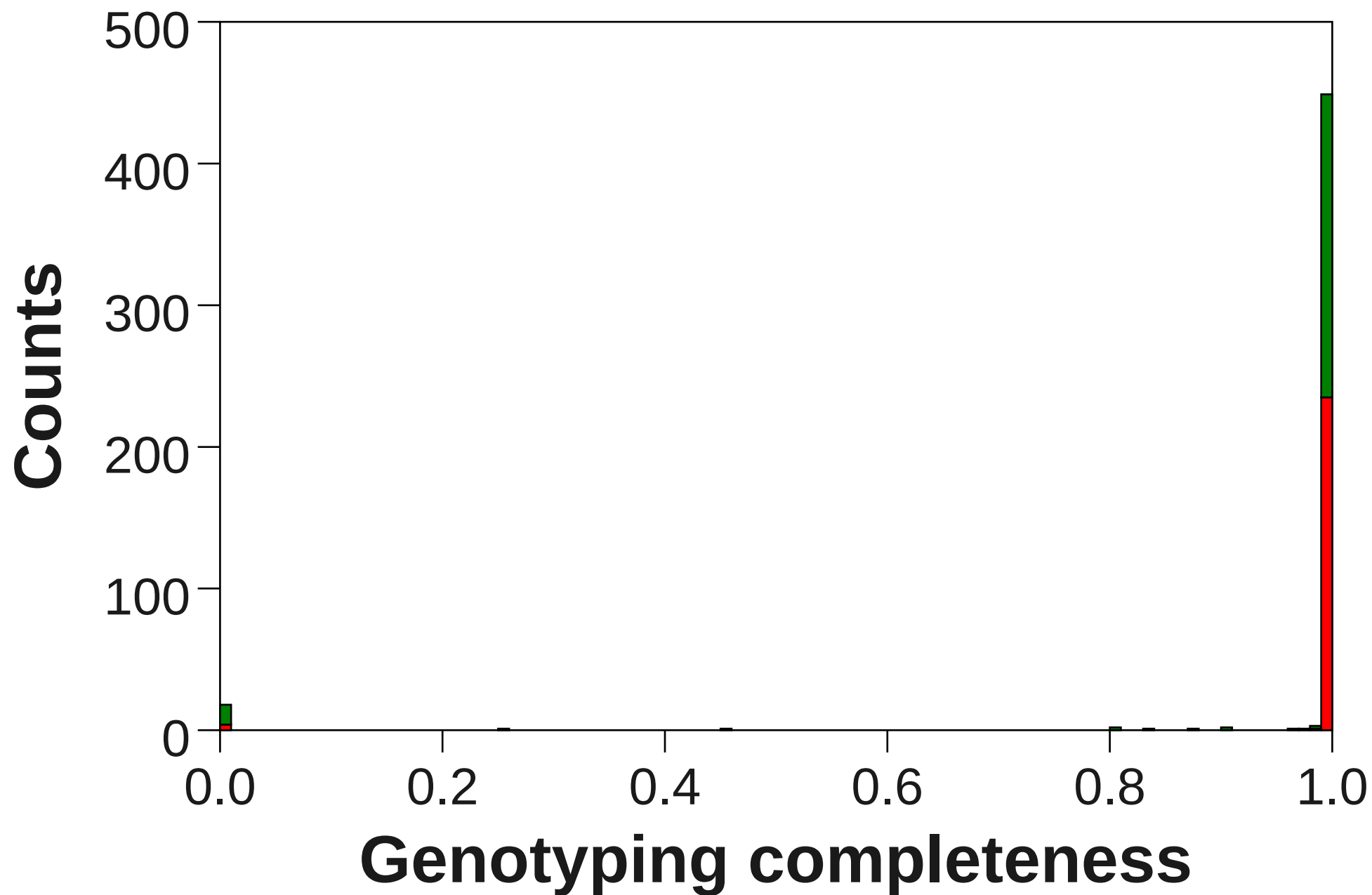
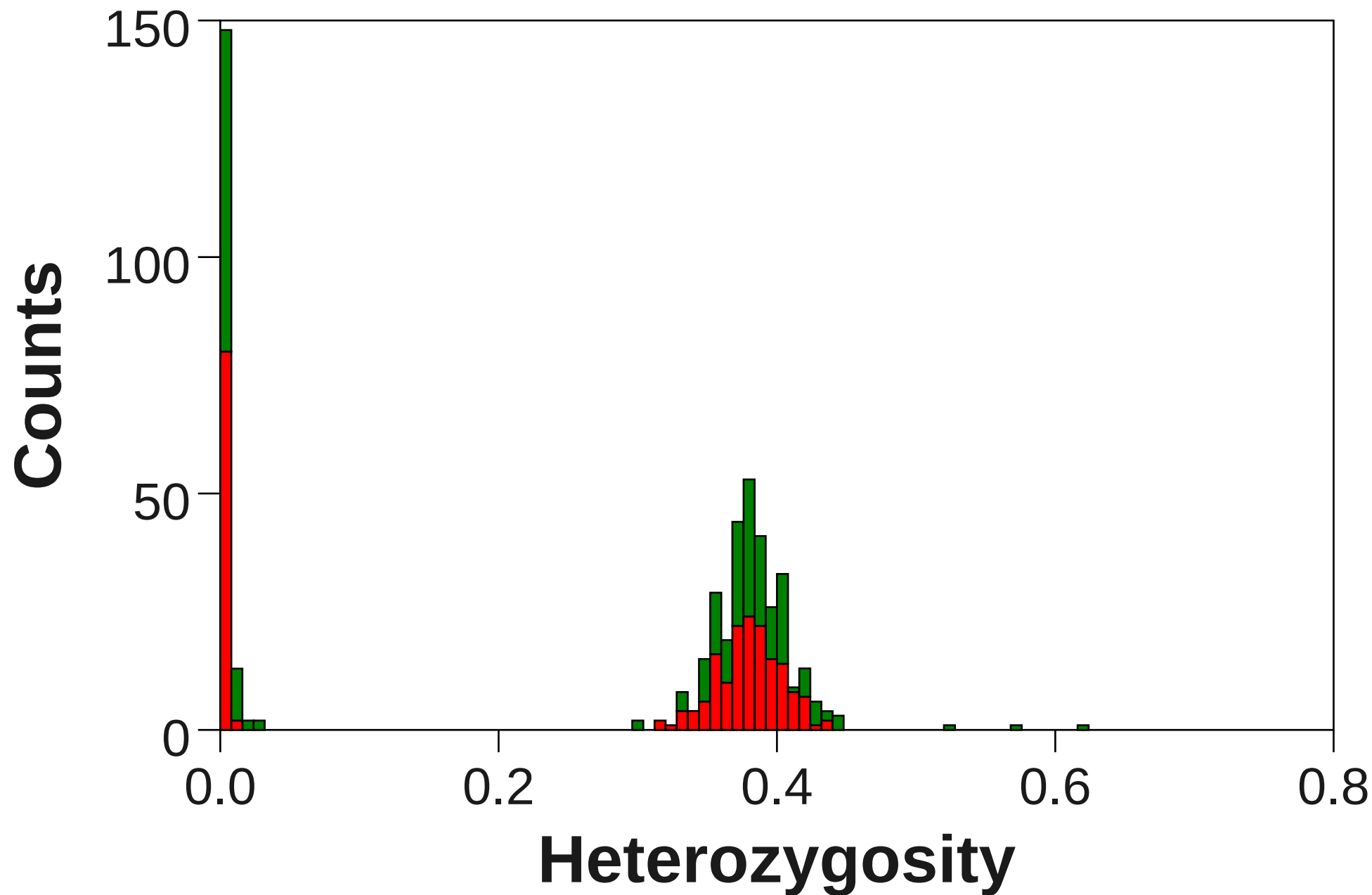


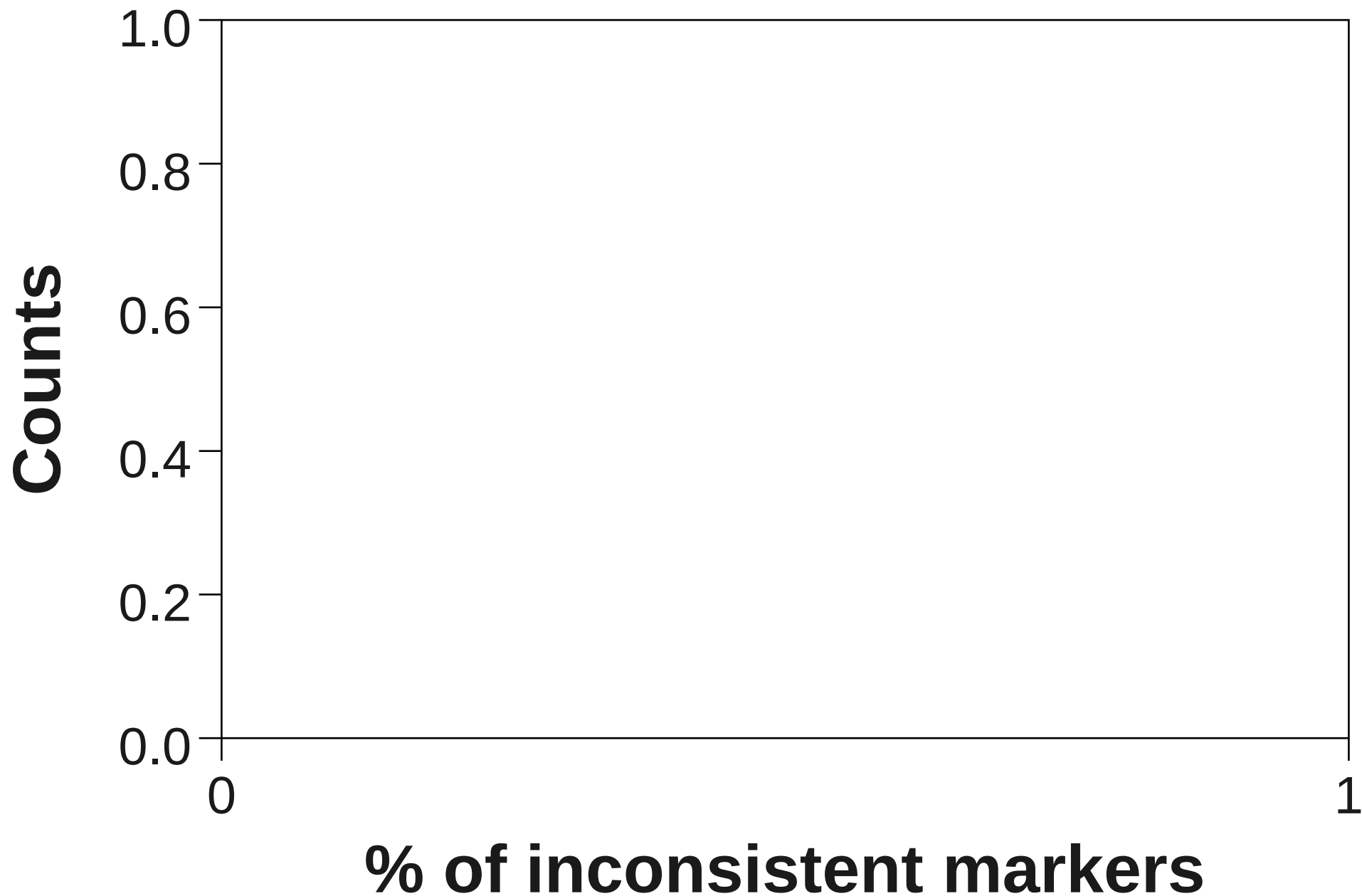
Sample genotyping completeness



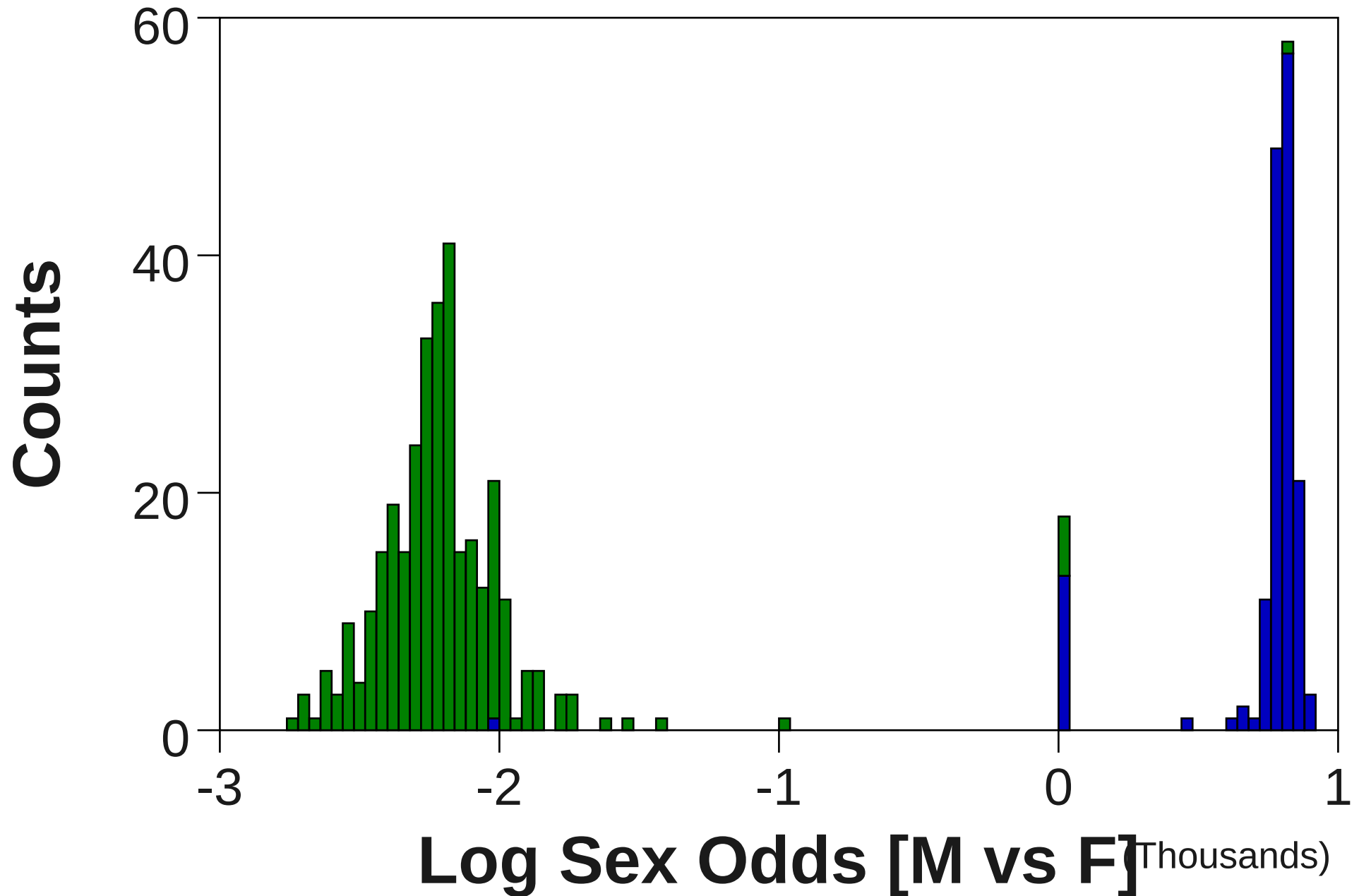
Sample heterozygosity



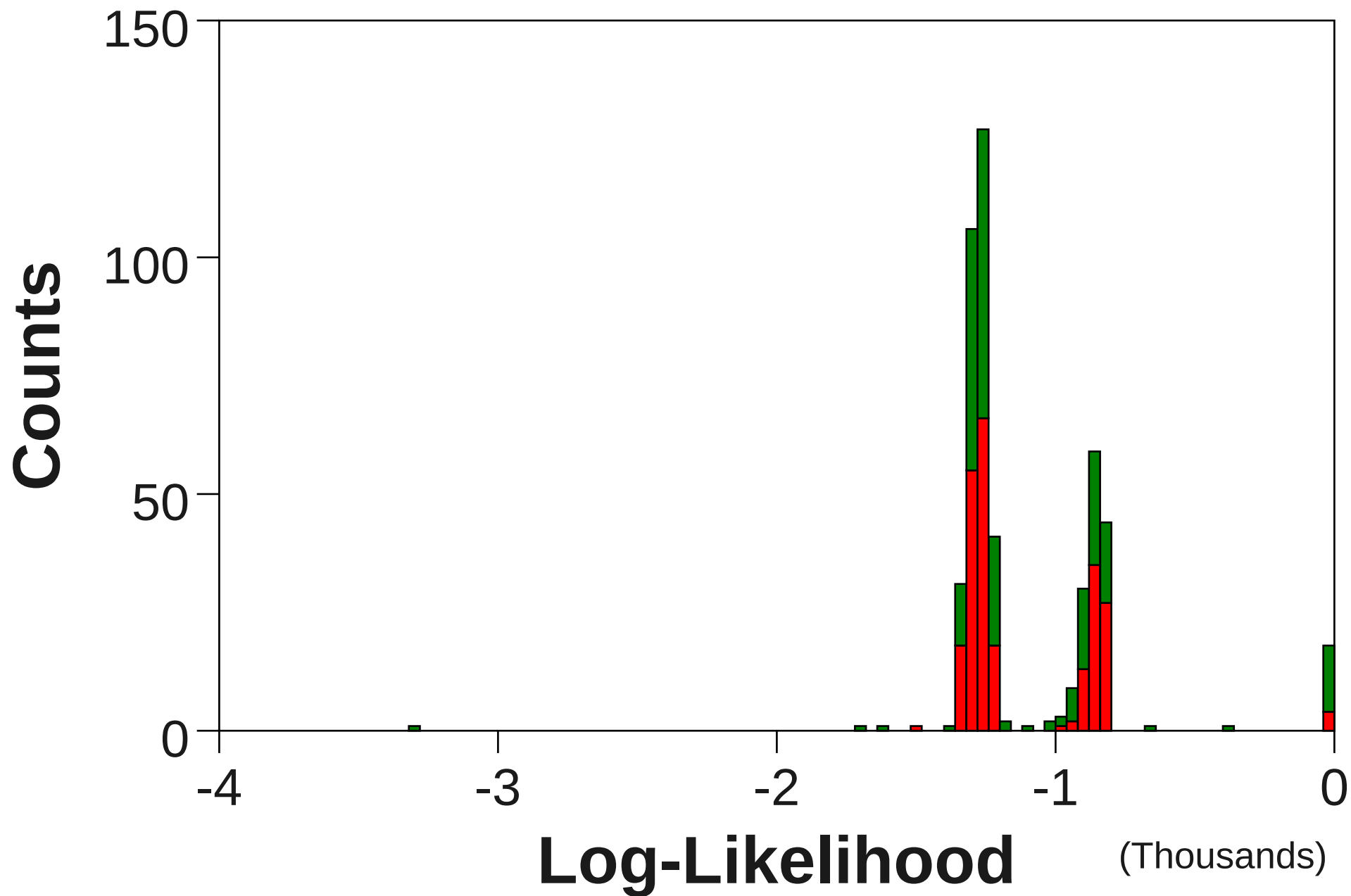
Sample mendelian inconsistencies



Sample Log Sex Odds



Sample Log-Likelihood



Sample Average Quality Score

