

Metabolite Data -> Sequence: OCC_R_SV_PRESS_35_prewake2#; Fitting algorithm: LCModel; Fitting Style: Separate; Selected subspecs: metab
 Fitting range: 0.5 to 4 ppm; Baseline knot spacing: 0.4 ppm; ph0: 11.00deg; ph1: 4.30deg; refShift: 0.01 Hz;

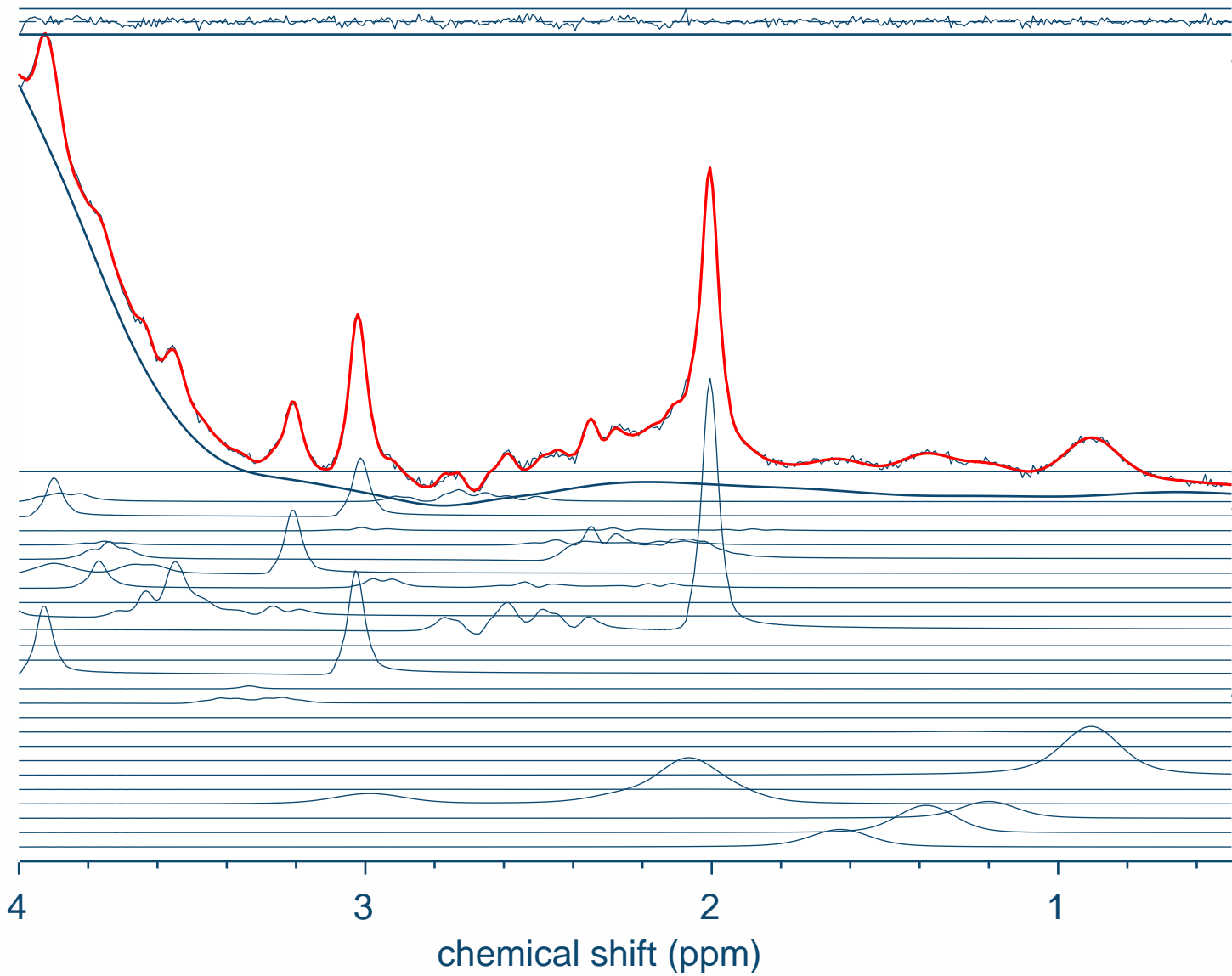


LCModel Separate metab fit plot: ses-prewakepostflicker-mrs_sub-019_act.SDAT

1.27e+02

4.19e+03

Raw Water Ratio	
Asp	8.11e+0
Cr	0
GABA	4.56e+0
Gln	0
Glu	1.16e+0
GPC	0
GSH	2.31e+0
Lac	0
ml	1.12e+0
NAA	1
NAAG	1.68e+0
PCh	0
PCr	2.65e+0
sl	0
Tau	0.00e+0
CrCH2	0
PCh_GPC	8.40e+0
Cr_PCr	1.99e+0
NAA_NA	1
AG	0.00e+0
Glu_Gln	0
Lip13a	0.00e+0
Lip13b	0
Lip09	8.04e+0
MM09	0
Lip20	1.07e-0
MM20	1.29e+0
MM12	0.00e+0
MM14	1.68e+0



Asp
 GABA
 Gln
 Glu
 GPC
 GSH
 Lac
 ml
 NAA
 NAAG
 PCh
 PCr
 sl
 Tau
 CrCH2
 PCh_GPC
 Cr_PCr
 NAA_NA
 AG
 Glu_Gln
 Lip13a
 Lip13b
 Lip09
 MM09
 Lip20
 MM20
 MM12
 MM14