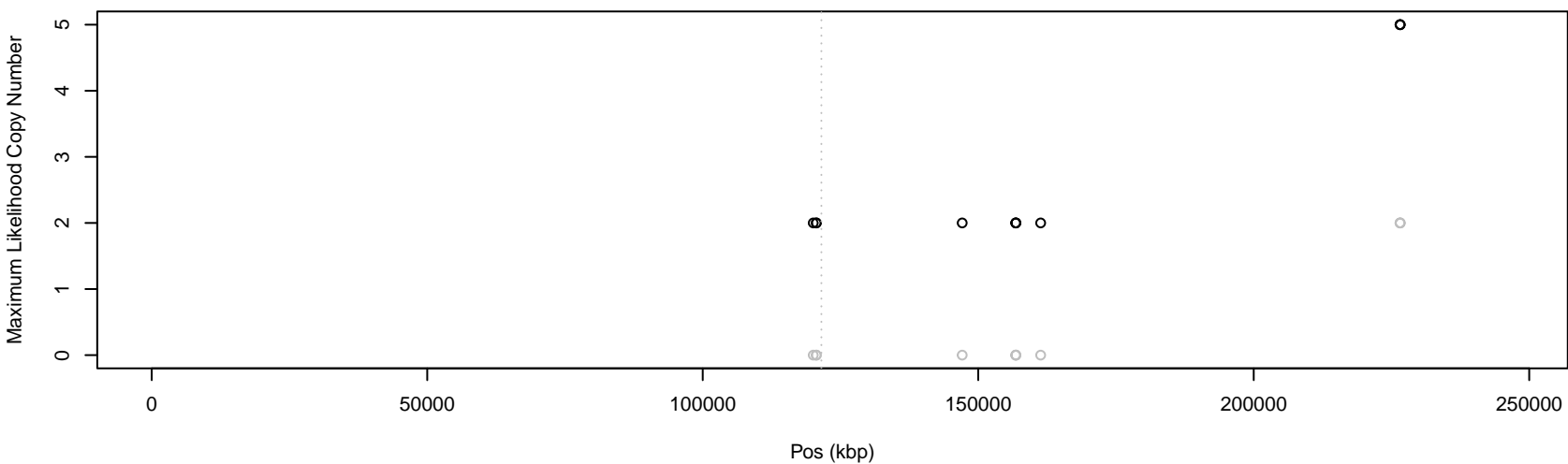
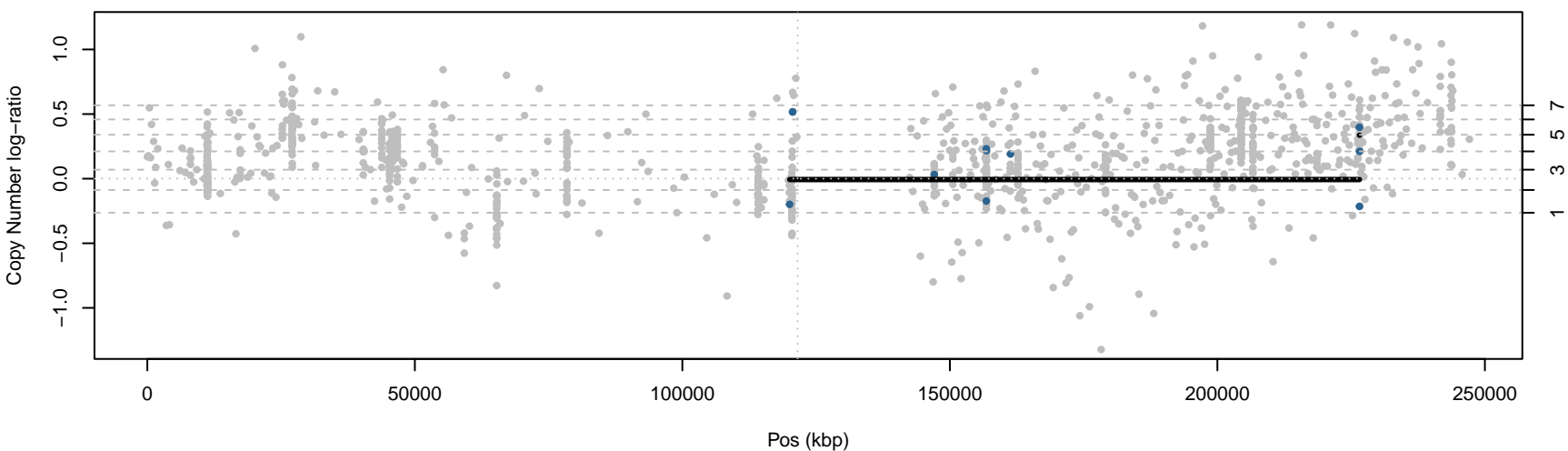
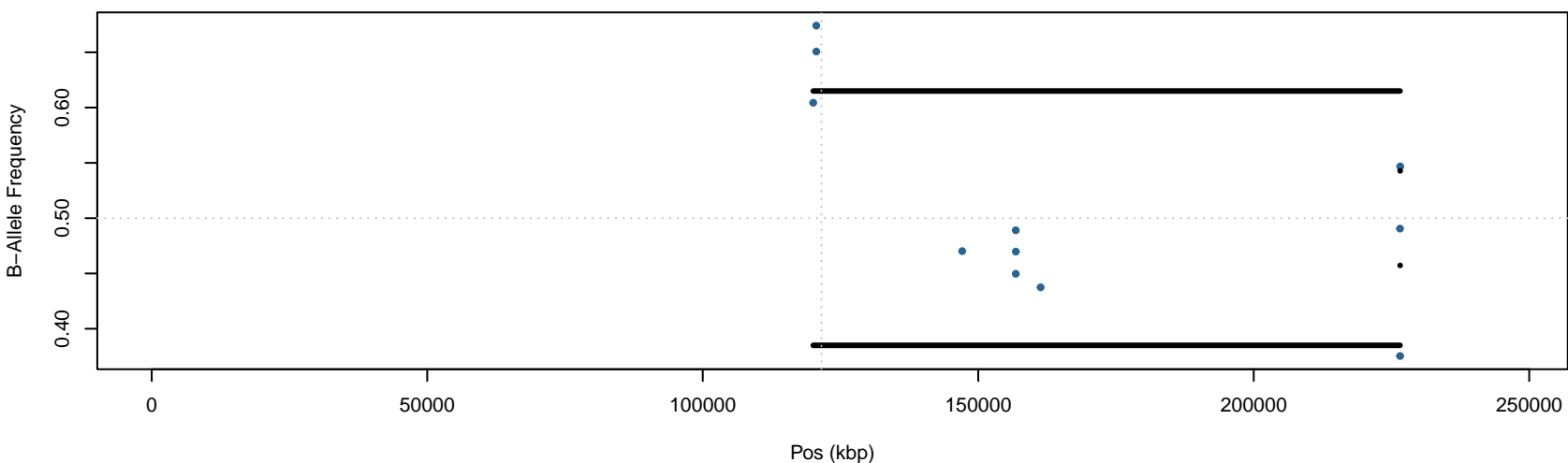
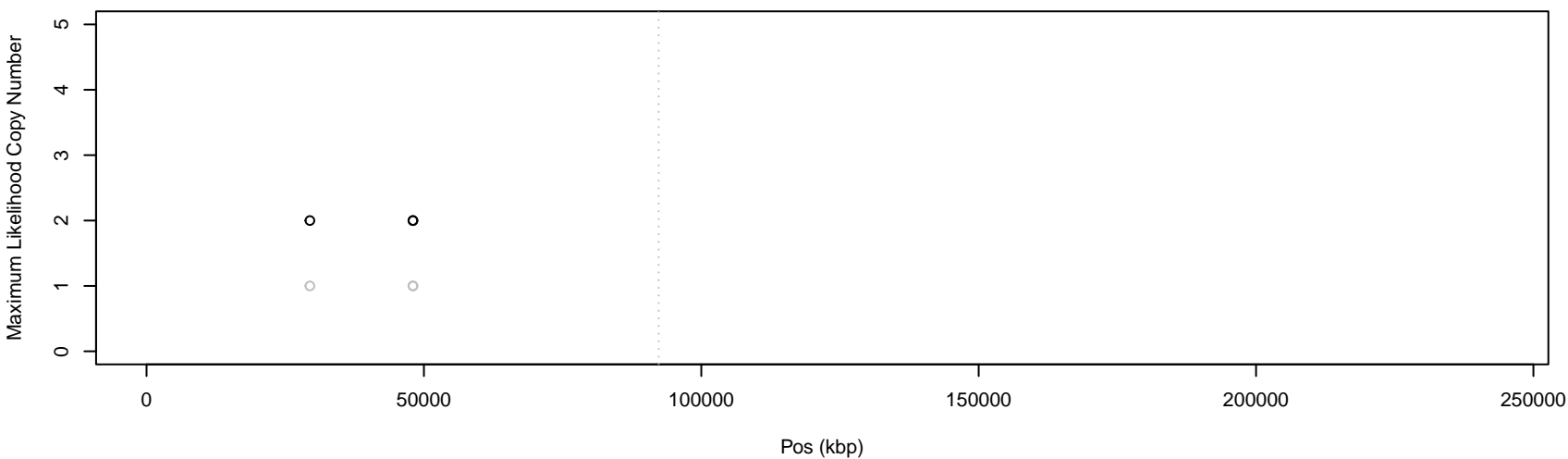
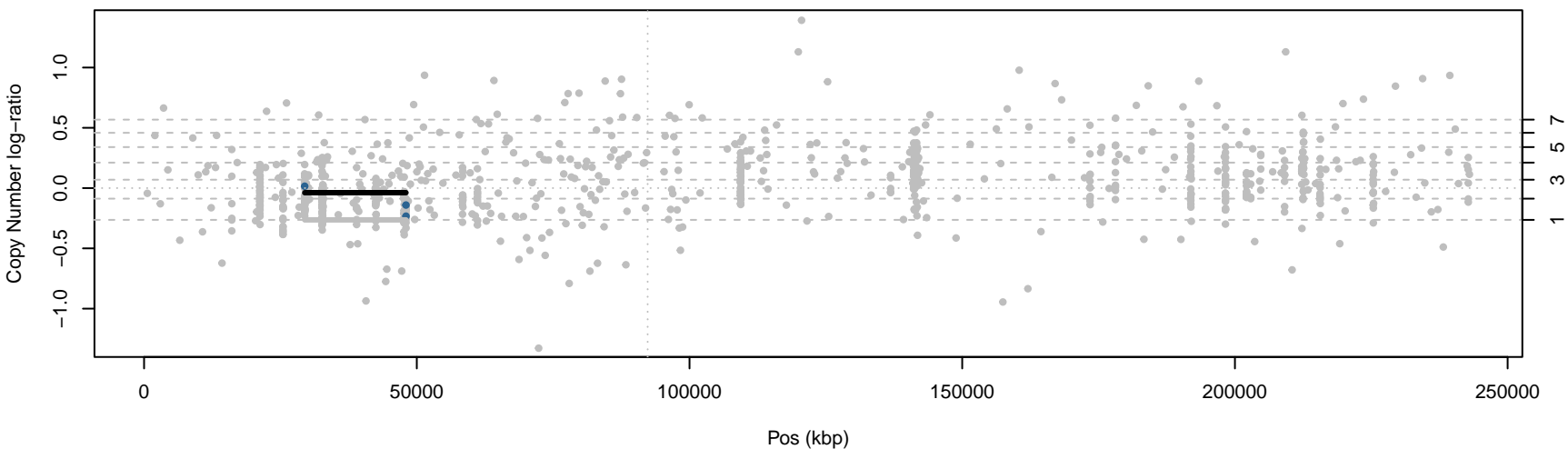
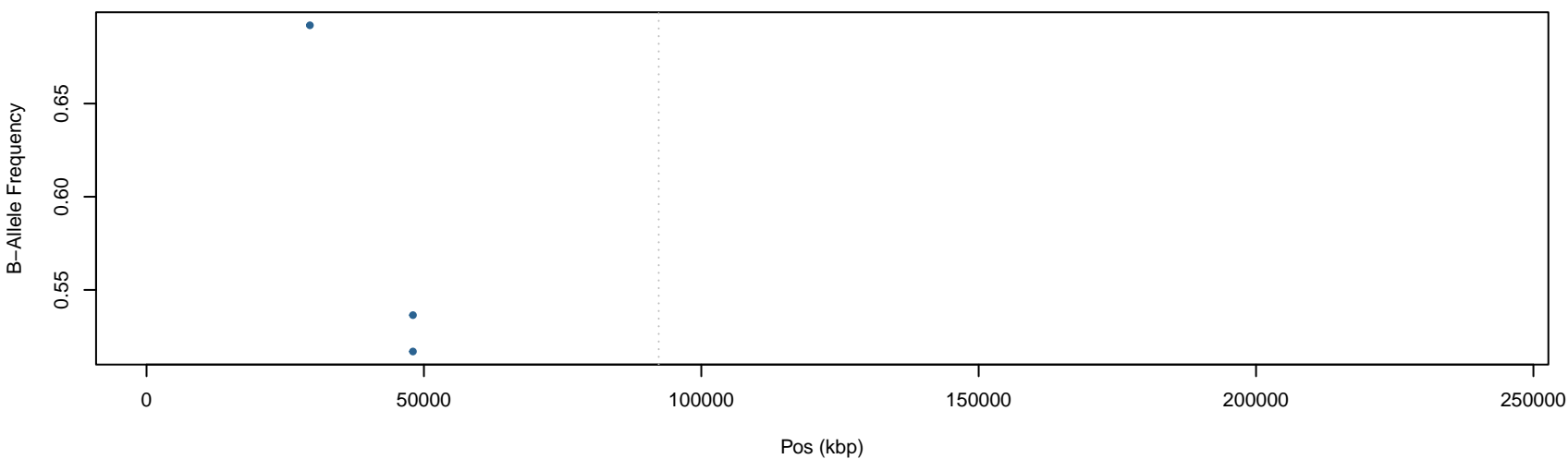


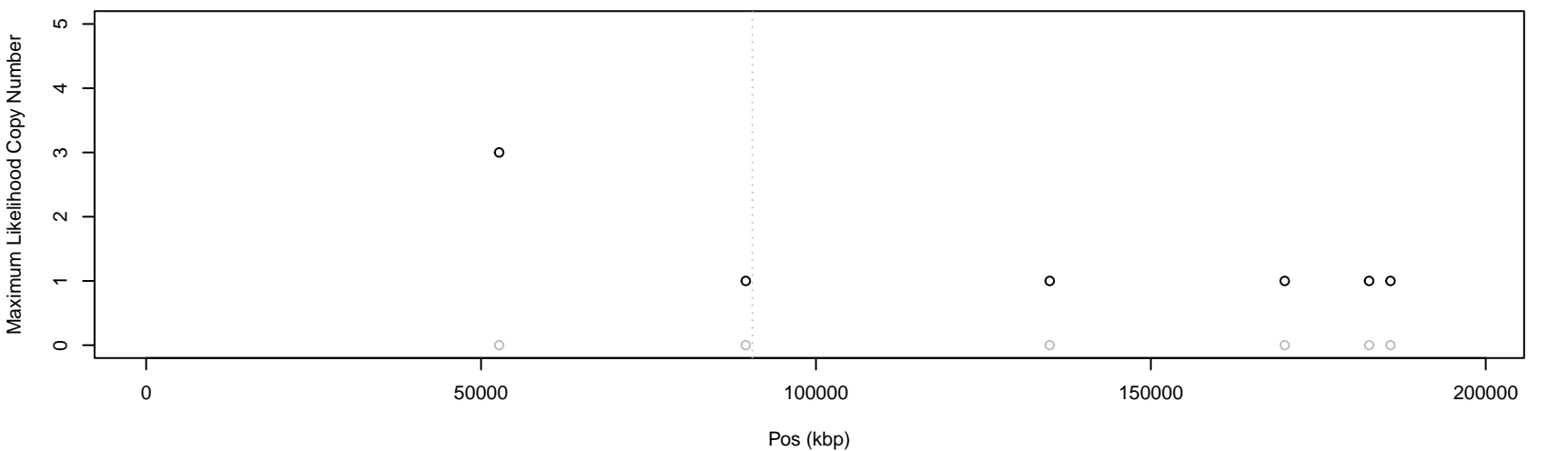
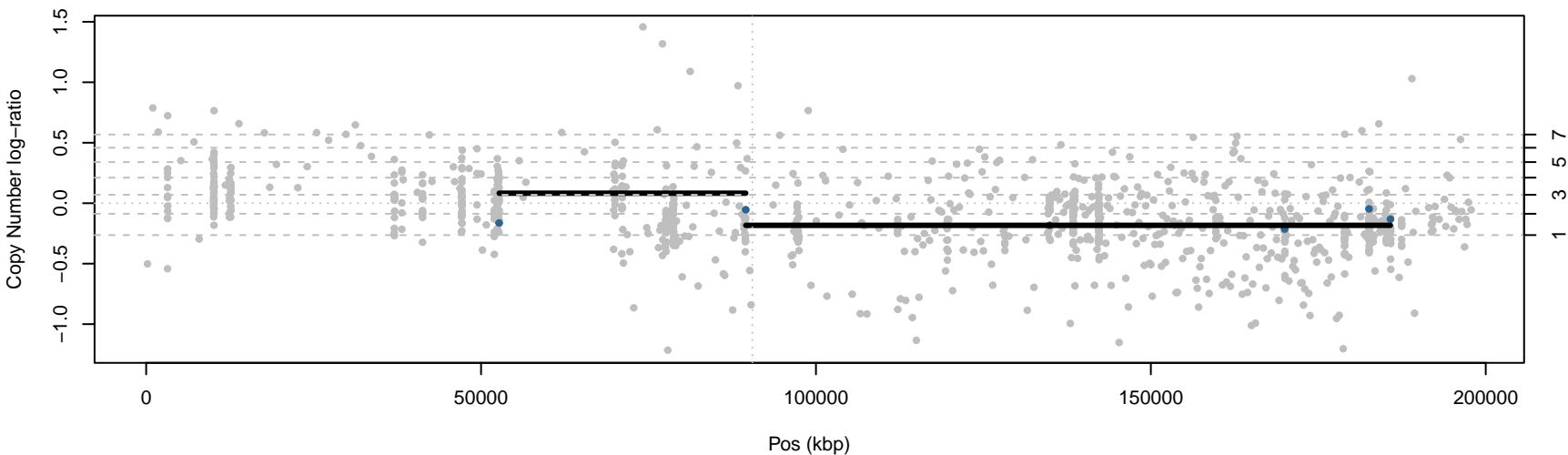
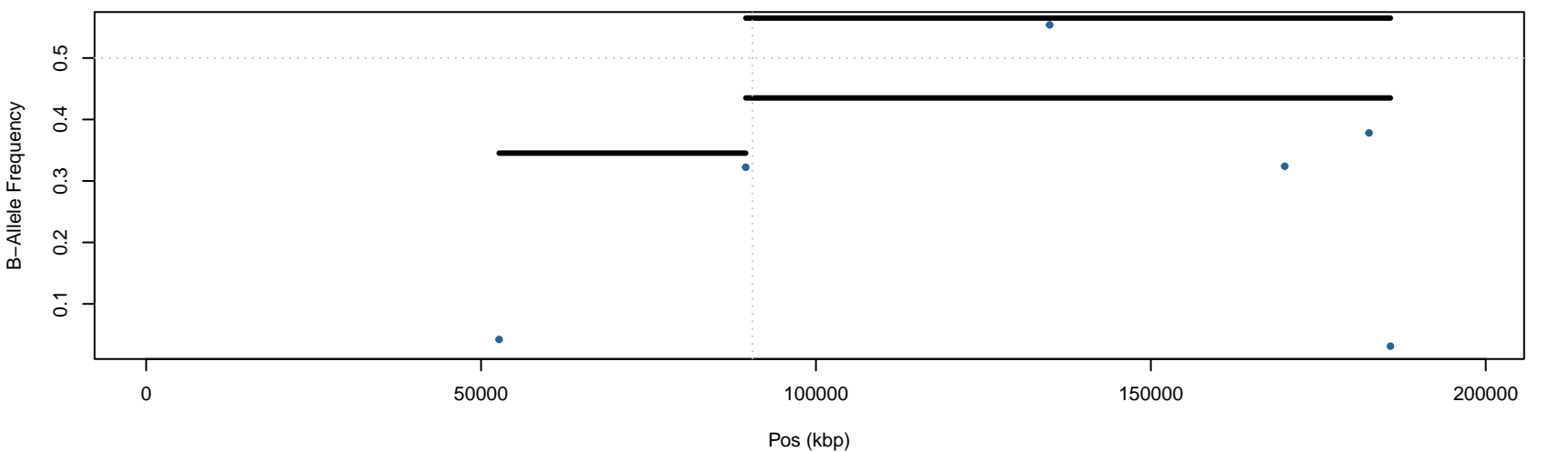
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 1



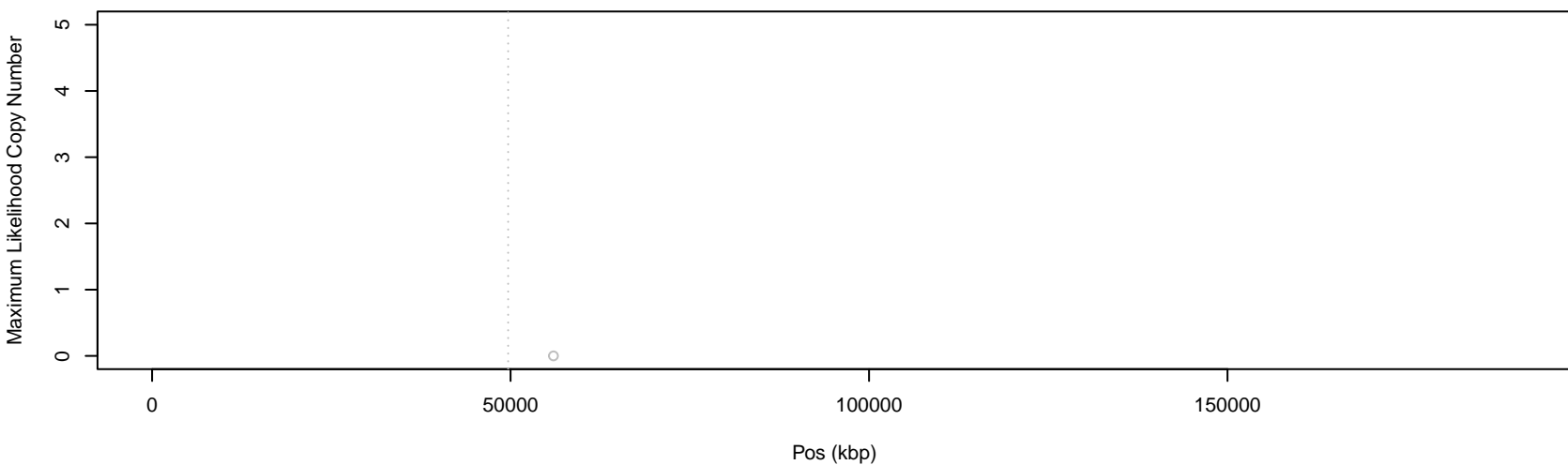
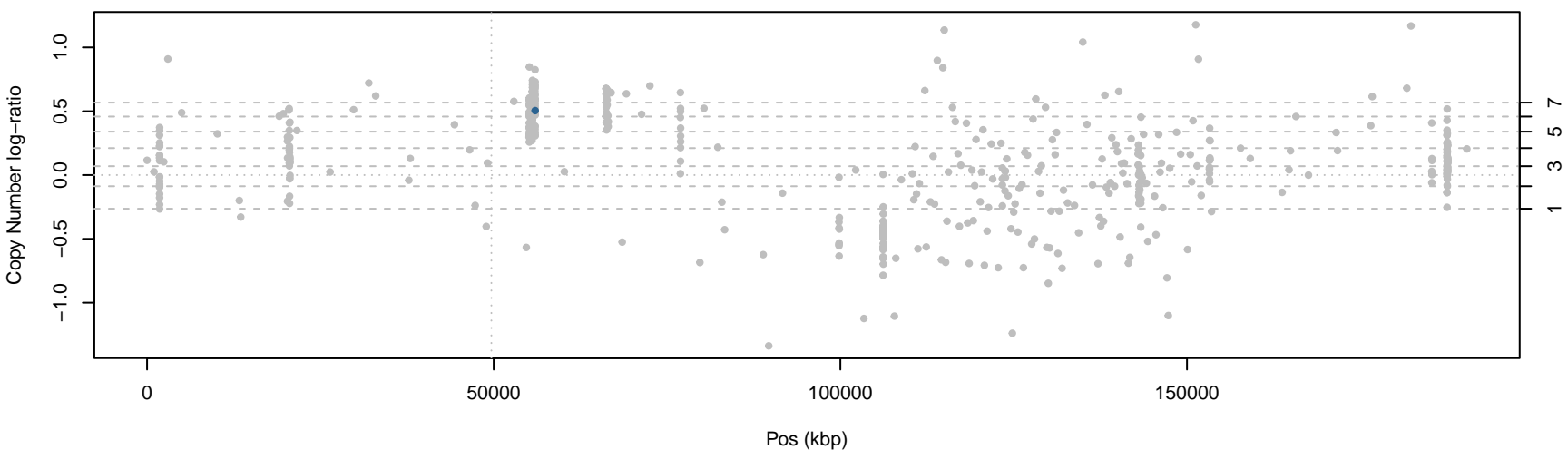
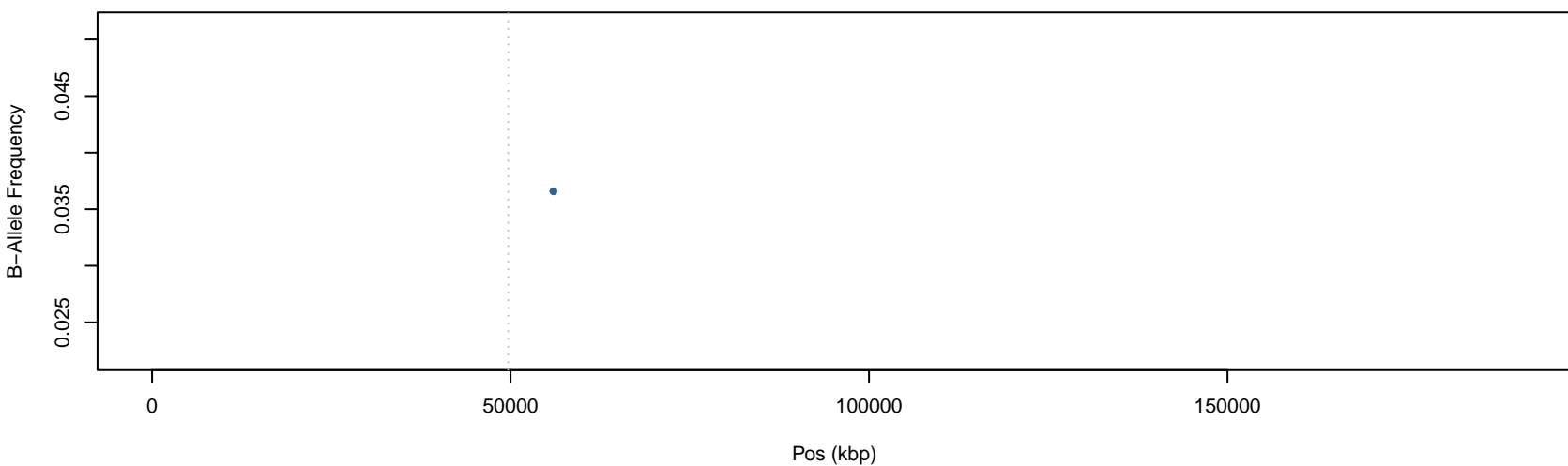
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 2



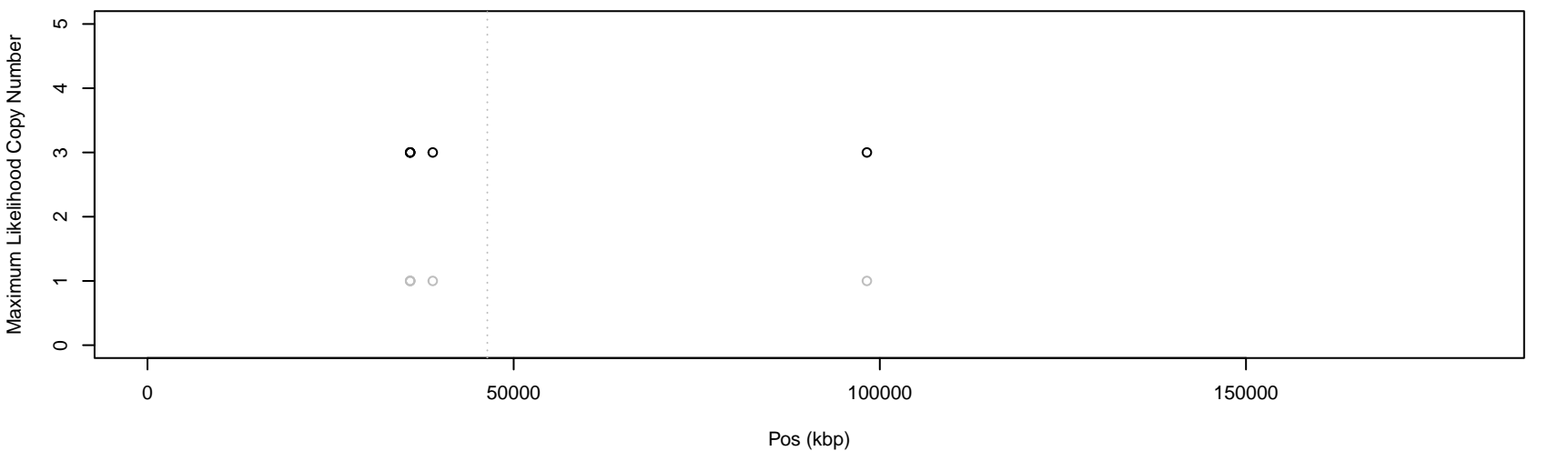
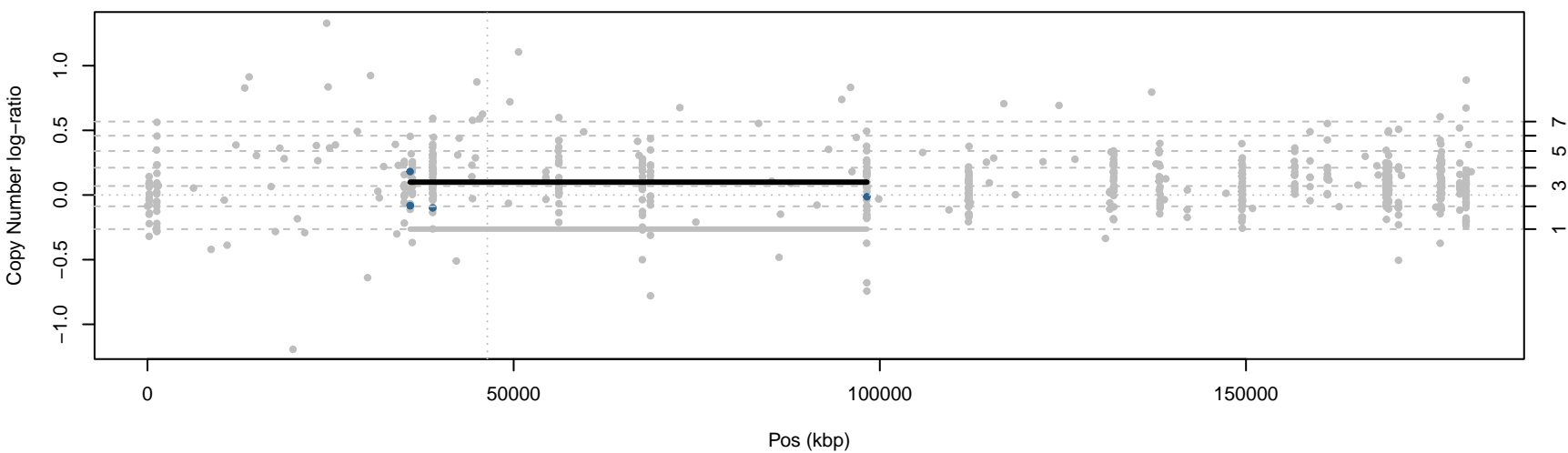
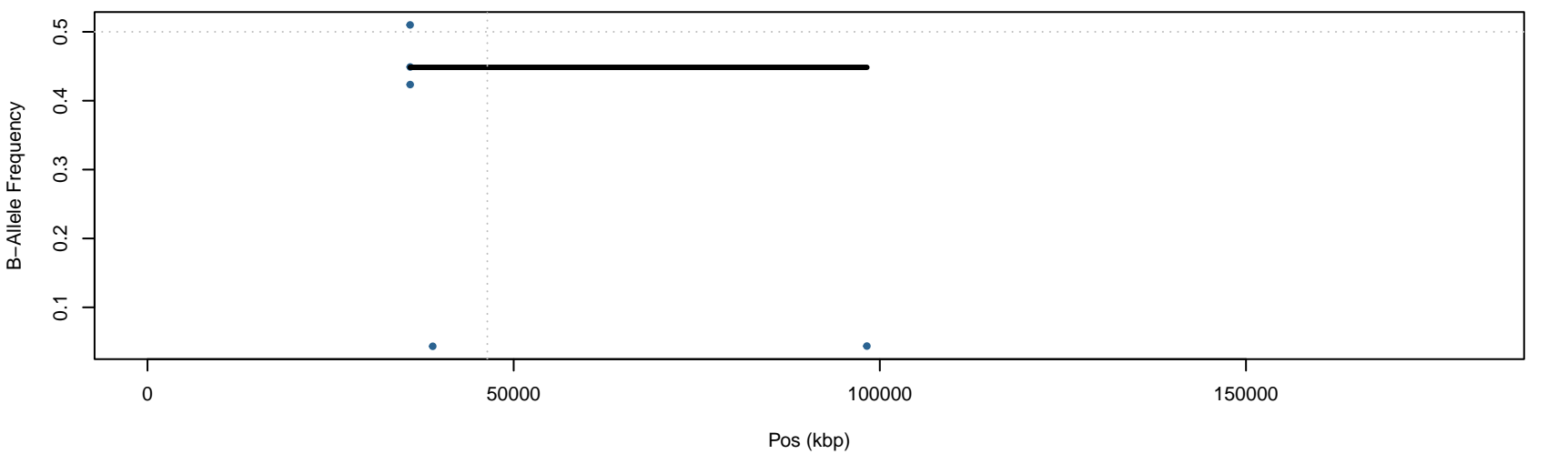
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 3



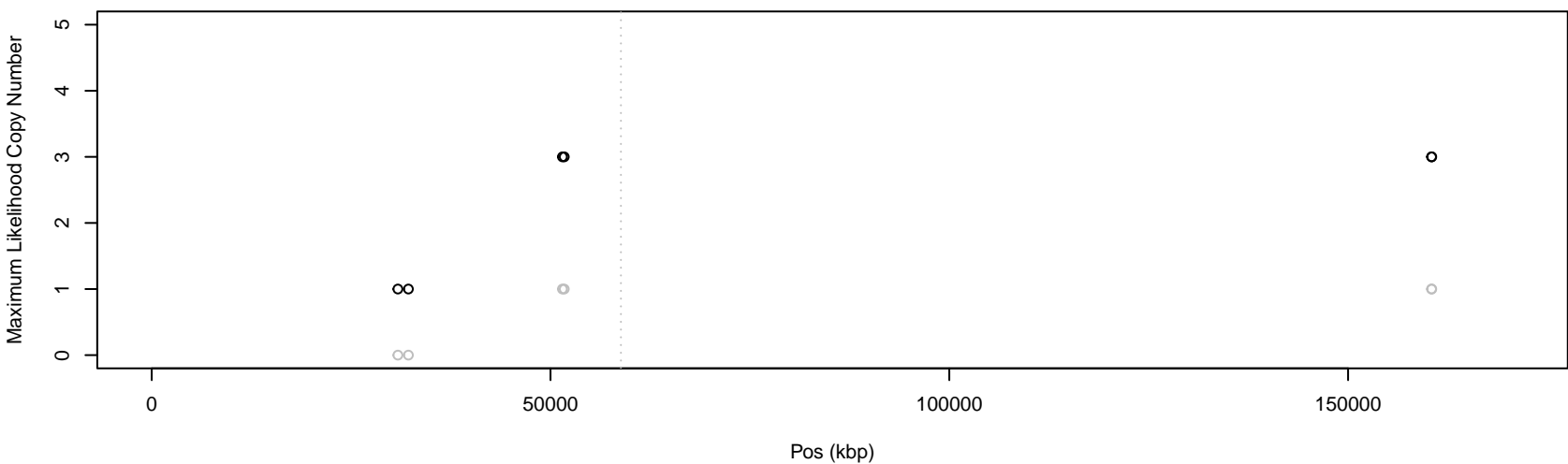
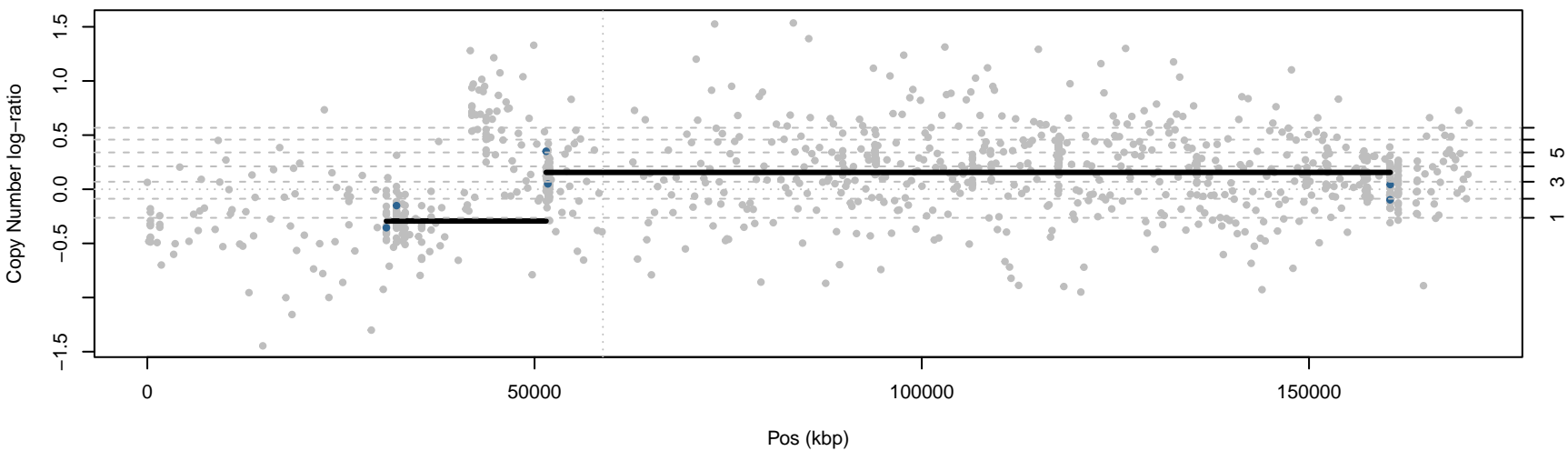
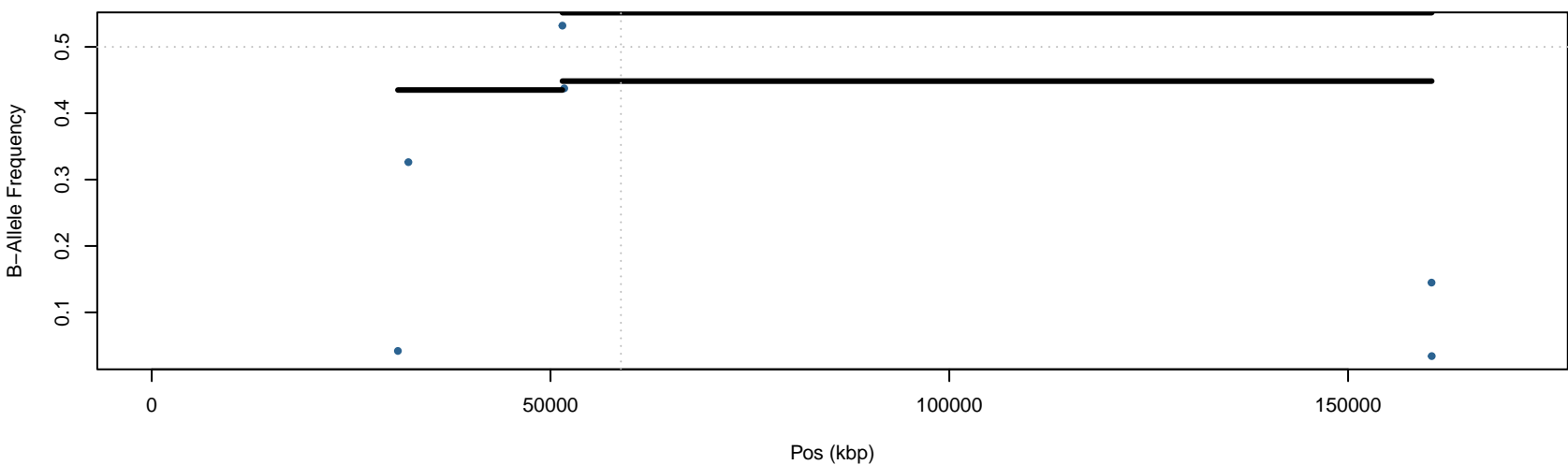
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 4



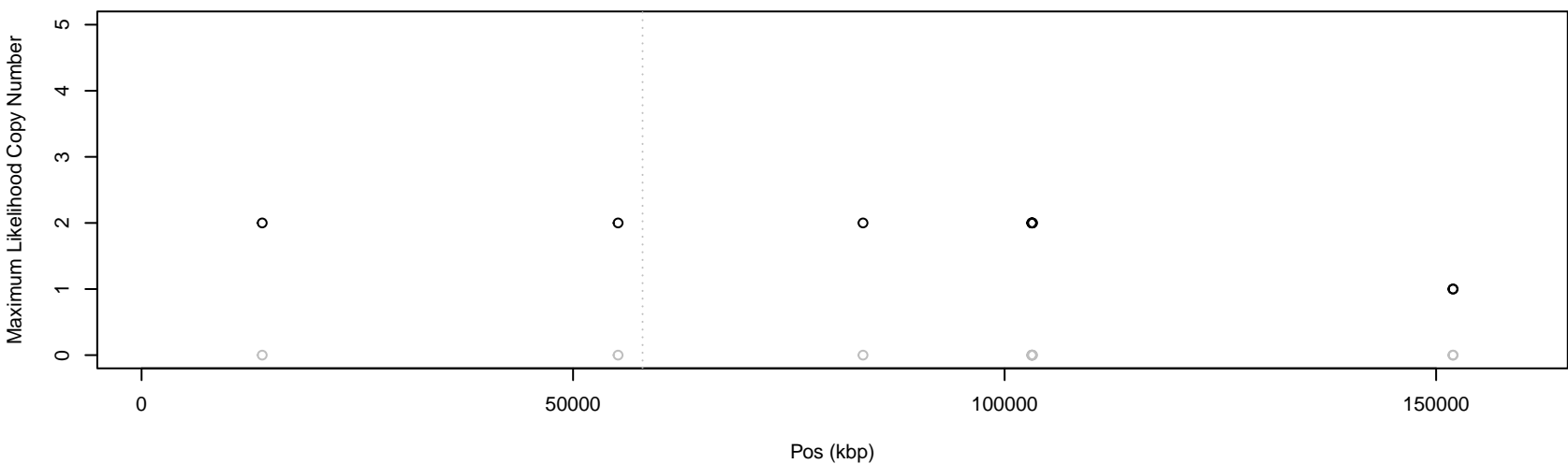
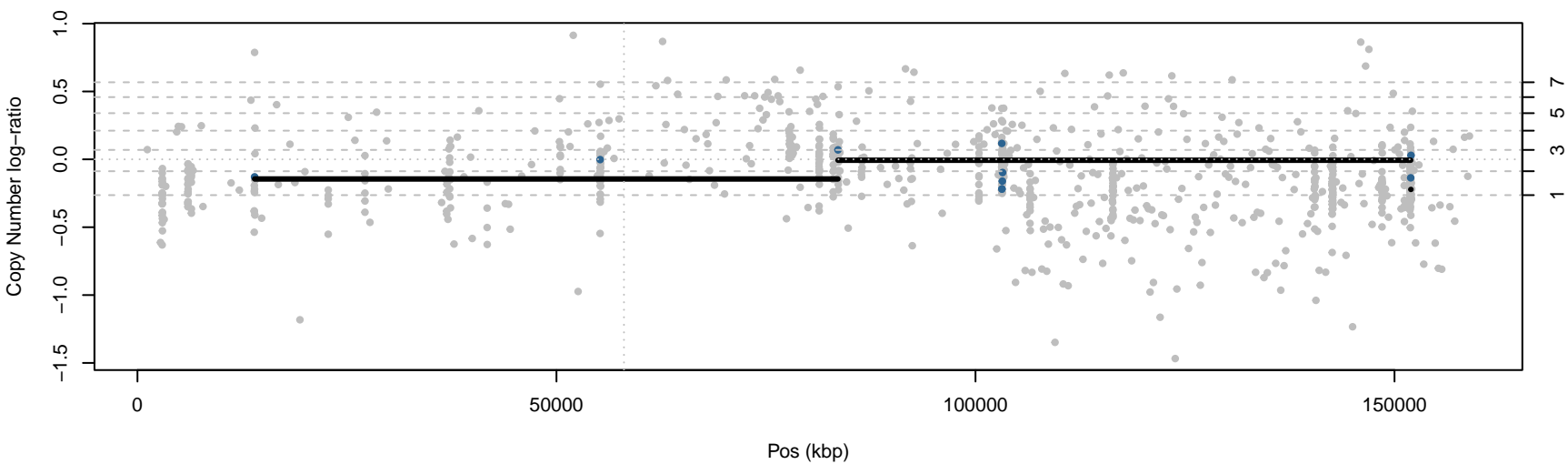
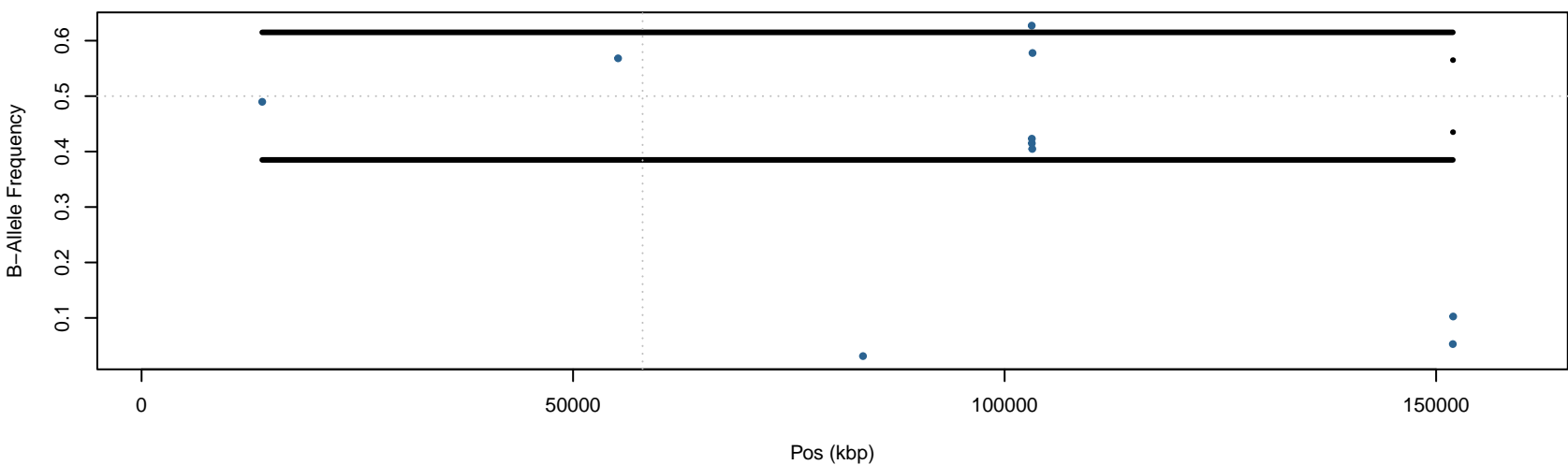
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 5



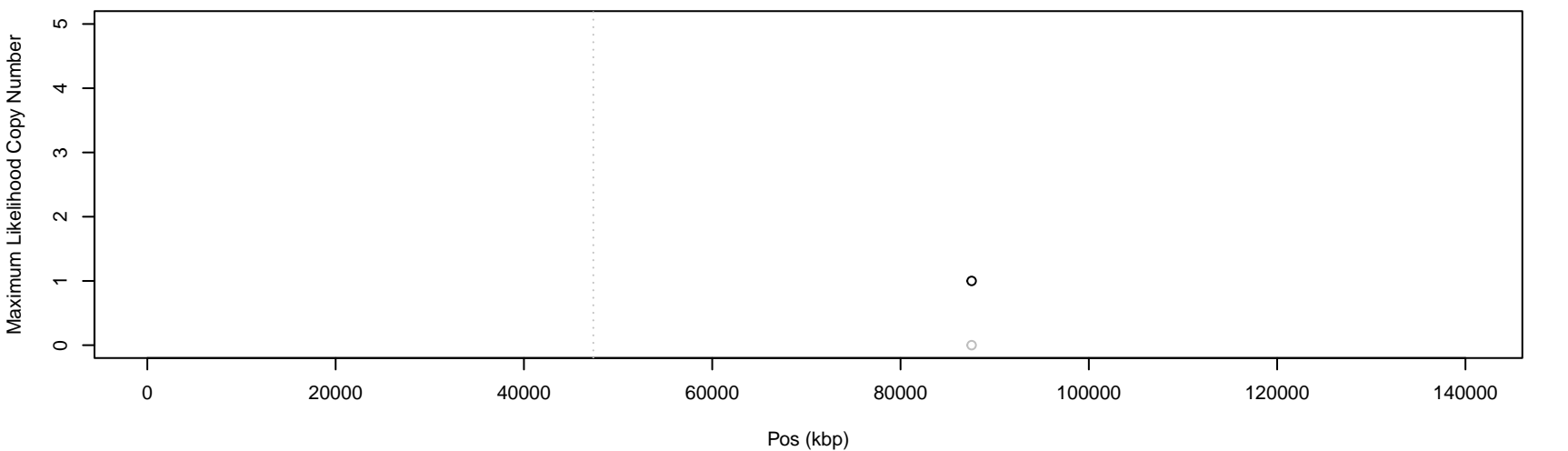
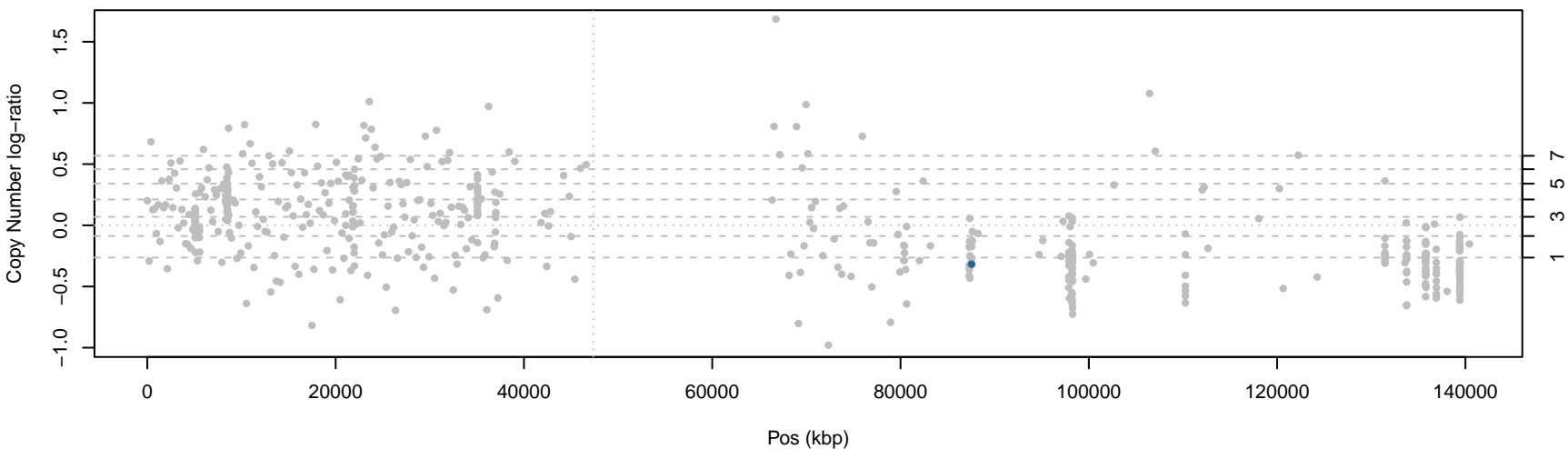
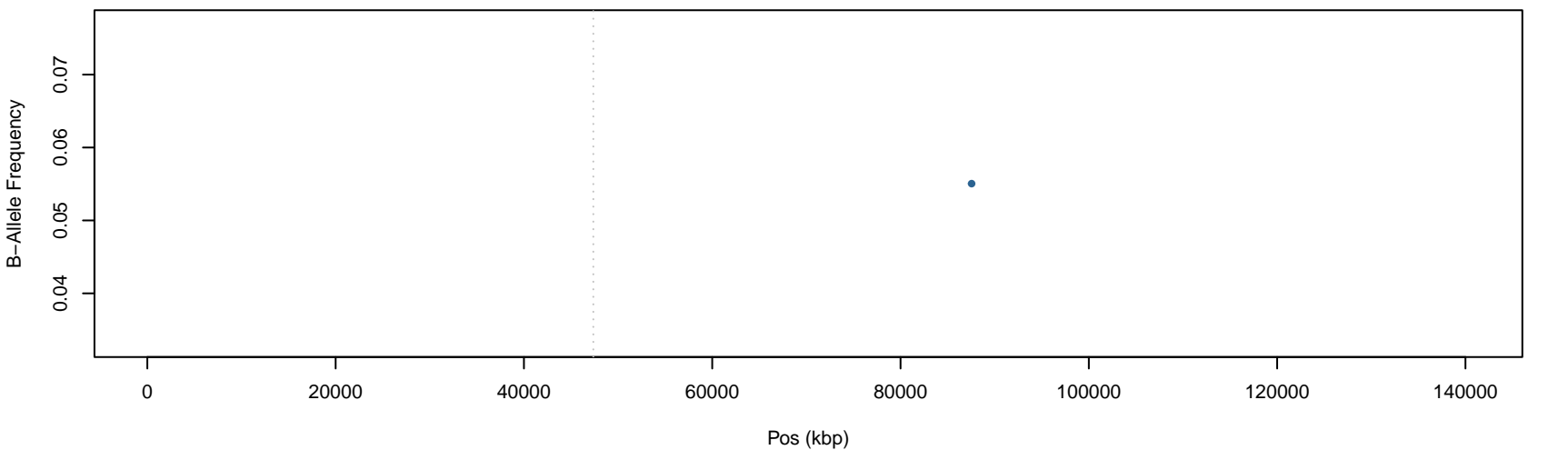
Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 6



Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 7

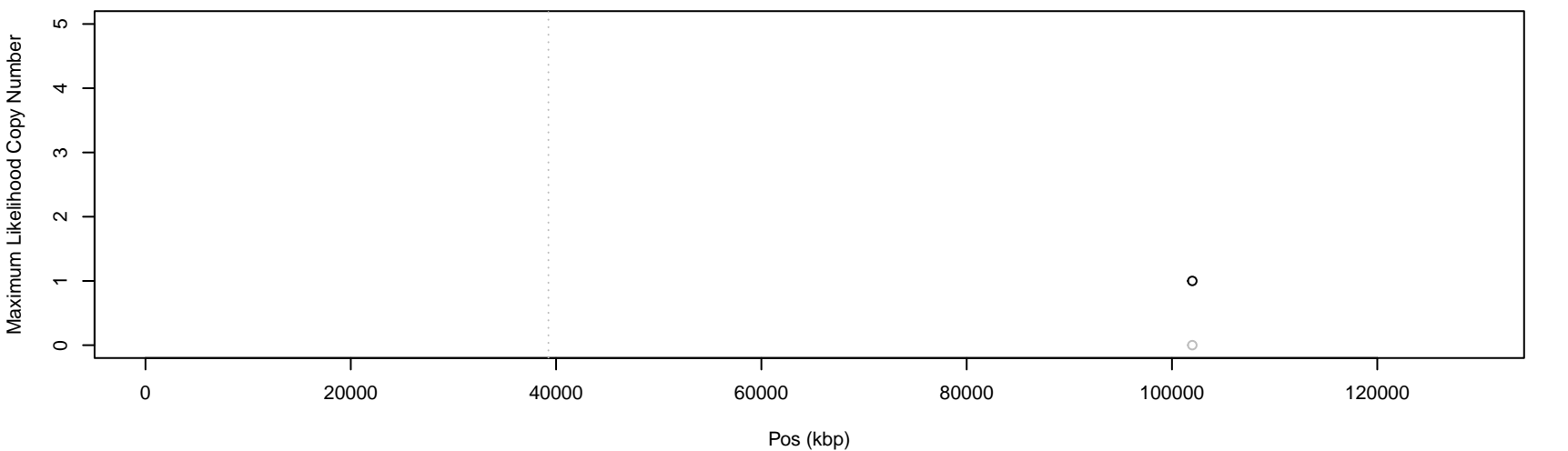
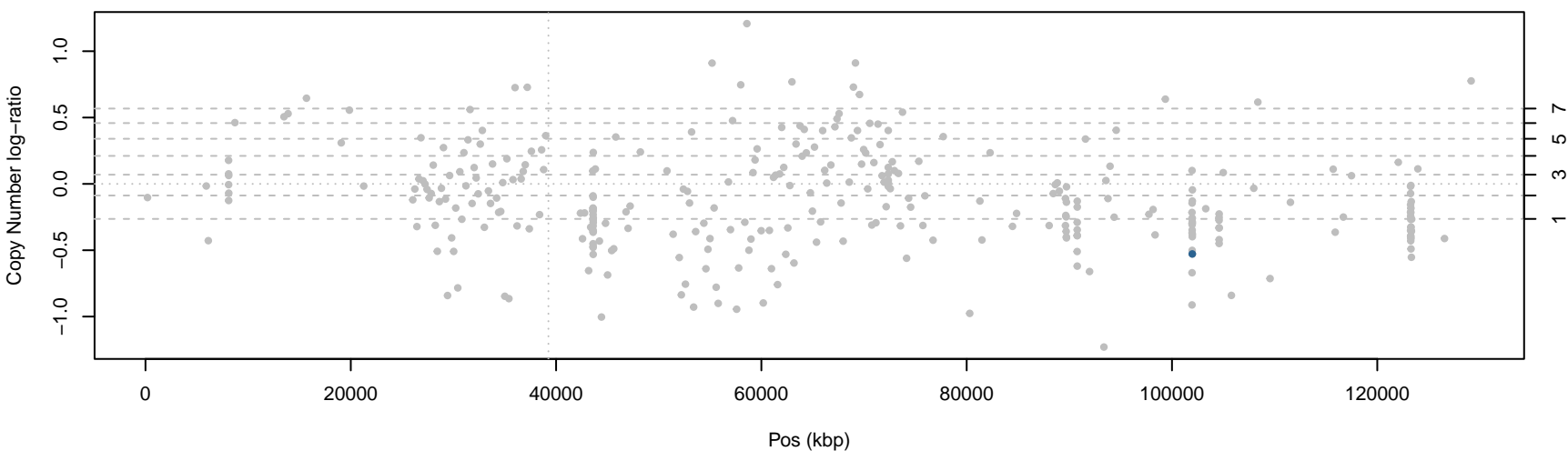
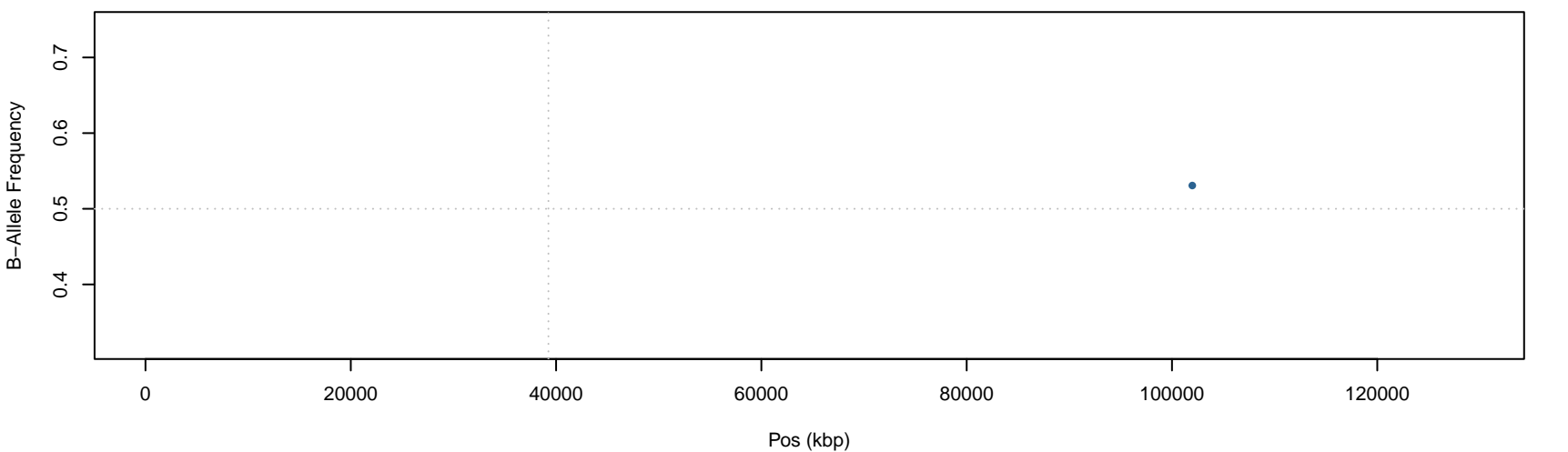


Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 9





Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 10



Purity: 0.23 Tumor ploidy: 2.544 SNV log-likelihood: -65.21 GoF: 91.7% Mean coverage: 542;511 Chromosome: 19

