

Supplementary Information

Methods

Materials: 2,5-dioxopyrrolidin-1-yl 4-azido-2-hydroxybenzoate (NHS-ASA) was purchased from ProChem. Inc (Rockford, IL). 6-Methyl-*N*¹-(4-(pyridin-3-yl)pyrimidin-2-yl)benzene-1,3-diamine and *N*-(4-methyl-3-(4-(pyridin-3-yl)pyrimidin-2-ylamino)phenyl)-4-(piperazin-1-ylmethyl)benzamide (*N*-desmethyl imatinib) were purchased from ChemPacific Inc (Baltimore, MD). 2,5-dioxopyrrolidin-1-yl 5-((3*a*S,4*S*,6*a*R)-2-oxo-hexahydro-1*H*-thieno[3,4-*d*]imidazol-4-yl)pentanoate (Biotin-OSu), *N*-(chloro(dimethylamino)methylene)-*N*-methylmethanaminium hexafluorophosphate (TCFH), trifluoroacetic acid (TFA), 1*H*-benzo[*d*][1,2,3]triazol-1-ol (HOBt) and *N,N*-diisopropylethyl amine (DIPEA) were purchased from Sigma-Aldrich (St. Louis, MO). *Tert*-butyl 2-(piperazin-1-yl)ethylcarbamate was purchased from Astatech Inc (Bristol, PA).

Synthesis of an imatinib derived photo-affinity label, G01: DIEPA (63 μ l, 0.36 mmol) was added to a solution of NHS-ASA (50 mg, 0.18 mmol), HOBt (25 mg, 0.18 mmol), and 6-Methyl-*N*¹-(4-(pyridin-3-yl)pyrimidin-2-yl)benzene-1,3-diamine (50 mg, 0.18 mmol) in DMF (2 ml). The reaction mixture was stirred at room temperature overnight under argon atmosphere. The generated crude product was purified by a semi-preparative HPLC to give 54 mg of the titled compound with a yield of 68%. The product, **G01**, 4-azido-2-hydroxy-*N*-(4-methyl-3-(4-(pyridin-3-yl)pyrimidin-2-ylamino)phenyl)benzamide, was confirmed by mass spectral analysis using an ESI-MS in the positive mode $[M+H]^+$, demonstrating a m/z of 439.1.

Radioiodination of G01 by ^{125}I was performed without carrier using a modification of a Chloramine-T procedure and the iodinated product was purified by HPLC. Specifically, in a UV protected "V" vial, total volume 0.9 ml, ~ 10 mCi of ^{125}I stock isotope (volume = 25 μl) was added to 200 μl of 0.2M phosphate buffer, pH 7.2. G01 was dissolved to 1 mg/ml in ethanol and 25 μl of this solution was combined with chloramine-T at 1 mg/mL in water (50 μl) and then added to the V-vial. The reaction proceeded for 1 min and was terminated by the addition of 50 μL of 1 mg/ml meta-bisulfite. The reaction mixture was chromatographed on a 25 cm Waters RP-C18 column, using 0.1% TFA in water as the "A" solvent and 0.1% TFA in acetonitrile as the "B" solvent. A gradient was run at 1 ml/min from 0% B to 50% B for 45 minutes and held at 50% B for 15 minutes. The product demonstrated a retention time of 54.5 min as followed by radiochemical detection, and had a specific activity of 2000 Curies per millimole. The I^{125} labeling experiment was performed by PerkinElmer Life and Analytical Sciences, Inc.

$^3\text{H-G01}$ was prepared by ViTrax Radiochemicals via catalytic tritium exchange of G01. The labeled product was purified by HPLC. The composition of the purified product was verified by co-injection of the tritium labeled product with its cold precursor and both compounds co-chromatographed on an analytical HPLC.

Cellular A β production assays and incubation with G01.

Neuroblastoma 2a cells stably overexpressing human APP695 were treated with 10 μM G01 for 3 hr. Cells treated with DMSO, or DMSO plus imatinib, were used as controls. After 3 hr, conditioned medium was collected and A β immunoprecipitation was conducted using 4G8 antibody. The immunoprecipitated A β was separated on 10-20% Tris-tricine gel, transferred to PVDF membrane and detected by 6E10 antibody.

Synthesis and kinase profiling of biotin-imatinib (active and inactive form): Inactive biotin-imatinib, (IC200001) was synthesized by reacting N-desmethyl imatinib with Biotin-OSu. **Active biotin-imatinib, (IC339239)** was synthesized from the key intermediates, tert-butyl 2-(piperazin-1-yl)ethylcarbamate and 6-methyl-*N*¹-(4-(pyridin-3-yl)pyrimidin-2-yl)benzene-1,3-diamine, via 4 steps, as shown in supplementary figure 1.

The kinase profiling was performed by Millipore Inc. using the standard assays for Abl kinase and PDGF receptor (ATP = 45 μ M). Compound IC200001 showed no significant inhibitory activity toward either kinase, while compound IC339239 had an IC₅₀ of 146 nM against Abl kinase (imatinib had an IC₅₀ of 79 nM) and an IC₅₀ of 6.6 μ M against PDGF receptor (imatinib had an IC₅₀ of 4.8 μ M). Thus, we refer to IC200001 as “inactive biotin-imatinib” and IC339239 as “active biotin-imatinib”.

Construction of APP-CTF truncated forms

APP-CTF-T1 is the truncated form of APP- β -CTF spanning from its N-terminus to HHGV⁶⁴; APP-CTF-T2 is the truncated form of APP- β -CTF spanning from its N-terminus to VMLKK⁵⁵. Both truncated forms were generated by introducing a stop codon in related positions of the CT100 (APP-CTF) in pcDNA4. Mutagenesis was performed using QuickChange Site-Directed Mutagenesis kit (Stratagene) according to the manufacturer's instructions. The primers used for APP-CTF-T1 are: Forward: 5' CATTTCATCATGGTGTGTAGGAGGTTGACGCCGC 3'. Reverse: 5' GCGGCGTCAACCTCCTACACACCATGATGAATG 3'. The primers used for APP-CTF-T2 are: Forward: 5' CTTGGTGATGCTGAAGAAGTAACAGTACACATCCATTC 3' Reverse: 5'

GAATGGATGTGTACTGTTACTTCTTCAGCATCACCAAG 3'. The presence of the stop codon and integrity of the cDNA were verified by sequencing.

Construction, expression, and analysis of APP/NICD and Notch Δ E/AICD

APP/NICD and Notch Δ E coding sequences were synthesized by Genscript Inc. and are illustrated in supplementary Fig. 10. Both sequences were then incorporated into a pcDNA3.1 vector coding for a C-terminal Myc tag. APP-CTF/NICD, APP-CTF, and Notch Δ E/AICD were overexpressed separately in HEK293 cells. Immunoprecipitation was conducted using gSAP antibody. APP/NICD and Notch Δ E/AICD were detected by c-myc antibody. APP-CTF was detected by 369 antibody.

APP-CTF/NICD, APP-CTF, and Notch Δ E/AICD were transfected into N2a cells with/without gSAP siRNA knockdown. NICD was detected by both myc antibody and cleavage specific Val1844 antibody (Cell Signaling); AICD was detected by myc antibody. AICD production from APP-CTF was detected by 369 antibody.

Mouse administration of a γ -secretase inhibitor dibenzazepine (DBZ)

gSAP RNAi mice (6 months old) were administered dibenzazepine (DBZ) (10 μ mol/kg) once daily for 5 days by intraperitoneal injection. DBZ was suspended finely in 0.5% (w/v) hydroxypropylmethylcellulose and 0.1% (w/v) Tween 80. Mice were treated with DBZ or with vehicle, with 4 mice in each group. After sacrificing, mouse brain was removed for A β ELISA assays (Invitrogen); Mouse intestine was processed for H&E and PAS staining.

Supplementary Figure Legends

Supplementary Figure 1. gSAP action on APP processing. Ternary complex of gSAP, APP and γ -secretase (top) is associated with elevated γ -cleavage ($A\beta$ production) and reduced ϵ -cleavage (AICD production). In the absence of gSAP (bottom), the binary complex of APP and γ -secretase is associated with decreased γ -cleavage and increased ϵ -cleavage.

Supplementary Figure 2. a: structures of imatinib, G01, and ^{125}I -G01. **b:** G01 significantly reduces levels of $A\beta$ in N2a cells.

Supplementary Figure 3. a: Procedures for the synthesis of biotinylated imatinib derivative, IC339239: reagents and conditions: **(a)** 4-(bromomethyl) benzoic acid, K_2CO_3 , DMF, room temperature, 2 h. **(b)** 6-methyl- N^1 -(4-(pyridin-3-yl)pyrimidin-2-yl)benzene-1,3-diamine, TCFH, DIPEA, DMF, room temperature, overnight. **(c)** TFA, CH_2Cl_2 , room temperature, 30 min. **(d)** Biotin-OSu, HOBt, DIPEA, room temperature, overnight, and then HPLC purification. Compound IC200001 was synthesized by reacting N-desmethyl imatinib with Biotin-OSu.

b: Kinase profiling results shows that IC339239 has activities comparable to those of imatinib, while IC200001 showed no activity. Therefore, IC339239 is designated as “active biotin-imatinib” and IC200001 as “inactive biotin-imatinib”.

Supplementary Figure 4. Alignments of gSAP sequences among species. Red: identical residues; Blue/Green: conservative substitutions. The C-terminal region of gSAP is highly conserved. The gSAP-16K region is underlined.

Supplementary Figure 5. gSAP mRNA expression levels in different tissues. Tissues from 3 month old wild type BL/6 mice were harvested and gSAP levels were quantitated

using real time PCR. Both actin and GAPDH served as internal controls (n=6). Tissue extracts were adjusted to the same protein levels prior to analysis.

Supplementary Figure 6. gSAP knockdown does not influence the generation of α -, or β -APP-CTF from full length APP. N2a cells overexpressing APP695 were pre-treated with the γ -secretase inhibitor, L685,458 [1 μ M] and the cleavages of APP were monitored by pulse-chase labeling (35 S-methionine) followed by immunoprecipitation. Proteins were separated by SDS-PAGE and transferred to PVDF membrane for autoradiography.

Supplementary Figure 7. Regulation of AICD production by gSAP. **a.** Either gSAP knockdown or imatinib treatment increases AICD levels in N2a cells overexpressing APP695 (**p < 0.01; n=3). **b.** Transfection of gSAP into HEK293 cells overexpressing APP-CTF led to reduced AICD production and increased A β production. Cleavage of APP-CTF was monitored by pulse-chase labeling at indicated time point. AICD and A β levels after the 3 hr chase were quantitated (**p < 0.01; n=3).

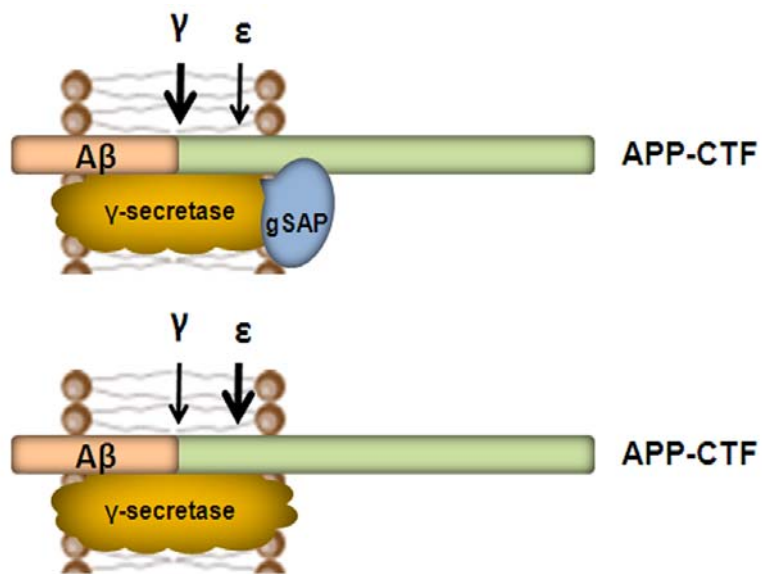
Supplementary Figure 8. After separation of organelles from N2a cells on a continuous sucrose gradient, endogenous gSAP co-localizes with γ -secretase in a Golgi-enriched fraction (No. 6), which also contains endosomes.

Supplementary Figure 9. Truncation of APP-CTF and immunoprecipitation using gSAP antibody through gSAP demonstrates that gSAP interacts with the juxtamenbrane region of APP-CTF. APP-CTF-T1 is the truncated form of APP- β -CTF spanning from its N-terminus to HHGV⁶⁴; APP-CTF-T2 is the truncated form of APP- β -CTF spanning from its N-terminus to VMLKK⁵⁵. Truncated forms were overexpressed in HEK293 cells and immunoprecipitated with gSAP antibody. 6E10 antibody was used for immuno-detection.

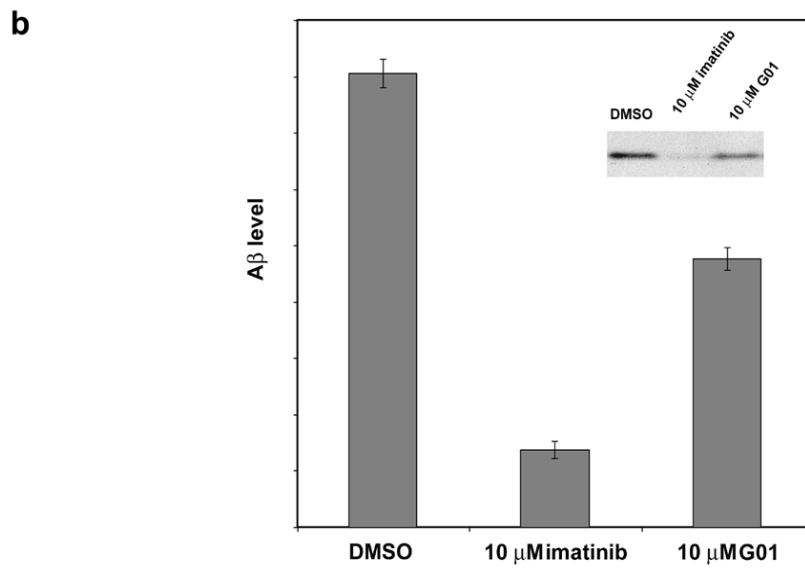
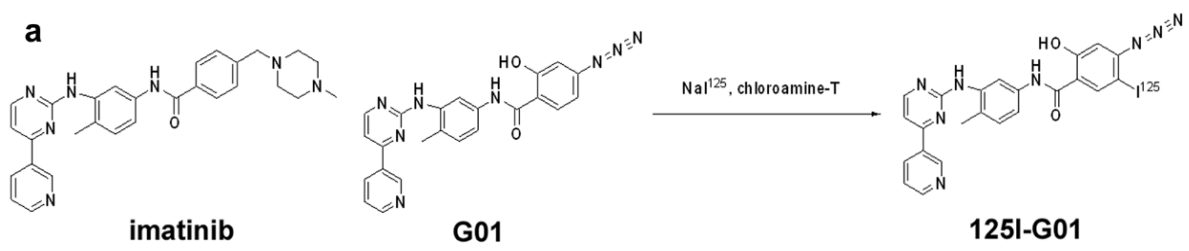
Supplementary Figure 10. Domain exchange studies demonstrate that gSAP regulates ϵ -cleavage of APP-CTF but not Notch Δ E, through selective interaction with AICD. **a.** Design of APP-CTF/NICD and Notch Δ E/AICD constructs. **b.** APP-CTF/NICD, APP-CTF, and Notch Δ E/AICD were overexpressed separately in HEK293 cells. Both APP-CTF and Notch Δ E/AICD interact with gSAP, while APP-CTF/NICD does not. **c.** Effects of gSAP knockdown on the cleavage of APP-CTF/NICD, APP-CTF, and Notch Δ E/AICD by γ -secretase. Individual constructs were overexpressed in N2a cells. Upon gSAP knockdown, AICD production from APP-CTF (center panel) as well as from Notch Δ E/AICD (right panel) increased, but NICD production from APP-CTF/NICD was not influenced (left panel).

Supplementary Figure 11. Comparison of the effects of either reducing gSAP or of a γ -secretase inhibitor on A β levels and histopathology. **a.** Mice were given a γ -secretase inhibitor DBZ (10 μ mol/kg) for 5 days. This resulted in reduced A β 40 and A β 42 levels of 44 \pm 9% and 47 \pm 5% respectively. **b.** No histopathological changes were observed in mouse small intestine after gSAP knockdown (H&E and PAS staining). However, increased amounts of violet-stained goblet cells were observed after DBZ administration, a finding typical of Notch inhibition.

Supplementary Figure 1

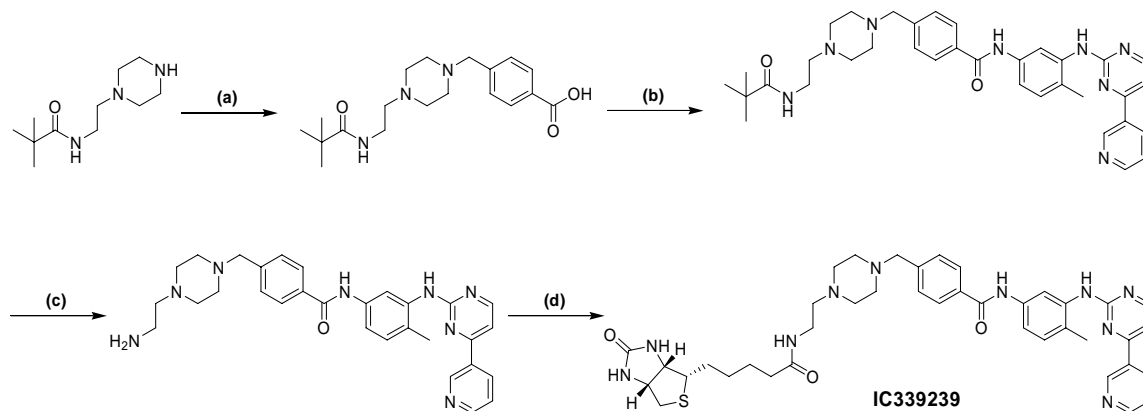


Supplementary Figure 2

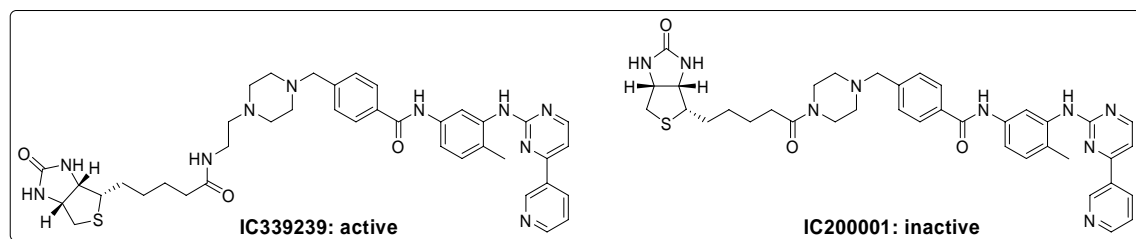
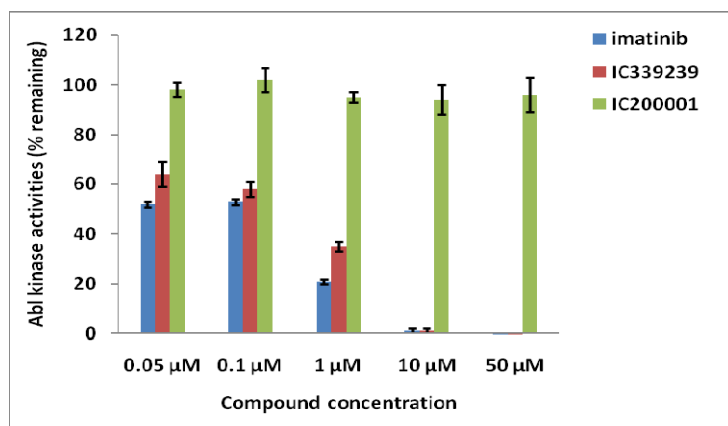


Supplementary Figure 3

a



b



Supplementary Figure 4

Homo	-----MALRLVADFDL GKDVLPWLR A
Canis	-----MTQNLSWPGH
Bos	-----MALRLIADFDLEKDVLPWLRV
Mus	-----MALRLVTHFDVLEDEVLP SLLT
Rattus	-----MALRLVTHFDVLADVLP SLLV
Gallus	MAVAAPQQPARCGGQRPECGRVGPRRLRALPSGGRRSQAGRESPRAAHGAASPLP LPSGPG
Homo	QRAVSEASGAGSG-----GADVLENDYES--
Canis	SSNGSCIRFPVAG-----GTAVL-----
Bos	QLAASAAAGARGG-----GPGVLENNYEC--
Mus	QAATTEDEGDRAGV-----LETTYG----S--
Rattus	QAATADEGDEGA-----ETTLG----S--
Gallus	RLEATGGRGNGGASGRPQLRGLSPPAPLPCGGCAGPELRGLTSLCGGSALDTSEKSSA
Homo	LHVLNVERNGNI IYTYKDDKGNVVFGLYDCQTRQNELLYTFEKDLQVFSVNSERTLLA
Canis	-----WQGAVSSIQGLGTADHELPTWRAYEQLPYADLA
Bos	LRVLNVERNRI IYTYKDNKGNVVFGLYDYQTKQNEHLYTFEKDLQVSSVNSKEKTLA
Mus	LRVLNIERNNGNI IYTYKDNKGNVAVFGLYDCQTRQNEHLYTFEKDMQAVSSVNSERTVLA
Rattus	LRVLNIERNGDI IYTYKDNKGNVAVFGLYDCQTRQNEHLYTFEKDMQAVSSVNSERTVLA
Gallus	LYIVNVERNGKI IYTWKGNQRSTHIGLYDLQTKENEHLYTFEKDLRIISVNSERTLLA
Homo	ASLVQSTK-EGKRNELQPGSKCLTLLVEIHPVNNVVKLVKAVDSYI WVQFLYPHIESHPLP
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Bos	TSLVQAAK-EGRSNELQPGSKCLTLLVEIHPINNPKLVKAVDSYI WVQFLYPHVESCPQP
Mus	ASFIQYTT-EGVKNDLQPGSKCLTLLVEIHPVNNVVKLVKAVDSCVWVQFLYPQAESHLLP
Rattus	ASFIQYT--EGVRSELQPGSKCLTLLVEIHPVNNVTVLKAVDSCVWVQFLYPQAESHLLA
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Gallus	ESHLLLVSEDKYIEQFDIHVAEEE-HRVVIQNSGQLPRARVADDLIWAQWDMTEQRIFYI
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Rattus	ELQESRSILKCVQFWADESFTIMFEMPLDISLSGLRFKLVNFGYDYRQDQAKLCHQPSLC
Gallus	VPKESRSILRCVQFYPDENFNSTLESQLDISVNDKRVKLVNFGYNDCEDRDVPPKSLNLQ
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Canis	VFTNHTGSLCVCYSPKFDSEKITYSVFYFHKGHSKFTTAALGSVDSLVTKG-----
Bos	VFTNHTGSLCVCYCPNFDSWEQITYSVFYFHKGHSKFTTTLGSVDSHVTKG-----
Mus	IFTNHTGSLCMCYSKSDSREEITYSVFYLHKGYRKIFTAAPGSADSQVTNGADSQVTDG
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Gallus	VFTNKAG-----FSKFTTASLERPETPQLKE-----
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Mus	IAFLNLDGYFVAVYSPGHFLHLLNIQHPDLVCHSLFLTGNNKIAAVLPPSPLQSLPGSLVL
Rattus	VTFLNLDGYFVAVYSPCRFLHLLNIRHPDLICHSLFLTGNNKTAAVLPPSPLQSLPGSLIL

Gallus VAFNLNDYYVAAYLPGQFLHLLNIQHPDLLCYSLFLTGEDARIDMLPNCISIQSPLVSTVL

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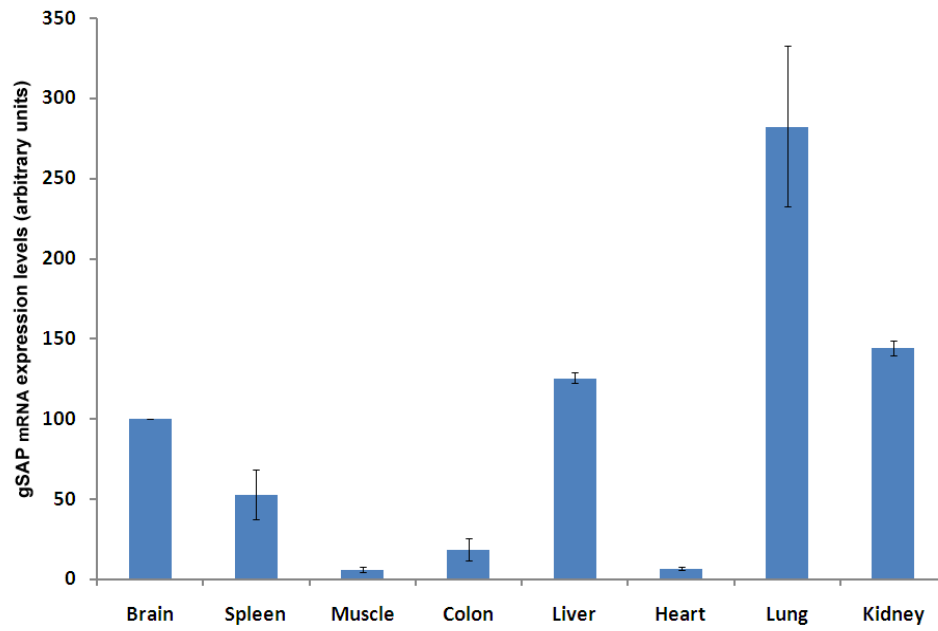
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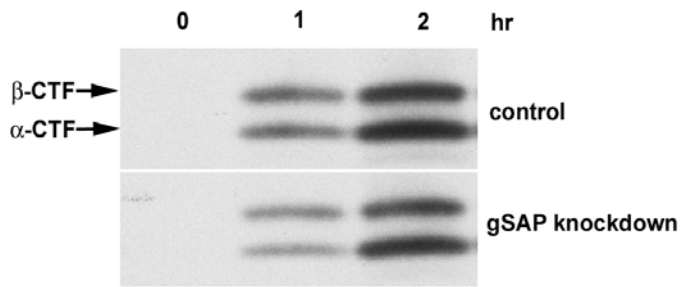
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Supplementary Figure 5.

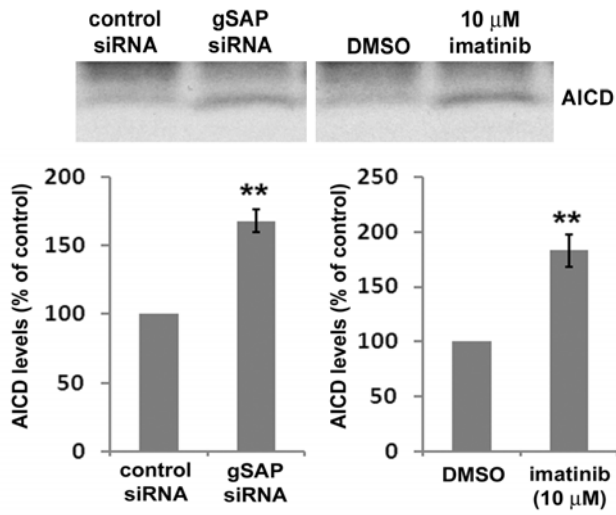


Supplementary Figure 6

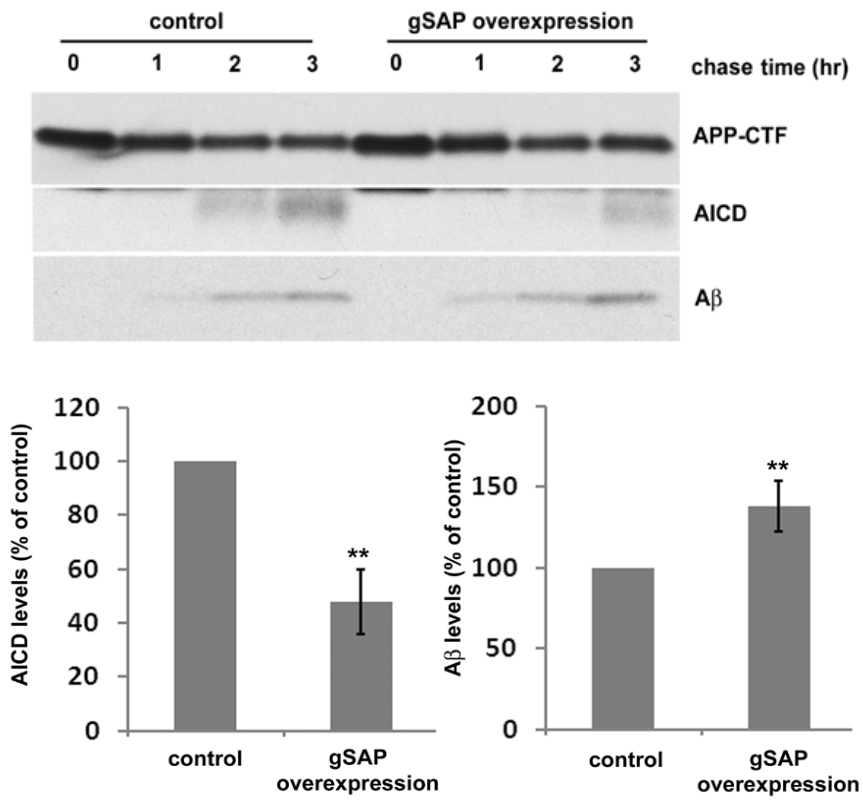


Supplementary Figure 7

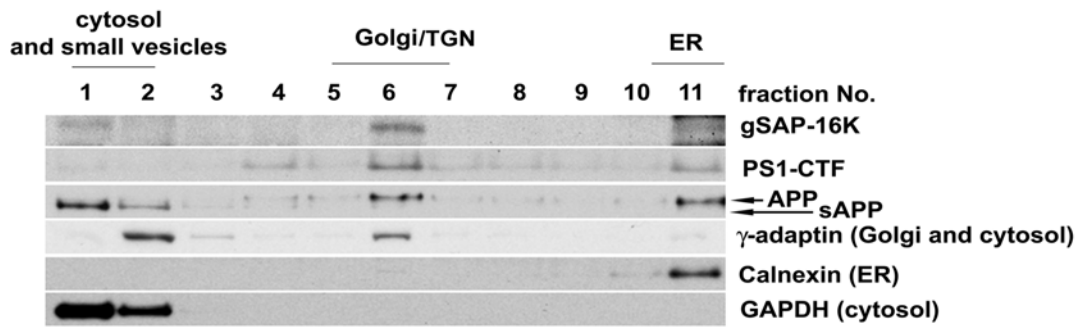
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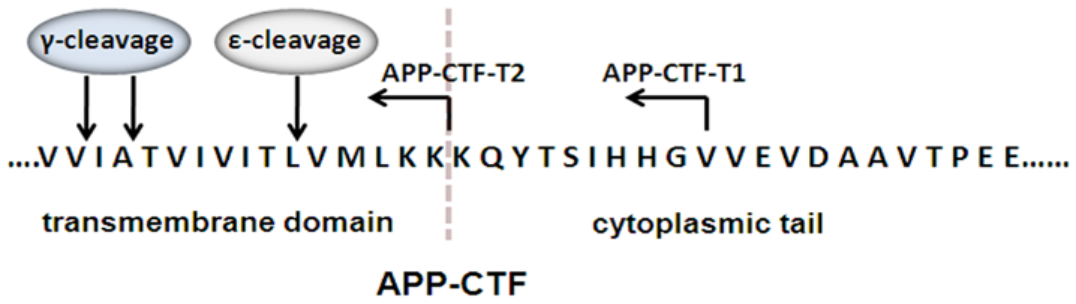
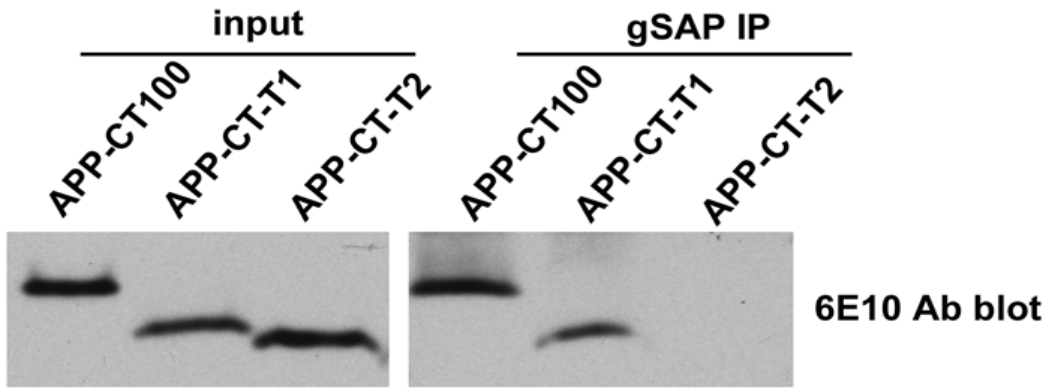
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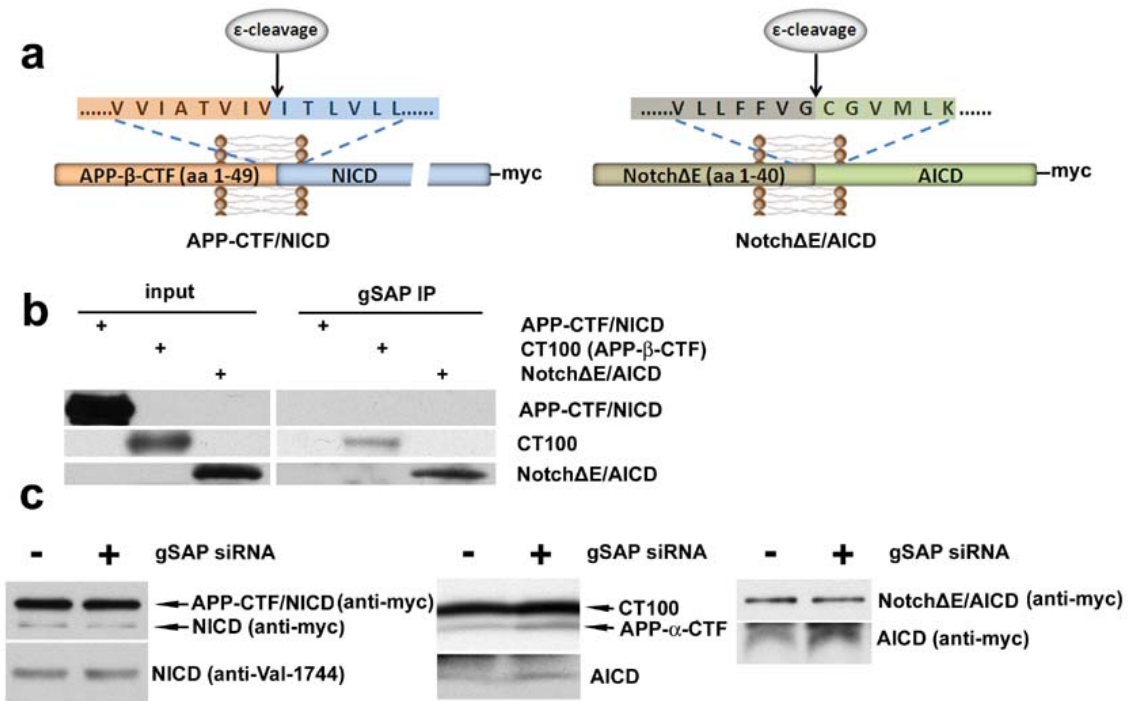
Supplementary Figure 8



Supplementary Figure 9

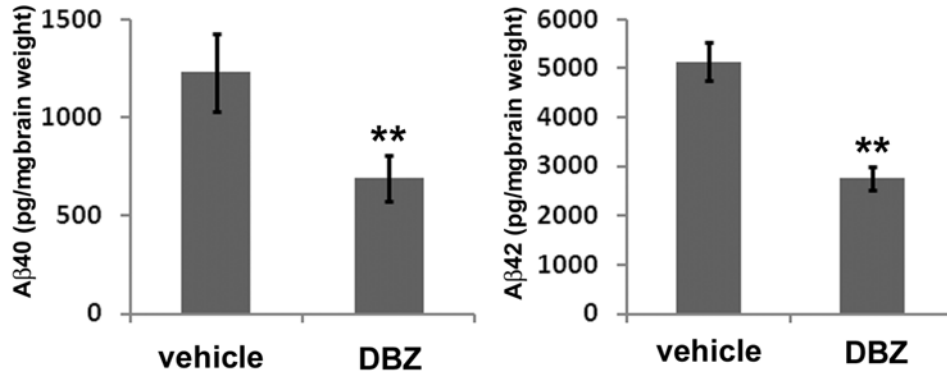


Supplementary Figure 10



Supplementary Figure 11

a



b

