	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	P32356	TREA_YEAST	Saccharomyces cerevisiae	Cytosolic neutral trehalase	24.475439	NaN	S20;S21;S23;T58;S60;S66;S83	33229814	MSQVNTSQGPVAQGRQRRLSSLSEF NDPFSNAEVYYGPPTDPRKQKQAKP AKINRTRTMSVFDNVSPFKKTGFGK LQQTRRGSEDDTYSSSQGNRRFFIE DVDKTLNELLAAEDTDKNYQITIEDT GPKVLKVGTANSYGYKHINIRGTYM LSNLLQELTIAKSFGRHQIFLDEARI NENPVNRLSRLINTQFWNSLTRRVD LNNVGEIAKDTKIDTPGAKNPRIYVP YDCPEQYEFYVQASQMHPSLKLEVE YLPKKITAEYVKSVNDTPGLLALAME EHFNPSTGEKTLIGYPYAVPGGRFN ELYGWDSYMMALGLLEANKTDVAR GMVEHFIFEINHYGKILNANRSYYLC RSQPPFLTEMALVVFKKLGGRSNPD AVDLLKRAFQASIKEYKTVWTASPRL DPETGLSRYHPNGLGIPPETESDHF DTVLLPYASKHGVTLDEFKQLYNDG KIKEPKLDEFFLHDRGVRESGHDTT YRFEGVCAYLATIDLNSLLYKYEIDIA DFIKEFCDDKYEDPLDHSITTSAMW KEMAKIRQEKITKYMWDDESGFFFD YNTKIKHRTSYESATTFWALWAGLA TKEQAQKMVEKALPKLEMLGGLAA CTERSRGPISISRPIRQWDYPFGWAP HQILAWEGLRSYGYLTVTNRLAYRW LFMMTKAFVDYNGIVVEKYDVTRGT DPHRVEAEYGNQGADFKGAATEGF GWVNASYILGLKYMNSHARRALGA CIPPISFFSSLRPQERNLYGL	None	None	None	None	None	None	None	None	None