

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
P15801	DPOG_YEAST	Saccharomyces cerevisiae	DNA polymerase gamma	9.365641	NaN	NaN	33229814	MTKLMVRSECMLRMVRRRPLRVQF CARWFSTKKNTAEAPRINPVGIQYL ESLQRQVFGSCGGKDEVEQSDKLM ELSKKSLKDHGLWGKKTLLITDPISFP LPPLQGRSLDEHFQKIGRFNSEPYKS FCEDKFTMVARPAEWLRKPGWVK YVPGMAPVEVAYPDEELVVDVETL YNVSDYPTLATALSSTAWYLWCSPFI CGGDDPAALPLNNTLNKEQVIIGHNV AYDRARVLEEYNFRDSKAFFLDTQS LHIASFGLCSRQPMFMKNNKKKE AEVESEVHPEISIEDYDDPWLNVSAL NSLKDVAKFHCKIDLKTDTRDFFAS TDKSTIENFQKLVNYCATDVTATSQ VFDEIFPVFLKKCPHPVSFAGLKSL KCILPTKLNWDNDYLNSSSESLYQOS KVQIESKIVQIIKDIVLLKDKPDFYLK DPWLSQLDWTTKPLRLTKKGVPAKC QKLPGFPEWYRQLFPSKDTVEPKITI KSRIIPILFKLSWENSPVIWSKESGW CFNVPHEQVETYKAKNYVLADSVSQ EEEEIRTHNLGLQCTGVLFKVPHPN GPTFNCTNLLTKSYNHFFEKGVLS ESELHQALQINSSGSYWMSARERI QSQFVVPSCKFPNEFQSLSAKSSLN NEKTNDLAIIPKIVPMGTITRAVEN AWLTASNAKANRIGSELKTQVKAPP GYCFVGADVSEELWIASLVGDSIF NVHGGTAIGWMCLEGTKNEGTDLH TKTAQILGCSRNEAKIFNYGRIYGAG AKFASQLLKRFNPSLTDEETKKIANK LYENTKGKTKRSKLFKKFWYGGSES ILFNKLESIAEQETPKTPVLGCGITYS LMKKNLRANSFLPSRINWAIQSSGV DYLHLLCCSMEYIHKYNLEARLCISI HDEIRFLVSEKDKYRAAMALQISNI WTRAMFCQQMGINELPQNCAFFSQ VDIDSVIRKEVNMDICITPSNKTAIPH GEALDINQLLDKSNKLGKPNLDID SKVSQYAYNYREPVFEEYNKSYTPEF LKYFLAMQVQSDKRVDNRLEDEYLR ECTSKEYARDGNTAEYSLLDYIKDVE K GKRTKVRIMGSNFLDGTKNAKAD QRIRLPVNMPDYPTLHKIANDSAIPE KQLLENRRKKENRIDDENKKKLTRK KNTTPMERKYKRVYGGKAFEAFYE CANKPLDYTLETEKQFFNIPIDGVID DVLNDKSNYKKKPSQARTASSSPIRK TAKAVHSHKLPARKSSTTNRNLVEL

