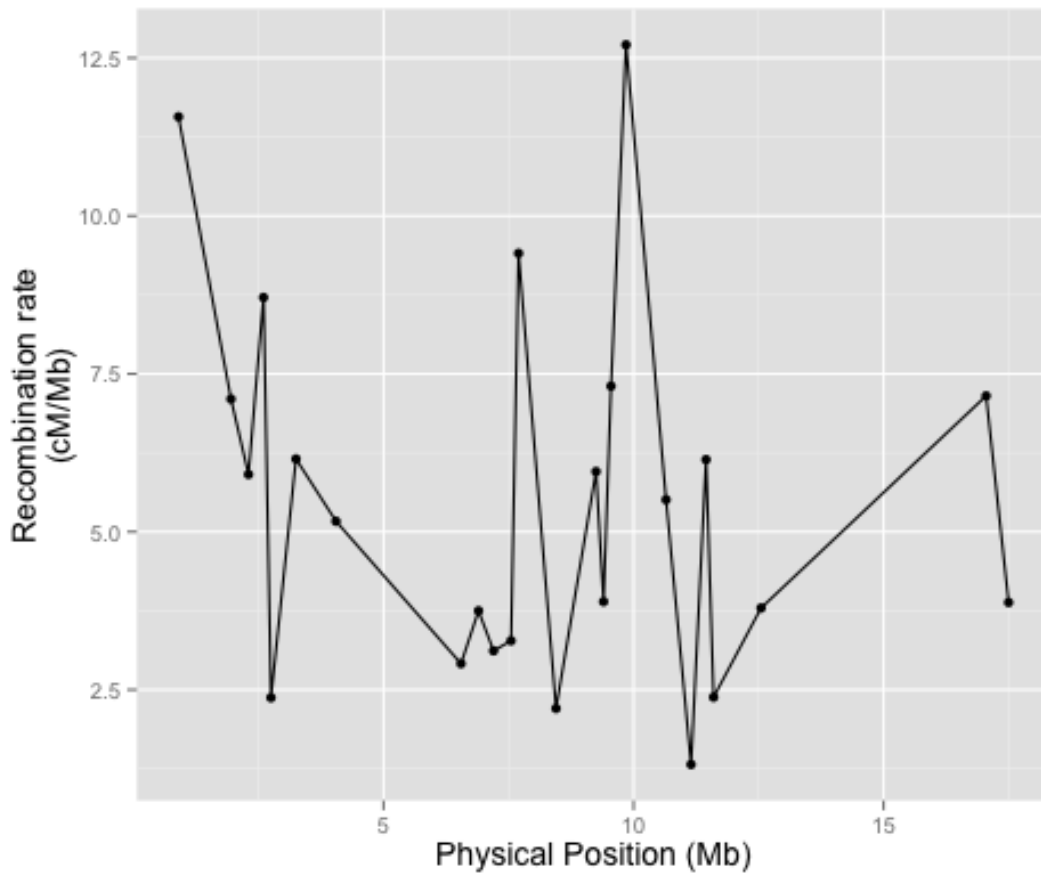
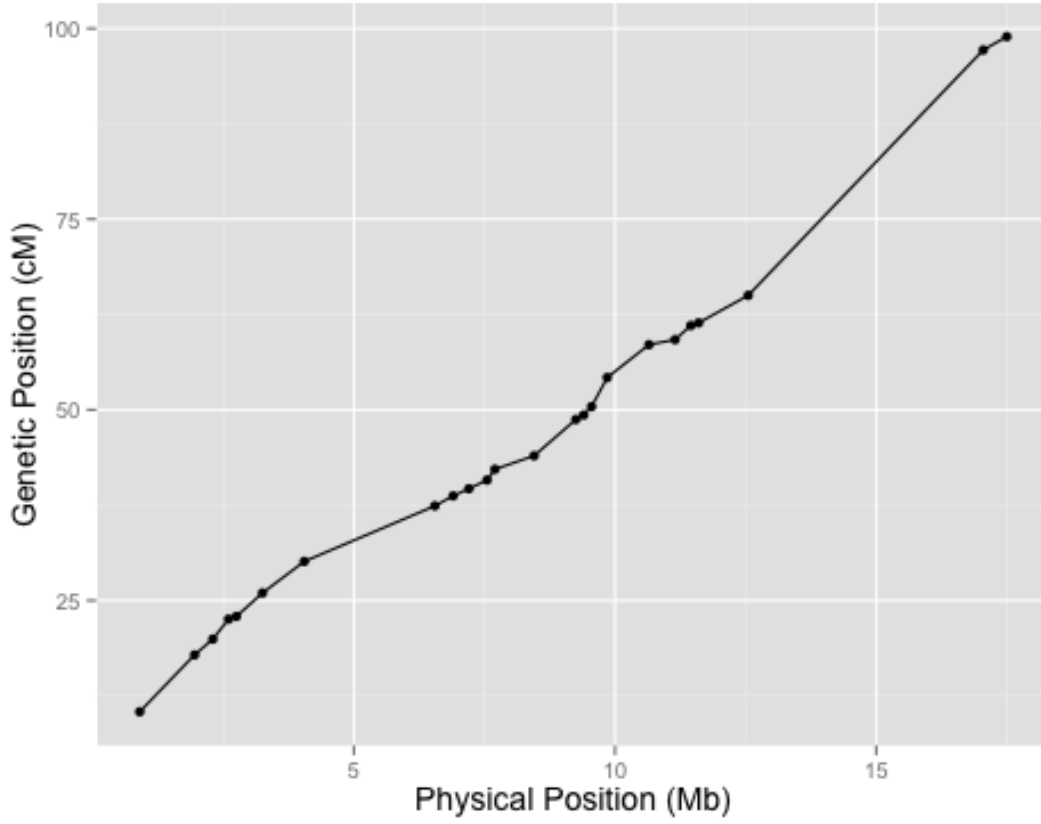
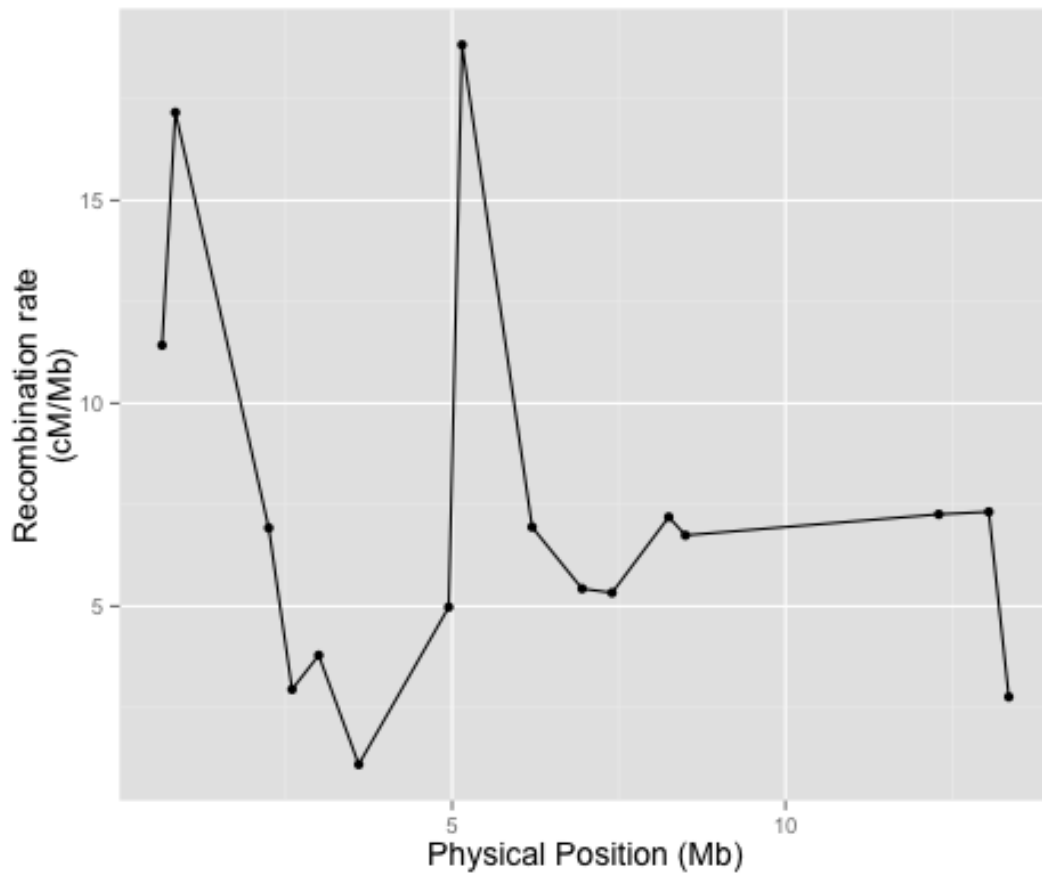
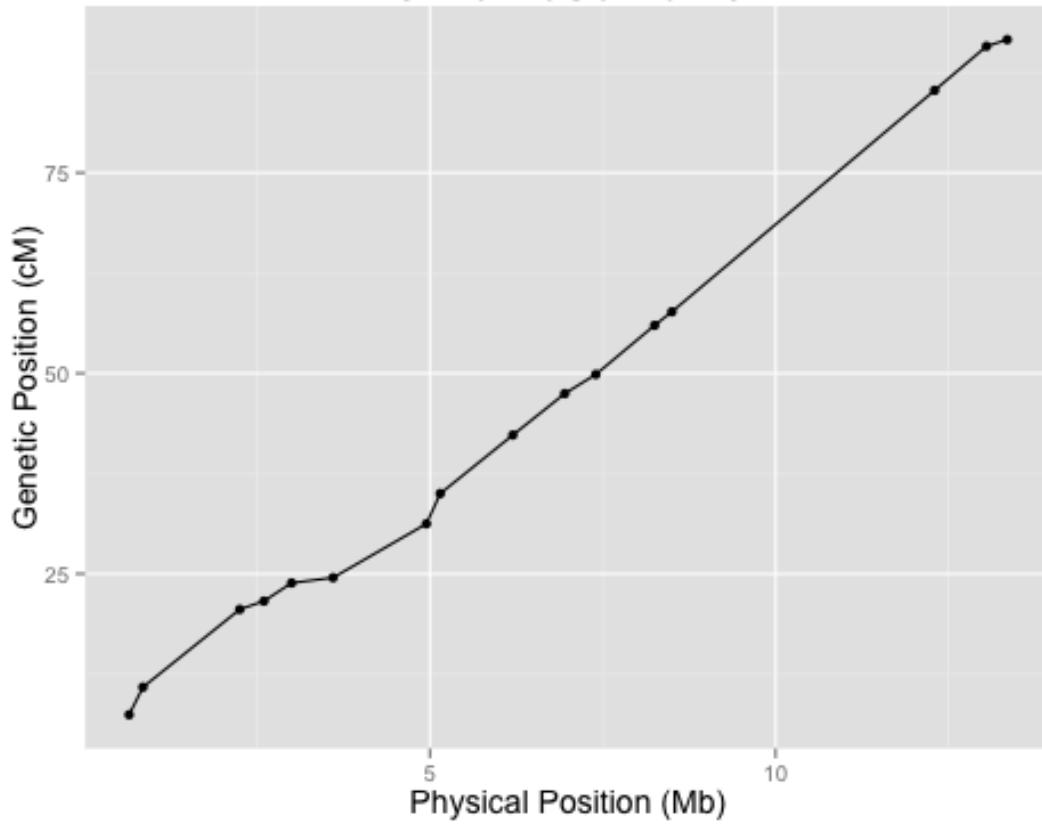


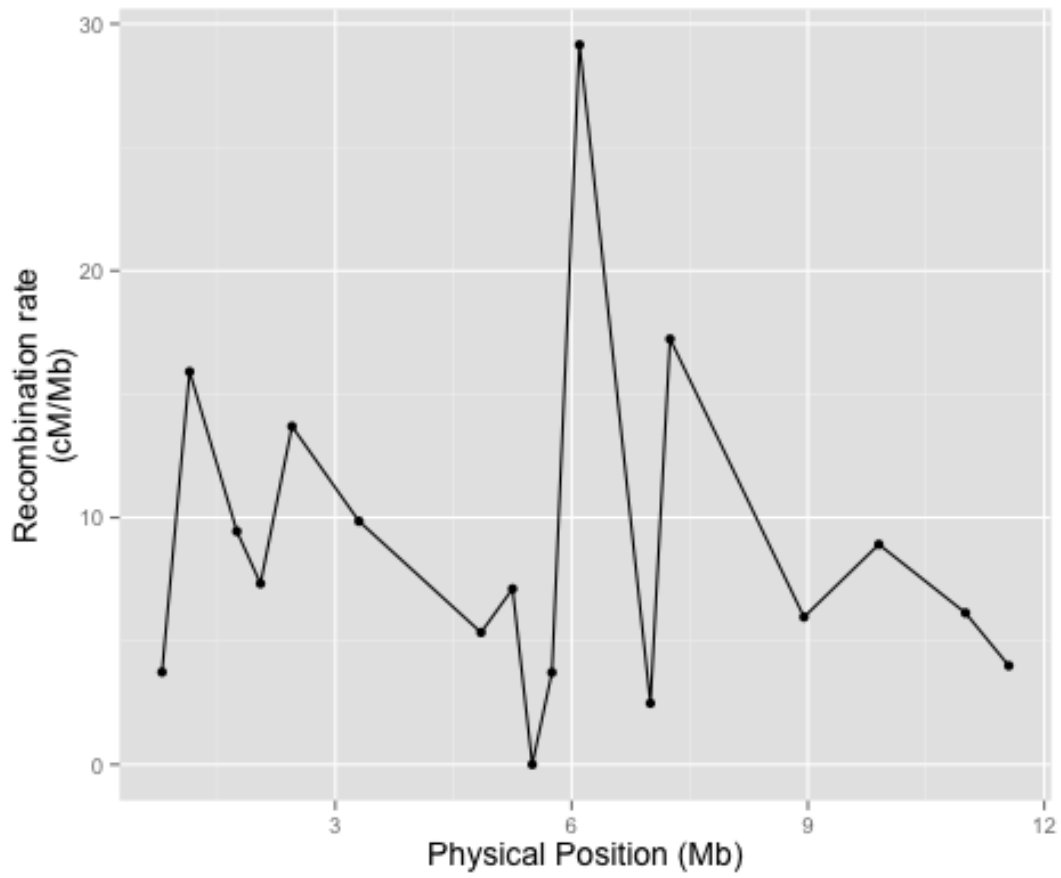
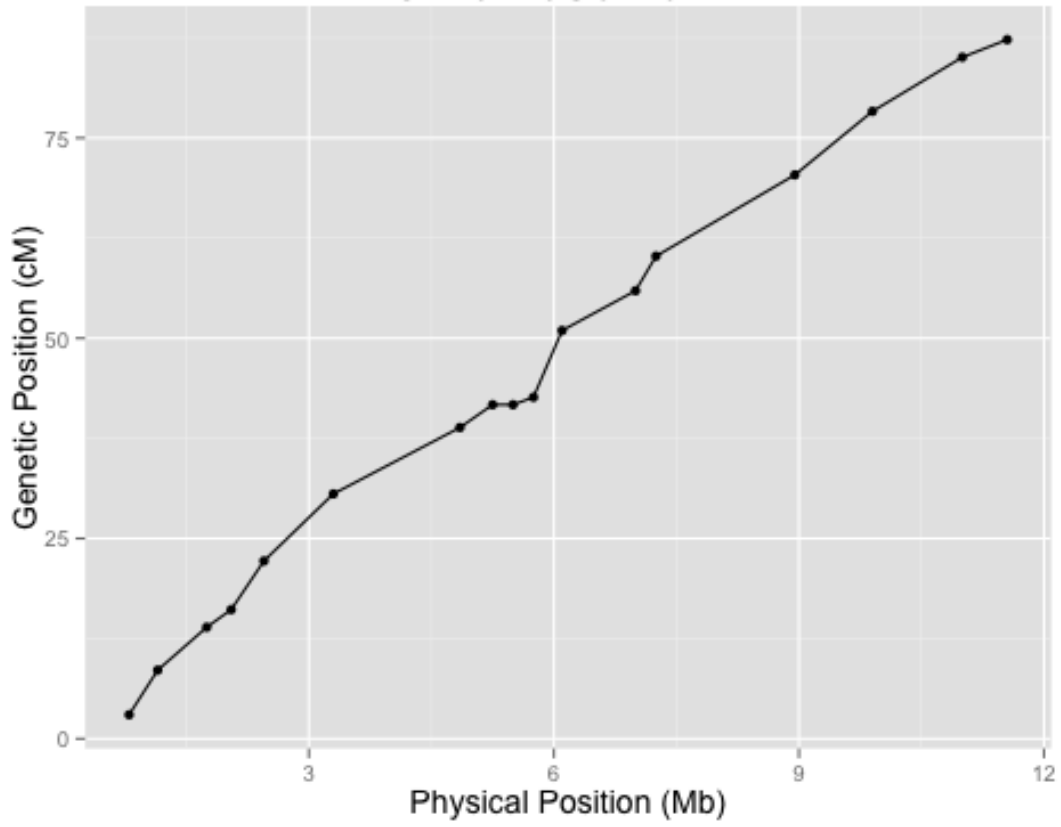
# Chromosome 14



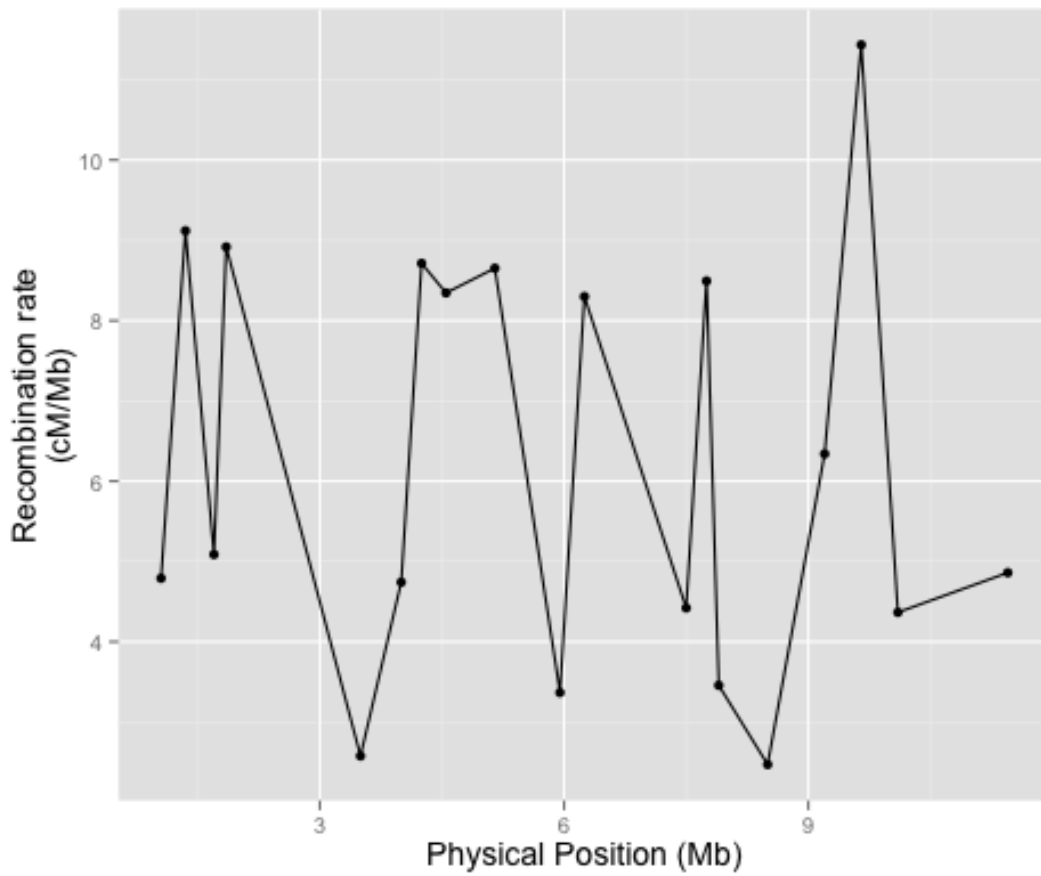
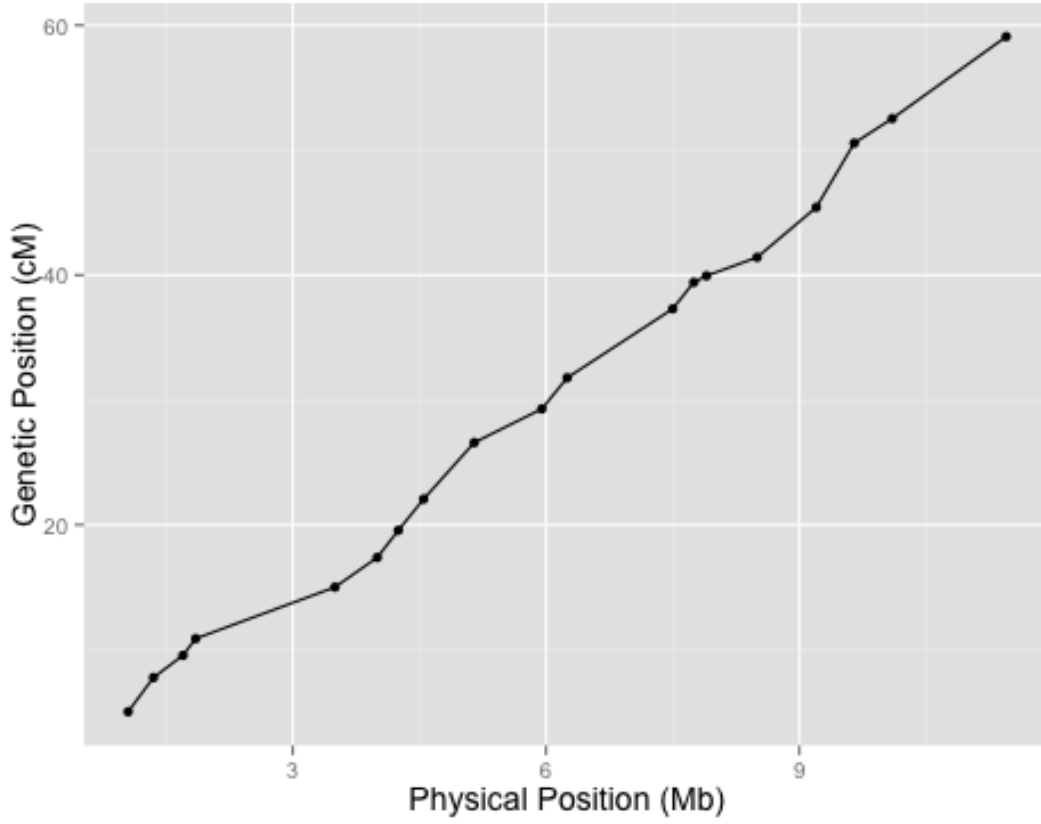
# Chromosome 13



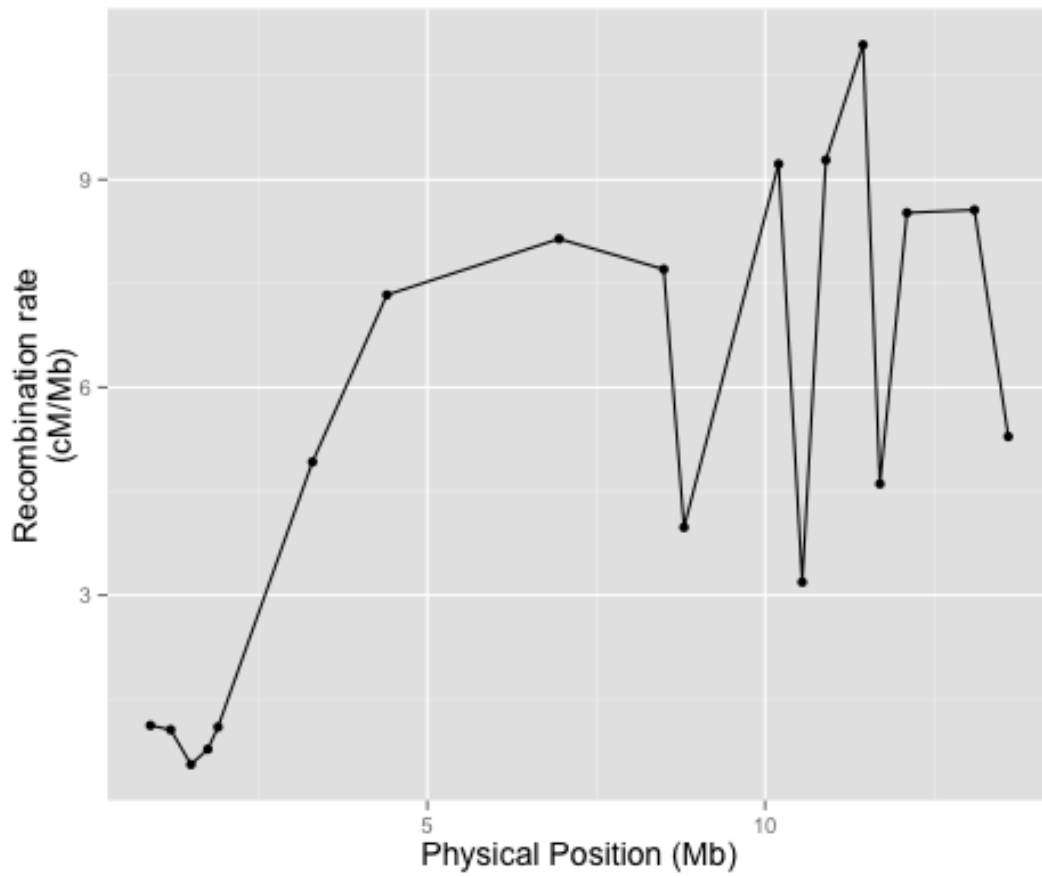
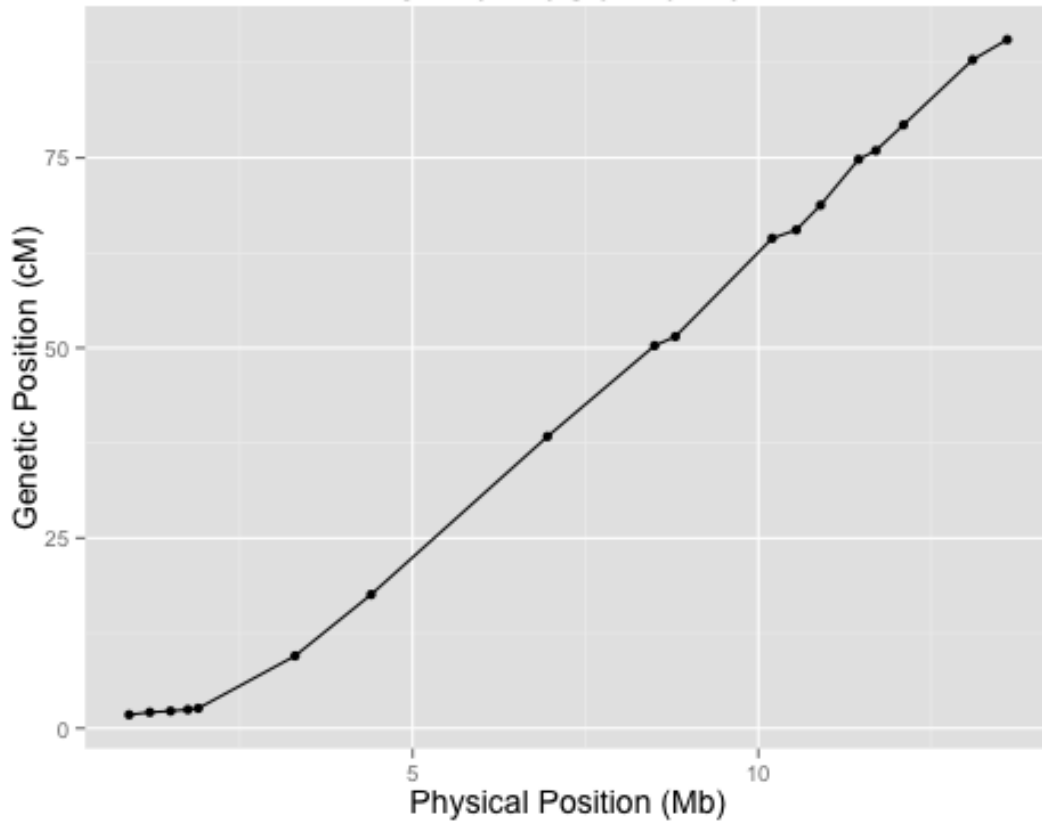
## Chromosome 12



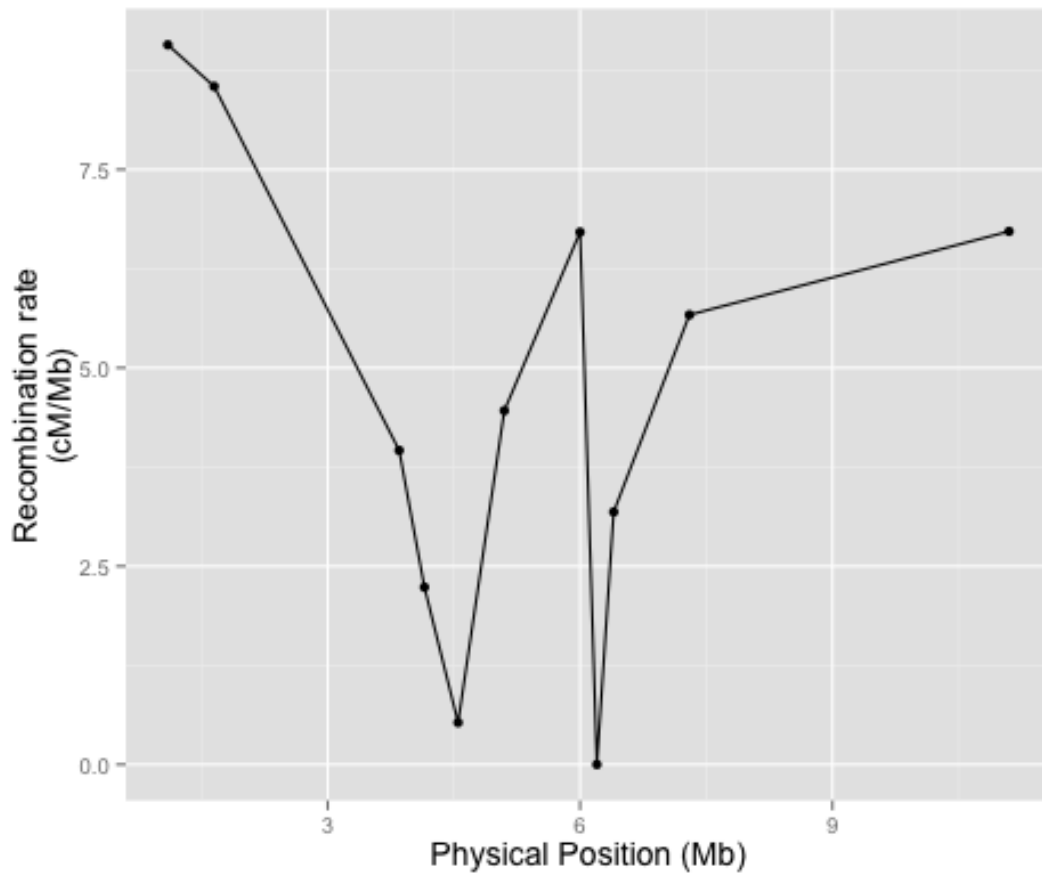
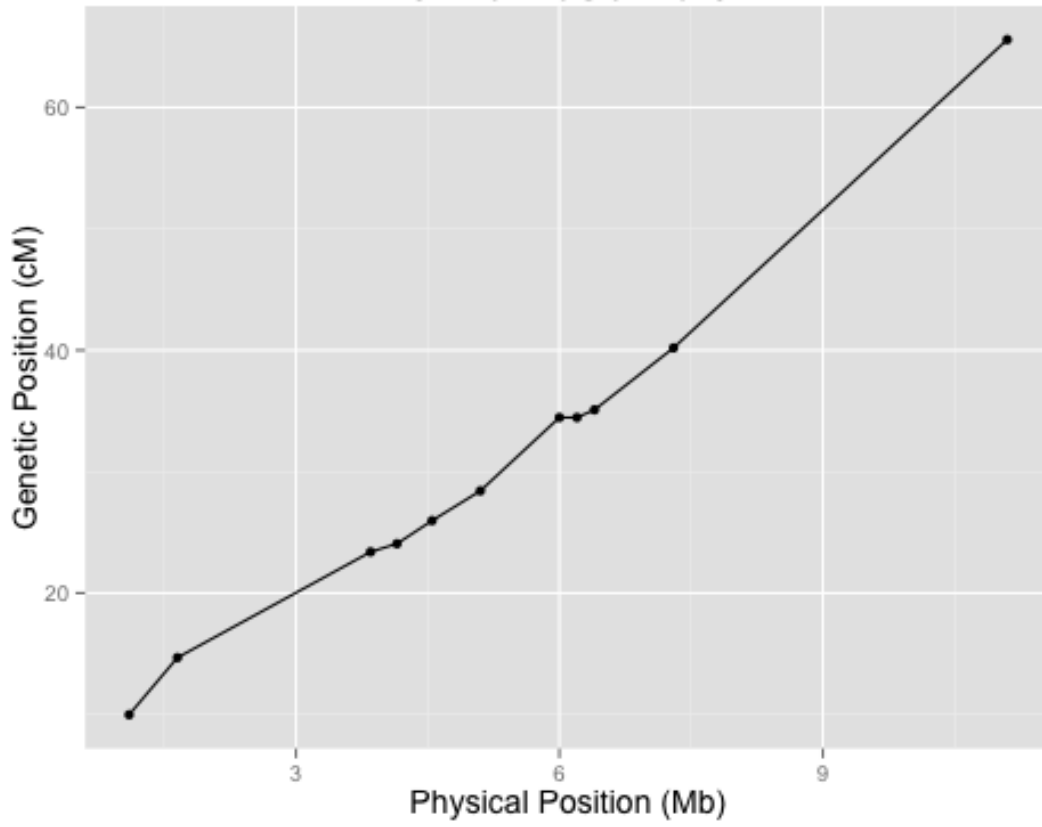
# Chromosome 11



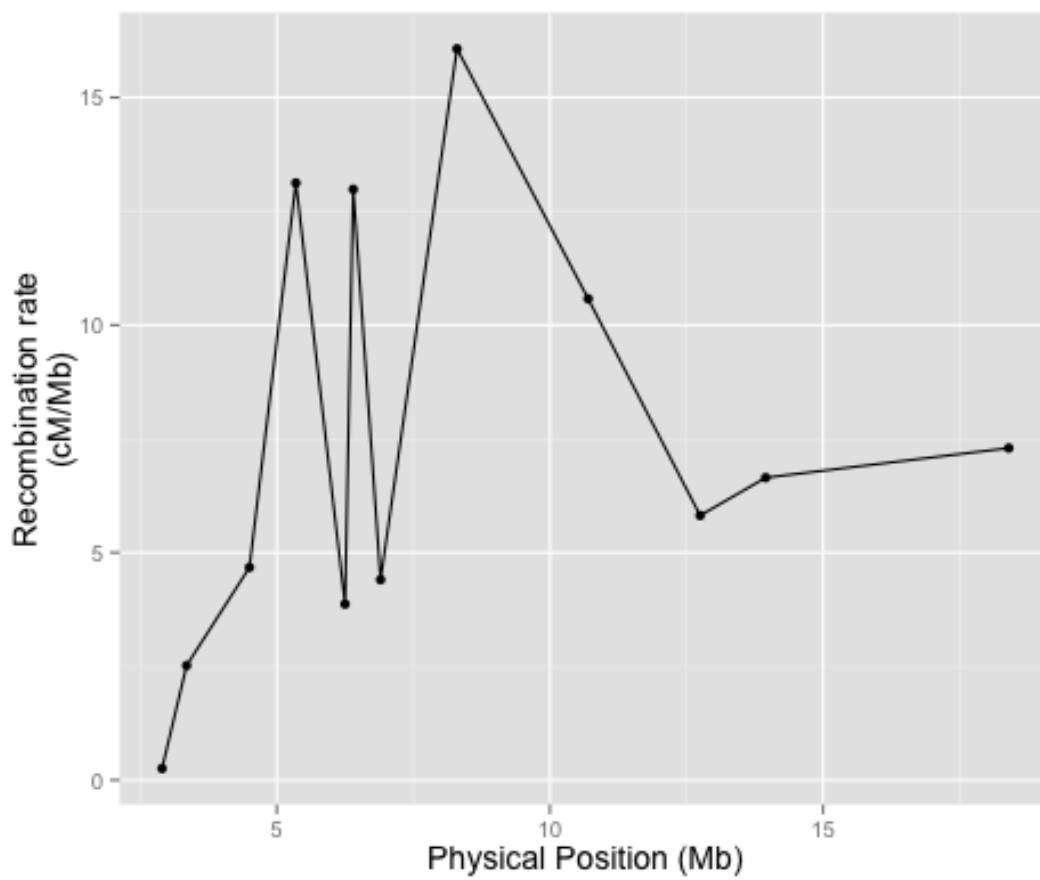
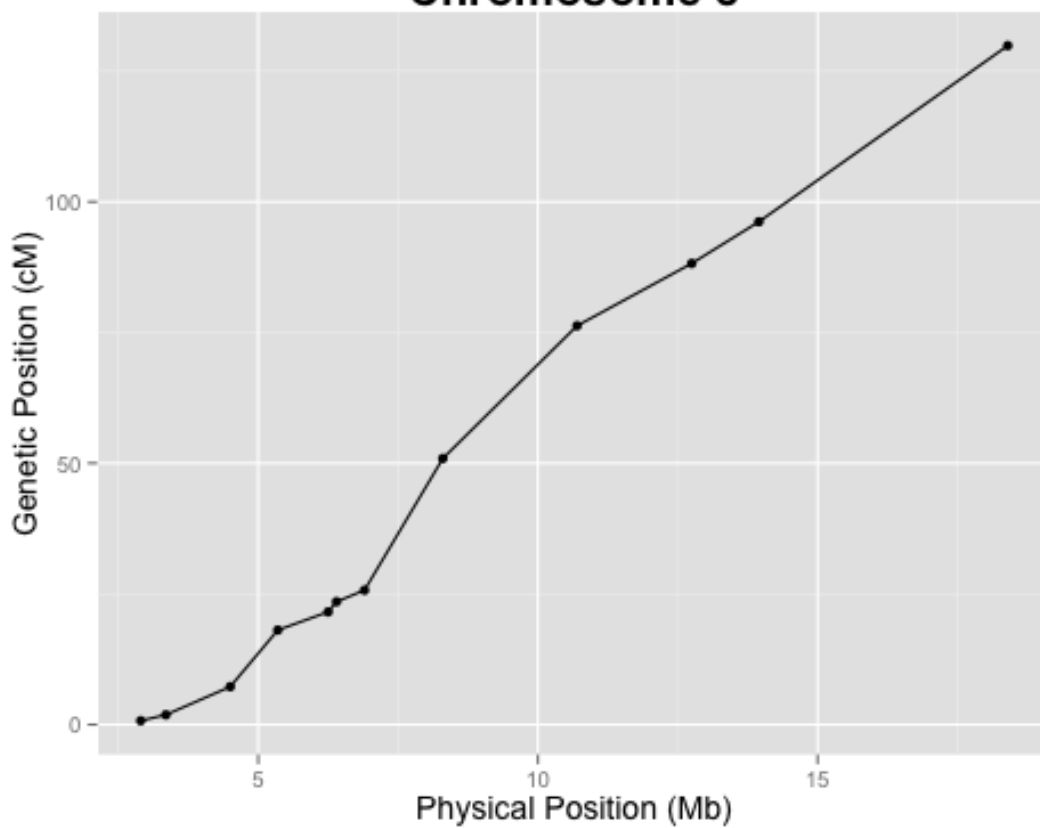
# Chromosome 10



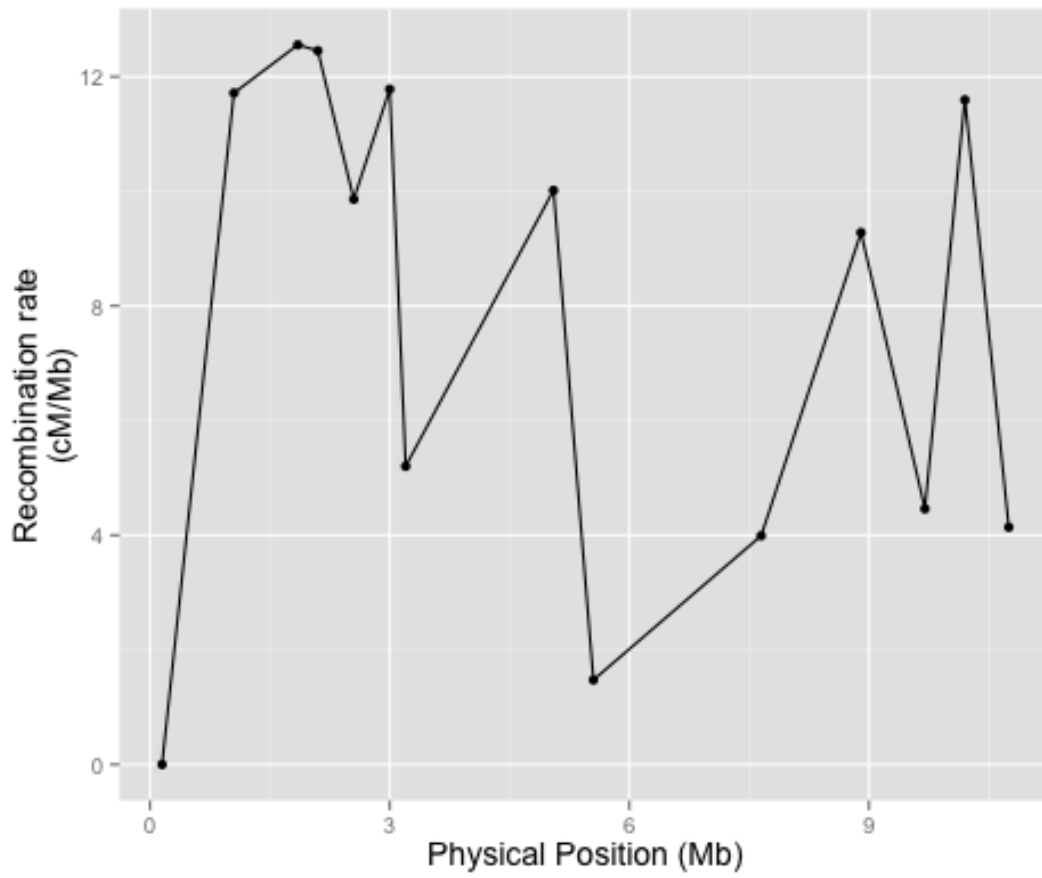
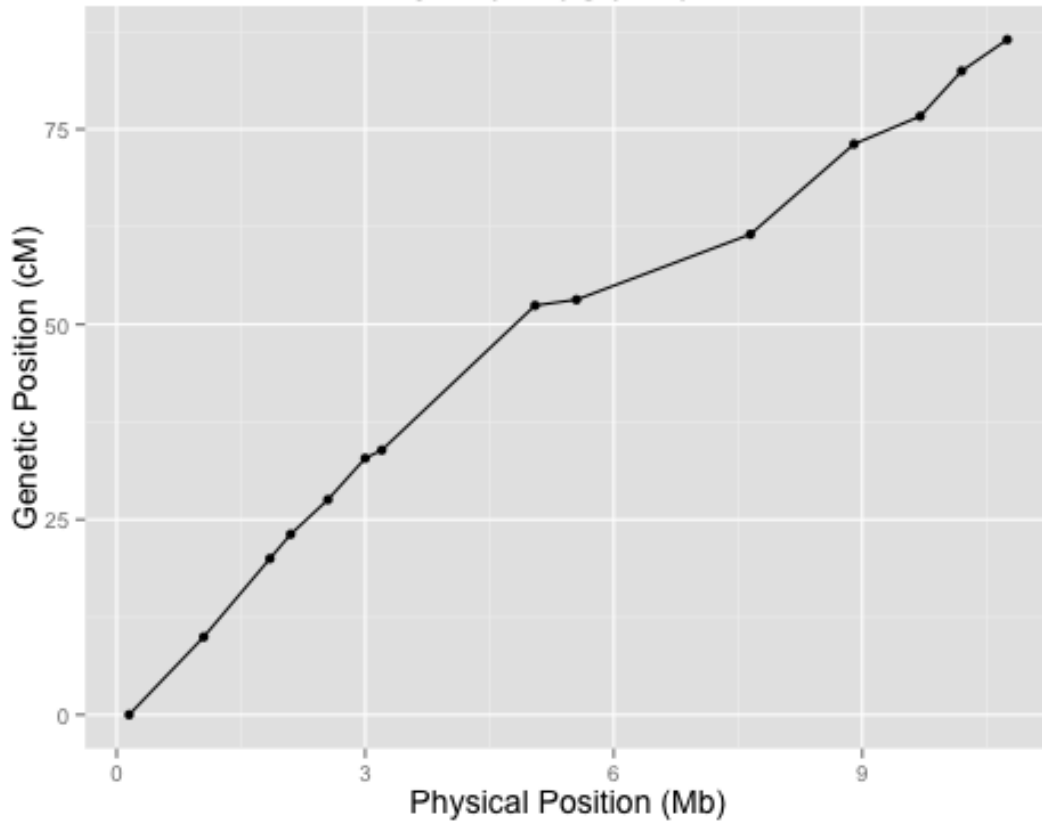
# Chromosome 9



## Chromosome 8

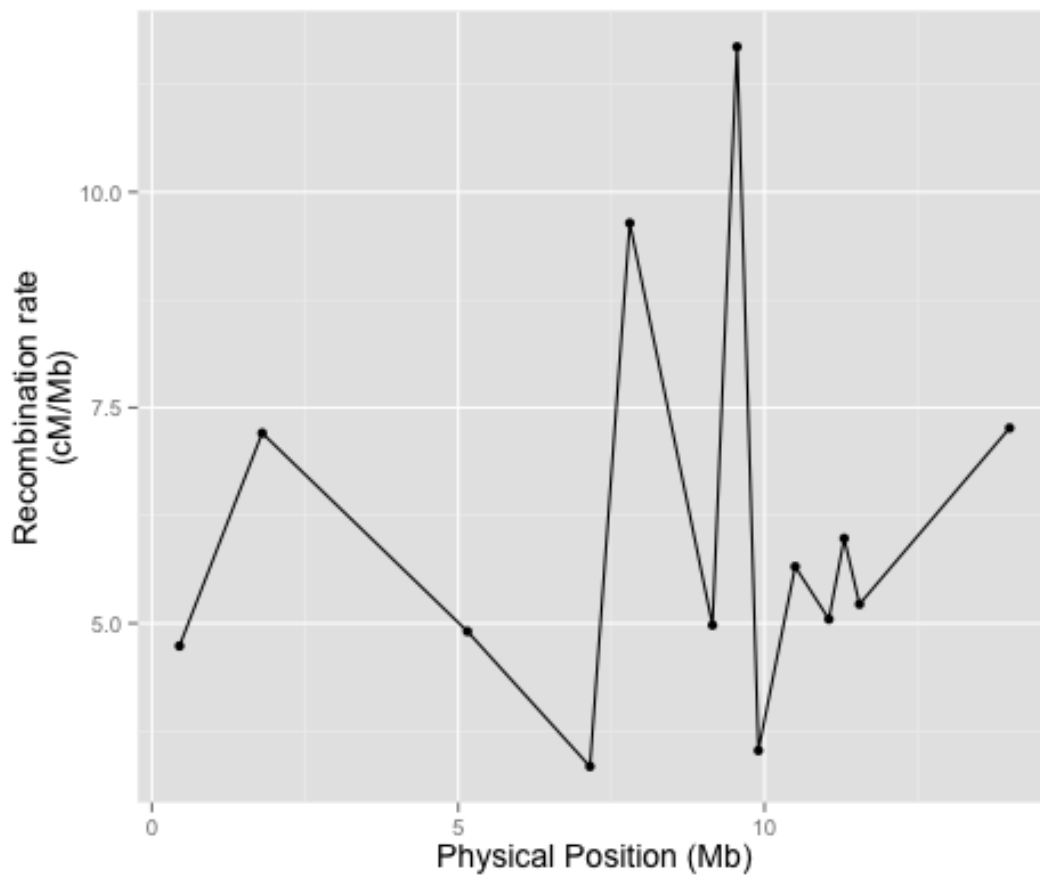
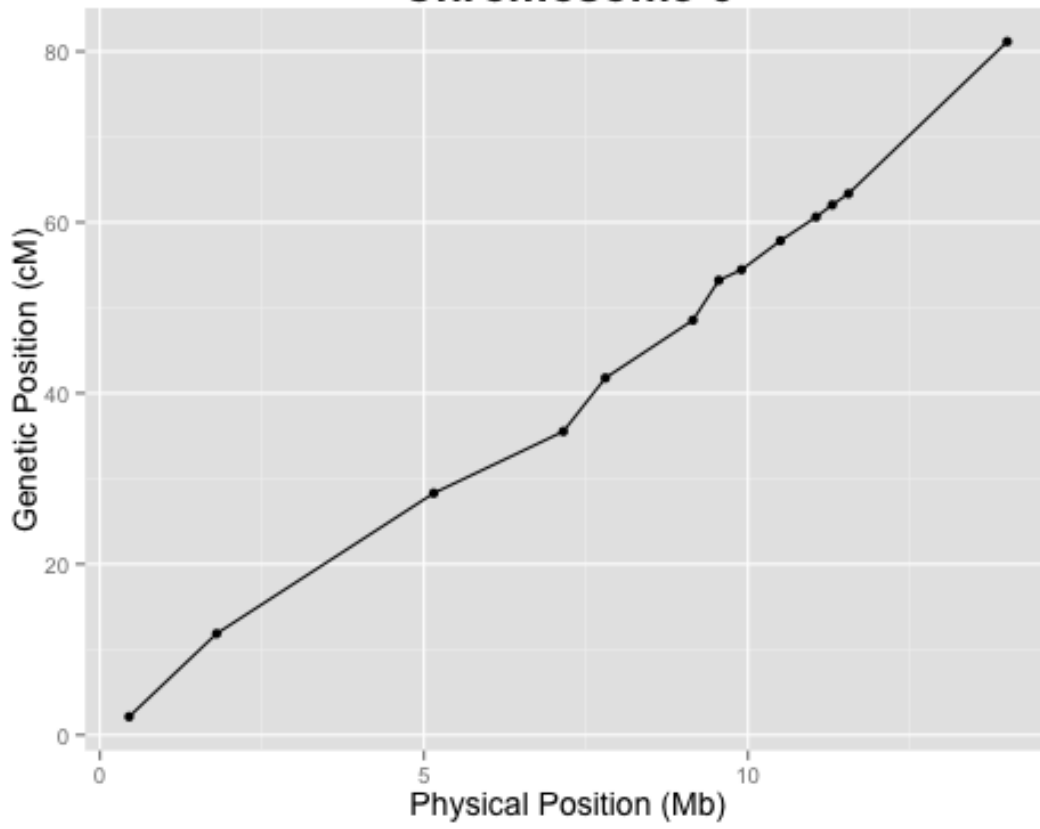


# Chromosome 7

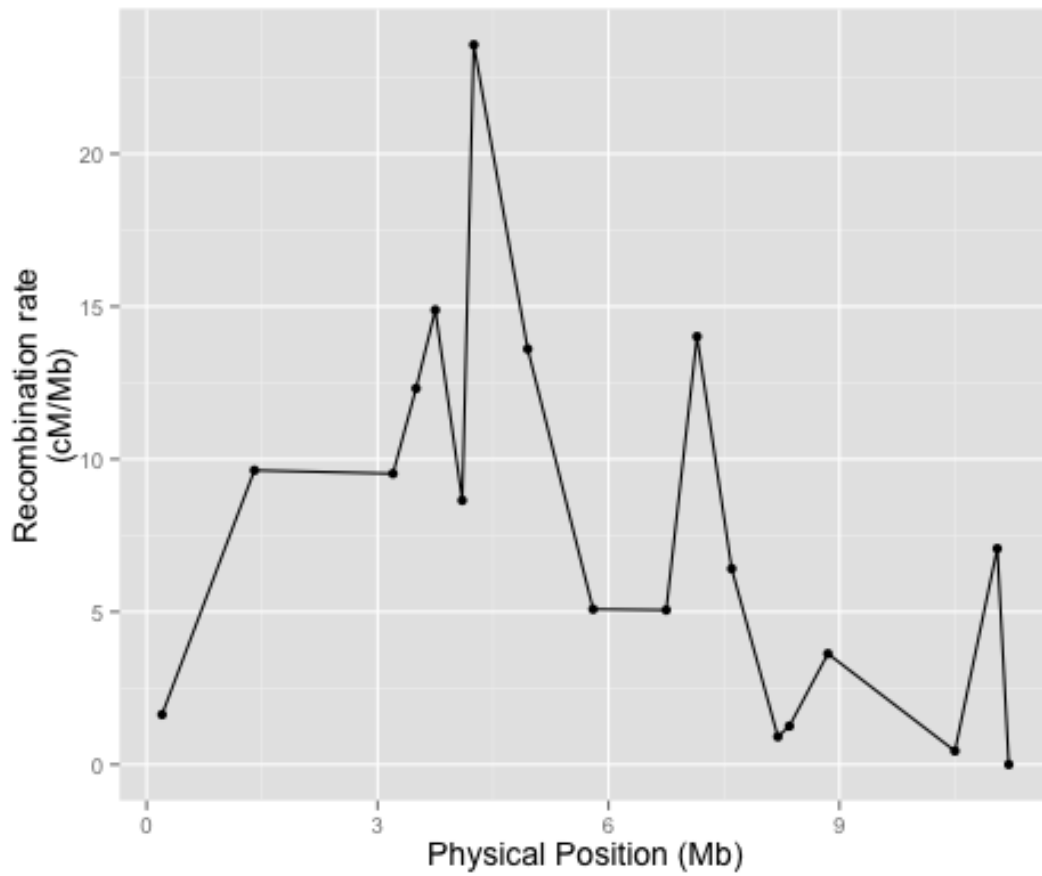
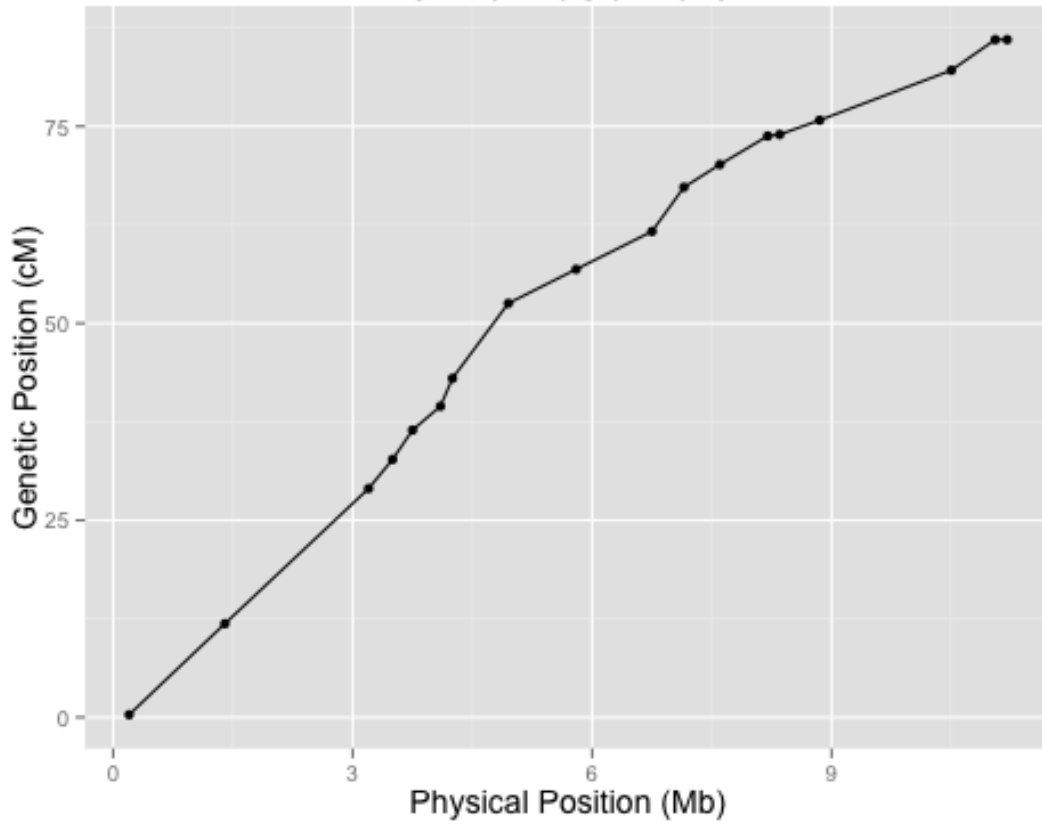




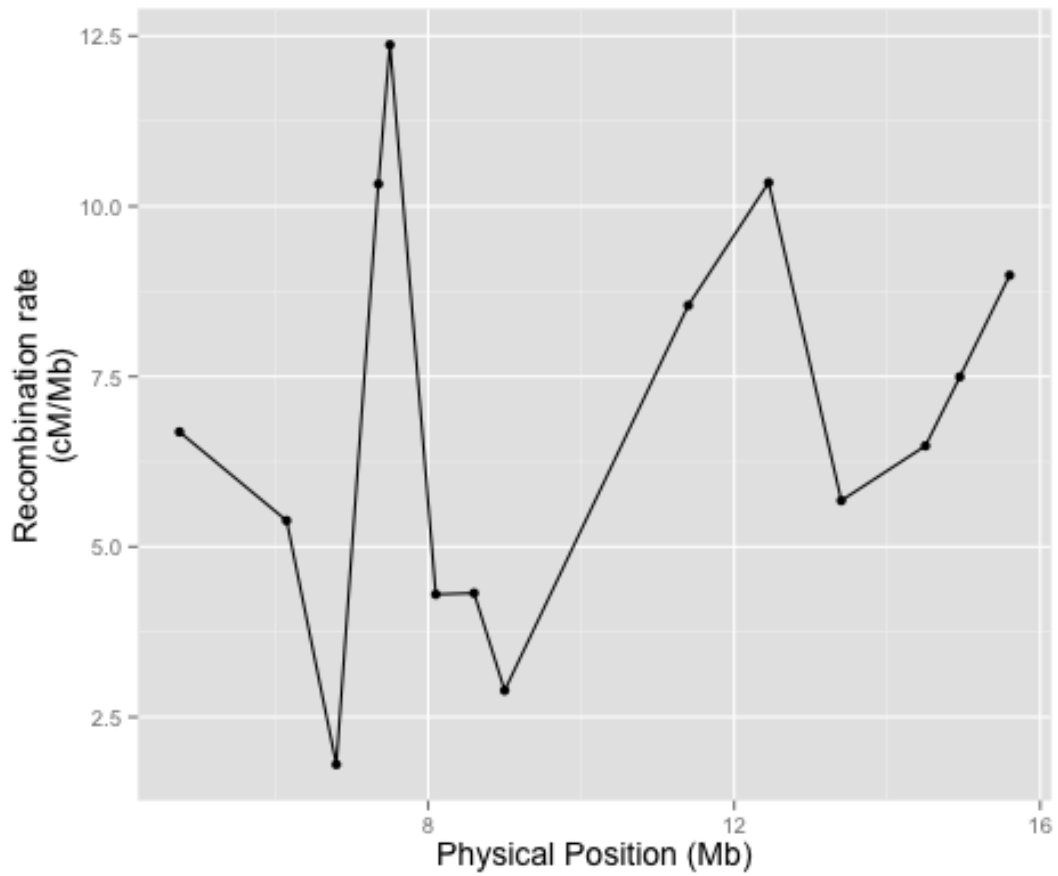
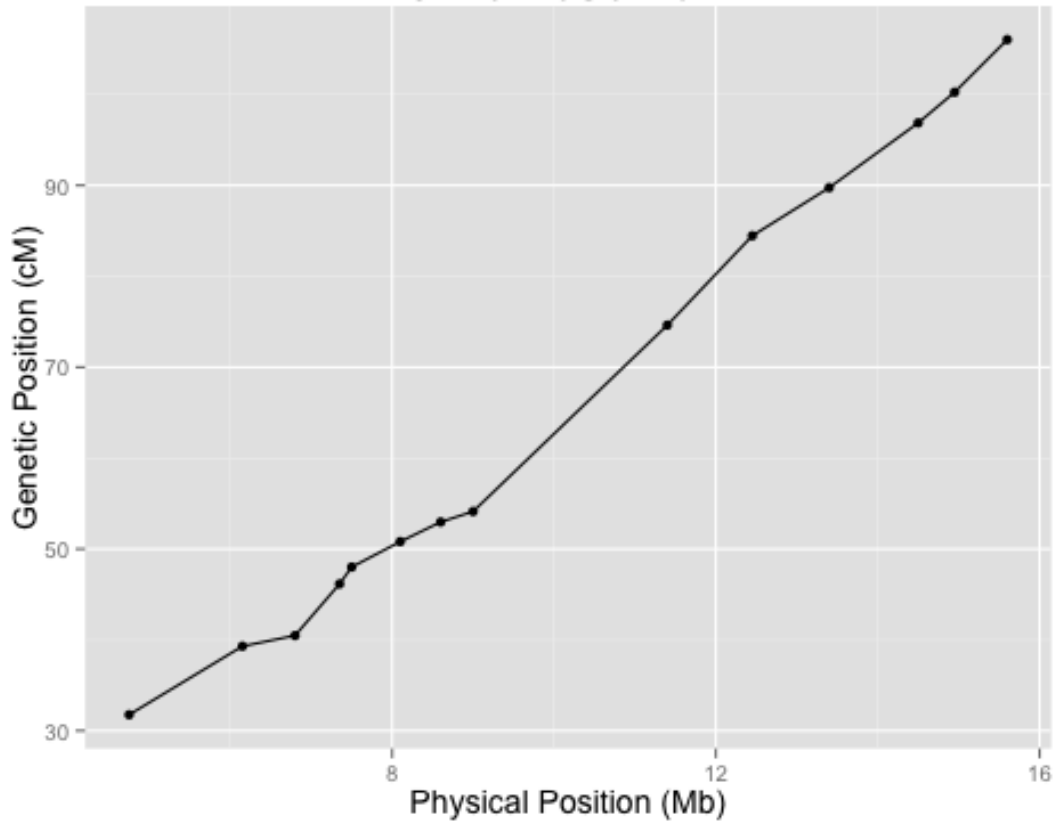
## Chromosome 6



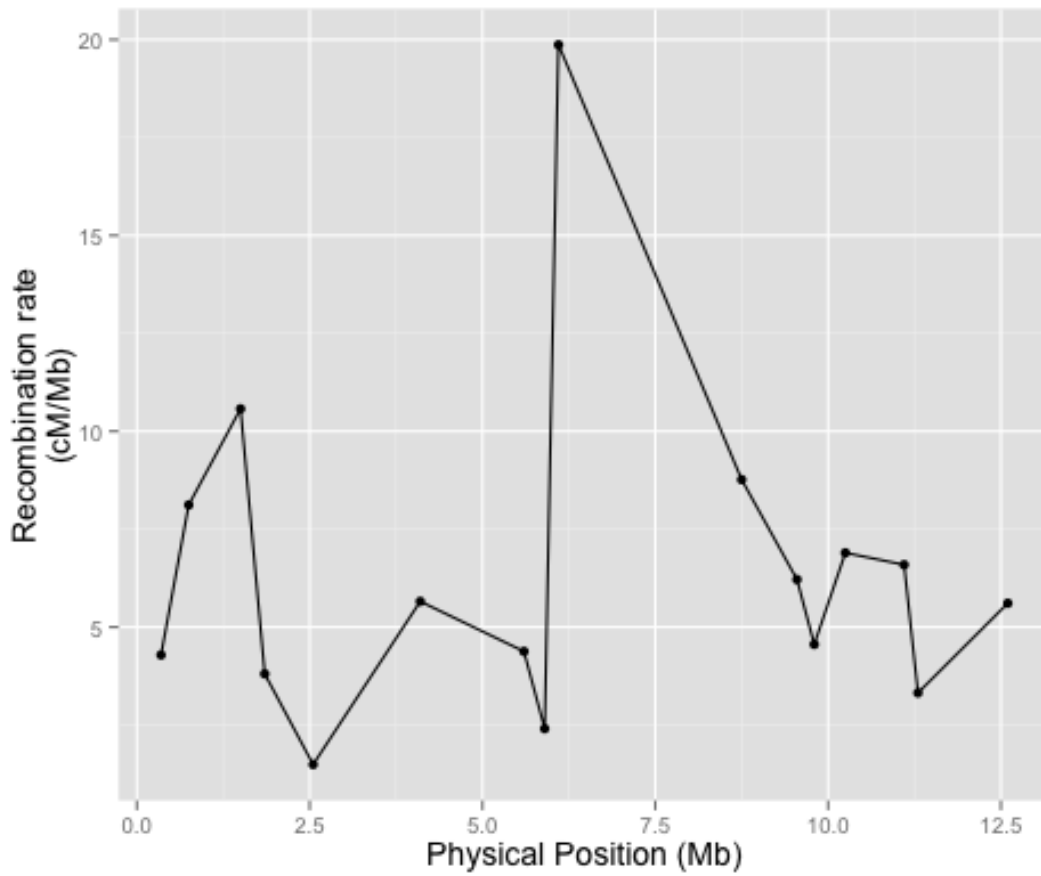
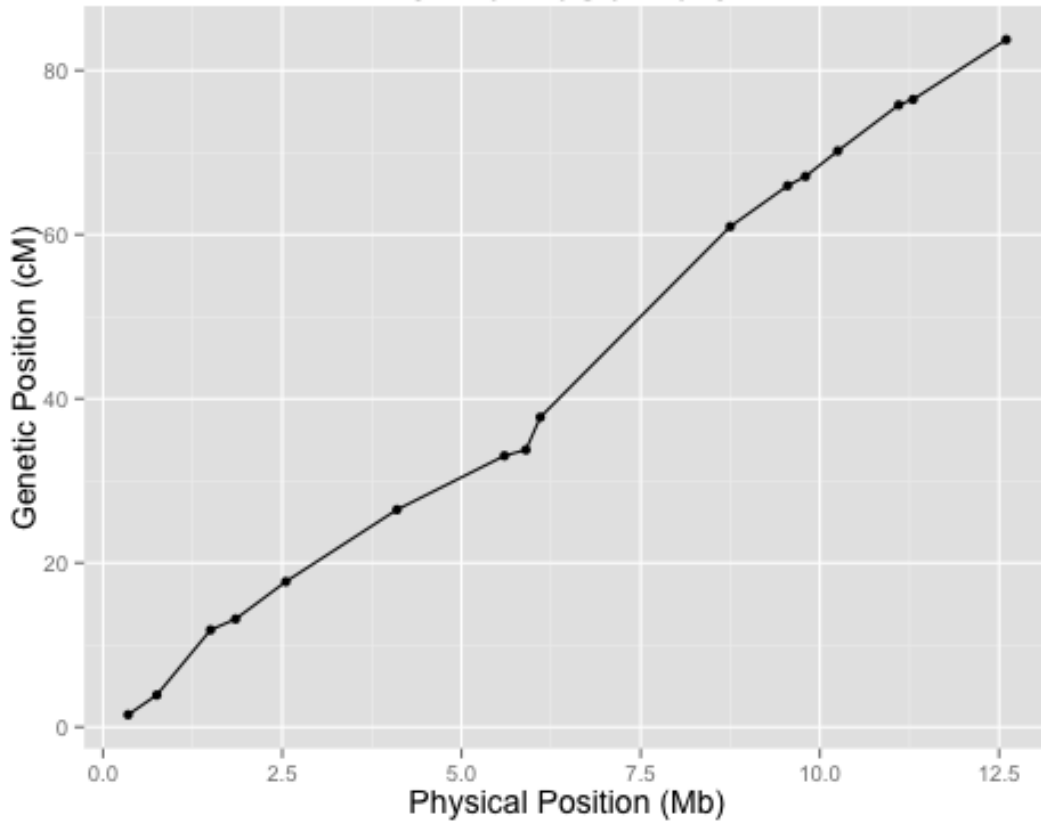
# Chromosome 5



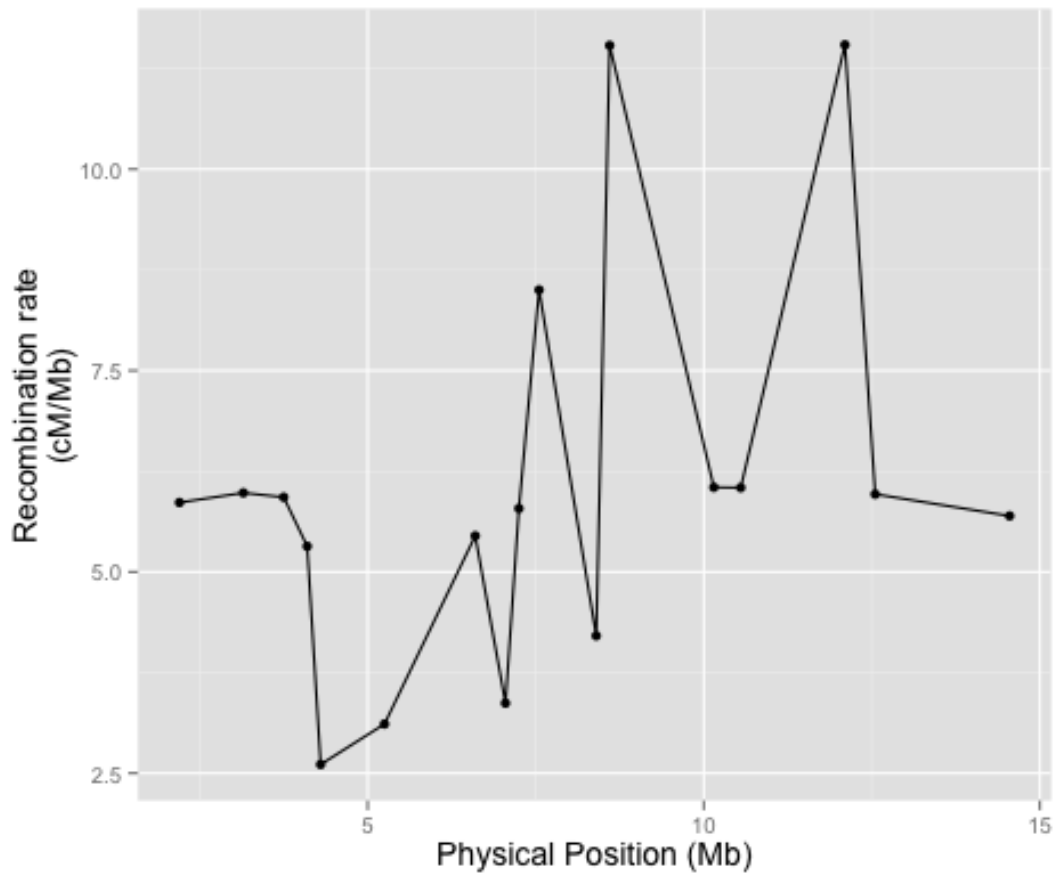
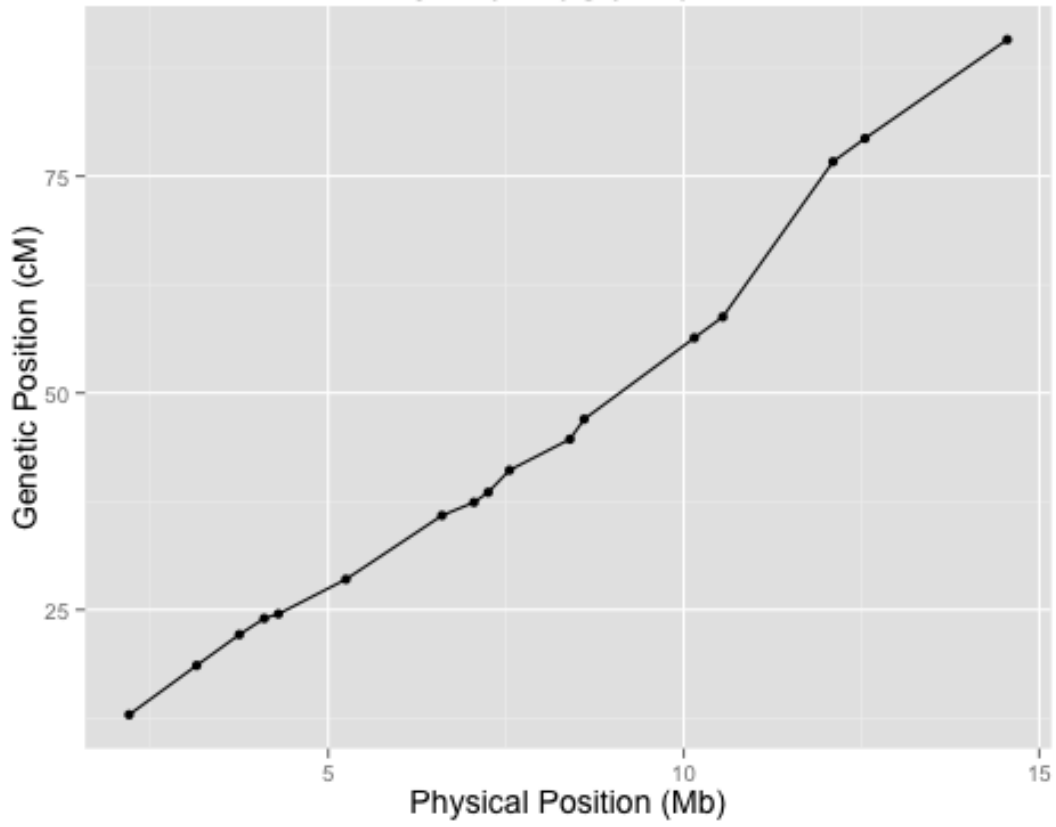
## Chromosome 4



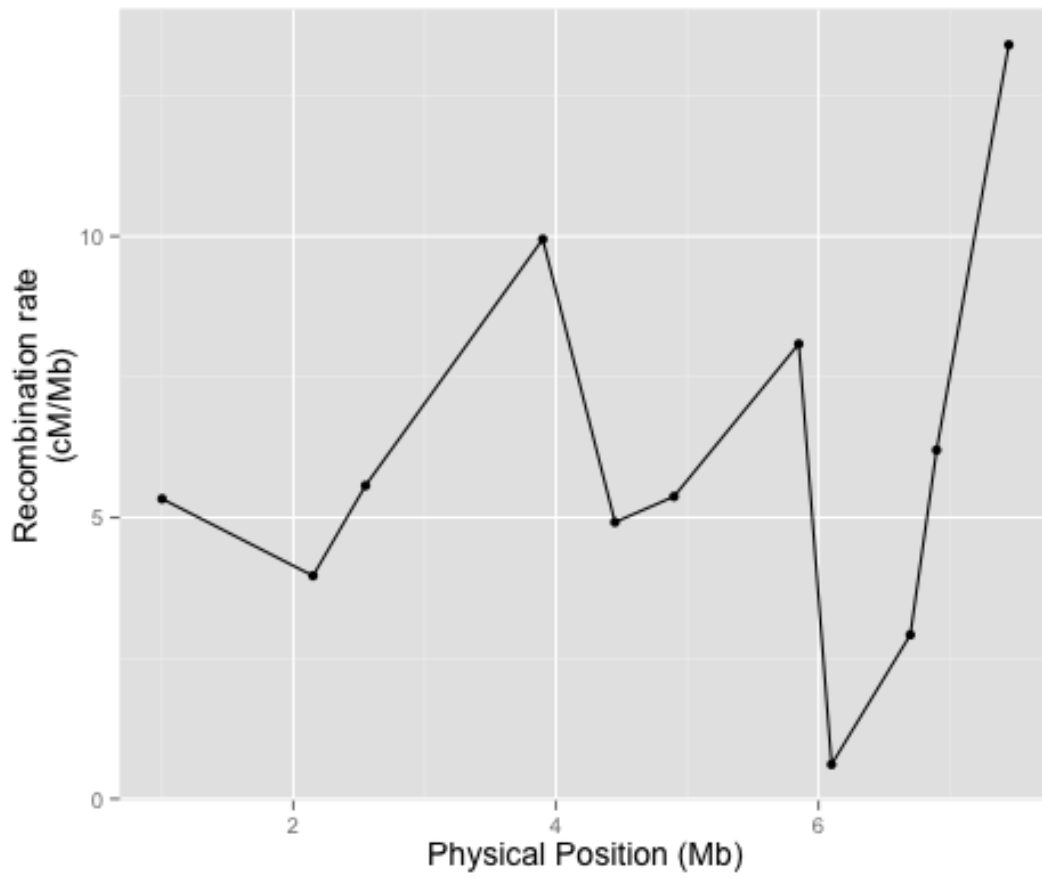
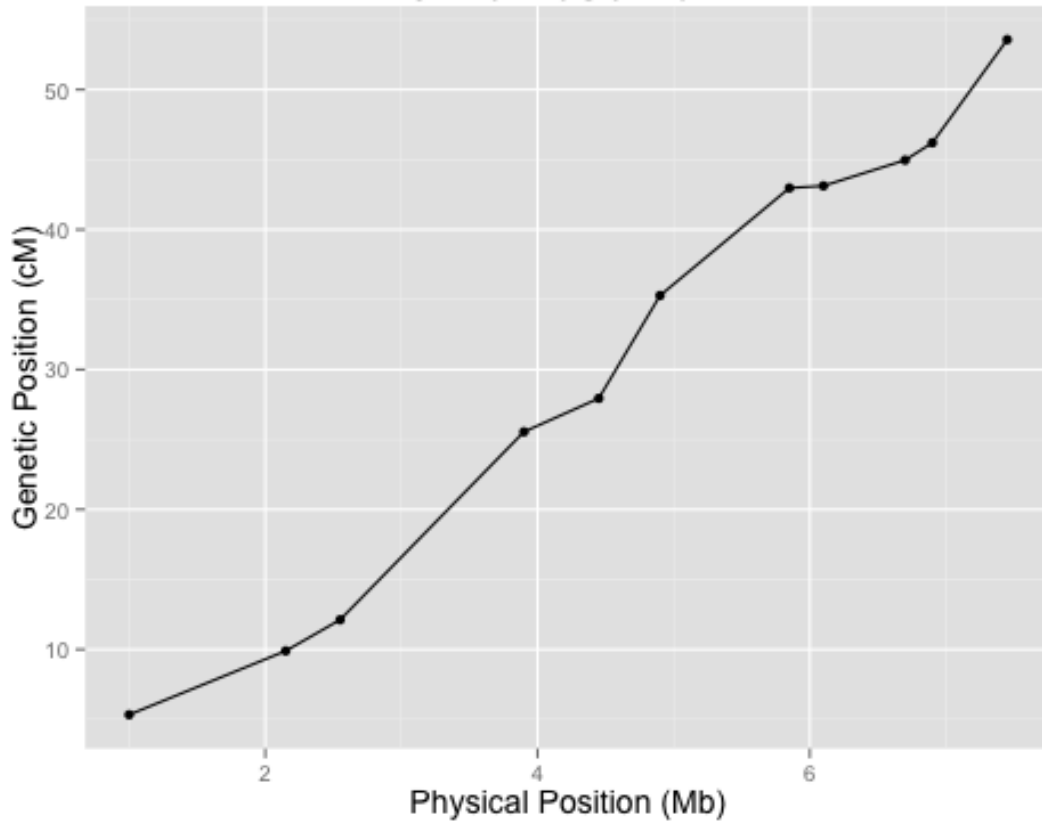
### Chromosome 3



## Chromosome 2



# Chromosome 1



**Figure S1** Depiction of recombination rate variation along each chromosome. Each point is an estimate from an entire scaffold. Scaffolds were aligned according to their order in the linkage map. Note that the apparently adjacent scaffolds are not physically adjacent as we do not account for distance between scaffolds. The distance between points corresponds to the length of the scaffolds. Only scaffolds >100 kb are depicted, and extreme observations were removed for better visualization.