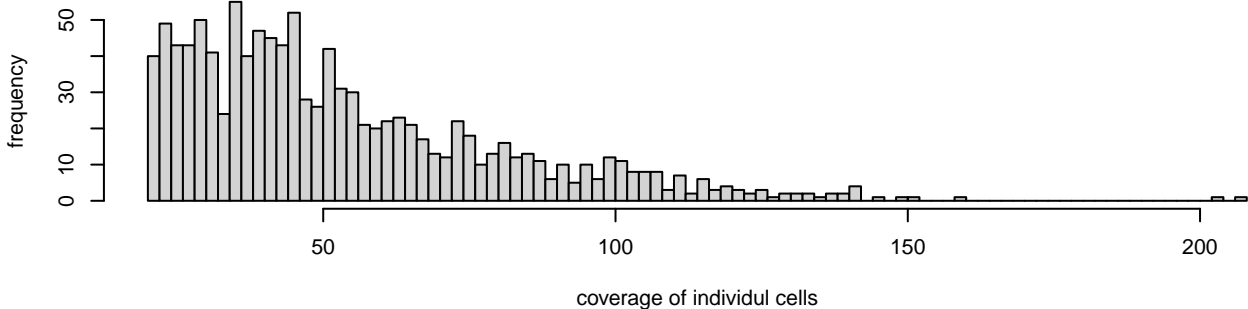
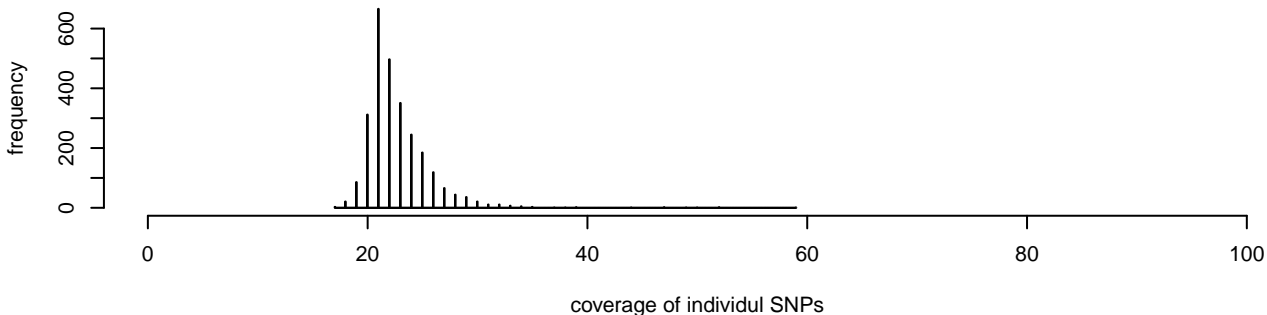


Sample scDNAseq chr4:57600000 coverage (1132 cells across 2701 SNPs)



Sample scDNAseq chr4:57600000 coverage (2701 SNPs across 1132 cells)



Histogram of VAF values

