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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 392114 GA660 RS01120 WP 005180729.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDAWTDKTTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYG.............QSQEAYQRPAQQSRQP   139
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYG.............QSQEAYQRPAQQSRQP   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMN...........VPAQQGSWGQPQQP....   144
NTDB id 392114 GA660 RS01120 WP 005180729.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSGRQQGE.QGDNGFSQPRFNNNQGGGYSNNNQGGYAPQAQGGFNN   153
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSG..W..NQAPAQTNYNQGGYSDNYAQ...N   146
consensus                                   !!*****!***** !**!*******       **!!***  ***  * * * *             **!***  ****  *

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ASDAPSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     APDAPSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ...AKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 392114 GA660 RS01120 WP 005180729.1 .NNAGGGYGNQGGYQQPKPAPAATPAPADLDDDLPF   188
NTDB id 1390 A4U84 RS00055 WP 021115958.1   .NNFNGGNA.TRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                     *  *     ***   * * *  *****!!!*!!
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X non conserved

X similar

X ≥ 50% conserved