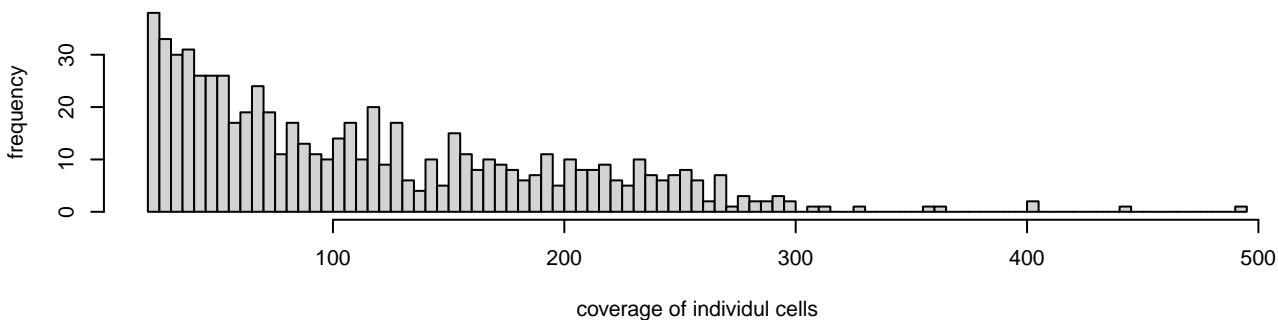
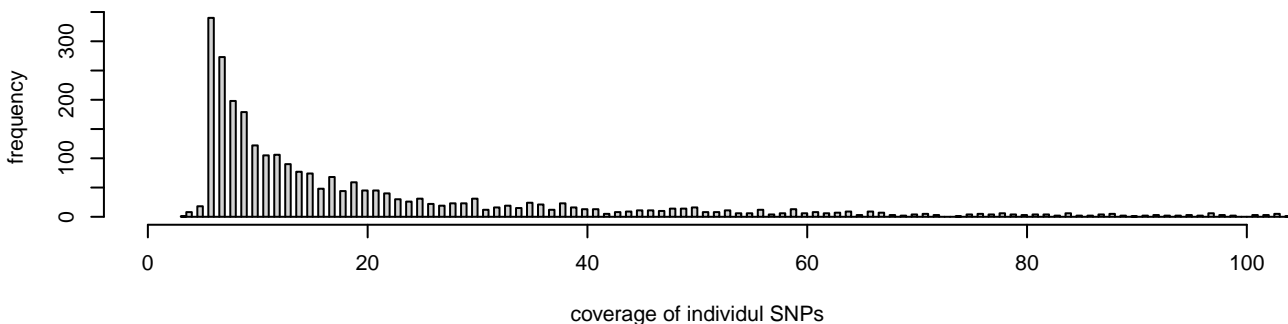


Sample scATACseq chr3:97670875 coverage (674 cells across 2807 SNPs)



Sample scATACseq chr3:97670875 coverage (2807 SNPs across 674 cells)



Histogram of VAF values

