```
logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1033396 AB8Z76 RS15210 WP 099770622.1 .MARGINKVILVGNLGNDPDTKYTQAGMAITRVSLATTSMRKDR.DGNNQERTEWHRVVFFGKLGEIAGEYLRKGSQVYV   78
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.......QGSWG...QPQQPAKQ  147
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQT...NYNQGGYS...DNYAQNNN  148
NTDB id 1033396 AB8Z76 RS15210 WP 099770622.1 EGELRYDKYTGQDGVEKYSTDIVA....NEMQMLGGRGEGGGGGMGGDRPQRSASRQQGGGGGQGGGYGGGGQDYAPRRQ  154
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYE...................EGYGQSQEAYQRPAQ  134
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYD...................EGYGQSQEAYQRPAQ  134
consensus                                     !!*****!*****!**********    ****!!**! ******           *         ****   * **   *

logo
NTDB id 1166 A1552VC RS00795 WP 000168289.1    HQ..PMQ.........QSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1      FNGGNAT.........RPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1033396 AB8Z76 RS15210 WP 099770622.1 ........QPAQQ............QSAPPMDDFADDDIPF   175
NTDB id 1103 NMB RS07590 WP 002212976.1       QSRQPASDAPSHPQEAPAAPRRQPVPAAAPVE.DIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1      QSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVE.DIDDDIPF   174
consensus                                        **   *         **   * * * *** * !!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved