Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 1

Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 2
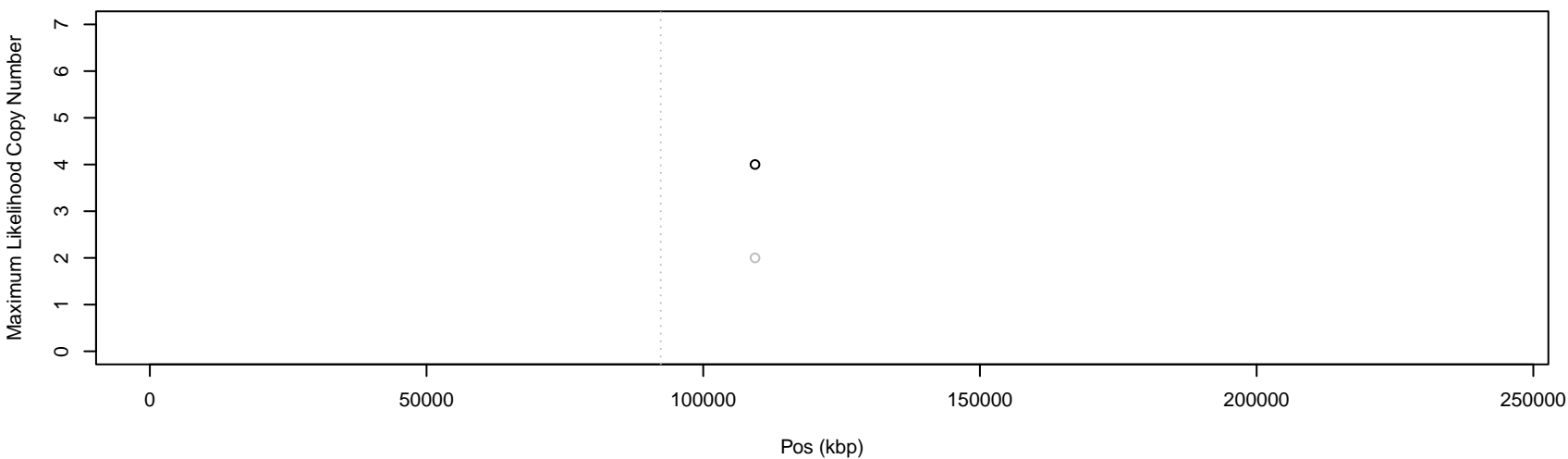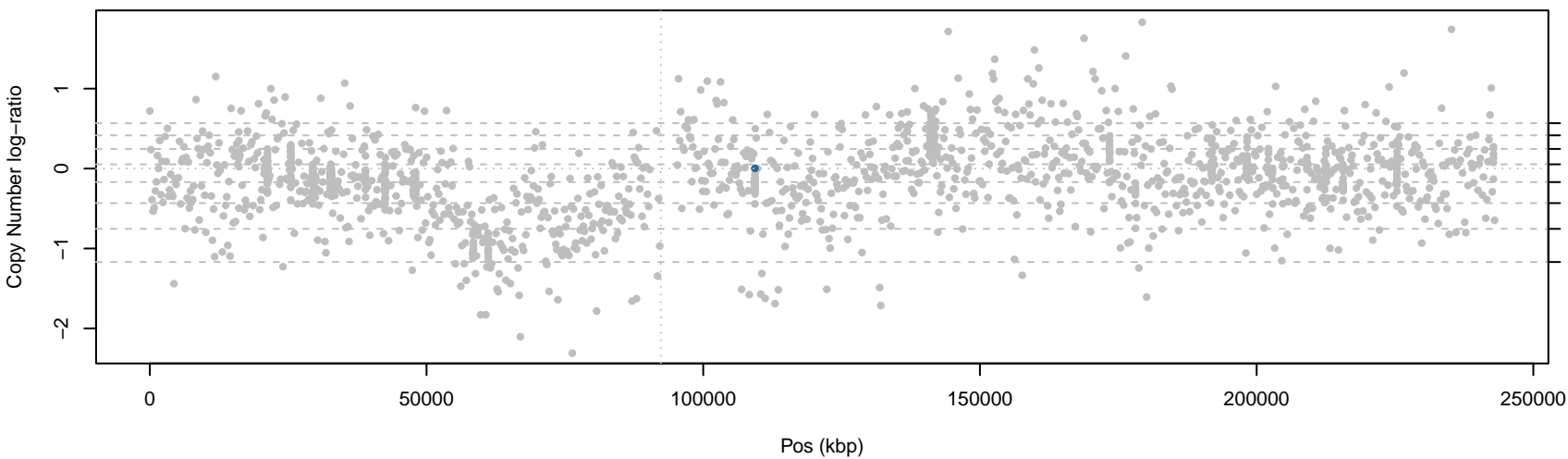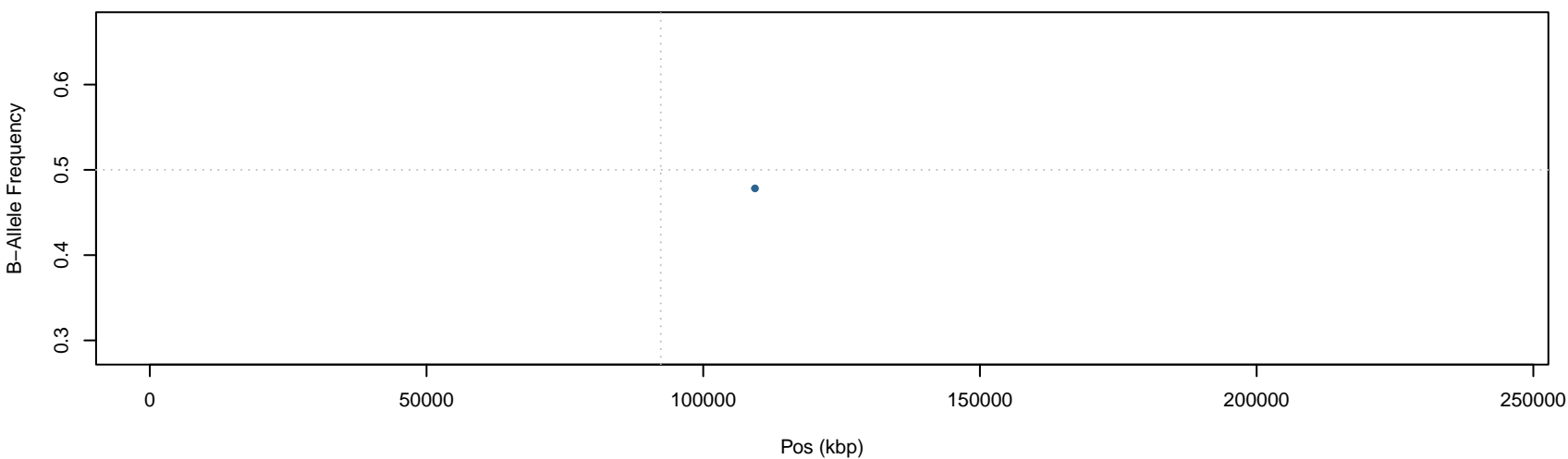
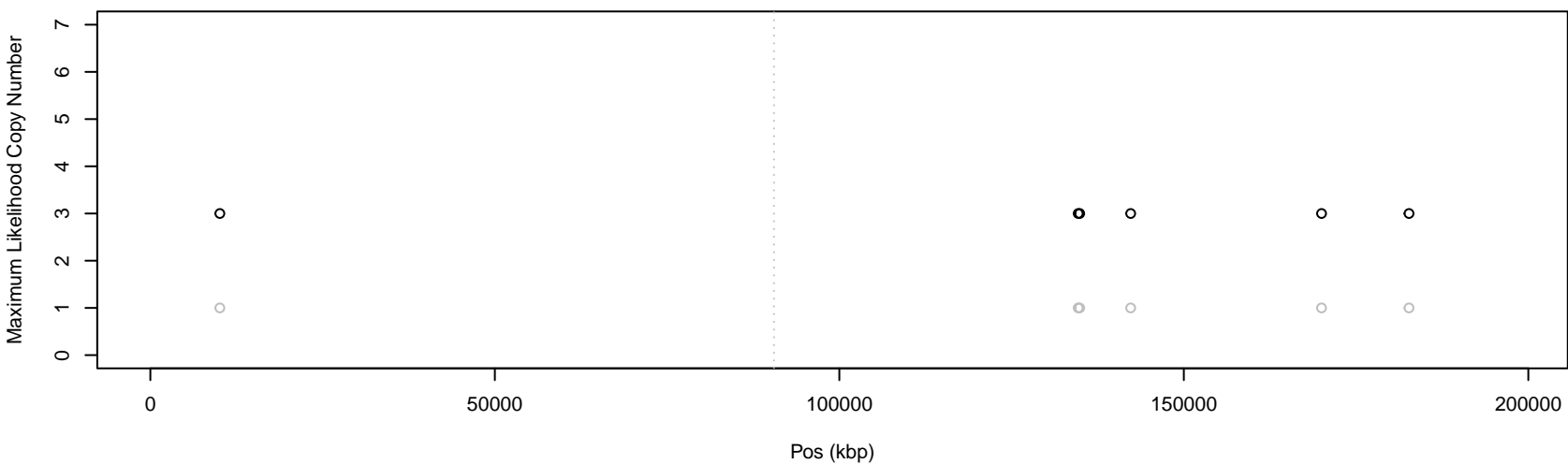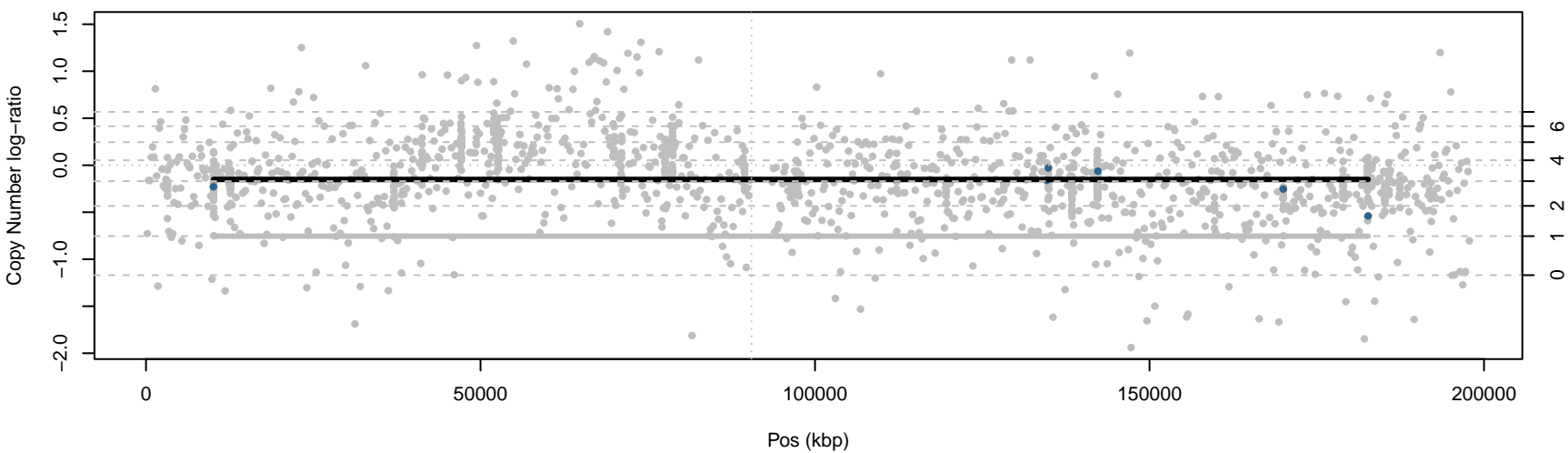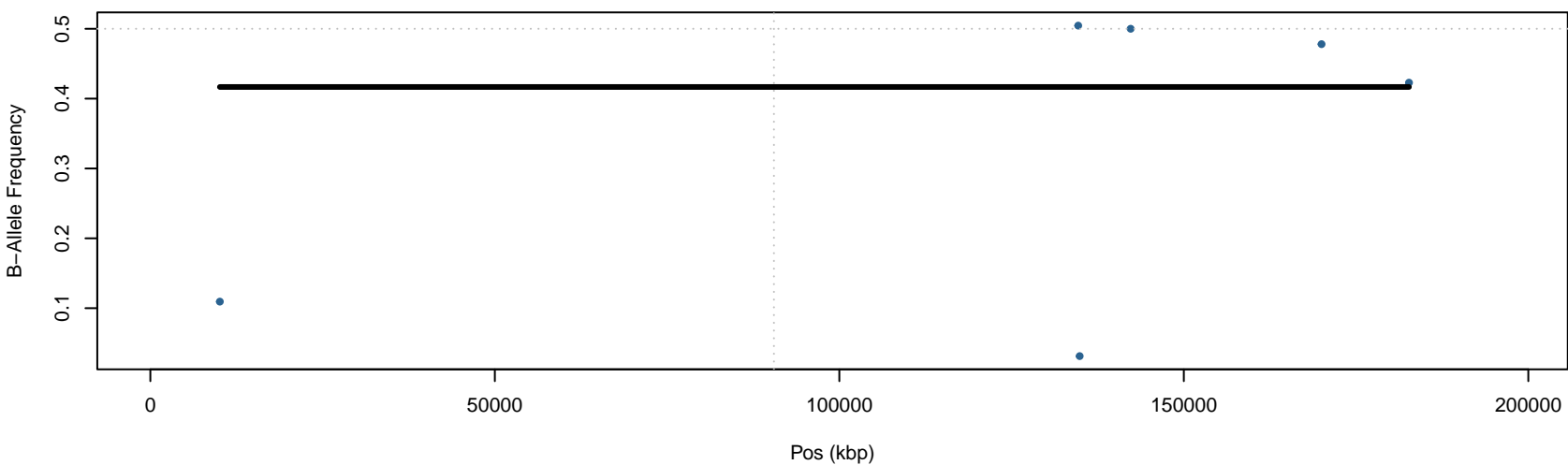**Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 3**
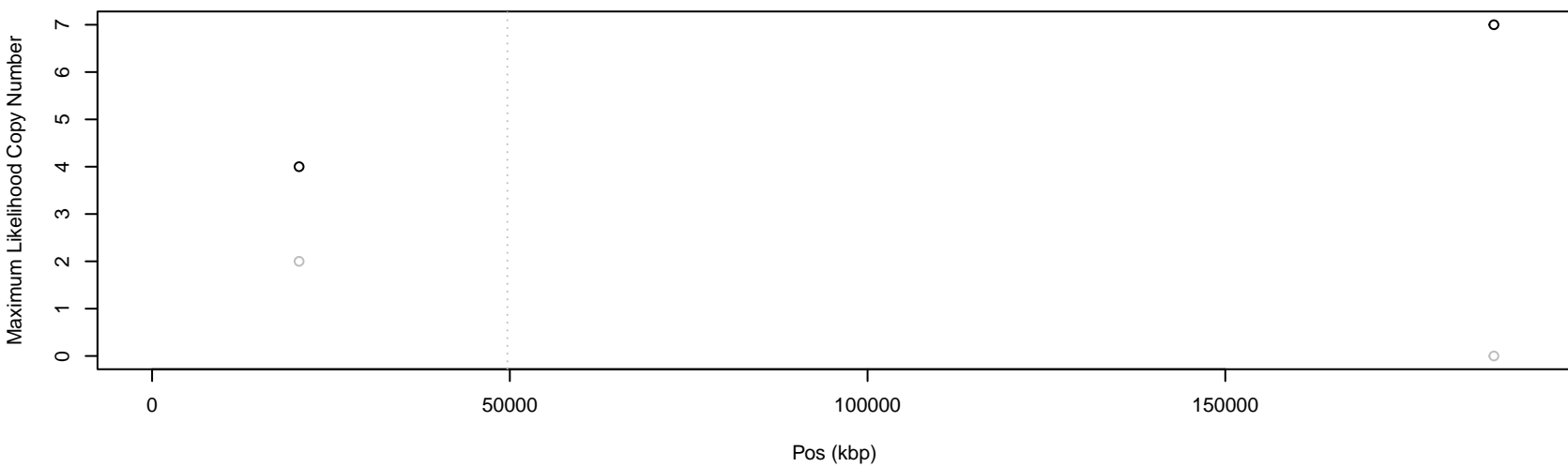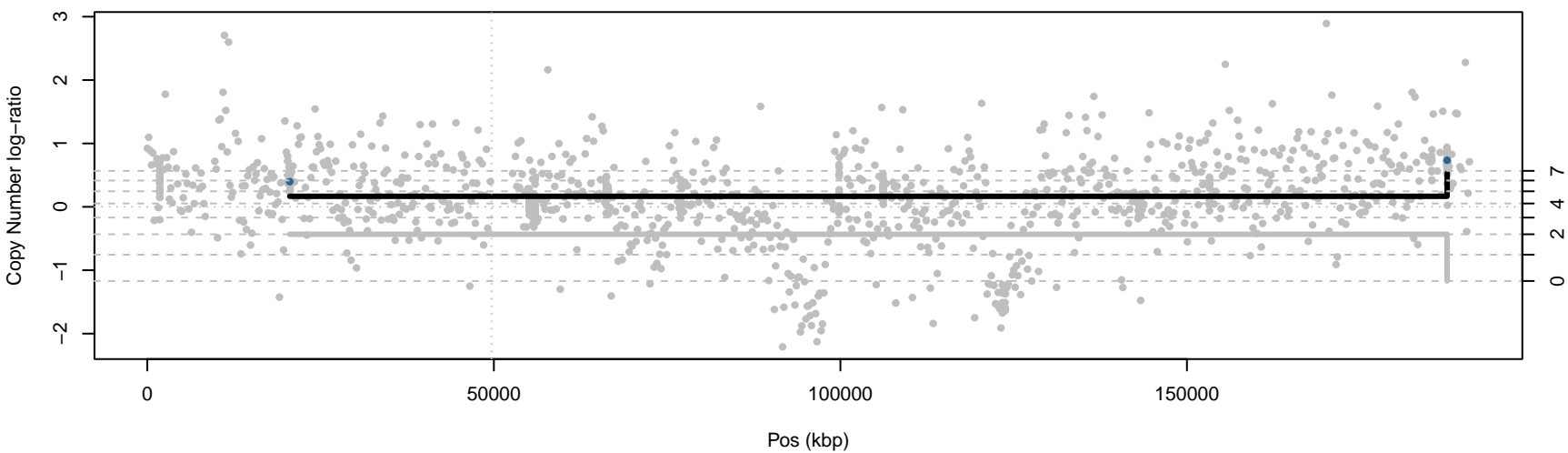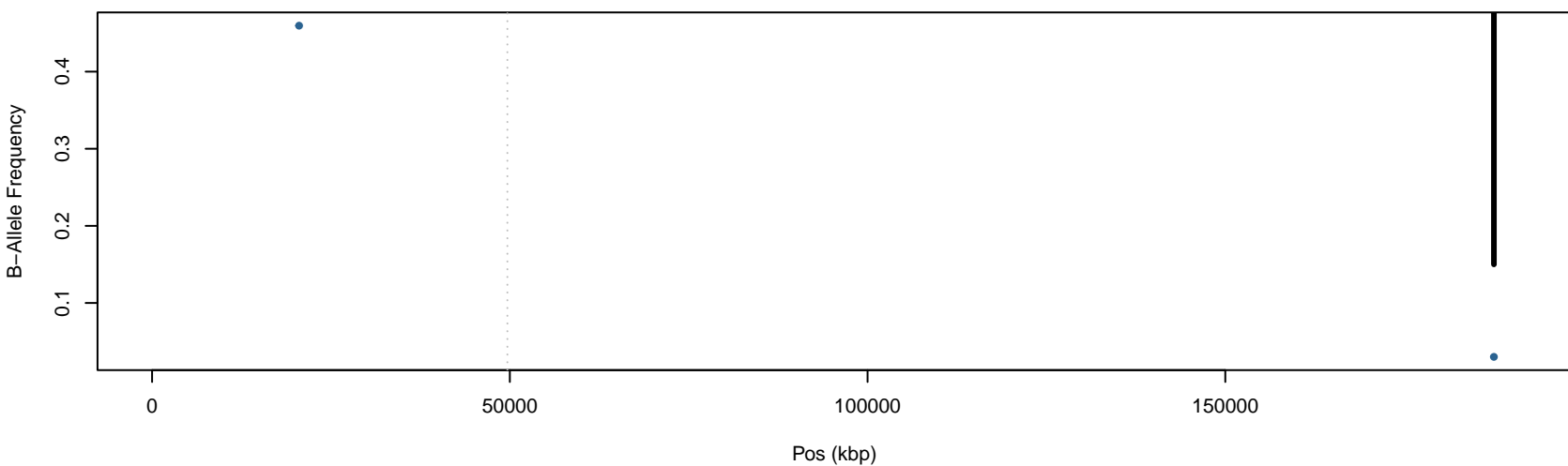
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 4
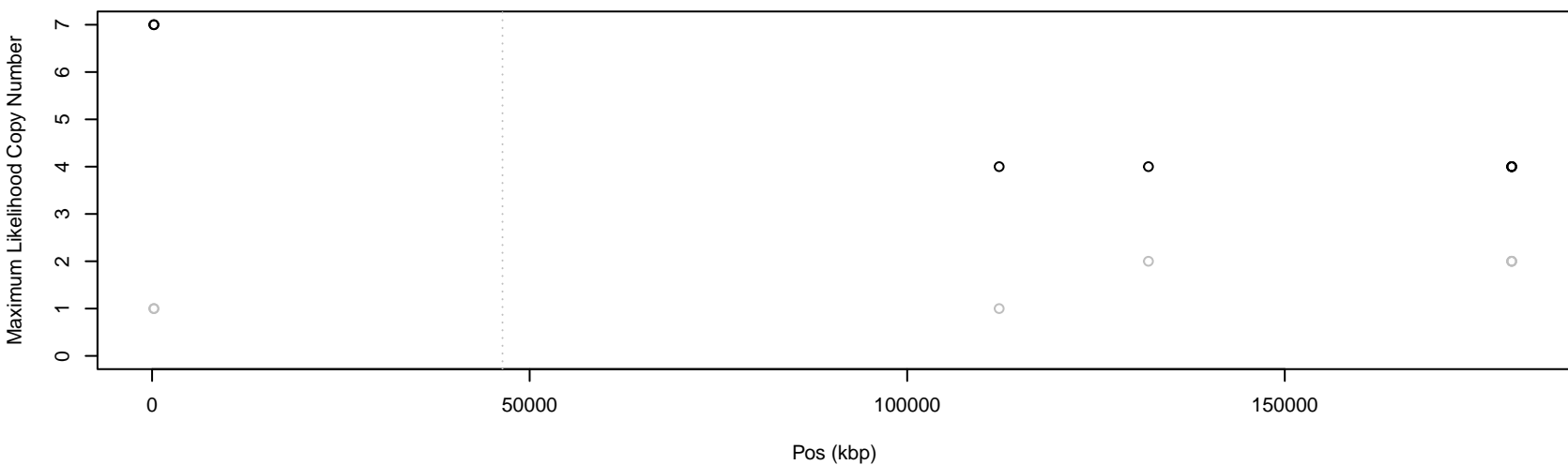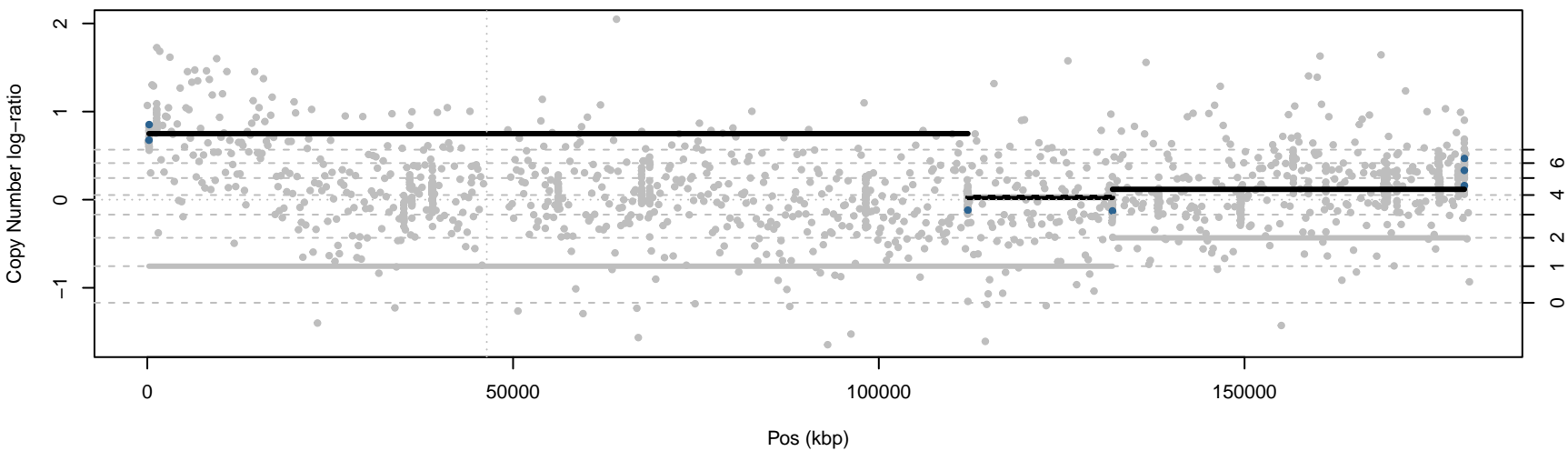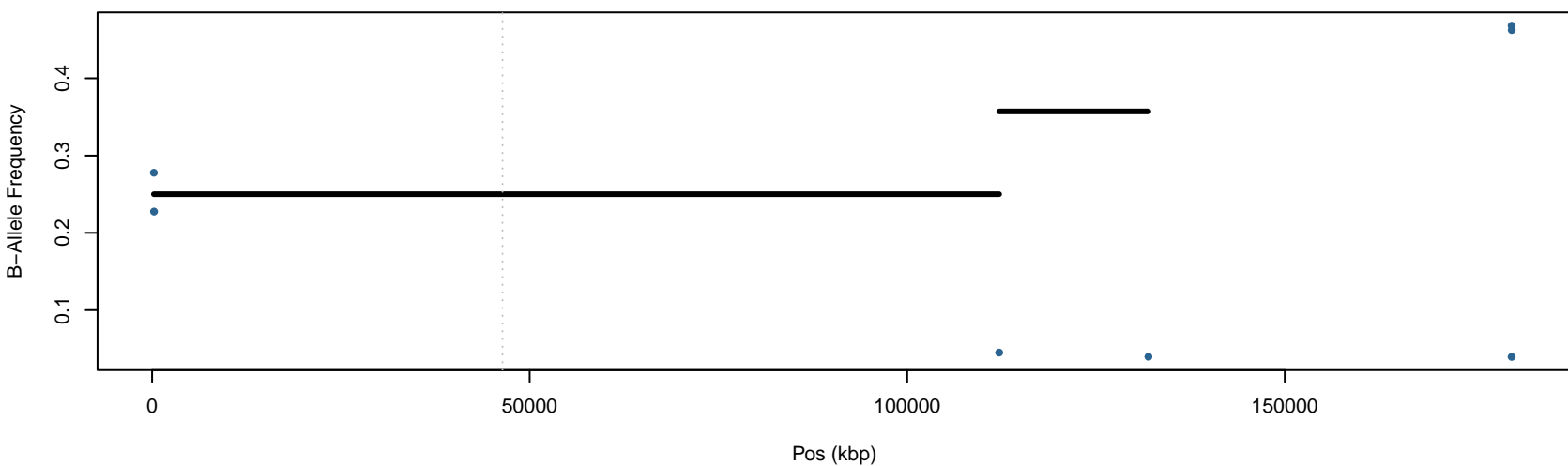
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 5
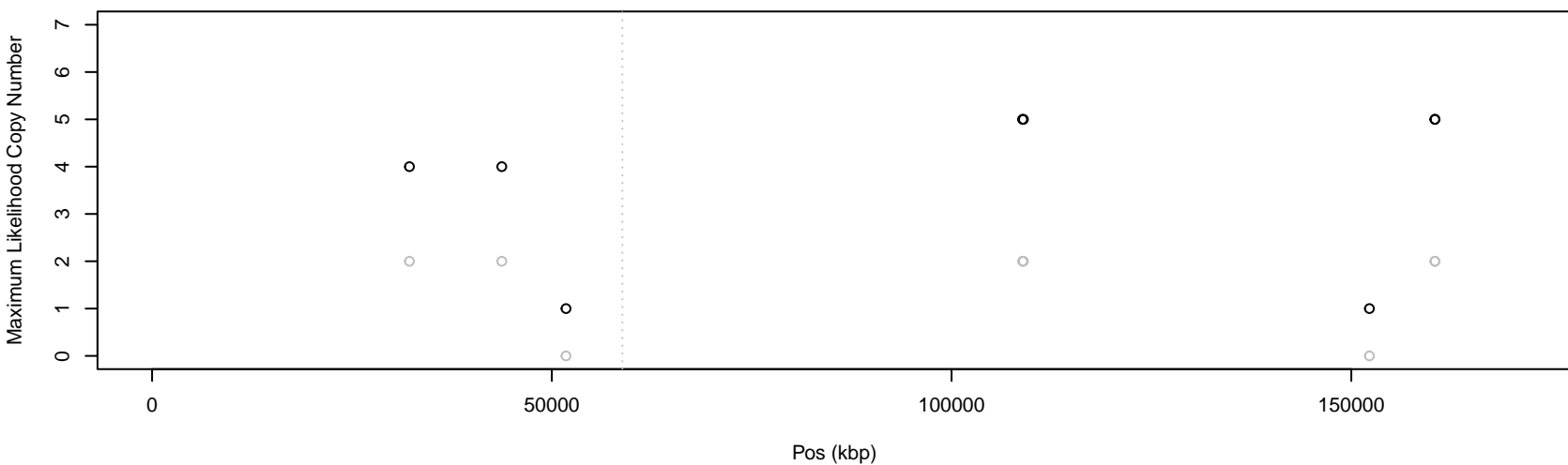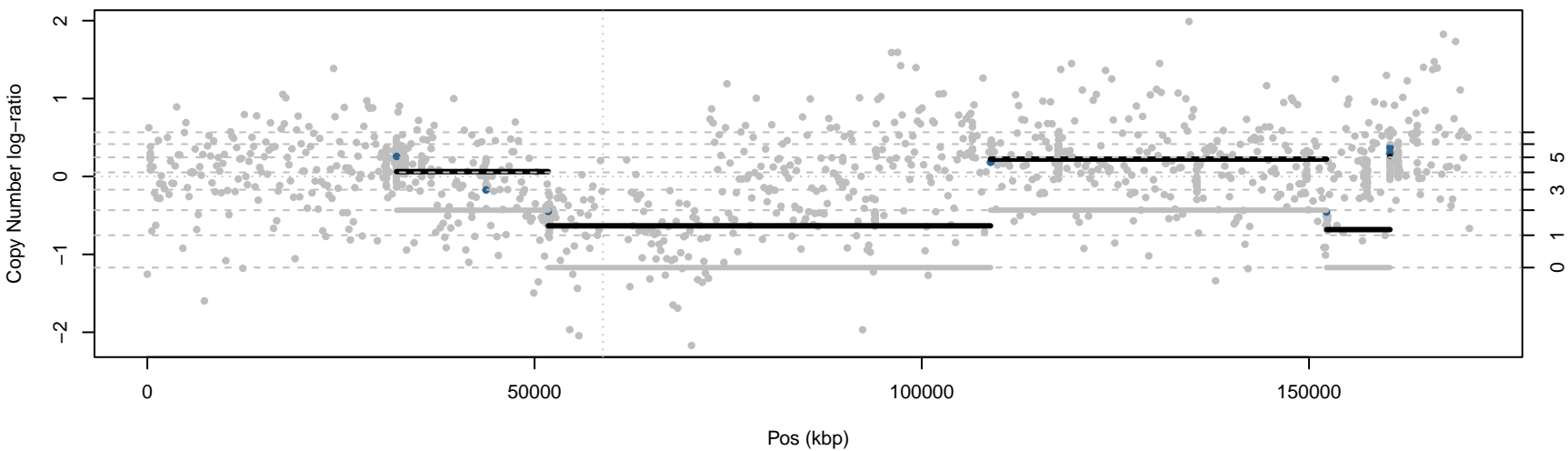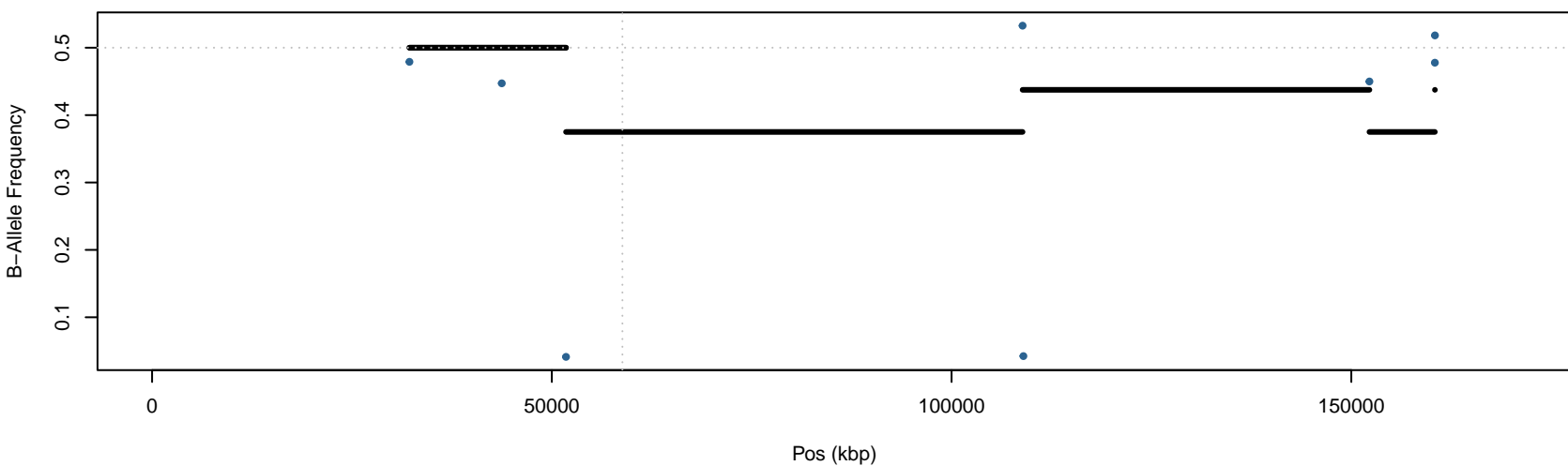
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 6
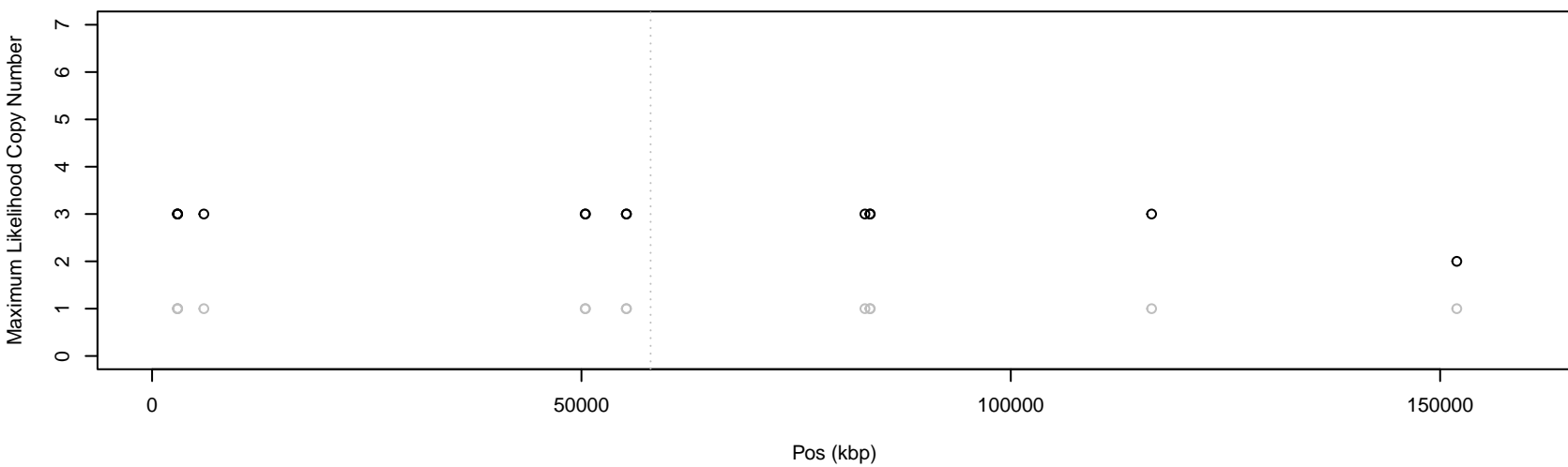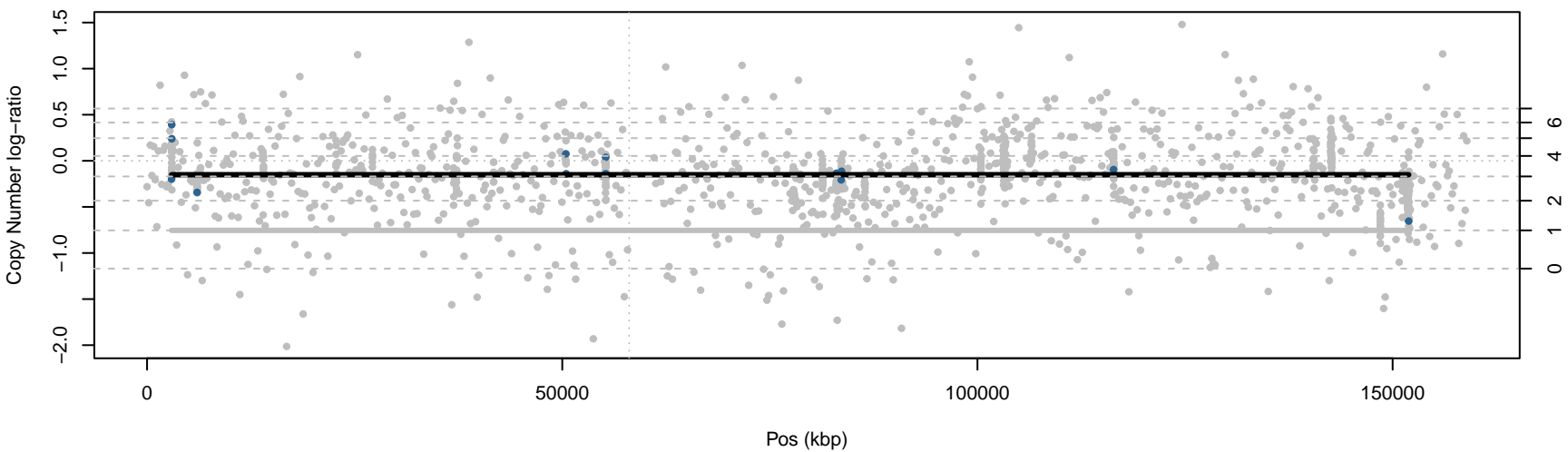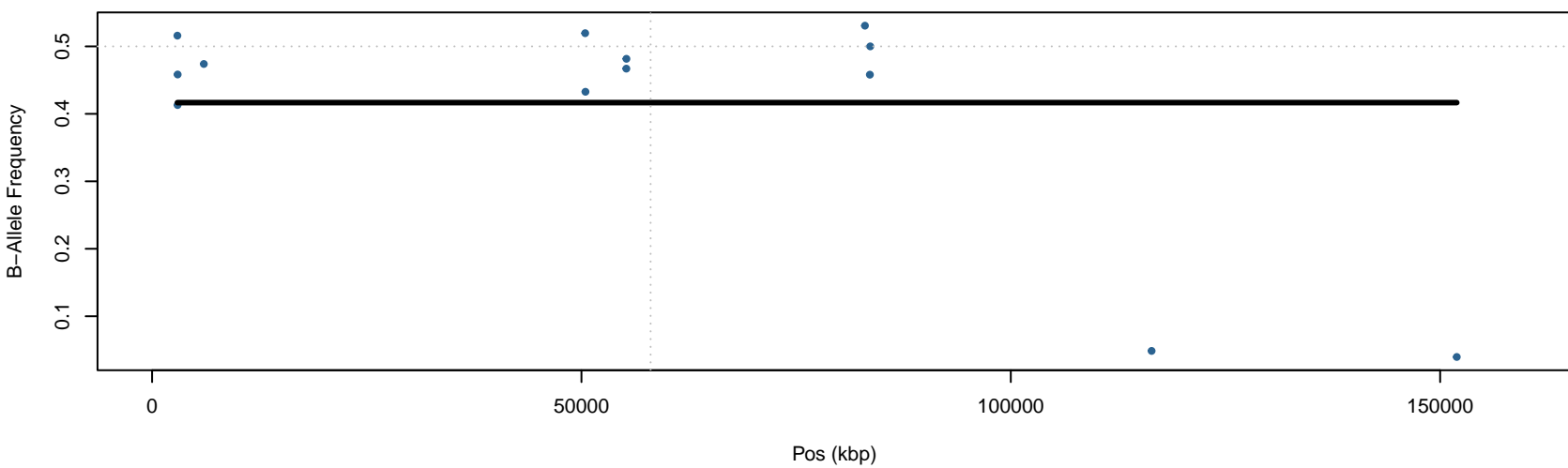
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 7
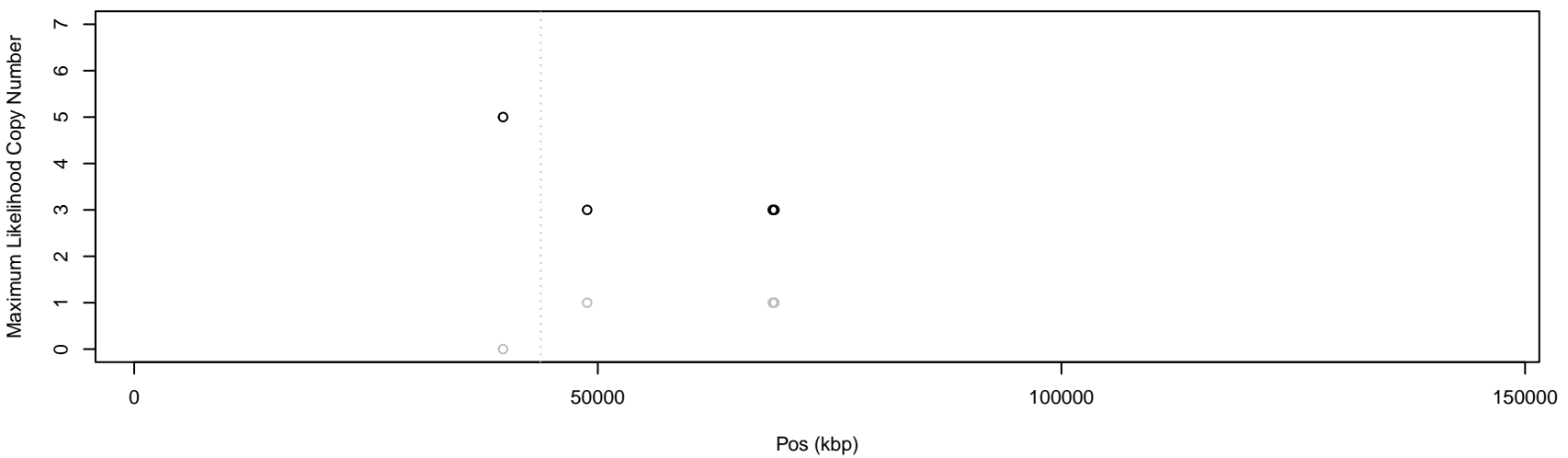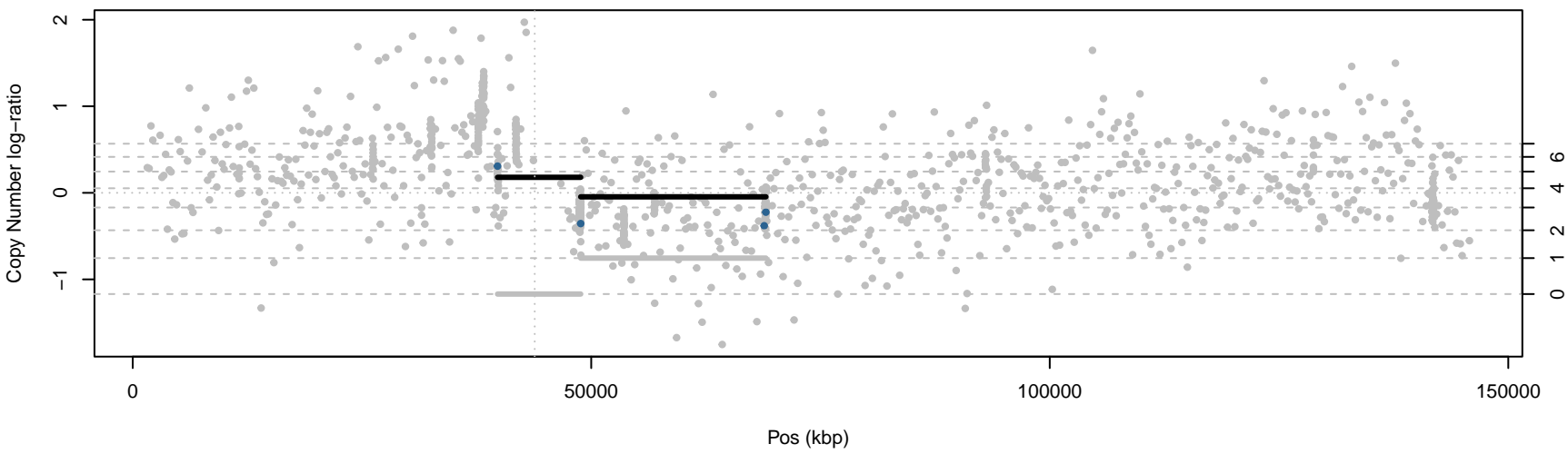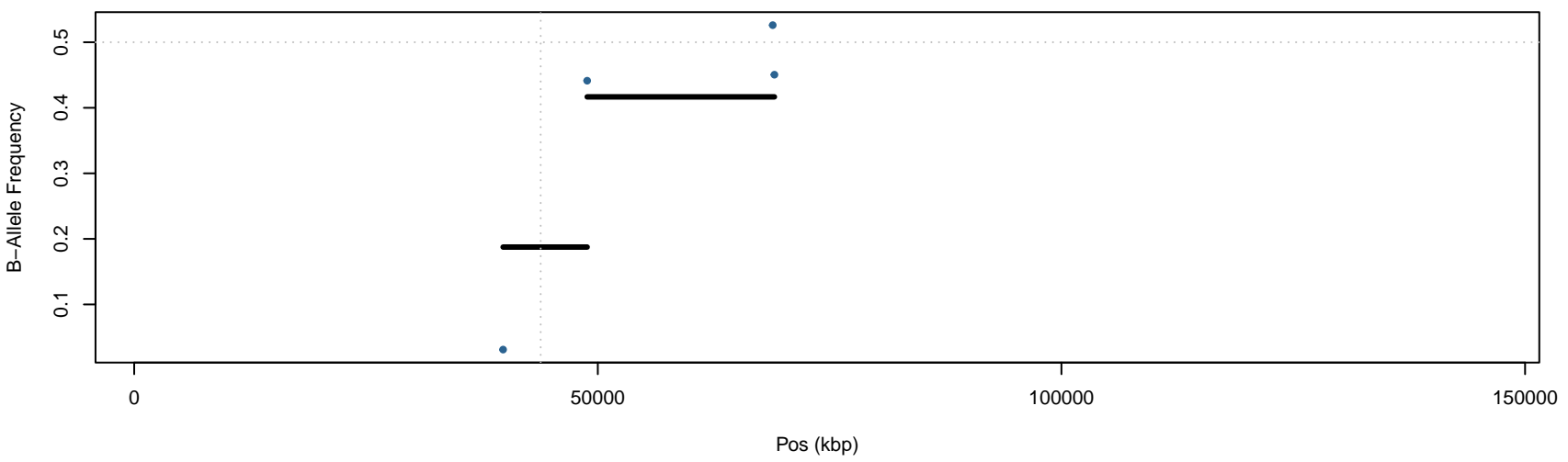
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 8

Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 9
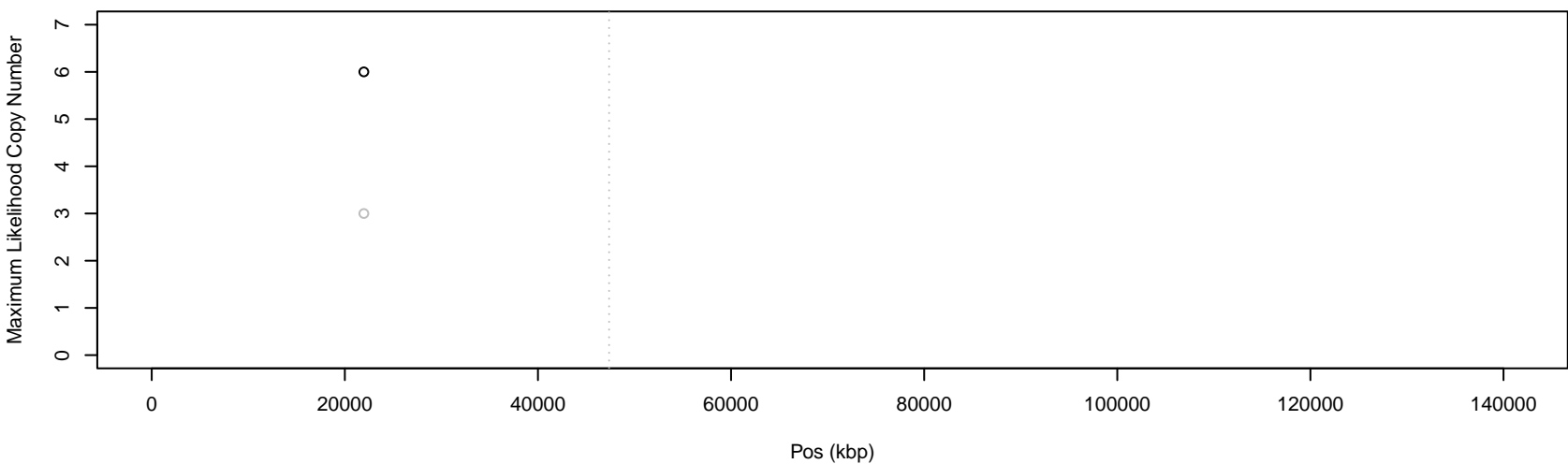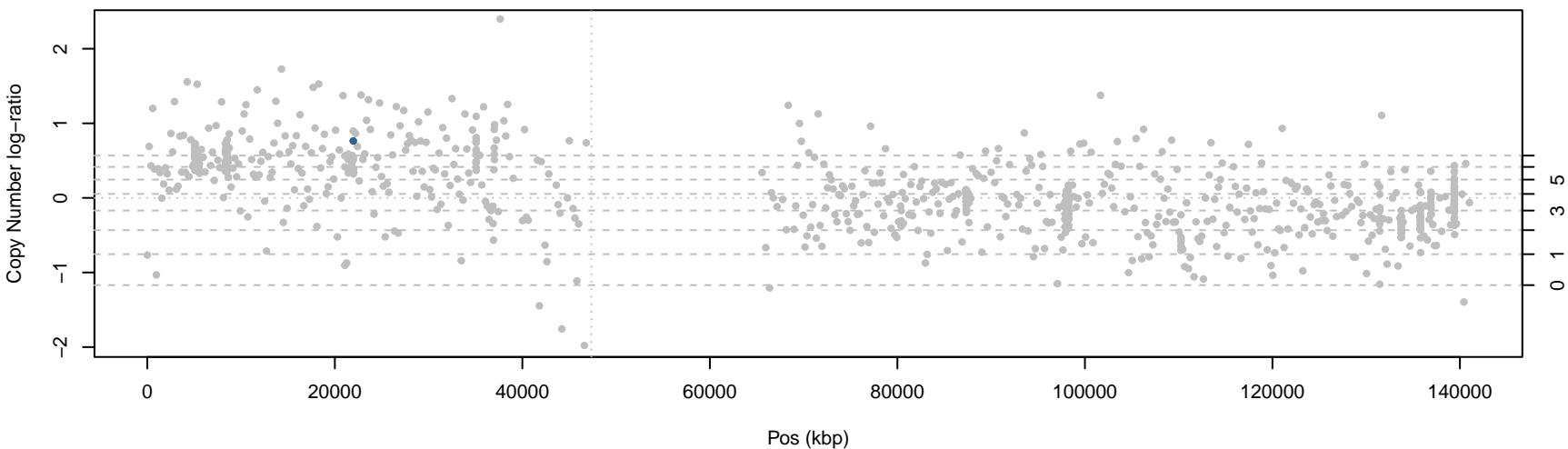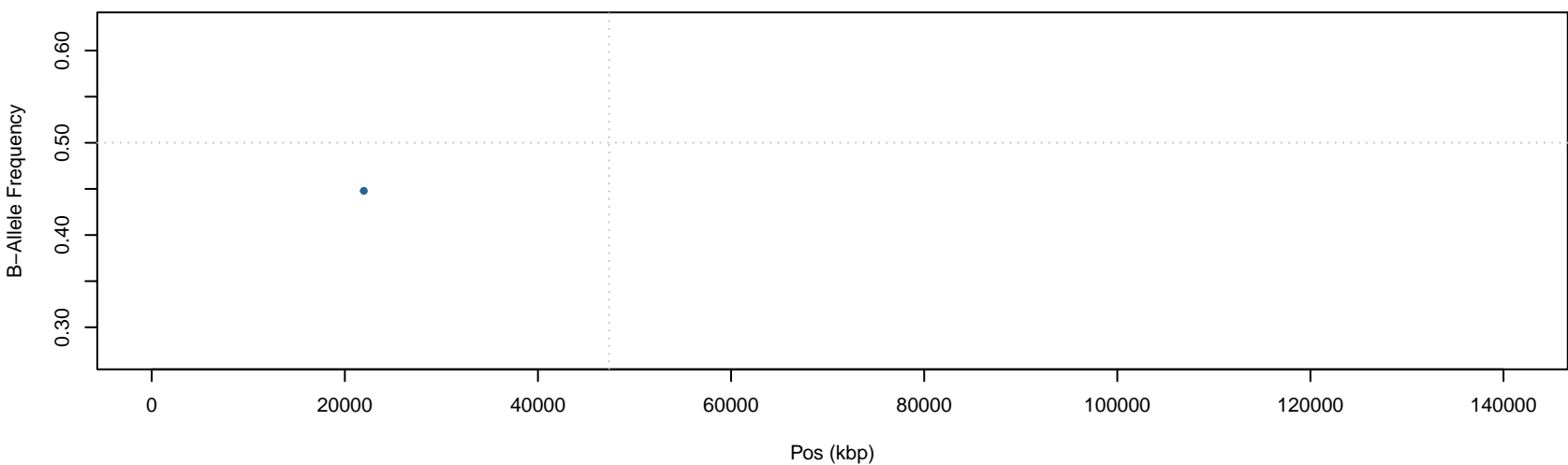
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 10
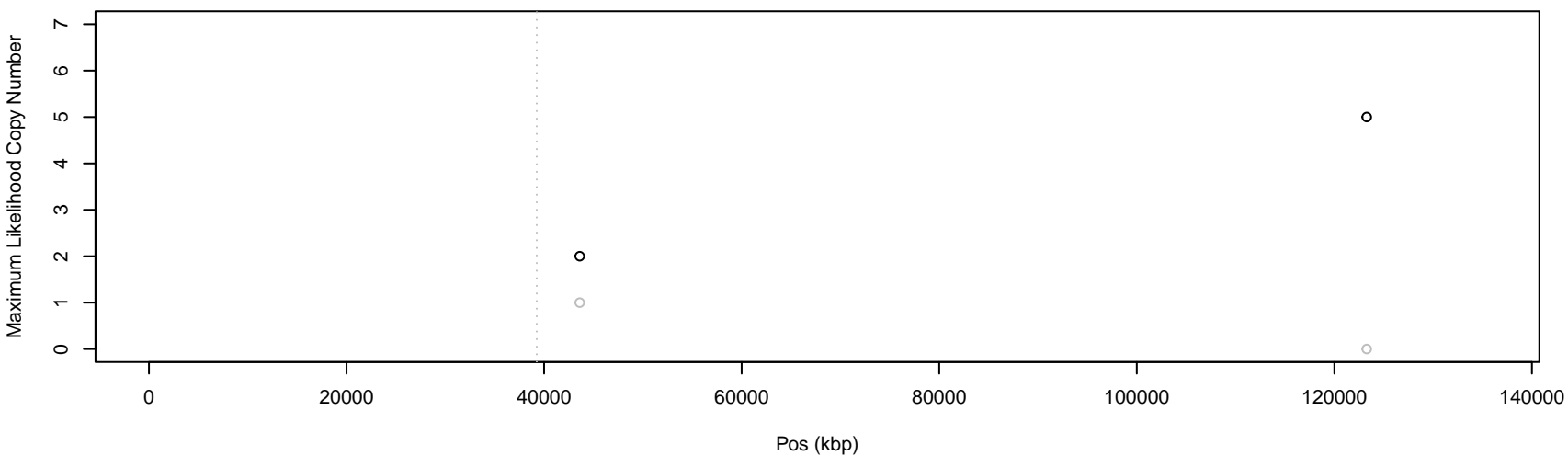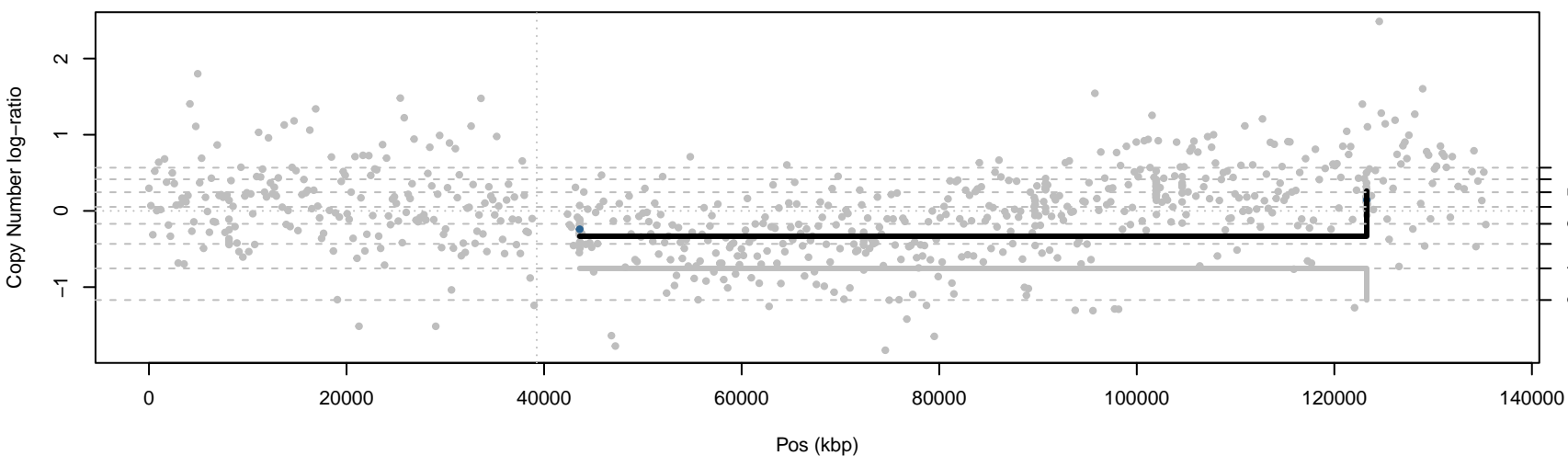
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 11

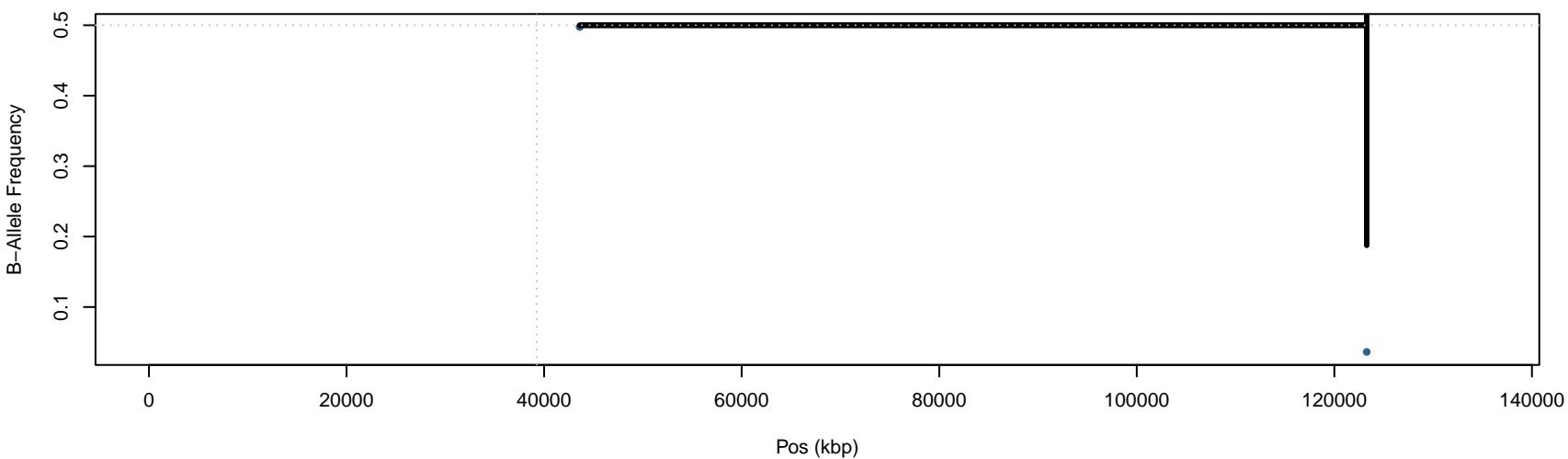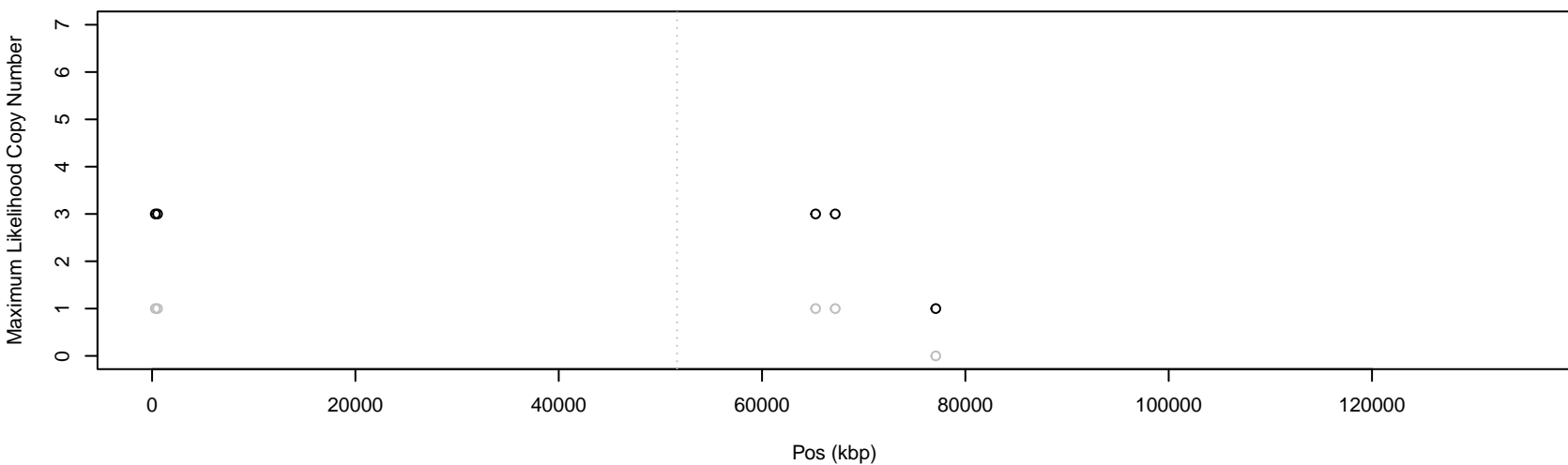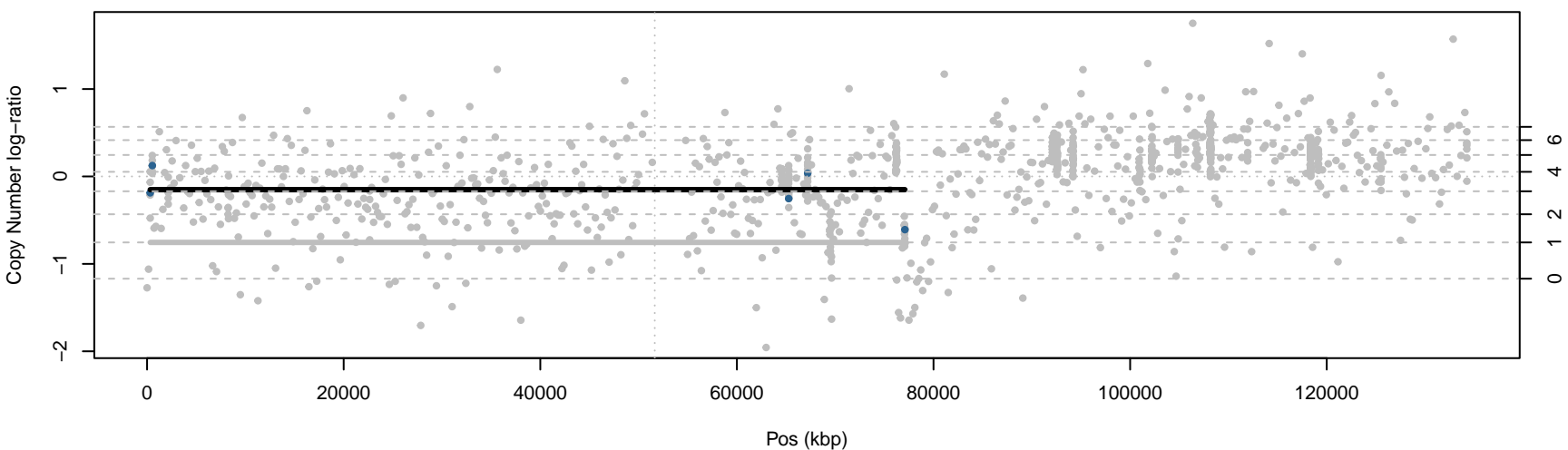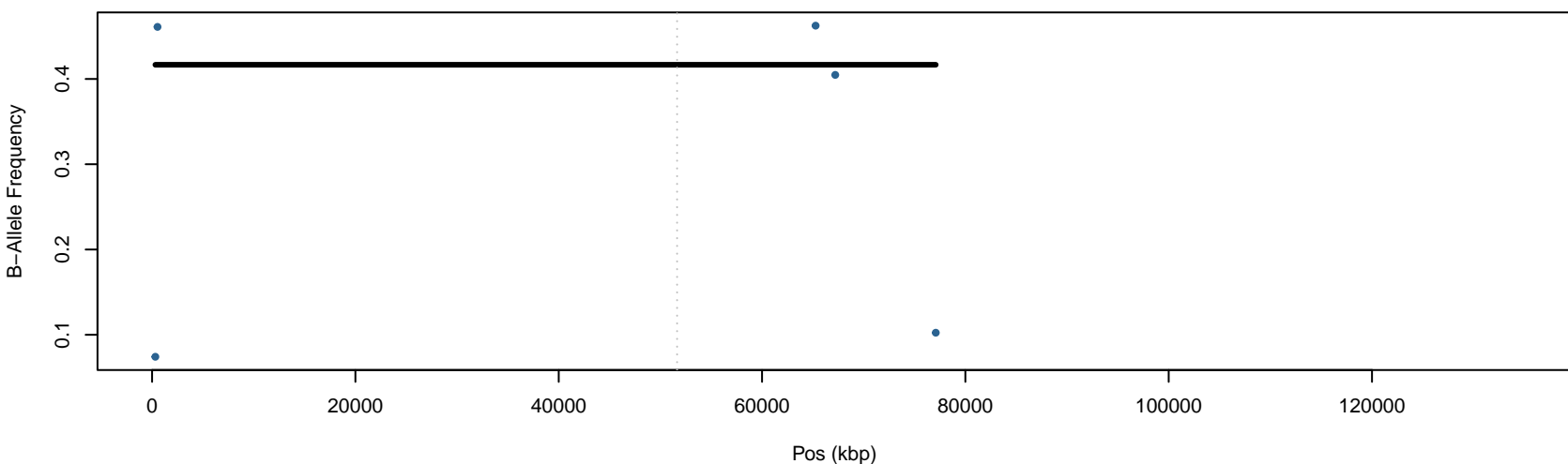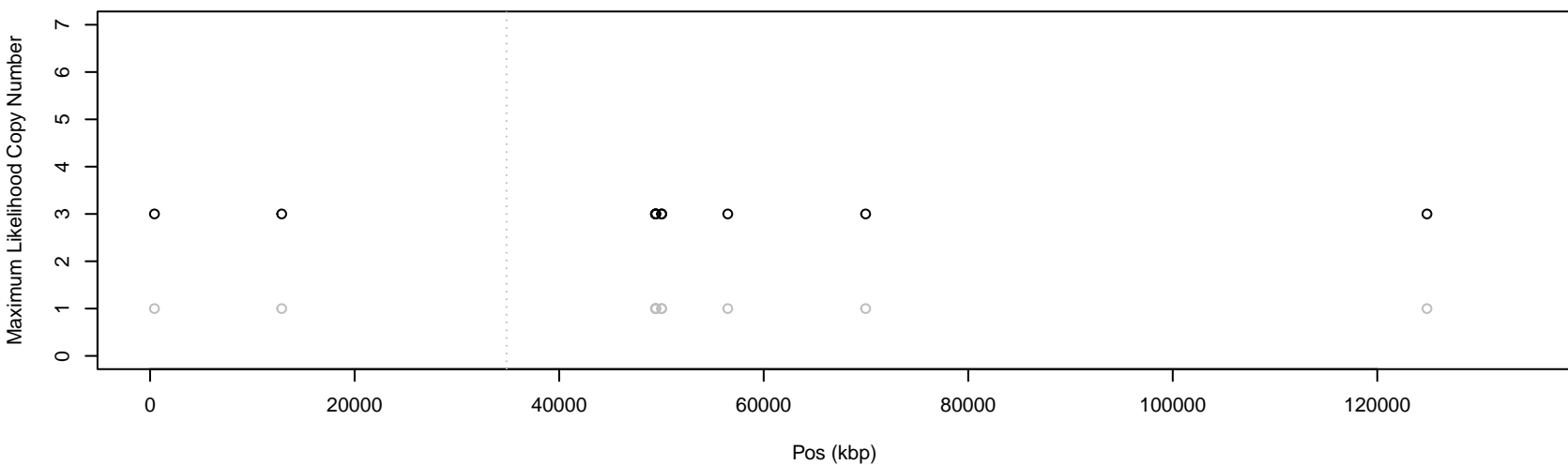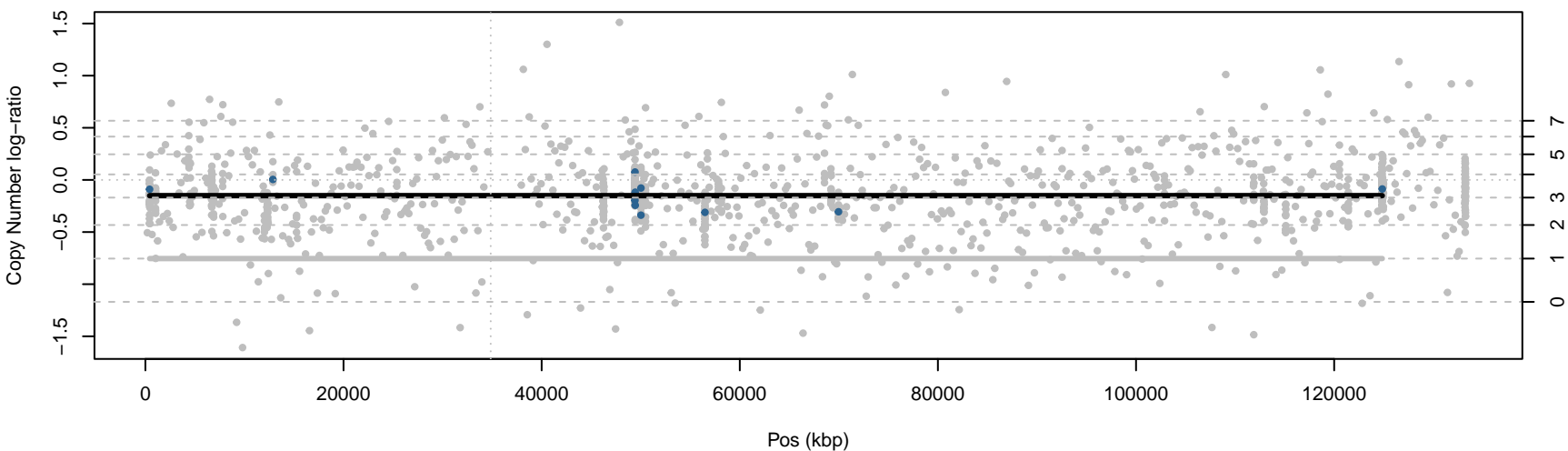**Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 12**

Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 15
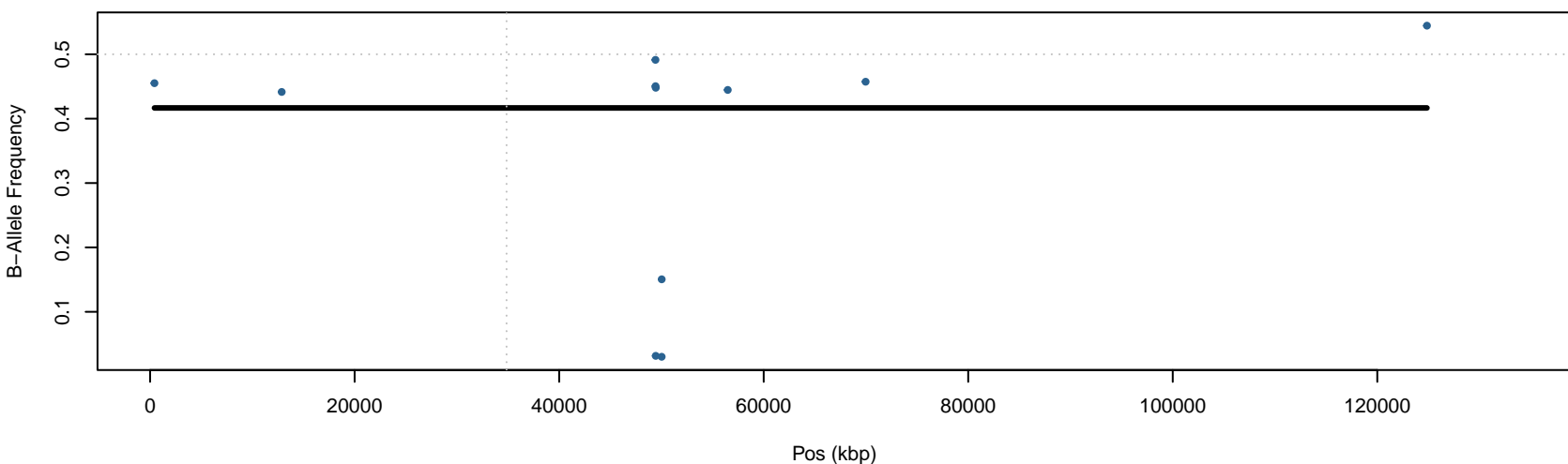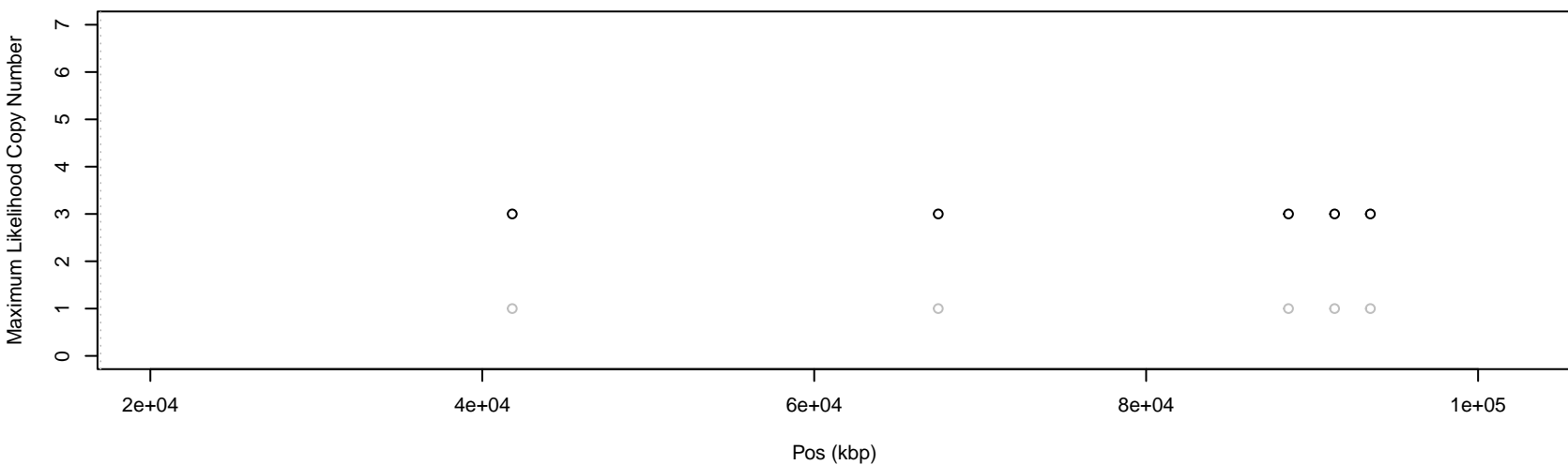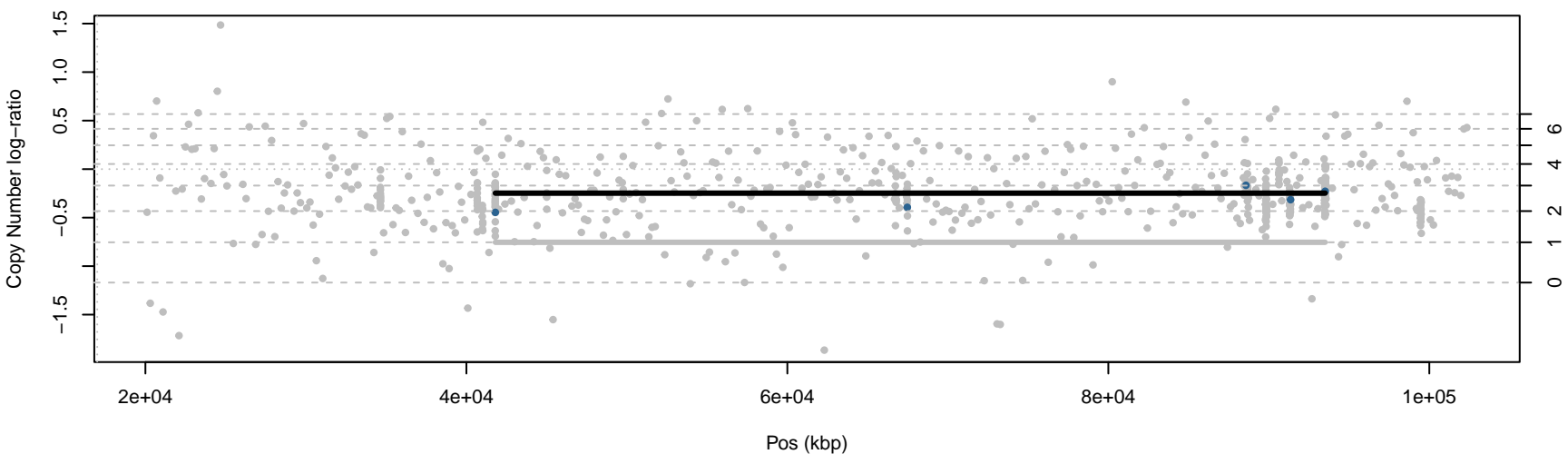
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 16
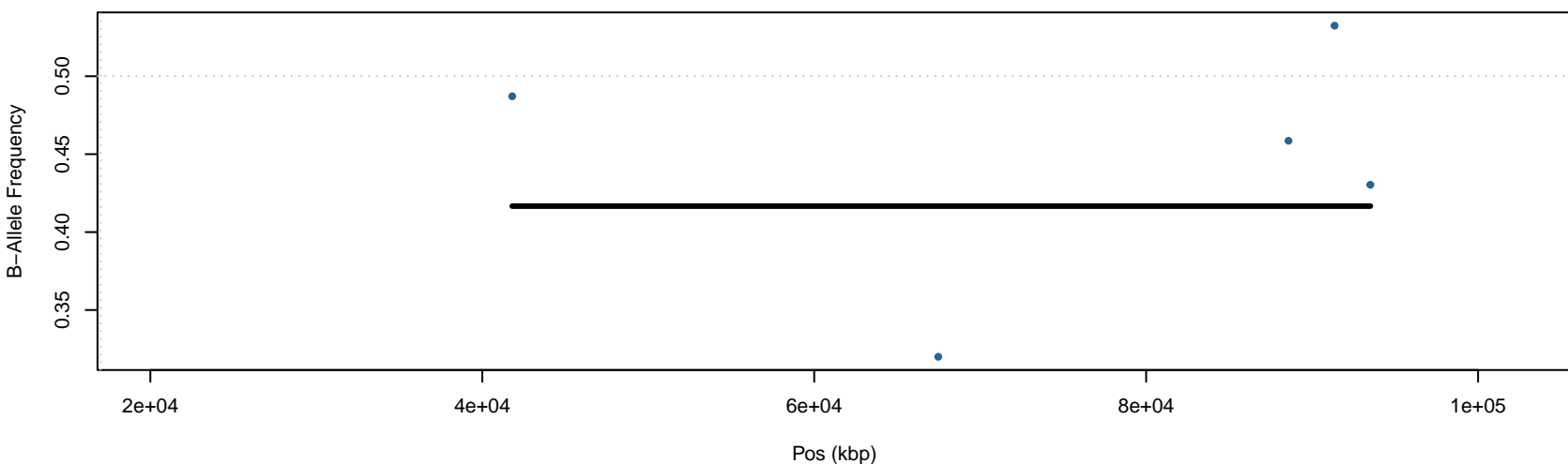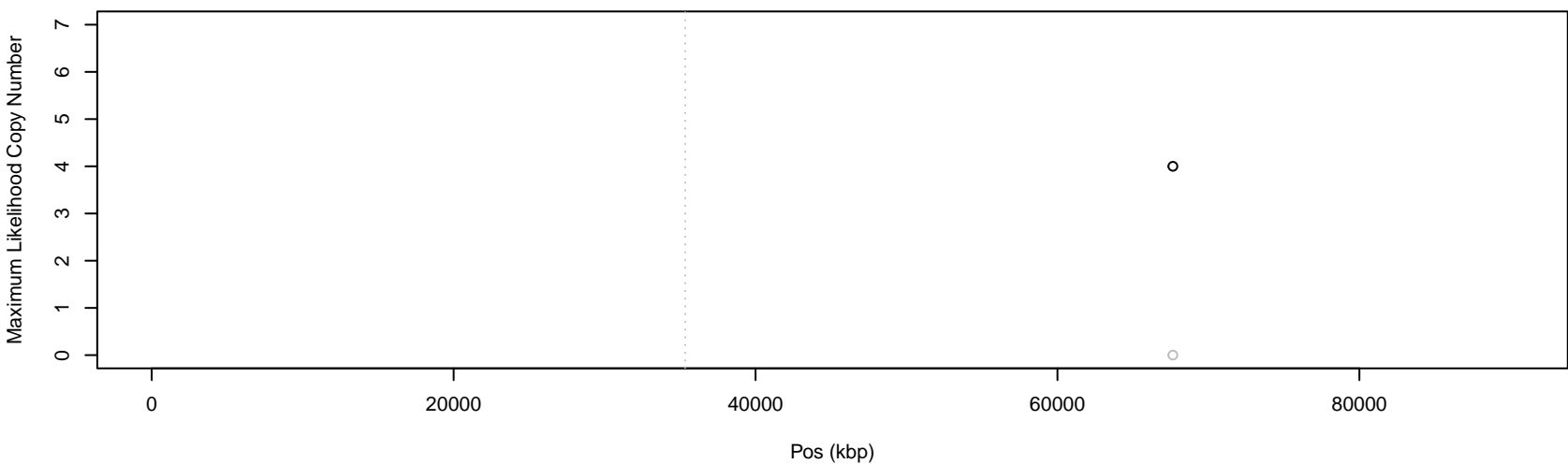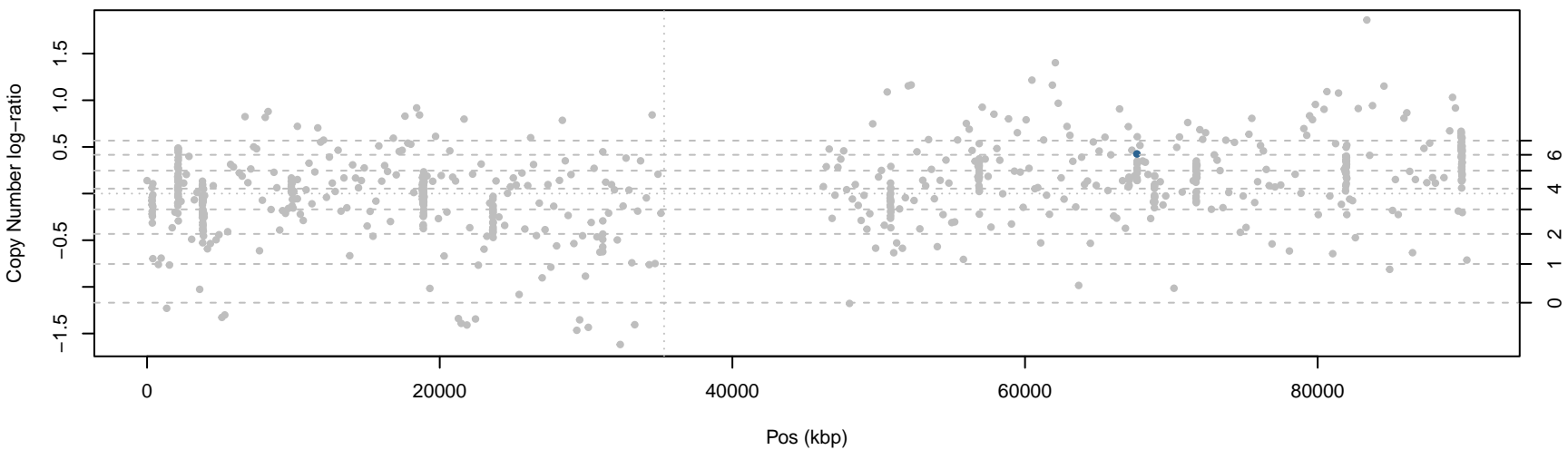
Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 19

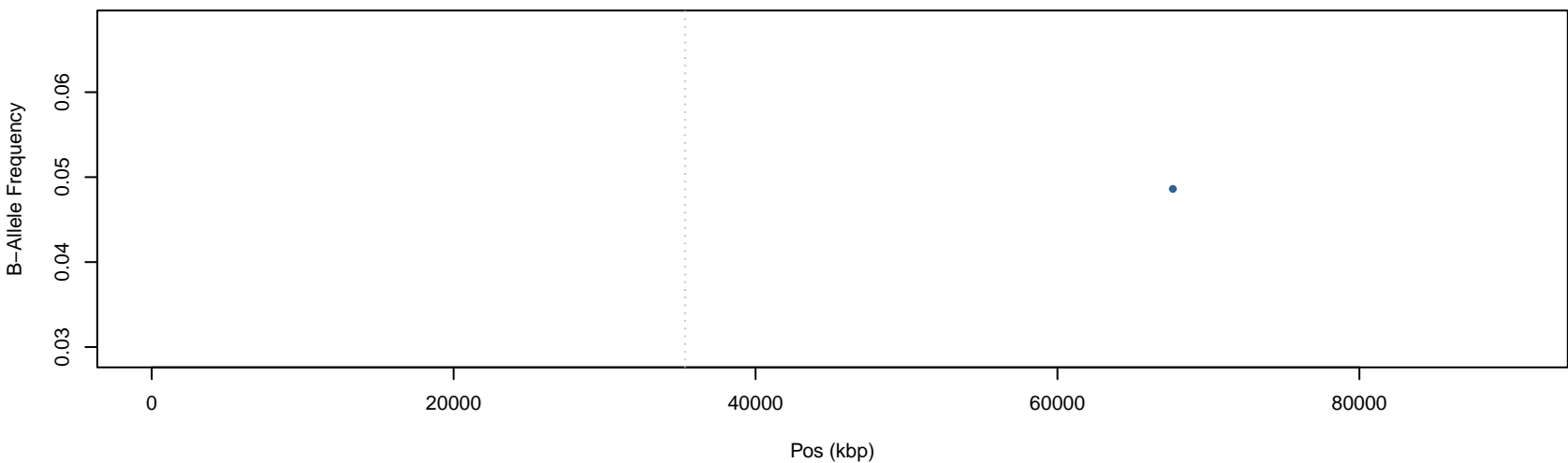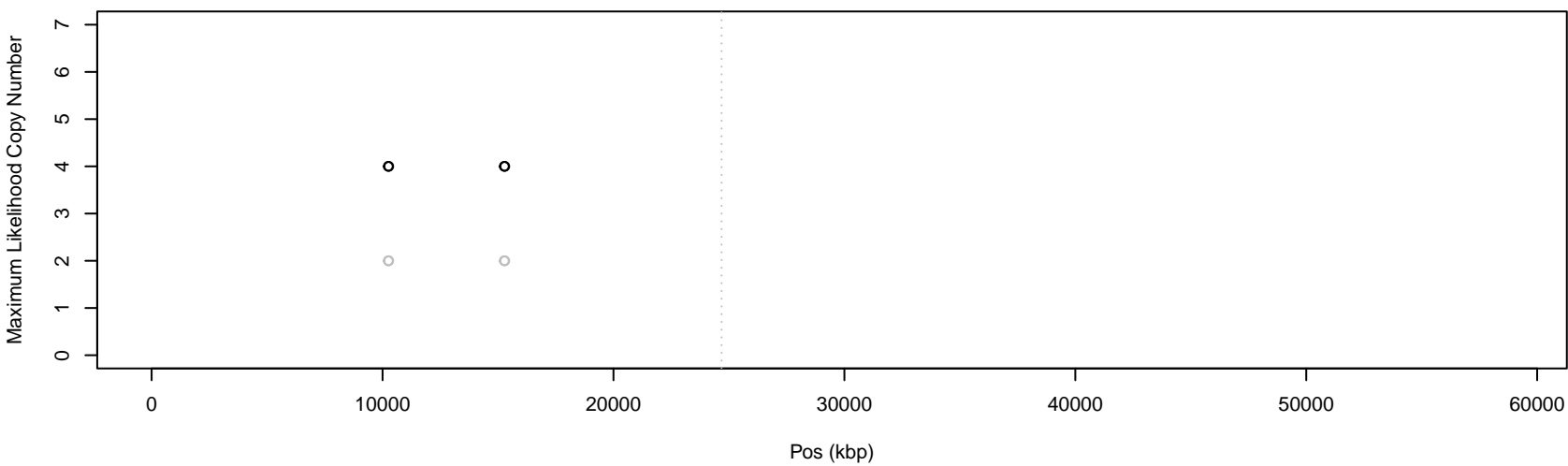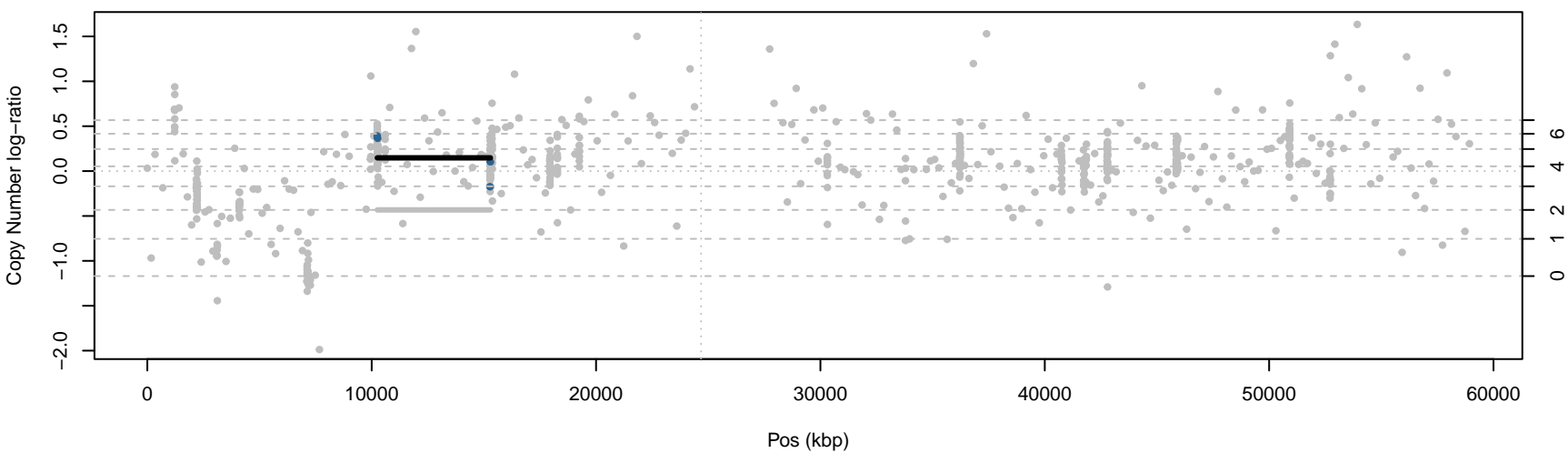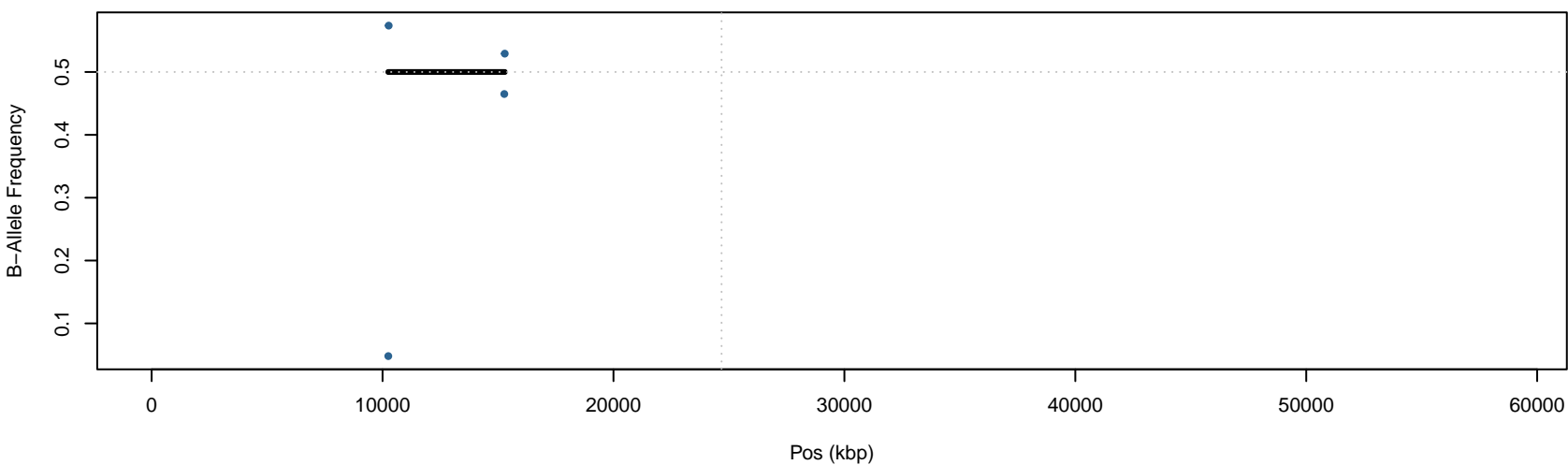**Purity: 0.4 Tumor ploidy: 3.748 SNV log−likelihood: −57.53 GoF: 87% Mean coverage: 505;489 Chromosome: 21**