

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
A0A1D5RM18	A0A1D5RM18_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MEPEDLPWPDELEEEEEEEEEEGEE EEGKKEVENASAAATEEALTSEESG RLEEFEEAGPDLDFNYESQRQESSD EEEDELAKAWLQAHPDRPGSAFSLP PPTPPPPPLSPRLRYTPVEHLGKT EVVPLTCRVWQQSSYQDNSRAQFS NSSTMLETGVRWVGSEEDQRTESW HCLPQERDSSQTLAMSQTEIGRVEG TEVPDLPSQEGGLPAQSQCPGKKPK LNVLCSPLLVIQDNFAAPDLLLTCL IQDQEEVEPDSLFOQSELEFAPLRGI PDKSEDSWELARPSEVSEALIQTSE TSSDLANSFCFSISQHLTGSTTFGSQ HSLFMFEQENNGETVPSNAPDELK SPKDSDDHDLVLCYSMSWPSEKDTQQ PEGNLADKDQVSTSTLSDISDESIAA KRNNSEFGVSCSCNQYWNQKALQQT EACLASKDHASFPCKITPLPKVYPS FLQRMLKRGKEAPLIADNHLNENVCL RTPAEDEMSLKNESLEGFEGNEDL QKELSQFFKQKETRNTWASSQVHT RYPEIKYIGTVAVPDSKKQVAEPYILY RGLDISDCPLPTVKSNLDETSEQLYF QEERNQKAASTFDEKNVDATENVA FEAIIASIESSNQESLISQTDTNTEL LADDSEEKEPLKPDLSPLNCKNEISF IDKLKYLKYQSTPGVFEPADSKSLLY KEDDASSLYLYPDLKSSINAPQNML NKPVSVTLELKSSEKACNQAVGL ETQSPSFQYGISNLDLTEKVGSSSIS FAMKIATSESWVLEDLQKQTFEEVA DSSDYKLEGLQGAESGLVTRCGDA KYSSLYENLGAQSERIAVLQREVGCS NLGISQASPSLPSFVPOEPTSEPEY HSSNLRMLRVSPDTLLTTHSAGS ADQKIGAAVVSSAYSQEIKPGSFHQE ELPDRHLNEEIRKVSALRTAGQKP EMLPVQSSYSKGMKSIFYOHPVSH GHQGKEPLSVSAVCGSAGNKAFHQ LSTLSDSLLTEETWPVSVIPGLGNQK TLPSEFSLSYSHRGKNLPEDVVKVS TDSGSAHKKADILTASSRTYQHKMK PANIYHQELPDSRVPIGTRKVAFESG PAGQKSGVSHPYGEMPSVFYQQGLP DRHSAKSPKTFIPGPADQKTDLSPV PPTSSSHAEKPVSPYQLTLPGSHLPE DVFKASSVCSSDELGITALTSASY SYKGRPNSSYQQKFPDLSHNEEAQK ILGTTGTVDQKTVPTMSSFLQKEK PSIFYQQTLPDGGLEEDLQVSAVP WPADQNIAPTIVTSAAFSQREKPRIF YQQLSVDRLPGEPLNVLTSGPPD QNTGAPTVPSSYFPGEESIIFYQAG FPGNTLSAMSFKVPRISSGSTEQTNV TTGSSSSYSVGEKSIIFYHQALPDGR LPQEASAPADLNTGEPMYLASCS VGVKPIIFYQQPMSSDSQRTKGHKE DVPGPTDQKTGIATVHSTSQSYIGRR TVSYQKEFPDLSEKALKVLGDVGSTE QKTQIPVVSSALLHKEGPSAYQEDLP DLTEEPLQILGVSEEVSSSYQRKLP DHIEVFLKSVGSGSADRKTGAQIVSS SREKSSGFHQELPNTGGDAVDAF HPEPVVQEVVRKVQTPGAPAGPSSSH FHKEKLSDYKASPHRDLTESSLKA STVPGLSDQKKKPAVSSGFCLHKEK HEISASALLNCQTAELLTVTQRSCLH REDPAISTVIKPDQKIPLPTTFHGSS DQKVKPVIFVQKQLRDRDQSEDIPKI STVSEPTVVNTVLPVLLPGSYSHREK SDSFYPQELPDGHLTEVDLKVSSGL GOADQISGLPTGIPGTYSHEKHQLI SEHVQELMDNLSSESSCLSVDSM PLNSQIDDGVIICKPESLGFANAGCE EMQNIDRGSKTLKEIQTLLEAENM ALKRCNFVPLVPRFDVNDVVFIRSK	None	None	None	None	None	None	None		

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HIEYVQKDIGTQTNLKYQRGVGNWE  
FISSATFRSPLQEAEGTARMAYDETF  
ROYKAARSVMRSEPEGCGSTGIGNKM  
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TAITFSSHEHSECISDSSGCKVGTA  
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