```
logo
NTDB id 391964 GBN93 RS16045 WP 234964893.1  MPMFSHFLHTAKQLLHKAQPCQLCSIVRADLHSVCLDCWQQLPLHPQTIEKQELSIHVAGQYQYPLDHLIQQFKYEQQLY   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFK.F.LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFK.F.LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
consensus                                      !!* ! *****!*****!!*!!*!******!*!**!!*!!!***!!!*****!**!!**!*!!****!!!!!!!!*!*

logo
NTDB id 391964 GBN93 RS16045 WP 234964893.1  WQPLLSGILQQIRLPKVQAIVPMPISTQRLAERGYNQSMILAKDLAKQLNVPIWQPVIRLHQHSQKGLSRLERLDQIHQQ  160
NTDB id 1042 HON27 RS01930 WP 000472273.1    YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
NTDB id 1071 ABD1 RS15375 WP 000472269.1     YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
consensus                                    *!*!!***!!!***!!!!!!!!!!!!!*!!!*!!!*!!!**!!**!*!!!**!*!!!!*!!**!!!!!!!!!!!***!*!!

logo
NTDB id 391964 GBN93 RS16045 WP 234964893.1  FKIDPQQRQYAKLYRRVLILDDVVTTGSSIQALTLQLEQLGCHKVSAACIAYALK...   215
NTDB id 1042 HON27 RS01930 WP 000472273.1    FVALTQ...EKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     FVALTQ...ENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                    !****!    * **!!!!!!*!!!*!!!!!!*!!***!*!!!!****!*!*!*!*****
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X non conserved

X similar

X $\geq$ 50% conserved