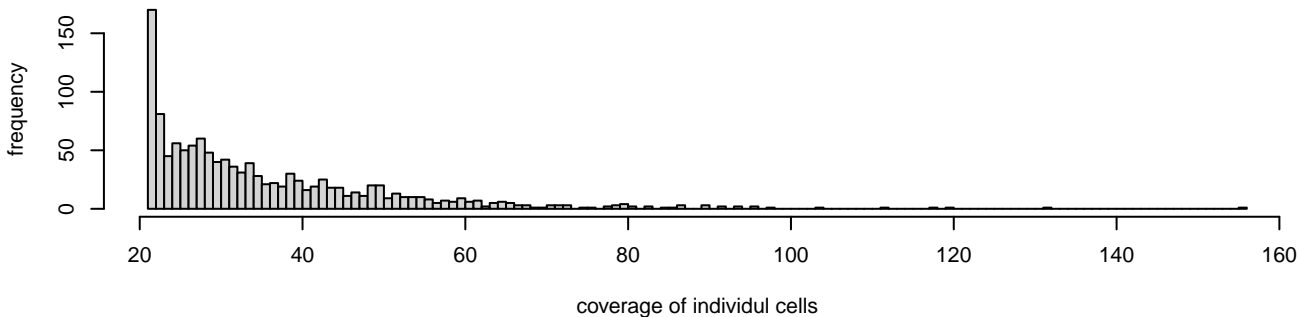
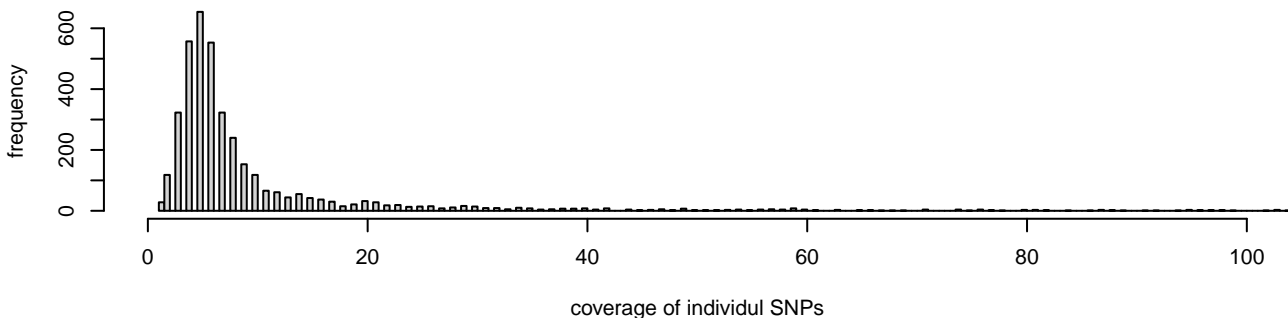


Sample scATACseq chr4:57600000 coverage (1239 cells across 3879 SNPs)



Sample scATACseq chr4:57600000 coverage (3879 SNPs across 1239 cells)



Histogram of VAF values

