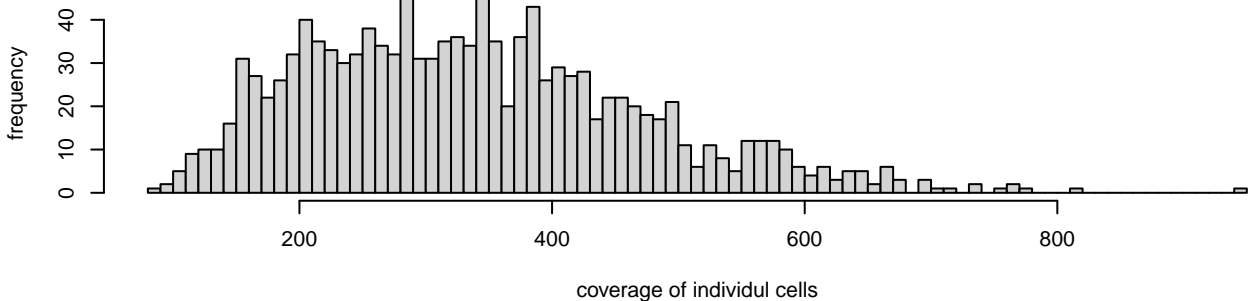
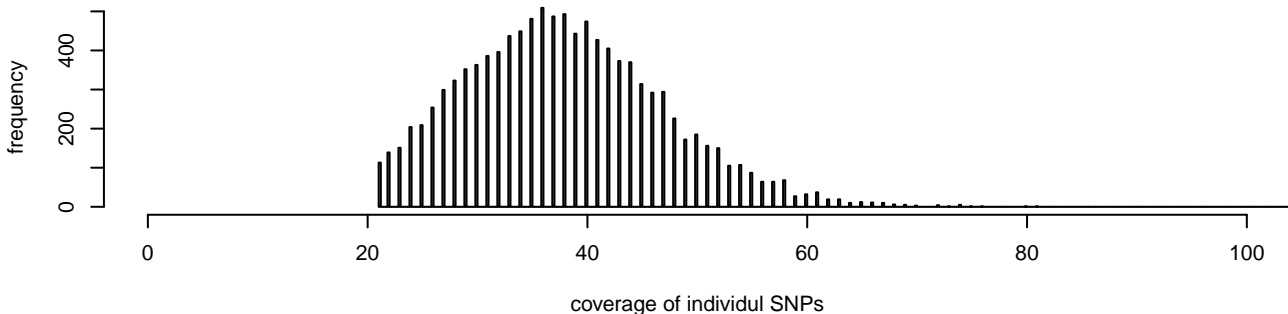


**Sample scDNAseq chr3:147340000 coverage (1244 cells across 11029 SNPs)**



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**Histogram of VAF values**

