```
logo

NTDB id 1142187 DQN30 RS03535 WP 111696320.1  MSMNLLPWRTNQHQKRLRRLAFYIALFILLAINLMLAFNNLIEQQKQNLQAQQKSFEQLNQQFHKTTMQIDQLRSAVKVS   80
NTDB id 1370 NTHI RS02660 WP 011272049.1      MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQTSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1344 HI 0438 AAC22097.1               MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRIAVKVG   80
consensus                                     !!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!*!!!!!!!!*!!!!!!!!!!!!*!!!!*

logo

NTDB id 1142187 DQN30 RS03535 WP 111696320.1  EVLQPLTNEQVQNSLQRLSELPLQAGELNRFKQDAHNLNLEGRAQDQMEFESVHQFLKKHFPNVKLSQVQPEQHTLFFRF  160
NTDB id 1370 NTHI RS02660 WP 011272049.1      EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
NTDB id 1344 HI 0438 AAC22097.1               EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
consensus                                     !!!****!!!!**!!!*!!!!!*!*!!!!*!!!!!*!!*!!!*!!!!*!!!**!!!!!!!!!!!!!!!!!!!!*!!!!*!

logo

NTDB id 1142187 DQN30 RS03535 WP 111696320.1  DVEEVGEDK   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