

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membr
Q8CH18	CCAR1_MOUSE	Mus musculus	Cell division cycle and apoptosis regulator protein 1	32.195558	NaN	S453;T624;T664;S682;S694;T858	39627609;29125738;34887587	MAQFGGQKNPPWATQFTATAVSQP AALGVQQPSLLGASPTIYTQOTALAA AGLTTQTPANYQLTQTAALQQQAAA VLQQQYSQPQALYSVQQQLQQPQ QTILTQPAVALPTSLSLSTPQAAQIT VSYPTPRSSQQQTQPQKQRVFTGVV TKLHDTFGFVDEDFVFLGAVKGGKT PQVGDVRLVEATYNPNMPPFKWNAQ RIQTLPNQNSQTQPLLKTPTAVIQP IVPQTTFGVQAQPPQSLQAQISAA SITPLLQTPQPLLQQPQKAGLLQP PVRIVSQPQPARLLDPPSRFSGRND RGDQVNRKDDRSRERDRERRRRSR ERSPQRKRSRERSPRRERERSPRRV RRVVPRYTVQFSKFLDPCSCDMME LRRRYONLYIPSDFFDAQFTWVDAF PLSRPFQLGNFCYFVMMHREVESLE KNMAVLDPPDADHLYSAKVMLMAS PSMEDLYHKSCALAEQPDLRDFG QHPARLVKFLVGMKGKDEAMAIGG HWSPSLDGFNPEKDPVLIKTAIRCC KALTGIDL SVCTQWYRFAEIRYHRPE ETHKGRVPAHVETVWVFFPDVWHC LPTREWETLSRGYKQLVEKLGQE RKKADGEQDEEEKDDGEVKEIATPT HWSKLDPKAMKVNDRKELESRAL SSKGLKSQLIARLTKLKIEEQKEEQ KELEKSEKEEDEDKKSSEDDKEEQ ERKROEEVERQRQERRYILPDEPAI VHPNWAAKSGKFDCSIMSLSVLLDY RLEDNKEHSFEVSLFAELFNEMLQR DFGVRIYKSLLSLPEKEDKKKKEKKS KKEERKDKKEEREDDIDEKPKRRK SGDDKDKKEDRDERKKEEKRRKDD KDDDETEEDNNQDEYDPMEAEAE DEDDREEEVVRDDKRDVSRICK DRPAKDKEKEKPMVTNDRLLMA FVYFDQSHCGYLLEKDLLEIILYTLGL HLSRAQVKLLNKVVLRESCFYRKL TDTSKDDENHESEALQEDMLGNR LLLPTPTIKQESKGEENVGLIVYNG AMVDVGSLLQKLEKSEKVAEVEQK LQLEEKTDDEGKTILNLENSNKS SGELREVKKDLGQLQENLEVS NLQFENQLNKTLRNLSTVMDDIHTV LKKDNVKSEDRDEKSKENGSGV	True	False	1.737	4.714	2.752	0.853	0.828	1.125