

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
Q9Y6X6	MYO16_HUMAN	Homo sapiens	Unconventional myosin-XVI	19.994096	NaN	NaN	37217939	MEIDQCLESPLGQRQRLVKRMRC EQIKAYYEREKAFQKQEGFLKRLKH AKNPKVHFNLTDMLQDAIIHHNDK EVLRLKKEGADPHTLVSSGGSLHL CARYDNAFIAEILDRGVNVNHQDE DFWTPMHIAACDNDPDIIVLLVLAG ANVLLQDVNGNIPLDYAVEGTESSSI LLTYLDENGVDLTSLRQMKLQRPMS MLTDVKHFLSSGGNVNEKNDEGVT LLHMACASGYKEVVSILEHGGDLN IVDDQYWTPLHLAAKYGQTNLVKLL LMHQANPHLVNCEEKASDIAASE FIEEMLLKAEIAWEEMKPELSASTL AQEEPYEIIIHDLPLVSSKLSPLVLP AKQDSLLEKDIMFKDATKGLCKQQS QDSIPENPMMSGSTKPEQVKLMPPA PNDDLATLSELNDGSLLYEIQKRFG NNQIYTFIGDILLVNPYKELPIYSSM VSQLYFSSGKLCSSLPPLHFCVER AFHQLFREQRPOCFILSGERGSKGS EASKQIIRHLTCRAGASRATLDSRFK HVVCILEAFGHAKTTLNDLSSCFIKY FELQFCERKQQLTGARIYTYLLEKSR LVSQPLGQSNFLIFYLMDGLSAEEK YGLHLNLCALHRYLNQTIQDDASTG ERSLNREKLAVLKRALNVVGFSSLE VENLFLVLAAILHLGDIRFTALNEGN SAFVSDLQLEQVAGMLQVSTDELA SALTTDIQYFKGDMIRRHITQIAEFF RDLLAKSLYSRLFSFLVNTMNSCLH SQDEQKSMQTLDIGILDIFGFEFQK NEFEQLCVNMTNEKMHYINEVLF LHEQVCEVQEGVTMETAYSPGNQN GVLDFFFQKPSGFLTLDEESQMIW SVESNFPKQLQSLLESNTNAVYSP MKDGNNGNVALKDHGTAFTIMHYAG RVMYDVVGAIEKNKDSLSQNLFFVM KTSENVVINHLFQSKLQSGSLVSAY PSFKFRGHKSALLSKKMTASSIIGEN KNYLELTKLKKGTSTFLQRLERL DPVTIASQLRKSMDIIGKLOKCTPH FIHCIRPNNSKLPDTFDNFYVSAQLQ YIGVLEMVKIFRYGYPVRLSFDLFS RYKPLADTFLREKKEQSAAEERCRLV QQCKLQGWQMGVRKVFYKWHAD QLNDLCLQLQRKIITCQKQVIRFLAR QHLLQRISIROQEVTSINSFLQNTED MGLKTYDALVIQNASDIARENDRLR SEMNPYHKEKLEVRNMQEEGSKR TDDKSGPRHFPSSMSVCAAVDGL GQCLVGPSPWSPSLHSVFSMDSSS LPSPRKQPPPKRDPNTRLSASYEA VSACLSAAREANEALARPSPHSD YSTMKKIPPRKPKRSPNTKLSGSYEE ISGSRPGDARPAAGAAAARVLTPTGT PQCALPPAAPPGDEDDSEPVYIEM GHAARPDSPDPGESVYEMKCCLPD DGGPGAGSFLHGCASPLLHRAPE EAAGPPGDACDIPPPFNLLPHRPPL LVFPPTVTCSPASDESPLTPLEVKK LPVLETNLKYPVQPEGSSPLSPQYSK SQKGDGRPASPLALFNGSGRASP PSTPPPPPPGPPPPAPYRCAHLAF PPEPAPVNAAGKAGPSAEAPKVPKP NSAPVAGPCSSFPKIPYSPVKATRAD ARKAGSSASPPAPYSPSSRPLSSPL DELASLFNSGRSVLRKSAAGRKIREA EGFETNMNISRDDPSTSEITSETQD RNANNHGIQLSNLSAITAENGNS ISNGLPEEDGYSRLSISGTGTSTFQR HRDSHTTQVIHQRLSENEVALQE LLDWRRKLCPEGQDWQQLHHAEP RVPPPPCKKPSLLKPEGASCNRLP SELWDTTI	None	None	None	None	None	None	None		