```
logo

NTDB id 1006874 ABLC01 RS01995 WP 016806074.1  MFNFLNFQHLIQLFSPCSLCELGMREKYSLCKDCWEQLPWLKQTIQRNNHSVLVACNYAYPVNRIIQQFKYEQKLHYQIL   80
NTDB id 1042 HON27 RS01930 WP 000472273.1      MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
consensus                                      !!*!!!*!*!*!!*!!!*!!!*!*!!!!!!!!*!!!!!!!!!!!!!!!!**!!!!!!*!!!!*!!!!!!!!!!!!!!!!*!

logo

NTDB id 1006874 ABLC01 RS01995 WP 016806074.1  LQEILKQLKFPKVQAIVPMPISNQRLIERGFNQSLLLANILSRHLKIPVWQPIQRLNEHSQKGLTRLERFENIEQQFLPH  160
NTDB id 1042 HON27 RS01930 WP 000472273.1      LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL  160
NTDB id 1071 ABD1 RS15375 WP 000472269.1       LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL  160
consensus                                      !*!*!*!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!!!!!!*!***!!!!!!!!*!!!!!!!!!!!!*!!!!!!!!!!!!***

logo

NTDB id 1006874 ABLC01 RS01995 WP 016806074.1  HQEKRRYRRVLIIDDVITTGSSVHALSQALKQLGCTSIHTVCLAATPKS..  209
NTDB id 1042 HON27 RS01930 WP 000472273.1      TQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       TQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                      *!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!**!!!!******
```

X non conserved

X similar

X $\geq$ 50% conserved