

logo

NTDB id 334235 EQH48 RS17335 WP 032008286.1  
NTDB id 1044 HON27 RS16325 WP 206677389.1  
consensus

MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ  
MQFPLNSSL SHTIFRLG TWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLGYSLLSLVQLL CFKFIATQATRQ  
!!

80  
80

logo

NTDB id 334235 EQH48 RS17335 WP 032008286.1  
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consensus

LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL  
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL  
LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGS LFDYNNLNTIGNSAL  
!!

160  
160

logo

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consensus

LAFLEFFVHGIQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPPQFANE  
LAFLEFFVHGIQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPPQFANE  
LAFLEFFVHGIQIAVQRFKLLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVSNPAACSL LGIPPPQFANE  
!!

240  
240

logo

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consensus

KYPLAKWHADLFEILKFGDLKEGDRFIFESR LSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS  
KYPLAKWHADLFEILKFGDLKEGDRFIFESR LSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS  
KYPLV KWHADLFEILKFGDLKEGDRFIFESR LSAYSINIKVQHLLVPQQTLLILQDAQQINQQAQQLKLAALGQLSAS  
!!

320  
320

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IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMI GKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK  
IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMI GKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK  
IAHEIRNPLAAIVQANELLKSDSPEQQNTRHMI GKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFID TLLEEDLFDVK  
!!

400  
400

logo

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HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
!!

480  
480

logo

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consensus

EINGTGLGLYLSHSFCEANHAKLTYVEQKQGA WFER  
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGA WFER  
EINGTGLGLYLSHSFCEANHAKLTYVEQKQGA WFER  
!!

515  
522

 non conserved  
 similar  
  $\geq 50\%$  conserved