```
logo

NTDB id 1008694 ABNP33 RS10015 WP 262439608.1  MIKNKSFRSPSSASPLAVRSEQMMRGYSEQSSSSCAQTKHCFKDSHTRNIVSHLATRSSSARLTRASGFTLVELLVSIAI   80
NTDB id 1076 ABD1 RS08075 WP 000594585.1       ....................................MIKNKYIHIRSIASRLAARSSSARLTRASGFTLVELLVAIAI   42
consensus                                      ************************************ ** !  ! ! ! !*!! !!!!!!!!!!!!!!!!!!!! !!!

logo

NTDB id 1008694 ABNP33 RS10015 WP 262439608.1  FAILSLLGWKVFDYLLKVRDRNTQHEVQLFELQDAYQQVLRDTLQIIPLSANQGGQLRPALEIDNQVLRFSKAGVTDPLK   160
NTDB id 1076 ABD1 RS08075 WP 000594585.1       FAVLSLLGWKIFDYLLKVRDRNAEHEVHLFELQDAYQQILRDTLQIIPLSANQGGQLHPALEIDNQILRFSKAGVTDPLK   122
consensus                                      !!*!!!!!!!*!!!!!!!!!!!  !!!*!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!*!!!!!!!!*!!!!!!!!!!!!!

logo

NTDB id 1008694 ABNP33 RS10015 WP 262439608.1  QGLSPFERIEYRYDPDQKKLYRLKYSNLNTLNREQPLSSTLLSQVEQFQIMVLTPQEVTKWPEVNIDPTKPDEFRKLPKG   240
NTDB id 1076 ABD1 RS08075 WP 000594585.1       QGLSPFERIEYRYDADQKKLYRLKYTNLNTSNREQPLSSTLLSQVDQYQIMVLTPQEVTKWPEVNIDPTKPNELKKLPKG   202
consensus                                      !!!!!!!!!!!!!!!*!!!!!!!!!!!*!!!!! !!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!!!! ! *!!!!!

logo

NTDB id 1008694 ABNP33 RS10015 WP 262439608.1  IKMQLTVAGVSYEWIYSLNQGDLSLSKQGGQ   271
NTDB id 1076 ABD1 RS08075 WP 000594585.1       IKIQLTVAGVNYEWIYSLNQGDLSLSQEGNS   233
consensus                                      !!*!!!!!!! !!!!!!!!!!!!!!!  !
```

X non conserved

X similar

X ≥ 50% conserved