```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
NTDB id 1049689 ACETKC RS12540 WP 271164175.1 .MAGSVNKVILVGNLGRDPEIRSMPNGDRIANLSIATSETWRDKSSGERKEKTEWHRVVIFNDNIVKVVENYVKKGSTVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                         **!!!!**!*!!*!!**!*!!*!*******!!!!*!*!* *!***!*!!!!*****  ********!**!!* !!

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQPASDA.....PSH   146
NTDB id 1131 NGFG RS05740 WP 003695064.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQPAPDA.....PSH   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1      VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGG   152
NTDB id 1049689 ACETKC RS12540 WP 271164175.1 IEGALQTRKWTDQQGVEKYSTEIVVSRFKGELTMLGGRAEGG.ASQG.GGYG..GGGGRG...DDDYS.........SGF   143
NTDB id 1166 A1552VC RS00795 WP 000168289.1   IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMN..VPAQQGSWGQPQQPAK.....QHQPM   151
consensus                                     *!!****!!**** !*********      ** !!**! *** * **  !**  * * *   **  ***  *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1049689 ACETKC RS12540 WP 271164175.1  ...STGGANKP..SGPKESYDLNDDIPF   166
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QQSAPQQYSQP..QYNEPPMDFDDDIPF   177
consensus                                       * ****  **   *  ******!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved