```
logo

NTDB id 1009467 ABJ384 RS01865 WP 349928568.1  MLQLIHPLLHKTMQSLFPCILCGIDQQRHHSVCTDCWQQLPWLKQTIQRQEIEIQAAFHYAYPMDRIIQKFKYEQQLHFQ   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1      MFKFLNPQ..YLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1       MFKFLNPQ..YLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                      !*****!*  ***!*!*!!*!!*!******!*!**!!*!!!!!!!!!!!!!!!!!*!!!!!**!!!!*!!!!!*!!*!

logo

NTDB id 1009467 ABJ384 RS01865 WP 349928568.1  NLLAGSLIGLKLPKIQAIVPMPISHQRLAERGYNQSLVLAKLLAKQLNIPVWQPVARLAQHSQKGLSRIERLDQIDQQFV  160
NTDB id 1042 H0N27 RS01930 WP 000472273.1      TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1       TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                      *!!!**!**!!*!!*!!!!!!!!!*!!!*!!!*!!!!*!!*!!*!!!*!!!!!!!!*!!**!!!!!!!!*!!***!*!!!!

logo

NTDB id 1009467 ABJ384 RS01865 WP 349928568.1  IATKTRVRYRKVLMLDDVVTTGSSLHALKQKLEALGCQKVYAACIASAEI...  210
NTDB id 1042 H0N27 RS01930 WP 000472273.1      ALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       ALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                      **!***!!!*!!**!!!*!!!!!*!!!*!*!**!!!****!*!*!*!*****
```

X non conserved

X similar

X $\geq$ 50% conserved