```
logo
NTDB id 24844 PFL01 RS24360 WP 011335993.1  MIAEADNADSKQAQRLLRLYHLYRLSVGITLVLLISSNMDNRLLTSANDEWLRNGSWLYLVLNILLVVFLENIRRPAQLFGLALV   85
NTDB id 1472 PAKAF RS23995 WP 016253893.1   MRAERLRLSEEQGQRILRLYHLYRLTIGLVLVLLISSELEDQVLKLVHPELFHVGSWCYLVFNILVALFLPPSRQLLPIFILALT   85
consensus                                   ! !!        !*!!*!!!!!!!!!**!* !!!!!!! **  *!   **!  * !!!*!!! !!!* *!!** ! * **! !!!

logo
NTDB id 24844 PFL01 RS24360 WP 011335993.1  DILLLCGVFYAAGGVASAIGNLLIVSVAISNTLLRRRIGLLIAAIATLGIVGLSFLLSFSHPLSANDYLQAGTLGALCFAASLLV   170
NTDB id 1472 PAKAF RS23995 WP 016253893.1   DVLMLCGLFYAGGGVPSGIGSLLVVAVAIANILLRGRIGLVIAAAASLGLLYLTFFLSLSSPDATNHYVQAGGLGTLCFAAALVI   170
consensus                                   !*!*!!!*!!!*!!!*!*!! !!*! !!! ! !!! !!!!*!!! !*!!** !*! !! !*!   !*!*!!! !! !!!!! !**

logo
NTDB id 24844 PFL01 RS24360 WP 011335993.1  QGLMRRLEVSENLAEQRASEVVGLEALNALILQRMRTGILVLDEQRRVQLANHSARVLLGQTHLEGDLIDEHSSALVERLQLWLN   255
NTDB id 1472 PAKAF RS23995 WP 016253893.1   QALVRRQEQTETLAEERAETVANLEELNALILQRMRTGILVVDSRQAILLANQAALGLLRQDDVQGASLGRHSPMLMHCMKQWRL   255
consensus                                   !*!*!! ! *! !!! !!  !  !! !!!!!!!!!!!!!!*!    * !!!* !  !! ! ** !  *  !!* !****  !

logo
NPTLRPQSLKIAGNGLELQPSFIALEQNPNRQTLVFLEDLAQIAQQAQQLKLAALGRLTAGIAHEIRNPLGAISHAAQLLQESEE   340
NPSLRPPTLKVVPDGPTVQPSFISLNREDDQHVLIFLEDISQIAQQAQQMKLAGLGRLTAGIAHEIRNPLGAISHAAQLLQESEE   340
consensus                                   !!*!!!**!!* * !* *!!!!! !   *  * !*!!!!* !!!!!!!!*!!!*!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved