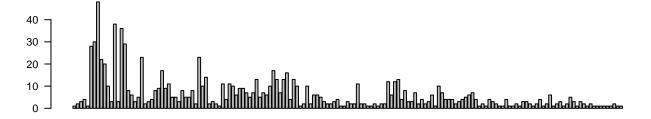
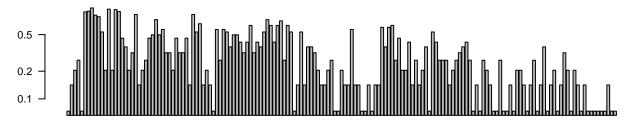
## n. mutated genes per sample



## p(one mutation in this pathway)



Expected number of samples with mutation in this pathway = 50

## Samples with mutations in this pathway = 113

