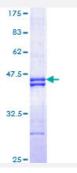


PSMC3 (Human) Recombinant Protein (Q01)

Catalog # H00005702-Q01 Size 25 ug, 10 ug

Applications



Specification	
Product Description	Human PSMC3 partial ORF (AAH08713, 53 a.a 152 a.a.) recombinant protein with GST-tag at N-t erminal.
Sequence	MNLLPNIESPVTRQEKMATVWDEAEQDGIGEEVLKMSTEEIIQRTRLLDSEIKIMKSEVLRVTHELQ AMKDKIKENSEKIKVNKTLPYLVSNVIELLDVD
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	36.63
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.
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 — PSMC3	
Entrez GenelD	<u>5702</u>
GeneBank Accession#	BC008713
Protein Accession#	AAH08713
Gene Name	PSMC3
Gene Alias	MGC8487, TBP1
Gene Description	proteasome (prosome, macropain) 26S subunit, ATPase, 3
Omim ID	<u>186852</u>
Gene Ontology	<u>Hyperlink</u>
Gene Summary	The 26S proteasome is a multicatalytic proteinase complex with a highly ordered structure composed of 2 complexes, a 20S core and a 19S regulator. The 20S core is composed of 4 rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are composed of 7 beta subunits. The 19S regulator is composed of a base, which contains 6 ATPase subunits and 2 non-ATPase subunits, and a lid, which contains up to 10 non-ATPase subunits. Proteasomes ar e distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ub iquitin-dependent process in a non-lysosomal pathway. An essential function of a modified protea some, the immunoproteasome, is the processing of class I MHC peptides. This gene encodes on e of the ATPase subunits, a member of the triple-A family of ATPases that have chaperone-like a ctivity. This subunit may compete with PSMC2 for binding to the HIV tat protein to regulate the inte raction between the viral protein and the transcription complex. A pseudogene has been identified on chromosome 9. [provided by RefSeq
Other Designations	Tat-binding protein 1 human immunodeficiency virus tat transactivator binding protein-1 proteaso me 26S ATPase subunit 3 proteasome subunit P50

Pathway



Proteasome