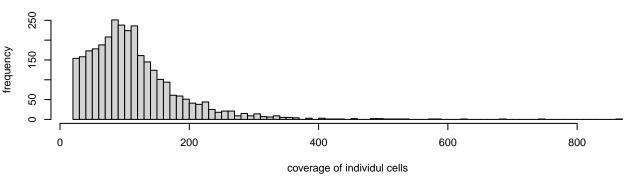
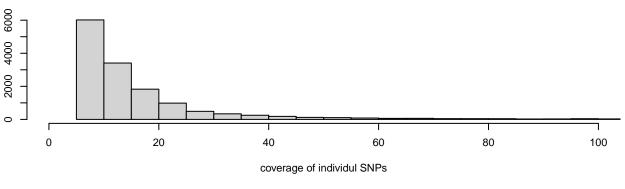
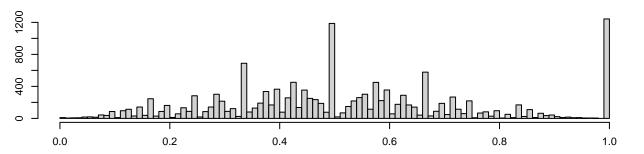
SNU601 scATACseq chr20:44820000 coverage (3120 cells across 14693 SNPs)



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af_all_sub

Histogram of VAF values