	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	Q08446	SGT1_YEAST	Saccharomyces cerevisiae	Protein SGT1	24.475439	NaN	S168;S171	33229814	MPVEKDLKTAYKALYDEKEPLKALH LYDEILKGSPTNLTALIFKAACLEKLY FGFSDWHSDATMENAKELLDKALM TAEGRGDRSKIGLVNFRYFVHFFNIK DYELAQSYFKKAKNLGYVDDTLPLW EDRLETKLNKKNKKQKDSTNKHTIK PVESIENRGDNNSSHSPISPLKIETA PQESPKFKIDWYQSSTSVTISLFTVN LPESKEQVNIYISPNDRRTLSISYQVP KSGSEFQYNAKLSHEVDPKAVSLKIF PKKLEITLSKIDSTQWKKLEEDILTES SRLSDEGKNSDSATRLLSAETASKE RLSYPSSSKKKIDWSKLDIDEEADEE AGSADSFFQKLYAGADPDTKRAMM KSFIESNGTALSTDWEDVSKGTVKT SPPEGMEPKHW	None	None	None	None	None	None	None	None	None