

logo

NTDB id 964079 NSQ12 RS13125 WP 212082276.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF
MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF
MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF

80
80

logo

NTDB id 964079 NSQ12 RS13125 WP 212082276.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVDITPKIDGRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDMEYLYRQHI
KAVDITPKIDGRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI
KAVDITPKIDGRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI

160
160

logo

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consensus

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

240
240

logo

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consensus

GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVHSATAICLSYIVLLLFNP
GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVHSATAICLSYIVLLLFNP
GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVHSATAICLSYIVLLLFNP

320
320

logo

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consensus

YHLFEAGFQLSFAVSFSLILSSSIHQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTCILPGA
YHLFEAGFQLSFAVSFSLILSSSIHQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTCILPGA
YHLFEAGFQLSFAVSFSLILSSSIHQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTCILPGA

400
400

logo

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consensus

VAGVLLLSLSxSFGRLFFSWFDLLISHTNRLITNIADVbVFTIMAH PAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

VAGVLLLSLSV[SFGRLFFSWFDLLISHTNRLITNIADVbVFTIMAH PAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

VAGVLLLSLSA[SFGRLFFSWFDLLISHTNRLITNIADVbVFTIMAH PAPVLLFLFTVTIILLMAIEKRSLSQLMVTGQ

*

480
480

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consensus

CCTVMFLLFYPCLSSEGEVDMIDIGGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLLFYPCLSSEGEVDMIDIGGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLLFYPCLSSEGEVDMIDIGGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG

560
560

logo

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consensus



640
640

logo

NTDB id 964079 NSQ12 RS13125 WP 212082276.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



720
720

logo

NTDB id 964079 NSQ12 RS13125 WP 212082276.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

 non conserved
 similar
 ≥ 50% conserved