```
logo
NTDB id 1122403 EL239 RS05825 WP 232001743.1  MRQKIASITQRFRIPAVCLLCGSYHRGCYAVCRTCEGLLTRVDHPCRYCGLPLIDESFAVCGHCIKQRPAFDRVLTAYRF   80
NTDB id 1270 LPP RS11540 WP 015961431.1       MLQKFLSITQNLRLPSICTLCNQFHKSQLAVCSNCMEFMKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF   80
NTDB id 1247 GCO85 RS11680 WP 015961431.1     ......................................MKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF   42
consensus                                     * **  ****  *** ** **  ***   ***  *   *******!*!!**!!*!****!!!*!***!!*!!****!!!!

logo
NTDB id 1122403 EL239 RS05825 WP 232001743.1  EGALRTLLHDYKYGQTLAIRTFLAKLMLDALPRQPYKPDCLIPVPLHPKRLRQRGFNQAVELAKLLAGQLNIPCYRTLCR  160
NTDB id 1270 LPP RS11540 WP 015961431.1       EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  160
NTDB id 1247 GCO85 RS11680 WP 015961431.1     EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  122
consensus                                     !**!!*!*!**!!***!****!!**!*!*!!!****!!!!!!!!!*!!!!!**!!!!!!**!**!!!*!!!!!*****!*

logo
NTDB id 1122403 EL239 RS05825 WP 232001743.1  KTINTPAQVGLSSQQRRKNLRHSFTIGPHNFQHVTLVDDLLTTGSTASELARQFKKQGVRRVDVWCCARTAERSCYPR  238
NTDB id 1270 LPP RS11540 WP 015961431.1       KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN....  234
NTDB id 1247 GCO85 RS11680 WP 015961431.1     KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN....  196
consensus                                     !*!!!**!**!***!!!!!!!!*!***!****!!**!!!!!!!!!!!!*!*!***!**!!*!!!**!!!!*****
```

X non conserved

X similar

X $\geq$ 50% conserved