



Figure S2: Correcting the underestimation of observed heterozygosity for simulated data with population structure. In each panel, a purple bar indicates the uncorrected observed heterozygosity averaged across all individuals in a simulated data set after applying allelic dropout; a green bar indicates the “true” observed heterozygosity averaged across all individuals in the same simulated data set before applying allelic dropout; and a striped black bar indicates the corrected observed heterozygosity averaged across all individuals and across 100 imputed data sets. The x-axis indicates values of the inbreeding coefficient that were set for different simulations. Different panels correspond to different values of the F parameter in the F -model for simulating structured populations. (A) $F = 0$; (B) $F = 0.04$; (C) $F = 0.08$; (D) $F = 0.12$; (E) $F = 0.16$; (F) $F = 0.20$.