logo

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NTDB id 1020488 ABU615 RS04220 WP 370389268.1  ...MSVNKVILVGRLGRDPETRYMPSGDAVTNFSIATDEQWRDR.NGERQTRTEWHNITLFGKLGEIANQYLRKGSQVFI   76
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 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        ***!!!!**!*!!*!!**!*!!*!*!! ****!!*!*!*!* *!** **!!!!**********!*!**!!*!!* !**
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NTDB id 1020488 ABU615 RS04220 WP 370389268.1  EGRIQSRKYSGKDGIERTAYDIIG....SEMKMLGSRNDTGAASGSYDQMGGGADYAQGGNFSQGGYDA.........ASA   144
NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYEEGYGQSQEAYQRPAQQSRQPASD   142
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSG..........GAPYDEGYGQSQEAYQRPAQQSRQPAPD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQA..PAQTNYNQGGYSDNYAQNNNFNGG   152
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGM...NV..PAQ....QGSWGQPQQPAKQHQ..   149
consensus                                      !!****!!*****!**!*******      ***!!**!*****  *       ** *  *   *!***  * ** ** *
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logo

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NTDB id 1020488 ABU615 RS04220 WP 370389268.1  ASSPPPAAPRRQAPRPAPAPAPIDDIDDDIPF   176
NTDB id 1103 NMB RS07590 WP 002212976.1        APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       APSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ....NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1    ....PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                      * * *  ***   * * ** *!****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved