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logo
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```
NTDB id 1131 NGFG RS05740 WP 003695064.1    ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 177985 A5892 RS19380 WP 064124179.1 .MARGVNKVILIGNLGQDPEVRFLPSGNPVCNLRLATTDTWNDRQSGQRQERTEWHTVVLFNKLAEIAQQYVKKGSRLYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                   **!!!!**!*!!*!!**!**!*!**!*****!!***!*!* *!** !*!!!! *******!!*!**!**!!* *!*
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1    EGRLQSRKYQGKDGIERTAYDIVANEMKMLGGRNEN....SGG..............A...PYDEGYGQSQEA....YQR   131
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVANEMKMLGGRNEN....SGG..............A...PYEEGYGQSQEA....YQR   131
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQGDVLQMLDSRSSGGD......FG.....GNQGSGWNQAPAQTNYNQGGYSDNYAQNN   147
NTDB id 177985 A5892 RS19380 WP 064124179.1 EGRLQTRKWQGQDGQDRYSTEIVANDMQMLDSRAGDFNPNNGGGQGGYQPRGNQGGGAPQDAPQGGYGGGQDYGQ.AARA   158
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRA.....QQGGMPAQ....GGMNVPAQQGSWGQPQQP.......   144
consensus                                   !!****!!*!***!**!*********** *       **  *          **   * * **** **       *
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1    PAQQSRQPAPDAPSHPQEA...PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     PAQQSRQPASDAPSHPQEA...PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NFNGGNATRPQPAQKPA.............AQAEPPMDNFDDDIPF   180
NTDB id 177985 A5892 RS19380 WP 064124179.1 PAHGGGQPRPAPNNAPQQGGYGAPQQNNNGAPNPGSFDDFDDEIPF   204
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....AKQHQPMQQSAPQ..QYSQPQY.......NEPPMDFDDDIPF   177
consensus                                   *** **** *   * !* *          *  ** ******!!*!!!
```

X non conserved
X similar
X ≥ 50% conserved