

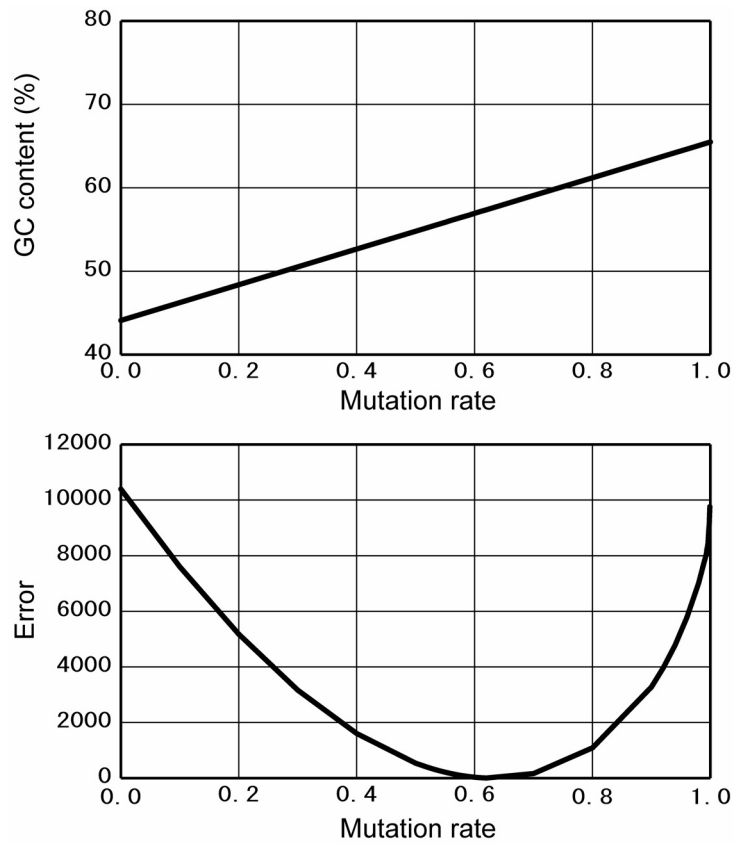


Figure S1. Variations of GC content (upper) and prediction error (lower) as a function of mutation rate in *A. thaliana*.

GC content and error are averages of 20 simulations for mutation correction. Error is measured as the χ^2 value between the observed and predicted intron phase distributions.