Method 301--Field Validation of Pollutant Measurement Methods from Various Waste Media

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Using Method 301

1.0 What is the purpose of Method 301?

Method 301 provides a set of procedures for the owner or operator of an affected source to validate a candidate test method as an alternative to a required test method based on established precision and bias criteria. These validation procedures are applicable under 40 CFR part 63 or 65 when a test method is proposed as an alternative test method to meet an applicable requirement or in the absence of a validated method. Additionally, the validation procedures of Method 301 are appropriate for demonstration of the suitability of alternative test methods under 40 CFR parts 59, 60, and 61. If, under 40 CFR part 63 or 60, you choose to propose a validation method other than Method 301, you must submit and obtain the Administrator’s approval for the candidate validation method.

2.0 What approval must I have to use Method 301?

If you want to use a candidate test method to meet requirements in a subpart of 40 CFR part 59, 60, 61, 63, or 65, you must also request approval to use the candidate test method according to the procedures in Section 16 of this method and the appropriate section of the part (§59.104, §59.406, §60.8(b), §61.13(h)(1)(ii), §63.7(f), or §65.158(a)(2)(iii)). You must receive the Administrator’s written approval to use the candidate test method before you use the candidate test method to meet the applicable federal requirements. In some cases, the Administrator may decide to waive the requirement to use Method 301 for a candidate test method to be used to meet a requirement under 40 CFR part 59, 60, 61, 63, or 65 in absence of a validated test method. Section 17 of this method describes the requirements for obtaining a waiver.

3.0 What does Method 301 include?

3.1 Procedures. Method 301 includes minimum procedures to determine and document systematic error (bias) and random error (precision) of measured concentrations from exhaust gases, wastewater, sludge, and other media. Bias is established by comparing the results of sampling and analysis against a reference value. Bias may be adjusted on a source-specific basis using a correction factor and data obtained during the validation test. Precision may be determined using a paired sampling system or quadruplicate sampling system for isotopic spiking. A quadruplicate sampling system is required when establishing precision for analyte spiking or when comparing a candidate test method to a validated method. If such procedures have not been established and verified for the candidate test method, Method 301 contains procedures for ensuring sample stability by developing sample storage procedures and limitations and then testing them. Method 301 also includes procedures for ruggedness testing and determining detection limits. The procedures for ruggedness testing and determining detection limits are required for candidate test methods that are to be applied to multiple sources and optional for candidate test methods that are to be applied at a single source.
3.2 Definitions.

Affected source means an affected source as defined in the relevant part and subpart under Title 40 (e.g., 40 CFR parts 59, 60, 61, 63, and 65).

Candidate test method means the sampling and analytical methodology selected for field validation using the procedures described in Method 301. The candidate test method may be an alternative test method under 40 CFR part 59, 60, 61, 63, or 65.

Paired sampling system means a sampling system capable of obtaining two replicate samples that are collected as closely as possible in sampling time and sampling location (collocated).

Quadruplicate sampling system means a sampling system capable of obtaining four replicate samples (e.g., two pairs of measured data, one pair from each method when comparing a candidate test method against a validated test method, or analyte spiking with two spiked and two unspiked samples) that are collected as close as possible in sampling time and sampling location.

Surrogate compound means a compound that serves as a model for the target compound(s) being measured (i.e., similar chemical structure, properties, behavior). The surrogate compound can be distinguished by the candidate test method from the compounds being analyzed.

4.0 How do I perform Method 301?

First, you use a known concentration of an analyte or compare the candidate test method against a validated test method to determine the bias of the candidate test method. Then, you collect multiple, collocated simultaneous samples to determine the precision of the candidate test method. Additional procedures, including validation testing over a broad range of concentrations over an extended time period are used to expand the applicability of a candidate test method to multiple sources. Sections 5.0 through 17.0 of this method describe the procedures in detail.

Reference Materials

5.0 What reference materials must I use?

You must use reference materials (a material or substance with one or more properties that are sufficiently homogenous to the analyte) that are traceable to a national standards body (e.g., National Institute of Standards and Technology (NIST)) at the level of the applicable emission limitation or standard that the subpart in 40 CFR part 59, 60, 61, 63, or 65 requires. If you want to expand the applicable range of the candidate test method, you must conduct additional test runs using analyte concentrations higher and lower than the applicable emission limitation or the anticipated level of the target analyte. You must obtain information about your analyte according to the procedures in Sections 5.1 through 5.4 of this method.

5.1 Exhaust Gas Test Concentration. You must obtain a known concentration of each analyte from an independent source such as a specialty gas manufacturer, specialty chemical company, or chemical laboratory. You must also obtain the manufacturer’s certification of traceability, uncertainty, and stability for the analyte concentration.

5.2 Tests for Other Waste Media. You must obtain the pure liquid components of each analyte from an independent manufacturer. The manufacturer must certify the purity, traceability, uncertainty, and shelf life of the pure liquid components. You must dilute the pure liquid components in the same type medium or matrix as the waste from the affected source.

5.3 Surrogate Analytes. If you demonstrate to the Administrator’s satisfaction that a surrogate compound behaves as the analyte does, then you may use surrogate compounds for highly toxic or
reactive compounds. A surrogate may be an isotope or compound that contains a unique element (e.g., chlorine) that is not present in the source or a derivation of the toxic or reactive compound if the derivative formation is part of the method’s procedure. You may use laboratory experiments or literature data to show behavioral acceptability.

5.4 Isotopically-Labeled Materials. Isotope mixtures may contain the isotope and the natural analyte. The concentration of the isotopically-labeled analyte must be more than five times the concentration of the naturally-occurring analyte.

Sampling Procedures

6.0 What sampling procedures must I use?

You must determine bias and precision by comparison against a validated test method using isotopic spiking or using analyte spiking (or the equivalent). Isotopic spiking can only be used with candidate test methods capable of measuring multiple isotopes simultaneously such as test methods using mass spectrometry or radiological procedures. You must collect samples according to the requirements specified in Table 301-1 of this method. You must perform the sampling according to the procedures in Sections 6.1 through 6.4 of this method.

6.1 Isotopic Spiking. Spike all 12 samples with isotopically-labelled analyte at an analyte mass or concentration level equivalent to the emission limitation or standard specified in the applicable regulation. If there is no applicable emission limitation or standard, spike the analyte at the expected level of the samples. Follow the applicable spiking procedures in Section 6.3 of this method.

6.2 Analyte Spiking. In each quadruplicate set, spike half of the samples (two out of the four samples) with the analyte according to the applicable procedure in Section 6.3 of this method. You should spike at an analyte mass or concentration level equivalent to the emission limitation or standard specified in the applicable regulation. If there is no applicable emission limitation or standard, spike the analyte at the expected level of the samples. Follow the applicable spiking procedures in Section 6.3 of this method.

6.3 Spiking Procedure.

6.3.1 Gaseous Analyte with Sorbent or Impinger Sampling Train. Sample the analyte being spiked (in the laboratory or preferably in the field) at a mass or concentration that is approximately equivalent to the applicable emission limitation or standard (or the expected sample concentration or mass where there is no standard) for the time required by the candidate test method, and then sample the stack gas stream for an equal amount of time. The time for sampling both the analyte and stack gas stream should be equal; however, you must adjust the sampling time to avoid sorbent breakthrough. You may sample the stack gas and the gaseous analyte at the same time. You must introduce the analyte as close to the tip of the sampling probe as possible.

6.3.2 Gaseous Analyte with Sample Container (Bag or Canister). Spike the sample containers after completion of each test run with an analyte mass or concentration to yield a concentration approximately equivalent to the applicable emission limitation or standard (or the expected sample concentration or mass where there is no standard). Thus, the final concentration of the analyte in the sample container would be approximately equal to the analyte concentration in the stack gas plus the equivalent of the applicable emission standard (corrected for spike volume). The volume amount of spiked gas must be less than 10 percent of the sample volume of the container.
6.3.3 Liquid or Solid Analyte with Sorbent or Impinger Trains. Spike the sampling trains with an amount approximately equivalent to the mass or concentration in the applicable emission limitation or standard (or the expected sample concentration or mass where there is no standard) before sampling the stack gas. If possible, do the spiking in the field. If it is not possible to do the spiking in the field, you must spike the sampling trains in the laboratory.

6.3.4 Liquid and Solid Analyte with Sample Container (Bag or Canister). Spike the containers at the completion of each test run with an analyte mass or concentration approximately equivalent to the applicable emission limitation or standard in the subpart (or the expected sample concentration or mass where there is no standard).

6.4 Probe Placement and Arrangement for Stationary Source Stack or Duct Sampling. To sample a stationary source, you must place the paired or quadruplicate probes according to the procedures in this subsection. You must place the probe tips in the same horizontal plane. Section 17.1 of Method 301 describes conditions for waivers. For example, the Administrator may approve a validation request where other paired arrangements for the probe tips or pitot tubes (where required) are used.

6.4.1 Paired Sampling Probes. For paired sampling probes, the first probe tip should be 2.5 centimeters (cm) from the outside edge of the second probe tip, with a pitot tube on the outside of each probe.

6.4.2 Quadruplicate Sampling Probes. For quadruplicate sampling probes, the tips should be in a 6.0 cm x 6.0 cm square area measured from the center line of the opening of the probe tip with a single pitot tube, where required, in the center of the probe tips or two pitot tubes, where required, with their location on either side of the probe tip configuration. Section 17.1 of Method 301 describes conditions for waivers. For example, you must propose an alternative arrangement whenever the cross-sectional area of the probe tip configuration is approximately five percent or more of the stack or duct cross-sectional area.

7.0 How do I ensure sample stability?

7.1 Developing Sample Storage and Threshold Procedures. If the candidate test method includes well-established procedures supported by experimental data for sample storage and the time within which the collected samples must be analyzed, you must store the samples according to the procedures in the candidate test method and you are not required to conduct the procedures specified in Section 7.2 or 7.3 of this method. If the candidate test method does not include such procedures, your candidate method must include procedures for storing and analyzing samples to ensure sample stability. At a minimum, your proposed procedures must meet the requirements in Section 7.2 or 7.3 of this method. The minimum duration between sample collection and storage must be as soon as possible, but no longer than 72 hours after collection of the sample. The maximum storage duration must not be longer than 2 weeks.

7.2 Storage and Sampling Procedures for Stack Test Emissions. You must store and analyze samples of stack test emissions according to Table 301-2 of this method. You may reanalyze the same sample at both the minimum and maximum storage durations for: (1) samples collected in containers such as bags or canisters that are not subject to dilution or other preparation steps, or (2) impinger samples not subjected to preparation steps that would affect stability of the sample such as extraction or digestion. For candidate test method samples that do not meet either of these criteria, you must analyze one of a pair of replicate samples at the minimum storage duration and the other replicate at the proposed storage duration but no later than 2 weeks of the initial analysis to identify the effect of storage duration on analyte samples. If you are using the isotopic spiking procedure, then you must analyze each sample for the spiked analyte and the native analyte.
7.3 Storage and Sampling Procedures for Testing Other Waste Media (e.g., Soil/Sediment, Solid Waste, Water/Liquid). You must analyze one of each pair of replicate samples (half the total samples) at the minimum storage duration and the other replicate (other half of samples) at the maximum storage duration or within 2 weeks of the initial analysis to identify the effect of storage duration on analyte samples. The minimum time period between collection and storage should be as soon as possible, but no longer than 72 hours after collection of the sample.

7.4 Sample Stability. After you have conducted sampling and analysis according to Section 7.2 or 7.3 of this method, compare the results at the minimum and maximum storage durations. Calculate the difference in the results using Equation 301-1.

\[ d_i = R_{\text{mini}} - R_{\text{maxi}} \]  

(Eq. 301-1)

Where:
- \( d_i \) = Difference between the results of the \( i^{\text{th}} \) replicate pair of samples.
- \( R_{\text{mini}} \) = Results from the \( i^{\text{th}} \) replicate sample pair at the minimum storage duration.
- \( R_{\text{maxi}} \) = Results from the \( i^{\text{th}} \) replicate sample pair at the maximum storage duration.

For single samples that can be reanalyzed for sample stability assessment (e.g., bag or canister samples and impinger samples that do not require digestion or extraction), the values for \( R_{\text{mini}} \) and \( R_{\text{maxi}} \) will be obtained from the same sample rather than replicate samples.

7.4.1 Standard Deviation. Determine the standard deviation of the paired samples using Equation 301-2.

\[ SD_d = \sqrt{\frac{\sum_{i} (d_i - d_m)^2}{n - 1}} \]  

(Eq. 301-2)

Where:
- \( SD_d \) = Standard deviation of the differences of the paired samples.
- \( d_i \) = Difference between the results of the \( i^{\text{th}} \) replicate pair of samples.
- \( d_m \) = Mean of the paired sample differences.
- \( n \) = Total number of paired samples.

7.4.2 T Test. Test the difference in the results for statistical significance by calculating the t-statistic and determining if the mean of the differences between the results at the minimum storage duration and the results after the maximum storage duration is significant at the 95 percent confidence level and \( n-1 \) degrees of freedom. Calculate the value of the t-statistic using Equation 301-3.
\[ t = \frac{|d_m|}{\left( \frac{SD_d}{\sqrt{n}} \right)} \]  
(Eq. 301-3)

Where:

\[ t \quad = \quad \text{t-statistic.} \]
\[ d_m \quad = \quad \text{The mean of the paired sample differences.} \]
\[ SD_d \quad = \quad \text{Standard deviation of the differences of the paired samples.} \]
\[ n \quad = \quad \text{Total number of paired samples.} \]

Compare the calculated t-statistic with the critical value of the t-statistic from Table 301-3 of this method. If the calculated t-value is less than the critical value, the difference is not statistically significant. Therefore, the sampling, analysis, and sample storage procedures ensure stability, and you may submit a request for validation of the candidate test method. If the calculated t-value is greater than the critical value, the difference is statistically significant, and you must repeat the procedures in Section 7.2 or 7.3 of this method with new samples using a shorter proposed maximum storage duration or improved handling and storage procedures.

**Determination of Bias and Precision**

8.0 What are the requirements for bias?

You must determine bias by comparing the results of sampling and analysis using the candidate test method against a reference value. The bias must be no more than ±10 percent for the candidate test method to be considered for application to multiple sources. A candidate test method with a bias greater than ±10 percent and less than or equal to ±30 percent can only be applied on a source-specific basis at the facility at which the validation testing was conducted. In this case, you must use a correction factor for all data collected in the future using the candidate test method. If the bias is more than ±30 percent, the candidate test method is unacceptable.

9.0 What are the requirements for precision?

You may use a paired sampling system or a quadruplicate sampling system to establish precision for isotopic spiking. You must use a quadruplicate sampling system to establish precision for analyte spiking or when comparing a candidate test method to a validated method. If you are using analyte spiking or isotopic spiking, the precision, expressed as the relative standard deviation (RSD) of the candidate test method, must be less than or equal to 20 percent. If you are comparing the candidate test method to a validated test method, the candidate test method must be at least as precise as the validated method as determined by an F test (see Section 11.2.2 of this method).

10.0 What calculations must I perform for isotopic spiking?

You must analyze the bias, RSD, precision, and data acceptance for isotopic spiking tests according to the provisions in Sections 10.1 through 10.4 of this method.
10.1 **Numerical Bias.** Calculate the numerical value of the bias using the results from the analysis of the isotopic spike in the field samples and the calculated value of the spike according to Equation 301-4.

\[ B = S_m - CS \]  

(Eq. 301-4)

Where:

- **B** = Bias at the spike level.
- **S_m** = Mean of the measured values of the isotopically-labeled analyte in the samples.
- **CS** = Calculated value of the isotopically-labeled spike level.

10.2 **Standard Deviation.** Calculate the standard deviation of the \( S_i \) values according to Equation 301-5.

\[ SD = \sqrt{\frac{\sum_{i=1}^{n} (S_i - S_m)^2}{n-1}} \]  

(Eq. 301-5)

Where:

- **SD** = Standard deviation of the candidate test method.
- **S_i** = Measured value of the isotopically-labeled analyte in the \( i^{th} \) field sample.
- **S_m** = Mean of the measured values of the isotopically-labeled analyte in the samples.
- **n** = Number of isotopically-spiked samples.

10.3 **T Test.** Test the bias for statistical significance by calculating the t-statistic using Equation 301-6. Use the standard deviation determined in Section 10.2 of this method and the numerical bias determined in Section 10.1 of this method.

\[ t = \frac{|B|}{SD} \left( \frac{1}{\sqrt{n}} \right) \]  

(Eq. 301-6)

Where:

- **t** = Calculated t-statistic.
- **B** = Bias at the spike level.
- **SD** = Standard deviation of the candidate test method.
- **n** = Number of isotopically spike samples.
Compare the calculated t-value with the critical value of the two-sided t-distribution at the 95 percent confidence level and n-1 degrees of freedom (see Table 301-3 of this method). When you conduct isotopic spiking according to the procedures specified in Sections 6.1 and 6.3 of this method as required, this critical value is 2.201 for 11 degrees of freedom. If the calculated t-value is less than or equal to the critical value, the bias is not statistically significant, and the bias of the candidate test method is acceptable. If the calculated t-value is greater than the critical value, the bias is statistically significant, and you must evaluate the relative magnitude of the bias using Equation 301-7.

\[ B_R = \left( \frac{B}{CS} \right) \times 100\% \]  
(Eq. 301-7)

Where:

\( B_R \) = Relative bias.
\( B \) = Bias at the spike level.
\( CS \) = Calculated value of the spike level.

If the relative bias is less than or equal to 10 percent, the bias of the candidate test method is acceptable for use at multiple sources. If the relative bias is greater than 10 percent but less than or equal to 30 percent, and if you correct all data collected with the candidate test method in the future for bias using the source-specific correction factor determined in Equation 301-8, the candidate test method is acceptable only for application to the source at which the validation testing was conducted and may not be applied to any other sites. If either of the preceding two cases applies, you may continue to evaluate the candidate test method by calculating its precision. If not, the candidate test method does not meet the requirements of Method 301.

\[ CF = \left( \frac{1}{1 + \frac{B}{CS}} \right) \]  
(Eq. 301-8)

Where:

\( CF \) = Source-specific bias correction factor.
\( B \) = Bias at the spike level.
\( CS \) = Calculated value of the spike level.

If the CF is outside the range of 0.70 to 1.30, the data and method are considered unacceptable.

10.4 **Precision.** Calculate the RSD according to Equation 301-9.

\[ RSD = \left( \frac{SD}{S_m} \right) \times 100 \]  
(Eq. 301-9)
Where:

\[ \text{RSD} \quad = \quad \text{Relative standard deviation of the candidate test method.} \]
\[ \text{SD} \quad = \quad \text{Standard deviation of the candidate test method calculated in Equation 301-5.} \]
\[ S_m \quad = \quad \text{Mean of the measured values of the spike samples.} \]

The data and candidate test method are unacceptable if the RSD is greater than 20 percent.

11.0 What calculations must I perform for comparison with a validated method?

If you are comparing a candidate test method to a validated method, then you must analyze the data according to the provisions in this section. If the data from the candidate test method fail either the bias or precision test, the data and the candidate test method are unacceptable. If the Administrator determines that the affected source has highly variable emission rates, the Administrator may require additional precision checks.

11.1 Bias Analysis. Test the bias for statistical significance at the 95 percent confidence level by calculating the t-statistic.

11.1.1 Bias. Determine the bias, which is defined as the mean of the differences between the candidate test method and the validated method \((d_i)\). Calculate \(d_i\) according to Equation 301-10.

\[
d_i = \frac{(V_{1i} + V_{2i})}{2} - \frac{(P_{1i} + P_{2i})}{2}
\]

(Eq. 301-10)

Where:

\( d_i \) = Difference in measured value between the candidate test method and the validated method for each quadruplicate sampling train.

\( V_{1i} \) = First measured value with the validated method in the \(i^{th}\) quadruplicate sampling train.

\( V_{2i} \) = Second measured value with the validated method in the \(i^{th}\) quadruplicate sampling train.

\( P_{1i} \) = First measured value with the candidate test method in the \(i^{th}\) quadruplicate sampling train.

\( P_{2i} \) = Second measured value with the candidate test method in the \(i^{th}\) quadruplicate sampling train.

Calculate the numerical value of the bias using Equation 301-11.

\[
B = \frac{\sum_{i=1}^{n} d_i}{n}
\]

(Eq. 301-11)

Where:

\( B \) = Numerical bias.
\[ d_i = \text{Difference between the candidate test method and the validated method for the } i^{th} \text{ quadruplicate sampling train.} \]

\[ n = \text{Number of quadruplicate sampling trains.} \]

**11.1.2 Standard Deviation of the Differences.** Calculate the standard deviation of the differences, SD\(_d\), using Equation 301-12.

\[
SD_d = \sqrt{\frac{\sum_{i=1}^{n} (d_i - d_m)^2}{n - 1}} 
\]  
(Eq. 301-12)

Where:

\[ SD_d = \text{Standard deviation of the differences between the candidate test method and the validated method.} \]

\[ d_i = \text{Difference in measured value between the candidate test method and the validated method for each quadruplicate sampling train.} \]

\[ d_m = \text{Mean of the differences, } d_i \text{ between the candidate test method and the validated method.} \]

\[ n = \text{Number of quadruplicate sampling trains.} \]

**11.1.3 T Test.** Calculate the t-statistic using Equation 301-13.

\[
t = \frac{|d_m|}{SD \sqrt{n}} 
\]  
(Eq. 301-13)

Where:

\[ t = \text{Calculated t-statistic.} \]

\[ d_m = \text{The mean of the differences, } d_i \text{ between the candidate test method and the validated method.} \]

\[ SD_d = \text{Standard deviation of the differences between the candidate test method and the validated method.} \]

\[ n = \text{Number of quadruplicate sampling trains.} \]

For the procedure comparing a candidate test method to a validated test method listed in Table 301-1 of this method, n equals six. Compare the calculated t-statistic with the critical value of the t-statistic, and determine if the bias is significant at the 95 percent confidence level (see Table 301-3 of this method). When six runs are conducted, as specified in Table 301-1 of this method, the critical value of the t-statistic is 2.571 for five degrees of freedom. If the calculated t-value is less than or equal to the critical value, the bias is not statistically significant and the data are acceptable. If the calculated t-value is greater than the critical value, the bias is statistically significant, and you must evaluate the magnitude of the relative bias using Equation 301-14.
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If the relative bias is less than or equal to 10 percent, the bias of the candidate test method is acceptable. On a source-specific basis, if the relative bias is greater than 10 percent but less than or equal to 30 percent, and if you correct all data collected in the future with the candidate test method for the bias using the correction factor, CF, determined in Equation 301-8 (using VS for CS), the bias of the candidate test method is acceptable for application to the source at which the validation testing was conducted. If either of the preceding two cases applies, you may continue to evaluate the candidate test method by calculating its precision. If not, the candidate test method does not meet the requirements of Method 301.

11.2 Precision. Compare the estimated variance (or standard deviation) of the candidate test method to that of the validated test method according to Sections 11.2.1 and 11.2.2 of this method. If a significant difference is determined using the F test, the candidate test method and the results are rejected. If the F test does not show a significant difference, then the candidate test method has acceptable precision.

11.2.1 Candidate Test Method Variance. Calculate the estimated variance of the candidate test method according to Equation 301-15.

\[ S_p^2 = \frac{\sum_{i=1}^{n} d_i^2}{2n} \]  
(Eq. 301-15)

Where:

\( S_p^2 \) = Estimated variance of the candidate test method.

\( d_i \) = The difference between the \( i \)th pair of samples collected with the candidate test method in a single quadruplicate train.

\( n \) = Total number of paired samples (quadruplicate trains).

Calculate the estimated variance of the validated test method according to Equation 301-16.

\[ S_v^2 = \frac{\sum_{i=1}^{n} d_i^2}{2n} \]  
(Eq. 301-16)

Where:
\( S_v^2 \) = Estimated variance of the validated test method.

\( d_i \) = The difference between the \( i^{th} \) pair of samples collected with the validated test method in a single quadruplicate train.

\( n \) = Total number of paired samples (quadruplicate trains).

11.2.2 The F test. Determine if the estimated variance of the candidate test method is greater than that of the validated method by calculating the F-value using Equation 301-17.

\[
F = \frac{S_p^2}{S_v^2}
\]

(Eq. 301-17)

Where:

\( F \) = Calculated F value.

\( S_p^2 \) = The estimated variance of the candidate test method.

\( S_v^2 \) = The estimated variance of the validated method.

Compare the calculated F value with the one-sided confidence level for F from Table 301-4 of this method. The upper one-sided confidence level of 95 percent for \( F_{(6,6)} \) is 4.28 when the procedure specified in Table 301-1 of this method for quadruplicate sampling trains is followed. If the calculated F value is greater than the critical F value, the difference in precision is significant, and the data and the candidate test method are unacceptable.

12.0 What calculations must I perform for analyte spiking?

You must analyze the data for analyte spike testing according to this section.

12.1 Bias Analysis. Test the bias for statistical significance at the 95 percent confidence level by calculating the t-statistic.

12.1.1 Bias. Determine the bias, which is defined as the mean of the differences between the spiked samples and the unspiked samples in each quadruplicate sampling train minus the spiked amount, using Equation 301-18.

\[
d_i = \frac{(S_{1i} + S_{2i})}{2} - \frac{(M_{1i} + M_{2i})}{2} - CS
\]

(Eq. 301-18)

Where:

\( d_i \) = Difference between the spiked samples and unspiked samples in each quadruplicate sampling train minus the spiked amount.

\( S_{1i} \) = Measured value of the first spiked sample in the \( i^{th} \) quadruplicate sampling train.

\( S_{2i} \) = Measured value of the second spiked sample in the \( i^{th} \) quadruplicate sampling train.
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\[ M_{1i} = \text{Measured value of the first unspiked sample in the } i^{th} \text{ quadruplicate sampling train.} \]

\[ M_{2i} = \text{Measured value of the second unspiked sample in the } i^{th} \text{ quadruplicate sampling train.} \]

\[ CS = \text{Calculated value of the spike level.} \]

Calculate the numerical value of the bias using Equation 301-19.

\[ B = \frac{\sum_{i}^{n} d_i}{n} \]  

(Eq. 301-19)

Where:

\[ B = \text{Numerical value of the bias.} \]

\[ d_i = \text{Difference between the spiked samples and unspiked samples in each quadruplicate sampling train minus the spiked amount.} \]

\[ n = \text{Number of quadruplicate sampling trains.} \]


\[ SD_d = \sqrt{\frac{\sum_{i}^{n} (d_i - d_m)^2}{n - 1}} \]  

(Eq. 301-20)

Where:

\[ SD_d = \text{Standard deviation of the differences of paired samples.} \]

\[ d_i = \text{Difference between the spiked samples and unspiked samples in each quadruplicate sampling train minus the spiked amount.} \]

\[ d_m = \text{The mean of the differences, } d_i, \text{ between the spiked samples and unspiked samples.} \]

\[ n = \text{Total number of quadruplicate sampling trains.} \]

12.1.3 T-test. Calculate the t-statistic using Equation 301-21, where \( n \) is the total number of test sample differences (\( d_i \)). For the quadruplicate sampling system procedure in Table 301-1 of this method, \( n \) equals six.

\[ t = \frac{|d_m|}{\left(\frac{SD_d}{\sqrt{n}}\right)} \]  

(Eq. 301-21)

Where:
\( t \) = Calculated t-statistic.

\( d_m \) = Mean of the difference, \( d_i \), between the spiked samples and unspiked samples.

\( \text{SD}_d \) = Standard deviation of the differences of paired samples.

\( n \) = Number of quadruplicate sampling trains.

Compare the calculated t-statistic with the critical value of the t-statistic, and determine if the bias is significant at the 95 percent confidence level. When six quadruplicate runs are conducted, as specified in Table 301-1 of this method, the 2-sided confidence level critical value is 2.571 for the five degrees of freedom. If the calculated t-value is less than the critical value, the bias is not statistically significant and the data are acceptable. If the calculated t-value is greater than the critical value, the bias is statistically significant and you must evaluate the magnitude of the relative bias using Equation 301-22.

\[
B_R = \frac{B}{\text{CS}} \times 100\% 
\]  
(Eq. 301-22)

Where:

\( B_R \) = Relative bias.

\( B \) = Bias at the spike level from Equation 301-19.

\( \text{CS} \) = Calculated value at the spike level.

If the relative bias is less than or equal to 10 percent, the bias of the candidate test method is acceptable. On a source-specific basis, if the relative bias is greater than 10 percent but less than or equal to 30 percent, and if you correct all data collected with the candidate test method in the future for the magnitude of the bias using Equation 301-8, the bias of the candidate test method is acceptable for application to the tested source at which the validation testing was conducted. Proceed to evaluate precision of the candidate test method.

12.2 Precision. Calculate the standard deviation using Equation 301-23.

\[
SD = \sqrt{\frac{\sum_{i=1}^{n} (S_i - S_m)^2}{n-1}} 
\]  
(Eq. 301-23)

Where:

\( SD \) = Standard deviation of the candidate test method.

\( S_i \) = Measured value of the analyte in the \( i^{th} \) spiked sample.

\( S_m \) = Mean of the measured values of the analyte in all the spiked samples.

\( n \) = Number of spiked samples.
Calculate the RSD of the candidate test method using Equation 301-9, where SD and Sm are the values from Equation 301-23. The data and candidate test method are unacceptable if the RSD is greater than 20 percent.

13.0 How do I conduct tests at similar sources?

If the Administrator has approved the use of an alternative test method to a test method required in 40 CFR part 59, 60, 61, 63, or 65 for an affected source, and you would like to apply the alternative test method to a similar source, then you must petition the Administrator as described in Section 17.1.1 of this method.

Optional Requirements

14.0 How do I use and conduct ruggedness testing?

Ruggedness testing is an optional requirement for validation of a candidate test method that is intended for the source where the validation testing was conducted. Ruggedness testing is required for validation of a candidate test method intended to be used at multiple sources. If you want to use a validated test method at a concentration that is different from the concentration in the applicable emission limitation under 40 CFR part 59, 60, 61, 63, or 65, or for a source category that is different from the source category that the test method specifies, then you must conduct ruggedness testing according to the procedures in Reference 18.16 of Section 18.0 of this method and submit a request for a waiver for conducting Method 301 at that different source category according to Section 17.1.1 of this method.

Ruggedness testing is a study that can be conducted in the laboratory or the field to determine the sensitivity of a method to parameters such as analyte concentration, sample collection rate, interferent concentration, collection medium temperature, and sample recovery temperature. You conduct ruggedness testing by changing several variables simultaneously instead of changing one variable at a time. For example, you can determine the effect of seven variables in only eight experiments. (W.J. Youden, Statistical Manual of the Association of Official Analytical Chemists, Association of Official Analytical Chemists, Washington, DC, 1975, pp. 33-36).

15.0 How do I determine the Limit of Detection for the candidate test method?

Determination of the Limit of Detection (LOD) as specified in Sections 15.1 and 15.2 of this method is required for source-specific method validation and validation of a candidate test method intended to be used for multiple sources.

15.1 Limit of Detection. The LOD is the minimum concentration of a substance that can be measured and reported with 99 percent confidence that the analyte concentration is greater than zero. For this protocol, the LOD is defined as three times the standard deviation, S\text{m}, at the blank level.

15.2 Purpose. The LOD establishes the lower detection limit of the candidate test method. You must calculate the LOD using the applicable procedures found in Table 301-5 of this method. For candidate test methods that collect the analyte in a sample matrix prior to an analytical measurement, you must determine the LOD using Procedure I in Table 301-5 of this method by calculating a method detection limit (MDL) as described in 40 CFR part 136, appendix B. For the purposes of this section, the LOD is equivalent to the calculated MDL. For radiochemical methods, use the Multi-Agency Radiological Laboratory Analytical Protocols (MARLAP) Manual (i.e., use the minimum detectable concentration (MDC) and not the LOD) available at https://www.epa.gov/radiation/marlap-manual-and-supporting-documents.
Other Requirements and Information

16.0 How do I apply for approval to use a candidate test method?

16.1 Submitting Requests. You must request to use a candidate test method according to the procedures in §63.7(f) or similar sections of 40 CFR parts 59, 60, 61, and 65 (§59.104, §59.406, §60.8(b), §61.13(h)(1)(ii), or §65.158(a)(2)(iii)). You cannot use a candidate test method to meet any requirement under these parts until the Administrator has approved your request. The request must include a field validation report containing the information in Section 16.2 of this method. You must submit the request to the Group Leader, Measurement Technology Group, U.S. Environmental Protection Agency, E143-02, Research Triangle Park, NC 27711.

16.2 Field Validation Report. The field validation report must contain the information in Sections 16.2.1 through 16.2.8 of this method.

16.2.1 Regulatory objectives for the testing, including a description of the reasons for the test, applicable emission limits, and a description of the source.

16.2.2 Summary of the results and calculations shown in Sections 6.0 through 16.0 of this method, as applicable.

16.2.3 Reference material certification and value(s).

16.2.4 Discussion of laboratory evaluations.

16.2.5 Discussion of field sampling.

16.2.6 Discussion of sample preparation and analysis.

16.2.7 Storage times of samples (and extracts, if applicable).

16.2.8 Reasons for eliminating any results.

17.0 How do I request a waiver?

17.1 Conditions for Waivers. If you meet one of the criteria in Section 17.1.1 or 17.1.2 of this method, the Administrator may waive the requirement to use the procedures in this method to validate an alternative or other candidate test method. In addition, if the EPA currently recognizes an appropriate test method or considers the candidate test method to be satisfactory for a particular source, the Administrator may waive the requirement for you to demonstrate the applicability of the method to your source.

17.1.1 Similar Sources. If the alternative or other candidate test method that you want to use was validated for source-specific application at another source and you can demonstrate to the Administrator’s satisfaction that your affected source is similar to that validated source, then the Administrator may waive the requirement for you to validate the alternative or other candidate test method. One procedure you may use to demonstrate the applicability of the method to your affected source is to conduct a ruggedness test as described in Section 14.0 of this method.

17.1.2 Documented Methods. If the bias, precision, LOD, or ruggedness of the alternative or other candidate test method that you are proposing have been demonstrated through laboratory tests or protocols different from this method, and you can demonstrate to the Administrator’s satisfaction that the
bias, precision, LOD, or ruggedness apply to your application, then the Administrator may waive the requirement to use this method or to use part of this method.

17.2 Submitting Applications for Waivers. You must sign and submit each request for a waiver from the requirements in this method in writing. The request must be submitted to the Group Leader, Measurement Technology Group, U.S. Environmental Protection Agency, E143-02, Research Triangle Park, NC 27711.

17.3 Information Application for Waiver. The request for a waiver must contain a thorough description of the candidate test method, the intended application, and results of any validation or other supporting documents. The request for a waiver must contain, at a minimum, the information in Sections 17.3.1 through 17.3.4 of this method. The Administrator may request additional information if necessary to determine whether this method can be waived for a particular application.

17.3.1 A Clearly Written Test Method. The candidate test method should be written preferably in the format of 40 CFR part 60, appendix A, Test Methods. Additionally, the candidate test must include an applicability statement, concentration range, precision, bias (accuracy), and minimum and maximum storage durations in which samples must be analyzed.

17.3.2 Summaries of Previous Validation Tests or Other Supporting Documents. If you use a different procedure from that described in this method, you must submit documents substantiating the bias and precision values to the Administrator’s satisfaction.

17.3.3 Ruggedness Testing Results. You must submit results of ruggedness testing conducted according to Section 14.0 of this method, sample stability conducted according to Section 7.0 of this method, and detection limits conducted according to Section 15.0 of this method, as applicable. For example, you would not need to submit ruggedness testing results if you will be using the method at the same affected source and level at which it was validated.

17.3.4 Applicability Statement and Basis for Waiver Approval. Discussion of the applicability statement and basis for approval of the waiver. This discussion should address as applicable the following: applicable regulation, emission standards, effluent characteristics, and process operations.

18.0 Where can I find additional information?

You can find additional information in the references in Sections 18.1 through 18.18 of this method.


18.18 Statistical Table, [http://www.math.usask.ca/~szafron/Stats244/f_table_0_05.pdf](http://www.math.usask.ca/~szafron/Stats244/f_table_0_05.pdf).

19.0 Tables.

### Table 301-1. Sampling Procedures

<table>
<thead>
<tr>
<th>If you are...</th>
<th>You must collect...</th>
</tr>
</thead>
<tbody>
<tr>
<td>Comparing the candidate test method against a validated method</td>
<td>A total of 24 samples using a quadruplicate sampling system (a total of six sets of replicate samples). In each quadruplicate sample set, you must use the validated test method to collect and analyze half of the samples.</td>
</tr>
<tr>
<td>Using isotopic spiking (can only be used with methods capable of measurement of multiple isotopes simultaneously)</td>
<td>A total of 12 samples, all of which are spiked with isotopically-labeled analyte. You may collect the samples either by obtaining six sets of paired samples or three sets of quadruplicate samples.</td>
</tr>
<tr>
<td>Using analyte spiking</td>
<td>A total of 24 samples using the quadruplicate sampling system (a total of six sets of replicate samples – two spiked and two unspiked).</td>
</tr>
</tbody>
</table>

### Table 301-2. Storage and Sampling Procedures for Stack Test Emissions

<table>
<thead>
<tr>
<th>If you are...</th>
<th>With...</th>
<th>Then you must...</th>
</tr>
</thead>
<tbody>
<tr>
<td>Using isotopic or analyte spiking procedures.</td>
<td>Sample container (bag or canister) or impinger sampling systems that are not subject to dilution or other preparation steps.</td>
<td>Analyze six of the samples within 7 days and then analyze the same six samples at the proposed maximum storage duration or 2 weeks after the initial analysis.</td>
</tr>
<tr>
<td></td>
<td>Sorbent and impinger sampling systems that require extraction or digestion.</td>
<td>Extract or digest six of the samples within 7 days and extract or digest six other samples at the proposed maximum storage duration or 2 weeks after the first extraction or digestion. Analyze an aliquot of the first six extracts (digestates) within 7 days and proposed maximum storage duration or 2 weeks after the initial analysis. This will allow analysis of extract storage impacts.</td>
</tr>
<tr>
<td>Comparing a candidate test method against a validated test method</td>
<td>Sample container (bag or canister) or impinger sampling systems that are not subject to dilution or other preparation steps.</td>
<td>Analyze six samples within 7 days. Analyze another set of six samples at the proposed maximum storage time or within 2 weeks of the initial analysis.</td>
</tr>
<tr>
<td></td>
<td>Sorbent sampling systems that require thermal desorption.</td>
<td>Analyze at least six of the candidate test method samples within 7 days and then analyze the same six samples at the proposed maximum storage duration or within 2 weeks of the initial analysis.</td>
</tr>
</tbody>
</table>
Table 301-2. Storage and Sampling Procedures for Stack Test Emissions

<table>
<thead>
<tr>
<th>If you are…</th>
<th>With…</th>
<th>Then you must…</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sorbent and impinger sampling systems that require</td>
<td>Extract or digest six of the candidate test</td>
<td>Extract or digest six of the candidate test method samples within 7 days and</td>
</tr>
<tr>
<td>extraction or digestion.</td>
<td>extraction method samples within 7 days and</td>
<td>digest six other samples at the proposed maximum storage duration or within 2</td>
</tr>
<tr>
<td></td>
<td>extract or digest six other samples at the</td>
<td>weeks of the first extraction or digestion. Analyze an aliquot of the first</td>
</tr>
<tr>
<td></td>
<td>proposed maximum storage duration or within</td>
<td>six extracts (digestates) within 7 days and an aliquot at the proposed</td>
</tr>
<tr>
<td></td>
<td>2 weeks of the first extraction or digestion.</td>
<td>maximum storage durations or within 2 weeks of the initial analysis. This</td>
</tr>
<tr>
<td></td>
<td></td>
<td>will allow analysis of extract storage impacts.</td>
</tr>
<tr>
<td>Sorbent systems that require thermal desorption.</td>
<td>Analyze six samples within 7 days. Analyze</td>
<td>Analyze six samples within 7 days. Analyze another set of six samples at the</td>
</tr>
<tr>
<td></td>
<td>another set of six samples at the proposed</td>
<td>proposed maximum storage duration or within 2 weeks of the initial analysis.</td>
</tr>
<tr>
<td></td>
<td>maximum storage duration or within 2 weeks</td>
<td></td>
</tr>
<tr>
<td></td>
<td>of the initial analysis.</td>
<td></td>
</tr>
</tbody>
</table>

Table 301-3. Critical Values of t for the Two-Tailed 95 Percent Confidence Limit

<table>
<thead>
<tr>
<th>Degrees of freedom</th>
<th>t₀.₅</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>12.706</td>
</tr>
<tr>
<td>2</td>
<td>4.303</td>
</tr>
<tr>
<td>3</td>
<td>3.182</td>
</tr>
<tr>
<td>4</td>
<td>2.776</td>
</tr>
<tr>
<td>5</td>
<td>2.571</td>
</tr>
<tr>
<td>6</td>
<td>2.447</td>
</tr>
<tr>
<td>7</td>
<td>2.365</td>
</tr>
<tr>
<td>8</td>
<td>2.306</td>
</tr>
<tr>
<td>9</td>
<td>2.262</td>
</tr>
<tr>
<td>10</td>
<td>2.228</td>
</tr>
<tr>
<td>11</td>
<td>2.201</td>
</tr>
<tr>
<td>12</td>
<td>2.179</td>
</tr>
<tr>
<td>13</td>
<td>2.160</td>
</tr>
<tr>
<td>14</td>
<td>2.145</td>
</tr>
<tr>
<td>15</td>
<td>2.131</td>
</tr>
<tr>
<td>16</td>
<td>2.120</td>
</tr>
<tr>
<td>17</td>
<td>2.110</td>
</tr>
<tr>
<td>18</td>
<td>2.101</td>
</tr>
<tr>
<td>19</td>
<td>2.093</td>
</tr>
<tr>
<td>20</td>
<td>2.086</td>
</tr>
</tbody>
</table>

¹ Adapted from Reference 18.17 in section 18.0.
Table 301-4. Upper Critical Values of the F Distribution for the 95 Percent Confidence Limit\(^1\)

<table>
<thead>
<tr>
<th>Numerator ((k_1)) and denominator ((k_2)) degrees of freedom</th>
<th>(F{F &gt; F_{0.05}(k_1,k_2)})</th>
</tr>
</thead>
<tbody>
<tr>
<td>1,1</td>
<td>161.40</td>
</tr>
<tr>
<td>2,2</td>
<td>19.00</td>
</tr>
<tr>
<td>3,3</td>
<td>9.28</td>
</tr>
<tr>
<td>4,4</td>
<td>6.39</td>
</tr>
<tr>
<td>5,5</td>
<td>5.05</td>
</tr>
<tr>
<td>6,6</td>
<td>4.28</td>
</tr>
<tr>
<td>7,7</td>
<td>3.79</td>
</tr>
<tr>
<td>8,8</td>
<td>3.44</td>
</tr>
<tr>
<td>9,9</td>
<td>3.18</td>
</tr>
<tr>
<td>10,10</td>
<td>2.98</td>
</tr>
<tr>
<td>11,11</td>
<td>2.82</td>
</tr>
<tr>
<td>12,12</td>
<td>2.69</td>
</tr>
<tr>
<td>13,13</td>
<td>2.58</td>
</tr>
<tr>
<td>14,14</td>
<td>2.48</td>
</tr>
<tr>
<td>15,15</td>
<td>2.40</td>
</tr>
<tr>
<td>16,16</td>
<td>2.33</td>
</tr>
<tr>
<td>17,17</td>
<td>2.27</td>
</tr>
<tr>
<td>18,18</td>
<td>2.22</td>
</tr>
<tr>
<td>19,19</td>
<td>2.17</td>
</tr>
<tr>
<td>20,20</td>
<td>2.12</td>
</tr>
</tbody>
</table>

\(^1\) Adapted from References 18.17 and 18.18 in section 18.0.
Table 301-5. Procedures for Estimating S₀

<table>
<thead>
<tr>
<th>Procedure I</th>
<th>Procedure II</th>
</tr>
</thead>
<tbody>
<tr>
<td>If the estimated LOD (LOD₁, expected approximate LOD concentration level) is no more than twice the calculated LOD or an analyte in a sample matrix was collected prior to an analytical measurement, use Procedure I as follows.</td>
<td>If the estimated LOD (LOD₁, expected approximate LOD concentration level) is greater than twice the calculated LOD, use Procedure II as follows.</td>
</tr>
<tr>
<td>Procedure I</td>
<td>Procedure II</td>
</tr>
<tr>
<td>Determine the LOD by calculating a method detection limit (MDL) as described in 40 CFR part 136, appendix B.</td>
<td>Prepare two additional standards (LOD₂ and LOD₃) at concentration levels lower than the standard used in Procedure I (LOD₁).</td>
</tr>
<tr>
<td>Sample and analyze each of these standards (LOD₂ and LOD₃) at least seven times.</td>
<td>Sample and analyze each of these standards (LOD₂ and LOD₃) at least seven times.</td>
</tr>
<tr>
<td>Calculate the standard deviation (S₂ and S₃) for each concentration level.</td>
<td>Calculate the standard deviation (S₂ and S₃) for each concentration level.</td>
</tr>
<tr>
<td>Plot the standard deviations of the three test standards (S₁, S₂ and S₃) as a function of concentration.</td>
<td>Plot the standard deviations of the three test standards (S₁, S₂ and S₃) as a function of concentration.</td>
</tr>
<tr>
<td>Draw a best-fit straight line through the data points and extrapolate to zero concentration. The standard deviation at zero concentration is S₀.</td>
<td>Draw a best-fit straight line through the data points and extrapolate to zero concentration. The standard deviation at zero concentration is S₀.</td>
</tr>
<tr>
<td>Calculate the LOD₀ (referred to as the calculated LOD) as 3 times S₀.</td>
<td>Calculate the LOD₀ (referred to as the calculated LOD) as 3 times S₀.</td>
</tr>
</tbody>
</table>