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Methanol-Chloroform-Water Precipitation

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Works for me dx.doi.org/10.17504/protocols.io.biknkcve

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ABSTRACT

Methanol-chloroform-water (MCW) precipitation is a rapid method for removing interferring species from protein samples prior to LC-MS analysis. MCW precipitation can effectively remove both ionic and non-ionic detergents and salts. Precipitating less than 1 µg protein is challenging yet possible with careful pipetting.

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KEYWORDS

protein, mass spectrometry, top-down proteomics

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Acetonitrile Optima™ LC/MS Grade Fisher Chemical Fisher

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Resuspension buffer: 94.9% water, 4.9% acetonitrile, 0.2% formic acid

SAFETY WARNINGS

Halogenated solvent waste should be disposed properly as it poses environmental hazards.

- 1 Dilute protein sample to 150 µL with Optima-grade water.
- 2 Precipitate protein at interface
 - 2.1 Add 600 µL Optima-grade methanol and mix well.
 - 2.2 Add 150 µL HPLC-grade chloroform and mix well.
 - 2.3 Centrifuge **20000** x g, 4°C, 00:10:00 and discard top layer while being careful to not disturb interface between layers. Use a gel-loading tip to remove last bit.
- 3 Pellet protein at interface
 - 3.1 Add 600 µL and homogenize the solvent, being careful not to break up the precipitated protein.
 - 3.2 Centrifuge **20000** x g, 4°C and discard supernatant, being careful not to disturb pellet.

- 4 Wash pellet with methanol. (This step can be repeated if a high level of contaminating species is anticipated)
 - 4.1 Add 600 μ L methanol and mix by inverting tube.
 - $\textbf{4.2} \quad \text{Centrifuge } \textcircled{\$20000 x g, 4°C} \text{ and discard supernatant, being careful not to disturb pellet.}$
- 5 Redissolve protein in LC buffer
 - 5.1 Evaporate excess methanol by laying tube on its side for 5 min at room temperature.
 - 5.2 Redissolve precipitated protein in 30 μL Buffer A. Incubate 10 min on ice. If protein is not completely dissolved, sample can be briefly sonicated with a bath sonicator.
 - 5.3 Centrifuge **20000** x g, 4°C and transfer supernatant to LC sample vial.