```
logo
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```
NTDB id 372953 FOG18 RS11790 WP 244946889.1  MLQIFASITQRLRFPTVCLLCKQYQRDAQFICKDCLCLFKPIKYGCKYCACPLPDAHFPVCGACSTKQPDFDRAIIAYPY  80
NTDB id 1270 LPP RS11540 WP 015961431.1      MLQKFLSITQNLRLPSICTLCNQFHKSQLAVCSNCMEFMKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF  80
NTDB id 1247 GCO85 RS11680 WP 015961431.1    ......................................MKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF  42
consensus                                    *** * **** ** **** ** ** *    **  **  *!*****!*!!!*!!*!****!!!*!**!*!*!!*!*!!!**
```

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logo
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```
NTDB id 372953 FOG18 RS11790 WP 244946889.1  EEPLRTLIHQFKYQEKLYLVSFLSQLMLPFLKECDEIPQCLIPVPMHPQRLKQRGFNQAAVLTKALAKKMHLPYQLQGCQ  160
NTDB id 1270 LPP RS11540 WP 015961431.1      EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  160
NTDB id 1247 GCO85 RS11680 WP 015961431.1    EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ  122
consensus                                    !!!!!*!!!!!!!***!!!*!!!*!!*!**!******!*!!!!!!!!!*!!!*!!!!!!!!!!**!!*****!!*!**!!
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logo
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```
NTDB id 372953 FOG18 RS11790 WP 244946889.1  KIINTKPQASLDKKQRAKNLRQAFTINTLPYQHIALVDDLLTTGNTANELARAFKKKGVLRVDVWCCARAVLA.  233
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 ≥ 50% conserved