```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
NTDB id 1105334 ACN2C7 RS11170 WP 419317027.1 .MAGSVNKVILVGNLGADPEIRSLGSGDRVANLRIATSESWRDRTSGERKEKTEWHRVVIFNDNLVKVAEQYLRKGSTVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                         **!!!!**!*!! !!**!****!**!*****!!!!*!*!* *!***!*!!!!******* *****!**!!*!!* !!

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP......YEE..GYGQS..QEAYQRPAQ.QSRQPA  140
NTDB id 1131 NGFG RS05740 WP 003695064.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP......YDE..GYGQS..QEAYQRPAQ.QSRQPA  140
NTDB id 1390 A4U84 RS00055 WP 021115958.1      VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFG.......GNQGSGWNQAPAQTNYNQGGYSDNYAQN  146
NTDB id 1105334 ACN2C7 RS11170 WP 419317027.1 IEGAIQTRKWTDNTGAEKYSTEIVLQKFRGELTMLGGRGDGAGASSGGGDDGYSSGGGGGG...............Y...  142
NTDB id 1166 A1552VC RS00795 WP 000168289.1   IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQP...............QQPA  145
consensus                                     *!!****!!**** ! ********     ** !!**! ******       *    ****   *  **  *     ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1105334 ACN2C7 RS11170 WP 419317027.1  ........GGGAPRSQPSGPRESFSADLDDEIPF   168
NTDB id 1166 A1552VC RS00795 WP 000168289.1    KQHQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                            *  ****  *   **  *****!!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved