```
logo
NTDB id 403811 GK020 RS13655 WP 156189545.1  MPMFKLLAHSRTLLQKLLPCQLCGSAAQQQYRVCADCLQQLPWLKANVCRQETEIQVACHYRYPVDRMIQQFKYEQQLHY   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                     !!!*! ****!*!*!*!!*!!*******!**!**!**!!!!!!****!******!!!!!*!!**!*!!!!!!!!!*!!!

logo
NTDB id 403811 GK020 RS13655 WP 156189545.1  QPLLAGSLCELAYPRVQAIVPMPVATERLAERGYHHTLLLAQVLARHLQLPIWQPVVRRHQHSQKGLSRLERLENLDQQF  160
NTDB id 1071 ABD1 RS15375 WP 000472269.1     QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1042 HON27 RS01930 WP 000472273.1    QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                    !*!!!**!**!**!*!!!!!!!!****!!*!!!****!!!!**!***!**!*!!!!*!***!!!!!!!!!!!!*!!**!!!

logo
NTDB id 403811 GK020 RS13655 WP 156189545.1  QLRPDTGRPYRRVLIVDDVVTTGSSIRALKQQLEQLGCHKVYAVCIASADG...  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1042 HON27 RS01930 WP 000472273.1    VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ****** !*!!!!!!*!!!*!!!!!!*!!*!*!*!!!!****!*!*!*!*****
```

X non conserved

X similar

X ≥ 50% conserved