```
logo
NTDB id 207782 BSR55 RS02210 WP 100884434.1  MLQFIHQLLHKGIQCISPCLLCGIDQQQAHSLCQDCWNQLPWYKQSIERHEQQILCAHHYDFPIDRIIQTYKYEQQLQYQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    MFKFLNP..QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNP..QYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                    !**!***  ****!**!!!!!!*!******!!!**!!*!!!!*!!*!*!**!**!*!*!!**!!*!!!!**!!!!*!*!!

logo
NTDB id 207782 BSR55 RS02210 WP 100884434.1  NVLAHSLLNLRFSRVQAIVPMPISTERLKERGYNQMLIVAKIMAKQLKIPVWQPVVRQAQHSQKGLSRVERLENIEHQFR  160
NTDB id 1042 HON27 RS01930 WP 000472273.1    TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1     TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                    **!!**!**!*!**!!!!!!!!!!**!!*!!!*!!*!**!***********!!!!!!!!!!!*!***!!!!!!!!*!!*!!!!*!!*

logo
NTDB id 207782 BSR55 RS02210 WP 100884434.1  TIQSEKRKYKKVLIVDDVVTTGSSIHALTQALQRLGCQQISSLCIAAAHPSMK  213
NTDB id 1042 HON27 RS01930 WP 000472273.1    ALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ****!*!*!**!!!*!!!*!!!!!!!!!!*!!!**!!!**!***!*!!!*****
```

X non conserved

X similar

X $\geq$ 50% conserved