

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A5YKK6	CNOT1_HUMAN	Homo sapiens	CCR4-NOT transcription complex subunit 1	20.096702	T746;S758;T764;S827;T880;T1038;T1041;T1042;S1043;T1044;T1051;T1055;T1058;S1061;T1418;S1775;T1780	S318;S1061	30379171;33214551;30059200;27655845;31492838;32119511;28657654;32574038;29351928;29237092;23301498;28510447	MNLDLSLALSQISYLVNLTCKNYRASQQEIQHIVNRHGPEADRHLLRC LFSHVDFSGDGKSSGKDFHQTFQLI QECALLITKPNFISTLSYAIIDNPLHYQ KSLKPAPHLFAQLSKVLKLSKVQVEVI FGLALLNSSSSDLRGFAAQFIKQKLP DLLRSYIDADVSGNQEGGFQDIAIEV LHLLSHLLFGQKGAFGVQEQIDA FLKTLRRDFPQERCVVVLAFLLYPEK RDILMDRILPDSGGVAKTMMESSLA DFMQEVGYGFCASIEECRNIIIVQFV REVTAQVARVLGMMARTHSGLTD GIPLQISAPGSGIWSGDKDKSDGAQ AHTWNVEVLIDVLKELNPSLNFKEV TYELDHPGFQIRDSKGLHNVVYGIQ RGLGMEVFPVDLIYRPWKHAEGQLS FIQHSLINPEIFCFADYPCHTVATDIL KAPPEDDNREIATWKSLLDIESLLRL AEVGQYEQVKQLFSFPIKHCPDMLV LALLQINTSWHTLRHELITLMPIFL GNHPNSAILHYAWHGQSQSPSIRQ LIMHAMAEWYMRGEQYDQAKLSRI LDVAQDLKALSMLLNGTPFAFVIDL AALASRREYLKLDKWLTDKIREHGE PFIQACMTFLKRRCP SILGGLAPEKD QPKSAQLPPETLATMLACLQACAGS VSQELSETILTMVANCSNVMNKAR QPPPGVMPKGRPPSASSLDAISPVQI DPLAGMTSLSIGGSAAPHTQSMQGF PPNLGSFSTPQSPAKAFPLSTPNQ TTAFFSGIGLSSQLPVGGLGTGSLTG IGTGALGLPAVNDPFPVQRKLGTSG LNQPTFQQSKMKPSDLSQVWPEAN QHFSKEIDDEANSYFQRIYNHPPHP TMSVDEVLEMLQRFKDKSTIKREREV FNCMLRNLFEYRFFPQYPDKELHI TACLFGGIIEKGLVTYMALGLALRYV LEALRKPFSGSKMYFYGIAALDRFKN RLKDYPOYCQHLSISHFMQFPHHL QEYIEYGQSRDPPVKMQGSITTPGS IALAQAQAQAQVPAKAPLAGQVSTM VTTSTTTTVAKTVTVTRPTGVSFKKD VPPSINTTNIDTLLVATDQTERIVEPP ENIQEKIAFIFNNLSQSNMTQKVEE LKETVKEEFMPWVSQYLVMKRVSIE PNFHSLYSNFLDTLKNPEFNKMVL NETYRNIKVLLTSDKAAANFSDRSLI KNLGHWLGMITLAKNKPIHTDLDV KSLLEAYVKGQQLLYVVPFVAKVL ESSIRSVVFRPPNPWTMAIMNVLAELHQEHDLKLNLFKFEIEVLCKNLALDI NELKPGNLLKDKDRLKNLDEQLSAP KKDVKQPEELPPITTTTTSTTPATNTT CTATVPPQPQYSYHDINVYSLAGLAP HITLNPITPLFQAHPQLKQCVRQAIE RAVQELVHPVVDRSIKIAMTTCEQIV RKDFALDSEESRMRIAHHMMRNL TAGMAMITCREPLLSISTNLKNSF

ASALRTASPPQREMMDQAAAQLAQ
DNCELACCFIQKTAVEKAGPEMDKR
LATEFELRKHARQEGRRYCDPVVLT
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EEFARNVPGFLPTNDLSQPTGFLAQ
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VEAVELLIRNHLVNMQQYDLHLAQS
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RSVAHVTEADLFHTIETLMRINAHS
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