```
logo
NTDB id 310159 CH582 RS02965 WP 116941733.1  MSMNLLPWRTYQHQNRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1370 NTHI RS02660 WP 011272049.1     MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQTSFEQLNQQLHKTTMQIDQLRSAVKVG   80
NTDB id 1344 HI 0438 AAC22097.1              MSMNLLPWRTYQHQKRLRRLAFYIALFILLAINLMLAFSNLIEQQKQNLQAQQKSFEQLNQQLHKTTMQIDQLRIAVKVG   80
consensus                                    !!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!

logo
NTDB id 310159 CH582 RS02965 WP 116941733.1  EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQFQPEQDTLFFHF  160
NTDB id 1370 NTHI RS02660 WP 011272049.1     EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
NTDB id 1344 HI 0438 AAC22097.1              EVLTSIPNEQVKKSLQQLSELPFQQGELNKFKQDANNLSLEGNAQDQTEFELIHQFLKKHFPNVKLSQVQPEQDTLFFHF  160
consensus                                    !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!

logo
NTDB id 310159 CH582 RS02965 WP 116941733.1  DVEQGAEK  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