

# Vented Column Technology Applied to Proteomic MudPIT Analysis on Long Capillary Columns

Andrew W. Guzzetta and Allis S. Chien

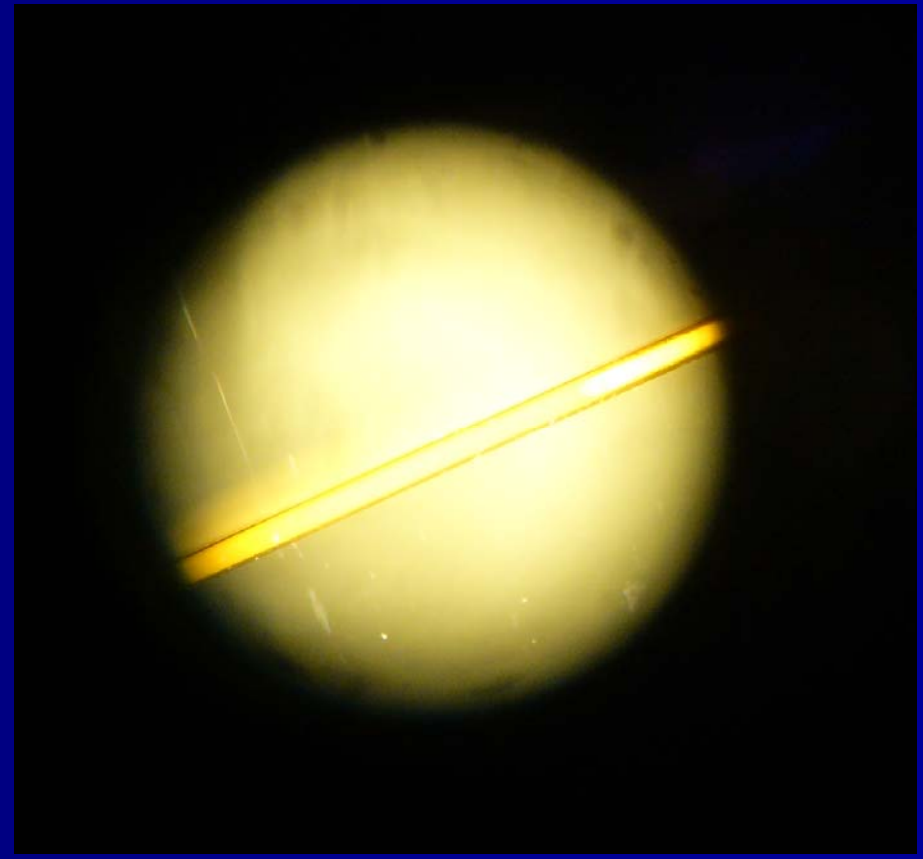
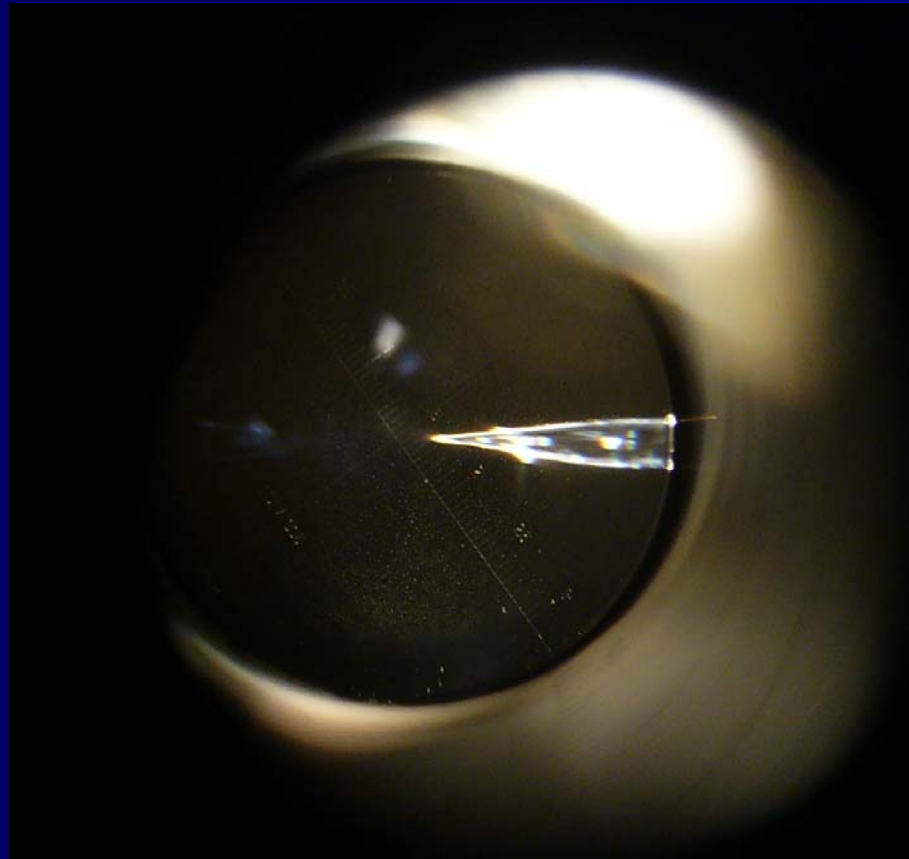
Vincent Coates Foundation  
Mass Spectrometry Laboratory  
Department of Chemistry  
Stanford University  
Stanford California

<http://mass-spec.stanford.edu>

Last year.... the long column and proteomics

# What if?

Space



# What if we could do MudPIT on a 60 cm column?

## Available Column Technologies

### Tri-Phasic MudPIT

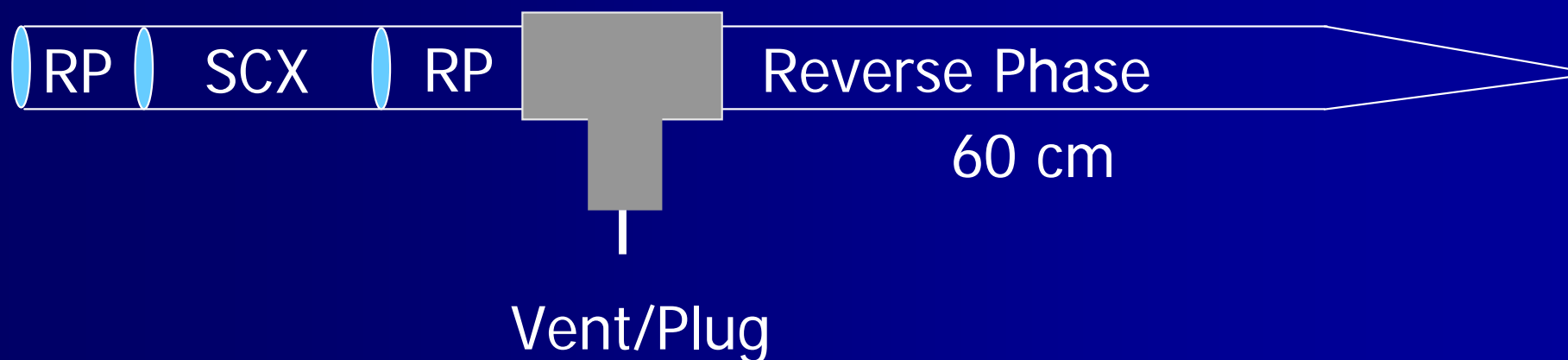


### Vented Technology



1. "Comparison of three directly coupled HPLC MS/MS strategies for identification of proteins from complex mixtures: single-dimension LC-MS/MS, 2-phase MudPIT, and 3-phase MudPIT." W. Hayes McDonald, Ryoma Ohi, David T. Miyamoto, Timothy J. Mitchison, and John R. Yates, III. *Int J. Mass Spectrom.* **2002**, 219 (1), 245-251.
2. "Automation of Nanoscale Microcapillary Liquid Chromatography-Tandem Mass Spectrometry a Vented Column." Lawrence J. Licklider, Carson C. Thoreen, Junmin Peng and Steven P. Gygi. *Anal. Chem.* **2002**, 74, 3076-3083.

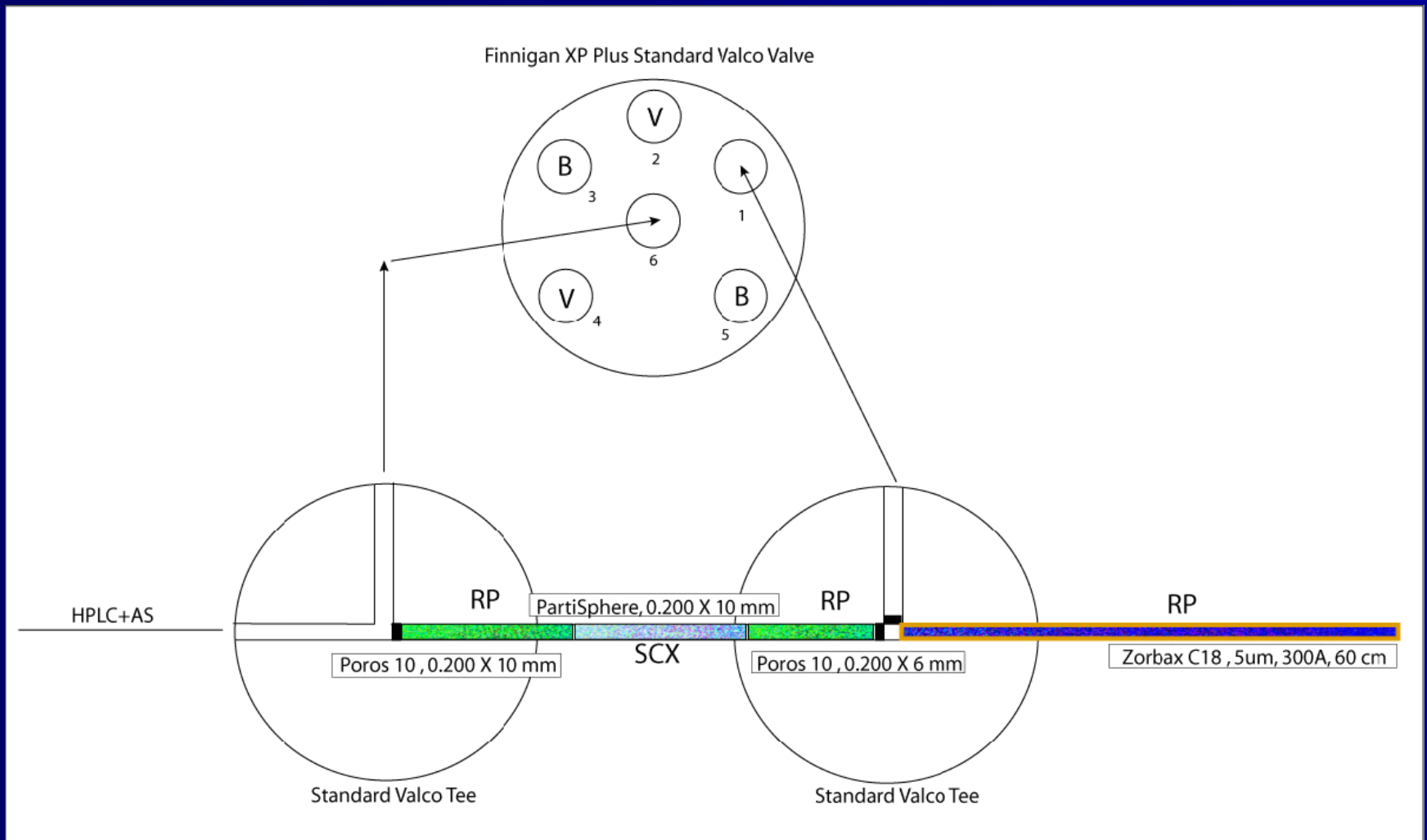
# MudPIT Meets Vented Column Technology Meets the Long Column



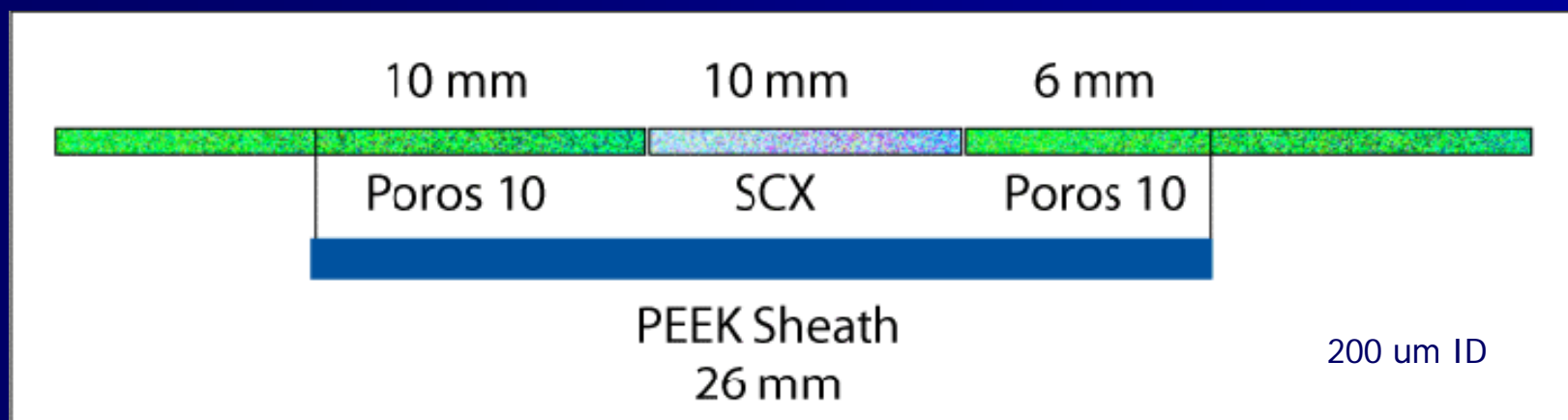
HPLC: Standard Flow Shimadzu 10ADVP with homemade splitter  
MS: XP Plus Ion Trap mass spectrometer

Guzzetta AW, Chien AS. Related Article: A double-vented tetraphasic continuous column approach to MudPIT analysis on long capillary columns demonstrates superior proteomic coverage.  
*J. Proteome Res.* **2005** Nov-Dec, 4 (6), 2412-9.

# Building the Double Vented Tetra Phasic MudPIT Device



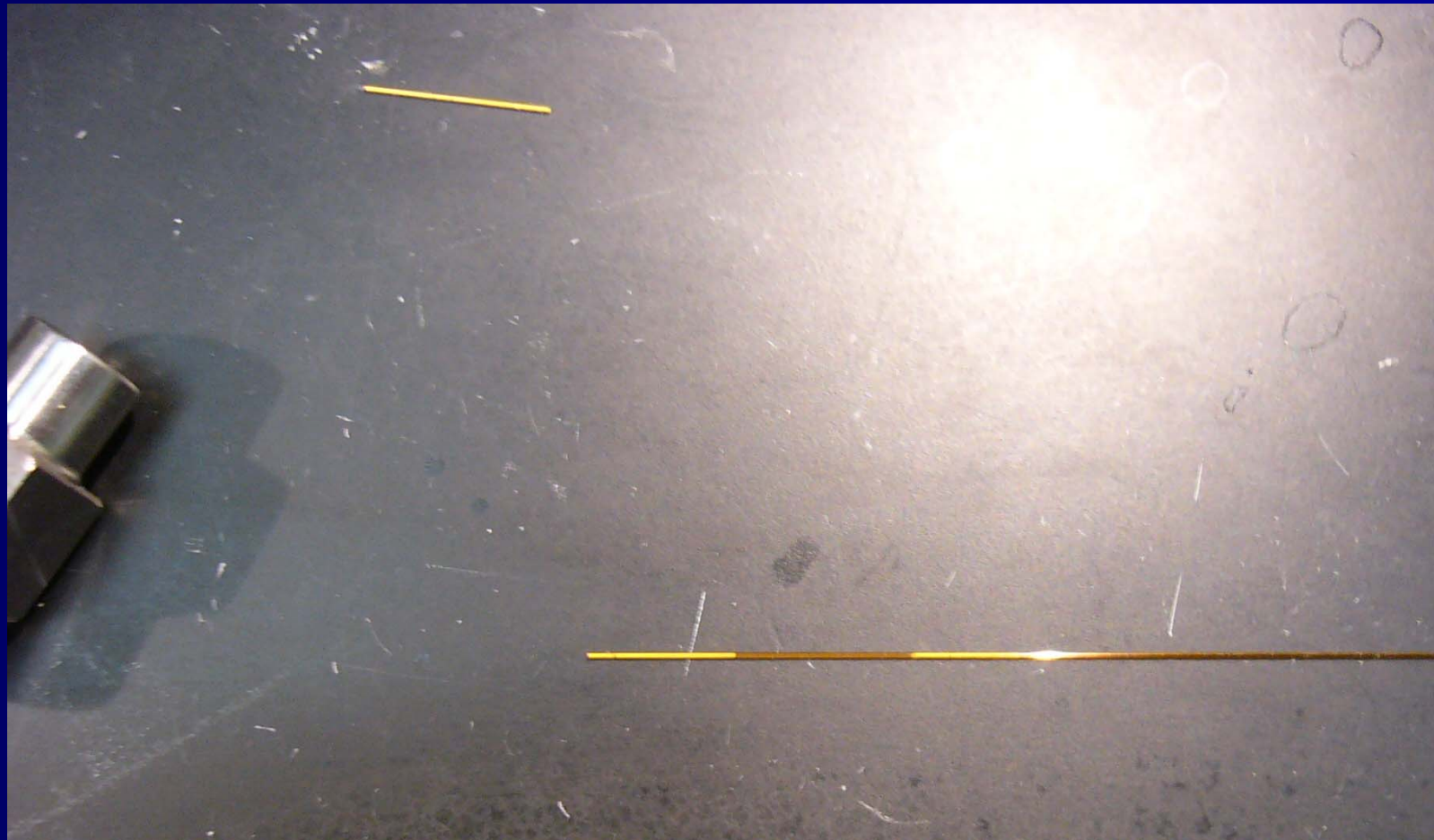
# Building the Triphasic MudPITter



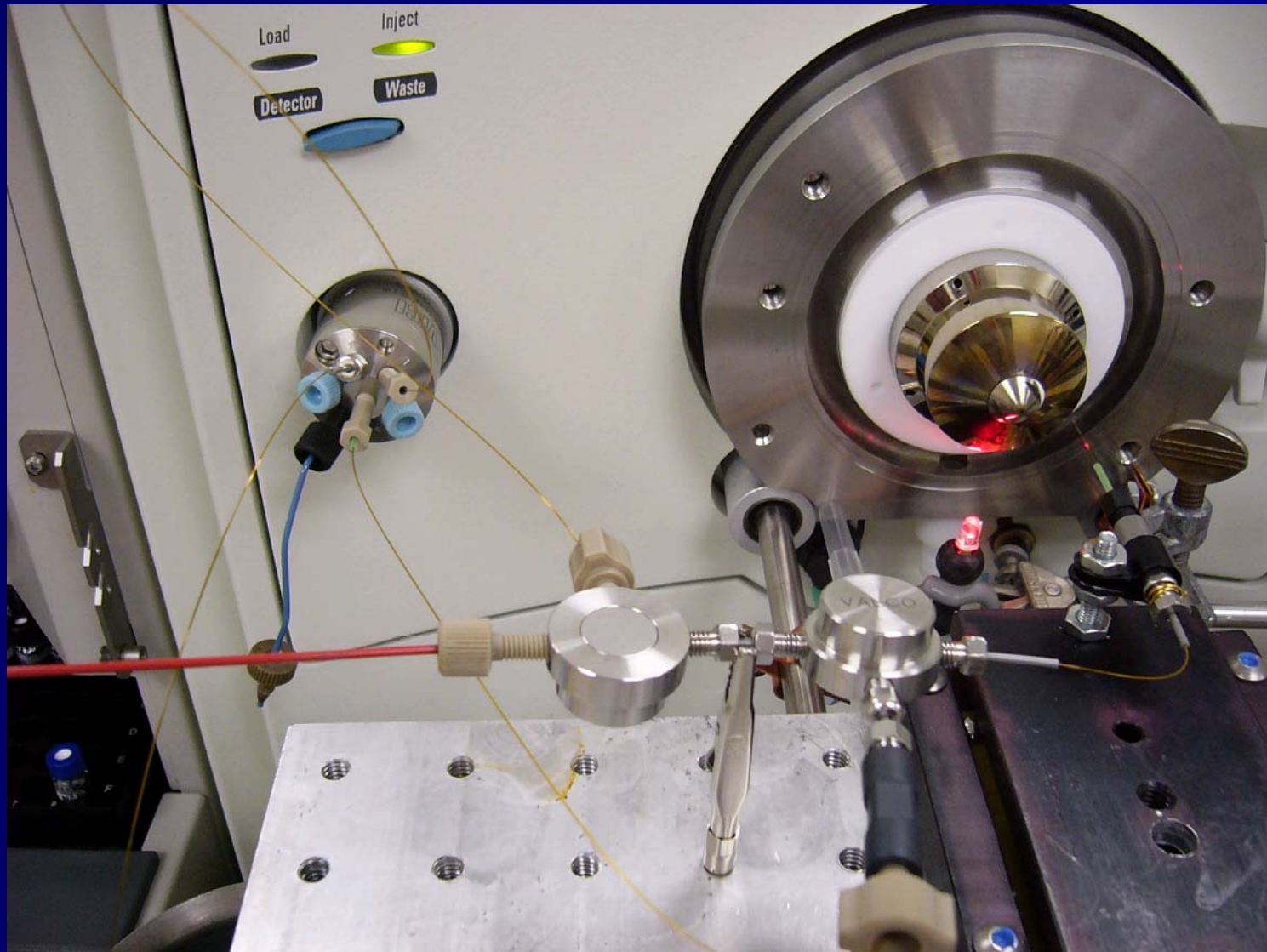
## Method

1. Place screen frit in the tee
2. Swage with short piece of PEEK
3. Pack Poros 10
4. Wash
5. Pack SCX (Whatman Partisphere, 5 μm)
6. Wash
7. Pack final section of Poros 10
8. Unswage
9. Cut first segment of Poros 10 to length
10. Reswage with new PEEK
11. Cut off final length of fused silica capillary

# Building the Triphasic MudPIT Column



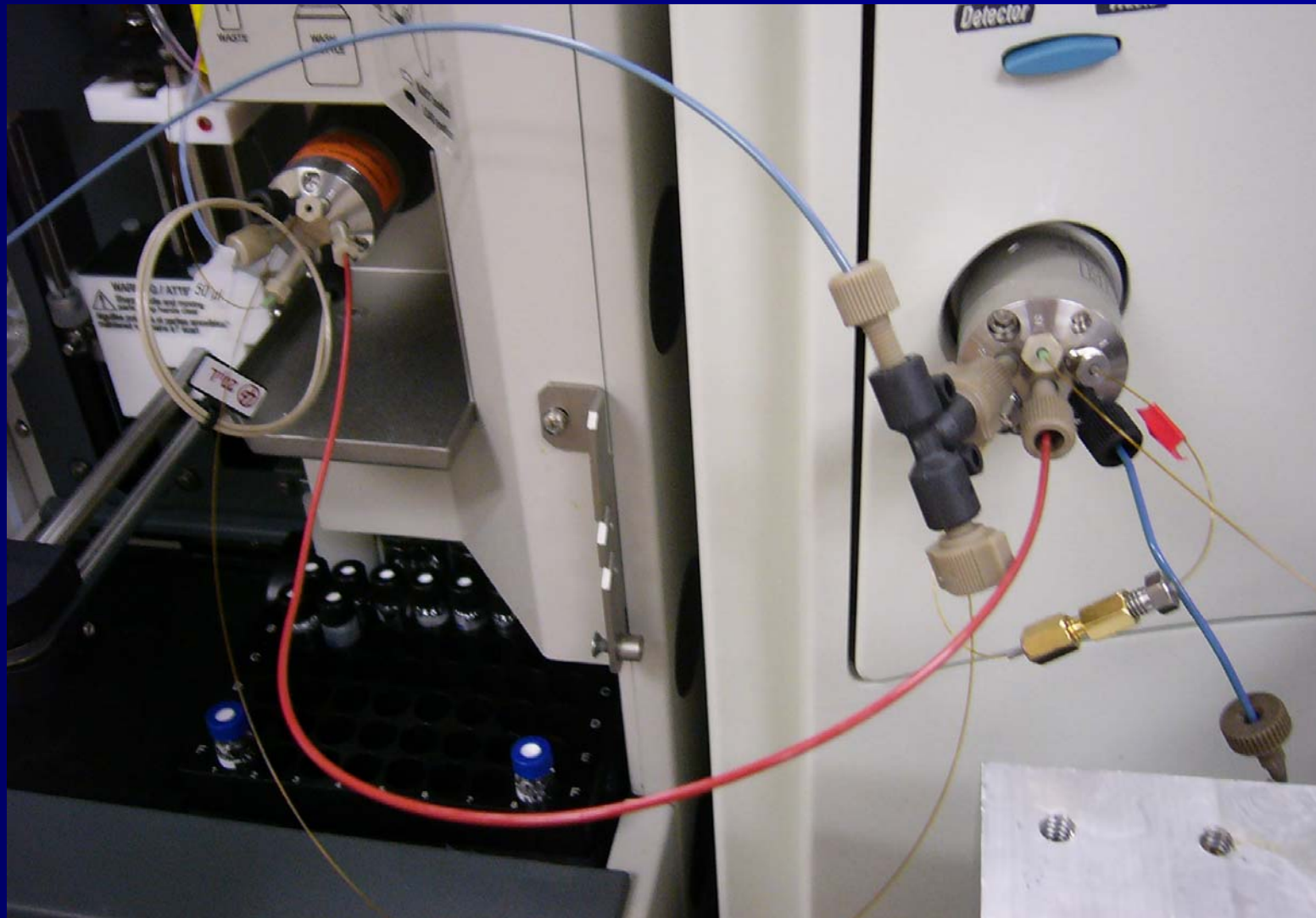
# Continuous Vent





# Discontinuous Vent

Tri-Phasic On The Valve



# MudPIT Gradients

Optimized for speed to test different models, not optimized for proteomic performance

Load  
Wash  
ACN bump

Gradient 17

Initial conditions  
30 $\mu$ l/min 0%B

T	%B
1	5
4	5
10	80
12	80
12.1	5
20	Stop

Purpose: Load sample onto first RP. Wash and elute onto SCX.

Salt &  
Separate

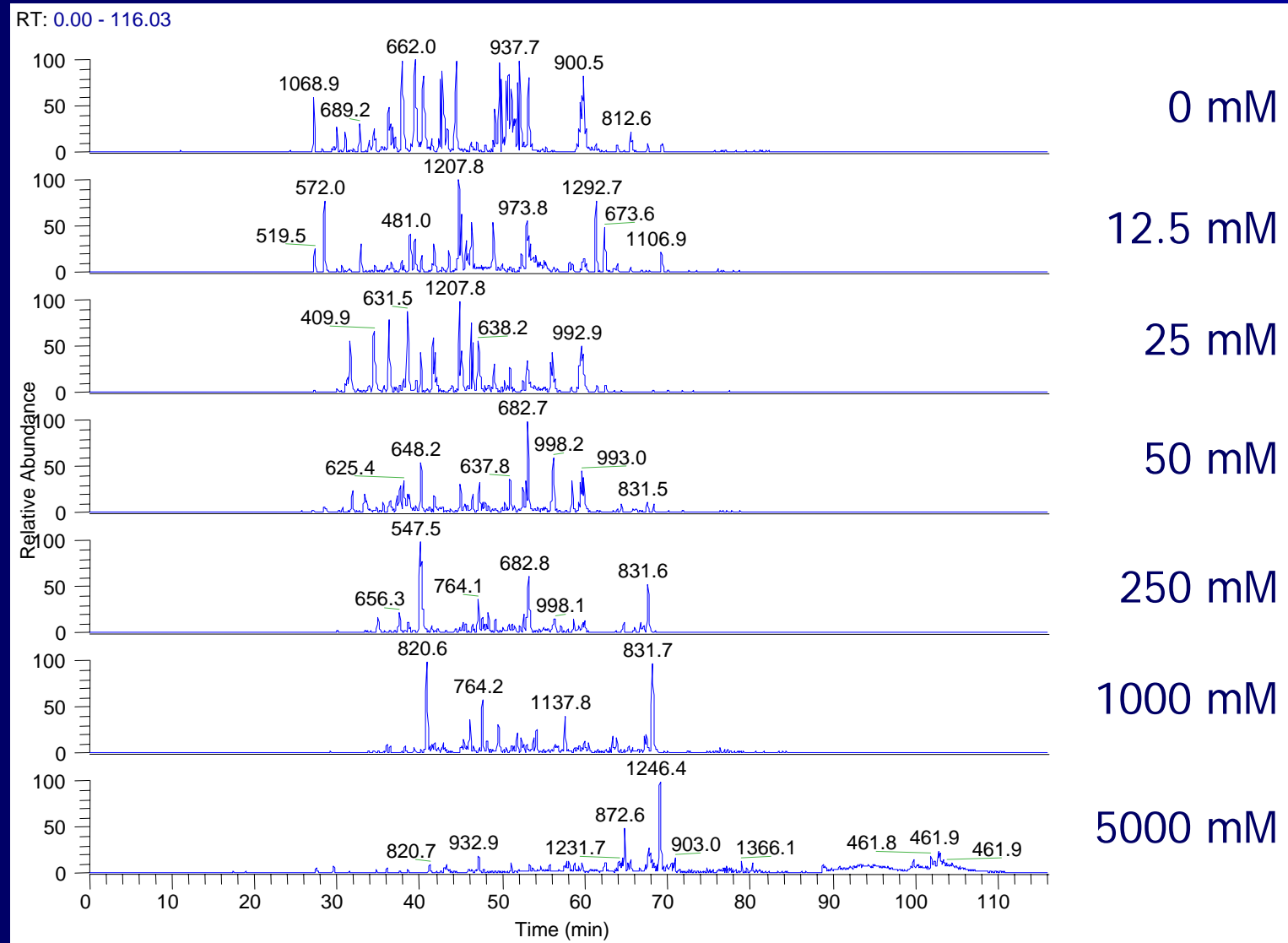
Gradient 18

Initial conditions  
30 $\mu$ l/min 0%B

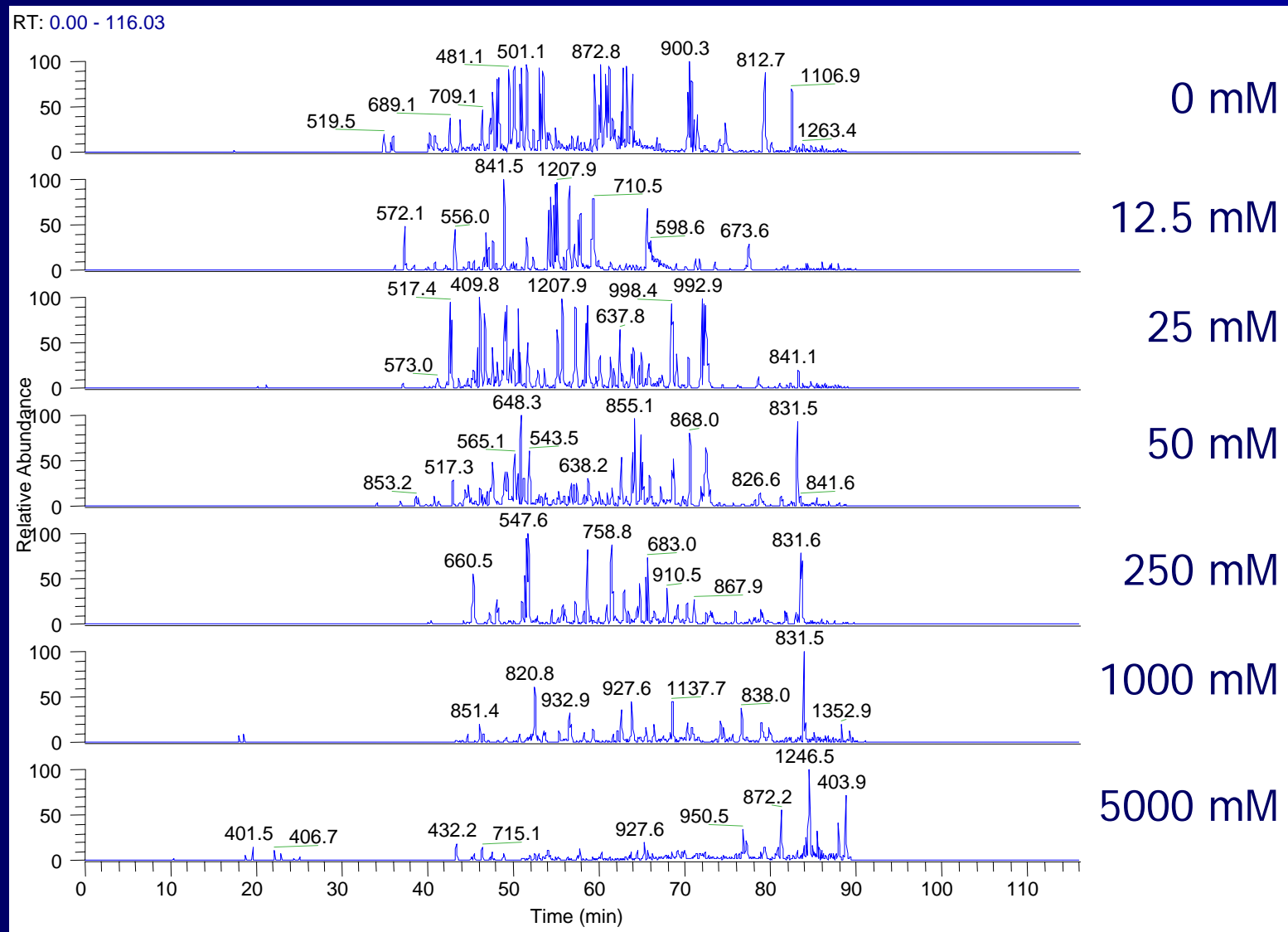
T	%B
1	5
10	5
70	40
80	80
80.1	5
120	Stop

Purpose: Salt bump. Separate peptides on the analytical column.

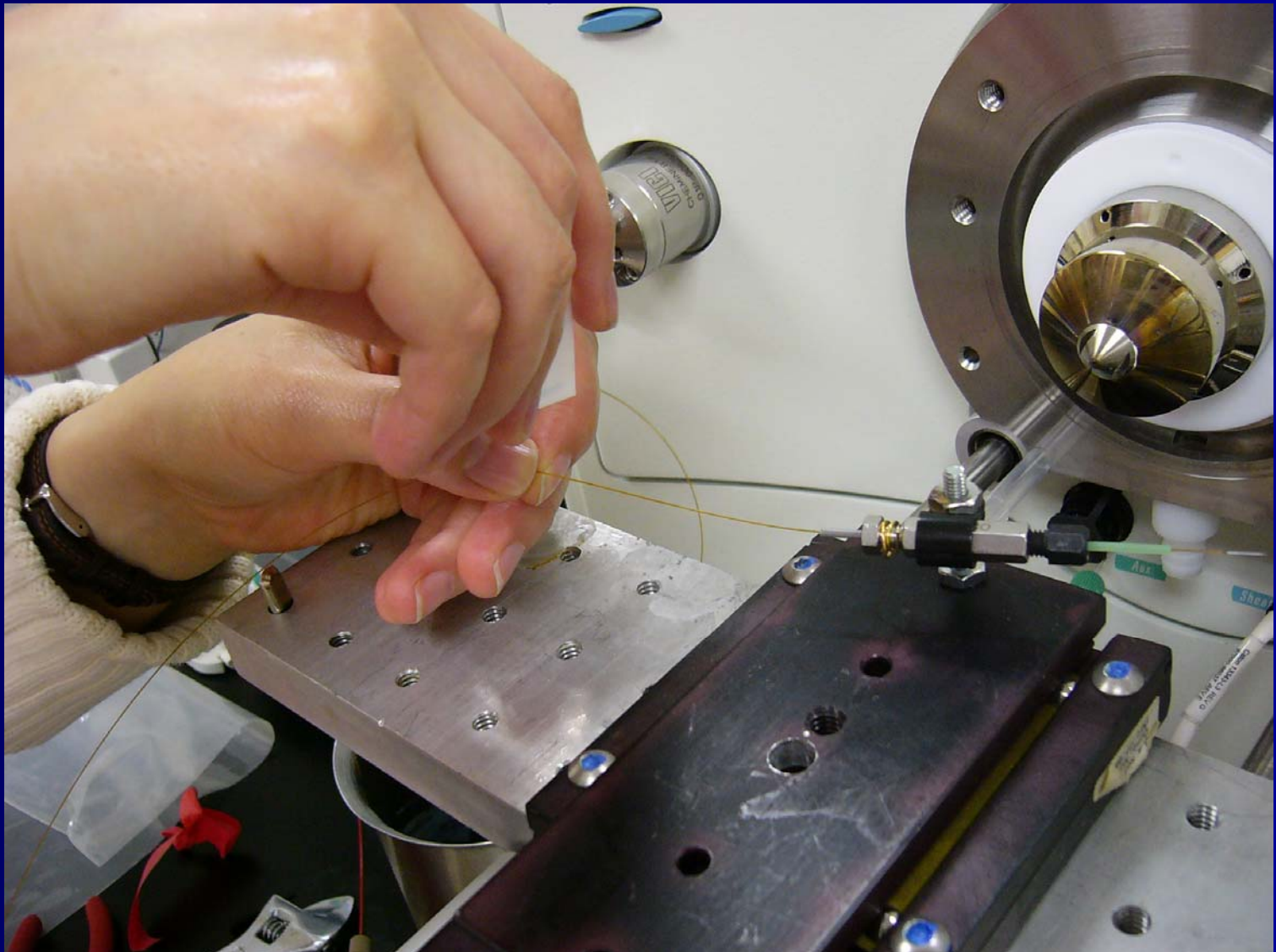
# 10 cm Continuous Vent Human Serum MudPIT



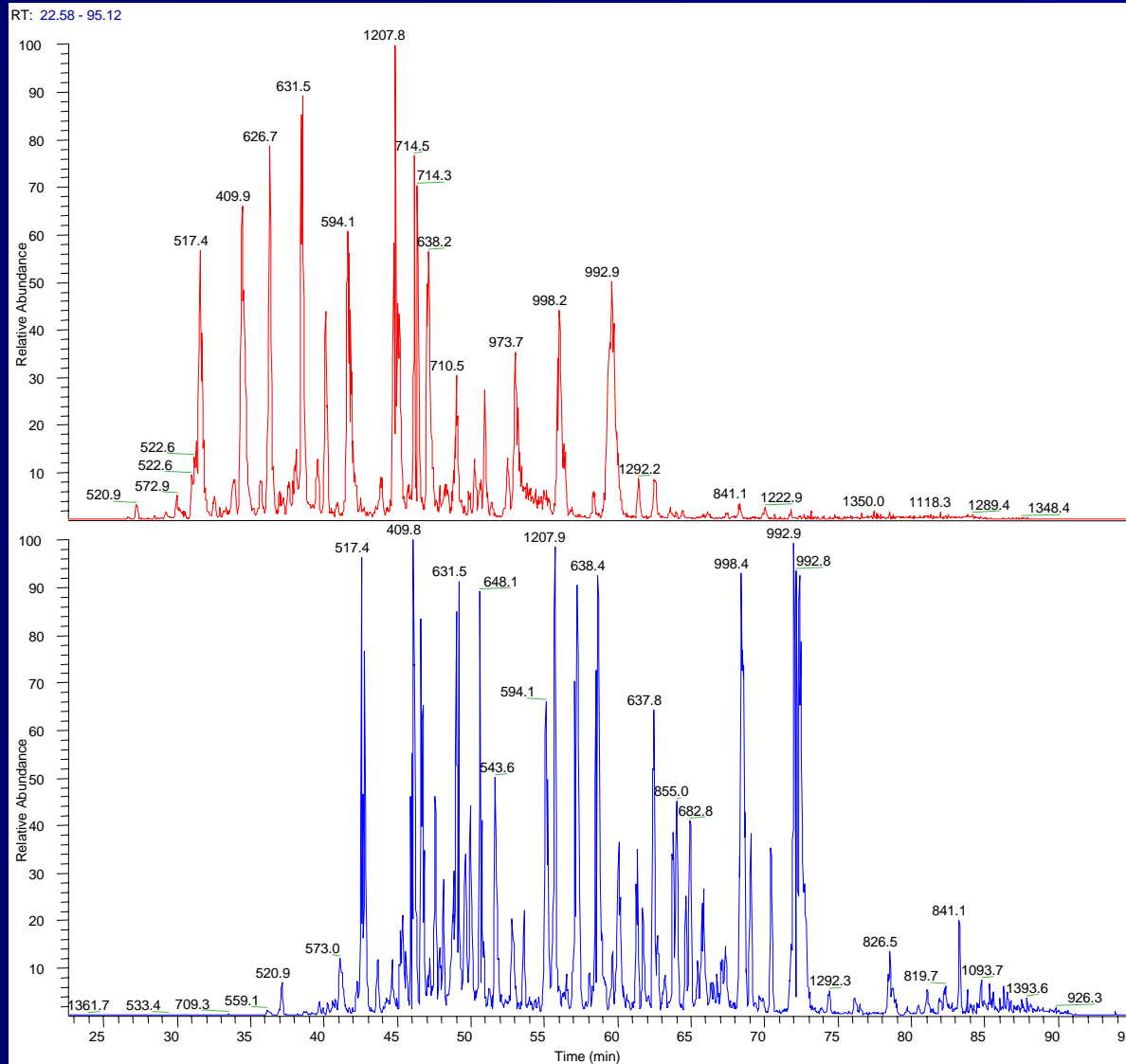
# 60 cm Continuous Vent Human Serum MudPIT



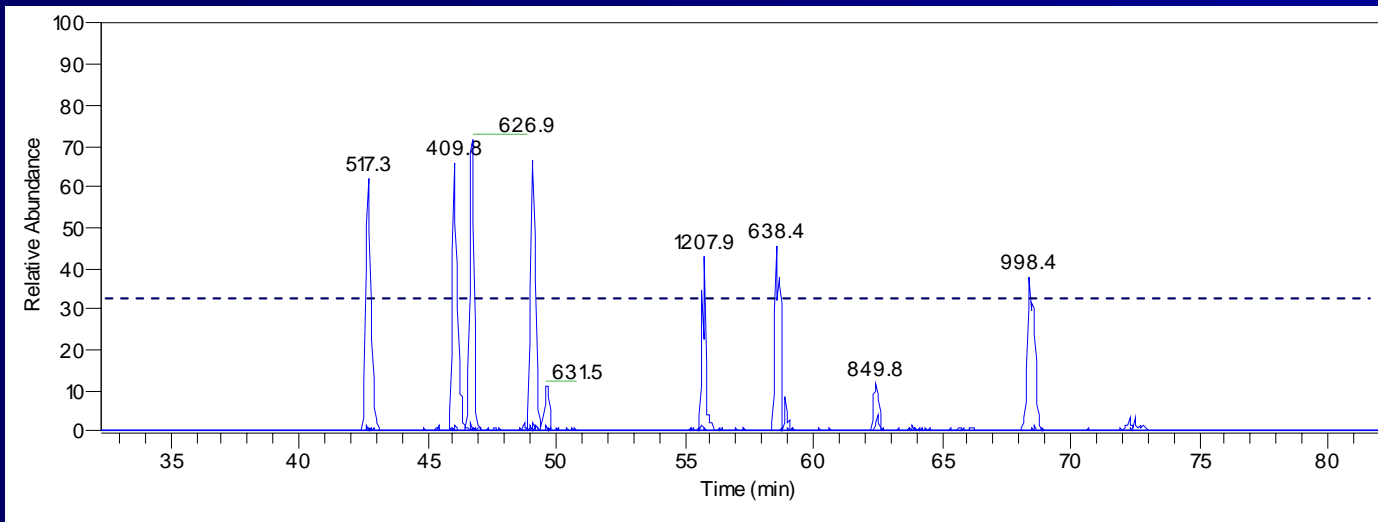
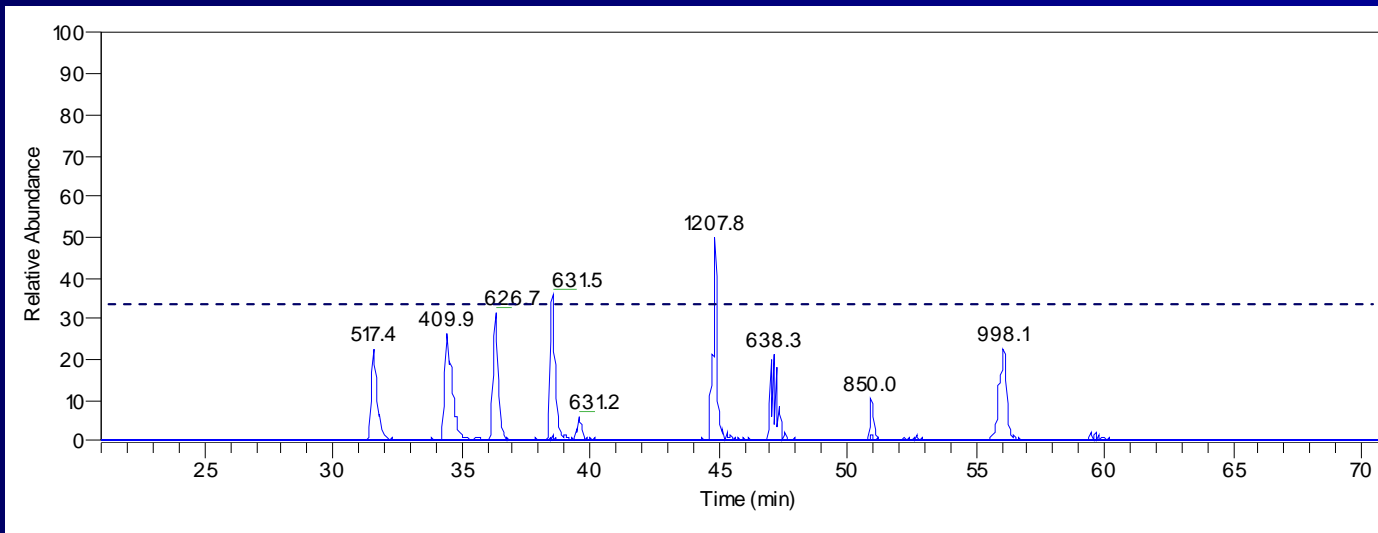
# Making the Cut



# Comparing 10 and 60 cm Columns at the 25 mM Salt Step using the Continuous Vent Method



# XIC Plots for 10 and 60 Columns at the 25 mM Salt Step using the Continuous Vent Method



# CV vs. DCV

Short vs. Long

	10 cm DCV	60 cm DCV	% Gain
Peptide Count	537	635	18.3
	10 cm CV	60 cm CV	
Peptide Count	593	739	24.6
% Gain	10.4	16.4	



# Our First Proteomics Evaluation Method: Counting Unique Peptides on the Mascot Significant Hits List

	10cmDCV	60cmDCV	10cmCV	60cmCV		10cmDCV	60cmDCV	10cmCV	60cmCV				
10cmDCV		60cmDCV	10cmCV	60cmCV	60cmCV	P08649	2	P10909	3	P02760	2	P02747	3
43232			47319		44212	P10909	2	P01031	3	P01764	2	P05546	3
P01024	53	P01024	61	P01024	75	P27169	2	P05546	3	P02748	2	O43866	3
P01023	50	P01023	50	P02768	60	P02766	2	P00760	2	P02763	2	P02753	3
P02768	46	P02768	45	P01023	51	P04220	1	P01786	2	P01617	2	P02748	3
P02787	25	P04114	34	P02787	35	P20742	1	O60841	2	P10643	2	P03952	3
P01028	23	P01028	31	P04114	32	P01766	1	P02766	2	Q28522	1	P01861	2
P00450	23	P02787	23	P01028	33	P01619	1	P04430	2	P01766	1	Q28372	2
P01857	18	P00450	19	P04114	29	P02743	1	P41339	2	P01786	1	P01625	2
P04114	18	P00747	17	P08603	17	P01612	1	P19652	2	P49065	1	O14791	2
P00747	15	Q14624	17	P01857	16	Q03591	1	P01019	2	P01742	1	P01766	2
P08603	14	P01857	15	P00450	15	P04433	1	P01605	2	P01765	1	P25311	2
P01871	13	P01871	15	P01009	14	P01742	1	P05155	2	Q59072	1	P01614	2
P02774	12	P01009	15	P02647	14	P18902	1	P02763	2	P01781	1	P01031	2
P02647	11	P02647	15	P00738	13	P01631	1	P01625	2	P01700	1	pos155	2
P01009	10	P02774	14	P00738	14	P13671	1	P04196	2	P05546	1	P01598	2
P19827	10	P19827	14	P00747	11	P11426	1	Q28522	1	P01593	1	P01611	2
P02790	10	P08603	13	P19823	10	P01765	1	P04220	1	Q6RE44	1	P02743	2
P01859	9	P19823	12	P02774	10	O18500	1	P35568	1	P01621	1	P13671	2
P01876	9	P01859	9	P04217	10	P01825	1	P97279	1	Q61838	1	P02775	2
P00738	9	P01876	9	P01876	9	P05156	1	P01596	1	P01784	1	P10643	2
Q14624	9	P04217	8	P00747	13	Q60841	1	P02749	1	P08649	1	Q28522	1
P04003	9	P02790	8	P04217	10	P04196	1	P46782	1	P01785	1	P02769	1
P19823	8	P02751	8	P01042	9	P01591	1	P01598	1	O63270	1	P04220	1
P01042	7	P04003	8	P20742	8	P01611	1	P01593	1	Q8S7P9	1	P01877	1
P01834	6	P01834	7	P01859	7	P01617	1	P01619	1	P01029	1	P49065	1
P01842	6	Q95LBO	7	P01011	8	P01700	1	P08649	1	P18527	1	P01766	1
P01011	6	P01842	6	P01842	7	P35827	1	P01717	1	P01973	1	P01762	1
P04217	6	P01011	6	P02765	7	P01784	1	P14227	1	Q9UGM5	1	P08649	1
P02749	6	P01042	6	P00734	7	P01763	1	P01700	1	P01700	1	P01770	1
P02751	6	P00738	6	P06727	6					P27169	1	P01742	1
P02765	5	P02765	5	P01860	5					Q9UGM5	1	P07360	1
P00734	5	P20742	5	P01011	5					P22752	1	P01779	1
P01860	4	P02760	5	P01620	4					P08185	1	P13635	1
P06396	4	P06396	5	P10909	6					P01765	1	P06310	1
O43866	4	P04004	4	P01019	4					Q28372	1	P01714	1
P01861	3	P00734	4	P04004	4					P04433	1	P01612	1
P80748	3	P09871	4	O43866	4					P98137	1	Q03591	1
P02652	3	P00739	4	P06396	4					P02745	1	P01700	1
P04004	3	O43866	4	P02790	3					P37093	1	P07225	1
P00760	2	P02748	4	P02751	3					Q03591	1	P01743	1
P01625	2	P00751	4	P02760	3					Q62910	1	pos090	1
P01597	2	P01861	3	P01019	4					P02746	1	P18982	1
P01620	2	P01860	3	P02751	3							Q864W0	1
P01019	2	P01597	3	P02763	3	537		635				P80108	1
P00739	2	P02652	3	P01787	3							O84877	1
P46782	2	P02753	3	P00751	3							Q59072	1
P01714	2	P01620	3	P02760	3							P01806	1
P01616	2	P03952	3	P01787	3							P50042	1
Q9UGM5	2	P06727	3	P00751	3							P00736	1
P01764	2	P80748	3	P01620	3							P01825	1
P08649	2	P10909	3	P02652	3							P01935	1
P10909	2	P01031	3	P02749	3							Q97RH2	1
P27169	2	P05546	3	P00739	2							P97313	1
P02766	2	P00760	2	P01861	2							P01717	1
P04220	1	P01786	2	P01606	3							Q09687	1
P20742	1	O60841	2	P80748	3							P04937	1
P01766	1	P02766	2	P00751	3							P01618	1
P01619	1	P04430	2	P02747	3							P01767	1
				P01031	3							P33449	1
				P02748	3							Q13619	1
				P05546	3								
				O43866	3								
				P02753	3								
				P02766	3								
				P01617	3								
				P03952	3								
				P01861	2								
				Q28522	1								
				P01766	2								
				P01766	2								

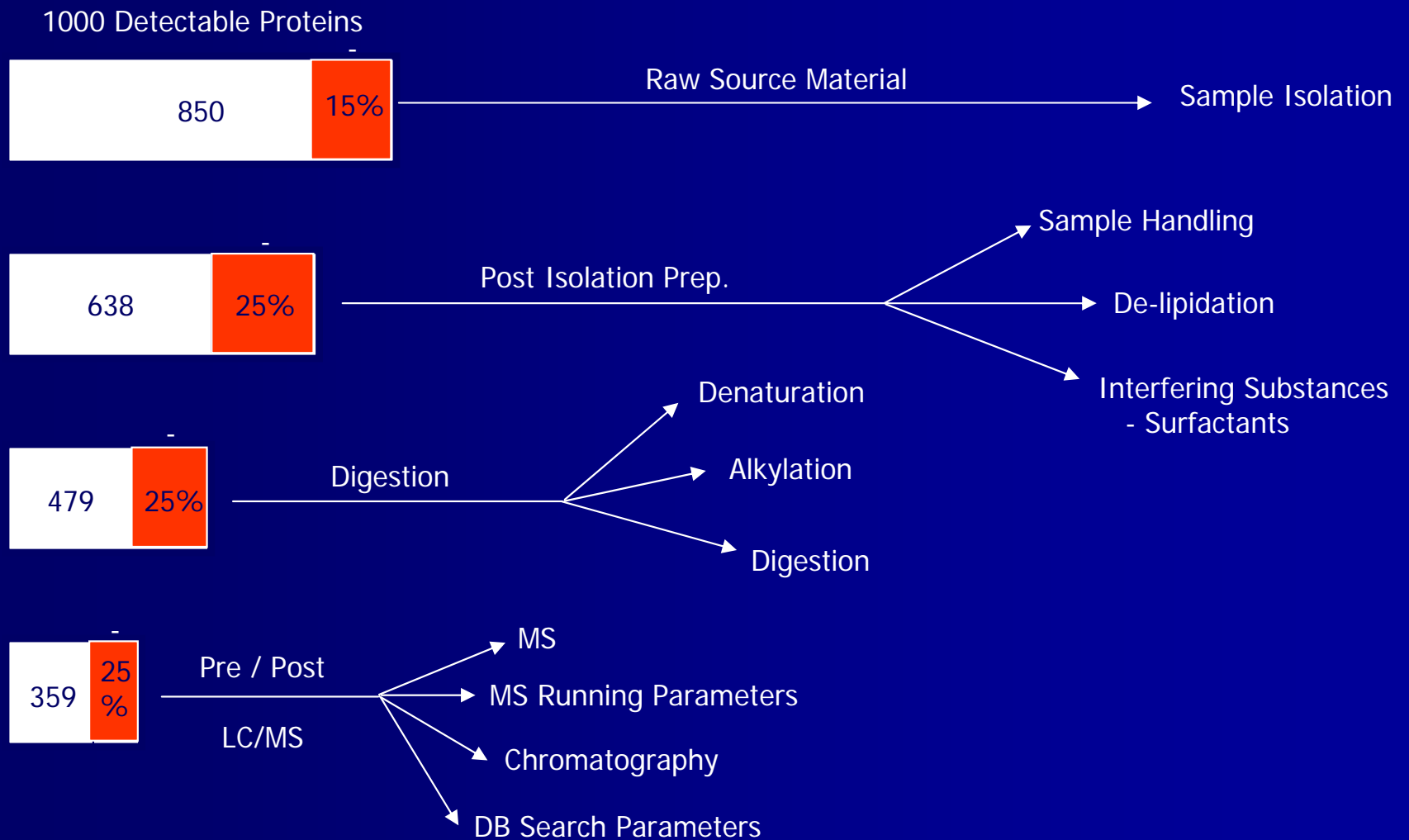
# CV vs. DCV

Short vs. Long

Peptide Count Category <small>(protein with n peptides)</small>	10 cm DCV	60 cm DCV	% Gain
6+	447	498	11.4
1-5	90	137	52.2
	10 cm CV	60 cm CV	
6+	486	589	21.2
1-5	107	150	40.2

# Improving the Efficiency of the Process

Worshipping at the altar of incrementalism

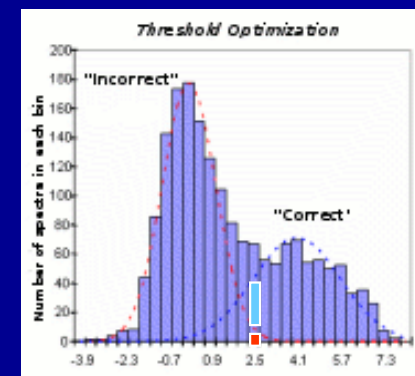
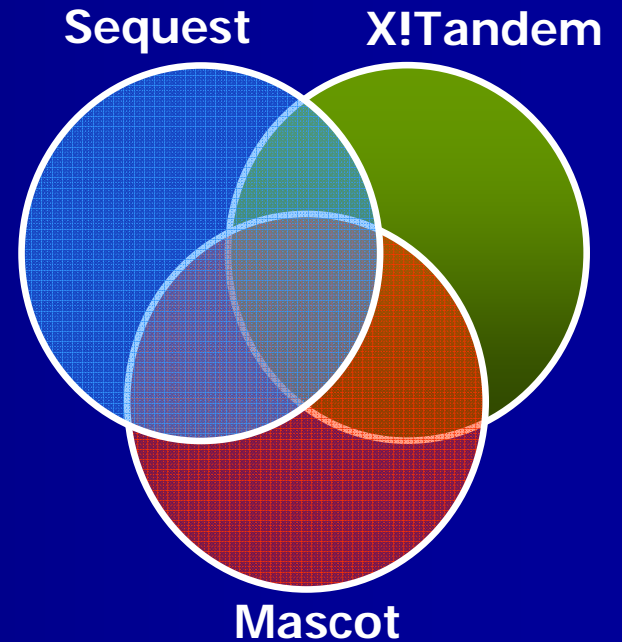


# Scaffold Proteomics Evaluation Method

The screenshot shows the Scaffold software interface with the following elements:

- Search Parameters:**
  - Min Protein: 95...
  - Min # Peptides: 1
  - Min Peptide: 80%
- Probability Legend:**
  - over 95% (Green)
  - 80% to 94% (Yellow)
  - 50% to 79% (Orange)
  - 20% to 49% (Red)
  - 0% to 19% (Dark Red)
- Bio View:**
  - # Identified Proteins (420)
- Table of Identified Proteins:**

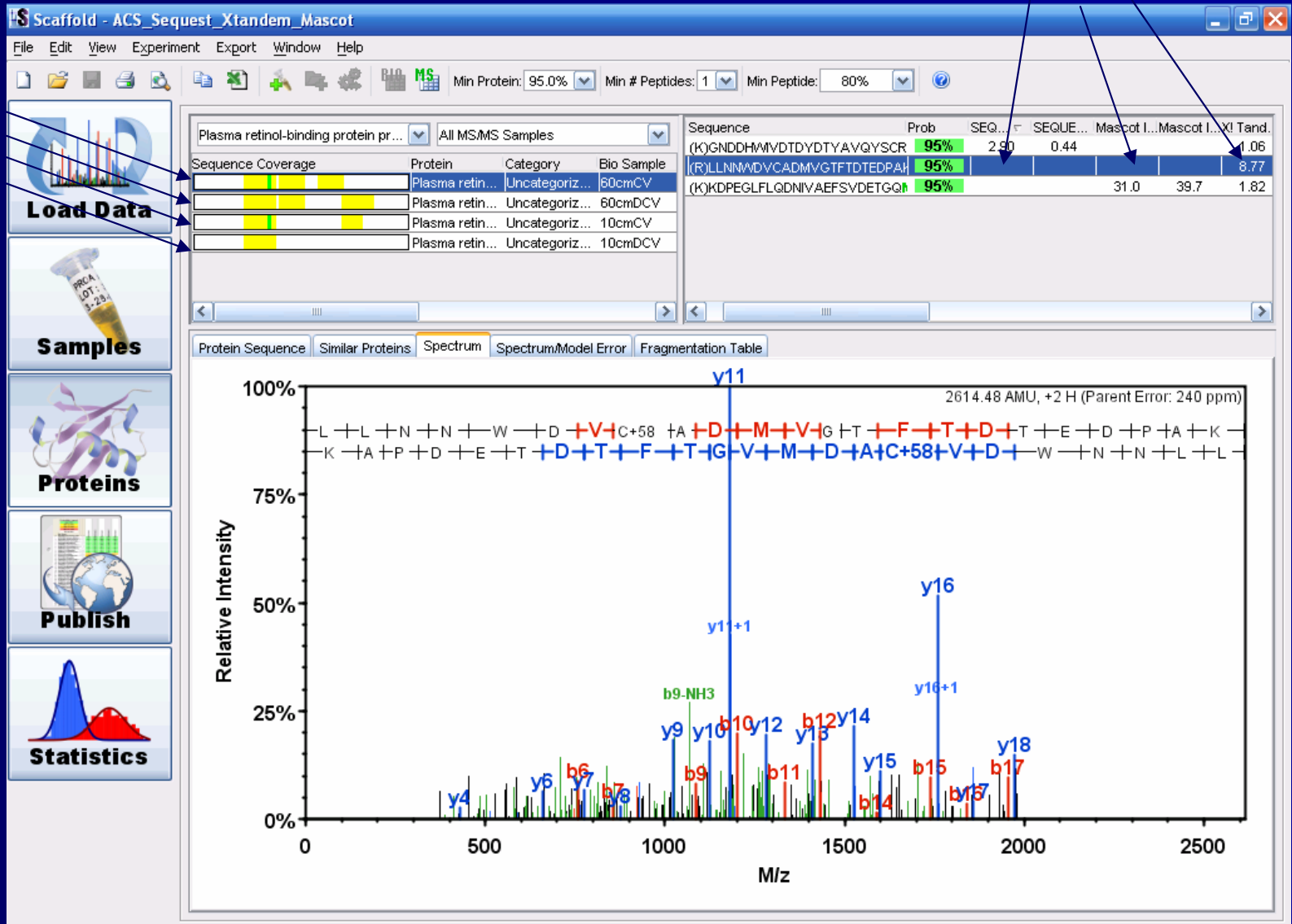
#	Protein Name	Accession	Weight	10cmDCV	10cmDCV	80cmDCV	80cmDCV
1	Complement C3 precursor [Contai...	CO3_HUMAN	187 kDa	55	67	58	77
2	Serum albumin precursor	ALBU_HUMAN	69 kDa	55	72	53	71
3	Alpha-2-macroglobulin precursor (... A2MG_HUMAN	A2MG_HUMAN	163 kDa	48	48	44	51
4	Serotransferrin precursor (Transf...	TRFE_HUMAN	77 kDa	32	47	27	43
5	Apolipoprotein B-100 precursor (A...	APOB_HUMAN	516 kDa	20	34	31	35
6	Complement C4-A precursor (Acid...	CO4A_HUMAN	193 kDa	26	25	32	34
7	Ceruloplasmin precursor (EC 1.16...	CERU_HUMAN	122 kDa	25	22	19	23
8	Complement factor H precursor (H...	CFAH_HUMAN	139 kDa	17	22	14	21
9	Apolipoprotein A-I precursor (Apo...	APOA1_HUM...	31 kDa	12	17	15	18
10	Alpha-1-antitrypsin precursor (Alp...	A1AT_HUMAN	47 kDa	13	15	13	17
11	Ig gamma-1 chain C region	IGHG1_HUMAN	36 kDa	16	12	13	16
12	Ig mu chain C region	MUC_HUMAN	50 kDa	12	13	14	16
13	Vitamin D-binding protein precurs...	VTDB_HUMAN	53 kDa	11	14	16	15
14	Inter-alpha-trypsin inhibitor heavy ...	ITI2_HUMAN	106 kDa	11	13	13	15
15	Inter-alpha-trypsin inhibitor heavy ...	ITI4_HUMAN	103 kDa	11	15	18	14
16	Inter-alpha-trypsin inhibitor heavy ...	ITI1_HUMAN	101 kDa	11	13	15	14
17	Haptoglobin precursor [Contains...	HPT_HUMAN	?	7	5	6	14
- Protein Information:**
  - Lookup Accession Number In: NCBI (ie: gi|135...
  - CO3\_HUMAN
  - Protein Name: Complement C3 precursor [Cont
- Sample Information:**
  - Biological Sample:
  - Sample Category:
  - Sample Description:
  - MSMS Sample:
  - MSMS Sample Notes:



# Example

50<sup>th</sup> on the list

60cmCV  
60cmDCV  
10cmCV  
10cmDCV



# Reworking the Data With Scaffold

## CV vs. DCV

Short vs. Long

	10 cm DCV	60 cm DCV	% Gain
Peptide Count	728	829	14 %
original Mascot numbers ->	537 (36%)	635 (31%)	
	10 cm CV	60 cm CV	
Peptide Count	787	894	14 %
original Mascot numbers ->	593 (33%)	739 (21%)	
% Gain	8 %	8 %	166 / 23%

# Conclusion

- Long columns are better at showing us a greater number of low level peptides.
- The CV method is superior to the DCV method.
- Be an incrementalist.

# Acknowledgements

- The Vincent & Stella Coates Foundation
- Mike MacCoss, University of Washington
- Andy Gieschen, Agilent
- ThermoFinnigan