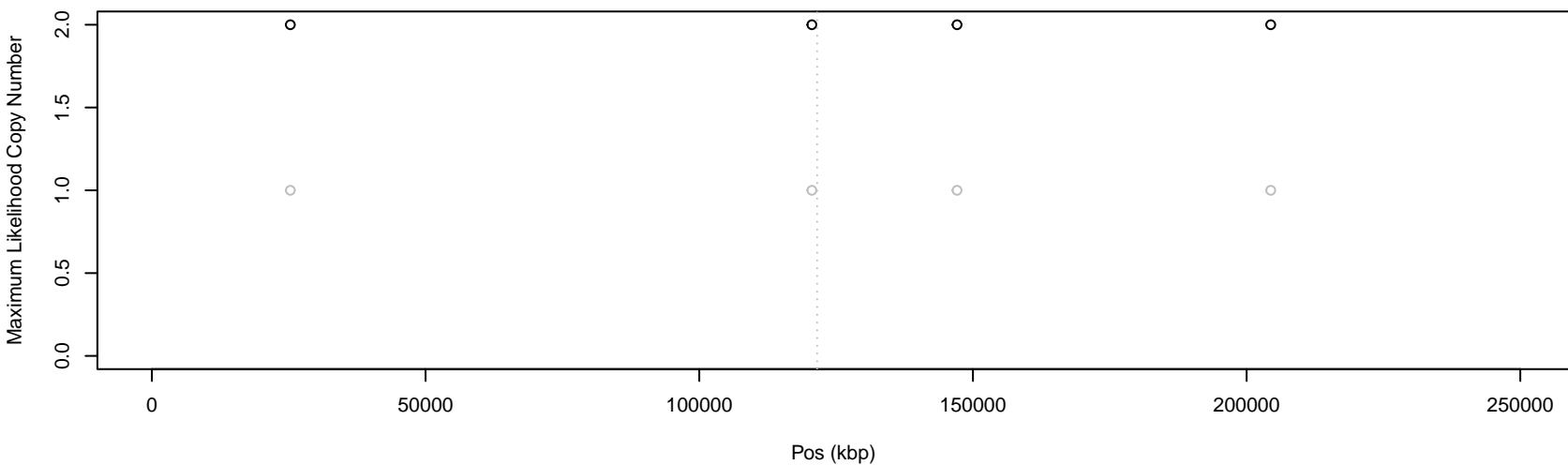
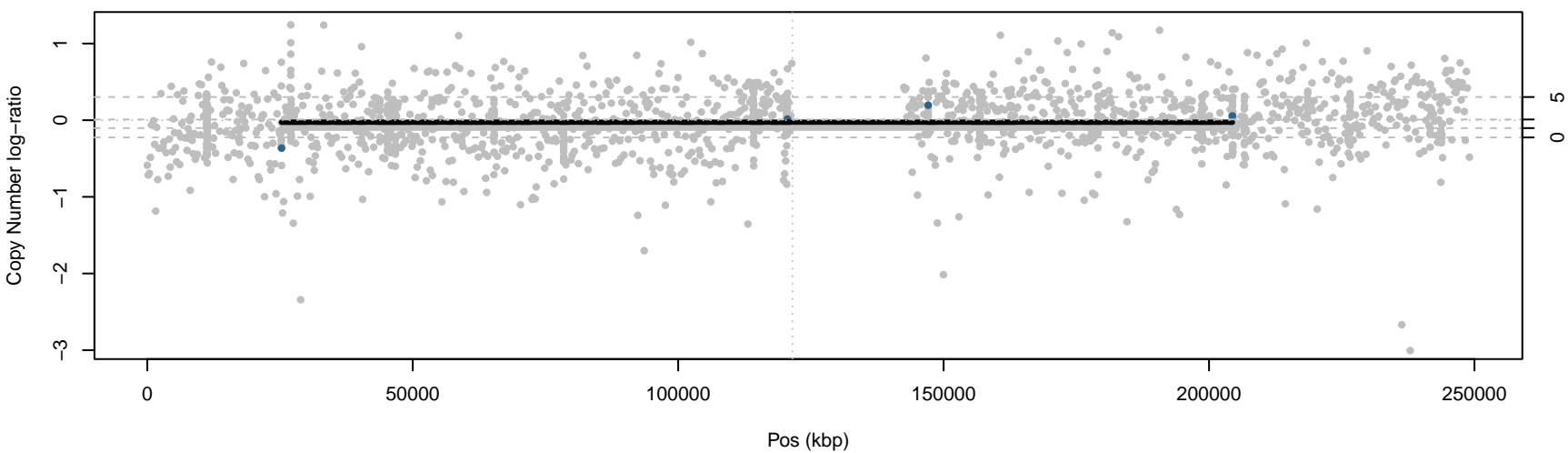
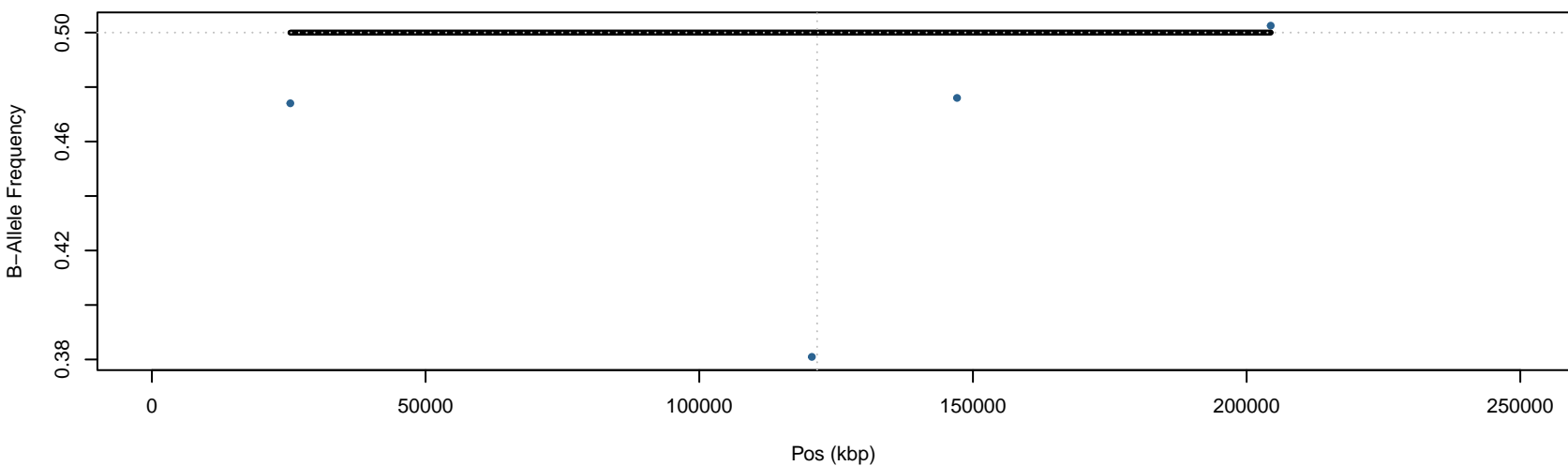
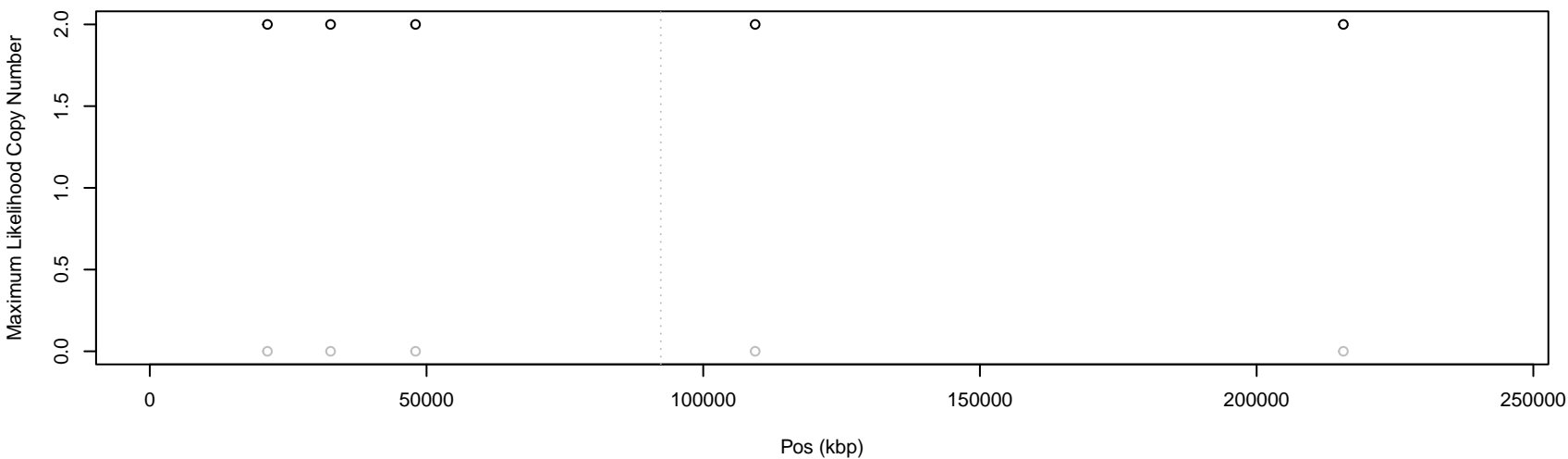
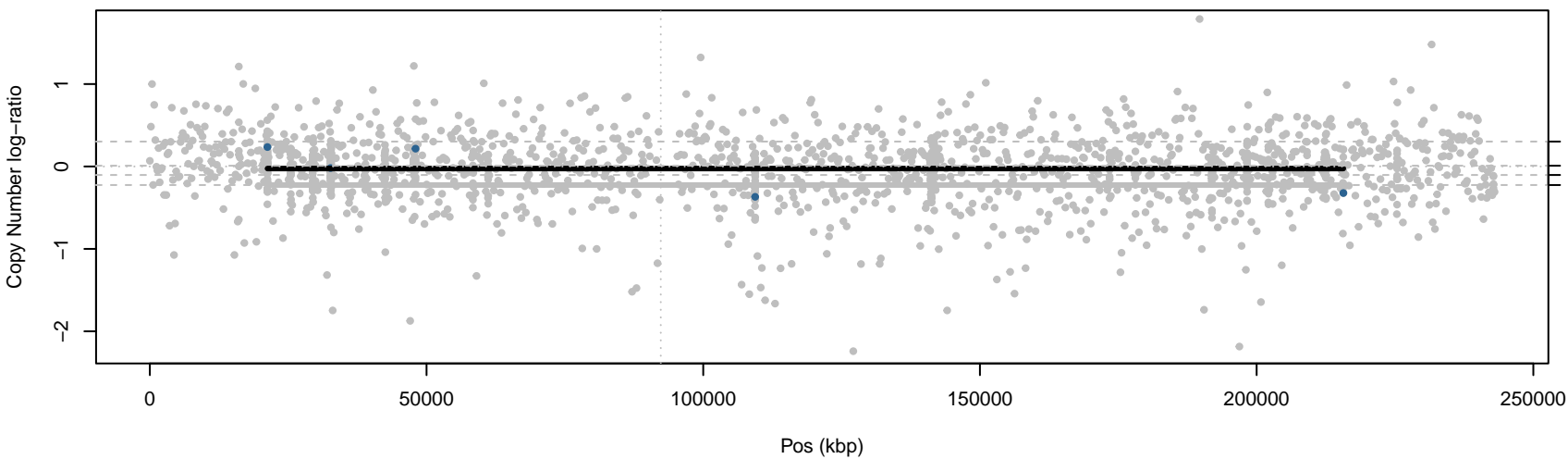
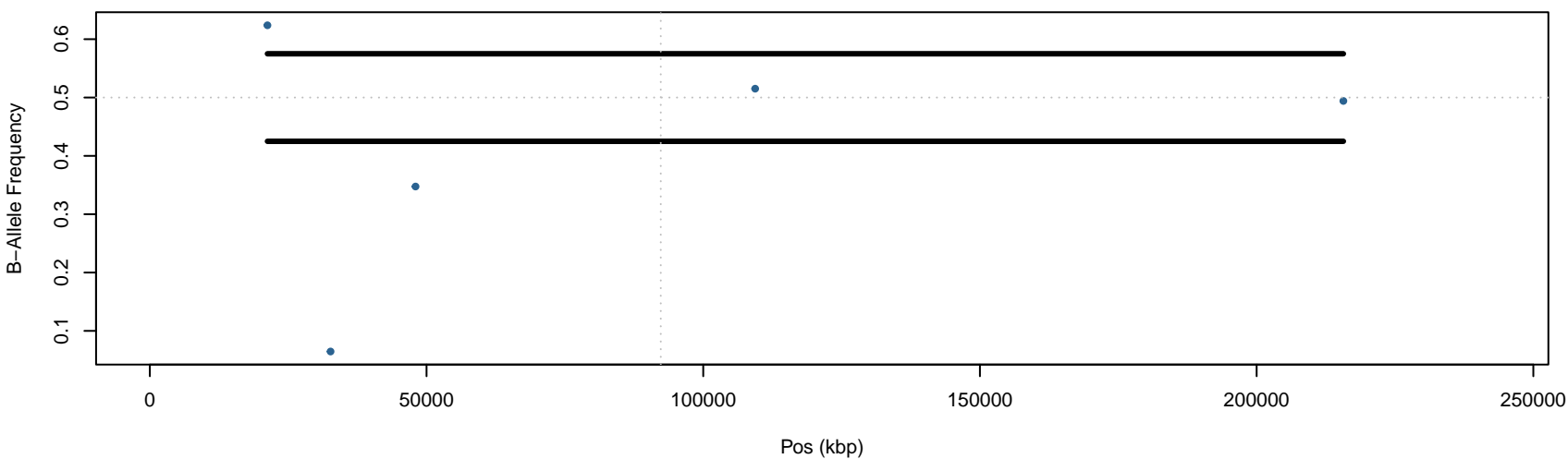


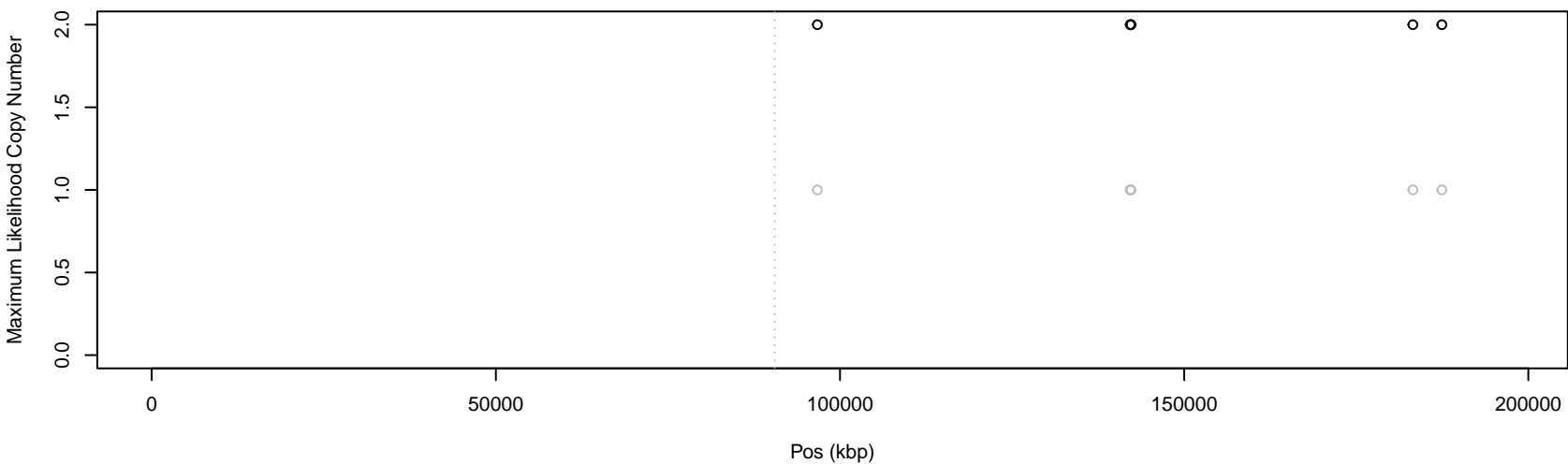
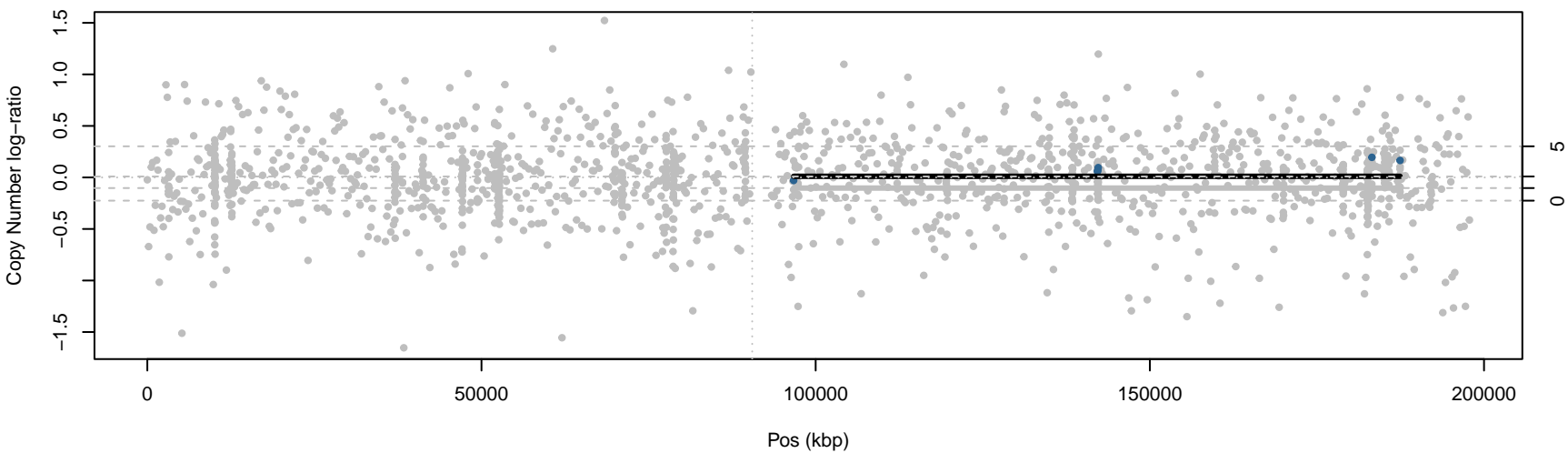
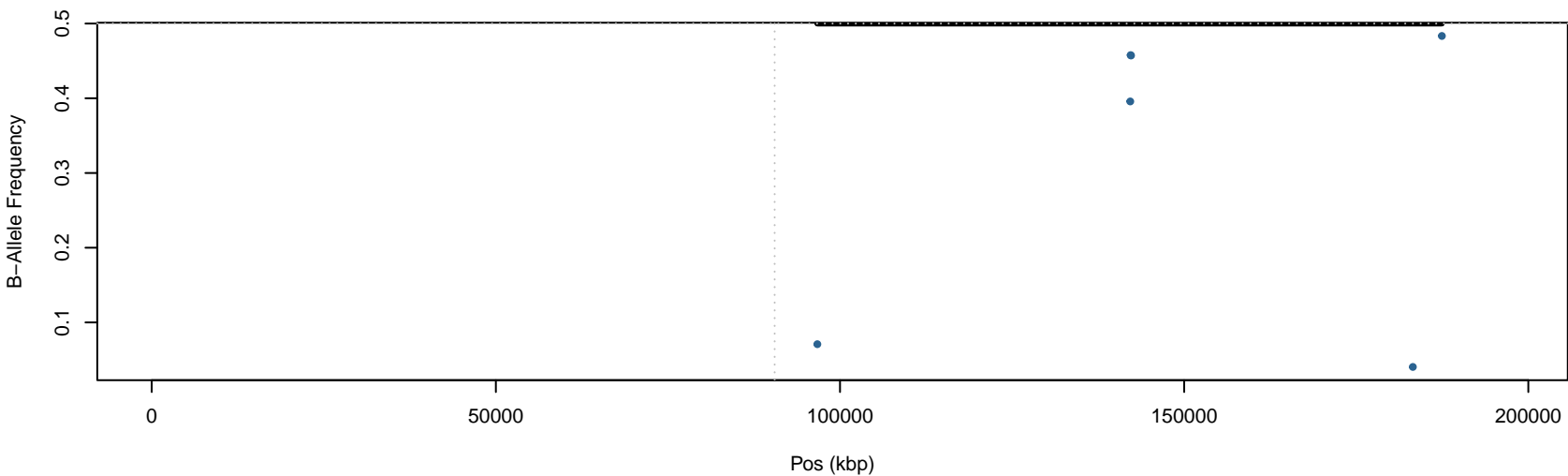
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 1



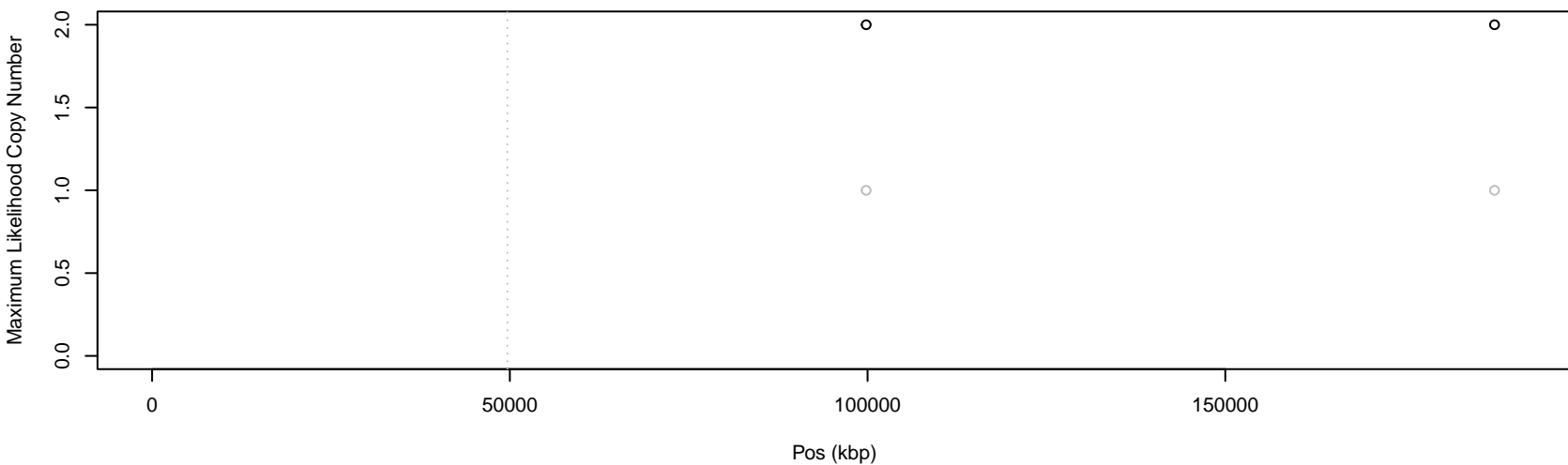
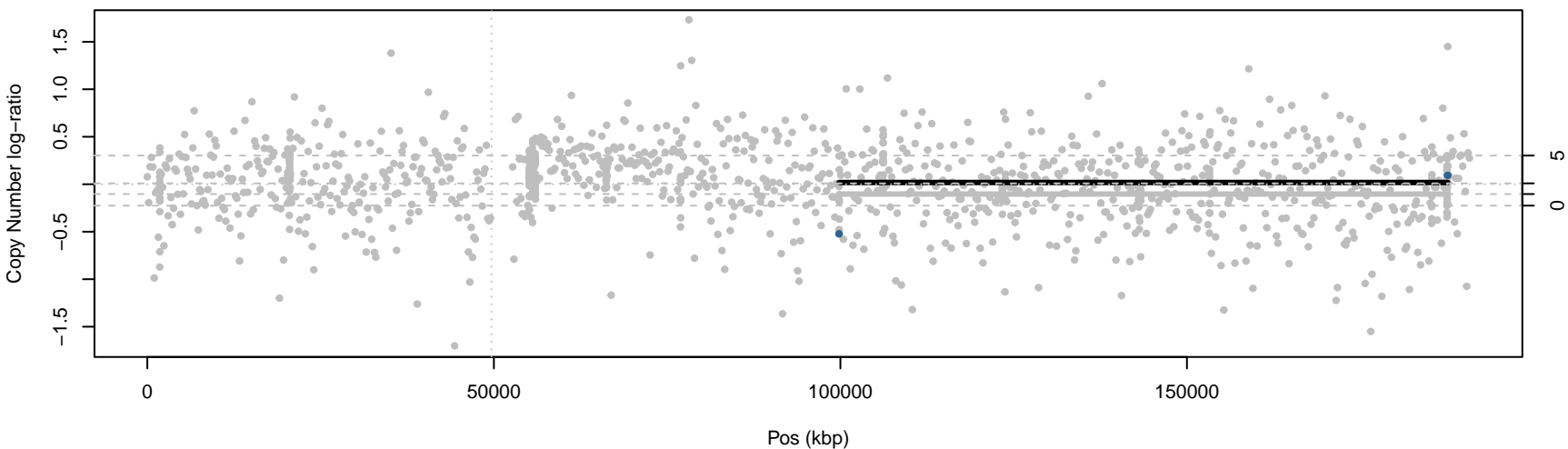
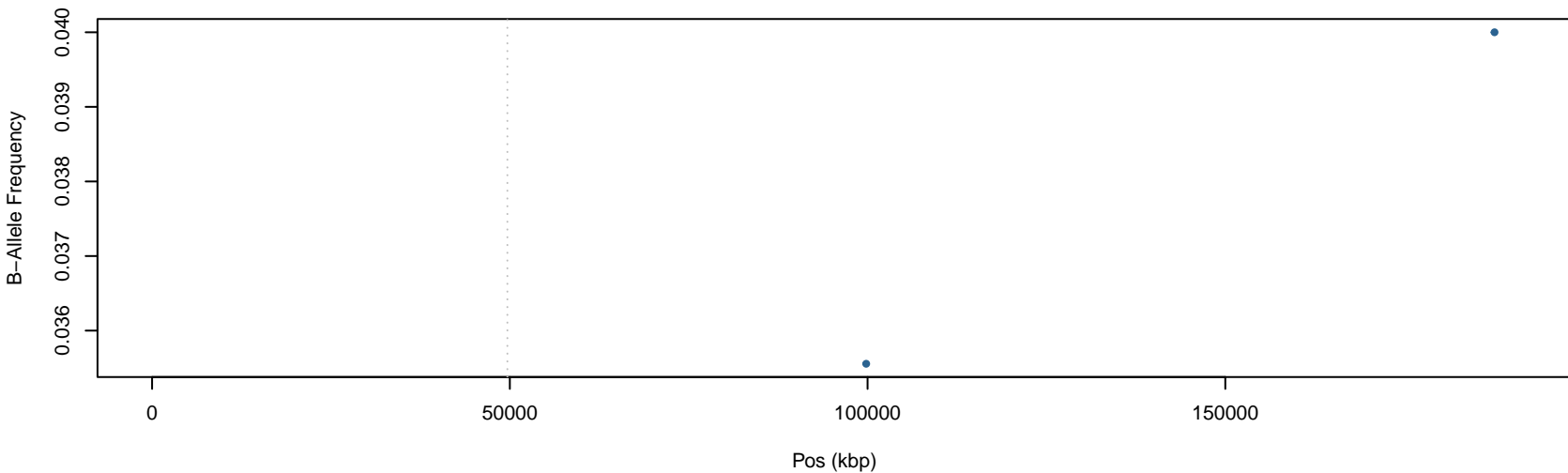
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 2



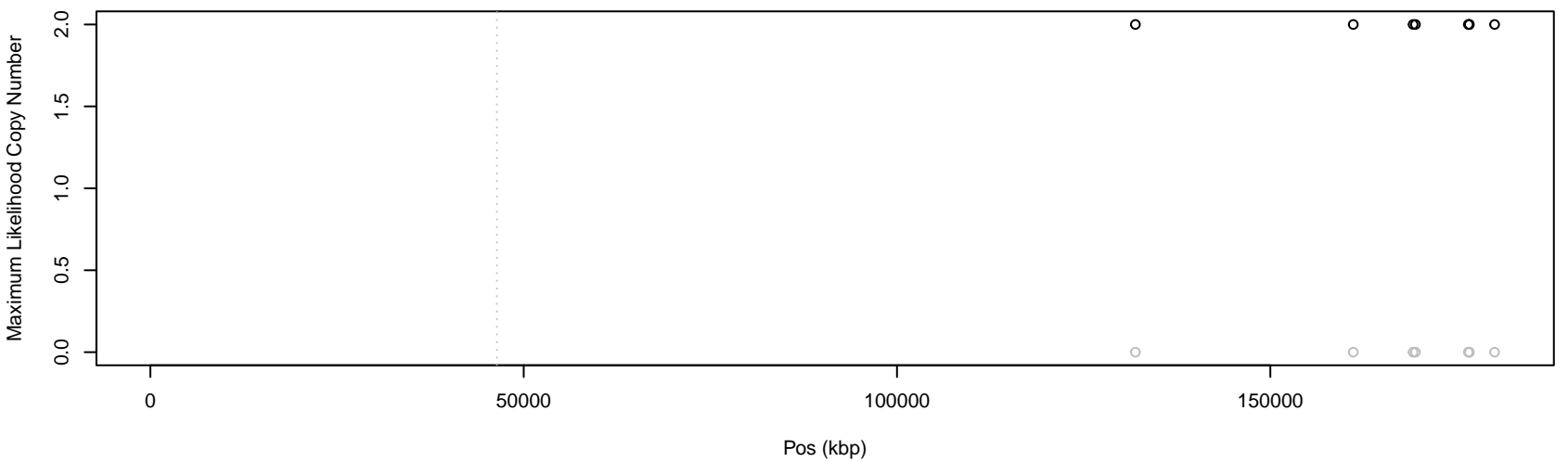
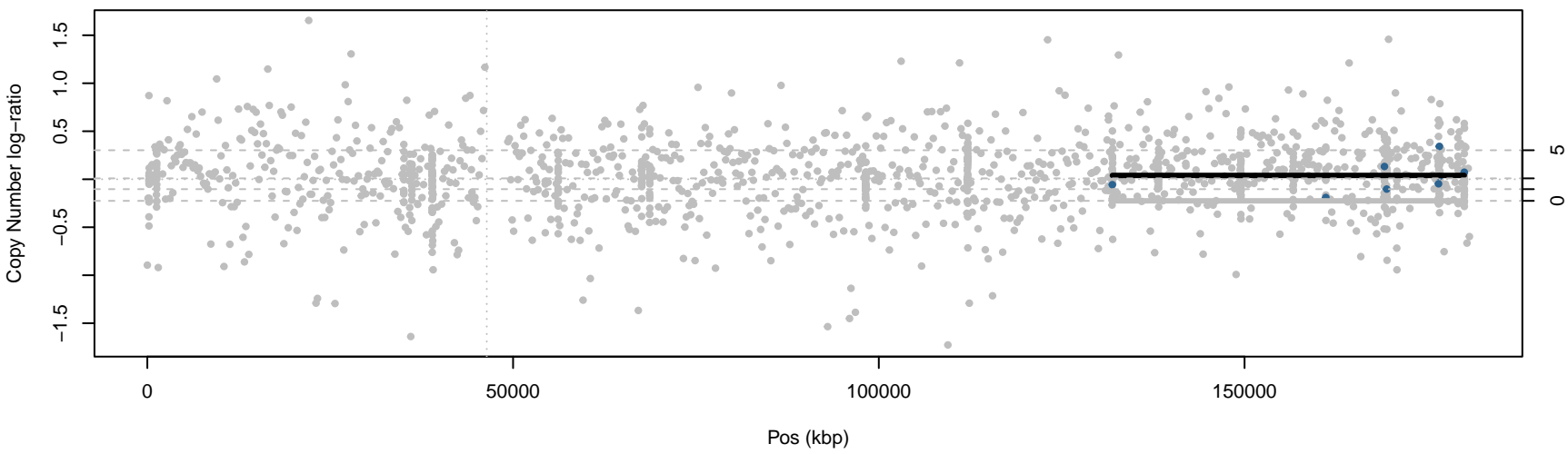
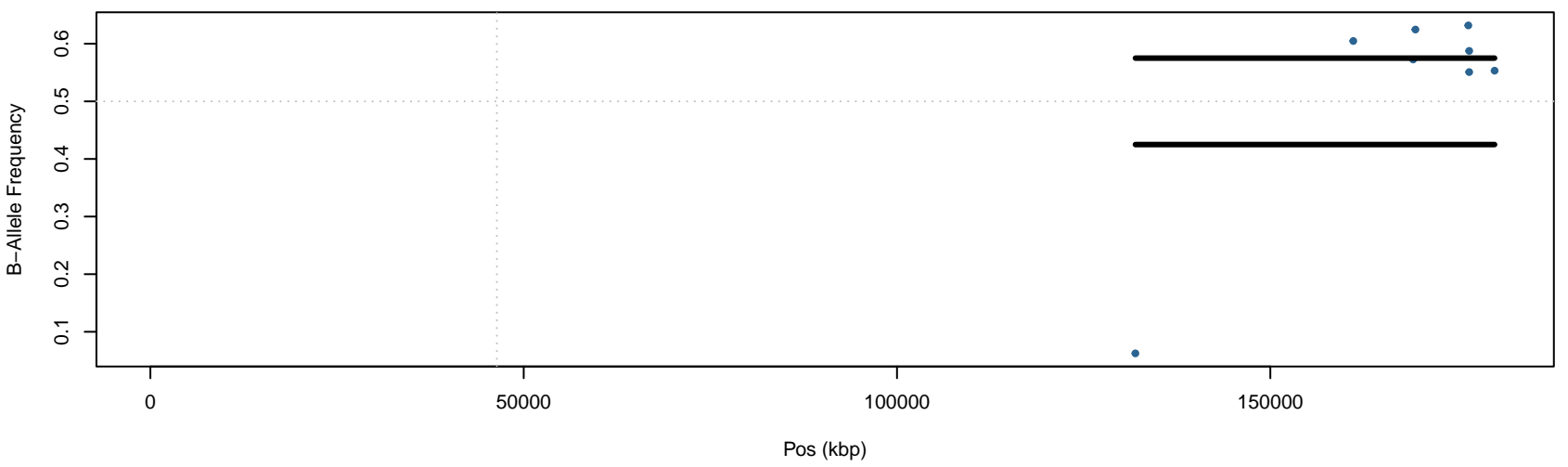
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 3



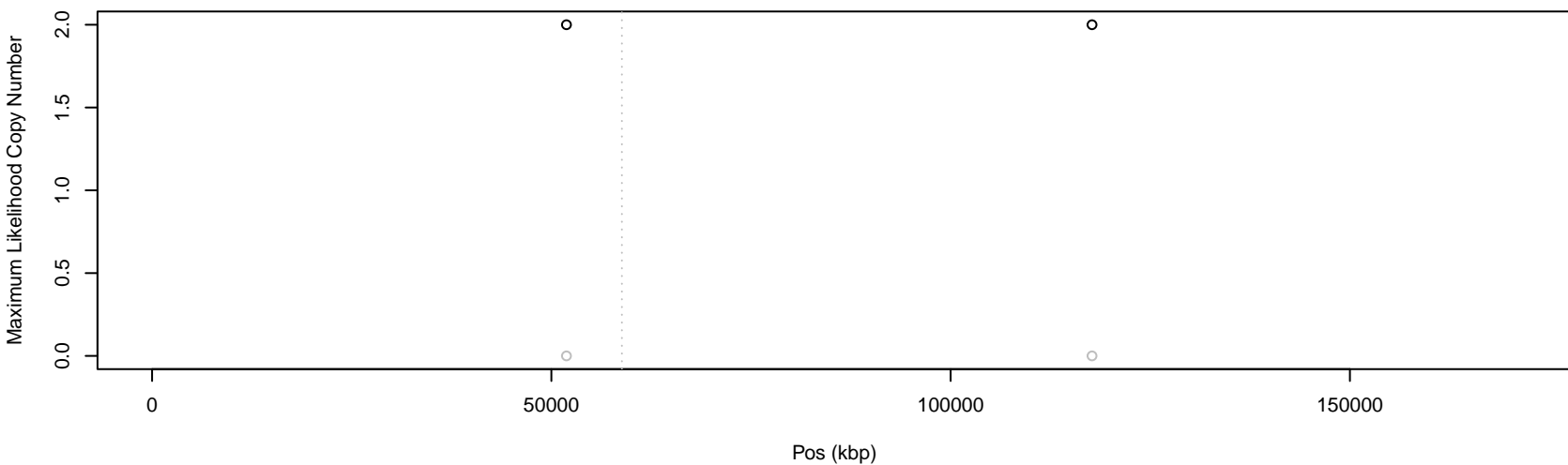
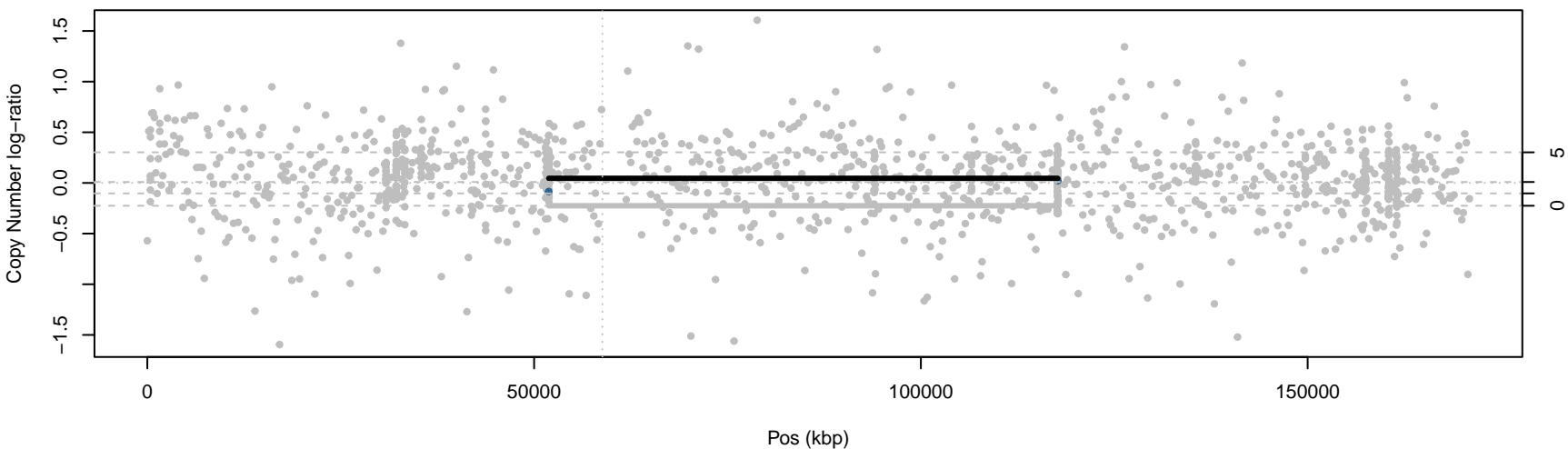
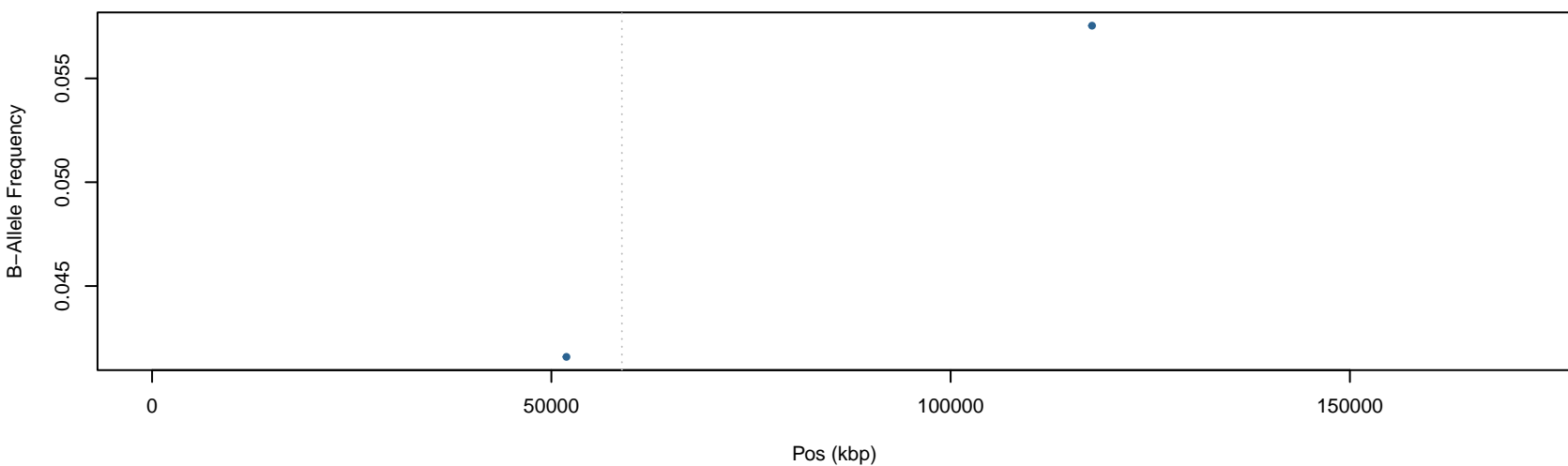
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 4



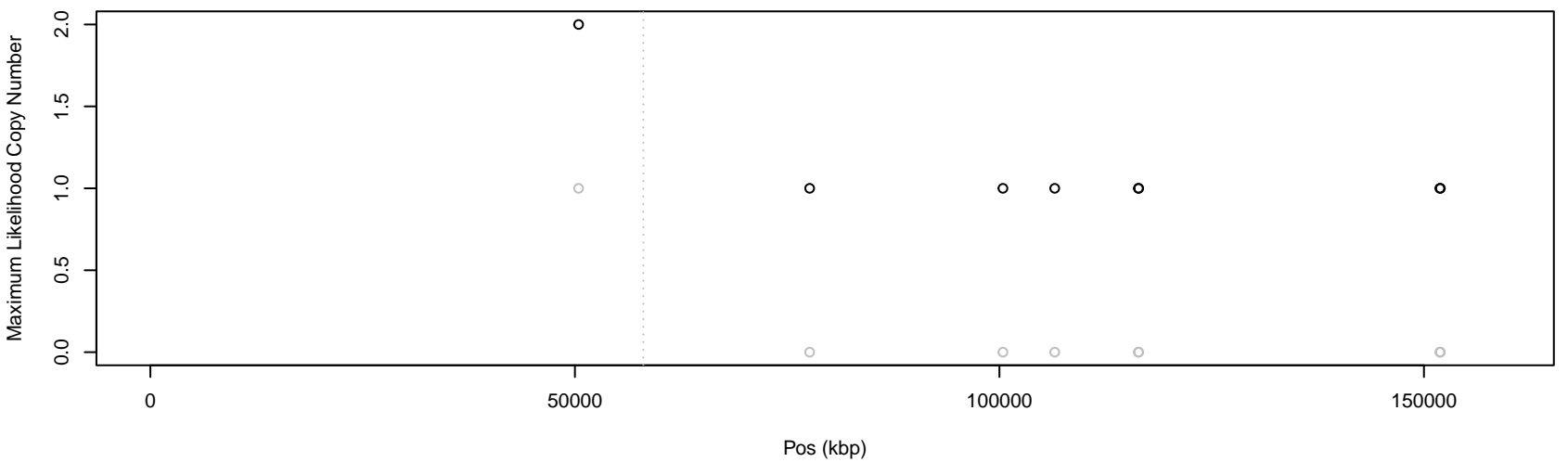
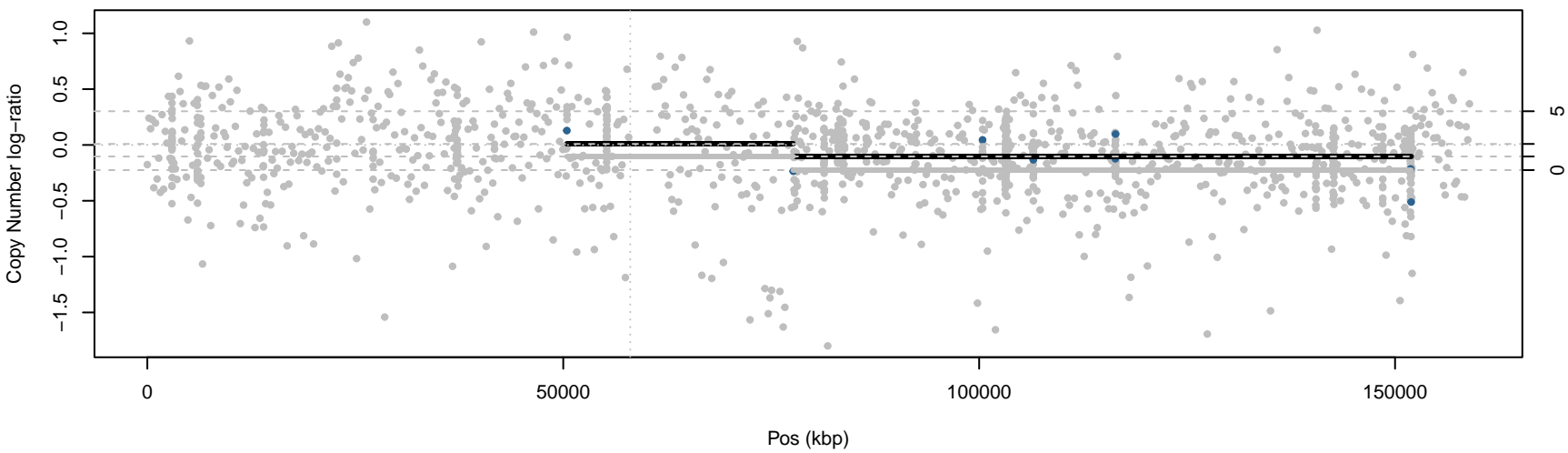
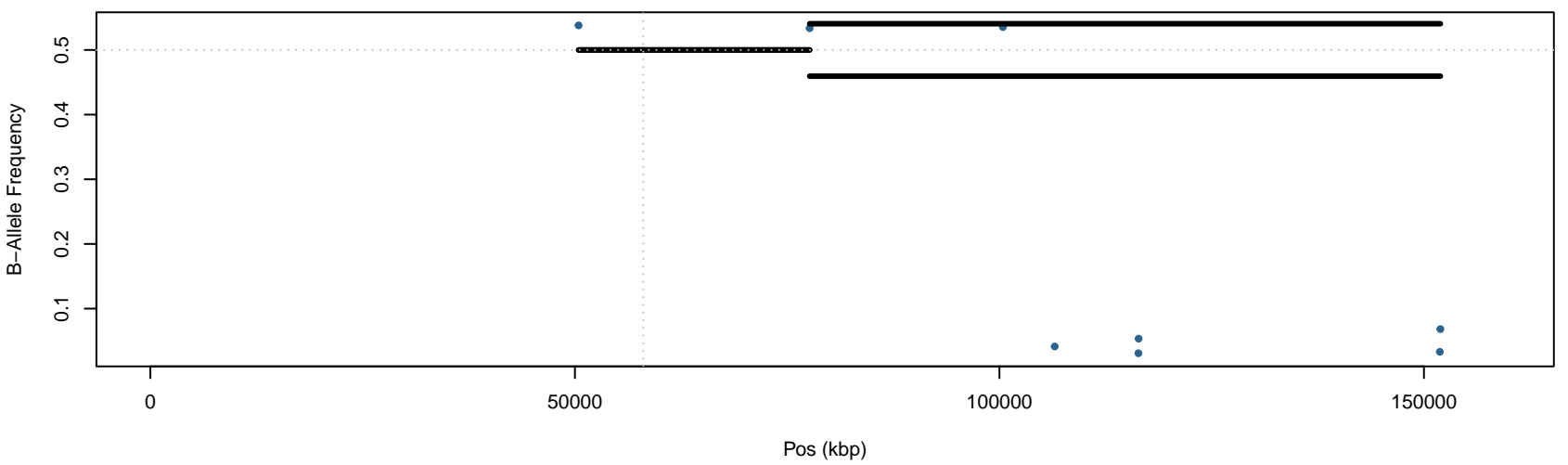
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 5



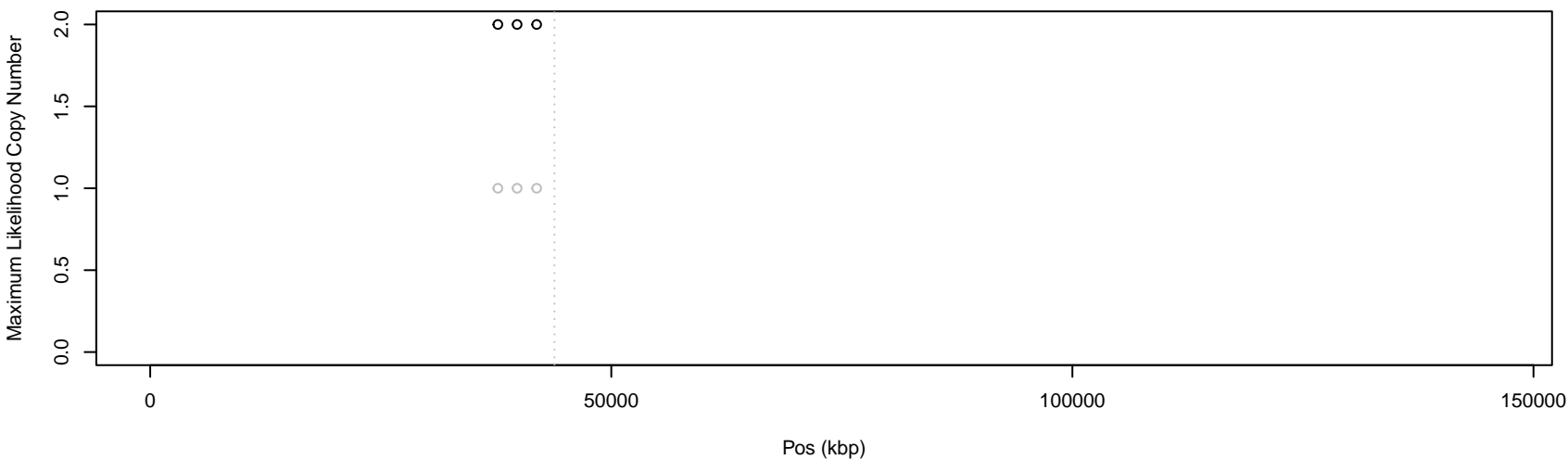
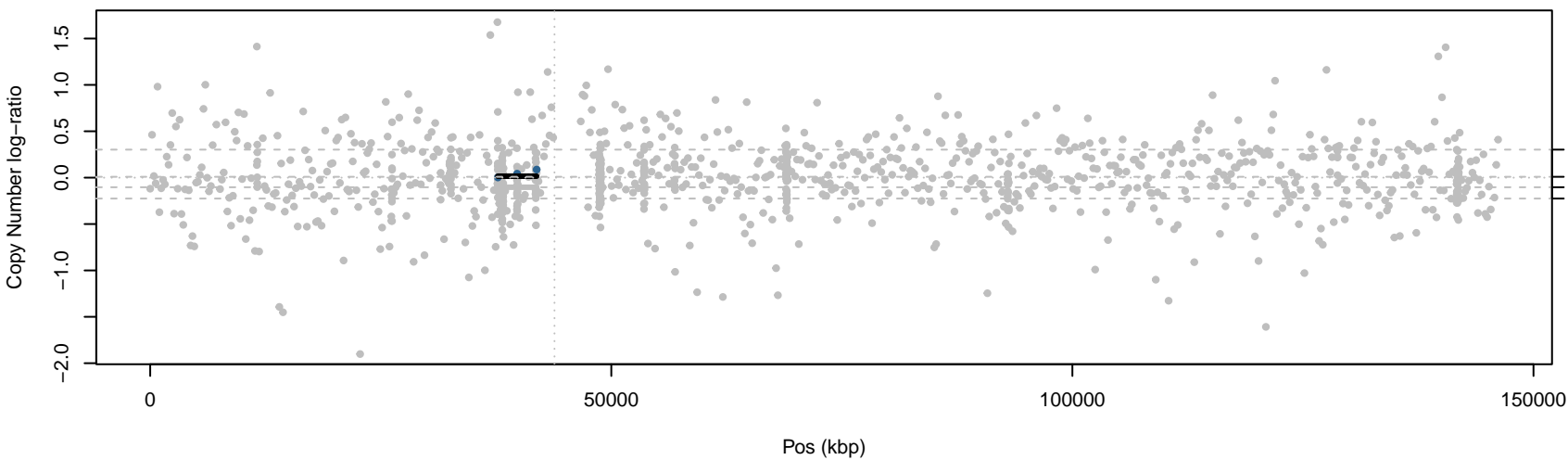
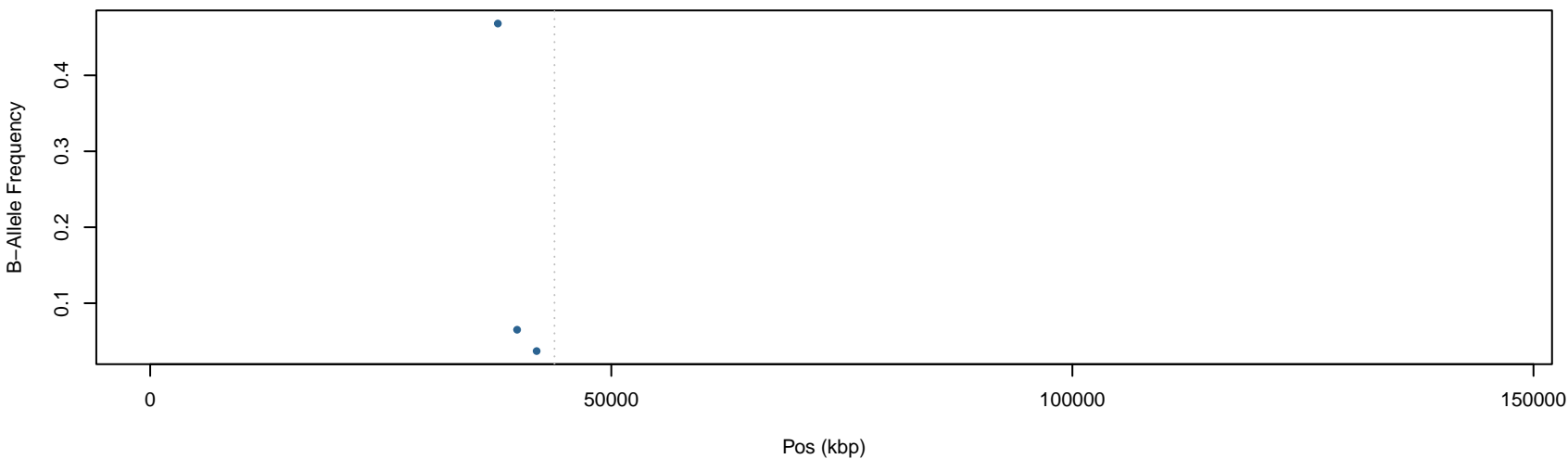
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 6



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 7

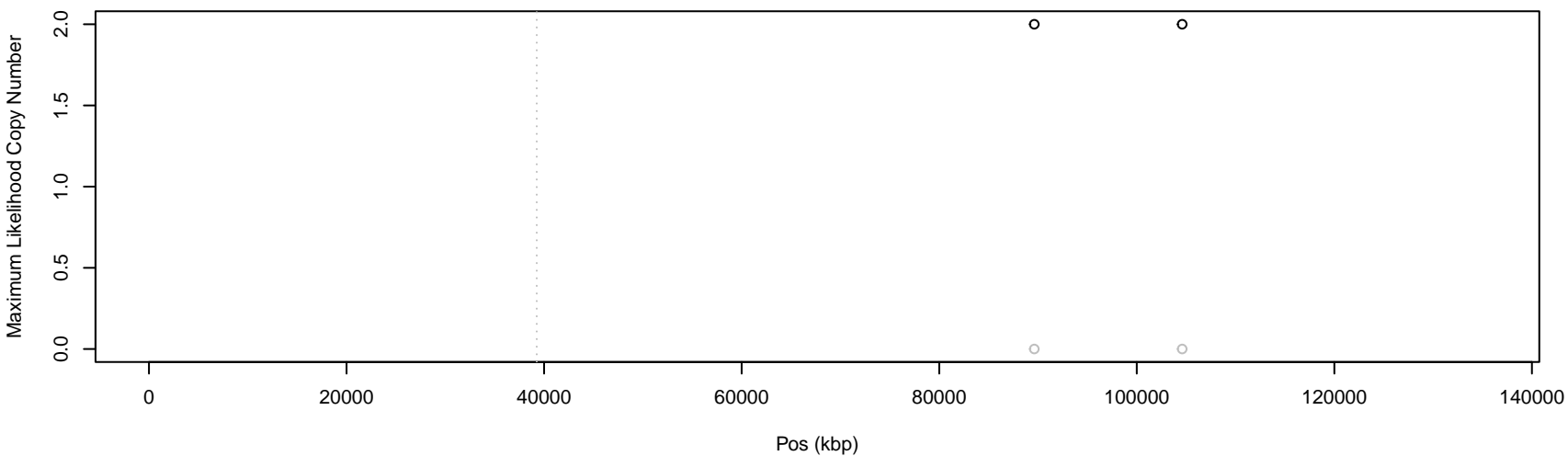
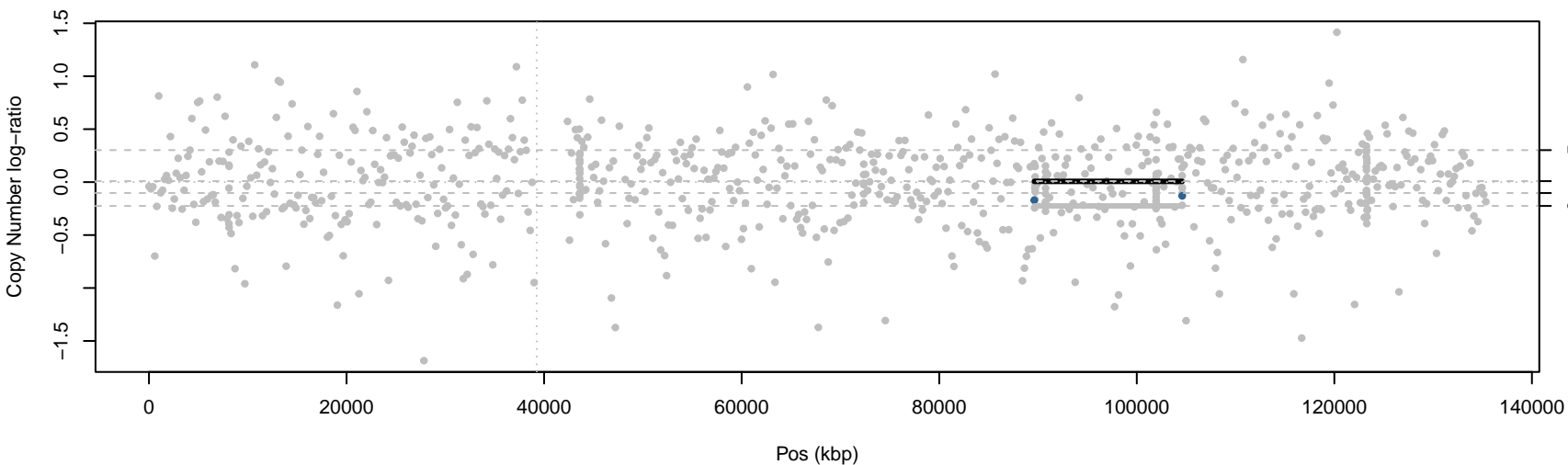
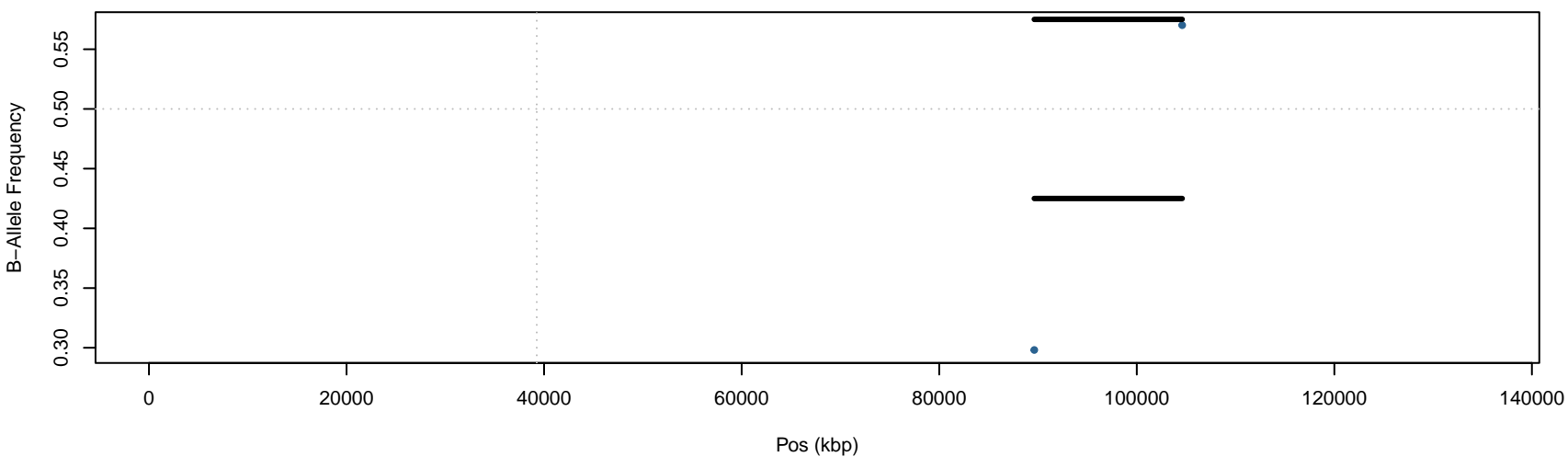


Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 8

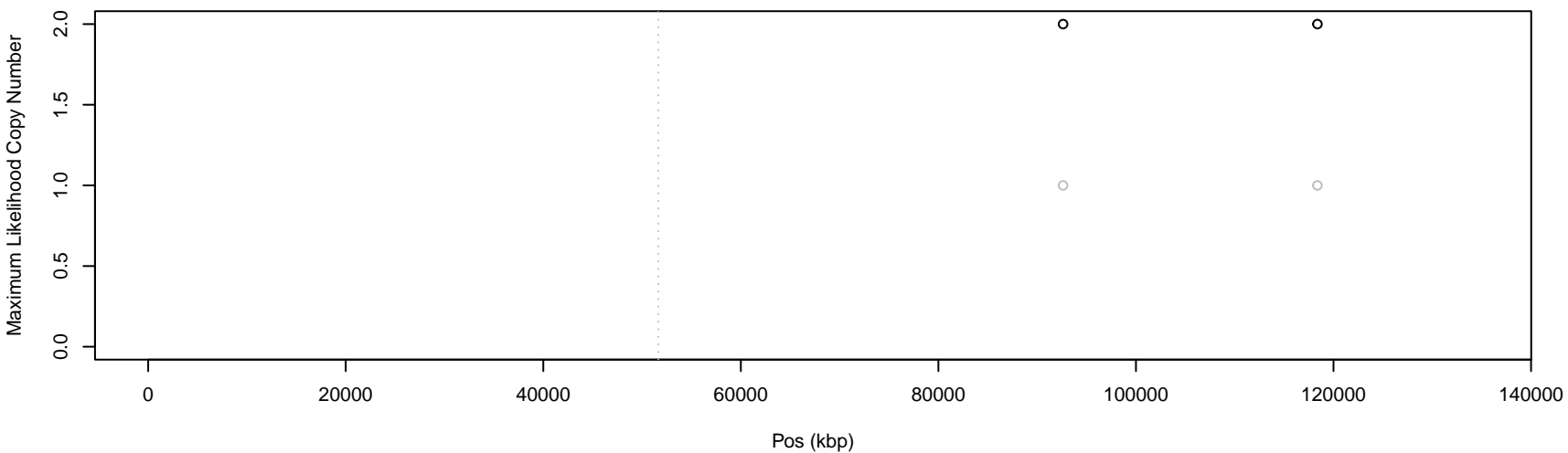
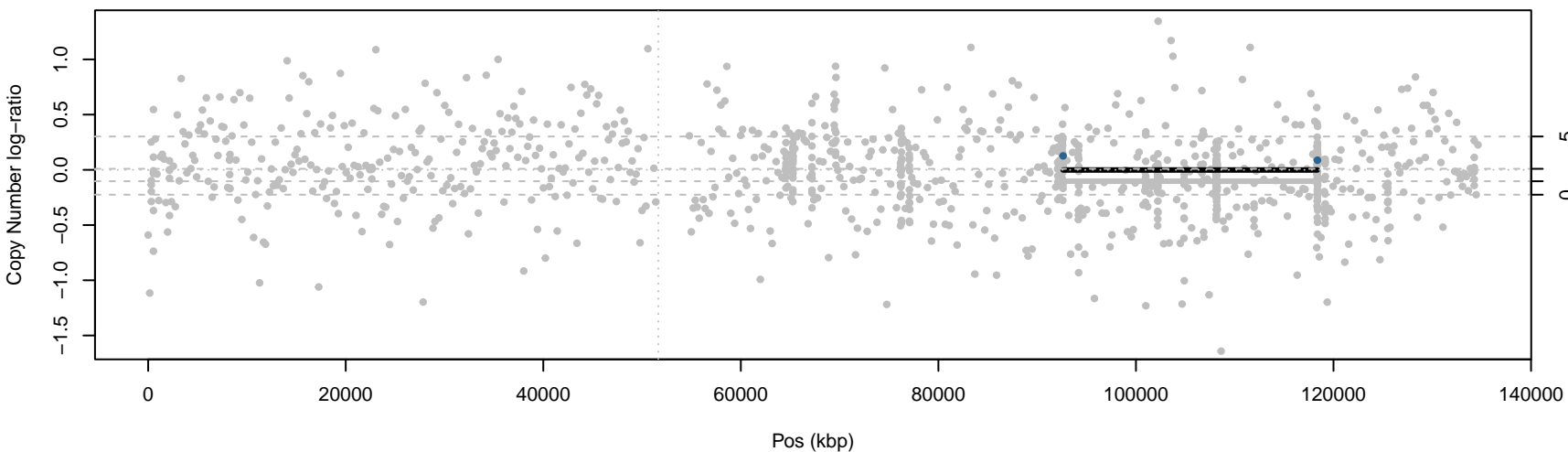
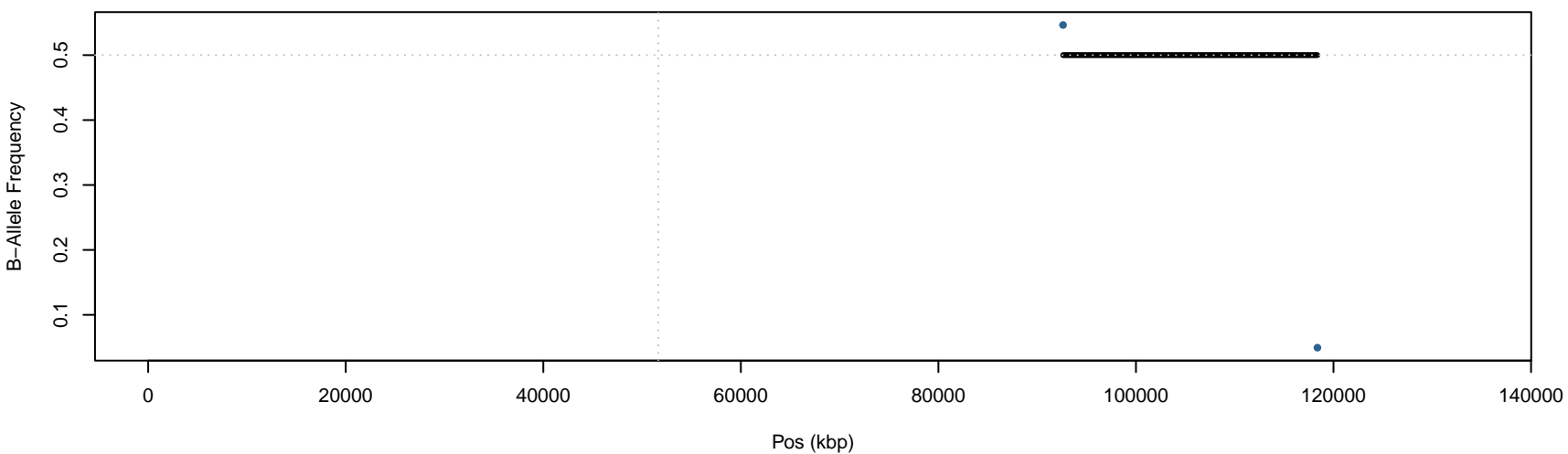




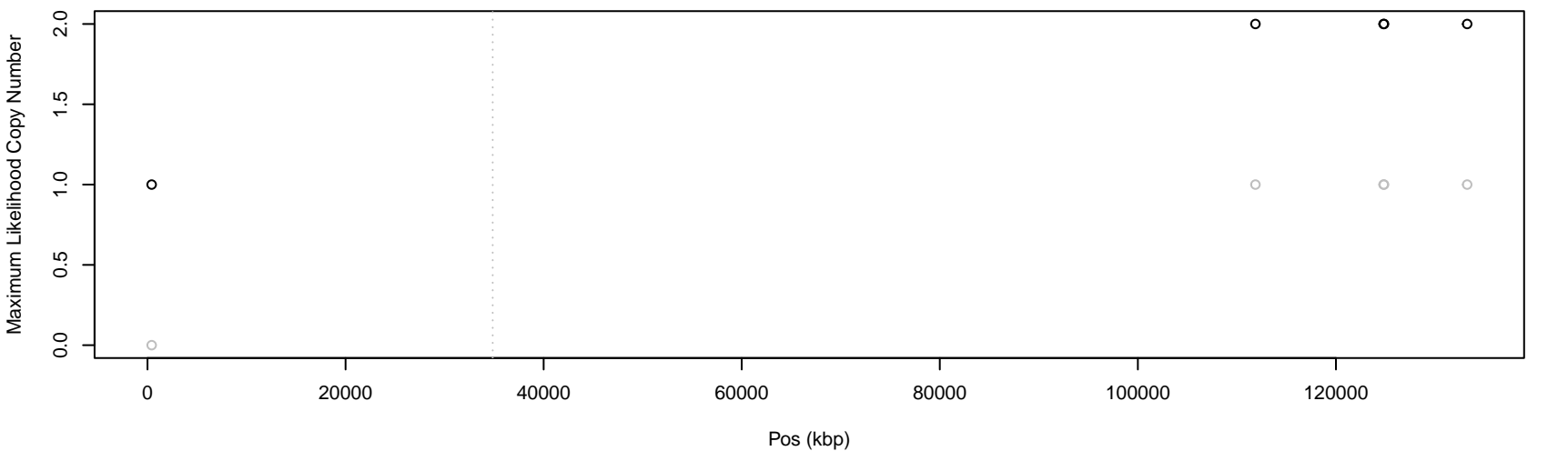
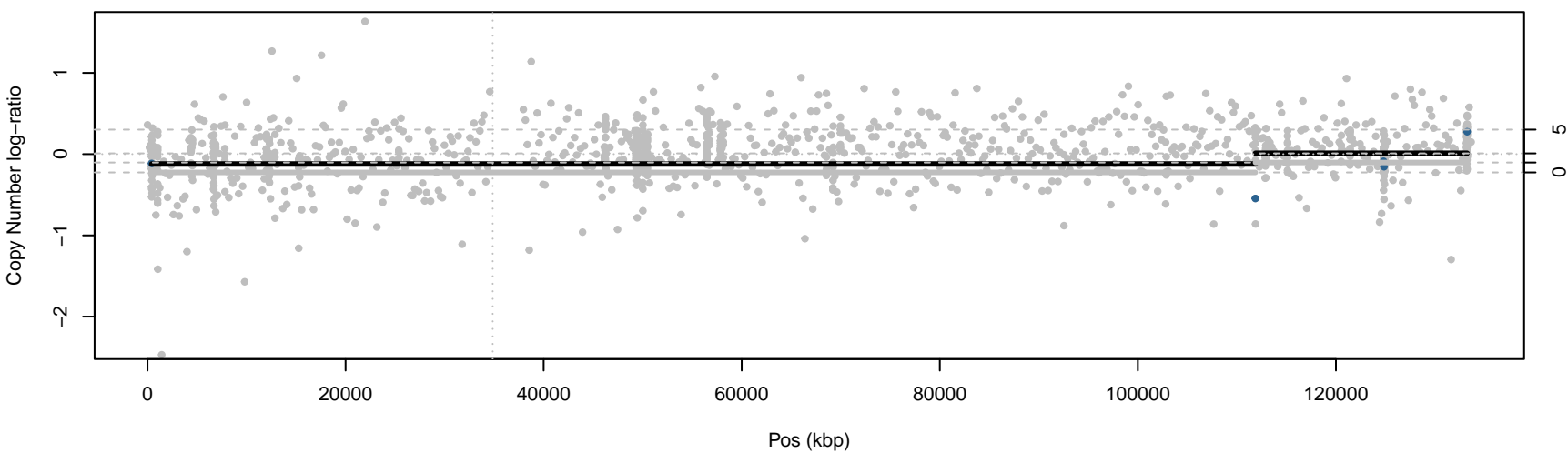
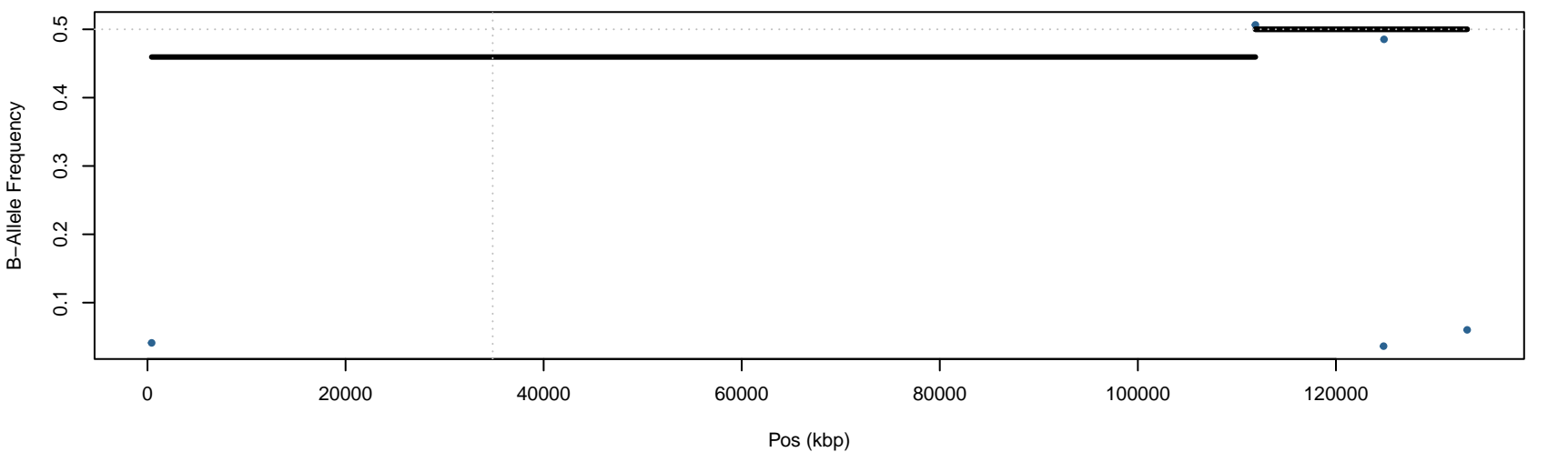
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 10



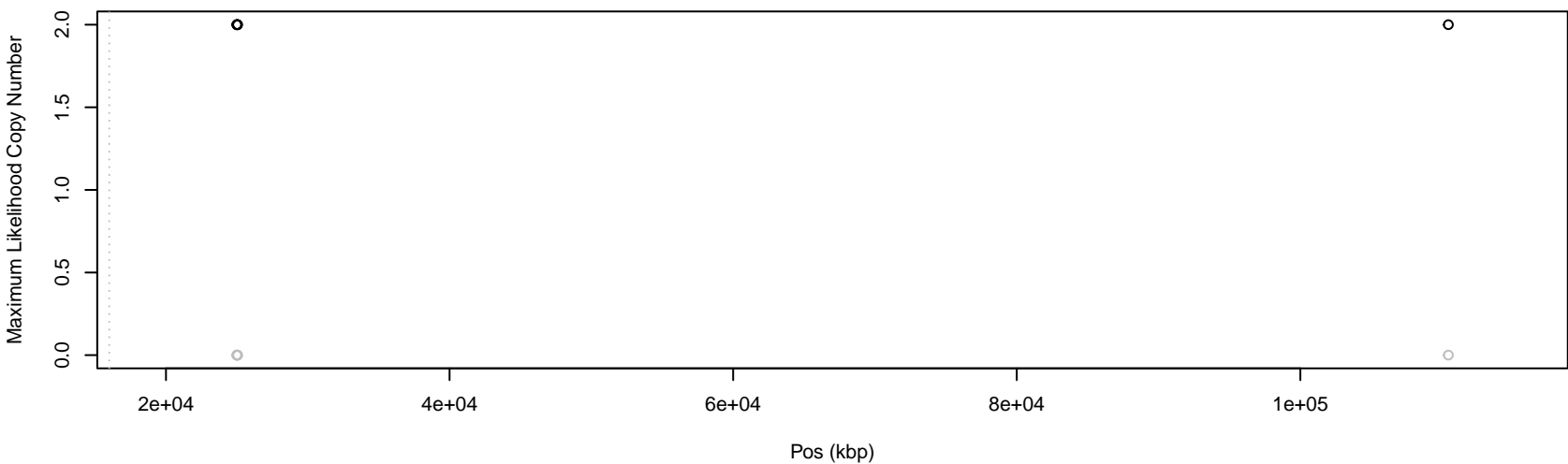
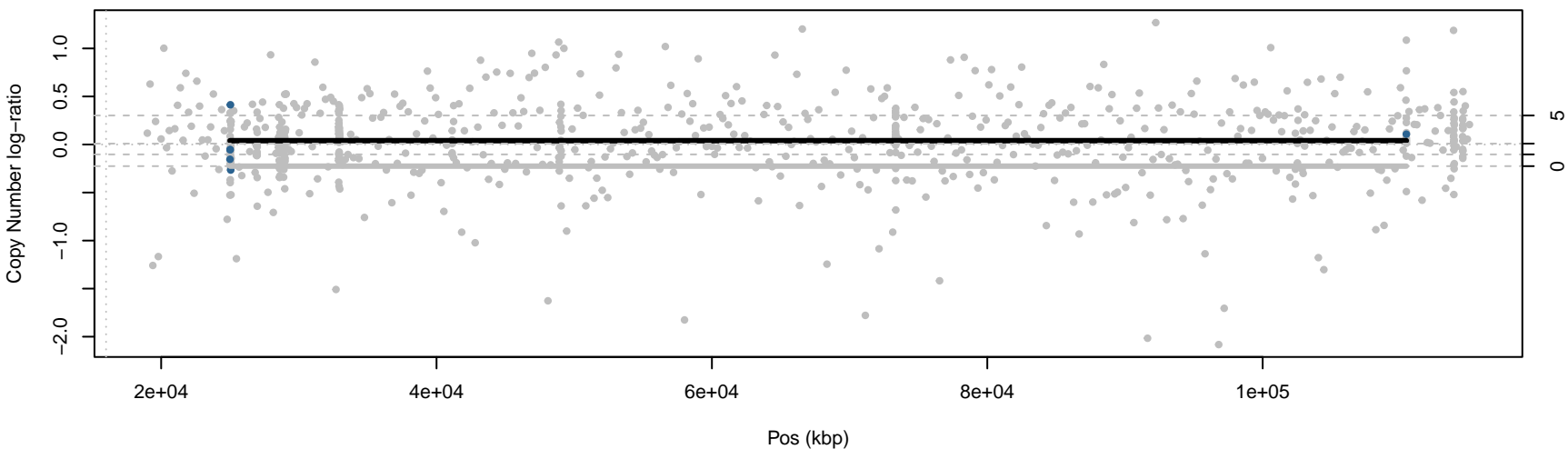
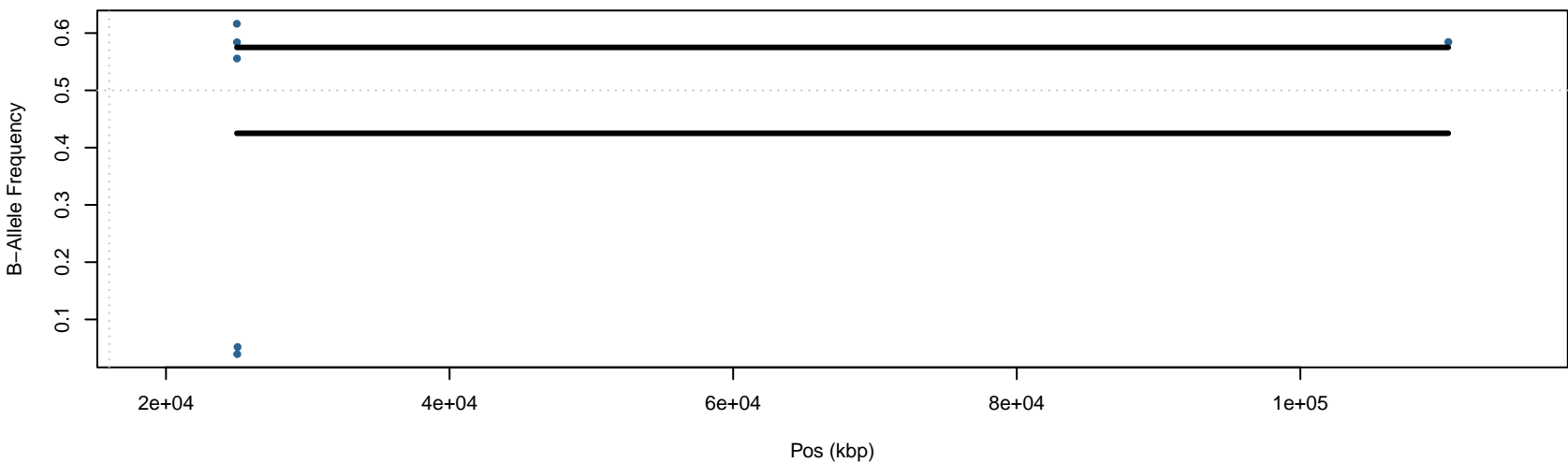
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 11



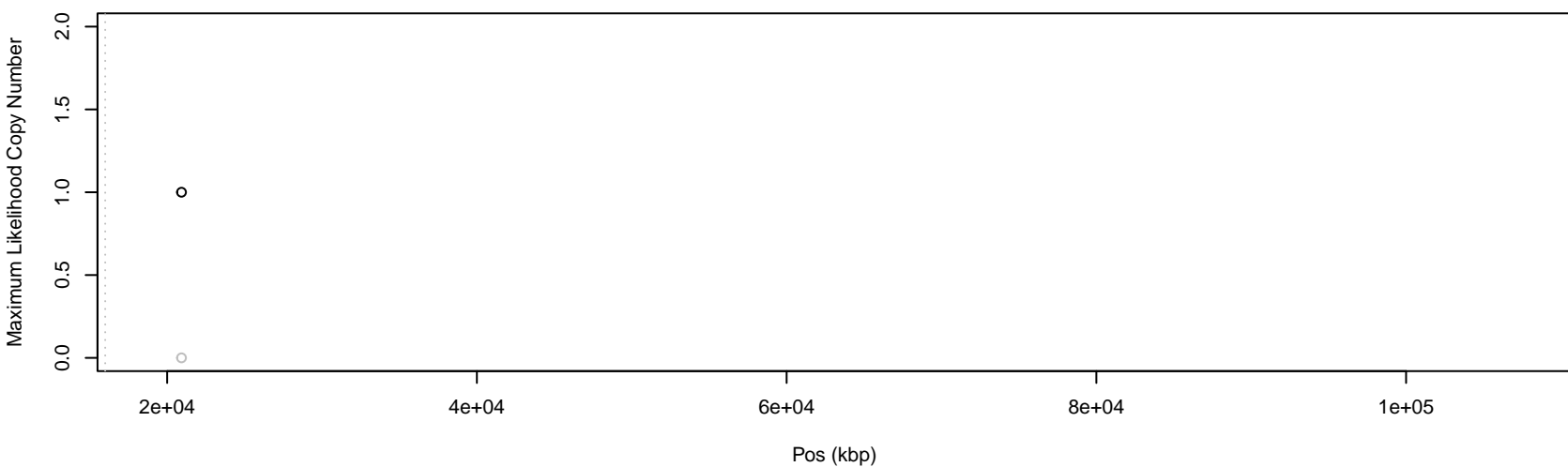
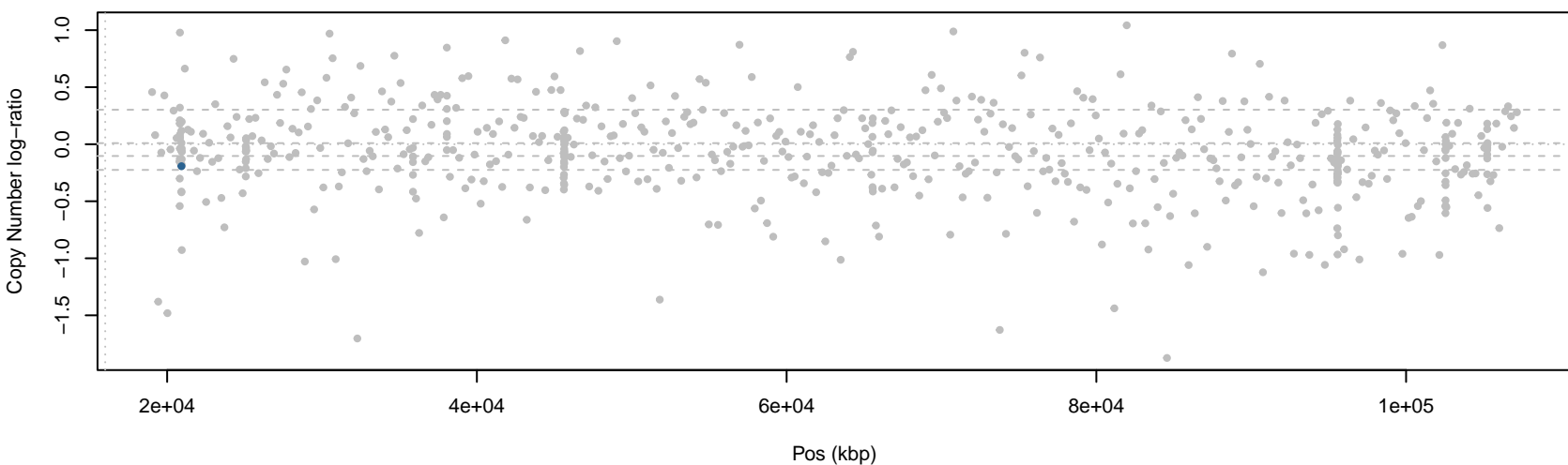
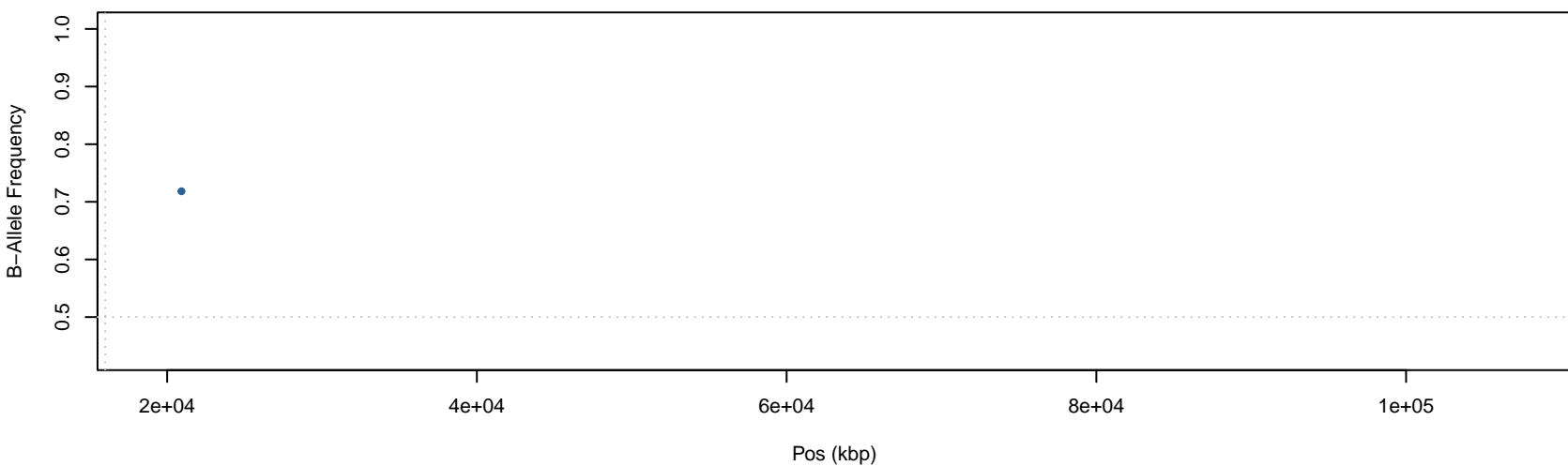
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 12



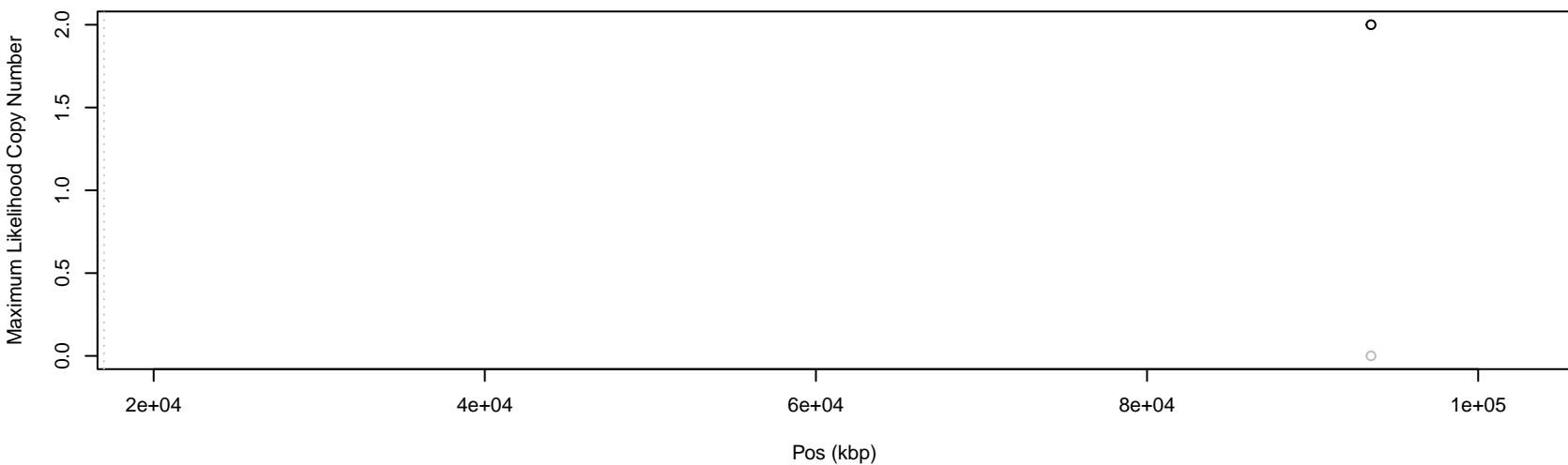
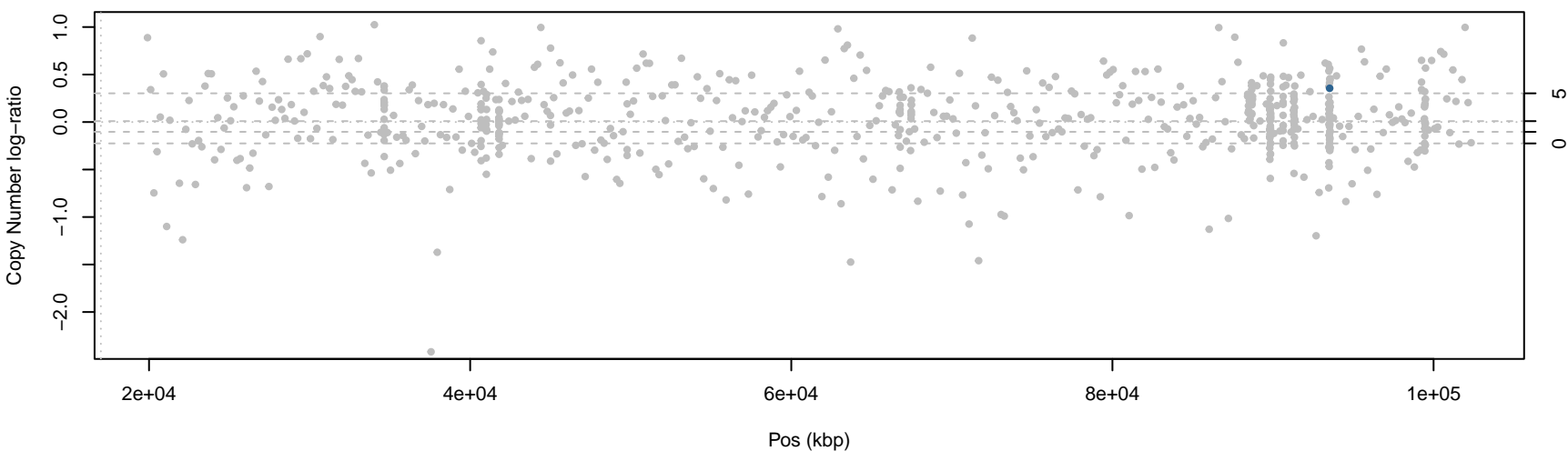
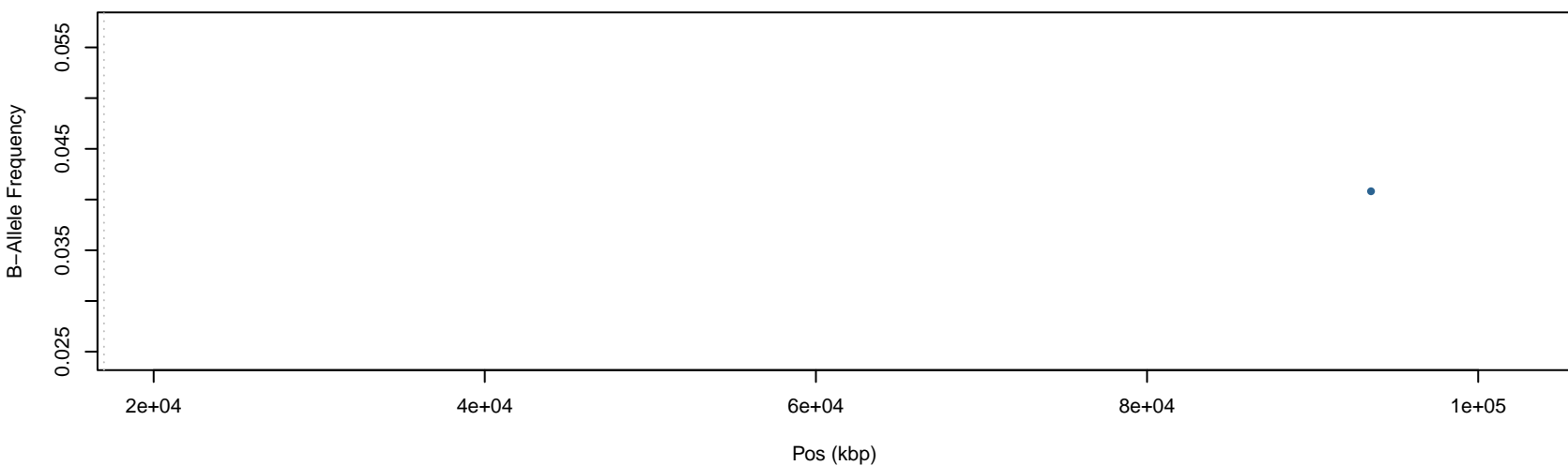
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 13



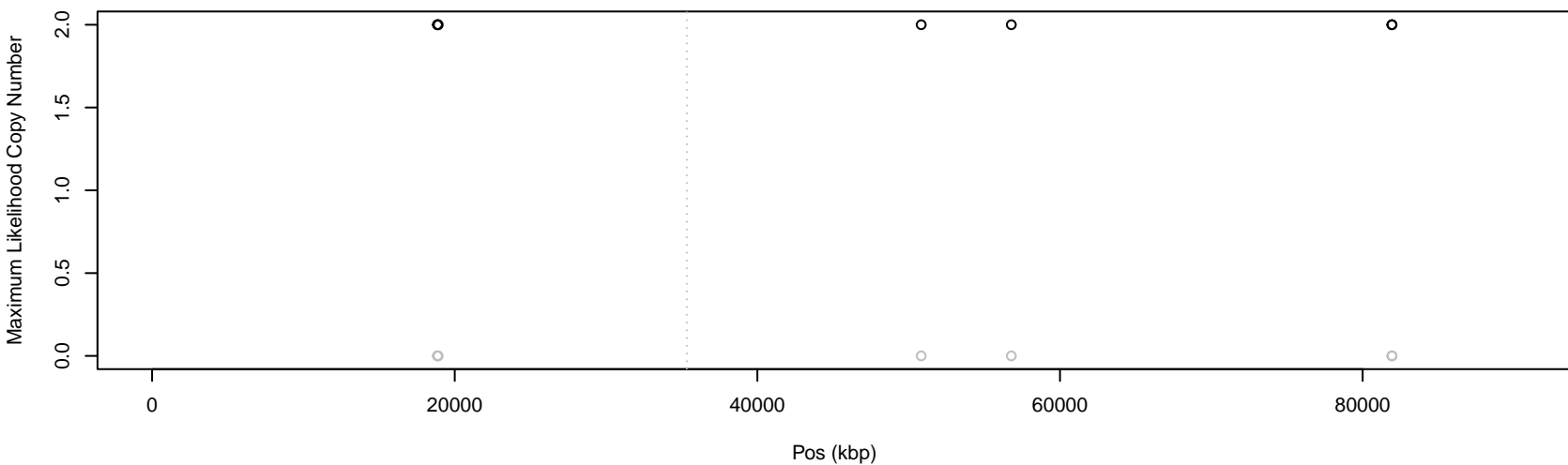
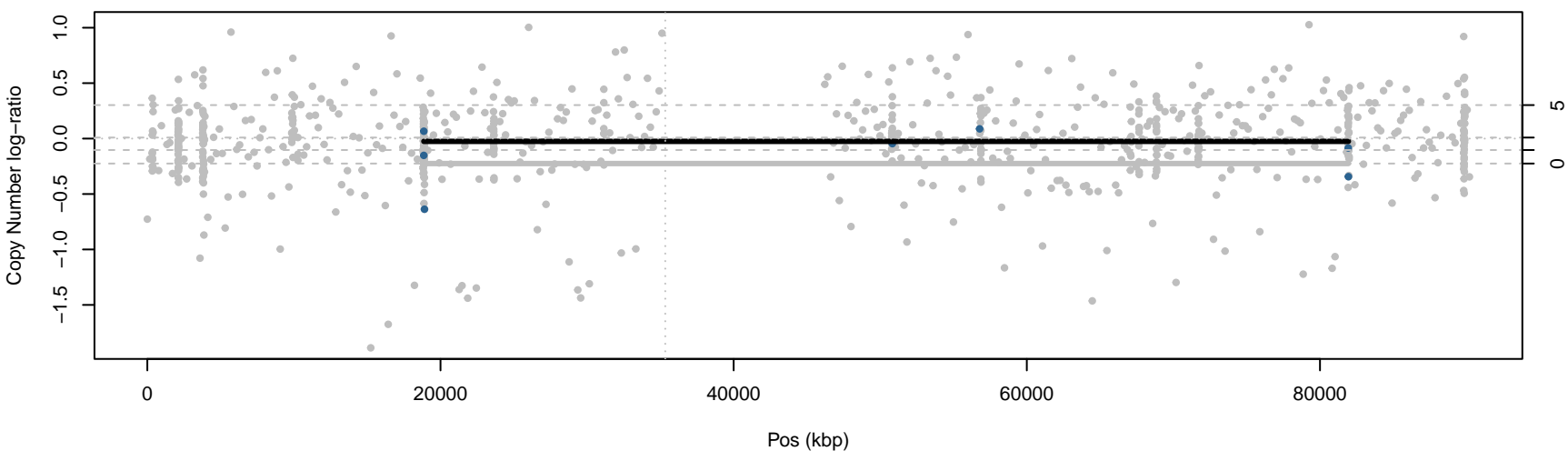
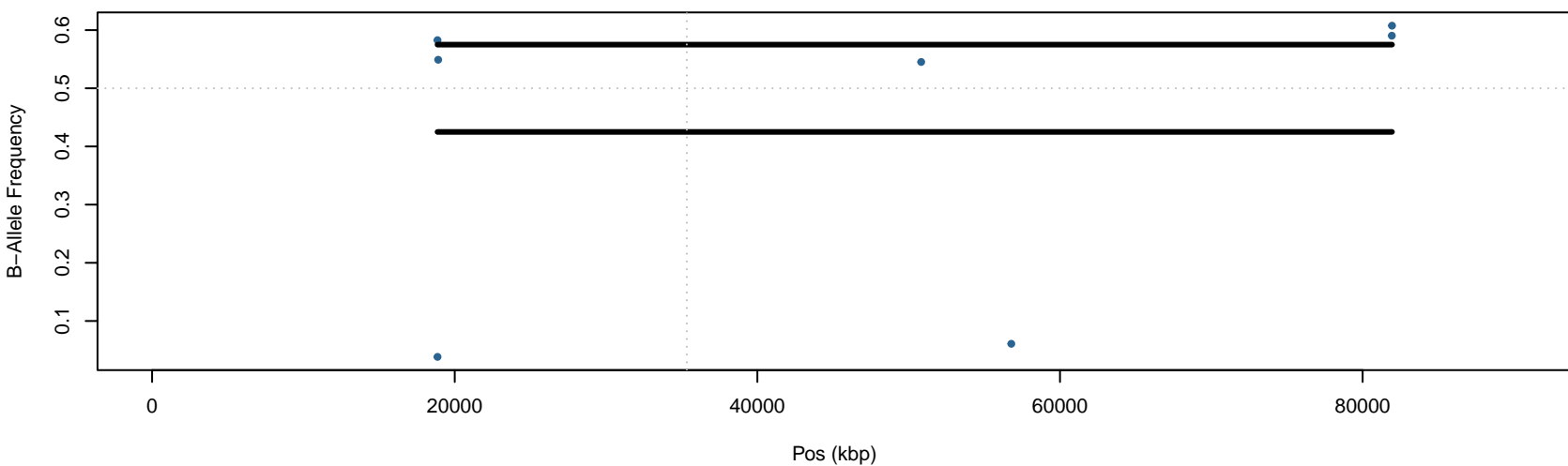
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 14



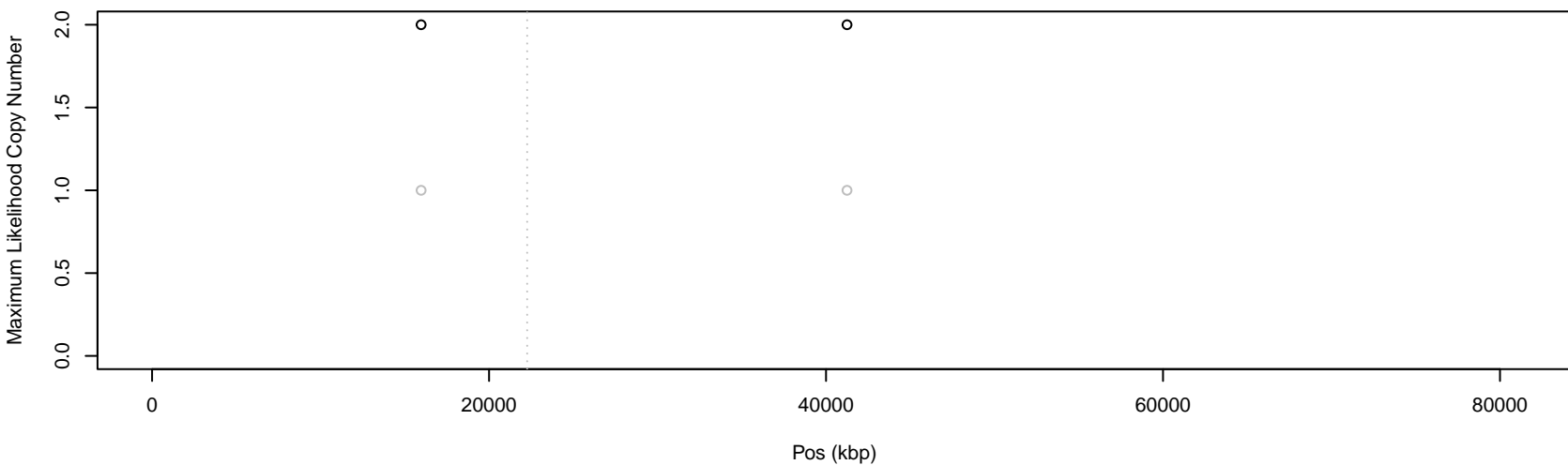
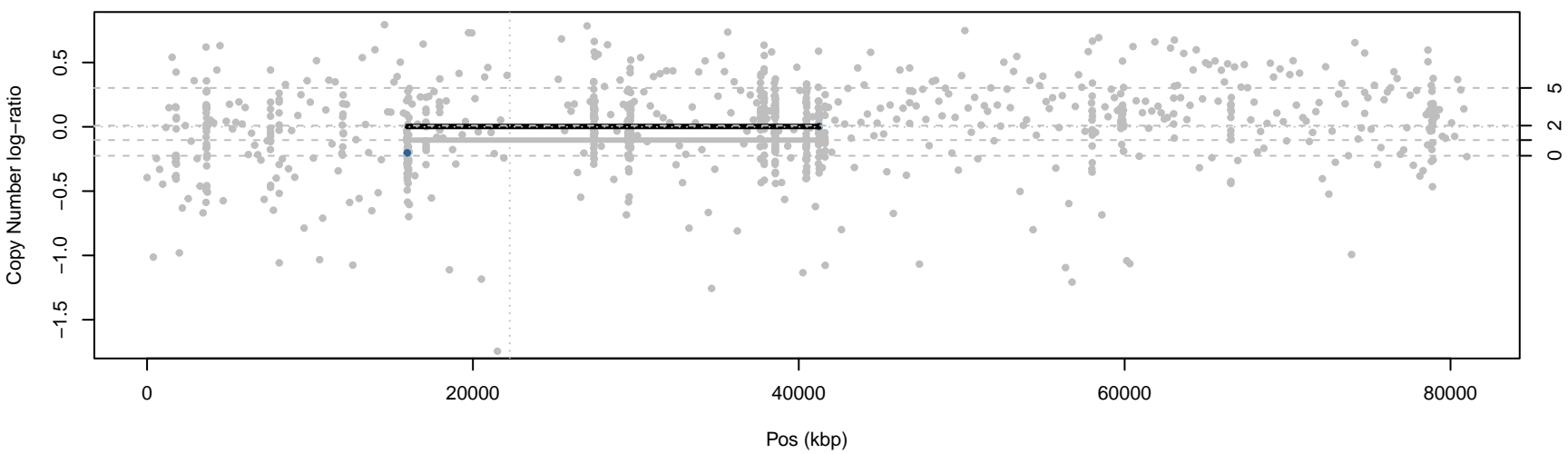
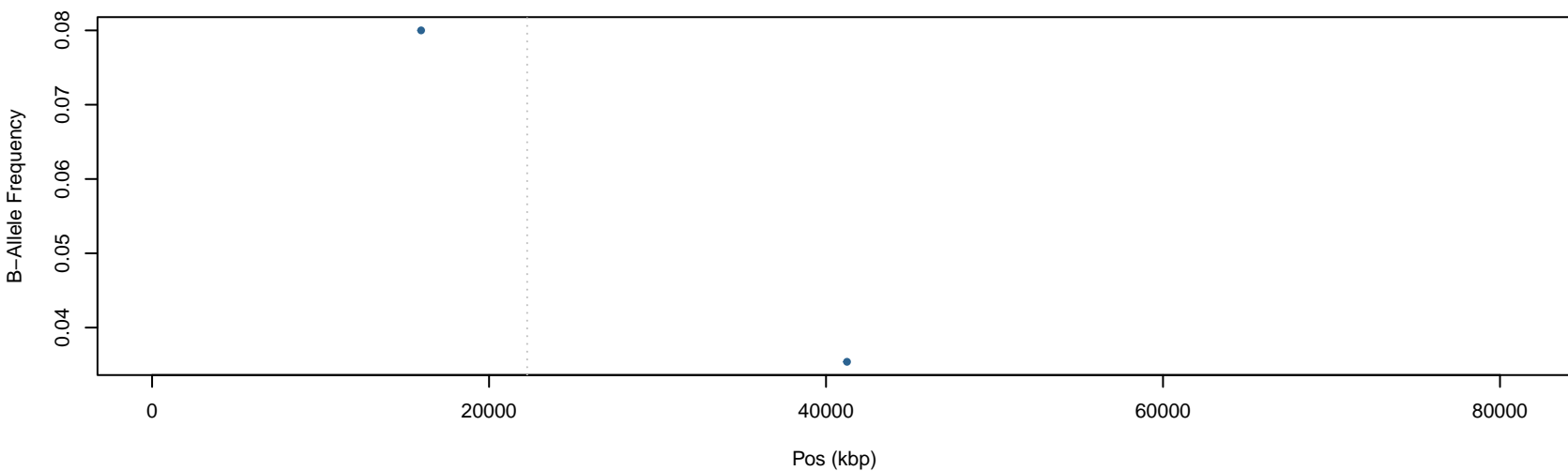
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 15



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 16

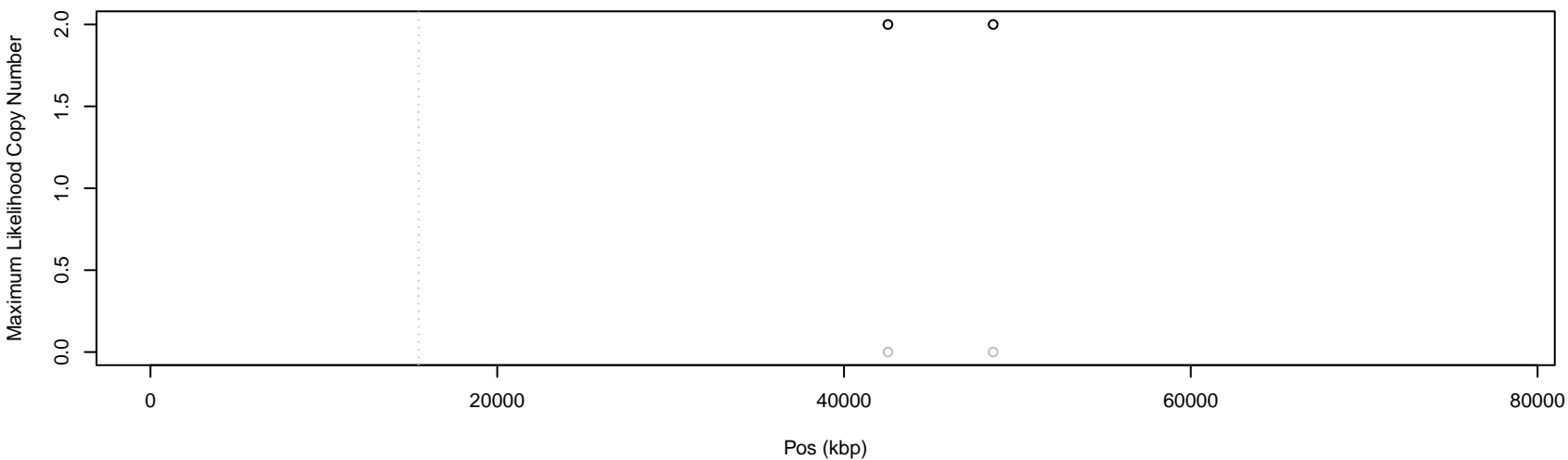
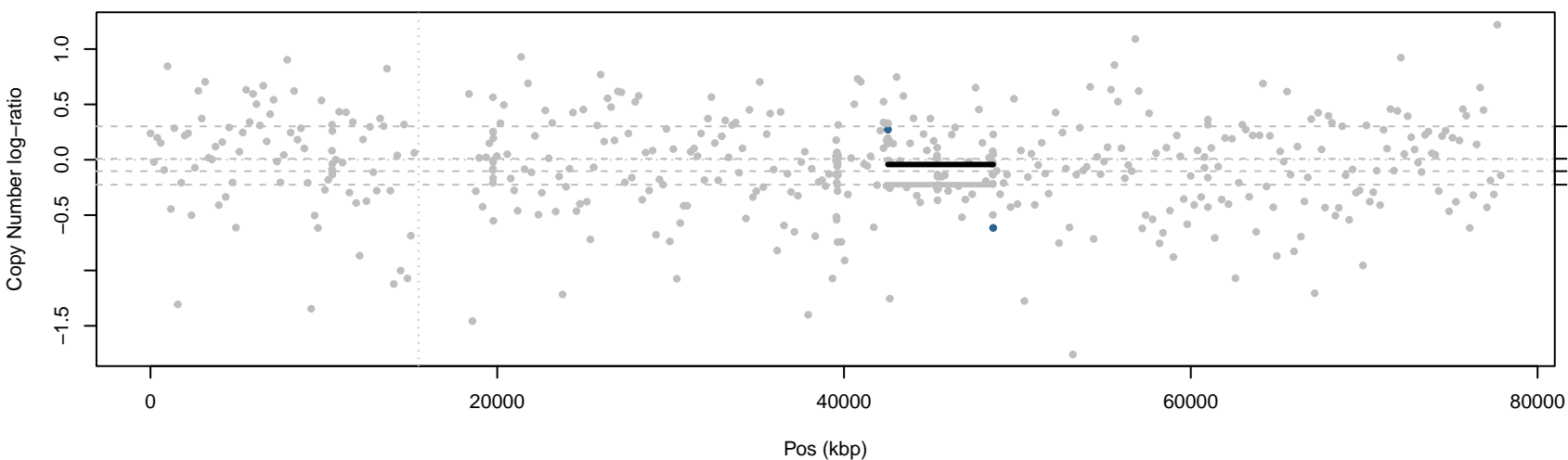
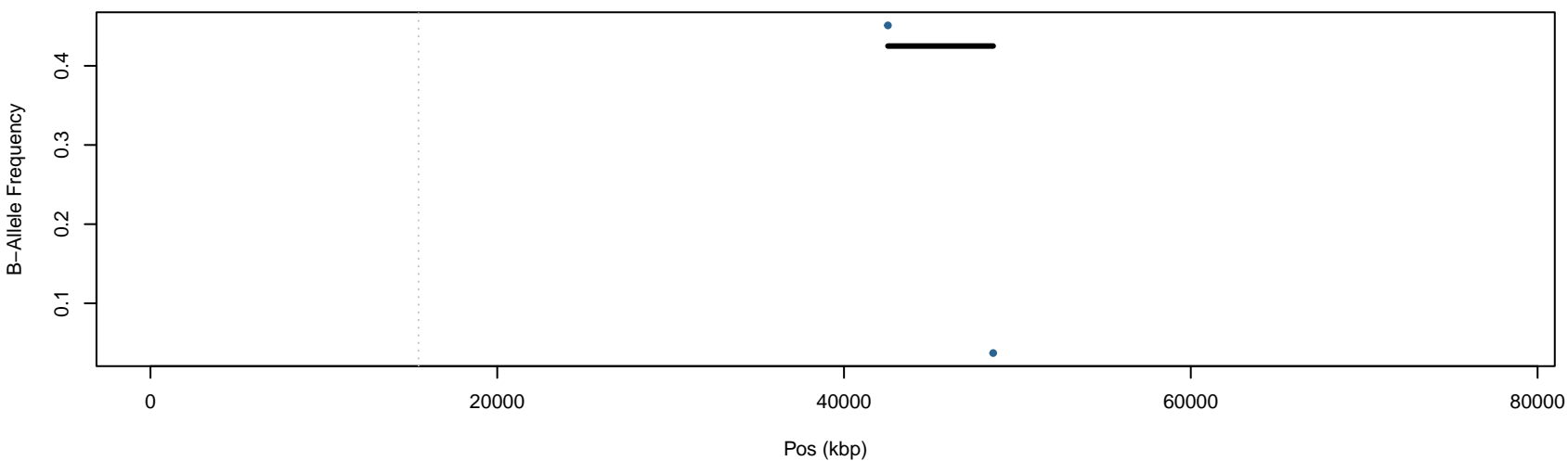


Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 17

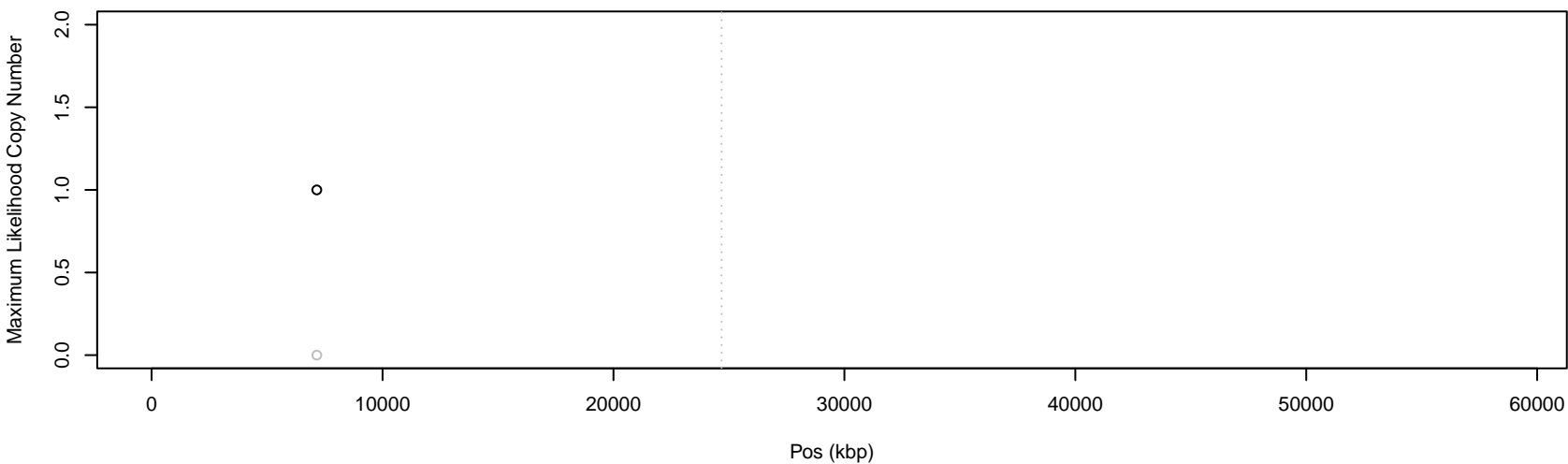
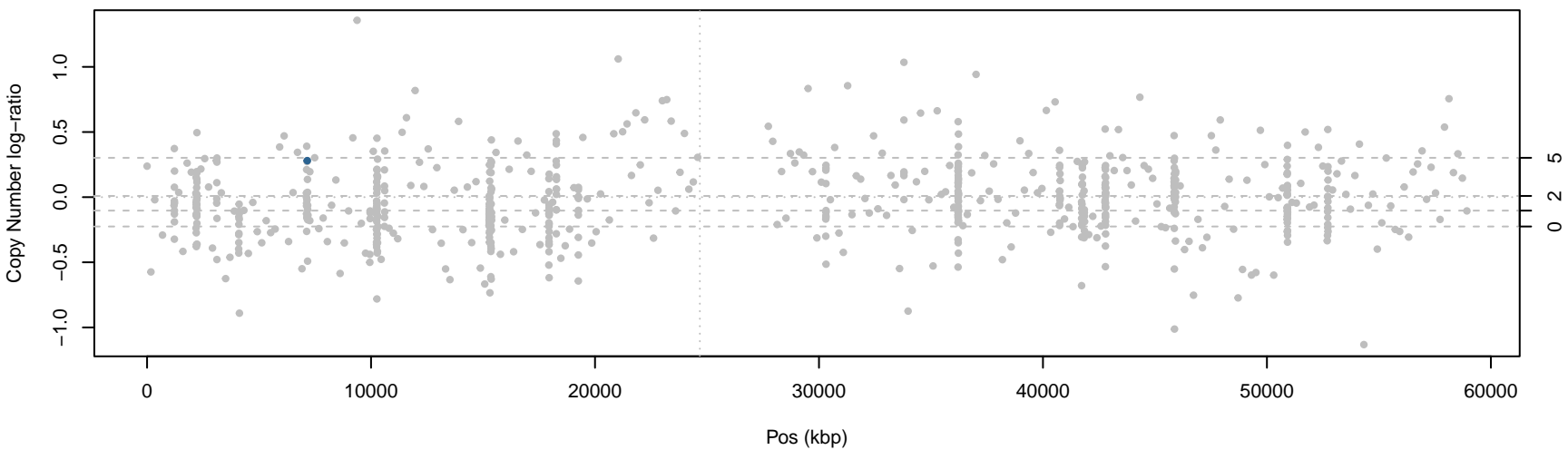
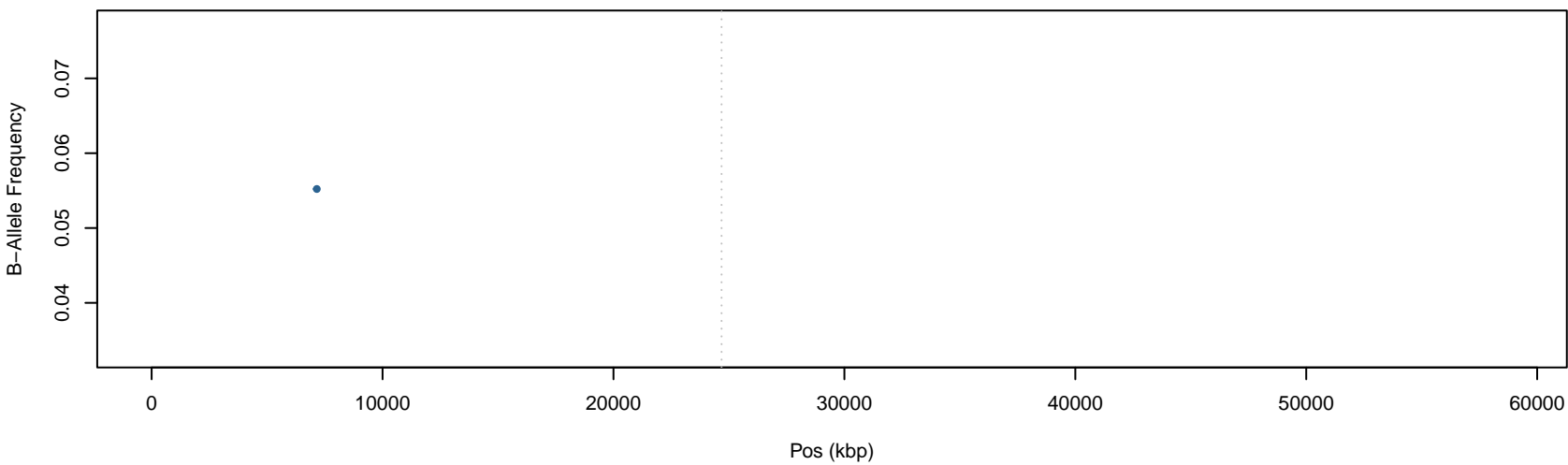




Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 18



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 19



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 22

