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Obsalting of Peptides to Prepare for Mass Spectrometry Analysis

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ABSTRACT

Prior to proteomic analysis, peptide samples are desalted and eluted with freshly prepared 50% acetonitrile, 0.1% trifluoroacetic acid, followed by concentration in a vacuum concentrator. Peptides are then resuspended in freshly prepared 5% acetonitrile, 0.1% formic acid.

Note

- The last step can be conducted at a mass spectrometry facility according to their own preferred methods.
- After desalting, the concentration of the peptide solution can be measured by spectrophotometry.

MATERIALS

Materials:

- 1. X HPLC-grade water Thermo Fisher Scientific Catalog #51140
- 2. X Trifluoro-acetic Acid Thermo Fisher Scientific Catalog #85183
- 3. Acetonitrile mass spectrometry grade Thermo Scientific Catalog #51101
- 4. S Formic Acid Thermo Scientific Catalog #28905

5.

EquipmentPeptide Desalting Spin ColumnsNAMEThermo ScientificBRAND89851SKUhttps://www.thermofisher.com/order/catalog/product/89851LINK

6.

Equipment	
Low Retention Tubes and Tips	NAME
Brand	BRAND
0000000	SKU

Note

■ The maximum volume for spin columns is 300 uL. If columns ever become unpacked, repeat the step that caused this by reloading the flowthrough and spinning at the recommended speed.

Reagents

Table 1: Reagent Preparation

Table Reagent Preparation	
Reagents	Amount
Priming buffer	100% Acetonitrile
Washing buffer	0.1% TFA in HPLC H_2O
Elution buffer	50% solution with 0.1% TFA in HPLC-grade water
Reconstitution buffer	5% solution with 0.1% Formic acid in HPLC-grade water

Column Preparation

4m

1 Column Preparation

Take a Pierce peptide desalting spin column and remove the white tip (do not remove the screw cap of the tube). Place in a 2mL tube and spin column at \mathfrak{g} for

1m

© 00:01:00

1m

1.3 Repeat this step once

2m

Note

• Note that if columns ever become unpacked, repeat that step as the columns will not work properly if unpacked.

Sample Loading

2m

- 2 Sample Loading
- 2.1 Place the spin column in a new low-retention 2 mL tube labeled "flowthrough".

2.2

Load \underline{A} 300 μL of peptide sample into the tube and spin at ζ 5



Note

- You can save the flow-through to ensure it does not contain any unbound peptides and that peptides are binding to the columns.
- **2.3** Based on the total sample volume, if:

 - 2.3b)Less than $\boxed{\text{\em L}}$ 300 μL -> reload the flow-through.
- Spin the sample at g for 00:01:00 . Store "FT" at 8 -80 °C for troubleshooting purpose.

1m

Wash

3m

- 3 Wash Sample
- Place the spin column in a new low-retention 2mL-tube and load $2 300 \,\mu$ L of 0.1% TFA in HPLC-grade H_2O . Centrifuge at g for g 00:01:00 Discard wash flow-through.
- **3.2** Repeat step 3.1 2 more times.

2m

Note

■ Note that if columns ever become unpacked, repeat that step as the columns will not work properly if unpacked.

Sample Elution

1m

- 4 Elute Samples
- 4.1 Place the spin column in a new 2mL low-retention tube labeled with the sample name.
- Load Δ 300 μL of 0.1% TFA, 50% acetonitrile in HPLC-grade H₂O. Spin at

(3000 x)

for 🕙 00:01:00

Note

- Note that if columns ever become unpacked, repeat that step as the columns will not work properly if unpacked.
- **4.3** Transfer the spin column to another 2mL-low retention tube and repeat the step.
- **4.4** Pool the two elution samples from 4.2 and 4.3. These are the desalted peptides.

4.5 Store at 1 -20 °C

Lyophilization and Reconstitution

- 5 Lyophilization
- **5.1** To remove reagents incompatible with mass spectrometry place tubes in SpeedVac™ until completely dry.
- Depending on the size of the peptide pellet, resuspend samples with Δ 20 -75 μL of 0.1% formic acid, 5% acetonitrile in HPLC-grade H_2O .
- **5.3** Vortex until completely resuspended.
- **5.4** Peptide concentration can be measures using a spectrophotometer or using peptide concentration measurement kits such as:

Pierce Quantitative Colorimetric Peptide Assay **Thermo Fisher Scientific Catalog** #23275