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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 295990 DM292 RS15215 WP 063544073.1 .MARGVNKVILIGNVGGDPETRYMPNGNAVTNITLATTDSWKDKQTGQLQERTEWHRVVLFGKVAEIAGEYLRKGSQCYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!**! !!**!*!!*!*!! ****!!***!*!* *!** !*!!!!********!!*!!*!!*!!* *!*

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYDEGYGQSQEAYQRPAQQSRQPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYEEGYGQSQEAYQRPAQQSRQPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG..   152
NTDB id 295990 DM292 RS15215 WP 063544073.1 EGRLQTREWEKD.GVKRYTTEIVV.DMNGTMQLLGGRGGNSDDA..........PRQPR....PQRE..PQQAPRQQA..   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM...NVPAQQG....SWGQ..PQQPAKQHQ..   149
consensus                                   !!****!***   !**!*******      ***!**!  *** *          ! * ** *   * * ****** ** *

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     APSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      APSHPQEAPAAPRRQPVP.AAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    .....NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 295990 DM292 RS15215 WP 063544073.1 .....QPQQPAARQQPAPDY....DSFDDDIPF   163
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .....PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                       *     ****** * *  *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved