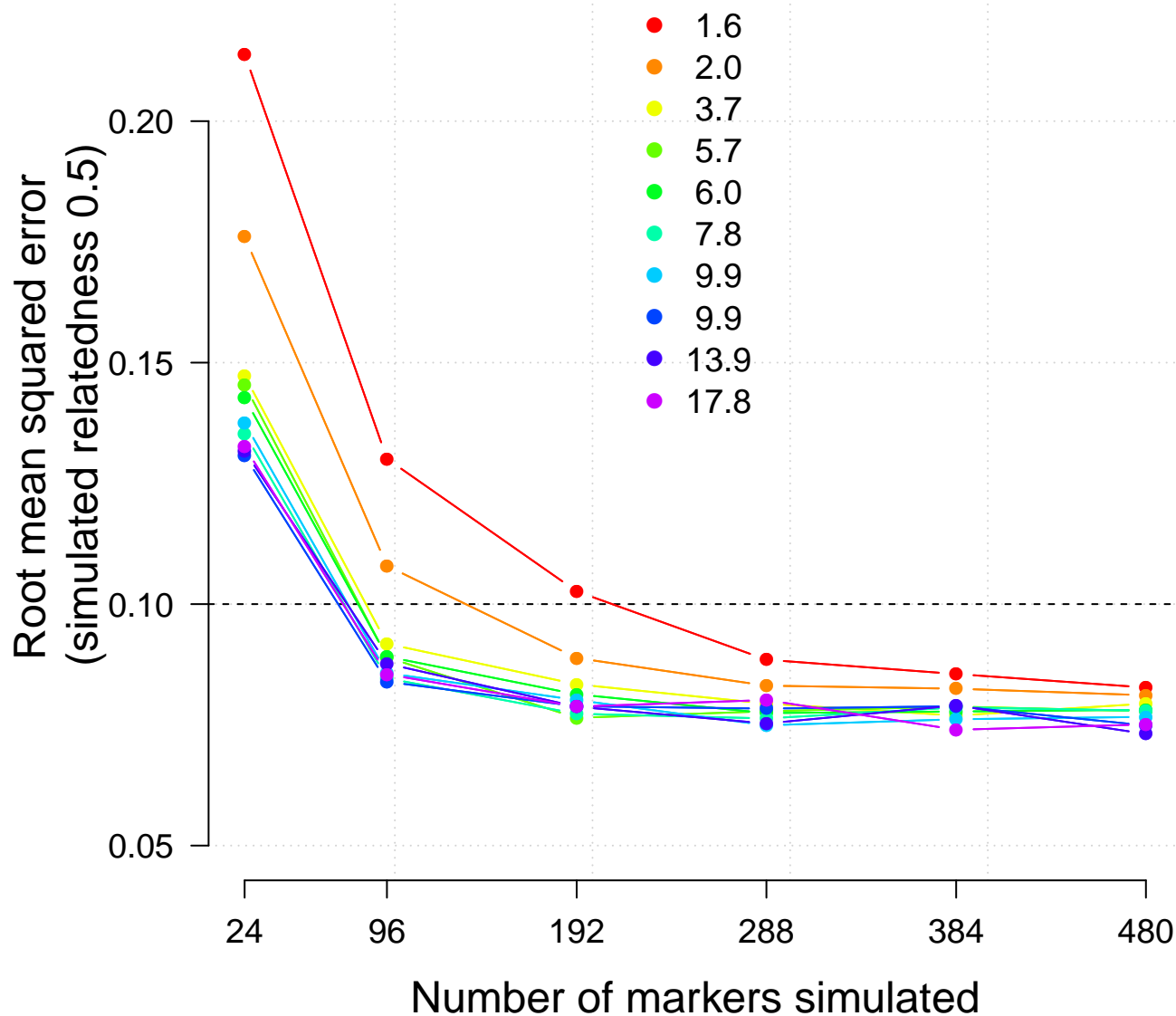


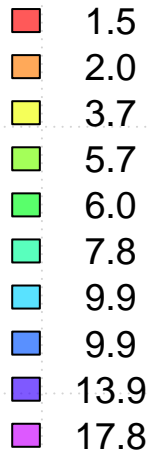
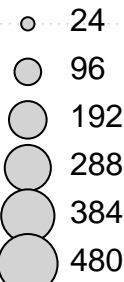
Average effective number of alleles simulated



Average effective number of alleles simulated

Root mean squared error  
(simulated relatedness 0.5)

Marker count



0.20  
0.15  
0.10  
0.05

0 2000 4000 6000 8000

Number of markers simulated multiplied by  
the average effective number of alleles simulated

