```
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 1171227 ACOQOE RS17755 WP 420554608.1  .MARGVNKVILVGNLGGDPEVRYMPNGNAVTNVTIATSESWKDKQTGQQQERTEWHRVVFFNRLAEIAGEYMRKGGKVYV   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                          **!!!!**!*!! !!**!*!!*!*!! *****!!!!*!*!* *!** !*!!!!*********!!*!!*!**!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSG.......G...APYDEGYGQSQEA......YQRPAQQS   136
NTDB id 1103 NMB RS07590 WP 002212976.1         EGRIQSRKYQGKDGIERTAYDIVANEM....KMLGGRNENSG.......G...APYEEGYGQSQEA......YQRPAQQS   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGG.......YSDNYAQN   146
NTDB id 1171227 ACOQOE RS17755 WP 420554608.1  EGALRTRKWQDQSGADRYTTEIVASEMQ....LLDSRGAGD......GGGYQQQPQAGGYAPQQSAPAPQQPASQPQYAP   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQ........QGGM...PAQGGMN....VPAQQGSWGQPQQPA   145
consensus                                      !!*****!!*!** ! *!********** *!**!            !   *!   **** * *      *  * **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        RQPAPDAPSHPQEAPAAP.RRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         RQPASDAPSHPQEAPAAP.RRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NN......FNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1171227 ACOQOE RS17755 WP 420554608.1  QQ......SKPAQRPAAPQQSAPAPQPAPGFDDFDDDIPF   183
NTDB id 1166 A1552VC RS00795 WP 000168289.1    KQ......HQPMQQSAPQQYS..QPQYNEPPMDFDDDIPF   177
consensus                                      **       * * ******       *** *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved