

Full-Length

SEPT5 (Human) Recombinant Protein (P01)

Catalog # H00005413-P01 Size 25 ug, 10 ug

Applications



Specification	
Product Description	Human SEPT5 full-length ORF (AAH25261, 1 a.a 369 a.a.) recombinant protein with GST-tag at N -terminal.
Sequence	MSTGLRYKSKLATPEDKQDIDKQYVGFATLPNQVHRKSVKKGFDFTLMVAGESGLGKSTLVHSLF LTDLYKDRKLLSAEERISQTVEILKHTVDIEEKGVKLKLTIVDTPGFGDAVNNTECWKPITDYVDQQF EQYFRDESGLNRKNIQDNRVHCCLYFISPFGHGLRPVDVGFMKALHEKVNIVPLIAKADCLVPSEIR KLKERIREEIDKFGIHVYQFPECDSDEDEDFKQQDRELKESAPFAVIGSNTVVEAKGQRVRGRLY PWGIVEVENQAHCDFVKLRNMLIRTHMHDLKDVTCDVHYENYRAHCIQQMTSKLTQDSRMESPIPI LPLPTPDAETEKLIRMKDEELRRMQEMLQRMKQQMQDQ
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	66.33
Interspecies Antigen Sequence	Mouse (99)
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.



Product Information

Storage Instruction	Store at -80°C. Aliquot to avoid repeated freezing and thawing.
Note	Best use within three months from the date of receipt of this protein.

Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — SEPT5	
Entrez GeneID	<u>5413</u>
GeneBank Accession#	BC025261
Protein Accession#	AAH25261
Gene Name	SEPT5
Gene Alias	CDCREL, CDCREL-1, CDCREL1, H5, PNUTL1
Gene Description	septin 5
Omim ID	602724
Gene Ontology	<u>Hyperlink</u>
Gene Summary	This gene is a member of the septin gene family of nucleotide binding proteins, originally describ ed in yeast as cell division cycle regulatory proteins. Septins are highly conserved in yeast, Droso phila, and mouse and appear to regulate cytoskeletal organization. Disruption of septin function di sturbs cytokinesis and results in large multinucleate or polyploid cells. This gene is mapped to 22 q11, the region frequently deleted in DiGeorge and velocardiofacial syndromes. A translocation in volving the MLL gene and this gene has also been reported in patients with acute myeloid leukem ia. Two transcripts of this gene, a major one of 2.2 kb and a minor one of 3.5 kb, have been obser ved. The 2.2 kb form results from the utilization of a non-consensus polyA signal (AACAAT). In the absence of polyadenylation from this imperfect site, the consensus polyA signal of the downstrea m neighboring gene (GP1BB; platelet glycoprotein lb) is used, resulting in the 3.5 kb transcript. A n alternatively spliced transcript variant with a different 5' end has also been identified, but its full-length nature has not been completely determined. [provided by RefSeq



Product Information

Other Designations

cell division control related protein 1|peanut-like 1|platelet glycoprotein lb beta chain