



Metabolite Data -> Sequence: MEGAPRESS; Fitting algorithm: Osprey; Fitting Style: Separate; Selected subspecs: diff1
 Fitting range: 0.5 to 4 ppm; Baseline knot spacing: 0.4 ppm; ph0: 16.80; ph1: 4.84; refShift: 0.84 Hz; refFWHM: 0.06 ppm
 Number of metabolites: 18; Number of MM/lipids: 8 scale: 0.00065252; initial ph0: 15.00; initial ph1: 4.93

Raw Water Ratio

Asc:	2.47e-06
Asp:	0.00e+00
Cr:	0.00e+00
CrCH2:	4.09e-06
GABA:	3.47e-05
GPC:	3.01e-05
GSH:	8.71e-05
Gln:	1.71e-04
Glu:	4.71e-04
Ins:	4.77e-06
Lac:	0.00e+00
NAA:	6.38e-04
NAAG:	4.78e-05
PCh:	3.45e-05
PCr:	0.00e+00
PE:	0.00e+00
Scyllo:	2.70e-06
Tau:	2.07e-06
MM09:	8.47e-05
MM12:	0.00e+00
MM14:	0.00e+00
MM17:	0.00e+00
MM20:	0.00e+00
Lip09:	0.00e+00
Lip13:	4.92e-21
Lip20:	1.52e-20

