```
logo
NTDB id 267011 C2U64 RS03520 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 267011 C2U64 RS03520 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 267011 C2U64 RS03520 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