	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	Q86VH2	KIF27_HUMAN	Homo sapiens	Kinesin-like protein KIF27	26.21016	T316;S322	S643;S646;S672;S675;S704;S999;S1367;S1389	29351928	MEEIPVKVAVRIRPLLCKEALHNHQ VCVRVIPNSQQVIIGRDRVFTFDFVF GKNSTQDEVYNTCIKPLVLSLIEGYN ATVFAYGQTGSGKTYTIGGGHIASVV EGQKGIIPRAIQEIFQSISEHPSIDFN VKVSYIEVYKEDLRDLLELETSMKDL HIREDEKGNTVIVGAKECHVESAGE VMSLLEMGNAARHTGTTQMNEHSS RSHAIFTISICQVHKNMEAAEDGSW YSPRHIVSKFHFVDLAGSERVTKTG NTGERFKESIQINSGLLALGNVISAL GDPRRKSSHIPYRDAKITRLLKDSLG GSAKTVMITCVSPSSSNFDESLNSLK YANRARNIRNKPTVNFSPESDRIDE MEFEIKLLREALQSQQAGVSQTTQI NREGSPDTNRIHSLEEQVAQLQGEC LGYQCCVEEAFTFLVDLKDTVRLNE KQQHKLQEWFNMIQEVRKAVLTSF RGIGGTASLEEGPQHVTVLQLKREL KKCQCVLAADEVVFNQKELEVKELK NQVQMMVQENKGHAVSLKEAQKV NRLQNEKIIEQQLLVDQLSEELTKLN LSVTSSAKENCGDGPDARIPERRPYT VPFDTHLGHYIYIPSRQDSRKVHTSP PMYSLDRIFAGFRTRSQMLLGHIEE QDKVLHCQFSDNSDDEESEGQEKS GTRCRSRSWIQKPDSVCSLVELSDT QDETQKSDLENEDLKIDCLQESQEL NLQKLKNSERILTEAKQKMRELTINI KMKEDLIKELIKTGNDAKSVSKQYSL KVTKLEHDAEQAKVELIETQKQLQE LENKDLSDVAMKVKLQKEFRKKMD AAKLRVQVLQKKQQDSKKLASLSIQ NEKRANELEQSVDHMKYQKIQLQR KLREENEKRKQLDAVIKRDQQKIKEI QLKTGQEEGLKPKAEDLDACNLKRR KGSFGSIDHLQKLDEQKKWLDEEVE KVLNQRQELEELEADLKKREAIVSK KEALLQEKSHLENKKLRSSQALNTD SLKISTRLNLLEQELSEKNVQLQTST AEEKTKISEQVEVLQKEKDQLQKRR HNVDEKLKNGRVLSPEEEHVLFQLE EGIEALEAAIEYRNESIQNRQKSLRA SFHNLSRGEANVLEKLACLSPVEIRT ILFRYFNKVVNLREAERKQQLYNEE MKMKVLERDNMVRELESALDHLKL QCDRRLTLQQKEHEQKMQLLLHHF KEQDGEGIMETFKTYEDKIQQLEKD LYFYKKTSRDHKKKLKELVGEAIRRQ LAPSEYQEAGDGVLKPEGGGMLSEE LKWASRPESMKLSGREREMDSSAS SLRTQPNPQKLWEDIPELPPIHSSLA PPSGHMLGNENKTETDDNQFTKSH SRLSSQIQVVGNVGRLHGVTPVKLC RKELRQISALELSLRRSSLGVGIGSM AADSIEVSRKPRDLKT	None	None	None	None	None	None	None	None	None