# SANTA CRUZ BIOTECHNOLOGY, INC.

# BACE (C-15): sc-10055



## BACKGROUND

Autosomal dominant Alzheimer's disease is caused by mutations in the gene encoding the  $\beta$ -amyloid protein precursor (APP). Amyloid  $\beta$ -peptide (A $\beta$ ), the major feature of amyloid plaques in Alzheimer's patients, is the product of APP cleavage by  $\beta$ - and  $\gamma$ -secretases. BACE is the transmembrane protease which cleaves A $\beta$  from APP. BACE and the related protein Asp1 are both widely expressed in human tissue with the highest levels in the pancreas. BACE is localized within Golgi and endosomes

#### REFERENCES

- 1. Kang, J., et al. 1987. The precursor of Alzheimer's disease amyloid A4 protein resembles a cell-surface receptor. Nature 325: 733-736.
- Goate, A., et al. 1991. Segregation of a missense mutation in the amyloid precursor protein gene with familial Alzheimer's disease. Nature 349: 704-706.
- 3. Mullan, M., et al. 1992. A pathogenic mutation for probable Alzheimer's disease in the APP gene at the N-terminus of  $\beta$ -amyloid. Nat. Genet. 1: 345-347.
- Selkoe, D.J. 1998. The cell biology of β-amyloid precursor protein and presenilin in Alzheimer's disease. Trends. Cell Biol. 8: 447-453.
- 5. Yan, R., et al. 1999. Membrane-anchored aspartyl protease with Alzheimer's disease  $\beta$ -secretase activity. Nature 402: 533-537.
- Vassar, R., et al. 1999. Beta-secretase cleavage of Alzheimer's amyloid precursor protein by the transmembrane aspartic protease BACE. Science 286: 735-741.
- 7. Hussain, I., et al. 1999 Identification of a novel aspartic protease (Asp 2) as  $\beta$ -secretase Molec. Cell Neurosci. 14: 419-427.

## CHROMOSOMAL LOCATION

Genetic locus: BACE1 (human) mapping to 11q23.3; Bace1 (mouse) mapping to 9 A5.2.

## SOURCE

BACE (C-15) is an affinity purified goat polyclonal antibody raised against a peptide mapping at the C-terminus of BACE of human origin.

## PRODUCT

Each vial contains 200  $\mu g$  IgG in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

Blocking peptide available for competition studies, sc-10055 P, (100  $\mu$ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% BSA).

#### **STORAGE**

Store at 4° C, \*\*D0 NOT FREEZE\*\*. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

## PROTOCOLS

See our web site at www.scbt.com or our catalog for detailed protocols and support products.

#### **APPLICATIONS**

BACE (C-15) is recommended for detection of BACE of mouse, rat and human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), immunoprecipitation [1-2  $\mu$ g per 100-500  $\mu$ g of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

BACE (C-15) is also recommended for detection of BACE in additional species, including equine, canine, bovine, porcine and avian.

Suitable for use as control antibody for BACE siRNA (h): sc-37224, BACE siRNA (m): sc-37225, BACE shRNA Plasmid (h): sc-37224-SH, BACE shRNA Plasmid (m): sc-37225-SH, BACE shRNA (h) Lentiviral Particles: sc-37224-V and BACE shRNA (m) Lentiviral Particles: sc-37225-V.

Molecular Weight of BACE: 70 kDa.

Positive Controls: SH-SY5Y cell lysate: sc-3812.

## **RECOMMENDED SECONDARY REAGENTS**

To ensure optimal results, the following support (secondary) reagents are recommended: 1) Western Blotting: use donkey anti-goat IgG-HRP: sc-2020 (dilution range: 1:2000-1:100,000) or Cruz Marker™ compatible donkey anti-goat IgG-HRP: sc-2033 (dilution range: 1:2000-1:5000), Cruz Marker™ Molecular Weight Standards: sc-2035, TBS Blotto A Blocking Reagent: sc-2333 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml). 3) Immunofluorescence: use donkey anti-goat IgG-FITC: sc-2024 (dilution range: 1:100-1:400) or donkey anti-goat IgG-TR: sc-2783 (dilution range: 1:100-1:400) with UltraCruz™ Mounting Medium: sc-24941.

## SELECT PRODUCT CITATIONS

- Fuso, A., et al. 2007. γ-secretase is differentially modulated by alterations of homocysteine cycle in neuroblastoma and glioblastoma cells. J. Alzheimers Dis. 11: 275-290.
- Fuso, A., et al. 2008. B-vitamin deprivation induces hyperhomocysteinemia and brain S-adenosylhomocysteine, depletes brain S-adenosylmethionine, and enhances PS1 and BACE expression and Amyloid-β deposition in mice. Mol. Cell. Neurosci. 37: 731-746.

#### **RESEARCH USE**

For research use only, not for use in diagnostic procedures.

## Try BACE (61-3E7): sc-33711 or BACE (A-12):

MONOS Satisfation Guaranteed

sc-365948, our highly recommended monoclonal aternatives to BACE (C-15). Also, for AC, HRP, FITC, PE, Alexa Fluor<sup>®</sup> 488 and Alexa Fluor<sup>®</sup> 647 conjugates, see **BACE (61-3E7): sc-33711**.