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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 400660 GJD93 RS13855 WP 096902231.1 ..MRGVNKVILVGTLGRDPETKTFPNGGSLTQFSIATSDVWTDKNTGERKEQTEWHRIVLHNRLGEIAQQYLRKGSKVYI    78
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                       **!!!!**! !!*!!*****!*!*** ****!!!**!*!***!***!*!!!!**********!*!**!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQ.......SQEAYQRPAQQSRQP......   139
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQ.......SQEAYQRPAQQSRQP......   139
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQ..GGMNVPA...QQGSWGQP.QQPAKQ......   147
NTDB id 400660 GJD93 RS13855 WP 096902231.1 EGSLRTRQWTDQNGQERYTTEIRG....DQMQMLDSANRGQSD.QQD..GGFAQPRFNNNQGGYGNQNQAPATGYANQNQ   151
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.GGNQGSGWN...............QAPA..QTNYNQ   136
consensus                                    !!*****!***** !**!*******       **!!**** * *  * * * !         * **  * !***

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ........ASDAPSH.PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     ........APDAPSH.PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ............HQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 400660 GJD93 RS13855 WP 096902231.1 GGFNQNQGGGYANNNPSGFAP....KPQAAPATAPADLDDDLPF   191
NTDB id 1390 A4U84 RS00055 WP 021115958.1    GGYSDNYAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                            *  * *  *  ***   *    ** *****!!!*!!
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X non conserved

X similar

X ≥ 50% conserved