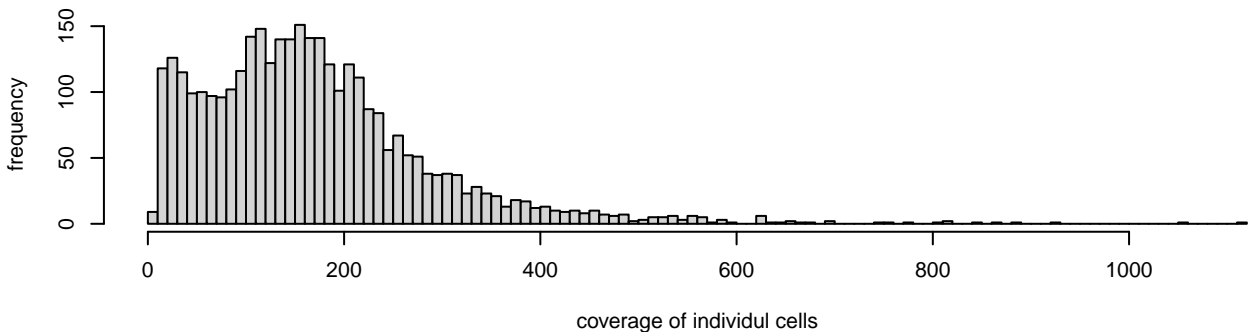
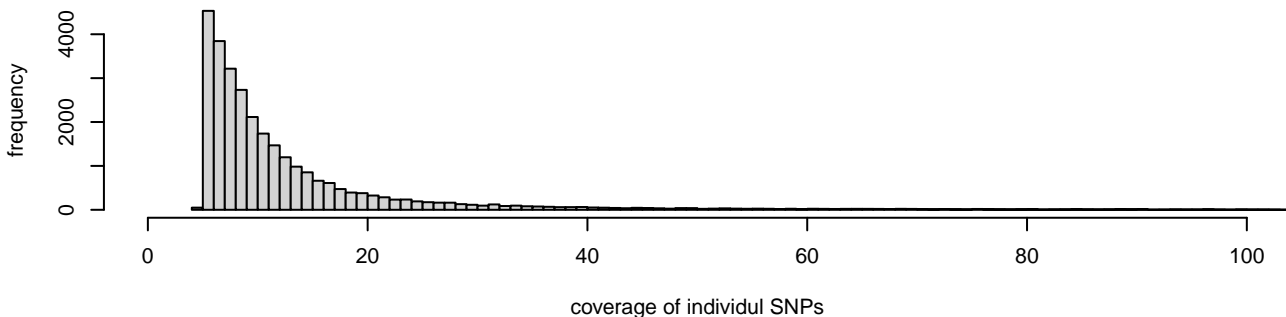


**Sample scATACseq chr7:0 coverage (3405 cells across 30000 SNPs)**



**Sample scATACseq chr7:0 coverage (30000 SNPs across 3405 cells)**



**Histogram of VAF values**

