

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38692	KIC1_YEAST	Saccharomyces cerevisiae	Serine/threonine-protein kinase KIC1	9.365641	NaN	S735	33229814	MTTKPQNSKQGLAEGEMDVSSLFK RTEVIGRGKFGVVYKGYNVKTGRVY AIKVLNLDSDSDEVEDVQREIQFLAS LKQISNITRYGSSYLKDTSLWIIMEH CAGGSLRSLLRPGKIDEKYIGVIMRE LLVALKCIHKDNVIHRDIKAANVLIT NEGNVKLCDFGVAAQVNQTSRRQ TMAGTPYWMAPEVIMEGVVYDTKV DIWSLGITTYEATGNPPYCDVEALR AMQLIISKPPRLEDRSYSTSLKEFIA LCLDEDPKERLSADDLLKSKFIRAHK ATPTSILKELISRYLLFRDKNKNKYKI EGSIPENEPSKPSSEAPKPSQNGGGD EAQKSIASNDNEIKRVNEGDMEMK WDFDSLSSDYIENINLDALAEDN NEWATAQHDLFNYAYPDEDSYYFDP TSHNTRPFVYQGTTIGKGYPGTIAQN STLNAPVTNNYTNISKYPSKMVAGTT NTSGTHTAGPMTSSKRLESKAPKQL LELFEDNEIITAENDVNTAPKISKSI SSLNAGNSSRDDFIPSISNEVNGNIN NNKMRPHLPLSSGNNYYSQSTPAL PLLQTKFNKTSKGPPTSGLTTAPTISI EIEIPEELPNSALPTPASADPVLIPST KARSSTVTAGTPSSSSSIQYKSPSNV PRRLTVSNNRPEHCPSTITNQKLG AVASNSGISSTPNNSNNYNNNTDSE NSRGSSGSNTANSTQMGITNPGNV TKLSTHKASSPSRPLFGVGTSPNRKP AGSPTQNIGHNSTHTNLAPPPTMKP MANSKDNKDILLQPLNSIPSSSTLNT ISGNSSNNLTSSNYFSNEKEGSRVN GDFKRNNPNLKLQMLPTPVVRNK LLDPNTATSQNNNGMPGSAGISTNE NINQFGFNTSSASNIPVSMTPISEKH IDFGGKIKRSQISNRKNSSASEHPL NILGSSVSGNVSGIGNNVGSNNNS GPNNVPLSANTGNTTIKANSTTIAT SSSAAASTTAPISQQTIPSGTQFNHIL SSAATAANSVNSLGFGMCPPPQSLQ MEMFLDLESFLPGKQRRVDRKPQV LKELENLLQMFEEGLPCIEHALKEQ LISTPIKDNEH