	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	Q04049	POLH_YEAST	Saccharomyces cerevisiae	DNA polymerase eta	24.475439	NaN	NaN	33229814	MSKFTWKELIQLGSPSKAYESSLACI AHIDMNAFFAQVEQMRCGLSKEDP VVCVQWNSIIAVSYAARKYGISRMDT IQEALKKCSNLIPIHTAVFKKGEDFW QYHDGCGSWVQDPAKQISVEDHKV SLEPYRRESRKALKIFKSACDLVERA SIDEVFLDLGRICFNMLMFDNEYEL TGDLKLKDALSNIREAFIGGNYDINS HLPLIPEKIKSLKFEGDVFNPEGRDL ITDWDDVILALGSQVCKGIRDSIKDIL GYTTSCGLSSTKNVCKLASNYKKPD AQTIVKNDCLLDFLDCGKFEITSFWT LGGVLGKELIDVLDLPHENSIKHIRE TWPDNAGQLKEFLDAKVKQSDYDR STSNIDPLKTADLAEKLFKLSRGRYG LPLSSRPVVKSMMSNKNLRGKSCN SIVDCISWLEVFCAELTSRIQDLEQE YNKIVIPRTVSISLKTKSYEVYRKSGP VAYKGINFQSHELLKVGIKFVTDLDI KGKNKSYYPLTKLSMTITNFDIIDLQ KTVVDMFGNQVHTFKSSAGKEDEE KTTSSKADEKTPKLECCKYQVTFTD QKALQEHADYHLALKLSEGLNGAEE SSKNLSFGEKRLLFSRKRPNSQHTA TPQKKQVTSSKNILSFFTRKK	None	None	None	None	None	None	None	None	None