

Metabolite Data -> Sequence: TE35#; Fitting algorithm: Osprey; Fitting Style: Separate; Selected subspecs: off

Fitting range: 0.2 to 4.2 ppm; Baseline knot spacing: 0.4 ppm; nh0: -12.45°; nh1: -6.26°; refShift:



Raw Water Ratio

Asc: 0.00e+0  
 Asp: 0  
 Cr: 5.50e-0  
 CrCH2: 5  
 5.58e-0  
 GABA: 5  
 GPC: 1.79e-0  
 GSH: 5  
 Gln: 6.66e-0  
 Glu: 7  
 H2O: 1.64e-0  
 Ins: 5  
 Lac: 2.83e-0  
 NAA: 5  
 NAAG: 8.89e-0  
 PCh: 6  
 PCr: 6.64e-0  
 PE: 5  
 Scyllo: 5.98e-0  
 Tau: 9  
 Tyros: 7.65e-0  
 MM09: 5  
 6.96e-0  
 MM12: 6  
 2.05e-0  
 MM14: 4  
 2.51e-0  
 MM17: 5  
 2.65e-0  
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## Osprey Separate off fit plot: MRH134\_005\_WIP\_TE35\_9\_1\_raw\_act.SDAT

