```
logo
NTDB id 1113916 LHA RS03870 WP 407927331.1  MRYKISSITHMLRFPSACVICSQYHRENCAVCKNCMTLLQPLGHACQYCAFPLPDKDFLVCGQCAKVKPAFDKAFVMYQFEEPLR   85
NTDB id 1270 LPP RS11540 WP 015961431.1     MLQKFLSITQNLRLPSICTLCNQFHKSQLAVCSNCMEFMKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRFEEPLR   85
NTDB id 1247 GCO85 RS11680 WP 015961431.1   ....................................MKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRFEEPLR   47
consensus                                   *  *  ***  ** ** * ** **** * *** ***  ***!!**!!!!!*!!*!***!!!!!!****!*!!*!***!*!!!!!!
```

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logo
NTDB id 1113916 LHA RS03870 WP 407927331.1  TLLHQYKYNGALYLRKFLVQLILSALPTEKTKTQCLIPVPLHPQRLRERGFNQAAEFSKHLAKTLKIPCELNLCKKILHTPAQVS   170
NTDB id 1270 LPP RS11540 WP 015961431.1     SLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQKIINTASQAN   170
NTDB id 1247 GCO85 RS11680 WP 015961431.1   SLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQKIINTASQAN   132
consensus                                   *!*!!*!!***!!!**!!*!!*!*!!!****!**!!!!!!*!!*!!**!!!!!!*****!!**!*!!**!**!*!!**!**!**
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logo
NTDB id 1113916 LHA RS03870 WP 407927331.1  LNGKARRNNLRHAFEIKNNAYRHITLVDDLLTTGSTANELAKLFKRQGVERVDIWCCARTC...   231
NTDB id 1270 LPP RS11540 WP 015961431.1     LDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN   234
NTDB id 1247 GCO85 RS11680 WP 015961431.1   LDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN   196
consensus                                   !*!**!!*!!!!!!******!*!***!!!!!!!!!!!!!!*!***!**!!*!!!!*!!!!*****
```

X non conserved

X similar

X ≥50% conserved