```
logo

NTDB id 275932 C3V42 RS02810 WP 065265618.1   MSMNLLPWRIYQYQKRLRRLAFYVALFILLAINLMLAFSNLIEQQKQNLQTQQKYFEQLNQQLHKTTTQIDLLRSAVKVG   80
NTDB id 1370 NTHI RS02660 WP 011272049.1      MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQTSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1344 HI 0438 AAC22097.1               MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRIAVKVG   80
consensus                                     !!!!!!!!!*!!*!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!**!!!!!!!!!!!!!*!!!*!!*!!!!!

logo

NTDB id 275932 C3V42 RS02810 WP 065265618.1   EVLHSLTNGQVQNSLQRLSELPLQKGELNRFKQDGHNLILEGRAQDQMEFESVHQFLKQHFPNVKLSQVQPEQHTLFFRF   160
NTDB id 1370 NTHI RS02660 WP 011272049.1      EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF   160
NTDB id 1344 HI 0438 AAC22097.1               EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF   160
consensus                                     !!!*!**!*!!**!!!*!!!!!*!*!!!!*!!!!**!!*!!!*!!!!*!!!**!!!!!*!!!!!!!!!!!!!!!*!!!!*!

logo

NTDB id 275932 C3V42 RS02810 WP 065265618.1   DMEQGEEK   168
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