```
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 1167940 ACLIIK RS28095 WP 062838810.1 ..MASVNKVILVGNLGRDPEVRYSPDGAAICNMSIATISSWKDKASGEKREETEWHRVVMYNRLAEIAGEYLKKGRSVYI   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         **!!!!**!*!!*!!**!**!*!*!*****!!***!*!* *!***!*!!!!********!!*!!*!!*!!  !!*

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPY.EEGYGQSQEAYQRPAQQ.........S.....   136
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGK.DGIERTAYDIVAN....EMKMLGGRNENSGGAPY.DEGYGQSQEAYQRPAQQ.........S.....   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQ.NGQDRYTTEIQGD....VLQMLDSRSSGGDFGGNQGS..GWNQAPAQTNYNQGGYSDNYAQNNNFNG   151
NTDB id 1167940 ACLIIK RS28095 WP 062838810.1 EGRLKTRKWQDKDTGADRYSTEIVAD....QMQMLGGREDGGGGGGSYGGGGGYDDAPRQQRA.................   137
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQ.SGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQ....GGMNVPAQQGSW................   138
consensus                                     !!****!!*!**  ! *!********     **!!**! *******    * !  * * !  ***

logo
NTDB id 1103 NMB RS07590 WP 002212976.1        RQPASDAPS.HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       RQPAPDAPS.HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      GNATRPQPAQKPAA.........QAEPPMDNFDDDIPF   180
NTDB id 1167940 ACLIIK RS28095 WP 062838810.1  APAQRPAPQQRPAA....PAAAPASSGANLADMDDDIPF   172
NTDB id 1166 A1552VC RS00795 WP 000168289.1    GQPQQPAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                      ***  ***   !    * *   *  ********!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved