Proteins



Product Data Sheet

BACE2 Protein, Mouse (HEK293, His)

Cat. No.: HY-P75588

Synonyms: Beta-secretase 2; Aspartyl protease 1; ASP1; Memapsin-1; Bace2

Species: Mouse Source: HEK293

Accession: Q9JL18 (A20-P462)

Gene ID: 56175

Molecular Weight: Approximately 55 kDa

PROPERTIES

AA Sequence	AVPALAPAPF TLPLQVAGAT NHRASAVPGL GTPELPRADG LALALEPVRA TANFLAMVDN LQGDSGRGYY LEMLIGTPPQ KVQILVDTGS SNFAVAGAPH SYIDTYFDSE SSSTYHSKGF DVTVKYTQGS WTGFVGEDLV TIPKGFNSSF LVNIATIFES ENFFLPGIKW NGILGLAYAA LAKPSSSLET FFDSLVAQAK IPDIFSMQMC GAGLPVAGSG TNGGSLVLGG IEPSLYKGDI WYTPIKEEWY YQIEILKLEI GGQNLNLDCR EYNADKAIVD SGTTLLRLPQ KVFDAVVEAV ARTSLIPEFS DGFWTGAQLA CWTNSETPWA YFPKISIYLR DENASRSFRI TILPQLYIQP MMGAGFNYEC YRFGISSSTN ALVIGATVME GFYVVFDRAQ RRVGFAVSPC AEIEGTTVSE ISGPFSTEDI ASNCVPAQAL	
Biological Activity	Measured by its ability to cleave a fluorogenic peptide substrate Mca-KPLGL-Dpa-AR-NH2. The specific activity is 2187.09 pmol/min/μg, as measured under the described conditions.	98
Appearance	Lyophilized powder	
Formulation	Lyophilized from a 0.2 μm filtered solution of PBS, pH 7.4.	
Endotoxin Level	<1 EU/μg, determined by LAL method.	
Reconsititution	It is not recommended to reconstitute to a concentration less than 100 μ g/mL in ddH ₂ O. For long term storage it is recommended to add a carrier protein (0.1% BSA, 5% HSA, 10% FBS or 5% Trehalose).	
Storage & Stability	Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.	
Shipping	Room temperature in continental US; may vary elsewhere.	

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DESCRIPTION

Background

The BACE2 protein assumes a critical role in the proteolytic processing of the amyloid precursor protein (APP), cleaving it between residues 690 and 691. This cleavage event results in the generation and extracellular release of beta-cleaved soluble APP and a corresponding cell-associated C-terminal fragment, later released by gamma-secretase. Additionally, BACE2 has been shown to cleave APP between residues 671 and 672. Beyond its involvement with APP, BACE2 plays a pivotal role in the proteolytic shedding of PMEL during early stages of melanosome biogenesis, cleaving PMEL within the Mbeta fragment to release the amyloidogenic PMEL luminal fragment comprising M-alpha and a small portion of M-beta N-terminus. This cleavage step is crucial for the subsequent processing and assembly of PMEL fibrils into amyloid sheets. Furthermore, BACE2 is responsible for the proteolytic processing of CLTRN in pancreatic beta cells, highlighting its diverse functions in cellular processes and contributing to our understanding of its significance in various physiological contexts.

Caution: Product has not been fully validated for medical applications. For research use only.

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