```
logo

NTDB id 149765 ACG33 RS01460 WP 066918087.1  MENLATTQATRLAGLKVLVIDDSKTIRRTAETLLSKEGCEVFTAVDGFDALSKIADHQPDIVFVDIMMPRLDGYQTCSLI   80
NTDB id 1047 HON27 RS03110 WP 000389061.1    .......MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALI   73
consensus                                    *******       !!!*!!!!!!!!!!!!!!!! *!!!!! !!!!!!*!!!!!!***!!!!!!!!!!!!!!!!!!!! !!

logo

NTDB id 149765 ACG33 RS01460 WP 066918087.1  KHNKVFRSTPVIMLSSKDGLFDRARGRIVGSEHYLTKPFTKDELLNAIEAHVGEVSR   137
NTDB id 1047 HON27 RS03110 WP 000389061.1    KNSQNYQNIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS...   127
consensus                                    !*   *   !!!!!!!!!!!!! !*!!*!!!**!!!!!!*!!!!!!!!  !!  ***
```

X non conserved

X similar

X $\geq$ 50% conserved