	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	P53330	RT102_YEAST	Saccharomyces cerevisiae	Regulator of Ty1 transposition protein 102	24.475439	NaN	S77;S122	33229814	MDPQTLITKANKVSYYGNPTSKESW RYDWYQPSKVSSNVQQPQQQLGDM ENNLEKYPFRYKTWLRNQEDEKNL QRESCEDILDLKEFDRRILKKSLMTS HTKGDTSKATGAPSANQGDEALSVD DIRGAVGNSEAIPGLSAGVNNDNTK ESKDVKMN	None	None	None	None	None	None	None	None	None