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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 160038 UZ73 RS11350 WP 035269622.1  ..MASVNKVILVGNLGRDPEVRYSAEGSAICNISIATISQWKDRTSGERREETEWHRVVFYNRLAEIAGEYLRKGRPVYV   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**!*!!*!**!****!*!******!!***!*!* *!***!*!!!!*********!!*!!*!!*!!  !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY..........EEGYGQSQ..EAYQRPAQ......   134
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY..........DEGYGQSQ..EAYQRPAQ......   134
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMP..AQGGMNVPAQQGSWGQPQ.............QPA   145
NTDB id 160038 UZ73 RS11350 WP 035269622.1  EGRLRTRKWTGQDGQERFTTEIIA....EQMQMLGGRDGGDMGGGSMGGGE........YG..............GGGGA   132
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGNA   154
consensus                                   !!****!!*****!**!*******       **!!**!  * ***     *          *** *  * *    *     *

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      QSRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     QSRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 KQHQPMQQSAPQQ........YSQPQYNEPPMDFDDDIPF   177
NTDB id 160038 UZ73 RS11350 WP 035269622.1  PAPQRAQRPAPQQQQQQQAP.RNNAPMSDNLADMDDDIPF   171
NTDB id 1390 A4U84 RS00055 WP 021115958.1   TRPQPAQKPAA..............QAEPPMDNFDDDIPF   180
consensus                                     !***  *   *    ** **  **  *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved