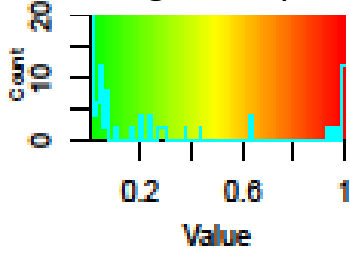
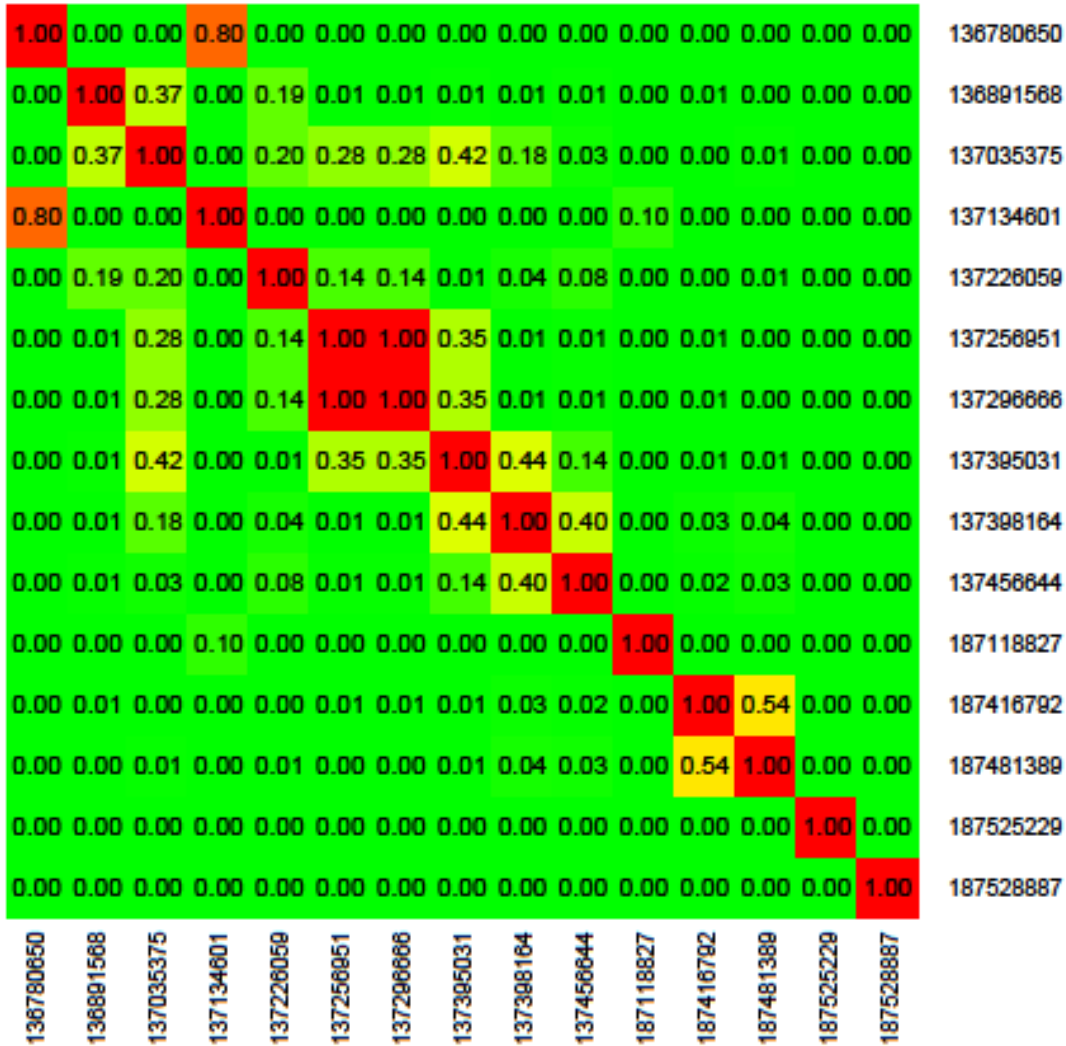


# Linkage disequilibrium $r^2$



Mb Position



Haplotype in Region 1.1

Haplotype in Region 1.2