```
logo
NTDB id 1141198 DQN52 RS10740 WP 003787468.1   ...MSLNKVMLIGRLGRDPETRYMPNGDAVCNFSVATSENWTDRNNGQRQERTEWHNITMYRRQAEIAAQYLRKGSQVYL   77
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      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         ***!!!***!*!!*!!**!*!!*!*!!*****!!!!*!*!* *!** !*!!!!********!!*!**!!*!!* !!*

logo
NTDB id 1141198 DQN52 RS10740 WP 003787468.1   EGKIQSRKYTDKTGAERTAYDIVC....DVMRMLDSKNGQQGGS.......SHVPYDNEYSAAP......PPARQQPAPA   140
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYDEGYGQSQEAYQRPAQQSRQPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYEEGYGQSQEAYQRPAQQSRQPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGG..YSDNYAQNNNFN.G   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM...NVPAQ....QGS..WGQPQQPAKQHQ..   149
consensus                                      !!****!!**** ! *!*******     ***!!**** * ***          !**  * **   *  * ** ****

logo
NTDB id 1141198 DQN52 RS10740 WP 003787468.1   APASEPPVAPQRRESKPVAPVEPVENIDEDIPF   173
NTDB id 1131 NGFG RS05740 WP 003695064.1       APS.HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1        APS.HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ....GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1    .....PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                      **   *  ***   * ** *  !****!*!!!!
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X non conserved

X similar

X ≥ 50% conserved