Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 1

Purity: 0.22 Tumor ploidy: 2.446 SNV log–likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 2
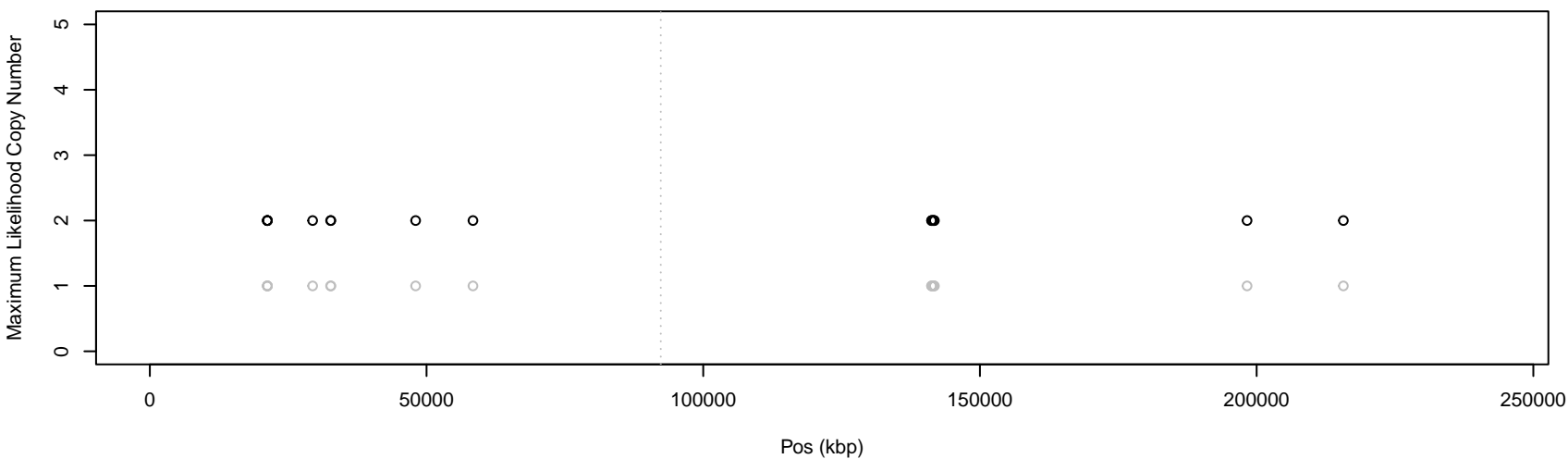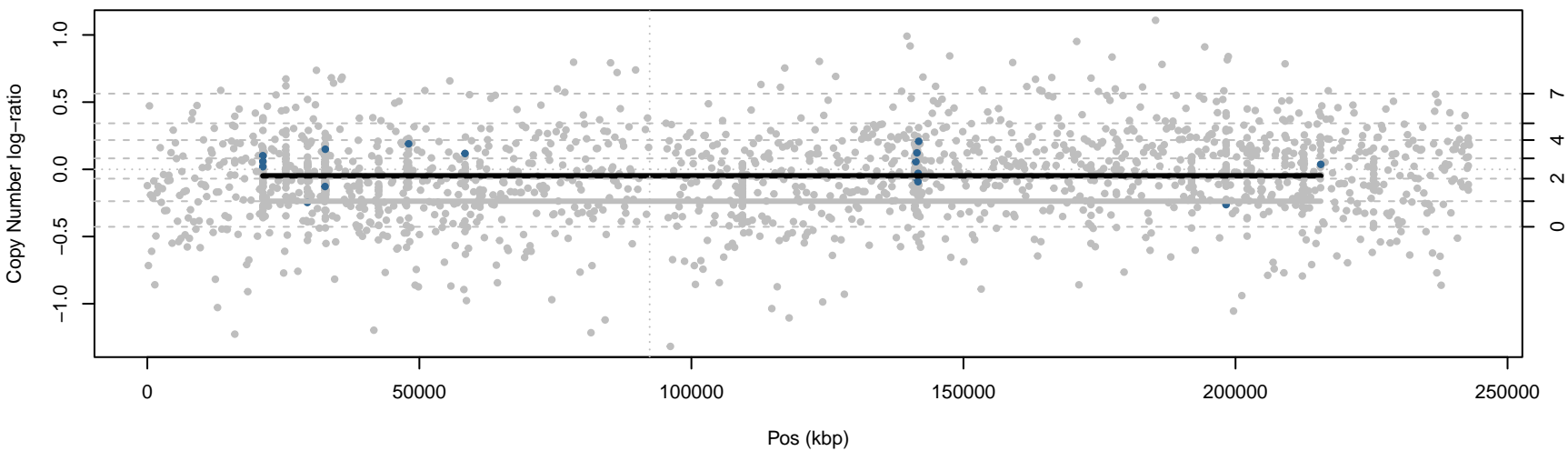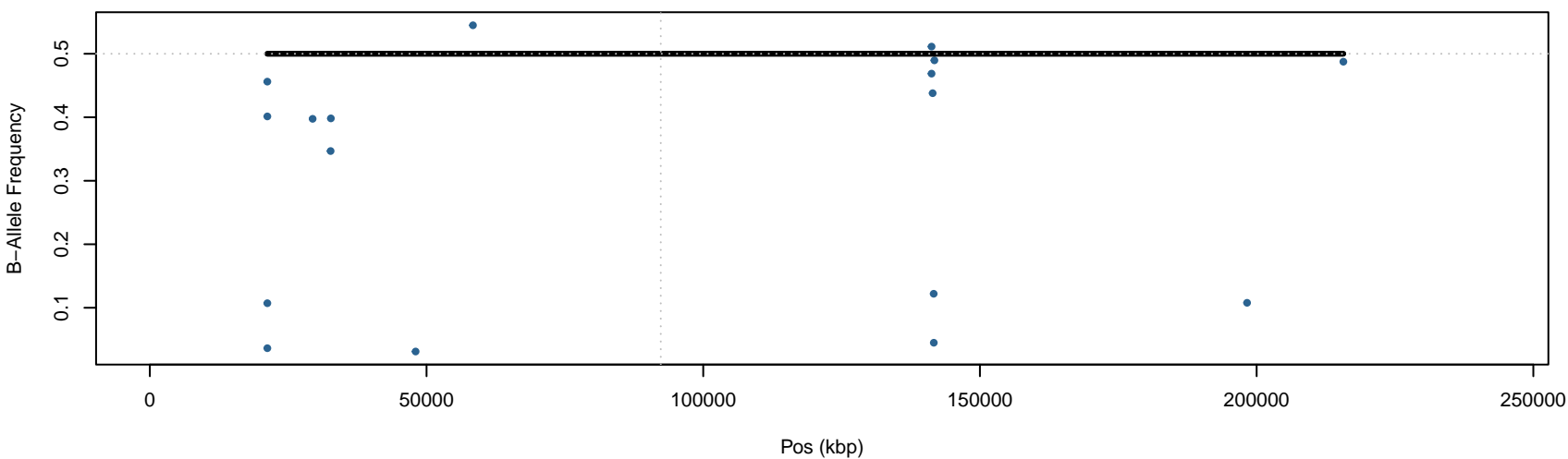
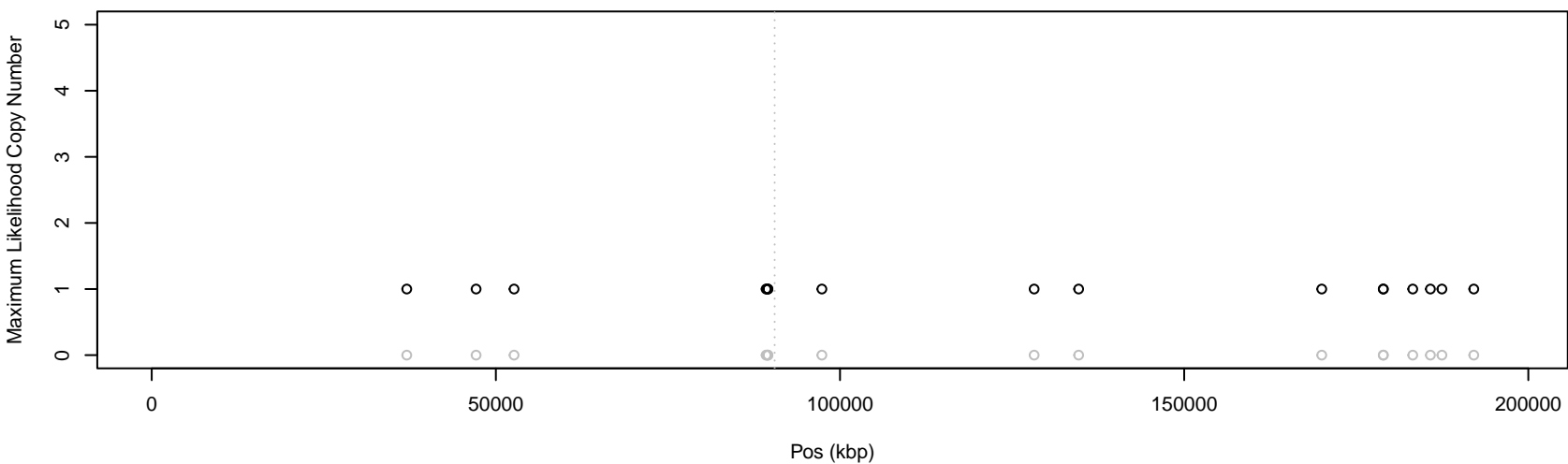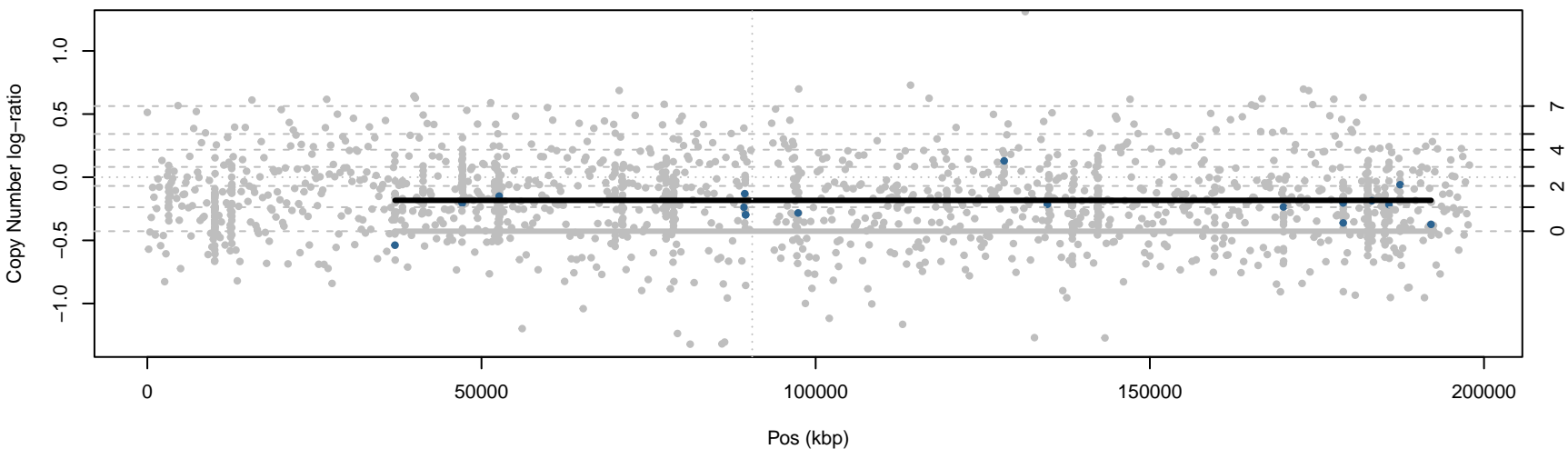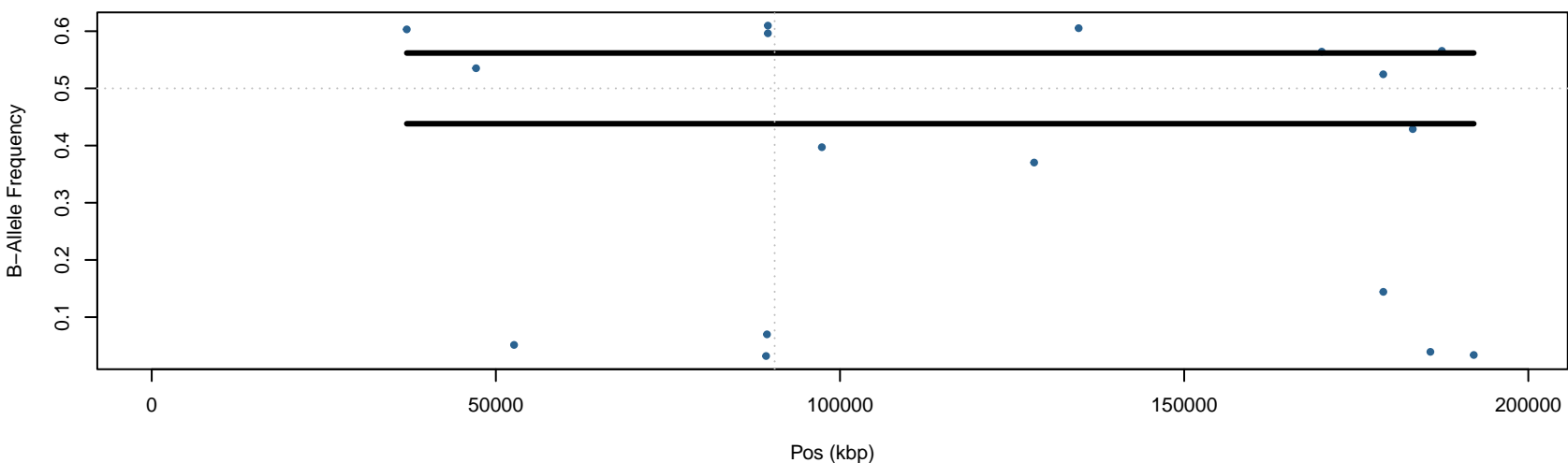**Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 3**

Purity: 0.22 Tumor ploidy: 2.446 SNV log–likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 5
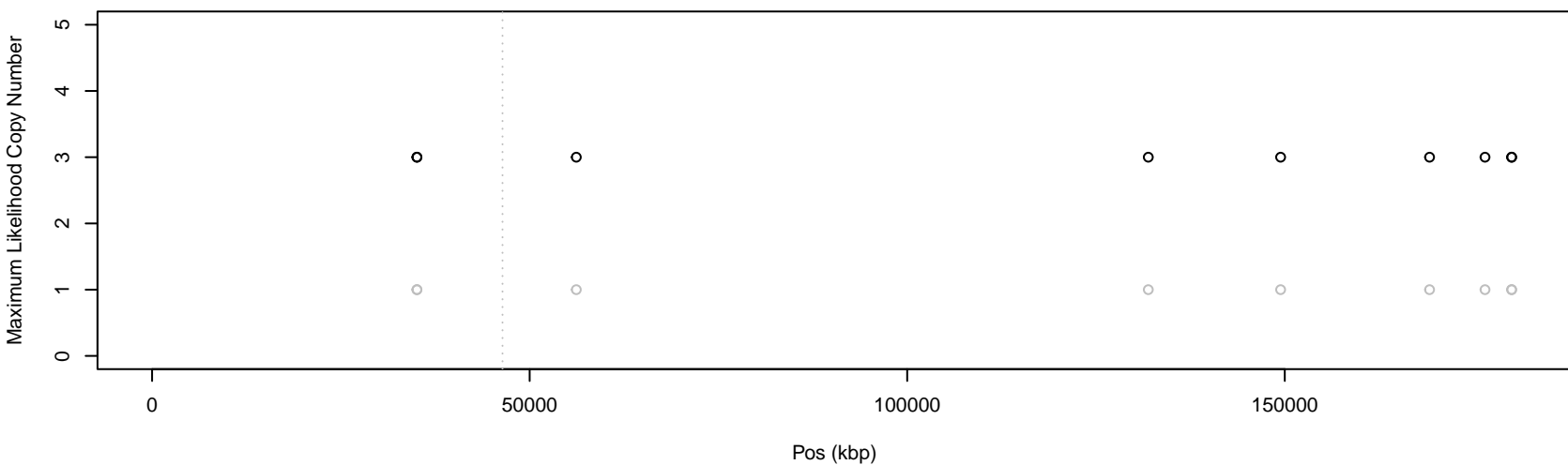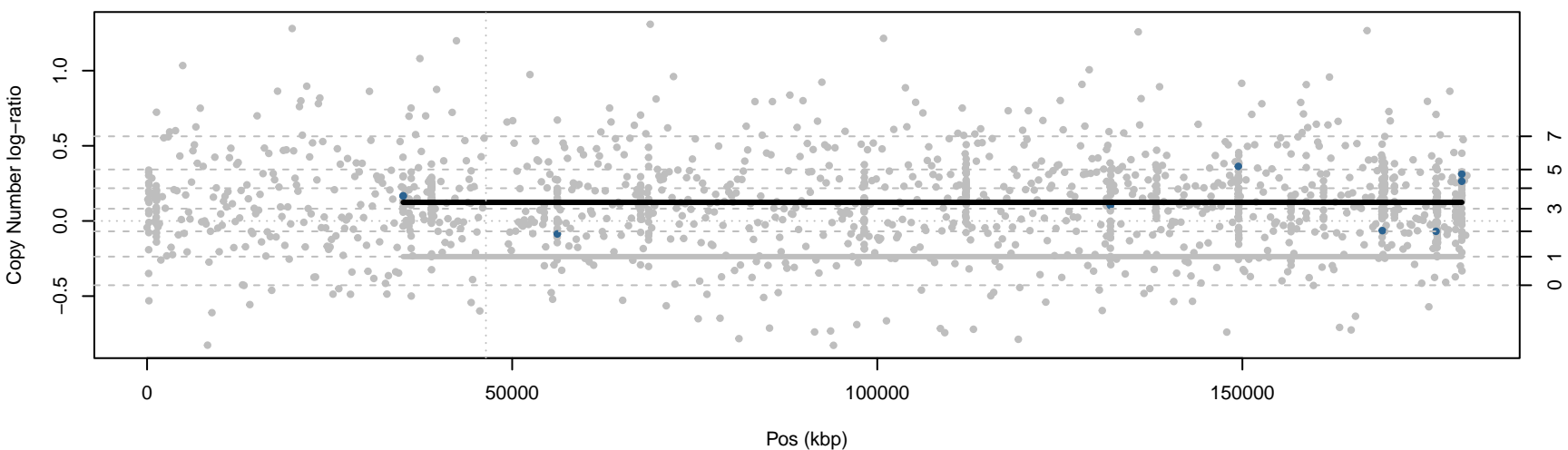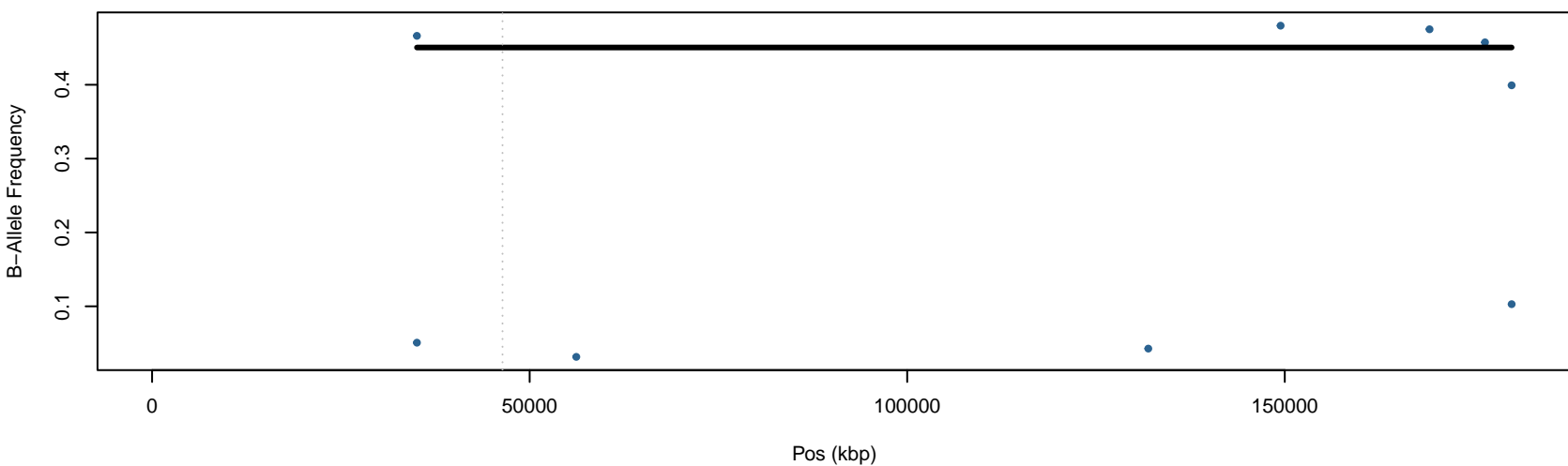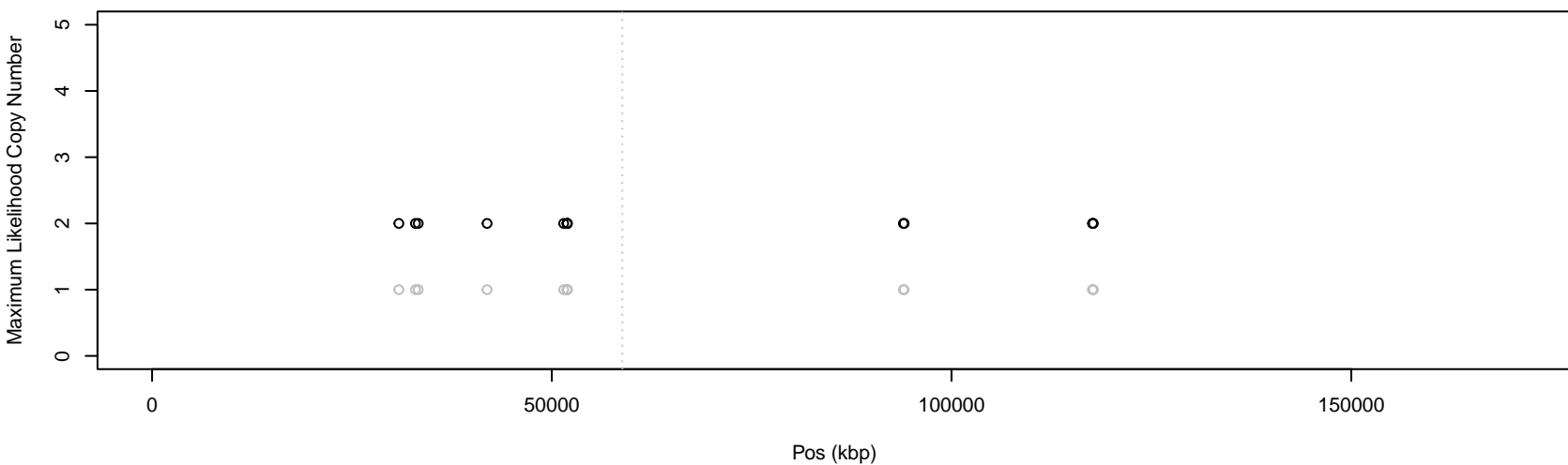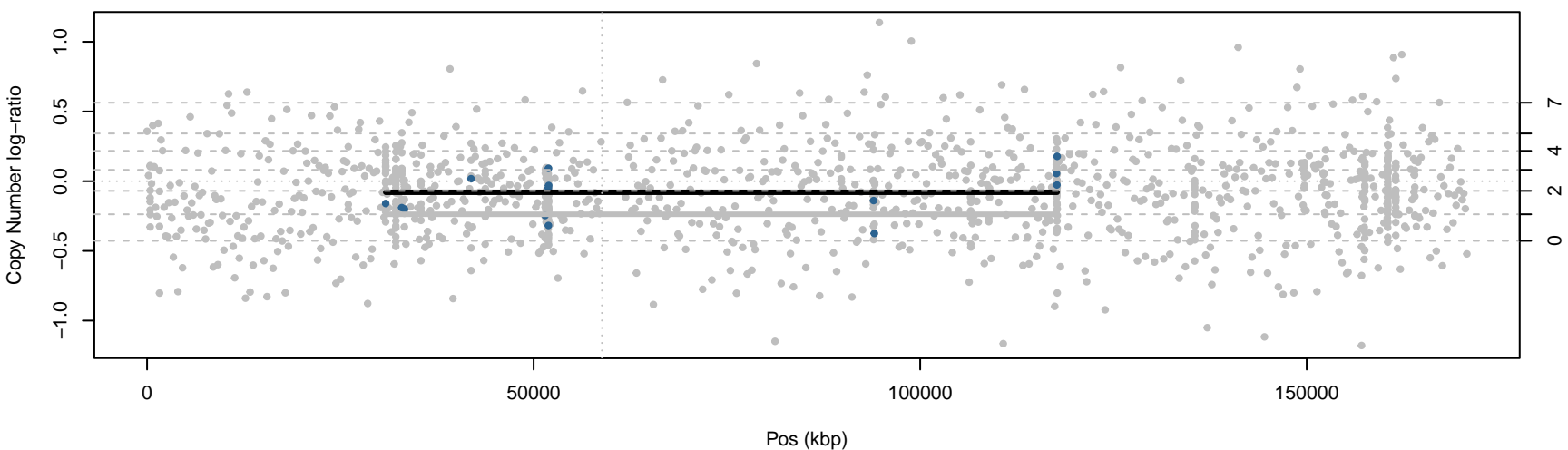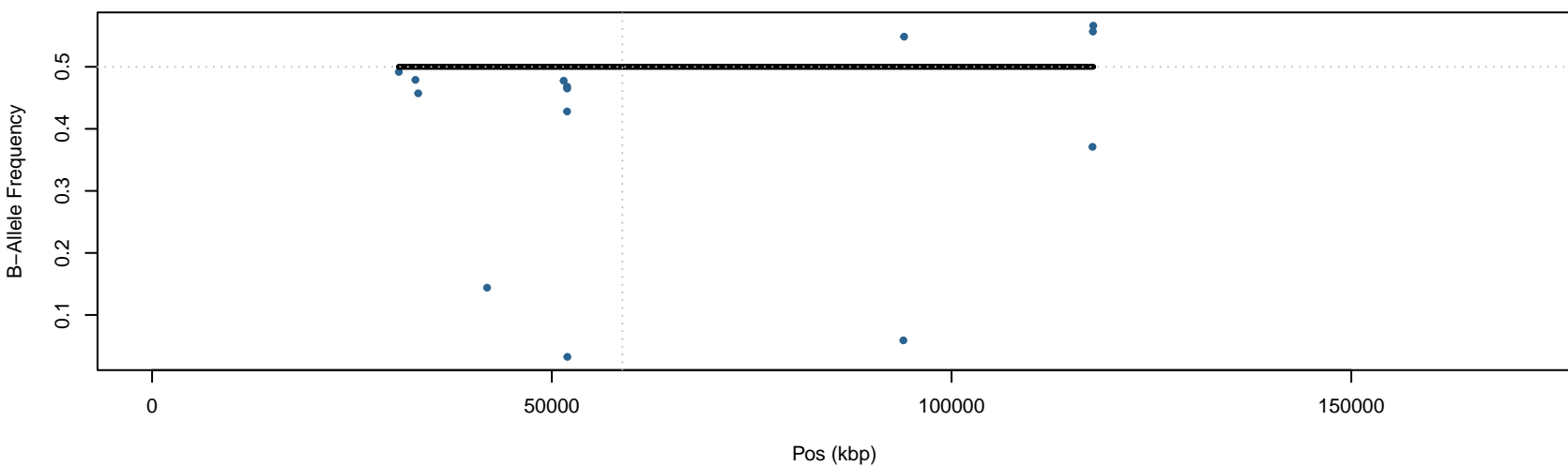
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 6
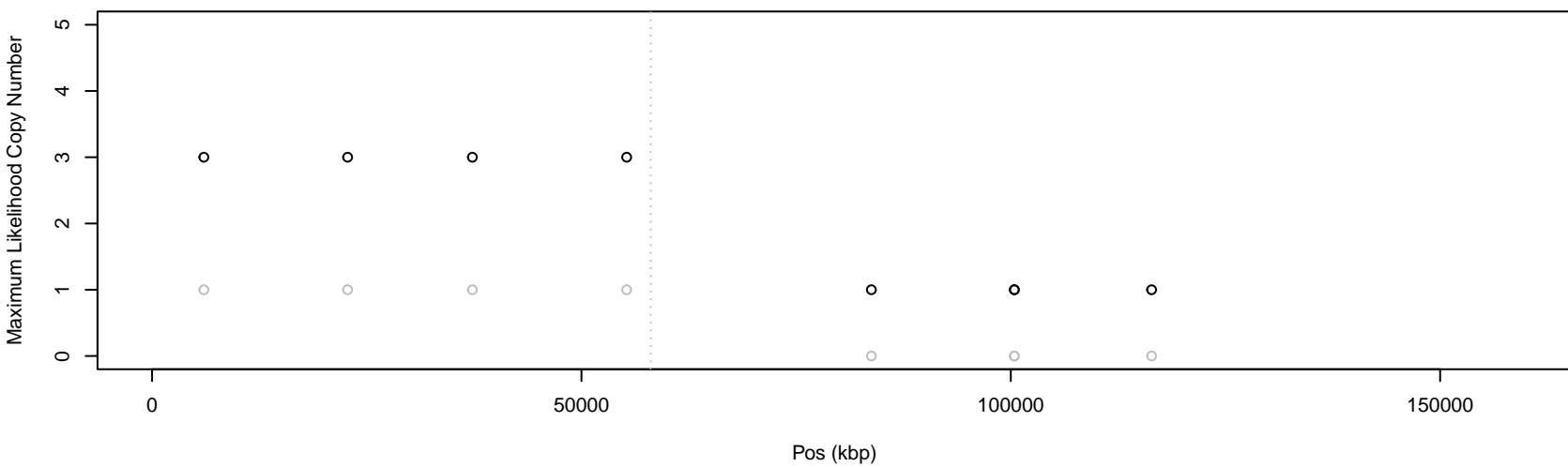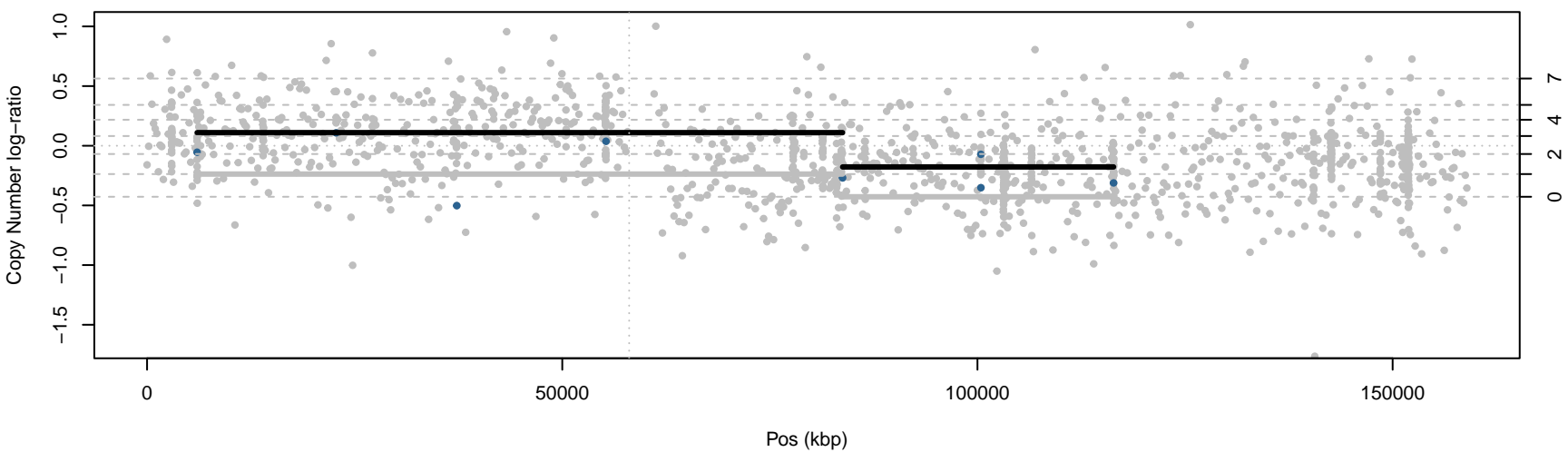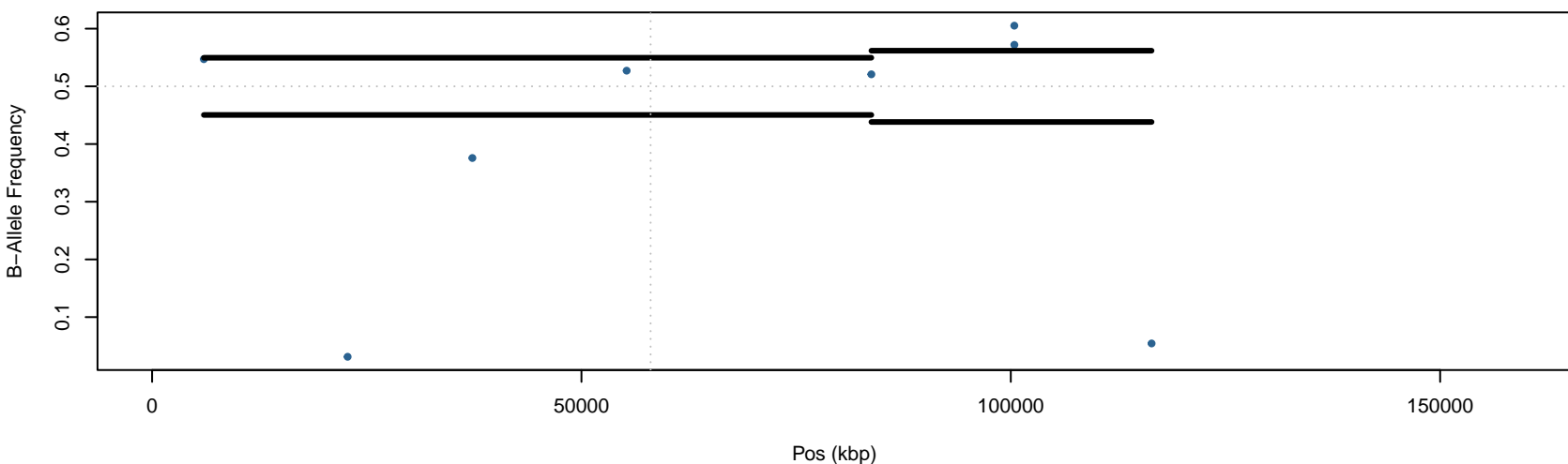
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 7
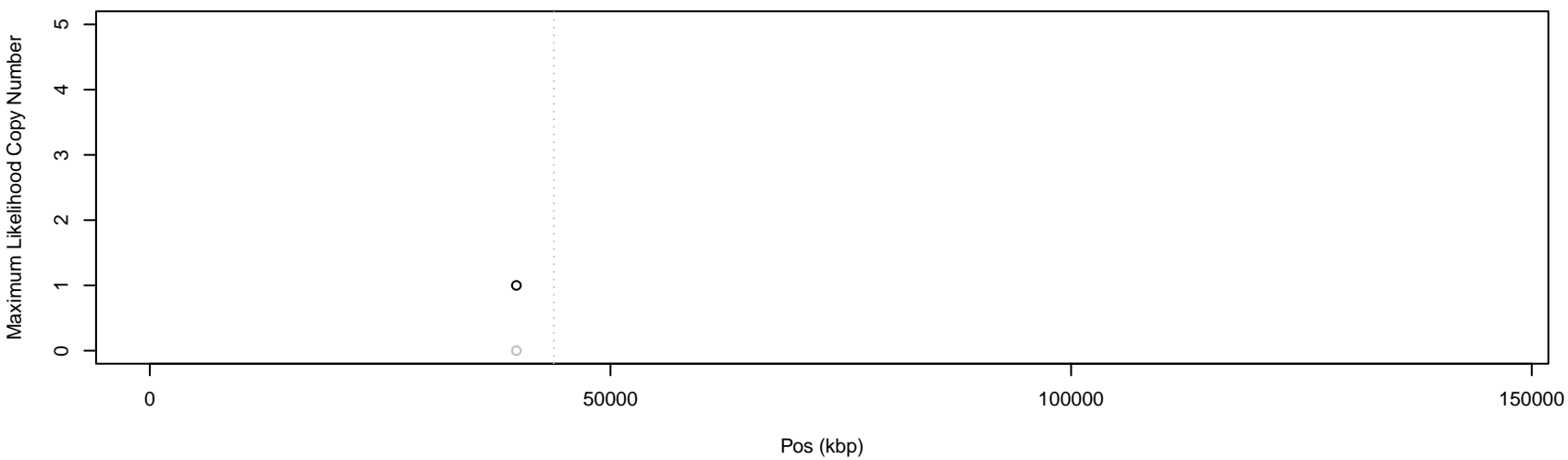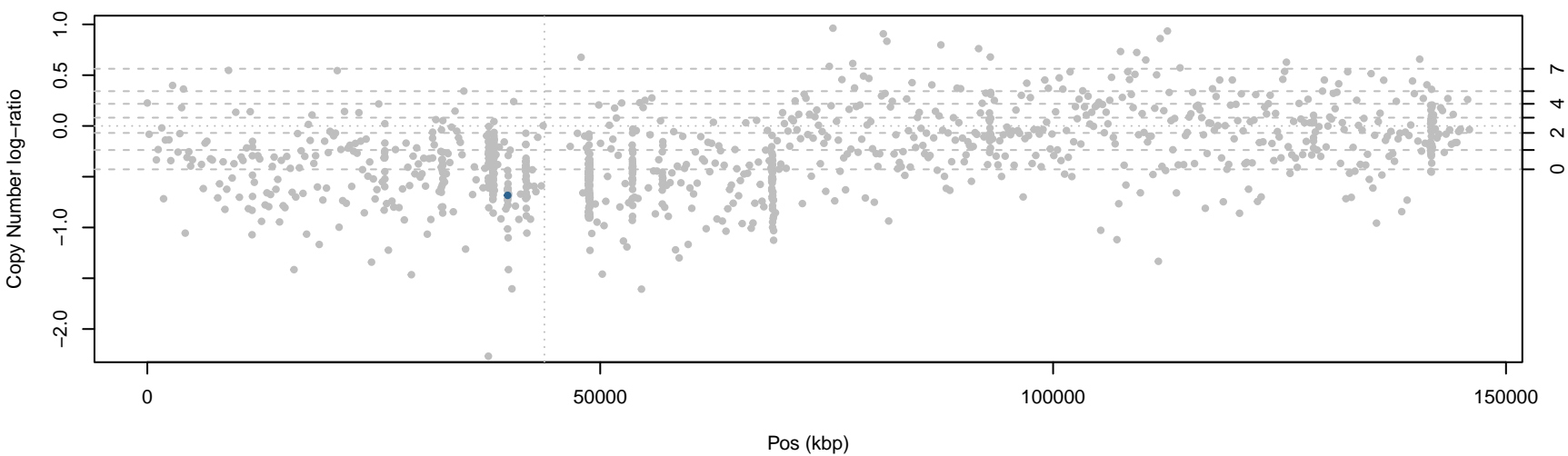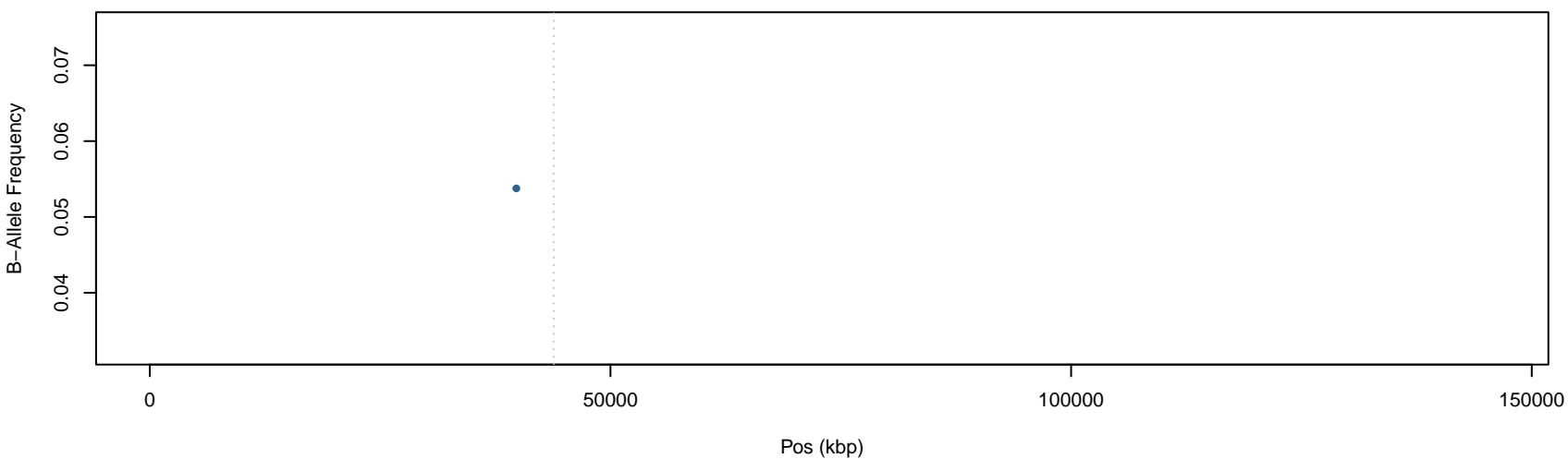
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 8

Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 9
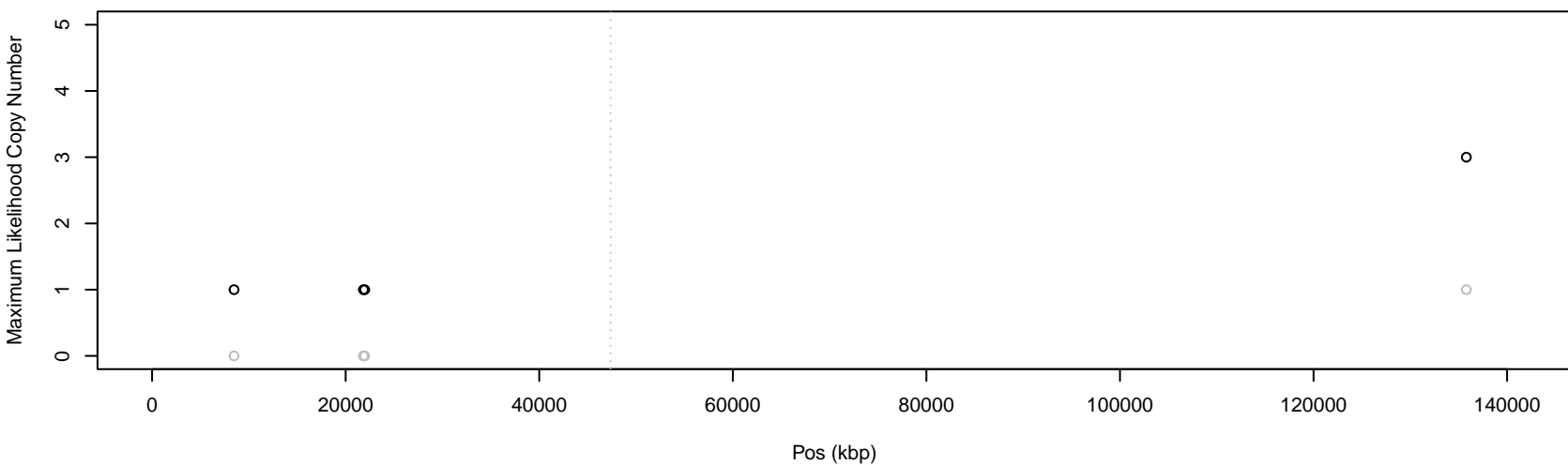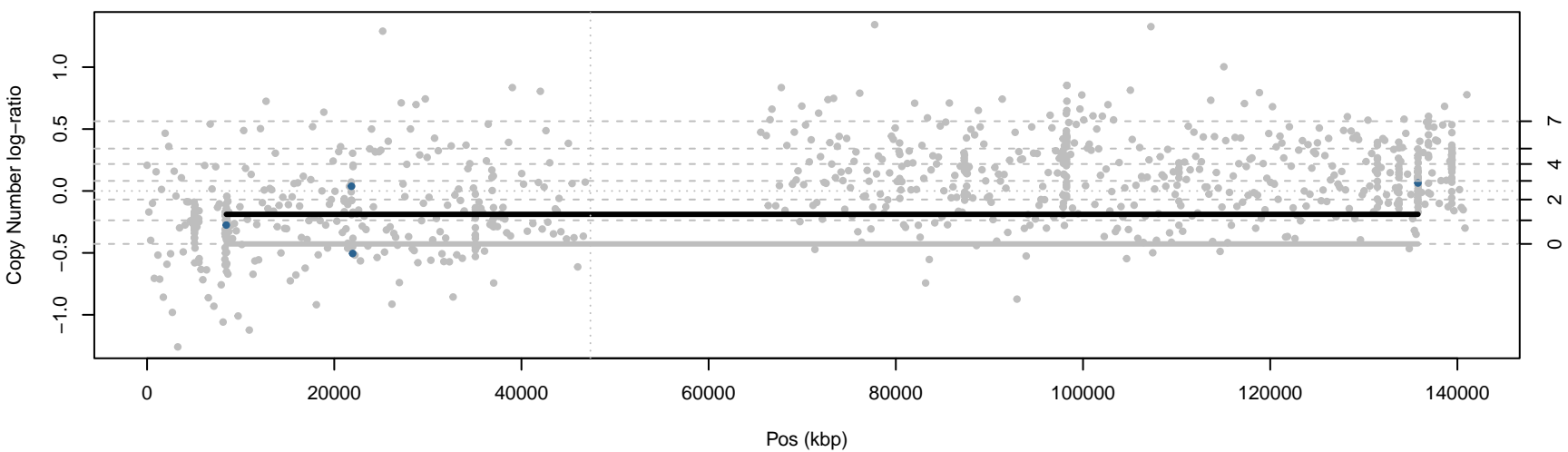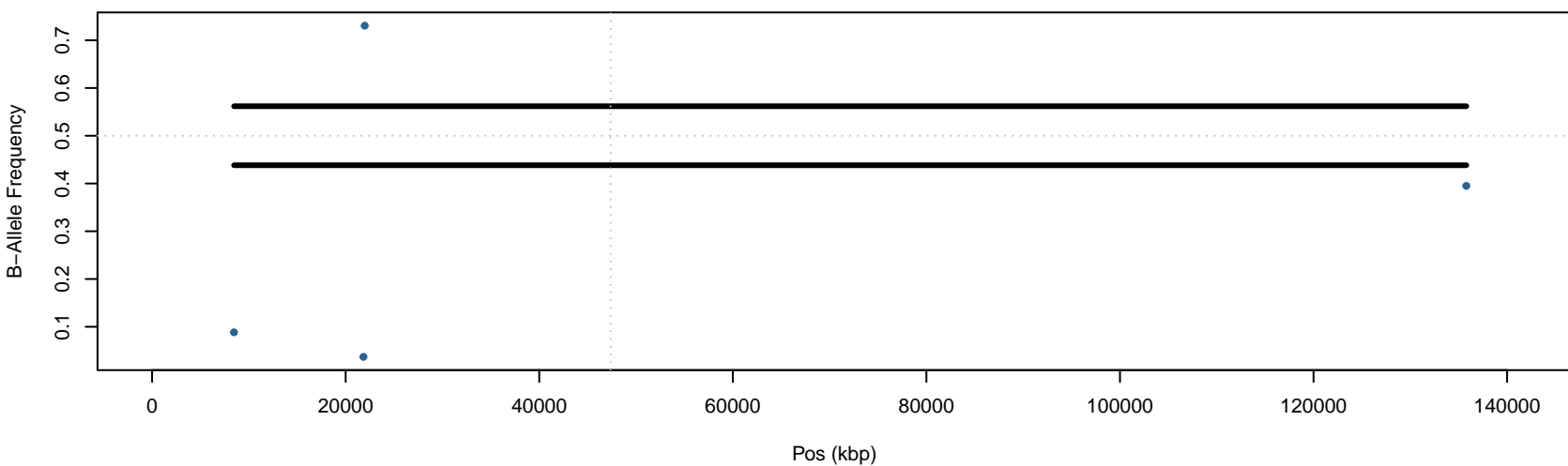
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 10
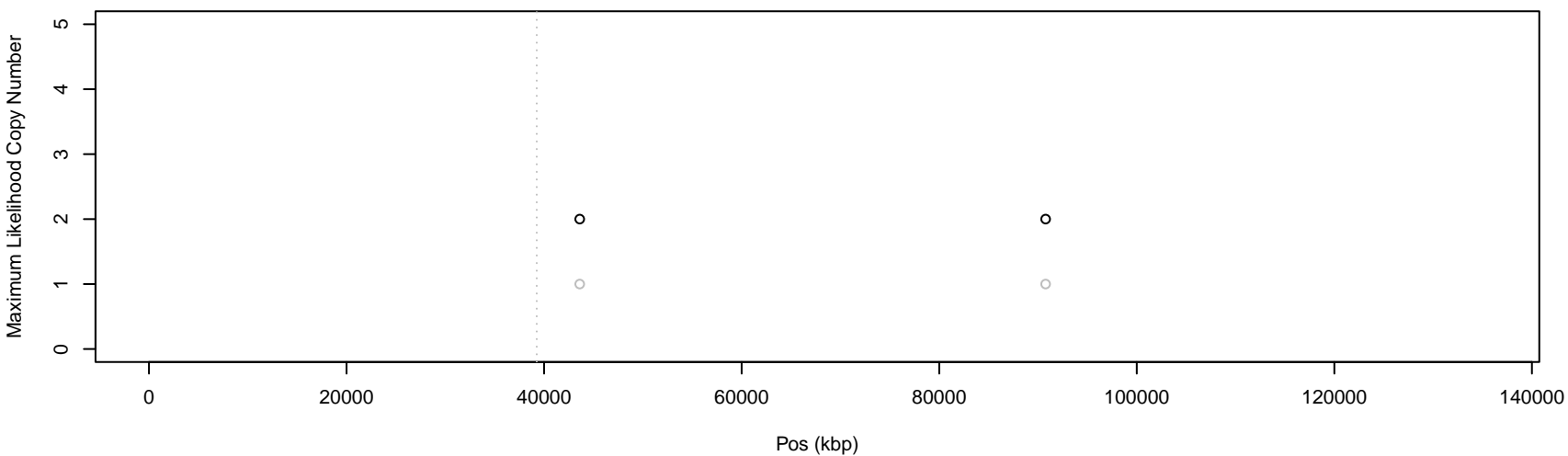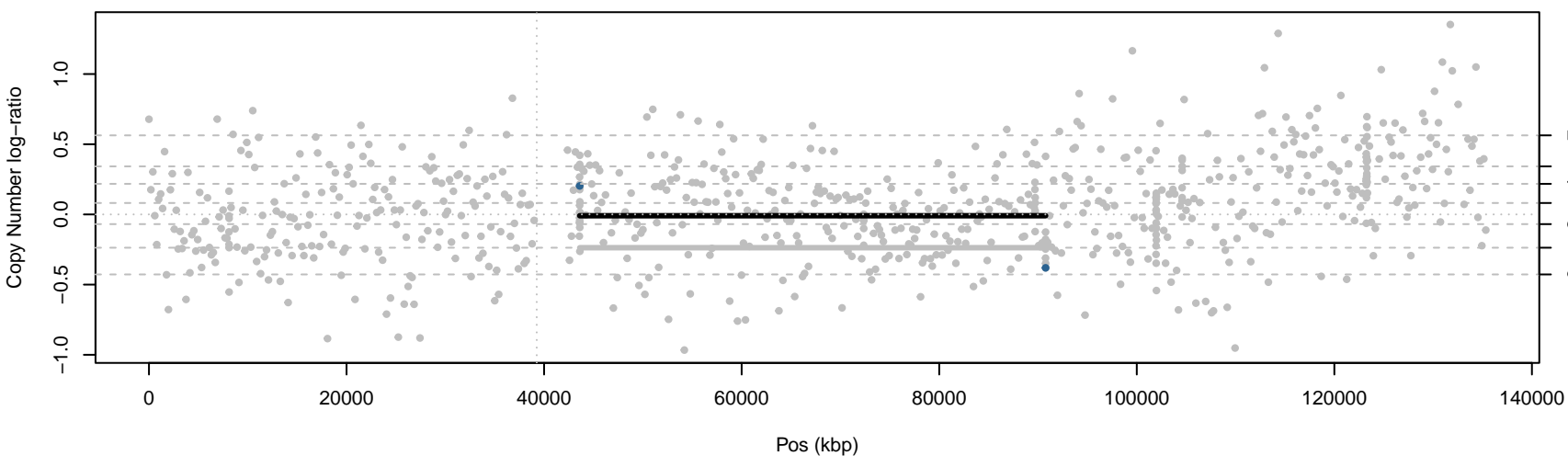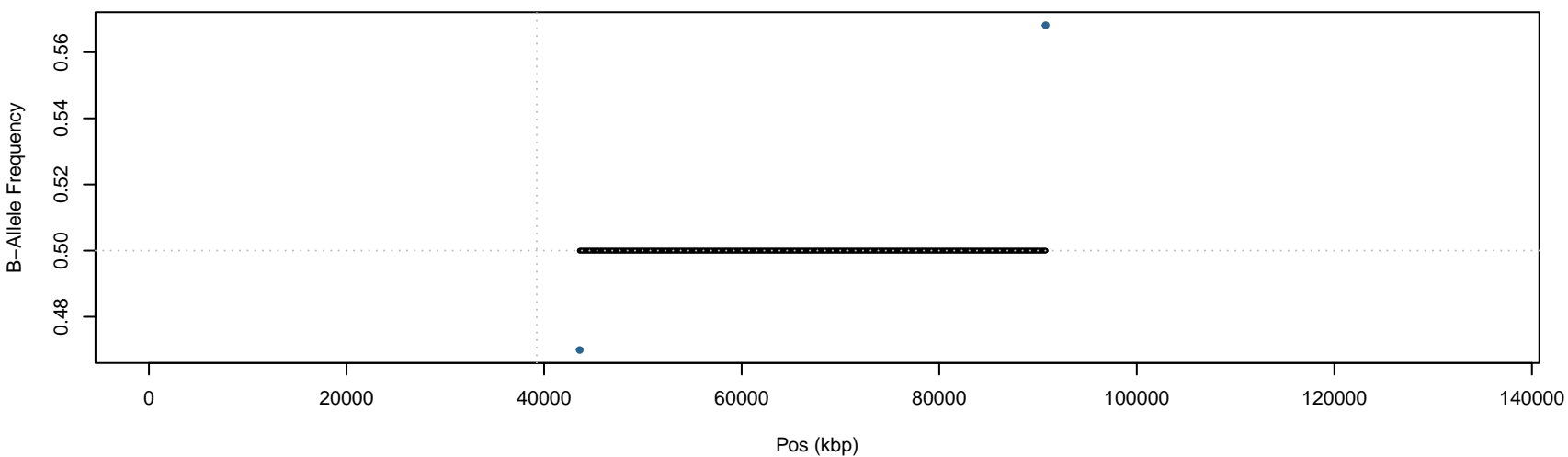
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 11
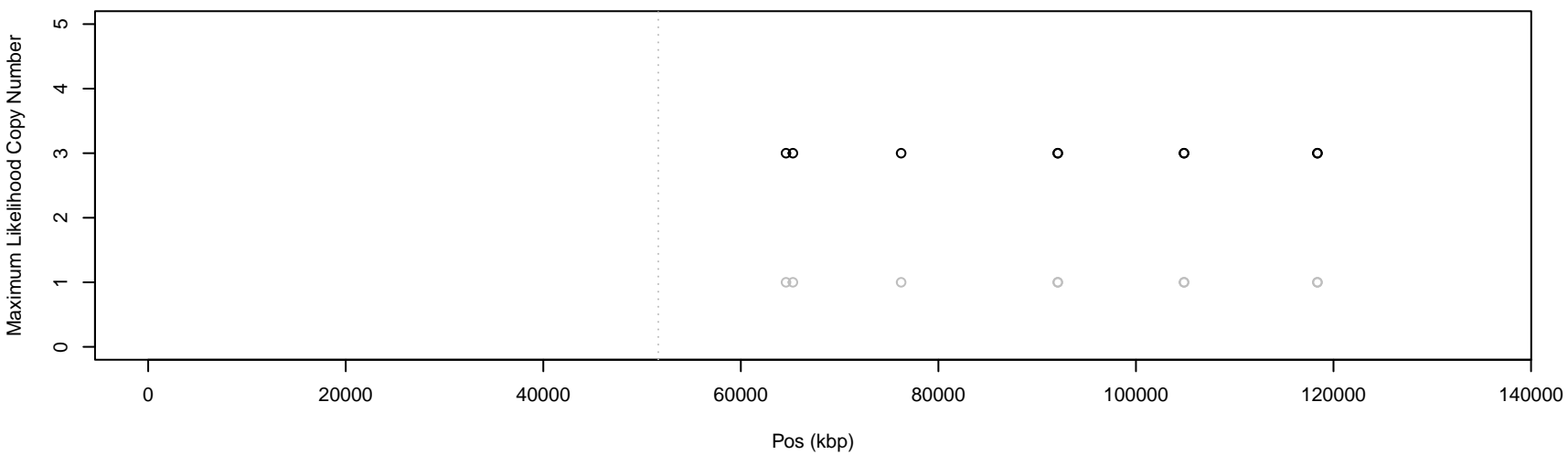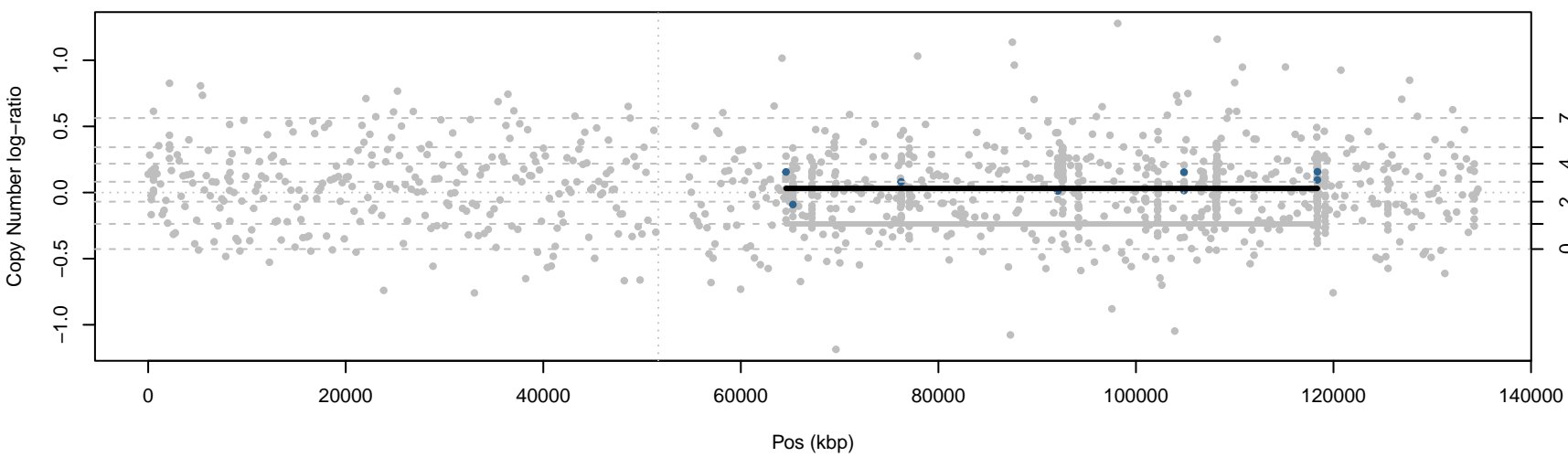
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 12

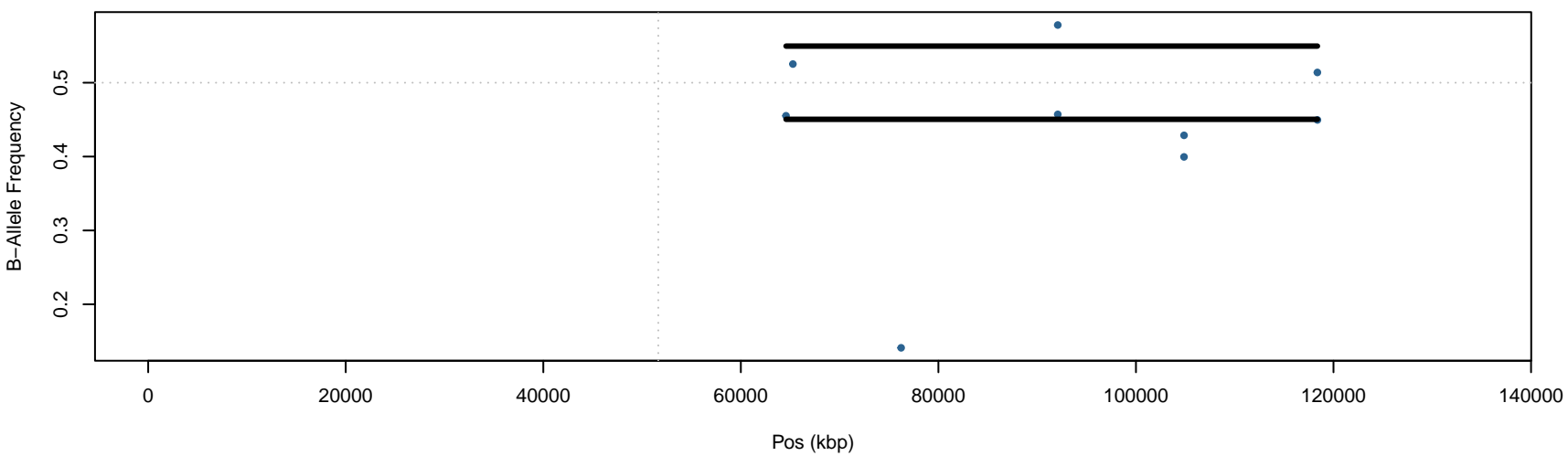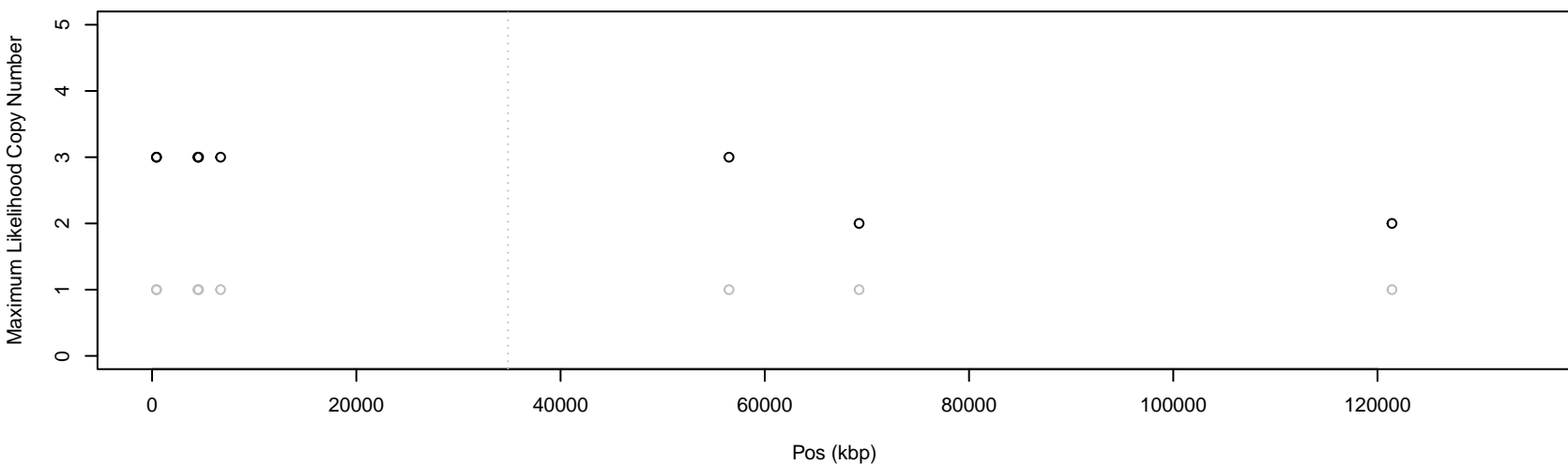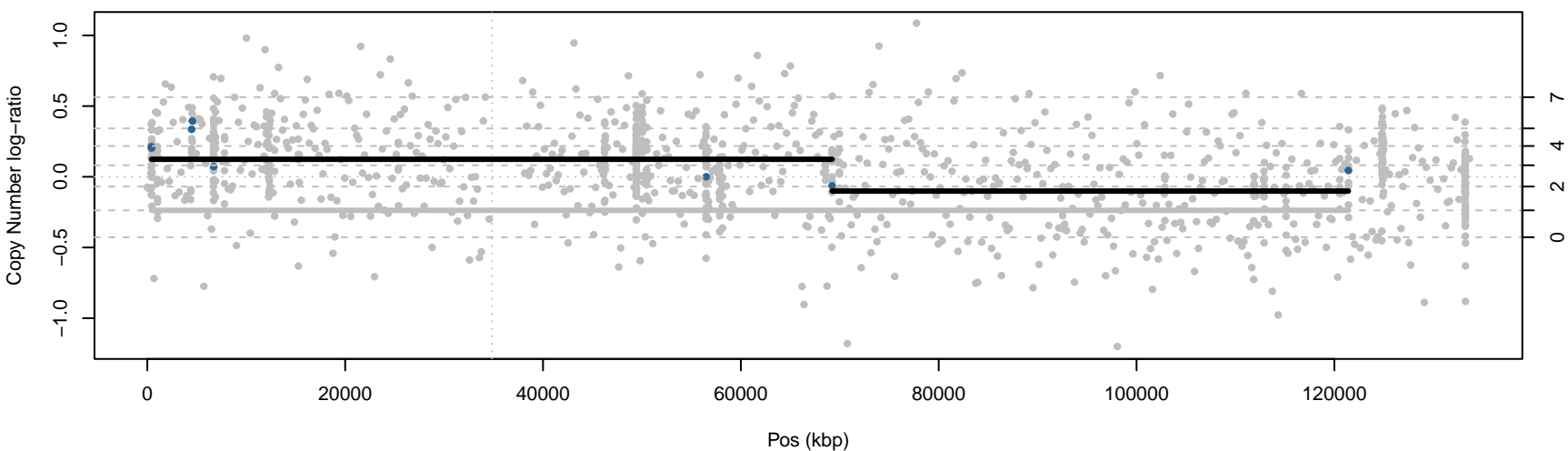**Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 13**
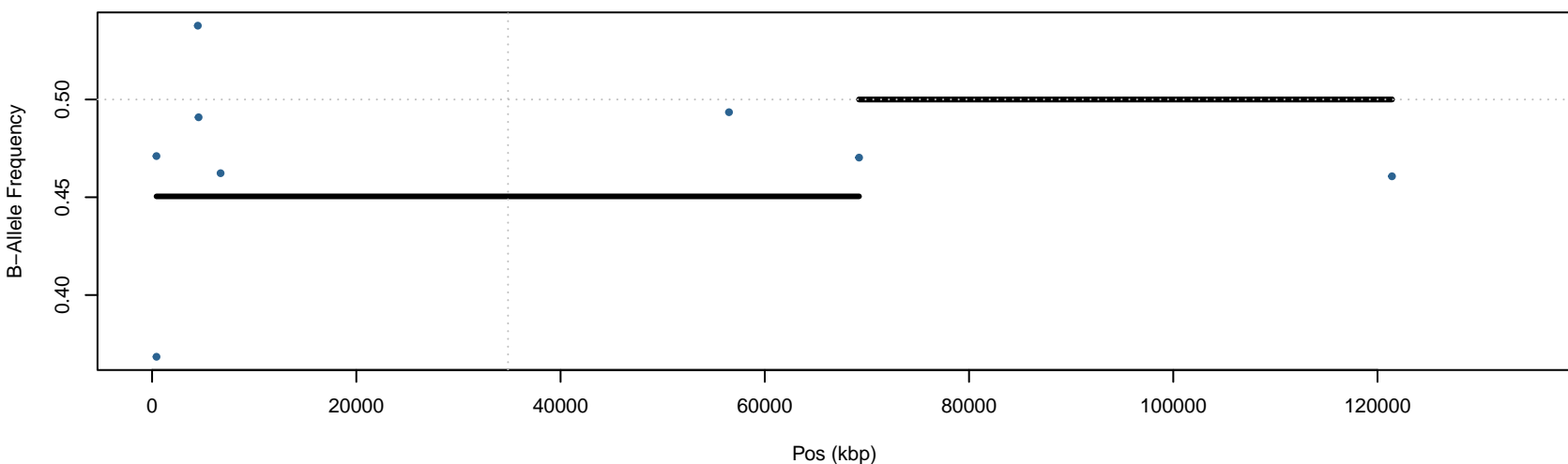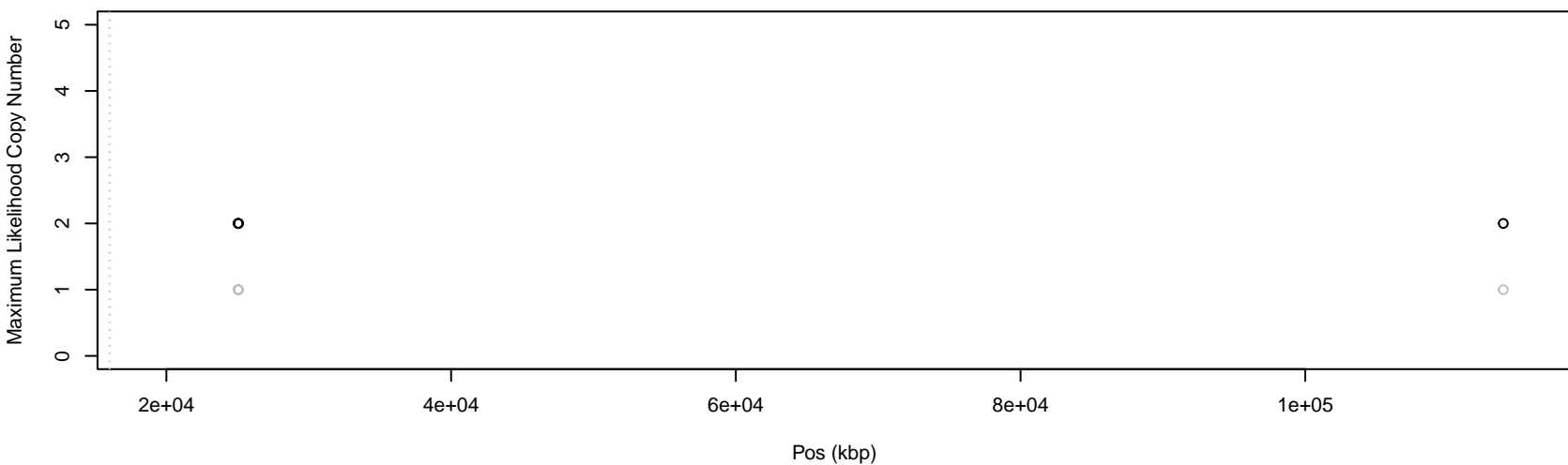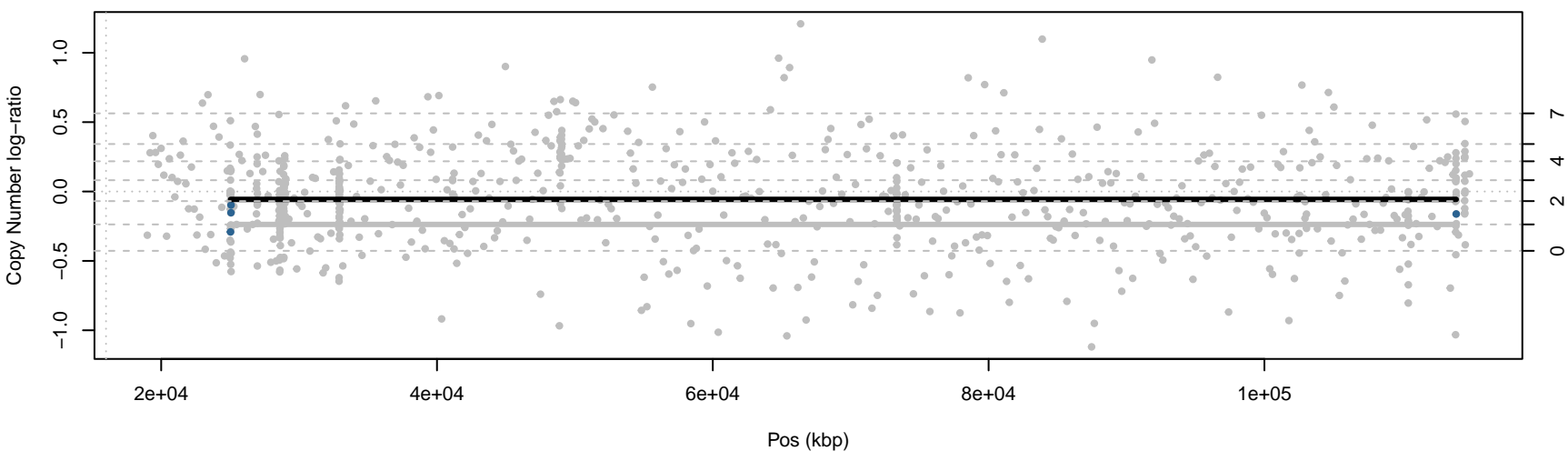
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 19
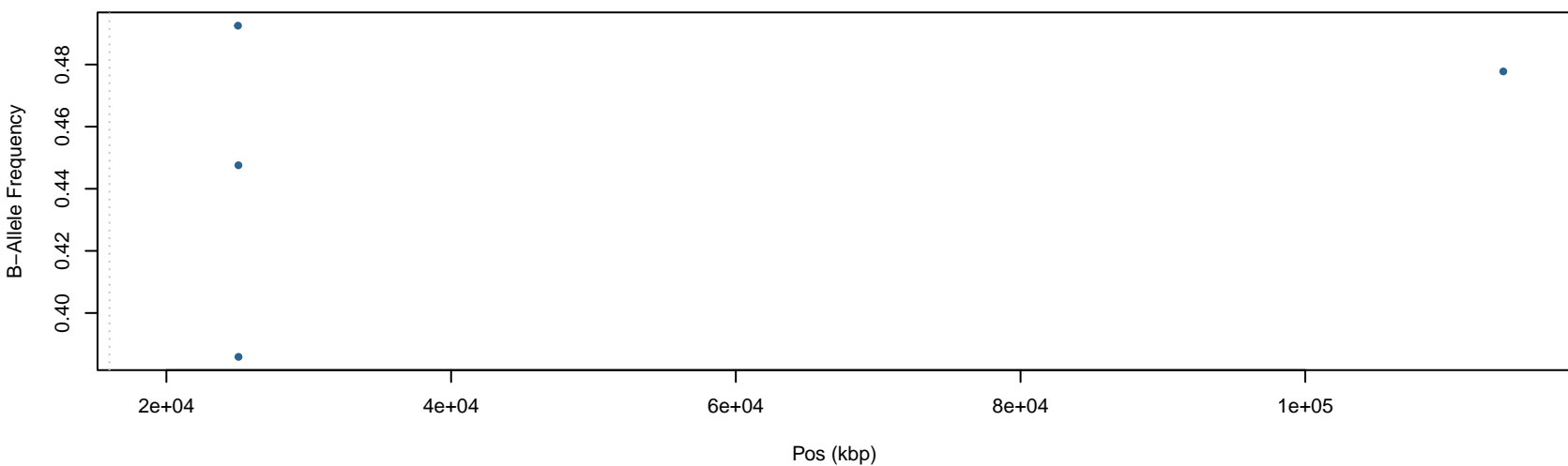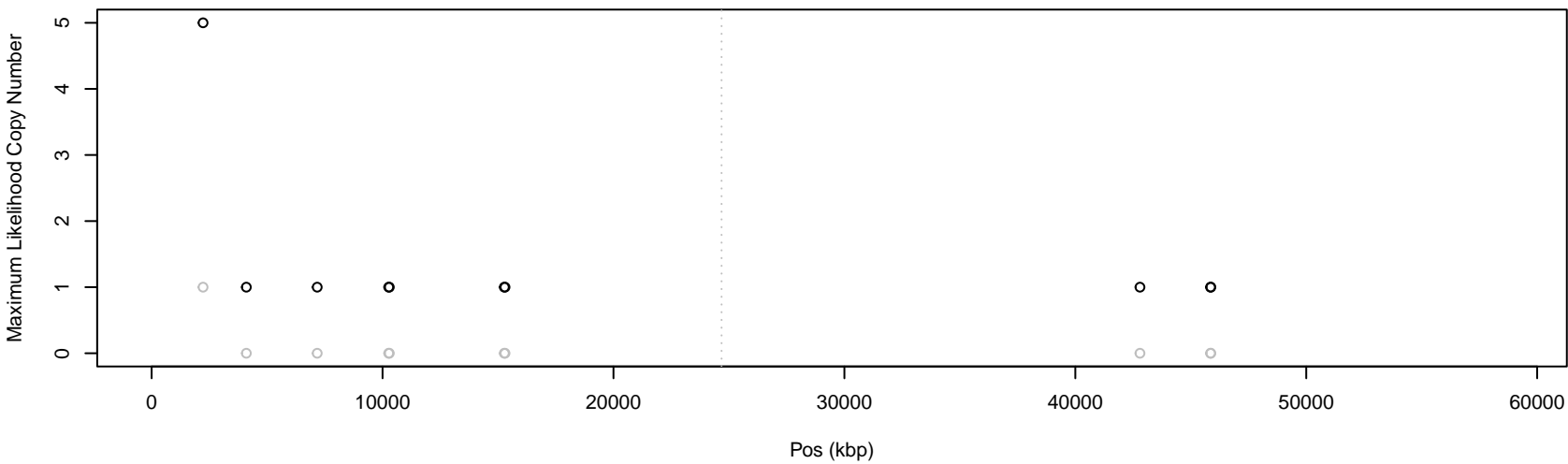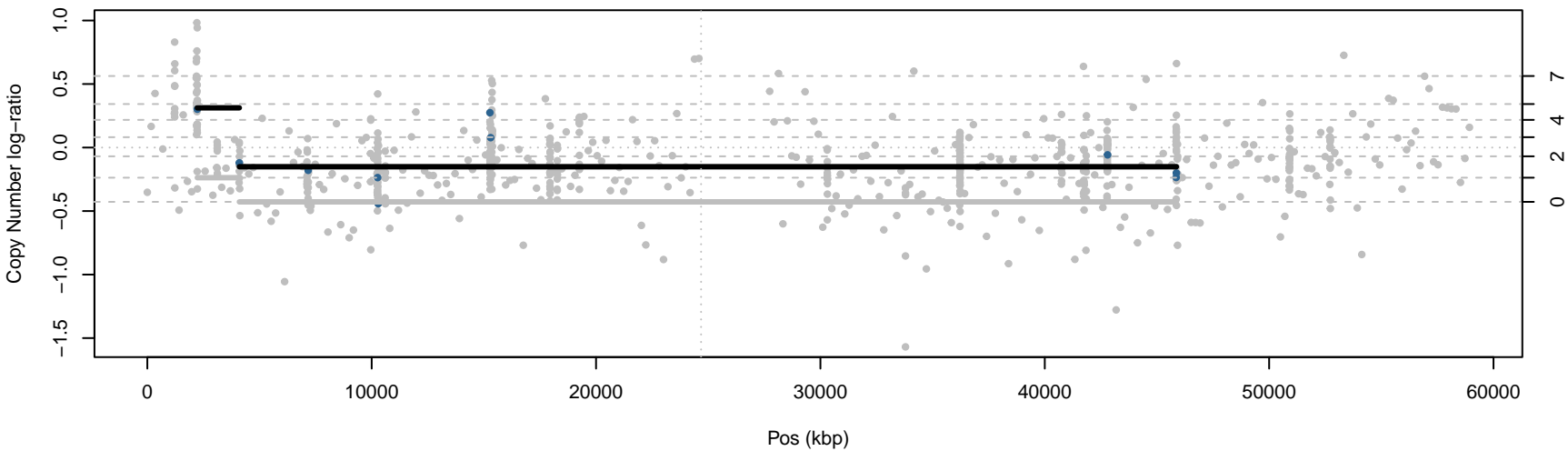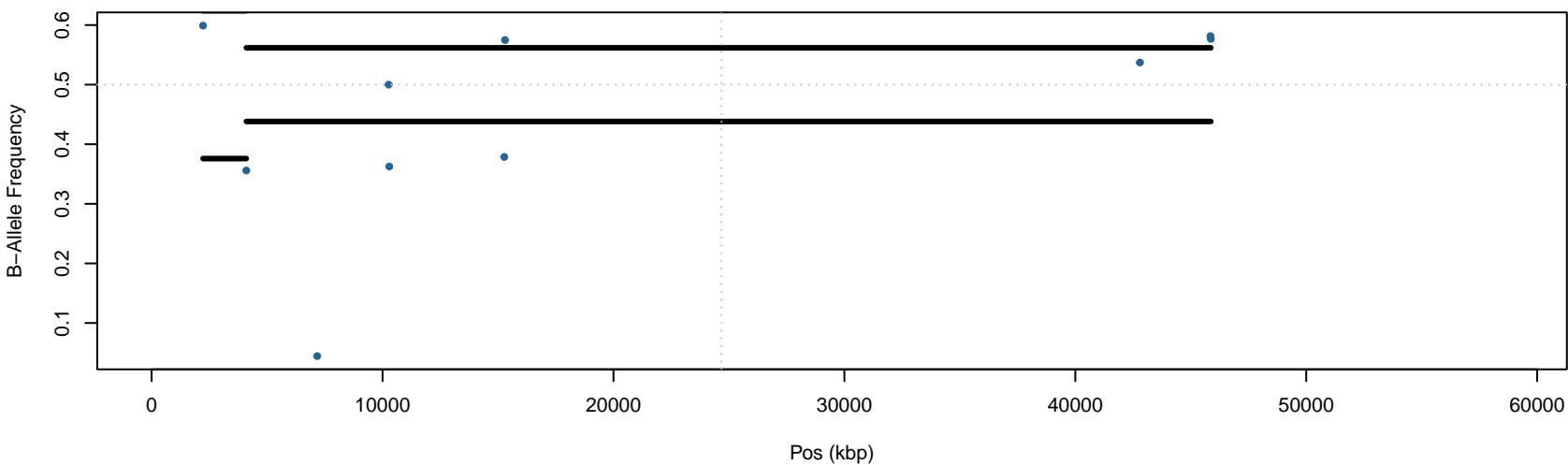
Purity: 0.22 Tumor ploidy: 2.446 SNV log−likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 20