

logo

NTDB id 410640 GE573 RS01775 WP 190664418.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGKS I I I DAYSLVGGRSSEFVRYGE_AKAELEGLFLL_DSGHPV_FKVC
MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGKS I I I DAYSLVGGRSSEFVRYGE_TKAELEGLFLL_DSGHPV_FKVC
MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGKS I I I DAYSLVGGRSSEFVRYGE_AKAELEGLFLL_DSGHPV_LVC

80
80

logo

NTDB id 410640 GE573 RS01775 WP 190664418.1
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consensus

160
160

logo

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NTDB id 352 BSU 24240 NP 390304.2
consensus

A V Q E G V Q R V Y K L L K K Y K K L S E S E Q E M A H R L D L I Q F Q L E E I E S A K L E L N E D E L L Q E E R K Q I S N F E K I Y E S L Q N A Y N A L R S E
 A Y H E G Y E R Y M K L L K K Y K K L S E S E Q E M A H R L D L I Q F Q L E E I E S A K L E L N E D E L L Q E E R K Q I S N F E K I Y E S L Q N A Y N A L R S E
 T Y Q E G Y Q R Y Y K L L K K Y K K L S E S E Q E M A H R L D L I Q F Q L E E I E S A K L E L N E D E L L Q E E R K Q I S N F E K I Y E S L Q N A Y N A L R S E
 * * *

240
240

logo

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consensus

QGGLDVWGMSAQL^QEDISDINEPL^QK^MSESVS^SSYLL^SEDATFQMRN^MLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
 QGGLDVWGMSAQL^QEDISDINEPL^QK^MSESVS^SSYLL^SEDATFQMRN^MLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
 QGGLDVWGMSAQL^QEDISDINEPL^QK^MSESVS^SSYLL^SEDATFQMRN^MLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
 * * * * *

320
320

logo

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consensus

EDILEY^GSKIEEEEIDQIENRDSHL^{QS}EAL^{EA}LKKELE^DSVGKDVAVEEAA^NY^SQ^KIRK^TAWAKKL^AEE^{DA}IR^HRELKSLY^MME^EKSTFDTEF^kLV

400
400

logo

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consensus

480
480

logo

NTDB id 410640 GE573 RS01775 WP 190664418.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

560
560

logo

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consensus

HAQELLKQAGQVKTTG

HAQELLKQAGQVKTTG

HAQELLKQADQVKTTG

!! !!!!!!! !!!!!!!

576

576

- X

non conserved
- X

similar
- X

≥ 50% conserved