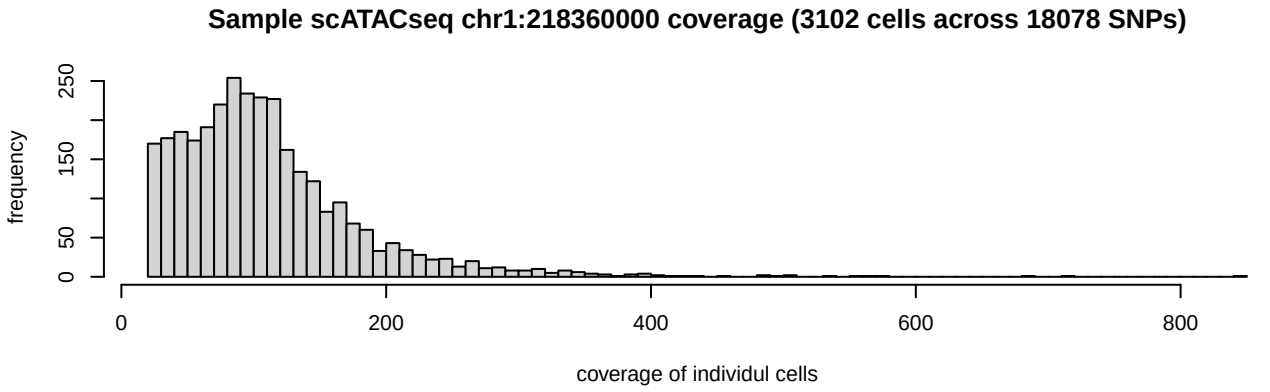
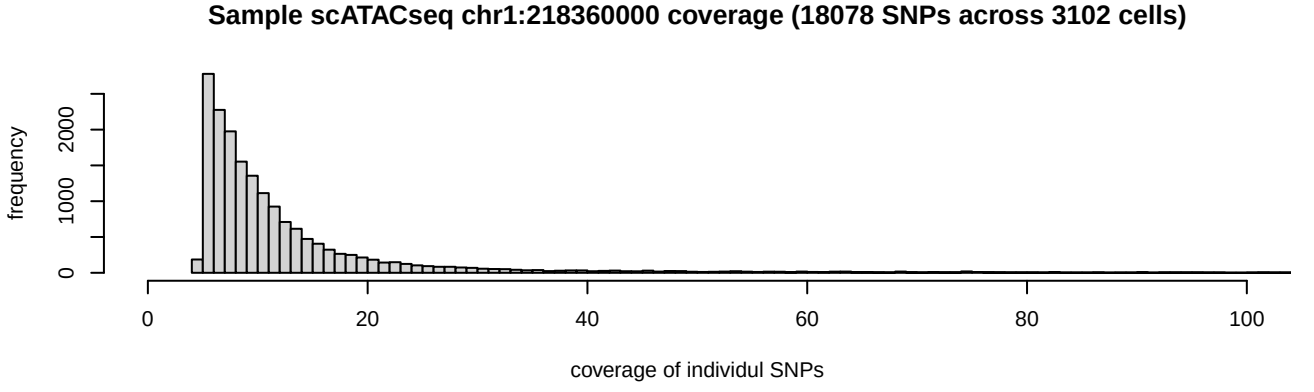
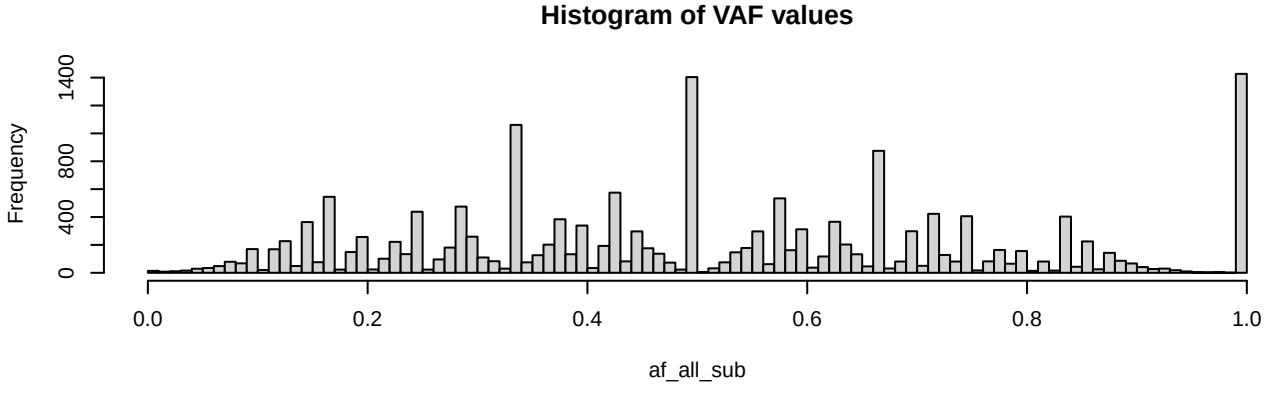

Sample scATACseq chr1:218360000 coverage (3102 cells across 18078 SNPs)
frequency
0
50
150
250
0
200
400
600
800
coverage of individul cells


Sample scATACseq chr1:218360000 coverage (18078 SNPs across 3102 cells)
frequency
0
1000
2000
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
400
800
1400
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub