

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
D4A0B4	D4A0B4_RAT	Rattus norvegicus	NaN	28.175057	NaN	NaN	18683930	MVETAAMEAYVLEDILEVESFILDQ DDLENPMLETASKLLLSGTADGADL RTVDPETQARLEALLEAAGIGKLSA DGKAFADPEVLRRLTSSVSCALDEA AAALTRMRAESTANAGQSDNRSLAE ACSEGDVNAVRLKLLIEGRSVNEHTE EGESLLCLACSAGYYELAQVLLAMH ANVEDRGIKGDITPLMAAANGGHVK IVKLLLAHKADVNAQSSGTNTALTYA CAGGYVDVVKVLLSESGASIEDHNEN GHTPLMEAGSAGHVEVARLLENG AGINTHSNEFKESALTLACYKGHLE MVRFLLEAGADQEHKTDEMHTALM EACMDGHVEVARLLLDGSAQVNMP ADSFESPLTLAACGGHVELAALLIER GASLEEVNDEGYTPLMEAAREGHEE MVALLLGGGANINAQTETQETALT LACCGGFLEVADFLIKAGADIELGCS TPLMEAAQEGHLELVKYLLAAGANV HATTATGDTALTYACENGHTDVADV LLQAGADLEHESEGGRTPLMKAARA GHVCTVQFLISKGANVNRTTANNND HTVLSLACAGGHLAVVELLLAHGAD PThrLkDgStmLIEAAKGgHtSVVC YLLDYPNNLLAAPPDVTQLTPPSHD LNRAPRVVQALPMVVPVQEPDKPP ANVAATLpvrnkaaskQksnshlpa NSQDVQGYITNQSPESIVEEAQGKLT ELEQRKEAIEKNAQLQSLLELAHADQ LTKEKIEELNKTRFQIQKQKILEE LQKVERELQKTKQQLKQOYLEVKA QRIQLQQQQQQQQQQQQQQQQQQ QQQQQQQQQQQQQSCQHLGLLTSV GVGEPLPEGDYARLQVDPVLLKDE PQQTAAQMGFAIQPLAMPQALPLA AGPLPPGSIANLTELQGVIVGQPVLG QAQLAGLGQGILTETQQGLMVASPA QTLNDTLDDIMAVSGRASAMSNTP HSIAASVSQPQTPTPSPIISPSAMLP YPAIDIDAQTESNHDTALTLACAGGH EELVQTLLERGASIEHRDKKGFPTLI LAATAGHVGVVEILLDNGADIEAQSE RTKDTPLSLACSGGRQEVVELLLAR GANKEHRNVSDYTPLSLAASGGYVN IIKILLNAGAEINSRTGSKLGISPLML AAMNGHTAAVKLLLDMGSDINAQIE TNRNTALTLACFQGRTEVVSLLDR KANVEHRAKTGLTPLMEAASGGYAE VGRVLLDKGADVNAAPPVSSRDAL TIAADKGHYKFCCELLIGRGAHIDVRN KKGNTPLWLAANGGHLDVQQLLVQ AGADVDAADNRKITPLMAAFRKGHV KVVRYLVKEVNQFSPDSECMRYIATI TDKLRRESRRLALAAREKRKEKRR KKKEEQRRLLEEIEAKNKENFELQA AQEKEKHKVEEPEVLTPEPSATTTT TIGISATWTTLAGSHGKRNNTTTTT SKRKNRKNKITPENVOIIFDDPLPISY SQPEKASGESKSSSTSESGDSNMR ISSCDESSNSNSRSDNHSPAVA TATVASKKQPSVLVTFPKEERKSVSG KASIKLSETVNEGTSNSLSTCTKSGP SPLSSPNGKLTVASPKRGPKREGW KEVVRRGGTESTRQATQLINALIKDP DKEIDELIPKNRLKSSANSKIGSSA PTTTAANSSLMGKMTSVALSSTSOT ATALTPAVSSASTHKTIKNPVNNVR PGFPVSLPLAYPPQFAHALLAAQTF QQIRPPRLPMTHTGGTFPPAQSTWG FPVVRPLSPARATNSPKPHMVRHS NQNSSGSQVNSAGSLTSSPTTTTSS SASAVPGTTSNGSPSSPSVRRQLFVT VVKTSNATTTTITASNNSTAPTNA	None	None	None	None	None	None	None		

TYPMPTAKEHYVSSPSSPSPAQPG
GVSRRNSPLDCGSASPNGASSSEQE
ASSPPVETTNSRPSHSSTSSGSSSG
HSTQQPPGSPVQEP RPPLQSQV
SPDVRMTVPPTATSSAPVVVPSTAP
MTYPMPTQMGCSQPPKMETPAIR
PPSHGTTAPHKNSAPVQNSSVAVLN
VNHKRPHSVPSVQLPSTLSTQSAC
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TLFENNPTNAHAFWGGPVVSSQSTP
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PGFRPPLQRPAPSPSESPAQSVSSGV
RAPSPAPSSVPLGSEKPSVSDRKY
PVPIGTERSARIRQTGTSAVSVGSNL
STSVGHSGIWSFEGIGGNQDKVDW
CNPGMGNPMIHRPMSDPGVFSQH
QAMERDSTGIVTPSGTFHQHVPAGY
MDFPKLGSMFPFSVYGNAMLPVAPI
PDGAGGPIFNGPHSADPSWNSLIK
VSSSTENNGPQTVVWTGPWAPHMNS
VHMNQLG