

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
Q12965	MYO1E_HUMAN	Homo sapiens	Unconventional myosin-1e	21.056457	NaN	S980;S1002	33214551;28510447	MGSKGVYQYHWQSHNVKHSVDD MVLKSKITENSIVENLKKRYMDDYIF TYIGSVLISVNPFFKQMPYFGEKEIEM YQGAAQYENPPHIYALADNMYRNM IDRENQCVIISGSGAGKTVAAYIM SYISRVSGGGTKVQHVKDILQSNPL LEAFGNAKTVRNNNSSRFKGFYFIQ FSPGGEFDGGKISNFFLEKSRVVMR NPGERSFHIFYQLIEGASAEQKHSLG ITSMDDYVYLSLSGSYKVVDDIDDRRE FQETLHAMNVIGIFAEEQTLVLQIVA GILHLGNISFKEVGNAAVESEEFLLA FPAYLLGINQDRLKEKLTSRQMDSK WGGKSESIHVTLNVEQACYTRDALA KALHARVDFLVDVSNKAMEKDHEE YNIGVLDIYGFEIFQKNGFEQFCINF VNEKLLQIFIELTLKAEQEEYVQEGI RWTPIEFNKNKIVCDLIENKVNPPGI MSILDDVCATMHAVGEGADQTLLO KLQMQIGSHEHFNSWNQGFIIHHY AGKVSVDMDGFCERNRDLVFMDLI ELMQSELFPFKSLFFENLQADKKG RPTTAGSKIKKQANDLVSTLMKCTP HYRCIKPNETTKPRDWEEERVKHQ VEYLGLENIRVRRAGYAYRRIFQKF LQRYAILTKATWPSWQGEKQGVLLH LLQSVNMDSDQFQLGRSKVFIKAP SLFLLLEEMREKYDGYARVIQKSWR KFVARKKYVQMREEASDLLLNKKER RRNSINRNFIGDYIGMEEHPELQQF VGKREKIDFADVTYKDRRFKGVKR DLLLLTPKCLYLIGREKVKQGPDKGLV KEVLKRKIEIERILSVSLSTMQDDIFI LHEQEYDSLLESVFKTEFLSLAKRY EEKTQKQLPLKFSNTLELKKENW GPWSAGGSRQVQFHQGFGLAVLK PSNKVLQVSIQGPLPKNSRPTRRNTT QNTGYSSTQANYPVRAAPPVPPGY HQNGVIRNQVYPYPHAPGSQRSNQ KSLYTSMARPPLPRQSTSSDRVSO TPESLDFLKVPDQGAAGVRRQTSR PPPAGGRPKPQPKPQVPOCKALY AYDAQDTDELSFNANDIHKEDPS GWWTGRLLRKGQGLFPNNYVTKI	False	False	3.277	1.537	1.175	0.763	0.822	3.648	4.273