

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q99460	PSMD1_HUMAN	Homo sapiens	26S proteasome non-ATPase regulatory subunit 1	25.810628	S829	T273;S290;T311;S315;T830;S834	30444036;26374642;16408927;36580660;35138101;34019948;32119511;28510447;34725712;33465208;38665916	<p> MITSAAAGIISLLDEDEPQLKEFALHK LNAVVNDFWAEISESVDKIEVLYEDE GFRSRQFAALVASKVFYHLGAFEEES LNYALGAGDLFNVNDNSEYVETIAK CIDHYTKQCVENADLPEGEKKPIDQ RLEGIVNKMFORCLDDHKYQAIIGI ALETRRLDVFEKTILESNDVPGMLAY SLKLCMSLMQNKQFRNKVLRVVKI YMNLEKPDFINVCQCLIFLDDPQAV SDILEKLVKEDNLLMAYQICFDLYES ASQQFLSSVIQNLRTVGTPIASVPGS TNTGTVPGEKSDSDMETEETSSA FVGKTPEASPEPKDQTLKMIKLSGE MAIELHLQFLIRNNNTDLMILKNTK DAVRNSVCHTATVIANSFMHCGTTS DQFLRDNLEWLRATNWKFTATA SLGVIHKGHEKEALQLMATYLPKDT SPGSAYQEGGGLYALGLIHANHGGD IIDYLLNQLKNASNDIVRHGGSLGLG LAAMGTARQDVYDLLKTNLYQDDAV TGEAAGLALGLVMLGSKNAQAIEDM VGYAQETQHEKILRGLAVGIALVY GRMEEADALIESLDRDKDPILRRSG MYTVAMAYCGSGNNAIRRLHVAV SDVNDVRRRAAVESLGFILFRTPEQ CPSVVLLSESYNPHVRYGAAMALGI CCAGTGNKEAINLLEPMTNDPVNYV ROGALIASALIMIQTEITCPKVNQF RQLYSKVINDKHDDVMAKFGAILAQ GILDAGGHNVTISLQSRGTGHTHMP VVGVLVFTQFWFVPLSHFLSLAYT PTCVIGLNKDLKMPKVQYKSNCKPS TFAYPAPLEVPKEKEKEKSTAVLSIT AKAKKKEKEKEKEKEKMEVDEAEK KEEKKKEPEPNFQLLDNPARVMP AQLKVLTMPETCRYOPFKPLSIGGIII LKDTSEDIELVEPVAAHGPKIEEEE QEPEPPPEFYIDD </p>	True	False	4.633	4.662	1.871	1.558	1.028