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C4 ZipTip Solid Phase Extraction

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1 Works for me



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ABSTRACT

Solid phase extraction for clean-up and concentration of proteins prior to introduction into the mass spectrometer.

PROTOCOL CITATION

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MATERIALS TEXT

C4 ZipTips

KimWipes

P20 and P200 pipettes with tips

LoBind Microcentrifuge tubes

C4 ZipTip Activation Buffer

C4 ZipTip Equilibration/Wash Buffer

C4 ZipTip Elution Buffer

HPLC Buffer A



- 1 Activate Ziptip by pipetting 10 µL of C4 ZipTip Activation Buffer and discarding onto a Kimwipe for a total of 6 times.
- 2 Equilibrate the Ziptip by pipetting 10 μL of C4 ZipTip Equilibration/Wash Buffer and discarding onto a Kimwipe for a total of 6 times.
- Remove C4 Ziptip from p20 pipette and place safely back into tip box to hold. Take a p200 pipette set at 200 μL and add a p200 pipette tip to the end. Take the p200 with pipette tip and then add the C4 ZipTip to the end. Carefully pipette the elution sample up and down for a total of 10 times per elution fraction. Avoid forcing air bubbles through the pipette tip as this will disrupt the resin and introduce oxidation to the target protein.
- 4 Remove C4 Ziptip from p200 pipette and place safely back into tip box to hold. Reattach to the p20 pipette and by pipet 10 μ L of C4 ZipTip Equilibration/Wash Buffer and discard onto a Kimwipe for a total of 10 times.
- 5 Pipette into $5 \mu L$ of C4 ZipTip Elution Buffer that is inside a clean LoBind tube a total of 10 times. Dilute the final volume up to $25 \mu L$ for LC-MS or $80 \mu L$ for I2MS.