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BAF_Protocol_008 Metabolomics: Soluble Metabolite Extraction

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Protocol status: Working

We use this protocol and it's working

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metabolomic, soluble metabolite extraction, standard metabolite sample prep, clean sample free of most lipid, extraction, clean sample, lipid, methanol, using chloroform, most lipid

Abstract

This is a standard metabolite sample prep using chloroform/methanol and keeping the soluble (aqueous) phase. This method will produce a clean sample free of most lipids and very hydrophobic molecules. The samples are ready for MS analysis or storage dry at -80C.

Guidelines

Make sure some type of normalization is considered before starting - dry weight, cell number, etc - so equal amounts of material can be injected into the MS. This can be done by counting the same number of cells into each sample tube or weighing the same amount of tissue/stool into each tube or if larger differences you may adjust the extraction volume to account of the differences.



Materials

Microtubes 1.5 mL - SEAL-RITE® 1.5 ML MICROCENTRIFUGE TUBES color: natural, USA Scientific
Reinforced tubes 2 mL- with screw caps and O-rings, Fisherbrand™, White/Opaque, part number 5-340-162.
Stainless steel balls - OMNI International 2.4mm Metal Bead Media 500g SKU 19-640
Pipette tips - Fisherbrand™, yellow, part number: 02-681-151.
Water - Fisher chemical, W64, Optima LC/MS
FA - Fisher chemical, A117-50, Formic Acid, Optima LC/MS
Methanol - Fisher chemical, A456-212, Methanol, Optima LC/MS
Chloroform - Millipore sigma, CX1050P-1, Chloroform, HPLC grade
2 to 20 µL Micropipette - Gilson™ F144056MT
20 to 200 µL Micropipette - Gilson™ F144058MT
100 to 1000 µL Micropipette - Gilson™ F144059MT
VWR Analog Vortex mixer - CAT No: 58816-121
Thermo Scientific™ integrated Speedvac Concentrator CAT No: SPD1030-115
Eppendorf 5415D Digital Centrifuge
Thermo Scientific™ Thermal Mixer with blocks, Block, 24 × 2.0mL microtubes, CAT No: 13687713
Fisherbrand™ bead Mill 24 homogenizer CAT No: 15-340-163
Autosampler vials - SureSTART™ 6ERV11-03PPC Vial 0.3mL CLR PP SNP CON
Vial caps - SureSTART™ 6ARC11ST1OR Cap snap 11mm orange polyethylene

Troubleshooting



Liquid Samples: Urine, Plasma, Cell Media

4h

- 1 To each sample containing 100 μ L add 750 μ L of -20°C cold Chloroform:methanol (2:1) mixture and vortex 1m
- 2 Shake tubes vigorously for 30 min at 4°C in temperature temperature-controlled thermal shaker 30m
- 3 Add 400 μ L of water, shake vigorously, and centrifuge for 10 min at 10000 rpm for phase separation. 11m
- 4 Recover the top aq. methanolic phase as metabolite mixture and transfer to Eppendorf tube. 1m
- 5 Dry the soluble metabolite samples under vacuum for 3-4 h. Store at -80°C until ready for LC-MS(/MS) analysis. 3h
- 6 *Note: save the lower phase if you need to extract lipids. Go for BAF_Protocol_009.
- 7 Before running, reconstitute samples in 100 μ L of 0.1% Formic acid in water containing 100X diluted Metabolomics QReSS heavy labeled standards.
(https://www.isotope.com/userfiles/files/assetLibrary/MET_RSCH_QReSS.pdf) 2m
- 8 Prepare a QC sample with 10 μ L of each sample and transfer 50 μ L of each sample to autosampler vials 5m
- 9 Injection volume of each sample: 10 μ L

Samples that need disruption/lysis: Tissue, Stools, Cell pellets

10m

- 10 Place the sample into reinforced tubes: frozen tissue slice or lyophilized stools. For cell pellets: mix well with 50-100 μ L of water transfer solution and suspended cells to reinforced tubes. 5m
- 11 To each sample add 5 stainless steel balls, 750 μ L of -20°C cold Chloroform:methanol (2:1) mixture 2m



- 12 Disrupt cells/tissues with Fisher Bead Mill 24 (speed: 5m/s, time: 20 sec, number of cycle: 3, dwell/pause between runs: 10 sec)
- 13 Follow steps #2 to #9.

3m