

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96RK0	CIC_HUMAN	Homo sapiens	Protein capicua homolog	35.106741	S13;S16;S530;S534;S535;S541;S542;S543;S566;T583;S649;S665;S734;T738;T752;T831;T880;S891;T967;S972;T973;S975;S982;T985;S991;S1001;T1002;T1143;T1149;S1154	S146;S173;S277;S362;S431;S436;S496;S700;S721;S739;T1291;S1294;S1351;S1373;S1378;S1382;S1389;S1397;T1398;S1402;S1409;S1595	23301498;32574038;29237092;31492838;30397120;35254053;28657654;20305658;33214551;27655845;30059200;35132862;34725712;21740066;32119511;35289036;35138101;34019948	MYSahrPLMPASSAASRGLGMFVW TNVEPRSVAVFPWHSLVPLFAPSQ DPSVQVSEAAQPPASHPVASNQSK AESAAVAHERPPGGTGSADPERPPG ATCPESPGPPHPLGVVESGKGGP PTTEEEASGPPGPRLDSETESDHD DAFLSIMSPEIQLPLPPGKRRTQSL ALPKERDSSSEKDRSPNKREKDH RRPMNAFMIFSKRHRALVHQRHPN QDNRTVSKILGEWWYALGPKEKQKY HDLAFQVKEAHFKAHPDWKWCNK DRKSSSEAKPTSLGLAGGHKRE RSMSETGTAAPGVSELLSVAQT LLSSDTKAPGSSSCGAERLHTVGGP GSARPRAFSHSGVHSLDGGEVDSQA LQELTQMVSGPASYSGPKPSTQYGA PGPFAAPGEGGALAATGRPPLPTRA SRSQRAASEDMTSDEERMVICEE DDDVIADDGFGTTDIDLKCKERTDS ESGDSSGEDPEGNKGFGRKVFSPV RSSFTHCRPPLDPEPPGPPDPPVAF KGYGSAPSSASSPASSASAATSFS LGSSTFKAQESGQSTAGPLRPPPP GAGGPATPSKATRFLPMDPATFRRK RPESVGGLEPPGVSIAAPPSSGGNI LQTLVLPNKEEQEGGGARVPSAPA PSLAYGAPAAPLSRPAATMVTNVVR PVSSSTPVIASKPFPTSGRAEASPN TAGARTEMGTGSRVPGGSPLGVSLV YSDKKSAAATSPAPHLVAGPLLGTVG KAPATVTNLLVGTTPGYGAPAPPVQF IAQGAPGGGTTAGSGAGAGSGPNGP VPLGILQPGALGKAGGITQVQYILPTL PQQLQVAPAPAPAGTKAAAPSGPAP TTSIRFTLPPGTSTNGKVLAAATPTP GIPILQSVSPAPPKQSVSPVQAPP GGSQQLLPGKVLVPLAAPSMSVRGG GAGQPLPLVSPFVSPVQNGAQPPS KIIQLTPVPVSTPSGLVPLSPATLPG PTSQPQKVLPLSSTRITYVQSAGGHA LPLGTSPASSQAGTVTSYGPTSSVAL GFTSLGSPGPAFVQPLLSAGQAPLLA PGQVGVSPVPSQLPPACAAPGGPVI TAFYSGSPAPTSSAPLAQPSQAPPSL VYTVATSTTPAATILPKGPPAPATAT PAPTSPFPSATAGSMTYSLVAPKAQR PSPKAPQKVKAALIASIPVGSFEAGAS GRPGPAPRQPLEPGVREPTAPESEL EGQTPPAPPPLPETWTPTARSSPPL PPPAERTSAKGPETMASKFPSSSSD WRVPGQGLNRRGEPPTPPSPAPAPA VAPGSSSESSGRAAGDTPERKEAA GTGKVKVRPPPLKKTFFDSVDNRVL SEVDFEERFAELPEFRPEEVLPSPTL QSLATSPRILGSYRKKRKNSTDLDL APEDPTSPKRKMRRSSCSSEPNT KSAKCEGDIFTFDRTGTEAEDVLGEL EYDKVPYSSLRRLDQRRALVMQLF QDHGFFPSAQATAAFQARYADIFPS KVCLQLKIREVRQKIMQAATPTEQPP GAEAPLPVPPPTGTAAPAPTSPAG

