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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
NTDB id 205799 BO996 RS04205 WP 071954098.1 ..MASVNKVIIVGNLGRDPEMRTFPSGDQVANVTIATTDRWRDKNTGENKEATEWHRVVFNGRLAEIVGQYLRKGSQVYV    78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                      **!!!!**!*!!*!!**!**!*!**!**!***!!***!*!***!***!*!!!!*********!!**!*!!*!!* !!*

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYDEGYGQSQEAYQRPAQ..QSRQP   139
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYEEGYGQSQEAYQRPAQ..QSRQP   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM..NV.PAQ....QGS..WGQPQ.......   142
NTDB id 205799 BO996 RS04205 WP 071954098.1 EGSLRTRKWTDQASGQERYATEIRA....DSMQMLGSRQGGGG..GQQGGGYGD....EGYGDSG..YDAPPQQQQQRRP   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQ.NGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGG..YSDNYAQNNNFNG   151
consensus                                   !!*****!!****  !**!*******      **!!**!* ***  ** *     **  ******   *  * *  * ***

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     APDAPSHPQ......EAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     ASDAPSHPQ......EAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....QPAKQHQPMQQSAPQQYSQ..PQYNEPPMDFDDDIPF   177
NTDB id 205799 BO996 RS04205 WP 071954098.1 APMAAPAPRQAPAQRPAPAPMAPPPQRAASGFDDMDDDIPF   187
NTDB id 1390 A4U84 RS00055 WP 021115958.1   GNAT............RPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                   *  *   **       *!*   * *  ** *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved