

logo

NTDB id 200895 KAB01 RS01400 WP 053530925.1  
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 GQIAVQRFKLLEALTFHQSI EYQLQNI NRYILEQIEEGYLVLDENYDIVSNPAACSL L GIPPFANE  
LAF LFFVHGI GQIAVQRFKLLEALTFHQSI EYQLQNI NRYILEQIEEGYLVLDENYDIVSNPAACSL L GIPPFANE  
LAF LFFVHGI GQIAVQRFKLLEALTFHQSI EYQLQNI NRYILEQIEEGYLVLDENYDIVSNPAACSL L GIPPFANE  
!!

240  
240

logo

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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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HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
!!

480  
480

logo

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consensus

EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECP I IY  
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECP I IY  
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECP I IY  
!!

522  
522

 non conserved  
 similar  
  $\geq 50\%$  conserved