```
logo
NTDB id 1056 ABD1 RS15860 WP 000076101.1      MSQDELQELSLEQLSAKKKKFNLDKFLQQFNTLDMNNYGSWPISVKITCWIFIFFAVLALGYFIVIQPKLQAIDNAQAQE   80
NTDB id 301314 DPR03 RS01375 WP 171260573.1  MMSEQFDELGQEAVVVPKNKMTVEKFFQQFNTLDPNNYGSWPLAVKITCWVFIVFLVLMLAYFGLIRGKIESITQANAQE   80
NTDB id 1012 ACIAD RS15185 WP 004923719.1    MKLDQLDKTPKESGQPKKKRMNVEKFFQQFNMLDMSNYGSWPLSVKITCWIFIFFLVCALGYFVVIRPQLDDIQTAQAQE   80
consensus                                    !  ******  ! *  *!******!!*!!!!*!!**!!!!!!**!!!!!!*!!*!*!**!*!!**!***** ! *!*!!!

logo
NTDB id 1056 ABD1 RS15860 WP 000076101.1      SNLLNEFREKDSKLRNLQQYQVQLQEMQANFNQQLEQLPKETEIPSLVEDINLTGVNSGLKFKNIRLEDEVKQEIFIEQP  160
NTDB id 301314 DPR03 RS01375 WP 171260573.1  QNLLNEFREKDSKLRNLQQYQQQLQDMEASFNQQLEQLPKETEIPGLVEDINLSGVNAGLKFKNIRLEPEVKQEFFIEQP  160
NTDB id 1012 ACIAD RS15185 WP 004923719.1    KNLLNEFREKQSKLRNLQQYQHQLEEMRASFNQQLTQLPKESEIPGLVEDINMTGVNSGLKFKNIRLENEVKQEFFIEQP  160
consensus                                     !!!!!!!!!*!!!!!!!!!!  !!**! !*!!!!!*!!!!!*!!!*!!!!!!**!!!*!!!!!!!!!!  !!!!!*!!!!!

logo
NTDB id 1056 ABD1 RS15860 WP 000076101.1      IAIEATGDYHAFGAFVSSIAALPRIVTMHDFTVDVSPAKDNKSDIPVLNYSIKAKTYRYVGATDTSEQTGKAPAASAST.  239
NTDB id 301314 DPR03 RS01375 WP 171260573.1  ISIEATGDYHAFGSFVSGIAGLSRIVTLHDFTITGTENKEKKTDIPVVDYTLKAKTYRYVGSANSASPDTTAANGQG...  237
NTDB id 1012 ACIAD RS15185 WP 004923719.1    ISIEATGDYHAFGAFVSGIAALPRIVTLHDFVIEAKQSK.EKTDIPEVDYAVKAKTYRYIGVDEQGNASAVQANTSTTTP  239
consensus                                    !*!!!!!!!!!!!!*!!!*!!*!*!!!!*!!!*****  !* !*!!!***!**!!!!!!!*!  ***  ** ********

logo
NTDB id 1056 ABD1 RS15860 WP 000076101.1      NTTVQPK   246
NTDB id 301314 DPR03 RS01375 WP 171260573.1  ...AQ..   239
NTDB id 1012 ACIAD RS15185 WP 004923719.1    AQGGTP.   245
consensus                                       ***
```

X non conserved

X similar

X ≥ 50% conserved