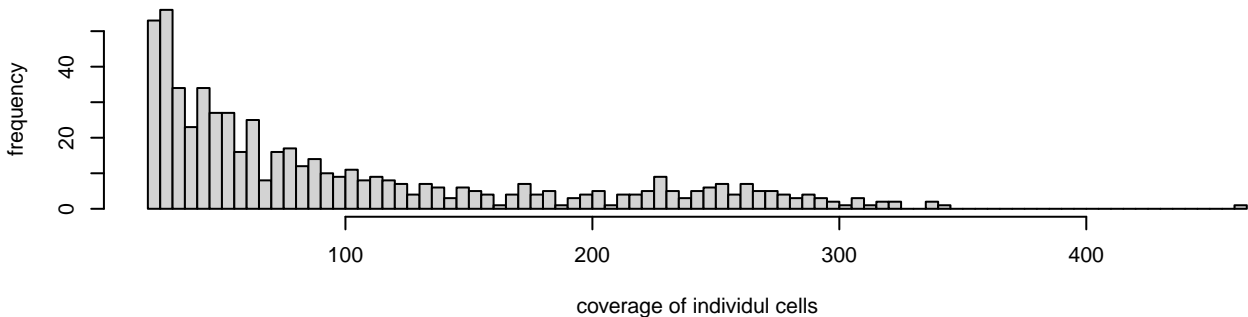
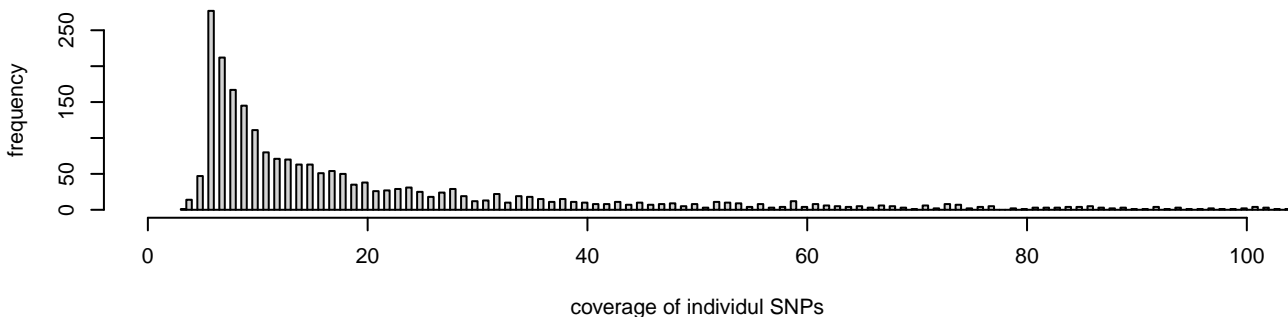


Sample scATACseq chr15:23467210 coverage (592 cells across 2326 SNPs)



Sample scATACseq chr15:23467210 coverage (2326 SNPs across 592 cells)



Histogram of VAF values

