```
logo
NTDB id 391520 GBN67 RS01030 WP 234963397.1  MPMFSHLLHTAKQLLHKAQPCQLCGIVRADLHSVCLDCWQQLPLHPQTIEKKELSIHVAGQYQYPLDHLIQQFKYEQQLH   80
NTDB id 1042 HON27 RS01930 WP 000472273.1    ..MFKF..LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKF..LNPQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
consensus                                      !!**  *****!*****!!*!!*!******!*!**!!*!!!***!!!*****!**!!**!*!!****!!!!!!!!*!!

logo
NTDB id 391520 GBN67 RS01030 WP 234963397.1  WQPLLSGILQQIRLPKVQAIVPMPISTQRLAERGYNQSMILAKDLAKQFNVPIWQPVIRLHQHSQKGLSRLERLDQIHQQ  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 391520 GBN67 RS01030 WP 234963397.1  FKIDPQQRQYAKRYRRVLILDDVVTTGSSIQALTLQLEQLGCHKVSAACIAYALE...   215
NTDB id 1042 HON27 RS01930 WP 000472273.1    FVALTQ...EKRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
NTDB id 1071 ABD1 RS15375 WP 000472269.1     FVALTQ...ENRRYRRVLIIDDVITTGSSIHALSQALKQLGCTSIHASCLAAASSTSY   211
consensus                                    !****!   * *!!!!!!!*!!!*!!!!!!*!!***!*!!!!****!*!*!*!*****
```

X non conserved

X similar

X $\geq$ 50% conserved