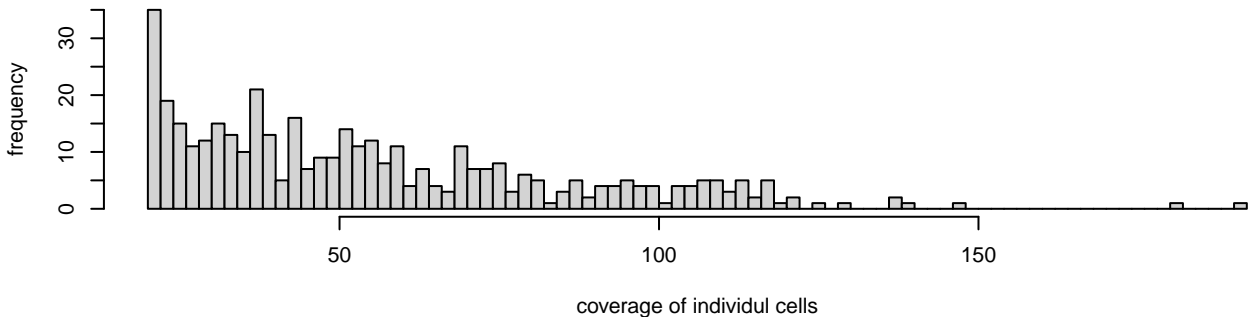
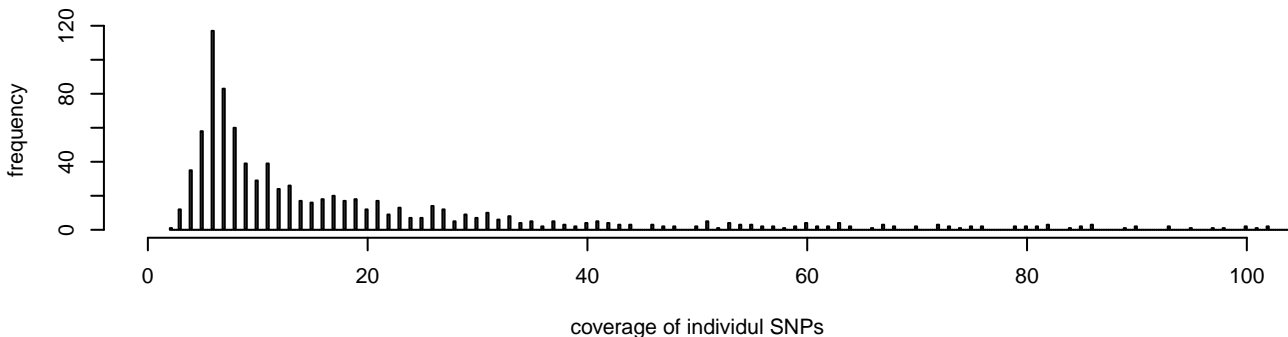


**Sample scATACseq chr21:16437159 coverage (408 cells across 938 SNPs)**



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**Histogram of VAF values**

