

**S9 Fig.** Allele-specific differential expression at neo-X and neo-Y SNP sites. Left panels, allele-specific read counts over strain-specific SNP sites (points) differentiating the neo-X and neo-Y chromosomes were used to calculate the fold difference. Right panels, histograms of the distribution of the log<sub>2</sub> fold differences. Red lines demarcate the median fold difference.