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logo
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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 1310753 D0B88 RS00780 WP 007644503.1 .MARGINKVILIGNIGQDPEVKYMPSGGAVTNVSVATSETWKDKNTGQPQERTEWHRVVFFNRLGEIAGEYLKKGSKVYI   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                      **!!!!**!**!*!!****!!*!*!! ****!!!!*!*!***!** !*!!!!**********!*!!*!!*!!* !!*
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logo
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NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS..QEAYQR......PAQQSRQPA....   140
NTDB id 1131 NGFG RS05740 WP 003695064.1    EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS..QEAYQR......PAQQSRQPA....   140
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMNVPAQQGSWGQPQ..................   142
NTDB id 1310753 D0B88 RS00780 WP 007644503.1 EGALRTRKWQAQDGTDRYTTEIVA....SEMQMLDGRGDGAGM.GNNMGAGGFDQTPQYNNQQHSNAPAQNAYNQAAAKP   154
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDF.GGNQGSGW.NQAP..........AQTNYNQGGYSD   141
consensus                                   !!*****!!*!***! *!*******     ***!!**! **** * * ! *   **  **       ****  * *
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logo
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NTDB id 1103 NMB RS07590 WP 002212976.1     ............SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1    ............PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ...........QPAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1310753 D0B88 RS00780 WP 007644503.1 A...ANNQGGGNQGGFGQPANRPAPQQNV.PPQPAGFDSFDDDIPF   196
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NYAQNNNFNGGN.ATRPQPAQKPAAQ......AEPPMDNFDDDIPF   180
consensus                                          * **  *! * *!**     ** ******!!!!!!
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X non conserved

X similar

X ≥ 50% conserved