

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
E9Q634	MYO1E_MOUSE	Mus musculus	Unconventional myosin-Ie	29.280344	NaN	S1001	29187734;34887587	MGSKGAYRYHWQSHNVKHSGVDD MVLSSKITESSIVENLKKRYMDDYIF TYIGSVLISVNPFKQMPYFGEKEVEM YQGAAQYENPPHIYALADSMYRNMI IDRENQCVIHSGEGAGKTVAAKYIM SYVSRVSGGGPKVQHVKDILQSNPL LEAFGNAKTVRNNNSSRFKGYFEIQ FSPGGEPDGGKISNFFLEKSRVVMR NPGERSFHIFYQLIEGASPEQKQSLG ITSMDDYYLSSGYSKVDIDDKRD FQETLHAMNVIGIFSEEQTLVLQIVA GILHLGNISFKEVGNAAVESEEFLLA FPAYLLGINQDRLKEKLTSRQMDSK WGGKSESIHVTLNVEQACYTRDALA KALHARVDFLVDLVSINKAMEKDHEE YNIGVLDIYGFEIFQKNGFEQFCINF VNEKLQOIFIELTLKAEQEEYVQEGI RWTPIEYFNKIVCDLIESKVNPPGI MSILDDVCATMHAVGEGADQTLLO KLQMIGSHEHFNSWNQGFIIHHY AGKVSVDMDGFCERNRDLVFMDLI ELMQSSELPFIKSLFPENLQADKKG RPTTAGSKIKKQANDLVSTLMKCTP HYIRCIPNETKKPKDWEESRVKHQ VEYLGLENIRVRRAGYARRVFQKF LQRYAILTKATWPVWRGDEKQGVLLH LLQSVNMDSDFQLGRSKVFIKAP SLFLLLEMRERKYDGYARVIQKTWR KFVARKKYVQMREEASDLLNKKER RRNSINRNFIGDYIGMEERPELQOF VGKREKIDFADTVTKYDRRFKGVKR DLLLTPKCLYLGREKVKQGPDKGVV KEVLKRRIEVERILSVSLTMQDDIFI LHEQEYDSLLESVFKTEFLSLAKRY EEKTKQLPLKFSNTLELKLKKNW GPWSAGSRQVQFHQGFGLAILKP SNKVLQVSIGPLPKNSRPTRRNTV TSRGYPGGTKNNYPMRAAPAPPGCH QNGVIRNQFVPPHAFGNQRSNQK SLYTSMARPPLPROQSTGSDRLSQT PESLDFLKVPDQCVAGVRRROTSSRP PPAGGRPKQPKPKQVPOCKALYA YDAQDTDELSFNANDIIDIKEDPSG WWTGRLRGKQGLFPNNYVTKI	False	False	3.341	1.974	1.135	0.963	0.93	3.666	1.074