

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9UBC5	MYO1A_HUMAN	Homo sapiens	Unconventional myosin-1a	22.903221	S257	NaN	38253038	MPLLEGSVGVEDLVLEPLVEESLLK NLQLRYENKEIYTYIGNVVISVNPYQ QLPIYGPEFIAKYQDYTFYELKPHIYA LANVAYQSLRDRDRDQCILITGESGS GKTEASKLVMSYVAAVCGKGEQVNS VKEQLLQSNPVLAEAFGNAKTIRNNN SSRFGKYMIEFDFKGSPLGGVITNY LLEKSRLVKQLKGERNFHIFYQLLAG ADEQLLKALKLERDITTYAYLNHEV SRVDGMDDASSFRAVQSAMAVIGFS EEEIROVLEVTSMVLKLGNVLVADE FQASGIPASGIRDGRGVREIGEMVGL NSEEVERALCSRTMETAKEKVVTAL NVMQAQYARDALAKNIYSRLFDWIV NRINESIKVIGIEKKKVMGVLDIYGF EILEDNSFEQFVINYCNEKLOQVFIE MTLKEEQEEYKREGIPWTKVDYFDN GIICKLIEHNQRGILAMLDEECLRPG VVSDFLAKLNQLFSKHGHYESKV TQNAQRQYDHTMGLSCFRICHYAG KVTYNVTSEFIDKNDLLFRDLLQAM WKAQHPLRLSLFPEGNPKQASLKR PTAGAQFKSSVAILMKNLYSKSPNYI RCIKPNEHQQRGQFSSDLVATQARY LGLLENVRRRAGYAHRQGYGPFLE RYRLLSRSTWPHWNGDREGVEKV LGELSMSSGELAFGKTIFIRSPKTL FYLEEQRRLRLQLATLIQIYRGWR CRTHYQLMRKSQILISSWFRGNMQ KKCYGKIKASVLLIQAFVRGWKARK NYRKYFRSEALTLADFIYKSMVQKF LLGLKNNLPSTNVLDKTPAAPYKC LSTANQELQLFYQWKCKRFRDQL SPKQVEILREKLCASELFGKKASYP QSVPIPFCDYIGLQGNPKLQKLKGG EEGPVLMAEAVKVNRRGNGKTSRI LLLTKGHVILDTKKSQAKIVIGLDN VAGVSVTSLKDGLFSLHSEMSSVG SKGDFLLVSEHVIELLTKMYRAVLDA TQRQLTVVTEKFSVRFKENSVAVK VVQGPAGGDNSKLRYKKKGSCHCLEV TVQ	None	None	None	None	None	None	None		

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