```
logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 975468 AADH43 RS01820 WP 015297396.1 MLYRINLLPWRENQREEHRRRFVGLVVLGVMVALGIQWGVGKYYEYQQDLQQERLDYLTTYISDLDKRIEAMKIAELEHS   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 975468 AADH43 RS01820 WP 015297396.1 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 975468 AADH43 RS01820 WP 015297396.1 HGKARFGKEFQTFKVSFMFNTVEEK..KEGKQHG   192
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASI...KGGKQHG   191
consensus                                    *!**!!!!*!!*!!!!!** *      **!***!
```

X non conserved

X similar

X $\geq$ 50% conserved