	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus
0	Q62448	IF4G2_MOUSE	Mus musculus	Eukaryotic translation initiation factor 4 gamma 2	38.093057	T505	S11;T89;S394;S442;T507;T513;S901	36852467;22645316;35822049;36288343	MESAIAEGGASRFSASSGGGGSRGA PQHYPKTAGNSEFLGKTPGQNAQK WIPARSTRRDDNSAANNSANEKER HDAIFRKVRGILNKLTPEKFDKLCLE LLNVGVESKLILKGVILLIVDKALEEP KYSSLYAQLCLRLAEDAPNFDGPAA EGQPGQKQSTTFRRLLISKLQDEFE NRTRNVDVYDKRENPLLPEEEEQRA IAKIKMLGNIKFIGELGKLDLIHESIL HKCIKTLLEKKKRVQLKDMGEDLEC LCQIMRTVGPRLDHERAKSLMDQYF ARMCSLMLSKELPARIRFLLQDTVE LREHHWVPRKAFLDNGPKTINQIRQ DAVKDLGVFIPAPMAQGRSDFFLEG PFMPPRMKMDRDPLGGLADMFGQ MPGSGIGTGPGVIQDRFSPTMGRHR SNQLFNGHGGHIMPPTQSQFGEMG GKFMKSQGLSQLYHNQSQGLLSQL QGQSKDMPPRFSKKGQLNADEISLR PAQSFLMNKNQVPKLQPQITMIPPS AQPPRTQTPPLGQTPQLGLKTNPPLI QEKPAKTSKKPPPSKEELLKLTEAVV TDYLNSGNANDAVSGVREMRAPKH FLPEMLSKVIILSLDRSDEDKEKASS LISLLKQEGIATSDNFMQAFLNVLEQ CPKLEVDIPLVKSYLAQFAARAIISEL VSISELAQPLESGTHFPLFLLCLQQL AKLQDREWLTELFQQSKVNMQKML PEIDQNKDRMLEILEGKGLSFLFPLL KLEKELLKQIKLDPSPQTIYKWIKDNI SPKLHVDKGFVNILMTSFLQYISSEV SPPSDETDSSSAPSKEQLEQEKQLLL SFKPVMQKFLHDHVDLQVSALYALQ VHCYNSSFPKGMLLRFFVHFYDMEI IEEEAFLAWKEDITQEFPGKGKALFQ VNQWLTWLETAEEEESEEEAD	None	None	None	None	None	None	None