USEPA CONTRACT LABORATORY PROGRAM

STATEMENT OF WORK FOR ORGANICS ANALYSIS

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Multi-Media, Multi-Concentration

Document Number 0LM01.0

STATEMENT OF WORK

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EPA 0657 1990 c.1

EXHIBIT A

SUMMARY OF REQUIREMENTS

SECTION I

GENERAL REQUIREMENTS

The Contractor shall use proven instruments and techniques to identify and measure the concentrations of volatile, semivolatile, and pesticide compounds listed on the Target Compound List (TCL) in Exhibit C. The Contractor shall employ state-of-the-art GC/MS and/or GC/EC procedures to perform all analyses, including the necessary preparations for analysis.

In Exhibit D, the EPA provides the Contractor with the specific analytical procedures to be used and defines the specific application of these procedures to this contract. For volatiles and semivolatiles, this includes instructions for sample preparation, gas chromatographic screening, mass spectrometric identification, and data evaluation. Specific ions used for searching the mass spectral data for each compound are included. For pesticides, this includes instructions for sample preparation, gas chromatography, confirmation of identification by gas chromatography and/or mass spectrometry, and data evaluation.

The Contractor shall prepare extracts and dilutions of samples. The Contractor shall screen extracts by methods of his choice (soil characterization mandatory; water characterization optional) at an initial extract concentration. Then, based on the screening response, the Contractor shall use the specific analytical methods described in Exhibit D to extract and concentrate samples to achieve the Contract Required Quantitation Limits (CRQL) listed in Exhibit C. Exhibit D lists the analytical methods and starting points to be utilized for each of the target compounds.

During preparation, the Contractor shall fortify all semivolatile and pesticide samples, blanks, matrix spikes, and matrix spike duplicates with the surrogate spiking compounds listed in Exhibit D. The Contractor shall fortify all volatile samples, blanks, matrix spikes, and matrix spike duplicates with the system monitoring compounds listed in Exhibit D. Additionally, all sample semivolatile extracts and aliquots for volatile organics analysis shall be spiked with the internal standard compounds listed in Exhibit D before injection or purging.

Additionally, for each sample analyzed by GC/MS, the Contractor shall conduct mass spectral library searches to determine the possible identity of up to ten (10) non-system monitoring compounds volatile components and up to twenty (20) non-surrogate semivolatile components that are not on the Target Compound List (Exhibit C).

Exhibit F contains chain-of-custody and sample documentation requirements which the Contractor must follow in processing samples under this contract, and specifies requirements for written laboratory standard operating procedures.

Sample analysis data, sample documentation and other deliverables shall be reported as specified in Exhibit B. Specifications for reporting data in computer-readable form appear in Exhibit H. To ensure proper understanding of language utilized in this contract, Exhibit G contains a glossary of terms. When a term is used in the text without explanation, the glossary meaning shall be applicable.

The samples to be analyzed by the Contractor are from known or suspected hazardous waste sites and, potentially, may contain hazardous organic and/or inorganic materials at high concentration levels. The Contractor should be aware of the potential hazards associated with the handling and analyses of these samples. It is the Contractor's responsibility to take all necessary measures to ensure the health and safety of its employees.

In addition, the Contractor must be aware of the importance of maintaining the integrity of the data generated under the contracts as it is used to make major decisions regarding public health and environmental welfare. In addition, it may be used in litigation against potentially responsible parties in the enforcement of Superfund legislation.

Prior to accepting any samples from the Agency, the Contractor shall have, in-house, the appropriate standards for <u>all</u> target compounds listed in Exhibit C.

SECTION II

SUMMARY OF REQUIREMENTS

- I. For each sample, the Contractor shall perform the following tasks:
 - A. Task I Receive and Prepare Hazardous Waste Samples.
 - 1. Receive and handle samples under the chain-of-custody procedures described in Exhibit F.
 - 2. Prepare samples as described in Exhibit D. VOA analysis of water or soil samples must be completed within 10 days of VTSR (Validated Time of Sample Receipt). Separatory funnel extractions for pesticides in water samples must be completed within 5 days of VTSR. Sonication extractions for pesticides and/or semivolatiles in soil samples must be completed within 5 days of VTSR. Continuous liquid-liquid extraction for semivolatile samples must be started within 5 days of VTSR.

Extracts of either water or soil samples must be analyzed within 40 days of VTSR. This does not release the Contractor from the data turnaround time specified in Exhibit B, Section I.

- B. Task II Analysis for Identification of Specific Organic Compounds.
 - 1. Extracts and aliquots prepared in Task I shall be analyzed by GC/EC and GC/MS techniques given in Exhibit D for the target compounds listed in Exhibit C.
 - 2. The target compounds listed in Exhibit C shall be identified as described in the methodologies given in Exhibit D. Automated computer programs may be used to facilitate the identification.
- C. <u>Task III</u> Qualitative Verification of the Compounds Identified in Task II.
 - 1. The volatile and semivolatile compounds analyzed by GC/MS techniques and initially identified in Task II shall be verified by an analyst competent in the interpretation of mass spectra by comparison of the suspect mass spectrum to the mass spectrum of a standard of the suspected compound. This procedure requires the use of multiple internal standards. Two criteria must be satisfied to verify the identifications:
 - a. Elution of the sample component at the same GC relative retention time as the standard component.
 - b. Correspondence of the sample component and standard component mass spectra.

2. For establishing correspondence of the GC relative retention time (RRT), the sample component RRT must compare within \pm 0.06 RRT units of the RRT of the standard component. For reference, the calibration standard must be run on the same 12-hour time period as the sample.

For comparison of standard and sample component mass spectra, mass spectra obtained on the Contractor's GC/MS are required. Once obtained, these standard spectra may be used for identification purposes only if the Contractor's GC/MS meets the DFTPP or BFB daily instrument performance check requirements specified in Exhibit D. The standard spectra used may be from a laboratory generated library on the same instrument or obtained from the calibration standard run used to obtain reference RRTs. The requirements for qualitative verification by comparison of mass spectra are as follows:

- a. All ions present in the standard mass spectrum at a relative intensity greater than 10 percent (most abundant ion in the spectrum equals 100 percent) must be present in the sample spectrum.
- b. The relative intensities of ions specified above must agree within ± 20 percent between the standard and sample spectra.
- c. Ions greater than 10 percent in the <u>sample</u> spectrum but not present in the <u>standard</u> spectrum must be considered and accounted for by the analyst making the comparison. When GC/MS computer data processing programs are used to obtain the sample component spectrum, both the processed and the raw spectra must be evaluated. In Task III, the verification process should favor false positives.
- 3. If a compound analyzed by GC/MS techniques and initially identified in Task II cannot be verified by all of the criteria in items 1 and 2 above, but in the technical judgement of the mass spectral interpretation specialist the identification is correct, then the Contractor shall report that identification, and proceed with quantification in Task IV.
- 4. The Pesticide/Aroclor compounds listed in Exhibit C and analyzed by GC/EC techniques shall have their identifications verified by an analyst competent in the interpretation of gas chromatograms. Two criteria must be satisfied to verify the identifications:
- a. Elution of the sample component within the retention time window (established by the procedures in Exhibit D) of the standard component analyzed on the same GC column and instrument, as specified in Exhibit D.

- b. Analysis of the sample and standard on a second GC column with a stationary phase with retention characteristics dissimilar to that used in a. above, and meeting the same criteria for elution of the sample component and the standard as in a. above.
- D. Task IV Quantification of Compounds Verified in Task III.
 - 1. The Contractor shall quantify components analyzed by GC/MS techniques, identified in Task II and verified in Task III by the internal standard method stipulated in Exhibit D. Where multiple internal standards are required by EPA, the Contractor shall perform quantitation utilizing the internal standards specified in Exhibit D.
 - 2. The Contractor shall determine response factors for each 12-hour time period of GC/MS analysis and shall include a calibration check of the initial five point calibration as described in Exhibit D.
 - 3. The Contractor shall quantify components analyzed by GC/EC techniques, identified in Task II and verified in Task III by the external standard method stipulated in Exhibit D.
 - 4. The Contractor shall perform an initial three-point calibration, verify its linearity, determine the breakdown of labile components, and determine calibration factors for all standards analyzed by GC/EC techniques as described in Exhibit D.
- E. Task V Tentative Identification of Non-target Sample Components.
 - For each analysis of a sample, the Contractor shall conduct 1. mass spectral library searches to determine tentative compound identifications as follows. For each volatile fraction, the Contractor shall conduct a search to determine the possible identity of up to ten (10) organic compounds of greatest concentration which are not system monitoring compounds and are not listed in Exhibit C. For each semivolatile fraction. the Contractor shall conduct a search to determine the possible identification of up to twenty (20) non-surrogate organic compounds of greatest concentration which are not listed in Exhibit C. In performing searches, the 1989 (or most recent) release of the NIST/EPA/MSDC mass spectral library (containing 50,000 spectra) must be used. NOTE: Substances with responses less than 10 percent of the nearest internal standard are not required to be searched in this fashion.
 - 2. Only after visual comparison of sample spectra with the spectra from the library searches will the mass spectral interpretation specialist assign a tentative identification. If the compound does not meet the identification criteria of Task III, it shall be reported as unknown. The mass spectral

specialist should give additional classification of the unknown compound, if possible (i.e., unknown aromatic, unknown hydrocarbon, unknown acid type, unknown chlorinated compound). If probable molecular weights can be distinguished, include them.

- 3. The Contractor shall <u>not</u> report as semivolatile tentatively identified compounds (TIC) any target compounds from the volatile fraction (i.e., do not report late eluting volatile compounds as TICs in the semivolatile analysis). However, the Contractor may report pesticide target compounds that appear as semivolatile tentatively identified compounds.
- F. Task VI Quality Assurance/Quality Control Procedures.
 - All specific quality assurance procedures prescribed in Exhibits D and E shall be strictly adhered to by the Contractor. Records documenting the use of the protocol shall be maintained in accordance with the document control procedures prescribed in Exhibit F, and shall be reported in accordance with Exhibit B, Reporting Requirements and Deliverables.
 - 2. The Contractor shall establish a Quality Assurance Plan (QAP) with the objective of providing sound analytical chemical measurements. This program shall incorporate the quality control procedures, any necessary corrective action, and all documentation required during data collection as well as the quality assessment measures performed by management to ensure acceptable data production.
 - 3. The Contractor shall perform one spiked sample analysis (matrix spike) and one duplicate spiked sample analysis (matrix spike duplicate) for each group of samples of a similar matrix (for water or soil samples) and concentration level (for volatile and semivolatile soil samples only) for the following, whichever is most frequent:
 - o Each Case of field samples received, OR
 - o Each 20 samples in a Case, OR
 - Each 14 calendar day period during which field samples in a Case were received (said period beginning with the receipt of the first sample in that Sample Delivery Group).

Matrix spikes and matrix spike duplicates shall be carried through the entire analytical process from extraction to final GC/MS or GC/EC analysis, including all Contract Performance/Delivery Requirements (see Contract Schedule).

- 4. The Contractor shall prepare and analyze one laboratory reagent blank (method blank) for each group of samples of a similar matrix (for water or soil samples), extracted by a similar method (separatory funnel, continuous liquid-liquid extraction, or sonication, as specified in Exhibit D), and a similar concentration level (for volatile and semivolatile soil samples only) for the following, whichever is most frequent:
 - o Each Case of field samples received, OR
 - o Each 20 samples in a Case, including matrix spikes and reanalyses, OR
 - Each 14 calendar day period during which field samples in a Case were received (said period beginning with the receipt of the first sample in that Sample Delivery Group), OR
 - o Whenever samples are extracted.

Volatile analysis requires one method blank for each 12-hour time period when volatile target compounds are analyzed.

Semivolatile and pesticide method blanks shall be carried through the entire analytical process from extraction to final GC/MS or GC/EC analysis, including all Contract Performance/Delivery Requirements (see Contract Schedule).

5. The Contractor shall verify instrument performance for each 12-hour time period, to include the following: Decafluorotriphenylphosphine (DFTPP) and/or Bromofluorobenzene (BFB) as applicable, and a specific calibration using standards of defined concentration to monitor response, retention time, and mass spectra.

Additional quality control shall be conducted in the form of the analysis of Performance Evaluation check samples submitted to the laboratory by the Agency. The results of all such control or PE check samples may be used as grounds for termination of noncompliant Contractors. "Compliant performance" is defined as that which yields correct compound identification and concentration values as determined by the Agency, as well as meeting the contract requirements for analysis (Exhibit D), quality assurance/quality control (Exhibit E), data reporting and other deliverables (Exhibits B and H), and sample custody, sample documentation and SOP documentation (Exhibit F).

B. The EPA has provided to the Contractor formats for the reporting of data (Exhibits B and H). The Contractor shall be responsible for completing and returning analysis data sheets and submitting computer-readable data on diskette in the format specified in this SOW and within the time specified in the Contract Performance/Delivery Schedule.

- 1. Use of formats other than those designated by EPA will be deemed as noncompliance. Such data are unacceptable. Resubmission in the specified format at no additional cost to the government will be required.
- 2. Computer generated forms may be submitted in the hardcopy data package(s) provided that the forms are in EXACT EPA FORMAT. This means that the order of data elements is the same as on each EPA required form, including form numbers and titles, page numbers and header information.
- 3. The data reported by the Contractor on the hardcopy data forms and the associated computer-readable data submitted by the Contractor must contain identical information. If during government inspection discrepancies are found, the Contractor shall be required to resubmit either or both sets of data at no additional cost to the government.
- C. The Contractor shall provide analytical equipment and technical expertise for this contract as specified by the following:
 - 1. The Contractor shall have sufficient gas chromatograph (GC) and gas chromatograph/mass spectrometer/data system (GC/MS/DS) capability to meet all the terms and conditions of the Contract. The Contractor shall maintain, at a minimum, all analytical equipment allocated for this contract at the time of contract award.
 - 2. The Contractor's instrument systems shall have the following:
 - a. The GC/MS shall be equipped with a glass jet separator when using packed columns.
 - b. The computer shall be interfaced by hardware to the mass spectrometer and be capable of acquiring continuous mass scans for the duration of the chromatographic program.
 - c. The computer shall be equipped with mass storage devices for saving all data from the GC/MS runs.
 - d. Computer software shall be available to allow searching GC/MS runs for specific ions and plotting the intensity of the ions with respect to time or scan number.
 - e. A computer data system must be interfaced to the mass spectrometer that allows the continuous acquisition and storage, on machine readable media, of all mass spectra obtained throughout the duration of the chromatographic program. The computer must have software that allows searching any GC/MS data file for ions of a specified mass and plotting such ion abundances versus time or scan number. This type of plot is defined as an Extracted Ion Current Profile (EICP). Software must also be available that allows integrating the abundance in any EICP between specified time or scan number limits. Also, for the non-target compounds,

software must be available that allows for the comparison of sample spectra against reference library spectra. The 1989 (or most recent) release of the NIST/EPA/MSDC mass spectral library shall be used as the reference library. The data system must be capable of flagging all data files that have been edited manually by laboratory personnel.

- f. The GC/MS shall be equipped with a GC to MS interface capable of extending a fused silica capillary column into the ion source. The column is to be 30 meters long by 0.25 or 0.32 mm inside diameter, bonded DB-5, fused silica, or equivalent.
- g. The GC for pesticide analysis shall be equipped with wide bore capillary columns and a suitable detector as described in Exhibit D.
- 3. The Contractor shall use a magnetic tape storage device capable of recording data and suitable for long-term, off-line storage. The Contractor shall retain all raw GC/MS data acquired under this contract on magnetic tape in appropriate instrument manufacturer's format. The Contractor is required to retain the magnetic tapes with associated hardcopy tape logbook identifying tape contents (see Exhibit B) for 365 days after data submission. During that time, the Contractor shall submit tapes and logbook within 7 days of request, as specified in the Contract Performance/Delivery Schedule.
- 4. The Contractor shall have a computerized MS library search system capable of providing a forward comparison, utilizing the standard spectra contained in the mass spectral library. The 1989 (or most recent) release of the NIST/EPA/MSDC mass spectral library, containing 50,000 spectra, must be used.
 - a. The system shall provide a numerical ranking of the standard spectra most closely corresponding to the sample spectra examined.
 - b. The data system shall have software capable of removing background signals from spectra.
- 5. The Contractor shall have, in-house and operable, a device capable of analyzing purgeable organics as described in Exhibit D.
- 6. The Contractor shall have, in-house, the appropriate standards for <u>all</u> target compounds listed in Exhibit C prior to accepting any samples from the Agency. Standards provided by the EPA for use in the Preaward Performance Evaluation may not contain all the target compounds and thus must <u>not</u> be used for routine analyses unless or until they have been supplemented with commercially-available standard materials.

- D. The Contractor shall have an IBM or IBM-compatible mini-computer or PC capable of recording required sample data on 5.25 inch floppy double-sided double-density 360 K-byte or 1.2 M-byte or a 3.5 inch double-sided, double density 720 K-byte or 1.44 M-byte diskette, in ASCII text file format, and in accordance with the file, record and field specifications listed in Exhibit H.
- E. The Contractor shall designate and utilize key personnel to perform the minimum functional requirements necessary to meet the terms and conditions of this contract. The EPA reserves the right to review personnel qualifications and experience. The minimum functional requirements are listed below:
 - GC/MS Laboratory Supervisor
 - GC/EC Laboratory Supervisor
 - Sample Preparation Laboratory Supervisor
 - Quality Assurance Officer
 - o Systems Manager
 - Programmer Analyst
 - GC/MS Operator
 - Mass Spectral Interpretation Specialist
 - GC/EC Operator
 - Pesticide Residue Analysis Expert.
 - Extraction/Concentration Expert
- F. The Contractor shall respond within seven days to requests from data recipients for additional information or explanations that result from the Government's inspection activities.
- G. The Contractor shall preserve all sample extracts after analysis in bottles/ vials with Teflon-lined septa and shall maintain stored extracts at 4°C (\pm 2°C). The Contractor is required to retain the sample extracts for 365 days after data submission. During that time, the Contractor shall submit the extracts within 7 days after request, as specified in the Contract Performance/Delivery Schedule.
- H. The Contractor shall adhere to chain-of-custody procedures described in Exhibit F. Documentation, as described therein, shall be required to show that all procedures are being strictly fellowed. This documentation shall be reported as the complete SDG file purge (see Exhibit B).

I. Sample shipments to the Contractor's facility will be scheduled and coordinated by the EPA CLP Sample Management Office (SMO) acting on behalf of the Administrative Project Officer. The Contractor shall communicate with SMO personnel by telephone, as necessary throughout the process of sample scheduling, shipment, analysis and data reporting, to ensure that samples are properly processed.

If there are problems with the samples (e.g., mixed media, containers broken or leaking) or sample documentation/paperwork (e.g., Traffic Reports not with shipment, sample and Traffic Report numbers do not correspond) the Contractor shall immediately contact SMO for resolution. The Contractor shall immediately notify SMO regarding any problems and laboratory conditions that affect the timeliness of analyses and data reporting. In particular, the Contractor shall notify SMO personnel in advance regarding sample data that will be delivered late and shall specify the estimated delivery date.

J. Sample analyses will be scheduled by groups of samples, each defined as a Case and identified by a unique EPA Case number assigned by SMO. A Case signifies a group of samples collected at one site or geographical area over a finite time period, and will include one or more field samples with associated blanks. Samples may be shipped to the Contractor in a single shipment or multiple shipments over a period of time, depending on the size of the Case.

A Case consists of one or more Sample Delivery Group(s). A Sample Delivery Group (SDG) is defined by the following, whichever is most frequent:

- o Each Case of field samples received, OR
- o Each 20 field samples within a Case, OR
- Each 14 calendar day period during which field samples in a Case are received (said period beginning with the receipt of the first sample in the Sample Delivery Group).

Samples may be assigned to Sample Delivery Groups by matrix (i.e., all soils in one SDG, all waters in another), at the discretion of the laboratory. Such assignment must be made at the time the samples are received, and may not be made retroactively.

All data (hardcopy and computer-readable format) for all samples in a Sample Delivery Group are due concurrently to all data recipients as stipulated in the Delivery Schedule in Exhibit B. Section I. Data for all samples in a Sample Delivery Group must be submitted together (in one package) in the order specified in Exhibit B. The Sample Delivery Group number is the EPA sample number of the first sample received in the SDG. When several samples are received together in the first SDG shipment, the SDG number shall be the lowest sample number (considering both alpha and numeric designations) in the first group of samples received under the SDG. The SDG number is reported on all data reporting forms. The SDG Receipt Date is the day the last sample in the SDG is received. Data for all samples in the SDG are due as stipulated in the Delivery Schedule in Exhibit B.

The Contractor is responsible for identifying each Sample Delivery Group as samples are received, through proper sample documentation (see Exhibit B) and communication with SMO personnel.

K. Each sample received by the Contractor will be labeled with an EPA sample number, and accompanied by a Traffic Report form bearing the sample number and descriptive information regarding the sample. The Contractor shall complete and sign the Traffic Report, recording the date of sample receipt and sample condition on the receipt for each sample container.

The Contractor shall submit signed copies of Traffic Reports for all samples in a Sample Delivery Group to SMO within <u>3 calendar days</u> following receipt of the last sample in the Sample Delivery Group. Traffic Reports shall be submitted in Sample Delivery Group sets (i.e., all Traffic Reports for a Sample Delivery Group shall be clipped together) with an SDG Cover Sheet containing information regarding the Sample Delivery Group, as specified in Exhibit B.

- L. EPA Case numbers (including SDG numbers) and EPA sample numbers shall be used by the Contractor in identifying samples received under this contract both verbally and in reports/correspondence.
- M. Samples will be shipped routinely to the Contractor through an overnight delivery service. However, as necessary, the Contractor shall be responsible for any handling or processing required for the receipt of sample shipments, including pick-up of samples at the nearest servicing airport, bus station or other carrier service within the Contractor's geographical area. The Contractor shall be available to receive sample shipments at any time the delivery service is operating, including Saturdays.
- N. The Contractor shall accept all samples scheduled by SMO, provided that the total number of samples received in any calendar month does not exceed the monthly limitation expressed in the contract. Should the Contractor elect to accept additional samples, the Contractor shall remain bound by all contract requirements for analysis of those samples accepted.

SECTION III

DETAILED TECHNICAL & MANAGEMENT REQUIREMENTS

As cited in Section II the Contractor shall have the following technical and management capabilities:

I. TECHNICAL CAPABILITY

A. <u>Technical Functions</u>

- 1. GC/MS Laboratory Supervisor
 - a. Responsible for all technical efforts of the GC/MS laboratory to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

(2) Experience:

Minimum of three years of laboratory experience, including at least one year of supervisory experience.

- 2. GC/EC Laboratory Supervisor
 - a. Responsible for all technical efforts of the GC/EC laboratory to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

(2) Experience:

Minimum of three years of laboratory experience, including at least one year of supervisory experience.

- 3. Sample Preparation Laboratory Supervisor
 - a. Responsible for all technical efforts of sample preparations to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

(2) Experience:

Minimum of three years of laboratory experience, including at least one year of supervisory experience.

- 4. Quality Assurance Officer
 - a. Responsible for overseeing the quality assurance aspects of the data and reporting directly to upper management to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

(2) Experience:

Minimum of three years of laboratory experience, including at least one year of applied experience with QA principles and practices in an analytical laboratory.

- 5. Systems Manager
 - a. Responsible for the management and quality control of all computing systems (hardware, software, documentation and procedures), generating, updating, and quality controlling automated deliverables to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree with four or more intermediate courses in programming, information management, database management systems, or systems requirements analysis. (2) Experience:

Minimum of three years experience in data or systems management or programming including one year of experience with the software being utilized for data management and generation of deliverables.

- 6. Programmer Analyst
 - a. Responsible for the installation, operation and maintenance of software and programs generating, updating and quality controlling analytical databases and automated deliverables to meet all terms and conditions of the EPA contract.
 - b. Qualifications:
 - (1) Education:

Minimum of Bachelor's degree with four or more intermediate courses in programming, information management, information systems, database management systems, or systems requirements analysis.

(2) Experience:

Minimum of two years experience in systems or applications programming including one year of experience with the software being utilized for data management and generation of deliverables.

7. GC/MS Operator Qualifications

One year of experience in operating and maintaining GC/MS/DS with a Bachelor's degree in chemistry or a scientific/engineering discipline, or in lieu of the Bachelor's degree, three years of experience in operating and maintaining the GC/MS and interpreting GC/MS data.

- 8. Mass Spectral Interpretation Specialist Qualifications
 - a. Education:
 - o Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.
 - o Training course(s) in mass spectral interpretation.
 - b. Experience:

Minimum of two years of experience in mass spectral interpretation.

9. GC/EC Operator Qualifications

One year of experience in operating and maintaining GC/EC with a Bachelor's degree in chemistry or a scientific/engineering discipline, or in lieu of the Bachelor's degree, three years of experience in operating and maintaining the GC/EC and interpreting GC/EC data.

- 10. Pesticide Residue Analysis Expert Qualifications
 - a. Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

b. Experience:

Minimum of two years of experience in operating and maintaining GC and interpreting GC chromatograms.

- 11. Extraction/Concentration Expert Qualifications
 - a. Education:

Minimum of High school diploma and a college level course in general chemistry.

b. Experience:

Minimum of one year of experience in extraction/concentration.

12. Technical Staff Redundancy

The bidder shall have a minimum of one (1) chemist available at any one time as a back-up technical person with the following qualifications, to ensure continuous operations to accomplish the required work as specified by EPA contract.

a. Education:

Minimum of Bachelor's degree in chemistry or any scientific/engineering discipline.

- b. Experience: Minimum of one year in each of the following areas
 - o GC operation and maintenance for volatiles and semivolatiles analyses.
 - o Mass spectral interpretation.
 - o Extraction.
 - o Pesticide/ Aroclors analysis.

B. Facilities

The adequacy of the facilities and equipment is of equal importance as the technical staff to accomplish the required work as specified by the EPA contract.

1. Sample Receipt Area

Adequate, contamination-free, well ventilated work space provided with chemical resistant bench top for receipt and safe handling of EPA samples.

2. Storage Area

Sufficient refrigerator space to maintain unused EPA sample volume for 60 days after data submission and sample extracts for 365 days after data submission. Samples must be stored in an atmosphere demonstrated to be free from all potential contaminants. Volatile samples must be stored in a refrigerator used only for storage of volatile samples from this contract.

3. Sample Preparation Area

Adequate, contamination-free, well-ventilated work space provided with the following:

- a. Benches with chemical resistant tops, exhaust hoods. NOTE: Standards must be prepared in a glove box or isolated area.
- b. Source of distilled or demineralized organic-free water.
- c. Analytical balance(s) located away from draft and rapid change in temperature.

C. <u>Instrumentation</u>

At a minimum, the Contractor shall have the following instruments operative at the time of the Preaward Site Evaluation and committed for the full duration of the contract. 1. Primary Instrument Requirements for up to 200 Samples/Month Capacity

 Fraction	No. of Instrument(s)	Type of Instrument
 Volatiles	1	GC/MS/DS with purge and trap device
Semivolatiles	2	GC/MS/DS
Pesticides/ Aroclors	2	GC/EC with dual column

NOTE: For 300-400 Samples/Month Capacity, twice as much instrumentation is needed as is listed in item 1.

2. Secondary Instrument Requirements for up to 200 Samples/Month Capacity

The Contractor shall have the following instruments in place and operational at any one time as a back-up system;

Ouentity	Instruments
One	GC/MS/DS
One	Purge and Trap Device
One	GC

These instruments must be included in the bidder's inventory of equipment along with those in 1, above.

In addition, the Contractor shall have an in-house stock of instrument parts and circuit boards to ensure continuous operation to meet contract-specified holding and turn-around times.

3. Instrument Specifications

Instrument specifications are described in detail in the Statement of Work (SOW) in the following Exhibits:

0	Purge and	trap	device	Exhibit	D	
0	GC/MS/DS			Exhibit	D	
0	GC/EC			Exhibit	D	

D. Data Management and Handling

1. Hardware - Contractor shall have an IBM or IBM-compatible minicomputer or PC capable of recording required sample data on 5.25 inch double-sided, double-density 360 K-byte or high density 1.2 Mbyte diskettes; or 3.5 inch double-sided, double-density 720 K-byte or 1.44 M-byte diskettes in ASCII text file format and in accordance with the file, record and field specifications listed in SOW, Exhibit H.

Other minimum requirements include:

- o Hard disk of at least 20 M-bytes.
- o Synchronous modem capable of at least 4,800 baud transmission speed.
- o PC hardware and software enabling IBM SNA 3270 emulation.
- 2. Software Software, utilized in generating, updating and quality controlling analytical databases and automated deliverables shall have the following additional capabilities:
 - o Editing and updating databases.
 - o QC of automated deliverables.
 - o Controlled access using user ID and file password protection.
- 3. The Contractor shall also be able to submit reports and data packages as specified in the Statement of Work Exhibit B. To complete this task, the Contractor shall be required to provide space, tables and adequate copy machines to meet the contract requirements.
- II. LABORATORY MANAGEMENT CAPABILITY

The Contractor must have an organization with well-defined responsibilities for each individual in the management system to ensure sufficient resources for the EPA contract(s) and to maintain a successful operation. To establish this capability, the Contractor shall designate personnel to carry out the following responsibilities for the EPA contract. Functions include, but are not limited to, the following:

A. <u>Technical Staff</u>

Responsible for all technical efforts for the EPA contract.

B. <u>Project Manager</u>

Responsible for overall aspects of EPA contract(s) (from sample receipt through data delivery) and shall be the primary contact for EPA Headquarters Administrative Project Officer and Regional Technical Project Officers.

C. <u>Sample Custodian</u>

Responsible for receiving the EPA samples (logging, handling and storage).

D. <u>Ouality Assurance Officer</u>

Responsible for overseeing the quality assurance aspects of the data and reporting directly to upper management.

E. Data Reporting and Delivery Officer

Responsible for all aspects of data deliverables: organization, packaging, copying, and delivery.

EXHIBIT B

REPORTING AND DELIVERABLES REQUIREMENTS

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SECTION I

CONTRACT REPORTS/DELIVERABLES DISTRIBUTION

The following table reiterates the Contract reporting and deliverables requirement specified in the Contract Schedule and specifies the distribution that is require for each deliverable. NOTE: Specific recipient names and addresses are subject change during the term of the contract. The Administrative Project Officer will notify the Contractor in writing of such changes when they occur.

		No.	Delivery	Di	strib	ution
	Item	Copies	Schedule	(1)	(2)	(3)
Α.	Updated SOPs	2	45 days after contract receipt.		x	x
*B.	Sample Traffic Reports	1	3 days after receipt of last sample in Sample Delivery Group (SDG).**	X		
***C.	Sample Data Summary Package	1	35 days after receipt of last sample in SDG.	X		
***D、	Sample Data Package	2	35 days after receipt of last sample in SDG.	X		x
E. *	Complete SDG File	1	35 days after receipt of last sample in SDG.		X	
*****F.	Quality Assurance Plan	1	Submit copy within 7 days of written request by APO	A	s dir	ected
G.	Data in Computer- Readable Form	1	35 days after receipt of last sample in SDG.	X		

Item	No. <u>Copies</u>	Delivery Schedule	Distribution (1) (2) (3)
H. GC/MS Tapes	Lot	Retain for 365 days after data submis- sion, or submit with- in 7 days after receipt of written request by APO and/or EMSL/LV.	
I. Extracts	Lot	Retain for 365 days after data submis- sion, or submit with- in 7 days after receipt of written request by APO or SMO	

Distribution:

(1) Sample Management Office (SMO)

(2) Region-Client (Technical Project Officer)

(3) EMSL-LV

* Also required in the Sample Data Summary Package.

- ** Sample Delivery Group (SDG) is a group of samples within a Case, received over a period of 14 days or less and not exceeding 20 samples. Data for all samples in the SDG are due concurrently. The date of delivery of the SDG or any samples within the SDG is the date that all samples have been delivered. (See Exhibit A for further description).
- <u>Concurrent</u> delivery required. Delivery shall be made such that all designated recipients receive the item on the same calendar day.
- **** Complete SDG File will contain the original sample data package plus all of the original documents described under Complete SDG File paragraph E.

***** See Exhibit E for a more detailed description.

NOTE: As specified in the Contract Schedule (G.6 Government Furnished Supplies and Materials), unless otherwise instructed by the CLP Sample Management Office, the Contractor shall dispose of unused sample volume and used sample bottles/containers no earlier than sixty (60) days following submission of analytical data.

Distribution Addresses:

- USEPA Contract Lab Program Sample Management Office (SMO)
 P. O. Box 818
 Alexandria, VA 22313
 For overnight delivery service, use street address: 300 North Lee Street Alexandria, VA 22313
- (2) USEPA Environmental Monitoring Systems Laboratory (EMSL-LV) P. O. Box 93478 Las Vegas, NV 89193-3478 ATTN: Data Audit Staff

For overnight delivery service, use street address: 944 E. Harmon, Executive Center Las Vegas, NV 89109 ATTN: Data Audit Staff

(3) USEPA REGIONS:

The CLP Sample Management Office, acting on behalf of the Administrative Project Officer, will provide the Contractor with the list of addressees for the ten EPA Regions. SMO will provide the Contractor with updated Regional address/name lists as necessary throughout the period of the contract and identify other client recipients on a case-by-case basis.

SECTION II

REPORT DESCRIPTIONS AND ORDER OF DATA DELIVERABLES

The Contractor laboratory shall provide reports and other deliverables as specified in the Contract Schedule (Performance/Delivery Schedule, Section F.1). The required content and form of each deliverable is described in this Exhibit.

All reports and documentation MUST BE as follows:

- o Legible,
- o Clearly lábeled and completed in accordance with instructions in this Exhibit,
- o Arranged in the order specified in this Section, and
- o Paginated.

If submitted documentation does not conform to the above criteria, the Contractor will be required to resubmit such documentation with deficiency(ies) corrected, at no additional cost to the Agency.

Whenever the Contractor is required to submit or resubmit data as a result of an on-site laboratory evaluation or through a APO/TFO action, the data must be clearly marked as ADDITIONAL DATA and must be sent to all three contractual data recipients (SMO, EMSL/LV, and Region). A cover letter shall be included which describes what data is being delivered, to which EPA Case(s) it pertains, and who requested the data.

Whenever the Contractor is required to submit or resubmit data as a result of Contract Compliance Screening (CCS) review by SMO, the data must be sent to all three contractual data recipients (SMO, EMSL/LV and Region), and in all three instances must be accompanied by a color-coded COVER SHEET (Laboratory Response To Results of Contract Compliance Screening) provided by SMO.

Section III of this Exhibit contains forms instructions to assist the Contractor in accurately providing the Agency with all required data. Section IV contains copies of the required data reporting forms in Agency-specified formats. Data elements with field parameters for reporting data in computer readable form are contained in Exhibit H.

Descriptions of the requirements for each deliverable item cited in the Contract Performance/Delivery Schedule (Contract Schedule, Section F.1) are specified in A-H of this Section. Items submitted concurrently MUST BE arranged in the order listed. Additionally, the components of each item MUST BE arranged in the order presented in this Section when the item is submitted.

Examples of specific data deliverables not included herein may be obtained by submitting a written request to the EPA Administrative Project Officer, stating the information requested, and signed by the Laboratory Manager.

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A. Updated SOPs

The Contractor shall submit updated copies of all required Standard Operating Procedures (SOPs) that were submitted with the prebid Performance Evaluation sample results. The updated SOPs must address any and all issues of laboratory performance and operation identified through the review of the Performance Evaluation sample data and the evaluation of Bidder-Supplied Documentation.

The Contractor must supply SOPs for the following:

- 1. Sample receipt and logging.
- 2. Sample storage.
- 3. Preventing sample contamination.
- 4. Security for laboratory and samples.
- 5. Standards purity/preparation.
- 6. Maintaining instrument records and logbooks.
- 7. Sample analysis and data control systems.
- 8. Glassware cleaning.
- 9. Technical and managerial review of laboratory operation and data package preparation.
- 10. Internal review of contractually-required quality assurance and quality control data for each individual data package.
- 11. Sample analysis, data handling and reporting.
- 12. Chain-of-Custody procedures and document control including SDG file preparation.
- 13. Laboratory data validation/laboratory selfinspection system.
- 14. Data Management and Handling

B. Sample Traffic Reports

Original Sample Traffic Report page marked "Lab Copy for Return to SMO" with lab receipt information and signed in original Contractor signature, for each sample in the Sample Delivery Group.

Traffic Reports (TRs) shall be submitted in Sample Delivery Group (SDG) sets (i.e., TRs for all samples in an SDG shall be clipped together), with an SDG Cover Sheet attached.

The SDG Cover Sheet shall contain the following items:

- o Lab name.
- o Contract number.
- o Sample Analysis Price full sample price from contract.
- o Case Number.
- List of EPA sample numbers of all samples in the SDG, identifying the <u>first</u> and <u>last</u> samples received, and their dates of receipt (LRDs).
 NOTE: When more than one sample is received in the first or last SDG shipment, the "first" sample received would be the lowest sample number (considering both alpha and numeric designations); the "last" sample received would be the highest sample number (considering both alpha and numeric designations).

In addition, <u>each</u> Traffic Report must be clearly marked with the SDG Number, the sample number of the first sample in the SDG (as described in the following paragraph). This information should be entered below the Lab Receipt Date on the TR. In addition, the TR for the <u>last</u> sample received in the SDG must be clearly marked "SDG - FINAL SAMPLE."

The EPA sample number of the first sample received in the SDG is the SDG number. When several samples are received together in the first SDG shipment, the SDG number shall be the lowest sample number (considering both alpha and numeric designations) in the first group of samples received under the SDG. The SDG number is also reported on all data reporting forms (see Section III, Forms Instruction Guide).

If samples are received at the laboratory with multi-sample Traffic Reports (TRs), all the samples on one multi-sample TR may not necessarily be in the same SDG. In this instance, the laboratory must make the appropriate number of photocopies of the TR, and submit one copy with each SDG cover sheet.

C. <u>Sample Data Summary Package</u>

As specified in the Delivery Schedule, one Sample Data Summary Package shall be delivered to SMO concurrently with delivery of other required sample data. The Sample Data Summary Package consists of copies of specified items from the Sample Data Package. These items are listed below and are described under part D, Sample Data Package.

The Sample Data Summary Package shall be ordered as follows and shall be submitted separately (i.e., separated by rubber bands, clips or other means) directly <u>preceding</u> the Sample Data Package. Sample data forms shall be arranged in increasing EPA sample number order, considering <u>both</u> letters and numbers. For example, BE400 is a lower sample number than BF100, as E precedes F in the alphabet.

The Sample Data Summary Package shall contain data for samples in one Sample Delivery Group of the Case, as follows:

- 1. Case Narrative
- 2. By fraction (VOA, SV, PEST) and by sample within each fraction tabulated target compound results (Form I) and tentatively identified compounds (Form I, TIC) (VOA and SV only)
- 3. By fraction (VOA, SV, PEST) surrogate spike analysis results (Forma II) by matrix (water and/or soil) and for soil, by concentration (low or medium)
- 4. By fraction (VOA, SV, PEST) matrix spike/matrix spike duplicate results (Form III)
- 5. By fraction (VOA, SV, PEST) blank data (Form IV) and tabulated results (Form I) including tentatively identified compounds (Form I, TIC) (VOA and SV only)
- 6. By fraction (VOA, SV only) internal standard area data (Form VIII)

D. Sample Data Package

The Sample Data Package is divided into the five major units described below. The last three units are each specific to an analytical fraction (volatiles, semivolatiles, pesticides/Aroclors). If the analysis of a fraction is not required, then that fraction-specific unit is not required as a deliverable.

The Sample Data Package shall include data for analyses of all samples in one Sample Delivery Group, including field samples, reanalyses, blanks, matrix spikes, and matrix spike duplicates.

The Contractor shall retain a copy of the sample Data Package for 365 days after final acceptance of data. After this time, the Contractor may dispose of the package.

1. SDG Narrative

This document shall be clearly labeled "SDG Narrative" and shall contain: laboratory name; Case number; sample numbers in the Sample Delivery Group (SDG), differentiating between initial analyses and re-analyses; SDG number; Contract number; and detailed documentation of any quality control, sample, shipment and/or analytical problems encountered in processing the samples reported in the data package.

Whenever data from sample re-analyses are submitted, the Contractor shall state in the SDG Narrative for <u>each</u> re-analysis, whether it considers the re-analysis to be billable, and if so, why.

The Contractor must also include any problems encountered: both technical and administrative, the corrective actions taken, and resolution and an explanation for all flagged edits (i.e., manual edits) on quantitation lists.

The SDG Narrative shall contain the following statement, <u>verbatim</u>: "I certify that this data package is in compliance with the terms and conditions of the contract, both technically and for completeness, for other than the conditions detailed above. Release of the data contained in this hardcopy data package and in the computer-readable data submitted on diskette has been authorized by the Laboratory Manager or his designee, as verified by the following signature." This statement shall be directly followed by signature of the Laboratory Manager or his designee with a typed line below it containing the signer's name and title, and the date of signature.

Additionally, the SDG Narrative itself must be signed in original signature by the Laboratory Manager or his designee and dated.

2. Traffic Reports

A copy of the Sample Traffic Reports submitted in Item A for all of the samples in the SDG. The Traffic Reports shall be arranged in increasing EPA sample number order, considering <u>both</u> letters and numbering in ordering samples. Copies of the SDG cover sheet is to be included with the copies of the Traffic Reports.

If samples are received at the laboratory with multi-sample Traffic Reports (TRs), all the samples on one multi-sample TR may not necessarily be in the same SDG. In this instance, the laboratory must make the appropriate number of photocopies of the TR so that a copy is submitted with each data package to which it applies. In addition, in any instance where samples from more than one multisample TR are in the same data package, the laboratory must submit a copy of the SDG cover sheet with copies of the TRs.

- 3. Volatiles Data
 - a. QC Summary
 - (1) System Monitoring Compound Summary (Form II VOA)
 - (2) Matrix Spike/Matrix Spike Duplicate Summary (Form III VOA)
 - (3) Method Blank Summary (Form IV VOA)

(If more than a single form is necessary, forms must be arranged in chronological order by date of analysis of the blank.)

(4) GC/MS Instrument Performance Check (Form V VOA)

In chronological order; by instrument.

(5) Internal Standard Area and RT Summary (Form VIII VOA)

In chronological order; by instrument.

b. Sample Data

Sample data shall be arranged in packets with the Organic Analysis Data Sheet (Form I VOA, including Form I VOA-TIC), followed by the raw data for volatile samples. These sample packets should then be placed in increasing EPA sample number order, considering both letters and numbers in ordering samples.

(1) Target Compound Results - Organic Analysis Data Sheet (Form I VOA).

Tabulated results (identification and quantitation) of the specified target compounds (Exhibit C). The validation and release of these results is authorized by a specific, signed statement in the SDG Narrative (reference E.1). In the event that the Laboratory Manager cannot validate all data reported for each sample, the Laboratory Manager shall provide a detailed description of the problems associated with the sample in the SDG Narrative.

(2) Tentatively Identified Compounds (Form I VOA-TIC).

This form must be included even if no compounds are found. If so, indicate this on the form by entering "O" in the field for "Number found."

Form I VOA-TIC is the tabulated list of the highest probable match for up to 10 organic compounds that are not system monitoring compounds and are not listed in Exhibit C (TCL). It includes the CAS (Chemical Abstracts Service) Registry Number, tentative identification, and estimated concentration.

(3) Reconstructed total ion chromatograms (RIC) for each sample or sample extract.

RICs must be normalized to the largest nonsolvent component, and must contain the following header information:

- EPA sample number.
- Date and time of analysis.
- GC/MS instrument ID.
- Lab file ID.

Internal standards and system monitoring compounds are to be labeled with the names of compounds, either directly out from the peak, or on a print-out of retention times if retention times are printed over the peak. If automated data system procedures are used for preliminary identification and/or quantification of the target compounds, the complete data system report must be included in all sample data packages, in addition to the reconstructed ion chromatogram. The complete data system report shall include all of the information listed below. For laboratories which do not use the automated data system procedures, a laboratory "raw data sheet," containing the following information, must be included in the sample data package in addition to the chromatogram.

- o EPA sample number.
- o Date and time of analysis.
- o RT or scan number of identified target compounds.
- o Ion used for quantitation with measured area.
- o Copy of area table from data system.
- o GC/MS instrument ID.
- o Lab file ID.

In all instances where the data system report has been edited, or where manual integration or quantitation has been performed, the GC/MS operator must identify such edits or manual procedures by initialing and dating the changes made to the report.

- (4) For each sample, by each compound identified, the following shall be included in the data package:
 - (a) Copies of raw spectra and copies of background-subtracted mass spectra of target compounds listed in Exhibit C (TCL) that are identified in the sample and corresponding background-subtracted target compound standard mass spectra. Spectra must be labeled with EPA sample number, lab file ID, date and time of analysis, and GC/MS instrument ID; compound names must be clearly marked on all spectra.
 - (b) Copies of mass spectra of organic compounds not listed in Exhibit C (TCL), Tentatively Identified Compounds, with associated best-match spectra (three best matches), labeled as in (4)(a) above.

c. Standards Data

- (1) Initial Calibration Data (Form VI VOA) in order by instrument, if more than one instrument used.
 - (a) VOA standard(s) reconstructed ion chromatograms and quantitation reports (or legible facsimile) for the

initial (five point) calibration, labeled as in b.(3) above. Spectra are not required.

- (b) All initial calibration data that pertain to samples in the data package must be included, regardless of when it was performed and for which Case. When more than one initial calibration is performed, the data must be put in chronological order, by instrument.
- (2) Continuing Calibration (Form VII VOA) in order by instrument, if more than one instrument used.
 - (a) VOA standard(s) reconstructed ion chromatograms and quantitation reports (or legible facsimile) for all continuing (12 hour) calibrations, labeled as in b.(3) above. Spectra are not required.
 - (b) When more than one continuing calibration is performed, forms must be in chronological order, within fraction and instrument.

d. Raw QC Data

- (1) BFB (for each 12-hour period, for each GC/MS system utilized)
 - (a) Bar graph spectrum, labeled as in b. (3) above.
 - (b) Mass listing, labeled as in b. (3) above.
 - (c) Reconstructed total ion chromatogram (RIC), labeled as in b.(3) above.
- (2) Blank Data in chronological order. NOTE: This order is different from that used for samples.
 - (a) Tabulated results (Form I VOA).
 - (b) Tentatively Identified Compounds (Form I VOA-TIC) even if none found.
 - (c) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b.(3) above.
 - (d) Target compound spectra with lab generated standard, labeled as in b.(4) above. Data systems which are incapable of dual display shall provide spectra in the following order:
 - o Raw target compound spectra.
 - o Enhanced or background subtracted spectra.
 - o Laboratory generated standard spectra.

- (e) GC/MS library search spectra for Tentatively Identified Compounds (TIC), labeled as in b.(4) above.
- (f) Quantitation/Calculation of Tentatively Identified Compound(s) (TIC) concentrations.
- (3) Matrix Spike Data
 - (a) Tabulated results (Form I VOA) of target compounds. Form I VOA-TIC not required.
 - (b) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b.(4) above. Spectra not required.
- (4) Matrix Spike Duplicate Data
 - (a) Tabulated results (Form I VOA) of target compounds. Form I VOA-TIC not required.
 - (b) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b.(4) above. Spectra not required.

4. Semivolatiles Data

- a. QC Summary
 - (1) Surrogate Percent Recovery Summary (Form II SV)
 - (2) Matrix Spike/Matrix Spike Duplicate Summary (Form III SV)
 - (3) Method Blank Summary (Form IV SV)

(If more than a single form is necessary, forms must be arranged in chronological order by date of analysis of the blank.)

(4) GC/MS Instrument Performance Check (Form V SV)

In chronological order; by instrument.

(5) Internal Standard Area and RT Summary (Form VIII SV)

In chronological order; by instrument.

b. Sample Data

Sample data shall be arranged in packets with the Organic Analysis Data Sheet (Form I SV, including Form I SV-TIC), followed by the raw data for semivolatile samples. These sample packets should then be placed in increasing EPA sample number order, considering both letters and numbers in ordering samples.

(1) Target Compound Results - Organic Analysis Data Sheet (Form I SV-1, SV-2).

Tabulated results (identification and quantitation) of the specified target compounds (Exhibit C). The validation and release of these results is authorized by a specific, signed statement in the SDG Marrative (reference E.1). In the event that the Laboratory Manager cannot validate all data reported for each sample, the Laboratory Manager shall provide a detailed description of the problems associated with the sample in the SDG Marrative.

(2) Tentatively Identified Compounds (Form I SV-TIC).

This form must be included even if no compounds are found. If so, indicate this on the form by entering "O" in the field for "Number found".

Form I SV-TIC is the tabulated list of the highest probable match for up to 20 of the non-surrogate organic compounds not listed in Exhibit C (TCL). It includes the CAS (Chemical Abstracts Service) Registry Number, tentative identification, and estimated concentration.

(3) Reconstructed total ion chromatograms (RIC) for each sample, sample extract, standard, blank, and spiked sample.

RICs must be normalized to the largest nonsolvent component, and must contain the following header information:

- o EPA sample number.
- o Date and time of analysis.
- o GC/MS instrument ID.
- o Lab file ID.

Internal standards and surrogates compounds are to be labeled with the names of compounds, either directly out from the peak, or on a print-out of retention times if retention times are printed over the peak.

If automated data system procedures are used for preliminary identification and/or quantification of the target compounds, the complete data system report must be included in all sample data packages, in addition to the reconstructed ion chromatogram. The complete data system report shall include all of the information listed below. For laboratories which do not use the automated data system procedures, a laboratory "raw data sheet." containing the following information, must be included in the sample data package in addition to the chromatogram.

- o EPA sample number.
- o Date and time of analysis.
- o RT or scan number of identified target compounds.
- o Ion used for quantitation with measured area.
- o Copy of area table from data system.
- o GC/MS instrument ID.
- o Lab file ID.

In all instances where the data system report has been edited, or where manual integration or quantitation has been performed, the GC/MS operator must identify such edits or manual procedures by initialing and dating the changes made to the report.

- (4) For each sample, by each compound identified, the following shall be included in the data package:
 - (a) Copies of raw spectra and copies of background-subtracted mass spectra of target compounds listed in Exhibit C (TCL) that are identified in the sample and corresponding background-subtracted target compound standard mass spectra. Spectra must be labeled with EPA sample number, lab file ID, date and time of analysis, and GC/MS instrument ID; compound names must be clearly marked on all spectra.
 - (b) Copies of mass spectra of nonsurrogate organic compounds not listed in Exhibit C (TCL), Tentatively Identified Compounds with associated best-match spectra (three best matches), labeled as in (4)(a) above.
- c. Standards Data
 - (1) Initial Calibration Data (Form VI SV-1, SV-2) in order by instrument, if more than one instrument used.
 - (a) Semivolatile standard(s) reconstructed ion chromatograms and quantitation reports (or legible facsimile) for the initial (five point) calibration, labeled as in b.(3) above. Spectra are not required.
 - (b) All initial calibration data that pertain to samples in the data package must be included, regardless of when it was performed and for which Case. When more than one initial calibration is performed, the data must be put in chronological order, by instrument.

- (2) Continuing Calibration (Form VII SV-1, SV-2) in order by instrument, if more than one instrument used.
 - (a) Semivolatile standard(s) reconstructed ion chromatograms and quantitation reports (or legible facsimile) for all continuing (12 hour) calibrations, labeled as in b.(3) above. Spectra are not required.
 - (b) When more than one continuing calibration is performed, forms must be in chronological order, by instrument.
- (3) Semivolatile GPC Calibration Data UV detector traces showing peaks that correspond to the compounds in the semivolatile GPC calibration mixture. Traces must be labeled with GPC column identifier, date of calibration, and with compound names labeled either directly out from the peak, or on a printout of retention times, if retention times are printed over the peak. Do not include Form IX Pest-2, as the compounds used on that form are not appropriate for semivolatile sample extracts.
- d. Raw QC Data
 - DFTPP (for each 12-hour period, for each GC/MS system utilized)
 - (a) Bar graph spectrum, labeled as in b(3) above.
 - (b) Mass listing, labeled as in b. (3) above.
 - (c) Reconstructed total ion chromatogram (RIC), labeled as in b.(3) above.
 - (2) Blank Data in chronological order. NOTE: This order is different from that used for samples.
 - (a) Tabulated results (Form I SV-1, SV-2)
 - (b) Tentatively Identified Compounds (Form I SV-TIC) even if none found.
 - (c) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b(3) above.
 - (d) Target compound spectra with lab generated standard, labeled as in b(4) above. Data systems which are incapable of dual display shall provide spectra in the following order:
 - o Raw target compound spectra.
 - Enhanced or background subtracted spectra.
 - o Laboratory generated standard spectra.

- (e) GC/MS library search spectra for Tentatively Identified Compounds (TIC), labeled as in b(4) above.
- (f) Quantitation/Calculation of Tentatively Identified Compound(s) (TIC) concentrations
- (3) Matrix Spike Data
 - (a) Tabulated results (Form I) of target compounds. Form 1 SV-TIC not required.
 - (b) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b(3) above. Spectra <u>not</u> required.
- (4) Matrix Spike Duplicate Data
 - (a) Tabulated results (Form I SV-1, SV-2) of target compounds. Form 1 SV-TIC not required.
 - (b) Reconstructed ion chromatogram(s) and quantitation report(s) or legible facsimile (GC/MS), labeled as in b(3) above. Spectra not required.

5. Pasticide/Aroclor Data

- a. QC Summary
 - (1) Surrogate Percent Recovery Summary (Form II PEST)
 - (2) Matrix Spike/Matrix Spike Duplicate Summary (Form III PEST)
 - (3) Method Blank Summary (Form IV PEST)

(If more than a single form is necessary, forms must be arranged in chronological order by date of analysis of the blank.)

b. Sample Data

Sample data shall be arranged in packets with the Organic Analysis Data Sheet (Form I PEST), followed by the raw data for pesticide samples. These sample packets should then be placed in increasing EPA sample number order, considering both letters and numbers in ordering samples.

(1) Target Compound Results - Organic Analysis Data Sheet (Form I PEST).

Tabulated results (identification and quantitation) of the specified target compounds (Exhibit C). The validation and release of these results is authorized by a specific, signed statement in the SDG Narrative (reference E.1). In

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the event that the Laboratory Manager cannot validate all data reported for each sample, the Laboratory Manager shall provide a detailed description of the problems associated with the sample in the SDG Narrative.

(2) Copies of pesticide chromatograms.

All chromatograms must be labeled with the following information:

- o EPA Sample Number.
- o Volume injected (uL).
- o Date and time of injection.
- o GC column identification (by stationary phase and internal diameter).
- o GC instrument identification.
- o Positively identified compounds must be labeled with the names of compounds, either directly out from the peak, or on a print-out of retention times if retention times are printed over the peak.
- (3) Copies of pesticide chromatograms from second GC column, labeled as in (2) above.
- (4) GC Integration report or data system printout.
- (5) Manual work sheets.
- (6) If pesticide/Aroclors are confirmed by GC/MS, the Contractor shall submit copies of raw spectra and copies of background-subtracted mass spectra of target compounds listed in Exhibit C (TCL) that are identified in the sample and corresponding background-subtracted TCL standard mass spectra. Compound names must be clearly marked on all spectra. For multicomponent pesticides/Aroclors confirmed by GC/MS, the Contractor shall submit mass spectra of 3 major peaks of multicomponent compounds from samples and standards.
- c. Standards Data
 - Initial Calibration of Single Component Analytes (Form VI PEST-1 and PEST-2) - all GC columns, all instruments, in chronological order by GC column and instrument.
 - (2) Initial Calibration of Multicomponent Analytes (Form VI PEST-3) - all GC columns, all instruments, in chronological order by GC column and instrument.
 - (3) Analyte Resolution Summary (Form VI PEST-4) all GC columns and instruments, in chronological order by GC column and instrument.

- (4) Calibration Verification Summary (Form VII PEST-1) for all Performance Evaluation Mixtures and Instrument blanks, on all GC columns and instruments, in chronological order by GC column and instrument.
- (5) Calibration Verification Summary (Form VII PEST-2) for all mid point concentrations of Individual Standard Mixtures A and B and Instrument blanks used for calibration verification, on all GC columns and instruments, in chronological order by GC column and instrument.
- (6) Analytical Sequence (Form VIII PEST) all GC columns and instruments, in chronological order by GC column and instrument.
- (7) Florisil Cartridge Check (Form IX PEST-1) for all lots of cartridges used to process samples in the SDG.
- (8) Pesticide GPC Calibration (Form IX PEST-2) all GPC columns, in chronological order by calibration date.
- (9) Pesticide Identification Summary for Single Component Analytes (Form X PEST-1) - for all samples with positively identified single component analytes, in order by EPA sample number.
- (10) Pesticide Identification Summary for Multicomponent Analytes (Form X PEST-2) - for all samples with positively identified multicomponent analytes, in order by EPA sample number.
- (11) Chromatograms and data system printouts are required for <u>all</u> standards including the following:
 - o Resolution Check Mixture.
 - o Performance Evaluation Mixtures, all.
 - o Individual Standard Mixture A, at three concentrations, each initial calibration.
 - o Individual Standard Mixture B, at three concentrations, each initial calibration.
 - o All multicomponent analytes (Toxaphene and Aroclors), each initial calibration.
 - o All mid point concentrations of Individual Standard Mixtures A and B used for calibration verification.
 - o Florisil cartridge check solution, all lots.
 - o Pesticide GPC Calibration Check Solution, all calibrations relating to samples in the SDG.
 - o All multicomponent analyte standards analyzed for confirmation.

- (12) A printout of retention times and corresponding peak areas or peak heights must accompany each chromatogram. In addition, all chromatograms are required to be labeled with the following:
 - o EPA Sample Number for the standard, i.e., INDA1, INDA2, etc. (See Forms Instructions for details.)
 - Label all standard peaks for all individual compounds either directly out from the peak or on the printout of retention times if retention times are printed over the peak.
 - o Total nanograms injected for each standard.
 - o Date and time of injection.
 - o GC column identification (by stationary phase and internal diameter).
 - o GC instrument identification.
- (13) A copy of the computer reproduction output covering the entire calibration range is required for each initial calibration.
- (14) Pesticide GPC Calibration Data UV detector traces showing peaks that correspond to the compounds in the pesticide GPC calibration mixture. Traces must be labeled with GPC column identifier, date of calibration, and with compound names labeled either directly out from the peak, or on a printout of retention times, if retention times are printed over the peak.
- d. Raw QC Data
 - Blank Data in chronological order, by type of blank (method, instrument, sulfur clean up). NOTE: This order is different from that used for samples.
 - (a) Tabulated results (Form I PEST).
 - (b) Chromatogram(s) and data system printout(s) (GC) for each GC column and instrument used for analysis, labeled as in b.(2) above.
 - (2) Matrix Spike Data
 - (a) Tabulated results (Form I PEST) of target compounds.
 - (b) Chromatogram(s) and data system printout(s) (GC), labeled as in b.(2) above.

- (3) Matrix Spike Duplicate Data
 - (a) Tabulated results (Form I PEST) of target compounds.
 - (b) Chromatogram(s) and data system printout(s) (GC), labeled as in b.(2) above.

E. <u>Complete SDG File</u>

As specified in the Delivery Schedule, one Complete SDG File (CSF) including the original Sample Data Package shall be delivered to the Region concurrently with delivery of the Sample Data Package to SMO and EMSL-LV. The contents of the CSF will be numbered according to the specifications described in Section III and IV of Exhibit B. The Document Inventory Sheet, Form DC-2, is contained in Section IV. The CSF will contain all original documents where possible. No copies will be placed in the CSF unless the originals are bound in a logbook which is maintained by the laboratory. The CSF will contain all original documents specified in Section III and IV, and Form DC-2 of Exhibit B of the SOW.

The CSF will consist of the following original documents in addition to the documents in the Sample Data Package:

- 1. Original Sample Data Package
- 2. A completed and signed Document Inventory Sheet (Form DC-2).
- 3. All original shipping documents, including, but not limited to, the following documents:
 - a. EPA Chain of Custody Record.
 - b. Airbills.
 - c. EPA Traffic Reports.
 - d. Sample Tags (if present) sealed in plastic bags.
- 4. All original receiving documents, including, but not limited to, the following documents:
 - a. Form DC-1.
 - b. Other receiving forms or copies of receiving logbooks.
 - c. SDG Cover Sheet.
- 5. All original laboratory records, not already submitted in the Sample Data Package, of sample transfer, preparation and analysis, including, but not limited to, the following documents:
 - a. Original preparation and analysis forms or copies of preparation and analysis logbook pages.
 - b. Internal sample and sample extract transfer chaip-of-custody records.

- c. Screening records.
- d. All instrument output, including strip charts from screening activities.
- 6. All other original SDG-specific documents in the possession of the laboratory, including, but not limited to, the following documents:
 - a. Telephone contact logs.
 - b. Copies of personal logbook pages.
 - c. All hand written case-specific notes.
 - d. Any other case-specific documents not covered by the above.

NOTE: All Case-related documentation may be used or admitted as evidence in subsequent legal proceedings. Any other Case-specific documents generated after the CSF is sent to EPA, as well as copies that are altered in any fashion, are also deliverables to EPA. (Original to the Region, and copies to SMO and EMSL-LV).

If the laboratory does submit SDG-specific documents to EPA after submission of the CSF, the documents should be numbered as an addendum to the CSF and a revised DC-2 form should be submitted, or the documents should be numbered as a new CSF and a new DC-2 form should be submitted to the Regions only.

F. Data in Computer-Readable Form

The Contractor shall provide a computer-readable copy of the data on data reporting Forms I-X for all samples in the Sample Delivery Group, as specified in the Contract Performance/Delivery Schedule. Computer-readable data deliverables shall be submitted on IBM or IBM-compatible, 5.25 inch floppy double-sided, double density 360 K-byte, a high density 1.2 M-byte or a 3.5 inch double-sided double density 720 K-byte or 1.44 M-byte diskette.

When submitted, diskettes shall be packaged and shipped in such a manner that the diskette(s) cannot be bent or folded, and will not be exposed to extreme heat or cold or any type of electromagnetic radiation. The diskette(s) must be included in the same shipment as the hardcopy data and shall, at a minimum, be enclosed in a diskette mailer. The data shall be recorded in ASCII, text file format, and shall adhere to the file, record and field specifications listed in Exhibit H, Data Dictionary and Format for Data Deliverables in Computer-Readable Format.

If the Contractor wishes to use a reporting format other than the one specified, equivalence must be demonstrated and approved by the Administrative Project Officer prior to the award of the contract.

G. GC/MS Tapes

The Contractor must store <u>all</u> raw and processed GC/MS data on magnetic tape, in appropriate instrument manufacturer's format. This tape must include data for samples, blanks, matrix spikes, matrix spike duplicates, initial calibrations, continuing calibrations, BFB and DFTPP, as well as all laboratory-generated spectral libraries and quantitation reports required to generate the data package. The Contractor shall maintain a written reference logbook of tape files to EPA sample number, calibration data, standards, blanks, matrix spikes, and matrix spike duplicates. The logbook should include EPA sample numbers and standard and blank ID's, identified by Case and Sample Delivery Group.

The Contractor is required to retain the GC/MS tapes for 365 days after data submission. During that time, the Contractor shall submit tapes and associated logbook pages within seven days after receipt of a written request from the Administrative Project Officer.

When submitting GC/MS tapes to the Agency, the following materials must be delivered in response to the request:

- 1. All associated raw data files for samples, blanks, matrix spikes, matrix spike duplicates, initial and continuing calibration standards, and instrument performance check solutions (BFB and DFTPP).
- 2. All processed data files and quantitation output files associated with the raw data files described above.
- 3. All associated identifications and calculation files used to generate the data submitted in the data package.
- 4. All laboratory-generated mass spectral library files (NIST/EPA/MSDC library not required).
- 5. A copy of the Contractor's written reference logbook relating tape files to EPA Sample Number, calibration data, standards, blanks, matrix spikes, and matrix spike duplicates. The logbook must include EPA Sample Numbers and Lab File identifiers for all samples, blanks, and standards, identified by Case and SDG.

The laboratory must also provide a statement attesting to the completeness of the GC/MS data tape submission, signed and dated by the Laboratory Manager. This statement must be part of a cover sheet that includes the following information relevant to the data tape submission:

- 1. Laboratory name
- 2. Date of submission
- 3. Case Number
- 4. SDG Number
- 5. GC/MS make and model number
- 6. Software version
- 7. Disk drive type (e.g. CDC, PRIAM, etc.)

- 8. File transfer method (e.g. DSD, DTD, FTP, Aquarius, etc.)
- 9. Names and telephone numbers of two laboratory contacts for further information regarding the submission.

H. Extracts

The Contractor shall preserve sample extracts at $4^{\circ}C$ ($\pm 2^{\circ}C$) in bottles/vials with Teflon-lined septa. Extract bottles/vials shall be labeled with EPA sample number, Case number and Sample Delivery Group (SDG) number. A logbook of stored extracts shall be maintained, listing EPA sample numbers and associated Case and SDG numbers.

The Contractor is required to retain extracts for 365 days following data submission. During that time, the Contractor shall submit extracts and associated logbook pages within seven days following receipt of a written request from the Administrative Project Officer or the Sample Management Office. SECTION III

FORMS INSTRUCTIONS

SECTION III

FORM INSTRUCTION GUIDE

This section includes specific instructions for the completion of all required forms. Each of the forms is specific to a given fraction (volatile, semivolatile, pesticide/Aroclor), and in some instances specific to a given matrix (water or soil) within each fraction. The contractor shall submit only those forms pertaining to the fractions analyzed for a given sample or samples. For instance, if a sample is scheduled for volatile analysis only, provide only VOA forms. There are two pages relating to the semivolatile fraction for Forms I, VI, VII, and VIII. Whenever semivolatiles are analyzed and one of the above-named forms is required, both pages (SV-1 and SV-2) must be submitted. These instructions are arranged in the following order:

- A. General Information and Header Information
- B. Organic Analysis Data Sheets (Form I, All Fractions)
- C. System Monitoring Compound Recovery (Form II VOA)
- D. Surrogate Recovery (Form II, SV and PEST)
- E. Matrix Spike/Matrix Spike Duplicate Recovery (Form III, All Fractions)
- F. Method Blank Summary (Form IV, All Fractions)
- G. GC/MS Instrument Performance Check and Mass Calibration (Form V VOA and SV)
- H. GC/MS Initial Calibration Data (Form VI VOA, SV-1, SV-2)
- I. GC Initial Calibration Data (Form VI PEST-1, PEST-2, PEST-3, PEST-4)
- J. GC/MS Continuing Calibration Data (Form VII VOA, SV-1, SV-2)
- K. GC/EC Continuing Calibration (Form VII PEST)
- L. Internal Standard Area Summary (Form VIII VOA and SV)
- M. Pesticide Analytical Sequence (Form VIII PEST)
- N. Pesticide Cleanup Procedures (Form IX PEST-1, PEST-2)
- 0. Pesticide/Aroclor Identification (Form X PEST-1, PEST-2)
- P. Sample Log-In Sheet (Form DC-1)
- Q. Document Inventory Sheet (Form DC-2)

A. <u>General Information and Header Information</u>

The data reporting forms presented in Section IV have been designed in conjunction with the computer-readable data format specified in Exhibit H, Data Dictionary and Format for Data Deliverables in Computer-Readable Format. The specific length of each variable for computer-readable data transmission purposes is given in the Data Dictionary (Exhibit H). Information entered on these forms must not exceed the size of the field given on the form, including such laboratory-generated items as Lab Name and Lab Sample ID.

Note that on the hardcopy forms (Section IV), the space provided for entries is greater in some instances than the length prescribed for the variable as written to diskette (see Exhibit H). Greater space is provided on the hardcopy forms for the sake of visual clarity.

Values must be reported on the hardcopy forms according to the individual form instructions in this Section. For example, results for concentrations of VOA target compounds must be reported to two significant figures if the value is greater than or equal to 10. Values can be written to the diskette file in any format that does not exceed the field specification as given in the record specifications and discussed in "Record Structure", paragraph 5 of Exhibit H.

All characters which appear on the data reporting forms presented in the contract (Exhibit B, Section IV) <u>must</u> be reproduced by the Contractor when submitting data, and the format of the forms submitted <u>must be identical</u> to that shown in the contract. No information may be added, deleted, or moved from its specified position without <u>prior</u> <u>written</u> approval of the EPA Administrative Project Officer. The names of the various fields and compounds (i.e., "Lab Code," "Chloromethane") <u>must</u> appear as they do on the forms in the contract, including the options specified in the form (i.e., "Matrix: (soil/water)" must appear, not just "Matrix"). For items appearing on the uncompleted forms (Section IV), the use of uppercase <u>and</u> lowercase letters is optional.

Alphabetic <u>entries</u> made onto the forms by the Contractor shall be in <u>ALL UPPERCASE</u> letters (i.e., "LOW", not "Low" or "low"). If an entry does not fill the entire blank space provided on the form, null characters shall be used to remove the remaining underscores that comprise the blank line. See Exhibit H for more detailed instructions. However, <u>do not</u> remove the underscores or vertical bar characters that delineate "boxes" on the forms. The only exception would be those underscores at the bottom of a "box" that are intended as a data entry line (for instance, on Form 2A, line 30, if data must be entered on line 30, it will replace the underscores).

Six pieces of information are common to the header sections of each data reporting form. They are Lab Name, Contract, Lab Code, Case No., SAS No., and SDG No. Except as noted below for SAS No., this information <u>must</u> be entered on every form and <u>must</u> match on every form.

The "Lab Name" shall be the name chosen by the Contractor to identify the laboratory. It may not exceed 25 characters.

The "Lab Code" is an alphabetical abbreviation of up to 6 letters, assigned by the EPA, to identify the laboratory and aid in data processing. This lab code shall be assigned by the EPA at the time a contract is awarded, and <u>shall not</u> be modified by the Contractor, except at the direction of the EPA. If a change of name or ownership occurs at the laboratory, the lab code will remain the same until the contractor is directed by the EPA to use another lab code assigned by the EPA.

The "Case No." is the EPA-assigned Case number associated with the sample, and reported on the Traffic Report.

The "Contract" is the number of the EPA contract under which the analyses were performed. In the case of multiple laboratories operating under a corporate-wide contract, the contract number entered shall be that of the corporate contract, regardless of the facility performing the analyses (see Lab Code, above).

The "SDG No." is the Sample Delivery Group number. The Sample Delivery Group (SDG) number is the EPA Sample Number of the first sample received in the SDG. When several samples are received together in the first SDG shipment, the SDG number shall be the lowest sample number (considering both alpha and numeric designations) in the first group of samples received under the SDG.

The "SAS No." is the EPA-assigned number for analyses performed under Special Analytical Services. If samples are to be analyzed under SAS only and reported on these forms, then enter SAS No. and leave Case No. blank. If samples are analyzed according to the "Routine Analytical Services" (IFB) protocols and have additional "SAS" requirements, list both Case No. and SAS No. on all forms. If the analyses have no SAS requirements, leave "SAS No." blank. NOTE: Some samples in an SDG may have a SAS No. while others do not.

The other information common to most of the forms is the "EPA Sample No.". This number appears either in the upper righthand corner of the form, or as the left column of a table summarizing data from a number of samples. When the "EPA Sample No." is entered into the triple-spaced box in the upper righthand corner of Form I, Form IV, or Form X, it should be entered on the middle line of the three lines that comprise the box.

<u>All</u> samples, matrix spikes, matrix spike duplicates, blanks, and standards shall be identified with an EPA Sample Number. For field samples, matrix spikes and matrix spike duplicates, the EPA Sample Number is the unique identifying number given in the Traffic Report that accompanied that sample. In order to facilitate data assessment, the following sample suffixes must be used:

XXXXX	-	EPA sample number
XXXXXMS		Matrix spike sample
XXXXXMSD	-	Matrix spike duplicate sample
XXXXXRE	-	Re-extracted and re-analyzed sample
XXXXXDL	-	Sample analyzed at a secondary dilution

Form VIII Pest requires that <u>all</u> samples analyzed in a given analytical sequence be listed, regardless of whether or not they are part of the SDG being reported. Therefore, use "ZZZZZ" as the EPA Sample No. for any sample analysis <u>not</u> associated with the SDG being reported.

For blanks and standards, the following identification scheme must be used as the "EPA Sample No."

- 1. Volatile blanks shall be identified as VBLK##.
- 2. Semivolatile blanks shall be identified as SBLK##.
- 3. Pesticide/Aroclor method blanks shall be identified as PBLK##.
- 4. Pesticide/Aroclor instrument blanks shall be identified as PIBLK##.

The "EPA Sample No." <u>must be unique</u> for each blank within an SDG. Within a fraction, a laboratory must achieve this by replacing the twocharacter "##" terminator of the identifier with one or two characters or numbers, or a combination of both. For example, possible identifiers for volatile blanks would be VBLK1, VBLK2, VBLKA1, VBLKB2, VBLK10, VBLKAB, etc.

Volatile and semivolatile standards shall be identified as FSTD###, where

F - Fraction (V for volatiles; S for semivolatiles).
STD - Indicates a standard.
- The concentration in ug/L of volatile standards
(i.e., 010, 020, 050, 100, and 200) or the amount
injected in ng for semivolatile standards
(i.e., 020, 050, 080, 120, and 160).

As for the blank identifiers, these designations will have to be concatenated with other information to uniquely identify each standard. For pesticide/Aroclor standards, the following scheme shall be used to enter "EPA Sample Number".

Name

EPA Sample Number

Individual Mix A (low point)	INDAL##
Individual Mix A (mid point)	INDAM##
Individual Mix A (high point)	INDAH##
Individual Mix B (low point)	INDBL##
Individual Mix B (mid point)	INDBM##
Individual Mix B (high point)	INDBH##
Resolution Check	RESC##
Performance Evaluation Mixture	PEM##
Toxaphene	TOXAPH##
Aroclor 1016	AR1016##
Aroclor 1221	AR1221##
Aroclor 1232	AR1232##
Aroclor 1242	AR1242##
Aroclor 1248	AR1248##
Aroclor 1254	AR1254##
Aroclor 1260	AR1260##

The permitted mixture of Aroclor 1016 and Aroclor 1260 shall be entered as AR1660##.

If the standards are injected onto both GC columns on the same instrument simultaneously, the same EPA Sample Number may be used for reporting data for the standards for both columns. If simultaneous injections are not made, then the same number may not be used.

Several other pieces of information are common to many of the Data Reporting Forms. These include Matrix, Sample wt/vol, Level, Lab Sample ID, and Lab File ID.

For "Matrix", enter "SOIL" for soil/sediment samples, and enter "WATER" for water samples. NOTE: The matrix <u>must</u> be spelled out. Abbreviations such as "S" or "W" shall not be used.

For "Sample wt/vol" enter the number of grams (for soil) or milliliters (for water) of sample used in the first blank line, and the units, either "G" or "ML", in the second blank.

For volatiles and semivolatiles, for "Level", enter the determination of concentration level made from the mandatory screening of soils. Enter as "LOW" or "MED", not "L" or "M". All water samples are "LOW" level and shall be entered as such. Note: There is no differentiation between low and medium soil samples for the Pesticide/Aroclor forms, and no level is entered on any of the these forms.

"Lab Sample ID" is an optional laboratory-generated internal identifier. Up to 12 alpha-numeric characters may be reported here. If the contractor does not have a Lab Sample ID, this field may be left blank. "Lab File ID" is the laboratory-generated name of the GC/MS data system file containing information pertaining to a particular analysis. Up to 14 alpha-numeric characters may be used here.

"Instrument ID" is common to many of the forms, particularly those containing calibration data. The identifier used by the laboratory must include some indication of the manufacturer and/or model of the instrument, and contain additional characters that differentiate between all instrument of the same type in the laboratory.

"GC Column" and "ID (mm)" are common to various other forms. These two fields are to be used to identify the stationary phase of the GC column (previously called GC Column ID), and the internal diameter of the GC column in millimeters (mm). For packed columns, convert the ID from inches to millimeters as necessary, and enter in the "ID" field.

Forms II, IV, V, VIII, IX, and X contain a field labeled "page _ of _" in the bottom lefthand corner. If the number of entries required on any of these forms exceeds the available space, continue entries on another copy of the same fraction-specific form, duplicating all header information. If a second page is required, number the pages consecutively, as "page 1 of 2" and "page 2 of 2". If a second page is not required, number the page "page 1 of 1." NOTE: These forms are fraction-specific, and often matrix-specific within fraction. For example, Form II VOA-1 and Form II VOA-2 are for different data. Therefore, <u>do not</u> number the pages of all six versions of Form II as "1 of 6, 2 of 6, etc." Number only pages within a fraction-specific and matrix-specific form.

For rounding off numbers to the appropriate level of precision, observe the following common rules. If the figure following those to be retained is less than 5, drop it (round down). If the figure is greater than 5, drop it and increase the last digit to be retained by 1 (round up). If the figure following the last digit to be retained equals 5, round up if the digit to be retained is odd, and round down if that digit is even.

B. Organic Analysis Data Sheet (Form I. All Fractions)

1. Form I VOA, Form I SV-1, Form I SV-2, Form I PEST

This form is used for tabulating and reporting sample analysis results for target compounds. If all fractions are not requested to be analyzed, only the pages specifically required must be submitted. If VOA analysis only is requested, Form I VOA and Form I VOA TIC must be submitted. If the pesticide/Aroclor analysis is the only analysis requested, only Form I Pest must be submitted for that sample.

Complete the header information on each page of Form I required, according to the instructions in Part A and as follows:

For volatiles, for "t moisture not dec.", enter the nondecanted percent moisture. This is the only percent moisture determination made for volatiles, as the entire contents of the VOA vial are considered as the sample.

For semivolatiles and pesticides/Aroclors, enter the values for the percent moisture determined during the analysis. In the field "decanted (Y/N)", enter "Y" if the sample had standing water above the soil/sediment that was decanted, or "N" if no water was decanted off the surface of the sample. Report percent moisture (decanted or not decanted) to the nearest whole percentage point (i.e., 5%, not 5.3%). Leave these fields blank for Form I for method blanks and instrument blanks.

For volatiles, enter the GC column identifier under "GC Column", and the internal diameters in millimeters (mm), to two decimal places, under "ID". For packed columns, convert the ID from inches to millimeters as necessary, and enter in this field.

For pesticides/Aroclors, enter the method of extraction as "SEPF" for separatory funnel, "CONT" for continuous liquid-liquid extraction, or "SONC" for sonication (soils only).

If gel permeation chromatography, "GPC Cleanup", was performed, enter "Y" for yes. Otherwise, enter "N" for no, if GPC was not performed. NOTE: GPC is <u>required</u> for all <u>soil</u> samples analyzed for semivolatiles and pesticides/Aroclors, therefore all soil sample forms will contain "Y" in this field.

For soil samples only, enter pH for semivolatiles and pesticides/Aroclors, reported to 0.1 pH units.

"Date Received" is the date of sample receipt at the laboratory, as noted on the Traffic Report (i.e., the VTSR). It should be entered as MM/DD/YY.

"Date Extracted" and "Date Analyzed" should be entered in a similar fashion. If continuous liquid-liquid extraction procedures are used, enter the date on which the procedure was <u>started</u> for "Date Extracted". If separatory funnel or sonication procedures are used, enter the date on which the procedure was <u>completed</u>. For pesticide/Aroclor samples, the date of analysis should be the date of the first GC analysis performed. The date of sample receipt will be compared with the extraction and analysis dates of each fraction to ensure that contract holding times were not exceeded.

For volatiles, if a medium soil sample is analyzed, under the "Soil Extract Volume" enter total volume of the methanol extract in <u>microliters</u>. This volume includes any methanol not collected from the filtration of the extract through glass wool, and is typically 10000 uL, i.e., the 10 mL of methanol use for the extraction. If a medium soil sample is analyzed, enter also the volume of the methanol extract added to the reagent water in the purge tube and analyzed, under "Soil Aliquot Volume". Enter this volume in microliters (uL). For semivolatiles and pesticides/Aroclors, enter the actual volume of the most concentrated sample extract, in microliters, under "Concentrated Extract Volume". This volume typically will be 1000 uL, or 500 uL when GPC was performed. If a dilution of the sample extract is made in a subsequent analysis, this volume will remain the same, but the dilution factor will change.

For semivolatiles and pesticides/Aroclors, enter the volume of the sample extract injected into the GC under "Injection Volume". Report this volume in microliters to one decimal place, i.e., 1.0 uL. Note: A 2.0 microliter injection is <u>required</u> for semivolatile analyses.

If a sample or sample extract has been diluted for analysis, enter the "Dilution Factor" as a single number, not a fraction, such as "100.0," for a 1 to 100 dilution of the sample. Enter 0.1 for a concentration of 10 to 1. If a sample was not diluted, enter "1.0." Report dilution factors to one decimal place.

For positively identified target compounds, the Contractor shall report the concentrations detected as <u>uncorrected</u> for blank contaminants.

For volatile and semivolatile results, report analytical results to one significant figure if the value is less than 10, and two significant figures if the value is above 10.

Report all pesticide/Aroclor results to two significant figures.

The appropriate concentration units, ug/L or ug/kg, must be entered.

If the result is a value greater than or equal to the quantitation limit, report the value.

Under the column labeled "Q" for qualifier, flag each result with the specific Data Reporting Qualifiers listed below. The Contractor is encouraged to use additional flags or footnotes. The definition of such flags must be explicit and must be included in the SDG Narrative.

For reporting results to the USEPA, the following contract specific qualifiers are to be used. The seven qualifiers defined below are not subject to modification by the laboratory. Up to five qualifiers may be reported on Form I for each compound.

The seven EPA-defined qualifiers to be used are as follows:

U - Indicates compound was analyzed for but not detected. The sample quantitation limit must be corrected for dilution and for percent moisture. For example, 10 U for phenol in water if the sample final volume is the protocol-specified final volume. If a 1 to 10 dilution of extract is necessary, the reported limit is 100 U. For a soil sample, the value must also be

adjusted for percent moisture. For example, if the sample had 24% moisture and a 1 to 10 dilution factor, the sample quantitation limit for phenol (330 U) would be corrected to

(<u>330 U</u>) x df where D = <u>100 - % moisture</u> D 100

and df = dilution factor

For example, at 24% moisture, $D = \frac{100-24}{100} = 0.76$

(330 U) x 10 - 4300 U rounded to the appropriate number of significant figures .76

For soil samples subjected to GPC clean-up procedures, the extract must be concentrated to 0.5 mL, and the sensitivity of the analysis is not compromised by the cleanup procedures. Therefore, the CRQL values in Exhibit C will apply to all samples, regardless of cleanup. However, if a sample extract cannot be concentrated to the protocol-specified volume (see Exhibit C), this fact must be accounted for in reporting the sample quantitation limit.

- J Indicates an estimated value. This flag is used either when estimating a concentration for tentatively identified compounds where a 1:1 response is assumed, or when the mass spectral data indicate the presence of a compound that meets the identification criteria but the result is less than the sample quantitation limit but greater than zero. For example, if the sample quantitation limit is 10 ug/L, but a concentration of 3 ug/L is calculated, report it as 3J. The sample quantitation limit must be adjusted for dilution as discussed for the U flag.
- N Indicates presumptive evidence of a compound. This flag is only used for tentatively identified compounds, where the identification is based on a mass spectral library search. It is applied to all TIC results.
- P This flag is used for a pesticide/Aroclor target analyte when there is greater than 25% difference for detected concentrations between the two GC columns (see Form X) The lower of the two values is reported on Form I and flagged with an "P".
- C This flag applies to pesticide results where the <u>identification</u> has been confirmed by GC/MS. If GC/MS confirmation was attempted but was unsuccessful, do <u>not</u> apply this flag, instead use a laboratory-defined flag, discussed below.

- B This flag is used when the analyte is found in the associated blank as well as in the sample. It indicates possible/probable blank contamination and warns the data user to take appropriate action. This flag must be used for a TIC as well as for a positively identified target compound.
- E This flag identifies compounds whose concentrations exceed the calibration range of the GC/MS instrument for that specific analysis. If one or more compounds have a response greater than full scale, except as noted in Exhibit D, the sample or extract must be diluted and re-analyzed according to the specifications in Exhibit D. All such compounds with a response greater than full scale should have the concentration flagged with an "E" on the Form I for the original analysis. If the dilution of the extract causes any compounds identified in the first analysis to be below the calibration range in the second analysis, then the results of both analyses shall be reported on separate copies of Form I. The Form I for the diluted sample shall have the "DL" suffix appended to the sample number. NOTE: For total xylenes, where three isomers are quantified as two peaks, the calibration range of each peak should be considered separately, e.g., a diluted analysis is not required for total xylenes unless the concentration of either peak separately exceeds 200 ug/L.
- D This flag identifies all compounds identified in an analysis at a secondary dilution factor. If a sample or extract is re-analyzed at a higher dilution factor, as in the "E" flag above, the "DL" suffix is appended to the sample number on the Form I for the diluted sample, and <u>all</u> concentration values reported on that Form I are flagged with the "D" flag. This flag alerts data users that any discrepancies between the concentrations reported may be due to dilution of the sample or extract.
- A This flag indicates that a TIC is a suspected aldol-condensation product.
- X Other specific flags may be required to properly define the results. If used, they must be fully described, and such description attached to the Sample Data Summary Package and the SDG Narrative. Begin by using "X". If more than one flag is required, use "Y" and "Z" as needed. If more than five qualifiers are required for a sample result, use the "X" flag to combine several flags, as needed. For instance, the "X" flag might combine the "A", "B", and "D" flags for some sample. The laboratory-defined flags are <u>limited to</u> the letters "X", "Y", and "Z".

The combination of flags "BU" or "UB" is expressly prohibited. Blank contaminants are flagged "B" only when they are detected in the sample.

2. Form I VOA-TIC and Form I SV-TIC

Fill in all header information as above.

Report Tentatively Identified Compounds (TICs) including CAS number, compound name, retention time, and the estimated concentration (criteria for reporting TICs are given in Exhibit D, Section IV). Retention time must be reported in minutes and decimal minutes, not seconds or minutes:seconds.

If, in the opinion of the mass spectral interpretation specialist, no valid tentative identification can be made, the compound shall be reported as unknown.

Include a Form I VOA-TIC or SV-TIC for every volatile and semivolatile fraction of every sample and method blank analyzed, <u>even</u> if no TICs are found. Total the number of TICs found, <u>including</u> aldol-condensation products (but see below), and enter this number in the "Number TICs found." If none were found, enter "0" (zero). Form I VOA-TIC or SV-TIC must be provided for <u>every analysis</u>, including required dilutions and reanalyses, even if no TICs are found.

If the name of a compound exceeds the 28 spaces in the TIC column, truncate the name to 28 characters. If the compound is an unknown, restrict description to no more than 28 characters (i.e., unknown hydrocarbon, etc.).

Peaks that are suspected as aldol-condensation reaction products (i.e., 4-methyl-4-hydroxy-2-pentanone and 4-methyl-3-pentene-2-one) shall be summarized on this form, flagged "A", and included in the total "Number TICs found," but not counted as part of the 20 most intense non-target semivolatile compounds to be searched.

C. System Monitoring Compound Recovery (Form II VOA)

For volatiles, Form II is used to report the recoveries of the system monitoring compounds added to each volatile sample, blank, matrix spike, and matrix spike duplicate prior to analysis. The system monitoring compounds, previously less than accurately termed volatile surrogates, are used to monitor the performance of the purge and trapgas chromatograph-mass spectrometer system as a whole. Form II VOA is matrix-specific, so that system monitoring compound recoveries for water samples are reported on a different version of Form II than the recoveries in soil samples. Soil sample recoveries are further differentiated by concentration level.

Complete the header information and enter EPA Sample Numbers as described in part A. For volatile soil samples <u>only</u>, specify the "level" as "LOW" or "MED", as on Form I. <u>Do not</u> mix low and medium level samples on one form. Complete one for each level. For each system monitoring compound, report the percent recovery to the nearest whole percentage point, and to the number of significant figures given by the QC limits at the bottom of the form. Flag each system monitoring compound recovery outside the QC limits with an asterisk (*). The asterisk must be placed in the last space in each appropriate column, under the "#" symbol. In the far righthand column, total the number of system monitoring compound(s) recoveries outside the QC limits for each sample. If no system monitoring compound(s) were outside the limits, enter "0".

If the system monitoring compound(s) are diluted out in any analysis, enter the calculated recovery, or "0" (zero) if the system monitoring compound(s) is not detected, and flag the system monitoring compound(s) recoveries with a "D" in the column under the "#" symbol. Do not include results flagged "D" in the total number of recoveries for each sample outside the QC limits.

Number all pages as described in part A.

D. Surrogate Recovery (Form II, SV and PEST)

Form II is used to report the recoveries of the surrogate compounds added to each semivolatile and pesticide/Aroclor sample, blank, matrix spike, and matrix spike duplicate. For semivolatiles, Form II is matrix-specific as well as fraction-specific, so surrogate recoveries for semivolatile water samples are reported on a different version of Form II than semivolatile soil sample surrogate recoveries.

Complete the header information and enter EPA Sample Numbers as described in part A. For semivolatile soil samples only, specify the "level" as "LOW" or "MED", as on Form I. <u>Do not</u> mix low and medium level samples on one form. Complete one for each level. For each surrogate, report the percent recovery to the nearest whole percentage point, and to the number of significant figures given by the QC limits at the bottom of the form.

Flag each surrogate recovery outside the QC limits with an asterisk (*). The asterisk must be placed in the last space in each appropriate column, under the "#" symbol. In the far righthand column, total the number of surrogate recoveries outside the QC limits for each sample. If no surrogates were outside the limits, enter "0".

If the surrogates are diluted out in any analysis, enter the calculated recovery, or "O" (zero) if the surrogate is not detected, and flag the surrogate recoveries with a "D" in the column under the "#" symbol. Do not include results flagged "D" in the total number of recoveries for each sample outside the QC limits.

The pesticide surrogate recoveries must be reported from <u>both</u> GC columns used for the analyses. Therefore, identify each GC column in the header, entering the stationary phase under "GC Column" (previously called GC Column ID), and the internal diameter (ID) of the column in millimeters under "ID". The assignment of columns as "1" and "2" is left to the discretion of the laboratory <u>if</u> the analyses are performed by simultaneous injection into a GC containing two columns. If so analyzed, the assignment of "GC Column 1" and "GC Column 2" must be consistent across all the reporting forms. If the analysis is not performed by simultaneous injection, then the assignment of GC Column number should be based on the chronological order of the two analyses.

The pesticide surrogate recovery limits are only advisory, but the Contractor must flag those recoveries outside the advisory QC limits or diluted out, nonetheless.

Number all pages as described in part A.

E. Matrix Spike/Matrix Spike Duplicate Recovery (Form III, All Fractions)

This form is used to report the results of the analyses of a matrix spike and matrix spike duplicate. The form is matrix-specific for volatiles and semivolatiles.

Complete the header information as instructed in Part A, including the EPA Sample Number for the matrix spike, without the suffixes MS or MSD.

For volatile and semivolatile soil samples, specify "level" as "LOW" or "MED", as on Form I. SDGs containing soil samples at both levels require MS/MSD at each level, therefore, for soils, prepare one form for each level.

All water samples are "LOW". Therefore, there is no MS/MSD for "medium level waters", and none shall be reported.

In the upper box in Form III, under "SPIKE ADDED", enter the calculated concentration in ug/L or ug/Kg (according to the matrix) that results from adding each spiked compound to the aliquot chosen for the matrix spike (MS). For instance, for base/neutral compounds in medium level soils, if 50 ug of spike are added to 1 g of soil, the resulting concentration is 50,000 ug/Kg. Enter the "SAMPLE CONCENTRATION", in 50,000 similar units, of each spike compound detected in the original sample. If a spike compound was not detected during the analysis of the original sample, enter the sample result as "0" (zero). Under "MS CONCENTRATION", enter the actual concentration of each spike compound detected in the matrix spike aliquot. Calculate the percent recovery of each spike compound in the matrix spike aliquot to the nearest whole percent, according to Exhibit E, and enter under "MS & REC". Flag all percent recoveries outside the QC limits with an asterisk (*). The asterisk must be placed in the last space of the percent recovery column, under the "#" symbol.

For pesticide/Aroclor matrix spikes and matrix spike duplicates, the concentration used for "MS CONCENTRATION" AND "MSD CONCENTRATION" must be the concentration of the spiked analyte reported on Form I that those analyses. Of the two concentrations calculated for each pesticide/Aroclor target compound, one on each GC column, the <u>lower</u> concentration is reported on Form I, and <u>both</u> concentrations are reported on Form X. The <u>lower</u> concentration is reported on Form III and used in the calculation of spike recovery, even if that concentration yields a recovery value that is outside the advisory QC limits.

Complete the lower box on Form III in a similar fashion, using the results of the analysis of the matrix spike duplicate (MSD) aliquot. Calculate the relative percent difference (RPD) between the matrix spike recovery and the matrix spike duplicate recovery, and enter this value in the lower box under "% RPD". Report the relative percent difference to the nearest whole percent. Compare the RPDs to the QC limits given on the form, and flag each RPD outside the QC limits with an asterisk (*) in the last space of the "% RPD" column, under the "#" symbol.

Summarize the values outside the QC limits at the bottom of the page. No further action is required by the laboratory. Performance-based QC limits will be generated and updated from recovery and RPD data.

F. Method Blank Summary (Form IV, All Fractions)

This form summarizes the samples associated with each method blank analysis. A copy of the appropriate Form IV is required for each blank.

Complete the header information on Form IV as described in Part A. The "EPA Sample No." entered in the box at the top of Form IV shall be the same number entered on the Form I for the blank itself.

For volatile blanks, enter the "Instrument ID", "Date Analyzed" "Time Analyzed", "GC Column", "ID", and "Heated Purge (Y/N)". Volatile samples analyzed by the same purging technique, i.e., ambient purge, or heated purge, may be reported together on the same Form IV, if the same method blank applies to those samples. Thus, water samples and medium soil sample may be combined on a single form, if run with a single blank.

For semivolatile blanks, enter the "Instrument ID", "Date Analyzed", "Matrix" and "Level". All water blanks are "LOW". The "Time Analyzed" shall be in military time.

For pesticide/Aroclor blanks, enter the method of extraction as "SEPF" for separatory funnel, "SONC" for sonication, or "CONT" for continuous liquid-liquid extraction.

For pesticide/Aroclor blanks, there is no differentiation between medium and low level soil samples, so no "Level" is entered on this form.

For semivolatile and pesticide/Aroclor method blanks, enter the date of extraction of the blank.

If the samples associated with pesticide/Aroclor blank are subjected to sulfur cleanup, then the blank must also be subjected to sulfur cleanup. If sulfur cleanup is employed, enter "Y" in the "Sulfur Cleanup" field, else, enter "N". If only some of the samples associated with the method blank are subjected to sulfur cleanup, a <u>Separate</u> sulfur cleanup blank is required (see Exhibit D PEST). If a <u>separate</u> sulfur cleanup blank is prepared, complete one version of Form IV associating all the samples with the method blank, and a <u>second</u> version of Form IV listing only those samples associated with the separate sulfur cleanup blank. Note: Subjecting all samples associated with a method blank to sulfur cleanup avoids the need for two forms.

Pesticide/Aroclor contaminants must meet the identification criteria in Exhibit D PEST, which requires analysis of the blank on two different GC Columns. Therefore, enter the date, time and instrument ID of both analyses of the blank on the pesticide method blank summary. The information on the two analyses is differentiated as Date Analyzed (1), Date Analyzed (2), etc. If the analyses were run simultaneously, the order of reporting is not important, but must be consistent with the information reported on all other pesticide forms. Otherwise, (1) shall be the first analysis, and (2) the second. Identify the GC Column and internal diameter as described previously.

Enter "Lab File ID" only if GC/MS confirmation was attempted. Otherwise, leave blank.

For all three fractions, as appropriate, summarize the samples associated with a given method blank in the table below the header, entering EPA Sample Number and Lab Sample ID. For volatiles, enter the Lab File ID and time of analysis of each sample. For semivolatiles, enter Lab File ID and Date Analyzed. For pesticides/Aroclors, enter the dates of both analyses as Date Analyzed (1) and Date Analyzed (2), as discussed above.

Number all pages as described in part A.

G. GC/MS Instrument Performance Check and Mass Calibration (Form V VOA and SV)

This form is used to report the results of GC/MS instrument performance check (previously known as "Tuning") for volatiles and semivolatiles, and to summarize the date and time of analysis of samples, standards, blanks, matrix spikes, and matrix spike duplicates associated with each analysis of the instrument performance check solution.

Complete the header information as in Instruction A. Enter the "Lab File ID" for the injection containing the instrument performance check solution (BFB for volatiles, DFTPP for semivolatiles). Enter the "Instrument ID". Enter the date and time of injection of the instrument performance check solution. Enter time as military time. For volatiles, indicate the purging method by entering "Y" for heated purge, and "N" for ambient temperature purge, as described previously. Water samples and medium soil sample extracts may be reported on the same Form V if analyzed together, as a single calibration may be applied to both sample types.

For each ion listed on the form, enter the percent relative abundance in the righthand column. Report relative abundances to the number of significant figures given for each ion in the ion abundance criteria column. Note that for both BFB and DFTPP, one or more of the high mass ions may exceed the abundance of the ion listed on the form as the nominal base peak, m/z 95 for BFB, and m/z 198 for DFTPP. Despite this possibility, all ion abundances are to be normalized to the nominal base peaks listed on Form V (see Exhibits D and E).

All relative abundances must be reported as a number. If zero, enter "0", not a dash or other non-numeric character. Where parentheses appear, compute the percentage of the ion abundance of the mass given in the appropriate footnote, and enter that value in the parentheses.

In the lower half of the form, list all samples, standards, blanks, matrix spikes, and matrix spike duplicates analyzed under that instrument performance check in <u>chronological order</u>, by time of analysis (in military time). Refer to Part A for specific instructions for identifying standards and blanks. Enter "EPA Sample No.", "Lab Sample ID", "Lab File ID", "Date Analyzed", and "Time Analyzed" for all standards, samples, blanks, matrix spikes, and matrix spike duplicates.

The GC/MS instrument performance check solution must be analyzed again twelve hours from the time of injection of the instrument performance check solution (BFB or DFTPP) listed at the top of the form. In order to meet these requirements, a sample, standard, blank, matrix spike, or matrix spike duplicate must be injected within twelve hours of the injection of the instrument performance check solution.

Number all pages as described in Instruction A.

H. GC/MS Initial Calibration Data (Form VI VOA, SV-1, SV-2)

After a GC/MS system has undergone an initial five-point¹ calibration at the specific concentration levels described in Exhibits D and E, and after all initial calibration criteria have been met, the laboratory must complete and submit a Form VI for each volatile or semivolatile target compound initial calibration performed which is relevant to the samples, blanks, matrix spikes, matrix spike duplicates in the SDG, regardless of when that calibration was performed.

Complete all header information as in Part A. Enter the "Case No." and "SDG No." for the current data package, regardless of the original Case for which the initial calibration was performed. Enter "Instrument ID" and the date(s) of the calibration. If the calendar date changes during the calibration procedure, the inclusive dates should be given on Form VI. Enter the injection times of the first and last of the standards analyzed under "Calibration Times".

¹ For Semivolatiles, eight compounds: 2,4-Dinitrophenol,

^{2,4,5-}Trichlorophenol, 2-Nitroaniline, 3-Nitroaniline, 4-Nitroaniline, 4-Nitrophenol, 4,6-Dinitro-2-Methylphenol, and Pentachlorophenol will only require a four-point initial calibration at 50, 80, 120, and 160 total nanograms because detection at less than 50 nanograms per injection is difficult. If a four-point calibration is performed for these compounds, leave RF20 blank.

For volatiles, enter Heated Purge, and GC Column, ID, as on Form V.

Enter the "Lab File ID" for each of the five calibration standards injected. Complete the response factor data for the five calibration points, and then calculate and report the average relative response factor (RRF) for all target compounds. For volatiles, report the response factors for the system monitoring compounds in the calibration standards. For semivolatiles, report the response factors for all surrogate compounds in the calibration standards. The laboratory must report the relative standard deviation (%RSD) for <u>all</u> compounds.

$$RSD - (SD/x) \times 100$$

Where

SD

 Standard Deviation of initial 5 response factors (per compound), is calculated as:

$$\sqrt{\sum_{i=1}^{n} \frac{(x_i - \bar{x})^2}{n-1}}$$

and \overline{x} - Mean of initial 5 response factors (per compound)

In order to be used for the analysis of samples or sample extracts, the volatile and semivolatile initial calibration must meet the acceptance criteria for relative response factors outlined in Exhibits D and E. The compounds for which criteria have been developed for minimum RRF and maximum %RSD are indicated on the form by an "*". All other compounds must meet a minimum RRF of 0.010.

I.

GC/EC Initial Calibration Data (Form VI PEST-1, PEST-2, PEST-3, PEST-4)

The initial calibration of pesticides and Aroclors involves the determination of retention times, retention time windows, and calibration factors. For single component pesticide target compounds, these data are calculated from the analyses of the Individual Standard Mixtures A and B at three different concentration levels. For the multicomponent target compounds, these data are calculated from a single point calibration.

For each set of three analyses of Individual Standard Mixture A (low point, mid point, and high point), and set of three analyses of Individual Standard Mixture B, during an initial calibration, complete one copy of Form VI for <u>each</u> GC column used. Thus, each initial calibration will require at least two forms for the Individual Mixture A analyses, and two for Individual Mixture B analyses. However, for each of the forms, half of the compounds will have no entries, as they are not in that mixture. Complete the header information as above. Enter the Instrument ID, GC Column, and ID as described previously. Enter the dates of analysis of the first and last of the three standards on each form under "Date(s) Analyzed". Under "Level (x low)", enter the concentration of the low point, mid point, and high point calibration standards as a multiplier of the low point. Therefore, for the low point, enter "1.0". The concentration of the mid point standard is specified in Exhibit D as ten times the low point, therefore, enter "10.0" for "mid". The high point standard must be at least 30 times the low point, but may be up to 100 times the low point, if that value lies within the linear range of the instrument, as specified in Exhibit D. Therefore, enter the appropriate multiplier to the high point standard concentration to one decimal place.

For each standard analyzed, enter the retention time of each applicable analyte in minutes and decimal minutes, under the appropriate concentration level. Calculate the mean retention time of each analyte from the three individual mixtures, and report it under "Mean RT". The retention time window of each analyte is defined as ± 0.04 minutes from the mean retention time, except for Methoxychlor, where the width of the window is ± 0.05 minutes. Calculate the retention time window for each analyte, using these specifications, and enter the lower limit of the window under RT Window "From", and the upper limit of the window under "To". The retention times of the surrogates are reported for <u>both</u> Individual mixtures, but the windows are only required to be calculated for Individual Mixture A.

For each three analyses of the same Individual Standard Mixture (A or B), the laboratory must also complete the calibration factor data on Form VI PEST-2. In a similar fashion as for the retention time data on Form VI PEST-1, prepare one form for each group of three standards, for each instrument and GC column used. Enter the concentration level of the standards in the same fashion as on Form VI PEST-1.

Enter the calibration factor for each compound in each of the standards, and calculate a mean calibration and a percent relative standard deviation (%RSD), and enter on the form. As with surrogate retention times, the calibration factors are only required from Individual Mixture A analyses.

In order to be used for sample analyses, the %RSD of the initial calibration factors must be less than or equal to 10.0 percent for all compounds except alpha-BHC, beta-BHC, gamma-BHC, DDT, Endrin, and Methoxyclor, where the %RSD must be less than or equal to 15.0 percent. These criteria apply to both GC columns.

For the multicomponent target compounds, the retention times, retention time windows, and calibration factor must be reported in a similar fashion for each single point calibration standard. For each multicomponent compound, the laboratory must select at least three peaks from each analyte, according to the specifications in Exhibit D. The retention and calibration factor data apply to <u>each peak</u>. Complete one version of Form VI PEST-3 for each GC column, for each initial calibration that applies to samples in the data package. Form VI is used also to report the results of analysis of the Resolution Check Solution that must begin each pesticide/Aroclor initial calibration sequence. The purpose of the Resolution Check Solution is to demonstrate for each initial calibration that the GC columns employed are capable of satisfactorily resolving the most difficult of the target analytes. One copy of Form VI PEST-4 is completed that covers both GC columns.

Complete the header information as described in Instruction A. Using the same assignment of first and second GC columns made for Form IV, enter the GC Column, ID, Instrument ID, and Date and Time Analyzed. Enter the "EPA Sample No." for the Resolution Check Standard. If simultaneous injections on a single GC are used, the EPA Sample No. may be the same for both Resolution Check Standards. If simultaneous injections were not used, use different suffixes to identify the standards.

In the boxes on the form, list each analyte, in retention time order, including both surrogate compounds. Thus, the order of analytes in the two boxes on a copy of this form will be different, due to the dissimilarity of the stationary phases of the two GC columns used. Enter the name of each target analyte in the Resolution Check Mixture as it appears on Form I PEST. Spell out the names of the surrogates as they appear on Form VII PEST-2.

Enter the retention time of each analyte from the analysis under "RT". Calculate the resolution between each pair of analytes according to the formulae in Exhibit D. The resolution is calculated as percentage of the height of the smaller of each pair of adjacent peaks. Enter the resolution between the first and second peaks on the line for the first analyte listed in the box. Enter the resolution between the second and third peaks on the line for the second analyte, and so on, until the resolutions of all possible pairs of adjacent analytes have been entered. NOTE: Only eight of the nine resolution fields will be filled. In order for these GC columns to be used for pesticide/Aroclor analyses, the resolution of all pairs of peaks listed on this form must be greater than or equal to 60.0%.

J. GC/MS Continuing Calibration Data (Form VII VOA, SV-1, SV-2)

For volatiles and semivolatiles, the Continuing Calibration Check form is used to report the calibration of the GC/MS system by the analysis of specific calibration standards. A Continuing Calibration Check form is required for each twelve (12) hour time period for both volatile and semivolatile target compound analyses.

The Contract laboratory must analyze calibration standards and meet all criteria outlined in Exhibits D and E for the minimum RRF and maximum percent difference between initial and continuing calibrations.

Complete all header information as in Instruction A. Enter instrument ID, date and time of continuing calibration, the Lab File ID of the continuing calibration standard, and dates and times of initial calibration (give inclusive dates if initial calibration is performed over more than one date). For volatiles, enter purge method and column as on Forms IV, V and VI. Using the appropriate Initial Calibration (volatile or semivolatile) fill in the average relative response factor (RRF) for each target compound, each system monitoring compound for volatiles, and each surrogate for semivolatiles. Report the relative response factor (RRF50) from the continuing calibration standard analysis. Calculate the Percent Difference (%D) for all compounds.

* Difference
$$-\frac{\overline{RRF_i} - RRF_c}{\overline{RRF_i}} \times 100$$

Where

RRF, - Average relative response factor from initial calibration.

RRF_c - Relative response factor from continuing calibration standard.

All semivolatile standards are analyzed at 50 total ng.

K. GC/EC Continuing Calibration (Form VII PEST)

The Calibration Verification Summary Form VII is used to report the results of the Performance Evaluation Mixtures (PEM), instrument blanks, and Individual Standard Mixtures A and B analyzed at the beginning and end of a twelve hour sequence. The laboratory must submit this form for each twelve hour sequence analyzed.

Complete the header information on each Form VII required according to the instructions in part A.

Enter the initial calibration date(s) analyzed. Give inclusive dates if initial calibration is performed over more than one date.

On Form VII PEST-1, enter the EPA Sample No., Lab Sample ID, Date Analyzed, and Time Analyzed for the instrument blank that preceded the twelve hour sequence (PIBLK). For the PEM that initiated or terminated the twelve hour sequence (PEM), enter the EPA Sample No., Lab Sample ID, Date Analyzed, and Time Analyzed.

In the table, report the retention time for each analyte in the PEM as well as the retention time windows. For each analyte in the PEM, enter the amount of the analyte calculated to be in the PEM, in nanograms to three decimal places, under "CALC AMOUNT". Enter the nominal amount of each analyte in the PEM under "NOM AMOUNT". Calculate the relative percent difference between the calculated amount and nominal amount for each analyte according to Exhibit D. Report the values under "RPD". Calculate the percent breakdown for endrin and 4,4'-DDT, and the combined percent breakdown in the PEM according to Exhibit D. Enter the values for the breakdown of endrin and 4,4'-DDT in their respective fields immediately under the table.

Form VII PEST-2 is used to report the results of the analyses of the instrument blank and the midpoint concentrations of Individual Standard Mixtures A and B that, along with the PEM, bracket each 12-hour period of sample analyses. One copy of Form VII PEST-2 must be completed each time the Individual Standard Mixtures are analyzed, for each GC column used. The form is completed in a fashion similar to Form VII, entering the EPA Sample No., Lab Sample ID, Date Analyzed, and Time Analyzed for the instrument blank immediately preceding the Individual Standard Mixtures A and B, and for the standards themselves. The upper table on the form contains the retention time and amount data for Individual Standard Mixture A compounds. The lower table contains the data for Mixture B. Enter the data in these tables in a fashion similar to that for the PEM. Complete copies of Form VII PEST-1 and PEST-2 for each standard reported on Form VIII PEST.

L. Internal Standard Area Summary (Form VIII VOA and SV)

This form is used to summarize the peak areas and retention times of the internal standards added to all volatile and semivolatile samples, blanks, matrix spikes, and matrix spike duplicates. The data are used to determine when changes in internal standard responses will adversely affect quantification of target compounds. This form must be completed each time a continuing calibration is performed, or when samples are analyzed under the same GC/MS instrument performance check as an initial calibration.

Complete the header information as in Instruction A. Enter the Lab File ID of the continuing calibration standard, as well as the date and time of analysis of the continuing calibration standard. If samples are analyzed immediately following an initial calibration, before another instrument performance check and a continuing calibration, Form VIII shall be completed on the basis of the internal standard areas of the 50 ug/L initial calibration standard for volatiles, and the 50 ng initial calibration standard for semivolatiles. Use the date and time of analysis of this standard, and its Lab File ID and areas in place of those of a continuing calibration standard.

For volatiles, enter purge method and column, as on Forms IV, V, VI, and VII.

From the results of the analysis of the continuing calibration standard, enter the area measured for each internal standard and its retention time (in decimal minutes) under the appropriate column in the row labeled "12 HOUR STD". For each internal standard, calculate the upper limit of the area as the area of the particular standard plus 100% of its area (i.e., two times the area in the 12 HOUR STD box), and the lower limit of the area as the area of the internal standard minus 50% of its area (i.e., one half the area in the 12 HOUR STD box). Report these values in the boxes labeled "UPPER LIMIT" and "LOWER LIMIT" respectively. Calculate the upper limit of the retention time as the retention of the internal standard plus 0.50 minutes (30 seconds), and the lower limit of the retention time as the retention time in the standard minus 0.50 minutes (30 seconds).

For each sample, blank, matrix spike, and matrix spike duplicate analyzed under a given continuing calibration, enter the EPA Sample Number and the area measured for each internal standard and its retention time. If the internal standard area is outside the upper or lower limits calculated above, flag that area with an asterisk (*). The asterisk must be placed in the far right hand space of the box for each internal standard area, directly under the "#" symbol. Similarly, flag the retention time of any internal standard that is outside the limits with an asterisk.

Number all pages as described in Instruction A.

M. Pesticide Analytical Sequence (Form VIII Pest)

This form is used to report the analytical sequence for pesticide analysis. At least one Form VIII PEST is required for each GC column used for pesticide/Aroclor analyses.

The laboratory shall complete all the header information as in Part A. Enter dates of analyses for the initial calibration, GC column, ID, and Instrument ID, as on Forms IV, VI, and VII.

At the top of the table, report the mean retention time for tetrachloro-m-xylene and decachlorobiphenyl calculated from the initial calibration sequence under "TCX" and "DCB", respectively. For every analysis associated with a particular analytical sequence starting with the initial calibration, enter the EPA Sample Number, Lab Sample ID, Date Analyzed, and Time Analyzed. Each sample analyzed as part of the sequence must be reported on Form VIII PEST even if it is not associated with the SDG. The laboratory may use the EPA Sample No. of "ZZZZZ" to distinguish all samples that are not part of the SDG being reported. Report the retention time of the surrogates for each analysis under "TCX RT" and "DCB RT". All sample analyses must be bracketed by acceptable analyses of instrument blanks, a PEM, and Individual Standard Mixtures A and B. Given the fact that the initial calibration may remain valid for some time (see Exhibit D), it is not necessary to report the data from 12-hour periods when no samples in an SDG were run. The laboratory must deliver the Form VIII for the initial calibration sequence, and Forms that include the PEMs and Individual Standard Mixtures that bracket any and all samples in the SDG. While the data for time periods between the initial calibration and samples in the SDG is not a routine deliverable, it must be made available on request during on-site evaluations, etc. Here again, non-EPA samples may be indicated with "ZZZZZ".

Flag all those values which do not meet the contract requirements by entering an asterisk (*) in the last column, under the "*". If the retention time cannot be calculated due to interfering peaks, leave the RT column <u>blank</u> for that surrogate, enter an asterisk in the last column, and <u>document</u> the problem in the SDG Narrative.

If more than a single copy of Form VIII PEST is required, enter the same header information on all subsequent pages for that GC Column and Instrument, and number each page as described in Part A.

Form VIII PEST is required for each for each GC system and for each GC column used to analyze target pesticides/Aroclors.

N. Pesticide Cleanup Summary (Form IX PEST-1, PEST-2)

This form summarizes the results of the checks performed for both cleanup procedures employed during the preparation of pesticide extracts for analysis. Form IX PEST-1 is used to report the results of the check of the Florisil cartridges used to process all sample extracts, and to associate the lot of cartridges with particular sample results. In this fashion, problems with a lot of cartridges may be tracked across many samples.

Complete the header information on each Form IX required, according to the instructions in Part A.

Enter the "Case No." and "SDG No." for the current data package, regardless of the original Case for which the cartridge check was performed. Enter the "Florisil Cartridge Lot Number". Enter under the "Date Analyzed" the date the Florisil cartridge check solution was analyzed.

Enter "GC Column" and "ID" for the GC columns used to determine the recovery of the analytes in the Florisil cartridge check solution, under "GC Column (1)", and "GC Column (2)", etc., as discussed previously.

In the upper table, enter the amount of spike added and spike recovered in nanograms for each analyte.

Calculate to the nearest whole percent, and enter the percent recovery in the "% REC" field. Flag each spike recovery outside the QC limits with an asterisk (*). The asterisk must be placed in the last space in the "% REC" column, under the "#" symbol.

In the lower table, enter the "EPA Sample No.", the "Lab Sample ID", and "Date Analyzed" for each sample and blank that was cleaned up using this lot of Florisil cartridges.

Number the Form IX pages as described in Part A.

Form IX PEST-2 summarizes the results of the calibration of the Gel Permeation Chromatography device (GPC) that must be used to process all soil sample extracts for pesticide/Aroclor analyses. Calibration of the GPC is required at least once every 7 days, and each time the GPC column is repacked. Complete all header information as in Part A. Enter an identifier for the GPC Column, and the date of calibration in the appropriate fields. Enter the two "GC Column" and "ID" fields, as discussed above.

For each of the pesticide matrix spike compounds listed in the box in the upper portion of the form, enter the amount of the spike added to the GPC column in ng, and the amount recovered, also in ng. Calculate the percent recovery of each analyte, and enter these values on the form, to the nearest percent. Compare the recoveries to the QC limits shown on the form, and flag all those values outside the limits with an asterisk (*) in the column under the "#" symbol.

For each sample in the data package that was subjected to GPC under this calibration, enter the EPA Sample No., Lab Sample ID, and the date of <u>both</u> analyses in the lower portion of the form.

If more than one copy of Form IX PEST-2 is required, number all pages as described in Instruction A.

0. Pesticide/Aroclor Identification (Form X PEST-1, PEST-2)

This form summarizes the quantitations of all target pesticides/Aroclors detected in a given sample. It reports the retention times of the compound on both columns on which it was analyzed, as well as the retention time windows of the standard for that compound on both of these columns. In addition, it is used to report the concentration determined from each GC column, and the percent difference between the two quantitative results. Separate copies of Form X are used for single component analytes and multicomponent analytes.

Copies of Form X are required for each sample, blank, matrix spike, and matrix spike duplicate in which target pesticides or Aroclors are detected. If none are detected in a given sample, no copy of Form X is required for that sample.

Complete the header information as in Instruction A. Enter the GC Column, and ID for each of the two columns, one as GC Column (1), the other as (2), as described previously. Enter the Instrument ID associated with each GC column directly below.

For each single component pesticide detected, enter the name of the compound under "ANALYTE" as it appears on Form I. Enter the retention times on each column of the compounds detected in the sample next to the appropriate column designation (1 or 2). Enter the retention time windows on each column from the initial calibration standard. These data must correspond with those on Form VI, and are entered in a similar manner. The lower value is entered under the "FROM" column, the upper value under the "TO" column.

Enter the concentration calculated from each GC column under the column labeled "CONCENTRATION". The units are the same as those used on Form I, ug/L for water samples, and ug/Kg for soil samples. However, do not enter any units on Form X. Calculate the percent difference between the concentrations entered, and report it to a tenth of a percent under "BD".

The <u>lower</u> of the two concentrations is reported on Form I for each pesticide compound. The lower concentration is used because, if present, co-eluting interferences are likely to increase the calculated concentration of any target compound. If the percent difference between the calculated concentrations is greater than 25.0 percent, flag the concentration on Form I, as described previously. This will alert the data user to the potential problems in quantitating this analyte.

If more pesticide compounds are identified in an individual sample than can be reported on one copy of Form X, then complete as many additional copies of Form X as necessary, duplicating all header information, and numbering the pages as described in Instruction A.

Multicomponent analytes detected in samples are reported on a separate version of Form X. Complete the header information and Instrument and GC Column fields as described above. For multicomponent analytes, it is necessary to report the retention time and concentration of each peak chosen for quantitation in the target analyte, in fashion similar to that for single component pesticides. The concentrations of all peaks quantitated (three are required, up to five may be used) are averaged to determine the mean concentration. Report the lower of the two mean concentrations on Form I. Flag this value as described previously, if the mean concentrations from the two GC columns differ by more than 25.0 percent.

If more multicomponent compounds are identified in an individual sample than can be reported on one copy of Form X, then complete as many additional copies of Form X as necessary, duplicating all header information, and numbering the pages as described in Instruction A.

> THE FOLLOWING ARE DOCUMENT CONTROL FORMS (To be submitted as hardcopy only)

P. Sample Log-In Sheet (Form DC-1)

This form is used to document the receipt and inspection of sample containers and samples. One original of Form DC-1 is required for each sample shipping container. If the samples in a single sample shipping container must be assigned to more than one Sample Delivery Group, the original Form DC-1 shall be placed with the deliverables for the Sample Delivery Group of the lowest Arabic number and a copy of Form DC-1 must be placed with the deliverables for the other Sample Delivery Group(s). The copies should be identified as "copy(ies)," and the location of the original should be noted on the copies.

Sign and date the airbill (if present). Examine the shipping container and record the presence/absence of custody seals and their condition (i.e., intact, broken) in item 1 on Form DC-1. Record the custody seal numbers in item 2. Open the container, remove the enclosed sample documentation, and record the presence/absence of chain-of-custody record(s), SMO forms (i.e., Traffic Reports, Packing Lists), and airbills or airbill stickers in items 3-5 on Form DC-1. Specify if there is an airbill present or an airbill sticker in item 5 on Form DC-1. Record the airbill or sticker number in item 6.

Remove the samples from the shipping container(s), examine the samples and the sample tags (if present), and record the condition of the sample bottles (i.e., intact, broken, leaking) and presence of absence of sample tags in items 7 and 8 on Form DC-1.

Review the sample shipping documents and complete the header information described in Instruction A. Compare the information recorded on all the documents and samples and circle the appropriate answer in item 9 on Form DC-1.

If there are no problems observed during receipt, sign and date (include time) Form DC-1, the chain-of-custody record, and Traffic Report, and write the sample numbers on Form DC-1. Record the appropriate sample tags and assigned laboratory numbers if applicable. The log-in date should be recorded at the top of Form DC-1 and the date and time of cooler receipt at the laboratory should be recorded in items 10 and 11. Record the fraction designation (if appropriate) and the specific area designation (e.g., refrigerator number) in the Sample Transfer block located in the bottom left corner of Form DC-1. Sign and date the Sample Transfer block. Cross out unused columns and spaces.

If there are problems observed during receipt or an answer marked with an asterisk (i.e., "absent*") was circled, contact SMO and document the contact as well as resolution of the problem on a CLP Communication Log. Following resolution, sign and date the forms as specified in the preceding paragraph and note, where appropriate, the resolution of the problem.

Q. Document Inventory Sheet (Form DC-2)

This form is used to record the inventory of the SDG File Purge documents and count of documents in the original Sample Data Package which is sent to the Region.

Organize all EPA-CSF documents as described in Exhibit B, Section II and Section III. Assemble the documents in the order specified on Form DC-2 and Section II and III, and stamp each page with a consecutive number. (Do not number the DC-2 form). Inventory the CSF by reviewing the document numbers and recording page numbers ranges in the columns provided in the Form DC-2. If there are no documents for a specific document type, enter an "NA" in the empty space. Certain laboratory specific documents related to the CSF may not fit into a clearly defined category. The laboratory should review DC-2 to determine if it is most appropriate to place them under No. 7, 8, 9, or 10. Category 10 should be used if there is no appropriate previous category. These types of documents should be described or listed in the blanks under each appropriate category. SECTION IV

DATA REPORTING FORMS

•

VOLATILE ORGAN	1A NICS ANALYSIS DATA SH		PA SAMPLE	NO.
Lab Name:	Contract:			
Lab Code: Case No.	: SAS No.: _	SDG No).:	•
Matrix: (soil/water)	La	b Sample ID: _		
Sample wt/vol:((g/mL) La	b File ID: _		
Level: (low/med)	Da	te Received: _		•
% Moisture: not dec	Da	te Analyzed: _		•
GC Column: ID:	(mm) Di	lution Factor:		-
Soil Extract Volume:	_(uL) So	oil Aliquot Vol	ume:	(uL)
CAS NO. COMPO		ATION UNITS: ug/Kg)	Q	
74-87-3	Dmethane 1 Chloride roethane			

3/90

1**B**

EPA SAMPLE NO.

SEMIVOLATI	1B LE ORGANICS ANALYSIS	5 DATA SHEET		SAMPLE NO.
Lab Name:	Coi	ntract:		
Lab Code:	Case No.: Si	AS No.:	SDG No.:	
Matrix: (soil/water)		Lab Sa	mple ID:	
Sample wt/vol:	(g/mL)		.le ID:	
Level: (low/med)			eceived:	
<pre>% Moisture:</pre>			xtracted:	
Concentrated Extract			.on Factor:	
Injection Volume:		011401	on ractor: _	
GPC Cleanup: (Y/N)	pr:			
CAS NO.	COMPOUND	CONCENTRATIO		Q
53-57-6 541-73-1	Phenol bis(2-Chloroethy 2-Chlorophenol 1,3-Dichlorobenze 1,2-Dichlorobenze 	ene ol ene loropropane) ropylamine ol enzene ene entadiene henol ene ene		
208-96-8 606-20-2 99-09-2	Acenaphthylene 2,6-Dinitrotolue 3-Nitroaniline			

EPA SAMPLE NO. 10 SEMIVOLATILE ORGANICS ANALYSIS DATA SHEET Lab Name:_____ Contract:_____ Lab Code: _____ Case No.: ____ SAS No.: ____ SDG No.: ____ Lab Sample ID: _____ Matrix: (soil/water)_____ Sample wt/vol: _____(g/mL)____ Lab File ID: _____ Date Received: Level: (low/med) % Moisture: _____ decanted: (Y/N) ____ Date Extracted: Date Analyzed: Concentrated Extract Volume: ____(uL) Dilution Factor: Injection Volume: ____(uL) GPC Cleanup: (Y/N) ____ pH:____ CONCENTRATION UNITS: 0 CAS NO. (ug/L or ug/Kg)_____ COMPOUND 51-28-5-----2,4-Dinitrophenol_____ 100-02-7-----4-Nitrophenol_____ 132-64-9----Dibenzofuran_ 121-14-2----2,4-Dinitrotoluene____ 84-66-2----Diethylphthalate 7005-72-3-----4-Chlorophenyl-phenylether___ 86-73-7----Fluorene 100-01-6----4-Nitroaniline 534-52-1-----4,6-Dinitro-2-methylphenol_ 86-30-6----Nitrosodiphenylamine (1) 101-55-3-----4-Bromophenyl-phenylether 118-74-1-----Hexachlorobenzene_ 87-86-5-----Pentachlorophenol___ 85-01-8----Phenanthrene 120-12-7----Anthracene 86-74-8----Carbazole 84-74-2----Di-n-butylphthalate___ 206-44-0----Fluoranthene 129-00-0----Pyrene 85-68-7----Butylbenzylphthalate 91-94-1-----3,3'-Dichlorobenzidine__ 56-55-3----Benzo(a)anthracene 218-01-9----Chrysene 117-81-7----bis(2-Ethylhexyl)phthalate 117-84-0----Di-n-octylphthalate_ 205-99-2----Benzo(b) fluoranthene 207-08-9----Benzo(k)fluoranthene__ 50-32-8----Benzo(a)pyrene_ 193-39-5-----Indeno(1,2,3-cd)pyrene 53-70-3-----Dibenz(a,h)anthracene 191-24-2----Benzo(g,h,i)perylene____

(1) - Cannot be separated from Diphenylamine

EPA SAMPLE NO.

PESTICII	1D DE ORGANICS ANAI	LYSIS DATA S	HEET	EPA	SAMPLE	N(
ab Name:		Contract:_		_		
ab Code: (Case No.:	SAS No.:	S	DG No.:		-
atrix: (soil/water)		L	ab Sample	ID:		
ample wt/vol:	(g/mL)	L	ab File ID):		
Moisture:	decanted: (Y/N)) D	ate Receiv	'ed:		
xtraction: (SepF/Co	ont/Sonc)	D	ate Extrac	:ted:		
oncentrated Extract	Volume:	_(uL) D	ate Analyz	ed:		
njection Volume:	(uL)		ilution Fa			
PC Cleanup: (Y/N)	pH:	s	ulfur Clea	nup: (Y/	'N)	
CAS NO.	COMPOUND		TRATION UNI		Q	
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Heptachlor eg Endosulfan I Dieldrin 4,4'-DDE Endrin Endosulfan I Endosulfan s 4,4'-DDT Endosulfan s Endrin keton Endrin aldeh Endrin aldeh gamma-Chlord	indane) poxide I ulfate e yde ane				

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		ALYSIS DATA		I 	EPA SAMP	LE NO.
		FIED COMPOUN				
Lab Name:						
Lab Code: Case		SAS No.:		SDG No	».:	
Matrix: (soil/water)			Lab Sampl	e ID: _		
Sample wt/vol:	(g/mL)		Lab File	ID: _		
Level: (low/med)			Date Rece	ived:		
<pre>% Moisture: not dec</pre>			Date Anal	yzed: _		. <u></u>
GC Column: ID:	(mm)	l i i i i i i i i i i i i i i i i i i i	Dilution	Factor:		
Soil Extract Volume:	(uL)		Soil Alig	uot Vol	ume:	(uL)
Number TICs found:			TRATION U or ug/Kg)	-		
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1F SEMIVOLATILE ORGANICS ANALYSIS DATA TENTATIVELY IDENTIFIED COMPOU	
Lab Name: Contract	
Lab Code: Case No.: SAS No.	SDG No.:
Matrix: (soil/water)	Lab Sample ID:
Sample wt/vol:(g/mL)	Lab File ID: '
Level: (low/med)	Date Received:
<pre>% Moisture: decanted: (Y/N)</pre>	Date Extracted:
Concentrated Extract Volume:(uL)	Date Analyzed:
Injection Volume:(uL)	Dilution Factor:
GPC Cleanup: (Y/N) pH:	
CONCE	INTRATION UNITS:

Number TICs found:

CONCENTRATION UNITS: (ug/L or ug/Kg)_____

CAS NUMBER	COMPOUND NAME	 RT	EST. CONC.	Q
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b Name:		(Contract	t:		
b Code:	Case No.:		SAS No	.:	SD	G No.:
	EPA SAMPLE NO.	(TOL) #	•	(DCE) #		OUT
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SM	C1 (TOL) = Tol C2 (BFB) = Bro C3 (DCE) = 1,2	mofluor	obenzen	(e (88-110) 86-115) 76-114))
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	Values outside					lits
D	System Monitor	ing Com	nound d	iluted		

2A WATER VOLATILE SYSTEM MONITORING COMPOUND RECOVERY 2B

SOIL VOLATILE SYSTEM MONITORING COMPOUND RECOVERY

Lab	Name:	· · · · · · · · · · · · · · · · · · ·	Contract:	
Lab	Code:	Case No.:	SAS No.:	SDG No.:

Level: (low/med)

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D System Monitoring Compound diluted out

page ____ of ____

FORM II VOA-2

2C WATER SEMIVOLATILE SURROGATE RECOVERY

Lab	Name:	······································	Contract:				
Lab	Code:	Case No.:	SAS No.:	SDG No.:			

EPA	51	S2	53	S4	S5	S6	S7	S8	TO'
SAMPLE NO.	(NBZ) #	(FBP)#		(PHL)#	(2FP)#	(TBP) #	(2CP) #	(DCB) #	jou
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QC LIMITS S1 (NBZ) = Nitrobenzene-d5 (35-114) S2 (FBP) = 2-Fluorobiphenyl (43-116) S3 (TPH) = Terphenyl-d14 (33 - 141)S4 (PHL) = Phenol-d5 (10 - 110)S5 (2FP) = 2-Fluorophenol (21 - 110)S6 (TBP) = 2,4,6-Tribromophenol (10-123) S7 (2CP) = 2-Chlorophenol-d4 (advisory) (33-110) S8 (DCB) = 1,2-Dichlorobenzene-d4 (16-110)(advisory)

Column to be used to flag recovery values
* Values outside of contract required QC limits
D Surrogate diluted out

page ____ of ____

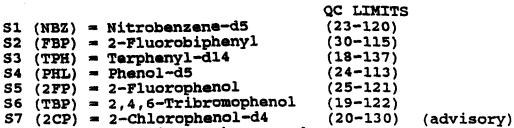
2D SOIL SEMIVOLATILE SURROGATE RECOVERY

Lab	Name:	Contract:

Lab Code: _____ Case No.: ____ SAS No.: ____ SDG No.: ____

Level: (low/med)

EPA	51	S2	S 3	S4	S5	S6	S7	S8	TOT
SAMPLE NO	• (NBZ) #	(FBP) #	(TPH) #	(PHL) #	(2FP) #	(TBP) #	(2CP) #	(DCB) #	I OUT
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S8 (DCB) = 1,2-Dichlorobenzene-d4 (20-130) (advisory)

Column to be used to flag recovery values
* Values outside of contract required QC limits
D Surrogate diluted out

2E WATER PESTICIDE SURROGATE RECOVERY

Lab	Name:		Contract:		
Lab	Code:	Case No.:	SAS No.:	SDG No.:_	
GC (Column(1):	ID:	(mm) GC Column((2):	ID:(mm)

EPA		TCX	1	TCX	2	DCB	1	DCB	2	OTHER	OTHER	TOT
SAMPLE	NO.	\$REC	#	%REC	#	\$REC	#	*REC	#	(1)	(2)	I OUI
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ADVISORY

			QC LIMITS
TCX	-	Tetrachloro-m-xylene	(60-150)
		Decachlorobiphenyl	(60-150)

Column to be used to flag recovery values
* Values outside of QC limits
D Surrogate diluted out

page __ of ___

FORM II PEST-1

2F SOIL PESTICIDE SURROGATE RECOVERY

Lab	Name:		Contract:	t:			
Lab	Code:	Case No.:	SAS No.:	SDG No.:			
GC	Column(1):	ID:()	mm) GC Column(2):	ID:(mm)			

EPA		TCX	1	TCX	2	DCB	1	DCB		OTHER	OTHER	TOT
SAMPLE	NO.	%REC	#	%REC	#	%REC	#	%REC	#	(1)	(2)	OUT
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4		i ——								l	1	1
5				i						l	1	
6			_							l	I	
7		1		i				1		l		
8		i				1				l	1	
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ADVISORY

			QC LIMITS
TCX	=	Tetrachloro-m-xylene	(60-150)
DCB	=	Decachlorobiphenyl	(60-150)

Column to be used to flag recovery values
* Values outside of QC limits
D Surrogate diluted out

page __ of ___

3A

WATER VOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:	·····	Contract:	
Lab	Code:	Case No.:	SAS No.:	SDG No.:
Mati	rix Spike - EPA	Sample No.:		·

COMPOUND	SPIKE	SAMPLE	MS	MS	QC.
	ADDED	CONCENTRATION	CONCENTRATION	*	LIMITS
	(ug/L)	(ug/L)	(UG/L)	REC #	REC.
1,1-Dichloroethene Trichloroethene Benzene Toluene Chlorobenzene					61-145 71-120 76-127 76-125 75-130

COMPOUND	SPIKE ADDED (ug/L)	MSD CONCENTRATION (Ug/L)	MSD % REC #	8 RPD #	RPD	IMITS
1,1-Dichloroethene Trichloroethene Benzene Toluene Chlorobenzene					11 13	61-145 71-120 76-127 76-125 75-130

Column to be used to flag recovery and RPD values with an asterisk

* Values outside of QC limits

 RPD:______out of _____outside limits

 Spike Recovery:______out of _____outside limits

3B

SOIL VOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:		Contract:				
Lab	Code:	Case No.:	SAS No.:	SDG No.:			
Mati	rix Spike - EPA	Sample No.:	Level:(lo	w/med)			

COMPOUND	SPIKE ADDED (ug/Kg)	(ug/Kg)	MS CONCENTRATION (ug/Kg)	MS % REC #	QC. LIMITS REC.
1,1-Dichloroethene Trichloroethene Benzene Toluene Chlorobenzene					59-172 62-137 66-142 59-139 60-133

COMPOUND	SPIKE ADDED (ug/Kg)	MSD CONCENTRATION (ug/Kg)	MSD % REC #	 RPD	IMITS REC.
1,1-Dichloroethene Trichloroethene Benzene Toluene Chlorobenzene				22 24 21 21 21 21	59-172 62-137 66-142 59-139 60-133

Column to be used to flag recovery and RPD values with an asterisk

* Values outside of QC limits

RPD:	out of	outside	limits	
Spike	Recovery:	out of	outside	limits

3C

WATER SEMIVOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:	Contract:	

Lab Code: _____ Case No.: _____ SAS No.: _____ SDG No.: _____

Matrix Spike - EPA Sample No.:

COMPOUND	SPIKE	SAMPLE	MS	MS	QC.
	ADDED	CONCENTRATION	CONCENTRATION	%	LIMITS
	(ug/L)	(ug/L)	(ug/L)	REC #	REC.
Phenol 2-Chlorophenol 1,4-Dichlorobenzene N-Nitroso-di-n-prop.(1) 1,2,4-Trichlorobenzene 4-Chloro-3-methylphenol Acenaphthene 4-Nitrophenol 2,4-Dinitrotoluene Pentachlorophenol Pyrene					12-110 27-123 36-97 41-116 39-98 23-97 46-118 10-80 24-96 9-103 26-127

COMPOUND	SPIKE ADDED (ug/L)	MSD CONCENTRATION (ug/L)	MSD % REC #	% RPD #		IMITS REC.
Phenol 2-Chlorophenol 1,4-Dichlorobenzene N-Nitroso-di-n-prop.(1) 1,2,4-Trichlorobenzene 4-Chloro-3-methylphenol Acenaphthene 4-Nitrophenol 2,4-Dinitrotoluene Pentachlorophenol Pyrene					42 40 28 38 28 42 31 50 38 50 31	12-110 27-123 36-97 41-116 39-98 23-97 46-118 10-80 24-96 9-103 26-127

(1) N-Nitroso-di-n-propylamine

Column to be used to flag recovery and RPD values with an asterisk
* Values outside of QC limits

RPD:______ out of ______ outside limits **Spike Recovery:**_____ out of ______ outside limits

3D SOIL SEMIVOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:	<u> </u>	Contract:			
Lab	Code:	Case No.:	SAS No.:	SDG No.:		
Mata	rix Spike - EPA	Sample No.:	Level: (low	/med)		

COMPOUND	SPIKE ADDED (ug/Kg)	SAMPLE CONCENTRATION (ug/Kg)	MS CONCENTRATION (ug/Kg)	MS t REC #	QC. LIMITS REC.
Phenol 2-Chlorophenol 1,4-Dichlorobenzene					26- 90 25-102 28-104
N-Nitroso-di-n-prop.(1) 1,2,4-Trichlorobenzene 4-Chloro-3-methylphenol					41-126 38-107 26-103
Acenaphthene 4-Nitrophenol 2,4-Dinitrotoluene					31-137 11-114 28- 89
Pentachlorophenol Pyrene					17-109 35-142

COMPOUND	SPIKE ADDED (ug/Kg)	MSD CONCENTRATION (ug/Kg)	MSD * REC #	* RPD #	QC L RPD	IMITS REC.
Phenol 2-Chlorophenol 1,4-Dichlorobenzene N-Nitroso-di-n-prop.(1) 1,2,4-Trichlorobenzene 4-Chloro-3-methylphenol Acenaphthene 4-Nitrophenol 2,4-Dinitrotoluene Pentachlorophenol Pyrene					35 50 27 38 23 33 19 50 47 47 36	26- 90 25-102 28-104 41-126 38-107 26-103 31-137 11-114 28- 89 17-109 35-142

(1) N-Nitroso-di-n-propylamine

Column to be used to flag recovery and RPD values with an asterisk
* Values outside of QC limits

RPD:______ out of ______ outside limits Spike Recovery:______ out of ______ outside limits

3E

WATER PESTICIDE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:		Contract:			
Lab	Code:	Case No.:	SAS No.:	SDG No.:		
Mati	rix Spike - EPA	Sample No.:				

COMPOUND	SPIKE	SAMPLE	MS	MS	QC.
	ADDED	CONCENTRATION	CONCENTRATION	t	LIMITS
	(ug/L)	(ug/L)	(Ug/L)	REC #	REC.
gamma-BHC (Lindane) Heptachlor Aldrin Dieldrin Endrin 4,4'-DDT					56-123 40-131 40-120 52-126 56-121 38-127

COMPOUND	SPIKE ADDED (ug/L)	MSD CONCENTRATION (ug/L)	MSD ¥ REC #	* RPD #		INITS
gamma-BHC (Lindane) Heptachlor Aldrin Dieldrin Endrin 4,4'-DDT					15 20 22 18 21 27	56-123 40-131 40-120 52-126 56-121 38-127

Column to be used to flag recovery and RPD values with an asterisk

* Values outside of QC limits

RPD: _____ out of _____ outside limits Spike Recovery: _____ out of _____ outside limits

3F

SOIL PESTICIDE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY

Lab	Name:_					Contract				
Lab	Code:			Case No.:		SAS No.	:	SDG	No.:	
Mati	rix Spi	ke -	EPA	Sample No.:	:					

	SPIKE	SAMPLE	MS	MS	QC.
	ADDED	CONCENTRATION	CONCENTRATION	*	LIMITS
COMPOUND	(ug/Kg)	(ug/Kg)	(ug/Kg)	REC #	REC.

gamma-BHC (Lindane)					46-127
Heptachlor		1			35-130
Aldrin					34-132
Dieldrin					31-134
Endrin					42-139
4,4'-DDT			l		23-134
					II

COMPOUND	SPIKE ADDED (ug/Kg)	MSD CONCENTRATION (ug/Kg)	MSD 8 REC #	* RPD #	QC L	IMITS
gamma-BHC (Lindane) Heptachlor Aldrin Dieldrin Endrin 4,4'-DDT					50 31 43 38 45 50	46-127 35-130 34-132 31-134 42-139 23-134

Column to be used to flag recovery and RPD values with an asterisk

* Values outside of QC limits

 RPD:
 out of
 outside limits

 Spike Recovery:
 out of
 outside limits

	VOLATILE MET	4A HOD BLANK	SUMMARY	EPA	SAMPLE	NO.
Lab Name:			Contract:			
Lab Code:	Case No.	:	SAS No.:	SDG No.:		-
Lab File ID:			Lab Sai	mple ID:		
Date Analyzed:			Time A	nalyzed:		
GC Column:	ID:	(mm)	Heated	Purge: (Y/N)		
Instrument ID:						

THIS METHOD BLANK APPLIES TO THE FOLLOWING SAMPLES, MS AND MSD:

		LAB	LAB	TIME
	EPA	SAMPLE ID	FILE ID	ANALYZED
	SAMPLE NO.			
01				
02				
03				
04				
05				
06				
07				
80	· · · · · · · · · · · · · · · · · · ·			
09				
10				
11				
12	1			
13	·			
14				·
15	·			
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18				
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22				
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27				1
28				1
29				·
30	and the second design of the s		I	

COMMENTS:

page ____ of ____

4B

SEMIVOLATILE METHOD BLANK SUMMARY

EPA SAMPLE NO.

1

Lab Name:		Contract	t:		
Lab Code:	Case No.:	SAS No.	• •	SDG	No.:
Lab File ID:			Lab Sample	ID:	. <u></u>
Instrument ID:			Date Extrac	ted	
Matrix: (soil/water)		Date Analyz	ed:	
Level: (low/med)			Time Analy:	ed:	

THIS METHOD BLANK APPLIES TO THE FOLLOWING SAMPLES, MS AND MSD:

	EPA	LAB	LAB	DATE
	SAMPLE NO.	SAMPLE ID	FILE ID	ANALYZED
01				
02				
03				
04				
06				
07				
08		· ·		
09		·	•	
10				
11				
12				
13				
14				
15				
16				
17				
18 19		 		l
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21		l		1
22	 	 		
23				·
24				
25	1			
26				
27	1			
28				
29				
30	l			

4C PESTICIDE METHOD BLANK	SUMMARY
Lab Name:C	ontract:
Lab Code: Case No.:	SAS No.: SDG No.:
Lab Sample ID:	Lab File ID:
Matrix:(soil/water)	Extraction: (SepF/Cont/Sonc)
Sulfur Cleanup: (Y/N)	Date Extracted:
Date Analyzed (1):	Date Analyzed (2):
Time Analyzed (1):	Time Analyzed (2):
Instrument ID (1):	Instrument ID (2):
	GC Column (2): ID:(mm)

THIS METHOD BLANK APPLIES TO THE FOLLOWING SAMPLES, MS AND MSD:

EPA		LAB		DATE			ATE	
SAMPLE		SAMPLE	ID	ANALYZED	1	ANAL	YZED	2
01	I							
02								
031	{			1				
04	1			· .				
05								
06	{			·				
07								
08				1				
09	!							
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11	{			/				
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14				'				
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17	!					¦		
18	!					¦		
19	1			!		¦		
20					_	۰ <u></u>		
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25				¦		¦		
26				I		J		

COMMENTS:

page __ of __

FORM IV PEST

5A VOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK BROMOFLUOROBENZENE (BFB)

Lab	Name:	Contract:			
Lab	Code: Case No.:	SAS No.: SDG No.:			
Lab	File ID:	BFB Injection Date:			
Inst	crument ID:	BFB Injection Time:			
GC (Column: ID:(mm)	Heated Purge: (Y/N)			

m/e	ION ABUNDANCE CRITERIA	<pre>% RELATIVE ABUNDANCE</pre>
50	8.0 - 40.0% of mass 95	
75	30.0 - 66.0% of mass 95	
95	Base peak, 100% relative abundance	
96	5.0 - 9.0% of mass 95	
173	Less than 2.0% of mass 174	
174	50.0 - 120.0% of mass 95	
175	4.0 - 9.0 % of mass 174	
176	93.0 - 101.0% of mass 174	
177	5.0 - 9.0% of mass 176	()2

1-Value is % mass 174 2-Value is % mass 176

THIS, CHECK APPLIES TO THE FOLLOWING SAMPLES, MS, MSD, BLANKS, AND STANDARDS:

EPA	LAB	LAB	DATE	TIME
SAMPLE NO		FILE ID	ANALYZED	ANALYZED
01;				
02				
03				
04				
05				· ····
06				
07	[
08				
09				
10				
11	!			
12				
13				
14				
15				
16	{			
17 18	{			
19				
201		· · · · · · · · · · · · · · · · · · ·		
21				
22				
22	l	, <u> </u>		

page ____ of ____

ŚΒ SEMIVOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK DECAFLUOROTRIPHENYLPHOSPHINE (DFTPP)

Lab	ab Name:			Contract:							
Lab	Code:		Case	No.:		SAS	No.:		SDG	No.:	<u> </u>
Lab	File I	D:				ſ	FTPP	Injection	1 Dat	:e:	

Instrument ID: _____

DFTPP Injection Time:

m/e	ION ABUNDANCE CRITERIA	& RELATIVE
51	30.0 - 80.0% of mass 198	
68	Less than 2.0% of mass 69	
69	Mass 69 relative abundance	;;;;;;
70	Less than 2.0% of mass 69	
127	25.0 - 75.0% of mass 198	
197	Less than 1.0% of mass 198	
198	Base Peak, 100% relative abundance	
199	5.0 to 9.0% of mass 198	
275	10.0 - 30.0% of mass 198	
365	Greater than 0.75% of mass 198	
441	Present, but less than mass 443	
442	40.0 - 110.0% of mass 198	
443	15.0 - 24.0% of mass 442	()2
I	1-Value is % mass 69 2-Value	is % mass 442

THIS CHECK APPLIES TO THE FOLLOWING SAMPLES, MS, MSD, BLANKS, AND STANDARDS:

!	EPA	LAB	LAB	DATE	TIME
	SAMPLE NO.	SAMPLE ID	FILE ID	ANALYZED	ANALYZED
01					
02				[
03					
04					
05					
06					
07					
08					
09					
10					
11					
12 13					
14				i	
15				·	
16					
17	[, 	······································	······	
18			·····	/	
19	i			1	
201	/		1	(
21					j
22	· · ·]

page ___ of ___

3/90

6A

VOLATILE ORGANICS INITIAL CALIBRATION DATA

Lab Name:		Contrac	::					
Lab Code: Case	No.:	SAS No	».:	si	OG No.:		-	
Instrument ID:	Calibrati	on Date	(s):				-	
Heated Purge: (Y/N)	Calibrati	on Times	5:					
GC Column: ID:	(mm)							
LAB FILE ID: RR	F10 =		RRF2	and the second se				
RRF50 = RR	F100=		RRF20	0=	· · · · · · · · · · · · · · · · · · ·			
COMPOUND	RRF10	1			RRF200		۶ RSD	
Chloromethane								
Dromonections		I			·			
Vinyl Chloride	**	I			l		·	
Chloroethane								
Methylene Chloride		!						
Acetone	!	!			[
Carbon Disulfide		ļ	·		·			
1,1-Dichloroethene	*	!						
1,1-Dichloroethane	*				·			
1,2-Dichloroethene (tota	1)_!		!					
Chloroform	**				!			
1,2-Dichloroethane	*		·				· `	
2-Butanone	I			!				
1,1,1-Trichloroethane	**		!					
Carbon Tetrachloride	*	I						
Bromodichloromethane	*			!		I		
1,2-Dichloropropane	I	<u> </u>			ļ	l		
cis-1,3-Dichloropropene	*	· /			I]		
Trichloroethene	*	l			I	l	I	
Dibromochloromethane	*		I			I	I	
1,1,2-Trichloroethane	*		I	I	I	I		
Benzene	**	1		!	1	[۱	
trans-1,3-Dichloroproper	ne*	<u> </u>				I	I	
Bromoform	*	.1					l	
4-Methyl-2-Pentanone			۱		I	I	ł	
2-Hexanone					1			
Tetrachloroethene	**				۱	1	1	
1,1,2,2-Tetrachloroethar	1e*		I			I		
Toluene	**						I	
Chlorobenzene	**	I				1	I	
Ethylbenzene	**	_					I	
Styrene	**	_]				<u> </u>	I	
Xylene (total)	**	_[!				l	
		=====		======			=====	
Toluene-d8	I				1	I	I	
Bromofluorobenzene	**	_ 1				1	۱	
1,2-Dichloroethane-d4		. I			1		I	
· · · · · · · · · · · · · · · · · · ·			1				1.	

* Compounds with required minimum RRF and maximum %RSD values. All other compounds must meet a minimum RRF of 0.010. 6B

SEMIVOLATILE ORGANICS INITIAL CALIBRATION DATA

Lab Name:		Contra	ct:						
Lab Code: Case No.:		SAS N	o.:	SI	DG No.:		-		
Instrument ID: Ca	librati	on Date	(s):				-		
	Calibration Times:								
Ca	librati	on Time	s:				-		
LAB FILE ID: RRF20	2		RRF50				l		
RRFSO = RRF120			RRF16	0=		_			
					1 1				
COMPOUND	RRF20	RRF50	RRF80		RRF160		RSD		
Phenol	*		i				*		
bis(2-Chloroethyl)ether	*]		! !		*		
[2-Ch]oropheno]	*			ļ					
11 3-Dichlorobenzene	*		!	!	! !				
1.4-Dichlorobenzene	*	1	ļ		\		~_		
1,4-Dichlorobenzene	*				! !				
13_Mathrilahana]	*	.1	.	¦	}}		¦î		
2,2'-oxybis(1-Chloropropane)			·	¦	¦ Ⅰ	· <u> </u>	¦		
	- L		.]	1	¦'		·		
N-Nitroso-di-n-propylamine	*				¦	~~~~~	*		
Hexachloroethane	*	.!	.	·]			*		
			·	·	i		·		
Isonhorone	*	_ [·	. [· {	í í		í *		
2-Nitrophenol	*	- !			;		,*		
2,4-Dimethylphenol	*	-	.)	·	i i		i — *		
bis(2-Chloroethoxy)methane	*		·		· · · · · · · · · · · · · · · · · · ·		*		
2,4-Dichlorophenol	_*	-	· /	· · · · · · · · · · · · · · · · · · ·			*		
1,2,4-Trichioropenzene	-*	- ¦	• 1	1			*		
Naphthalene	_^	-	· '						
4-Chloroaniline	-		. /						
Hexachlorobutadiene		-]	1				I*		
4-Chloro-3-methylphenol		_			I	I	*		
2-Methylnaphthalene	-			1	I	l			
Hexachlorocyclopentadiene	_ *	-		1	I		*		
2,4,6-Trichlorophenol	*		1				·*		
2,4,5-Trichlorophenol	*				· · · · · · · · · · · · · · · · · · ·	!	*		
2-Chloronaphthalene	- 1		1						
2-Nitroaniline	-			. !			!!		
Dimethylphthalate	*					!	!*		
Acenaphthylene	*					!	!*		
3-Nitroaniline	1	_	. !			¦			
Acenaphthene	*			·		¦	¦"		
2,4-Dinitrophenol			.	.		¦			
4-Nitrophenol	1		.	·	·	¦	¦!		
Dibenzofuran	*		•	•		¦	¦"		
2,4-Dinitrotoluene	*	_ !	·			¦	"		
	1	i		I	I	· ·	Ii		

* Compounds with required minimum RRF and maximum %RSD values. All other compounds must meet a minimum RRF of 0.010.

6C

SEMIVOLATILE ORGANICS INITIAL CALIBRATION DATA

ab Name:		Contra	ct:					
ab Code: Case No.:		SAS N	o.:	SI	DG No.:		-	
nstrument ID: Ca	alibrati	on Date	(s):				-	
Ca	Calibration Times:							
LAB FILE ID: RRF20			RRF50			,	ſ	
RRF80 = RRF120			= RRF16					
						!		
COMPOUND		 RRF50		 RRF120	RRF160	RRF (% RSD	
			1	İ				
Diethylphthalate 4-Chlorophenyl-phenylether Fluorene	*							
	*	1			i ——— i			
4-Nitroaniline		- i	1		i			
4,6-Dinitro-2-methylphenol_	-	-	· /		ii			
N-Nitrosodiphenylamine (1)	- ;		1		i			
4-Bromophenyl-phenylether	*	- '						
Hexachlorobenzene	*		· ·		i — i			
Pentachlorophenol	- *	-	·		i i		,	
Phenanthrene	*	-	· ;	1			1	
Anthracene	-*							
Carbazole	- , <u></u>	- ¦	-	·	i i		[
Di-n-butylphthalate	- ¦		- ¦	·	i		i	
Fluoranthene	_1 *	···	-	- <u>`</u>				
Pyrene	-*		-	• i	i I		¦	
Butylbenzylphthalate			- '					
3,3'-Dichlorobenzidine			-	· .	i i		¦	
Benzo(a) anthracene	! *		-	- /	· []		¦	
Chrysene	-*		-		· · · · · · · · · · · · · · · · · · ·		¦	
bis(2-Ethylhexyl)phthalate			- '	• [·		¦	
				• ¦	·		¦	
Di-n-octylphthalate Benzo(b)fluoranthene		- ¦	- !		·		¦	
Benzo(k) fluoranthene			•	- ¦	·		¦	
Benzo(a) pyrene		- !	- ¦		·			
Indeno(1,2,3-cd) pyrene			- '	- ¦	·		{	
Dibenz(a, h) anthracene	^**	_		-			¦	
Benzo(g,h,i)perylene			-	-	·]]		¦	
Nitrobenzene-d5	1	1	1	1	1		1	
2-Fluorobiphenyl			- ¦	• ¦	·		¦	
Terphenyl-dl4	-*	- ¦	- i	- ¦	¦		¦	
Phenol-d5	*		- ¦	•¦	¦}		¦	
2-Fluorophenol			- ¦	-	·			
2,4,6-Tribromophenol		-	-	- ¦	¦}			
2-Chlorophenol-d4		- ¦	-	- ¦	·			
1,2-Dichlorobenzene-d4		-	-		¦			
1,2-DICHIOLOBENZENE-04			- ¦		¦			
	_ 1		<u> </u>	. 1	· /		1	

6D

- .

PESTICIDE INITIAL CALIBRATION OF SINGLE COMPONENT ANALYTES

Lab Name:	Contract:					
Lab Code:	Case No.:	SAS No.:	SDG No.:			
Instrument ID:	Level (X	(low): low mid	high			
GC Column:	ID:(mm)	Date(s) Analyzed:				

	RTC	F STANE	DARDS	MEAN	MEAN RT WINDO	
COMPOUND	LOW	MID	HIGH	RT	FROM	TO
alpha-BHC	======	; ;				
beta-BHC	}	• }	•]	·	· [
delta-BHC	¦	·╎·		 	·	
gamma-BHC (Lindane)	 	• {	• 1			
Heptachlor	¦	• {	·	¦		
Aldrin		·	·	¦		
Heptachlor epoxide		·	¦			
Endosulfan I		·				
Dieldrin			{	;		
4,4'-DDE		¦			·	
Endrin			·			i
Endosulfan II						i
4,4'-DDD		1				1
Endosulfan sulfate						
4,4'-DDT						
Methoxychlor		1				
Endrin ketone		I		1	1	1
Endrin aldehyde		· i		!		1
alpha-Chlordane			I		!	I
gamma-Chlordane				!		!

Tetrachloro-m-xylene			!		!	!
Decachlorobiphenyl			!			!
			1		ł.	

* Surrogate retention times are measured from Standard Mix A analyses.

Retention time windows are \pm 0.04 minutes for all compounds except Methoxychlor, at \pm 0.05 minutes.

6E

PESTICIDE INITIAL CALIBRATION OF SINGLE COMPONENT ANALYTES

Lab	Name:	Contract:					
Lab	Code:	Case	No.:	SAS No.:		SDG No.:	
Ins	trument I	D:	Level (x	low): low	mid	high	
GC	Column: _	ID:	(mm)	Date(s)	Analyzed:		

	CALIBRATION FACTORS									
COMPOUND	LOW	MID	HIGH	MEAN	*RSD					
alpha-BHC										
beta-BHC										
delta-BHC										
gamma-BHC (Lindane)										
Heptachlor										
Aldrin										
Heptachlor epoxide										
Endosulfan I										
Dieldrin										
4,4'-DDE										
Endrin										
Endosulfan II]	·					
4,4'-DDD										
Endosulfan sulfate				I						
4,4'-DDT				I						
Methoxychlor										
Endrin ketone					l					
Endrin aldehyde					I					
alpha-Chlordane					I					
gamma-Chlordane										

Tetrachloro-m-xylene					<u> </u>					
Decachlorobiphenyl										
II					l					

* Surrogate calibration factors are measured from Standard Mix A analyses.

%RSD must be less than or equal 10.0 % for all compounds except alpha-BHC beta-BHC, gamma-BHC, DDT, Endrin, and Methoxyclor, where **%RSD** must be less than or equal to 15.0%

6F PESTICIDE INITIAL CALIBRATION OF MULTICOMPONENT ANALYTES

Lab	Name:		Contract:					
Lab	Code: _		Case No.:		SAS No	o.:	_ SDG No.	;
Inst	rument	ID: _			Date(s)	Analyzed:		

GC Column: _____ ID: ____(mm)

	AMOUNT				VINDOW CALIBRATIO	
COMPOUND	(ng)	PEAK		FROM	TO	FACTOR

Toxaphene		*1				
		*2				
		*3				
		4				
		5				
Aroclor 1016		*1				
1		*2 *3				· · · · · · · · · · · · · · · · · · ·
		4				
		5				· /
Aroclor 1221		*1		·		·]
		*4				
		4	· · · · · · · · · · · · · · · · · · ·			·
	Į	5	·	·		
				·		·
Aroclor 1232		*2		¦ '		
		*3	ـــــــــــــــــــــــــــــــــــــ			
	{	4)	·		1
		5	·			· .
	·		·	·		
Aroclor 1242	!	×2				
1	ļ	*3				
		4		1		
	(5				
l	·	*1	·			
Aroclor 1248	{	*2	1	1		1
	1	*3	1			
	1	4	1	1		1
	1	i 5			l	1
	{	*1			I	<u> </u>
Aroclor 1254	l	*2		1	l	l
) 	*3			í	
	{	4	1	l		
	1	5_				
Aroclor 1260		*1		1		
A A A A A A A A A A A A A A A A A A A		*2	Í	·	·	
		*3]]		
	1	4	l			
	1	5				
		1				

* Denotes required peaks

6G PESTICIDE ANALYTE RESOLUTION SUMMARY

				ntract:	Co	ame:	Lab Name:_
EPA Sample No. (Standard 1): Lab Sample ID (1): Date Analyzed (1): Time Analyzed (1): ANALYTE RT (%) 01 02 03 04 05 06 07 08 09 EPA Sample No. (Standard 2): Lab Sample ID (2):		No.:	SDG	AS No.: _	Case No.: S	ode: Case	Lab Code:
ANALYTE RT (%) 01			ent ID (1):	Instrume	ID:(mm)	lumn (1):	GC Column
ANALYTE RT (%) 01	_		ple ID (1): .	Lab Samp	dard 1):	ample No. (Standard	EPA Sample
ANALYTE RT (%) 01			alyzed (1):	Time Ana		Analyzed (1):	Date Analy
01			(%)	RT			
EPA Sample No. (Standard 2): Lab Sample ID (2):						02 03 04 05 06 07 08	
Date Analyzed (2): Time Analyzed (2):			ple ID (2):	Lab Samp	ndard 2):	Sample No. (Standar	EPA Sample
ANALYTE RT (%)			RESOLUTION (%)	 RT	(TE	ANALYTE	Date Analy
01						01 02 03 04 05 05 06 07 08	

Resolution of two adjacent peaks must be calculated as a percentage of the height of the smaller peak, and must be greater than or equal to 60.0%.

7A VOLATILE CONTINUING CALIBRATION CHECK

Lab Name:	Contract:	
Lab Code:	Case No.: SAS No.:	SDG No.:
Instrument ID:	Calibration Date:	Time:
Lab File ID:	Init. Calib. Date(s):	
Heated Purge: (Y/N)	Init. Calib. Times:	
GC Column:	ID:(mm)	

COMPOUND	RRF	RRF50	MIN RRF	 %D	MAX %D
Chloromethane	******		=====	======	
Bromomethane			0.100	¦	25.0
Vinyl Chloride			0.100		25.0
Chloroethane			1001200	·	123.0
Methylene Chloride			1 ·	/	
Acetone			1	¦	
Carbon Disulfide			1		1
1,1-Dichloroethene	í		0.100		125 0
1,1-Dichloroethane			0.200		25.0
1,2-Dichloroethene (total)	/		00200		25.0
Chloroform			0.200		125 01
1,2-Dichloroethane		And in case of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the local division of the loc	0.100		25.0
2-Butanone		ا محمد محمد معمد ا ا	0.100		23.0
1,1,1-Trichloroethane	¦ ·		0.100		25.0
Carbon Tetrachloride	<u>}</u> -		0.100		25.0
Bromodichloromethane	{ -		0.200		25.0
1,2-Dichloropropane	{ -	/	0.200		25.0
cis-1,3-Dichloropropene	¦ -	i	0.2001	······	25 01
Trichloroethene	/ /		0.300		25.0
Dibromochloromethane			0.100		25.0
1,1,2-Trichloroethane	; _		0.100	{	25.0
Benzene	/	and the second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second se	0.5001		25.01
trans-1,3-Dichloropropene		and the second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second second se	0.100		25.01
Bromoform			0.100		25.0
4-Methy1-2-Pentanone	;			! `	2.01
2-Hexanone	·	/	-		
Tetrachloroethene	i	ic	.2001	; ·	25.0
1,1,2,2-Tetrachloroethane	;		.500		25.0
Toluene	i		.4001	The state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the state of the s	25.0
Chlorobenzene			.500		25.0
Ethylbenzene	i	io			
Styrene			.3001		5.01
Xylene (total)			.300j ⁻		5.0
Toluene-d8	İ	i	. 1		
Bromofluorobenzene	j	io	.200		5.0
1,2-Dichloroethane-d4	i			~	
All other compounds must meet	a minim			i	i

All other compounds must meet a minimum RRF of 0.010.

7B SEMIVOLATILE CONTINUING CALIBRATION CHECK

Lab	Name:		_ Contract:		
Lab	Code:	Case No.:	SAS No.:	SDG No.:	
Inst	trument ID:	Calib	ration Date:	Time:	
Lab	File ID:	Init.	Calib. Date(s):		
		Init.	Calib. Times:		

] .	1	MIN		MAXI
COMPOUND	RRF	RRF50	RRF	%D	% D

Phenol	İ	I	0.800		25.0
bis(2-Chloroethyl)ether			0.700		25.0
2-Chlorophenol			0.800		25.0
1,3-Dichlorobenzene			0.600		25.0
1,4-Dichlorobenzene			0.500		25.0
1,2-Dichlorobenzene			0.400		25.0
2-Methylphenol	1		0.700		25.0
2,2'-oxybis(1-Chloropropane)	1		l		ÍÍ
4-Methylphenol		1	0.600		25.0
N-Nitroso-di-n-propylamine			0.500		25.0
Hexachloroethane			0.300	1	25.0
Nitrobenzene		1	0.200	1	25.0
Isophorone	i		0.400		25.0
2-Nitrophenol	i		0:100	1	25.0
2,4-Dimethylphenol	1	1	0.200		25.0
bis(2-Chloroethoxy)methane	i		0.300		25.0
2,4-Dichlorophenol			0.200	1	25.0
1,2,4-Trichlorobenzene	1	1	0.200		25.0
Naphthalene		i	0.700	1	25.0
4-Chloroaniline	i	i	ĺ	1	Ì
Hexachlorobutadiene	· i	1	ĺ	1	i i
4-Chloro-3-methylphenol			j0.200		25.0
2-Methylnaphthalene		·	0.400		25.0
Hexachlorocyclopentadiene		·	'i		1
2,4,6-Trichlorophenol	1		0.200		25.0
2,4,5-Trichlorophenol	1		10.800		25.0
2-Chloronaphthalene	· ·		10.900		25.0
2-Nitroaniline	·			1	1
Dimethylphthalate	·	·		1	i i
Acenaphthylene	·		1.400	1	25.0
2,6-Dinitrotoluene	- ¦	·	0.200	1	25.0
3-Nitroaniline	1	·	'i	1	
Acenaphthene	· ;		0.900		25.0
2,4-Dinitrophenol	·			i	1
4-Nitrophenol		· •		i	i
Dibenzofuran		;	0.300	i	25.0
2,4-Dinitrotoluene	·	· [0.200		25.0
	· [·	1	¦	1
	· •	· ·	· ·	·	·

All other compounds must meet a minimum RRF of 0.010.

7C

SEMIVOLATILE CONTINUING CALIBRATION CHECK

Lab Name:	Contract:
Lab Code:	Case No.: SAS No.: SDG No.:
Instrument ID:	Calibration Date: Time:
Lab File ID:	Init. Calib. Date(s):
	Init. Calib. Times:

		 RRF50	MIN RRF	۶D	MA: %D
COMPOUND	RRF	1			-
Niethylphthalate		1			
-Chlorophenyl-phenylether				And the Owner of Concession of Concession of Concession, Name	25.
luorene		1	1.100		25.
-Nitroaniline			ļ (l 1
,6-Dinitro-2-methylphenol	1				
-Nitrosodiphenylamine (1)	1				25.
-Bromophenyl-phenylether	1		0.100		25.
Iexachlorobenzene	1		0.100		25.
Pentachlorophenol			0.100		25.
Phenanthrene	1		0.700		25.
nthracene	I				23.
Carbazole			/ 1 1		}
Di-n-butylphthalate			0.600		25.
luoranthene	I				25.
					1
Butylbenzylphthalate	ļ	.	, , , , , , , , , , , , , , , , , , ,		1
3,3'-Dichlorobenzidine			0.900		25.
Senzo(a) anthracene			0.900		25.
hrvene		.]			
is(2-Ethylhexyl)phthalate	!				ĺ
)i-n-octylphthalate	!	.!	0.700		25.
Senzo(b)fluoranthene	[.	0.700		25.
Senzo(k)fluoranthene	!	.	0.600		25.
Benzo(a) pyrene			10.400		25.
Indeno(1,2,3-cd)pyrene			10 (01)		143.
Dibenz(a,h)anthracene		-!	0.200		25.
Dibenz(a,h)anthracene Benzo(g,h,i)perylene			======		====
	 				100
Nitrobenzene-d5 2-Fluorobipheny1	1		0.700		25.
e-tractopthenit	1	1	0.500		25.
Terphenyl-d14	1	1	0.800		25.
Phenol-d5 2-Fluorophenol]	. !	0.600		25.
2,4,6-Tribromophenol					25.
2-Chlorophenol-d4			0.800		25.
1,2-Dichlorobenzene-d4			0.400		143.
			I		ا
l) Cannot be separated from All other compounds must me	Dipheny	lamine	· .		

7D PESTICIDE CALIBRATION VERIFICATION SUMMARY

Lab	Name:		Contract:	
Lab	Code:	Case No.:	SAS No.: SDG	No.:
GC	Column:	ID:(mm)	Init. Calib. Date(s):	
EPA	Sample	No.(PIBLK):	Date Analyzed	•
Lab	Sample	ID (PIBLK):	Time Analyzed	•
EPA	Sample	No.(PEM):	Date Analyzed	:
Lab	Sample	ID (PEM):	Time Analyzed	ŧ

PEM COMPOUND	RT	RT WI	INDOW TO	CALC AMOUNT	NOM AMOUNT	RPD
				(ng)	(ng)	
alpha-BHC						İİ
gamma-BHC (Lindane)				- <u></u>		
Endrin						
Methoxychlor						

4,4'-DDT % breakdown (1): _____ Endrin % breakdown (1): _____ Combined % breakdown (1):

QC LIMITS:

RPD of amounts in PEM must be less than or equal to 25.0% 4,4'-DDT breakdown must be less than or equal to 20.0% Endrin breakdown must be less than or equal to 20.0% Combined breakdown must be less than or equal to 30.0%

7E PESTICIDE CALIBRATION VERIFICATION SUMMARY

Lab Name:		Contr	act:			
Lab Code: Case No.: _						
GC Column: ID:	(mm)	Init.	Calib. D	ate(s):		
EPA Sample No.(PIBLK):				Analyzed		
Lab Sample ID (PIBLK):			Time	Analyzed	:	
			Date	Analyzed	:	
EPA Sample No.(INDA):			M ima	Analyzed		
Lab Sample ID (INDA):						
INDIVIDUAL MIX A COMPOUND	RT	RT W FROM		(ng)	NOM AMOUNT (ng)	RPD
alpha-BHC						
Heptachlor						
Heptachlor			.			
Dielarin			•	-		
Endrin			-			
4,4'-000				· · · ·		
4,4'-DDT						
Methoxychlor						
Tetrachloro-m-xylene Decachlorobiphenyl						
EPA Sample No.(INDB):				Analyzed		
Lab Sample ID (INDB):				Analyzed		
INDIVIDUAL MIX B			INDOW	CALC	NOM	
COMPOUND	RT	FROM	TO	AMOUNT	AMOUNT	RPD
				(ng)	(ng)	

beta-BHC			-]
delta-BHC			-			
Aldrin			-			
Heptachlor epoxide			-			
4,4'-DDE			-			
Endosulfan II						
Endosulfan sulfate						
Endrin ketone						
Endrin aldehyde						
alpha-Chlordane						
Tetrachloro-m-xylene			_			
Decachlorobiphenyl			-			
Secaontor os thread a			- ا			

QC LIMITS: RPD of amounts in the Individual Mixes must be less than or equal to 25.0%.

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8A VOLATILE INTERNAL STANDARD AREA AND RT SUMMARY

Lab	Name:	Contract:				
Lab	Code: Case No.:	SAS No.: SDG No.:				
Lab	File ID (Standard):	Date Analyzed:				
Inst	trument ID:	Time Analyzed:				
GC	Column: ID:(mm)	Heated Purge: (Y/N)				

	IS1(BCM) AREA #	RT #	IS2(DFB) AREA #	RT #	IS3(CBZ) AREA #	RT
12 HOUR STD	مالى دىر الجني موجود بور مو					
UPPER LIMIT						
LOWER LIMIT						
EPA SAMPLE						
NO.						

	· · · ·					
	· · · · · · · · · · · · · · · · · · ·					
	·····					
	······································					
1			1			

IS1 (BCM) = Bromochloromethane IS2 (DFB) = 1,4-Difluorobenzene IS3 (CBZ) = Chlorobenzene-d5

AREA UPPER LIMIT = +100% of internal standard area AREA LOWER LIMIT = - 50% of internal standard area RT UPPER LIMIT = +0.50 minutes of internal standard RT RT LOWER LIMIT = -0.50 minutes of internal standard RT

Column used to flag values outside QC limits with an asterisk. * Values outside of QC limits.

8B SEMIVOLATILE INTERNAL STANDARD AREA AND RT SUMMARY

Lab	Name:	Contract:
Lab	Code: Case No.:	SAS No.: SDG No.:
Lab	File ID (Standard):	Date Analyzed:
	trument ID:	Time Analyzed:

·		S1 (DCB)		IS2(NPT)		IS3 (ANT)	
		AREA	# RT #	AREA #	RT #	AREA #	RT #
12 HOU							
UPPER			-				
EPA SA							
NO		للتجريف الناري والمتحد					
01							
02			_				
03			-[
04			-				
06							
07							
08							
09 10			-				
11							
12							
13 <u>·</u>			-				
15					·		
16							
17							
18 19							
20							
21			-				
22			_]		·	·	
IS1 (DCB)	= 1 4-[hichloro	benzene-d	4			
TC2 (NDT)	= Nanht	halene-	uð				
IS3 (ANT)	= Acena	aphthene	-010				
	יים דערדיד	= +1.00%	of inter	nal standar nal standar	d area		
AREA UPPE	R LIMIT	= - 50%	of inter	nal standar internal s	a area Fandard 1	איי	
RT UPPER	LIMIT =	+0.50 .	inutes of	internal s internal s	tandard I	RT	
RT LOWER	I.TMIT =	-0.50 #	THUCCCO ()				
# Column	need to	flag in	ternal st	andard area	values v	with an ast	erisk.
* Values	outside	of QC 1	imits.				
page of	-						

3/90

8C

SEMIVOLATILE INTERNAL STANDARD AREA AND RT SUMMARY

Lab	Name:		Contract:	Contract:				
Lab	Code:	Case No.:	SAS No.:	SDG No.:				
Lab	File ID (Standa:	rd):	Date A	inalyzed:				
Inst	trument ID:		Time A	nalyzed:				

	IS4(PHN)		IS5(CRY)		IS6(PRY)	
i	AREA #	RT #		RT #	AREA #	RT #

12 HOUR STD					1	I
UPPER LIMIT						
LOWER LIMIT						

EPA SAMPLE	Ì		1		1	
NO.			I			
01		l	l			
02			l			<u> </u>
03			1			
04	l		l			
05						!!
061					·	!!
07		ļ				
08						!]
09						!!
10	ļ			·		! !
11				¦		!!
12	!	·		¦	!	!!
13	·		l]
14	·	ļ				¦
16	·	·			1	
17	l			¦]¦
18	 			 	1] [
19	 	¦	l	1		
20		1 1				
20		۱ ۱		1		¦
22		 	1		 	
44	I	I	I	۱	I	۱ <u> </u>
IS4 (PHN) = Pho	enanthrene-	016				
IS5 (CRY) = Ch		420				
	rylene-d12					
AREA UPPER LIM	IT = +100%	of inter	nal standar	d area		
AREA LOWER LIM						
RT UPPER LIMIT					RT	
RT LOWER LIMIT						

Column used to flag internal standard area values with an asterisk.
* Values outside of QC limits.

page __ of __

8D PESTICIDE ANALYTICAL SEQUENCE

Lab	Name:	Contract:
Lab	Code:	Case No.: SAS No.: SDG No.:
GC	Column:	ID:(mm) Init. Calib. Date(s):

Instrument ID:

THE ANALYTICAL SEQUENCE OF PERFORMANCE EVALUATION MIXTURES, BLANKS, SAMPLES, AND STANDARDS IS GIVEN BELOW:

EPA	LAB	DATE	TIME ANALYZED	TCX RT #	DCB
SAMPLE NO.	SAMPLE ID	ANALYZED		nt T Rasses	
SAMPLE NO.			i		1
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					1
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Contraction of the second second second second second second second second second second second second second s					
Construction of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of the Party of th		l			·]
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					}
		QC LIM	IITS		
	aro-m-vvlene	(± 0.04 M	IINUTES)		
CCX = Tetrachl CB = Decachlo	unchinhenv]	(± 0.04 M	IINUTES)		

* Values outside of QC limits.

9A PESTICIDE FLORISIL CARTRIDGE CHECK

Lab Name:		Contract:		
Lab Code:	Case No.:	SAS No.:	SDG No.:	
Florisil Cartridge	Lot Number:	Date of Analy	/sis:	
GC Column(1):	ID:(m	m) GC Column(2): _	<u></u>	ID:(mm)

COMPOUND	SPIKE ADDED (ng)	SPIKE RECOVERED (ng)	* REC	 #	QC LIMITS
alpha-BHC gamma-BHC (Lindane) Heptachlor Endosulfan I Dieldrin					80-110 80-110 80-110 80-110 80-110
Endrin 4,4'-DDD 4,4'-DDT Methoxychlor			 		80-110 80-110 80-110 80-110
Tetrachloro-m-xylene Decachlorobiphenyl	 			 	80-110 80-110

Column to be used to flag recovery with an asterisk.

* Values outside of QC limits.

THIS CARTRIDGE LOT APPLIES TO THE FOLLOWING SAMPLES, BLANKS, MS, AND MSD:

1	EPA		LAB		DATE		DATE	
i	SAMPLE	NO.	SAMPLE	ID	ANALYZED	1	ANALYZED	2
i				*===	=========	= 3		==
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02								
03								
04								
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06					1		l	
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17			·				l	
18		1						
19	- <u></u>							
20							l	
21								
22								
23								

9B PESTICIDE GPC CALIBRATION

Lab	Name:			(Contract:			
Lab	Code:	Case	No.:		SAS No.:	SDG No.:	·····	
GPC	Column:				Calibration Date:		-	
GC (Column(1):		ID:	(mm)	GC Column(2):		ID:	_(mm)

COMPOUND	SPIKE ADDED (ng)	SPIKE RECOVERED (ng)	REC #	QC. LIMITS REC.
gamma-BHC (Lindane) Heptachlor Aldrin Dieldrin Endrin 4,4'-DDT				80-110 80-110 80-110 80-110 80-110 80-110

Column to be used to flag recovery values with an asterisk
* Values outside of QC limits

THIS GPC CALIBRATION APPLIES TO THE FOLLOWING SAMPLES, BLANKS, MS AND MSD:

EPA	I LAB		DATE		DATE	
SAMPLE NO	4.	ID	ANALYZED	1	ANALYZED	2
SAMPLE NO						
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021			1		·	i
041			·			¦
051			/	i	·	¦
061			1			¦
071	/			j		¦
07			۱	¦		{
08			!	-;		¦
09			·			
10			/	-1		¦
11/	/		I	-¦		¦
12			·	-¦		
13	/		 	-¦		-¦
14		I		-;		¦
15		······		-¦		-¦
16/	_!	{				
17	/	······································		-¦		
18		¦		-1		
19		I				
20	_ !	{		-¦·		-!
21/	_ [-¦·		
22				-¦-		-!
23		}		-¦-		-!
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25		·		-¦-		-!
26		······································		_!_		_1

page __ of ___

PESTICIDE For singi					EPA SAM	PLE NO.
Lab Name:		Co	ntract:			
Lab Code: Case No.	SDG No.:	<u> </u>				
Lab Sample ID :	-		Date(s)) Analy	zed:	
Instrument ID (1):			Instru	ment ID	(2):	
GC Column(1): ID:		_(mm)	GC Col	umn(2):	ID:	(mm)
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ANALYTE 	COL		FROM		CONCENTRATION	%D
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page ____ of ____

10B Pesticide identi For multicompo	IFICATION SUMMARY ONENT ANALYTES	o. — _!
Lab Name:	Contract:	
Lab Code: Case No.:	SAS No.: SDG No.:	
Lab Sample ID :	Date(s) Analyzed:	
Instrument ID (1):	Instrument ID (2):	
GC Column(1): ID:	(mm) GC Column(2): ID:	(11111)

			RTW	INDOW		MEAN	
ANALYTE	PEAK	RT	FROM	TO		CONCENTRATION	*D
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At least 3 peaks are required for identification of multicomponent analytes page __ of ___

		SAMPLE L	OG-IN SHEET		
Lab Name:			·		Page of
Received By (Print Nam				Log-in Date:	
Received By (Signature)	:			<u> </u>	
Case Number:			CORRE	SPONDING	
Sample Delivery Group No.:		EPA Sample	SAMPLE TAG	ASSIGNED LAB	REMARKS: CONDITION OF SAMPLE
REMARKS:		*			SHIPMENT, ETC.
1. Custody Seal(s)	Present/Absent* Intect/Broken				
2. Custody Seal Nos.:					
3. Chaim-of-Custody Records	Present/Absent*				
4. Traffic Reports or Packing List	Present/Absont"				
5. Airbill	Airbill/Sticker Present/Absent*				
6. Airbill No.:					
7. Sample Tags	Present/Absont*			<u></u>	
Sampia Tag Numbers	Listed/Not Listed on Chain-of-				
8. Sample Condition:	Custody				
s. Sampa Colousou:	Leaking				
9. Does information on					
custody records, traff reports, and sample					
tags agros?	Yes/No*				
10. Date Received at Lab					
11. Time Received:			·		
Sample 1	'ransfer		ļ		
Fraction:			ļ		
Area #:					
By:					
On:					

* Contact SMO and attach record of resolution

Logbook No.: _____ Logbook Page No: __

Reviewed By: Date: ___

FORM DC-1

ORGANICS COMPLETE SDG FILE (CSF) INVENTORY SHEET

LABORATORY NAME		CITY/STATE				
		SOG NOS. TO FOLLOW				
CONTRACT NO.			10			
All docume where poss	nts delivered ible. (REFEREN	in the complete SDG file m ICE EXHIBIT B, SECTION II and	nust be or: SECTION II	iginal [.)	documer	lts
			PAGE	NOs	CH LAB	
Inventory	Sheet (Form DC-	2) (Do not number)				
SDG Case N	arrative					
Traffic Ret						
Volatiles						
0.00						
S	Devent Pa	covery Summary (Form II VOA)				
Lab Co	ontrol Samole R	ecovery (From 111 Vor)				
Method	i Blank Summary	(Form IV VUA)				
Tuning	g and Mass Cali	bration (Form V VOA)			·	
b. Sample	Data					~ <u>-</u> -
Tentra	esults - (Form tively Identifi	I VOA) ed Compounds (Form I VOA-TIC) ion chromatograms (RIC)		·		
for	each sample	Tou curoactore				
Raw		ckground-subtracted target compounds identified				
Masi	s spectra of al est library mat	1 reported flos with these			·	<u>-</u> -
c. Standa	ds Data (All In	nstruments)				
T	al Calibration	Data (FOTH VI VUR)		~~~~		
PTCe .	and Oven Report	s for all scandards			····	
Conti	oving Calibrati	on (FOLM VIL VOR)				
DT.C	and Owen Banart	e for all Scallaros				
Inter	nal Standard Ar	ea and RT Summary	· · · · · · · · · · · · · · · · · · ·			
(Fo:	rm VIII VOA)					
d. QC Data	L Constant					
BFB	_					
Blank						
Matri	x Spike Data	ta Data			·	
Matri	x Spike Duplica					
<u>Semivolati</u>						
a. QC Summ	pary	Summary (Form II SV)				
Surro	gate Percent Re	covery Summary (Form II SV)				
MC /MC	D Summary (FOID					
Metho	i Blank Summary	bration (Form V SV)	····			
Tuning	g and Mass Call	DIALLON (*******				······································

CASE NO.	SÓG NO	SDG NOS. TO FOLLOW	SAS N)	
			PAGE NOS	Cl	HECK
			FROM TO	LAB	EPA
Semi	ivolatiles Data (con	nt.)			
Ъ.	Sample Data				······
	TCL Results (Form	I SV)			
	Tentatively Ident:	ified Compounds (Form I SV-TIC	3)		
	Reconstructed tota	al ion chromatograms (RIC)			
	for each sample				
	For each sample:				
	Raw spectra and	background-subtracted			
	mass spectra (of target compounds			
		TICs with three best library	matches		
	GPC chromatog	rams (if GPC performed)			
.c.	Standards Data (All				
		on Data (Form VI SV)			
		orts for all Standards			
		ation (Form VII SV)			
		orts for all Standards			
		Area and RT Summary			
	(Form VIIIB and	Form VIIIC SV)		<u> </u>	····
d.	QC Data	а. С			
	DFTPP				
	Blank Data				
	Matrix Spike Data				
	Matrix Spike Dupl	lcate Data		<u>-</u>	
Pes	ticides				
a.	QC Summary				
	Surrogate Percent	Recovery Summary (Form II PES	T)		
	MS/MSD Duplicate	Summary (Form III PEST)			
	method Blank Summ	ary (Form IV PEST)			
L	Comple Det-				
D.	Sample Data				
	(Form I PEST)	anic Analysis Data Sheet			
	Chromatograms (Pr	imarry Column)			
	Chromatograme fra	mary column) m second GC column confirmatio	-	- <u></u>	
	GC Integration re	port or data system printout a	ind		
	calibration plo				
	Manual work sheet				
	UV traces from GP	C (if available)			
	ror pesticides/Ar	oclors confirmed by GC/MS, cop	ies		
	or raw spectra	and copies of background-subtr et compounds (samples & standa	acted mass		

ORGANICS COMPLETE SDG FILE (CSF) INVENTORY SHEET (Cont.)

	CASE NO SDG NO SDG NOS. TO	FOLLOW	- <u> </u>		SAS NO.		
				PAGE FROM	NOs TO	CH LAB	IECK EPA
6.	Pesticides (cont.)						
	c. Standards Data Pesticides Evaluation Standards (Form VIII PEST-1)	Summary					
	Pesticides Evaluation Standards (Form VIII, PEST-2)	Summary					
	Pesticide/Aroclor Standards Sur Pesticide/Aroclor Identification Standard chromatograms and data for all Standards	on (Form X PI	LST)				
	For pesticides/Aroclors confirm of spectra for standards used		, c opies				
	d. QC Data Blank Data Matrix Spike Data				<u></u>		
	Matrix Spike Duplicate Data						
7	<u>Miscellaneous Data</u>						
	Original preparation and analysis preparation and analysis logbook Internal sample and sample extract	pages	ies of				
	chain-of-custody records Screening records						
	All instrument output, including s from screening activities (descr						
		• •					
8	. EPA Shipping/Receiving Documents	<u>,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,</u>					
	Airbills (No. of shipments)						
	Chain-of-Custody Records						
	Sample Tags Sample Log-In Sheet (Lab & DC1)						
	SDG Cover Sheet						
	Miscellaneous Shipping/Receiving R (describe or list)	ecords					
9	. <u>Internal Lab Sample Transfer Record</u> (describe or list)	s and Tracki	ng Sheets				

ORGANICS COMPLETE SDG FILE (CSF) INVENTORY SHEET (Cont.)

CASE NO.	SDG NO	SDG NOS. TO FOLLOW		SAS NO	•	
				PAGE NOS	CHI	
				FROM TO	LAB	EPA
<u>Other Reco</u>	<u>rds</u> (describe	or list)				
Telephone	Communication	Log				
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<u>Comments:</u>				·····		-
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pleted by:						
(CLP Lab)	(Signat	lre)	(Printed	Name/Title)		(Date)
dited by:						

ORGANICS COMPLETE SDG FILE (CSF) INVENTORY SHEET (Cont.)

EXHIBIT C

TARGET COMPOUND LIST (TCL) AND CONTRACT REQUIRED QUANTITATION LIMITS (CRQL)

NOTE: The values in these tables are quantitation limits, not absolute detection limits. The amount of material necessary to produce a detector response that can be identified and reliably quantified is greater than that needed to simply be detected above the background noise. The quantitation limits in these tables are set at the concentrations in the sample equivalent to the concentration of the lowest calibration standard analyzed for each analyte.

Specific quantitation limits are highly matrix dependent. The quantitation limits listed herein are provided for guidance and may not always be achievable.

The CRQL values listed on the following pages are based on the analysis of samples according the specifications given in Exhibit D. For each fraction and matrix, a brief synopsis of the sampling handling and analysis steps is given, along with an example calculation for the CRQL value. All CRQL values are rounded to two significant figures. For soil samples, the moisture content of the samples is not considered in these example calculations.

OLM01.0

	COMPOUND	LIST	(TCL)	AND	CONTRACT	REQUIRED	QUANTITATION	LIMITS	(CROL)
TARGET	CONLOOND		(102)		OOUTING	KEGO LEED	QUART THE TON	14TLTT 1.3	(UNQL)

		Quanti	tation I	Limits*	
			Low	Med.	On
		Water	Soil	Soil	Column
Volatiles	CAS Number	ug/L		ug/Kg	(ng)
V014C1400					
1. Chloromethane	74-87-3	10	10	1200	(50)
2. Bromomethane	74-83-9	10	10	1200	(50)
3. Vinyl Chloride	75-01-4	10	10	1200	(50)
4 Chloroethane	75-00-3	10	10	1200	(50)
5. Methylene Chloride	75-09-2	10	10	1200	(50)
-					
6. Acetone	67-64-1	10	10	1200	(50)
7. Carbon Disulfide	75-15-0	10	10	1200	(50)
8. 1,1-Dichloroethene	75-35-4	10	10	1200	(50)
9. 1,1-Dichloroethane	75-34-3	10	10	1200	(50)
10. 1,2-Dichloroethene (total)	540-59-0-	10	10	1200	(50)
44	67-66-3	10	10	1200	(50)
11. Chloroform	107-06-2	10	10	1200	(50)
12. 1,2-Dichloroethane	78-93-3	10	10	1200	(50)
13. 2-Butanone	71-55-6	10	10	1200	(50)
14. 1,1,1-Trichloroethane	56-23-5	10	10	1200	(50)
15. Carbon Tetrachloride	30-23-3	10	10	1200	(50)
16. Bromodichloromethane	75-27-4	10	10	1200	(50)
17. 1,2-Dichloropropane	78-87-5	10	10	1200	(50)
18. cis-1,3-Dichloropropene	10061-01-5	10	10	1200	(50)
19. Trichloroethene	79-01-6	10	10	1200	(50)
20. Dibromochloromethane	124-48-1	10	10	1200	(50)
					(/
21. 1,1,2-Trichloroethane	79-00-5	10	10	1200	(50)
22. Benzene	71-43-2	10	10	1200	(50)
23. trans-1,3-Dichloropropene		10	10	1200	(50)
24. Bromoform	75-25-2	10	10	1200	(50)
25. 4-Methyl-2-pentanone	108-10-1	10	10	1200	(50)
		10			
26. 2-Hexanone	591-78-6	10	10	1200	(50)
27. Tetrachloroethene	127-18-4	10	10	1200	(50)
28. Toluene	108-88-3	10	10	1200	(50)
29. 1,1,2,2-Tetrachloroethane		10	10	1200	(50)
30. Chlorobenzene	108-90-7	10	10	1200	(50)
31. Ethyl Benzene	100-41-4	10	10	1200	(50)
32. Styrene	100-42-5	10	10	1200	(50)
33. Xylenes (Total)	1330-20-7	10	10	1200	(50)
infinite (local)	1990-20-1	74	10	1200	(50)

* Quantitation limits listed for soil/sediment are based on wet weight. The quantitation limits calculated by the laboratory for soil/sediment, calculated on dry weight basis as required by the contract, will be higher.

Note that the CRQL values listed on the preceding page may <u>not</u> be those specified in previous CLP Statements of Work. These values are set at concentrations in the sample equivalent to the concentration of the lowest calibration standard specified in Exhibit D VOA. Lower quantitation limits may be achievable for water samples by employing the Statement of Work for Low Concentration Water for Organic Analyses.

VOLATILES

Water Samples

A 5 mL volume of water is purged with an inert gas at ambient temperature. The volatiles are trapped on solid sorbents, and desorbed directly onto the GC/MS. For a sample with compound X at the CRQL of 10 ug/L:

(10 ug/L) (5 mL) $(10^{-3} \text{ L/mL}) = 50 \times 10^{-3} \text{ ug} = 50 \text{ ng on the GC column}$

Low Level Soil/Sediment Samples

A 5 g aliquot of the soil/sediment sample is added to a volume of water in a purge tube, heated, and purged with an inert gas. The volatiles are trapped, and later desorbed directly onto the GC/MS. For a sample with compound X at the CRQL of 10 ug/Kg:

(10 ug/Kg) (5 g) $(10^{-3} \text{ Kg/g}) = 50 \times 10^{-3} \text{ ug} = 50 \text{ ng on the GC column}$

Medium Level Soil/Sediment Samples

A 4 g aliquot of soil/sediment is extracted with 10 mL of methanol, and filtered through glass wool. Only 1 mL of the methanol extract is taken for screening and analysis. Based on the results of a GC/FID screen, an aliquot of the methanol extract is added to 5 mL of reagent water and purged at ambient temperature. The largest aliquot of extract considered in Exhibit D is 100 uL. For a sample with compound X at the CRQL of 1200 ug/Kg:

(1200 ug/Kg) (4 g) $(10^{-3} \text{ Kg/g}) = 4800 \text{ x} 10^{-3} \text{ ug} = 4800 \text{ ng}$

This material is contained in the 10 mL methanol extract:

(4800 ng)/ 10 mL = 480 ng/mL

Of which, 100 uL are purged from the reagent water.

(480 ng/mL) (100 uL) (10^{-3} mL/uL) = 480 x 10^{-1} ng = 50 ng on the GC column

Note that for both low and medium soil/sediment samples, while it may affect the purging efficiency, the volume of reagent water used in the purging process does not affect the calculations.

TARGET COMPOUND LIST (TCL) AND CONTRACT REQUIRED QUANTITATION LIMITS (CRQL)

		Quanti	tation		_
			Low	Med.	On
		<u>Water</u>	<u>Soil</u>	<u>Soil</u>	<u>Column</u>
<u>Semivolatiles</u>	CAS Number	ug/L	ug/Kg	ug/Kg	<u>(ng)</u>
34. Phenol	108-95-2	10	330	10000	(20)
	111-44-4	10	330	10000	(20)
35. bis(2-Chloroethyl) ether	95-57-8	10	330	10000	(20)
36. 2-Chlorophenol			330		(20)
37. 1,3-Dichlorobenzene	541-73-1	10		10000	(20)
38. 1,4-Dichlorobenzene	106-46-7	10	330	10000	(20)
39. 1,2-Dichlorobenzene	95-50-1	10	330	10000	(20)
40. 2-Methylphenol	95-48-7	10	330	10000	(20)
41. 2,2'-oxybis					(20)
(1-Chloropropane) [#]	108-60-1	10	330	10000	(20)
42. 4-Methylphenol	106-44-5	10	330	10000	(20)
43. N-Nitroso-di-n-	200 44 5				(20)
dipropylamine	621-64-7	10	330	10000	(20)
arbrob) ramino	021 04 /			20000	(20)
44. Hexachloroethane	67-72-1	10	330	10000	(20)
45. Nitrobenzene	98-95-3	10	330	10000	(20)
46. Isophorone	78-59-1	10	330	10000	(20)
47. 2-Nitrophenol	88-75-5	10	330	10000	(20)
48. 2,4-Dimethylphenol	105-67-9	10	330	10000	(20)
					()
49. bis(2-Chloroethoxy)					
methane	111-91-1	10	330	10000	(20)
50 2,4-Dichlorophenol	120-83-2	10	330	10000	(20)
51. 1,2,4-Trichlorobenzene	120-82-1	10	330	10000	(20)
52. Naphthalene	91-20-3	10	330	10000	(20)
53. 4-Chloroaniline	106-47-8	10	330	10000	(20)
54. Hexachlorobutadiene	87-68-3	10	330	10000	(20)
55. 4-Chloro-3-methylphenol	59-50-7	10	330	10000	(20)
56. 2-Methylnaphthalene	91-57-6	10	330	10000	(20)
57. Hexachlorocyclopentadiene	77-47-4	10	330	10000	(20)
58. 2,4,6-Trichlorophenol	88-06-2	10	330	10000	(20)
59. 2,4,5-Trichlorophenol	95-95-4	50	1700	50000	(100)
60. 2-Chloronaphthalene	91-58-7	10	330		(100)
61. 2-Nitroaniline	88-74-4	50	1700	10000	(20)
62. Dimethylphthalate	131-11-3	10	330	50000	(100)
	208-96-8	10		10000	(20)
63. Acenaphthylene	200-90-0	10	330	10000	(20)
64. 2,6-Dinitrotoluene	606-20-2	10	330	10000	(20)
65. 3-Nitroaniline	99-09-2	50	1700	50000	(100)
66. Acenaphthene	83-32-9	10	330	10000	(20)
67. 2,4-Dinitrophenol	51-28-5	50	1700	50000	(100)
68. 4-Nitrophenol	100-02-7	50	1700	50000	(100)
	T00 02 1	20	1,00	50000	(100)

Previously known by the name bis(2-Chloroisopropyl) ether

(continued)	<u>Quantitation Limits*</u> Low Med. On				
Semivolatiles	CAS Number	<u>Water</u> ug/L	<u>Soil</u> ug/Kg	<u>Soil</u> ug/Kg	<u>Column</u> (ng)
69. Dibenzofuran	132-64-9	10	330	10000	(20)
70. 2,4-Dinitrotoluene	121-14-2	10	330	10000	(20)
71. Diethylphthalate	84-66-2	10	330	10000	(20)
72. 4-Chlorophenyl-phenyl					
ether	7005-72-3	10	330	10000	(20)
73. Fluorene	86-73-7	10	330	10000	(20)
74. 4-Nitroaniline	100-01-6	50	1700	50000	(100)
75. 4,6-Dinitro-2-methylphenol		50	1700	50000	(100)
76. N-nitrosodiphenylamine	86-30-6	10	330	10000	(20)
77. 4-Bromophenyl-phenylether	101-55-3	10	330	10000	(20)
78. Hexachlorobenzene	118-74-1	10	330	10000	(20)
70 Bentachlanach I					(100)
79. Pentachlorophenol	87-86-5	50	1700	50000	(100)
80. Phenanthrene	85-01-8	10	330	10000	(20)
81. Anthracene	120-12-7	10	330	10000	(20)
82. Carbazole	86-74-8	10	330	10000	(20)
83. Di-n-butylphthalate	84-74-2	10	330	10000	(20)
84. Fluoranthene	206-44-0	10	330	10000	(20)
85. Pyrene	129-00-0	10	330	10000	(20)
86. Butylbenzylphthalate	85-68-7	10	330	10000	(20)
87. 3,3'-Dichlorobenzidine	91-94-1	10	330	10000	(20)
88. Benzo(a)anthracene	56-55-3	10	330	10000	(20)
• • • • • • • • • • • • • • • • • • • •					
89. Chrysene	218-01-9	10	330	10000	(20)
90. bis(2-Ethylhexyl)phthalate	117-81-7	10	330	10000	(20)
91. Di-n-octylphthalate	117-84-0	10	330	10000	(20)
92. Benzo(b)fluoranthene	205-99-2	10	330	10000	(20)
93. Benzo(k)fluoranthene	207-08-9	10	330	10000	(20)
					-
94. Benzo(a)pyrene	50-32-8	10	330	10000	(20)
95. Indeno(1,2,3-cd)pyrene	193-39-5	10	330	10000	(20)
96. Dibenz(a,h)anthracene	53-70-3	10	330	10000	(20)
97. Benzo(g,h,i)perylene	191-24-2	10	330	10000	(20)
-					

* Quantitation limits listed for soil/sediment are based on wet weight. The quantitation limits calculated by the laboratory for soil/sediment, calculated on dry weight basis as required by the contract, will be higher.

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SEMIVOLATILES

Water Samples

A 1 L volume of water is extracted three times with 100 mL of methylene chloride at a pH of approximately 2. This extract is reduced in volume to 1.0 mL, and a 2 uL volume is injected onto the GC/MS for analysis. For a sample with compound X at the CRQL of 10 ug/L:

(10 ug/L) (1 L) = 10 ug in the original extract

When the extract is concentrated, this material is contained in the 1 mL concentrated extract, of which 2 uL are injected into the instrument:

(10 ug/mL) (2 uL) $(10^{-3} \text{ mL/uL}) = 20 \text{ x} 10^{-3} \text{ ug} = 20 \text{ ng on the GC column}$

Low Soil Samples

A 30 g soil sample is extracted three times with methylene chloride at ambient pH, by sonication. The extract is reduced in volume to 1.0 mL, and a 2 uL volume is injected onto the GC/MS for analysis. For a sample with compound X at the CRQL of 330 ug/Kg:

(330 ug/Kg) (30 g) $(10^{-3} \text{ Kg/g}) = 9900 \times 10^{-3} \text{ ug} = 9.9 \text{ ug}$

When the sample extract is to be subjected to Gel Permeation Chromatography (<u>required</u>) to remove high molecular weight interferences, the volume of the extract is initially reduced to 10 mL. This 10 mL is put through the GPC column, and only 5 mL are collected off the GPC. That 5 mL volume is reduced to 0.5 mL prior to analysis. Therefore:

(9.9 ug/10 mL) (5 mL) = 4.95 ug

This material is contained in the 0.5 mL extract, of which 2 uL are injected into the instrument:

(4.95 ug/0.5 mL) (2 uL) (10⁻³ mL/uL) = 1.98 x 10⁻² ug \approx 20 ng on the GC column

Medium Soil Samples

A 1 g soil sample is extracted once with 10 mL of methylene chloride, which is filtered through glass wool to remove particles of soil. The filtered extract is then subjected to GPC clean up, and only 5 mL of extract are collected after GPC. This extract is reduced in volume to 0.5 mL, of which 2 uL are injected onto the GC/MS. For a sample with compound X at the CRQL of 10,000 ug/Kg:

(10,000 ug/Kg) (1g) $(10^{-3} \text{ Kg/g}) = 10 \text{ ug}$

(continued)

Semivolatiles, Medium Soil, continued -

This material is contained in the 10 mL extract, of which only 5 mL are collected after GPC:

(10 ug) (5 mL/10 mL) - 5 ug

The volume of this extract is reduced to 0.5 mL, of which 2 uL are injected into the instrument:

(5 ug/0.5 mL) (2 uL) $(10^{-3} \text{ mL/uL}) = 20 \times 10^{-3} \text{ ug} = 20 \text{ ng on the GC column}$

TARGET COMPOUND LIST (TCL) AND CONTRACT REQUIRED QUANTITATION LIMITS (CRQL)

			tation L	
		Water	<u>Soil</u>	<u>On Column</u>
Pesticides/Aroclors	CAS Number	ug/L	ug/Kg	<u>(pg)</u>
	319-84-6	0.05	1.7	5
98. alpha-BHC	319-84-6	0.05	1.7	5
99. beta-BHC	319-85-7	0.05	1.7	5
100. delta-BHC	58-89-9	0.05	1.7	5
101. gamma-BHC (Lindane)	76-44-8	0.05	1.7	5
102. Heptachlor	/0-44-0	0.05	1./	5
103. Aldrin	309-00-2	0.05	1.7	5
104. Heptachlor epoxide	1024-57-3	0.05		5
105. Endosulfan I	959-98-8	0.05	1.7	5
106. Dieldrin	60-57-1	0.10	3.3	10
107. 4,4'-DDE	72-55-9			10
2071 4,4 552			•	
108. Endrin	72-20-8	0.10	3.3	10
109. Endosulfan II	33213-65-9	0.10	3.3	10
110. 4,4'-DDD	72-54-8	0.10	3.3	10
111. Endosulfan sulfate	1031-07-8	0.10	3.3	10
112. 4,4'-DDT	50-29-3	0.10	3.3	10
·				
113. Methoxychlor	72-43-5	0.50	17.0	50
114. Endrin ketone	53494-70-5	0.10	3.3	10
115. Endrin aldehyde	7421-36-3	0.10	3.3	10
116. alpha-Chlordane	5103-71-9	0.05	1.7	. 5
117. gamma-Chlordane	5103-74-2	0.05	1.7	5
118. Toxaphene	8001-35-2	5.0	170.0	500
119. Aroclor-1016	12674-11-2	1.0	33.0	100
120. Aroclor-1221	11104-28-2	1.0	33.0	100
121. Aroclor-1232	11141-16-5	2.0	67.0	200
122. Aroclor-1242	53469-21-9	1.0	33.0	100
123. Aroclor-1248	12672-29-6	1.0	33.0	100
124. Aroclor-1254	11097-69-1	1.0	33.0	100
125. Aroclor-1260	11096-82-5	1.0	33.0	100

* Quantitation limits listed for soil/sediment are based on wet weight. The quantitation limits calculated by the laboratory for soil/sediment, calculated on dry weight basis as required by the contract, will be higher.

There is no differentiation between the preparation of low and medium soil samples in this method for the analysis of Pesticides/Aroclors.

OLMO1.0

PESTICIDES/AROCLORS

Water Samples

A 1 L volume of water is extracted three times with 100 mL of methylene chloride. This extract is reduced in volume to approximately 3-5 mL, and diluted up to 10.0 mL with clean solvent. Although not required, if Gel Permeation Chromatography is performed, only 5 of the 10 mL of extract are collected after GPC.

Regardless of whether GPC is performed, only 1.0 mL of the 10.0 mL of the original extract is taken through the remaining clean up steps (Florisil and sulfur removal). The final volume of the extract after the clean up steps depends on the requirements of the autosampler. If the autosampler can handle 1.0 mL final extract volumes, this is the final volume. If the autosampler cannot reliably handle 1.0 mL volumes, the final volume is 2.0 mL. When using an autosampler, the injection volume may be 1.0 or 2.0 uL. Manual injections must use a 2.0 uL injection volume.

For a sample with compound X at the CRQL of 0.05 ug/L:

(0.05 ug/L) (1 L) = 0.05 ug in the original extract

This material is contained in the 10.0 mL of extract:

(0.05 ug)/(10.0 mL) = 0.005 ug/mL

Of which, only 1.0 mL is carried through the remaining clean up steps. For a final extract volume of 1.0 mL and a 1 uL injection volume:

 $(0.005 \text{ ug/mL})(1 \text{ uL})(10^{-3} \text{ mL/uL}) = 5 \times 10^{-6} \text{ ug} = 5 \text{ pg on the GC column}$

Soil Samples

There is no differentiation between the preparation of low and medium soil samples in this method for the analysis of pesticides/Aroclors. A 30 g soil sample is extracted three times with methylene chloride by sonication. The extract is reduced in volume to 10.0 mL and subjected to Gel Permeation Chromatography. After GPC, only 5.0 mL of extract are collected. However, as with the water sample described above, only 1.0 mL of that extract is subjected to the other clean up steps, so no loss of sensitivity results from the use of GPC. From this point on, the soil sample extract is handled in the same fashion as the extract of a water sample. For a sample with compound X at the CRQL of 1.7 ug/Kg:

(1.7 ug/Kg) (30 g) $(10^{-3} \text{ Kg/g}) = 51 \times 10^{-3} \text{ ug} = 51 \text{ ng in the original extract}$

This material is contained in the 10.0 mL of extract:

(51 ng) / 10 mL = 5.1 ng/mL

(continued)

Pesticides/Aroclors, continued -

Of which, only 1.0 mL is carried through the remaining clean up steps. For a final extract volume of 1.0 mL and a 1 uL injection volume:

 $(5.1 \text{ ng/mL})(1 \text{ uL})(10^{-3} \text{ mL/uL}) = 5.1 \times 10^{-3} \text{ ng} = 5 \text{ pg}$ on the GC column.

For either water or soil samples, if the autosampler used requires a 2.0 mL final volume, the concentration in the 10.0 mL of extract above remains the same.

Using a 2 uL injection volume, twice the total number of picograms are injected onto the GC column. However, because the injection volume must be the same for samples and standards, twice as much material is injected onto the column during calibration, and thus the amount of compound X injected from the sample extract is equivalent to the amount of compound X injected from the calibration standard, regardless of injection volume.

EXHIBIT D

ANALYTICAL METHODS FOR VOLATILES

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SECTION I

INTRODUCTION

The analytical methods that follow are designed to analyze water, sediment and soil from hazardous waste sites for the organic compounds on the Target Compounds List (TCL, see Exhibit C). The methods are based on EPA Method 624 (Purgeables).

The methods are divided into the following sections: sample preparation, screening, and analysis. Sample preparation covers sample storage, sample holding times, and medium level sample extraction. As described in the screening section, a portion of a hexadecane extract may be screened on a gas chromatograph with appropriate detectors to determine the concentration level of organics. The analysis section contains the GC/MS analytical methods for organics. The purge and trap technique, including related sample preparation, is included in the analysis section because GC/MS operation and the purge and trap technique are interrelated.

1. Method for the Determination of Volatile (Purgeable) Organic Compounds

1.1 Scope and Application

This method covers the determination of the target volatile (purgeable) organics as listed in Exhibit C. The contract required quantitation limits are also listed in Exhibit C. The method includes an optional hexadecane screening procedure. The extract is screened on a gas chromatograph/flame ionization detector (GC/FID) to determine the approximate concentration of organic constituents in the sample. The actual analysis is based on a purge and trap gas chromatographic/mass spectrometer (GC/MS) method. For soil/sediment samples, the purge device is heated.

- 1.2 Problems have been associated with the following compounds analyzed by this method:
 - o Chloromethane, vinyl chloride, bromomethane, and chloroethane can display peak broadening if the compounds are not delivered to the GC column in a tight band.
 - o Acetone, hexanone, 2-butanone, and 4-methyl-2-pentanone have poor purge efficiencies.
 - o 1,1,1-Trichloroethane and all the dichloroethanes can dehydrogenate during storage or analysis.
 - o Tetrachloroethane and 1,1-dichloroethane can be degraded by contaminated transfer lines in purge and trap systems and/or active sites in trapping materials.
 - o Chloromethane can be lost if the purge flow is too fast.
 - Bromoform is one of the compounds most likely to be adversely affected by cold spots and/or active sites in the transfer lines. Response of its quantitation ion (m/z 173) is directly affected by the tuning of the GC/MS to meet the instrument performance criteria for BFB at ions m/z 174/176. Increasing the m/z 174/176 ratio may improve bromoform response.
- 1.3 Because performance data indicate erratic and poor linearity, the following compounds have no Maximum %RSD, or Maximum %Difference criteria; however, these compounds <u>must</u> meet a minimum RRF criterion of 0.010:

Acetone	1,2-Dichloropropane
2-Butanone	2-Hexanone
Carbon disulfide	Methylene chloride
Chloroethane	4-Methyl-2-pentanone
Chloromethane	Toluene-dg
1,2-Dichloroethene (total)	1,2-Dichloroethane-d ₄

SECTION II

SAMPLE PREPARATION AND STORAGE

• • •

PART A - SAMPLE STORAGE AND HOLDING TIMES

1. <u>Procedures for Sample Storage</u>

The samples must be protected from light and refrigerated at $4^{\circ}C$ ($\pm 2^{\circ}C$) from the time of receipt until 60 days after delivery of a complete sample data package to the Agency. After 60 days, the samples may be disposed of in a manner that complies with all applicable regulations.

The samples must be stored in an atmosphere demonstrated to be free of all potential contaminants and in a refrigerator used only for storage of purgeable samples received under this contract.

Samples and standards must be stored separately.

2. <u>Contract Required Holding Times</u>

Analysis of water samples must be completed within 10 days of validated time of sample receipt (VTSR), and analysis of soil/sediment samples must be completed within 10 days of VTSR.

PART B - PROTOCOLS FOR HEXADECANE EXTRACTION OF VOLATILES FROM WATER AND SOIL/SEDIMENT FOR OPTIONAL SCREENING

- 1. <u>Summary of Method</u>
- 1.1 Matrices
 - 1.1.1 Water a 40 mL aliquot of sample is extracted with 2 mL of hexadecane. This provides a minimum quantitation limit (MQL) as follows:

Compounds	MOL ug/L
non-halogenated aromatics	40- 50
halogenated methanes	800-1000
halogenated ethanes	400- 500

- 1.1.2 Soil/sediment 40 mL of reagent water are added to 10 g (wet weight) of soil and shaken. The water phase is in turn extracted with 2 mL of hexadecane. This provides a minimum quantitation limit of approximately four times higher than those listed for water.
- 1.2 The hexadecane extraction and screening protocols for purgeables are optional. These protocols are included to aid the analyst in deciding whether a sample is low or medium level. The use of these or other screening protocols could prevent saturation of the purge and trap system and/or the GC/MS system. It is recommended that these or other screening protocols be used, particularly if there is some doubt about the level of organics in a sample. This is especially true in soil/sediment analysis.
- 2. Limitations

These extraction and preparation procedures were developed for rapid screening of water samples from hazardous waste sites. The design of the methods thus does not stress efficient recoveries or low limits of quantitation. Rather, the procedures were designed to screen at moderate recovery and sufficient sensitivity for a broad spectrum of organic chemicals. The results of the analyses thus may reflect only a minimum of the amount actually present in some samples. This is especially true if water soluble solvents are present.

3. <u>Interferences</u>

3.1 Method interferences may be caused by contaminants in solvents, reagents, glassware, and other sample processing hardware that lead to discrete artifacts and/or elevated baselines in the total ion current profiles. All of these materials must be routinely demonstrated to be free from interferences under the conditions of the analysis by running laboratory reagent blanks. Matrix interferences may be caused by contaminants that are coextracted from the sample. The extent of matrix interferences will vary considerably from source to source, depending upon the nature and diversity of the site being sampled.

4. Apparatus and Materials

- 4.1 Vials and caps 2 mL for GC auto sampler.
- 4.2 Volumetric flask 50 mL with ground glass stopper.
- 4.3 Pasteur pipets disposable.
- 4.4 Centrifuge tube 50 mL with ground glass stopper or Teflon-lined screw cap.
- 4.5 Balance analytical, capable of accurately weighing \pm 0.0001 g.

5. <u>Reagents</u>

- 5.1 Hexadecane and methanol pesticide residue analysis grade or equivalent.
- 5.2 Reagent water defined as water in which an interferent is not observed at the CRQL of each parameter of interest.
- 5.3 Standard mixture #1 containing benzene, toluene, ethyl benzene and xylene. Standard mixture #2 containing n-nonane and n-dodecane.
 - 5.3.1 Stock standard solutions (1.00 ug/uL) prepared from pure standard materials or purchased as certified solutions.
 - 5.3.1.1 Prepare stock standard solutions by accurately weighing about 0.0100 g of pure material. Dissolve the material in methanol and dilute to volume in a 10 mL volumetric flask. Larger volumes can be used at the convenience of the analyst. If compound purity is certified at 96% or greater, the weight can be used without correction to calculate the concentration of the stock standard.
 - 5.3.1.2 Transfer the stock standard solutions into multiple Teflon-sealed screw-cap vials. Store, with no head-space, at -10°C to -20°C, and protect from light. Stock standard solutions should be checked frequently for signs of degradation or evaporation, especially just prior to preparing calibration standards from them. These solutions must be replaced after six months, or sooner, if comparison with quality control check samples indicates a problem. Standards prepared from gases or reactive compounds such as styrene must be replaced after two months, or sooner, if comparison with quality control check samples indicates a problem.
 - 5.3.2 Prepare working standards of mixtures #1 and #2 at 100 ng/uL of each compound in methanol. Store these solutions as in 5.3.1.2 above.

6. Sample Extraction

6.1 Water

- 6.1.1 Allow the contents of the 40 mL sample vial to come to room temperature. Quickly transfer the contents of the 40 mL sample vial to a 50 mL volumetric flask. Immediately add 2.0 mL of hexadecane, cap the flask, and shake vigorously for 1 minute. Let phases separate. Open the flask and add sufficient reagent water to bring the hexadecane layer into the neck of the flask.
- 6.1.2 Transfer approximately 1 mL of the hexadecane layer to a 2.0 mL GC vial. If an emulsion is present after shaking the sample, break it by doing the following:
 - Pulling the emulsion through a small plug of Pyrex glass wool packed in a pipet, or
 - Transferring the emulsion to a centrifuge tube and centrifuging for several minutes.
- 6.1.3 Add 200 uL of working standard mixture #1 and #2 to separate 40 mL portions of reagent water. Follow steps 6.1.1 6.1.2, starting with the immediate addition of 2.0 mL of hexadecane.

6.2 Soil/Sediment

- 6.2.1 Add approximately 10 g of soil (wet weight) to 40 mL of reagent water in a 50 mL centrifuge tube with a ground glass stopper or teflon-lined cap. Cap and shake vigorously for one minute. Centrifuge the capped flask briefly. Quickly transfer supernatant water to a 50 mL volumetric flask equipped with a ground-glass stopper.
- 6.2.2 Follow 6.1, starting with the addition of 2.0 mL of hexadecane.

7. <u>Sample Analysis</u>

The sample is ready for GC/FID screening. Proceed to Section III, Optional Screening of Hexadecane Extracts for Volatiles. SECTION III

OFTIONAL SCREENING OF HEXADECANE EXTRACTS FOR VOLATILES

1. <u>Summary of Method</u>

The hexadecane extracts of water and soil/sediment are screened on a gas chromatograph/flame ionization detector (GC/FID). The results of the screen will determine if volatile organics are to be analyzed by low or medium level GC/MS procedures if the sample is a soil/sediment, or to determine the appropriate dilution factor if the sample is water.

2. Apparatus and Materials

- 2.1 Gas chromatograph an analytical system complete with gas chromatograph suitable for on-column injection and all required accessories including syringes, analytical columns, gases, detector, and strip-chart recorder. A data system is recommended for measuring peak areas.
 - 2.1.1 Above-described GC, equipped with flame ionization detector.
 - 2.1.2 GC column 3 m x 2 mm ID glass column packed with 10% OV-101 on 100-120 mesh Chromosorb W-HP (or equivalent). The column temperature should be programmed from 80°C to 280°C at 16°C/min. and held at 280°C for 10 minutes.

3. <u>Reagents</u>

Hexadecane - pesticide residue analysis grade or equivalent.

- 4. Limitations
- 4.1 The flame ionization detector varies considerably in sensitivity when comparing aromatics and halogenated methanes and ethanes. Halomethanes are approximately 20x less sensitive than aromatics and haloethanes approximately 10x less sensitive. Low molecular weight, water soluble solvents, e.g., alcohols and ketones, will not extract from the water, and therefore will not be detected by the GC/FID.
- 4.2 Following are two options for interpreting the GC/FID chromatogram.
 - 4.2.1 Option A is to use standard mixture #1 containing the aromatics to calculate an approximate concentration of the aromatics in the sample. Use this information to determine the proper dilution for purge and trap if the sample is a water, or whether to use the low or medium level GC/MS purge and trap methods if the sample is a soil/sediment (see Table 1, paragraph 6.2.1.3 for guidance). This should be the best approach; however, the aromatics may be absent or obscured by higher concentrations of other purgeables. In these cases, Option B may be the best approach.
 - 4.2.2 <u>Option B</u> is to use standard mixture #2 containing n-nonane and n-dodecane to calculate a factor. Use the factor to calculate a dilution for purge and trap of a water sample or to determine whether to use the low or medium level GC/MS purge and trap methods for soil/sediment samples (see Table 1, paragraph

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6.2.1.3 for guidance). All purgeables of interest have retention times less than the n-dodecane.

- 5. <u>Extract Screening</u>
- 5.1 External standard calibration Standardize the GC/FID each 12 hr. shift for half scale response. This is done by injecting 1-5 uL of the extracts that contain approximately 10 ng/uL each of mix #1 and mix #2 compounds, as prepared in 5.3, Section II, Part B. Use the GC conditions specified in paragraph 2.1.2.
- 5.2 Inject the same volume of hexadecane extract as the extracted standard mixture in 5.1. Use the GC conditions specified in 2.1.2.
- 6. Analytical Decision Point
- 6.1 Water
 - 6.1.1 Compare the chromatograms of the hexadecane extract of the sample with those of the reagent blank and extract of the standard.
 - 6.1.1.1 If no peaks are noted, other than those also in the reagent blank, analyze a 5 mL water sample by purge and trap GC/MS.
 - 6.1.1.2 If peaks are present prior to the n-dodecane and the aromatics are distinguishable, follow Option A (4.2.1).
 - 6.1.1.3 If peaks are present prior to the n-dodecane but the aromatics are absent or indistinguishable, use Option B as follows: if all peaks are ≤3% of the n-nonane, analyze a 5 mL water sample by purge and trap GC/MS. If any peaks are ≥3% of the n-nonane, measure the peak height or area of the major peak and calculate the dilution factor as follows:

peak area of sample major peak x 50 = dilution peak area of n-nonane factor

The water sample will be diluted using the calculated factor just prior to purge and trap GC/MS analysis.

6.2 Soil/Sediment

- 6.2.1 Compare the chromatograms of the hexadecane extract of the sample with those of the reagent blank and extract of the standard.
 - 6.2.1.1 If no peaks are noted, other than those also in the reagent blank, analyze a 5 g sample by low level GC/MS.

- 6.2.1.2 If peaks are present prior to the n-dodecane and the aromatics are distinguishable, follow Option A (4.2.1) and the concentration information in Table 1, paragraph 6.2.1.3, to determine whether to analyze by low or medium level method.
- 6.2.1.3 If peaks are present prior to the n-dodecane but the aromatics are absent or indistinguishable, and using Option B as follows, calculate a factor using the following formula:

<u>peak area of sample major peak</u> - X Factor peak area of n-nonane

Table 1 - Determination of GC/MS Purge & Trap Method

X Factor	Analyze by	Approximate Concentration Range* (ug/kg)	
0-1.0	low level method	0-1,000	
>1.0	medium level method	>1,000	

* This concentration range is based on the response of aromatics to GC/FID. When comparing GC/FID responses, the concentration for halomethanes is 20x higher, and that for haloethanes is 10x higher.

6.3 Sample Analysis

Proceed to Section IV, GC/MS Analysis of Volatiles.

SECTION IV

GC/MS ANALYSIS OF VOLATILES

1. <u>Summary of Methods</u>

1.1 Water Samples

An inert gas is bubbled through a 5 mL sample contained in a specifically designed purging chamber at ambient temperature. The purgeables are efficiently transferred from the aqueous phase to the vapor phase. The vapor is swept through a sorbent column where the purgeables are trapped. After purging is completed, the sorbent column is heated and backflushed with the inert gas to desorb the purgeables onto a gas chromatographic column. The gas chromatograph is temperature programmed to separate the purgeables which are then detected with a mass spectrometer.

An aliquot of the sample is diluted with reagent water when dilution is necessary. A 5 mL aliquot of the dilution is taken for purging.

1.2 Soil/Sediment Samples

- 1.2.1 Low level an inert gas is bubbled through a mixture of a 5 g sample and reagent water contained in a suggested specially designed purging chamber at elevated temperatures. The purgeables are efficiently transferred from the aqueous phase to the vapor phase. The vapor is swept through a sorbent column where the purgeables are trapped. After purging is completed, the sorbent column is heated and backflushed with the inert gas to desorb the purgeables onto a gas chromatographic column. The gas chromatograph is temperature programmed to separate the purgeables, which are then detected with a mass spectrometer.
- 1.2.2 Medium level a measured amount of soil is extracted with methanol. A portion of the methanol extract is diluted to 5 mL with reagent water. An inert gas is bubbled through this solution in a specifically designed purging chamber at ambient temperature. The purgeables are effectively transferred from the aqueous phase to the vapor phase. The vapor is swept through a sorbent column where the purgeables are trapped. After purging is completed, the sorbent column is heated and backflushed with the inert gas to desorb the purgeables onto a gas chromatographic column. The gas chromatograph is temperature programmed to separate the purgeables, which are then detected with a mass spectrometer.

2. <u>Interferences</u>

2.1 Impurities in the purge gas, organic compounds out-gassing from the plumbing ahead of the trap, and solvent vapors in the laboratory account for the majority of contamination problems. The analytical system must be demonstrated to be free from contamination under the conditions of the analysis by running laboratory reagent blanks as described in Exhibit E. The use of non-TFE tubing, non-TFE thread sealants, or flow controllers with rubber components in the purging device should be avoided.

- 2.2 Samples can be contaminated by diffusion of volatile organics (particularly fluorocarbons and methylene chloride) through the septum seal into the sample during storage and handling. A holding blank prepared from reagent water and carried through the holding period and the analysis protocol serves as a check on such contamination. One holding blank per case should be analyzed. Data must be retained by the laboratory and be made available for inspection during on-site evaluations.
- 2.3 Contamination by carry-over can occur whenever high level and low level samples are sequentially analyzed. To reduce carryover, the purging device and sampling syringe must be rinsed with reagent water between sample analyses. Whenever an unusually concentrated sample is encountered, it should be followed by an analysis of reagent water to check for cross contamination. For samples containing large amounts of water soluble materials, suspended solids, high boiling compounds or high purgeable levels, between analyses, it may be necessary to wash out the purging device with a detergent solution, rinse it with distilled water, and then dry it in a 105°C oven. The trap and other parts of the system are also subject to contamination; therefore, frequent bakeout and purging of the entire system may be required.
- 2.4 The laboratory where volatile analysis is performed should be completely free of solvents.
- 3. Apparatus and Materials
- 3.1 Micro syringes 25 uL and larger, 0.006 inch ID needle.
- 3.2 Syringe valve two-way, with Luer ends (three each), if applicable to the purging device.
- 3.3 Syringe 5 mL, gas-tight with shut-off valve.
- 3.4 Balance analytical, capable of accurately weighing \pm 0.0001 g, and a top-loading balance capable of weighing \pm 0.1 g.
- 3.5 Glassware
 - o Bottle 15 mL, screw cap, with Teflon cap liner.
 - o Volumetric flasks class A with ground-glass stoppers.
 - o Vials 2 mL for GC autosampler.
- 3.6 Purge and trap device consists of three separate pieces of equipment: the sample purger, trap, and the desorber. Several complete devices are now commercially available.
 - 3.6.1 The sample purger must be designed to accept 5 mL samples with a water column at least 3 cm deep. The gaseous head space between the water column and the trap must have a total volume of less than 15 mL. The purge gas must pass through the water column as finely divided bubbles, each with a diameter of less

than 3 mm at the origin. The purge gas must be introduced no more than 5 mm from the base of the water column.

- 3.6.2 The trap must be at least 25 cm long and have an inside diameter of at least 0.105 inch. The trap must be packed to contain the following minimum lengths of absorbents: 15 cm of 2,6-diphenylene oxide polymer (Tenax-GC, 60/80 mesh) and 8 cm of silica gel (Davison Chemical, 35/60 mesh, grade 15, or equivalent).
- 3.6.3 The desorber should be capable of rapidly heating the trap to 180°C. The polymer section of the trap should not be heated higher than 180°C and the remaining sections should not exceed 220°C during bakeout mode.
- 3.6.4 The purge and trap device may be assembled as a separate unit or be coupled to a gas chromatograph.
- 3.6.5 A heater or heated bath capable of maintaining the purge device at $40^{\circ}C \pm 1^{\circ}C$ is to be used.

3.7 GC/MS system

- 3.7.1 Gas Chromatograph the gas chromatograph (GC) system must be capable of temperature programming and have a flow controller that maintains a constant column flow rate throughout desorption and temperature program operations. The system must include or be interfaced to a purge and trap system as specified in paragraph 3.6 and have all required accessories including syringes, analytical columns, and gases. All GC carrier gas lines must be constructed from stainless steel or copper tubing. Non-polytetrafluoroethylene (PTFE) thread sealants, or flow controllers with rubber components are not to be used. If capillary columns are to be used (see below), the column oven must be cooled to 10°C; therefore, a subambient oven controller is required.
- 3.7.2 Gas Chromatography Columns
 - 3.7.2.1 Packed Columns 6 ft long x 0.1 in ID glass, packed with 1% SP-1000 on Carbopack B (60/80 mesh) or equivalent.

NOTE: Capillary columns may be used for analysis of volatiles, as long as the Contractor uses the instrumental parameters in EPA Method 524.2 as guidelines, uses the internal standards and surrogates specified in this contract, and demonstrates that the analysis meets all of the performance and QA/QC criteria contained in this contract.

- 3.7.2.2 Capillary Columns
 - o 30 m long x 0.53 mm ID VOCOL (Supelco, Inc., or equivalent) fused silica wide-bore capillary column with 3 um film thickness.

OR

- o 30 m long x 0.53 mm ID DB-624 fused silica widebore (J&W Scientific, Inc., or equivalent) column with 3 um film thickness.
- Mass Spectrometer must be capable of scanning from 35 to 300 3.7.3 amu every 1 second or less, utilizing 70 volts (nominal) electron energy in the electron impact ionization mode, and producing a mass spectrum which meets all the instrument performance acceptance criteria when 50 ng of p-bromofluorobenzene (BFB) is injected through the gas chromatograph inlet. The instrument conditions required for the acquisition of the BFB mass spectrum are given in paragraph 6.4.4. NOTE: BFB criteria must be met before any sample extracts are analyzed. Any samples analyzed when BFB criteria have not been met will require reanalysis at no cost to the Agency. To ensure sufficient precision of mass spectral data, the MS scan rate should allow acquisition of at least five spectra while a sample compound elutes from the GC. The purge and trap GC/MS system must be in a room whose atmosphere is demonstrated to be free of all potential contaminants which will interfere with the analysis. The instrument must be vented to the outside of the facility or to a trapping system which prevents the release of contaminants into the instrument room.
- 3.7.4 GC/MS interface any gas chromatograph to mass spectrometer interface that gives acceptable calibration points, at 50 ng or less per injection, for each of the parameters of interest and achieves all acceptance criteria may be used. Gas chromatograph to mass spectrometer interfaces constructed of all-glass or glass-lined materials are recommended. Glass can be deactivated by silanizing with dichlorodimethylsilane.
- 3.7.5 Data system a computer system must be interfaced to the mass spectrometer that allows the continuous acquisition and storage, on machine readable media, of all mass spectra obtained throughout the duration of the chromatographic program. The computer must have software that allows searching any GC/MS data file for ions of a specified mass and plotting such ion abundances versus time or scan number. This type of plot is defined as an Extracted Ion Current Profile (EICP). Software must also be available that allows integrating the abundance in any EICP between specified time or scan number limits. Also, for the non-target compounds, software must be available that allows for the comparison of sample spectra against reference library spectra. The 1989 (or most recent)

OLM01.0

release of the NIST/EPA/MSDC mass spectral library shall be used as the reference library. The data system must be capable of flagging all data files that have been edited manually by laboratory personnel.

3.7.6 Magnetic tape storage device - must be capable of recording data and must be suitable for long-term, off-line storage.

4. Reagents

- 4.1 Reagent water defined as water in which an interferent is not observed at or above the CRQL of the parameters of interest.
 - 4.1.1 Reagent water may be generated by passing tap water through a carbon filter bed containing about 453 g of activated carbon (Calgon Corp., Filtrasorb-300 or equivalent).
 - 4.1.2 A water purification system (Millipore Super-Q or equivalent) may be used to generate reagent water.
 - 4.1.3 Reagent water may also be prepared by boiling water for 15 minutes. Subsequently, while maintaining the temperature at 90°C, bubble a contaminant-free inert gas through the water for one hour. While still hot, transfer the water to a narrow-mouth screw-cap bottle and seal with a Teflon-lined septum and cap.
- 4.2 Sodium thiosulfate (ACS) granular.
- 4.3 Methanol pesticide quality or equivalent.

5. <u>Standards</u>

- 5.1 The Contractor must provide all standards to be used with this contract. These standards may be used only after they have been certified according to the procedure in Exhibit E. The Contractor must be able to verify that the standards are certified. Manufacturer's certificates of analysis must be retained by the Contractor and presented upon request.
- 5.2 Stock Standard Solutions

Stock standard solutions may be purchased or may be prepared in methanol from pure standard materials.

5.2.1 Prepare stock standard solutions by placing about 9.8 mL of methanol into a 10.0 mL ground-glass stoppered volumetric flask. Allow the flask to stand, unstoppered, for about 10 minutes, or until all alcohol wetted surfaces have dried. Weigh the flask to the nearest 0.1 mg.

- 5.2.2 Add the assayed reference material as described below.
 - 5.2.2.1 If the compound is a liquid, using a 100 μ L syringe, immediately add two or more drops of assayed reference material to the flask, then reweigh. The liquid must fall directly into the alcohol without contacting the neck of the flask.
 - 5.2.2.2 If the compound is a gas at room temperature, fill a 5 mL valved gas-tight syringe with the reference standard to the 5.0 mL mark. Lower the needle to 5 mm above the methanol meniscus. Slowly introduce the reference standard above the surface of the liquid. The gas will rapidly dissolve in the methanol. This may also be accomplished by using a lecture bottle equipped with a Hamilton Lecture Bottle Septum (#86600). Attach Teflon tubing to the side-arm relief valve and direct a gentle stream of the reference standard into the methanol meniscus.
- 5.2.3 Reweigh, dilute to volume, stopper, then mix by inverting the flask several times. Calculate the concentration in micrograms per microliter from the net gain in weight. When compound purity is assayed to be 97 percent or greater, the weight may be used, without correction, to calculate the concentration of the stock standard. If the compound purity is assayed to be less than 97 percent, the weight must be corrected when calculating the concentration of the stock solution.
- 5.2.4 Prepare fresh stock standards every two months for gases or for reactive compounds such as styrene. All other stock standards for non-gases/non-reactive purgeable compounds must be replaced after six months, or sooner if standard has degraded or evaporated.
- 5.3 Secondary Dilution Standards
 - 5.3.1 Using stock standard solutions, prepare secondary dilution standards in methanol that contain the compounds of interest, either singly or mixed together. Secondary dilution standard solutions should be prepared at concentrations that can be easily diluted to prepare working standard solutions.
 - 5.3.2 Prepare fresh secondary dilution standards for gases and for reactive compounds such as styrene every month, or sooner, if standard has degraded or evaporated. Secondary dilution standards for the other purgeable compounds must be replaced after six months, or sooner if standard has degraded or evaporated.

5.4 Working Standards

5.4.1 Instrument Performance Check Solution - p-Bromofluorobenzene (BFB)

> Prepare a 25 ng/ μ L solution of BFB in methanol. Prepare fresh BFB solution every six months, or sooner, if the solution has degraded or evaporated. NOTE: The 25 ng/ μ L concentration is used with a 2 μ L injection volume. The laboratory may prepare a 50 ng/ μ L solution of BFB if a 1 μ L injection volume is used.

5.4.2 Calibration Standard Solution

Prepare the working calibration standard solution containing all of the purgeable target compounds in methanol. The recommended concentration of the target compounds is 100 ug/mL. Prepare fresh working calibration standard solutions weekly, or sooner, if solutions have degraded or evaporated.

5.4.3 Internal Standard Spiking Solution

Prepare an internal standard spiking solution containing Bromochloromethane, Chlorobenzene-d₅, and 1,4-Difluorobenzene in methanol at the concentration of 25.0 ug/mL for each internal standard. Add 10 μ L of this spiking solution into 5.0 mL of sample or calibration standard for a concentration of 50 μ g/L. Prepare fresh spiking solution weekly or sooner if the solution has degraded or evaporated.

5.4.4 System Monitoring Compound (SMC) Spiking Solution

Prepare a system monitoring compound spiking solution containing Toluene-d₈, p-Bromofluorobenzene, and 1-2-Dichloroethane-d₄ in methanol at a concentration of 25.0 μ g/mL. Add 10.0 μ L of this spiking solution into 5.0 mL of sample, for a concentration of 50 μ g/L. Prepare fresh spiking solution every three months, or sooner if the solution has degraded or evaporated.

- 5.4.5 Volatile Matrix Standard Spiking Solution
 - 5.4.5.1 Prepare a spiking solution in methanol that contains the following compounds at a concentration of 25.0 μ g/mL: 1,1-Dichloroethene, Trichloroethene, Chlorobenzene, Toluene, and Benzene.
 - 5.4.5.2 Matrix spikes also serve as duplicates; therefore, add an aliquot of this solution to each of two portions from one sample chosen for spiking.

5.5 Aqueous Calibration Standard Solutions

- 5.5.1 Prepare five aqueous initial calibration standard solutions containing all of the purgeable target compounds and system monitoring compounds at the 10, 20, 50, 100, and 200 μ g/L levels. Note: These are not the same levels as have been used in previous Statements of Work.
- 5.5.2 Aqueous calibration standards may be prepared in a volumetric flask or in the syringe used to inject the standard into the purging device.
 - 5.5.2.1 Volumetric flask add an appropriate volume of working calibration standard solution to an aliquot of reagent water in a volumetric flask. Use a microsyringe and rapidly inject the alcohol standard into the expanded area of the filled volumetric flask. Remove the needle as quickly as possible after injection. Bring to volume. Mix by inverting the flask three times only. Discard the contents contained in the head of the flask.
 - 5.5.2.2 Syringe remove the plunger from a 5 mL "Luerlock" syringe. Pour reagent water into the syringe barrel to just short of overflowing. Replace the syringe plunger and compress the water. Invert the syringe, open the syringe valve and vent any residual air. Adjust the water volume to 5.0 mL minus the amount of calibration standard to be added. Withdraw the plunger slightly and add an appropriate volume of working calibration standard through the valve bore of the syringe. Close the valve and invert three times.
- 5.5.3 The 50 μ g/L aqueous calibration standard solution is the continuing calibration standard.
- 5.5.4 The methanol purged in each of the aqueous calibration standards must not exceed 1% by volume.
- 5.6 Storage of Standards
 - 5.6.1 Store the stock standards in Teflon-sealed screw-cap bottles with zero headspace at -10°C to -20°C. Protect the standards from light. Once one of the bottles containing the stock standard solution has been opened, it may be used for no longer than one week.
 - 5.6.2 Store secondary dilution standards in Teflon-sealed screw-cap bottles with minimal headspace at -10°C to -20°C. Protect the standards from light. The secondary dilution standards must be checked frequently for signs of degradation or evaporation, especially just prior to preparing the working calibration standards from them.

- 5.6.3 Aqueous standards may be stored for up to 24 hours if held in Teflon-sealed screw-cap vials with zero headspace at 4°C. Protect the standards from light. If not so stored, they must be discarded after one hour unless they are set up to be purged by an autosampler. When using an autosampler, the standards may be kept for up to 12 hours in purge tubes connected via the autosampler to the purge and trap device.
- 5.6.4 Purgeable standards must be stored separately from other standards.

6. Instrument Operating Conditions

6.1 Purge and Trap Device

The following are the purge and trap analytical conditions:

Purge Conditions:

Purge	Gas:	Helium or Nitrogen
Purge	Time:	11.0 ±0.1 min
Purge	Flow Rate:	25-40 mL/min
Purge	Temperature:	Ambient

Desorb Conditions:

Desorb	Temperature:	180°C
Desorb	Flow Rate:	15 mL/min
Desorb	Time:	4.0 ±0.1 min

Trap Reconditioning Conditions:

Reconditioning Temperature:	180°C
Reconditioning Time:	7.0 min ±0.1 min

Before initial use, condition the trap overnight at 180°C by backflushing with at least 20 mL/min flow of inert gas. Vent the trap effluent to the room and not to the analytical column. Prior to daily use, condition the trap by heating at 180°C for 10 minutes while backflushing. The trap may be vented to the analytical column during daily conditioning; however, the column must be run through the temperature program prior to analysis of samples.

6.2 Gas Chromatograph

The following are the recommended gas chromatographic analytical conditions:

6.2.1 Packed Columns

Carrier Gas:	Helium
Flow Rate:	30 mL/min
Initial Temperature:	45°C
Initial Hold Time:	3 min
Ramp Rate:	8°C/min
Final Temperature:	220 [•] C
Final Hold Time:	15 min
Transfer Line Temperature:	250-300°C

6.2.2 Capillary Columns

Carrier Gas:	Helium
Flow Rate:	15 mL/min
Initial Temperature:	10°C
Initial Hold Time:	1.0 - 5.0 min (± 0.1 min precision)
Ramp Rate:	6°C/min
Final Temperature:	160°C
Final Hold Time:	Until all target compounds elute

6.2.3 Optimize GC conditions for compound separation and sensitivity.

6.3 Mass Spectrometer

The following are the required mass spectrometer conditions:

Electron Energy:	70 Volts (nominal)
Mass Range:	35-300 anu
Scan Time:	To give at least 5 scans per peak, not to exceed 1 second per scan

- 6.4 The GC/MS system must be tuned to meet the manufacturer's specifications, using a suitable calibrant such as FC-43 or perfluorokerosene (PFK). The mass calibration and resolution of the GC/MS system are verified by the analysis of the instrument performance check solution (paragraph 5.4.1).
 - 6.4.1 Prior to the analyses of any samples, blanks, or calibration standards, the Contractor must establish that the GC/MS system meets the mass spectral ion abundance criteria for the instrument performance check solution containing pbromofluorobenzene (BFB).

- 6.4.2 The analysis of the instrument performance check solution may be performed as follows:
 - o As an injection of up to 50 ng of BFB into the GC/MS.
 - o By adding 50 ng of BFB to 5.0 ml of reagent water and analyzing the resulting solution as if it were an environmental sample (see section 8 below).

BFB may not be analyzed simultaneously with a calibration standard.

- 6.4.3 The mass spectrum of BFB must be acquired in the following manner. Three scans (the peak apex scan and the scans immediately preceding and following the apex) are acquired and averaged. Background subtraction is required, and must be accomplished using a single scan prior to the elution of BFB. NOTE: All instrument conditions must be identical to those used in the sample analysis.
- 6.4.4 The analysis of the instrument performance check solution must meet the ion abundance criteria given below.

			TABL	E 1		
BFB	KEY	IONS	AND	ION	ABUNDANCE	CRITERIA

Mass	Ion Abundance Criteria
50	8.0 - 40.0 percent of mass 95
75	30.0 - 66.0 percent of mass 95
95	base peak, 100 percent relative abundance
96	5.0 - 9.0 percent of mass 95 (see note)
173	less than 2.0 percent of mass 174
174	50.0 - 120.0 percent of mass 95
175	4.0 - 9.0 percent of mass 174
176	93.0 - 101.0 percent of mass 174
177	5.0 - 9.0 percent of mass 176

Note: All ion abundances must be normalized to m/z 95, the nominal base peak, even though the ion abundance of m/z 174 may be up to 120 percent that of m/z 95.

6.4.5 The criteria listed above are based on adherence to the acquisition specifications identified in paragraph 6.4.3, and were developed for the specific target compound list associated with this Statement of Work. The criteria are based on performance characteristics of instruments currently utilized in routine support of Program activities. These specifications, in conjunction with relative response factor criteria for 23 target compounds (see Table 2), are designed to control and monitor instrument performance associated with the requirements of this Statement of Work. 6.4.6 The instrument performance check solution must be injected once at the beginning of each 12-hour period, during which samples or standards are to be analyzed. The twelve (12) hour time period for GC/MS Instrument Performance Check (BFB), standards calibration (initial or continuing calibration criteria) and method blank analysis begins at the moment of injection of the BFB analysis that the laboratory submits as documentation of a compliant instrument performance check. The time period ends after twelve (12) hours has elapsed according to the system clock.

7. <u>Calibration</u>

- 7.1 Prior to the analysis of samples and required blanks, and after the instrument performance check solution criteria have been met, each GC/MS system must be calibrated at a minimum of five concentrations to determine instrument sensitivity and the linearity of GC/MS response for the purgeable target compounds.
- 7.2 Assemble a purge and trap device that meets the specification in 3.6. Condition the trap overnight at 180°C in the purge mode with an inert gas flow of at least 20 mL/min. Daily, prior to use, condition the traps for 10 minutes while backflushing at 180°C with the column at 220°C.
- 7.3 Connect the purge and trap device to a gas chromatograph. The gas chromatograph must be operated using temperature and flow rate parameters equivalent to those in 6.2. Calibrate the purge and trap-GC/MS system using the internal standard technique (7.4).
- 7.4 Internal standard calibration procedure. The three internal standards are Bromochloromethane, 1,4-Difluorobenzene, and Chlorobenzene-d₅, at 50 ug/L at time of purge. Separate initial and continuing calibrations must be performed for water samples, and low level soil samples (unheated purge vs. heated purge). Extracts of medium level soil samples may be analyzed using the calibrations for water samples.
 - 7.4.1 Prepare calibration standards at a minimum of five concentration levels for each target compound and system monitoring compound, as specified in 5.5. Standards may be stored up to 24 hours, following the procedures in paragraph 5.6.3.
 - 7.4.2 Prepare a spiking solution containing each of the internal standards using the procedures described in paragraph 5.4.3.
 - 7.4.3 Verify that the GC/MS system meets the instrument performance criteria in paragraph 6.4 by injecting BFB. Analyze each calibration standard, according to paragraph 7.1, adding 10 uL of internal standard spiking solution directly to the syringe.

Tabulate the area response of the characteristic ions in the extracted ion current profile (EICP) against concentration for each compound and internal standard and calculate relative response factors (RRF) for each compound using Equation 1.

RRF -
$$\frac{A_x}{A_{is}}$$
 x $\frac{C_{is}}{C_x}$ EQ.1

Where

- Ax Area of the characteristic ion (EICP) for the compound to be measured (see Table 4)
- A_{is} Area of the characteristic ion (EICP) for the specific internal standard (see Table 3)
- C_{ie} Concentration of the internal standard

$$C_{\rm x}$$
 - Concentration of the compound to be measured

7.4.4 The average relative response factor (RRF) must be calculated for all compounds. Calculate the % Relative Standard Deviation (%RSD) of RRF values over the working range of the curve.

RSD - <u>Standard deviation</u> x 100 mean

7.4.5 Response factor criteria have been established for the calibration of 23 of the volatile target compounds and one volatile system monitoring compound. The compounds listed below (Table 2) must meet the minimum RRF and maximum %RSD criteria for the initial calibration, with allowance made for up to two volatile compounds. However, the RRFs for those two compounds must be greater than or equal to 0.010, and the %RSD of those two compounds must be less than or equal to 40.0% for the initial calibration to be acceptable.

	TABLE 2	
RELATIVE RESPONS	FACTOR CRITERIA FOR INITIAL AND CONTIN	UING
CALIBR	TION OF VOLATILE ORGANIC COMPOUNDS	

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Volatile	Minimum	Maximum	Maximum
	RRF	<u> %RSD</u>	&Diff
Compound			
Bromomethane	0.100	20.5	25.0
	0.100	20.5	25.0
Vinyl chloride 1,1-Dichloroethene	0.100	20.5	25.0
1,1-Dichloroethane	0.200	20.5	25.0
Chloroform	0.200	20.5	25.0
1,2-Dichloroethane	0.100	20.5	25.0
1,1,1-Trichloroethane	0.100	20.5	25.0
Carbon tetrachloride	0.100	20.5	25.0
Bromodichloromethane	0.200	20.5	25.0
	0.200	20.5	25.0
cis-1,3-Dichloropropene Trichloroethene	0.300	20.5	25.0
Dibromochloromethane	0.100	20.5	25.0
1,1,2-Trichloroethane	0.100	20.5	25.0
L, L, 2-IIICHIOFOGCHANG Benzene	0.500	20.5	25.0
trans-1,3-Dichloropropene	0.100	20.5	25.0
Bromoform	0.100	20.5	25.0
Tetrachloroethene	0.200	20.5	25.0
1,1,2,2-Tetrachloroethane	0.500	20.5	25.0
Toluene	0.400	20.5	25.0
Chlorobenzene	0.500	20.5	25.0
Ethylbenzene	0.100	20.5	25.0
Styrene	0.300	20.5	25.0
Xylenes (total)	0.300	20.5	25.0
Bromofluorobenzene	0.200	20.5	25.0

7.4.6 Because performance data indicate erratic and poor linearity, the following compounds have no Maximum %RSD, or Maximum %Difference criteria; however, these compounds <u>must</u> meet a minimum RRF criterion of 0.010:

Acetone	1,2-Dichloropropane
2-Butanone	2-Hexanone
Carbon disulfide	Methylene chloride
Chloroethane	4-Methyl-2-pentanone
Chloromethane	Toluene-dg
1,2-Dichloroethene (total)	1,2-Dichloroethane-d ₄

7.4.7 A check of the calibration curve must be performed once every 12 hours (see paragraph 6.4.6 for the definition of the twelve hour time period). Check the relative response factors of those compounds for which RRF values have been established. If these criteria are met, the relative response factors for all compounds are calculated and reported. A percent difference of the daily relative response factor (12 hour) compared to the average relative response factor from the initial curve is calculated. Calculate the percent difference for each compound and compare with the maximum percent difference criteria listed above. As with the initial calibration, up to two volatile compounds in Table 2 may fail to meet the minimum RRF or maximum %D criteria, but the RRFs of those two compounds must be greater than or equal to 0.010, and the percent differences must be less than or equal to 40.0% for the continuing calibration to be acceptable.

- 7.4.8 Internal standard responses and retention times in all standards must be evaluated during or immediately after data acquisition. If the retention time for any internal standard changes by more than 0.50 minutes (30 seconds) from the latest daily (12 hour) calibration standard, the chromatographic system must be inspected for malfunctions, and corrections made as required. The extracted ion current profile (EICP) of the internal standards must be monitored and evaluated for each standard. If the EICP area for any internal standard changes by more than a factor of two (-50% to +100%), the mass spectrometric system must be inspected for malfunction, and corrections made as appropriate. When corrections are made, re-analysis of samples analyzed while the system was malfunctioning is necessary.
- 7.5 Each GC/MS system must be calibrated upon award of the contract, whenever the Contractor takes corrective action which may change or affect the initial calibration criteria (i.e., ion source cleaning or repair, column removal or replacement; etc.), or if the continuing calibration acceptance criteria have not been met.
- 7.6 If time remains in the 12 hour time period after meeting the acceptance criteria for the initial calibration, samples may be analyzed. It is not necessary to analyze a continuing calibration standard, if the initial calibration meets the calibration acceptance criteria above. A method blank <u>is</u> necessary. Quantify all sample results against the initial calibration standard that is the same concentration as the continuing calibration standard (50 ug/L).
- 7.7 If time does <u>not</u> remain in the 12-hour period beginning with the injection of the instrument performance check solution, a new injection of the instrument performance check solution must be made. If the new injection meets the ion abundance criteria for BFB, then a continuing calibration standard may be injected.
- 7.8 The concentrations of volatile target compounds in the continuing calibration standard are given in paragraph 5.5.3.
- 7.9 The response factors for the continuing calibration standard must meet the criteria given in paragraph 7.4.5 prior to the analysis of any blanks or samples.

8. Sample Analysis

8.1 Water Samples

- 8.1.1 All water samples must be allowed to warm to ambient temperature before analysis.
- 8.1.2 Prior to the analysis of samples, establish the appropriate GC/MS operating conditions, as outlined in paragraphs 6-6.4.6, analyze the instrument performance check solution (6.4), and calibrate the GC/MS system according to paragraphs 7-7.7.3.
- 8.1.3 If time remains in the 12-hour period (as described in paragraph 7.6), samples may be analyzed without analysis of a continuing calibration standard.
- 8.1.4 If time does not remain in the 12-hour period since the injection of the instrument performance check solution, both the instrument performance check solution and the continuing calibration standard must be analyzed before sample analysis may begin (see paragraphs 7.7-7.9).
- 8.1.5 Adjust the purge gas (helium) flow rate to 25-40 mL/min. Variations from this flow rate may be necessary to achieve better purging and collection efficiencies for some compounds, particularly Chloromethane and Bromoform.
- Remove the plunger from a 5 mL syringe and attach a closed 8.1.6 syringe valve. Open the sample or standard bottle which has been allowed to come to ambient temperature, and carefully pour the sample into the syringe barrel to just short of overflowing. Replace the syringe plunger and compress the sample. Open the syringe valve and vent any residual air while adjusting the sample volume to 5.0 mL. This process of taking an aliquot destroys the validity of the sample for future analysis so, if there is only one VOA vial, the analyst should fill a second syringe at this time to protect against possible loss of sample integrity. This second sample is maintained only until such time as the analyst has determined that the first sample has been analyzed properly. Filling one 5 mL. syringe would allow the use of only one syringe. If an analysis is needed from the second 5 mL syringe, it must be performed within 24 hours. Care must also be taken to prevent air from leaking into the syringe.
- 8.1.7 The purgeable organics screening procedure (Section III), if used, will have shown the approximate concentrations of major sample components. If a dilution of the sample was indicated, this dilution shall be made just prior to GC/MS analysis of the sample. All steps in the dilution procedure must be performed without delays until the point at which the diluted sample is in a gas tight syringe.

- 8.1.7.1 The following procedure will allow for dilutions near the calculated dilution factor from the screening procedure:
 - 8.1.7.1.1 All dilutions are made in volumetric flasks (10 mL to 100 mL).
 - 8.1.7.1.2 Select the volumetric flask that will allow for the necessary dilution. Intermediate dilutions may be necessary for extremely large dilutions.
 - 8.1.7.1.3 Calculate the approximate volume of reagent water which will be added to the volumetric flask selected and add slightly less than this quantity of reagent water to the flask.
 - 8.1.7.1.4 Inject the proper aliquot from the syringe prepared in paragraph 8.1.6 into the volumetric flask. Aliquots of less than 1 mL increments are prohibited. Dilute the flask to the mark with reagent water. Cap the flask, invert, and shake three times.
 - 8.1.7.1.5 Fill a 5 mL syringe with the diluted sample as in paragraph 8.1.6.
 - 8.1.7.1.6 If this is an intermediate dilution, use it and repeat the above procedure to achieve larger dilutions.
- 8.1.8 Add 10.0 uL of the system monitoring compound spiking solution (paragraph 5.4.4) and 10.0 uL of the internal standard spiking solution (paragraph 5.4.3) through the valve bore of the syringe, then close the valve. The system monitoring compounds and internal standards may be mixed and added as a single spiking solution. The addition of 10 uL of the system monitoring compound spiking solution to 5 mL of sample is equivalent to a concentration of 50 ug/L of each system monitoring compound.
- 8.1.9 Attach the syringe-syringe valve assembly to the syringe valve on the purging device. Open the syringe valves and inject the sample into the purging chamber.
- 8.1.10 Close both values and purge the sample for 11.0 ± 0.1 minutes at ambient temperature.
- 8.1.11 At the conclusion of the purge time, attach the trap to the chromatograph, adjust the device to the desorb mode, and begin the gas chromatographic temperature program. Concurrently, introduce the trapped materials to the gas chromatographic

column by rapidly heating the trap to 180°C while backflushing the trap with an inert gas between 20 and 60 mL/min for four minutes. If this rapid heating requirement cannot be met, the gas chromatographic column must be used as a secondary trap by cooling it to 30°C (or subambient, if problems persist) instead of the recommended initial temperature of 45°C.

- 8.1.12 While the trap is being desorbed into the gas chromatograph, empty the purging chamber. Wash the chamber with a minimum of two 5 mL flushes of reagent water to avoid carryover of target compounds.
- 8.1.13 After desorbing the sample for four minutes, recondition the trap by returning the purge and trap device to the purge mode. Wait 15 seconds, then close the syringe value on the purging device to begin gas flow through the trap. The trap temperature should be maintained at 180°C. Trap temperatures up to 220°C may be employed, however the higher temperature will shorten the useful life of the trap. After approximately seven minutes, turn off the trap heater and open the syringe value to stop the gas flow through the trap. When cool, the trap is ready for the next sample.
- 8.1.14 Each analytical run must be checked also for saturation. The level at which an individual compound will saturate the detection system is a function of the overall system sensitivity and the mass spectral characteristics of that compound. The initial calibration requires that the system should not be saturated for high response compounds at 200 ug/L for VOA target compounds. Secondary ion quantitation is allowed only when there are sample matrix interferences with the primary ion. If secondary ion quantitation is performed, document the reasons in the SDG Narrative. When a sample is analyzed that has saturated ions from a compound, this analysis must be followed by the analysis of a reagent water blank. If the blank is not free of interferences, the system must be decontaminated. Sample analysis may not resume until a blank has been analyzed that is free of interferences.
- 8.1.15 To prepare a matrix spike and matrix spike duplicate for water samples, add 10 uL of the matrix spike solution (paragraph 5.4.5) to each of the 5 mL aliquots of the sample chosen for spiking. Disregarding any dilutions, this is equivalent to a concentration of 50 ug/L of each matrix spike compound. The frequency of MS/MSD analysis is given in paragraph 10.8.
- 8.1.16 A volatile method blank must be analyzed at least once during every twelve hour time period, on each GC/MS system used for volatile analysis (see paragraph 6.4.6 for the definition of the twelve hour time period).
 - 8.1.16.1 For water samples, a volatile method blank consists of a 5 mL volume of reagent water (paragraph 4.1) spiked with the system monitoring compounds and

internal standards, and carried through the analytical procedure.

- 8.1.16.2 An acceptable volatile method blank for water samples must contain less than or equal to five times (5x) the Contract Required Quantitation Limit (CRQL, see Exhibit C) of Methylene chloride, Acetone, and 2-Butanone, and less than or equal to the CRQL of any other volatile target compound.
- 8.1.16.3 <u>All</u> volatile analyses associated with a blank that does not meet the requirements above, (i.e., a contaminated blank) <u>must</u> be repurged, reanalyzed, and reported at no additional cost to the Agency.
- 8.1.16.4 The volatile method blank <u>must</u> be analyzed <u>after</u> the calibration standards, to ensure that there is no carryover of material from the standards into samples.

8.2 Soil/Sediment Samples

Two approaches may be taken to determine whether the low level or medium level method must be followed.

- Assume the sample is low level and analyze a 5 g sample.
- Use the X factor calculated from the optional hexadecane screen (Section III, paragraph 6.2.1.3) to determine the appropriate method for analysis.

If peaks are saturated from the analysis of a 5 g sample, a smaller sample size must be analyzed to prevent saturation. However, the smallest sample size permitted is 1 g. If smaller than 1 g sample size is needed to prevent saturation, the medium level method <u>must</u> be used.

8.2.1 Low Level Soil Method

The low level soil method is based on purging a heated sediment/soil sample mixed with reagent water containing the system monitoring compounds and the internal standards. Analyze all method blanks and standards under the same conditions as the samples.

Use 5 grams of sample, or use the X Factor to determine the sample size for purging.

- o If the X Factor is 0 (no peaks noted on the hexadecane screen), analyze a 5 g sample.
- o If the X Factor is between 0 and 1.0, analyze a minimum of a 1 g sample.
- 8.2.1.1 The GC/MS system should be set up as in paragraphs 7-7.7.3. This should be done prior to the preparation of the sample to avoid loss of volatiles

from standards and sample. A heated purge calibration curve must be prepared and used for the quantitation of all samples analyzed with the low-level method. Follow the initial and daily calibration instructions (7.4 and 7.7), but increase the purge temperature to 40° C.

- 8.2.1.2 To prepare the reagent water containing the system monitoring compounds and the internal standards, remove the plunger from a 5 mL "Luerlock" type syringe equipped with a syringe valve and fill until overflowing with reagent water. Replace the plunger and compress the water to vent trapped air. Adjust the volume to 5.0 mL. Add 10 uL of the system monitoring compound spiking solution and 10 uL of the internal standard solution to the syringe through the valve.
- 8.2.1.3 The sample (for volatile organics) consists of the <u>entire</u> contents of the sample container. Do not discard any supernatant liquids. Mix the contents of the sample container with a narrow metal spatula. Weigh the amount determined in paragraph 8.2.1 into a tared purge device. Use a top loading balance. Note and record the actual weight to the nearest 0.1 g.
- 8.2.1.4 Immediately after weighing the sample, weigh 5-10 g of the sediment into a tared crucible. Determine the percent moisture by drying overnight at 105°C. Allow to cool in a desiccator before weighing. Concentrations of individual analytes will be reported relative to the dry weight of sediment.

<u>g of wet sample-g of dry sample</u> g of wet sample x 100 = % moisture

- 8.2.1.5 Add the spiked reagent water to the purge device and connect the device to the purge and trap system. NOTE: Prior to the attachment of the purge device, the steps in paragraphs 8.2.1.2 and 8.2.1.3 must be performed rapidly to avoid loss of volatile organics. These steps must be performed in a laboratory free of solvent fumes.
- 8.2.1.6 Heat the sample to $40^{\circ}C \pm 1^{\circ}C$ and purge the sample for 11.0 \pm 0.1 minutes.
- 8.2.1.7 Proceed with the analysis as outlined in paragraphs 8.1.10 - 8.1.13. Requirements for dilution of samples are given in paragraphs 8.2 and 10.7.

- 8.2.1.8 To prepare a matrix spike and matrix spike duplicate for low level soils/sediment, add 10 uL of the matrix spike solution (5.4.4) to the 5 mL of water added to each of the two aliquots of the soil from the sample chosen for spiking (paragraph 8.2.1.2). The concentration for a 5 g sample would be equivalent to 50 ug/kg of each matrix spike compound. The frequency of MS/MSD analysis is given in paragraph 10.8.
- 8.2.1.9 A volatile method blank must be analyzed at least once during every twelve hour time period, on each GC/MS system used for volatile analysis (see paragraph 6.4.6 for the definition of the twelve hour time period).
 - 8.2.1.9.1 For low level soil/sediment samples, a volatile method blank consists of a 5 g of a purified solid matrix added to reagent water, spiked with the system monitoring compounds and internal standards, and carried through the analytical procedure.
 - 8.2.1.9.2 An acceptable volatile method blank for low level soil samples must contain less than or equal to five times (5x) the Contract Required Quantitation Limit (CRQL, see Exhibit C) of Methylene chloride, Acetone, and 2-Butanone, and less than or equal to the CRQL of any other volatile target compound.
 - 8.2.1.9.3 <u>All volatile analyses associated with a blank that does not meet the requirements above, (i.e., a contaminated blank) must be repurged, reanalyzed, and reported at no additional cost to the Agency.</u>
 - 8.2.1.9.4 The volatile method blank <u>must</u> be analyzed <u>after</u> the calibration standards, to ensure that there is no carryover of material from the standards into samples.

8.2.2 Medium Level Soil Method

The medium level soil method is based on extracting the soil/ sediment sample with methanol. An aliquot of the methanol extract is added to reagent water containing the system monitoring compounds and the internal standards. The reagent water containing the methanol extract is purged at ambient

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temperature. All samples with an X Factor >1.0 must be analyzed by the medium level method. If saturated peaks occurred, or would occur, when a 1 g sample was analyzed, the medium level method must be used.

- 8.2.2.1 The GC/MS system should be set up as in paragraphs 7-7.7.3. This should be done prior to the addition of the methanol extract to reagent water. Because the methanol extract and reagent water mixture is purged at <u>ambient</u> temperature, the instrument performance check, initial calibration, and continuing calibration for water samples may be used for analyses of medium soil sample extracts.
- 8.2.2.2 The sample (for volatile organics) consists of the entire contents of the sample container. Do not discard any supernatant liquids. Mix the contents of the sample container with a narrow metal spatula. Weigh 4 g (wet weight) into a tared 15 mL vial. Use a top loading balance. Note and record the actual weight to the nearest 0.1 g. Determine the percent moisture as in paragraph 8.2.1.4.
- 8.2.2.3 Quickly add 9.0 mL of methanol to the vial. Then add 1.0 mL of the system monitoring compound spiking solution to the vial. Cap and shake for 2 minutes. NOTE: The steps in paragraphs 8.2.2.2 and 8.2.2.3 must be performed rapidly to avoid loss of volatile organics. These steps must be performed in a laboratory free of solvent fumes.
- 8.2.2.4 Using a disposable pipette, transfer approximately 1 mL of extract into a GC vial for storage. The remainder may be discarded. Transfer approximately 1 mL of the reagent methanol to a GC vial for use as the method blank for each Case, SDG, or day on which medium soil sample extractions are performed, whichever is most frequent. These extracts may be stored in the dark at 4°C (±2°C) prior to analysis.
- 8.2.2.5 The following table can be used to determine the volume of methanol extract to add to the 5 mL of reagent water for analysis. If the Hexadecane screen procedure was followed, use the X factor (Option B) or the estimated concentration (Option A) to determine the appropriate volume. Otherwise, estimate the concentration range of the sample from the low level analysis to determine the appropriate volume. If the sample was submitted as a medium level sample, start with 100 uL.

All dilutions must keep the response of the major constituents (previously saturated peaks) in the upper half of linear range of the curve.

X Factor	Estimated <u>Concentration Range</u> 1 ug/kg	Take this Volume of <u>Methanol Extract</u> ² uL
0.25 - 5.0 0.5 - 10.0	500 - 10,000 1000 - 20,000	100
2.5 - 50.0 12.5 - 250	5000 - 100,000 25,000 - 500,000	10 100 of 1/50 dilution ³

Calculate appropriate dilution factor for concentrations exceeding the table.

- ¹ Actual concentration ranges could be 10 to 20 times higher than this if the compounds are halogenated and the estimates are from GC/FID.
- ² The volume of methanol added to the 5 mL of water being purged should be kept constant. Therefore, add to the 5 mL syringe whatever volume of methanol is necessary to maintain a volume of 100 uL added to the syringe.
- ³ Dilute an aliquot of the methanol extract and then take 100 uL for analysis.
 - 8.2.2.6 Remove the plunger from a 5 mL "Luerlock" type syringe equipped with a syringe valve and fill until overflowing with reagent water. Replace the plunger and compress the water to vent trapped air. Adjust the volume to 4.9 mL. Full the plunger back to 5 mL to allow volume for the addition of sample and standards. Add 10 uL of the internal standard solution. Also add the volume of methahol extract determined in paragraph 8.2.2.5 and a volume of clean methanol to total 100 uL (excluding methanol in standards).
 - 8.2.2.7 Attach the syringe-syringe valve assembly to the syringe valve on the purging device. Open the syringe valve and inject the water/methanol sample into the purging chamber.
 - 8.2.2.8 Proceed with the analysis as outlined in paragraph
 8. Analyze all method blanks on the same instrument as the samples. Requirements for dilution of samples are given in paragraphs
 8.2 and 10.7:
 - 8.2.2.9 To prepare a matrix spike and matrix spike duplicate for the medium level soil/sediment samples, add 8.0 mL of methanol, 1.0 mL of the system monitoring compound spiking solution, and 1.0 mL of matrix spike solution (paragraph 5.4.4) as in paragraph 8.2.2.3, to each of the two aliquots of the soil sample chosen for spiking. This results in a 6,200 ug/kg concentration of each matrix spike compound when added to a 4 g sample. Add a 100 uL aliquot of this extract to 5 mL of water for purging (as per

paragraph 8.2.2.6). The frequency of MS/MSD analysis is given in paragraph 10.8.

- 8.2.2.10. A volatile method blank must be analyzed at least once during every twelve hour time period, on each GC/MS system used for volatile analysis (see paragraph 6.4.6 for the definition of the twelve hour time period).
 - 8.2.2.10.1 For medium level soil/sediment samples, a volatile method blank consists of a 1 g of a purified solid matrix spiked with the system monitoring compounds, extracted with methanol, and carried through the analytical procedure.
 - 8.2.2.10.2 An acceptable volatile method blank for medium level soil/sediment samples must contain less than or equal five times (5x) the Contract Required Quantitation Limit (CRQL, see Exhibit C) of Methylene chloride, Acetone, and 2-Butanone, and less than or equal ot the CRQL of any other volatile target compound.
 - 8.2.2.10.3 <u>All</u> volatile analyses associated with a blank that does <u>not</u> meet the requirements above; (i.e. a contaminated blank) <u>must</u> be repurged, reanalyzed, and reported at no additional cost to the Agency.
 - 8.2.2.10.4 The volatile method blank <u>must</u> be analyzed <u>after</u> the calibration standards, to ensure that there is no carryover of material from the standards into samples.

9. <u>Qualitative Analysis</u>

- 9.1 The compounds listed in the Target Compound List (TCL), Exhibit C, shall be identified by an analyst competent in the interpretation of mass spectra (see Bidder Responsibility description) by comparison of the sample mass spectrum to the mass spectrum of a standard of the suspected compound. Two criteria must be satisfied to verify the identifications: (1) elution of the sample component at the same GC relative retention time as the standard component, and (2) correspondence of the sample component and standard component mass spectra.
 - 9.1.1 For establishing correspondence of the GC relative retention time (RRT), the sample component RRT must compare within \pm 0.06 RRT units of the RRT of the standard component. For reference,

the standard must be run in the same 12-hour time period as the sample. If coelution of interfering components prohibits

accurate assignment of the sample component RRT from the total ion chromatogram, the RRT should be assigned by using extracted ion current profiles for ions unique to the component of interest.

- 9.1.2 For comparison of standard and sample component mass spectra, mass spectra obtained on the Contractor's GC/MS are required. Once obtained, these standard spectra may be used for identification purposes, <u>only</u> if the Contractor's GC/MS meets the daily instrument performance requirements for BFB. These standard spectra may be obtained from the run used to obtain reference RRTs.
- 9.1.3 The requirements for qualitative verification by comparison of mass spectra are as follows:
 - 9.1.3.1 All ions present in the standard mass spectra at a relative intensity greater than 10% (most abundant ion in the spectrum equals 100%) must be present in the sample spectrum.
 - 9.1.3.2 The relative intensities of ions specified in paragraph 9.1.3.1 must agree within ± 20% between the standard and sample spectra. (Example: For an ion with an abundance of 50% in the standard spectra, the corresponding sample abundance must be between 30 and 70 percent).
 - 9.1.3.3 Ions greater than 10% in the <u>sample</u> spectrum but not present in the standard spectrum must be considered and accounted for by the analyst making the comparison. In Task III, the verification process should favor false positives. All compounds meeting the identification criteria must be reported with their spectra. For all compounds below the CRQL report the actual value followed by a "J", e.g., "3J."
- 9.1.4 If a compound cannot be verified by all of the criteria in paragraph 9.1.3.3, but in the technical judgement of the mass spectral interpretation specialist, the identification is correct, then the Contractor shall report that identification and proceed with quantification in paragraph 10.
- 9.2 A library search shall be executed for non-target sample components for the purpose of tentative identification. For this purpose, the 1989 (or more recent) release of the NIST/EPA/MSDC mass spectral library, containing 50,000 spectra, shall be used. Computer generated library search routines must not use normalization routines that would misrepresent the library or unknown spectra when compared to each other.

- 9.2.1 Up to 10 organic compounds of greatest apparent concentration not listed in Exhibit C for the purgeable organic fraction, excluding the system monitoring compounds, shall be tentatively identified via a forward search of the NIST/EPA/MSDC Library (substances with responses less than 10% of the internal standard are not required to be searched in this fashion). Only after visual comparison of sample spectra with the nearest library searches will the mass spectral interpretation specialist assign a tentative identification. Computer generated library search routines must not use normalization routines that would misrepresent the library or unknown spectra when compared to each other.
- 9.2.2 Guidelines for making tentative identification:
 - 9.2.2.1 Relative intensities of major ions in the reference spectrum (ions greater than 10% of the most abundant ion) should be present in the sample spectrum.
 - 9.2.2.2 The relative intensities of the major ions should agree within ± 20%. (Example: For an ion with an abundance of 50 percent of the standard spectra, the corresponding sample ion abundance must be between 30 and 70 percent.)
 - 9.2.2.3 Molecular ions present in reference spectrum should be present in sample spectrum.
 - 9.2.2.4 Ions present in the sample spectrum but not in the reference spectrum should be reviewed for possible background contamination or presence of co-eluting compounds.
 - 9.2.2.5 Ions present in the reference spectrum but not in the sample spectrum should be reviewed for possible subtraction from the sample spectrum because of background contamination or co-eluting compounds. Data system library reduction programs can sometimes create these discrepancies.
- 9.2.3 If, in the technical judgement of the mass spectral interpretation specialist, no valid tentative identification can be made, the compound should be reported as <u>unknown</u>. The mass spectral specialist should give additional classification of the unknown compound, if possible (i.e., unknown aromatic, unknown hydrocarbon, unknown acid type, unknown chlorinated compound). If probable molecular weights can be distinguished, include them.

10. <u>Quantitative Analysis</u>

10.1 Target components identified shall be quantified by the internal standard method. The internal standard used shall be that which is assigned in Table 5 of this Section. The EICP area of the

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characteristic ions of analytes listed in Tables 3 and 4 in this Section are used.

- 10.2 Internal standard responses and retention times in all standards must be evaluated during or immediately after data acquisition. If the retention time for any internal standard changes by more than 30 seconds from the latest daily (12 hour) calibration standard, the chromatographic system must be inspected for malfunctions, and corrections made as required. The extracted ion current profile (EICP) of the internal standards must be monitored and evaluated for each sample, blank, matrix spike, and matrix spike duplicate. If the EICP area for any internal standard changes by more than a factor of two (-50% to +100%), the mass spectrometric system must be inspected for malfunction and corrections made as appropriate. When corrections are made, reanalysis of samples analyzed while the system was malfunctioning is necessary.
 - 10.2.1 If after re-analysis, the EICP areas for all internal standards are inside the contract limits (-50% to +100%), then the problem with the first analysis is considered to have been within the control of the laboratory. Therefore, <u>submit only</u> data from the analysis with EICPs within the contract limits. This is considered the <u>initial</u> analysis and must be reported as such on all data deliverables.
 - 10.2.2 If the re-analysis of the sample does not solve the problem, i.e., the EICP areas are outside the contract limits for both analyses, then submit the EICP data and sample data from both analyses. Distinguish between the initial analysis and the re-analysis on all data deliverables, using the sample suffixes specified in Exhibit B. Document in the SDG Narrative all inspection and corrective actions taken.

10.3 The relative response factor (RRF) from the continuing calibration standard is used to calculate the concentration in the sample. Use the relative response factor as determined in paragraph 7.4.3 and the equations below. When target compounds are below contract required quantitation limits (CRQL), but the spectra meet the identification criteria, report the concentration with a "J." For example, if CRQL is 10 ug/L and concentration of 3 ug/L is calculated, report as "3J."

Water

Concentration

$$ug/L = \frac{(A_{\chi})(I_{s})(Df)}{(A_{is})(RRF)(V_{o})}$$

Where

- A_x Area of the characteristic ion (EICP) for the compound to be measured (see Table 4)
- A_{is} Area of the characteristic ion (EICP) for the specific internal standard (see Table 3)
- I = Amount of internal standard added in nanograms (ng)
- RRF = Relative response factor from the <u>ambient</u> temperature purge of the calibration standard.
- Vo Volume of water purged in milliliters (mL)
- Df Dilution factor. The dilution factor for analysis of water samples for volatiles by this method is defined as the ratio of the number of milliliters (mL) of water purged (i.e., V_0 above) to the number of mL of the original water sample used for purging. For example, if 2.5 mL of sample is diluted to 5.0 mL with reagent water and purged, Df = 5.0 mL/2.5 mL = 2.0. If no dilution is performed, Df = 1.0.

Low Soil

Concentration
(Dry weight basis)
$$ug/Kg = \frac{(A_x)(I_s)}{(A_{is})(RRF)(W_s)(D)}$$

Where

A_x, I_s, A_{is} RRFs are as given for water.

 W_s - Weight of sample added to the purge tube, in grams (g)

Medium Soil

Concentration (Dry weight basis) $ug/Kg = \frac{(A_{\chi})(I_{s})(V_{t})(1000)(Df)}{(A_{1s})(RRF)(V_{a})(W_{s})(D)}$

Where

A., Ais, Is, RRFs are as given for water above.

- Vt = Total volume of the methanol extract in milliliters (mL). NOTE: This volume is typically 10.0 mL, even though only 1.0 mL is transferred to the vial in paragraph 8.2.2.4.
- V_a = Volume of the aliquot of the methanol extract in microliters (uL) added to reagent water for purging

W_g = Weight of soil extracted, in grams (g)

Df - Dilution factor. The dilution factor for analysis of soil/sediment samples for volatiles by the <u>medium level</u> method is defined as the ratio of the number of microliters (uL) of methanol added to the reagent water for purging i.e., V_a above, to the number of microliters of the methanol <u>extract</u> of the sample contained in that volume V_a . The dilution factor is equal to 1.0 in all cases other than those requiring dilution of the methanol extract. Dilution of the extract is required when the "X" factor (paragraph 8.2.2.5) is ≥ 12.5 .

The factor of 1,000 in the numerator converts the value of V_t from mL to uL.

10.4 An estimated concentration for non-target components tentatively identified shall be determined by the internal standard method. For quantification, the nearest internal standard <u>free of interferences</u> shall be used.

The formula for calculating concentrations is the same as in paragraph 10.3. Total area counts (or peak heights) from the total ion chromatograms are to be used for both the compound to be measured and the internal standard. A relative response factor (RRF) of one (1) is to be assumed. The resulting concentration shall be qualified as "J" (estimated, due to lack of a compound-specific response factor), and "N" (presumptive evidence of presence), indicating the quantitative and qualitative uncertainties associated with this non-target component. An estimated concentration should be calculated for all tentatively identified compounds as well as those identified as unknowns.

10.5 Xylenes (o-,m-, & p- isomers) are to be reported as Xylenes (total). Since o- and p-Xylene overlap, the Xylenes must be quantitated as m-Xylene. The concentration of all Xylene isomers must be added together to give the total.

- 10.6 Both trans and cis stereoisomers of 1,2-Dichloroethene are to be reported as 1,2-Dichloroethene (total). The concentrations of both isomers must be added together to give the total.
- 10.7 If the on-column concentration of any compound in any sample exceeds the initial calibration range, that sample must be diluted, the internal standard concentration re-adjusted, and the sample repurged. Guidance in performing dilutions, and <u>exceptions</u> to this requirement are given below.
 - 10.7.1 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
 - 10.7.2 The dilution factor chosen should keep the response of the largest analyte peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.
 - 10.7.3 Do <u>not</u> submit data for more than two analyses, i.e., the original sample and <u>one</u> dilution, or, if the volatile screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
 - 10.7.4 Do not dilute MS/MSD samples solely to get non-spiked analytes within calibration range. The spiking level of each compound in the volatile matrix spiking solution should not require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.
 - 10.7.5 For total Xylenes, where three isomers are quantified as two peaks, the calibration of <u>each peak</u>, should be considered separately, i.e., a diluted analysis is <u>not</u> required for total Xylenes unless the concentration of either peak separately exceeds 200 ug/L.
- 10.8 Calculate the recovery of each system monitoring compound in all samples, blanks, matrix spikes, and matrix spike duplicates. Determine if the recovery is within limits (see Table 6), and report on appropriate form.
 - 10.8.1 Calculate the concentrations of the system monitoring compounds using the same equations as used for target compounds. Calculate the recovery of each system monitoring compound as follows:
 - %R = Concentration (or amount) found x 100 Concentration (or amount) spiked

- 10.8.2 If the recovery of any one system monitoring compound is not within limits, the following are required:
 - Check to be sure that there are no errors in calculations, formulation of the system monitoring compound spiking solutions, and internal standards. Also check instrument performance.
 - Reanalyze the sample if none of the above steps reveal a problem.
 - Do not reanalyze diluted samples if the system monitoring compound recoveries are outside the limits.
 - Never reanalyze the matrix spike or matrix spike duplicate (MS/MSD), even if the system monitoring compound recoveries are outside the limits.
 - If the sample associated with the matrix spike and matrix spike duplicate does not meet specifications, it should be reanalyzed only if the MS/MSD system monitoring compound recoveries are within the limits. If the sample and associated MS/MSD show the same pattern (i.e., outside the limits), then the sample does not require reanalysis and a reanalysis must not be submitted. Document in the narrative the similarity in recoveries of the system monitoring compounds in the sample and associated MS/MSD.
- 10.8.3 If the reanalysis of the sample solves the problem, then the problem was within the laboratory's control. Therefore, submit only data from the analysis with system monitoring compound recoveries within the limits. This shall be considered the initial analysis and shall be reported as such on all data deliverables.
- 10.8.4 If the reanalysis of the sample does not solve the problem (i.e., the system monitoring compound recoveries are outside the limits for both analyses), then submit the data from both analyses. Distinguish between the initial analysis and the reanalysis on all data deliverables, using the sample suffixes supplied in Exhibit B.
- 10.8.5 For medium level soil analyses, involving methanol extraction, the treatment of system monitoring compound recoveries is similar to that for semivolatile surrogate recoveries. If any system monitoring compound recovery is outside the limits, reanalyze the methanol extract first, to determine if the problem was with the analysis. If reanalysis of the extract does not solve the problem, then reextract the medium soil sample and analyze the second extract. Follow paragraphs 10.8.3 and 10.8.4 when determining which analyses to submit.
- 10.8.6 If the recovery of any one system monitoring compound in a method blank is outside the limits, then the method <u>and</u> all associated samples must be reanalyzed at no additional cost to the Agency.

- 10.9 A matrix spike and matrix spike duplicate must be performed for each group of samples of a similar matrix, for the following, whichever is most frequent:
 - o Each Case of field samples received, OR
 - o Each 20 field samples in a Case, OR
 - o Each group of field samples of a similar concentration level (soils only), OR
 - o Each 14 calendar day period during which field samples in a Case were received (said period beginning with the receipt of the first sample in that Sample Delivery Group).
 - 10.9.1 Calculate the concentrations of the matrix spike compounds using the same equations as used for target compounds. Calculate the recovery of each matrix spike compound as follows:
 - %R = <u>Concentration (or amount) found</u> x 100 Concentration (or amount) spiked
 - 10.9.2 Calculate the relative percent difference (RPD) of the recoveries of each compound in the matrix spike and matrix spike duplicate as follows:
 - $\frac{\text{RPD} |\text{MSR} \text{MSDR}|}{(1/2)(\text{MSR}+\text{MSDR})} \times 100$

Where

MSR - Matrix Spike Recovery

MSDR - Matrix Spike Duplicate Recovery

The vertical bars in the formula above indicate the absolute value of the difference, hence RPD is always expressed as a positive value.

- 10.9.3 The limits for matrix spike compound recovery and RPD are given in Table 7. As these limits are only advisory, no further action by the laboratory is required, however, frequent failures to meet the limits for recovery or RPD warrant investigation by the laboratory, and may result in questions from the Agency.
- 10.10 Determine the concentrations of any target compounds detected in the volatile method blank, using the equations in paragraph 10.3. The method blank must contain less than or equal to the Contract Required Quantitation Limit (CRQL) of the volatile target compounds in Exhibit C, except Methylene chloride, Acetone, and 2-Butanone, which must be less than or equal to five times (5x) the CRQL. For soil/sediment method blanks, CRQL value must be adjusted for percent moisture (see Exhibit B).

If a laboratory method blank exceeds these criteria, the Contractor must consider the analytical system to be out of control. The source of the contamination must be investigated and appropriate corrective measures <u>MUST</u> be taken and documented before further sample analysis proceeds. All samples processed with a method blank that is out of control (i.e., contaminated) <u>MUST</u> be reextracted/repurged and reanalyzed at no additional cost to the Agency. The Laboratory Manager, or his designee, must address problems and solutions in the SDG Narrative (Exhibit B).

TABLE 3 CHARACTERISTIC IONS FOR SYSTEM MONITORING COMPOUNDS AND INTERNAL STANDARDS FOR VOLATILE ORGANIC COMPOUNDS

Compound	Primary Ion	Secondary Ion(s)
SYSTEM MONITORING COMPOUNDS		
4-Bromofluorobenzene	95	174, 176
1,2-Dichloroethane-d-4	65	102
Toluens-d-8	98	70, 100
INTERNAL STANDARDS		
Bromochloromethane	128	49, 130, 51
1,4-Difluorobenzene	114	63, 88
Chlorobenzene-d-5	117	82, 119

			SLE 4		
CHARACTERISTIC	IONS	FOR	VOLATILE	TARGET	COMPOUNDS

		Secondary
Analyte	Primary Ion*	Ion(s)
Chloromethane	50	52
Bromomethane	94	96
	62	64
Vinyl chloride	64	66
Chloroethane	84	49, 51, 86
Methylene chloride	43	58
	76	78
Carbon disulfide	96	61, 98
1,1-Dichloroethene	63	65, 83, 85, 98, 100
1,1-Dichloroethane	96	61, 98
1,2-Dichlorosthene	83	85
Chloroform	62	64, 100, 98
1,2-Dichloroethane	62 43**	57
2-Butanone		99 , 117, 119
1,1,1-Trichloroethane	97	119, 121
Carbon tetrachloride	117	85
Bromodichloromethane	83	85, 131, 133, 166
1,1,2,2-Tetrachloroethane	83	65, 114
1,2-Dichloropropane	63	77
trans-1,3-Dichloropropene	75	95, 97, 132
Trichloroethene	130	
Dibromochloromethane	129	208, 206
1,1,2-Trichloroethane	97	83, 85, 99, 132, 134
Benzene	78	77
cis-1,3-Dichloropropene	75	77
Bromoform	173	171, 175, 250, 252, 254, 256
2-Hexanone	43	58, 57, 100
4-Methyl-2-pentanone	43	58, 100
Tetrachloroethene	164	129, 131, 166
Toluene	91	92
Chlorobenzene	112	114
Ethyl benzene	106	91
Styrene	104	78, 103
Total Xylenes	106	91

* The primary ion should be used unless interferences are present, in which case, a secondary ion may be used.

** m/z 43 is used for quantitation of 2-Butanone, but m/z 72 must be present for positive identification.

-

TABLE 5

VOLATILE INTERNAL STANDARDS WITH CORRESPONDING TARGET COMPOUNDS AND SYSTEM MONITORING COMPOUNDS ASSIGNED FOR QUANTITATION

Bromochloromethane	1,4-Difluorobenzene	Chlorobenzene-d ₅
Chloromethane Bromomethane Vinyl Chloride Chloroethane Methylene Chloride Acetone Carbon Disulfide 1,1-Dichloroethene 1,2-Dichloroethene(tot.) Chloroform 1,2-Dichloroethene 2-Butanone 1,2-Dichloroethane 2-Butanone 1,2-Dichloroethane-d ₄ (smc)	1,1,1-Trichlorosthane Carbon Tetrachloride Bromodichloromethane 1,2-Dichloropropane trans-1,3-Dichloropropene Trichlorosthene Dibromochloromethane 1,1,2-Trichlorosthane Benzene cis-1,3-Dichloropropene Bromoform	2-Hexanone 4-Methyl-2-Pentanone Tetrachloroethene 1,1,2,2-Tetrachloroethane Toluene Chlorobenzene Ethylbenzene Styrene Xylene (total) Bromofluorobenzene (smc) Toluene-d ₈ (smc)

(smc) - system monitoring compound

TABLE 6 SYSTEM MONITORING COMPOUND RECOVERY LIMITS

	\$R	%R
Compound	Water	Soil
Toluene-dg	88-110	84-138
Bromofluorobenzene	86-115	59-113
1,2-Dichloroethane-d4	76-114	70-121

TABLE 7MATRIX SPIKE RECOVERY ANDRELATIVE PERCENT DIFFERENCE LIMITS

Compound	SR Water	RPD Water	%R Soil	RPD Soil
1,1-Dichloroethane	61-145	14	59-172	22
Trichloroethene	71-120	14	62-137	24
Benzene	76-127	11	66-142	21
Toluene	76-125	13	59-139	21
Chlorobenzene	75-130	13	60-133	21

EXHIBIT D

ANALYTICAL METHODS FOR SEMIVOLATILES

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SECTION I

INTRODUCTION

The analytical methods that follow are designed to analyze water, soil and sediment from hazardous waste sites for the organic compounds on the Target Compound List (TCL, see Exhibit C). The methods are based on EPA Method 625 (Bases/Neutrals and Acids).

The methods are divided into the following sections: sample preparation, screening, and analysis. Sample preparation covers sample extraction and cleanup techniques. As described in the screening section, a portion of the extracts may be screened on a gas chromatograph with appropriate detectors to determine the concentration level of organics. The analysis section contains the GC/MS analytical methods for organics.

1. <u>Method for the Determination of Extractable Semivolatile Organic</u> <u>Compounds</u>.

1.1 Scope and Application

This method covers the determination of a number of organic compounds that are partitioned into an organic solvent and are amenable to gas chromatography. These target compounds and the contract required quantitation limits are listed in Exhibit C.

Problems have been associated with the following compounds analyzed by this method:

- o Dichlorobenzidine and 4-Chloroaniline can be subject to oxidative losses during solvent concentration.
- Hexachlorocyclopentadiene is subject to thermal decomposition in the inlet of the gas chromatograph, chemical reactions in acetone solution, and photochemical decomposition.
- o N-Nitrosodiphenylamine decomposes in the gas chromatograph inlet forming diphenylamine and, consequently, may be detected as diphenylamine.
- o Benzoic acid exhibits poor extraction recoveries.
- 1.2 Because performance data indicate erratic and poor linearity and sensitivity, the following compounds have Maximum %RSD, or Maximum %Difference criteria; however, these compounds <u>must</u> meet a minimum RRF criterion of 0.010:

2,2'-oxybis(1-Chloropropane) 4-Chloroaniline Hexachlorobutadiene Hexachlorocyclopentadiene 2-Nitroaniline Dimethylphthalate 3-Nitroaniline 2,4-Dinitrophenol 4-Nitrophenol Diethylphthalate 4-Nitroaniline 4,6-Dinitro-2-methylphenol N-Nitrosodiphenylamine Di-n-butylphthalate Butylbenzylphthalate 3,3'-Dichlorobenzidine bis(2-Ethylhexyl)phthalate Di-n-octylphthalate 2,4,6-Tribromophenol

In addition, Carbazole, a new target compound for which there is insufficient performance data, must only meet a minimum RRF criterion of 0.010.

1.3 The method involves solvent extraction of the matrix sample, characterization to determine the appropriate analytical protocol to be used, and GC/MS analysis to determine semivolatile organic compounds present in the sample.

SECTION II

SAMPLE PREPARATION AND STORAGE

PART A - SAMPLE STORAGE AND HOLDING TIMES

1. <u>Procedures for Sample Storage</u>

The samples must be protected from light and refrigerated at 4°C (\pm 2°C) from the time of receipt until 60 days after delivery of a complete data package to the Agency. After 60 days the samples may be disposed of in a manner that complies with all applicable regulations.

Samples must be stored in an atmosphere demonstrated to be free of all potential contaminants.

Samples, sample extracts, and standards must be stored separately.

2. Procedure for Sample Extract Storage

Sample extracts must be protected from light and stored at $4^{\circ}C$ ($\pm 2^{\circ}C$) until 365 days after delivery of a complete data package to the Agency.

Samples must be stored in an atmosphere demonstrated to be free of all potential contaminants.

Samples, sample extracts, and standards must be stored separately.

3. Contract Required Holding Times

Extraction of water samples by continuous liquid-liquid procedures shall be started within 5 days of VTSR (Validated Time of Sample Receipt). Extraction of soil/sediment samples by sonication procedures shall be completed within 10 days of VTSR. NOTE: Separatory funnel extraction procedures are <u>not</u> permitted.

Extracts of either water or soil/sediment samples must be analyzed within 40 days following extraction.

PART B - SAMPLE PREPARATION FOR EXTRACTABLE SEMIVOLATILES (BNA) IN WATER

1. <u>Summary of Sample Preparation Method</u>

1.1 A one liter aliquot of sample is acidified to pH 2 and extracted with methylene chloride using a continuous liquid-liquid extractor. Separatory funnel extraction is NOT permitted. The methylene chloride extract is dried and concentrated to a volume of 1.0 mL.

2. Interferences

Contaminants in solvents, reagents, glassware, and other sample processing hardware may cause method interferences such as discrete artifacts and/or elevated baselines in the total ion current profiles (TICPs). All of these materials routinely must be demonstrated to be free from interferences under the conditions of the analysis by running laboratory method blanks. Matrix interferences may be caused by contaminants that are coextracted from the sample. The extent of matrix interferences will vary considerably from source to source.

3. Apparatus and Materials

- 3.1 Glassware (brand names and catalog numbers are included for illustration purposes only).
 - 3.1.1 Continuous liquid-liquid extractors equipped with Teflon or glass commecting joints and stopcocks requiring no lubrication (Hershberg-Wolf Extractor, Ace Glass Company, Vineland, NJ, P/N 6841-10 or equivalent).
 - 3.1.2 Drying column 19 mm ID chromatographic column with coarse frit (substitution of a small pad of Pyrex glass wool for the frit will prevent cross contamination of sample extracts).
 - 3.1.3 Concentrator tube Kuderna-Danish, 10 mL, graduated (Kontes K-570050-1025 or equivalent). Calibration must be checked at the volumes employed in the test. Ground-glass stoppers are used to prevent evaporation of extracts.
 - 3.1.4 Evaporative flask Kuderna-Danish, 500 mL (Kontes K-570001-0500 or equivalent). Attach to concentrator tube with springs.
 - 3.1.5 Snyder column Kuderna-Danish, three-ball macro (Kontes K-503000-0121 or equivalent).
 - 3.1.6 Snyder column Kuderna-Danish, two-ball micro (Kontes K569001-0219 or equivalent).
 - 3.1.7 Vials amber glass, 2 mL capacity with Teflon-lined screw cap.
- 3.2 Silicon carbide boiling chips approximately 10/40 mesh. Heat to 400°C for 30 minutes or Soxhlet extract with methylene chloride.

- 3.3 Water bath heated, with concentric ring cover, capable of temperature control (± 2°C). The bath should be used in a hood.
- 3.4 Balance analytical, capable of accurately weighing \pm 0.0001 g.
- 3.5 Nitrogen evaporation device equipped with a water bath that can be maintained at 35-40°C. The N-Evap by Organomation Associates, Inc., South Berlin, MA (or equivalent), is suitable.

4. <u>Reagents</u>

- 4.1 Reagent water defined as water in which an interferent is not observed at or above the CRQL of each parameter of interest.
- 4.2 Sodium thiosulfate (ACS) granular.
- 4.3 Sulfuric acid solution (1+1) slowly add 50 mL of H₂SO₄ (sp gr 1.84) to 50 mL of reagent water.
- 4.4 Acetone, methanol, methylene chloride pesticide residue analysis grade or equivalent.
- 4.5 Sodium sulfate (ACS) powdered, anhydrous. Purify by heating at 400°C for four hours in a shallow tray, cool in a desiccator and store in a glass bottle (Baker anhydrous powder, catalog #73898, or equivalent).
- 4.6 Surrogate standard spiking solution.
 - 4.6.1 Surrogate standards are added to all samples and calibration solutions; the compounds specified for this purpose are Phenol-d₅, 2,4,6-Tribromophenol, 2-Fluorophenol, Nitrobenzene-d₅, Terphenyl-d₁₄, 2-Fluorobiphenyl, 2-Chlorophenol-d₄, and 1,2-Dichlorobenzene-d₄. Additional surrogates may be added at the laboratory's discretion.
 - 4.6.2 Prepare a surrogate standard spiking solution that contains Nitrobenzene-d₅, Terphenyl-d₁₄, 2-Fluorobiphenyl, and 1,2-Dichlorobenzene-d₄ at a concentration of 100 ug/mL; Phenol-d₅, 2,4,6-Tribromophenol, 2-Fluorophenol, and 2-Chlorophenol-d₄ at a concentration of 200 ug/mL. Store the spiking solutions at $4^{\circ}C$ ($\pm 2^{\circ}C$) in Teflon-sealed containers. The solutions should be checked frequently for stability. These solutions must be replaced after twelve months, or sooner if comparison with quality control check samples indicates a problem.

4.7 BNA Matrix standard spiking solution - the matrix spike solution consists of the following:

Bases/Neutrals	Acids
1,2,4-Trichlorobenzene	Pentachlorophenol
Acenaphthene	Phenol
2,4-Dinitrotoluene	2-Chlorophenol
Pyrene	4-Chloro-3-methylphenol
N-Nitroso-di-n-propylamine	4-Nitrophenol
1,4-Dichlorobenzene	-

Prepare a spiking solution that contains each of the base/neutral compounds above at 100 ug/1.0 mL in methanol and the acid compounds at 200 ug/1.0 ml in methanol. Analyze duplicate aliquots of a sample spiked with BNA matrix spiking solution.

- 5. <u>Water Sample Extraction</u>
- 5.1 Continuous liquid-liquid extraction is used to extract the samples.
 - 5.1.1 Add methylene chloride to the bottom of the extractor and fill it to a depth of at least 1 inch above the bottom side arm.
 - 5.1.2 Using a 1 liter graduated cylinder, measure out a 1.0 liter sample aliquot. Transfer the 1 liter sample aliquot to the continuous extractor. Pipet 0.5 mL of surrogate standard spiking solution into the sample and mix well. Check the pH of the sample with wide range pH paper and adjust the pH to 2.0 with 1:1 H₂SO₄.
 - 5.1.3 Following the procedures in 5.1.1 and 5.1.2 above, prepare two additional 1.0 Liter aliquots of the sample chosen for spiking. Add 0.5 mL of the BNA Matrix Spiking Solution to each of the additional aliquots. The frequency of MS/MSD analysis is given in Section IV, paragraph 8.6.
 - 5.1.4 Add 500 mL of methylene chloride to the distilling flask. Add sufficient reagent water to ensure proper operation. Extract for 18 hours. Allow to cool, then detach the distilling flask, and label the flask.
 - 5.1.5 Prepare a method blank with each group of water samples extracted. For semivolatile analyses, a method blank for water samples consists of a 1 L volume of reagent water (see paragraph 4.1), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section IV, paragraph 8.7
- 5.2 Concentrating the Extracts
 - 5.2.1 Assemble a Kuderna-Danish (K-D) concentrator by attaching a 10 mL concentrator tube to a 500 mL evaporative flask. Other concentration devices or techniques may be used in place of the

K-D, if equivalency is demonstrated for all the semivolatile target compounds listed in Exhibit C.

- 5.2.2 Transfer the extract by pouring the extract through a drying column containing about 10 cm of anhydrous granular sodium sulfate, and collect the extract in a K-D concentrator. Rinse the Erlenmeyer flask and column with 20 to 30 mL of methylene chloride to complete the quantitative transfer.
- 5.2.3 Add one or two clean boiling chips and attach a three-ball Snyder column to the evaporative flask. Pre-wet the Snyder column by adding about 1 mL methylene chloride to the top of the column. Place the K-D apparatus on a hot water bath (60°C to 80°C) so that the concentrator tube is partially immersed in the hot water, and the entire lower rounded surface of the flask is bathed with hot vapor. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 10 to 15 minutes. At the proper rate of distillation, the balls of the column will chatter actively, but the chambers will not flood with condensed solvent. When the apparent volume of liquid reaches 1 mL, remove the K-D apparatus from the water bath and allow it to drain and cool for at least 10 minutes. Remove the Snyder column and rinse the flask and its lower joint into the concentrator tube with 1 to 2 mL of methylene chloride. A 5 mL syringe is recommended for this operation.
- 5.2.4 Two different concentration techniques are permitted to obtain the final 1.0 mL volume: micro Snyder column and nitrogen blowdown techniques.
 - 5.2.4.1 Micro Snyder Column Technique

Add another one or two clean boiling chips to the concentrator tube and attach a two-ball micro Snyder column. Pre-wet the Snyder column by adding about 0.5 mL of methylene chloride to the top of the column. Place the K-D apparatus on a hot water bath (60°C to 80°C) so that the concentrator tube is partially immersed in the hot water. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 5 to 10 minutes. At the proper rate of distillation, the balls of the column will chatter actively, but the chambers will not flood with condensed solvent. When the apparent volume of liquid reaches about 0.5 mL, remove the K-D apparatus from the water bath and allow it to drain for at least 10 minutes while cooling. Remove the Snyder column and rinse its flask and its lower joint into the concentrator tube with 0.2 mL of methylene chloride. Adjust the final volume to 1.0 mL with methylene chloride. Transfer the extract to a Teflon-sealed screw-cap bottle, label the bottle, and store at 4°C (± 2 °C).

5.2.4.2 Nitrogen Blowdown Technique (taken from ASTM Method D3086)

Place the concentrator tube with an open micro Snyder attached in a warm water bath (30°C to 35°C) and evaporate the solvent volume to just below 1 mL by blowing a gentle stream of clean, dry nitrogen (filtered through a column of activated carbon) above the extract. CAUTION: Gas lines from the gas source to the blowdown apparatus must be stainless steel, copper, or Teflon tubing. The internal wall of the concentrator tube must be rinsed down several times with methylene chloride during the operation and the final volume brought to 1.0 mL with methylene chloride. During evaporation, the tube solvent level must be kept below the water level of the bath. The extract must never be allowed to become dry. Transfer the extract to a Teflon-sealed screw-cap bottle, label the bottle and store at 4°C (±2°C).

6. The sample extracts are ready for GC/MS analysis. Proceed to Section IV, GC/MS Analysis of Semivolatiles. If high concentrations are suspected (e.g., highly colored extracts), the optional GC/FID screen in Section III is recommended.

<u>PART C</u> - <u>SAMPLE PREPARATION FOR EXTRACTABLE SEMIVOLATILES (BNA) IN</u> <u>SOIL/SEDIMENT</u>

It is mandatory that all soil/sediment samples be characterized as to concentration level so that the appropriate analytical protocol is chosen to ensure proper quantitation limits for the sample. Note that the terms "low level" and "medium level" are not used here as a judgement of degree of contamination but rather as a description of the concentration ranges that are encompassed by the "low" and "medium" level procedures.

The laboratory is at liberty to determine the method of characterization. The following two screening methods may be used for soil/sediment sample characterization:

- Screen an aliquot from the "low level" 30 g extract or an aliquot from the "medium level" 1 g extract.
- o Screen using either GC/FID or GC/MS as the screening instrument.

The concentration ranges covered by these two procedures may be considered to be approximately 330 ug/kg - 10,000 ug/kg for the low level analysis and >10,000 ug/kg for medium level analysis for semivolatile extractables.

Screen from the Medium Level Method

Take 5.0 mL from the 10.0 mL total extract and concentrate to 1.0 mL and screen. If the sample concentration is >10,000 ug/kg, proceed with GC/MS analysis of the organics. If the sample concentration is <10,000 ug/kg, discard the medium level extract and follow the low level method.

Screen from the Low Level Method

Take 5.0 mL from the 300 mL (approximate) total extract from the 30 g sample and concentrate to 1.0 mL and screen. If the original sample concentration is >10,000 ug/kg, discard the 30 g extract and follow the medium level methods for organics, using medium level surrogates. If the sample concentration is <10,000 ug/kg, proceed with concentration and the remainder of the low level method.

Mandatory GPC Clean Up

Regardless of the concentration level, all soil/sediment sample extracts must be subjected to clean up by Gel Permeation Chromatography (GPC). Because the effectiveness of GPC can be adversely affected by the amount of material loaded onto the GPC column, it may be advisable to screen the sample extracts described here prior to employing GPC.

1. Medium Level Preparation for Screening and Analysis of Semivolatiles

1.1 Scope and Application

This procedure is designed for the preparation of sediment/soil samples which may contain organic chemicals at a level greater than 20,000 ug/kg.

- 1.1.1. The extracts and sample aliquots prepared using this method are screened by GC/MS or FID, using capillary columns for semivolatile priority pollutants, and related organic chemicals. The results of these screens will determine whether sufficient quantities of pollutants are present to warrant analysis by low or medium protocol.
- 1.1.2 If the screenings indicate no detectable pollutants at the lower limits of quantitation, the sample should be prepared by the low level protocol in Section II, Part C, beginning at paragraph 2.

1.2 Summary of Method

- 1.2.1. Approximately 1 g portions of sediment/soil are transferred to vials and extracted with methylene chloride. The methylene chloride extract is screened for extractable organics by GC/FID or GC/MS.
- 1.2.2 If organic compounds are detected by the screen, the methylene chloride extract is subjected to GPC clean up and analyzed by GC/MS for extractable organics.
- 1.2.3 If no organic compounds are detected by the medium level screen, then a low level sample preparation is required.

1.3 Interferences

1.3.1. Method interferences may be caused by contaminants in solvents, reagents, glassware, and other sample processing hardware that lead to discrete artifacts and/or elevated baselines in the total ion current profiles. All of these materials routinely must be demonstrated to be free from interferences under the conditions of the analysis by running laboratory reagent blanks. Matrix interferences may be caused by contaminants that are coextracted from the sample. The extent of matrix interferences will vary considerably from source to source.

1.4 Limitations

1.4.1. The procedure is designed to allow quantitation limits for screening purposes as low as 20,000 ug/kg for extractable organics. For analysis purposes, the quantitation limits are 20,000 ug/kg for extractable organics. If peaks are present based on the GC/FID screen, the sample is determined to require a medium level analysis by GC/MS. Some samples may contain

high concentrations of chemicals that interfere with the analysis of other components at lower levels; the quantitation limits in those cases may be significantly higher.

1.4.2 These extraction and preparation procedures were developed for rapid and safe handling of high concentration hazardous waste samples. The design of the methods thus does not stress efficient recoveries or low limits of quantitation of all components. Rather, the procedures were designed to screen, at moderate recovery and sufficient sensitivity, a broad spectrum of organic chemicals. The results of the analyses thus may reflect only a minimum of the amount actually present in some samples.

1.5 Reagents

- 1.5.1. Sodium Sulfate anhydrous powdered reagent grade, heated at 400°C for four hours, cooled in a desiccator, and stored in a glass bottle (Baker anhydrous powder, catalog # 73898 or equivalent).
- 1.5.2 Acetone, Methanol, Methylene chloride pesticide residue analysis grade or equivalent.
- 1.5.3 Base/Neutral and Acid Surrogate Spiking Solution

Surrogate standards are added to all samples and calibration solutions. The compounds specified are Phenol-d₅, 2,4,6-Tribromophenol, 2-Fluorophenol, Nitrobenzene-d₅, Terphenyl-d₁₄, 2-Fluorobiphenyl, 2-Chlorophenol-d₄, and 1,2-Dichlorobenzene-d₄. Prepare a surrogate standard spiking solution that contains Nitrobenzene-d₅, Terphenyl-d₁₄, 2-Fluorobiphenyl, and 1,2-Dichlorobenzene-d₄ at a concentration of 100 ug/mL; Phenol-d₅, 2,4,6-Tribromophenol, 2-Fluorophenol, and 2-Chlorophenol-d₄ at a concentration of 200 ug/mL. Store the spiking solutions at 4°C (\pm 2°C) in Teflon-sealed containers. The solutions must be replaced after twelve months, or sooner if comparison with quality control check samples indicates a problem.

1.5.4 Base/Neutral and Acid Matrix Spiking solution

Prepare a spiking solution in methanol that contains the following compounds at a concentration of 100 ug/mL for base/neutrals and 200 ug/mL for acids. Store the spiking solutions at $4^{\circ}C$ ($\pm 2^{\circ}C$) in Teflon-sealed containers. The solutions should be checked frequently for stability. These solutions must be replaced after twelve months, or sconer if comparison with quality control check samples indicates a problem.

Bases/ Neutrals

<u>Acids</u>

1,2,4-Trichlorobenzene Acenaphthene 2,4-Dinitrotoluene Pyrene N-Nitroso-di-n-propylamine 1,4-Dichlorobenzene

Pentachlorophenol Phenol 2-Chlorophenol 4-Chloro-3-methylphenol 4-Nitrophenol

1.6 Equipment

- 1.6.1. Glass scintillation vials at least 20 mL, with screw cap and teflon or aluminum foil liner.
- 1.6.2 Spatula stainless steel or Teflon.
- 1.6.3 Balance capable of weighing 100 g to \pm 0.01 g.
- 1.6.4 Vials and caps 2 mL for GC auto sampler.
- 1.6.5 Disposable pipets Pasteur; glass wool rinsed with methylene chloride.
- 1.6.6 Concentrator tubes 15 mL.
- 1.6.7 Ultrasonic cell disruptor Heat Systems, Ultrasonics, Inc., Model W-385 SONICATOR (475 Watt with pulsing capability, No. 200, 1/2 inch tapped disruptor horn and No. 419, 1/8 inch standard tapered MICROTIP probe), or equivalent device with a minimum of 375 Watt output capability. NOTE: In order to ensure that sufficient energy is transferred to the sample during extraction, the MICROTIP probe <u>must</u> be replaced if the tip begins to erode. Erosion of the tip is evidenced by a rough surface.
- 1.6.8 Sonabox acoustic enclosure recommended with above disruptors for decreasing cavitation sound.
- 1.6.9 Test tube rack.
- 1.6.10 Oven drying.
- 1.6.11 Desiccator.
- 1.6.12 Crucibles porcelain.
- 1.7 Medium Level Sample Preparation.
 - 1.7.1. Transfer the sample container into a fume hood. Decant and discard any water layer on a sediment sample. Mix samples thoroughly, especially composited samples. Discard any foreign objects such as sticks, leaves, and rocks. Transfer approximately 1 g (record weight to the nearest 0.1 g) of

OLM01.0

sample to a 20-mL vial. Wipe the mouth of the vial with a tissue to remove any sample material. Record the exact weight of sample taken. Cap the vial before proceeding with the next sample to avoid any cross-contamination.

- 1.7.1.1 Transfer 50 g of soil/sediment to a 100 mL beaker. Add 50 mL of water and stir for 1 hour. Determine pH of sample with glass electrode and pH meter while stirring. Report pH value on appropriate data sheets. If the pH of the soil is greater than 11 or less than 5, contact the Technical Project Officer cited in the contract for instructions on how to handle the sample. Document the instructions in the SDG Narrative. Discard this portion of sample. NOTE: If limited sample volume is received, use 5 g of soil and 5 mL of water for the pH determination. Note this in the SDG Narrative.
- 1.7.2 Immediately after weighing the sample for extraction, weigh 5-10 g of the sediment into a tared crucible. Determine the percent moisture by drying overnight at 105°C. Allow to cool in a desiccator before weighing. Concentrations of individual analytes will be reported relative to the dry weight of sediment.

<u>g of sample - g of dry sample</u> x 100 - % moisture

- 1.7.3 Add 2.0 g of anhydrous <u>powdered</u> sodium sulfate to the sample in the 20 mL vial from paragraph 1.7.1 and mix well.
- 1.7.4 Surrogates are added to all samples, spikes, and blanks. Add 0.5 mL of surrogate spiking solution to sample mixture.
- 1.7.5 Add 0.5 mL of matrix standard spiking solution to each of two 1 g portions from the sample chosen for spiking. The frequency of MS/MSD analysis is given in Section IV, paragraph 8.6.
- 1.7.6 Immediately add 9.5 mL of methylene chloride to the sample and disrupt the sample with the 1/8 inch tapered MICROTIP ultrasonic probe for 2 minutes at output control setting 5, in continuous mode (if using a sonicator other than Models W-375 or W-385, contact the Project Officer for appropriate output settings). Before extraction, make certain that the sodium sulfate is free flowing and not a consolidated mass. As required, break up large lumps with a clean spatula or, very carefully, with the tip of the unenergized probe.

Add only 9.0 mL of methylene chloride to the matrix spike samples to achieve a final volume of 10 mL.

1.7.7 Prepare a method blank with each group of medium soil/sediment samples extracted. For semivolatile analyses, a method blank for medium soil/sediment samples consists of 1 g of sodium sulfate (see paragraph 1.5.1), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section IV, paragraph 8.7

- 1.7.8 Loosely pack disposable Pasteur pipets with 2-3 cm glass wool plugs. Filter the extract through the glass wool and collect at least 8.0 mL in a concentrator tube.
- 1.7.9 If the extract is to be screened <u>prior to</u> GPC, concentrate 5.0 mL of the extract collected in paragraph 1.7.7 to 1.0 mL using the nitrogen blowdown technique described in paragraph 3.6.2. Transfer the concentrate to an autosampler vial for GC/FID or GC/MS for screening. The quantitation limits for the screening procedure in Section III are approximately 20,000 ug/Kg.
- 1.7.10 If the extract is to be cleaned up using GPC without screening, take at least 8.0 mL of the extract in paragraph 1.7.7 and proceed to paragraph 3 of this section. Following GPC, the 5.0 mL of extract collected must be concentrated to 0.5 mL by the nitrogen blowdown technique described in paragraph 3.6.2, and screened according to the procedures in Section III. In this case, the quantitation limits for the screening procedures in Section III are approximately 10,000 ug/Kg.

2. Low Level Preparation for Screening and Analysis of Semivolatiles

2.1 Summary of Method

A 30 gram portion of sediment is mixed with anhydrous powdered sodium sulfate and extracted with 1:1 methylene chloride/acetone using an ultrasonic probe. If the optional low level screen is used, a portion of this dilute extract is concentrated fivefold and is screened by GC/FID or GC/MS. If peaks are present at greater than 20,000 ug/kg, discard the extract and prepare the sample by the medium level method. If no peaks are present at greater than 20,000 ug/kg, the entire extract is concentrated, subjected to GPC clean up, and analyzed by GC/MS for extractable organics.

2.2 Interferences

Method interferences may be caused by contaminants in solvents, reagents, glassware, and other sample-processing hardware that lead to discrete artifacts and/or elevated baselines in the total ion current profiles. All of these materials routinely must be demonstrated to be free from interferences under the conditions of the analysis by running laboratory reagent blanks. Matrix interferences may be caused by contaminants that are coextracted from the sample. The extent of matrix interferences will vary considerably from source to source.

- 2.3 Apparatus and Materials
 - 2.3.1 Apparatus for determining percent moisture
 - 2.3.1.1 Oven drying.

2.3.1.2 Desiccator.

2.3.1.3 Crucibles - porcelain.

- 2.3.2 Disposable Pasteur glass pipets 1 mL.
- 2.3.3 Ultrasonic cell disruptor, Heat Systems, Ultrasonics, Inc., Model W-385 SONICATOR (475 Watt with pulsing capability, No. 305, 3/4 inch tapped high gain "Q" disruptor horn, or No. 208, 3/4 inch standard solid disruptor horn), or equivalent device with a minimum of 375 Watt output capability. NOTE: In order to ensure that sufficient energy is transferred to the sample during extraction, the horn <u>must</u> be replaced if the tip begins to erode. Erosion of the tip is evidenced by a rough surface.

Sonabox acoustic enclosure - recommended with above disruptors for decreasing cavitation sound.

- 2.3.4 Beakers 400 mL.
- 2.3.5 Vacuum filtration apparatus.

2.3.5.1 Buchner funnel.

- 2.3.5.2 Filter paper Whatman No. 41 or equivalent.
- 2.3.6 Kuderna-Danish (K-D) apparatus.
 - 2.3.6.1 Concentrator tube 10 mL, graduated (Kontes K-570040-1025 or equivalent).
 - 2.3.6.2 Evaporative flask 500 mL (Kontes K-570001-0500 or equivalent).
 - 2.3.6.3 Snyder column three-ball macro (Kontes K-503000-0121 or equivalent).
 - 2.3.6.4 Snyder column two-ball micro (Kontes K-569001-0219 or equivalent).
- 2.3.7 Silicon carbide boiling chips approximately 10/40 mesh. Heat to 400°C for 30 minutes or Soxhlet extract with methylene chloride.
- 2.3.8 Water bath heated, with concentric ring cover, capable of temperature control $(\pm 2^{\circ}C)$. The bath should be used in a hood.
- 2.3.9 Balance capable of accurately weighing ± 0.01 g.
- 2.3.10 Vials and caps 2 mL for GC auto sampler.
- 2.3.11 Balance analytical, capable of accurately weighing \pm 0.0001g.

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- 2.3.12 Nitrogen evaporation device equipped with a water bath that can be maintained at 35-40°C. (N-Evap by Organomation Associates, Inc., South Berlin, MA, or equivalent).
- 2.3.13 Pyrex glass wool.
- 2.3.14 Pasteur pipets disposable.

2.4 Reagents

- 2.4.1 Sodium Sulfate anhydrous <u>powdered</u> reagent grade, heated at 400°C for four hours, cooled in a desiccator, and stored in a glass bottle (Baker anhydrous powder, catalog #73898 or equivalent).
- 2.4.2 Methylene chloride, methanol, acetone, isooctane, 2-propanol, and benzene - pesticide residue analysis grade or equivalent.
- 2.4.3 Reagent water defined as water in which an interferent is not observed at or above the CRQL of each parameter of interest.
- 2.4.4 Sodium Sulfite reagent grade.
- 2.4.5 Base/Neutral and Acid Surrogate Spiking Solution

Surrogate standards are added to all samples and calibration solutions. The compounds specified are Phenol- d_5 , 2,4,6-Tribromophenol, 2-Fluorophenol, Nitrobenzene- d_5 , Terphenyl- d_{14} , 2-Fluorobiphenyl, 2-Chlorophenol- d_4 , and 1,2-Dichlorobenzene- d_4 . Prepare a surrogate standard spiking solution that contains Nitrobenzene- d_5 , Terphenyl- d_{14} , 2-Fluorobiphenyl, and 1,2-Dichlorobenzene- d_4 at a concentration of 100 ug/mL; Phenol- d_5 , 2,4,6-Tribromophenol, 2-Fluorophenol, and 2-Chlorophenol- d_4 at a concentration of 200 ug/mL. Store the spiking solutions at 4°C (\pm 2°C) in Teflon-sealed containers. The solutions should be checked frequently for stability. These solutions must be replaced after twelve months, or sooner if comparison with quality control check samples indicates a problem.

2.4.6 Base/Neutral and Acid Matrix Spiking solution

Prepare a spiking solution in methanol that contains the following compounds at a concentration of 100 ug/mL for base/neutrals and 200 ug/mL for acids. Store the spiking solutions at 4°C (\pm 2°C) in Teflon-sealed containers. The solutions should be checked frequently for stability. These solutions must be replaced after twelve months, or sooner if comparison with quality control check samples indicates a problem.

Bases/ Neutrals

<u>Acids</u>

1,2,4-Trichlorobenzene Acenaphthene 2,4-Dinitrotoluene Pyrene N-Nitroso-di-n-propylamine 1,4-Dichlorobenzene Pentachlorophenol Phenol 2-Chlorophenol 4-Chloro-3-methylphenol 4-Nitrophenol

- 2.5 Low Level Sample Preparation
 - 2.5.1 Decant and discard any water layer on a sediment sample. Mix samples thoroughly, especially composited samples. Discard any foreign objects such as sticks, leaves, and rocks.

Transfer 50 g of soil/sediment to 100 mL beaker. Add 50 mL of water and stir for 1 hour. Determine pH of sample with glass electrode and pH meter while stirring. Report pH value on appropriate data sheets. If the pH of the soil is greater than 11 or less than 5, contact the Technical Project Officer cited in the contract for instructions on how to handle the sample. Document the instructions in the SDG Narrative. Discard this portion of sample. NOTE: If limited sample volume is received, use 5 g of soil and 5 mL of water for the pH determination. Note this in the SDG Narrative.

- 2.5.2 The following steps should be performed rapidly to avoid loss of the more volatile extractables. Weigh approximately 30 g of sample to the nearest 0.1 g into a 400-mL beaker and add 60 g of anhydrous powdered sodium sulfate. Mix well. The sample should have a sandy texture at this point. Immediately, add 100 mL of 1:1 methylene chloride-acetone to the sample, then add the surrogates according to paragraph 2.5.2.3.
 - 2.5.2.1 Immediately after weighing the sample for extraction, weigh 5-10 g of the sediment into a tared crucible. Determine the percent moisture by drying overnight at 105°C. Allow to cool in a desiccator before weighing. Concentrations of individual analytes will be reported relative to the dry weight of sediment.

g of sample - g of dry sample g of sample x 100 - % moisture

2.5.2.2 Weigh out two 30 g (record weight to nearest 0.1 g) portions for use as matrix and matrix spike duplicates according to paragraph 2.5.2. Add 0.5 mL of the BNA matrix spike solution to each of two portions. The frequency of MS/MSD analysis is given in Section IV, paragraph 8.6.

- 2.5.2.3 Add 0.5 mL of base/neutral and acid surrogate standard to the sample and each of the aliquots in 2.5.2.2.
- 2.5.2.4 Prepare a method blank with each group of low soil/sediment samples extracted. For semivolatile analyses, a method blank for low soil/sediment samples consists of 30 g of sodium sulfate (see paragraph 2.4.1), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section IV, paragraph 8.7
- 2.5.3 Place the bottom surface of the tip of the 3/4 inch disruptor horn about 1/2 inch below the surface of the solvent but above the sediment layer.
- 2.5.4 Sonicate for 1 1/2 minutes with the W-385 (or 3 minutes with the W-375), using No. 208, 3/4 inch standard disruptor horn with output control knob set at 10 (or No. 305, 3/4 inch tapped high gain "Q" disruptor horn at 5) and mode switch on "1 sec. pulse" and % duty cycle knob set at 50% (if using a sonicator other than Models W-375 or W-385, contact the Project Officer for appropriate output settings). Do NOT use MICROTIP probe.
- 2.5.5 Decant and filter extracts through Whatman #41 filter paper using vacuum filtration or centrifuge and decant extraction solvent.
- 2.5.6 Repeat the extraction two more times with 2 additional 100 mL portions of 1:1 methylene chloride-acetone. Before each extraction, make certain that the sodium sulfate is free flowing and not a consolidated mass. As required, break up large lumps with a clean spatula or, <u>very carefully</u>, with the tip of the probe. Decant the extraction solvent after each sonication. On the final sonication, pour the entire sample into the Buchner funnel and rinse with 1:1 methylene chloride-acetone.
 - 2.5.6.1 If the sample is to be screened from the low level method <u>prior to</u> GPC, take 5.0 mL and concentrate to 1.0 mL following paragraph 3.6.1 or 3.6.2, but note that the final volume for screening is 1.0 mL, not 0.5 mL. Screen the extract as per Section III, paragraph 1., "Screening of Extractable Organic Extracts."
 - 2.5.6.2 After GC/FID or GC/MS screening, transfer the remainder of the 1 mL back to the total extract from paragraph 2.5.6. GAUTION: To minimize sample loss, autosamplers which pre-flush samples through the syringe should not be used.

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2.6 Concentration and Solvent Exchange

- 2.6.1 Low level soil/sediment samples prepared by the procedures in paragraph 2.5 will result in extracts containing a mixture of acetone and methylene chloride. Because all soil/sediment sample extracts <u>must</u> be subjected to GPC clean up prior to analysis, the majority of the acetone must be removed from the extract, otherwise it will have adverse effects on the GPC column. To remove the acetone from the sample extract, follow the steps in 2.6.2.
- 2.6.2 Transfer the extract to a Kuderna-Danish (K-D) concentrator consisting of a 10 mL concentrator tube and a 500 mL evaporative flask. Other concentration devices or techniques may be used if equivalency is demonstrated for all extractable compounds listed in Exhibit C.
- 2.6.3 Add one or two clean boiling chips to the evaporative flask and attach a three-ball Snyder column. Pre-wet the Snyder column by adding about 1 mL methylene chloride to the top. Place the K-D apparatus on a hot water bath (60 to 80°C) so that the concentrator tube is partially immersed in the hot water and the entire lower rounded surface of the flask is bathed with hot vapor. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 10 to 15 minutés. At the proper rate of distillation the balls of the column will actively chatter but the chambers will not flood with condensed solvent. When the apparent volume of liquid reaches 1 mL, remove the K-D apparatus and allow it to drain and cool for at least 10 minutes.
- 2.6.4 Dilute the extract to 10.0 mL with methylene chloride, and proceed with GPC clean up (see paragraph 3).

3. Extract Cleanup by Gel Permeation Chromatography

Gel permeation chromatography (GPC) is a size exclusion cleanup procedure using organic solvents and hydrophobic gels in the separation of synthetic macromolecules. The packing gel is porous and is characterized by the range or uniformity (exclusion range) of that pore size. In the choice of gels, the exclusion range must be larger than the molecular size of the molecules to be separated. A cross-linked divinyl benzenestyrene copolymer (SX-3 Bio Beads or equivalent) is specified for this method.

GPC is <u>required</u> for all soil/sediment samples, regardless of concentration level, for the elimination of lipids, polymers, copolymers, proteins, natural resins and polymers, cellular components, viruses, steroids, and dispersed high-molecular-weight compounds from the sample extract. GPC is appropriate for both polar and non-polar analytes, therefore, it can be used effectively to clean up extracts containing a broad range of analytes. Normally, this method is most efficient for removing high boiling materials that condense in the injection port area of a gas chromatograph (GC) or in the front of the GC column. This residue ultimately will reduce the chromatographic separation efficiency or column capacity because of adsorption of the target analytes on the active sites. Pentachlorophenol especially is susceptible to this problem.

In the event that the Laboratory fails to appropriately employ GPC clean-up procedures, the Agency will require the clean up and reanalysis of all affected samples or sample extracts at <u>no additional</u> <u>cost</u> to the Agency.

- 3.1 Apparatus and Materials
 - 3.1.1 Gel permeation chromatography (GPC) cleanup device. NOTE: GPC cleanup is <u>required</u> for all soil/sediment extracts.

Gel permeation chromatography system - GPC Autoprep Model 1002 A or B (Analytical Biochemical Laboratories, Inc., or equivalent) Systems that perform very satisfactorily also have been assembled from the following components - an HPLC pump, an auto sampler or a valving system with sample loops, and a fraction collector. All systems, whether automated or manual, must meet the calibration requirements of paragraph 3.4.

- 3.1.1.1 Chromatographic column 700 mm x 25 mm i.d. glass column. Flow is upward. To simplify switching from the UV detector during calibration to the GPC collection device during extract cleanup, an optional double 3-way valve (Rheodyne Type 50 Teflon Rotary Valve #10-262 or equivalent) may be attached so that the column exit flow can be shunted either to the UV flow-through cell or to the GPC collection device.
- 3.1.1.2 Guard column (Optional) 5 cm, with appropriate fittings to connect to the inlet side of the analytical column (Supelco 5-8319 or equivalent).
- 3.1.1.3 Bio Beads (S-X3) 200-400 mesh, 70 gm (Bio-Rad Laboratories, Richmond, CA, Catalog 152-2750 or equivalent). An additional 5 gm of Bio Beads is required if the optional guard column is employed. The quality of Bio Beads may vary from lot to lot because of excessive fines in some lots. In addition to fines having a detrimental effect on chromatography, they also can pass through the column screens and damage the valve.
- 3.1.1.4 Ultraviolet detector fixed wavelength (254 nm) with a semi-prep flow-through cell.

- 3.1.1.5 Strip chart recorder, recording integrator or laboratory data system.
- 3.1.1.6 Syringe 10-mL with Luerlok fitting.
- 3.1.1.7 Syringe filter assembly, disposable Bio-Rad "Prep Disc" sample filter assembly #343-0005, 25 mm, and 5 micron filter discs or equivalent. Check each batch for contaminants. Rinse each filter assembly (prior to use) with methylene chloride if necessary.
- 3.1.1.8 A description of a manual system assembled from parts can be found in Wise, R.H., Bishop, D.F., Williams, R.T. & Austern, B.M. "Gel Permeation Chromatography in the GC/MS Analysis of Organics in Sludges" U.S. EPA, Municipal Environmental Research Laboratory, Cincinnati, Ohio, 45268.

3.2 Reagents

3.2.1 GPC Calibration Solution - prepare a calibration solution in methylene chloride containing the following analytes (in elution order):

Compound	mg/mL
corn oil	25.0
bis(2-ethylhexyl)phthalate	1.0
methoxychlor	0.2
perylene	0.02
sulfur (optional)	0.08

NOTE: If used, sulfur is not very soluble in methylene chloride, however, it is soluble in warm corn oil. Therefore, one approach is to weigh out the corn oil, warm it and transfer the weighed amount of sulfur into the warm corn oil. Mix it and then transfer into a volumetric flask with methylene chloride, along with the other calibration compounds.

Store the calibration solution in an amber glass bottle with a Teflon-lined screw-cap at 4°C, and protect from light (refrigeration may cause the corn oil to precipitate. Before use, allow the calibration solution to stand at room temperature until the corn oil dissolves). Replace the calibration standard solution every 6 months, or more frequently if necessary.

3.3 Column Preparation

3.3.1 Weigh out 70 gm of Bio Beads (SX-3). Transfer them to a quart bottle with a Teflon-lined cap or a 500 mL separatory funnel with a large bore stopcock, and add approximately 300 mL of methylene chloride. Swirl the container to ensure the wetting of all beads. Allow the beads to swell for a minimum of 2 hours. Maintain enough solvent to cover the beads sufficiently at all times. If a guard column is to be used, repeat the above with 5 gm of Bio Beads in a 125 mL bottle or a beaker, using 25 mL of methylene chloride.

- 3.3.2 Turn the column upside down from its normal position, and remove the inlet bed support plunger (the inlet plunger is longer than the outlet plunger). Position and tighten the outlet bed support plunger as near the end as possible, but no closer than 5 cm (measured from the gel packing to the collar).
- 3.3.3 Raise the end of the outlet tube to keep the solvent in the GPC column, or close the column outlet stopcock. Place a small amount of solvent in the column to minimize the formation of air bubbles at the base of poured column packing.
- 3.3.4 Swirl the bead/solvent slurry to get a homogeneous mixture and, if the wetting was done in a quart bottle, quickly transfer it to a 500 mL separatory funnel with a large bore stopcock. Drain the excess methylene chloride directly into the waste beaker, and then start draining the slurry into the column by placing the separatory funnel tip against the column wall. This will help to minimize bubble formation. Swirl occasionally to keep the slurry homogeneous. Drain enough to fill the column. Place the tubing from the column outlet into a waste beaker below the column, open the stopcock (if attached), and allow the excess solvent to drain. Raise the tube to stop the flow, and close the stopcock when the top of the gel begins to look dry. Add additional methylene chloride to just rewet the gel.
- 3.3.5 Wipe any remaining beads and solvent from the inner walls of the top of the column with a laboratory tissue. Loosen the seal slightly on the other plunger assembly (long plunger) and insert it into the column. Make the seal just tight enough so that any beads on the glass surface will be pushed forward, but loose enough so that the plunger can be pushed forward.

CAUTION: Do not tighten the seal if beads are between the seal and the glass surface because this can damage the seal and cause leakage.

- 3.3.6 Compress the column as much as possible without applying excessive force. Loosen the seal and gradually pull out the plunger. Rinse and wipe off the plunger. Slurry any remaining beads and transfer them into the column. Repeat the step in paragraph 3.3.5 and reinsert the plunger. If the plunger cannot be inserted and pushed in without allowing beads to escape around the seal, continue compression of the beads without tightening the seal, and loosen and remove the plunger as described. Repeat this procedure until the plunger is inserted successfully.
- 3.3.7 Push the plunger until it meets the gel, then compress the column bed about four centimeters.

- 3.3.8 Pack the optional 5 cm column with approximately 5 gm of preswelled beads (different guard columns may require different amounts). Connect the guard column to the inlet of the analytical column.
- 3.3.9 Connect the column inlet to the solvent reservoir (reservoir should be placed higher than the top of the column) and place the column outlet tube in a waste container. Placing a restrictor in the outlet tube will force air out of the column more quickly. A restrictor can be made from a piece of capillary stainless steel tubing of 1/16" OD x 10/1000" ID x 2". Pump methylene chloride through the column at a rate of 5 mL/min for one hour.
- 3.3.10 After washing the column for at least one hour, connect the column outlet tube, without the restrictor, to the inlet side of the UV detector. Connect the system outlet to the outlet side of the UV detector. A restrictor (same size as the one in paragraph 3.3.9) in the outlet tube from the UV detector will prevent bubble formation which causes a noisy UV baseline. The restrictor will not effect flow rate. After pumping methylene chloride through the column for an additional 1-2 hours, adjust the inlet bed support plunger until approximately 6-10 psi backpressure is achieved. Push the plunger in to increase pressure or slowly pull outward to reduce pressure.
- 3.3.11 When the GPC column is not to be used for several days, connect the column outlet line to the column inlet to prevent column drying and/or channeling. If channeling occurs, the gel must be removed from the column, reswelled, and repoured as described above. If drying occurs, methylene chloride should be pumped through the column until the observed column pressure is constant and the column appears wet. Always recalibrate after column drying has occurred to verify retention volumes have not changed.
- 3.4 Calibration of the GPC Column
 - 3.4.1 Using a 10 mL syringe, load sample loop #1 with calibration solution (paragraph 3.2). With the ABC automated system, the 5 mL sample loop requires a minimum of 8 mL of the calibration solution. Use a firm, continuous pressure to push the sample onto the loop. Switch the valve so that GPC flow is through the UV flow-through cell.
 - 3.4.2 Inject the calibration solution and obtain a UV trace showing a discrete peak for each component. Adjust the detector and/or recorder sensitivity to produce a UV trace that meets the following requirements. Differences between manufacturer's cell volumes and detector sensitivities may require a dilution of the calibration solution to achieve similar results. An

analytical flow-through detector cell will require a much less concentrated solution than the semi-prep cell and, therefore, the analytical cell is not acceptable for use.

- Peaks must be observed and should be symmetrical for all compounds in the calibration solution.
- Corn oil and phthalate peaks must exhibit >85% resolution.
- Phthalate and methoxychlor peaks must exhibit >85% resolution.
- Methoxychlor and perylens peaks must exhibit >85% resolution.
- Perylene and sulfur peaks must not be saturated and must exhibit >90% baseline resolution.
- 3.4.3 Using the information from the UV trace, establish appropriate collect and dump time periods to ensure collection of all target analytes. Initiate column eluate collection just before elution of bis(2-ethylhexyl)phthalate and after the elution of the corn oil. Stop eluate collection shortly after the elution of perylens. Collection should be stopped before sulfur elutes. Use a "wash" time of 10 minutes after the elution of sulfur. Each laboratory is required to establish its specific time sequences.
- 3.4.4 Verify the flow rate by collecting column eluate for 10 minutes in a graduated cylinder and measure the volume, which should be 45-55 mL (4.5-5.5 mL/min). If the flow rate is outside of this range, corrective action must be taken, as described above. Once the flow rate is within the range of 4.5-5.5 mL/min, record the column pressure (should be 6-10 psi) and room temperature. Changes in pressure, solvent flow rate, and temperature conditions can affect analyte retention times and must be monitored. If the flow rate and/or column pressure do not fall within the above ranges, a new column should be prepared. A UV trace that does not meet the criteria in paragraph 3.4.2 would also indicate that a new column should be prepared. It may be necessary to obtain a new lot of Bio Beads if the column fails all the criteria.
- 3.4.5 Reinject the calibration solution after appropriate collect and dump cycles have been set, and the solvent flow and column pressure have been established.
 - 3.4.5.1 Measure and record the volume of collected GPC eluate in a graduated cylinder. The volume of GPC eluate collected for each sample extract processed may be used to indicate problems with the system during sample processing.
 - 3.4.5.2 The retention times for bis(2-ethylhexyl)phthalate and perylene must not vary more than ±5% between calibrations. If the retention time shift is >5%,

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take corrective action. Excessive retention time shifts are caused by the following:

- Poor laboratory temperature control or system leaks.
- o An unstabilized column that requires pumping methylene chloride through it for several more hours or overnight.
- Excessive laboratory temperatures causing outgassing of the methylens chloride.
- 3.4.6 Analyze a GPC blank by loading 5 mL of methylene chloride into the GPC. Concentrate the methylene chloride that passes through the system during the collect cycle using a Kuderna-Danish (KD) evaporator. Analyze the concentrate by GC/MS. If the blank exceeds one half the CRQL of any analyte, pump additional methylene chloride through the system for 1-2 hours. Analyze another GPC blank to ensure the system is sufficiently clean. Repeat the methylene chloride pumping if necessary.
- 3.5 Sample Extract Cleanup

It is very important to have consistent laboratory temperatures during an entire GPC run, which could be 24 hours or more. If temperatures are not consistent, retention times will shift, and the dump and collect times determined by the calibration standard no longer will be appropriate. The ideal laboratory temperature to prevent outgassing of the methylene chloride is 72°F.

- 3.5.1 In order to prevent overloading of the GPC column, highly viscous sample extracts must be diluted prior to cleanup. Any sample extract with a viscosity greater than that of a 1:1 glycerol:water solution must be diluted and loaded into several loops. Similarly, extracts containing more than 500 mg of nonvolatile residue per 5 mL of extract must be diluted and loaded into several loops. The nonvolatile residue may be determined by evaporating a 100 uL aliquot of the extract to dryness in a tared aluminum weighing pan, or other suitable container.
- 3.5.2 Particles greater than 5 microns may scratch the valve, which may result in a system leak and cross contamination of sample extracts in the sample loops. To avoid such problems, filter the extract through a 5 micron filter disc by attaching a syringe filter assembly containing the filter disc to a 10 mL syringe. Draw the sample extract through the filter assembly and into the 10 mL syringe. Disconnect the filter assembly before transferring the sample extract into a small glass container, e.g., a 15 mL culture tube with a Teflon lined screw cap. Alternatively, draw the extract into the syringe without the filter assembly. Attach the filter assembly and force the

extract through the filter and into the glass container. Draw a minimum of 8 mL of extract into a 10 mL syringe.

3.5.3 Attach the syringe to the turn lock on the injection port. Use firm, continuous pressure to push the sample onto the 5-mL sample loop. If the sample is difficult to load, some part of the system may be blocked. Take appropriate corrective action. If the back pressure is normal (6-10 psi) the blockage is probably in the valve. Blockage may be flushed out of the valve by reversing the inlet and outlet tubes and pumping solvent through the tubes (this should be done before sample loading).

> NOTE: Approximately 2 mL of the extract remains in the lines between the injection port and the sample loop; excess sample also passes through the sample loop to waste.

- 3.5.4 After loading a loop, and before removing the syringe from the injection port, index the GPC to the next loop. This will prevent loss of sample caused by unequal pressure in the loops.
- 3.5.5 After loading each sample loop, wash the loading port with methylene chloride in a PTFE wash bottle to minimize cross contamination. Inject approximately 10 mL of methylene chloride to rinse the common tubes.
- 3.5.6 After loading all the sample loops, index the GPC to the OO position, switch to the "RUN" mode and start the automated sequence. Process each sample using the collect and dump cycle times established in 3.4.
- 3.5.7 Collect each sample in a 250-mL Erlenmeyer flask, covered with aluminum foil to reduce solvent evaporation, or directly into a Kuderna-Danish evaporator. Monitor sample volumes collected. Changes in sample volumes collected may indicate one or more of the following problems:
 - Change in solvent flow rate, caused by channeling in the column or changes in column pressure.
 - Increase in column operating pressure due to the absorption of particles or gel fines onto either the guard column or the analytical column gel, if a guard column is not used.
 - Leaks in the system or significant variances in room temperature.
- 3.5.8 Concentrate the extract as per paragraphs 3.6.1 or 3.6.2.
- 3.5.9 Calibrate the GPC at least once per week, following the procedure outlined in 3.4. The UV trace must meet the requirements in paragraph 3.4.2. In addition, the retention times of the calibration compounds must be within ±5% of their retention times in the previous calibration. A copy of the UV

trace of the calibration solution must be submitted with the data for the associated samples.

- 3.5.10 If the requirements in paragraphs 3.4.2 and 3.5.9 cannot be met, the column may be cleaned by processing several 5 mL volumes of butyl chloride throught the system. Butyl chloride removes the discoloration and particles that may have precipitated out of the methylene chloride extracts. If a guard column is being used. replace it with a new one. This may correct the problem. If column maintenance does not restore the performance of the column, the column must be repacked with new packing and recalibrated.
- 3.6 Final Concentration of Extract
 - Transfer the sample extract to a K-D evaporator, attach the 3.6.1 micro-Snyder column to the concentrator tube and add a silicon carbide boiling chip to the concentrator tube. Pre-wet the Snyder column with 0.5 mL of methylene chloride. Place the K-D apparatus on the hot water bath (80-90°C) so that the concentrator tube is partially immersed in the hot water. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 5 to 10 minutes. When the apparent volume of the liquid reaches 0.4 mL, remove the K-D apparatus from the water bath and allow it to drain for at least 10 minutes while cooling. Remove the Snyder column and rinse the lower joint into the concentrator tube with 0.1 mL of methylene chloride. Adjust the final volume to 0.5 mL with methylene chloride. Concentrating the extract to 0.5 mL will account for the volume of extract (5 mL) not recovered after GPC.
 - 3.6.2 Nitrogen blowdown technique (taken from ASTM Method D 3086)

The following method may be used for final concentration of the semivolatile extract instead of the procedures in paragraph 3.6.1. Place the concentrator tube in a warm water bath (35°C) and evaporate the solvent volume to below 0.5 mL using a gentle stream of clean, dry nitrogen (filtered through a column of activated carbon). CAUTION: New plastic tubing must not be used between the carbon trap and the sample, since it may introduce interferences.

The internal wall of the tube must be rinsed down several times with methylene chloride during the operation. During evaporation, the tube solvent level must be kept below the water level of the bath. The extract must never be allowed to become dry. Concentrating the extract to 0.5 mL will account for the volume of extract (5 mL) not recovered after GPC.

3.7 Store all extracts at $4^{\circ}C$ ($\pm 2^{\circ}C$) in the dark in Teflon-sealed containers.

3.8 If the extract was <u>not</u> screened prior to GPC, proceed to Section III for the screening procedures. If the extract <u>was</u> screened prior to GPC, proceed with the GC/MS analysis in Section IV.

SECTION III

SCREENING OF SEMIVOLATILE ORGANIC EXTRACTS

1. <u>Summary of Method</u>

1.1 The solvent extracts of water and soil/sediment are screened on a gas chromatograph/flame ionization detector (GC/FID) using a fused silica capillary column (FSCC). For water samples, the results of the screen may be used to determine an appropriate dilution factor for the GC/MS analysis of the sample extract. For soil/sedimant samples, the results of the screen are used to determine which of the two sample preparation procedures (low or medium) is required, and to determine an appropriate dilution factor for GC/MS analysis. The results of the screen may be used also to assist the analyst in performing Gel Permeation Chromatography (GPC) clean up procedures on extracts of either water or soil/sediment samples.

2. Apparatus and Materials

- 2.1 Gas chromatograph an analytical system complete with a temperature programmable gas chromatograph and all required accessories including syringes, analytical columns, and gases. The injection port must be designed for on-column injection when using packed columns and for splitless injection when using capillary columns.
 - 2.1.1 Above GC equipped with flame ionization detector.
 - 2.1.2 GC column 30 m x 0.32 mm, 1 micron film thickness, silicone coated, fused silica capillary column (J & W Scientific DB-5 or equivalent).

3. Reagents

- 3.1 Methylene chloride pesticide residue analysis grade or equivalent.
- 3.2 GC calibration standard prepare a standard solution containing phenol, phenanthrens and di-n-octylphthalate.
 - 3.2.1 Stock standard solutions (1.00 ug/uL) Stock standard solutions can be prepared from pure standard materials or purchased solutions.
 - 3.2.1.1 Prepare stock standard solutions by accurately weighing about 0.0100 g of pure material. Dissolve the material in pesticide quality methylene chloride and dilute to volume in a 10 mL volumetric flask. Larger volumes may be used at the convenience of the analyst. If compound purity is assayed at 96% or greater, the weight may be used without correction to calculate the concentration of the stock standard. Commercially prepared stock standards may be used at any concentration if they are certified by the manufacturer or by an independent source (see Exhibit E).

- 3.2.1.2 Transfer the stock standard solutions into Teflonsealed screw-cap bottles. Store at -10°C to -20°C and protect from light. Stock standard solutions should be checked frequently for signs of degradation or evaporation, especially just prior to preparing calibration standards from them. Stock standard solutions must be replaced after six months, or sooner, if comparison with quality control check samples indicates a problem. Standards prepared from gases or reactive compounds such as styrene must be replaced after two months, or sooner, if comparison with quality control check samples indicates a problem.
- 3.2.2 Prepare a working standard mixture of the three compounds in methylene chloride. The concentration must be such that the volume injected equals 50 ng of each compound. The storage and stability requirements are the same as specified in paragraph 3.2.1.2.

4. <u>GC Calibration</u>

- 4.1 At the beginning of each 12 hour shift, inject the GC calibration standard. The following criteria must be met:
 - 4.1.1 The GC must be standardized for half scale response from 50 ng of phenanthrene.
 - 4.1.2 The GC must adequately separate phenol from the solvent front.
 - 4.1.3 A minimum of quarter scale response for 50 ng of di-n-octylphthalate must be exhibited.

5. <u>GC/FID Screening</u>

- 5.1 Suggested GC operating conditions are as follows:
 - o Initial Column Temperature Hold 50°C for 4 minutes.
 - o Column Temperature Program 50-280°C at 8 degrees/min.
 - o Final Column Temperature Hold 280°C for 8 minutes.
 - Injector Grob-type, splitless.
 - o Sample Volume 1-2 uL.
 - Carrier Gas Helium at 30 cm³/sec.
- 5.2 Inject the GC calibration standard and ensure the criteria specified in 4 are met before injecting samples. Estimate the response for 10 ng of phenanthrene.
- 5.3 Inject the appropriate extracts from Section II, including blanks.

6. Interpretation of Chromatograms

6.1 Water

- 6.1.1 If no sample peaks are detected, or all are less than full scale deflection, the undiluted extract is analyzed on GC/MS.
- 6.1.2 If any sample peaks are greater than full scale deflection, calculate the dilution necessary to reduce the major peaks to between half and full scale deflection. Use this dilution factor to dilute the extract for GC/MS analysis.

6.2 Soil/Sediment

- 6.2.1 If no sample peaks from the extract (from low or medium level preparation) are detected, or all are less than 10% full scale deflection, the sample must be prepared by the low level protocol, Section II, Part C, beginning at 2.
- 6.2.2 Peaks are detected at greater than 10% full scale deflection and less than or equal to full scale deflection.
 - 6.2.2.1 If the screen is from the medium level extract, proceed with GC/MS analysis of this extract with appropriate dilution if necessary.
 - 6.2.2.2 If screen is from the low level extract, discard extract and prepare sample by medium level method for GC/MS analysis.
- 6.2.3 Peaks are detected at greater than full scale deflection.
 - 6.2.3.1 If the screen is from the medium level preparation, calculate the dilution necessary to reduce the major peaks to between half and full scale deflection. Use this dilution factor to dilute the extract. This dilution is analyzed by GC/MS for extractable organics.
 - 6.2.3.2 If the screen is from the low level preparation, discard the extract and prepare a sample by the medium level method for GC/MS analysis.

7. <u>GC/MS Analysis</u>

- 7.1 Use the information from paragraph 6 (Interpretation of Chromatograms) to perform the GC/MS analysis, beginning Section IV, GC/MS Analysis of Semivolatiles.
- 7.2 The information from paragraph 6 may be usefull also in processing sample extracts through GPC clean up.

NOTE: The choice of screening sample extracts before or after GCP clean up is left to the laboratory.

SECTION IV

GC/MS ANALYSIS OF SEMIVOLATILES

1. Summary of Method

This method is to be used for the GC/MS analysis of semivolatiles screened by Section III protocols and for confirmation of pesticides/Aroclors identified by GC/EC, if concentrations permit.

2. Apparatus and Materials

- 2.1 Gas chromatograph/mass spectrometer system.
 - 2.1.1 Gas chromatograph an analytical system complete with a temperature programmable gas chromatograph suitable for splitless injection and all required accessories including syringes, analytical columns and gases.
 - 2.1.2 Column 30 m x 0.25 mm ID (or 0.32 mm) bonded-phase silicone coated fused silica capillary column (J&W Scientific DB-5 or equivalent). A film thickness of 1.0 micron is recommended because of its larger capacity. A film thickness of 0.25 micron may be used.
 - 2.1.3 Mass Spectrometer capable of scanning from 35 to 500 amu every 1 second or less, utilizing 70 volts (nominal) electron energy in the electron impact ionization mode and producing a mass spectrum which meets all the instrument performance criteria in Table 1 when 50 ng of decafluorotriphenylphosphine (DFTPP) is injected through the GC inlet. The instrument conditions required for the acquisition of the DFTPP mass spectrum are given in paragraph 4.3.4. NOTE: DFTPP criteria must be met before any sample extracts are analyzed. Any samples analyzed when DFTPP criteria have not been met will require reanalysis at no cost to the Agency.
 - 2.1.4 GC/MS interface any gas chromatograph to mass spectrometer interface that gives acceptable calibration points, at 50 ng or less per injection, for each of the parameters of interest, and achieves all acceptable performance criteria (Exhibit E), may be used. Gas chromatograph to mass spectrometer interfaces constructed of all-glass or glass-lined materials are recommended. Glass can be deactivated by silanizing with dichlorodimethylsilane.
 - 2.1.5 Data system a computer system must be interfaced to the mass spectrometer that allows the continuous acquisition and storage on machine readable media of all mass spectra obtained throughout the duration of the chromatographic program. The computer must have software that allows searching any GC/MS data file for ions of a specific mass and plotting such ion abundances versus time or scan number. This type of plot is defined as an Extracted Ion Current Profile (EICP). Software must be available that allows integrating the abundance in any EICP between specified time or scan number limits. Also, for the non-target compounds, software must be available that allows comparing sample spectra against reference spectra. The

1989 (or most recent) release of the NIST/EPA/MSDC mass spectral library shall be used as the reference library. The data system must be capable of flagging all data files that have been editted manually by laboratory personnel.

3. <u>Reagents</u>

3.1 Internal standards - 1,4 Dichlorobenzene-d₄, Naphthalene-d₈, Acenaphthene-d₁₀, Phenanthrene-d₁₀, Chrysene-d₁₂, Perylene-d₁₂.

An internal standard solution can be prepared by dissolving 100 mg of each compound in 50 mL of methylene chloride. It may be necessary to use 5 to 10 percent benzene or toluene in this solution and a few minutes of ultrasonic mixing in order to dissolve all the constituents. The resulting solution will contain each standard at a concentration of 2000 ng/uL. A 10 uL portion of this solution should be added to each 1 mL of sample extract. This will result in 40 ng of each internal standard in the 2 uL volume of extract injected into the GC/MS.

3.2 Prepare calibration standards at a minimum of five concentration levels (20, 50, 80, 120, and 160 total ng per 2 uL). Each calibration standard should contain each compound of interest and each surrogate. Nine compounds, Benzoic acid, 2,4-Dinitrophenol, 2,4,5-Trichlorophenol, 2-Nitroaniline, 3-Nitroaniline, 4-Nitroaniline, 4-Nitrophenol, 4,6-Dinitro-2-methylphenol, and Pentachlorophenol will require only a fourpoint initial calibration at 50, 80, 120, and 160 total ng, since detection at less that 50 ng per injection is difficult. Great care must be taken to maintain the integrity of all standard solutions. Store all standard solutions at -10°C to -20°C in screw-cap amber bottles with teflon liners. Fresh standards should be prepared every twelve months at a minimum. The continuing calibration standard (50 ng) should be prepared weekly and stored at 4°C (±2°C).

In order to facilitate the confirmation of pesticides and Aroclors from the semivolatile library search data (see Exhibit D PEST, paragraph 17), the laboratory may wish to include the pesticide/Aroclor target compounds listed in Exhibit C in the semivolatile continuing calibration standard. The laboratory may add any or all of these compounds to the semivolatile continuing calibration standard, but at a concentation of 10 ng/uL or less. If added to this GC/MS standard, these additional analytes are <u>not</u> reported on the semivolatile calibration form (Form VII), but must be included in the quantitation report for the continuing calibration standard. As only a single point calibration would be performed, no %RSD or percent difference criteria would apply to these additional analytes.

3.3 Instrument performance check solution - prepare a solution of decafluorotriphenylphosphine (DFTPP), such that a 2 uL injection will contain 50 ng of DFTPP. The DFTPP also be included in the calibration standards at this level.

4. Instrument Operating Conditions

4.1 Gas Chromatograph

The following are the recommended gas chromatographic analytical conditions:

Initial Column Temperature Hold	-	40°C for 4 minutes
Column Temperature Program	-	40-270°C at 10 degrees/min.
Final Column Temperature Hold	-	270°C for 10 minutes
Injector Temperature	-	250-300°C
Transfer Line Temperature	-	250-300°C
Source Temperature	-	according to manufacturer's specifications
Injector	-	Grob-type, splitless
Sample Volume	-	2 uL
Carrier Gas	-	Helium at 30 cm ³ /sec

4.2 Mass Spectrometer

The following are the recommended mass spectrometer analytical conditions:

Electron Energy - 70 volts (nominal) Mass Range - 35 to 500 amu Scan Time - not to exceed 1 second per scan

- 4.3 The GC/MS system must be tuned to meet the manufacturer's specifications, using a suitable calibrant such as FC-43 or perfluorokarosene (PFK). The mass calibration and resolution of the GC/MS system are verified by the analysis of the instrument performance check solution (paragraph 3.3).
 - 4.3.1 Prior to the analysis of any samples, blanks, or calibration standards, the Contractor must establish that the GC/MS system meets the mass spectral ion abundance criteria for the instrument performance check solution containing decafluorotriphenylphosphine (DFTPP).

- 4.3.2 The analysis of the instrument performance check solution may be performed as follows:
 - As an injection of up to 50 ng of DFTPP into the GC/MS
 - o By adding 50 ng of DFTPP to the calibration standards (paragraph 3.2) and analyzing the calibration standard.
- 4.3.3 The analysis of the instrument performance check solution must meet the ion abundance criteria given below.

TABLE 1

DFTPP KEY IONS AND ION ABUNDANCE CRITERIA FOR QUADRAPOLE MASS SPECTROMETERS

<u>Mass</u>	Ion Abundance Criteria
51	30.0 - 80.0 percent of mass 198
68	Less than 2.0 percent of mass 69
69	Present
70	Less than 2.0 percent of mass 69
127	25.0 - 75.0 percent of mass 198
197	Less than 1.0 percent of mass 198
198	Base peak, 100 percent relative abundance (see note)
199	5.0 - 9.0 percent of mass 198
275	10.0 - 30.0 percent of mass 198
365	Greater than 0.75 percent of mass 198
441	Present but less than mass 443
442	40.0 - 110.0 percent of mass 198
443	15.0 - 24.0 percent of mass 442

NOTE: All ion abundances MUST be normalized to m/z 198, the nominal base peak, even though the ion abundances of m/z 442 may be up to 110 percent that of m/z 198.

4.3.4 The abundance criteria listed above must be met for a 50 ng injection of DFTPP. The mass spectrum of DFTPP must be acquired in the following manner. Three scans (the peak apex scan and the scans immediately preceding and following the apex) are acquired and averaged. Background subtraction is required, and must be accomplished using a single scan prior to the elution of DFTPP.

4.3.5 The criteria above are based on adherence to the acquisition specifications identified in paragraph 4.3.4. The criteria are based on performance characteristics of instruments currently utilized in routine support of Program activities. These specifications, in conjunction with relative response factor criteria for 54 target compounds (see Table 2), are designed to control and monitor instrument performance associated with the requirements of this Statement of Work.

4.3.6 The instrument performance check solution must be analyzed once at the beginning of each 12-hour period during which samples or standards are analyzed.

> The twelve (12) hour time period for a GC/MS system instrument performance check and standards calibration (initial or continuing calibration criteria) begins at the moment of injection of the DFTPP analysis that the laboratory submits as documentation of a compliant instrument performance check. The time period ends after twelve (12) hours has elapsed according to the system clock.

5. <u>Calibration</u>

- 5.1 Prior to the analysis of samples and required blanks, and after the instrument performance check solution criteria have been met, each GC/MS system must be calibrated at a minimum of five concentrations to determine instrument sensitivity and the linearity of GC/MS response for the semivolatile target compounds.
- 5.2 The internal standards are added to all calibration standards and all sample extracts (including blanks, matrix spikes, and matrix spike duplicates) just prior to analysis by GC/MS. A 10 uL aliquot of the internal standard solution should be added to a 1 mL aliquot of calibration standards. The internal standards specified in paragraph 3.1 should permit most of the semivolatile target compounds to have relative retention times of 0.80 to 1.20, using the assignments of internal standards to target compounds given in Table 2.
- 5.3 The quantitation ions for each internal standard are given in Table 3. Use the primary ion listed in Table 3 for quantitation, unless interferences are present. If interferences prevent the use of the primary ion for a given internal standard, use the secondary ion(s) listed in Table 3.

TABLE 2 SEMIVOLATILE INTERNAL STANDARDS WITH CORRESPONDING TARGET COMPOUNDS AND SURROGATES ASSIGNED FOR QUANTITATION

1,4-Dichlorobenzene-d ₄	Naphthalene-d ₈	Acenaphthene-d ₁₀	Phenanthrene-d ₁₀	Chrysene-d ₁₂	Perylene-d ₁₂
Phenol bis(2-Chloroethyl) ether 2-Chlorophenol 1,3-Dichlorobenzene 1,4-Dichlorobenzene 2-Methylphenol 2,2'-oxybis- (1-Chloropropane) 4-Methylphenol N-Nitroso-Di-n- propylamine Hexachloroethane 2-Fluorophenol (surr) Phenol-d5 (surr) 2-Chlorobenzene-d4 (surr) 1,2-Dichlorobenzene-d4 (surr)	Nitrobenzene Isophorone 2-Nitrophenol 2,4-Dimethyl- phenol bis(2-Chloro- ethoxy)methane 2,4-Dichloro- phenol 1,2,4-Trichloro- benzene Naphthalene 4-Chloroaniline Hexachloro- butadiene 4-Chloro-3- methylphenol 2-Methylnaphth- alene Nitrobenzene-d ₅ (surr)	Hexachlorocyclo- pentadiene 2,4,6-Trichloro- phenol 2,4,5-Trichloro- phenol 2-Chloronaphthalene 2-Nitroaniline Dimethyl Phthalate Acenaphthylene 3-Nitroaniline Acenaphthene 2,4-Dinitrophenol 4-Nitrophenol Dibenzofuran 2,4-Dinitrotoluene 2,6-Dinitrotoluene Diethyl phthalate 4-Chlorophenyl phenyl ether Fluorene 4-Nitroaniline 2-Fluorobiphenyl (surr) 2,4,6-Tribromo- phenol (surr)	4,6-Dinitro-2- methylphenol N-nitrosodi- phenylamine 4-Bromophenyl phenyl ether Hexachloro- benzene Pentachloro- phenol Phenanthrene Carbazole Anthracene Di-n-butyl- phthalate Fluoranthene	Pyrene Butylbenzyl phthalate 3,3'-Dichloro- benzidine Benzo(a)- anthracene bis(2-Ethyl- hexyl)phthalate Chrysene Terphenyl-d ₁₄ (surr)	Di-n-octyl- phthalate Benzo(b)fluor- anthene Benzo(k)fluor- anthene Benzo(a)pyrene Indeno(1,2,3-cd) pyrene Dibenz(a,h)- anthracene Benzo(g,h,i)- perylene

V

surr - surrogate compound

D-42/SV

EQ 1

INTERNAL STANDARDS	Primary Ion	Secondary Ions
1,4-Dichlorobenzene-d ₄	152	115
Naphthalene-dg	136	68
Acenapthens-d ₁₀	164	162, 160
Phenanthrene-d ₁₀	188	94, 80
Chrysene-d ₁₂	240	120, 236
Perylens-d ₁₂	264	260, 265

TABLE 3 CHARACTERISTIC IONS FOR INTERNAL STANDARDS FOR SEMIVOLATILE COMPOUNDS

5.4 Prepare calibration standards at a minimum of five concentration levels for each target compound and surrogate, as specified in paragraph 3.2. Analyze 2 uL of each calibration standard and tabulate the area of the primary characteristic ion against concentration for each compound including the surrogate compounds. A 2 uL injection is <u>required</u>. Calculate relative response factors (RRF) for each compound using Equation 1.

$$RRF = \frac{A_x}{A_{is}} \times \frac{C_{is}}{C_x}$$

Where

- A_x Area of the characteristic ion for the compound to be measured (see Table 4)
- A_{is} Area of the characteristic ion for the specific internal standard (see Table 3)
- C_{is} Concentration of the internal standard (ng/uL)
- C_{y} = Concentration of the compound to be measured (ng/uL)
- 5.5 The average relative response factor (RRF) must be calculated for all compounds. Calculate the % Relative Standard Deviation (%RSD) of the RRF values for the initial calibration.

TABLE 4 CHARACTERISTIC IONS FOR SEMIVOLATILE TARGET COMPOUNDS AND SURROGATES

Parameter	Primary Ion	Secondary Ion(s)
Phenol	94	65, 66
bis(2-Chloroethyl)ether	93	63, 95
2-Chlorophenol	128	64, 130
1,3-Dichlorobenzene	146	148, 113
1,4-Dichlorobenzene	146	148, 113
1,2-Dichlorobenzene	146	148, 113
2-Methylphenol	108	107
2,2'-oxybis(1-Chloropropane)	45	77, 79
4-Methylphenol	108	107
N-Nitroso-di-propylamine	70	42, 101, 130
Hexachloroethane	117	201, 199
Nitrobenzene	77	123, 65
Isophorone	82	95, 138
2-Nitrophenol	139	65, 109
2,4-Dimethylphenol	107	121, 122
bis(2-Chloroethoxy)methane	93	95, 123
2,4-Dichlorophenol	162	164, 98
1,2,4-Trichlorobenzene	180	182, 145
Naphthalene	128	129, 127
4-Chloroaniline	127	129
Hexachlorobutadiene	225	223, 227
4-Chloro-3-methylphenol	107	144, 142
2-Methylnaphthalene	142	141
Hexachlorocyclopentadiene	237	235, 272
2,4,6-Trichlorophenol	196	198, 200
2,4,5-Trichlorophenol	196	198, 200
2-Chloronaphthalene	162	164, 127
2-Nitroaniline	65	92, 138
Dimethyl phthalate	163	194, 164
Acenaphthylene	152	151, 153
3-Nitroaniline	138	108, 92
Acenaphthene	153	152, 154
2,4-Dinitrophenol	184	63, 154
4-Nitrophenol	109	139, 65
Dibenzofuran	168	139
2,4-Dinitrotoluene	165	63, 182
2,6-Dinitrotoluene	165	89, 121

(continued)

TABLE 4 (continued) CHARACTERISTIC IONS FOR SEMIVOLATILE TARGET COMPOUNDS AND SURROGATES

?arameter	Primary Ion	Secondary Ion(s)
)iethylphthalate	149	177, 150
-Chlorophenyl-phenylether	204	206, 141
luorene	166	165, 167
-Nitroaniline	138	92, 108
,6-Dinitro-2-methylphenol	198	182, 77
N-Nitrosodiphenylamine	169	168, 167
-Bromophenyl-phenylether	248	250, 141
iexachlorobenzene	284	142, 249
Pentachlorophenol	266	264, 268
Phenanthrene	178	179, 176
Anthracene	178	179, 176
Carbazole	167	166, 139
)i-n-butylphthalate	149	150, 104
Fluoranthene	202	101, 100
Pyrene	202	101, 100
Butylbenzylphthalate	149	91, 206
3,3'-Dichlorobenzidine	252	254, 126
Senzo(a)anthracene	228	·229, 226
is(2-Ethylhexyl)phthalate	149	167, 279
Chrysene	228	226, 229
Di-n-Octyl phthalate	149	
Senzo(b)fluoranthene	252	253, 125
Senzo(k)fluoranthene	252	253, 125
Benzo(a)pyrene	252	253, 125
Indeno(1,2,3-cd)pyrene	276	138, 227
Dibenz(a,h)anthracene	278	139, 279
Senzo(g,h,i)perylene	276	138, 277
SURROGATES		
Phenol-d ₅	99	42, 71
2-Fluorophenol	112	64
2,4,6-Tribromophenol	330	332, 141
Nitrobenzene-d ₅	82	128, 54
2-Fluorobiphenyl	172	171
Terphenyl	244	122, 212
2-Chlorophenol-d ₄	132	68, 134
L,2-Dichlorobenzene-d ₄	152	115, 150

- 5.6 Response factor criteria have been established for the calibration of 45 of the semivolatile target compounds and 6 semivolatile surrogate compounds.
 - 5.6.1 The compounds listed in Table 5 must meet the minimum RRF and maximum %RSD criteria for the initial calibration, with allowance made for up to four semivolatile target and surrogate compounds. However, the RRFs for those four compounds must be greater than 0.010, and the %RSD of those four compounds must be less than or equal to 40.0% for the initial calibration to be acceptable.

TABLE	5
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RELATIVE RESPONSE FACTOR CRITERIA FOR INITIAL AND CONTINUING CALIBRATION OF SEMIVOLATILE TARGET COMPOUNDS

Semivolatile	Minimum	Maximum	Maximum
Compounds	RRF	%RSD	%Diff
Phenol	0.800	20.5	25.0
bis(-2-Chloroethyl)ether	0.700	20.5	25.0
2-Chlorophenol	0.800	20.5	25.0
1,3-Dichlorobenzene	0.600	20.5	25.0
l,4-Dichlorobenzene	0.500	20.5	25.0
1,2-Dichlorobenzene	0.400	20.5	25.0
2-Methylphenol	0.700	20.5	25.0
4-Methylphenol	0.600	20.5	25.0
N-Nitroso-Di-propylamine	0.500	20.5	25.0
Hexachloroethane	0.300	20.5	25.0
Nitrobenzene	0.200	20.5	25.0
Isophorone	0.400	20.5	25.0
2-Nitrophenol	0.100	20.5	25.0
2,4-Dimethylphenol	0.200	20.5	25.0
bis(-2-Chloroethoxy)methane	0.300	20.5	25.0
2,4-Dichlorophenol	0.200	20.5	25.0
1,2,4-Trichlorobenzene	0.200	20.5	25.0
Naphthalene	0.700	20.5	25.0
4-Chloro-3-methylphenol	0.200	20.5	25.0
2-Methylnaphthalene	0.400	20.5	25.0
2,4,6-Trichlorophenol	0.200	20.5	25.0
2,4,5-Trichlorophenol	0.200	20.5	25.0
2-Chloronaphthalene	0.800	20.5	25.0
Acenaphthylene	1.300	20.5	25.0
2,6-Dinitrotoluene	0.200	20.5	25.0
Acenaphthene	0.800	20.5	25.0
Dibenzofuran	0.800	20.5	25.0
2,4-Dinitrotoluene	0.200	20.5	25.0
-Chlorophenyl-phenylether	0.400	20.5	25.0
Fluorene	0.900	20.5	25.0
-Bromophenyl-phenylether	0.100	20.5	25.0
lexachlorobenzene	0.100	20.5	25.0
Pentachlorophenol	0.050	20.5	25.0

(continued)

TABLE 5 (continued)			
RELATIVE RESPONSE FACTOR CRITERIA FOR INITIAL AND CONTINUING			
CALIBRATION OF SEMIVOLATILE TARGET COMPOUNDS			

Semivolatile Compounds	Minimum RRF	Maximum &RSD	Maximum &Diff
Phenanthrene	0.700	20.5	25.0
Anthracene	0.700	20.5	25.0
Fluoranthene	0.600	20.5	25.0
Pyrene	0.600	20.5	25.0
Benzo(a)anthracene	0.800	20.5	25.0
Chrysene	0.700	20.5	25.0
Benzo(b)fluoranthene	0.700	20.5	25.0
Benzo(k)fluoranthene	0.700	20.5	25.0
Benzo(a)pyrene	0.700	20.5	25.0
Indeno(1,2,3-cd)pyrene	0.500	20.5	25.0
Dibenzo(a,h)anthracene	0.400	20.5	25.0
Benzo(g,h,i)perylene	0.500	20.5	25.0
Nitrobenzene-d ₅	0.200	20.5	25.0
2-Fluorobiphenyl	0.700	20.5	25.0
Terphenyl-d ₁₄	0.500	20.5	25.0
Phenol-d ₅	0.800	20.5	25.0
2-Fluorophenol	0.600	20.5	25.0
2-Chlorophenol-d ₄	0.800	20.5	25.0
1,2-Dichlorobenzene-d	0.400	20.5	25.0

5.6.2 Because performance data indicate erratic and poor linearity and sensitivity, the following compounds have no Maximum %RSD, or Maximum %Difference criteria; however, these compounds <u>must</u> meet a minimum RRF criterion of 0.010:

2,2'-oxybis(1-Chloropropane)
4-Chloroaniline
Hexachlorobutadiene
Hexachlorocyclopentadiene
2-Nitroaniline
Dimethylphthalate
3-Nitroaniline
2,4-Dinitrophenol
4-Nitrophenol
Diethylphthalate

4-Nitroaniline 4,6-Dinitro-2-methylphenol N-Nitrosodiphenylamine Di-n-butylphthalate Butylbenzylphthalate 3,3'-Dichlorobenzidine bis(2-Ethylhexyl)phthalate Di-n-octylphthalate 2,4,6-Tribromophenol

In addition, Carbazole, a new target compound for which there is insufficient performance data, must only meet a minimum RRF criterion of 0.010.

5.7 A check of the calibration curve must be performed once every 12 hours (see paragraph 4.3.6 for the definition of the twelve-hour time period). Check the relative response factors of those compounds for which RRF values have been established. If these criteria are met, the relative response factors for all compounds are calculated and reported. A percent difference of the daily relative response factor (12 hour) compared to the average relative response factor from the initial curve is calculated. Calculate the percent difference for each compound and compare with the maximum percent difference criteria listed above.

As with the initial calibration, up to four semivolatile target compounds in Table 5 may fail to meet the minimum RRF or maximum %D criteria, but the RRFs of those four compounds must be greater than or equal to 0.010, and the percent differences must be less than or equal to 40.0% for the continuing calibration to be acceptable.

- 5.8 Internal standard responses and retention times in all standards must be evaluated during or immediately after data acquisition. If the retention time for any internal standard changes by more than 0.50 minutes (30 seconds) from the latest daily (12 hour) calibration standard, the chromatographic system must be inspected for malfunctions, and corrections made as required. The extracted ion current profile (EICP) of the internal standards must be monitored and evaluated for each standard. If the EICP area for any internal standard changes by more than a factor of two (-50% to +100%), the mass spectrometric system must be inspected for malfunction, and corrections made as appropriate. When corrections are made, re-analysis of samples analyzed while the system was malfunctioning is necessary.
- 5.9 Each GC/MS system must be calibrated upon award of the contract, whenever the Contractor takes corrective action which may change or affect the initial calibration criteria (i.e., ion source cleaning or repair, column removal or replacement, etc.), or if the continuing calibration acceptance criteria have not been met.
- 5.10 If time remains in the 12 hour time period after meeting the acceptance criteria for the initial calibration, samples may be analyzed. It is not necessary to analyze a continuing calibration standard, if the initial calibration standard that is the same concentration as the continuing calibration standard meets the continuing calibration acceptance criteria. Quantify all sample results against the initial calibration standard that is the same concentration as the continuing calibration standard that is the same concentration as the continuing calibration standard (50ng/2uL).
- 5.11 If time does NOT remain in the 12-hour period beginning with the injection of the instrument performance check solution, a new injection of the instrument performance check solution must be made. The DFTPP may be included in the continuing calibration standard.
- 5.12 If the injection of the instrument performance check solution meets the criteria in Table 1, calculate the response factors for the continuing calibration standard and the percent difference of the response factors from the mean response factors in the initial calibration.
- 5.13 The response factors from the continuing calibration standard must meet the criteria in Table 5 prior to the analysis of any blanks or samples.
- 6. <u>Sample Analysis</u>
- 6.1 Sample extracts may be analyzed only after the GC/MS system has met the instrument performance check, initial calibration, and continuing

calibration requirements above. The same instrument conditions must be employed for the analysis of samples as were used for calibration.

- 6.2 Internal standard solution is added to each sample extract. Add 10 uL of internal standard solution to each accurately measured 1.0 mL of sample extract.
- 6.3 Make any extract dilution indicated by characterization prior to the addition of internal standards. If any further dilutions of water or soil/sediment extracts are made, additional internal standards must be added to maintain the required 40 ng (20 ng/uL) of each internal standard in the extract volume.
- 6.4 Inject 2 uL of the sample extract, containing 40 ng of each internal standard, into the GC/MS.

7. <u>Oualitative Analysis</u>

- 7.1 The compounds listed in the Target Compound List (TCL), Exhibit C, shall be identified by an analyst competent in the interpretation of mass spectra (see Bidder Qualification description) by comparison of the sample mass spectrum to the mass spectrum of a standard of the suspected compound. Two criteria must be satisfied to verify the identifications:
 - o Elution of the sample component at the GC relative retention time as the standard component.
 - o Correspondence of the sample component and standard component mass spectra.
 - 7.1.1 For establishing correspondence of the GC relative retention time (RRT), the sample component RRT must compare within ± 0.06 RRT units of the RRT of the standard component. For reference, the standard must be run on the same shift as the sample. If coelution of interfering components prohibits accurate assignment of the sample component RRT from the total ion chromatogram, the RRT should be assigned by using extracted ion current profiles for ions unique to the component of interest.
 - 7.1.2 For comparison of standard and sample component mass spectra, mass spectra obtained on the contractor's GC/MS are required. Once obtained, these standard spectra may be used for identification purposes, <u>only</u> if the contractor's GC/MS meets the DFTPP daily instrument performance requirements. These standard spectra may be obtained from the run used to obtain reference RRTs.
 - 7.1.3 The requirements for qualitative verification by comparison of mass spectra are as follows:
 - 7.1.3.1 All ions present in the standard mass spectra at a relative intensity greater than 10% (most abundant ion in the spectrum equals 100%) must be present in the sample spectrum.

- 7.1.3.2 The relative intensities of ions specified in 7.1.3.1 must agree within ±20% between the standard and sample spectra. (Example: For an ion with an abundance of 50% in the standard spectra, the corresponding sample ion abundance must be between 30 and 70 percent.)
- 7.1.3.3 Ions greater than 10% in the <u>sample</u> spectrum but not present in the <u>standard</u> spectrum must be considered and accounted for by the analyst making the comparison. In Task III, the verification process should favor false positives. All compounds meeting the identification criteria must be reported with their spectra. For all compounds below the CRQL report the actual value followed by "J", e.g., "3J."
- 7.1.4 If a compound cannot be verified by all of the criteria in 7.1.3, but in the technical judgement of the mass spectral interpretation specialist, the identification is correct, then the Contractor shall report that identification and proceed with quantification in paragraph 8.
- 7.2 A library search shall be executed for non-target sample components for the purpose of tentative identification. For this purpose, the 1989 (or most recent) release of the NIST/EPA/MSDC mass spectral library, containing 50,000 spectra, shall be used.
 - 7.2.1 Up to 20 nonsurrogate organic compounds of greatest apparent concentration not listed in Exhibit C for the semivolatile fraction shall be identified tentatively via a forward search of the NIST/EPA/MSDC mass spectral library. Substances with responses less than 10% of the nearest internal standard are not required to be searched in this fashion. Only after visual comparison of sample spectra with the nearest library searches will the mass spectral interpretation specialist assign a tentative identification. NOTE: Computer generated library search routines must not use normalization routines that would misrepresent the library or unknown spectra when compared to each other.
 - 7.2.2 Guidelines for making tentative identification.
 - 7.2.2.1 Relative intensities of major ions in the reference spectrum (ions greater than 10% of the most abundant ion) should be present in the sample spectrum.
 - 7.2.2.2 The relative intensities of the major ions should agree within \pm 20%. (Example: For an ion with an abundance of 50% in the standard spectra, the corresponding sample ion abundance must be between 30 and 70 percent.)
 - 7.2.2.3 Molecular ions present in reference spectrum should be present in sample spectrum.

- 7.2.2.4 Ions present in the sample spectrum but not in the reference spectrum should be reviewed for possible background contamination or presence of co-eluting compounds.
- 7.2.2.5 Ions present in the reference spectrum but not in the sample spectrum should be reviewed for possible subtraction from the sample spectrum because of background contamination or coeluting compounds. NOTE: Data system library reduction programs sometimes can create these discrepancies.
- 7.2.3 If, in the technical judgement of the mass interpretation spectral specialist, no valid tentative identification can be made, the compound should be reported as <u>unknown</u>. The mass spectral specialist should give additional classification of the unknown compound, if possible (i.e., unknown phthalate, unknown hydrocarbon, unknown acid type, unknown chlorinated compound). If probable molecular weights can be distinguished, include them.

8. <u>Ouantitation</u>

8.1 Target components identified shall be quantified by the internal standard method. The internal standard used shall be the one nearest the retention time to that of a given analyte (see Table 2). The EICP area of characteristic ions of analytes listed in Table 4 are used for quantitation.

Internal standard responses and retention times in all samples must be evaluated during or immediately after data acquisition. If the retention time for any internal standard changes by more than 0.50 minutes (30 seconds) from the latest daily (12 hour) calibration standard, the chromatographic system must be inspected for malfunctions, and corrections made as required. The extracted ion current profile (EICP) of the internal standards must be monitored and evaluated for each sample, blank, matrix spike, and matrix spike duplicate. The criteria are described in detail in the instructions for Form VIII, Internal Standard Area Summary. If the EICP area for any internal standard changes by more than a factor of two (-50% to +100%), the mass spectrometric system must be inspected for malfunction and corrections made as appropriate. If the analysis of a subsequent sample or standard indicates that the system is functioning properly, then corrections may not be required. The samples or standards with EICP areas outside the limits must be re-analyzed, and treated according to paragraphs 8.1.1 and 8.1.2 below. If corrections are made, then the laboratory must demonstrate that the mass spectrometric system is functioning properly. This must be accomplished by the analysis of a standard or sample that does meet the EICP criteris. After corrections are made, the re-analysis of samples analyzed while the system was malfunctioning is required.

8.1.1 If after re-analysis, the EICP areas for all internal standards are inside the contract limits (-50% to +100%), then the problem with the first analysis is considered to have been within the control of the laboratory. Therefore, submit <u>only</u> data from the analysis with EICPs within the contract limits. This is considered the <u>initial</u> analysis and must be reported as such on all data deliverables.

- 8.1.2 If the re-analysis of the sample does not solve the problem, i.e., the EICP areas are outside the contract limits for both analyses, then submit the EICP data and sample data from both analyses. Distinguish between the initial analysis and the re-analysis on all data deliverables, using the sample suffixes specified in Exhibit B. Document in the SDG Narrative all inspection and corrective actions taken.
- 8.1.3 Do not re-analyze MS/MSD samples that do not meet the EICP area limits.
- 8.2 The relative response factor (RRF) from the daily standard analysis is used to calculate the concentration in the sample. Secondary ion quantitation is allowed ONLY when there are sample interferences with the primary ion. If secondary ion quantitation is performed, document the reasons in the SDG Narrative. The area of a secondary ion cannot be substituted for the area of a primary ion unless a relative response factor is calculated using the secondary ion.

When target compounds are below contract required quantitation limits (CRQL) but the spectrum meets the identification criteria, report the concentration with a "J." For example, if CRQL is 10 ug/L and concentration of 3 ug/L is calculated, report as "3J."

Calculate the concentration in the sample using the relative response factor (RRF) as determined in 4.3 and the following equation:

Water

Concentration ug/L -
$$\frac{(A_{x})(I_{s})(V_{t})(Df)}{(A_{is})(RRF)(V_{o})(V_{i})}$$

A_x = Area of the characteristic ion for the compound to be measured

A_{is} = Area of the characteristic ion for the internal standard

- I_s = Amount of internal standard injected in nanograms (ng)
- V_o = Volume of water extracted in milliliters (mL)

 V_i - Volume of extract injected in microliters (uL)

 V_t - Volume of the concentrated extract in microliters (uL)

Df = Dilution Factor. The dilution factor for analysis of water samples for semivolatiles by this method is defined as follows:

uL most conc. extract used to make dilution + uL clean solvent uL most conc. extract used to make dilution

If no dilution is performed, Df = 1.0.

Soil/Sediment

Concentration ug/kg =
$$\frac{(A_x)(I_s)(V_t)(Df)}{(A_{is})(RRF)(V_i)(W_s)(D)}$$

Where

A_x, I_g, A_{ig} are as given for water, above.

- V_t Volume of the concentrated extract in microliters (uL)
- Df Dilution Factor. The dilution factor for analysis of soil samples for semivolatiles by this method is defined as follows:
- uL most conc. extract used to make dilution + mL clean solvent uL most conc. extract used to make dilution
- If no dilution is performed, Df = 1.0.
- V_i Volume of extract injected in microliters (uL)
- D = <u>100 % moisture</u> 100
- W_e Weight of sample extracted in grams (g)
- 8.3 An estimated concentration for non-target components tentatively identified shall be quantified by the internal standard method. For quantification, the nearest internal standard <u>free of interferences</u> shall be used. The formula for calculating concentrations is the same as in paragraph 8.2. Total area counts (or peak heights) from the total ion chromatograms are to be used for both the compound to be measured and the internal standard. A relative response factor (RRF) of one (1) is to be assumed. The resulting concentration shall be qualified as "J" (estimated, due to lack of a compound-specific response factor), and "N" (presumptive evidence of presence), indicating the quantitative and qualitative uncertainties associated with this non-target component. An estimated concentration should be calculated for all tentatively identified compounds as well as those identified as unknowns.
- 8.4 If the on-column concentration of any compound in any sample exceeds the initial calibration range, that sample must be diluted, the internal standard concentration readjusted, and the sample extract reanalyzed. Guidance in performing dilutions and exceptions to this requirement are given below.
 - 8.4.1 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
 - 8.4.2 The dilution factor chosen should keep the response of the largest peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.

- 8.4.3 Do not submit data for more than two analyses, i.e., the original sample extract and <u>one</u> dilution, or, if the semivolatile screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
- 8.4.4 Do not dilute MS/MSD samples solely to get non-spiked analytes within the calibration range. The spiking level of each compound in the semivolatile matrix spiking solution should not require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.
- 8.5 Calculate surrogate standard recovery on all samples, blanks, and spikes. Determine if recovery is within limits (see Table 6) and report on appropriate form.
 - 8.5.1 Calculate the concentrations of the surrogate compounds using the same equations as used for the target compounds. Calculate the recovery of each surrogate as follows:
 - %R = Concentration (or amount) found x 100 Concentration (or amount) spiked
 - 8.5.2 Determine if the sample surrogate recovery meets specifications as follows:
 - The eight semivolatile surrogates can be divided into three groups: base/neutral compounds (Nitrobenzene-d₅, 2-Fluorobiphenyl, and Terphenyl-d₁₄); acid compounds (Phenol-d₅, 2-Fluorophenol, and 2,4,6-Tribromophenol); and compounds with advisory QC limits (2-Chlorophenol-d₄ and 1,2-Dichlorobenzene-d₄.
 - If a single surrogate recovery from any group is not within the contract windows, the sample does not require reanalysis or reextraction.
 - If a single surrogate recovery from the base/neutral group and a single surrogate recovery from the acid group are not within the contract windows, the sample does not require reanalysis or reextraction.
 - Do not reanalyze or reextract if only surrogates with advisory QC limits are not within the contract windows.
 - 8.5.3 If the sample surrogate recovery does not meet specifications (i.e., if two base/neutral or two acid surrogates are out of limits or if recovery of any one base/neutral or acid surrogate is below 10%), the following are required:
 - o Check to be sure that there are no errors in calculations, surrogate solutions, and internal standards. Also check instrument performance.
 - o Reanalyze the sample if none of the above reveal a problem.
 - o If surrogate recoveries in a blank do not meet specifications, the blank may be reanalyzed alone.

- Do not reanalyze <u>dilutions</u> if surrogate recoveries are outside the limits.
- Never reanalyze the matrix spike or matrix spike duplicate (MS/MSD), even if surrogate recoveries are outside the limits.
- If the sample associated with the matrix spike and matrix spike duplicate does not meet specifications, it should be reanalyzed only if the MS/MSD surrogate recoveries are within the limits. If the sample and associated MS/MSD show the same pattern (i.e., outside the limits), then the sample does not require reanalysis and must not be submitted.

Document in the narrative the similarity in surrogate recoveries.

- 8.5.4 If the reanalysis of the sample solves the problem, then the problem was within the laboratory's control. Therefore, submit only data from the analysis with surrogate spike recoveries within the contract windows. This shall be considered the initial analysis and shall be reported as such on all data deliverables.
- 8.5.5 If none of the steps in paragraph 8.4.1 or 8.4.2 solves the problem, then, except as noted below, <u>reextract</u> and <u>reanalyze</u> the sample. If the reextraction and reanalysis of the sample solves the problem, then the problem was within the laboratory's control. Therefore, submit <u>only</u> data from the analysis with surrogate recoveries <u>within</u> the contract windows. This shall be considered the <u>initial</u> analysis and shall be reported as such on all data deliverables.
 - If surrogate recoveries in a blank do not meet specifications even after reanalysis, <u>all</u> of the samples associated with that blank must be reextracted along with the blank. The blank is intended to detect contamination in samples processed at the same time.
 - o Do not reextract diluted samples if surrogate recoveries are outside the limits.
 - Never reextract the matrix spike or matrix spike duplicate (MS/MSD), even if surrogate recoveries are outside the limits.
 - o If the sample associated with the matrix spike and matrix spike duplicate does not meet specifications after reanalysis, it should be reextracted only if the reanalysis surrogate recoveries are not within the limits and MS/MSD surrogate recoveries are within the limits. If the sample and associated MS/MSD show the same pattern (i.e., outside the limits), then the sample does not require reanalysis and must not be submitted.

Document in the narrative the similarity in surrogate recoveries.

8.5.6 If the reextraction and reanalysis of the sample does not solve the problem (i.e., the surrogate recoveries are outside the contract limits for both analyses), then submit the surrogate recovery data and sample analysis data from the initial analysis of <u>both</u> sample extracts. Distinguish between the initial analysis and the reanalysis on all data deliverables, using the sample suffixes supplied in Exhibit B.

	₹R.	% R	
Compound	<u>Water</u>	Soil	
Nitrobenzene-d ₅	35-114	23-120	
2-Fluorobiphenyl	43-116	30-115	
Terphenyl-d ₁₄	33-141	18-137	
Phenol-d ₅	10-110	24-113	
2-Fluorophenol	21-110	25-121	
2,4,6-Tribromophenol	10-123	19-122	
2-Chlorophenol-d4	33-110	20-130 (advisor	y)
1,2-Dichlorobenzene-d4	16-110	20-130 (advisor	

TABLE 6 SURROGATE RECOVERY LIMITS

- 8.6. A matrix spike and matrix spike duplicate must be performed for each group of samples of a similar matrix, for the following, whichever is most frequent:
 - o Each Case of field samples received, OR
 - o Each 20 field samples in a Case, OR
 - Each group of field samples of a similar concentration level (soils only), OR
 - Each 14 calendar day period during which field samples in a Case were received (said period beginning with the receipt of the first sample in that Sample Delivery Group).

Calculate the recovery of each matrix spike compound in the matrix spike and matrix spike duplicate and report on appropriate form.

- 8.6.1. Calculate the concentrations of the matrix spike compounds using the same equations as used for target compounds. Calculate the recovery of each matrix spike compound as follows:
 - %R = Concentration (or amount) found x 100 Concentration (or amount) spiked

8.6.2 Calculate the relative percent difference of the <u>recoveries</u> of each compound in the matrix spike and matrix spike duplicate as follows:

 $\frac{\text{RPD} = |MSR - MSDR|}{(1/2)(MSR+MSDR)} \times 100$

Where

RPD - Relative Percent Difference

MSR - Matrix Spike Recovery

MSDR - Matrix Spike Duplicate Recovery

The vertical bars in the formula above indicate the absolute value of the difference, hence RPD is always expressed as a positive value.

8.6.3 The limits for matrix spike compound recovery and RPD are given in Table 7. As these limits are only advisory, no further action by the laboratory is required, however, frequent failures to meet the limits for recovery or RPD warrant investigation by the laboratory, and may result in questions from the Agency.

	*R	RPD	ŧR	RPD
Compound	Water	Water	Soil	<u>Soil</u>
Phenol	12-110	42	26- 90	35
2-Chlorophenol	27-123	40	25-102	50
1,4-Dichlorobenzene	36- 97	28	28-104	27
N-Nitroso-di-n-propylamine	41-116	38	41-126	38
1,2,4-Trichlorobenzene	39- 98	28	38-107	23
4-Chloro-3-methylphenol	23- 97	42	26-103	33
Acenaphthene	46-118	31	31-137	19
4-Nitrophenol	10- 80	50	11-114	50
2,4-Dinitrotoluene	24- 96	38	28- 89	47
Pentachlorophenol	9-103	50	17-109	47
Pyrene	26-127	31	35-142	36

TABLE 7 MATRIX SPIKE RECOVERY AND RELATIVE PERCENT DIFFERENCE LIMITS

- 8.7 Method blank analysis must be performed once for the following, on each GC/MS system used to analyze samples, whichever is most frequent:
 - o Each Case, OR
 - Each 14 calendar day period during which samples in a Case are received (said period beginning with the receipt of the first sample in that Sample Delivery Group), OR

- o Each 20 samples in a Case, including matrix spikes and reanalyses, that are of similar matrix (water or soil) or similar concentration (soil only), OR
- o Whenever samples are extracted by the same procedure (continuous liquid-liquid extraction or sonication).

Determine the concentrations of any target compounds detected in the semivolatile method blank, using the equations in paragraph 8.2. The method blank must contain less than or equal to the Contract Required Quantitation Limit (CRQL) of the semivolatile target compounds in Exhibit C, except the phthalate esters, which must be less than or equal to five times (5x) the CRQL. For soil/sediment method blanks, CRQL value must be adjusted for percent moisture (see Exhibit B).

If a laboratory method blank exceeds these criteria, the Contractor must consider the analytical system to be out of control. The source of the contamination must be investigated and appropriate corrective measures <u>MUST</u> be taken and documented before further sample analysis proceeds. All samples processed with a method blank that is out of control (i.e., contaminated) <u>MUST</u> be reextracted and reanalyzed at no additional cost to the Agency. The Laboratory Manager, or his designee, must address problems and solutions in the SDG Narrative (Exhibit B).

9. <u>GC/MS Confirmation of Pesticides and Aroclors</u>

The requirements for GC/MS confirmation of pesticides and Aroclors are given in paragraph 17 of Exhibit D PEST. When performed, the characteristic ions to be used for these analytes are given in Table 8. Also see paragraph 3.2 of this section regarding the inclusion of these analytes in the semivolatile continuing calibration standard.

TABLE 8

Parameter	Primary Ion	Secondary Ion(s)	
alpha-BHC			
beta-BHC	183	181, 109	
delta-BHC	181	183, 109	
gamma-BHC (Lindane)	183	181, 109	
Heptachlor	183	181, 109	
Aldrin	100	272, 274	
Heptachlor epoxide	66	263, 220	
Endosulfan I	353	355, 351	
Dieldrin	195	339, 341	
4,4'-DDE	79	263, 279	
Endrin	246	248, 176	
Endrin ketone	263	82, 81	
Endrin aldehyde	317	67, 319	
Endosulfan II	67	250, 345	
4,4'-DDD	337	339, 341	
e,e Endosulfan sulfate	235	237, 165	
4,4'-DDT	272	387, 422	
Methoxychlor	235	237, 165	
•	. 227	228	
Chlordane (alpha and/or gamma)	373	375, 377	
Toxaphene	159	231, 233	
Aroclor-1016	222	260, 292	
Aroclor-1221	190	222, 260	
Aroclor-1232	190	222, 260	
Aroclor-1242	222	256, 292	
Aroclor-1248	292	362, 326	
Aroclor-1254	292	362, 326	
Aroclor-1260	360	362, 394	

CHARACTERISTIC IONS FOR PESTICIDES/AROCLORS

EXHIBIT D

ANALYTICAL METHODS FOR PESTICIDES/AROCLORS

EXHIBIT D

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SECTION I

INTRODUCTION

The analytical method that follows is designed to analyze water, sediment and soil from hazardous waste sites to determine the presence and concentration of the chlorinated pesticides and Aroclors found in the Target Compound List (Exhibit C). The method can be used for determining analyte concentrations in the range from the contract required quantitation limits (CRQL) to one million times the CRQL in these matrices. The method is based on EPA Method 608.

The method is divided into three sections: Introduction, Sample Preparation, and Analysis. Sample preparation covers sample extraction and cleanup techniques. The analysis section contains the specific GC/EC analytical methods for pesticides and Aroclors.

1. <u>Summary of the Method</u>

- 1.1 Continuous liquid-liquid or separatory funnel extraction procedures are employed for aqueous samples. Sonication extraction is required for soil/sediment samples (Section II, beginning at 6.2). The method specifies GPC, adsorption column cleanup, and sulfur cleanup techniques (Section II, beginning at 7).
- 1.2 The chlorinated pesticides and Aroclors listed in Exhibit C are determined by a two-column GC/ECD technique.
- 1.3 Sample extracts, standards, and blanks must be analyzed within an analytical sequence as defined in Section III. GC/ECD analysis begins with an initial demonstration of instrument performance and the calibration of all pesticides and Aroclors. Acceptable initial calibration is defined in Section III, beginning at 6. Initial calibration must be repeated whenever the calibration verification stipulated in Section III, 7, fails or when major instrument maintenance or modification is performed.
- 1.4 An instrument blank and a Performance Evaluation Mixture are analyzed no less than once in every 12 hour analytical sequence in order to monitor retention times, calibration factors, and column performance. Data can be collected only as long as the results for the Performance Evaluation Mixtures and instrument blanks fall within the limits defined in Section III, 7. If two consecutive unacceptable Performance Evaluation Mixtures are run, all extracts run since the previous acceptable Performance Evaluation Mixture must be reanalyzed. Additional Performance Evaluation Mixtures and blanks are recommended when highly contaminated samples are suspected.
- 1.5 Calibration and analysis sequence specifications of the GC/EC method apply independently to both GC columns.

- 1.6 Matrix spike and a matrix spike duplicate analyses must be prepared and analyzed at least once for each matrix type or once per Sample Delivery Group (SDG), whichever is most frequent.
- 1.7 Analysis of a sample on both GC columns is required for <u>all</u> samples, blanks, matrix spikes, and matrix spike duplicates.
- 1.8 A single component pesticide is identified if a peak is detected within its appropriate retention time window on each of two columns. Toxaphene and Aroclors are identified primarily by pattern recognition, but RTs of three to five major peaks must also be taken into consideration. Guidance on quantitation of Aroclors is given in Section III, paragraph 13.
- 1.9 Standards for all tentatively identified Aroclors must be run within 72 hours of the sample analysis in which they were observed. These standards are used to verify identification only; quantitation is based on the standard analyzed during initial calibration.
- 1.10 Quantitative analysis of pesticides/Aroclors must be accomplished by the external standard method. Three-point calibration curves for single component analytes and the surrogates must be generated during the initial calibration. A linear response range must be demonstrated from the CRQL to a high point at least 30 times greater than the CRQL. Single-point calibrations for multicomponent analytes are sufficient for quantitation by this method.
- 1.11 The ECD response for single component analytes must be within the three-point calibration range in order for quantitative measurements to be made. The ECD response for the Aroclors/toxaphene must not be larger than the response for the high point calibration analysis of the single component analytes. The extracts must be diluted if the ECD response exceeds the calibration range. Quantitation must be performed and reported for both GC columns.
- 1.12 Absolute retention times (RTs) are used for the identification of pesticides/Aroclors. The absolute retention time window is calculated during initial calibration as ± 0.04 minutes of the mean RT of the standard for all analytes except Methoxychlor, where the RT window is ± 0.05 minutes of the mean RT of the standard. For the multicomponent analytes, the absolute retention time window is calculated for each peak (3 to 5) as ± 0.04 minutes of the RT of the same peak in the initial calibration standard analysis.
- 1.13 The surrogates, 2,4,5,6-Tetrachloro-m-xylene and decachlorobiphenyl, must be added to all samples, blanks, matrix spikes, and matrix spike duplicates prior to extraction. The retention time of both surrogates must fall within the retention time windows for an analysis to be acceptable. The surrogate recoveries will be determined in all of these samples and will be reported to the Agency as a measure of method performance.

- 1.14 The criteria in Section III, paragraph 14, are used to determine whether an analysis is complete or whether additional cleanup, dilution, or reextraction is required.
- 1.15 Resolution difficulties have been associated with the following pairs of compounds using this method:
 - o On a DB-608 or equivalent column, DDE and Dieldrin; Methoxychlor and Endrin ketone; and Endosulfan I and gamma-Chlordane.
 - o On a DB-1701 or equivalent column, Endosulfan I and gamma-Chlordane; and Methoxychlor and Endosulfan sulfate.

SECTION II

SAMPLE PREPARATION AND STORAGE

SECTION II

PART A - SAMPLE STORAGE AND HOLDING TIMES.

1. <u>Procedures for Sample Storage</u>

The samples must be protected from light and refrigerated at $4^{\circ}C$ ($\pm 2^{\circ}C$) from the time of receipt until 60 days after delivery of a complete sample data package to the Agency. After 60 days the samples may be disposed of in a manner that complies with all applicable regulations.

The samples must be stored in an atmosphere demonstrated to be free of all potential contaminants and in a refrigerator used only for storage of samples received under this contract.

Samples and standards must be stored separately.

2. <u>Contract Required Holding Times</u>

The extraction of water samples by separatory funnel procedures must be completed within five days of the Validated Time of Sample Receipt (VTSR). Extraction of water samples by continuous liquid-liquid extraction procedures must be started within five days of VTSR. Extraction of soil/sediment samples by sonication must be completed within 10 days of VTSR.

Analysis of samples must be completed within 35 days following the start of extraction.

PART B - SAMPLE PREPARATION FOR EXTRACTABLE PESTICIDES AND AROCLORS

1. <u>Summary of Sample Preparation Methods</u>

1.2 Water Samples

A 1-L volume of sample is spiked with the surrogate solution and is extracted with methylene chloride by using a separatory funnel or a continuous extractor. The methylene chloride extract is dried and concentrated (5.3). The extract is then cleaned up by GPC (optional), exchanged to hexane, cleaned up by Florisil cartridge, and adjusted to a final volume of 10.0 mL (5.0 mL if GPC cleanup is performed) as described beginning at paragraph 7.2.

1.3 Soil/sediment Samples

A 30 g aliquot of sample is spiked with the surrogate solution and then mixed with sodium sulfate and extracted with a 1:1 acetons/methylene chloride solvent mixture by sonication. The extract is then filtered, dried, concentrated by K-D, and the solvent exchanged into methylene chloride (6.3). The extract is then cleaned up by GPC (<u>mandatory</u>), exchanged to hexane, cleaned up by Florisil cartridge, and adjusted to a final volume of 5.0 mL (7.2).

2. <u>Interferences</u>

- 2.1 Method interferences may be caused by contaminants in solvents, reagents, glassware, and other sample processing hardware. These contaminants lead to discrete artifacts or to elevated baselines in gas chromatograms. Routinely, all of these materials must be demonstrated to be free from interferences under the conditions of the analysis by running reagent blanks. Interferences caused by phthalate esters can pose a major problem in pesticide analysis. Common flexible plastics contain varying amounts of phthalates which are easily extracted during laboratory operations, so cross-contamination of glassware frequently occurs when plastics are handled. Interferences from phthalates can best be minimized by avoiding the use of such plastics in the laboratory.
- 2.2 Matrix interferences may be caused by contaminants that are coextracted from the sample. The extent of matrix interferences will vary considerably from source to source, depending upon the nature of the site being sampled. The cleanup procedures must be used to remove such interferences in order to achieve the Contract Required Quantitation Limits.
- 3. <u>Apparatus and Materials</u> (Brand names, suppliers, and part numbers are for illustrative purposes only)
- 3.1 Continuous liquid-liquid extractors with Teflon or glass connecting lines for use with methylene chloride, (Hershberg-Wolf Extractor, Ace Glass Company, Vineland, NH, P/N 6841-10 or equivalent).
- 3.2 Separatory funnel 2000 mL with Teflon stopcock.

- 3.3 Apparatus for determining percent moisture
 - 3.3.1 Oven drying.
 - 3.3.2 Desiccator.
 - 3.3.3 Crucibles porcelain (optional).
 - 3.3.4 Aluminum weighing pans (optional).
- 3.4 Sonic cell disruptor Heat Systems, Ultrasonics, Inc., Model W-385 (475 watt with pulsing capability, No. 207 3/4-inch tapped disruptor horn), or equivalent device with a minimum 375 Watt output capability. NOTE: In order to ensure that sufficient energy is transferred to the sample during extraction, the horn must be replaced if the tip begins to erode. Erosion of the tip is evidenced by a rough surface.
- 3.5 Sonabox (or equivalent) for use with disruptor to decrease noise level.
- 3.6 Beakers 400-mL.
- 3.7 Kuderna-Danish (K-D) apparatus.
 - 3.7.1 Concentrator tube 10-mL, graduated (Kontes K-570040-1029, or equivalent).
 - 3.7.2 Evaporative flask 500-mL (Kontes K-470001-0500, or equivalent).
 - 3.7.3 Snyder column three-ball macro (Kontes K-503000-0121, or equivalent).
- 3.8 Funnels and Filter Paper.
 - 3.8.1 Powder funnels 10-cm diameter (optional), for filtration/drying.
 - 3.8.2 Buchner funnels 9-cm diameter, for filtration (optional).
 - 3.8.3 Filter paper No. 41 Whatman (or equivalent), 9-cm circles (optional).
- 3.9 Boiling chips.
 - 3.9.1 Silicon carbide boiling chips approximately 10 to 40 mesh. Heat the chips to 400°C for 30 minutes or solvent rinse before use.
 - 3.9.2 Teflon boiling chips (optional) solvent rinse the chips before use.
- 3.10 Water bath heated, with concentric ring cover, capable of temperature control. NOTE: The water bath should be used in a hood.

- 3.11 Top loading balance capable of weighing accurately to \pm 0.01 g.
- 3.12 Balance analytical, capable of weighing accurately to \pm 0.0001 g. The balance must be calibrated with class S weights once per each 12hour workshift. The balances must also be checked annually by a certified technician.
- 3.13 Nitrogen evaporation device equipped with a heated bath that can be maintained at 35 to 40°C, N-Evap by Organomation Associates, Inc., South Berlin, MA (or equivalent).
- 3.14 Vials and caps 2-mL for GC auto sampler.
- 3.15 Gel permeation chromatography (GPC) cleanup device. NOTE: GPC cleanup is <u>required</u> for all extracts for <u>all</u> soils (see paragraph 7.1.1).

Gel permeation chromatography system - GPC Autoprep Model 1002 A or B, Analytical Biochemical Laboratories, Inc., or equivalent. Systems that perform very satisfactorily also have been assembled from the following components - an HPLC pump, an auto sampler or a valving system with sample loops, and a fraction collector. All systems, whether automated or manual, must meet the calibration requirements of 7.1.3.

- 3.15.1 Chromatographic column 700 mm x 25 mm i.d. glass column. Flow is upward. To simplify switching from the UV detector during calibration to the GPC collection device during extract cleanup, an optional double 3-way valve (Rheodyne Type 50 Teflon Rotary Valve #10-262 or equivalent) may be attached so that the column exit flow can be shunted either to the UV flowthrough cell or to the GPC collection device.
- 3.15.2 Guard column (Optional) 5 cm, with appropriate fittings to connect to the inlet side of the analytical column (Supelco 5-8319 or equivalent).
- 3.15.3 Bio Beads (S-X3) 200-400 mesh, 70 gm (Bio-Rad Laboratories, Richmond, CA, Catalog 152-2750 or equivalent). An additional 5 gm of Bio Beads is required if the optional guard column is employed. The quality of Bio Beads may vary from lot to lot because of excessive fines in some lots. In addition to fines having a detrimental effect on chromatography, they also can pass through the column screens and damage the valve.
- 3.15.4 Ultraviolet detector fixed wavelength (254 nm) with a semiprep flow-through cell.
- 3.15.5 Strip chart recorder, recording integrator or laboratory data system.
- 3.15.6 Syringe 10-mL with Luerlok fitting.

- 3.15.7 Syringe filter assembly, disposable Bio-Rad "Prep Disc" sample filter assembly #343-0005, 25 mm, and 5 micron filter discs or equivalent. Check each batch for contaminants. Rinse each filter assembly (prior to use) with methylene chloride if necessary.
- 3.15.8 A description of a manual system assembled from parts can be found in Wise, R.H., Bishop, D.F., Williams, R.T. & Austern, B.M. "Gel Permeation Chromatography in the GC/MS Analysis of Organics in Sludges" U.S. EPA, Municipal Environmental Research Laboratory, Cincinnati, Ohio, 45268.
- 3.16 Florisil 500-mg or 1-g cartridges with stainless steel or Teflon frits, (Catalog No. 694-313, Analytichem, 24201 Frampton Ave., Harbor City, CA, or equivalent).
- 3.17 Vacuum system for eluting multiple cleanup cartridges.
 - 3.17.1 Vac Elute Manifold Analytichem International, J.T. Baker, or Supelco (or equivalent). The manifold design must ensure that there is no contact between plastics containing phthalates and sample extracts.
 - 3.17.2 Vacuum trap made from a 500-mL sidearm flask fitted with a onehole stopper and glass tubing.
 - 3.17.3 Vacuum pressure gauge.
 - 3.17.4 Rack for holding 10-mL volumetric flasks in the manifold.
- 3.18 Pyrex glass wool rinsed with methylene chloride and dried before use.
- 3.19 Bottle or test tube 20-mL with Teflon-lined screw cap for sulfur removal.
- 3.20 Drying column chromatographic column approximately 400 mm long x 19 mm ID, with coarse frit. (Substitution of a small pad of disposable Pyrex glass wool for the frit will help prevent cross-contamination of sample extracts.)
- 3.21 Glass vials minimum of 20-mL, with screw cap and Teflon or aluminum foil liner.
- 3.22 Spatula stainless steel or Teflon.
- 3.23 pH Paper wide range, (Hydrion Papers, Microessential Laboratory, Brooklyn, NY, or equivalent).
- 3.24 Pipet Volumetric 1.00-mL or 2.00-mL (optional).
- 3.25 Syringe 1.00-mL or 2.00-mL (optional).
- 3.26 Flask Volumetric 10.00-mL.

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- 3.27 Flask Volumetric 1.00-mL or 2.00-mL (optional).
- 3.28 Vials 10-mL, with screw cap and Teflon liner (optional).
- 3.29 Tube centrifuge, 12- to 15-mL with 19-mm ground glass joint (optional).
- 3.30 Snyder Column micro two or three ball with a 19-mm ground glass joint.
- 3.31 Centrifuge table top (optional).
- 3.32 Vortex mixer Genie, Model 550-6, Scientific Industrial, Inc., Bohemia, NY, or equivalent.
- 3.33 pH Meter with a combination glass electrode.
- 3.34 Magnetic stirrer motor Model PC353, Corning Co., Corning, NY, or equivalent.
- 3.35 Magnetic stir bar Teflon coated, at least 4 cm long.
- 3.36 Graduated cylinder 1 L capacity.

4. <u>Reagents</u>

- 4.1 Sodium sulfate granular-anhydrous reagent grade, heated at 400°C for 4 hours, or at 120°C for 16 hours, cooled in a desiccator, and stored in a glass bottle. Each lot must be extracted with hexane and analyzed by GC/ECD to demonstrate that it is free of interference before use. Baker anhydrous granular, Catalog No. 3375, or equivalent. CAUTION: An open container of sodium sulfate may become contaminated during storage in the laboratory.
- 4.2 Methylene chloride, hexane, acetone, toluene, iso-octane, and methanol (optional) - pesticide quality or equivalent. It is recommended that each lot of solvent used be analyzed to demonstrate that it is free of interference before use. Methylene chloride must be certified as acid free or must be tested to demonstrate that it is free of hydrochloric acid. Acidic methylene chloride must be passed through basic alumina and then demonstrated to be free of hydrochloric acid.
- 4.3 Mercury triple distilled, for sulfur clean-up.
- 4.4 Copper powder (optional) fine, granular (Mallinckrodt 4649 or equivalent). Copper may be used instead of mercury for sulfur cleanup. Remove oxides by treating with dilute nitric acid, rinse with distilled water to remove all traces of acid, rinse with acetone, and dry under a stream of nitrogen.
- 4.5 Sodium hydroxide solution (10 N) Carefully dissolve 40 g of NaOH in reagent water and dilute the solution to 100 mL.
- 4.6 Concentrated sulfuric acid 18 N.

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- 4.7 Reagent water defined as a water in which no interferent is observed at one-half the CRQL of any pesticide/Aroclor when one liter of the reagent water is extracted and prepared by using the same workup procedure as for a water sample.
- 4.8 Ten percent acetone in hexane (v/v) prepare by adding 10.0 mL of acetone to 90.0 mL of hexane. NOTE: Prepare this mixture accurately or the results from the Florisil cartridge cleanup will be adversely affected. Water in the acetone also will adversely affect Florisil performance.
- 4.9 Standards
 - 4.9.1 The Contractor must provide all standards to be used with this contract. These standards may be used only after they have been certified according to the procedure in Exhibit E. The Contractor must be able to verify that the standards are certified. Manufacturer's certificates of analysis must be retained by the Contractor and presented upon request.
 - 4.9.2 Stock standard solutions (1.00 ug/uL) can be prepared from pure standard materials or purchased as certified solutions.
 - 4.9.2.1 Prepare stock standard solutions by accurately weighing about 0.0100 g of pure material. Dissolve the material in toluene, dilute to volume in a 10-mL volumetric flask with toluene or acetone.
 - 4.9.2.2 Transfer the stock standard solutions into a bottle/vial with Teflon-lined cap or septa. Store at 4°C (±2°C) and protect from light. Stock standard solutions must be replaced after six months or sooner, if comparison with check standards indicates a problem.
 - 4.9.3 GPC calibration solution prepare a solution in methylene chloride that contains the following analytes in the concentrations listed below:

Analyte	mg/mL
corn oil	25
bis-2-ethylhexyl phthalate	1.0
methoxychlor	0.2
perylene	0.02
sulfur	0.08

NOTE: Sulfur is not very soluble in methylene chloride, however, it is soluble in warm corn oil. Therefore, one approach is to weigh out the corn oil, warm it and transfer the weighed amount of sulfur into the warm corn oil. Mix it and then transfer into a volumetric flask with methylene chloride, along with the other calibration compounds. Store the calibration solution in an amber glass bottle with a Teflon lined screw-cap at 4°C, and protect from light. (Refrigeration may cause the corn oil to precipitate. Before use, allow the calibration solution to stand at room temperature until the corn oil dissolves.) Replace the calibration standard solution every six months, or more frequently if necessary.

- 4.9.4 Surrogate solution the surrogates, Tetrachloro-m-xylene and Decachlorobiphenyl, are added to all standards, samples, matrix spikes, and blanks. Prepare a surrogate spiking solution of 0.2 ug/mL of each of the two compounds in acetone. The solution should be checked frequently for stability. The solution must be replaced after six months, or sooner, if comparison with quality control check samples indicates a problem. CAUTION: Analysts must allow all spiking solutions to equilibrate to room temperature before use.
- 4.9.5 Pesticide matrix spiking solution prepare a spiking solution in acetone or methanol that contains the following pesticides in the concentrations specified:

<u>Pesticide</u>	<u>ug/mL</u>
gamma-BHC (Lindane)	1.0
4,4'-DDT	4.0
Endrin	4.0
Heptachlor	1.0
Aldrin	1.0
Dieldrin	4.0

The solution must be prepared every six months, or sooner if the solution has degraded or concentrated.

4.9.6 Florisil cartridge check solution.

Prepare a solution of 2,4,5-trichlorophenol in acetone, at a concentration of 0.1 ug/mL.

5. <u>Extraction of Water Samples</u>

Water samples may be extracted by either a separatory funnel procedure or a continuous liquid-liquid extraction procedure. If an emulsion prevents acceptable solvent recovery with the separatory funnel procedure, continuous liquid-liquid extraction <u>must</u> be employed.

5.1 Separatory Funnel Extraction

5.1.1 Measure out each 1.0 L sample aliquot in a separate graduated cylinder. Measure and record the pH of the sample with wide range pH paper and adjust the pH to between 5 and 9 with 10 N sodium hydroxide or conc. sulfuric acid, if required. Samples requiring pH adjustment must be noted in the SDG Narrative. Place the sample into a 2-L separatory funnel.

- 5.1.2 For each sample selected for matrix spike and matrix spike duplicate analyses, measure out two additional 1-L portions and transfer those portions into separate funnels. Adjust the pH of each, if required, and fortify each with 1.0 mL of matrix spike solution before continuing the extraction. The frequency of MS/MSD analysis is given in Section III, paragraph 16.
- 5.1.3 Using a syringe or a volumetric pipet, add 1.0 mL of the surrogate solution to all water samples, matrix spikes, and blanks.
- 5.1.4 Add 60 mL methylene chloride to the separatory funnel and extract the sample by shaking the funnel for two minutes, with periodic venting to release excess pressure. Allow the organic layer to separate from the water phase for a minimum of 10 minutes. If the emulsion interface between layers is more than one-third the volume of the solvent layer, the analyst must employ mechanical techniques to complete the phase separation. The optimum technique depends upon the sample, and may include stirring, filtration of the emulsion through glass wool, centrifugation or other physical means. Drain the methylene chloride into a 250 mL Erlenmeyer flask.
- 5.1.5 Add a second 60 mL volume of methylene chloride to the sample bottle and repeat the extraction procedure a second time, combining the extracts in the Erlenmeyer flask. Perform a third extraction in the same manner.
- 5.1.6 Prepare a method blank with each group of water samples extracted. For pesticide/Aroclor analyses, a method blank for water samples consists of a 1 L volume of reagent water (see paragraph 4.7), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section III, paragraph 15.
- 5.2 Continuous Liquid-Liquid Extraction
 - 5.2.1 Add methylene chloride (100 to 250 mL) to the bottom of the extractor and fill it to a depth of at least one inch above the bottom sidearm.
 - 5.2.2 Measure out each 1.0 L sample aliquot in a separate graduated cylinder. Measure and record the pH of the sample with wide range pH paper and adjust the pH to between 5 and 9 with 10 N sodium hydroxide or concentrated sulfuric acid, if required. Samples requiring pH adjustment must be noted in the SDG Narrative. Place the sample into the continuous extractor.
 - 5.2.3 With some samples it may be necessary to place a layer of glass wool between the methylene chloride and the water layers in the extractor to prevent precipitation of suspended solids into the methylene chloride during extraction.

- 5.2.4 For each sample selected for matrix spike and matrix spike duplicate analyses, measure out two additional 1-L portions and transfer those portions into separate funnels. Adjust the pH of each, if required, and fortify each with 1.0 mL of matrix spike solution before continuing the extraction. The frequency of MS/MSD analysis is given in Section III, paragraph 16.
- 5.2.5 Using a syringe or a volumetric pipet, add 1.0 mL of the surrogate solution to all water samples, matrix spikes, and blanks.
- 5.2.6 Adjust the level of methylene chloride in the extractor so that the bottom sidearm is half filled with solvent.
- 5.2.7 Add sufficient methylene chloride to the distilling flask to ensure proper solvent cycling during operation and extract the solution for 18 hours.
- 5.2.8 Prepare a method blank with each group of water samples extracted. For pesticide/Aroclor analyses, a method blank for water samples consists of a 1 L volume of reagent water (see paragraph 4.7), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section III, paragraph 15.
- 5.3 Extract Drying and Concentration
 - 5.3.1 Assemble a Kuderna-Danish (K-D) concentrator by attaching a 10 mL concentrator tube to a 500 mL evaporative flask. Other concentration devices or techniques may be used in place of the K-D if equivalency is demonstrated for all the pesticide/Aroclor target compounds listed in Exhibit C.
 - 5.3.2 Pour the extract through a drying column containing about 10 cm of anhydrous granular sodium sulfate and collect the extract in the K-D concentrator. Rinse the Erlenneyer flask and the column with at least two additional 20 to 30 mL portions of methylene chloride to complete the quantitative transfer.
 - 5.3.3 Add one or two clean boiling chips to the evaporative flask and attach a three-ball Snyder column. Pre-wet the Snyder column by adding about 1 mL of methylene chloride to the top of the column. Place the K-D apparatus on a hot water bath (60° -80°C) so that the concentrator tube is partially immersed in the hot water and the entire lower rounded surface of the flask is bathed with hot vapor. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 15 to 30 minutes. At the proper rate of distillation, the balls of the column will actively chatter, but the chambers will not flood with condensed solvent. When the apparatus. Allow it to drain and cool for at least 10 minutes. DO NOT ALLOW THE EVAPORATOR TO GO DRY.

5.3.4 If no GPC cleanup is required, proceed with the hexane exchange described in paragraph 7.2. If GPC cleanup is to be used, remove the Snyder column, rinse the flask and its lower joint and collect the rinsate in the concentrator tube, adjust the volume to 10.0 mL with methylene chloride. Proceed to 7.1.

6. Extraction of Soil/Sediment Samples

- 6.1 Sample Preparation
 - 6.1.1 Mix samples thoroughly, especially composited samples. Discard any foreign objects such as sticks, leaves and rocks. Also, decant and discard any standing aqueous phase.
 - 6.1.2 pH Determination transfer 50 g of soil/sediment to a 100-mL beaker. Add 50 mL of water and stir the solution with a magnetic stirrer for 1 hour. Determine the pH of the sample by using a glass electrode and the pH meter while the sample is stirred. Report pH value on the appropriate data sheet. If the pH of the soil is > 9 or < 5, document any subsequent problems in the analysis related to pH in the SDG Narrative, but do not attempt to adjust the pH of the sample. Discard the portion of the sample used for pH determination.</p>

NOTE: If insufficient volume of soil is received, use 5 g of soil and 5 mL of water for the pH determination and note in the SDG Narrative.

6.1.3 Percent Moisture - Weigh 5 to 10 g of the sediment to the nearest 0.01 g into a tared crucible or aluminum weighing pan. Determine the weight percent volatilized by drying overnight at 105°C (hereafter referred to as percent moisture). After the sample is dry, remove the sample and pan and allow them to cool in a desiccator before weighing. Calculate the percent moisture according to Equation 1 below. Concentrations of individual analytes will be reported relative to the dry weight of sediment. CAUTION: Gases volatilized from some soil/sediment samples may require that this drying procedure be carried out in a hood.

Percent Moisture - Wt of Sample - Wt of Dry Sample Wt of sample x 100 EQ. 1

6.2 Extraction with Sonication

- 6.2.1 Tune the sonicator according to the manufacturer's directions prior to extracting samples by this procedure.
- 6.2.2 Weigh approximately 30 g of sample (to the nearest 0.1 g) into a 250 or 400-mL beaker and add 60 g of anhydrous sodium sulfate (granular).

- 6.2.3 For a sample to be used for matrix spike and matrix spike duplicate analysis, weigh out two additional 30 g (record weight to nearest 0.1 g) portions of sample and add 2.0 mL of the pesticide matrix spike solution to each soil aliquot. The frequency of MS/MSD analysis is given in Section III, paragraph 16.
 - 6.2.4 Add 2.0 mL of surrogate solution to all soil samples, matrix spikes, and blanks by using a volumetric pipet or a syringe. Mix the solution well. The sample and the added sodium sulfate should be a homogeneous, granular mixture at this point.
 - 6.2.5 Immediately add 80 to 100 mL of 1:1 methylene chloride/acetone to the sample.
 - 6.2.6 Place the bottom surface of the sonicator probe about 1/2 inch below the surface of the solvent but above the sediment layer.
 - 6.2.7 Sonicate for 3 minutes using a 3/4-inch horn at full power (output control knob at 10) with pulse on and percent duty cycle knob set at 50 percent. Do not use a microtip.

NOTE: These settings refer to the Model W-385. When using a sonicator other than Model W-385, refer to the instructions provided by the manufacturer for appropriate output settings.

- 6.2.8 The extracted sample can be filtered by using gravity or vacuum filtration.
 - 6.2.8.1 For gravity filtration, prepare a filtration/drying bed by placing a plug of glass wool in the neck of a 10-cm powder funnel and filling the funnel to approximately half its depth (4 or 5 cm) with anhydrous sodium sulfate (80-100 g). Decant the extract through the packed funnel and collect it in a 500-mL evaporation (K-D) flask.
 - 6.2.8.2 For vacuum filtration, use Whatman No. 41 paper in the Buchner funnel. Pre-wet the paper with methylene chloride/acetone before decanting the solvent.
- 6.2.9 Repeat the extraction two more times with additional 80 to 100 mL portions of the 1:1 methylene chloride/acetone. Before each extraction, thoroughly mix the solid residue, and make certain that the sodium sulfate is free flowing and not a consolidated mass. As required, break up large lumps with a clean spatula. Decant and filter the extraction solvent after each sonication by using the same funnel described in paragraph 6.2.8. After the final sonication, pour the entire sample into the funnel and rinse the beaker and funnel with 60 mL of 1:1 methylene chloride/acetone.

6.2.10 Prepare a method blank with each group of soil/sediment samples extracted. For pesticide/Aroclor analyses, a method blank for soil/sediment samples consists of 30 g of sodium sulfate (see paragraph 4.1), spiked with the surrogates and carried through the entire analytical procedure. The frequency of method blank analysis is given in Section III, paragraph 15.

6.3 Soil Extract Concentration

- 6.3.1 Add one or two clean boiling chips to the evaporative flask and attach a three-ball macro Snyder column. Pre-wet the Snyder column by adding about 1 mL of methylene chloride to the top. Place the K-D apparatus on a hot water bath (60 to 80°C) so that the concentrator tube is partially immersed in the hot water and the entire lower rounded surface of the flask is bathed with hot vapor. Adjust the vertical position of the apparatus and the water temperature as required to complete the concentration in 15 to 30 minutes. At the proper rate of distillation, the balls of the column will actively chatter, but the chambers will not flood with condensed solvent. Reduce the volume of liquid to less than 10 mL. Remove the K-D apparatus and allow it to drain and cool for at least 10 minutes. DO NOT ALLOW THE EVAPORATOR TO GO DRY.
- 6.3.2 In order to remove most of the acetone, it is absolutely necessary to further reduce the volume of all soil/sediment extracts to 1.0 mL. This is best accomplished using the nitrogen blowdown technique (7.3.2). The presence of acetone will cause a dead volume to develop in the GPC column and thus will cause loss of surrogates and analytes during GPC cleanups.
- 6.3.3 Adjust the extract volume to 10.0 mL with methylene chloride. Proceed to 7, below, for <u>mandatory</u> GPC and Florisil cartridge cleanup of soil extracts.

7. <u>Extract Cleanup</u>

There are three cleanup procedures specified in this method: GPC, Florisil cartridge, and sulfur cleanup. GPC <u>must</u> be performed for all soil extracts. GPC <u>must</u> be performed for water extracts that contain higher molecular weight contaminants that interfere with the analysis of the target analytes. Florisil cartridge cleanup is <u>mandatory</u> for <u>all</u> extracts. Sulfur cleanup must be performed for all sample extracts contaminated with sulfur. Blanks and matrix spike and matrix spike duplicate samples must be subjected to the same cleanup as the unspiked samples.

- 7.1 Extract Cleanup by Gel Permeation Chromatography (GPC)
 - 7.1.1. GPC cleanup is <u>mandatory</u> for all soil/sediment extracts. GPC must be performed for water extracts that contain higher molecular weight contaminants that interfere with the analysis of the target analytes. Gel permeation chromatography (GPC) is

OLM01.0

a size exclusion cleanup procedure using organic solvents and hydrophobic gels in the separation of synthetic macromolecules. The packing gel is porous and is characterized by the range or uniformity (exclusion range) of that pore size. In the choice of gels, the exclusion range must be larger than the molecular size of the molecules to be separated. A cross-linked divinyl benzenestyrene copolymer (SX-3 Bio Beads or equivalent) is specified for this method.

GPC is recommended for the elimination from the sample of lipids, polymers, copolymers, proteins, natural resins and polymers, cellular components, viruses, steroids, and dispersed high-molecular-weight compounds. GPC is appropriate for both polar and non-polar analytes, therefore, it can be used effectively to clean up extracts containing a broad range of analytes.

Normally, this method is most efficient for removing high boiling materials that condense in the injection port area of a gas chromatograph (GC) or in the front of the GC column. This residue ultimately will reduce the chromatographic separation efficiency or column capacity because of adsorption of the target analytes on the active sites. Pentachlorophenol especially is susceptible to this problem. GPC system performance must be validated at least once every seven days by demonstrating 80-100 percent recovery of the pesticide matrix spike mixture and an Aroclor 1016/1260 mixture.

7.1.2 GPC Column Preparation

- 7.1.2.1 Weigh out 70 gm of Bio Beads (SX-3). Transfer them to a quart bottle with a Teflon-lined cap or a 500 mL separatory funnel with a large bore stopcock, and add approximately 300 mL of methylene chloride. Swirl the container to ensure the wetting of all beads. Allow the beads to swell for a minimum of 2 hours. Maintain enough solvent to cover the beads sufficiently at all times. If a guard column is to be used, repeat the above with 5 gm of Bio Beads in a 125 mL bottle or a beaker, using 25 mL of methylene chloride.
- 7.1.2.2 Turn the column upside down from its normal position, and remove the inlet bed support plunger (the inlet plunger is longer than the outlet plunger). Position and tighten the outlet bed support plunger as near the end as possible, but no closer than 5 cm (measured from the gel packing to the collar).

- 7.1.2.3 Raise the end of the outlet tube to keep the solvent in the GPC column, or close the column outlet stopcock. Place a small amount of solvent in the column to minimize the formation of air bubbles at the base of poured column packing.
- 7.1.2.4 Swirl the bead/solvent slurry to get a homogeneous mixture and, if the wetting was done in a quart bottle, quickly transfer it to a 500 mL separatory funnel with a large bore stopcock. Drain the excess methylene chloride directly into the waste beaker, and then start draining the slurry into the column by placing the separatory funnel tip against the column wall. This will help to minimize bubble formation. Swirl occasionally to keep the slurry homogeneous. Drain enough to fill the column. Place the tubing from the column outlet into a waste beaker below the column, open the stopcock (if attached), and allow the excess solvent to drain. Raise the tube to stop the flow, and close the stopcock when the top of the gel begins to look dry. Add additional methylene chloride to just rewet the gel.
- 7.1.2.5 Wipe any remaining beads and solvent from the inner walls of the top of the column with a laboratory tissue. Loosen the seal slightly on the other plunger assembly (long plunger) and insert it into the column. Make the seal just tight enough so that any beads on the glass surface will be pushed forward, but loose enough so that the plunger can be pushed forward.

CAUTION: Do not tighten the seal if beads are between the seal and the glass surface because this can damage the seal and cause leakage.

- 7.1.2.6 Compress the column as much as possible without applying excessive force. Loosen the seal and gradually pull out the plunger. Rinse and wipe off the plunger. Slurry any remaining beads and transfer them into the column. Repeat the step in paragraph 7.1.2.5 and reinsert the plunger. If the plunger cannot be inserted and pushed in without allowing beads to escape around the seal, continue compression of the beads without tightening the seal, and loosen and remove the plunger as described. Repeat this procedure until the plunger is inserted successfully.
- 7.1.2.7 Push the plunger until it meets the gel, then compress the column bed about four centimeters.

- 7.1.2.8 Pack the optional 5 cm column with approximately 5 gm of preswelled beads (different guard columns may require different amounts). Connect the guard column to the inlet of the analytical column.
- 7.1.2.9 Connect the column inlet to the solvent reservoir (reservoir should be placed higher than the top of the column) and place the column outlet tube in a waste container. Placing a restrictor in the outlet tube will force air out of the column more quickly. A restrictor can be made from a piece of capillary stainless steel tubing of 1/16" OD x 10/1000" ID x 2". Pump methylene chloride through the column at a rate of 5 mL/min for one hour.
- 7.1.2.10 After washing the column for at least one hour, connect the column outlet tube, without the restrictor, to the inlet side of the UV detector. Connect the system outlet to the outlet side of the UV detector. A restrictor (same size as the one in paragraph 7.1.2.9) in the outlet tube from the UV detector will prevent bubble formation which causes a noisy UV baseline. The restrictor will not effect flow rate. After pumping methylene chloride through the column for an additional 1-2 hours, adjust the inlet bed support plunger until approximately 6-10 psi backpressure is achieved. Push the plunger in to increase pressure or slowly pull outward to reduce pressure.
- 7.1.2.11 When the GPC column is not to be used for several days, connect the column outlet line to the column inlet to prevent column drying and/or channeling. If channeling occurs, the gel must be removed from the column, reswelled, and repoured as described above. If drying occurs, methylene chloride should be pumped through the column until the observed column pressure is constant and the column appears wet. Always recalibrate after column drying has occurred to verify retention volumes have not changed.

NOTE: The description of solvent flow rate and column pressure applies only to the ABC GPC apparatus. Laboratories using equivalent equipment must develop the parameters for their apparatus which give acceptable performance as described in 7.1.4. 7.1.2.12 The GPC calibration procedure is based on monitoring the elution of standards with a UV detector connected to the GPC column. Care must be taken to account for any difference in volume (elution time) between the GC column and the detector and between the GPC column and the collection vial.

> NOTE: The UV detector calibration procedure described in 7.1.3 is to be used for the analyses of organochlorine pesticides and Aroclors listed in Exhibit C. IT MUST NOT BE USED FOR THE ANALYSIS OF GC/MS EXTRACTABLES OR OTHER ANALYTES WITHOUT A RECOVERY STUDY.

- 7.1.3 Calibration of the GPC Column
 - 7.1.3.1 Using a 10 mL syringe, load sample loop #1 with calibration solution (paragraph 4.9.3). With the ABC automated system, the 5 mL sample loop requires a minimum of 8 mL of the calibration solution. Use a firm, continuous pressure to push the sample onto the loop. Switch the valve so that GPC flow is through the UV flow-through cell.
 - 7.1.3.2 Inject the calibration solution and obtain a UV trace showing a discrete peak for each component. Adjust the detector and/or recorder sensitivity to produce a UV trace that meets the following requirements. Differences between manufacturer's cell volumes and detector sensitivities may require a dilution of the calibration solution to achieve similar results. An analytical flow-through detector cell will require a much less concentrated solution than the semi-prep cell and, therefore, the analytical cell is not acceptable for use.
 - o Peaks must be observed and should be symmetrical for all compounds in the calibration solution.
 - o Corn oil and phthalate peaks must exhibit >85% resolution.
 - Phthalate and methoxychlor peaks must exhibit >85% resolution.
 - o Methoxychlor and perylene peaks must exhibit >85% resolution.
 - o Perylene and sulfur peaks must not be saturated and must exhibit >90% baseline resolution.
 - 7.1.3.3 Determine the elution times for the phthalate, methoxychlor, and perylene. Phthalate will elute first, perylene, last.

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- 7.1.3.4 Choose a "DUMP" time which removes > 85 percent of the phthalate. Choose a "COLLECT" time so that > 95 percent of the methoxychlor is collected, and continue to collect until just prior to the elution time of sulfur. Use a "WASH" time of 10 minutes.
- 7.1.3.5 NOTE: The DUMP and COLLECT times must be adjusted to compensate for the difference in volume of the lines between the detector and the collection flask.
- 7.1.3.6 Verify the flow rate by collecting column eluate for 10 minutes in a graduated cylinder and measure the volume. which should be 45-55 mL (4.5-5.5 mL/min). If the flow rate is outside of this range. corrective action must be taken, as described above. Once the flow rate is within the range of 4.5-5.5 nL/min. record the column pressure (should be 6-10 psi) and room temperature. Changes in pressure, solvent flow rate, and temperature conditions can affect analyte retention times and must be monitored. If the flow rate and/or column pressure do not fall within the above ranges. a new column should be prepared. A UV trace that does not meet the criteria in paragraph 7.1.3.2 would also indicate that a new column should be prepared. It may be necessary to obtain a new lot of Bio Beads if the column fails all the criteria.
- 7.1.3.7 Reinject the calibration solution after appropriate collect and dump cycles have been set, and the solvent flow and column pressure have been established.
 - 7.1.3.7.1 Measure and record the volume of collected GPC eluate in a graduated cylinder. The volume of GPC eluate collected for each sample extract processed may be used to indicate problems with the system during sample processing.
 - 7.1.3.7.2 The retention times for bis(2ethylhexyl)phthalate and perylene must not vary more than ±5% between calibrations. If the retention time shift is >5%, take corrective action. Excessive retention time shifts are caused by the following:
 - o Poor laboratory temperature control or system leaks.

- o An unstabilized column that requires pumping methylene chloride through it for several more hours or overnight.
- o Excessive laboratory temperatures causing outgassing of the methylene chloride.
- 7.1.3.8 Analyze a GPC blank by loading 5 mL of methylene chloride into the GPC. Concentrate the methylene chloride that passes through the system during the collect cycle using a Kuderna-Danish (KD) evaporator. Analyze the concentrate by GC/MS. If the blank exceeds one half the CRQL of any analyte, pump additional methylene chloride through the system for 1-2 hours. Analyze another GPC blank to ensure the system is sufficiently clean. Repeat the methylene chloride pumping if necessary.
- 7.1.4 GPC Calibration Check

No Florisil cleanup is used in the GPC calibration check.

- 7.1.4.1 At least once every 7 days, the calibration of the GPC must be verified with two check mixtures. The first mixture is prepared by concentrating 2.0 mL of the matrix spiking solution (paragraph 4.9.5) to less than 1 mL under a stream of nitrogen (7.3.2), and adjusting the final volume to 10.0 mL with methylene chloride. The second mixture is prepared with 2 ug of Aroclor 1016 and 2 ug of Aroclor 1260 in a final volume of 10.0 mL methylene chloride.
- 7.1.4.2 Load the first 5.0 mL sample loop by using a 10 mL syringe containing 8 mL of the diluted pesticide matrix spike solution (paragraph 7.1.4.1). The Aroclor mixture is loaded into Loop 2 in the same manner. Fractions are collected in an auto sequence by using the GPC program established by the UV detector calibration procedure (Paragraph 7.1.3).
- 7.1.4.3 The collected GPC calibration fraction is transferred to a K-D apparatus, and the collection vessel is rinsed with two additional 10-mL portions of methylene chloride to complete the transfer. The volume of methylene chloride is reduced (described in 7.3.2). After cooling, the solvent is exchanged to hexane according to the instruction in 7.2. The final volume is adjusted to 10.0 mL, and the sample is analyzed by GC according to the procedures in Section III. The analysis must be performed on at least one of the GC columns used for samples analysis.

- 7.1.4.4 The pattern of the Aroclor quantitation peaks and the recovery of each single component analyte must be determined for evaluation and reporting purposes. If the recovery of each of the analytes is 80 to 110 percent and if the Aroclor pattern is the same as with previously run standards, then the analyst may continue to use the column. If recoveries are out of the acceptance window or if changes in the relative peak heights of the patterns of the Aroclor are observed, the column must be replaced and recalibrated according to the instructions in 7.1.3.
- 7.1.4.5 Some samples may contaminate the SX-3 Bio Beads and change the retention volume of the GPC column. Therefore system calibration and analyte recovery must be checked whenever a sample causes significant discoloration of the GPC column. Even if no darkening is visible, GPC calibration must be checked not less than once every seven days. In many cases, the SX-3 Bio Beads may be used for several months as long as the column calibration and flow rate remain constant.
- 7.1.5 Daily UV calibration check (optional)

The calibration of the GPC may be monitored daily by use of the UV-GPC calibration solution (paragraph 4.9.3) and the UV Detector Calibration Procedure (7.1.3). The UV detector should be used to monitor the elution times for the phthalate, methoxychlor, and perylene, in that order. The precalibrated GPC program should "DUMP" > 85 percent of the phthalate and should "COLLECT" > 95 percent of the methoxychlor and perylene. Significant changes in elution times of the analytes (e.g., > 0.5 minutes) indicate that the column is out of calibration and must be recalibrated or replaced.

7.1.6 Sample Extract Cleanup

It is very important to have consistent laboratory temperatures during an entire GPC run, which could be 24 hours or more. If temperatures are not consistent, retention times will shift, and the dump and collect times determined by the calibration standard no longer will be appropriate. The ideal laboratory temperature to prevent outgassing of the methylene chloride is 72°F.

7.1.6.1 In order to prevent overloading of the GPC column, highly viscous sample extracts must be diluted prior to cleanup. Any sample extract with a viscosity greater than that of a 1:1 glycerol:water solution must be diluted and loaded into several loops. Similarly, extracts containing more than 500 mg of nonvolatile residue per 5 mL of extract must be diluted and loaded into several loops. The nonvolatile residue may be determined by evaporating a 100 uL aliquot of the extract to dryness in a tared aluminum weighing pan, or other suitable container.

- Particles greater than 5 micron may scratch the 7.1.6.2 valve, which may result in a system leak and cross contamination of sample extracts in the sample loops. To avoid such problems, filter the extract through a 5 micron filter disc by attaching a syringe filter assembly containing the filter disc to a 10 mL syringe. Draw the sample extract through the filter assembly and into the 10 mL syringe. Disconnect the filter assembly before transferring the sample extract into a small glass container, e.g., a 15 mL culture tube with a Teflon lined screw cap. Alternatively, draw the extract into the syringe without the filter assembly. Attach the filter assembly and force the extract through the filter and into the glass container. Draw a minimum of 8 mL of extract into a 10 mL syringe. INTRODUCTION OF PARTICULATES OR GLASS WOOL INTO THE GPC SWITCHING VALVES MAY REQUIRE FACTORY REPAIR OF THE APPARATUS.
- 7.1.6.3 Prior to loading samples, put the GPC into the "LOAD" mode, set the instrument terminal for the number of loops to be loaded, and set the "DUMP", "COLLECT", and "WASH" times for the values determined by the calibration procedure described in 7.1.3.
- 7.1.6.4 Attach the syringe to the turn lock on the injection port. Use firm, continuous pressure to push the sample onto the 5-mL sample loop. If the sample is difficult to load, some part of the system may be blocked. Take appropriate corrective action. If the back pressure is normal (6-10 psi) the blockage is probably in the valve. Blockage may be flushed out of the valve by reversing the inlet and outlet tubes and pumping solvent through the tubes (this should be done before sample loading).

NOTE: Approximately 2 mL of the extract remains in the lines between the injection port and the sample loop; excess sample also passes through the sample loop to waste.

7.1.6.5 After loading a loop, and before removing the syringe from the injection port, index the GPC to the next loop. This will prevent loss of sample caused by unequal pressure in the loops.

- 7.1.6.6 After loading each sample loop, wash the loading port with methylene chloride in a PTFE wash bottle to minimize cross contamination. Inject approximately 10 mL of methylene chloride to rinse the common tubes.
- 7.1.6.7 After loading all the sample loops, index the GPC to the OO position, switch to the "RUN" mode and start the automated sequence. Process each sample using the collect and dump cycle times established in 7.1.3.
- 7.1.6.8 Collect each sample in a 250-mL Erlenmeyer flask, covered with aluminum foil to reduce solvent evaporation, or directly into a Kuderna-Danish evaporator. Monitor sample volumes collected. Changes in sample volumes collected may indicate one or more of the following problems:
 - o Change in solvent flow rate, caused by channeling in the column or changes in column pressure.
 - o Increase in column operating pressure due to the absorption of particles or gel fines onto either the guard column or the analytical column gel, if a guard column is not used.
 - o Leaks in the system or significant variances in room temperature.
- 7.1.6.9 After the appropriate GPC fraction has been collected for each sample, exchange the sample extract solvent to hexane, as described beginning at 7.2. DO NOT ALLOW THE EXTRACT TO GO TO DRYNESS. Adjust the final volume of the hexane extract to 5.0 mL.
- 7.1.6.10 Any samples that were loaded into two or more loops should be recombined and adjusted to a final volume of 5.0 mL in hexane before proceeding to 7.3, Florisil Cartridge Procedure.
- 7.1.6.11 NOTE: A final volume of 5.0 mL is required for extract concentrates after GPC cleanup because only half of the sample is used in the GPC sample loop (5 mL of the 10 mL described in paragraph 5.3.4 or 6.3.3).

7.2 Solvent exchange into hexane

This procedure applies to both extracts of water samples and extracts of soil samples.

- 7.2.1 Momentarily remove the Snyder column, add 50 mL of hexane and a new boiling chip, and reattach the Snyder column. Pre-wet the column by adding about 1 mL of hexane to the top. Concentrate the solvent extract as before. When the apparent volume of liquid reaches 3 to 5 mL, remove the K-D apparatus and allow it to drain and cool for at least 10 minutes. DO NOT ALLOW THE EVAPORATOR TO GO DRY.
- 7.2.2 Remove the Snyder column; using 1 to 2 mL of hexane, rinse the flask and its lower joint into the concentrator tube. Complete quantitative transfer of the extract to a vial by using hexane.
- 7.2.3 Adjust the volume of the hexane extract to 10.0 mL. Proceed to 7.3 for Florisil cartridge cleanup.
- 7.3 Florisil Cartridge Procedure

Florisil cartridge cleanup is required for all extracts. Cleanup significantly reduces matrix interferences caused by polar compounds.

- 7.3.1 Cartridge Performance Check - every lot number of Florisil cartridges must be tested by the following procedure before they are used for sample cleanup. Add 0.5 mL of 2,4,5trichlorophenol solution (0.1 mg/mL in acetone) and 0.5 mL of Standard Mixture A, midpoint concentration (Section III, paragraph 3.3) to 4 mL of hexane. Reduce the final volume to 0.5 mL using nitrogen (paragraph 7.3.2). Place the mixture onto the top of a washed Florisil cartridge (paragraph 7.3.4.4), and elute it with 9 mL of hexane/acetone [(90:10)(V/V)]. Use two additional 1-mL hexane rinses to ensure quantitative transfer of standard from the cartridge. Reduce the final volume to 1.0 mL using nitrogen (7.3.2) and analyze the solution by GC/ECD. The recovery of each analyte must be determined for evaluation and reporting purposes. The lot of Florisil cartridges is acceptable if all pesticides are recovered at 80 to 110 percent, if the recovery of trichlorophenol is less than 5%, and if no peaks interfering with the target analytes are detected.
- 7.3.2 Nitrogen blowdown technique (taken from ASTM Method D 3086)
 - 7.3.2.1 Place the concentrator tube with an open mini-Snyder column attached in a heating bath (30 to 35°C) and evaporate the solvent to the final volume by blowing a gentle stream of clean, dry nitrogen (filtered through a column of activated carbon) onto the solvent. DO NOT ALLOW THE EXTRACT TO GO TO DRYNESS.

- 7.3.2.2 New plastic tubing must not be used between the carbon trap and the sample as it may introduce interferences. The internal wall of new tubing must be rinsed several times with hexane and then dried prior to use.
- 7.3.3 Extract preparation before Florisil cartridge cleanup

For samples which have been run through the GPC cleanup, exchange the solvent to hexane (7.2) and adjust the hexane extract volume to 5.0 mL by using the nitrogen blowdown technique described in 7.3.2. For those aqueous samples not passed through the GPC step, adjust the hexane extract volume to 10.0 mL. The different extract volumes are required because only half of the methylene chloride concentrates are used in the GPC cleanup.

- 7.3.4 Florisil cartridge cleanup
 - 7.3.4.1 Attach the vacuum manifold to a water aspirator or to a vacuum pump with a trap installed between the manifold and the vacuum source. Adjust the vacuum pressure in the manifold to between 5 and 10 pounds of vacuum.
 - 7.3.4.2 Place one Florisil cartridge into the vacuum manifold for each sample extract.
 - 7.3.4.3 The required Florisil cartridge size and the final volume of the extract after Florisil cleanup are a function of the GC autosampler that a laboratory uses. If the autosampler operates reliably with 1.0 mL of sample extract, then a 500-mg cartridge is used and the required final volume is 1.0 mL. If the autosampler requires more sample, prepare 2.0 mL of sample extract using a 1-g cartridge. Manual injection requires only a 1.0 mL final extract volume and a 500-mg cartridge.
 - 7.3.4.4 Prior to cleanup of samples, the cartridges must be washed with hexane/acetone (90:10). This is accomplished by placing the cartridge in the vacuum manifold, by pulling a vacuum, and by passing at least 5 mL of the hexane/acetone solution through the cartridge. While the cartridges are being washed, adjust the vacuum applied to each cartridge so that the flow rate through each cartridge is approximately equal. DO NOT ALLOW THE CARTRIDGES TO GO DRY AFTER THEY HAVE BEEN WASHED.
 - 7.3.4.5 After the cartridges in the manifold are washed, the vacuum is released, and a rack containing labeled 10-mL volumetric flasks is placed inside the manifold. Care must be taken to ensure that the

solvent line from each cartridge is placed inside of the appropriate volumetric flask as the manifold top is replaced.

- 7.3.4.6 After the volumetric flasks are in place, vacuum to the manifold is restored, and a volume of extract equal to the required final volume (1.0 or 2.0 mL) from each sample, blank or matrix spike extract is transferred to the top frit of the appropriate Florisil cartridge.
- 7.3.4.7 Because the volumes marked on concentrator tubes are not necessarily accurate at the 1-mL level, the use of a syringe or a volumetric pipet is required to transfer the extract to the cleanup cartridge.
- 7.3.4.8 The pesticides/Aroclors in the extract concentrates are then eluted through the column with 9 mL of hexane/acetone (90:10) and are collected into the 10-mL volumetric flasks held in the rack inside the vacuum manifold.
- 7.3.4.9 Transfer the eluate in each volumetric flask to a clean centrifuge tube or 10-mL vial. Use two additional 1-mL hexane rinses to ensure quantitative transfer of the cartridge eluate.
- 7.3.4.10 Concentrate the extract to 1.0 or 2.0 mL as required in paragraph 7.3.4.3 by using either nitrogen blowdown (7.3.2) or a micro-Snyder column. Measure the final volume with a syringe or by transferring the extract to a volumetric flask.
- 7.3.4.11 Sulfur contamination will cause a rise in the baseline of the chromatogram that may interfere with the analyses of the later eluting pesticides. If crystals of sulfur are evident or if the presence of sulfur is suspected, proceed to 7.4. Sample analyses showing the presence of sulfur are not acceptable and must be cleaned up and reanalyzed.
- 7.3.4.12 If sulfur is not present, transfer the sample to a GC vial and label the vial. The extract is ready for GC/ECD analysis. Proceed to Section III. Store the extracts at 4°C in the dark.

7.4 Sulfur Removal

Sulfur can be removed by one of two methods, according to laboratory preference. Interference which is due to sulfur is not acceptable. If the sulfur concentration is such that crystallization occurs in the concentrated extract, centrifuge the extract to settle the sample extract with a disposable pipette, leaving the excess sulfur in the centrifuge tube. Transfer the extract to a clean centrifuge tube or clean concentrator tube before proceeding with further sulfur cleanup.

7.4.1 If only part of a set of samples requires sulfur cleanup, then two method blanks are required for that set: one that is shaken with mercury or copper, and one that is not.

Sulfur cleanup blank - add 1.0 mL of surrogate to 10 mL of hexane in a clean centrifuge tube or 10-mL vial. Concentrate the solution to 2.0 mL by using either nitrogen blowdown or a micro Snyder column. The concentrated volume of the blank must be the same as the final volume of the samples associated with the blank. Measure the volume with a syringe or by transferring the solution to a volumetric flask. Proceed with the sulfur removal using the same technique (mercury or copper) as the samples associated with the blank.

7.4.2 Mercury technique

Add one to three drops of mercury to each hexane extract in a clean vial. Tighten the top on the vial and agitate the sample for 30 seconds. Filter or centrifuge the extract. Pipet the extract to another vial and leave all solid precipitate and liquid mercury. If the mercury appears shiny, proceed to Section III and analyze the extract. If the mercury turns black, repeat sulfur removal as necessary. CAUTION: Waste containing mercury should be segregated and disposed of properly.

NOTE: Mercury is a highly toxic metal and therefore must be used with great care. Prior to using mercury, it is recommended that the analyst become acquainted with proper handling and cleanup techniques associated with this metal.

7.4.3 Copper technique

Add approximately 2 g of cleaned copper powder to the extract in the centrifuge or concentrator tube. (2 g will fill the tube to about the 0.5 mL mark). Mix the copper and extract for at least 1 minute on a mechanical shaker. Separate the extract from the copper powder by drawing off the extract with a disposable pipet, and transfer the extract to a clean vial. The extract transferred to the vial still represents the 2.0 mL final volume. The separation of the extract from the copper powder is necessary to prevent degradation of the pesticides. If the copper appears bright, proceed to Section III and analyze the extract. If the copper changes color, repeat the sulfur removal procedure as necessary. SECTION III

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GC/EC ANALYSIS OF PESTICIDES AND AROCLORS

SECTION III

GC/EC ANALYSIS OF PESTICIDES AND AROCLORS

- 1. <u>Summary of GC/EC Analysis</u>
- 1.1 The analysis of samples is accomplished by using a wide-bore (0.53 mm ID) fused silica capillary column.
- 1.2 Sample extracts, standards, and blanks must be analyzed within an analytical sequence as defined in 5. GC/ECD analysis begins with an initial demonstration of instrument performance and the calibration of all pesticides and Aroclors. Acceptable initial calibration is defined in paragraph 6. Initial calibration must be repeated whenever the calibration verification stipulated in paragraph 7 fails, or when major instrument maintenance or modification is performed.
- 1.3 An instrument blank, a Performance Evaluation Mixture, and a second instrument blank and the midpoint concentration of Individual Standard Mixtures A and B are analyzed no less than once in every 12 hour analytical sequence in order to monitor retention times, calibration factors, and column performance. Data can be collected only as long as the results for these standards and instrument blanks fall within the limits defined in paragraph 7. If two consecutive unacceptable standards are run, all extracts run since the previous acceptable standard must be reanalyzed. Additional standards and blanks are recommended when highly contaminated samples are suspected.
- 1.4 Calibration and run sequence specifications of the GC/EC method apply independently to both GC columns.
- 1.5 Matrix spike and a matrix spike duplicate analyses must be prepared and analyzed at least once for each matrix type or once per Sample Delivery Group (SDG), whichever is most frequent.
- 1.6 Analysis of a sample on both GC columns is required for all samples, blanks, matrix spikes, and matrix spike duplicates.
- 1.7 A single component pesticide is identified if a peak is detected within its appropriate retention time window on each of two columns. Toxaphene and Aroclors are identified primarily by pattern recognition, but RTs of three to five major peaks must also be taken into consideration. Guidance on quantitation of Aroclors is given beginning at 13.9.
- 1.8 Standards for all tentatively identified Aroclors must be run within 72 hours of the sample analysis in which they were observed. These standards are used to verify identification only; quantitation is based on the standard analyzed during initial calibration.
- 1.9 Quantitative analysis of pesticides/Aroclors must be accomplished by the external standard method. Three-point calibration curves for single component analytes and the surrogates must be generated during

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the initial calibration. A linear response range must be demonstrated from the CRQL to a high point at least 30 times greater than the CRQL. Single-point calibrations for multicomponent analytes are sufficient for quantitation by this method.

- 1.10 The ECD response for single component analytes must be within the three-point calibration range in order for quantitative measurements to be made. The ECD response for the Aroclors/toxaphene must not be larger than the response for the high point calibration analysis of the single component analytes. The extracts must be diluted if the ECD response exceeds the calibration range. Quantitation must be performed and reported for both GC columns.
- 1.11 Absolute retention times (RTs) are used for the identification of pesticides/Aroclors. The absolute retention time window is calculated during initial calibration as ± 0.04 minutes of the mean RT of the standard for all analytes except Methoxychlor, where the RT window is ± 0.05 minutes of the mean RT of the standard. For the multicomponent analytes, the absolute retention time window is calculated for each peak (3 to 5) as ± 0.04 minutes of the RT of the same peak in the initial calibration standard analysis.

2. <u>Gas Chromatograph/Election Capture Detector (GC/EC)</u>

- 2.1 Gas Chromatograph
 - 2.1.1 The gas chromatograph (GC) system must adequately regulate temperature in order to give a reproducible temperature program and have a flow controller that maintains a constant column flow rate throughout temperature program operations. The system must be suitable for splitless injection and have all required accessories including syringes, analytical columns, and gases.
 - 2.1.2 Gas chromatographs that are available from some manufacturers may have difficulty in meeting certain method QC requirements because of Endrin and DDT breakdown in the injector. This problem can be minimized by operating the injector at 200-205°C, using a <u>Pyrex</u> (not quartz) methyl silicone deactivated injector liner, and deactivating any metal parts in the injector with dichlorodimethyl silane. In some cases, using a 1/4 inch packed column injector converted for use with 0.53 mm capillary columns works better than a Grob-type injector. If a Grob-type injector is used, a 4 mm liner may be required to meet breakdown criteria.
- 2.2 Gas Chromatograph Columns
 - 2.2.1 Two wide-bore (0.53 mm ID) fused silica GC columns are required. A separate detector is required for each column. The specified analytical columns are a DB-1701, 30 m x 0.53 mm ID, 1.0 um film thickness, (J&W Scientific, Folsom, CA, or equivalent), and a DB-608, 30 m x 0.53 mm ID, 0.5 to 1.0 um film thickness (J&W Scientific, or equivalent). Equivalent

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columns may be employed if they meet the requirements for resolution, initial calibration, and calibration verification listed in this section.

- 2.2.2 Columns are mounted in 0.25-inch injector ports by using glass adapters available from a variety of commercial sources (J&W Scientific, Supelco, Inc., Hewlett-Packard, Varian, Inc., Perkin Elmer, or equivalent). The two columns may be mounted into a single injection port with a tee adapter (Supelco, Inc., Bellefonte, PA, Catalog No. 2-3660, or equivalent). Use of this adapter allows simultaneous injection onto both columns. The laboratory should follow manufacturer's recommendations for mounting 0.53 mm capillary columns in injector ports.
- 2.3 The carrier gas for routine applications is helium. Laboratories may choose to use hydrogen as a carrier gas, but they must clearly identify its use in the SDG Narrative and on all divider pages preceding raw chromatographic data in submissions to the Agency. Laboratories that choose to use hydrogen are advised to exercise caution in its use. Use of a hydrogen leak detector is highly recommended when hydrogen is used as the carrier gas. All GC carrier gas lines must be constructed from stainless steel or copper tubing. Non-polytetrafluoroethylene (PTFE) thread sealants or flow controllers with rubber components are not to be used.
- 2.4 Electron Capture Detector the makeup gas must be P-5, P-10 (argon/methane) or nitrogen according to the instrument specification. The GC/ECD system must be in a room in which the atmosphere has been demonstrated to be free of all contaminants which may interfere with the analysis. The instrument must be vented to outside the facility or to a trapping system which prevents the release of contaminants into the instrument room.
- 2.5 Data System a data system must be interfaced to the GC/ECD. The data system must allow the continuous acquisition of data throughout the duration of the chromatographic program and must permit, at the minimum, the output of time vs. intensity (peak height or peak area) data. Also, the data system must be able to rescale chromatographic data in order to report chromatograms meeting the requirements listed within this method.
- 3. <u>Calibration Standards</u>
- 3.1 Resolution Check Mixture prepare the mixture of pesticides in hexane or iso-octane at the concentrations listed below. The mixture must be prepared every six months, or sooner if the solution has degraded or concentrated.

gamma-Chlordane	10.0 ng/mL	Endrin ketone	20.0 ng/mL
Endosulfan I	10.0 ng/mL	Methoxychlor	100.0 ng/mL
p,p'-DDE	20.0 ng/mL	Tetrachloro-m-xylene	20.0 ng/mL
Dieldrin	20.0 ng/mL	Decachlorobiphenyl	20.0 ng/m
Endosulfan sulfate	20.0 ng/mL	_	.

3.2 Performance Evaluation Mixture (PEM) - prepare the PEM in hexane or iso-octane at the concentration levels listed below. The PEM must be prepared weekly, or more often if the solution has degraded or concentrated.

gamma - BHC	10.0 ng/mL	Endrin	50.0 ng/mL
alpha-BHC	10.0 ng/mL	Methoxychlor	250.0 ng/mL
4.4'-DDT	100.0 ng/mL	Tetrachloro-m-xylene	20.0 ng/mL
beta-BHC	10.0 ng/mL	Decachlorobiphenyl	20.0 ng/mL

3.3 Individual Standard Mixtures A and B - the single component pesticide standards must be prepared in hexane or iso-octane at three concentrations for each analyte, including the surrogates. Two separate calibration mixtures, A and B, (listed below) are used to ensure that each peak is adequately resolved. The low point concentration corresponds to the CRQL for each analyte. The midpoint concentration must be 10 times the low point concentration. The high point concentration must be at least 30 times that of the low point, but a higher concentration may be chosen by the Contractor. The high point concentration defines the upper end of the concentration range for which the concentration is valid. The solution must be prepared every six months, or sooner if the solution has degraded or concentrated.

Individual Standard Mixture A - Low Point Concentration

alpha-BHC	5.0 ng/	'mL
Heptachlor	5.0 ng/	
gamma-BHC	5.0 ng/	
Endosulfan I	5.0 ng/	
Dieldrin	10.0 ng/	
Endrin	10.0 ng/	
P,P'-DDD	10.0 ng/	
P,P'-DDT	10.0 ng/	
Methoxychlor	50.0 ng/	
Tetrachloro-m-xylene	20.0 ng/	
Decachlorobiphenyl	20.0 ng/	

Individual Standard Mixture B- Low Point Concentration

beta-BHC delta-BHC Aldrin Heptachlor expoxide alpha-Chlordane gamma-Chlordane p,p'-DDE Endosulfan sulfate	5.0 5.0 5.0 5.0 10.0 10.0	ng/mL ng/mL ng/mL ng/mL ng/mL ng/mL ng/mL
	10.0	ng/mL
Endrin aldehyde		ng/mL ng/mL
Endrin ketone		ng/mL
Endosulfan II		ng/mL
Tetrachloro-m-xylene Decachlorobiphenyl		ng/mL ng/mL
200401101001010101	20.0	ш <u>к/ш</u> г

3.4 Multicomponent Standards - Toxaphene and Aroclor standards must be prepared individually except for Aroclor 1260 and Aroclor 1016 which may be combined in one standard mixture. The calibration standards for the Aroclors must be prepared at concentrations of 100 ng/mL, except for Aroclor 1221 which must be prepared at 200 ng/mL. Toxaphene must be prepared at 500 ng/mL. All multicomponent standards must contain the surrogates at 20.0 ng/mL. The Aroclor and Toxaphene solutions must be prepared in hexane or iso-octane. Each solution must be prepared every six months, or sooner if the solution has degraded or evaporated.

4. <u>Gas Chromatograph Operating Conditions</u>

The following are the gas chromatographic analytical conditions. The conditions are recommended unless otherwise noted.

Carrier Gas:	Helium (Hydrogen may be used, see 2.3)
Column Flow:	5 mL/min
Make-up Gas:	P-5/P-10 or N ₂ (required)
Injector Temperature:	\geq 200°C (see 4.1)
Injection:	On-column
Injection Volume:	1 or 2 uL (see 4.2)
Injector:	Grob-type, splitless
Initial Temperature:	150°C
Initial Hold Time:	1/2 min
Temperature Ramp:	5°C to 6°C/min
Final Temperature:	275°C
Final Hold Time:	Until after Decachlorobiphenyl has eluted (approximately 10 minutes)

Optimize GC conditions for analyte separation and sensitivity.

- 4.1 Manual injections must be 2.0 uL. Auto injectors may use 1.0 uL volumes. The same injection volume <u>must</u> be used for all standards, blanks, and samples.
- 4.2 Cold (ambient temperature) on-column injectors that allow injection directly onto a 0.53 mm ID column may be used as long as the acceptance criteria for resolution, calibration, and analyte breakdown are met.

5. Analysis Sequence for Standards and Samples

5.1 All acceptable samples must be analyzed within a valid analysis sequence as given below.

Time	Injection #	Material Injected
	1 - 15	First 15 steps of the Initial Calibration
0 hr.	16	Instrument Blank at end of Initial Calibration
	17	PEM at end of Initial Calibration
	18	First Sample
	0	
	0	Subsequent Samples
	0	
12 hr.	0	Last sample
	lst injection past 12:00 hr.	Instrument Blank
	2nd and 3rd	Individual Standard Mixtures A and B
	injections	
	past 12:00 hr.	
	0	Sample
	0	
	0	Subsequent Samples
	0	
	0	
Another 12 hr.	0	Last Sample
	<pre>1st injection past 12:00 hr.</pre>	Instrument Blank
	2nd injection	Performance Evaluation Mixture
	0	Sample
	0	
	0	Subsequent Samples
	0	
1 10 h	0	
Another 12 hr.	0 let inication	Last Sample
	lst injection past 12:00 hr.	Instrument Blank
	2nd and 3rd injections past 12 hr.	Individual Standard Mixtures A and B
	0	Sample
	0	
	0	Subsequent Samples
	0	
	0	
	etc.	

NOTE: The first 12 hours are counted from the injection #16 (the Instrument Blank at the end of the initial calibration sequence), not from injection #1. Samples may be <u>injected</u> until 12:00 hours have elapsed. All subsequent 12-hour periods are timed from the injection of the instrument blank that bracketts the front end of the samples. Because the 12-hour time period is timed from injection of the instrument blank until the <u>injection</u> of the last sample each 12-hour period may be separated by the length of one chromatographic run, that of the analysis of the last sample. While the 12-hour period may not be exceeded, the laboratory may run instrument blanks and standards <u>more</u> frequently, for instance to accomodate staff working on 8-hour shifts.

- 5.2 Before any samples are analyzed, it is necessary for the Contractor to complete an acceptable initial calibration sequence (see paragraph 6).
- 5.3 After the initial calibration, the analysis sequence may continue as long as acceptable instrument blanks, Performance Evaluation Mixtures, and Individual Standard Mixtures A and B are analyzed at the required frequency (see paragraph 7). This analysis sequence shows only the minimum required blanks and standards. More blanks and standards may be run at the discretion of the Contractor; these must also satisfy the criteria presented in paragraph 7 in order to continue the run sequence.
- 5.4 An analysis sequence must also include all required matrix spike/matrix spike duplicate analyses and method blanks, but the Contractor may decide at what point in the sequence they are to be analyzed.
- 5.5 A standard of any identified Aroclor must be run within 72 hours of its detection in a sample chromatogram.

6. Initial Calibration

- 6.1 Initial Calibration Sequence
 - 6.1.1 Before any samples are analyzed, it is necessary for the Contractor to complete the initial calibration sequence given below.

NOTE: Steps 16 and 17 are used as part of the calibration verification as well (see paragraph 7).

INITIAL CALIBRATION SEQUENCE

- 1. Resolution Check
- 2. Performance Evaluation Mixture
- 3. Aroclor 1016/1260
- 4. Aroclor 1221
- 5. Aroclor 1232
- 6. Aroclor 1242
- 7. Aroclor 1248
- 6. Aroclor 1254
- 9. Toxaphene
- 10. Low Point Standard A
- 11. Low Point Standard B
- 12. Midpoint Standard A
- 13. Midpoint Standard B
- 14. High Point Standard A
- 15. High Point Standard B
- 16. Instrument Blank
- 17. Performance Evaluation Mixture
- 6.1.2 Samples may be analyzed only after the initial calibration acceptance criteria (6.2) are met. Otherwise, the analytical system is not functioning adequately for use with this protocol.
- 6.1.3 The initial calibration may continue to be used as long as the analytical system remains under control. The proof that the analytical system is under control is provided by the analyses of the Performance Evaluation Mixtures. If those analyses do not meet the criteria described in paragraph 7, appropriate corrective action must be taken, and the initial calibration sequence must be repeated. The calibration sequence must also be repeated if any major change in instrument hardware or instrument parameters is made (e.g., if a new column is installed or if the detector temperature is changed).
- 6.2 Initial Calibration Acceptance Criteria
 - 6.2.1 The initial calibration sequence must be analyzed in the order listed in paragraph 6.1 using the GC/ECD operating conditions described in paragraph 4. The standards must be prepared according to paragraph 3. Calculate the calibration factors and retention times according to paragraphs 8-10.

- 6.2.2 The resolution criterion is that the height of the valley between two adjacent peaks in the Resolution Check Mixture must not be greater than 60% of the height of the shorter peak. The poorest resolution on the DB-608 column probably will be between DDE and Dieldrin, between Methoxychlor and Endrin ketone and between Endosulfan I and gamma-Chlordane. On the DB-1701 column, resolution difficulties most frequently occur between Endosulfan I and gamma-Chlordane, and between Methoxychlor and Endosulfan Sulfate.
- 6.2.3 The breakdown of DDT and Endrin in both of the Performance Evaluation Mixtures must be less than 20.0 percent, and the combined breakdown of DDT and Endrin must be less than 30.0 percent where

EQ.1

% Breakdown DDT - Amount found in ng (DDD+DDE) * 100 Amount in ng of DDT injected

EQ.2

% Breakdown Endrin Amount found in ng (Endrin aldehyde + Endrin ketone) * 100
Amount of Endrin injected in ng

EQ.3

Combined % Breakdown = %Breakdown DDT + %Breakdown Endrin

- 6.2.4 All peaks in both of the Performance Evaluation Mixtures must be 100 percent resolved on both columns.
- 6.2.5 The absolute retention times of each of the single component pesticides and surrogates in both of the PEMs must be within ± 0.02 minutes of their mean absolute retention time determined from the three-point initial calibration, except Methoxychlor. The absolute retention time for Methoxychlor in both of the PEMs must be within ± 0.025 minutes of its mean absolute retention time determined from the three-point initial calibration.
- 6.2.6 The relative percent difference of the calculated amount and the true amount for each of the single component pesticides and surrogates in both of the PEMs must be less than or equivalent to 25.0 percent, using equation 4.
- 6.2.7 At least one chromatogram from each of the two Individual Standard Mixtures A and B, run during the initial calibration, must yield peaks that give recorder deflections of 50 to 100 percent of full scale.
- 6.2.8 The retention time of each single component pesticide in each of the Individual Standard Mixtures must be within ±0.02 minutes of the mean RT calculated for that compound.

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- 6.2.9 The RT of each single component pesticide in each of the three calibration standards must be within 0.02 minutes of the mean RT calculated for that compound.
- 6.2.10 The resolution between any two adjacent peaks in the midpoint concentrations of Individual Standard Mixtures A and B in the initial calibration must be greater than or equal to 90.0 percent.
- 6.2.11 The % RSD of the calibration factors for each single component analyte and surrogate must be less than 10.0 percent, except for alpha-BHC, beta-BHC, gamma-BHC, DDT, Endrin, and Methoxychlor, where the % RSD must be less than 15.0 percent.

6.3 Corrective Action.

- 6.3.1 If the technical acceptance criteria for the initial calibration are not met, inspect the system for problems. It may be necessary to change the column, bake out the detector, clean the injection port, or take other corrective actions to achieve the acceptance criteria.
- 6.3.2 Contamination should be suspected as a cause if the detector cannot achieve acceptable linearity using this method. In the case of light contamination, baking out the detector at an elevated temperature (350°C) should be sufficient to achieve acceptable performance. In the case of heavy contamination, passing hydrogen through the detector 1-2 hours at an elevated temperature may correct the problem. In the case of severe contamination, the detector may require servicing by the ECD manufacturer. DO NOT OPEN THE DETECTOR. THE ECD CONTAINS RADIOCHEMICAL SOURCES.
- 6.3.3 If a laboratory cleans out a detector using an elevated temperature, the ECD electronics must be turned off during the bake out procedure.
- 6.3.4 After bake out or hydrogen reduction, the detector must be recalibrated using the initial calibration sequence.
- 6.3.5 Initial calibration technical acceptance criteria MUST be met before any samples or required blanks are analyzed. Any samples or required blanks analyzed after the initial calibration criteria have not been met will require reanalysis at no additional cost to the Agency.

7. <u>Galibration Verification</u>

7.1 Three types of analyses are used to verify the calibration and evaluate instrument performance. The analyses of instrument blanks, Performance Evaluation Mixtures (PEM), and the mid point concentration of Individual Standard Mixtures A and B constitute the continuing calibration. Sample data are not acceptable unless bracketed by acceptable analyses of instrument blanks, PEM, and both Individual Standard Mixtures A and B.

- 7.2 An instrument blank and the Performance Evaluation Mixture must bracket one end of a 12-hour period during which sample data are collected, and a second instrument blank and the the mid point concentration of Individual Standard Mixtures A and B must bracket the other end of the 12-hour period.
- 7.3 For the 12-hour period immediately following the initial calibration sequence, the instrument blank and the PEM that are the last two steps in the initial calibration sequence bracket the front end of that 12hour period. The injection of the instrument blank starts the beginning of that 12-hour period (see paragraph 5.1). Samples may be <u>injected</u> for 12 hours from the injection of the instrument blank. The three injections <u>immediately after</u> that 12-hour period must be an instrument blank, Individual Standard Mixture A, and Individual Standard Mixture B. The instrument blank must be analyzed first, before either standard. The Individual Standard Mixtures may be analyzed in either order (A,B or B,A).
- 7.4 The analyses of the instrument blank and Individual Standard Mixtures A and B immediately following one 12-hour period may be used to begin the subsequent 12-hour period, provided that they meet the acceptance criteria in paragraphs 7.8-7.14. In that instance, the subsequent 12hour period must be bracketed by the acceptable analyses of an instrument blank and a PEM, in that order. Those two analyses may in turn be used to bracket the front end of yet another 12-hour period. This progression may continue every 12 hours until such time as any of the instrument blanks, PEMs, or Individual Standard Mixtures fails to meet the acceptance criteria in paragraphs 7.8-7.14. The 12-hour time period begins with the injection of the instrument blank. Standards (PEM or Individual Standard Mixtures), samples and required blanks may be injected for 12:00 hours from the time of injection of the instrument blank.
- 7.5 If more than 12 hours have elapsed since the injection of the instrument blank that bracketed a previous 12-hour period, an acceptable instrument blank and PEM <u>must</u> be analyzed in order to start a new sequence. This requirement applies even if no analyses were performed since that standard(s) was injected.
- 7.6 After a break in sample analyses, the laboratory may only resume the analysis of samples using the current initial calibration for quantitation by analyzing an acceptable instrument blank and a PEM.
- 7.7 If the entire 12-hour period is not required for the analyses of all samples to be reported and all data collection is to be stopped, the incomplete sequence <u>must</u> be ended with either the instrument blank/PEM combination or the instrument blank/Individual Standard Mixtures A and B combination, whichever was due to be performed at the end of 12-hour period.

- 7.8 All single component pesticides and surrogates in the Performance Evaluation Mixtures used to demonstrate continuing calibration must be 100 percent resolved. The resolution between any two adjacent peaks in the midpoint concentrations of Individual Standard Mixtures A and B in the initial calibration must be greater than or equal to 90.0 percent.
- 7.9 The absolute retention time for each of the single component pesticides (except methoxychlor) and surrogates in the PEMs and mid point concentration of the Individual Standard Mixtures used to demonstrate continuing calibration must be within \pm 0.02 minutes of their mean absolute retention times determined from the three-point initial calibration. The absolute retention time for Methoxychlor in the PEM and Individual Standard Mixture must be within \pm 0.025 minutes of its mean absolute retention time determined from the three-point initial calibration. NOTE: This is half the retention time window described in paragraph 8.4.
- 7.10 The relative percent difference of the calculated amount and the true amount for each of the single component pesticides and surrogates in the PEM and mid point concentration of the Individual Standard Mixtures used to demonstrate continuing calibration must be less than or equal to 25.0 percent, using Equation 4.

$$RPD = \frac{C_{nom} - C_{calc}}{C_{nom}} \times 100$$
 EQ. 4

C_{nom} - nominal concentration of each analyte

C_{calc} - calculated concentration of each analyte from the analyses of the standard

- 7.11 The percent breakdown of DDT and Endrin in the PEM must be less than or equal to 20.0 percent on <u>both</u> columns. The combined breakdown of DDT and Endrin must be less than or equal to 30.0 percent on <u>both</u> columns.
- 7.12 All instrument blanks must meet the acceptance criteria in paragraph 15.3.
- 7.13 Analysts are cautioned that running an instrument blank and a performance evaluation mixture once every 12 hours is the minimum contract requirement. Late eluting peaks may carry over from one injection to the next if highly complex samples are analyzed or if the GC conditions are unstable. Such carryover is unacceptable. Therefore, it may be necessary to run instrument blanks and performance evaluation mixtures more often to avoid discarding data.
- 7.14 The requirements for running the instrument blanks, Performance Evaluation Mixture, and Individual Standard Mixtures A and B are waived when no samples, method blanks, or matrix spikes are run during that 12-hour period. After a break in sample analysis, a laboratory may resume the analysis of samples, method blanks, and matrix spikes and may use the current initial calibration for quantitation <u>only</u> after an

acceptable PEM is run (paragraphs 7.2 - 7.6). If a successful PEM cannot be run after an interruption, an acceptable initial calibration must be run before sample data may be collected. All acceptable sample analyses must be bracketed by acceptable performance evaluation mixtures and instrument blanks.

8. <u>Determination of Absolute Retention Times.</u>

- 8.1 During the initial calibration sequence, absolute retention times (RT) are determined for all single response pesticides, the surrogates, and at least three major peaks of each multicomponent analyte.
- 8.2 For single component pesticides, an RT is measured in each of three calibration standards and the mean RT is calculated as the average of the three values. An RT is measured for the surrogates in each of the three analyses of Individual Mixture A during the initial calibration and the mean RT is calculated as the average of the three values. The RT of each single component pesticide in each of the three calibration standards must be within 0.02 minutes of the mean RT calculated for that compound.
- 8.3 A retention time window is calculated for each single component analyte and surrogate by using the list in paragraph 8.4. Windows are centered around the mean absolute retention time for the analyte established during the initial calibrations.

8.4 Retention time windows for single and multicomponent analytes and surrogates.

Compound	Retention Time Window in Minutes
alpha-BHC	± 0.04
beta-BHC	± 0.04
gamma-BHC	± 0.04
delta-BHC	± 0.04
Heptachlor	± 0.04
alpha-Chlorodane	± 0.04
gamma-Chlorodane	± 0.04
Aldrin	± 0.04
Heptachlor epoxide	± 0.04
Dieldrin	± 0.04
Endrin	± 0.04
Endrin aldehyde	± 0.04
Endrin ketone	± 0.04
DDD	± 0.04
DDE	± 0.04
DDT	± 0.04
Endosulfan I	± 0.04
Endosulfan II	± 0.04
Endosulfan sulfate	± 0.04
Methoxychlor	± 0.05
Aroclors	± 0.04
Toxaphene	± 0.04
Tetrachloro-m-xylene	± 0.04
Decachlorobiphenyl	± 0.04

- 8.5 For each multicomponent analyte, the RTs for three to five peaks are calculated from the initial calibration standard analysis. An RT window of ±0.04 minutes is used for all multicomponent analyte peaks.
- 8.6 Analytes are identified when peaks are observed in the RT window for the compound on both GC columns.
- 9. <u>Calibration Factors for Single Component Pesticides.</u>
- 9.1 During the initial calibration sequence, the Contractor must establish the magnitude of the linear ECD response range for each single component pesticide and surrogate on each column and for each GC system. This is accomplished by analyzing the Individual Standard Mixtures A and B at three concentrations during the initial calibration sequence in paragraph 6.
- 9.2 The linearity of the instrument is determined by calculating a percent relative standard deviation (%RSD) of the calibration factors from a three-point calibration curve for each single component pesticide and surrogate. Either peak area or peak height may be used to calculate calibration factors used in the %RSD equation. For example, it is permitted to calculate linearity for Endrin based on peak area and to calculate linearity for Aldrin based on peak height. It is not

permitted within a %RSD calculation for an analyte to use calibration factors calculated from both peak area and peak height. For example, it is not permitted to calculate the calibration factor for the low point standard for endrin using peak height and calculate the midpoint and high point standard calibration factors for endrin using peak area.

- 9.2.1 Calculate the calibration factor for each single component pesticide and surrogate over the initial calibration range using Equation 5.
- 9.2.2 Calculate the mean and the *RSD of the calibration factors for each single component pesticide and surrogate over the initial calibration range using Equations 6 and 7.

$$\overline{CF} - \sum_{i=1}^{n} \frac{CF_i}{n}$$

* RSD - SD
$$\overline{CF}$$
 x 100

EQ. 7

EQ. 6

Where
$$SD = \sqrt{\frac{\sum_{i=1}^{n} (CF_{i} - \overline{CF})^2}{n-1}}$$
 and n-3

- 9.2.3 The linearity of the calibration is considered acceptable when the % RSD of the three point calibration is less than 10.0 percent for all single component pesticides except alpha-BHC, beta-BHC, gamma-BHC, DDT, Endrin, and Methoxychlor, where the % RSD must be less than 15.0 percent.
- 9.2.4 If the linearity requirements listed above are met, the calibration factor from the mid point concentration standard is used for quantitation of each single component pesticide.
- 9.3 Sample analysis may not proceed until a satisfactory calibration has been demonstrated.

10. <u>Calibration Factors for Toxaphene and Aroclors</u>

- 10.1 Toxaphene and Aroclors require only a single-point calibration and they present special analytical difficulties. Because of the alteration of these materials in the environment, it is probable that samples which contain multicomponent analytes will give patterns similar to, but not identical with, those of the standards.
- 10.2 A set of three to five major peaks is selected for each multicomponent analyte. Retention times (see 8.4) and calibration factors are

determined from the initial calibration analysis for each peak. Guidance for the choice of which peaks to use is given in paragraph 13.9

11. Acceptance Criteria for Chromatograms of Calibration Standards

The identification of single component pesticides by gas chromatographic methods is based primarily on rentention time data. The retention time of the apex of a peak can be verified only from an on-scale chromatogram. The identification of multicomponent analytes is based primarily on recognition of patterns of retention times displayed on a chromatogram. Therefore, the following requirements apply to all data presented for single component and multicomponent analytes.

- 11.1 The chromatograms that result from the analyses of the Resolution Check Mixture, the Performance Evaluation Mixture, and Individual Standard Mixtures A and B during the initial calibration sequence must display the single component analytes present in each standard at greater than 10 percent of full scale but less than 100 percent of full scale.
- 11.2 The chromatograms, for at least one of the three analyses each of Individual Standard Mixtures A and B from the initial calibration sequence, must display the single component analytes at greater than 50 percent and less than 100 percent of full scale.
- 11.3 The chromatograms of the standards for the multicomponent analytes analyzed during the initial calibration sequence must display the peaks chosen for identification of each analyte at greater than 25 percent and less than 100 percent of full scale.
- 11.4 For any standard containing alpha-BHC, the baseline of the chromatogram must return to below 50 percent of full scale before the elution time of alpha-BHC, and return to below 25 percent of full scale after the elution time of alpha-BHC and before the elution time of decachlorobiphenyl.
- 11.5 If a chromatogram is replotted electronically to meet requirements, the scaling factor used must be displayed on the chromatogram.
- 11.6 If the chromatogram of any standard needs to be replotted electronically to meet these requirements, both the initial chromatogram and the replotted chromatogram must to submitted in the data package.
- 12. <u>Sample Analysis</u>
- 12.1 Unless ambient temperature on-column injection is used (see paragraph 4.2), the injector must be heated to at least 200°C. The gas chromatographic conditions from paragraph 4 must be used.
- 12.2 The injection must be made on-column by using either automatic or manual injection. If autoinjectors are used, 1.0 uL injection volumes may be used. Manual injections shall use at least 2.0 uL injection

volumes. The same injection volume must be used for all standards, samples, and blanks associated with the same initial calibration.

- 12.3 Analysis of a sample on both GC columns is required for <u>all</u> samples, blanks, matrix spikes, and matrix spike duplicates.
- 12.4 The requirements for the analysis sequence apply to both GC columns and for all instruments used for these analyses.
- 12.5 The laboratory will identify and quantitate analyte peaks based on RT and calibration factor established during the initial calibration sequence, as long as an acceptable calibration verification (see paragraph 7) is performed every 12 hours.
- 12.6 The protocol is intended to achieve the quantitation limits shown in Exhibit C whenever possible. If sample chromatograms have interfering peaks, a high baseline, or off-scale peaks, then those samples must be reanalyzed following dilution, further cleanup, or reextraction. Samples which cannot be made to meet the given specifications after one reextraction and three-step cleanup (GPC, Florisil, and sulfur removal) are reported in the SDG Narrative and do not require further analysis. No limit is placed on the number of reextractions of samples that may be required because of contaminated method blanks.
- 12.7 The sample must be analyzed at the most concentrated level that is consistent with achieving satisfactory chromatography (defined below). If dilution is employed solely to bring a peak within the calibration range or to get a multicomponent pattern on scale, the results for both the more and the less concentrated extract must be reported. The resulting changes in quantitation limits and surrogate recovery must be reported also for the diluted samples.
- 12.8 If the Contractor has reason to believe that diluting the final extracts will be necessary, an undiluted run may not be required. If an acceptable chromatogram (as defined below) is achieved with the diluted extract, an additional extract 10 times the concentration of the dilute sample must be injected and reported with the sample data.
- 12.9 No target analyte concentrations may exceed the upper limit of the initial calibration.
- 12.10 A standard for any identified multicomponent analyte must be analyzed on the same instrument within 72 hours of its detection in a sample.
- 12.11 The identification of single component pesticides by gas chromatographic methods is based primarily on retention time data. The retention time of the apex of a peak can be verified only from an onscale chromatogram. The identification of multicomponent analytes is based primarily on recognition of patterns of retention times displayed on a chromatogram. Therefore, the following requirements apply to all data presented for single component and multicomponent analytes.

- 12.11.1 When no analytes are identified in a sample, the chromatograms from the analyses of the sample extract must use the same scaling factor as was used for the low point standard of the initial calibration associated with those analyses.
- 12.11.2 Chromatograms must display single component pesticides detected in the sample at less than full scale.
- 12.11.3 Chromatograms must display the largest peak of any multicomponent analyte detected in the sample at less than full scale.
- 12.11.4 If an extract must be diluted, chromatograms must display single component pesticides between 10 and 100 percent of full scale.
- 12.11.5 If an extract must be diluted, chromatograms must display the peaks chosen for quantitation of multicomponent analytes between 25 and 100 percent of full scale.
- 12.11.6 For any sample, the baseline of the chromatogram must return to below 50 percent of full scale before the elution time of alpha-BHC, and return to below 25 percent of full scale after the elution time of alpha-BHC and before the elution time of decachlorobiphenyl.
- 12.11.7 If a chromatogram is replotted electronically to meet these requirements, the scaling factor used must be displayed on the chromatogram.
- 12.11.8 If the chromatogram of any sample needs to be replotted electronically to meet these requirements, both the initial chromatogram and the replotted chromatogram must be submitted in the data package.

13. <u>Quantitation of Analytes</u>

- 13.1 Quantitation must be performed and reported on both columns.
- 13.2 Analytes must be quantitated with an electronic integrator or with a laboratory data system. The analyst can use either peak height or peak area as the basis for quantitation. The use of an electronic integrator or a laboratory data system is required.
- 13.3 The chromatograms of all samples must be reviewed by a qualified pesticide analyst before they are reported.
- 13.4 In order to be quantitated, the detector response (peak area or peak height) of all of the single component analytes must lie between the response of the low and high concentrations in the initial calibration. If the analytes are detected below the CRQL, they are reported as present below the CRQL, and flagged according to the instructions in Exhibit B. If they are detected at a level greater than the high calibration point, the sample must be diluted either to a maximum of

1:100,000 or until the response is within the linear range established during calibration. Guidance in performing dilutions and exceptions to this requirement are given below.

- 13.4.1 If the response is still above the high calibration point after the dilution of 1:100,000, the Contractor shall contact the SMO immediately.
- 13.4.2 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
- 13.4.3 The dilution factor chosen should keep the response of the largest peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.
- 13.4.4 Do <u>not</u> submit data for more than two analyses, i.e., the original sample extract and <u>one</u> dilution, or, if a screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
- 13.4.5 Do not dilute MS/MSD samples solely to get non-spiked analytes within the calibration range. The spiking level of each compound in the pesticide/Aroclor matrix spiking solution should not require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.
- 13.5 The concentrations of the single component pesticides are calculated by using the following equations:
 - 13.5.1 Water

Concentration
$$ug/L = \frac{(A_x)(V_t)(Df)}{(CF)(V_c)(V_i)}$$
 EQ. 8

Where

- A_x Area of the peak for the compound to be measured
- CF Calibration factor for the mid point concentration external standard (area per ng)
- V_o Volume of water extracted in milliliters (mL)
- V_i Volume of extract injected in microliters (uL)
- V_+ Volume of the concentrated extract in microliters (uL)
- Df Dilution Factor. The dilution factor for analysis of water samples by this method is defined as follows:

uL most conc. extract used to make dilution + uL clean solvent uL most conc. extract used to make dilution

If no dilution is performed, Df = 1.0.

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13.5.2 Soil/Sediment

Concentration ug/Kg =
$$\frac{(A_x)(V_t)(Df)}{(CF)(V_i)(W_s)(D)}$$
 EQ. 9
Where

 A_x and CF are as given for water, above.
 V_t - Volume of the concentrated extract in microliters (uL)
 V_i - Volume of extract injected in microliters (uL)
 D = 100 - % moisture 100
 W_s - Weight of sample extracted in grams (g)
 Df - Dilution Factor. The dilution factor for analysis of soil samples by this method is defined as follows:
 uL most conc. extract used to make dilution

If no dilution is performed, Df = 1.0.

- 13.5.3 Note that the calibration factors used for the quantitation of the single component pesticides are the calibration factors from the mid point concentration standard for each analyte.
- 13.6 The concentrations of the surrogates are calculated in a similar manner as the other analytes, using Equations 8. and 9. The recoveries of the surrogates are calculated according to Equation 10.

Surrogate Percent Recovery -
$$\frac{Q_d}{Q_a}$$
 x 100 EQ. 10

Where

 Q_d = Quantity determined by analysis Q_a = Quantity added to sample/blank

The limits for the recovery of the surrogates are 60-150 percent for both surrogate compounds. As these limits are only advisory, no further action is required by the laboratory is required, however, frequent failures to meet the limits for surrogate recovery warrant investigation by the laboratory, and may result in questions from the Agency.

13.7 The quantitative determination of Toxaphene or Aroclors is somewhat different from that of single component pesticides. Quantitation of peaks within the detector linear range CRQL to > 30 times CRQL is based on a single calibration point assuming linear detector response. Alternatively, a linear calibration range may be established during a run sequence by a three-point calibration curve for any multicomponent analyte.

- 13.8 The reporting requirements for multicomponent analytes are similar to those for single component analytes. If the concentration is calculated to be 10⁶ times the CRQL, the Contractor shall contact the SMO immediately.
- 13.9 The quantitation of toxaphene or Aroclors must be accomplished by comparing the heights or the areas of each of the three to five major peaks of the multicomponent analyte in the sample with the calibration factor for the same peaks established during the initial calibration sequence. The concentration of multicomponent analytes is calculated by using Equations 8 and 9, where A_x is the area for each of the major peaks of the multicomponent analyte. The concentration of each peak is determined and then an average concentration for three to five major peaks is determined and reported on Form I (Exhibit B). The following table lists the number of potential quantitation peaks for each Aroclor and Toxaphene.

<u>Analyte</u>	No. of Potential <u>Quantitation Peaks</u>	
Aroclor 1016/1260	5/5	
Aroclor 1221	3	
Aroclor 1232	4	
Aroclor 1242	5	
Aroclor 1248	5	
Aroclor 1254	5	
Toxaphene	4	

- 13.10 The choice of the peaks used for multicomponent quantitation and the recognition of those peaks may be complicated by the environmental alteration of the Toxaphene or Aroclors, and by the presence of coeluting analytes or matrix interferences, or both.
- 13.11 If more than one multicomponent analyte is observed in a sample, the Contractor must choose separate peaks to quantitate the different multicomponent analytes. A peak common to both analytes present in the sample must not be used to quantitate either compound.

14. Sample Data Acceptance Criteria

- 14.1 The requirements above apply to <u>both</u> columns, and quantitation must be performed on both GC columns and reported.
- 14.2 All samples must be be analyzed as part of a valid analysis sequence (paragraph 5). They must be bracketed by acceptable instrument blanks (paragraph 15.3), acceptable Performance Evaluation Mixtures, and acceptable Individual Standard Mixtures A and B (paragraph 7) that were analyzed at the required frequency.
- 14.3 The retention times for both of the surrogates must be within the retention time windows as calculated in paragraph 8.

- 14.4 Reportable data for a sample must include a chromatogram in which a baseline returns to below 50 percent of full scale before the elution time of alpha-BHC, and to below 25 percent of full scale after alpha-BHC and before decachlorobiphenyl.
- 14.5 If dilution has been applied and if no peaks are detected above 25 percent of full scale, analysis of a more concentrated sample is required.
- 14.6 Reportable sample data must include chromatogram(s) which meet the criteria in paragraph 12.11.

15. Blanks

There are two types of blanks required by this method: the method blank and the instrument blank. A separate sulfur cleanup blank <u>may</u> be required if all samples associated with a given method blank are not subjected to sulfur cleanup. Samples that are associated with a sulfur cleanup blank are also associated with the method blank with which they were extracted. Both the method and sulfur cleanup blanks must meet the respective acceptance criteria for the sample analysis acceptance criteria to be met.

- 15.1 Method blank
 - 15.1.1 Method blanks are spiked with the surrogate solution, extracted, cleaned up, and analyzed by following the same procedure that is used with the samples. A water method blank is one liter of reagent water treated as the water sample aliquot. A soil method blank is 30 g of sodium sulfate treated as the soil sample aliquot.

Method blank analysis must be performed once for the following, whichever is most frequent, and analyzed on each GC/EC system used to analyze samples:

- o Each Case, OR
- Each 14 calendar day period during which samples in a Case are received (said period beginning with the receipt of the first sample in that Sample Delivery Group), OR
- Each 20 samples in a Case, including matrix spikes and reanalyses, that are of similar matrix (water or soil), OR
- Whenever samples are extracted by the same procedure (separatory funnel, continuous liquid-liquid extraction, or sonication).
- 15.1.2 In order to be acceptable, a method blank analysis cannot contain any of the analytes listed in Exhibit C at greater than the CRQL. The surrogate retention times must be within the retention time windows calculated from the initial calibration sequence mean retention time for both tetrachloro-m-xylene and decachlorobiphenyl.

- 15.1.3 All samples associated with an unacceptable method blank (see Form IV) must be reextracted and reanalyzed at no additional cost to the Agency.
- 15.2 Sulfur Cleanup Blank.
 - 15.2.1 The sulfur cleanup blank is a modified form of the method blank. The sulfur cleanup blank is hexane spiked with the surrogates and passed through the sulfur cleanup procedure (see Section II, paragraph 7.4).
 - 15.2.2 The sulfur cleanup blank is prepared when only part of a set of samples extracted together requires sulfur removal. A method blank is associated with the entire set of samples. The sulfur cleanup blank is associated with the part of the set which required sulfur cleanup. If all the samples associated with a given method blank are subjected to sulfur cleanup, then the method blank must be subjected to sulfur cleanup, and <u>no</u> separate sulfur cleanup blank is required.
 - 15.2.3 In order to be acceptable, a sulfur blank analysis cannot contain any of the analytes listed in Exhibit C at greater than the CRQL. The surrogate retention times must be within the retention time windows calculated from the initial calibration sequence mean retention time for both tetrachloro-m-xylene and decachlorobiphenyl.
 - 15.2.4 All samples associated with an unacceptable sulfur blank (see Form IV) must be reextracted and reanalyzed at no additional cost to the Agency.
- 15.3 Instrument blank
 - 15.3.1 An instrument blank is a hexane or iso-octane solution containing 20.0 ng/mL of tetrachloro-m-xylene and decachlorobiphenyl.
 - 15.3.2. The first analysis in a 12-hour analysis sequence must be an instrument blank. All acceptable samples analyses are to be bracketed by acceptable instrument blanks, as described in paragraph 5.1.
 - 15.3.3 An acceptable instrument blank must be analyzed within a 12hour analysis sequence and must demonstrate that no analyte in Exhibit C is detected at greater than 0.5 times the CRQL and that the surrogate retention times are within the retention time windows.
 - 15.3.4 If analytes are detected at greater than half the CRQL or the surrogate RTs are outside the RT windows, all data collection must be stopped, and corrective action must be taken. Data for samples which were run between the last acceptable instrument blank and the unacceptable blank are considered suspect. An acceptable instrument blank must be run before additional data

are collected. After an acceptable instrument blank is run, all samples which were run after the last acceptable instrument blank must be reinjected during a valid run sequence at no additional cost to the Agency and must be reported.

- 15.3.5 Analysts are cautioned that running an instrument blank once every 12 hours is the minimum contract requirement. Late eluting peaks may carry over from one injection to the next if highly complex samples are analyzed or if the GC conditions are unstable. Such carryover is unacceptable. Therefore, it may be necessary to run instrument blanks more often to avoid discarding data.
- 16. <u>Matrix Spike/Matrix Spike Duplicate</u>
- 16.1 A matrix spike and matrix spike duplicate must be extracted and analyzed at least once with every 20 samples of each matrix. NOTE: There is no differentiation between "low" and "medium" soil samples in this method. Therefore only one soil MS/MSD is to be submitted.
- 16.2 The surrogate retention times must be within the retention time windows specified.
- 16.3 The percent recoveries and the relative percent difference between the recoveries of each of the 6 compounds in the matrix spike samples will be calculated and reported by using the following equations:

Matrix Spike Recovery =
$$\frac{SSR - SR}{SA}$$
 x 100
Eq. 11

Where

SSR - Spike sample result SR - Sample result SA - Spike added

$$RPD = \frac{|MSR - MSDR|}{\frac{1}{2}(MSR + MSDR)} \times 100$$
 Eq. 12

Where

RPD = Relative percent difference
MSR = Matrix spike recovery
MSDR = Matrix spike duplicate recovery

The vertical bars in the formula above indicate the absolute value of the difference, hence RPD is always expressed as a positive value.

16.4 The Contractor shall report matrix spike and matrix spike duplicate recoveries and percent difference values with the analytical results (see Exhibit B). The limits for matrix spike compound recovery and RPD are given below. As these limits are only advisory, no further action by the laboratory is required, however, frequent failures to meet the limits for recovery or RPD warrant investigation by the laboratory, and may result in questions from the Agency.

Compound	%R	RPD	%R	RPD
	Water	Water	Soil	Soil
gamma-BHC (Lindane)	56-123	50	46-127	15
Heptachlor	40-131	31	35-130	20
Aldrin	40-120	43	34-132	22
Dieldrin	52-126	38	31-134	18
Endrin	56-121	45	42-139	21
4,4'-DDT	38-127	50	23-134	27

MATRIX SPIKE RECOVERY AND RELATIVE PERCENT DIFFERENCE LIMITS

17. GC/MS Confirmation of Pesticides and Aroclors

- 17.1 Any pesticide or Aroclor analyte listed in Ex. C for which a concentration is reported from a GC/EC analysis must have the identification confirmed by GC/MS if the concentration is sufficient for that purpose. The following paragraphs are to be used as guidance perform GC/MS confirmation. If the the Contractor fails to reanalysis of any affected samples at no additional cost to the Agency.
 - 17.1.1 The GC/MS confirmation may be accomplished by one of three general means.
 - Examination of the semivolatile GC/MS library search results (i.e. TIC data)
 - A second analysis of the the semivolatile extract
 - Analysis of the pesticide/Aroclor extract, following any solvent exchange and concentration steps that may be necessary.
 - 17.1.2 The semivolatile GC/MS analysis procedures outlined in Ex. D SV are based on the injection into the instrument of approximately 20 ng of a target compound in a 2 uL volume. The semivolatile CRQL values in Ex. C are based on the sample concentration that corresponds to an extract concentration of 10 ng/uL of target analyte. However, these are <u>quantitation</u> limits, and the <u>detection</u> of analytes and generation of reproducible mass The sample concentration corresponding to 10 ng/uL in extract will depend on the sample matrix.
 - For water samples, 20 ng/ 2 uL corresponds to a sample concentration of 10 ug/L.

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- For soil samples prepared according to the semivolatile low level soil method (i. e. 30 g of soil), the corresponding sample concentration is 330 ug/Kg.
- For soil samples prepared according to the semivolatile medium level soil method (i. e. 1 g of soil), the corresponding sample concentration is 10,000 ug/Kg.

Therefore, based on the values given above, any pesticide sample in which the sample concentration is greater than or equal to an extract concentration of 10 ng/uL should enable the laboratory to confirm the pesticide/Aroclor by GC/MS analysis of the semivolatile extract.

- 17.1.3 In order to confirm the identification of the target pesticide/Aroclor, the laboratory must also analyze a reference standard for the analyte. In order to demonstrate the ability of the GC/MS system to identify the analyte in question, the concentration of the standard should be no greater than 10 ng/uL.
 - 17.1.3.1 To facilitate the confirmation of the pesticide/Aroclor analytes from the semivolatile library search data, the laboratory may wish to include these analytes in the semivolatile continuing calibration standard at a concentration of 10 ng/uL or less. If added to this GC/MS standard, the response factors, retention times, etc. for these analytes would be reported on the GC/MS quantitation report, but <u>not</u> on the GC/MS calibration data reporting forms. As only a single concentration of each analyte would be analyzed, no linearity (%RSD) or percent difference criteria would be applied to the response factors for these additional analytes.
 - 17.1.3.2 The laboratory is advised that library search results from the NIST/EPA/MSDC mass spectral library will not likely list the name of the pesticide/Aroclor analyte as it appears in this SOW, hence, the mass spectral interpretation specialist is advised to compare the CAS Registry numbers for the pesticides/Aroclors to those from the library search routine.
- 17.1.4 If the analyte cannot be confirmed from the semivolatile library search data for the original semivolatile GC/MS analysis, the laboratory may analyze another aliquot of the semivolatile sample extract after further concentration of the aliquot. This second aliquot must either be analyzed as part of a routine semivolatile GC/MS analysis, including instrument performance checks (DFTPP), calibration standards containing the pesticides/Aroclors as described in paragraph 17.1.3, etc.,

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or it must be analyzed along with separate reference standards for the analytes to be confirmed.

- 17.1.5 If the analyte cannot be confirmed by either the procedures in paragraphs 17.1.3 or 17.1.4, then an aliquot of the extract prepared for the GC/EC analysis must be analyzed by GC/MS, following any necessary solvent exchange and concentration steps. As in paragraph 17.1.4, analysis of a reference standard is required if the GC/MS continuing calibration standard does not contain the analyte to be confirmed.
- 17.1.6 Regardless of which of the three approaches above is used for GC/MS confirmation, the appropriate blank must also be analyzed by GC/MS to demonstrate that the presence of the analyte was not the result of laboratory contamination. If the confirmation is based on the analysis of the semivolatile extract, then the semivolatile method blank extracted with the sample must also be analyzed. If the confirmation is based on the analysis, then the pesticide method blank extracted with the sample must be analyzed.
- 17.2 If the identification of the analyte can not be confirmed by any of the GC/MS procedures above and the concentration calculated from the GC/EC analysis is greater than or equal to the concentration of the reference standard analyzed by GC/MS, then report the analyte as undetected, adjust the sample quantitation limit (the value associated with the "U" qualifier) to a sample concentration equivalent to the concentration of the GC/MS reference standard, and qualify the result on Form I with one of the laboratory-defined qualifiers ("X", "Y", or "Z"). In this instance, define the qualifier explicitly in the SDG Narrative, and describe the steps taken to confirm the analyte in the Narrative.
- 17.3 For GC/MS confirmation of single component analytes, the required deliverables are copies of the library search results (three best TIC matches) or analyte spectrum and the spectrum of the reference standard. For multicomponent analytes, spectra of three characteristic peaks are required for both the sample component and the reference standard.
- 17.4 The purpose of GC/MS analysis is for confirmation of identification, not quantitation. Therefore, the concentrations of all pesticides/Aroclors shall be based on the GC/EC results. The exception noted in paragraph 17.2 applies only to analytes that cannot be confirmed at a concentration above that of the reference standard.

EXHIBIT E

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QUALITY ASSURANCE/QUALITY CONTROL REQUIREMENTS

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OVERVIEW

Quality assurance and quality control are integral parts of the Environmental Protection Agency's (EPA) Contract Laboratory Program $(CLP)^{(1,2,4,5,6)}$. The quality assurance (QA) process consists of management review and oversight at the planning, implementation, and completion stages of the environmental data collection activity, to ensure that data provided are of the quality required. The quality control (QC) process includes those activities required during data collection to produce the data quality desired and to document the quality of the collected data⁽¹⁾.

During the planning of an environmental data collection program, QA activities focus on defining data quality criteria and designing a QC system to measure the quality of data being generated. During the implementation of the data collection effort, QA activities ensure that the QC system is functioning effectively, and that the deficiencies uncovered by the QC system are corrected. After environmental data are collected, QA activities focus on assessing the quality of data obtained to determine its suitability to support enforcement or remedial decisions (1,2,3).

The purpose of this Exhibit is to describe the overall quality assurance/quality control operations and the processes by which the Program meets the QA/QC objective defined above. This contract requires a variety of QA/QC activities. These contract requirements are the minimum QC operations necessary to satisfy the analytical requirements associated with the determination of the different method analytes. These QC operations are designed to facilitate laboratory comparison by providing the EPA with comparable data from all Contractors. These requirements do not release the analytical Contractor from maintaining their own QC checks on method and instrument performance.

SECTION I

INTRODUCTION

Appropriate use of data generated under the great range of analytical conditions encountered in environmental analyses requires reliance on the quality control procedures and criteria incorporated into the methods. The methods in this contract have been validated on samples typical of those received by the laboratories in the Contract Laboratory Program (CLP). However the validation of these methods does not guarantee that they perform equally well for all sample matrices encountered. Inaccuracies can also result from causes other than unanticipated matrix effects, such as sampling artifacts, equipment malfunctions, and operator error. Therefore, the quality control component of each method is indispensable.

The data acquired from quality control procedures are used to estimate and evaluate the information content of analytical results and to determine the necessity for or the effect of corrective action procedures. The means used to estimate information content include precision, accuracy, detection limit, and other quantitative and qualitative indicators. In addition, it gives an overview of the activities required in an integrated program to generate data of known and documented quality required to meet defined objectives.

The necessary components of a complete QA/QC program include internal QC criteria that demonstrate acceptable levels of performance, as determined by QA review. External review of data and procedures is accomplished by the monitoring activities of the National Program Office, Regional data users, Sample Management Office, NEIC, and EMSL/LV. Each external review accomplishes a different purpose. These reviews are described in specific sections of this Exhibit. Performance evaluation samples and magnetic tape audits provide an external QA reference for the program. A laboratory onsite evaluation system is also part of the various review functions to the contract laboratories through direct communications with the Administrative Project Officers and Technical Project Officers.

This Exhibit is not a guide to constructing quality assurance project plans, quality control systems, or a quality assurance organization. It is, however, an explanation of the quality control and quality assurance requirements of the program. It outlines some minimum standards for QA/QC programs. It also includes specific items that are required in a QA Plan and by the QA/QC documentation detailed in this contract. Delivery of this documentation provides the Agency with a complete data package which will stand alone, and limits the need for contact with the Contractor or with an analyst, at a later date, if some aspect of the analysis is questioned.

In order to assure that the product delivered by the Contractor meets the requirements of the contract, and to improve interlaboratory data comparison, the Agency requires the following from the Contractor:

o A written Quality Assurance Plan, the elements of which are designated in Section II.

- Written preparation of and adherence to QA/QC Standard Operating Procedures (SOPs) as described in Section III.
- Adherence to the analytical methods and associated QC requirements specified in the contract.
- o Verification of analytical standard and documentation of the purity of neat materials and the purity and accuracy of solutions obtained from private chemical supply houses.
- o Submission of all raw data and pertinent documentation for Regional review.
- o Participation in the analysis of Laboratory Evaluation Samples, including adherence to corrective action procedures.
- o Submission, upon request, of GC/MS tapes and applicable documentation for tape audits.
- o Participation in On-Site Laboratory Evaluations, including adherence to corrective action procedures.
- o Submission of all original documentation generated during sample analyses for Agency review.

SECTION II

QUALITY ASSURANCE PLAN

The Contractor shall establish a quality assurance program with the objective of providing sound analytical chemical measurements. This program shall incorporate the quality control procedures, any necessary corrective action, and all documentation required during data collection as well as the quality assessment measures performed by management to ensure acceptable data production.

As evidence of such a program, the Contractor shall prepare a written Quality Assurance Plan (QAP) which describes the procedures that are implemented to achieve the following:

- o Maintain data integrity, validity, and useability.
- o Ensure that analytical measurement systems are maintained in an acceptable state of stability and reproducibility .
- o Detect problems through data assessment and establishes corrective action procedures which keep the analytical process reliable.
- o Document all aspects of the measurement process in order to provide data which are technically sound and legally defensible.

The QAP must present, in specific terms, the policies, organization, objectives, functional guidelines, and specific QA and QC activities designed to achieve the data quality requirements in this contract. Where applicable, SOPs pertaining to each element shall be included or referenced as part of the QAP. The QAP must be available during On-Site Laboratory evaluation and upon written request by the Administrative Project Officer. Additional information relevant to the preparation of a QAP can be found in EPA and ASTM publication^(2,4).

Elements of a Quality Assurance Plan

- A. Organization and Personnel
 - 1. QA Policy and Objectives
 - 2. QA Management
 - a. Organization
 - b. Assignment of QC and QA Responsibilities
 - c. Reporting Relationships
 - d. QA Document Control Procedures
 - e. QA Program Assessment Procedures

- 3. Personnel
 - a. Resumes
 - b. Education and Experience Pertinent to This Contract
 - c. Training Progress
- B. Facilities and Equipment
 - 1. Instrumentation and Backup Alternatives
 - 2. Maintenance Activities and Schedules
- C. Document Control
 - 1. Laboratory Notebook Policy
 - 2. Samples Tracking/Custody Procedures
 - 3. Logbook Maintenance and Archiving Procedures
 - 4. Case File Organization, Preparation and Review Procedures.
 - 5. Procedures for Preparation, Approval, Review, Revision, and Distribution of SOPs.
 - 6. Process for Revision of Technical or Documentation Procedures
- D. Analytical Methodology
 - 1. Calibration Procedures and Frequency
 - 2. Sample Preparation/Extraction Procedures
 - 3. Sample Analysis Procedures
 - 4. Standards Preparation Procedures
 - 5. Decision Processes, Procedures, and Responsibility for Initiation of Corrective Action
- E. Data Generation
 - 1. Data Collection Procedures
 - 2. Data Reduction Procedures
 - 3. Data Validation Procedures
 - 4. Data Reporting and Authorization Procedures
- F. Quality Control
 - 1. Solvent, Reagent and Adsorbent Check Analysis
 - 2. Reference Material Analysis
 - 3. Internal Quality Control Checks
 - 4. Corrective Action and Determination of QC Limit Procedures
 - 5. Responsibility Designation

G. Quality Assurance

- 1. Data Quality Assurance
- 2. Systems/Internal Audits
- 3. Performance/External Audits
- 4. Corrective Action Procedures
- 5. Quality Assurance Reporting Procedures
- 6. Responsibility Designation

SECTION III

STANDARD OPERATING PROCEDURES

In order to obtain reliable results, adherence to prescribed analytical methodology is imperative. In any operation that is performed on a repetitive basis, reproducibility is best accomplished through the use of Standard Operating Procedures (SOPs). As defined by the EPA, an SOP is a written document which provides directions for the step-by-step execution of an operation, analysis, or action which is commonly accepted as the method for performing certain routine or repetitive tasks⁽²⁾.

SOPs prepared by the Contractor must be functional: i.e., clear, comprehensive, up-to-date, and sufficiently detailed to permit duplication of results by qualified analysts. All SOPs, as presented to the Agency, must reflect activities as they are currently performed in the laboratory. In addition, all SOPs must:

- o Be consistent with current EPA regulations, guidelines, and the CLP contract's requirements (3,4,5,6,7).
- o Be consistent with instruments manufacturers's specific instruction manuals.
- Be available to the EPA during an On-Site Laboratory Evaluation. A complete set of SOP's shall be bound together and available for inspection at such evaluations. During On-Site evaluations, laboratory personnel may be asked to demonstrate the application of the SOPs.
- o Provide for the development of documentation that is sufficiently complete to record the performance of all tasks required by the protocol.
- o Demonstrate the validity of data reported by the Contractor and explain the cause of missing or inconsistent results.
- o Describe the corrective measures and feedback mechanism utilized when analytical results do not meet protocol requirements.
- Be reviewed regularly and updated as necessary when contract, facility, or Contractor procedural modifications are made.
- o Be archived for future reference in usability or evidentiary situations.
- o Be available at specific work stations as appropriate
- o Be subject to a document control procedure which precludes the use of outdated or inappropriate SOPs.

SOP FORMAT:

The format for SOPs may vary depending upon the kind of activity for which they are prepared, however, at a minimum, the following sections must be included:

- o Title Page
- o Scope and Application
- o Definitions
- o Procedures
- o QC Limits
- o Corrective Action Procedures, Including Procedures for Secondary Review of Information Being Generated
- o Documentation Description and Example Forms
- o Miscellaneous Notes and Precautions
- o References

SOPs Required:

The following SOPs are required by the Agency:

1. Evidentiary SOP

Evidentiary SOPs required are discussed in Exhibit F, "Specification for Written Standard Operating Procedures"

- 2. Sample Receipt and logging
- 3. Sample storage
- 4. Preventing sample contamination
- 5. Security for laboratory and samples
- 6. Standards purity/preparation
- 7. Maintaining instrument records and logbooks
- 8. Sample analysis and data control systems
- 9. Glassware cleaning
- 10. Technical and managerial review of laboratory operation and data package preparation
- 11. Internal review of contractually-required quality assurance and quality control data for each individual data package
- 12. Sample analysis, data handling and reporting
- 13. Chain-of-custody procedures and document control including SDG file preparation
- 14. Laboratory data validation/laboratory self inspection system
 - a. Data flow and chain-of-command for data review
 - b. Procedures for measuring precision and accuracy
 - c. Evaluation parameters for identifying systematic errors⁽⁹⁾
 - d. Procedures to assure that hardcopy and diskette deliverables are complete and compliant with the requirements in SOW Exhibits B and H.

- e. Procedures to assure that hardcopy deliverables are in agreement with their comparable diskette deliverables.
- f. Demonstration of internal QA inspection procedure (demonstrated by supervisory sign-off on personal notebooks, internal PE samples, etc.).
- g. Frequency and type of internal audits (eg., random, quarterly, spot checks, perceived trouble areas).
- h. Demonstration of problem identification-corrective actions and resumption of analytical processing. Sequence resulting from internal audit (i.e., QA feedback).
- i. Documentation of audit reports, (internal and external), response, corrective action, etc.

15. Data Management and Handling

- a. Procedures for controlling and estimating data entry errors.
- b. Procedures for reviewing changes to data and deliverables and ensuring traceability of updates.
- c. Lifecycle management procedures for testing, modifying and implementing changes to existing computing systems including hardware, software, and documentation or installing new systems.
- d. Database security, backup and archival procedures including recovery from system failures.
- e. System maintenance procedures and response time.
- f. Individuals(s) responsible for system operation, maintenance, data integrity and security.
- g. Specifications for staff training procedures.

SOPs Delivery Requirements:

Within forty-five (45) days of contract receipt, a complete set of SOPs, relevant to this contract shall be sent to the Technical Project Officer, SMO and EMSL/LV. Also, during the term of performance of the contract, copies of SOPs which have been amended or new SOPs which have been written shall be sent to the Technical Project Officer, EMSL/LV (quality assurance SOPs) and NEIC (evidentiary SOPs).

SECTION IV VOA VOLATILE QA/QC REQUIREMENTS

INTRODUCTION

Sections II and III of this exhibit outline the requirements for the quality assurance program that each laboratory must establish under this contract. The purpose of Section III is to outline the minimum quality control (QC) operations necessary to satisfy the analytical requirements associated with the determination of the volatile organic target compounds listed in Exhibit C using the procedures in Exhibit D VOA for water and soil/sediment samples. This section is <u>not</u> intended as a comprehensive quality control document, but rather as a guide to the specific QC operations that must be considered for volatile analyses. At a minimum, the laboratory is expected to address these operations in preparing the quality assurance plan and QA/QC Standard Operating Procedures discussed in Section II.

These operations include the following:

- o GC/MS Mass Calibration and Ion Abundance Patterns
- o GC/MS Initial and Continuing Calibration
- o Stability of Internal Standard Responses and Retention Times
- Method Blank Analysis
- o System Monitoring Compound Recoveries
- o Matrix Spike and Matrix Spike Duplicate Analyses
- o Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

Not discussed in this section are the requirements for quality assurance of the data reporting aspects of volatile analyses which are described in general terms in Section II and III of this exhibit.

1. GC/MS Mass Calibration and Ion Abundance Patterns

Prior to initiating any data collection activities involving samples, blanks, or standards, it is necessary to establish that a given GC/MS system meets the instrument performance criteria specified in Exhibit D VOA, Section IV, paragraph 6. The purpose of this instrument performance check is to assure correct mass calibration, mass resolution, and mass transmission. This is accomplished through the analysis of Bromofluorobenzene (BFB).

- 1.1 The required frequency of BFB analysis (once every 12 hours on each GC/MS system) is described in detail in Ex. D VOA, Section IV, paragraph 6.4.
- 1.2 The key ions produced during the analysis of BFB and their respective ion abundance criteria are given in Table 1, Ex. D VOA, Section IV, paragraph 6.4.4.

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- 1.3 The documentation includes Form V VOA, and a mass listing and bar graph spectrum of each BFB analysis.
- 2. GC/MS Initial Calibration for Target Compounds and System Monitoring Compounds

Prior to the analysis of samples and required blanks and after instrument performance criteria have been met, the GC/MS system must be initially calibrated at a minimum of five concentrations to determine the linearity of response utilizing target compound and system monitoring compound standards.

- 2.1 The concentrations of the initial calibration standards for volatile target compounds and system monitoring compounds are 10, 20, 50, 100, and 200 ug/L, as déscribed in Ex. D VOA, Section IV, paragraph 5.5.
- 2.2 The standards are to be analyzed according to the procedures given in Ex. D VOA, Section IV, paragraph 7, and at the frequency given in that paragraph.
- 2.3 The relative response factors (RRFs) are determined according to the procedures in Ex. D VOA, Section IV, paragraph 7.4, using the assignment of internal standard to target compounds and system monitoring compounds given in Ex. D VOA, Section IV, paragraph 7.4, and Tables 3 and 4.
- 2.4 The calibration of the GC/MS is evaluated on the basis of the magnitude and stability of the relative response factors of each target compound and system monitoring compound. The minimum RRF of each compound in the initial calibration and the percent relative standard deviation (%RSD) across all five points must meet the criteria given Ex. D VOA, Section IV, paragraphs 7.4.5 and 7.4.6, and Table 2. Allowance is made for any two volatile compounds that fail to meet these criteria. The minimum RRFs of those two compounds must be greater than or equal to 0.010, and the %RSD must be less than or equal to 40.0% for the initial calibration to be acceptable.
- 2.5 The documentation includes Form VI VOA, a GC/MS data system printout for the analysis of each volatile calibration standard, and the mass spectrum of each target compound and system monitoring compound.
- 3. GC/MS Continuing Calibration for Target Compounds and System Monitoring Compounds.

Once the GC/MS system has been calibrated, the calibration must be verified each twelve (12) hour time period for each GC/MS system.

3.1 The concentration of the continuing calibration standard for volatile target compounds and system monitoring compounds is 50 ug/L, as described in Ex. D VOA, Section IV, paragraph 5.5.3.

- 3.2 The standard is to be analyzed according to the procedures given in Ex. D VOA, Section IV, paragraph 7, and at the frequency given in that paragraph.
- 3.3 The continuing calibration of the GC/MS system is evaluated on the basis of the magnitude of the relative response factors and the percent difference between the <u>average</u> RRF of each compound from the initial calibration and the RRF of that compound in the continuing calibration standard. The minimum RRF of each compound in the continuing calibration and the percent difference must meet the criteria given Ex. D VOA, Section IV, paragraphs 7.4.5 and 7.4.7, and Table 2. Allowance is made for any two volatile compounds that fail to meet these criteria. The minimum RRFs of those two compounds must be greater than or equal to 0.010, and the percent difference must be less than or equal to 40.0% for the continuing calibration to be acceptable.
- 3.4 The documentation includes Form VII VOA, a GC/MS data system printout for the analysis of the volatile calibration standard, and the mass spectrum of each target compound and system monitoring compound.
- 4. Internal Standard Responses and Retention Times

The response of each of the internal standards in all calibration standards, samples, and blanks is crucial to the provision of reliable analytical results, because the quantitative determination of volatile compounds by these procedures is based on the use of internal standards added immediately prior to analysis.

- 4.1 The specific compounds used as internal standards are given in Ex. D VOA, Section IV, paragraph 5.4.3. The concentration of each internal standard in the aliquot of the sample analyzed by GC/MS must be 50 ug/L at the time of purging.
- 4.2 The retention time and the extracted ion current profile (EICP) of each internal standard must be monitored for all analyses.
- 4.3 The area response of each internal standard from the EICP and the retention time of the internal standard are evaluated for stability, according to the procedures in Ex. D VOA, Section IV, paragraph 10.2. The area of the internal standard in a sample must not vary by more than a factor of 2 (i.e. -50% to +100%) from the area of the same internal standard in the associated continuing calibration standard. Likewise, the retention time of an internal standard must be within \pm 0.50 minutes (30 seconds) of its retention time in the continuing calibration standard (see Ex.D VOA, Section IV, paragraph 10.2).
- 4.4 Requirements for reanalysis of samples when internal standards do not meet specifications are given in Ex. D VOA, Section IV, paragraph 10.2.

- 4.5 The documentation includes Form VIII VOA, and the GC/MS data system printout for the analysis of each sample, blank, matrix spike, matrix spike duplicate, and standard.
- 5. Method Blank Analysis

A method blank is a volume of a clean reference matrix (deionized distilled water for water samples, or a purified solid matrix for soil/sediment samples) that is carried through the entire analytical procedure. The volume or weight of the reference matrix must be approximately equal to the volume or weight of samples associated with the blank. The purpose of a method blank is to determine the levels of contamination associated with the processing and analysis of samples.

- 5.1 For volatile analysis, a method blank must be analyzed once every 12 hours on each GC/MS system, as described in detail in Ex. D VOA, Section IV, paragraphs 8.1.16, 8.2.1.9, and 8.2.2.10.
- 5.2 For the purposes of this protocol, an acceptable method blank must meet the criteria in paragraphs 5.2.1. and 5.2.2 below.
 - 5.2.1 A method blank for volatile analysis must contain less than or equal to five times (5x) the Contract Required Quantitation Limit (CRQL, see Exhibit C) of Methylene chloride, Acetone, and 2-Butanone.
 - 5.2.2 For all other target compounds not listed above, the method blank must contain less than or equal to the Contract Required Quantitation Limit (CRQL, see Exhibit C) of any single target compound.
- 5.3 If a method blank exceeds the limits for contamination above, the Contractor must consider the analytical system out of control. The source of the contamination must be investigated and appropriate corrective actions taken and documented before further sample analysis proceeds. The requirements for reanalysis of associated samples are given in Ex. D VOA, Section IV, paragraph 10.10.
- 5.4 The documentation includes Form I VOA for the blank analysis, Form IV VOA, associating the samples and the blank, and a GC/MS data system printout for the analysis of the method blank.
- 6. System Monitoring Compound Recoveries

The recoveries of the three system monitoring compounds are calculated from the analysis of each sample, blank, matrix spike and matrix spike duplicate. The purpose of the system monitoring compounds is to evaluate the performance of the entire purge and trap-gas chromatograph-mass spectrometer system. Poor purging efficiency, leaks, and cold spots in transfer lines are only a few of the potential causes of poor recovery of these compounds.

- 6.1 The system monitoring compounds are added to each sample, blank, matrix spike, and matrix spike duplicate prior to purging or extraction (medium soils only), at the concentration described in Ex. D VOA, Section IV, paragraph 5.4.4.
- 6.2 The recoveries of the system monitoring compounds are calculated according to the procedures in Ex. D VOA, Section IV, paragraph 10.8.1.
- 6.3 The recoveries must be within the quality control limits given in Ex. D VOA,. Section IV, Table 5. If the recovery of any one system monitoring compound is outside these limits, the Contractor must follow the steps outlined in Ex. D VOA, Section IV, paragraphs 10.8.2 to 10.8.6.
- 6.4 The documentation includes Form II VOA, and a GC/MS data system printout for the analysis of each sample, blank, matrix spike, and matrix spike duplicate.
- 7. Matrix Spike and Matrix Spike Duplicate Analysis

In order to evaluate the effects of the sample matrix on the methods used for volatile analyses, the Agency has prescribed a mixture of volatile target compounds to be spiked into two aliquots of a sample, and analyzed in duplicate.

- 7.1 The frequency of matrix spike and matrix spike duplicate (MS/MSD) analysis is described in Ex. D VOA, Section IV, paragraph 10.9.
- 7.2 The recoveries of the matrix spike compounds are calculated according to the procedures in Ex. D VOA, Section IV, paragraph 10.9.1. The relative percent difference between the results for each spiked analyte of the matrix spike and the matrix spike duplicate are calculated according to the procedures in Ex. D VOA, Section IV, paragraph 10.9.2.
- 7.3 The quality control limits for recovery and relative percent difference are given in Ex. D VOA, Section IV, Table 6. These limits are only advisory at this time, and no further action is required when the limits are exceeded.
- 7.4 The documentation includes Form I VOA for both the MS and MSD analyses, Form III VOA, and a GC/MS printout for each analysis.
- 8. Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

If the on-column concentration of any sample exceeds the initial calibration range, that sample must be diluted and reanalyzed, as described in Ex. D VOA, Section IV, paragraph 10.7. Guidance in performing dilutions and <u>exceptions</u> are given in that paragraph, and reiterated here.

- 8.1 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
- 8.2 The dilution factor chosen should keep the response of the largest analyte peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.
- 8.3 Do <u>not</u> submit data for more than two analyses, i.e., the original sample and <u>one</u> dilution, or, if the volatile screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
- 8.4 Do <u>not</u> dilute MS/MSD samples solely to get non-spiked analytes within calibration range. The spiking level of each compound in the volatile matrix spiking solution should <u>not</u> require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.
- 8.5. For total Xylenes, where three isomers are quantified as two peaks, the calibration of <u>each peak</u>, should be considered separately, i.e., a diluted analysis is <u>not</u> required for total Xylenes unless the concentration of either peak separately exceeds 200 ug/L.

SECTION IV SV SEMIVOLATILE QA/QC REQUIREMENTS

INTRODUCTION

Sections II and III of this exhibit outline the requirements for the quality assurance program that each laboratory must establish under this contract. The purpose of Section III is to outline the minimum quality control (QC) operations necessary to satisfy the analytical requirements associated with the determination of the semivolatile organic target compounds listed in Exhibit C using the procedures in Exhibit D SV for water and soil/sediment samples. This section is not intended as a comprehensive quality control document, but rather as a guide to the specific QC operations that must be considered for semivolatile analyses. At a minimum, the laboratory is expected to address these operations in preparing the quality assurance plan and QA/QC Standard Operating Procedures discussed in Section II.

These operations include the following:

- o GC/MS Mass Calibration and Ion Abundance Patterns
- o GC/MS Initial and Continuing Calibration
- o Stability of Internal Standard Responses and Retention Times
- o Method Blank Analysis
- o Surrogate Recoveries
- Matrix Spike and Matrix Spike Duplicate Analyses
- o Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

Not discussed in this section are the requirements for quality assurance of the data reporting aspects of semivolatile analyses which are described in general terms in Section II and III of this exhibit.

1. GC/MS Mass Calibration and Ion Abundance Patterns

Prior to initiating any data collection activities involving samples, blanks, or standards, it is necessary to establish that a given GC/MS system meets the instrument performance criteria specified in Exhibit D SV, Section IV, paragraph 4. The purpose of this instrument performance check is to assure correct mass calibration, mass resolution, and mass transmission. This is accomplished through the analysis of Decafluorotriphenyl phosphine (DFTPP).

- 1.1 The required frequency of DFTPP analysis (once every 12 hours on each GC/MS system) is described in detail in Ex. D SV, Section IV, paragraph 4.3.6.
- 1.2 The key ions produced during the analysis of DFTPP and their respective ion abundance criteria are given in Table 1, Ex. D SV, Section IV, paragraph 4.3.3.

- 1.3 The documentation includes Form V SV, and a mass listing and bar graph spectrum of each DFTPP analysis.
- 2. GC/MS Initial Calibration for Target Compounds and Surrogates.

Prior to the analysis of samples and required blanks, and after instrument performance criteria have been met, the GC/MS system must be initially calibrated at a minimum of five concentrations to determine the linearity of response utilizing target compound and surrogate standards.

- 2.1 The levels of the initial calibration standards for semivolatile target compounds and surrogates are 20, 50, 80, 100, and 160 ng, in a 2 uL injection volume, as described in Ex. D SV, Section IV, paragraph 3.2.
- 2.2 The standards are to be analyzed according to the procedures given in Ex. D SV, Section IV, paragraph 5, and at the frequency given in that paragraph.
- 2.3 The relative response factors (RRFs) are determined according to the procedures in Ex. D SV, Section IV, paragraph 5.4, using the assignment of internal standard to target compounds and surrogates given in Ex. D SV, Section IV, paragraph 5.4, and Tables 3 and 4.
- 2.4 The calibration of the GC/MS is evaluated on the basis of the magnitude and stability of the relative response factors of each target compound and surrogate. The minimum RRF of each compound in the initial calibration and the percent relative standard deviation (%RSD) across all five points must meet the criteria given Ex. D SV, Section IV, paragraph 5.6, and Table 5. Allowance is made for any four semivolatile compounds that fail to meet these criteria. The minimum RRFs of those four compounds must be greater than or equal to 0.010, and the %RSD must be less than or equal to 40.0% for the initial calibration to be acceptable.
- 2.5 The documentation includes Form VI SV, a GC/MS data system printout for the analysis of each semivolatile calibration standard, and the mass spectrum of each target compound and surrogate.
- 3. GC/MS Continuing Calibration for Target Compounds and Surrogates.

Once the GC/MS system has been calibrated, the calibration must be verified each twelve (12) hour time period for each GC/MS system.

3.1 The level of the continuing calibration standard for semivolatile target compounds and surrogates is 50 ng, in a 2 uL injection volume, as described in Ex. D SV, Section IV, paragraph 3.2.

- 3.2 The standard is to be analyzed according to the procedures given in Ex. D SV, Section IV, paragraph 5, and at the frequency given in that paragraph.
- 3.3 The continuing calibration of the GC/MS system is evaluated on the basis of the magnitude of the relative response factors and the percent difference between the <u>average</u> RRF of each compound from the initial calibration and the RRF of that compound in the continuing calibration standard. The minimum RRF of each compound in the continuing calibration and the percent difference must meet the criteria given Ex. D SV, Section IV, paragraphs 5.6, 5.7, and Table 5. Allowance is made for any four semivolatile compounds that fail to meet these criteria. The minimum RRFs of those four compounds must be greater than or equal to 0.010, and the %RSD must be less than or equal to 40.0% for the continuing calibration to be acceptable.
- 3.4 The documentation includes Form VII SV, a GC/MS data system printout for the analysis of the semivolatile calibration standard, and the mass spectrum of each target compound and surrogate.
- 4. Internal Standard Responses and Retention Times

The response of each of the internal standards in all calibration standards, samples, and blanks is crucial to the provision of reliable analytical results because the quantitative determination of semivolatile compounds by these procedures is based on the use of internal standards added immediately prior to analysis.

- 4.1 The specific compounds used as internal standards are given in Ex. D SV, Section IV, paragraph 3.1. The amount of each internal standard in the injection volume (2 uL) of the sample extract analyzed by GC/MS must be 40 ng (20 ng/uL).
- 4.2 The retention time and the extracted ion current profile (EICP) of each internal standard must be monitored for all analyses.
- 4.3 The area response of each internal standard from the EICP and the retention time of the internal standard are evaluated for stability, according to the procedures in Ex. D SV, Section IV, paragraph 8.1. The area of the internal standard in a sample must not vary by more than a factor of 2 (i.e. -50% to +100%) from the area of the same internal standard in the associated continuing calibration standard. Likewise, the retention time of an internal standard must be within \pm 0.50 minutes (30 seconds) of its retention time in the continuing calibration standard (see Ex.D SV, Section IV, paragraph 8.1).
- 4.4 Requirements for reanalysis of samples when internal standards do not meet specifications are given in Ex. D SV, Section IV, paragraph 8.1.

- 4.5 The documentation includes Form VIII SV, and the GC/MS data system printout for the analysis of each sample, blank, matrix spike, matrix spike duplicate, and standard.
- 5. Method Blank Analysis

A method blank is a volume of a clean reference matrix (deionized distilled water for water samples, or a purified sodium sulfate for soil/sediment samples) that is carried through the entire analytical procedure. The volume or weight of the reference matrix must be approximately equal to the volume or weight of samples associated with the blank. The purpose of a method blank is to determine the levels of contamination associated with the processing and analysis of samples.

- 5.1 For semivolatile analysis, one method blank must be extracted with each group of samples of a similar matrix and concentration level (soils only), as described in Ex. D SV, Section IV, paragraph 8.7.
- 5.2 For the purposes of this protocol, an acceptable method blank must meet the criteria in paragraphs 5.2.1. and 5.2.2 below.
 - 5.2.1 A method blank for semivolatile analysis must contain less than or equal to five times (5x) the Contract Required Quantitation Limit (CRQL, see Ex. C) of the phthalate esters listed in Ex. C.
 - 5.2.2 For all other target compounds not listed above, the method blank must contain less than or equal to the Contract Required Quantitation Limit (CRQL, see Exhibit C) of any single target compound.
- 5.3 If a method blank exceeds the limits for contamination above, the Contractor must consider the analytical system out of control. The source of the contamination must be investigated and appropriate corrective actions taken and documented before further sample analysis proceeds. The requirements for reextraction and reanalysis of associated samples are given in Ex. D SV, Section IV, paragraph 8.7.
- 5.4 The documentation includes Form I SV for the blank analysis, Form IV SV, associating the samples and the blank, and a GC/MS data system printout for the analysis of the method blank.
- 6. Surrogate Recoveries

The recoveries of the eight surrogates are calculated from the analysis of each sample, blank, matrix spike and matrix spike duplicate. The purpose of the surrogates is to evaluate the preparation and analysis of samples.

- 6.1 The surrogates are added to each sample, blank, matrix spike, and matrix spike duplicate prior to extraction, at the concentrations described in Ex. D SV, Section II, Part B paragraph 4.6 and Part C paragraph 1.5.3 and 2.4.5.
- 6.2 The recoveries of the surrogates are calculated according to the procedures in Ex. D SV, Section IV, paragraph 8.5.1.
- 6.3 The recoveries must be within the quality control limits given in Ex. D SV, Section IV, Table 6. If the recovery of any surrogate is outside these limits, the Contractor must follow the steps outlined in E. D SV, Section IV, paragraphs 8.5.2 to 8.5.6 to determine whether or not reextraction and/or reanalysis is required.
- 6.4 The documentation includes Form II SV, and a GC/MS data system printout for the analysis of each sample, blank, matrix spike, and matrix spike duplicate.
- 7. Matrix Spike and Matrix Spike Duplicate Analysis

In order to evaluate the effects of the sample matrix on the methods used for semivolatile analyses, the Agency has prescribed a mixture of semivolatile target compounds to be spiked into two aliquots of a sample, and analyzed in duplicate.

- 7.1 The frequency of matrix spike and matrix spike duplicate (MS/MSD) analysis is described in Ex. D SV, Section IV, paragraph 8.6.
- 7.2 The recoveries of the matrix spike compounds are calculated according to the procedures in Ex. D SV, Section IV, paragraph 8.6.1. The relative percent difference between the results for each spiked analyte of the matrix spike and the matrix spike duplicate is calculated according to the procedures in Ex. D SV, Section IV, paragraph 8.6.2.
- 7.3 The quality control limits for recovery and relative percent difference are given in Ex. D SV, Section IV, Table 7. These limits are only advisory at this time, and no further action is required when the limits are exceeded.
- 7.4 The documentation includes Form I SV for both the MS and MSD analyses, Form III SV, and a GC/MS printout for each analysis.
- 8. Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

If the on-column concentration of any sample exceeds the initial calibration range, that sample must be diluted and reanalyzed, as described in Ex. D SV, Section IV, paragraph 8.4. Guidance in performing dilutions and <u>exceptions</u> are given in that paragraph, and reiterated here.

- 8.1 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
- 8.2 The dilution factor chosen should keep the response of the largest analyte peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.
- 8.3 Do <u>not</u> submit data for more than two analyses, i.e., the original sample and <u>one</u> dilution, or, if the semivolatile screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
- 8.4 Do not dilute MS/MSD samples solely to get non-spiked analytes within calibration range. The spiking level of each compound in the semivolatile matrix spiking solution should not require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.

SECTION IV PEST PESTICIDE/AROCLOR QA/QC REQUIREMENTS

INTRODUCTION

Sections II and III of this exhibit outline the requirements for the quality assurance program that each laboratory must establish under this contract. The purpose of Section III is to outline the minimum quality control (QC) operations necessary to satisfy the analytical requirements associated with the determination of the pesticide/Aroclor target compounds listed in Exhibit C using the procedures in Exhibit D PEST for water and soil/sediment samples. This section is <u>not</u> intended as a comprehensive quality control document, but rather as a guide to the specific QC operations that must be considered for pesticide/Aroclor analyses. At a minimum, the laboratory is expected to address these operations in preparing the quality assurance plan and QA/QC Standard Operating Procedures discussed in Section II.

These operations include the following:

- o GC Column Resolution
- GC/EC Initial and Continuing Calibration
- * o Determination of Retention Times and Retention Time Windows
 - Analytical Sequence
 - Blank Analyses
 - o Surrogate Recoveries
 - Matrix Spike and Matrix Spike Duplicate Analyses
 - Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

Not discussed in this section are the requirements for quality assurance of the data reporting aspects of pesticide/Aroclor analyses which are described in general terms in Section II and III of this exhibit.

1. GC Column Resolution

Prior to initiating any data collection activities involving samples, blanks, or standards, it is necessary to establish that a given GC column meets the analyte resolution criteria specified in Exhibit D PEST, Section III, paragraph 6.2.2. The purpose of this resolution check is to demonstrate that at the time of the initial calibration, the GC column is capable of chromatographically resolving the target compounds. This is accomplished through the analysis of the Resolution Check Mixture (see Ex. D, Section III, paragraph 3.1), which contains the nine target compounds that are most difficult to resolve.

1.1 The Resolution Check Mixture must be analyzed at the beginning of every initial calibration sequence, on each GC column and instrument used for analysis.

- 1.2 Additional resolution criteria apply to the target compounds in the standards used for initial calibration and calibration verification, as described in Ex. D Section III, paragraphs 6.2.4, 6.2.10, and 7.8.
- 1.3 The documentation includes Form VI PEST-4, chromatograms and data system printouts for the analysis of the Resolution Check Mixture on each GC column and instrument used for analysis.
- 2. GC/EC Initial Calibration for Target Compounds and Surrogates.

Prior to the analysis of samples and required blanks, the GC/EC system must be initially calibrated at a minimum of three concentrations to determine the linearity of response utilizing single component target compound and surrogate standards. Multicomponent target compounds are calibrated at a single point.

- 2.1 The concentrations of the low point initial calibration standards for single component pesticide target compounds and surrogates are described in Ex. D PEST, Section III, paragraph 3.3. The concentration of the mid point initial calibration standards is specified in Ex. D, Section III, paragraph 3.3 as 10 times the low point concentration. The concentration of the high point initial calibration standard must be at least 30 times the low point concentration, and may be as high as 100 times the low point concentration, as described in Ex. D PEST, Section III, paragraph 3.3.
- 2.2 The standards are to be analyzed according to the procedures given in Ex. D PEST, Section III, using the GC operating conditions in paragraphs 4 and 6, and at the frequency given in paragraph 6.1.
- 2.3 The calibration factors are determined according to the procedures in Ex. D PEST, Section III, paragraphs 9 and 10.
- 2.4 The initial calibration of the GC/EC is evaluated on the basis of the stability of the calibrations factors and retention times of each target compound and surrogate, described in Ex. D PEST, Section III, paragraphs 6.2.5 to 6.2.9.
- 2.5 The calibration is also evaluated on the basis of the extent of breakdown of two target compounds, Endrin and 4,4'-DDT, as described in Ex. D PEST, Section III, paragraph 6.2.3.
- 2.6 The documentation includes Form VI PEST, chromatograms and data system printouts of all standards for the pesticide/Aroclor calibration standards.

3. GC/EC Continuing Calibration for Target compounds and Surrogates.

Once the GC/EC system has been calibrated, the calibration must be verified each twelve (12) hour time period for each GC column and instrument used for analysis. The calibration is verified through the analysis of instrument blanks, Performance Evaluation Mixtures (PEM), and the mid point concentrations of Individual Standard Mixtures A and B.

- 3.1 The concentrations of the PEM and Individual Standard Mixtures used for continuing calibration are given in Ex. D PEST, Section III, paragraphs 3.2 and 3.3.
- 3.2 The instrument blank is described in Ex. D PEST, Section III, paragraph 15.3.
- 3.3 The instrument blank and the standards must be analyzed once every twelve hours according to the procedures in Ex. D PEST, Section III, paragraph 5, bracketing the sample analyses, as described in Ex. D PEST, Section III, paragraph 7.
- 3.4 The continuing calibration is evaluated on the basis of the stability of the retention times of the target compounds in the standards.
- 3.5 The continuing calibration is evaluated on the basis of the stability of the instrument response to the target compounds in the PEM, as judged by the reproducibility of the determinations of the concentrations of these compounds in the standard, as described in Ex. D PEST, Section III, paragraph 7.10.
- 3.6 The continuing calibration is evaluated on the basis of the extent of breakdown of two target compounds in the PEM, Endrin and 4,4'-DDT, as described in Ex. D PEST, Section III, paragraph 7.11.
- 3.7 The continuing calibration is evaluated on the basis of the levels of contamination that are found in the instrument blank, as described in Ex. D PEST, Section III paragraph 15.3.
- 3.8 The documentation includes Form VII PEST, Form VIII PEST, chromatograms and data system printouts for all standards and instrument blanks analyzed.
- 4. Determination of Retention Times and Retention Time Windows

The identification of single component pesticides by gas chromatographic methods is based primarily on retention time data. The identification of multicomponent analytes is based primarily on recognition of patterns of retention times displayed on a chromatogram. Therefore, the determination of retention times and retention time windows is crucial to the provision of valid data for these target compounds.

- 4.1 The identification of all target compounds analyzed by he procedures described in Ex. D PEST is based on the use of absolute retention time. The mean retention time of each target compound, or each peak in a multicomponent target compound, is determined from the initial calibration standards, according to the procedures outlined in Ex. D PEST. Section III, paragraph 8.
- 4.2 The retention time window of each target compound peak is determined as \pm 0.04 minute of the mean retention time for all analytes except Methoxychlor, where the retention time window is \pm 0.05 minutes, as described in Ex. D, Section III, paragraph 8.4.
- 4.3 The retention time of each analyte in each standard in the initial and continuing calibrations must be within a retention time window half as large as those described above, as detailed in Ex D PEST, Section III, paragraphs 6.2.9 and 8.2.
- 4.4 The retention time shifts of the surrogates are used to evaluate the stability of the gas chromatgraphic system during analysis of samples and standards. The retention time of the surrogates must be within the retention time windows determined during the initial calibration (Ex. D PEST, Section III, paragraph 8.4.
- 4.5 The documentation includes Form VI PEST, Form VII PEST, Form VIII PEST, chromatograms and data system printouts for all standards for the Pesticide/Aroclor initial and continuing calibrations, on each instrument and GC column used for analysis.
- 5. Analytical Sequence

The standards and samples analyzed according to the procedures in Ex. D PEST Section III must be analyzed in a sequence described in paragraphs ⁵ and 6. This sequence includes requirements that apply to the initial and continuing calibrations, as well as to the analysis of samples. The documentation includes Form VIII PEST.

6. Blank Analysis

Two types of blanks are required for analyses using the procedures in Ex. D PEST. They are method blanks and instrument blanks. A third type of blank, a sulfur clean up blank, may be required.

6.1 A method blank is a volume of a clean reference matrix (deionized distilled water for water samples, or a purified sodium sulfate for soil/sediment samples) that is carried through the entire analytical procedure. The volume or weight of the reference matrix must be approximately equal to the volume or weight of samples associated with the blank. The purpose of a method blank is to determine the levels of contamination associated with the processing and analysis of samples.

- 6.1.1 The frequency of method blank extraction is described in Ex. D PEST, Section III, paragraph 15.1.1.
- 6.1.2 The method blank must be analyzed on each GC column and instrument used for the analysis of associated samples.
- 6.1.3 For the purposes of this protocol, an acceptable method blank must meet the criteria in Ex. D PEST, Section, III, paragraph 15.1.2.
- 6.2 The instrument blank is a volume of clean solvent spiked with the surrogates and analyzed on each GC column and instrument used for sample analysis. The purpose of the instrument blank is to determine the levels of contamination associated with the instrumental analysis itself, particularly with regard to the carry-over of analytes from standards or highly contaminated samples into other analyses.
 - 6.2.1 The frequency of instrument blank analysis is part of the initial and continuing calibration requirements described in Ex. D, Section III, paragraphs 5,6, and 7.
 - 6.2.2 For the purposes of this protocol, an acceptable instrument blank must meet the criteria in Ex. D PEST, Section, III, paragraph 15.3.3.
- 6.3 The sulfur clean up blank is a volume of clean solvent spiked with the surrogates and carried through the sulfur clean up and analysis steps. The purpose of the sulfur clean up blank is to determine the levels of contamination associated with the separate sulfur clean up steps.
 - 6.3.1 The sulfur clean up blank is only required when all the samples associated with a particular method blank are not subjected to sulfur clean up, as described in Ex. D PEST, Section III, paragraph 15.2.2.
 - 6.3.2 The sulfur clean up blank must be analyzed on all GC column and instruments used for analysis of samples that received sulfur clean up.
 - 6.3.3 For the purposes of this protocol, an acceptable sulfur clean up blank must meet the criteria in Ex. D PEST, Section, III, paragraph 15.2.3.

- 6.4 If a method blank exceeds the limits for contamination above, the Contractor must consider the analytical system out of control. The source of the contamination must be investigated and appropriate corrective actions taken and documented before further sample analysis proceeds. The requirements for reextraction and reanalysis of associated samples are given in Ex. D PEST, Section III, paragraph 15.1.3.
- 6.5 If an instrument blank exceeds the limits for contamination above, the Contractor must consider the analytical system out of control. The source of the contamination must be investigated and appropriate corrective actions taken and documented before further sample analysis proceeds. The requirements for reanalysis of associated samples are given in Ex. D PEST, Section III, paragraph 15.3.4.
- 6.6 If a sulfur clean up blank exceeds the limits for contamination above, the Contractor must consider the analytical system out of control. The source of the contamination must be investigated and appropriate corrective actions taken and documented before further sample analysis proceeds. The requirements for reextraction and reanalysis of associated samples are given in Ex. D PEST, Section III, paragraph 15.2.4.
- 6.7 The documentation includes Form I PEST for the analysis of each type of blank; Form IV PEST, associating the samples and the method and sulfur clean up blank; Form VIII PEST, associating the samples and the instrument blanks; and a GC/EC data system printout for the analysis of each blank.

7. Surrogate Recoveries

The recoveries of the two surrogates are calculated from the analysis on each GC column of each sample, blank, matrix spike and matrix spike duplicate. The purpose of the surrogates is to evaluate the preparation and analysis of samples.

- 7.1 The surrogates are added to each sample, blank, matrix spike, and matrix spike duplicate prior to extraction, at the concentrations described in Ex. D PEST, Section II, paragraph 4.9.4.
- 7.2 The recoveries of the surrogates are calculated according to the procedures in Ex. D PEST, Section III, paragraph 13.6.
- 7.3 The quality control limits for surrogate recovery, given in Ex. D PEST,. Section III, paragraph 13.6, are 60-150 percent. These limits are only advisory, and no further action by the laboratory is required if the limits are exceeded, however, frequent failures to meet the limits for surrogate recovery warrant investigation by the laboratory, and any result in questions from the Agency.

- 7.4 The documentation includes Form II PEST, a chromatogram and a GC/EC data system printout for the analysis of each sample, blank, matrix spike, and matrix spike duplicate.
- 8. Matrix Spike and Matrix Spike Duplicate Analysis

In order to evaluate the effects of the sample matrix on the methods used for pesticide/Aroclor analyses, the Agency has prescribed a mixture of pesticide/Aroclor target compounds to be spiked into two aliquots of a sample, and analyzed in duplicate.

- 8.1 The frequency of matrix spike and matrix spike duplicate (MS/MSD) analysis is described in Ex. D PEST, Section III, paragraph 16.1.
- 8.2 The recoveries of the matrix spike compounds are calculated according to the procedures in Ex. D PEST, Section III, paragraph 16.3. The relative percent difference for each spiked analyte between the results of the matrix spike and the matrix spike duplicate are calculated according to the procedures in Ex. D PEST, Section III, paragraph 16.3.
- 8.3 The quality control limits for recovery and relative percent difference are given in Ex. D PEST, Section III, paragraph 16.4. These limits are only advisory at this time, and no further action is required when the limits are exceeded.
- 8.4 The documentation includes Form I PEST for both the MS and MSD analyses, Form III PEST, and chromatograms and a GC/EC data system printout for each analysis.
- 9. Dilution of Samples, Matrix Spikes, and Matrix Spike Duplicates

If the on-column concentration of any sample exceeds the initial calibration range, that sample must be diluted and reanalyzed, as described in Ex. D PEST, Section III, paragraph 13.4. Guidance in performing dilutions and <u>exceptions</u> are given in that paragraph, and reiterated here.

- 9.1 If the response is still above the high calibration point after the dilution of 1:100,000, the Contractor shall contact the SMO immediately.
- 9.2 Use the results of the original analysis to determine the approximate dilution factor required to get the largest analyte peak within the initial calibration range.
- 9.3 The dilution factor chosen should keep the response of the largest analyte peak for a <u>target compound</u> in the upper half of the initial calibration range of the instrument.

- 9.4 Do not submit data for more than two analyses, i.e., the original sample and <u>one</u> dilution, or, if the pesticide/Aroclor screening procedure was employed, from the most concentrated dilution analyzed and one further dilution.
- 9.5 Do <u>not</u> dilute MS/MSD samples solely to get non-spiked analytes within calibration range. The spiking level of each compound in the pesticide/Aroclor matrix spiking solution should <u>not</u> require the dilution of the MS/MSD unless the sample from which these aliquots were taken contains high levels of the spiked analytes.

SECTION V

ANALYTICAL STANDARDS REQUIREMENTS

Overview

The U.S. Environmental Protection Agency will not supply analytical reference standards either for direct analytical measurements or for the purpose of traceability. All contract laboratories will be required to prepare from neat materials or purchase from private chemical supply houses those standards necessary to successfully and accurately perform the analyses required in this protocol.

A. Preparation of Chemical Standards from the Neat High Purity Bulk Material

A laboratory may prepare their chemical standards from neat materials. Commercial sources for neat chemical standards pertaining to compounds listed on the Compound Target List are given in the Appendix C of the "Quality Assurance Materials Bank: Analytical Reference Standards" Seventh Edition, January 1988. Laboratories should obtain the highest purity possible when purchasing neat chemical standards; standards purchased at less than 97% purity must be documented as to why a higher purity could not be obtained.

- 1. Neat chemical standards must be kept refrigerated when not being used in the preparation of standard solutions. Proper storage of neat chemicals is essential in order to safeguard them from decomposition.
- 2. The purity of a compound can sometimes be misrepresented by a chemical supply house. Since knowledge of purity is needed to calculate the concentration of solute in a solution standard, it is the contract laboratory's responsibility to have analytical documentation ascertaining that the purity of each compound is correctly stated. Purity confirmation, when performed, should use either differential scanning calorimetry, gas chromatography with flame ionization detection, high performance liquid chromatography, infrared spectrometry, or other appropriate techniques. Use of two or more independant methods is recommended. The correction factor for impurity when weighing neat materials in the preparation of solution standards is:

Equation 1

weight of impure compound - (percent purity/100)

where "weight of pure compound" is that required to prepare a specific volume of a solution standard of a specified concentration.

OLMO1.0

- 3. Mis-identification of compounds occasionally occurs and it is possible that a mislabeled compound may be received from a chemical supply house. It is the contract laboratory's responsibility to have analytical documentation ascertaining that all compounds used in the preparation of solution standards be correctly identified. Identification confirmation, when performed, should use, gas chromatographic/mass spectrometry analysis on at least two different analytical columns, or other appropriate techniques.
- 4. Calculate the weight of material to be weighed out for a specified volume taking into account the purity of the compound and the desired concentration. A second person must verify the accuracy of the calculations. Check balances for accuracy with a set of standard weights. All weighing should be performed on an analytical balance to the nearest 0.1 mg and verified by a second person. The solvent used to dissolve the solute should be compatible with the protocol in which the standard is to be used; the solute should be soluble, stable, and nonreactive with the solvent. In the case of a multicomponent solution, the components must not react with each other.
- 5. Transfer the solute to a volumetric flask and dilute to the specified solution volume with solvent after ensuring dissolution of the solute in the solvent. Sonication or warming may be performed to promote dissolution of the solute. This solution is to be called the primary standard and all subsequent dilutions must be traceable back to the primary standard.
- 6. Log notebooks are to be kept for all weighing and dilutions. All subsequent dilutions from the primary standard and the calculations for determining their concentrations are to be recorded and verified by a second person. All solution standards are to be refrigerated when not in use. All solution standards are to be clearly labeled as to the identity of the compound or compounds, concentration, date prepared, solvent, and initials of the preparer.
- B. Purchase of chemical standards already in solution

Solutions of analytical reference standards can be purchased by Contractors provided they meet the following criteria:

- 1. Laboratories must maintain the following documentation to verify the integrity of the standard solutions they purchase:
 - a. mass spectral identification confirmation of the neat material
 - b. purity confirmation of the neat material
 - c. chromatographic and quantitative documentation that the solution standard was QC checked according to the following section

2. The Contractor must purchase standards for which the quality is demonstrated statistically and analytically by a method of the supplier's choice. One way this can be demonstrated is to prepare and analyze three solutions; a high standard, a low standard, and a standard at the target concentration (see parts a and b below). The supplier must then demonstrate that the analytical results for the high standard and low standard are consistent with the difference in theoretical concentrations. This is done by the Student's t-test in part "d". If this is achieved, the supplier must then demonstrate that then demonstrate that the demonstrate that the demonstrate that the demonstrate in part "d". If this is achieved, the supplier must then demonstrate that the concentration of the target standard lies midway between the concentrations of the low and high standards. This is done by the Student's t-test in part e. Thus the standard is certified to be within 10 percent of the target concentration.

If the procedure above is used, the supplier must document that the following have been achieved:

- a. Two solutions of identical concentration must be prepared independently from neat materials. An aliquot of the first solution must be diluted to the intended concentration (the "target standard"). One aliquot is taken from the second solution and diluted to a concentration ten percent greater than the target standard. This is called the "high standard". One further aliquot is taken from the second solution and diluted to a concentration 10 percent less that the target standard. This is called the "low standard".
- b. Six replicate analyses of each standard (a total of 18 analyses) must be performed in the following sequence: low standard, target, high standard, low standard, target standard, high standard,
- c. The mean and variance of the six results for each solution must be calculated.

Equation 2 MEAN - $(Y_1 + Y_2 + Y_3 + Y_4 + Y_5 + Y_6)/6$

Equation 3

VARIANCE - $(Y_1^2 + Y_2^2 + Y_3^2 + Y_4^2 + Y_5^2 + Y_6^2 - (6*MEAN)^2)/5$

The values Y_1 , Y_2 , Y_3 , ..., represent the results of the six analyses of each standard. The means of the low, target, and high standards are designated M_1 , M_2 , and M_3 , respectively. The variances of the low, target, and high standards are designated V_1 , V_2 , and V_3 , respectively. Additionally, a pooled variance, V_p , is calculated.

Equation 4

 $v_p = (v_1/(0.81) + v_2 + v_3 / (1.21))/3$

If the square root of V_p is less than one percent of M_2 , then M_2^2 /10,000 is to be used as the value of V_p in all subsequent calculations.

d. The test statistic must be calculated:

Equation 5 TEST STATISTIC - $|(M_3 / 1.1) - (M_1 / 0.9)| / (V_p / 3)^{0.5}$

If the test statistic exceeds 2.13 then the supplier has failed to demonstrate a twenty percent difference between the high and low standards. In such a case, the standards are not acceptable.

e. The test statistic must be calculated:

Equation 6 TEST STATISTIC - $|M_2 - (M_1 / 1.8) - (M_3 / 2.2)| / (V_p / 4)^{0.5}$

If the test statistic exceeds 2.13, the supplier has failed to demonstrate that the target standard concentration is midway between the high and low standards. In such a case, the standards are not acceptable.

f. The 95 percent confidence intervals for the mean result of each standard must be calculated:

Equation 7 Interval for Low Standard - $M_1 \pm (2.13) (V_p / 6)^{0.5}$ Equation 8 Interval for Target Standard - $M_2 \pm (2.13) (V_p / 6)^{0.5}$ Equation 9 Interval for High Standard - $M_3 \pm (2.13) (V_p / 6)^{0.5}$

These intervals must not overlap. If overlap is observed, then the supplier has failed to demonstrate the ability to discriminate the 10 percent difference in concentrations. In such a case, the standards are not acceptable.

In any event, the laboratory is responsible for the quality of the standards employed for analyses under this contract.

C. Requesting Standards From the EPA Standards Repository

Solutions of analytical reference materials can be ordered from the U.S. EPA Chemical Standards Repository, depending on availability. The Contractor can place an order for standards only after demonstrating that these standards are not available from commercial vendors either in solution or as a neat material.

D. Documentation of the Verification and Preparation of Chemical Standards

It is the responsibility of each laboratory to maintain the necessary documentation to show that the chemical standards they have used in the performance of CLP analysis conform to the requirements previously listed. Weighing logbooks, calculations, chromatograms, mass spectra, etc, whether produced by the laboratory or purchased from chemical supply houses, must be maintained by the laboratory and may be subject to review during on-site inspection visits. Documentation of standards preparation may be required to be sent to EPA for verification of contract compliance. In those cases where the documentation is supportive of the analytical results of data packages sent to EPA, such documentation is to be kept on file by the laboratories for a period of one year.

SECTION VI

CONTRACT COMPLIANCE SCREENING

Contract Compliance Screening (CCS) is one aspect of the Government's contractual right of inspection of analytical data. CCS examines the Contractor's adherence to the contract requirements based on the sample data package delivered to the Agency.

CCS is performed by the Sample Management Office (SMO) under the direction of the EPA. To assure a uniform review, a set of standardized procedures have been developed to evaluate the sample data package submitted by a Contractor against the technical and completeness requirements of the contract.

CCS results are mailed to the Contractor and all other data recipients. The Contractor has a period of time to correct deficiencies. The Contractor must send all corrections to the Regional Client, EMSL/LV, and SMO.

CCS results are used in conjunction with other information to measure overall Contractor performance and to take appropriate actions to correct deficiencies in performance.

SECTION VII

REGIONAL DATA REVIEW

Contract laboratory data are generated to meet the specific needs of the Regions. In order to verify the useability of data for the intended purpose, each Region reviews data from the perspective of end-user, based upon functional aspects of data quality. General guidelines for data review have been developed jointly by the Region and the National Program Office. Each Region uses these guidelines as the basis for data evaluation. Individual Regions may augment the basic guideline review process with additional review based on Region-specific or site-specific concerns. Regional reviews, like the sites under investigation, vary based on the nature of the problems under investigation and the Regional response appropriate to the specific circumstances.

Regional data reviews, relating useability of the data to a specific site, are part of the collective assessment process. They complement the review done at the Sample Management Office, which is designed to identify contractual discrepancies and the review done at EMSL/LV which is designed to evaluate Contractor and method performance. These individual evaluations are integrated into a collective review that is necessary for program and laboratory administration and management and may be used to take appropriate action to correct deficiencies in the Contractor's performance.

SECTION VIII

LABORATORY EVALUATION SAMPLES

Although intralaboratory QC may demonstrate Contractor and method performance that can be tracked over time, an external performance evaluation program is an essential feature of a QA program. As a means of measuring Contractor and method performance, Contractors participate in interlaboratory comparison studies conducted by the EPA. Results from the analysis of these laboratory evaluation samples will be used by the EPA to verify the Contractor's continuing ability to produce acceptable analytical data. The results are also used to assess the precision and accuracy of the analytical methods for specific analytes.

Sample sets may be provided to participating Contractors as frequently as on an SDG-by-SDG basis as a recognizable QC sample of known composition; as a recognizable QC sample of unknown composition; or not recognizable as a QC material. The laboratory evaluation samples may be sent either by the Regional client or the National Program Office, and may be used for contract action.

Contractors are required to analyze the samples and return the data package and all raw data within the contract required turnaround time.

At a minimum, the results are evaluated for compound identification, quantification, and sample contamination. Confidence intervals for the quantification of target compounds are based on reported values using population statistics. EPA may adjust the scores on any given laboraotry evaluation sample to compensate for unanticipated difficulties with a particular sample. Normally, a fraction of the compounds spiked into the sample are not specifically listed in the contract. Contractors are required to use the NIST/EPA/MSDC mass spectral library to tentatively identify a maximum number of non-target compounds in each fraction that are present above a minimal response. Tentative identification of these compounds, based on contractually described spectral interpretation procedures, is evaluated and integrated into the evaluation process.

A Contractor's results on the laboratory evaluation samples will determine the Contractor's performance as follows:

1. Acceptable, No Response Required (Score greater than or equal to 90 percent):

Data meets most or all of the scoring criteria. No response is required.

2. Acceptable, Response Explaining Deficiency(ies) Required (Score greater than or equal to 75 percent but less than 90 percent):

Deficiencies exist in the Contractor's performance.

Within 14 days of receipt of notification from EPA, the Contractor shall describe the deficiency(ies) and the action(s) taken to correct the deficiency(ies) in a letter to the Administrative Project Officer, the Technical Project Officer and EMSL/LV.

3. Unacceptable Performance, Response Explaining Deficiency(ies) Required (Score less than 75 percent):

Deficiencies exist in the Contractor's performance to the extent that the National Program Office has determined that the Contractor has not demonstrated the capability to meet the contract requirements.

Within 14 days of receipt of notification from EPA, the Contractor shall describe the deficiency(ies) and the action(s) taken to correct the deficiency(ies) in a letter to the Administrative Project Officer, the Technical Project Officer and EMSL/LV.

The Contractor shall be notified by the Administrative Project Officer or Technical Project Officer concerning the remedy for their unacceptable performance. A Contractor may expect, but EPA is not limited to, the following actions: Reduction of the number of samples sent under the contract, suspension of sample shipment to the Contractor, a site visit, a full data audit, analysis of remedial PE samples, and/or a contract sanction, such as a Cure Notice.

Note: A Contractor's prompt response demonstrating that corrective actions have been taken to ensure the Contractor's capability to meet contract requirements will facilitate continuation of full sample delivery.

SECTION IX

GC/MS TAPE AUDITS

Periodically, EPA requests from Contractors the GC/MS magnetic tapes corresponding to a specific case in order to accomplish tape audits. Generally, tape submissions and audits are requested for the following reasons:

- o Program overview
- o Indication of data quality problems from EMSL/LV, SMO, or Regional data reviews
- o Support for on-site audits
- o Specific Regional requests

Depending upon the reason for an audit, the tapes from a recent case, a specific case, or a laboratory evaluation sample may be requested. Tape audits provide a mechanism to assess adherence to contractual requirements and to ensure the consistency of data reported on the hardcopy/floppy diskettes with that generated on the GC/MS tapes. This function provides external monitoring of Program QC requirements and checks adherence of the Contractor to internal QA procedures. In addition, tape audits enable EPA to evaluate the utility, precision, and accuracy of the analytical methods.

The GC/MS tape shall include raw data and quantitation reports for samples, blanks, laboraotry evaluation samples, initial calibrations, continuing calibration, BFB and DFTPP associated with the case requested. The specific requirements for submissions of GC/MS tapes are discussed in Exhibit B.

Upon request of the Administrative Project Officer or EMSL/LV, the required tapes and all necessary documentation shall be submitted to EPA within seven (7) days of notification.

SECTION X

ON-SITE LABORATORY EVALUATIONS

At a frequency dictated by a contract laboratory's performance, the Administrative Project Officer, Technical Project Officer or their authorized representative will conduct an on-site laboratory evaluation. On-site laboratory evaluations are carried out to monitor the Contractor's ability to meet selected terms and conditions specified in the contract. The evaluation process incorporates two separate categories: Quality Assurance Evaluation, and an Evidentiary Audit.

- A. Quality Assurance On-Site Evaluation
 - Quality assurance evaluators inspect the Contractor's facilities to verify the adequacy and maintenance of instrumentation, the continuity of personnel meeting experience or education requirements, and the acceptable performance of analytical and QC procedures. The Contractor should expect that items to be monitored will include, but not be limited to the following items.
 - Size and appearance of the facility
 - Quantity, age, availability, scheduled maintenance and performance of instrumentation
 - Availability, appropriateness, and utilization of SOPs
 - Staff qualifications, experience, and personnel training programs
 - o Reagents, standards, and sample storage facilities
 - Standard preparation logbooks and raw data
 - Bench sheets and analytical logbook maintenance and review
 - Review of the Contractor's sample analysis/data package inspection procedures

Prior to an on-site evaluation, various documentation pertaining to performance of the specific Contractor is integrated in a profile package for discussion during the evaluation. Items that may be included are previous on-site reports, laboratory evaluation sample scores, Regional review of data, Regional QA materials, GC/MS tape audit reports, results of CCS, and date trend reports.

B. Evidentiary Audit

Evidence auditors conduct an on-site laboratory evaluation to determine if laboratory policies and procedures are in place to satisfy evidence handling requirements as stated in Exhibit F. The evidence audit is comprised of the following three activities

1. Procedural Audit

The procedural audit consists of review and examination of actual standard operating procedures and accompanying documentation for the following laboratory operations: sample receiving, sample storage, sample identification, sample security, sample tracking (from receipt to completion of analysis) and analytical project file organization and assembly.

2. Written SOPs Audit

The written SOPs audit consists of review and examination of the written SOPs to determine if they are accurate and complete for the following laboratory operations: sample receiving, sample storage, sample identification, sample security, sample tracking (from receipt to completion of analysis) and analytical project file organization and assembly.

3. Analytical Project File Evidence Audit

The analytical project file evidence audit consists of review and examination of the analytical project file documentation. The auditors review the files to determine:

- The accuracy of the document inventory
- The completeness of the file
- The adequacy and accuracy of the document numbering system
- Traceability of sample activity
- Identification of activity recorded on the documents
- Error correction methods
- C. Discussion of the On-Site Team's Findings

The quality assurance and evidentiary auditors discuss their findings with the Administrative Project Officer/Technical Project Officer prior to debriefing the Contractor. During the debriefing, the auditors present their findings and recommendations for corrective actions necessary to the Contractor personnel.

- D. Corrective Action Reports For Follow-Through to Quality Assurance and Evidentiary Audit Reports
 - o Following an on-site evaluation, quality assurance and evidentiary audit reports which discuss deficiencies found during the on-site will be forwarded to the Contractor. The Contractor must discuss the corrective actions taken to resolve the deficiencies discussed during the on-site visit and discussed in the on-site reports in a letter to the Administrative Project Officer, Technical Project Officer, EMSL/LV (response to the quality assurance report) and NEIC (response to the evidentiary report) within 14 days of receipt of the finding or within the time agreed upon between the Administrative Project Officer/Technical Project Officer and the Contractor. If SOPs are

required to be written or SOPs are required to be amended, the Contractor must provide the SOPs to the Technical Project Officer, EMSL/LV (quality assurance/technical SOPs) and NEIC (evidentiary SOPs) within 30 days of receipt of the finding or within the time agreed upon between the Administrative Project Officer/Technical Project Officer and the Contractor.

 o If the Contractor fails to take appropriate corrective action to resolve the deficiencies discussed in the on-site reports, a Contractor may expect, but the Agency is not limited to, the following actions: reduction of the number of samples sent under the contract, suspension of sample shipment to the Contractor, a followup site visit, a full data audit, analysis of remedial PE samples and/or contract sanction, such as a Cure Notice.

SECTION II

QUALITY ASSURANCE AND DATA TREND ANALYSIS

Data submitted by laboratories are subject to review from several aspects: compliance with contract-required QC, useability, and full data package evaluation. Problems resulting from any of these reviews may determine the need for a GC/MS tape audit, an on-site laboratory evaluation and/or a remedial laboratory evaluation sample. In addition, QC prescribed in the methods provides information that is continually used by the Agency to assess sample data quality, Contractor data quality and Program data quality via data trend analysis. Trend analysis is accomplished by entering data into a computerized data base. Statistical reports that evaluate specific anomalies or disclose trends in many areas, including the following, are generated from this data base:

- o Surrogate Spike Recovery
- o Laboratory Evaluation Sample
- o Blanks
- o GC/MS Instrument Performance Checks (BFB and DFTPP)
- o Initial and Continuing Calibration Data
- o Other QC and Method Parameters

Program-wide statistical results are used to rank laboratories in order to observe the relative performance of each Contractor using a given protocol against its peers. The reports are also used to identify trends within laboratories. The results of many of these trends analyses are included in overall evaluation of a Contractor's performance, and are reviewed to determine if corrective action or an on-site laboratory evaluation is indicated in order to meet the QA/QC requirements of the contract.

Contractor performance over time is monitored using these trend analysis techniques to detect departures of Contractor output from required or desired levels of quality control, and to provide an early warning of Contractor QA/QC problems which may not be apparent from the results of an individual case.

As a further benefit to the Program, the data base provides the information needed to establish performance-based criteria in updated analytical protocols, where advisory criteria has been previously used. The vast empirical data set produced by contract laboratories is carefully analyzed, with the results augmenting theoretical and research-based performance criteria. The result is a continuously monitored set of quality control and performance criteria specifications of what is routinely achievable and expected of environmental chemistry laboratories in mass production analysis of environmental samples. This, in turn, assists the Agency in meeting its objectives of obtaining data of known and documented quality.

SECTION XII

DATA MANAGEMENT

Data management procedures are defined as procedures specifying the acquisition or entry, update, correction, deletion, storage and security of computer readable data and files. These procedures should be in written form and contain a clear definition for all databases and files used to generate or resubmit deliverables. Key areas of concern include: system organization (including personnel and security), documentation operations, traceability and quality control.

Data manually entered from hard-copy must be quality controlled and the error rates estimated. Systems should prevent entry of incorrect or out-of-range data and alert data entry personnel of errors. In addition, data entry error rates must be estimated and recorded on a monthly basis by reentering a statistical sample of the data entered and calculating discrepancy rates by data element.

The record of changes in the form of corrections and updates to data originally generated, submitted, and/or resubmitted must be documented to allow traceablilty of updates. Documentation must include the following for each change:

- Justification or rationale for the change.
- Initials of the person making the change or changes. Data changes must be implemented and reviewed by a person or group independent of the source generating the deliverable.
- Change documentation must be retained according to the schedule of the original deliverable.
- Resubmitted diskettes or other deliverables must be reinspected as a part of the laboratories' internal inspection process prior to resubmission. The entire deliverable, not just the changes, must be inspected.
- The Laboratory Manager must approve changes to originally submitted deliverables.
- Documentation of data changes may be requested by laboratory auditors.

Lifecycle management procedures must be applied to computer software systems developed by the laboratory to be used to generate and edit contract deliverables. Such systems must be thoroughly tested and documented prior to utilization.

- A software test and acceptance plan including test requirements, test results and acceptance criteria must be developed, followed, and available in written form.
- System changes must not be made directly to production systems generating deliverables. Changes must be made first to a development system and tested prior to implementation.

- o Each version of the production system will be given an identification number, date of installation, date of last operation and archived.
- o System and operations documentation must be developed and maintained for each system. Documentation must include a users manual and an operations and maintenance manual.

Individual(s) responsible for the following functions must be identified:

- o System operation and maintenance including documentation and training.
- o Database integrity, including data entry, data updating and quality control.
- o Data and system security, backup and archiving.

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EXHIBIT F

CHAIN-OF-CUSTODY, DOCUMENT CONTROL, AND STANDARD OPERATING PROCEDURES

1. SAMPLE CHAIN-OF-CUSTODY

A sample is physical evidence collected from a facility or from the environment. Controlling evidence is an essential part of the hazardous waste investigation effort. To accomplish this, Contractors are required to develop and implement the following sample identification, chain-of-custody, sample receiving, and sample tracking procedures.

1.1 Sample Identification

To assure traceability of the samples while in possession of the Contractor, the Contractor shall have a specified method for maintaining identification of samples throughout the laboratory.

Each sample and sample preparation container shall be labeled with the EPA number or a unique laboratory identifier. If a unique laboratory identifier is used, it shall be cross-referenced to the EPA number.

1.2 Chain-of-Custody Procedures

Because of the nature of the data being collected, the custody of EPA samples must be traceable from the time the samples are collected until they are introduced as evidence in legal proceedings. The Contractor shall have procedures ensuring that EPA sample custody is maintained and documented. A sample is under custody if:

- o It is in your possession, or
- o It is in your view after being in your possession, or
- o It was in your possession and you locked it up, or
- o It is in a designated secure area. (Secure areas shall be accessible only to authorized personnel.)
- 1.3 Sample Receiving Procedures
 - 1.3.1 The Contractor shall designate a sample custodian responsible for receiving all samples.
 - 1.3.2 The Contractor shall designate a representative to receive samples in the event that the sample custodian is not available.
 - 1.3.3 The condition of the shipping containers and sample bottles shall be inspected upon receipt by the sample custodian or his/her representative.
 - 1.3.4 The condition of the custody seals (intact/not intact) shall be inspected upon receipt by the sample custodian or his/her representative.
 - 1.3.5 The sample custodian or his/her representative shall check for the presence or absence of the following documents accompanying the sample shipment:

- o Airbills or airbill stickers
- o Custody seals
- o EPA custody records
- o EPA traffic reports or SAS packing lists
- o Sample tags
- 1.3.6 The sample custodian or his/her representative shall sign and date all forms (e.g., custody records, traffic reports or packing lists, and airbills) accompanying the samples at the time of sample receipt.
- 1.3.7 The Contractor shall contact the Sample Management Office (SMO) to resolve discrepancies and problems such as absent documents, conflicting information, broken custody seals, and unsatisfactory sample condition (e.g., leaking sample bottle).
- 1.3.8 The Contractor shall record the resolution of discrepancies and problems on Telephone Contact Logs.
- 1.3.9 The following information shall be recorded on Form DC-1 (See Exhibit B) by the sample custodian or his/her representative as samples are received and inspected:
 - o Condition of the shipping container
 - Presence or absence and condition of custody seals on shipping and/or sample containers
 - o Custody seal numbers, when present
 - o Condition of the sample bottles
 - o Presence or absence of airbills or airbill stickers
 - o Airbill or airbill sticker numbers
 - o Presence or absence of EPA custody records
 - Presence or absence of EPA traffic reports or SAS packing lists
 - o Presence or absence of sample tags
 - Sample tag identification numbers cross-referenced to the EPA sample numbers
 - Verification of agreement or non-agreement of information recorded on shipping documents and sample containers
 - o Problems or discrepancies

1.4 Sample Tracking Procedures

The Contractor shall maintain records documenting all phases of sample handling from receipt to final analysis.

2. DOCUMENT CONTROL PROCEDURES

The goal of the laboratory document control program is to assure that all documents for a specified Sample Delivery Group (SDG) will be accounted for when the project is completed. Accountable documents used by contract laboratories shall include, but not be limited to, logbooks, chain-of-custody records, sample work sheets, bench sheets, and other documents relating to the sample or sample analyses. The following document control procedures have been established to assure that all laboratory records are assembled and stored for delivery to EPA or are available upon request from EPA prior to the delivery schedule.

- 2.1 Preprinted Laboratory Forms and Logbooks
 - 2.1.1 All documents produced by the Contractor which are directly related to the preparation and analysis of EPA samples shall become the property of the EPA and shall be placed in the complete sample delivery group file (CSF). All observations and results recorded by the laboratory but not on preprinted laboratory forms shall be entered into permanent laboratory logbooks. When all data from a SDG is compiled, all original laboratory forms and copies of all SDG-related logbook entries shall be included in the documentation package.
 - 2.1.2 The Contractor shall identify the activity recorded on all laboratory documents which are directly related to the preparation and analysis of EPA samples.
 - 2.1.3 Pre-printed laboratory forms shall contain the name of the laboratory and be dated (month/day/year) and signed by the person responsible for performing the activity at the time an activity is performed.
 - 2.1.4 Logbook entries shall be dated (month/day/year) and signed by the person responsible for performing the activity at the time an activity is performed.
 - 2.1.5 Logbook entries shall be in chronological order. Entries in logbooks, with the exception of instrument run logs and extraction logs, shall include only one SDG per page.
 - 2.1.6 Pages in both bound and unbound logbooks shall be sequentially numbered.

2.1.7 Instrument run logs shall be maintained so as to enable a reconstruction of the run sequence of individual instruments.

Because the laboratory must provide copies of the instrument run logs to EPA, the laboratory may exercise the option of using only laboratory or EPA sample identification numbers in the logs for sample ID rather than government agency or commercial client names to preserve the confidentiality of commercial clients.

2.1.8 Corrections to supporting documents and raw data shall be made by drawing a single line through the error and entering the correct information. Corrections and additions to supporting documents and raw data shall be dated and initialed. No information shall be obliterated or rendered unreadable.

All notations shall be recorded in ink.

Unused portions of documents shall be "z'd" out.

2.2 Consistency of Documentation

The Contractor shall assign a document control officer responsible for the organization and assembly of the CSF.

All copies of laboratory documents shall be complete and legible.

Original documents which include information relating to more than one SDG shall be filed in the CSF of the lowest SDG number. The copy(s) shall be placed in the other CSF(s) and the Contractor shall record the following information on the copy(s) in red ink:

"COPY

ORIGINAL IS FILED IN CSF

The Contractor shall sign and date this addition to the copy(s).

Before releasing analytical results, the document control officer shall assemble and cross-check the information on samples tags, custody records, lab bench sheets, personal and instrument logs, and other relevant deliverables to ensure that data pertaining to each particular sample or sample delivery group is consistent throughout the CSF.

2.3 Document Numbering and Inventory Procedure

In order to provide document accountability of the completed analysis records, each item in the CSF shall be inventoried and assigned a serialized number as described in Exhibit B).

All documents relevant to each sample delivery group, including logbook pages, bench sheets, mass spectra, chromatograms, screening records, re-preparation records, re-analysis records, records of failed or attempted analysis, custody records, library research results, etc. shall be inventoried.

The Document Control Officer (DCO) shall be responsible for ensuring that all documents generated are placed in the CSF for inventory and are delivered to the appropriate EPA region or other receiver as designated by EPA. The DCO shall place the sample tags in plastic bags in the file.

2.4 Storage of EPA Files

The Contractor shall maintain EPA laboratory documents in a secure location.

2.5 Shipment of Deliverables

The Contractor shall document shipment of deliverables packages to the recipients. These shipments require custody seals on the containers placed such that they cannot be opened without damaging or breaking the seal. The Contractor shall document what was sent, to whom, the date, and the method (carrier) used.

A copy of the transmittal letter for the CSF shall be sent to the NEIC/CEAT and the SMO.

3. SPECIFICATIONS FOR WRITTEN STANDARD OPERATING PROCEDURES

The Contractor shall have written standard operating procedures (SOPs) for receipt of samples, maintenance of custody, sample identification, sample storage, sample tracking, and assembly of completed data. An SOP is defined as a written narrative stepwise description of laboratory operating procedures including examples of laboratory documents. The SOPs shall accurately describe the actual procedures used in the laboratory, and copies of the written SOPs shall be available to the appropriate laboratory personnel. These procedures are necessary to ensure that analytical data produced under this contract are acceptable for use in EPA enforcement case preparation and litigation. The Contractor's SOPs shall provide mechanisms and documentation to meet each of the following specifications and shall be used by EPA as the basis for laboratory evidence audits.

- 3.1 The Contractor shall have written SOPs describing the sample custodian's duties and responsibilities.
- 3.2 The Contractor shall have written SOPs for receiving and logging in of the samples. The procedures shall include but not be limited to documenting the following information:
 - 3.2.1 Presence or absence of EPA chain-of-custody forms
 - 3.2.2 Presence or absence of airbills or airbill stickers

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- 3.2.3 Presence or absence of traffic reports or SAS packing lists
- 3.2.4 Presence or absence of custody seals on shipping and/or sample containers and their condition
- 3.2.5 Custody seal numbers, when present
- 3.2.6 Airbill or airbill sticker numbers
- 3.2.7 Presence or absence of sample tags
- 3.2.8 Sample tag ID numbers
- 3.2.9 Condition of the shipping container
- 3.2.10 Condition of the sample bottles
- 3.2.11 Verification of agreement or non-agreement of information on receiving documents and sample containers
- 3.2.12 Resolution of problems or discrepancies with the SMO
- 3.2.13 An explanation of any terms used by the laboratory to describe sample condition upon receipt (e.g., good, fine, OK)
- 3.3 The Contractor shall have written SOPs for maintaining identification of EPA samples throughout the laboratory.

If the Contractor assigns unique laboratory identifiers, written SOPs shall include a description of the method used to assign the unique laboratory identifier and shall include a description of the document used to cross-reference the unique laboratory identifier to the EPA sample number.

If the Contractor uses prefixes or suffixes in addition to sample identification numbers, the written SOPs shall include their definitions.

- 3.4 The Contractor shall have written SOPs describing all storage areas for samples in the laboratory. The SOPs shall include a list of authorized personnel who have access or keys to secure storage areas.
- 3.5 The Contractor shall have written SOPs describing the method by which the laboratory maintains samples under custody.
- 3.6 The Contractor shall have written SOPs describing the method by which the laboratory maintains the security of any areas identified as secure.
- 3.7 The Contractor shall have written SOPs for tracking the work performed on any particular samples. The tracking SOP shall include:

- A description of the documents used to record sample receipt, sample storage, sample transfers, sample preparations, and sample analyses.
- A description of the documents used to record calibration and QA/QC laboratory work.
- Examples of document formats and laboratory documents used in the sample receipt, sample storage, sample transfer, and sample analyses.
- A narrative step-wise description of how documents are used to track samples.
- 3.8 The Contractor shall have written SOPs for organization and assembly of all documents relating to each SDG. Documents shall be filed on a sample delivery group-specific basis. The procedures shall ensure that all documents including logbook pages, sample tracking records, chromatographic charts, computer printouts, raw data summaries, correspondence, and any other written documents having reference to the SDG are compiled in one location for submission to EPA. The written SOPs shall include:
 - o A description of the numbering and inventory method.
 - A description of the method used by the laboratory to verify consistency and completeness of the CSF.
 - Procedures for the shipment of deliverables packages using custody seals.

4. HANDLING OF CONFIDENTIAL INFORMATION

A Contractor conducting work under this contract may receive EPAdesignated confidential information from the agency. Confidential information must be handled separately from other documentation developed under this contract. To accomplish this, the following procedures for the handling of confidential information have been established.

4.1 All confidential documents shall be under the supervision of a designated document control officer (DCO).

4.2 Confidential Information

Any samples or information received with a request of confidentiality shall be handled as "confidential." A separate locked file shall be maintained to store this information and shall be segregated from other nonconfidential information. Data generated from confidential samples shall be treated as confidential. Upon receipt of confidential information, the DCO will log these documents into a Confidential Inventory Log. The information will then be available to authorized personnel but only after it has been signed out to that person by the DCO. The documents shall be returned to the locked file at the conclusion of each working day. Confidential information may not be reproduced except upon approval by the EPA Technical and Administrative Project Officer. The DCO will enter all copies into the document control system described above. In addition, this information may not be disposed of except upon approval by the EPA project officer. The DCO shall remove and retain the cover page of any confidential information disposed of for one year and shall keep a record on the disposition in the Confidential Inventory Log.

EXHIBIT G

GLOSSARY OF TERMS

GLOSSARY OF TERMS

ALIQUOT - a measured portion of a sample taken for analysis.

ANALYSIS DATE/TIME - the date and military time of the <u>injection</u> of the sample, standard, or blank into the GC/MS or GC system.

BAR GRAPH SPECTRUM - a plot of the mass-to-charge ratio (m/e) versus relative intensity of the ion current.

BLANK - see Method Blank

4-BROMOFLUOROBENZENE (BFB) - compound chosen to establish mass spectral instrument performance for volatile analyses.

CASE - a finite, usually predetermined number of samples collected over a given time period from a particular site. Case numbers are assigned by the Sample Management Office. A Case consists of one or more Sample Delivery Groups.

CHARACTERIZATION - a determination of the approximate concentration range of compounds of interest used to choose the appropriate analytical protocol.

CONCENTRATION LEVEL (low or medium) - characterization of soil samples or sample fractions as low concentration or medium concentration is made on the basis of the laboratory's preliminary screen, <u>not</u> on the basis of information entered on the Traffic Report by the sampler.

CONTINUING CALIBRATION - analytical standard run every 12 hours to verify the calibration of the GC/MS system.

CONTINUOUS LIQUID-LIQUID EXTRACTION - used herein synonymously with the terms continuous extraction, continuous liquid extraction, and liquid extraction.

DAY - unless otherwise specified, day shall mean calendar day.

DECAFLUOROTRIPHENYLPHOSPHINE (DFTPP) - compound chosen to establish mass spectral instrument performance for semivolatile analysis.

EXTRACTABLE - a compound that can be partitioned into an organic solvent from the sample matrix and is amenable to gas chromatography. Extractables include semivolatile (BNA) and pesticide/Aroclor compounds.

IN-HOUSE - at the Contractor's facility.

INITIAL CALIBRATION - analysis of analytical standards for a series of different specified concentrations; used to define the linearity and dynamic range of the response of the mass spectrometer to the target compounds.

INTERNAL STANDARDS - compounds added to every standard, blank, matrix spike, matrix spike duplicate, sample (for VOAs), and sample extract (for semivolatiles) at a known concentration, prior to analysis. Internal standards are used as the basis for quantitation of the target compounds.

LABORATORY - synonymous with Contractor as used herein.

MATRIX - the predominant material of which the sample to be analyzed is composed. For the purpose of this SOW, a sample matrix is either water or soil/sediment. Matrix is <u>not</u> synonymous with phase (liquid or solid).

MATRIX SPIKE - aliquot of a matrix (water or soil) fortified (spiked) with known quantities of specific compounds and subjected to the entire analytical procedure in order to indicate the appropriateness of the method for the matrix by measuring recovery.

MATRIX SPIKE DUPLICATE - a second aliquot of the same matrix as the matrix spike (above) that is spiked in order to determine the precision of the method.

METHOD BLANK (previously termed reagent blank) - an analytical control consisting of all reagents, internal standards and surrogate standards, that is carried through the entire analytical procedure. The method blank is used to define the level of laboratory background contamination.

NARRATIVE (SDG Narrative) - portion of the data package which includes laboratory, contract, Case and sample number identification, and descriptive documentation of any problems encountered in processing the samples, along with corrective action taken and problem resolution. Complete SDG Narrative specifications are included in Exhibit B.

PERCENT MOISTURE - an approximation of the amount of water in a soil/sediment sample made by drying an aliquot of the sample at 105°C. The percent moisture determined in this manner also includes contributions from all compounds that may volatilize at 105°C, including water. Percent moisture is determined from decanted samples and from samples that are not decanted.

PROTOCOL - describes the exact procedures to be followed with respect to sample receipt and handling, analytical methods, data reporting and deliverables, and document control. Used synonymously with Statement of Work (SOW).

PURGE AND TRAP (DEVICE) - analytical technique (device) used to isolate volatile (purgeable) organics by stripping the compounds from water or soil by a stream of inert gas, trapping the compounds on a porous polymer trap, and thermally desorbing the trapped compounds onto the gas chromatographic column.

REAGENT WATER - water in which an interferent is not observed at or above the minimum quantitation limit of the parameters of interest.

RECONSTRUCTED ION CHROMATOGRAM (RIC) - a mass spectral graphical representation of the separation achieved by a gas chromatograph; a plot of total ion current versus retention time.

RECOVERY - a determination of the accuracy of the analytical procedure made by comparing measured values for a fortified (spiked) sample against the known spike values. Recovery is determined by the following equation:

Rec = measured value *Rec = x 100*

RELATIVE RESPONSE FACTOR (RRF) - a measure of the relative mass spectral response of an analyte compared to its internal standard. Relative Response Factors are determined by analysis of standards and are used in the calculation of concentrations of analytes in samples. RRF is determined by the following equation:

$$RRF = \frac{A_x}{A_{is}} \times \frac{C_{is}}{C_x}$$

Where

A - area of the characteristic ion measured

C = concentration

- is internal standard
- x analyte of interest

RESOLUTION - also termed separation, the separation between peaks on a chromatogram, calculated by dividing the depth of the valley between the peaks by the peak height of the smaller peak being resolved, multipled by 100.

SAMPLE - a portion of material to be analyzed that is contained in single or multiple containers and identified by a unique sample number.

SAMPLE DELIVERY GROUP (SDG) - a unit within a single Case that is used to identify a group of samples for delivery. An SDG is a group of 20 or fewer field samples within a Case, received over a period of up to 14 calendar days. Data from all samples in an SDG are due concurrently. A Sample Delivery Group is defined by one of the following, whichever occurs first:

- o Case; or
- o Each 20 field samples within a Case; or
- o Each 14-day calendar period during which field samples in a Case are received, beginning with receipt of the first sample in the Case or SDG.

Samples may be assigned to Sample Delivery Groups by matrix (i.e., all soils in one SDG, all waters in another), at the discretion of the laboratory.

SAMPLE NUMBER (EPA Sample Number) - a unique identification number designated by EPA for each sample. The EPA sample number appears on the sample Traffic Report which documents information on that sample. SEMIVOLATILE COMPOUNDS - compounds amenable to analysis by extraction of the sample with an organic solvent. Used synonymously with Base/Neutral/Acid (BNA) compounds.

SOIL - used herein synonymously with soil/sediment and sediment.

STANDARD ANALYSIS - an analytical determination made with known quantities of target compounds; used to determine response factors.

SURROGATES (Surrogate Standard) - for semivolatiles and pesticides/Aroclors, compounds added to every blank, sample, matrix spike, matrix spike duplicate, and standard; used to evaluate analytical efficiency by measuring recovery. Surrogates are brominated, fluorinated, or isotopically labelled compounds not expected to be detected in environmental media.

SYSTEM MONITORING COMPOUNDS - compounds added to every blank, sample, matrix spike, matrix spike duplicate, and standard for volatile analysis, and used to evaluate the performance of the entire purge and trap-gas chromatographmass spectrometer system. These compounds are brominated or deuterated compounds not expected to be detected in environmental media.

TARGET COMPOUND LIST (TCL) - a list of compounds designated by the Statement of Work (Exhibit C) for analysis.

TENTATIVELY IDENTIFIED COMPOUNDS (TIC) - compounds detected in samples that are not target compounds, internal standards, system monitoring compounds, or surrogates. Up to 30 peaks (those greater than 10% of peak areas or heights of nearest internal standards) are subjected to mass spectral library searches for tentative identification.

TIME - when required to record time on any deliverable item, time shall be expressed as Military Time, i.e., a 24-hour clock.

TRAFFIC REPORT (TR) - an EPA sample identification form filled out by the sampler, which accompanies the sample during shipment to the laboratory and which documents sample condition and receipt by the laboratory.

TWELVE-HOUR TIME PERIOD - The twelve (12) hour time period for GC/MS system instrument performance check, standards calibration (initial or continuing calibration), and method blank analysis begins at the moment of injection of the DFTPP or BFB analysis that the laboratory submits as documentation of instrument performance. The time period ends after 12 hours has elapsed according to the system clock.

VALIDATED TIME OF SAMPLE RECEIPT (VTSR) - the date on which a sample is received at the Contractor's facility, as recorded on the shipper's delivery receipt and Sample Traffic Report.

VOLATILE COMPOUNDS - compounds amenable to analysis by the purge and trap technique. Used synonymously with purgeable compounds.

WIDE BORE CAPILLARY COLUMN - a gas chromatographic column with an internal diameter (ID) that is <u>greater</u> than 0.32 mm. Columns with lesser diameters are classified as narrow bore capillaries.

EXHIBIT H

DATA DICTIONARY AND FORMAT FOR DATA DELIVERABLES IN COMPUTER-READABLE FORMAT

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SECTION I

DESCRIPTION OF DELIVERABLE

1. Introduction

- 1.1 Format A and Format B are the file formats specified for delivery of computer-readable data. Format A is oriented to the structure of the hardcopy reporting forms required by the contract. Format B is organized by analysis, with groups of analyses aggregated to produce files corresponding to an SDG. Information sufficient to generate required hardcopy forms is contained in each format.
- 1.2 The file(s) for a Sample Delivery Group (SDG, see Exhibit A, Section I, B) must be submitted on diskette(s) (see Deliverable, 2.1).
- 1.3 Format A consists of variable length ASCII records.

Format B consists of fixed length ASCII records.

1.4 All information for one SDG must be in one and only one file for Format A.

Format B data may be contained in more than one file as necessary.

2. <u>Deliverable</u>

2.1 The file must be submitted on 5-1/4 inch floppy diskette(s), which may be either double-sided, double density, 360 K-byte or high capacity 1.2 M-byte diskette(s). IBM-compatible, 3.5 inch double-sided, double density 720 K-byte or high density 1.44 M-byte diskettes may also be submitted. The diskettes must be formatted and recorded using MS-DOS Operating System. The diskettes must contain all information relevant to one and only one SDG, and must accompany the hardcopy package for the SDG submitted to the Sample Management Office (see Exhibit B).

Format B data from an entire SDG may not fit onto a single diskette. If a single production run is being split onto multiple diskettes, then all files must start with a type 10 record, and the multiple type 10 records for each file of the same production run must be identical. If it is necessary to split the data from a single sample onto multiple diskettes, then the type 20 (and following) type records for that sample must be repeated. In this situation, columns 7-30, which collectively identify the sample, must be identical on each diskette.

- 2.2 Information on the diskett <u>must correspond exactly</u> with information submitted in the hardcopy data package and on the hardcopy data package forms. Blank or unused records in either format must not be included on the diskettes. If the information submitted in the hardcopy data package forms is changed, the information in the diskette file must be changed accordingly, and a complete diskette containing all the information for the SDG must be resubmitted along with the hard copy at no additional cost to the EPA.
- 2.2 Each diskette must be identified with an external label containing (in this order) the following information:

Disk Density File Name(s) Laboratory Name (optional) Laboratory Code Case Number SAS Number (where applicable)

The format for the File Name must be XXXXX.01A

where XXXXX is the SDG identifier

Dimensions of the label must be in the range 4-3/4" to 5" long by 1-1/4 to 1-1/2" wide.

SECTION II

FORMAT A SPECIFICATION

1. Format Characteristics

- 1.1 Format A is based upon the structure of the hardcopy reporting forms required by the contract. Except for Record Type and the Variable Type, all fields in the format must correspond directly with entries or items on the hardcopy forms.
- 1.2 Format A includes detailed specifications for the required format of each reporting Form's HEADER and DETAIL records. The exact column in which each field is to be contained is specified, as well as the length of the field. Each field 's required contents are specified either as a literal (contained in single quotes) which must appear exactly as shown (without the quotes), or as a variable for which a format is listed in the format column. Each field's required format is specified either as an option of two or more choices (separated by slashes), as MM/DD/YY for a date, as HHMM for time, as a CHARACTER field, or as NUMERIC field.
- 1.3 All Format A fields listed as CHARACTER may contain standard ASCII characters, and must be left-justified and padded to the right with blanks. Formats listed as NUMERIC may contain numeric digits, a decimal place, and a leading minus sign. Numeric entries must be rightjustified and padded to the left with blanks. A positive sign is assumed if no negative sign is entered in a numeric field. A positive sign must not be entered in a numeric field.

Requirements for significant figures and number of decimal places are specified in Exhibit B. The numeric field length are specified such that all possible numeric values can be written to the file. The size of the numeric field indicates the maximum number of digits, decimal, and negative sign if appropriate that can appear in the field at the same time. Therefore, the number reported may need to be rounded (using EPA Rounding Rules) to fit into the field. The rounding must maintain the greatest significance possible providing the field length limitation. In addition, the rounded number that appears on the form, and therefore the feild in the diskette file, must be used in any calculation that may result in other numbers reported on the same form or other forms in the SDG.

The following examples describe the format NUMERIC with a field size of 6:

(Quotation marks indicate the limits of the field described and are not included in the format.)

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RAW DATA VALUE	REPORTED NUMBER
1000000	*999999*
1000	* 1000*
100.1	* 100.1*
100.11	"100.11"
10.1	" 10.1"
0.29	" 0.29 "
0.125559	".12556"
-10.1	" -10.1 "
-1000.1	-1000
-10000.1	*-10000*
-1000000	*-99999*

2. <u>Record Types</u>

- 2.1 Format A consists of variable length ASCII records. The last two bytes of each record must contain "carriage return" and "line feed", respectively. Unused bytes in partially filled fields must be blankfilled.
- 2.2 Format A has three types of records: Header Records, Detail Records and Comment Records.

Туре	Type ID	Contents
Header	Н	Nonrepeating fields which together are unique to the associated hardcopy form
Detail	D	A group of fields that are repeated on a form, and are uniquely positioned by (e.g.) CAS Number or Sequence Number
Comment	C	Nonrepeating fields containing text that comments on information reported on the form

Although there are forms with HEADER records which are not unique per page, the Header record must be included as a place holder, indicating that the DETAIL records follow.

The format for COMMENT records is the same for all forms, and is described after all other formats.

2.3 The first 3 columns in each record contain the FORM ID, identifying the hard copy Low Concentration Organics Reporting Form for which the record contains data. The ID must be left-justified in the feild.

FORM DESCRIBTION

1A	VOLATILE ORGANIC ANALYSIS DATA SHEET
1BC	SEMIVOLATILE ORGANIC ANALYSIS DATA SHEET
1D	PESTICIDE ORGANIC ANALYSIS DATA SHEET
1E	VOLATILE ORGANIC ANALYSIS DATA SHEET (TIC)
1F	SEMIVOLATILE ORGANIC ANALYSIS DATA SHEET (TIC)
2A	WATER VOLATILE SYSTEM MONITORING COMPOUND RECOVERY
2B	SOIL VOLATILE SYSTEM MONITORING COMPOUND RECOVERY
2C	WATER SEMIVOLATILE SURROGATE RECOVERY
2D	SOIL SEMIVOLATILE SURROGATE RECOVERY
2E	WATER PESTICIDE SURROGATE RECOVERY
2F	SOIL PESTICIDE SURROGATE RECOVERY
2.r 3A	WATER VOLATILE MATRIX SPIKE/DUPLICATE RECOVERY
	SOIL VOLATILE MATRIX SPIKE/DUPLICATE RECOVERY
3B	SOIL VOLATILE MAINIA SPIKE/DUPLICATE RECOVERY
30	WATER SEMIVOLATILE MATRIX SPIKE/DUPLICATE RECOVERY
3D	SOIL SEMIVOLATILE MATRIX SPIKE/DUPLICATE RECOVERY
3E	WATER PESTICIDE MATRIX SPIKE/DUPLICATE RECOVERY
3F	SOIL PESTICIDE MATRIX SPIKE/DUPLICATE RECOVERY
4A	VOLATILES METHOD BLANK SUMMARY
4B	SEMIVOLATILES METHOD BLANK SUMMARY
4C	PESTICIDES METHOD BLANK SUMMARY
5A	VOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK
5B	SEMIVOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK
6A	VOLATILE ORGANIC INITIAL CALIBRATION DATA
6B	SEMIVOLATILE ORGANIC INITIAL CALIBRATION DATA
6C	SEMIVOLATILE ORGANIC INITIAL CALIBRATION DATA
6D	PESTICIDES INITIAL CALIBRATION OF SINGLE
	COMPONENT ANALYTES
6E	PESTICIDES INITIAL CALIBRATION OF SINGLE
	COMPONENT ANALYTES
6F	PESTICIDES INITIAL CALIBRATION OF MULTICOMPONENT
	ANALYTES
6G	PESTICIDES RESOLUTION CHECK SUMMARY
7 A	VOLATILES CONTINUING CALIBRATION SUMMARY
7B	SEMIVOLATILES CONTINUING CALIBRATION SUMMARY
7C	SEMIVOLATILES CONTINUING CALIBRATION SUMMARY
7D	PESTICIDES CALIBRATION VERIFICATION SUMMARY
7E	PESTICIDES CALIBRATION VERIFICATION SUMMARY
8A	VOLATILES INTERNAL STANDARD AREA AND RT SUMMARY
8B	SEMIVOLATILES INTERNAL STANDARD AREA AND RT
•••	SUMMARY
8C	SEMIVOLATILES INTERNAL STANDARD AREA AND RT
00	SUMMARY
8D	PESTICIDES ANALYTICAL SEQUENCE
9A	PESTICIDES FLORISIL CARTRIDGE CHECK
9B	PESTICIDES GPC CALIBRATION
10A	PESTICIDES IDENTIFICATION SUMMARY FOR SINGLE
TOU	COMPONENT ANALYTES
10B	PESTICIDES IDENTIFICATION SUMMARY FOR
TOD	MULTICOMPONENT ANALYTES

3. <u>Record Length</u>

Table 3.1 summarizes the length and (in parentheses) the number of records in Format A. The maximum number of detail and comment records is shown, corresponding to a submission of hardcopy forms on which information is written on all possible lines. The Form Totals are the maximum lengths (excluding carriage return/line feed) required for a complete set of each type of form.

Form	Record					
	<u>Header</u>	<u>Detail</u>		Comment		
	<u>H1</u>	<u>D1</u>	<u>D2</u>	<u>C1</u>	<u>C2</u>	
1A	179 ^a (1) ^b	35(33)				
1B	179(1)	35(33)				
10	179(1)	35(31)				
1D	181(1)	35(28)				
1E	21(1)	71(30)				
1 F	21(1)	71(30)				
2 A	11(1)	46(30)			·	
2B	14(1)	46(30)				
2C	11(1)	70(30)				
2D	11(1)	70(30)				
2E	39(1)	52(30)				
2F	39(1)	52(30)				
3 A	27(1)	70(5)	61(5)	77(1)	77(1)	
3B	30(1)	70(5)	61(5)	77(1)	77(1)	
3C	27(1)	70(11)	61(11)	77(1)	77(1)	
3D	30(1)	70(11)	61(11)	77(1)	77(1)	
3E	27(1)	70(6)	61(6)	77(1)	77(1)	
3F	27(1)	70(6)	61(6)	77(1)	77(1)	
4A	86(1)	51(30)		77(1)	77(1)	
4B	87(1)	55(30)		77(1)	77(1)	
4C	139(1)	49(26)		77(1)	77(1)	
5 A	62(1)	20(9)	59(22)			
5B	47(1)	21(13)	59(22)			
6 A	126(1)	69(36)				
6B	111(1)	69(35)				
6C	111(1)	69(35)				
6D	62(1)	64(22)				
6E	62(1)	79(22)				
6F	47(1)	68(40)				
6G	127(1)	45(9)	45(9)			

Table	3.1	Format	A Summe	IY

7A 7B 7C 7D 7E	82(1) 67(1) 67(1) 124(1) 145(1)	58(36) 58(35) 58(35) 70(6) 70(11)	70(13)		
8A 8B 8C 8D	62(1) 47(1) 47(1) 57(1)	57(3) 57(3) 57(3) 57(32)	66(22) 66(22) 66(22)		
9A 9B	57(1) 57(1)	62(11) 62(6)	49(23) 49(26)	77(1) 77(1)	77(1) 77(1)
10A 10B	99(1) 99(1)	97(8) 91(15)	65(3)		

a - length of record in bytes (excluding carriage return/line feed)

b - maximum number of records required for a form.

4. Form Suffix

The fourth and fifth bytes of each record contain the form suffix (AA-ZZ), which must be unique (within a type of form (e.g., Form 1A, Form 2C, etc.,) for each set of records that corresponds to one hardcopy form. For example, the form suffix for records for the first occurrence in the file of a Form 1C must be AA. The second occurrence must be AB, and the twenty-eighth must be BA.

5. <u>Record Listing</u>

The remainder of this section contains detailed specifications for every record required for a full set of hardcopy forms.

FORM I FILE DESCRIPTION (FORM 1)

HEADER RECORD 1 (H1)

COLUMN	I(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'1 <u>A</u> '
4 -			FORM SUFFIX	'AA'-'ZZ'
6 -	7	2 2	RECORD TYPE	'H1'
	19		EPA SAMPLE NO.	CHARACTER
20 -	44	25	LAB NAME	CHARACTER
45 -	55	11	CONTRACT	CHARACTER
56 -	61	6	LAB CODE	CHARACTER
62 -	66	5	CASE NO.	CHARACTER
67 -	72	5 6	SAS NO.	CHARACTER
73 -	77	5 5	SDG NO.	CHARACTER
78 -	82	5	MATRIX	CHARACTER
83 -		12	LAB SAMPLE ID	CHARACTER
95 -	99	5	SAMPLE WT/VOL	NUMERIC
100 -	101	2	SAMPLE WT/VOL UNITS	'G' OR 'ML'
102 -	115	14	LAB FILE ID	CHARACTER
116 -	118	3	LEVEL	'LOW'/'MED'
119 -	126	8	DATE RECEIVED	MM/DD/YY
127 -		2	<pre>% MOISTURE: NOT DEC. DATE ANALYZED</pre>	NUMERIC
129 -	136	8	DATE ANALYZED	MM/DD/YY
137 -	146	10	GC COLUMN	CHARACTER
147 -	150	4	GC COLUMN ID	NUMERIC
151 -			DILUTION FACTOR	NUMERIC
159 -			SOIL EXTRACT VOLUME	NUMERIC
167 -		8 5	SOIL ALIQUOT VOLUME	NUMERIC
175 -		5	CONCENTRATION UNITS	'UG/L' OR 'UG/KG'

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3	FORM NUMBER	'1A'
	2	FORM SUFFIX	'AA'-'2Z'
	2	RECORD TYPE	'D1'
	10	CAS NO.	CHARACTER
	13	RESULT	NUMERIC
	5	QUALIFIER (Q)	CHARACTER

SEMIVOLATILE ORGANICS ANALYSIS DATA SHEET - (FORM 1B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'1B'
4 - 5		FORM SUFFIX	'AA'-'ZZ'
6 - 7		RECORD TYPE	'H1'
8 - 19		EPA SAMPLE NO.	CHARACTER
		LAB NAME	CHARACTER
		CONTRACT	CHARACTER
56 - 61		LAB CODE	CHARACTER
62 - 66	5	CASE NO.	CHARACTER
67 - 72	6	CASE NO. SAS NO.	CHARACTER
73 - 77	5	SDG NO. Matrix	CHARACTER
78 - 82	5	MATRIX	CHARACTER
83 - 94	12		CHARACTER
95 - 99	5	SAMPLE WT/VOL	NUMERIC
100 - 101	2	SAMPLE WT/VOL UNITS	'G' OR 'ML'
102 - 115	14	LAB FILE ID	CHARACTER
116 - 118	3	LEVEL	'LOW'/'MED' MM/DD/YY
119 - 126	8	DATE RECEIVED	MM/DD/YY
127 - 128	2	% MOISTURE	NUMERIC
129 - 129	1	DATE RECEIVED % MOISTURE % MOISTURE (DECANTED) DATE EXTRACTED	'Y'/'N'
130 - 137	8	DATE EXTRACTED	MM/DD/YY
138 - 145	8	CONC. EXTRACT VOLUME	NUMERIC
146 - 153	8	DATE ANALYSED	MM/DD/YY
154 - 161	8	INJECTION VOLUME	NUMERIC
162 - 169	8	DILUTION FACTOR	NUMERIC
170 - 170	1	GPC CLEANUP	'Y'/'N'
171 - 174		PH	NUMERIC
175 - 179	5	CONCENTRATION UNITS	'UG/L' OR 'UG/KG'

DETAIL RECORD 1 (D1)

.

COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'1B'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	'D1'
8 -	17	10	CAS NO.	CHARACTER
18 -	30	13	RESULT	NUMERIC
31 -	35	5	QUALIFIER (Q)	CHARACTER

H-11

SEMIVOLATILE ORGANICS ANALYSIS DATA SHEET - (FORM 1C)

HEADER RECORD 1 (H1)

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'10'
4 - 5	2 2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19		EPA SAMPLE NO.	CHARACTER
20 - 44		LAB NAME	CHARACTER
45 - 55	11	CONTRACT	CHARACTER
56 - 61		LAB CODE	CHARACTER
62 - 66		CASE NO.	CHARACTER
67 - 72		SAS NO.	CHARACTER
73 - 77	5 5	SDG NO.	CHARACTER
78 - 82	5	MATRIX	CHARACTER
83 - 94	12	LAB SAMPLE ID	CHARACTER
95 - 99	5	SAMPLE WT/VOL	NUMERIC
100 - 101	2	SAMPLE WT/VOL UNITS	'G' OR 'ML'
102 - 115	14	LAB FILE ID	CHARACTER
116 - 118	3	LEVEL	'LOW'/'MED'
119 - 126		DATE RECEIVED	MM/DD/YY
127 - 128	2 1	% MOISTURE	NUMERIC
129 - 129		<pre>% MOISTURE (DECANTED)</pre>	'Y'/'N'
130 - 137	8	DATE EXTRACTED	MM/DD/YY
138 - 145		CONC. EXTRACT VOLUME	NUMERIC
146 - 153	8	DATE ANALYSED	MM/DD/YY
154 - 161		INJECTION VOLUME	NUMERIC
162 - 169		DILUTION FACTOR	NUMERIC
170 - 170	1	GPC CLEANUP	'Y'/'N'
171 - 174	4	PH	NUMERIC
175 - 179	5	CONCENTRATION UNITS	'UG/L' OR 'UG/KG'

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 10 13 5	FORM NUMBER FORM SUFFIX RECORD TYPE CAS NO. RESULT QUALIFIER (Q)	'1C' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC CHARACTER

PESTICIDE ORGANICS ANALYSIS DATA SHEET - (FORM 1D)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 4 - 5 6 - 7	3	FORM NUMBER FORM SUFFIX RECORD TYPE EPA SAMPLE NO. LAB NAME CONTRACT	'1D'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19	12	EPA SAMPLE NO.	CHARACTER
20 - 44	25	LAB NAME	CHARACTER
45 - 55	11	CONTRACT	CHARACTER
56 - 61	6	LAB CODE	CHARACTER
62 - 66	5	LAB CODE CASE NO. SAS NO. SDG NO. MATRIX	CHARACTER
67 - 72	6	SAS NO.	CHARACTER
73 - 77	5	SDG NO.	CHARACTER
78 - 82	5	MATRIX	CHARACTER
83 - 94	12	LAB SAMPLE ID	CHARACTER
95 - 99	5	SAMPLE WT/VOL	NUMERIC
100 - 101	2	SAMPLE WT/VOL UNITS	'G' OR 'ML'
102 - 115	14	LAB FILE ID	CHARACTER
116 - 117	2	% MOISTURE	NUMERIC
118 - 118	1	<pre>% MOISTURE (DECANTED)</pre>	'Y'/'N'
119 - 126	8	DATE RECEIVED	MM/DD/YY
127 - 130	4	EXTRACTION	CHARACTER
131 - 138	8	DATE EXTRACTED	MM/DD/YY
139 - 146	8	CONC. EXTRACT VOLUME	NUMERIC
147 - 154	8	DATE ANALYZED	MM/DD/YY
155 - 162	8	INJECTION VOLUME	NUMERIC
163 - 170	8	DILUTION FACTOR	NUMERIC
171 - 171	1	GPC CLEANUP	יצי/יאי
172 - 175	4	PH	NUMERIC
176 - 176	1	SULFUR CLEANUP	'Y'/'N'
177 - 181	5	SDG NO. MATRIX LAB SAMPLE ID SAMPLE WT/VOL SAMPLE WT/VOL UNITS LAB FILE ID & MOISTURE & MOISTURE (DECANTED) DATE RECEIVED EXTRACTION DATE EXTRACTED CONC. EXTRACTED CONC. EXTRACT VOLUME DATE ANALYZED INJECTION VOLUME DILUTION FACTOR GPC CLEANUP PH SULFUR CLEANUP CONCENTRATION UNITS	'UG/L' OR 'UG/KG'

COLUI	MN ((S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 .	-	3 5	3	FORM NUMBER FORM SUFFIX	'1D'
6		5	2 2	RECORD TYPE	'AA'-'ZZ' 'Dl'
8 ·	-	17	10	CAS NO.	CHARACTER
20	-	30	13	RESULT	NUMERIC
31 ·		35	5	QUALIFIER (Q)	CHARACTER

VOLATILE ORGANICS ANALYSIS DATA SHEET - (FORM 1E)

TENTATIVELY IDENTIFIED COMPOUNDS

HEADER RECORD 1 (H1)

COLUMN(S) LE	INGTH CONTENT	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$		IFFIX 'AA'-'ZZ'

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	2 2 10 28 6	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER CAS NO. COMPOUND NAME RT	'1E' 'AA'-'ZZ' 'D1' NUMERIC CHARACTER CHARACTER NUMERIC
54 - 66 67 - 71	_	ESTIMATED CONCENTRATION QUALIFIER (Q)	NUMERIC CHARACTER

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SEMIVOLATILE ORGANICS ANALYSIS DATA SHEET - (FORM 1F)

TENTATIVELY IDENTIFIED COMPOUNDS

HEADER RECORD 1 (H1)

COLUMN (S) LENGTH CONTENTS

FORMAT/CONTENTS

1 -	3	3	FORM NUMBER	'1F'
4 -	5	2	FORM SUFFIX	'AA'-' ZZ'
6 -	7	2	RECORD TYPE	'H1'
8 -	19	12	EPA SAMPLE NO.	CHARACTER
20 -	21	2	NUMBER OF TICS FOUND	NUMERIC

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'1 F'
4 -	5	2	Form suffix	' AA'-' ZZ'
6 -	7	2	RECORD TYPE	'D1'
8 -	9	2	SEQUENCE NUMBER	NUMERIC
10 -	19	10	CAS NO.	CHARACTER
20 -	47	28	COMPOUND NAME	CHARACTER
48 -	53	6	RT	NUMERIC
54 -	66	13	ESTIMATED CONCENTRATION	NUMERIC
67 -	71	5	QUALIFIER (Q)	CHARACTER

FORM II FILE DESCRIPTION (FORM 2)

WATER VOLATILE SYSTEM MONITORING COMPOUND RECOVERY - (FORM 2A)

HEADER RECORD 1 (H1)

COLUMN (S) LENGTH CONTENTS

	•••			•
1 -	3	3	FORM NUMBER	'2A'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
б -	7	2	RECORD TYPE	'H1'
8 -	10	3	LEVEL	CHARACTER
11 -	12	2	PAGE	NUMERIC
13 -	14	2	OF	NUMERIC

DETAIL RECORD 1 (D1)

.

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'2A'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 9	2	SEQUENCE NUMBER	NUMERIC
10 - 21	12	EPA SAMPLE NO.	CHARACTER
22 - 26	5	TOL % RECOVERY	NUMERIC
27 - 27	1	TOL % RECOVERY FLAG	**'/BLANK
28 - 32	5	BFB % RECOVERY	NUMERIC
33 - 33	1	BFB % RECOVERY FLAG	'*'/BLANK
34 - 38	5	DCE % RECOVERY	NUMERIC
39 - 39	1 -	DCE % RECOVERY FLAG	'*' /BLANK
40 - 45	6	OTHER	NUMERIC
46 - 46	1	TOTAL OUT	NUMERIC

FORMAT/CONTENTS

SOIL VOLATILE SYSTEM MONITORING COMPOUND RECOVERY - (FORM 2B)

HEADER RECORD 1 (H1)

COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'2B'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	'H1'
8 -	9	2	PAGE	NUMERIC
10 -	11	2	OF	NUMERIC

COLUMN (S	5) LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{r} 1 - \\ 4 - \\ 6 - \\ 8 - \\ 10 - 2 \\ 22 - 2 \\ 27 - 2 \\ \end{array} $	3 3 5 2 7 2 9 2 21 12 26 5 27 1 22 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. TOL % RECOVERY TOL % RECOVERY FLAG BFB % RECOVERY	FORMAT/CONTENTS '2B' 'AA'-'2Z' 'D1' NUMERIC CHARACTER NUMERIC '*'/BLANK NUMERIC
39 - 3 40 - 4	3 1 8 5 9 1 .5 6 .6 1	BFB % RECOVERY FLAG DCE % RECOVERY DCE % RECOVERY FLAG OTHER TOTAL OUT	'*'/BLANK NUMERIC '*'/BLANK NUMERIC NUMERIC

WATER SEMIVOLATILE SURROGATE RECOVERY - (FORM 2C)

HEADER RECORD 1 (H1)

1 -	3	3	FORM NUMBER	'2C'
4 -	5	2	FORM SUFFIX	'AA'-' ZZ'
6 -	7	2	RECORD TYPE	'H1'
8 -	9	2	PAGE	NUMERIC
10 -	11	2	op	NUMERIC

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'2C'
4 - 5	2	Form suffix	' AA'-' ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 9	2	SEQUENCE NUMBER	NUMERIC
10 - 21	12	EPA SAMPLE NO.	CHARACTER
22 - 26	5	NBZ & RECOVERY	NUMERIC
27 - 27	1 .	NBZ 🕻 RECOVERY FLAG	'*'/BLANK
28 - 32	5	FBP & RECOVERY	NUMERIC
33 - 33	1	FBP & RECOVERY FLAG	'*'/BLANK
34 - 38	5	TPH & RECOVERY	NUMERIC
39 - 39	1	TPH & RECOVERY FLAG	'*'/BLANK
40 - 44	5	PHL & RECOVERY	NUMERIC
45 - 45	1	PHL & RECOVERY FLAG	'*'/BLANK
46 - 50	5	2FP & RECOVERY	NUMERIC
51 - 51	1	2FP & RECOVERY FLAG	'*'/BLANK
52 - 56	5	TBP & RECOVERY	NUMERIC
57 - 57	1	TBP & RECOVERY FLAG	'*'/BLANK
58 - 62	5	2CP & RECOVERY	NUMERIC
63 - 63	1	2CP & RECOVERY FLAG	'*'/BLANK
64 - 68	5	DCB & RECOVERY	NUMERIC
69 - 69	ī	DCB & RECOVERY FLAG	
70 - 70	ī	TOTAL OUT	'*'/BLANK
	-		NUMERIC

SOIL SEMIVOLATILE SURROGATE RECOVERY - (FORM 2D)

HEADER RECORD 1 (H1)

1 -	3	3	FORM NUMBER	201
4 -	5	2	Form Suffix	'AA'+'22'
6 -	7	2	RECORD TYPE	'H1'
8 -	9	2	PAGE	NUMERIC
10 -	11	2	of	NUMERIC

COLUMN (S)	LENGTH	Contents	FORMAT/CONTENTS
$ \begin{array}{rrrrr} 1 & - & 3 \\ 4 & - & 5 \\ 6 & - & 7 \\ 8 & - & 9 \\ 10 & - & 21 \\ \end{array} $	3 2 2 12 5 1 5 1 5 1 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. NBZ & RECOVERY NBZ & RECOVERY FBP & RECOVERY FBP & RECOVERY FBP & RECOVERY FBP & RECOVERY FDH & RECOVERY FLAG PHL & RECOVERY FLAG 2FP & RECOVERY FLAG	'2D' 'AA'-'ZZ' 'D1' NUMERIC CHARACTER NUMERIC '*'/BLANK NUMERIC '*'/BLANK NUMERIC '*'/BLANK
52 - 56 57 - 57 58 - 62	5	2FP & RECOVERY TBP & RECOVERY TBP & RECOVERY TBP & RECOVERY 2CP & RECOVERY 2CP & RECOVERY FLAG	NUMERIC '*'/BLANK NUMERIC '*'/BLANK NUMERIC
63 - 63 64 - 68 69 - 69 70 - 70	2	2CP & RECOVERY FLAG DCB & RECOVERY DCB & RECOVERY FLAG TOTAL OUT	'*'/BLANK NUMERIC '*'/BLANK NUMERIC

HEADER RECORD 1 (H1)

1	-	3	3	FORM NUMBER	'2E'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'H1'
8	-	17	10	GC COLUMN 1	CHARACTER
18	-	21	4	GC COLUMN ID 1	NUMERIC
22	-	31	10	GC COLUMN 2	CHARACTER
32	-	35	4	GC COLUMN ID 2	NUMERIC
36	-	37	2	PAGE	NUMERIC
38	-	39	2	of	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$\begin{array}{rrrr} 1 & - & 3 \\ 4 & - & 5 \\ 6 & - & 7 \\ 8 & - & 9 \\ 10 & - & 21 \\ 22 & - & 26 \\ 27 & - & 27 \\ 28 & - & 32 \\ 33 & - & 33 \\ 34 & - & 38 \\ 39 & - & 39 \\ 40 & - & 44 \\ 45 & - & 45 \end{array}$	3 2 2 12 5 1 5 1 5 1 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. TCX (1) % RECOVERY	FORMAT/CONTENTS '2E' 'AA'-'ZZ' 'D1' NUMERIC CHARACTER NUMERIC '*'/BLANK NUMERIC '*'/BLANK NUMERIC '*'/BLANK NUMERIC '*'/BLANK
46 - 50 51 - 51 52 - 52	5 1 1	OTHER (1) OTHER (2) TOTAL OUT	NUMERIC NUMERIC NUMERIC

SOIL PESTICIDE SURROGATE RECOVERY - (FORM 2F)

HEADER RECORD 1 (H1)

1 -	3	3	FORM NUMBER	'2F'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	'Hl'
8 -	17	10	GC COLUMN 1	CHARACTER
18 -	21	4	GC COLUMN ID 1	NUMERIC
22 -	31	10	GC COLUMN 2	CHARACTER
32 -	35	4	GC COLUMN ID 2	NUMERIC
36 -	37	2	PAGE	NUMERIC
38 -	39	2	OF	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'2F'
4 - 5	2	FORM SUFFIX	'AA'-'2Z'
6 - 7	2	RECORD TYPE	'D1'
8 - 9	2	SEQUENCE NUMBER	NUMERIC
10 - 21 22 - 26	12 5	EPA SAMPLE NO. TCX (1) % RECOVERY TCX (1) % RECOVERY FLAG	CHARACTER NUMERIC
27 - 27 28 - 32 33 - 33	1 5 1	TCX (2) % RECOVERY TCX (2) % RECOVERY FLAG	'*'/BLANK NUMERIC '*'/BLANK
34 - 38	5	DCB (1) % RECOVERY	NUMERIC
39 - 39	1	DCB (1) % RECOVERY FLAG	'*'/BLANK
40 - 44	5	DCB (2) % RECOVERY	NUMERIC
45 - 45	1	DCB (2) % RECOVERY FLAG	'*'/BLANK
46 - 50	5	OTHER (1)	NUMERIC
51 - 51	1	OTHER (2)	NUMERIC
52 - 52	1	TOTAL OUT	NUMERIC

FORM III FILE DESCRIPTION (FORM 3)

WATER VOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3A)

HEADER RECORD 1 (H1)

	COLUMN(S)	(S) LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 33FORM NOMBER'3A'4 - 52FORM SUFFIX'AA'-'ZZ'6 - 72RECORD TYPE'H1'8 - 1912EPA SAMPLE NO.CHARACTER20 - 212RPD: # OUTSIDE LIMITSNUMERIC22 - 232RPD: TOTALNUMERIC24 - 252SPIKE RECOVERY: OUTSIDENUMERIC26 - 272SPIKE RECOVERY: TOTALNUMERIC	6 - 7 8 - 19 20 - 21 22 - 23 24 - 25	5 2 7 2 19 12 21 2 23 2 25 2	RECORD TYPE EPA SAMPLE NO. RPD: # OUTSIDE LIMITS RPD: TOTAL SPIKE RECOVERY: OUTSIDE	'3A' 'AA'-'ZZ' 'H1' CHARACTER NUMERIC NUMERIC NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 24 9 13 13 3 1	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) SAMPLE CONC. (UG/L) MS CONC. (UG/L) MS % RECOVERY MS % RECOVERY FLAG	'3A' 'AA'-'ZZ' 'DI' CHARACTER NUMERIC NUMERIC NUMERIC '*'/BLANK

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 24 9 13 3 1 3 1	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) MSD CONC. (UG/L) MSD % RECOVERY MSD % RECOVERY FLAG % RPD % RPD OUT FLAG	'3A' 'AA'-'ZZ' 'D2' CHARACTER NUMERIC NUMERIC '*'/BLANK NUMERIC '*'/BLANK

SOIL VOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19	12	EPA SAMPLE NO.	CHARACTER
20 - 22	3	LEVEL	'LOW'/'MED'
23 - 24	2	RPD: # OUTSIDE LIMITS	NUMERIC
25 - 26	2	RPD: TOTAL	NUMERIC
27 - 28	2	SPIKE RECOVERY: OUTSIDE	NUMERIC
29 - 30	2	SPIKE RECOVERY: TOTAL	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	יםטי
8 - 31	24	COMPOUND	CHARACTER
32 - 40	9	SPIKE ADDED (UG/L)	NUMERIC
41 - 53	13	SAMPLE CONC. (UG/KG)	NUMERIC
54 - 66	13	MS CONC. (UG/KG)	NUMERIC
67 - 69	3	MS & RECOVERY	NUMERIC
70 - 70	1	MS & RECOVERY FLAG	'*'/BLANK

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 24 9 13 3	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) MSD CONC. (UG/KG) MSD % RECOVERY	'3B' 'AA'-'ZZ' 'D2' CHARACTER NUMERIC NUMERIC NUMERIC
57 - 57 58 - 60 61 - 61	1 3 1	MSD % RECOVERY FLAG % RPD % RPD OUT FLAG	'*'/BLANK NUMERIC '*'/BLANK

WATER SEMIVOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3C)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 12 2 2	FORM NUMBER FORM SUFFIX RECORD TYPE EPA SAMPLE NO. RPD: # OUTSIDE LIMITS RPD: TOTAL	'3C' 'AA'-'ZZ' 'H1' CHARACTER NUMERIC NUMERIC
24 - 25 26 - 27	2 2	SPIKE RECOVERY: OUTSIDE SPIKE RECOVERY: TOTAL	NUMERIC NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3C'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 31	24	COMPOUND	CHARACTER
32 - 40	9	SPIKE ADDED (UG/L)	NUMERIC
41 - 53	13	SAMPLE CONC. (UG/L)	
54 - 66	13	MS CONC. (UG/L)	NUMERIC
67 - 69	3	MS % RECOVERY	NUMERIC
70 - 70	1	MS & RECOVERY FLAG	'*'/BLANK

COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 4 -	3	3	FORM NUMBER	'3C'
	5	2	FORM SUFFIX	'AA'-'2Z'
6 -	7	2	RECORD TYPE	'D2 '
8 -	31	24	COMPOUND	CHARACTER
32 -	40	9	SPIKE ADDED (UG/L)	NUMERIC
41 -	53	13	MSD CONC. (UG/L)	NUMERIC
	56	3	MSD % RECOVERY	NUMERIC
54 - 57 -	57	1	MSD & RECOVERY FLAG	'*'/BLANK
58 -	60	3	% RPD	NUMERIC
61 -	61	1	% RPD OUT FLAG	'*'/BLANK

SOIL SEMIVOLATILE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3D)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3D'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19	12	EPA SAMPLE NO.	CHARACTER
20 - 22	3	LEVEL	'LOW'/'MED'
23 - 24	2	RPD: # OUTSIDE LIMITS	NUMERIC
25 - 26	2	RPD: TOTAL	NUMERIC
27 - 28	2	SPIKE RECOVERY: OUTSIDE	NUMERIC
29 - 30	2	SPIKE RECOVERY: TOTAL	NUMERIC

DETAIL RECORD 1 (D1)

COLI	JMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'3D'
4		5	2	FORM SUFFIX	'AA'-'ZZ'
б	-	7	2	RECORD TYPE	'D1'
8	-	31	24	COMPOUND	CHARACTER
32	-	40	9	SPIKE ADDED (UG/L)	NUMERIC
41	-	53	13	SAMPLE CONC. (UG/KG)	NUMERIC
54	-	66	13	MS CONC. (UG/KG)	NUMERIC
67	-	6 9	3	MS & RECOVERY	NUMERIC
70	-	70	1	MS & RECOVERY FLAG	'*'/BLANK

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$1 - 3 \\ 4 - 5 \\ 6 - 7 \\ 8 - 31 \\ 32 - 40 \\ 41 - 53 \\ 54 - 56 \\ 57 - 57 \\ 58 - 60 \\ $	3 2 24 9 13 3 1 3	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) MSD CONC. (UG/KG) MSD % RECOVERY MSD % RECOVERY FLAG % RPD	'3D' 'AA'-'ZZ' 'D2' CHARACTER NUMERIC NUMERIC '*'/BLANK NUMERIC
61 - 61	T	% RPD OUT FLAG	'*' /BLANK

WATER PESTICIDE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3E)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 12 2 2 2 2 2 2	FORM NUMBER FORM SUFFIX RECORD TYPE EPA SAMPLE NO. RPD: # OUTSIDE LIMITS RPD: TOTAL SPIKE RECOVERY: OUTSIDE SPIKE RECOVERY: TOTAL	'3E' 'AA'-'ZZ' 'H1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC

DETAIL RECORD 1 (D1)

COLUM	N(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 4 - 8 - 32 - 41 - 54 - 57 - 70 -	3 5 7 31 40 53 66 69 70	3 2 24 9 13 13 3	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) SAMPLE CONC. (UG/L) MS CONC. (UG/L) MS % RECOVERY MS % RECOVERY FLAG	'3E' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC
/0 -	,0	-		'*'/BLANK

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 24 9 13 3 1 3 1	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED (UG/L) MSD CONC. (UG/L) MSD % RECOVERY MSD % RECOVERY MSD % RECOVERY FLAG % RPD % RPD OUT FLAG	'3E' 'AA'-'2Z' 'D2' CHARACTER NUMERIC NUMERIC '*'/BLANK NUMERIC '*'/BLANK

SOIL PESTICIDE MATRIX SPIKE/MATRIX SPIKE DUPLICATE RECOVERY (FORM 3F)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3F'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19	12	EPA SAMPLE NO.	CHARACTER
20 - 21	2	RPD: # OUTSIDE LIMITS	NUMERIC
22 - 23	2	RPD: TOTAL	NUMERIC
24 - 25	2	SPIKE RECOVERY: OUTSIDE	NUMERIC
26 - 27	2	SPIKE RECOVERY: TOTAL	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3F'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 31	24	COMPOUND	CHARACTER
32 - 40	9	SPIKE ADDED (UG/L)	NUMERIC
41 - 53	13	SAMPLE CONC. (UG/KG)	NUMERIC
54 - 66	13	MS CONC. (UG/KG)	NUMERIC
67 - 69	3	MS & RECOVERY	NUMERIC
70 - 70	1	MS & RECOVERY FLAG	'*'/BLANK

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3F'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D2 '
8 - 31	24	COMPOUND	CHARACTER
32 - 40	9	SPIKE ADDED (UG/L)	NUMERIC
41 - 53	13	MSD CONC. (UG/KG)	NUMERIC
54 - 56	3	MSD & RECOVERY	NUMERIC
57 - 57	1	MSD & RECOVERY FLAG	'*'/BLANK
58 - 60	3	% RPD	NUMERIC
61 - 61	1	% RPD OUT FLAG	'*'/BLANK

FORM IV FILE DESCRIPTION (FORM 4)

VOLATILE METHOD BLANK SUMMARY - (FORM 4A)

HEADER RECORD 1 (H1)

			•	•	
COLU	JMIN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'4A'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'Hl'
8	-	19	12	EPA SAMPLE NUMBER	CHARACTER
20	-	33	14	LAB FILE ID	CHARACTER
34	-	45	12	LAB SAMPLE ID	CHARACTER
46	-	53	8	DATE ANALYZED	MM/DD/YY
54	-	57	4	TIME ANALYZED	HHMM
58	-	67	10	GC COLUMN	CHARACTER
	-	71	4	GC COLUMN ID	NUMERIC
	-	72	1	HEATED PURGE	'Y'/'N'
73	-	82	10	INSTRUMENT ID	CHARACTER
83	-	84	2	PAGE	NUMERIC
85	-	86	2	OF	NUMERIC

COLUMN (S	5) LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3 3	FORM NUMBER	'4A'
4 -	5 2	FORM SUFFIX	'AA'-'ZZ'
6 -	7 2	RECORD TYPE	'D1'
8 -	9 2	SEQUENCE NUMBER	NUMERIC 2
10 - 2	21 12	EPA SAMPLE NO.	CHARACTER
22 - 3	33 12	LAB SAMPLE ID	CHARACTER
34 - 4	47 14	LAB FILE ID	CHARACTER
48 - 5	51 4	TIME ANALYZED	HHMM

SEMIVOLATILE METHOD BLANK SUMMARY - (FORM 4B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 $4 - 5$ $6 - 7$ $8 - 19$ $20 - 33$ $34 - 45$ $46 - 55$ $56 - 63$ $64 - 68$ $69 - 76$ $77 - 79$ $80 - 83$ $84 - 85$ $86 - 87$	3 2 12 14 12 10 8 5 8 3 4 2 2	FORM NUMBER FORM SUFFIX RECORD TYPE EPA SAMPLE NUMBER LAB FILE ID LAB SAMPLE ID INSTRUMENT ID DATE EXTRACTED MATRIX DATE ANALYZED LEVEL TIME ANALYZED PAGE OF	'4B' 'AA'-'ZZ' 'H1' CHARACTER CHARACTER CHARACTER MM/DD/YY 'SOIL'/'WATER' MM/DD/YY 'LOW'/'MED' HHMM NUMERIC NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 2 12 12 14 8	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. LAB SAMPLE ID LAB FILE ID DATE ANALYZED	'4B' 'AA'-'ZZ' 'D1' NUMERIC 2 CHARACTER CHARACTER CHARACTER MM/DD/YY

PESTICIDE METHOD BLANK SUMMARY - (FORM 4C)

HEADER RECORD 1 (H1)

COLUMN(S) LENGTH CONTENTS

		• •			
1	÷	3	3	FORM NUMBER	'4C'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'H1'
8	-	19	12	EPA SAMPLE NUMBER	CHARACTER
20	-	31	12	LAB SAMPLE ID	CHARACTER
32	-	45	14	LAB FILE ID	CHARACTER
46	-	50	5	MATRIX	'SOIL'/'WATER'
51	-	54	4	EXTRACTION	CHARACTER
55	-	55	1	SULFUR CLEANUP	'Y'/'N'
56	-	63	8	DATE EXTRACTED	MM/DD/YY
64	-	71	8	DATE ANALYZED (1)	MM/DD/YY
72	-	79	8	DATE ANALYZED (2)	MM/DD/YY
80	-	83	4	TIME ANALYZED (1)	HHMM
84	-	87	4	TIME ANALYZED (2)	HHMM
88	-	97	10	INSTRUMENT ID (1)	CHARACTER
98	-	107	10	INSTRUMENT ID (2)	CHARACTER
108	-	117	10	GC COLUMN (1)	CHARACTER
118	-	121	4	GC COLUMN ID (1)	NUMERIC
122	-	131	10	GC COLUMN (2)	CHARACTER
132	-	135	4	GC COLUMN ID (2)	NUMERIC
136	-	137	2	PAGE	NUMERIC
138	-	139	2	OF	NUMERIC
					NOREKIC

DETAIL RECORD 1 (D1)

COLUMN(S)LENGTHCONTENTS1 - 33FORM NUMBER4 - 52FORM SUFFIX6 - 72RECORD TYPE8 - 92SEQUENCE NUMBER10 - 2112EPA SAMPLE NO.22 - 3312LAB SAMPLE ID34 - 418DATE ANALYZED 142 - 498DATE ANALYZED 2

FORMAT/CONTENTS

.

FORMAT/CONTENTS

'4C'
'AA'-'ZZ'
'D1'
NUMERIC
CHARACTER
CHARACTER
MM/DD/YY
MM/DD/YY
• •

FORM V FILE DESCRIPTION (FORM 5)

VOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK - (FORM5A)

BROMOFLUOROBENZENE (BFB)

HEADER RECORD 1 (H1)

COLUM	N(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	• 3	3	FORM NUMBER	'5A'
4 -	- 5	2	FORM SUFFIX	'AA'-' ZZ'
6 -	• 7	2	RECORD TYPE	'H1'
8 -	· 21	14	LAB FILE ID	CHARACTER
22 -	29	8	BFB INJECTION DATE	MM/DD/YY
30 -	. 39	10	INSTRUMENT ID	CHARACTER
40 -	· 43	4	BFB INJECTION TIME	HHMM
44 -	53	10	GC COLUMN	CHARACTER
54 -	· 57	4	GC COLUMN ID	NUMERIC
58 -	- 58	1	HEATED PURGE	יאי/יאי
59 -	• 60	2	PAGE	NUMERIC
61 -	• 62	2	OF	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	15A 1
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	'D1'
8 -	10	3	M/E	NUMERIC
11 -	15	5	% RELATIVE ABUNDANCE	NUMERIC
16 -	20	5	<pre>% MASS (WHERE APPLICABLE)</pre>	NUMERIC

COLUMN(S)		LENGTH	CONTENTS	FORMAT/CONTENTS	
1	-	3	3	FORM NUMBER	'5A'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D2 '
8	-	9	2	SEQUENCE NUMBER	NUMERIC
10		21	12	EPA SAMPLE NO.	CHARACTER
22	-	33	12	LAB SAMPLE ID	CHARACTER
34	-	47	14	LAB FILE ID	CHARACTER
48	-	55	8	DATE ANALYZED	MM/DD/YY
56	~	59	4	TIME ANALYZED	HHMM

SEMIVOLATILE ORGANIC INSTRUMENT PERFORMANCE CHECK - (FORM5B)

DECAFLUOROTRIPHENYLPHOSPHINE (DFTPP)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 14 8 10 4 2	FORM NUMBER FORM SUFFIX RECORD TYPE LAB FILE ID DFTPP INJECTION DATE INSTRUMENT ID DFTPP INJECTION TIME PAGE	'5B' 'AA'-'ZZ' 'H1' CHARACTER MM/DD/YY CHARACTER HHMM NUMERIC
46 - 47	2	of	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3	FORM NUMBER	'5B'
	2	FORM SUFFIX	'AA'-'ZZ'
	2	RECORD TYPE	'D1'
	3	M/E	NUMERIC
	6	% RELATIVE ABUNDANCE	NUMERIC
	5	% MASS (WHERE APPLICABLE)	NUMERIC

COLU	IMN	(S)	LENGTH	Contents	FORMAT/CONTENTS
1 4	-	3 5	3	FORM NUMBER FORM SUFFIX	'5B' 'AA'-'ZZ'
6	-	79	2 2	RECORD TYPE SEQUENCE NUMBER	'D2' NUMERIC
10	-	21	12 12	EPA SAMPLE NO. LAB SAMPLE ID	CHARACTER
	-	33 47	14	LAB FILE ID	CHARACTER CHARACTER
48 56		55 59	8 4	DATE ANALYZED TIME ANALYZED	MM/DD/YY HHMM

FORM VI FILE DESCRIPTION (FORM 6)

VOLATILE ORGANICS INITIAL CALIBRATION DATA - (FORM 6A)

HEADER RECORD 1 (H1)

COLUMN(S) I	ENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 $4 - 5$ $6 - 7$ $8 - 17$ $18 - 25$ $26 - 33$ $34 - 34$ $35 - 38$ $39 - 42$ $43 - 52$ $53 - 56$ $57 - 70$ $71 - 84$ $85 - 98$ $99 - 112$ $113 - 126$	3 2 2 10 8 8 1 4 4 10 4 14 14 14 14 14	FORM NUMBER FORM SUFFIX RECORD TYPE INSTRUMENT ID CALIBRATION DATE 1 CALIBRATION DATE 2 HEATED PURGE CALIBRATION TIME 1 CALIBRATION TIME 1 CALIBRATION TIME 2 GC COLUMN GC COLUMN ID RRF10 LAB FILE ID RRF20 LAB FILE ID RRF50 LAB FILE ID RRF100 LAB FILE ID RRF100 LAB FILE ID	'6A' 'AA'-'ZZ' 'H1' CHARACTER MM/DD/YY MM/DD/YY 'Y'/'N' HHMM HHMM CHARACTER NUMERIC CHARACTER CHARACTER CHARACTER CHARACTER CHARACTER CHARACTER

COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{r} 1 - \\ 4 - \\ 8 - \\ 35 - \\ 40 - \\ 50 - \\ 55 - \\ 60 - \\ 65 - \\ \end{array} $	3 5 7 34 39 49 59 59 69	3 2 27 5 5 5 5 5 5 5 5 5 5	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND RRF10 RRF20 RRF50 RRF100 RRF100 RRF200 AVERAGE RRF % RSD	'6A' 'AA'-'2Z' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC

SEMIVOLATILE ORGANICS INITIAL CALIBRATION DATA - (FORM 6B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'6B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 17	10	INSTRUMENT ID	CHARACTER
18 - 25	8	CALIBRATION DATE 1	MM/DD/YY
26 - 33	8	CALIBRATION DATE 2	MM/DD/YY
34 - 37	4	CALIBRATION TIME 1	HHMM
38 - 41	4	CALIBRATION TIME 2	HHMM
42 - 55	14	RRF20 LAB FILE ID	CHARACTER
56 - 69	14	RRF50 LAB FILE ID	CHARACTER
70 - 83	14	RRF80 LAB FILE ID	CHARACTER
84 - 97	14	RRF120 LAB FILE ID	CHARACTER
98 - 111	14	RRF160 LAB FILE ID	CHARACTER

COLU	IMN	(S)	LENGTH	Contents	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'6B'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D1'
8	-	34	27	COMPOUND	CHARACTER
35	-	39	5	RRF20	NUMERIC
40	-	44	5	RRF50	NUMERIC
45	-	49	5	RRF80	NUMERIC
50	-	54	5	RRF120	NUMERIC
55	-	59	5	RRF160	NUMERIC
60	-	64	5	AVERAGE RRF	NUMERIC
65	-	69	5	* RSD	NUMERIC

SEMIVOLATILE ORGANIC INITIAL CALIBRATION DATA - (FORM 6C)

HEADER RECORD 1 (H1)

COLUMN (S) LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3 3	FORM NUMBER	'6C'
4 -	52	FORM SUFFIX	'AA'-'ZZ'
6 -	7 2	RECORD TYPE	'H1'
8 - 1	7 10	INSTRUMENT ID	CHARACTER
18 - 2	58	CALIBRATION DATE 1	MM/DD/YY
26 - 3	38	CALIBRATION DATE 2	MM/DD/YY
34 - 3	7 4	CALIBRATION TIME 1	HHMM
38 - 4	1 4	CALIBRATION TIME 2	HHMM
42 - 5	5 14	RRF20 LAB FILE ID	CHARACTER
56 - 6	9 14	RRF50 LAB FILE ID	CHARACTER
70 - 8	3 14	RRF80 LAB FILE ID	CHARACTER
84 - 9	7 14	RRF120 LAB FILE ID	CHARACTER
98 - 11	1 14	RRF160 LAB FILE ID	CHARACTER

1 - 3 3 FORM NUMBER '6C' 4 - 5 2 FORM SUFFIX 'AA'-'ZZ' 6 - 7 2 RECORD TYPE 'D1' 8 - 34 27 COMPOUND CHARACTER 35 - 39 5 RRF20 NUMERIC 40 - 44 5 RRF50 NUMERIC 45 - 49 5 RRF80 NUMERIC 50 - 54 5 RRF120 NUMERIC 55 - 59 5 RRF160 NUMERIC 60 - 64 5 AVERAGE RRF NUMERIC	COLUMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
65 - 69 5 % RSD NUMERIC	$\begin{array}{r} 4 \\ - \\ 6 \\ 35 \\ - \\ 40 \\ - \\ 45 \\ - \\ 55 \\ - \\ 55 \\ - \\ 60 \\ - \end{array}$	5 7 34 39 44 49 54 59 64	2 27 5 5 5 5 5 5 5 5	FORM SUFFIX RECORD TYPE COMPOUND RRF20 RRF50 RRF50 RRF80 RRF120 RRF160 AVERAGE RRF	'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC

PESTICIDES INITIAL CALIBRATION OF SINGLE COMPONENT ANALYTES - (FORM 6D)

HEADER RECORD 1 (H1)

COLUMN (S)LENGTHCONTENTSFORMAT/CONTENTS1 - 33FORM NUMBER'6D'

4	-	5	2	FORM SUFFIX		' <u>AA'-'</u> ZZ'
6	-	7	2	RECORD TYPE		'H1'
8	-	17	10	INSTRUMENT ID		CHARACTER
18	-	22	5	LOW LEVEL		NUMERIC
23	-	27	5	MID LEVEL		NUMERIC
28	-	32	5	HIGH LEVEL		NUMERIC
33	-	42	10	GC COLUMN		CHARACTER
43	-	46	4	GC COLUMN ID		NUMERIC
47	-	54	8	DATE ANALYZED	1	MM/DD/YY
55	-	62	8	DATE ANALYZED	2	MM/DD/YY

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'6D'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 34	27	COMPOUND	CHARACTER
35 - 39	5	RT OF LOW STANDARD	NUMERIC
40 - 44	5	RT OF MID STANDARD	NUMERIC
45 - 49	5	RT OF HIGH STANDARD	NUMERIC
50 - 54	5	MEAN RT	NUMERIC
55 - 59	5	RT WINDOW LOWER LIMIT	NUMERIC
60 - 64	5	RT WINDOW UPPER LIMIT	NUMERIC

PESTICIDES INITIAL CALIBRATION OF SINGLE COMPONENT ANALYTES - (FORM 6E)

HEADER RECORD 1 (H1)

COLUM	1(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{r} 1 - \\ 4 - \\ 8 - \\ 18 - \\ 23 - \\ 23 - \\ 33 - \\ 43 - \\ 47 - \\ 55 - \\ \end{array} $	3 5 7 17 22 27 32 42 46 54 54	3 2 10 5 5 10 4 8 8	FORM NUMBER FORM SUFFIX RECORD TYPE INSTRUMENT ID LOW LEVEL MID LEVEL HIGH LEVEL GC COLUMN GC COLUMN ID DATE ANALYZED 1 DATE ANALYZED 2	'6E' 'AA'-'ZZ' 'H1' CHARACTER NUMERIC NUMERIC CHARACTER NUMERIC MM/DD/YY MM/DD/YY

COLU	IMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 4 8 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5		3 5 7 34 44 54 64 74 79	3 2 27 10 10 10 10 5	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND LOW CAL. FACTOR MID CAL. FACTOR HIGH CAL. FACTOR MEAN CAL. FACTORS % RSD	'6E' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC

PESTICIDES INITIAL CALIBRATION OF MULTICOMPONENT ANALYTES - (FORM 6F)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'6F'
4 - 5	2	FORM SUFFIX	'AA'-' ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 17	10	INSTRUMENT ID	CHARACTER
18 - 25	8	DATE ANALYZED 1	MM/DD/YY
26 - 33	8	DATE ANALYZED 2	MM/DD/YY
34 - 43	10	GC COLUMN	CHARACTER
44 - 47	4	GC COLUMN ID	NUMERIC

COLU	JMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1		3	3	FORM NUMBER	'6F'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D1'
8	-	34	27	COMPOUND	CHARACTER
35	-	42	8	AMOUNT	NUMERIC
43	-	43	1	Peak	NUMERIC
44	-	48	5	RT	NUMERIC
49	-	53	5	RT WINDOW LOWER LIMIT	NUMERIC
54	-	58	5	RT WINDOW UPPER LIMIT	NUMERIC
59	-	68	10	CALIBRATION FACTOR	NUMERIC

PESTICIDES RESOLUTION CHECK SUMMARY (FORM 6G)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 4 - 5 6 - 7 8 - 17	3 2 2 10	FORM NUMBER FORM SUFFIX RECORD TYPE GC COLUMN 1 GC COLUMN 1 ID	'6G' 'AA'-'ZZ' 'H1' CHARACTER
$18 - 21 \\ 22 - 31 \\ 32 - 43 \\ 44 - 55 \\ 56 - 63 \\ 64 - 67 \\ 68 - 77 \\ 78 - 81$	4 10 12 12 8 4 10 4	INSTRUMENT ID 1 EPA SAMPLE NO. (STD 1) LAB SAMPLE ID 1 DATE ANALYZED 1 TIME ANALYZED 1 GC COLUMN 2 GC COLUMN 2 ID	NUMERIC CHARACTER CHARACTER MM/DD/YY HHMM CHARACTER NUMERIC
82 - 91 92 - 103 104 - 115 116 - 123 124 - 127	10 12 12 8 4	INSTRUMENT ID 2 EPA SAMPLE NO. (STD 2) LAB SAMPLE ID 2 DATE ANALYZED 2 TIME ANALYZED 2	CHARACTER CHARACTER CHARACTER MM/DD/YY HHMM

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 1 27 5 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER ANALYTE RT RESOLUTION (%)	'6G' 'AA'-'2Z' 'D1' CHARACTER CHARACTER NUMERIC NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 1 27 5 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER ANALYTE RT RESOLUTION (%)	'6G' 'AA'-'ZZ' 'D2' CHARACTER CHARACTER NUMERIC NUMERIC

FORM VII FILE DESCRIPTION (FORM 7)

VOLATILE CONTINUING CALIBRATION SUMMARY - (FORM 7A)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$\begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 10 8 4 14 8 8 1 4 4 10 4	FORM NUMBER FORM SUFFIX RECORD TYPE INSTRUMENT ID CALIBRATION DATE CALIBRATION TIME LAB FILE ID INIT. CALIB. DATE 1 INIT. CALIB. DATE 1 INIT. CALIB. TIME 1 INIT. CALIB. TIME 1 INIT. CALIB. TIME 2 GC COLUMN GC COLUMN ID	'7A' 'AA'-'2Z' 'H1' CHARACTER MM/DD/YY HHMM CHARACTER MM/DD/YY 'Y'/'N' HHMM HHMM CHARACTER NUMERIC

COLUM	I(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{r} 1 - \\ 4 - \\ 6 - \\ 8 - \\ 35 - \\ 40 - \\ 45 - \\ 50 - \\ 55 - \\ \end{array} $	3 5 7 34 39 44 49 54 58	3 2 27 5 5 5 5 4	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND AVERAGE RRF RRF50 MIN RRF % D MAX % D	'7A' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC

SEMIVOLATILE CONTINUING CALIBRATION SUMMARY - (FORM 7B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'7B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 17	10	INSTRUMENT ID	CHARACTER
18 - 25	8	CALIBRATION DATE	MM/DD/YY
26 - 29	4	CALIBRATION TIME	HHMM
30 - 43	14	LAB FILE ID	CHARACTER
44 - 51	8	INIT. CALIB. DATE 1	MM/DD/YY
52 - 59	8	INIT. CALIB. DATE 2	MM/DD/YY
60 - 63	4	INIT. CALIB. TIME 1	HHMM
64 - 67	4	INIT. CALIB. TIME 2	HHMM

DETAIL RECORD 1 (D1)

COL	JMN	(S)	LENGTH	CONTENTS	FORMAT/CONT
1	-	3	3	FORM NUMBER	'7B'
4		5	2	FORM SUFFIX	'AA'-' ZZ'
6	-	7	2	RECORD TYPE	'D1'
8	-	34	27	COMPOUND	CHARACTER
35	-	39	5	AVERAGE RRF	NUMERIC
40	-	44	5	RRF50	NUMERIC
45	-	49	5	MIN RRF	NUMERIC
50	-	54	5	% D	NUMERIC
55	-	58	4	MAX & D	NUMERIC

/CONTENTS

HEADER	RECORD	1 ((H1)	
--------	--------	-----	------	--

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$\begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3	FORM NUMBER	'7C'
	2	FORM SUFFIX	'AA'-'ZZ'
	10	RECORD TYPE	'H1'
	8	INSTRUMENT ID	CHARACTER
	4	CALIBRATION DATE	MM/DD/YY
	14	CALIBRATION TIME	HHMM
	8	LAB FILE ID	CHARACTER
	8	INIT. CALIB. DATE 1	MM/DD/YY
	4	INIT. CALIB. DATE 1	MM/DD/YY
	4	INIT. CALIB. TIME 1	HHMM
	4	INIT. CALIB. TIME 2	HHMM

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
45 - 50 -	3 5 7 39 49 58 58	3 2 27 5 5 5 5 4	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND AVERAGE RRF RRF50 MIN RRF % D MAX % D	'7C' 'AA'-'2Z' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC
55 -	58	4	Maa 5 u	NUMERIC

PESTICIDES CALIBRATION VERIFICATION SUMMARY - (FORM 7D)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'7D'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 17	10	GC COLUMN	CHARACTER
18 - 21	4	GC COLUMN ID	NUMERIC
22 - 29	8	INIT. CALIB. DATE 1	MM/DD/YY
30 - 37	8	INIT. CALIB. DATE 2	MM/DD/YY
38 - 49	12	EPA SAMPLE NO. (PIBLK)	CHARACTER
50 - 61	12	LAB SAMPLE ID. (PIBLK)	CHARACTER
62 - 73	12	EPA SAMPLE NO. (PEM)	CHARACTER
74 - 85	12	LAB SAMPLE ID. (PEM)	CHARACTER
86 - 93	8	DATE ANALYZED (PIBLK)	MM/DD/YY
94 - 101	8	DATE ANALYZED (PEM)	MM/DD/YY
102 - 105	4	TIME ANALYZED (PIBLK)	HIHMM
106 - 109	4	TIME ANALYZED (PEM)	HHMM
110 - 114	5	4,4'-DDT & BREAKDOWN (1)	NUMERIC
115 - 119	5	ENDRIN & BREAKDOWN (1)	NUMERIC
120 - 124	5	COMBINED & BREAKDOWN (1)	NUMERIC

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
4 - 9	9 5 4 5 9 5 7 8	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND RT RT LOWER LIMIT RT UPPER LIMIT CALCULATED AMOUNT NOMINAL AMOUNT	'7D' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC
66 - 70	0 5	RPD	NUMERIC

PESTICIDES CALIBRATION VERIFICATION SUMMARY - FORM 7E)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'7E'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 17	10	GC COLUMN	CHARACTER
18 - 21	4	GC COLUMN ID	NUMERIC
22 - 29	8	INIT. CALIB. DATE 1	MM/DD/YY
30 - 37	8	INIT. CALIB. DATE 2	MM/DD/YY
38 - 49	12	EPA SAMPLE NO. (PIBLK)	CHARACTER
50 - 61	12	LAB SAMPLE ID. (PIBLK)	CHARACTER
62 - 73	12	EPA SAMPLE NO. (INDA)	CHARACTER
74 - 85	12	LAB SAMPLE ID. (INDA)	CHARACTER
86 - 97	12	EPA SAMPLE NO. (INDB)	CHARACTER
98 - 109	12	LAB SAMPLE ID. (INDB)	CHARACTER
110 - 117	8	DATE ANALYZED (PIBLK)	MM/DD/YY
118 - 125	8	DATE ANALYZED (INDA)	MM/DD/YY
126 - 133	8	DATE ANALYZED (INDB)	MM/DD/YY
134 - 137	4	TIME ANALYZED (PIBLK)	HHMM
138 - 141	4	TIME ANALYZED (INDA)	HHMM
142 - 145	4	TIME ANALYZED (INDB)	HHMM

1 - 3 3 FORM NUMBER '7E' 4 - 5 2 FORM SUFFIX 'AA'-'ZZ' 6 - 7 2 RECORD TYPE 'D1' 8 - 34 27 COMPOUND CHARACTER 35 - 39 5 RT NUMERIC 40 - 44 5 RT LOWER LIMIT NUMERIC 45 - 49 5 RT UPPER LIMIT NUMERIC	COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
50 - 578CALCULATED AMOUNTNUMERIC58 - 658NOMINAL AMOUNTNUMERIC66 - 705RPDNUMERIC	$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	2 27 5 5 5 8 8	FORM SUFFIX RECORD TYPE COMPOUND RT RT LOWER LIMIT RT UPPER LIMIT CALCULATED AMOUNT NOMINAL AMOUNT	'7E' 'AA'-'ZZ' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC

COLU	MN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'7E'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D2 '
8	-	34	27	COMPOUND	CHARACTER
35	-	39	5	RT	NUMERIC
40	-	44	5	RT LOWER LIMIT	NUMERIC
45	-	49	5	RT UPPER LIMIT	NUMERIC
50		57	8	CALCULATED AMOUNT	NUMERIC
58	-	65	8	NOMINAL AMOUNT	NUMERIC
66	-	70	5	RPD	NUMERIC

FORM VIII FILE DESCRIPTION (FORM 8)

VOLATILE INTERNAL STANDARD AREA AND RT SUMMARY - (FORM 8A)

HEADER RECORD 1 (H1)

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'8A'
4 -	5	2	FORM SUFFIX	' AA'-' ZZ'
6 -	7	2	RECORD TYPE	'H1'
8 -	21	14	LAB FILE ID (STANDARD)	CHARACTER
22 -	29	8	DATE ANALYZED	MM/DD/YY
30 -	39	10	INSTRUMENT ID	CHARACTER
40 -	43	4	TIME ANALYZED	HIHIMM
44 -	53	10	GC COLUMN	CHARACTER
54 -	57	4	GC COLUMN ID	NUMERIC
58 -	58	1	HEATED PURGE	'Y'/'N'
59 -	60	2	PAGE	NUMERIC
61 -	62	2	OF	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 4 - 5	3	FORM NUMBER FORM SUFFIX	'8A' 'AA'-'22'
6 - 7 8 - 18	2 11	RECORD TYPE ENTRY TYPE	'D1' '12 HOUR STD'/ 'UPPER LIMIT'/
19 - 26 27 - 31	8 5	IS1 (BCM) AREA IS1 (BCM) RT	'LOWER LIMIT' Numeric Numeric
32 - 39 40 - 44 45 - 52	8 5 8	IS2 (DFB) AREA IS2 (DFB) RT IS3 (CBZ) AREA	NUMERIC NUMERIC NUMERIC
53 - 57	5	IS3 (CBZ) RT	NUMERIC

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3 $4 - 5$ $6 - 7$ $8 - 9$ $10 - 21$ $22 - 29$ $30 - 30$ $31 - 35$ $36 - 36$ $37 - 44$ $45 - 45$ $46 - 50$ $51 - 51$ $52 - 59$ $60 - 60$ $61 - 65$	3 2 2 2 12 8 1 5 1 8 1 5 1 8 1 5	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. IS1 (BCM) AREA IS1 (BCM) AREA IS1 (BCM) RT IS1 (BCM) RT IS1 (BCM) RT IS1 (BCM) RT IS1 (BCM) RT IS1 (BCM) RT IS2 (DFB) AREA IS2 (DFB) AREA IS2 (DFB) RT IS2 (DFB) RT IS2 (DFB) RT IS2 (DFB) RT IS2 (DFB) RT IS3 (CBZ) AREA IS3 (CBZ) RT	'8A' 'AA'-'ZZ' 'D2' NUMERIC CHARACTER NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC
66 - 66	1	IS3 (CBZ) RT FLAG	BLANK / '*'

SEMIVOLATILE INTERNAL STANDARD AREA AND RT SUMMARY - (FORM 8B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'8B'
4 - 5	2	FORM SUFFIX	'AA'-'2Z'
6 - 7	2	RECORD TYPE	'H1'
8 - 21	14	LAB FILE ID (STANDARD)	CHARACTER
22 - 29	8	DATE ANALYZED	MM/DD/YY
30 - 39	10	INSTRUMENT ID	CHARACTER
40 - 43	4	TIME ANALYZED	HHMM
44 - 45	2	PAGE	NUMERIC
46 - 47	2	OF	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'8B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 18	11	ENTRY TYPE	'12 HOUR STD'/ 'UPPER LIMIT'/
			'LOWER LIMIT'
19 - 26	8	IS1 (DCB) AREA	NUMERIC
27 - 31	5	ISI (DCB) RT	NUMERIC
32 - 39	8	IS2 (NPT) AREA	NUMERIC
40 - 44	5	IS2 (NPT) RT	NUMERIC
45 - 52	8	IS3 (ANT) AREA	NUMERIC
53 - 57	5	IS3 (ANT) RT	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 $4 - 5$ $6 - 7$ $8 - 9$ $10 - 21$ $22 - 29$ $30 - 30$ $31 - 35$ $36 - 36$ $37 - 44$ $45 - 45$ $46 - 50$ $51 - 51$ $52 - 59$ $60 - 60$ $61 - 65$ $66 - 66$	3 2 2 12 8 1 5 1 8 1 5 1 8 1 5 1 8 1 5 1	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NO. IS1 (DCB) AREA IS1 (DCB) AREA IS1 (DCB) RT IS1 (DCB) RT IS1 (DCB) RT IS1 (DCB) RT IS1 (DCB) RT IS2 (NPT) AREA IS2 (NPT) AREA IS2 (NPT) AREA IS2 (NPT) RT IS2 (NPT) RT IS2 (NPT) RT IS3 (ANT) AREA IS3 (ANT) AREA IS3 (ANT) RT IS3 (ANT) RT IS3 (ANT) RT FLAG	'8B' 'AA'-'ZZ' 'D2' NUMERIC CHARACTER NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC BLANK / '*' NUMERIC
	-		BLANK / '*'

SEMIVOLATILE INTERNAL STANDARD AREA AND RT SUMMARY - (FORM 8C)

HEADER RECORD 1 (H1)

COLUMN	I(S)	LENGTH	Contents	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'8C'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	'H1'
8 -	21	14	LAB FILE ID (STANDARD)	CHARACTER
22 -	29	8	DATE ANALYZED	MM/DD/YY
30 -	39	10	INSTRUMENT ID	CHARACTER
40 -	43	4	TIME ANALYZED	HHIMM
44 -	45	2	PAGE	NUMERIC
46 -	47	2	of	NUMERIC

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	1801
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	יומי
8 - 18	11	ENTRY TYPE	'12 HOUR STD'/ 'UPPER LIMIT'/
			'LOWER LIMIT'
19 - 26	. 8	IS4 (PHN) AREA	NUMERIC
27 - 31	5	IS4 (PHN) RT	NUMERIC
32 - 39	8	IS5 (CRY) AREA	NUMERIC
40 - 44	5	IS5 (CRY) RT	NUMERIC
45 - 52	8	IS6 (PRY) AREA	NUMERIC
53 - 57	5	IS6 (PRY) RT	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	18C1
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D2'
8 - 9	2	SEQUENCE NUMBER	NUMERIC
10 - 21	12	EPA SAMPLE NO.	CHARACTER
22 - 29	8	IS4 (PHN) AREA	NUMERIC
30 - 30	1	IS4 (PHN) AREA FLAG	BLANK / '*'
	5	IS4 (PHN) RT	NUMERIC
	1	IS4 (PHN) RT FLAG	BLANK / '*'
	8	IS5 (CRY) AREA	NUMERIC
	1	IS5 (CRY) AREA FLAG	BLANK / '*'
45 - 45	5	IS5 (CRY) RT	NUMERIC
46 - 50	1	IS5 (CRY) RT FLAG	BLANK / '*'
51 - 51	1	IS6 (PRY) AREA	NUMERIC
52 - 59	8	ISG (PRY) AREA FLAG	BLANK / '*'
60 - 60			NUMERIC
61 - 65	5		BLANK / '*'
66 - 66	1	IS6 (PRY) RT FLAG	DIMIN / *

PESTICIDE ANALYTICAL SEQUENCE - (FORM 8D)

HEADER RECORD 1 (H1)

COLUMN(S) LENGTH CONTENTS

1	-	3	3	FORM NUMBER	'8D'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'H1'
8	-	17	10	GC COLUMN	CHARACTER
18	-	21	4	GC COLUMN ID	NUMERIC
22	-	29	8	INIT. CALIB. DATE 1	MM/DD/YY
30	-	37	8	INIT. CALIB. DATE 2	MM/DD/YY
38	-	47	10	INSTRUMENT ID	CHARACTER
48	-	52	5	MEAN RT (TCX)	NUMERIC
53	-	57	5	MEAN RT (DCB)	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN (5) L	ength	CONTENTS	FORMAT/CONTENTS
1 -	3	3	FORM NUMBER	'8D'
4 -	5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	7	2	RECORD TYPE	' <u>D1</u> '
8 -	9	2	SEQUENCE NUMBER	NUMERIC
10 - 3	21	12	EPA SAMPLE NUMBER	CHARACTER
22 - 3	33	12	LAB SAMPLE ID	CHARACTER
34 - 4	41	8	DATE ANALYZED	MM/DD/YY
42 - 4	45	4	TIME ANALYZED	HIHMM
46 - 3	50	5	TCX RT	NUMERIC
51 - !	51	1	TCX RT FLAG	CHARACTER
52 - 3	56	5	DCB RT	NUMERIC
57 - 9	57	1	DCB RT FLAG	CHARACTER

FORMAT/CONTENTS

FORM IX FILE DESCRIPTION (FORM 9)

PESTICIDE FLORISIL CARTRIDGE CHECK - (FORM 9A)

HEADER RECORD 1 (H1)

COLU	JMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'9 A '
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'H1'
8	-	17	10	FLOR. CART. LOT NO.	CHARACTER
18	-	25	8	DATE OF ANALYSIS	MM/DD/YY
26	-	35	10	GC COLUMN 1	CHARACTER
36	-	39	4	GC COLUMN 1 ID	NUMERIC
40	-	49	10	GC COLUMN 2	CHARACTER
50	-	53	4	GC COLUMN 2 ID	NUMERIC
54	-	55	2	PAGE	NUMERIC
56	-	57	2	OF	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	Contents	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'9 A '
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 34	27	COMPOUND	CHARACTER
35 - 42	8	SPIKE ADDED	NUMERIC
43 - 50	8	SPIKE RECOVERED	NUMERIC
51 - 55	5	\$ REC	NUMERIC
56 - 56	1	% REC FLAG	CHARACTER
57 - 62	6	QC LIMIT	'80-110'

COLU	JMN	(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1		3	3	FORM NUMBER	'9A'
4		5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D2 '
8	-	9	2	SEQUENCE NUMBER	NUMERIC
10	-	21	12	EPA SAMPLE NUMBER	CHARACTER
22	-	33	12	LAB SAMPLE ID	CHARACTER
34	-	41	8	DATE ANALYZED 1	MM/DD/YY
42	-	49	8	DATE ANALYZED 2	MM/DD/YY

PESTICIDE GPC CALIBRATION - (FORM 9B)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER FORM SUFFIX	'98' 'AA'-'22'
4 - 5 6 - 7	2	RECORD TYPE	'H1'
8 - 17 18 - 25	10 8	GPC COLUMN CALIBRATION DATE	CHARACTER MM/DD/YY
26 - 35 36 - 39	10 4	GC COLUMN 1 GC COLUMN 1 ID	CHARACTER NUMERIC
40 - 49	10	GC COLUMN 2 GC COLUMN 2 ID	CHARACTER NUMERIC
50 - 53 54 - 55	2	PAGE	NUMERIC
56 - 57	2	OF	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 27 8 8 5 1 6	FORM NUMBER FORM SUFFIX RECORD TYPE COMPOUND SPIKE ADDED SPIKE RECOVERED % REC % REC FLAG QC LIMIT	'9B' 'AA'-'2Z' 'D1' CHARACTER NUMERIC NUMERIC NUMERIC CHARACTER '80-110'

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
$ \begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$	3 2 2 12 12 8 8	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER EPA SAMPLE NUMBER LAB SAMPLE ID DATE ANALYZED 1 DATE ANALYZED 2	'9B' 'AA'-'ZZ' 'D2' NUMERIC CHARACTER CHARACTER MM/DD/YY MM/DD/YY

FORM X FILE DESCRIPTION (FORM 10)

PESTICIDE IDENTIFICATION SUMMARY FOR SINGLE COMPONENT ANALYTES - (FORM 10A)

HEADER RECORD 1 (H1)

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'10A'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'H1'
8 - 19	12	EPA SAMPLE NO.	CHARACTER
20 - 31	12	LAB SAMPLE ID	CHARACTER
32 - 39	8	DATE ANALYZED 1	MM/DD/YY
40 - 47	8	DATE ANALYZED 2	MM/DD/YY
48 - 57	10	INSTRUMENT ID 1	CHARACTER
58 - 67	10	INSTRUMENT ID 2	CHARACTER
68 - 77	10	GC COLUMN 1	CHARACTER
78 - 81	4	GC COLUMN 1 ID	NUMERIC
82 - 91	10	GC COLUMN 2	CHARACTER
92 - 95	4	GC COLUMN 2 ID	NUMERIC
96 - 97	2	PAGE	NUMERIC
98 - 99	2	OF	NUMERIC

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3 $4 - 5$ $6 - 7$ $8 - 9$ $10 - 36$ $37 - 41$ $42 - 46$ $47 - 51$ $52 - 64$ $65 - 69$ $70 - 74$ $75 - 79$	LENGIA 3 2 2 27 5 5 5 13 5 5 5 13	FORM NUMBER FORM SUFFIX RECORD TYPE SEQUENCE NUMBER ANALYTE COLUMN 1 RT COLUMN 1 RT LOWER LIMIT COLUMN 1 RT UPPER LIMIT COLUMN 2 RT COLUMN 2 RT COLUMN 2 RT LOWER LIMIT COLUMN 2 RT UPPER LIMIT COLUMN 2 CONCENTRATION	'10A' 'AA'-'ZZ' 'D1' NUMERIC CHARACTER NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC NUMERIC
80 - 92 93 - 97	13 5	* D	NUMERIC

PESTICIDE IDENTIFICATION SUMMARY FOR MULTICOMPONENT ANALYTES - (FORM 10B)

HEADER RECORD 1 (H1)

COLUMN(S) LENGTH CONTENTS

•	-	•		11001
1 -	• 3	3	FORM NUMBER	'10B'
4 -	• 5	2	FORM SUFFIX	'AA'-'ZZ'
6 -	· 7	2	RECORD TYPE	'H1'
8 -	• 19	12	EPA SAMPLE NO.	CHARACTER
20 -	· 31	12	LAB SAMPLE ID	CHARACTER
32 -	• 39	8	DATE ANALYZED 1	MM/DD/YY
40 -	47	8	DATE ANALYZED 2	MM/DD/YY
48 -	· 57	10	INSTRUMENT ID 1	CHARACTER
58 -	• 67	10	INSTRUMENT ID 2	CHARACTER
68 -	· 77	10	GC COLUMN 1	CHARACTER
78 -	· 81	4	GC COLUMN 1 ID	NUMERIC
82 -	• 91	10	GC COLUMN 2	CHARACTER
92 -	· 95	4	GC COLUMN 2 ID	NUMERIC
96 -	• 97	2	PAGE	NUMERIC
98 -	• 99	2	OF	NUMERIC

DETAIL RECORD 1 (D1)

COLUMN (S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'10B'
4 - 5	2	FORM SUFFIX	'AA'-'ZZ'
6 - 7	2	RECORD TYPE	'D1'
8 - 34	27	ANALYTE	CHARACTER
35 - 35	1	PEAK	NUMERIC
36 - 40	5	COLUMN 1 RT	NUMERIC
41 - 45	5	COLUMN 1 RT LOWER LIMIT	NUMERIC
46 - 50	5	COLUMN 1 RT UPPER LIMIT	NUMERIC
51 - 63	13	COLUMN 1 CONCENTRATION	NUMERIC
64 - 68	5	COLUMN 2 RT	NUMERIC
69 - 73	5	COLUMN 2 RT LOWER LIMIT	NUMERIC
7 4 - 78	5	COLUMN 2 RT UPPER LIMIT	NUMERIC
7 9 - 91	13	COLUMN 2 CONCENTRATION	NUMERIC

DETAIL RECORD 2 (D2)

COLUMN(S)			LENGTH	CONTENTS	FORMAT/CONTENTS
1	-	3	3	FORM NUMBER	'10B'
4	-	5	2	FORM SUFFIX	'AA'-'ZZ'
6	-	7	2	RECORD TYPE	'D2 '
8	-	34	27	ANALYTE	CHARACTER
35	-	47	13	COLUMN 1 MEAN CONC.	NUMERIC
48	-	60	13	COLUMN 2 MEAN CONC.	NUMERIC
61	-	65	5	ኛ D	NUMERIC

FORMAT/CONTENTS

COMMENT RECORDS

COMMENT records are optional for some Forms as specified in Table 3.1 in Section I of this Exhibit, and as appears on the hard copy of the Reporting Forms in Section IV of Exhibit B. They must come after the DETAIL records and be formatted as follows:

COLUMN(S)	LENGTH	CONTENTS	FORMAT/CONTENTS
1 - 3	3	FORM NUMBER	'3A'/'3B'/'3C' '3D'/'3E'/'3F' '4A'/'4B'/'4C' '9A'
4 - 5 6 - 7 8 - 77	2 2 70	FORM SUFFIX RECORD TYPE COMMENT	'AA'-' ZZ' 'C1'/'C2' CHARACTER

SECTION III

FORMAT B SPECIFICATION

1. Format Characteristics

- 1.1 Format B is an implementation of the EFA Agency Standard for Electronic Data Transmission based upon analytical results and ancillary information required by the contract. All data generated by a single analysis are grouped together, and the groups are aggregated to produce files that report data from an SDG. Because Format B is based on the Agency Standard, some non-CLP fields have been replaced by blanks.
- 1.2 Format B includes detailed specifications for the required format of each record. The exact position in the fixed-length record where each field is to be contained is specified, as well as the length of the field. Each field's required contents are specified as literal (contained in quotes) which must appear exactly as shown (without quotes), or as a variable for which format and/or descriptions are listed in the format/contents column. Options and examples are listed for most fields. For fields where more than three options are available, a list and description of options are supplied on a separate page following the record descriptions. Fields are separated from each other by blanks.
- 1.3 All Format B character fields must be left-justified and padded to the right with blanks. Numeric fields may contain numeric digits, a decimal place, and a leading minus sign. A positive sign is assumed if no negative sign is entered in a numeric field and must not be entered into any numeric field. Numeric entries must be right justified and padded to the left with blanks. Allowance is made within the Agency Standard for exponential notation.

Requirements for significant figures and number of decimal places are specified in Exhibit B. The numeric field length are specified such that all possible numeric values can be written to the file. The size of the numeric field indicates the maximum number of digits, decimal, and negative sign if appropriate that can appear in the field at the same time. Therefore, the number reported may need to be rounded (using EPA Rounding Rules) to fit into the field. The rounding must maintain the greatest significance possible providing the field length limitation. In addition, the rounded number that appears on the form, and therefore the field in the diskette file, must be used in any calculation that may result in other numbers reported on the same form or other forms in the SDG.

2. <u>Record Types</u>

2.1 Format B consists of fixed-length 80-byte ASCII records. The last two bytes of each record must contain "carriage return" and "line feed", respectively. Unused bytes in partially filled fields must be blank-filled. 2.2 Format B consists of eleven record types than can be summarized in four groups, designated by the first record type in each group:

Type	<u>Type ID</u>	Contents	
Run Header	10	Information pertinent to a group of samples processed in a continuous sequence; usually several per SDG	
Sample Header	20	Sample identifying, qualifying, and linking information	
Results Record	30	Analyte results and qualifications	
Comments Record	90	Free form comments and flags	

A separate run header is used for volatiles, semivolatiles, and each column analysis for pesticides (four Type 10 series for VOA/SV/PEST SDG). The 20 series records are used to link samples within an SDG to the corresponding calibrations, blanks, and so on for screening purposes. The 30 series records contain the actual analytical results by analyte within each sample. The 10, 20, and 30 records are associated with each other by their position in the file (i.e., 30 series records follow the corresponding 20 series, which in turn follow the 10

2.3 The sequence of records for Format B files is as follows: A Run Header (type 10) record must be present once and once only (per file) as the first record in a file. Therefore, a complete VOA/SV/PEST SDG will consist of sevral files.

Each environmental sample, calibration, or quality control sample is represented by a group composed of type 20, 21, 22, 23, and 27 records, which hold sample level identifying information, followed by type 30, 31, and 32 records for each method analyte or standard in the sample. The type 20 record holds a count for the number of method analytes being determined, and includes all target compounds plus any tentatively identified compounds. Type 20 records must occur in the order of sample analysis. The type 20 records for quality control items have the additional rule that the LF1 record must occur before the LF2 record, but the records need not be adjacent (matrix spike data). In addition, a type 20 record is used as a header for any additional run-wide data that must be reported for each method analyte (such as mean response factors). Unique identifiers given on page H-85 are used in place of "QC codes" to indicate the type of data that follows. Type 30 records for each analyte must occur in the order specified on

Type 90 comment records may be defined to occupy any position except before the type 10 (header) record.

EXAMPLE OF THE SEQUENCE OF RECORD TYPES IN A FILE

10						
11		Contains additional run-wide information if required.				
	20		Occurs once for each sample, calibration, mean response factor, matrix spike duplicate result, etc Acts as a header.			
	21					
	22		Contains additional information for samples.			
	23					
	27					
	27	30	Occurs once for each final analytical result. Reports			
		50	the value being determined as defined by the type 20.			
			• • • • •			
			33 Reports component names if necessary.			
		30	Values for the next analyte or parameter being measured.			
			31 Additional data may vary for each parameter, and records			
			32 may occur in any order. Multiple occurrences of the			
			32 same record type, however, must be consecutive.			
			33			
		30	Continues for as many as are necessary.			
			31			
			32			
			33			
		30				
			31			
			32			
			33			
•	20		Next Sample Header record - The following applies to the next			
	21		sample or other group of data.			
	22					
		30				
			31			
			32			
			33			
		30				
			31			
			32			
			33			
			etc.			
	20					
	21					
		30				
			31			
			32			
			33			
			etc.			

3. <u>File/Record Integrity</u>

All record types shall contain the following check fields to ensure file and record integrity:

Record	Field	Field	<u>Remarks</u>
<u>Position</u>	<u>Length</u>	<u>Contents</u>	
1-2 72-74	2 3	Record type Record sequence number within file	"10" or as appropriate 000-999, repeated as necessary
75-78	4	Record checksum	Four hexadecimal digits(*)
79-80	2	Must contain CR and LF	

(*) The checksum is the sum of the thirty-five Integers that make up the data in columns 1 to 70, when data are represented in the format 35A2 on processors which store data bytes in left to right order. The sum is taken modulo 65536 (2^{16}) and represented as four hexadecimal digits. For processors which use an A70 character representation of data, the checksum is the sum of all the even character position values plus 256 times the sum of all the odd character position values.

4. Dates and Times

Date or time-of-day information consists of successive groups of one or two decimal digits, each separated by blanks. Dates are given in the order YY MM DD, and times as HH MM. All hours must be given as 0 to 23 using a 24 hour clock and must be local time.

5. Record Listing

Following is a listing of every record type required to report data from a single SDG.

PRODUCTION RUN HEADER RECORD (TYPE 10)

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3-18	2 16	RECORD TYPE blank	"10"
19-23	5	MEASUREMENT TYPE	"GC/MS" for VOA and SV
24-26	3	blank	"GC/EC" for Pesticides.
27-30	4	METHOD NUMBER	"6241" for Volatiles; "625C" for Semivolatiles; "6081"for Pesticides
31-35	5	blank	
36-41 42-51	6 10	LAB CODE blank	Character
52-61 62	10 1	CONTRACT NUMBER blank	Character
63-68	6	INSTRUMENT ID	e.g. 59951G; provided by contract lab; left justi- fied; must be unique and permanent within lab. Firsi

contract lab; left justified; must be unique and permanent within lab. First four characters are vendors designator; fifth and sixth are lab assigned so each instrument is unique. Use: To describe chromatograph conditions. Must be present for volatiles and pesticides. Is optional for semivolatiles.

Position: Follows type 10.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	RECORD TYPE blank	"11"
4-11	8	GC COLUMN	Left justified. e.g., SP2330
12-17	6	blank	e.g., 512550
18-21	4	GC COLUMN ID	mm e.g., 2 or .3 Right justified

.

SAMPLE HEADER DATA RECORD (TYPE 20)

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3-6	2 4	RECORD TYPE blank	"20"
7-13	7	EPA SAMPLE NO.	Left justified. Raw Sample No. only; no suffixes.
14-15	2	blank	NO. ONLY, NO SULLINES.
16	1	MATRIX	"O" if not applicable (calibrations, tunes, etc.) "1" for water
17	1	blank	"H" for soil
18-20	3	QC CODE	Codes type of data to be
21	1	blank	reported (see page H-85)
22-24	3	SAMPLE QUALIFIER	"RIN" = reanalyzed "REX" = reprepared
25	1	blank	"REJ" - rejected "SRN" - diluted
26-30	5	CASE NUMBER	Required for blanks,
31-33	3	blank	EPA samples, optional otherwise
34-38 39	5	SDG NO. blank	Left justified
40-47	8	DATE ANALYZED	YY MM DD
48	1	blank	
49-53	5	TIME ANALYZED	HH MM
54-56	3	blank	
57	1	SAMPLE WT/VOL UNITS	"L" - liters for water
58	1	blank	"K" - kilograms for soil
59-66	8	SAMPLE WT/VOL	right justified; see Note
67	1	blank	
68-70	3	ANALYTE COUNT	Numeric; 1-3 decimal digits; right justified. Counts all analytes including TIC's. For calibrations, count all analytes, surrogates, and DETER (SER if mixed in

NOTE: SAMPLE WT/VOL is the volume in liters for liquids and the wet weight in kilograms for solids. The Sample Units Code indicates which units are in use for the current sample. Leave zero or blank if not applicable. SAMPLE WT/VOL includes purge volume

DFTPP/BFB if mixed in

injection.

SAMPLE HEADER DATA RECORD (TYPE 21)

Use: Continuation of type 20.

Position: Follows the type 20 to which it applies.

13-175Diam"N" - Sonication18-236SAS NO. blankLeave blank if none. Optional for calibration241blankGrow instrument data system. Lab File ID for or Semivolatiles; Lab Sa ID for Pesticides.361blank37-448 blankDATE OF EXTRACTION blank37-448 blankDATE OF EXTRACTION blank48-558 blankDATE RECEIVED56-649 blankINJECTION/ALIQUOT VOLUME ul, Injection Volume for				
1-2 1 blank 3 1 blank 4 1 PURGE "H" for heated 5 1 blank "L" = low 6 1 LEVEL "L" = low 7 1 blank (See note). 8 1 CLEAN-UP "G" for GPC or blank 9-11 3 blank "S" - Separatory Funnel 12 1 EXTRACTION "S" - Separatory Funnel 13-17 5 blank "C" - Continuous Liq-Liq 13-17 5 blank Optional for calibration 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Se ID for Pesticides. 36 1 blank Joing Joing 37-44 8 DATE OF EXTRACTION YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank Joing Joing	POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
4 1 blank 5 1 blank 6 1 LEVEL "L" = low "M" = medium 7 1 blank (See note). 8 1 CLEAN-UP "G" for GPC or blank 9-11 3 blank "S" - Separatory Funnel 12 1 EXTRACTION "S" - Separatory Funnel 13-17 5 blank "C" - Continuous Liq-Liq 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Se ID for Pesticides. 36 1 blank Jor Pesticides. 36 1 blank YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank JinkCTION/ALIQUOT VOLUME ul, Injection Volume for				."21"
6 1 LEVEL "M" - medium 7 1 blank (See note). 8 1 CLEAN-UP "G" for GPC or blank 9-11 3 blank "S" - Separatory Funnel 12 1 EXTRACTION "S" - Separatory Funnel 13-17 5 blank "C" - Continuous Liq-Liq 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Sa ID for Pesticides. 36 1 blank Joank 37-44 8 DATE OF EXTRACTION YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank Joank		-		"H" for heated
7 1 Dimin 8 1 CLEAN-UP "G" for GPC or blank 9-11 3 blank "S" - Separatory Funnel 12 1 EXTRACTION "S" - Separatory Funnel 13-17 5 blank "C" - Continuous Liq-Liq 13-17 5 blank "S" - Separatory Funnel 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Sa ID for Pesticides. 36 1 blank Jon Festicides. 36 1 blank YMM DD 45-47 3 blank YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank ul, Injection Volume for	6	1	LEVEL	
8 1 Diank 9-11 3 blank 12 1 EXTRACTION "S" - Separatory Funnel 13-17 5 blank "C" - Continuous Liq-Liq 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Sa ID for Pesticides. 36 1 blank Jone Fextraction YY MM DD 45-47 3 blank YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank UNIECTION/ALIQUOT VOLUME ul, Injection Volume for	7	1	blank	(See note).
12 1 Infinition 13-17 5 blank "C" - Continuous Liq-Liq 18-23 6 SAS NO. Leave blank if none. 24 1 blank Optional for calibration 25-35 11 LAB FILE/SAMPLE ID From instrument data system. Lab File ID for or Semivolatiles; Lab Sa ID for Pesticides. 36 1 blank Joint Content of the system of the system of the system of the system. 37-44 8 DATE OF EXTRACTION YY MM DD 45-47 3 blank YY MM DD 48-55 8 DATE RECEIVED YY MM DD 56-64 9 blank ul, Injection Volume for				"G" for GPC or blank
18-23 246SAS NO.241blankOptional for calibration25-3511LAB FILE/SAMPLE IDFrom instrument data system. Lab File ID for or Semivolatiles; Lab Sa 				"C" - Continuous Liq-Liq
25-3511LAB FILE/ SINCL Construction361blank37-448DATE OF EXTRACTION45-473blank48-558DATE RECEIVED56-649blank6INJECTION/ALIQUOT VOLUMEul, Injection Volume for				Leave blank if none. Optional for calibrations
361DATE OF EXTRACTIONYY MM DD37-448DATE OF EXTRACTIONYY MM DD45-473blankYY MM DD48-558DATE RECEIVEDYY MM DD56-649blankJNJECTION/ALIQUOT VOLUMEul, Injection Volume for	25-35	11	LAB FILE/SAMPLE ID	system. Lab File ID for VOA or Semivolatiles; Lab Sample
37-443blank45-473blank48-558DATE RECEIVED56-649blank6INJECTION/ALIQUOT VOLUMEul, Injection Volume for	36	1	blank	
48-55 8 DATE RECEIVED TO LONG THE SECONDATE STATE OF SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SECONDATE SEC				YY MM DD
56-64 JULICTION/ALIQUOT VOLUME ul, Injection Volume for	48-55	8	DATE RECEIVED	YY MM DD
INJECTION/ALIQUOT VOLUME ul, Injection Volume for	56-64	9	blank	
65-70 BNA's and PEST's Volume aliquot for VOA's.	65-70	6	INJECTION/ALIQUOT VOLUME	ul, Injection Volume for BNA's and PEST's Volume of aliquot for VOA's.

Note: The LEVEL is an estimate of overall level for all analytes. It is blank for calibrations and tunes.

SAMPLE CONDITIONS RECORD (TYPE 22)

Use: Continuation of type 20. Used to describe additional sample conditions. Position: Follows the type 20 and 21 to which it applies.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	RECORD TYPE blank	*22*
4-11	8	CALIBRATION DATE	YY MM DD; see Note 1 (Date of Source of the
12	1	blank	response factor used)
13-17	5	CALIBRATION TIME	HH MM
18	1	blank	
19-29	11	CALIBRATION FILE ID	Lab File ID of associated continuing calibration, or "AVERAGE", position 21-27 and padded with blanks (Note 2)
30	1	blank	
31-34	4	PH	XX or XX.X or zero, right justified
35	1	blank	
36-37 38	2 1	* MOISTURE blank	decanted or nondecanted value used to correct to dry weight; use zero if not applicable
39-40	2	AUXILLIARY PERCENT MOISTURE	report nondecanted % water if water was decanted; use
41	1	blank	zero if not applicable
42-46	5	CONC. EXTRACT VOLUME	ul, e.g., 1,000 or 500 (See Note 3)
47	1	blank	
48-54	7	DILUTION FACTOR	Right justified; from Form 1 e.g., 2000

Note 1: For average, use the date and time average was calculated.

Note 2: If "AVERAGE" is entered, then Form 8 will be constructed using the initial calibration with a QC code of "CLD". This field must match columns 25-35 of record type 21 for the associated calibration.

Note 3: Use the initial extract volume adjusted (multiplied) by all contract-mandated dilutions that are to be excluded from the dilution factor.

ASSOCIATED INJECTION AND COUNTER RECORD (TYPE 23)

Use: Continuation of type 20. Used to identify associated blanks and tunes, and to count the number of surrogates and spikes outside of QC limits and the number of TIC compounds. Used for Forms 3, 4, and 5.

Position: Follows the type 20, 21, and 22 to which it applies.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
	2	RECORD TYPE	"23"
1-2	1	blank	
3 4	1	TUNE LABEL	"P" or blank Labels data as "tune" data.
5	1	blank	
6-13	8	INJECTION DATE	YY MM DD. Acquisition date of tune to be linked with
14	1	blank	<pre>sample (if applicable).</pre>
15-19 20	5 1	INJECTION TIME blank	HH MM (if applicable)
21-31 32	11 1	DFTPP/BFB LAB FILE ID blank	From instrument data system
33	1	BLANK LABEL	"B" or blank Labels data as "blank" data.
34	1	blank	(use blank if none)
35-42	8	BLANK INJECTION DATE	YY MM DD. Acquisition date of method blank to be linked
43	1	blank	with sample (if applicable).
44-48 49	5 1	BLANK INJECTION TIME blank	HH MM (if applicable)
50-60 61	11 1	BLANK LAB FILE/SAMPLE ID blank	From instrument data system Lab File ID - VOA/BNA Lab Sample ID - Pest
62	1	RECOVERY NO. LABEL	Identifies following counter "P" - # of % Recoveries
63	1	blank	Outside of QC limits
64-65	2	NO. OF RECOVERIES OUT	Use the Counter from Form 2 or 3 for each sample.
66	1	blank	-
67	1	TIC/RPD LABEL	"T" - # of TIC compounds. blank - TIC's not searched "R" - # of % RPD's outside
68	1	blank	limits
69-70	2	NO. TIC/RPD	From Form 1-E or 1-F or Form 3

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SAMPLE CLEANUP RECORD (TYPE 27)

Use: Continuation of Type 20. Used to identify sample clean-up procedures and QC results. Used for Form 9.

Position: Follows Type 20, 21, 22 and 23 to which it applies

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	RECORD TYPE blank	"27"
4 5	1 1	FIRST CLEANUP TYPE blank	"G" for GPC
6-13	8	GPC CALIBRATION CHECK DATE	YY MM DD. Check of GPC calibration to be linked
14	1	blank	with this sample
15-19 20	4 1	GPC CALIBRATION CHECK TIME blank	HHMM
21-31 32 33 34	10 1 1 1	LAB DATA DESCRIPTOR blank SECOND CLEANUP TYPE blank	Lab Sample ID for GPC check "F" for Florisil
35-42	8	FLORISIL LOT CHECK DATE	YY MM DD. Check of Florisil lot check cartridge lots to be linked with this sample
43	1	blank	
44-48	4	FLORISIL LOT CHECK TIME	HHMM
49	1	blank	
50-60	10	LAB DATA DESCRIPTOR	Lab Sample ID for Florisil lot check
61	1	blank	
62	ī	SULFUR CLEANUP	"S" or blank

RESULTS	DATA RECORD	(TYPE 30)	
POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	RECORD TYPE blank	*30*
4	1	ANALYTE LABEL	"C" - CAS Number "I" - Combined DDT and
5	1	blank	Endrin breakdown. "M" - mean conc for Form 10B
6-14	9	CAS NUMBER	Right justified. "COMBINED" - combined DDT
15	1	blank	and Endrin breakdown. "MEANCONC" - mean concentration (Form 10B)
16-24	9	INTERNAL STANDARD CAS NUMBER	Right justified - blank for pesticides
25	1	blank	
26-30	5	CONCENTRATION UNITS	"UG/KG" for Soil; "UG/L " for Water;
31	1	blank	"PERCT" for percent (See Note).
32-34 35	3 1	RESULT QUALIFIER blank	See page H-87. First flag.
36-41	6	RESULT	Right justified; fixed point or scientific
42	1	blank or 'E'	notation
43-45	3 1	<u>EXPONENT</u> blank	Blanks in field will be interpreted as "zeros"
46	1	FIRST VALUE LABEL	Describes following value
47 48	1	blank	(See page H-89)
49-54	6	FIRST RELATED VALUE	Value of item described above. Format
55	1	blank or "E"	same as 36-45.
56-58	3	EXPONENT	
59	1	blank	
60	1	SECOND VALUE LABEL	Describes following value (See page H-89)
61	1	BLANK	
62-66	5 1	SECOND RELATED VALUE blank or 'E'	Value of item described above.
67 68-70	3	EXPONENT	

Note: Units apply to the results field for samples and the related calculated value for calibrations.

0LM01.0

CALCULATION OF RESULTS ON TYPE 30 RECORDS

The following equations are used to calculate all results reported on type 30 records. RELATIVE RESPONSE FACTORS OR CALIBRATION FACTORS

- $\frac{RRF}{(I.S. AREA)} (I.S. AMOUNT)}{(I.S. AREA)} (BNA'S AND VOA'S)$
- CF = (ANALYTE AREA) (PEST's) (ANALYTE AMOUNT [ng injected])

UNKNOWN CONCENTRATIONS

- SAMPLE SIZE _ Sample Volume (in L) for water Sample Weight (in Kg) X (100 - Water) for soil 100
- CONC (ANALYTE AREA) (I.S. injected) (Extract Vol) (D.F. (BNA, VOA) (I.S. AREA) (R.F.) (Injected Vol) (SAMPLE SIZE)
- CONC <u>(ANALYTE AREA)</u> (Extract Vol) (D.F.) (PEST's) (R.F.) (Injected Vol) (SAMPLE SIZE)

CRQL CORRECTION

CRQL = (<u>CROL [ng injected]</u>) (<u>Extract Vol</u>) (D.F.) (Injected Vol) SAMPLE SIZE)

The Sample Volume or Weight is in Type 20 Col. 59-66. The % Water is in Type 22 Col. 36-37. The Extract Vol is in Type 22 Col. The Dilution Factor is in Type 22 Col. 48-54. The Injection Volume is in Type 21 Col. 65-70.

Use: To describe peak areas for internal standards to be used in Form 8 and DFTPP/BFB percent abundances for Form 5.

Position: Follows type 30 for internal standards and DFTPP/BFB data.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	RECORD TYPE	*31*
3	1	blank	
4	1	MASS LABEL	"M" - mass for DFTPP/BFB data
5	1	blank	"I" - internal standard
6	1	VALUE LABEL	A - Area (Internal Standards). P - % Abundance of base peak (DFTPP/BFB);
7-9	3	blank	S - & Abundance of secondary ion.
10-17	8	FIRST MASS	Gives the DFTPP/BFB masses (right justified), e.g., 442.
18	1	blank	
19-28 29	10 1	FIRST VALUE blank	up to 10 decimal digits, right justified (% abundance or area)
30-37	8	SECOND MASS (DFTPP/BFB)	
38	1	blank	
39-48	10	SECOND PERCENT ABUNDANCE	
49	1	blank	
50-57	8	THIRD MASS	
58	1	blank	

59-68 10 THIRD PERCENT ABUNDANCE

Note 1: Columns 30-68 will be blank on records reporting internal standard data.

Note 2: Up to three masses and percent abundances may be reported on each record for DFTPP/BFB data. Those ions that require two percent abundance values must be listed twice.

AUXILLARY DATA RECORD (TYPE 32)

Use: Used to report scan number and retention time (in minutes) for Internal Standards and for TIC compounds. Used to report retention time data and percent breakdown for Pesticides.

Position: Follows type 30. (Record will only be required as specified above.)

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS	
1-2 3-8	2 6	RECORD TYPE blank	*32*	
9-10 11	2 1	RETENTION TIME LABEL blank	"RT"	
12-21 22	10 1	RETENTION TIME blank	In decimal minutes	
23-24	2	FIRST LIMIT LABEL	"SC" - scan number for	
25	1	blank	GC/MS "RF" - RT "From" Value for Pesticides (Form 7D).	
26-35 36	10 1	RT WINDOW LOWER LIMIT blank	In decimal minutes.	
37-38	2	SECOND LIMIT LABEL	"RO" - RT "To" Value	
39	1	blank	for Pesticides (Form 7D)	
40-49 50	10 1	RT WINDOW UPPER LIMIT blank	In decimal minutes.	
51-52	2	THIRD LIMIT LABEL	"PB" indicates	
53	1	blank	<pre>% breakdown for Pesticides *PD* indicates % difference</pre>	
54-63	10	S BREAKDOWN or S DIFFERENCE	in quantitation from two columns (Form 10)	
64	1	blank		
65	1	PEAK	Peak identification (1 through 5)	

.

NAME RECORD (TYPE 33)

Use: To carry an analyte name for TIC compounds.Position:Follows type 30 for TIC compounds.POSITIONLENGTHCONTENTSFORMAT/CONTENTS1-22RECORD TYPE"33"31blank"33"4-7067NAME OF COMPOUND

COMMENT RECORD (TYPE 90)

Use: To provide for Operator-Entered Comments.

Position: May occur anywhere.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	RECORD TYPE blank	*90 *
4-70	67	ANY COMMENT	

FLAGS RECORD (TYPE 91)

Use: To provide for multiple Form 1 Results Qualifier Flags.

Position:	Immediately	follows	the	type	30	record	to	which	ĺt	applies.
-----------	-------------	---------	-----	------	----	--------	----	-------	----	----------

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	RECORD TYPE	"91 "
3	1	blank	
4-6	3	SECOND FLAG	Result Qualifiers
		FLAG	(See page H-87)
7	1	blank	
8-10	3	THIRD FLAG	
11	1	blank	
12-14	3	FOURTH FLAG	
15	1	blank	
16-18	3	FIFTH FLAG	
19	1	blank	
20-22	3	SIXTH FLAG	

6. Definitions of Codes Used in Format B Records

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6.1 Quality Control and Related Codes (QCC) in Type 20 Records

Note: These codes appear in the QC code fields of type 20 records. They are used to indicate the type of data that are being reported.

000	Name	Definition
LRB	LABORATORY (REAGENT) BLANK	The "Method Blank" (See Exhibit G).
LIB	LABORATORY INSTRUMENT BLANK	The "Instrument Blank" (Pesticides)
LSB	LABORATORY SULFUR BLANK	If different from "Method Blank" (Pesticides)
LSD	LABORATORY SPIKE DUPLICATE BACKGROUND (ORIGINAL) VALUES	An environmental sample which is analyzed according to the analytical method, and subsequently used for the matrix spike and the matrix spike duplicate (See Exhibit G).
LF1	LABORATORY SPIKED SAMPLE - FINAL - FIRST MEMBER	The "Matrix Spike" (See Exhibit G) - must preceed LF2
LF2	LABORATORY SPIKED SAMPLE - FINAL - SECOND MEMBER	The "Matrix Spike Duplicate" (See Exhibit G)
LPC	LABORATORY PERFORMANCE CHECK SOLUTION	A solution of DFTPP (BNA) or BFB (VOA) or method analytes (PEST/PCB) used to evaluate the performance of an instrument with respect to a defined set of criteria (Tune or Resolution Check Sample) (See Exhibit G).
FLO	FLORISIL CHECK SOLUTION	A solution of pesticides used to check recovery from each lot of Florisil cartridges.
GPC	GPC CHECK Solution	A solution of pesticides used to check recovery from each new GPC calibration.
CLM	INITIAL CALIBRATION - MULTI POINT	The Initial Calibration for GC/MS (See Exhibit G), or the Initial Evaluation Standard Mixes (A, B) for Pesticides (See Exhibit D PEST). Response factors (GC/MS) or Calibration Factors (Pesticides) rather than concentrations will be reported on the following type 30 records.
CLS	INITIAL CALIBRATION SINGLE POINT	The Initial Individual Standard/Toxaphene/Aroclor Mixes used to determine all calibration factors. (See Exhibit D PEST).

000	Name	Definition
CLE	CONTINUING PERFORMANCE CHECK	The subsequent Individual/Evaluation Standard Mixes (A,B) and Performance Evaluation Mixture for Pesticides (See Exhibit D PEST).
CLC	CONTINUING CHECK CALIBRATION	The Continuing Calibration for GC/MS (See Exhibit G.)
CLD	DUAL PURPOSE CALIBRATION	A calibration solution as above used both as an initial calibration (CLM) and a continuing check (CLC). [50 level initial calibration if needed for Form 8]
blank		Sample, not associated with any quality control item.

The following QCC values are used on type 20 records which act as a header, and indicate that additional (usually calculated) analyte specific data will be present on type 30 (and following type) records. Usually these data will apply to an entire production run, in which case they will appear immediately following the type 10 record. If the data apply to only a portion of the samples in the run, they should be placed immediately preceding the samples to which they apply. Much of the rest of the information in the type 20 record may be blank, indicating that these data do not apply to these results.

MNC	MEAN VALUES FROM CALIBRATIONS	The data following represent mean values and percent RSD's from the initial calibribration (GC/MS) or the evaluation mixes (Pesticides).
SDR	MATRIX SPIKE DUPLICATE CALCULATED RESULTS	The data following represent calculated QC results for the triplicate of samples LSD, LF1, and LF2. Data will consist of the LF1 percent recovery, the LF2 percent recovery, and the percent RSD for each method analyte that was spiked according to the analytical method as is present on hardcopy deliverable Form 3.

6.2 Result Qualifiers in Type 30 Records

A sample qualifier or a result qualifier (also called a non-numeric result) consists of 3 characters which act as an indicator of the fact and the reason that the subject analysis (a) did not produce a numeric result, or (b) produced a numeric result but it is qualified in some respect relating to the type or validity of the result.

BDL	BELOW DETECTABLE LIMITS	Indicates compound was analyzed for but not detected; (Form 1 "U" Flag).
NAR	NO ANALYSIS RESULT	There is no analysis result required for this subject parameter.
AVG	AVERAGE VALUE	Average value - used to report a range of values; e.g., relative response factors.
CBC	CANNOT BE CALCULATED	The analysis result cannot be calculated because an operand value is qualified. Identifies analytes whose Internal Standard is not found.
LTL	LESS THAN LOWER CALIBRATION LIMIT	Actual value is known to be less than the lower calibration range due to dilution. (Form 1 "D" Flag)
GTL	GREATER THAN UPPER CALIBRATION LIMIT	Actual value is known to be greater than the upper calibration range. (Form 1 "E" Flag)
LLS	LESS THAN LOWER STANDARD	The analysis result is less than the sample quantitation limit. (Form 1 "J" Flag)
tie	TENTATIVELY IDENTIFIED -ESTIMATED VALUE	The indicated analyte is a tentatively identified analyte; its concentration has been estimated. (Form 1-E or 1-F "J" Flag)
REJ	REJECTED	Results rejected by the laboratory.
STD	INTERNAL STANDARD	The indicated compound is an internal standard. There is no analysis result to report.
STB	INTERNAL STANDARD BELOW DETECTION LIMITS	A combination of "STD" and "BDL".
FBK	FOUND IN BLANK	The indicated compound was found in the associated method blank (LRB) as well as the sample. (Form 1 "B" flag)
MSP	PERCENT RECOVERY	The following value represents the percent recovery for the "MS" sample. The remaining two values give the "MSD" percent recovery and the Percent RPD.

CON	CONFIRMED	Pesticide identification has been confirmed by GC/MS (Form 1 "C" Flag).
TFB	TENTATIVELY IDENTIFIED AND FOUND IN BLANK	A Combination of "TIE" and "FBK" (Form 1- E or 1-F "B" flag).
ALC	ALDOL CONDENSATION	Labels a suspected Aldol Condensation- product for TIC's (Form 1-E or 1-F "A" Flag).
EST	ESTIMATED VALUE	Present above detection limit but not quantified within expected limits of precision (Form 1-D "P" flag).
PRE	PRESUMPTIVE PRESENCE	Presumptive evidence of presence of material; tentative identification (Form 1-E or 1-F "N" flag).

6.3 Value Labels in Type 30 Records

These codes appear in column 47 (FIRST VALUE LABEL) of Type 30 records to identify the value in columns 49-58.

<u>Oualifier</u>	Full Name	Definition
A	AMOUNT ADDED	Identifies the amount of matrix spike analyte added (for QC codes "LF1" and "LF2") or nominal amount for pesticide standards.
P	PERCENT RECOVERY	Identifies the Percent Recovery of the "MSD" Sample in the Matrix Spike Results Record (for QC Code "SDR")

These codes appear in column 60 (SECOND VALUE LABEL) of Type 30 records to identify the value in columns 62-70.

<u>Oualifier</u>	Full Name	Definition
U	UNDETECTED	Value is the corrected sample quantitation limit (Form 1 "U" Value).
R	PERCENT RSD	Value is the Percent RPD for the Matrix Spike and Matrix Spike Duplicate (QC Code "SDR") or RSD for the Mean Response Factors (QC Code "MNC").
S	SURROGATE RECOVERY	Value is the Percent Recovery for the indicated Surrogate or Spike (Form 2, Form 9A, Form 9B).
D	PERCENT DIFFERENCE	Value is the Percent Difference of the Result of the Continuing Check from that of the Initial Calibration (as on Form 7).
P	PESTICIDE LEVEL	Value is the concentration of pesticide standard (LOW, MED, HIGH) as a factor of the LOW (e.g., LOW-1, MED-10)

7. Format of Records for Specific Uses

7.1 Format of the SAMPLE HEADER DATA RECORD (Type 20) for Mean Relative Response Factors

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	"20"	Record Type
3-15	13	blank	
16	1	"O"	All matrices
17	1	blank	
18-20	3	"MNC"	Identifies Mean Response
21-66	46	blank	Factors
67-69	3	ANALYTE COUNT	Numeric; 1-3 decimal digits; right justified.

7.2 Format of the RESULTS DATA RECORD (Type 30) for Mean Relative Response Factors

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	"30"	Record Type
3	1	bl ank	
4	1	"C"	
5	1	blank	
6-14	9	CAS NUMBER	Right justified.
15	1	blank	
16-24	9	INTERNAL STANDARD CAS NUMBER	Right justified
25-31	7	blank	
32-34	3	"AVG"	Indicates Average Value
35	1	blank	
36-41	6	RESULT	Mean Relative Response Factor Right justified; fixed
42	1	blank or 'E'	or scientific notation
43-45	3	EXPONENT	Blank field will be interpreted as "+00"
46-59	14	blank	
60	1	"R"	Indicates Percent RSD
61	1	blank	
62-66	5	SECOND RELATED VALUE	Percent RSD
67	1	blank or 'E'	
68-70	3	EXPONENT	

7.3 Format of the SAMPLE HEADER DATA RECORD (Type 20) for Matrix Spike Duplicates

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	*20*	Record Type
3-6	4	blank	
7-11	5	EPA SAMPLE NO.	Left justified. Raw
12-15	4	blank	Sample I.D. only; no suffixes.
16	1 1	MATRIX	"l" for water
17	1	blank	"H" for soil
18-20	3 5	"SDR"	Identifies Matrix Spike
21-25	5	blank	Duplicate Results
26-30	5	CASE NUMBER	
31-33	3	blank	
34-38	5	SDG NO.	
39-66	2	blank	
67-69	3	ANALYTE COUNT	Numeric; 1-3 decimal digits; right justified. Counts number of spiked

7.4 Format of the COUNTER RECORD (Type 23) for Matrix Spike Duplicates

Position: Follows the type 20 to which it applies.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3-61	2 59	"23" bl ank	Record Type
62 63	1 1	"P" blank	Identifies Number of % Recoveries outside of limits.
64-65	2	NO. RECOVERIES OUT	Use the counter from
66	1	blank	Form 3 for each sample.
67	1	"R"	# of % RPD's outside
68	1	blank	limits.
69-70	2	NO. RPD LIMITS	From Form 3.

analytes.

7.5 Format of the RESULTS DATA RECORD (Type 30) for Matrix Spike Duplicates

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	*30*	Record type
3	1	blank	
4	1	*C*	
5	1	blank	
6-14	9	CAS NUMBER	Right justified.
15	1	blank	
16-24	9	INTERNAL STANDARD CAS NUMBER	Right justified
25	1	blank	
26-30	5	*PERCT*	Units are "Percent"
31	1	blank	
32-34	3	"MSP"	Indicates Matrix Spike
32-3 4 35	1	blank	Percent recovery
36-41	6	RESULT	MS Percent Recovery Right justified; fixed
42	1	blank or 'E'	or scientific notation
43-45	3	EXPONENT	Blank field is
43-45	1	blank	interpreted as "+00"
47	1	"P"	Indicates Matrix Spike
47 48	1	blank	Duplicate Percent Recovery
49-54	6	FIRST RELATED VALUE	MSD Percent Recovery Format same as 36-46.
55	1	blank or 'E'	
56-58	3	EXPONENT	
59	1	blank	
60	1	"R"	Indicates Percent RSD
61	1	blank	
62-66	5	SECOND RELATED VALUE	Percent RPD
67	1	blank or 'E'	
68-70	3	EXPONENT	

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7.6 Format of the SAMPLE HEADER DATA RECORD (Type 20) for Pesticide Evaluation Mix B

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3-6	2 4	"20" blank	Record Type
7-11 12-15	5 4	"EVALB" blank	Sample I.D.
16 17	1 1	"0" blank	All matrices
18-20	3	"CLC"	Indicates Continuing Check (Pesticide Standard)
21-25	5	blank	
26-30	5	CASE NUMBER	
31-38	8	blank	
39-46	8	DATE ANALYZED	YY MM DD
47	1	blank	
48-52	5	TIME ANALYZED	HH MM
53-66	14	blank	
67 - 69	3	ANALYTE COUNT	Numeric; 1-3 decimal digits; right justified.

7.7 Format of the RESULTS DATA RECORD (Type 30) for Pesticide Evaluation Mix B (Percent Breakdown Data From Form 8D)

The following type 30 record occurs once if reporting "Combined" breakdown only, or three times if reporting separate breakdowns for Endrin and DDT along with the Combined value.

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	"30" blank	Record Type
4	1	"C" or "I"	Use "C" - CAS Number unless identifying combined DDT and Endrin, in which case
5	1	blank	use "I"
6-14	9	CAS NUMBER	Right justified. Use "COMBINED" for combined
15-25	11	blank	DDT and Endrin.
26-30 31-46	5 16	"PERCT" blank	Units
47 48	1 1	"B" blank	Identifies Percent Breakdown
49 - 54	6	FIRST RELATED VALUE	Percent Breakdown of Indicated Compound(s) Right justified; fixed scientific notation
55 56-58	1 3	blank or 'E' Exponent	Blank field will be interpreted as "+00"

7.8 Format of the SAMPLE HEADER DATA RECORDS (Type 20-23) and RESULTS DATA RECORD (Type 30) for Continuing Checks (GC/MS Methods)

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	*20*	
3-6	4	blank	
	_		
7-13	7	Identifier	e.g., VTD050
14	1	Daily Sequence Number	From Exhibit B
15	1	blank	
16	1	"O"	All matrices
17	1	blank	
	-		
18-20	3	"CLC"	Indicates
21	1	blank	Continuing Check
22-24	3	SAMPLE QUALIFIER	See page H-85
25	1	blank	see hafe u-os
26-30	5	CASE NUMBER	
31-38	8	blank	
39-46	8	DATE ANALYZED	
39-40 47	8 1	blank	YY MM DD
47	Ŧ	UTAUK	
48-52	5	TIME ANALYZED	HH MM
53-66	14	blank	
	-		
67-69	3	ANALYTE COUNT	Numeric; 1-3
			decimal digits;
			right justified.
POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2	2	*21*	Record Type
3-13	11	blank	••
1/ 16	3	"H"	
14-16	3	n n	Indicates heated purge
17	1	blank	
an f	-		
18-23	6	SAS NUMBER	Leave blank
24	1	blank	if none.
25-35	11	LAB FILE I.D.	

POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1-2 3	2 1	"23" blank	Record Type
4 5	1 1	"P" blank	Labels data as "tune" data.
6-13	8	INJECTION DATE	YY MM DD. Aquisition date
14	1	blank	of tune to be linked with this calibration.
15-19 . 20	5 1	INJECTION TIME blank	HH MM
21-31	11	DFTPP/BFB LAB FILE ID	From instrument data system.
POSITION	LENGTH	CONTENTS	FORMAT/CONTENTS
1 0	2	*30*	
1-2 3	1	blank	
4	1	"C"	
5	1	blank	
6-14	9	CAS NUMBER	Right justified.
15	1	blank	
16-24	9	INTERNAL Standard Cas Number	Right justified.
25-31	7	blank	
32-34 35	3 1	RESULT QUALIFIER blank	See page H-87;
36-41	6	RESULT	Response Factor Right justified;
42	1	blank or 'E'	fixed or scientific notation
43-45	3	EXPONENT	Blank field will be
46-59	14	blank	interpreted as "+00".
60	1	"D"	Identifies Percent
61	ī	blank	Difference.
62-66	5	SECOND RELATED VALUE	RRF Percent Difference From Initial
67	1	blank or 'E'	Calibration
68-70	3	EXPONENT	(from Form 7).