

**BACE2/Asp1 Antibody**  
**Rabbit Polyclonal Antibody**  
**Catalog # ABV11100****Specification**

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**BACE2/Asp1 Antibody - Product Information**

|                   |                        |
|-------------------|------------------------|
| Application       | WB                     |
| Primary Accession | <a href="#">O9Y5Z0</a> |
| Reactivity        | Human, Mouse, Rat      |
| Host              | Rabbit                 |
| Clonality         | Polyclonal             |
| Isotype           | Rabbit IgG1            |
| Calculated MW     | 56180                  |

**BACE2/Asp1 Antibody - Additional Information****Gene ID** 25825**Application & Usage****Western Blot: 1-4 µg/ml. However, the optimal conditions should be determined individually.****Other Names**

BACE2, Asp1

**Target/Specificity**

BACE2

**Antibody Form**

Liquid

**Appearance**

Colorless liquid

**Formulation**

100 µg (0.5 mg/ml) affinity purified rabbit anti-BACE2 polyclonal antibody in phosphate buffered saline (PBS), pH 7.2, containing 30% glycerol, 0.5% BSA, 0.01% thimerosal.

**Handling**

The antibody solution should be gently mixed before use.

**Reconstitution & Storage**

-20 °C

**Background Descriptions****Precautions**

BACE2/Asp1 Antibody is for research use only and not for use in diagnostic or therapeutic procedures.

## **BACE2/Asp1 Antibody - Protein Information**

**Name** BACE2

**Synonyms** AEPLC, ALP56, ASP21

### **Function**

Responsible for the proteolytic processing of the amyloid precursor protein (APP). Cleaves APP, between residues 690 and 691, leading 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. It has also been shown that it can cleave APP between residues 671 and 672 (PubMed:<a href="http://www.uniprot.org/citations/10591213" target="\_blank">10591213</a>, PubMed:<a href="http://www.uniprot.org/citations/11083922" target="\_blank">11083922</a>, PubMed:<a href="http://www.uniprot.org/citations/11423558" target="\_blank">11423558</a>, PubMed:<a href="http://www.uniprot.org/citations/15857888" target="\_blank">15857888</a>, PubMed:<a href="http://www.uniprot.org/citations/16816112" target="\_blank">16816112</a>). Involved in the proteolytic shedding of PMEL at early stages of melanosome biogenesis. Cleaves PMEL within the M-beta fragment to release the amyloidogenic PMEL luminal fragment containing M-alpha and a small portion of M-beta N-terminus. This is a prerequisite step for subsequent processing and assembly of PMEL fibrils into amyloid sheets (PubMed:<a href="http://www.uniprot.org/citations/23754390" target="\_blank">23754390</a>). Responsible also for the proteolytic processing of CLTRN in pancreatic beta cells (PubMed:<a href="http://www.uniprot.org/citations/21907142" target="\_blank">21907142</a>).

### **Cellular Location**

Cell membrane; Single-pass type I membrane protein. Golgi apparatus. Endoplasmic reticulum. Endosome Melanosome. Note=Colocalizes with PMEL in stage I and II melanosomes.

### **Tissue Location**

Brain. Present in neurons within the hippocampus, frontal cortex and temporal cortex (at protein level). Expressed at low levels in most peripheral tissues and at higher levels in colon, kidney, pancreas, placenta, prostate, stomach and trachea. Expressed at low levels in the brain. Found in spinal cord, medulla oblongata, substantia nigra and locus coeruleus. Expressed in the ductal epithelium of both normal and malignant prostate.

## **BACE2/Asp1 Antibody - Protocols**

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

## **BACE2/Asp1 Antibody - Images**

## **BACE2/Asp1 Antibody - Background**

Accumulation of the amyloid-beta (A $\beta$ ) plaque in the cerebral cortex is a critical event in the pathogenesis of Alzheimer's disease. A $\beta$  peptide is generated by proteolytic cleavage of the

beta-amyloid protein precursor (APP) at beta- and gamma-sites by proteases. The long-sought beta-secretase was recently identified by several groups independently and designated beta-site APP cleaving enzyme (BACE) and aspartyl protease 2 (Asp2). A BACE homolog was recently cloned and designated BACE2, Asp1, DRAP (for Down region aspartic protease), and memapsin 1. BACE2 also cleaves APP at beta-site and at a different site within Abeta. BACE2 locates on chromosome 21q22.3, the so-called 'Down critical region', suggesting that BACE2 and Abeta may also contribute to the pathogenesis of Down syndrome.