	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P46735	MYO1B_MOUSE	Mus musculus	Unconventional myosin-Ib	25.295579	NaN	S60	37453647;33300544	MAKMEVKSSLLDNMIGVGDMVLLE PLNEETFIDNLKKRFDHNEIYTYIGS VVISVNPYRSLPIYSPEKVEDYRNRN FYELSPHIFALSDEAYRSLRDQDKDQ CILITGESGAGKTEASKLVMSYVAAV CGKGAEVNQVKEQLLQSNPVLEAFG NAKTVRNDNSSRFGKYMDIEFDFKG DPLGGVISNYLLEKSRVVKQPRGER NFHVFYQLLSGASEELLYKLKLERD FSRYNYLSLDSAKVNGVDDAANFRT VRNAMQIVGFLDHEAEAVLEVVAAV LKLGNIEFKPESRVNGLDESKIKDKN ELKEICELTSIDQVVLERAFSFRTVE AKQEKVSTTLNVAQAYYARDALAKN LYSRLFSWLVNRINESIKAQTKVRKK VMGVLDIYGFEIFEDNSFEQFIINYC NEKLQQIFIELTLKEEQEEYIREDIE WTHIDYFNNAIICDLIENNTNGILAM LDEECLRPGTVTDETFLEKLNQVCA THQHFESRMSKCSRFLNDTTLPHS CFRIQHYAGKVLYQVEGFVDKNNDL LYRDLSQAMWKAGHSLIKSLFPEGN PAKVNLKRPPTAGSQFKASVATLMR NLQTKNPNYIRCIKPNDKKAAHIFN ESLVCHQIRYLGLLENVRVRRAGYA FRQAYEPCLERYKMLCKQTWPHWK GPARSGVEVLFNELEIPVEEHSFGRS KIFIRNPRTLFQLEDLRKQRLEDLAT LIQKIYRGWKCRTHFLLMKRSQVVIA AWYRRYAQQKRYQQIKSSALVIQSYI RGWKARKILRELKHQKRCKEAATTI AAYWHGTQARRELKRLKEEARRKH AVAVIWAYWLGLKVRREYRKFFRAN AGKKIYEFTLQRIVQKYLLEMKNKM PSLSPIDKNWPSRPYLFLDSTHKELK RIFHLWRCKKYRDQFTDQQKLIYEE KLEASELFKDKKALYPSSVGQPFQG AYLEINKNPKYKKLKDAIEEKIIIAEV VNKINRANGKSTSRIFLLTNNNLLLA DQKSGQIKSEVPLVDVTKVSMSSQN DGFFAVHLKEGSEAASKGDFLFSSD HLIEMATKLYRTTLSQTKQKLNIEIS DEFLVQFRQDKVCVKFIQGNQKNGS VPTCKRKNNRLLEVAVP	None	None	None	None	None	None	None	None	None