

# Product datasheet for PH318938

## PSMF1 (NM\_006814) Human Mass Spec Standard

### **Product data:**

#### Product Type: Mass Spec Standards **Description:** PSMF1 MS Standard C13 and N15-labeled recombinant protein (NP\_006805) Species: Human **HEK293 Expression Host:** RC218938 **Expression cDNA Clone** or AA Sequence: Predicted MW: 29.6 kDa >RC218938 representing NM\_006814 **Protein Sequence:** Red=Cloning site Green=Tags(s) MAGLEVLFASAAPAITCRQDALVCFLHWEVVTHGYFGLGVGDQPGPNDKKSELLPAGWNNNKDLYVLRYE YKDGSRKLLVKAITVESSMILNVLEYGSQQVADLTLNLDDYIDAEHLGDFHRTYKNSEELRSRIVSGIIT PIHEQWEKANVSSPHREFPPATAREVDPLRIPPRHPHTSRQPPWCDPLGPFVVGGEDLDPFGPRRGGMIV DPLRSGFPRALIDPSSGLPNRLPPGAVPPGARFDPFGPIGTSPPGPNPDHLPPPGYDDMYL 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 006805 **RefSeq Size:** 3241 **RefSeq ORF:** 813 Synonyms: PI31 9491 Locus ID: UniProt ID: Q92530, A0A140VJT2



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### OriGene Technologies, Inc.

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	1 (NM_006814) Human Mass Spec Standard – PH318938	
Cytogenetics:	20p13	
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. This gene encodes a protein that inhibits the activation of the proteasome by the 11S and 19S regulators. Alternative transcript variants have been identified for this gene. [provided by RefSeq, Jul 2008]	
Protein Pathway	t <b>ein Pathways:</b> Proteasome	

### **Product images:**

188	_	
98	_	
62	_	
49	_	
38	_	
28	_	
17	_	
14		
63	=	

Coomassie blue staining of purified PSMF1 protein (Cat# [TP318938]). The protein was produced from HEK293T cells transfected with PSMF1 cDNA clone (Cat# [RC218938]) using MegaTran 2.0 (Cat# [TT210002]).

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