Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 1

Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 2
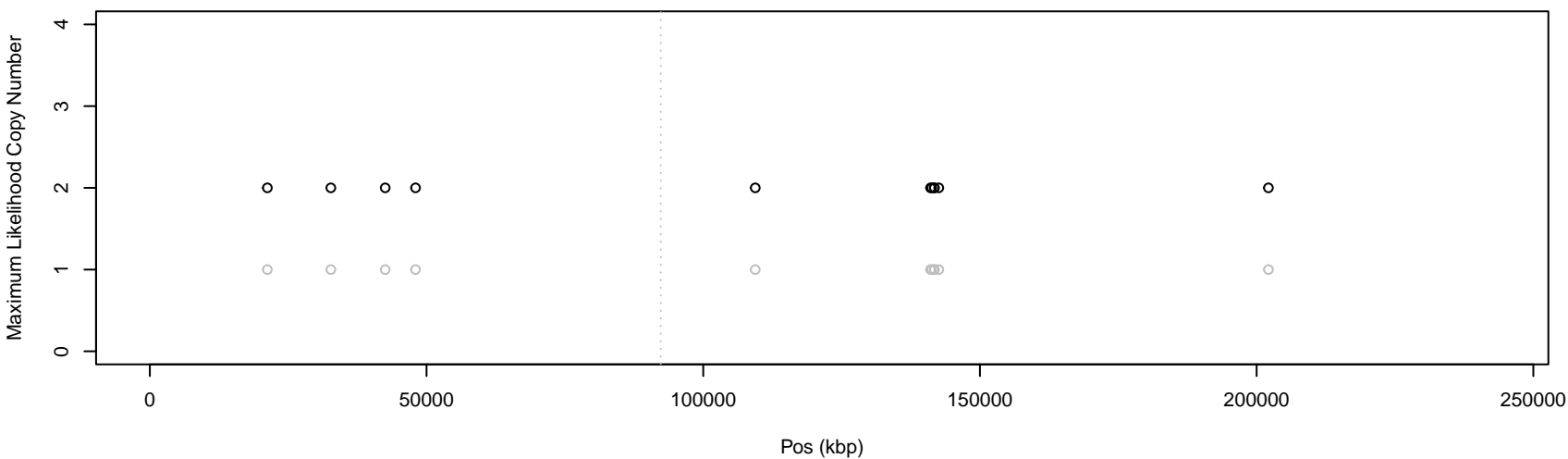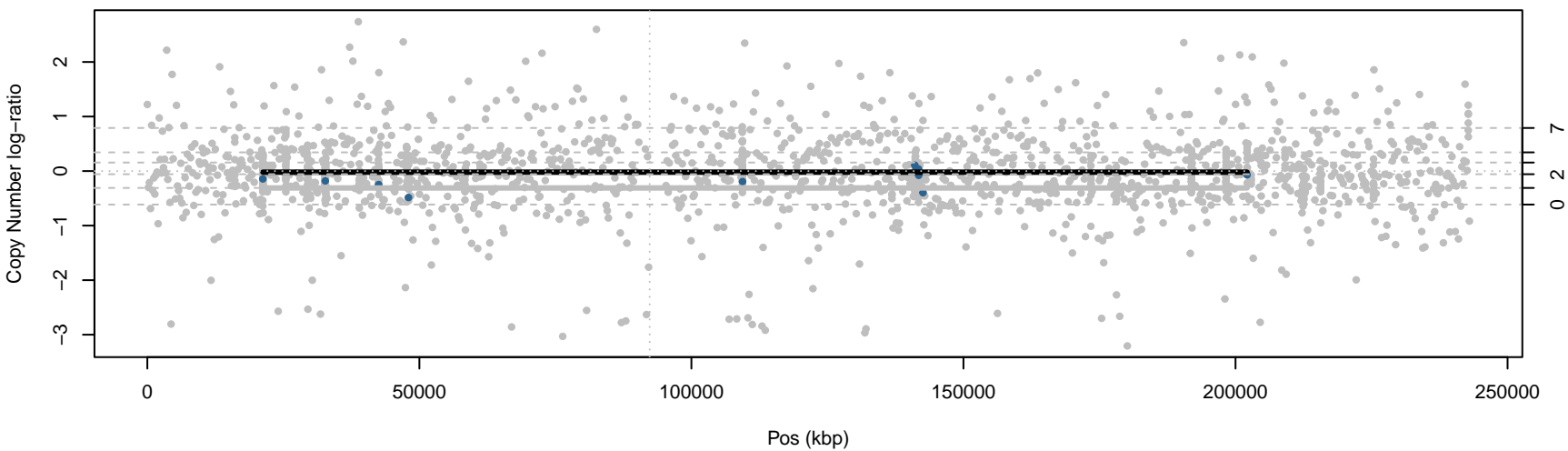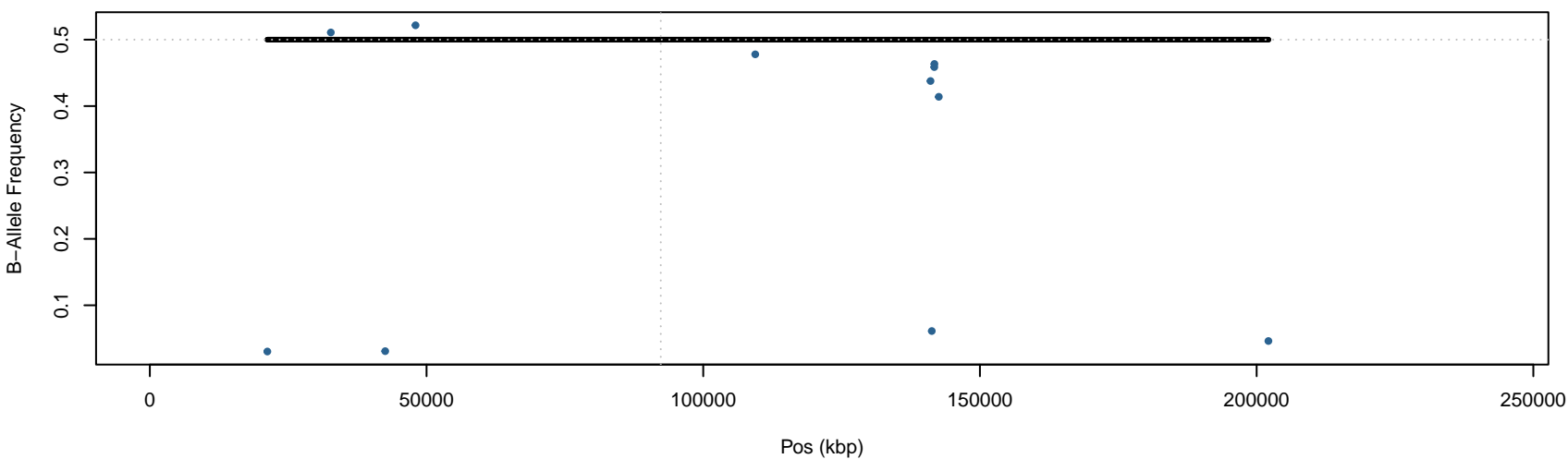
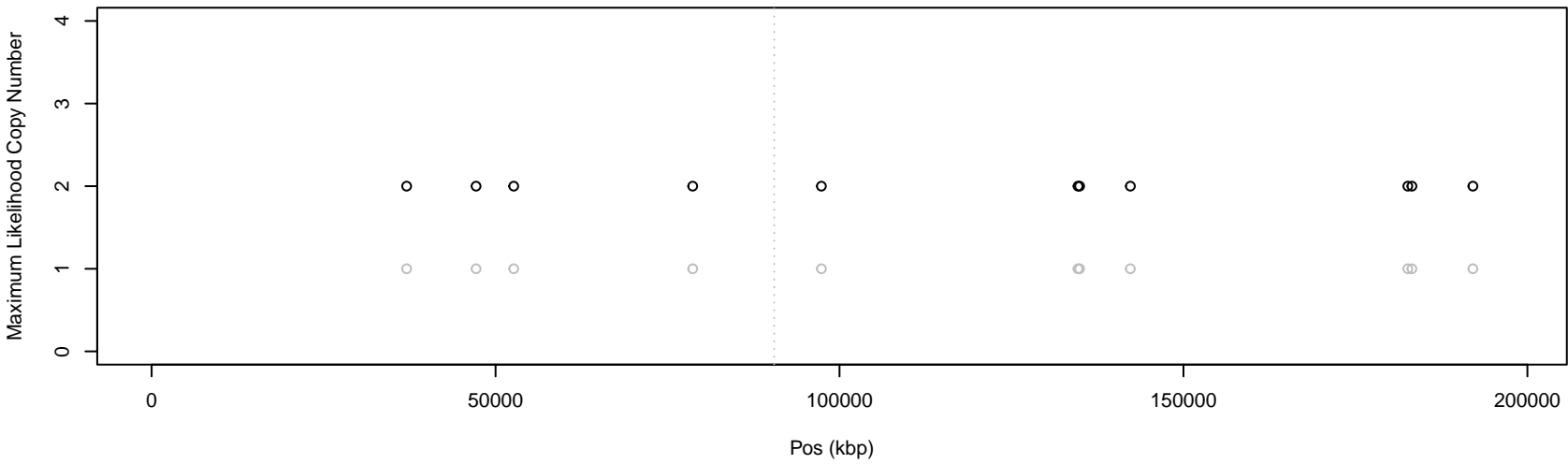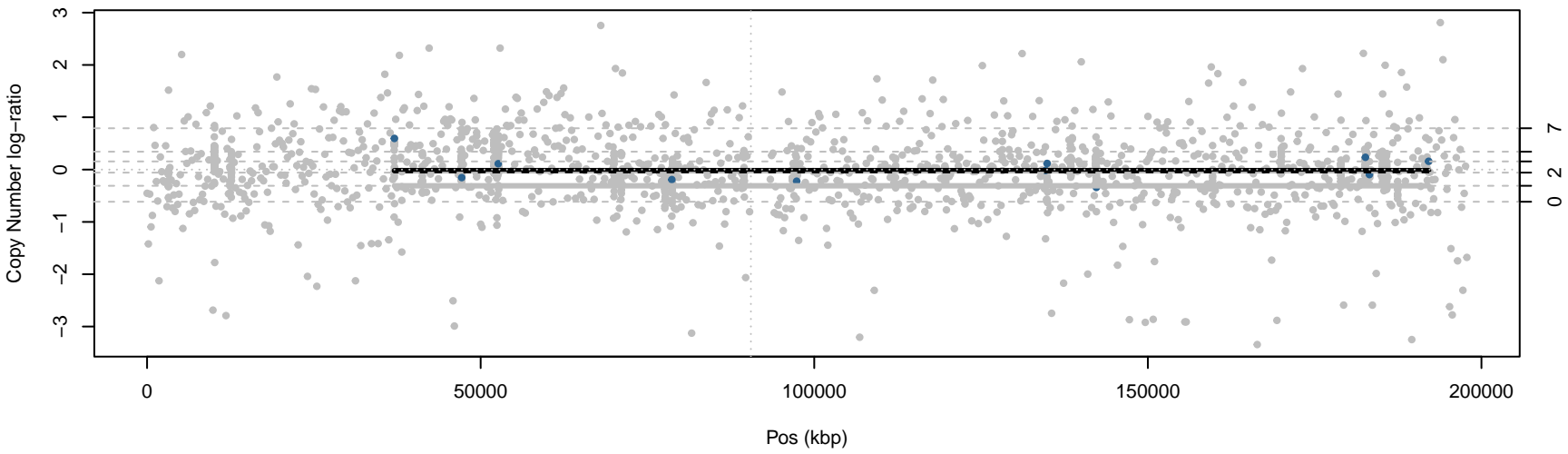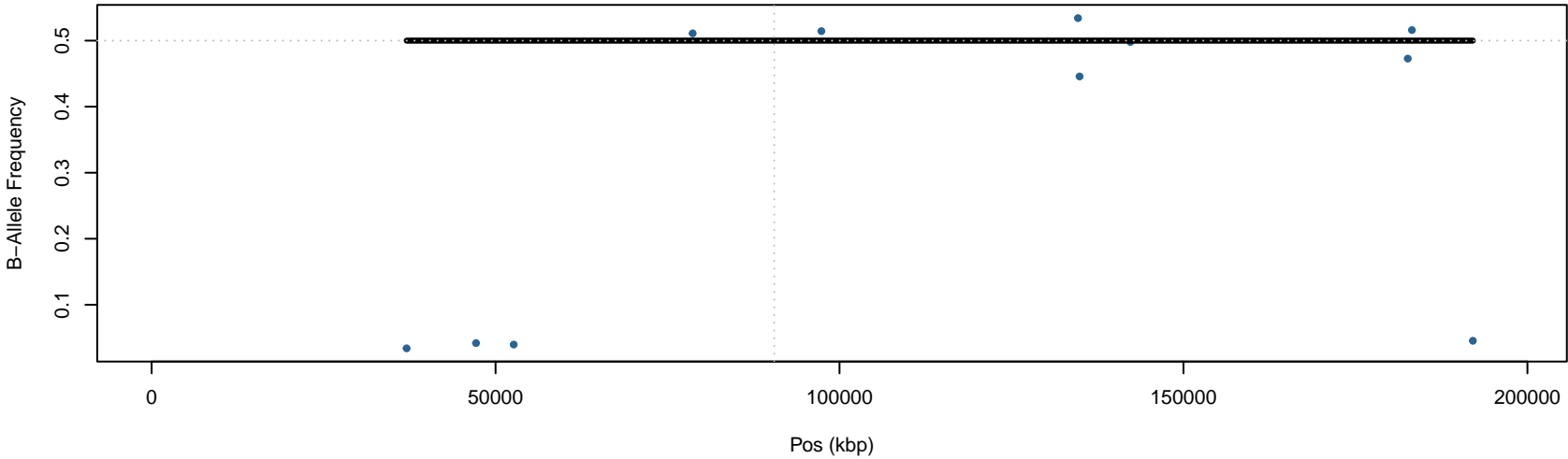**Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 3**
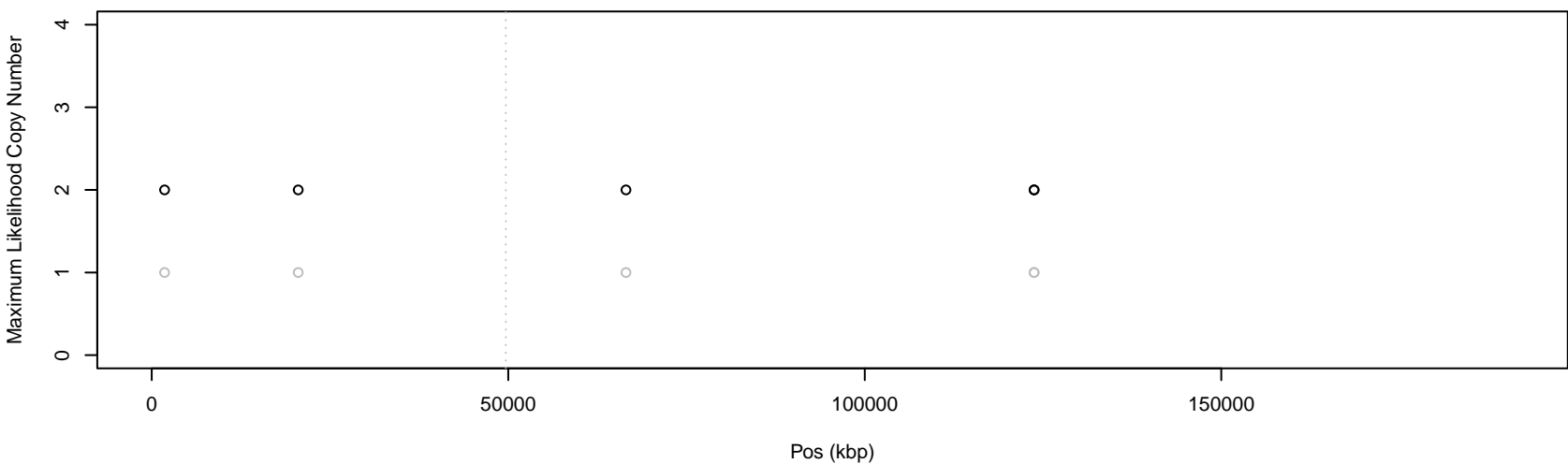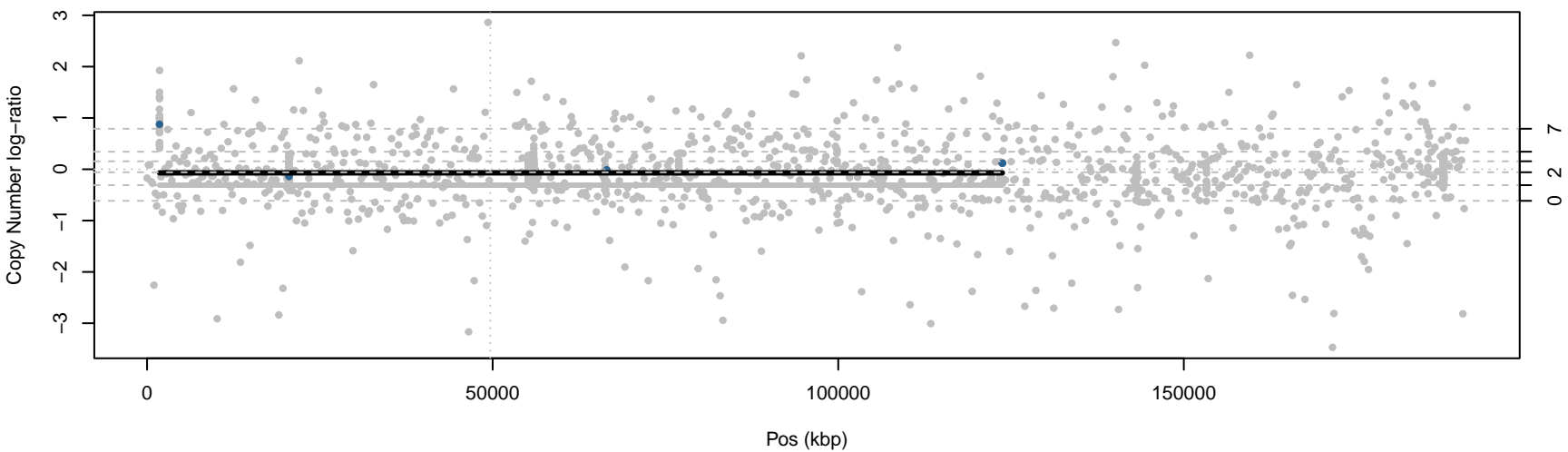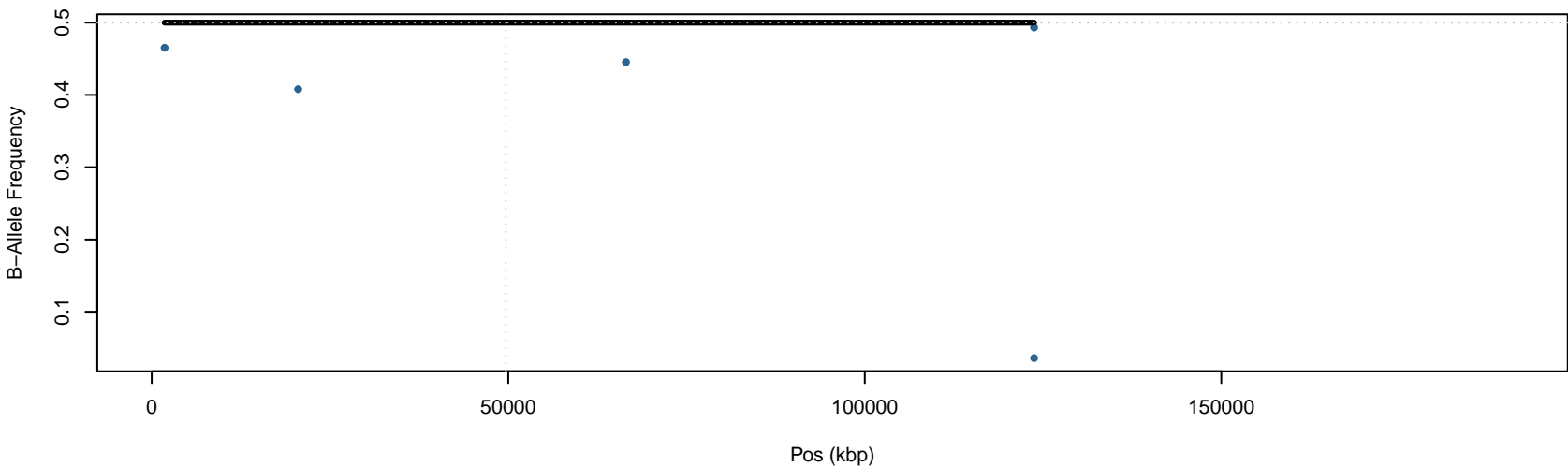
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 4
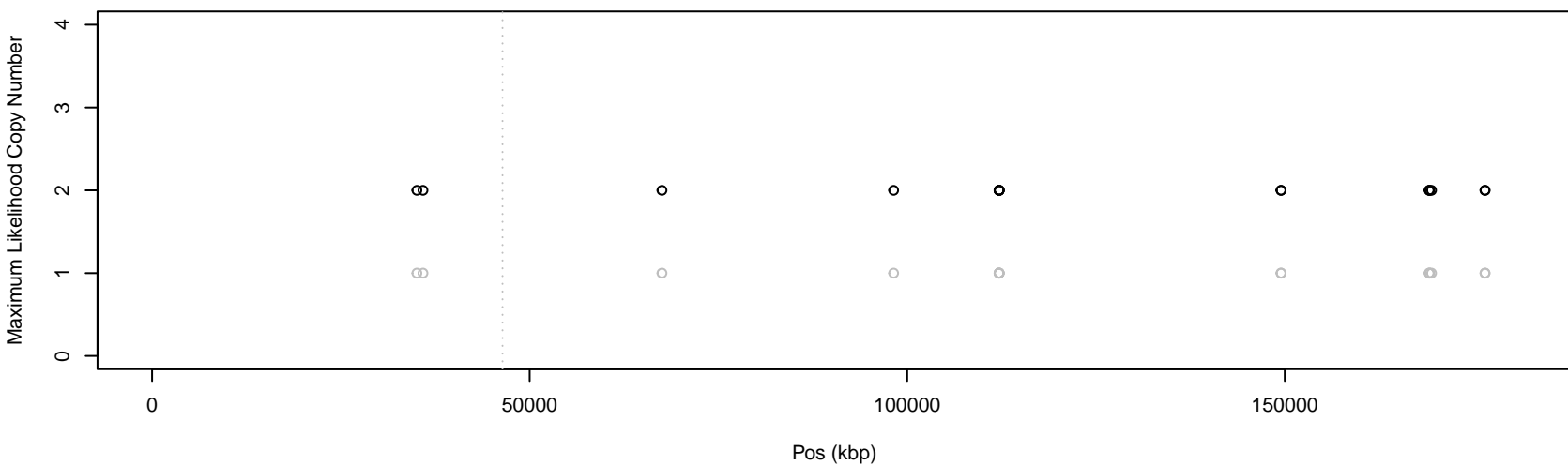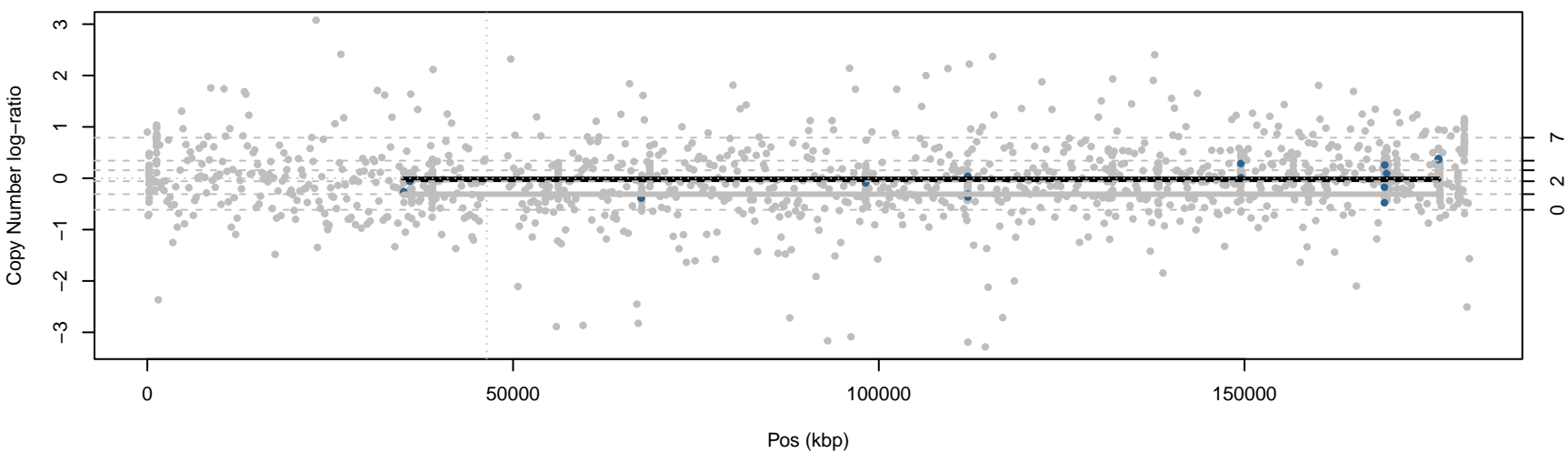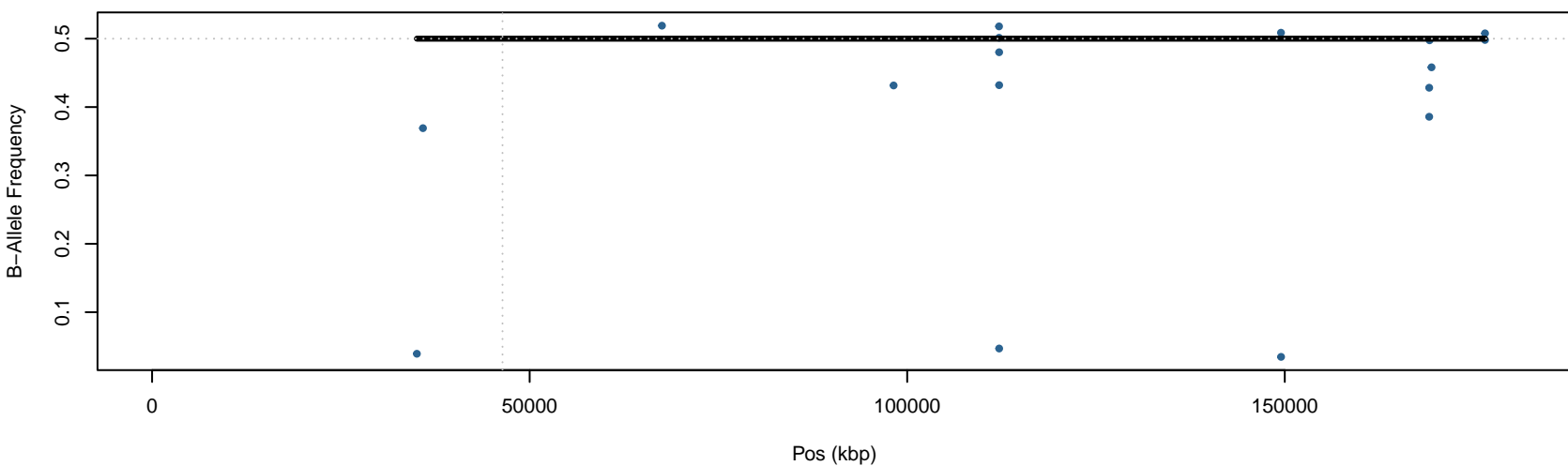
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 5

Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 6

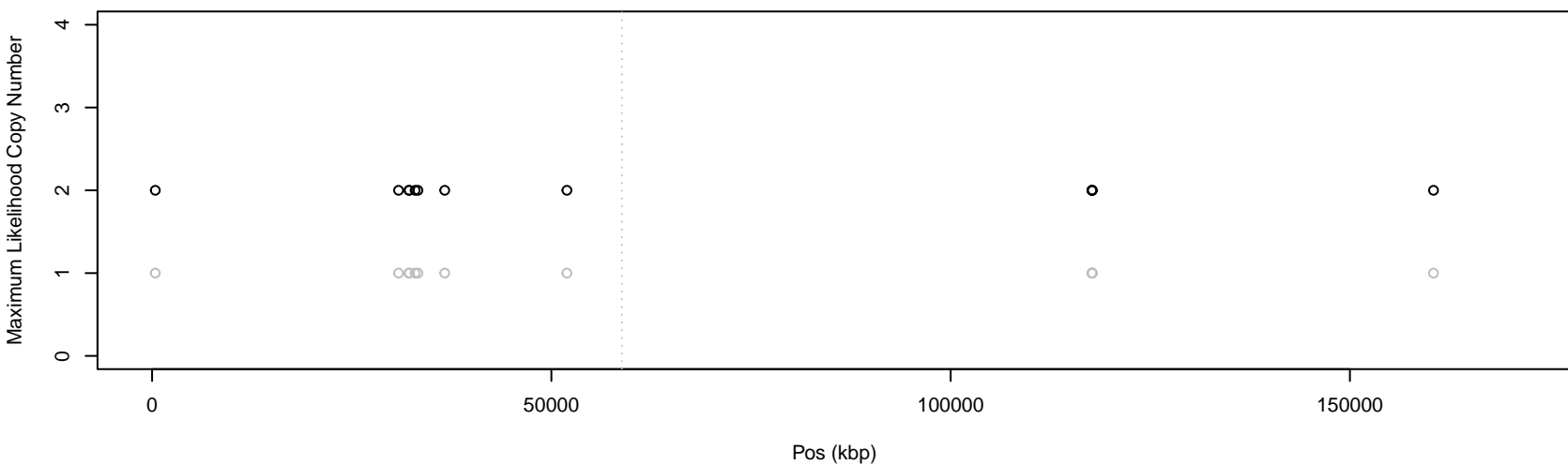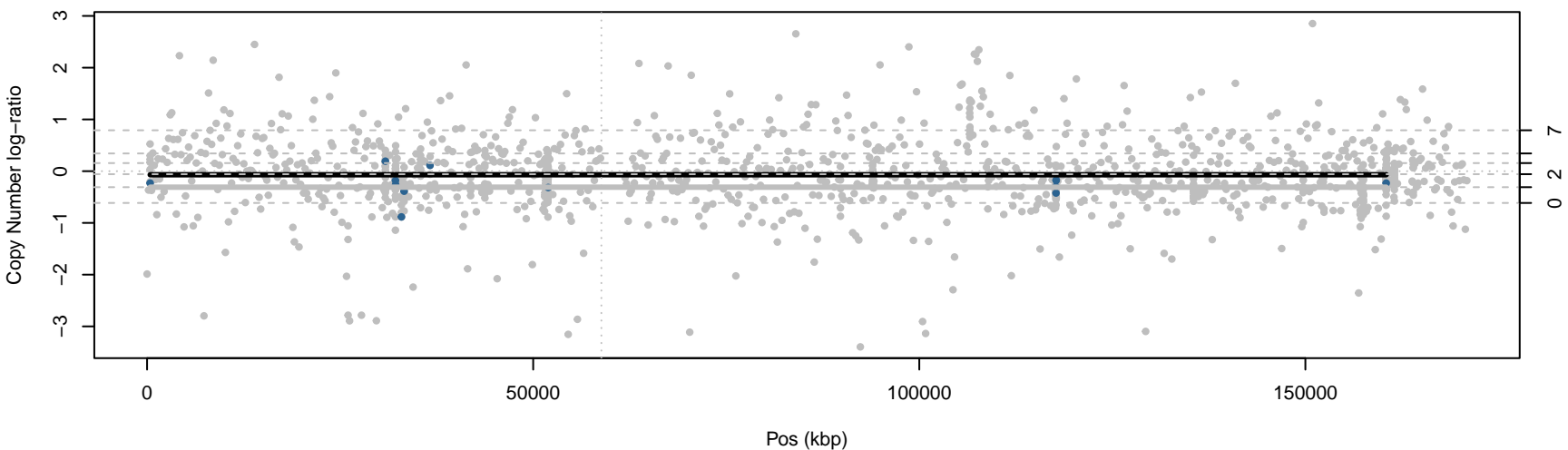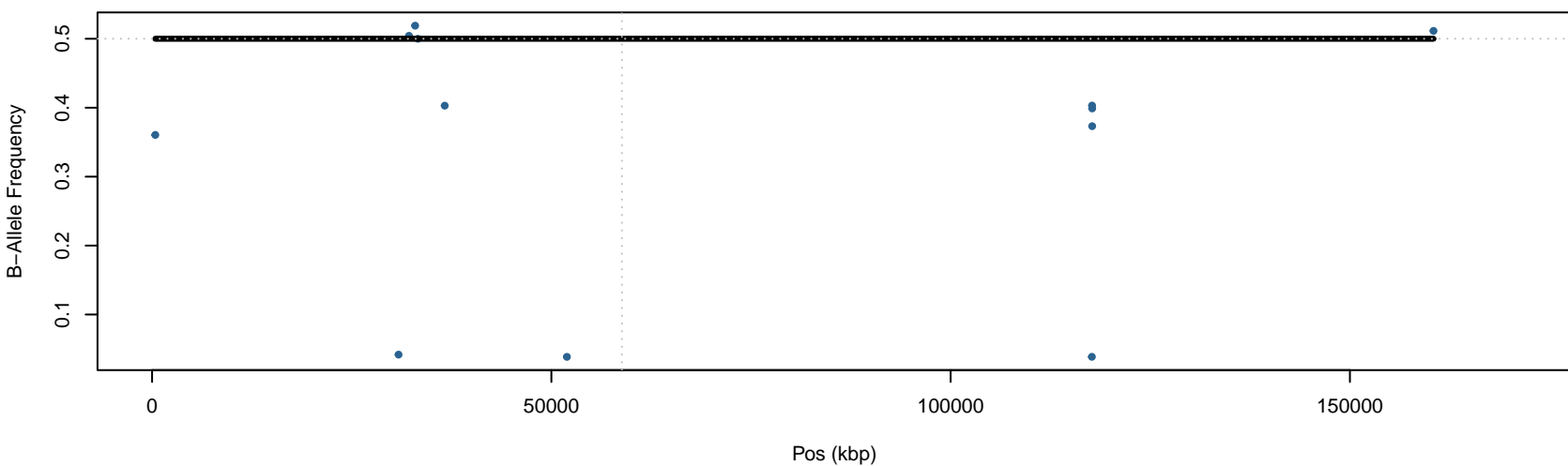**Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 7**

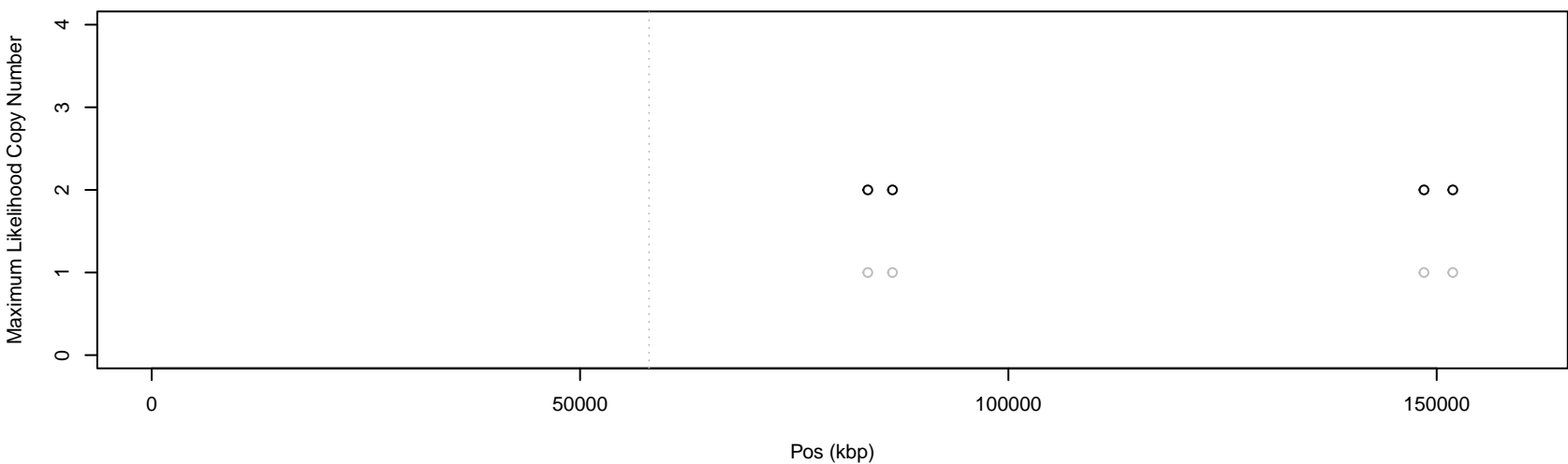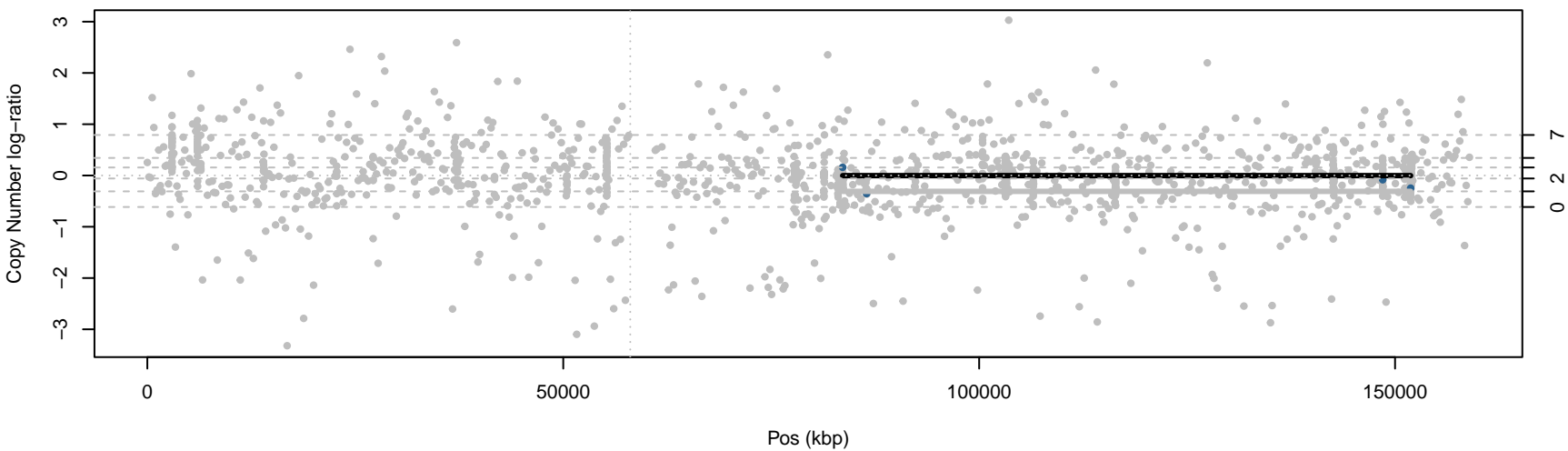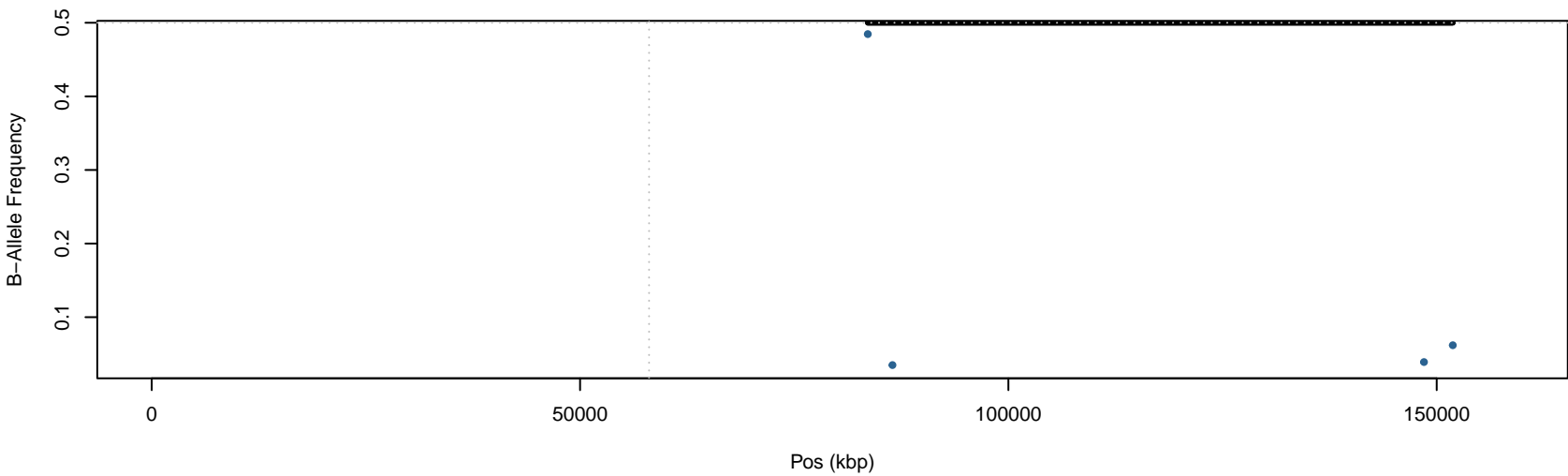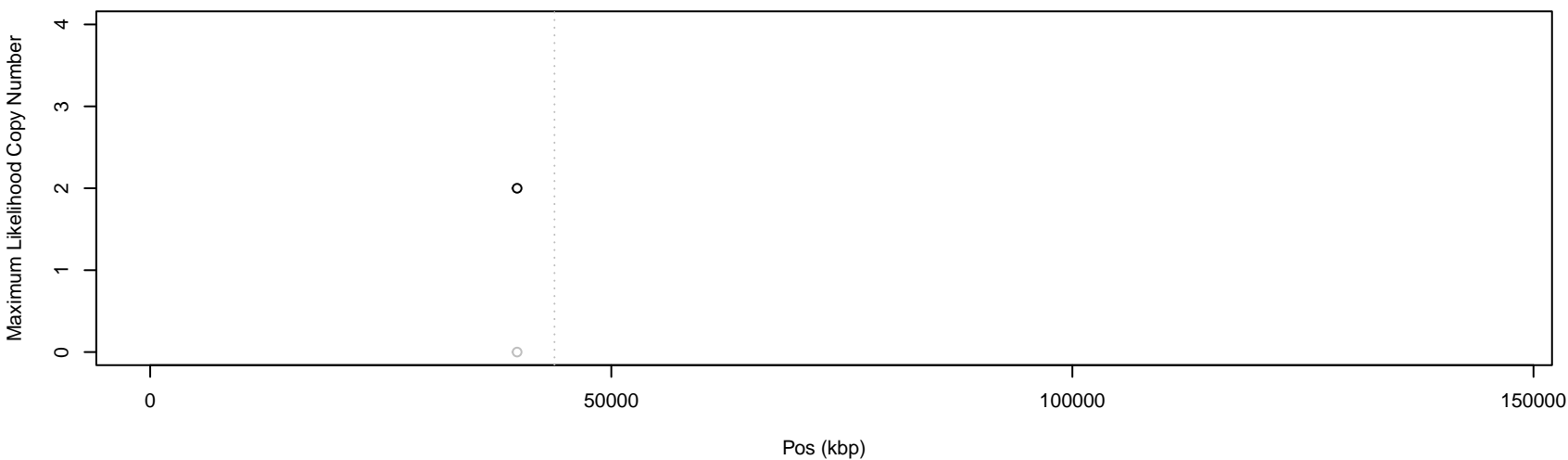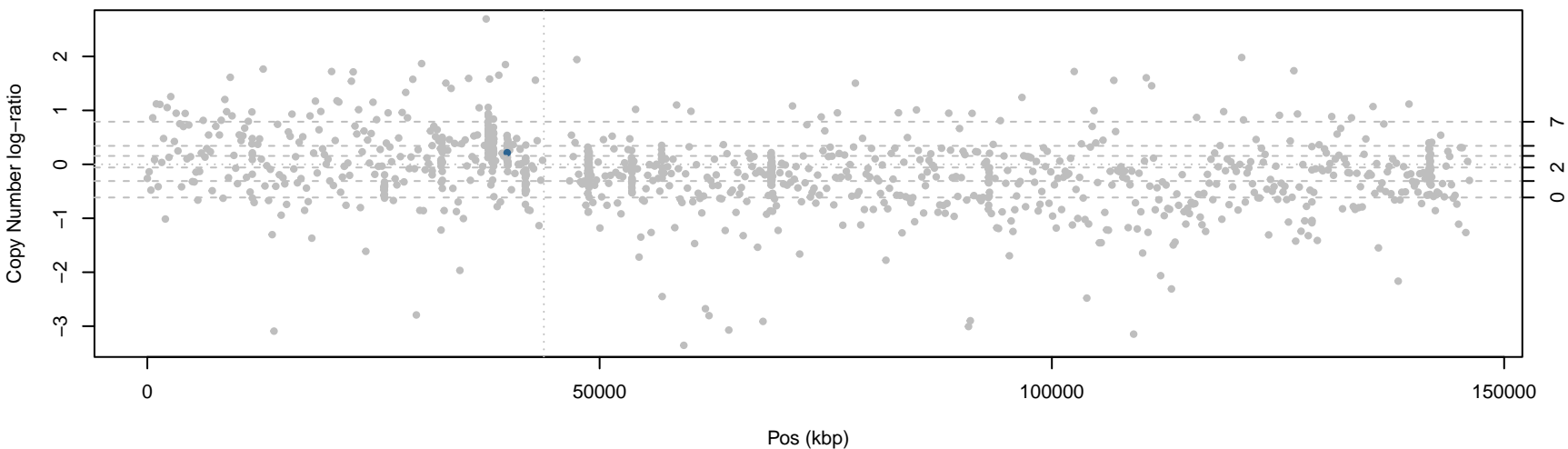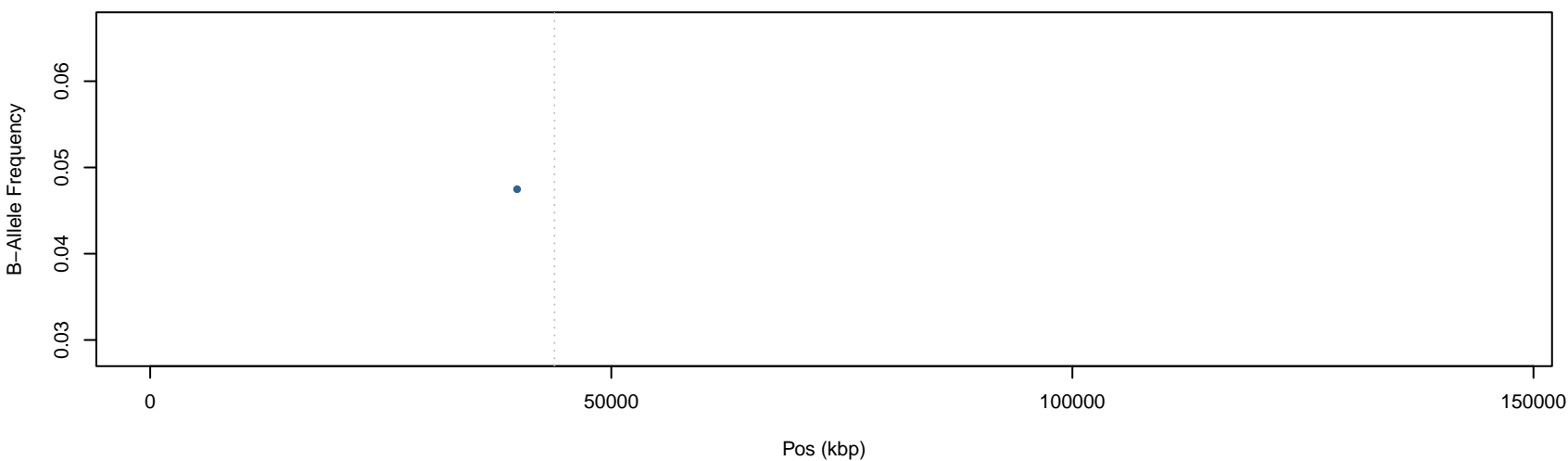**Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 8**

Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 9
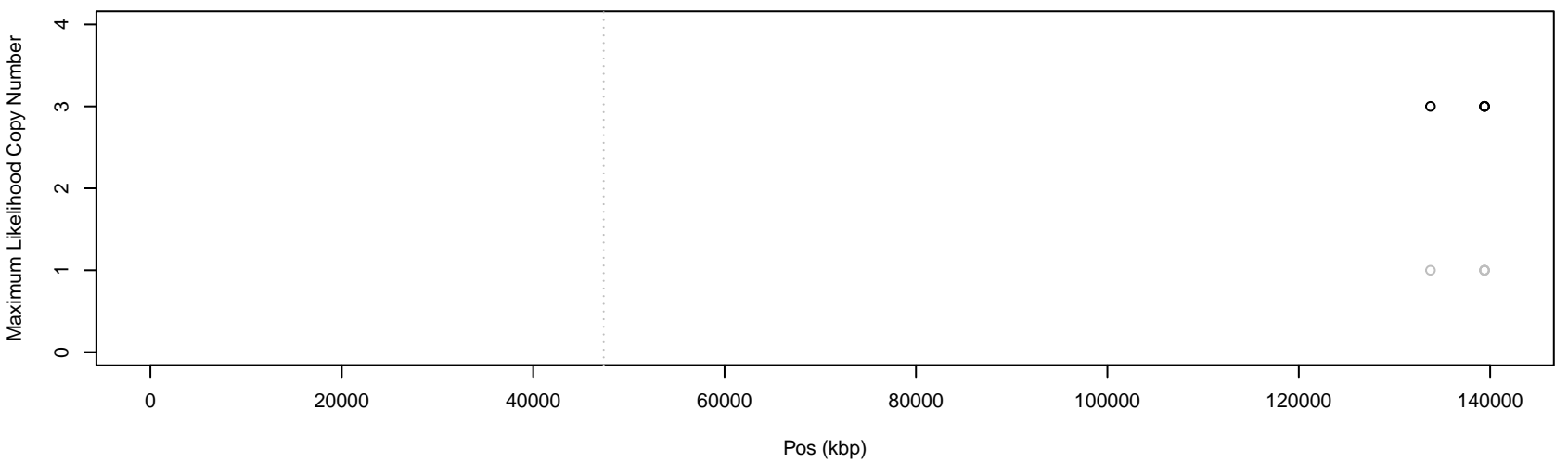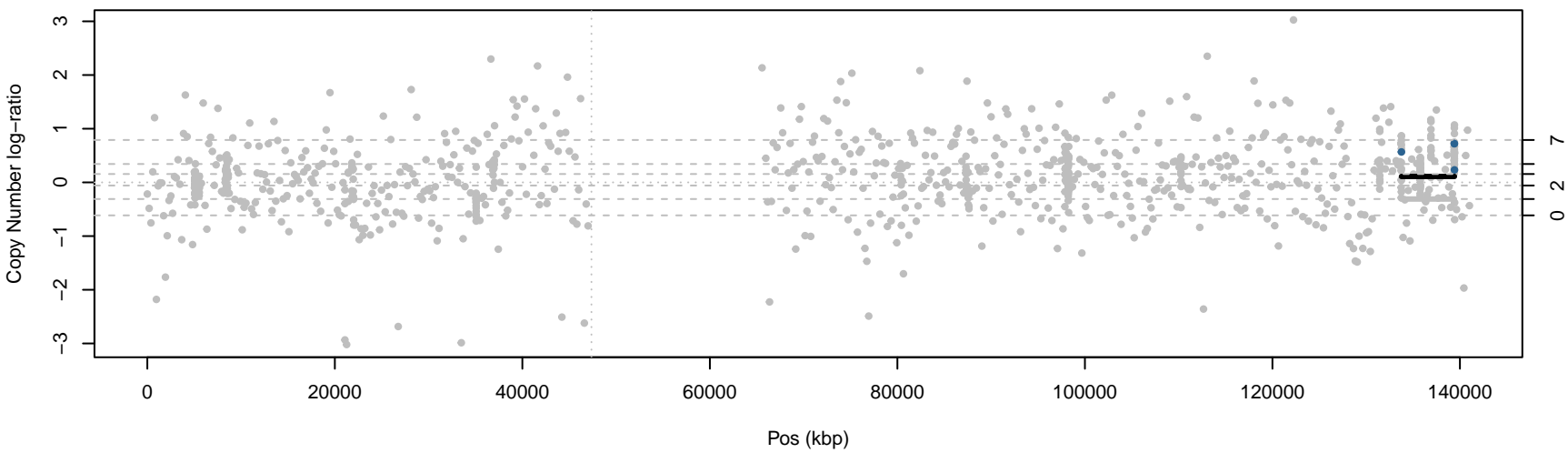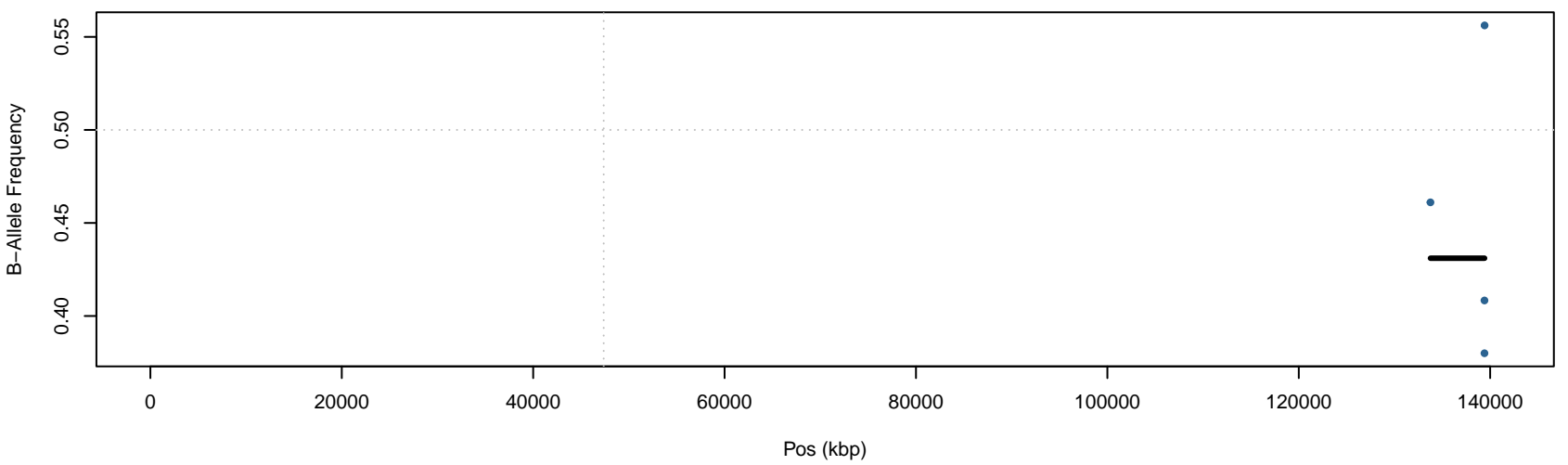
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 10
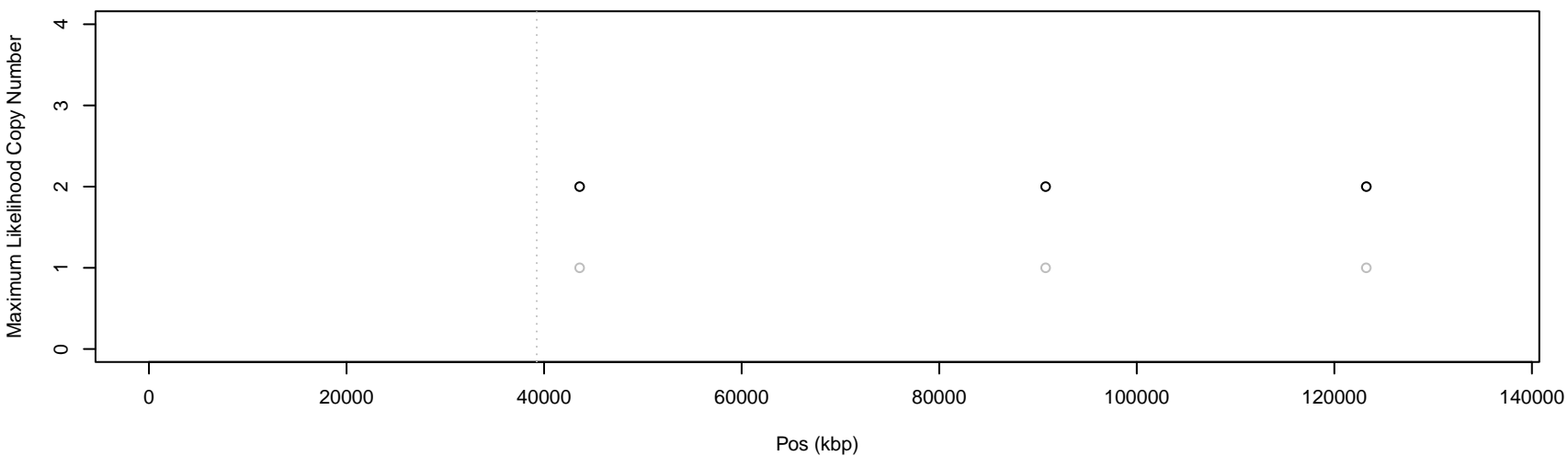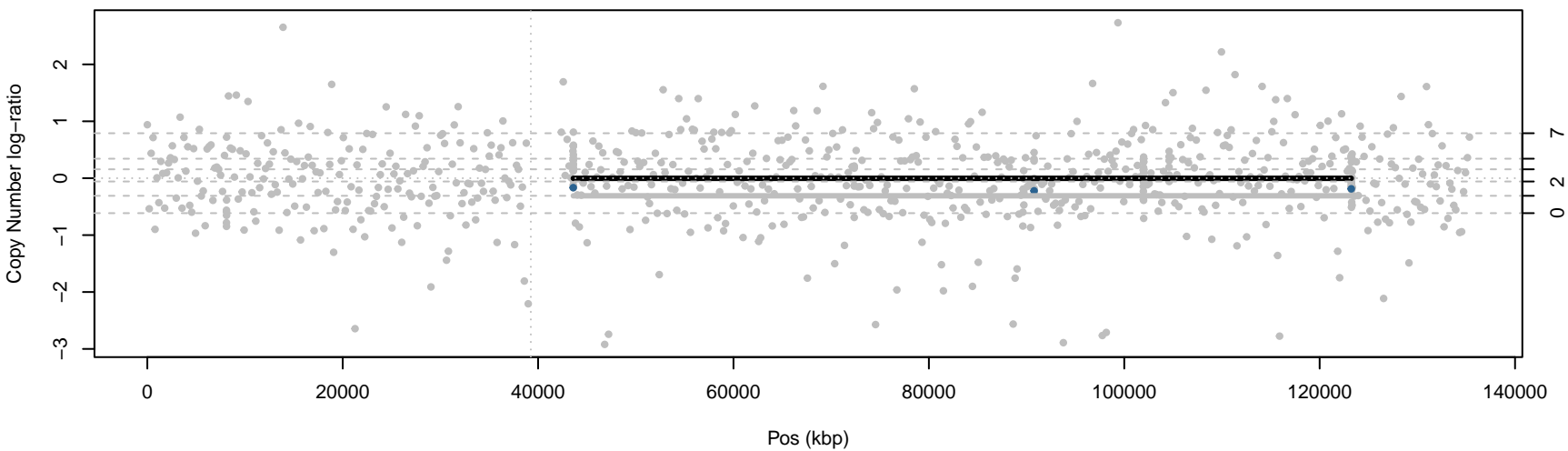
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 12

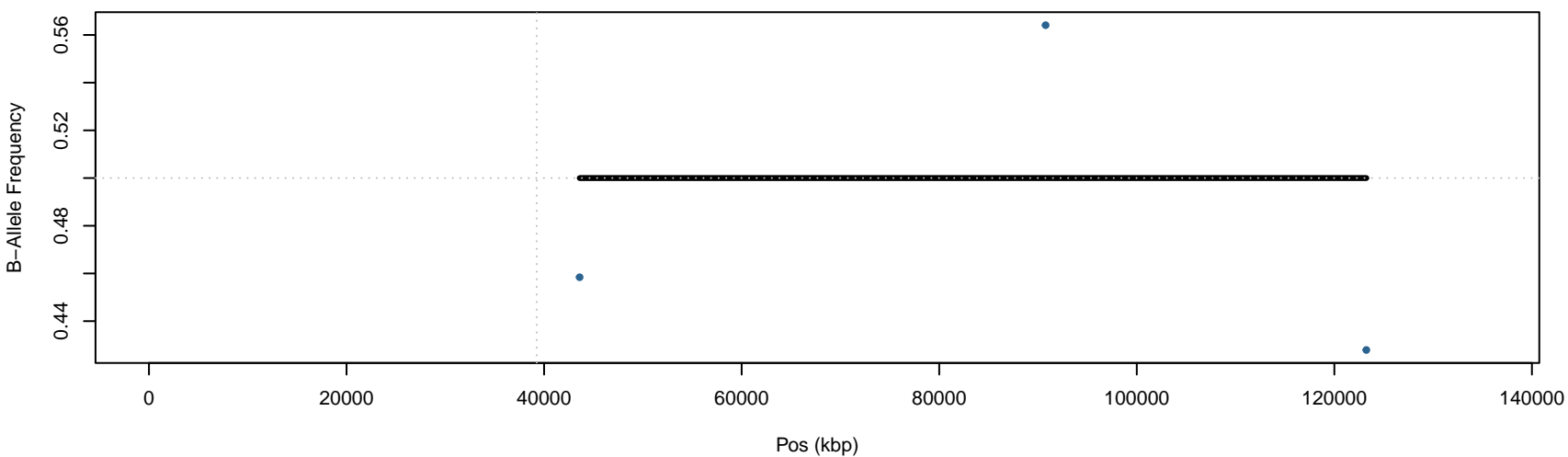**Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 13**
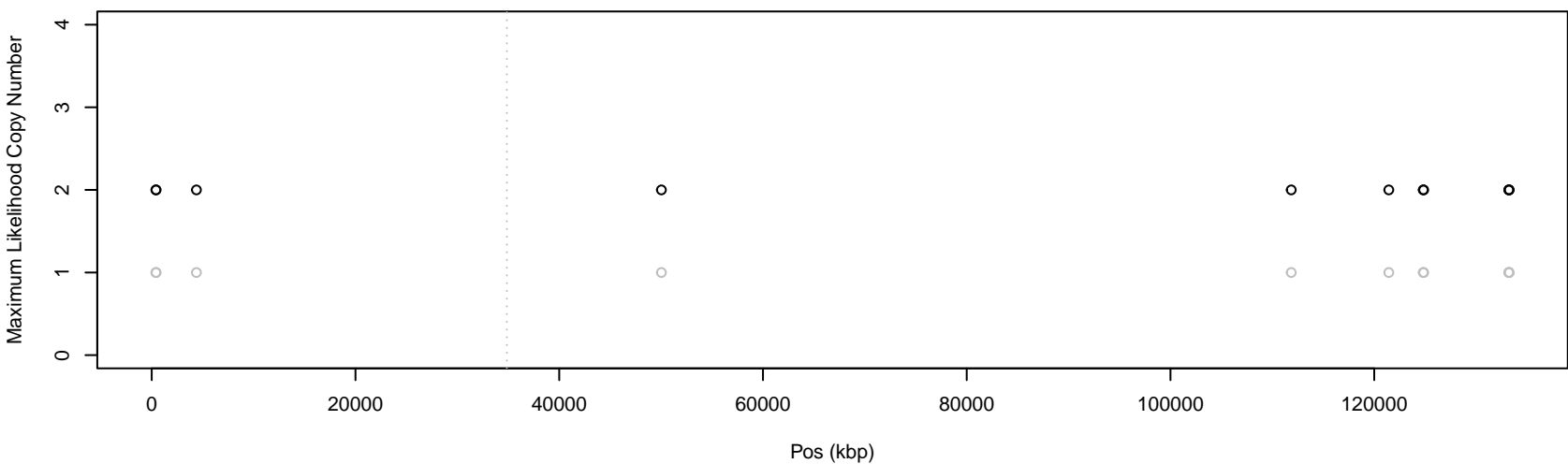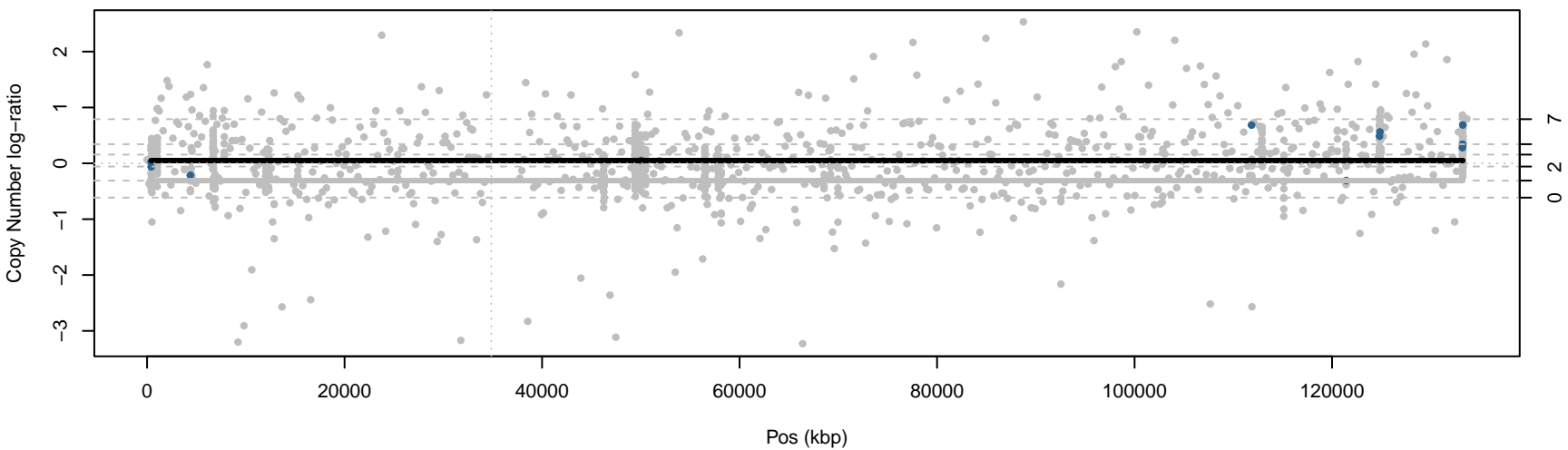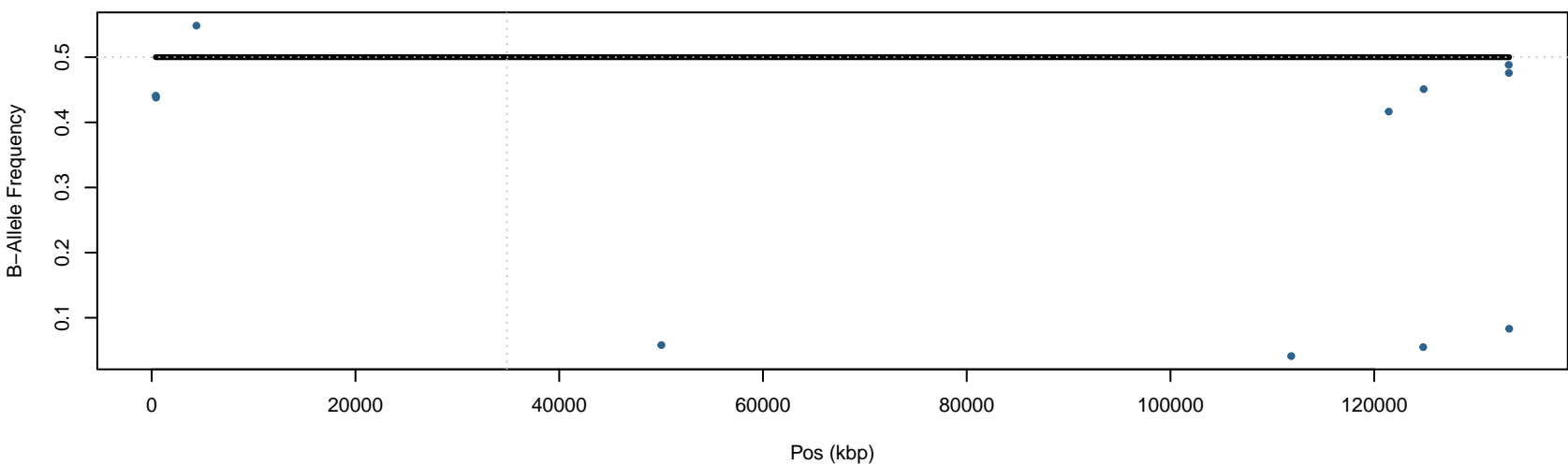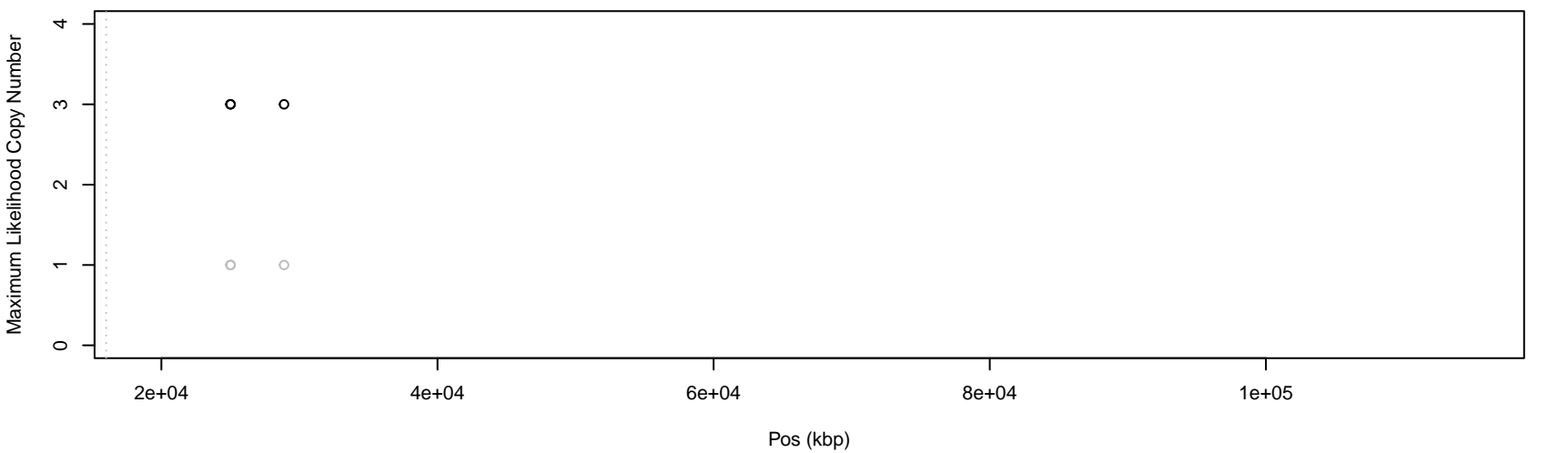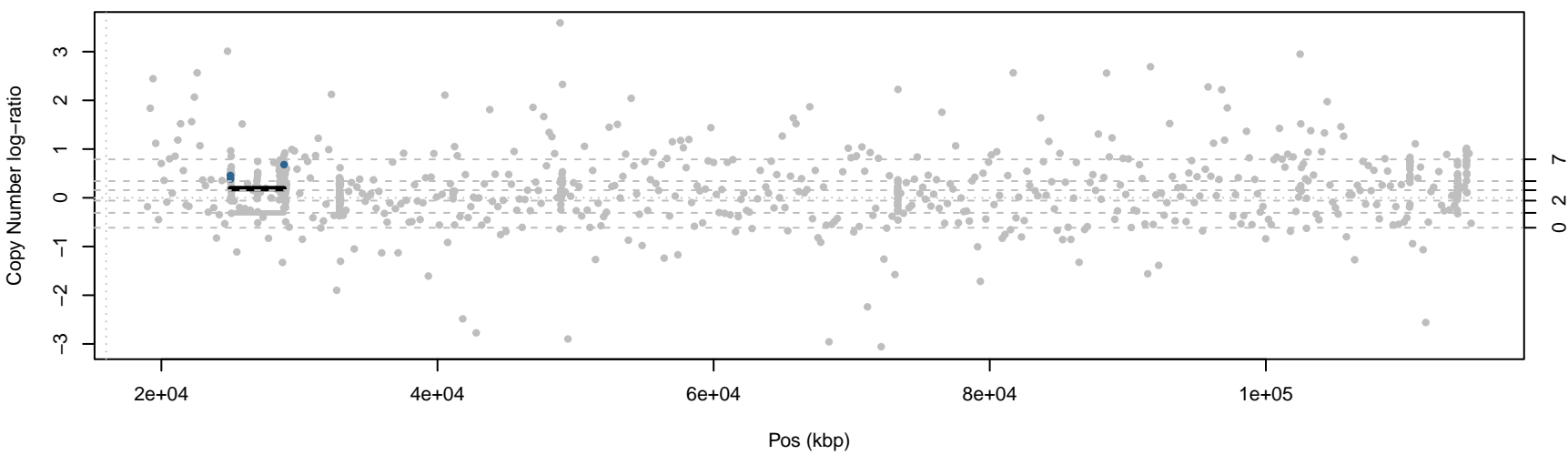
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 14
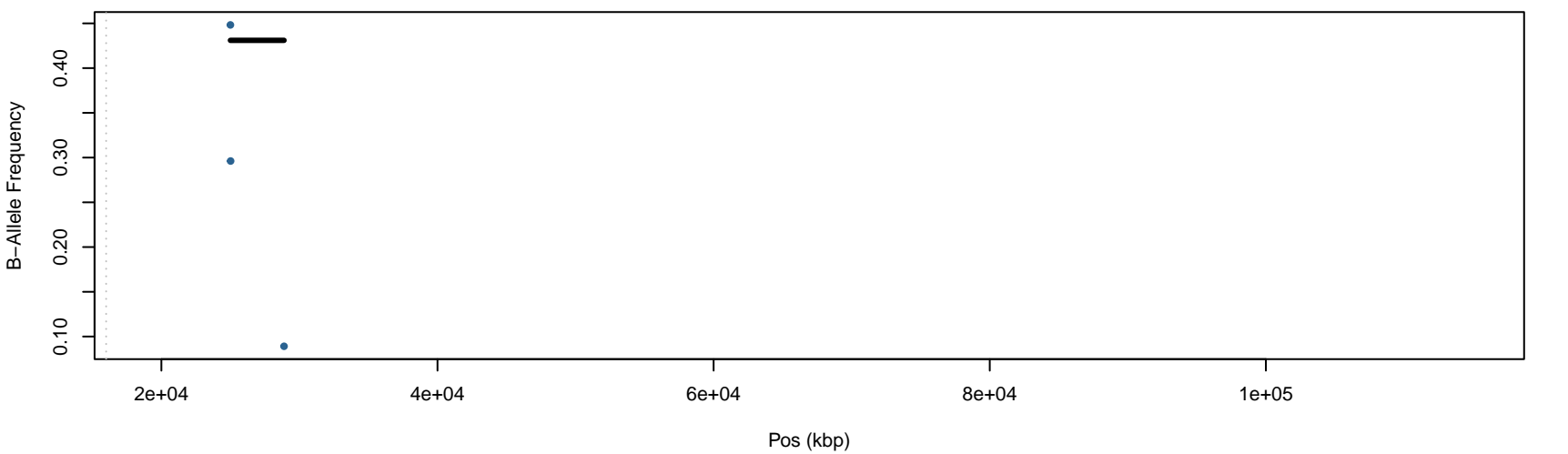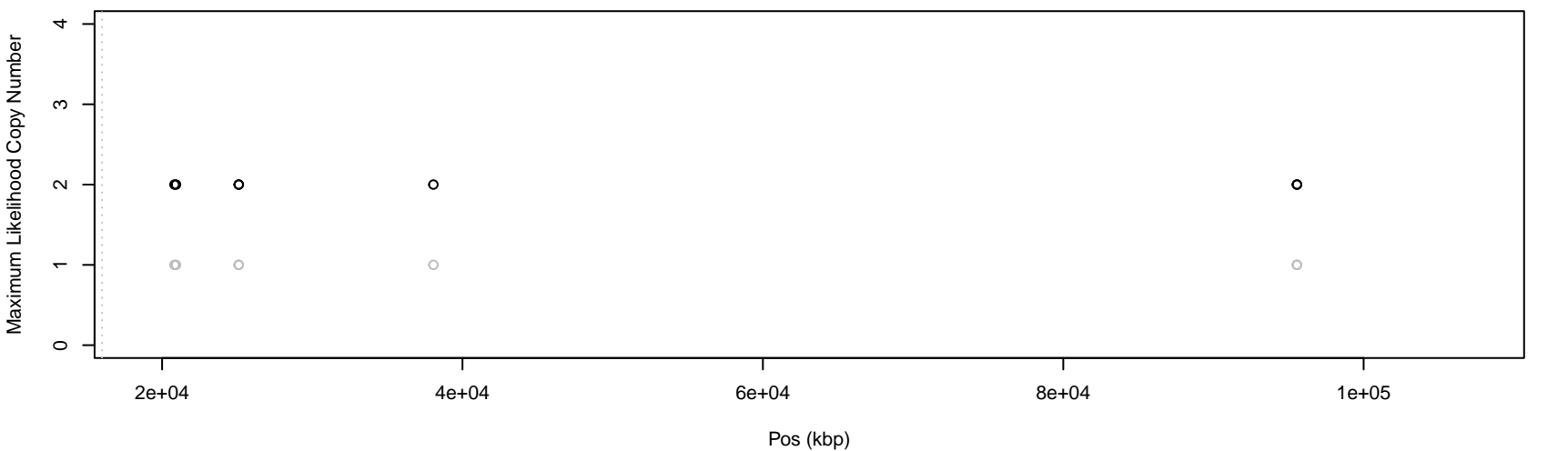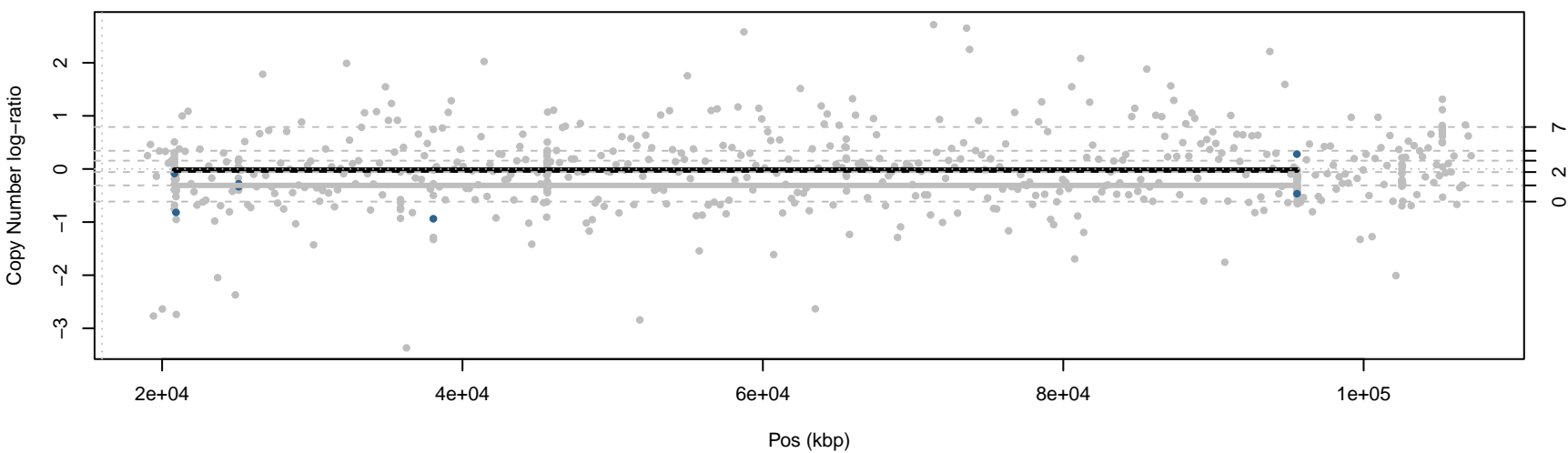
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 15
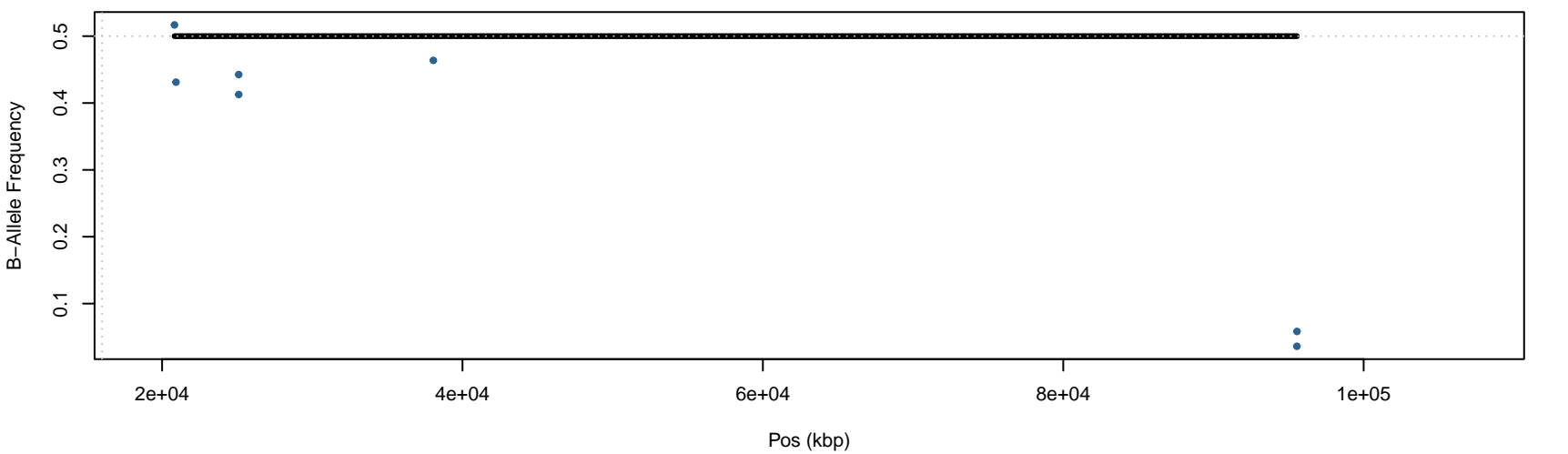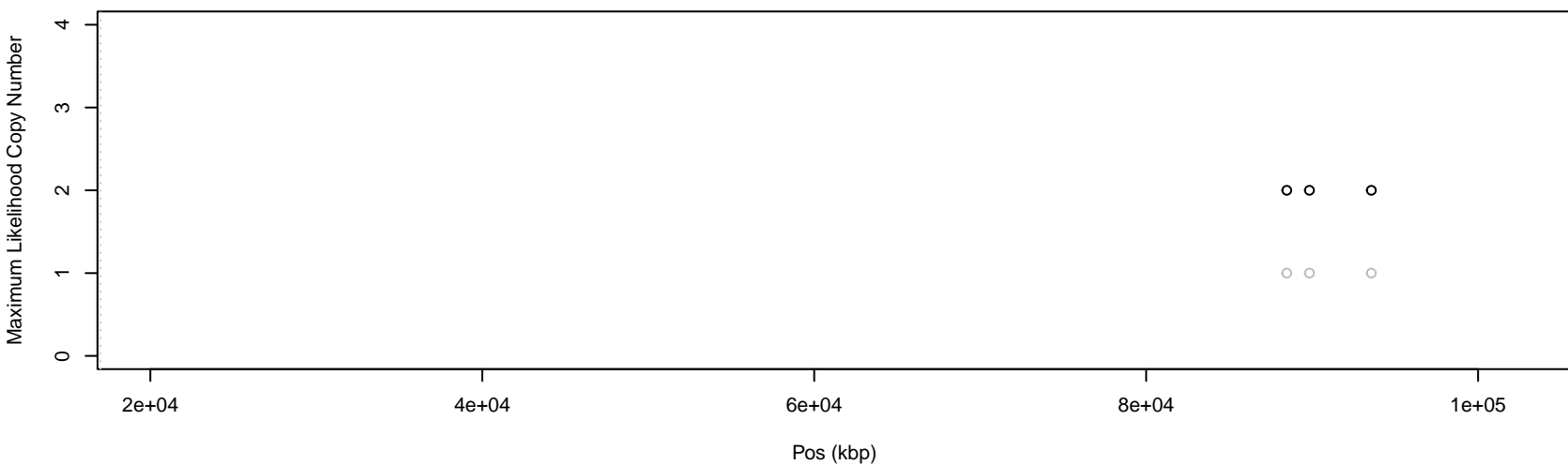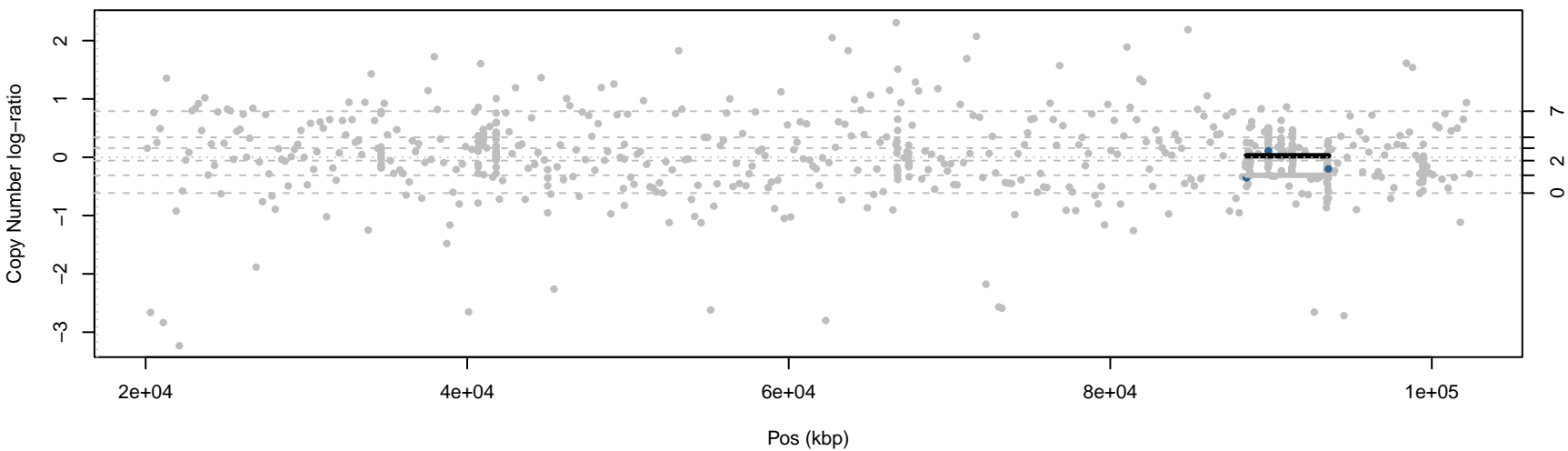
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 16
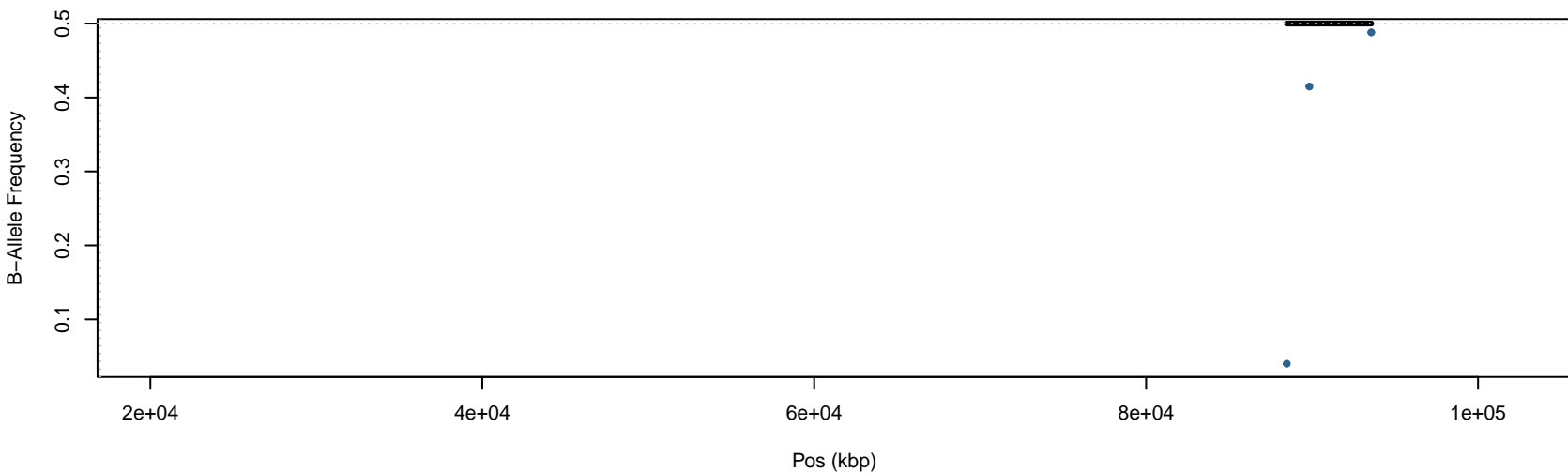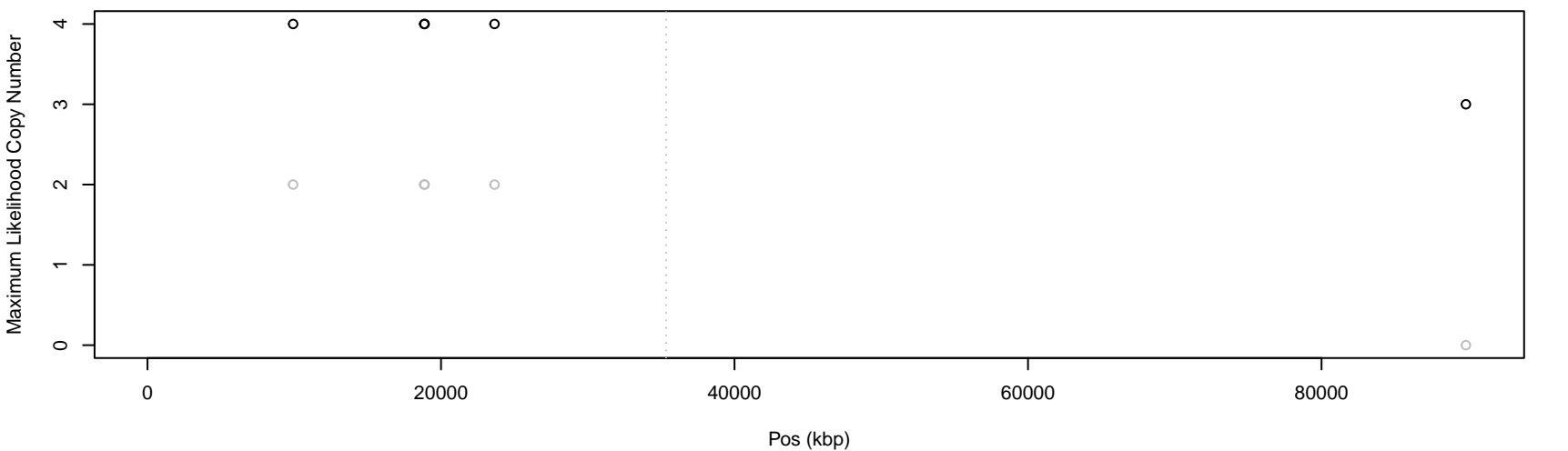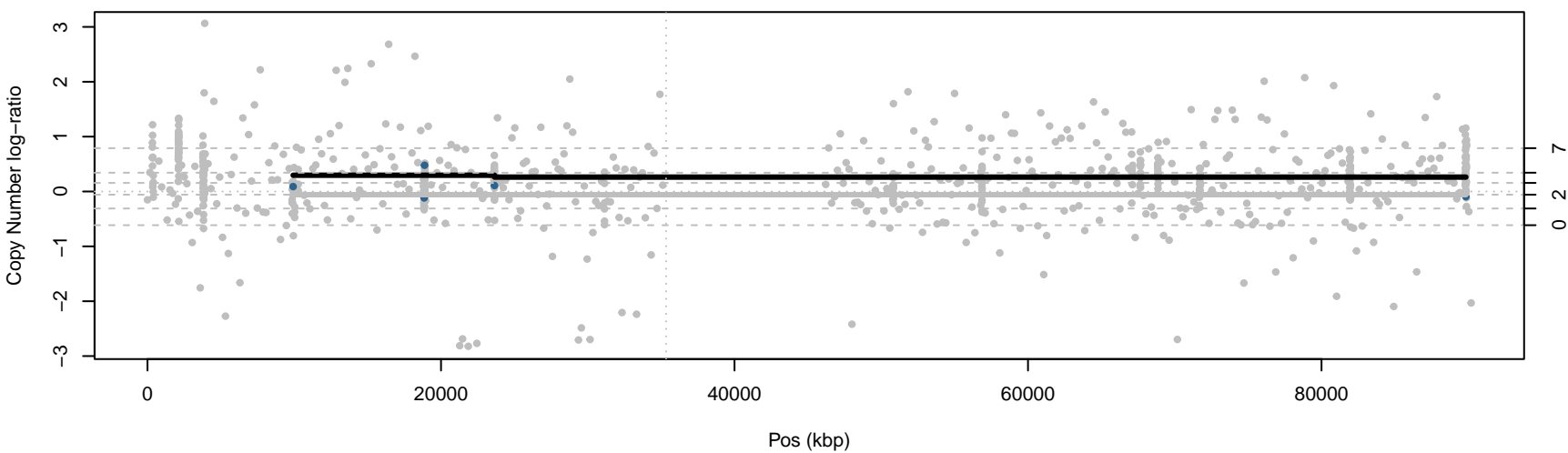
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 18
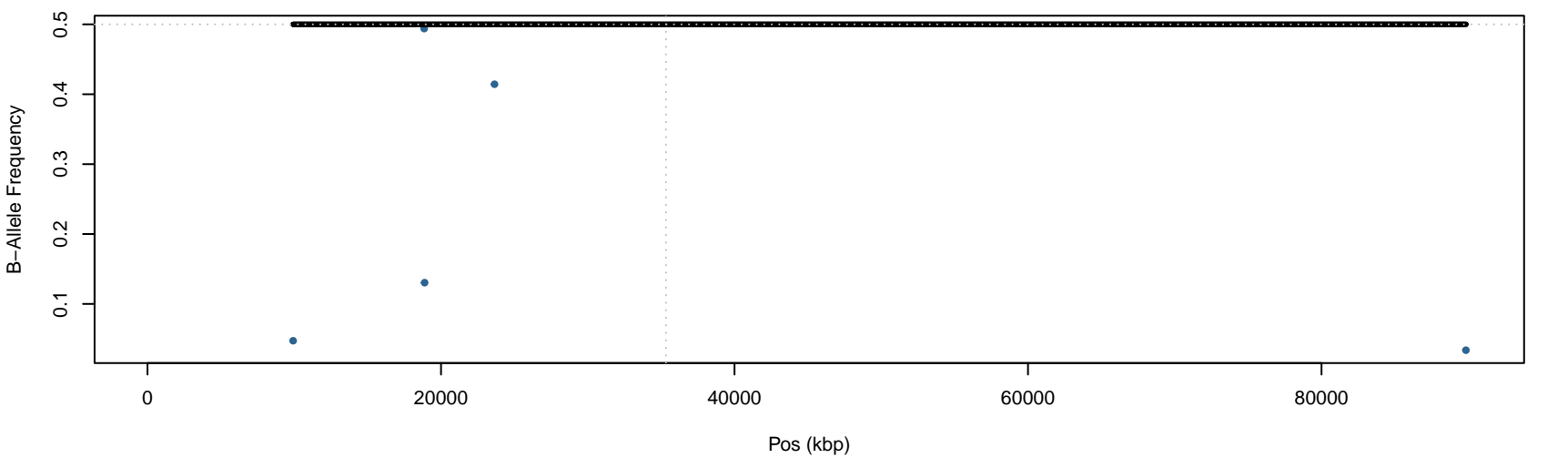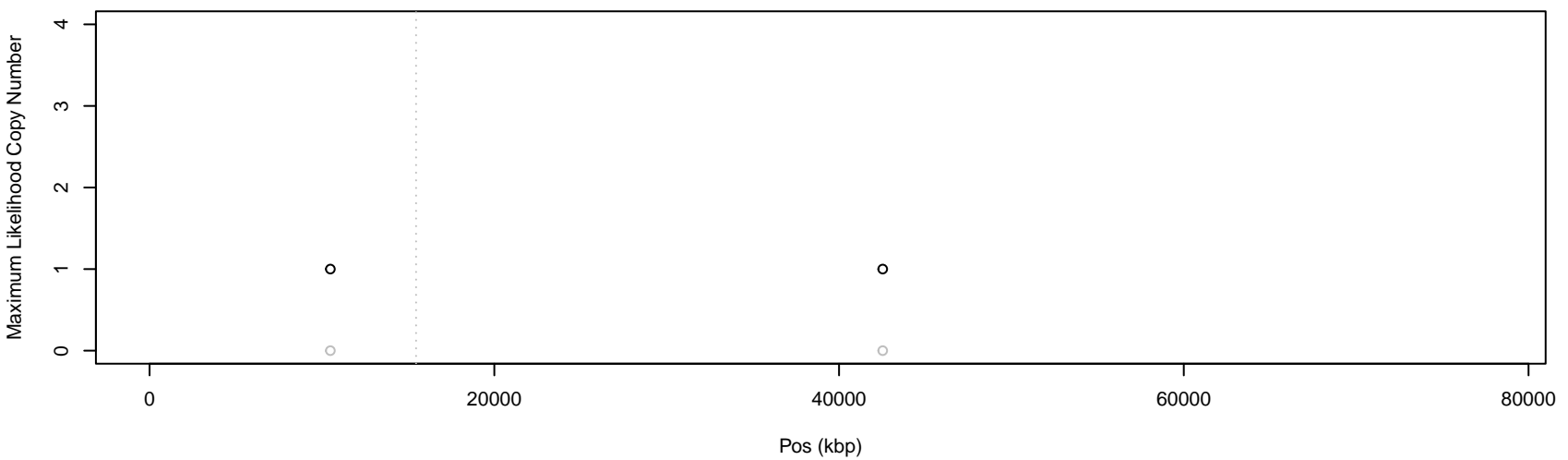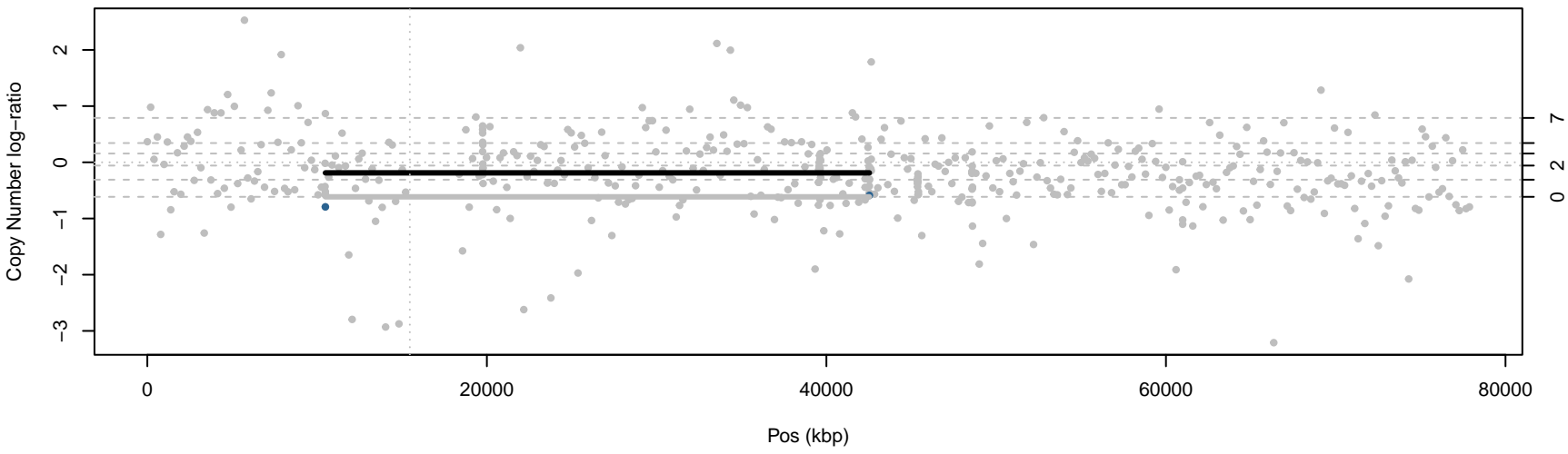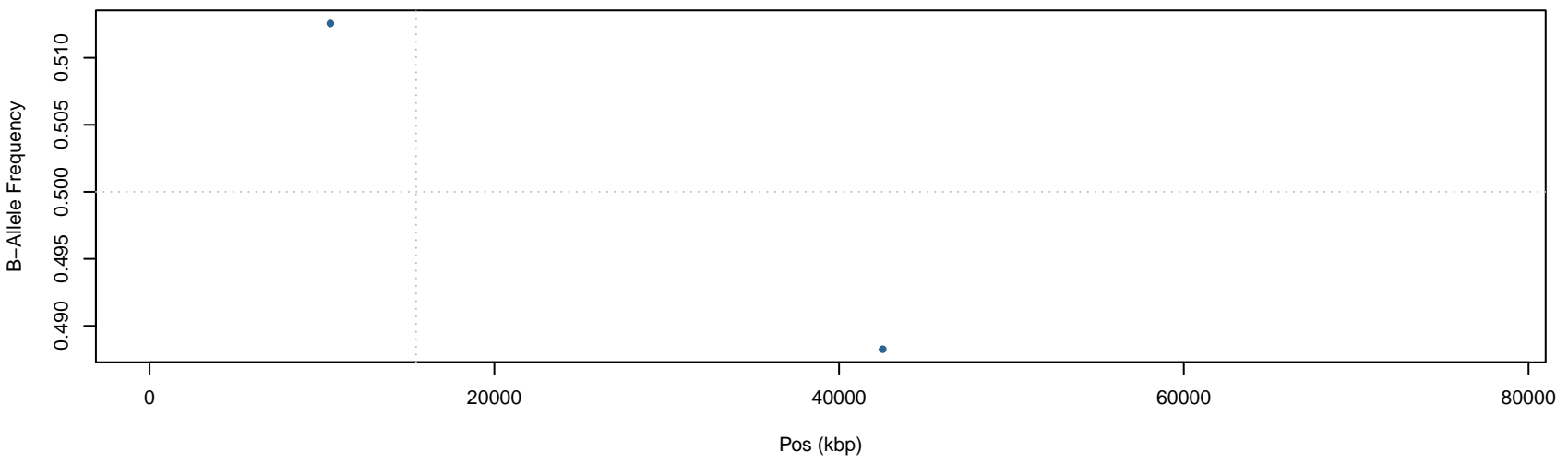
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 19

Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 20
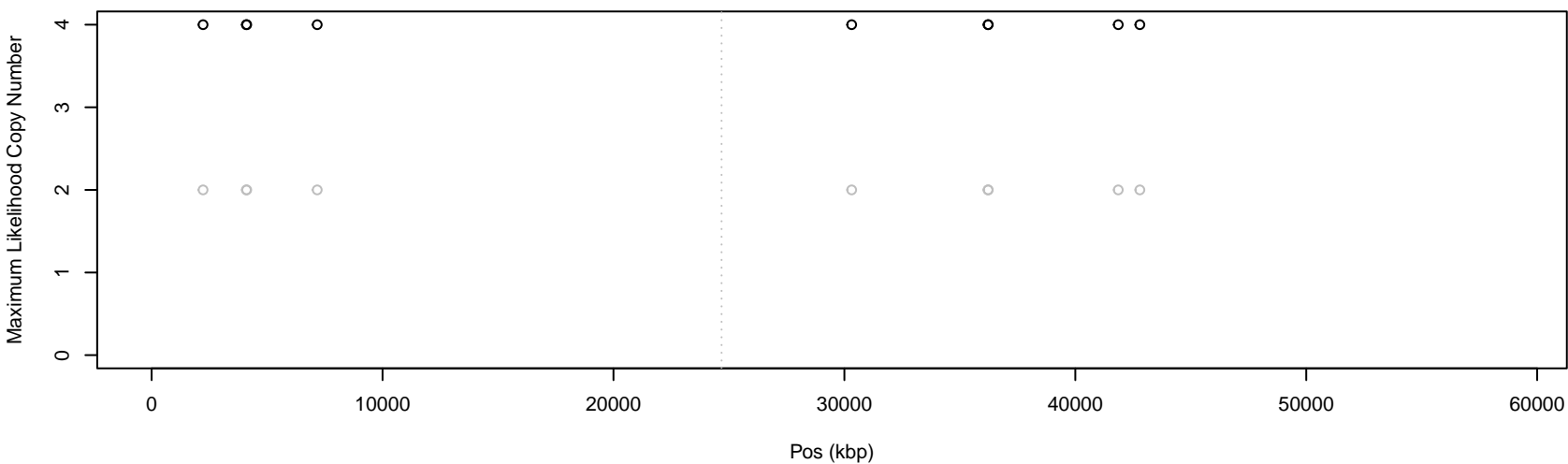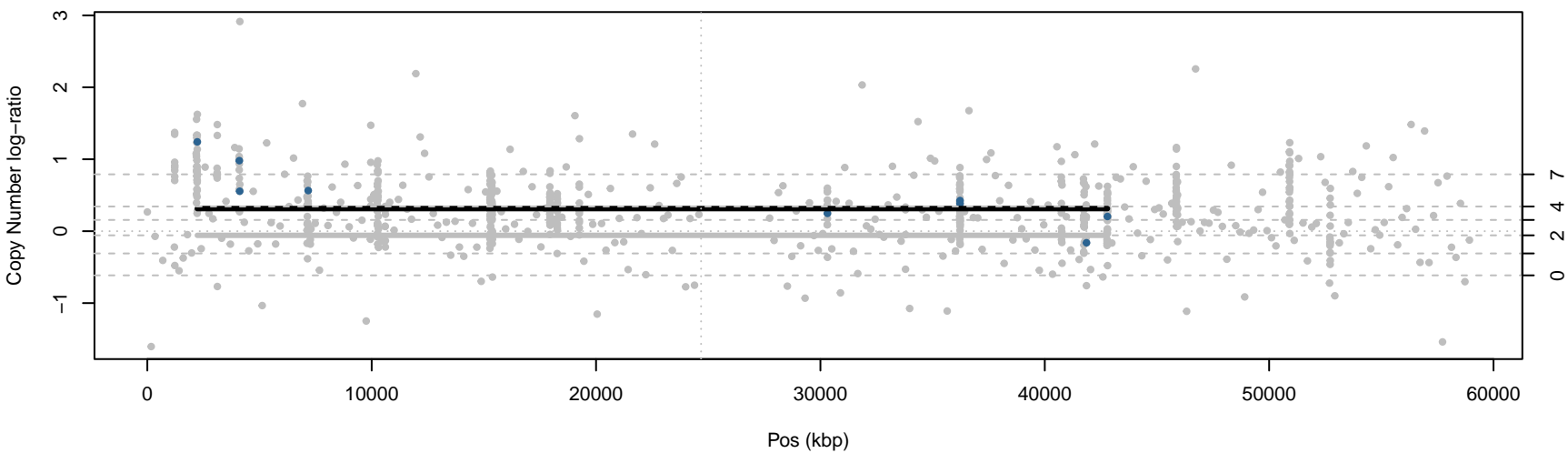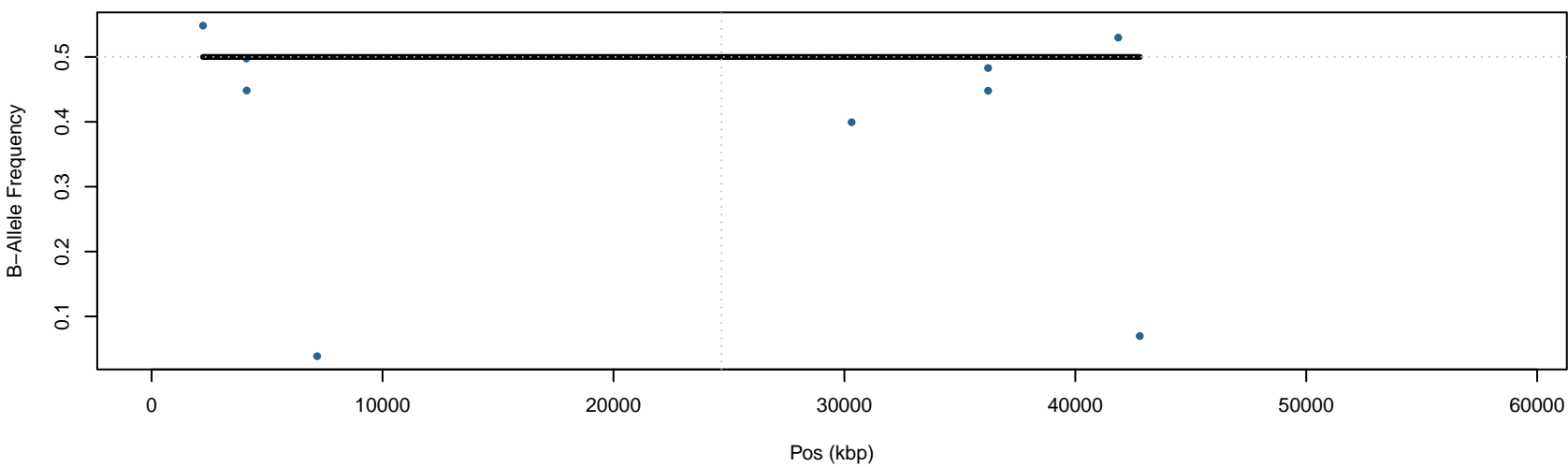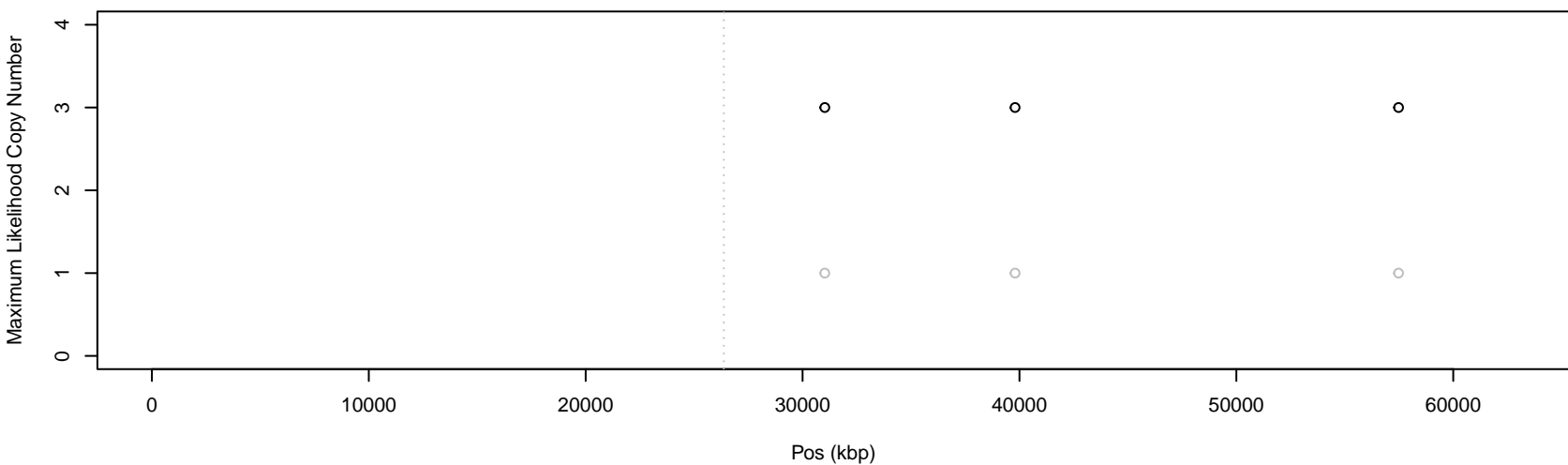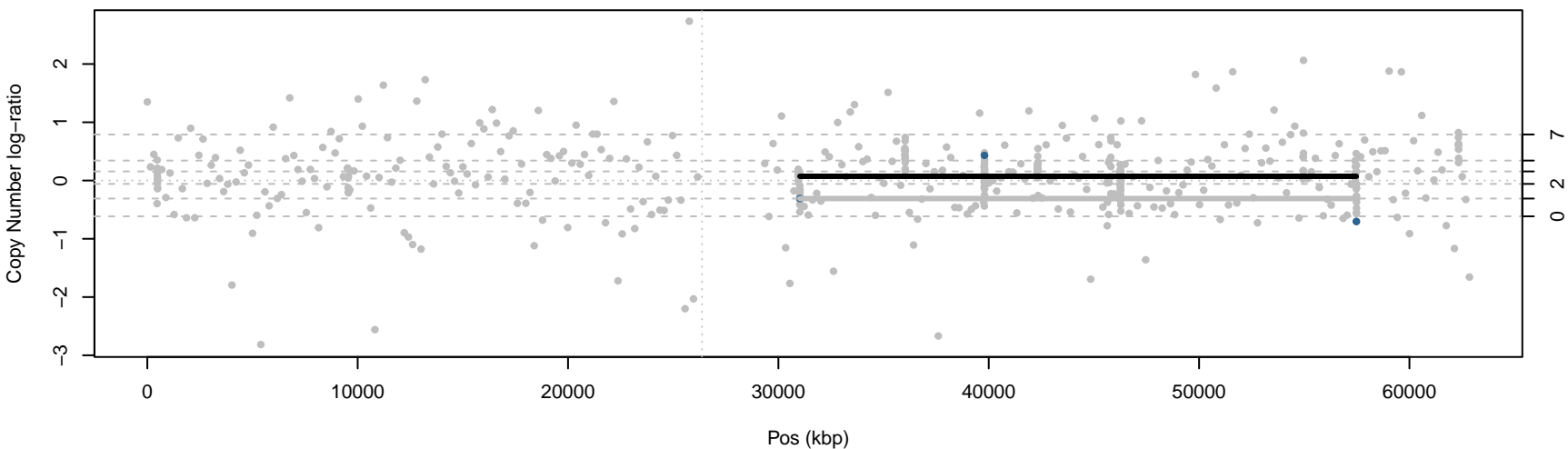
Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 21

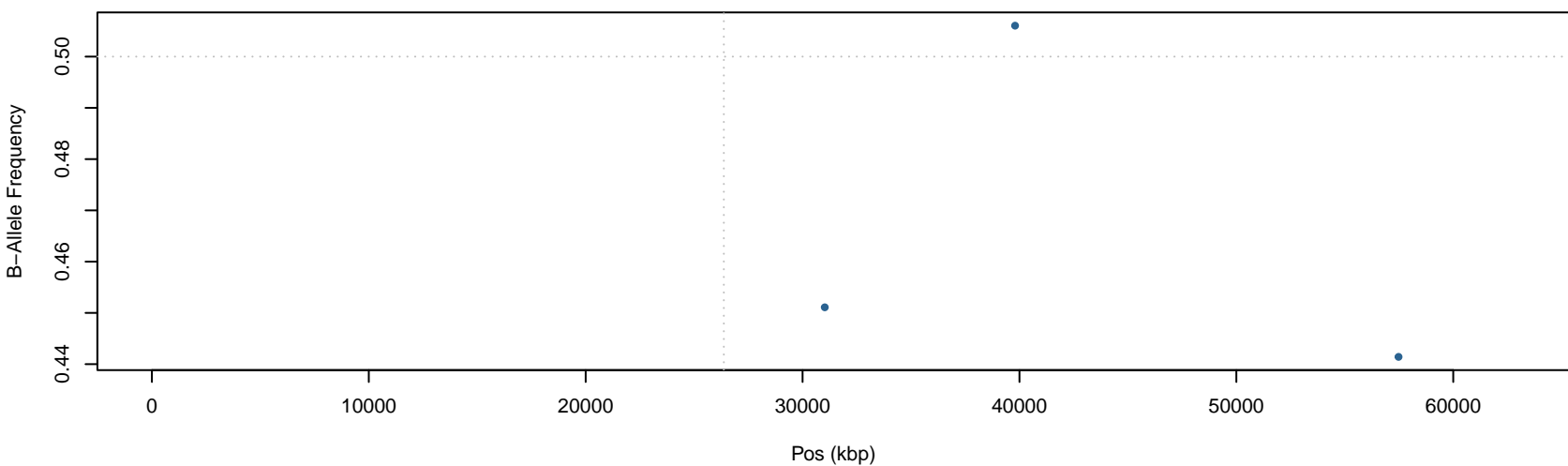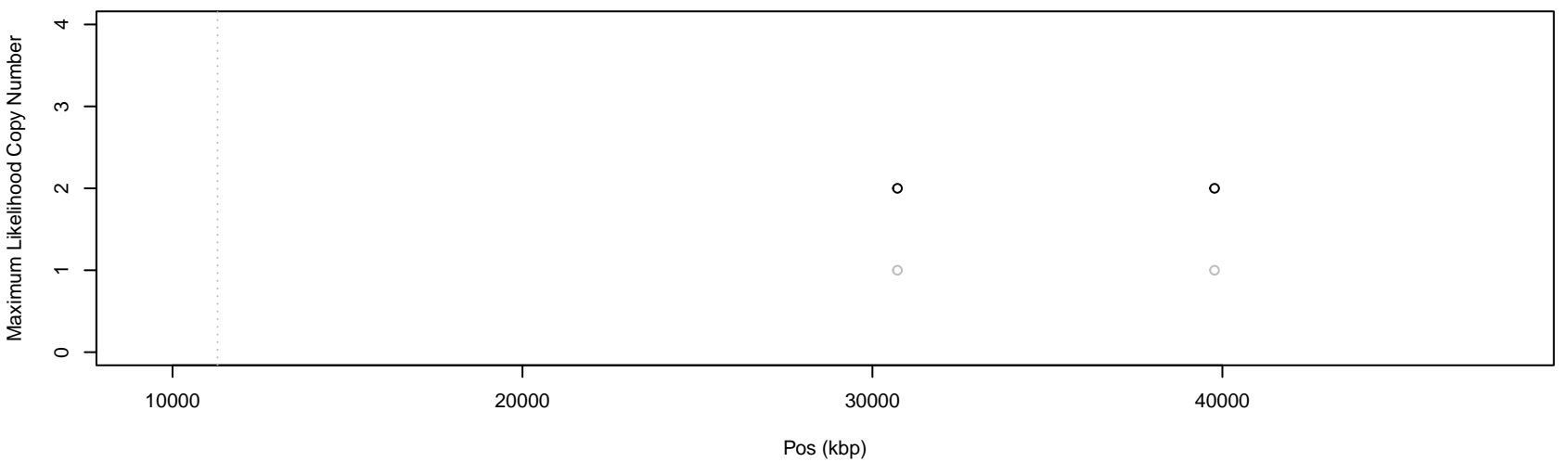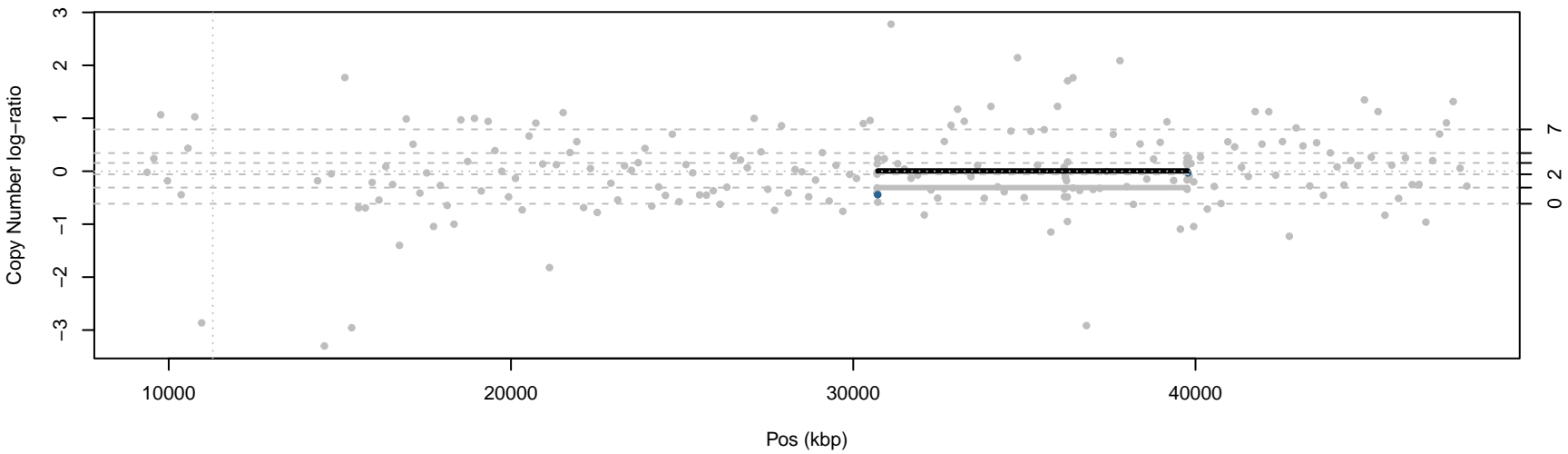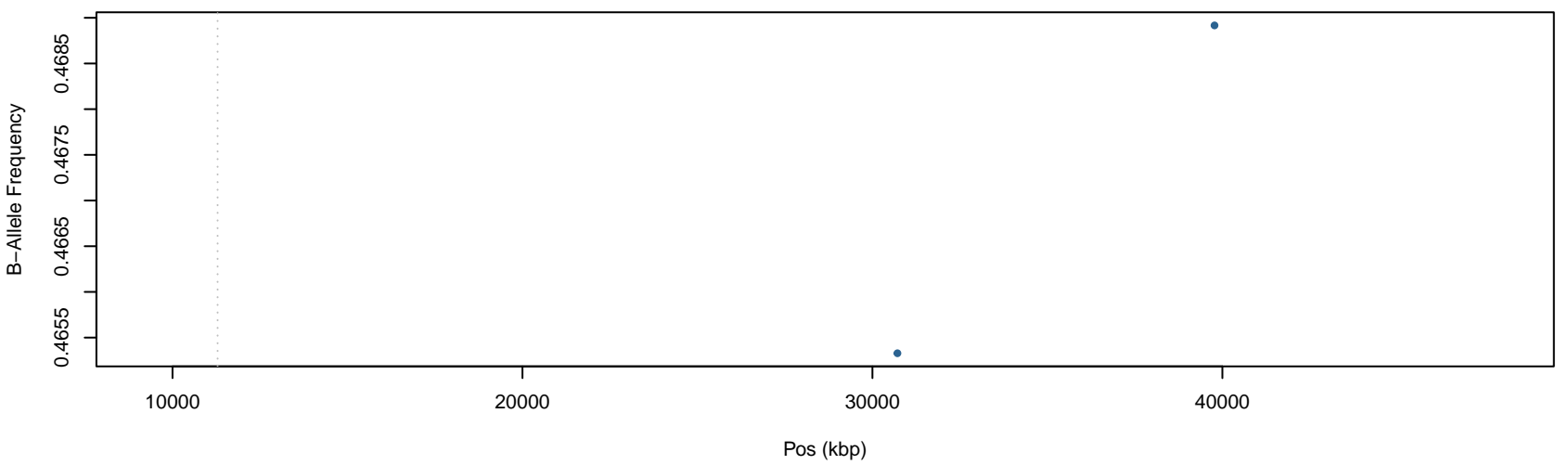**Purity: 0.32 Tumor ploidy: 2.256 SNV log−likelihood: 30.2 GoF: 86% Mean coverage: 575;453 Chromosome: 22**