```
logo
NTDB id 238186 CFH90 RS02330 WP 126035174.1  MPMFSHLRHTAKQLLHKAQPCQLCGIVRADLHSVCLDCWQQLPLHPQTVEKQELSIHVAGQYQYPLDHLIQQFKYEQQLH   80
NTDB id 1042 H0N27 RS01930 WP 000472273.1    ..MFKFLN..PQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
NTDB id 1071 ABD1 RS15375 WP 000472269.1     ..MFKFLN..PQYLFQLLSPCLLCEIGTREKYSLCKECWEQLPWLKQTIQRNDQSVLVACHYAYPINRIIQQFKYEQKLH   76
consensus                                     !!**!*  ***!*****!!*!!*!******!*!**!!*!!!***!!******!**!!**!*!!****!!!!!!!!*!!

logo
NTDB id 238186 CFH90 RS02330 WP 126035174.1  WQPLLSGVLQQIRLPKVQAIVPMPISTQRLAERGYNQSMILAKDLAKQFNVPIWQPVIRLHQHSQKGLSRLERLDQIHQQ  160
NTDB id 1042 H0N27 RS01930 WP 000472273.1    YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLSKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
NTDB id 1071 ABD1 RS15375 WP 000472269.1     YQTLLAEVLQQLKFPKVQAIVPMPISKQRLTERGFNQSLLLANLLGKQLKIPVWQPVQRLNEHSQKGLSRLERFENIEQQ  156
consensus                                    *!*!!**!!!!***!!!!!!!!!!!!!*!!!*!!!*!!!**!!**!*!!***!*!!!!*!!**!!!!!!!!!!!***!*!!

logo
NTDB id 238186 CFH90 RS02330 WP 126035174.1  FKIDPQQRQYAKRYRRVLILDDVVTTGSSIQALTLQLEQLGCHKISAACIAYALK...   215
NTDB id 1042 H0N27 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