

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P46682	AP3B_YEAST	Saccharomyces cerevisiae	AP-3 complex subunit beta	9.365641	NaN	S693;S698;S724;S726	33229814	MVDSIHRIASALDTAKVITREAAAVA TSKLGESSYTYYSQNINPQQLVTLN SRNSREVRDAMKRIISIMASDDDSID VQLYFADVVKNITTNDTKVKRLIHL LLRFAENDPNLTLLSINSLQKSLSDS NSELRCFALSALSDMKMSSLAPIILH TVKKLVTDPSAMVRGEVALAIKLYR AGKNDYHEELLDKELMADTDPKV ISCAVLAYKECYADHLELLHGHFRR YCRIIKQLDSWSQSYLELLIKYCKQY LPKPTVVDKSSEGS PRSCPLPKYNE IEYPSYEVVNDPDLDFLQSLNCLY SSNPTVILSCCNALYQLASPLQMKN TKFIEALVRTVTMTENQGNKEMLLQ AIHFLSILDQTLFLPYTKKFYVFPKDP IVASIWKIQLSTLINESNVKEIFKELK YYVASAHFPENVVIMAVKSLSRCGQ LSTSWESHVMKWLIDHMESHNL SVLDAYVNVIRMLVQKNPTKHLRIIF KLADLLTVQTS LADNARAGIVWLF EIASIEFKICPDVLRRLIQNFSNEGPE TRCQILVLSAKLLSYDIDNFKQAQVT GSEENNQNPPYYDFSGSRISQMYNA VLYLAKYDDEFDIRDRARMISLFD GKYEIVSLLLQAPKPTARSDDFIVSA RLEHTPEIKEFFRMLPWNTTEITEVG ETGNDIREGAELKDYNKYKSFSSQ SFITNNSARSFTSSSNKLTGINDGD SNSISGKGNVNTFTSQNGKKYRLQS LDEFFSDIPERKSKPRKIIKVVES EDEDESEESSDDDEYS DSSLGTSS GTSSSHLEL

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