	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	Q04067	EIF3G_YEAST	Saccharomyces cerevisiae	Eukaryotic translation initiation factor 3 subunit G	24.475439	NaN	S131	33229814	MSEVAPEEIIENADGSRSIITYKIEDG VKYKITQKVKEVKVLEKVHKSVAERK NWHKYGSEKGSPAGPSAVTARLGEE VELRLSRNWKQAEEERIQKEKASLT KTGLQCRLCGNDHMTMNCPFKTIL SELSALEDPATNEGGVEAASEEKAG QVGGAGSIPGQYVPPSRRAGARDPS SDAYRDSRERDDMCTLKIMQVNEN ADENSLREELLFPFAPIPRVSVVRNK ETGKSRGLAFVTFSSEEVAEQALRFL DGRGYMNLILRVEWSKPKVKE	None	None	None	None	None	None	None	None	None