

logo

NTDB id 110498 ACAK33 RS03160 WP 001160337.1  
NTDB id 1044 HON27 RS16325 WP 206677389.1  
consensus

MQFPLNSSL SHTIFRLGTWYGLYRLIIAVSLNII LVL TDAQTDNSLQQPALYSY TLLGYSLLSLVQLL CFKFIATQATRQ  
MQFPLNSSL SHTIFRLGTWYGLYRLIIAVSLNII LVL TDAQTDNSLQQPALYSY TLLGYSLLSLVQLL CFKFIATQATRQ  
!!

80  
80

logo

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consensus

LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL  
LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL  
LILFFIVDIIICLSLLTFSVGEPNLQSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLEFDYNNLNTIGNSAL  
!!

160  
160

logo

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consensus

LAFLEFFVHGIQIAVQRFKLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPQFANE  
LAFLEFFVHGIQIAVQRFKLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPQFANE  
LAFLEFFVHGIQIAVQRFKLEALTFHQSI ELYQLQINRYILEQIEEGYLVLDENYDIVVSNPAACSLGIPPQFANE  
!!

240  
240

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KYPLAKWHTDLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS  
KYPLAKWHTDLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS  
KYPLAKWHTDLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPPQTLTLLILQDAQQINQQAQQLKLAALGQLSAS  
!!

320  
320

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IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK  
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK  
IAHEIRNPLAAIVQANELLKSDSPEQQNTRLRHIGKQTKRIDSIVQDTLGLARSERTHP IQIDVKHFIDTLLLEEDLFDVK  
!!

400  
400

logo

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consensus

HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPIYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPIYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPIYITINIHSQTNKIYIDVIDYGEVSKRDISQLFKPFFST  
!!

480  
480

logo

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consensus

EINGTGLGLYLSSHFC EANHAKLTYVEQKQGACFRIECP IY  
EINGTGLGLYLSSHFC EANHAKLTYVEQKQGACFRIECP IY  
EINGTGLGLYLSSHFC EANHAKLTYVEQKQGACFRIECP IY  
!!

522  
522

 non conserved  
 similar  
  $\geq 50\%$  conserved