

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
E9QPE7	E9QPE7_MOUSE	Mus musculus	NaN	26.781227	NaN	NaN	36064721;34887587	MAQKQQLSDDEKFLFVDKNFMNSP MAQADWVAKKLVVWPSEKQGFEEA SIKEEKGDEVVVELVENGKKVTVGK DDIQKMNPPKFSKVEDMAELTCLN EASVLHNLRRERYFSGLIYTYSGLFCV VVPYKYLPIYSEKIVDMYKGGKRHE MPPHIYAIADTAYRSMLQDREDQSIL CTGESGAGKTENTKKVIQYLAVVASS HKGKKDSSITGELEKQLQANPILEA FGNAKTVKNDNSSRFKGFIRINFDV TGYIVGANIETYLLEKSRAIRQARDE RTFFHIFYLLAGAKEKMKSDLLESF NSYTFLSNGFVPIPAAQDDEMFQET LEAMSIMGFNEEEQLAILKVVSSVL QLGNIVFKKERNTDQASMPDNTAA QKVCHLVGINVTDFTRAILTPRIKVG RDVVQKAQTKEQADFAIEALAKATY ERLFRWILSRVNKALDKTHRQGASF LGILDIAGFEIFEVNSFEQLCINYTNE KLQQLFNHTMFILEQEEYQREGIEW NFIDFGLDLQPCIELIERPNNPPGVL ALLDEECWFPKATDKSFVEKLCSEQ GNHPKFKPKQLDKKTEFSIHYAG KVDYNASAWLTKNMDPLNDNVTSL LNASSDKFVADLWKDVRIVGLDQ MAKMTESSLPSASKTKKGMFRTVG QLYKEQLGKLMTTLRNTTPNFVRCII PNHEKRSGLDAFLVLEQLRCNGVL EGIRICRQGFNRIQFQEFQRQYEILA ANAIKGFMDGQACILMIKALELD PNLYRIGQSKIFFRTGVLAHLEEERD LKITDVIMAFQAMCRGYLARKAFTK RQQQLTAMKVIQRNCAAYLKLNRW QWWRLFTVKPLLOVTRQEEEMQA KEEEMQKIKERQQAETELKELEQK HTQLAEEKTLLEQLQAETELYAEA EEMRVRLAACKQLEELHEMEARL EEEEDRGQQLQAERKKMAQQMLDL EEQLEEEAARQKLQLEKVTAEAKIK KLEDDILVMDQNSKLSKERKLLLE RVSDLTTNLAEEEEKAKNLTCLKSK HESMISELEVRLKKEEKSRQLEKLE KRKLEGDASDFHEQIADLQAQIAEL KMQLAKKEEELQAALARLDEEIAQK NNALKKIRELEGHISDLQEDLDSER AARNKAQKQKRDLEGELEALKTELE DTLDSTATQQLRAKREQEVTVLKK ALDEETRSHEAQVQEMROKHTQAV EELTEQLEQFKRAKANLDKSKQTLLE KENADLAGELRVLGQAKQVEVEHKK KKLEVQLQDLQSKSDGERARAELS DKVHKLQNEVESVTGMLNEAEGKAI KLAKDVASLGSQLODTQELLQEETR QKLVNSTKLRQLEDERNLSLQDQLDE EMEAKQNLERHVSTLNIQLSDSKKK LQDFASTIEVMEEGKRLQKEMEGL SQYEEKAAAYDKLEKTKNRLQQL DDLVDLDNRQQLVSNLEKKQKFF DQLLAEEKNISKYADERDRAEAEA REKETKALSLARALEEALAEKEELER TNKMLKAEEDLVSSKDDVGNVH ELEKSKRALETQMEEMKTQLEELED ELQATEDAKLRLEVNMQALKGQFE RDLQARDEQNEEKRRQLQRLHEY ETELEDERKQRAAAAANKKLEGLD KDLELQADSAIKGEEAIKQLRKLQA QMKDFQRELDARASRDEIFATSK NEKKAASLEADLMQLQEDLAAAERA RKQADLEKEELAEELASSLSGRNTL QDEKRRLEARIAQLEEEEEEQGNM EAMSDRVRKATLQAEQLSNELATER STAQKNESARQQLERQNKELRSKLQ EVEGAVKAKLKSTVAALEAKIAQLEE QVEQEAAREKQAATKSLKQDKKLE VLLQVEDERKMAEQYKEQAEKGN KVKQLKRQLEEEAEEESQRINANRRK LQRELDEATESNEAMGREVNALKS	None	None	None	None	None	None	None	None	

