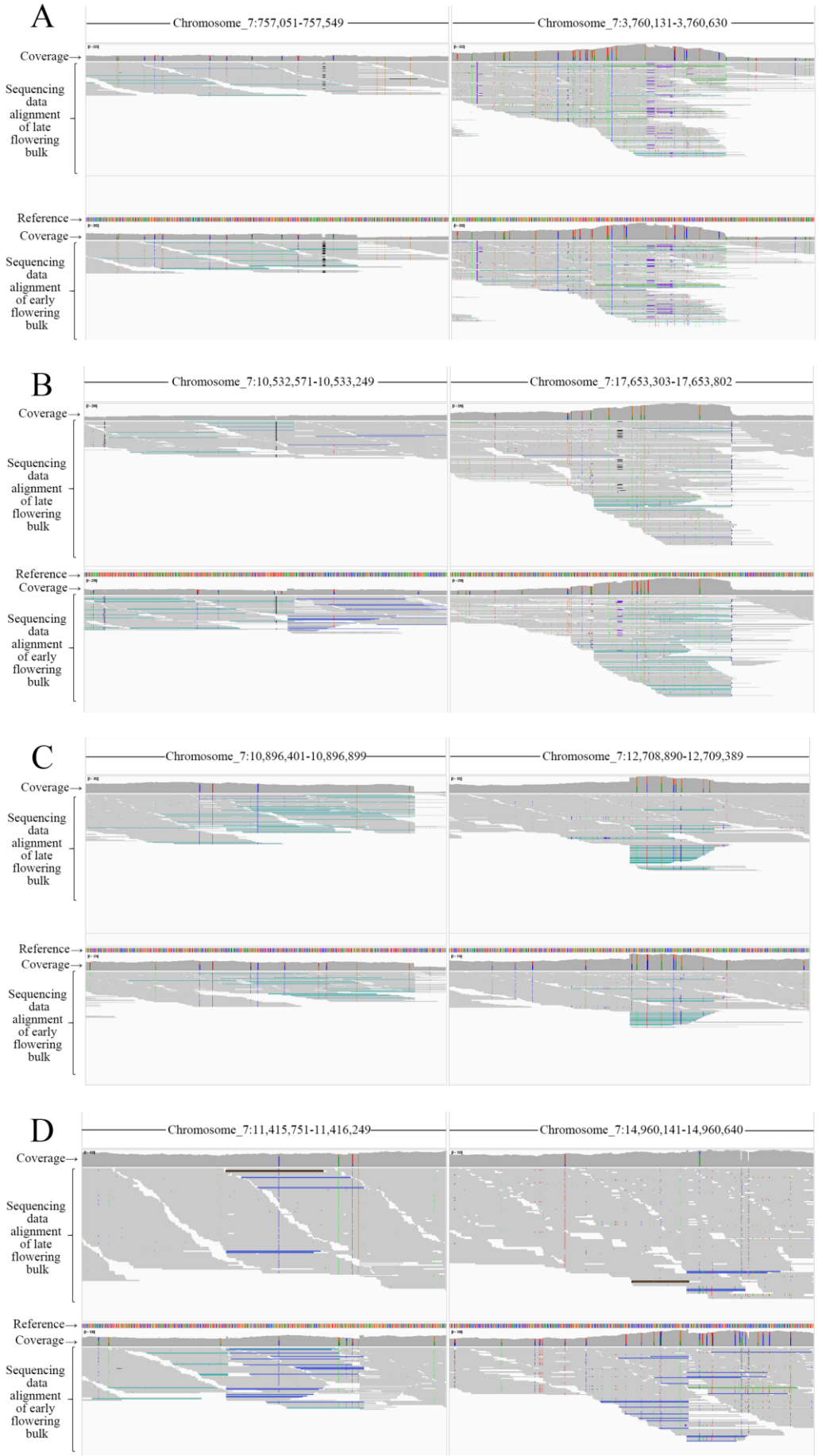

A
Chromosome_7:757,051-757,549
Chromosome_7:3,760,131-3,760,630
Coverage→
Sequencing data alignment of late flowering bulk
Reference→
Coverage→
Sequencing data alignment of early flowering bulk
B
Chromosome_7:10,532,571-10,533,249
Chromosome_7:17,653,303-17,653,802
Coverage→
Sequencing data alignment of late flowering bulk
Reference→
Coverage→
Sequencing data alignment of early flowering bulk
C
Chromosome_7:10,896,401-10,896,899
Chromosome_7:12,708,890-12,709,389
Coverage→
Sequencing data alignment of late flowering bulk
Reference→
Coverage→
Sequencing data alignment of early flowering bulk
D
Chromosome_7:11,415,751-11,416,249
Chromosome_7:14,960,141-14,960,640
Coverage→
Sequencing data alignment of late flowering bulk
Reference→
Coverage→
Sequencing data alignment of early flowering bulk

Figure S1 Visualization of four other inversions on chromosome 7 detected from Illumina sequencing data mapping files of two extreme bulks for flowering time in an F2 population between *Melastoma candidum* and *M. normale* using Integrative Genomics Viewer. To visualize, we used the options of "set color alignments by pair orientation" and "view mate region in split screen". Paired-end reads with blue and cyan support inversion. The region ranges containing the inversion breakpoints on chromosome 7 are indicated on the top of each panel.