

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
G3V6E1	G3V6E1_RAT	Rattus norvegicus	NaN	25.563465	NaN	NaN	34502162	MSSDSEMAVFGAAPYLKSEKERI EAQNRPFDAKTSVVFVAEPKESFVKG TIQSKDAGKVTVKTEAGATLTVKEDQ IFPMNPPKYDKIEDMAMMTHLHEP AVLYNLKERYAAWMIYYSGLFCVT VNPYKWLVPVYNPEVVAAYRGKKRQE APPHIFSISDNAYQFMLTDRENQSIL ITGESGAGKTVNTKRVIQYFATIAVT GEKKKEEVTSGKMQGTLEDQIISAN PLLEAFGNAKTVRNDNSSRFGKFIRI HFGTTGKLASADIETYLLEKSRVTFQ LKAERSYHIFYQITSNKKPELIEMLLI TTNPYDYPPFVSQGEISVASIDDQEEL MATDSAIDILGFTNDEKVSIIYKLTGA VMHYGNMKFKQKQREEQAEPDGTE VADKAAYLQGLNSADLLKALCYPRV KVGNEYVTKGQTVQVTVNAV GALAK AMYEKMFLWMVTRINQQLDTKQPR QYFIGVLDIAGFEIFDFNSLEQLCINF TNEKLQOFFNHHMFVLEQEEYKKE GIEWTFIDFGMDLAAACIELIEKPMGI FSILEEECMFPKATDTSFKNKLYEQ HLGKSANFQKPKVVKGAEAHFSLI HYAGTVDYNTGWLDKNKDPLNETV VGLYQKSSMKTLAYLFSGAQTAEAE ASSGGAACKGAKKKGSSFTVSALF RENLNKLMNTLRSTHPHFVRCIIPN ETKTPGAMEHELVLHLRCNGVLE GIRICRKGFP SRILYADFQRYKVLN ASAIPEGQYIDSKKASEKLLGSIDIDH TQYKFGHTKVFFKAGLLGLLEEMRD DKLAQLITRTQAMCRGFLARVEYQK MVERRESIFCIQYNIRAFMNVKHWP WMKLEFFKIKPLLKSAETEKEMATMK EEFQTKDELAKSEAKRKELEEKMV SLLKEKNDLQVQAEAEGLADAE RCDQLIKTKIQLEAKIKEVTERAEDE EEINAELTAKKRKLEDECSSELKDDID DLELTLAKVEKEKHATENKVNLTE EMAGLDETIKLTKEKKALQEAHQ TLDDLQAEEDKVNLTAKAKIKLEQ VDDLEGSLEQEKKLRMDLERAKRKL EGDLKLAQESIMDIENEKQQLDERL KKKEFEMS NLQSKIEDEQAIGIQLK KIKELQARIEELEEIEAERASRAKAE KQRS DLSRELEIEISERLEEAGGATSA QIEMNKKREAEFQKMRRDLEEATL QHEATAATLRKKHADSV AELGEQID NLQ RVKQKLEKEKSEMMEIDDLA SNVETVSKAKGNLEKMCRTLEDQVS ELKSKEEQRLINDLTTQRGRLQT ESGEFSRQLDEKEALVSQLSRGKQA FTQIEELKRQLEEEVKAKNALAHA LQSSRHDCDLLREQYEEQESKAE QRALSKANSEVAQWR TKYETDAIQ TEELEEAKKLAQRLQAAEEHVEAV NAK CASLEKTQRLQNEVEDLMLD VERTNAACAALDKQRNFDKILA EW KQKYEETHAELEASQKEARSLGTEL FKMKNAYEESLDQLETLKRENKLN QQEISDLTEQIAEGGKRIHELEKIKK QVEQEKCELQAAL EEAASLEHEEG KILRIQLELNQVKSEIDRKIAEKDEEI DQLKRNHIRVVESMQSTLDAEIRSR NDAIRIKKKMEGDLNEMEIQLNHA NRMAAEALRNYRNTQGILKDTQLHL DDALRGQEDLKEQLAMVERRANLL QAEIEELRATLEQTERRKIAEQELL DASERVQLLHTQNTSLINTKKKLET DISQIQGEMEDIVQEARNAEKKAKK AITDAAMMAEELKKEQD TSAHLER MKKNMEQTVKDLQLRLDEAEQLAL KGGKKQIQKLEARVRELEGEVESEQ	None	None	None	None	None	None	None		

