



Supplemental Fig. 7 A complex event (deletion + inverted spacer). (a) Dot plot of the alignment of the contig to chr3:162807456-162830372 on NCBI build v38 using nucmer (Delcher et al. 1999) (b) Screenshot of Integrative Genomics Viewer (IGV) of reference alignment of Illumina reads on the same focal region.

Reference

Delcher AL, Kasif S, Fleischmann RD, Peterson J, White O, Salzberg SL. 1999. Alignment of whole genomes. *Nucleic Acids Res* **27**: 2369-2376.