```
logo
NTDB id 106791 ACAM42 RS28355 WP 021012487.1  MGPNGSGYKVLVIDDSNTIRRSAEIFLKQGGHEVLLAEDGFDALSKVNDHKPHLIFCDILMPRLDGYQTCAIIKRNAHFS   80
NTDB id 1047 HON27 RS03110 WP 000389061.1     MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ   80
consensus                                     ! *      !!*!!!!! !!!!*!!   !    ! !!* ! !!!*!!!!* ** !***!*!!*!!!!!!!!!!!*!!    **

logo
NVPVVMLSSKDGVFDKARGRMVGSQDYLTKPFTKDQLLQAVQQFGIVQQEVQ
NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....
consensus                                     !*!!*!!!!!!!*!! !*!!*!!! *!!!!!!*!! !!*!* **   *****
NTDB id 106791 ACAM42 RS28355 WP 021012487.1  NVPVVMLSSKDGVFDKARGRMVGSQDYLTKPFTKDQLLQAVQQFGIVQQEVQ   132
NTDB id 1047 HON27 RS03110 WP 000389061.1     NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....   127
consensus                                     !*!!*!!!!!!!*!! !*!!*!!! *!!!!!!*!! !!*!* **   *****
```

X non conserved

X similar

X $\geq$ 50% conserved