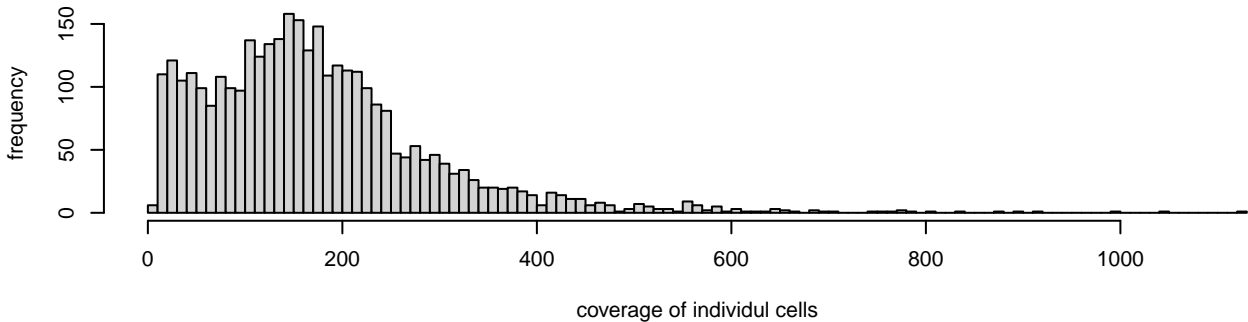
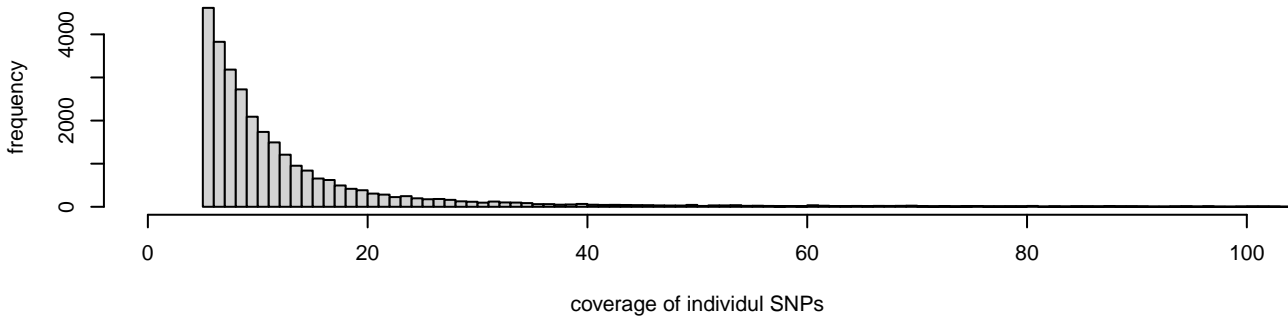


SNU601 scATACseq chr7:0 coverage (3405 cells across 30000 SNPs)



SNU601 scATACseq chr7:0 coverage (30000 SNPs across 3405 cells)



Histogram of VAF values

