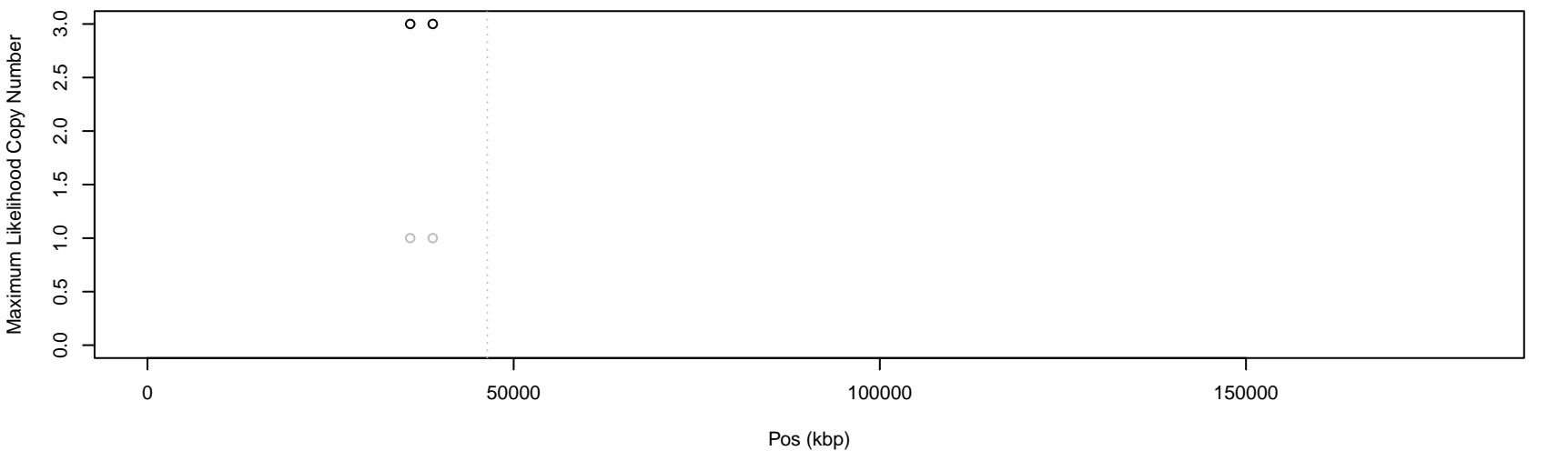
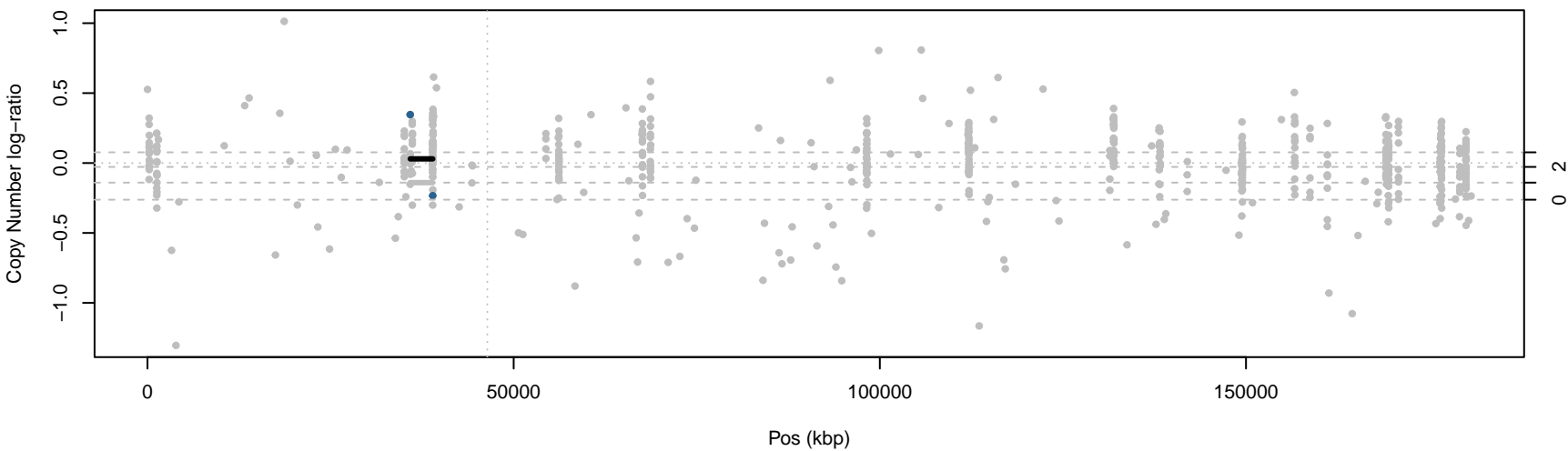
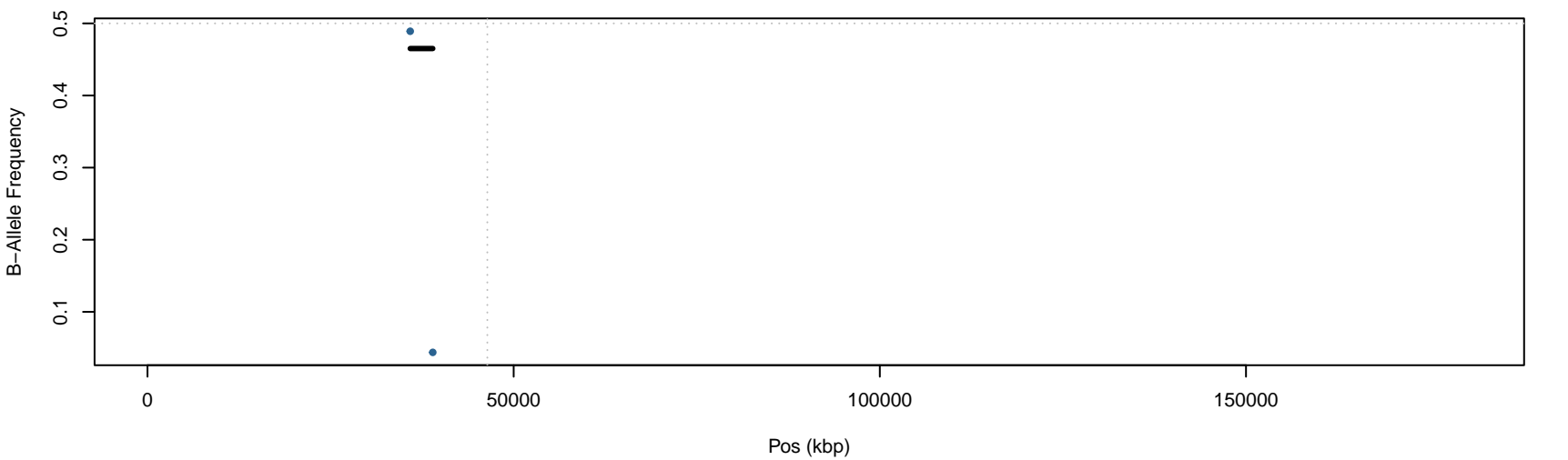
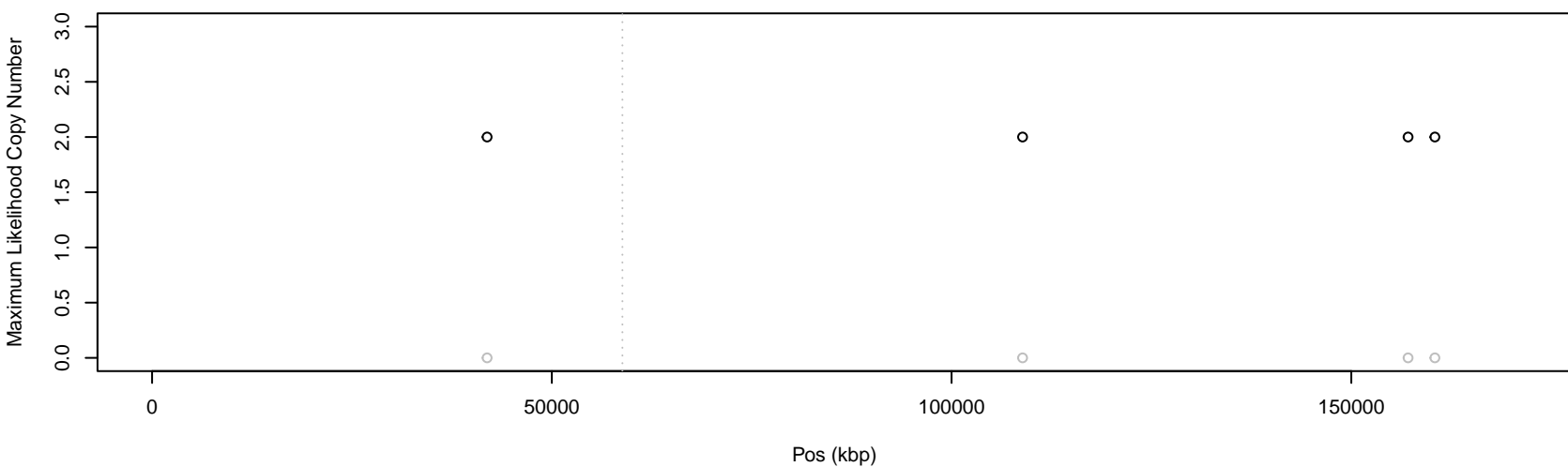
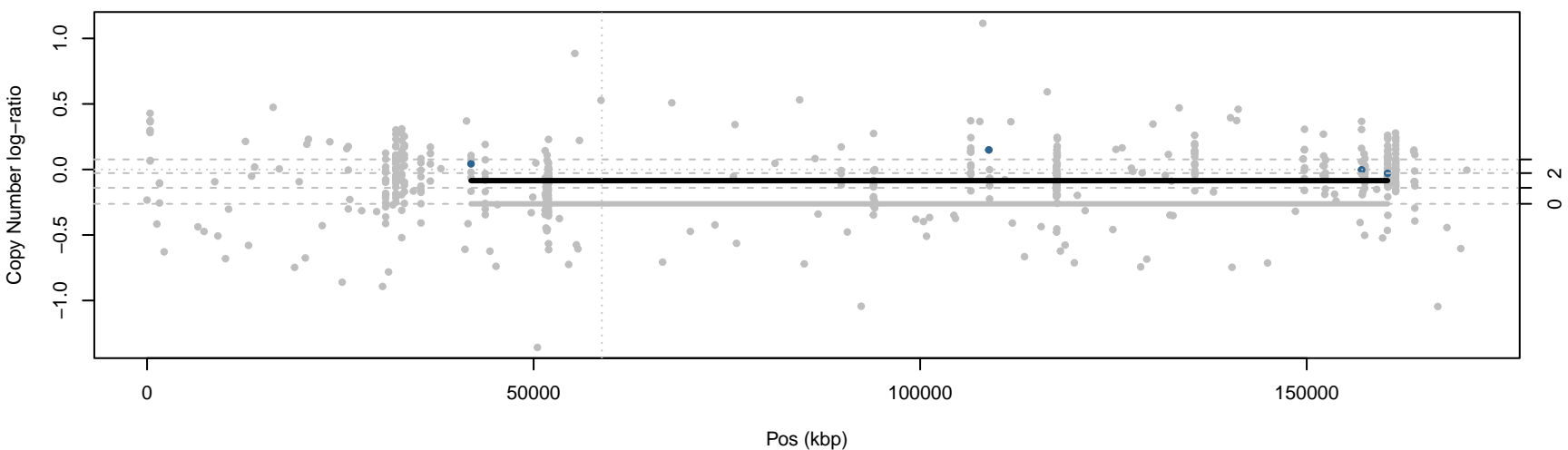
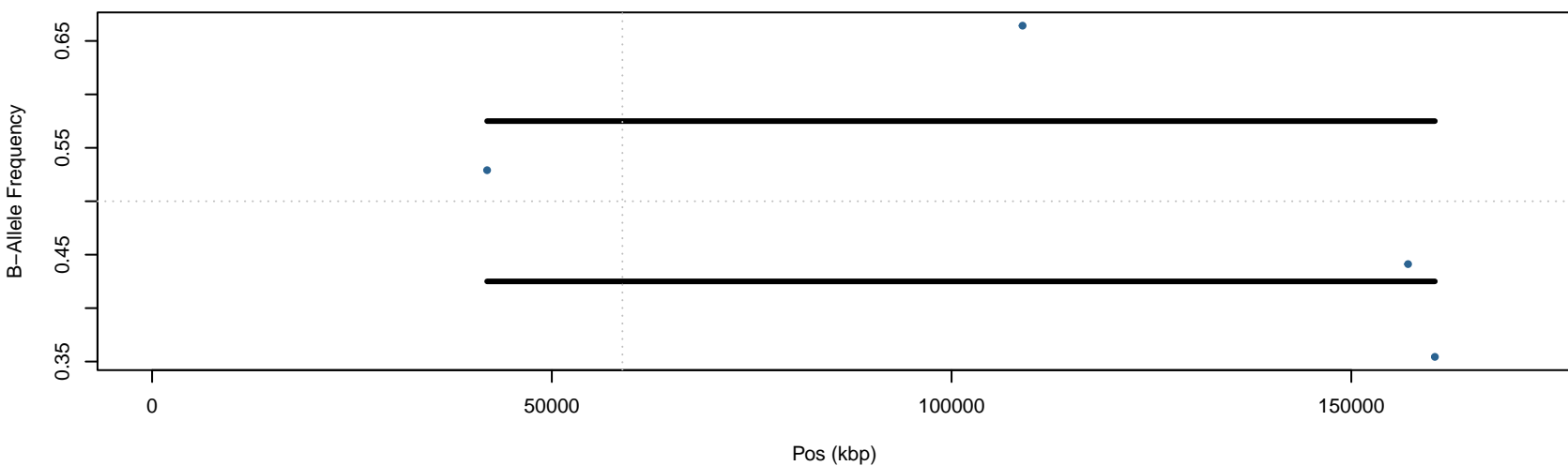


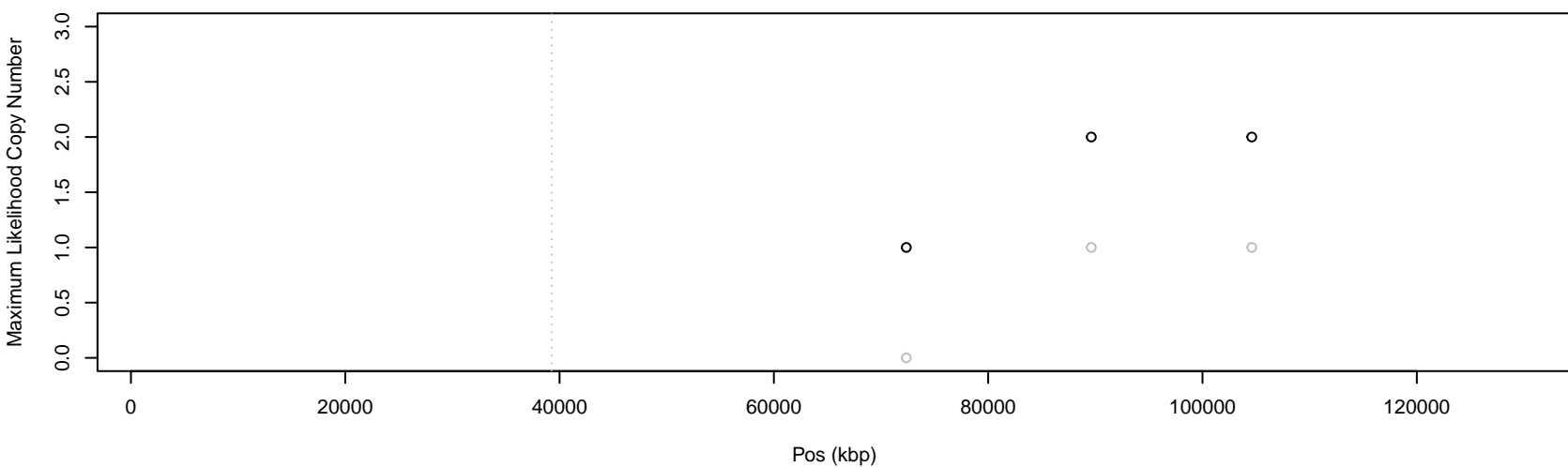
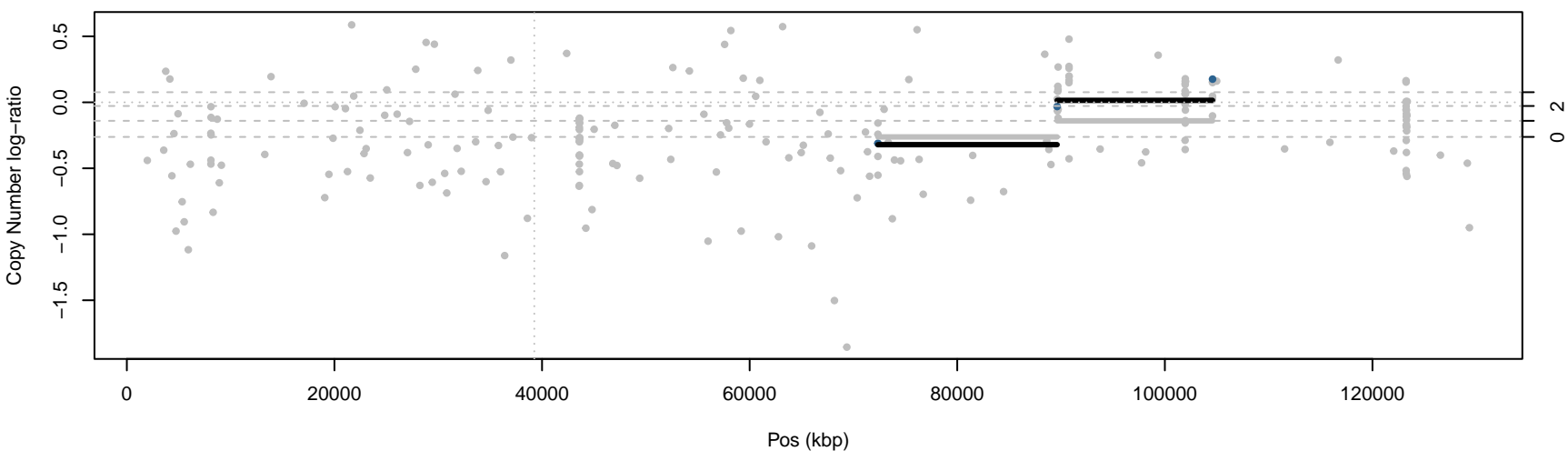
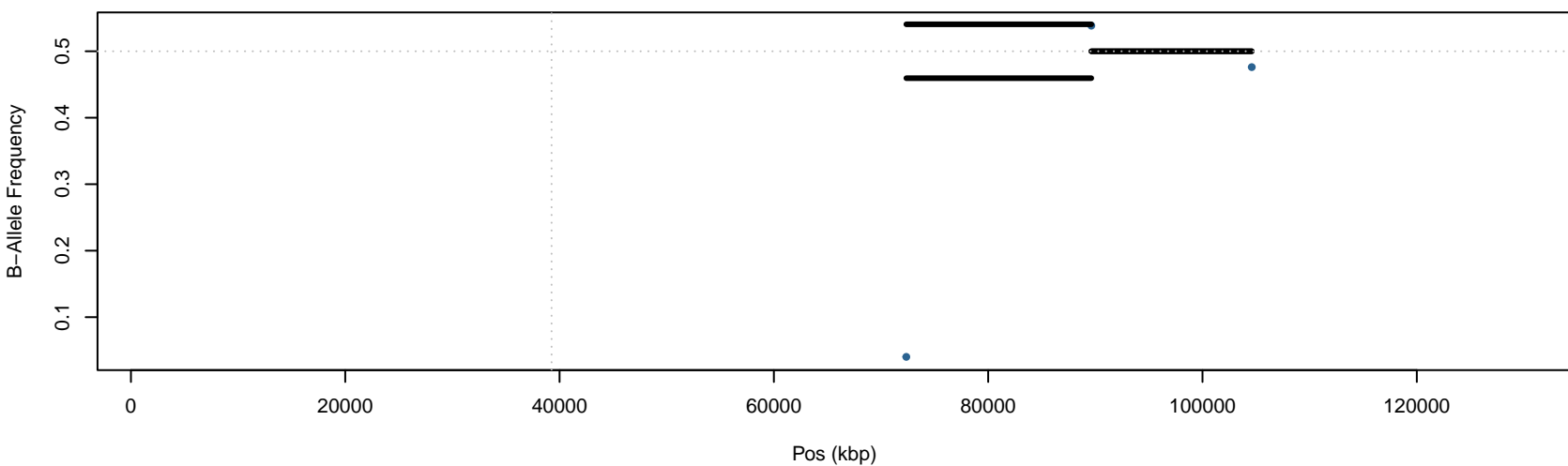
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 5



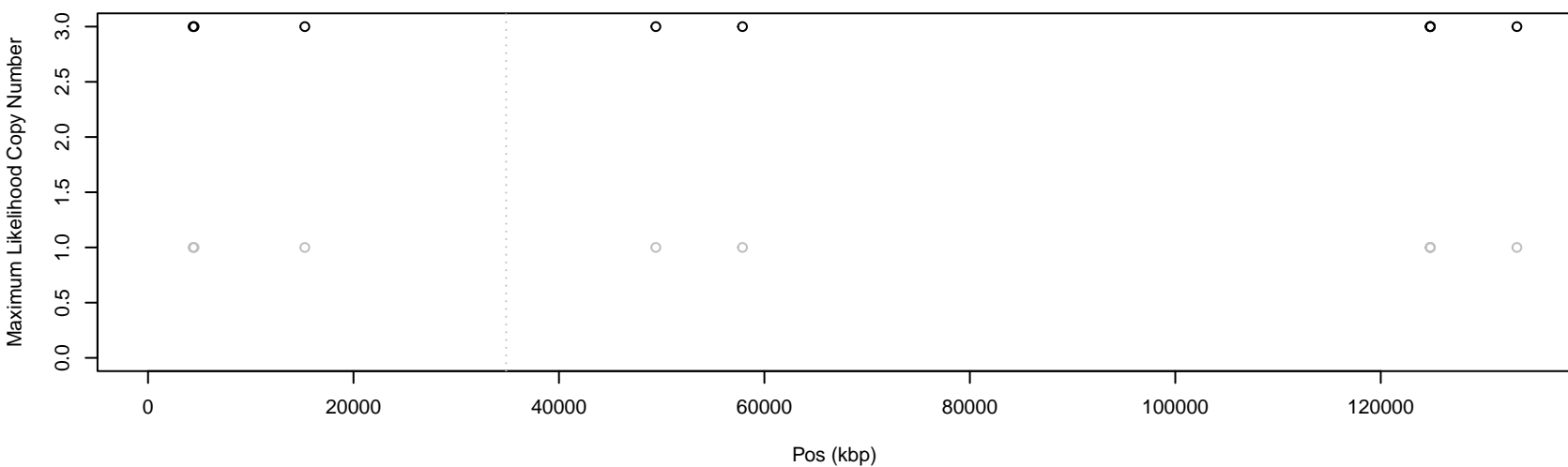
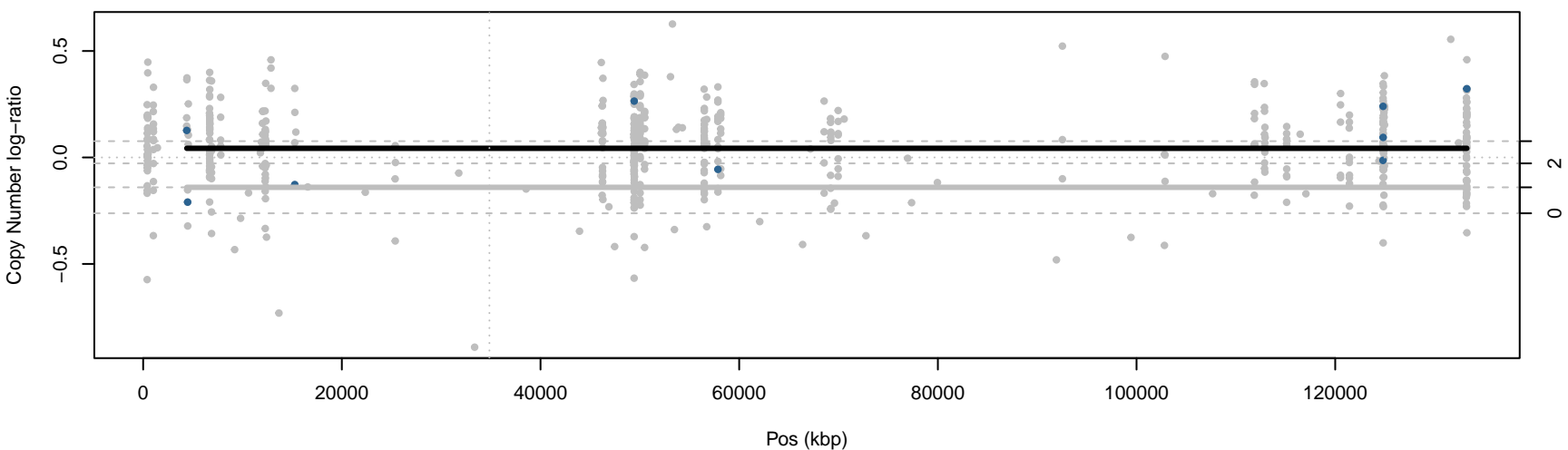
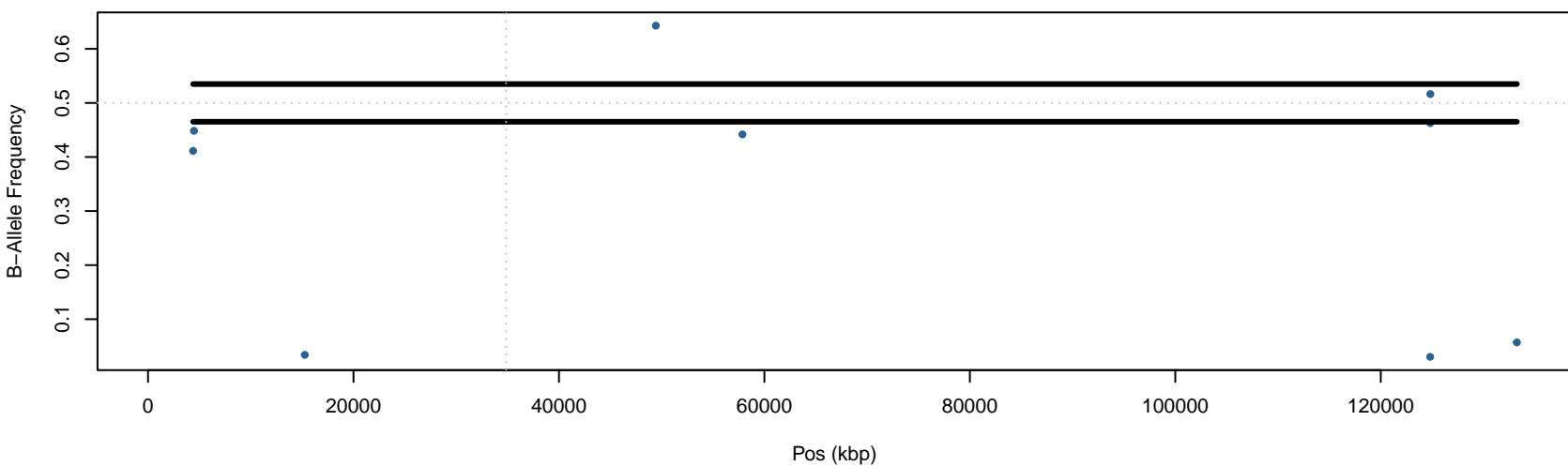
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 6



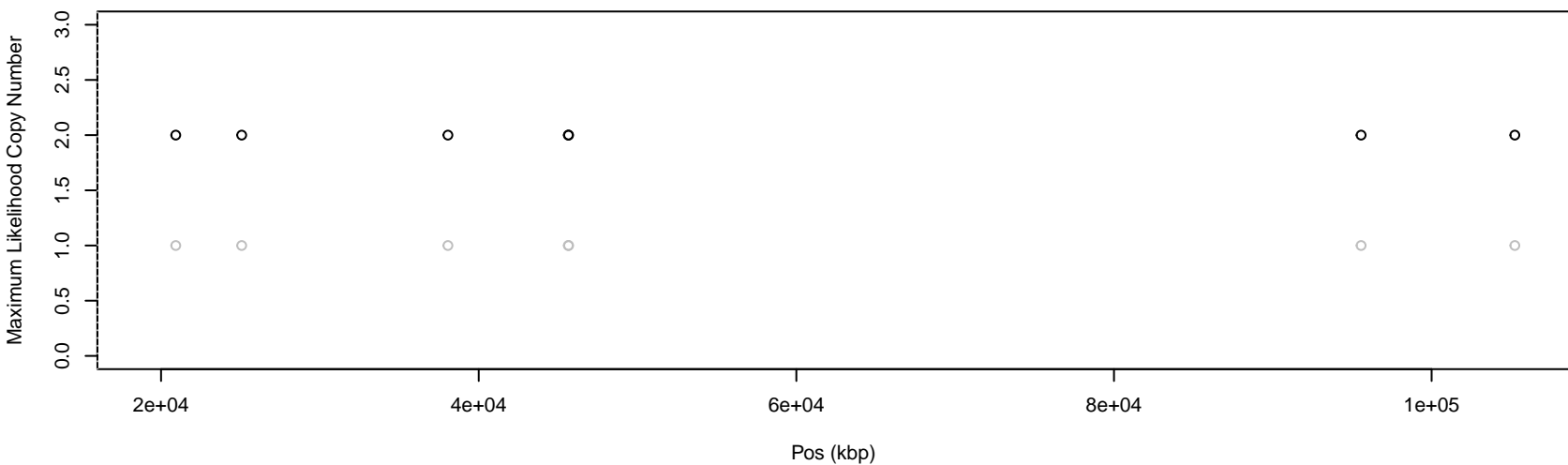
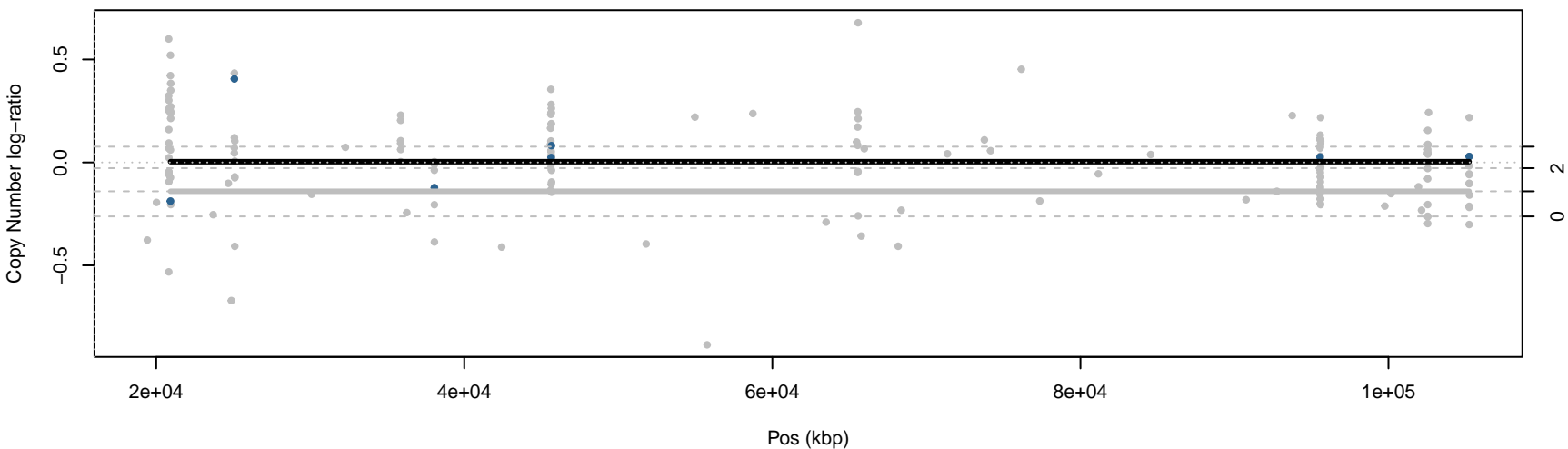
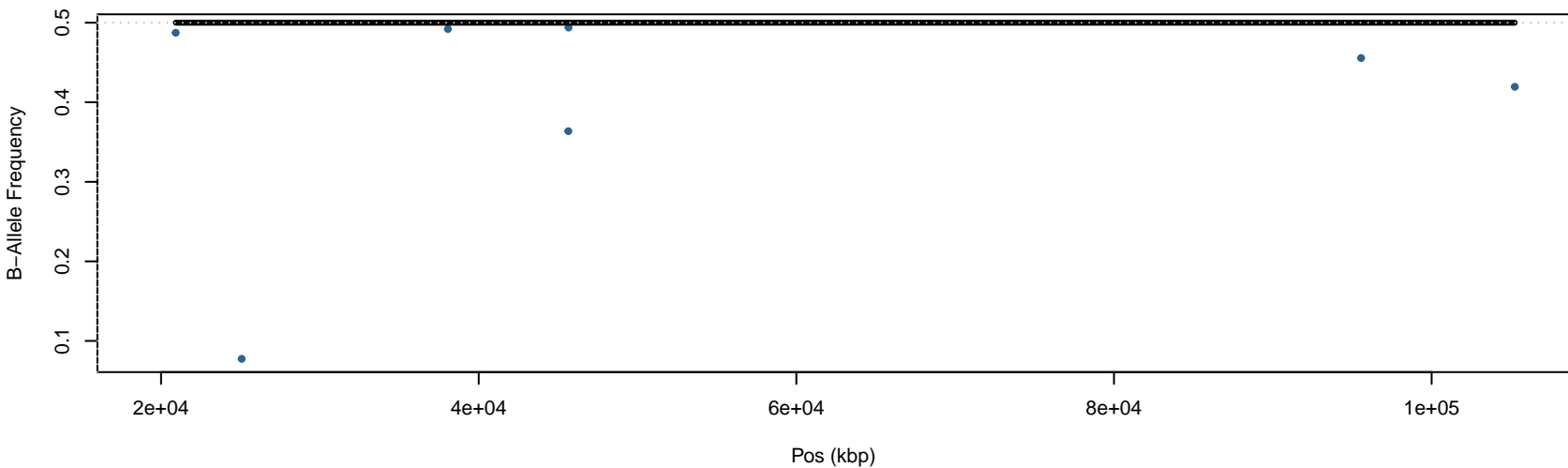
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 10



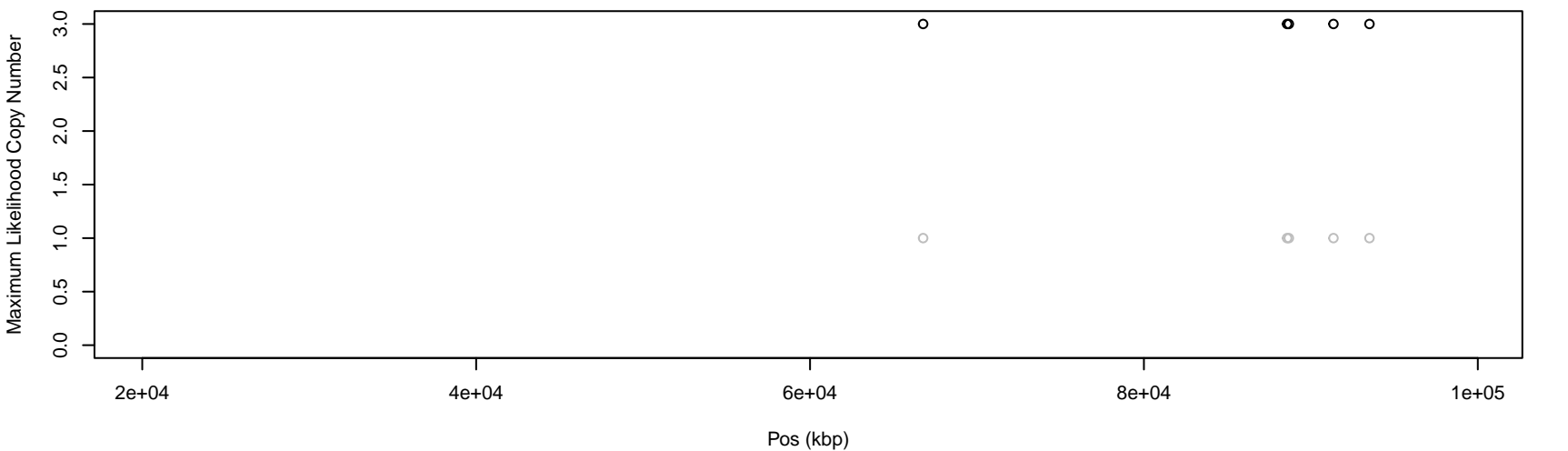
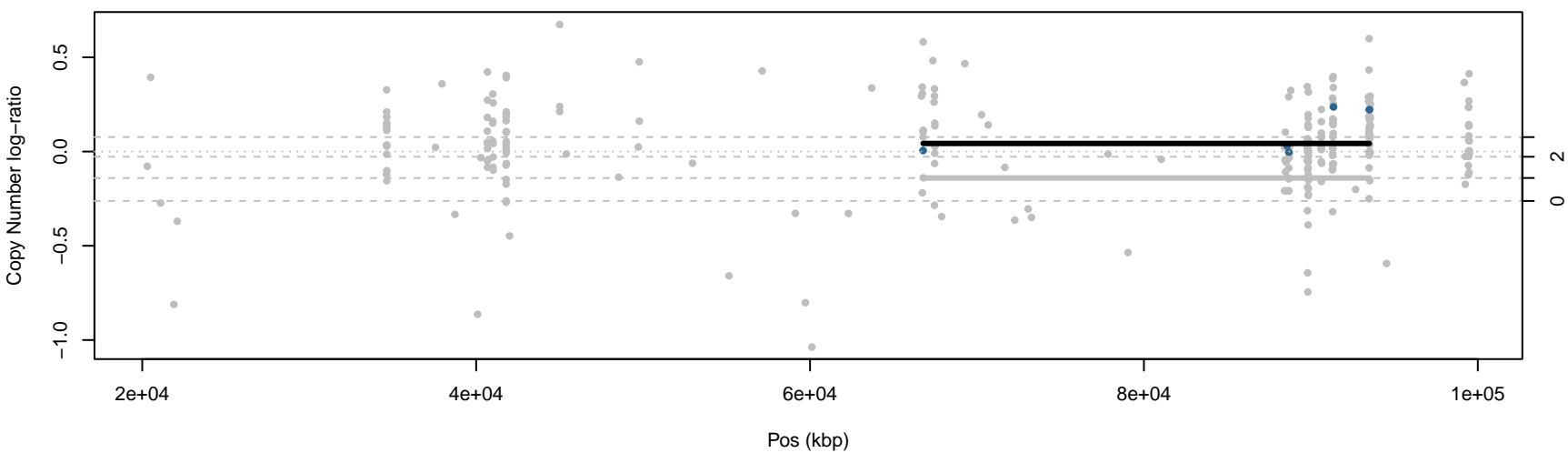
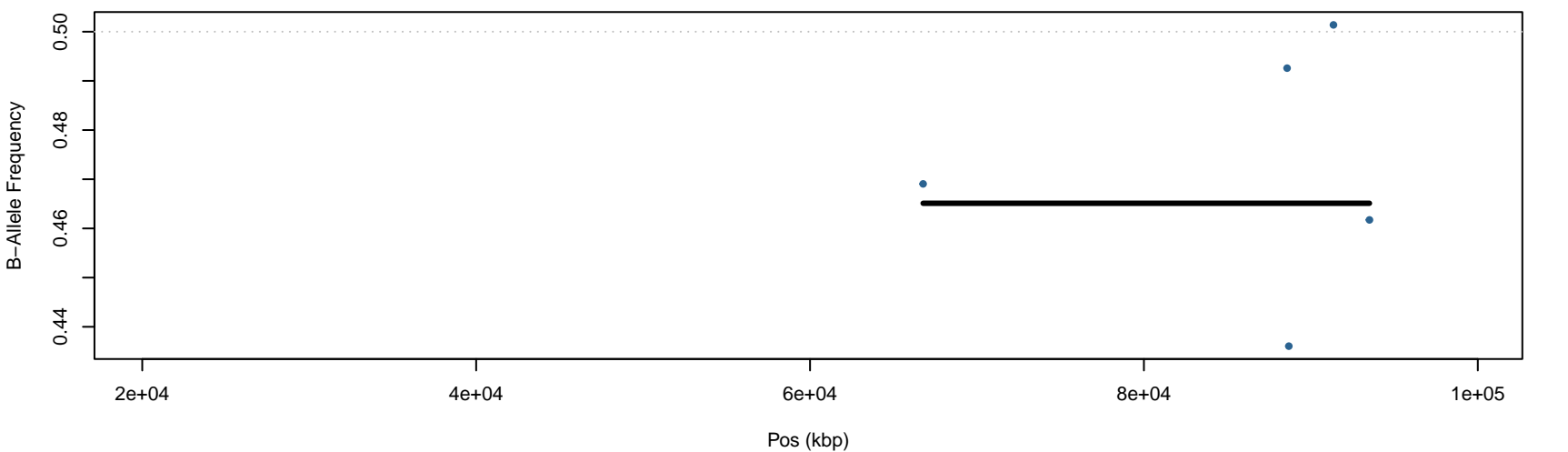
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 12



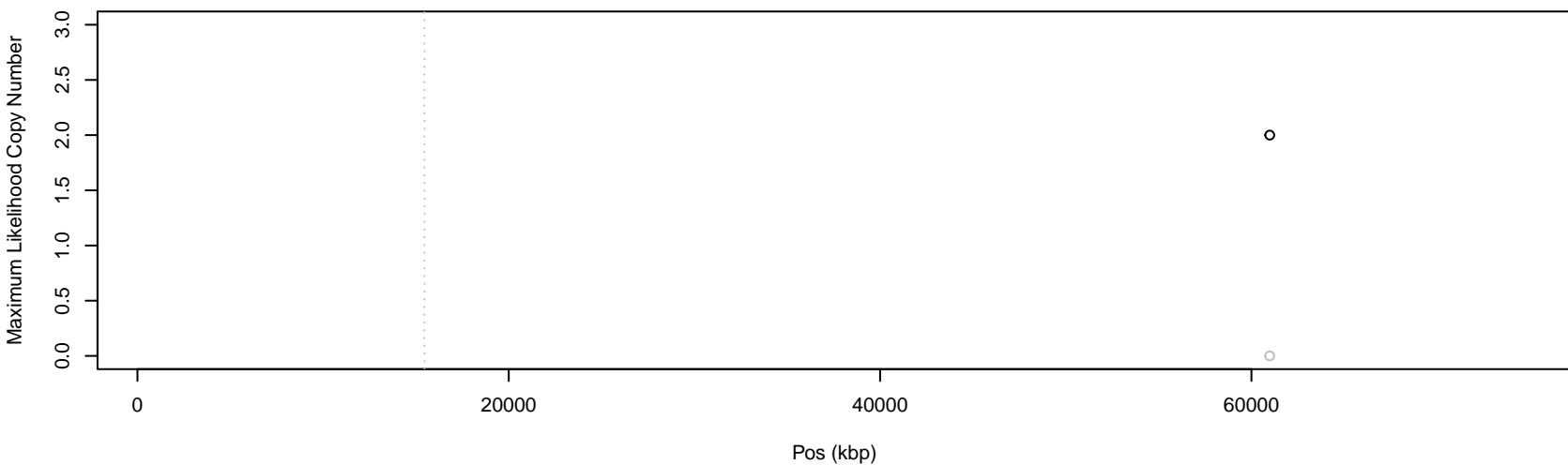
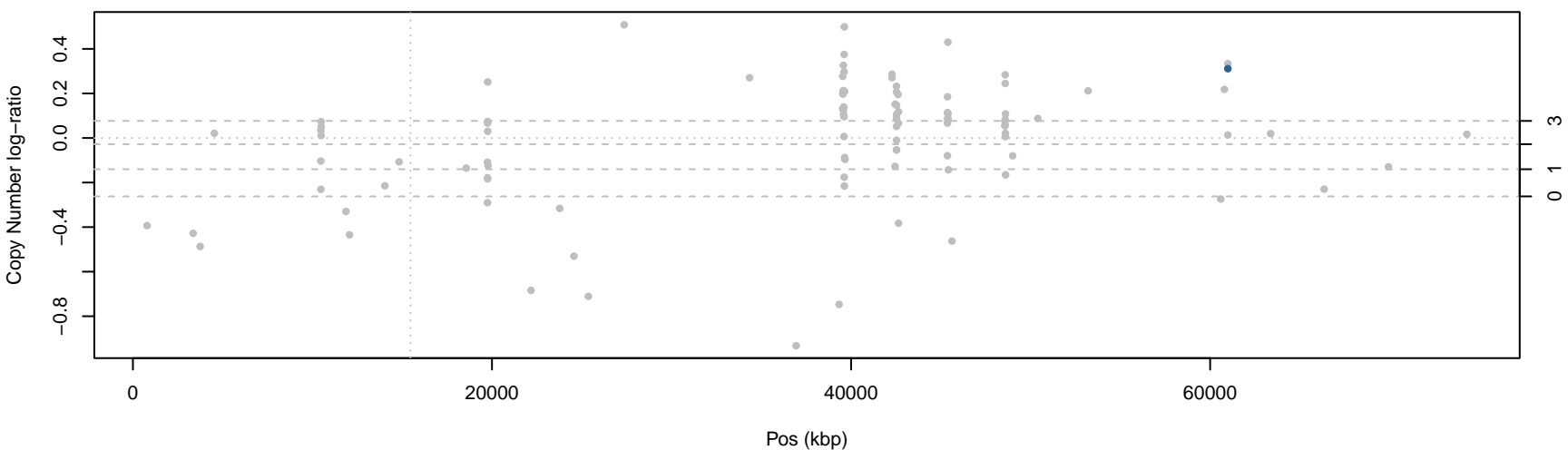
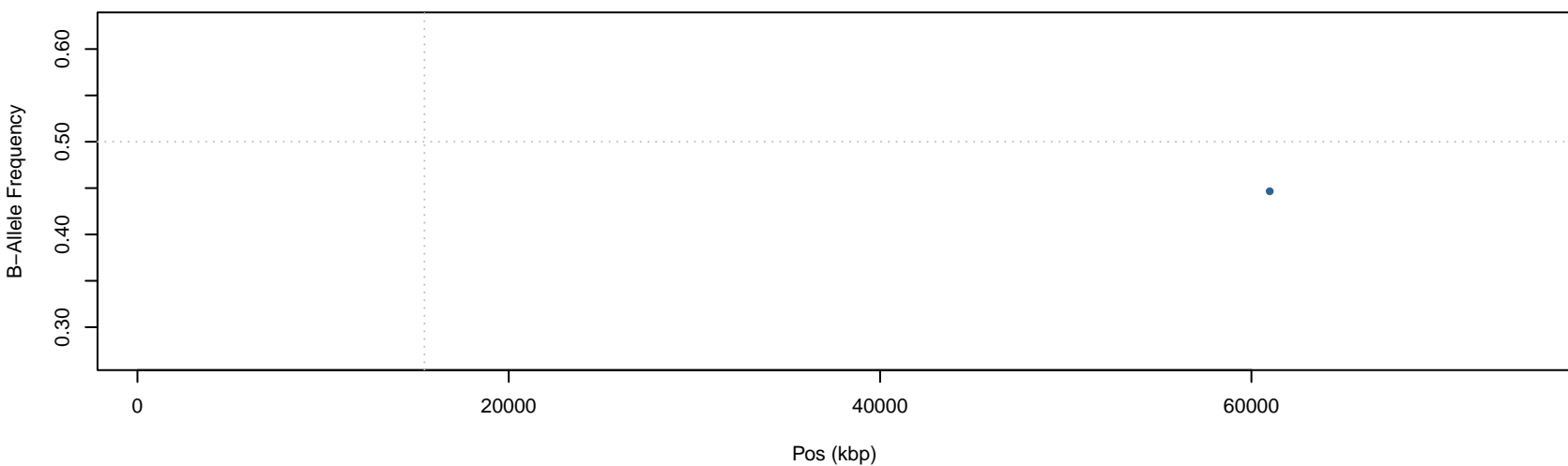
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 14



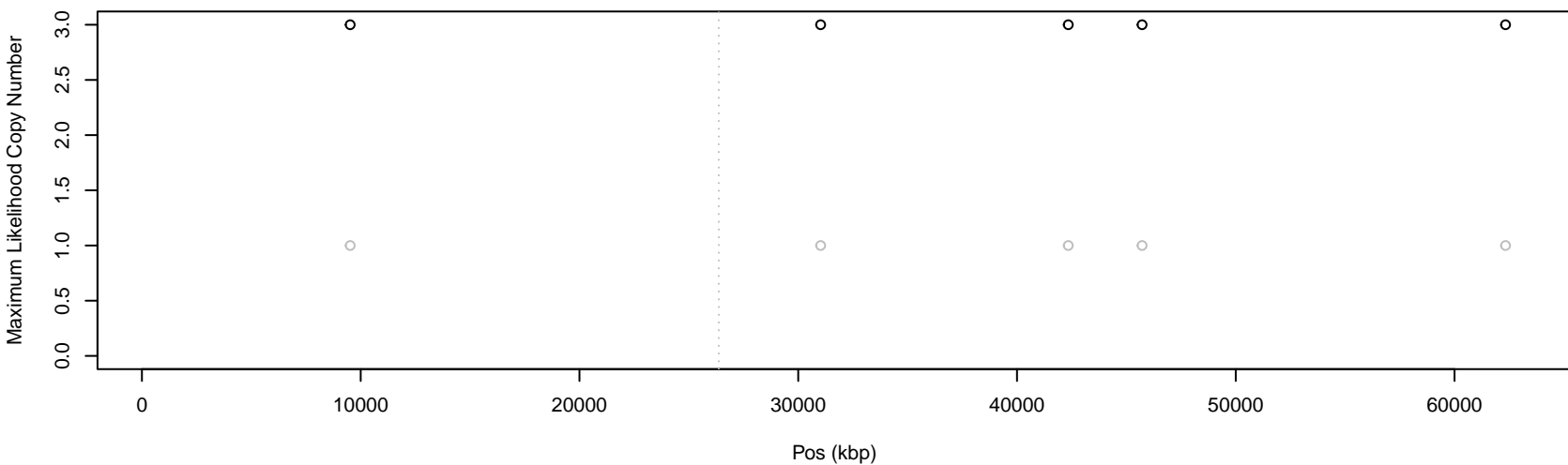
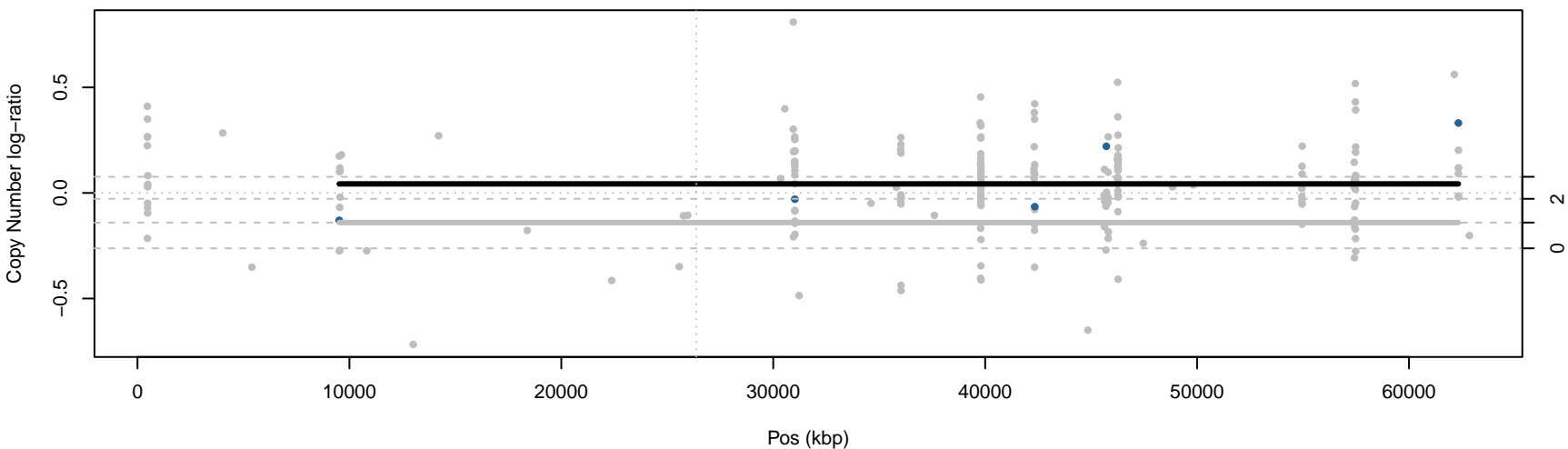
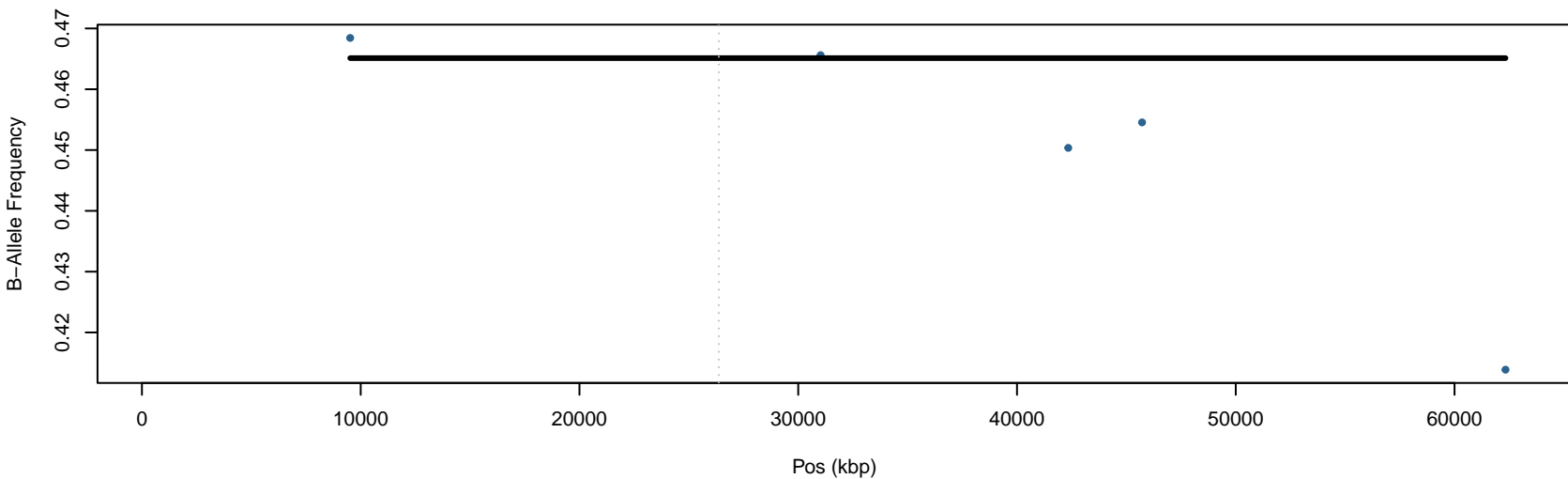
Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 15



Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 18



Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 20





Purity: 0.15 Tumor ploidy: 2.259 SNV log-likelihood: -16.35 GoF: 95.8% Mean coverage: 315;324 Chromosome: 22

