

BACE1 Antibody

Rabbit mAb Catalog # AP91310

Product Information

Application WB, IP **Primary Accession** P56817

Reactivity Rat, Human, Mouse

Clonality Monoclonal

Other Names Beta-secretase 1; Aspartyl protease 2; ASP2; Asp 2; Memapsin-2; BACE1; BACE;

IsotypeRabbit IgGHostRabbitCalculated MW55764

Additional Information

Dilution WB 1:500~1:2000 IP 1:40 **Purification** Affinity-chromatography

Immunogen A synthesized peptide derived from human BACE1

Description Responsible for the proteolytic processing of the amyloid precursor protein

(APP). Cleaves at the N-terminus of the A-beta peptide sequence, between residues 671 and 672 of APP, leads to the generation and extracellular release of beta-cleaved soluble APP, and a corresponding cell-associated C-terminal

fragment which is later released by gamma-secretase.

Storage Condition and Buffer Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% sodium

azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term.

Avoid freeze / thaw cycle.

Protein Information

Name BACE1 (HGNC:933)

Synonyms BACE, KIAA1149

Function Responsible for the proteolytic processing of the amyloid precursor protein

(APP). Cleaves at the N-terminus of the A-beta peptide sequence, between residues 671 and 672 of APP, leads to the generation and extracellular release of beta-cleaved soluble APP, and a corresponding cell-associated C-terminal fragment which is later released by gamma-secretase (PubMed:10656250, PubMed:10677483, PubMed:20354142). Cleaves CHL1 (By similarity).

Cellular Location Cell membrane; Single-pass type I membrane protein Golgi apparatus,

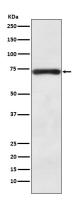
trans-Golgi network. Endoplasmic reticulum. Endosome. Cell surface. Cytoplasmic vesicle membrane; Single-pass type I membrane protein. Membrane raft {ECO:0000250|UniProtKB:P56818}. Lysosome. Late endosome. Early endosome. Recycling endosome. Cell projection, axon

{ECO:0000250 | UniProtKB:P56818}. Cell projection, dendrite {ECO:0000250 | UniProtKB:P56818}. Note=Predominantly localized to the later Golgi/trans-Golgi network (TGN) and minimally detectable in the early Golgi compartments. A small portion is also found in the endoplasmic reticulum, endosomes and on the cell surface (PubMed:11466313, PubMed:17425515). Colocalization with APP in early endosomes is due to addition of bisecting N-acetylglucosamine which blocks targeting to late endosomes and lysosomes (By similarity) Retrogradly transported from endosomal compartments to the trans-Golgi network in a phosphorylation- and GGA1- dependent manner (PubMed:15886016). {ECO:0000250 | UniProtKB:P56818, ECO:0000269 | PubMed:11466313, ECO:0000269 | PubMed:15886016, ECO:0000269 | PubMed:17425515}

Tissue Location

Expressed at high levels in the brain and pancreas. In the brain, expression is highest in the substantia nigra, locus coruleus and medulla oblongata.

Images



Western blot analysis of BACE1 expression in SH-SY5Y cell lysate.

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