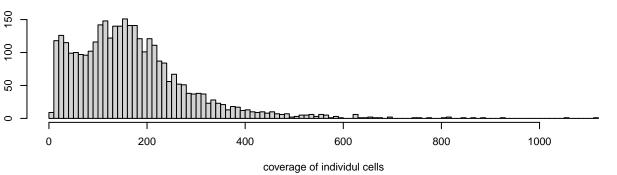
Sample scATACseq chr7:0 coverage (3405 cells across 30000 SNPs)

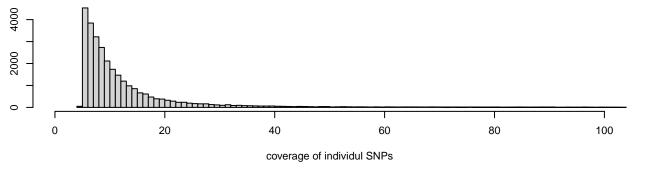


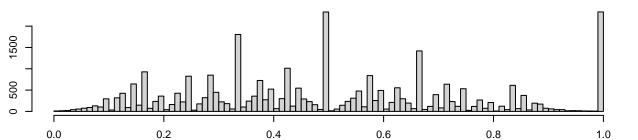
frequency

frequency

Frequency

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





af_all_sub

Histogram of VAF values