```
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 1162532 AABF56 RS07230 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 1162532 AABF56 RS07230 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
NATRPAAPRRQPVPAAAPVEDIDDDIPF    174
NTDB id 1103 NMB RS07590 WP 002212976.1         ....PAAPRRQPVPAAAPVEDIDDDIPF    174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NATRPQPAQKPAAQAEPPMDNFDDDIPF    180
NTDB id 1162532 AABF56 RS07230 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