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logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 409951 GQR89 RS20695 WP 158771976.1 MASKGVNKVILVGNLGQDPEVRYMPNGNAVANLSIATSESWKDQQ.GQMQERTEWHRLTMYRRLAEVAGEYLRKGSQIYV   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!** !** !*!!!!********!!*!!*!!*!!* *!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENS..GGAPYE..........EGYG..QSQEAYQRPAQQSR.   137
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENS..GGAPYD..........EGYG..QSQEAYQRPAQQSR.   137
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGDFGGNQGSGWNQ..APAQTNYNQGGYSDNYAQNN.....   147
NTDB id 409951 GQR89 RS20695 WP 158771976.1 EGKLQTRKWQDQQGQDKYTTEIIVDQ....MQMLGGKGGGQSEGGFQGQQRPAAAAPQQRAQNSGGQQPGYQQAPNQGYN   155
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQ...QGGMPAQGGMN..VPAQQGSWGQPQQPAKQHQP.....   150
consensus                                   !!****!!*!** !***********      **!!***       !! **         * * **      *****    **

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      .QPASDAP.SHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     .QPAPDAP.SHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ......NFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 409951 GQR89 RS20695 WP 158771976.1 QAPQGGQPNQGFNQAPQGGQNKPAPMAEPDFDFDDDIPF   194
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ..........MQQ.SAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                     *    *    * **** ** * *** !****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved