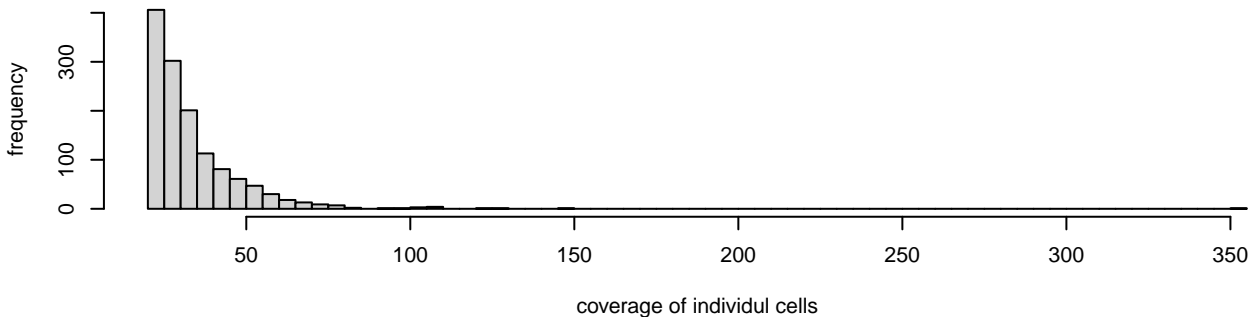
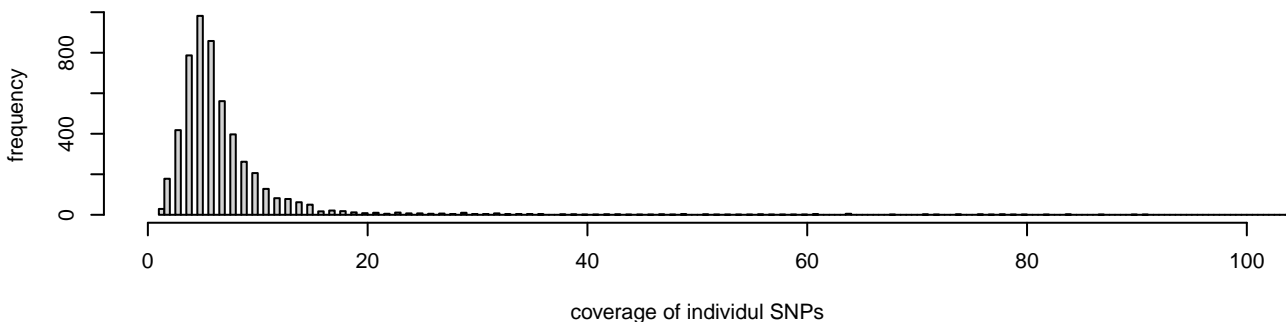


Sample scATACseq chr13:80640000 coverage (1303 cells across 5332 SNPs)



Sample scATACseq chr13:80640000 coverage (5332 SNPs across 1303 cells)



Histogram of VAF values

