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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 103835 QUF06 RS16035 WP 015872740.1  MASRGVNKVILIGNLGQDPEVRYMPNGGAVANLTLATSDSWRDKQTGEQKERTEWHRVALYGKLAEIAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!*!!*!!**!*!!*!*!!*****!!!**!*!* *!***!*!!!!*********!!*!!*!!*!!* !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP........YEEGYGQSQEAY..QRP......AQQS   136
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP........YDEGYGQSQEAY..QRP......AQQS   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGGN.......QGSGWNQAPAQTNYNQGGYSDNYAQNN   147
NTDB id 103835 QUF06 RS16035 WP 015872740.1  EGQLRTRKWQDQSGQDRYTTEVVVD.ISGSMQMLGGRQGGAP......MGNQGQQGGWGQPQQPQ.............GA   140
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPA.............KQ   147
consensus                                    !!****!!*!** !**!********      **!!**!* *** *          *  ***! *      *         ***
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NTDB id 1103 NMB RS07590 WP 002212976.1      RQPASDA.PS..HPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     RQPAPDA.PS..HPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NFNGGNAT.........RPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 103835 QUF06 RS16035 WP 015872740.1  AFSGGQAAPRQQSQQGGYSAPAQPAAPQSSEPPMDFDDDIPF   182
NTDB id 1166 A1552VC RS00795 WP 000168289.1  HQP............MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                     ***  * *      * *  *! *    **  !****!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved