

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
Q12427	STB3_YEAST	Saccharomyces cerevisiae	Protein STB3	24.475439	NaN	S254	33229814	MSENQKEVSPPOAISVKSEASSSIFS KPISTSSPAGLAAAQRVTPGKLSTLLL EMGPLAIRHITQTLCLDIPCFKDLSS SKQRRLIMSAMESGDKEKSVVFEKI GWGQWSAKRVDPANFDKELEATNF ANAKVKDLISQESQRRKSNNSNSNS GGKVEMPMKVEHNITNIDGATTPPT AVASTTIPVNIKRSKSPLAANVVYID ENALASEDEDEEFDEDDHLLHYQN KSRNSSNNGKSSNGDPYSFGRRRS QVVFADSTPENIEHEIIAQKIRPLLRN RRRSSIKPHTPFISKLNTHQDSSYLS PNTTSTTTSPNNNSNSNQAKIDLEK LTATSEPTSRASRLSVSKESSIRSTL FPNKNYLIVTTNPNSKATSVSTSPKL EEQMNVSNNPIVLSKKEKHRVASQH LNGESSPQLVPHSHHQPHSDTDEE DWESIGAASLRNNSLAPNIDSVASST NGVVSPKPTNPSFTNSQNGDIEPPL QHDQQKHEQPRNGEDNSAAFL MSLKS	None	None	None	None	None	None	None	None	

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