

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A0A0R4J0K2	A0A0R4J0K2_MOUSE	Mus musculus	NaN	15.841233	NaN	NaN	37507081	MGDDSEWLKLPVDQKCEHKLWKAR LSGYEEALKIFQKIKDEKSPEWSKYL GLIKKFVTDNAVQKLGLEAALVYV ENAHVAGKTTGEVVSQVSKVFNQP KAKAKELGIEICLMEVEIEKGESVQE ELLKGLDNKNPKIIVACIETLRKALSE FGSKIISLKPIIKVLPKLFESRDKAVR DEAKLFAIEIYRWNRDVAKHTLQNI NSVQLKELEEEWVKLPTGAPKPSRF LRSQQELEAKLEQQQSAGGDAEGG GDDGDEVPQVDAYELLDVEILSKLP KDFYDKIEAKKWQERKEALEAVEVL VKNPKLEAGDYADLVKALKKVVGKD TNVMLVALAAKCLTGLAVGLRKKFG QYAGHVVPVPTILEKFKKPKQVVQAL QEIDAIFLTTTLQNISEDVLAVMDN KNPTIKQQTSLFIARSFRHCTSSTLP KSLKPFCAALLKHINDSAPEVRDAA FEALGTALKVVGEKSVNPFLADVDK LKLDRICEKSEKVELVHGKKSGLATE KKESKPLPGRASGAAGDKDTKDV SGPKPGPLKKTPTAKAGGPSKKGKTT APGGSASAGTKNKKGLETKEIVEPEL SIEVCEEKASAVLPPTCIQLLDSSNW KERLACMEEFQKAVELMERTEMPC QALVKMLAKKPGWKETNFQVMQM KLHIVALIAQKGNFSKTSAQIVLDGL VDKIGDVKCGNNAKEAMTAIAEACM LPWTAEQVMSMAFSQKNPKNQSET LNWLSNAIKEFGFSELNVKAFISNV KTALAATNPAVRTSAITLLGVMYLYV GPSLRMIFEDEKPAALLSQIDAETFQK MQGQSPAPTRGIAKHSTSATDEGE DGEEPGEGGNDVVDLLPRIEISDKIT SELVSKIGDKNWKIRKEGLDEVAGII NEAKFIQPNIGELPTALKGRLNDSN KILVQQLNILQQLAVAMGANIRQH VKNLGPVITVLGDSKNNVRAAALAT VNAWAEQTGMKEWLEGEDLSEELK KENPFLRQELLGWLAEKPLTLRSTP TDLILCVPHLYSCLEDRNGDVRKKA QDALPFFMMHLGYEKMAKATGKLLK PTSKDQVLAMLEKAKANMPSKPAAP AKAMSKPMGGSAPAKTQPIAPVED SVSSTIEAKPDLKKAKAPGVSSKAKS VQGKKVPSKTTLKEDDDKSGPIFIVV PNGKEQRMREKGLKVLKWNFTTP RDEYIEQLKTQMSTCVAKWLQDEM FHSDFQHHNKALAVMVDHLESEKD GVISCLDLILKWLTLRFFDTNTSVLM

KALEYLKLLFTLLSEEEYHLTENEAS  
SFIPYLILKVGEPKDVIRKDVRAILNR  
MCLVYPASKMFPPFIMEGTKSKNSKQ  
RAECLEELGCLIESYGMNVCQPTPG  
KALKEIAIHIGDRDNAVRNAALNTIV  
TVYNVHGDQVFKLIGNLSEKDMSM  
LEERIKRSAKRPSAAPVKQAEKQOR  
TONINSNANMLRKGPAEDMSSKLN  
QARSLSGHPEAAQMVRREFQLDLD  
EIENDNGTVRCPELVQHKLDDIF  
EPVLIPEPKIRAVSPHFDDMHSNTAS  
TINFIISQVASGDINTSIQALTOIDEVL  
RQEDKAEAMSGHIDQFLIATFMQLR  
LIYSTMMADEKLDKDEIIKLYSCHIIGN  
MISLFQIESLAREASTGVLKDLMHG  
LITLMLDSRIEDLEEGQQVIRSVNLL  
VVKVLEKSDQTNILSALLVLLQDSLL  
ATASSPKFSELVMKCLWRMVRLLPD  
TINSINLDRILLDIHIFMKVFPKEKIK  
QCKSEFPRTLKTLHTLCKLKGPKIL  
DHLTMIDNKNESELEAHLCRMVKH  
SMDQTGSKSDKETEKGASRIDEKSS  
KAKVNDFLAEIFKKIGSKENTKEGLA  
ELYEYKKYSDTDIEPFLKNSSQFFQ  
SYVERGLRVIEMERESKGRIPSTGN  
TNGEEVGPSVYLERLKILRQRCGLD  
NTKQDDRPLTSLLSKPAIPVASST  
DMLHSKLSQLRESREQHQHSDLDS  
NQTHSAGTMTSSSSTTNIIDLKRL  
ERIKSSRK