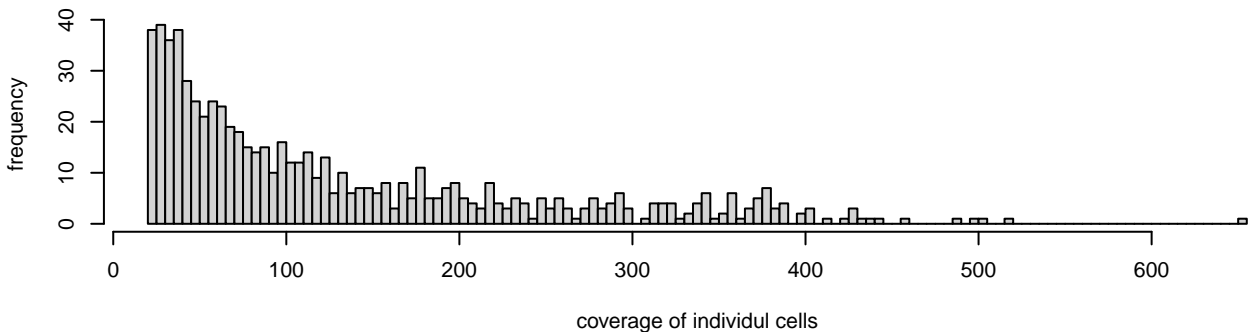
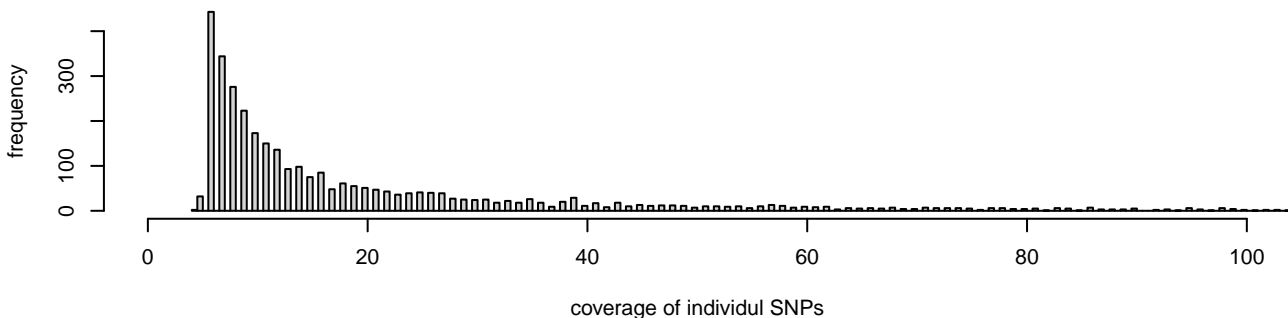


**Sample scATACseq chr12:0 coverage (695 cells across 3469 SNPs)**



**Sample scATACseq chr12:0 coverage (3469 SNPs across 695 cells)**



**Histogram of VAF values**

