

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac 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	25.161187	NaN	T457;T925	34887587	MASQPSSLKKKEEKGRNIQVVVRCR PFNLAERKANAHSVVECDHARKEVS VRTAGLTDKTSKTTYTFDMVFGAST KQIDVYRSVVCPILEDEVIMGYNCTIF AYGQTGTGKFTTMEGERSPNEVYT WEEDPLAGIIPRTLHQIFEKLTDNGT EFSVKVSLLEIYNEELFDLLSPSSDV SERLQMFDDPRNKRGVIIKGLEEITV HNKDEVYQILEKGAARRTAATLMN AYSSRSHSVFSVTIHMKETTIDGEEL VKIGKLNLDLAGSENIGRSGAVDK RAREAGNINQSLTLGRVITALVERT PHIPYRESKLTRIQSLGGRTRTSII ATISPAFNLEETLSTLEYAHRAKNI MNKPEVNQKLTKKALIKYEETEEIERL KRDLAAAREKNGVYISEESFRAMNG KVTVOEEQIVELVEKIAVLEEELSKA TELFMDSKNELDQCKSDLQTKTQEL ETTQKHLQETKLQLVKEEYVSSALE RTEKTLHDTASKLLNTVKETTRAVS GLHSKLDRKRAIDEHNAEAQESFGK NLNSLFNNMELIKDGSQKQKAML DVHKTFLFGNLMSSSVSALDTITTTAL ESLVSIPENV SARVSQISDMILEEQS LAAQSKSVLQGLIDELVTDLFTSLKTI VAPSVVILNINKQLQHIFRASSTVA EKVEDQKREIDSFLSILCNNLHELRE NTVSSLVESQKLCGDLTEDLKTIKET HSQELCQLSSWAERFCALEKKYEN IQPLNSIQENTELRSTDIINKTTVHS KKILAESDGLLQELRHFNQEGTQLV EESVGHCSLNSNLETVSQEITQKC GTLNTSTVHFSDQWASCLSKRKEEL ENLMEFVNGCCKASSEITKKVREQ SAAVANQHSSFVAQMTSDEESCKA GSELDKTIKTGLTKLNCFLKQDLKL DIPTGMTPERKKLYPTTLVRTEPRE QLLDQLQKQPPMMLNSSEASKET SQDMDEEREALEQCTEELVSPETTE HPSADCSSRGLPFFQRKKPHGKDK ENRGLNPVEKYKVEEASDLSISK SRL PLHTSINL	True	False	4.142	4.398	1.654	1.396	1.472	1.815	1.538

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