```
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 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 140857 UC35 RS11100 WP 061499347.1    ..MASVNKVIIVGNLGRDPEMRSFPSGDRVANVTIATTDKWKDKQTGEMKEATEWHRVVFNGRLAEIAGEYLRKGSQVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                        **!!!!**!*!!*!!**!**!*!**!**!***!!***!*!* *!***!*!!!!*********!!*!!*!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQPAPDAPSH....PQ   148
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQPASDAPSH....PQ   148
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQ...GGMNVPAQQ....GSWGQPQQPAKQ.HQPM   151
NTDB id 140857 UC35 RS11100 WP 061499347.1    EGSLRTRKWTDQQGVEKYTTEIRA....DTMQMLGSRQGMGG.PAPDEGDGGMRR.............AQ........PA   132
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
consensus                                     !!*****!!**** !**********       **!!**!* ***  * *** !  *   *   **        **       **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1      EAPAAPR.RQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1       EAPAAPR.RQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1   QQ.SAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 140857 UC35 RS11100 WP 061499347.1    ARSAPAASKPASKPSTGFDDMDDDIPF   159
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                        ***  *    *  *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved