

Description of Additional Supplementary Files

Supplementary Data 1 and 2:

Protein label-free quantification of CSF samples from 21.5 months old APPPS1 and WT mice with and without BACE inhibition (for mouse IDs see Supplementary Data 1; for Mass-Spec results see Supplementary Data 2). The protein LFQ intensities were log₂ transformed. A one-way ANOVA was applied to evaluate significant differences between the experimental groups. Afterwards, two-sided T-tests were applied between the individual groups. A permutation based FDR estimation ($p=0.05$; $s_0=0.1$) was applied to correct for multiple hypotheses.

Source File

Reporting Summary