```
logo
                                                  ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1103 NMB RS07590 WP 002212976.1
NTDB id 1131 NGFG RS05740 WP 003695064.1          ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 32327 AFE RS07340 WP 012607100.1          ..MAGVNKVILLGHLGRDPEMRYMPDGTAVANMTIATSESYKDR.DGNRQERTEWHRVALFARTAEVAGEYLRKGSQAYI    77
NTDB id 1390 A4U84 RS00055 WP 021115958.1         ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
NTDB id 1166 A1552VC RS00795 WP 000168289.1       MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
consensus                                             **!!!!**! !!*!!**!*!!*!*!!**!***!!!!***!*  !** !*!!!!********!!*!!*!!*!!* *!*

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NTDB id 1103 NMB RS07590 WP 002212976.1           EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE.........AYQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1          EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE.........AYQRPAQQSRQPAPDA   143
NTDB id 32327 AFE RS07340 WP 012607100.1          EGRIRTRKWTDKDGQERYSTEIVG....DQLQLIGGRRDTPE.SGS.......................PAPQH   123
NTDB id 1390 A4U84 RS00055 WP 021115958.1         EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYA....QNNNF   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1       EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPAQ....QGSWGQPQQ....PAKQH   148
consensus                                         !!****!!*****!**!*******      ******! ** *  *   * *  *          **  * *    ** *

logo

NTDB id 1103 NMB RS07590 WP 002212976.1           PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1131 NGFG RS05740 WP 003695064.1          PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 32327 AFE RS07340 WP 012607100.1          GGGG.........GGDMGAAPMEEFDDDIPFITKAETICQPRWRLPRVRF   164
NTDB id 1390 A4U84 RS00055 WP 021115958.1         NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.................   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1       Q..PMQQSAPQQYSQPQYNEPPMDFDDDIPF.................   177
consensus                                            *   **   *    ***!****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved