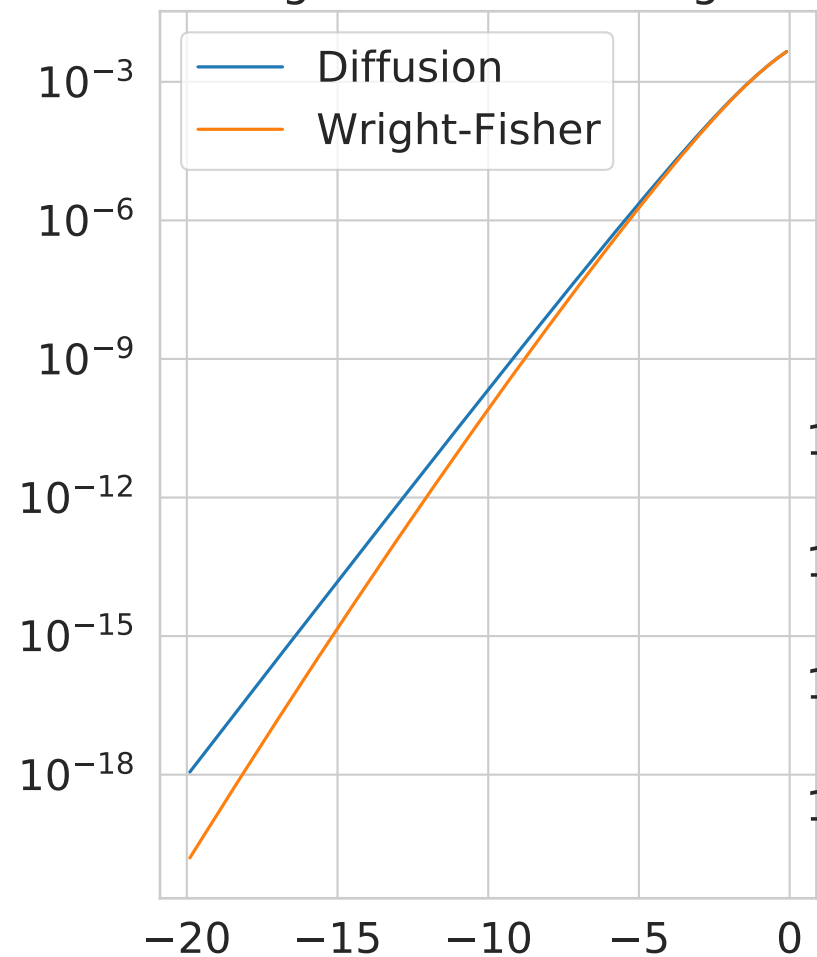
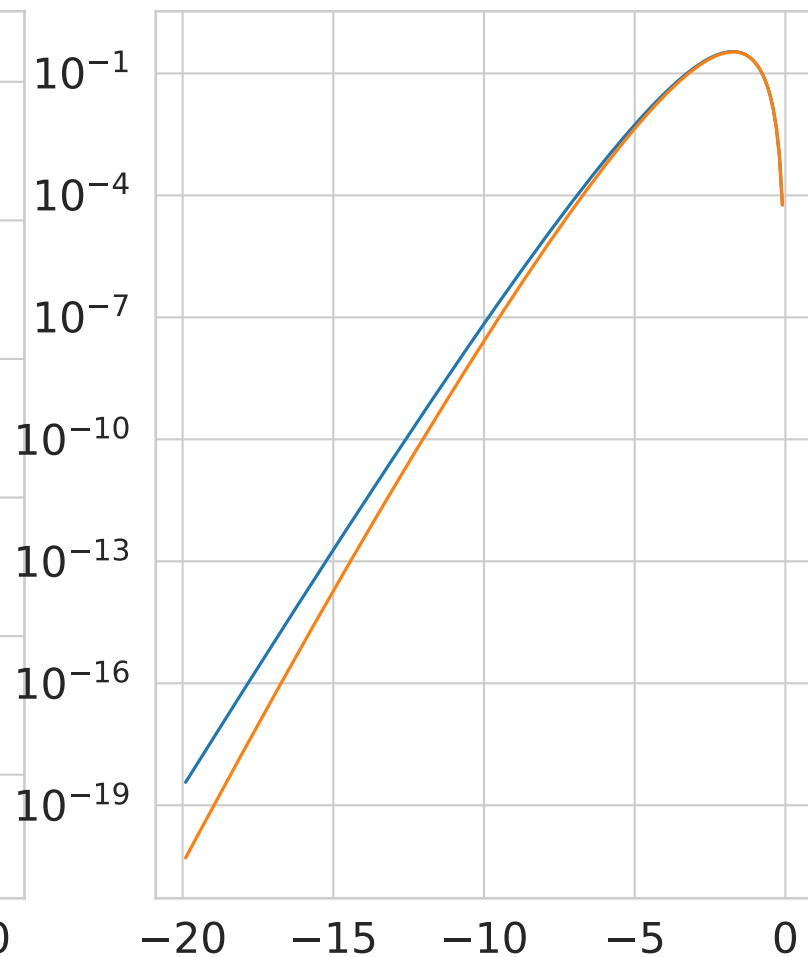


A Probability of fixation
of a single variant in the genome



B Fixation load



C Relative error of fixation probabilities

