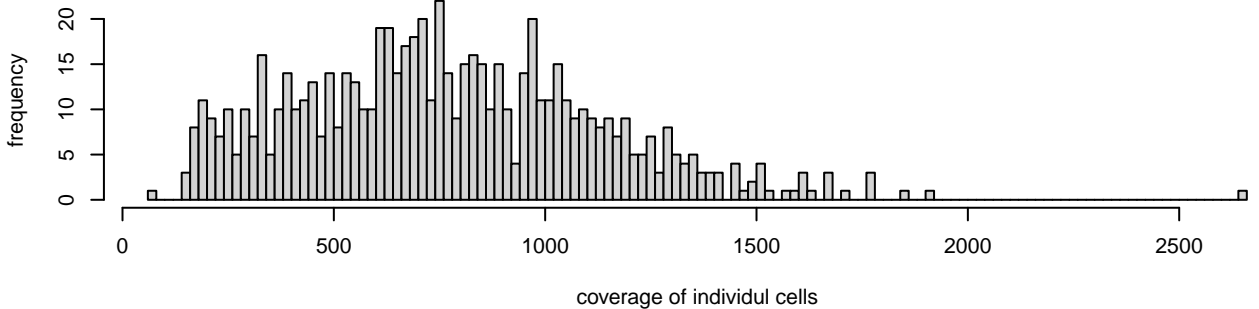
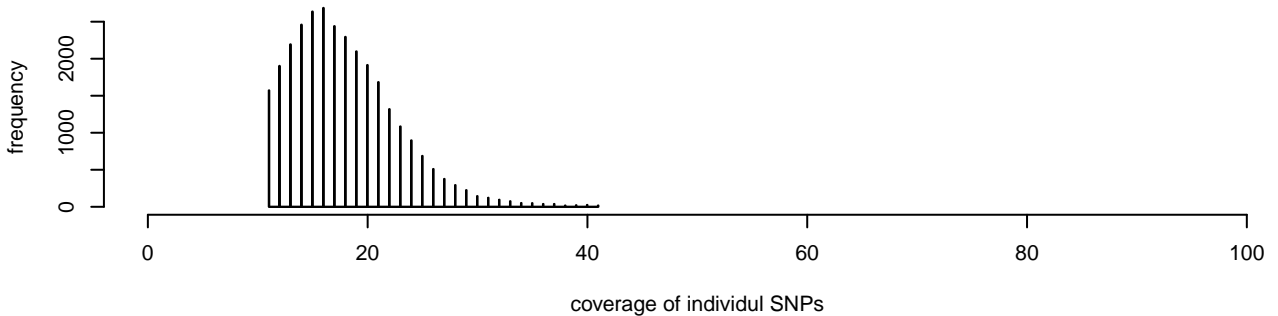


P5931 scDNAseq chr1 coverage (696 cells across 30000 SNPs)



P5931 scDNAseq chr1 coverage (30000 SNPs across 696 cells)



Histogram of VAF values

