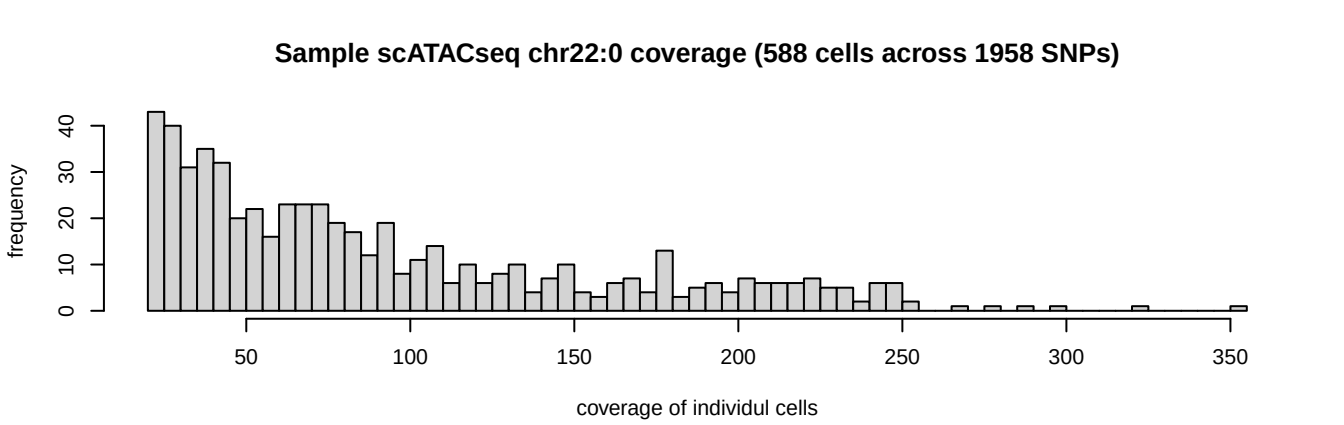
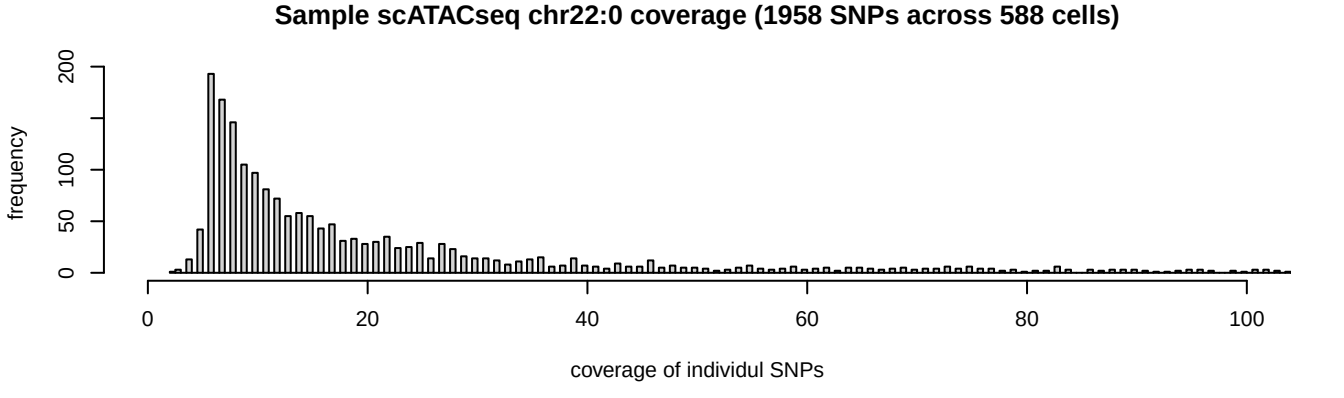
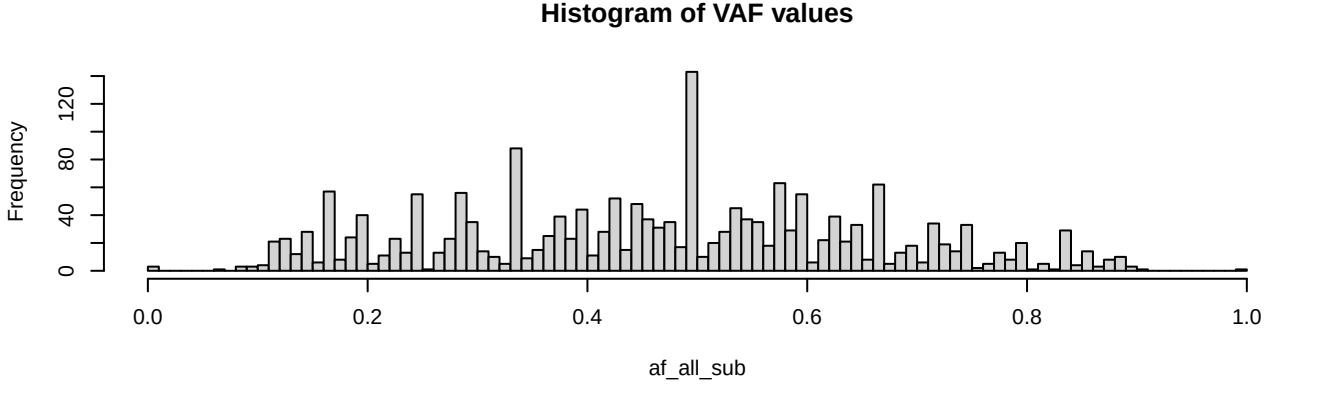

Sample scATACseq chr22:0 coverage (588 cells across 1958 SNPs)
frequency
0
10
20
30
40
50
100
150
200
250
300
350
coverage of individul cells


Sample scATACseq chr22:0 coverage (1958 SNPs across 588 cells)
frequency
0
50
100
200
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
40
80
120
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub