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AT10033 WP 162467455.1 NC 005838:58866-59282      MRYLLDTNVVSEAVKRRPHPKVVA.....FLRHLKVGEAYLSALTLGELVQGVARAPEARRAILSRWLSGLKASFSGRIL    75
AT10197 WP 001261276.1 NC 023314:c57660-57430     ..........................................................................    0
AT6112 YP 009066163.1 NC 025100:c160993-160763    ..........................................................................    0
AT6111 YP 009066160.1 NC 025100:c156298-155900    ..................MPDICIFMVYAYPHHLHMLNLYICLTSA...YTKHMHTPGAM.................NII    42
AT652 WP 001192482.1 NC 004342:c1008551-1008321   ..........................................................................    0
AT3858 WP 012030622.1 NC 009446:1076865-1077095   ..........................................................................    0
AT3856 WP 012030622.1 NC 009446:296317-296547     ..........................................................................    0
AT1070 WP 005648011.1 NC 000907:c1008816-1008583  ..........................................................................    0
AT4922 WP 005616131.1 NC 010278:1929188-1929418   ..........................................................................    0
AT439 NP 461951.1 NC 003197:c3195787-3195560      ..........................................................................    0
AT10054 WP 066619609.1 NZ CP014301:308578-308829  ..........................................................................    0
consensus
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AT10033 WP 162467455.1 NC 005838:58866-59282      PLDAEVMELWGELTGNAMREGKPLSPLDAMLAATALRHGLILATRNTEEFR.................HVPVPLVNPWEG.   138
AT10197 WP 001261276.1 NC 023314:c57660-57430     ............................MRTVSIFKNGNNRAIRLPRDLDFD.GVSELEIVREGDSIILRPVRPTWGSF    50
AT6112 YP 009066163.1 NC 025100:c160993-160763    ............................MRTVSIFKNGNNRAIRLPRDLDFE.GVSELEIVREGDRIILRPLRPSWSSF    50
AT6111 YP 009066160.1 NC 025100:c156298-155900    P..............DMHHISTLTRERIMRTVSIFKNGNNRAIRLPRDLDFE.GVSELEIVREGDSIILRPVRPTWGSF   106
AT652 WP 001192482.1 NC 004342:c1008551-1008321   ............................MQTAKLFINGRSQAVRLPKEFQFT..GDDVLIQKVGEAVILVPKNKAWNVF    49
AT3858 WP 012030622.1 NC 009446:1076865-1077095   ............................MKVAKVFTTGRSQAVRIPKAFQFD..TKEVIIQRFGNGILLIPKNSDWQGF    49
AT3856 WP 012030622.1 NC 009446:296317-296547     ............................MKVAKVFTTGRSQAVRIPKAFQFD..TKEVIIQRFGNGILLIPKNSDWQGF    49
AT1070 WP 005648011.1 NC 000907:c1008816-1008583  ............................MIEASVFMTNRSQAVRLPAEVRFSEEIKKLSVRVSGSDRILSPLNQSWDSF    51
AT4922 WP 005616131.1 NC 010278:1929188-1929418   ............................MEASVFMTNRSQAVRLPAEVRFASDVKKLSVRVVGNERILAPINQSWDSF    50
AT439 NP 461951.1 NC 003197:c3195787-3195560      ............................MHTTLFFSNRTQAVRLPKSISFPEDVKHVEIIAVGRSRIITPVGESWDSW    50
AT10054 WP 066619609.1 NZ CP014301:308578-308829  ........................MPAKARATTKLFVTNRTQAVRLPKELAFPPEVTEVEIIRQGSGVLIVPKGKGWDDF    56
consensus                                                                 *  ****  *****!*!*****  *   *  **  *  *  *  ***  !*  *!  **
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AT10033 WP 162467455.1 NC 005838:58866-59282      ..........................   138
AT10197 WP 001261276.1 NC 023314:c57660-57430     AQLD.RADPDFMAEREDVVSDEGRFEP    76
AT6112 YP 009066163.1 NC 025100:c160993-160763    LELE.KADPGFMTEREDVVCDEGRFNL    76
AT6111 YP 009066160.1 NC 025100:c156298-155900    LELE.KADPDFMTEREDVVSDEGRVNL   132
AT652 WP 001192482.1 NC 004342:c1008551-1008321   LEGLNGFSDDFFKEEREQPKSDKREKL    76
AT3858 WP 012030622.1 NC 009446:1076865-1077095   LEALNEFEPDFKIEREPQIEQVREDFL    76
AT3856 WP 012030622.1 NC 009446:296317-296547     LEALNEFEPDFKIEREPQIEQVREDFL    76
AT1070 WP 005648011.1 NC 000907:c1008816-1008583  FLNDQAVSDDFMNEREIAFQPE.REAL    77
AT4922 WP 005616131.1 NC 010278:1929188-1929418   FLGSATVSDDFMTEREISTQAE.REEL    76
AT439 NP 461951.1 NC 003197:c3195787-3195560      FDGE.GASTDFMSTREQPAVQE.REGF    75
AT10054 WP 066619609.1 NZ CP014301:308578-308829  FDNGPFASEDFMVTRDQPLAAEEREEF    83
consensus                                          *       ***  ***   *   *  **  *
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X non conserved
X similar
X ≥ 50% conserved