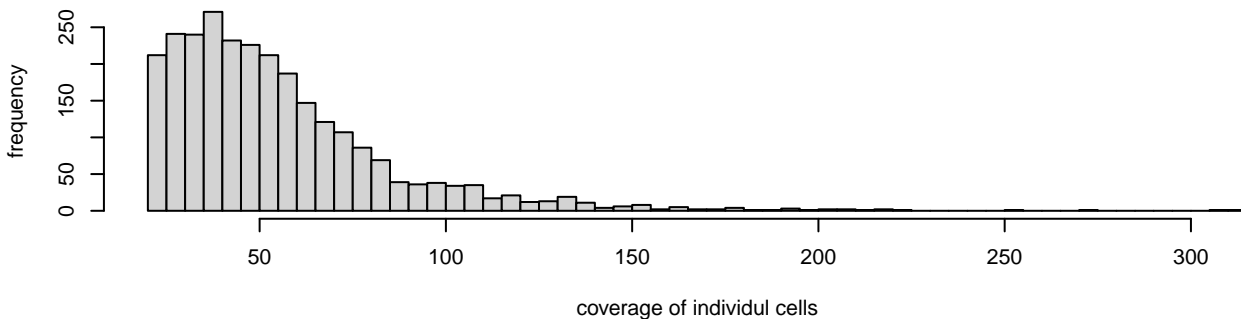
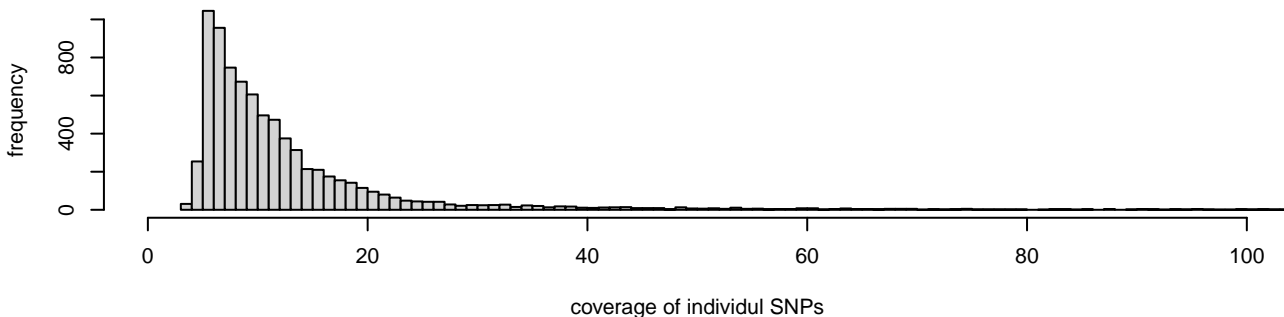


Sample scATACseq chr3:147340000 coverage (2677 cells across 8117 SNPs)



Sample scATACseq chr3:147340000 coverage (8117 SNPs across 2677 cells)



Histogram of VAF values

