

logo

NTDB id 326681 EGM95 RS19360 WP 001160330.1
NTDB id 1044 HON27 RS16325 WP 206677389.1
consensus

MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ
MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ
!!

80
80

logo

NTDB id 326681 EGM95 RS19360 WP 001160330.1
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consensus

LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL
!!

160
160

logo

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consensus

LAFLEFFVHGI GQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPPQFANE
LAFLEFFVHGI GQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPPQFANE
LAFLEFFVHGI GQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSL LGIPPPQFANE
!!

240
240

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consensus

KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLV KWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS
!!

320
320

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consensus

IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLL EEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLL EEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHMIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLL EEDLFDVK
!!

400
400

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consensus

HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITVNIHSQTNKIYIDVIDYEGEVSKRDISQLFKPFFST
!!

480
480

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consensus

EINGTGLGLYLSHSFCEANHAKLT YVEQKQGACFRIECP I IY
EINGTGLGLYLSHSFCEANHAKLT YVEQKQGACFRIECP I IY
EINGTGLGLYLSHSFCEANHAKLT YVEQKQGACFRIECP I IY
!!

522
522

 non conserved
 similar
 $\geq 50\%$ conserved