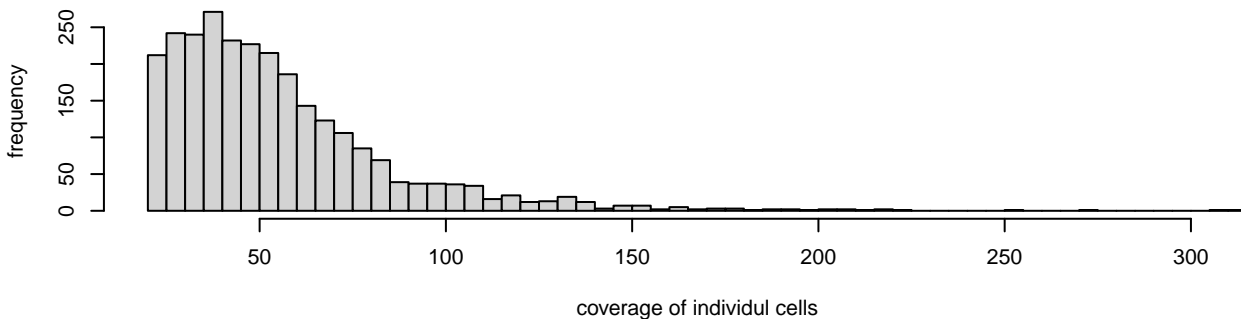
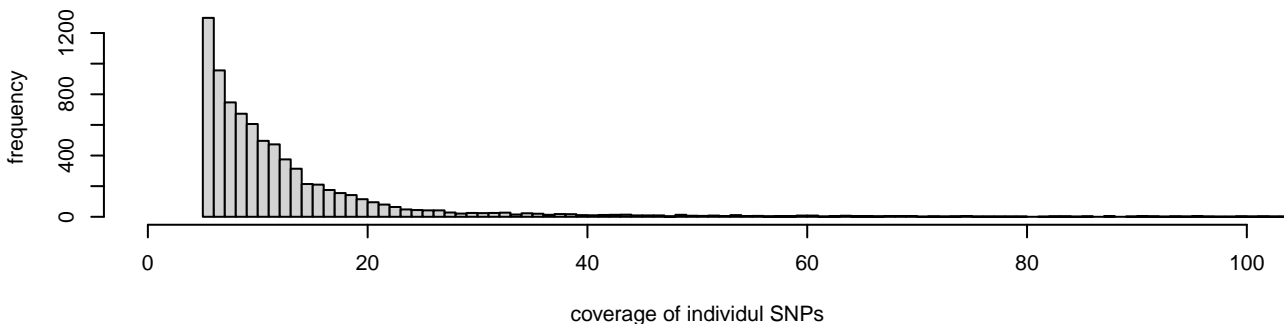


SNU601 scATACseq chr3:147340000 coverage (2677 cells across 8086 SNPs)



SNU601 scATACseq chr3:147340000 coverage (8086 SNPs across 2677 cells)



Histogram of VAF values

