

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
K3W4R2	K3W4R2_MOUSE	Mus musculus	NaN	24.557655	NaN	NaN	36064721;37507081	MAAVTMSVSRGKVASRPGVPVEAAQ SFLYAPRTPNVGGPGGQVEWTARR MVVWVPSLHGFEAAALRDEGEDEEA EVELAESGRRLRLPRDQIQRMNPPK FSKAEDMAELTCLNEASVLHNLRE YSGLIYTYSGLFCVVINPYKQLPIYT EAIVEMYRGKKRHEVPPHYAVTEG AYRSMQLDREDQSILCTGESGAGKT ENTKKVIQYLAAHVASSPKGRKEPGVP VSHTALGHGELERQLLOANPILEAF GNAKTVKNDNSSRFGKFRINFDIAG YIVGANIETYLLEKSRRAIRQAKDECSF HIFYQLLGGAGEQLKADLLEPCSHY RFLTNGPSSSPGQERELFQETLES LR VLGLLPEITAMLRTVSAVLQFGNIV LKKERNTDQATMPDNTAAQKLCRL LGLGVTDFSRALLTPRIKVGDRDYVQK AQTKEQADFALEALAKATYERLFRW LVLRLNRALDRSPRQASFLGILDIA GFEIFQLNSFEQLCINYTNEKLLQFL NHTMFVLEQEEYQREGIPWTFDFG LDLQPCIDLIERPANPPGLLALLDEE CWFPKATDKSFVEKVAQEQGSHPKF QRPRNLRDQADFVSLHYAGKV DYKA SEWLKMNMDPLNDNVAALLHQST DRLTAEIWKDVEGIVGLEQVSSLGD GPPGGRPRGMFRITVQGLYKESLSR LMATLSNTNPSFVRCIVPNHEKRAG KLEPRLVLDQLRCNGVLEGRICRQG FPNRILFQEFRQRYEILTPNAIPKGF MDGKQACEKMIQALELDPNLYRVG QSKIFFRAGVLAQLEERDLKVTDII VSFQAAARGYLARRAFQRRQQQSA LRVMQRNCAAYLKLNRWQWWRLF KVKPLLVQVTRQDEVLQARAQELQKV QELQQSAREVGEVLEQGRVAQLEEEER TRLAEQLRAEALCSEAEETRARLAA RKQLELVVTELEARVGEEECSRQ LQSEKRLQHQHIELESHLEAEEGA RQKLQLEKVTTEAKMKFEDL LLL EDQNSKLSKERRLLEERLAEFSSQA AEKKEKVKSLNKLRLKYEATISDME DRLKKEEKGRQELEKLRRLDGESE ELQEQMVEQKQRAEELLAQLGRKE DELQAALLRAEEEGGARAQLLKS LR EAQAGLAEAEQEDLEAERVARAKAEK QRRDLGEELEALRGELEDTL DSTNA QQELRSKREQEVTELEKKALEESRA HEVSMQELRQRHSQALVEMAEQLE QARRGKGVWEKTRLSLEAEVSELKA ELSSLQTSRQEGEGKRRRLESQLE VQGRSSDSEARSEAAEKLQRAQAE LESVSTALSEAESKAIKRLGKELSSAE SQLHDTQELLQEETRAKLALGSRVR ALEAEAAGLREQMEEVVARERAGR ELQSTQAQLSEWRRRQEEEA VLEA GEEARRRAAREATLTQRLAEKTEA VERLERARRRLQQLDDATVDLGQQ KQLLSTLEKKQRKFDQLLAEEKAAV LRAVEDREIEAEGRERARALS LTR ALEEEQEAARELERQNRALRAE LEA LLSSKDDVGNVHELERARAKAEQA ASDLRTQVTELEDELTAEDAKLRLE VTVQALKAQHERDLQGRDDAGEER RRQLAKQLRDAEVRDEERKQORAL MAARKLELELEELKAQTSAAAGQK EEAVKQLKMQVQMKELWREVEET RSSRDEMFTLSRENEKLLKGLEAEV LRLQEEAASDRARRQAQQRDEM AEEVASGNLSKAATLEEKRLQLEGR SLEEELEEEQNNSELLKDHYRKL LQVESLTTLSAERSFSAKAESGRQ QLERQIQELRARLGEE DAGARAROK MLIAALESKLAQAEQLEQESRERIL SGKLVRAEKRLKEVVQVDEERRV ADQVRDQLEKSNLRLKQLEKQLEEA EEEASRAQARRRLQRELEDTVTEA ESMNREVTTLRNLRRLRGLTFTTTRT	None	None	None	None	None	None	None		

