	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	Q5TEA3	DAAF9_HUMAN	Homo sapiens	Dynein axonemal assembly factor 9	28.275761	NaN	NaN	39894887	MDVYPPRRQGLPRARSPGGSSRGSP SVSCSRLRQVQSILTQSSKSRPDGIL CILGIDSRYNEGCRELANYLLFGLYN QNTSDFEKTGFSEEVLDDVIILIKSD SVHLYCNPVNFRYLLPYVAHWRNL HFHCMTENEYEDEEAAEEFKITSFV DMVRDCSRIGIPYSSQGHLQIFDMF VVEKWPIVQAFALEGIGGDGFFTMK YELQDVSLNLWNVYSKMDPMSLES LLSDDLVAFEHQWTSFFANFDTEIP FLLELSESQAGEPFRSYFSHGMISS HITENSPNRQPFVLFGNHSTRENLN AGNFNFPSEGHLVRSTGPGGSFAKH MVAQCVSPKGPLACSRTYFFGATHV PYLGGDSKLPKKTEQIRLLSQIYAAVI EAVLAGIACYAKTSSLTKAKEVAEQT LGSGLDSFELIPFKAALRSKMTFHIH AVNNQGRIVPLDSEDSLSFVKTACM AVYDIPDLLGGNGCLGSVVFSESFLT SQILVKEKDGTVTTETSSVVLTAAVP RFCSWLVEDNEVKLSEKTQQAVRG DESFLGTYLTGGEGAYLYSSNLQSW PEEGNVHFFSSGLLFSHCRHRSIIIS KDHMNSISFYDGDSTSTVAALLIDFK SSLLPHLPVHFHGSSNFLMIALFPKS KIYQAFYSEVFSLWKQQDNSGISLKV IQEDGLSVEQKRLHSSAQKLFSALS QPAGEKRSSLKLLSAKLPELDWFLQ HFAISSISQEPVMRTHLPVLLQQAEI NTTHRIESDKVIISIVTGLPGCHASEL CAFLVTLHKECGRWMVYRQIMDSS ECFHAAHFQRYLSSALEAQQNRSAR QSAYIRKKTRLLVVLQGYTDVIDVVQ ALQTHPDSNVKASFTIGAITACVEPM SCYMEHRFLFPKCLDQCSQGLVSNV VFTSHTTEQRHPLLVQLQSLIRAANP AAAFILAENGIVTRNEDIELILSENSF SSPEMLRSRYLMYPGWYEGKLNAG SVYPLMVQICVWFGRPLEKTRFVAK CKAIQSSIKPSPFSGNIYHILGKVKFS DSERTMEVCYNTLANSLSIMPVLEG PTPPPDSKSVSQDSSGQQECYLVFIG CSLKEDSIKDWLRQSAKQKPQRKAL KTRGMLTQQEIRSIHVKRHLEPLPA GYFYNGTQFVNFFGDKTDFHPLMD QFMNDYVEEANREIEKYNQELEQQ EYHDLFELKP	False	False	0.81	2.625	0.441	0.188	False	False	False