

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q13200	PSMD2_HUMAN	Homo sapiens	26S proteasome non-ATPase regulatory subunit 2	22.549522	S409;S413	S16;T24;S29;S147;T194;S361;S363	39990658;30397120;23301498;20305658;33214551;16408927;40596516;26853435;38665916;34725712;32119511;40914422;34019948;35138101;38253038	MEEGGRDKAPVQPQSPAAAPGGT DEKPSGKERRDAGDKDKEQELSEED KQLQDELEMLVERLGEKDTSLYRPA LEELRRQIRSSSTTSMTSVPKPLKFLR PHYGKLKEIYENMAPGENKRFAADII SVLAMTMSGERECLKYRLVGSQEEL ASWGHEYYRHLAGEVAKEWQELDD AEKVQREPLLTLVKEIVPYNMAHNA EHEACDLLMEIEQVDMLEKIDIDENA YAKVCLYLTSCVNYVPEPENSALLRC ALGVFRKFSRFPALRLALMLNDME LVEDIFTCKDQVVKQMAFMLGR HGVFLELSEEDVEEYEDLTEIMSNVQ LNSNFLALARELDIMEPKVPDDIYKT HLENNRFGSGSQVDSARMNLASS FVNGFVNAAFGQDKLLTDDGNKWL YKNKDHGMLSAAAALGMILLWDVD GGLTQIDKYLSSSEDIKSGALLACGI VNSGVRNECDPALALLSDYVLLHNSN TMRGSIIFGLGLAYAGSNREDVLTLL LPVMGDSKSSMEVAGVTALACGMIA VGSCNGDVTSTILOIMEKSETELKD TYARWLPLGLNLHLGKGEAIEAAILA ALEVVSEPFPSFANTLVDVCAAYAGS GNVLKVQQLLHICSEHFDSKEKEED KDKKEKDKKKEAPADMGAHQGV AVLGIALIAMGEEIGAEMALRTFGHL LRYGEPTLRRVPLALALISVSNPRL NILDTLSKFSHDADPEVSYNSIFAMG MVGSGTNNARLAAMLRLQLAQYHAK DPNNLFMVRLAQGLTHLGKGTLTLC PYHSDRQLMSQVAVAGLLTVLVFSL DVRNIIKGSHYVLYGLVAAMQPRM LVTDFEELRPLPVSVRVGAQVDVVG QAGKPKTITGFQHTTTPVLLAHGERA ELATEEFLPVTPILEGFVILRKNPNY DL	True	False	4.69	4.611	1.635	2.139	1.303