```
logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 240646 BSG32 RS00170 WP 021449760.1  MLYRINLLPWRENQREEHRRRFVGLVVLGVMVALGIQWGVSKYYEYQQDLQQERLDYLTTYISDLDKRIEAMKIAELEHS   80
NTDB id 1405 DSB67 RS13995 WP 010643496.1    MLYHINLLAWRENQREEHRRRFIGLILLGAIAALGVQWGIGKYYQYQQDQQQQRLDYLAHYISQLDTRIEAMKIAEQEHS   80
consensus                                    !!**!!!*!!***!!*!!!*!!**!!***!***!*****!* *!***!!*!**!!  *!* !! *!***!!!! !!*

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   ALLTRLTVVEQLQQKRNKTTEFMNQMPNLIPEGVYVDKIKMNGHQIEITGISDSTARLATMLDNLEKSDKLTDVEMHEIV   160
NTDB id 240646 BSG32 RS00170 WP 021449760.1  KILERLKVVEALQNGRNKTTDFMNLMPAVIPEGVYVDKIKMNDLEIEISGISDSTPRLATMLDNMERSAKLLDVEMHSIV   160
NTDB id 1405 DSB67 RS13995 WP 010643496.1    KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV   160
consensus                                    **!*!!**!!*!!**!!!!!*!!!*!! *!!!!!!!!!!!!!* *!!!*!!!!!! !!!!!!!!*!*!*!!*!!*!!*!!

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   SGNKRFGKQFQSFKVSFQILTPASNPQAGGAHNG   194
NTDB id 240646 BSG32 RS00170 WP 021449760.1  HGKARFGKEFQTFKVSFMFNTVEE..QKEGKQHG   192
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDAS...IKGGKQHG   191
consensus                                    *!**!!!!*!!*!!!!!** *      ***!***!
```

X non conserved

X similar

X $\geq$ 50% conserved