```
logo                                                 MSNLQNYKVMVIDDSNTIRRTAETFLQREGCEVITAYDGFEALSKIAEHNPDYYFVDYMMPRLDGYQTCSLIKNSQRYQ
                                                     MEDAFAGLKVMVIDDSKTIRRSAEIFLGQAGCEVILAEDGFDALAKIADAHPAYIFVDIMMPRLDGYQTCALIKKNPNFK

NTDB id 1006239 ABHF33 RS06845 WP 348946222.1  .MSNLAGVKVMVIDDSNTIRRSAEIFLGQAGCEVILAEDGFDALAKIADHHPAVIFVDVMMPRLDGYQTCSLIKKNPRFK   79
NTDB id 1047 H0N27 RS03110 WP 000389061.1      MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ   80
consensus                                      *       *!!!!!!!! !!!!*!!  !   !!!!! ! !!!*!! !!!***! **!!!*!!!!!!!!!!! !!!  * *

logo                                                 NTPVVMLSSKDGLFDRARGRMVGSDEYLTKPFTKDSLLNAYRVHVSS
                                                     ATPVVMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHAGIV

NTDB id 1006239 ABHF33 RS06845 WP 348946222.1  ATPVVMLSSKDGLFDRARGRMVGSDEYLTKPFTKDSLLAAVRVHAGIV   127
NTDB id 1047 H0N27 RS03110 WP 000389061.1      NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.   127
consensus                                        !!*!!!!!!!!!! !*!!*!!!!!!!!!!!!*!! !! !*! !   *
```

X non conserved

X similar

X $\geq$ 50% conserved