

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	p
Q9WTI7	MYO1C_MOUSE	Mus musculus	Unconventional myosin-Ic	30.973994	NaN	S408;S536;S864;S1041;S10	29187734;34887587;39627609;36064721;37507081	MALQVELIPTGEIIRVVHPHRPCKLA LGSDGVRVTMESALTARDRVGVQDF VLENFTSEAAFIENLRFRRENLIY TYIGPVLVSVNPNYRDLQIYSRQHMER YRGVSFYEVPPHFLFAVDTVYRALRT ERRDQAVMISGESGAGKTEATKRLL QFYAETCPAPERGGAVDRLLQSNP VLEAFGNAKTLRNDNSSFQKYM VQDFDKGAPVGGHILSYLLEKSRVV HQNHGERNFHFVYQLLEGGEETL RRLGLERNPQSYLYLVKQCAKVSSI NDKSDWVMRKALSVIDFTEDEVE DLLSIVASVLHLGNIHFAAEDSNA QVTTENQLKYLTRLLGVEGTTREAL THRKHAKGEELLSPLNLEQAAYARD ALAKAVYSRTFTWLVKRKINRSLASKD AESPSWRSTTVLGLLDIYGFVQHS NSFEQFCINYCNEKLQQLFIELTLKS EQEEYEAEGIAWEPVQYFNNKIICDL VEEKFKGHSILDEECLRPGEATDLTF LEKLEDTVKPHPHFLTHKLADQKTR KSLDRGEFRLHYAGEVTVSFTGFL DKNNDLLFRNLKETMCSSMNPIMA QCFDKSELSDKRPETVATQFKMSL LQLVEILRSKEPAYIRCIPNDAKQP GRFDEVLRHQVKYLGLENLRRVRR AGFAYRRKYEAFQRYKSLCPETWVP MWAGRQDGVAVLVRHLGKPEEY KMGRTKIFIRFPKTLFATEDSLEVRR QSLATKIQAAWRGFHWRQKFLRVK RSAICIQSWWRGTLGRRKAARKKWA AQTIRRLIRGFILRHSPRCPENAFFL DHVRASFLNLRRQLPRNVLDTSWP TPPPALREASELLRELCMKNMVVK YCRSISPEWKQQLQKAVASEIFKG KKDNYQSVPRFLFISTRGTEEISPR VLQSLGSEPIQYAVPVVKYDRKGYKP RPRQLLTPSAVVIVEDAKVKQRIDY ANLTGISVSSLSDFVLHVQREDN KQKGDVVLQSDHVIEITLKTALSAD RVNNININQGSITFAGGPRDGIIDF TSGSELLITKAKNGHLAVVAPRLNSR	True	True	4.673	4.358	1.214	1.032	1.246	5.