CAB1985



Product Information

Product SKU:	CAB1985	Gene ID:	5701	Size:	20uL, 100uL	
Clone No:	-	Host Species:	Rabbit	Reactivity :	Human, Mouse, Rat	
Additional Information						

Observed MW:49kDaConjugate:UnconjugatedCalculated MW:49kDaIsotype:IgG

Immunogen Information

Background	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. An essential function of a modified proteasome, the immunoproteasome, is the
	processing of class I MHC peptides. This gene encodes one of the ATPase subunits, a member of the
	triple-A family of ATPases which have a chaperone-like activity. This subunit has been shown to interact
	with several of the basal transcription factors so, in addition to participation in proteasome functions,
	this subunit may participate in the regulation of transcription. This subunit may also compete with
	PSMC3 for binding to the HIV tat protein to regulate the interaction between the viral protein and the
	transcription complex. Alternative splicing results in multiple transcript variants encoding distinct
	isoforms.
Recommended Dilution:	WB,1:500 - 1:2000
Synonyms:	S7; MSS1; RPT1; Nbla10058; PSMC2
Purifcation Method:	Affinity purification
Immunogen:	Recombinant fusion protein containing a sequence corresponding to amino acids 1-160 of human
	PSMC2 (NP_002794.1).
Storage:	Store at -20°C. Avoid freeze / thaw cycles.Buffer: PBS with 0.02% sodium azide,50% glycerol,pH7.3.