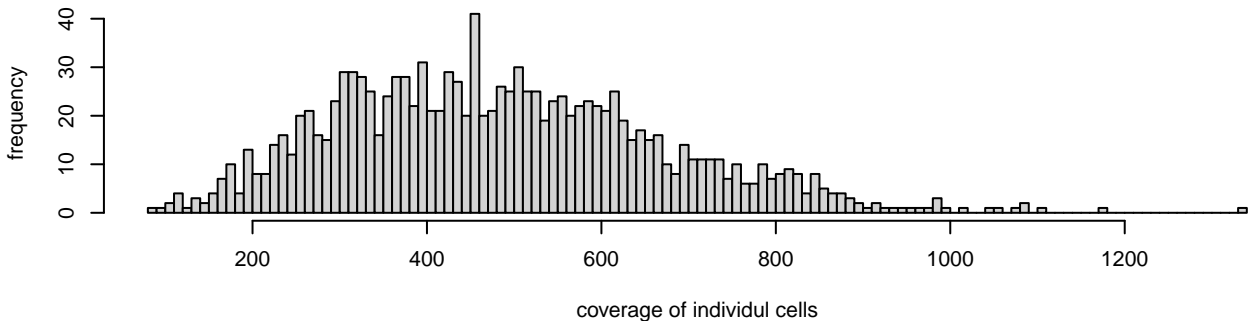
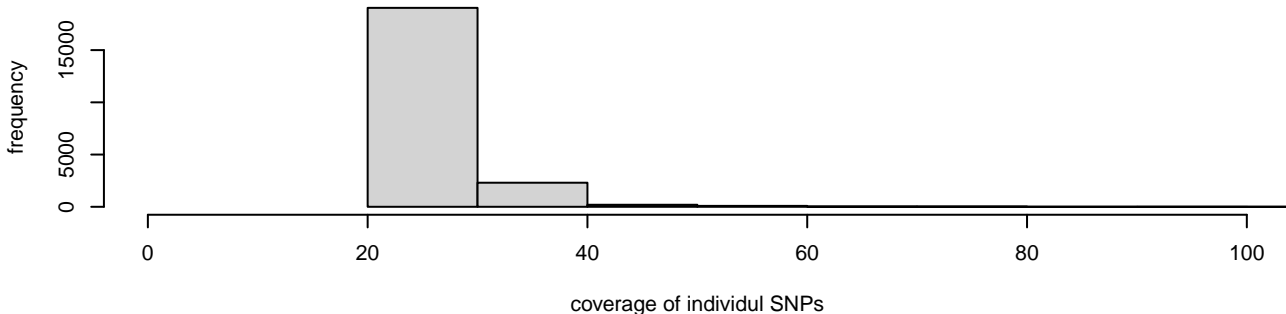


Sample scDNAseq chr10:82420000 coverage (1276 cells across 21771 SNPs)



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Histogram of VAF values

