```
logo
NTDB id 351512 E3H47 RS12405 WP 005018309.1  .MPMFNPSSLLRPISRLLPCPLCSSSLQMQHGVCHDCWEQLPWYRQTVERQELNIHVACGYQYPIDRIIQQFKYEQQLNY   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 351512 E3H47 RS12405 WP 005018309.1  QQLLTASILTLRLPKVQAIVPMPISTERLIERGYNQSLVIAKILARELNVPVWQPVIRLAQHSQKGLSRLERLNQIEMQF  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 351512 E3H47 RS12405 WP 005018309.1  QPILTERRRYRRVLMLDDVVTTGSSLSALAQALTQLGCQHIFAACIAAAETA..  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