# antibodies .- online.com





# anti-PSMB1 antibody (Middle Region)



Image



Publication



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**Target Details** 

PSMB1

Target:

Quantity:	100 μL
Target:	PSMB1
Binding Specificity:	Middle Region
Reactivity:	Human, Mouse, Rat, Rabbit, Saccharomyces cerevisiae, Cow, Dog, Guinea Pig, Horse, Zebrafish (Danio rerio)
Host:	Rabbit
Clonality:	Polyclonal
Conjugate:	This PSMB1 antibody is un-conjugated
Application:	Western Blotting (WB)
Product Details	
Product Details Immunogen:	The immunogen is a synthetic peptide directed towards the middle region of human PSMB1
	The immunogen is a synthetic peptide directed towards the middle region of human PSMB1  NNKAMTTGAI AAMLSTILYS RRFFPYYVYN IIGGLDEEGK GAVYSFDPVG
lmmunogen:	
Immunogen: Sequence:	NNKAMTTGAI AAMLSTILYS RRFFPYYVYN IIGGLDEEGK GAVYSFDPVG  Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit:

Alternative Name:	PSMB1 (PSMB1 Products)
Background:	The proteasome is a multicatalytic proteinase complex with a highly ordered ring-shaped 20S
	core structure. The core structure 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. 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.
	PSMB1 is a member of the proteasome B-type family, also known as the T1B family, that is a
	20S core beta subunit. This gene is tightly linked to the TBP (TATA-binding protein) gene in
	human and in mouse, and is transcribed in the opposite orientation in both species. The
	proteasome is a multicatalytic proteinase complex with a highly ordered ring-shaped 20S core
	structure. The core structure 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. 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. This gene is tightly linked to the TBP (TATA-binding protein) gene in
	human and in mouse, and is transcribed in the opposite orientation in both species. Publication
	Note: This RefSeq record includes a subset of the publications that are available for this gene.
	Please see the Entrez Gene record to access additional publications.
	Alias Symbols: FLJ25321, HC5, KIAA1838, PMSB1, PSC5
	Protein Interaction Partner: MAPRE3, UBC, TRAF1, TRIM27, KRT15, KRT13, KIFC3, CDKN1B,
	PSMA7, HAUS2, PSMD14, PSEN1, PSMG3, PSMB8, PSMB7, PSMB6, PSMB5, PSMB4, PSMB3,
	PSMB2, PSMA6, PSMA5, PSMA4, PSMA3, PSMA1, HNRNPA2B1, DDB2, SMEK1, AP3M1,
	PSME4, RPP38, RPP30, SEC13, PYCRL, HECW2, PSMA2
	Protein Size: 241
Molecular Weight:	26 kDa
Gene ID:	5689
NCBI Accession:	NM_002793, NP_002784
UniProt:	P20618
Pathways:	Mitotic G1-G1/S Phases, DNA Replication, Synthesis of DNA

## **Application Details**

Application Details		
Application Notes:	Optimal working dilutions should be determined experimentally by the investigator.	
Comment:	Antigen size: 241 AA	
Restrictions:	For Research Use only	
Handling		
Format:	Liquid	
Concentration:	Lot specific	
Buffer:	Liquid. Purified antibody supplied in 1x PBS buffer with 0.09 % (w/v) sodium azide and 2 % sucrose.	
Preservative:	Sodium azide	
Precaution of Use:	This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only.	
Handling Advice:	Avoid repeated freeze-thaw cycles.	
Storage:	-20 °C	
Storage Comment:	For short term use, store at 2-8°C up to 1 week. For long term storage, store at -20°C in small aliquots to prevent freeze-thaw cycles.	

### **Publications**

Product cited in:

Ewing, Chu, Elisma, Li, Taylor, Climie, McBroom-Cerajewski, Robinson, OConnor, Li, Taylor, Dharsee, Ho, Heilbut, Moore, Zhang, Ornatsky, Bukhman, Ethier, Sheng, Vasilescu, Abu-Farha, Lambert, Duewel et al.: "Large-scale mapping of human protein-protein interactions by mass spectrometry. ..." in: **Molecular systems biology**, Vol. 3, pp. 89, (2007) (PubMed).

90 kDa\_\_ 65 kDa\_\_ 40 kDa\_\_ 29 kDa\_\_ 22 kDa\_\_

### **Western Blotting**

Image 1. WB Suggested Anti-PSMB1 Antibody Titration:

0.2-1 ug/ml

**ELISA Titer:** 1:312500

Positive Control: 721\_B cell lysate