

'Web course "LC-MS Method Validation"

University of Tartu

[https://sisu.ut.ee/lcms\\_method\\_validation/](https://sisu.ut.ee/lcms_method_validation/)

**NB!** “This table is meant as a general “big picture” comparison between the different guidelines.

The wording in this table does not necessarily exactly match the wording in the guidelines.

For the definitive wording the user **must** consult the original documents.”

Validation parameter	FDA 2018 bioanalysis [1]	EMA 2022 [2]	ICH 2005 [3]	Eurachem 2014 [4]
<b>Selectivity</b>	Blank from at least 6 individual sources;  <b>Acceptance Criteria:</b> Blank and zero: no interference at analyte and IS RTs. Spiked samples: $\pm 20\%$ LLOQ. Blank: IS response $< 5\%$ of Cals and QCs average IS responses.	Blank from at least 6 individual sources;  <b>Acceptance Criteria:</b> Analyte response $< 20\%$ of LLOQ; IS response $< 5\%$		Test samples and RMs Candidate and other independent methods  Also test samples with suspected interferences
<b>Specificity</b>	Assess for interference by cross-reacting molecules, concomitant medications, bio-transformed species, etc.  <b>Acceptance Criteria:</b> Same as Selectivity.		Blanks, matrix-matched samples. <b>Impurities, if applicable:</b> Spiking blanks with impurities and/or excipients. Degradation experiments: light, heat, humidity, acid/base hydrolysis and oxidation	
<b>Carryover</b>	Assess impact of carryover on accuracy.  <b>Acceptance Criteria:</b> $< 20\%$ of LLOQ	Inject blank after a high concentration sample or calibration;  <b>Acceptance Criteria:</b> Blank response: $< 20\%$ of LLOQ IS response: $< 5\%$		
<b>Linearity / Calibration Curve</b>	Matrix-matched, Blank, zero, 6 level (inc. LLOQ),  <b>Acceptance Criteria:</b> LLOQ $\pm 20\%$ ,	Matrix-matched, Blank, zero, 3 runs x 6 levels (inc. LLOQ and ULOQ) (optional, 2 parallels),  <b>Acceptance Criteria:</b>	$\geq 5$ levels  <b>Range:</b> assay: from 80 to 120 % of target conc.;	Instrument and method working range.  <b>Range of interest:</b> Blank, 6 - 10 levels evenly spaced

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	<p>others <math>\pm 15\%</math>, 75% (or min. 6 calibrator levels) should meet the criteria, Cal runs need to be reproducible</p>	<p>LLOQ <math>\pm 20\%</math>, others <math>\pm 15\%</math>, 75% (min. 6 calibrator levels) should meet the criteria 50% per level should meet the criteria</p>	<p>content uniformity: from 70 to 130 % of target conc.,  dissolution testing: <math>\pm 20\%</math> over the specified range;  impurity: from reporting level to 120% of the specification;</p>	<p>expected range <math>\pm 10\%</math> / <math>\pm 20\%</math>.  <b>Linear range:</b> Blank, 2-3 parallels x 6-10 levels evenly spaced  <b>Determine if linear range is fit:</b> Blank, reference materials or spiked sample blanks, 2-3 parallels x 6-10 levels evenly spaced  regression plot, residuals plot, regression statistics</p>
<b>Accuracy and Precision (A &amp; P)</b>	<p>3 runs x 4 levels (LLOQ, L, M, H QC) x 5 parallels, Over several days,  <b>Within-run and between runs</b>  <b>Accuracy:</b> LLOQ: <math>\pm 20\%</math> from nominal conc. others: <math>\pm 15\%</math> from nominal conc.  <b>Precision:</b> LLOQ: <math>\pm 20\%</math> RSD others: <math>\pm 15\%</math> RSD</p>	<p><b>Within-run QCs</b> 4 levels (LLOQ, L, M, H) x 5 parallels LLOQ: <math>\pm 20\%</math> others: <math>\pm 15\%</math> (mean conc. from nominal value or RSD)  <b>Between runs QCs</b> 3 runs (LLOQ, L, M, H) 2 different days LLOQ: <math>\pm 20\%</math> others: <math>\pm 15\%</math> (mean conc. from nominal value or RSD)</p>	<p><b>Repeatability:</b> 3 levels x 3 parallels OR 6 determinations at target conc.  <b>Intermediate precision:</b> Several days, analysts, equipment, etc.  Not necessary to study effects individually.  Experiment design is encouraged.</p>	<p><b>Accuracy:</b> Blanks, CRMs and/or spiked samples (if RM not available).  10 parallels per level  Alternatively: RM/test sample using candidate method and alternative method.  <b>Precision:</b> RMs, surplus test samples or spiked sample blanks at various levels  <b>Repeatability:</b> Same analyst, equipment, laboratory, short timescale.  6-15 parallels  <b>Intermediate precision:</b> Different analysts, equipment, same laboratory, extended timescale.  6-15 parallels for each material.  <b>Repeatability and intermediate precision in one study:</b></p>

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				Different analysts, equipment, same laboratory, extended timescale.  6-15 runs x 2 parallels  ANOVA to calculate repeatability standard deviation and intermediate precision standard deviation
<b>Matrix effect</b>	Matrix effects should be assessed and eliminated.	≥ 6 lots of individual blank matrices. 3 parallels. Pooled matrix should not be used.  CV: ≤ 15 % at L and H levels		
<b>Recovery</b>	QC (L, M, H) extracted samples vs blank extracts spiked post extraction	typically at 3 conc. (L, M, H)		
<b>Stability</b>	Stock solution, freeze-thaw, bench-top, long-term, processed sample, auto-sampler  3 parallels at L and H  <b>Acceptance Criteria:</b> Accuracy: ± 15% of nominal conc.	Stock and working solution, freeze and thaw, short term, long term, processed sample, on-instrument/auto-sampler, At L and H levels  <b>Acceptance Criteria:</b> Mean conc. at each level: ±15% of the nominal conc.		
<b>Sensitivity / LLOQ</b>	LLOQ: Lowest non-zero standard  <b>Acceptance Criteria:</b> Response at LLOQ ≥ 5 x zero response Accuracy ± 20% & Precision ≤ 20% (3 runs x ≥ 5 parallels)	Lowest calibration standard  <b>Acceptance Criteria:</b> Accuracy ± 20% and precision ≤ 20%	<b>LoD and LoQ:</b> Visual evaluation  OR  LoD: S/N of 3 or 2:1 LoQ: S/N of 10:1  OR  Based on response SD and Slope: $LoD = 3.3 \times \frac{\sigma}{slope}$ $LoQ = 10 \times \frac{\sigma}{slope}$  σ determined from: 1) standard deviation of several blanks	<b>CC<sub>α</sub>, CC<sub>β</sub></b> refer to EU Commission Decision 2021/808/EC and ISO 11843-2:2007  <b>LoD and LoQ:</b> Blank samples, test samples or spiked samples, concentrations of analyte close to or below the expected LOD  6 - 15 parallels  $LoD = 3 \times s_0'$ $LoQ = k_Q \times s_0'$  For calculation of s <sub>0</sub> '(modified standard deviation) refer to the guide, Other alternatives suggested as well.

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			2) calibration graph in LoD region, residual standard deviation or y-intercept standard deviation	
<b>Robustness / Ruggedness</b>			<p>System suitability parameters should be established.</p> <p><b>Examples for study:</b></p> <ul style="list-style-type: none"> <li>- stability of analytical solutions;</li> <li>- extraction time.</li> <li>- influence of variations of pH in a mobile phase;</li> <li>- influence of variations in mobile phase composition;</li> <li>- different columns (different lots and/or suppliers);</li> <li>- temperature;</li> <li>- flow rate.</li> </ul>	<p>Variables with significant effects must be identified,</p> <p>RMs or test samples,</p> <p>Most effective with experimental designs: e.g. Plackett-Burman experimental design for start.</p> <p>Rank the variables in order of the greatest effect on method performance.</p> <p>Significance tests to determine whether observed effects are statistically significant.</p>
<b>Other Validation Runs</b>	<p>3 QCs (L, M, H) in duplicates</p> <p><b>Run Acceptance Criteria:</b></p> <p>Cals: Same as calibration curve.</p> <p>QCs: ≥ 67% of QCs ± 15% ≥ 50% of QCs per level ± 15%</p>			
<b>Quality Controls (QC)</b>	<p><b>Accuracy and Precision:</b></p> <p>4 lvs: LLOQ, L, M, H 3 runs x 5 parallels</p> <p><b>Other runs:</b></p> <p>At L, M, and H levels QCs in duplicates Nr of QCs: 5% or 6, whichever is higher</p> <p><b>Acceptance Criteria:</b></p> <p>≥ 67% of QCs ± 15% ≥ 50% of QCs per level ± 15%</p>	<p><b>All runs (also after validation):</b></p> <p>Blank, zero Cals: 6 levels QC: ≥ 3 levels (L, M, H) x 2 parallels or ≥ 5% of study samples, whichever is higher</p> <p><b>Acceptance Criteria:</b></p> <p><b>Cals:</b> LLOQ ± 20% Other: ± 15% 75% (min. 6 levels) of Cals</p> <p><b>QCs:</b> ≥ 67% of QCs ± 15% ≥ 50% of QCs per level ± 15%</p>		<p>Every batch should have QCs, stable test samples, blanks and/or standard solutions, control charts are recommended,</p>

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<b>Dilution</b>	QCs for planned dilutions, 5 replicates per dilution factor, A & P: $\pm 15\%$ of nominal conc. or RSD	Cover the dilution applied to the study samples. Spiking matrix above the ULOQ and diluting with blank matrix ( $\geq 5$ determinations per dilution factor).  <b>A &amp; P:</b> $\pm 15\%$ of nominal conc. or RSD		
<b>Incurred Sample Reanalysis (ISR)</b>	Must reanalyze samples for control: first 1000: 10% remaining: 5%  <b>Sample selection:</b> $C_{max}$ and in the elimination phase  <b>Acceptance Criteria:</b> 67% of repeated samples within $\pm 20\%$ of the mean	Must reanalyze samples for control: first 1000: 10% remaining: 5%  <b>Sample selection:</b> $C_{max}$ and in the elimination phase  <b>Acceptance Criteria:</b> 67% of repeated samples within $\pm 20\%$ of the mean		
<b>Repeat Analysis</b>	No re-analysis of individual calibrators and QCs is permitted.  Reanalysis should be based on reasons At least the same number of replicates for repeats as originally tested	Example cases: Run did not fulfil the acceptance criterias,  IS response significantly differing from cal. or QCs response (if criteria pre-defined),  Improper sample injection or malfunction of equipment,  Obtained concentration above ULOQ or below LLOQ,  Analyte levels in blanks too high,  Poor chromatography		
<p>Parallels – Samples that have been taken through the entire measurement procedure (each has had independent sample pretreatment)</p> <p>EMA: LLOQ, L: within three times the LLOQ (low QC), M: around 30 - 50% of the calibration curve range (medium QC), H: at least at 75% of the upper calibration curve range (high QC).</p> <p>FDA: LLOQ, low (L: defined as three times the, LLOQ), mid (M: defined as mid-range), high (H: defined as high-range)</p>				

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<p><b>References:</b></p> <ol style="list-style-type: none"> <li>1. U.S. Department of Health and Human Services Food and Drug Administration, Bioanalytical Method Validation, Guidance for Industry, 2018: <a href="https://www.fda.gov/media/70858/download">https://www.fda.gov/media/70858/download</a></li> <li>2. ICH guideline M10 on bioanalytical method validation and study sample analysis. European Medicines Agency 2022. <a href="https://www.ema.europa.eu/en/documents/scientific-guideline/ich-guideline-m10-bioanalytical-method-validation-step-5_en.pdf">https://www.ema.europa.eu/en/documents/scientific-guideline/ich-guideline-m10-bioanalytical-method-validation-step-5_en.pdf</a></li> <li>3. ICH harmonized tripartite guideline: validation of analytical procedures: text and methodology Q2(R1), International Conference of harmonization of technical requirements for registration of pharmaceuticals for human use 2005: <a href="https://database.ich.org/sites/default/files/Q2_R1__Guideline.pdf">https://database.ich.org/sites/default/files/Q2_R1__Guideline.pdf</a></li> <li>4. B. Magnusson and U. Örnemark (eds.) Eurachem Guide: The Fitness for Purpose of Analytical Methods - A Laboratory Guide to Method Validation and Related Topics, (2nd ed. 2014): <a href="https://www.eurachem.org/images/stories/Guides/pdf/MV_guide_2nd_ed_EN.pdf">https://www.eurachem.org/images/stories/Guides/pdf/MV_guide_2nd_ed_EN.pdf</a></li> </ol> <p style="text-align: right;"> <a href="https://sisu.ut.ee/lcms_method_validation/">University of Tartu</a>  <a href="https://sisu.ut.ee/lcms_method_validation/">https://sisu.ut.ee/lcms_method_validation/</a> </p>				