```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1089753 ACLVWN RS06580 WP 010907419.1 ..MAGVNKVIIVGNLGNDPEIRTMPNGEAVANISVATSESWIDKNTNERREVTEWHRIVFYRRQAEVAGEYLRKGSKVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!*******!*!!!!*********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEG..........YGQ..SQEAYQRPAQQSRQPA   140
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEG..........YGQ..SQEAYQRPAQQSRQPA   140
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM...PAQGGMNVPAQ....QGS....WGQPQQPAKQHQ   149
NTDB id 1089753 ACLVWN RS06580 WP 010907419.1 EGRLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRNERQQTGGYAP.....QTAAPQYNA.........PTGSY....   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNG..G   152
consensus                                      !!****!!*!** !**!******     ***!!**!***** * *  *         *   * *    * *  * *  ** *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        SDAPSHPQEAPAAPRRQPVPAAAPVE...DIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PDAPSHPQEAPAAPRRQPVPAAAPVE...DIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    PMQQSAPQQ......YSQPQYNEPPMD...FDDDIPF   177
NTDB id 1089753 ACLVWN RS06580 WP 010907419.1 GAQPSRPAT......K.PAPQNEPPMDMGFEEDNIPF   166
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NATRPQPAQ......K.PAAQAEPPMDN..FDDDIPF   180
consensus                                         ** !*        * * * **!***     *!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved