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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 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 314728 CDG62 RS03775 WP 087526717.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDSWTDKSTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!**!*!* *!***!*!!!!**********!*!**!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQ........EAYQRPAQQ........S   136
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQ........EAYQRPAQQ........S   136
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM..NVP....AQQGSWGQPQ...........   142
NTDB id 314728 CDG62 RS03775 WP 087526717.1 EGSLRTRQWTDQNGQERYTTEIRG....EQMQMLDNNRQQGE.QMQGGNDGYAQPRFNNNQGGGYNNN.....NNQGGYG   148
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FG.............GNQGSGWNQAPAQTNYNQGGYS   140
consensus                                   !!****!***** !**!*******      **!!***  ***  *   * *  *      *  *****  *        *

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NTDB id 1103 NMB RS07590 WP 002212976.1      RQPASDA.......PSHPQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     RQPAPDA.......PSHPQEAPAAPRRQP.VPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  .QPAK..........QHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 314728 CDG62 RS03775 WP 087526717.1 GNPATGNSNNQQGGYANNNPSGFAPKAAPQSAPATAPADMDDDLPF   194
NTDB id 1390 A4U84 RS00055 WP 021115958.1   DNYAQNN..NFNGGNA....TRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                    **!            * *    *!*  *   ** *****!!!*!!
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X non conserved

X similar

X ≥ 50% conserved