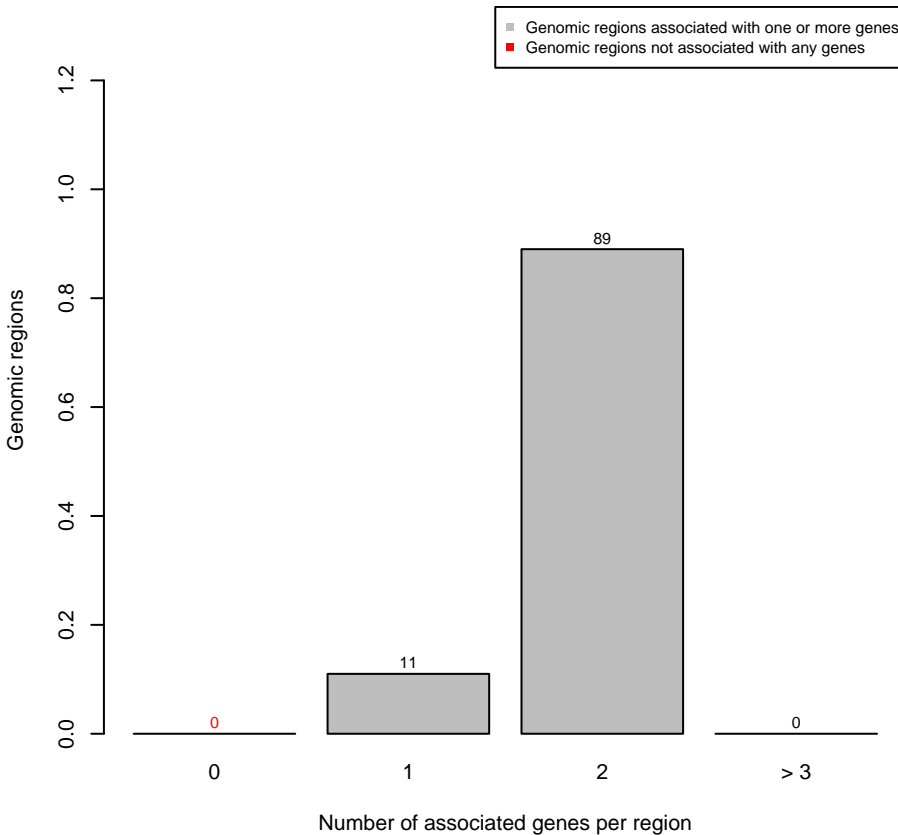
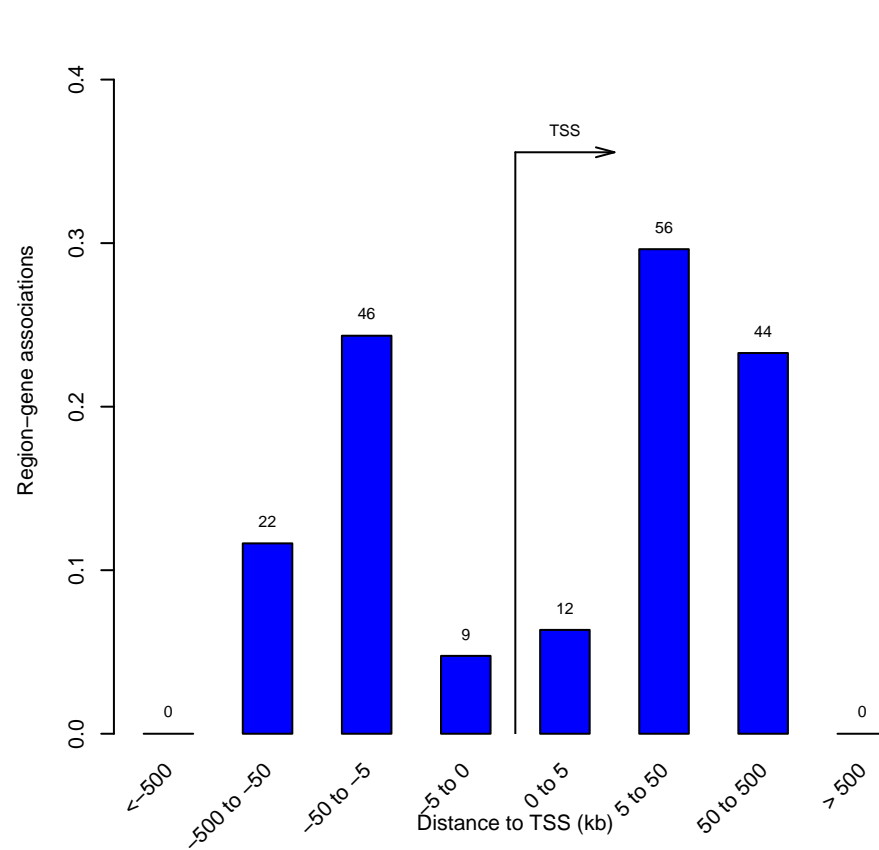


Number of associated genes per region



Binned by orientation and distance to TSS



Binned by absolute distance to TSS

