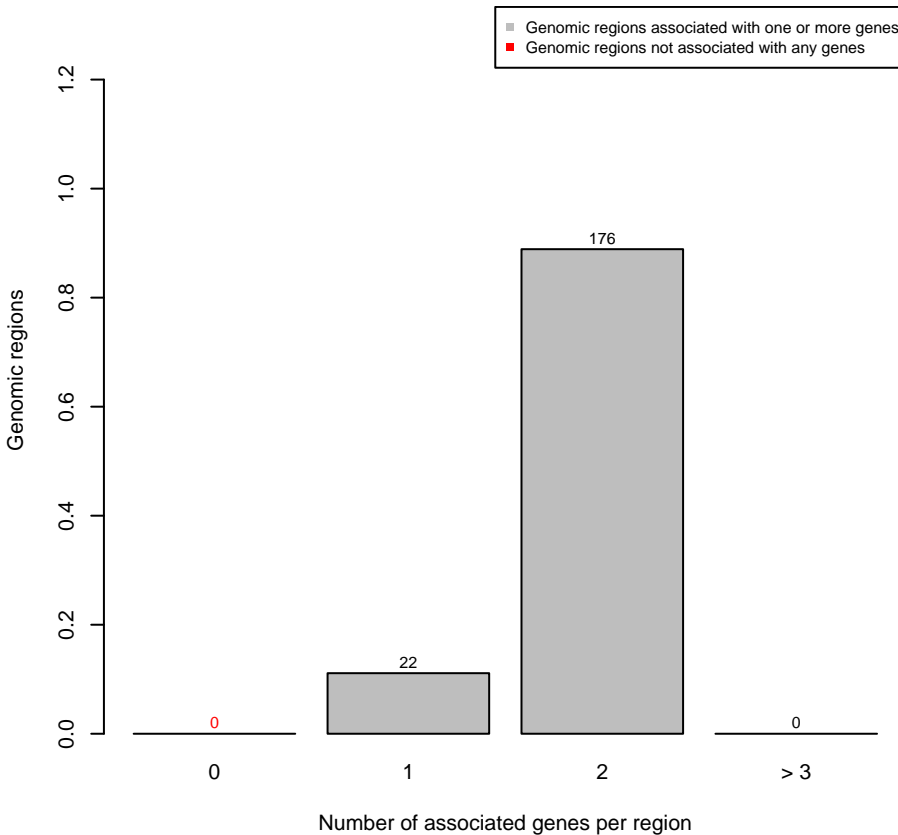
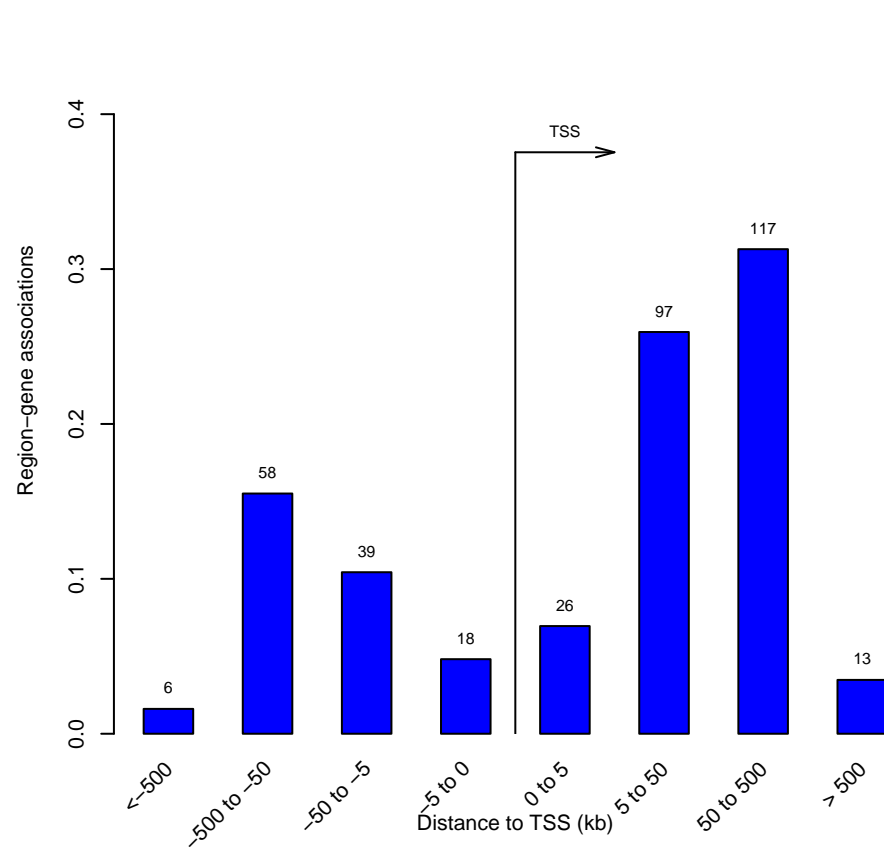


### Number of associated genes per region



### Binned by orientation and distance to TSS



### Binned by absolute distance to TSS

