```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY   75
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY   75
NTDB id 1029624 AB6B37 RS09240 WP 371395485.1 .MAGSVNKVILIGNVGKDPEVRTFGNGGKVCNFSIATSESWKDKQTGERKEKTDWHNIAIFNEGLVGIVERYVKKGTKVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGK.LAEVAGEYLRKGSQVY   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRR.QAEVAGQYLRKGSKVY   77
consensus                                         **!!!!**!**!*!!**!****!**!*****!!!!*!*!* *!***!*!*!!******* *******!**!!* !!

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEA.........YQRPAQQSRQPA..   140
NTDB id 1131 NGFG RS05740 WP 003695064.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEA.........YQRPAQQSRQPA..   140
NTDB id 1029624 AB6B37 RS09240 WP 371395485.1 IEGALKTRKWQDRDGNDRYTTEVVLQGYNGNLTLLDSRNSQGG.GGGGGDYGNQGGGYGGGGGQGNYGND.......GGG   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1   IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPAQ....QGSWGQPQQPAKQHQ..   149
NTDB id 1390 A4U84 RS00055 WP 021115958.1     VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG   152
consensus                                     *!!*****!!*!***!**!******       * *!**!* *** ***  ****** * *      ** *  * ** ** *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1029624 AB6B37 RS09240 WP 371395485.1 GS.......RSASA....MEGPADNFDLDDEIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1   PMQQSAPQQYSQPQ....YNEPP..MDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NATRPQPAQKPAAQ....AEPPM..DNFDDDIPF   180
consensus                                         * **  ***     * ***  ***!!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved