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Myo5_R265    1    MAPSKKAGKKGAVGGLLSGASKPQK----VQKADWSEGFTKKKAAGVPDMTLLSTITNEAINENLKARFQNQEIYTYIAHVLISVNPFRDLGIYTNDVLNSYRGKNRLEMSPHVFAIAES  116
Myo1_Sp      1    MAILKRTNRAKAATAAAPNSTGKSN---GIKKAVYTS--TRKKTVGVDDLTLLSKITDEEINKNLELRFRNGEIYTYIGHVLISVNPFRDLGIYTMDILKSYQGKNRLETSPHVYAIAEN  115
Myo5_Sc      1    MAILKRGARKKVHQE--PAKRSAN-----IKKATFDS--SKKKEVGVSDLTLLSKISDEAINENLKKRFLNATIYTYIGHVLISVNPFRDLGIYTDAVMNEYKGKNRLEVPPHVFAIAES  111
Myo5_Ca      1    MAIVKRGGRTKTKQQQVPAKSSGGGSSGGIKKAEFDI--TKKKEVGVSDLTLLSKITDEAINENLHKRFMNDTIYTYIGHVLISVNPFRDLGIYTLENLNKYKGRNRLEVPPHVFAIAES  118
myosin-le    1          MGSKG-------------------VYQYHWQS--HNVKHSGVDDMVLLSKITENSIVENLKKRYMDDYIFTYIGSVLISVNPFKQMPYFGEKEIEMYQGAAQYENPPHIYALADN   94
myosin-lf    1          MGSK----------------------ERFHWQS--HNVKQSGVDDMVLLPQITEDAIAANLRKRFMDDYIFTYIGSVLISVNPFKQMPYFTDREIDLYQGAAQYENPPHIYALTDN   92

Myo5_R265  117  AYYRMTTEKENQCVIISGESGAGKTEAAKRIMQYIAAVSGGAEGSGIEGIKEMVLATNPLLESFGCAKTLRNDNSSRHGKYLEIMFNG-MGQPVGAQITNYLLEKNRVVGQIDDERDFHI  235
Myo1_Sp    116  AYYQMKSYHENQCIIISGESGAGKTEAAKRIMQYITHVSK-SVGTEIERVSEIILATNPLLESFGCAKTLRNNNSSRHGKYLEMIFNS-GGVPVGAKITNYLLEKNRIVNQVRNERNFHI  233
Myo5_Sc    112  MYYNMKSYNENQCVIISGESGAGKTEAAKRIMQYIAAASS-THTESIGKIKDMVLATNPLLESFGCAKTLRNNNSSRHGKYLEIKFNN-QFEPCAGNITNYLLEKQRVVSQIKNERNFHI  229
Myo5_Ca    119  MYYNLKSYGENQCVIISGESGAGKTEAAKQIMQYIANVSVNQDNVEISKIKDMVLATNPLLESFGCAKTLRNNNSSRHGKYLEIKFSEGNYQPIAAHITNYLLEKQRVVSQITNERNFHI  238
myosin-le   95  MYRNMIIDRENQCVIISGESGAGKTVAAKYIMSYISRVSG--GGTKVQHVKDIILQSNPLLEAFGNAKTVRNNNSSRFGKYFEIQFSP-GGEPDGGKISNFLLEKSRVVMRNPGERSFHI  211
myosin-lf   93  MYRNMLIDCENQCVIISGESGAGKTVAAKYIMGYISKVSG--GGEKVQHVKDIILQSNPLLEAFGNAKTVRNNNSSRFGKYFEIQFSR-GGEPDGGKISNFLLEKSRVVMQNENERNFHI  209

Myo5_R265  236  FYQFTKGASAEMKEAFGLQGPEAYAYTSRSGCLDVKSVNDISDFKETLRAMQVIGLTSDEQNSIFRILATILWLGNIDFVEGDDGNAAISDSGVADFAAYLLKVDSAQLQKVLLTRIMET  355
Myo1_Sp    234  FYQFTKSAPQKYRDTYGIQGPENYVYTSACQCLSVDGISDEKDFQGTMNAMKVIGITEPEQDEIFRMLSIILWLGNIQFQEGQDGGSVISDKSITEFLGYLIGVPVAAIERALTIRIMQT  353
Myo5_Sc    230  FYQFTKGASDAYRQTFGVQKPEQYVYTAAAGCISAETIDDLQDYQETLKAMRVIGLGQEEQDQIFRMLAAILWIGNVSFIENEEGNAQVRDTSVTDFVAYLLQIDSQLLIKSLVERIMET  349
Myo5_Ca    239  FYQFTKHCPPQYQQMFGIQGPETYVYTSAAKCINVDGVDDAKDFQDTLNAMKIIGLTQQEQDNIFRMLASILWIGNISFVEDENGNAAIRDDSVTNFAAYLLDVNPEILKKAIIERTIET  358
myosin-le  212  FYQLIEGASAEQKHSLGITSMDYYYYLSLSGSYKVDDIDDRREFQETLHAMNVIGIFAEEQTLVLQIVAGILHLGNISFKEVGN-YAAVESEEFLAFPAYLLGINQDRLKEKLTSRQMDS  330
myosin-lf  210  YYQLLEGASQEQRQNLGLMTPDYYYYLNQSDTYQVDGTDDRSDFGETLSAMQVIGIPPSIQQLVLQLVAGILHLGNISFCEDGN-YARVESVDLLAFPAYLLGIDSGRLQEKLTSRKMDS  328

Myo5_R265  356  QRGGRRGSVYEVPQNVAQASSGRDALAKALYNNLFEWIVSRVNISMK-PQTVSEYVIGVLDIYGFEIFQDNSFEQLSINYVNEKLQQIFIELTLKAEQEEYVREQIKWTPIKFFDNSVVC  474
Myo1_Sp    354  QHGARRGSVYEVPLNPTQALAVRDALSMAIYNCLFDWIVERVNKALVTSDNSVSNSIGILDIYGFEIFENNSFEQLCINYVNEKLQQIFIELTLKTEQEEYVREQIAWTPIKYFNNKVVC  473
Myo5_Sc    350  NHGMKRGSVYHVPLNIVQADAVRDALAKAIYNNLFDWIVSRVNKSLQAFPG-AEKSIGILDIYGFEIFEHNSFEQICINYVNEKLQQIFIQLTLKSEQETYEREKIQWTPIKYFDNKVVC  468
Myo5_Ca    359  SHGMRRGSTYHSPLNIVQATAVRDALAKGIYNNLFEWIVERVNISLAGSQQQSSKSIGILDIYGFEIFERNSFEQICINYVNEKLQQIFIQLTLKAEQDEYVQEQIKWTPIDYFNNKVVC  478
myosin-le  331  KWGG-KSESIHVTLNVEQACYTRDALAKALHARVFDFLVDSINKAMEKDHE--EYNIGVLDIYGFEIFQKNGFEQFCINFVNEKLQQIFIELTLKAEQEEYVQEGIRWTPIEYFNNKIVC  447
myosin-lf  329  RWGG-RSESINVTLNVEQAAYTRDALAKGLYARLFDFLVEAINRAMQKPQE--EYSIGVLDIYGFEIFQKNGFEQFCINFVNEKLQQIFIELTLKAEQEEYVQEGIRWTPIQYFNNKVVC  445

Myo5_R265  475  SLIEDRRP-AGIFATLNDATATAHADPSAADNSFIQRSSML-ASNPNFEARGSKFLIKHYAGDVLYTVAGMTDKNKDTLIKDILDLIEGSKDPFL-HTLFPEKVDH---TSKKRPPTAGD  588
Myo1_Sp    474  DLIESKRP-PGLFAAMNDAIATAHADSAAADSAFAQRLNFL-SSNPHFEQRQNQFIVKHYAGDVTYSITGMTDKNKDQLATDILNLIHSSNNEFM-KSIFPVAEES---NSRRRPPTAGD  587
Myo5_Sc    469  DLIEARRP-PGIFAAMNDSVATAHADSNAADQAFAQRLNLF-TTNPHFDLRSNKFVIKHYAGDVTYDIDGITDKNKDQLQKDLVELIGTTTNTFL-ATIFPDTVDR---ESKRRPPTAGD  582
Myo5_Ca    479  DLIEATRPQPGLFAALNDSIKTAHADSEAADQVFAQRLSMVGASNRHFEDRRGKFIIKHYAGDVTYDVAGMTDKNKDAMLRDLLELVSTSQNSFINQVLFPPDLLAQLTDSRKRPETASD  598
myosin-le  448  DLIENKVNPPGIMSILDDVCATMHAVGEGADQTLLQKLQMQIGSHEHFNSWNQGFIIHHYAGKVSYDMDGFCERNRDVLFMDLIELMQSSELPFI-KSLFPENLQA---DKKGRPTTAGS  563
myosin-lf  446  DLIENKLSPPGIMSVLDDVCATMHATGGGADQTLLQKLQAAVGTHEHFNSWSAGFVIHHYAGKVSYDVSGFCERNRDVLFSDLIELMQTSEQAFL-RMLFPEKLDG---DKKGRPSTAGS  561

Myo5_R265  589  RIKLSANLLVENLMNCQPHYIRTIKPNQHRSPTEYDDKAILHQIKYLGLQENIRVRRAGFAYRAEFSKMIQRFYLLSPATSYAGDYIWTGDDRSGCERILTDAKIKREEWQMGVTKAFIK  708
Myo1_Sp    588  RIKTSANDLVETLMKCQPSYIRTIKPNQTKSPNDYDQQMVLHQIKYLGLQENIRIRRAGFAYRQAFDTFAQRFAVLSGKTSYAGEYTWQGDDKSACEQILKDTNIPSSEYQMGTSKVFIK  707
Myo5_Sc    583  KIIKSANDLVETLSKAQPSYIRTIKPNETKSPNDYDDRQVLHQIKYLGLQENVRIRRAGFAYRQVFEKFVERFYLLSPHCSYAGDYTWQGDTLDAVKYILQDSSIPQQEYQLGVTSVFIK  702
Myo5_Ca    599  KIKKSANILVDTLSQCTPSYIRTIKPNQTKKPRDYDNQQVLHQIKYLGLKENVRIRRAGFAYRSTFERFVQRFYLLSPATGYAGDYIWRGDDISAVKEILKSCHIPPSEYQLGTTKVFIK  718
myosin-le  564  KIKKQANDLVSTLMKCTPHYIRCIKPNETKKPRDWEESRVKHQVEYLGLKENIRVRRAGYAYRRIFQKFLQRYAILTKATWPS----WQGEEKQGVLHLLQSVNMDSDQFQLGRSKVFIK  679
myosin-lf  562  KIKKQANDLVATLMRCTPHYIRCIKPNETKRPRDWEENRVKHQVEYLGLKENIRVRRAGFAYRRQFAKFLQRYAILTPETWPR----WRGDERQGVQHLLRAVNMEPDQYQMGSTKVFVK  677

Myo5_R265  709  NPETLFYLEGERDRYWHTMASRIQRAWRAYVRRKHEAATKIQRFWRNQREALVYERKRDYGHQVLAGRKERRRFSLLGMRKFMGDYLDIAGGSAQGEMLRNAATISSAEQIHFSSRAELL  828
Myo1_Sp    708  NPETLFALEDMRDKFWDTMATRIQRAWRSYVRRRSEAAACIQKLWNRNKVNMELERVRNEGTKLLQGKKQRRRYSILGSRKFYGDYLSASKPNG-TLWNTCGLSQN--DHVIFSMRCEVL  824
Myo5_Sc    703  TPETLFALEHMRDRYWHNMAARIQRAWRRFLQRRIDAATKIQRTIRERKEGNKYEKLRDYGTKVLGGRKERRSMSLLGYRAFMGDYLSCNESKSKGAYIKRQVSIK--EKVIFSIHGEAL  820
Myo5_Ca    719  TPETLFALEDMRDKYWHNMAARIQRAWRRYVKRKEDAAKTIQNAWRIKKHGNQFEQFRDYGNGLLQGRKERRRMSMLGSRAFMGDYLGCNYKSGYGRFIINQVGIN--ESVIFSSKGEIL  836
myosin-le  680  APESLFLLEEMRERKYDGYARVIQKSWRKFVARK-----------------KYVQMREEASDLLLNKKERRRNSIN--RNFIGDYIGMEEHPELQQFVGKR------EKIDFADT----  769
myosin-lf  678  NPESLFLLEEVRERKFDGFARTIQKAWRRHVAVR-----------------KYEEMREEASNILLNKKERRRNSIN--RNFVGDYLGLEERPELRQFLGKR------ERVDFADS----  767

Myo5_R265  829  VSKLGRSSKLSPRFLIITDKAVYFVVSQAKDG-----RVSTTLERKIPLVTIRAISMTNLRDDFVALNVNACE--EGDPIFTCVFKTEMVTVILTLTGGNMSVNIGPTIDYAKKK-----  936
Myo1_Sp    825  VHKLGRTSKPSPRQLVLTKKNLYLVITKIVDQ-----KLTQQVEKKFAVSSIDSVGLTNLQDDWVAIRNKSSQ--NGDMFLRCFFKTEFITTLKRIN-RNIQVIVGPTIQYCRKP-----  931
Myo5_Sc    821  HTKFGRSAQRLKKTFLLTPTTLYIVGQTLVQN-----AMTYTQDYKIDVRNIQAVSLTNLQDDWVAIKLASSG--QPDPLINTYFKTELITHLKRLN-DKIQIKIGSAIEYQKKP-----  927
Myo5_Ca    837  LSKFGRSSKRLPRIFIVTKTSIYIIAEVLVEK-----RLQLQKEFTIPISGINYLGLSTFQDNWVAISLHSPTPTTPDVFINLDFKTELVAQLKKLN-PGITIKIGPTIEYQKKP-----  945
myosin-le  770  VTKYDRRFKGVKRDLLLTPKCLYLIGREKVKQGPDKGLVKEVLKRKIEIERILSVSLSTMQDDIFILHEQEYDS-LLESVFKTEFLSLLAKRYEEKTQKQLPLKFSNTLELKLKKENWGP  888
myosin-lf  768  VTKYDRRFKPIKRDLILTPKCVYVIGREKVKKGPEKGQVCEVLKKKVDIQALRGVSLSTRQDDFFILQEDAADS-FLESVFKTEFVSLLCKRFEEATRRPLPLTFSDTLQFRVKKEGWG-  885

Myo5_R265  937  ---DKRAVIKAQKNEA---VRGDATYKSHTIQVGSGEPPSSLSNPMPPRKPKVKKAAKTVSSSRPINSGRPAAVALPGATK-----PAAPAALSGMPSYTPVVAKPIATPAATTGAARAP  1045
Myo1_Sp    932  ---GKVQTVKTAKDET---TKDYDYYKSGTIHVGTGLPPTSKS----KPFPRLATGGSTAAARGPRPVVQNKPAATKPVSM-----PAAKSKPAPMANPVSTAQQTQNRPPAPAMQARPN  1036
Myo5_Sc    928  ---GKLHSVKCQINES--APKYGDIYKSSTISVRRGNPPNSQV----HKKPRKKSSIS--SGYH-------ASSSQATRR-----PVSIAAAQHVPTAPASRHSKKPAPPPGMQNKAA  1023
Myo5_Ca    946  ---GKFHTVKFIIGAGPEIPNNGDHYKSGTVSVKQGLPASSKN----PKRPRGVSSKVDYSKYYNRGAARKTAAAAQATPRYNQPTPVANSGYSAQPAYPIPQQPQQYQPQQSQQQTPYP  1058
myosin-le  889  WSAGGSRQVQFHQGFG-DLAVLKPSNKVLQVSIGPGLPKNSRP-----TRRNTTQNTGYSSGTQNANYPVRAAPPPP-GYHQNGV---IRNQYVPYPHAP--GSQRSNQKSLYTSMARPP  996
myosin-lf  886  --GGGTRSVTFSRGFG-DLAVLKVGGRTLTVSVGDGLPKSSKP-----TRKGMAKGKPRRS----SQAPTRAAPAPPRGMDRNGVPPSARGGPLPLEIMSGGGTHRPPRGPPSTSLGASR  993

Myo5_R265  1046  PSIPG---RAGAPPPPPPPPPAGPPKEIYKALYNFTGQEGEMNLVKGEEVEVKEKDDNGWWMVVKNGQEGWAPSNYLKKVEHAPP------PPPPPPASRPMPARPPAASSAPAAPDVT  1156
Myo1_Sp    1037  TT----------QAAAPVTSTTTTIKQATTVSASKPAPSTVTSAASSPSNISKPSAPVANN----------VSKPSAVPP------PPPPPPAEVEK---KDLYLALYDFAGR-  1120
Myo5_Sc    1024  TR----------RSVPNPASTLTASQSNARPSPPTAATRATPAATPAAAAMGSG----------RQANIPP-----PPPPPPPSSKPK---EPMFEAAYDFPGSG  1100
Myo5_Ca    1059  TQSSIPSVNQNQSRQPQRKVPPPAPSLQVSAAQAALGKSPTQQRQTPAHNPVASPNRPASTTIATTTSH----TSRPVKKTAPAPPVKKTAPPPPPPTLVKPK---FPTYKAMFDYDGSV  1171
myosin-le  997  LP----------RQQSTSSDRVSQTPESLDFLKVPDQGAAGVRRQTTS----------RPPPAGG------RPKPQPKPKPQ---VPQCKALYAYDAQ-  1065
myosin-lf  994  RP----------RARPPSE-------HNTEFLNVPDQGMAGMQRKRSV----------GQRPVPG------VGRPKPQPRTH---GPRCRALYQYVGQ-  1055

Myo5_R265  1157  NGSAVPSWKAKNTASGTPSAASTPPTSRPASSASKVPPVIKAKPAIPAKPQVGAKPALAIG---------GKPPVPTAPKIQPKAASKPGQVAQPAKAPGQLDLAAAFAKRAAR------  1261
Myo1_Sp    1121  SPNEMTIKKDEIIEIVQKEPSGWWLA--LKNGAEGWVPATYVTEYKGS----------TPQTTASSTNVAAQANNNASPAEVN-NLAGSLADALRMRAS-------  1206
Myo5_Sc    1101  SPSELPLKKGDVIYITREEPSGWSLGKLLDGSKEGWVPTAYMKPHSGN---NNIPTPPQ----------NRDVPKPVLNSVQHDN-TSANVIPAAAQASLGDGLANALAARAN-------  1199
Myo5_Ca    1172  AGS-IPLVKDTVYYVTQVN-GKWGLVKTMDETKEGWSPIDYLKECSPNETQKSAPPPPPPPAATASAGANGASNPISTTTSTNTTTSSHTTNATSNGSLGNGLADALKAKKQEETTLAG  1289
myosin-le  1066  DTDELSFNANDIIDIIKEDPSGWWTG--RLRGKQGLFPNNYVTKI----------  1108
myosin-lf  1056  DVDELSFNVNEVIEILMEDPSGWWKG--RLHGQEGLFPGNYVEKI----------  1098

Myo5_R265  1262  ----------AQQEED  1267
Myo1_Sp    1207  ----------AVRGSD-----EEEDW  1217
Myo5_Sc    1200  -------KMRLESDDEEANEDEEEDDW  1219
Myo5_Ca    1290  SLADALKKRQGATRDSDDEEEEDDDDW  1316
myosin-le  1109  1108
myosin-lf  1099  1098
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