

CLUSTAL O(1.2.4) multiple sequence alignment

A0A0M2ADL2 | A0A0M2ADL2_ENTFL MKLTFWEVAQAVEASNDWQQWDFPLTGIEFDSRKIVKGNLFVPLQGENDGHRFIESAMA 60
R3JFX3 | R3JFX3_ENTFL MKLTFWEVAQAVEASNDWQQWDFPLTGIEFDSRKIVKGNLFVPLQGENDGHRFIESAMA 60
C7CY18 | C7CY18_ENTFL MKLTFWEVAQAVEASNDWQQWDFPLTGIEFDSRKIVKGNLFVPLQGENDGHRFIESAMA 60
V7ZJZ7 | V7ZJZ7_ENTFL MKLTFWEVAQAVEASNDWQQWDFPLTGIEFDSRKIVKGNLFVPLQGENDGHRFIESAMA 60
Q837J3 | Q837J3_ENTFA MKLTFWEVAQAVEASNDWQQWDFPLTGIEFDSRKIVKGNLFVPLQGENDGHRFIESAMA 60

A0A0M2ADL2 | A0A0M2ADL2_ENTFL NGCQAAFQWQDLVDAPQQPLVPLHVTDPVAMQKLATYYLNKMQPNVIAVTGSNGKTTTKD 120
R3JFX3 | R3JFX3_ENTFL NGCQAAFQWQDLVDAPQQPLVPLHVTDPVAMQKLATYYLNKMQPNVIAVTGSNGKTTTKD 120
C7CY18 | C7CY18_ENTFL NGCQAAFQWQDLVDAPQQPLVPLHVTDPVAMQKLATYYLNKMQPNVIAVTGSNGKTTTKD 120
V7ZJZ7 | V7ZJZ7_ENTFL NGCQAAFQWQDLVDAPQQPLVPLHVTDPVAMQKLATYYLNKMQPNVIAVTGSNGKTTTKD 120
Q837J3 | Q837J3_ENTFA NGCQAAFQWQDLVDAPQQPLVPLHVTDPVAMQKLATYYLNKMQPNVIAVTGSNGKTTTKD 120

A0A0M2ADL2 | A0A0M2ADL2_ENTFL LIAAVLSEKFTVYKIQGNYNNQIGLPYTIHMPDETEKLILEMGMDHAQEISFSLNLAQP 180
R3JFX3 | R3JFX3_ENTFL LIAAVLSEKFTVYKIQGNYNNQIGLPYTIHMPDETEKLILEMGMDHAQEISFSLNLAQP 180
C7CY18 | C7CY18_ENTFL LIAAVLSEKFTVYKIQGNYNNQIGLPYTIHMPDETEKLILEMGMDHAQEISFSLNLAQP 180
V7ZJZ7 | V7ZJZ7_ENTFL LIAAVLSEKFTVYKIQGNYNNQIGLPYTIHMPDETEKLILEMGMDHAQEISFSLNLAQP 180
Q837J3 | Q837J3_ENTFA LIAAVLSEKFTVYKIQGNYNNQIGLPYTIHMPDETEKLILEMGMDHAQEISFSLNLAQP 180

A0A0M2ADL2 | A0A0M2ADL2_ENTFL EVAAITMIGEAHVENLGSRAGIAQAKMEIVDGLAKDGSFLVPSDEPLLEPLVEKVTQQVI 240
R3JFX3 | R3JFX3_ENTFL EVAAITMIGEAHVENLGSRAGIAQAKMEIVDGLAKDGSFLVPSDEPLLEPLVEKVTQQVI 240
C7CY18 | C7CY18_ENTFL EVAAITMIGEAHVENLGSRAGIAQAKMEIVDGLAKDGSFLVPSDEPLLEPLVEKVTQQVI 240
V7ZJZ7 | V7ZJZ7_ENTFL EVAAITMIGEAHVENLGSRAGIAQAKMEIVDGLAKDGSFLVPSDEPLLEPLVEKVTQQVI 240
Q837J3 | Q837J3_ENTFA EVAAITMIGEAHVENLGSRAGIAQAKMEIVDGLAKDGSFLVPSDEPLLEPLVEKVTQQVI 240

A0A0M2ADL2 | A0A0M2ADL2_ENTFL TFGFSQESQLQGIVTNEAKEQTSFKISDTQMTFTIPVPGTYNVTNALIAIGIGRYFQLTD 300
R3JFX3 | R3JFX3_ENTFL TFGFSQESQLQGIVTNEAKEQTSFKISDTQMTFTIPVPGTYNVTNALIAIGIGRYFQLTD 300
C7CY18 | C7CY18_ENTFL TFGFSQESQLQGIVTNEAKEQTSFKISDTQMTFTIPVPGTYNVTNALIAIGIGRYFQLTD 300
V7ZJZ7 | V7ZJZ7_ENTFL TFGFSQESQLQGIVTNEAKEQTSFKISDTQMTFTIPVPGTYNVTNALIAIGIGRYFQLTD 300
Q837J3 | Q837J3_ENTFA TFGFSQESQLQGIVTNEAKEQTSFKISDTQMTFTIPVPGTYNVTNALIAIGIGRYFQLTD 300

A0A0M2ADL2 | A0A0M2ADL2_ENTFL SEIQKGLATAELTKNTEWLKAANGAEILSDVYNANPTAMGLVLDSFSKMSKVGKRIAVL 360
R3JFX3 | R3JFX3_ENTFL SEIQKGLATAELTKNTEWLKAANGAEILSDVYNANPTAMGLVLDSFSKMSKVGKRIAVL 360
C7CY18 | C7CY18_ENTFL SEIQKGLATAELTKNTEWLKAANGAEILSDVYNANPTAMGLVLDSFSKMSKVGKRIAVL 360
V7ZJZ7 | V7ZJZ7_ENTFL SEIQKGLATAELTKNTEWLKAANGAEILSDVYNANPTAMGLVLDSFSKMSKVGKRIAVL 360
Q837J3 | Q837J3_ENTFA SEIQKGLATAELTKNTEWLKAANGAEILSDVYNANPTAMGLVLDSFSKMSKVGKRIAVL 360

A0A0M2ADL2 | A0A0M2ADL2_ENTFL GDMLELGPDSAAHQAMAQHLAPDAIEEVFLYGSEMAYLADKLQETYPANIHSFKKEEK 420
R3JFX3 | R3JFX3_ENTFL GDMLELGPDSAAHQAMAQHLAPDAIEEVFLYGSEMAYLADKLQETYPANIHSFKKEEK 420
C7CY18 | C7CY18_ENTFL GDMLELGPDSAAHQAMAQHLAPDAIEEVFLYGSEMAYLADKLQETYPANIHSFKKEEK 420
V7ZJZ7 | V7ZJZ7_ENTFL GDMLELGPDSAAHQAMAQHLAPDAIEEVFLYGSEMAYLADKLQETYPANIHSFKKEEK 420
Q837J3 | Q837J3_ENTFA GDMLELGPDSAAHQAMAQHLAPDAIEEVFLYGSEMAYLADKLQETYPANIHSFKKEEK 420
*****:*** *****:*****

A0A0M2ADL2 | A0A0M2ADL2_ENTFL NELIEAVKKVLQPEDMVVLKGSNGMGLREVIDALLEK 457
R3JFX3 | R3JFX3_ENTFL NELIEAVKKVLQPEDMVVLKGSNGMGLREVIDALLEK 457
C7CY18 | C7CY18_ENTFL NELIEAVKKVLQPEDMVVLKGSNGMGLREVIDALLEK 457
V7ZJZ7 | V7ZJZ7_ENTFL NELIEAVKKVLQPEDMVVLKGSNGMGLREVIDALLEK 457
Q837J3 | Q837J3_ENTFA NELIEAVKKVLQPEDMVVLKGSNGMGLREVIDALLEK 457
