



Figure 10: Length distribution of small RNAs mapping to the *P*-element. The abundance of small RNAs was normalized to 1 million piRNAs (smRNAs/mil.piRNA). Data are shown for three replicates (top panels) and several generations during the invasion (right panel). Sense RNAs are on the positive y-axis and antisense RNAs on the negative y-axis. small RNA data were either generated for whole bodies of female flies (light grey panels) or ovaries (dark grey panels).