```
logo

NTDB id 1140657 DQN24 RS02935 WP 111695365.1  MKTFFNDPFTPFGKWLSQPVYIQSLTFLLLLSAVIFRPVLDYIEASEHFAETEYELAKKSSEFLHQQKILTSLQQQSESR   80
NTDB id 1345 HI 0437 AAC22096.1               MKAFFNDPFTPFGKWLSQPFYVHGLTFLLLLSAVIFRPVLDYIEGSSRFHEIENELAVKRSELLHQQKILTSLQQQSESR   80
NTDB id 1371 NTHI RS02655 WP 011272048.1      MKAFFNDPFTPFGKWLSQPFYVHGLTFLLLLSAVIFRPVLDYIEGSSRFHEIENELAVKRSELLHQQKILTSLQQQSESR   80
consensus                                     !!*!!!!!!!!!!!!!!!!!*!***!!!!!!!!!!!!!!!!!!!!!!!*!**!*!*!*!!!*!*!!*!!!!!!!!!!!!!!!!!!

logo

NTDB id 1140657 DQN24 RS02935 WP 111695365.1  KLSPELAAQIIPLNKQIQRLAVRNGLSQHLRWEMGQQPILHLQLIGHFEKTKTFLTALLASTSQLSVSRLQFMKPEDSPL   160
NTDB id 1345 HI 0437 AAC22096.1               KLSPELAAQIIPLNKQIQRLAARNGLSQHLRWEMGQKPILHLQLTGHFEKTKTFLSALLANSSQLSVSRLQFMKPEDGPL   160
NTDB id 1371 NTHI RS02655 WP 011272048.1      KLSPELAAQIIPLNKQIQRLAARNGLSQHLRWEMGQKPILHLQLTGHFEKTKTFLTALLANSSQLSVSRLQFIKPEDNPL   160
consensus                                     !!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!*!!!!!!!*!!!!!!!!!!!*!!!!**!!!!!!!!!!*!!!!  !!

logo

NTDB id 1140657 DQN24 RS02935 WP 111695365.1  QTEIIFQLEKETK   173
NTDB id 1345 HI 0437 AAC22096.1               QTEIIFQLDKETK   173
NTDB id 1371 NTHI RS02655 WP 011272048.1      QTEIIFQLDKETK   173
consensus                                     !!!!!!!!*!!!!
```

X non conserved

X similar

X ≥ 50% conserved