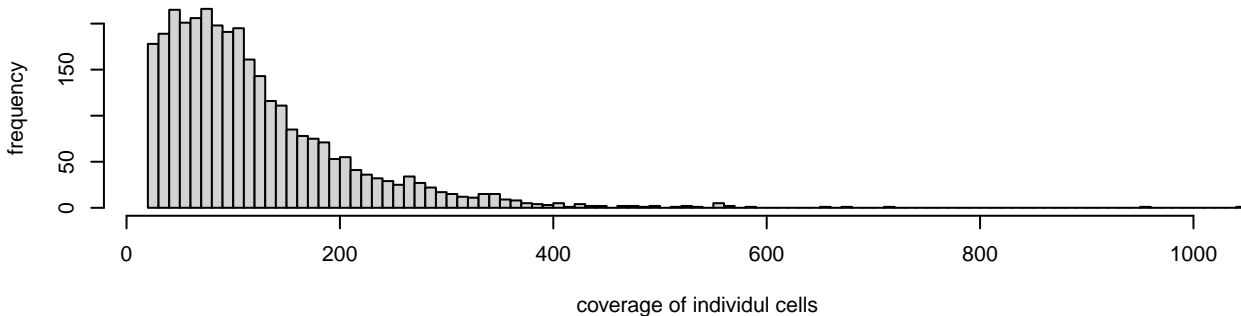
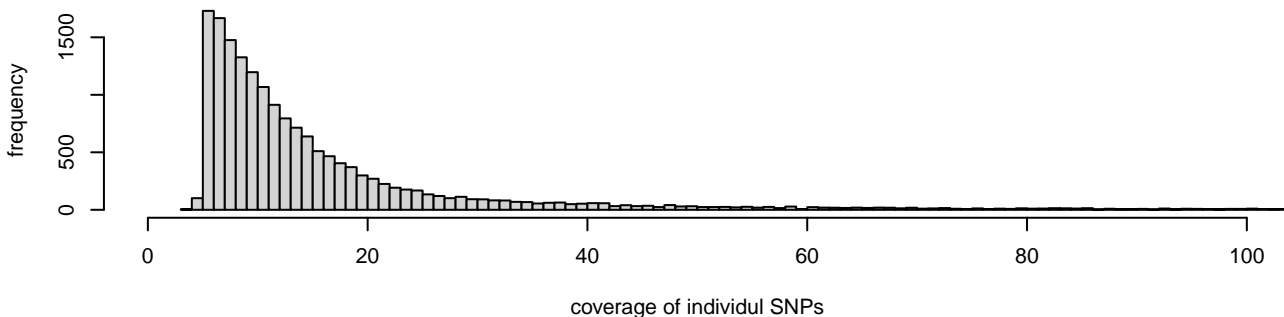


**Sample scATACseq chr20:26300000 coverage (3135 cells across 17556 SNPs)**



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**Histogram of VAF values**

