
Enhanced Analytical Development for Complex Antibody Formats

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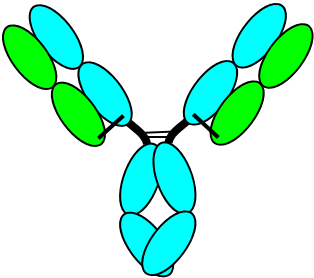




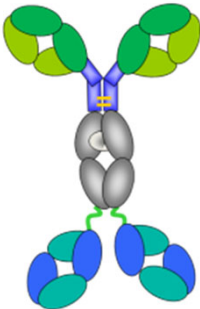
Outline

- Introduction
- Tools for Enhanced Method Development
- Case Study 1
- Case Study 2

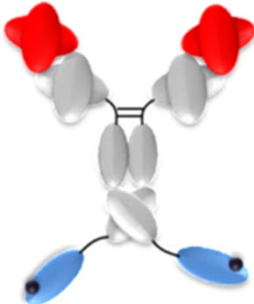
More Complex Antibody Formats



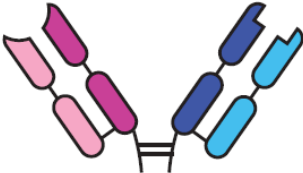
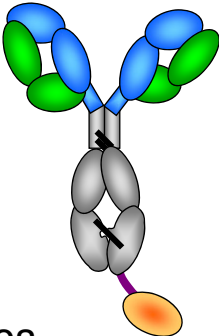
MAb



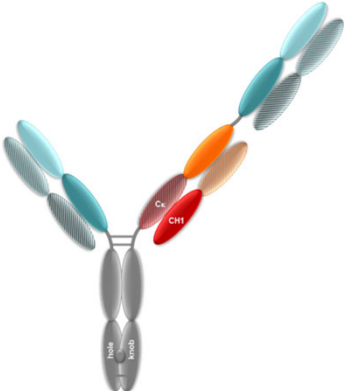
Appended BsAb



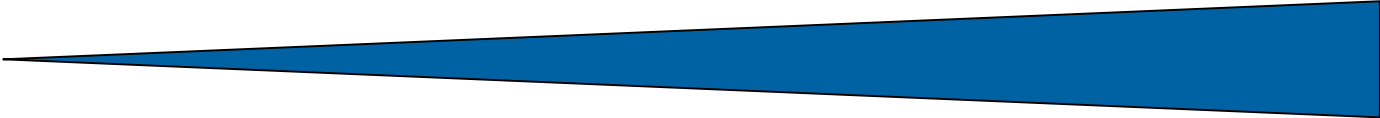
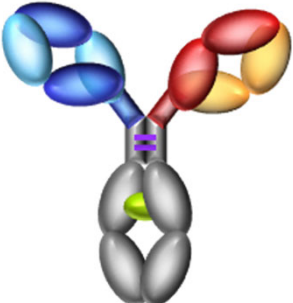
Conjugates



BsAb Fragments



Bispecific CrossMAb



Increase of complexity within the last 10 years

Complex Antibody Formats - Consequences

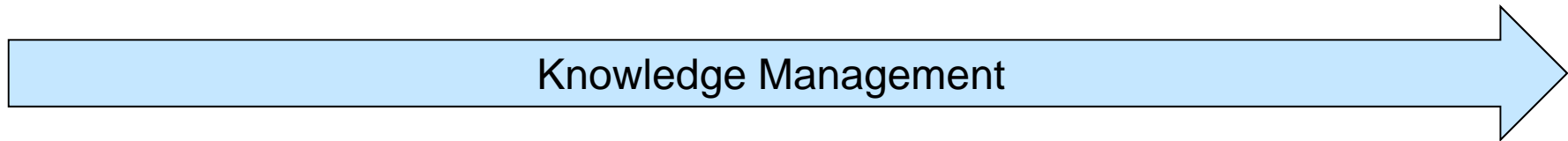
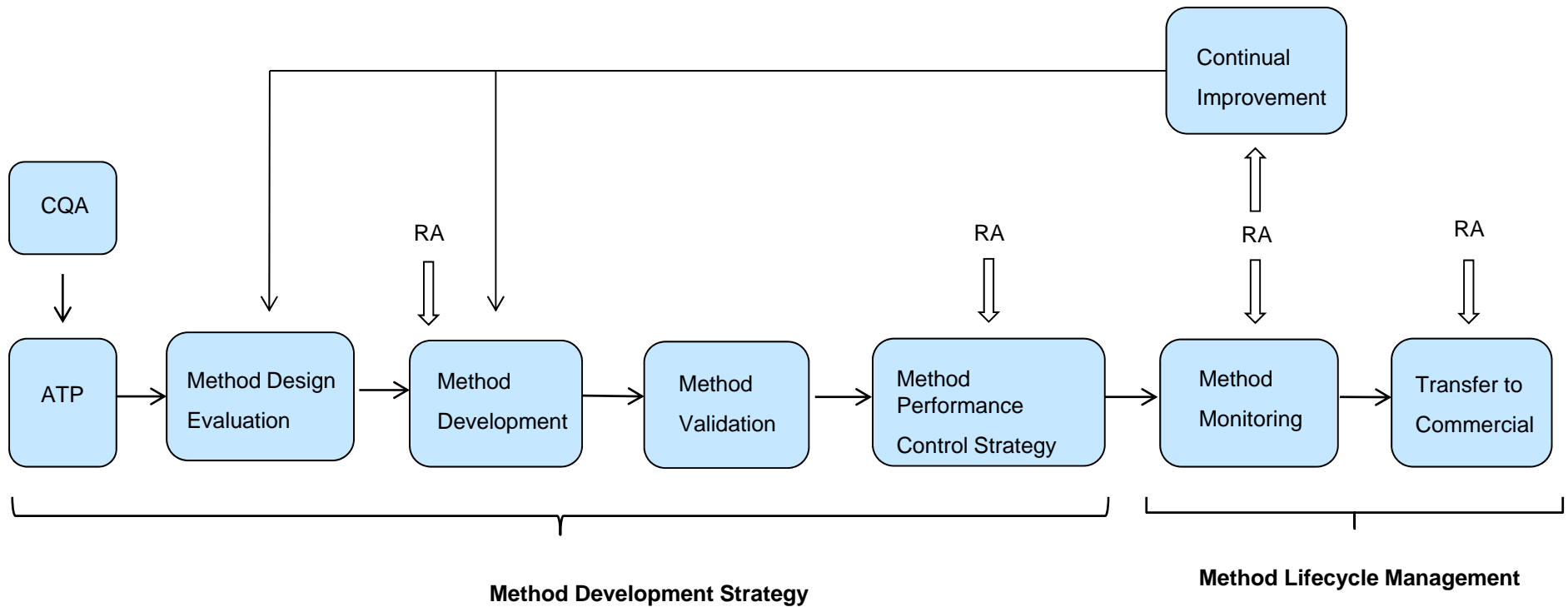
- New product related substances and impurities
 - New critical quality attributes (CQAs)
 - Less prior knowledge and more unknowns
 - More analytical challenges have been observed with platform methods
 - Insufficient resolution
 - Carry over
 - Insufficient sample stability
 - Insufficient robustness
 - Method induced artifacts
- > Need to switch from platform methods to product specific methods
- > Increased investment in analytical development
- > Enhanced analytical development is needed in order to address complexity and unknowns



Analytical Quality by Design Principles (AQbD)

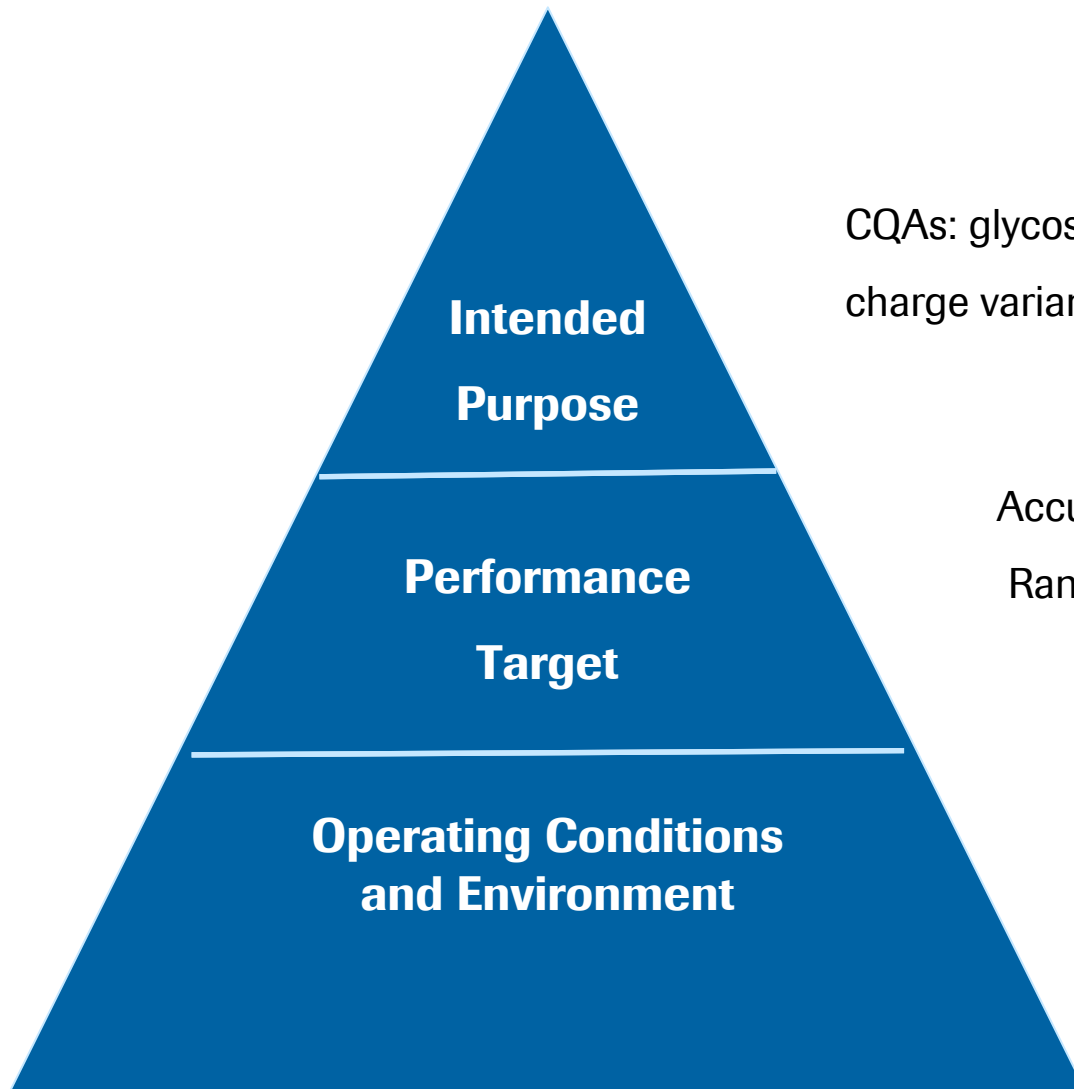
- Use of quality by design (QbD) principles according to the guidelines of international conference on harmonization ICH Q8-11
 - predefined objective
 - science and risk-based development
 - use multivariate DoE studies to define method operational limits
 - control strategy
 - lifecycle management
 - continual improvement
- > Analytical quality by design (AQbD) helps to develop robust methods which are applicable throughout the lifecycle of the method.
- > The objective is to understand, reduce and control sources of variability.

Method Lifecycle



CQA: critical quality attribute, ATP: Analytical Target Profile, RA: risk assessment

Analytical Target Profile (ATP)



CQAs: glycosylation, size variants
charge variants, oxidation, potency etc

Accuracy, Precision, Sensitivity,
Range etc / ICH Q2

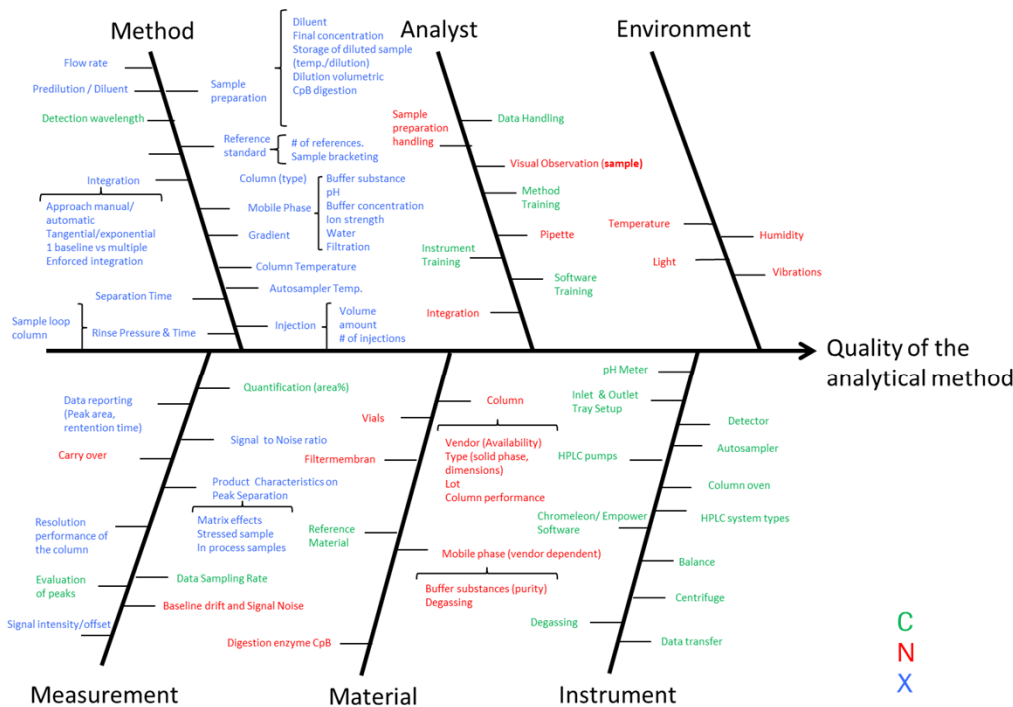
Stable test procedure ($\geq 48\text{h}$)
No harmful ingredients
Operator safety
User requirements: low costs, speed etc

Risk Assessment



Risk assessments are performed in order to identify and assess critical method variables and parameters that can impact the ATP. These structured risk assessments are used to guide experimental plans.

Ishikawa Diagram



Preliminary Hazard Analysis (PHA)

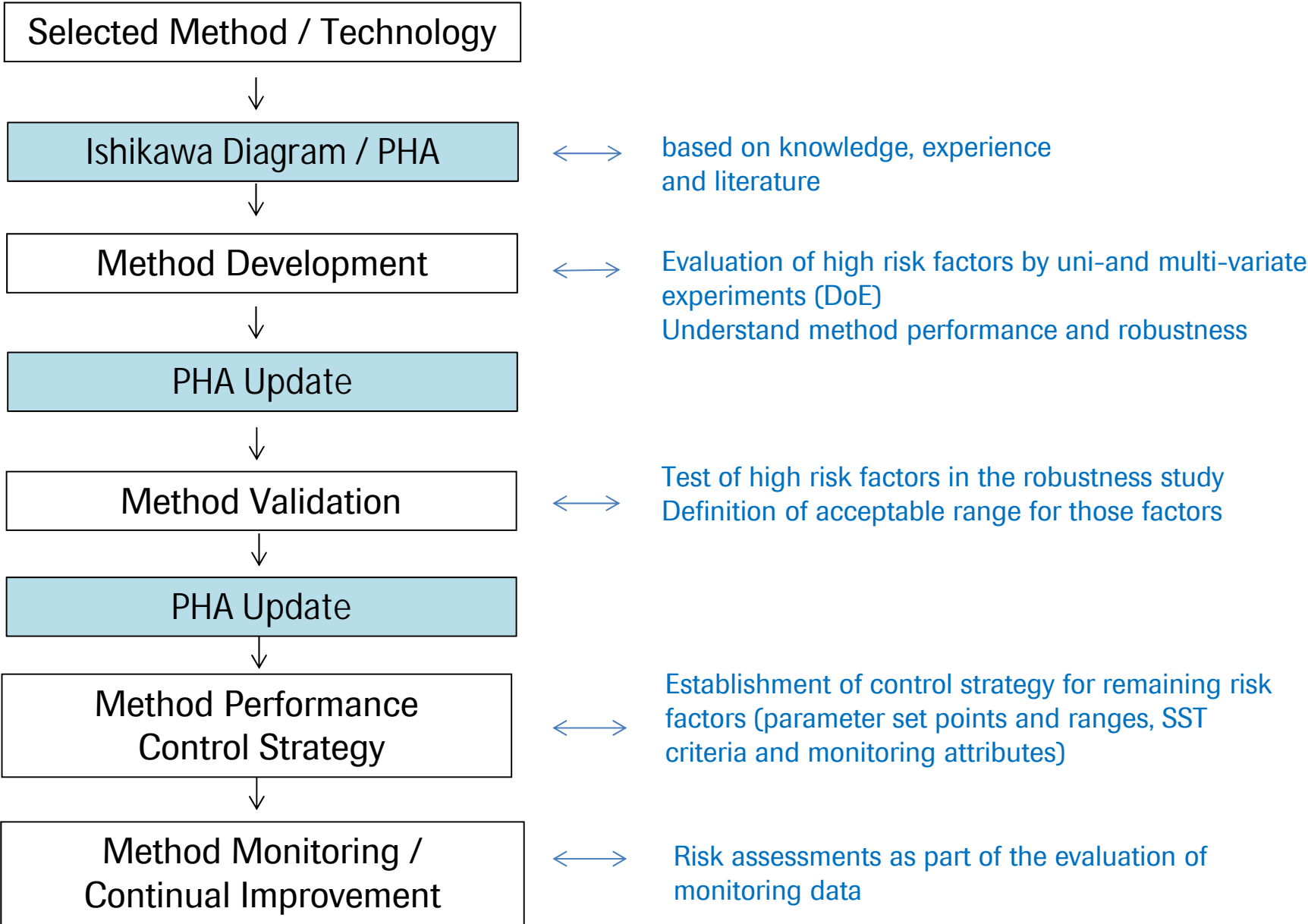
#	Category	Factor	PHA1 Classif.	PHA2 PRN	PHA1 PRN	PHA2 PRN
1	Method	flow rate	X	C	12	12
2	Method	predilution	X	X	12	12
3	Method	detection wavelength	C	C	12	12
4	Method	sample preparation: diluent	X	C	60	12
5	Method	sample preparation: final concentration	X	X	36	12
6	Method	sample preparation: storage of diluted sample - temperature	X	C	60	12
7	Method	sample preparation: storage of diluted sample - time	X	C	36	12
8	Method	sample preparation: volumetric dilution	X	X	12	12
9	Method	sample preparation: CpB digestion	X	X	12	12
10	Method	RS: # of references	X	X	12	12
11	Method	RS: sample bracketing	X	X	12	12
12	Method	integration: approach manual/ automatic	X	C	36	12
13	Method	integration: tangential/ exponential	X	C	12	12
14	Method	integration: one baseline vs. multiple enforced integration	X	C	60	36
15	Method	mobile phase: buffer substance	X	C	60	12
16	Method	mobile phase: pH	X	C	60	12
17	Method	mobile phase: buffer concentration	X	C	36	12
18	Method	mobile phase: ionic strength	X	C	60	12
19	Method	mobile phase: water	X	C	12	12
20	Method	mobile phase: filtration	X	C	36	36
21	Method	gradient	X	C	60	12
22	Method	column temperature	X	C	60	12
23	Method	autosampler temperature	X	C	12	12
24	Method	injection: volume	X	X	12	12
25	Method	injection: amount	X	C	36	36
26	Method	injection: No. of sample injections per sequence	X	X	12	12
27	Method	separation time	X	C	36	12
28	Method	column: rinse pressure & time	X	X	12	12
29	Method	sample loop: rinse pressure & time	X	X	12	12
30	Method	column (type)	X	C	60	12

A tool known as CNX is used to classify the identified factors. It must be decided which factors can be controlled (C), which are potential noise factors (N) and which should be assessed experimentally (X) to determine acceptable conditions and ranges

A preliminary hazard analysis (PHA) is used for the further risk assessment. Relevant factors like X-factors and critical N-factors which might have an influence on the method performance should be selected and assessed.

High risk factors should be evaluated during method development by using a Design of Experiments approach (DoE).

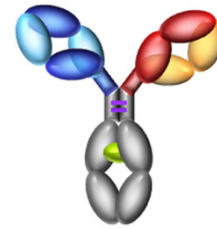
Risk Assessments



PHA: Preliminary Hazard Analysis

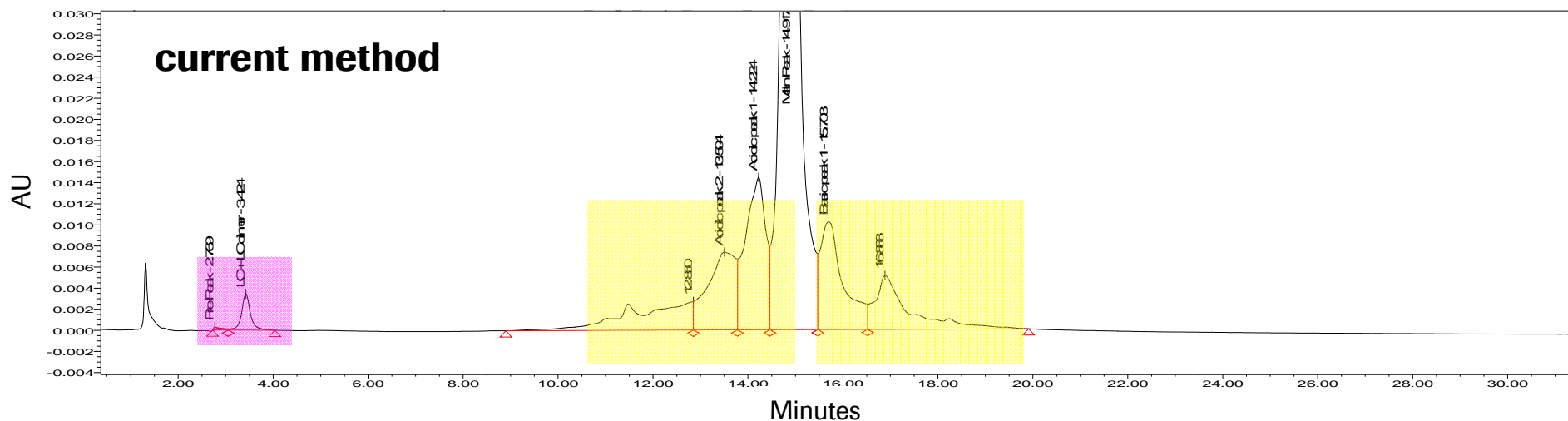
Case Study 1

- IEC method development for a BsAb



«Triggers» to develop a new IEC Method

- need for improved resolution of acidic and basic species
- need for increased method robustness regarding:
 - retention of LC/LC₂ signal
 - baseline / peak (group) delimiters
 - column lot-to-lot and system-to-system variability




Analytical Target Profile

Ion Exchange Chromatography



Topic	Examples
Method-operational Intent	<p>Separation of Acidic Region (AR) and Basic Region (BR) from Main Peak (MP)</p> <p>Separation of impurity peaks among each other, detectable and quantifiable</p>
Method Performance Criteria	<ul style="list-style-type: none"> • MP, AR, BR: ≤ 6.0 % RSD repeatability • MP, AR, BR: 94.0-106.0 % recovery
Range	<ul style="list-style-type: none"> • MP: $\geq 80\%$-120% of nominal protein concentration • Other components: QL- 120% of upper spec limit
Operating Conditions and Environment	<ul style="list-style-type: none"> • Suitable for HPLC platforms used in QC network • Column from established vendor and globally available • No carry over • Acceptable method performance for at least <ul style="list-style-type: none"> • two column types • 3 different resin batches • Short sample to sample run time, ≥ 48h of consecutive analyses • Preferably no harmful chemicals used

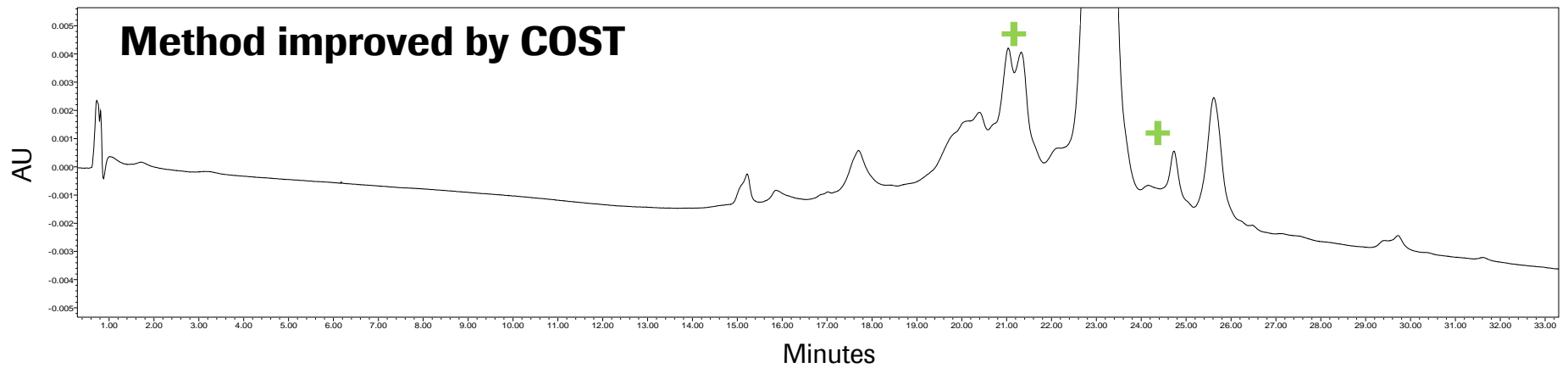
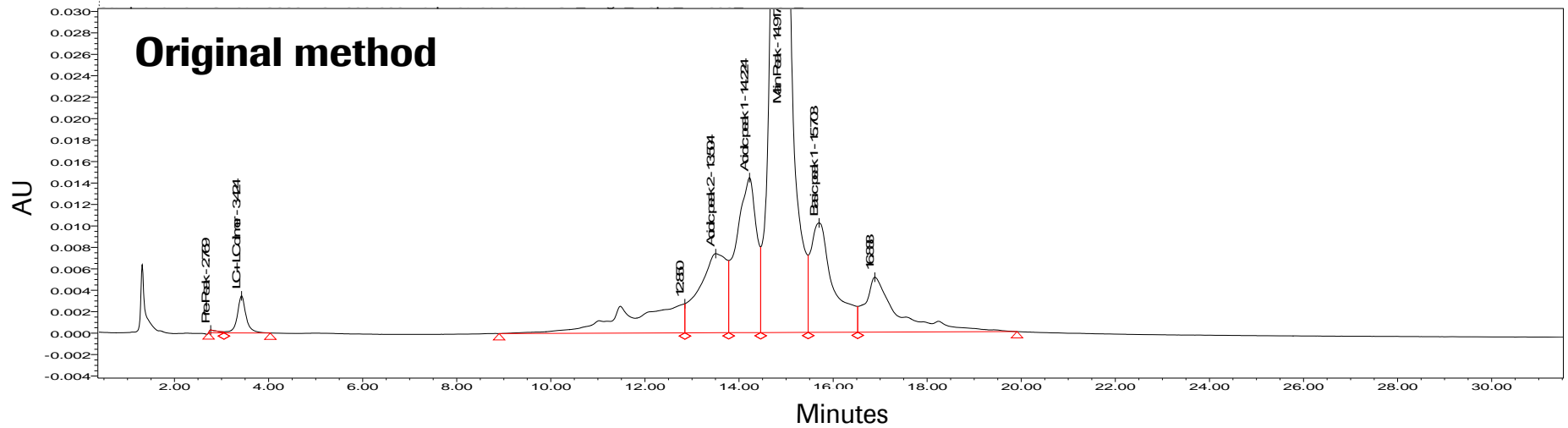
Method Development Approach

Factor	Screening with COST 	Optimization with DoE
stationary phase	<ul style="list-style-type: none"> 10 different columns (weak/strong cation exchange, source, vendor, length, particle size etc.) temperature 	<ul style="list-style-type: none"> Set to strong cation exchange (YMC, BioPro SP-F) temperature range: 28 - 45°C
mobile phase	<ul style="list-style-type: none"> buffer component (4 different buffer systems) pH range flow rate 0.7 - 1.1 salt content (up to 750 mM) 	<ul style="list-style-type: none"> buffer component: BES Buffer concentration: 15 - 25 mM pH range: 6.6 - 7.2 flow rate: 0.7 - 0.9 salt content: 488 mM NaCl
gradient	<ul style="list-style-type: none"> salt gradient gradient slope 	<ul style="list-style-type: none"> shape endpoint of slope 12 - 18%
injection		<ul style="list-style-type: none"> injection amount: 20 - 210 µg

COST = **C**hange **O**ne **S**ingle Factor a **T**ime

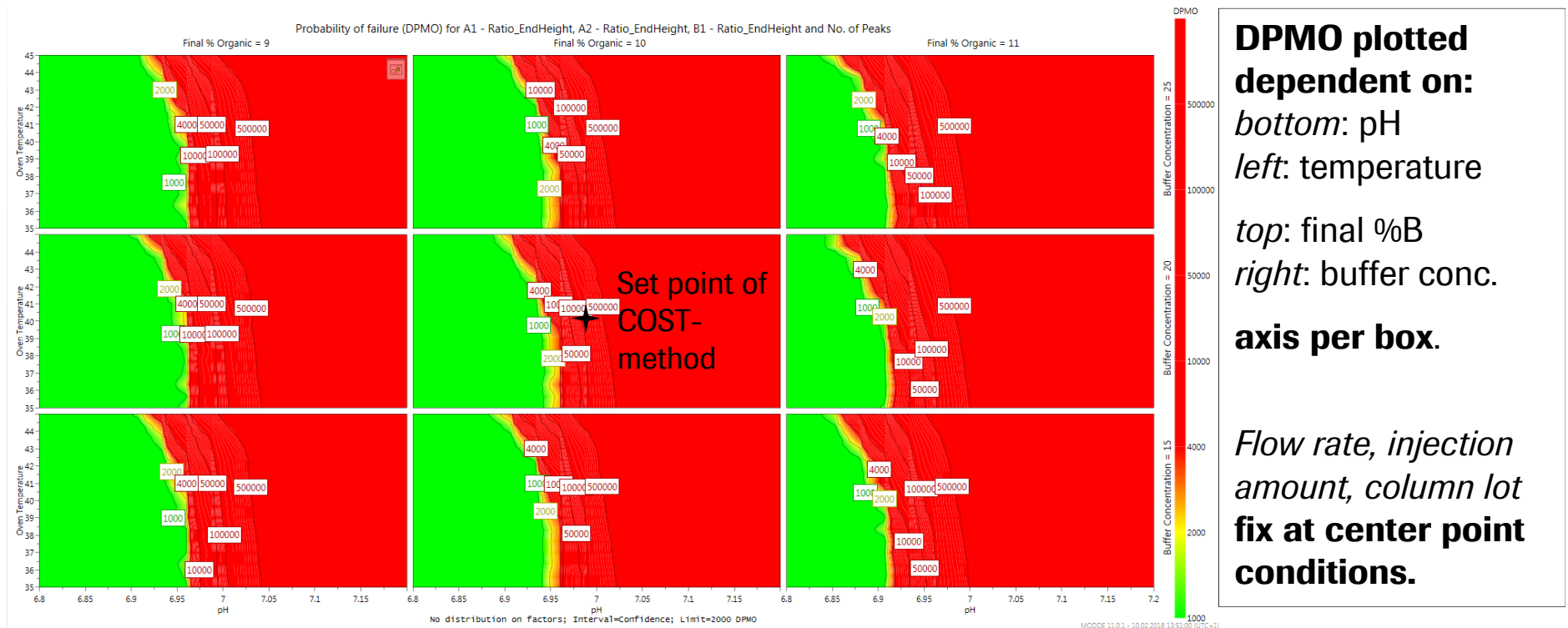
high impact factors from PHA1

Comparison Chromatograms: Original Method vs. Method improved by COST



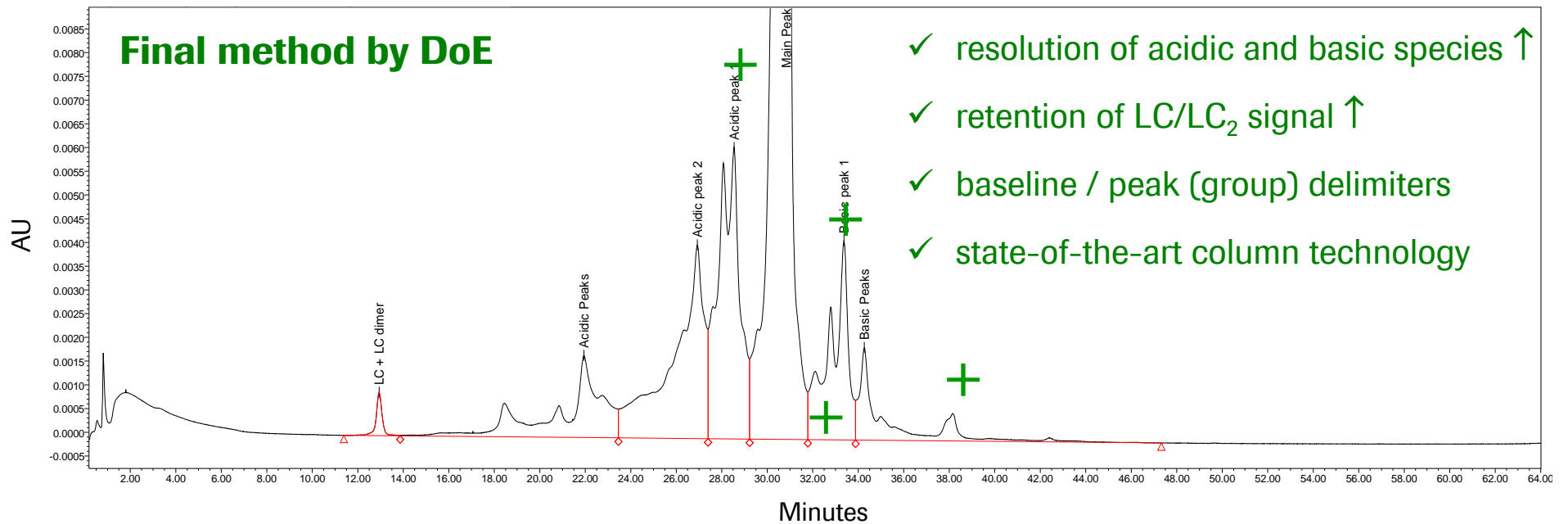
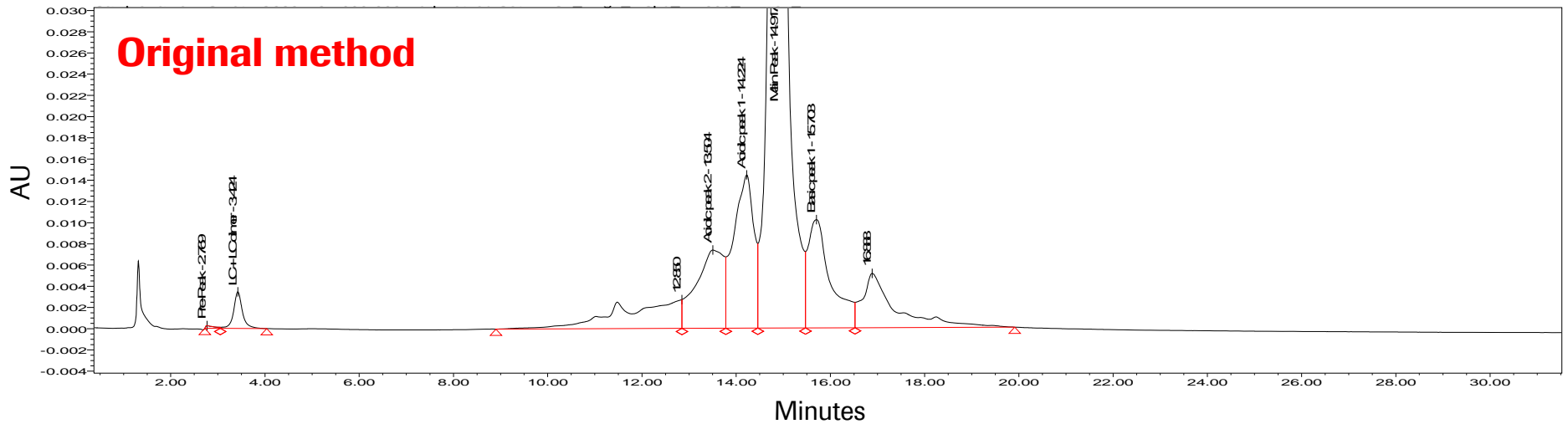
Method optimization: COST or DoE-based?

Design Space Plot

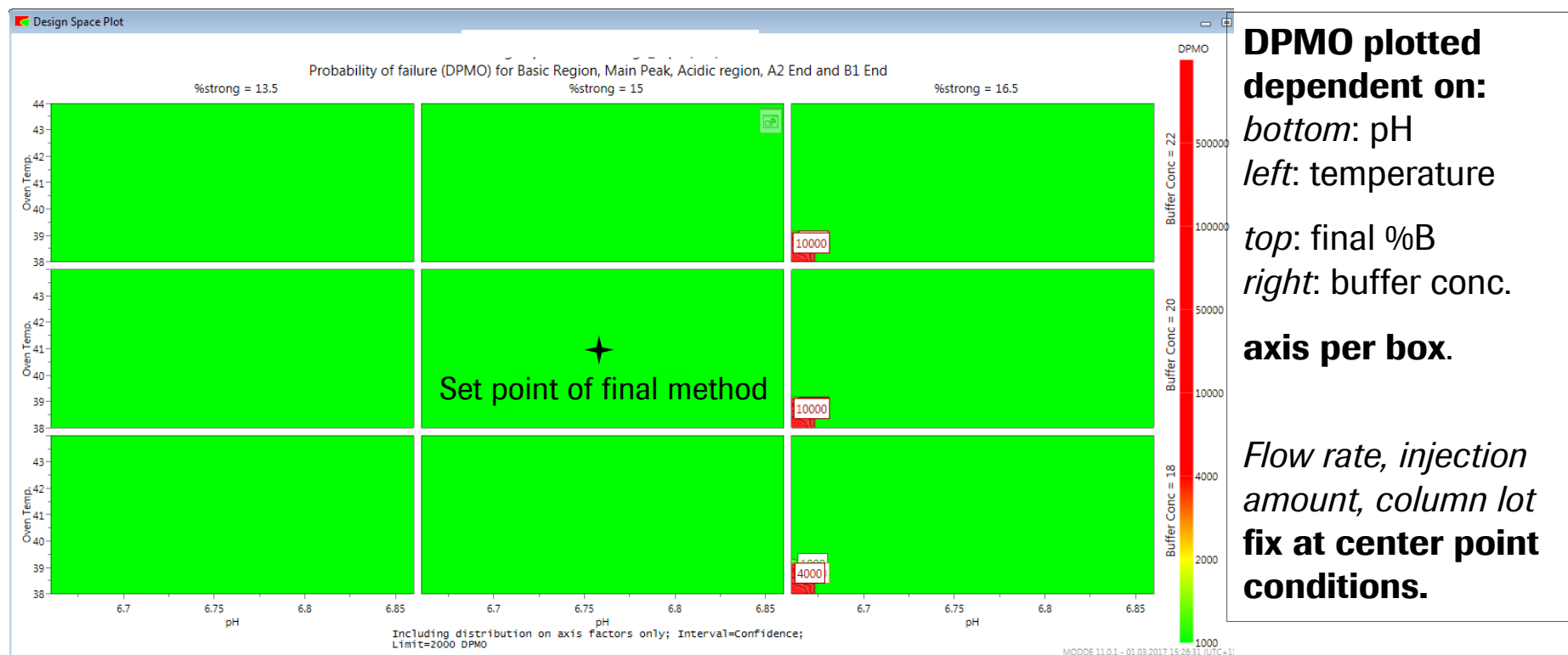


- Opportunity for further improvement elucidated by DoE modelling
- As result much better resolution could be obtained

Comparison Chromatograms: Original Method vs. Final Method improved by DOE



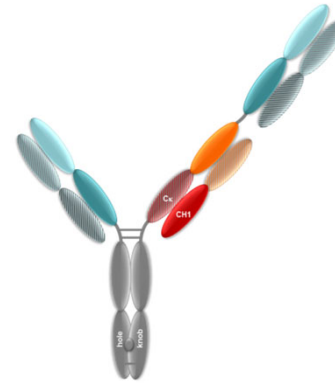
Final Method: *Design Space Plot*



- Very low probability of failure
- Highly robust method developed by using DoE

Case Study 2

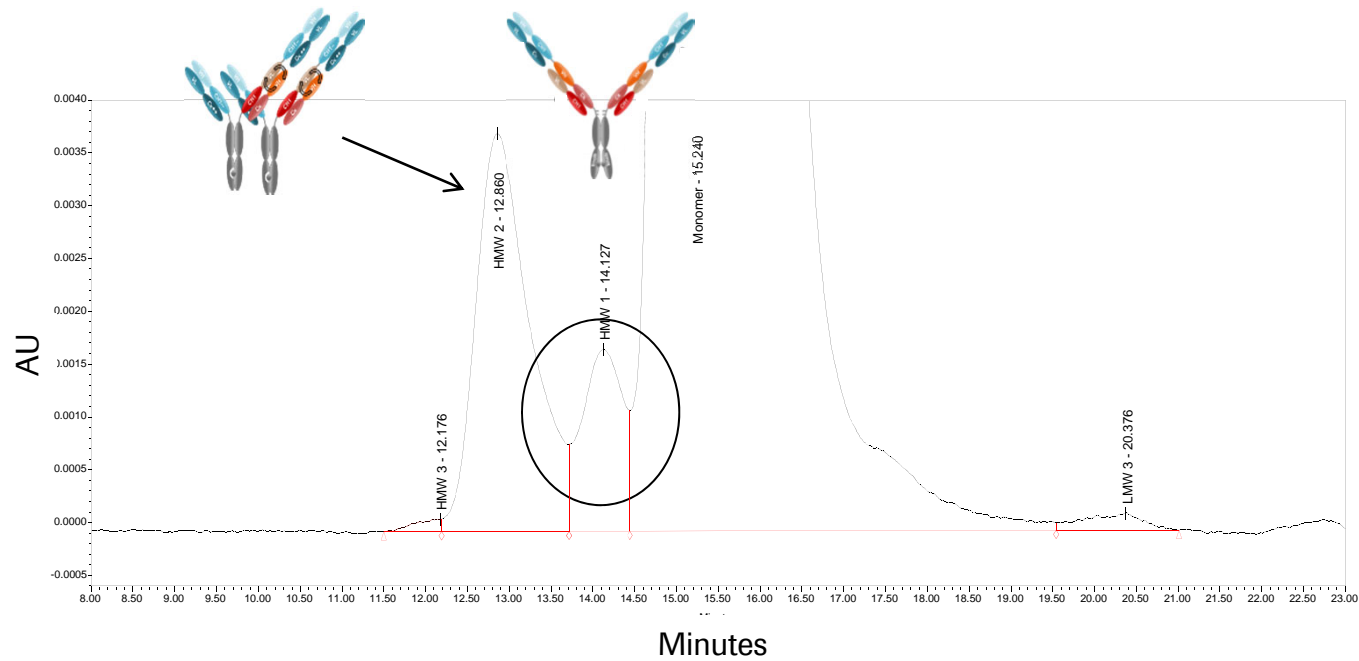
- SEC Method development for a BsAb



«Triggers» to develop a new SEC Method



Starting point: platform SEC method



- need for improved resolution of potential critical species (HMW 1, HMW 2)

Analytical Target Profile

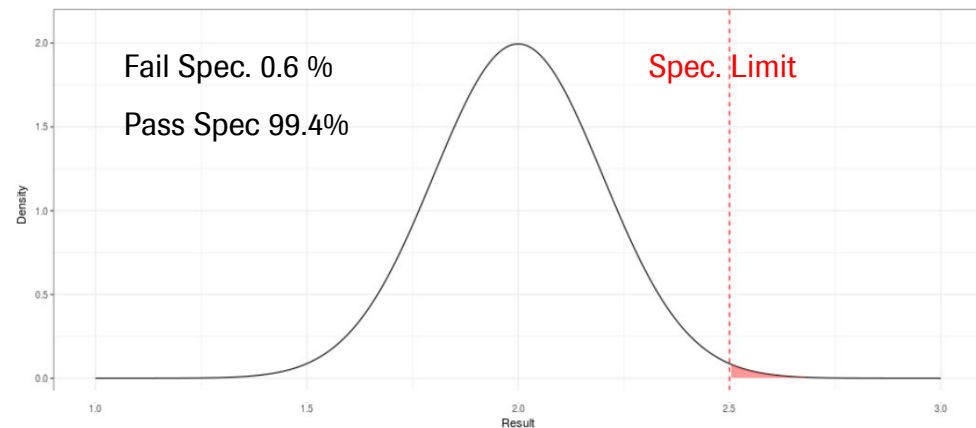
Size Exclusion Chromatography



Topic	Examples
Method-operational Intent	Separation of critical size variants i.e. High Molecular Weight species including HMW 1 from Main Peak detectable and quantifiable
Critical Quality Attribute (CQA)	<ul style="list-style-type: none"> • HMW species (e.g. HMW 1, HMW 2)
Method Performance Criteria	<ul style="list-style-type: none"> • Main Peak: ≤ 6.0 % RSD repeatability • HMWs: ≤ 0.2 SD repeatability • Main Peak: 94.0-106.0 % recovery • HMWs: assumed true value ± 0.2 (area%)
Range	<ul style="list-style-type: none"> • Main Peak: at least 80%-120% of nominal protein concentration • HMWs: QL- 3.0 %

Limitations:

Target values and process variability usually unknown at time of method development!



Probability of iOOS for Normally Distributed Data

Method Development Approach



Factor	Screening with COST	→	Optimization with DoE
Column	YMC Pack Diol 200 YMC Pack Diol 300 TSKgel Super SW mAb HR TSKgel Super SW mAb HTP TSKgel Ultra SW mAb Aggregate UPLC BEH200		UPLC BEH200
Column temperature			25°C – 45°C
Buffer composition	Sodium phosphate		
Buffer concentration	0.05 – 0.3 M		0.1 – 0.3 M
Mobile phase pH	pH range		6.2 – 7.8
Protein load	10 -150 µg		
Injection volume	2 – 30 µL		
Flow rate	0.1 – 0.5 mL/min		

Method optimization - *Design Space Plot*

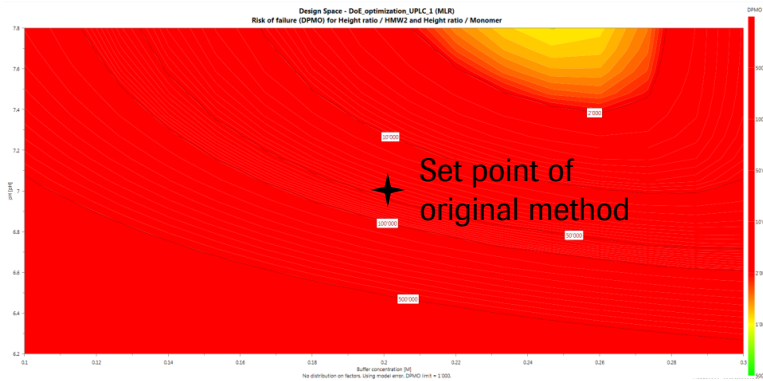


Response	Max
Height Ratio/ HMW2	0.4
Height Ratio/ Monomer	0.3

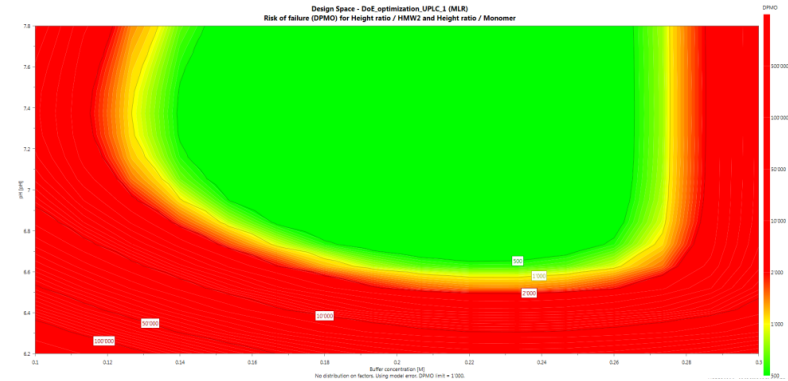
Factors at the plot axes: Low: High:

1st axis: Buffer concentration 0.1 0.3

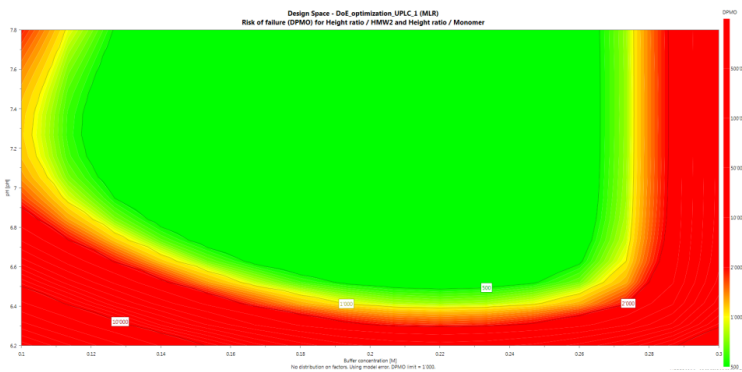
2nd axis: pH 6.2 7.8



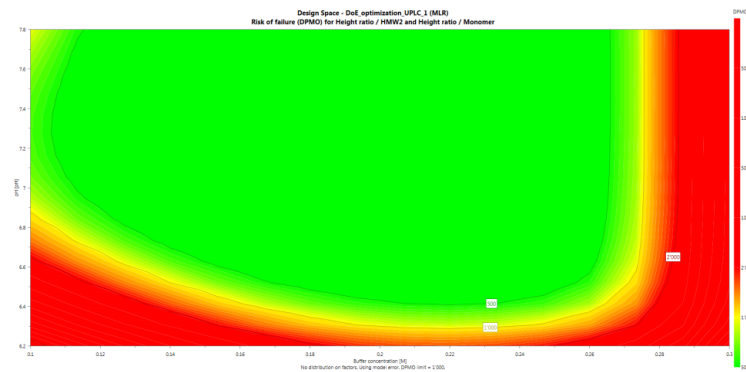
Column temperature = 25°C



Column temperature = 35°C



Column temperature = 40°C

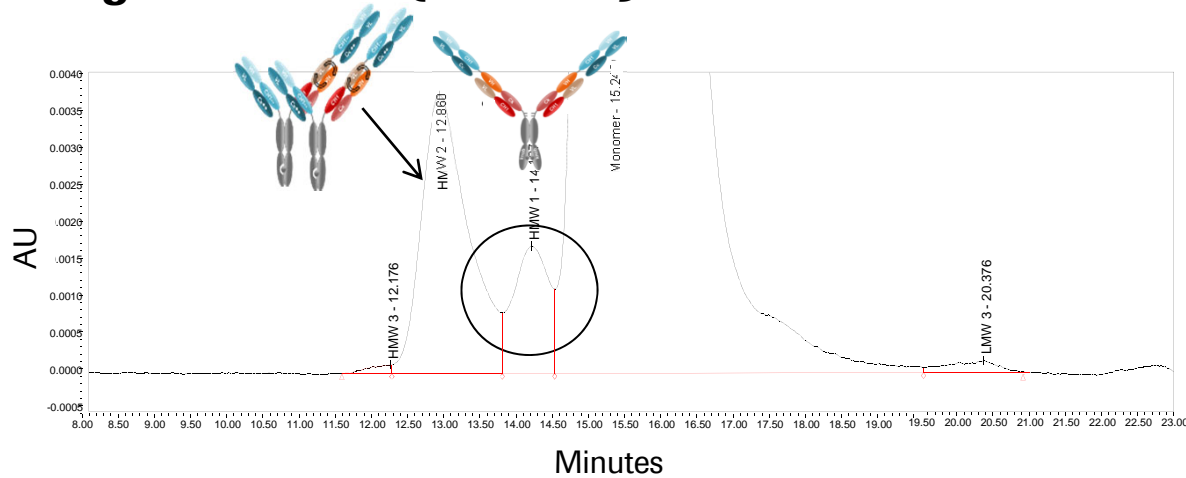


Column temperature = 45°C

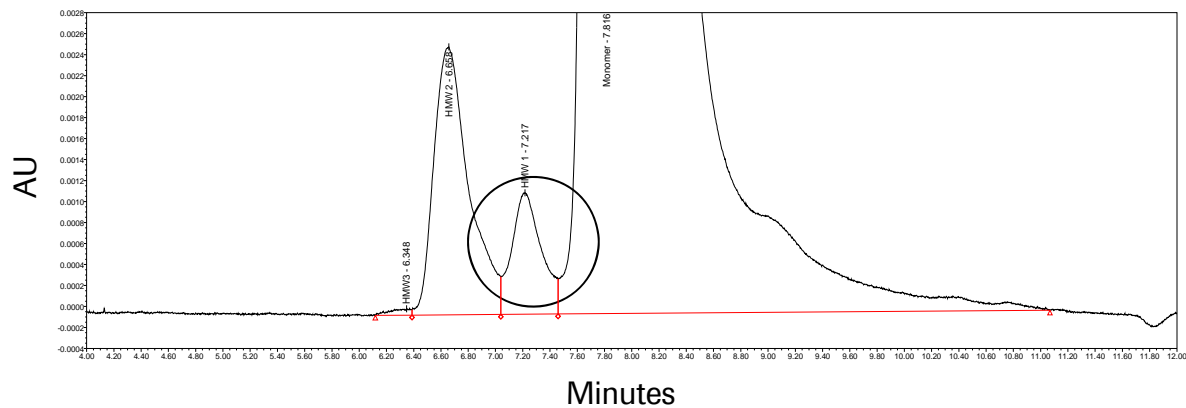
Comparison Chromatograms: Original Method vs. Final Method improved by DoE



Original method (SE-HPLC)



Final method (SE-UPLC)



Modified factors

1. Column material & dimensions
2. Column temperature
3. Buffer concentration
4. Salt concentration
5. Flow rate
6. pH

Benefits of Enhanced Method Development

- AQbD tools provide a science- and risk-based framework for developing enhanced understanding of analytical methods
- Structured risk assessments are used to guide experimental plans
- Enhanced method robustness and ruggedness through the lifecycle
- Less analytical method related Out-of-Specification and failure investigations
- More robust method knowledge transfer due to enhanced analytical method understanding



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- Mahdi Adibzadeh



Doing now what patients need next