

logo

NTDB id 267387 C2U32 RS15805 WP 032014308.1
NTDB id 1044 HON27 RS16325 WP 206677389.1
consensus

MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTL L GYSLLSLVQLL C FKF IATQATRQ
MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTL L GYSLLSLVQLL C FKF IATQATRQ
!!

80
80

logo

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consensus

LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
!!

160
160

logo

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consensus

LAF LFFVHGI GQ IAVQRFKLLEALTFHQSI ELYQLQNI NRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPFANE
LAF LFFVHGI GQ IAVQRFKLLEALTFHQSI ELYQLQNI NRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPFANE
LAF LFFVHGI GQ IAVQRFKLLEALTFHQSI ELYQLQNI NRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPFANE
!!

240
240

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KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLV KWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
!!

320
320

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IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLEEDLFDVK
!!

400
400

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consensus

HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
!!

480
480

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EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPPIY
!!

522
522

 non conserved
 similar
 $\geq 50\%$ conserved