logo

```
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL  76
NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL  76
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI  80
NTDB id 1002619 ABEF85 RS00860 WP 034588730.1  ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSESWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI  78
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV  78
consensus                                          **!!!!**! !!*!!*****!*!*** ****!!!!*!*!***!***!*!!!!*********!*!**!!*!!* !!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQE...............AYQRPAQQSR  137
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQE...............AYQRPAQQSR  137
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQ..GGMNVP...........AQQGSWGQPQQPAK  146
NTDB id 1002619 ABEF85 RS00860 WP 034588730.1  EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDSGRAQGG.EQSD.NGGFAQPRFNNNNQQGGYANQGGYGAPQQGGF  152
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSD............  141
consensus                                      !!****!***** !**!*******      **!!***  ***  * * * !  *                **  * *  *
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NTDB id 1131 NGFG RS05740 WP 003695064.1       Q...PAPDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF  174
NTDB id 1103 NMB RS07590 WP 002212976.1        Q...PASDAPSHPQEAPAAPRRQPV.PAAAPVEDIDDDIPF  174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    QH.........QPMQQSAPQQYSQPQYNEPPMDFDDDIPF  177
NTDB id 1002619 ABEF85 RS00860 WP 034588730.1  NNNAPAGNGY.GNNPGGFAPKAAPA.AVAAPAADLDDDLPF  191
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..NYAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF  180
consensus                                      *   **  *    * *  *!*  *   ***!****!!!*!!
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X non conserved

X similar

X ≥50% conserved