

logo

NTDB id 319250 D9C10 RS11260 WP 033881047.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLLIVCFFSFILFFVLYAVTDSQNVSSYRQGT YQF
MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLLIVCFFSFILFFVLYAVTDSQNVSSYRQGT YQF
MRNSRLLLPMAAA SATAGITAAAYFPAIFLFILFLLIILIKTRHAFLLIVCFFSFILFFVLYAVTDSQNVSSYRQGT YQF

80
80

logo

NTDB id 319250 D9C10 RS11260 WP 033881047.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDMEYLYRQHI
KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI
KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI

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

CCTVMFLLFYIYPCLSSEGEVDMIDIGQGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLLFYIYPCLSSEGEVDMIDIGQGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLLFYIYPCLSSEGEVDMIDIGQGDSMFVGVAPHQRGRVLI DTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG

560
560

logo

NTDB id 319250 D9C10 RS11260 WP 033881047.1
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consensus



640
640

logo

NTDB id 319250 D9C10 RS11260 WP 033881047.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



720
720

logo

NTDB id 319250 D9C10 RS11260 WP 033881047.1
NTDB id 107 BSU 25570 NP 390435.1
consensus



776
776

 non conserved
 similar
 ≥ 50% conserved