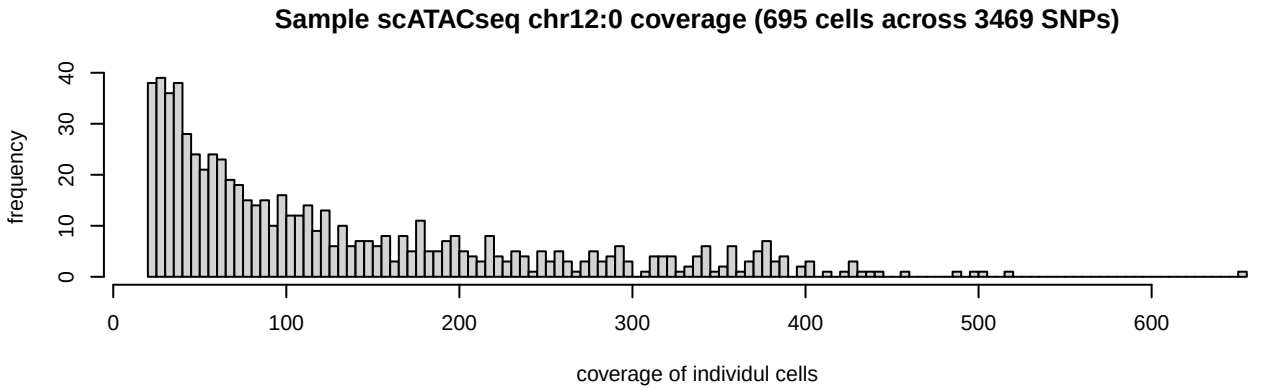
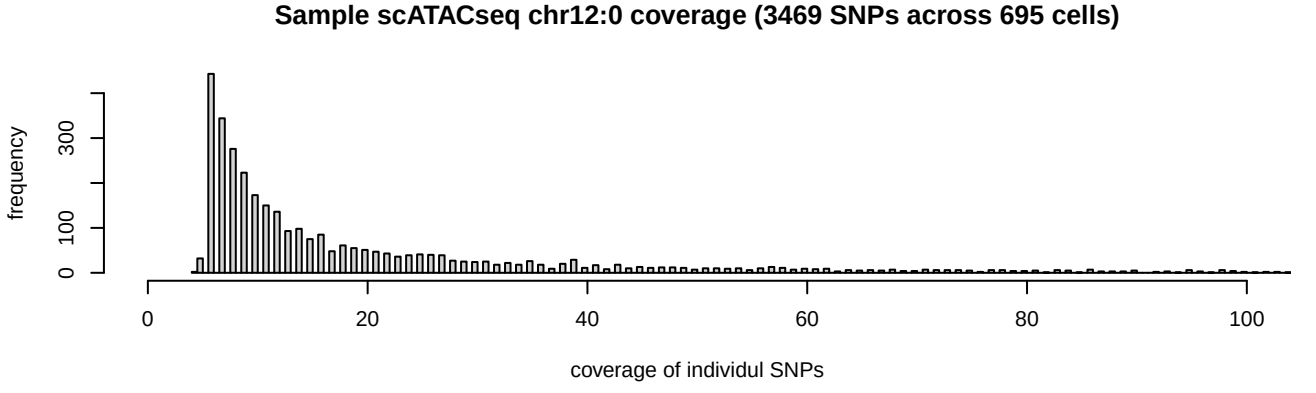
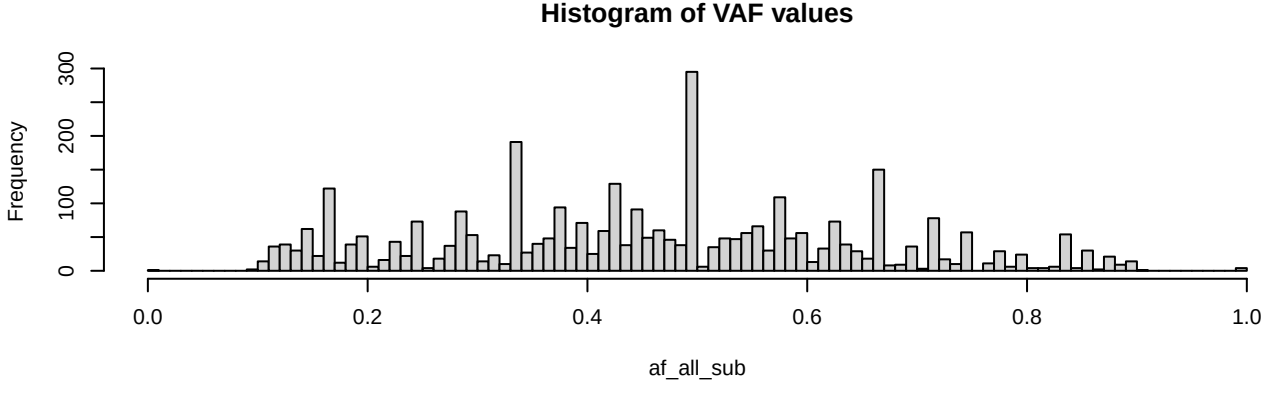

Sample scATACseq chr12:0 coverage (695 cells across 3469 SNPs)
frequency
0
10
20
30
40
0
100
200
300
400
500
600
coverage of individul cells


Sample scATACseq chr12:0 coverage (3469 SNPs across 695 cells)
frequency
0
100
300
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
100
200
300
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub