## Supplementary Information for

GSAP modulates γ-secretase specificity by inducing conformational change in PS1

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## This PDF file includes:

Figs. S1-S3

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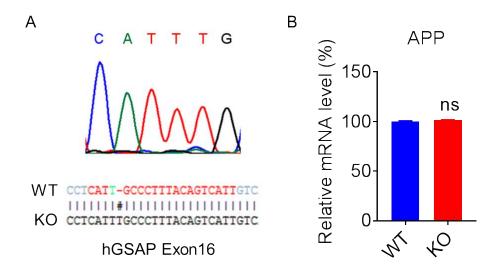


Figure S1. **HEK-APP GSAP KO conformation and APP expression.** CRISPR gRNA targeting exon 16 (A), sequence chromatogram of HEK-APP GSAP KO in the targeted region (upper panel) and sequence alignment of HEK-APP WT versus KO in exon16 (lower panel, CRISPR gDNA in red and last PAM nucleotide in green). APP mRNA levels in HEK-APP WT versus HEK-APP GSAP KO cells (B).

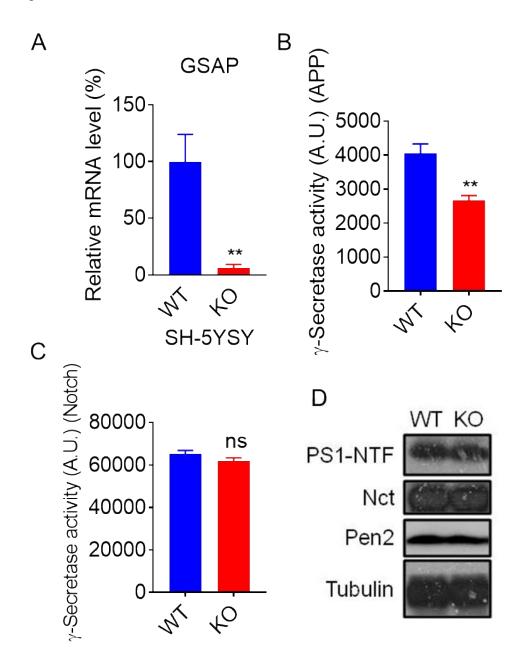


Figure S2. **SH-5YSY GSAP KO reduces** γ-secretase activity for **APP.** GSAP mRNA levels in SH-5YSY cells versus CRISPR GSAP KO cells (A). γ-Secretase activity levels toward recombinant APP (B) are decreased in GSAP KO SH-5YSY compared to GSAP WT counterpart but remained unchanged for recombinant Notch (C). All data represent means  $\pm$ SEM; n=6. (D) Western blot analysis of γ-secretase subunits PS1-NTF, Nct, and Pen2 in SH5-YSY WT and KO (p-value \*\*< 0.01).

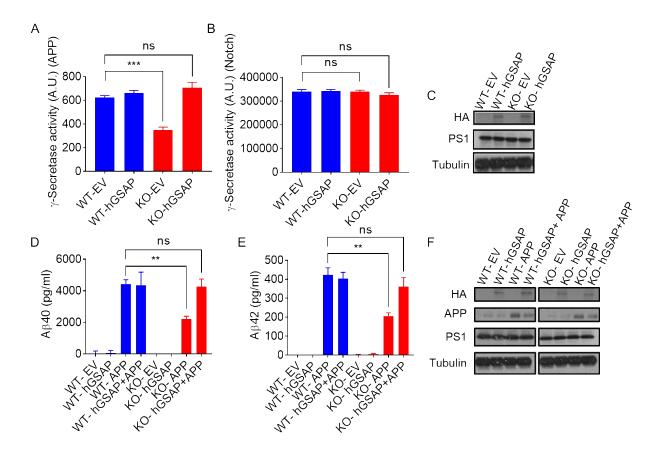


Figure S3. Overexpression of hGSAP in SH-5YSY GSAP KO cells rescues  $\gamma$ -secretase activity and A $\beta$  secretion. Membrane fractions prepared 48 hours post transfection of EV and hGSAP in SH-5YSY GSAP WT or KO were assayed for  $\gamma$ -secretase activity using recombinant APP (A) or Notch (B). Western blot analysis of HA (hGSAP) and PS1-NTF in SH-5YSY WT and KO confirming GSAP expression which had no effect on PS1 (C). A $\beta$  analysis was done on SH-5YSY WT or KO transfected with Empty vector (EV) or full-length human GSAP (hGSAP). To detect A $\beta$  levels the cells were transfected either with APP or hGSAP+APP. A $\beta$  secretion was measured 48 hours post-transfection using MSD A $\beta$  detection kit; A $\beta$ 40 (D) and A $\beta$ 42 (E). All data represent means  $\pm$ SEM; n=3. (F) Western blot analysis of HA (hGSAP), APP and PS1-NTF in SH-5YSY WT and KO transfected with: EV, hGSAP, APP or hGSAP+APP confirming the expression of hGSAP or APP in the indicated cells (p-value \*\* < 0.01; \*\*\* < 0.001).