```
logo

NTDB id 1172 A1552VC RS12120 WP 000902745.1   MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 373033 VV7356 RS21615 WP 011079353.1 MQHKINLLPWREHKREEHKRRFVSLVALGALIAVGIQWGIGQYFSYQQRLQQERLDYLTHYIAQLDQRIQAMNIAKEEHS   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 373033 VV7356 RS21615 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   SGNKRFGKQFQSFKVSFQILTPASNPGA...GGAHNG   194
NTDB id 373033 VV7356 RS21615 WP 011079353.1 HDKKRFGKSFQTFTVSFVFDEHADSANTLAKEGNH.G   196
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASIKGGK.....QH.G   191
consensus                                    ****!!!! !!*!*!!!** * *  **      **! !
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X non conserved

X similar

X $\geq$ 50% conserved