Datasheet for ABIN7164970 anti-PSMB1 antibody (AA 29-241) (FITC)

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Overview

Quantity:	100 µg
Target:	PSMB1
Binding Specificity:	AA 29-241
Reactivity:	Human
Host:	Rabbit
Clonality:	Polyclonal
Conjugate:	This PSMB1 antibody is conjugated to FITC
Application:	Please inquire

Product Details

Immunogen:	Recombinant Human Proteasome subunit beta type-1 protein (29-241AA)
Isotype:	lgG
Cross-Reactivity:	Human
Purification:	>95%, Protein G purified

Target Details

Target:	PSMB1
Alternative Name:	PSMB1 (PSMB1 Products)
Background:	Background: The proteasome is a multicatalytic proteinase complex which is characterized by
	its ability to cleave peptides with Arg, Phe, Tyr, Leu, and Glu adjacent to the leaving group at

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Storage Comment:	Upon receipt, store at -20°C or -80°C. Avoid repeated freeze.
Storage:	-20 °C,-80 °C
Precaution of Use:	This product contains ProClin: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only.
Preservative:	ProClin
Buffer:	Preservative: 0.03 % Proclin 300 Constituents: 50 % Glycerol, 0.01M PBS, PH 7.4
Format:	Liquid
Handling	
Restrictions:	For Research Use only
Application Details	
Pathways:	Mitotic G1-G1/S Phases, DNA Replication, Synthesis of DNA
UniProt:	P20618
	antibody, PSC5 antibody, PSMB 1 antibody, PSMB1 antibody
	Proteasome subunit beta type-1 antibody, Proteasome subunit HC5 antibody, PSB1_HUMAN
	antibody, Proteasome gamma chain antibody, Proteasome subunit beta type 1 antibody,
	subunit beta type 1 antibody, Proteasome beta 1 subunit antibody, Proteasome component C5
	Aliases: FLJ25321 antibody, HC5 antibody, KIAA1838 antibody, Macropain subunit C5 antibody, Multicatalytic endopeptidase complex subunit C5 antibody, Proteasome (prosome macropain)
	the inside of the barrel. Interacts with SERPINB2. Interacts with HIV-1 TAT protein.
	rings are each formed by seven beta subunits. The catalytic chamber with the active sites is on
	structure. The two end rings are each formed by seven alpha subunits, and the two central
	composed of 28 subunits that are arranged in four stacked rings, resulting in a barrel-shaped
	a 20S proteasome core and two 19S regulatory subunits. The 20S proteasome core is
	activity.Cleavage of peptide bonds with very broad specificity.The 26S proteasome consists of
	neutral or slightly basic pH . The proteasome has an ATP-dependent proteolytic