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logo

AT6083 AAN67206.1 AE015451:c1777857-1777558          MLKNGMRPIHPGEILREEFQKEMGFSAAALARALGVATPTVNNILRERGGVSADMALRLSICLDTTPEFWLNLQT   75
AT10161 WP 003133769.1 NZ CP127126:c5517867-5517562  MATNGMRPIHPGEILRDEFLMELDISPAALARALKVSAPTVNDIVREQRGISADMAIRLGRYFDTSAQFWMNLQS   75
AT10032 NP 253363.1 NC 002516:c5242559-5242254       MATNGMRPIHPGEILRDEFLMEFDISPAALARALKVSAPTVNDIVREQRGISADMAIRLGRYFDTSAQFWMNLQS   75
AT171 WP 001232701.1 NC 002506:369724-370041         MRKTKRRPVSVGEMLKVEFLEPMGITSKALAEAMGVHRNTVSNLINGG.VLTAPVAIKLAAALGNTPEFWLNIQH   74
AT6124 WP 005650217.1 NC 007146:c1756423-1756100     ..MMTRKPTSVGEILQEEFLEPLSLKISDLAQILDVHRNTASNIVNNSSRITLEMAVKLAKVFDTTPEFWLNLQT   73
consensus                                            *  ****!***!!*!**!!* ** ** **!!*** !  *!*******  ******!**!** ******!!*!*!*
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logo

AT6083 AAN67206.1 AE015451:c1777857-1777558          AFDLRTAEQQHGDE.IIGSVQRLVA.........   99
AT10161 WP 003133769.1 NZ CP127126:c5517867-5517562  EYSLATAYAANGEQ.IEHEIEPLLAHG.......  101
AT10032 NP 253363.1 NC 002516:c5242559-5242254       EYSLATAYAANGKQ.IEHEIEPLLAHG.......  101
AT171 WP 001232701.1 NC 002506:369724-370041         AVDLWDTRNR..YQEEAKFVKPLFVSLEQSART.  105
AT6124 WP 005650217.1 NC 007146:c1756423-1756100     RIDLWDLEHNKRFQQSLANVKPALHRHDTSTFAM  107
consensus                                             **! **    * * *   * ****
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X non conserved

X similar

X $\geq$50% conserved