```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1115015 BN2297 RS22990 WP 003064700.1   ..MASVNKVIIVGNLGRDPEMRTFPSGDQVANVTIATTDRWRDKNTGENREATEWHRVVFNGKLAEIVGQYLRKGSQVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**!*!!*!!**!**!*!**!**!***!!***!*!***!***!*!!!!*********!!**!*!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYDEGYGQSQEAYQRPAQQSRQPAP   141
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGK.DGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYEEGYGQSQEAYQRPAQQSRQPAS   141
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGG...MNVPAQ....Q..GSWGQPQQ....PAK   146
NTDB id 1115015 BN2297 RS22990 WP 003064700.1   EGSLRTRKWTDQASGQERYATEIRA....DAMQMLGSRQGGGQQQGGYGSDD.......GYGE..SSYEAPRRQAPAPMQ   145
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQ.NGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQ..GGYSDNYAQNNN.F.   149
consensus                                       !!*****!****  !**!*******      **!!**!* ***  *  *     **   ****   **  * **  ***

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        DAPSH....PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         DAPSH....PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1     Q...HQPMQQ.SA..PQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1115015 BN2297 RS22990 WP 003064700.1   QRPAPAPMQQRPAPAPMAPPPQRAASGFDDMDDDIPF   182
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ......NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                         * *     *  ****  * *  ** *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved