

1 **Proposed Revisions 2-2021:**

2 **Appendix L: AOAC Recommended Guidelines for Stakeholder Program**
3 **on Infant Formula and Adult Nutritionals (SPIFAN) Single-Laboratory**
4 **Validation**
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6

7 **1 Definitions**

- 8 (a) Reference Material: A sufficiently stable, homogeneous sample matrix containing a specified analyte or
9 group of analytes with a content that is reliable and reproducible [1]. The sample has been established
10 to be fit for its intended use in a measurement process between two or more laboratories [2]. The
11 measurable quantity is the mean of a specified population of measurements [11]. The stability and
12 homogeneity may be determined as described elsewhere [3].
- 13 (b) Certified Reference Material (CRM): A reference material characterized by a recognized procedure for
14 determining analyte concentration accompanied by a certificate issued by an authoritative body that
15 provides the value of the concentration, its associated uncertainty, and a statement of metrological
16 traceability [4].
- 17 (c) Reference Standard: A substance of known identity and purity with accompanying certificate of
18 analysis from an authoritative body and used to prepare calibration standards and/or for the
19 calibration of other measurement standards. Moisture content should be monitored to ensure stability
20 and purity.
- 21 (d) Limit of Detection (LoD): the lowest concentration of analyte that can be confidently detected.
- 22 (e) Limit of Quantification (LoQ): the lowest concentration of analyte that can be determined with
23 acceptable precision and accuracy.
- 24 (f) Matrix Blank: A product matrix that does not contain the analyte of interest (it may contain
25 endogenous levels below LOD/LOQ) but does contain all the same components as the sample solution.
- 26 (g) Method Reagent Blank: Blank (e.g. water, buffer, solvent, or any other diluent which is free from the
27 analyte) analyzed by the method in place of a sample. It identifies the amount of the signal that is due
28 to the reagents used in the preparation of the samples.

29
30 **2 General**

- 31 (a) All methods for (a) given analyte(s) will be subjected to a common SLV protocol utilizing available
32 SPIFAN matrices. When SPIFAN matrices are not available, sample types to be considered can be
33 found in the associated SMPR.
- 34 (b) Assessment of the various parameters for single laboratory validation are described elsewhere [5].
- 35 (c) SLV protocols may vary somewhat between analytes, depending on the specific demands
36 associated with each.
- 37 Study directors (SDs) for each analyte will agree on final details of the required SLV protocol.
- 38 (d) System Suitability criteria indicating method/system performance is acceptable and will be
39 generated during SLV [6, Annex B].
- 40 (e) Ruggedness of an analytical method could be evaluated to measure the capacity to remain

1 unaffected by small but deliberate variations in method parameters [7, 8, 9]. By evaluating
2 ruggedness, one provides an indication of the method's robustness during normal usage. If
3 robustness/ruggedness was part of the method development phase, results of this can be
4 documented in the SLV report.

5 (f) Units of measure for reported values and figures of merit must be consistent with those stated in
6 the associated Standard Method Performance Requirement (SMPR).

7

8 **3 Materials**

9 (a) Use of a CRM where available is recommended to assess method accuracy as bias. The CRM should be
10 accompanied by documentation (certificate) issued by an authoritative body.

11 (b) If a variety of matrices with different physical and chemical properties are defined in the SMPR, then a
12 CRM of each type of matrix shall be included if available, otherwise see 3C.

13 (c) Where a CRM is not available, the concentration of the analyte(s) being studied in a reference material
14 are assessed using preferably two appropriate orthogonal techniques. Statistically equivalent results
15 from these analyses are requested with a minimum of two independent analyses, preferably determined
16 by different laboratories. The completed SLV report should be accompanied by assessment protocols
17 and results.

18 (d) Any reference standard used needs to be accompanied with a certificate of analysis, stating supplier,
19 identity, batch number, purity and basis for the purity statement in the SLV report. The purity of the
20 reference standards used should be established, understood and fit for purpose. If a non-commercial
21 reference standard is used, its origin needs to be clearly identified along with all pertinent information
22 demonstrating purity and/or analyte concentration and the means with which these were determined.

23 (e) If a variety of matrices with different physical and chemical properties are defined in the SMPR, the
24 number of matrices needs to be at least one for each matrix type. The matrix sources should cover the
25 range of expected concentrations of the analyte(s) of interest. If only a single matrix is studied, then ≥ 3
26 sources are recommended, preferably with different attributes (e.g. maturity, varieties, age).

27 **4 Linearity/Calibration Fit**

28 (a) Minimum of six (6) levels that span the desired working range as described in the SMPR.

29 (b) Assessment of heteroscedasticity should be performed; for example, calculation of relative error of
30 back-calculated concentrations¹

31 (c) Minimum of three (3) independent experiments to confirm the linearity/calibration fit.

32 **5 LOD/LOQ**

33 To determine the LOD and LOQ, the Standard Deviation and Blank Mean of sample measurements is
34 obtained from repeated measurements of samples with a relevant low concentration (e.g. a
35 homogeneous Matrix Blank or a Method Reagent Blank spiked at low level)³ [12, 13]. It is
36 recommended that a minimum of twenty (20) sample measurements be used and that these
37 measurements, where possible, come from a variety of samples and over several days.

38 $LOD = \text{Blank Mean} + 3 \text{ Standard Deviations}$

¹ No specific criterion in SMPR; recommend calibration errors to be <15%. Along the whole range. Higher calibration error at LLOQ can be accepted (see FDA 2018, Bioanalytical Method Validation, guidance for industry).

1 LOQ = Blank Mean + 10 Standard Deviations

2 The LOD/LOQ could also be determined via the signal/noise ratio procedure. Relevant low
3 concentration samples (close to expected LOD/LOQ in at least 3 concentration levels) are analyzed
4 three times. The signal/noise ratio can be calculated for each replicate of the spiked samples. A graph
5 can be plotted of the signal/noise ratio versus the concentration. After a linear regression analysis, the
6 intercept and slope determine the LOD/LOQ:

$$7 \quad LOD = \frac{3 - \text{Intercept}}{\text{Slope}}$$

$$8 \quad LOQ = \frac{10 - \text{Intercept}}{\text{Slope}}$$

9 Alternative approaches are possible as suggested in literature elsewhere [13].

10 11 **6 Selectivity**

- 12 (a) Selectivity is the degree to which the method can quantify the target analyte in the presence of other
13 analytes, matrices, or other potentially interfering materials. No explicit proposals for evaluating
14 selectivity are suggested, however methods must be tested in the presence of accompanying analytes or
15 matrices most likely to interfere. The freedom from effects of interfering materials can be studied using
16 various samples, ranging from pure measurement standards to mixtures with complex matrices. The
17 recovery of the analyte(s) of interest should be determined and the influences of suspected
18 interferences stated [6]. Examples of selectivity tests for chromatographic methods are described
19 elsewhere [10, 13]. Methods with a known interference should be modified prior to SLV. If method
20 selectivity was part of the method development phase, results of this can be documented in the SLV
21 report.
- 22 (b) Useful strategies for completing selectivity vary from analyte to analyte. Therefore, the Study Directors
23 for each analyte will decide on an acceptable practice.

24 25 **7 Precision**

26 All samples selected for precision studies will be analyzed in duplicate on each of six (6) days using multiple
27 analysts and instruments as practical for the different days. Fresh reagents and working standards should
28 be used each day. Reports will include information of number of analysts, instruments, etc. The number
29 of matrices may vary between analytes.

30 (a) Precision data using a CRM or reference material should be included for all candidate methods.

31 (b) Estimate within-day repeatability, between-day repeatability (intermediate precision) and an,
32 estimation of measurement uncertainty for each sample type. Estimates pooled across the sample
33 types may also be useful, if appropriate.

34 **8 Accuracy (Trueness)**

35 (a) *Analysis of CRM or Reference Material.*

- 36 • A minimum of nine (9) independent replicates of CRM should be tested across three (3) days (e.g.
37 3 x 3) and compared to certified or reference values provided.
- 38 • Where a CRM is not available, a suitable reference material may be substituted and compared to

1 the established levels (see paragraph 3C). A *t*-test or other statistical test should be used to
2 evaluate method bias.

3 (b) *Spike recovery*.

4 (1) Recovery will be determined from an appropriate sampling of a range of infant formula and adult
5 nutritional matrices (use of the recognized SPIFAN kit is recommended where appropriate, otherwise
6 the Study Directors may agree on the samples to be included).

7 (2) Each selected matrix will be spiked at three levels. Use spike levels covering the analytical range
8 specified in the SMPR.

9 (3) Spiked and unspiked samples will be analyzed in triplicates on each of three (3) days.

10 (4) The daily mean of unspiked samples will be used for calculating individual recoveries.

11 12 References

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