

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q5T655	CFA58_HUMAN	Homo sapiens	Cilia- and flagella-associated protein 58	30.442579	S719	NaN	39894887;36240223	MAEEKGGKQVLEESAFEEMERDFQ GVLHELSDGDKSLEKFRIEYERLHAV MKKSYDNEKRLMAKCRELNAEIVV NSAKVATALKLSQDDQTTIASLKKEI EKAWKMDVSAYDKEQKAKETILALK EEIVNLTKLVEQGSGLSMDQHSNIR DLLRFKEEVTKERDQLLSEVVKLRE SLAQTEQQQETERSKEEAHAISQ FQQEIQQRQNEASREFRKKKELEKE LKQIQADMDSRQTEIKALQQYVQKS KEELQKLEQQLKEQKILNERAAKEL EQFQMRNAKLQQENEQHSLVCEQL SQENQKALELKAKEEEVHOMRLDI GKLNKIREQIHKKLHHTEDQKAEVE QHKETLKNQIVGLEREVEASKKQAE LDRKAMDELLRERDILNKNMLKAV NATQKQTDLVKLHEQAKRNLEGEIQ NYKDEAQKQRKIIHFLEKERDRYINQ ASDLTQKVLNMEDIKVRQIFDY RKKIAESEIKLQQQNLYEAVRSDR NLYSKNLVEAQDEITDMKRKLKIMI HOVDELKEDISAKESALVKLHLEQQ RIEKEKETLKAELQKLRQQALETKHF IEKQAEERKLLRIIAEADGERLRQK KELDQVISERDILGSQLVRRNDELAL LYEKIKIQQSVLNKGESQYNQRLED MRILRLEIKLRREKGLARSMANVE ELRQEFFHMORELLKERTRCRALEE ELENPLNVHRWRKLEASDPNAYELI QKIHTLQKRLISKTEEVVEKELLQE KEKLYMELKHVLRQPGPEAAEQLK LYRRTLHDKKQQLKVLSSSELNMYEV QSKEYKYEVEKLTNELQNLKKKYLA QKRKEQLQKNDTAPMDNTFLMVK PNGPGFTGGGFPLRSTKMTF	False	False	1.734	3.291	False	False	1.078	False	4.0