```
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 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 327510 EHF44 RS03795 WP 124682517.1 ..MASVNKVILVGNLGADPETRYMPSGDAVTNIRLATTDRYKDKQSGEFKEATEWHRIAFFGKLAEIAGQYLRKGSSVYI   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!*!! !!**!*!!*!*!! ****!!*****!* *!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYEEGYGQ...SQEAYQRPAQQSRQPA.........  140
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYDEGYGQ...SQEAYQRPAQQSRQPA.........  140
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGD.FGGNQGSG.......WNQAPAQTNYNQGGYSDNYAQN  146
NTDB id 327510 EHF44 RS03795 WP 124682517.1 EGRIRTRKWQDQSGQDKYSTEIVADQ....MQMLGARQGGGG.GGDEGGYGGGGGGGGGYSRESQGGSGGAGYGGGRGQG  153
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMN..........VPAQQGSWG........QP  141
consensus                                   !!****!!*!** !***********      **!!**!* *** ** * !**        ******!*  * *       *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      .SDA.PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     .PDA.PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNFNG..GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 327510 EHF44 RS03795 WP 124682517.1  GGGAGGSG...GQQGGQRRQQQSSNGFEDMDDDIPF   186
NTDB id 1166 A1552VC RS00795 WP 000168289.1  QQPAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                       *  * *  **** **   **  *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved