

CLUSTAL O(1.2.4) multiple sequence alignment

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B7I325|MURD_ACIB5      MLIQRGGLKVVAGLGISGVS AVNFLHEQGYQVAVTDSRPTPPGHDQIPAGVKTSFGQLDQ      60
A3M1C6|MURD_ACIBT      MLIQRGGLKVVAGLGISGVS AVNFLHEQGYQVAVTDSRPTPPGHDQIPAGVKTSFGQLDQ      60
B2I1V6|MURD_ACIBC      MLIQRGGLKVVAGLGISGVS AVNFLHEQGYQVAVTDSRPTPPGHDQIPAGVKTSFGQLDQ      60
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B7I325|MURD_ACIB5      ELLLQAEI IILSPGLAPQLPEIQAAIAKGISVVGDIQLLRRATDVPIVAITGSNAKSTVT      120
A3M1C6|MURD_ACIBT      ELLLQAEI IILSPGLAPQLPEIQAAIAKGISVVGDIQLLRRATDVPIVAITGSNAKSTVT      120
B2I1V6|MURD_ACIBC      ELLLQAEI IILSPGLAPQLPEIQAAIAKGISVVGDIQLLRRATDVPIVAITGSNAKSTVT      120
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B7I325|MURD_ACIB5      TLIGLMAKDAGKKVAVGGNLRPALDLLKDKPELLVLELSSFQLETTSHLNAEVAVVLNM      180
A3M1C6|MURD_ACIBT      TLIGLMAKDAGKKVAVGGNLRPALDLLKDKPELLVLELSSFQLETTSHLNAEVAVVLNM      180
B2I1V6|MURD_ACIBC      TLIGLMAKDAGKKVAVGGNLRPALDLLKDKPELLVLELSSFQLETTSHLNAEVAVVLNM      180
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B7I325|MURD_ACIB5      SEDHLDRHGNMLGYHQAKHRIFQGAKKVVFNRRDALSRLVPDTPMQSFGLNAPDLNQY      240
A3M1C6|MURD_ACIBT      SEDHLDRHGNMLGYHQAKHRIFQGAKKVVFNRRDALSRLVPDTPMQSFGLNAPDLNQY      240
B2I1V6|MURD_ACIBC      SEDHLDRHGNMLGYHQAKHRIFQGAKKVVFNRRDALSRLVPDTPMQSFGLNAPDLNQY      240
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B7I325|MURD_ACIB5      GVLRDADGTLWLARGLQRLIKSSDLYIQGMHNVANALACLALGEAIGLPMESMLETLKQF      300
A3M1C6|MURD_ACIBT      GVLRDADGTLWLACGLQRLIKSSDLYIQGMHNVANALACLALGEAIGLPMESMLETLKQF      300
B2I1V6|MURD_ACIBC      GVLRDADGTLWLARGLQRLIKSSDLYIQGMHNVANALACLALGEAIGLPMESMLETLKRF      300
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B7I325|MURD_ACIB5      KGLEHRCEYVKTVHDVRYYNDSKGTNVGATLAAIDGLGAAIEVKKGKVALILGGQKGQD      360
A3M1C6|MURD_ACIBT      KGLEHRCEYVKTVHDVRYYNDSKGTNVGATLAAIDGLGAAIEVKKGKVALILGGQKGQD      360
B2I1V6|MURD_ACIBC      KGLEHRCEYVKTVRDVRYYNDSKGTNVGATLAAIDGLGAAIEVKKGKVALILGGQKGQD      360
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B7I325|MURD_ACIB5      FGPLRSSIEKYAKVVVLIGEDAPVIEQAIQGATKILHAATLKEAVELCQRETQAEDVLL      420
A3M1C6|MURD_ACIBT      FSPLRSSIEKYAKVVVLIGEDAPVIEQAIQGATKILHAATLKEAVELCQRETQAEDVLL      420
B2I1V6|MURD_ACIBC      FSPLRSSIEKYAKVVVLIGEDAPVIEQAIQGATKILHAATLKEAVELCQRETQAEDVLL      420
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B7I325|MURD_ACIB5      SPACASFDMFKSYNDRGQQFVACVNSLV      448
A3M1C6|MURD_ACIBT      SPACASFDMFKSYNDRGQQFVACVNSLV      448
B2I1V6|MURD_ACIBC      SPACASFDMFKSYNDRGQQFVACVNSLV      448
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