```
logo
NTDB id 1007669 ABL848 RS05990 WP 093205631.1  MGPNGSGYKVLVIDDSNTIRRSAEIFLKQGGHEVLLAEDGFDALSKVNDHKPHLIFCDILMPRLDGYQTCAIIKRNAHFS   80
NTDB id 1047 H0N27 RS03110 WP 000389061.1      MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ   80
consensus                                      ! *     !!*!!!!! !!!!*!!  !   ! !!* ! !!!*!!!!* ** !***!*!!*!!!!!!!!!!!*!!    **

logo
NTDB id 1007669 ABL848 RS05990 WP 093205631.1  NVPVVMLSSKDGVFDKARGRMVGSQDYLTKPFTKDQLLQAVQQFGIVQQEVQ   132
NTDB id 1047 H0N27 RS03110 WP 000389061.1      NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....   127
consensus                                      !*!!*!!!!!!!*!! !*!!*!!! *!!!!!!*!! !!*!* ** * *****
```

X non conserved

X similar

X $\geq$ 50% conserved