

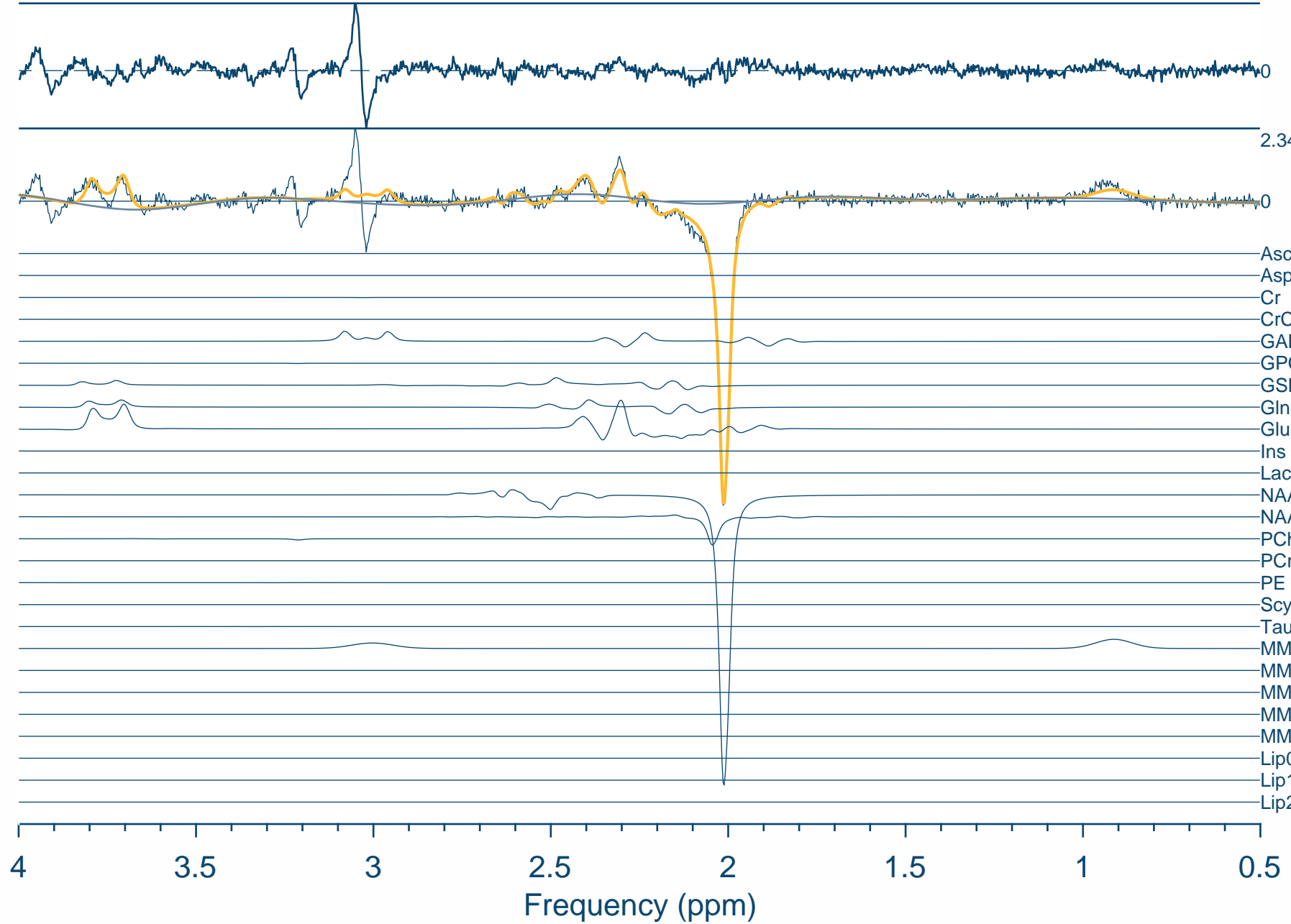


Metabolite Data -> Sequence: MEGAPRESS; Fitting algorithm: Osprey; Fitting Style: Separate; Selected subspecs: diff1
Fitting range: 0.5 to 4 ppm; Baseline knot spacing: 0.4 ppm; ph0: 1.67; ph1: -0.61; refShift: 0.90 Hz; refFWHM: 0.04 ppm
Number of metabolites: 18; Number of MM/lipids: 8 scale: 0.00093492; initial ph0: 0.40; initial ph1: -0.38

Raw Water Ratio

Asc:	2.89e-06
Asp:	0.00e+00
Cr:	1.74e-05
CrCH2:	2.28e-05
GABA:	9.08e-05
GPC:	2.10e-05
GSH:	1.01e-04
Gln:	1.51e-04
Glu:	4.93e-04
Ins:	0.00e+00
Lac:	0.00e+00
NAA:	5.54e-04
NAAG:	6.00e-05
PCh:	1.02e-04
PCr:	0.00e+00
PE:	2.26e-05
Scyllo:	0.00e+00
Tau:	6.82e-06
MM09:	5.14e-05
MM12:	1.04e-05
MM14:	1.03e-05
MM17:	1.51e-06
MM20:	0.00e+00
Lip09:	0.00e+00
Lip13:	0.00e+00
Lip20:	3.19e-20

Osprey Separate diff1 fit plot: meas_MID00026_FID24639_eja_svs_mpress_Gaba_VIS_1.dat



2.16

2.34

0

0

Asc

Asp

Cr

CrCH

GAB

GPC

GSH

Gln

Glu

Ins

Lac

NAA

NAAG

PCh

PCr

PE

Scyll

Tau

MM0

MM1

MM1

MM1

MM1

MM2

Lip0

Lip1

Lip2