```
logo
NTDB id 393336 GCS73 RS03290 WP 012978820.1  MRQKIWSLTQSLRIYSICILCKQYHNDKMAVCSHCIELMPLLGPGCHHCAYPLPDTQFLVCGHCIKNPPSFDRAFIKYTF   80
NTDB id 1270 LPP RS11540 WP 015961431.1      MLQKFLSITQNLRLPSICTLCNQFHKSQLAVCSNCMEFMKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF   80
NTDB id 1247 GC085 RS11680 WP 015961431.1    ......................................MKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF   42
consensus                                    * **  **** *** *** ** ***   ***** *** !**!!**!**!!!!!*!***!!!!*!****!!!!*!*!*!*!

logo
NTDB id 393336 GCS73 RS03290 WP 012978820.1  EEPLRTLLHQFKYHNGLYLTSFLSYLIFKSLSSQVEMPQCLIPVPMHPQRIKRRGFNQAAILARSLAKKFKLPCDLFSCQ  160
NTDB id 1270 LPP RS11540 WP 015961431.1      EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  160
NTDB id 1247 GC085 RS11680 WP 015961431.1    EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  122
consensus                                    !!!!!*!*!!!!!!!!!!!*!!!**!****!******!*!!!!!!!!!*!*!!!!!!!!!*!*!*!!*****!*!!**!!

logo
NTDB id 393336 GCS73 RS03290 WP 012978820.1  KTLNTKPQAGLDEKQRKINLRSAFVTHQLPYQHVAIIDDLLTTGNTANELAYTLKKSGVKQVDVWCCARTIEKH  234
NTDB id 1270 LPP RS11540 WP 015961431.1      KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN  234
NTDB id 1247 GC085 RS11680 WP 015961431.1    KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN  196
consensus                                    !**!!**!!*!!**!!**!!!*!!******!*!!*!*!!!!!!!*!!!!!*!!!!**!!!*!!**!!!!***!*
```

X non conserved

X similar

X $\geq$ 50% conserved