```
logo
NTDB id 1039307 ACDJ02 RS02825 WP 004995146.1  MPMFDNIKASMQQLGQKLLPCRLCESSPRLKHYVLCKSCWQQLPWLKQPIHRHELQIMAACYYHYPIDRILQQFKYEQKL   80
NTDB id 1042 HON27 RS01930 WP 000472273.1      ..MFKFLN..PQYLFQLLSPCLLCEIGTR.EKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKL   75
NTDB id 1071 ABD1 RS15375 WP 000472269.1       ..MFKFLN..PQYLFQLLSPCLLCEIGTR.EKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKL   75
consensus                                        !!****  *!*!*!*!*!!*!!!***! **!*!!!*!!*!!!!!!!*!*!*******!!*!*!!!*!!*!!!!!!!!!

logo
NTDB id 1039307 ACDJ02 RS02825 WP 004995146.1  HHQPLLDQLLAQVRHPKVQAIVPMPISDARLIERGYNQSLLLAQQLAKKLNVPIWQPIHRLEQHSQKGLNRLERIQGIEQ  160
NTDB id 1042 HON27 RS01930 WP 000472273.1      HYQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQ  155
NTDB id 1071 ABD1 RS15375 WP 000472269.1       HYQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQ  155
consensus                                      !*!*!!***!*!***!!!!!!!!!!!!!**!!*!!!*!!!!!!!**!*!*!**!*!!!**!!**!!!!!!*!!!!***!!!

logo
NTDB id 1039307 ACDJ02 RS02825 WP 004995146.1  QFVVLEKPKIRYRKVLIIDDVVTTGSSIYALHQALQQLGCQQIYACCIAAATNQNAKT  218
NTDB id 1042 HON27 RS01930 WP 000472273.1      QFVALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY..  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1       QFVALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY..  211
consensus                                      !!!*!*****!!!*!!!!!!!*!!!!!!*!!*!!!*!!!!**!*!*!*!!!*****
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X non conserved

X similar

X ≥ 50% conserved