

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
Q6P9P6	KIF11_MOUSE	Mus musculus	Kinesin-like protein KIF11	27.796441	NaN	T457;T925	39627609;34887587	MASQPSSLLKKEEKGRNIQVWVRCR PFNLAERKANAHSVVECDHARKEVS VRTAGLTDKTSKKTYYTFDMVFGAST KQIDVYRSVVCPIIDEVIMGYNCTIF AYGQTGTGKTFMEGERSPNEVYT WEEDPLAGIIPRTLHQIFELTDNGT EFSVKVSLLEIYNEELFDLLSPSSDV SERLQMFDDPRNKRGVIIKGLEEITV HNKDEVYQILEKGAARKRTTAATLMN AYSSRSHSVFVSTIHMKETTIDGEEEL VKIGKLNLDLAGSENIGRSGAVDK RAREAGNINQSLTLGRVITALVERT PHIPYRESKLRILQDSLGGRTSII ATISPAFNLLEETLSTLEYAHRAKNI MNKPEVNQKLTKKALIKKEYTEEIERL KRDAAAAREKNGVYISEESFRAMNG KVTVOEEQIVELVEKIAVLEEEELSKA TELFMDSKNELDQCKSDLQTKTQEL ETTQKHLQETKQLVKEEYVSSALE RTEKTLHDTASKLLNTVKETTRAVS GLHSLDRKRAIDEHNAEAQESFGK NLNSLFNNMEELIKDGS AKQKAML DVHKTFLFGNLMSSSVSALDTITTTAL ESLVSIPENV SARVSQISDMILEEQS LAAQSKSVLQGLIDELVTDLFTSLKTI VAPSVV SILNINKLOLHIFRASSTVA EKVEDQKREIDSFLSILCNLHELRE NTVSSLVESQKLCGDLTEDLKTIKET HSQELCQLSSSWAERFCALEKKYEN IQKPLNSIQENTELRSTDIINKTTVHS KKILAESDGLLQELRHFNQEGTQLV EESVGHCSLNSNLETVSEQEITQKC GTLNTSTVHFSDQWASCLSKRKEEL ENLMEFVNGCCKASSEITKKVREQ SAAVANQHSSFVAQMTSDEESCKA GSLELDKTIKTLTKLNCFLKQDLKL DIPTGMTPERKKYLYPTTLVRTEPRE QLDQLQKKQPPMMLNSSEASKET SQDMDEEREALQCTEELVSPETTE HPSADCSRRGLPFFQRKKPHGKDK ENRGLNPVEKYKVEEASDLSIKSRL PLHTSINL	True	False	4.143	3.495	1.657	1.396	1.473	1.815	1.537

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