

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
G3V8B0	G3V8B0_RAT	Rattus norvegicus	NaN	9.894158	NaN	NaN	34502162	MADAEMAAFGAAAPFLRKSEKERLE AQTRPFDLKKDVFVPDDKEEFVKAKI VSREGGKVTAETENGKTVTVKEDQV MQQNPPKFDKIEDMAMLTFLHEPA VLYNLKERYASWMIYTYSGLFCVTV NPYKWLPVYNAEVVAAYRGKKRSEA PPHIFSISDNAYQYMLTDRENQSILIT GESGAGKTVNTRKVIQYFAVIAAIGD RSKKDQTPGKGTLEDQIIQANPALEA FGNAKTVRNDNSSRFGKFIHFGA TGKLASADIETYLLEKSRVIFQLKAER DYHIFYQILSNKKPELLDMLLITNPNP YDYAFISQGETTVASIDDSEELMATD SAFDVLGFTPEEKNSIYKLTGAIMHF GNMKFKQKQREEQAEPDGTTEADK SAYLMGLNSADLLKGLCHPRVKVG NEYVTKGQNVQQVAYAGALAKSVY EKMFNWMVTRINATLETQPRQYFI GVLDIAGFEIFDFNSFEQLCINFTNE KLQQFFNHMHMFVLEQEEYKKEGIE WTFIDFGMDLQACIDLIEKPMGIMSI LEEEMFPKATDMTFKAKLYDNHL GKSNNFQKPRNIKKGQEAHFSLIHY AGTVDYNILGWLQKNKDPLNETVVG LYQKSSLKLLSNLFANYAGADAPVD KGKGKAKKGSSFQTVSALHRENLNK LMTNLRSTHPIHFVRCIIPNETKSPGV MDNPLVMHQLRCNGVLEGRICRK GFPNRILYGDFRQRYRILNPAAPPEG QFIDSRKGAEKLLGSLDIDHNQYKF GHTKVFFKAGLLGLLEEMRDERLSR IITRIQAQSRGVLSRMEFKKLLERRD SLLIIQWNIRAFMGVKNWPWMKLY FKIKPLLKSAETEKEMANMKEEFGR VKDALEKSEARRKELEEKMVSLLOE KNDLQLQVQAEQDNLADAEERCDQ LIKNIQLEAKVKEMTERLEDEEEM NAELTAKKRKLEDECESELKRDIDDL ELTLAKVEKEKHATENKVKNLTEEM AGLDEIIVKLTKEKKALQEAHQALD DLQAEEDKVNTLTKAKVKLEQQVDD LEGSLEQEKVVRMDLERAKRKLEGD LKLTQESIMDLENDKQQLDERLKKK DFELNALNARIEDEQALGSQLOKKL KELQARIEELEELEAERTARAKVEK LRSDLSRELEEISERLEEAGGATSVQ IEMNKKREAEFQKMRRDLEEATLQ HEATAAALRKKHADSVAELGEQIDN LQRVKQKLEKEKSEFKLELDDVTSN MEQIIKAKANLEKMCRTLEDQMNE

HRSKAEETQRSVNDLTSQRAKLQTE
NGELSRQLDEKEALISQLTRGKLYT
QQLEDLKRQLEEEVKAKNALAHALQ
SARHDCDLLREQYEEETEAKAELQR
VLSKANSEVAQWRTKYETDAIQRTE
ELEEAKKKLAQRLQDAEEAVEAVNA
KCSSLEKTKHRLQNEIEDLMVDVER
SNAAAAALDKKQRNFDKILAEWKQK
YEESQSELESSQKEARSLSTELFKLK
NAYEESLEHLETFKRENKNLQEEIS
DLTEQLGSTGKSIHELEKIRKQLEAE
KLELQSALEEAASLEHEEGKILRAQ
LEFNQIKAEIERKLAEKDEEMEQAK
RNHLRVVDSLQTSLDAETRNRNEAL
RVKKKMEGDLNEMEIQLSHANRMA
AEAQKQVKSLSLLKDTQIQLDDAV
RANDDLKENIAIVERRNLLQAELE
ELRAVVEQTERRSRKLAEQELIETSER
VQLLSQNTSLINQKKKMDADLSQL
QTEVEEAVQECRNAEEKAKKAITDA
AMMAEELKKEQDTS AHLERMKKN
MEQTIKDLQHRLDEAEQIALKGGKK
QLQKLEARVRELENELEAEQKRNAE
SVKGMRKSERRIKELTYQTEEDRKN
LLRLQDLVDKLQLKV KAYKRQAEEA
EEQANTNLSKFRKVQHELDEAEERA
DIAESQVNKLRAKSRDIGAKGLNEE