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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 25279 EL2594 RS00190 WP 011413009.1 .MAGSLNKVMLIGNLGQDPEVRSFQNGGKVCNLRIATSETWKDRNSGERQERTEWHTVAIFSEGLVNVAERYLKKGSKVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                       **!!!***!*!!*!!**!****!**!*****!!!!*!*!***!** !*!!!! ***** *****!**!!*!!* !!

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP.YDEGYGQSQEAYQRPAQQSRQPAPDAPS.....   145
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP.YEEGYGQSQEAYQRPAQQSRQPASDAPS.....   145
NTDB id 1390 A4U84 RS00055 WP 021115958.1    VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQA..PAQTNYNQGGYSDNYAQNNNFNG   151
NTDB id 25279 EL2594 RS00190 WP 011413009.1 IEGQLQTRKWQDQSGNDRYSTEVVLRGYNGTLTMLDGAGGGQGGGGG..GGYGGG..GQRSGGNNGGWDQGGGGQSGGQ.   154
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.......QG..GMNVPAQQGSWGQPQQPAKQHQ.   149
consensus                                   *!!****!!*!** !**!*******       * !!***  * ****    *****  * * ***** *    **    *

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 25279 EL2594 RS00190 WP 011413009.1 .SGGG......SGGGGGSNYDDLDDDIPF   176
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                    *  ***   *    ** *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved