```
logo
NTDB id 392220 GA636 RS05095 WP 160232300.1  MPMFKLLAHSRTLLQKLLPCQLCGSAAQQQYRVCADCLQQLPWLKANVYRQETEIQVACHYRYPVDRMIQQFKYEQQLHY   80
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFKFL.NPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLHY   77
consensus                                     !!!*! ****!*!*!*!!*!!*******!**!**!**!!!!!!****!******!!!!!*!!**!*!!!!!!!!!*!!!

logo
NTDB id 392220 GA636 RS05095 WP 160232300.1  QSLLAGSLCELTYPRVQAIVPMPVATERLAERGYHHTLLLAQVLARHLQLPIWQPVVRRHQHSQKGLSRLERLENLDQQF  160
NTDB id 1071 ABD1 RS15375 WP 000472269.1     QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
NTDB id 1042 HON27 RS01930 WP 000472273.1    QTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQF  157
consensus                                    !*!!!**!**!**!*!!!!!!!!****!!*!!!****!!!!**!*****!*!!!!*!***!!!!!!!!!!!!*!!**!!!

logo
NTDB id 392220 GA636 RS05095 WP 160232300.1  QLRPDAGRPYRRVLIVDDVVTTGSSIRALKQQLEQLGCHKVYAVCIASADG...  211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     VALTQENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
NTDB id 1042 HON27 RS01930 WP 000472273.1    VALTQEKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY  211
consensus                                    ****** !*!!!!!!*!!!*!!!!!!*!!*!*!*!!!!****!*!*!*!*****
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X non conserved

X similar

X $\geq$ 50% conserved