

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
G3V7K5	G3V7K5_RAT	Rattus norvegicus	NaN	23.439218	NaN	NaN	38843836	MGAHHPALGLLLLLLCPAQVFSQS CVWYGECEGVAFGDKRYNCEYSGPP KPLPKDGYDLVQELCPGLFFGNVSL CCDVQQLQTLKSNLQLPLQFLSRCP SCFYNLMTLFCELTCSPHQSQFLNV TATEDYFDPETRENKTNVKELEYV GRSFANAMYNACRDVEAPSSNEKA LGLLCGRDARACNATNWIEYMFNK DNGQAPFTIHPVFSDSLVLGMEPMR NATKGCNESVDEVTGPCSCQDCSAV CGPKPQPPPPVPWRILGLDAMYVI MWVAYMAFLVLFVFGLLAVWCHRR RYFVSEYTPIDSNIAFSSNSSDKGEA SCCDPLGVAFDDCLRRMFTKWGAF CVRNPTCIIFFSLVFIAACSSGLVFVQ VTTNPVELWSAPHSQARLEKEYFDK HFGPFFRTEQLIIQAPNSSVHIYEPYP SGADVPPFGPPLNKEILHQVLDLQIAI ESITASYNNETVTLQDICVAPLSPYN KNCTIMSVLNYFQNSHAVLDNQVG DDFVYADYHTHFLYCVRAPASLND TSLHGPCLGTFGGPPVFPWLVLGGY DDQNYNNATALVITFPVNNYNDTE KLQRAWAWEKEFINFVKNYKNPNL TISFTAERSIEDELNRESNSDVFTVLI SYAVMFLYISLALGHIQSWNRVLVD SKISLGIAGLIVLSSVTCSLGIFSYIG MPLTLVIEVIPFLVLAVGVDNIFILV QTYQRDERLQEETLDQQLGRILGEV APTMFLSSFSETSAFFFGALSSMPA VHTFSLFAGMAVLIDFLLQITCFVSF LGLDIKROEKNRDLILCCVRGPDDG QESQASESYLFRFFKNAPFLLTD WLRPVVMAVFGILSFSIAVVKVEI GLDQSLMPNDSYVIDYFKSLGQYL HSGPPVYFVLEEGYNYSSRRGQNMV CGGMGCDNDSLQVQIFNAAELDTYT RVGFAPSSWIDDYFDVWVSPQSSCCR LYNVTHQFCNASVIDPTCVRCRPLTP EGKQRPOGKEFMKFLPMFLSDNPN PKCGKGGHAAYSsavNIMGDDTYIG ATYFMTYHTILKTSADYIDALKKARLI ASNITETMRKSGSDYRVFPYSVFYVF YEQYLTIIDDAIFNLSVSLGSIFLVTLV ALGCELWSAVIMCLTIAMILVNMFG VMWLWGISLNAVSLVNLVMTCGISV EFCSHITRAFTMSTKGSRVSRAEAL AHMGSSVFSGITLTKFGGIVVLAFK SQIFEIFYFRMYLAMVLLGATHGLIF LPVLLSYIGPSVNAKARHTTQERYKG TERERLLNF	True	True	2.113	4.405	2.094	4.462	4.427	4.787	4.452