```
logo

NTDB id 121954 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 121954 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 121954 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                                    *!**!!!!*!!*!!!!!***********!!***!
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X non conserved

X similar

X $\geq$ 50% conserved