

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
P33314	BUD2_YEAST	Saccharomyces cerevisiae	Inhibitory regulator protein BUD2/CLA2	9.365641	NaN	S854	33229814	MSSNNEPAQSRTSYFKLNEFLSNVK HYKNTFKGEIQWCNNLSLNDWKTH YLQITSTGALTHSIDELTADSTNIQPII KHLQQCRIEIHKDKHSSFKDINANCN FIIQVNTSGKDNKVYLRVKSWSDFK KLLTCLIWSSMKTNGIFNKFQVSR PLEFKSKMAKPESLLVYKLNFGPI VKNIVLPPATNILESPDIINDDNSV GWFSAMGVLKSNGLDLLLLQSDGS LIYSLNISQLLRSEIRILDSSVLQSEN SLFLGELPLLRSQLGLEKFRIENIASA ATNSSDISQEIIVEFPLRIDLEDCFIAL QSFARSEYLSITGSDKSNDMKISNSF KISILEANFQSINLNDKNNTWPWSIFT DITAWGHTWARTSMVSNSSNPFWR EEFQFNELLRLTNSYLEIKQLFHDL NNKKRLRLIGKIKITQEIINDTRYNKE TRLPIMDVDNKNFQIGTICKISSNL NFILPSTNFKLEKLLMNANLSMVS NLIYKSSSSMENDNKLTQTSIIFLDIF QLSRIEWFHVLIDKELAKIDGTVS RINQKNLDSKHVFNLSFRGNSILTK SIEQYFFRVGNEYLSKALSAILKEIIE SNKSCELDPARVKEKDEVKKRKIIAD NYKRLYSWVTKIWKRLYATSNLPIE IRNVLKIFRQKLEIICIDDTLQIILNGI SGLLFLRFFCPVILNPKLFKYVSQNL NETARRNLTLSKVLLNLSTLTQFAN KEPWLMKMNNFIDKRHNDLLDYID KMTQKKLDFNSKILNLSSTISRPKLA IEQTMLDDLQIPYLLDKNLRETEFV NLIVNFSQEDMTKMEKYNHMDNG GKGELIEEGLLSGSSLNLSVDKKDL DSPIEVKPEIGELEFEKITENNTEIFG DDLMLLKSDDVGSRSRDLNNGAN SGIKFNSIIPKAEEKHAMKELEQES CLLYNRINHIRKRLSGYECASSTLFE DKKYSISLSHKIFYEEIKEGKEIVLKL LNKPTNENSSARLQKFFTKGVSSKS NNTVGD SYCKFLTIDVSDENPKSSN KTSVHGTSSENGAKDDYLTLPNSQG KGNLGNRFSPTKLSRIMRKPKNADV PKEQNSRKLTRWFKKKKETGGS