

CLUSTAL O(1.2.4) multiple sequence alignment

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G2JLK8 | G2JLK8_ACIBA      MFKNRFFNTMQIQNQVQLKPFNTLSLDVTASHYTKVKSIEDI EEALAFAKEHELNVLVLS      60
B0VRK7 | MURB_ACIBS      MFKNRFFNTMQIQNQVQLKPFNTLSLDVTASHYTKVKSIEDI EEALAFAKEHELNVLVLS      60
B0V744 | MURB_ACIBY      MFKNRFFNTMQIQNQVQLKPFNTLSLDVTASHYTKVKSIEDI EEALAFAKEHELNVLVLS      60
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G2JLK8 | G2JLK8_ACIBA      GGSNMLLPQQINALVIHLDIQGI DVLSEDQDFIRVKVGAGQVWHDFVLYSTKQNWFG LQN      120
B0VRK7 | MURB_ACIBS      GGSNMLLPQQINALVIHLDIQGI DVLSEDQDFIRVKVGAGQVWHDFVLYTTKQNWFG LQN      120
B0V744 | MURB_ACIBY      GGSNMLLPQQINALVIHLDIQGI DVLSEDQDFIRVKVGAGQVWHDFVLYTTKQNWFG LQN      120
*****;*****

G2JLK8 | G2JLK8_ACIBA      LALIPGLVGASPVQNIGAYGVEVGEFIESVQVYDRLLKQTGSISAADCHFSYRHSIFKDD      180
B0VRK7 | MURB_ACIBS      LALIPGLVGASPVQNIGAYGVEVGEFIESVQVYDRLLKQTGSISAADCHFSYRHSIFKDE      180
B0V744 | MURB_ACIBY      LALIPGLVGASPVQNIGAYGVEVGEFIESVQVYDRLLKQTGSISAADCHFSYRHSIFKDD      180
*****;

G2JLK8 | G2JLK8_ACIBA      PARYIITHVTFKLLKQANLKLNYGDLKQAVGDNLTAENLQNQVIHIRQSKLPDPKEYPNV      240
B0VRK7 | MURB_ACIBS      PARYIITHVTFKLLKQANLKLNYGDLKQAVGDNLTAENLQNQVIHIRQSKLPDPKEYPNV      240
B0V744 | MURB_ACIBY      PARYIITHVTFKLLKQANLKLNYGDLKQAVGDNLTAENLQNQVIHIRQSKLPDPKEYPNV      240
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G2JLK8 | G2JLK8_ACIBA      GSFFKNPIVNTQEFERLIAQFSTIPHYPQANGNVKIAAGWLIDQAGWKGKQLGVVGMFHK      300
B0VRK7 | MURB_ACIBS      GSFFKNPIVNTQEFERLIAQFSTIPHYPQANGNVKIAAGWLIDQAGWKGKQLGVVGMFHK      300
B0V744 | MURB_ACIBY      GSFFKNPIVNTQEFERLIAQFSTIPHYPQANGNVKIAAGWLIDQAGWKGKQLGVVGMFHK      300
*****

G2JLK8 | G2JLK8_ACIBA      QALVLVNYANASLADVKKTYQAVQHDVEQRFQIMLEPEPVL YNNLGLIENHTE      353
B0VRK7 | MURB_ACIBS      QALVLVNYANASLADVKKTYQAVQHDVEQRFQIMLEPEPVL YNNLGLIENHTE      353
B0V744 | MURB_ACIBY      QALVLVNYANASLADVKKTYQAVQHDVEQRFQIMLEPEPVL YNNLGLIENHTE      353
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