```
logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   MLHKVNLLPWRDARREAHKRRFLGLVTLGVLLAVLMQFAAGEYLGGQMALQQERIGYLQQHIFSLDQQIAKLKIAEEEHK   80
NTDB id 1054714 BCU28 RS15530 WP 010435288.1 MLHQINLLSWRDEIRAQHKKRFIHLVILGVIIALAGQWAVGHYFHDQQAKQQARLSYLDKYIAVLDRQIQSLKVAEQEHK   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 1054714 BCU28 RS15530 WP 010435288.1 AILTRLAVVESLQFGRNKTTDFMNLMPELIPEGVYVDKIKMNGQEIEMSGISDSTARLATMLDNLERSDSLKEVEMHSIV   160
NTDB id 1405 DSB67 RS13995 WP 010643496.1    KILERLKTVEGLQNGRNKTTEFMNLMPSVIPEGVYVDKIKMNDYEIEISGISDSTSRLATMLDNMERSAKLLDVDMHSIV   160
consensus                                    **!*!! *!! !!*!!!!!*!!!*!! *!!!!!!!!!!!!!* *!!*!!!!!!*!!!!!!!!*!*!***! *!*!!*!!

logo
NTDB id 1172 A1552VC RS12120 WP 000902745.1   SGNKRFGKQFQSFKVSFQILTPASNPQ.A.........GGAHNG   194
NTDB id 1054714 BCU28 RS15530 WP 010435288.1 HNRKRFNKEFQSFKVSFVFSPVSLENTTAEGTTANIKEKGANHG   204
NTDB id 1405 DSB67 RS13995 WP 010643496.1    HGKARFGKEFQTFKVSFMFKDASIKGGKQHG............   191
consensus                                    ****!!*!*!!*!!!!!**   **     * *         **  *
```

X non conserved

X similar

X $\geq$ 50% conserved