```
logo
NTDB id 276617 C6I29 RS12925 WP 034669897.1  .MPMFNPSSLLRPISRLLPCPLCSSSLQMQHGVCHDCWEQLPWYRQTVERQELDIHVACGYQYPIDRIIQQFKYEQQLNY   79
NTDB id 1042 H0N27 RS01930 WP 000472273.1    MFKFLNPQYL...FQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNPQYL...FQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                    *****!!**!   ***!*!!*!!**********!**!!!!!!!**!!**!******!!!*!*!!!*!!!!!!!!!!!*!*!

logo
NTDB id 276617 C6I29 RS12925 WP 034669897.1  QQLLTASIQTLRLPKVQAIVPMPISTERLIERGYNQSLVIAKILARELNVPVWQPVIRLAQHSQKGLSRLERLNQIEMQF  159
NTDB id 1042 H0N27 RS01930 WP 000472273.1    QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1071 ABD1 RS15375 WP 000472269.1     QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                    !*!!****!*!**!!!!!!!!!!!!!**!!*!!!*!!!!**!**!*****!**!!!!!!*!!**!!!!!!!!!!!!***!!*!!

logo
NTDB id 276617 C6I29 RS12925 WP 034669897.1  QPILTECRRYRRVLMLDDVVTTGSSLSALAQALTRLGCQHIFAACIAVAETA..  211
NTDB id 1042 H0N27 RS01930 WP 000472273.1    VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    *****! !!!!!!!!**!!!*!!!!!**!!*!!!**!!!**!*!*!*!*!*****
```

X non conserved

X similar

X $\geq$ 50% conserved