

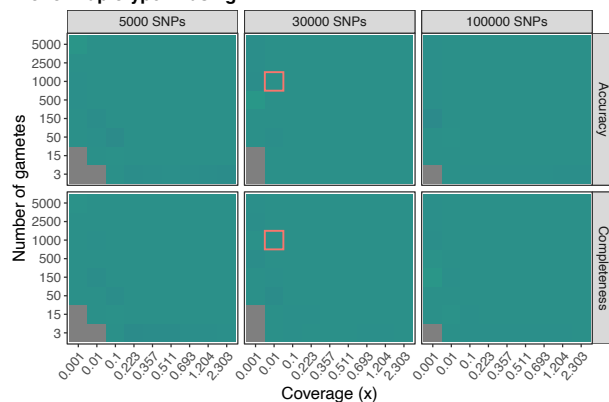
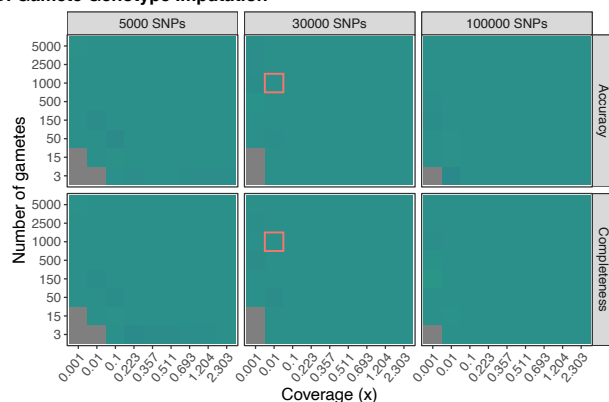
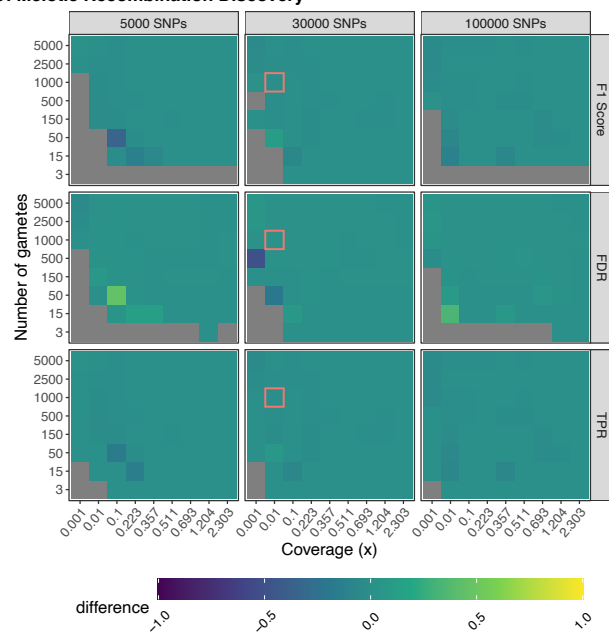
Overestimated Genotyping Error Rate**A: Donor Haplotype Phasing****B: Gamete Genotype Imputation****C: Meiotic Recombination Discovery**

Figure 2-figure supplement 8 Model robustness when genotyping error is overestimated. Plotted values are the difference between the mean performance (across 3 independent trials) when the generative model and rhapsodi use the same parameters (Fig. 2) and the mean performance (across 3 independent trials) when the rhapsodi genotyping error rate (0.005) is overestimated compared to the rate used by the generative model (0.001). The average recombination rate is 1 per chromosome in both models.