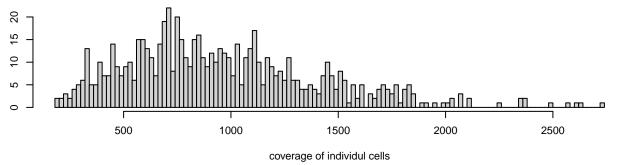
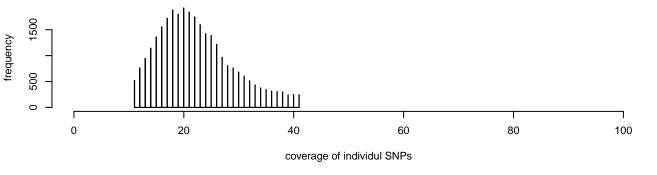
P5931 scDNAseq chr20 coverage (696 cells across 30000 SNPs)



frequency

P5931 scDNAseq chr20 coverage (30000 SNPs across 696 cells)



Histogram of VAF values

