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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 413786 GTQ55 RS14565 WP 161859397.1 .MARGVNKVILIGNLGGDPETRYLPSGGAVTNITVATSETWKDKQTGQQQERTEWHRVVFFNRLAEIAGEYLRKGSKVYL   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!*!! !!**!**!*!*!! ****!!!!*!*!* *!** !*!!!!********!!*!!*!!*!!* !!*

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGY..GQSQEAYQ....................   130
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGY..GQSQEAYQ....................   130
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQ.NGQDRYTTEIQGD....VLQMLDSRSSGGDFG........GNQGSGWNQAPAQTNYNQGGYSDNYAQ   145
NTDB id 413786 GTQ55 RS14565 WP 161859397.1 EGSLRTRKWQDKTSGQDRYTTEIVAS....EMQMLDGRGEQGGGYQQGGGQGFQQGGGYAQQ....GPQGGGYQDEYSQ   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMN.VPAQQGSWGQ..................   140
consensus                                   !!*****!!*!**  !**!*******    ***!!**! ******    *   **** ****

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1     .....RPAQQSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      .....RPAQQSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NNNFNGGN..ATRPQPAQ..KP......AA....QAEPPMDNFDDDIPF   180
NTDB id 413786 GTQ55 RS14565 WP 161859397.1 GRSAPSPMAPSNQPQPGQPQNK......PAGNQPPAGGFDNSFDDDIPF   194
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ......PQQPAKQHQPMQQSAP......QQYSQPQYNEPPMDFDDDIPF   177
consensus                                      * * ****** *** *       *   *  *********!!!!!!
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X non conserved

X similar

X ≥ 50% conserved