```
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 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 395897 GFH30 RS00825 WP 153370262.1 ..MRGVNKVILVGTLGKDPETKTFANGGSLTQFSIATSDSWTDKSTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                      **!!!!**! !!*!!*******!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS..........QEAYQRPAQQSR.....   137
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS..........QEAYQRPAQQSR.....   137
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPA................QGGMNVPAQQGSWGQPQ   142
NTDB id 395897 GFH30 RS00825 WP 153370262.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSNRQQGE.QGD...TGFNQPRFNNNQNNQGGYNNNANQGGYNNAP   150
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQT.NYNQGGYSDNYAQ.......   145
consensus                                   !!*****!***** !**!*******       **!!***  ***  *   * *            !**** ***!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     .....QPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     .....QPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 .Q......PAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 395897 GFH30 RS00825 WP 153370262.1 QQGGFNNAPANGYGNNNPSGFAPKAPAPTTAPAADLDDDLPF   192
NTDB id 1390 A4U84 RS00055 WP 021115958.1   .NNNFNG......GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                    *   * **     **  **   * * ** *!****!!!*!!
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X non conserved

X similar

X $\geq$ 50% conserved