```
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1058870 ACFYG5 RS12395 WP 395116683.1 .MARGVNKVILIGNLGSDPEVRSTGSGTTVCSFSLATSESWTDKQSGEKQERTEWHRIKIFGRLGEIAGEYLKKGRQVYI   79
NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
consensus                                        **!!!!**!*!! !!**!****!**!*****!!!!*!*!* *!** !*!!!!*****!*!!*!!*!!  !!*

logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM...NVPAQ....Q.......GSWGQPQQPAK   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQ.......GGYSDNYAQNN   147
NTDB id 1058870 ACFYG5 RS12395 WP 395116683.1 EGSIRTDKFTGKDGVEKYFTDIIA....NEMQMLGSNEGGGGGG...........YQRERPQGGQRQGGGNYGGQSRGND   144
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYEEGYGQ.....SQEAYQRPAQQSR   137
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGA..........PYDEGYGQ.....SQEAYQRPAQQSR   137
consensus                                     !!*****!*****!**** *****    ****!!***  *****          ***  * !        ***  * ** *

logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QHQ......PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NFNGG....NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1058870 ACFYG5 RS12395 WP 395116683.1 DRGSSRP.............SSPAPADNGFTDDEIPF   168
NTDB id 1103 NMB RS07590 WP 002212976.1       QPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1      QPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
consensus                                     *  *   *  *   **   *   ** ***** !!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved