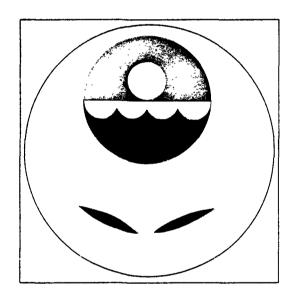
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METHOD OF STANDARD ADDITIONS
AND EFFECTS OF DILUTION

May 1980

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Method of Standard Additions and Effects of Dilution

May 1980

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Region III

U.S. Environmental Protection Agency

The method of standard additions is useful when the matrices of standards and samples differ significantly. The method of additions may involve additions of small quantities of the analyte to the sample with no significant dilution of the sample or cases which involve significant sample dilution. In general, a plot is made of absorbance versus added concentration and the value of the X-intercept is taken as the analyte concentration.

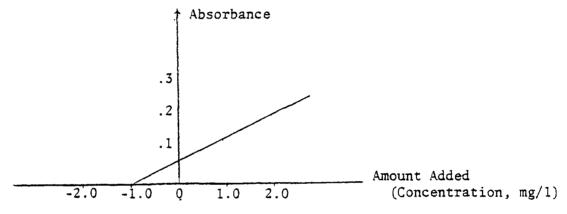
The method of standard addition eliminates interferences that cause constant multiplicative errors in the concentration of analyte measured. As an example of this type of interference, consider a sample with a true analyte concentration of 1.0 ppm. The analyst assays the sample and determines a concentration of 0.5 mg/l indicating an interference factor of .5. (This is a negative interference since the interference factor is less than 1.) The sample is spiked with small quantities of the analyte to produce an added concentration of 1.0 ppm and 2.0 ppm and values of 1 ppm and 1.5 ppm, respectively, are obtained. From these results the analyst decides there is an interference and plots the data (Figure 1).

Figure 1. Data and Plot of Method of Standard Additions

Concentration of analyte	*Measured	**Measured
added after dilution	Concentration	Absorbance
by sample		
(mg/1)	(mg/1)	•
0	.5	.05
1	1.0	.10
2	1.5	.15

*This represents an interference factor of .5.

^{**}The standard curve had a slope of .1 and a y-intercept of zero.

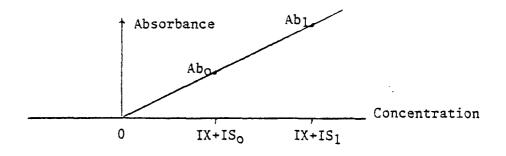


Scale of negative concentration values and positive concentration values must be the same.

The extrapolation of the line to the X-intercept generates the actual concentration of the sample (ignoring the negative sign).

The addition of small known quantities of the analyte or "spiking" to determine sample concentration is based upon the Beer's Law relationship: the absorbance is directly proportional to the concentration of the analyte present (Figure 2).

Figure 2. Plot of Absorbance vs Concentration with Application of Beer's Law



Beer's Law:

$$\frac{Ab_0}{IX + IS_0} = \frac{Ab_1}{IX + IS_1}$$
or
$$X = \frac{(Ab_0)(S_1)}{Ab_1 - Ab_0} \frac{I}{I}$$
or
$$X = \frac{Ab_0S_1}{Ab_1 - Ab_0} = \frac{I}{I}$$

where: Ab_O = Absorbance of solution with concentration = IX + IS_O

Ab₁ = Absorbance of solution with concentration = IX + IS₁

I = Interference factor (constant
 multiplicative error)

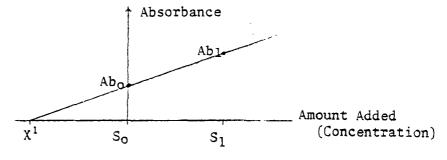
 $S_0 = 0$ spike addition = 0

S₁ = Concentration of analyte added
 after dilution by the total
 volume (sample volume + spike
 volume)

X = Concentration of analyte in sample

The sample concentration may be determined by solving for χ in the proportion (Figure 2). The method of standard additions plot may be generalized (Figure 3):

Figure 3. Generalized Plot of Method of Standard Additions



y = mX + bat y = 0 $X^{1} = -b/m$ $b = Ab_{0}$ $m = (Ab_{1} - Ab_{0}) / S_{1}$ $X^{1} = -Ab_{0}S_{1} / (Ab_{1} - Ab_{0})$ Ab_O = Absorbance of solution with zero analyte addition

Ab₁ = Absorbance of solution associated with addition of spike of S₁

 $S_0 = 0$ spike addition

S₁ = Concentration of analyte added after dilution by the total volume (sample volume + spike volume)

X¹ = Method of standard addition
 determined concentration of
 analyte in sample

3.

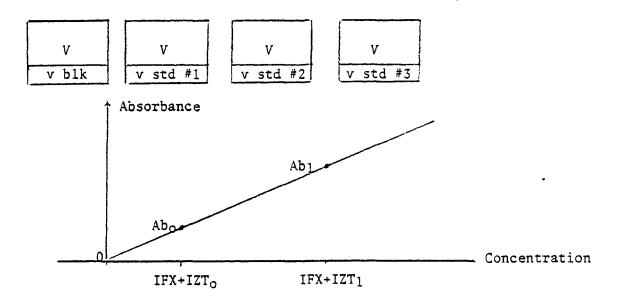
Again, a linear relationship based upon Beer's Law is assumed. A linear relation is defined: y = mX+b; where y is the absorbance, X is the concentration, m is the slope, and b is the y-intercept. At the X-intercept, y = 0 and $X^1 = -b/m$, or $X^1 = -Ab_0S_1/Ab_1-Ab_0$. The extrapolation generates the same numerical value (but negative in sign) as the true result and is therefore applicable to additions of small volumes of analyte. The interference free (true) result was obtained from the Beer's Law relation because of the constant nature of the interference.

A problem arises in using the extrapolation method when significant and varying dilutions of the sample by the addition of the spike are involved. The spiked samples are prepared by adding a significant volume of analyte and diluting to a pre-set volume with the sample (i.e., 1. establish a pre-set volume, for example, 100 ml; 2. for the first spiked sample, add 2 ml of analyte of a known concentration and add 98 ml of sample to bring the total volume to 100 ml; 3. for the second spiked sample, add 4 ml of analyte of the same known concentration and add 96 ml of sample to bring the total volume to 100 ml; etc.). The results obtained from this method of standard additions are incorrect because the concentration of the interferent varies in each spiked sample aliquot. In order to accurately determine the analyte concentration, the concentration of the interferent must be kept constant. This can be accomplished by following the EPA method of standard additions.

The results of the EPA method of standard additions outlined in the 1974 and 1979 editions of the EPA <u>Manual of Methods of Chemical Analysis</u> of <u>Water and Wastes*</u> may be effected by dilution. The EPA method requires that a constant volume of sample be added to a constant volume of standards and blank. If the volume of sample consistently used is V and the volume of standards and blank is consistently v where $V \neq V$, this experiment may be generalized to that presented in Figures 4(1) and 4(2).

^{*}Metal's section page 12, 1979 EPA Manual of Methods of Chemical Analysis of Water and Wastes

Figure 4(1). Correct result obtained using Beer's Law coupled with the Dilution and Interference Factors and Using the Procedure outlined by EPA



$$\frac{Ab_0}{IFX + IZT_0} = \frac{Ab_1}{IFX + IZT_1}$$

or
$$X = \frac{Z}{F} \frac{-Ab_0T_1}{Ab_0-Ab_1}$$

but; Z/F = v/V

therefore;
$$X = \frac{V}{V} = \frac{Ab_0T_1}{Ab_1-Ab_0}$$

where: V = Volume of sample

v = Volume of standard or blank

F = V/(V+v)

Z = v/(V+v)

I = Any interference factor

 T_0 = Concentration of the analyte in the blank = 0

 T_1 = Concentration of the analyte in standard #1 (S_1 = ZT_1)

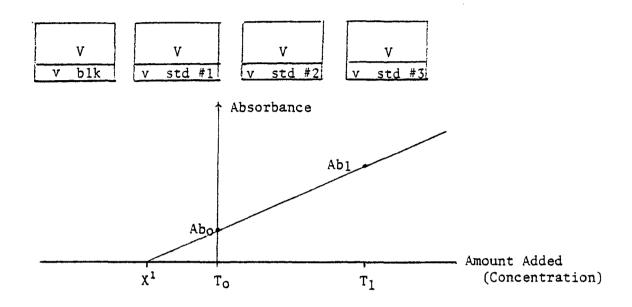
Abo = Absorbance associated with the concentration:

IFX+IZT_O
Ab₁ = Absorbance associated with

the concentration: IFX+IZT₁

X = Sample concentration

Figure 4(2). Incorrect result obtained using the Method of Standard Additions as outlined by EPA, where $V \neq v$, (an example is included in the appendix).



 $m = \Delta y/\Delta X = (Ab_1 - Ab_0)T_1 - T_0$

where:

 $b = y-intercept = Ab_0$ at y=0, $X^1=-b/m$ therefore; $X^1 = (-Ab_0)$

 $=(Ab_1-Ab_0)T_1$

 $X^1 = -Ab_0T_1$ Ab1-Abo

 $y = mX^1 + b$

V = Volume of sample

v = Volume of standard or blank T_0 = Concentration of the analyte

in the blank = 0

 T_1 = Concentration of the analyte in standard #1

Abo = Absorbance associated with zero analyte addition

 Ab_1 = Absorbance associated with addition of a concentration of T_1

 X^1 = Method of standard addition determined concentration of analyte in sample

These results indicate that if $v \neq V$, this method of standard additions does not yield accurate results. The results of extrapolation may be corrected by a factor, -(v/V):

$$X = (X^{1})(v/V)(-1)$$
True

If the "Amount Added" is interpreted as the concentration of standard corrected for dilution by the sample, the true result remains:

$$X = Z \quad Ab_0T_1$$
True F Ab_1-Ab_0

the extrapolation becomes:

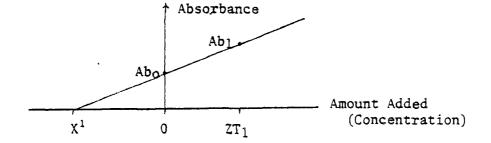
y = mX + b

 $b = Ab_{O}$ at y=0, $X^{1}=-b/m$ $X^{1}=(\frac{-Ab_{O}}{Ab_{1}-Ab_{O}}$

 $m = \Delta y/\Delta X = (Ab_1 - Ab_0)/ZT_1$

$$X^1 = \frac{-Ab_0 ZT_1}{Ab_1 - Ab_0}$$
 (See Figure 5)

Figure 5. Incorrect result obtained using the Method of Standard Additions when Dilution of the Sample by the Spike is Disregarded $(V \neq v)$



where: V

V = Volume of sample

v = Volume of standard or blk

F = V/(V+v) Z = v/(V+v)

 T_1 = Concentration of the analyte in standard #1 (S_1 = ZT_1)

Abo = Absorbance associated with zero standard addition

Ab₁ = Absorbance associated with addition of a concentration of ZT₁

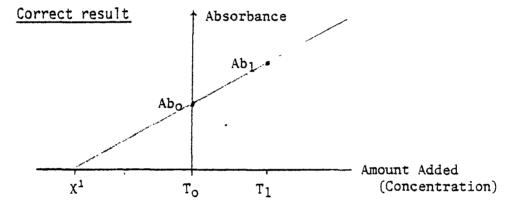
X¹ = Method of standard addition determined concentration of analyte in sample The result of the extrapolation, $X^1 = \frac{(-Ab_0)(ZT_1)}{Ab_1 - Ab_0}$, may be corrected by a factor: $X = (X^1)(\frac{-1}{F})$, where F = V/(v+V).

The EPA method of standard additions yields accurate results when v = V. In this case $F = \frac{V}{V+V} = \frac{v}{V+V}$ and Beer's Law with correction for dilution yields the result:

$$\frac{Ab_0}{IFX} = \frac{Ab_1}{IFX + IFT_1}$$
 or $X = \frac{Ab_0T_1}{Ab_1 - Ab_0}$

which is the same result obtained from a plot of absorbance vs the concentration of the analyte in the added standard (See Figure 6).

Figure 6. Plot of Absorbance vs Concentration Added when V=V



where:

$$y = mX+b$$

at $y = 0$, $X^1 = -b/m$
 $b = Ab_0$
 $m = \Delta y/\Delta X = (Ab_1 - Ab_0)/T_1$
 $X^1 = (-Ab_0)(T_1) / (Ab_1 - Ab_0)$
 $X = (X^1)(v/V)(-1)$ and $v = V$;
True
 $X = (X^1)(-1)$
True

V = Volume of sample

v = Volume of standard or blank
T1 = Concentration of the analyte

in standard #1

Abo = Absorbance associated with zero standard addition

Ab₁ = Absorbance associated with addition of a concentration of T₁

X = True concentration of analyte in sample

X¹ = Method of standard addition determined concentration of analyte in sample The EPA method of standard additions does not account for the dilution of sample by the standard except when:

- 1. The volume of sample used is the same as the volume of standard, V = V.
- 2. Zero analyte was initially present, $Ab_0 = 0$.
- 3. No dilution of sample by standard occurs (insignificant volumes, e.g. microliter of standard and blank are used and diluted to a constant volume with sample).

If the volumes of sample and standard are significant and unequal, the extrapolation will only yield accurate results if a correction factor is employed (an example is included in the Appendix).

In conclusion, the method of standard addition does not necessarily correct for dilutions resulting from the addition of the analyte.

Appendix

A 1.0 mg/l phenol reference sample was analyzed using the 4-amino antipyrine method. The results obtained from a standard calibration curve and from the method of standard additions follow.

Standard Calibration Curve

Concentration (mg/1)	Absorbance	
0	.000	
.2	.029	
.5	.070	
.7	.098	correlation coefficient = .999
1.0	.142	slope = .140 y-intercept =000346
2.0	.272	
3.0	.421	
4.0	.562	

The reference sample had an absorbance of 0.142 which corresponds to 1.0 mg/l based upon the standard calibration curve.

Standard Method of Additions

Twenty milliliters of sample were placed into each of four flasks. To flask #1, 80 ml of deionized water were added. To flask #2, 80 ml of a 0.5 mg/l phenol solution were added. To flask #3, 80 ml of a 1.0 mg/l phenol solution were added. To flask #4, 80 ml of a 2.0 mg/l phenol solution were added. The concentrations added and absorbances obtained are tabulated in Table I.

Table I

Flask	No. Concentration (mg/1)	Added Absorbanc	ce
1	0	.027	
2	.5		<pre>correlation coefficient = .9999 slope = .112 y-intercept = .027</pre>
3	1.0		at $y = 0$, $x =240$
4	2.0	.252	

The result obtained from a plot of the data in Table I is 0.240 mg phenol/1. This value is incorrect due to the dilution of the sample by the spike. The actual concentration may be obtained by multiplying the value of the x-intercept by v/V (see Equation 1), where v is the spike volume and V is the sample volume.

Actual Concentration = (Value of the x-intercept) (-1)(v/V) Equation 1 In this example, the actual concentration = (-.240)(-1)(80/20) = 0.96 mg/l.

Alternatively, the concentrations of the standards added may be corrected for dilution by the sample. In Table II, the added concentrations corrected for dilution by the sample and corresponding absorbances are listed. (Tables I and II are identical except that the concentrations added have been corrected for dilution by the sample in Table II).

Table II

Flask No.	Added Conc. Corrected for Dilution (mg/l)	Absorbance	
1	0	.027	
2	. 4	correlation coefficient = .084 slope = .140 y-intercept = .027	.999
3	.8	.138 at $y = 0, x =192$	
4	1.6	. 252	

The result obtained from a plot of the data in Table II is 0.192 mg/l. Again an incorrect result is generated. The actual value may be obtained by multiplying the x-intercept by (v+V)/v, where v is the spike volume and V is the sample volume (see Equation 2).

Actual Concentration = (Value of the x-intercept) (-1)(v+V)/v Equation 2

In this example, the actual concentration = (-.192)(-1)(100/20) = 0.96 mg/1.

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16. ABSTRACT

The basis and limitation of the Method of Standard Addition is discussed with particular emphasis placed on the effects of dilution often encountered in using this analytical tool. Dilutions employed when using Standard Additions may effect the sample matrix. This problem may be avoided by maintaining a constant sample dilution. If significant dilution of the sample is involved, a dilution factor must be applied to obtain the correct result by the Method of Standard Additions.

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