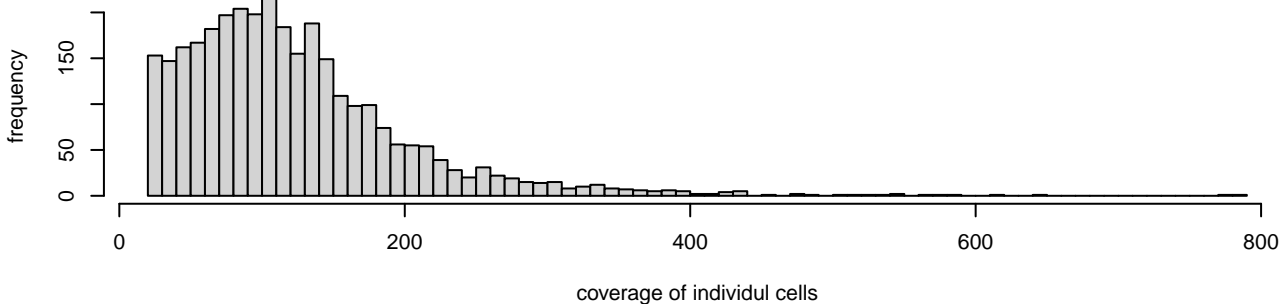
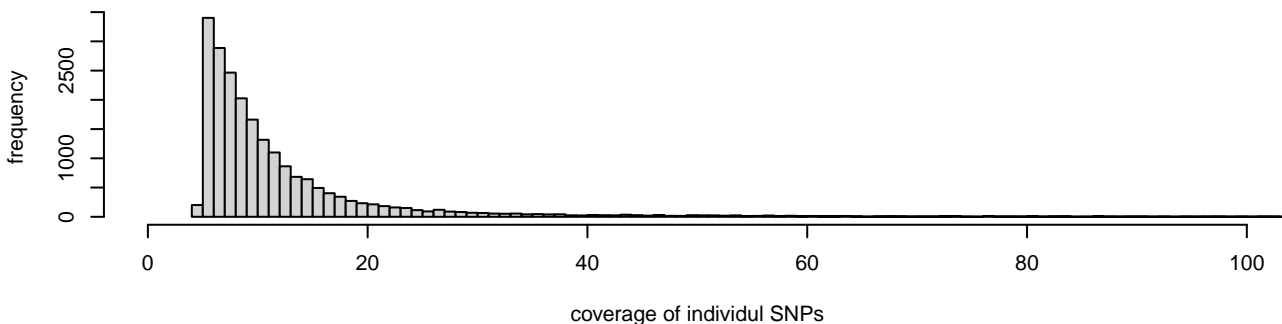


**Sample scATACseq chr3:96160000 coverage (3148 cells across 21931 SNPs)**



**Sample scATACseq chr3:96160000 coverage (21931 SNPs across 3148 cells)**



**Histogram of VAF values**

