

CLUSTAL O(1.2.4) multiple sequence alignment

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Q830P3 | MURB_ENTFA      MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
R3KRK1 | R3KRK1_ENTFL    MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
A0A0M2A9A4 | A0A0M2A9A4_ENTFL MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
A0A0M2AQJ4 | A0A0M2AQJ4_ENTFL MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
C7WZB7 | C7WZB7_ENTFL    MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
A0A0M2CJ94 | A0A0M2CJ94_ENTFL MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
C7D0R3 | C7D0R3_ENTFL    MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
C7VBG0 | C7VBG0_ENTFL    MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
V7ZMX2 | V7ZMX2_ENTFL    MNTKAMLETLNEITLLVDEPLKNVFTFKTGGPADVLALPKTKKEVEEIVAYCREQGLSWL 60
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Q830P3 | MURB_ENTFA      VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
R3KRK1 | R3KRK1_ENTFL    VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
A0A0M2A9A4 | A0A0M2A9A4_ENTFL VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
A0A0M2AQJ4 | A0A0M2AQJ4_ENTFL VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
C7WZB7 | C7WZB7_ENTFL    VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
A0A0M2CJ94 | A0A0M2CJ94_ENTFL VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
C7D0R3 | C7D0R3_ENTFL    VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
C7VBG0 | C7VBG0_ENTFL    VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
V7ZMX2 | V7ZMX2_ENTFL    VLGNASNLIVRDGGIRDVVIMLTEMKEIKVAGTTMIVDAGAKLIDTTYEALAADLTGFEEF 120
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Q830P3 | MURB_ENTFA      ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
R3KRK1 | R3KRK1_ENTFL    ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
A0A0M2A9A4 | A0A0M2A9A4_ENTFL ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
A0A0M2AQJ4 | A0A0M2AQJ4_ENTFL ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
C7WZB7 | C7WZB7_ENTFL    ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
A0A0M2CJ94 | A0A0M2CJ94_ENTFL ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
C7D0R3 | C7D0R3_ENTFL    ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
C7VBG0 | C7VBG0_ENTFL    ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
V7ZMX2 | V7ZMX2_ENTFL    ACGIPGSVGGAVYMNAGAYGGEIKDVFQSAEVLADGTIQTMTKEDLNFRYRHSEIQELH 180
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Q830P3 | MURB_ENTFA      CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
R3KRK1 | R3KRK1_ENTFL    CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
A0A0M2A9A4 | A0A0M2A9A4_ENTFL CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
A0A0M2AQJ4 | A0A0M2AQJ4_ENTFL CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
C7WZB7 | C7WZB7_ENTFL    CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
A0A0M2CJ94 | A0A0M2CJ94_ENTFL CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
C7D0R3 | C7D0R3_ENTFL    CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
C7VBG0 | C7VBG0_ENTFL    CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
V7ZMX2 | V7ZMX2_ENTFL    CIVLQATFALEKGNHAEIKAQMDELTELRELKQPLEYPCSGSVFKRPVGHFTGKLIQDAG 240
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Q830P3 | MURB_ENTFA      LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
R3KRK1 | R3KRK1_ENTFL    LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
A0A0M2A9A4 | A0A0M2A9A4_ENTFL LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
A0A0M2AQJ4 | A0A0M2AQJ4_ENTFL LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
C7WZB7 | C7WZB7_ENTFL    LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
A0A0M2CJ94 | A0A0M2CJ94_ENTFL LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
C7D0R3 | C7D0R3_ENTFL    LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
C7VBG0 | C7VBG0_ENTFL    LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
V7ZMX2 | V7ZMX2_ENTFL    LQGLKWGGAQISEKHAGFIVNIDHATATDYVELIAHIQEVIKEKFDVDELQTEVRIIGEEV 300
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