

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



80  
80

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



160  
160

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
NTDB id 1263 GC085 RS09730 WP 080450539.1  
consensus



240  
240

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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consensus



320  
320

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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consensus



400  
400

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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consensus



480  
480

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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consensus



560  
560

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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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 RQKSLMNKHSPIVLITASSSDLSSIDMKKSGVDFCFQKPIDEKQLLIQILRTV DKTTHAA  
MILLDLHMPKLNGL EAAKMI RQKSLMNKHSPIVLITASSSDLSSIDMKKSGVDFCFQKPIDEKQLLIQILRTV DKTTHAA  
MILLDLHMPKLNGL EAAKMI RQKSLMNKHSPIVLITASSSDLSSIDMKKSGVDFCFQKPIDEKQLLIQILRI VDKTKHAA  
!!

800  
800

logo

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IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLEKLARRT  
IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLEKLARRT  
IDWQLCVQKVS GNQALAE EFLAKFIEELYKNREEFI ELMHQKNVKGLADLAHKLHGACCF CGVPILQKRVAQLEKLARRT  
!!

880  
880

logo

NTDB id 193158 BE844 RS03055 WP 080477646.1  
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consensus

ANADNL<sub>A</sub>EAFTDLIQSIDAVISEYENQYSQ  
ANADNL<sub>A</sub>EAFTDLIQSIDAVISEYENQYSQ  
ANADNL<sub>A</sub>EAFTDLIQSIDAVISEYENQYSQ  
!!

910  
910

⊠ non conserved  
⊠ similar  
⊠ ≥ 50% conserved