```
logo
NTDB id 113 BSU 40900 NP 391970.1                 ....MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQS.GE..READFINCVTWRRQAENVANFLKKGSLAGV   73
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         ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFRRQAEVAGQYLRKGSKVYV   78
NTDB id 195367 EnteroDNA1 RS15580 WP 029740211.1 MASRGVNKVILVGNLGQDPEVRYMPSGGAVANITLATSESWRDKATGEMKEQTEWHRVVLFGKLAEVAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1      MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                             *!*!***! !* !!**!**!*! !!** **!***** **  !*  * **** * *****!!*****!*!!* ***

logo
NTDB id 113 BSU 40900 NP 391970.1                 DGRLQTRNYENQQGQRVFVTEVQ....AESVQFLEPKNGGGSGSGGYNEGN.....SGGGQYFGGGQNDNPFGGNQNNQR  144
NTDB id 1103 NMB RS07590 WP 002212976.1           EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGA.PYEEGYGQSQEAYQRPA.........QQSRQPASD  142
NTDB id 1131 NGFG RS05740 WP 003695064.1          EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGA.PYDEGYGQSQEAYQRPA.........QQSRQPAPD  142
NTDB id 1390 A4U84 RS00055 WP 021115958.1         EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSG.GDF.GGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG  152
NTDB id 195367 EnteroDNA1 RS15580 WP 029740211.1 EGQLRTRKWTDQAGAEKYTTEVVVN.VGGTMQMLGGRQG..GGA.PA..GGG..QSQQQGGW...GQPQQPQGGNQFSGG  149
NTDB id 1166 A1552VC RS00795 WP 000168289.1      EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQ.GGM.PA..QGGMNVPAQQGSW...GQPQQPAKQHQPM..  151
consensus                                        *!****!*** * ! *** ****    *  ***!****   **  *   * *  *   *  *   *    *     *

logo
NTDB id 113 BSU 40900 NP 391970.1                 ...RNQ.GNSFNDDPFANDGKPIDISDDDLPF   172
NTDB id 1103 NMB RS07590 WP 002212976.1           APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1          APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1         NATRPQ....PAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 195367 EnteroDNA1 RS15580 WP 029740211.1  AQSRPQ....QQSAPAPSNEPPMD.FDDDIPF   176
NTDB id 1166 A1552VC RS00795 WP 000168289.1       QQSAPQ....QY.SQPQYNEPPMD.FDDDIPF   177
consensus                                           * *!               !**  !!!*!!
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X non conserved

X similar

X ≥ 50% conserved