```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 32539 BUAPTUC7 RS02755 WP 009874495.1 MASRGVNKVILIGHLGQDPEVRYMPNGNAVVNMTLATSENWKDKNTGENKEKTEWHRIVLFGKLAEIAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**! !!*!!**!*!!*!*!! ****!!!!*!*!***!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGYGQSQE.............AYQRPAQQSRQP  139
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGYGQSQE.............AYQRPAQQSRQP  139
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDF.GGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG.  152
NTDB id 32539 BUAPTUC7 RS02755 WP 009874495.1 EGSLQTRKWQDQNGLERYTTEIIVN.IGGTMQMLGNRNSNLQA.MTVNDKN...ST.......GIKIK.KTDAVEFDK..  145
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMN...VPAQ....QGSWGQ.PQQPAKQHQ..  149
consensus                                     !!****!!*!** !**!******    *  **!!**!* * ** *   * *  *         *     ** **** *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF  174
NTDB id 1131 NGFG RS05740 WP 003695064.1       APDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF  174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      .......NATRPQPAQKPAAQAEPPMDNFDDDIPF  180
NTDB id 32539 BUAPTUC7 RS02755 WP 009874495.1 .......KI..ETHESNKDLTHSPSEIDFDDEIPF  171
NTDB id 1166 A1552VC RS00795 WP 000168289.1   .......PMQQSAPQQYSQPQYNEPPMDFDDDIPF  177
consensus                                            *   **   *  * *  *****!!*!!!
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X non conserved
X similar
X $\geq$ 50% conserved