```
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 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 43932 SBAL678 RS26345 WP 006083192.1   MASRGVNKVILVGNLGQDPEVRYMPNGNAVANITVATSESWKDQ.QGQQQERTEWHRVVLFGKLAEITGEYLRKGSQVYL   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                          **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!* *!** !*!!!!********!!**!*!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENS..GGAPYDEGYGQSQ.......................   126
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENS..GGAPYEEGYGQSQ.......................   126
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSG.........WNQAPAQTNYNQGGYSD....   141
NTDB id 43932 SBAL678 RS26345 WP 006083192.1   EGKLQTRKWKDQSGQDRYSTEVVIDQ.SGSMQMLGSRGQGGQAQGAPMGGGMPQNA..GYQSAPQQAAPAQNQYAPAPQA   156
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG...GMPAQGGMNVPAQQGSWGQPQQPA.........KQH   148
consensus                                      !!****!!**** !**!*******       **!!**!  **  *!**  **  *         * *

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        .EAYQRPAQQSR.QPA.......................PDA.PSHPQEAPAAPRRQPVP......AAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         .EAYQRPAQQSR.QPA.......................SDA.PSHPQEAPAAPRRQPVP......AAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..........NYA............QNNNFNGGNATRPQPAQKPAAQ..................AEPPMDNFDDDIPF   180
NTDB id 43932 SBAL678 RS26345 WP 006083192.1   APAYQAPAQQQYAAPAPAQQQYGQQQAQPQQGGYAPKPQAAPAPAYQAPAAPAQRPAPQPQQNFTPDLDDGWDDDIPF   234
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QPMQ.QSAPQQYS...........................QPQYNE....................PPMDFDDDIPF   177
consensus                                        *** ******  **                        * * * *  ******   * *      *  *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved