



Figure 2-figure supplement 6 Model robustness when genotyping error is underestimated. 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 (0.005) is underestimated compared to the rate used by the generative model (0.05). The average recombination rate is 1 per chromosome in both models.