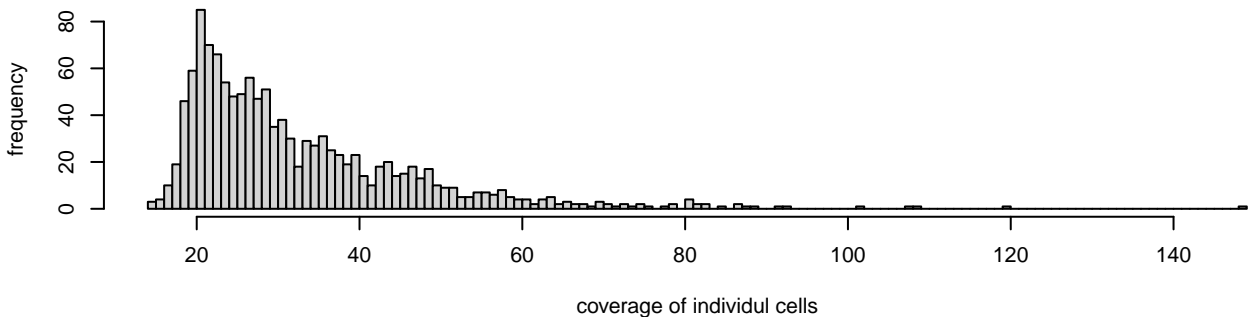
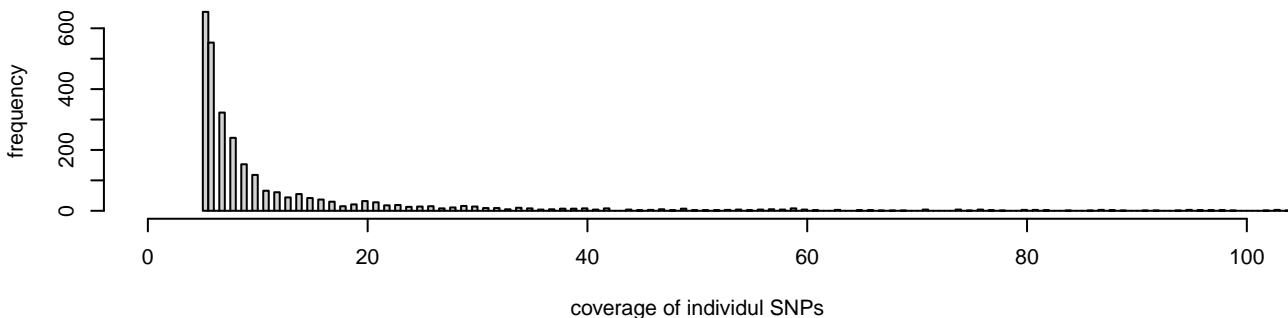


**SNU601 scATACseq chr4:57600000 coverage (1239 cells across 2853 SNPs)**



**SNU601 scATACseq chr4:57600000 coverage (2853 SNPs across 1239 cells)**



**Histogram of VAF values**

