

logo

NTDB id 200988 KAB03 RS01550 WP 053530925.1
NTDB id 1044 HON27 RS16325 WP 206677389.1
consensus

MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ
MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ
!!

80
80

logo

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consensus

LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
!!

160
160

logo

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consensus

LAFLEFFVHGIQIAVQRFKLLEALTFHQSI EYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPFANE
LAFLEFFVHGIQIAVQRFKLLEALTFHQSI EYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPFANE
LAFLEFFVHGIQIAVQRFKLLEALTFHQSI EYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPFANE
!!

240
240

logo

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consensus

KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLV KWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
!!

320
320

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IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK
!!

400
400

logo

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consensus

HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
!!

480
480

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consensus

EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
!!

522
522

 non conserved
 similar
 $\geq 50\%$ conserved