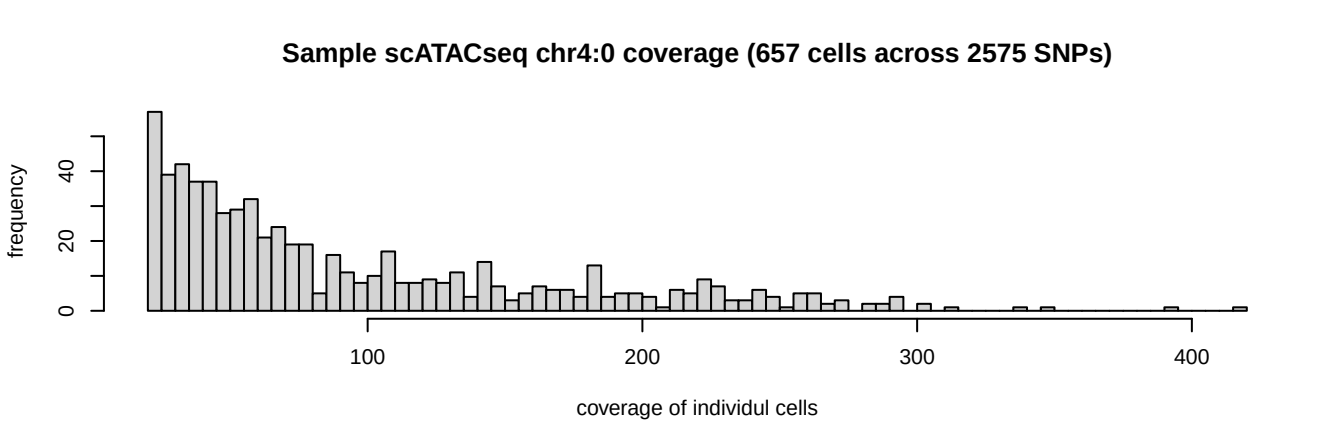
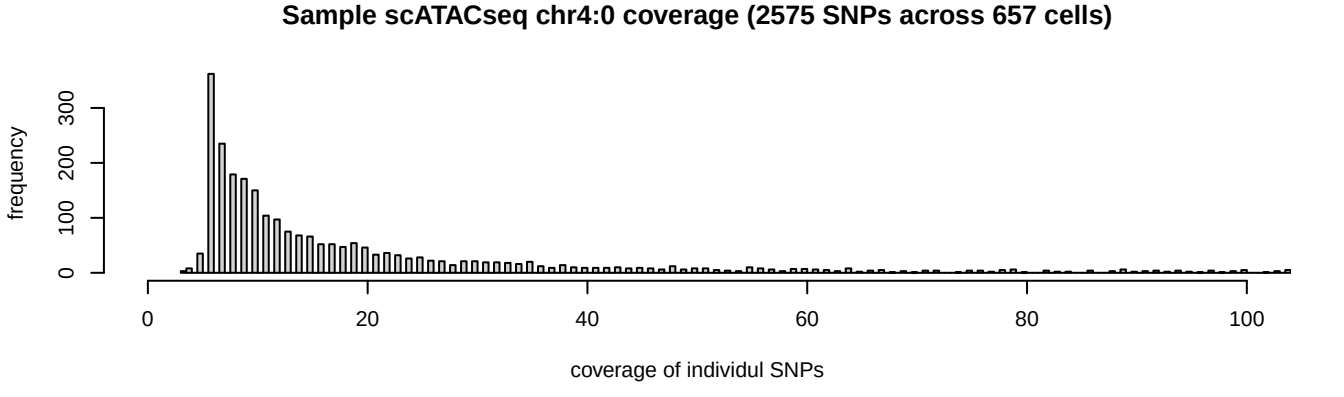
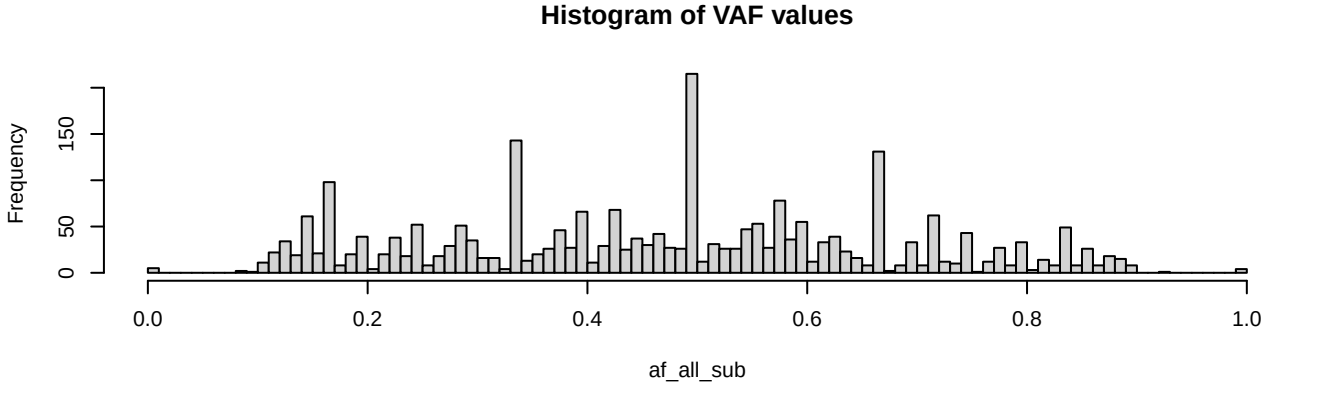

Sample scATACseq chr4:0 coverage (657 cells across 2575 SNPs)
frequency
0
20
40
100
200
300
400
coverage of individul cells


Sample scATACseq chr4:0 coverage (2575 SNPs across 657 cells)
frequency
0
100
200
300
0
20
40
60
80
100
coverage of individul SNPs


Histogram of VAF values
Frequency
0
50
150
0.0
0.2
0.4
0.6
0.8
1.0
af_all_sub