

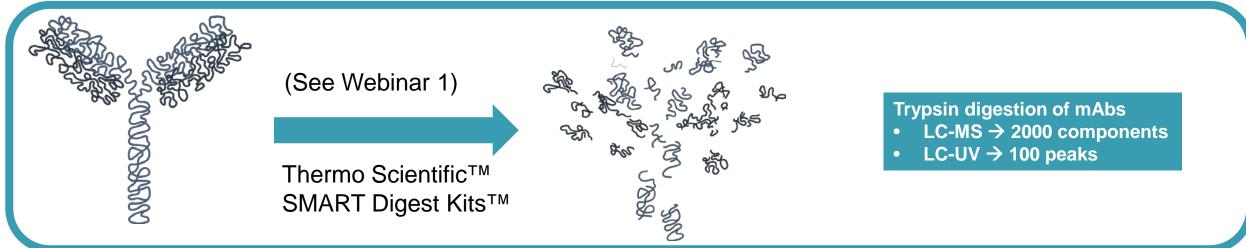
#### **ThermoFisher** SCIENTIFIC

# Peptide separations with pinpoint precision

Mauro De Pra, PhD

The world leader in serving science

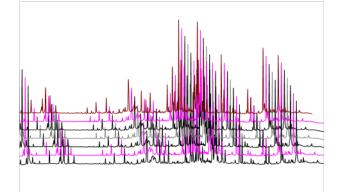
#### Identity confirmation or purity analysis



#### **Method requirements**

High resolving power in reasonable time

- 30-45 minutes run time
- Theoretically capable of resolving > 300 peaks

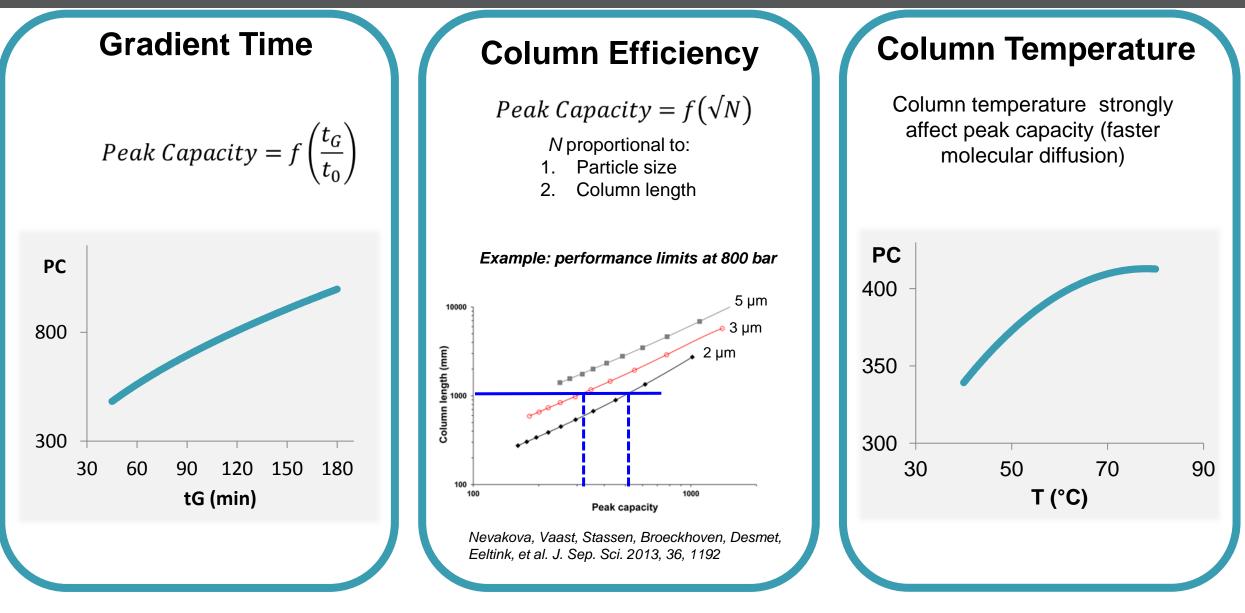


Comparison against reference

Highly reproducible method



# How do we achieve high resolving power?





# High resolving power without compromising on time

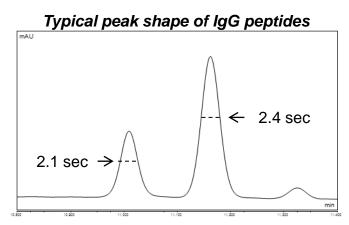
Thermo Scientific<sup>™</sup> Acclaim<sup>™</sup> VANQUISH<sup>™</sup> C18

- 2.2 µm particles
- 120 Å
- 250 mm length
- 2.1 mm ID
- 60 °C
- Excellent peak shape with FA and TFA

Thermo Scientific™ Vanquish™ Flex UHPLC Quaternary System

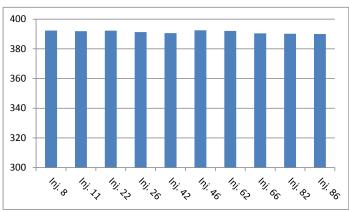


- 1000 bar (15,000 psi)
- Quaternary pump
- Advanced temperature control
- Low dispersion fluidics
- SmartInject
- Seamless integration
  with HRAM MS



2x Acclaim VANQUISH C18 25 cm (50 cm total) 30 minutes gradient @ 400 µL/min 0.05% TFA

#### Peak capacity for BSA digest

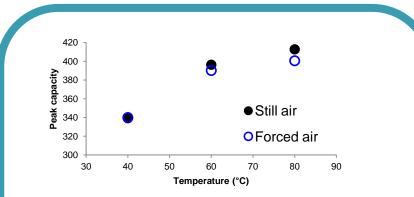


Acclaim VANQUISH long lasting performances

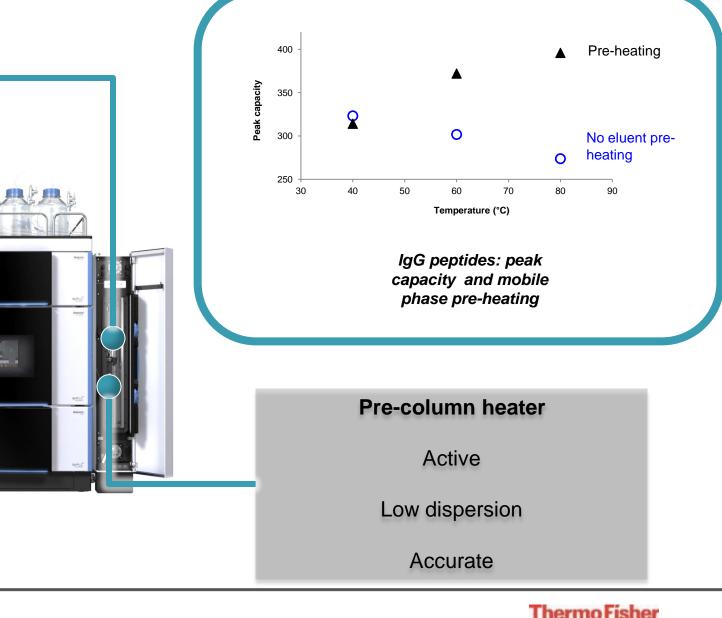


# Column thermostatting



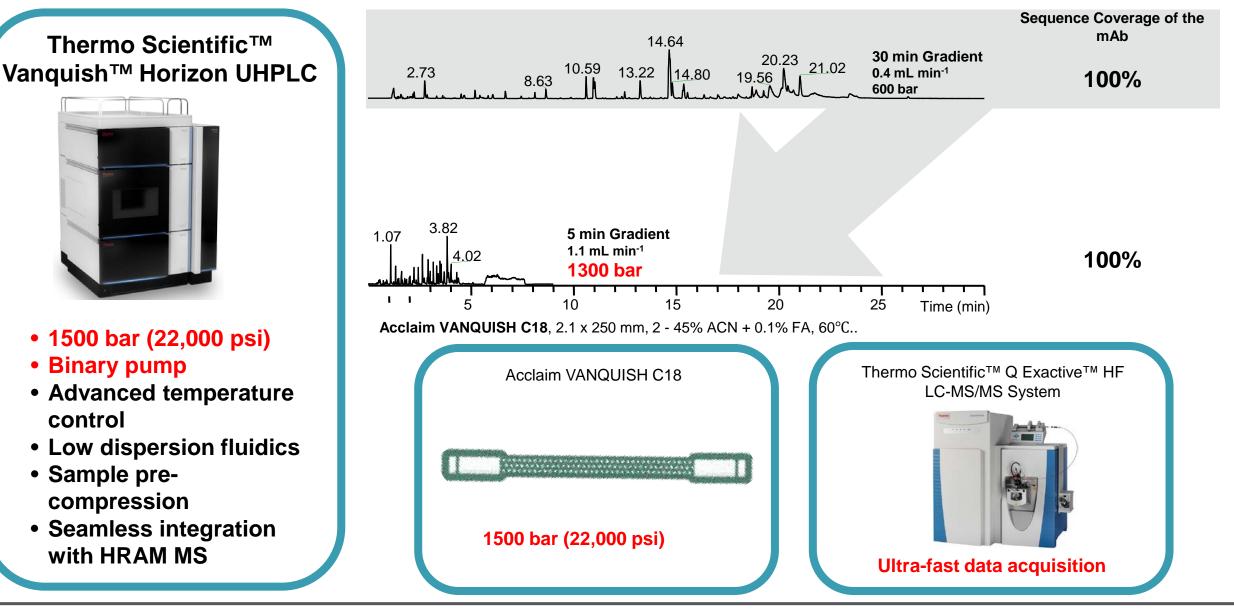


- Slight capacity increase with still-air
- Dual heating mode facilitates method transfer from wide range of instruments
- Forced-air heating beneficial for fast system preparation



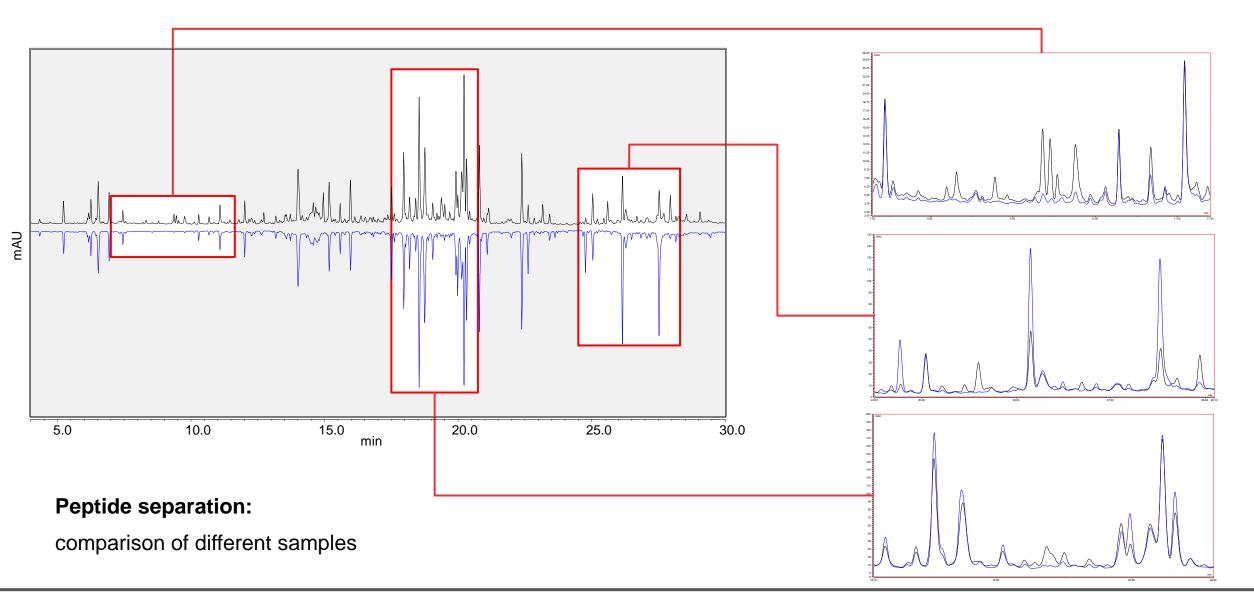
SCIENTIFIC

# Vanquish Horizon UHPLC System – Ultra High Throughput



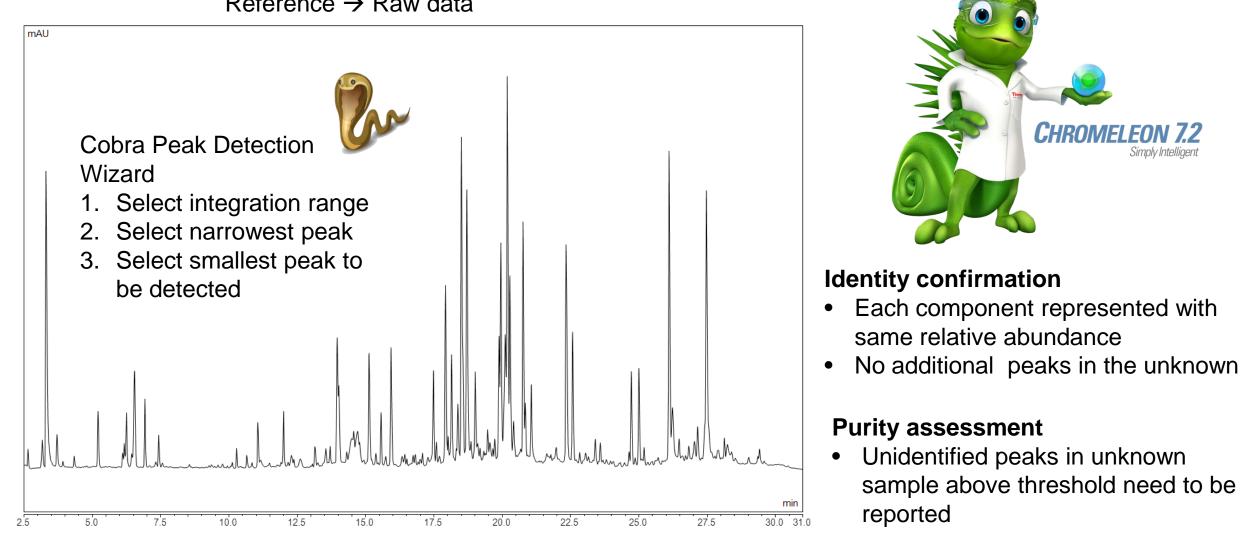


### Visual comparison reference & unknown



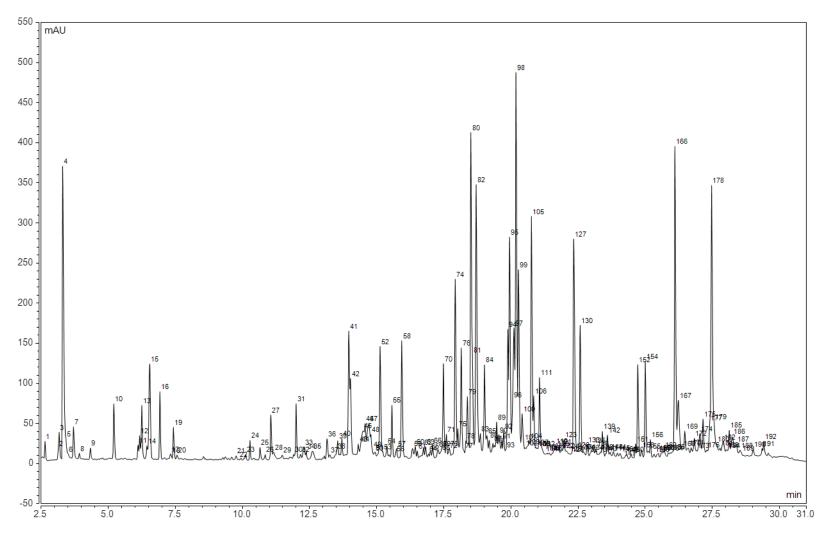


Reference  $\rightarrow$  Raw data





Reference  $\rightarrow$  Integrated





#### **Identity confirmation**

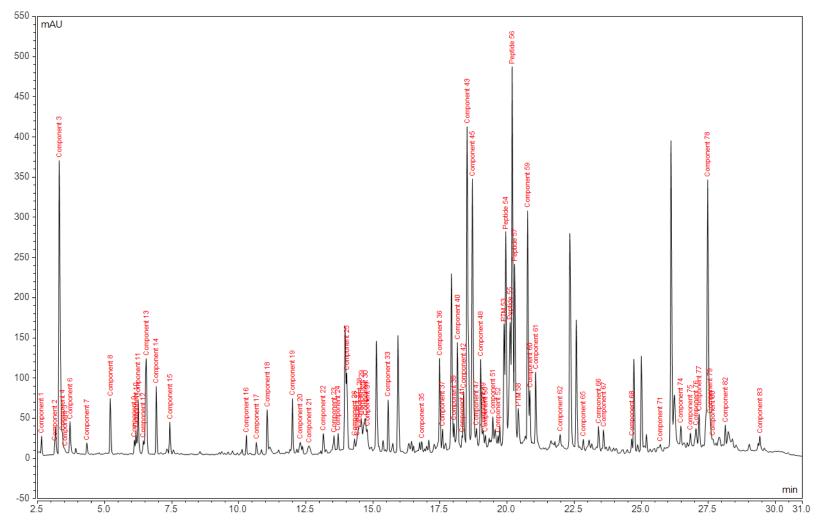
- Each component represented with same relative abundance
- No additional peaks in the unknown

#### **Purity assessment**

 Unidentified peaks in unknown sample above threshold need to be reported



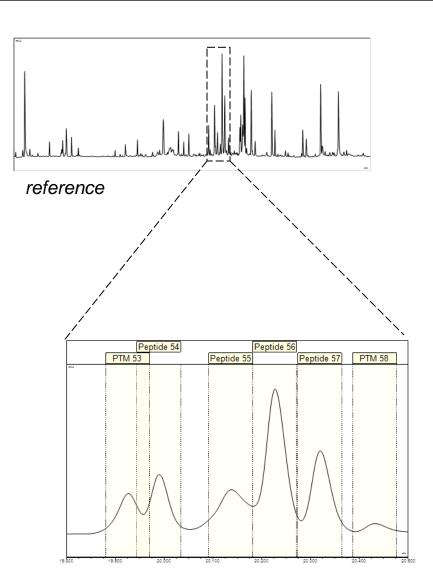
Reference  $\rightarrow$  Peak assigned

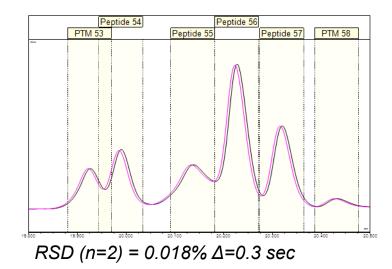




- Generic assignment or identity based on MS data
- Peak assignment saved as Processing Method
- Check if all assigned peaks are present in the unknown sample
- Check if non-assigned peaks are present in the unknown sample







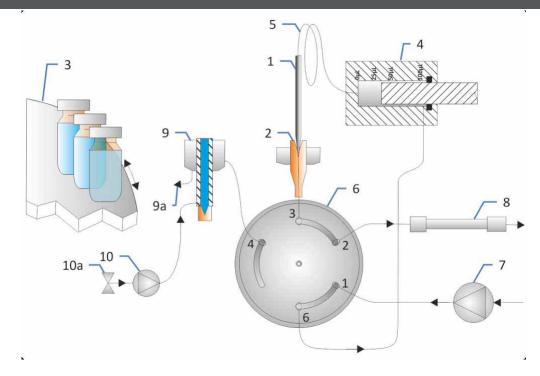
	Peptide 54		Peptide 56		
PTM 5	3	Peptide 55		Peptide 57	PTM 58
19,900	20.000	20.100	20.200	20.300	20.400 2
RSD (n:	=2) =	0.160%	∆= -3	.8sec	

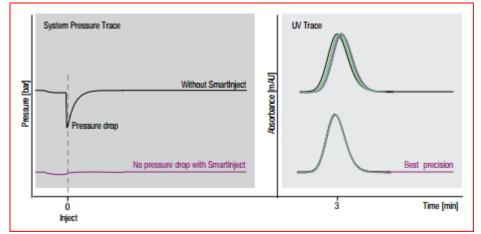
Component	Reference Rel. Area (%)	Unknown Rel. Area (%)
PTM 53	1.9	1.8
Peptide 54	2.6	2.4
Peptide 55	3.1	2.9
Peptide 56	6.7	6.3
Peptide 57	3.8	3.5
PTM 58	0.7	0.6

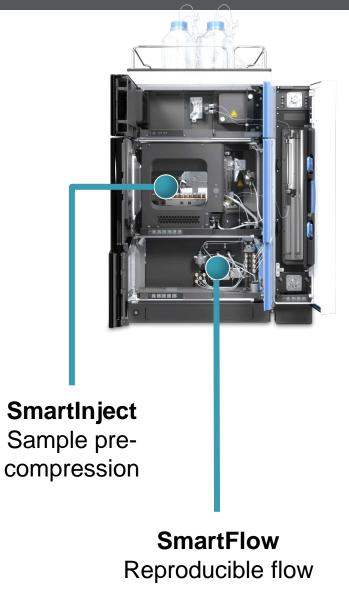
Component	Reference Rel. Area (%)	Unknown Rel. Area (%)
PTM 53	1.9	2.4
Peptide 54	2.6	-
Peptide 55	3.1	6.3
Peptide 56	6.7	3.5
Peptide 57	3.8	-
PTM 58	0.7	-



# Retention time precision Vanquish Flex Quaternary UHPLC System



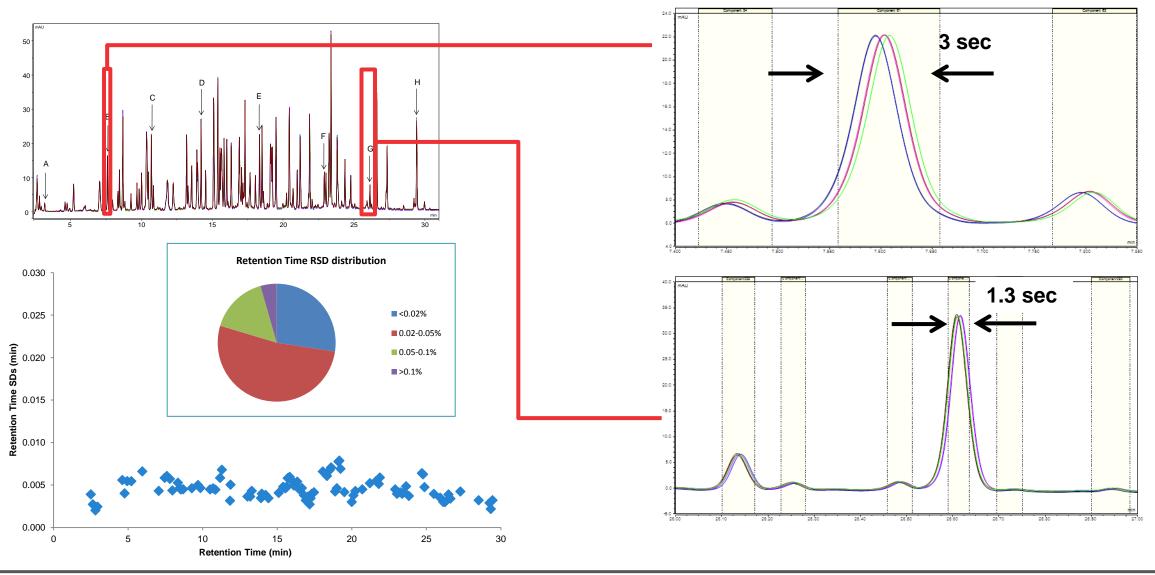






# What are the precision we expect from Vanquish Flex Quaternary?

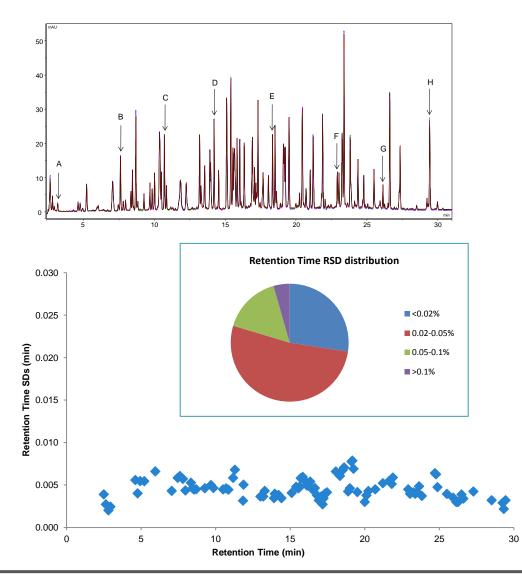
BSA tryptic digest (n=5) with Vanquish Flex Quaternary





# What are the precision we expect from Vanquish Flex Quaternary?

BSA tryptic digest (n=5) with Vanquish Flex Quaternary



Peak ID	Retention Time (min)	Retention Time RSD (%)	RSD Area (%)
А	3.171	0.077	0.61
В	7.601	0.077	0.31
С	10.702	0.042	0.22
D	14.217	0.028	0.24
Е	18.345	0.036	0.77
F	22.912	0.018	0.79
G	26.137	0.013	0.20
Н	29.438	0.007	2.10



1. Optimize resolving power of the peptide separation

- 2. Streamline peak assignment for efficient comparisons of different chromatograms
- 3. Importance of retention time precision on peak assignment
- 4. Ultra fast separation of peptides with great reproducibility



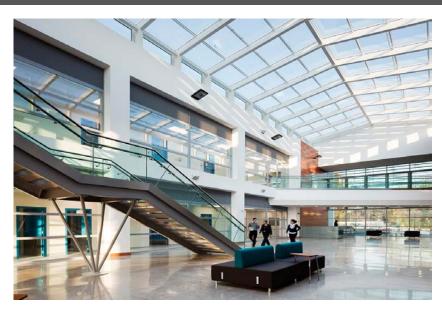


**Biopharmaceutical Peptide Mapping – A User Perspective** 



Amy Farrell, PhD

# About NIBRT





- World-class facility dedicated to address the training and research needs of the global biopharmaceutical industry.
- Unique competency based training experience in an environment that replicates the most modern industrial bioprocessing facilities.
- Research with impact developing solutions to address real challenges faced within the biopharmaceutical industry.
- CCL complete characterisation of bioprocesses and expressed therapeutic glycoproteins





# Benefits of Thermo Scientific Platform for Peptide Mapping Analysis



#### **UHPLC System Set-Up**

- Smart start-up, standby and shut-down
- eWorkflows https://appslab.thermofisher.com

# **SMART Digest**

- Time-Saving 30 minute digestion for broad sequence coverage
- User friendly reduction in sample preparation steps compared to traditional digest methods

# Vanquish Flex Quarternary UHPLC – Acclaim 120 C18 5 µm 2.1 x 250 mm analytical column

- Reproducibility
- Separation Efficiency





# **SMART** Digest Time Course Study – Light Chain Sequence Coverage

#### **Sample Preparation:**

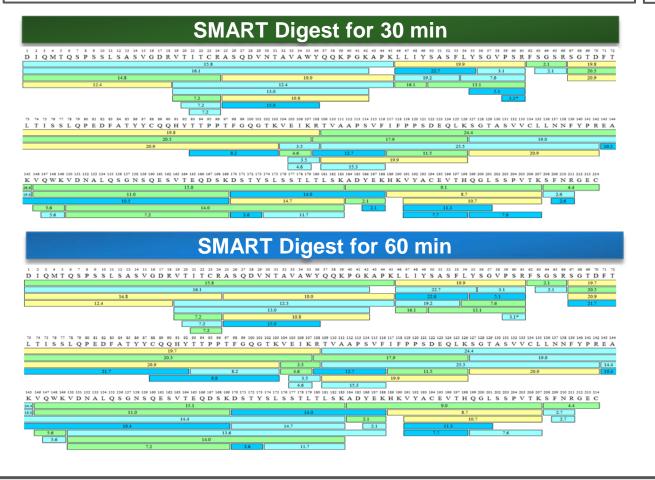
- 50 μL of 2 mg.mL<sup>-1</sup> IgG1 sample digested.
- Reduction with 50 mM TCEP preformed after digestion with Thermo Scientific SMART Digest Kits.

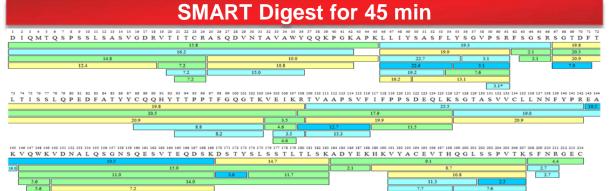
#### Sample Analysis:

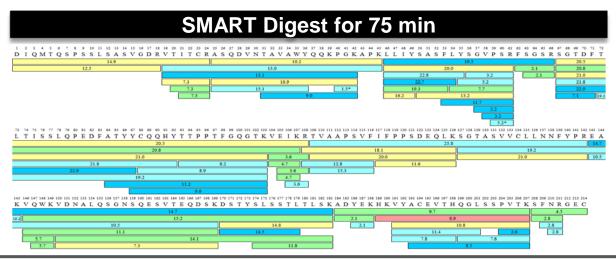
Acclaim 120 C18 2.1 x 250 mm.



- Thermo Scientific<sup>™</sup> Q Exactive<sup>™</sup> BioPharma platform
- Thermo Scientific<sup>™</sup> BioPharma Finder<sup>™</sup>

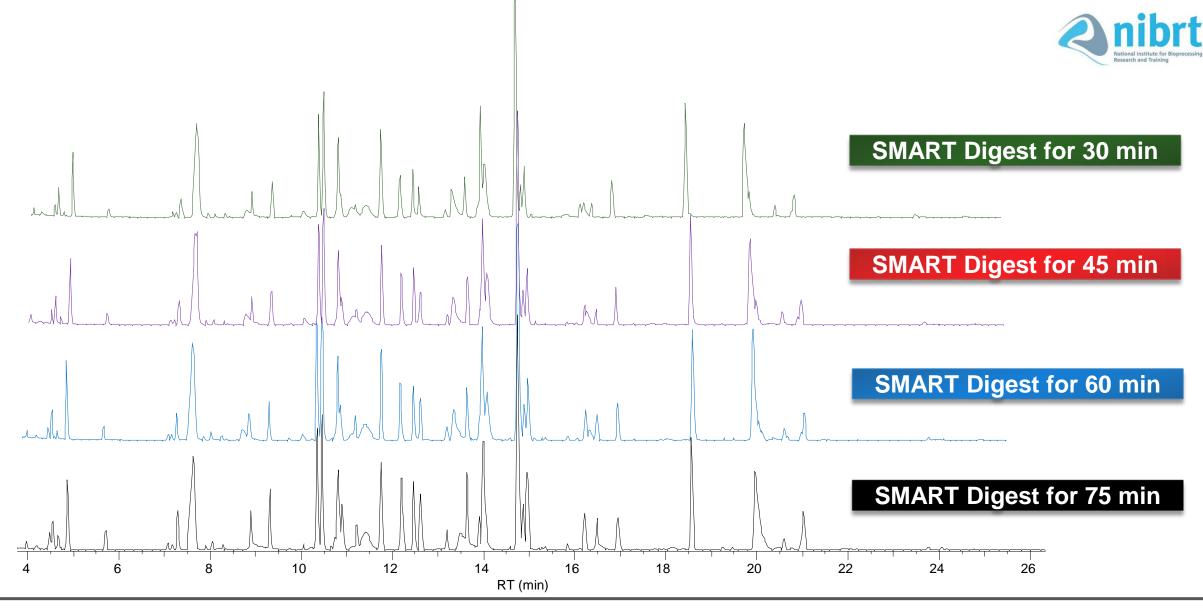






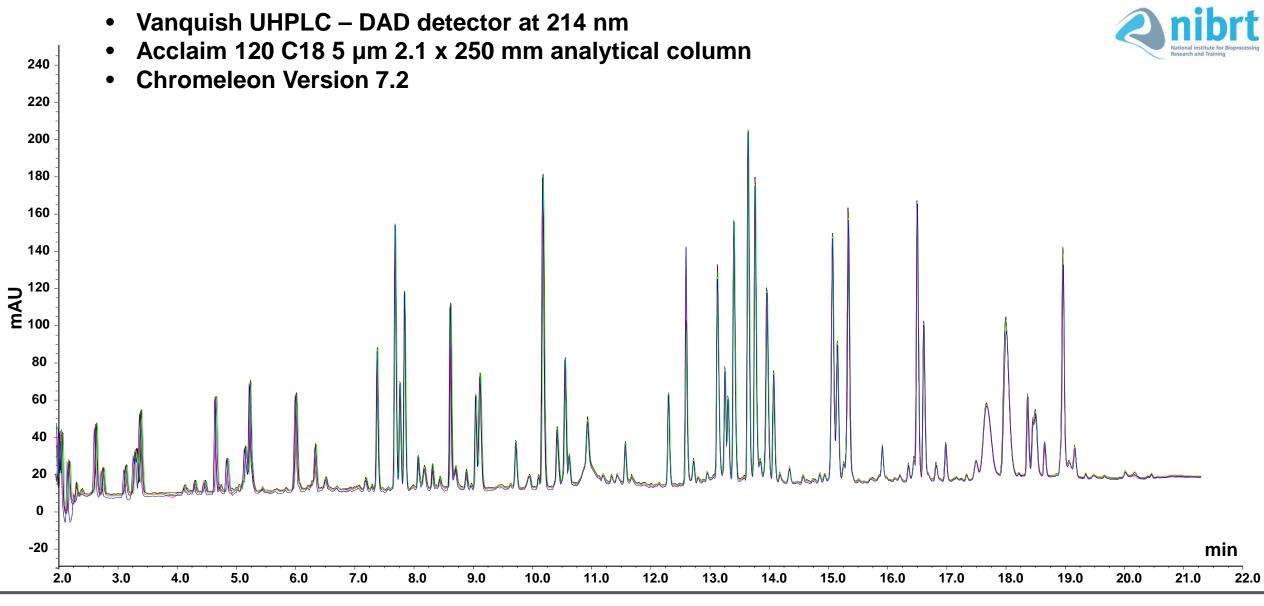


## **SMART** Digest Kits - Time Course Study – Total Ion Chromatograms





# Vanquish UHPLC - Reproducible Sample Separation





# Summary

**Identified beneficial attributes of the Chromeleon 7.2:** 

- ✓ Minimal hands-on time for system set-up
- ✓ Ability to download developed methods from AppsLab

## **Identified beneficial attributes of the Vanquish:**

✓ Retention time precision across multiple analyses

## **Identified beneficial attributes of SMART digest:**

- ✓ Efficient Monoclonal Antibody digestion in 30 minutes
- ✓ Full sequence coverage



