```
logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1     MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 1024024 AB2S62 RS01310 WP 367987937.1  MLHSINLLPWREELRERHKKRFLSLLILGILVGIGIQWGIGLYISNETEKQQSRLSYLNSYIAELDRRIAALKQVEQDHA   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 1024024 AB2S62 RS01310 WP 367987937.1  ALLTRLAVVEKLQKQRNKTTDFMNLMPGLIPEGVYVDKIKMGGNEIEIAGISDTTSRLATMLDNLEKSSKLNEVEMHSIV  160
NTDB id 1405 DSB67 RS13995 WP 010643496.1      KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV  160
consensus                                      **!*!! *!! !!* !!!!!*!!!*!! *!!!!!!!!!!!!** *!!!*!!!!*!*!!!!!!!!*!*! !! *!*!!*!!

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1     SGNKRFGKQFQSFKVSFQILTPA..SNPQAGGAHNG   194
NTDB id 1024024 AB2S62 RS01310 WP 367987937.1  HGKVRFGKKFQTFKVSFNFMTDELPQTPANGGKRNG   196
NTDB id 1405 DSB67 RS13995 WP 010643496.1      HGKARFGKEFQTFKVSFMFKDASIK.....GGKQHG   191
consensus                                      *!* !!!! !!*!!!!!****  *   *  !!* *!
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X non conserved

X similar

X ≥ 50% conserved