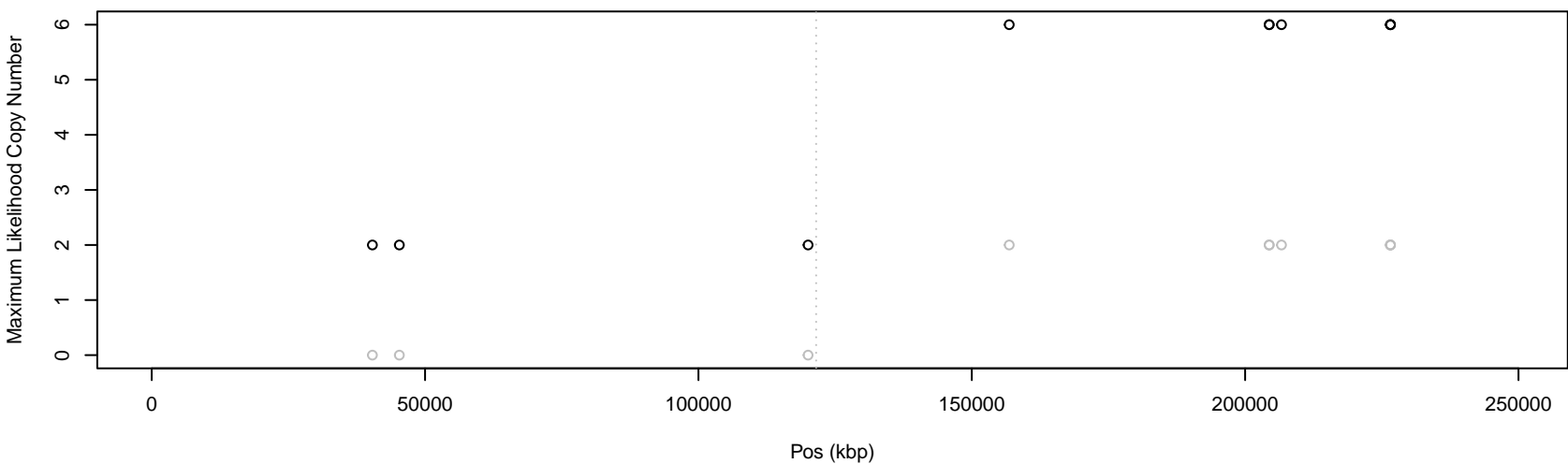
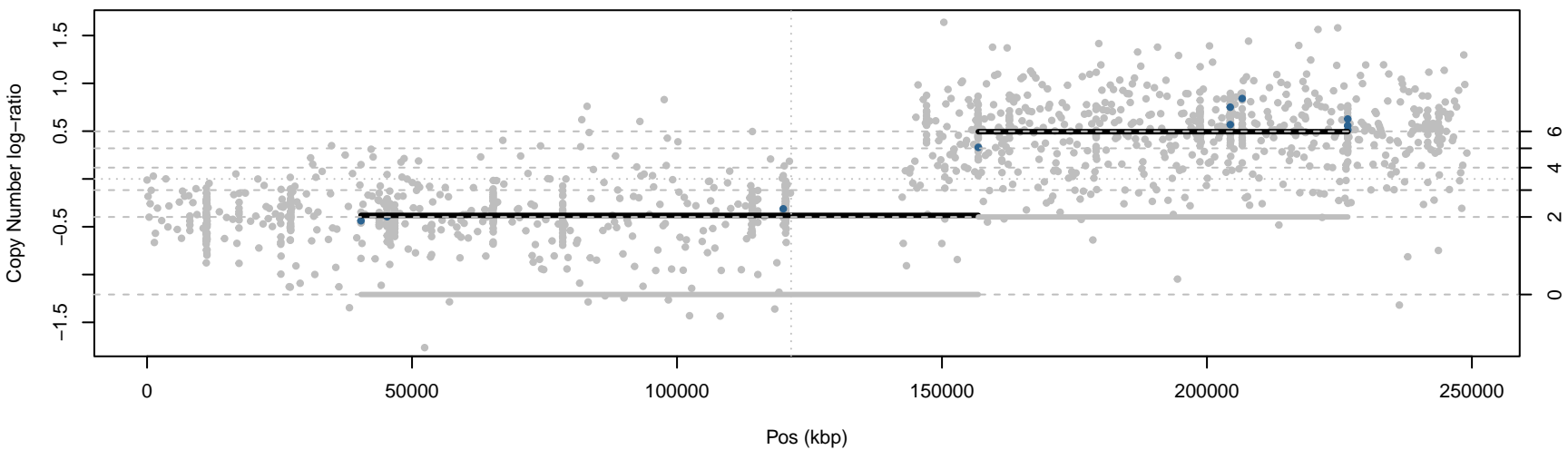
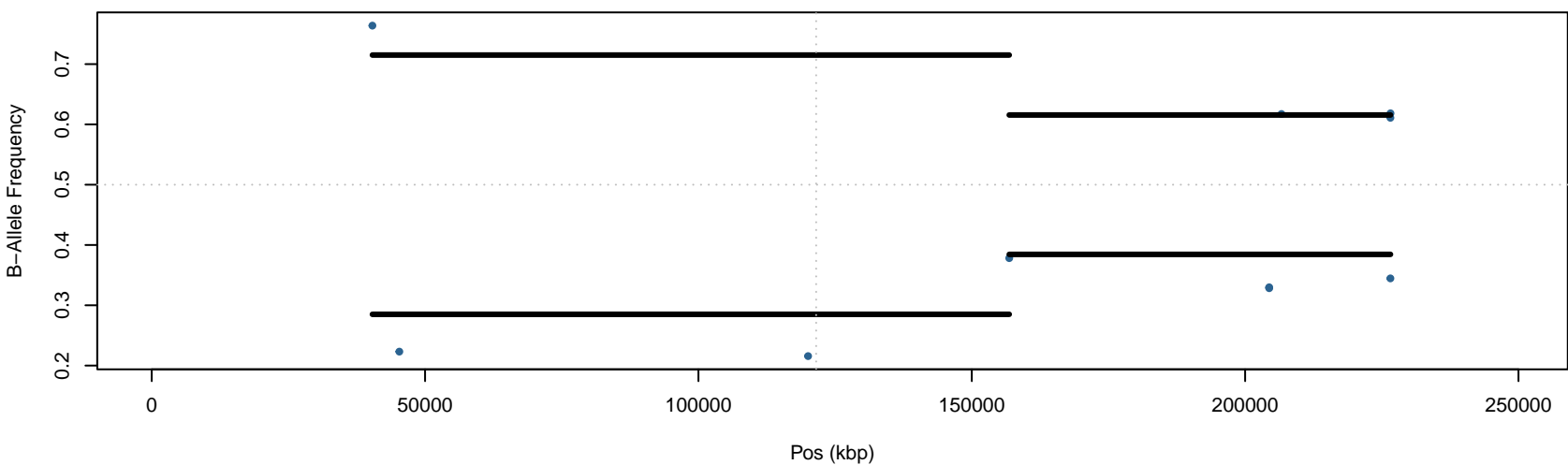
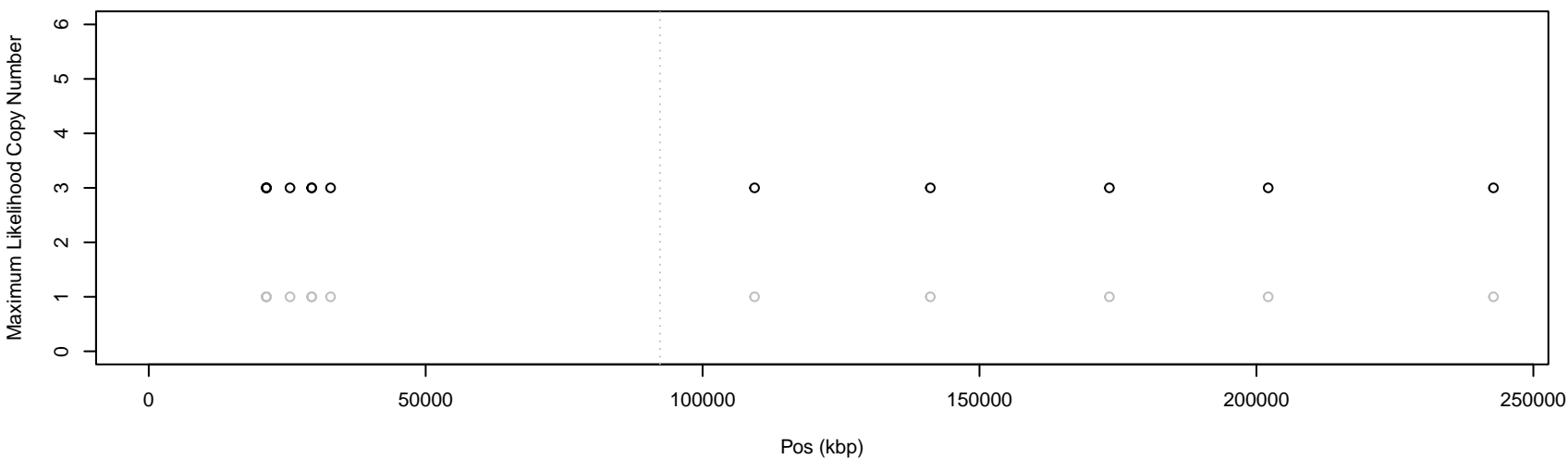
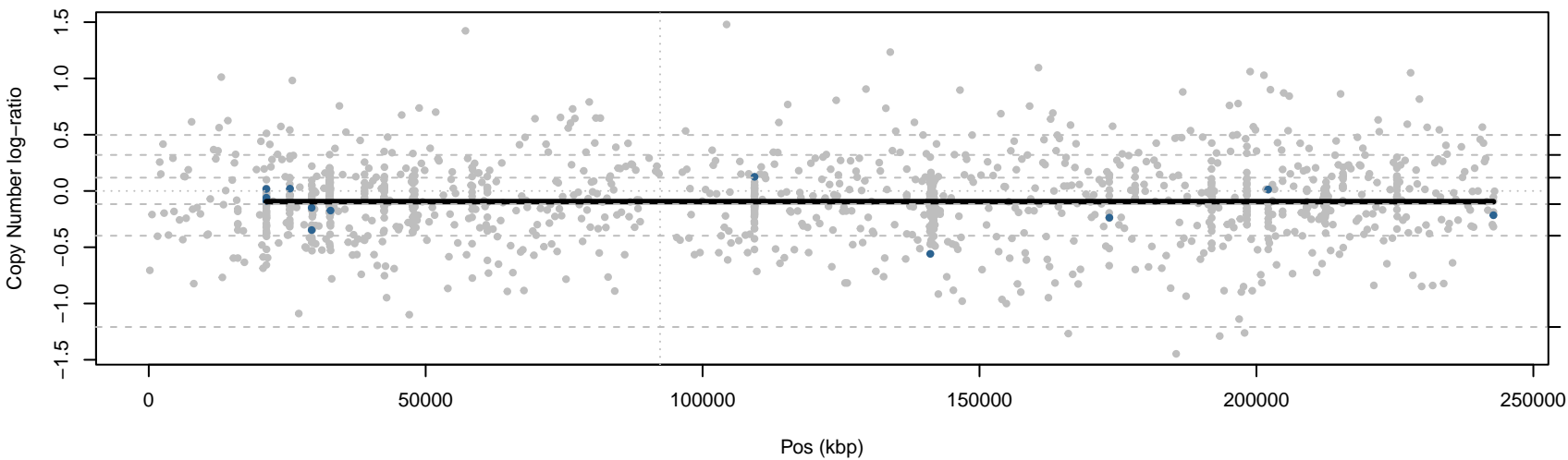
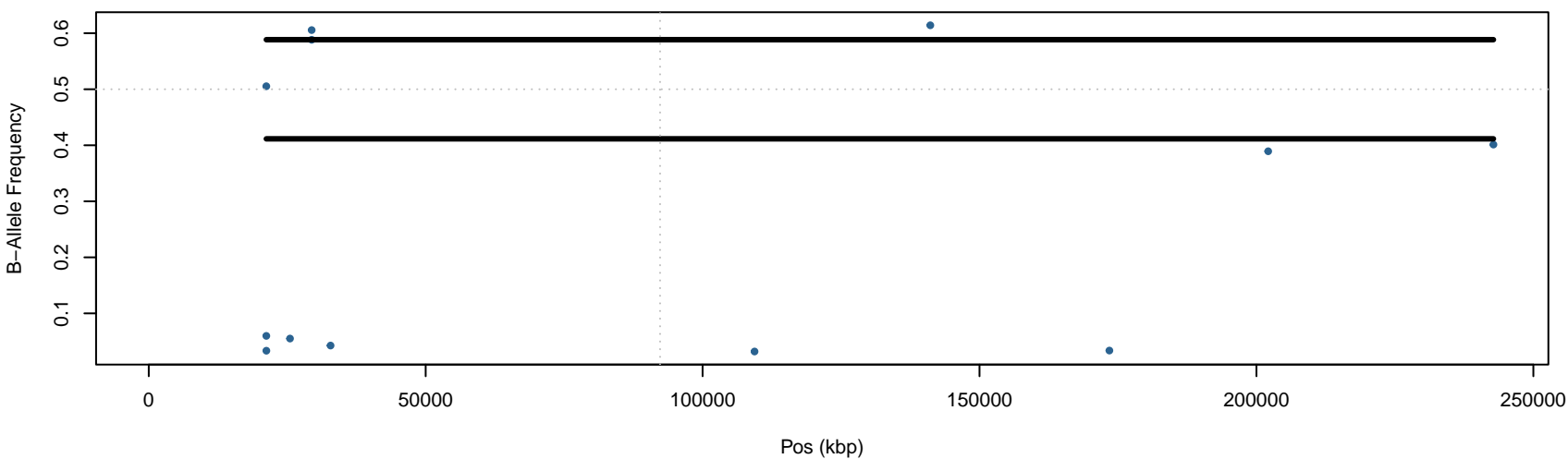


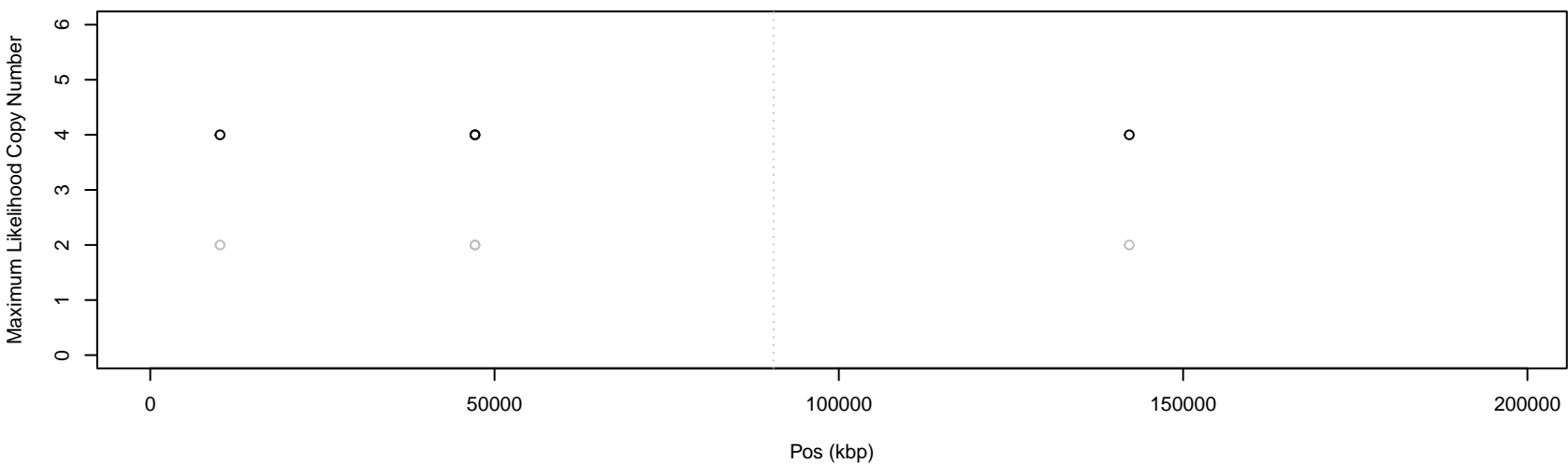
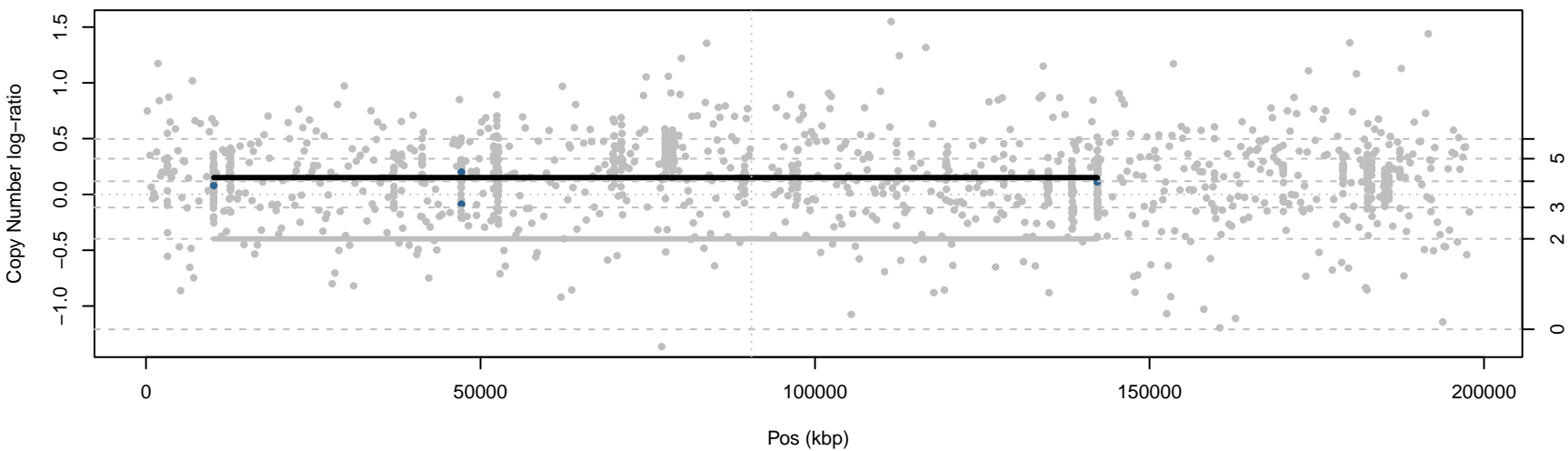
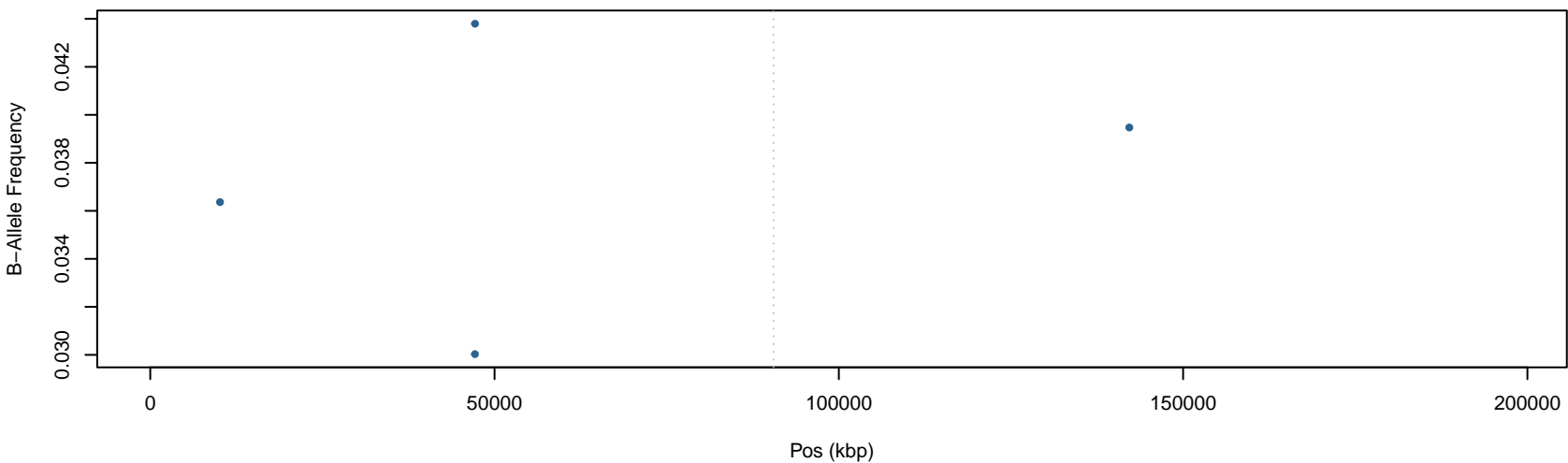
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 1



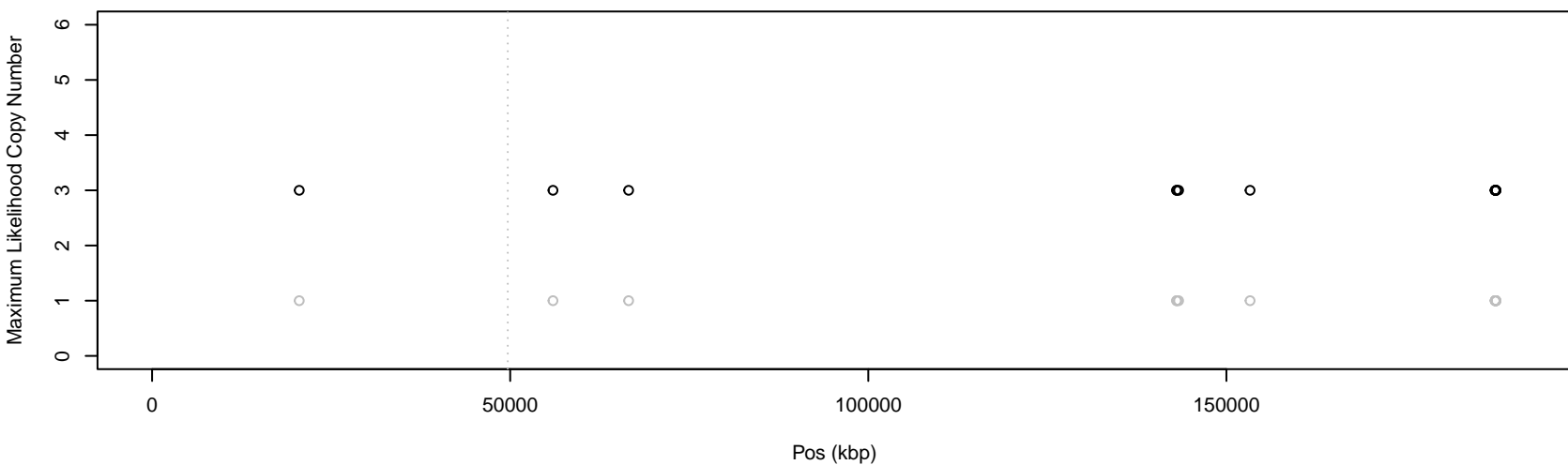
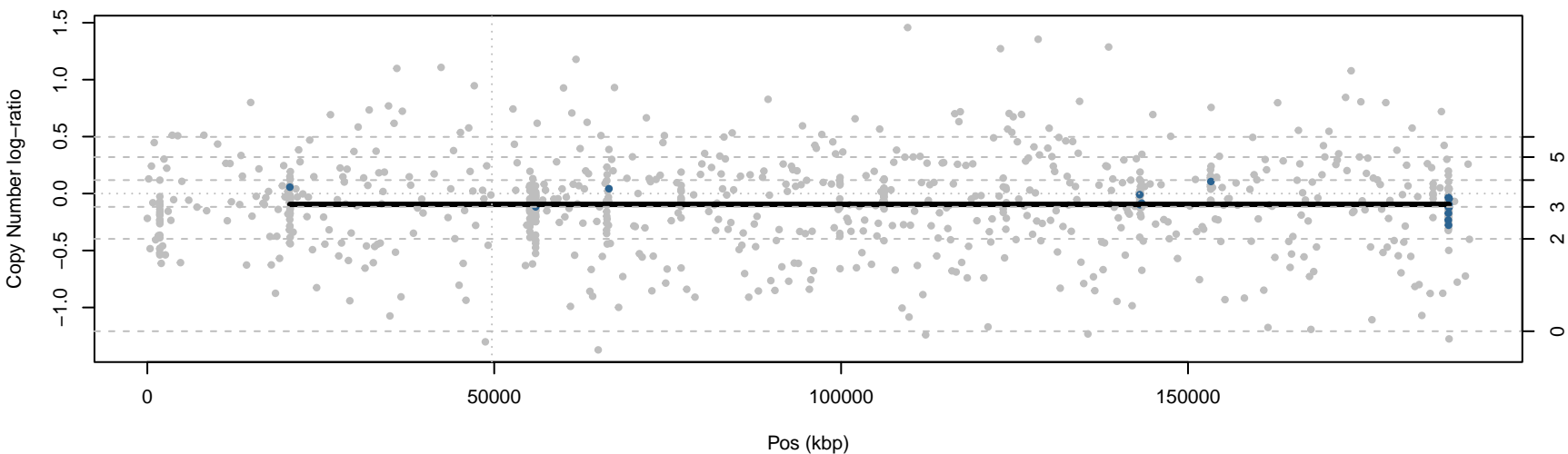
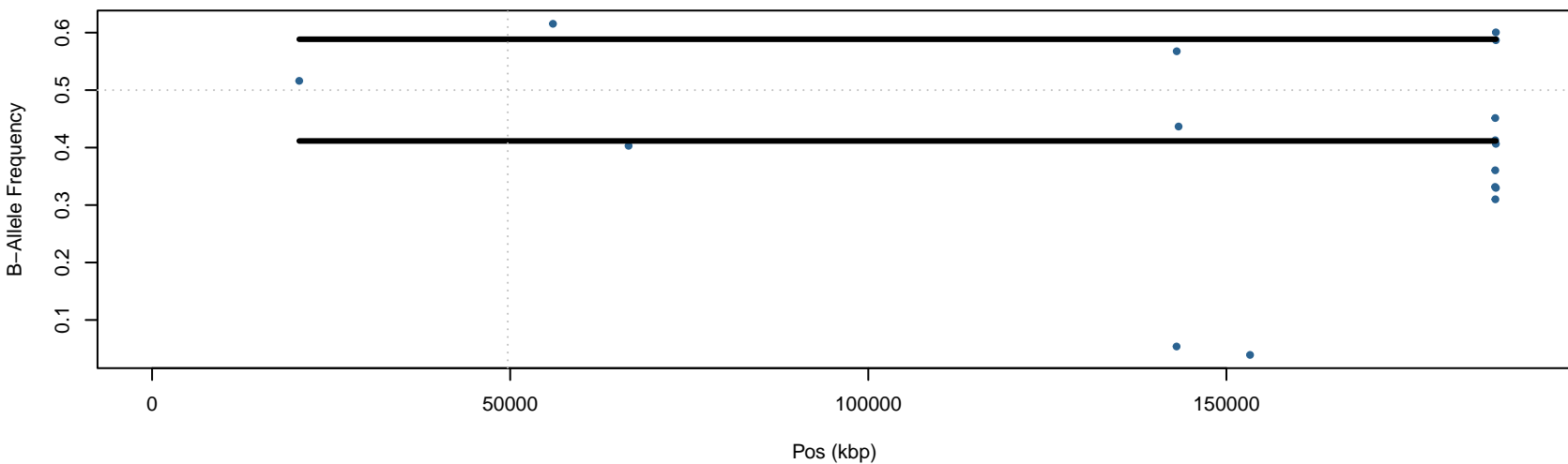
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 2



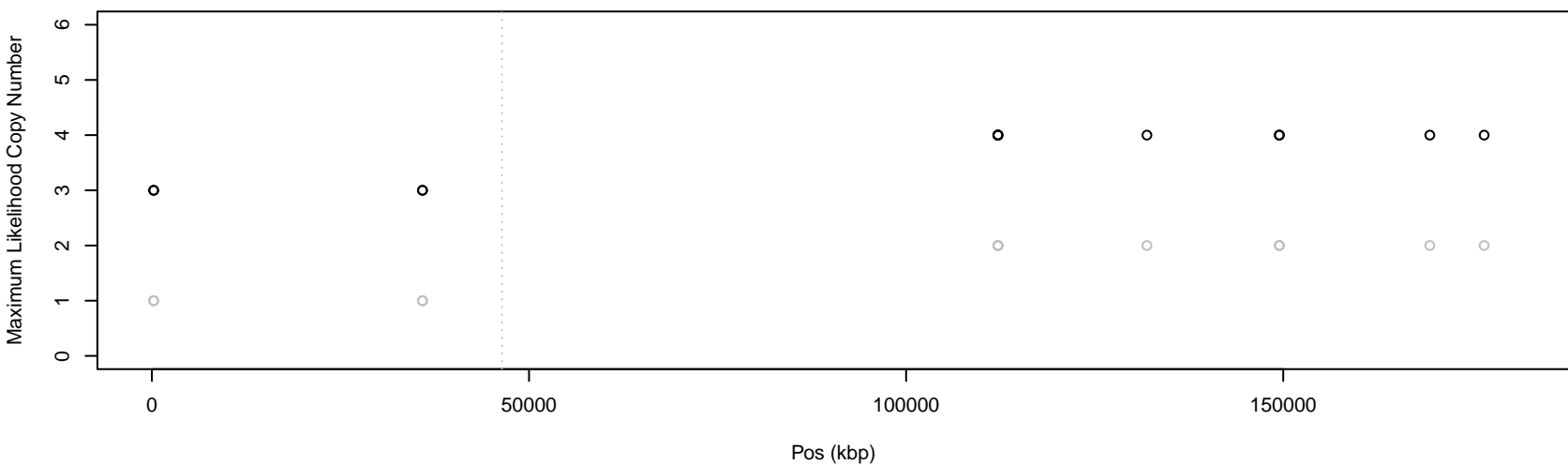
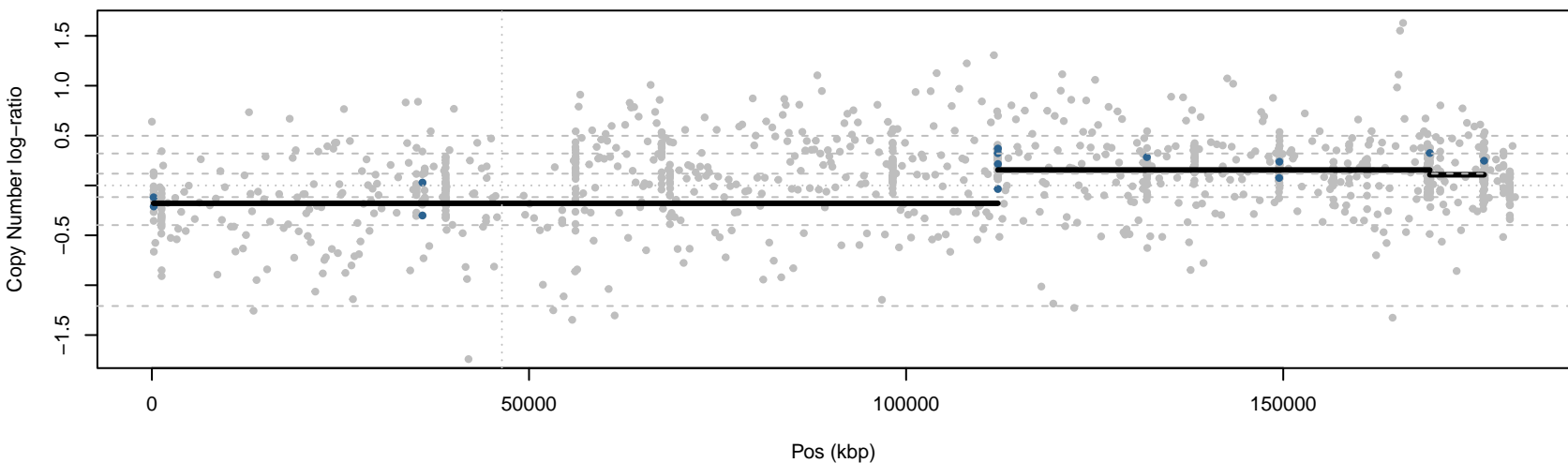
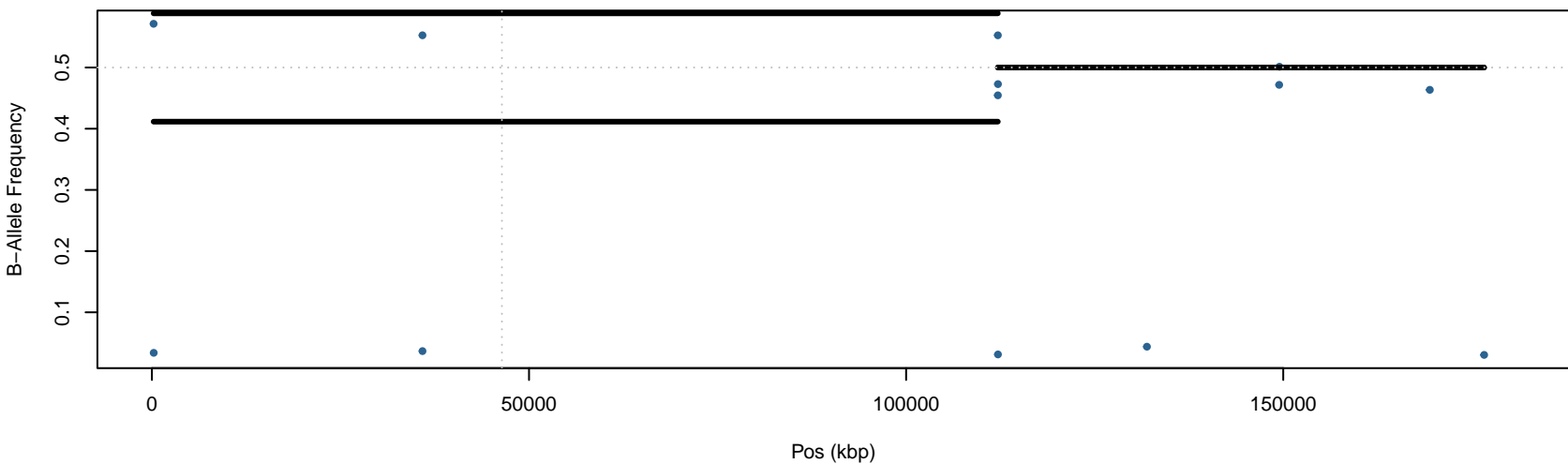
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 3



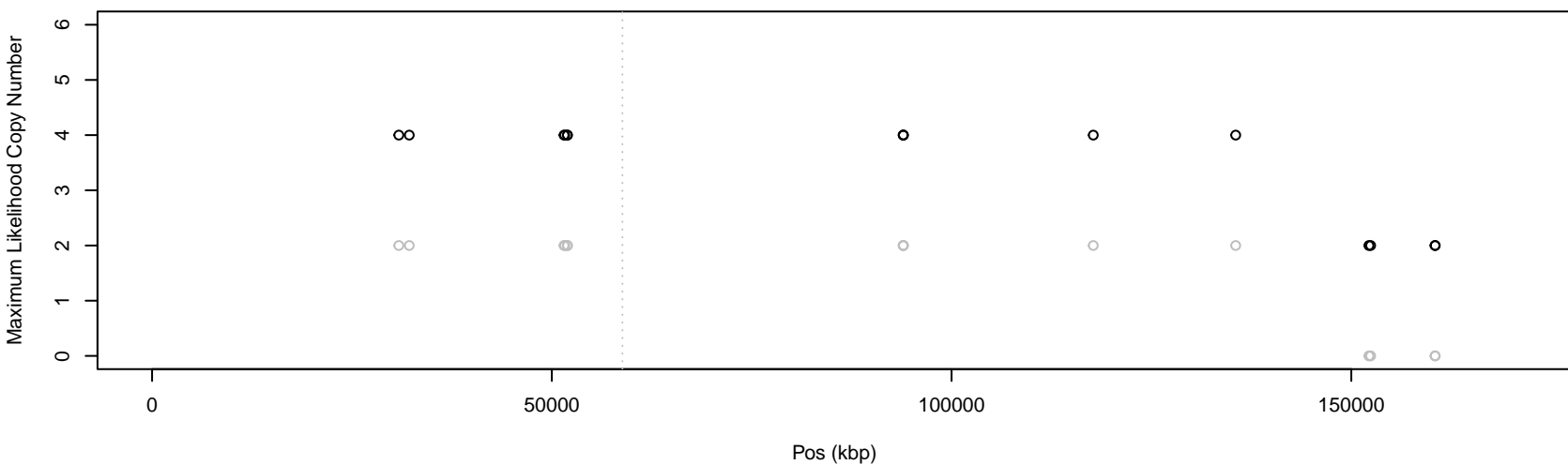
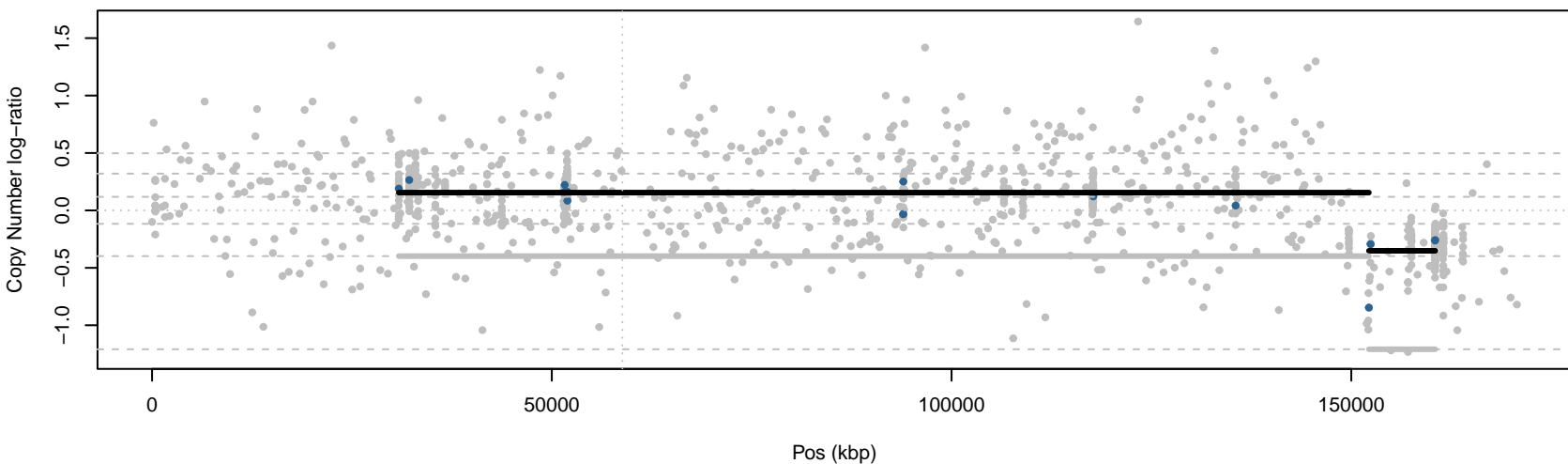
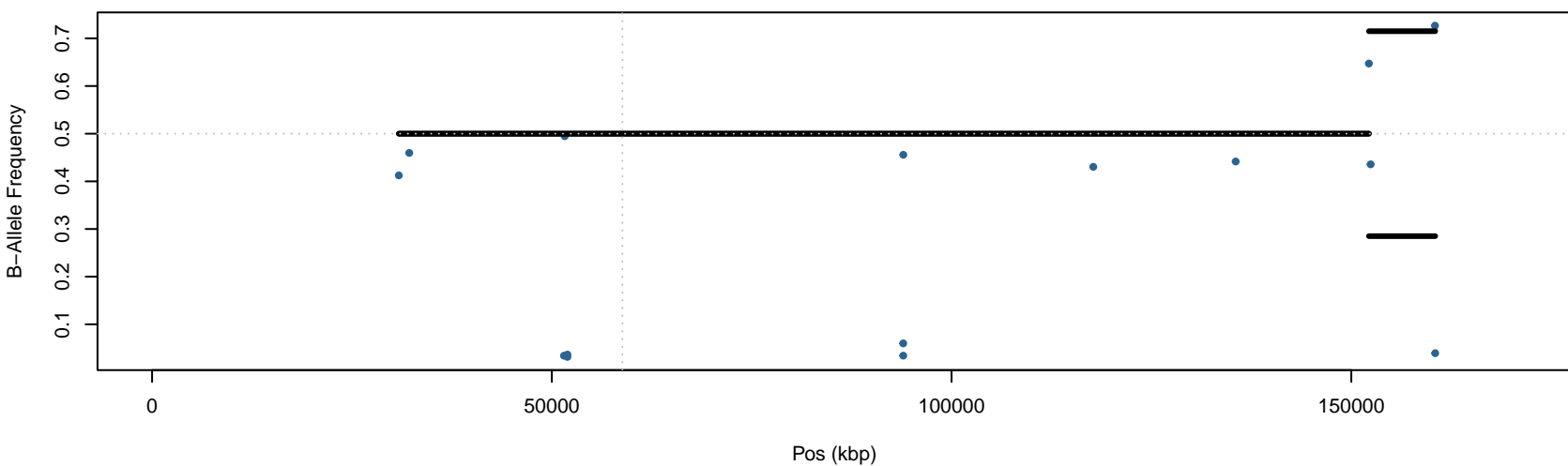
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 4



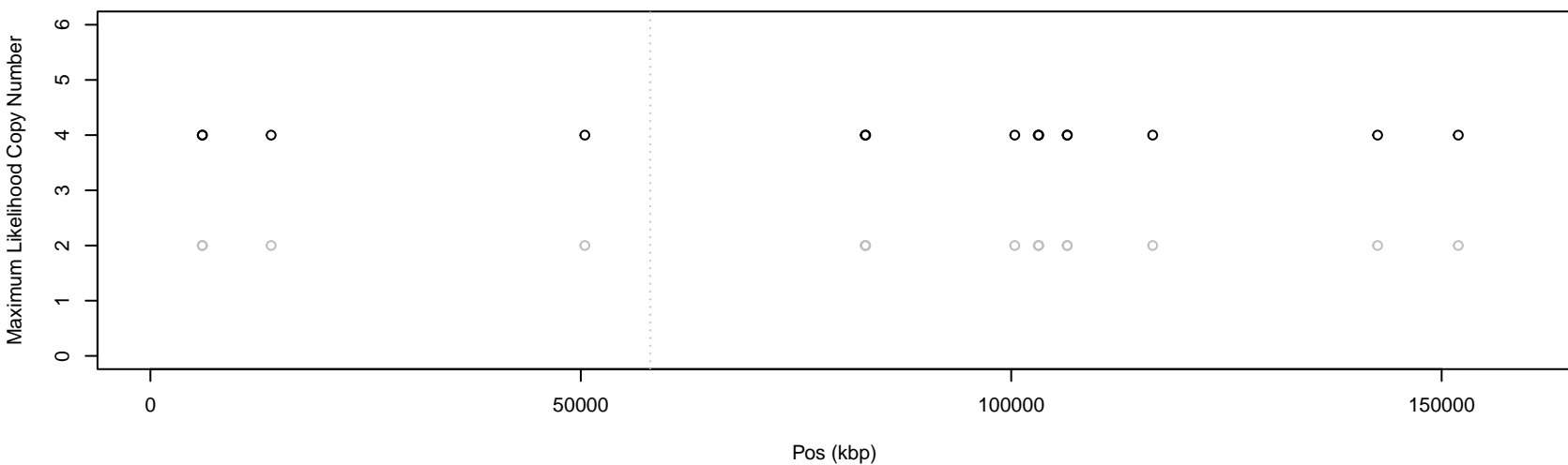
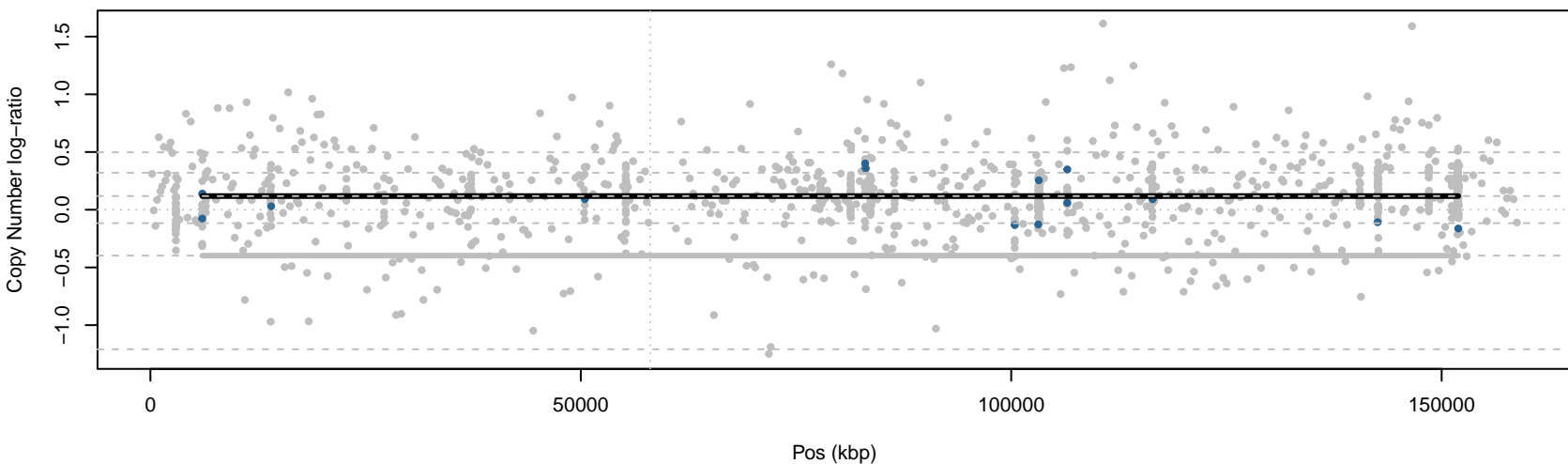
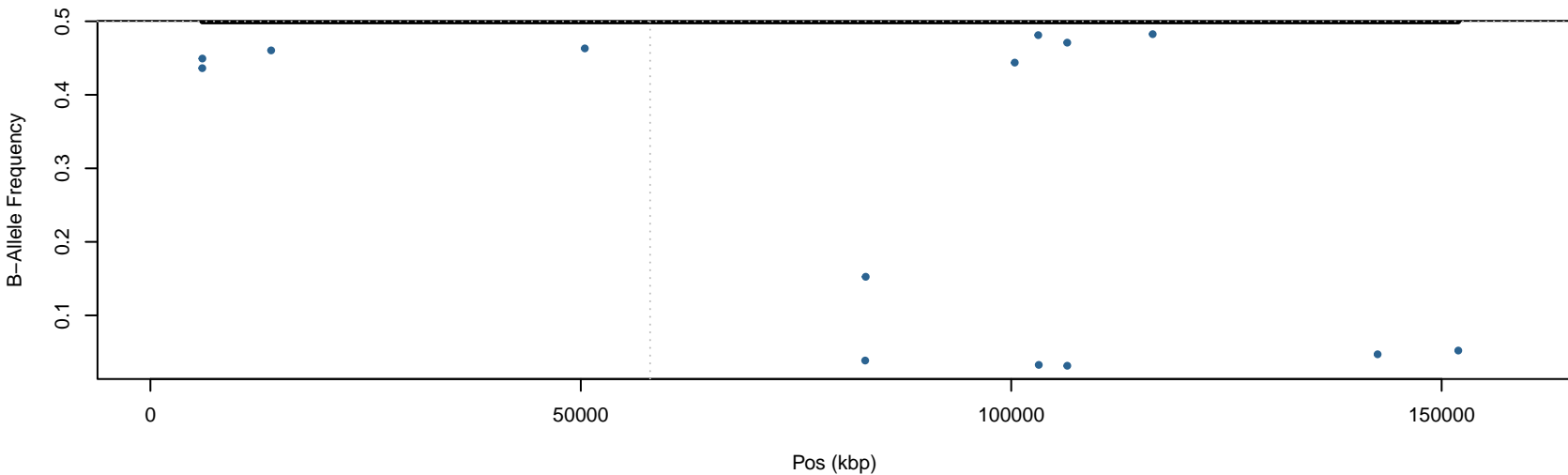
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 5



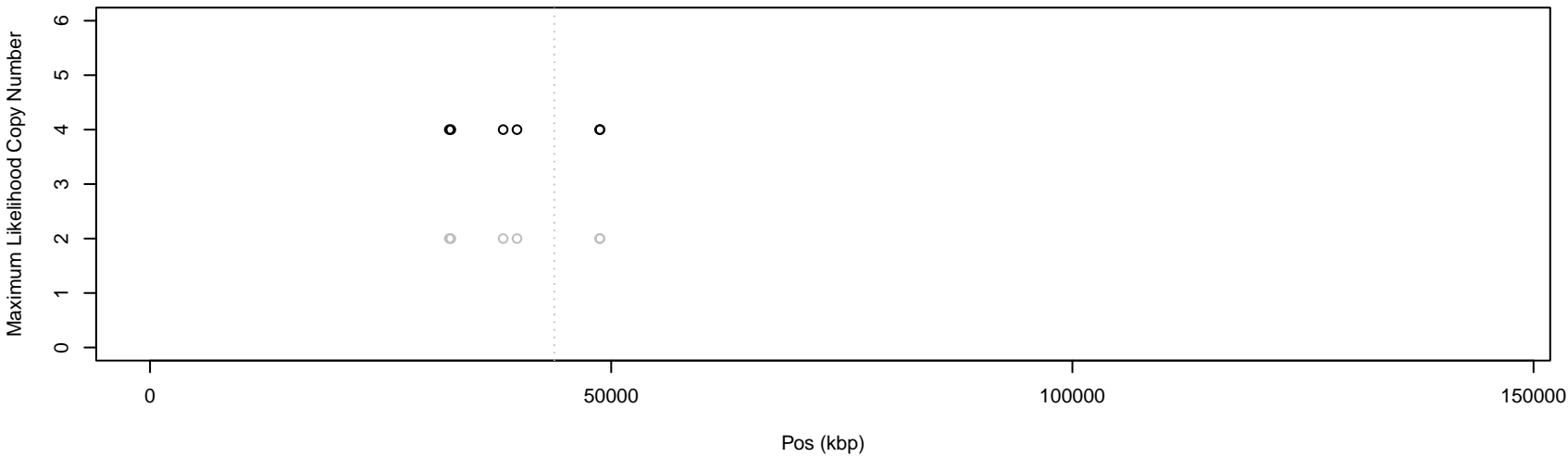
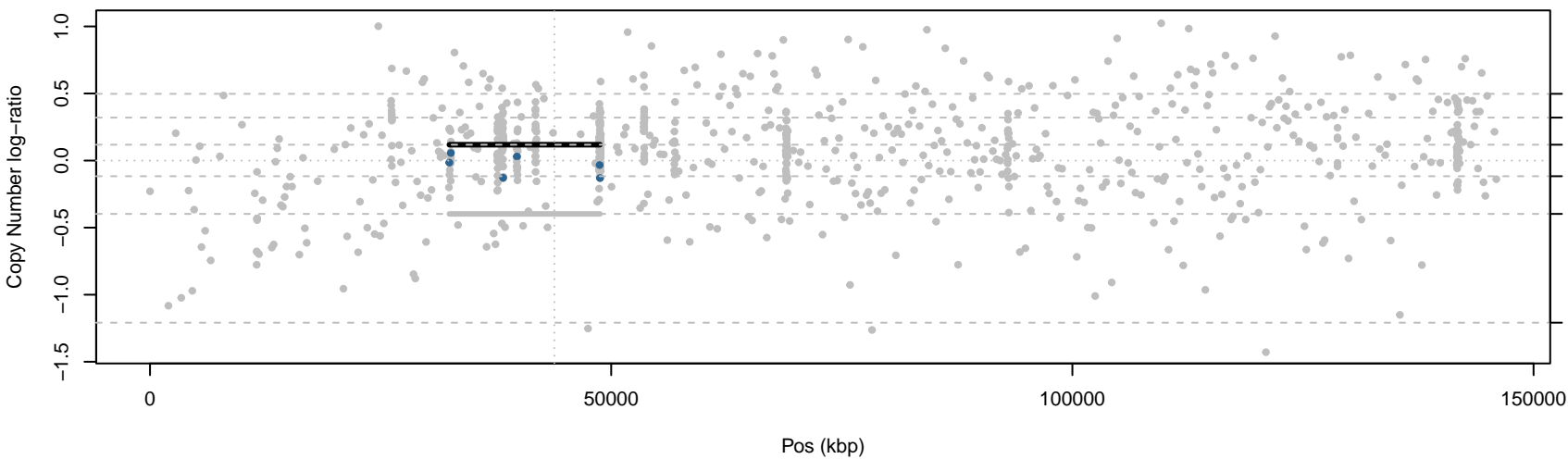
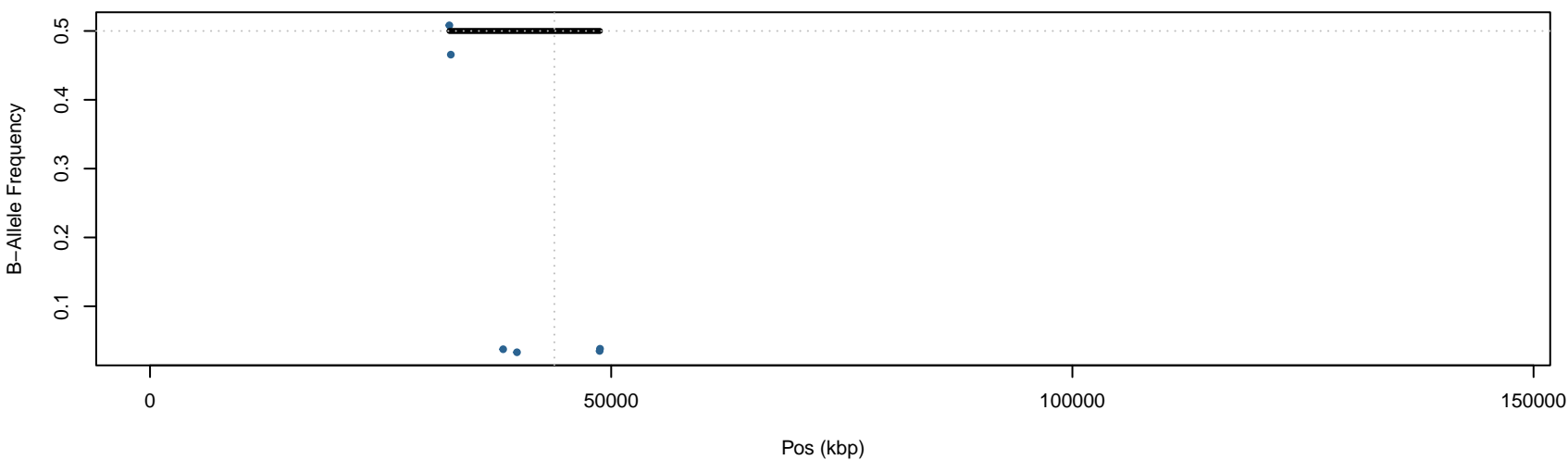
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 6



Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 7

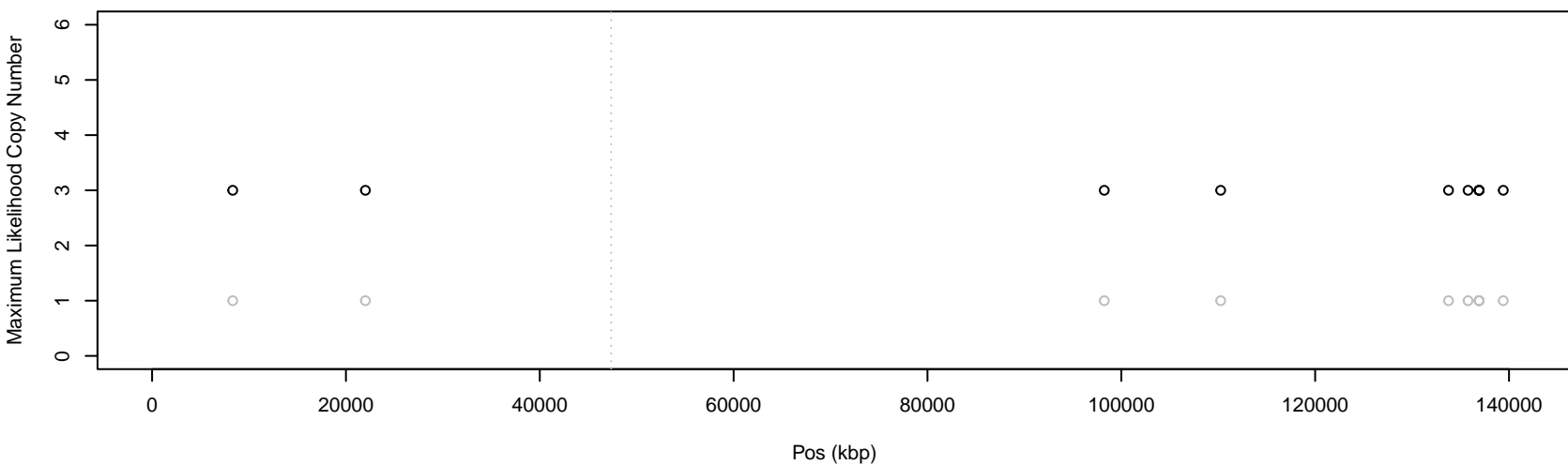
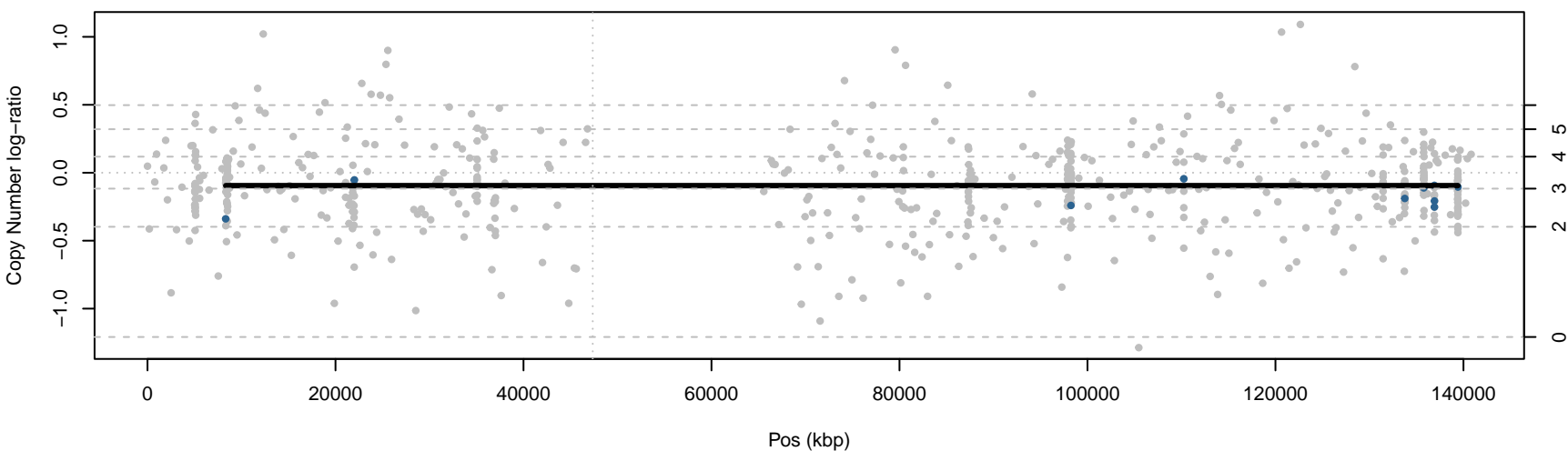
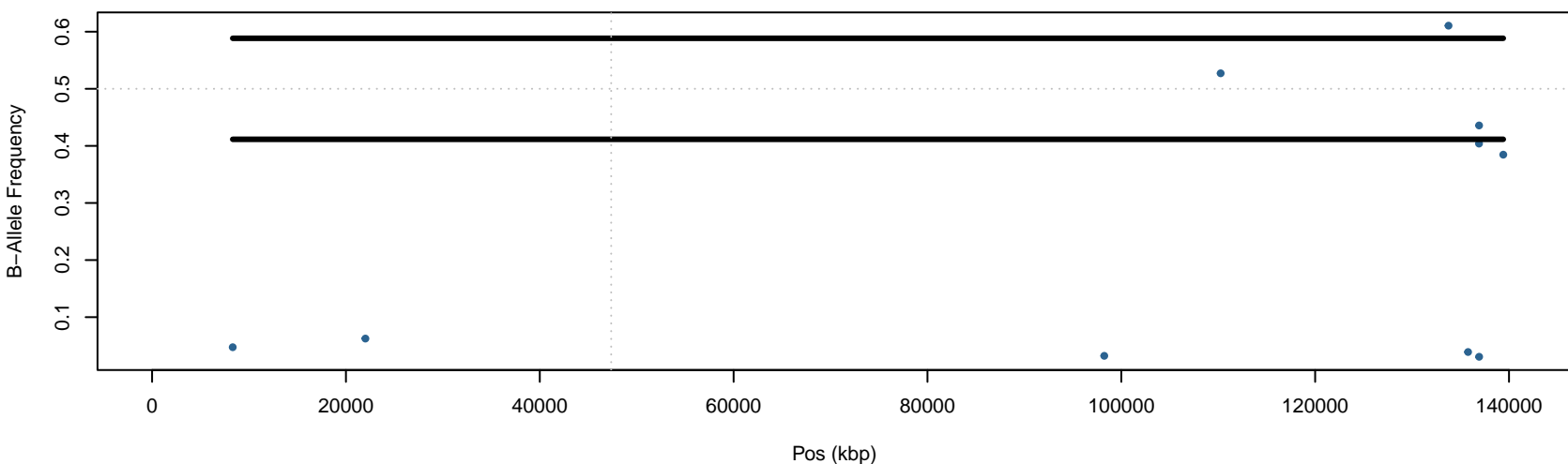


Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 8

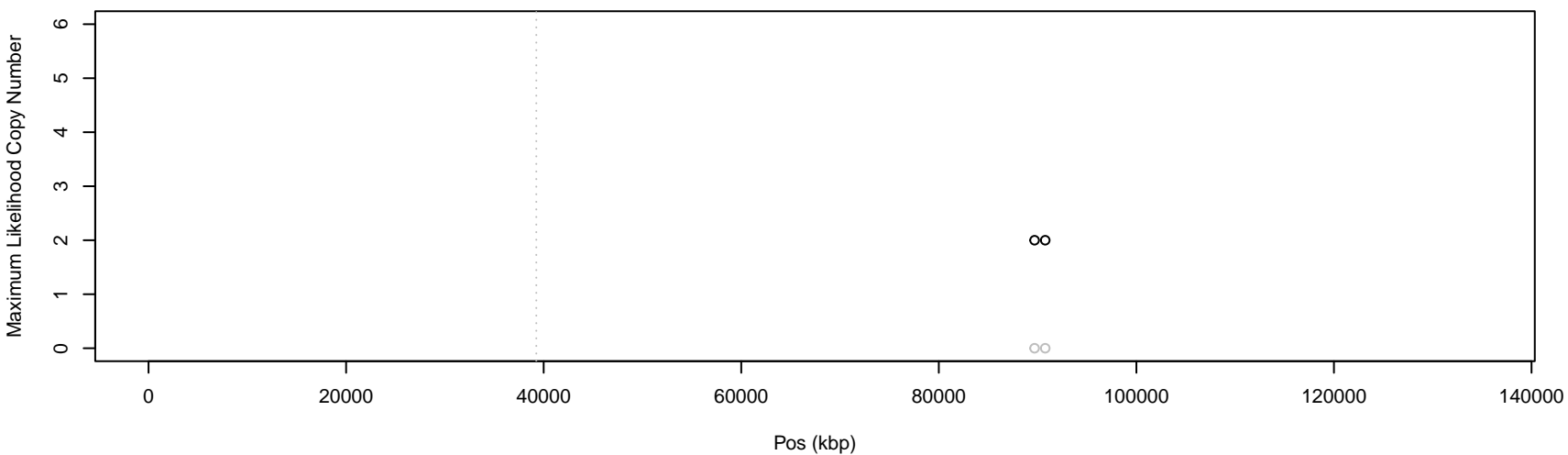
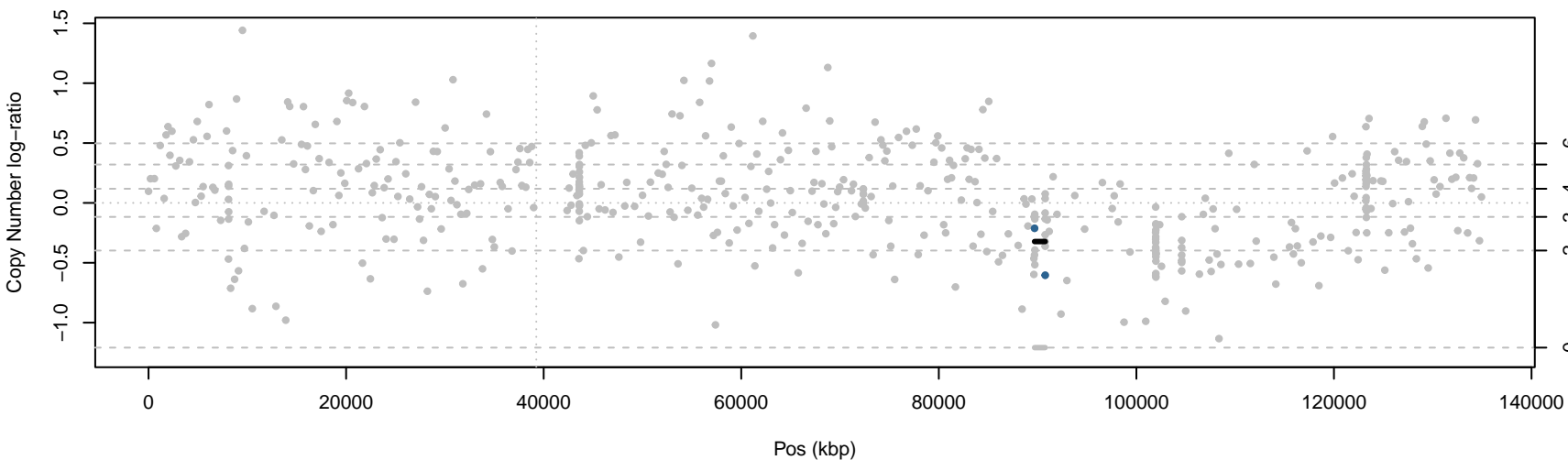
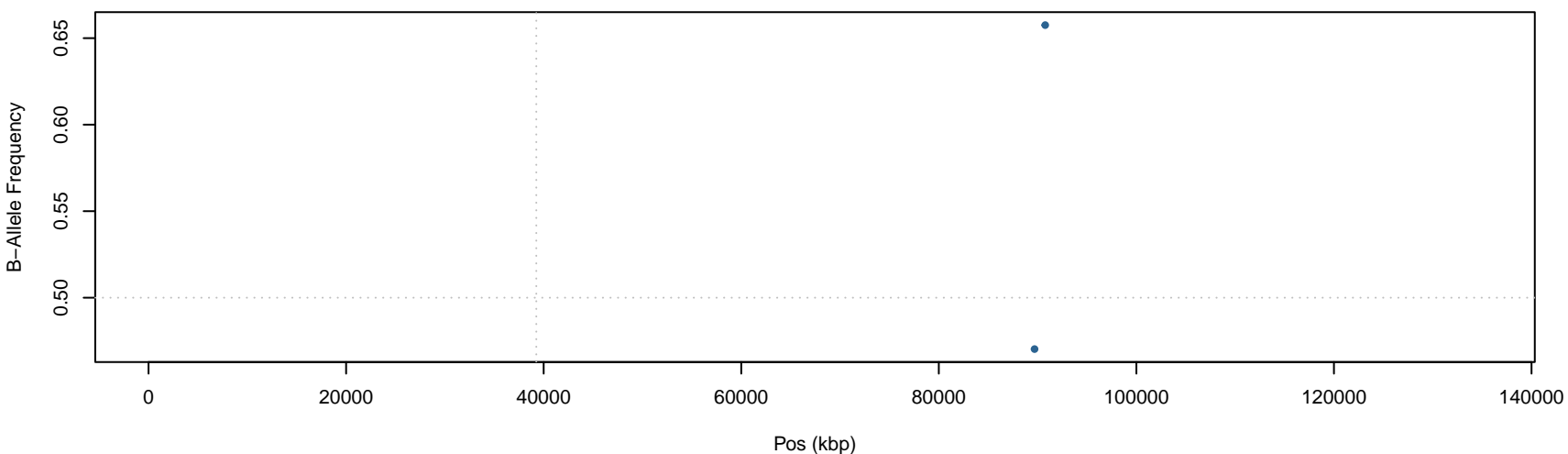




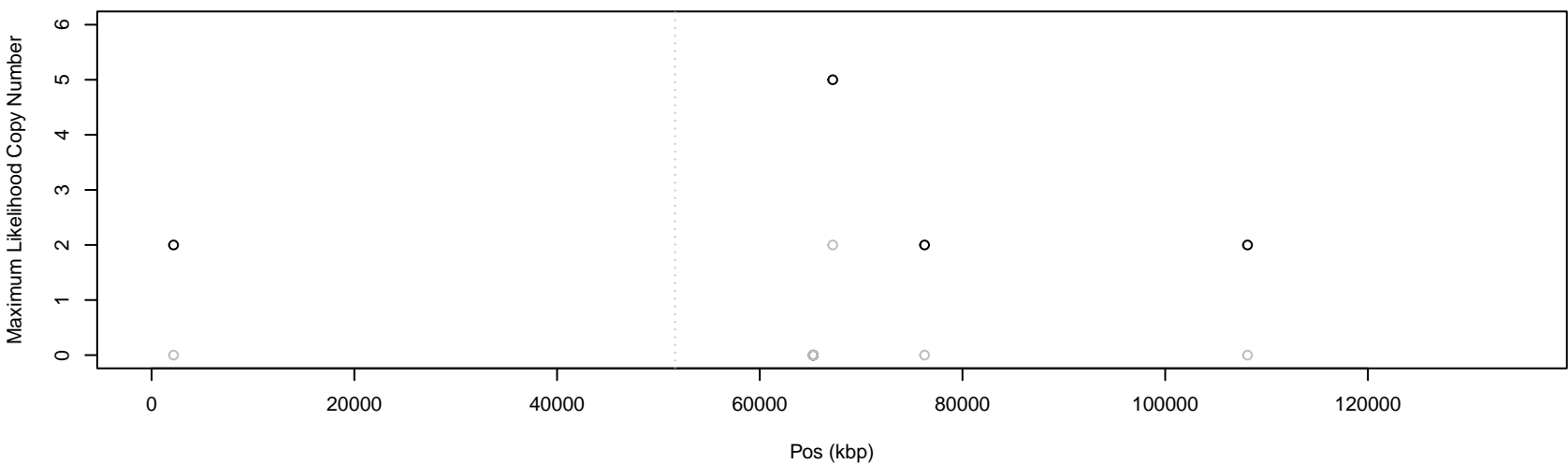
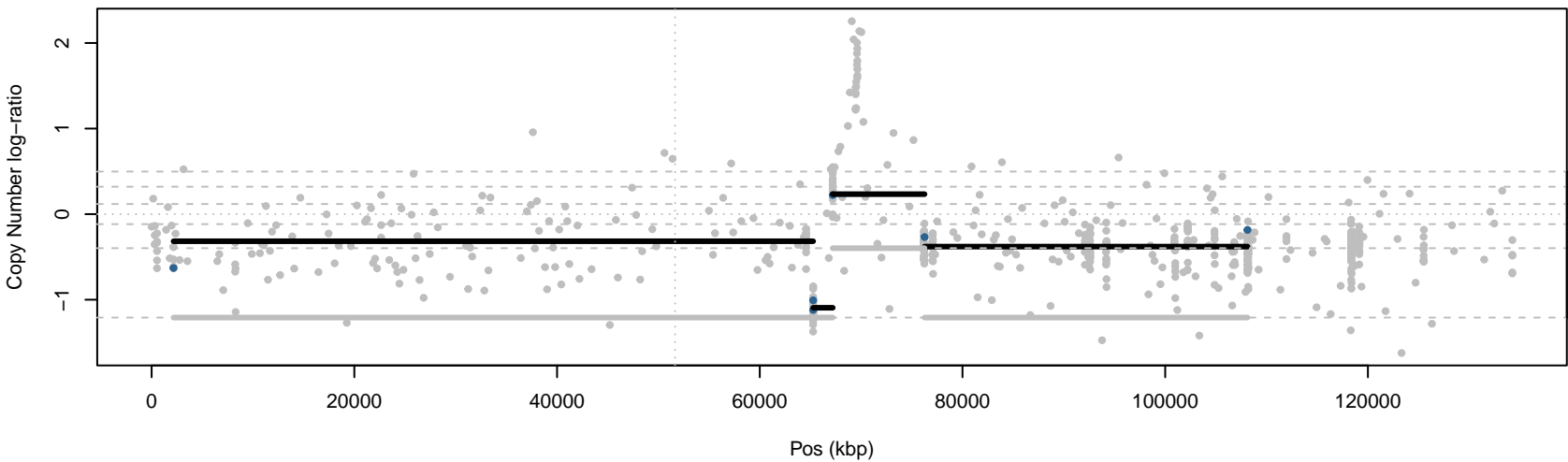
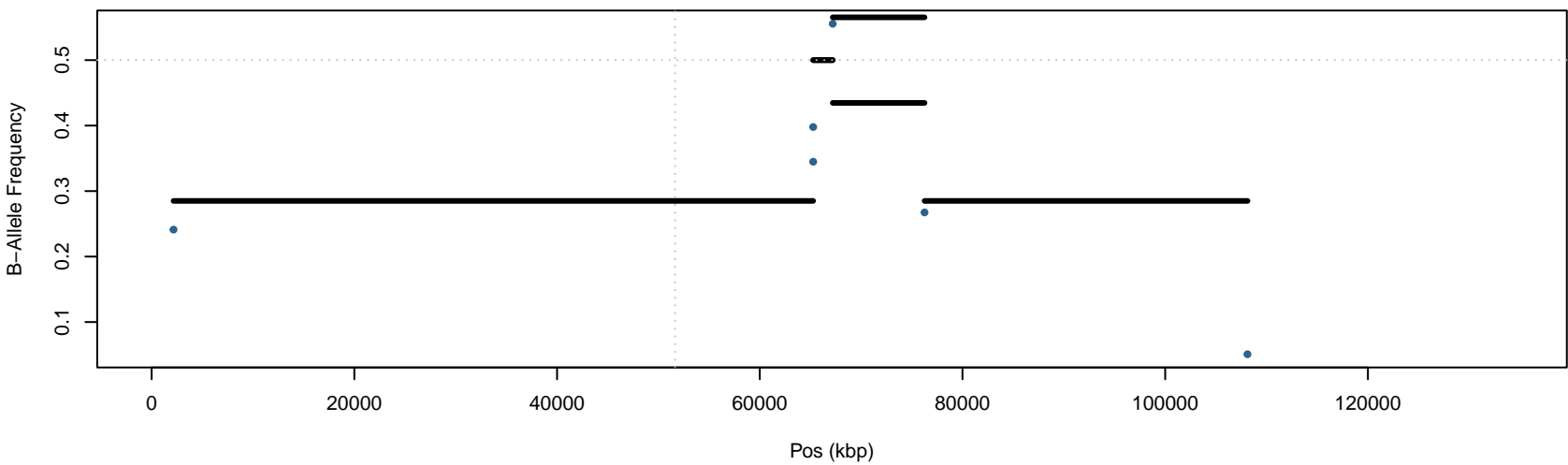
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 9



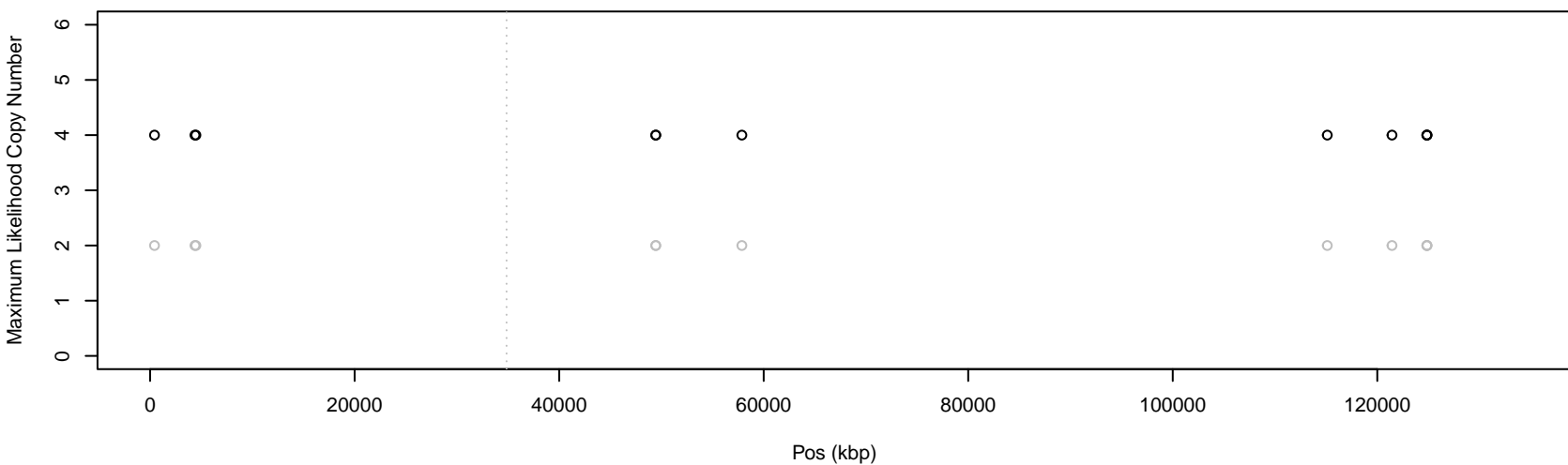
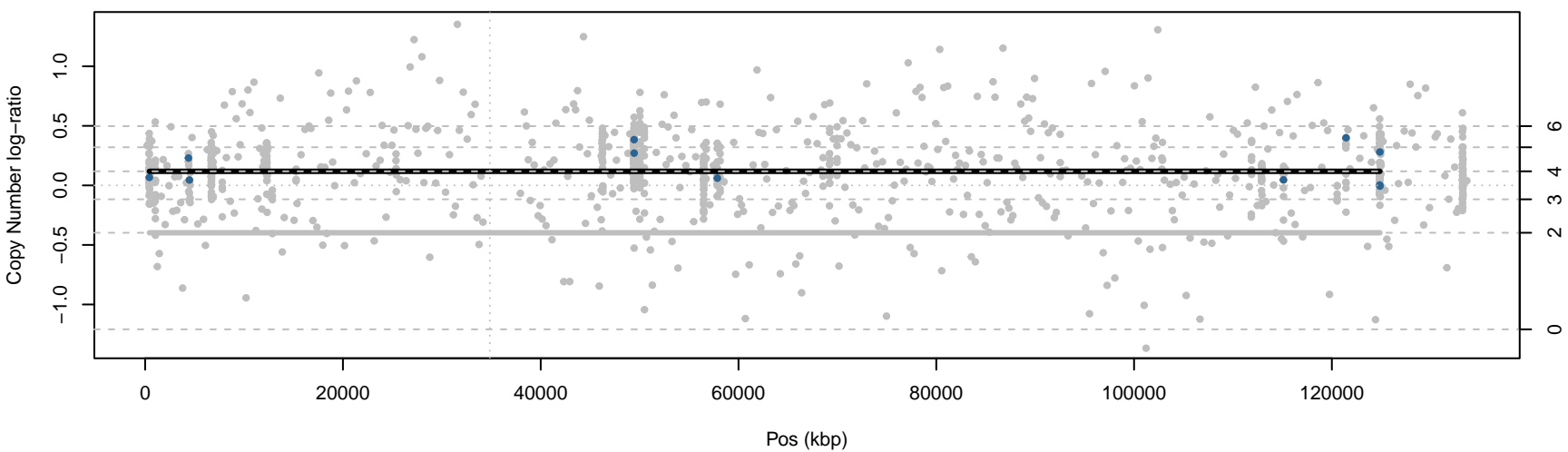
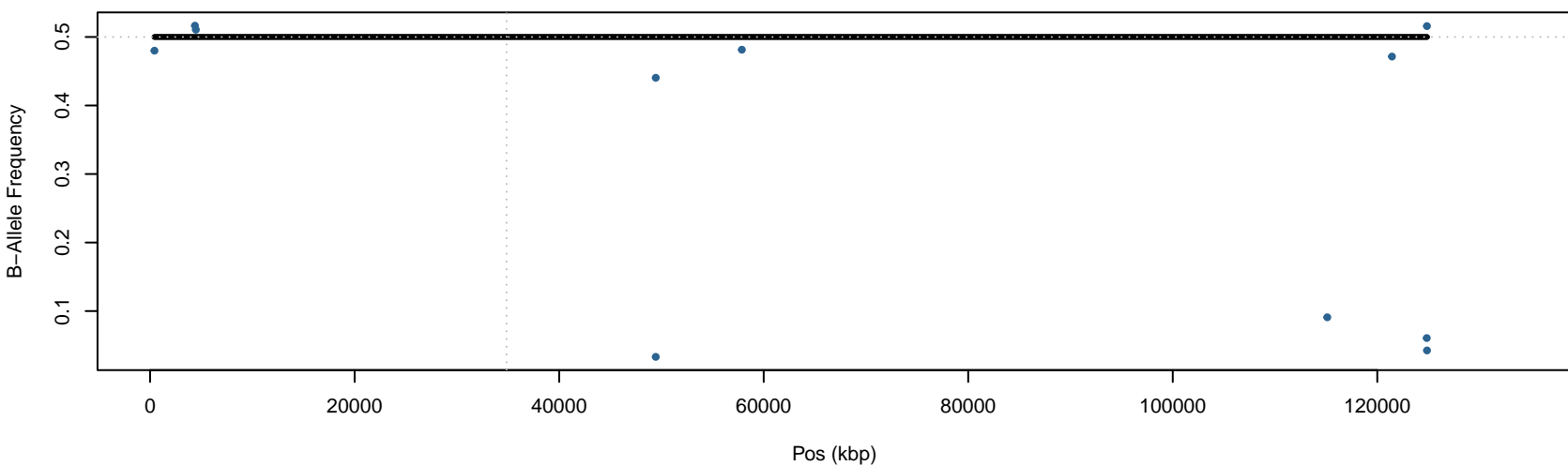
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 10



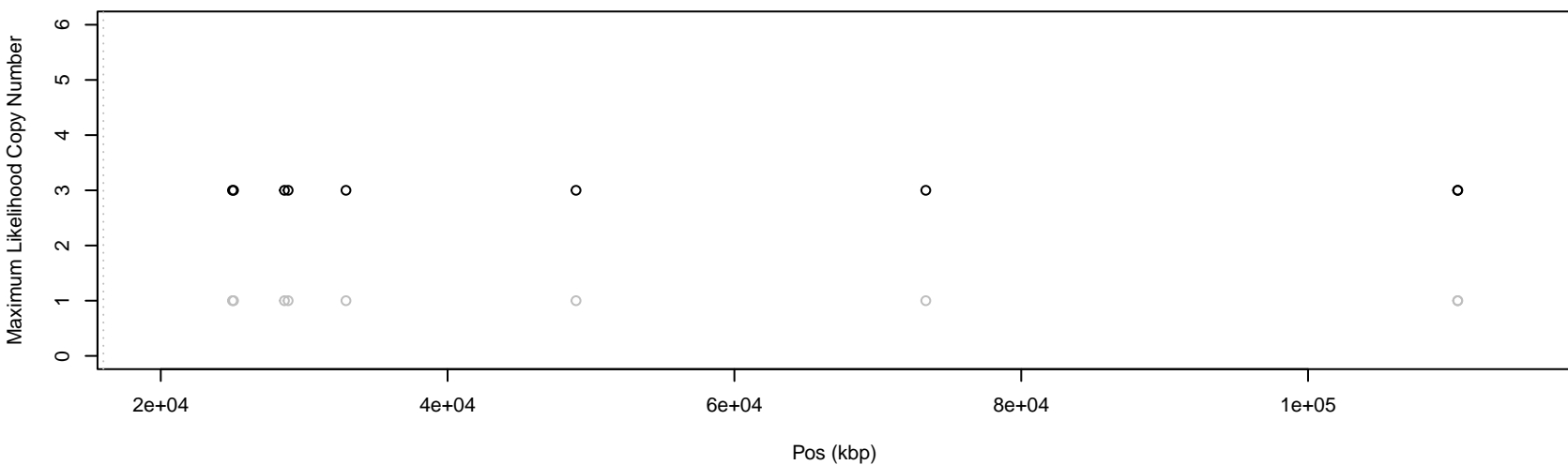
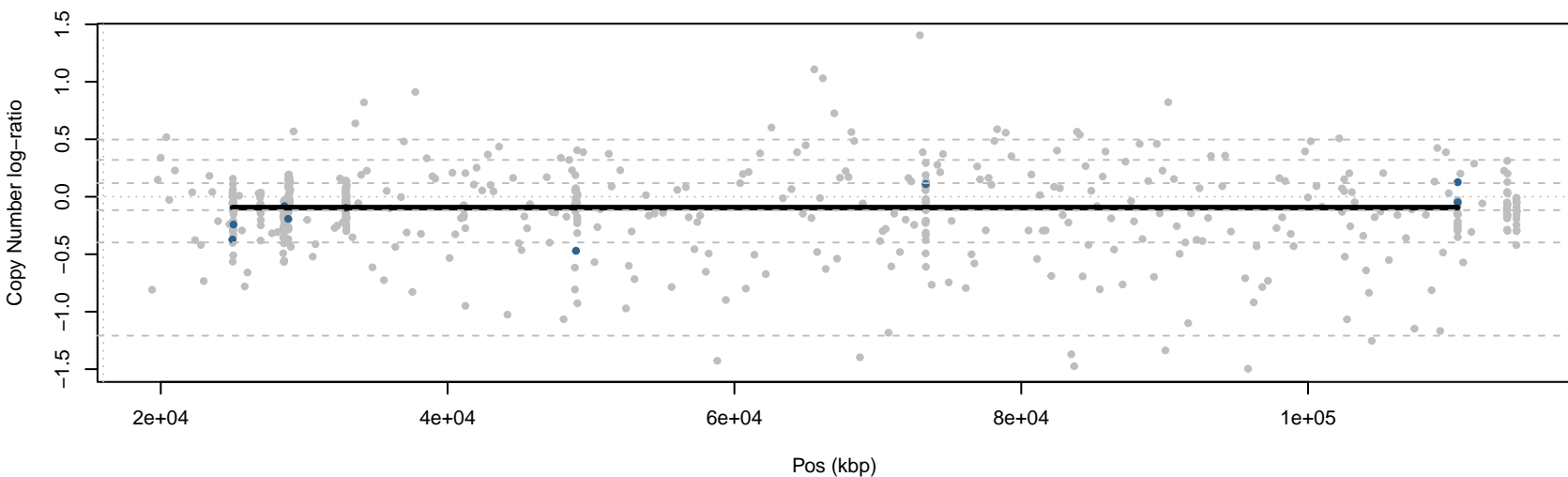
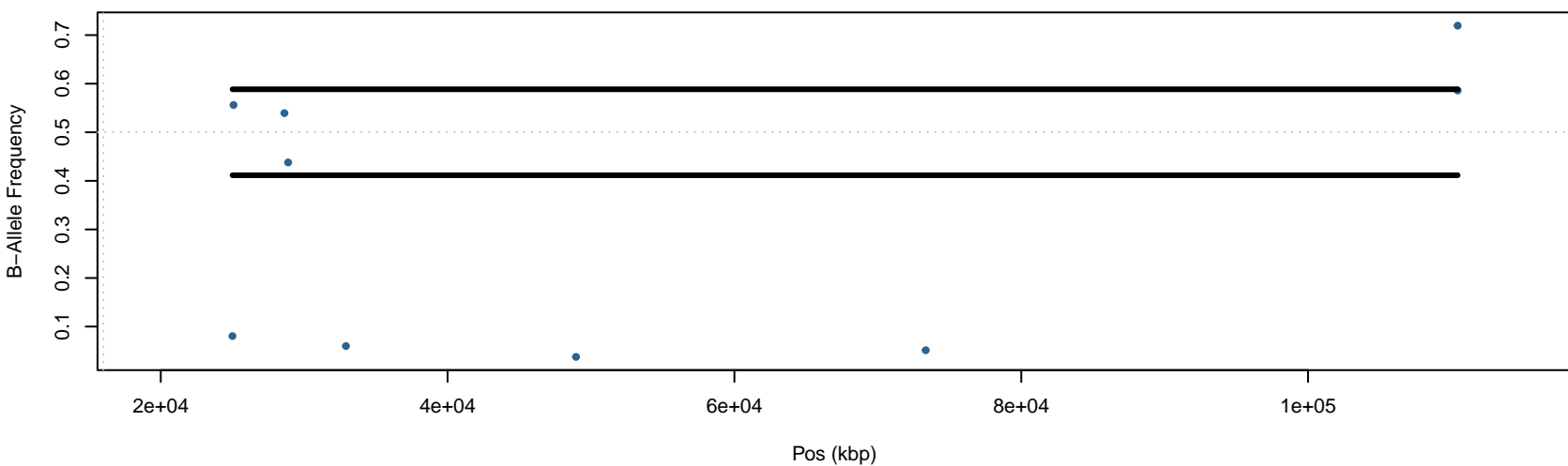
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 11



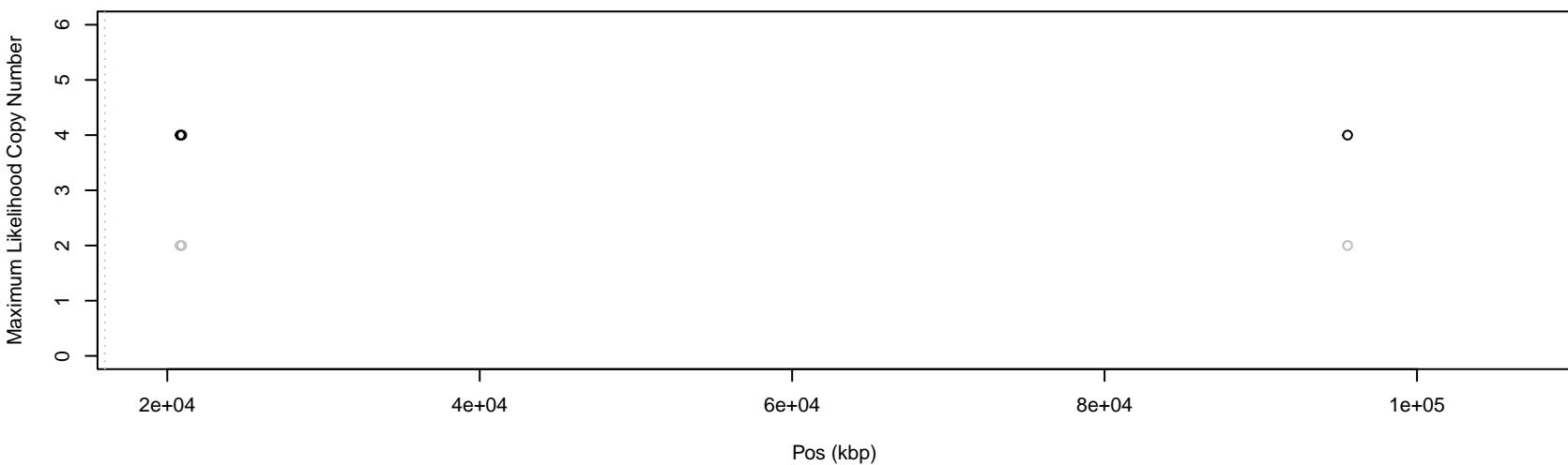
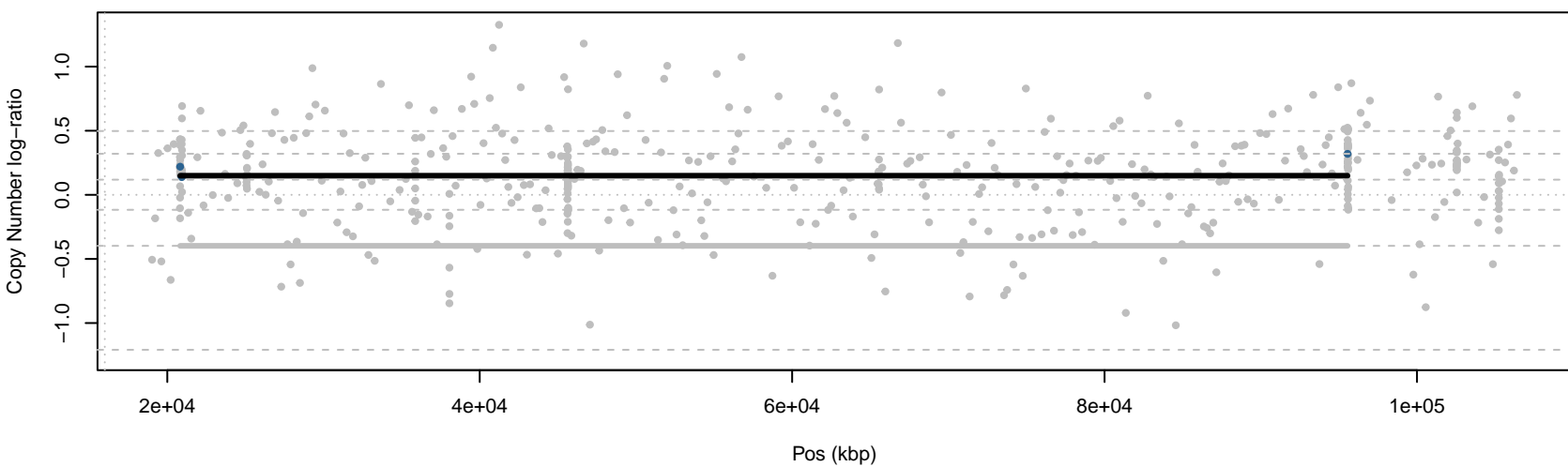
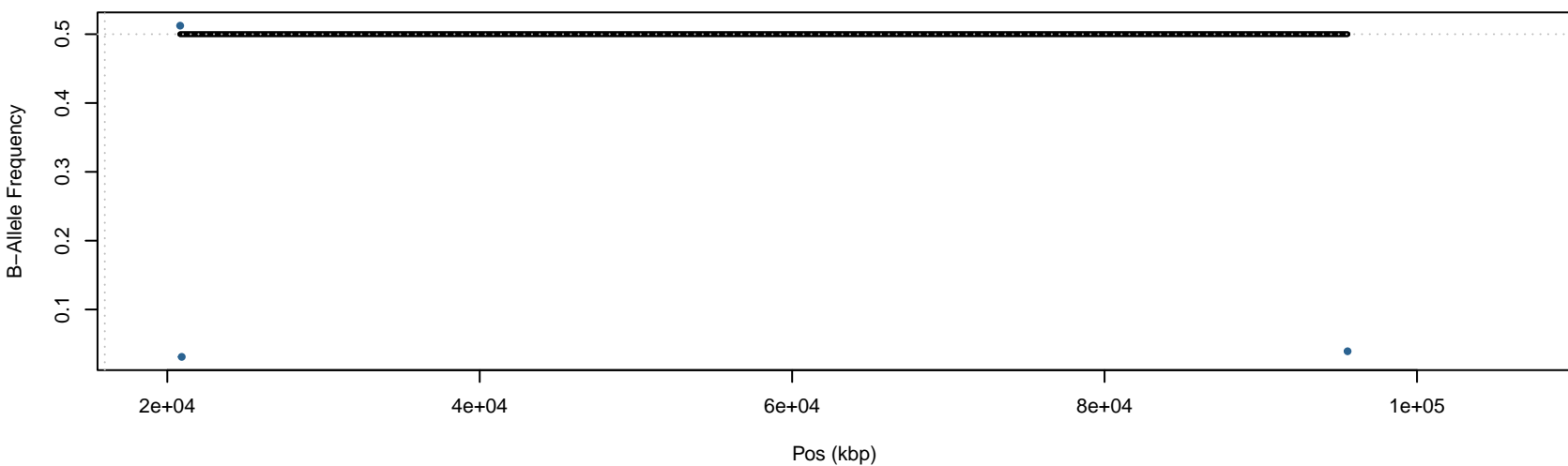
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 12



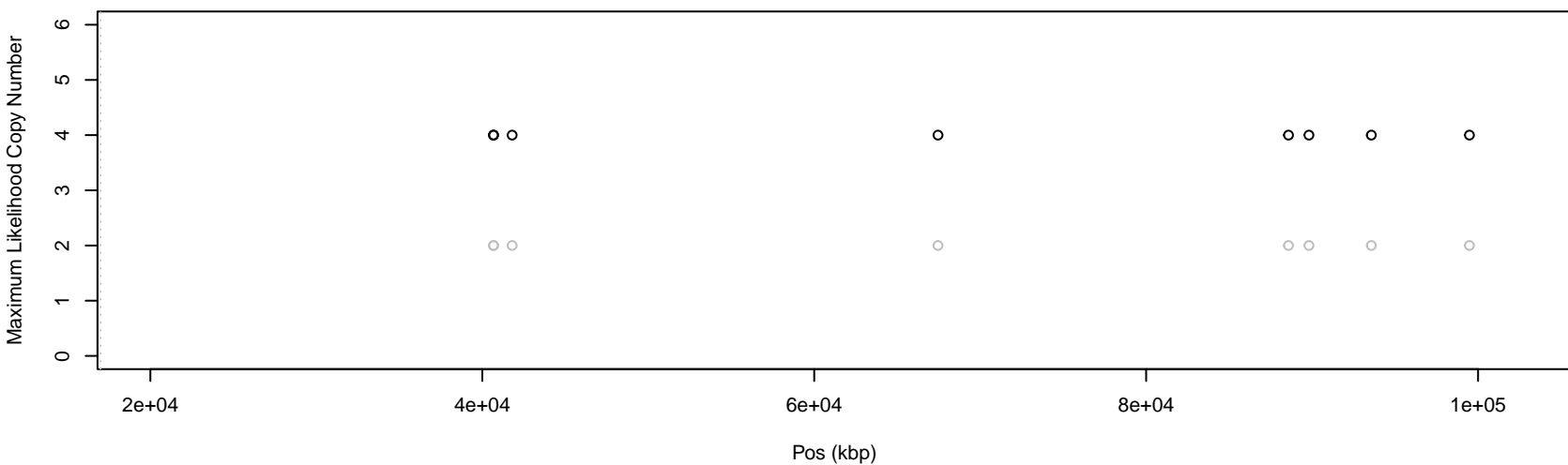
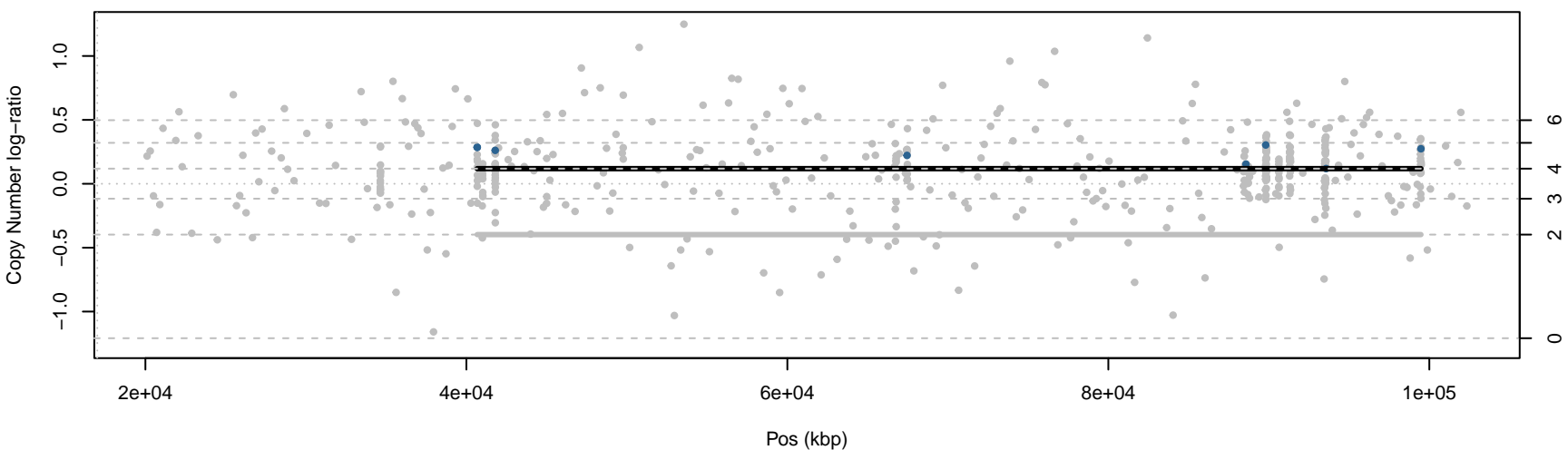
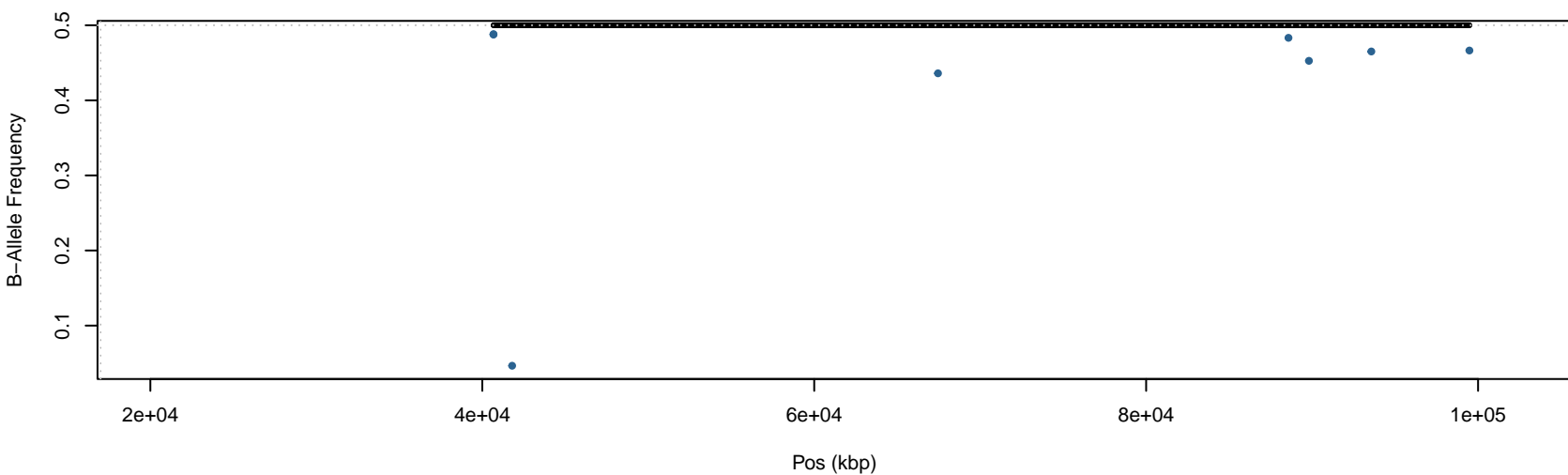
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 13



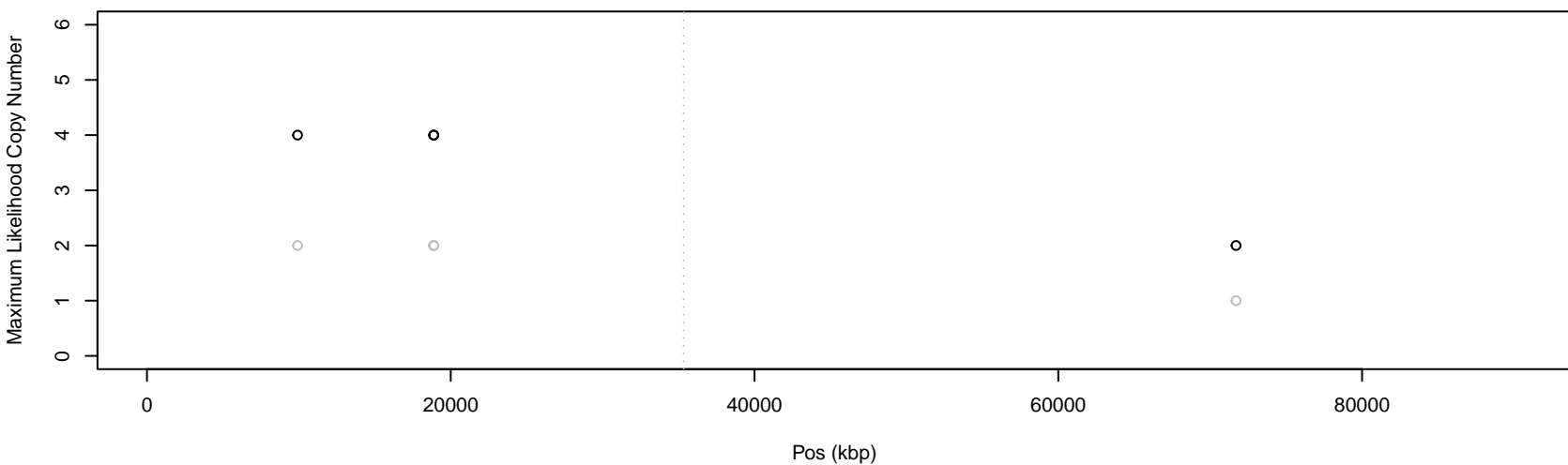
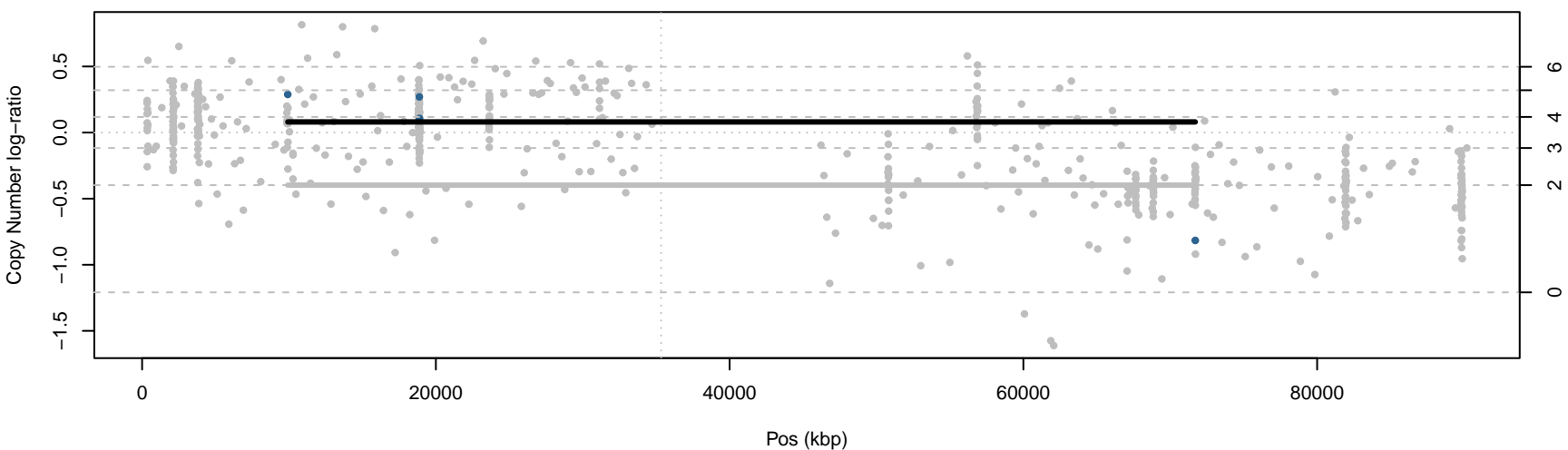
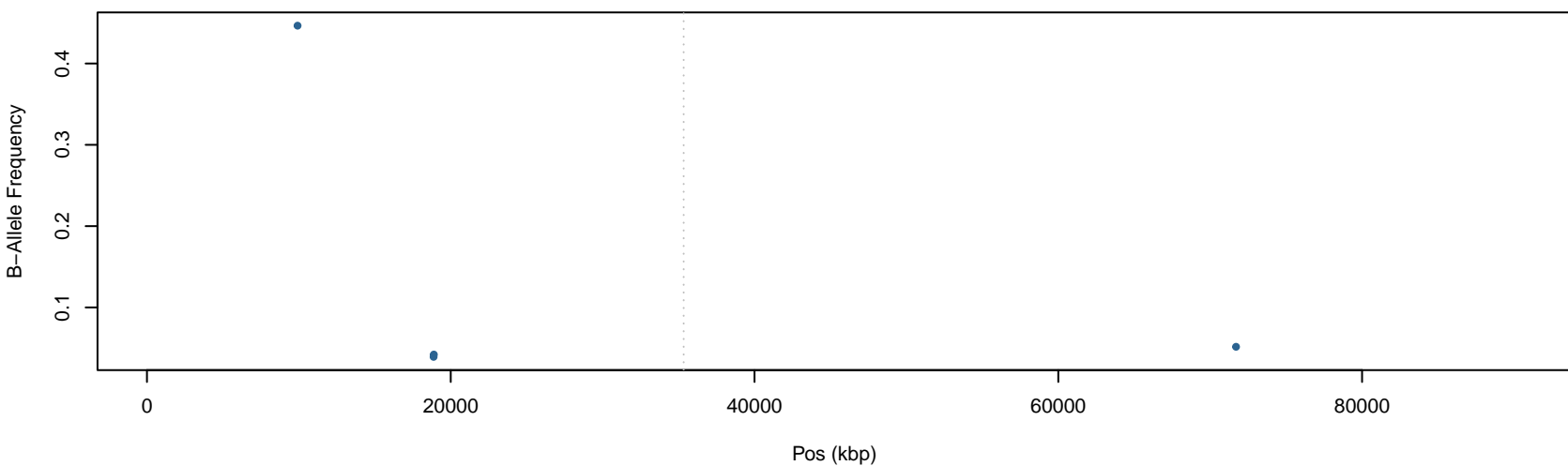
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 14



Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 15

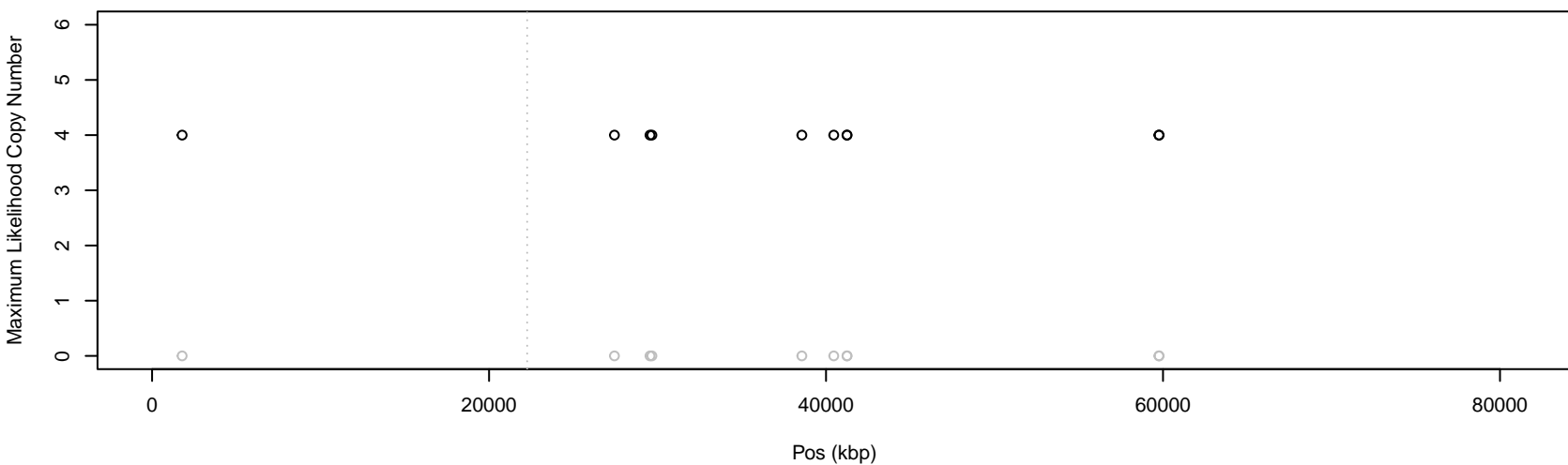
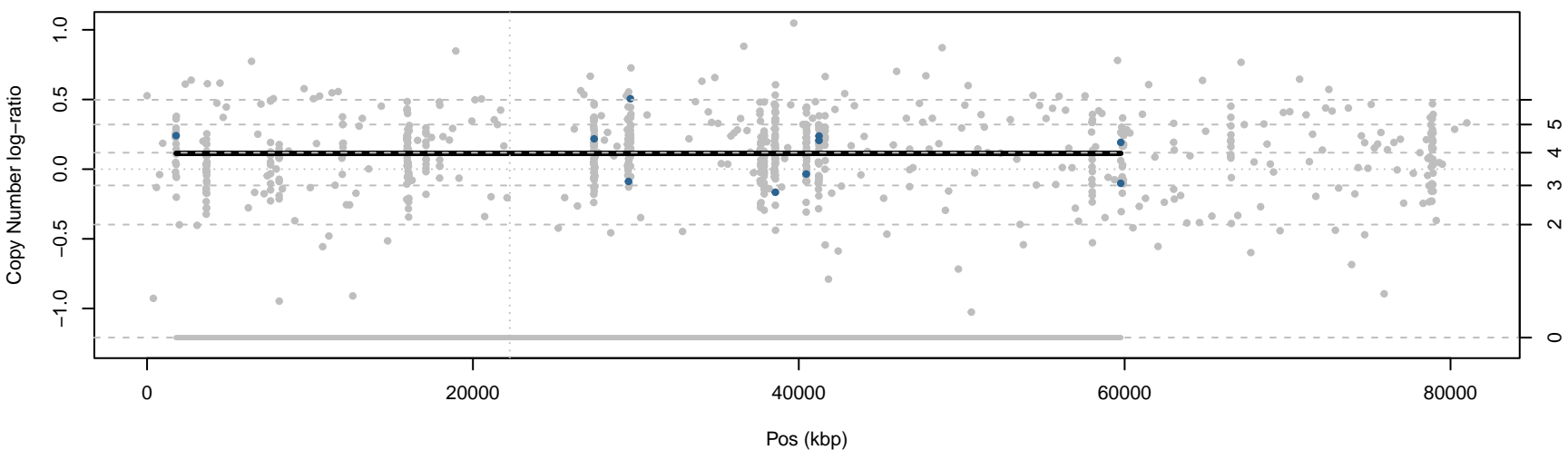
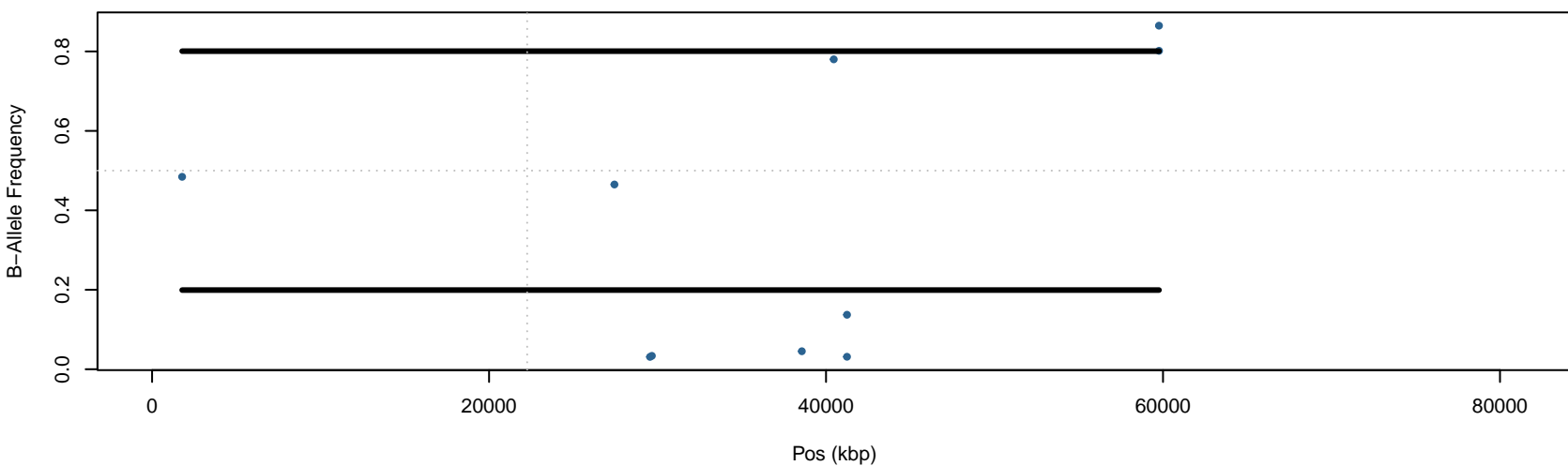


Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 16

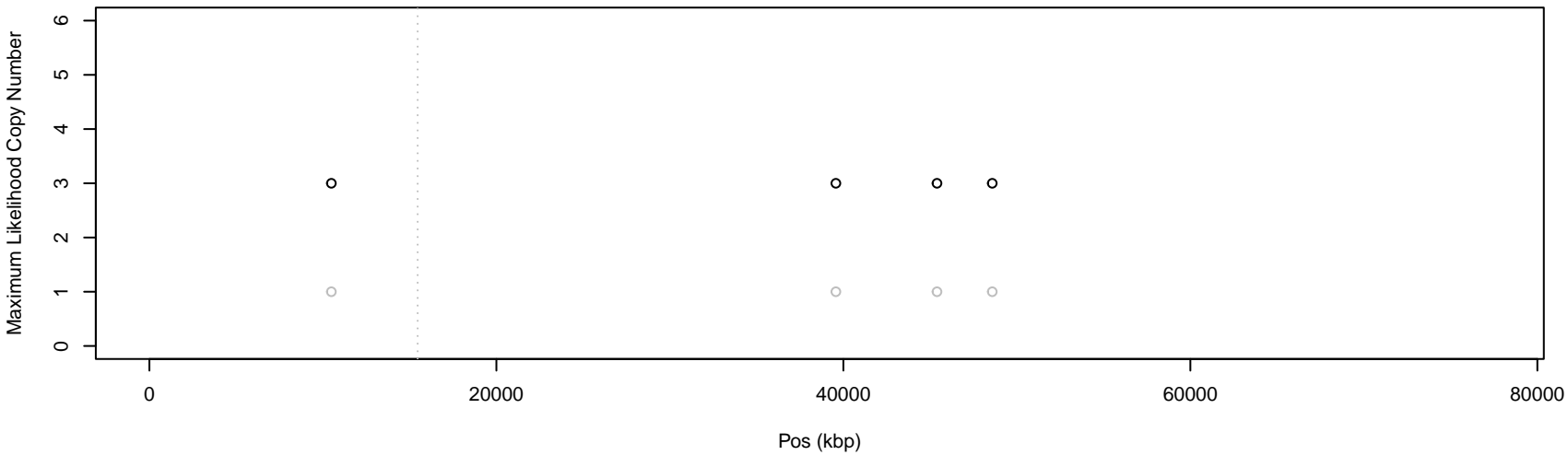
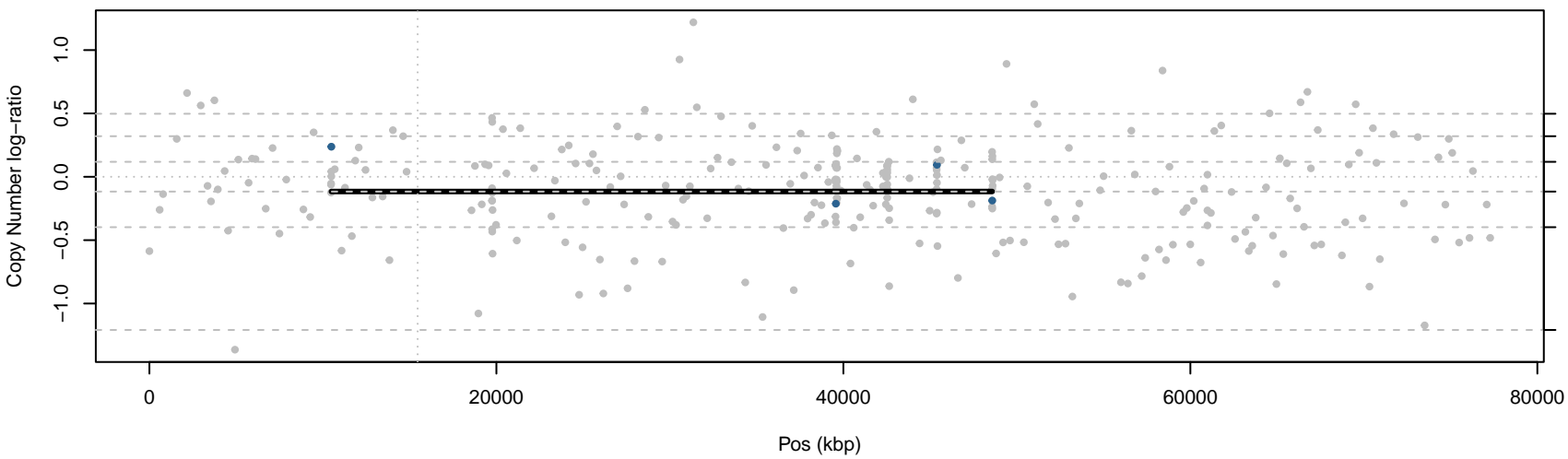
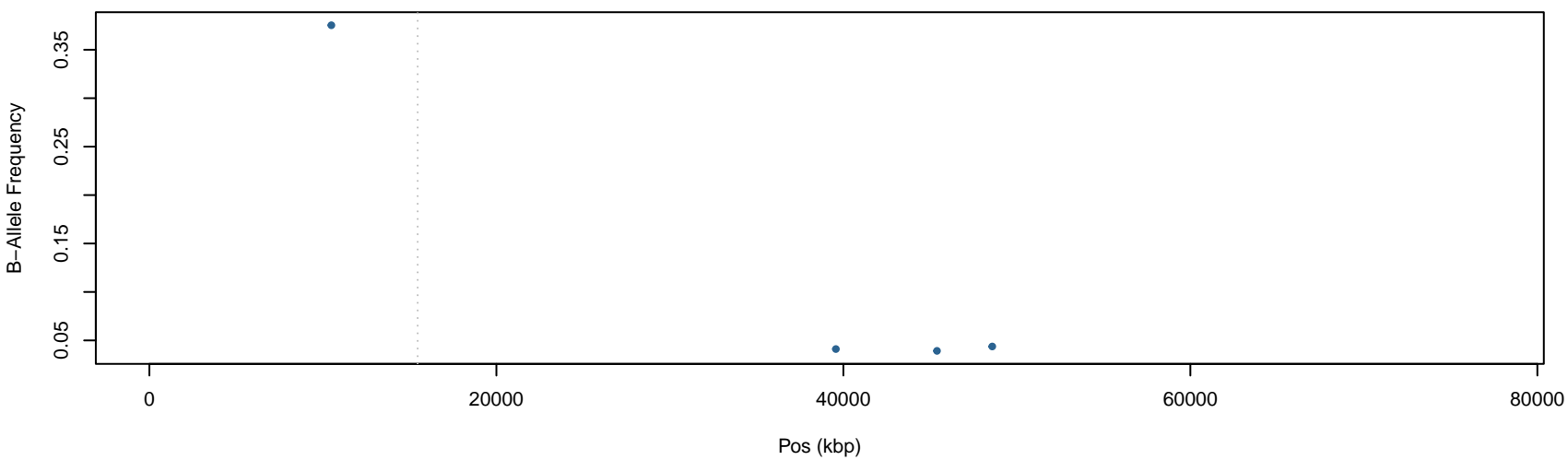




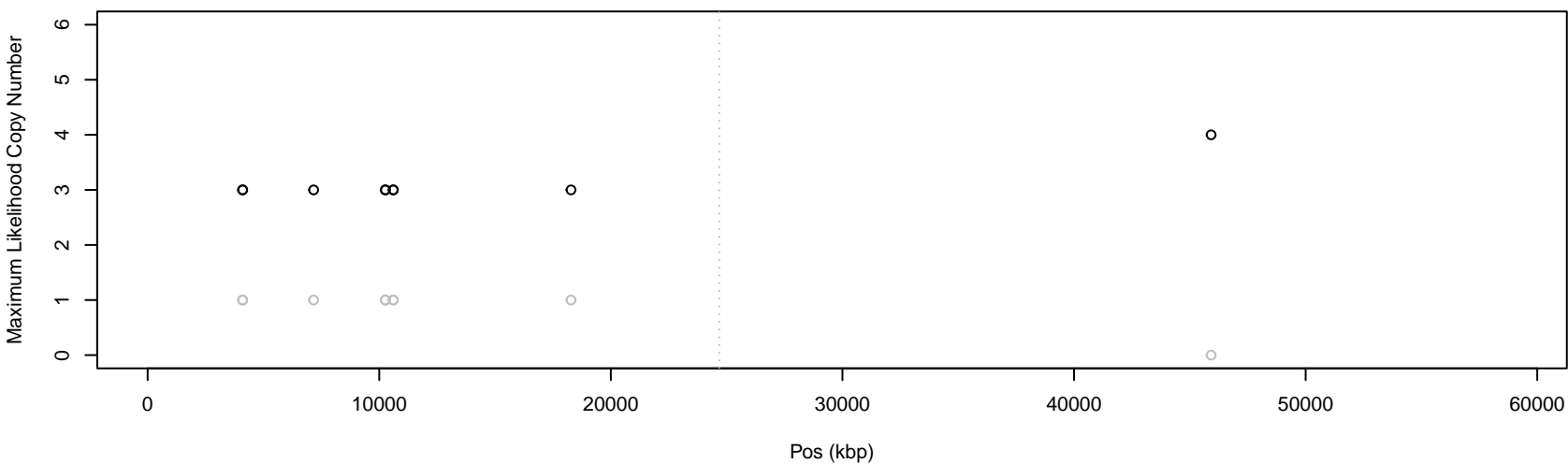
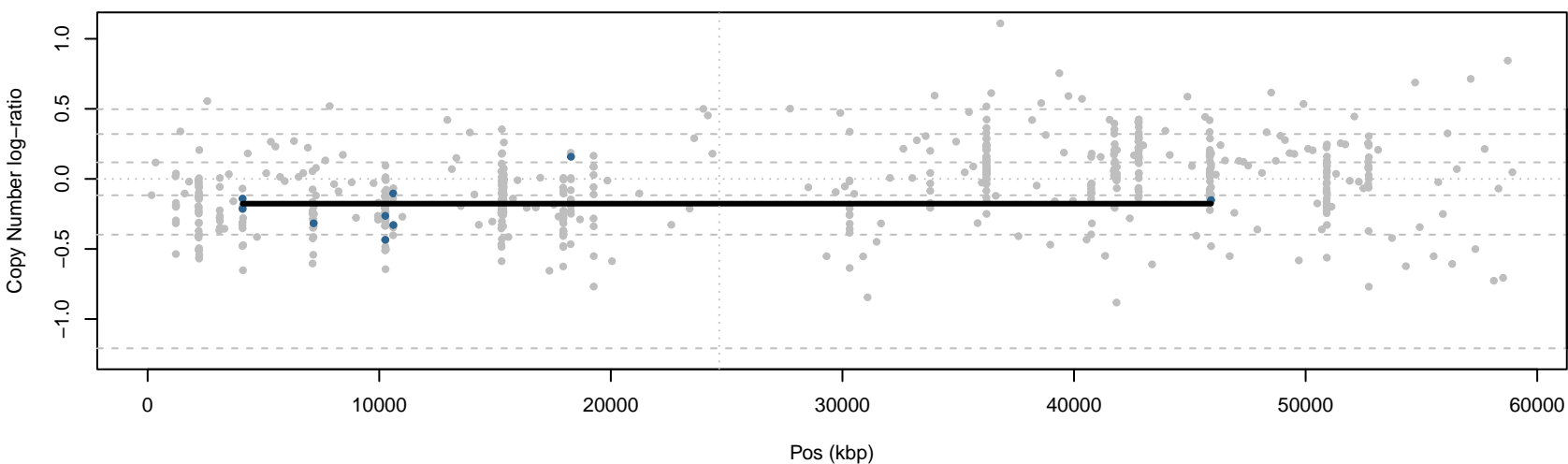
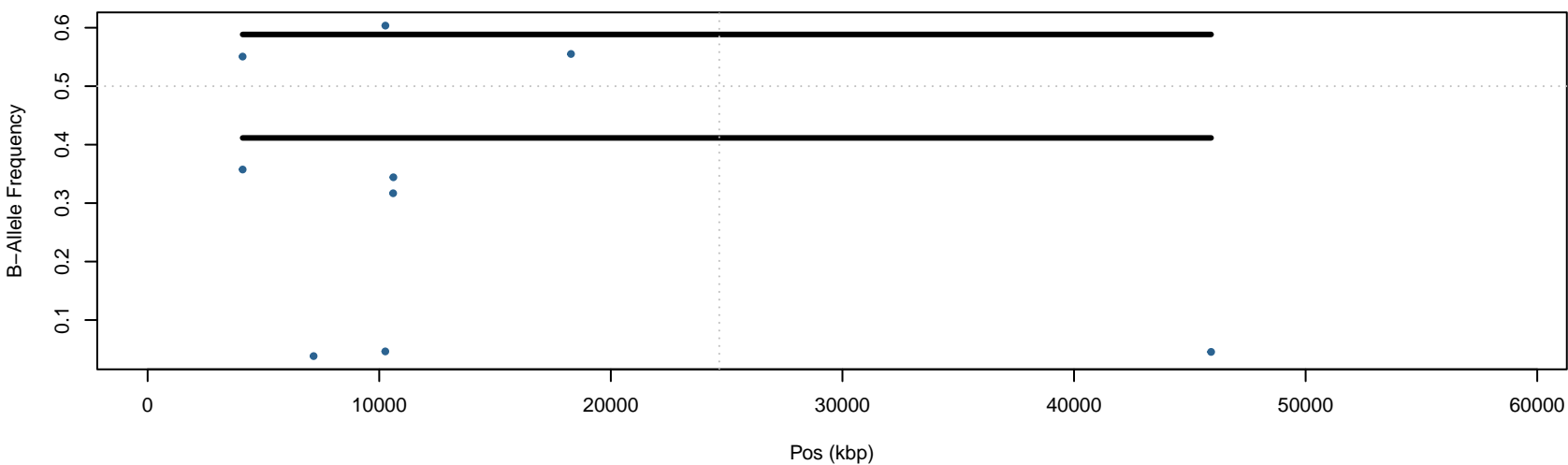
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 17



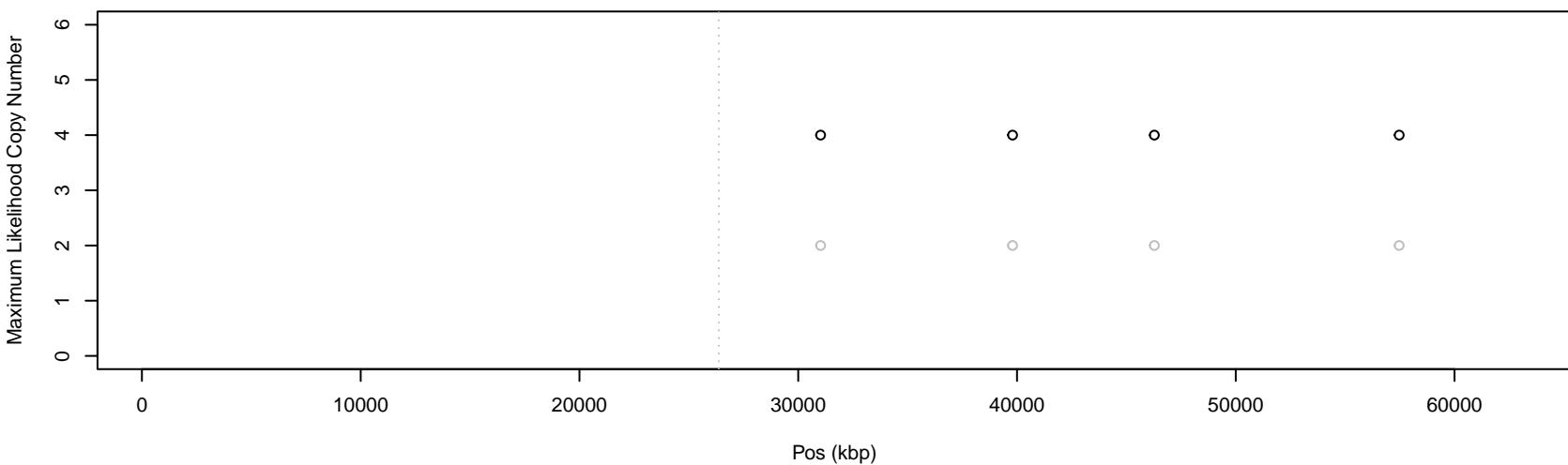
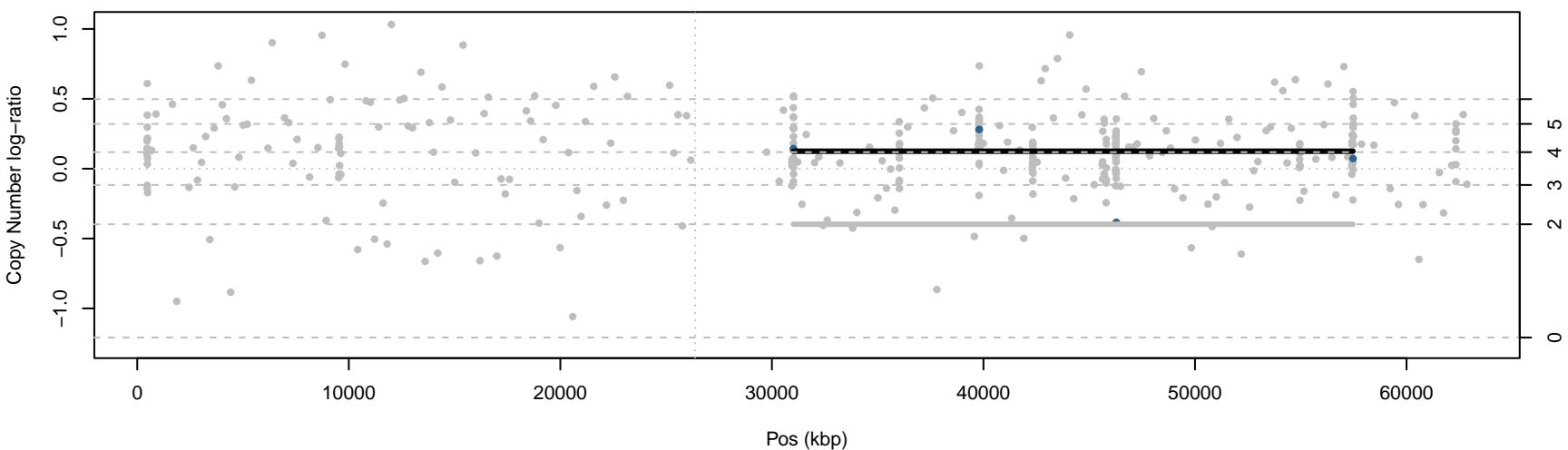
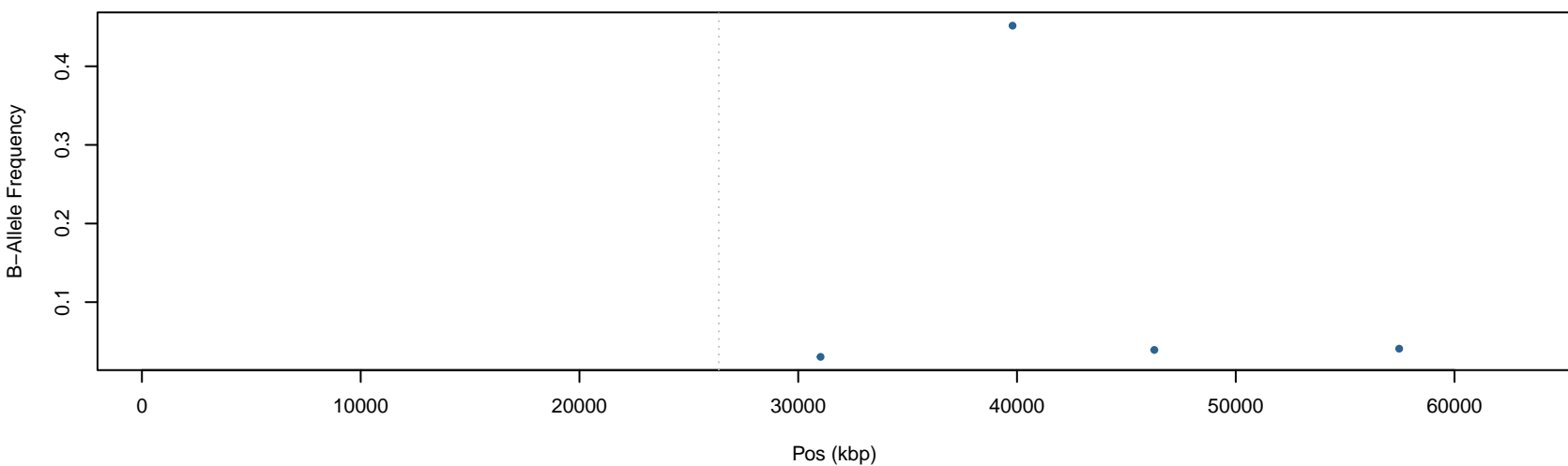
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 18



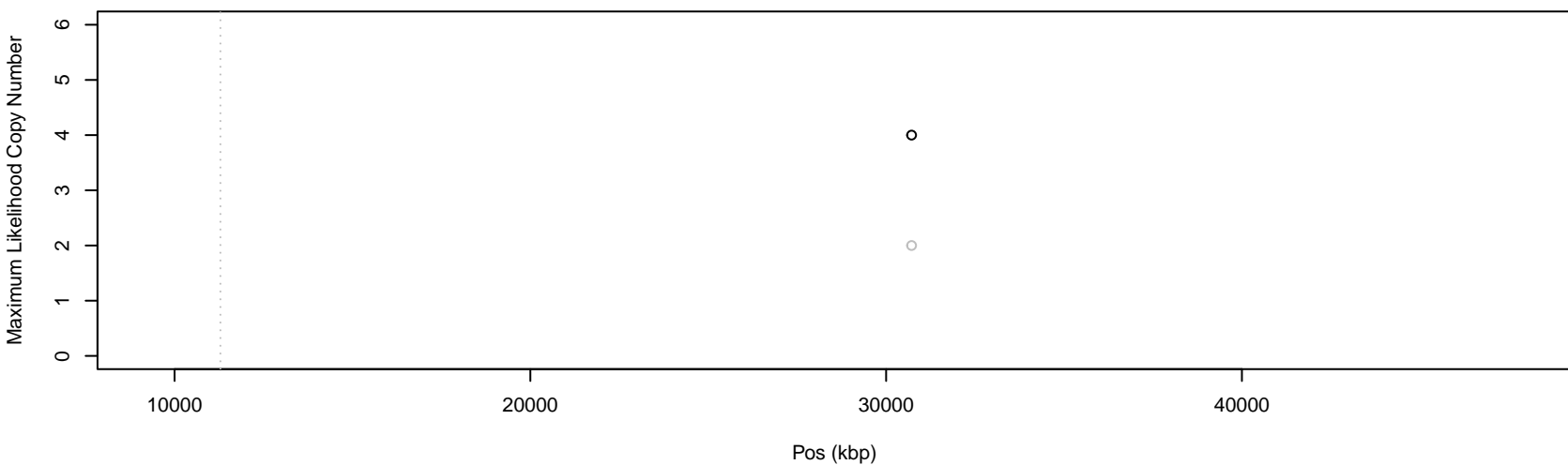
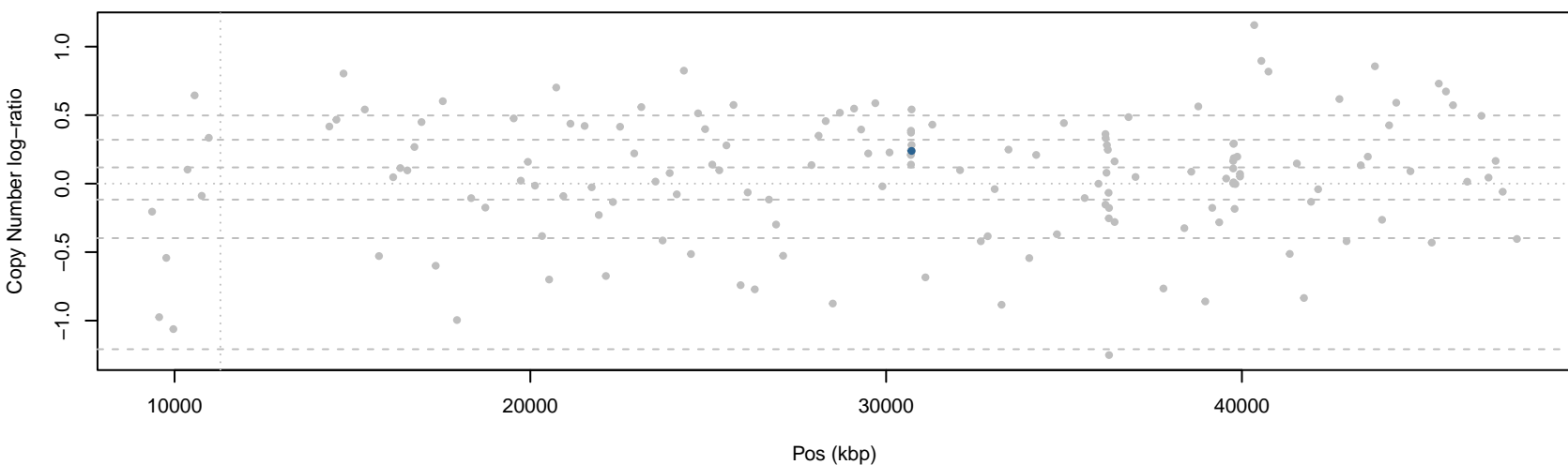
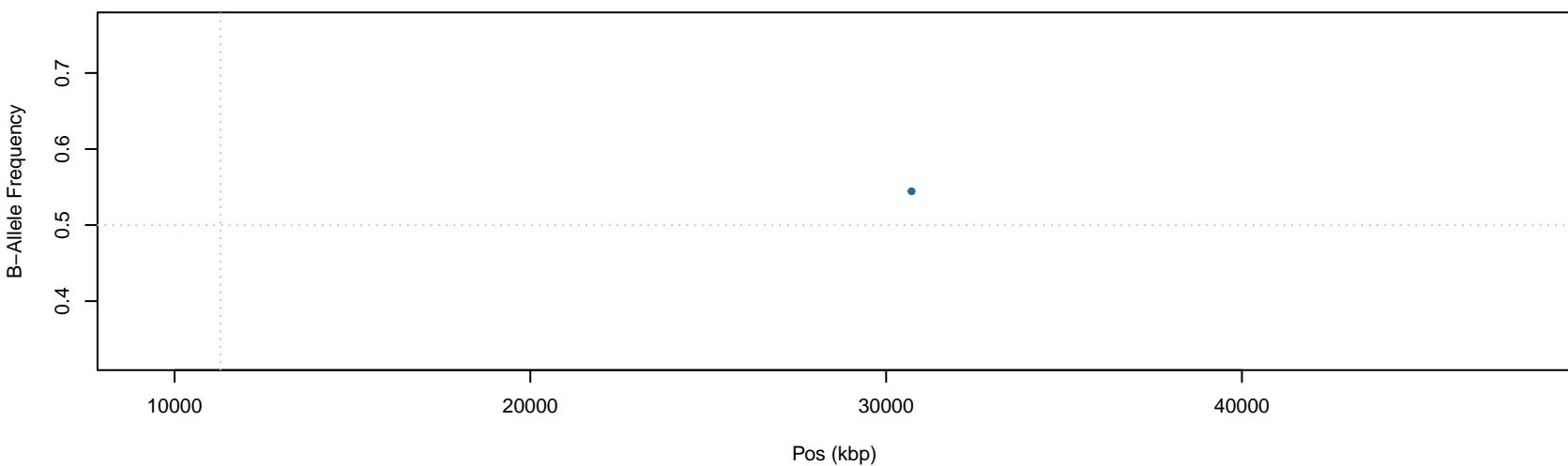
Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 19



Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 20



Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 21



Purity: 0.43 Tumor ploidy: 3.477 SNV log-likelihood: -76.1 GoF: 81.7% Mean coverage: 299;387 Chromosome: 22

