	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	Q12329	HSP42_YEAST	Saccharomyces cerevisiae	Heat shock protein 42	24.475439	NaN	S182;S213;S214;S215;S223	33229814	MSFYQPSLSLYDVLNALSNQTGQRG QQGYPRQPQRPQRYHPHYGQVHVG GHHPRHHPLYSRYNGVPNTYYYQFP GQAYYYSPEYGYDDEDGEEEDQDED MVGDSGTTRQEDGGEDSNSRRYPS YYHCNTARNNRTNQQANSLNDLLT ALIGVPPYEGTEPEIEANTEQEGEKG EEKDKKDKSEAPKEEAGETNKEKPL NQLEESSRPPLAKKSSSFAHLQAPSP IPDPLQVSKPETRMDLPFSPEVNVYD TEDTYVVVLALPGANSRAFHIDYHPS SHEMLIKGKIEDRVGIDEKFLKITEL KYGAFERTVKFPVLPRIKDEEIKATY NNGLLQIKVPKIVNDTEKPKPKKRIA IEEIPDEELEFEENPNPTVEN	None	None	None	None	None	None	None	None	None