```
logo

NTDB id 309984 CH613 RS02955 WP 011961685.1   MSMNLLPWRTYQHQNRLRRLAFYIALFILLAINLTLAFSNLIEQQKQNLQAQQTSFEQLNQQLYKTAKQIEQLRSAVQVG   80
NTDB id 1370 NTHI RS02660 WP 011272049.1      MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQTSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1344 HI 0438 AAC22097.1               MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRIAVKVG   80
consensus                                     !!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!*!!**!!*!!!*!!*!!

logo

NTDB id 309984 CH613 RS02955 WP 011961685.1   EILTSIPNEQVKKSLQLLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF   160
NTDB id 1370 NTHI RS02660 WP 011272049.1      EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF   160
NTDB id 1344 HI 0438 AAC22097.1               EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF   160
consensus                                     !*!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!
```