

logo

NTDB id 411126 GRT15 RS05390 WP 159164900.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

80
80

logo

NTDB id 411126 GRT15 RS05390 WP 159164900.1
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consensus

160
160

logo

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KLHDL_GILYQQMER_RSLA_GCHYLHSE_DYLTLLA_QQ_LPLA_EDDV_KGAR_VYVDG_FYQ_FT_PQ_ELRVLEQL_MHA_EHYTF_SLTAD_SP_K
 KLHDL_GILYQQMER_RSLA_GCHYLHSE_DYLTLLA_QQ_LPLA_EDDV_KGAR_VYVDG_FYQ_FT_PQ_ELRVLEQL_MHA_EHYTF_SLTAD_SP_K
 KLHDL_SILYQQME_KSLA_DCHYLHSE_DYLTLLA_EHL_IPLA_EDDV_KGAR_VYVDG_FYQ_FT_PQ_ELRVLEQL_MHA_EHYTF_SLTAD_KP

240
240

logo

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SYEREPHLELFRMTGKSTYRQLYQAKELNLDITYKELSGTKRRHTKVPELAHLAESQVVRPAIPYTGGKEAFATVMQAQNR
SAEQAPDELDLFRMTGSYYKLTYQTAKELNADISCKELHGTKRRRHVPCLACIESQYDVRPAAAYTGGKEAFATVMQAQNR
SYEREPHELELFRMTGKYRLLHQAKELNLDTIKELSGTERHTKTPELAHLEAQYEARPAIPYAEEKQEALATVMQAANR

320
320

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RAELEGIAREIQSLVRDGGYRYKDMAILLRQPEDYKDLLEKVFADYGLTPYFIDGKASMLQHPLIEFIRSSLDVVKGNWRY
RAELEGIAREIHALVREKGYRYKDVAILLRQPEDYKDMVLEKVFADYETLPYFIDGKASMLNHPLIEFIRSSLDVLKGNWRY

400
400

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EAVFRCAKTELLFPLNQDEKQKQYREQVDQLENYCIAYGIGKERWNTNGERFQYRRFVSLDQDFAQTDQEIEMENMLNETREW
 EAVFRCAKTELLFPLDQPEQKQYREQVDQLENYCIAYGIGKERWNTNGERFQYRRFVSLDQDFAQTDQEIEMEHMLNETREW
 EAVFRCAKTELLFPLNEPKAKQYREQVDQLENYCIAYGIGKERWNTNGERFQYRRFVSLDQDFAQTDQEIEMENMLNETRDW

480
480

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[illegible]560
560

logo

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640
640

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720
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800
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880
880

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960
960

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consensus



1040
1040

logo

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consensus



1120
1120

logo

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consensus

CTYCAFKSVCQFDESLEENEYRSLKAEKDGTTLEWIKKEADGNEHS

CTYCAFKSVCQFDESLEENEYRSLKAEKDGTTLEWIKKEADDDANS

!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!*!*!!!!!!*!

1166

1166

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved