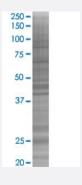


## PSMB5 293T Cell Transient Overexpression Lysate(Denatured)

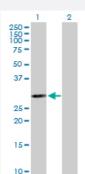
Catalog # H00005693-T03 Size 100 uL

### **Applications**



#### SDS-PAGE Gel

PSMB5 transfected lysate.



#### Western Blot

Lane 1: PSMB5 transfected lysate (29.04 KDa)

Lane 2: Non-transfected lysate.

| Specification           |  |
|-------------------------|--|
| Transfected Cell Line   | 293T   |
| Plasmid                 | pCMV-PSMB5 full-length   |
| Host                    | Human  |
| Theoretical MW (kDa)    | 29.04  |
| Quality Control Testing | Transient overexpression cell lysate was tested with Anti-PSMB5 antibody ( <u>H00005693-B01P</u> ) by W estern Blots.  SDS-PAGE Gel PSMB5 transfected lysate.  Western Blot Lane 1: PSMB5 transfected lysate (29.04 KDa) Lane 2: Non-transfected lysate. |



### **Product Information**

| Storage Buffer      | 1X Sample Buffer (50 mM Tris-HCl, 2% SDS, 10% glycerol, 300 mM 2-mercaptoethanol, 0.01% Bro mophenol blue) |
|---------------------|--|
| Storage Instruction | Store at -80°C. Aliquot to avoid repeated freezing and thawing.  |

## **Applications**

Western Blot

| Gene Info — PSMB5   |   |
|---------------------|---|
| Entrez GenelD       | <u>5693</u>   |
| GeneBank Accession# | BC057840  |
| Protein Accession#  | AAH57840  |
| Gene Name           | PSMB5   |
| Gene Alias          | LMPX, MB1, MGC104214, X   |
| Gene Description    | proteasome (prosome, macropain) subunit, beta type, 5   |
| Omim ID             | 600306  |
| Gene Ontology       | <u>Hyperlink</u>  |
| Gene Summary        | The proteasome is a multicatalytic proteinase complex with a highly ordered ring-shaped 20S cor e structure. The core structure is composed of 4 rings of 28 non-identical subunits; 2 rings are co mposed of 7 alpha subunits and 2 rings are composed of 7 beta 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 member of the proteasome B-type family, also known as the T1B family, that is a 20S core beta subunit in the proteasome. This catalytic subunit is not present in the immunoproteasome and is replaced by catalytic subunit 3i (proteasome beta 8 subunit). Multiple transcript variants encoding different is oforms have been found for this gene. [provided by RefSeq |
| Other Designations  | PSX large multifunctional protease X macropain epsilon chain multicatalytic endopeptidase comp lex epsilon chain proteasome beta 5 subunit proteasome catalytic subunit 3 proteasome chain 6 p roteasome epsilon chain proteasome subunit MB1 proteasome subunit  |

# Pathway



Proteasome