

logo

NTDB id 187911 BBJ33 RS11905 WP 003153174.1
NTDB id 352 BSU 24240 NP 390304.2
consensus

MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGK S I I I D A ySLLVGG RGSSEFVR YGE T KAELEGLFLL DSGHPV F E VC
MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGK S I I I D A ySLLVGG RGSSEFVR YGE T KAELEGLFLL DSGHPV F E VC
MLAELS IKNFAI I EELT ySFERGLTVLTGETGAGK S I I I D A ySLLVGG RGSSEFVR YGE A KAELEGLFLL DSGHPV LG VC

80
80

logo

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consensus

R EQGID ASDMIVMRRD STSGKSVCRVNGKLVLTIA SLREIGRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALQ
 A EQGID ASDMIVMRRD NAGTSGKSVCRVNGKLVLTIA ALREIGRLLLDIHGQHDNQLLMEDDKHLELLDKFAGAEAESALQ
 A EQGID VSDMIVMRRD STSGKSVCRVNGKLVLTIAS LREIGRLLLDIHGQHDNQLLMEDENHLQLLDKFAGAEVESALK

160
160

logo

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A Y H E G V 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 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 V 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 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 V Q R Y Y K L L K K L K Q 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 Q E E R Q Q I S N F E K I Y E S L Q N A Y N A L R S E
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240
240

logo

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consensus

QGGLDVWGMSAQL^QEDISDINEPL^QKK^MSESVS^SSYLL^SEDATFQMRN^MLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
 QGGLDVWGMSAQL^QEDISDINEPL^QKK^KSESVS^SSYLL^SEDATFQMRN^NLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
 QGGLDVWGMSAQL^QEDISDINEPL^QKK^KSESVS^SSYLL^SEDATFQMRN^NLD^LLE^FYDPERLN^FY^FETRLNEIKQLKRKYGATV
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320
320

logo

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consensus

EDILEY^GSKIEEEIDQENRDSHL^{QS}EAL^{EA}LKKE^{ES}VGKDVAVEAAN^{YS}Q^QIRK^{TA}WAKKL^{AD}EE^{EE}IR^{HQ}EL^{QS}LY^{ME}KS^{KS}TFDTEF^{LV}
EDILEY^GSKIEEEIDQENRDSHL^{QS}EAL^{EA}LKKE^{ES}VGKDVAVEAAN^{YS}Q^QIRK^{TA}WAKKL^{AD}EE^{EE}IR^{HQ}EL^{QS}LY^{ME}KS^{KS}TFDTEF^{LV}
EDILEY^GASKIEEEIDQENRDSHL^{QS}EAL^{EA}LKKE^{ES}VGKDVAVEAAN^{YS}Q^QIRK^{TA}WAKKL^{AD}EE^{EE}IR^{HQ}EL^{QS}LY^{ME}KS^{KS}TFDTEF^{LV}

400
400

logo

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R T D S S N S E A P V N G Q P V Q L T K Q G I D L V K F L I S T N T G E P L K S L S K V A S G G E L S R V M L A M K S I F S S Q Q D V T S I I F D E V D T G V
K T D P S A S E A P V N G Q P V Q L T K Q G I D L V K F L I S T N T G E P L K P L S K V A S G G E L S R V M L A M K S I F S S Q Q D V T S I I F D E V D T G V
R T A S R N E E A P V N G Q P V Q L T E Q G I D L V K F L I S T N T G E P L K S L S K V A S G G E L S R V M L A M K S I F S S Q Q D V T S I I F D E V D T G V

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480
480

logo

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consensus

SGRVAQAIAEKIHKVSTGSQVLCITHLPQVAAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMIAGVEVTDLTKR
SGRVAQAIAEKIHKVSTGSQVLCITHLPQVAAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMIAGVEVTDLTKR
SGRVAQAIAEKIHKVSTGSQVLCITHLPQVAAMADTHLYIAKEFKDGRITTHVTPLSKQDKVAEIGRMIAGVEVTDLTKR

560
560

logo

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consensus

HAQELLKQAGQVKTTG

HAQELLKQAGQVKTTG

HAQELLKQADQVKTTG

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

576

576

- X

non conserved
- X

similar
- X

≥ 50% conserved