```
logo
NTDB id 275933 C3V42 RS02815 WP 106013563.1   MQFSSKNHRTLQIGIHRKQGNFDFVWFDELEQPQCYQASINDSDLKNRFLRHLKTQFQGKTFPLQFVASISAHLTWSKVL   80
NTDB id 1369 NTHI RS02665 WP 011272050.1      MQFSLKNYRTLQIGIHRKQGYFDFVWFDDLEQPQSYQIFVNDRDFKNRFLQQLKTQYQGKTFPLQFVASIPAHLTWSKVL   80
NTDB id 1343 HI 0439 AAC22098.1               MQFSLKNYRTLQIGIHRKQSYFDFVWFDDLEQPQSYQIFVNDRYFKNRFLQQLKTQYQGKTFPLQFVASIPAHLTWSKVL   80
consensus                                     !!!!*!!*!!!!!!!!!!!**!!!!!!!*!!!!!*!!***!***!!!!!**!!!!!*!!!!!!!!!!!!!!*!!!!!!!!!

logo
NTDB id 275933 C3V42 RS02815 WP 106013563.1   MLPQLLNSQECHQQCKFVIEKELPIPLEELWFDYISTPLKQGFRLEITAIREASAQTYLQDFQPFKINVLDIVPHSILRA   160
NTDB id 1369 NTHI RS02665 WP 011272050.1      MLPQVLNAQECHQQCKFVIEKELPISLNELWFDYRSTSLKQGFRLDVTAIRKSTAQTYLQDFQPFKINVLDVASNAILRA   160
NTDB id 1343 HI 0439 AAC22098.1               MLPQVLNAQECHQQCKFVIEKELPIFLEELWFDYRSTPLKQGFRLEVTAIRKSSAQTYLQDFQPFNINILDVASNAVLRA   160
consensus                                     !!!!*!!*!!!!!!!!!!!!!!!!!!  !*!!!!!!*!!*!!!!!!!**!!!!***!!!!!!!!!!!!*!!*!!******!!!

logo
NTDB id 275933 C3V42 RS02815 WP 106013563.1   FQYLLNEQVRSENTLFLFQEDDYCLAICERAQQSQILQSHENLTVLYEQFTERFEGKLEQVFVYQIPSSHTPLPEHWHRV   240
NTDB id 1369 NTHI RS02665 WP 011272050.1      FQYLLNEQVRSENTLFLFQEDDYCLAICERSQQSQILQSHENLTALYEQFTERFEGQLEQVFVYQIPSSHTPLPENWQRV   240
NTDB id 1343 HI 0439 AAC22098.1               FQYLLNEQVRSENTLFLFQEDDYCLAICERSQQSQILQSHENLTALYEQFTERFEGQLEQVFVYQIPSSHTPLPENWQRV   240
consensus                                     !!!!!!!!!!!!!!!!!!!
```