

CLUSTAL O(1.2.4) multiple sequence alignment

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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      MLEPLRLSQLTVALDARLIGDDAVFSAVSTDSRAIGPGQLFIALSGPRFDGHDYLAEVAA 60
W1MPY5 | W1MPY5_PSEAI              MLEPLRLSQLTVALDARLIGEDAVFSAVSTDSRAIGPGQLFIALSGPRFDGHDYLSAEVAA 60
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     MLEPLRLSQLTVALDARLIGEDAVFSAVSTDSRAIGPGQLFIALSGPRFDGHDYLAEVAA 60
Q9HVZ7 | Q9HVZ7_PSEAE              MLEPLRLSQLTVALDARLIGEDAVFSAVSTDSRAIGPGQLFIALSGPRFDGHDYLAEVAA 60
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      KGAVAALVEREVADAPLPQLLVRDTRAALGRLGALNRRKFTGPLAAMTGSSGKTTVKEML 120
W1MPY5 | W1MPY5_PSEAI              KGAVAALVEREVADAPLPQLLVRDTRAALGRLGALNRRKFTGPLAAMTGSSGKTTVKEML 120
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     KGAVAALVEREVADAPLPQLLVRDTRAALGRLGALNRRKFTGPLAAMTGSSGKTTVKEML 120
Q9HVZ7 | Q9HVZ7_PSEAE              KGAVAALVEREVADAPLPQLLVRDTRAALGRLGALNRRKFTGPLAAMTGSSGKTTVKEML 120
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      ASILRTQAGDAESVLATRGNLNNDLGVPLTLLQLAPQHRSAVIELGASRIGEIAITVELT 180
W1MPY5 | W1MPY5_PSEAI              ASILRTQAGDAESVLATRGNLNNDLGVPLTLLQLAPQHRSAVIELGASRIGEIAITVELT 180
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     ASILCTQAGDAESVLATRGNLNNDLGVPLTLLQLAPQHRSAVIELGASRIGEIAITVELT 180
Q9HVZ7 | Q9HVZ7_PSEAE              ASILRTQAGDAESVLATRGNLNNDLGVPLTLLQLAPQHRSAVIELGASRIGEIAITVELT 180
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      RPHVAIITNAGTAHVGEFVGPEKIVEAKGEILEGLAADGTAVLNLDKAFDTWKARASGR 240
W1MPY5 | W1MPY5_PSEAI              RPHVAIITNAGTAHVGEFVGPEKIVEAKGEILEGLAADGTAVLNLDKAFDTWKARASGR 240
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     RPHVAIITNAGTAHVGEFVGPEKIVEAKGEILEGLAADGTAVLNLDKAFDTWKARASGR 240
Q9HVZ7 | Q9HVZ7_PSEAE              RPHVAIITNAGTAHVGEFVGPEKIVEAKGEILEGLAADGTAVLNLDKAFDTWKARASGR 240
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      PLLTFSLDRPQADFRAADLQRDARGCMGFRLQGVAEGAQVQLNLLGRHNVANALAAAAAA 300
W1MPY5 | W1MPY5_PSEAI              PLLTFSLDRPQADFRAADLQRDARGCMGFRLQGVAEGAQVQLNLLGRHNVANALAAAAAA 300
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     PLLTFSLDRPQADFRAADLQRDARGCMGFRLQGVAEGAQVQLNLLGRHNVANALAAAAAA 300
Q9HVZ7 | Q9HVZ7_PSEAE              PLLTFSLDRPQADFRAADLQRDARGCMGFRLQGVAEGAQVQLNLLGRHNVANALAAAAAA 300
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      HALGVPLSGIVAGLQALQPVKGRAVAQLTASGLRVIDDSYNANPASMLAIDILSGFSGR 360
W1MPY5 | W1MPY5_PSEAI              HALGVPLDGIIVAGLQALQPVKGRAVAQLTASGLRVIDDSYNANPASMLAIDILSGFSGR 360
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     HALGVPLDGIIVAGLQALQPVKGRAVAQLTASGLRVIDDSYNANPASMLAIDILSGFSGR 360
Q9HVZ7 | Q9HVZ7_PSEAE              HALGVPLDGIIVAGLQALQPVKGRAVAQLTASGLRVIDDSYNANPASMLAIDILSGFSGR 360
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      TVLVLGDMGELGSWAEQAHREVGAYAAGKVSALYAVGPLMAHAVQAFGATGRHFADQASL 420
W1MPY5 | W1MPY5_PSEAI              TVLVLGDMGELGSWAEQAHREVGAYAAGKVSALYAVGPLMAHAVQAFGATGRHFADQASL 420
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     TVLVLGDMGELGSWAEQAHREVGAYAAGKVSALYAVGPLMAHAVQAFGATGRHFADQASL 420
Q9HVZ7 | Q9HVZ7_PSEAE              TVLVLGDMGELGSWAEQAHREVGAYAAGKVSALYAVGPLMAHAVQAFGATGRHFADQASL 420
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A0A1C7BHE4 | A0A1C7BHE4_PSEAI      IGALATEQPTTILIKGSRSAAMDKVVAALCGSSEESH 458
W1MPY5 | W1MPY5_PSEAI              IGALATEQPTTILIKGSRSAAMDKVVAALCGSSEESH 458
A0A0H3R1R0 | A0A0H3R1R0_PSEAI     IGALATEQPTTILIKGSRSAAMDKVVAALCGSSEESH 458
Q9HVZ7 | Q9HVZ7_PSEAE              IGALATEQPTTILIKGSRSAAMDKVVAALCGSSEESH 458
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