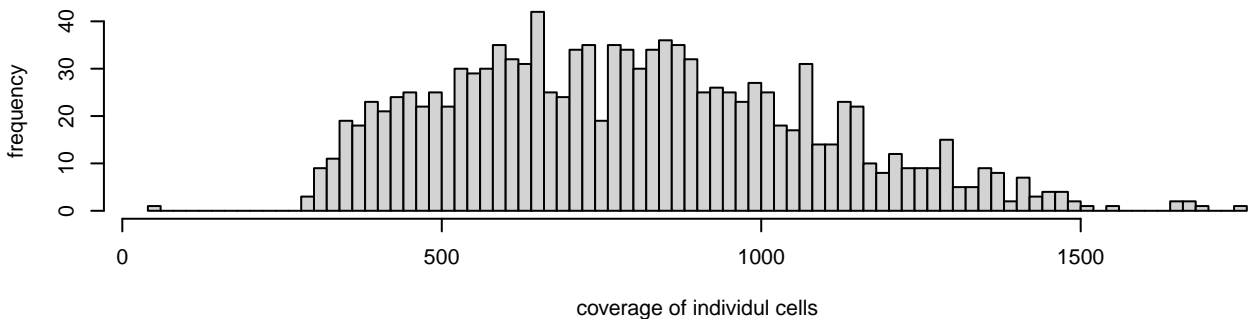
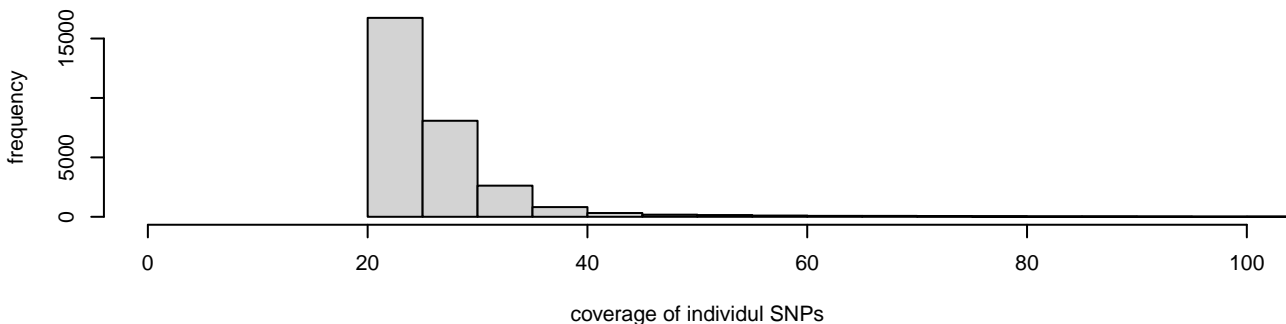


Sample scDNAseq chr10:0 coverage (1244 cells across 30000 SNPs)



Sample scDNAseq chr10:0 coverage (30000 SNPs across 1244 cells)



Histogram of VAF values

