

Metabolite Data -> Sequence: MEGAPRESS; Fitting algorithm: Osprey; Fitting Style: Separate; Selected subspecs: metab
 Fitting range: 0.5 to 4 ppm; Baseline knot spacing: 0.55 ppm; ph0: 15.94deg; ph1: 4.41deg; refShift: 0.46 Hz; refFWHM: 0.0
 Number of metabolites: 18; Number of MM/lipids: 8 scale: 0.00022392



Raw Water Ratio	
Asc	3.14e-06
Asp	8.99e-06
Cr	3.65e-06
CrCH2	0.00e+00
GABA	5.77e-05
GPC	9.76e-06
GSH	7.43e-06
Gln	2.26e-04
Glu	1.83e-04
ml	3.00e-06
Lac	0.00e+00
NAA	3.24e-04
NAAG	8.17e-05
PCh	6.78e-06
PCr	6.80e-06
PE	0.00e+00
sl	4.08e-07
Tau	0.00e+00
MM09	4.03e-05
MM12	0.00e+00
MM14	0.00e+00
MM17	0.00e+00
MM20	0.00e+00
Lip09	1.01e-20
Lip13	0.00e+00
Lip20	3.93e-21

Osprey Separate metab fit plot: sub-03_svs_mpress.dat

