```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 1162799 AABQ55 RS13930 WP 244765530.1   .MAGSVNKVILIGNVGADPEIRRTQDGRPIANLRIATSETWRDRNSGERREKTEWHNVVVFNEGLCKVVEQYVKKGAKLY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                          **!!!!**!**! !!**!****!*******!!!!*!*!***!***!*!!!!******* ********!**!!* *!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQPAPDAPSHPQEA.   150
NTDB id 1103 NMB RS07590 WP 002212976.1         LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQPASDAPSHPQEA.   150
NTDB id 1390 A4U84 RS00055 WP 021115958.1       VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG   152
NTDB id 1162799 AABQ55 RS13930 WP 244765530.1   IEGALQTRKWQDQTGNDRYTTEVVLQGFNSTLTMLDGRGEGGG.ERRGGGD.FGAAPNYDDYGSR..P..AS......G.   146
NTDB id 1166 A1552VC RS00795 WP 000168289.1     IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM.NVP.AQQGSWGQPQQP..AKQHQPMQQ.   153
consensus                                       *!!****!!*!** !**!*******      * !!**! ****  *   ! ** * * *  ***  **   *   *  *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ....PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         ....PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1162799 AABQ55 RS13930 WP 244765530.1   ....G...GGSSRSSGSLSRDMDDDIPF   167
NTDB id 1166 A1552VC RS00795 WP 000168289.1     ....SAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                          **   *    ** *****!!!!!!
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X non conserved
X similar
X $\geq$ 50% conserved