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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL    76
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
NTDB id 398922 GH975 RS09665 WP 153714325.1  .MARGINKVILVGNLGNDPEMRATNNGSQIANLSIATSESWRDKNTGEMQERTEWHRVVFFNRLAEICGQYLKKGSKVYV    79
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
consensus                                        **!!!!**!*!!*!!**!****!********!!!!*!*!***!** !*!!!!*********!!**!*!!*!!* !!*

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQG.KDGIERTAYDIV....ANEMKMLGGRNENSGGA.PYDEGYGQSQEA..YQ.........R..P..AQQ   135
NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQG.KDGIERTAYDIV....ANEMKMLGGRNENSGGA.PYEEGYGQSQEA..YQ.........R..P..AQQ   135
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQD.QSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQ..GGMNVPAQQGSW................GQ   140
NTDB id 398922 GH975 RS09665 WP 153714325.1  EGSLRTRSWDDAQSGQKRYTTEIV....GGEMQMLDGRGDNMGGGAPMGGGYQQNNAPQ..............AP...SA   138
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQD.QNGQDRYTTEIQ....GDVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
consensus                                     !!*****!**** * !**!******    * ***!!**! ** *****   !*** *    **           *  * *

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      SRQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1       SRQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1   PQQPAKQHQPMQQS..APQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 398922 GH975 RS09665 WP 153714325.1  PRQAAPAPAPA.......QAASAPPPGFADDIGDDDIPF   170
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ATRPQPAQKPA.......AQAEPPMDN.....FDDDIPF   180
consensus                                      *****   *  *   *     *** * * ** !!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved