

Supporting Information for Publication

Toward ‘Omic’ Scale Metabolite Profiling: A Dual Separation – Mass Spectrometry Approach for Coverage of Lipids and Central Carbon Metabolism

Julijana Ivanisevic^{†,‡}, Zhengjiang Zhu^{†,‡}, Lars Plate[§], Ralf Tautenhahn[†], Stephen Chen[†], Peter J. O’Brien[#], Caroline H. Johnson[†], Michael A. Marletta[§], Gary J. Patti^{||,*} and Gary Siuzdak^{†,*}

[†]Scripps Center for Metabolomics and Mass Spectrometry, The Scripps Research Institute, 10550 North Torrey Pines Road, La Jolla, CA 92037

[§]Department of Chemistry, The Scripps Research Institute, 10550 North Torrey Pines Road, La Jolla, CA 92037

^{||}Department of Chemistry, Washington University, One Brookings Drive, St. Louis, MO 63130

[#]Pfizer Worldwide Research and Development, La Jolla Laboratories, San Diego, CA 92121

AUTHOR INFORMATION

Corresponding Authors

E-mail: gipattij@wustl.edu, cell: (314) 604-6616, siuzdak@scripps.edu, phone: (858) 784-9415

Author Contributions

[‡]These authors contributed equally

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Sample preparation. For untargeted analysis of the bacterial metabolic response to nitric oxide, *Shewanella oneidensis* MR-1 samples were cultured in MM mineral medium, supplemented with 0.1 % casamino acids, 50 mM sodium fumarate and 50 mM sodium D/L-lactate.¹⁻² Replicate batch cultures were inoculated at a 1:100 dilution from an overnight culture in LB medium, and were grown anaerobically in sealed serum vials at 30 °C. When cultures reached mid-log phase (OD₆₀₀ between 0.2 – 0.3), each sample was split into two separate Hungate tubes. One half was treated with the slow-release nitric oxide donor PAPA NONOate (10 µM, Cayman Chemicals). Anaerobic NONOate stocks were freshly prepared in 10 mM NaOH inside an anaerobic glovebox (MBraun). The second half of each sample was treated with 0.5 mM of the nitric oxide scavenger 2-Phenyl-4,4,5,5-tetramethylimidazoline-3-oxide-1-oxyl (PTIO, TCI America). The cultures were incubated at 30 °C for an additional 60 min, and subsequently chilled on ice. The supernatant was removed by centrifugation at 2,000 x g for 20 min at 4 °C. The cell pellets were rinsed once with ice-cold 0.6 % NaCl and then flash-frozen in liquid nitrogen. Pellets were stored at -80 °C until extraction of metabolites.

For the evaluation of comprehensive, single extraction – dual separation workflow, the following three types of samples were used: standard human plasma samples, *Escherichia coli* cell pellets and one human cancer (Raji Burkitt's) cell line.

The *E. coli* strain W3110 was cultured in three batch culture replicates in complete LB medium from a single overnight culture. The cells were grown at 37 °C, and 15 mL aliquots were harvested from each culture at mid-exponential growth (OD₆₀₀ nm = 0.2 - 0.3). The supernatant was removed by centrifugation for 5 min at 3000 rpm and 4 °C and the cells were rinsed once with 1.5 mL of 0.6 % NaCl solution to wash off the media. Cell pellets were stored at -80 °C until metabolite extraction.

Raji Burkitt's lymphoma cells (CCL-86; ATCC, Manassas, VA, USA) were propagated at a density of 150,000 to 2.0 million cells per mL in complete growth medium consisting of RPMI-1640 medium (#A10491-01; Invitrogen, Carlsbad, CA, USA) supplemented with 10% heat-inactivated Fetal Bovine Serum (PAA #A15-704; GE Healthcare Bio-Sciences, Piscataway, NJ, USA), 100 I.U./ml penicillin, 100 µg/ml streptomycin, and 2mM L-Glutamine (Invitrogen #15070-063) at 37 °C in a 5 % CO₂ incubator. Cell count and viability was monitored daily using the Guava Viacount[®] assay (#4000-0040; Millipore, Billerica, MA, USA). The day before harvest, cells were placed in fresh growth medium, and were greater than 93% viable at the time of sample collection. Samples within each experiment were collected from a common batch of cells as follows: 50 million cells were collected in 50 mL polypropylene centrifuge tubes (#352070; Becton Dickinson, Franklin Lakes, NJ), and immediately centrifuged for 3 minutes, at 450 x g and 4 °C. Supernatants were immediately decanted by inverting the tubes, and cell pellets were gently but rapidly re-suspended by pipetting in 50 mL of ice cold, cell culture-grade Dulbecco's phosphate buffered saline (PBS, #14190; Invitrogen). After centrifugation as above, cell pellets were re-suspended in 25 mL ice cold PBS and 5 mL aliquots (~ 10 million cells per replicate) were immediately dispensed into 15 mL polypropylene tubes (#352097, Becton Dickinson), centrifuged and immediately decanted as above. Residual PBS from this final wash was removed by gently tapping each tube on paper towels to reduce residual PBS without disturbing cell pellets. Samples were snap-frozen in a dry ice-ethanol bath, and stored at -80 °C until extraction.

Table S1. List of metabolites in hydrophilic standard mixture.

NAME	EXACT MASS	[M+H] ⁺	[M-H] ⁻	FORMULA
L-Glutamic Acid	147.0532	148.0602	146.0454	C5H9NO4
D-Aspartic acid	133.0375	134.0445	132.0297	C4H7NO4
D-Ala-D-Ala	160.0848	161.0918	159.0770	C6H12N2O3
L-Arginine	174.1117	175.1187	173.1039	C6H14N4O2
DL-Homoserine	119.0582	120.0652	118.0504	C4H9NO3
Phenylalanine	165.0790	166.0860	164.0712	C9H11NO2
S-(5'-Adenosyl)-L-Homocysteine	384.1216	385.1286	383.1138	C14H20N6O5S
L-Tryptophan	204.0899	205.0969	203.0821	C11H12N2O2
Taurine	125.0147	126.0217	124.0069	C2H7NO3S
Spermine	202.2157	203.2227	201.2079	C10H26N4
Melatonin	232.1212	233.1282	231.1134	C13H16N2O2
Adenine	135.0545	136.0615	134.0467	C5H5N5
Adenosine	267.0967	268.1037	266.0889	C10H13N5O4
Inosine	268.0808	269.0878	267.0730	C10H12N4O5
Guanosine	283.0917	284.0987	282.0839	C10H13N5O5
Thymidine	242.0903	243.0973	241.0825	C10H14N2O5
Deoxyuridine	228.0746	229.0816	227.0668	C9H12N2O5
5'-CMP	323.0519	324.0589	322.0441	C9H14N3O8P
ATP	506.9958	508.0028	505.9880	C10H16N5O13P3
AMP	347.0631	348.0701	346.0553	C10H14N5O7P
UMP	324.0359	325.0429	323.0281	C9H13N2O9P
D-Erythrose-4-phosphate	200.0086	201.0156	199.0008	C4H9O7P
D-Ribose 5-phosphate	230.0192	231.0262	229.0114	C5H11O8P
D-glucose-6-phosphate	260.0297	261.0367	259.0219	C6H13O9P
D-Fructose-6-Phosphate	260.0297	261.0367	259.0219	C6H13O9P
Fructose-1,6-bisphosphate	339.9961	341.0031	338.9883	C6H14O12P2
Phosphoenolpyruvate	167.9824	168.9894	166.9746	C3H5O6P
UDP-D-glucose	566.0550	567.0620	565.0472	C15H24N2O17P2
D-Ribulose-5-phosphate	230.0192	231.0262	229.0114	C5H11O8P
Nicotinuric acid	180.0535	181.0605	179.0457	C8H8N2O3
2-Ethyl-2-Hydroxybutyric acid	132.0786	133.0856	131.0708	C6H12O3
Citric Acid	192.0270	193.0340	191.0192	C6H8O7
Retinoic acid	300.2089	301.2159	299.2011	C20H28O2
Succinic Acid	118.0266	119.0336	117.0188	C4H6O4
Ascorbic acid	176.0321	177.0391	175.0243	C6H8O6
Nicotinic acid	123.0320	124.0390	122.0242	C6H5NO2
Aconitic acid	174.0164	175.0234	173.0086	C6H6O6
Oxaloacetic acid	132.0059	133.0129	130.9981	C4H4O5
Malic acid	134.0215	135.0285	133.0137	C4H6O5
Lactic acid	90.0317	91.0387	89.0239	C3H6O3
Acetyl-CoA	809.1257	810.1327	808.1179	C23H38N7O17P3S
Malonyl-CoA	853.1156	854.1226	852.1078	C24H38N7O19P3S
NAD ⁺	663.1091	664.1161	662.1013	C21H27N7O14P2
NADPH	743.0755	744.0825	742.0677	C21H28N7O17P3
Glutathione oxidized	612.1520	613.1590	611.1442	C20H32N6O12S2
Alpha ketoglutarate	146.0215	147.0285	145.0137	C5H6O5
FAD	785.1571	786.1641	784.1493	C27H33N9O15P2
R-(-)-Mandelic acid	152.0473	153.0543	151.0395	C8H8O3
PGA1	336.2301	337.2371	335.2223	C20H32O4
Palmitoyl dopamine	391.3086	392.3156	390.3008	C24H41NO3

Table S2. List of metabolites in hydrophobic standard mixture.

NAME	EXACT MASS	[M+H] ⁺	[M-H] ⁻	FORMULA
Creatinine	113.0589118	114.0662118	112.0516118	C4H7N3O
Caffeic Acid	180.04226	181.04956	179.03496	C9H8O4
Caffeine	194.080376	195.087676	193.073076	C8H10N4O2
1-Tetradecanol	214.2297	215.237	213.2224	C14H30O
Acetyl-DL-Glucosamine	221.089933	222.097233	220.082633	C8H15NO6
Melatonin	232.1211716	233.1284716	231.1138716	C13H16N2O2
Alizarin	240.0422587	241.0495587	239.0349587	C14H8O4
Palmitic Acid	256.2402172	257.2475172	255.2329172	C16H32O2
1-Octadecanol	270.2923	271.2996	269.285	C18H38O
Estradiole	272.17763	273.18493	271.17033	C18H24O2
Linoleic Acid	280.2402	281.2475	279.2329	C18H32O2
Oleic Acid	282.25588	283.26318	281.24858	C18H34O2
Stearamide	283.2874992	284.2947992	282.2801992	C18H37NO
Stearic Acid	284.2715	285.2788	283.2642	C18H36O2
trans-Retinol(Vitamin A)	286.229653	287.236953	285.222353	C20H30O
B-Octylglucoside	292.1885788	293.1958788	291.1812788	C14H28O6
Sphingosine	299.282429	300.289729	298.275129	C18H37NO2
Retinoic acid	300.2089188	301.2162188	299.2016188	C20H28O2
Arachidonic Acid	304.24023	305.24753	303.23293	C20H32O2
Progesterone	314.2245802	315.2318802	313.2172802	C21H30O2
Decanoyl-DL-Carnitine	315.241	316.2483	314.2337	C17H33NO4
Docosahexanoic Acid	328.2402	329.2475	327.2329	C22H32O2
Eicosapentaenoic Acid	334.2144	335.2217	333.2071	C20H30O4
PGA1	336.23006	337.23736	335.22276	C20H32O4
Erucamide	337.334464	338.341764	336.327164	C22H43NO
Erucic acid (13(Z)-Docosenoic Acid)	338.3184632	339.3257632	337.3111632	C22H42O2
Lauroylcarnitine	343.2722	344.2795	342.2649	C19H37NO4
Rosmarinic Acid	360.0845136	361.0918136	359.0772136	C18H16O8
Hydrocortisone	362.209313	363.216613	361.202013	C21H30O5
Misoprostol	368.256275	369.263575	367.248975	C21H36O5
7-dehydrocholesterol	384.3391974	385.3464974	383.3318974	C27H44O
Cholesterol	386.3548661	387.3621661	385.3475661	C27H46O
Palmitoyl Dopamine	391.3086276	392.3159276	390.3013276	C24H41NO3
C-6 Ceramide	397.3555944	398.3628944	396.3482944	C24H47NO3
Palmitoyl-DL-Carinitine	399.3349	400.3422	398.3276	C23H45NO4
Delta-Tocopherol	402.34978	403.35708	401.34248	C27H46O2
Cholic Acid	408.2876	409.2949	407.2803	C24H40O5
Squalene	410.39123	411.39853	409.38393	C30H50
alpha-Tocopherol	430.3810808	431.3883808	429.3737808	C29H50O2
Folic Acid	441.1397	442.147	440.1324	C19H19N7O6
Geranylgeranyl pyrophosphate	450.1936166	451.2009166	449.1863166	C20H36O7P2
PC (16:0/0:0)	495.332492	496.339792	494.325192	C24H50NO7P
N-Oleoyl-D-Erythro-Sphingosine (C18:1 Ceramide)	563.5277164	564.5350164	562.5204164	C36H69NO3
Cholesteryl oleate	650.6002	651.6075	649.5929	C45H78O2
PS (16:0/16:0)	735.505	736.5123	734.4977	C38H74NO10P
Palmitoyl Thio-PC	749.539282	750.546582	748.531982	C40H80NO7PS
L-Thyroxine	776.6867126	777.6940126	775.6794126	C15H11I4NO4
Coenzyme Q10	862.6839114	863.6912114	861.6766114	C59H90O4
Phosphatidylinositol 3-phosphate, dipalmitoyl	890.492134	891.499434	889.484834	C41H80O16P2
Decanoyl-CoA	921.2509	922.2582	920.2436	C31H54N7O17P3S

Table S3. List of tested mobile phase conditions.

HILIC ESI±MS		
Additives in A = 95% H₂O	Additives in B = 95% ACN	pH
20 mM NH ₄ Ac + 0.1% FA	/	4-4.5
20 mM NH ₄ Ac	/	6.5-7
20 mM NH ₄ OH / 20 mM NH ₄ Ac	/	9.8
50 mM NH ₄ OH / 50 mM NH ₄ Ac	/	9.8
10 mM NH ₄ OH / 10 mM NH ₄ Ac	/	9.8
5 mM NH ₄ OH / 5 mM NH ₄ Ac	/	9.8
HILIC ESI-MS		
Additives in A = 95% H₂O	Additives in B = 95% ACN	pH
10 mM NH ₄ OH / 10 mM NH ₄ HCOO	/	9.6
10 mM NH ₄ OH / 10 mM (NH ₄) ₂ CO ₃	/	9.6
10 mM NH ₄ OH / 10 mM NH ₄ Ac + 0.1m MNH ₄ F	0.1m MNH ₄ F	9.7
10 mM NH ₄ OH / 9 mM NH ₄ Ac + 1 mM NH ₄ F	/	9.5
5 mM NH ₄ OH / 5 mM NH ₄ Ac	5 mM NH ₄ OH / 5 mM NH ₄ Ac	9.7
RPLC ESI-MS		
Additives in A = 100% H₂O	Additives in B = 100% ACN	
5 mM NH ₄ Ac	/	7.8
1 mM NH ₄ F	/	7.5
0.5 mM NH ₄ F	/	7.5
0.1 mM NH ₄ F	/	7.9
0.5 mM NH ₄ F	0.5 mM NH ₄ F	8.5
0.1 mM NH ₄ F	0.1 mM NH ₄ F	8.6

Further optimization of HILIC/MS mobile phase. To further examine the effect of salt additives, ammonium acetate was replaced by ammonium formate and ammonium carbonate in the same concentration mixture with ammonium hydroxide at high pH (Table S3). Both ammonium formate and ammonium carbonate buffers resulted in longer retention times, deterioration of peak shape, and signal suppression on the MS detection in comparison to ammonium acetate, thus giving significantly lower performance scores (Figure S1).

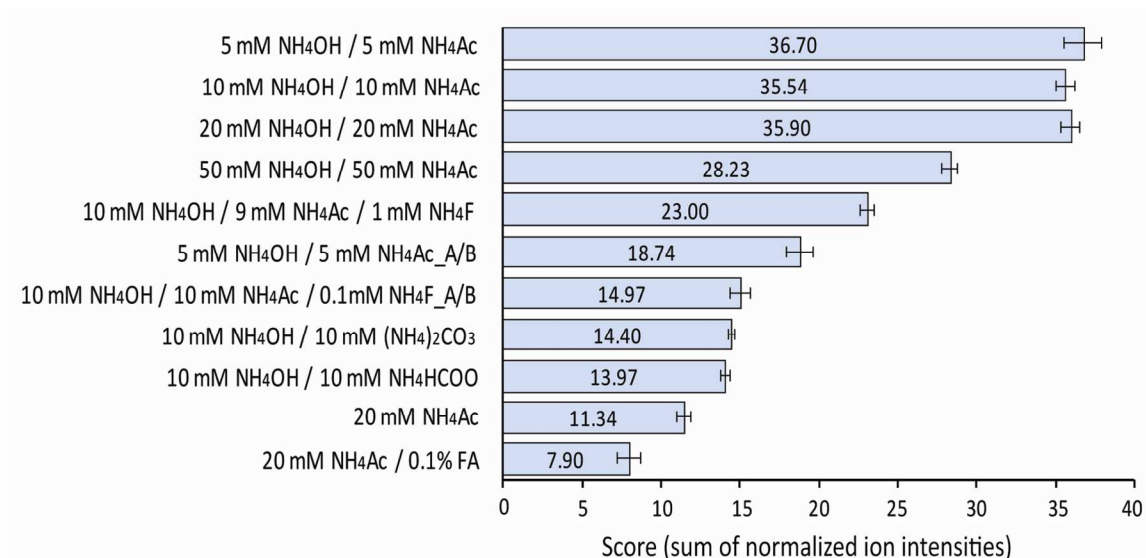


Figure S1. Further optimization of analysis conditions in HILIC ESI negative mode. Each tested analysis condition (Table S3) is presented by its overall score, calculated as a sum of normalized intensities of 50 metabolites from the hydrophilic standard mixture (Table S1).

Ammonium fluoride (NH₄F), which has previously been shown to promote ionization efficiency in negative mode in RPLC,³ was also tested in HILIC ESI negative mode. Fluoride ions, which have stronger basicity than acetate ions in gas phase, effectively promote analyte deprotonation and enhance ionization efficiency³. In RPLC/MS, ammonium fluoride significantly increased the intensity of most metabolites by as much as five times compared to

ammonium acetate. As seen in Figure S2, the optimal concentration of ammonium fluoride additive in RPLC/MS was 0.5 mM. However, the addition of fluoride to the mobile phase in HILIC/MS had deleterious effects for polar metabolite ionization (Figure S1). The overall performance score decreased upon addition of NH_4F from 35.5 (10 mM NH_4Ac and 10 mM NH_4OH) to 23.0 (9 mM NH_4Ac + 1 mM NH_4F and 10 mM NH_4OH). The observed sensitivity decrease may be due to competition of ammonium fluoride with ammonium acetate which may interfere with the ionization process thus decreasing the MS signal intensity.

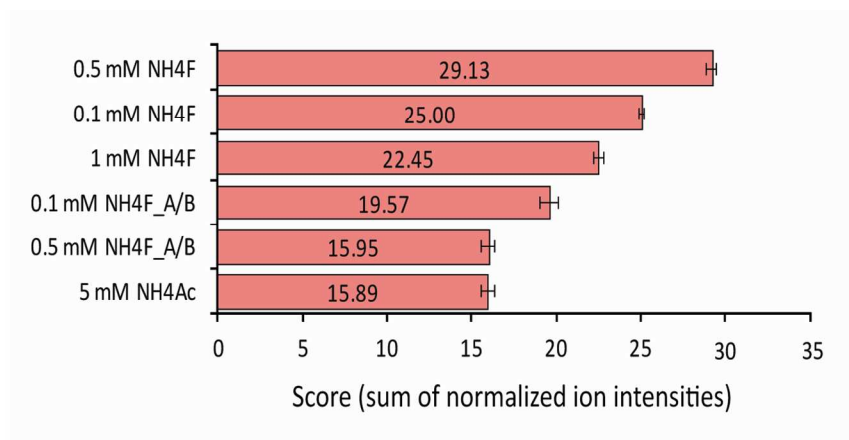


Figure S2. Optimization of analysis conditions in RPLC ESI negative mode. Each tested analysis condition is presented by its overall score, calculated as a sum of normalized intensities of 50 metabolites from the hydrophobic standard mixture (Table S2).

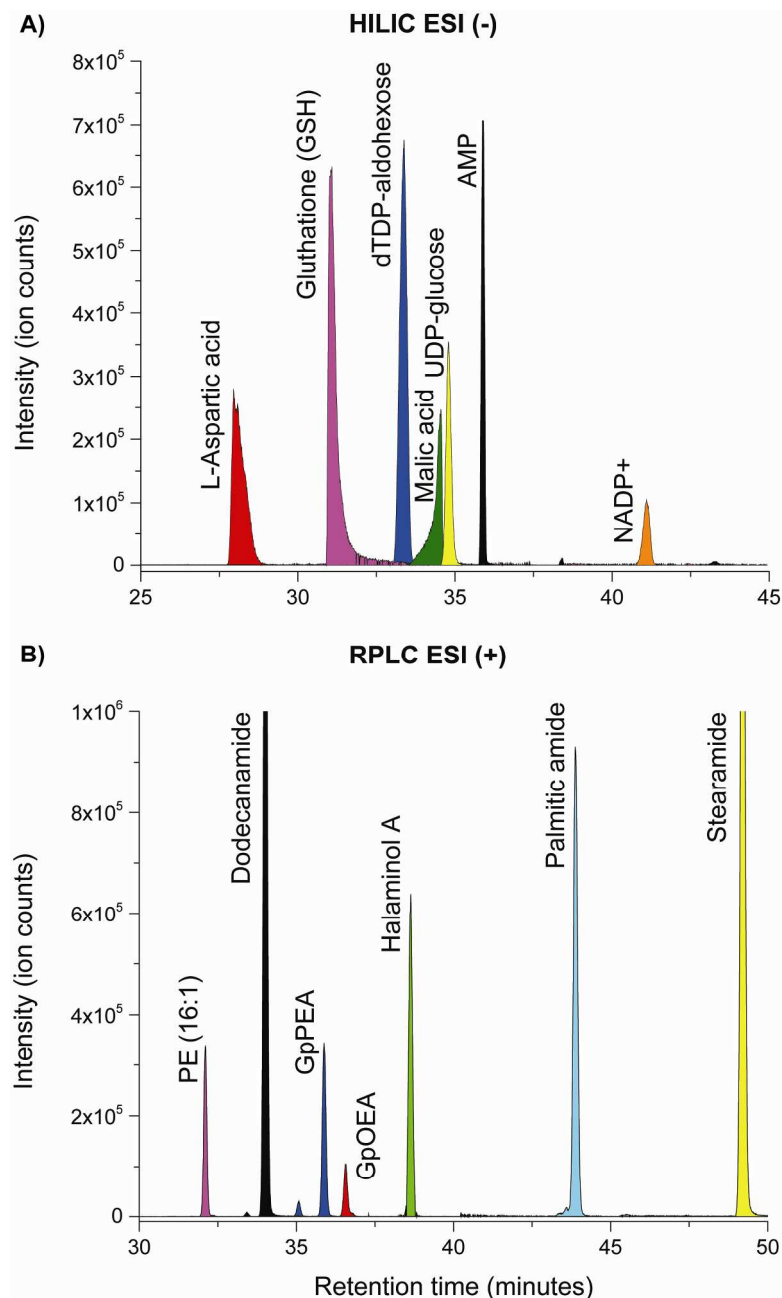


Figure S3. Extracted ion chromatograms from HILIC ESI negative and RPLC ESI positive mode profiling of *E. coli* extract. Indicated compounds have been putatively identified using METLIN database ($\Delta m/z < 5$ ppm and/or MS-MS spectra matching) and retention time. GSH = Glutathione reduced, dTDP-aldohexose = Deoxythymidine diphosphate aldohexose, UDP-glucose = Uridine diphosphate glucose, AMP = Adenosine monophosphate, NADP⁺ = Nicotinamide Adenine Dinucleotide Phosphate, PE = Phosphoethanolamine, GpPEA = Glycerophospho-N-Palmitoyl Ethanolamine, GpOEA = Glycerophospho-N-Oleoyl Ethanolamine

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