

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q3TXS7	PSMD1_MOUSE	Mus musculus	26S proteasome non-ATPase regulatory subunit 1	22.380232	NaN	T273;S290;T311;S315;T830;S834	37507081	MITSAAAGIISLLDEEEPOLKEFALHK LNAVVNDFWAEISESVDKIEVLYEDE GFRSRQFAALVASKVVFYHLGAFEEES LNYALGAGDLFNVDNSENSEYVETIIAK CIDHYTKQCVENADLPEGEKKPIDQ RLEGIVNKMFQRCLDDHKYKQAI ALETRRLDVFEKILESNVDPGMLAY SLKLCMSLMQNKQFRNKVLRVVKI YMNLEKVPDFINVCQCLIFLDDPQAV SDILEKLVKEDNLLMAYQICFDLYES ASQQFLSSVIQNLRTVGTPIASVPGS TNTGTVPGEKSDPMETEEKTASA VAGKTPDASPEPKDQTLKMIKILSGE MAIELHLQFLIRNNNTDLMILKNTK DAVRNSVCHTATVIANSEFMHCGTTS DQFLRDNLEWLARATNWAKFTATA SLGVIIHKGHEKEALQLMATYLPKDT SPGSAYQEGGGLYALGLIHANHGGD IIDYLLNQLKNASNDIVRHGGSLGLG LAAMGTARQDVYDLLKTNLYQDDAV TGEAAGLALGLVMLGSKNAQAIEDM VGYAQETOHEKILRGLAVGIALVMY GRMEEADALIESLCRDKDPILRRSG MYTVAMAYCGSGNNKAIRRLHVAV SDVNDVRRAAVESLGFILFRTPPEQ CPSVVSLLSESYNPHVRYGAAMALGI CCAGTGKNEAINLLEPMTNDPVNYV RQGALIASALMIOQTEITCPKVNQF RQLYSKVINDKHDDVMAKFGAILAQ GILDAGGHNVTISLQSRGTGHTHMPS VVGVLVFTQFWFWFPLSHFLSLAYT PTCVIGLNKDLKMPKVQYKSNCKPS TFAYPAPLEVPKEKEKEKVESTAVLSIT AKAKKKEKEKEKEKEKMEVDEAEK KEEKKEKPEPNFQLLDNPARVMP AQLKVLVSMETCRYPFKPLSIGGIII LKDTSEVVEELVEPVAAHGPKIEEEE QEPEPEPEFYIDD	True	False	3.155	4.039	1.559	1.438	0.907	1.097	0.933