

iScience, Volume 25

Supplemental information

Familial natural short sleep mutations

reduce Alzheimer pathology in mice

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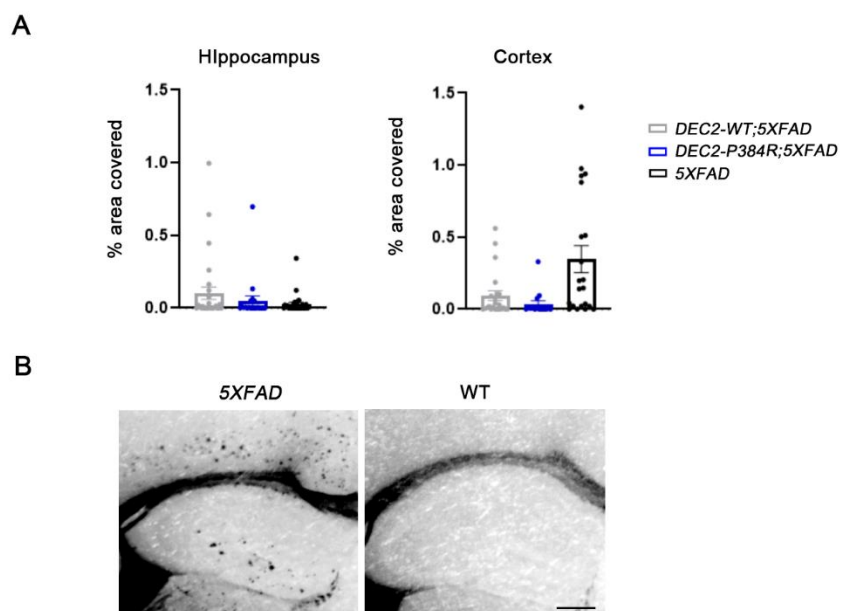


Figure S1 Plaques in 3- and 6-month-old mouse, related to Figure 1.

(A) % area of silver-stained brain slice from hippocampus and cortex covered by plaques in 3 months old mice. Hippocampus: *DEC2-WT;5XFAD*, $n = 4$ ($N = 28$ slices); *DEC2-P384R;5XFAD*, $n = 3$ ($N = 21$ slices); *5XFAD*, $n = 4$ ($N = 28$ slices); cortex: *DEC2-WT;5XFAD*, $n = 4$ ($N = 22$ slices); *DEC2-P384R;5XFAD*, $n = 3$ ($N = 15$ slices); *5XFAD*, $n = 4$ ($N = 21$ slices). (B) Representative silver staining of plaques for 6-month-old hippocampus in *5XFAD* and WT mice at high magnification. Scale bars, 500 μm . Data expressed as mean \pm SEM with individual measures displayed.

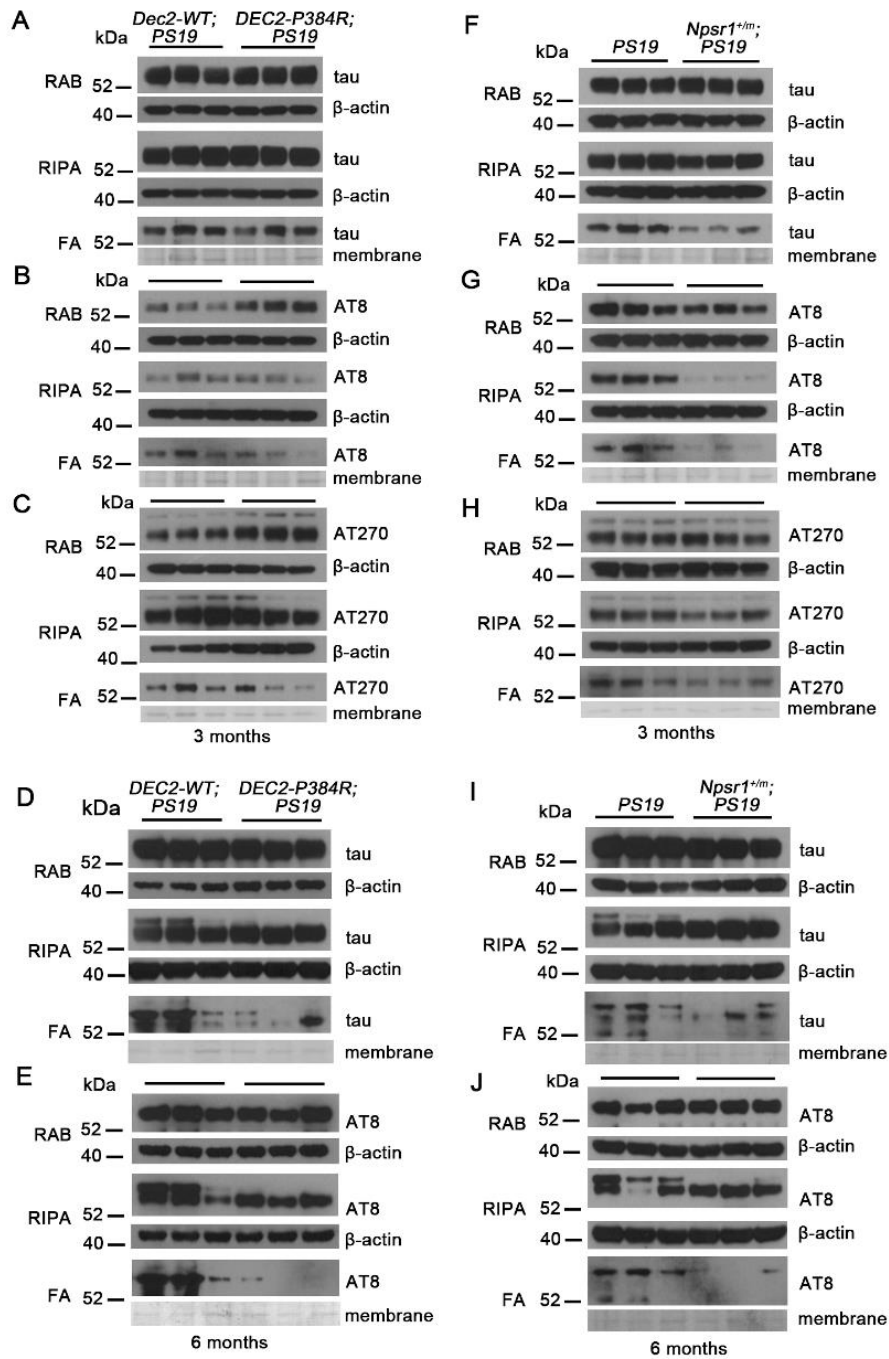


Figure S2 Representative Western blot for tau protein in *FNSS;PS19* mice, related to Figure 2.

(A-J) Representative Western blot analysis of the hippocampus of three-month-old and six-month-old mice for total tau or phosphorylated tau in soluble (RAB), less soluble (RIPA) and insoluble (FA) fractions, respectively. β -actin served as loading control for RAB and RIPA fractions. Membranes stained with Coomassie brilliant blue (CBB) served as loading control for FA fractions.

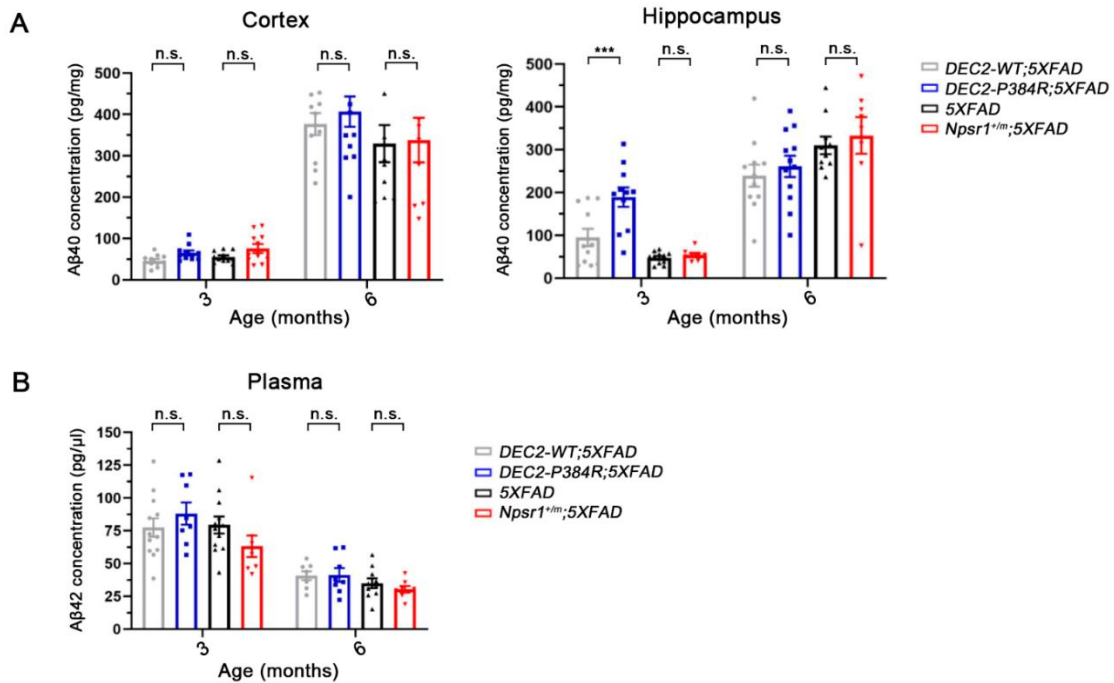


Figure S3 A β 40 and A β 42 from *FNSS:5XFAD* measured by ELISA, related to **Figure 3**.

(A) Soluble A β 40 from mouse cortex (left) of indicated genotypes measured by ELISA at 3 months of age ($n = 11$ mice per group) and 6 months of age (*DEC2-WT;5XFAD*, $n = 11$; *DEC2-P384R;5XFAD*, $n = 12$; *5XFAD*, $n = 10$; *Npsr1^{+/-};5XFAD*, $n = 9$); soluble A β 40 from hippocampus (right) at 3 months of age (*DEC2-WT;5XFAD*, $n = 11$; *DEC2-P384R;5XFAD*, $n = 11$; *5XFAD*, $n = 12$; *Npsr1^{+/-};5XFAD*, $n = 8$) and 6 months of age (*DEC2-WT;5XFAD*, $n = 11$; *DEC2-P384R;5XFAD*, $n = 12$; *5XFAD*, $n = 10$; *Npsr1^{+/-};5XFAD*, $n = 8$). (B) Plasma A β 42 from indicated genotypes measured by ELISA at 3 months of age (*DEC2-WT;5XFAD*, $n = 12$; *DEC2-P384R;5XFAD*, $n = 8$; *5XFAD*, $n = 12$; *Npsr1^{+/-};5XFAD*, $n = 8$) and 6 months of age (*DEC2-WT;5XFAD*, $n = 8$; *DEC2-P384R;5XFAD*, $n = 8$; *5XFAD*, $n = 10$; *Npsr1^{+/-};5XFAD*, $n = 8$). Data expressed as mean \pm SEM with individual measures displayed. Statistical analysis performed through one-way ANOVA with post-hoc Tukey's multiple comparisons test between relevant conditions. *** $p < 0.001$; n.s., not significant.

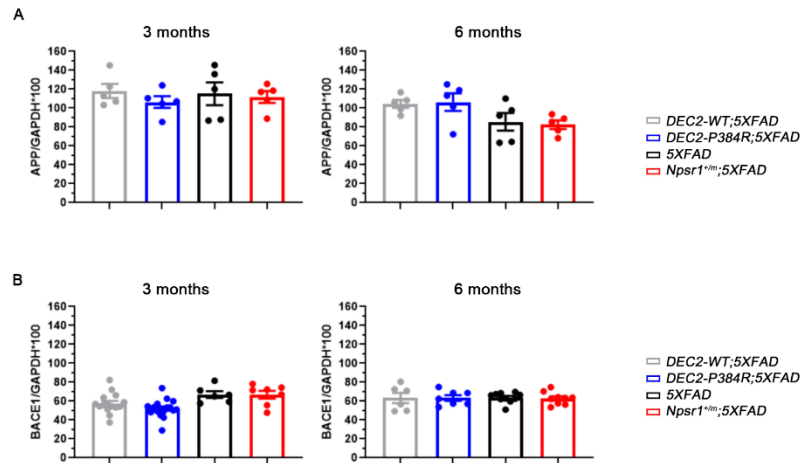


Figure S4 Quantitative results from Western blots for APP and BACE1, related to Figure 3.

(A) APP levels at 3 and 6 months of age ($n = 5$ mice per group). (B) BACE1 levels at 3 months of age (*DEC2-WT*;5XFAD, $n = 15$; *DEC2-P384R*;5XFAD, $n = 17$; 5XFAD, $n = 6$; *Npsr1*^{+/*m*};5XFAD, $n = 7$) and 6 months of age (*DEC2-WT*;5XFAD, $n = 6$; *DEC2-P384R*;5XFAD, $n = 7$; 5XFAD, $n = 10$; *Npsr1*^{+/*m*};5XFAD, $n = 10$). Data expressed as mean \pm SEM with individual measures displayed.

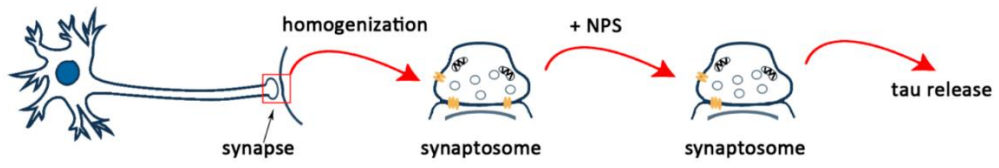


Figure S5 Schematic for synaptosome preparation and treatment with NPS, related to Figure 4.