

logo

NTDB id 988820 AAHB48 RS17765 WP 001160333.1
NTDB id 1044 HON27 RS16325 WP 206677389.1
consensus

MQFPLNSSL SHTIFRLGTWYGLYRLIIAVSLNIIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLLCKFKFIATQATRQ
MQFPLNSSL SHTIFRLGTWYGLYRLIIAVSLNIIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLLCKFKFIATQATRQ
!!

80
80

logo

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consensus

LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL
LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL
LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL
!!

160
160

logo

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consensus

LAFLEFFVHGIQIAVQRFKLEALTFHQSIELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPFANE
LAFLEFFVHGIQIAVQRFKLEALTFHQSIELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPFANE
LAFLEFFVHGIQIAVQRFKLEALTFHQSIELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPFANE
!!

240
240

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consensus

KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS
KYPLAKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS
!!

320
320

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IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMIKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMIKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK
IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMIKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK
!!

400
400

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HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST
!!

480
480

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consensus

EINGTGLGLYLSSHFC EANHAKLTYVEKQKQACFRIECP IY
EINGTGLGLYLSSHFC EANHAKLTYVEKQKQACFRIECP IY
EINGTGLGLYLSSHFC EANHAKLTYVEKQKQACFRIECP IY
!!

522
522

 non conserved
 similar
 $\geq 50\%$ conserved