```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1094554 ACLBLI RS01225 WP 414829372.1 MASRGVNKVILVGNLGNDPEVRYMPNGNAVANLSLATSESWKDQQ.GQVQERTEWHRLTMYRRLAEVAGEYLKKGSQIYV   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!** !** !*!!!!********!!*!!*!!*!!* *!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGA.....P..............YEEGYGQSQEAYQRPA  133
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGA.....P..............YDEGYGQSQEAYQRPA  133
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNN.NF....  149
NTDB id 1094554 ACLBLI RS01225 WP 414829372.1 EGKLQTRKWQDQQGQDRYTTEIIVDQ....MQMLGGREGGGSNAGGGG.GYQQRP...QGNQGGYSNQGGGQQGGYGNQG  151
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGG.MNV..................PAQQGSWGQPQ  142
consensus                                     !!****!!*!** !**!********     **!!**!  *** **  *                * * ***** **  **

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        QQSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       QQSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      .NGGNA.....TRPQPA....QKPAAQAEPPMDNFDDDIPF   180
NTDB id 1094554 ACLBLI RS01225 WP 414829372.1 GPQGNP.....QGGRPQSGNQQPKQPPMAEPDFDFDDDIPF   187
NTDB id 1166 A1552VC RS00795 WP 000168289.1   QPAKQ......HQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                     ** ***      * ** * *  **   *** !****!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved