Pumilio2 regulates synaptic plasticity via translational repression of synaptic receptors in mice

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Coronal sections of mouse brain from heterozygote *Pum2* gene trap mice *Pum2*^{XE772} was stained with X-gal. LacZ Positive cells staining blue color, indicative of *Pum2* mRNA expression, were seen in many brain regions of the mice, including the hippocampus (A–D), frontal cortex (E–G) and the amygdala (H). Nuclei stained red in the images).



Supplementary Figure 2: Western blot analysis of synaptic related protein expression in the hippocampus in the absence of Pum2. (A) PSD95, GLUR2, and phospho-CAMK2 were significantly increased in $Pum2^{-/-}$ mice. (B) Western blot signal intensity was quantified showing significant increases in $Pum2^{-/-}$ mutant tissues for PSD-95, GLUR2, and CAMK2. *means p < 0.05. Data represent as mean \pm SEM.



Supplementary Figure 3: Schematic illustration of mouse *Glur2* 3'UTR showing two putative PUM2 binding elements (red and black boxes) and their conservation in different species. (A) Schematic illustration of mouse *Glur2* 3'UTR showing two putative PUM2 binding elements (black boxes) and their conservation in different species. The numbers indicate positions of PUM2 PBE elements in the mouse *Glur2* 3'UTR. (B) Schematic representation of wildtype *Glur2* 3'UTR showing the location of PBE and mutated PBE in mutated UTR. (C) Luciferase assay performed as in B. Results of luciferase assay showing repression of PUM2 on wt 3' UTR and mutated 3'UTR of *Glur2*. (D) *p27*, as a known PUM target, serves as positive control of Luciferase assay.