FIGURE S6.—Distribution of MLEs for the three growth parameters inferred using the SFS from Pop AA using different size data sets. “Original” denotes the original data sets presented in Figure 5 using a sample size of $n = 24$ chromosomes. “Filtered” denotes data sets consisting of a sample size of $n = 100$ chromosomes. Here we filtered SNPs so that the number of SNPs per data set was approximately equal to the mean number of SNPs in the original data sets. “Full” denotes the data sets consisting of a sample size of $n = 100$ without filtering any SNPs. Solid horizontal bars denote the true parameter values for Pop A ($N_A = 20,000; N_B/N_A = 0.5; t_{cur} = 4000$ generations).