

## Product datasheet for TP301201M

### PSMD11 (NM\_002815) Human Recombinant Protein

#### Product data:

|                                       |  |
|---------------------------------------|--|
| Product Type:                         | Recombinant Proteins   |
| Description:                          | Recombinant protein of human proteasome (prosome, macropain) 26S subunit, non-ATPase, 11 (PSMD11), 100 µg                                |
| Species:                              | Human  |
| Expression Host:                      | HEK293T  |
| Expression cDNA Clone or AA Sequence: | >RC201201 representing NM_002815<br><span style="color: red;">Red</span> =Cloning site <span style="color: green;">Green</span> =Tags(s) |

MAAAAVVEFQRAQSLLSTDREASIDILHSIVKRDIQENDEEAVQVKEQSILELGSLLAKTGQAAELGGLL  
 KYVRPFLNSISKAKAARLVRSLLDLFLDMEAATGQEVELCLEIEWAKSEKRTFLRQALEARLVSLYFDT  
 KRYQEALHLGSQLLRELKKMDDKALLVEVQLLESKTYHALSNLPKARAALTSARTTANAIYCPKQLQATL  
 DMQSGIIHAAEEKDWKTAYSIFYEAFEGYDSIDSPKAITSLKYMILLCKIMLNTPEDVQALVSGKLALRYA  
 GRQTEALKCVAQASKNRSADFELKALTDYRAELRDDPIISTHLAKLYDNLLEQNLIRVIEPFSRVQIEHI  
 SSLIKLSKADVERKLSQMILDKKFHGILDQGEGLIIFDEPPVDKTYEAALETIQNMSKVVDLSLYNKAKK  
 LT

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

|                |  |
|----------------|--|
| Tag:           | C-Myc/DDK  |
| Predicted MW:  | 47.3 kDa   |
| Concentration: | >0.05 µg/µL as determined by microplate BCA method   |
| Purity:        | > 80% as determined by SDS-PAGE and Coomassie blue staining  |
| Buffer:        | 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol   |
| Preparation:   | Recombinant protein was captured through anti-DDK affinity column followed by conventional chromatography steps.                                     |
| Note:          | For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process. |
| Storage:       | Store at -80°C.  |
| Stability:     | Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.        |


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RefSeq: [NP\\_002806](#)

Locus ID: 5717

UniProt ID: [O00231](#)

RefSeq Size: 1598

Cytogenetics: 17q11.2

RefSeq ORF: 1266

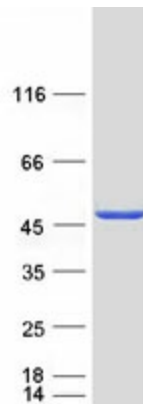
Synonyms: p44.5; Rpn6; S9

**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. This gene encodes a member of the proteasome subunit S9 family that functions as a non-ATPase subunit of the 19S regulator and is phosphorylated by AMP-activated protein kinase. Alternatively spliced transcript variants have been observed for this gene. [provided by RefSeq, Jul 2012]

**Protein Families:** Stem cell - Pluripotency

**Protein Pathways:** Proteasome

## Product images:



Coomassie blue staining of purified PSMD11 protein (Cat# [TP301201]). The protein was produced from HEK293T cells transfected with PSMD11 cDNA clone (Cat# [RC201201]) using MegaTran 2.0 (Cat# [TT210002]).