```
logo
NTDB id 314735 CDG62 RS04605 WP 087527368.1  MPMFSHLLHTAKQLLHRAQPCQLCGIVRADLHAVCLDCWQQLPLHPQTIEKQALSIHVAGQYQYPLDHLIQQFKYEQQLH   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 314735 CDG62 RS04605 WP 087527368.1  WQPLLSGILQQIRLPKVQAIVPMPISSQRLAERGYNQSMILAKDLAKQLNVPIWQPVIRLHQHSQKGLSRLERLDQIQQQ  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 314735 CDG62 RS04605 WP 087527368.1  FKIDPQQRQYAKRYRRVLILDDVVTTGSSIQALTLELEHLGCHKVSAACIAYAQE...   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