```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 375092 FPL17 RS15150 WP 005297510.1 ..MRGVNKVILVGTLGKDPETKTFPNGGSLTQFSIATSESWMDKSTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!!*!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP..Y........EEGYGQSQEAYQRPAQQSR.....   137
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP..Y........DEGYGQSQEAYQRPAQQSR.....   137
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP.................AQGGMNVPAQQGSWGQPQ   142
NTDB id 375092 FPL17 RS15150 WP 005297510.1 EGSLRTRQWTDQNGQERYSTEIRG....DQMQMLDSRQQGEQSGGDFNQPRFNNNNNQGGGYQNTGYNNNNNQGGYGQGG   154
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGN...........QGSGWNQAPAQTNYNQGGYSDNY   143
consensus                                   !!*****!***** !**!*******      **!!**!* * * *             * *  * *** ***!* *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      QPASDAPS.....HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     QPAPDAPS.....HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  QPAKQ.......HQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 375092 FPL17 RS15150 WP 005297510.1  GNQGNYAGSPQAGNGFNTPKAAPQPAA...AAPADLDDDLPF   193
NTDB id 1390 A4U84 RS00055 WP 021115958.1    AQNNNFNGG.....NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                    *** *          *   ***  *    * ******!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved