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logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  MDIR.RTQAELNQYHLGLWYGAYRLIIAFCLLLIYLLTTQTLNHDYQHPRLYAASLVLYFIFSCAQLLYFRYVRHRIPLQ   79
NTDB id 1044 HON27 RS16325 WP 206677389.1      MQFPLNSSLSHTIFRLGTWYGLYRLIIAVSLNIILVLTDAQTDNSLQQPALYSYTLLGYSLLSLVQLLCFKFIATQATRQ   80
consensus                                      !  ** *   *  **!! !!! !!!!!! *! *! *!!     *  !*! !!  *!* ! * !* !!!*!*** *  * !

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  LTLIFSVDVLVLSSLTFALDGPSLHLSLLRVIAIFAATLLLEKSKALFITLVAMICVTYQHFVGTLLDFSSLDRIGQSAL  159
NTDB id 1044 HON27 RS16325 WP 206677389.1      LILFFIVDIICLSLLTFSVGEPNLQLSLLYVIAIFTSAILLSARMSLLITLLAVIAVIYQRFVGSLFDYNNLNTIGNSAL  160
consensus                                      ! ! ! !!***!! !!! *  ! !*!!!!*!!!!!   *!!     ! !!!*!*!*! !!*!!!*! !*  !  !!*!!!

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  LSFLFFIIYRFGQIAVRRFQLLEHLNFSQSLELNRLKNINRYILEQTDMGYLVLDENCHIMVSNPAACHLLGINPLYAFD  239
NTDB id 1044 HON27 RS16325 WP 206677389.1      LAFLFFVVHGIGQIAVQRFKLLEALTFHQSIELYQLQNINRYILEQIEEGYLVLDENYDIVVSNPAACSLLGIPPQFANE  240
consensus                                      ! !!!!***  !!!!! !! *!!*! !*!!*!!  ! !!!!!!!!! * !!!!!!!!**!*!!!!!!!*!!!!*! *!  *

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  KYPLYKVQPDLFQLLQFDQLSNGEKFQFESQLSHFLIQIQVQKLVPQQTLTLLVLQDARQINQRVQQLKLAALGQLSAS  319
NTDB id 1044 HON27 RS16325 WP 206677389.1      KYPLVKWHADLFEILKFGDLKEGDRFIFESRLSAYSINIKVQHLLVPQQTLTLLILQDAQQINQQAQQLKLAALGQLSAS  320
consensus                                      !!!! ! **!!! *! !  !  !**! !!! !!** !*! !!*!*!!!!!!!!!*!!!! !!!!  !!!!!!!!!!!!!!

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  IAHEIRNPLAAIVQANRLLAGSDTDQQQMLSQMIDKQSQRIDKIIDDTLSMVKNKETTPVQIDLARFIPQLIQDDLADLA  399
NTDB id 1044 HON27 RS16325 WP 206677389.1      IAHEIRNPLAAIVQANELLKDSDPEQQNTLRHMIGKQTKRIDSIVQDTLGLARSERTHPIQIDVKHFIDTLLEEDLFDVK  400
consensus                                      !!!!!!!!!!!!!!!! !!  !!**!!* ! *!! !!* !!! !* !!! * *   !*!*!!!* *!!* !* *!! !*

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  SQIQVSQ.PEPLQILFDESQLRQVLINLIRNAQRHNAPDHPI.ELQLYRQQDTVYMDVKDYGSGVASHDIRYLFQPFFST  477
NTDB id 1044 HON27 RS16325 WP 206677389.1      HSIQLKISDSSLKFLFDEKQLRQVMINLVRNALRHNAPDSPYITINIHSQTNKIYIDVIDYGEGVSKRDISQLFKPFFST  480
consensus                                      * !!*  ** *!  !!!! !!!!!*!!!*!!! !!!!!!*!  * **** !   *!*!! !!! !!  *!!  !! !!!!!

logo

NTDB id 1106875 ACLKM6 RS00845 WP 104489117.1  EISGTGLGLYLSHSFCEANRAKLRYVEQKQGACFRIECSPVDVNK  522
NTDB id 1044 HON27 RS16325 WP 206677389.1      EINGTGLGLYLSHSFCEANHAKLTYVEQKQGACFRIECPIIY...  522
consensus                                      !! !!!!!!!!!!!!!!!!*!!! !!!!!!!!!!!!!!!*** ***
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X non conserved
X similar
X $\geq$50% conserved