	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	P25332	RBSK_YEAST	Saccharomyces cerevisiae	Ribokinase	24.475439	NaN	NaN	33229814	MGITVIGSLNYDLDTFTDRLPNAGET FRANHFETHAGGKGLNQAAAIGKLK NPSSRYSVRMIGNVGNDTFGKQLKD TLSDCGVDITHVGTYEGINTGTATILI EEKAGGQNRILIVEGANSKTIYDPKQ LCEIFPEGKEEEEYVVFQHEIPDPLSI IKWIHANRPNFQIVYNPSPFKAMPK KDWELVDLLVVNEIEGLQIVESVFD NELVEEIREKIKDDFLGEYRKICELLY EKLMNRKKRGIVVMTLGSRGVLFCS HESPEVQFLPAIQNVSVVDTTGAGD TFLGGLVTQLYQGETLSTAIKFSTLA SSLTIQRKGAAESMPLYKDVQKDA	None	None	None	None	None	None	None	None	None