```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWN.DRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWN.DRNGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1132453 H7A87 RS13525 WP 119127850.1  .MARGINKVILVGNLGNDPDTKYTQAGMAITRVSLATTSMR.KDREGNNQERTEWHRVVFFGKLGEIAGEYLRKGSQVYV   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                        **!!!!**!*!!*!!*******!*!* ****!!***** ** !** !*!!!!*********!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQ......SRQPASDAPSH   146
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQ......SRQPAPDAPSH   146
NTDB id 1132453 H7A87 RS13525 WP 119127850.1  EGELRYDKYTGQDGVEKYSTDIVA....NEMQMLGGRGEGGGGGG.AGMGGDRPQRSQAPRQQQGGG........GQGGG   145
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.QGGM...NVPAQ.....QGSWGQPQQPAKQHQ..   149
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGG.NQGSGWNQAPAQTN.YNQGGYSDNYAQNNNFNGG   152
consensus                                    !!*****!*****!**********    ****!!**! *******   * *  *   ! * ****      ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1       PQEAPAAPRRQPVPAAAPVEDI.DDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1      PQEAPAAPRRQPVPAAAPVEDI.DDDIPF   174
NTDB id 1132453 H7A87 RS13525 WP 119127850.1  QDYAPRRQQPAQQQSAPPMDDFADDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  PMQQSAPQQYSQPQYNEPPMDF.DDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NATRPQPAQKPAAQAEPPMDNF.DDDIPF   180
consensus                                    *  ***  **   *** !**** !!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved