```
logo
NTDB id 66369 AS4 RS09745 WP 096734012.1  ..........MMNLIKHKKNQTVYAQKSTGFTLIELLVAIAIFAVLSALGWKIFDYLIKVKDRNVIHEENLGQLQGAYQQIQRDT   75
NTDB id 1076 ABD1 RS08075 WP 000594585.1  MIKNKYIHIRSIASRLAARSSSARLTRASGFTLVELLVAIAIFAVLSLLGWKIFDYLLKVRDRNAEHEVHLFELQDAYQQILRDT   85
consensus                                 ********** *    * *  *    * *!!!!*!!!!!!!!!!!!! !!!!!!!!!*!!*!!!  !! *!  !! !!!!! !!!

logo
NTDB id 66369 AS4 RS09745 WP 096734012.1  LQIVPISARSGEEIEPALGLDNQKLSFSKTGVTDPLKQGLSPYERIEYVYNAQEKKLYRLKYSNLNRNNSEQPLSSVLLADVDQY  160
NTDB id 1076 ABD1 RS08075 WP 000594585.1  LQIIPLSANQGGQLHPALEIDNQILRFSKAGVTDPLKQGLSPFERIEYRYDADQKKLYRLKYTNLNTSNREQPLSSTLLSQVDQY  170
consensus                                 !!!*!*!!  !  **!!! *!!! ! !!! !!!!!!!!!!!!*!!!!! ! !  !!!!!!!!*!!!  ! !!!!!! !!  !!!!

logo
NTDB id 66369 AS4 RS09745 WP 096734012.1  RIDVLNPTELDRWPNDMIKN.....DGQLPRGIKIKVTIRDIEYEWIFSLLNSDYLINNGAAN   218
NTDB id 1076 ABD1 RS08075 WP 000594585.1  QIMVLTPQEVTKWPEVNIDPTKPNELKKLPKGIKIQLTVAGVNYEWIYSLNQGDLSLSQEGNS   233
consensus                                  ! !! ! !* *!!   ! ******   !!*!!!! *!*  * !!!!*!! * !  * * *
```

X non conserved

X similar

X ≥ 50% conserved