	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	Q06505	IWS1_YEAST	Saccharomyces cerevisiae	Transcription factor SPN1	24.475439	NaN	T15;S23;S40;S85;T86;S89	33229814	MSTADQEQPKVVEATPEDGTASSQK STINAENENTKQNQSMEPQETSKGT SNDTKDPDNGEKNEEAAIDENSNV EAAERKRKHISTDFSDDDLEKEEHN DQSLQPTVENRASKDRDSSATPSSR QELEEKLDRILKKPKVRRTRRDEDD LEQYLDEKILRLKDEMNIAAQLDIDT LNKRIETGDTSLIAMQKVKLLPKVVS VLSKANLADTILDNNLLQSVRIWLEP LPDGSLPSFEIQKSLFAALNDLPVKT EHLKESGLGRVVIFYTKSKRVEAQLA RLAEKLIAEWTRPIIGASDNYRDKRI MQLEFDSEKLRKKSVMDSAKNRKK KSKSGEDPTSRGSSVQTLYEQAAAR RNRAAAPAQTTTDYKYAPVSNLSAV PTNARAVGVGSTLNNSEMYKRLTSR LNKKHK	None	None	None	None	None	None	None	None	None