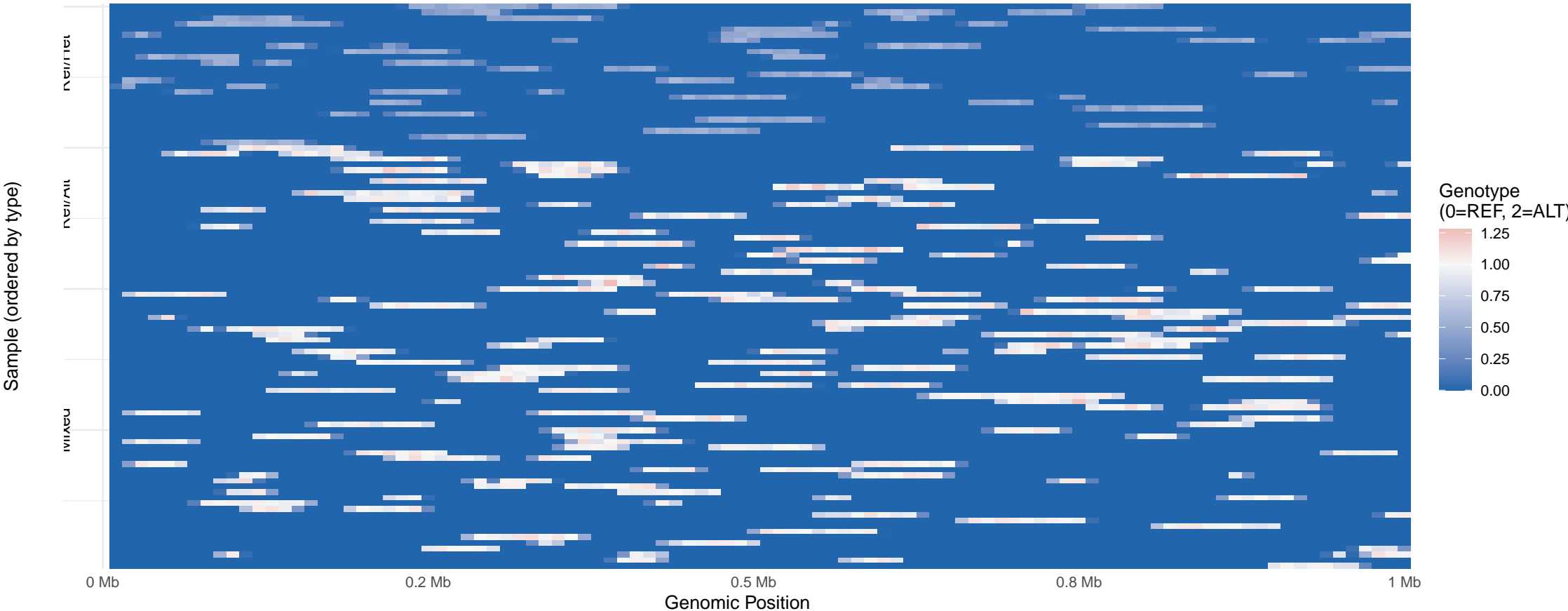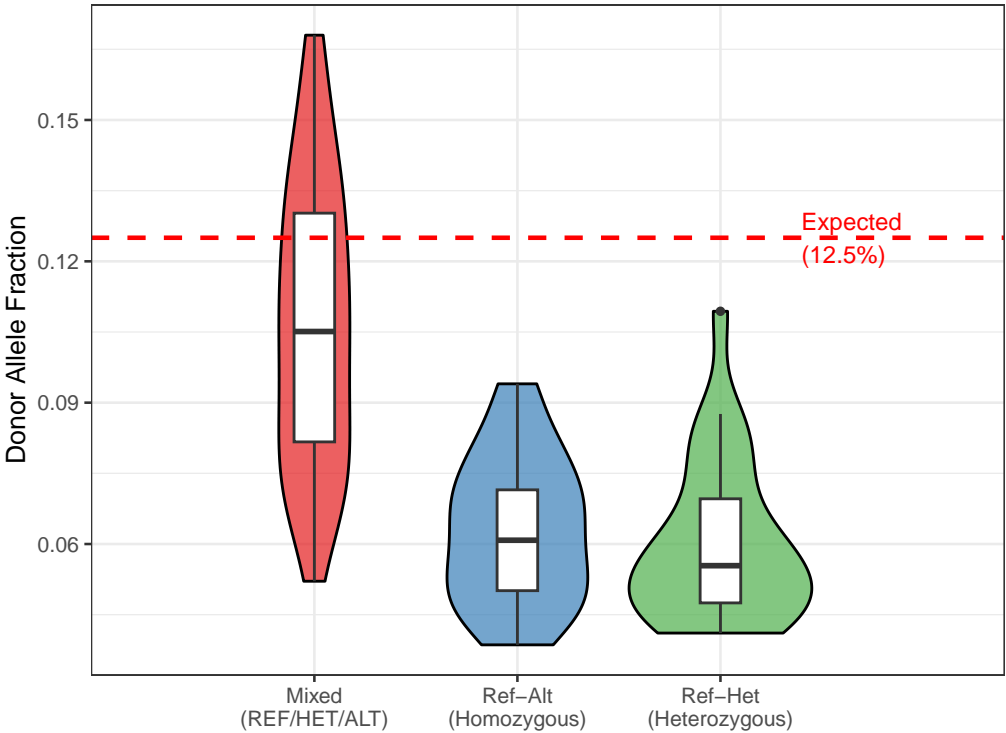# Figure 5: Cross–Species Introgression Simulation

BC2S3–like population: 100 samples, 1000 kb genome, 10000 SNPs
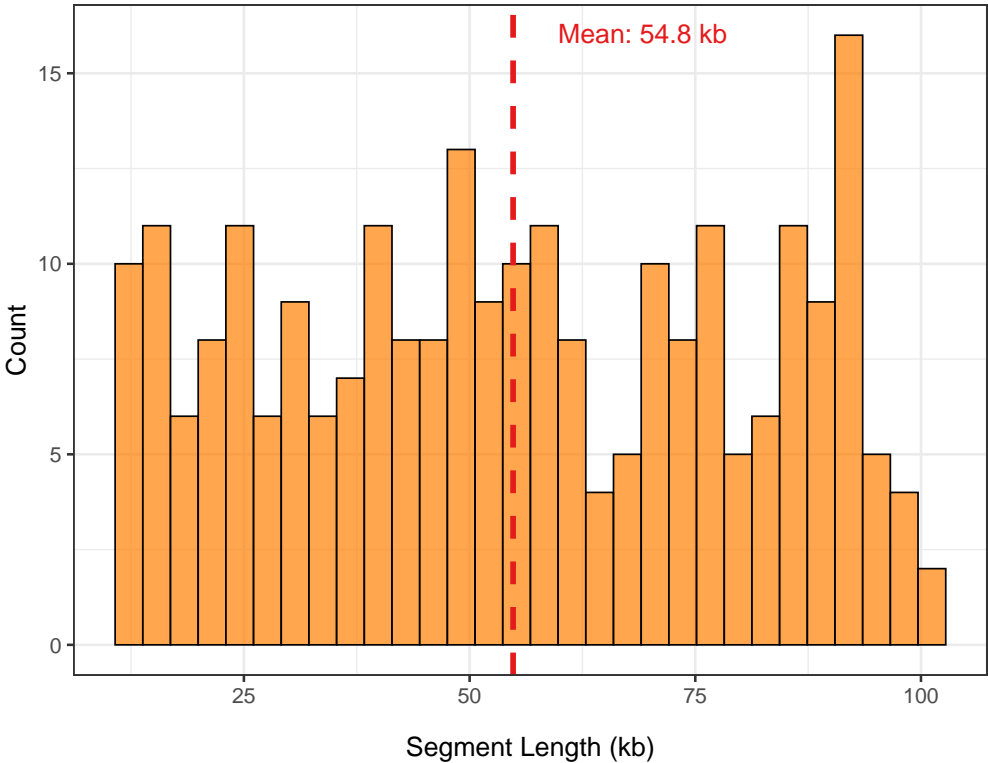
A. Introgression Landscape Across Samples

B. Donor Fraction by Sample Type

C. Introgression Segment Size Distribution

D. Genotype Calling Accuracy vs Coverage