

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q03735	NAB6_YEAST	Saccharomyces cerevisiae	RNA-binding protein NAB6	9.365641	NaN	S464;S467	33229814	MSNSNSKPKPVANYAYRQQDYNGM NAMVGNPMMYHPVDFVNGAGQYG PSQHPAYYTN SPLNIPPTPFDTAYG ASLFPSHLLMGSPFVSSPNMQSGYN SARSSNLKRKAYSRPVSNHNGYNG NSNSNQNTNNGMVTSPSNYYRMG RNSFSRNNNSTRNVTHNNKGCCT RNNSGRRTFARNNIFDDILPEMLLQ RPFYCINVKVLP TGDDAYRTRSLLEN VDHSIDLHSIVKNFVKSNTLESAYLI EGGKSDDSKDVETKNLSILISFLT DCLNFYNNILQRLSEFKTFLKSEAL NLKFVCLNYDPKCLPTFIESEALTE AEEADITNGSTMISASLHHNIANKD ATRSIIIEFKSPVEKSDLFKKKLQFLD RSKNKRYILESIDL VNTDVP SNQFPE NYAVLTFNLISMAIEVLDYLKKYSKN LGISKCFYVSLAPLVSSARSSVANIY EGKTSTHRLSVPSVTAGNNNDSNN NGNNNKS NMSGITLNNNSSIGVS VYGHSNMSLTSLSSSVSLNEEIDML ATKLQGVELDGTYLEIN YRDYQTPTI EEHSTHLSNVKISKTTENS RQFSQDI PSPLPLNEHMF MNDSNQSN GAIIPQ QLIATPSPVSPNLQMNQRVLPNPITQ SLEQNFNVSAKVASSMGSDIGNRTI YIGNINPRSKAEDICNVVRGGILQSIK YIPEKKICFVTFIEAPSAVQFYANSFI DPIVLHGNMLRVGWGHYSGPLPKLI SLAVTIGASRN VYVSLPEFAFKEKFI HDPQYKKLHETLSLPDAEQLREDF TYGDIEQINYLSDSHCCWINFMNISS AISLVEEMNKESTVQNESGEVTLKR ATEEKFGGRYKGLLINYGKDRCGNI NKNLIAGKNSRFYKKVKRPSYNIRLS KLEEKRRQNEIDEKEKAFDKPLNLE SLGISLDAHKDNGGGETGTANNTG HENESELEAENENGNETGSFGGLG LAVASSDVKRATSD ETDYEDIFNKSS GSSDSSSDVEVIMHSPSDPEYALKS QTLRSSSQTVINSKRPKIEDEEEAV GMSQLNYRSSLRQAPP RAPSTLSYN HSKNNETPMQDIFTNGETANNRKK KRGSFARHRTIPGSDVMAQYLAQVQ HSTFMYAANILGASAEDNTHPDE