

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



80  
80

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



160  
160

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



240  
240

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
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consensus



320  
320

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
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consensus



400  
400

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
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consensus



480  
480

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
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consensus



560  
560

logo

NTDB id 183677 A9E98 RS09315 WP 010947629.1  
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consensus

LGFWGI EAIRVDSFNKLSR T LTKHKDCKIAFINVNQGGERQAAELIAKHKQIPFVLI SKWPINDFAALGARGFLYKPI SI  
LGFWGIEAIRVDSFNKLSR T LTKHKDCKIAFINVNQGGERQAAELIAKHKQIPFVLI SKWPINDFAALGARGFLYKPI SI  
!!

640  
640

logo

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consensus

QKLQDL IESIANENQTEKNTNQELDTLREQLRFLHPEILIAEDNPVNKMLLTSLLNNNANITTVDDGEMAVTACEDKKFD  
QKLQDLIESIANENQTEKNTNQELDTLREQLRFLHPEILIAEDNPVNKMLLTSLLNNNANITTVDDGEMAVTACEDKKFD  
QKLQDLIESIANENQTEKNTNQELDTLREQLRFLHPEILIAEDNPVNKMLLTSLLNNNANITTVDDGEMAVTACEDKKFD  
!!

720  
720

logo

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consensus

MILLDLHMPKLNGL EAAKMI RQKSLMNKHSPIVLITASSSDLSSIDMKKYGVD FCFQKPID EKQLLIQILRIVDKTKHAA  
MILLDLHMPKLNGL EAAKMI RQKSLMNKHSPIVLITASSSDLSSIDMKKYGVD FCFQKPID EKQLLIQILRIVDKTKHAA  
MILLDLHMPKLNGL EAAKMI RQKSLMNKHSPIVLITASSSDLSSIDMKKSGVD FCFQKPID EKQLLIQILRIVDKTKHAA  
!!

800  
800

logo

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consensus

IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLE RLARRT  
IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLE RLARRT  
IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLE RLARRT  
!!

880  
880

logo

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consensus

ANADNL T EAFTDL IQS I DAVI SEYENQYSQ  
ANADNL T EAFTDL IQS I DAVI SEYENQYSQ  
ANADNL T EAFTDL IQS I DAVI SEYENQYSQ  
!!

910  
910

⊠ non conserved  
✖ similar  
⊠ ≥ 50% conserved