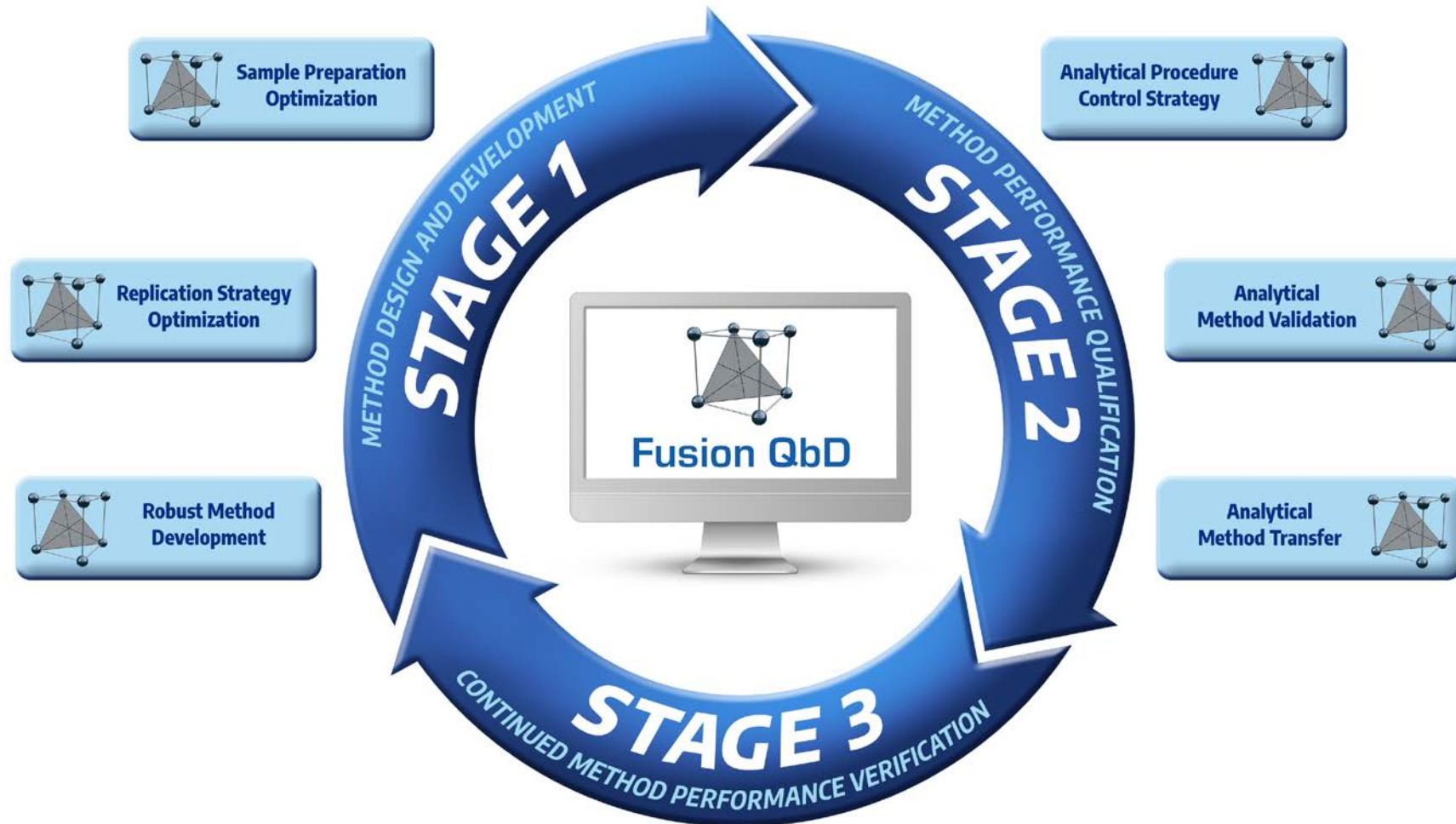


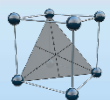
Fusion QbD[®]

Automated Method Validation & Transfer

A Complete Solution for APLM Stages 1 and 2

Analytical Procedure Lifecycle Management Workflow





INTERNATIONAL COUNCIL FOR HARMONISATION OF TECHNICAL
REQUIREMENTS FOR PHARMACEUTICALS FOR HUMAN USE

ICH HARMONISED GUIDELINE

ANALYTICAL PROCEDURE DEVELOPMENT Q14

Final Version
Adopted on 1 November 2023

This Guideline has been developed by the appropriate ICH Expert Working Group and has been subject to consultation by the regulatory parties, in accordance with the ICH Process. At Step 4 of the Process the final draft is recommended for adoption to the regulatory bodies of ICH regions.



INTERNATIONAL COUNCIL FOR HARMONISATION OF TECHNICAL
REQUIREMENTS FOR PHARMACEUTICALS FOR HUMAN USE

ICH HARMONISED GUIDELINE

VALIDATION OF ANALYTICAL PROCEDURES Q2(R2)

Final Version
Adopted on 1 November 2023

This Guideline has been developed by the appropriate ICH Expert Working Group and has been subject to consultation by the regulatory parties, in accordance with the ICH Process. At Step 4 of the Process the final draft is recommended for adoption to the regulatory bodies of ICH regions.

EUROPEAN PHARMACOPOEIA 11.6

2.2.46. Chromatographic separation techniques



01/2025:20246

2.2.46. CHROMATOGRAPHIC SEPARATION TECHNIQUES⁽¹⁾

INTRODUCTION

Chromatographic separation techniques are multi-stage separation methods in which the components of a sample are distributed between 2 phases, one of which is stationary, while the other is mobile. The stationary phase may be a solid or a liquid supported on a solid or a gel. The stationary phase may be packed in a column, spread as a layer, or distributed as a film, etc. The mobile phase may be gaseous or liquid. The separation may be based on adsorption, mass distribution (partition), ion exchange, etc., or may be based on differences in the physico-chemical properties of the molecules such as size, mass, volume, etc.

This chapter contains definitions and calculations of common parameters and generally applicable requirements for system suitability. Principles of separation, equipment and methods are given in the corresponding general chapters:

- 0- paper chromatography (2.2.26);
- thin-layer chromatography (2.2.27);
- gas chromatography (2.2.28);
- liquid chromatography (2.2.29);
- size-exclusion chromatography (2.2.30).

DEFINITIONS

The system suitability and acceptance criteria in monographs have been set using parameters as defined below. With some equipment, certain parameters such as signal-to-noise ratio

and resolution can be calculated using software provided by the manufacturer. It is the responsibility of the user to ensure that the calculation methods used in the software are equivalent to the requirements of the European Pharmacopoeia and to make any necessary corrections if this is not the case.

Chromatogram

A graphical or other representation of detector response, effluent concentration or other quantity used as a measure of effluent concentration, versus time or volume. Ideally, chromatograms are represented as a sequence of Gaussian peaks on a baseline (Figure 2.2.46-1).

V_M = hold-up volume;

t_M = hold-up time;

V_{R1} = retention volume of peak 1;

t_{R1} = retention time of peak 1;

V_{R2} = retention volume of peak 2;

t_{R2} = retention time of peak 2;

W_h = peak width at half-height;

W_i = peak width at the inflexion point;

h = height of the peak;

$h/2$ = half-height of the peak.

Distribution constant (K_d)

In size-exclusion chromatography, the elution characteristics of a component in a particular column may be given by the distribution constant (also referred to as distribution coefficient), which is calculated using the following equation:

$$K_d = \frac{t_R - t_0}{t_0 - t_M}$$

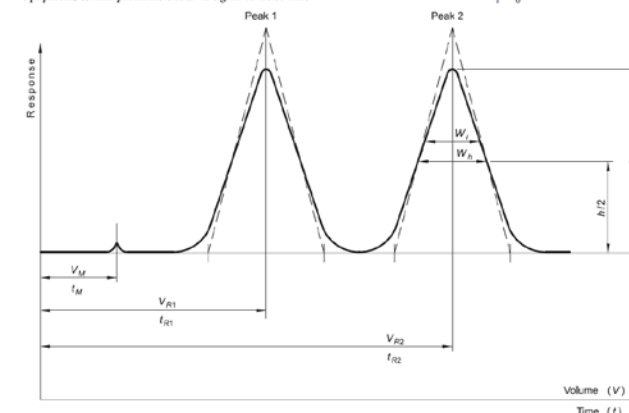


Figure 2.2.46-1.

⁽¹⁾ This chapter has undergone pharmacopoeial harmonisation. See chapter 3.8. Pharmacopoeial harmonisation.

General Notices (1) apply to all monographs and other texts

6109



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DOI: https://doi.org/10.1002/USJUPNF.M10175_02_01
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1

Add the following:

▲(1220) ANALYTICAL PROCEDURE LIFE CYCLE

INTRODUCTION

This general chapter holistically considers the validation activities that take place across the entire life cycle of an analytical procedure and provides a framework for the implementation of the life cycle approach.

The analytical procedure life cycle approach described here is consistent with the quality by design concepts described in International Council for Harmonisation (ICH) guidelines. The procedure life cycle approach emphasizes the importance of sound scientific approaches and quality risk management for the development, control, establishment, and use of analytical procedures. Total error is used in this chapter; however, measurement uncertainty can also be used.

The procedure life cycle approach is applicable to all types of analytical procedures, and the extent of effort should be consistent with the complexity of the procedure and the criticality of the quality attribute to be measured. The life cycle approach can be considered optional, and any of the elements can be applied on the basis of how the procedure is used. Elements of the life cycle approach can also be applied retrospectively if deemed useful or in early stages of development with the appropriate modifications.

Elements of life cycle management of analytical procedures are also discussed in *Analytical Procedures and Methods Validation for Drugs and Biologics* (Guidance for Industry, FDA 2015).

Validation of an analytical procedure is the process by which it is established, through laboratory studies, that the performance of the procedure meets the requirements for the intended analytical applications. Validation, or demonstration that a procedure is suitable for the intended purpose, takes place during the entire procedure life cycle, beginning during the initial procedure design activities and extending through routine use. These activities include the formal procedure validation, verification, and transfer of procedures, as well as establishing and assuring adherence to an appropriate set of procedure controls and system suitability requirements.

The procedure life cycle is comprised of the analytical target profile (ATP) and three stages, which are introduced below and shown in Figure 1.

The ATP defines the criteria for the procedure performance characteristics that are linked to the intended analytical application and the quality attribute to be measured. It applies to all stages of the procedure life cycle. For quantitative procedures, the ATP describes the required quality of the reportable value since the reportable value generated using a qualified analytical procedure provides the basis for key decisions regarding compliance of a test article with regulatory, compendial, and manufacturing limits. The acceptable level of risk of making an incorrect decision can also be considered when establishing ATP criteria.

Stage 1: Procedure design encompasses procedure development, which consists of the analytical technology and sample preparation. It includes understanding gained through knowledge gathering, systematic procedure development experiments, and risk assessments and associated lab experiments. The output of Stage 1 includes:

1. A set of procedure conditions that minimizes procedure bias to a suitable level, can provide acceptable precision, and can meet the ATP criteria
2. An understanding of the effect of procedure parameters (e.g., temperature, wavelength, flow rate, etc.) on procedure performance
3. Optimization of performance characteristics of the analytical procedure such as accuracy, precision, the appropriateness of any calibration model, specificity and limit of quantitation (as far as applicable); this includes a preliminary replication strategy for samples and standards
4. An initial analytical control strategy (ACS), which is a set of controls (system suitability tests [SSTs] and other procedure-specific controls) needed to ensure proper performance

Stage 2: Procedure performance qualification consists of studies designed to demonstrate that the procedure is suitable for its intended purpose. This involves confirmation that the reportable values generated by application of the analytical procedure meet the ATP criteria as well as confirmation of procedure performance characteristics through the traditional validation, verification, or transfer studies. Data generated during Stage 1 can be used if available and suitable. At the end of Stage 2, the replication strategy and the performance of the procedure is confirmed to meet the ATP and other criteria.

Stage 3: Ongoing procedure performance qualification involves monitoring the analytical procedure during routine use and confirming that the performance continues to meet ATP criteria. Monitoring ensures that the performance of the procedure is maintained at an acceptable level over the procedure lifetime. It can also provide an early indication of potential performance issues or adverse trends and aid in identifying required changes for the analytical procedure. Confirming procedure performance after changes ensures that the modified procedure will produce reportable values that meet the criteria defined in the ATP.

More details about the procedure life cycle are described in the subsequent sections.

https://online.uspcg.com/suppl/document/1_GUID-35D7E47E-65E5-46B7-84C0-4D06F42A30B21_2_en-US

1/12

A Complete Solution for APLM Stage 2



METHOD VALIDATION MODULE

- Full Validation Experiment Suite
- Instant Analysis and Reporting
- Advanced Method Transfer Support
- Meets all Regulatory Requirements

Critical QbD Capability

FMV

Complete Method Validation Experiment Suite



Simple Experiment Workflows



Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- Analytical Method Transfer

All the Critical QbD Capabilities You Need

Critical QbD Capability

FMV

Complete Method Validation Experiment Suite



Simple Experiment Workflows



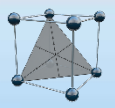
Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- Analytical Method Transfer



- Replication Strategy*
- Specificity
- Filter Validation
- Sample Solution Stability
- Accuracy*
- Linearity & Range
- Repeatability*
- Accuracy / Linearity / Repeatability*
[Combined as per ICH Q2(R1)]
- LOQ*, LOD*
- Intermediate Precision and Reproducibility
- Validation Robustness – LC
- Validation Robustness – Non-LC
[e.g. Sample Preparation, Dissolution]
- Method Transfer Study Support*

* – integration of USP <1210> Tolerance & Prediction Intervals]

All the Critical QbD Capabilities You Need

Critical QbD Capability

FMV

Complete Method Validation Experiment Suite



Simple Experiment Workflows



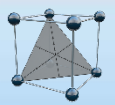
Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- Analytical Method Transfer



Example: Accuracy / Linearity / Repeatability – Combined Experiment

Create New Work File

Project

Select Project: User Defined

Project Name: Project 1

☒ Audit Logging Enabled

Instrument

Fusion QbD Demo H_Class

Instrument Type: LC
Data System: Waters Empower
Pump Module: Quaternary

Sample Compound Type

☒ Small Molecule ☐ Large Molecule

Experiment Phase

Final Phase Method Validation

Experiment Type

- Analytical Capability
- Specificity
- Accuracy
- Linearity and Range
- Repeatability
- Accuracy, Linearity, Repeatability**
- Robustness
- Robustness Non-LC

OK Cancel ?

1. Simple Experiment Setup Template

Experiment Setup

Replication Settings

Global Compound Settings

Assay Type

Assay Type Name

No. of Compounds

2

No. of Levels per Compound

5

100% Std. Level

Level 3

Compound Name	Units		Level Settings
Compound 1	%	<div>-0.00</div> <div>+0.00</div>	<div>Level 1</div> <div>80</div>
			<div>Level 2</div> <div>90</div>
			<div>Level 3</div> <div>100</div>
			<div>Level 4</div> <div>110</div>
			<div>Level 5</div> <div>120</div>

Compound Name	Units		Level Settings
Compound 2	%	<div>-0.00</div> <div>+0.00</div>	<div>Level 1</div> <div>80</div>
			<div>Level 2</div> <div>90</div>
			<div>Level 3</div> <div>100</div>
			<div>Level 4</div> <div>110</div>
			<div>Level 5</div> <div>120</div>

Create and Maintain Templates.

Set Automatic E-Review and E-Approve Loops.

2. Standards Setup Options

Standards Setup

Standards Strategy
 Calibration and Check Standards
 <None Selected>
 Bracketing - NonOverlap
 Grand Average
 Calibration and Check Standards
 Multi-level Bracketing - Overlap

No. of Repeat Injections per Level 1

Check Standards Scheme
 No. of Standards per Group 1
 No. of Injections Between Groups 5

Experiment Design

	Run No.	API	Impurity 1	Impurity 2	Degradant A	Degradant B
1	CAL - L1.1	---	---	---	---	---
2	CAL - L2.1	---	---	---	---	---
3	CAL - L3.1	---	---	---	---	---
4	CAL - L4.1	---	---	---	---	---
5	CAL - L5.1	---	---	---	---	---
6	Chk - 1.a	100	100	100	100	100
7	1.a	80	80	80	80	80
8	1.b	80	80	80	80	80
9	1.c	80	80	80	80	80
10	1.d	80	80	80	80	80
11	1.e	80	80	80	80	80
12	Chk - 1.b	100	100	100	100	100
13	2.a	90	90	90	90	90

Validation Status: Your settings are valid.

Clear Reset Next >> Cancel ?

Flexible setup of the required Standards Strategies.

3. Auto-generated Experiment Design

Analytical Method Validation - Untitled1.smae

File Edit Activity Tools Window Help

Design of Experiments

- Create a Design
- Design Reports**

Data Entry / Analysis

- Data Entry
- Data Analysis

Reporting Toolkit

- Fusion Reporter
- Audit Log Reporter

Design Reports

- Experiment Design
- Instrument Report
- Experiment Design**
- Experiment Setup

Comprehensive, automated reporting.

Name: Administrator
Company: S-Matrix
Project: User Defined
Date: August 19, 2016 1:09:19 PM PDT [GMT-07:00]

Experiment Design - Accuracy/Linearity/Repeatability

Experiment Design Matrix

Run No.	API (%)	Impurity 1 (%)	Impurity 2 (%)	Degradant A (%)	Degradant B (%)
CAL - L1.1	100	100	100	100	100
CAL - L2.1	100	100	100	100	100
CAL - L3.1	100	100	100	100	100
CAL - L4.1	100	100	100	100	100
CAL - L5.1	100	100	100	100	100
Chk - 1.a	100	100	100	100	100
1.a	80	80	80	80	80
1.b	80	80	80	80	80
1.c	80	80	80	80	80
2.a	90	90	90	90	90

Ready modified

4. Analysis Wizard for CDS Imported Results

Method Validation - Small Molecule Data Analysis

Accuracy | **Linearity** | Repeatability

Select Response for Analysis
Peak Area

API

☒ Perform Data Analysis

Compound-based Acceptance Criteria

☒ Linearity Regression ($r \geq$) 0.9998

☒ Intercept % Bias Calculation Options

☒ Data Based ☐ Model Based

☒ Intercept |% Bias| \leq 2.00

LOQ / LOD

Calculation Method(s)

☐ ICH-Q2B ☒ USP <1210>

Significance Level 5 %

LOQ Calculation(s)

☒ Residual Std. Dev. ☐ Intercept Std. Dev.

LOD Calculation(s)

☒ Residual Std. Dev. ☐ Intercept Std. Dev.

Level-based Acceptance Criteria

Computed Results

☒ Include Response Factor

Level	Linearity % Bias of Residuals \leq	Response Factor % Bias \leq
1.000	5.00	
2.000	5.00	
4.000	5.00	
5.000	5.00	

Source Data

Level	Individual Results Spec. Lower Limit	Individual Results Spec. Upper Limit
1.000	1627663	1798996
2.000	3340993	3512326
4.000	6767652	6938985
5.000	8480023	8652215

The settings are valid.

Back Finish Cancel

Associate responses with Analyses:
For Example – Amount for Accuracy and Area for Linearity.

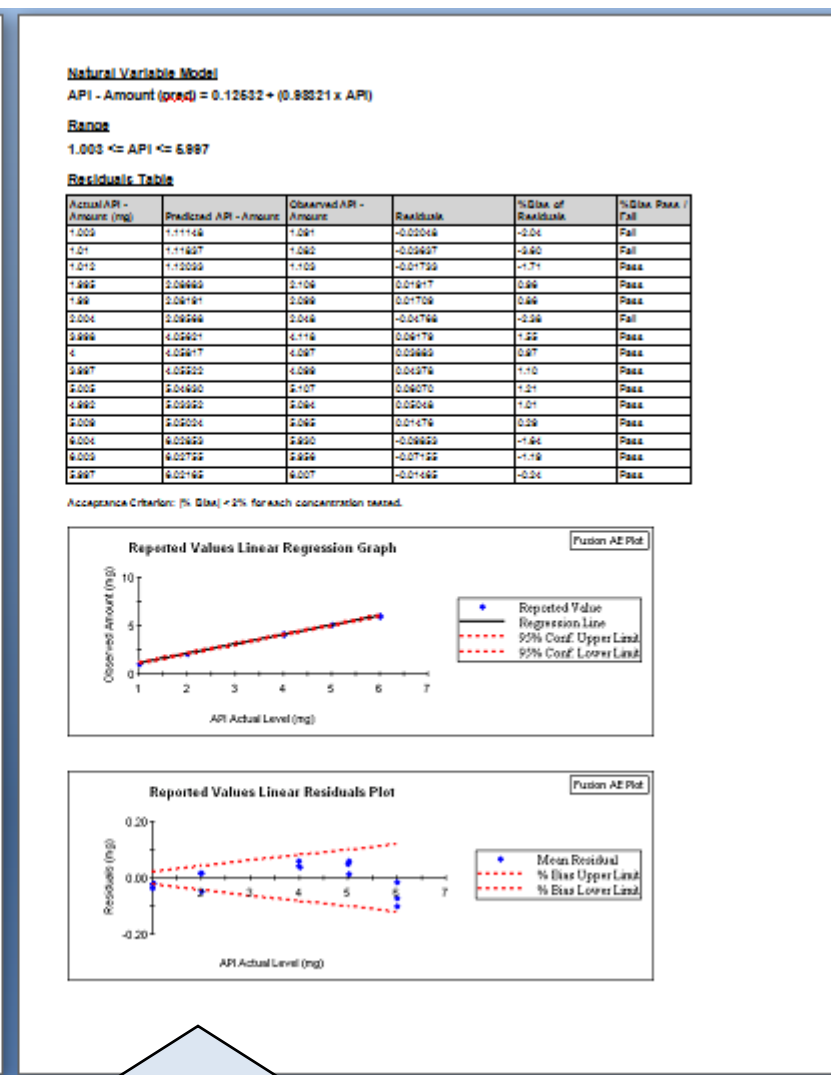
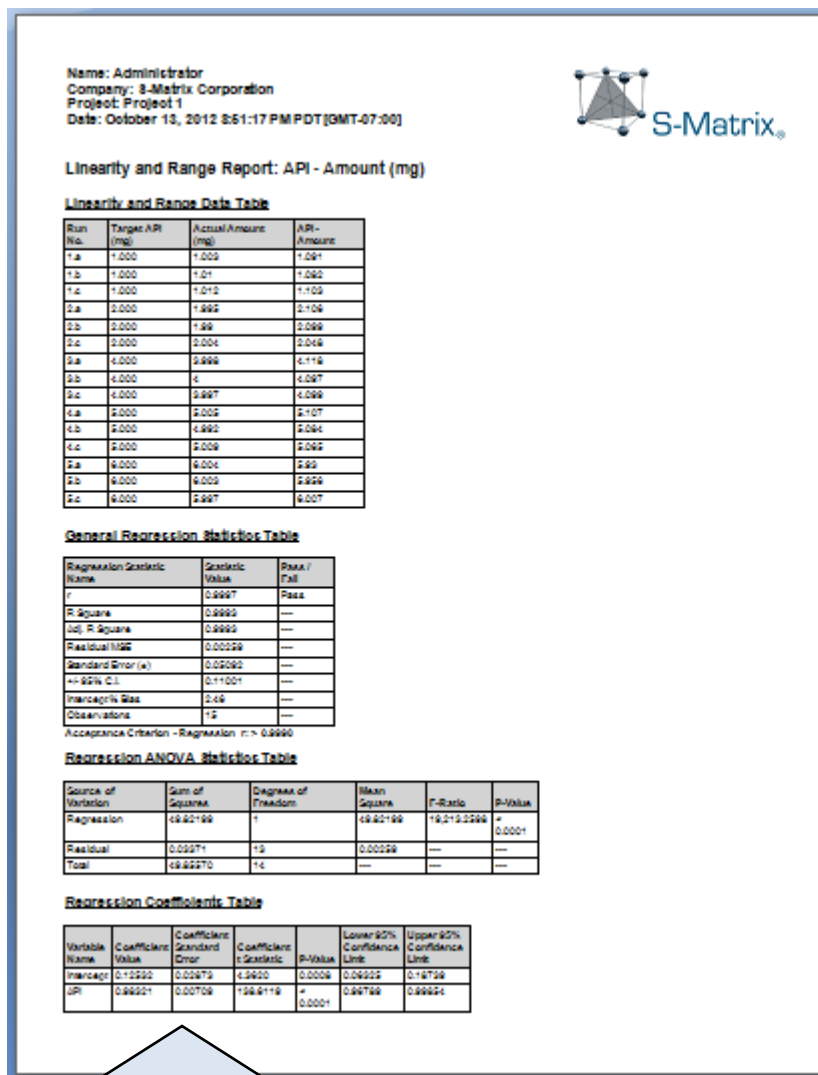
Set Global and
Level-specific
Acceptance
Criteria,
including Spec
Limits for Data

Include LOQ and LOD.
Select Calculation
Method Options.

5. Instant Analysis, Graphing, and Reporting

ICH Q2(R2):

Data derived from the regression line may help to provide mathematical estimates of the linearity. A plot of the data, the correlation coefficient or coefficient of determination, y-intercept and slope of the regression line should be provided. An analysis of the deviation of the actual data points from the regression line is helpful for evaluating linearity (e.g., for a linear response, the impact of any non-random pattern in the residuals plot from the regression analysis should be assessed).



Fusion QbD instantly creates formal reports with all required tables and graphs.

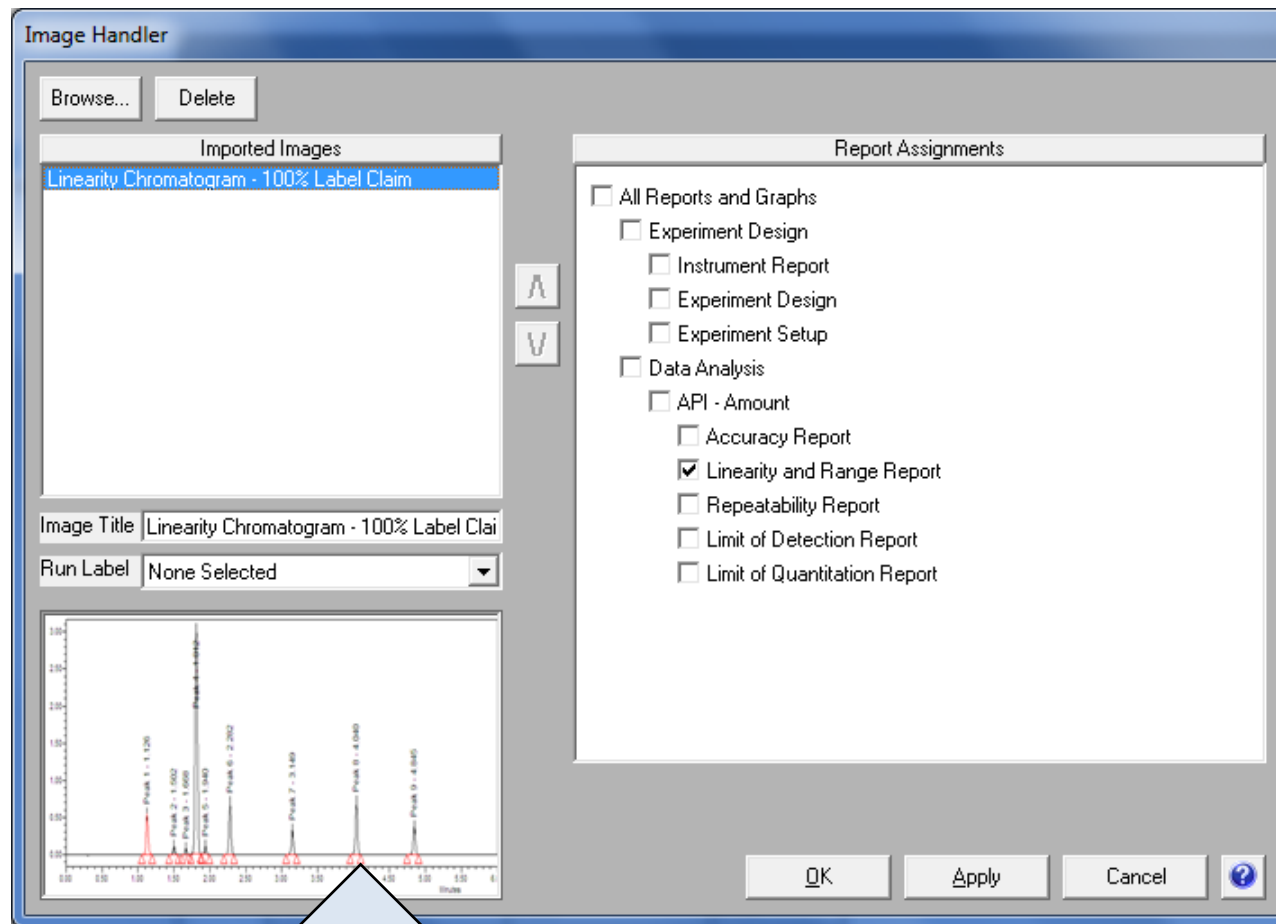


5. Instant Analysis, Graphing, and Reporting

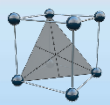
ICH Q2(R2):

Representative data (e.g., chromatograms, electropherograms, spectra, biological response) should be used to demonstrate specificity and relevant components should be labelled, if appropriate.

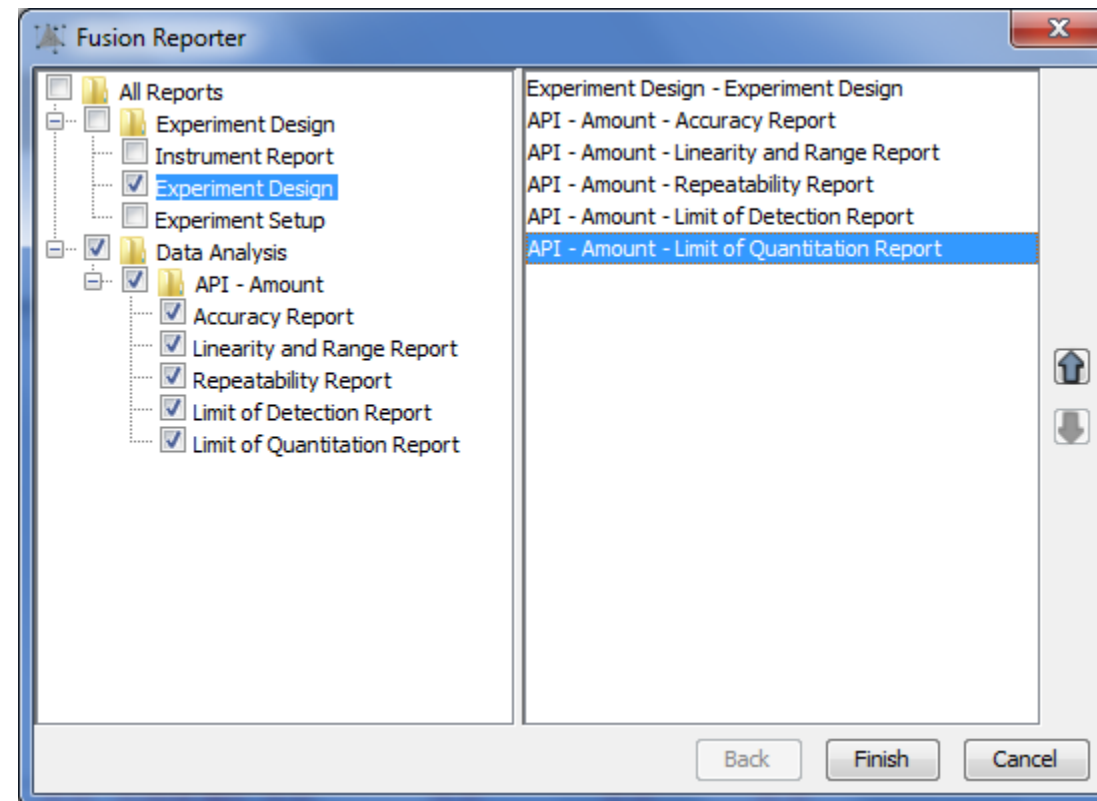
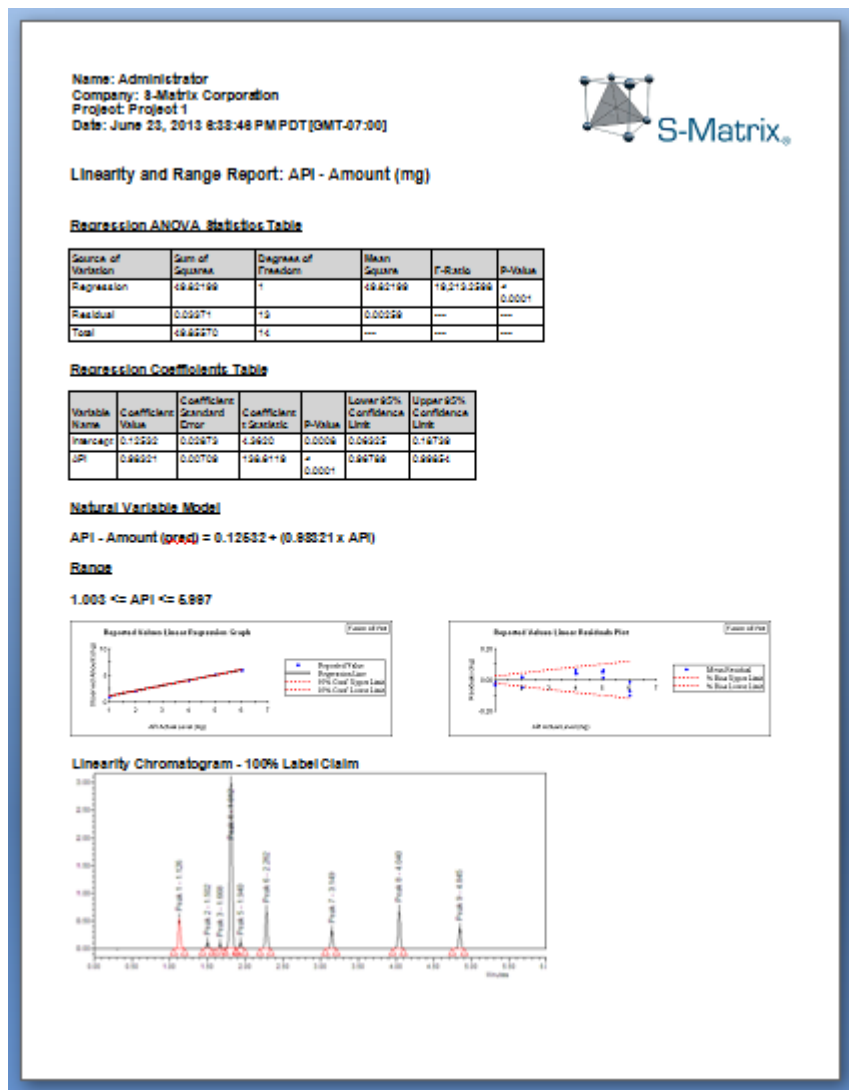
For a purity or impurity test, discrimination can be established by stressing or spiking product to achieve appropriate levels of impurities or related substances and demonstrating the absence of interference.



Reports can be augmented with images of relevant chromatograms.



5. Instant Analysis, Graphing, and Reporting



Reports meet all output format requirements:

.TXT / .RTF / .DOC / .PDF / .HTML / XLSX

All the Critical QbD Capabilities You Need

Critical QbD Capability

FMV

Complete Method Validation Experiment Suite



Simple Experiment Workflows



Full LC Experiment Automation



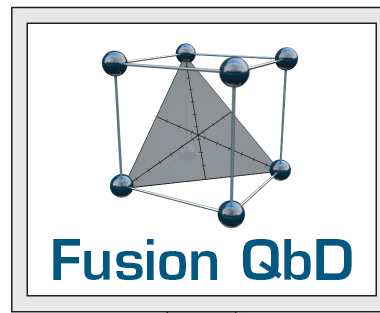
USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- Analytical Method Transfer

Automated Experiment Workflow

Steps 1 and 2



Generates Selected
Validation Experiment











Automatically Builds
Sequence with
Standards Protocol and
Assigns Method

Step 3

Chromatography Data Software (CDS)

Text Mix pH in S-Matrix - MD DemoLC Tutorial - Sample Workup as System/Administrator - Sample Set Method Editor

File Edit View Help



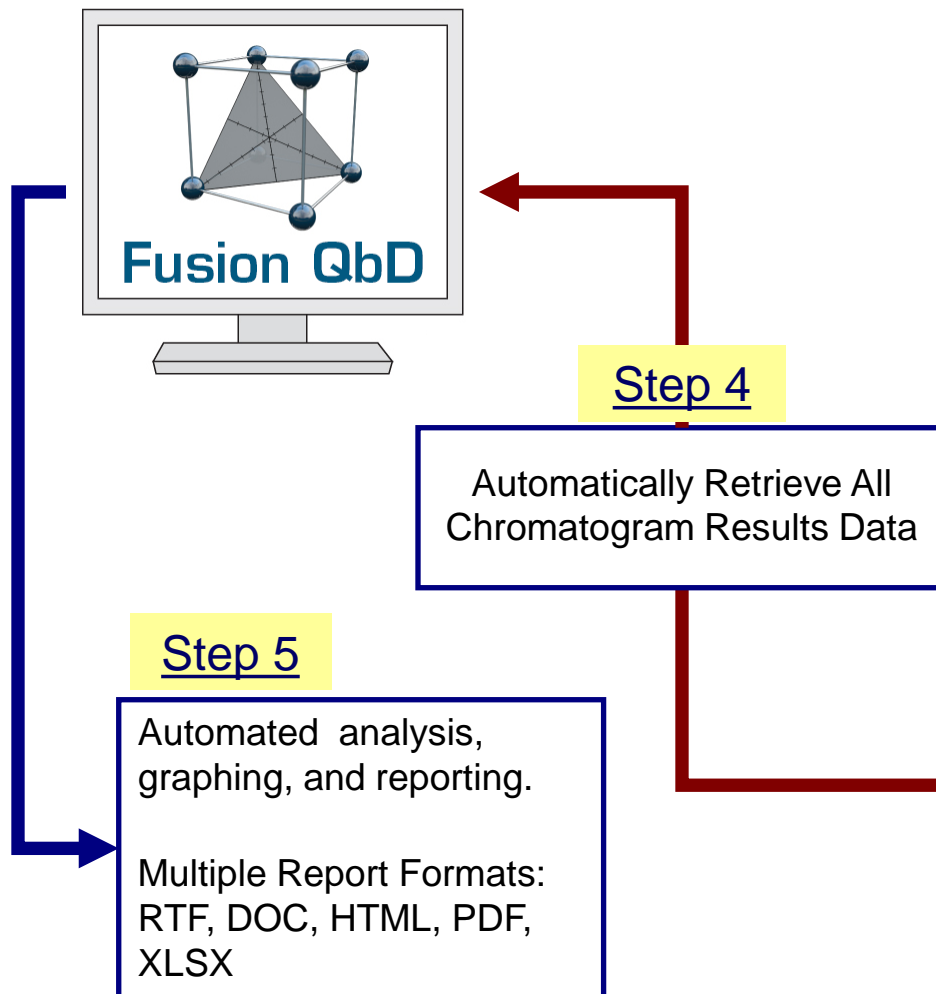
Apply Table Preferences

Sample Set Method

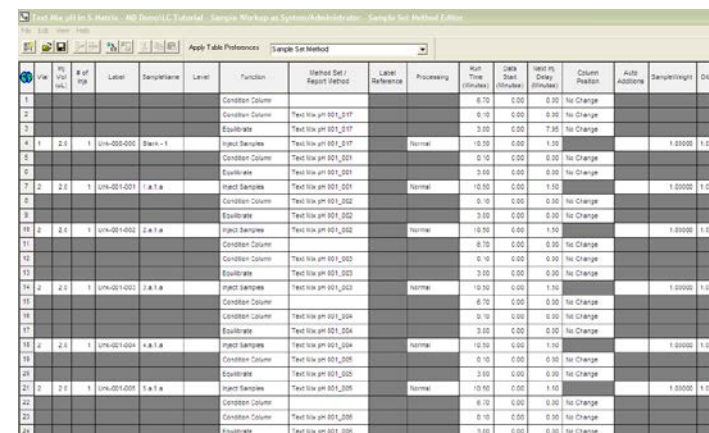
	Vial	Inj Vol (uL)	# of Injs	Label	SampleName	Level	Function	Method Set / Report Method	Label Reference	Processing	Run Time (Minutes)	Data Start (Minutes)	Next Inj Delay (Minutes)	Column Position	Auto Additions	Sample/Weight	Dilution
1							Condition Column				6.70	0.00	0.00	No Change			
2							Condition Column	Text Mix pH 001_017			0.10	0.00	0.00	No Change			
3							Equilibrate	Text Mix pH 001_017			3.00	0.00	7.95	No Change			
4	1	2.0	1	Unk-000-000	Blank - 1		Inject Samples	Text Mix pH 001_017		Normal	10.50	0.00	1.50			1.00000	1.00000
5							Condition Column	Text Mix pH 001_001			0.10	0.00	0.00	No Change			
6							Equilibrate	Text Mix pH 001_001			3.00	0.00	0.00	No Change			
7	2	2.0	1	Unk-001-001	1.a.1.a		Inject Samples	Text Mix pH 001_001		Normal	10.50	0.00	1.50			1.00000	1.00000
8							Condition Column	Text Mix pH 001_002			0.10	0.00	0.00	No Change			
9							Equilibrate	Text Mix pH 001_002			3.00	0.00	0.00	No Change			
10	2	2.0	1	Unk-001-002	2.a.1.a		Inject Samples	Text Mix pH 001_002		Normal	10.50	0.00	1.50			1.00000	1.00000
11							Condition Column				6.70	0.00	0.00	No Change			
12							Condition Column	Text Mix pH 001_003			0.10	0.00	0.00	No Change			
13							Equilibrate	Text Mix pH 001_003			3.00	0.00	0.00	No Change			
14	2	2.0	1	Unk-001-003	3.a.1.a		Inject Samples	Text Mix pH 001_003		Normal	10.50	0.00	1.50			1.00000	1.00000
15							Condition Column				6.70	0.00	0.00	No Change			
16							Condition Column	Text Mix pH 001_004			0.10	0.00	0.00	No Change			
17							Equilibrate	Text Mix pH 001_004			3.00	0.00	0.00	No Change			
18	2	2.0	1	Unk-001-004	4.a.1.a		Inject Samples	Text Mix pH 001_004		Normal	10.50	0.00	1.50			1.00000	1.00000
19							Condition Column	Text Mix pH 001_005			0.10	0.00	0.00	No Change			
20							Equilibrate	Text Mix pH 001_005			3.00	0.00	0.00	No Change			
21	2	2.0	1	Unk-001-005	5.a.1.a		Inject Samples	Text Mix pH 001_005		Normal	10.50	0.00	1.50			1.00000	1.00000
22							Condition Column				6.70	0.00	0.00	No Change			
23							Condition Column	Text Mix pH 001_006			0.10	0.00	0.00	No Change			
24							Equilibrate	Text Mix pH 001_006			3.00	0.00	0.00	No Change			

Automated, Audited Data Exchange
Preserves Data Integrity

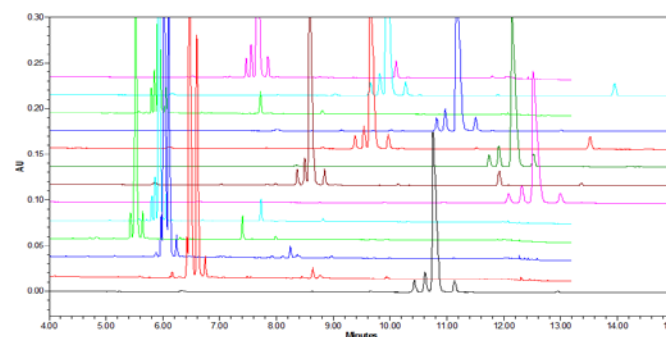
Automated Experiment Workflow



Chromatography Data Software (CDS)



Run	File	# of Pkts	Label	Sample Name	Level	Function	Method Set / Report Method	Label Reference	Processing	Run Time (minutes)	Data Size (MB)	Method ID	Column Position	Auto Addition	Sample Name	Run
1						Condition Column	Test file pH 01_017			8.70	0.00	0.00	No Change			
2						Condition Column	Test file pH 01_017			8.10	0.00	0.00	No Change			
3						Equilibrant	Test file pH 01_017			3.00	0.00	7.95	No Change			
4	1	2.0	1	UPLC-005-000	Blank - 1	Inject Samples	Test file pH 01_017		Normal	10.00	0.00	1.00			1.00000	1.00000
5						Condition Column	Test file pH 01_021			8.10	0.00	0.00	No Change			
6						Equilibrant	Test file pH 01_021			3.00	0.00	0.00	No Change			
7	2	2.0	1	UPLC-001-001	1 x 1 x	Inject Samples	Test file pH 01_021		Normal	10.00	0.00	1.00			1.00000	1.00000
8						Condition Column	Test file pH 01_022			8.10	0.00	0.00	No Change			
9						Equilibrant	Test file pH 01_022			3.00	0.00	0.00	No Change			
10	2	2.0	1	UPLC-001-002	2 x 1 x	Inject Samples	Test file pH 01_022		Normal	10.00	0.00	1.00			1.00000	1.00000
11						Condition Column	Test file pH 01_023			8.70	0.00	0.00	No Change			
12						Equilibrant	Test file pH 01_023			3.00	0.00	0.00	No Change			
13						Equilibrant	Test file pH 01_023			3.00	0.00	0.00	No Change			
14	2	2.0	1	UPLC-001-003	2 x 1 x	Inject Samples	Test file pH 01_023		Normal	10.00	0.00	1.00			1.00000	1.00000
15						Condition Column	Test file pH 01_024			8.70	0.00	0.00	No Change			
16						Equilibrant	Test file pH 01_024			3.00	0.00	0.00	No Change			
17	2	2.0	1	UPLC-001-004	4 x 1 x	Inject Samples	Test file pH 01_024		Normal	10.00	0.00	1.00			1.00000	1.00000
18						Condition Column	Test file pH 01_025			8.10	0.00	0.00	No Change			
19						Equilibrant	Test file pH 01_025			3.00	0.00	0.00	No Change			
20	2	2.0	1	UPLC-001-005	5 x 1 x	Inject Samples	Test file pH 01_025		Normal	10.00	0.00	1.00			1.00000	1.00000
21						Condition Column	Test file pH 01_026			8.70	0.00	0.00	No Change			
22						Equilibrant	Test file pH 01_026			3.00	0.00	0.00	No Change			



Automated, Audited Data Exchange
Preserves Data Integrity

Full Automation for Robustness Studies



- ✓ Solvent Selection Valves
- ✓ Column Switching Valves

Alliance HPLC



Alliance iS HPLC



Acquity Binary



Acquity H-Class



Acquity Arc



Acquity UPC²



Full Automation for Robustness Studies



OpenLab –
ChemStation
Edition



Solvent Selection Valves



Column Switching Valves

Agilent 1100s
And 1200s



Agilent 1260
Infinity Series



Agilent 1260
Infinity II Series



Agilent 1290
Infinity Series



Agilent 1290
Infinity II Series

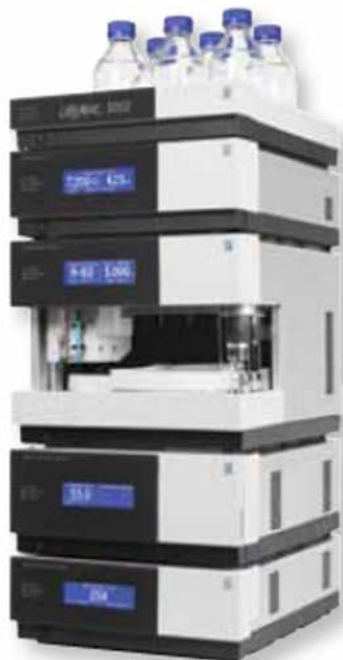


Full Automation for Robustness Studies



- ✓ Solvent Selection Valves
- ✓ Column Switching Valves

UltiMate LCs



Vanquish Horizon
And Flex LCs



Critical QbD Capability

FMV

Complete Method Validation Experiment Suite



Simple Experiment Workflows



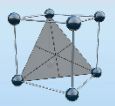
Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- Analytical Method Transfer



2. CONSIDERATIONS PRIOR TO VALIDATION

How many individual determinations will compose the reportable value, and how will they be aggregated?

- To answer this question, it is necessary to understand the contributors to the procedure variance and the ultimate purpose of the procedure.

Estimation of variance components during pre-validation provides useful information for making this decision.

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Replication Strategy Optimization



ICH Q14

Reportable Result: the result as generated by the analytical procedure after calculation or processing and applying the described sample replication. (ICH Q2). (Pg. 19)

ICH Q2(R2)

The experimental design of the validation study should reflect the number of replicates used in routine analysis to generate a reportable result. (Pg. 2)

USP <1220>

Stage 1:

Optimization of performance characteristics of the analytical procedure such as accuracy, precision, ...; this includes a preliminary replication strategy for samples and standards. (Pg. 1)

Replication Strategy Experiment

Define your Proposed Replication Strategy, Target Result Value, Acceptance Limits, Desired TAE Limits, and your Desired Probability and Tolerance (Confidence Interval).

Replication Strategy Analysis Setup

REFERENCE TABLE

Sigma	FPT	CP
1.00	322.17	0.33
2.00	44.43	0.67
3.00	2.7	1.00
4.00	0.07	1.33
5.00	0.00054	1.67
6.00	0.000002	2.00

Replication Strategy

Number of preparations x Number of injections per preparation.
Together these settings define the data which are averaged into the reportable value for the documented method.

Total Analytical Error (TAE) Limits

The \pm value = the minimum allowable \pm distance of a given reportable value from the corresponding acceptance limit.

Select a Replication Strategy

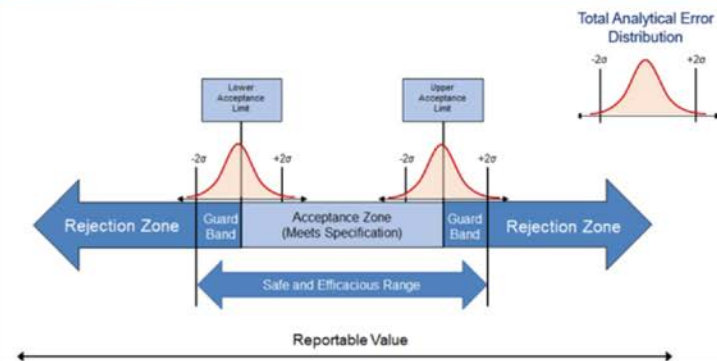
No. of preparation replicates per sample No. of injections per preparation replicate

Enabled	Responses	Target Value	\pm Acceptance Limits	\pm Total Analytical Error (TAE) Limits	TAE $\pm\sigma$ Width	Interval Type	Desired Probability (%)	Tolerance Alpha (%)
<input checked="" type="checkbox"/>	API - Amount	100.000	2.000	0.300	2σ	Tolerance	95.00	5.00

The settings are valid.

S-Matrix

Total Analytical Error and Guard Bands



The diagram illustrates the Total Analytical Error (TAE) distribution. It shows a central 'Acceptance Zone (Meets Specification)' flanked by 'Guard Bands' and 'Rejection Zones'. The 'Safe and Efficacious Range' is indicated within the acceptance zone. The 'Reportable Value' is shown as a horizontal line. The distribution is centered around the target value, with -2σ and $+2\sigma$ limits marked. A smaller inset graph shows the 'Total Analytical Error Distribution' with -2σ and $+2\sigma$ limits.

Replication Strategy for the Reportable Value

Between Variables Components of Variation

Variable Name	Variance	Standard Deviation	Degrees of Freedom	t-table Value	(+/-) 95% Confidence Limits	Error Contribution (%)
Sample Preparation	0.008	0.092	4	2.7764	0.25	94.17
Injection	0.001	0.023	20	2.0860	0.04	5.83

Overall Error in a Single Determination

Statistic	Value
Mean	100.051
Variance	0.009
Standard Deviation	0.094
% RSD	0.094

TOST Analysis Results Summary

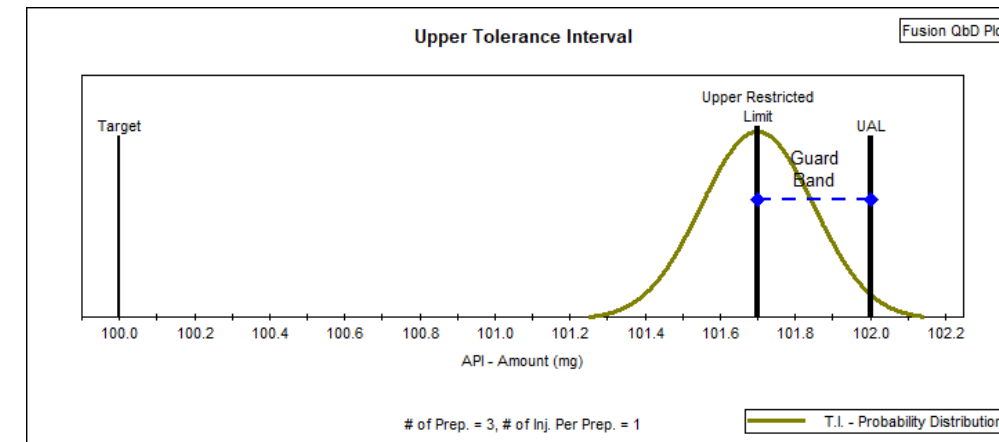
Statistic	Value	Pass/Fail
TAE Width (2σ) - Target	±0.300	
Computed TAE Width (2σ)	±0.156	Pass
FPT	<0.0001	
Cp	12.2271	
Variance	0.003	
Standard Deviation	0.055	
% RSD	0.05	
% CV	0.05	

Tolerance Interval Analysis Results

Interval Setting	Value	Number of Preparations	Number of Injections per Preparation
Target	100.000	3	1
Acceptance Limits	±2.000		
Desired Probability %	95.00		
Tolerance Alpha %	5.00		
Grand Mean	100.051		
Computed Tolerance Interval	±0.293	Pass	
Required Guard Band Width	±0.300		

The computed Tolerance Interval falls within the defined Total Analytical Error Limits.

No. of Injections	No. of Preparations										
		1	2	3	4	5	6	7	8	9	10
1	±2σ	0.2710	0.1911	0.1564	0.1355	0.1212	0.1106	0.1024	0.0958	0.0903	0.0857
	T.I.	0.6210	0.3801	0.2927	0.2455	0.2151	0.1936	0.1774	0.1647	0.1543	0.1456
2	±2σ	0.2670	0.1888	0.1541	0.1335	0.1194	0.1090	0.1009	0.0944	0.0890	0.0844
	T.I.	0.5299	0.3421	0.2698	0.2295	0.2029	0.1838	0.1693	0.1577	0.1482	0.1402
3	±2σ	0.2657	0.1878	0.1534	0.1328	0.1188	0.1085	0.1004	0.0939	0.0886	0.0840
	T.I.	0.4971	0.3288	0.2620	0.2240	0.1988	0.1806	0.1665	0.1553	0.1461	0.1384
4	±2σ	0.2650	0.1874	0.1530	0.1325	0.1185	0.1082	0.1002	0.0937	0.0883	0.0838
	T.I.	0.4801	0.3221	0.2580	0.2213	0.1968	0.1789	0.1652	0.1542	0.1451	0.1375
5	±2σ	0.2646	0.1871	0.1528	0.1323	0.1183	0.1080	0.1000	0.0935	0.0882	0.0837
	T.I.	0.4697	0.3180	0.2557	0.2197	0.1955	0.1779	0.1644	0.1535	0.1445	0.1369
6	±2σ	0.2643	0.1869	0.1526	0.1322	0.1182	0.1079	0.0999	0.0934	0.0881	0.0836
	T.I.	0.4626	0.3152	0.2541	0.2186	0.1947	0.1773	0.1638	0.1530	0.1441	0.1366
7	±2σ	0.2641	0.1868	0.1525	0.1321	0.1181	0.1078	0.0998	0.0934	0.0880	0.0835
	T.I.	0.4576	0.3133	0.2529	0.2178	0.1941	0.1768	0.1634	0.1527	0.1438	0.1363
8	±2σ	0.2640	0.1867	0.1524	0.1320	0.1181	0.1078	0.0998	0.0933	0.0880	0.0835
	T.I.	0.4537	0.3118	0.2521	0.2172	0.1937	0.1764	0.1631	0.1524	0.1436	0.1361
9	±2σ	0.2639	0.1866	0.1523	0.1319	0.1180	0.1077	0.0997	0.0933	0.0880	0.0834
	T.I.	0.4507	0.3106	0.2514	0.2167	0.1933	0.1762	0.1629	0.1522	0.1434	0.1360
10	±2σ	0.2638	0.1865	0.1523	0.1319	0.1180	0.1077	0.0997	0.0933	0.0879	0.0834
	T.I.	0.4483	0.3097	0.2509	0.2164	0.1931	0.1759	0.1627	0.1521	0.1433	0.1358



Replication Strategy for the Reportable Value

Fusion QbD reports the Components of Variation and the Corresponding % Contributions to Total Analytical Error.

Between Variables Components of Variation

Variable Name	Variance	Standard Deviation	Degrees of Freedom	t-table Value	(+/-) 95% Confidence Limits	Error Contribution (%)
Sample Preparation	0.008	0.092	4	2.7764	0.254	94.17
Injection	0.001	0.023	20	2.0860	0.046	5.83

Overall Error in a Single Determination

Statistic	Value
Mean	100.051
Variance	0.009
Standard Deviation	0.094
% RSD	0.094

Fusion QbD also reports the TOST Results (Traditional Precision Only) and the USP <1210> Interval Results (Combined Precision + Bias).

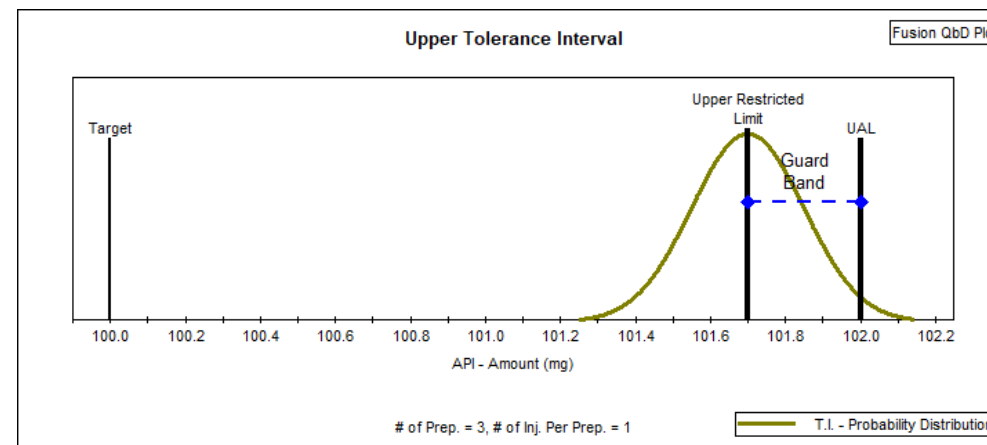
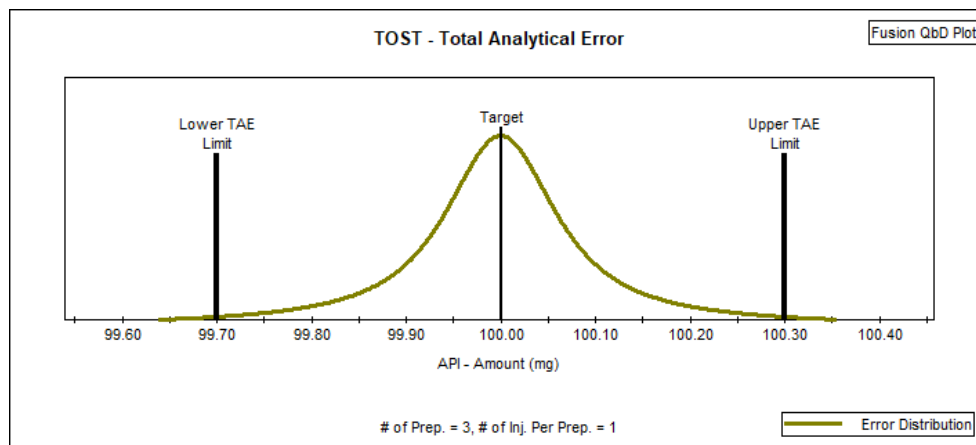
TOST Analysis Results Summary

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Computed TAE Width (2σ)	±0.156	Pass
FPT	<0.0001	
Cp	12.2271	
Variance	0.003	
Standard Deviation	0.055	
% RSD	0.05	
% CV	0.05	

Tolerance Interval Analysis Results

Interval Setting	Value	Number of Preparations	Number of Injections per Preparation
Target	100.000	3	1
Acceptance Limits	±2.000		
Desired Probability %	95.00		
Tolerance Alpha %	5.00		
Grand Mean	100.051		
Computed Tolerance Interval	±0.293	Pass	
Required Guard Band Width	±0.300		

The computed Tolerance Interval falls within the defined Total Analytical Error Limits.



Critical QbD Capability

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Full 21 CFR Part 11 Compliance Support



Complete Method Validation Experiment Suite



Simple Experiment Workflows



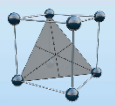
Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- **Accuracy and Repeatability**
- Analytical Method Transfer

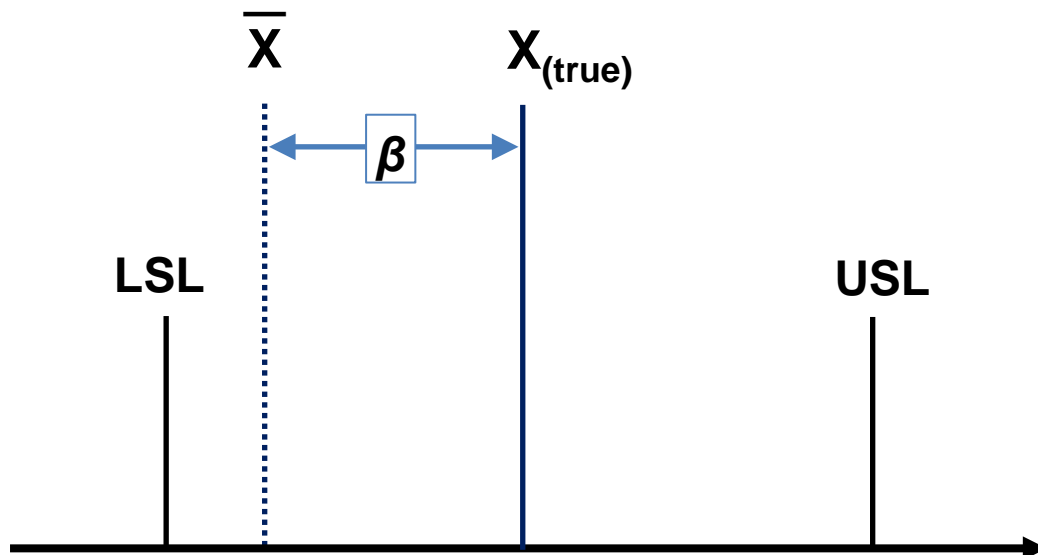


3. ACCURACY AND PRECISION

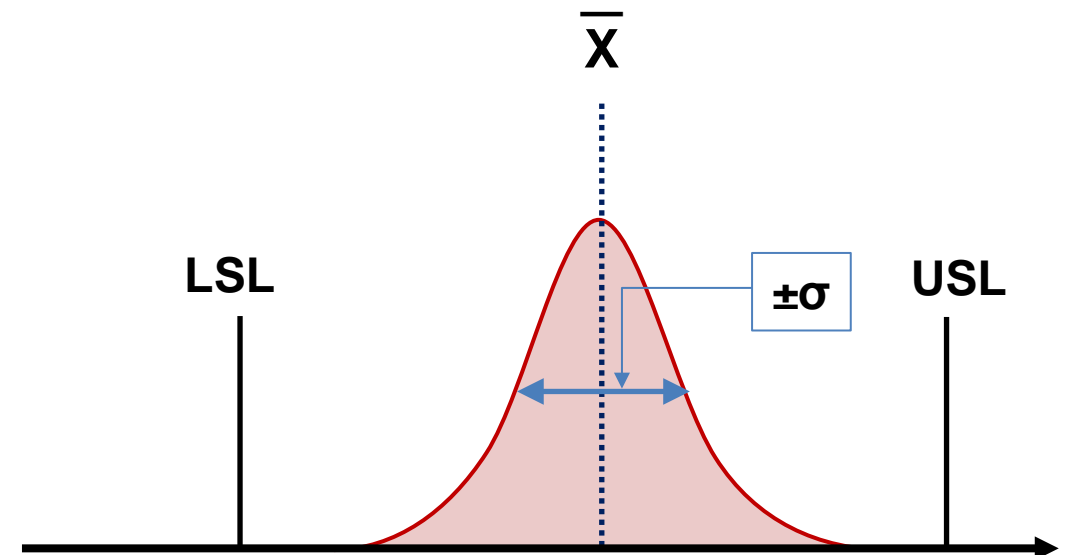
3.2 Combined Validation of Accuracy and Precision

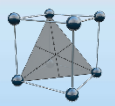
The illustration below shows that the method will pass System Suitability performance for the Critical Quality Attribute (CQA) being tested SST when Accuracy (β – bias estimate) and Precision (σ – variation estimate) are assessed independently (= High Risk Approach).

β = Bias



σ = Precision

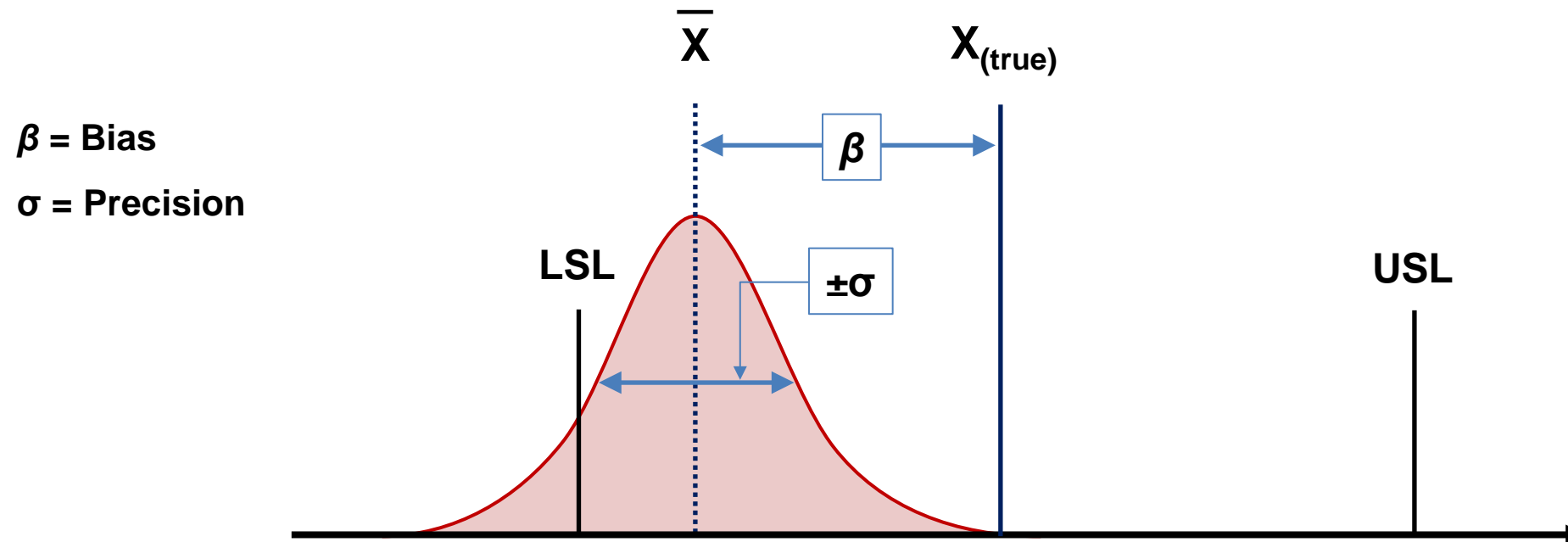




3. ACCURACY AND PRECISION

3.2 Combined Validation of Accuracy and Precision

However, as the illustration below shows – the method does not have acceptable System Suitability performance for the Critical Quality Attribute (CQA) being tested when both Accuracy (β – bias estimation) and Precision (σ – variation estimation) **are assessed together (= Low Risk Approach).**



Simple Analysis Setup Wizard

Define your Acceptance Limits:

- Compound-based – USP <1210>
- Computed Results
- Source Data

Small Molecule Data Analysis

Accuracy | Linearity | Repeatability

Select Response for Analysis
Amount

API

☒ Perform Data Analysis

Response Treatment
☐ % Recovered (Relative) ☒ Difference from Mean (Absolute)

Compound-based Acceptance Criteria
☒ Tolerance / Prediction Interval

Interval Type
☒ Tolerance ☐ Prediction

Name	Value	Unit
[Specification Limits] <=	0.20	mg
Desired Probability	95.00	%
Tolerance Alpha	5.00	%

Level-based Acceptance Criteria

Computed Results

Target Level	Accuracy [Bias (%) <=]
1.000	5.000
2.000	5.000
4.000	5.000
5.000	5.000
6.000	5.000

Source Data

Set Limit 2.000 % Apply

Mean Value	Individual Result LSL (mg)	Individual Result USL (mg)
1.055	1.034	1.076
2.084	2.043	2.126
4.105	4.023	4.187
5.085	4.984	5.187
5.964	5.845	6.084

The settings are valid.

Back Finish Cancel

Automated Reporting – all Results and Graphs for Accuracy, Linearity, Repeatability, and USP <1210> Intervals.

General Regression Statistics

Regression Statistic Name	Statistic Value
R Square	0.9999
Adj. R Square	0.9999
Residual MSE	682,072,000
Standard Error (±)	26,117
+/- 95% C.I.	56,421
Observations	15

General Validation Acceptance Criteria

Regression Statistic Name	Statistic Value	Validation	Pass / Fail
R	1.0000	0.9998	Pass
Intercept % Bias - Data Based	-0.17	2.00	Pass

Regression Coefficients

Variable Name	Coefficient Value	Coefficient Standard Error	Coefficient t Statistic	P-Value	Lower 95% Confidence Limit	Upper 95% Confidence Limit
Intercept	-11,549	14,735	< 0.0001	0.4472	-43,382	20,283
API	1,715,593	3,638	471.5873	< 0.0001	1,707,734	1,723,452

Natural Variable Model

API - Area (pred) = -11,549 + (1,715,593 x API - Weight Amount)

Range

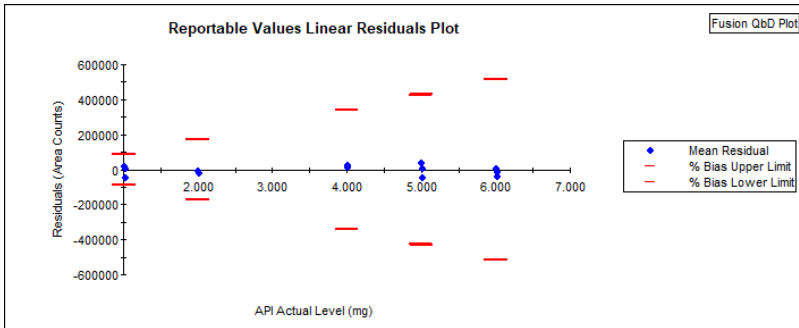
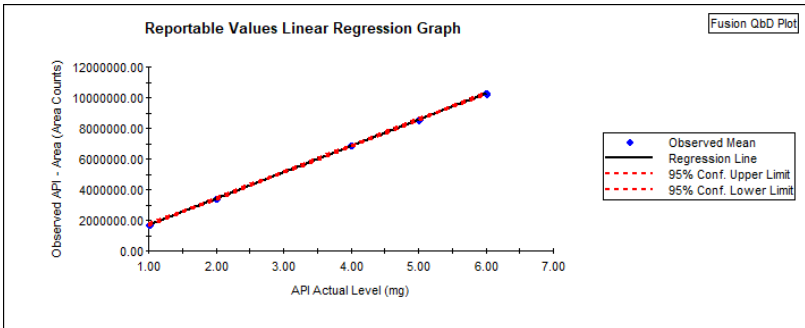
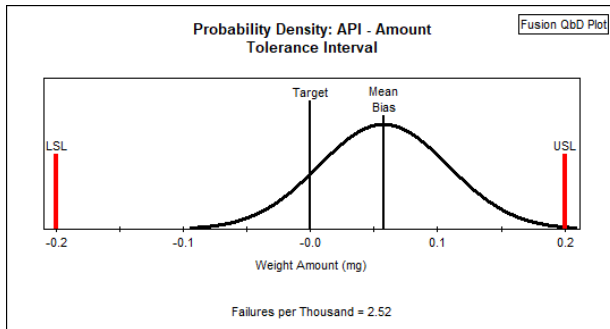
1.003 (mg) <= API - Weight Amount <= 6.004 (mg)

Accuracy Results

Target API (mg)	Mean Observed API - Amount (mg)	Standard Deviation	Lower 95% Confidence Limit	Upper 95% Confidence Limit	RSD (%)	Mean % Bias	Accuracy [% Bias] <=	% Bias Pass/Fail
1.000	1.055	0.031	0.921	1.190	2.96	4.654	5.000	Pass
2.000	2.084	0.032	1.948	2.221	1.52	4.412	5.000	Pass
4.000	4.105	0.012	4.055	4.155	0.28	2.659	5.000	Pass
5.000	5.085	0.021	4.995	5.176	0.41	1.666	5.000	Pass
6.000	5.964	0.039	5.796	6.133	0.66	-0.616	5.000	Pass

Tolerance Interval

Name	Value
Desired Probability %	95.00
Tolerance Alpha %	5.00
Target	0.00
Mean (Pooled)	0.058
Specification Limits (mg)	-0.20 <= Target <= 0.20
Computed Interval (mg)	-0.04 <= Mean <= 0.16
Result	Pass



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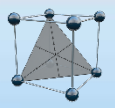
Full LC Experiment Automation



USP 1210> Tolerance and Prediction Interval Metrics



- Replication Strategy and Total Analytical Error
- Accuracy and Repeatability
- **Analytical Method Transfer**



Comparative Testing

Comparative testing requires the analysis of a predetermined number of samples of the same lot by both the sending and the receiving units. Other approaches may be valid, e.g., if the receiving unit meets a predetermined acceptance criterion for the recovery of an impurity in a spiked product. Such analysis is based on a preapproved transfer protocol that stipulates the details of the procedure, the samples that will be used, **and the predetermined acceptance criteria, including acceptable variability**. Meeting the predetermined acceptance criteria is necessary to assure that the receiving unit is qualified to run the procedure.

Analytical Method Transfer Example

Transferring Lab



Fusion QbD
Sequence
Execution

Chromatography
Data Software

ALR
Design



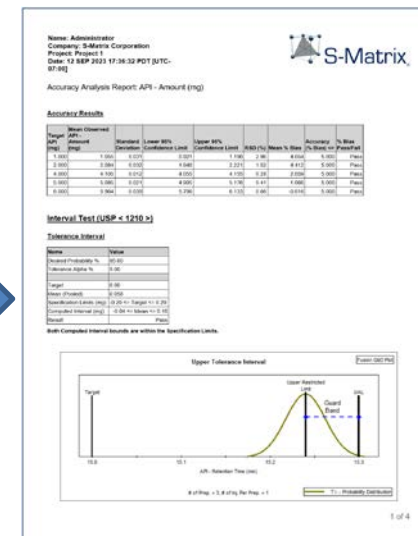
Fusion QbD
Sequence
Execution

Chromatogram
Results Data

Receiving Lab



1. Fusion QbD – Exports experiment to the CDS as Ready-to-Run sequence, methods, standards
2. Sequence is run at both labs.
3. Fusion QbD – Imports results for instant and complete analysis and reporting.



Accuracy
Linearity
Repeatability
Tolerance Interval
Pass/Fail Results

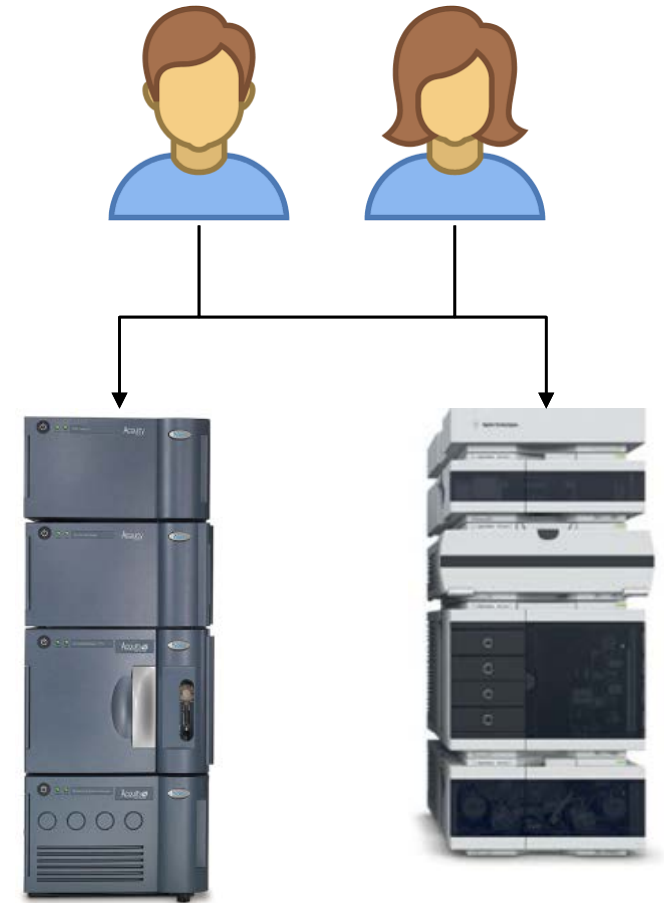
Analytical Method Transfer

Automation Makes it Easy to Extend the Analysis to Address Bias Concerns:

- **Analyst**
- **Equipment**
- **Day**
- **Etc.**

For example, each analyst could run the sequence on **each LC on each Day.**

Each results set could then be imported into Fusion QbD for direct comparison analysis.



Key Benefits of FMV

1. Consistency – Workflow and Reporting.

Work is standardized – done the same way every time. Reporting is standardized, complete, easy to communicate.

2. Simplicity

Tremendous ease of use. Very brief learning curve. Clearly defined templatable workflows with built-in workflow management.

3. Speed (Productivity)

Automation and simplified workflows dramatically increase productivity. Review process is minimized and simplified.

4. Regulatory Alignment and Completeness

All required validation experiment types are supported. Reporting meets regulatory requirements. Reports can be attached to Project specific narrative documents.

5. Platform Independence

Support for Empower, ChemStation, and Chromeleon means that the standardized workflows and reporting can be easily extended to users of other platforms at other sites or other companies (e.g. CMOs).

6. Customer Support

Our support is top-rated worldwide. S-Matrix and our local distributors have a multi-year history of proven ability to meet all our customer's support needs.

End of Presentation

Analytical Procedure Lifecycle Management Workflow

