

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
F1M949	F1M949_RAT	Rattus norvegicus	NaN	12.481722	NaN	NaN	18683930	MGDDSEWLKLPVDQCEHKLWKR LSGYEEALKIFQKIKDEKSPEWSKYL GLIKKFVTDSDNAVQLKGLEAALVYV ENAHVAGKTTGEVVSQVSKVFNQP KAKAKELGIEICLMYVEIEKGESVQE ELLKGLDNKNPKIIVACIETLRKALSE FGSKIILLKPIIKVLPKLFESRDKAVR DEAKLFAIEIYRWNRDVAKHTLQNI NSVQLKELEEEWVKLPTGAPKPSRF LRSQQELEAKLEQQQSAGGDAEGG GDDGDEVPOIDAYELLEAVEILSKLP KDFYDKIEAKKWQERKEALEAVEGL VRSPRLEAGDYAELVKALKKVVGKD TNVMLVALAAKCLTGLAVGLRKKFG QYAGHVVPTILEKFKEKKPQVVQAL QEIDAIFLTTTLQNISEDVLAVMND KNPTIKQQTSLFIARFRHCTSSTLP KSLKPFCAALLKHINDSAPEVRDAA FEALGTALKVVGKTVNPFADVDK LKLDKIKECSEKVELVHGKKSGLAAE KESKPLPGRAAASGAAGDKDTKDV SGPKPGPLKKAPATKAGGPPKKGKT TAPGGSASSGTKNKKGLETKEIAEPE LSIEVCEEKASAVLPPTCIQLDSSN WKERLACMEEFQKAVELMERSEMP CQALVKMLAKKPGWKETNFQVMQ MKLHIVALIAQKGNFSKTSAQIVLDG LVDKIGDVKCGNNAKEALTAIAEAC MLPWTAEQVMSMAFSQKNPKNQS ETLNWLSNAIKEFGFSELNVKAFIS NVKTALAATNPAVRTSAITLLGVMYL YVGPSLRMIFEDEKPALLSQIDAQEFQ KMQQQSPPAPTRGISKHSTSGTDEG DDGDEPGDGSNDVVDLLPRVEISDK ITSELVSKIGDKNWKIRKEGLDEVAG IINEAKFIQPNIGELPTALKGRLNDS NKILVQQLNILQQLAVAMGPNVKQ HVKNLGIPLTVLGDSKNNVRAAALA TVNAWAEQTGMKEWLEGEDLSEEL KKENPFLRQELLGWLAEKLPMLRST PTDLILCVPHLYSCLEDRNGDVRKK AQDALPFFMMHLGYEKMAKATGKL KPTSKDQVLAMLEKAKANMPSKPAA PSKAVSKPMGGSAPAKTQPIAPVED SVSNTMEAKPDPKAKAPGVSSKAK SAQGKKVPSKTTTLKEDDDKSGPIFIV VPNGKEQRMREDEKGLKVLKWNFTT PRDEYIEQLKTQMSSCVAKWLQDE MFHADFQHHNKALAVMVDHLESE KDGVISCLDLILKWLTLRFFDTNTSV

LMKALEYLKLLFTLLSDEEYHLTEN
EASSFIPYLILKVGEPKDVIRKDVRAI
LNRMCCLVYPASKMFPPFIMEGTKSKN
SKQRAECLEELGCLIESYGMNVCQP
TPGKALKEIAIHIGDRDNAVRNAALN
TIVTVYNVHGDQVFKLIGNLSEKDM
SMLEERIKRSAKRPSAVPVKQAEKPK
QRTQNINSNANMLRKGPAEDMSSK
LNQARSMGHPAAQMVRREFQLD
LDEIENDNGTVRCEMPELVQHKLD
DIFEPVLIPEPKIRAVSPHFDDMHSN
TASTINFIISQVASGDISTSIALTQID
EVLREQEDKAEAMSGHIDQFLIATFM
QLRLIYSTHMADEKLDKDEIHKLYSCI
IGNMISLFQIESLAREASTGVKDLML
HGLITLMLDSRIEDLEEGQQVIRSVN
LLVVKVLEKSDQTNILSALLVLLQDS
LLATASSPKFSELVMKCLWRMVRL
PDTINSINLDRILLDIHIFMKVFPKEK
LKQCKSEFPRTLKTLHTLCKLKGP
KILDHLT MIDNKNESELEAHL CRM
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EKSSKAKVNDFLAEIFKKIGSKENTK
EGLAELYEYKKKYSDDIEPFLKNSS
QFFQSYVERGLRVIEMERESKGRIP
SAGISPMQMEVTCVPTPTSTVSSLGNT
NGEEVGPSVYLERLRLKILRQRCGLDN
TKQDDRPPLTSLLSKPAIPPVASTD
MLHSKLSQLRESREQHQHSDLDSN
QTHSAGTMTSSSSTANIDDLKRRLE
RIKSSRK