

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
B2RQG2	B2RQG2_MOUSE	Mus musculus	NaN	41.574385	NaN	NaN	40997131;28528544	MDIVDTFNHLIPTEHLDDALFLGSN LENEVCEDFSTSQNVLEDSLKNMLS DKDPMLGASASQFCLPVLDSNDPN FQMPGSTVVGGLDDIMDEGVVKESE NDTIDEEELILPNRSLRDRVEDNSVR SPRKSRLMAQEQVRSRQSTIAKR SNAATLSTKKPSGKTLSTSKVGVKPA ERCQKGEVYASLKSEHPKESRRSG RHAEQMDVAPEVSASSVSSVSSCA GMKEEAFFDPKHACNNQGEVNVPS PELDCPLLSETSASVEEKNIEMEC KAKTNSPFLFKFPVREDEQSDLVSG ELNDTIEGKDAAGGKPDQESSEVVKFP CEGDQTAEEPESSDVSSDSACANKN KAEKNEGAECHLELKNTVDIVDKPE NSPQRNELETLYGEDTESNDARLQ STEFNKSDLEEVDACAFEPEASTLE NTICDVLQNSKQLNITQSIKMETAL NLQDDRSGLEPKNIKPKHIKSVTHS KQSMTTTETPRKTVAAKHEVGHSKTK SNVKAVKRNSGEPQPDQRPKV RKKQGDKVVKSQCNSGVKSVKSQ AHSVLRMPQDQNTMQISKPLTHP HSDKLHGHSFGFSKEPPHPVQTGHL VHSSQKQSQKQQAAAVGKGSQSHV KDEHDPVSEHLKEDDKLPRKPD RNLQPRRRSSRSFSLDEPPLFIPDN IATVKKEGSDQTTIESKYMWTSPKQ CGFCKKPHGNRFMVGCRDDWF HGDCVGLSLSQAQMGEEKEYVC VRCCAEDKKTDLDEIFEAQAPIE AHSEDKRMECGKLTSSKHAVTDDK HRHSDDPGKHVKILKRESGEGKTS SDSRDNEIKKWQLAPLRKLSQPHLP RRSSEKSEKIAKESTALASTGERVA RSGTHEKQETKKKMEKGGPNVHP PAATSKPSADQIRQSVRHSKLDILMK RLTDSNLKIPKAAKAVATKIEKELF SFFRDTDAKYKNKYRSLMFNLKDPK NNILFKVVKGEVTPDHLIRMSPEEL ASKELAAWRRRNRHTIEMIEKEQR EVERRPITKITHKGEIESDAPMKE QEAAIEIQEPSANKSMEKPDVSEKQ KEEVDSTSKDTTSQHRQHLFDLNCK ICIGRMAPPIDDLSPKTVKVVVGAR KHSNDNEAESLADALSSTTNILTSDF EEEKQESPKSTFSPTPRPEMPGTVE VESTFLARLNFIWGFINMPVAKF VTKAYPVGSPPEYLTELDLPSIQVGG RISPQTVWDYVEKIKASGTKEICVVR FTPVTEEDQISYTLFAYFSSRKYR VAANNMKQVDMYLIPLGAADKIPH PLVPFDGPGLELHRPNLLGLIIRQK LKRPHSASAGPSHTGETPESAPIVLP PDKKGMESCTEAAEESDFFNSF TTVLHKQRNKPSQPLQEDLPTAAEP LMEVTKQEPKPLRFLPGVLIGWDN QPSTLELANKPLPVDLILQSLGTTG QVYEQAQPLVEQSTLKEIFINDQAN PKVEKIDKVEVTEGDAKEIKVAENI SVSTSKNSGEETSSVSSSISPGPLA SLSLRGKPPDVSTEAFLTNLSIPSKQ EESVENKERTLKRLLQDQENSLQD NRTSSDSPCWPGTGKGMDDGDSG SGSGSGSGSEGPVANTRAPQFINLK RDRPQAAGRSQQTASESKDAESCRN GDKHAASAPPHNKEPLAEAVGGEGK LPSQEKSSCQVQNDDEAAPNSSSV ENLSSQAEQANPSQEDVLTQNIET VHPFRRSAPTSSRFEGGNTCQSEF PSKSVSFTCRSSSPRASTNFSMPMRP QQPNLQHLKSSPPGFPPGFQNFPP QNMFGFPPHLSPLLPPPGFGFPON PPMVPPVHVPGQPORMMGPLSQ ASRYMGPQNFYQVDIRRPERRHS PWGRQDQQPDRPFNRGKDRQRF YSDSHHLKRERHDKDWEQESERHR HRDRSQERDRDRKSEAAAAHKDK	True	False	1.641	4.359	0.769	False	False	False	False

