```
logo

NTDB id 422156 G0028 RS02510 WP 174493382.1  MLQFINQLLYKGVQSISPCLLCGIDRQQQHSLCNDCWEQLPWFKQHTIRHEQNILCAHHYDFPINRIIQTYKYEQQLQYQ   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    MFKFLNPQY..LFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
NTDB id 1071 ABD1 RS15375 WP 000472269.1     MFKFLNPQY..LFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHYQ   78
consensus                                    !**!*!***  **!**!!!!!!*!******!!**!!!!!!!*!!***!**!**!*!*!!**!!!!!!!**!!!!*!*!!

logo

NTDB id 422156 G0028 RS02510 WP 174493382.1  NLLAHSLLNLRLPKVQAIVPMPISTERLKDRGYNQMLLIAKIMARELKVPVWQPVIRAAQHSQKGLSRIERLENIENQFQ  160
NTDB id 1042 HON27 RS01930 WP 000472273.1    TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
NTDB id 1071 ABD1 RS15375 WP 000472269.1     TLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQFV  158
consensus                                    *!!!**!**!**!!!!!!!!!!!!!**!!**!!*!!*!!*!***********!*!!!!!!*!***!!!!!!!!*!!*!!!!*!!*

logo

NTDB id 422156 G0028 RS02510 WP 174493382.1  IIPAEKRKYKKVLIIDDVVTTGSSIHALNQALEKLGCQQIYSACIAVAGTK..  211
NTDB id 1042 HON27 RS01930 WP 000472273.1    ALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ****!*!*!**!!!!!!!*!!!!!!!!!!*!!**!!!**!***!*!*!*****
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X non conserved

X similar

X $\geq$ 50% conserved