



Fusion QbD

***Powerful, Flexible, Automated
Design of Experiments***

Risk Assessment



ICH Q14

Risk assessment and prior knowledge should be used to identify parameters, attributes and appropriate associated ranges to be investigated experimentally.

(Pg. 5)

USP <1220>

For variables where there may be higher risk, one way to reduce risk is to gain additional knowledge about the influence of those parameters using modeling and/or experimentation. (Pg. 8)



Risk Assessment – Parameter Selection

Sources of Risk for Bias and/or Variation in Current Method

Element	Presumed CMPs*	CMAs						Category (C, N, X)
		Resolution USP	S/N	Tailing USP	Area % RSD - API	K-Prime - 1st Peak	K-Prime - Last Peak	
Chemistry	Column Type	5	1	1	3	5	5	X-S
	Strong Solvent	5	1	1	3	5	5	X-S
	Aqueous solvent	5	5	5	1	5	5	X-S
	pH	5	5	5	3	5	5	X-S-O
Process	Pump Flow Rate	3	1	5	3	5	5	X-O
	Injection Volume	3	5	3	5	1	1	C
	Oven Temperature	5	1	3	3	5	5	X-O
Gradient Program	Initial Hold Time	1	1	1	1	5	1	C or X-O
	Gradient Slope	5	1	5	3	5	5	X-S-O
Detection	Wavelength	5	5	1	5	1	1	C
	Sampling Rate	3	5	1	5	1	1	C
	Precision	1	3	1	3	1	1	C

C = Controlled Factor, **X** = eXperimental Factor (**S** = Screening, **O** = Optimization)

Impact Severity

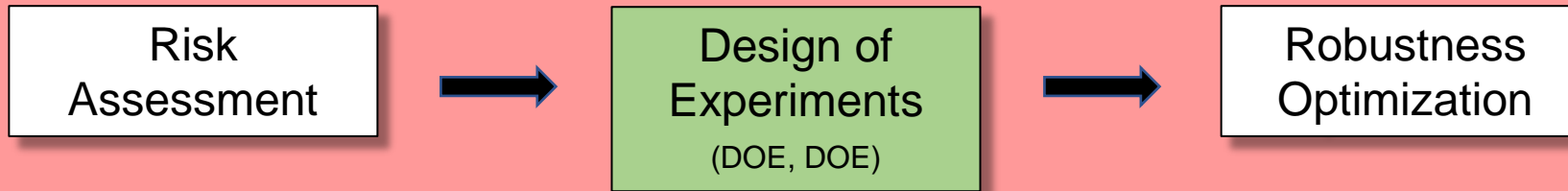
Low = 1

Medium = 3

High = 5

* – CMPs can change depending on the nature of sample compounds and the separation mode.

Design of Experiments (DOE, DoE)



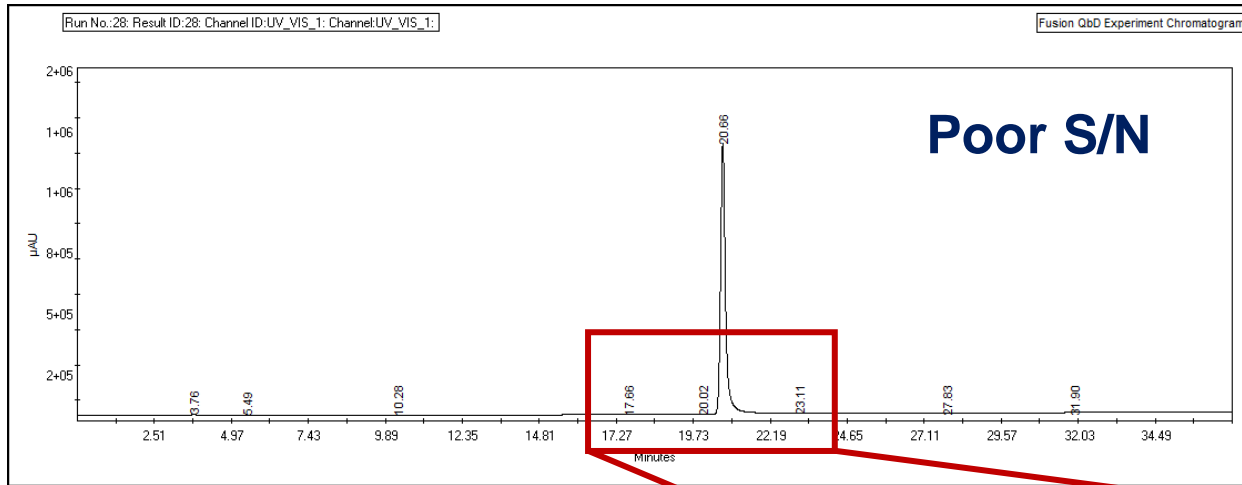
ICH Q14

In an enhanced approach, the ranges for the **relevant parameters and their interactions** can be investigated in multivariate experiments (DoE). (Pg. 5)

USP <1220>

Experimentation is a direct way of generating data that can be used to assess the impact of procedure parameters on performance, and **the use of statistical design of experiments (DOE) is an effective way to do this.** (Pg. 8)

DOE – The Critical First Step

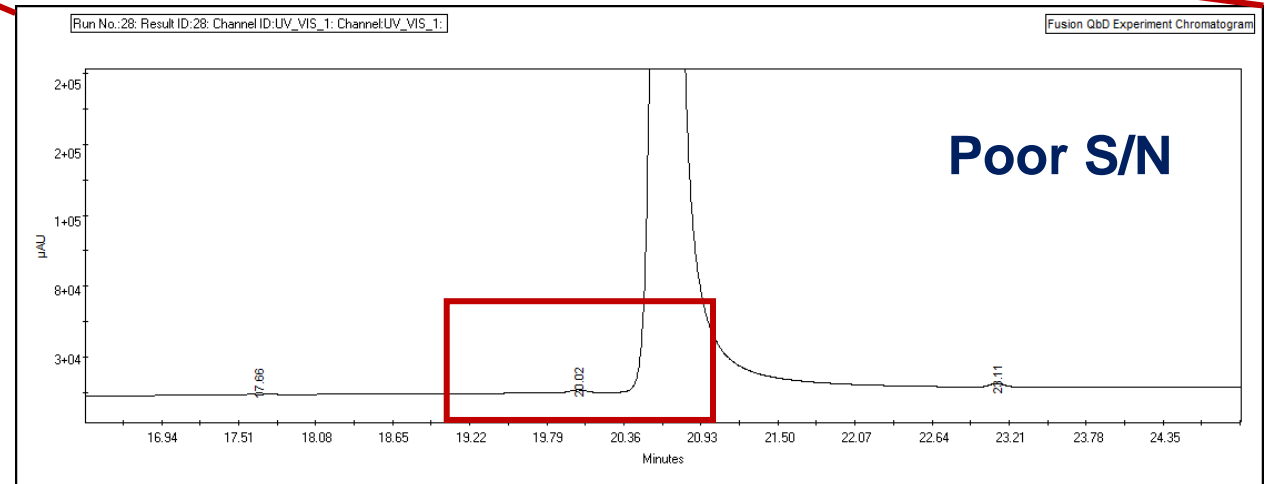


Problems with Determining the Cause of Peak Loss Corrupts the Data.

Co-elution or Just Poor Signal ?

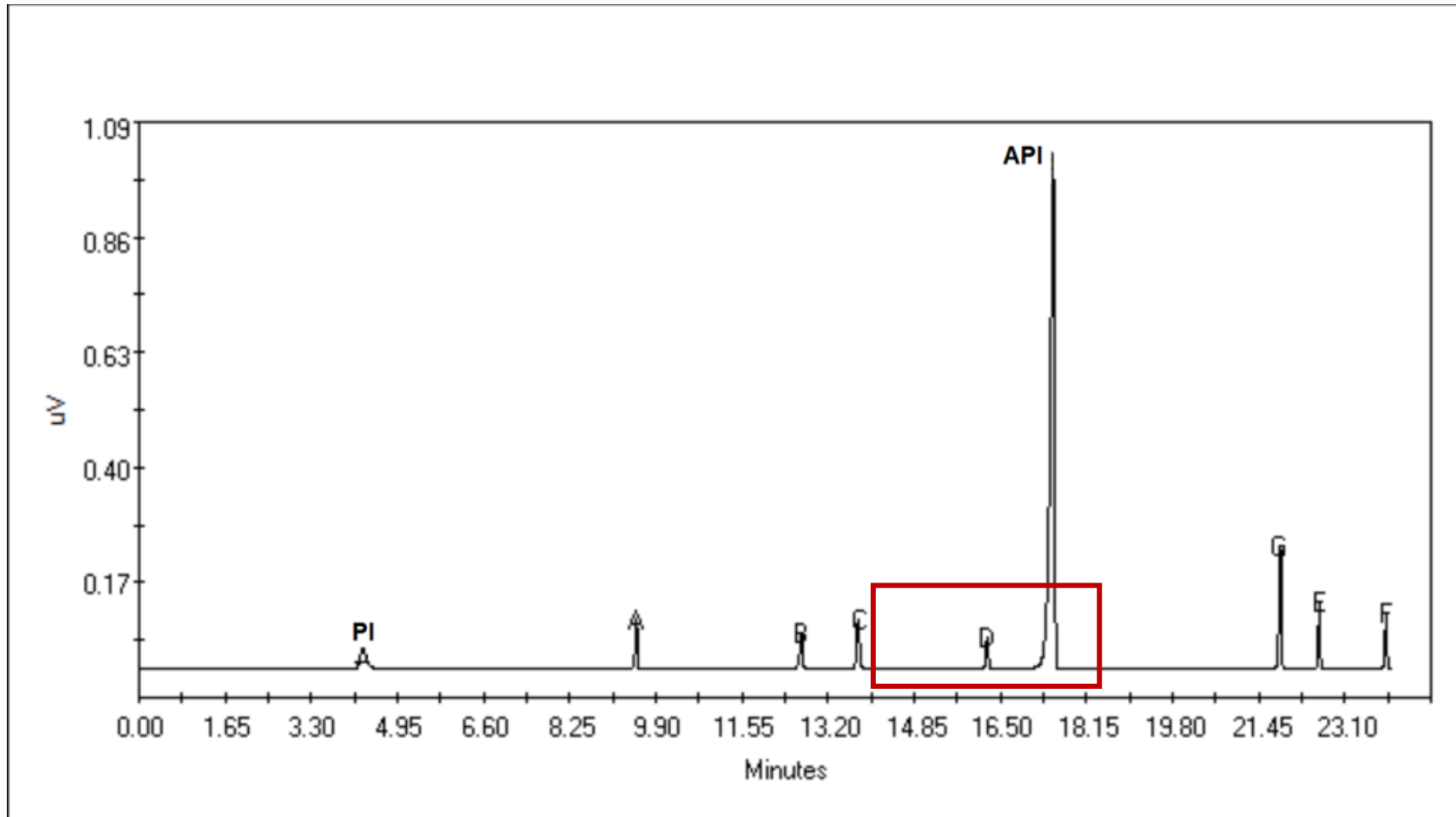
A Method Development Sample should not look like a final QC preparation – it should provide a very good S/N Ratio for all peaks of interest.

S/N Goal: Min = 20, Optimum ≥ 50



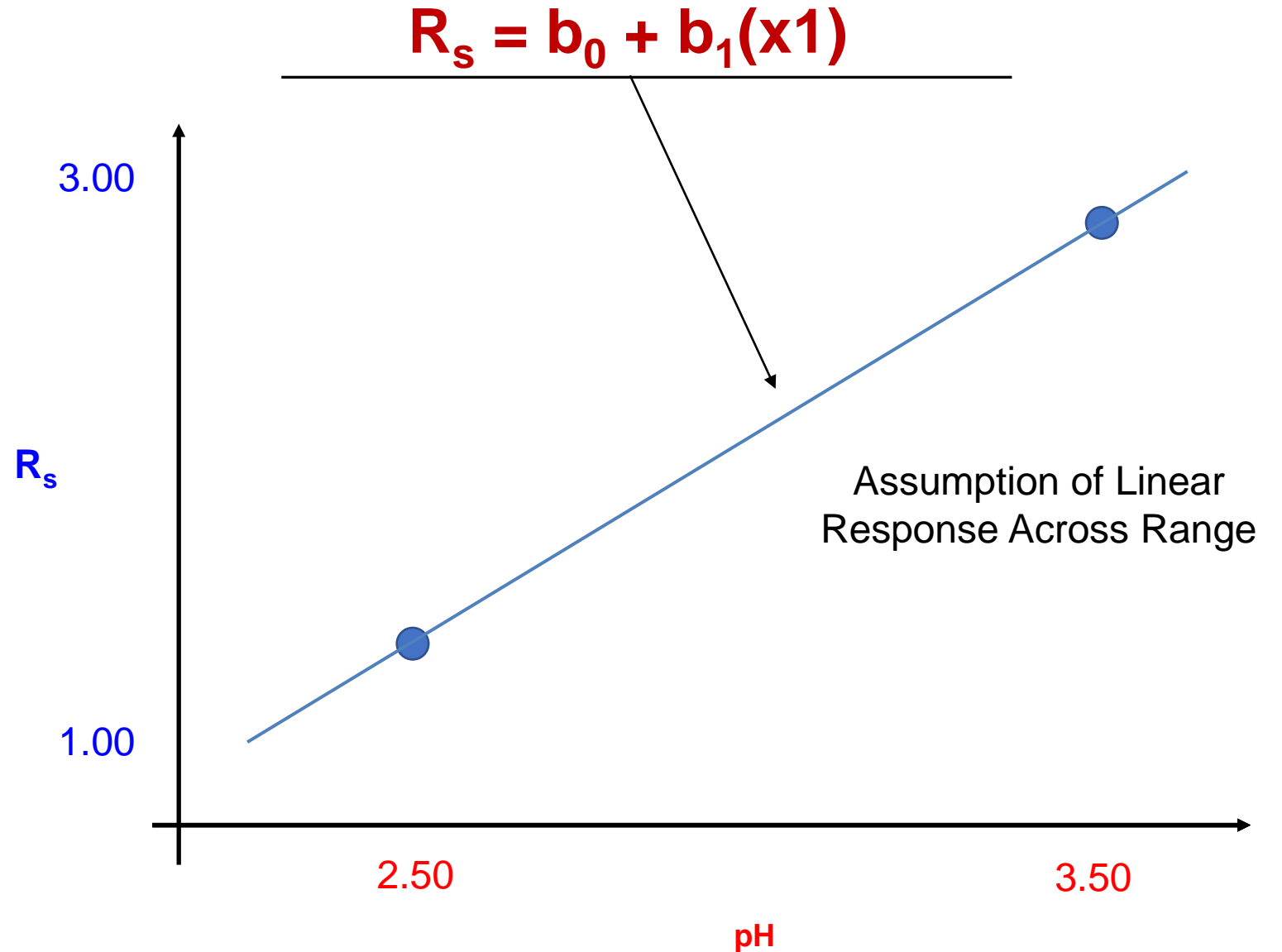
DOE – The Critical First Step

Final QC Preparation Should Provide a Very Good S/N Ratio for all Peaks of Interest



OFAT (Univariate) Experiment

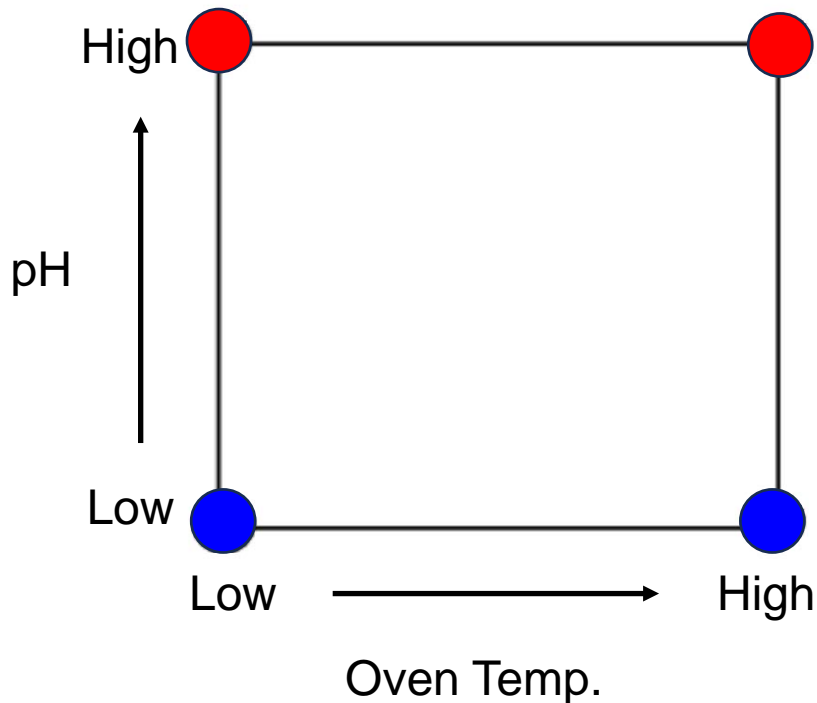
All Factors but One
Held Constant



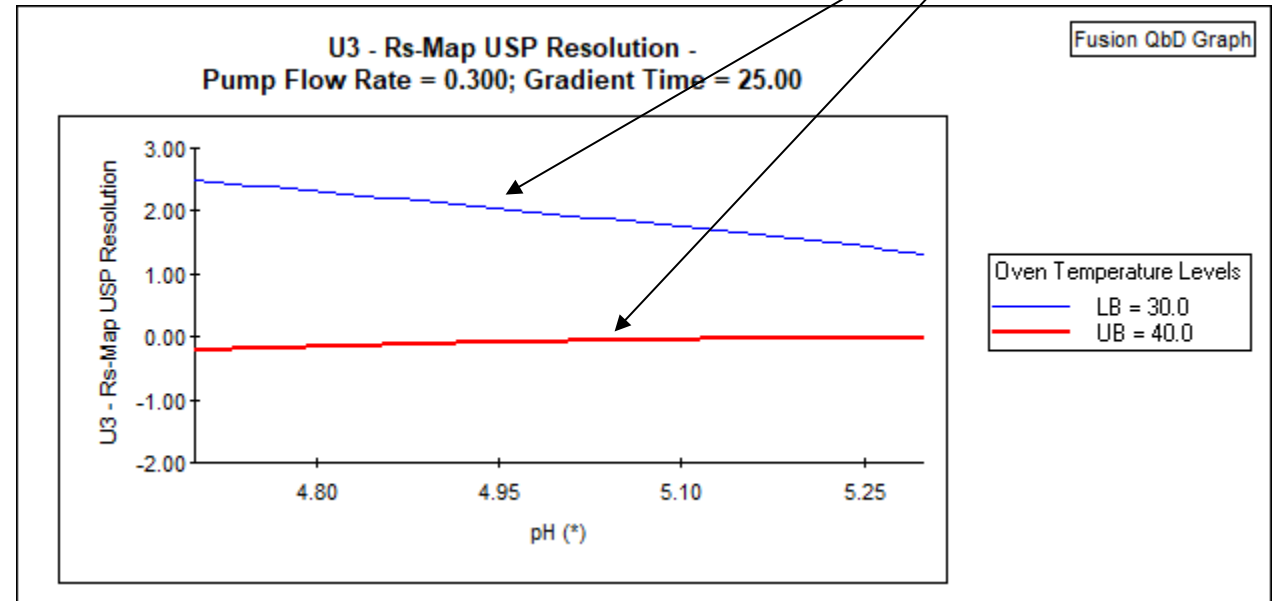
DoE – A Model Building Methodology

Interaction Effect: slope change

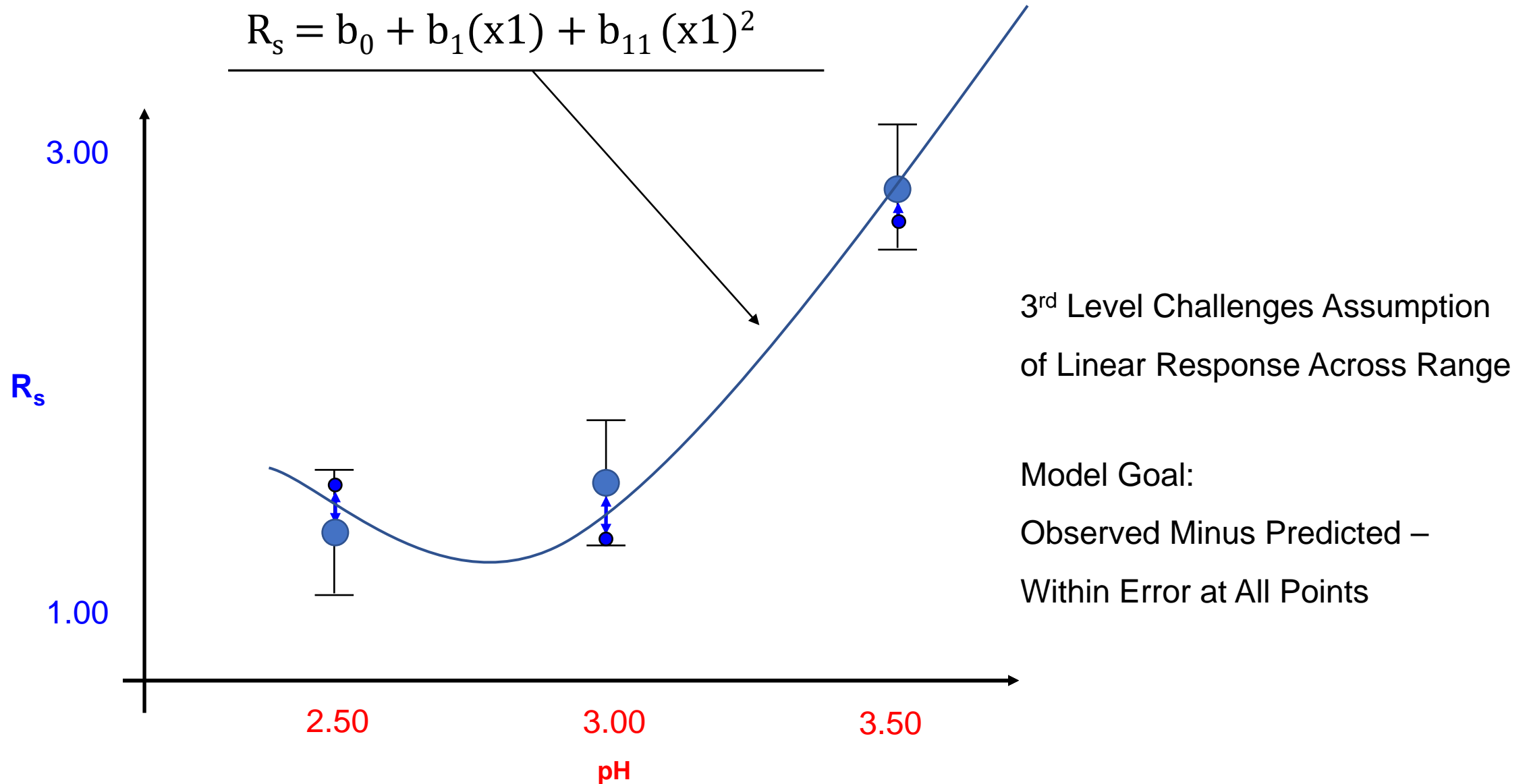
DoE for 2 Variables at 2 Levels –
Simplest Case



$$R_s = b_0 + b_1(x1) + b_2(x2) + b_{12}(x1 * x2)$$

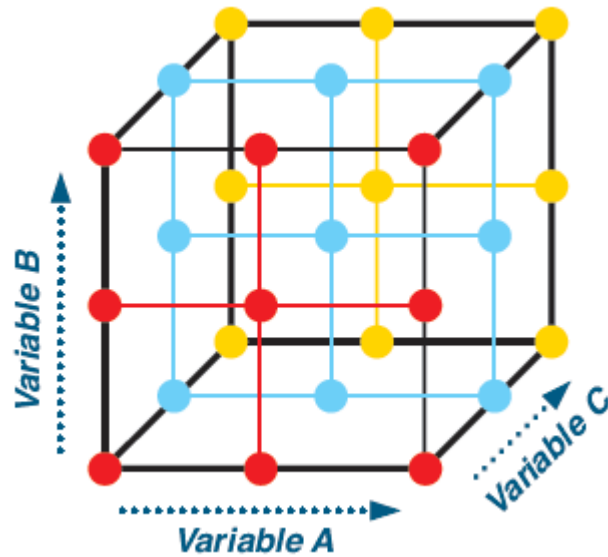


DoE – A Model Building Methodology

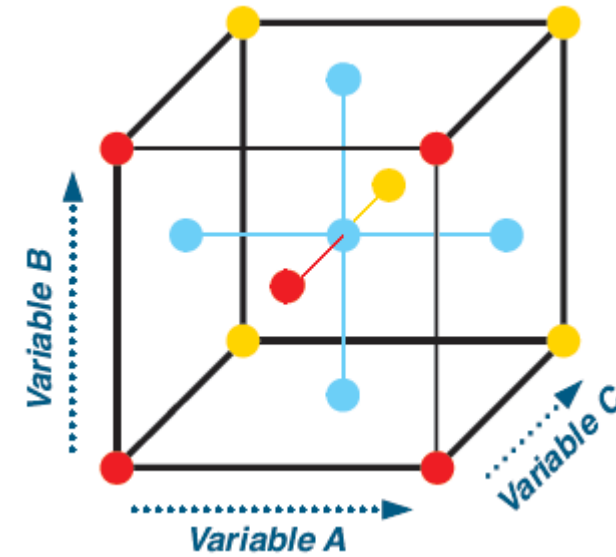
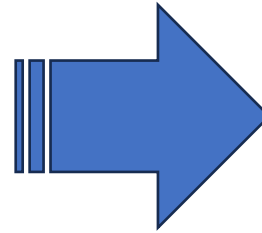


DoE Efficiency

**DoE uses Statistical Sampling of All Possible Combinations to Support
Accurate Estimation of Study Factor Effects**



All Possible Combinations
= 27 Runs



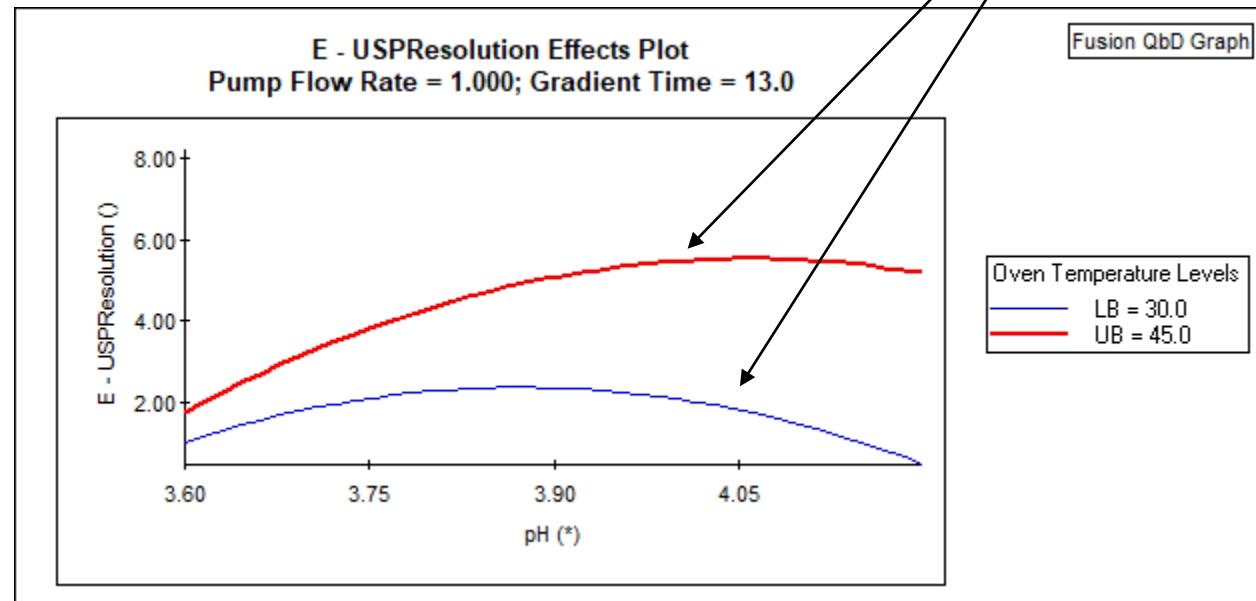
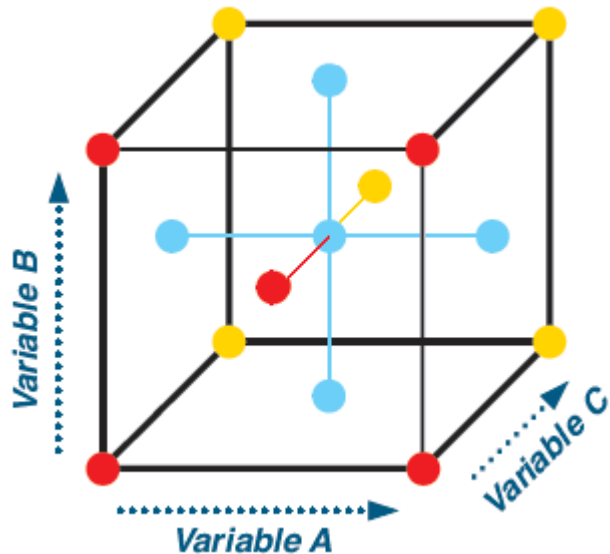
DoE: Central Composite Design
= 15 Runs

DoE – A Model Building Methodology

Squared Interaction Effect: Slope + Curvature Change

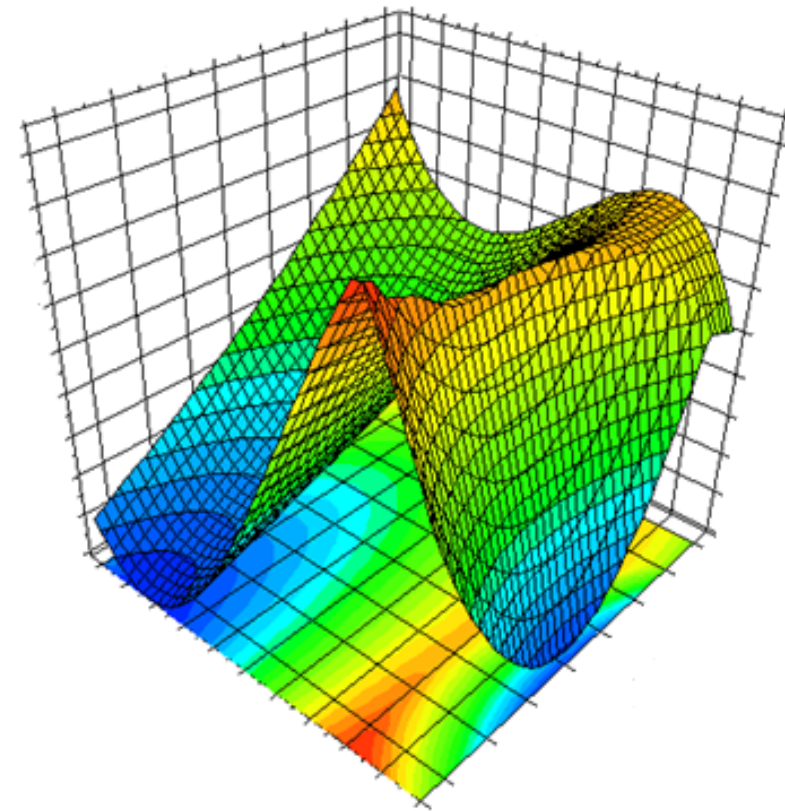
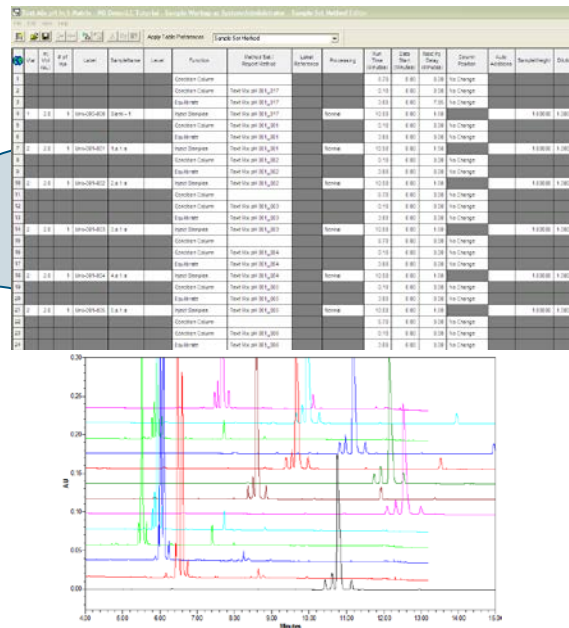
$$R_s = b_0 + b_1(x1) + b_2(x2) + b_3(x3) + b_{11}(x1)^2 + \dots + b_{12}(x1 * x2) \dots + \boxed{b_{112}(x1)^2(x2)} + \dots$$

Central Composite Design



DoE (DOE) – A Model Building Methodology

Turning Chromatograms into Knowledge



$$\# - R_s \geq 2.00 = 9.3 + 4.2(\text{PFR}) - 5.4(\Delta t_G)^2 + 12.7(\Delta t_G * \text{pH}) + 1.3(\text{pH} * \Delta T) + 1.6[(\Delta T)^2 * \Delta t_G] + \dots$$

Linear
Effect

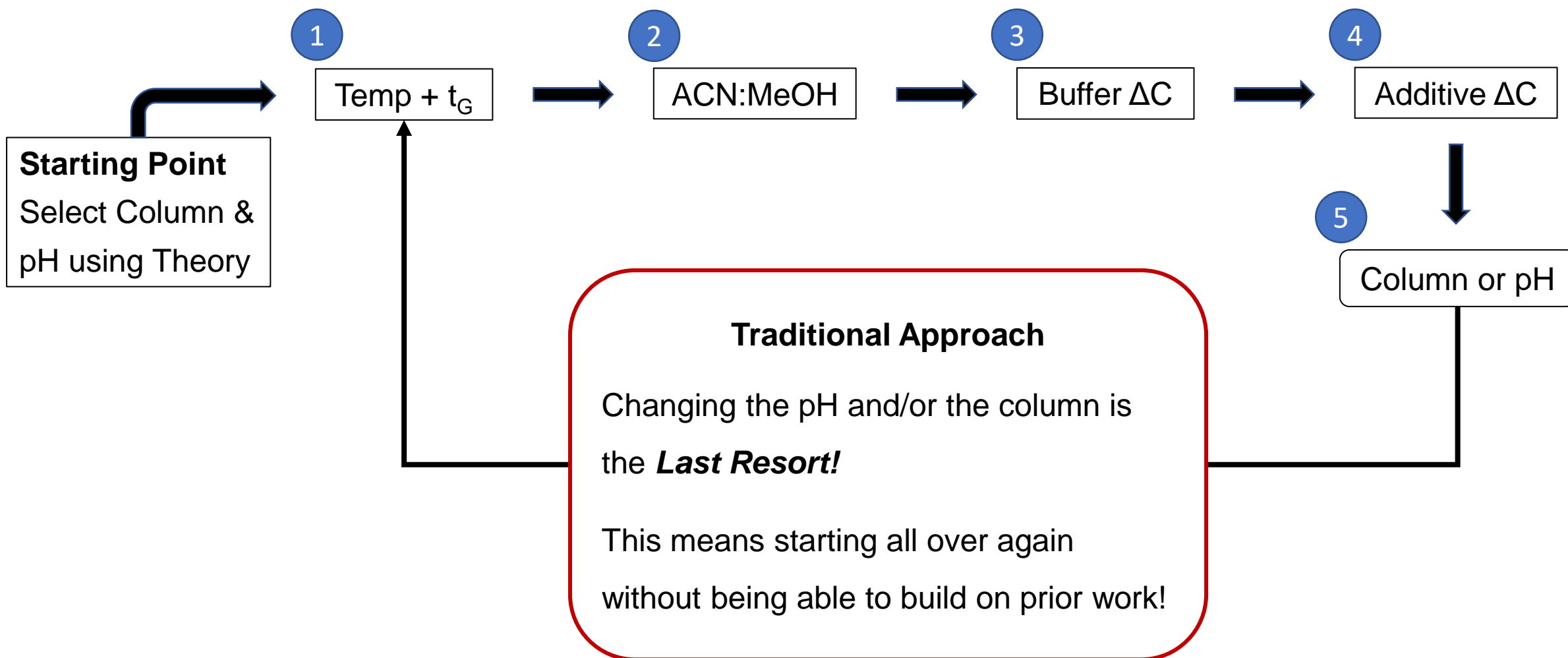
Curvature
Effect

Interaction Effects

Complex Effect

Before Fusion QbD

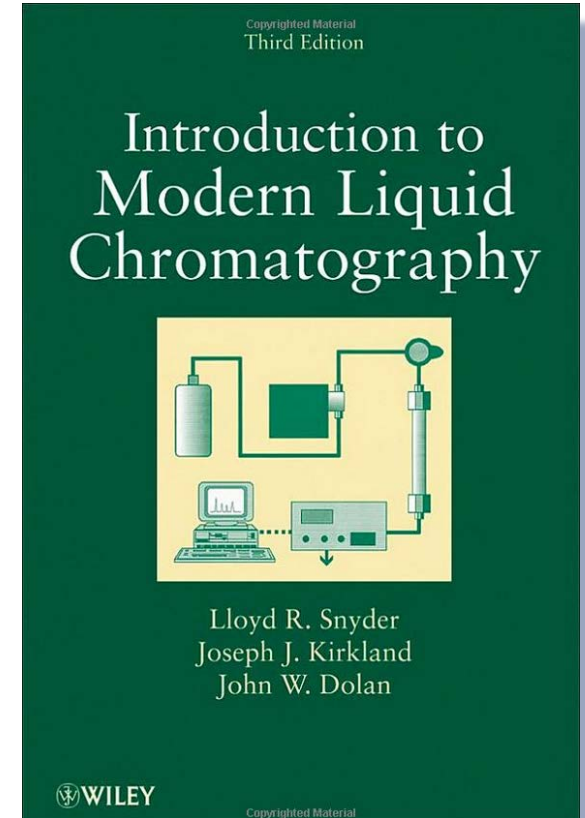
Before Fusion QbD: One-Factor-At-a-Time (OFAT) Approach



Current Thinking by Recognized Experts

“For methods involving a large number of samples, and where adequate resolution must be combined with run times that are as short as possible, **it can be profitable to spend more time initially on “scouting” experiments.**

- Different **columns**
- Different **B-solvents**
- Variations in **pH** and **temperature**
- Use of **Gradient elution** during the experiments can help avoid the need to separately optimize values of %B for each variable studied.”

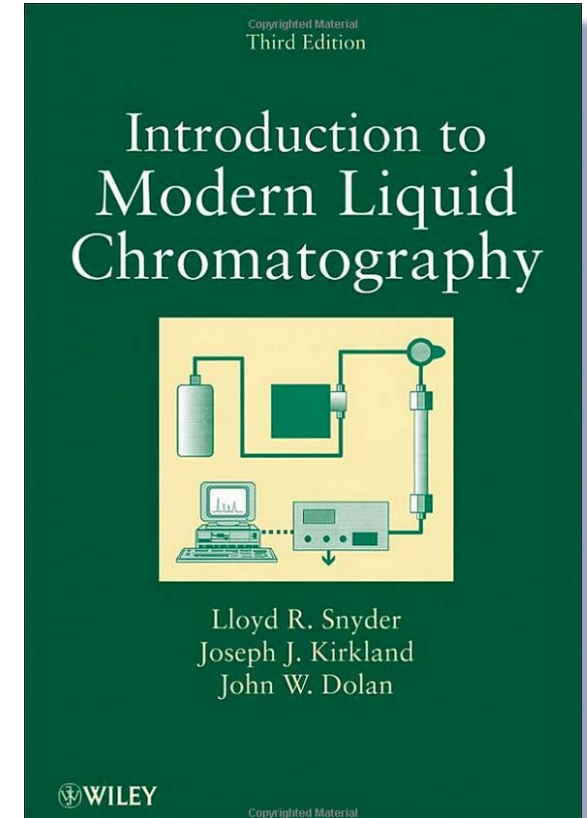


Snyder, Kirkland, and Dolan. (2010). *Introduction to Modern Liquid Chromatography*, 3rd Edition; John Wiley & Sons, Inc., Hoboken, New Jersey (p. 67)

Current Thinking by Recognized Experts

“Still another approach is to **search the literature** for separation of the same or similar sample. **Trial-and-error** modifications of conditions are then followed until an acceptable separation is achieved. ***We do not recommend this approach**** because possible deficiencies in literature methods can delay subsequent attempts at achieving a final, acceptable separation.”

* – italics added by Snyder, Kirkland, and Dolan in book text to emphasize the point.



Snyder, Kirkland, and Dolan. (2010). *Introduction to Modern Liquid Chromatography*, 3rd Edition; John Wiley & Sons, Inc., Hoboken, New Jersey (p. 67)

DOE Supports Study Parameter Flexibility

Full utilization of Quaternary Pumps, Solvent Selection Valves, and Column Switching Valves.

Study any combination of LC parameters which can interactively effect method performance!

- Isocratic and Gradient Methods
- Strong Solvent Type
- Any pump program steps – e.g.
 - Equilibration Time & %
 - Isocratic Hold Time & %
 - Gradient Time / Slope
 - Initial / Final Hold Time & %
 - Re-equilibration Time & %
- Column Temperature
- Column Type
- Flow Rate
- Injection Volume
- pH
- Mobile Phase Blends
- Salt, Buffer, Additive – Type & ΔC
- Wavelength



Experiment Setup – Column Type

Column Settings

* - Sufficient Conditioning Time for 10 column volumes is recommended.

	Name	Valve Position	pH Upper Limit	Flow Rate	Diameter (mm)	Length (mm)	Time Required for One Column Volume (min)	Conditioning Time (min)*
1	HSS T3	Position 1 ▼	8.00	0.400	2.10	100.00	0.9	9.0
2	CSH Phenyl-Hexyl	Position 2 ▼	11.00	0.400	2.10	100.00	0.9	9.0
3	BEH C8	Position 3 ▼	12.00	0.400	2.10	100.00	0.9	9.0
4	BEH Shield RP18	Position 4 ▼	11.00	0.400	2.10	100.00	0.9	9.0

Chemistry Intelligence –

- Blocks design on Column Temp when it is a study factor
- Groups runs by MP Chemistry (e.g., pH, Strong Solvent)
- Incorporates column conditioning between MP Chemistry changes

Valve Intelligence – Automatically generates multiple sequences as needed when # of columns in exceeds # of available valve positions.



Experiment Setup – Mobile Phase Blending

Experiment Setup

Replication Settings

Method Type: Gradient

Available Variables: Isocratic, Gradient

Gradient Curve
Gradient Slope
Sample Concentration
Additive Concentration
Additive Type

◀ ▶

Included Variables: Pump Flow Rate
Injection Volume
Oven Temperature
Wavelength
Column Type

☒ Activate Online Preparation
☒ pH
☐ Buffer Concentration
☐ Additive Concentration

Solvent Settings

No. of Strong Solvents: 1 No. of Weak Solvents: 3

☐ OK to Blend Strong Solvents ☒ OK to Blend Weak Solvents

Mobile Phase Precision: 0.00 0.00

Mobile Phase Name	Solvent Type	State	Lower Bound	Upper Bound	Reservoir
Acetonitrile	Strong (Organic)	---	---	---	---
Buffer	Weak (Aqueous)	Variable	85.5	89.5	---
Methanol	Weak (Aqueous)	Variable	3.5	6.5	---
IPA	Weak (Aqueous)	Variable	5.5	9.5	---

Available Reservoirs

☒ A ☒ D

☒ B ☒ D-1 ☒ D-2 ☒ D-3

☒ C ☒ D-4 ☒ D-5 ☒ D-6

Experiment Setup – pH

Select One of the Built-in Buffer Systems or Enter Your Own

Buffer Selector...

pH Online Blending Mode: **One Acid, Base Pair**

pH Buffer Settings

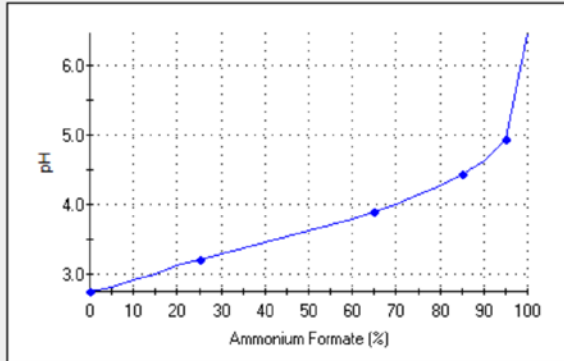
Buffer	Buffer Name	pH Level
Acid	Formic Acid (20 mM)	2.73
Base	Ammonium Formate (20 mM)	3.29
		3.88
		4.42
		4.93

No. of Levels: 5

Buffer Selector

Select Buffer System
pH 2.74 - 6.45 [Formate System (20 mM)]

Buffer Solutions
Formic Acid (20 mM) - pH 2.74
Ammonium Formate (20 mM) - pH 6.45



	Included	pH	Formic Acid (%)	Ammonium Formate (%)
1	<input checked="" type="checkbox"/>	2.74	100.00	0.00
2	<input type="checkbox"/>	2.80	95.00	5.00
3	<input type="checkbox"/>	2.91	90.00	10.00
4	<input type="checkbox"/>	3.00	85.00	15.00
5	<input type="checkbox"/>	3.11	80.00	20.00
6	<input checked="" type="checkbox"/>	3.20	75.00	25.00
7	<input type="checkbox"/>	3.29	70.00	30.00
8	<input type="checkbox"/>	3.37	65.00	35.00
9	<input type="checkbox"/>	3.45	60.00	40.00
10	<input type="checkbox"/>	3.53	55.00	45.00
11	<input type="checkbox"/>	3.61	50.00	50.00
12	<input type="checkbox"/>	3.69	45.00	55.00
13	<input type="checkbox"/>	3.78	40.00	60.00
14	<input checked="" type="checkbox"/>	3.88	35.00	65.00
15	<input type="checkbox"/>	3.99	30.00	70.00
16	<input type="checkbox"/>	4.14	25.00	75.00
17	<input type="checkbox"/>	4.27	20.00	80.00
18	<input checked="" type="checkbox"/>	4.42	15.00	85.00
19	<input type="checkbox"/>	4.62	10.00	90.00
20	<input checked="" type="checkbox"/>	4.93	5.00	95.00
21	<input type="checkbox"/>	6.45	0.00	100.00

Select All Select None OK Cancel

Select Desired pH Levels from the Built-in Buffer Curve or Use Your Own Curve



Experiment Setup – Gradient Time

Pump Program

No. of Gradient Steps: 1

Time Precision ± 0.00

	No.	Step Name	Time State	Time - Lower Bound	Time - Upper Bound	% Strong Solvent
<input checked="" type="checkbox"/>	1	Equilibration	Constant	3.00	---	5.0
	2	Initial Hold	Constant	1.00	---	5.0
	3	Gradient	Variable	10.00	25.00	---
	4	Final Hold	Constant	1.00	---	95.0
	5	Ramp Up to Wash	Constant	0.50	---	---
<input checked="" type="checkbox"/>	6	Column Wash	Constant	2.00	---	99.0
	7	Ramp Down from Wash	Constant	0.50	---	---
<input checked="" type="checkbox"/>	8	Re-equilibration	Constant	2.00	---	5.0

Program duration: Min = 20.00 minutes, Max = 35.00 minutes

Gradient Study Factor: Gradient Time

Setting Mode

☒ Time☐ Slope

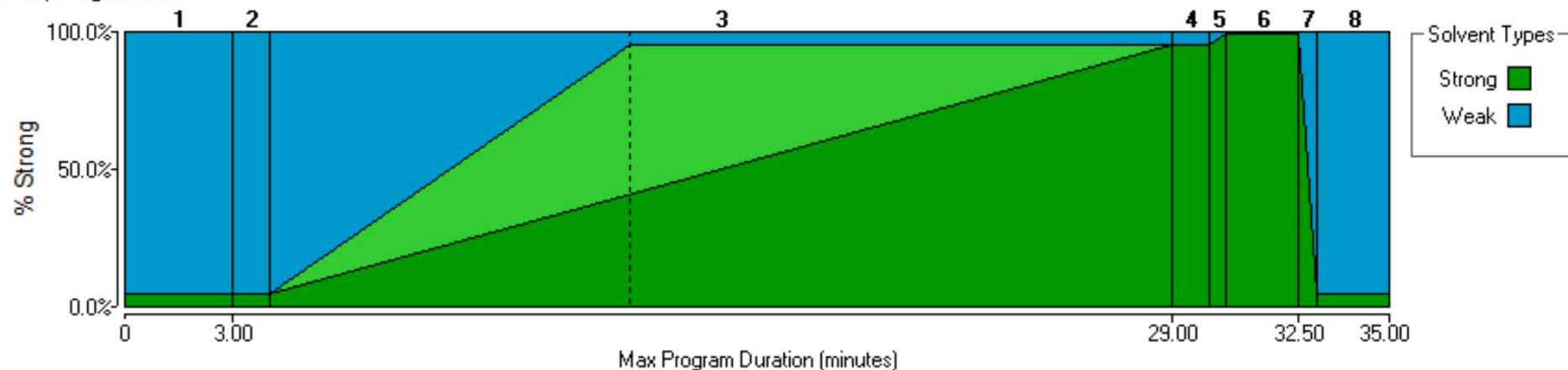
Update

Gradient

No. of Levels 3

Time (min)	Slope (%/min)
10.00	9.00
17.50	5.14
25.00	3.60

Pump Program Chart



Defining the Best Gradient Time Study Range

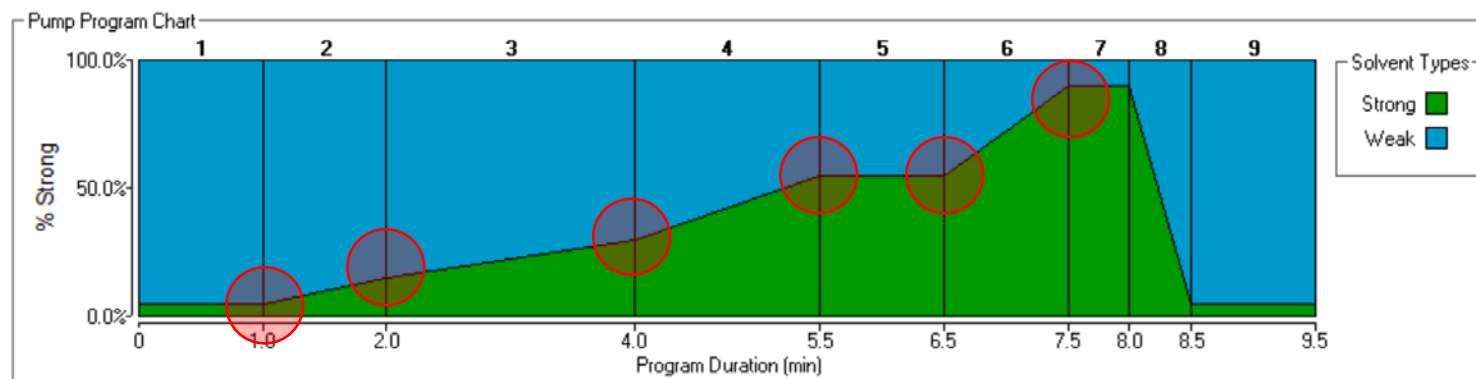
Whereas it is frequently recommended that the slopes of scanning gradients used to obtain retention data should vary by a factor of three or so, we do not see any evidence in our results that support this guideline. That is, similar retention prediction errors were obtained from models based on scanning gradients with slopes varying by a factor of three compared to models based on gradients with slopes varying by as little as 1.25. We also observe that the speed (i.e., absolute analysis or gradient time) does not have a strong impact on prediction error. On the other hand, the data show that **the proximity of the slope of a gradient, for which retention will be predicted, to one of the scanning gradients, used to build the model, is far more determinant of retention prediction error.** With decreasing proximity, it is more important that the slope of the target gradient lies between the slopes of the scanning gradients (i.e., interpolation is better than extrapolation, as one would expect).

These findings have obvious implications for the design of experiments; using scanning gradients with a large variation in slopes is not required per se, but **using a large range of slopes enables prediction of retention for a wider array of gradients without extrapolating.**

What about Multi-segment Gradient Optimization

Most LC Method Development software relies primarily on localized gradient slope-based optimization. This drives the user to a multi-segment gradient method.

Multi-segment Gradients = Multiple Regions of POOR Robustness!



Localized Slope-Based Optimization is Now Recognized as High Risk.

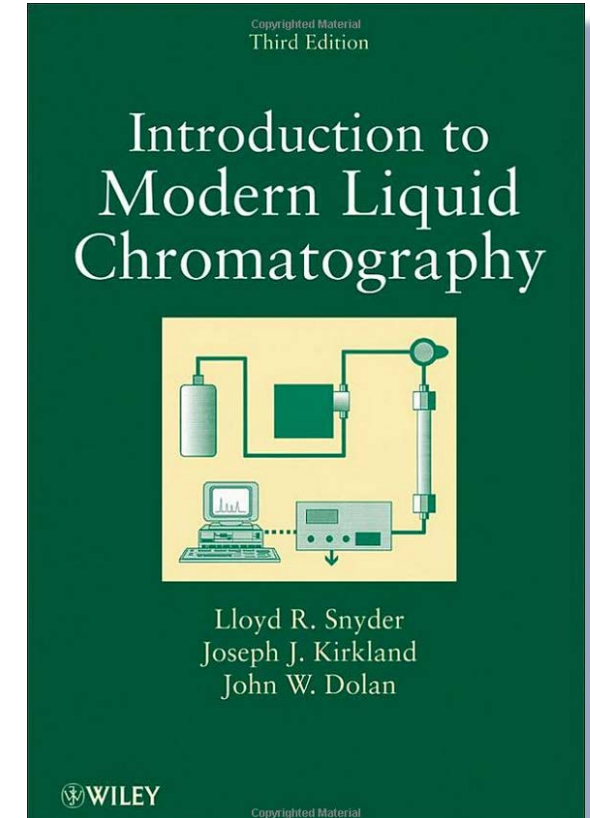
Fusion QbD Does Not Rely on This Approach!

(See the Next Slide →)

Current Thinking by Recognized Experts

Issues with a Multi-step Gradient Approach to Method Optimization

“Increasing resolution by adjusting selectivity for different parts of the chromatogram can sometimes be achieved with a segmented gradient; ... **Segmented gradients are not often used for improving resolution ... because their ability to enhance resolution without increasing run time is usually limited**... However, there are other – generally more useful – means of optimizing resolution by changing selectivity and relative retention. **Also, separations that use segmented gradients to improve resolution are likely to be less reproducible when transferred to another piece of equipment.**”



Snyder, Kirkland, and Dolan. (2010). *Introduction to Modern Liquid Chromatography*, 3rd Edition; John Wiley & Sons, Inc., Hoboken, New Jersey (p. 427-28)



Full Support for Forced Degradation Studies

☒ Forced Degradation Study

No. of Unique Degradation Path Samples

No. of Injection Repeats Per Sample

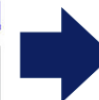
Path #	Degradation Path Description
	Sample Compound Mix
1	Oxidation

Experiment Design Matrix

Run No.	Pump Flow Rate (mL/min)	Gradient Time (min)	Oven Temperature (°C)	pH
Conditioning_Run_1	0.400	2.0	30.0	4.60
1.a	0.300	8.0	30.0	4.60
1.b	0.300	8.0	30.0	4.60
2.a	0.500	8.0	30.0	4.60
2.b	0.500	8.0	30.0	4.60
3.a	0.300	16.0	30.0	4.60
3.b	0.300	16.0	30.0	4.60

Simple Setup integrates the replication scheme into the DoE Study, and automatically assigns a separate vial position to each replicate injection.

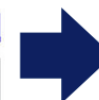
1.a Vial



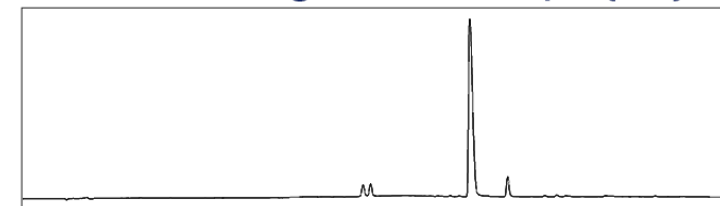
Sample Compound Mix (1.a)



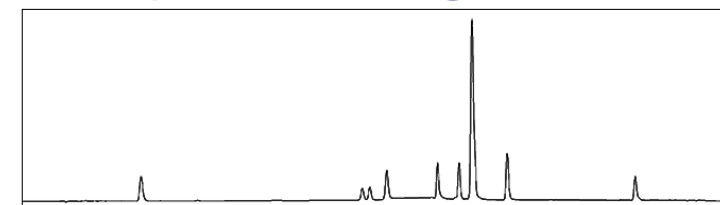
1.b Vial



Oxidation Degradation Sample (1.b)



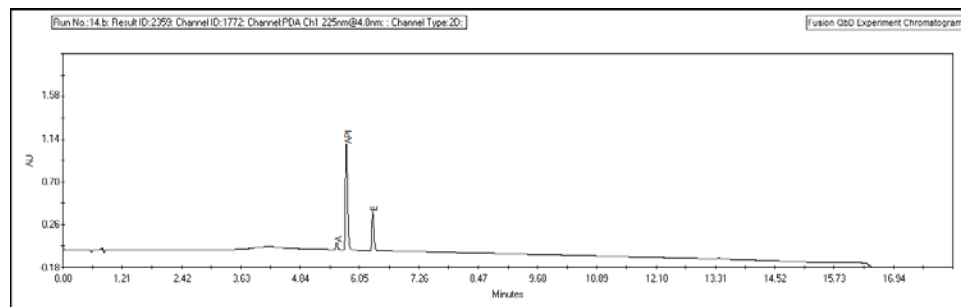
Composite Chromatogram – Run 1



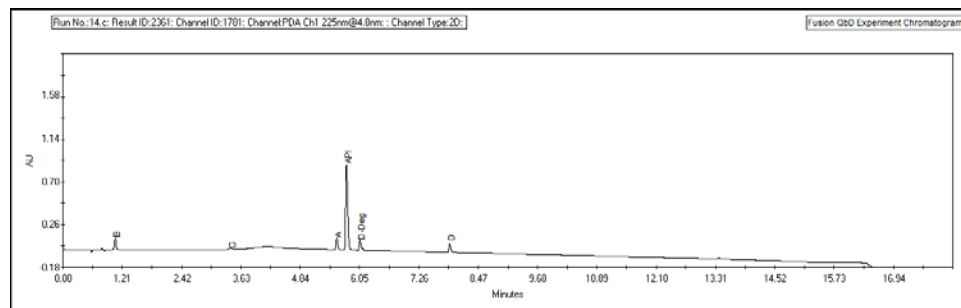
Fusion QbD tracks all peaks in all replicate chromatograms for each run and generates a *composite chromatogram* for each run containing all unique peaks from all replicate injections.

FDS Composite Chromatogram – 3 Degradation Path Example

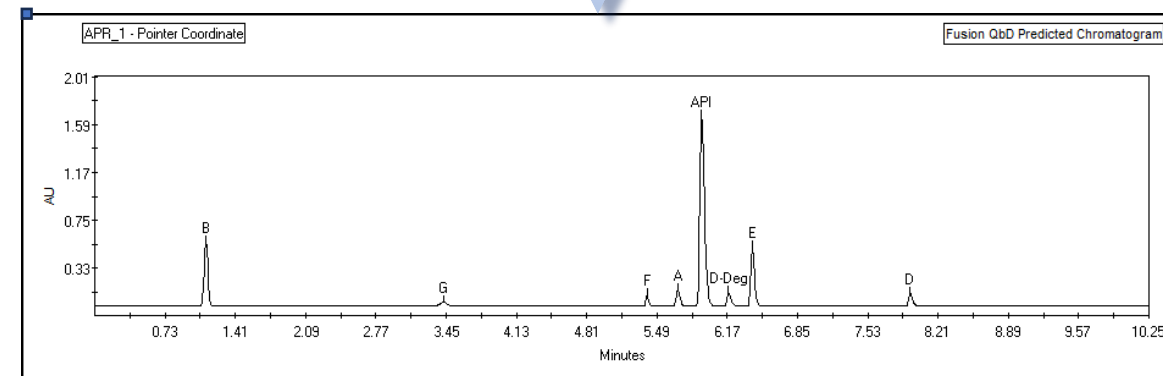
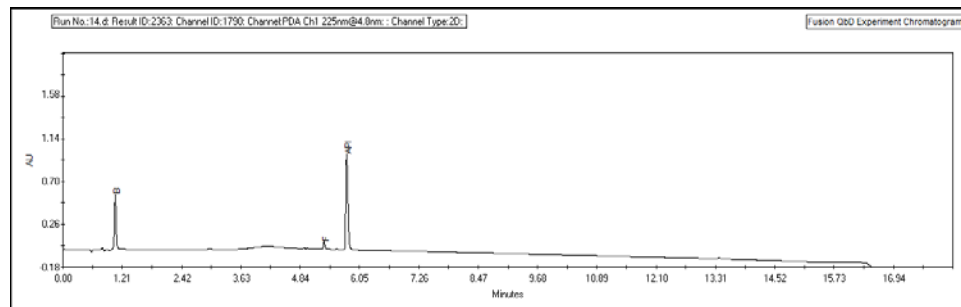
Acid Degradation Path



Base Degradation Path



Peroxide Degradation Path



Fusion QbD Composite Chromatogram

Fusion QbD – DoE Efficiency Example

5 levels of Gradient Time

5 levels of pH

4 levels of Column Type

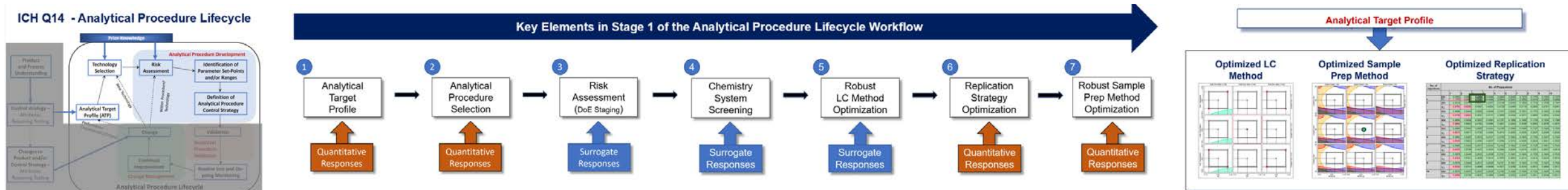
$5 \times 5 \times 4 = 100$ possible combinations

Fusion QbD design = 30 runs (plus 6 repeats)

> 3x efficiency.

End of Presentation

Fusion QbD is the Only LC Method Development Software Which Completely Supports the AQbD / APLM Workflow in the Regulatory Guidances



ICH Q2(R2) / ICH Q14 / USP <1210> / USP <1220> / EP 11.60