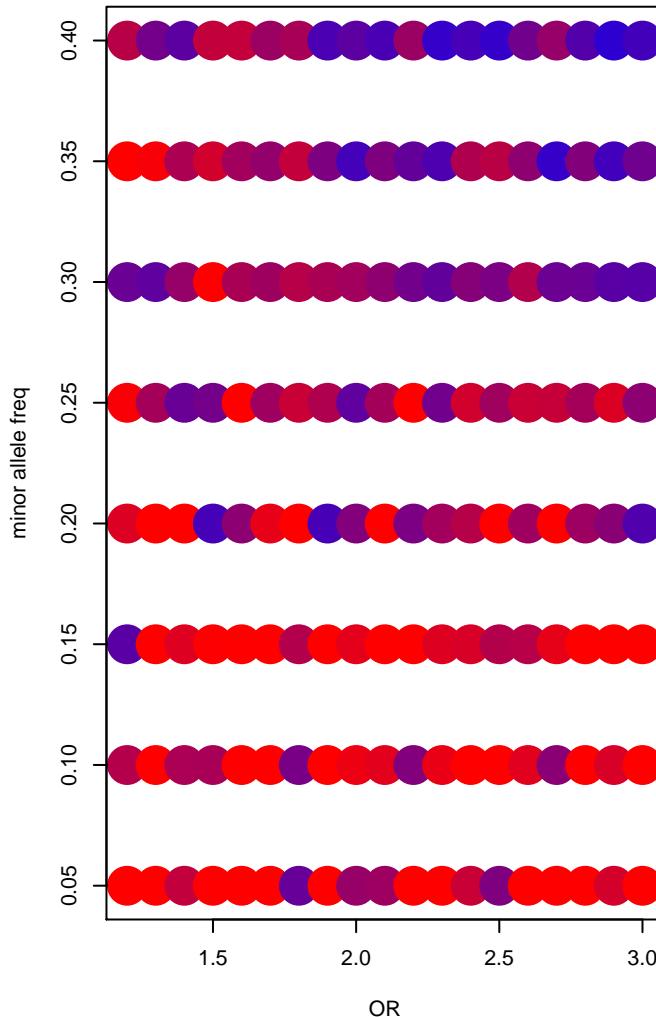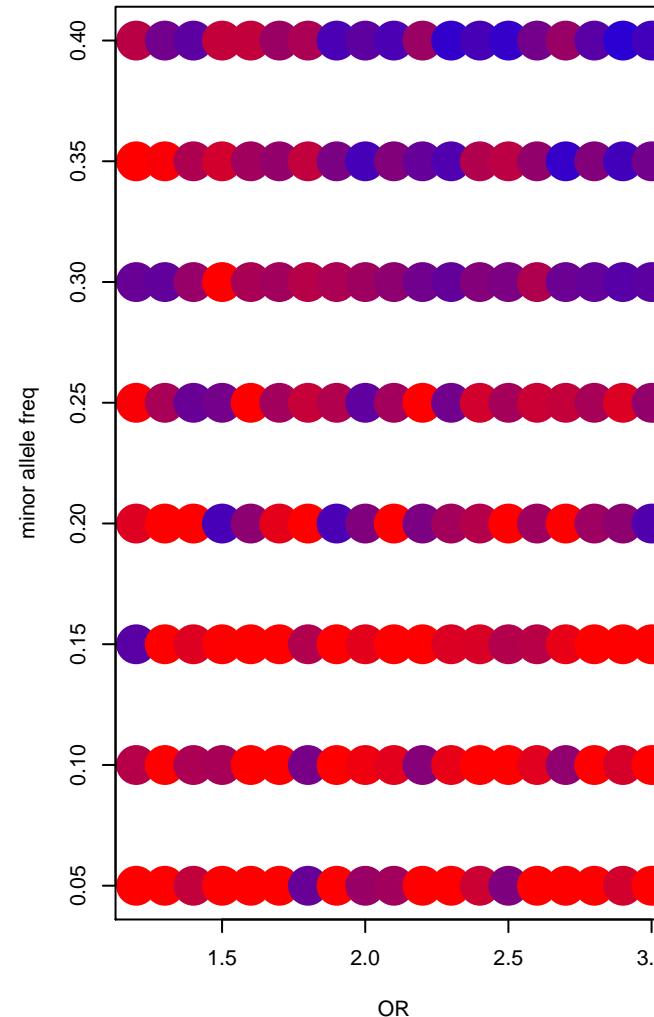Color coded correlations among data sets at each combination
of minor allele frequency and OR, cor<=0.6 => red, cor=1 => blue
N=5,000, n=2,000, pen.0=0.05, no.sets=40

cor(geno, trend), model='rec'   **cor(geno, allele), model='rec'**   **cor(trend, allele), model='rec'**

minor allele freq

OR