```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1091275 ACL2XQ RS09045 WP 187514540.1 MSSRGVNKVILIGFLGQEPEVRYLPNGMAVTNISLATSESWRDKQTGEEKERTEWHRVVLFGKLAEIAGEYLHKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**! !!**!**!**!*!*!! ****!!!!*!*!* *!***!*!!!!********!!*!!*!!*!!* !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEA.........YQRPAQQSRQPASDA   143
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEA.........YQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQN..NNF..   149
NTDB id 1091275 ACL2XQ RS09045 WP 187514540.1 EGQLQTRKWTDNAGVERWTTEIVVKQ.QGTMQMLGSNNRTN.TS....GSGQ...PAN....KQ..HSAAA....PAQ..   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMNV...PAQ....QGSWGQPQQ....PAK..   146
consensus                                     !!****!!**** !**!******       **!!**** *  * *   ! ** * ***        *  ****   **

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.................   180
NTDB id 1091275 ACL2XQ RS09045 WP 187514540.1  SGTPQSQSAP....QSMGNEPPMDFEDDIPFLGYGYGVCRGALYGIS   182
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF.................   177
consensus                                         **  ***  *     *  !*****!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved