

Product datasheet for PH301260

OriGene Technologies, Inc.

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PSME1 (NM_006263) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: PSME1 MS Standard C13 and N15-labeled recombinant protein (NP_006254)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC201260

or AA Sequence:

Protein Sequence:

Predicted MW: 28.7 kDa

>RC201260 protein sequence
Red=Cloning site Green=Tags(s)

MAMLRVQPEAQAKVDVFREDLCTKTENLLGSYFPKKISELDAFLKEPALNEANLSNLKAPLDIPVPDPVK EKEKEERKKQQEKEDKDEKKKGEDEDKGPPCGPVNCNEKIVVLLQRLKPEIKDVIEQLNLVTTWLQLQIP RIEDGNNFGVAVQEKVFELMTSLHTKLEGFHTQISKYFSERGDAVTKAAKQPHVGDYRQLVHELDEAEYR

DIRLMVMEIRNAYAVLYDIILKNFEKLKKPRGETKGMIY

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-Myc/DDK

Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining

Concentration: >0.05 μg/μL as determined by microplate BCA method

Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3

Storage: Store at -80°C. Avoid repeated freeze-thaw cycles.

Stability: Stable for 3 months from receipt of products under proper storage and handling conditions.

RefSeq: NP 006254

RefSeq Size: 1023 RefSeq ORF: 747

Synonyms: HEL-S-129m; IFI5111; PA28A; PA28alpha; REGalpha

Locus ID: 5720

UniProt ID: Q06323, A0A0K0K1L8





Cytogenetics:

14q12

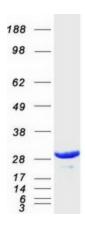
Summary:

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 are distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An essential function of a modified proteasome, the immunoproteasome, is the processing of class I MHC peptides. The immunoproteasome contains an alternate regulator, referred to as the 11S regulator or PA28, that replaces the 19S regulator. Three subunits (alpha, beta and gamma) of the 11S regulator have been identified. This gene encodes the alpha subunit of the 11S regulator, one of the two 11S subunits that is induced by gamma-interferon. Three alpha and three beta subunits combine to form a heterohexameric ring. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Jul 2013]

Protein Pathways:

Antigen processing and presentation, Proteasome

Product images:



Coomassie blue staining of purified PSME1 protein (Cat# [TP301260]). The protein was produced from HEK293T cells transfected with PSME1 cDNA clone (Cat# [RC201260]) using MegaTran 2.0 (Cat# [TT210002]).