```
logo
NTDB id 58348 AVCA6 RS01505 WP 012699006.1  MEHHSDGLKVMVIDDSKMIRRTAETLLKRVGCEVITAVDGFDALAKIADGHPRIIFVDIAMPRLDGYQTCALIKNNSAFKSTPVI   85
NTDB id 1047 HON27 RS03110 WP 000389061.1   MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
consensus                                   !!**   !!!!!!!!!! !!!!!!!!! ! !!!!!!!!!!!*!! !!!***! !*!!!! !!!!!!!!!!!!!!!    *    !!!

logo
NTDB id 58348 AVCA6 RS01505 WP 012699006.1  MLSSRDGLFDKAKGRIVGSEQYLTKPFGKEELLGAIRAHVPDFVPLEQAS   135
NTDB id 1047 HON27 RS03110 WP 000389061.1   MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS........   127
consensus                                   !!!!*!!!!! !!!!*!!!* !!!!!!! !*!!! !!! !!* ********
```

X non conserved

X similar

X $\geq$ 50% conserved