```
logo
NTDB id 251141 BS636 RS06935 WP 171266066.1  MMNEQFDELGQEAVVAPKNKMTVEKFFQQFNTLDPNNYGSWPLAVKITCWIFIVFLVLMLAYFGLIRGKIESITQANAQE   80
NTDB id 1056 ABD1 RS15860 WP 000076101.1     MSQDELQELSLEQLSAKKKKFNLDKFLQQFNTLDMNNYGSWPISVKITCWIFIFFAVLALGYFIVIQPKLQAIDNAQAQE   80
NTDB id 1012 ACIAD RS15185 WP 004923719.1    MKLDQLDKTPKESGQPKKKRMNVEKFFQQFNMLDMSNYGSWPLSVKITCWIFIFFLVCALGYFVVIRPQLDDIQTAQAQE   80
consensus                                    ! *******  ! * **!******!!*!!!!*!!**!!!!!!**!!!!!!!!!*!*!**!*!!**!***** ! *!*!!!

logo
NTDB id 251141 BS636 RS06935 WP 171266066.1  QNLLNEFREKDSKLRNLQQYQQQLQDMEASFNQQLEQLPKETEIPGLVEDINLSGVNAGLKFKNIRLEPEVKQEFFIEQP  160
NTDB id 1056 ABD1 RS15860 WP 000076101.1     SNLLNEFREKDSKLRNLQQYQVQLQEMQANFNQQLEQLPKETEIPSLVEDINLTGVNSGLKFKNIRLEDEVKQEIFIEQP  160
NTDB id 1012 ACIAD RS15185 WP 004923719.1    KNLLNEFREKQSKLRNLQQYQHQLEEMRASFNQQLTQLPKESEIPGLVEDINMTGVNSGLKFKNIRLENEVKQEFFIEQP  160
consensus                                     !!!!!!!!!*!!!!!!!!!! !!**! !*!!!!!*!!!!!*!!!*!!!!!!**!!!*!!!!!!!!!! !!!!!*!!!!!

logo
NTDB id 251141 BS636 RS06935 WP 171266066.1  ISIEATGDYHAFGSFVSGIAGLSRIVTLHDFTITGTENKEKKTDIPVIDYTLKAKTYRYVGAQDRSQMDQQSDVDKSGDK  240
NTDB id 1056 ABD1 RS15860 WP 000076101.1     IAIEATGDYHAFGAFVSSIAALPRIVTMHDFTVDVSPAKDNKSDIPVLNYSIKAKTYRYVGATDTSEQTGKAPAASAS..  238
NTDB id 1012 ACIAD RS15185 WP 004923719.1    ISIEATGDYHAFGAFVSGIAALPRIVTLHDFVIEAKQS.KEKTDIPEVDYAVKAKTYRYIGVDEQGNASAVQ....AN..  233
consensus                                    !*!!!!!!!!!!!!*!!!*!!*!*!!!!*!!!*****  ** !*!!!***!**!!!!!!!*!* * ** **       *

logo
NTDB id 251141 BS636 RS06935 WP 171266066.1  KNSKKVSKAKNGGKK   255
NTDB id 1056 ABD1 RS15860 WP 000076101.1     ..T.NTTVQPK....   246
NTDB id 1012 ACIAD RS15185 WP 004923719.1    ..T.STTTPAQGGTP   245
consensus                                      *  **   ***
```

X non conserved

X similar

X ≥ 50% conserved