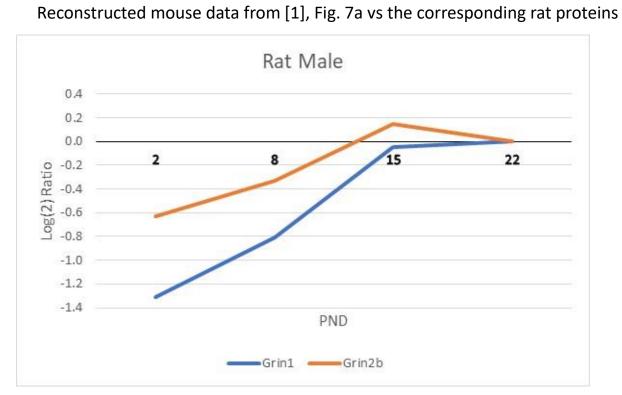
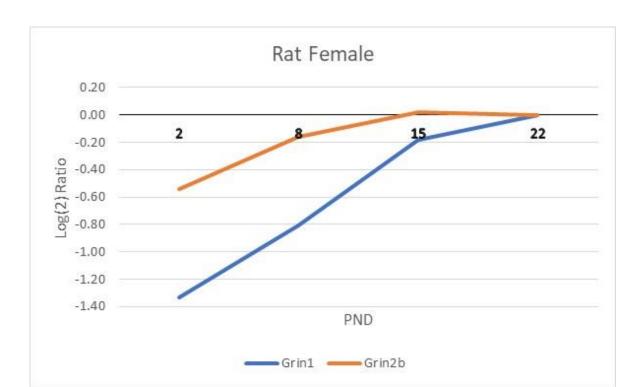
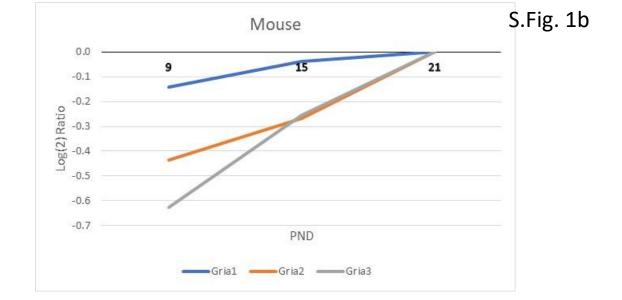
Caption for the supplementary S.Figures 1a-2c: The top left quadrants in S.Fig. 1a-2c show the mouse protein PND trendlines constructed from the S.Table 1 data in [1] for the proteins in the corresponding figures 7a-8c in [1] which match the rat protein gene symbols in the current work. The bottom graphs show the male and the female rat trendlines matching the PND trendlines above (current work). The reference point for the mouse P9/P21, P15/P21 logarithmic protein abundance ratios is P21, and for the rat (current work) P2/P22, P8/P22, P15/P22 logarithmic protein abundance ratios the reference point is P22. The upper right quadrants of these figures show, whenever available, the reconstructed trendlines which match rat protein gene symbols in the current work for the quantifiable synaptic cortex rat proteins calculated from Supplementary Table 5: Statistically significant changes in synaptosomal cortical proteome during development in [7], for which the PND protein ratio reference point is P20.

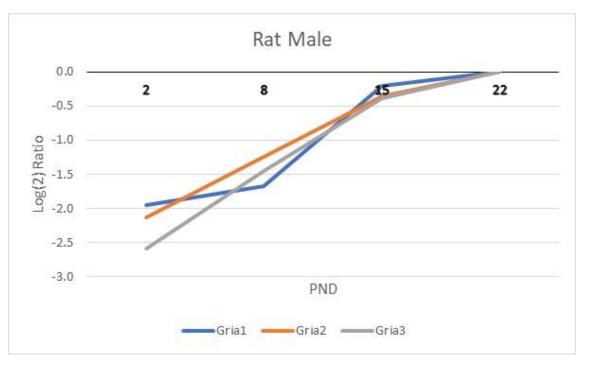


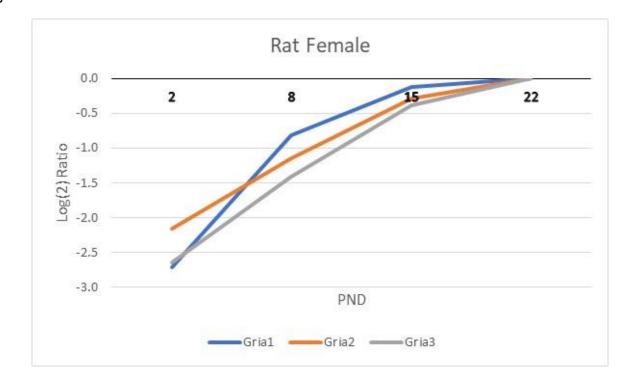






Reconstructed mouse data from [1], Fig. 7b vs the corresponding rat proteins

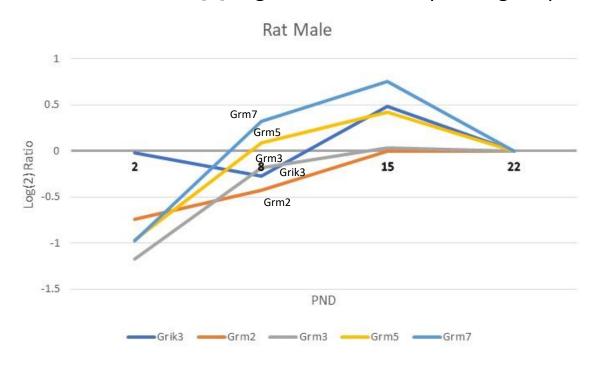


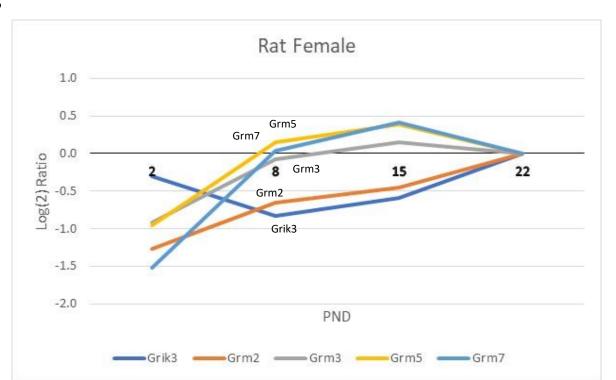


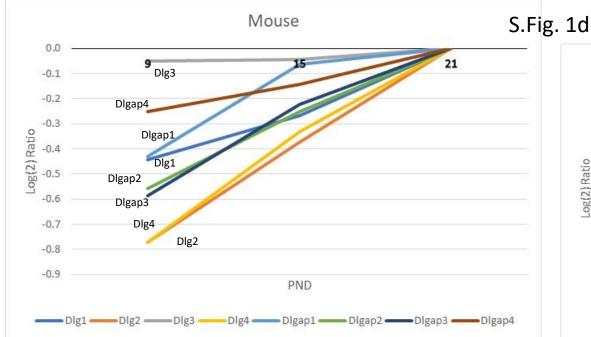


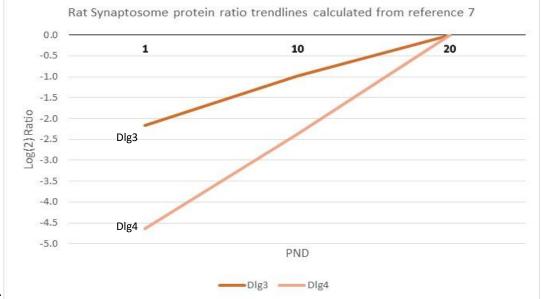
S.Fig. 1c

Mouse data from [1], Fig. 7c vs the corresponding rat proteins

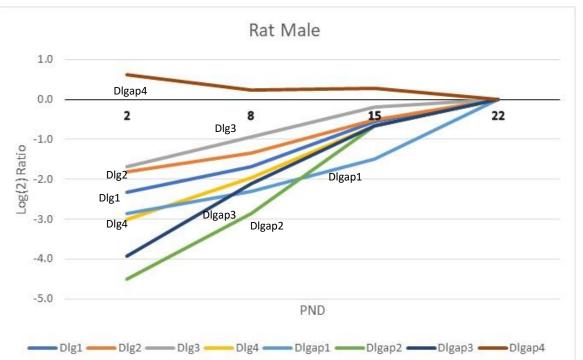


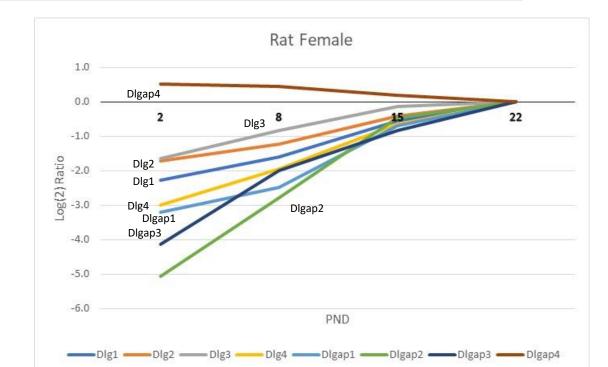


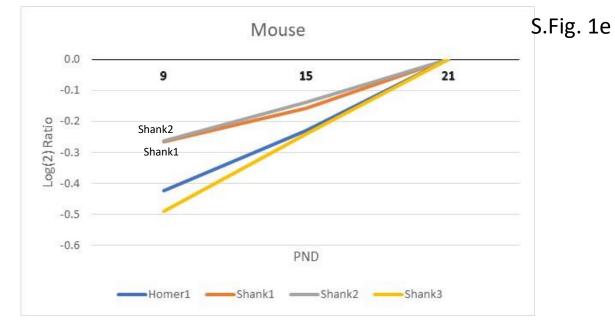




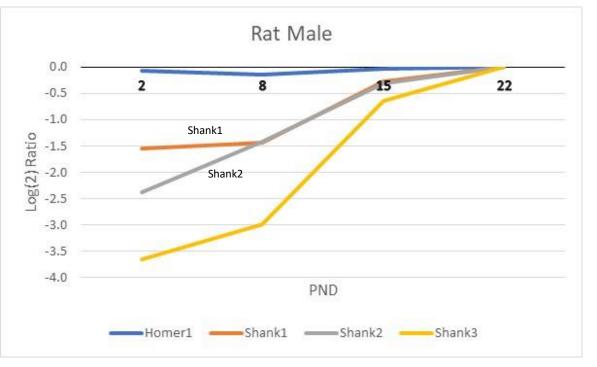


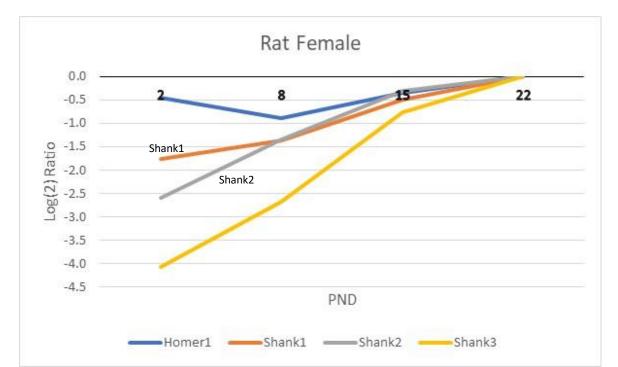


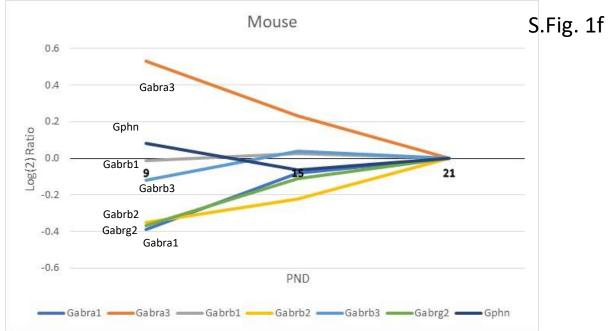




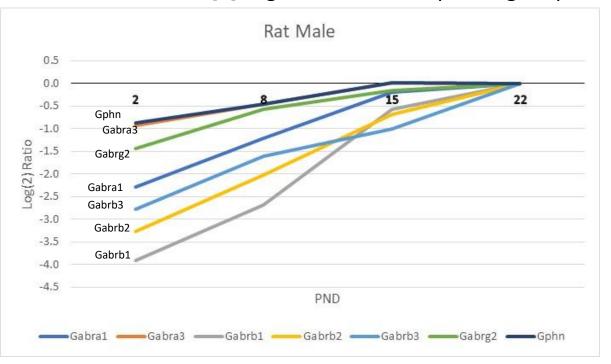
Mouse data from [1], Fig. 7e vs the corresponding rat proteins

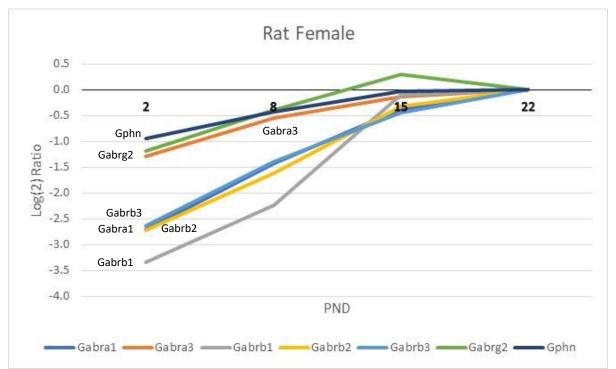


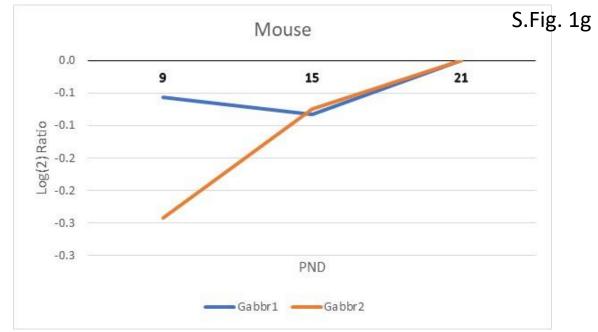




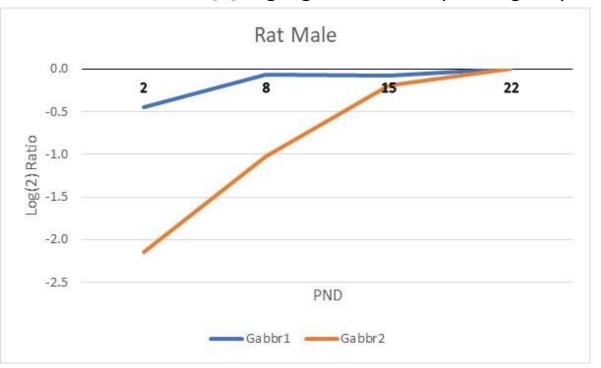
Mouse data from [1], Fig. 7f vs the corresponding rat proteins

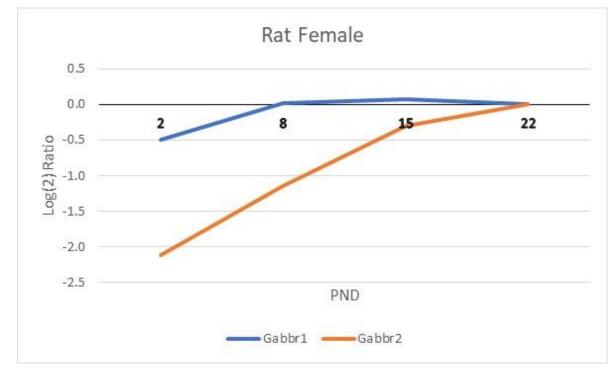


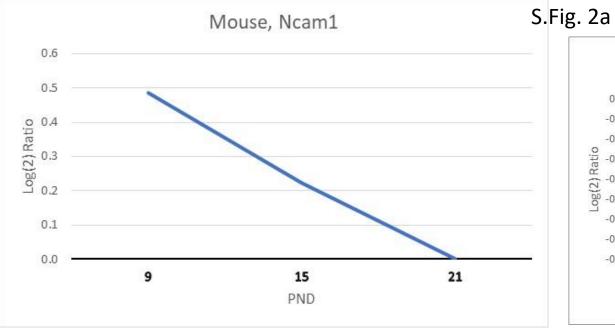


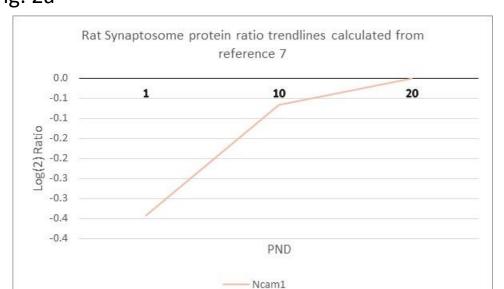


Mouse data from [1], Fig. 7g vs the corresponding rat proteins

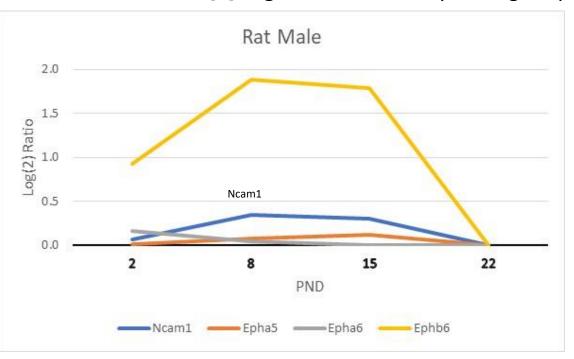


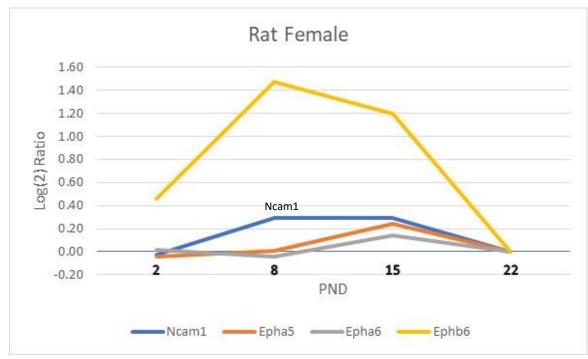


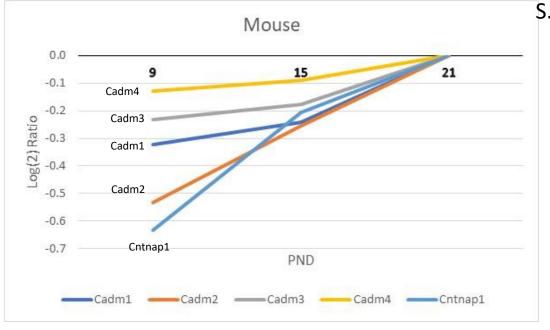




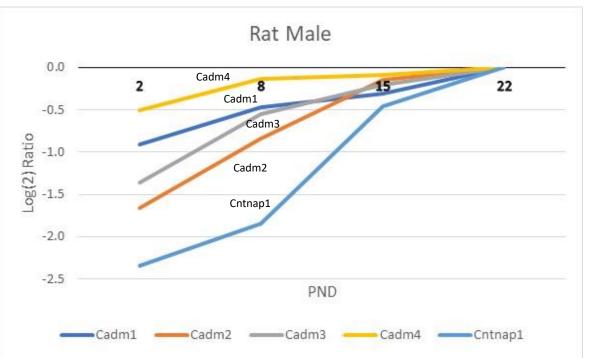
Mouse data from [1], Fig. 8a vs the corresponding rat proteins



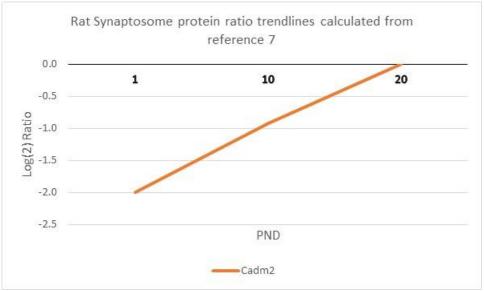




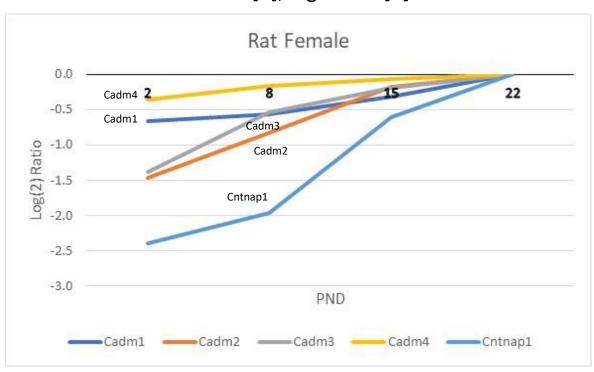
Mouse data from [1], Fig. 8b vs the corresponding rat proteins

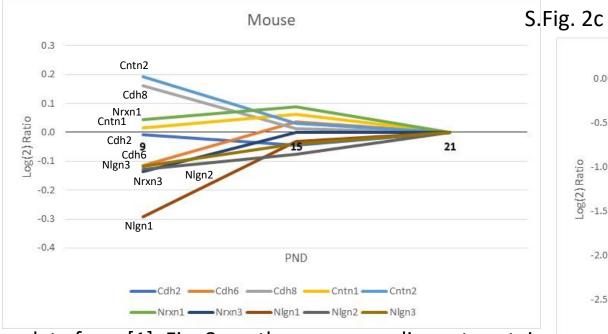


S.Fig. 2b

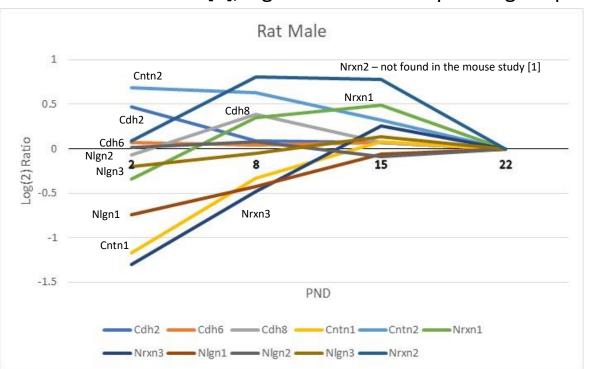


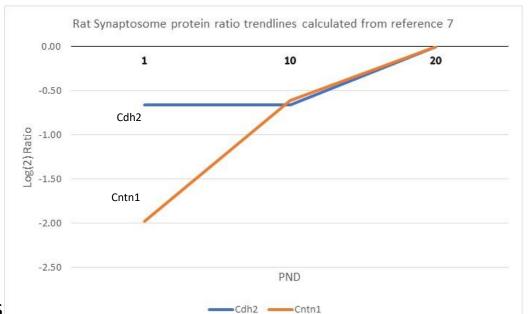
Mouse data[1], Fig. 8b in [1]

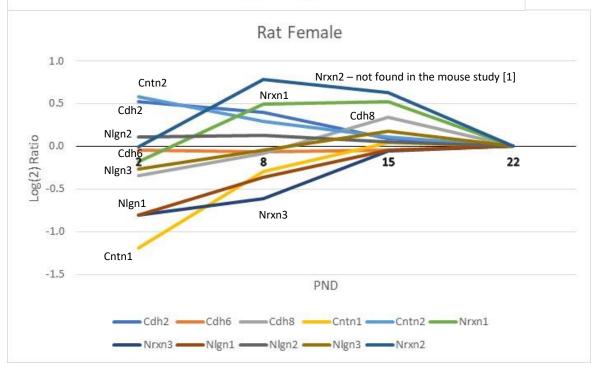




Mouse data from [1], Fig. 8c vs the corresponding rat proteins







Caption for the supplementary S.Figures 3a-c: The top left quadrants in S.Fig. 3a-c show the mouse protein PND trendlines constructed from the S.Table 1 data in [1] for the proteins in Table 1 representing high fold mouse PND protein ratio differences in [1], which match the rat protein gene symbols in the current work. The bottom graphs show the male and the female rat trendlines matching the PND trendlines above (current work). The reference point for the mouse P9/P21, P15/P21 logarithmic protein abundance ratios is P21, and for the rat (current work) P2/P22, P8/P22, P15/P22 logarithmic protein abundance ratios the reference point is P22. The upper right quadrants of these figures show, whenever available, the reconstructed trendlines which match rat protein gene symbols in the current work for the quantifiable synaptic cortex rat proteins calculated from Supplementary Table 5: Statistically significant changes in synaptosomal cortical proteome during development in [7], for which the PND protein ratio reference point is P20.

2.0 Mouse [1] Eef1a1 Atat1 Cat Cxadr Dcakd Dhcr7 Dpysl3 Bdh1 15 -0.5 Atp1a1 Camk2a — Dcakd — Dcx — Dhcr7 — Dpysl3 — Dpysl5 — Eef1a1

Rat Male

5.0

Dcx

Atat1

Dpysl3

Dcakd

-Atat1 -Atp1a1 -Bdh1 -Camk2a -Cat -Cxadr

—Dcakd —Dcx —Dhcr7 —Dpysl3 —Dpysl5 —Eef1a1

Dhcr7

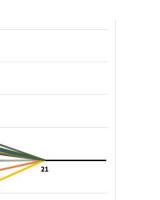
Cxadr

Bdh1

Camk2a

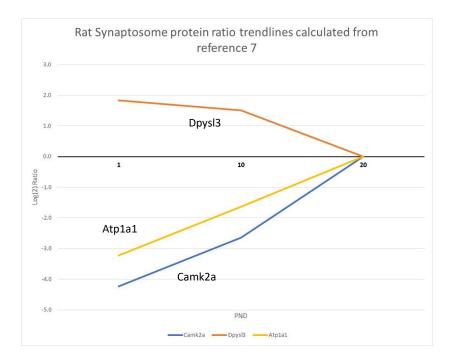
Atp1a1

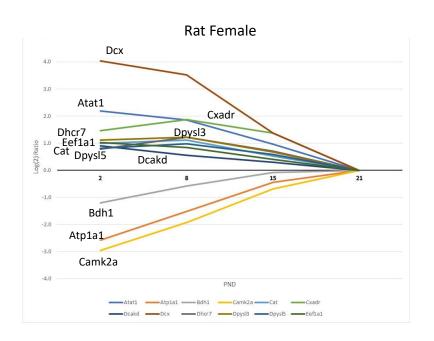
Lo Eef1a1 EDpysl5 Cat



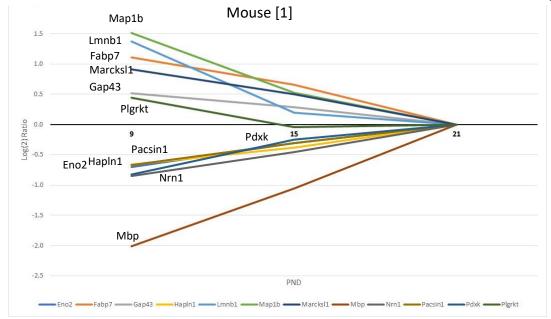


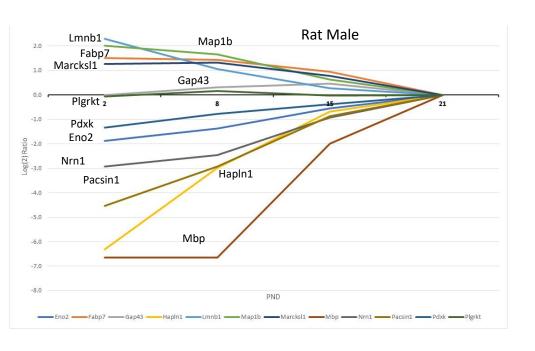


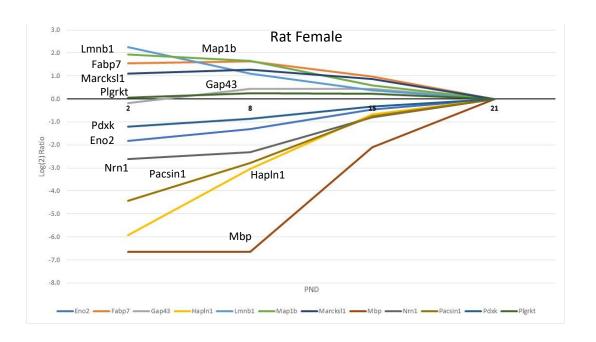


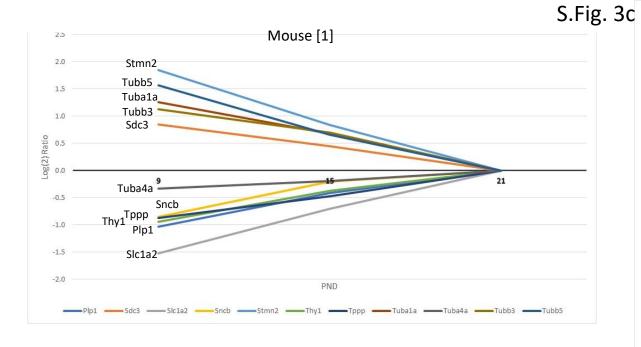


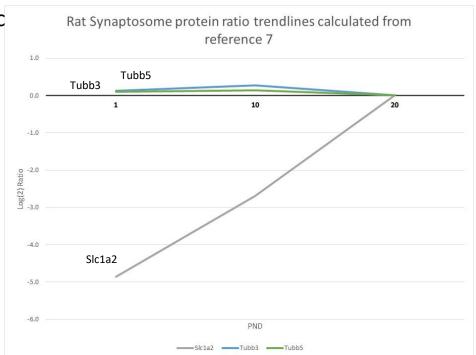
S.Fig. 3b

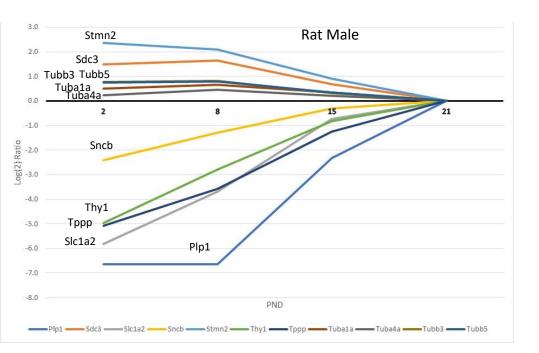


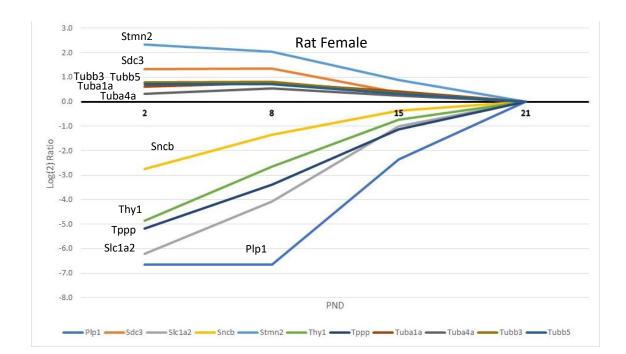












S.Fig. 4a

Box and whisker plot showing the log2 Pn/P22 (male and female rat rostral cortex) and log2 Px/P21 (mouse cortex [1]) ribosomal protein abundance ratio data point distributions for the PND points: n=2, 8, 15, 22 (rat), and x=9, 15, 21 (mouse)

S.Fig. 4b

Box and whisker plot showing the log2 Pn/P22 (male and female rat rostral cortex) and log2 Px/P21 (mouse cortex [1]) proteasome protein abundance ratio data point distributions for the PND points: n= 2, 8, 15, 22 (rat), and x= 9, 15, 21 (mouse)

S.Fig. 4c

Box and whisker plot showing the log2 Pn/P22 (male and female rat rostral cortex) mRNA splicing and mitochondrial protein abundance ratio data point distributions for the PND points: n= 2, 8, 15, 22

S.Fig. 4d

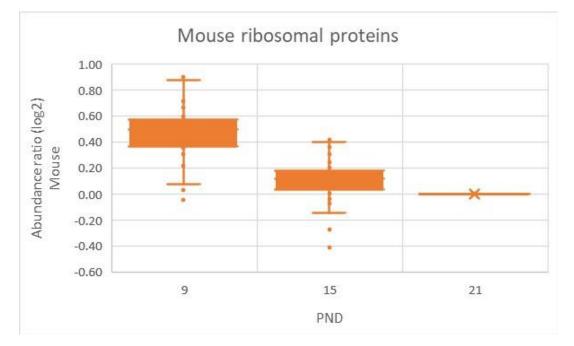
Box and whisker plot showing the log2 Pn/P22 (male and female rat rostral cortex) DNA, RNA associated and apoptosis-related protein abundance ratio data point distributions for the PND points: n= 2, 8, 15, 22

S.Fig. 5

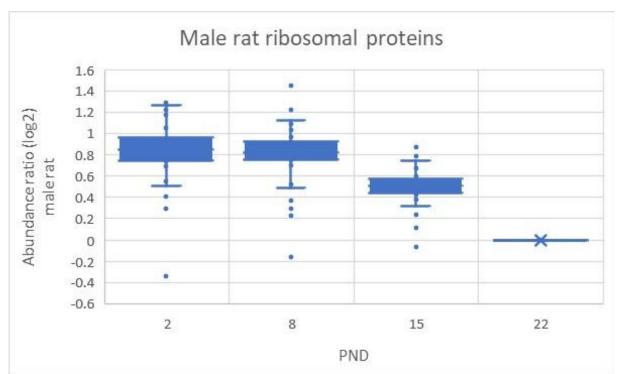
Box and whisker plot showing the log2 Pn/P22 (male and female rat rostral cortex) significantly rising and falling trend protein abundance ratio data point distributions for the PND points: n= 2, 8, 15, 22

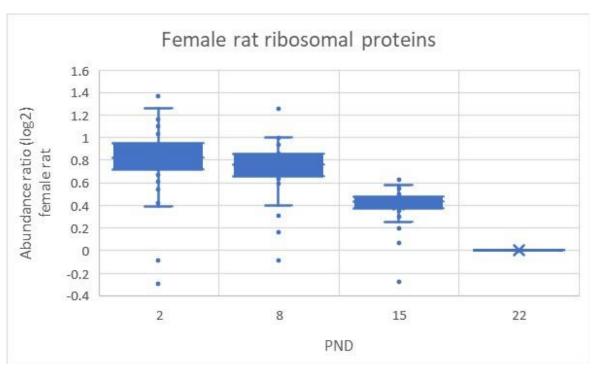
S.Fig 6

PCA grouping of the rat P2, P8, P15, P22 experimental PND male and female groups; six animals per group

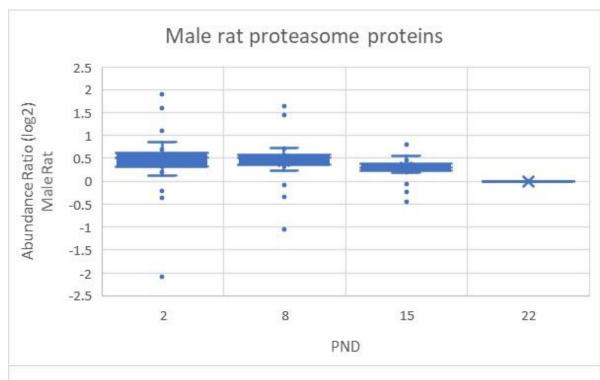


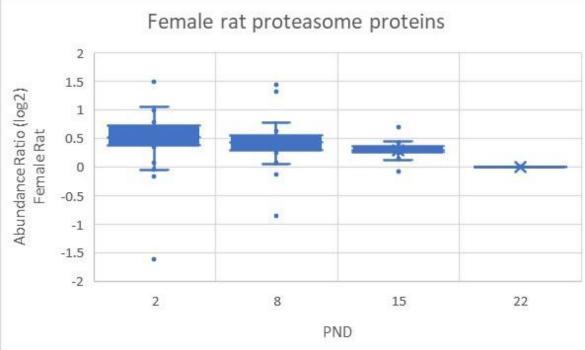
S.Fig. 4a

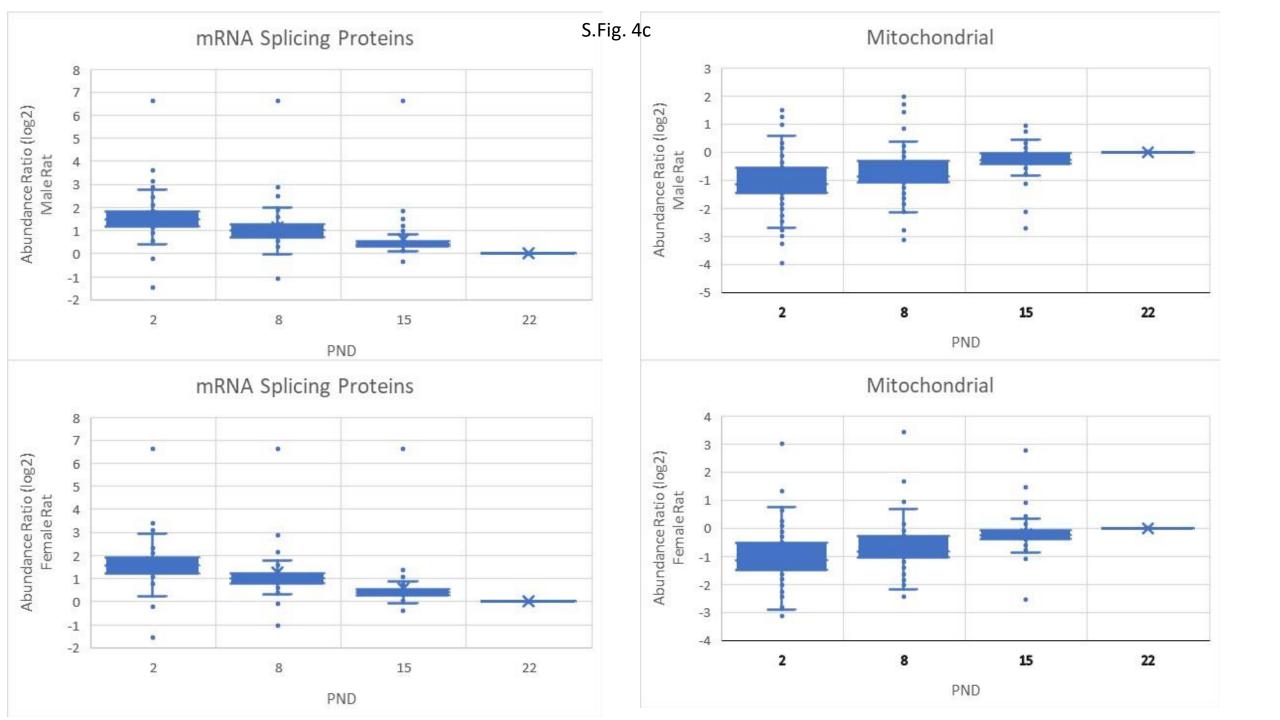


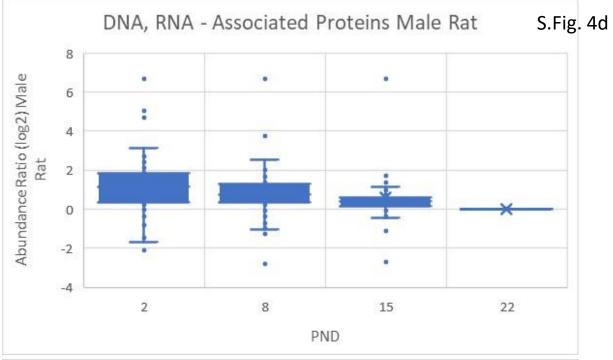


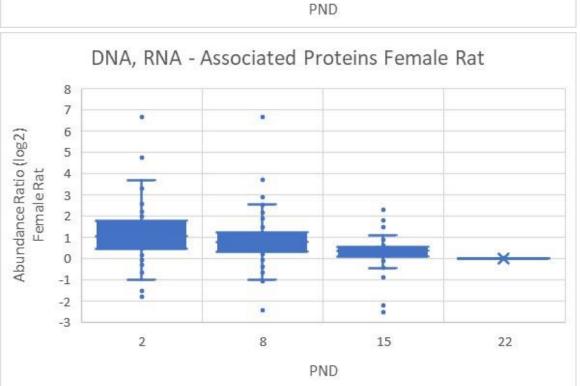
S.Fig. 4b

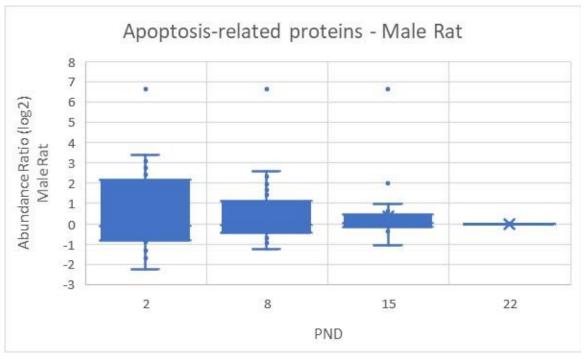


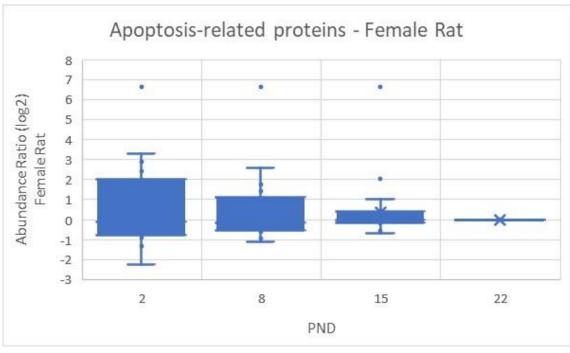


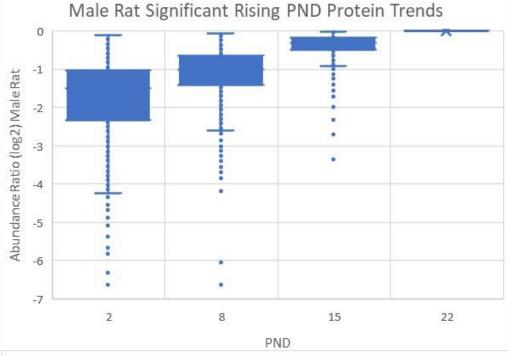


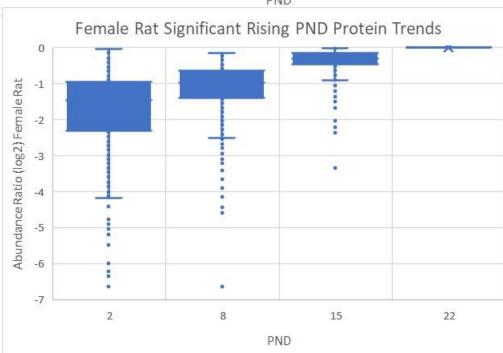


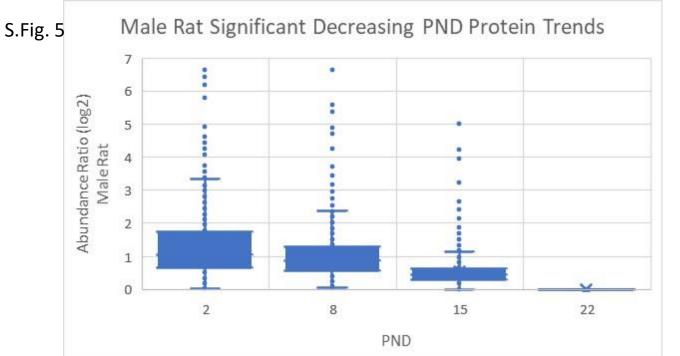


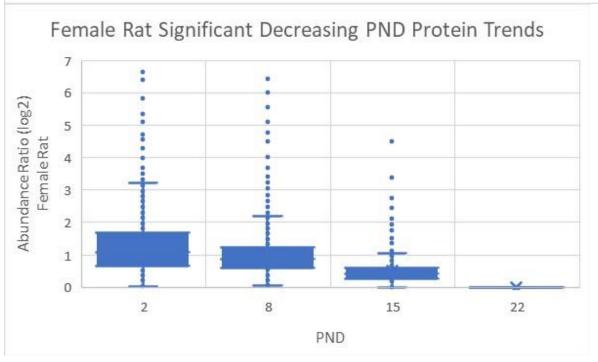












S.Fig. 6

