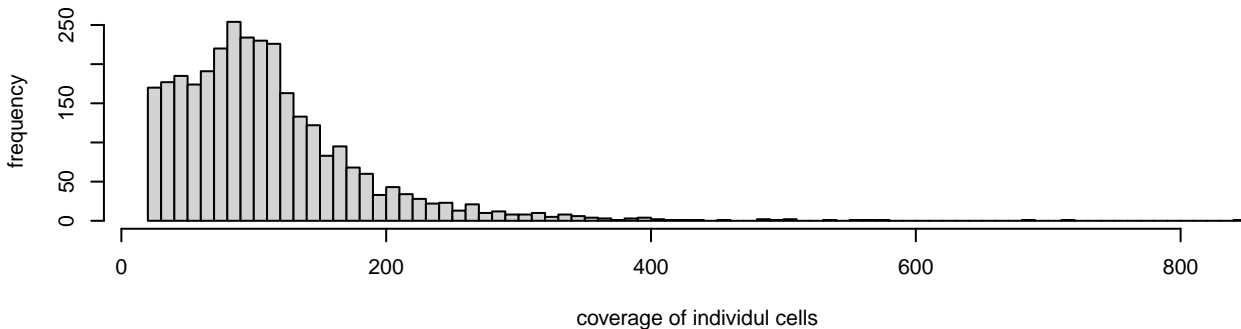
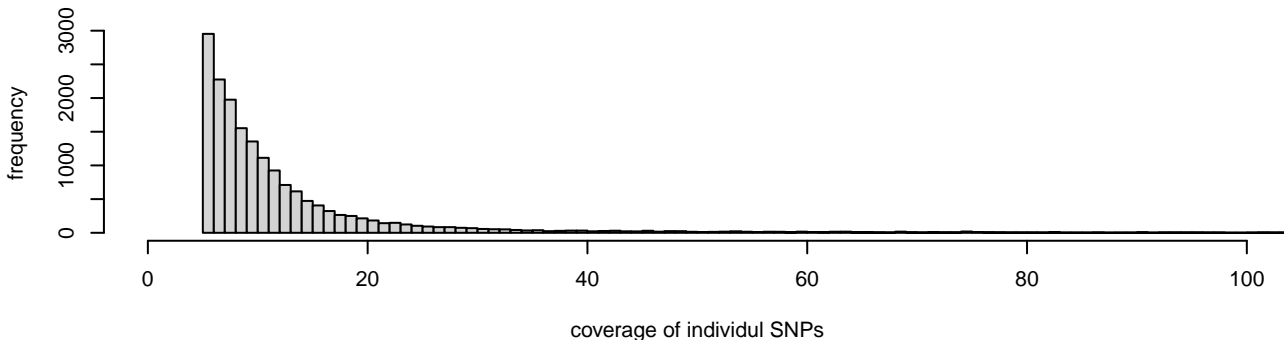


SNU601 scATACseq chr1:218360000 coverage (3102 cells across 18066 SNPs)



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Histogram of VAF values

