



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: 11.54deg; ph1: 1.57deg; refShift: 0.75 Hz; refFWHM: 0.1
 Number of metabolites: 18; Number of MM/lipids: 8 scale: 0.00057096

Raw Water Ratio	
Asc	2.75e-06
Asp	1.61e-05
Cr	1.42e-04
CrCH2	8.31e-05
GABA	0.00e+00
GPC	7.32e-05
GSH	4.21e-05
Gln	4.63e-06
Glu	7.17e-05
ml	9.10e-05
Lac	1.34e-06
NAA	3.02e-04
NAAG	8.52e-05
PCh	9.88e-07
PCr	1.39e-04
PE	0.00e+00
sl	6.26e-06
Tau	2.06e-05
MM09	1.45e-05
MM12	6.16e-06
MM14	1.60e-05
MM17	1.38e-06
MM20	7.98e-05
Lip09	1.10e-06
Lip13	2.01e-05
Lip20	1.14e-05

Osprey Separate metab fit plot: sub-07_svs_mpress.dat

