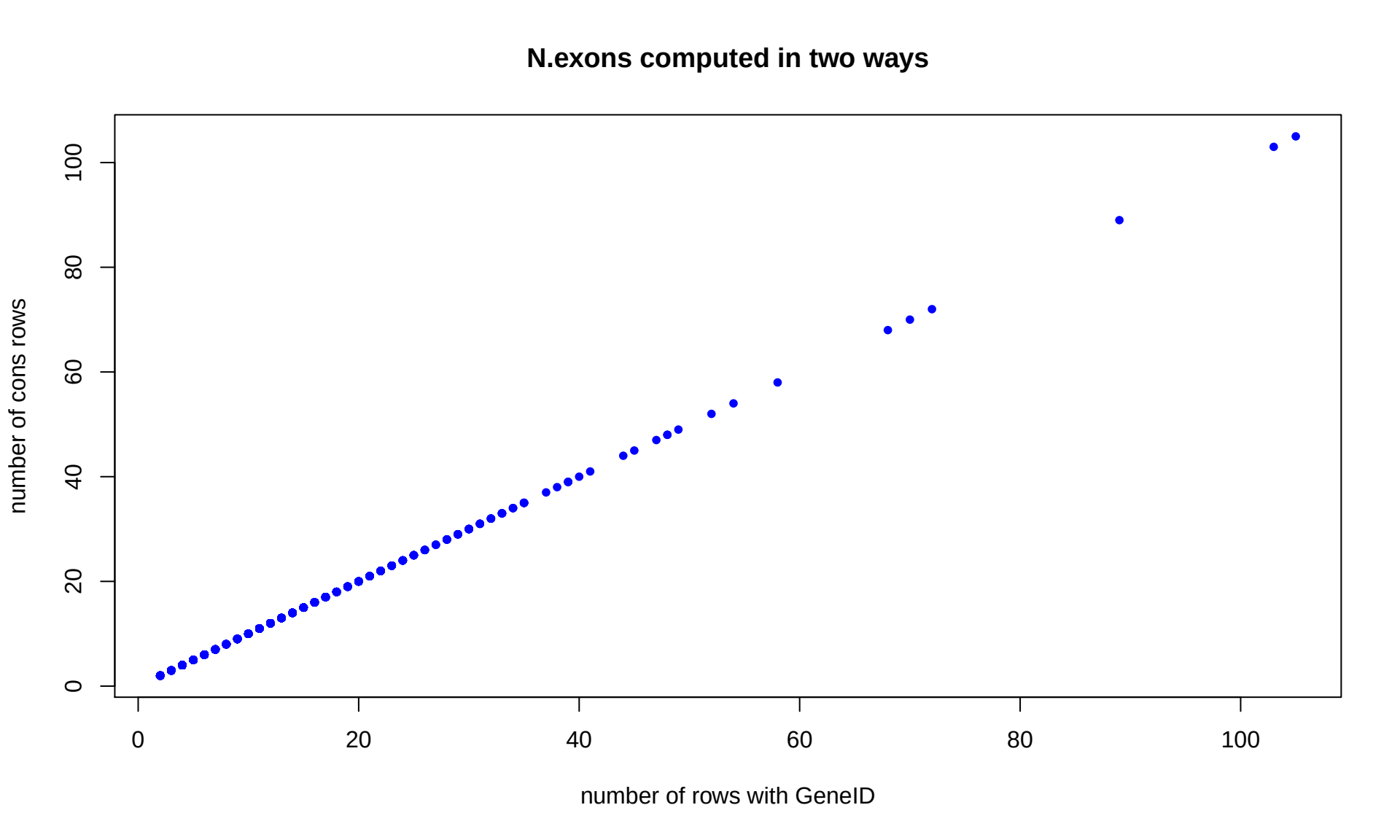
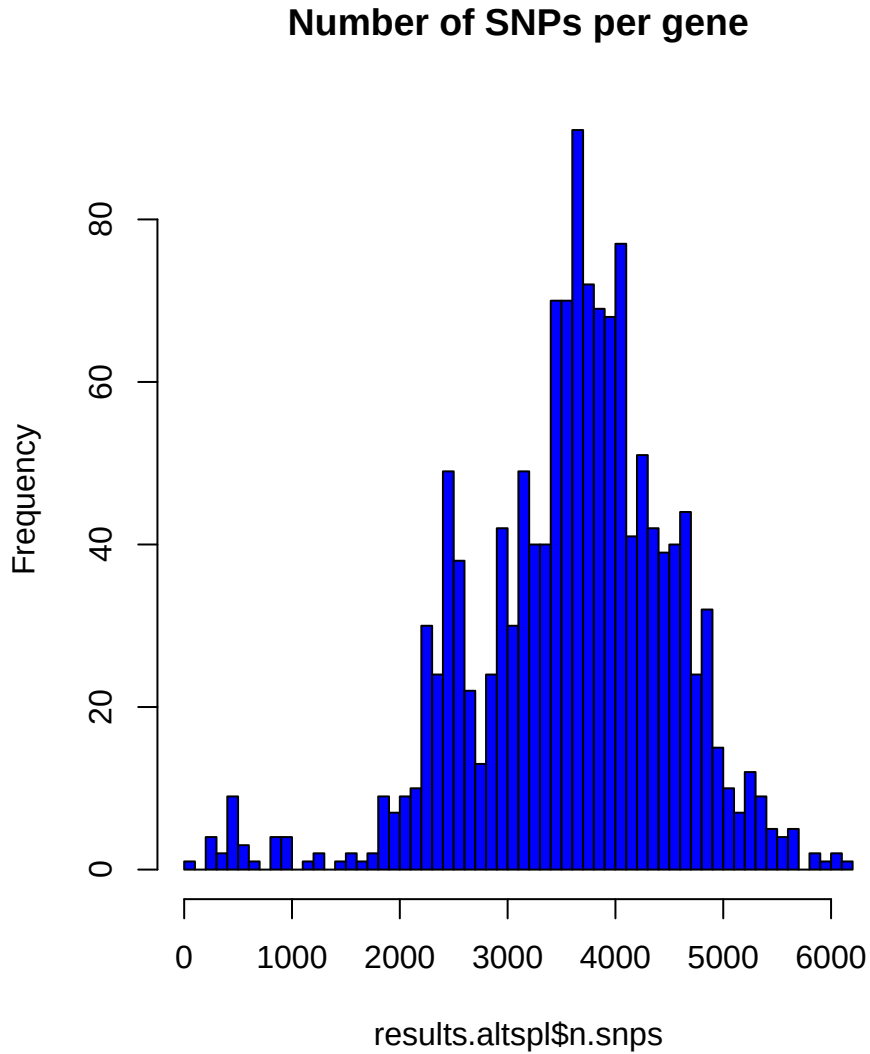
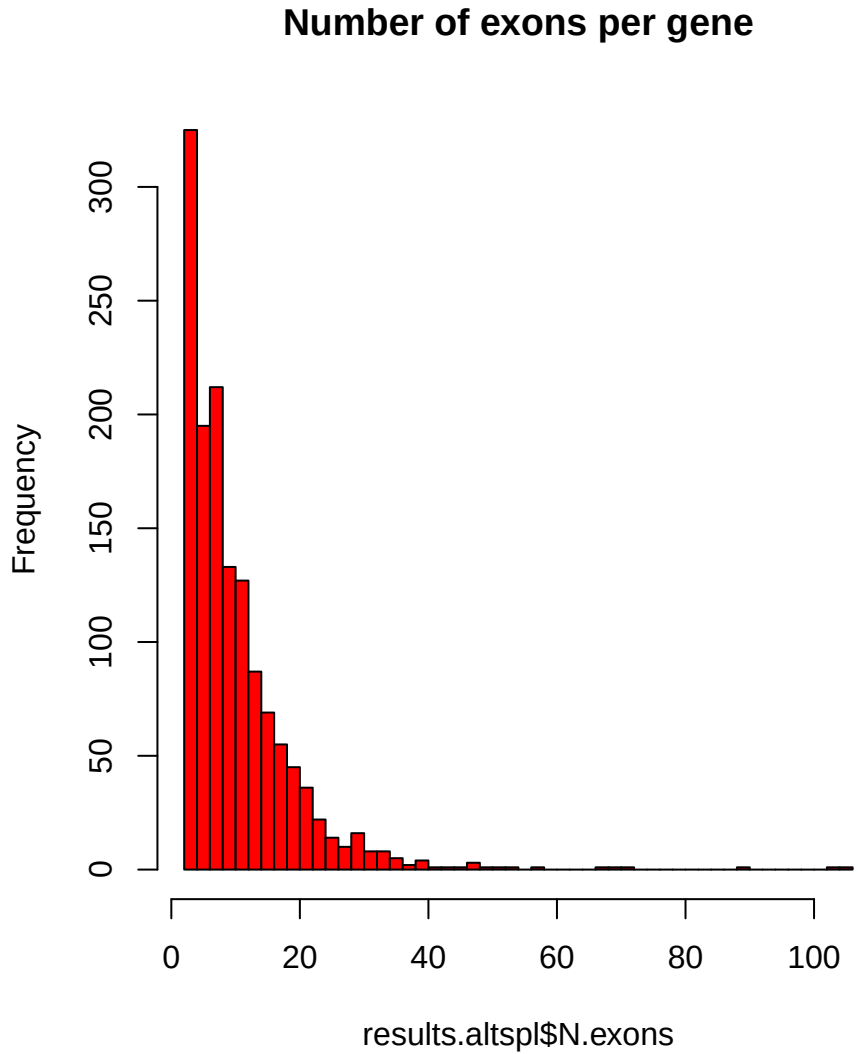
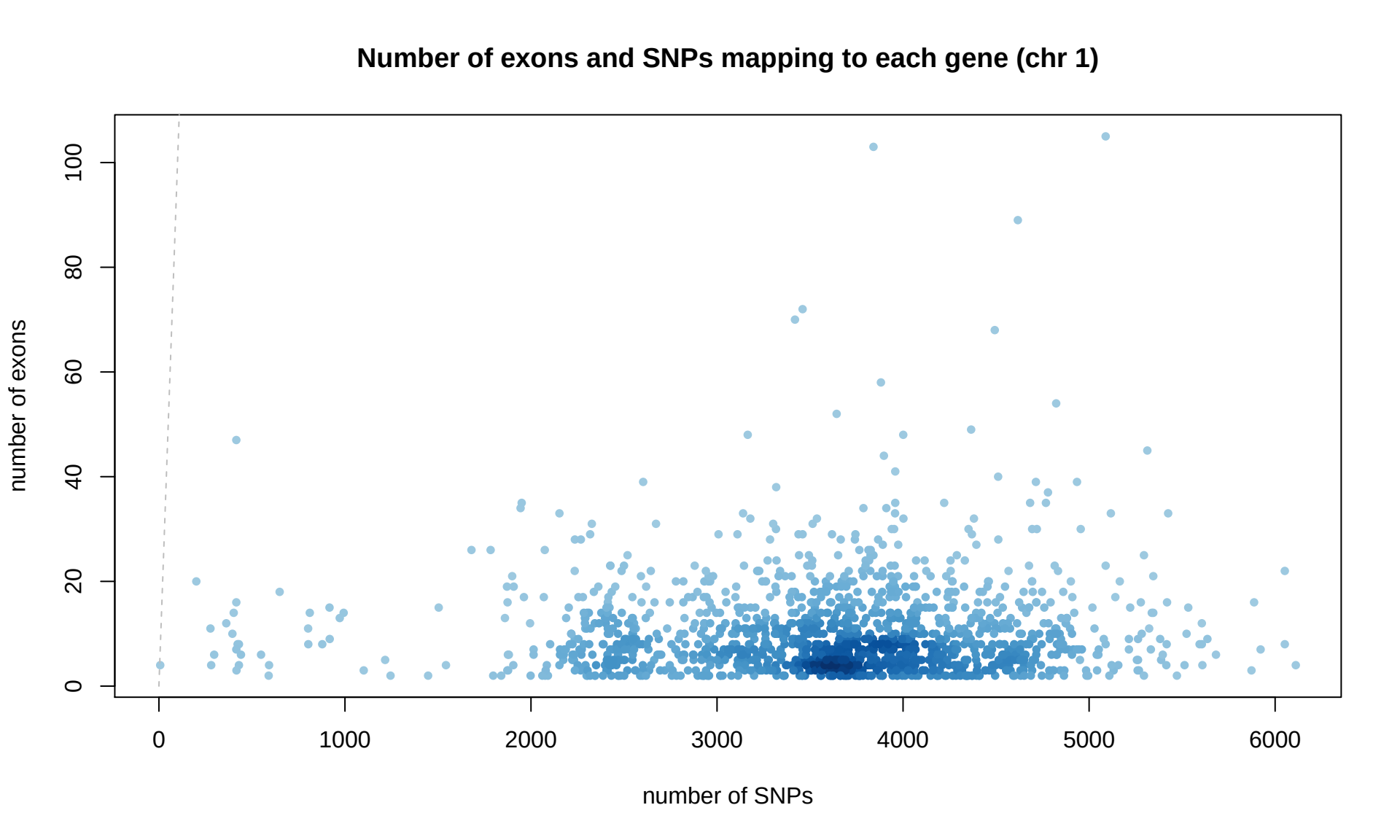
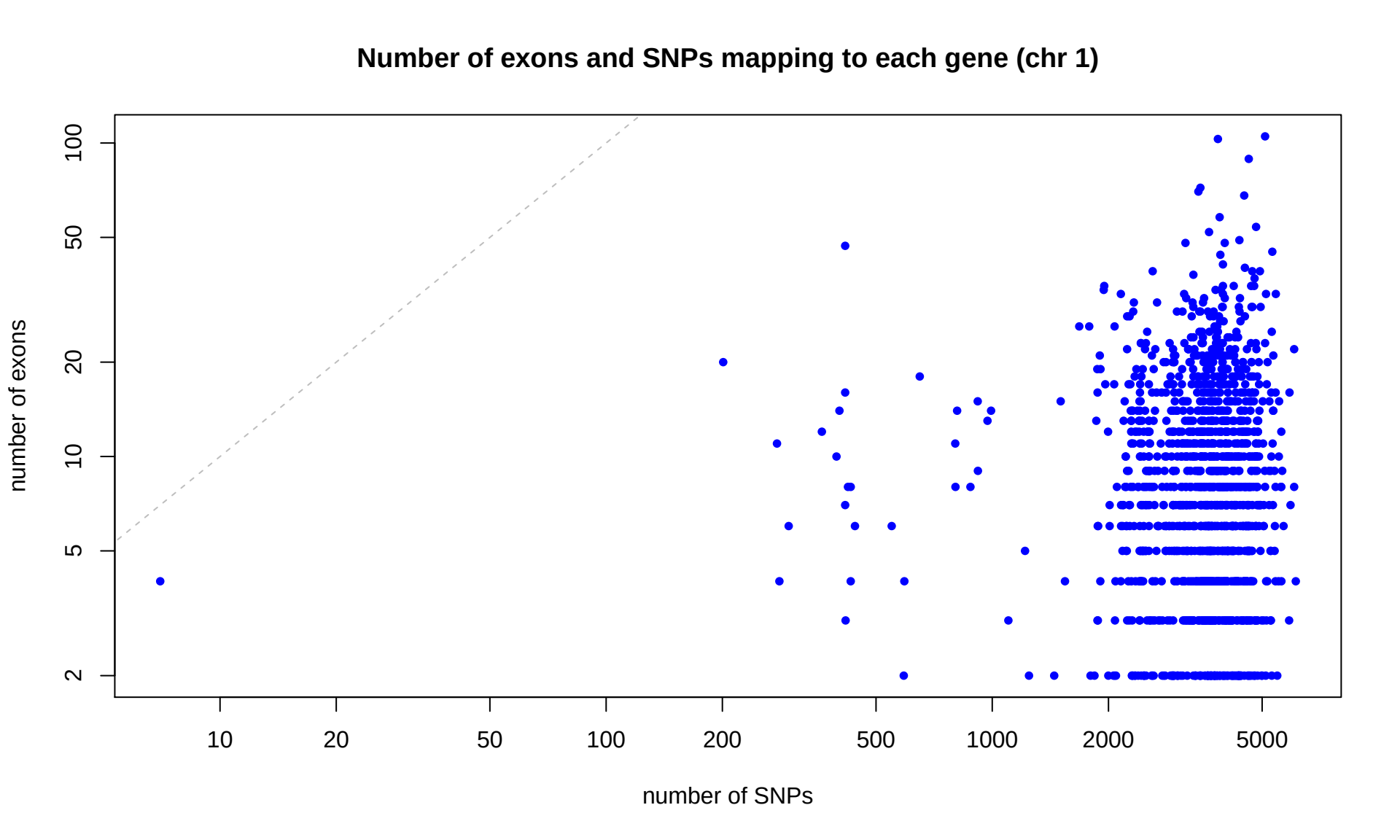

N.exons computed in two ways
number of cons rows
0
20
40
60
80
100
number of rows with GeneID
0
20
40
60
80
100


Number of SNPs per gene
Frequency
0
20
40
60
80
0
1000
2000
3000
4000
5000
6000
results.altspl$n.snps


Number of exons per gene
Frequency
0
50
100
150
200
250
300
0
20
40
60
80
100
results.altspl$N.exons


Number of exons and SNPs mapping to each gene (chr 1)
number of exons
0
20
40
60
80
100
0
1000
2000
3000
4000
5000
6000
number of SNPs


Number of exons and SNPs mapping to each gene (chr 1)
number of exons
2
5
10
20
50
100
number of SNPs
10
20
50
100
200
500
1000
2000
5000