```
logo
NTDB id 1122607 EL220 RS02705 WP 414092931.1  MRQKIWSLAQSLRIYSICILCNQYHNNKMAVCSHCLELMPLLGPGCRYCAYPLPDTHFLVCGNCIKKPPSFDRAFIKYTF   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 1122607 EL220 RS02705 WP 414092931.1  EEPLRSLLHQFKYHEGLYLTSFLSYLIFKSLSNQADMPQCLIPVPMHSQRIKQRGFNQAAVLARSLAKKLQLPCNLFSCQ  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 1122607 EL220 RS02705 WP 414092931.1  KTLNTKPQAGLDEKQRRKNLRSAFITQKLPYQHVAIIDDLLTTGNTANELARTLKKSGVKQVDVWCCARTIEKQ  234
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