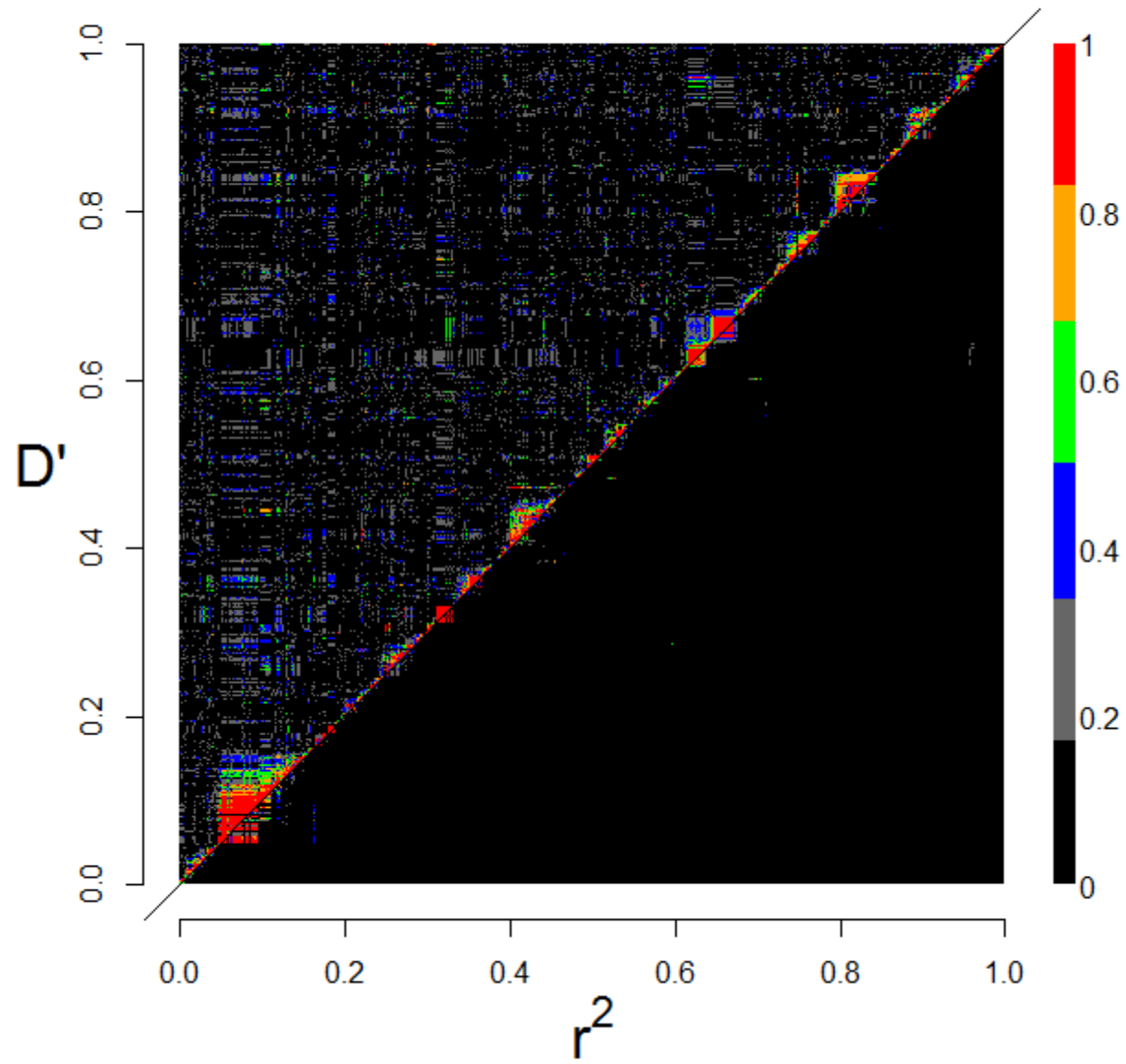
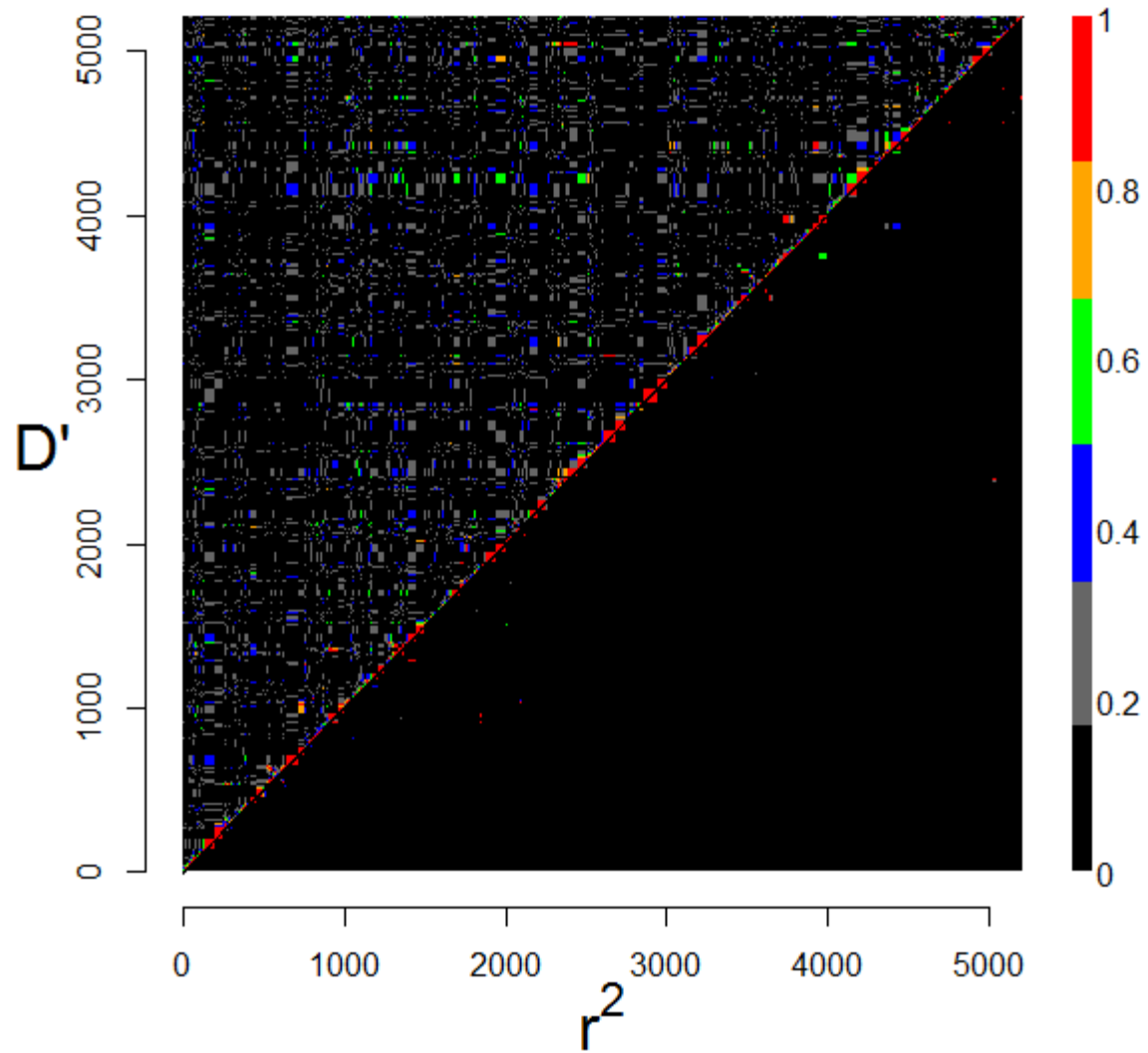


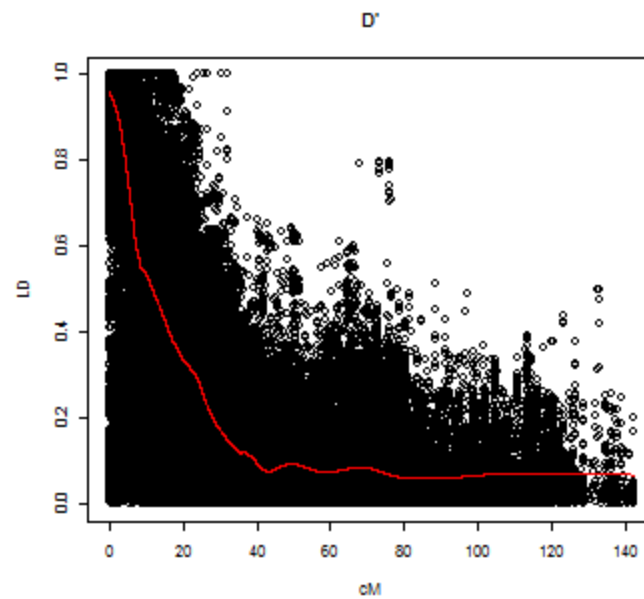
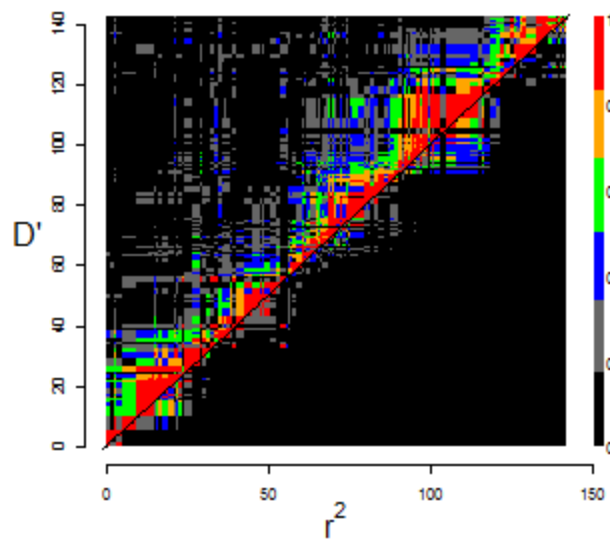
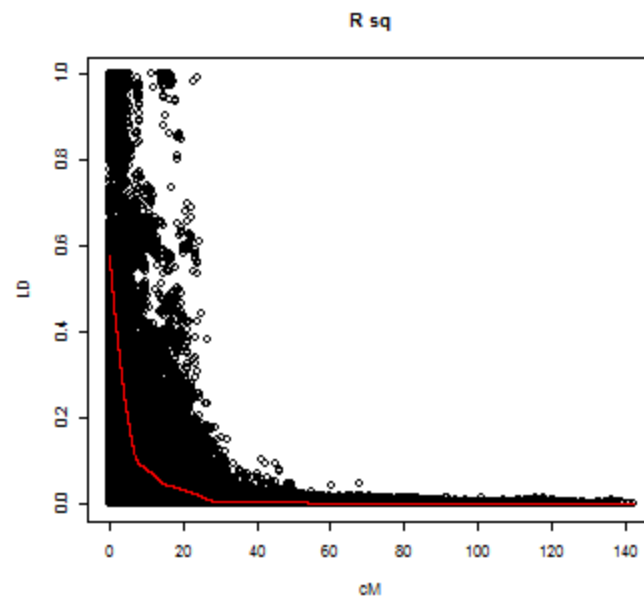
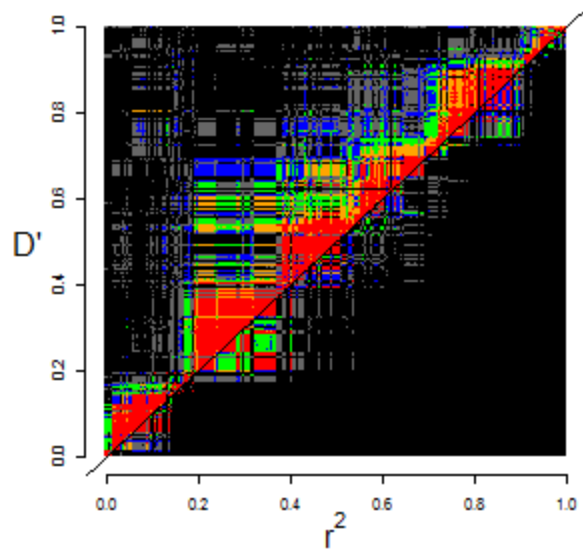
Figure S1 MAGIC progeny LD heatmaps and LD decay plots for the 21 wheat chromosomes based on 13074 mapped and segregating SNPs. For each chromosome: Top left: heatmap of LD among ordered markers. D' above diagonal, r^2 below. Bottom left: heatmap scaled by genetic map. Top right LD decay (r^2) plotted against genetic distance between marker pairs. Bottom right: LD decay (D') plotted against genetic distance between marker pairs.

All, no map

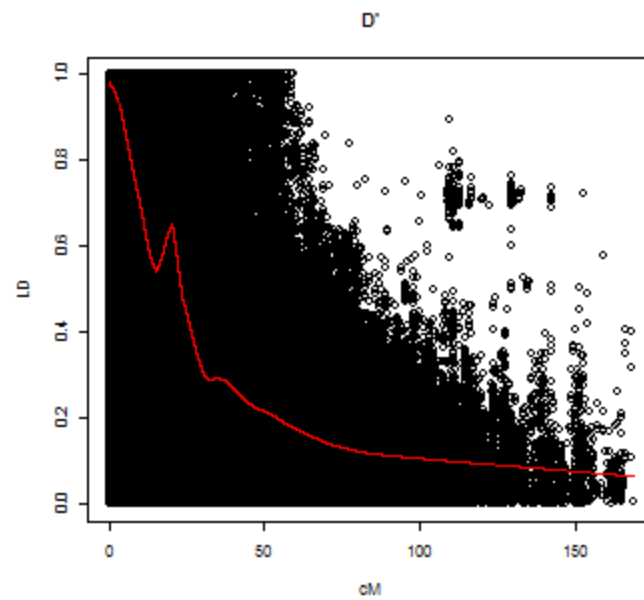
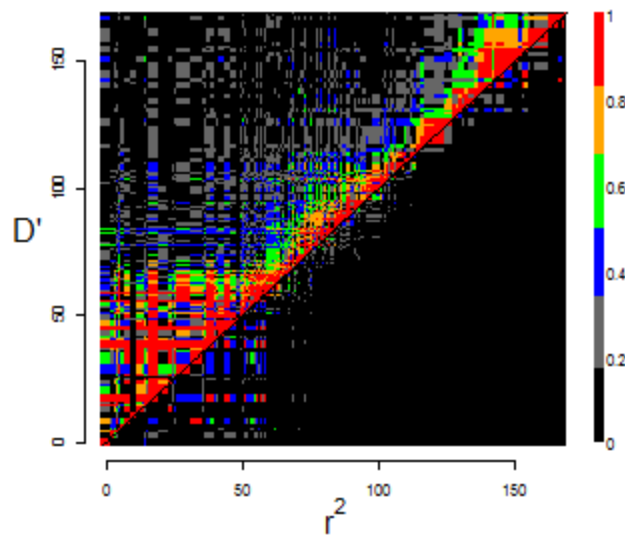
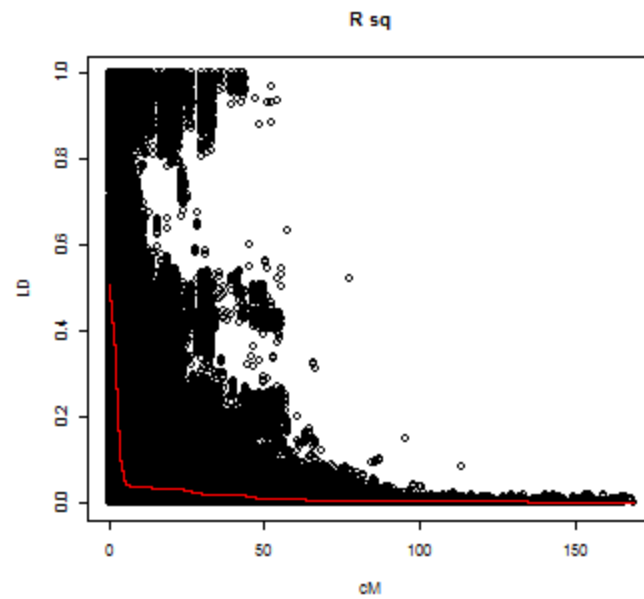
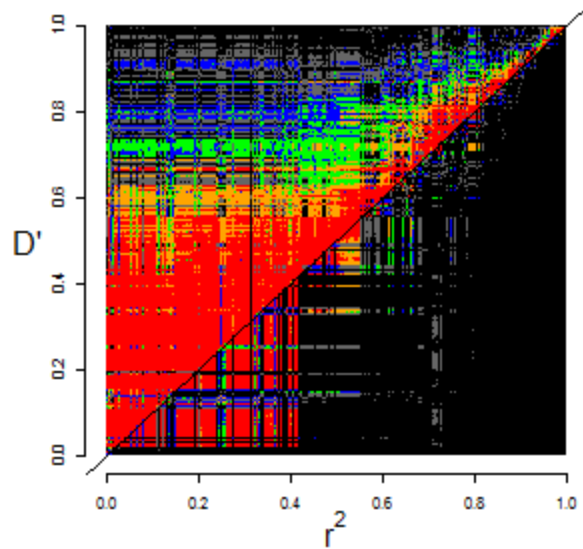


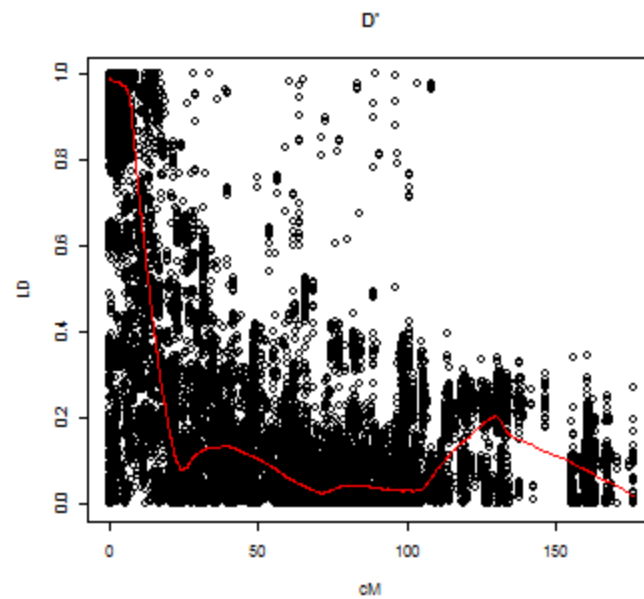
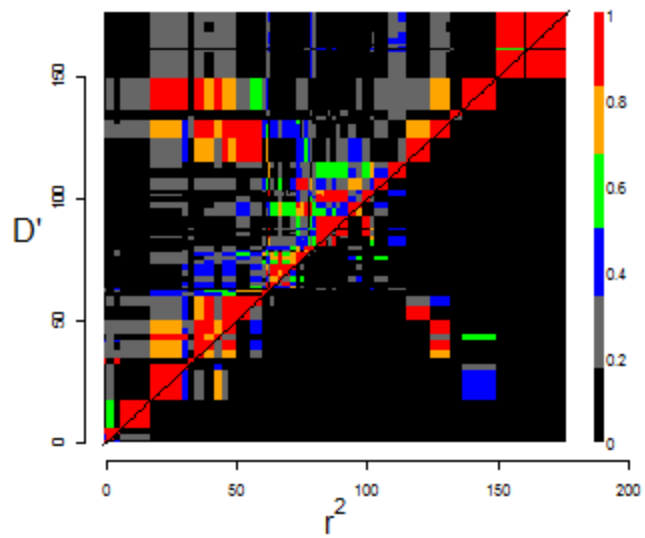
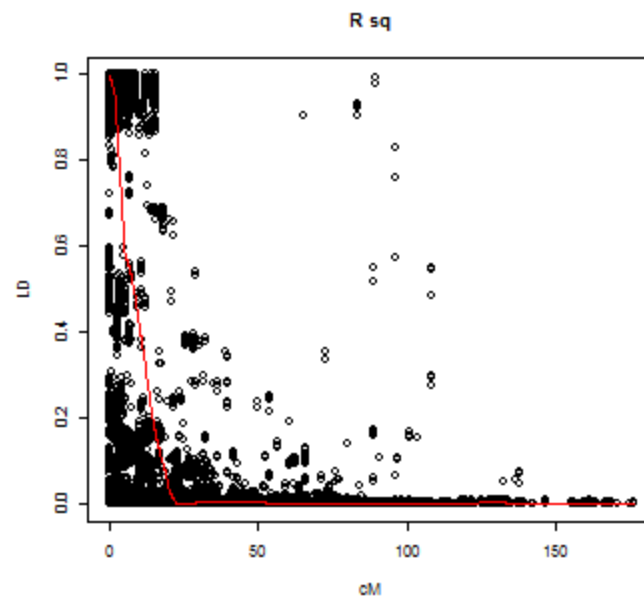
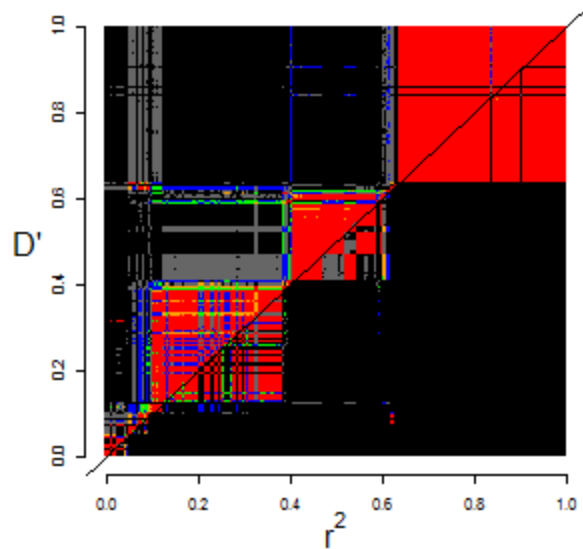
All, with map, change chromosome every 250 cM

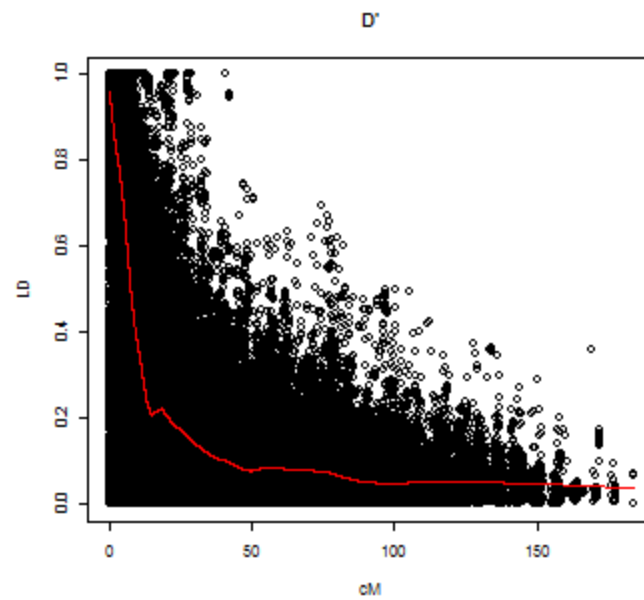
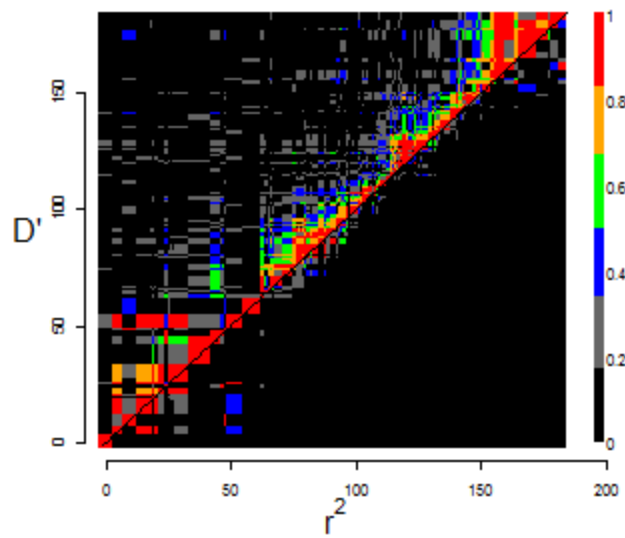
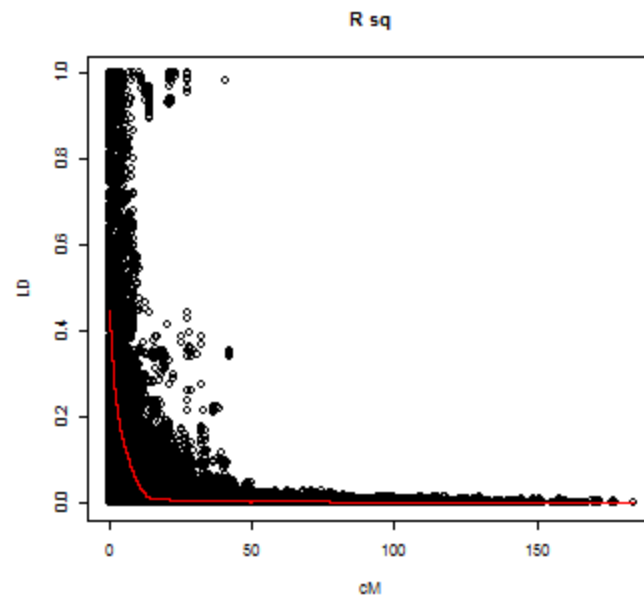
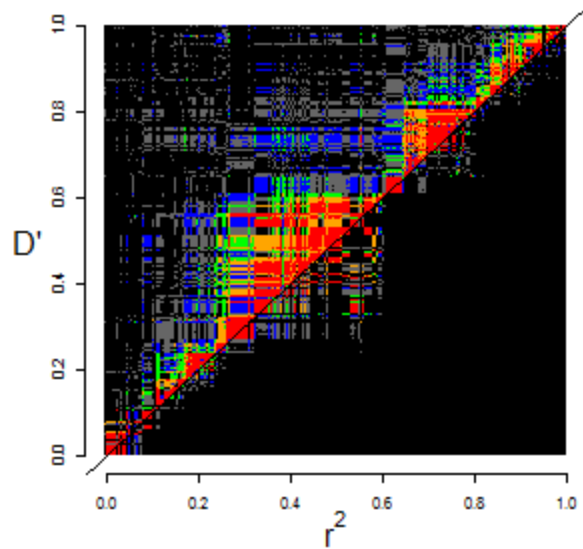


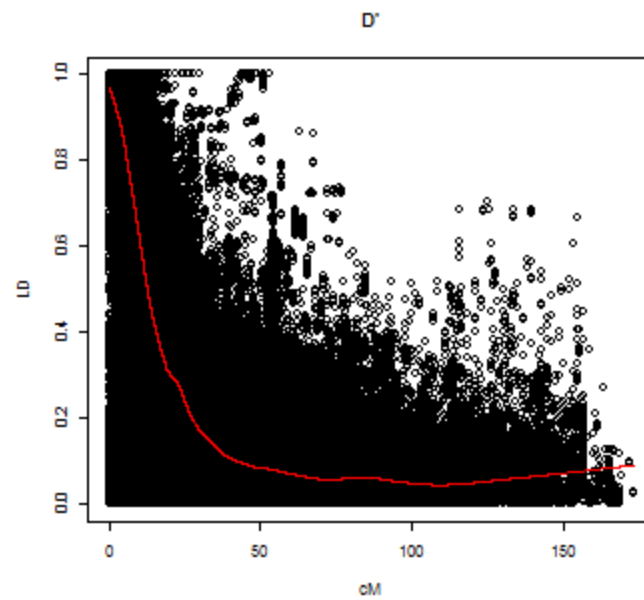
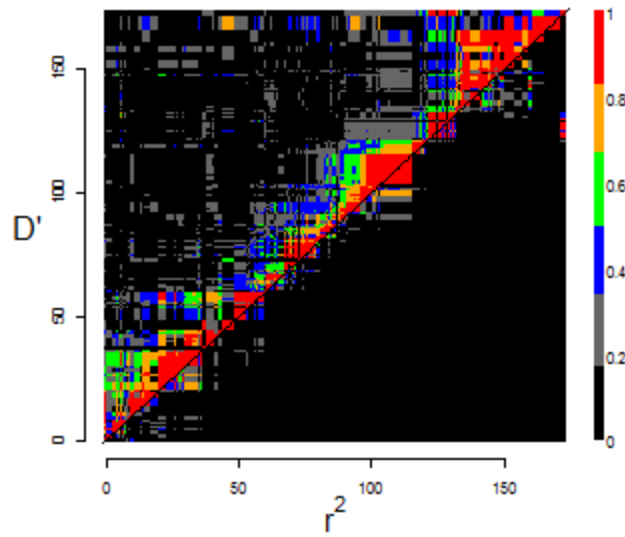
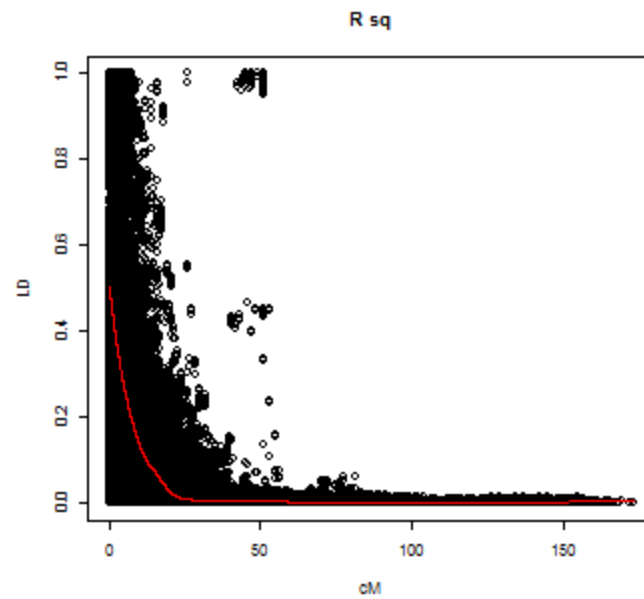
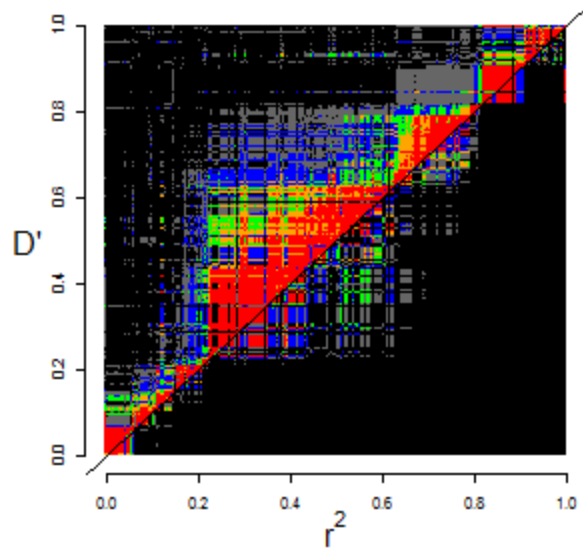


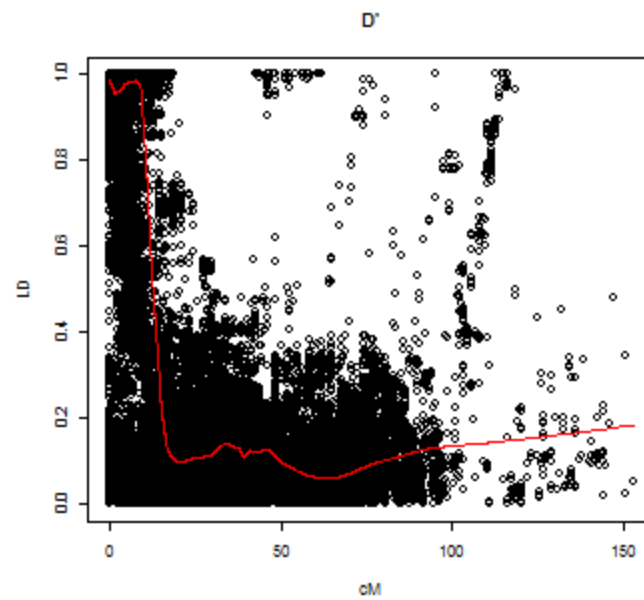
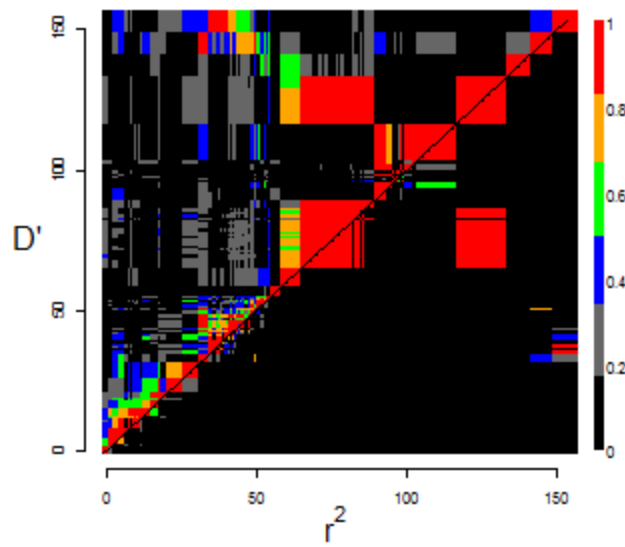
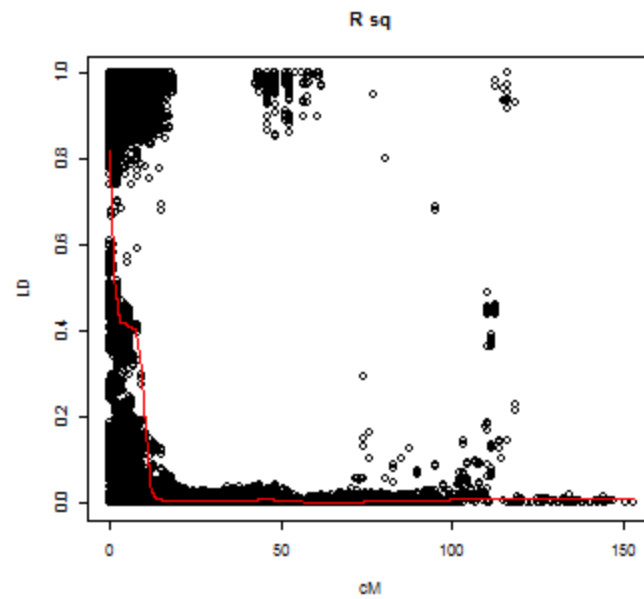
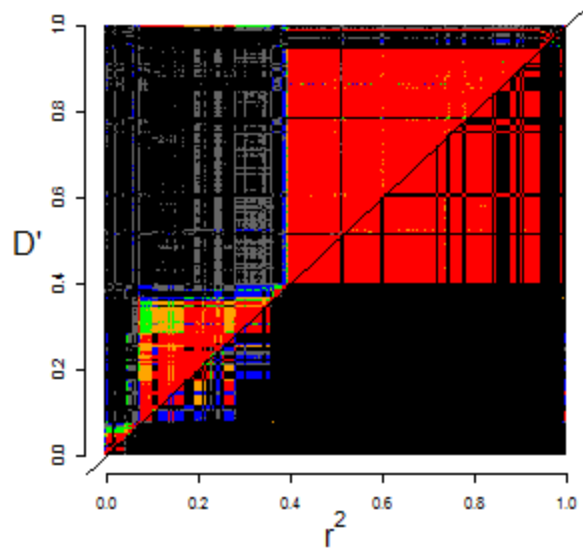
1A



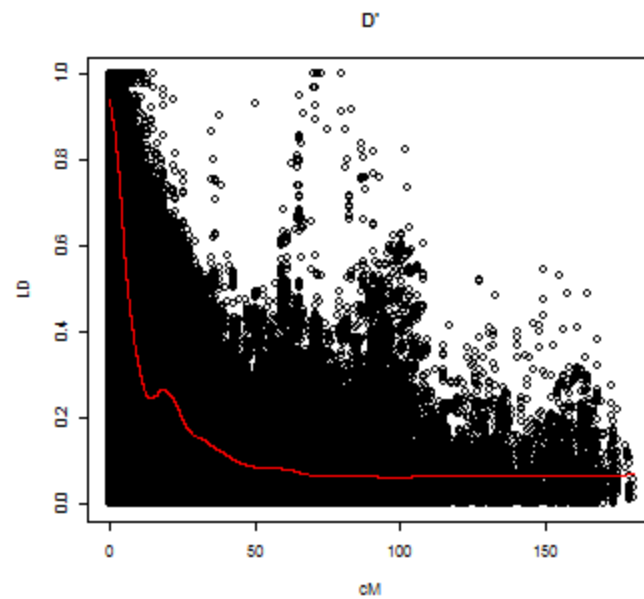
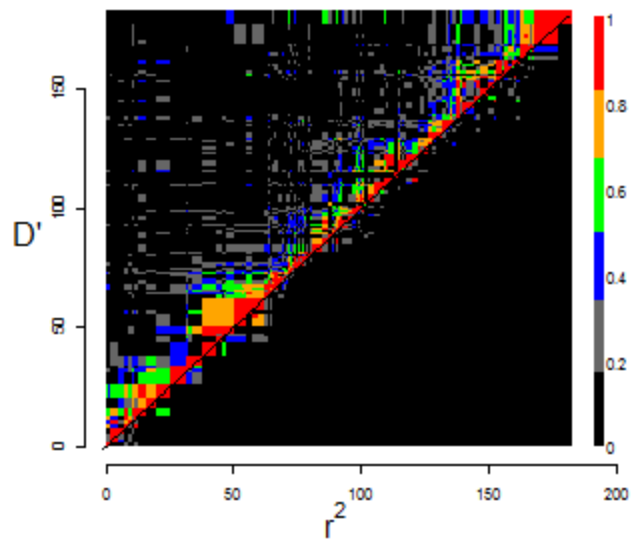
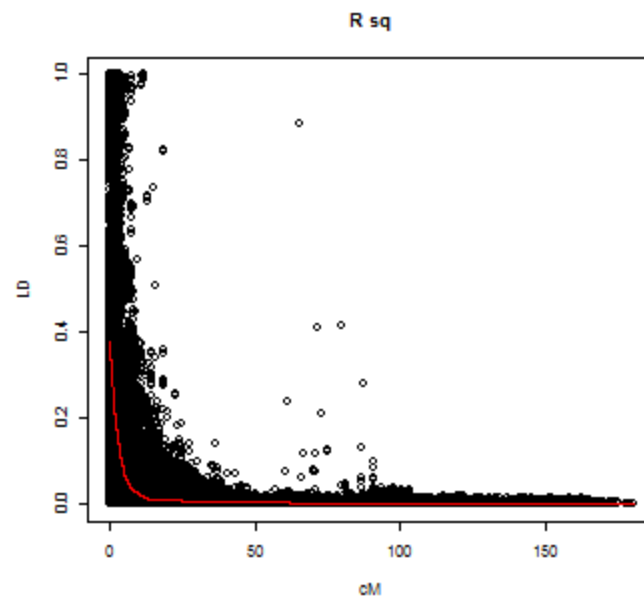
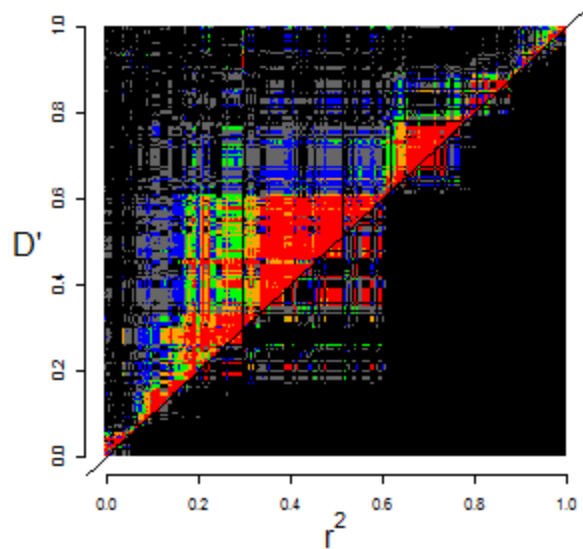




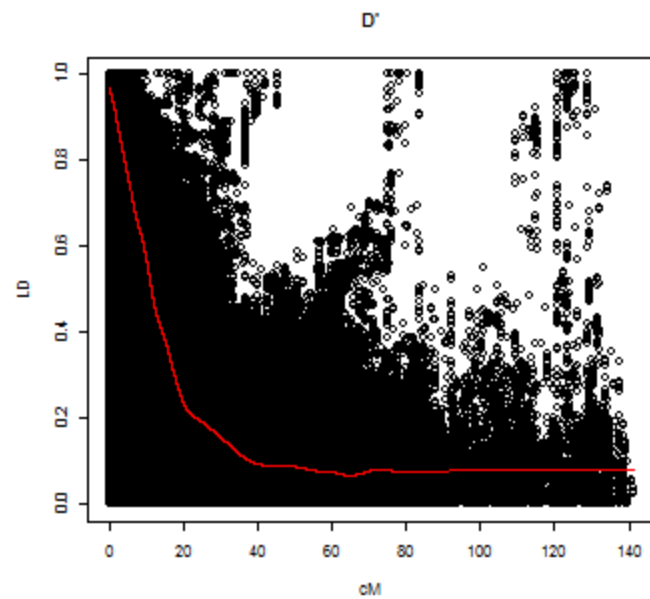
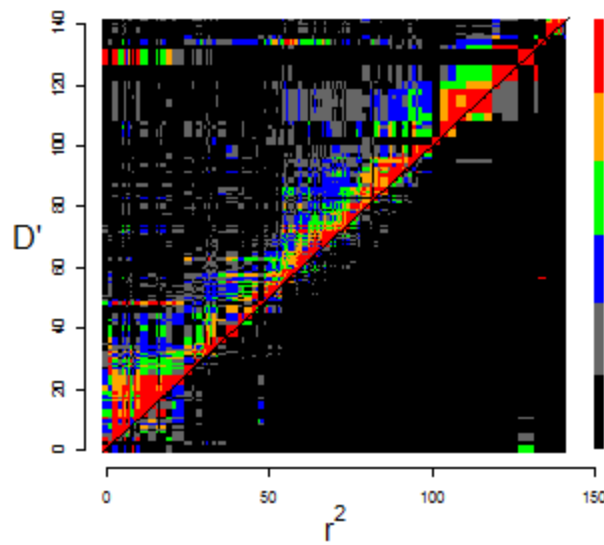
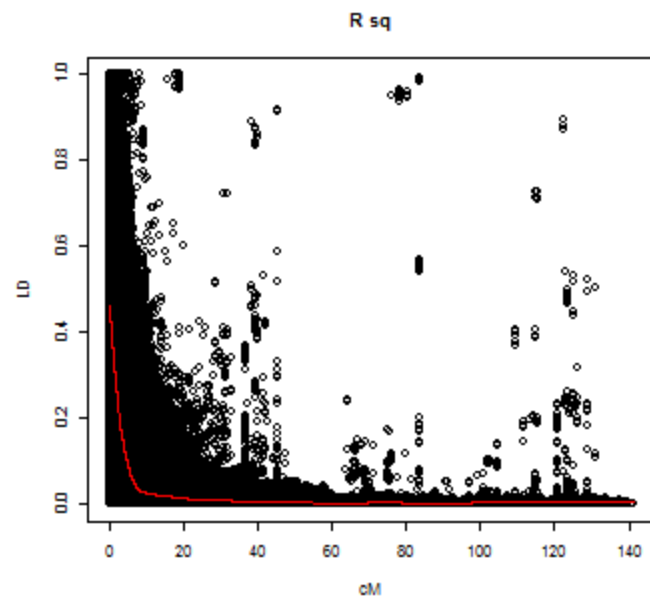
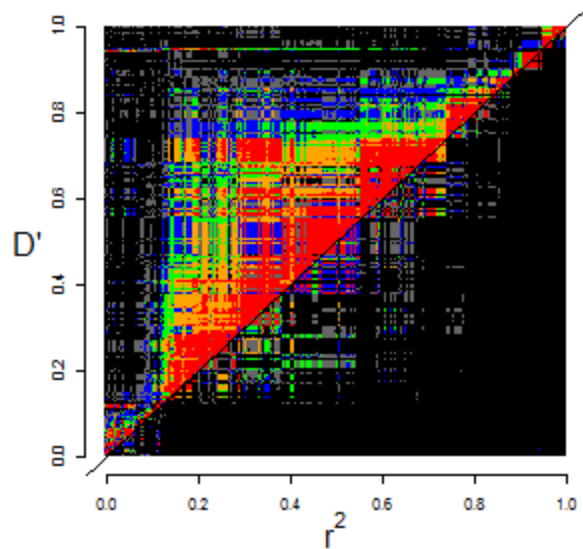




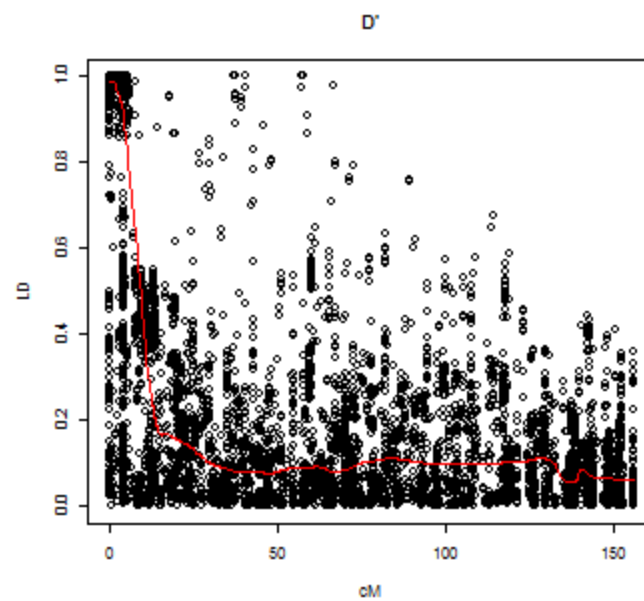
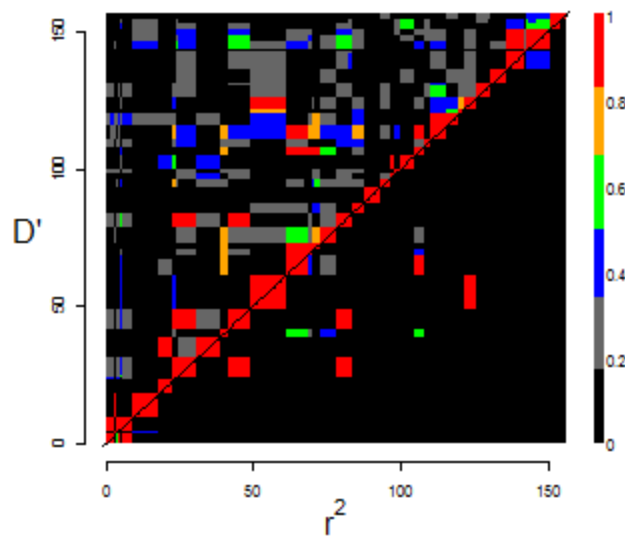
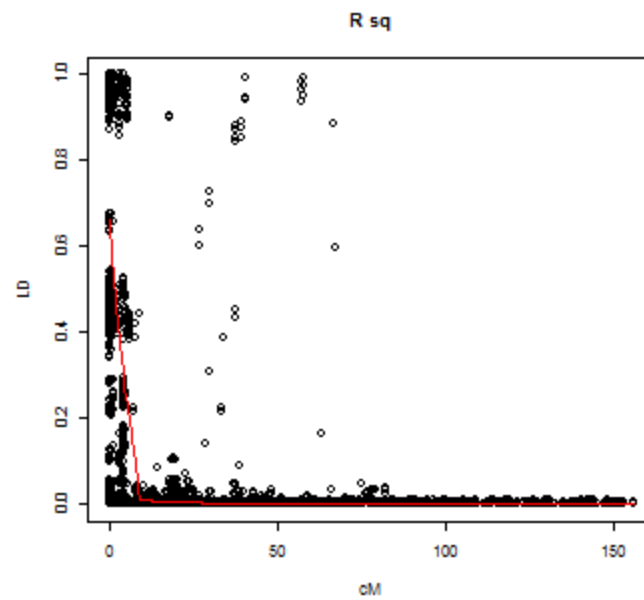
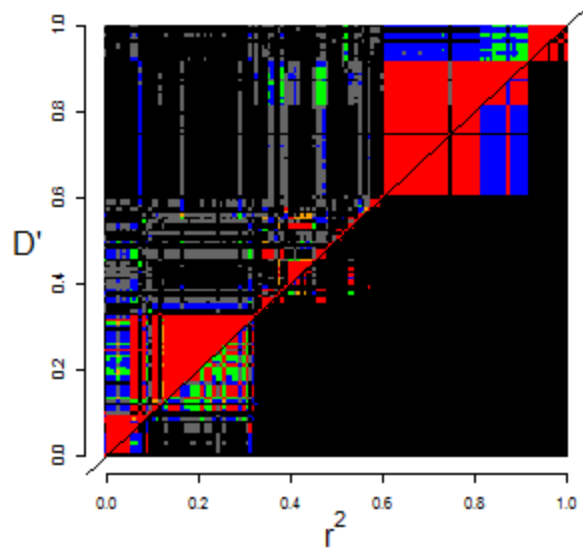
2D

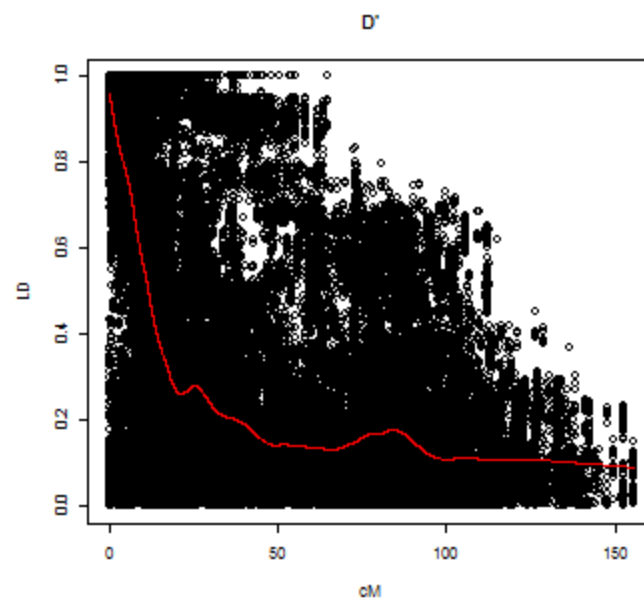
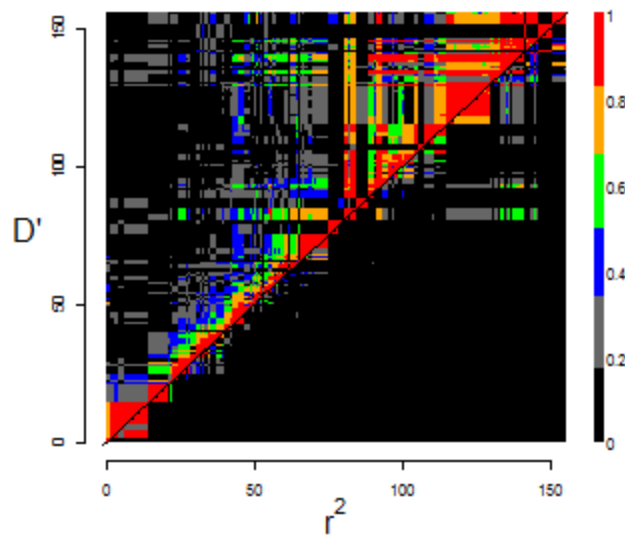
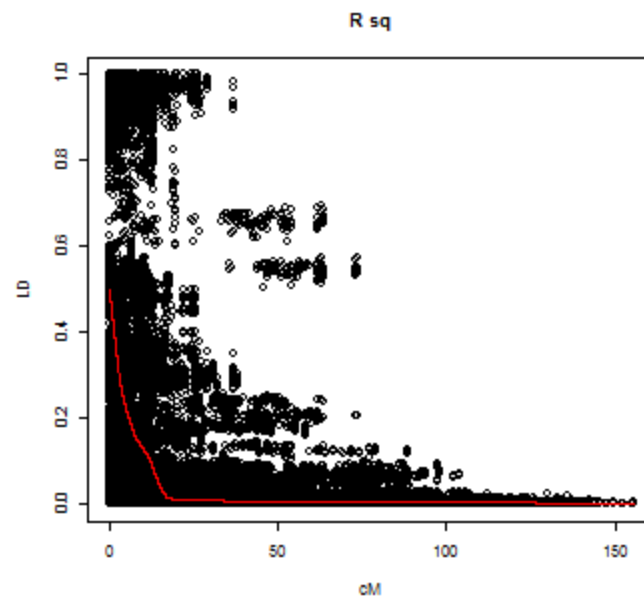
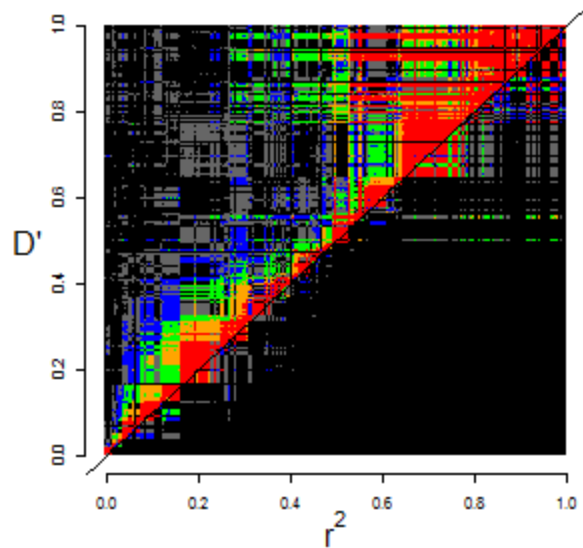


3A

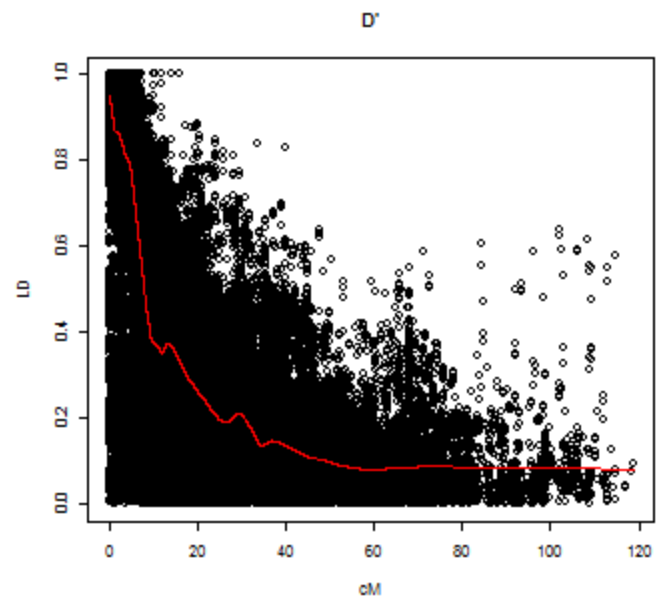
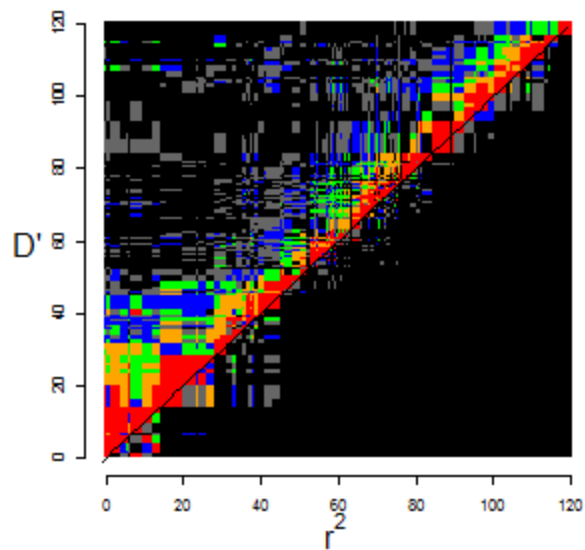
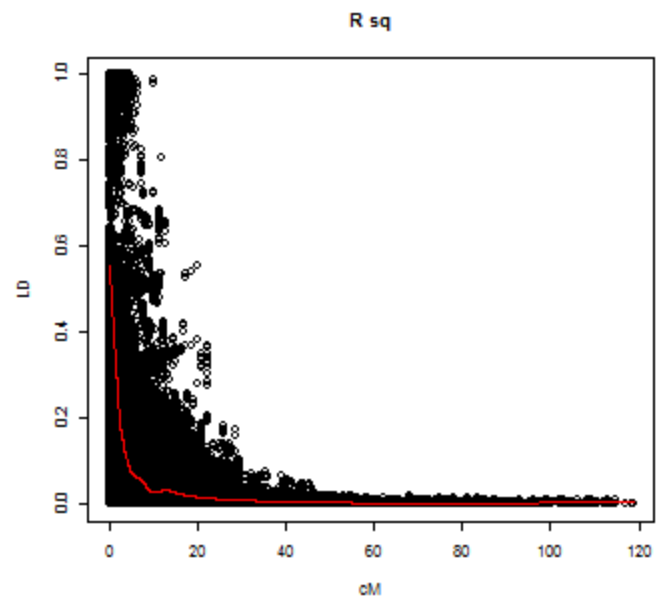
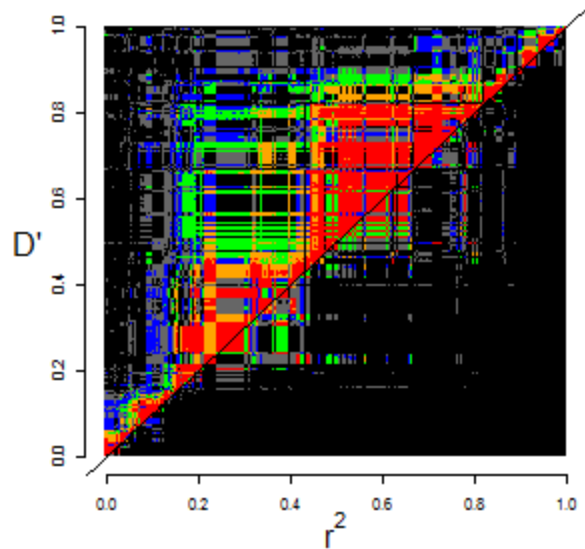


3B

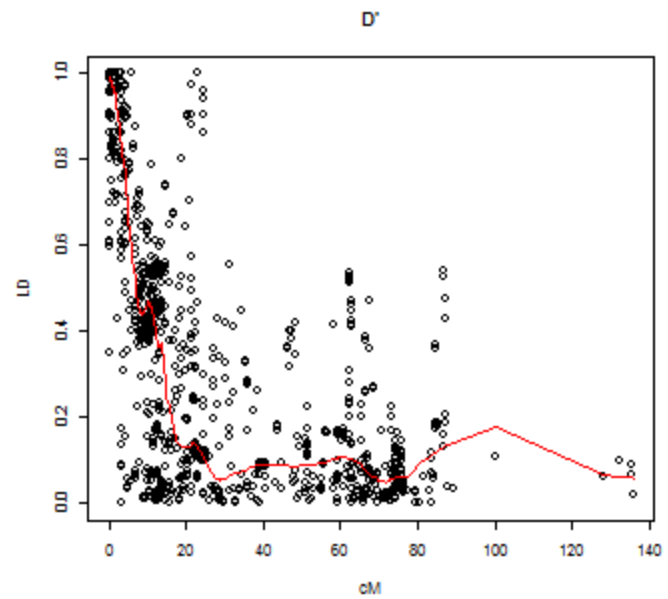
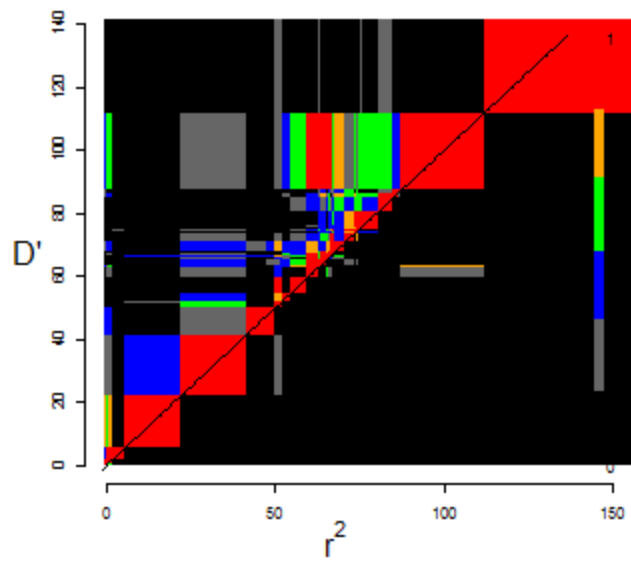
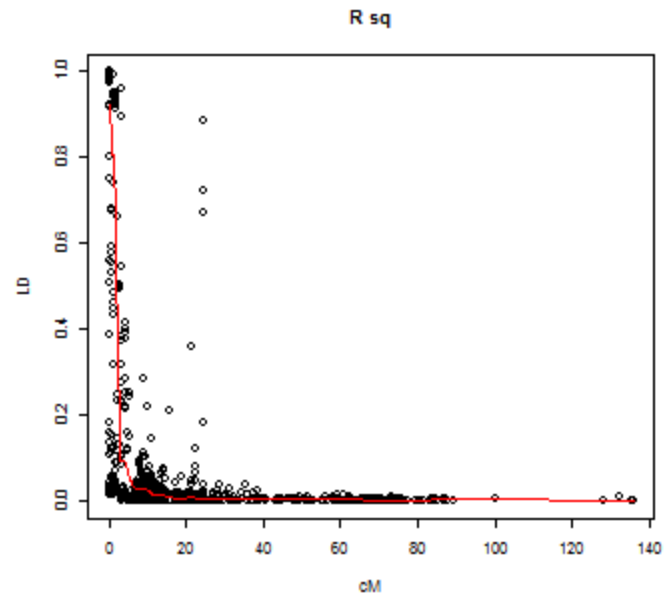
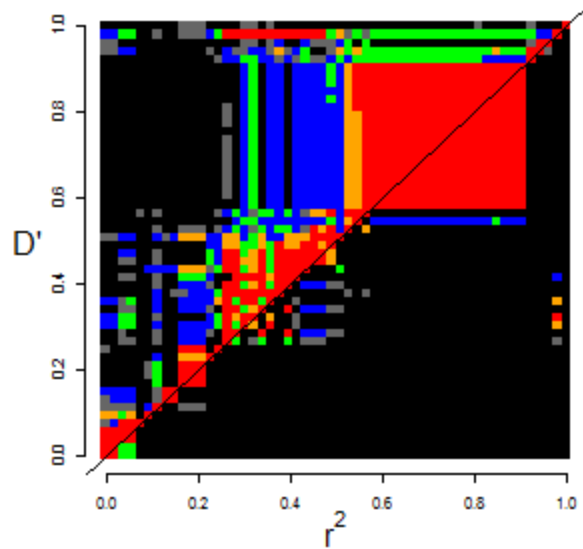


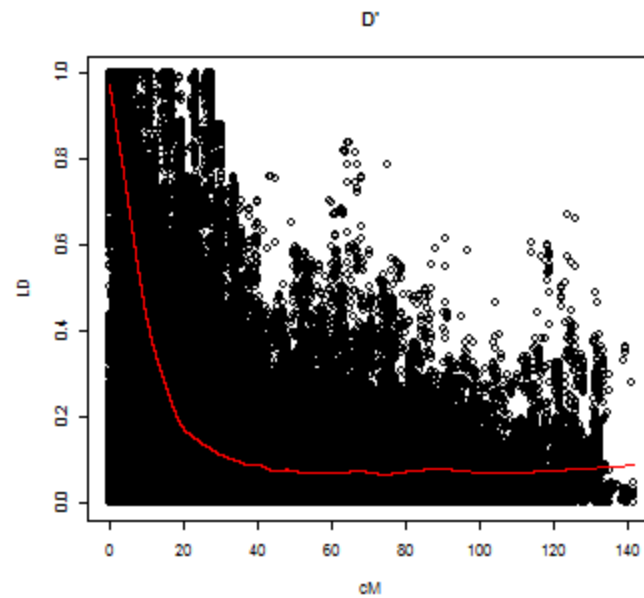
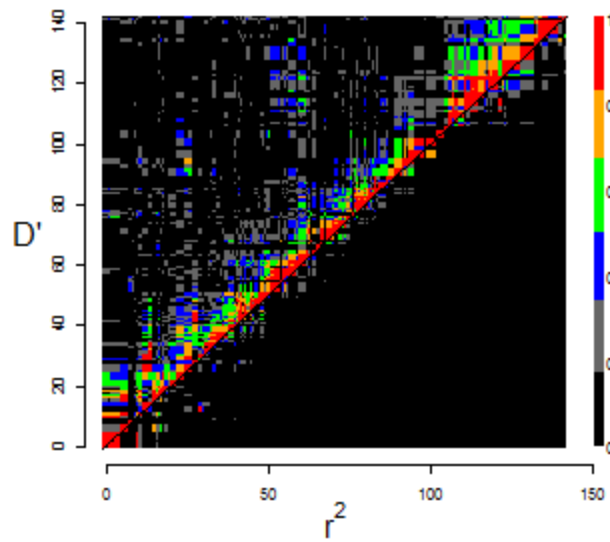
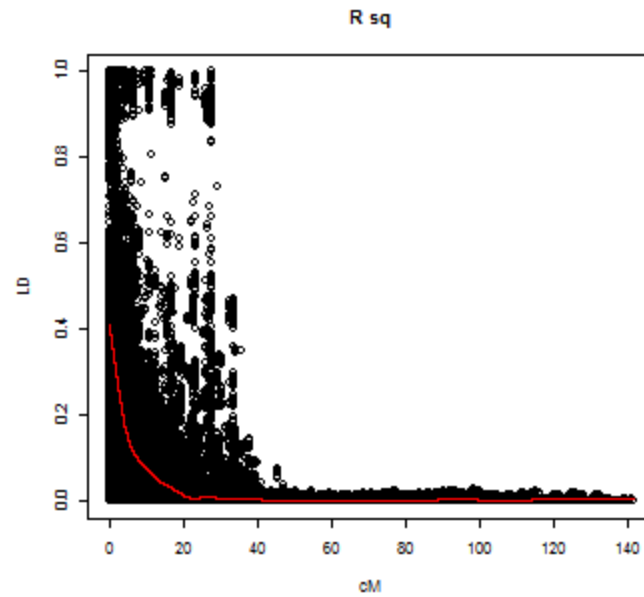
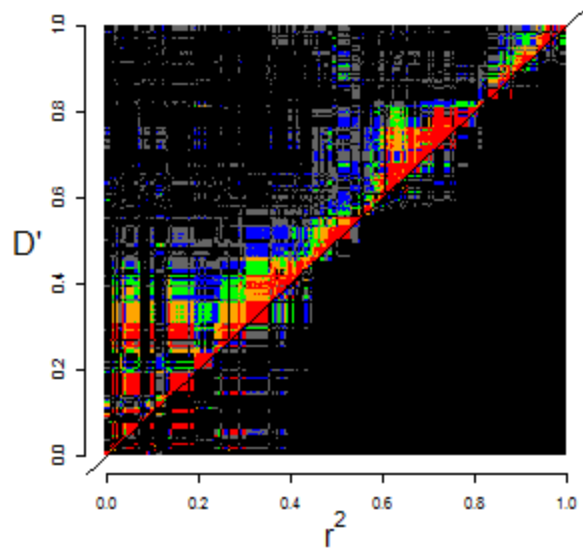


4A

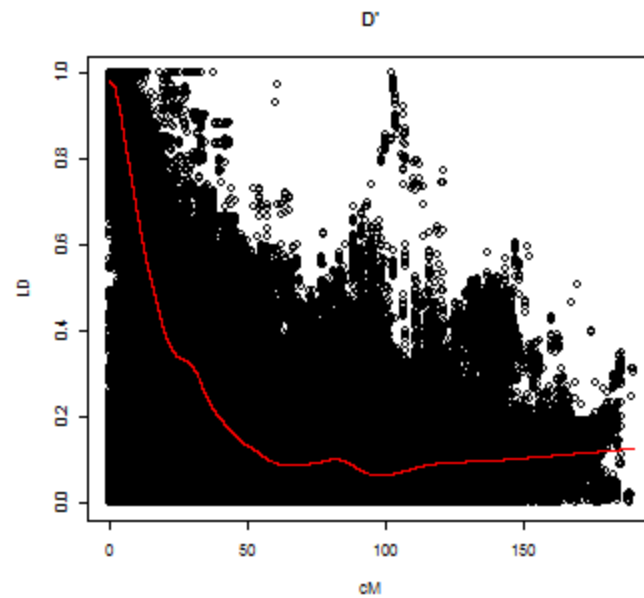
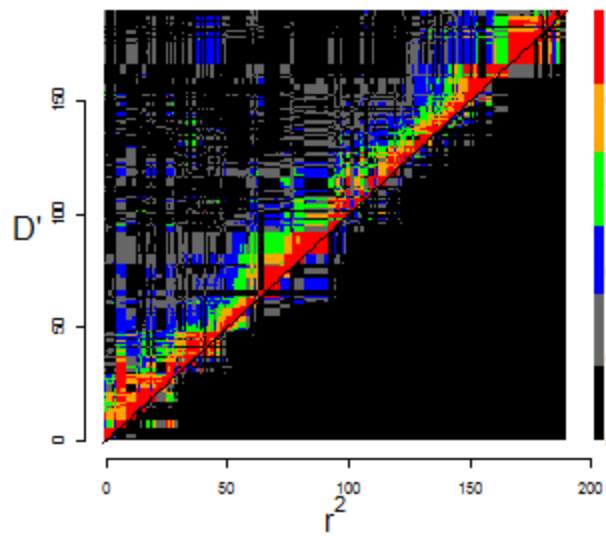
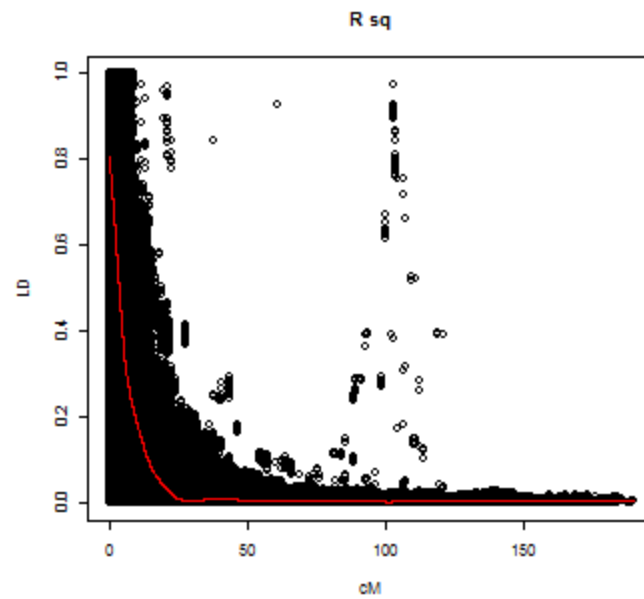
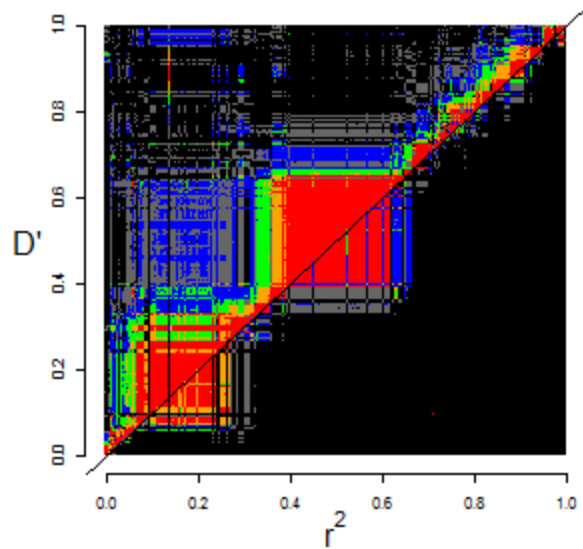


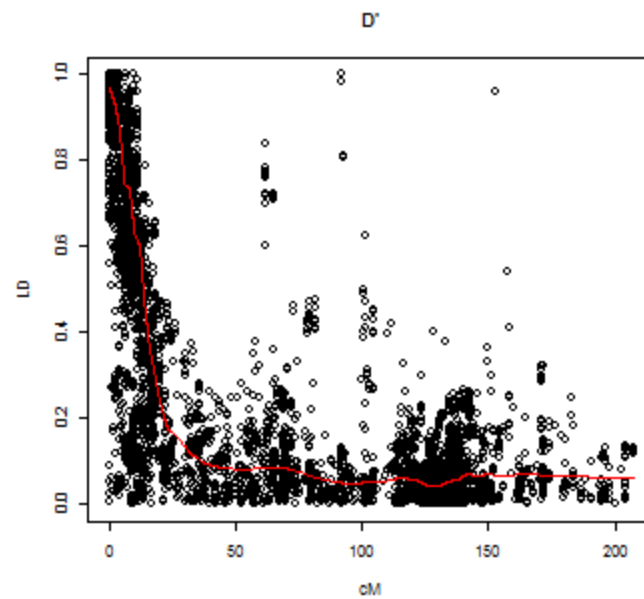
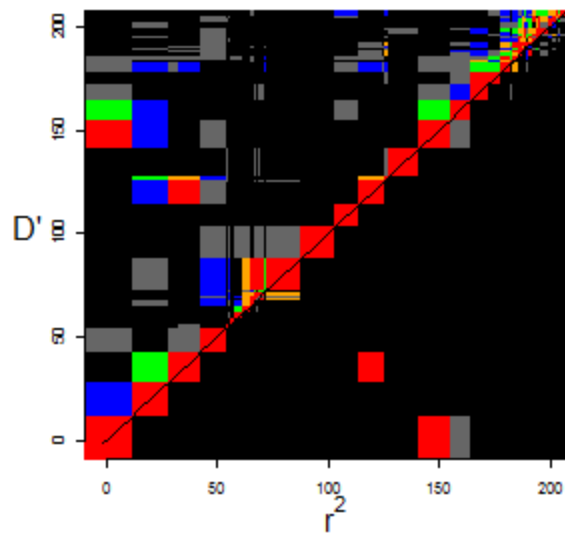
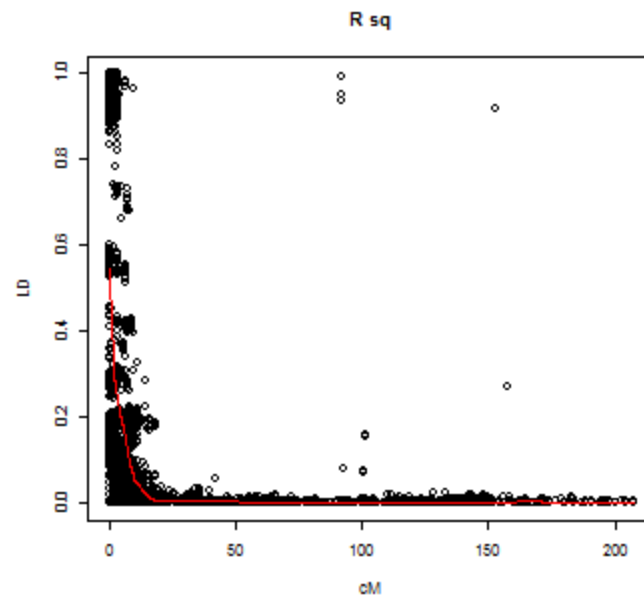
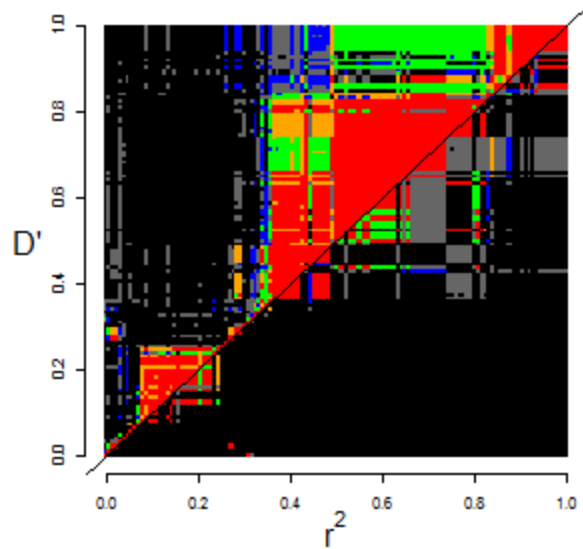
4B

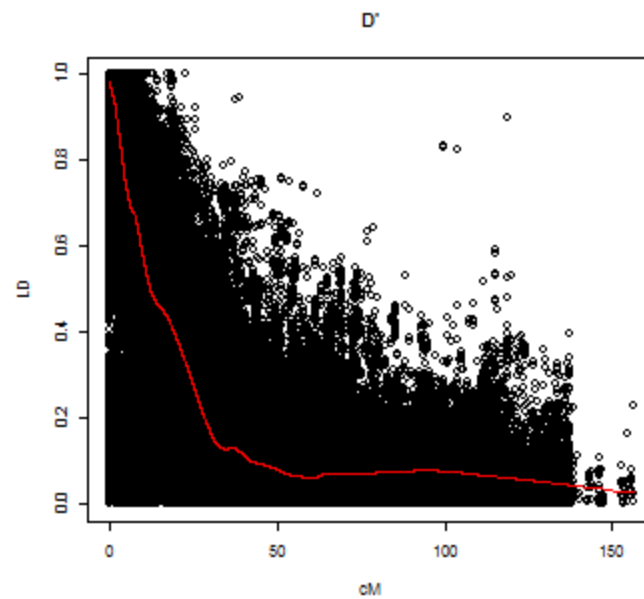
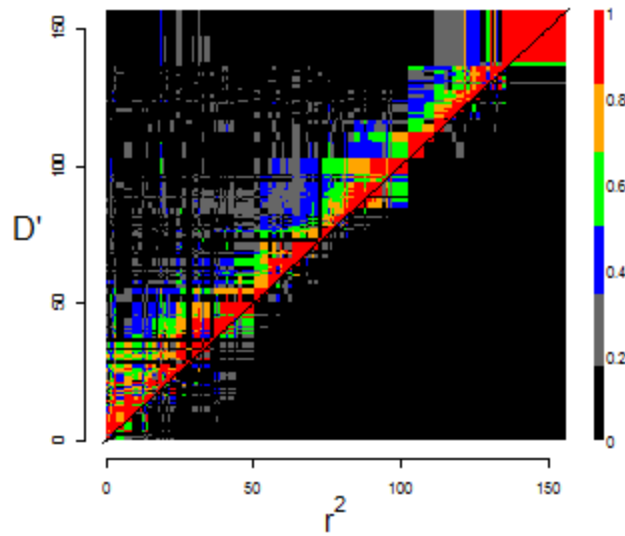
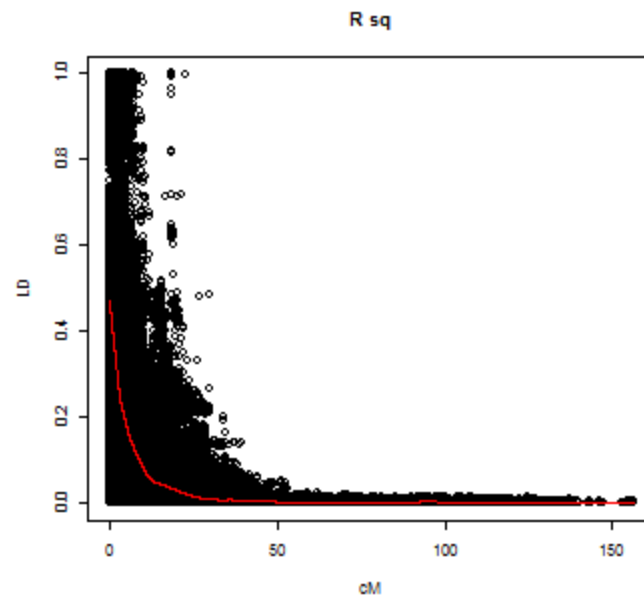
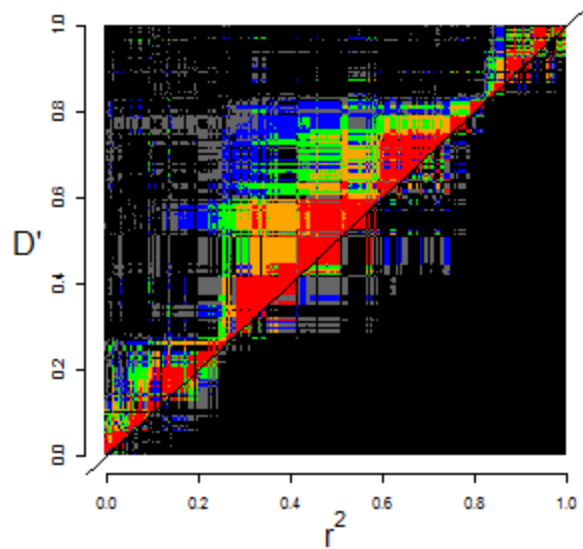




5A







6A

