

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
O43896	KIF1C_HUMAN	Homo sapiens	Kinesin-like protein KIF1C	22.903221	T1010	S295;S494;S674;S676;S915;S1033;T1083;S1092	38253038	MAGASVKAVVRVRFNARETSQDAK CVVSMQGNNTTSIINPKQSKDAPKSF TFDYSYWSHTSTEDPQFASQQQVYR DIGEEMLLHAFEGYNVCIFAYGQTG AGKSYTMMGRQEPGQQGIVPQLCE DLFSRVSENSAQLSYSVEVSYMEIY CERVRLDLLNPKSRGSLRVREHPILG PYVQDLKSLAVTSYADIADLMDCGN KARTVAATNMNETSSRSHAVFTIVF TQRCHDQLTGLDSEKVSISLVDLA GSERADSSGARGMRLKEGANINKSL TTLGKVISALADMQSKKRKSDFIPYR DSVLTWLLKENLGGNSRTAMIAALS PADINYEETLSTLRYADRTKQIRCNA IINEDPNARLIRELQEEVARLRELLM AQGLSASALEGLKTEEGSVRGALPA VSSPPAPVSPSSPTTHNGELEPSFSP NTESQIGPEEAMERLQETEKIAELN ETWEEKLRKTEALRMEREALLAEM GVAVREDGGTVGVFSPKTPHLVNL NEDPLMSECLLYHIKDGVTRVGQVD MDIKLTGQFIREQHCLEFRSIPQPDGE VVVTLEPCEGAETYVNGKLVTEPLVL KSGNRIVMGKNHVFRFNHPEQARL ERERGVPPPPGPPSEPVDFWFAQKE LLEQQGIDIKLEMEKRLQDLENOYR KEKEEADLLEQQRLYADSDSGDSDS DKRSCESWRLISSLRQLPPTTVQT IVKRCGLPSSGKRRAPRRVYQIPQRR RLQGDPRWATMADLKMQAVKEIC YEVALADFRHGRAEIEALAALKMRE LCRTYKPDGPGDAWRAVARVVD TVGEEEGGGAGSGGSEEGARGAE VEDLRAHIDKLTGILQEVKLQNSSKD RELQALRDRMLRMERVIPLAQDHE DENEEGGEVPWAPPEGSEAAEEAAP SDRMPSARPPSPPLSSWERVSRLEME EDPAFRGRRLRWLQEQRLRLQGLQ GSGGRGGGLRRPPARFVPPHDCKLR FPFKSNPQHRESWPGMGSGEAFPT LQPEEVTPHPATPARRPPSPRRSH HPRRNSLDGGRSRGAGSAQPEPQ HFQPKKHNSYPQPYPQARPPPGP RYPPYTPPRMRRQRSAPDLKESGA AV	False	False	3.913	3.825	1.888	4.267	4.338	1.317	0.864