	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	extracellul region
0	P04821	CDC25_YEAST	Saccharomyces cerevisiae	Cell division control protein 25	24.475439	NaN	S151;S154;S423;S580;S596;S632;T635;S649	33229814	MSDTNTSIPNTSSAREAGNASQTPSI SSSSNTSTTTNTESSSASLSSSPSTS ELTSIRPIGIVVAAYDFNYPIKKDSSS QLLSVQQGETIYILNKNSSGWWDGL VIDDSNGKVNRGWFPQNFGRPLRD SHLRKHSHPMKKYSSSKSSRRSSLN SLGNSAYLHVPRNPSKSRRGSSTLS ASLSNAHNAETSSGHNNTVSMNNS PFSAPNDASHITPQSSNFNSNASLS QDMTKSADGSSEMNTNAIMNNNE TNLQTSGEKAGPPLVAEETIKILPLEE IEMIINGIRSNIASTWSPIPLITKTSDY KLVYYNKDLDIYCSELPLISNSIMES DDICDSEPKFPPNDHLVNLYTRDLR KNANIEDSSTRSKQSESEQNRSSLL MEKQDSKETDGNNNSINDDDNNN ENNKNEFNEAGPSSLNSLSAPDLTQ NIQSRVVAPSRSSILAKSDIFYHYSRD IKLWTELQDLTVYYTKTAHKMFLKE NRLNFTKYFDLISDSIVFTQLGCRLM QHEIKAKSCSKEIKKIFKGLISSLSRI SINSHLYFDSAFHRKKMDTMNDKD NDNQENNCSRTEGDDGKIEVDSVH DLVSVPLSGKRNVSTSTTDTLTPMR SSFSTVNENDMENFSVLGPRNSVN SVVTPRTSIQNSTLEDFSPSNKNFKS AKSIYEMVDVEFSKFLRHVQLLYFVL QSSVFSDDNTLPQLLPRFFKGSFSG GSWTNPFSTFITDEFGNATKNKAVT SNEVTASSSKNSSISRIPPKMADAIA SASGYSANSETNSQIDLKASSAASGS VFTPFNRPSHNRTFSRARVSKRKKK YPLTVDTLNTMKKKSSQIFEKLNNA TGEHLKIISKPKSRIRNLEINSSTYEQ INQNVLLLEILENLDLSIFINLKNLIK TPSILLDLESEEFLVHAMSSVSSVLT EFFDIKQAFHDIVIRLIMTTQQTTLD DPYLFSSMRSNFPVGHHEPFKNISN TPLVKGPFHKKNEQLALSLFHVLVS QDVEFNNLEFLNNSDDFKDACEKY VEISNLACIIVDQLIEERENLLNYAAR MMKNNLTAELLKGEQEKWFDIYSE DYSDDDSENDEAIIDDELGSEDYIER KAANIEKNLPWFLTSDYETSLVYDSR GKIRGGTKEALIEHLTSHELVDAAFN VTMLITFRSILTTREFFYALIYRYNLY PPEGLSYDDYNIWIEKKSNPIKCRVV NIMRTFLTQYWTRNYYEPGIPLILNF AKMVVSEKIPGAEDLLQKINEKLINE NEKEPVDPKQQDSVSAVVQTTKRDN KSPIHMSSSSLPSSASSAFFRLKKLK LLDIDPYTYATQLTVLEHDLYLRITM FECLDRAWGTKYCNMGGSPNITKFI ANANTLTNFVSHTIVKQADVKTRSK LTQYFVTVAQHCKELNNFSSMTAIV SALYSSPIYRLKKTWDLVSTESKDLL KNLNNLMDSKRNFVKYRELLRSVT DVACVPFFGVYLSDLTFTFVGNPDFL HNSTNIINFSKRTKIANIVEEIISFKR FHYKLKRLDDIQTVIEASLENVPHIE KQYQLSLQVEPRSGNTKGSTHASSA SGTKTAKFLSEFTDDKNGNFLKLGK KKPPSRLFR	None	None	None	None	None	None	None	None	None