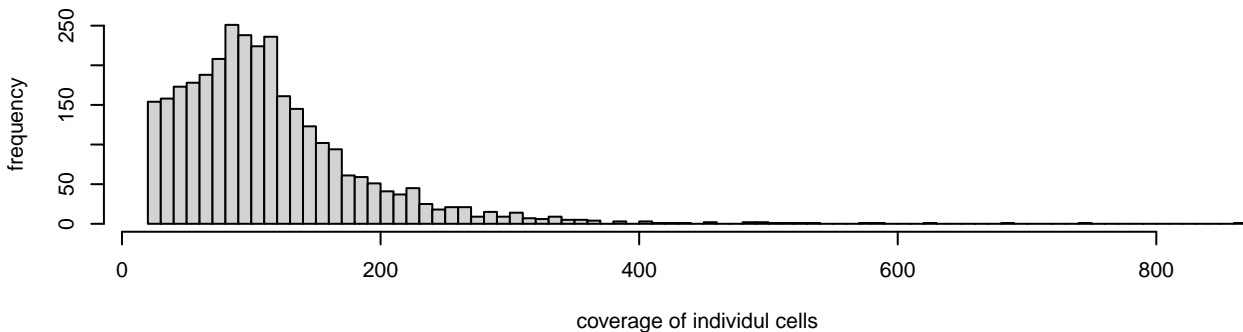
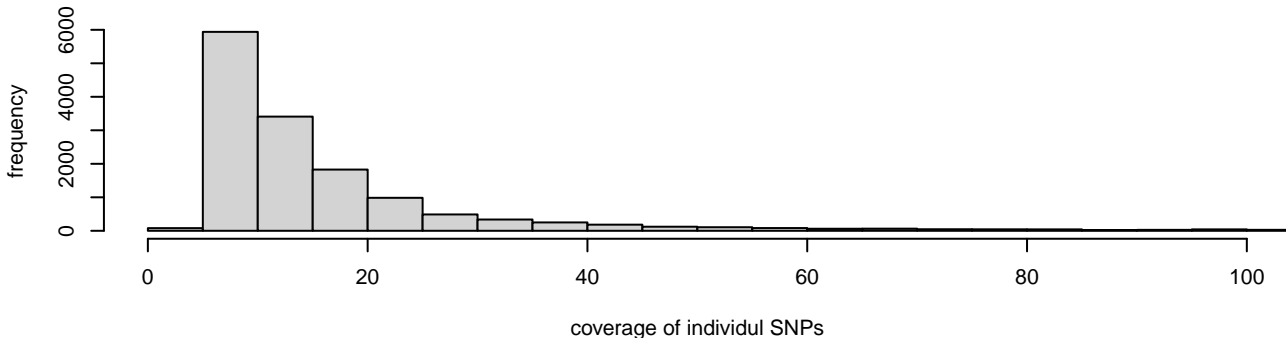


Sample scATACseq chr20:44820000 coverage (3120 cells across 14696 SNPs)



Sample scATACseq chr20:44820000 coverage (14696 SNPs across 3120 cells)



Histogram of VAF values

