	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endo reticu
0	Q96BY6	DOC10_HUMAN	Homo sapiens	Dedicator of cytokinesis protein 10	25.414088	S1000;S1318;T2165;S2169	T196;S302;T368;S877;S1232;S1257;S1292;S1295;S1318;T1406;T1440	37217939;40307207;30379171;40914422;29351928	MAGERTRRFTRSLLRPGQAAELRHS AASAAAVAVSSRQQQRQEKPRLLEP LDYETVIEELEKTYRNDPLQDLLFFP SDDFSAATVSWDIRTLYSTVPEDAE HKAENLLVKEACKFYSSQWHVVNY KYEQYSGDIRQLPRAEYKPEKLPSHS FEIDHEDADKDEDTTSHSSSKGGGG AGGTGVFKSGWLYKGNFNSTVNNT VTVRSFKKRYFQLTQLPDNSYIMNF YKDEKISKEPKGCIFLDSCTGVVQNN RLRKYAFELKMNDLTYFVLAAETES DMDEWIHTLNRILQISPEGPLQGRR STELTDLGLDSLDNSVTCECTPEET DSSENNLHADFAKYLTETEDTVKTT RNMERLNLFSLDPDIDTLKLQKKDL LEPESVIKPFEEKAAKRIMIICKALNS NLQGCVTENENDPITNIEPFFVSVAL YDLRDSRKISADFHVDLNHAAVRQM LLGASVALENGNIDTITPRQSEEPHI KGLPEEWLKFPKQAVFSVSNPHSEI VLVAKIEKVLMGNIASGAEPYIKNPD SNKYAQKILKSNRQFCSKLGKYRMP FAWAVRSVFKDNQGNVDRDSRFSP LFRQESSKISTEDLVKLVSDYRRADR ISKMQTIPGSLDIAVDNVPLEHPNCV TSSFIPVKPFNMMAQTEPTVEVEEF VYDSTKYCRPYRVYKNQIYIYPKHLK YDSQKCFNKARNITVCIEFKNSDEE SAKPLKCIYGKPGGPLFTSAAYTAVL HHSQNPDFSDEVKIELPTQLHEKHH ILFSFYHVTCDINAKANAKKEALET SVGYAWLPLMKHDQIASQEYNIPIAT SLPPNYLSFQDSASGKHGGSDIKWV DGGKPLFKVSTFVVSTVNTQDPHVN AFFQECQKREKDMSQSPTSNFIRSC KNLLNVEKIHAIMSFLPIILNQLFKV LVQNEEDEITTTVTRVLTDIVAKCHE EQLDHSVQSYIKFVFKTRACKERTV HEELAKNVTGLLKSNDSTTVKHVLK HSWFFFAIILKSMAQHLIDTNKIQLP RPQRFPESYQNELDNLVMVLSDHVI WKYKDALEETRRANHSVARFLKRCF TFMDRGYVFKMVNNYISMFSSGDL KTLCQYKFDFLQEVCQHEHFIPLCL PIRSANIPDPLTPSESTQELHASDMP EYSVTNEFCRKHFLIGILLREVGFAL QEDQDVRHLALAVLKNLMAKHSFD DRYREPRKQAQIASLYMPLYGMLLD NMPRIYLKDLYPFTVNTSNQGSRDD LSTNGGFQSQTAIKHANSVDTSFSK DVLNSIAAFSSIAISTVNHADSRASLA SLDSNPSTNEKSSEKTDNCEKIPRPL SLIGSTLRFDKLDQAETRSLLMCFLH IMKTISYETLIAYWQRAPSPEVSDFF SILDVCLQNFRYLGKRNIIRKIAAAFK FVQSTQNNGTLKGSNPSCQTSGLLS QWMHSTSSHEGHKQHRSQTLPIIR GKNALSNPKLLQMLDNTMTSNSNE IDIVHHVDTEANIATEVCLTILDLLSL FTQTHQRQLQQCDCQNSLMKRVFD TYMLFFQVNQSATALKHVFASLRLF VCKFPSAFFQGPADLCGSFCYEVLK CCNHRSRSTQTEASALLYFFMRKNF EFNKQKSIVRSHLQLIKAVSQLIADA GIGGSRFQHSLAITNNFANGDKQMK NSNFPAEVKDLTKRIRTVLMATAQM KEHEKDPEMLVDLQYSLANSYASTP ELRRTWLESMAKIHARNGDLSEAA MCYIHIAALIAEYLKRKGYWKVEKIC TASLLSEDTHPCDSNSLLTTPSGGS MFSMGWPAFLSITPNIKEEGAMKED SGMQDTPYNENILVEQLYMCVEFL WKSERYELIADVNKPIIAVFEKQRDF KKLSDLYYDIHRSYLKVAEVVNSEKR LFGRYYRVAFYGQGFFEEEEGKEYIY KEPKLTGLSEISQRLLKLYADKFGAD NVKIIQDSNKVNPKDLDPKYAYIQVT YVTPFFEEKEIEDRKTDFEMHHNIN RFVFETPFTLSGKKHGGVAEQCKRR TILTTSHLFPYVKKRIQVISQSSTELN	True	False	4.497	5.0	1.924	1.012

									PIEVAIDEMSKKVSELNQLCTMEEV DMIRLQLKLQGSVSVKVNAGPMAYA RAFLEETNAKKYPDNQVKLLKEIFR QFADACGQALDVNERLIKEDQLEYQ EELRSHYKDMLSELSTVMNEQITGR DDLSKRGVDQTCTRVISKATPALPTV SISSSAEV						