	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	P53861	ELOA1_YEAST	Saccharomyces cerevisiae	Elongin-A	24.475439	NaN	S235;S278	33229814	MKSLQTLCEISLMRNHSNIQSVSNV PYHLLKRILQKVKIPQLLKLEKSNVL LIFDDDELWLEFLRQDFPTNVHEQF VSKRDIICKYYFDFVKENDIELYHSN QDLLKSCVRQSVVKDIRNNKYRIPYR MLYSKYQQEVEKKQEESAERLRLE MQKLQQEREKKQTIVVDHTVYFKKR NTKKTTRLHNEPHSQLYMKSLKDH ESRLKHFKDGGFNIAKRHAQRVAFG GQAGGQSSSPKKGPLSIKPEPVKVN RQMDNVTAEKKDVTQPITPVKKRRS ESPSIFLNRKKPALFRPTPKTNAAGS RPHTTAITNDHRTTSHPYPHKDVVT SISSVTANPVTKGHKKKKSGIFVRNA GSDGDSFPHVTATGPTTRPYIYEPRK	None	None	None	None	None	None	None	None	None