

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
D3ZKV7	D3ZKV7_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MDGRDFGQSRVHGPPPLLSGLA MDSHRVGAATAGRLPSSGLPGPPP GKYMAGLNLHPHPGFSHLPGLYPS YLHLNHLEPPSSGSPLLSQLGQPSIF DTQKDGFFLPAAGTLHAHTPSSRTP SGHSSGGTAKGSSREGTKDRSSRG GDPPLFGKKDPRAREEVSGPRGVV DLTQEVRAEGRQDRGSSRLAERLSP FLAEAKAKGALQPSALSLCNGVDA GLVAELSRGGAKEVARQEENARLLR RAEALLPAARPCGSLPPPPPLPK GPPAPSSTPAGVYTVFREPGRHRV VAPTFVPSVEAFDERVGPQIASQAR DVRAREREPGRPGVLQGGSPRLE RPEVLREKSSVIRSLKRPPSDGPPA ARSSRSPDARAYLPPKELLKPEADP RPCERAPRGSSAAQQAALFGL PSRPPGPEHKWKPFELGNFATTQM AVLAAQHHAASRAEEAAVATASKK AYLDPGGAMPASATCGRPGADLHS AAAHGPGASAMQSLIKYSGSFAR AVAVRPGGCGKSPFGLGTMKPEP IPTSAGPPRAQARLTHPGVPTAGGGR QLKRDPERPESAKTFGREGSGAQGE AEVRHPPVGIAVAVARQKDSGSSRL GPGLIDQERSLSLNNVKGHGRTEDD CDRARHREDRLGTRLDQEKLLR ESKELADLARLHPTSCAPNGLNPNL MVTGGPTLAGSGRWSADPAHLAT HPWLPRSGTSMWLAGHPYGLGPP SLHQGMAPAFPPGLGGSLPSAYQFV RDPQSGQLVVIPSDHLPFAELMER ATVPPLWPALYPPGRSPLHHAQQLQ LFSQOHFLRQOELLYLQQAQAQALE LQSAQLVQERLKAQEHRTMEEKI TKRGLETTGKAGLSATGPGLLPQKSA GLATGPTGTHGKAMSPPSPRASPV TSLKAKVIQKVEDVSKPPAYTYPATP SSHPSPPASPPPTPGLTRKEAPE NVVEKDDLEKEAPSPFQALFTDIP PRYPFQALPPHYGRYPFLQPTAAA DADGLAPDVPLPADGPERLALSPEP KPICLSPSKIPEPPRDSQEEERLADR EVKAEIEDIEEGPTELPRLESPLALPV PETMVAVSPAGGCGGSPLEAQUALST AGPGCREPSEVSDFAQVAEPQIELPS RTEPHMAALELGTQLTPEPLVEAKE EPVEVPLDVPMEEPVAETGPEDSLP QPPLTEPPPSLELSDCDLVPPEGQCL NLEAQEAAPAPASTYLEETHPESLLP GLDDPLAGMNALAAAAELPQARPLP SLGTGVPGGKLDTAPSLVLEHSFL QGITLLSEIAELEDRRGQEVGPEP NLVVRPSLESLLAASSHMLKEVLES PLSDSLKNLRLPRELNSNKKYSWM QKKEERMFAKSSLEMDALELDF RMRLAEVQRRYKEKQRELVKLQRR RDSGDRHEDAHRSLTRRGPGRPRK RTHTPSALSPPCKRGKSHSGSGKLS SKSLTSDDYDLGAGIRKRHKGPKEE EQEALVSMGKARSRNQSWDDHESS SDFMSQLKIKKKMSSDQEQSLASKL DRALSLTKQDKLSPFKFSDSPGGK WKTSGGCGRFLTQCDLLGKDRKAL AKGLGLSLKPSREKHKRASKARKM EGGFKARGQPKSVHSPFASEVSSH YNTDSEDEDELLKNSWSAQGPSSSK LTSLLGMVAKNSKATGPKLTKRG PSGPRTLKPEATSRKQSFCLLLREA EAHSSFDSSAEDSFDQDDSSEEEE EEEEEEEEEEEEGIGSYRLGAGEQ ALSPSLEESGLGLLARFAASALPSPV VGPPLSVVQLEAEQKARKKEERQSL	False	False	1.594	3.046	False	False	False	False	

MGTEFEYTDSESEVVKVQSPAGLL
RPPKGAGEPGQSLAAPVPGTRASGP
TSPDKAKLASEKGRKARKIRGSKEPG
FEAGPEASDDDLWTRRRSERIFLHD
ASAAVQAPSNAAPATKPSRCGRGGA
PSPRKDTGRAKDRKDPKRRKRGKEA
GSAATLPSPRVSTLPSRAPHPGALA
TAKRSKAKARGKEAKKENRGKGGAV
SKLMECMAAEEDFEPNQDSSFS
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SPDIYRVVVEGERGNRPHIYCLEQLL
QEAIIDVRPASTRFLPQGTRIAAYWS
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CAEPSALLVPSAKRRSRKTSKDTGE
GKEGAATGPQEATGGKARGRGRKPS
TKAKADRAVVLEEGAATNELPSAPL
ALEPVSTPNSKSTPEPVDKRARAPK
ARSVSVQSPGPPTFSSCPAPEPFGE
LPTPATAPLVTMPVTMPATRPKPKKA
RATEGSGAKGPRRPGEDDELLVKLD
HEGVMSPKSKKAKEALLREDPGPG
AWPESTGLLSLGSYSPTVGSSEPKAT
WPKGLDGDLTQEPGPLEDPGNS
KSPDKAQAEQDGAEESETSSSSSS
SSSSSSSSSSSSSSSSSGSETEGE
EDAEKNREDGRGAGGRTCSAASSRA
SSPASSSSSSSSSSSSSSSSSSSS
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SSCSSDDEAAPAPAAGPSTQFALPTK
VSKPPSKARASAHSPGKAPPTQPP
PQPPPPQQTLPKQTGAGAKSRP
KKREGVHLPTTKELAKRORLPSVEN
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MKGKARKLFYKAIVRGKEMIRIGDC
AVFLSAGRPNLPYIGRIQSMWESWG
NNMVVRVKWFYHPEETSPGKQFHE
GQHWDQKSGHNLPAALRASSORKD
FMERALYQSSHVDENDVQTVSHKC
LVVGLEQYEQMLTKKYQDSEGLYY
LAGTYEPTGMIFSTDGVPVLC