```
logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 207809 BSR56 RS00915 WP 100833074.1 ..MRGVNKVILVGTLGKDPETKTFPNGGSLTQFSIATSESWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!!*!*!***!***!*!!!!**********!*!**!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP............YEEGYGQSQE...........AY   129
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAP............YDEGYGQSQE...........AY   129
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVP....AQQGSWGQPQQPA..........   145
NTDB id 207809 BSR56 RS00915 WP 100833074.1 EGSLRTRQWTDQNGQERYSTEIRG....DQMQMLDSRQQSEQSG..GDFNQPRFNNNQGGGYQNTGHNN..N....QNGY   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFG..G.........NQGSGWNQAPAQTNYNQGGYSDNY   143
consensus                                   !!*****!***** !**!******       **!!**!* * * *  *          *  ******            **

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      QRPAQQSRQPASDAPSHPQEAPAA.PRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     QRPAQQSRQPAPDAPSHPQEAPAA.PRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  .......KQH......QPMQQ.SAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 207809 BSR56 RS00915 WP 100833074.1 GQGGGFNGGNQGNYAGSPQAGNGFNSPKAPQPAATAPADLDDDLPF   192
NTDB id 1390 A4U84 RS00055 WP 021115958.1   AQNNNFNGGNATR....PQPA.....QKPAAQAEPPMDNFDDDIPF   180
consensus                                      **  ** *      !* * **   * * *** *****!!!*!!
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X non conserved

X similar

X ≥ 50% conserved