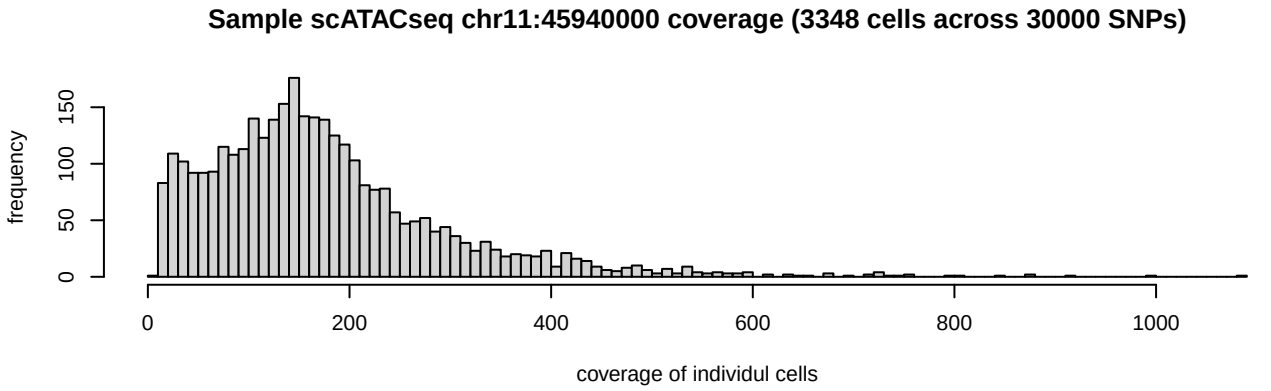
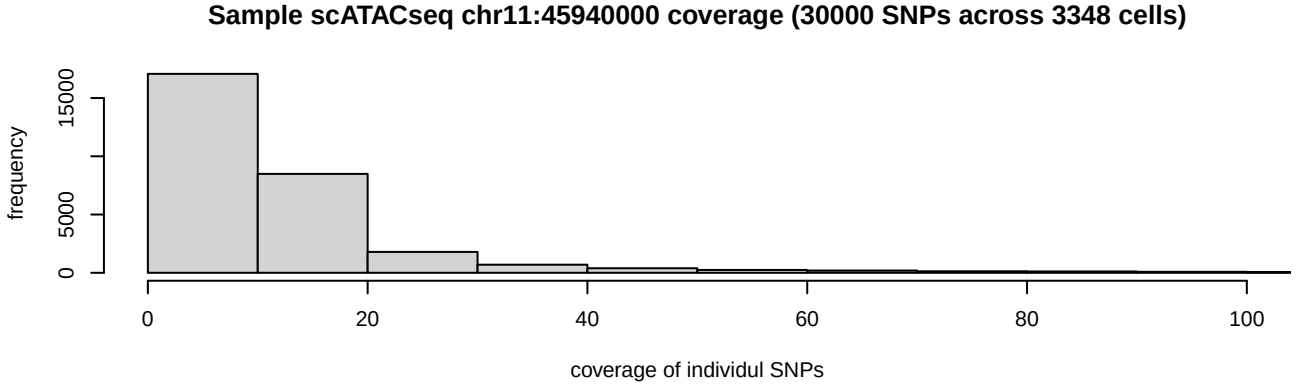
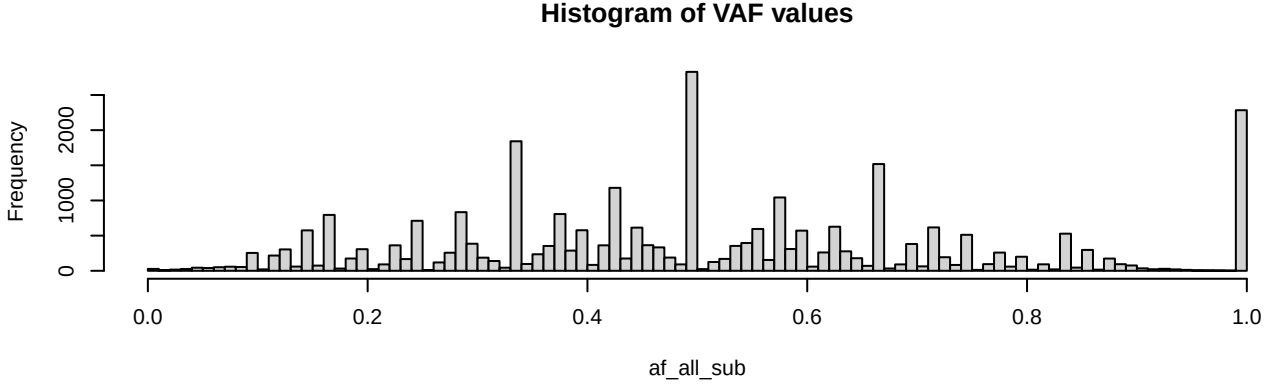

Sample scATACseq chr11:45940000 coverage (3348 cells across 30000 SNPs)
frequency
0
50
100
150
0
200
400
600
800
1000
coverage of individul cells


Sample scATACseq chr11:45940000 coverage (30000 SNPs across 3348 cells)
frequency
0
5000
15000
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
1000
2000
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub