

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
088329	MYO1A_MOUSE	Mus musculus	Unconventional myosin-1a	25.875327	S723	NaN	30059200	MPLLEGPVGVEDLILLEPLDEESLIK NLQLRYENKEIYTYIGNVVISMNPYE QLPIYGPEFIAKYRDYTFYELKPHIYA LANVAYQSLKDRDRDQCILITGESGA GKTEASKLVMSYVAAVCGKGEQVNS VKEQLLQSNPVLAEAFGNAKTIRNNN SSRFGKYMIDIEFDKGSPLGGVITNY LLEKSRVVKQLKGERNFHIFYQLLA GADAQLLKALKLEEDTSVYGYLNGE VSKVNGMDDASNFRAVQHAMSIVIG FSEEEIROVLEVTALVVLKLGNVKLT EFQANGIPASGICDGKGIQEIEMMM GLNSTELERALCSRTMETGKEKVVV VLNVTQAQYARDALAKNIYSRLFDWI VKRINESIKVGTGEKKKVMGVLDIYG FEILEDNSFEQFVINYCNERLQQVFI ELTLKEEQEEYKREGIPWTKVEYFD NGIICNLIEHSORGILAMLDEECLRP GVVSDSTFLAKLNQLFSKSHSHYESK VSQNAQRQYDRTMGLSCFRISHYAG KVITYNVTGFIDKNNDLLFRDLSQTM WKAQHPLLKSLFPEGNPKEASLKR PTAGTQFKNSVAVLMKNLYSKNPNY IRCIKPNDDQQKGRFTSEMVMVQAR YLGLENVRRVRRAGYAFRQGYKPF ERYRLLSRSTWPRWNGDDREGVEK VLGSLTSSSEELAYGKTIFIRSPKTL FYLEEQRRLRLQQLATLIQKVYRGW RCRTHYQQMRKSQILISAWFRGNKQ KKHYGKIRSSVLLIQAFVRGWRARK NYRKYFRSGAALTLANFIYQSMQAK FLLNLKKNLPSTKVDNTWPAAPYR CFNTANQELQRLFYQWCKKFRDQ LSPKQVQTLREKLCASELFGKKASY PQSVPIPFPGDYIGLQGNPKLQRLKG REEGPVLVADTVKKNRNGKTSAR ILLTKGHVILTDAKKSQAQIVIGLED VAGVSVSSLQDGLFSLHSEMSSAV SKGDILLVSDHVVELLTKMYQAVLD ATORQLSVTVTEKFSVRFKEGSVAV KVIQPEGGNRKLIKCKKGSNAME VTVR	False	True	3.179	2.465	1.165	0.796	1.128	5.0	1.184