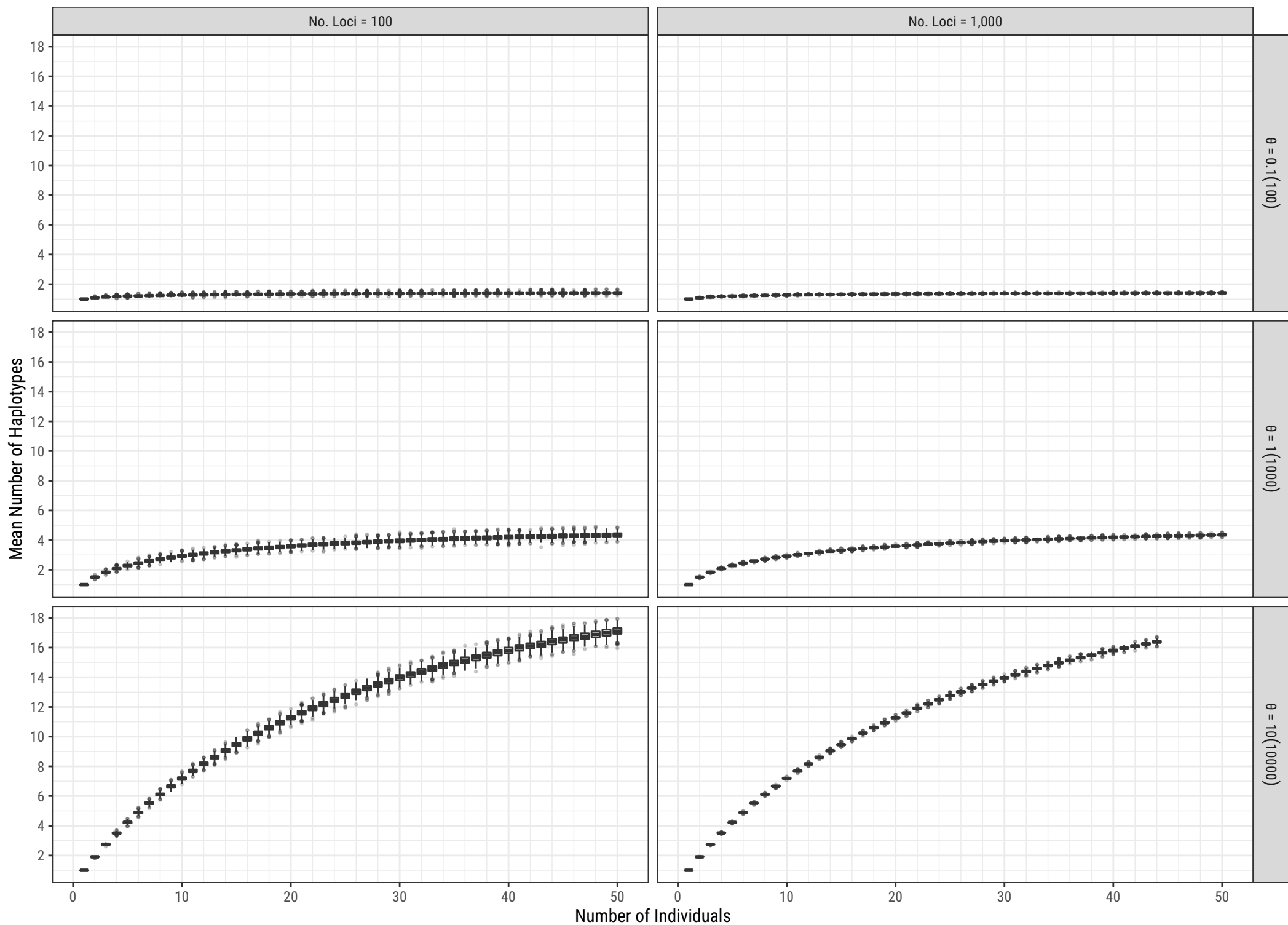Simulations of Mean Haplotypes at 100x Coverage

based on 1000 replicate simulations for each
parameter combination, performed in the program *ms*