

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P33609	DPOLA_MOUSE	Mus musculus	DNA polymerase alpha catalytic subunit	30.962534	NaN	T180;S192;S215	39627609;28528544	MAPMHEEDCKLEASAVSDSGSFAA SRARREKSKKGRQEALERLKKAKA GEKYKYEVEDLTSVYEEVDEEQYSKL VQARQDDDDWVDDDGIGYVEDGREI FDDDLDDALDTCGKGSDBGKAHRK DRKDVKKPSVTKPNNIKAMFIASAG KKTDDKAVDLSKDDLLGDILQDLNT ETAQITPPPVLIPKKRSTGALLNPFS VHTPKAIPSGKPASPVLNRNEPLLTPIP LKRAELAGELAQPECEDEQELGVM EFEDGDFDESMDTEKVDKPVTAKT WDQETEPVERVEHEADPERGTTSYL ENFLPDVSCWIDIDQDDESIPQEVQV DSSNLPVKGADDEQVFQFYWLDAY EDPYNQPGVVFLFGKVWIESVKTTHV SCCVMVKNIERTLYFLPREMKFDLN TGKETAIPVTMKDVYEEFDSKISAKY KIMFKFSKIVEKNYAFEIPDVPEKSE YLEVRYSAEVPQLPQNLKGETFSHV FGTNTSSLEFLMNRKIKGPCWLEV KNPQLLNQPISWCKFEVMALKPDLV NVIKDVSPPPLVVMFSFMSKMTQNV QNHQHEIIAAMALVHHSFALDKAPP EPPFQTHFCVVSFKPKCIFPCDFKEV ISKKNMKVEIAATERTLIGFFLAKVH KIDPDILVGHNICSFELEVLQRINE CKVPYWSKIGRLRRSNMPKLGSRSG FGERNATCGRMICDVEISAKELIHCK SYHLSLVQOILKTERIVIPTENIRN MYSESSYLLYLEHIWKDARFILQIM CELNVLPLALQITNIAGNIMSRTLGMG GRSERNEFLLLHAFYENNYIVPDKQI FRKPQQKLGDEDEEIDGDTNKYKKG RKKATYAGGLVLDPKVGFYDKFILLL DFNSLYPSIIQEFNICFTTVQRTSEV QKATEDEEQEIQPELDPNLEMGILP REIRKLVERRKQVKQLMKQDLNPD LVLQYDIRQKALKLTANSMYGCLGF SYSRFYAKPLAALVYKGREILMHTK DMVQKMNLEVIYGDTSIMINTNST NLEEVFKLGNKVKSEVNKLYKLEI DIDAVFKSLLLLKKKYAALVVEPTS DGNYYTKQELKGLDIVRRDWCDLAK DTGNFVIGQILSDQSRDTIVENIQKR LIEIGENVLNGSVVPSQFEINKALTK DPQDYPDRKSLPHVHVWALWINSQGG RKVKAGDTVSYVICQDGSNLTATQR AYAPEQLQKLDNLAIDTQYYLAQQIH PVVARICEPIDGIDAVLIALWGLDST QFRVHQYHKDEENDALLGGPAQLT DEEKYKDCFKCLCPSCGTENIYD NVFEGSGLDMEPSLYRCSNVDCKVS PLTFMVQLSNKLIMDIRRCIKKYDYG WLICEEPTCCSRLRRLPLHFSRNGP LCPVCMKAVLRPEYSDKSLYTLQCF YRYIFDADCALEKLTEHEKDKLKKQF FPLRVLQDYRKVKNIAEQFLSWSGY SEVNLSKLFANYAGKS	True	False	4.486	5.0	1.519	0.868	False	1.021	1.203