



# Electronic Chromatogram Review at Biogen Idec: Past, Present, and Future

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# Outline

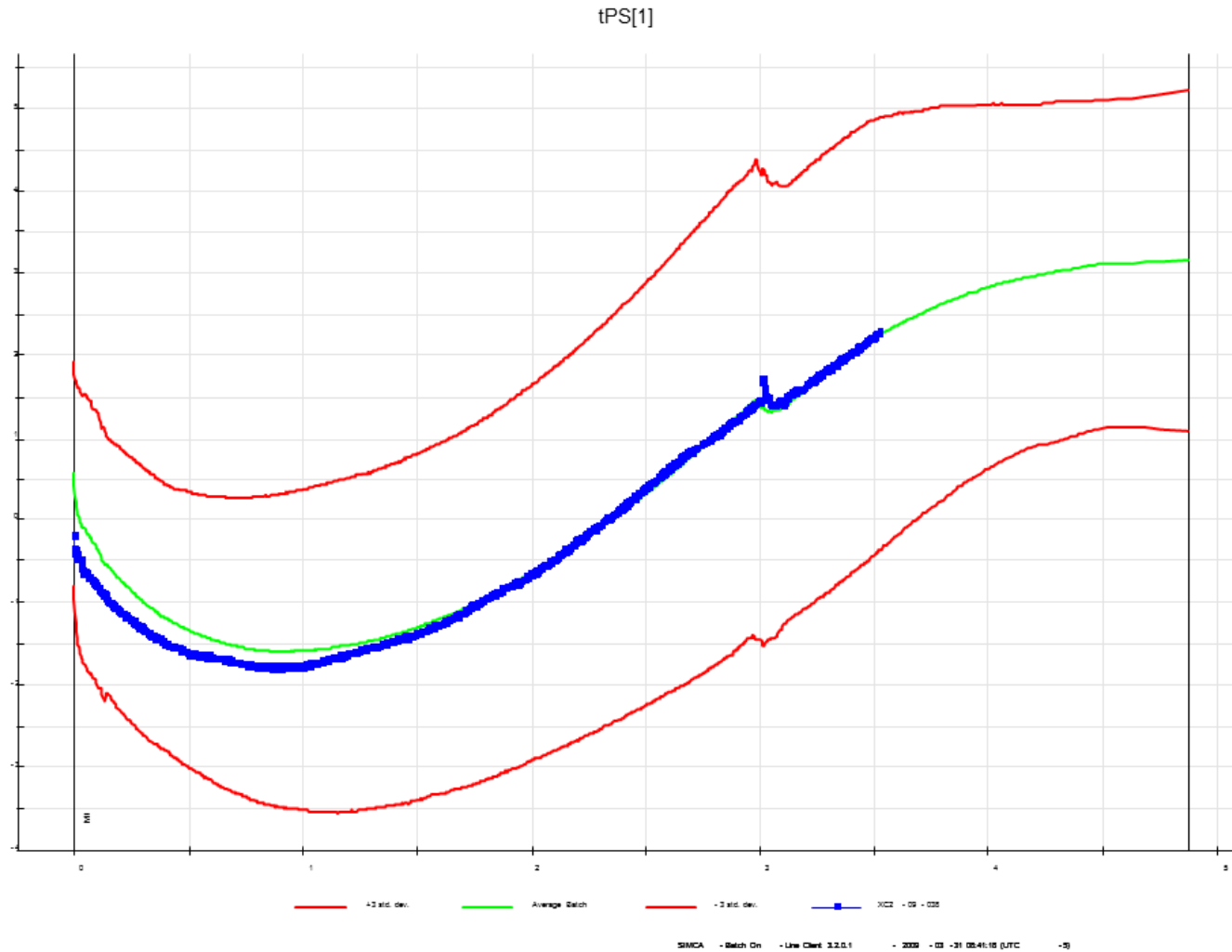
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- Introduction
- Biogen Idec Multivariate Analysis Journey
- Case Studies
- Summary

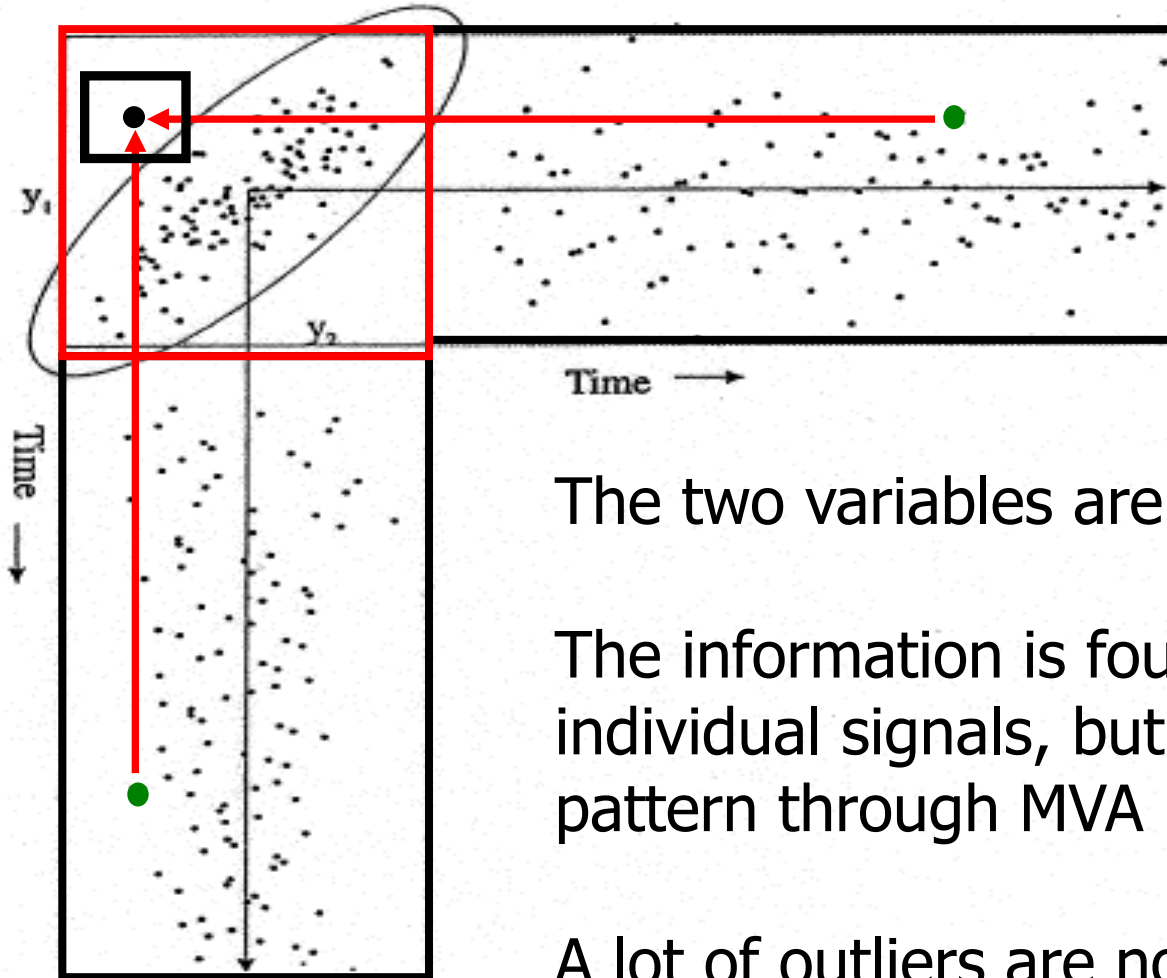




# This Looks Better



# Multivariate Analysis: What & Why



The two variables are correlated

The information is found NOT in the individual signals, but in the correlation pattern through MVA

A lot of outliers are not detected unless all the variables are analysed together

# Biogen Idec Advanced Process Control

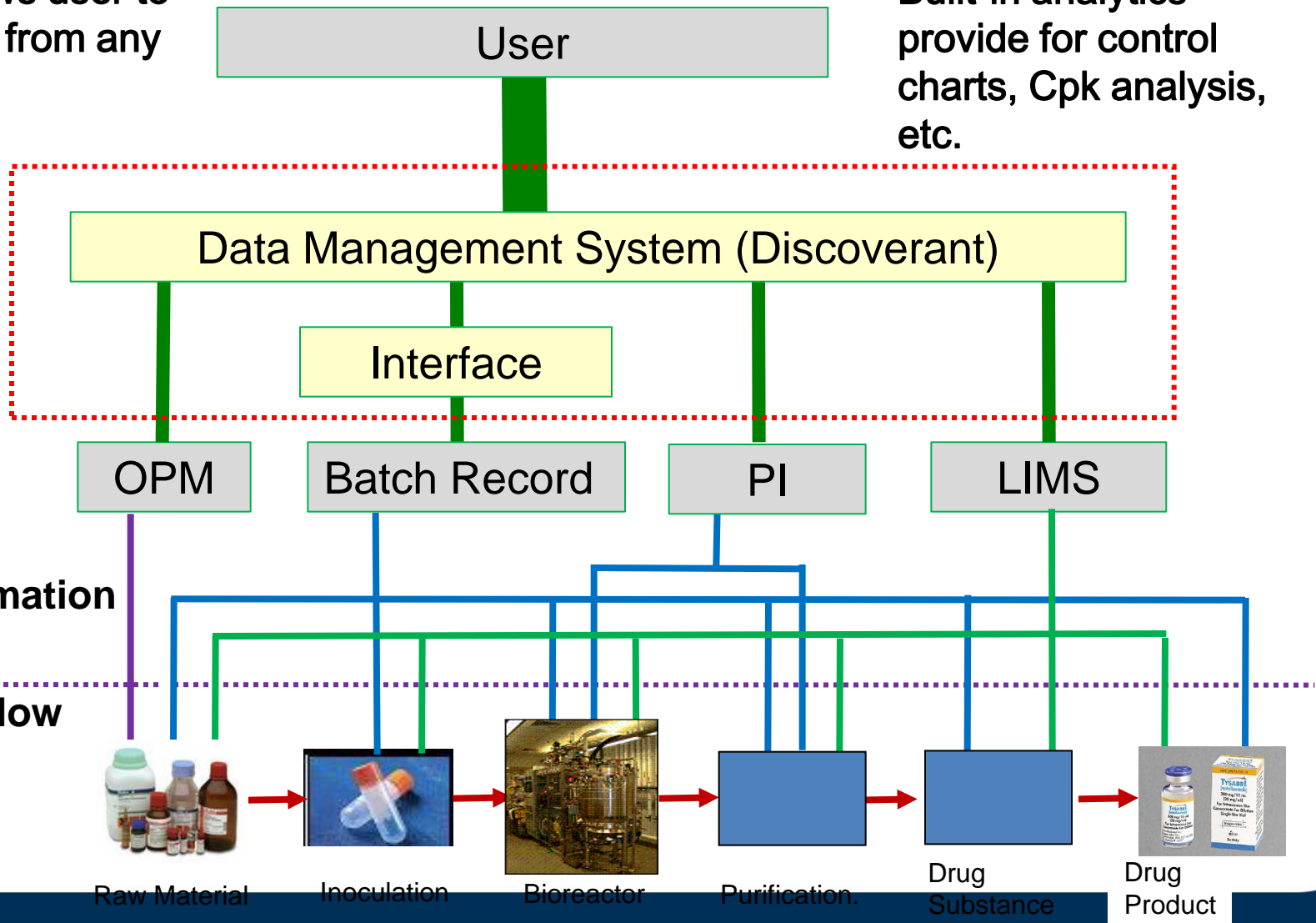
- Established an [infrastructure](#) for data centralization and real time process monitoring
  - Achieved [full scale](#) utilization of Discoverant and SBOL
    - 2003: 1<sup>st</sup> Discoverant Hierarchy & 1<sup>st</sup> Cell Culture SBOL
    - 2006: SBOL on Manufacturing Floor
    - 2007: 1<sup>st</sup> Purification SBOL model
    - 2013: Approved patent on “Systems and Methods for Evaluating Chromatography Column Performance”, US008410928B2, 02Apr2013
- Built a [culture](#) of advanced process monitoring
  - Shift Trend Review + Routine Trend Review



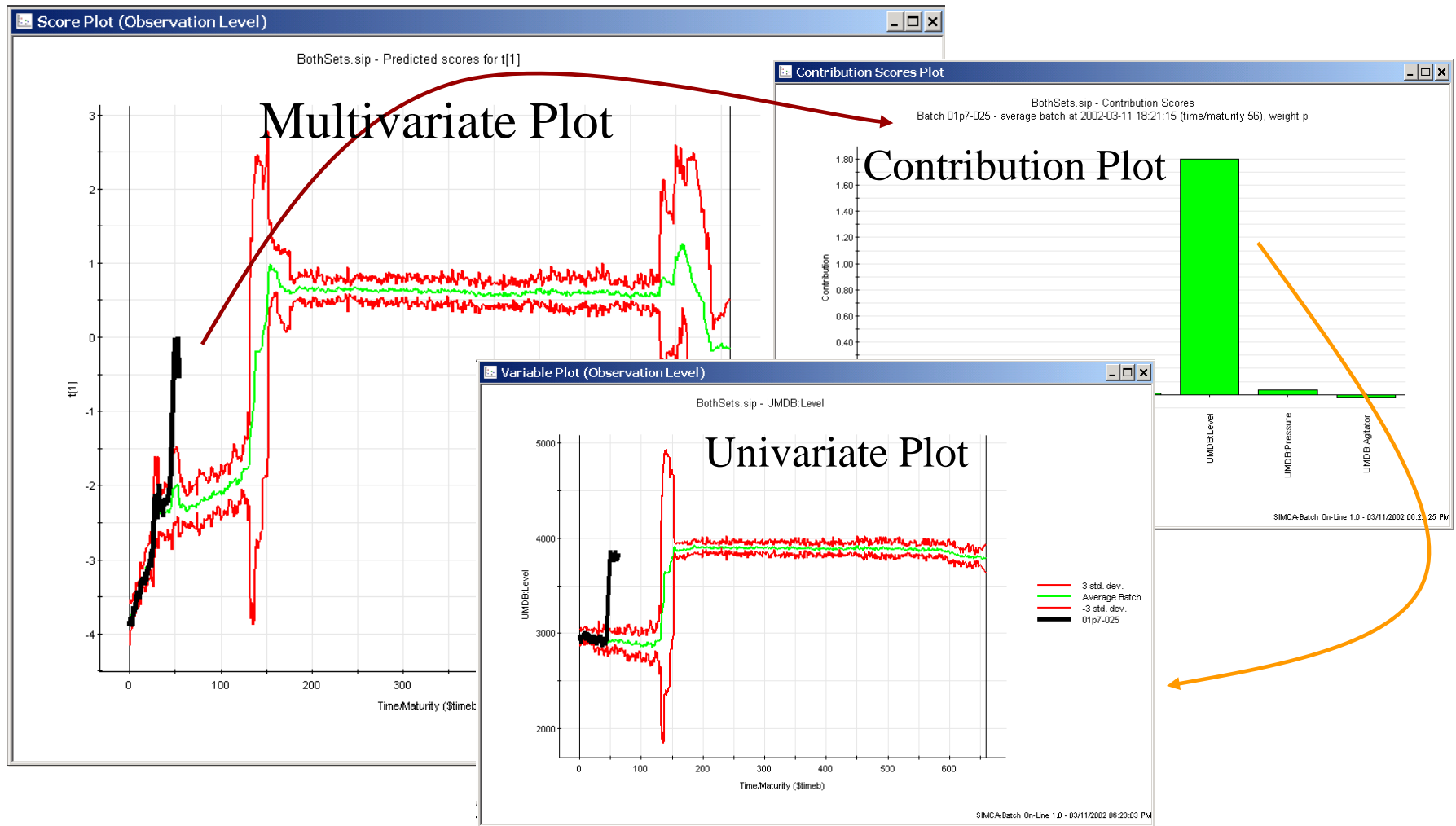
# Centralized Data Management System

System allows user to access data from any system

Built-in analytics provide for control charts, Cpk analysis, etc.



# Multi-variate Monitoring System - SBOL

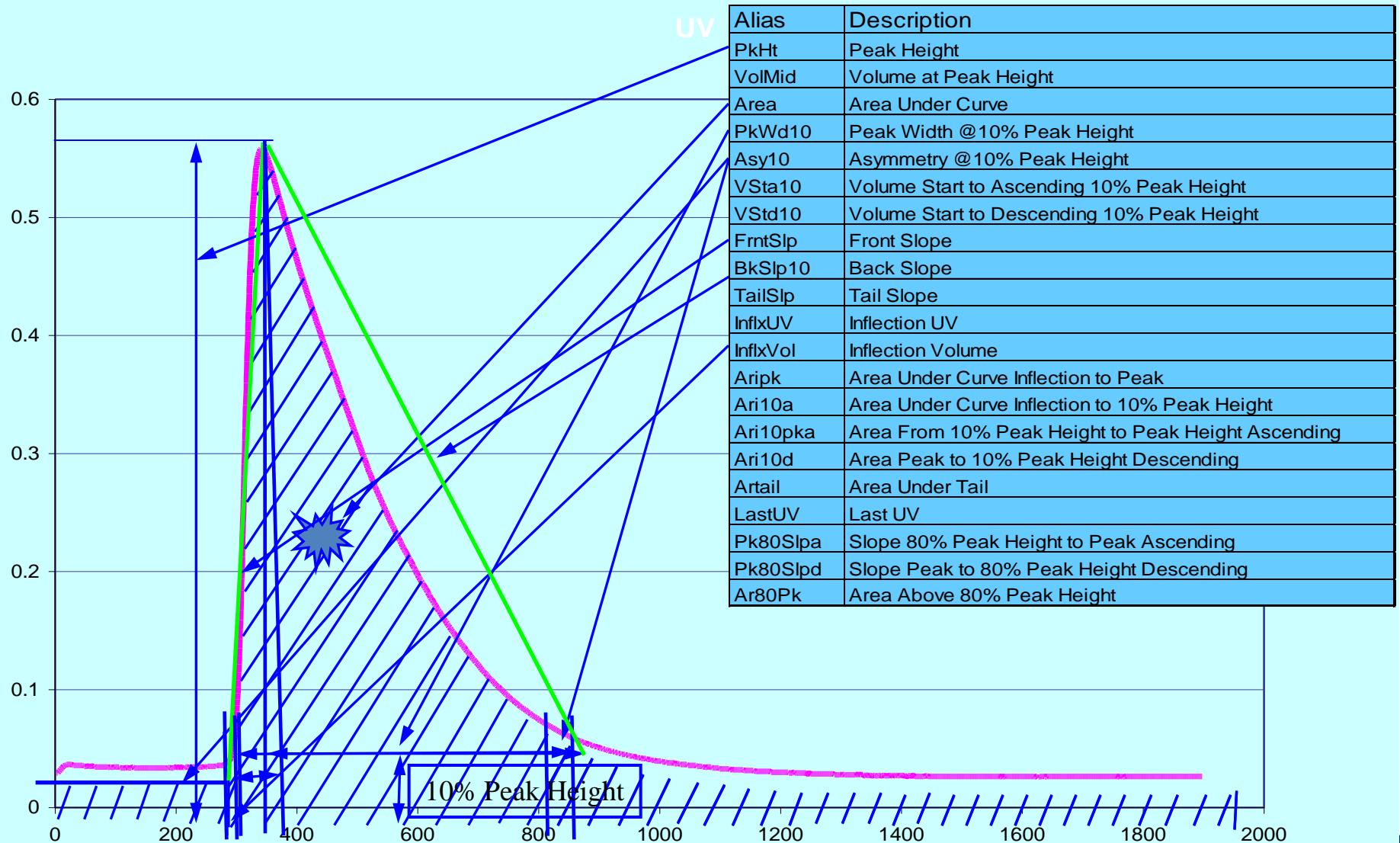


# Multivariate Chromatogram Analysis

- Motivation
  - Chromatograms contain a lot of information that we are not utilizing
  - Exploit already available continuous data
    - UV, conductivity, pH, pressure, volumetric flow
- Goals
  - Non-subjective, quantitative method for evaluating chromatograms
- Example: Analysis of elution peak UV tracing
  - MVA model using discrete parameters that describe the characteristics of the elution UV peak

# MVA of Chromatograms

## Elution Peak Parameters



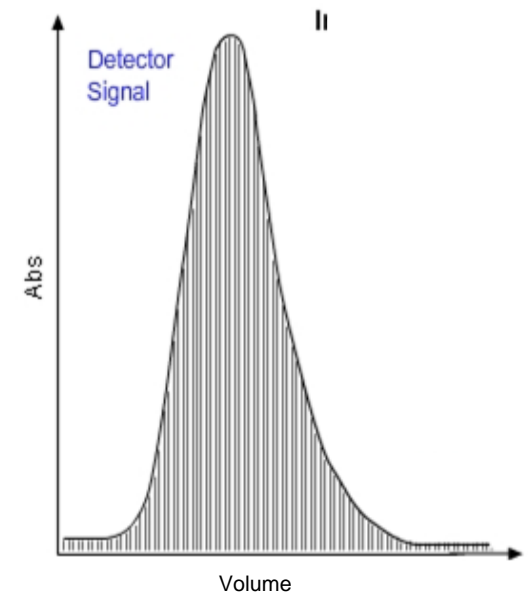
# Case Study #1: AUP

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- Area under peak (AUP) of elution UV peak was proposed as an alternative to quantify protein concentration
- Ratios of AUP to total offline A280 across manufacturing sites were shown to be different
- Investigation was Launched
  - To understand root cause and harmonize practice

# Background

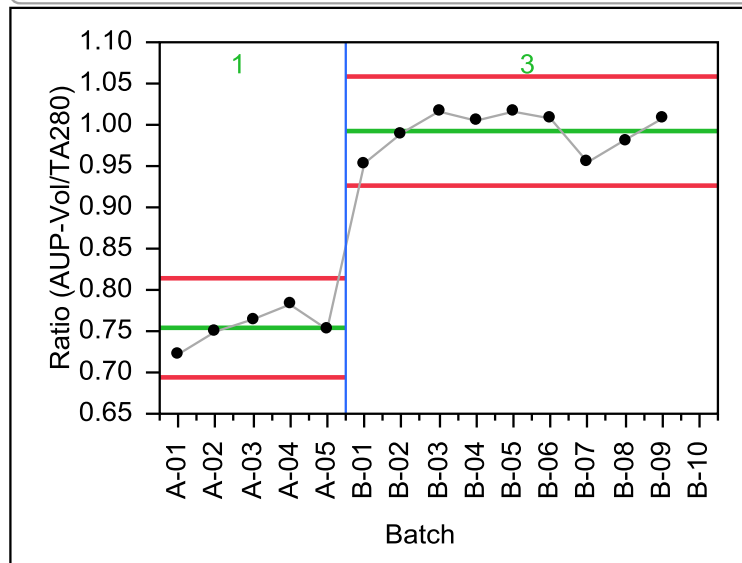
- Current practice: offline sampling for TA280
- TA280 information is contained in the elution peak
  - $\int A_{280} dV = \text{TA}_{280}$
  - Concept has been used in the past to assess incorrect A280 sampling deviations
  - Assumption: A280 signal is proportional to protein concentration



# Inconsistent Ratios X-Site (Before)

## Control Chart

### Individual Measurement of Ratio (AUP-Vol/TA280)

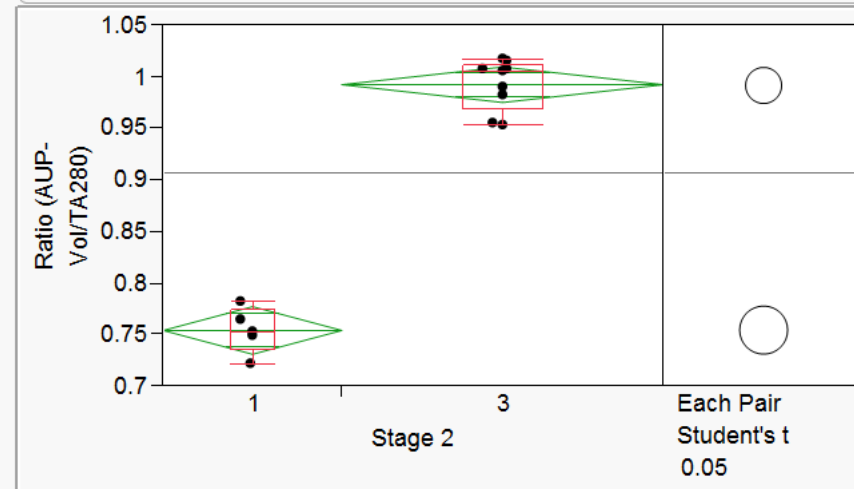


## Phase Limits

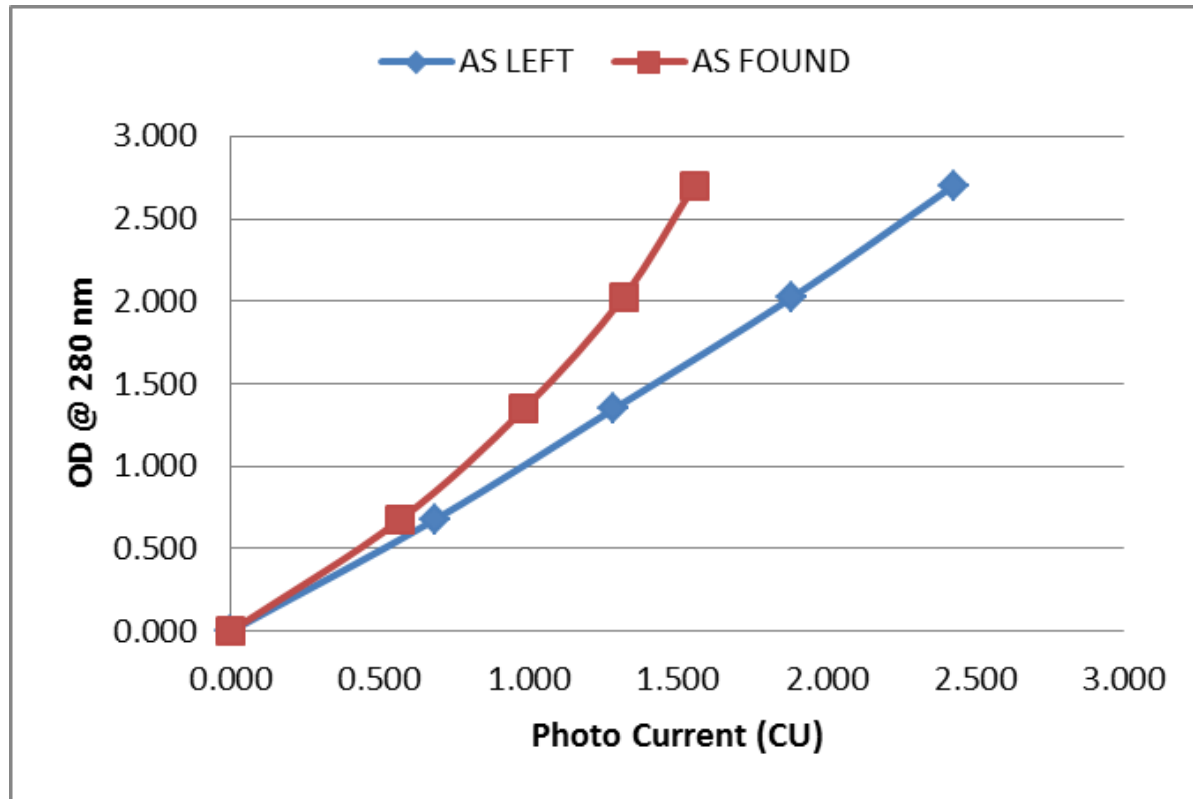
Phase	LCL	Avg	UCL
1	0.694062	0.754089	0.814115
3	0.926336	0.992396	1.058455

- 1: Data from site A per “As Found” linearization curve
- 3: Data from site B – Linearization curve normal

## Oneway Analysis of Ratio (AUP-Vol/TA280) By Stage 2



# Root Cause - Linearization Curve Not Linear

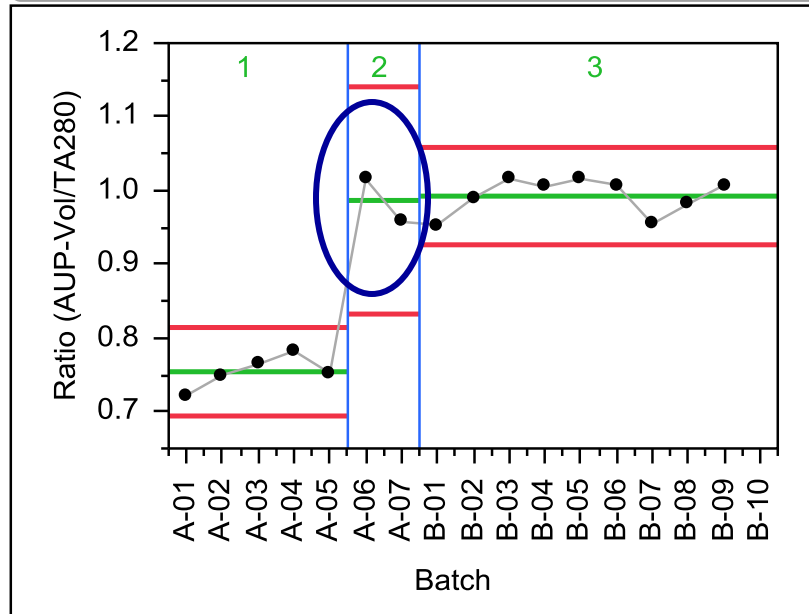


- “As Found” curve was not linear
- “As Left” curve returned to linear after replacing interference filter

# Consistent AUP/TA280 X-Site (After)

## Control Chart

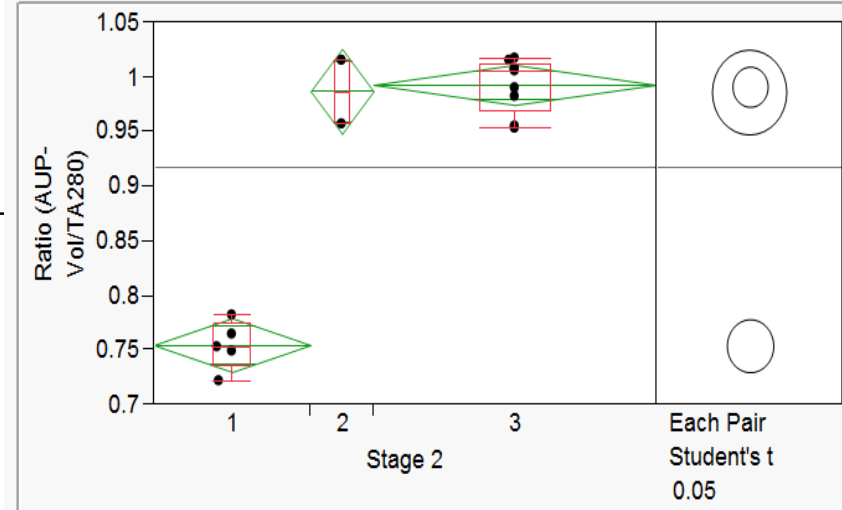
### Individual Measurement of Ratio (AUP-Vol/TA280)



### Phase Limits

Phase	LCL	Avg	UCL
1	0.694062	0.754089	0.814115
2	0.832367	0.986524	1.140681
3	0.926336	0.992396	1.058455

### Oneway Analysis of Ratio (AUP-Vol/TA280) By Stage 2



1. Data from site A per "As Found" linearization curve
2. Data from site A per "As Left" linearization curve
3. Data from site B – Linearization curve normal

# Lessons Learned

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- Area under peak of elution could add significant value to real time chromatography performance monitoring
- Question: Is UV representative of protein concentration?
- Answer: Correct linearization table is critical to UV meter performance and success of Chromatogram analysis (including AUP)

# Case Study #2 – Elution UV Chromatogram Analysis

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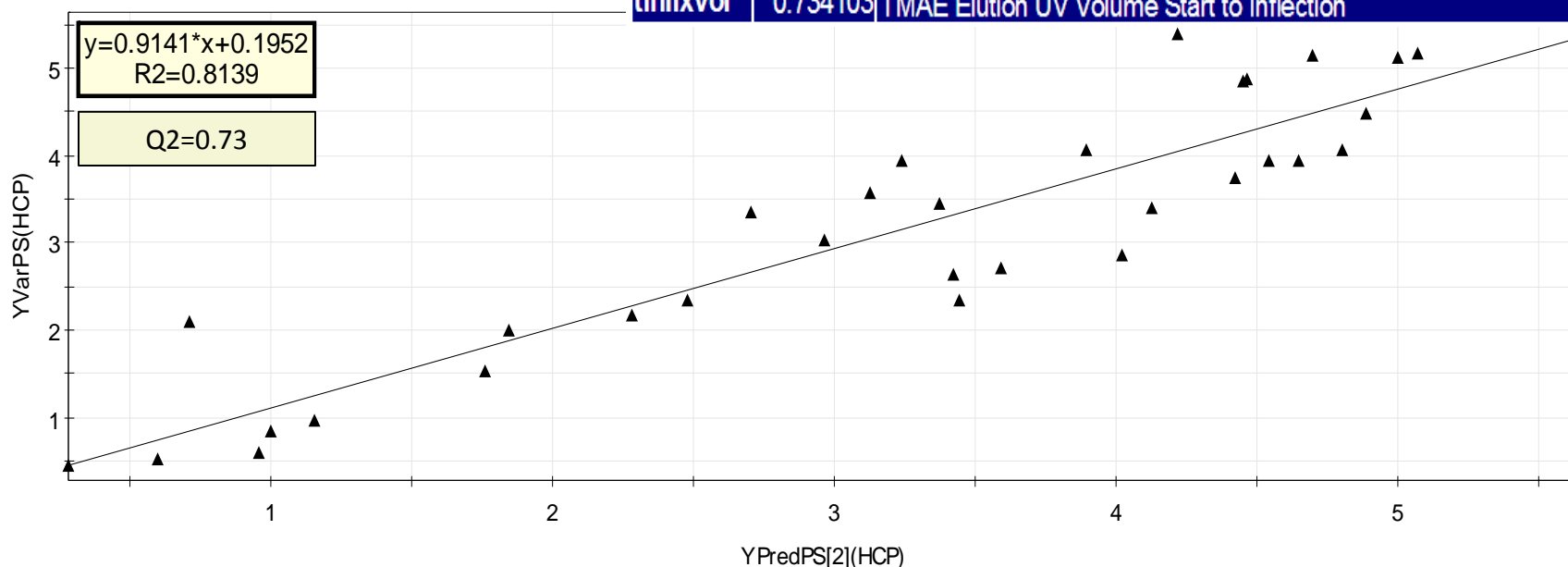
- Residual host cell protein was showing a gradual upward trend in the drug substance
- ALEX step (column 2) was suspected root cause
- Investigation was launched
  - Started by building a multivariate process model (approximately 60 variables covering cell culture and purification)

# AIEX PLS Model to Predict HCP

- Eight chromatogram parameters predicts well HCP level in the Drug Substance

Var ID (Primary)	M10.VIP[2]	
tFmtSlp	1.16213	TMAE Elution UV Front Slope
tPkHt	1.14422	TMAE Elution UV Peak Height
tBkSlp10	1.13214	TMAE Elution UV Back Slope (Pk to 10% Pk Height Decending)
tVolMid	1.00968	TMAE Elution UV Volume Start to Peak Max
tVStd10	0.961922	TMAE Elution UV Volume Start to 10% Peak Height Decending
tPkWd10	0.944189	TMAE Elution UV Volume @ 10% Peak Height
tTailSlp	0.826636	TMAE Elution UV Slope 10% Peak Height Decending to End
tInflxVol	0.734103	TMAE Elution UV Volume Start to Inflection

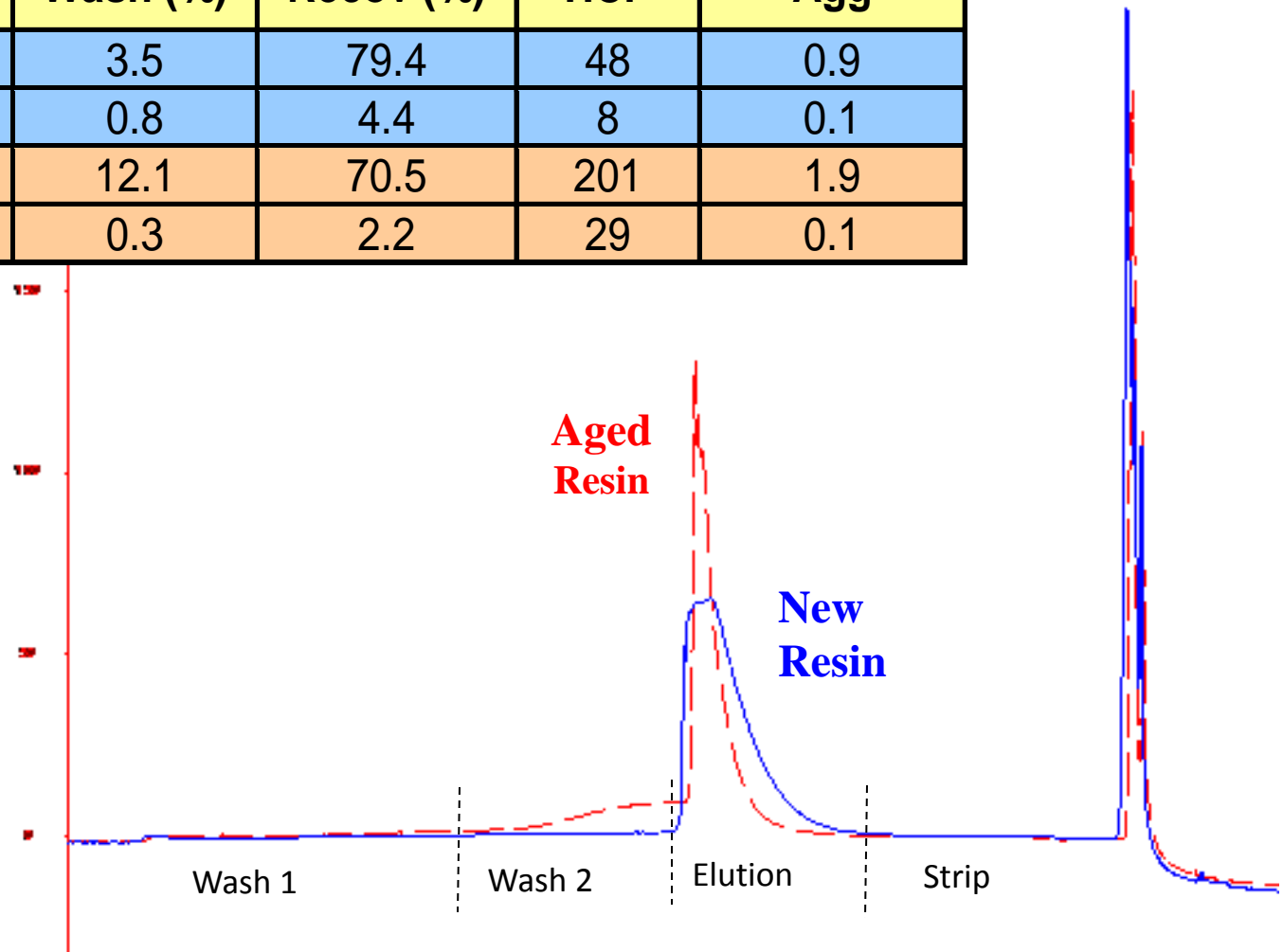
HCP Model2.  
YPredPS[Las



Further correlation analysis indicated: Chromatogram shapes changed as resin aged

# Confirmation via Lab Studies: Change in Chromatogram Shape due to Resin Age

Resin		Wash (%)	Recov (%)	HCP	Agg
New	Average	3.5	79.4	48	0.9
	S.D.	0.8	4.4	8	0.1
Aged	Average	12.1	70.5	201	1.9
	S.D.	0.3	2.2	29	0.1



# Chromatogram MVA Summary

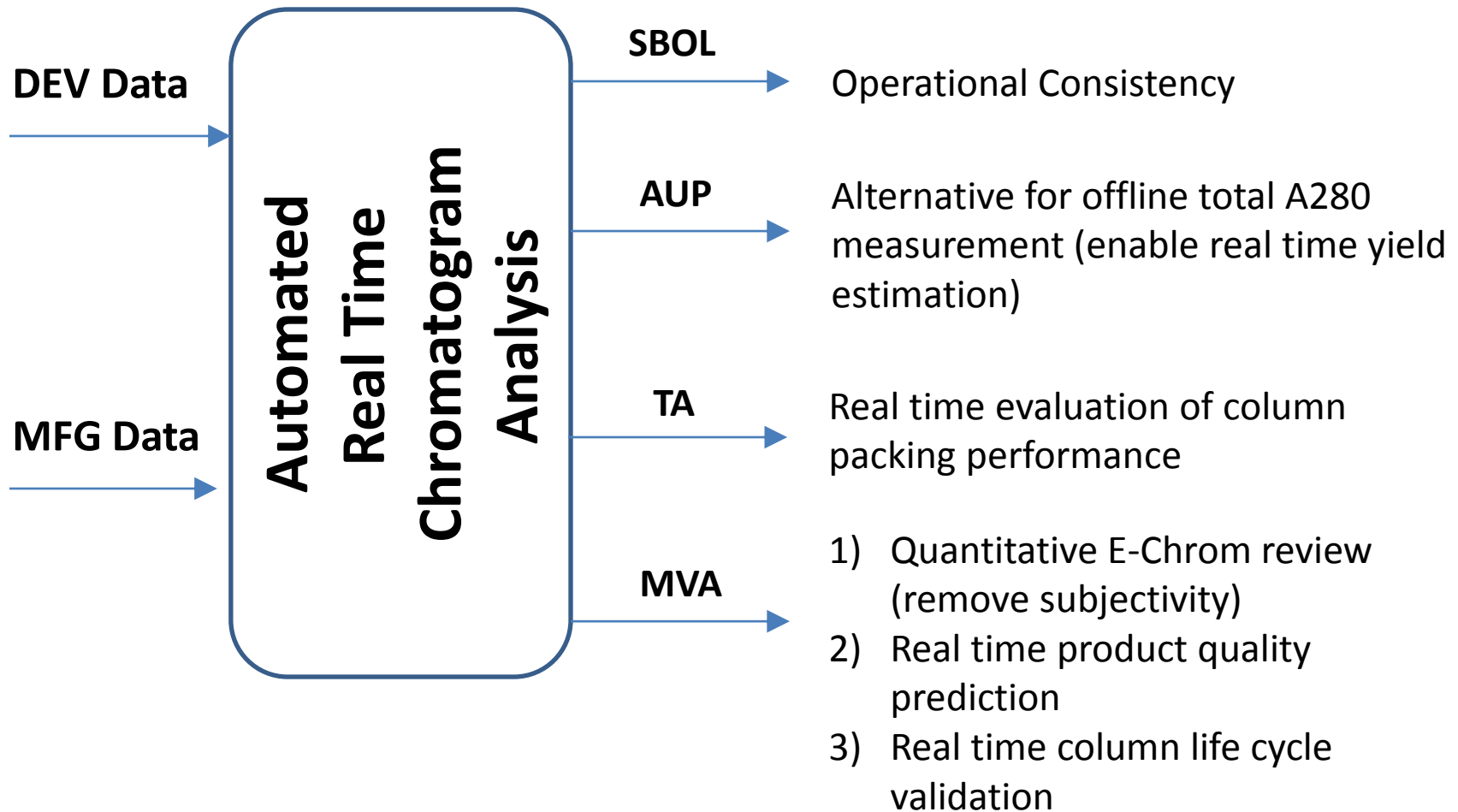
- Chromatogram MVA enabled successful root cause investigation of the HCP upward shift
  - Multivariate analysis of cell culture and chromatogram parameters indicated that chromatogram parameters were strongly correlated with HCP
  - Further correlation analysis indicated that resin age contributed to the chromatogram shape shift
  - Lab studies confirmed that resin age contributed to the column performance shift and the gradual upward trend of HCP

# Summary and Conclusions

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- Biogen Idec's E-Chrom Review
  - Our Goal is to develop non-subjective, quantitative method for evaluating chromatograms
- Data Accuracy is Key
  - Adequate instrument calibration/maintenance ensures accurate data and successful electronic chromatogram analysis
- Multivariate Chromatogram Review
  - More objective and quantitative than visual inspection
  - Relates chromatogram characteristics to column performance and product quality attributes

# Future of Chromatogram Review at Biogen Idec



## Challenges:

Equipment design, Instrument, and interaction of solution to resin



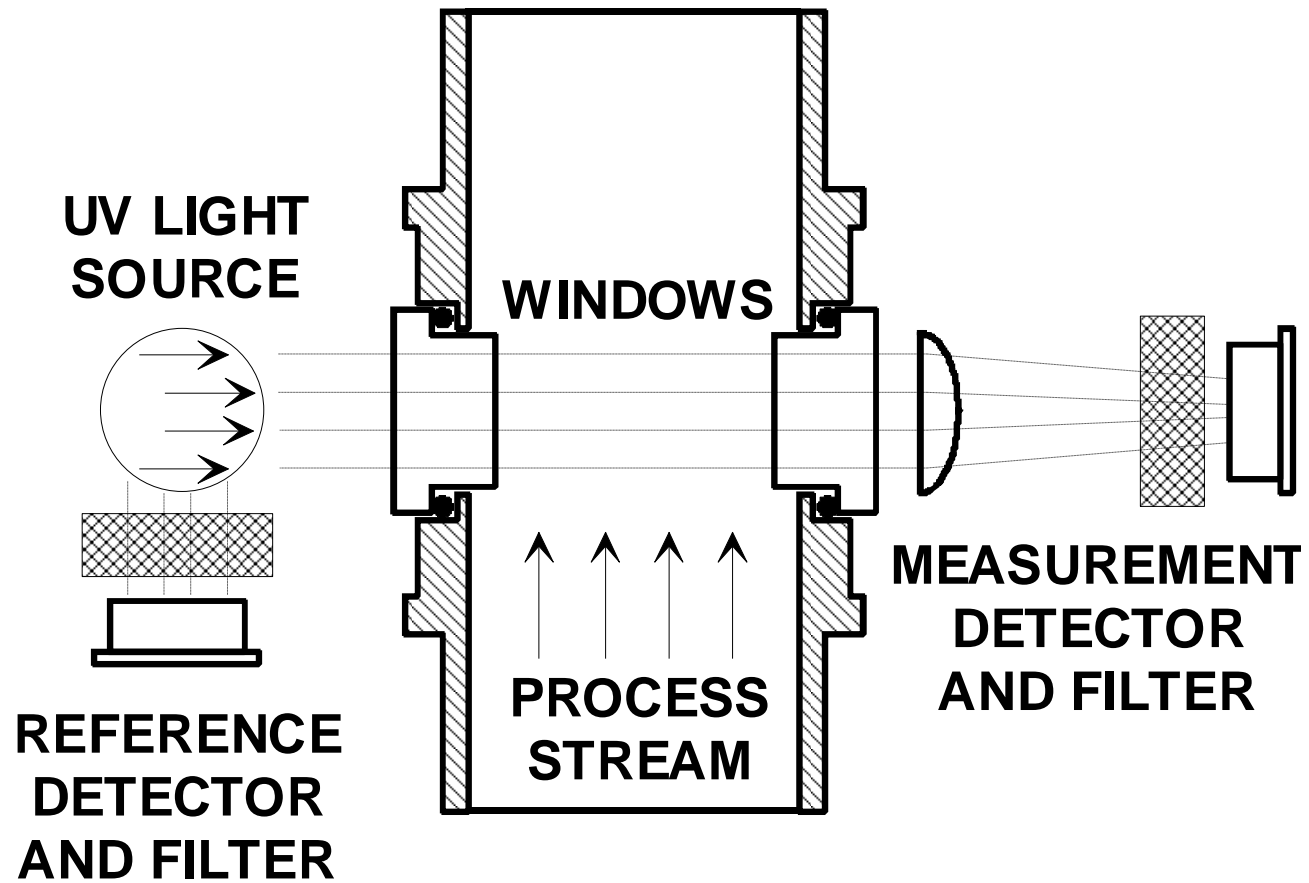
# Acknowledgement

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- Doug Cecchini
- Joydeep Ganguly
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- Ben Gilbert
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- Jeff Simeone
- Jorg Thommes
- Andre Walker
- Sarah Yuan

# Excess slides

# UV Absorbance



# MVA Principal Component Analysis

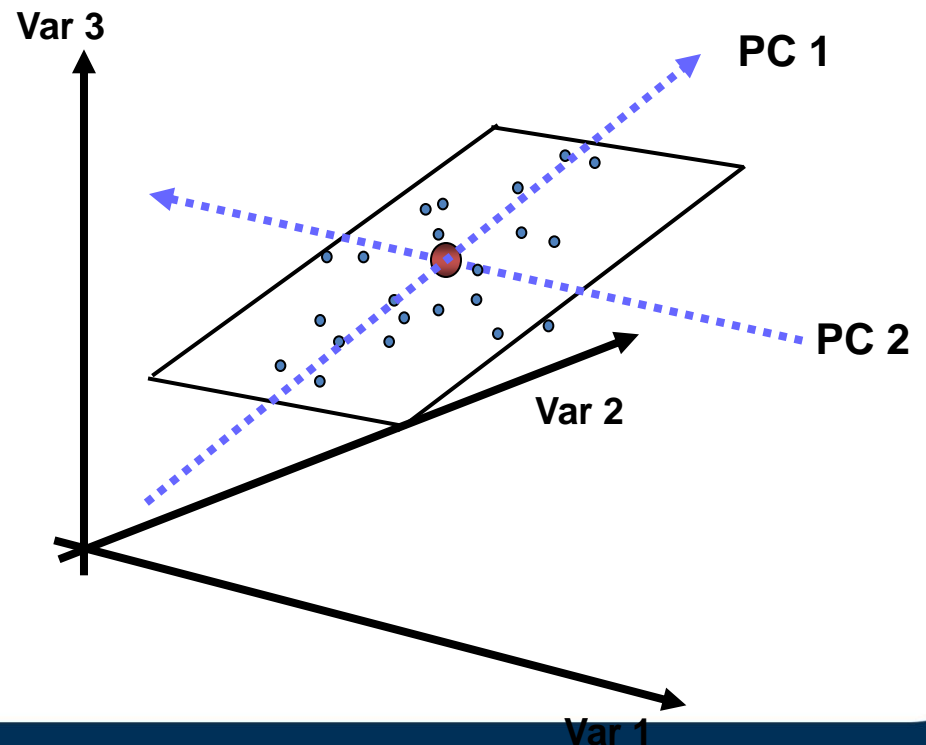
Variables					
N	1	$X_1$	$X_2$	$X_3$	$X_4$
	2	$X_1$	$X_2$	$X_3$	$X_4$
	3	$X_1$	$X_2$	$X_3$	$X_4$
	4	$X_1$	$X_2$	$X_3$	$X_4$
	5	$X_1$	$X_2$	$X_3$	$X_4$
	6	$X_1$	$X_2$	$X_3$	$X_4$
	7	$X_1$	$X_2$	$X_3$	$X_4$
	8	$X_1$	$X_2$	$X_3$	$X_4$
	9	$X_1$	$X_2$	$X_3$	$X_4$
	10	$X_1$	$X_2$	$X_3$	$X_4$

Scores	
$t_{1-1}$	$t_{2-1}$
$t_{1-2}$	$t_{2-2}$
$t_{1-3}$	$t_{2-3}$
$t_{1-4}$	$t_{2-4}$
$t_{1-5}$	$t_{2-5}$
$t_{1-6}$	$t_{2-6}$
$t_{1-7}$	$t_{2-7}$
$t_{1-8}$	$t_{2-8}$
$t_{1-9}$	$t_{2-9}$
$t_{1-10}$	$t_{2-10}$

- Data is scaled and centered
- Plotted in K dimensional space
- Fit 1st Principal components (least squares)
- Fit 2nd Principal components (least squares)
- Project data onto two dimensional plane

Weight	$P_{1-1}$	$P_{1-2}$	$P_{1-3}$	$P_{1-4}$
	$P_{2-1}$	$P_{2-2}$	$P_{2-3}$	$P_{2-4}$

SIMCA-P+ (Umetrics)

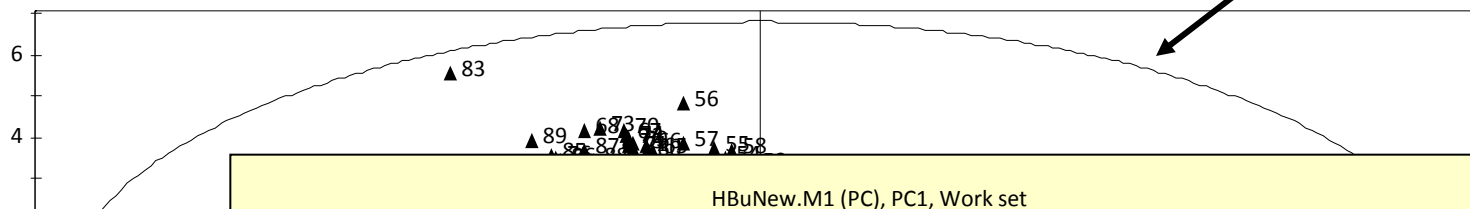


# Score Plot (t1 vs t2)

HBuNew.M1 (PC), PC1, Work set

Scores:  $t[1]/t[2]$

95 % Confidence  
limit



HBuNew.M2 (PLS), PLS1, Workset  
Contribution Scores, AVG-Obs48, Dif X scaled, weight= $w^*$ , Comp2

