

В

TEL top sequence

AML1 top sequence

1 GCCTCTCTGCTATATCCCAGCTGCTTCAGGTTTTCTGAACATGCCAGGCCCCTACCTGCCCCTAGATCCCTGGACATGC 80
81 CTTTACCTCCTCCGAAATTCCCTCGGTCCTCTCTCTCCCTTGGCTAGATACCCCTTGTTAAAGAGAATATTCACATCTGC 160
161 CATGGCTCCCTCAGCCTGGGTAGGCGCCCTGCCCTGTGCATTCTGGTTGGCCCTATGAGTCTGTTCCCTTCTGTGTCACC 240
241 CCCAGTGGGGCCCTTGATGACATGGCTGCCATGCAGTTTACCTCTACATAGCAGCTACGGGCACTTGGCTGACTTTATC 320
321 TGACCAAACTGCATAGCTATCACTATGTGCCCGGCAGCATTCCTAGCAGGGATTACGATGATGATGATGATGATGATGATGATGAT
401 GATGATGGTGATGATGACTGTTATTTCTGTTTTACAGGTGGAAAAGCAGAGGCACAAAGCGGTGATGTGACTTGCCTGAA 480
481 GTCACACGGCTACAAAGTGGTAGATAGGGGAACGGCTCCCACGTCTGTATGGTCTGAACTGGGTTGTGTAGGTTCCCCA 560
561 AAGTTCATGTCCACATAGAACCTCAGAATGTGACCTTTATTTGGATATAGCGTTTTTTGAAGAAGAAAAAACGGACAGACGAGGAAGA 720
721 CCTCATATGATGCTGGAGGCAGAACTGGGCTGATGTAGCCCAGAAGCTAAGGAGTGCTGGCAACTGCTGTAAGGTAGAA 800
801 AGAGGCAAGGAAGGATTTCCCCCTAGAGCCTGCACAGGGGACAGACCCTTGATTTTGGACTTCTGGCGT 880
881 CCAAAGCTGTGAAAATAATTTCTGTTTTTTCAGCCACCCAGTTTGTAGTAATGTGTTATGGCAACATGGGAAA

Figure 2 Numata et al

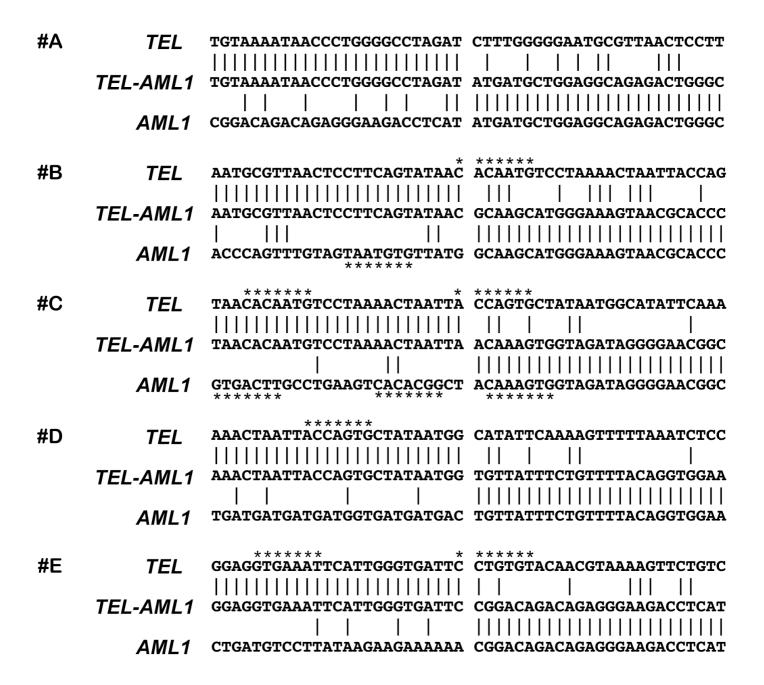
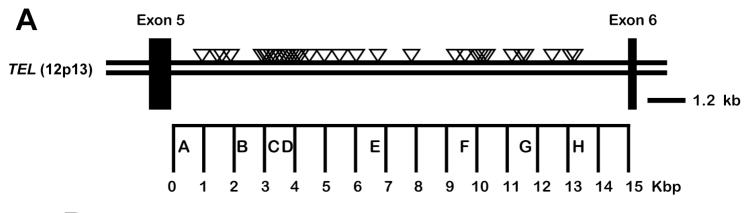
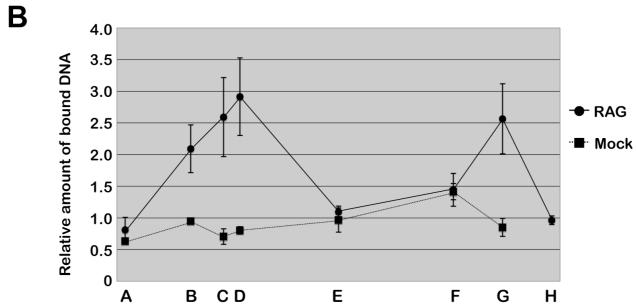
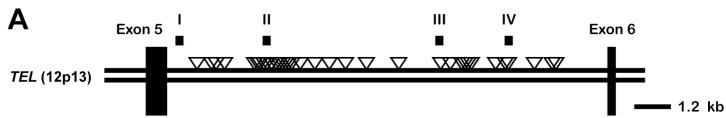


Figure 3 Numata et al





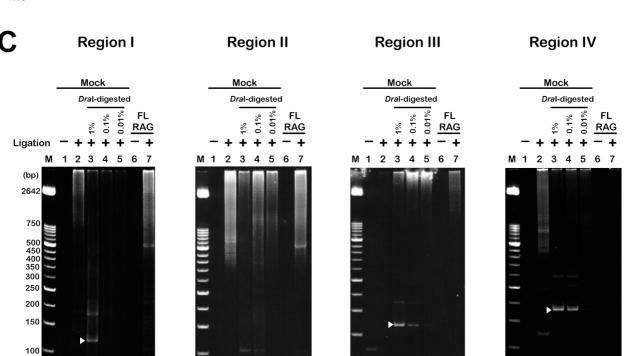




B Region I Region II Region III **Region IV**

26 TGGGATGGAGTAGTTAATGAGCCTC TTTTTGGGGGAATGCGTTAACTCCTT AAATTCCTATTGCTGTCTCCGTTAG GCAATAGATACGGATTGCTCCCACAG 51 AGAAATGTTAAGAAACAAATGTCCT CAGTATAACACAATGTCCTAAAACT TGACTAATATGTCCTGAATG TTCTGGGGGCTGGGGAGATCAAGGC 76 ACGTCCAGCTTACAAGGAGAGTCAC AATTACCAGTGCTATAATGGCATAT TCAACGTATTTTATTACTCAGCTA CCTGGCAGATGAGGCATCTGGTGAG 101 ATCAGAATCAAGGCTAAGCGAAAAC TCAAAAGTT**TTTAAA**TCTCCCAATA AAGTTGGCTTTGCTGGATTAAATGT GGCCCACTTTCTGGTTCATAGACAG $126 \ \mathtt{ATTTAAA} \mathtt{ATAAAAGGTTTATGAGCA} \quad \mathtt{TATATTTAACTAATGCAACCAAAT} \quad \mathtt{CATTTAAA} \mathtt{ATGATCAATTGGCAGCT} \quad \mathtt{CACCTTCTTGCTGTGTCTTCACTTG}$ 151 TGCTAATGGCCCTGTCCTGAAGTTT CCAAATTTAACTCATTCCACTGAAA TGTTTGGGGTTGTTTTCACATTTAT GTGGAAGGGTGAGGGGTCTCTCTCA

1 CCTAATGGCTTGGTTAATGAGCCGC GTAAAATAATCCTGGGGCCTAGATC GCTTGAAACAACGCTTTATGCCACA CTGCCGTAGACTGGGTAGTT<u>GGTAC</u> 176 CCAGCAGCTAATTAATGACCAGACA GGCATTGTTTTTCTCCGAAGAAGGT GAAAGGAAAAGTTTGCCGGATACAA GGCCTCT**TTTAAA**AGGGCACTAATC



* 3.4%