

Agilent 1290 Infinity II 2D-LC Solution Biopharmaceutical Polymer Analysis

> WCBP Jan 2017 Washington, DC

#### **Overview**

- Resolving power and how to measure it
- Why two-dimensional LC?
- Setup of a 2D-LC System
- Different modes of 2D-LC
  - Heart-Cutting 2D-LC
  - Multiple Heart-Cutting 2D-LC
  - High Resolution Sampling 2D-LC
  - Comprehensive 2D-LC
- One software for all 2D-LC modes
- Online 2DLC for Biopharmaceuticals 4 case studies

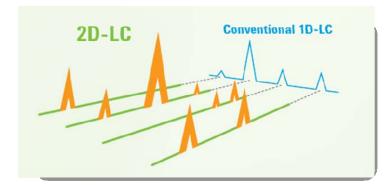


#### Increasing Lab Efficiency UHPLC and Other Solutions





- Automation
  - Automated sample preparation
  - Valve solutions
- Increasing peak capacity
  - Longer columns
  - Online 2D-LC





#### Why Two-Dimensional LC? General Thoughts in Separation Science



JOURNAL OF CHROMATOGRAPHY A

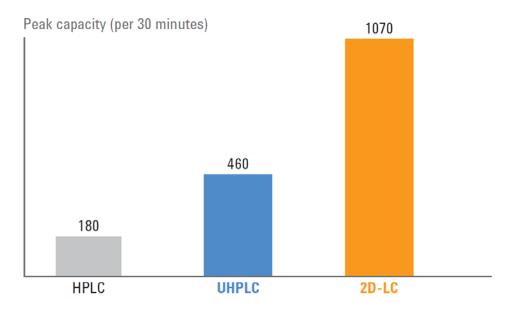
Journal of Chromatography A, 778 (1997) 3-21

Review

#### Some reflections on speed and efficiency of modern chromatographic methods

H. Poppe Amsterdam Institute for Molecular Studies (AIMS), Laboratory for Analytical Chemistry, University of Amsterdam, Nieuwe Achtergracht 166, 1018 WV Amsterdam, Netherlands

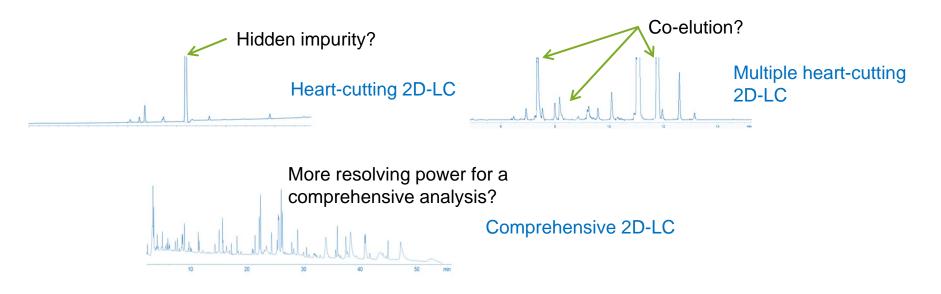
## *"Resolving power is what it is all about in analytical separation science."*





#### Why Two-Dimensional LC?

- Increased peak capacity
- Further resolution of a complex mixture that cannot be separated on a single column
- Sample cleanup by removing matrix or interfering compounds
- Increase sample throughput (two separations going on at once)



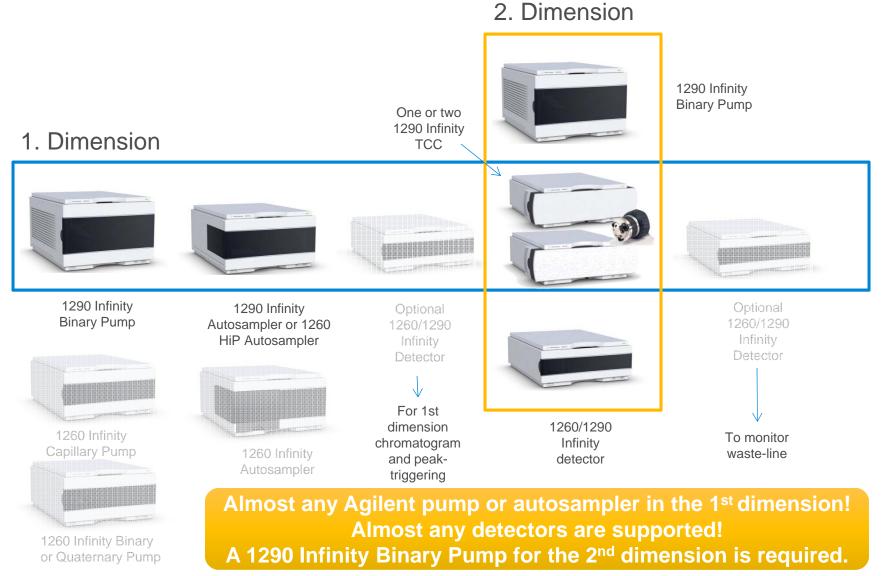


### Agilent's 2D-LC Solution



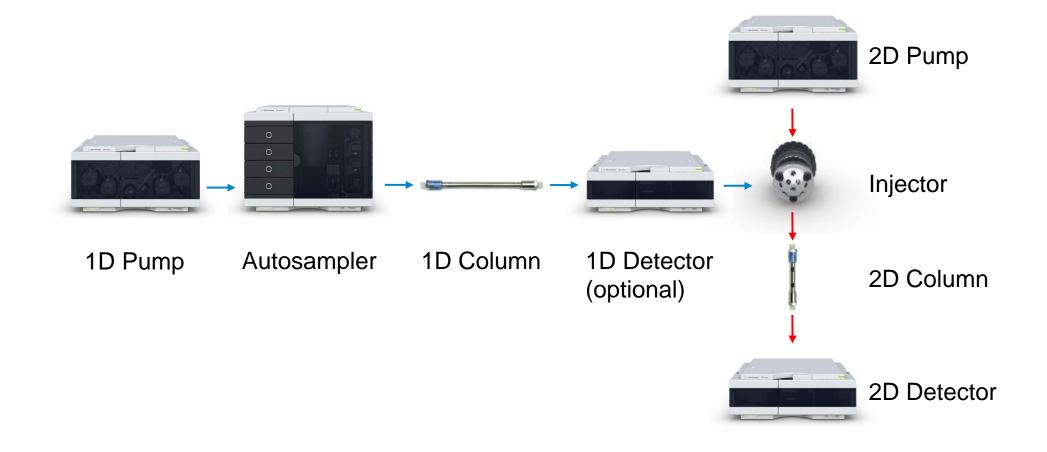


#### **2D LC Modules**



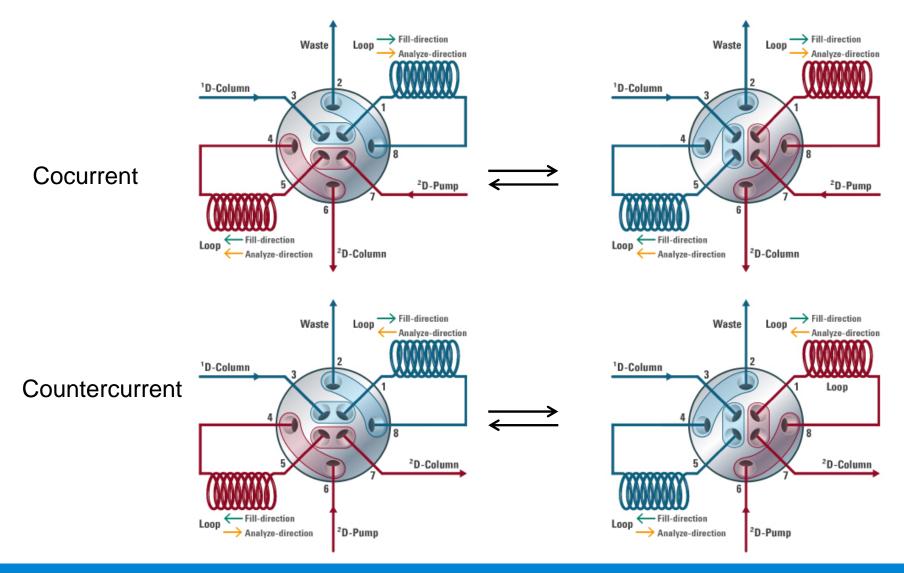


### **2D-LC System Configuration**





### **Unique Agilent 2D-LC Valve**





### Consider Dedicated Bio-inert HPLC Protein Analysis for First Dimension

Unspecific surface binding

- Low throughput, low resolution
- Peak tailing for critical proteins
- Corrosion and pH issues with standard LC
- Decreased column lifetime and chromatographic performance
- Metal-free chromatography needs (e.g. Cr Speciation)
- High pH applications
- Phosphorylated compounds, oligonucleotides



#### Agilent 1260 Infinity II Bio-Inert Quaternary LC The New Standard in Bioanalysis



#### 100% Bio-inert

BIO

inert

- Precious sample does not touch metal surfaces
- ✓ pH range 1-13 (shortterm 14)
- 2 M salt, 8 M urea
- No stainless steel in mobile phase flow path
- New capillary technology

#### **UHPLC** capability

🗸 600 bar

#### **Ease of Use and Robustness**

- Corrosion resistant
- Active seal wash
- Quaternary buffer mixing
- Bio-HPLC columns for biotherapeutic characterization



#### The choice for both, bioanalytical and biopurification up to 10 ml/min



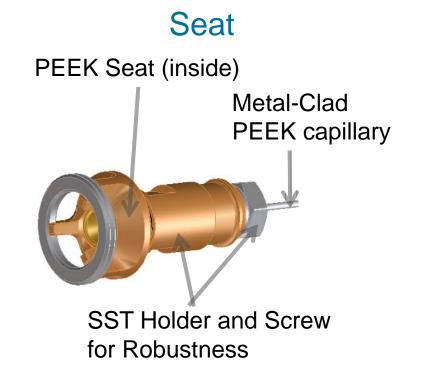
### **Bio-Inert Components**

Inert materials: Titanium (mechanical parts), PEEK, Ceramics, PTFE, FFKM

SST Housing for Robustness

PEEK and
 ceramic for metal free sample path

Needle





### New Capillary Design Ensures Bio-Inertness

Capillaries BIO



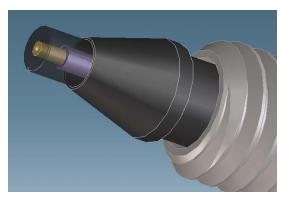
BIO inert

Newest fitting design: Hybrid technique

(Titanium Capillary also available)

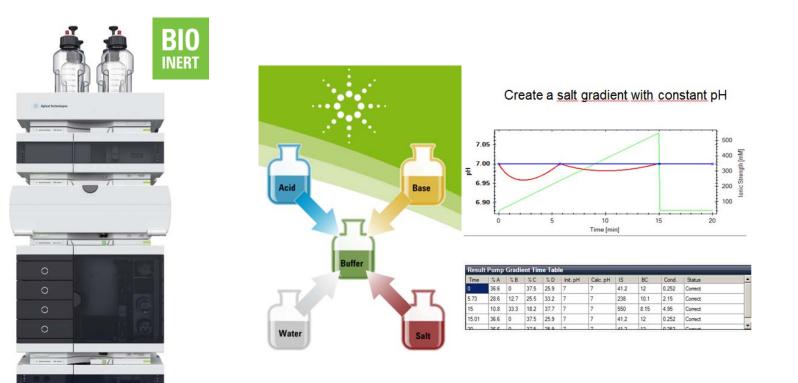
Metal-clad PEEK capillary

New capillary technolgy enables **600 bar** AND is completely metal free !!



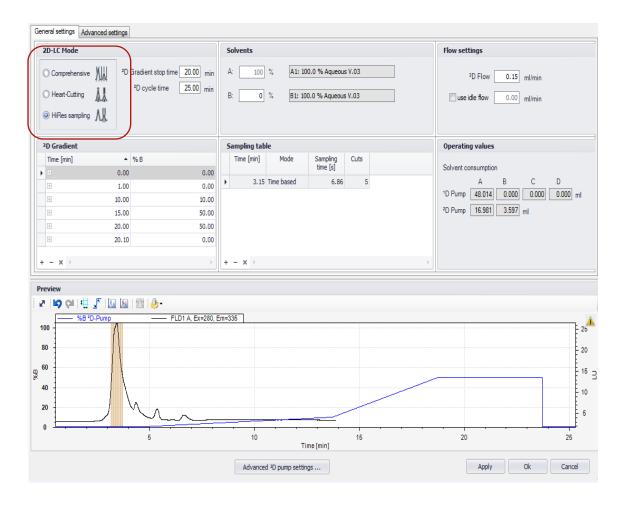


#### **Quaternary Buffer Mixing with Buffer Advisor** Use with Ion Exchange Chromatography





#### 2D-LC Acquisition Software One easy-to-use software for all operation modes



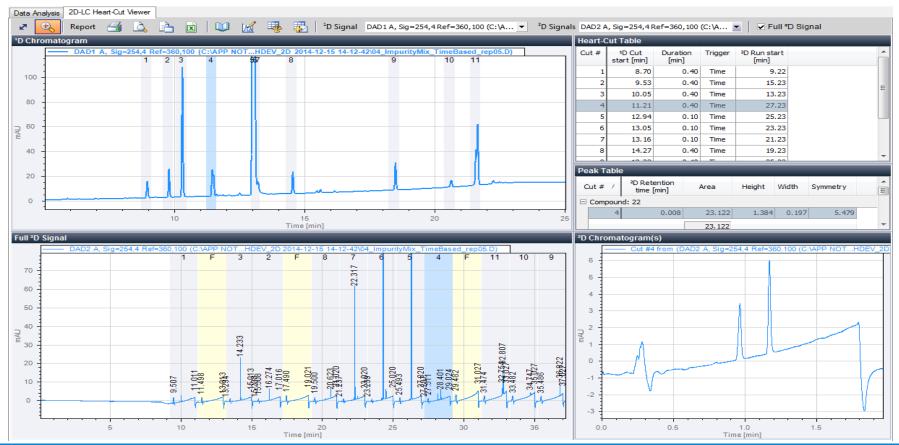
Most intuitive software to set up and edit methods within seconds:

- 1<sup>st</sup> dimension gradient
- 2<sup>nd</sup> dimension gradient
- Gradient shift
- Time-segments
- Method parameter
- Method set-up calculator
- Reference chromatogram overlay for heart-cutting
- Time-based or peak-based mode



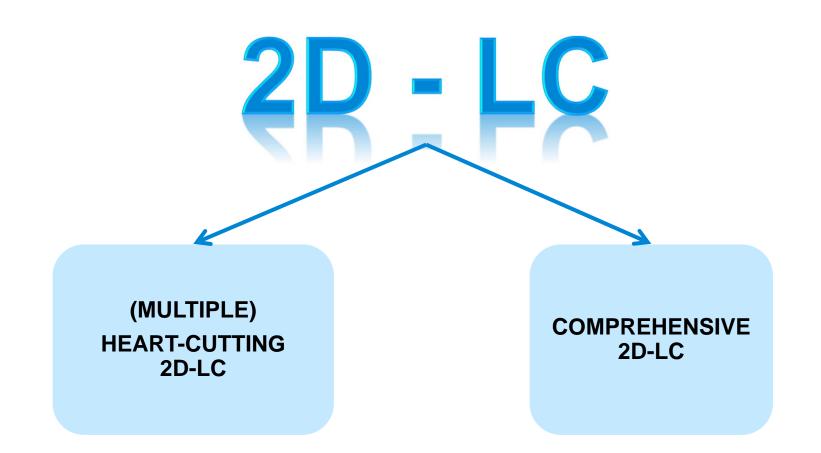
#### Data Analysis (Multiple) Heart-Cutting and High Resolution Sampling

The 2D-LC Heart-Cut Viewer allows straightforward data analysis of (multiple) heart-cutting and High resolution sampling 2D-LC data.





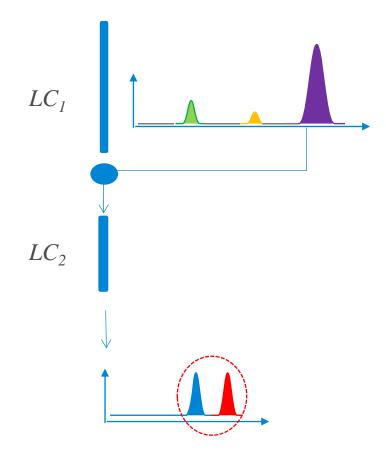
#### **2D-LC Major Types of Operation**





### 2D-LC – Heart-Cutting

#### Heart-cutting 2D-LC (LC-LC):



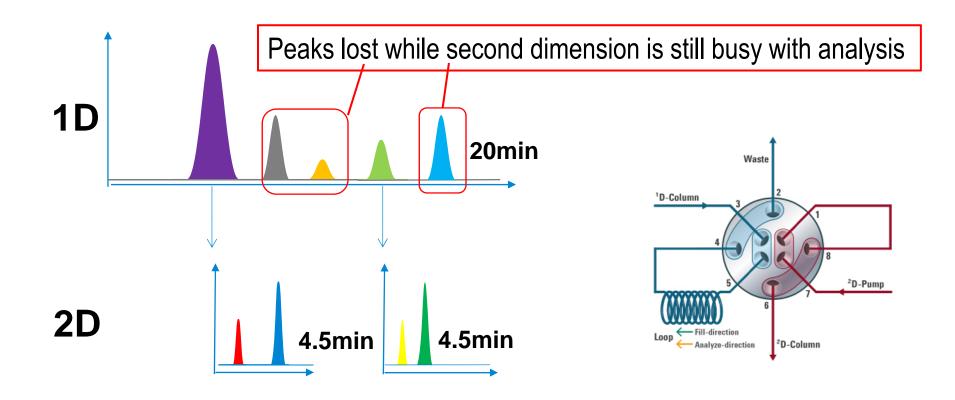
Parts of the 1D effluent are injected onto 2D system

Long 2D gradients possible  $\rightarrow$  good data quality

Limited 2D information

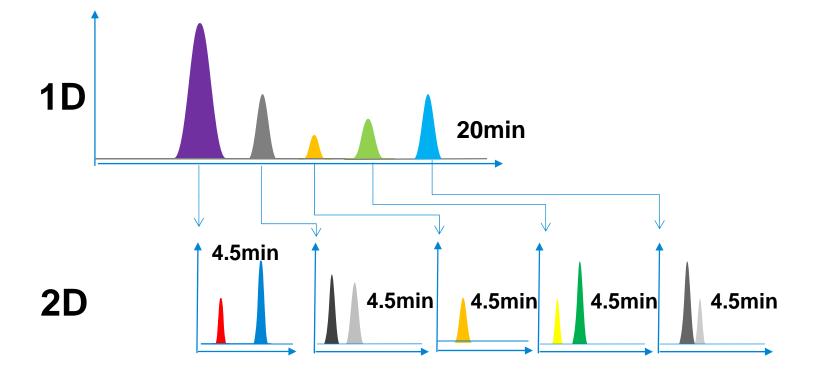


### Limitation of Heart-Cutting 2D-LC





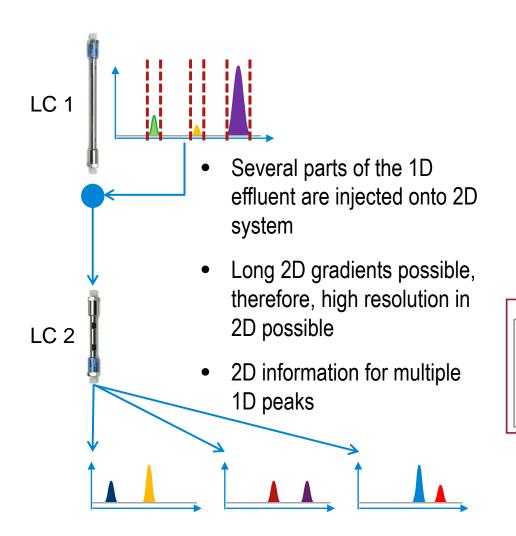
### From Single to Multiple Heart-Cutting 2D-LC



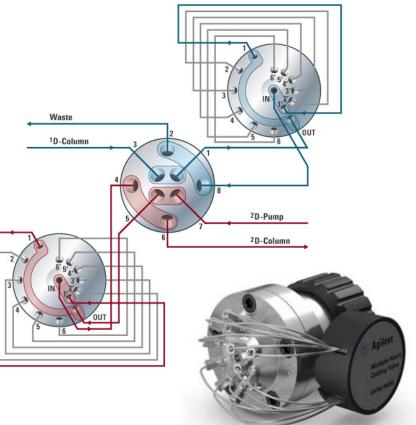
Idea: De-couple the first and second dimensions by intermediately storing peaks eluted from the first dimension for later analysis in the second dimension.



### Multiple Heart-Cutting 2D-LC



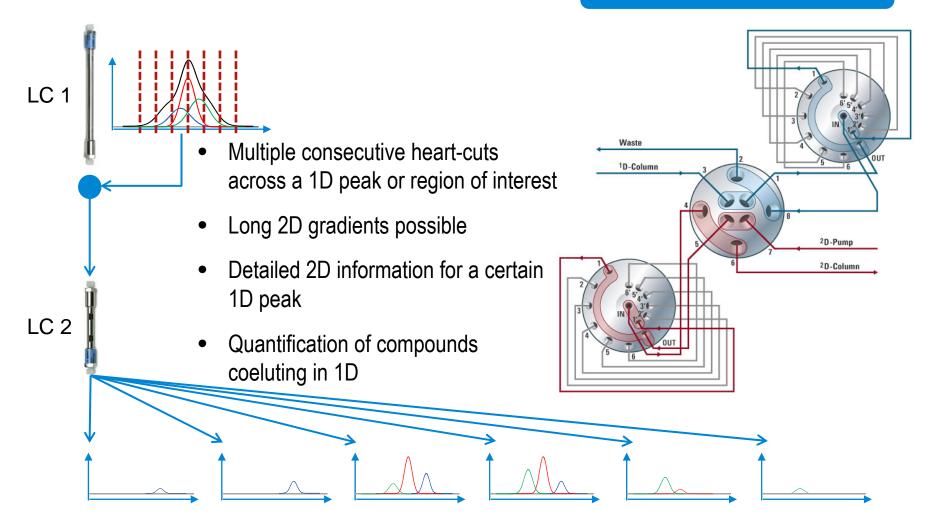
#### Valve and loop configuration



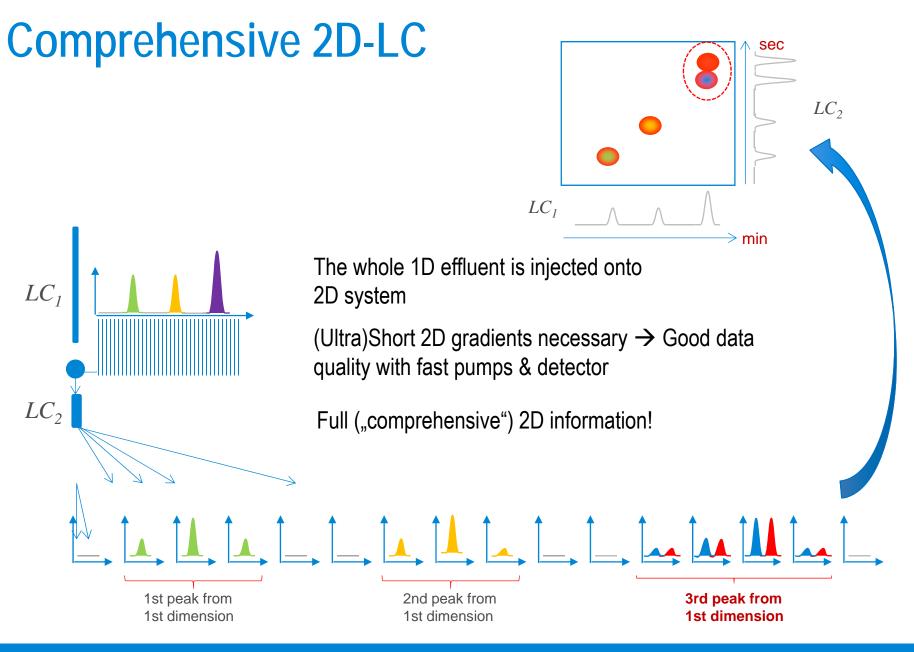


### **High-Resolution Sampling 2D-LC**

#### Valve and loop configuration

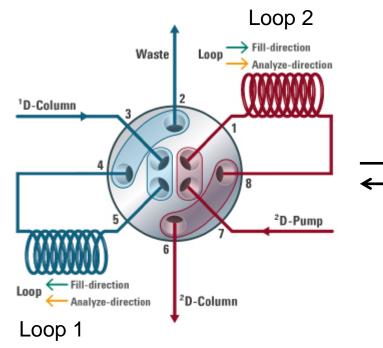




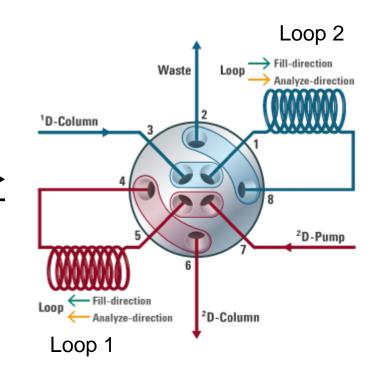




#### Comprehensive 2D-LC How it works



- Collection of effluent from the first dimension column in loop 1
- Analysis of the content from loop 2



- Collection of effluent from the first dimension column in loop 2
- Analysis of the content from loop 1



### Applications of Comprehensive 2D-LC

#### **Finger Printing and Profiling Analysis**

#### Analysis of very complex samples

Traditional Chinese Medicine (TCM)/Chinese Herbal Medicine (CHM):

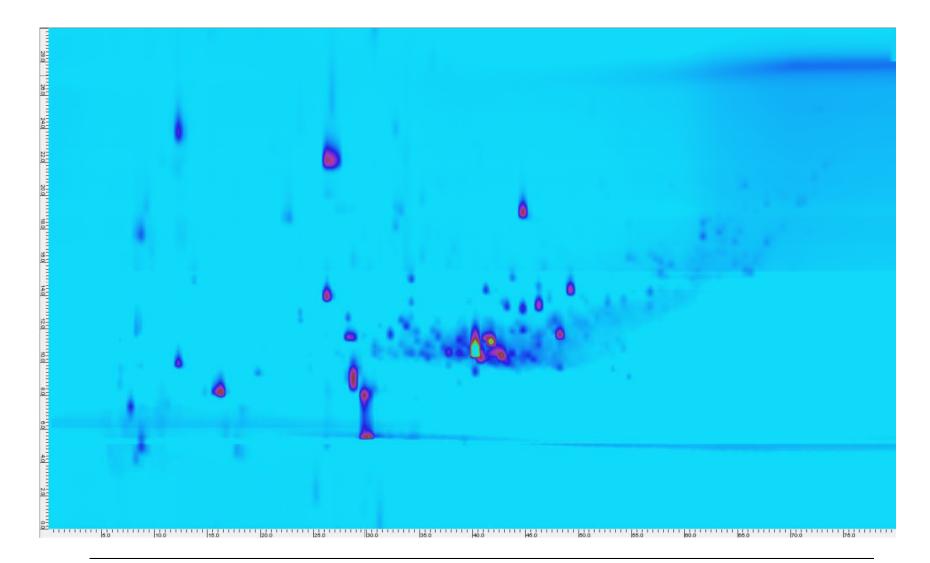
- Holistic healthcare system
- Pharmaceutical efficacy regarded to depend on synergistic effects of multiple components of the plants

#### Biopharmaceutical analysis:

- Peptide maps
- Complex glycan pattern



#### **Comprehensive 2D-LC – Data View**





#### Case Studies Online 2DLC of Monoclonal Antibodies

- Monoclonal Antibody Digests HILIC × RPLC-MS
   → Focus: Peak capacity
- Characterization of Monoclonal Antibodies Protein A x WCX
   → Focus: Combination of two different workflows
- Characterization of Monoclonal Antibodies SEC x WCX
   → Focus: Combination of two different workflows
- Characterization of Charge Variants WCX x RP
   → Focus: Desalting after IEX prior to MS



### Case Study 1 Comprehensive 2D-LC Orthogonal Separations of mAb Tryptic Digest

- Aggregation Studies
- Charge Variant Analysis
- Peptide Mapping
- Glycan Profiling
- Titer Analysis
- And others...



#### Analysis of Monoclonal Antibody Digests with the Agilent 1290 Infinity 2D-LC Solution



#### Focus: High Peak Capacity

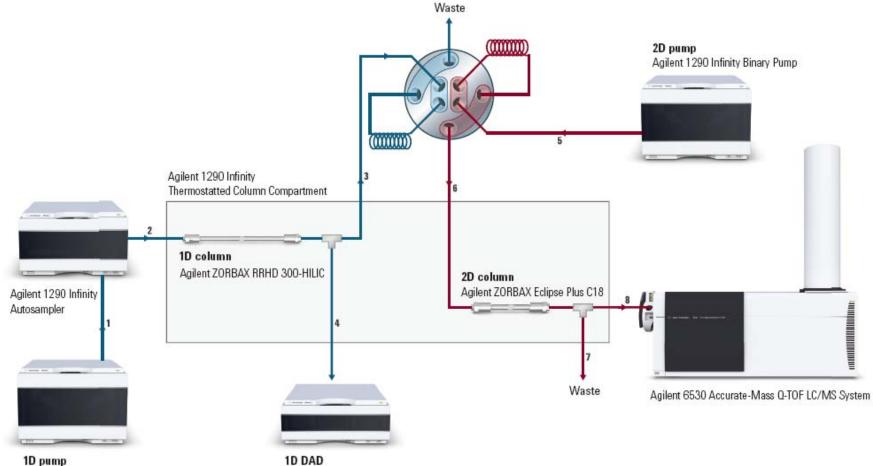


# Structure and Amino Acid Sequence of Trastuzumab

T1       T2       T3       T4         DIQMTQSPSSLSASVGDR       VTITCR       ASQDVNTAVAWYQQKPGK       APK         T5       T6       T7         LLIYSASFLYSGVPSR       FSGSR       SGTDFTLTISSLQPEDFATYYCQQHYTTPPTFGQGTK         T8       T9       T10       T11       T12       T13         VEIK       R       TVAAPSVFIFPPSDEQLK       SGTASVVCLLNNFYPR       EAK       VQWK         T14       T15       T16       T17         VDNALQSGNSQESVTEQDSK       DSTYSSTLTLSK       ADYEK       HK         T18       T19       T20         VYACEVTHQGLSSPVTK       SFNR       GEC	EVOLVESGGGLVOPGGSLRLSCAASGFN IKDTYIHWVROAPGKGLEW N-ter Hc Lc
T21 EVOLVESGGGLVOPGGSLR LSCAASGFNIK DTYIHWVR OAPGK GLEWVAR T26 T27 T28 T29 T30 T31 IYPTNGYTR YADSVK GR FTISADTSK NTAYLOMNSLR AEDTAVYYCSR T32 WGGDGFYAMDYWGOGTLVTVSSASTK GPSVFPLAPSSK STSGGTAALGCLVK T35 DYFPEPVTVSWNSGALTSGVHTFPAVLOSSGLYSLSSVVTVPSSSLGTOTYICNVNHKPSNTK +	C-ter C-ter OGGNVFSCSVMHEALHNHYTOKSLSLSPG
DYFPEPVTVSWNSGALTSGVHTFPAVLOSSGLYSLSSVVTVPSSSLGTOTYICNVNHKPSNTK T36 T37 T38 T39 T40 T41 VDK K VEPK SCDK THTCPPCPAPELLGGPSVFLFPPKPK DTLMISR T42 T43 T44 T45 TPEVTCVVVDVSHEDPEVK FNWYVDGVEVHNAK TKPR EEQYNSTYR T46 T47 T48 T49 T50 T51 T52 T53 T54 VVSVLTVLHODWLNGK EYK CK VSNK ALPAPIEK TISK AK GOPR EPOVYTLPPSR T55 T56 T57 T58 T59 EEMTK NOVSLTCLVK GFYPSDIAVEWESNGOPENNYK TTPPVLDSDGSFFLYSK LTVDK	<ul> <li>62 identity peptides</li> <li>Modifications</li> <li>Incomplete and aspecific cleavages</li> <li></li> <li>&gt; 100 peptides</li> </ul>
T60 T61 T62 SR WQQGNVFSCSVMHEALHNHYTQK SLSLSPG	



#### **Instrument Setup**

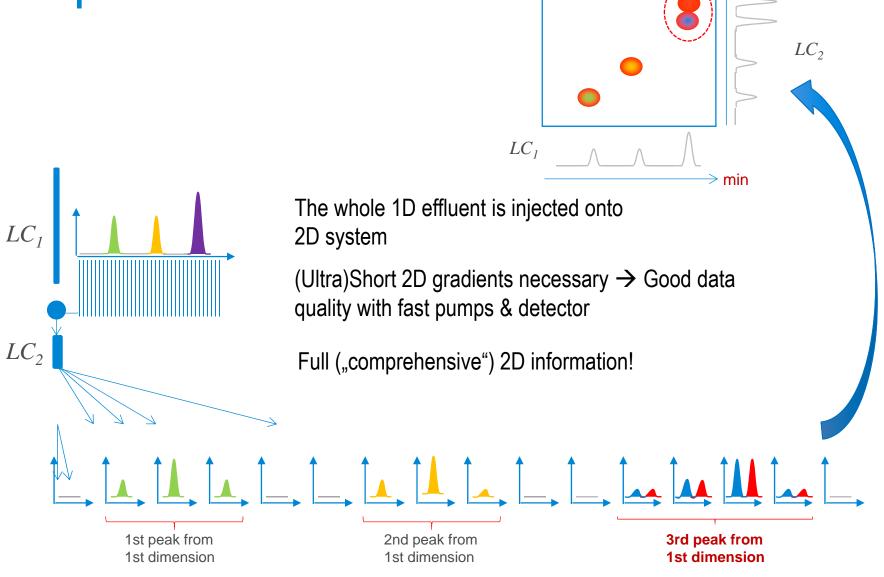


Agilent 1290 Infinity Binary Pump

1D DAD Agilent 1290 Infinity Diode Array Detector



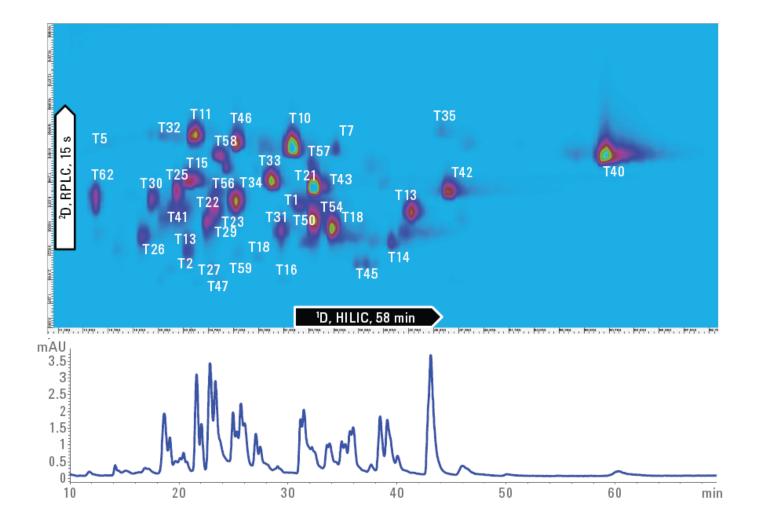
### **Comprehensive 2D-LC**





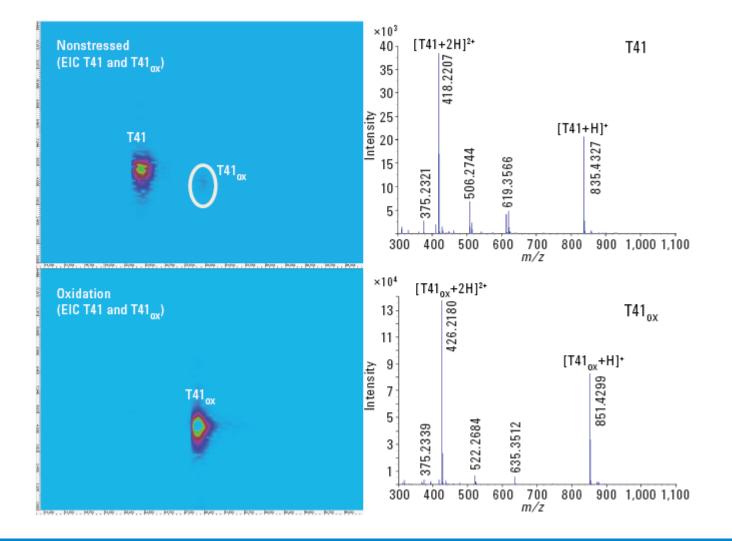
sec

#### **2D-LC Contour Plot**



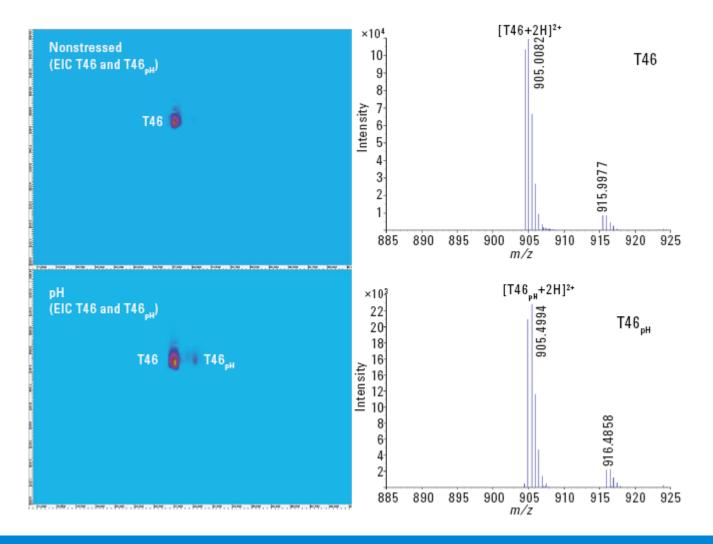


#### **2D-LC Trastuzumab Oxidation**





#### **2D-LC Trastuzumab Degradation**





## Comprehensive 2D-LC mAb Tryptic Digest Summary

- Higher peak capacity with 2D-LC
- Improved separation using othogonal column combination: HILIC x RP
- Proof of concept with the clear differentiation of stressed and non-stressed antibody samples
- Ideal combination of Agilent 1290 Infinity 2D-LC Solution coupled to an Agilent 6530 Accurate-Mass Q-TOF LC/MS

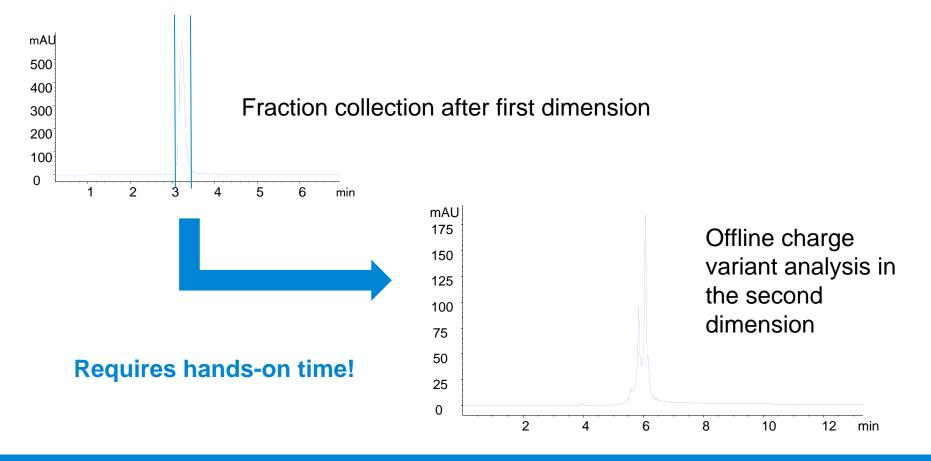


### Case Study 2 Combining Two Techniques Protein A and WCX

- Aggregation Studies
- Charge Variant Analysis
- Peptide Mapping
- Glycan Profiling
- Titer Analysis
- And others...

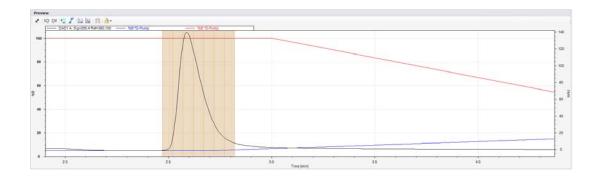


#### Combination of Two Analyses Offline Approach

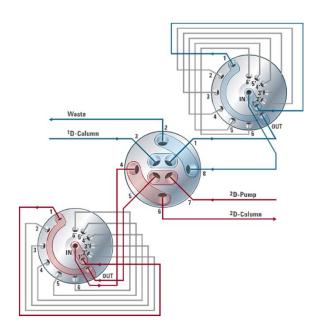




#### Characterization of Monoclonal Antibodies Protein A and Weak Cation Exchange Chromatography



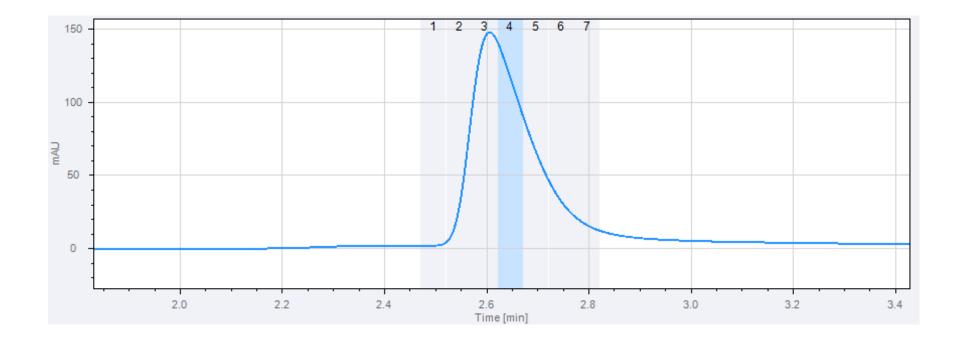
High Resolution Multiple Heart Cuts



#### Focus: Combination of two important Quality Attribute Analyses



#### 2D-LC with High Resolution Sampling Protein A Titer Analysis (First Dimension)

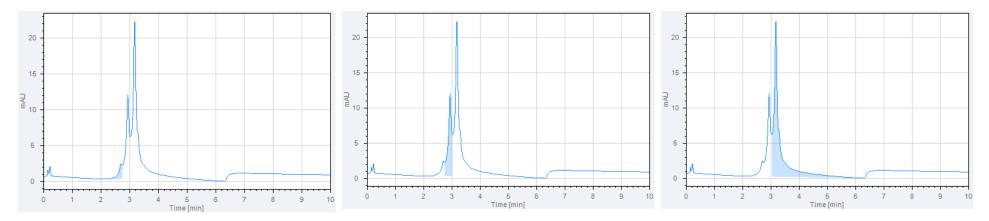


Consecutive Cuts Across the Peak



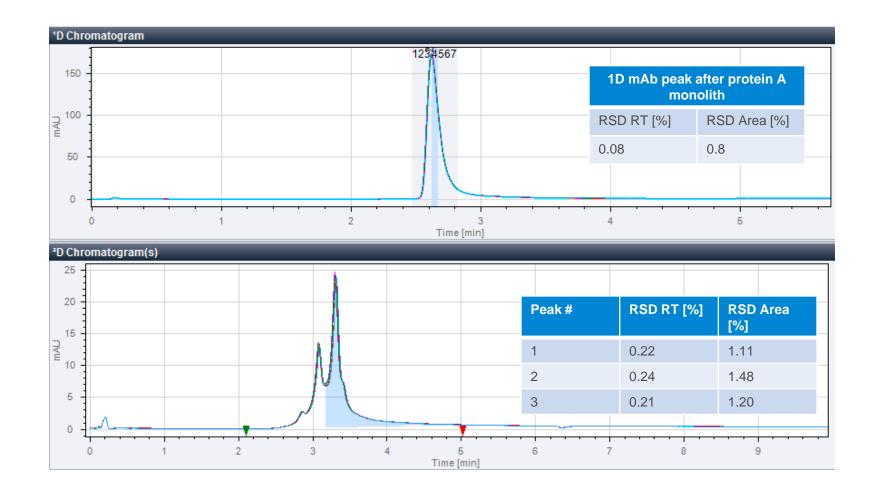
#### 2D-LC with High Resolution Sampling Weak Cation Exchange Chromatography (Second Dimension)

Peak table (²D)						
Cut # 🔺	<sup>2</sup> D Retention time [min]	Area	Height	Width	Symmetry	
Compound: 3						
4	2.946	78.259	9.293	0.115	1.494	
		78.259				
Compound: 4						
4	3.184	250.323	17.263	0.188	0.424	
		250.323				
Compound: 5						
4	2.706	16.746	1.827	0.123	2.747	
		16.746				



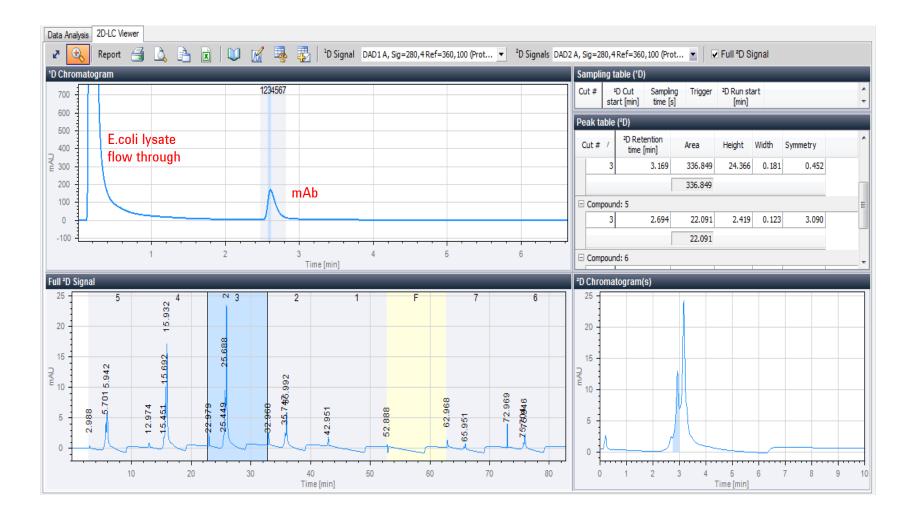


#### **Precision of Retention Time and Area**





#### Mimicking a Cell Lysate mAb Process Control Sample



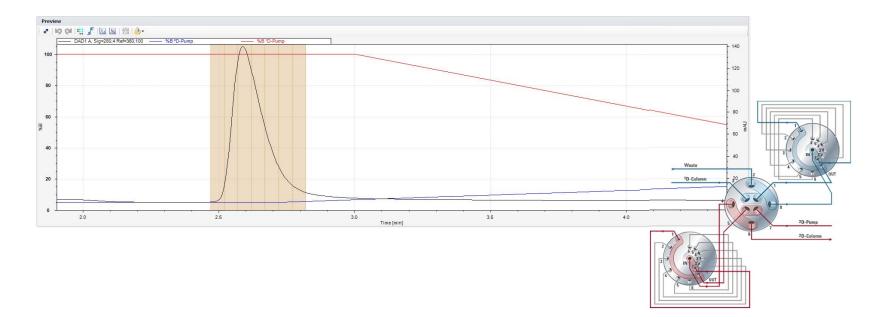


## High Resolution Multiple Heart Cuts Protein A - WCX Summary

- Fully automated combination of two different quality attribute analyses
   Protein A titer analysis and charge variants
- No hands-on time required between the two analyses
- High precision of retention time and area found in both dimensions
- Also suitable for process control samples



#### Characterization of Monoclonal Antibodies Size Exclusion and Weak Cation Exchange Chromatography



#### Focus: Combination of two important Quality Attribute Analyses

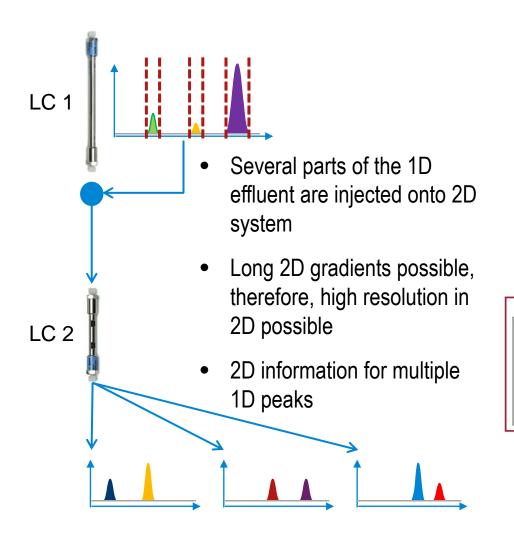


# Case Study 3 Combining Two Techniques SEC and WCX

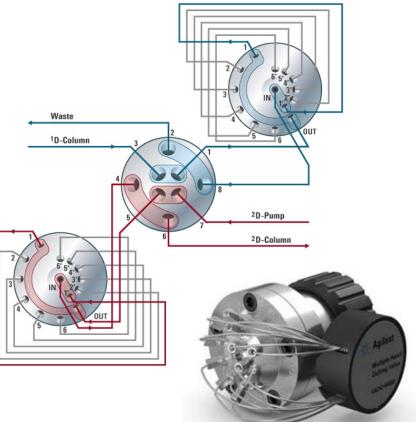
- Aggregation Studies
- Charge Variant Analysis
- Peptide Mapping
- Glycan Profiling
- Titer Analysis
- And others...



### Multiple Heart-Cutting 2D-LC

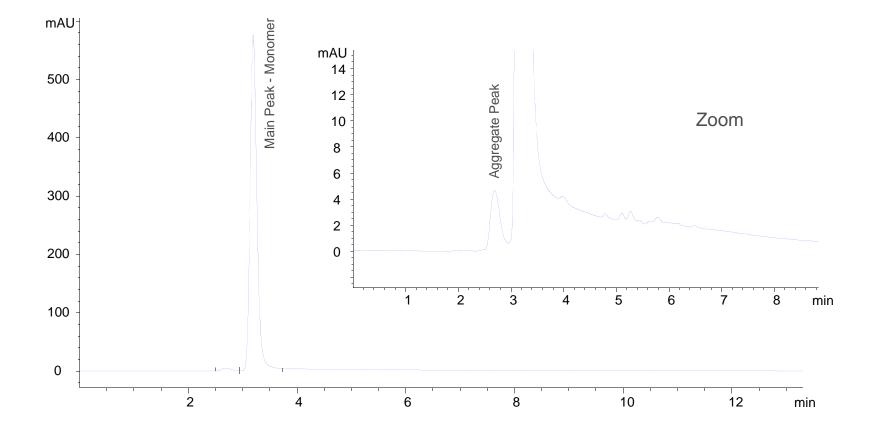


#### Valve and loop configuration



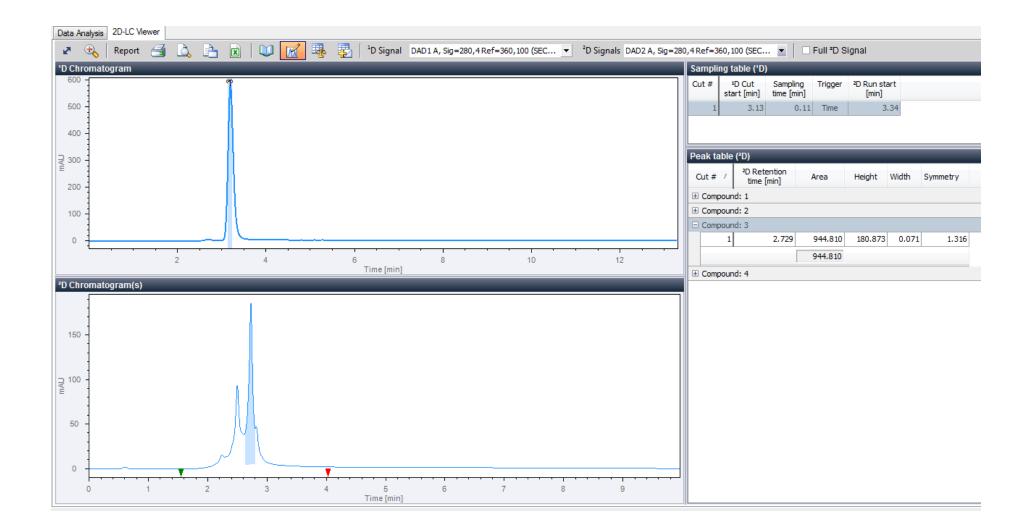


### Aggregation Analysis of mAb SEC in the First Dimension



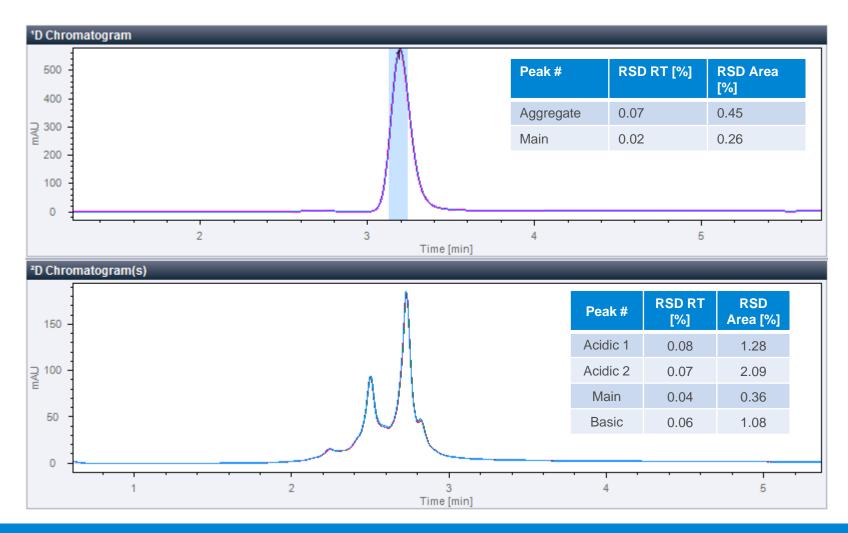


#### Heart Cuts from SEC Sent to WCX





#### 2D LC Multiple Heart Cut SEC – WCX Precision of Retention Time and Area



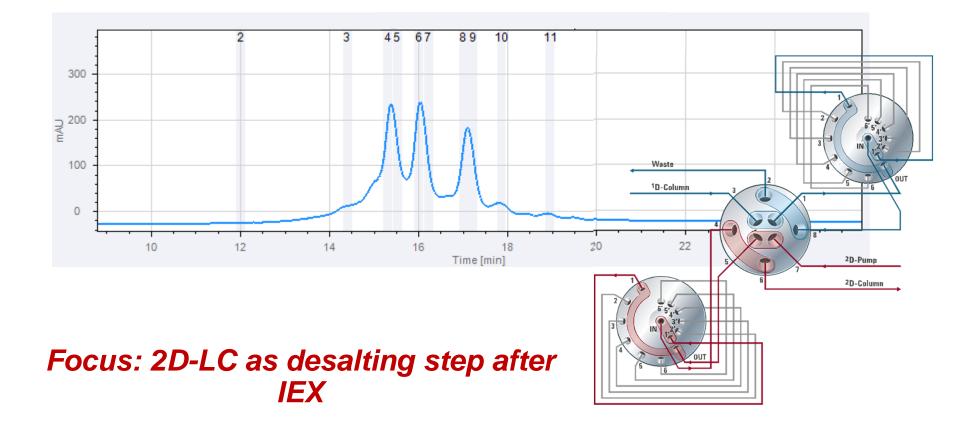


# Case Study 3 Combining SEC and WCX Summary

- Fully automated combination of two different quality attribute analyses
   Size Exclusion and Charge Variants
- No hands-on time required between the two analyses
- High precision of retention time and area found in both dimensions

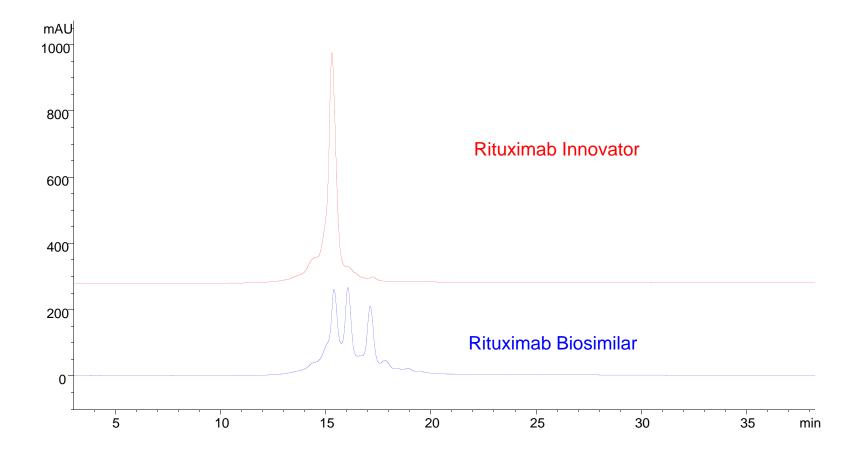


#### Case Study 4 Charge Variants Analysis LCMS Ion Exchange and Reversed-Phase Chromatography





#### Charge profiles of Rituximab Innovator and Biosimilar



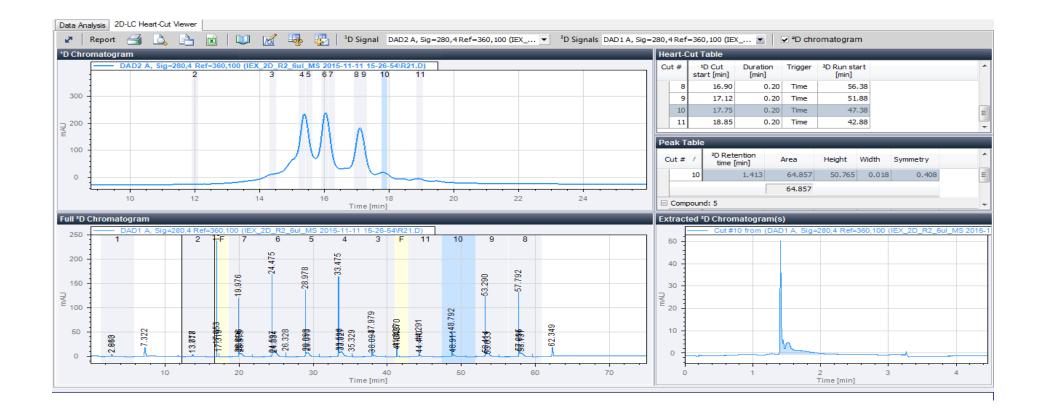


#### WCX (first dimension) and Reverse Phase (second dimension)

etup <sup>2</sup> D-Pump: (G4220A DE92900288)							
General settings Advanced settings							
2D-LC Mode	Solvents	Flow settings					
○ Comprehensive       2D Gradient stop time       3.00 min         ◎ Heart-Cutting       ▲       2D cycle time       4.50 min         ○ HiRes sampling       ▲       ▲       ▲	A:         90         %         A1: 100.0 % Water V.03           B:         10         %         B1: 100.0 % Acetonitrile V.03	<sup>2</sup> D Flow 1.00 ml/min v use idle flow 0.10 ml/min					
2D Gradient	<sup>2</sup> D Time segments	Operating values					
Time [min]         ▲         % B           ▶         0.00         10.00	Time [min]         Mode         Sampling time [min]         Loop filling [%]         Prioritize         2D + in	Solvent consumption					
2.50 60.00	▶ 12.00 Time based 0.20 100	A B					
2.75 90.00	14.80 Time based 0.20 100	<sup>1</sup> D Pump 13.900 1.100 ml					
	15.35 Time based 0.20 100	<sup>2</sup> D Pump 48.363 17.925 ml					
	15.78 Time based 0.20 100						
	16.00 Time based 0.20 100						
	16.48 Time based 0.20 100						
	16.70 Time based 0.20 100 -						
+ - x (	+ - x (						
Preview		- %B*D]					
100 80 60 0 0 0 10 20 30 Time [0	40 50 60 70	D 1 2 3 4 Time [min]					
•	Advanced <sup>2</sup> D pump settings	Apply Ok Cancel					

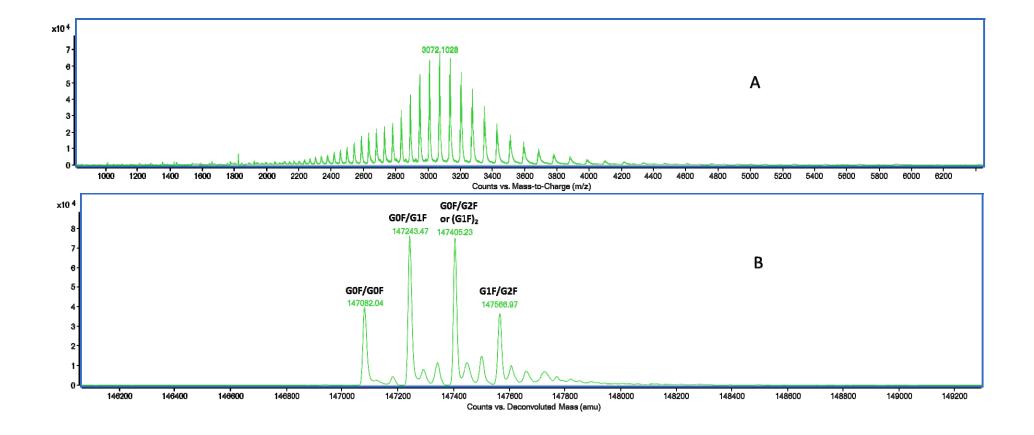


#### **2D LC Data View of UV Results**





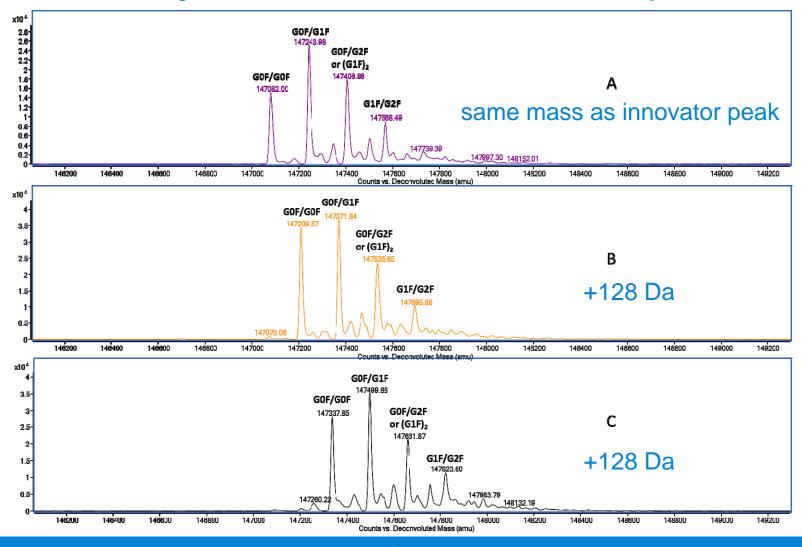
#### Online MS detection Intact mass analysis of Innovator Rituximab – 1 peak





#### **Online MS detection**

Intact Mass Analysis of Biosimilar Rituximab – 3 peaks





### Case Study 4 Charge Variants Analysis LCMS Summary

- Innovator and Biosimilar Rituximab charge variants were analyzed using weak cation exchange chromatography
- Online 2DLC-MS qualification was enabled by automated desalting and denaturation
- The three peaks of the biosimilar vs one peak of the innovator were qualified as c-terminal lysine variants (mass shift of 128 Da)

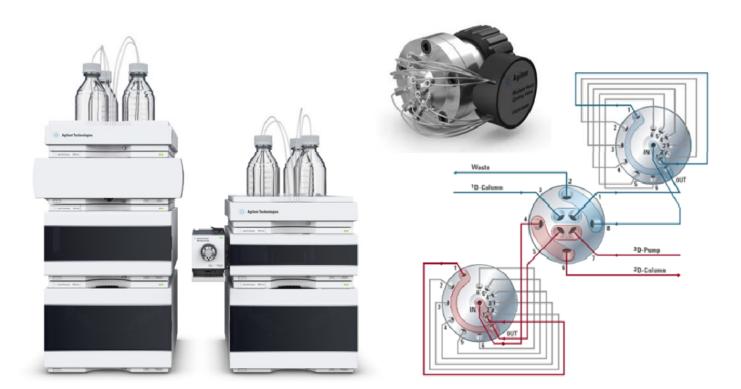


#### Further information can be found...

- Schneider, S. 2D-LC/MS Characterization of Charge Variants Using Ion Exchange and Reversed-Phase Chromatography, Agilent Technologies Application Note, publication number 5991-6673EN, 2016.
- Schneider, S. Online 2D-LC Characterization of Monoclonal Antibodies Using Protein A and Weak Cation Exchange Chromatography, Agilent Technologies Application Note, publication number 5991-6848EN, 2016.
- Schneider, S. Online 2D-LC Characterization of Monoclonal Antibodies with Size Exclusion and Weak Cation Exchange Chromatography, Agilent Technologies Application Note, publication number 5991-6906EN, **2016.**
- Vanhoenacker, G et al.; Analysis of Monoclonal Antibody Digests with the Agilent 1290 Infinity 2D-LC Solution Part 2: HILIC × RPLC-MS, Agilent Technologies Application Note, publication number 5991-4530EN, 2014.
- Schneider, S; Naegele, E; Krieger, S. Online 2D-LC Analysis of Complex N-Glycans in Biopharmaceuticals Using the Agilent 1290 Infi nity 2D-LC Solution Agilent Technologies Application Note, publication number 5991-5349EN, **2015**.



## Online 2D-LC Analysis of Complex N-Glycans in Biopharmaceuticals





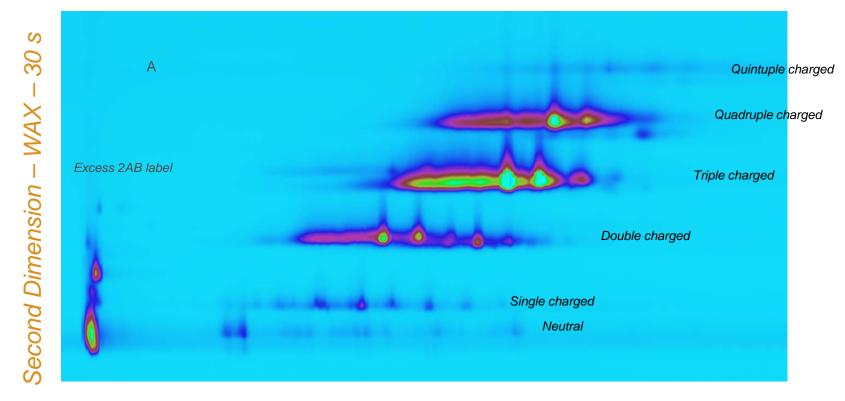
### **Comprehensive 2D-LC of Glycans**

#### HILIC / WAX

- First Dimension:
  - Agilent AdvanceBio Glycan Mapping column, 2.1 × 150 mm, 1.8 μm Total run time 165 min (Stoptime + Posttime)
- Second Dimension:
  - Agilent Bio WAX column, 2.1 × 50 mm, 5 µm 30 seconds 2D gradients



## Comprehensive 2D-LC of Fetuin HILIC/WAX



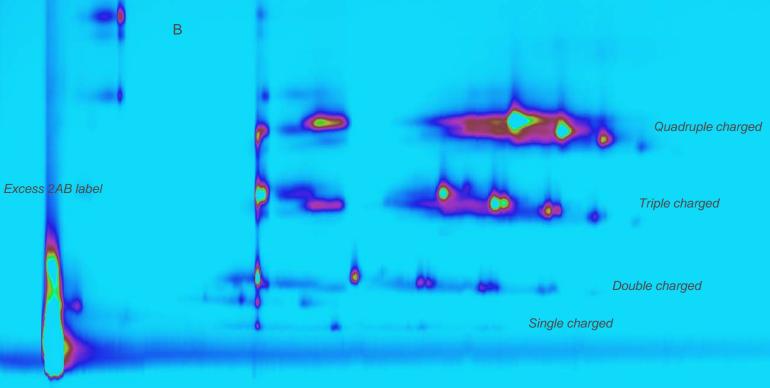
#### First Dimension – HILIC – 110 minutes

http://web.expasy.org/glycomod/ for the determination of glycan structures



#### **Comprehensive 2D-LC of EPO** HILIC/WAX

S 30 Second Dimension – WAX –



First Dimension – HILIC – 110 minutes



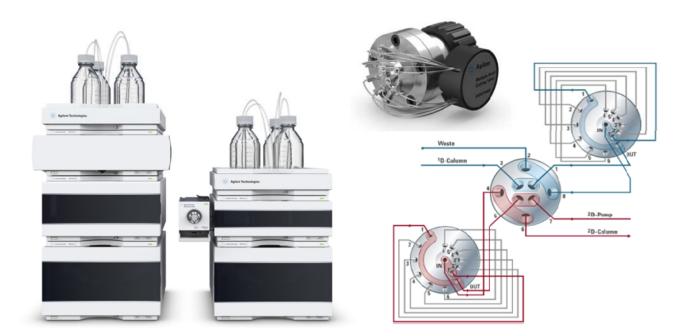
**Agilent Technologies** 

### Comprehensive 2D-LC Analysis of Glycans Summary

- Comprehensive HILIC/WAX high peak capacity
- Complete automation possible within a run time of about 110 minutes
- Easy data analysis and interpretation due to the grouping of the glycans according to their charge in the second dimension, enabling simultaneous charge profiling



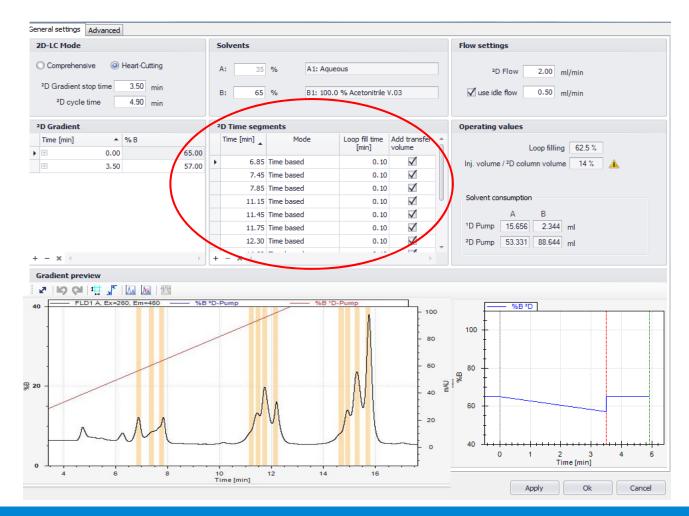
# Multiple Heart-Cutting 2D-LC of Glycans





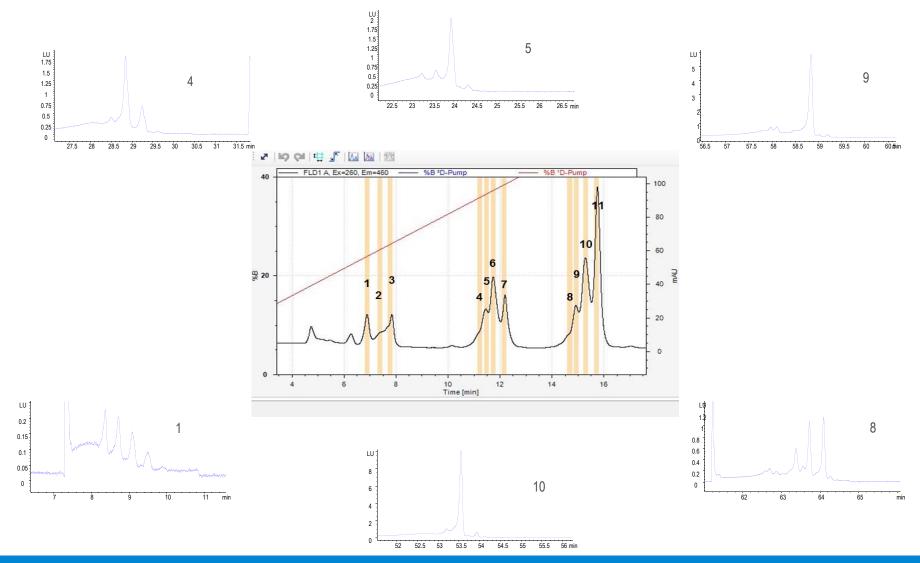
**Agilent Technologies** 

#### Time-Based Multiple Heart-Cutting of EPO 2D-LC Method Setup





#### **Time-Based Multiple Heart-Cutting of EPO**





### Time-Based Multiple Heart-Cutting of EPO Summary

- Multiple heart-cutting WAX/HILIC High Resolution
- Higher flexibility no limitation to super short gradients and high flow rates in the second dimension
- Facilitates the combination of WAX/HILIC having WAX in the first and HILIC in the second dimension enabling partly fill of the sample loops
- Time saving of over 70 % compared to the offline WAX/HILIC analysis with 4 hours vs only 70 minutes



## Thank you for your attention

### **Questions?**





**Agilent Technologies**