```
logo
NTDB id 76072 EN18 RS10330 WP 000902745.1     MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 1172 A1552VC RS12120 WP 000902745.1  MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 1405 DSB67 RS13995 WP 010643496.1    MLYHINLLAWRENQREEHRRRFIGLILLGAIAALGVQWGIGKYYQYQQDQQQQRLDYLAHYISQLDTRIEAMKIAEQEHS   80
consensus                                    !!***!!!*!!***!!*!*!!!*!!**!!***!***!***!*!***!***!!*!**!!***!**!!**!***!!!!*!!*

logo
NTDB id 76072 EN18 RS10330 WP 000902745.1     ALLTRLTVVEQLQQKRNKTTEFMNQMPNLIPEGVYVDKIKMNGHQIEITGISDSTARLATMLDNLEKSDKLTDVEMHEIV   160
NTDB id 1172 A1552VC RS12120 WP 000902745.1  ALLTRLTVVEQLQQKRNKTTEFMNQMPNLIPEGVYVDKIKMNGHQIEITGISDSTARLATMLDNLEKSDKLTDVEMHEIV   160
NTDB id 1405 DSB67 RS13995 WP 010643496.1    KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV   160
consensus                                    **!*!!**!!*!!**!!!!!!!!!*!!**!!!!!!!!!!!!!!***!!!*!!!!!!*!!!!!!!!!*!*!*!!*!!*!!*!!

logo
NTDB id 76072 EN18 RS10330 WP 000902745.1     SGNKRFGKQFQSFKVSFQILTPASNPQAGGAHNG   194
NTDB id 1172 A1552VC RS12120 WP 000902745.1  SGNKRFGKQFQSFKVSFQILTPASNPQAGGAHNG   194
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASI...KGGKQHG   191
consensus                                    *!**!!!!*!!*!!!!!***********!!***!
```

X non conserved

X similar

X $\geq$ 50% conserved