

### The Applications of Empore<sup>TM</sup> Membrane SPE for Proteomics

**May 2020** 

**CDS Analytical** 

#### **Outline**

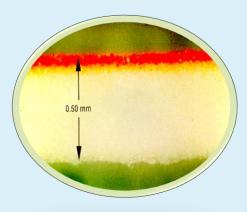


- 1. Empore SPE Introduction
- 2. Application in Proteomics

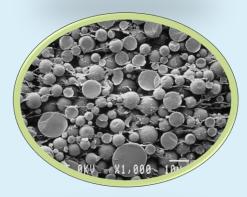
## **Empore™ Technology Highlights**



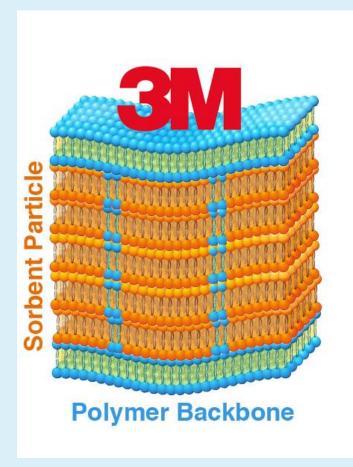
3M™ patented SPE particle-loaded membrane



Best Uniformity



High Density





**Least Elution Volume** 









Reduce Particle Fines

CDS Analytical, LLC

#### **Empore™ Portfolio**





- Empore<sup>™</sup> by Package Style
  - Disks
  - Cartridges
  - Plates
  - StageTips
- Empore<sup>™</sup> by Sorbent Style

Reverses Phases: C8, C18, UR, SDB-

XC, Activated Carbon, Oil & Grease

Mixed Phases: MPC, UR, SDB-RPS

Ion Exchange: Cation, Anion, Chelating,

SDB-RPS



#### **Empore™ Unique Features**



- Ultra-fast flow rate up to 700ml/min 2X faster than other leading brands
- Least elution volume 1/3 of other leading brands and 1/10 of loose-packed SPE
- Highest consistency & reproducibility 10-15% higher than Waters Oasis plates
- Least particle fines in eluates -1/10 of other leading brands to increase efficiency (reducing tube clogging & system downtime)



#### **Recommended in Dozens of EPA Methods:**

1664 (Rev. A) - N- Hexane Extrachable Material (HEM; Oil and Grease) 506 - Phthalate and Adipate Esters in Drinking Water 507 - Nitrogen- and Phosphorous-Containing Pesticides in Water - Chlorinated Pesticides, Herbicides, and Organohalides in Water 508.1 512.2 - Chlorinated Acids in Water 525.3 - Organic Compounds in Drinking Water 549.1 - Diquat and Paraquat in Drinking Water - Polycyclic Aromatic Hydrocarbons in Drinking Water 550.1 552.1 - Haloacetic Acids and Dalapon in Drinking Water - Benzidines and Nitrogen-Containing Pesticides in Water 553 - Tetra-Through Octa- Chlorinated Dioxins and Furans by 1613 (Rev. B) **Isotope Dilution e.g. in Water** SW846 method 3535 – Test Methods for TCLP Leachates - Aqueous Phases Quick Turnaround Methods **QTM** - PAH - Phenols - Pesticides & PCBs

#### **Typical Customers:**

Environmental











Food and Agricultural







Pharmaceutical/Clinical











Research









# CDS has rebuilt Empore production line in a new-construct, GMP-compliant, clean room facility at Oxford, PA.





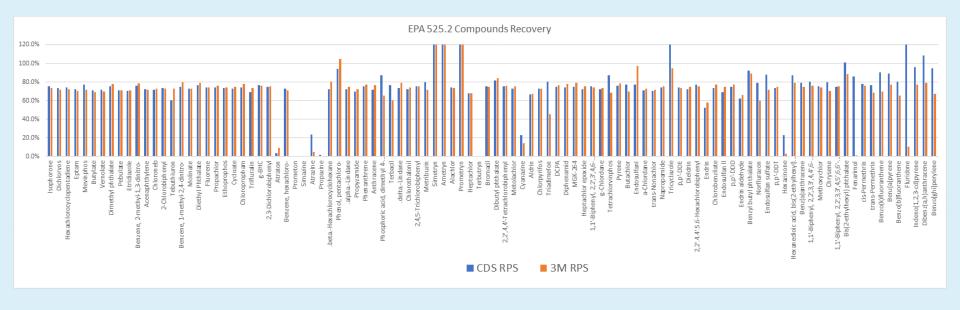


Empore production line at 3M

Empore new production line at CDS

### **Empore Disks with Improved Quality at CDS than at 3M**





EPA 8141.B Organophosphorous Pesiticides Recoveries with CDS Empore SDB-RPS Disk





### Part 2 Empore Disks' Applications in Proteomics

- 1. Empore StageTips Basics
- 2. Applications in Proteomics
- 3. High-pH Fractionation for Global Proteomics

#### **Empore StageTips Basics**



### Preparation of Single-Disk Stop and Go Extraction Tips (StageTips)

- Empore<sup>™</sup> 47 mm disk for various functionality
  - C18
  - C8
  - Cation exchange
  - Anion exchange
  - SDB-XC
- Self-made tool for customization
- Ease-of-use
- Good recovery
- Reproducible and robust

Rappsilber, J., Mann, M., and Ishihama, Y. "Protocol for Micro-purification, Enrichment, Pre-fractionation and Storage of Peptides for Proteomics using StageTips." *Nature Protocols* **2**, 1896-1906 (2007) Published online: 2 August 2007

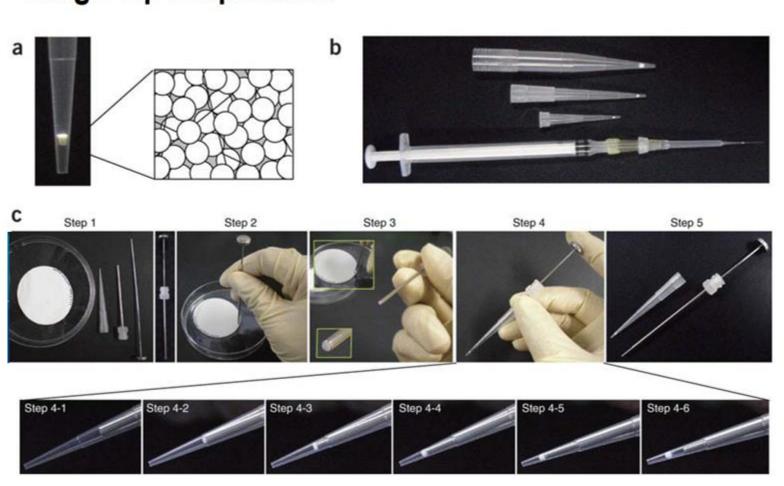


#### StageTips Preparation

- Small disk is stamped out using a blunt-ended syringe needle (cutter)
  - Particles held together for easy handling
  - Size of disk adjusted by using different gauge needle
- Place the cutter inside a pipette tip and release the disk using a correct size needle plunger
- Press the disk gently into place using the plunger
- Remove cutter
- Additional disks can be layered to provide multiple functionality



#### **Stage Tip Preparation**



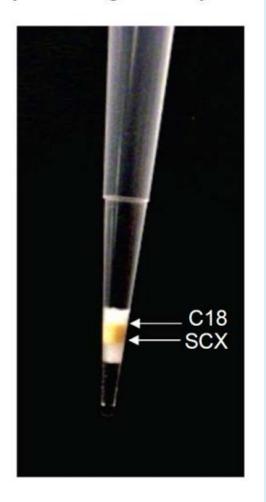


#### Multiple Layers for Additional Separating Ability

- C18 allows for desalting of the protein
- Fractionation done on the cation exchange disk.
- Can be followed up by another desalting step with C18
- Very clean sample for mass spec analysis

Ishihama, Y., Rappsilber, J., and Mann, M., "Modular Stop and Go Extraction Tips with Stacked disks for Parallel and Multidimensional Peptide Fractionation in Proteomics." *J. Proteome Res.*, **2006**, 5 (4), pp 988–994

Publication Date (Web): March 07, 2006 (Technical Note)





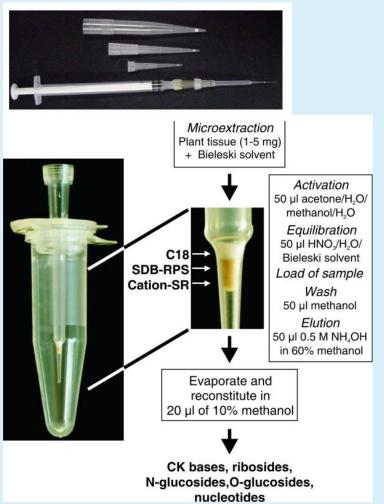
### Why Desalting is Important for LCMS Based Proteomics?

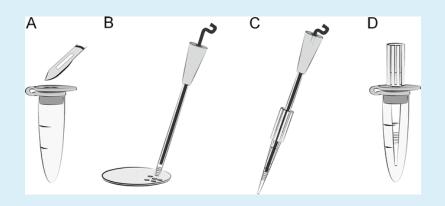
Salt crystals can damage switching valves in LC system, cause instable backpressure and potential sample loss!

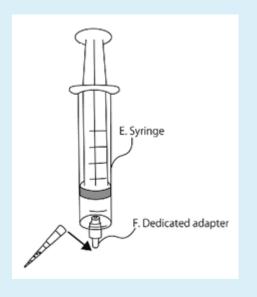




#### Manual Push, NO!









#### Automatable StageTips

Pipette tip adaptors for spinable StageTipping! Hands-free, scalable, high throughput!





# Empore StageTips QA/QC: SDB-RPS StageTips Lot-to-Lot Consistency



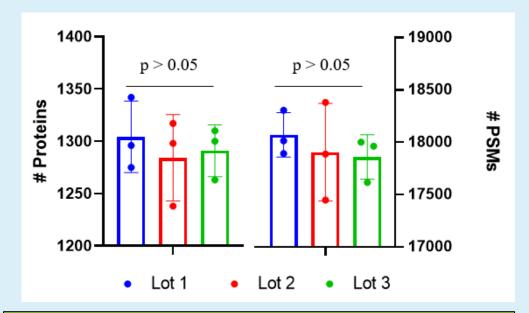


#### **Identification Rate**

	Tip	# Proteins	#PSMs
Lot1 (760179D)	Tip1	1,296	17,893
	Tip2	1,342	18,014
	Tip3	1,275	18,306
Lot2 (760214D)	Tip1	1,298	17,448
	Tip2	1,238	17,888
	Tip3	1,317	18,380
Lot3 (760002C)	Tip1	1,310	18,001
	Tip2	1,263	17,615
	Tip3	1,300	17,963

Cell lysate: BMDM cells. Digestion: STrap method Desalting: SDB tips

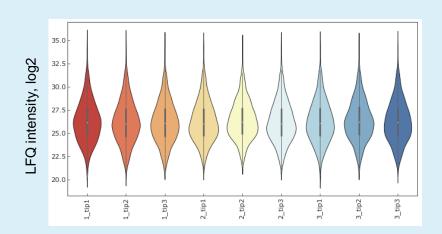
LCMS: Q Exactive (150-min)

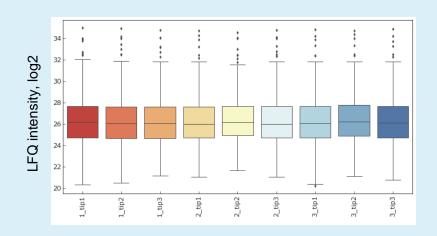


No significant variations were found between different tips and different lots, suggestion excellent reliability and reproducibility of our manufacturing procedure.



#### **Quantitation Performance**

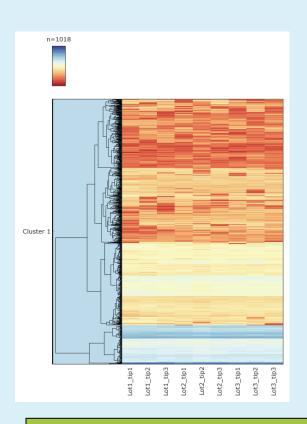


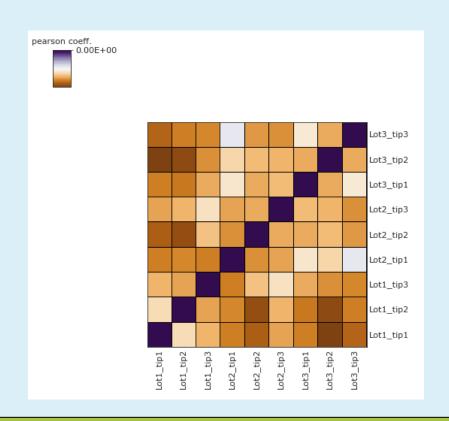


Proteins showed highly similar intensity distribution, suggesting excellent quantatitation performance!



#### **Heatmap and Correlation**



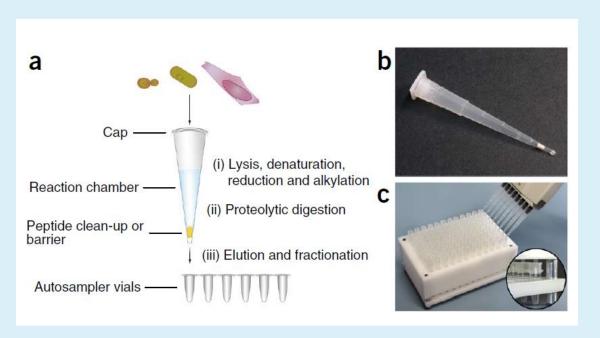


Quantified proteins showed high correlation (Pearson r > 0.93) between different tips and different lots, suggesting again excellent quantitation performance!

CDS Analytical Performance! 20

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#### 2. Applications in Proteomics



SCX (24 h) 94,000 24,000 29,000 19,000 SDB-RPS (12 h) 78,000 (2 × 12 h) 93,000

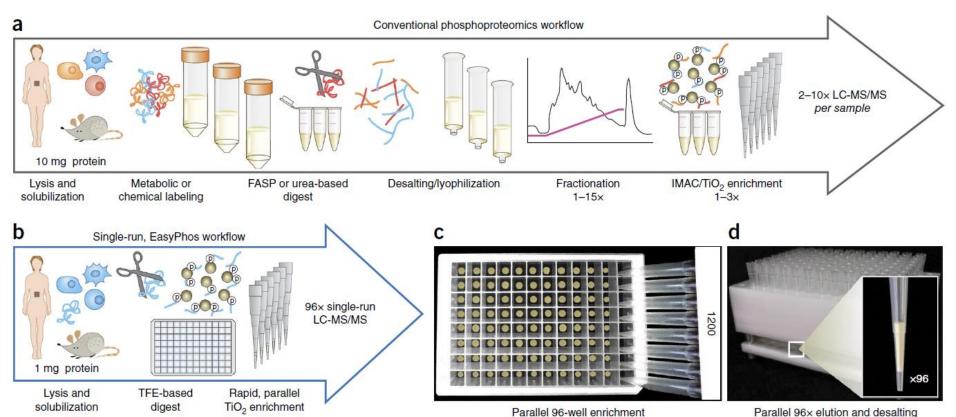
**HeLa Cells** 

Peptide identifications from 20 µg starting material fractionated by SAX, SCX and SDB-RPS StageTips.

Nils A Kulak, Garwin Pichler, Igor Paron, Nagarjuna Nagaraj & Matthias Mann, Nature Method, 2014, 11, 319-326.

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High-throughput phosphoproteomics: HePa 1-6 cells

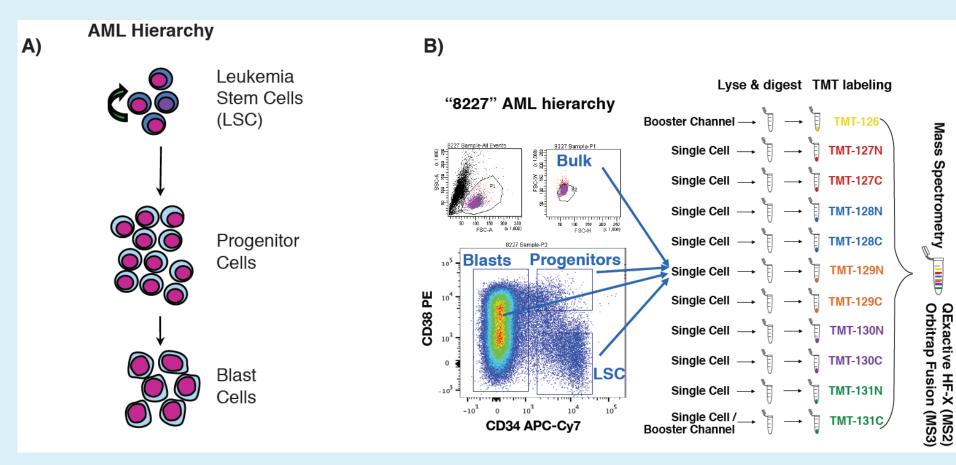
**StageTips V. S. Traditional method:** 

10 times less starting material; 3 times less measurement time; identifying 3 times of the quantified sites.

Sean J Humphrey, S Babak Azimifar & Matthias Mann, Nature Biotech., 2015, 33, 990.

## Analytical A LabTech Company

### Single-cell Proteomics for AML Hierarchy

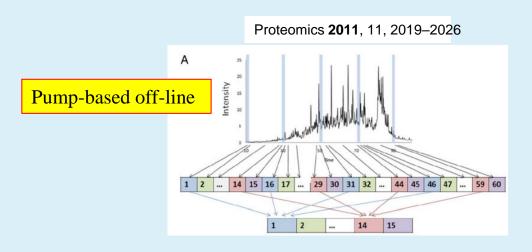


StageTips in sample preparation to help desalting and peptide purification.

### 3. High-pH Fractionation for Global Proteomics



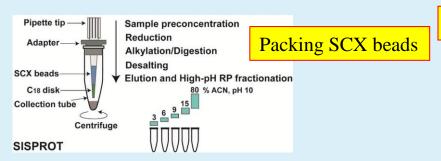
#### Current High pH Fractionation Approaches



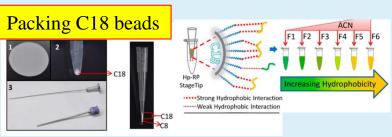
- 1. Pump-based approach (e.g., using off-line HPLC) is sophisticated and expensive.
- Beads-based approaches are unreliable (e.g., need to weight certain amount of beads material and do packing every time).

Our approach, C18 membrane tip based, is *simple, reliable and robust.* 

Anal. Chem., 2016, 88 (9), pp 4864-4871

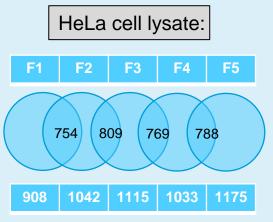


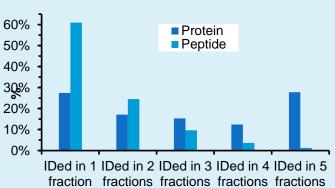
Anal. Chem., 2015, 87 (24), pp 12016-12023



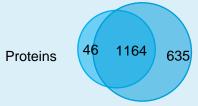


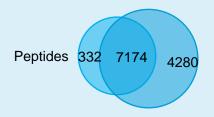
#### Comparison among Fractions

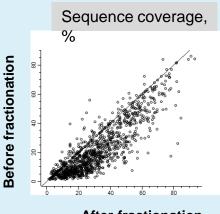






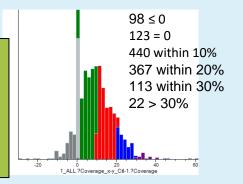






After fractionation

Fractionation significantly increases identification rate on both protein and peptide levels.



# Conclusions of Empore C18 StageTips for High-pH Fractionation



- 1. High pH fractionation increased the number of protein identification by > 50%.
- 2. The sequence coverage of most of the identified proteins increased > 10%.
- 3. At least five or six fractions should be applied (5%, 10%, 15%/25%, 40%, 80% acetonitrile in 10 mM  $NH_4F$ , pH=10).
- 4. C18 StageTip-based high pH fractionation is convenient, easy-to-manipulate and pump-free.
- 5. Current high pH fractionation methods usually apply off-line HPLC pump, or pack C18 resins into tips; C18 StageTip is much simple and cost-effective.
- 6. This method has not been reported for global proteomics before.

#### **Summary**



- 1. Empore StageTips is one of the most important methods for protein desalting and peptide purification in Proteomics research.
- 2. Empore C18 StageTips is a much simpler, costeffective and efficient sample preparation method in high-pH fractionation in proteomics than off-line HPLC pump and packed C18 resins into tips.