```
logo
NTDB id 291803 DIR46 RS17215 WP 109346326.1  MTTSPQSPALTIMVIDDSSTIRRSAEIFLNQAGYQVVLADDGFDALAKINDHHPALVFCDILMPRLDGYQTCALIKKSAR   80
NTDB id 1047 HON27 RS03110 WP 000389061.1    ..MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQN   78
consensus                                    **  *  * ! *!!!!!! !!!!*!!  !*  !* !* ! !!!*!! !! ***! *!!*!!*!!!!!!!!!!!!!! !

logo
NTDB id 291803 DIR46 RS17215 WP 109346326.1  FRATPVLMLSSKDGLFDRARGAMVGASAYLTKPFTKDSLLAAVREHAMPQA   131
NTDB id 1047 HON27 RS03110 WP 000389061.1    YQNIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS..   127
consensus                                    *   !!*!!!!!!!!!! !*! *!!   !!!!!!*!! !! !*! !  ***
```

X non conserved

X similar

X $\geq$ 50% conserved