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logo
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NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 210966 BUQ72 RS20345 WP 079218202.1   ..MASVNKVIIVGNLGRDPETRYMPNGEAVTNIAVATTESWKDKNSGEKKELTEWHRITFYRKLAEIAGQYLKKGSQIYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                        **!!!!**!*!!*!!**!*!!*!*!! ****!!*!*!*!**!***!*!!!!*********!!*!!*!!*!!* *!*
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logo
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NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMNVPAQQGSWGQP..............QQPA   145
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGY.GQSQE.AYQ.RPAQQSRQPA....PDAPS   145
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGY.GQSQE.AYQ.RPAQQSRQPA....SDAPS   145
NTDB id 210966 BUQ72 RS20345 WP 079218202.1   EGRLQTRKWQDKEGVERYTTEITA....DTMQMLGSRQGGGGGGGSMDEGSYGGGGGGGYGGGAPRQNAGGGGG......G   148
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD........FGGNQGSGWNQAPAQTNYNQGGYSDNYAQN   146
consensus                                     !!****!!*!***!**!*******       **!!**!* ****** ***  *  *  ***  ****  * *       **
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logo
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NTDB id 1166 A1552VC RS00795 WP 000168289.1   KQHQP..MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1131 NGFG RS05740 WP 003695064.1      HPQEAPAAPR.....RQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1       HPQEAPAAPR.....RQPVPAAAPVEDIDDDIPF   174
NTDB id 210966 BUQ72 RS20345 WP 079218202.1   GMGGGGGAPRQQPAQRPASN......FEDDDIPF   176
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                        * ****  *  **   **   **** !!!!!!
```

X non conserved
X similar
X ≥ 50% conserved