

Product datasheet

Recombinant Human beta Catenin protein ab158230

1 Image

Description

Product name	Recombinant Human beta Catenin protein
Expression system	Wheat germ
Protein length	Full length protein
Animal free	No
Nature	Recombinant
Species	Human

Sequence	<p>MATQADLMELDMAMEPDRKAAVSHWQQQSYLDSGIH SGATTTAPSLSGKG NPEEEDVDTSQVLYEWEQGFSQSFTQE QVADIDGQY AMTRAQRVRAAMFP ETLDEGMQIPSTQFDAAHPTNVQRLAEPSQMLKHAVV NLINYQDDAELAT RAIPELTKLLNDEDQVVVNKAAVMVHQLSKKEASRHAI MRSPQMVSAVR TMQNTNDVETARCTAGTLHNLSSHREGLLAIFKSGGIP ALVKMLGSPVDS VLFYAITTLHNLHLLHQEGAKMAVRLAGGLQKMVALLNK TNVKFLAITTDC LQILAYGNQESKLIILASGGPQALVNIMRTYTYEKLLWTT SRVLKVL SVC SSNKPAIMEAGGMQALGLHLTDPSQRLVQNCLWTLRN LSDAATKQEGMEG LLGTLVQLLGSDDINVVTCAAGILSNLTCNNYKNKMMV CQVGGIEALVRT VLRAGDREDITEPAICALRHLTSRHQEAEMAQNAVRLH YGLPVVVKLLHP PSHWPLIKATVGLIRNLALCPANHAPLREQGAIPRLVQL LVRAHQDTQRR TSMGGTQQQFVEGVRMEEVEGCTGALHILARDVHNRI VIRGLNTIPLFV QLLYSPIENIQRVAAGVLCELAQDKEAAEAIEAEGATAP LTELLHSRNEG VATYAAAVLFRMSEDKPQDYKKRLSVELTSSLFRTEP MAWNETADLGLDI GAQGEPLGYRQDDPSYRSFHSGGYGQDALGMDPMM</p>
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Amino acids	1 to 781
Tags	GST tag N-Terminus

Specifications

Our [Abpromise guarantee](#) covers the use of **ab158230** in the following tested applications.

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

Applications	ELISA Western blot
Form	Liquid
Additional notes	Protein concentration is above or equal to 0.05 mg/ml.

Preparation and Storage

Stability and Storage	Shipped on dry ice. Upon delivery aliquot and store at -80°C. Avoid freeze / thaw cycles. pH: 8.00 Constituents: 0.31% Glutathione, 0.79% Tris HCl
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General Info

Function	<p>Key downstream component of the canonical Wnt signaling pathway. In the absence of Wnt, forms a complex with AXIN1, AXIN2, APC, CSNK1A1 and GSK3B that promotes phosphorylation on N-terminal Ser and Thr residues and ubiquitination of CTNNB1 via BTRC and its subsequent degradation by the proteasome. In the presence of Wnt ligand, CTNNB1 is not ubiquitinated and accumulates in the nucleus, where it acts as a coactivator for transcription factors of the TCF/LEF family, leading to activate Wnt responsive genes.</p> <p>Involved in the regulation of cell adhesion. The majority of beta-catenin is localized to the cell membrane and is part of E-cadherin/catenin adhesion complexes which are proposed to couple cadherins to the actin cytoskeleton.</p>
Tissue specificity	Expressed in several hair follicle cell types: basal and peripheral matrix cells, and cells of the outer and inner root sheaths. Expressed in colon.
Involvement in disease	<p>Defects in CTNNB1 are associated with colorectal cancer (CRC) [MIM:114500]. Note=Activating mutations in CTNNB1 have oncogenic activity resulting in tumor development. Somatic mutations are found in various tumor types, including colon cancers, ovarian and prostate carcinomas, hepatoblastoma (HB), hepatocellular carcinoma (HCC). HBs are malignant embryonal tumors mainly affecting young children in the first three years of life.</p> <p>Defects in CTNNB1 are a cause of pilomatixoma (PTR) [MIM:132600]; a common benign skin tumor.</p> <p>Defects in CTNNB1 are a cause of medulloblastoma (MDB) [MIM:155255]. MDB is a malignant, invasive embryonal tumor of the cerebellum with a preferential manifestation in children.</p> <p>Defects in CTNNB1 are a cause of susceptibility to ovarian cancer (OC) [MIM:167000]. Ovarian cancer common malignancy originating from ovarian tissue. Although many histologic types of ovarian neoplasms have been described, epithelial ovarian carcinoma is the most common form. Ovarian cancers are often asymptomatic and the recognized signs and symptoms, even of late-</p>

stage disease, are vague. Consequently, most patients are diagnosed with advanced disease. Note=A chromosomal aberration involving CTNNB1 is found in salivary gland pleiomorphic adenomas, the most common benign epithelial tumors of the salivary gland. Translocation t(3;8) (p21;q12) with PLAG1.

Sequence similarities

Belongs to the beta-catenin family.
Contains 12 ARM repeats.

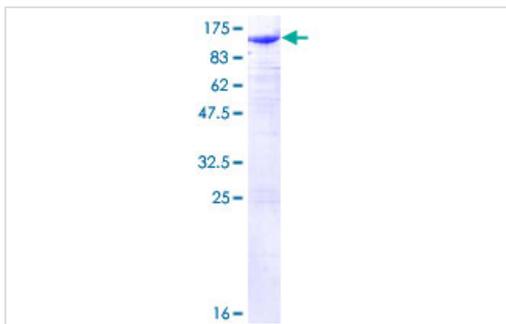
Post-translational modifications

Phosphorylation by GSK3B requires prior phosphorylation of Ser-45 by another kinase. Phosphorylation proceeds then from Thr-41 to Ser-37 and Ser-33. EGF stimulates tyrosine phosphorylation. Phosphorylation on Tyr-654 decreases CDH1 binding and enhances TBP binding. Ubiquitinated by the SCF(BTRC) E3 ligase complex when phosphorylated by GSK3B, leading to its degradation. Ubiquitinated by a E3 ubiquitin ligase complex containing UBE2D1, SIAH1, CACYBP/SIP, SKP1, APC and TBL1X, leading to its subsequent proteasomal degradation.

Cellular localization

Cytoplasm. Nucleus. Cytoplasm > cytoskeleton. Cell junction > adherens junction. Cell junction. Cell membrane. Cytoplasmic when it is unstabilized (high level of phosphorylation) or bound to CDH1. Translocates to the nucleus when it is stabilized (low level of phosphorylation). Interaction with GLIS2 and MUC1 promotes nuclear translocation. Interaction with EMD inhibits nuclear localization.

Images



ab158230 on a 12.5% SDS-PAGE stained with Coomassie Blue.

SDS-PAGE - Recombinant Human beta Catenin protein (ab158230)

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