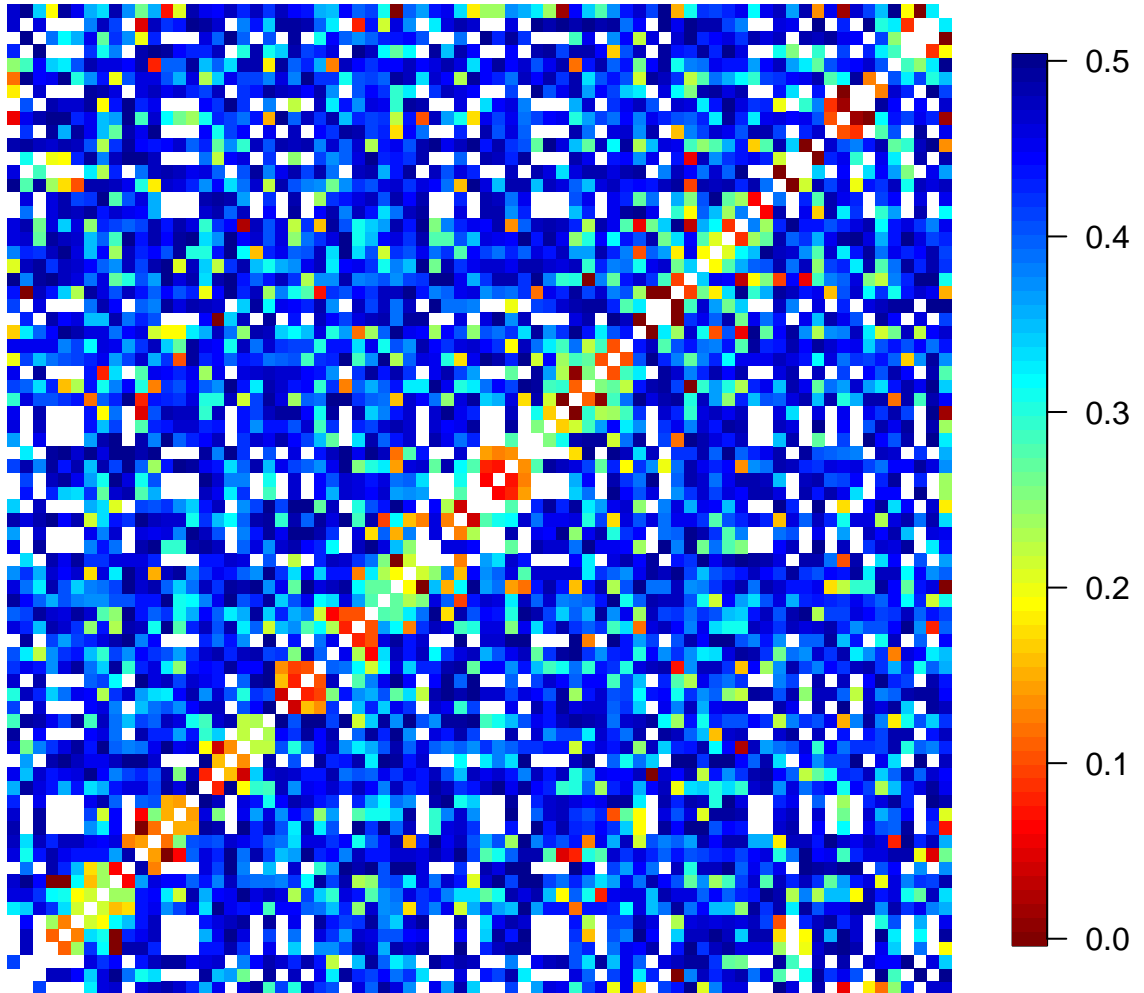
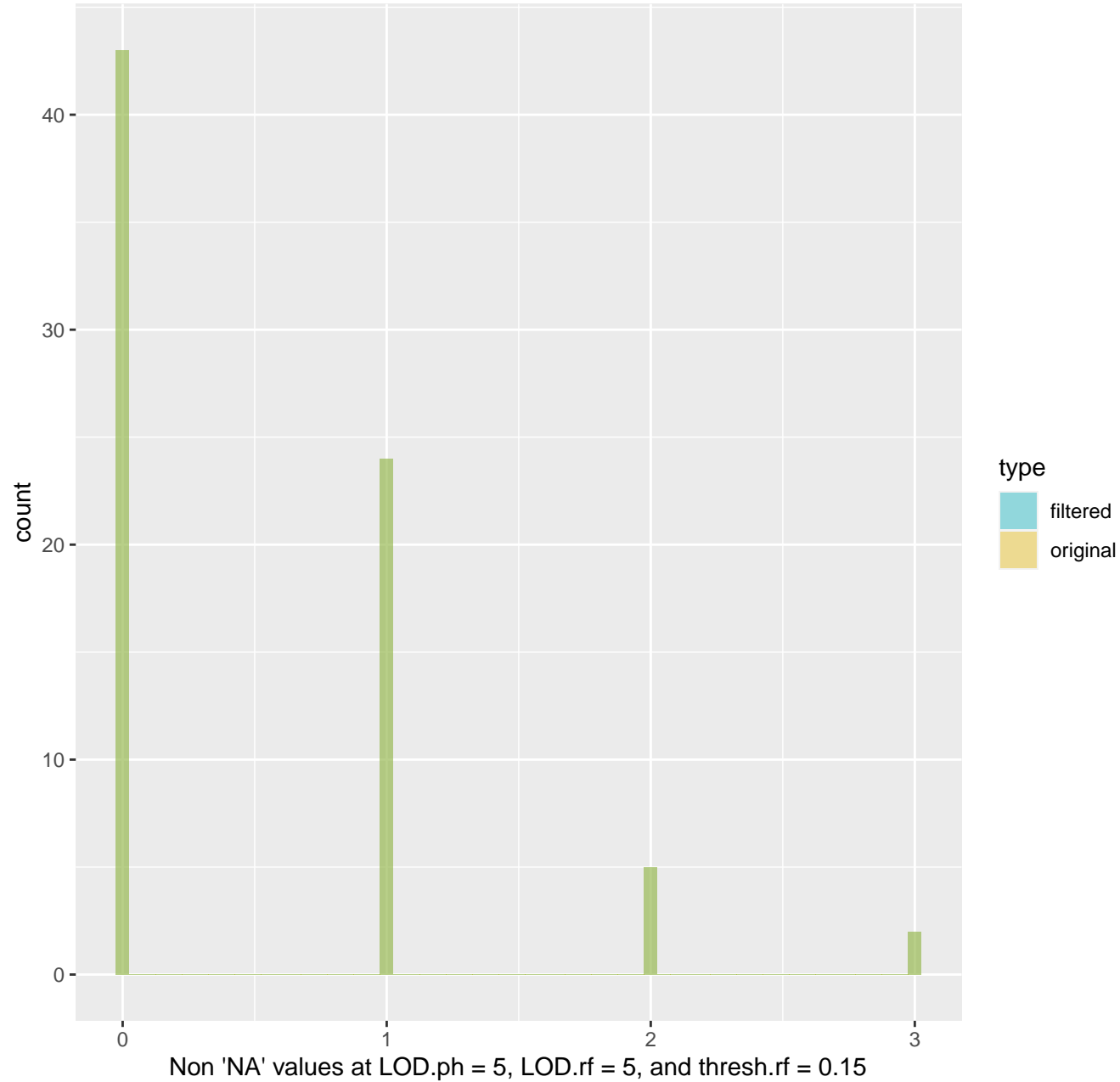


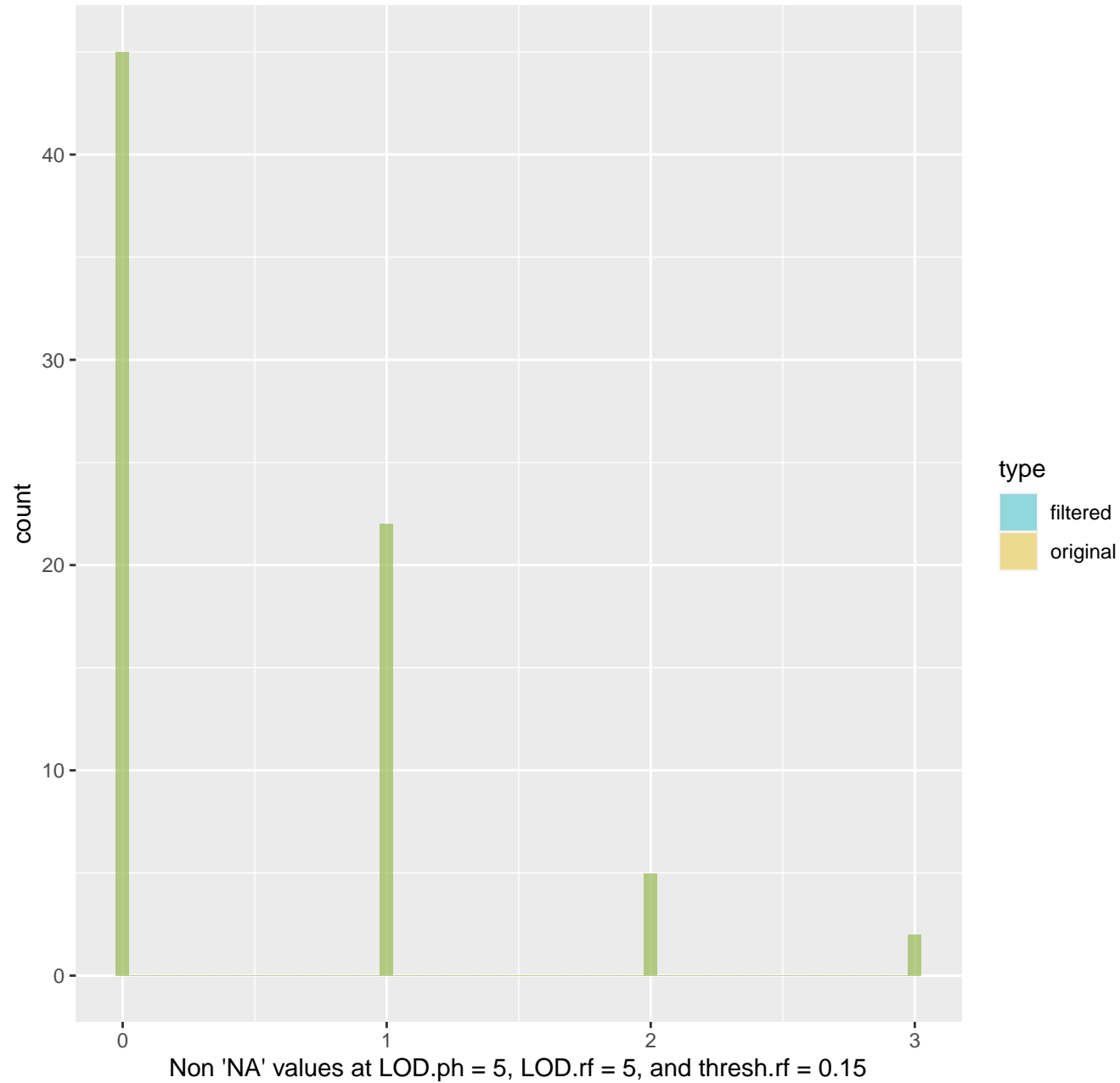
Recombination fraction matrix



Filtering probs: [0.05 : 1] – Non NA values by row in rf matrix – b width: 0.05

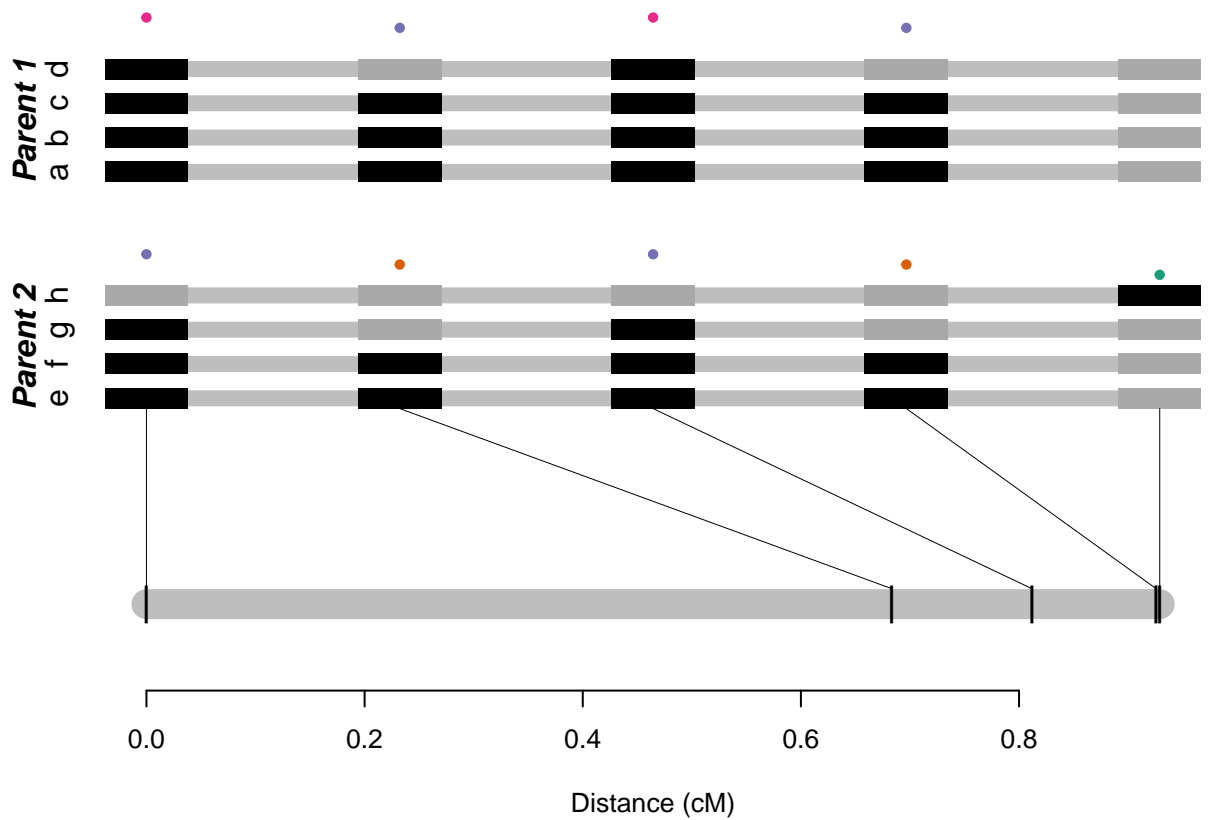


Filtering probs: [0.05 : 1] – Non NA values by row in rf matrix – b width: 0.05



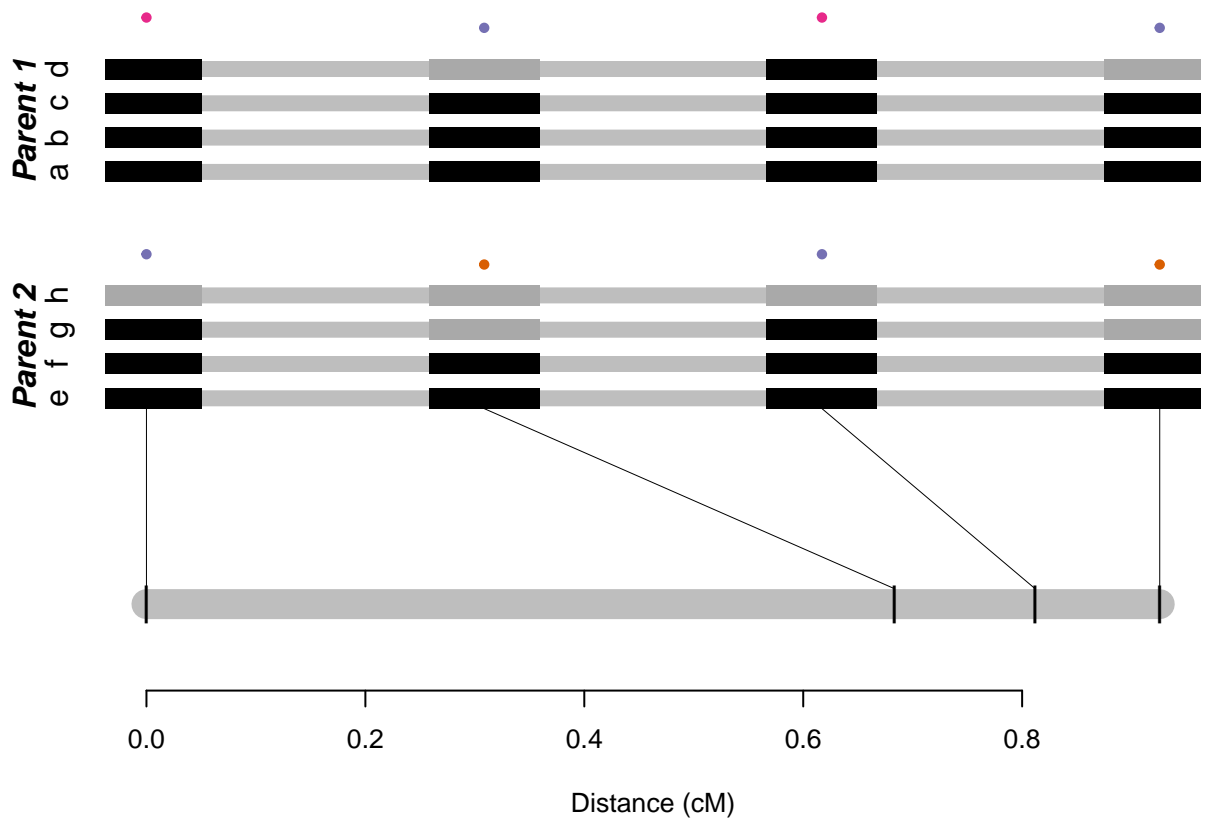
Variants
■ A ■ B

Doses
● 1 ● 3
● 2 ● 4



Variants
 ■ A ■ B

Doses
 ● 1 ● 3
 ● 2 ● 4



Linkage group

2

1



0.0

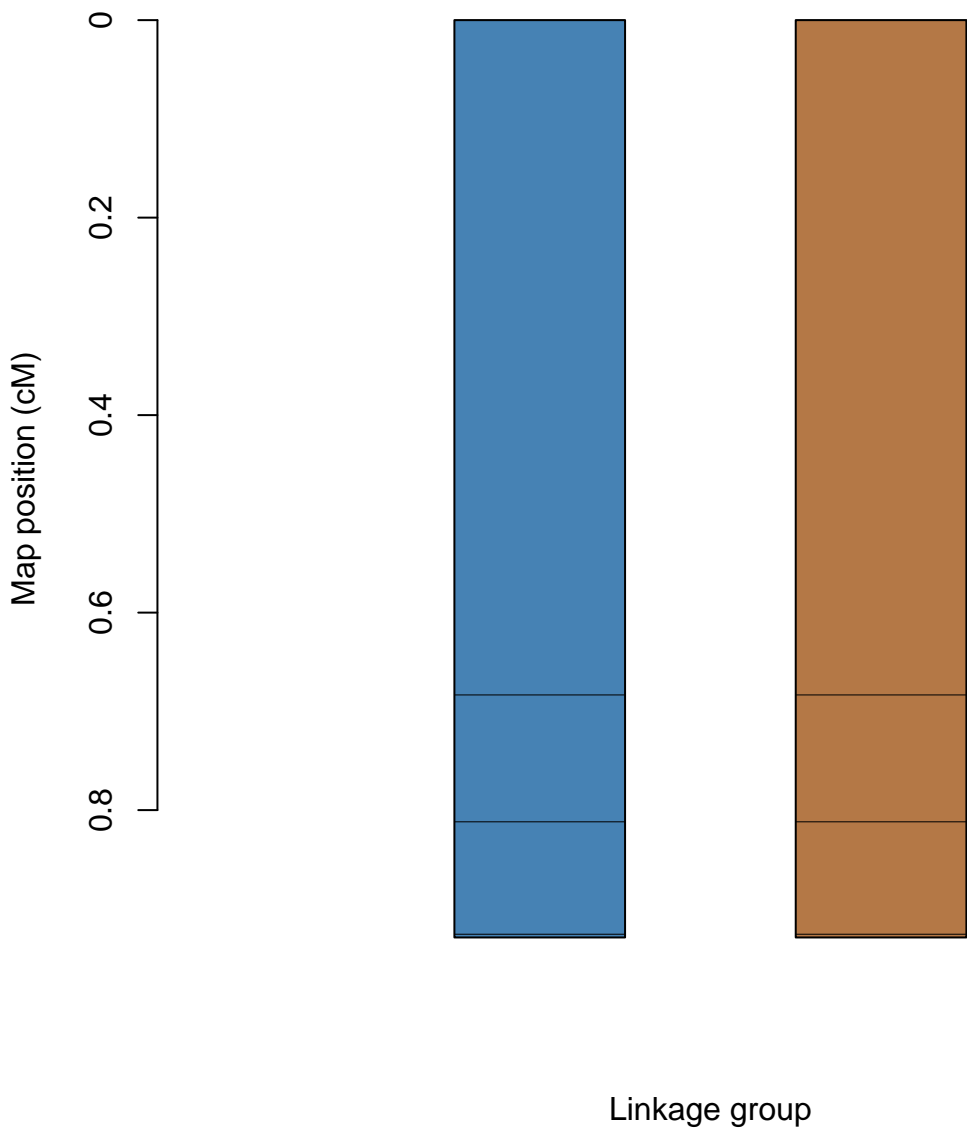
0.2

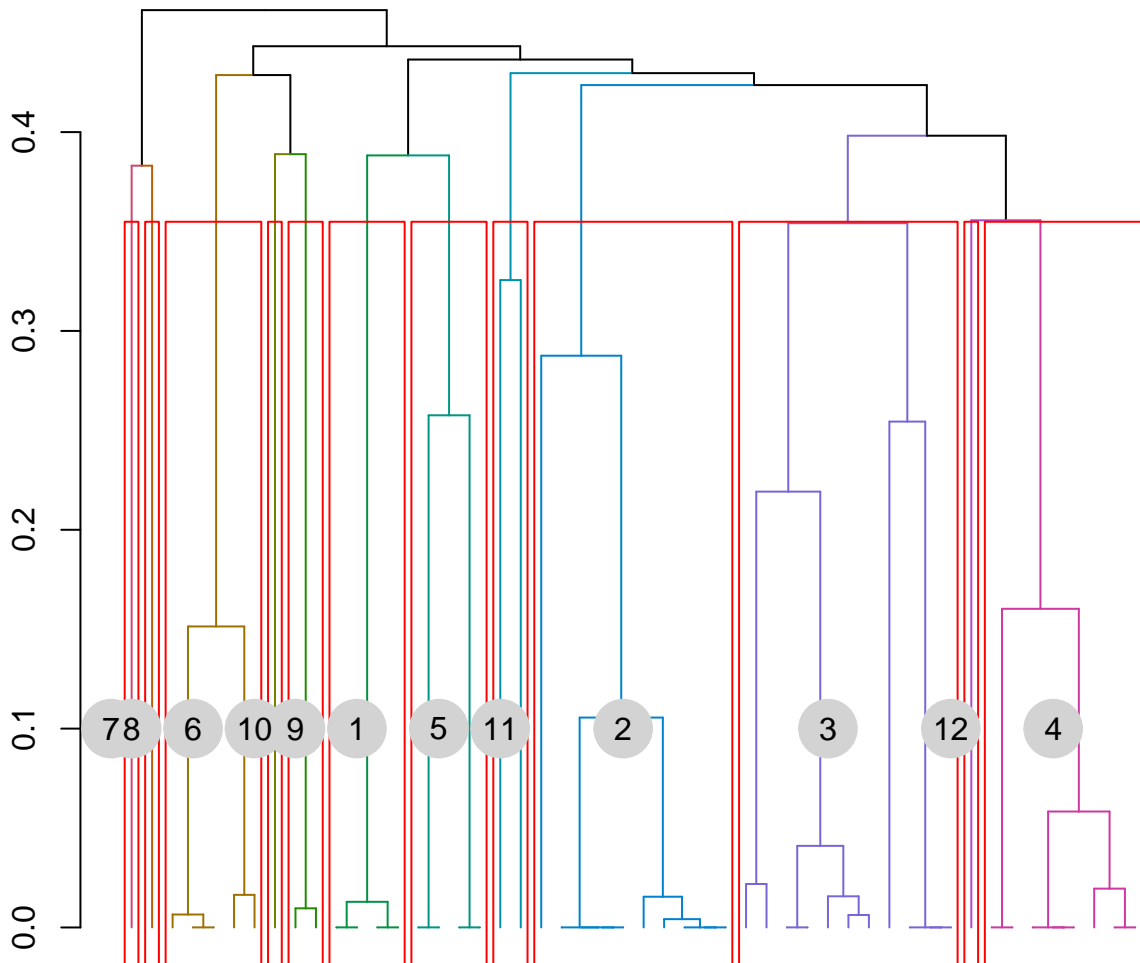
0.4

0.6

0.8

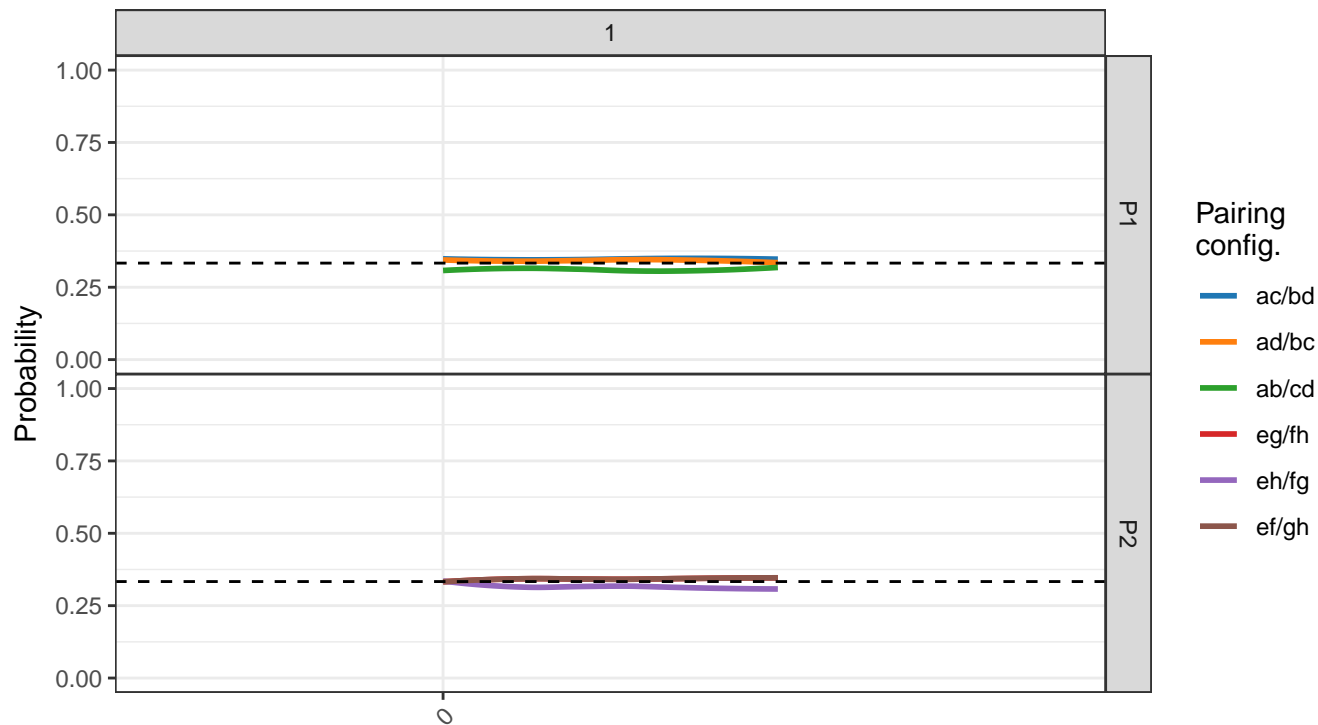
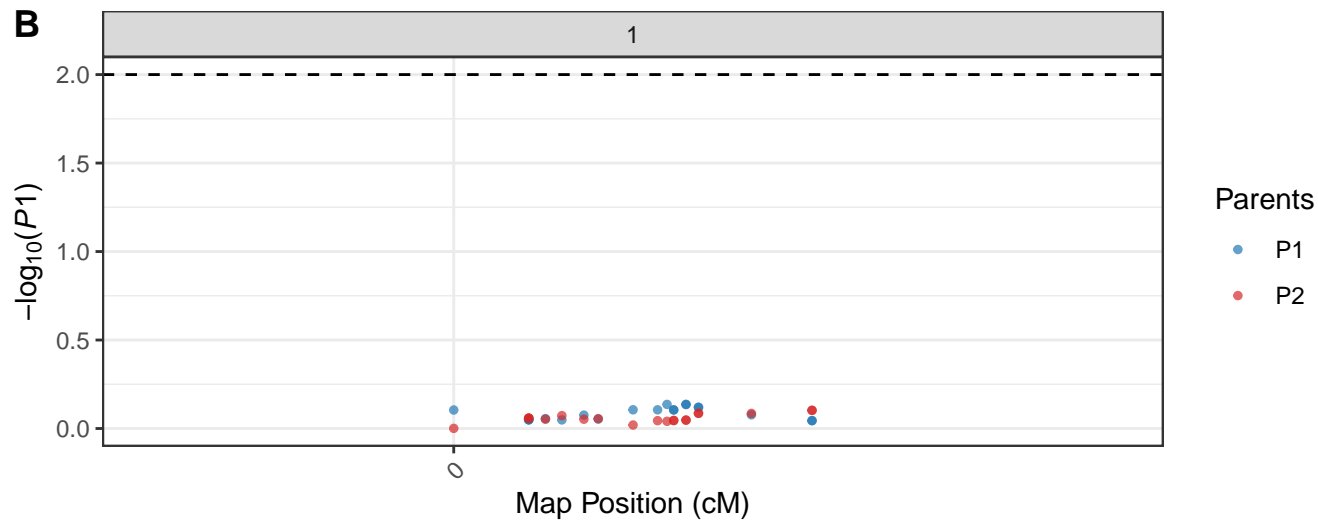
Map position (cM)





A

Linkage group

**B**

solcap_snp_c2_41437

marker #: 1

Dose in P1: 3

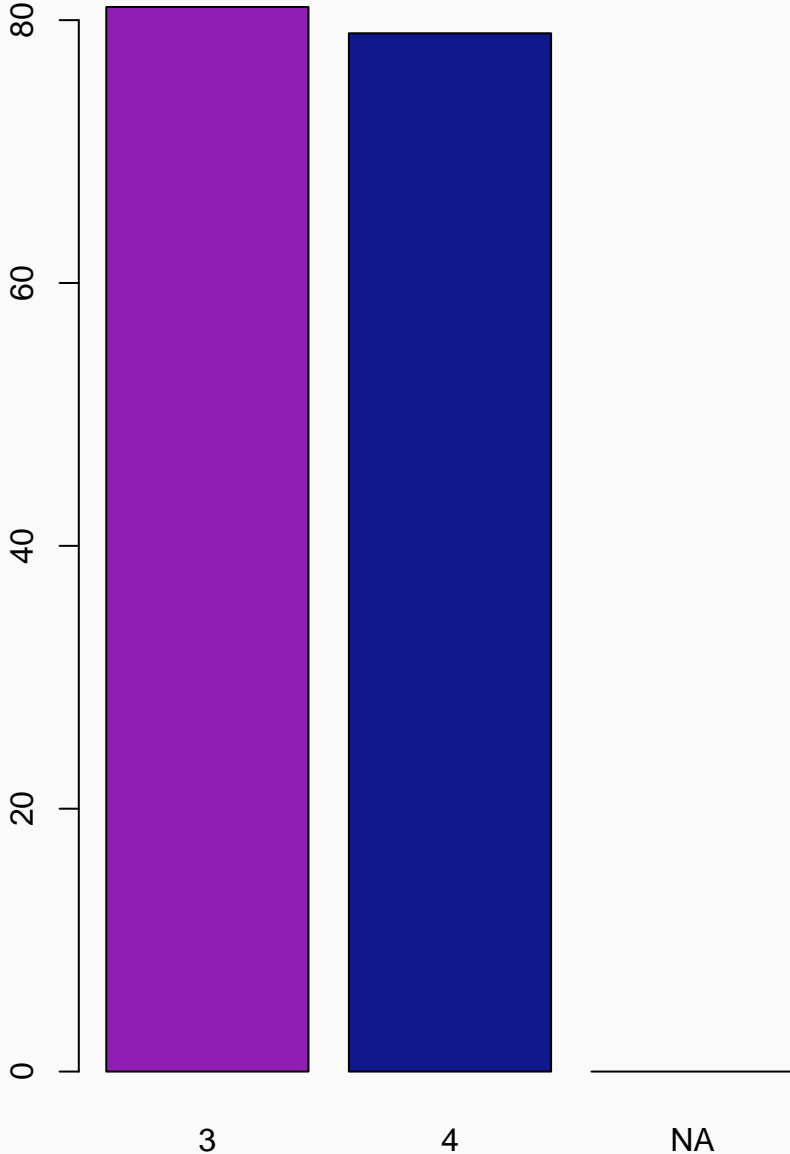
Dose in P2: 4

Missing: 0%

p-value: $8.74e-01$

sequence: NA

seq. position: 80527



solcap_snp_c2_41437

marker #: 1

Dose in P1: 3

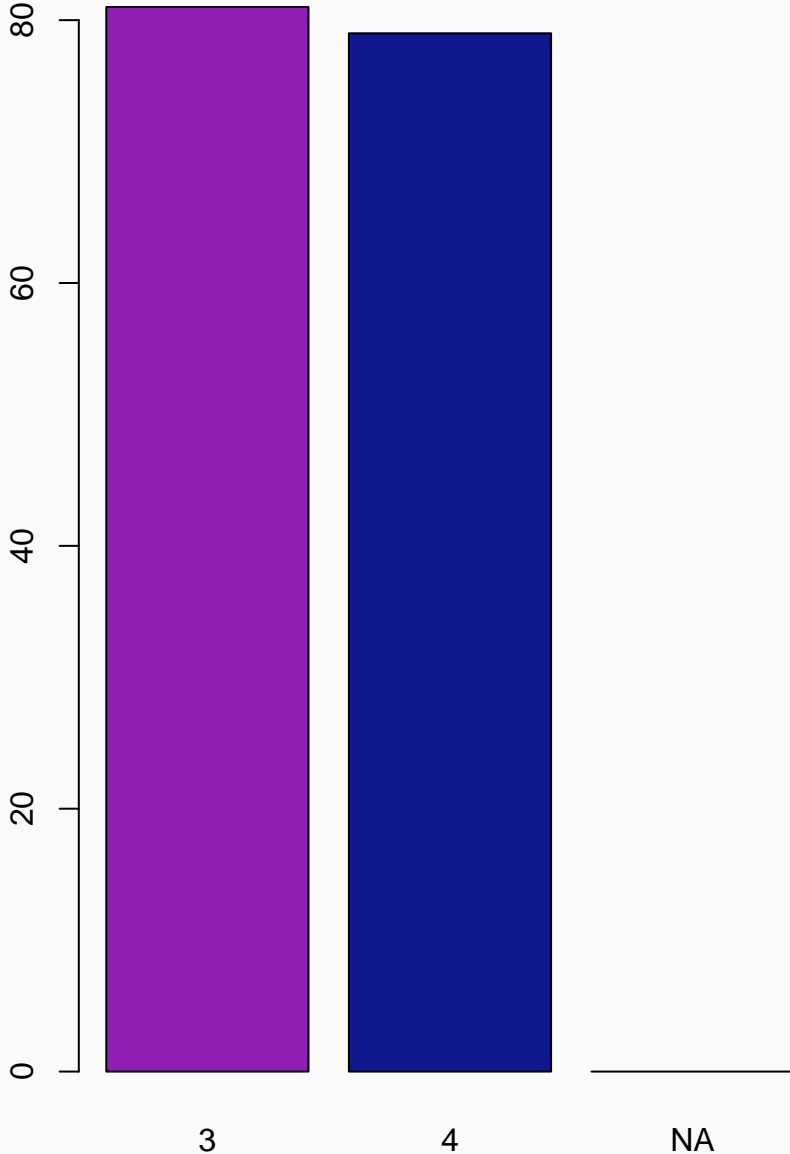
Dose in P2: 4

Missing: 0%

p-value: $8.74e-01$

sequence: NA

seq. position: 80527



solcap_snp_c2_41437

marker #: 1

Dose in P1: 3

Dose in P2: 4

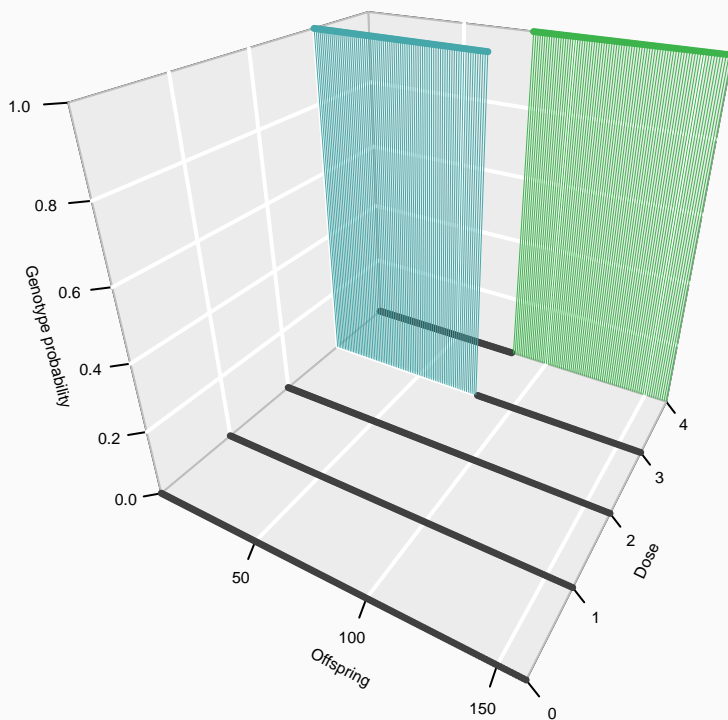
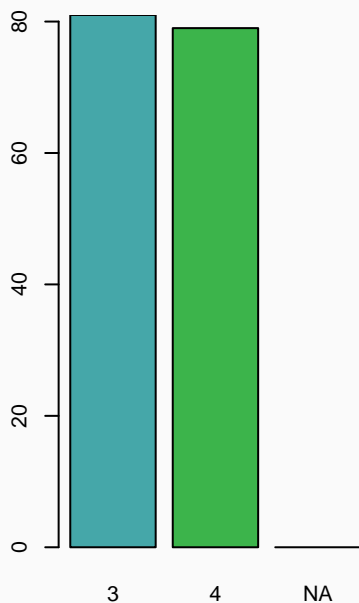
Missing: 0%

p-value: 8.74e-01

sequence: NA

seq. position: 805279

prob. threshold: 0.95



solcap_snp_c2_41437

marker #: 1

Dose in P1: 3

Dose in P2: 4

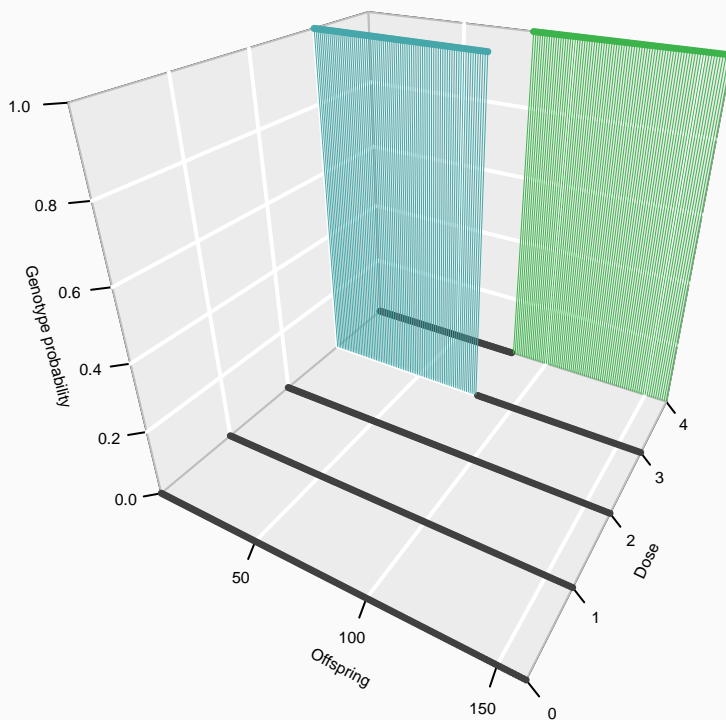
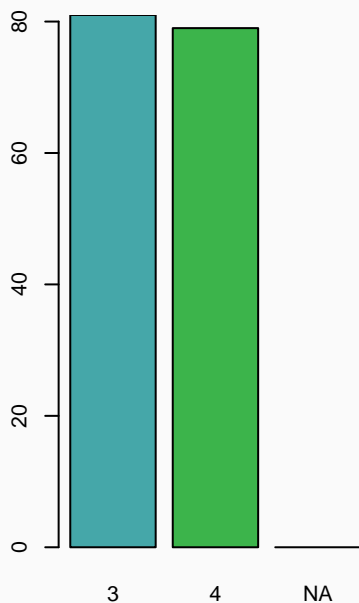
Missing: 0%

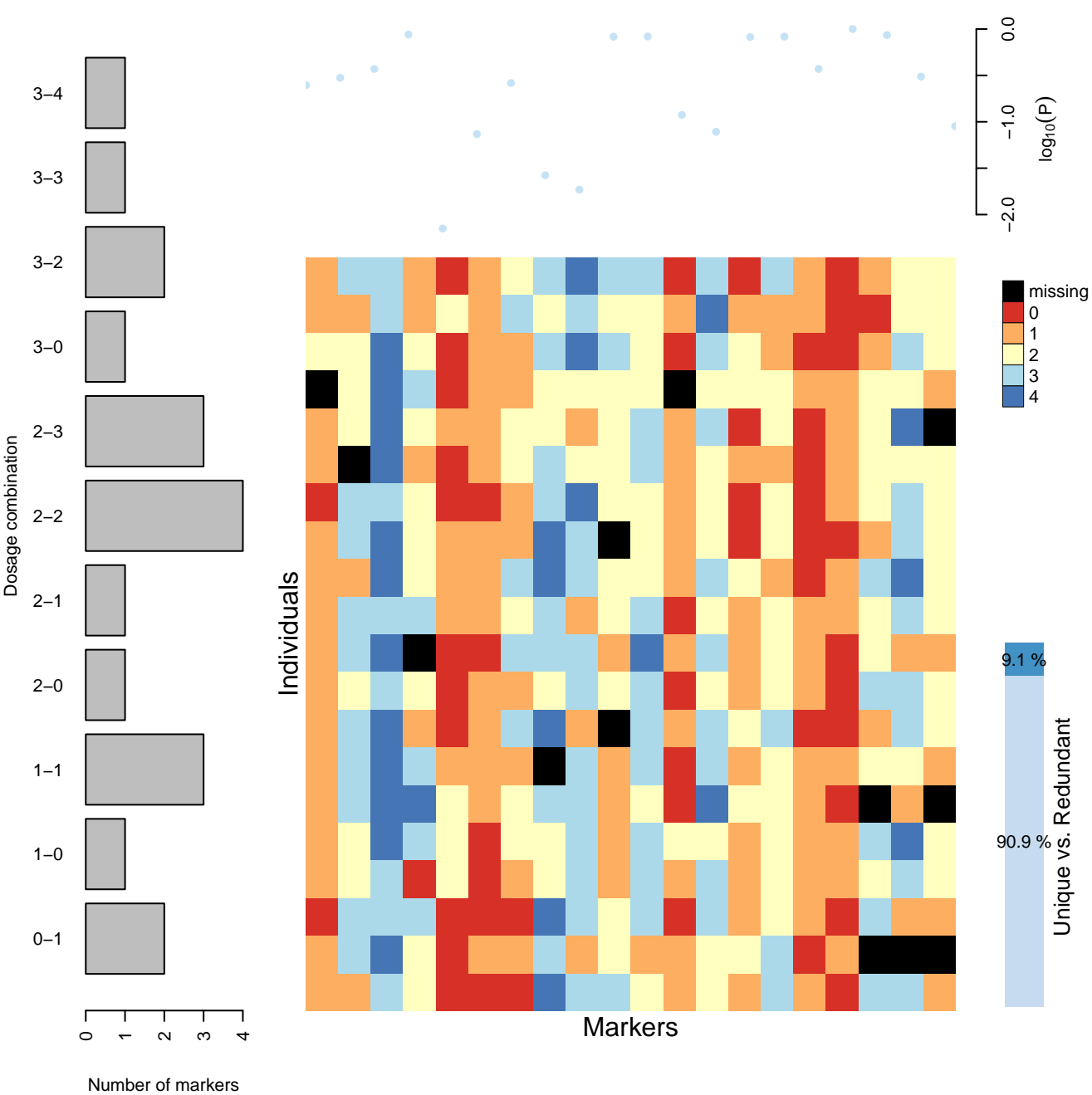
p-value: 8.74e-01

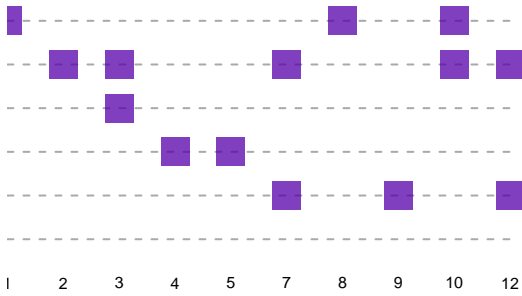
sequence: NA

seq. position: 805279

prob. threshold: 0.95







X

