

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
E9QAQ7	E9QAQ7_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MAAQVAPAAASLGNPPPPSELKK AEQQQREEAGGEAAAAAERGMK AAAGQESEGPVGPPLGKELQDG AESNGGGGGGAGSGGGPGAEPDL KNSNGNAGPRPALNNLPEPPGGG GGGGSSSDGVGAPPHSAAAALPPP AYGFGQAYGRSPSAVAAAAAVFHQ QHGGQQSPGLAALQSGGGGLEPY AGPQONSHDHGFPNHQYNSYYPNR SAYPPPPQAYALSSPRGGTPGSGAAA AAGSKPPSSASASSSSSFAQQRF GAMGGGGPSAAGGGTPQPTATPTLN QLLTSPSSARGYQGYPGGDYGGGPQ DGGAGKGPADMASQCWGAAAAAAA AAAAVSGGAQQRSHHAPMSPGSSG GGGQLARTPQSSSPMDQMGKMRP QPYGGTNPYSQQQPPSGPQQGHG YPGQPYGSQTPQRYPMTMQGRAQS AMGSLSYAQIPPYGQQGSPAYGQQ GOTPYYNQQSPHPQQQPPYAQQPPS QTPHAQPSYQQPQTQQPQLQSSQP PYSQQPSQPPHQQSPTPYPSQQSTT QQHPQSQPPYSQPAQSPYQQQP QQPASSLSQAAYPQPQSQSQQT AYSQQRFPQPQLSQDSFGSQASSA PSMTSSKGGQEDMNLSQLSRPSSLP DLSGSDDLPMGTGALSPGVSTSGI SSSQGEQSNPAQSPFSPHTSPHLPGI RGPSFSPVGSASVAQSRSGPLSPAA VPGNQMPRPPSGQSDSIMHPSMN QSSIAQDRGYMQRNPQMPQYTSPO PGSALSPRQSGGMHSGVGSYQQ NSMGSYGPQGSQYGPQGGYPRQPN YNALPNANYPNAGMAGSMNPMGA GGQMHGQPGIPPYGLPPGRMAHA SMGNRPYGNMANMPPQVGS GMC PPPGGMNRKTOESAVAMHVAANSI QNRPPGYPNMNQGGMMGTGPPYG QGINSMAGMINPQGGPPYPMGGTMA NNSAGMAASPEMMGLGDVKLTPAT KMNNKADGTPKTESKSKSSSTTT NEKITKLYELGGEPEPKMWVDRYLA FTEEKAMGMTNLPVGRKPLDLYRL YVSVKEIGGLTQVKNKKWRELATN LNVGTSSAASLKKQYIQCLYAFEC KIERGEDPPPDFAAADSKKSQPKIQ PPSPAGSGSMQGPQTPQSTSSMAE GGDLKPPTASTPHSQIPPLPMRS NSVGIQDAFPDGSDFTFQKRNSMTP NPGYQPSMNTSDMMGRMSYEPNK DPYGSMRKAPGSDPFMSSGQGPNG GMGDPYSRAAGPGLGSVAMGPRQH YPYGGPYDRVRTEPGIGPEGNMGTG APQPNLMPSTPDSGMYSPSRYPPQQ QQQQQQQQQRHDSYGNQFSTQ GTPSSSPFSQQTTMYQQQQQNYKR PMDGTYGPPAKRHEGEMYSVPYSAG QQQPQQQLPAAQSQPASQPAAAQ PSPQQDVYNQYSNAYPASATAADR RPAGGPQNQFPFQFGRDRVSAPPQS SAQQNMPMMGGPIQASAEVAQQ GTMWQGRNDMTYNYANRQNTGSA TQGPAYHGVNRTDEMLHTDQRANH EGPWPSHGTRQPPYGPSAPVPPMTR PPPSNYQPPPSMPNHIPQVSSPAPLP RPMENRTSPSKSPFLHSGMKMQKA GPPVPASHIAPTVPQPPMIRRDITFPP GSVEATQPVLKQRRRLTMKDITPE AWRVMSLSKSGLLAESTWALDTINI LLYDDNSIMTFNLSQLPGLLELLVEY FRRCLIEIFGILKEYEVDGQRTLLD PGRFTKVYSPAHEEEEEHLDPKL EEEEEEGVGNDEEMAF LGDKDPSS ENNEEKLVSKFDKLPVKIVQRNDPF	None	None	None	None	None	None	None		

VVDCSDKLGRVQEFDSGLLHWRIGG
GDTTEHIQTHFESKIELLPSRPYVPC
PTPPRKHLTTVEGTPGTTEQEGPPPD
GLPEKRITATMDDMLSTRSSTLTDE
GAKSAEATKESKFPFGISPAQSHRN
IKILEDEPHSKDETPLECTLLDWQDSL
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KHPGLLLILGKLLHKKHPERKQAP
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SICLPVLDGLLHWAVCPSEAEADPF
STLGPNAVLSPQRLVLETLKLSIQD
NNVDLILATPPFSRLEKLYSTMVRFL
SDRKNPVCREMAVVLLANLAQGDS
LAARAIQVQKGSIGNLLGFLEDSLAA
TQFQSQASLLHMQNPPFEPTSVD
MMRRAARALLALAKVDENHSEFTL
YESRLDISVPLMNSLVSQVICDVL
FLIGQS