	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	E9Q175	E9Q175_MOUSE	Mus musculus	Unconventional myosin-VI	26.781227	NaN	NaN	36064721;34887587	MEDGKPVWAPHPTDGFQMGNIVDI GPDSLTIEPLNQKGKTFLALINQVFP AEEDSKKDVEDNCSLMYLNEATLLH NVKVRYSKDRIYTYVANILIAVNPYF DIPKIYSSDTIKSYQGKSLGTMPPHV FAIADKAFRDMKVLKMSQSIIVSGES GAGKTENTKFVLRYLTESYGTGQDID DRIVEANPLLEAFGNAKTVRNNNSS RFGKFVEIHFNEKSSVVGGFVSHYL LEKSRICVQGKEERNYHIFYRLCAGA SEDIREKLHLSSPDNFRYLNRGCTR FFANKETDKQILQNRKSPEYVKAGS LKDPLLDDHGDFIRMCTAMKKIGLD DEEKLDLFRVVAGVLHLGNIDFEEA GSTSGGCNLKNKSAPSLEYCAELLG LDQDDLRVSLTTRVMLTTAGGTKGT VIKVPLKVEQANNARDALAKTVYSH LFDHVVNRVNQCFPFETSSYFIGVL DIAGFEYFEHNSFEQFCINYCNEKL QQFFNERILKEEQELYQKEGLGVNE VHYVDNQDCIDLIEVKLVGILDILDE ENRLPQPSDQHFTSVVHQKHKDHF RLTIPRKSKLAVHRNLRDDEGFIIRH FAGAVCYETTQFVEKNNDALHMSL ESLICESRDKFIRALFESSTNNNKDT KQKAGKLSFISVGNKFKTQLNLLLD KLRSTGASFIRCIKPNLKMTSHHFE GAQILSQLQCSGMVSVLDLMQGGFP SRASFHELYNMYKKYMPEKLARLDP RLFCKALFKALGLNEVDYKFGLTKV FFRPGKFAEFDQIMKSDPDHLAELV KRVNLWLVCSRWKKVQWCSLSVIK LKNKIKYRAEACIKMQKTIRMWLCK RRHKPRIDGLVKVGTLKKRLDKFNE VVSALKDGKPEVNRQIKNLEISIDAL MAKIKSTMMTREQIQKEYDALVKSS EDLLSALQKKKQQEEEAERLRRIQE EMEKERKRREEDEERRRKEEEERR MKLEMEAKRKQEEEERKKREDDEK RIQAEVEAQLARQREEESQQQAVLA QECRDRELALRIAQNESELISDEAQG DMALRRGPAVQATKAAAGTKKHDLS KWKYAELRDTINTSCDIELLAACREE FHRRLKVYHAWKSKNKKRNTETEQ RAPKSVTDYAQQNPAAQLPARQQEI DMKRQQRFFRIPFIRPADQYKDPQN KKKGWWYAHFDGPWIARQMELHP DKPPILLVAGKDDMEMCELNLEETG LTRKRGAEILPRQFEEIWERCGGIQY LQSAIESRQARPTYATAMLQNLLK	None	None	None	None	None	None	None	None	None