

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	extrace region
P52732	KIF11_HUMAN	Homo sapiens	Kinesin-like protein KIF11	22.998462	NaN	T458;T925;T926;S1033	34229054;23301498;28510447	MASQPNSSAKKKEEGKNIQVVVRC RPFNLAERKASAHSIVECDPVRKEVS VRTGGLADKSSRKTYTFDMVFGAST KQIDVYRSVVCILDEVIMGYNCTIF AYGQTGTGKTFMEGERSPNEEYT WEEDPLAGIIPRTLHQIFEKLTNDGT EFSVKVSLLEIYNEELFDLLNPSSDV SERLQMFDDPRNKRGVIIKGLEEITV HNKDEVYQILEKGAARKRTAATLMN AYSSRSHSVFVTHMKETTIDGEEEL VKIGKLNLDLAGSENIGRSGAVDK RAREAGNINQSLTLGRVITALVERT PHVPYRESKLTIRIQDSLGGRTTSII ATISPASLNLEETLSTLEYAHRANKNL NKPEVNOKLTKKALIKYEETEEIERLK RDAAAAREKNGVYISEENFRVMSGK LTVQEEQIVELIEKIGAVEEELNRVT ELFMDNKNELDQCKSDLQNKQTQEL ETTQKHLQETKLQLVKEEYITSALES TEEKLHDAASKLLNTVEETTKDVSG LHSKLDRKAVDQHNAEAQDIFGK NLNSLFNNMEELIKDGSSKQKAML EVHKTLFGNLLSSVSALDTITVAL GSLTIPENVSTHVSQIFNMLKEQS LAAESKTVLQELINVLKTDLLSLEM ILSPTVVLSILKINSQLKHIFKTSLTVA DKIEDQKKELDGFLSILCNNLHELQ ENTICSLVESQKQCGNLTEDLKTIKQ THSQELCKLMNLWTERFCALEEKC ENIQKPLSSVQENIQKSKDIVNKM TFHSQKFCADSDGFSQELRNFNQE GTKLVEESVKHSDKLNGLNLEKISQE TEQRCELNTRTVYFSEQWVSSLNE REQELHNLLEVVSQCCEASSSDITE KSDGRKAAHEKQHNIFLDQMTIDED KLIAQNLELNETIKIGLTKLNCFLEQ DLKLDIPTGTTTPQRKSYLYPSTLVRT EPREHLLDQLKRKQPELLMMLNCS ENNKEETIPDQVVEEAVLGQYTEEP LSQEPSVDAGVDCSSIGGVPFQHK KSHGKDKENRGINTLERSKVEETTE HLVTKSRLPLRAQINL	True	False	4.914	4.304	1.737	1.427	1.553	1.893	1.669