	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	extracellula region
0	Q69YH5	CDCA2_HUMAN	Homo sapiens	Cell division cycle-associated protein 2	21.731224	NaN	S98;S120;S126;S131;S210;S291;S309;T312;S400;S407;T412;S437;S591;S614;S710;S756;S936;S977;S1000	23301498	MDANSKDKPPETKESAMNNAGNAS FILGTGKIVTPQKHAELPPNPCTPDT FKSPLNFSTVTVEQLGITPESFVRNS AGKSSSYLKKCRRRSAVGARGSPET NHLIRFIARQQNIKNARKSPLAQDSP SQGSPALYRNVNTLRERISAFQSAF HSIKENEKMTGCLEFSEAGKESEMT DLTRKEGLSACQQSGFPAVLSSKRR RISYQRDSDENLTDAEGKVIGLQIFN IDTDRACAVETSVDLSEISSKLGSTQ SGFLVEESLPLSELTETSNALKVADC VVGKGSSDAVSPDTFTAEVSSDAVP DVRSPATPACRRDLPTPKTFVLRSVL KKPSVKMCLESLQEHCNNLYDDDG THPSLISNLPNCCKEKEAEDEENFE APAFLNMRKRKRVTFGEDLSPEVFD ESLPANTPLRKGGTPVCKKDFSGLS SLLLEQSPVPEPLPQPDFDDKGENL ENIEPLQVSFAVLSSPNKSSISETLSG TDTFSSSNNHEKISSPKVGRITRTSN RRNQLVSVVEESVCNLLNTEVQPCK EKKINRRKSQETKCTKRALPKKSQV LKSCRKKKGKGKKSVQKSLYGERDI ASKKPLLSPIPELPEVPEMTPSIPSIR RLGSGYFSSNGKLEEVKTPKNPVKR KDLLRHDPDLHMHQGYDKYDVSEF CSYIKSSSSLGNATSDEDPNTNIMNI NENKNIPKAKNKSESENEPKAGTDS PVSCASVTEERVASDSPKPALTLQQG QEFSAGGQNAENLCQFFKISPDLNI KCERKDDFLGAAEGKLQCNRLMPN SQKDCHCLGDVLIENTKESKSQSED LGRKPMESSSVVSCRDRKDRRRSM CYSDGRSLHLEKNGNHTPSSSVGSS VEISLENSELFKDLSDAIEQTFQRRN SETKVRRSTRLQKDLENEGLVWISL PLPSTSQKAKRRTICTFDSSGFESMS PIKETVSSRQKPQMAPPVSDPENSQ GPAAGSSDEPGKRRKSFCISTLANTK ATSQFKGYRRRSSLNGKGESSLTAL ERIEHNGERKQ	True	False	4.551	4.812	1.068	0.983	0.631	1.091	0.915