	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P09884	DPOLA_HUMAN	Homo sapiens	DNA polymerase alpha catalytic subunit	18.379595	NaN	T174;S186;S190;S209;T406	30379171	MAPVHGDDSLSDSGSFVSSRARREK KSKKGRQEALERLKKAKAGEKYKYE VEDFTGVYEEVDEEQYSKLVQARQD DDWIVDDDGIGYVEDGREIFDDDLE DDALDADEKGKDGKARNKDKRNVK KLAVTKPNNIKSMFIACAGKKTADK AVDLSKDGLLGDILQDLNTETPQITP PPVMILKKKRSIGASPNPFSVHTATA VPSGKIASPVSRKEPPLTPVPLKRAE FAGDDVQVESTEEEQESGAMEFED GDFDEPMEVEEVDLEPMAAKAWDK ESEPAEEVKQEADSGKGTVSYLGSF LPDVSCWDIDQEGDSSFSVQEVQVD SSHLPLVKGADEEQVFHFYWLDAYE DQYNQPGVVFLFGKVWIESAETHVS CCVMVKNIERTLYFLPREMKIDLNT GKETGTPISMKDVYEEFDEKIATKYK IMKFKSKPVEKNYAFEIPDVPEKSEY LEVKYSAEMPQLPQDLKGETFSHVF GTNTSSLELFLMNRKIKGPCWLEVK SPQLLNQPVSWCKVEAMALKPDLV NVIKDVSPPPLVVMAFSMKTMQNA KNHQNEIIAMAALVHHSFALDKAAP KPPFQSHFCVVSKPKDCIFPYAFKEV IEKKNVKVEVAATERTLLGFFLAKVH KIDPDIIVGHNIYGFELEVLLQRINVC KAPHWSKIGRLKRSNMPKLGGRSG FGERNATCGRMICDVEISAKELIRCK SYHLSELVQQILKTERVVIPMENIQN MYSESSQLLYLLEHTWKDAKFILQI MCELNVLPLALQITNIAGNIMSRTL MGGRSERNEFLLLHAFYENNYIVPD KQIFRKPQQKLGDEDEEIDGDTNKY KKGRKKAAYAGGLVLDPKVGFYDKFI LLLDFNSLYPSIIQEFNICFTTVQRVA SEAQKVTEDGEQEQIPELPDPSLEM GILPREIRKLVERRKQVKQLMKQQD LNPDLILQYDIRQKALKLTANSMYGC LGFSYSRFYAKPLAALVTYKGREILM HTKEMVQKMNLEVIYGDTDSIMINT NSTNLEEVFKLGNKVKSEVNKLYKL LEIDIDGVFKSLLLLKKKKYAALVVEP TSDGNYVTKQELKGLDIVRRDWCDL AKDTGNFVIGQILSDQSRDTIVENIQ KRLIEIGENVLNGSVPVSQFEINKAL TKDPQDYPDKKSLPHVHVALWINSQ GGRKVKAGDTVSYVICQDGSNLTAS QRAYAPEQLQKQDNLTIDTQYYLAQ QIHPVVARICEPIDGIDAVLIATWLGL DPTQFRVHHYHKDEENDALLGGPA QLTDEEKYRDCERFKCPCPTCGTEN IYDNVFDGSGTDMEPSLYRCSNIDC KASPLTFTVQLSNKLIMDIRRFIKKYY DGWLICEEPTCRNRTRHLPLQFSRT GPLCPACMKATLQPEYSDKSLYTQL CFYRYIFDAECALEKLTTDHEKDKLK KQFFTPKVLQDYRKLKNTAEQFLSR SGYSEVNLSKLFAGCAVKS	None	None	None	None	None	None	None	None	None