

Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: Sample metadata for all proteomic samples.

File name: Supplementary Data 2

Description: Statistical analyses using limma linear regression modeling, postprocessed with DEqMS (Methods), for independent control-vs-SCZ analyses per Layer set from Figures 1b-d and between-Layers comparison from Figure 2a.

File name: Supplementary Data 3

Description: pQTL significant hits at FDR adjusted p-value < 0.05 from Figure 2b.

File name: Supplementary Data 4

Description: Stratified LDSC analyses from Figure 2c.

File name: Supplementary Data 5

Description: GO analyses using GOAT from Figure 3.

File name: Supplementary Data 6

Description: SynGO analysis via www.syngoportal.org from Figure 3f.