	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	O35071	KIF1C_MOUSE	Mus musculus	Kinesin-like protein KIF1C	25.385059	NaN	S295;S494;S674;S676;S914;S1031;T1080;S1089	28528544	MAGASVKVAVRVRPFNARETSQDAK CVVSMQGNTTSIINPKQSKDAPKSF TFDYSYWSHTSVEDPQFASQQQVYR DIGEEMLLHAFEGYNVCIFAYGQTG AGKSYTMMGRQEPGQQGIVPQLCE DLFSRVNVNQSAQLSYSVEVSYMEI YCERVRDLLNPKSRGSLRVREHPIL GPYVQDLSKLAVTSYADIADLMDCG NKARTVAATNMNETSSRSHAVFTIV FTQRSHDQLTGLDSEKVSKISLVDLA GSERADSSGARGMRLKEGANINKSL TTLGKVISALADLQSKKRKSDFIPYR DSVLTWLLKENLGGNSRTAMIAALS PADINYEETLSTLRYADRTKQIRCNA VINEDPNARLIRELQEEVARLRDLL MAQGLSASALGGLKVEEGSPGGVLP PASSPPAPASPSSPPPHNGELEPSFS PSAEPQIGPEEAMERLQETEKIIAEL NETWEEKLRKTEALRMEREALLAE MGVAVREDGGTVGVFSPKKTPHLV NLNEDPLMSECLLYHIKDGVTRVGQ VDVDIKLTGQFIREQHCLFRSIPQPD GEVMVTLEPCEGAETYVNGKLVTEP LVLKSGNRIVMGKNHVFRFNHPEQ ARLERERGVPPPPGPPSEPVDWNFA QKELLEQQGIDIKLEMEKRLQDLEN QYRKEKEEADLLLEQQRLYADSDSG EDSDKRSCEESWRLISSLREQLPPTT VQNIVKRCGLPSSGKRRAPRRVYQIP QRRRLQGKDPRWATMADLKMQAVK EICYEVALADFRHGRAEIEALAALKM RELCRTYGKPEGPGDAWRAVARDV WDTVGEEEGCGGGGGGSEEGARGA EVEDLRAHIDKLTGILQEVKLQNSSK DRELQALRDRMLRMERVIPLTQDLE DDNDESGLVTWAPPEGPEAVEETVP NDHSPAVRPTSPPLSSWERVSRLME EDPAFRRGRLRWLKQEQLRLQGLQ GAGGRGGGLRRPPARFVPPHDCKLR FPFKSNPQHRESWPGMGSGEAPAP QPPEEVTVPPAPPNRRPPSPRRPHRS RRNSLDGGSRSRGGGSTQPEPQHL RPQKHNGYPQQPQPSPAQRPGPRYP PYTTPPRMRRQRSAPDLKESGAAV	None	None	None	None	None	None	None	None	None