

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P50875	SPT20_YEAST	Saccharomyces cerevisiae	Transcription factor SPT20	24.475439	NaN	S446;T516	33229814	MSANSPTGNDPHVFGIPVNATPSN MGSPGSPVNVPPMNPAPANVNH VMRTNSNSNANEGTRTLTREIQQL QQRQRLLQQRLLLEQQRKQALQN YEAQFYQMLMTLNKRPKRLYNFVE DADSILKKYEQYLHSFEFHIYENNYK ICAPANSRLQQQKQPELTSGLILT KNNETLKEFLEYVARGRIPDAIMEVL RDCNIQFYEGNLILQVYDHTNTVDV TPKENKPNLNSSSSPSNNSTQDN SKIQQPSEPNSGVANTGANTANKKA SFKRPRVYRTLLKPNDLTTYDMMS YADNARFSDSIYQFSEILTLTKRN LSLSVPLNPYEHRDMLEETAFASEPH WDSEKKSFIHEHRAESTREGTKGVV GHIEERDEFPOHSSNYEQLMLIMNE RTTTITNSTFAVSLTKNAMEIASSSS NGVRGASSTSNASNTRNNSLAN GNQVALAAAAAAAVGSTMGNDNN QFSRLKFIEQWRINKEKRKQALSA NINPTPFNARISMTAPLTPQQLLQR QQQALEQQQNGGAMKNANKRSGN NATSNNNNNNNLDKPKVKRPRK NAKSESGTPAPKKRMTKKKQSAS STPSSTTMS	None	None	None	None	None	None	None	None	

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