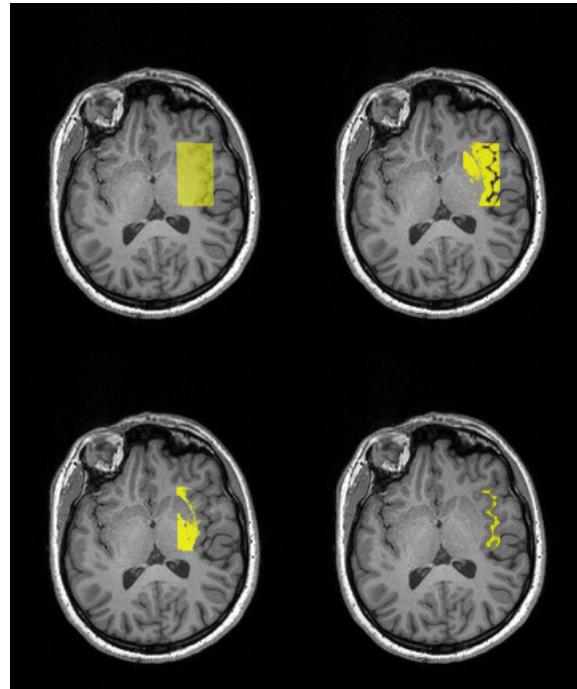


Voxel from meas_MID0019...A_INS.nii.gz on CBU_MPRAGE_3...series002.nii

Filename: meas_MID0019...A_INS.nii.gz

Anatomical image: CBU_MPRAGE_3...series002.nii

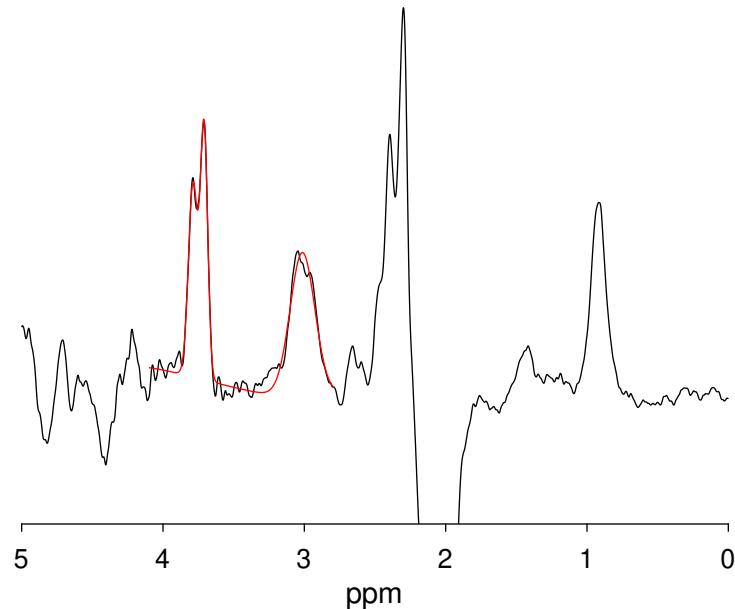
Relaxation-, tissue-corrected (Gasparovic et al. method)

GABA+/Water: 3.30 i.u.

Glx/Water: 12.20 i.u.

Relaxation-, tissue-, alpha-corrected (Harris et al. method)GABA+/Water ($\alpha = 0.5$): 3.04 i.u.Glx/Water ($\alpha = 0.5$): 11.24 i.u.**Relaxation-, tissue-, alpha-corrected; group-average-normalized (Harris et al. method)**GABA+/Water ($\alpha = 0.5$): 2.39 i.u.Glx/Water ($\alpha = 0.5$): 8.84 i.u.**Difference Spectrum and Model Fit**

QuantifyVer: 230314



3.3.1