```
logo
NTDB id 278703 C7A15 RS03295 WP 002116884.1  MFNFLNFKHLIKLFSPCSLCELDTREKYSLCKDCWEQLPWLKQTIQRNNQSVLVACNYAYPVNRIIQQFKYEQKLHYQIL   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1    MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQTL   80
consensus                                    !!*!!!***!**!*!!!*!!!**!!!!!!!!!*!!!!!!!!!!!!!!!!!*!!!!!!!*!!!!*!!!!!!!!!!!!!!!!!*!

logo
NTDB id 278703 C7A15 RS03295 WP 002116884.1  LQEILKQLKFPKVQAIVPMPISNQRLIERGFNQSLLLANILSRHLKIPVWQPIQRLNEHSQKELTRLERFENIEQQFLPH   160
NTDB id 1042 H0N27 RS01930 WP 000472273.1    LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL   160
NTDB id 1071 ABD1 RS15375 WP 000472269.1     LAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFVAL   160
consensus                                    !*!*!*!!!!!!!!!!!!!!!!!*!!!*!!!!!!!!!!!!!*!***!!!!!!!!*!!!!!!!!!*!*!!!!!!!!!!!!!***

logo
NTDB id 278703 C7A15 RS03295 WP 002116884.1  HQEKRRYRRVLIIDDVITTGSSVHALSQALKQLGCTSIHTACLAATLKS..   209
NTDB id 1042 H0N27 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