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logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1103 NMB RS07590 WP 002212976.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 288539 HYN69 RS09505 WP 108435532.1  .MAGSVNKVIIVGNLGRDPEVRSFPNGGKVVNLRIATSENWRDKATGERKERTEWHSVAIFNENLAKIAEQYLRKGSTVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                        **!!!!**!*!!*!!**!**!*!**! ****!!!!*!*!* *!***!*!!!! ***** **!**!**!!*!!* !!

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQSQEAYQRPAQQSRQPAPDAP.......   144
NTDB id 1103 NMB RS07590 WP 002212976.1       LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQSQEAYQRPAQQSRQPASDAP.......   144
NTDB id 1390 A4U84 RS00055 WP 021115958.1     VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGG.NQGSGWN....QAPAQTNYNQGGYSDN......   142
NTDB id 288539 HYN69 RS09505 WP 108435532.1  IEGQLETRKWQDQSGADRYTTEVVLRAFRGELTLLGGRGEGGGGGG.GGGD.......DRGGYDDYQGGGSASG..GGGA   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1  IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA.QGGM.......NVPAQQGSWGQPQQPAKQHQPM   151
consensus                                    *!!****!!*!** ! *!*******      ** *!**! *******    ! *       ******  * *  **

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1      ..SHPQEA.......PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1       ..SHPQEA.......PAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1     .YAQNNNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 288539 HYN69 RS09505 WP 108435532.1  RSGGGGGYGGGSSG......GGGGGGGGRANDMDDEIPF   182
NTDB id 1166 A1552VC RS00795 WP 000168289.1  QQSAPQQYS.............QPQYNEPPMDFDDDIPF   177
consensus                                      * ** *        *     *     ********!!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved