```
logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMY.RKLAEIAGQYLKKGGLVY    75
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMY.RKLAEIAGQYLKKGGLVY    75
NTDB id 333457 RMHFA RS12040 WP 075822127.1  .MAGSVNKVILVGNLGRDPEVRNFQNGGKVVNLRIATSENYKDR.EGNRQERTEWHSVAIFNERLGEVAEKYLRKGSTVY    78
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLY.GKLAEVAGEYLRKGSQVY    79
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFF.RRQAEVAGQYLRKGSKVY    77
consensus                                        **!!!!**!*!!*!**!****!**! ****!!!!***!*  !** !*!!!! **** ****!*!**!!*!!* !!

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGY.......GQS.......QEAYQRPAQQSR   137
NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGY.......GQS.......QEAYQRPAQQSR   137
NTDB id 333457 RMHFA RS12040 WP 075822127.1  IEGKLETRKWQDKDGQDRYTTEIVLRNFGGELTMIGGRASGGEGG.EGGGYSGGGGGFGGGPRAGGGERGGA........   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP.AQGGM.......NVPAQ....QGSWGQPQQPA.   145
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFG.GNQGS.....GWNQAPAQTNYNQGGYSDNYAQN.   146
consensus                                    *!!****!!*!***!**!*******      ** !***!  *****    !*       ** *      ****  * **

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      QPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     QPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 333457 RMHFA RS12040 WP 075822127.1  .........SGGRSPSRGGGGGGWDAPKGDLDDDIPF   177
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ...KQHQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ...NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                             *   **    *     * *!****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved