

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96JM3	CHAP1_HUMAN	Homo sapiens	Chromosome alignment-maintaining phosphoprotein 1	14.014672	S190;S325;S344;T362;S364;S366;S367;S369	S87;S108;S173;S184;S204;S214;S217;S244;S247;S253;S264;S275;S282;S286;S297;S308;S319;S344;S355;S376;S382;S386;T403;S405;S416;S427;S432;S436;S443;S445;S452;T458;S459;S462;S472;S476;S507;S512;S542;S572;S603;S615;S626;S627;S632;S651;S652;S653;S675;S736	32574038;29351928;34019948;35289036;32119511;33214551	MEAFQELRKPSARLECDHCSFRGTD YENVQIHMGTIHPFCDEMDAGGL GKMIFYQKSAKLFHCHKCFFTSKMY SNVYYHITSKHASPKWNDKPKNQL NKETDPVKSPLPEHQKIPCNSAEP KSIPALSMETQKLGSVLSPESPKPTP LTPLEPQKPGSVVSPQLQTPSPSEP SKPASVSSPEPKSVVPCESQKLPV PSPEPQKPAPVSPESVKATLSNPKPQ KQSHFPETLGPSSASSPESVLAASP EPWGPSPAASPESRKSARTTSPEPRK PSPSESEPWKPFPAVSPERRPAPA VSPGSKPGPPGSPRPWKSNSASS GPWKPAKPAPSVSPGPWKPIPSVSP GPWKPTPSVSSASWKSSSVSPSSWK SPPASPESWKSPPPELRKTAPTLSPE HWKAVPPVSPELRKPGLPLSPEIRSP AGSPELRKPSGSPDLWKLSPDQRKT SPASLDFPESQKSSRGGSPDLWKSS FFIEPQKPVFPETRKPGSPGSPESPK AASDIWKPVLSIDTEPRKPALEPAPA KTAPPASPEARLALFPEPRKHALFP ELPKSALFSESQKAVELGDELQIDAI DDQKCDILVQEELLASPKKLEDTLF PSSKKLKKDNQESSDAELSSSEYIKT DLDAMDIKQESSSDQEQVDVESID FSKENKMDMTSPEQSRNVLQFTEE KEAFISSEEEIAKYMKRGKGYCKIC CCRAMKKGAVLHHLVNHVHSPY KCTICGKAFLLESLLKNHVAAHGQS LLKPCRCNFESNFRGFKKHLTHCQ SRHNEEANKKLEALEPPLLEEQQI

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