logo



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NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1035587 ABE954 RS17175 WP 371025626.1   MASKGVNKVILIGHLGQDPEIRYMPAGGAVANLTLATSESWRDKQSGEMREKTEWHRVVIFGKLAEVAGEYLKKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                           **!!!!**! !!*!!**!*!!*!*!!*****!!!!*!*!* *!***!*!!!!*******!!*!!*!!*!!* !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYA....QNNNF   149
NTDB id 1035587 ABE954 RS17175 WP 371025626.1   EGSLQTRKWQDQSGQDRYTTEVVVN.IGGSMQMLGGNGGNQ......AGSR...................K....PA...   127
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMN...VPA....QQGSWGQPQQ....PAKQH   148
consensus                                       !!****!!*!** !**!*******      **!!***  * *  *   ! *           * **  * *    **
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NTDB id 1103 NMB RS07590 WP 002212976.1         PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1        PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1035587 ABE954 RS17175 WP 371025626.1   .....RQPQQ.SQQAPQNEPPMDWDDQGIPF   152
NTDB id 1166 A1552VC RS00795 WP 000168289.1     QPMQQSAPQQ.YSQPQYNEPPMDFDD.DIPF   177
consensus                                           * *!*   ***  ***!**  !**!!!
```

X non conserved
X similar
X $\geq 50\%$ conserved