logo

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NTDB id 1103 NMB RS07590 WP 002212976.1   ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1  ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 109887 ACGHM3 RS20670 WP 345427009.1 .MARGINKVILVGNCGQDPETKYMPSGGAVTNLSIATSETWKDKQTGQPQERTEWHRVVFFNRLAEIAGEYLRKGSQVYV   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                  **!!!!**!**!*!!*****!!*!*!! ****!!!!*!*!* *!** !*!!!!*********!!*!!*!!*!!* !!*
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NTDB id 1103 NMB RS07590 WP 002212976.1   EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGG.............APYEEGYGQSQEAYQ..........   130
NTDB id 1131 NGFG RS05740 WP 003695064.1  EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGG.............APYDEGYGQSQEAYQ..........   130
NTDB id 1390 A4U84 RS00055 WP 021115958.1 EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGD..FGGNQGSGWNQAPAQTNYNQGGYSD...........   141
NTDB id 109887 ACGHM3 RS20670 WP 345427009.1 EGSLRTRKWQDQSGQDRYSTEIVANE....MQMLGGRTEAAAGQVRPAQG....GYGNQGGMNQGGVNQGGPQGGGMAPP   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQ.........QG....GMPAQGGM..............N..   130
consensus                                  !!*****!!*!** !**!**********     **!!**! *  **      **      ** * ** *
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logo

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NTDB id 1103 NMB RS07590 WP 002212976.1   ....RPAQQSRQPASDAP.SHPQEA.PAAPRR.QPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1  ....RPAQQSRQPAPDAP.SHPQEA.PAAPRR.QPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1 ....NYAQNNNFNGGNA..TRPQ......PAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 109887 ACGHM3 RS20670 WP 345427009.1 MNQGAPAQGGGYGGGQAPASRPQQ....PP...QQPQQPPGMDSFDDDIPF   195
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....VPAQQGSWGQPQQPAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                      *!!*  * * *** * *!     *!   *  **  *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved