

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion
Q8IX12	CCAR1_HUMAN	Homo sapiens	Cell division cycle and apoptosis regulator protein 1	27.733592	T17;T19;S22;S33;T54;T55;T115;S116;S118;S120;T121;T125;T129;S131;T134;S137;S138;T142;T179;T222;T240;S246;S253;S256;T258;T263;S287;S428;S655	S456;T627;T667;S685;S697;T861	20305658;30379171;31492838;38665916;33214551;32870666;23301498;31373491;35138101;34019948;27655845;21740066;32119511;34725712;33465208;29351928	MAQFGGQKNPPWATQFTATAVSP AALGVQPSLLGASPTTYTQQTALAA AGLTTQTPANYQLTQTAALQQQAAA AAAAALQQQYSQPQALYSVQQQLQQ PQQTLLTQPAVALPTSLSLSTPQPTA QITVSYPTPRSSQQQTQPKQKRVFTG VVTKLHDTFGFVDEDFVQLSAVKG KTPQVGDRLVLEATYNPNMPPFKWN AQRIQTLPNQNSQTQPLLKTPPAV LQPIAPQTTFGVQTQPQPSLLQAQI SAASITPLLQTPQPLLQPPQKAGL LQPPVRVRSQPQARRLDPSPRFSGR NDRGDQVPNRKRDRSRERERERRR SRERSPQRKRSRERSPRRERERSPR RVRRVVPRTYVQFSKFSLDCPCSDM MELRRRYQNLIPSDFDAQFTWVD AFPLSRPFQLGNYNCFVVMHREVES LEKNMAILDPPDADHLYSAKVMMLM ASPSMEDLYHKSCALAEQPQLRDG FQHPARLVKFLVGMKKGKDEAMAIG GHWSPSLDGPDPEKDPVSLIKTAIRC CKALTGIDLSVCTQWYRFAEIRYHRP EETHKGRTPAHVETVVLFFPDVWH CLPTRSEWETLSRGYKQLVEKLOG ERKEADGEQDEEEKDDGEAKEISTP THWSKLDPKTMKVNDLRKELESRA LSSKGLKSQLIARLTKQLKVEEQKEE QKELEKSEKEEDEDKRSKEDDKEE EERKRQEEIERQRERRRYILPDEPAI VHPNWAAKSGKFDCSIMSLSVLLDY RLEDNKEHSFEVSLFAELFNEMLQR DFGVRIYKSLSLPEKEDKKEKDKKS KKDERKDKEERDDEDEPKPKRRK SGDDKDKKEDRDERKKEDKRKDDS KDDDETEEDNNQDEYDPMEEAEAE DEEDDRDEEEMTKRDDKRINRYC KERPSKDKEKEKTQMITINRDLLMA FVYFDQSHCGYLLEKDLLEIILYTLGL HLSRAQVKLLNKVVLRESCFYRKL TDTSKDEENHEESESLQEDMLGNNR LLLPTPTVKQESKDVENVGLIVYNG AMVDVGSLLQKLEKSEKVAEVEQK LQLEEKTDDEKTIENLNSNKSLS GELREVKDLSQLQENLKISENMNL QFENQMNTIRNLSTVMDEIHTVLK KDNVKNEDKQKSKENGASV	True	False	1.781	5.0	2.852

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