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Datasheet for ABIN2786720 anti-PSMB5 antibody (Middle Region)

1 Image

1 Publication



Overview

000101000	
Quantity:	100 µL
Target:	PSMB5
Binding Specificity:	Middle Region
Reactivity:	Human, Mouse, Rat, Cow, Dog, Horse, Rabbit, Zebrafish (Danio rerio), Guinea Pig,
	Saccharomyces cerevisiae
Host:	Rabbit
Clonality:	Polyclonal
Conjugate:	This PSMB5 antibody is un-conjugated
Application:	Western Blotting (WB)
Product Details	
Immunogen:	The immunogen is a synthetic peptide directed towards the middle region of human PSMB5
Sequence:	IVAADSRATA GAYIASQTVK KVIEINPYLL GTMAGGAADC SFWERLLARQ
Sequence: Predicted Reactivity:	IVAADSRATA GAYIASQTVK KVIEINPYLL GTMAGGAADC SFWERLLARQ Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit:
	Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit:
Predicted Reactivity:	Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit: 100%, Rat: 100%, Yeast: 92%, Zebrafish: 100%
Predicted Reactivity:	Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit: 100%, Rat: 100%, Yeast: 92%, Zebrafish: 100% This is a rabbit polyclonal antibody against PSMB5. It was validated on Western Blot using a
Predicted Reactivity: Characteristics:	Cow: 100%, Dog: 100%, Guinea Pig: 100%, Horse: 100%, Human: 100%, Mouse: 100%, Rabbit: 100%, Rat: 100%, Yeast: 92%, Zebrafish: 100% This is a rabbit polyclonal antibody against PSMB5. It was validated on Western Blot using a cell lysate as a positive control.

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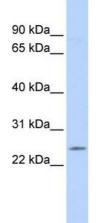
Target Details		
Alternative Name:	PSMB5 (PSMB5 Products)	
Background:	The proteasome is a multicatalytic proteinase complex with a highly ordered ring-shaped 20S	
	core structure. The core structure is composed of 4 rings of 28 non-identical subunits, 2 rings	
	are composed of 7 alpha subunits and 2 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.	
	PSMB5 is 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).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 in the proteasome. This catalytic subunit is not present in the	
	immunoproteasome and is replaced by catalytic subunit 3i (proteasome beta 8 subunit).	
	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: LMPX, MB1, MGC104214, X	
	Protein Interaction Partner: UBC, AURKA, PSMD14, BMI1, PSMG3, PSMB8, PSMB7, PSMB6,	
	PYCRL, AP3M1, RPP38, PSMB4, PSMB3, PSMB2, PSMB1, PSMA7, PSMA6, PSMA5, PSMA4,	
	PSMA3, PSMA1, KIF5B, EPB41L1, CAST, XRN1, CDK20, CDKL1, PSMA2, PARK2, BAG3, IQCB1,	
	HSP90AB1, HSP90AA1, FN1, IKBKAP, PSMA8, P	
	Protein Size: 263	
Molecular Weight:	28 kDa	
Gene ID:	5693	
NCBI Accession:	NM_002797, NP_002788	
UniProt:	P28074	
Pathways:	Mitotic G1-G1/S Phases, DNA Replication, Synthesis of DNA	

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Application Details		
Application Notes:	Optimal working dilutions should be determined experimentally by the investigator.	
Comment:	Antigen size: 263 AA	
Restrictions:	For Research Use only	

Handling

sucrose. Preservative: Sodium azide Precaution of Use: This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE will 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 aliquots to prevent freeze-thaw cycles. Publications Ewing, Chu, Elisma, Li, Taylor, Climie, McBroom-Cerajewski, Robinson, OConnor, Li, Tay Dharsee, Ho, Heilbut, Moore, Zhang, Ornatsky, Bukhman, Ethier, Sheng, Vasilescu, Abu-		
Buffer: Liquid. Purified antibody supplied in 1x PBS buffer with 0.09 % (w/v) sodium azide and sucrose. Preservative: Sodium azide Precaution of Use: This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE will 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 aliquots to prevent freeze-thaw cycles. Publications Ewing, Chu, Elisma, Li, Taylor, Climie, McBroom-Cerajewski, Robinson, OConnor, Li, Tay Dharsee, Ho, Heilbut, Moore, Zhang, Ornatsky, Bukhman, Ethier, Sheng, Vasilescu, Abu-Lambert, Duewel et al.: "Large-scale mapping of human protein-protein interactions by	Format:	Liquid
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Product cited in: Ewing, Chu, Elisma, Li, Taylor, Climie, McBroom-Cerajewski, Robinson, OConnor, Li, Tay Dharsee, Ho, Heilbut, Moore, Zhang, Ornatsky, Bukhman, Ethier, Sheng, Vasilescu, Abu- Lambert, Duewel et al.: "Large-scale mapping of human protein-protein interactions by	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.
Dharsee, Ho, Heilbut, Moore, Zhang, Ornatsky, Bukhman, Ethier, Sheng, Vasilescu, Abu- Lambert, Duewel et al.: "Large-scale mapping of human protein-protein interactions by	Publications	
Lambert, Duewel et al.: "Large-scale mapping of human protein-protein interactions by	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,
spectrometry" in: Molecular systems biology , Vol. 3, pp. 89, (2007) (PubMed).		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).



Western Blotting

Image 1. WB Suggested Anti-PSMB5 Antibody Titration:

0.2-1 ug/ml

ELISA Titer: 1:62500

Positive Control: MCF7 cell lysate

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