

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38041	BOB1_YEAST	Saccharomyces cerevisiae	Protein BOB1	9.365641	NaN	S104;S106;S128;T151;T158;S209;S393;S412;S525;S528;S589;S590;S593;S644;S655;S735;T919	33229814	MSLEGNTLGKGAJSFPLYIAVNOQYS KRMEDELNMKPGDKIKVITDDGEYN DGWYYGRNLRRTKEEGLYPAVFTKRI AIEKPENLHKSPTQESGNSGVKYGN LNDSASNIGKVSSHQENRYTSLKS TMSDIDKALEELRSGSVEQEVSKSPT RVPEVSTPQLQDEQTLIQEKTRNEE NTTHDSLFSSTADLNLSSSELKNISK SNISTKSLEPSSSVRQLDLKMAKS WSPEEVTDYFSLVGFQSTCNKFKE HQVSGKILLELELEHLKELEINSFGI RFQIFKEIRNIKS AIDSSSNKLDADYS TFAFENQAAQLMPAATVNRDEIQQQ ISSKCNKLSSSESDRKS SVTTELQR PSSVVVNP NFKLHDPAEQILDMTEV PNLFADKDIFESPGRAPKPPSPV QPPQSPSFNNRYTNNNARFPPQTTY PPKNKNPTVYSNGLIPNSSTSSDNST GKFKFPAMNGHDSNSRKTTLTSATI PSINTVNTDESLPAISNISSNATSHH PNRNSVVYNNHKRTESGSSFVDF NRISMLSPVKSSFDEEETKQPSKAS RAVFD SARRKSSYGHSRDASLSEMK KHRRNSSILSFFSSKSQSNPTSPTKQ TFTIDPAKMTSHRSQSNSYSHARS QSYSHSRKHSLVTSPLKTSLSPINSK SNIALAHSETPTSSNNKEAVSQPSE GKHKHKHKHKSKHKHKNSSSKDGS SEEKSKKKLFSSTKESFVGSKEFKRS PSELTQKSTKSILPRSNAKKQQTSAF TEGIRSITAKESMQTADCSGWMSKK GTGAMGTWKQRFRTLHGTRLSYFT NTNDEKERGLIDITAHRVLPASDDD RLISLYAASLGKGYCFKLVPPQPGS KKGLTFTEPRVHYFAVENKSEMKA WLSAIKATIDIDTSVPVISSYATPTIP LSKAQTLLEEARLQTLRDAEEEEG RDQFGWDDTQNKRN SNYPIDQDQF ETSDYLESSAFEYPPGGRL