```
logo

NTDB id 395911 GFH30 RS07695 WP 153371673.1  MIAKLRACQISNKPQEKLSSRQKTRGFLHSALGFTLVELLVAIAIFAVLSALGWKVFDHIISTKDQNTMHEQRLSAVQQG   80
NTDB id 1076 ABD1 RS08075 WP 000594585.1     MIKN.KYIHI..RSIASRLAARSSSARLTRASGFTLVELLVAIAIFAVLSLLGWKIFDYLLKVRDRNAEHEVHLFELQDA   77
consensus                                    !!   ** **!****         * * !* ! !!!!!!!!!!!!!!!!! !!!!*!!***  *! !  !! *!  *! *

logo

NTDB id 395911 GFH30 RS07695 WP 153371673.1  YQQILRDTVQIVPLTSNLNGEIRPALVLQNGRLSFSKTGVTDPLQQGISPDERVEYQYRQDEKKLYRLRYQNLNQTGREQ   160
NTDB id 1076 ABD1 RS08075 WP 000594585.1     YQQILRDTLQIIPLSANQGGQLHPALEIDNQILRFSKAGVTDPLKQGLSPFERIEYRYDADQKKLYRLKYTNLNTSNREQ   157
consensus                                    !!!!!!!!*!!*!!* !  ! **!!! * !  ! !!! !!!!!! !!*!! !!*!! !  ! !!!!!!*! !!! * !!!

logo

NTDB id 395911 GFH30 RS07695 WP 153371673.1  PDSSVLLENVDQFQIMVLNPDELSEWPKNGVDLEQIIEKQKLPKGIKINLTIHDVEYEWLFSLLNTDFLNQTKVNV   236
NTDB id 1076 ABD1 RS08075 WP 000594585.1     PLSSTLLSQVDQYQIMVLTPQEVTKWPEVNIDPTKPNELKKLPKGIKIQLTVAGVNYEWIYSLNQGDLSLSQEGNS   233
consensus                                    ! !! !! *!!!*!!!!! ! !** !!   *!*  * !  !!!!!!!!*!!** ! !!!**!! * !        !
```

X non conserved

X similar

X ≥ 50% conserved