```
logo
NTDB id 92692 HQS1 RS26635 WP 016447937.1  MTTSGAPLRVLVVDDSNTIRRSAEIFLKQGGHEVLLADDGFDALSKVNDYQPQLIFCDILMPKLDGYQTCAIIKRNARFADTPVV   85
NTDB id 1047 HON27 RS03110 WP 000389061.1  MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQNIPVI   85
consensus                                  !      *!*!*!*!!! !!!!*!!  !   ! !!* ! !!!*!!!!* * *! **!*!!*!!*!!!!!!!!*!!     *   !!*

logo
NTDB id 92692 HQS1 RS26635 WP 016447937.1  MLSSKDGVFDKARGRMVGCQDYLTKPFTKDQLLQAVEQFGKQTIGAL   132
NTDB id 1047 HON27 RS03110 WP 000389061.1  MLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS.....   127
consensus                                  !!!!!!!*!! !*!!*!!* *!!!!!!*!! !!*!* **   *****
```

X non conserved

X similar

X $\geq$ 50% conserved