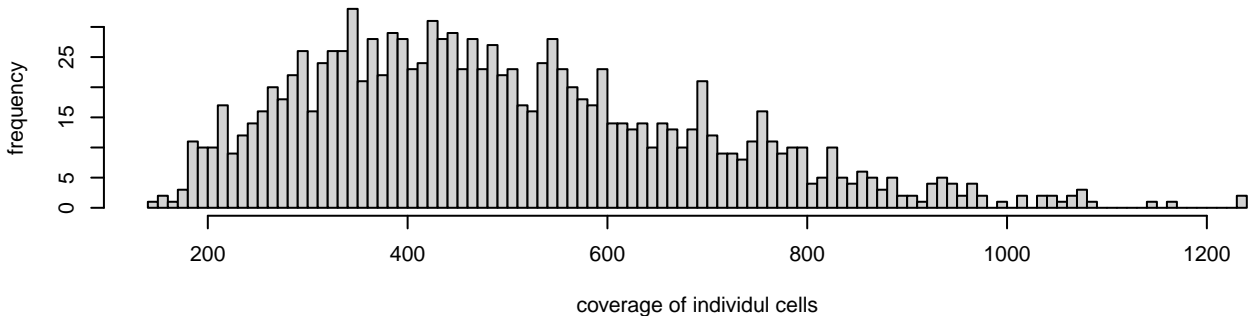
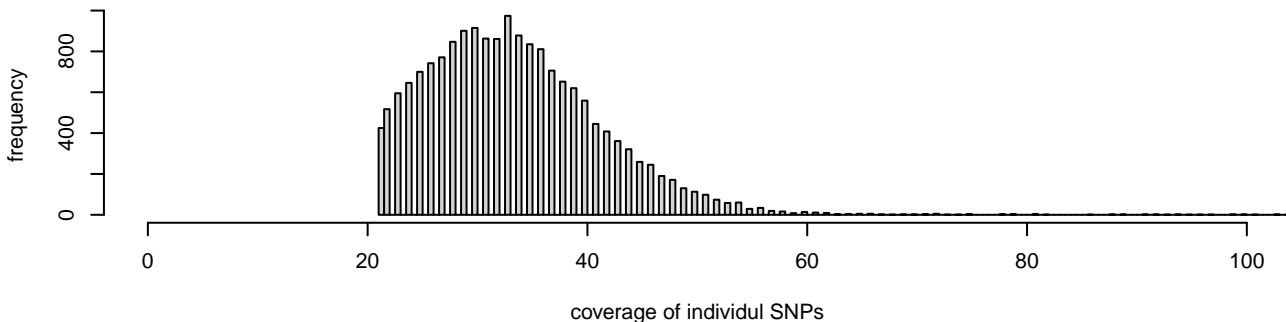


Sample scDNAseq chr20:44820000 coverage (1244 cells across 18034 SNPs)



Sample scDNAseq chr20:44820000 coverage (18034 SNPs across 1244 cells)



Histogram of VAF values

