

## **Reducing hippocampal extracellular matrix reverses early memory deficits in a mouse model of Alzheimer's disease**

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### **Supplementary Table 2**

**Suppl. Table 2** Functional enrichment of differentially expressed proteins at 3 months of age. Significantly enriched (Fisher's exact *p*-value <0.05) GO terms are listed. Enriched GO terms were assigned to two functional groups: extracellular matrix and presynaptic neurotransmitter release

| GO term  | Proteins  | Fold enrichment | Fisher's exact <i>p</i> -value |
|--|---|-----------------|--------------------------------|
| <b>Extracellular matrix (DAVID enrichment score: 1.30)</b>                 |   |                 |                                |
| GO:0031012~extracellular matrix  | HAPLN1, SLC1A3, TNR, BCAN, NCAN   | 4.4             | 0.0098                         |
| GO:0005578~proteinaceous extracellular matrix                              | HAPLN1, SLC1A3, TNR, BCAN, NCAN   | 4.4             | 0.0098                         |
| GO:0001871~pattern binding   | NCAM1, HAPLN1, APP, BCAN, NCAN  | 3.7             | 0.0233                         |
| GO:0030247~polysaccharide binding  | NCAM1, HAPLN1, APP, BCAN, NCAN  | 3.7             | 0.0233                         |
| GO:0005539~glycosaminoglycan binding                                       | NCAM1, HAPLN1, APP, BCAN, NCAN  | 3.7             | 0.0233                         |
| GO:0005576~extracellular region  | HAPLN1, SLC1A3, PPIA, ADAM23, TNR, CFL1, BCAN, NCAN   | 2.2             | 0.0434                         |
| <b>Presynaptic neurotransmitter release (DAVID enrichment score: 0.91)</b> |   |                 |                                |
| GO:0003001~generation of a signal involved in cell-cell signaling          | RAB3A, STX1A, SYN1, SYNJ1, SYN2, CACNB4, RPH3A, SNAP25  | 2.7             | 0.0120                         |
| GO:0007269~neurotransmitter secretion                                      | RAB3A, STX1A, SYN1, SYNJ1, SYN2, RPH3A, SNAP25  | 2.8             | 0.0190                         |
| GO:0016192~vesicle-mediated transport                                      | RAB3A, STX1A, CLTB, NAPG, ICAM5, SYNJ1, NAPB, RPH3A, CLTC, STX1B, AMPH, SYP, PACSIN1, APP, ARF3, RAC1 | 1.7             | 0.0210                         |