

logo

NTDB id 319526 D9C14 RS21905 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

[illegible]

80
80

logo

NTDB id 319526 D9C14 RS21905 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTTLEEPNHATVPGAFDYNEYLYRQHI
KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTTLEEPNHATVPGAFDYNEYLYRQHI
KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTTLEEPNHATVPGAFDYNEYLYRQHI

60
60

logo

NTDB id 319526 D9C14 RS21905 WP 009967776.1
NTDB id 107 BSU 25570 NP 390435.1
consensus

240
240

logo

NTDB id 319526 D9C14 RS21905 WP 009967776.1
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consensus

GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFP

20
20

logo

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YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA
YHLFEAGFQLSFAVSFSLILSSSIFQQVKTSLGQLTIVSLIAQLGSLPILLYHFHQFSIISVPMNMLMVPFYTFCILPGA

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:00

logo

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VAGVLLLSLSASFGRFFSFDLLISWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG
VAGVLLLSLSASFGRFFSFDLLISWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG
VAGVLLLSLSASFGRFFSFDLLISWINRLITNIADVDFVTIMIAHPAPVLLFLFTVTIILLMAIEKRSLSQLMVTGG

:80
:80

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ICCTVMFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLIDTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLIDTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG
 ICCTVMFLFIYPCLSSEGEVDMIDIGQGDSMFVGAPHQRGRVLIDTGGTLSYSSEPWREKQHPFSLGEKVLIPFLTAKG

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60

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640
640

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consensus



720
720

logo

NTDB id 319526 D9C14 RS21905 WP 009967776.1
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consensus



776
776

 non conserved
 similar
 ≥ 50% conserved