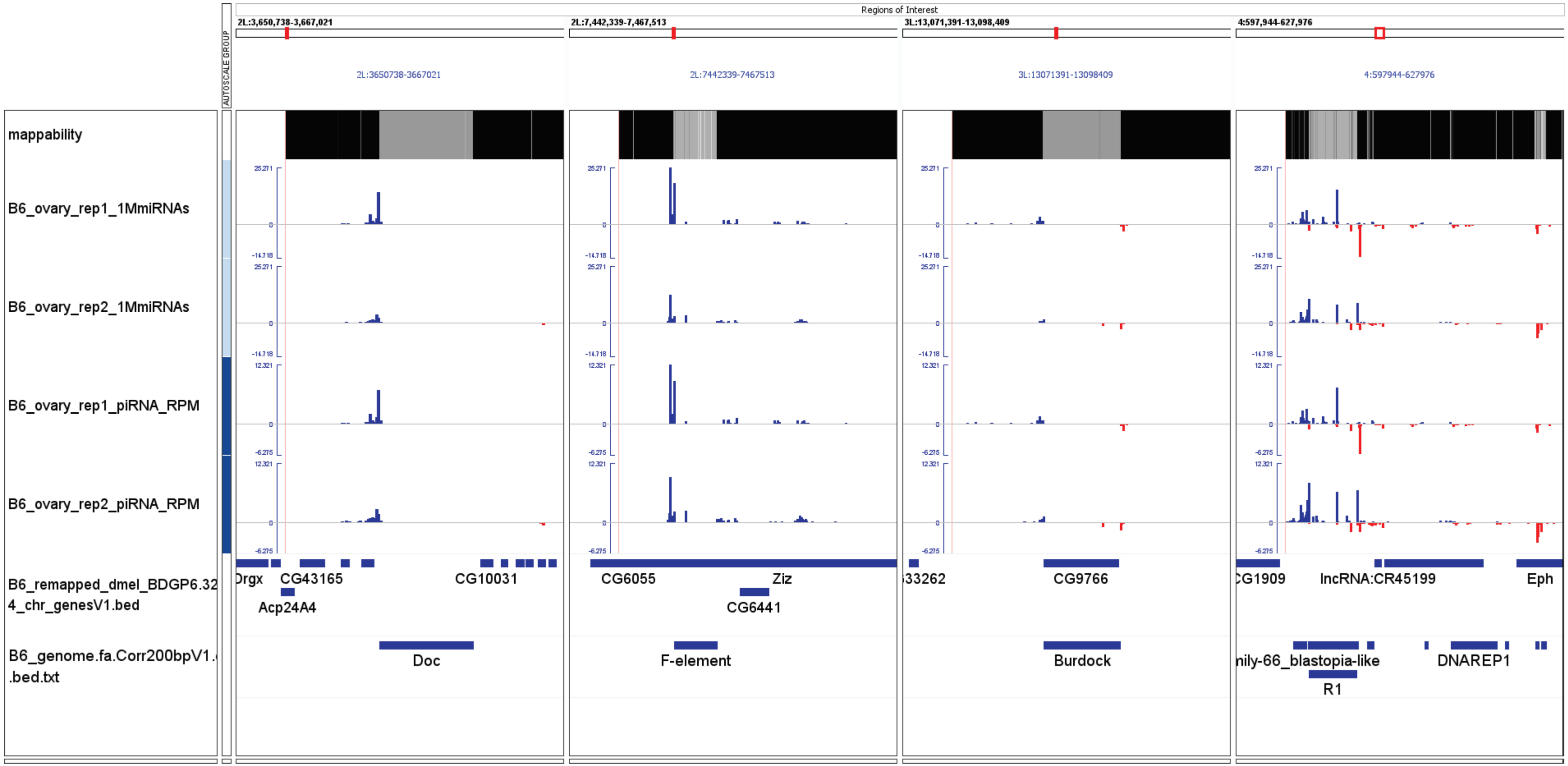

Supplementary_Fig_S6. IGV browsershots of examples of four distinct piCs with miRNA and total read depth normalized piRNA coverage tracks for strain B6. *Top to bottom* - Theoretical mappability scores for 25nt alignments are shown in 500bp windows. Next two tracks display, piRNA read coverage tracks for B6 small RNA library replicates normalized to per million miRNA reads. Next two tracks display, piRNA read coverage tracks for B6 small RNA library replicates normalized to per million total piRNA reads sequenced. Next, remapped iso-1 gene annotations are displayed in B6 genome coordinates. Finally, RepeatMasker annotations of TE insertions (>200bp) are shown to highlight TE insertions associated with each piC region.