```
logo

NTDB id 1140658 DQN24 RS02940 WP 111695366.1  MSMNLLPWRTYQHQNRLRRLAFYIALFILLAINLTLAFSNLIEQQKQHLQAQQTSFEQLNQQLYKTAKKIEQLRSAVQVG   80
NTDB id 1370 NTHI RS02660 WP 011272049.1      MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQTSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1344 HI 0438 AAC22097.1               MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRIAVKVG   80
consensus                                     !!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!*!!!!!*!!!!!!!!!*!!***!*!!!*!!*!!

logo

NTDB id 1140658 DQN24 RS02940 WP 111695366.1  EILTSIPKEQVKKSLQLLSELPFQQGELNKFKQDAHNLSLEGLAQDQREFELIHQFLKKHFPNVKLSQFQPEQQSLYFLF  160
NTDB id 1370 NTHI RS02660 WP 011272049.1      EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
NTDB id 1344 HI 0438 AAC22097.1               EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
consensus                                     !*!!!!!*!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!*!!!!*!!!!!!!!!!!!!!!!!!!!!!*!!!!**!*!*!

logo

NTDB id 1140658 DQN24 RS02940 WP 111695366.1  DVEQVGEDK   169
NTDB id 1370 NTHI RS02660 WP 011272049.1      DVEQGAEK.  168
NTDB id 1344 HI 0438 AAC22097.1               DVEQGAEK.  168
consensus                                     !!!!**!*
```

X non conserved

X similar

X ≥ 50% conserved