```
logo

NTDB id 305661 DV115 RS10315 WP 003752754.1  MCTRKQQGFTLIELLIVMVIAAIMAMIALPSMSRWIASRRIVSHAEQVANLLRFSRGEAVRLNIPVYICPVQVKKDGTVN   80
NTDB id 1136 NGFG RS02420 WP 003687912.1     MCTRKQQGFTLTELLIVMAIAAIMATIALPNMSGWIASRRIASHAEQVANLLRFSRGEAVRLNLPVYICPVQVKKDGAPN   80
consensus                                    !!!!!!!!!!! !!!!!! !!!!!! !!!! !! !!!!!!! !!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!  *!

logo

NTDB id 305661 DV115 RS10315 WP 003752754.1  NQCDFSKKGQGVLAFGDKNGNKAYDGDAADVFLRSVVLNDAADSRINYAFNHIAFGKSQPTADRVVWTFNQNGAFGYSPD  160
NTDB id 1136 NGFG RS02420 WP 003687912.1     NRCDFSKKGRGMLAFGDKNGNKAYDNDTVDVFLRSVVLNDTDDSRINYAFNHIAFGSSQPTADRVVWTFNQNGTFGYLPD  160
consensus                                    ! !!!!!!! !*!!!!!!!!!!!!!! !  !!!!!!!!!!!  !!!!!!!!!!!!!! !!!!!!!!!!!!!!!! !!! !!

logo

NTDB id 305661 DV115 RS10315 WP 003752754.1  QNLARTSRFVYSDGYIQIVLTDAKAVSADEKKFRSAVVLIDSSGRVEVCPRNDTRAVCQHQ   221
NTDB id 1136 NGFG RS02420 WP 003687912.1     QNLKDNSKFVYSDGYIQIVLTDARAVSDADKKFRSAVVLINSSGRVEVCRKNDTRAVCKH.   220
consensus                                    !!!   !*!!!!!!!!!!!!!!!!*!!!  *!!!!!!!!!! !!!!!!!!**!!!!!!! !*
```

X non conserved

X similar

X $\geq$ 50% conserved