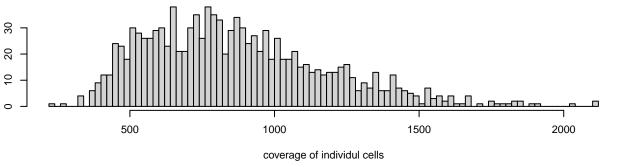
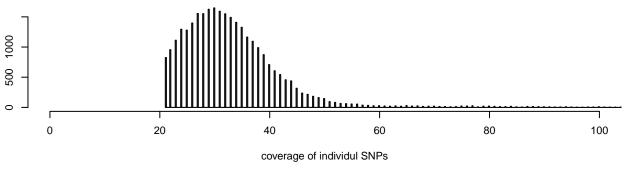
Sample scDNAseq chr3:163840000 coverage (1155 cells across 30000 SNPs)



frequency

Frequency

Sample scDNAseq chr3:163840000 coverage (30000 SNPs across 1155 cells)



Histogram of VAF values

