	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	P36037	DOA1_YEAST	Saccharomyces cerevisiae	Protein DOA1	24.475439	NaN	S332	33229814	MGYQLSATLKGHDQDVRDVVAVDD SKVASVSRDGTVRLWSKDDQWLGT VVYTGQGFLNSVCYDSEKELLLFGG KDTMINGVPLFATSGEDPLYTLIGHQ GNVCSLSFQDGVVISGSWDKTAKV WKEGSLVYNLQAHNASVWDAKVVS FSENKFLTASADKTIKLWQNDKVIKT FSGIHNDVVRHLAVVDDGHFISCSN DGLIKLVDMHTGDVLRTYEGHESFV YCIKLLPNGDIVSCGEDRTVRIWSKE NGSLKQVITLPAISIWSVDCMSNGDI IVGSSDNLVRIFSQEKSRWASEDEIN ELSTQVEKSTISSKTIEFDESKLSPYE ILQSPGRKEGQIVVVKSPQGTIEAHQ FSNSSWKKVGDVVGAGATGNDKKIE FEGKTYDYVFDVDIEDGKPPLKLPIN VSDNPYTAADNFLARYELPMSYRDQ VVQFILKNTNGISLDQPNDNASSSA VSPSKTSVMKVLPVKQYLIMENYNP DTIFNGIVKINSNEKTFDDEILAQIGG ALHDIDESWELLLSFANTIRSNWEIK TPAYDIVRLIVKKLPYSSDIKDYIEEG LGNKNITLTMLTVRILVNCFNNENW GVKLLESNQVYKSIFETIDTEFSQAS AKQSQNLAIAVSTLIFNYSALVTKGN SDLELLPIVADAINTKYGPLEEYQEC EEAAYRLTVAYGNLATVEPTLRQFA NSVTWLANIKRSYGNVPRFKDIFDD LS	None	None	None	None	None	None	None	None	None