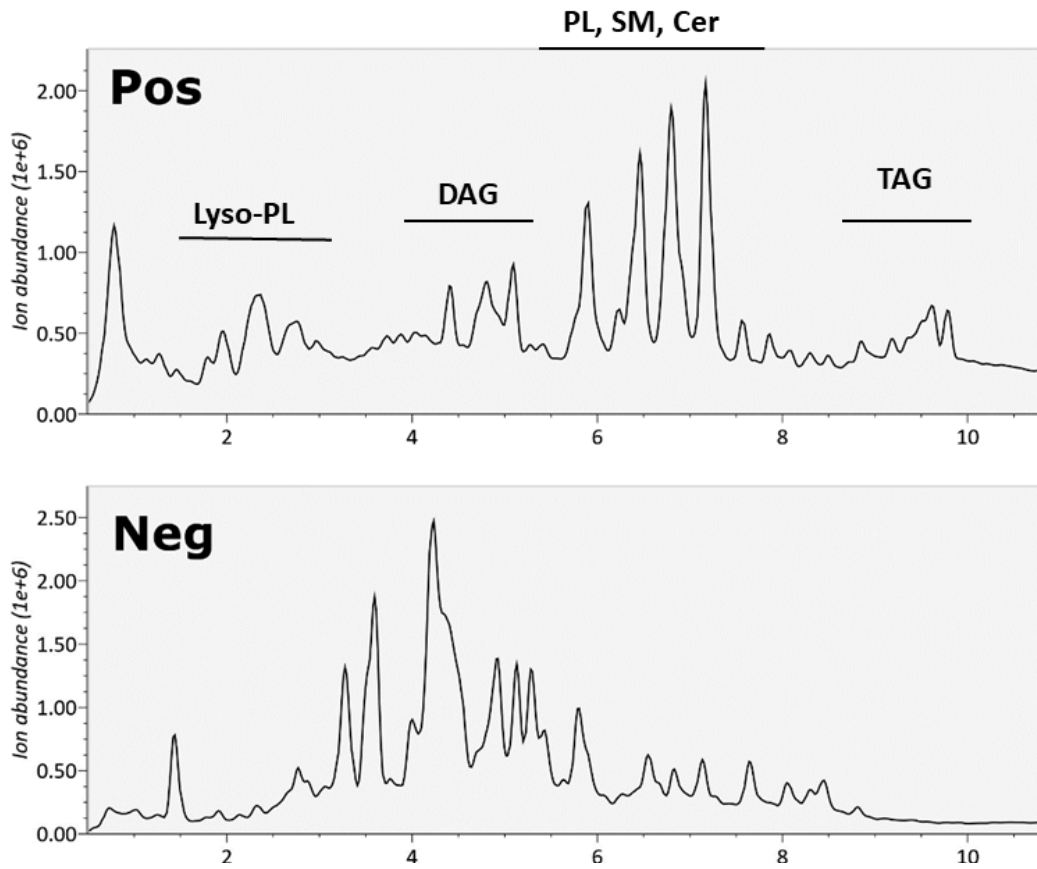
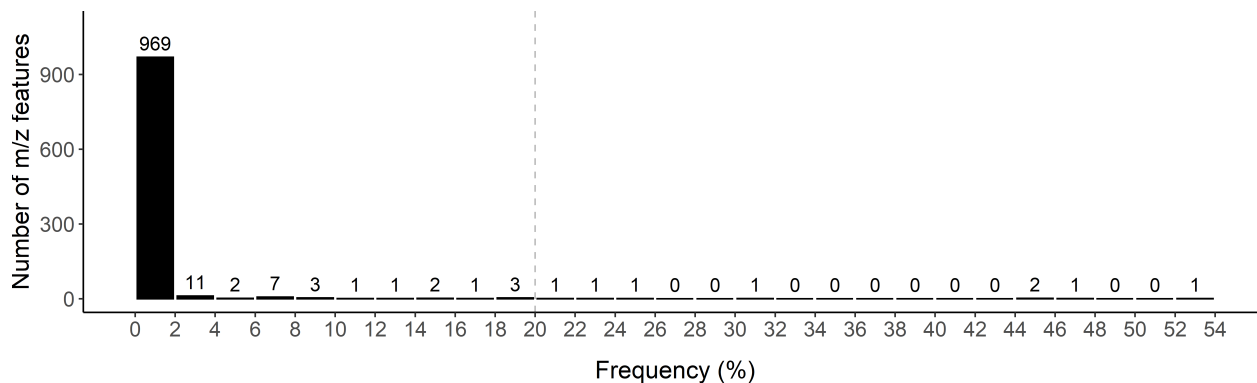


# Supplementary Material

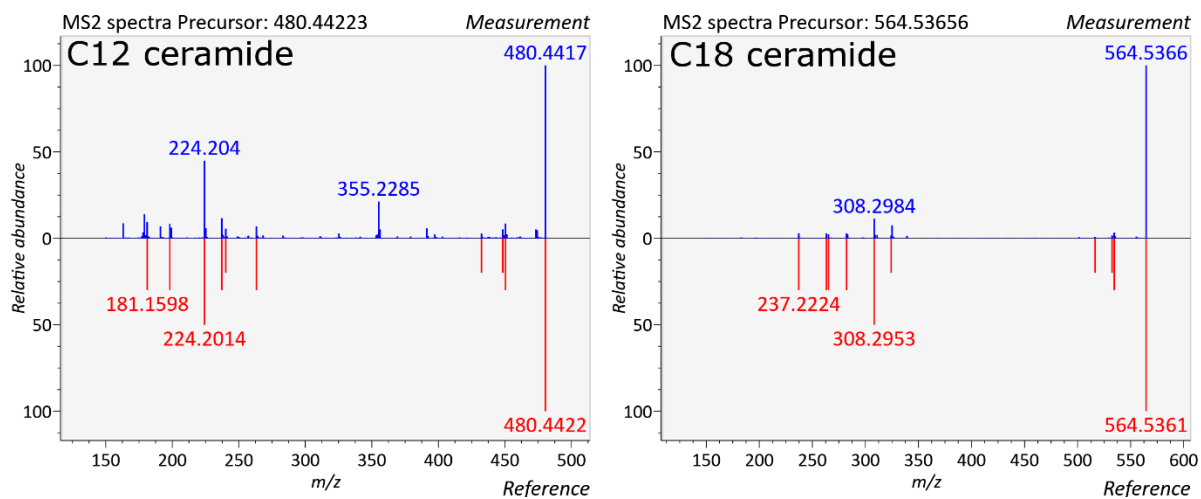
## Cerebrospinal Fluid C18 Ceramide Associates with Markers of Alzheimer's Disease and Inflammation at the Pre- and Early Stages of Dementia



**Supplementary Figure 1.** Total ion chromatographs from CSF extracts, both in negative and positive mode. Labels show roughly the distribution of major lipid classes according to retention time. PL, phospholipids; SM, sphingomyelins; Cer, ceramides; DAG, diacylglycerols; TAG, triacylglycerols; Lyso-PL, lysophospholipids.



**Supplementary Figure 2.** Distribution of m/z features significantly distinguishing between CSF profiles (AD and non-AD), in both discovery and validation sets, after 1000 bootstrap replications. Vast majority (969 of 1008 features) did only distinguish between the profiles in 0-2% of the replications (870 in less than 1% and 99 in between 1-2%). Of the remaining features, 31 distinguished between the profiles in 2-20% of the replications and 8 in 20% of the replications or higher. The vertical dotted line presents the cut-off point for selection of features.



**Supplementary Figure 3.** Negative mode MS/MS (MS2) spectra of C12 and C18 ceramide standards as measured (Blue) compared to a spectral reference provided by LipidBlast library (Red). C12 ceramide was added to all samples as an internal standard, C18 ceramide was run separately.

**Supplementary Table 1a.** LipidBlast annotations of all measured m/z features in positive ionization mode

Average RT (min)	Feature/peak (m/z)	Annotated lipid species
4.12	330.2805	ACar 11:0; [M] <sup>+</sup>
3.39	328.2675	ACar 11:1; [M] <sup>+</sup>
1.72	356.2661	ACar 13:1; [M] <sup>+</sup>
4.33	384.3145	ACar 15:1; [M] <sup>+</sup>
1.91	382.3138	ACar 15:2; [M] <sup>+</sup>
8.51	412.3458	ACar 17:1; [M] <sup>+</sup>
5.46	410.3392	ACar 17:2; [M] <sup>+</sup>
3.73	408.3291	ACar 17:3; [M] <sup>+</sup>
6.81	454.3737	ACar 20:1; [M] <sup>+</sup>
5.53	444.2991	ACar 20:6; [M] <sup>+</sup>
6.56	470.422	ACar 21:0; [M] <sup>+</sup>
4.46	462.3705	ACar 21:4; [M] <sup>+</sup>
3.49	482.4069	ACar 22:1; [M] <sup>+</sup>
3.16	472.3647	ACar 22:6; [M] <sup>+</sup>
3.81	496.4225	ACar 23:1; [M] <sup>+</sup>
2.43	488.3601	ACar 23:5; [M] <sup>+</sup>
2.88	502.3761	ACar 24:5; [M] <sup>+</sup>
3.17	500.3621	ACar 24:6; [M] <sup>+</sup>
3.1	516.3904	ACar 25:5; [M] <sup>+</sup>
3.86	538.4984	ACar 26:1; [M] <sup>+</sup>
6.7	532.4372	ACar 26:4; [M] <sup>+</sup>
2.63	300.2366	ACar 9:1; [M] <sup>+</sup>
5.68	670.4631	BMP 27:0; BMP 5:0-22:0; [M+NH <sub>4</sub> ] <sup>+</sup>
2.21	708.4869	BMP 30:2; BMP 14:1-16:1; [M+NH <sub>4</sub> ] <sup>+</sup>
2.88	736.5172	BMP 32:2; BMP 13:1-19:1; [M+NH <sub>4</sub> ] <sup>+</sup>
7.69	886.6542	BMP 43:4; BMP 23:0-20:4; [M+NH <sub>4</sub> ] <sup>+</sup>
2.25	664.613	CE 18:3; [M+NH <sub>4</sub> ] <sup>+</sup>
6.01	468.4431	Cer-NDS d29:1; Cer-NDS d15:0/14:1; [M+H] <sup>+</sup>
7.82	594.5826	Cer-NDS d38:1; Cer-NDS d22:0/16:1; [M+H] <sup>+</sup>
4.6	606.5599	Cer-NDS d39:2; Cer-NDS d19:0/20:2; [M+H] <sup>+</sup>
2.73	634.6001	Cer-NDS d41:2; Cer-NDS d21:0/20:2; [M+H] <sup>+</sup>
4.05	424.3601	Cer-NS d26:2; Cer-NS d14:1/12:1; [M+H] <sup>+</sup>
6.01	482.4583	Cer-NS d30:1; Cer-NS d18:1/12:0; [M+H] <sup>+</sup>
4.65	476.3901	Cer-NS d30:4; Cer-NS d14:3/16:1; [M+H] <sup>+</sup>
5.03	492.4216	Cer-NS d31:3; Cer-NS d15:1/16:2; [M+H] <sup>+</sup>
8.19	506.4733	Cer-NS d32:3; Cer-NS d14:3/18:0; [M+H] <sup>+</sup>
3.87	506.4607	Cer-NS d32:3; Cer-NS d17:3/15:0; [M+H] <sup>+</sup>
5.93	504.4403	Cer-NS d32:4; Cer-NS d17:3/15:1; [M+H] <sup>+</sup>
5.41	544.4728	Cer-NS d35:5; Cer-NS d15:3/20:2; [M+H] <sup>+</sup>
7.47	566.5524	Cer-NS d36:1; Cer-NS d20:1/16:0; [M+H] <sup>+</sup>
2.33	576.5523	Cer-NS d37:3; Cer-NS d21:1/16:2; [M+H] <sup>+</sup>
5.96	572.5032	Cer-NS d37:5; Cer-NS d15:3/22:2; [M+H] <sup>+</sup>
2.77	590.5693	Cer-NS d38:3; Cer-NS d25:3/13:0; [M+H] <sup>+</sup>
2.29	620.5825	Cer-NS d40:2; Cer-NS d21:1/19:1; [M+H] <sup>+</sup>
7.82	616.5629	Cer-NS d40:4; Cer-NS d20:3/20:1; [M+H] <sup>+</sup>
8.56	650.6429	Cer-NS d42:1; Cer-NS d19:1/23:0; [M+H] <sup>+</sup>
6.83	760.7443	Cer-NS d50:2; Cer-NS d15:2/35:0; [M+H] <sup>+</sup>
3.58	292.2317	DAG 11:0; DAG 5:0-6:0; [M+NH <sub>4</sub> ] <sup>+</sup>
2.07	334.2782	DAG 14:0; DAG 6:0-8:0; [M+NH <sub>4</sub> ] <sup>+</sup>
2.14	348.2934	DAG 15:0; DAG 5:0-10:0; [M+NH <sub>4</sub> ] <sup>+</sup>
1.61	362.2707	DAG 16:0; DAG 6:0-10:0; [M+NH <sub>4</sub> ] <sup>+</sup>
2.69	376.3246	DAG 17:0; DAG 7:0-10:0; [M+NH <sub>4</sub> ] <sup>+</sup>
1.93	376.2858	DAG 17:0; DAG 8:0-9:0; [M+NH <sub>4</sub> ] <sup>+</sup>

4.69	400.2915	DAG 19:2; DAG 2:0-17:2; [M+NH4] <sup>+</sup>
1.91	400.3239	DAG 19:2; DAG 3:0-16:2; [M+NH4] <sup>+</sup>
1.62	396.2929	DAG 19:4; DAG 3:0-16:4; [M+NH4] <sup>+</sup>
1.89	418.3344	DAG 20:0; DAG 5:0-15:0; [M+NH4] <sup>+</sup>
3.28	416.3544	DAG 20:1; DAG 6:0-14:1; [M+NH4] <sup>+</sup>
2.75	412.2878	DAG 20:3; DAG 2:0-18:3; [M+NH4] <sup>+</sup>
2.19	432.3497	DAG 21:0; DAG 9:0-12:0; [M+NH4] <sup>+</sup>
5.31	428.3221	DAG 21:2; DAG 5:0-16:2; [M+NH4] <sup>+</sup>
1.65	424.324	DAG 21:4; DAG 5:0-16:4; [M+NH4] <sup>+</sup>
4.96	452.3371	DAG 23:4; DAG 7:0-16:4; [M+NH4] <sup>+</sup>
3.06	468.3869	DAG 24:3; DAG 6:0-18:3; [M+NH4] <sup>+</sup>
2.84	464.3538	DAG 24:5; DAG 4:0-20:5; [M+NH4] <sup>+</sup>
4.15	484.416	DAG 25:2; DAG 9:0-16:2; [M+NH4] <sup>+</sup>
2.14	476.3177	DAG 25:6; DAG 3:0-22:6; [M+NH4] <sup>+</sup>
3.1	512.416	DAG 27:2; DAG 11:0-16:2; [M+NH4] <sup>+</sup>
4.11	510.438	DAG 27:3; DAG 7:0-20:3; [M+NH4] <sup>+</sup>
3.44	526.4323	DAG 28:2; DAG 14:1-14:1; [M+NH4] <sup>+</sup>
3.76	540.4475	DAG 29:2; DAG 13:0-16:2; [M+NH4] <sup>+</sup>
5.01	538.4621	DAG 29:3; DAG 7:0-22:3; [M+NH4] <sup>+</sup>
2.12	536.4103	DAG 29:4; DAG 13:1-16:3; [M+NH4] <sup>+</sup>
5.54	536.4109	DAG 29:4; DAG 5:0-24:4; [M+NH4] <sup>+</sup>
3.8	532.4153	DAG 29:6; DAG 7:0-22:6; [M+NH4] <sup>+</sup>
4.06	554.4628	DAG 30:2; DAG 14:1-16:1; [M+NH4] <sup>+</sup>
3.86	554.4572	DAG 30:2; DAG 6:0-24:2; [M+NH4] <sup>+</sup>
5.03	552.4813	DAG 30:3; DAG 8:0-22:3; [M+NH4] <sup>+</sup>
2.82	546.4022	DAG 30:6; DAG 8:0-22:6; [M+NH4] <sup>+</sup>
4.36	568.4781	DAG 31:2; DAG 15:1-16:1; [M+NH4] <sup>+</sup>
4.62	566.463	DAG 31:3; DAG 9:0-22:3; [M+NH4] <sup>+</sup>
3.06	560.4155	DAG 31:6; DAG 13:1-18:5; [M+NH4] <sup>+</sup>
4.98	596.5092	DAG 33:2; DAG 11:0-22:2; [M+NH4] <sup>+</sup>
6.06	612.5336	DAG 34:1; DAG 13:0-21:1; [M+NH4] <sup>+</sup>
6.31	604.5151	DAG 34:5; DAG 12:0-22:5; [M+NH4] <sup>+</sup>
4.53	600.4768	DAG 34:7; DAG 16:2-18:5; [M+NH4] <sup>+</sup>
5.63	620.5088	DAG 35:4; DAG 15:1-20:3; [M+NH4] <sup>+</sup>
3.66	628.4977	DAG 36:7; DAG 18:2-18:5; [M+NH4] <sup>+</sup>
6.15	648.5393	DAG 37:4; DAG 19:1-18:3; [M+NH4] <sup>+</sup>
4.25	656.5286	DAG 38:7; DAG 16:1-22:6; [M+NH4] <sup>+</sup>
5.22	656.5074	DAG 38:7; DAG 22:3-16:4; [M+NH4] <sup>+</sup>
4.5	654.5128	DAG 38:8; DAG 16:2-22:6; [M+NH4] <sup>+</sup>
5.31	668.5377	DAG 39:8; DAG 17:2-22:6; [M+NH4] <sup>+</sup>
4.85	684.5598	DAG 40:7; DAG 22:2-18:5; [M+NH4] <sup>+</sup>
4.34	682.5427	DAG 40:8; DAG 20:3-20:5; [M+NH4] <sup>+</sup>
6.6	696.5535	DAG 41:8; DAG 19:2-22:6; [M+NH4] <sup>+</sup>
2.64	722.6609	DAG 42:2; DAG 18:0-24:2; [M+NH4] <sup>+</sup>
5.34	708.5588	DAG 42:9; DAG 22:4-20:5; [M+NH4] <sup>+</sup>
7.29	724.6053	DAG 43:8; DAG 21:2-22:6; [M+NH4] <sup>+</sup>
6.39	732.5512	DAG 44:11; DAG 22:5-22:6; [M+NH4] <sup>+</sup>
6.02	736.5986	DAG 44:9; DAG 24:4-20:5; [M+NH4] <sup>+</sup>
7.19	762.5958	DAG 46:10; DAG 24:4-22:6; [M+NH4] <sup>+</sup>
7.91	792.6667	DAG 48:9; DAG 26:4-22:5; [M+NH4] <sup>+</sup>
4.84	413.2693	Diocetyl phthalate (also known as the production of plastic)
6.31	744.6244	HexCer-NDS d37:0; HexCer-NDS d21:0/16:0; [M+H] <sup>+</sup>
3.35	614.4825	HexCer-NS d28:2; HexCer-NS d16:1/12:1; [M+H] <sup>+</sup>
6.74	644.5001	HexCer-NS d30:1; HexCer-NS d16:1/14:0; [M+H] <sup>+</sup>
3.96	642.5135	HexCer-NS d30:2; HexCer-NS d18:1/12:1; [M+H] <sup>+</sup>
2.25	664.4615	HexCer-NS d32:5; HexCer-NS d14:3/18:2; [M+H] <sup>+</sup>

2.68	678.478	HexCer-NS d33:5; HexCer-NS d15:3/18:2; [M+H] <sup>+</sup>
8.57	754.631	HexCer-NS d38:2; HexCer-NS d14:1/24:1; [M+H] <sup>+</sup>
8.66	1084.9532	HexCer-NS d62:5; HexCer-NS d26:3/36:2; [M+H] <sup>+</sup>
1.72	434.2415	LPC 10:0; [M+Na] <sup>+</sup>
3.41	426.2667	LPC 11:0(SN2); [M+H] <sup>+</sup>
4.61	440.2678	LPC 12:0(SN2); [M+H] <sup>+</sup>
4.16	482.3297	LPC 15:0(SN2); [M+H] <sup>+</sup>
4.93	480.2951	LPC 15:1(SN2); [M+H] <sup>+</sup>
3.26	496.3411	LPC 16:0(SN2); [M+H] <sup>+</sup>
1.67	494.3277	LPC 16:1(SN2); [M+H] <sup>+</sup>
4.69	510.3595	LPC 17:0(SN2); [M+H] <sup>+</sup>
4.28	508.3435	LPC 17:1(SN2); [M+H] <sup>+</sup>
3.88	506.3289	LPC 17:2(SN2); [M+H] <sup>+</sup>
3.64	524.3748	LPC 18:0(SN2); [M+H] <sup>+</sup>
4.43	529.408	LPC 18:1(d7); [M+H] <sup>+</sup>
3.21	534.3411	LPC 19:2(SN2); [M+H] <sup>+</sup>
3.04	588.4109	LPC 21:0; [M+Na] <sup>+</sup>
3.53	578.4047	LPC 22:1(SN2); [M+H] <sup>+</sup>
2.33	576.4109	LPC 22:2(SN2); [M+H] <sup>+</sup>
3	604.4413	LPC 24:2(SN2); [M+H] <sup>+</sup>
3.07	600.4105	LPC 24:4(SN2); [M+H] <sup>+</sup>
4.99	420.2201	LPC 9:0; [M+Na] <sup>+</sup>
2.75	356.2258	LPE 10:0e; [M+H] <sup>+</sup>
2.45	384.2568	LPE 12:0e; [M+H] <sup>+</sup>
2.76	398.2721	LPE 13:0e; [M+H] <sup>+</sup>
1.91	440.3159	LPE 16:0e; [M+H] <sup>+</sup>
4.35	450.2462	LPE 16:2; [M+H] <sup>+</sup>
2.75	434.2691	LPE 16:3e; [M+H] <sup>+</sup>
3.16	468.3092	LPE 17:0; [M+H] <sup>+</sup>
3.3	468.3336	LPE 18:0e; [M+H] <sup>+</sup>
3.49	487.3626	LPE 18:1(d7); [M+H] <sup>+</sup>
4.99	476.2818	LPE 18:3; [M+H] <sup>+</sup>
5.75	482.365	LPE 19:0e; [M+H] <sup>+</sup>
5.64	494.364	LPE 20:1e; [M+H] <sup>+</sup>
3.3	490.3154	LPE 20:3e; [M+H] <sup>+</sup>
3.4	538.3901	LPE 22:0; [M+H] <sup>+</sup>
4.71	532.3436	LPE 22:3; [M+H] <sup>+</sup>
3.88	528.3094	LPE 22:5; [M+H] <sup>+</sup>
6.75	550.4262	LPE 24:1e; [M+H] <sup>+</sup>
3.13	544.3851	LPE 24:4e; [M+H] <sup>+</sup>
3.88	580.4359	LPE 25:0; [M+H] <sup>+</sup>
6.71	566.4569	LPE 25:0e; [M+H] <sup>+</sup>
7.22	594.4876	LPE 27:0e; [M+H] <sup>+</sup>
2.41	306.2837	MAG 13:0; [M+NH4] <sup>+</sup>
4.71	320.2964	MAG 14:0; [M+NH4] <sup>+</sup>
4.51	334.3143	MAG 15:0; [M+NH4] <sup>+</sup>
4.54	332.2967	MAG 15:1; [M+NH4] <sup>+</sup>
3.14	348.291	MAG 16:0; [M+NH4] <sup>+</sup>
2.41	346.2756	MAG 16:1; [M+NH4] <sup>+</sup>
4.21	342.2504	MAG 16:3; [M+NH4] <sup>+</sup>
2.6	340.2676	MAG 16:4; [M+NH4] <sup>+</sup>
3.08	362.3088	MAG 17:0; [M+NH4] <sup>+</sup>
4.26	358.3139	MAG 17:2; [M+NH4] <sup>+</sup>
5.57	376.3585	MAG 18:0; [M+NH4] <sup>+</sup>
4.26	381.3511	MAG 18:1(d7); [M+NH4] <sup>+</sup>
4.85	370.2812	MAG 18:3; [M+NH4] <sup>+</sup>

6.4	390.3738	MAG 19:0; [M+NH4] <sup>+</sup>
6.95	418.4045	MAG 21:0; [M+NH4] <sup>+</sup>
5.61	422.345	MAG 22:5; [M+NH4] <sup>+</sup>
5.92	446.4379	MAG 23:0; [M+NH4] <sup>+</sup>
2.75	180.107	MAG 4:0; [M+NH4] <sup>+</sup>
1.89	236.1694	MAG 8:0; [M+NH4] <sup>+</sup>
3.16	462.2151	PC 11:0; PC 4:0-7:0; [M+Na] <sup>+</sup>
3.46	476.2308	PC 12:0; PC 5:0-7:0; [M+Na] <sup>+</sup>
4.15	538.3902	PC 19:0e; PC 14:0e/5:0; [M+H] <sup>+</sup>
4.87	548.3204	PC 19:2; PC 3:0-16:2; [M+H] <sup>+</sup>
4.06	564.3719	PC 20:1; PC 2:0-18:1; [M+H] <sup>+</sup>
3.61	564.373	PC 20:1; PC 5:0-15:1; [M+H] <sup>+</sup>
5.69	590.3658	PC 22:2; PC 2:0-20:2; [M+H] <sup>+</sup>
4.36	592.4362	PC 23:1e; PC 14:1e/9:0; [M+H] <sup>+</sup>
3.44	622.4301	PC 24:0; PC 11:0-13:0; [M+H] <sup>+</sup>
2.29	620.4363	PC 24:1; PC 11:0-13:1; [M+H] <sup>+</sup>
2.73	634.4525	PC 25:1; PC 10:0-15:1; [M+H] <sup>+</sup>
2.98	632.4365	PC 25:2; PC 6:0-19:2; [M+H] <sup>+</sup>
6.25	632.43	PC 25:2; PC 8:0-17:2; [M+H] <sup>+</sup>
2.96	648.4661	PC 26:1; PC 12:0-14:1; [M+H] <sup>+</sup>
5.18	660.4972	PC 28:2e; PC 16:2e/12:0; [M+H] <sup>+</sup>
6.28	706.5356	PC 30:0; PC 15:0-15:0; [M+H] <sup>+</sup>
3.27	702.5328	PC 31:2e; PC 14:1e/17:1; [M+H] <sup>+</sup>
3.28	710.4811	PC 31:5; PC 15:1-16:4; [M+H] <sup>+</sup>
7.21	708.5081	PC 32:6e; PC 14:1e/18:5; [M+H] <sup>+</sup>
6.65	784.5828	PC 34:0; PC 12:0-22:0; [M+Na] <sup>+</sup>
6.48	782.5665	PC 34:1; PC 18:0-16:1; [M+Na] <sup>+</sup>
7.09	746.6017	PC 34:1e; PC 14:0e/20:1; [M+H] <sup>+</sup>
6.52	758.5668	PC 34:2; PC 10:0-24:2; [M+H] <sup>+</sup>
7.06	744.5869	PC 34:2e; PC 14:0e/20:2; [M+H] <sup>+</sup>
4.16	744.5804	PC 34:2e; PC 14:1e/20:1; [M+H] <sup>+</sup>
6.76	756.5488	PC 34:3; PC 18:0-16:3; [M+H] <sup>+</sup>
6.41	754.5337	PC 34:4; PC 17:2-17:2; [M+H] <sup>+</sup>
2.17	752.5128	PC 34:5; PC 16:0-18:5; [M+H] <sup>+</sup>
4.26	770.5947	PC 35:3; PC 15:1-20:2; [M+H] <sup>+</sup>
2.61	766.5291	PC 35:5; PC 17:1-18:4; [M+H] <sup>+</sup>
3.54	752.5598	PC 35:5e; PC 18:3e/17:2; [M+H] <sup>+</sup>
7.09	812.6159	PC 36:0; PC 12:0-24:0; [M+Na] <sup>+</sup>
4.12	788.6062	PC 36:1; PC 10:0-26:1; [M+H] <sup>+</sup>
6.91	810.5982	PC 36:1; PC 10:0-26:1; [M+Na] <sup>+</sup>
4.75	772.6116	PC 36:2e; PC 14:1e/22:1; [M+H] <sup>+</sup>
6.84	782.5642	PC 36:4; PC 18:2-18:2; [M+H] <sup>+</sup>
6.77	768.5869	PC 36:4e; PC 14:0e/22:4; [M+H] <sup>+</sup>
6.18	780.5516	PC 36:5; PC 18:0-18:5; [M+H] <sup>+</sup>
6.44	780.5506	PC 36:5; PC 20:2-16:3; [M+H] <sup>+</sup>
6.71	766.5721	PC 36:5e; PC 18:0e/18:5; [M+H] <sup>+</sup>
6.62	804.6611	PC 37:0; PC 16:0-21:0; [M+H] <sup>+</sup>
7.22	776.5559	PC 37:7e; PC 18:5e/19:2; [M+H] <sup>+</sup>
4.7	816.6381	PC 38:1; PC 17:0-21:1; [M+H] <sup>+</sup>
3.7	810.6019	PC 38:4; PC 18:1-20:3; [M+H] <sup>+</sup>
7.16	810.5968	PC 38:4; PC 19:2-19:2; [M+H] <sup>+</sup>
6.45	796.6216	PC 38:4e; PC 14:0e/24:4; [M+H] <sup>+</sup>
6.8	794.6022	PC 38:5e; PC 16:0e/22:5; [M+H] <sup>+</sup>
6.4	806.5672	PC 38:6; PC 16:2-22:4; [M+H] <sup>+</sup>
6.6	806.566	PC 38:6; PC 18:3-20:3; [M+H] <sup>+</sup>
6.76	792.5862	PC 38:6e; PC 16:0e/22:6; [M+H] <sup>+</sup>

6.5	804.5497	PC 38:7; PC 18:2-20:5; [M+H] <sup>+</sup>
7.75	832.6656	PC 39:0; PC 15:0-24:0; [M+H] <sup>+</sup>
3.08	842.5604	PC 39:6; PC 17:1-22:5; [M+Na] <sup>+</sup>
2.57	810.5549	PC 40:11e; PC 18:5e/22:6; [M+H] <sup>+</sup>
6.84	834.5987	PC 40:6; PC 20:3-20:3; [M+H] <sup>+</sup>
6.95	832.5811	PC 40:7; PC 16:3-24:4; [M+H] <sup>+</sup>
6.5	832.5827	PC 40:7; PC 18:2-22:5; [M+H] <sup>+</sup>
6.43	828.5494	PC 40:9; PC 18:3-22:6; [M+H] <sup>+</sup>
4.08	832.6329	PC 41:7e; PC 20:5e/21:2; [M+H] <sup>+</sup>
2.54	854.5814	PC 42:10; PC 20:5-22:5; [M+H] <sup>+</sup>
7.03	864.6849	PC 43:5e; PC 18:5e/25:0; [M+H] <sup>+</sup>
4.04	876.6595	PC 43:6; PC 21:1-22:5; [M+H] <sup>+</sup>
4.65	860.6649	PC 43:7e; PC 22:5e/21:2; [M+H] <sup>+</sup>
3.8	890.6583	PC 44:6; PC 20:2-24:4; [M+H] <sup>+</sup>
9.66	902.8162	PC 45:0e; PC 18:0e/27:0; [M+H] <sup>+</sup>
4.61	904.6923	PC 45:6; PC 23:0-22:6; [M+H] <sup>+</sup>
7.53	932.7495	PC 48:6e; PC 22:4e/26:2; [M+H] <sup>+</sup>
3.98	926.6876	PC 48:9e; PC 22:5e/26:4; [M+H] <sup>+</sup>
4.12	984.7329	PC 49:5; PC 27:0-22:5; [M+Na] <sup>+</sup>
5.07	548.3201	PE 22:2; PE 3:0-19:2; [M+H] <sup>+</sup>
3.24	592.3992	PE 25:1; PE 12:0-13:1; [M+H] <sup>+</sup>
2.77	590.4275	PE 26:2e; PE 18:2e/8:0; [M+H] <sup>+</sup>
5.49	664.5337	PE 31:0e; PE 14:0e/17:0; [M+H] <sup>+</sup>
3.36	666.4554	PE 32:6e; PE 14:1e/18:5; [M+H] <sup>+</sup>
6	692.5635	PE 33:0e; PE 14:0e/19:0; [M+H] <sup>+</sup>
6.76	734.567	PE 35:0; PE 17:0-18:0; [M+H] <sup>+</sup>
3.98	708.496	PE 35:6e; PE 18:4e/17:2; [M+H] <sup>+</sup>
6.61	746.5673	PE 36:1; PE 23:0-13:1; [M+H] <sup>+</sup>
5.84	728.559	PE 36:3e; PE 22:3e/14:0; [M+H] <sup>+</sup>
6.83	760.5833	PE 37:1; PE 18:0-19:1; [M+H] <sup>+</sup>
7.04	774.5973	PE 38:1; PE 21:0-17:1; [M+H] <sup>+</sup>
7.27	788.6135	PE 39:1; PE 19:0-20:1; [M+H] <sup>+</sup>
6.98	786.5978	PE 39:2; PE 18:0-21:2; [M+H] <sup>+</sup>
6.86	786.5983	PE 39:2; PE 20:0-19:2; [M+H] <sup>+</sup>
6.92	808.5805	PE 41:5; PE 19:0-22:5; [M+H] <sup>+</sup>
6.52	808.5828	PE 41:5; PE 23:0-18:5; [M+H] <sup>+</sup>
2.14	796.5389	PE 42:11e; PE 22:6e/20:5; [M+H] <sup>+</sup>
8.08	816.6984	PE 42:1e; PE 16:0e/26:1; [M+H] <sup>+</sup>
7	812.6591	PE 42:3e; PE 18:1e/24:2; [M+H] <sup>+</sup>
3.59	446.2206	PG 11:0; PG 5:0-6:0; [M+NH <sub>4</sub> ] <sup>+</sup>
6.83	850.5532	PG 41:8; PG 19:2-22:6; [M+NH <sub>4</sub> ] <sup>+</sup>
2.59	276.2368	Phytophingosine 15:0; [M+H] <sup>+</sup>
1.83	304.3065	Phytophingosine 17:0; [M+H] <sup>+</sup>
2.69	332.336	Phytophingosine 19:0; [M+H] <sup>+</sup>
3.39	360.366	Phytophingosine 21:0; [M+H] <sup>+</sup>
5.47	388.3611	Phytophingosine 23:0; [M+H] <sup>+</sup>
1.98	402.3765	Phytophingosine 24:0; [M+H] <sup>+</sup>
6.03	416.3914	Phytophingosine 25:0; [M+H] <sup>+</sup>
6.61	444.4223	Phytophingosine 27:0; [M+H] <sup>+</sup>
5.54	668.375	PS 28:6; PS 6:0-22:6; [M+H] <sup>+</sup>
2.64	722.5031	PS 31:0; PS 10:0-21:0; [M+H] <sup>+</sup>
3.14	798.5334	PS 37:4; PS 21:1-16:3; [M+H] <sup>+</sup>
3.21	754.5072	SHexCer d32:0; SHexCer d18:0/14:0; [M+H] <sup>+</sup>
4.34	832.5496	SHexCer d38:3; SHexCer d20:1/18:2; [M+H] <sup>+</sup>
3	587.4152	SM d26:3; SM d14:3/12:0; [M+H] <sup>+</sup>
7.56	605.4787	SM d27:1; SM d14:0/13:1; [M+H] <sup>+</sup>



5.51	599.4279	SM d27:4; SM d14:3/13:1; [M+H] <sup>+</sup>
5.03	641.4554	SM d30:4; SM d14:2/16:2; [M+H] <sup>+</sup>
8.05	659.5166	SM d31:2; SM d14:2/17:0; [M+H] <sup>+</sup>
5.72	675.5406	SM d32:1; SM d14:1/18:0; [M+H] <sup>+</sup>
8.95	671.535	SM d32:3; SM d14:1/18:2; [M+H] <sup>+</sup>
4.28	681.4922	SM d33:5; SM d17:3/16:2; [M+H] <sup>+</sup>
6.25	703.5724	SM d34:1; SM d14:1/20:0; [M+H] <sup>+</sup>
8.44	703.5994	SM d34:1; SM d21:0/13:1; [M+H] <sup>+</sup>
5.85	701.5561	SM d34:2; SM d21:2/13:0; [M+H] <sup>+</sup>
9.2	699.566	SM d34:3; SM d14:1/20:2; [M+H] <sup>+</sup>
7.25	697.5207	SM d34:4; SM d14:2/20:2; [M+H] <sup>+</sup>
8.75	695.5353	SM d34:5; SM d14:3/20:2; [M+H] <sup>+</sup>
8.13	715.56	SM d35:2; SM d14:1/21:1; [M+H] <sup>+</sup>
8.52	713.574	SM d35:3; SM d16:2/19:1; [M+H] <sup>+</sup>
7.51	711.5306	SM d35:4; SM d15:2/20:2; [M+H] <sup>+</sup>
6.75	731.6035	SM d36:1; SM d14:1/22:0; [M+H] <sup>+</sup>
6.37	729.5886	SM d36:2; SM d14:1/22:1; [M+H] <sup>+</sup>
6.27	727.5991	SM d36:3; SM d16:1/20:2; [M+H] <sup>+</sup>
6.25	725.5546	SM d36:4; SM d16:2/20:2; [M+H] <sup>+</sup>
7.33	723.5354	SM d36:5; SM d14:3/22:2; [M+H] <sup>+</sup>
6.51	739.5695	SM d37:4; SM d17:2/20:2; [M+H] <sup>+</sup>
6.75	753.5848	SM d38:4; SM d14:2/24:2; [M+H] <sup>+</sup>
7.97	753.582	SM d38:4; SM d14:3/24:1; [M+H] <sup>+</sup>
6.88	795.6607	SM d41:4; SM d15:2/26:2; [M+H] <sup>+</sup>
7.6	813.6812	SM d42:2; SM d21:1/21:1; [M+H] <sup>+</sup>
7.29	811.6606	SM d42:3; SM d14:1/28:2; [M+H] <sup>+</sup>
9.1	821.6562	SM d43:5; SM d15:3/28:2; [M+H] <sup>+</sup>
9.66	883.7721	SM d47:2; SM d14:1/33:1; [M+H] <sup>+</sup>
8.67	881.7552	SM d47:3; SM d15:1/32:2; [M+H] <sup>+</sup>
8.49	879.7394	SM d47:4; SM d15:2/32:2; [M+H] <sup>+</sup>
9.66	907.7718	SM d49:4; SM d21:2/28:2; [M+H] <sup>+</sup>
9.51	905.7562	SM d49:5; SM d15:3/34:2; [M+H] <sup>+</sup>
3.89	300.2941	Sphingosine 18:1; [M+H] <sup>+</sup>
4.56	328.3252	Sphingosine 20:1; [M+H] <sup>+</sup>
3.63	342.3194	Sphingosine 21:1; [M+H] <sup>+</sup>
7	652.5474	TAG 36:2; TAG 12:0-12:0-12:2; [M+NH4] <sup>+</sup>
7.23	650.549	TAG 36:3; TAG 12:0-12:0-12:3; [M+NH4] <sup>+</sup>
6.46	646.5173	TAG 36:5; TAG 12:0-12:2-12:3; [M+NH4] <sup>+</sup>
8.95	666.58	TAG 37:2; TAG 12:0-12:2-13:0; [M+NH4] <sup>+</sup>
3.3	658.508	TAG 37:6; TAG 12:3-12:3-13:0; [M+NH4] <sup>+</sup>
6.98	680.5734	TAG 38:2; TAG 12:2-13:0-13:0; [M+NH4] <sup>+</sup>
7.58	678.5791	TAG 38:3; TAG 12:3-13:0-13:0; [M+NH4] <sup>+</sup>
6.63	676.5695	TAG 38:4; TAG 12:1-12:3-14:0; [M+NH4] <sup>+</sup>
7.19	672.53	TAG 38:6; TAG 12:2-12:3-14:1; [M+NH4] <sup>+</sup>
3.62	672.5263	TAG 38:6; TAG 12:3-12:3-14:0; [M+NH4] <sup>+</sup>
9.2	694.6113	TAG 39:2; TAG 12:0-13:0-14:2; [M+NH4] <sup>+</sup>
8.96	692.5953	TAG 39:3; TAG 12:1-12:2-15:0; [M+NH4] <sup>+</sup>
8.74	690.5798	TAG 39:4; TAG 12:1-12:2-15:1; [M+NH4] <sup>+</sup>
5.38	690.5788	TAG 39:4; TAG 12:2-13:1-14:1; [M+NH4] <sup>+</sup>
3.91	686.5381	TAG 39:6; TAG 12:2-12:3-15:1; [M+NH4] <sup>+</sup>
2.92	692.4913	TAG 40:10; TAG 12:2-12:3-16:5; [M+NH4] <sup>+</sup>
4.2	700.5544	TAG 40:6; TAG 12:0-12:1-16:5; [M+NH4] <sup>+</sup>
9.42	722.642	TAG 41:2; TAG 12:0-14:0-15:2; [M+NH4] <sup>+</sup>
4.8	728.5857	TAG 42:6; TAG 12:0-14:1-16:5; [M+NH4] <sup>+</sup>
4.3	726.5685	TAG 42:7; TAG 14:2-14:2-14:3; [M+NH4] <sup>+</sup>
7.59	748.6354	TAG 43:3; TAG 12:1-13:0-18:2; [M+NH4] <sup>+</sup>

6.7	740.6002	TAG 43:7; TAG 12:3-14:1-17:3; [M+NH4] <sup>+</sup>
7.15	736.5405	TAG 43:9; TAG 12:3-13:1-18:5; [M+NH4] <sup>+</sup>
2.6	766.6912	TAG 44:1; TAG 12:0-14:1-18:0; [M+NH4] <sup>+</sup>
6.1	758.6394	TAG 44:5; TAG 12:2-14:0-18:3; [M+NH4] <sup>+</sup>
7.14	768.6309	TAG 45:7; TAG 14:2-14:3-17:2; [M+NH4] <sup>+</sup>
9.44	796.7357	TAG 46:0; TAG 14:0-16:0-16:0; [M+NH4] <sup>+</sup>
9.25	794.7197	TAG 46:1; TAG 14:0-16:0-16:1; [M+NH4] <sup>+</sup>
9.51	810.7521	TAG 47:0; TAG 15:0-16:0-16:0; [M+NH4] <sup>+</sup>
9.11	798.6728	TAG 47:6; TAG 12:0-13:0-22:6; [M+NH4] <sup>+</sup>
9.62	824.767	TAG 48:0; TAG 16:0-16:0-16:0; [M+NH4] <sup>+</sup>
9.46	822.7516	TAG 48:1; TAG 14:0-16:0-18:1; [M+NH4] <sup>+</sup>
9.29	820.7357	TAG 48:2; TAG 14:0-16:1-18:1; [M+NH4] <sup>+</sup>
9.18	812.6887	TAG 48:6; TAG 12:2-14:3-22:1; [M+NH4] <sup>+</sup>
9.78	852.7989	TAG 50:0; TAG 16:0-16:0-18:0; [M+NH4] <sup>+</sup>
9.63	850.7829	TAG 50:1; TAG 16:0-16:0-18:1; [M+NH4] <sup>+</sup>
9.48	848.7679	TAG 50:2; TAG 16:0-16:1-18:1; [M+NH4] <sup>+</sup>
9.31	846.7516	TAG 50:3; TAG 16:0-16:1-18:2; [M+NH4] <sup>+</sup>
7.65	842.6277	TAG 51:12; TAG 12:3-17:3-22:6; [M+NH4] <sup>+</sup>
9.92	880.8313	TAG 52:0; TAG 16:0-18:0-18:0; [M+NH4] <sup>+</sup>
7.26	878.8082	TAG 52:1; TAG 12:0-18:0-22:1; [M+NH4] <sup>+</sup>
9.81	878.8157	TAG 52:1; TAG 16:0-18:0-18:1; [M+NH4] <sup>+</sup>
7.92	856.6433	TAG 52:12; TAG 12:3-20:4-20:5; [M+NH4] <sup>+</sup>
9.65	876.7997	TAG 52:2; TAG 16:0-16:0-20:2; [M+NH4] <sup>+</sup>
9.5	874.7842	TAG 52:3; TAG 16:0-18:1-18:2; [M+NH4] <sup>+</sup>
9.36	872.768	TAG 52:4; TAG 16:0-16:1-20:3; [M+NH4] <sup>+</sup>
7.94	870.6589	TAG 53:12; TAG 12:3-19:3-22:6; [M+NH4] <sup>+</sup>
7.62	868.6436	TAG 53:13; TAG 12:3-19:5-22:5; [M+NH4] <sup>+</sup>
3.85	868.6453	TAG 53:13; TAG 14:3-19:5-20:5; [M+NH4] <sup>+</sup>
7.35	866.6274	TAG 53:14; TAG 12:3-19:5-22:6; [M+NH4] <sup>+</sup>
8.08	884.6762	TAG 54:12; TAG 12:3-20:5-22:4; [M+NH4] <sup>+</sup>
9.82	904.8318	TAG 54:2; TAG 16:0-18:1-20:1; [M+NH4] <sup>+</sup>
9.53	900.8006	TAG 54:4; TAG 18:1-18:1-18:2; [M+NH4] <sup>+</sup>
9.37	898.7849	TAG 54:5; TAG 18:1-18:2-18:2; [M+NH4] <sup>+</sup>
9.19	896.7692	TAG 54:6; TAG 18:1-18:2-18:3; [M+NH4] <sup>+</sup>
8.29	898.6913	TAG 55:12; TAG 14:3-19:3-22:6; [M+NH4] <sup>+</sup>
8.34	938.7243	TAG 58:13; TAG 16:4-20:3-22:6; [M+NH4] <sup>+</sup>
4.57	948.7203	TAG 59:15; TAG 16:4-21:4-22:7; [M+NH4] <sup>+</sup>
8.6	966.7564	TAG 60:13; TAG 17:1-21:5-22:7; [M+NH4] <sup>+</sup>
7.97	1000.8161	TAG 62:10; TAG 20:1-20:2-22:7; [M+NH4] <sup>+</sup>
4.52	992.7492	TAG 62:14; TAG 20:5-20:6-22:3; [M+NH4] <sup>+</sup>
8.7	1006.7911	TAG 63:14; TAG 21:4-21:5-21:5; [M+NH4] <sup>+</sup>
4.25	1042.7799	TAG 66:17; TAG 22:5-22:5-22:7; [M+NH4] <sup>+</sup>
5.12	313.1418	w/o MS2: Benzyl butyl phthalate (also known as chemical reagent)
4.61	622.5173	w/o MS2:LPC 26:0e; [M+H] <sup>+</sup>
5.32	440.4484	w/o MS2:Sphingosine 28:1; [M+H] <sup>+</sup>

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**Supplementary Table 1b.** LipidBlast annotations of all measured m/z features in negative ionization mode

Average RT (min)	Feature/peak (m/z)	Annotated lipid species
4	319.2303	12-HETE
7.85	869.6244	AcylGlcADG 40:0; AcylGlcADG 12:0-12:0-16:0; [M-H]-
7.71	867.6087	AcylGlcADG 40:1; AcylGlcADG 12:0-14:0-14:1; [M-H]-
7.43	865.595	AcylGlcADG 40:2; AcylGlcADG 12:0-12:0-16:2; [M-H]-
6.02	863.6027	AcylGlcADG 40:3; AcylGlcADG 12:0-12:0-16:3; [M-H]-
7.87	897.6705	AcylGlcADG 42:0; AcylGlcADG 12:0-12:0-18:0; [M-H]-
8.18	897.6548	AcylGlcADG 42:0; AcylGlcADG 14:0-14:0-14:0; [M-H]-
8.08	895.6412	AcylGlcADG 42:1; AcylGlcADG 12:0-14:1-16:0; [M-H]-
8.48	925.6866	AcylGlcADG 44:0; AcylGlcADG 12:0-16:0-16:0; [M-H]-
8.04	981.7627	AcylGlcADG 48:0; AcylGlcADG 14:0-16:0-18:0; [M-H]-
7.87	979.7463	AcylGlcADG 48:1; AcylGlcADG 14:0-14:1-20:0; [M-H]-
8.62	1033.7963	AcylGlcADG 52:2; AcylGlcADG 16:0-16:1-20:1; [M-H]-
6.89	564.4536	Cer-AP t34:3; Cer-AP t14:2/20:1; [M-H]-
7.15	636.4813	Cer-AP t36:4; Cer-AP t18:2/18:2; [M+FA-H]-
7.59	664.511	Cer-AP t38:4; Cer-AP t22:2/16:2; [M+FA-H]-
6.78	678.5439	Cer-AP t39:4; Cer-AP t23:2/16:2; [M+FA-H]-
6.7	566.5142	Cer-AS d35:1; Cer-AS d16:1/19:0; [M-H]-
8.44	664.6243	Cer-AS d42:1; Cer-AS d18:1/24:0; [M-H]-
6.84	646.5189	Cer-BS d38:5; Cer-BS d14:3/24:2; [M+FA-H]-
7.64	676.5648	Cer-BS d40:4; Cer-BS d14:3/26:1; [M+FA-H]-
8.03	704.5949	Cer-BS d42:4; Cer-BS d15:3/27:1; [M+FA-H]-
7.47	610.5344	Cer-NS d36:1; Cer-NS d18:1/18:0; [M+FA-H]-
7.47	564.5366	Cer-NS d36:1; Cer-NS d18:1/18:0; [M-H]-
1.74	265.1504	Dodecyl sulfate
5.14	325.191	Dodecylbenzenesulfonic acid
4.69	255.2359	FA 16:0; [M-H]-
6.76	253.2196	FA 16:1; [M-H]-
1.62	245.1573	FA 16:5; [M-H]-
7.19	269.2523	FA 17:0; [M-H]-
5.39	283.2668	FA 18:0; [M-H]-
4.81	281.251	FA 18:1; [M-H]-
4.34	279.2356	FA 18:2; [M-H]-
3.89	277.2204	FA 18:3; [M-H]-
5.22	297.2839	FA 19:0; [M-H]-
5.55	311.298	FA 20:0; [M-H]-
5.35	309.282	FA 20:1; [M-H]-
4.69	305.2505	FA 20:3; [M-H]-
4.51	303.2357	FA 20:4; [M-H]-
4.07	301.2196	FA 20:5; [M-H]-
3.6	299.2061	FA 20:6; [M-H]-
3.98	325.3206	FA 21:0; [M-H]-
6.7	339.3284	FA 22:0; [M-H]-
4.93	333.2827	FA 22:3; [M-H]-
2.94	327.2339	FA 22:6; [M-H]-
2.97	325.2134	FA 22:7; [M-H]-
2.1	325.2374	FAHFA 19:1; FAHFA 4:0/15:1; [M-H]-
1.94	337.2393	FAHFA 20:2; FAHFA 4:0/16:2; [M-H]-
2.05	335.2249	FAHFA 20:3; FAHFA 4:0/16:3; [M-H]-
2.83	331.1933	FAHFA 20:5; FAHFA 18:5/2:0; [M-H]-
3.88	349.2413	FAHFA 21:3; FAHFA 18:3/3:0; [M-H]-
3.16	359.2233	FAHFA 22:5; FAHFA 18:5/4:0; [M-H]-
4.46	377.2747	FAHFA 23:3; FAHFA 3:0/20:3; [M-H]-
3.02	385.2401	FAHFA 24:6; FAHFA 22:6/2:0; [M-H]-

2.9	399.256	FAHFA 25:6; FAHFA 22:6/3:0; [M-H]-
3.99	433.3326	FAHFA 27:3; FAHFA 11:0/16:3; [M-H]-
4.55	433.3336	FAHFA 27:3; FAHFA 18:3/9:0; [M-H]-
3.47	427.2863	FAHFA 27:6; FAHFA 22:6/5:0; [M-H]-
2.76	453.3875	FAHFA 28:0; FAHFA 18:0/10:0; [M-H]-
4.4	461.3631	FAHFA 29:3; FAHFA 20:3/9:0; [M-H]-
3.93	507.4396	FAHFA 32:1; FAHFA 12:0/20:1; [M-H]-
3.46	495.3436	FAHFA 32:7; FAHFA 16:4/16:3; [M-H]-
5.52	525.3958	FAHFA 34:6; FAHFA 16:3/18:3; [M-H]-
5.4	523.3845	FAHFA 34:7; FAHFA 18:5/16:2; [M-H]-
5.96	521.3665	FAHFA 34:8; FAHFA 18:5/16:3; [M-H]-
5.78	539.4122	FAHFA 35:6; FAHFA 18:5/17:1; [M-H]-
5.3	537.394	FAHFA 35:7; FAHFA 22:6/13:1; [M-H]-
4.12	563.497	FAHFA 36:1; FAHFA 14:1/22:0; [M-H]-
6.11	553.4328	FAHFA 36:6; FAHFA 16:4/20:2; [M-H]-
5.53	551.4111	FAHFA 36:7; FAHFA 22:6/14:1; [M-H]-
6.48	549.4042	FAHFA 36:8; FAHFA 18:5/18:3; [M-H]-
6.22	565.4265	FAHFA 37:7; FAHFA 20:5/17:2; [M-H]-
5.78	565.4263	FAHFA 37:7; FAHFA 22:6/15:1; [M-H]-
6.63	581.4538	FAHFA 38:6; FAHFA 22:6/16:0; [M-H]-
7.24	579.4412	FAHFA 38:7; FAHFA 18:4/20:3; [M-H]-
6.06	579.4413	FAHFA 38:7; FAHFA 18:5/20:2; [M-H]-
6.84	595.4725	FAHFA 39:6; FAHFA 22:5/17:1; [M-H]-
6.63	593.459	FAHFA 39:7; FAHFA 22:6/17:1; [M-H]-
7.21	591.4449	FAHFA 39:8; FAHFA 22:6/17:2; [M-H]-
6.12	601.4247	FAHFA 40:10; FAHFA 18:5/22:5; [M-H]-
7.11	609.4901	FAHFA 40:6; FAHFA 18:1/22:5; [M-H]-
7.74	607.472	FAHFA 40:7; FAHFA 18:5/22:2; [M-H]-
6.9	607.473	FAHFA 40:7; FAHFA 20:4/20:3; [M-H]-
6.57	607.4756	FAHFA 40:7; FAHFA 22:5/18:2; [M-H]-
7.34	605.4567	FAHFA 40:8; FAHFA 18:5/22:3; [M-H]-
5.99	605.457	FAHFA 40:8; FAHFA 20:4/20:4; [M-H]-
7.02	603.4427	FAHFA 40:9; FAHFA 18:5/22:4; [M-H]-
7.34	623.4977	FAHFA 41:6; FAHFA 24:4/17:2; [M-H]-
6.43	619.4747	FAHFA 41:8; FAHFA 22:6/19:2; [M-H]-
4.75	629.4535	FAHFA 42:10; FAHFA 20:5/22:5; [M-H]-
7.55	637.5183	FAHFA 42:6; FAHFA 24:4/18:2; [M-H]-
8.07	635.5027	FAHFA 42:7; FAHFA 20:3/22:4; [M-H]-
7.56	633.4908	FAHFA 42:8; FAHFA 20:4/22:4; [M-H]-
8.45	663.532	FAHFA 44:7; FAHFA 22:3/22:4; [M-H]-
6.87	715.5743	FAHFA 48:9; FAHFA 26:4/22:5; [M-H]-
4.02	631.3989	GlcADG 24:0; GlcADG 12:0-12:0; [M-H]-
4	659.4298	GlcADG 26:0; GlcADG 12:0-14:0; [M-H]-
4.29	657.4215	GlcADG 26:1; GlcADG 12:0-14:1; [M-H]-
4.51	673.4442	GlcADG 27:0; GlcADG 12:0-15:0; [M-H]-
6.46	669.4136	GlcADG 27:2; GlcADG 13:1-14:1; [M-H]-
6.87	701.4918	GlcADG 29:0; GlcADG 14:0-15:0; [M-H]-
4.5	697.4395	GlcADG 29:2; GlcADG 12:0-17:2; [M-H]-
4.08	693.4136	GlcADG 29:4; GlcADG 13:0-16:4; [M-H]-
5.52	707.4457	GlcADG 30:4; GlcADG 14:0-16:4; [M-H]-
4.49	707.4302	GlcADG 30:4; GlcADG 14:1-16:3; [M-H]-
4.59	705.4153	GlcADG 30:5; GlcADG 12:0-18:5; [M-H]-
4.55	729.5056	GlcADG 31:0; GlcADG 15:0-16:0; [M-H]-
4.68	727.4928	GlcADG 31:1; GlcADG 15:0-16:1; [M-H]-
5.08	725.4856	GlcADG 31:2; GlcADG 15:0-16:2; [M-H]-
4.17	723.4688	GlcADG 31:3; GlcADG 14:1-17:2; [M-H]-

4.93	721.4446	GlcADG 31:4; GlcADG 15:0-16:4; [M-H]-
4.51	743.5303	GlcADG 32:0; GlcADG 16:0-16:0; [M-H]-
6.78	739.4828	GlcADG 32:2; GlcADG 15:1-17:1; [M-H]-
5	735.4609	GlcADG 32:4; GlcADG 16:1-16:3; [M-H]-
5.47	733.457	GlcADG 32:5; GlcADG 14:0-18:5; [M-H]-
4.64	727.4023	GlcADG 32:8; GlcADG 16:4-16:4; [M-H]-
7.27	755.5382	GlcADG 33:1; GlcADG 16:0-17:1; [M-H]-
5.04	761.4749	GlcADG 34:5; GlcADG 16:0-18:5; [M-H]-
5.13	759.4586	GlcADG 34:6; GlcADG 12:0-22:6; [M-H]-
4.49	757.4513	GlcADG 34:7; GlcADG 18:3-16:4; [M-H]-
5.31	773.4754	GlcADG 35:6; GlcADG 13:0-22:6; [M-H]-
5.23	797.5666	GlcADG 36:1; GlcADG 14:0-22:1; [M-H]-
5.36	795.5563	GlcADG 36:2; GlcADG 15:1-21:1; [M-H]-
4.69	791.5179	GlcADG 36:4; GlcADG 18:1-18:3; [M-H]-
7.7	809.5734	GlcADG 37:2; GlcADG 15:0-22:2; [M-H]-
7.66	823.5833	GlcADG 38:2; GlcADG 20:0-18:2; [M-H]-
5.47	817.5394	GlcADG 38:5; GlcADG 22:1-16:4; [M-H]-
5.33	815.5218	GlcADG 38:6; GlcADG 16:1-22:5; [M-H]-
7.88	839.6127	GlcADG 39:1; GlcADG 18:0-21:1; [M-H]-
7.94	837.599	GlcADG 39:2; GlcADG 17:0-22:2; [M-H]-
5.67	831.5523	GlcADG 39:5; GlcADG 17:1-22:4; [M-H]-
5.88	829.5375	GlcADG 39:6; GlcADG 21:1-18:5; [M-H]-
8	851.6138	GlcADG 40:2; GlcADG 20:0-20:2; [M-H]-
7.96	849.5994	GlcADG 40:3; GlcADG 22:0-18:3; [M-H]-
7.69	849.5986	GlcADG 40:3; GlcADG 24:0-16:3; [M-H]-
6	847.6072	GlcADG 40:4; GlcADG 19:2-21:2; [M-H]-
7.41	847.5842	GlcADG 40:4; GlcADG 24:0-16:4; [M-H]-
3.96	843.5658	GlcADG 40:6; GlcADG 18:3-22:3; [M-H]-
5.95	865.6301	GlcADG 41:2; GlcADG 19:1-22:1; [M-H]-
8.12	865.6304	GlcADG 41:2; GlcADG 20:0-21:2; [M-H]-
6.35	861.6121	GlcADG 41:4; GlcADG 21:0-20:4; [M-H]-
7.08	853.5469	GlcADG 41:8; GlcADG 19:2-22:6; [M-H]-
5.8	879.6522	GlcADG 42:2; GlcADG 22:0-20:2; [M-H]-
8.3	879.6455	GlcADG 42:2; GlcADG 25:0-17:2; [M-H]-
5.8	869.5674	GlcADG 42:7; GlcADG 22:2-20:5; [M-H]-
8.07	891.645	GlcADG 43:3; GlcADG 22:1-21:2; [M-H]-
7.4	883.6053	GlcADG 43:7; GlcADG 21:1-22:6; [M-H]-
8.6	907.6776	GlcADG 44:2; GlcADG 20:1-24:1; [M-H]-
8.35	919.6759	GlcADG 45:3; GlcADG 24:1-21:2; [M-H]-
8.15	917.6619	GlcADG 45:4; GlcADG 19:0-26:4; [M-H]-
7.85	915.6487	GlcADG 45:5; GlcADG 21:1-24:4; [M-H]-
6.58	933.6898	GlcADG 46:3; GlcADG 26:0-20:3; [M-H]-
6.28	923.6148	GlcADG 46:8; GlcADG 24:2-22:6; [M-H]-
8.62	947.7083	GlcADG 47:3; GlcADG 26:1-21:2; [M-H]-
8.49	959.7078	GlcADG 48:4; GlcADG 24:2-24:2; [M-H]-
8.72	987.7383	GlcADG 50:4; GlcADG 26:0-24:4; [M-H]-
8.48	985.722	GlcADG 50:5; GlcADG 26:1-24:4; [M-H]-
8.27	983.7084	GlcADG 50:6; GlcADG 26:2-24:4; [M-H]-
6.84	618.4316	HexCer-AP t26:1; HexCer-AP t14:0/12:1; [M-H]-
7.56	674.4922	HexCer-AP t30:1; HexCer-AP t17:0/13:1; [M-H]-
8.13	730.5532	HexCer-AP t34:1; HexCer-AP t22:1/12:0; [M-H]-
6.89	906.6996	HexCer-AP t47:4; HexCer-AP t23:2/24:2; [M-H]-
2.94	602.4553	HexCer-NDS d27:0; HexCer-NDS d15:0/12:0; [M-H]-
6.5	644.5029	HexCer-NDS d30:0; HexCer-NDS d14:0/16:0; [M-H]-
3.58	670.5217	HexCer-NDS d32:1; HexCer-NDS d18:0/14:1; [M-H]-
5.5	652.4893	HexCer-NS d31:3; HexCer-NS d14:3/17:0; [M-H]-

7.67	760.565	HexCer-NS d39:5; HexCer-NS d23:3/16:2; [M-H]-
2.14	367.1934	LPA 13:0; [M-H]-
3.35	395.2249	LPA 15:0; [M-H]-
2.42	409.2394	LPA 16:0; [M-H]-
4.74	403.1958	LPA 16:3; [M-H]-
2.67	401.1734	LPA 16:4; [M-H]-
2.89	423.256	LPA 17:0; [M-H]-
2.37	421.2267	LPA 17:1; [M-H]-
2.58	437.2712	LPA 18:0; [M-H]-
2.67	435.255	LPA 18:1; [M-H]-
3.59	433.2422	LPA 18:2; [M-H]-
2.97	451.2809	LPA 19:0; [M-H]-
3.32	449.2709	LPA 19:1; [M-H]-
3.09	465.2983	LPA 20:0; [M-H]-
3.6	463.2841	LPA 20:1; [M-H]-
3.61	461.2696	LPA 20:2; [M-H]-
5.13	459.2461	LPA 20:3; [M-H]-
3.66	457.2402	LPA 20:4; [M-H]-
5.05	479.3191	LPA 21:0; [M-H]-
3.73	477.3026	LPA 21:1; [M-H]-
3.05	475.2884	LPA 21:2; [M-H]-
3.72	493.3321	LPA 22:0; [M-H]-
3.48	491.3165	LPA 22:1; [M-H]-
4.09	489.3014	LPA 22:2; [M-H]-
4.79	507.3499	LPA 23:0; [M-H]-
4.72	519.3471	LPA 24:1; [M-H]-
4.03	517.3295	LPA 24:2; [M-H]-
6.15	535.3806	LPA 25:0; [M-H]-
4.7	545.3575	LPA 26:2; [M-H]-
6.66	563.416	LPA 27:0; [M-H]-
3.39	496.3093	LPC 14:1e; [M+FA-H]-
2.45	524.3001	LPC 15:1; [M+FA-H]-
2.72	538.3502	LPC 17:1e; [M+FA-H]-
3.73	573.3939	LPC 18:1(d7); [M+FA-H]-
2.77	552.3687	LPC 18:1e; [M+FA-H]-
2.94	566.3846	LPC 19:1e; [M+FA-H]-
3.6	582.4146	LPC 20:0e; [M+FA-H]-
2.79	592.4003	LPC 21:2e; [M+FA-H]-
4.76	604.4003	LPC 22:3e; [M+FA-H]-
4.97	624.4628	LPC 23:0e; [M+FA-H]-
6.11	680.5259	LPC 27:0e; [M+FA-H]-
2.45	344.1549	LPC 3:0e; [M+FA-H]-
2.64	400.1777	LPC 6:0; [M+FA-H]-
2.39	386.1991	LPC 6:0e; [M+FA-H]-
3.09	400.2152	LPC 7:0e; [M+FA-H]-
3.54	428.2295	LPC 9:0e; [M+FA-H]-
3.36	368.189	LPE 10:0; [M-H]-
2.83	354.1955	LPE 10:0e; [M-H]-
2.21	368.219	LPE 11:0e; [M-H]-
2.44	396.2236	LPE 12:0; [M-H]-
2.44	382.2419	LPE 12:0e; [M-H]-
1.72	410.2344	LPE 13:0; [M-H]-
2.71	396.256	LPE 13:0e; [M-H]-
2.54	424.2506	LPE 14:0; [M-H]-
2.7	410.2715	LPE 14:0e; [M-H]-
2.45	408.2546	LPE 14:1e; [M-H]-

2.98	438.2685	LPE 15:0; [M-H]-
2.89	424.2849	LPE 15:0e; [M-H]-
2.48	452.283	LPE 16:0; [M-H]-
2.4	450.2657	LPE 16:1; [M-H]-
3.11	448.2501	LPE 16:2; [M-H]-
3.44	432.2563	LPE 16:3e; [M-H]-
3.56	444.2044	LPE 16:4; [M-H]-
2.59	466.2966	LPE 17:0; [M-H]-
1.8	452.318	LPE 17:0e; [M-H]-
3.94	472.2365	LPE 18:4; [M-H]-
1.64	492.3125	LPE 19:1; [M-H]-
5.54	476.3221	LPE 19:2e; [M-H]-
3.21	498.2688	LPE 20:5; [M-H]-
3.12	522.3567	LPE 21:0; [M-H]-
3.37	536.3737	LPE 22:0; [M-H]-
3.35	534.3591	LPE 22:1; [M-H]-
4.11	520.38	LPE 22:1e; [M-H]-
6.55	532.3448	LPE 22:2; [M-H]-
4.76	530.3297	LPE 22:3; [M-H]-
2.04	516.3512	LPE 22:3e; [M-H]-
4.53	514.3322	LPE 22:4e; [M-H]-
3.57	550.3901	LPE 23:0; [M-H]-
3.75	564.405	LPE 24:0; [M-H]-
3.85	578.4206	LPE 25:0; [M-H]-
5.6	564.4438	LPE 25:0e; [M-H]-
6.29	592.4727	LPE 27:0e; [M-H]-
2.34	326.1831	LPE 8:0e; [M-H]-
2.31	469.2599	LPG 15:0; [M-H]-
1.94	467.2462	LPG 15:1; [M-H]-
2.57	525.3226	LPG 19:0; [M-H]-
2.67	523.3064	LPG 19:1; [M-H]-
3.15	521.2985	LPG 19:2; [M-H]-
2.73	539.3308	LPG 20:0; [M-H]-
2.68	537.3215	LPG 20:1; [M-H]-
2.59	535.3069	LPG 20:2; [M-H]-
3.65	567.3704	LPG 22:0; [M-H]-
3.18	565.3563	LPG 22:1; [M-H]-
6.43	563.3282	LPG 22:2; [M-H]-
4.49	561.3216	LPG 22:3; [M-H]-
4.05	595.4016	LPG 24:0; [M-H]-
3.23	593.3836	LPG 24:1; [M-H]-
3.58	591.3652	LPG 24:2; [M-H]-
3.44	587.3403	LPG 24:4; [M-H]-
3.64	615.3688	LPG 26:4; [M-H]-
5.75	709.4324	LPI 26:1; [M-H]-
4.48	501.3157	Medicagenic acid
3.43	565.3229	MGDG 17:0; MGDG 7:0-10:0; [M+FA-H]-
6.63	579.3444	MGDG 18:0; MGDG 9:0-9:0; [M+FA-H]-
2.43	573.2901	MGDG 18:3; MGDG 2:0-16:3; [M+FA-H]-
5.56	593.3677	MGDG 19:0; MGDG 3:0-16:0; [M+FA-H]-
6.83	593.3596	MGDG 19:0; MGDG 9:0-10:0; [M+FA-H]-
6	607.3752	MGDG 20:0; MGDG 10:0-10:0; [M+FA-H]-
3.17	607.3704	MGDG 20:0; MGDG 9:0-11:0; [M+FA-H]-
6.33	597.2894	MGDG 20:5; MGDG 2:0-18:5; [M+FA-H]-
3.6	621.3913	MGDG 21:0; MGDG 10:0-11:0; [M+FA-H]-
5.5	621.3939	MGDG 21:0; MGDG 9:0-12:0; [M+FA-H]-

6.5	611.3046	MGDG 21:5; MGDG 3:0-18:5; [M+FA-H]-
5.02	635.4092	MGDG 22:0; MGDG 10:0-12:0; [M+FA-H]-
5.74	635.4092	MGDG 22:0; MGDG 9:0-13:0; [M+FA-H]-
3.6	631.3694	MGDG 22:2; MGDG 4:0-18:2; [M+FA-H]-
6.67	625.3224	MGDG 22:5; MGDG 2:0-20:5; [M+FA-H]-
5.98	649.4228	MGDG 23:0; MGDG 11:0-12:0; [M+FA-H]-
6.15	663.4405	MGDG 24:0; MGDG 10:0-14:0; [M+FA-H]-
6.23	663.4398	MGDG 24:0; MGDG 12:0-12:0; [M+FA-H]-
5.85	677.4552	MGDG 25:0; MGDG 10:0-15:0; [M+FA-H]-
6.32	677.4551	MGDG 25:0; MGDG 11:0-14:0; [M+FA-H]-
5.81	677.4564	MGDG 25:0; MGDG 12:0-13:0; [M+FA-H]-
3.98	675.444	MGDG 25:1; MGDG 11:0-14:1; [M+FA-H]-
5.59	675.4399	MGDG 25:1; MGDG 12:0-13:1; [M+FA-H]-
6.04	691.4697	MGDG 26:0; MGDG 11:0-15:0; [M+FA-H]-
6.72	689.455	MGDG 26:1; MGDG 12:0-14:1; [M+FA-H]-
4.67	685.4199	MGDG 26:3; MGDG 6:0-20:3; [M+FA-H]-
6.93	703.4695	MGDG 27:1; MGDG 9:0-18:1; [M+FA-H]-
6.55	733.5151	MGDG 29:0; MGDG 11:0-18:0; [M+FA-H]-
6.75	733.5183	MGDG 29:0; MGDG 14:0-15:0; [M+FA-H]-
7.27	747.5253	MGDG 30:0; MGDG 12:0-18:0; [M+FA-H]-
7.88	747.5317	MGDG 30:0; MGDG 7:0-23:0; [M+FA-H]-
6.35	747.5275	MGDG 30:0; MGDG 9:0-21:0; [M+FA-H]-
5.86	743.5011	MGDG 30:2; MGDG 12:0-18:2; [M+FA-H]-
6.84	761.5477	MGDG 31:0; MGDG 15:0-16:0; [M+FA-H]-
6.39	775.563	MGDG 32:0; MGDG 13:0-19:0; [M+FA-H]-
6.44	771.5242	MGDG 32:2; MGDG 16:1-16:1; [M+FA-H]-
7.4	829.6093	MGDG 36:1; MGDG 20:0-16:1; [M+FA-H]-
7.1	845.6404	MGDG 37:0; MGDG 13:0-24:0; [M+FA-H]-
6.98	843.6244	MGDG 37:1; MGDG 20:0-17:1; [M+FA-H]-
6.94	913.7015	MGDG 42:1; MGDG 22:0-20:1; [M+FA-H]-
7.6	913.7023	MGDG 42:1; MGDG 23:0-19:1; [M+FA-H]-
7.5	911.6871	MGDG 42:2; MGDG 24:0-18:2; [M+FA-H]-
6.59	983.6745	MGDG 48:8; MGDG 24:4-24:4; [M+FA-H]-
8.4	1049.8262	MGDG 52:3; MGDG 26:1-26:2; [M+FA-H]-
8.31	1047.8103	MGDG 52:4; MGDG 26:2-26:2; [M+FA-H]-
1.64	453.1941	MGDG 9:0; MGDG 3:0-6:0; [M+FA-H]-
2.99	339.2028	Norethisterone acetate
6.56	868.569	OxPC 38:5+1O; OxPC 18:1-20:4+1O; [M+FA-H]-
5.13	804.5173	OxPE 40:7+1O; OxPE 18:1-22:6+1O ; [M-H]-
6.41	781.5237	OxPG 34:0+2O; OxPG 16:0-18:0+2O; [M-H]-
6.09	813.5334	OxPG 38:4+1O; OxPG 18:0-20:4+1O; [M-H]-
5.93	800.5051	OxPS 36:3+1O; OxPS 18:0-18:3+1O; [M-H]-
3.43	381.1715	PA 13:0; PA 4:0-9:0; [M-H]-
2.33	421.193	PA 16:1; PA 3:0-13:1; [M-H]-
2.77	437.241	PA 17:0; PA 2:0-15:0; [M-H]-
2.99	451.2503	PA 18:0; PA 2:0-16:0; [M-H]-
2.47	449.235	PA 18:1; PA 4:0-14:1; [M-H]-
3.22	465.2658	PA 19:0; PA 2:0-17:0; [M-H]-
2.57	477.2688	PA 20:1; PA 7:0-13:1; [M-H]-
2.77	475.2475	PA 20:2; PA 2:0-18:2; [M-H]-
2.8	493.2953	PA 21:0; PA 10:0-11:0; [M-H]-
3.19	493.2956	PA 21:0; PA 2:0-19:0; [M-H]-
2.95	493.2946	PA 21:0; PA 5:0-16:0; [M-H]-
3.76	493.2962	PA 21:0; PA 7:0-14:0; [M-H]-
3.96	507.3121	PA 22:0; PA 10:0-12:0; [M-H]-
2.83	507.312	PA 22:0; PA 7:0-15:0; [M-H]-



2.79	505.2991	PA 22:1; PA 7:0-15:1; [M-H]-
3.15	503.2806	PA 22:2; PA 2:0-20:2; [M-H]-
4.32	521.3281	PA 23:0; PA 11:0-12:0; [M-H]-
3.86	521.3255	PA 23:0; PA 6:0-17:0; [M-H]-
3.28	521.3232	PA 23:0; PA 7:0-16:0; [M-H]-
3.46	521.3283	PA 23:0; PA 9:0-14:0; [M-H]-
3.13	519.3093	PA 23:1; PA 2:0-21:1; [M-H]-
3.66	519.3137	PA 23:1; PA 5:0-18:1; [M-H]-
2.85	519.3101	PA 23:1; PA 9:0-14:1; [M-H]-
4.48	517.3016	PA 23:2; PA 3:0-20:2; [M-H]-
3.08	535.3412	PA 24:0; PA 12:0-12:0; [M-H]-
3.82	535.3423	PA 24:0; PA 9:0-15:0; [M-H]-
3.1	533.3215	PA 24:1; PA 5:0-19:1; [M-H]-
3.81	549.3577	PA 25:0; PA 12:0-13:0; [M-H]-
5.25	547.3424	PA 25:1; PA 10:0-15:1; [M-H]-
3.9	547.3425	PA 25:1; PA 9:0-16:1; [M-H]-
3.12	545.3246	PA 25:2; PA 6:0-19:2; [M-H]-
4.51	545.3276	PA 25:2; PA 9:0-16:2; [M-H]-
3.4	543.3105	PA 25:3; PA 9:0-16:3; [M-H]-
4.09	563.3644	PA 26:0; PA 12:0-14:0; [M-H]-
4.05	561.3606	PA 26:1; PA 13:0-13:1; [M-H]-
3.48	561.3579	PA 26:1; PA 7:0-19:1; [M-H]-
2.86	561.3481	PA 26:1; PA 8:0-18:1; [M-H]-
4.81	559.3411	PA 26:2; PA 10:0-16:2; [M-H]-
5.04	557.3309	PA 26:3; PA 6:0-20:3; [M-H]-
4.47	577.399	PA 27:0; PA 11:0-16:0; [M-H]-
3.58	575.3725	PA 27:1; PA 9:0-18:1; [M-H]-
3.44	573.3579	PA 27:2; PA 13:1-14:1; [M-H]-
3.83	571.342	PA 27:3; PA 9:0-18:3; [M-H]-
4.12	589.3894	PA 28:1; PA 14:0-14:1; [M-H]-
4.29	587.3737	PA 28:2; PA 10:0-18:2; [M-H]-
3.63	587.3728	PA 28:2; PA 14:1-14:1; [M-H]-
3.87	585.3573	PA 28:3; PA 10:0-18:3; [M-H]-
3.78	583.344	PA 28:4; PA 2:0-26:4; [M-H]-
4.74	603.4064	PA 29:1; PA 15:0-14:1; [M-H]-
4.43	601.3889	PA 29:2; PA 9:0-20:2; [M-H]-
4.05	599.3728	PA 29:3; PA 9:0-20:3; [M-H]-
3.85	597.3582	PA 29:4; PA 13:1-16:3; [M-H]-
4.71	615.4027	PA 30:2; PA 11:0-19:2; [M-H]-
4.73	615.4028	PA 30:2; PA 13:0-17:2; [M-H]-
3.83	613.3889	PA 30:3; PA 14:1-16:2; [M-H]-
4.07	611.3722	PA 30:4; PA 14:1-16:3; [M-H]-
4.42	633.4504	PA 31:0; PA 14:0-17:0; [M-H]-
4.44	629.4177	PA 31:2; PA 12:0-19:2; [M-H]-
4.69	627.4034	PA 31:3; PA 13:0-18:3; [M-H]-
3.78	627.4028	PA 31:3; PA 13:1-18:2; [M-H]-
5.33	645.4501	PA 32:1; PA 12:0-20:1; [M-H]-
4.85	641.4207	PA 32:3; PA 16:1-16:2; [M-H]-
4.48	639.4043	PA 32:4; PA 12:0-20:4; [M-H]-
4.36	655.4343	PA 33:3; PA 15:1-18:2; [M-H]-
4.59	653.4193	PA 33:4; PA 17:1-16:3; [M-H]-
4.46	651.4039	PA 33:5; PA 13:0-20:5; [M-H]-
7.59	675.4974	PA 34:0; PA 16:0-18:0; [M-H]-
5.32	669.4573	PA 34:3; PA 17:1-17:2; [M-H]-
4.95	667.4335	PA 34:4; PA 18:1-16:3; [M-H]-
5.35	687.4963	PA 35:1; PA 19:0-16:1; [M-H]-

4.48	681.4496	PA 35:4; PA 13:1-22:3; [M-H]-
5.69	679.4341	PA 35:5; PA 17:0-18:5; [M-H]-
4.78	679.4353	PA 35:5; PA 19:1-16:4; [M-H]-
5.29	677.4186	PA 35:6; PA 13:0-22:6; [M-H]-
5.13	701.5115	PA 36:1; PA 22:0-14:1; [M-H]-
3.61	683.3616	PA 36:10; PA 18:5-18:5; [M-H]-
4.74	697.4778	PA 36:3; PA 18:1-18:2; [M-H]-
4.43	695.4655	PA 36:4; PA 18:2-18:2; [M-H]-
5.65	693.4503	PA 36:5; PA 16:1-20:4; [M-H]-
3.98	691.4354	PA 36:6; PA 16:1-20:5; [M-H]-
5.27	691.4343	PA 36:6; PA 18:1-18:5; [M-H]-
4.89	689.4174	PA 36:7; PA 20:3-16:4; [M-H]-
6.26	685.3851	PA 36:9; PA 16:4-20:5; [M-H]-
6.06	713.5107	PA 37:2; PA 18:0-19:2; [M-H]-
4.99	711.4964	PA 37:3; PA 15:0-22:3; [M-H]-
6.46	711.4925	PA 37:3; PA 17:0-20:3; [M-H]-
5.2	707.4641	PA 37:5; PA 17:0-20:5; [M-H]-
6.1	707.4658	PA 37:5; PA 19:0-18:5; [M-H]-
5.24	705.4504	PA 37:6; PA 19:1-18:5; [M-H]-
4.7	725.5107	PA 38:3; PA 18:1-20:2; [M-H]-
5.48	725.5109	PA 38:3; PA 20:1-18:2; [M-H]-
5.4	721.4776	PA 38:5; PA 18:1-20:4; [M-H]-
5.36	719.465	PA 38:6; PA 22:2-16:4; [M-H]-
5.18	717.4515	PA 38:7; PA 18:3-20:4; [M-H]-
5.28	739.5264	PA 39:3; PA 19:0-20:3; [M-H]-
5.45	733.4752	PA 39:6; PA 17:2-22:4; [M-H]-
6.33	731.4639	PA 39:7; PA 17:1-22:6; [M-H]-
6.06	755.5585	PA 40:2; PA 22:0-18:2; [M-H]-
6.33	741.4474	PA 40:9; PA 20:4-20:5; [M-H]-
5.87	769.5728	PA 41:2; PA 19:0-22:2; [M-H]-
6.48	783.5858	PA 42:2; PA 18:0-24:2; [M-H]-
5.7	779.5589	PA 42:4; PA 21:2-21:2; [M-H]-
5.37	793.573	PA 43:4; PA 23:0-20:4; [M-H]-
7.08	795.4931	PA 44:10; PA 22:4-22:6; [M-H]-
6.89	805.5759	PA 44:5; PA 20:1-24:4; [M-H]-
6.36	797.5306	PA 44:9; PA 22:3-22:6; [M-H]-
6.09	797.5125	PA 44:9; PA 26:4-18:5; [M-H]-
6.71	823.6207	PA 45:3; PA 27:0-18:3; [M-H]-
5.61	817.5704	PA 45:6; PA 19:2-26:4; [M-H]-
7.96	817.5668	PA 45:6; PA 23:0-22:6; [M-H]-
6.51	837.6357	PA 46:3; PA 26:0-20:3; [M-H]-
6.36	847.6218	PA 47:5; PA 25:0-22:5; [M-H]-
8.44	845.6101	PA 47:6; PA 25:0-22:6; [M-H]-
5.97	861.6332	PA 48:5; PA 22:1-26:4; [M-H]-
6.31	859.6201	PA 48:6; PA 26:1-22:5; [M-H]-
6.5	875.6539	PA 49:5; PA 27:0-22:5; [M-H]-
7.27	891.681	PA 50:4; PA 24:2-26:2; [M-H]-
7.1	905.696	PA 51:4; PA 25:0-26:4; [M-H]-
8.76	905.6967	PA 51:4; PA 27:0-24:4; [M-H]-
6.62	915.6813	PA 52:6; PA 26:2-26:4; [M-H]-
7.1	911.6505	PA 52:8; PA 26:4-26:4; [M-H]-
2.57	538.318	PC 16:1e; PC 14:1e/2:0; [M+FA-H]-
5.82	710.4338	PC 29:6e; PC 22:6e/7:0; [M+FA-H]-
6.87	804.572	PC 34:1; PC 16:0-18:1; [M+FA-H]-
6.53	802.5593	PC 34:2; PC 16:0-18:2; [M+FA-H]-
7.36	832.5991	PC 36:1; PC 17:0-19:1; [M+FA-H]-

6.82	830.5889	PC 36:2; PC 18:0-18:2; [M+FA-H]-
6.51	828.569	PC 36:3; PC 16:0-20:3; [M+FA-H]-
6.57	826.5565	PC 36:4; PC 16:0-20:4; [M+FA-H]-
7.22	824.5411	PC 36:5; PC 14:0-22:5; [M+FA-H]-
4.76	802.501	PC 36:9e; PC 20:5e/16:4; [M+FA-H]-
7.21	848.6301	PC 37:0; PC 13:0-24:0; [M+FA-H]-
6.99	854.5801	PC 38:4; PC 18:0-20:4; [M+FA-H]-
4.37	830.5316	PC 38:9e; PC 20:4e/18:5; [M+FA-H]-
4.76	930.6204	PC 44:8; PC 22:4-22:4; [M+FA-H]-
2.9	452.2789	PE 16:0e; PE 14:0e/2:0; [M-H]-
2.67	464.2816	PE 17:1e; PE 14:1e/3:0; [M-H]-
2.97	480.3084	PE 18:0e; PE 16:0e/2:0; [M-H]-
3.93	490.2469	PE 18:2; PE 2:0-16:2; [M-H]-
2.91	522.3219	PE 20:0; PE 5:0-15:0; [M-H]-
3	520.3073	PE 20:1; PE 6:0-14:1; [M-H]-
3.87	536.3392	PE 21:0; PE 7:0-14:0; [M-H]-
4.52	532.2897	PE 21:2; PE 3:0-18:2; [M-H]-
4.72	518.3334	PE 21:2e; PE 16:2e/5:0; [M-H]-
4.51	516.3115	PE 21:3e; PE 18:3e/3:0; [M-H]-
2.92	528.2775	PE 21:4; PE 5:0-16:4; [M-H]-
4.8	546.319	PE 22:2; PE 2:0-20:2; [M-H]-
5.09	546.3052	PE 22:2; PE 6:0-16:2; [M-H]-
1.83	532.3453	PE 22:2e; PE 18:2e/4:0; [M-H]-
4.08	564.3707	PE 23:0; PE 8:0-15:0; [M-H]-
3.93	568.2957	PE 24:5; PE 4:0-20:5; [M-H]-
3.2	590.3836	PE 25:1; PE 10:0-15:1; [M-H]-
3.62	576.4034	PE 25:1e; PE 16:1e/9:0; [M-H]-
4.36	574.3979	PE 25:2e; PE 18:2e/7:0; [M-H]-
4.14	582.314	PE 25:5; PE 3:0-22:5; [M-H]-
2.04	586.3917	PE 26:3e; PE 22:3e/4:0; [M-H]-
3.86	596.3252	PE 26:5; PE 4:0-22:5; [M-H]-
2.57	600.4056	PE 27:3e; PE 22:3e/5:0; [M-H]-
2.43	616.4374	PE 28:2e; PE 14:1e/14:1; [M-H]-
5.23	648.4593	PE 29:0; PE 14:0-15:0; [M-H]-
5.94	634.4779	PE 29:0e; PE 14:0e/15:0; [M-H]-
5.13	642.4591	PE 30:3e; PE 18:3e/12:0; [M-H]-
6.89	642.4546	PE 30:3e; PE 22:3e/8:0; [M-H]-
5.77	676.4924	PE 31:0; PE 6:0-25:0; [M-H]-
6.43	662.5117	PE 31:0e; PE 16:0e/15:0; [M-H]-
5.43	672.4601	PE 31:2; PE 13:0-18:2; [M-H]-
6.21	664.4426	PE 32:6e; PE 16:2e/16:4; [M-H]-
6.03	678.4559	PE 33:6e; PE 22:6e/11:0; [M-H]-
4.02	694.4747	PE 34:5e; PE 14:0e/20:5; [M-H]-
4.04	692.4662	PE 34:6e; PE 14:1e/20:5; [M-H]-
4.7	692.4653	PE 34:6e; PE 22:6e/12:0; [M-H]-
5.87	712.529	PE 35:3e; PE 18:1e/17:2; [M-H]-
5.81	710.5197	PE 35:4e; PE 20:4e/15:0; [M-H]-
3.82	722.4736	PE 35:5; PE 13:1-22:4; [M-H]-
4.83	720.4617	PE 35:6; PE 13:0-22:6; [M-H]-
5.39	720.4681	PE 35:6; PE 15:1-20:5; [M-H]-
3.97	706.4811	PE 35:6e; PE 22:6e/13:0; [M-H]-
6.9	744.5522	PE 36:1; PE 18:0-18:1; [M-H]-
3.92	736.4902	PE 36:5; PE 16:1-20:4; [M-H]-
5.57	762.5154	PE 38:6; PE 18:1-20:5; [M-H]-
7.64	768.5824	PE 39:3e; PE 18:3e/21:0; [M-H]-
5.37	794.5742	PE 40:4; PE 16:0-24:4; [M-H]-

6.56	780.5847	PE 40:4e; PE 22:3e/18:1; [M-H]-
6.44	778.5813	PE 40:5e; PE 18:0e/22:5; [M-H]-
5.39	784.4921	PE 40:9; PE 20:4-20:5; [M-H]-
6.7	808.5805	PE 41:4; PE 15:0-26:4; [M-H]-
6.28	820.5244	PE 44:12e; PE 22:6e/22:6; [M-H]-
6.38	848.6224	PE 44:5; PE 18:1-26:4; [M-H]-
6.63	930.7006	PE 50:6; PE 26:2-24:4; [M-H]-
3.2	563.3635	PEtOH 24:0; PEtOH 12:0-12:0; [M-H]-
3.87	603.4037	PEtOH 27:1; PEtOH 14:0-13:1; [M-H]-
4.59	619.4352	PEtOH 28:0; PEtOH 14:0-14:0; [M-H]-
4.02	643.4348	PEtOH 30:2; PEtOH 14:1-16:1; [M-H]-
4.28	643.4358	PEtOH 30:2; PEtOH 15:1-15:1; [M-H]-
4.08	641.4208	PEtOH 30:3; PEtOH 14:1-16:2; [M-H]-
4.7	657.4499	PEtOH 31:2; PEtOH 13:1-18:1; [M-H]-
4.06	657.4498	PEtOH 31:2; PEtOH 14:1-17:1; [M-H]-
3.82	655.4362	PEtOH 31:3; PEtOH 15:1-16:2; [M-H]-
4.57	671.465	PEtOH 32:2; PEtOH 16:1-16:1; [M-H]-
5.61	761.5099	PEtOH 39:6; PEtOH 19:2-20:4; [M-H]-
6.78	781.5732	PEtOH 40:3; PEtOH 24:1-16:2; [M-H]-
5.9	775.526	PEtOH 40:6; PEtOH 20:3-20:3; [M-H]-
5.89	807.5874	PEtOH 42:4; PEtOH 21:2-21:2; [M-H]-
6.58	817.5784	PEtOH 43:6; PEtOH 19:2-24:4; [M-H]-
6.38	815.5577	PEtOH 43:7; PEtOH 21:2-22:5; [M-H]-
6.24	829.5742	PEtOH 44:7; PEtOH 24:2-20:5; [M-H]-
8.62	891.6869	PEtOH 48:4; PEtOH 26:1-22:3; [M-H]-
6.94	915.6832	PEtOH 50:6; PEtOH 24:2-26:4; [M-H]-
2.59	539.3056	PG 19:0; PG 9:0-10:0; [M-H]-
2.44	553.3187	PG 20:0; PG 9:0-11:0; [M-H]-
5.52	593.3417	PG 23:1; PG 3:0-20:1; [M-H]-
6.04	593.3389	PG 23:1; PG 5:0-18:1; [M-H]-
5.73	591.3242	PG 23:2; PG 2:0-21:2; [M-H]-
3.37	603.3292	PG 24:3; PG 2:0-22:3; [M-H]-
6.02	677.4284	PG 29:1; PG 11:0-18:1; [M-H]-
3.83	675.4288	PG 29:2; PG 13:1-16:1; [M-H]-
4.05	671.3913	PG 29:4; PG 5:0-24:4; [M-H]-
6.19	691.4426	PG 30:1; PG 14:0-16:1; [M-H]-
4.33	689.4402	PG 30:2; PG 10:0-20:2; [M-H]-
5.83	709.4075	PG 32:6; PG 16:3-16:3; [M-H]-
4.72	725.444	PG 33:5; PG 15:1-18:4; [M-H]-
4	723.4182	PG 33:6; PG 11:0-22:6; [M-H]-
3.67	723.4218	PG 33:6; PG 17:2-16:4; [M-H]-
4.6	743.4999	PG 34:3; PG 14:0-20:3; [M-H]-
6.72	763.5539	PG 35:0; PG 9:0-26:0; [M-H]-
4.94	757.4999	PG 35:3; PG 13:1-22:2; [M-H]-
4.54	751.4551	PG 35:6; PG 13:0-22:6; [M-H]-
5.73	777.5786	PG 36:0; PG 17:0-19:0; [M-H]-
6.73	833.6393	PG 40:0; PG 20:0-20:0; [M-H]-
5.84	825.5623	PG 40:4; PG 16:0-24:4; [M-H]-
6.4	845.6381	PG 41:1; PG 27:0-14:1; [M-H]-
7.36	839.5782	PG 41:4; PG 25:0-16:4; [M-H]-
7.76	853.5942	PG 42:4; PG 16:2-26:2; [M-H]-
7.79	851.5789	PG 42:5; PG 16:1-26:4; [M-H]-
8.01	881.624	PG 44:4; PG 20:0-24:4; [M-H]-
8.1	879.6109	PG 44:5; PG 26:1-18:4; [M-H]-
8.34	965.7302	PG 50:4; PG 24:2-26:2; [M-H]-
6.29	599.2897	PI 17:0; PI 5:0-12:0; [M-H]-

6.5	613.3027	PI 18:0; PI 9:0-9:0; [M-H]-
6.65	627.3179	PI 19:0; PI 8:0-11:0; [M-H]-
5.86	641.3214	PI 20:0; PI 10:0-10:0; [M-H]-
6.61	631.2513	PI 20:5; PI 2:0-18:5; [M-H]-
6.34	683.3839	PI 23:0; PI 9:0-14:0; [M-H]-
7.34	809.5236	PI 32:0; PI 16:0-16:0; [M-H]-
6.07	807.4951	PI 32:1; PI 12:0-20:1; [M-H]-
7.44	823.5392	PI 33:0; PI 9:0-24:0; [M-H]-
7.55	837.5554	PI 34:0; PI 17:0-17:0; [M-H]-
5.16	865.579	PI 36:0; PI 10:0-26:0; [M-H]-
6.65	893.6118	PI 38:0; PI 18:0-20:0; [M-H]-
6.26	885.5521	PI 38:4; PI 19:2-19:2; [M-H]-
8.44	949.6769	PI 42:0; PI 17:0-25:0; [M-H]-
8.07	1047.7783	PI 49:0; PI 24:0-25:0; [M-H]-
4.24	575.3961	PMeOH 26:1; PMeOH 12:0-14:1; [M-H]-
3.77	601.3855	PMeOH 28:2; PMeOH 14:1-14:1; [M-H]-
4.11	629.4202	PMeOH 30:2; PMeOH 14:1-16:1; [M-H]-
5.11	703.437	PMeOH 36:7; PMeOH 16:3-20:4; [M-H]-
4.49	701.4249	PMeOH 36:8; PMeOH 18:4-18:4; [M-H]-
4.8	725.4235	PMeOH 38:10; PMeOH 18:5-20:5; [M-H]-
5.51	759.4957	PMeOH 40:7; PMeOH 22:2-18:5; [M-H]-
5.73	791.5576	PMeOH 42:5; PMeOH 20:1-22:4; [M-H]-
6.65	783.4962	PMeOH 42:9; PMeOH 20:3-22:6; [M-H]-
6.62	815.5591	PMeOH 44:7; PMeOH 22:1-22:6; [M-H]-
6.77	813.5434	PMeOH 44:8; PMeOH 22:4-22:4; [M-H]-
6.4	811.5266	PMeOH 44:9; PMeOH 24:4-20:5; [M-H]-
7.22	847.6261	PMeOH 46:5; PMeOH 24:0-22:5; [M-H]-
6.53	843.5895	PMeOH 46:7; PMeOH 20:3-26:4; [M-H]-
6.66	867.585	PMeOH 48:9; PMeOH 26:4-22:5; [M-H]-
6.6	929.6949	PMeOH 52:6; PMeOH 26:2-26:4; [M-H]-
4.6	630.3476	PS 25:3; PS 5:0-20:3; [M-H]-
6.09	678.4382	PS 28:0; PS 13:0-15:0; [M-H]-
4.36	710.4035	PS 31:5; PS 9:0-22:5; [M-H]-
4.69	708.3926	PS 31:6; PS 9:0-22:6; [M-H]-
4.61	724.4221	PS 32:5; PS 10:0-22:5; [M-H]-
7.07	742.4598	PS 33:3; PS 13:0-20:3; [M-H]-
7.88	854.5961	PS 41:3; PS 19:1-22:2; [M-H]-
3.99	297.2453	Ricinoleic acid
4.58	664.3855	SHexCer d26:2; [M-H]-
5.43	694.4133	SHexCer d28:1; [M-H]-
5.4	720.4339	SHexCer d30:2; [M-H]-
5.49	736.4553	SHexCer d31:1; [M-H]-
5.26	732.4462	SHexCer d31:3; [M-H]-
5.52	818.5316	SHexCer d37:2; [M-H]-
8.42	920.6796	SHexCer d44:0; [M-H]-
6.39	635.4495	SM d26:1; SM d14:0/12:1; [M+FA-H]-
3.71	633.4202	SM d26:2; SM d14:2/12:0; [M+FA-H]-
6.93	651.4777	SM d27:0; SM d14:0/13:0; [M+FA-H]-
4.08	649.4461	SM d27:1; SM d14:1/13:0; [M+FA-H]-
6.36	647.4469	SM d27:2; SM d14:1/13:1; [M+FA-H]-
3.92	661.4447	SM d28:2; SM d16:2/12:0; [M+FA-H]-
6.79	691.5032	SM d30:1; SM d16:0/14:1; [M+FA-H]-
6.92	689.4907	SM d30:2; SM d14:0/16:2; [M+FA-H]-
7.3	705.5204	SM d31:1; SM d19:0/12:1; [M+FA-H]-
4.02	701.483	SM d31:3; SM d15:1/16:2; [M+FA-H]-
4.78	717.5062	SM d32:2; SM d20:2/12:0; [M+FA-H]-

7.23	749.5853	SM d34:0; SM d17:0/17:0; [M+FA-H]-
8.54	749.5832	SM d34:0; SM d21:0/13:0; [M+FA-H]-
6.6	761.5782	SM d35:1; SM d19:1/16:0; [M+FA-H]-
7.93	759.5673	SM d35:2; SM d22:1/13:1; [M+FA-H]-
6.87	775.5937	SM d36:1; SM d19:1/17:0; [M+FA-H]-
7.62	817.6441	SM d39:1; SM d20:1/19:0; [M+FA-H]-
9.92	873.7043	SM d43:1; SM d22:1/21:0; [M+FA-H]-
8.6	931.7809	SM d47:0; SM d17:0/30:0; [M+FA-H]-
3.82	721.4098	SQDG 27:1; SQDG 12:0-15:1; [M-H]-
5.93	741.3986	SQDG 29:5; SQDG 13:1-16:4; [M-H]-
4.14	755.3959	SQDG 30:5; SQDG 12:0-18:5; [M-H]-
4.67	775.4549	SQDG 31:2; SQDG 15:1-16:1; [M-H]-
5.29	821.5532	SQDG 34:0; SQDG 12:0-22:0; [M-H]-
7.51	849.5674	SQDG 36:0; SQDG 17:0-19:0; [M-H]-
7.22	847.559	SQDG 36:1; SQDG 14:0-22:1; [M-H]-
7.56	841.4991	SQDG 36:4; SQDG 18:2-18:2; [M-H]-
7.12	863.5823	SQDG 37:0; SQDG 13:0-24:0; [M-H]-

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