	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi appara
0	O75420	GGYF1_HUMAN	Homo sapiens	GRB10-interacting GYF protein 1	36.005466	S155;S634	S24;S28;S137;S157;S230;S341;S406;S538;S638;S862	28657654;23301498;40307207;31492838	MAAETLNFGPEWLRALSGGGSVASP PPSPAMPKYKLADYRYGREEMLALY VKENKVPEELQDKEFAAVLQDEPLQ PLALEPLTEEEQRNFSLSVNSVAVLR LMGKGAGPPLAGTSRGRGSTRSRGR GRGDSCFYQRSIEEGDGAFGRSPREI QRSQSWDDRGERRFEKSARRDGAR CGFEEGGAGPRKEHARSDSENWRS LREEQEEEEEGSWRLGAGPRRDGD RWRSASPDGGPRSAGWREHGERRR KFEFDLRGDRGGCGEEEGRGGGGS SHLRRCRAPEGFEEDKDGLPEWCL DDEDEEMGTFDASGAFLPLKKGPKE PIPEEQELDFQGLEEEEEPSEGLEEE GPEAGGKELTPLPPQEEKSSSPSPLP TLGPLWGTNGDGDETAEKEPPAAED DIRGIQLSPGVGSSAGPPGDLEDDEG LKHLQQEAEKLVASLQDSSLEEEQF TAAMQTQGLRHSAAATALPLSHGAA RKWFYKDPQGEIQGPFTTQEMAEW FQAGYFSMSLLVKRGCDEGFQPLGE VIKMWGRVPFAPGPSPPPLLGNMD QERLKKQQELAAAALYQQLQHQQFL QLVSSRQLPQCALREKAALGDLTPPP PPPPQQQQQQLTAFLQQLQALKPPR GGDQNLLPTMSRSLSVPDSGRLWD VHTSASSQSGGEASLWDIPINSSTQ GPILEQLQLQHKFQERREVELRAKR EEEERKRREEKRRQQQQEEQKRRQ EEEELFRRKHVRQQELLLKLLQQQQ AVPVPPAPSSPPPLWAGLAKQGLSM KTLLELQLEGERQLHKQPPPREPAR AQAPNHRVQLGGLGTAPLNQWVSE AGPLWGGPDKSGGGSSGLGLWEDT PKSGGSLVRGLGLKNSRSSPSLSDSY SHLSGRPIRKKTEEEEKLLKLLQGIP RPQDGFTQWCEQMLHTLSATGSLD VPMAVAILKEVESPYDVHDYIRSCLG DTLEAKEFAKQFLERRAKQKASQQR QQQQEAWLSSASLQTAFQANHSTK LGPGEGSKAKRRALMLHSDPSILGY SLHGSSGEIESVDDY	False	False	3.619	2.391	False	False	False