	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9NJH0	EF1G_DROME	Drosophila melanogaster	Elongation factor 1-gamma	27.146871	NaN	NaN	15952742	MVKGTLYTYPENFRAYKALIAAQYSG AQVKVADNFKFGETNKSAEFLKKFP GGKVPAFETAEGQYLSESNAIAYLLA NEQLRGGKCPFVQAQVQQWISFAD NEIVPASCAWVFPLLGILPQQKNSTA KQEAEAVLQQLNQKLQDATFLAGER ITLADIVVFSSLLHLYEYVLEPSVRSA FGNVNRWFVTILNQKQVQAVVKDY KLCEKALVFDPKKYAEFQAKTGAAK PQQQAQQQKQEKKPKEKKEAPKKA AEPAEELDAADEALAAEPKSKDPFD ALPKGTFNFDDFKRVYSNEDEAKSI PYFFDKFDAENYSIWFGEYKYNEEL SKVFMSCNLITGMFQRLDKMRKAA FASVCLFGEDGNSTISGIWVWRGQD LAFTLSPDWQIDYEVYDWKKLDAKS EETKKLVTQYFSWSGTDKDGRKFNQ GKIFK	None	None	None	None	None	None	None	None	None