



Figure 19: Position and population frequency of *P-element* insertions in replicate 1 based on short-read data. The contigs are in the top panel and the generations in the right panels. *P-element* insertions in piRNA clusters are shown in red. The likely order of *D. erecta* contigs was inferred by aligning the contigs to the *D. melanogaster* genome using Mauve [Darling et al., 2004]. m million base pairs