

logo

NTDB id 391671 F9277 RS04260 WP 005444697.1
NTDB id 1157 A1552VC RS03895 WP 001888250.1
consensus

[illegible]80
66

logo

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consensus

PDVP|YFMTAHGS|DTAVEAMRHG^SQDFL|KPCEADRLRVTVNNAIRKA^SKLKN^{EV}DN^KKNQYQGF|GSSQTMQ^QVYRT
 PDVPI^IFMTAHGSIDTAVEAMRHG^SQDFLIKPC^EADRLRVTVNNAIRKA^SKLKN^EADNP^GKNQYQGF|GSSQTMQ^QVYRT
 PDVPI^IFMTAHGSIDTAVEAMRHG^AQDFLIKPC^EADRLRVTVNNAIRKA^SKLKN^EVDN[.]KNQYQGF|GSSQTMQ^AVYRT
 * * * * *

160
145

logo

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consensus

| D S A A S S K A S | F I T G E S G T G K E V C A E A I H A A S K R G D K P F I A I N C A A I P K D L I E S E L F G H V K G A F T G A A N D R R Q G A A E L A D G
 I D S A A S S K A S I F I T G E S G T G K E V C A E A I H A A S K R G D K P F I A I N C A A I P K D L I E S E L F G H V K G A F T G A A N D R R Q G A A E L A D G
 I D S A A S S K A S I F I T G E S G T G K E V C A E A I H A A S K R G D K P F I A I N C A A I P K D L I E S E L F G H V K G A F T G A A T E R R Q G A A E A A D G
 *

240
225

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GTLFLDELCEMDLDLQTKLLRFIQTGTGTFQKVGSSKMKSVDRVFCATNRDPWKEVQEGRFREDLYYRLYVIPHLPLPPLR_E
 GTLFLDELCEMDLDLQTKLLRFIQTGTGTFQKVGSSKMKSVDRVFCATNRDPWKEVQEGRFREDLYYRLYVIPHLPLPPLR_E
 GTLFLDELCEMDLDLQTKLLRFIQTGTGTFQKVGSSKMKSVDRVFCATNRDPWKEVQEGRFREDLYYRLYVIPHLPLPPLR_A

320
305

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RGE_DDI_EIA_IY_SLL_GY_FMS_HEE_GKS_DFY_RF_SQ_EEV_FER_SYE_IPG_NVR_QLQ_NVL_RN_YW_LN_GRE_ITL_DML_PPP_LN_QPV_WR_QS_MSA_PIN_N
 RGE_DDI_EIA_IY_SLL_GY_FMS_HEE_GKS_DFY_RF_SQ_EEV_FER_SYE_IPG_NVR_QLQ_NVL_RN_YW_LN_GRE_ITL_DML_PPP_LN_QPV_WR_QS_MSA_PIN_N
 RGD_DDI_EIA_IY_SLL_GF_MS_KEE_GK_DFV_RL_SAE_VER_FRQ_YEW_PGN_VR_QLQ_NVL_RN_YW_LN_GRE_ITL_DML_PPP_LN_QMS_APIN_N

400
385

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ASPLIHPNKVTSVSEIMPLWTEKQAI EQAI EQACDGNIPRAATYLDVSPSTIYRKLQTAINSRVKKQNV
 RALKFAEEDILSHDFPLWTEKQAI EQAI EQACDGNIPRAATYLDVSPSTIYRKLQTAINSRVKKQNV
 . ASKFI EPDI LTVSDI M PLWTEK M AIEQAI QACDGNIPRAATYLDVSPSTIYRKLQAWNSRDEKQNV .
 RALPLAHENKVSVEHIFPLWTEKQAIEQAI EACDGNIPRAATYLDVSPSTIYRKLQTWNEKVKEKEKER
 * * * * *

467
455

- X non conserved
- X similar
- X > 50% conserved