

logo

NTDB id 327500 EG882 RS18735 WP 015239537.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

80
80

logo

NTDB id 327500 EG882 RS18735 WP 015239537.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

160
160

logo

NTDB id 327500 EG882 RS18735 WP 015239537.1
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consensus

240
240

logo

NTDB id 327500 EG882 RS18735 WP 015239537.1
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consensus

SYEREPHELELFRLMTGSTYYRLYQTAKELNLDITYKELSGTKRHKTPELAHLAESQYEVRRPAIPYTGKQEAFTVMQAQNR
 AQAQADELDLFRMTGKTYRLLHKAKELNLDISCKELHGTGRHQHAPELACIESQYDVRPAAAYTGGQEAFTVMQAQNR
 SAEQAPDELDLFRMTGSTYYKLYQTAKELNADISCKELHGTGRHQHAPELACIESQYDVRPAAAYTGGQEAFTVMQAQNR
 SYEREPHELELFRLMTGKTYRLLHKAKELNLDITYKELSGTKRHTKTPELAHLEAQYEVRRPAIPYAEKQEAFTVMQAQNR
 *

320
320

logo

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RAELEGIAREI^{QS}SLVR^{EG}GGYRYKDMAIL^{LA}LRQPEDYKDM^{LL}KEVFADY^{GL}ET^{ET}PFYFDGKASML^{QH}NHPLIEFIRSSLDV^{ML}KGNWRY

RAELEGIAREI^{QS}SLVR^{EG}GGYRYKDMAIL^{LA}LRQPEDYKDM^{LL}KEVFADY^{GL}ET^{ET}PFYFDGKASML^{QH}NHPLIEFIRSSLDV^{ML}KGNWRY

RAELEGIAREI^{QS}SLVR^{EG}GGYRYKDMAIL^{LA}LRQPEDYKDM^{LL}KEVFADY^{GL}ET^{ET}PFYFDGKASML^{QH}NHPLIEFIRSSLDV^{ML}KGNWRY

400
400

logo

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EAVFRCAKTELLFPLDQPEKQYREQVDQLENYCIAYGIGKERWTSGERFQYRRFVSLDDFAQTDQEIEMENMLNETKEW
 EAVFRCAKTELLFPLDQPEKQYREQVDQLENYCIAYGIGKERWTSGERFQYRRFVSLDDFAQTDQEIEMENMLNETKEW
 EAVFRCAKTELLFPLDQPEKQYREQVDQLENYCIAYGIGKERWTSGERFQYRRFVSLDDFAQTDQEIEMENMLNETKEW

480
480

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560
560

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640
640

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720
720

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800
800

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880
880

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consensus



960
960

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1040
1040

logo

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consensus



1120
1120

logo



- ☐ non conserved
- ☒ similar
- ☒ ≥ 50% conserved