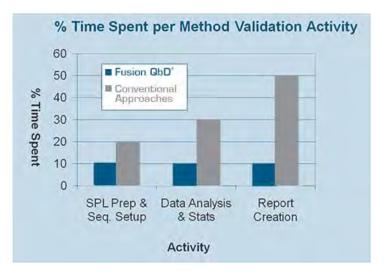
Analytical Method Validation



The Only Software That Has It All!

- Calculations and reporting meet all current FDA/ICH/USP validation guidances including the new USP <1210>!
- Can be used for LC and Non-LC methods (e.g. GC, CE, Q-NMR)!
- Automates LC method validation experiments on multiple instruments and CDS systems!
- Regulatory accepted validation for both Small & Large Molecules!
- Statistically rigorous and defensible robustness testing!
- Handles multiple compounds creates complete reports for each!
- Shortens your LC method validation time by as much as 75%!



Automated Experimentation for LC Method Validation

The objective of Method Validation is to provide documented evidence and a high degree of assurance that an analytical method employed for a specific test is suitable for its intended use. Method Validation is a regulatory requirement as much as a scientific necessity.

Key Benefits

- Full Automation for LC Method Validation multiple LCs and CDS systems
- > Aligned with FDA and ICH guidances
- 21 CFR 11 compliance support toolset Including E-records and E-signatures, full audit logging Workflow management system with E-review and E-approve loops
- Easy setup of experiments –
 Create standardized workflow templates
 Facilitate rigorous practice and defensibility
- Simple documentation review and reporting –
 Easy to defend and communicate
 Reports meet all FDA and ICH guidelines

Method Validation Experiment Suite

- Analytical Capability and System Suitability
- Specificity
- Filter Validation
- Sample Solution Stability (stability for a given time period under prescribed conditions)
- Accuracy
- Linearity and Range
- Repeatability (intra-assay precision)
- Accuracy/Linearity and Range/Repeatability Combined Design
- [ICH-Q2(R1) Accuracy, Linearity, and Repeatability can be done together as a single combined experiment]
- LOQ, LOD
- Intermediate Precision and Reproducibility (USP Ruggedness)
- Robustness done the right way!

Non-LC Method Validation Experiments

Used successfully for Non-LC methods such as GC, CE, Q-NMR, as well as hyphenated methods (e.g. LC-MS). Accepted in customer regulatory submittals.

Automated LC Method Validation – Five Step Workflow

- 1. You complete a simple experiment setup template.
- 2. Fusion QbD creates the Validation Experimental Design and exports it to the CDS.
- 3. The CDS runs the validation experiment sequence.
- 4. Fusion QbD imports and analyzes the results.
- 5. Fusion QbD automatically creates final reports and graphs.

Example Workflow - Combined Accuracy / Linearity / Repeatability

Step 1 – You Complete the Simple Template

Fusion LC Method Validation Software (FMV) has simple experiment setup templates for each type of validation experiment. The simple Linearity and Range template is shown below with user definable settings:

User-definable Settings - Basic Setup

- No. of Compounds
- No. of Levels per Compound
- 100% Standard Level
- Compound Name, Units, and Levels

User-definable Settings – Standards Setup

FMV has a flexible Standards Setup wizard which enables you to select your desired standards strategy for results quantitation within the CDS:

- Bracketing Overlap
- Bracketing Non-overlap
- Grand Average
- Calibration and Check Standards
- Multi-level Bracketing Overlap

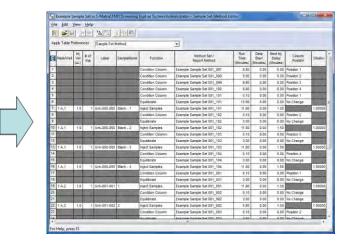
| Assay Type No. of Com | Potency | | |
|-----------------------|---------------|----------------|----|
| No. of Com | | | |
| | pounds 3 💌 | | |
| No. of Levels per Cor | npound 5 👻 | | |
| | | | |
| 100% Std. Le | vel Level 3 💌 | | |
| | | | |
| Compound Nama | Linite | Lovel Settings | |
| Compound Name | Units | Level Settings | |
| Compound Name API | Units | Level Settings | 80 |
| | | 0 +.0 | 80 |
| | | Level 1 | |
| | | Level 2 | 90 |

| one S scketi and A bielo Jti-leve | on and Chec relected ng - NonOve verage on and targe el Bracketing of Repeat Ine | rlap Stan I - O ve | derds. | <u>.</u> | Standards Scheme No. ni Standards per Group 1 💌 Intections: Between Groups 3 👻 |
|---|--|--------------------------|--------------------|------------|--|
| Expe | iment Design | | | | |
| | Run No. | API | Impurity A | Impurity B | <u>+</u> |
| 1 | CAL-L1.1 | | 100 | E. | |
| 2 | CAL-L21 | | | - | |
| 3 | CHK-1a | 100 | 100 | 100 B0 | |
| 4 | 1.8 | 80 | 80 | 80 | |
| 5 | 1.0 | 80 | 80 | 80 | |
| 7 | | 100 | 100 | 100 | |
| 8 | 2.0 | 90 | 90 | 90 | |
| 9 | 26 | 90 | 90 | 30 | |
| 10 | 2.0 | 90 | 90 | 90 | |
| 41 | CHK-1cc | 100 | 100 | 100 | 1 |
| | | | | | |
| | tion Status Y | OUX SE | ettings are valid. | | |
| alida | | | | | |

Step 2 – Fusion QbD Creates the Validation Experimental Design and Exports it to the CDS

FMV automatically constructs the validation experiment designs within the CDS as ready-to-run sequences/sample with the proper Vial No. and Injection Type designations for Samples, Standards, and Blanks.

| Project: F | : S-Matr | ix Corporation | on 0T [UTC-07:00] |
|------------|----------|----------------|----------------------|
| | | n Matrix | eriment 1 |
| Run No. | (mg) | (%) | (%) |
| CAL - 11.1 | | - | |
| CAL - L2.1 | | - | - |
| CHK - 1.a | 100 | 100 | 100 |
| 1.a | 60 | 60 | 80 |
| 1.b | 80 | 80 | BQ |
| 1.c | 60 | 80 | 80 |
| CHK - 1.b | 100 | 100 | 100 |
| 2.8 | 90 | 90 | 90 |
| 2.b | 90 | 90 | 90 |
| 2.c | 90 | 90 | 90 |
| CHK-1.c | 100 | 100 | 100 |
| 3.a | 100 | 100 | 100 |
| 3.b | 100 | 100 | 100 |
| 3.c | 100 | 100 | 100 |
| | | | 100 |



Step 3 - CDS runs the Validation Experiment

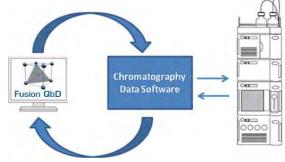
FMV sequences run automatically on the CDS. FMV even enables you to include a Shutdown method as the last method run so that you can execute FMV sequences overnight while you sleep!

Step 4 - Fusion QbD Imports and Analyzes the Chromatogram Results

FMV automatically imports the required peak result data from the CDS, and re-maps the results to the design for automated analysis, graphing, and reporting. This is a key feature ensuring quality, as manual transcription is a common source of error and risk.

| Hestbo | nse Name | Rea | iponse Units | | | |
|--------|----------|------------|--------------|--------------|--------------|---|
| Amou | ri | 100 | | - | | |
| | Run | AFITTarget | API 1 Actual | API 2 Target | API 2 Actual | * |
| 1 | La | 1.000 | 1.003 | 0.2500 | | - |
| 2 | 1.b | 1.000 | 1,01 | 0.2900 | | |
| 5 | 1.c | 1.000 | 5,012 | 0.2500 | | |
| - 4 | 24 | 2,000 | 1.995 | 0.3500 | | |
| 5 | 2.6 | 2,900 | 1 99 | 0.3500 | | |
| 6 | Zc | 2,000 | 2.004 | 0.3500 | | |
| 7. | 3.a | 4,000 | 3.998 | 0.3600 | | |
| 8 | 3.6 | 4.000 | 4.002 | 0.3600 | | |
| 3 | 30 | 4,000 | 3.997 | 0.3600 | | |
| | A.a. | 5 000 | 5.005 | 0.4000 | | |
| | 46 | 5.000 | 4 992 | C 4000 | | |
| | 4c | 5 000 | 5.009 | 0.4000 | | |
| | 5.4 | 6.000 | 6,004 | 0.4500 | | |
| 14 | 5.0 | 6,000 | 6.003 | 0.4500 | | |
| 15 | 5c | E 000 | 5.997 | 0.4500 | | |

| | | 54nm@1.2nm | | | 7 | | |
|--------|---------|----------------------|---------|-----------------|----------|--|------------------|
| 1 Im | port C | Chromatogram Trace D |)ata | | | | |
| | | | | | | | |
| rend F | Respo | inses | | | | - Named Compounds in CDS | |
| A | bbb | (i)/dete | Undo Ch | anges Restore I | Defaults | Available | Included API |
| | Γ | Operator | Value | Response | - | A B C D E F | Impurity A |
| 1 | 1.00 | No. 5' Field | - | | | D | Impany s |
| 2 | 100 | die of Frank | 100 | | - | F | 1 |
| 3 | 112 | He is then a | | | - | | |
| 4 | 12 | In a famous of | 10.00 | - | • | 1 | |
| - | 1 | | - | - | - | Response Data | |
| 5 | - | | • | | - | V Show All CDS Responses | |
| | | | | | | Available 2ndDerivativeApex | Included Area |
| | | | | | | 2Sigma 3Sigma | Amount |
| | | | | | | 4Sigma | ×1 |
| | | | | | | 5Sigma AboveldentificationThreshold | |
| • | T | | | | | AboveMaximumThreshold AboveQualificationThreshold | |
| Sele | ect All | Select None | | | omplete | AboveReportingThreshold | |
| - | - | | | D = Du | plicate | AsymAt10 * | 1 |



Flexible Data Analysis Setup Wizard

- Associate different responses with different analyses e.g.
 - Associate **Amount** results data with analysis of Accuracy
 - Associate Area results data with analysis of Linearity
- Include LOD and LOQ and select Calculation Method(s)
- Set Global and Level-specific Acceptance Criteria
- Include Level-specific Spec Limits for Raw Data

| imount | × | | | | | | | |
|----------------|-------------------------------------|------------------------------|-------------|--------------------------------------|--------------------------|---|-------------------------|--|
| Peak Area | | | | | | | | |
| API | | | | | | | | |
| Perform Data | a Analysis | | | | | | | |
| Intercept % B | as Calculation Options | | | LOQ / | | | | |
| Data Bases | d O Model Based | | | late LOD | Calculate L | | | |
| Compound Bas | ed Criteria | | | | ise Residual Standard De | viation Use Re | esidual Standard Deviat | |
| | Regression r >=) 0.9998 | 758 238 | | Use Intercept Standard Deviation | | | | |
| | | 100.01 (2001) | | | | | | |
| [Intercept | t % Blas <= 2.00 | 1.68 1.68 | | | | | | |
| | | | | | | | | |
| Level Based Cr | iteria | | | | | | | |
| -Computed Re | sults | | Source Data | | | | | |
| Include | Response Factor | | | | | | | |
| | 748 248 | 12.12 | | | | | | |
| Level | Linearity % Bias of Residuals <= | Response Factor % Blas <= | Level | Individual Result Spec. Lower Lim | it . | Individual Results Spec. Upper Limit | | |
| | 5.00 | | 1.000 | | 1627663 | | 1798996 3512326 | |
| 1.000 | | | 4.000 | - | 6767652 | | 6938985 | |
| 2.000 | 5.00 | | 5.000 | | 8480982 | | 8652315 | |
| | 5.00 | | 3.000 | | | 1 | 0002023 | |

Step 5 – Fusion QbD Automatically Creates Final Reports and Graphs

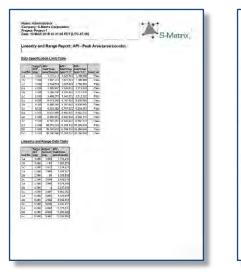
ICH Q2(R1). LINEARITY

... If there is a linear relationship, test results should be evaluated by appropriate statistical methods, for example, by calculation of a regression line by the method of least squares...

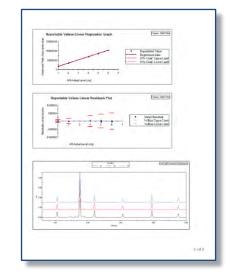
The correlation coefficient, y-intercept, slope of the regression line, and residual sum of squares should be submitted. A plot of the data should be included...:

- Correlation Coefficient
- Y Intercept
- Slope of the Regression Line

- Residual Sum of Squares
- Linear Regression Plot
- Residuals Data Table and Plot



<section-header>



FMV also enables you to include images of representative chromatograms into your final reports.

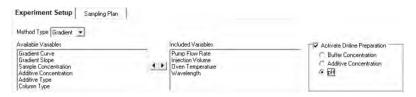
ICH Q2(R1):

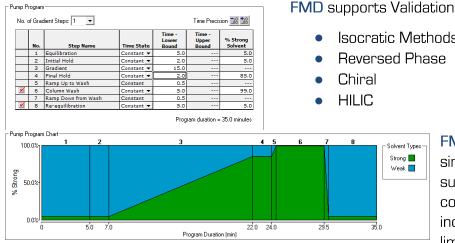
For chromatographic procedures, representative chromatograms should be used to demonstrate specificity, and individual components should be appropriately labeled...

Robustness Validation – DONE RIGHT!

Experiment Setup – LC Robustness Example

You select the parameters to include in the FMV robustness experiment. FMV will automatically generate the robustness design, re-construct it in the CDS as ready-to-run methods and sequence, import the chromatogram results directly from the CDS, re-map them to the robustness study, and instantly analyze, graph, and report the results.





FMD supports Validation Robustness studies for:

- Isocratic Methods
- **Gradient Methods**
- Normal Phase
- Ion Exchange
- Size Exclusion

FMD provides visual displays to simplify setup for complex settings such as required pump program conditions and key settings for each included column such as pH upper limit and conditioning time.

The FMV Difference Lowers your Field Failure Risk

FMV robustness experiments let you use valid experiment ranges for accurate, defensible estimates of parameter effects.

This avoids the risks associated with setting ranges equal to the expected variation ranges of your instrument parameters.

FMV robustness analysis wizard lets you set:

- expected parameter variation ranges •
- acceptable performance limits for each • key response

The wizard then accurately determines and reports the method's true robustness.

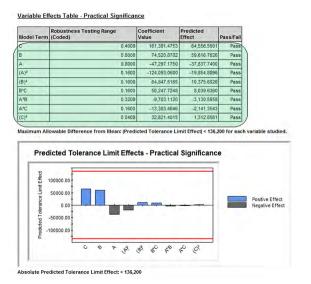
| Maximum Expected Variation: The ±3 gingma value defines the "total" variation experiment variable) around its defined setpoint occur on Nanster and normal use due to statistic | that is expected | to (a30 Venil) | 14,000 |
|--|------------------|--|--------|
| Experiment Variable | Units | Maximum Expected Variation (±3 Sigma Value) | - |
| Pump Flow Rate | mL/min | | |
| Final % Strong Solvent | % | 2 | |
| Oven Temperature | °C | 2 | |
| < Contraction of the second se | | : Back: Next>> Finish Cano | |

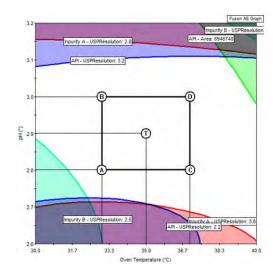
| The Maximu differences response) I he respons evaluated, normal use | llowable Difference from Mean: um Allowable Difference limit values from the mean for a given critical que beyond which the response values is to be considered robust in terms of the variation in the response measus must be encompassed by the Maxin imit values. | ality attribute unacceptable. For if the parameters rements obtained in | Tomore in writing block will be drawed based based of drawed based of the drawed of the draw | - |
|--|--|--|--|---|
| Enabled | Response | Maximum Allow | able Difference from Mean | - |
| v | API - USPResolution | 0.5 | (1.1.000) | |
| V | API - Peak Retention Time | 0.1 | | |
| Select, | All Select None | | | 1 |

Robustness Validation - Statistical Significance Testing - Model Coefficients

Robustness Report: API - Area (*) Coded Variable Name Key Coded Variable Name Actual Variable Name Initial % Organic Predicted Tolerance Limit Effects - Statistical Significance Oven Temperature pH Variable Effects Table - Statistical Significance Testing inge ded) Effect t statistic Erro Effect 4.640 64,556.5 13,911. 74,520.878 59,616.7 13,794.16 4.32 ict < 2.7622 -47 297 1754 -37,837.74 14,136,945 -2.67 0.160 124.093.06 19.854.89 14,136.945 (B)² 0.1600 64,847.5165 10,375.60 13,794.1618 0.752 Pat 0.1600 50,247,7248 8,039.64 13,714.4961 0.586 Pas -9,783.1120 -3,130.60 13,874.0259 -0.225 A*B 0.3200 Pas A*C 0.1600 -13.383.4646 -2,141.35 14,022.6463 -0.152 Pas 0.0400 32,821.4015 1,312.86 13,911.0838 0.0944 Pas able Value: |Predicted Tolerance Limit t statistic| < 2.2622 for each variable studied Maximum All

Robustness Validation - Practical Significance Testing - Effects Magnitude



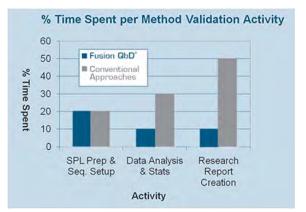


Automated LC Method Validation – Proven ROI

International Pharma Co. Benchmarking Project

Realized Time Savings = 85%.

Using historical records* and adjusting for project complexity



Copyright (©) 2019 S-Matrix Corporation, All Rights Reserved.

Minimum Expected Time Savings = 60%.

S-Matrix Software Products and Support

S-Matrix Corporation develops advanced Design of Experiment based-software that automates R&D experimental work according to Quality-by-Design principles and methodologies. S-Matrix's Fusion QbD platform automates and redefines experimentation in Analytical R&D, Chemical and Process R&D, Formulation, and Product R&D.

Fusion QbD Software System Product Suite

Fusion LC Method Development

Fully automated QbD experimenting on your LC system, integrated DOE, automated robustness simulation & chromatography data modeling. Chemistry screening without the need for peak tracking.

Fusion Analytical Method Validation

Meet regulatory guidelines with a best-practices approach toward LC method validation with comprehensive reporting. Also supports formal validation of Non-LC methods (e.g. GC, CE, Q-NMR).

Fusion Inhaler Testing

Create sampling plans, export and import data from your CDS via validated data exchange, calculate particle size distribution results, and generate reports according to USP 601, Ph.Eur. 2.9.18, and ISO 27427.

Fusion Product Development

The perfect QbD software for formulation & product development – automated experimental design selection, sophisticated analysis tools, including automated modeling and simulation, comprehensive reporting, with a full 21 CFR 11 compliance toolset.

Sales and Support

Sales: Tel: 800-336-8428 (Outside the USA: 707-441-0406). Email: <u>Sales@smatrix.com</u> Customer Support: Tel: 707-441-0407. Fax: 707-441-0410. Email: <u>Support@smatrix.com</u>

On-site and Web Training

S-Matrix offers on-site training programs for installed systems. Training includes experiment strategies, experimental design (DOE), data analysis, graphical visualization and ranking of effects, numerical and graphical optimization, and QbD Reporting.

S-Matrix also offers interactive web training which covers software features and operation, along with general principles of DOE and QbD. Web training programs can be tailored to suit your individual focus and information requirements.

To arrange an on-site or web-based training program, call 707-441-0406.

All trademarks are the property of their respective owners

S-Matrix Corporation 1594 Myrtle Avenue Eureka, CA 95501 USA www.smatrix.com