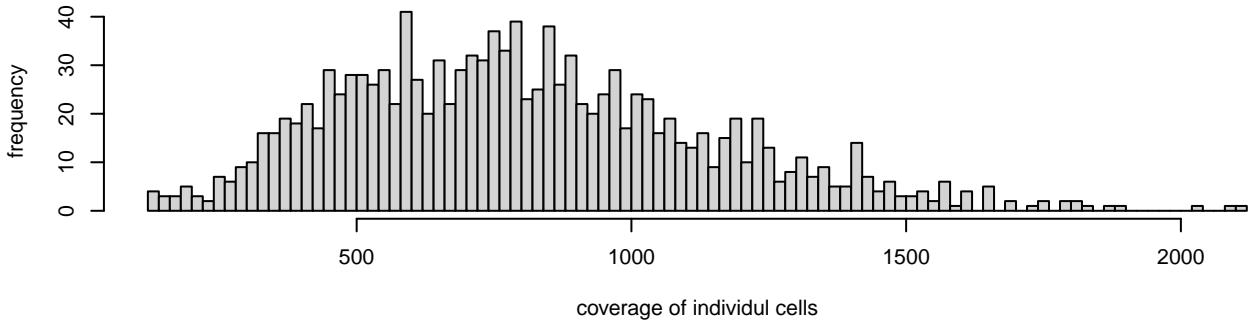
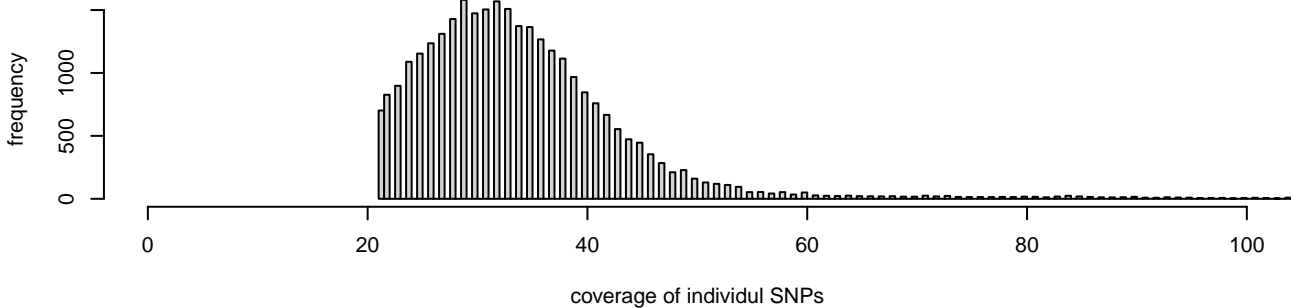


Sample scDNAseq chr3:0 coverage (1279 cells across 30000 SNPs)



Sample scDNAseq chr3:0 coverage (30000 SNPs across 1279 cells)



Histogram of VAF values

