

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appara
Q6Y7W8	GGYF2_MOUSE	Mus musculus	GRB10-interacting GYF protein 2	47.035857	S672;T681	S19;S26;S30;S140;S161;S190;S237;T383;S594;S995;S1276	22645316;40885482;35822049;36852467	MAAETQTLNFGPEWLRALSSGGGIT SPLSPALPKYLADYRYGREEMLAL FLKDYKIPFDLLEKEFLPILQEEPLPP LALVPFTEEEQRNFSMSVNSAAVLR LTGRGGGGGTVVGAPRGRSSSRGRG RGRGECGFYQRSFDEVEGVFGRGG GREMHRSSQWEERDRRFEKPGRK DVGRPNFEEESGPTSVGRKHEFIRSE SENWRIFREEQNGEDEDGGWRLAG SRRDGERWRPHSPDGPSTGWREH MERRRRFEFDRDRDDERGYRRVR SGSGSIDDDRSLPEWCLEDAEEEM GTFDSSGAFLSLKKVQKEPIPEEQE MDFRPVEEGEERSDSDSSHNEEAK EPDKTNRREGEKTDRAAGAEASEEVP QTSLSARPGTPSDHQPEATQFER KDEPKAEQVEKAEENRSENLSAK VPSRGDETVPASQQPSTPLPPDTASP LLILSPVPTPSSASRPVTAAVEAPG MSSVSTEPDDEEGLKHLEQQAEM VAYLQDSALDDERLTSKLOEHRAGK VSIPLMHEAMQKWWYKDPQGEIQGP FNNQEMAWEWFQAGYFTMSLLVKRA CDESFQPLGDIKMMWGRVPFSPG APPPHMGELDQERLTRQQLTALYQ MQHLQYQQFLIQQYAQVLAQQQK AALSSQQQQLALLLQFQALKMR MSDQNIIPSVTRSVSPDTGSIWELQ PAASQPAVWEGGSVVDLPLDTTAPG PSLEQLQOLEKAKAAKLEQERREAE MRAKREEEERKQEEELRRQEEILR RQEEERKRREEEELARRKQEEALR RQEEERKRRQEEERQEEALR RRLEERRREEEERRKQEEELRKQEE EAAKWAREEEEEAQRREENRLRME EEAARLRHEEEERKRKELELQORQD LMRQRQQQEQEALRRLQQQQQQQ LAQMKLPSSSTWGGQSNATCQSQ ATLSLAEIQKLEERERQLREEORRQ QRELMKALQQQQQQQKLSGW GNVSKPAGTTKSLLEIQQEEARQMQ KQQQQQQQQQHQSNRNARNST HSNLHTSLGNSVWGSINTGPSNQW ASELVSSIWSNADTKNSNMGFWDD AVKEVGPARNSTNKNKNASLSKSV GVSNRQNKVVEEELKLLKLFQGVN KAQDGFQWCEQMLHALNTANNL DVPTFVSFLKEVESPYEVHDYTRAYL GDTSEAKFAKQFLERRAKQVNNQ RQQQQQQQQQDQSVWGMNHSTLH SVFQTNQSNQSNFEAVQSGKKK KKQKMRADPSSLGFSVNASERLN MGEIETLDDY	False	False	3.441	3.349	1.307	2.663	2.0