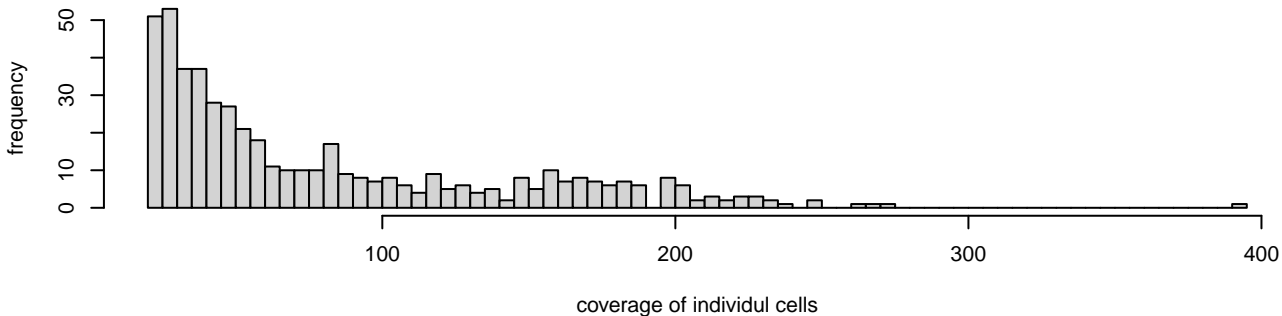
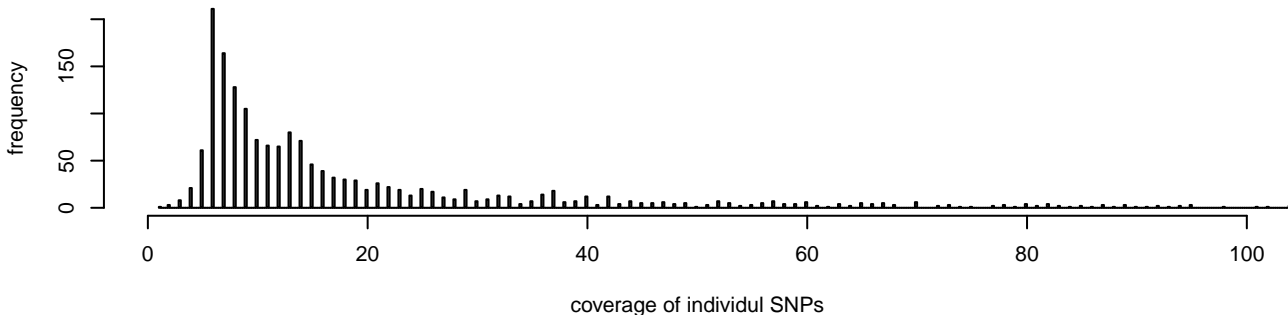


Sample scATACseq chr6:65612425 coverage (503 cells across 1760 SNPs)



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Histogram of VAF values

