```
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 366704 FF090 RS02470 WP 138855223.1 ..MASVNKVIVLGNLGRDPELRYTPSGAAVCNVSIATTRNWKSKDSGERQEETEWHRVVFYDRLAEIAGEYLKKGRPVYV    78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                       **!!!!**!*!!*!!**!**!*!*!!**!***!!***!*** *!** !*!!!!*********!!*!!*!!*!!  !!*

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDE...........GYGQS..QEAYQRPAQQSRQP   139
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEE...........GYGQS..QEAYQRPAQQSRQP   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMN....VPA....QQGSWG....QPQQPAKQH   148
NTDB id 366704 FF090 RS02470 WP 138855223.1 EGRLKTRKWQDKEGRDQYTTEIIA....EQMQLLGGREDGGGGGGSGYGGGSRGAP......QGG.GDDYGHDAPAERPA   147
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGW.NQAPAQTNYNQGGYSDNYAQNNNFN..G   151
consensus                                   !!****!!*!***! ********      ***!**! *******          *    * !*   * *  ***  **

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1     APDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1     ASDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QPMQ......QSAPQ.QYSQPQYNEPPMDFDDDIPF   177
NTDB id 366704 FF090 RS02470 WP 138855223.1 RPAA......RPAPQRPAPAPKPASSFDDMDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1   GNAT......RPQPA.QKPAAQAEPPMDNFDDDIPF   180
consensus                                   ** *        *** *  *   ** *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved