



Metabolite Data -> Sequence: HERCULES; Fitting algorithm: Osprey; Fitting Style: Separate;  
 Selected subspecs: sum  
 Fitting range: 0.5 to 4 ppm; Baseline knot spacing: 0.4 ppm; nb0: 7.80deg; nb1: 2.82deg;

Raw Water Ratio

Asc	1.50e-
Asp	04
Cr	2.79e-
GABA	05
	1.78e-
GPC	04
GSH	6.87e-
Gln	06
Glu	3.22e-
Ins	05
Lac	4.22e-
NAA	05
NAAG	2.36e-
	05
PCh	1.56e-
PCr	04
PE	2.85e-
Scyllo	04
Tau	3.43e-
MM09	04
	4.65e-
MM12	04
	7.80e-
MM14	05
	2.29e-
MM17	05
	1.37e-

## Osprey Separate sum fit plot: svs.dat

