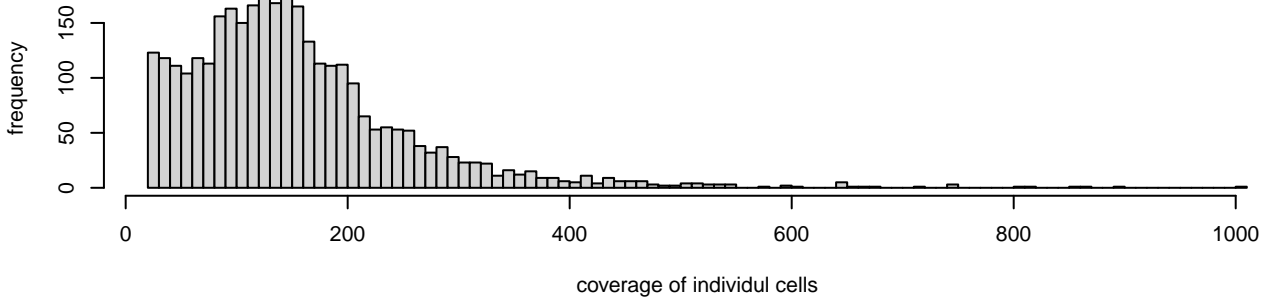
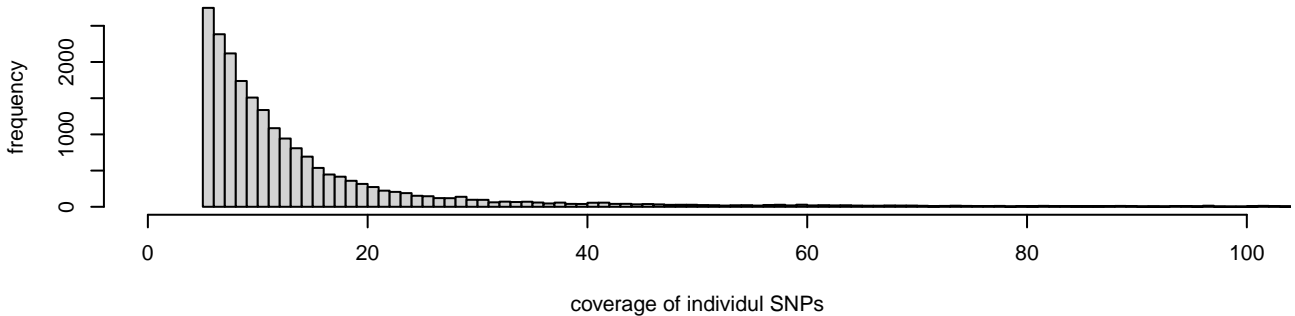


SNU601 scATACseq chr3:163840000 coverage (3225 cells across 21539 SNPs)



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Histogram of VAF values

