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Objective. LC-MS analysis of non-polar (and polar) metabolites from cells.					
Chemicals and Tools	Vendor	Part#	Hazards/Notes		
LC-MS grade methanol (MeOH)	Fisher	A456-4			
 LC-MS grade water (H₂O) 	Fisher	W6-4			
 HPLC grade chloroform (CHCl₃) 	Sigma	650471	Do NOT use amylene stabilized CHCl3		
• 2.5 mM pre-mixed Heavy Amino Acid (AA) mix (U- ¹³ C, ¹⁵ N)	CIL	MSK-A2-1.2	Add heavy ISTDs to methanol if you are submitting the polar layer for metabolite profiling		
• -80°C Freezer					
• Vials	Eppendorf	022431081	LoBind tubes		
VortexerBenchtop centrifuge					

Untargeted analysis of lipids

Lipids from various classes including ceramides, phospholipids, gangliosides, sphingosines, acylcarnitines, fatty acids and many others are measured using untargeted LC-MS/MS methods. The lipids are separated using reverse-phase chromatography (C18) and the mass to charge ratio (m/z) is measured in both positive and negative ionization modes (separate injections)— typically with mass accuracies ≤5 ppm. Raw data is searched against the Thermo Scientific™ LipidSearch™ Software which contains > 1.5 million lipid ions and their predicted fragment ions.

I. The extraction process will result in a methanol phase-which contains polar metabolites, and a chloroform phase-which contains non-polar metabolites. We can carry out targeted metabolomics on the methanol extract (polar metabolites) but some 'less polar' metabolites may partition into the chloroform phase.

	Procedure.	Examples, Tricks & Comments
1	Collect blood into lithium heparin tube and vortex the sample. Centrifuge the tube for 5-10 minutes at max speed at room temperature. Transfer the supernatant into an Eppendorf tube, snapfreeze on dry ice or liquid nitrogen and store at -80°C till extraction or extract immediately.	Prepare the plasma/serum sample as you see fit. Step 1 is a general guideline. Use the same type of vacutainer tube for the entire experiment.



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2	Prepare the MeOH extraction solution : This solution consists of 100% LC-MS grade methanol. Add the heavy AA mix to a final concentration of 3 μ M if the methanol layer will be retained for metabolite profiling. Example Preparation: 30 mL MeOH + 36 μ L of 2.5 mM heavy AA mix (Refer to Table1). This solution can be stored long-term at -20°C. It is recommended that this solution and the CHCl ₃ are pre-chilled at -20°C. The H ₂ O can be pre-chilled at 4°C.	Deuterated lipid standards like Avanti Splash ® (product #330707, #330709 or d5-DG ISTD Mix I) can be added to the chloroform, prior to extraction, to serve as internal standards for the non-polar phase.
3	 Add 180 μL of the MeOH extraction solution to a 10 μL aliquot of serum and vortex for 20 seconds. Add 390 μL of CHCl₃ to the mixture and vortex for 20 seconds. Chloroform can be substituted for GC/HPLC-grade dichloromethane (DCM). Add 120 μL of water to the mixture and vortex for 10 minutes at 4°C. Centrifugation the samples at 16,000 RCF (or max. speed) at 4°C for 10 minutes to separate the 3 phases (CHCl₃– Precipitate– MeOH). Carefully collect the two layers separately and transfer each layer into a new Eppendorf tube. Avoid the protein interface between the chloroform (bottom) and methanol (top) layers. 	Note. You can save the cell pellet to measure protein concentration for sample normalization or proteomic analysis. Pipette tips and Eppendorf tubes do not have high chemical resistance to chloroform. Avoid polycarbonate-based consumables. Propylene/polyethylene-based tubes can be used short term (≤4°C). All steps must be performed on ice or at 4°C.
	OPTIONAL: Each layer can be divided equally amongst two Eppendorf tubes. One vial can be stored at -80°C, to serve as a back-up, post evaporation .	
4	Dry the samples using nitrogen air or a temperature controlled centrifugal evaporator. Store the dried extracts at -80°C until LC-MS analysis.	Drying the chloroform layer will take ~1 hour. The methanol layer will take ~2-3 hours.
5	Fill out the metabolomics/lipidomics submission form and submit the dried extracts to the PRC. [https://formspolicies.rockefeller.edu/getfile.php?type=Form&file=Proteomics_Metabolomics_submission_form_xlsx]	Required information; • Cell line/ Cell count • List of specific metabolites (or full profiling) • ISTD composition/ concentration • Cell treatment (e.g. labels, inhibitors, etc.)



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Comments.

The heavy amino acid mix (MSK-A2-1.2) is used as an internal standard for the polar phase. Refer to **Table 1** for the composition of the MSK-A2-1.2 product.

Information regarding the mixed heavy lipid standards can be found in Tables 2-4 and here; https://avantilipids.com/product/330709 and https://avantilipids.com/product/lm6001 . You can substitute these with other isotopically labelled standard(s) so long as the extraction buffer does not contain any endogenous metabolites. Note: For lipidomics, it is best to use deuterated internal standards. Deuterated fatty acids can be purchased from Cambridge Isotope Laboratories on a gram scale (<\$300/1 G, e.g. Item #DLM-215-PK, DLM-208-PK, DLM-379-PK).

Samples can be normalized via cell count, protein concentration or DNA concentration. Note that the biological samples (dry extracts) will be treated identically upon submission to the PRC.

If you are treating the samples with reducing/oxidizing agents, drugs or any other compounds that can be extracted during the extraction step, the reagent name and the final concentration (in the dry pellet) needs to be listed in the submission form.



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Table 1. Composition of the Cambridge Isotope Laboratories MSK-A2-1.2 mixture.

Name	Product identifier
WATER UNLABELED	(CAS-No.) 7732-18-5 (EC-No.) 231-791-2
HYDROCHLORIC ACID	(CAS-No.) 7647-01-0 (EC-No.) 231-595-7 (EC Index-No.) 017-002-00-2
L-ALANINE (13C3, 99%; 15N, 99%)	(CAS-No.) 312623-85-1 (EC-No.) 200-273-8 (Unlabeled) (EC Index-No.)
L-LYSINE:2HCL (13C6, 99%; 15N2, 99%)	(CAS-No.) 657-26-1 (Unlabeled) (EC-No.) 211-518-3 (Unlabeled)
L-HISTIDINE:HCL:H2O (<5% D) (13C6, 97-99%; 15N3, 97-99%)	(CAS-No.) 5934-29-2 (Unlabeled)
L-ARGININE:HCL (13C6, 99%; 15N4, 99%)	(CAS-No.) 202468-25-5 (EC-No.) 214-275-1 (Unlabeled)
L-TYROSINE (13C9, 99%; 15N, 99%)	(CAS-No.) 202407-26-9 (EC-No.) 200-460-4 (Unlabeled)
L-PHENYLALANINE (13C9, 99%; 15N, 99%)	(CAS-No.) 63-91-2 (Unlabeled) (EC-No.) 200-568-1 (Unlabeled)
L-METHIONINE (13C5, 99%; 15N, 99%)	(CAS-No.) 63-68-3 (Unlabeled) (EC-No.) 200-562-9 (Unlabeled)
L-GLUTAMIC ACID (13C5, 99%; 15N, 99%)	(CAS-No.) 56-86-0 (Unlabeled) (EC-No.) 200-293-7 (Unlabeled)
L-ASPARTIC ACID (13C4, 99%; 15N, 99%)	(CAS-No.) 202468-27-7 (EC-No.) 200-291-6 (Unlabeled)
L-LEUCINE (13C6, 99%; 15N, 99%)	(CAS-No.) 202406-52-8 (EC-No.) 200-522-0 (Unlabeled)
L-ISOLEUCINE (13C6, 99%; 15N, 99%)	(CAS-No.) 73-32-5 (Unlabeled) (EC-No.) 200-798-2 (Unlabeled)
L-VALINE (13C5, 99%; 15N, 99%)	(CAS-No.) 72-18-4 (Unlabeled) (EC-No.) 200-773-6 (Unlabeled)
L-THREONINE (13C4, 97-99%; 15N, 97-99%)	(CAS-No.) 72-19-5 (Unlabeled) (EC-No.) 200-774-1 (Unlabeled)
L-CYSTINE (13C6, 99%; 15N2, 99%)	(CAS-No.) 1252803-65-8 (EC-No.) 200-296-3 (Unlabeled) (EC Index-No.)
L-PROLINE (13C5, 99%; 15N, 99%)	(CAS-No.) 147-85-3 (Unlabeled) (EC-No.) 205-702-2 (Unlabeled)
L-SERINE (13C3, 99%; 15N, 99%)	(CAS-No.) 202407-34-9 (EC-No.) 200-274-3 (Unlabeled)
GLYCINE (13C2, 99%; 15N, 99%)	(CAS-No.) 211057-02-2 (EC-No.) 200-272-2 (Unlabeled)



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Table 2. Composition of the Avanti SPLASH LipidoMIX[™] product # 330707.

Compound Name	Molecular Weight	Exact Mass	Chemical Formula	Concentration (µg/mL)*
15:0-18:1(d7) PC	753.11	752.61	C ₄₁ H ₇₃ D ₇ NO ₈ P	150.6
15:0-18:1(d7) PE	711.03	710.56	C ₃₈ H ₆₇ D ₇ NO ₈ P	5.3
15:0-18:1(d7) PS (Na Salt)	777.02	776.53	C39H66D7NNaO10P	3.9
15:0-18:1(d7) PG (Na Salt)	764.02	763.54	C ₃₉ H ₆₇ D ₇ NaO ₁₀ P	26.7
15:0-18:1(d7) PI (NH4 Salt)	847.13	846.60	C42H75D7NO13P	8.5
15:0-18:1(d7) PA (Na Salt)	689.94	689.50	C ₃₆ H ₆₁ D ₇ NaO ₈ P	6.9
18:1(d7) Lyso PC	528.72	528.39	C ₂₆ H ₄₅ D ₇ NO ₇ P	23.8
18:1(d7) Lyso PE	486.64	486.35	C23H39D7NO7P	4.9
18:1(d7) Chol Ester	658.16	657.64	C ₄₅ H ₇₁ D ₇ O ₂	329.1
18:1(d7) MAG	363.59	363.34	C21H33D7O4	1.8
15:0-18:1(d7) DAG	587.98	587.55	C ₃₆ H ₆₁ D ₇ O ₅	8.8
15:0-18:1(d7)-15:0 TAG	812.37	811.77	C ₅₁ H ₈₉ D ₇ O ₆	52.8
d18:1-18:1(d9) SM	738.12	737.64	C41H72D9N2O6P	29.6
Cholesterol (d7)	393.71	393.40	C ₂₇ H ₃₉ D ₇ O	98.4

^{*}Concentrations are based on the isotopic purity of each individual compound



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Table 3. Composition of the Avanti SPLASH II LipidoMIX™ product # 330709.

Compound Name	Molecular Weight	Exact Mass	Chemical Formula	Conc. (µg/mL)*	Conc. µM*
15:0-18:1(d7) PC	753.11	752.61	C41H73D7NO8P	158.2	
15:0-18:1(d7) PE	711.03	710.56	C ₃₈ H ₆₇ D ₇ NO ₈ P	5.0	7
15:0-18:1(d7) PS (Na Salt)	777.02	776.53	C ₃₉ H ₆₆ D ₇ NNaO ₁₀ P	7.8	10
15:0-18:1(d7) PI (NH₄ Salt)	847.13	846.60	C42H75D7NO13P	8.5	10
18:1(d7) Lyso PC	528.72	528.39	C ₂₆ H ₄₅ D ₇ NO ₇ P	23.8	45
18:1(d7) Lyso PE	486.64	486.35	C ₂₃ H ₃₉ D ₇ NO ₇ P	0.5	1
18:1(d7) Chol Ester	658.16	657.64	C45H71D7O2	348.8	530
C18(Plasm)-18:1(d9) PC	781.19	780.67	C ₄₄ H ₇₇ D ₉ NO ₇ P	7.8	10
15:0-18:1(d7) DAG	587.98	587.55	C ₃₆ H ₆₁ D ₇ O ₅	11.8	20
15:0-18:1(d7)-15:0 TAG	812.37	811.77	C ₅₁ H ₈₉ D ₇ O ₆	56.9	70
d18:1-18:1(d9) SM	738.12	737.64	C ₄₁ H ₇₂ D ₉ N ₂ O ₆ P	29.5	40
C18(Plasm)-18:1(d9) PE	739.11	738.62	C41H71D9NO7P	0.07	0.1



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Table 4. Composition of the Avanti d5-DG Internal Standard Mix I product # LM6001.

Component Name	LipidMAPS ID	Exact Mass	Concentration by LC/MS
1,3-14:0 DG-d5	LMGL02010309	517.84	4.01 μM (2.07 μg/mL)
1,3-15:0 DG-d5	LMGL02010310	545.51	4.00 μM (2.18 μg/mL)
1,3-16:0 DG-d5	LMGL02010311	573.54	4.09 μM (2.34 μg/mL)
1,3-17:0 DG-d5	LMGL02010312	601.57	4.06 μM (2.45 μg/mL)
1,3-19:0 DG-d5	LMGL02010313	657.63	4.06 μM (2.67 μg/mL)
1,3-20:0 DG-d5	LMGL02010314	685.66	4.06 μM (2.78 μg/mL)
1,3-20:2 DG-d5	LMGL02010315	677.60	4.19 μM (2.84 μg/mL)
1,3-20:4 DG-d5	LMGL02010316	669.54	4.48 μM (3.00 μg/mL)
1,3-20:5 DG-d5	LMGL02010308	665.51	4.47 μM (2.97 μg/mL)