```
logo
NTDB id 1012 ACIAD RS15185 WP 004923719.1       MKLDQLDKTPKESGQPKKKRMNVEKFFQQFNMLDMSNYGSWPLSVKITCWIFIFFLVCALGYFVVIRPQLDDIQTAQAQE   80
NTDB id 1048990 ACEQ85 RS09545 WP 060464993.1  MSPDELHELSLEQAPSKKKKITLEKFLQQFNTLDMNSYGSWPLSVKITCWIFIFFAVLALGYFVAIQPKLQAIDNAQAQE   80
NTDB id 1056 ABD1 RS15860 WP 000076101.1       MSQDELQELSLEQLSAKKKKFNLDKFLQQFNTLDMNNYGSWPISVKITCWIFIFFAVLALGYFIVIQPKLQAIDNAQAQE   80
consensus                                      !* !*! ****!**  !!!*****!!*!!!!*!!!**!!!!!*!!!!!!!!!!!!!*!*!!!!!**!*!*!**!**!!!!!

logo
NTDB id 1012 ACIAD RS15185 WP 004923719.1       KNLLNEFREKQSKLRNLQQYQHQLEEMRASFNQQLTQLPKESEIPGLVEDINMTGVNSGLKFKNIRLENEVKQEFFIEQP  160
NTDB id 1048990 ACEQ85 RS09545 WP 060464993.1  SNLLNEFREKDSKLRNLQQYQLQLQEMQANFNQQLEQLPKETEIPSLVEDINLTGVNSGLKFKNIRLEDEVKQEIFIEQP  160
NTDB id 1056 ABD1 RS15860 WP 000076101.1       SNLLNEFREKDSKLRNLQQYQVQLQEMQANFNQQLEQLPKETEIPSLVEDINLTGVNSGLKFKNIRLEDEVKQEIFIEQP  160
consensus                                      *!!!!!!!!!*!!!!!!!!!!*!!*!!*!*!!!!!*!!!!!*!!!*!!!!!!*!!!!!!!!!!!!!!!!!*!!!!!*!!!!!

logo
NTDB id 1012 ACIAD RS15185 WP 004923719.1       ISIEATGDYHAFGAFVSGIAALPRIVTLHDFVIEAKQSK.EKTDIPEVDYAVKAKTYRYIGVDEGGNASAVQANTSTITP  239
NTDB id 1048990 ACEQ85 RS09545 WP 060464993.1  ITIEATGDYHAFGAFVSSIAALPRIVTMHDFIVEAAPVKDGKSDIPVLNYSIKAKTYRYMGAVENPEHQEKNASASSVAP  240
NTDB id 1056 ABD1 RS15860 WP 000076101.1       IAIEATGDYHAFGAFVSSIAALPRIVTMHDFTVDVSPAKDNKSDIPVLNYSIKAKTYRYVGATDTSEQTGKAPAASAS..  238
consensus                                      !*!!!!!!!!!!!!!!!!*!!!!!!!!!!*!!!**** * !* !*!!!***!**!!!!!!!*!* ** * ***** *!** *

logo
NTDB id 1012 ACIAD RS15185 WP 004923719.1       AQGGTP....   245
NTDB id 1048990 ACEQ85 RS09545 WP 060464993.1  DSSTANTQPK   250
NTDB id 1056 ABD1 RS15860 WP 000076101.1       ..TNTTVQPK   246
consensus                                        * *  ***
```

X non conserved

X similar

X $\geq$ 50% conserved