	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	P00931	TRP_YEAST	Saccharomyces cerevisiae	Tryptophan synthase	24.475439	NaN	S540;S683	33229814	MSEQLRQTFANAKKENRNALVTFM TAGYPTVKDTVPILKGFQDGGVDIIEL GMPFSDPIADGPTIQLSNTVALQNG VTLPQTLEMVSQARNEGVTVPIILM GYYNPILNYGEERFIQDAAKAGANG FIIVDLPPEEALKVRNYINDNGLSLIP LVAPSTTDERLELLSHIADSFVYVVS RMGTTGVQSSVASDLDELISRVRKY TKDTPLAVGFGVSTREHFQSVGSVA DGVVIGSKIVTLCGDAPEGKRYDVAK EYVQGILNGAKHKVLSKDEFFAFQK ESLKSANVKKEILDEFDENHKHPIRF GDFGGQYVPEALHACLRELEKGFDE AVADPTFWEDFKSLYSYIGRPSSLHK AERLTEHCQGAQIWLKREDLNHTGS HKINNALAQVLLAKRLGKKNVIAET GAGQHGVATATACAKFGLTCTVFM GAEDVRRQALNVFRMRILGAKVIAV TNGTKTLRDATSEAFRFWVTNLKTT YYVVGSAIGPHPYPTLVRTFQSVIGK ETKEQFAAMNNGKLPDAVVACVGG GSNSTGMFSPFEHDTSVKLLGVEAG GDGVDTKFHSATLTAGRPGVFHGVK TYVLQDSDGQVHDTHSVSAGLDYPG VGPELAYWKSTGRAQFIAATDAQAL LGFKLLSQLEGIIPALESSHAVYGAC ELAKTMKPDQHLVINISGRGDKDVQ SVAEVLPKLGPKIGWDLRFEEDPSA	None	None	None	None	None	None	None	None	None