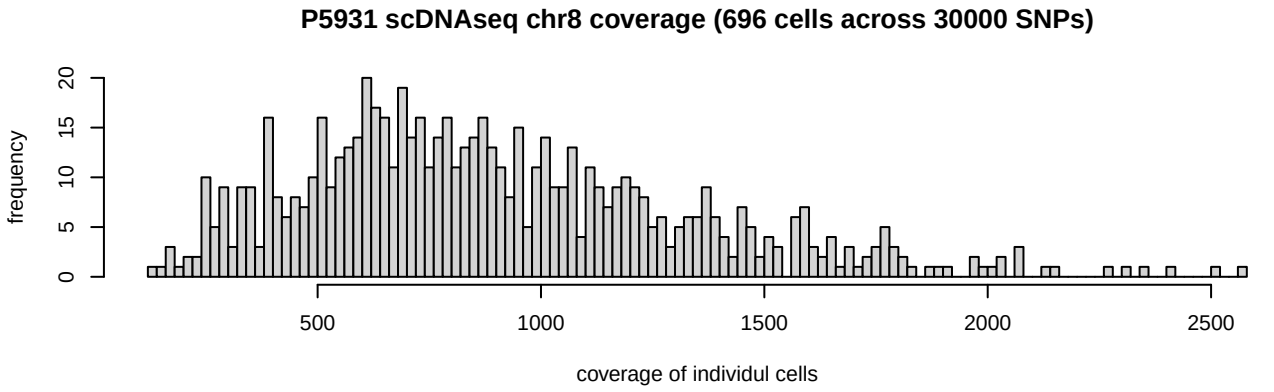
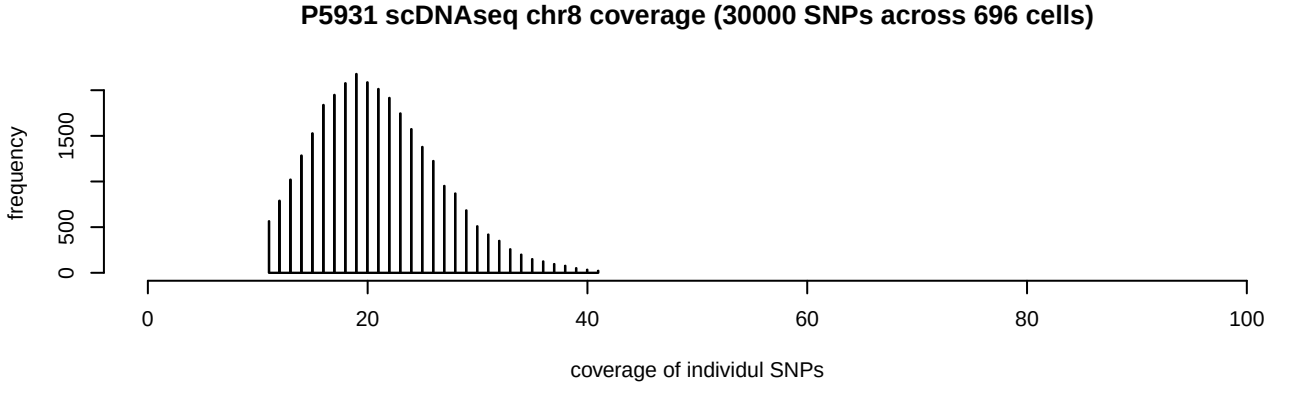
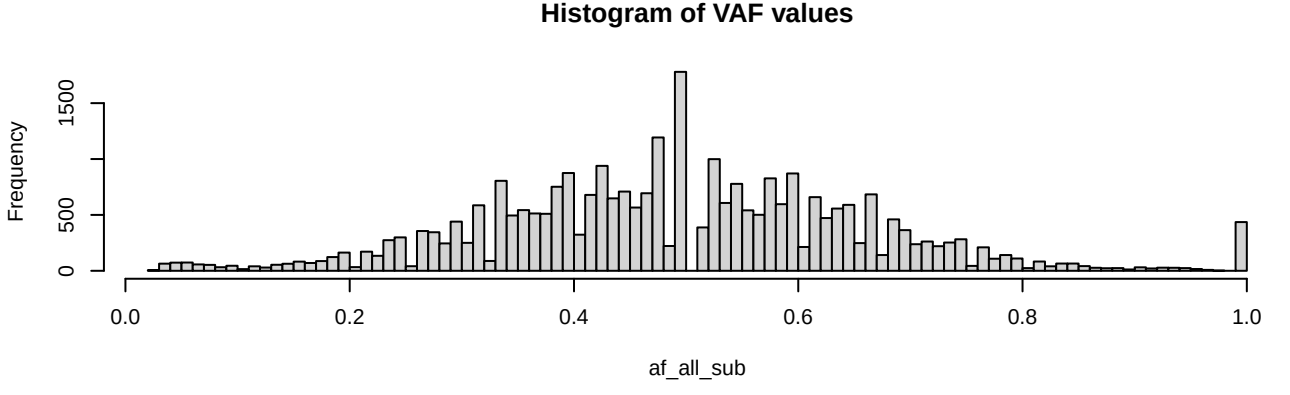

P5931 scDNAseq chr8 coverage (696 cells across 30000 SNPs)
frequency
0
5
10
15
20
500
1000
1500
2000
2500
coverage of individul cells


P5931 scDNAseq chr8 coverage (30000 SNPs across 696 cells)
frequency
0
500
1500
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
500
1500
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub