```
logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 39659 VVMO6 RS01725 WP 011079353.1   MQHKINLLPWREHKREEHKRRFVSLVALGALIAVGIQWGIGQYFSYQQRLQQERLDYLTHYIAQLDQRIQAMNIAKEEHS   80
NTDB id 1405 DSB67 RS13995 WP 010643496.1    MLYHINLLAWRENQREEHRRFIGLILLGAIAALGVQWGIGKYYQYQQDQQQQRLDYLAHYISQLDTRIEAMKIAEQEHS   80
consensus                                    !****!!!*!!* *!!*!*!!!**!* !!***!***!***! !* *!* *!!*!**!! **! *!!**! ***!!**!!*

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   ALLTRLTVVEQLQQKRNKTTEFMNQMPNLIPEGVYVDKIKMNGHQIEITGISDSTARLATMLDNLEKSDKLTDVEMHEIV   160
NTDB id 39659 VVMO6 RS01725 WP 011079353.1   QILARLKVVESLQAERNKTTAFMNLMTELIPEGVYVDKIKMNTNDVELAGISDSTARLATMLDNLERSTFLKEVEMHSIV   160
NTDB id 1405 DSB67 RS13995 WP 010643496.1    KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV   160
consensus                                     *! !!**!! !!* !!!!!*!!!*!* *!!!!!!!!!!!!!  **!**!!!!!!*!!!!!!!!*!*! *! *!*!!*!!

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   SGNKRFGKQFQSFKVSFQILTPASNPQA...GGAHNG   194
NTDB id 39659 VVMO6 RS01725 WP 011079353.1   HDKKRFGKSFQTFTVSFVFDEHADSANTLAKEGNH.G   196
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASIKGGK.....QH.G   191
consensus                                    ****!!!! !!*!*!!!** * *  **      **! !
```

X non conserved

X similar

X $\geq$ 50% conserved