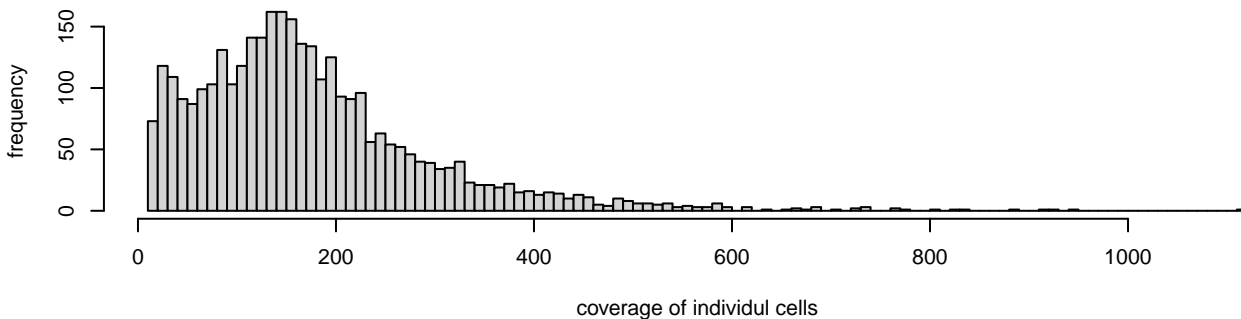
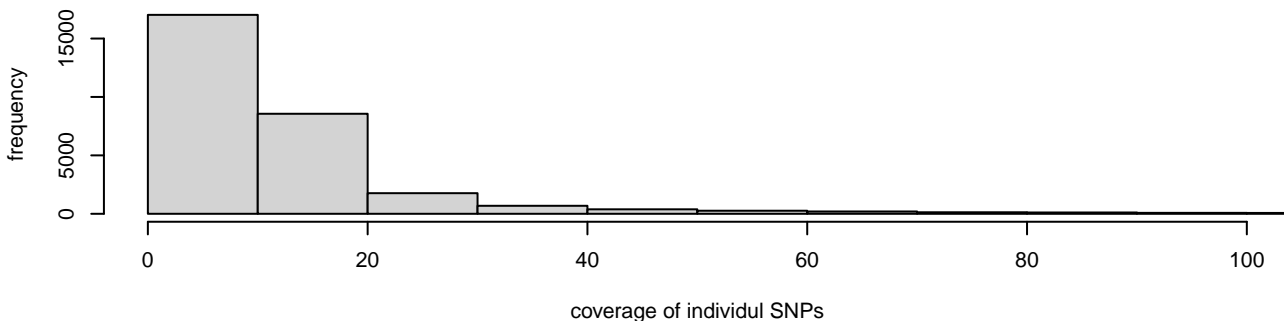


SNU601 scATACseq chr11:45940000 coverage (3348 cells across 30000 SNPs)



SNU601 scATACseq chr11:45940000 coverage (30000 SNPs across 3348 cells)



Histogram of VAF values

