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FIGURE S1.—Log$_{10}$ *P*-value of the goodness-of-fit test comparing the *H$_{pair}$* statistic under different SNP ascertainment schemes (shown on the x-axis) to that with complete ascertainment for the complex demographic model. Here a sample size of 40 chromosomes from each population is used. The solid horizontal line denotes the 5% significance cutoff. *P*-values <10^{-200} are set to 10^{-200}.