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Product datasheet

Recombinant Mouse CD13 protein (His tag) ab276741

1 Image

Description

Product name Recombinant Mouse CD13 protein (His tag)

Purity > 97 % SDS-PAGE.

Endotoxin level < 1.000 Eu/µg
Expression system HEK 293 cells

Accession P97449

Protein length Protein fragment

Animal free No

Nature Recombinant

Species Mouse

Sequence

DESKPWNQYR LPKTLIPDSY RVILRPYLTP NNQGLYIFQG NSTVRFTCNQ TTDVIIIHSK KLNYTLKGNH RVVLRTLDGT

PAPNIDKTEL VERTEYLVVH LQGSLVEGRQ YEMDSQFQGE LADDLAGFYR SEYMEGDVKK

VVATTQMQAA DARKSFPCFD EPAMKAMFNI TLIYPNNLIA

LSNMLPKESK PYPEDPSCTM TEFHSTPKMS

TYLLAYIVSE FKNISSVSAN GVQIGIWARP SAIDEGQGDY ALNVTGPILN FFAQHYNTSY PLPKSDQIAL PDFNAGAMEN

WGLVTYRESS LVFDSQSSSI SNKERVVTVI AHELAHQWFG NLVTVAWWND LWLNEGFASY VEYLGADYAE PTWNLKDLMV LNDVYRVMAV

DALASSHPLS SPADEIKTPD QIMELFDSIT YSKGASVIRM

LSSFLTEDLF KKGLSSYLHT YQYSNTVYLD LWEHLQKAVN QQTAVQPPAT VRTIMDRWIL

QMGFPVITVN TNTGEISQKH FLLDSKSNVT RPSEFNYIWI

APIPFLKSGQ EDHYWLDVEK NQSAKFQTSS

NEWILLNINV TGYYLVNYDE NNWKKLQNQL QTDLSVIPVI NRAQIIHDSF NLASAKMIPI TLALDNTLFL VKEAEYMPWQ

AALSSLNYFT LMFDRSEVYG PMKRYLKKQV

TPLFFYFONR TNNWVNRPPT LMEQYNEINA ISTACSSGLK

ECRDLVVELY SQWMKNPNNN TIHPNLRSTV YCNAIAFGGE EEWNFAWEQF RNATLVNEAD

KLRSALACSK DVWILNRYLS YTLNPDYIRK QDTTSTIISI

ASNVAGHPLV WDFVRSNWKK LFENYGGGSF

1

SFANLIQGVT RRFSSEFELQ QLEQFKADNS ATGFGTGTRA LEQALEKTRA NIDWVKENKD AVFKWFTENS S

Predicted molecular weight 104 kDa including tags

Molecular weight information The recombinant mouse ANPEP consists of 909 amino acids and has a calculated molecular

mass of 104 kDa. As a result of glycosylation, the recombinant protein migrates as an

approximately 120-130 kDa protein in SDS-PAGE under reducing conditions.

Amino acids 66 to 966

Tags His tag C-Terminus

Additional sequence information Predicted N-terminus: Lys 66.

Specifications

Our Abpromise guarantee covers the use of ab276741 in the following tested applications.

The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.

Applications SDS-PAGE

Form Lyophilized

Preparation and Storage

Stability and Storage Shipped at Room Temperature. Upon delivery aliquot. Store at -20°C or -80°C. Avoid freeze /

thaw cycle.

pH: 7.40

Constituent: 100% PBS

Reconstitution This information is lot specific. Please contact our technical Support team for details.

General Info

Function Broad specificity aminopeptidase. Plays a role in the final digestion of peptides generated from

hydrolysis of proteins by gastric and pancreatic proteases. May play a critical role in the pathogenesis of cholesterol gallstone disease. May be involved in the metabolism of regulatory peptides of diverse cell types including small intestinal and tubular epithelial cells, macrophages, granulocytes and synaptic membranes from the CNS. Found to cleave antigen peptides bound to major histocompatibility complex class II molecules of presenting cells and to degrade

neurotransmitters at synaptic junctions. Is also implicated as a regulator of IL-8 bioavailability in the endometrium, and therefore may contribute to the regulation of angiogenesis. Is used as a

marker for acute myeloid leukemia and plays a role in tumor invasion. In case of human coronavirus 229E (HCoV-229E) infection, serves as receptor for HCoV-229E spike glycoprotein.

Mediates as well human cytomegalovirus (HCMV) infection.

Tissue specificity Expressed in epithelial cells of the kidney, intestine, and respiratory tract; granulocytes,

monocytes, fibroblasts, endothelial cells, cerebral pericytes at the blood-brain barrier, synaptic membranes of cells in the CNS. Also expressed in endometrial stromal cells, but not in the endometrial glandular cells. Found in the vasculature of tissues that undergo angiogenesis and in malignant gliomas and lymph node metastases from multiple tumor types but not in blood vessels of normal tissues. A soluble form has been found in plasma. It is found to be elevated in plasma

and effusions of cancer patients.

Sequence similarities Belongs to the peptidase M1 family.

Domain Amino acids 260-353 are essential to mediate susceptibility to infection with HCoV-229E (in

porcine/human chimeric studies) and more specifically amino acids 288-295 (mutagenesis

studies).

Post-translational Sulfated.

modifications N- and O-glycosylated.

May undergo proteolysis and give rise to a soluble form.

Cellular localization Cell membrane. Cytoplasm > cytosol. A soluble form has also been detected.

Images



SDS-PAGE analysis of ab276741

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