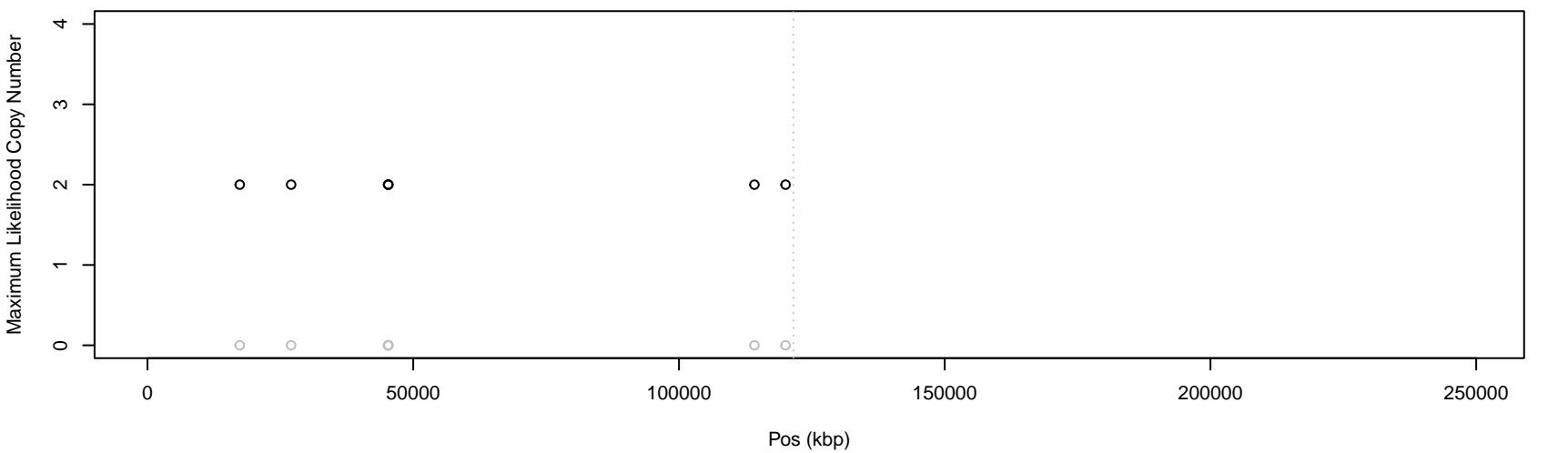
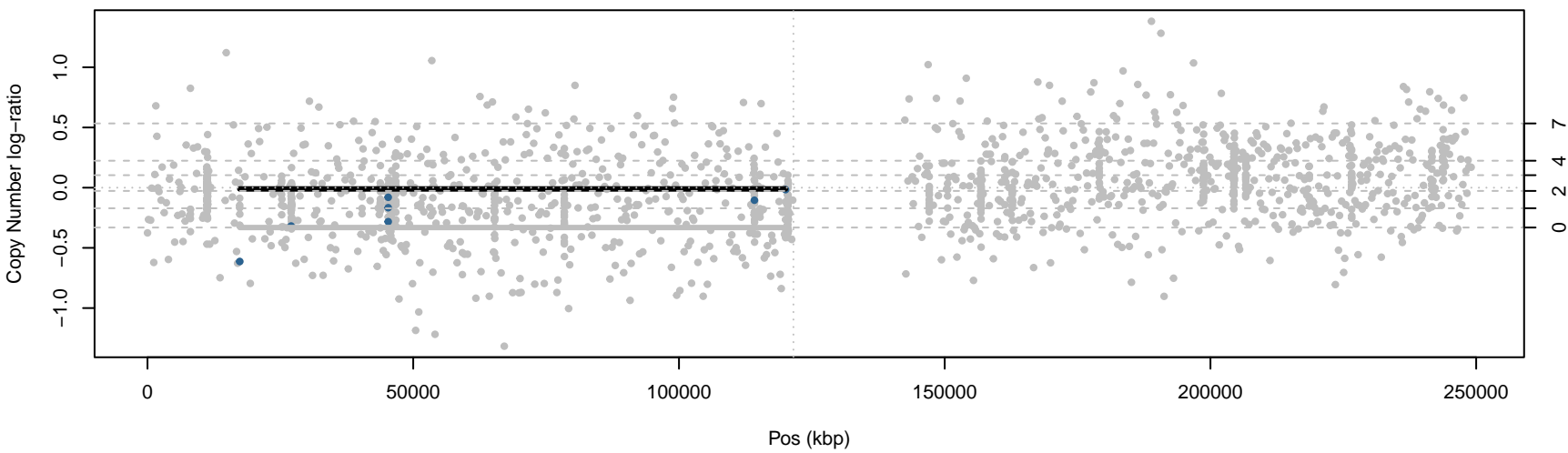
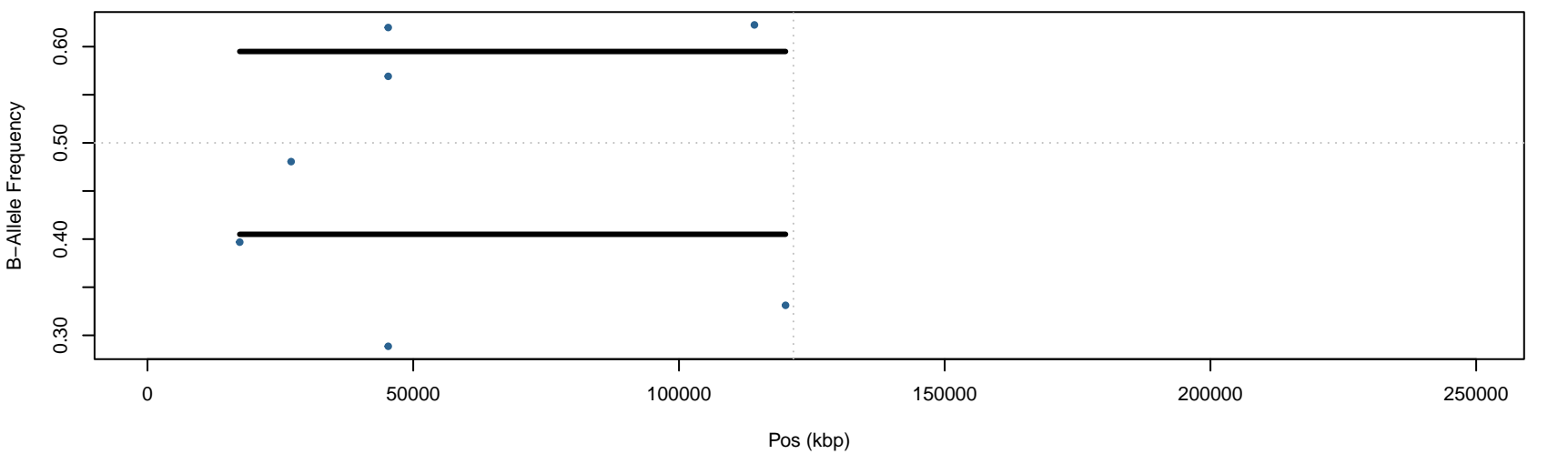
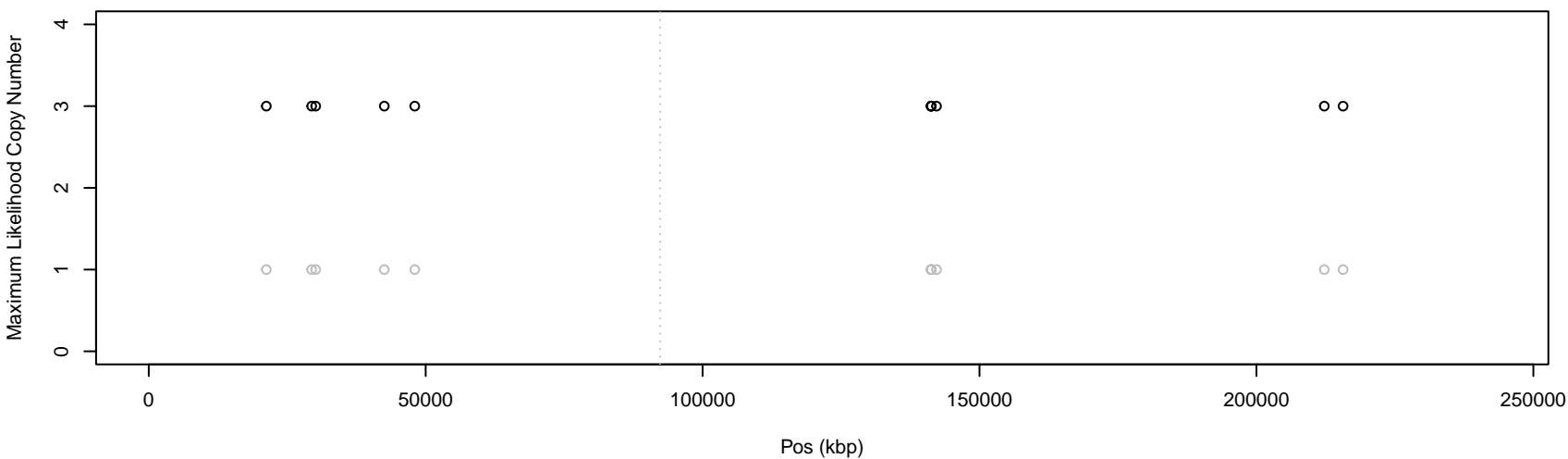
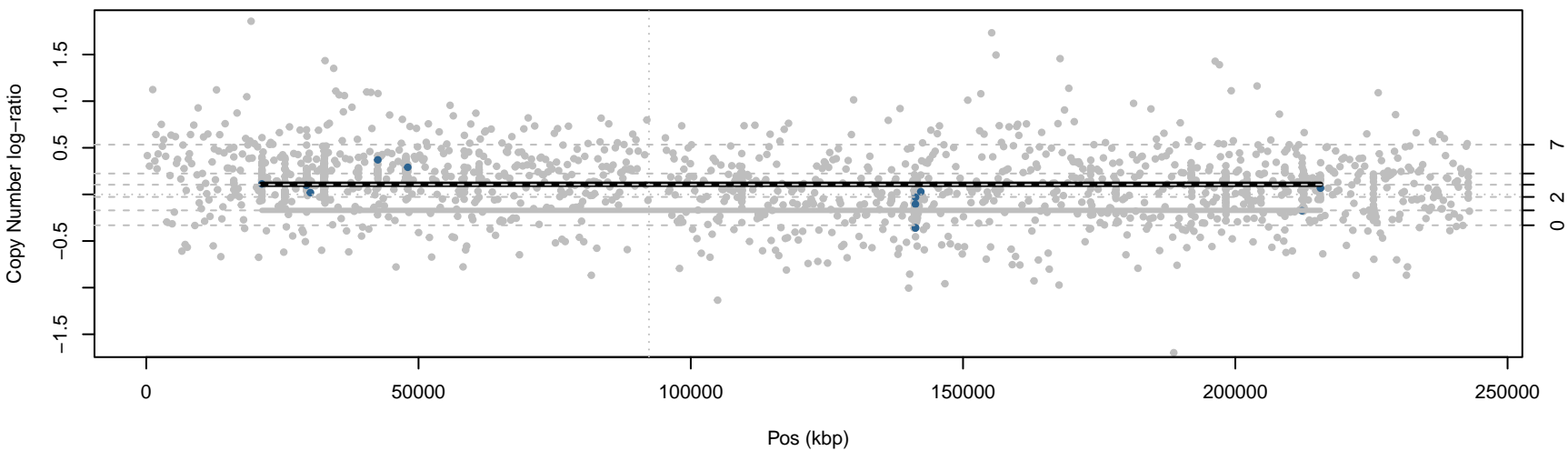
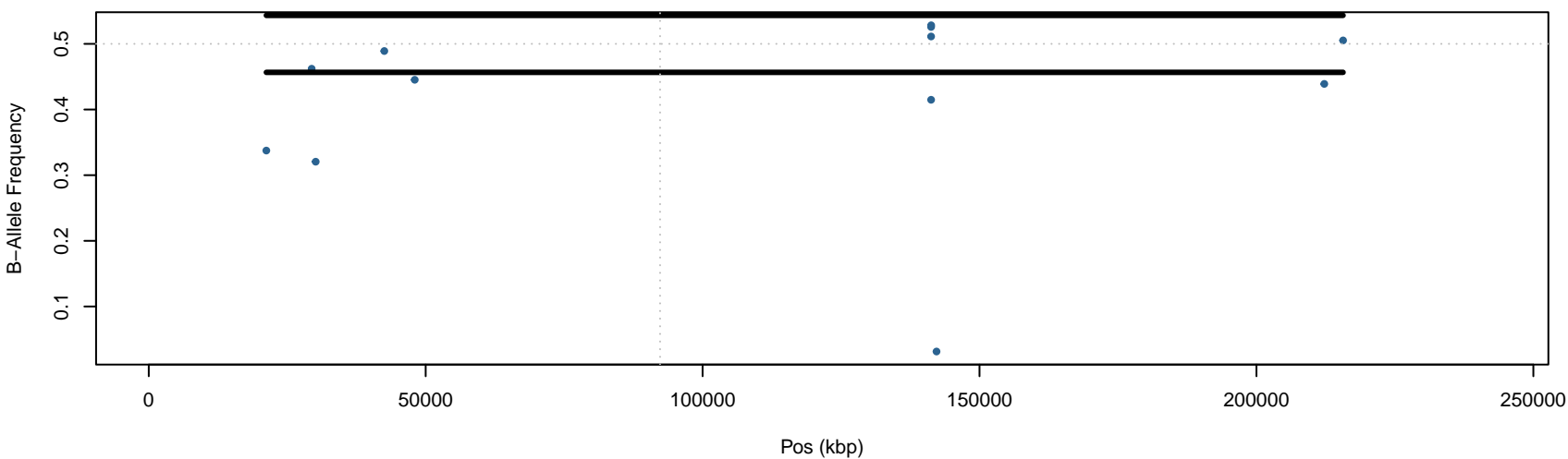


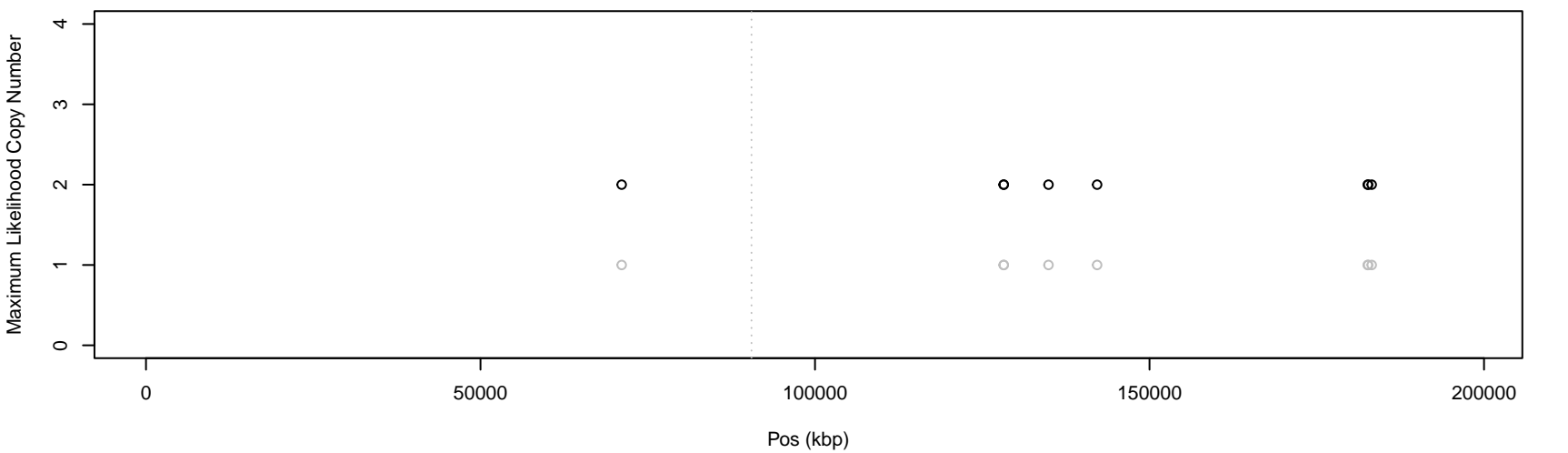
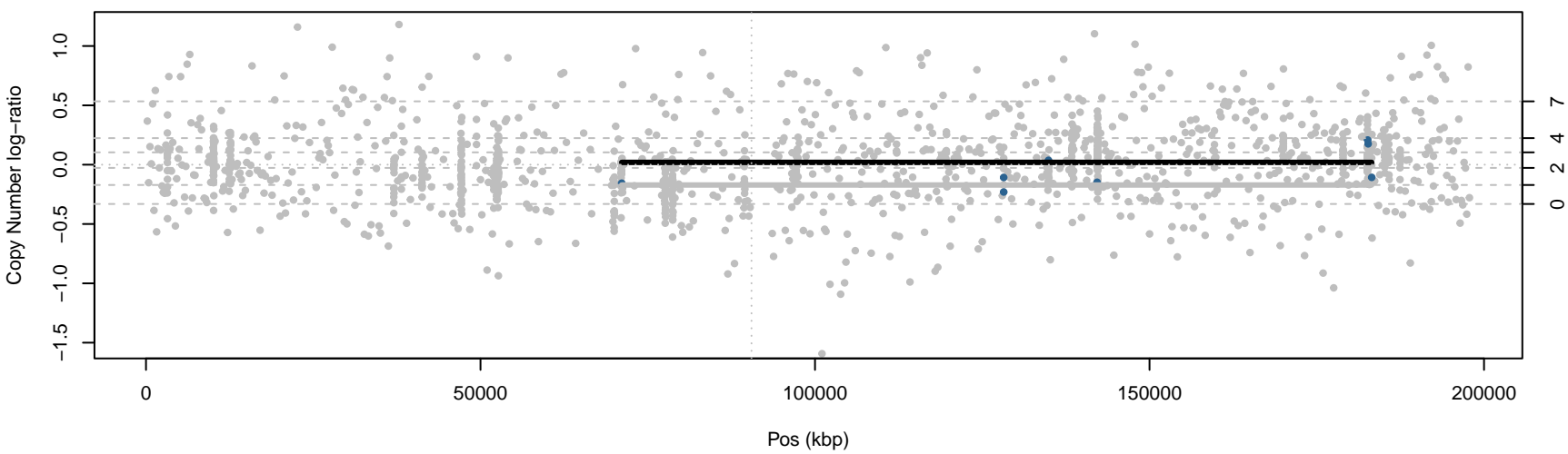
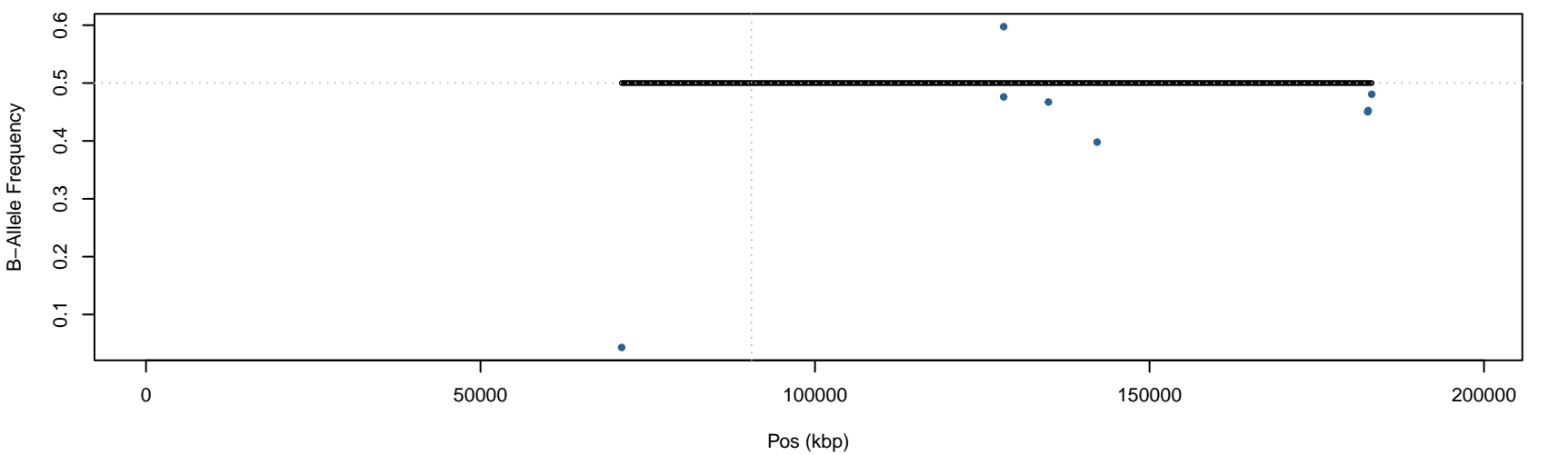
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 1



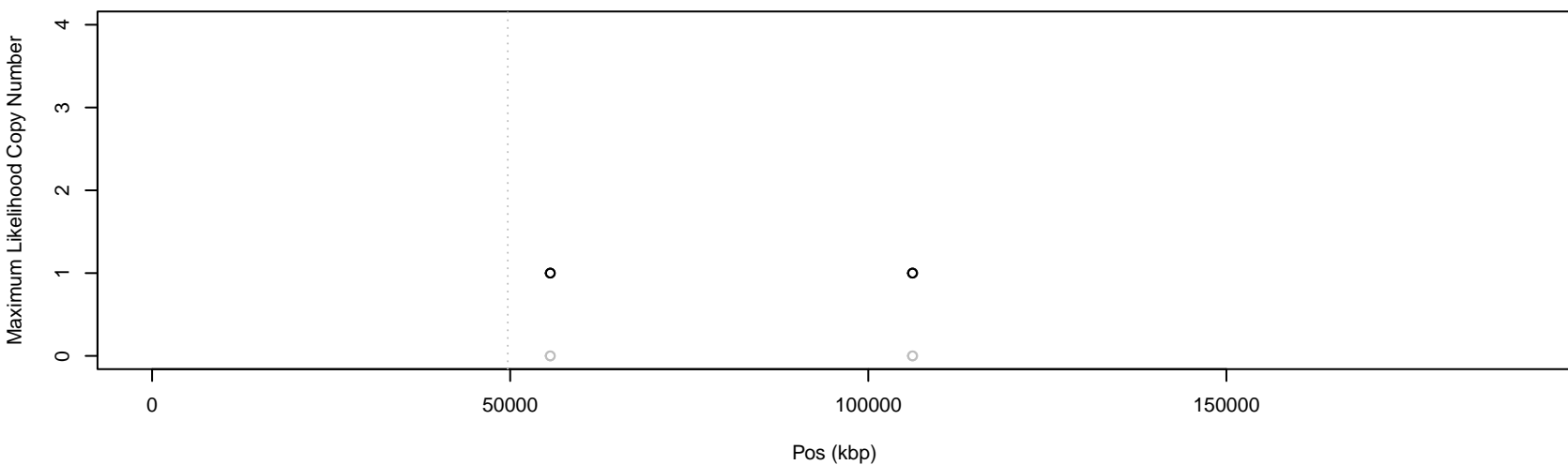
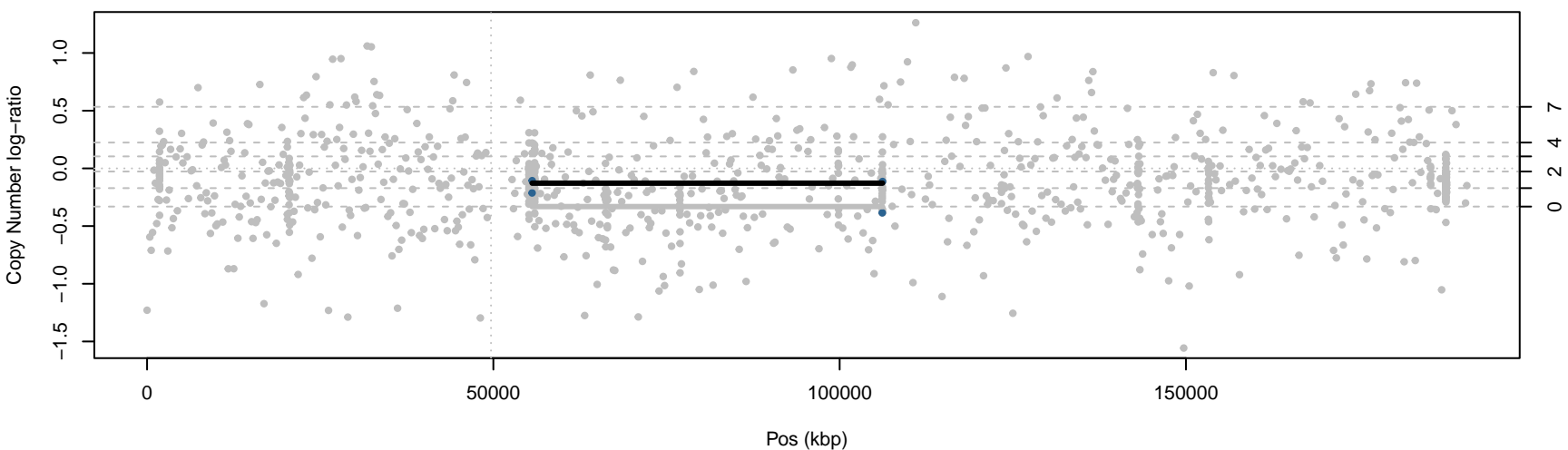
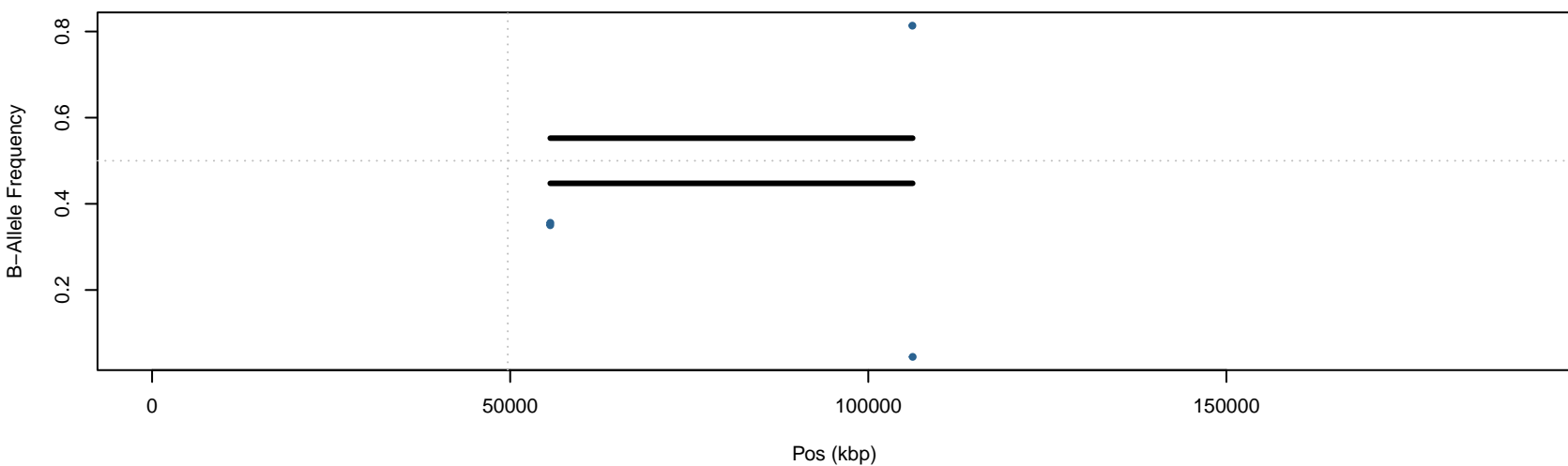
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 2



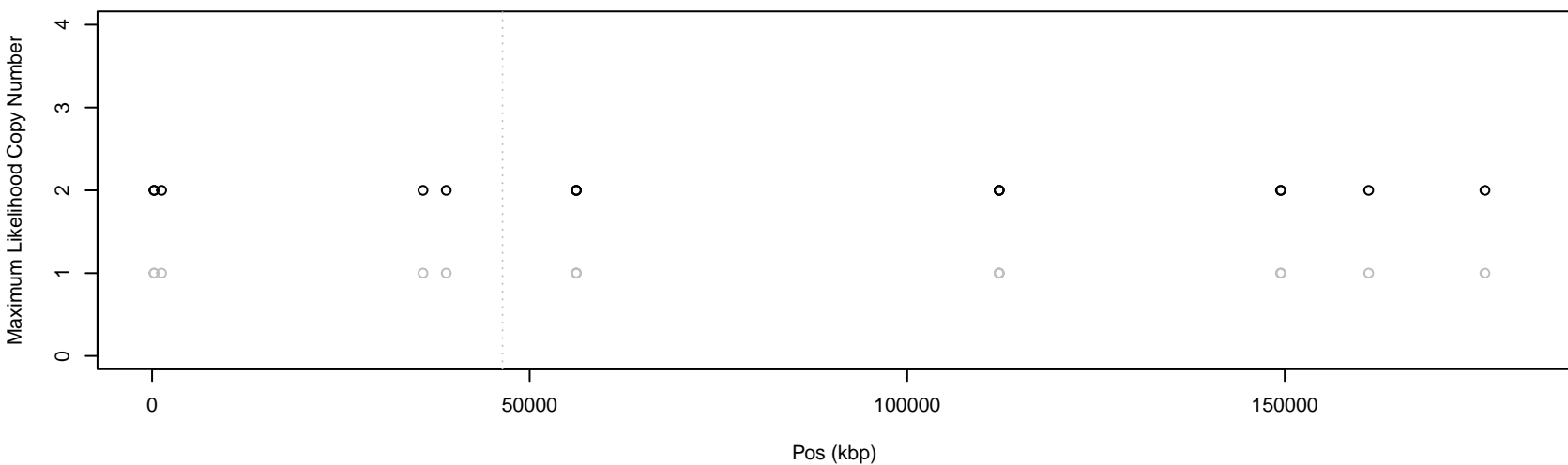
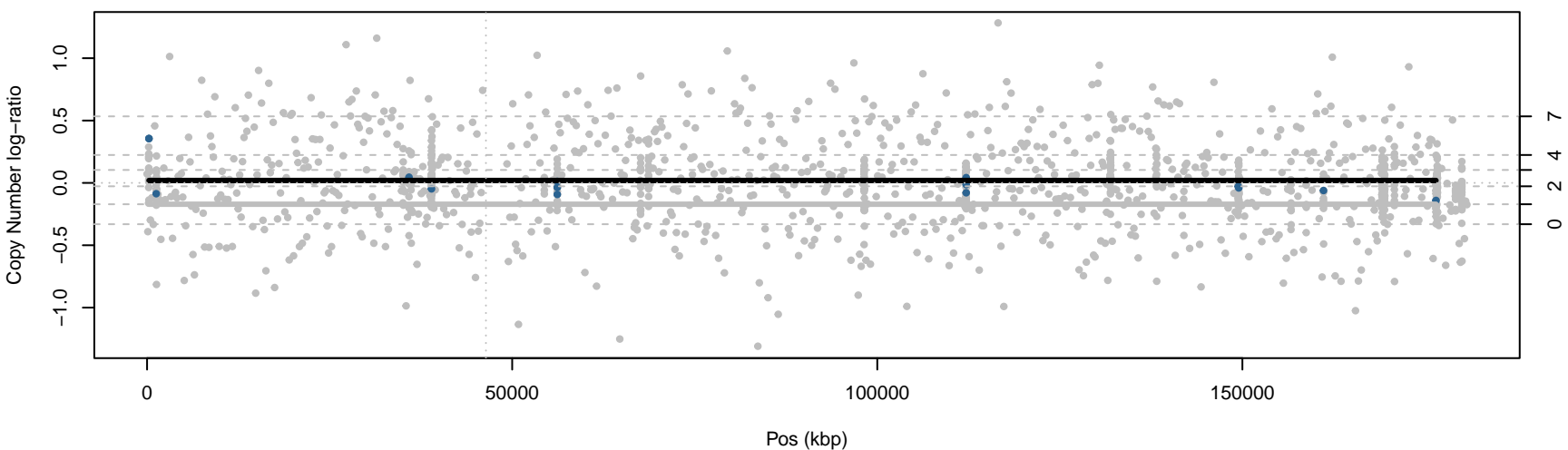
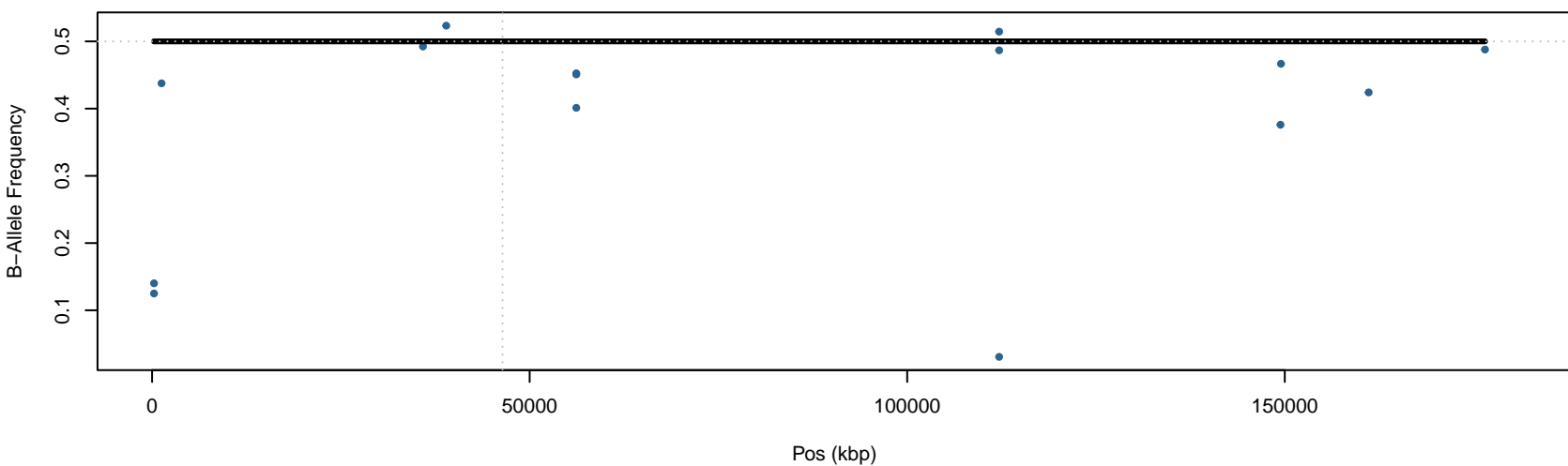
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 3



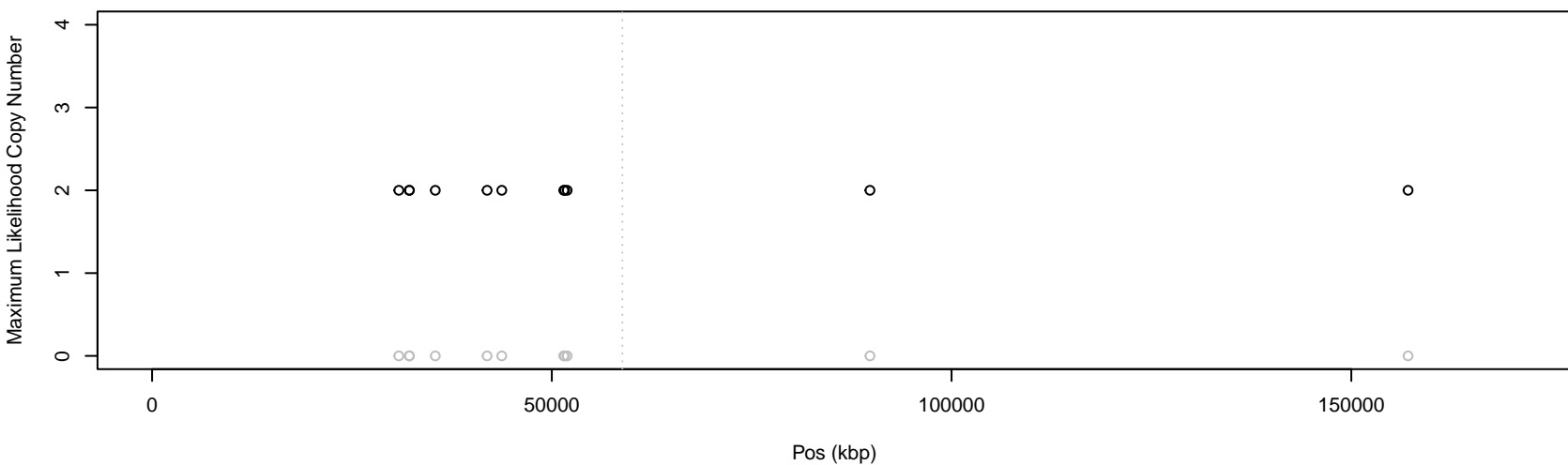
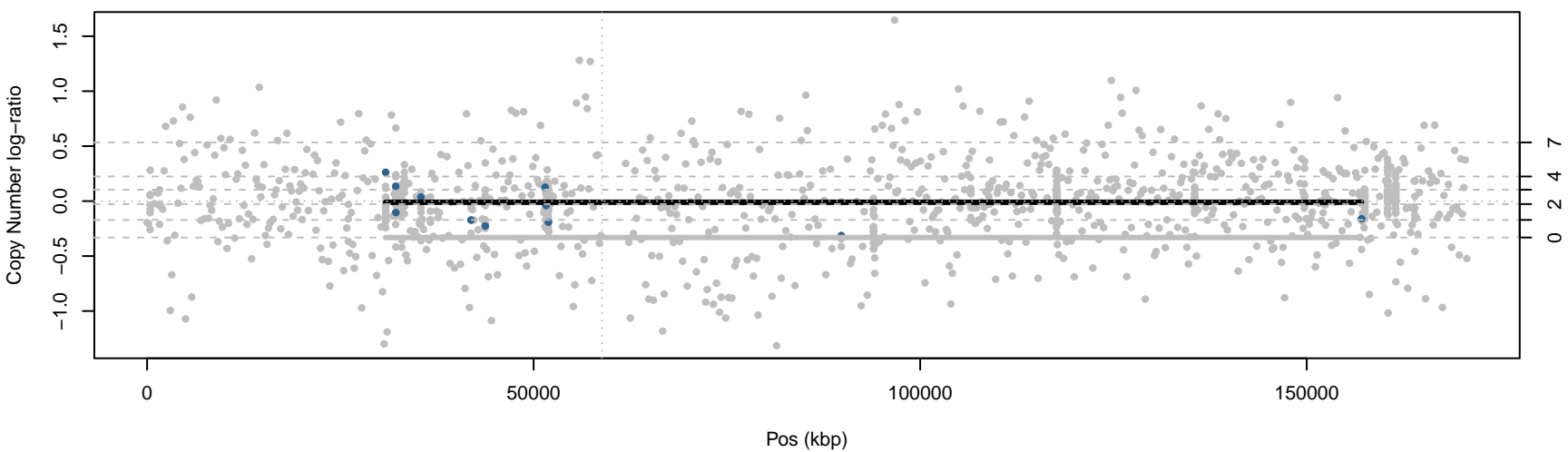
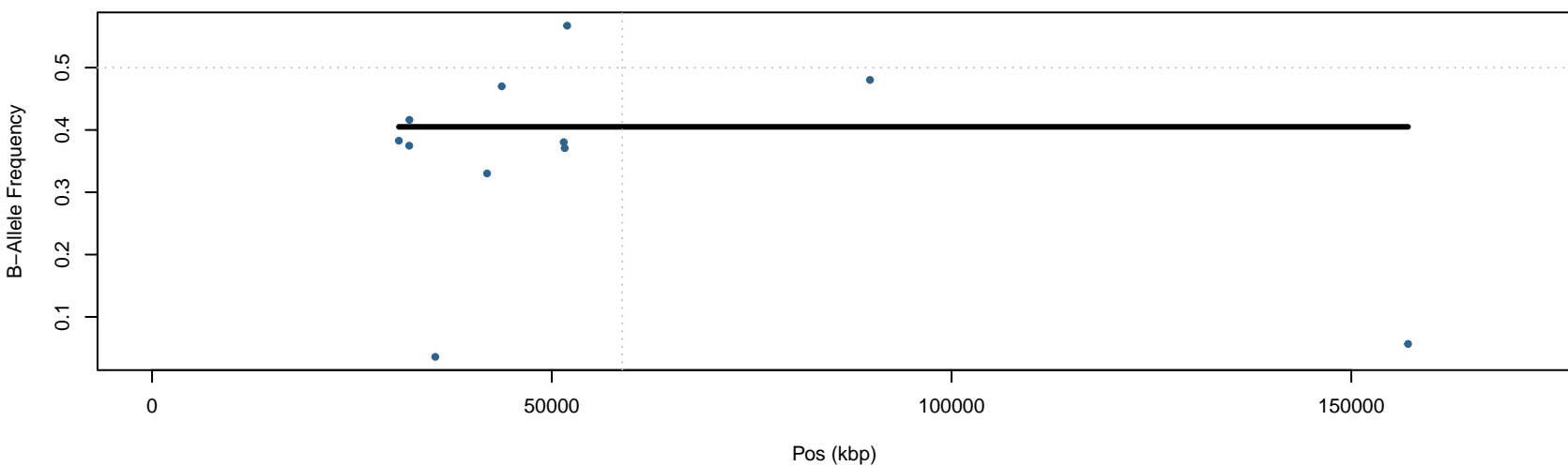
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 4



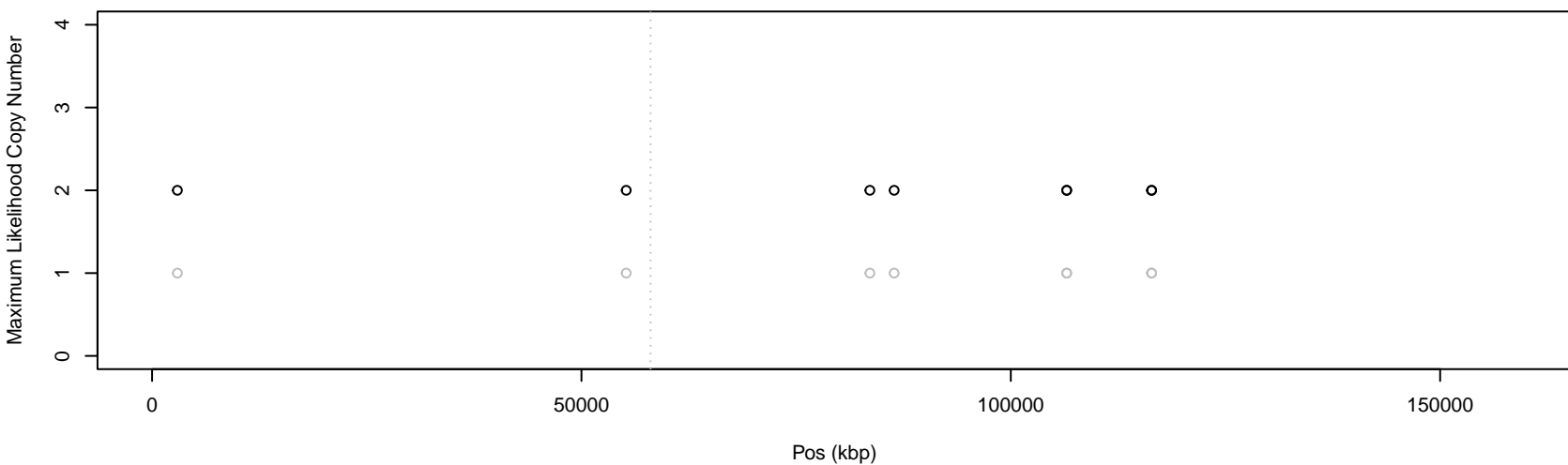
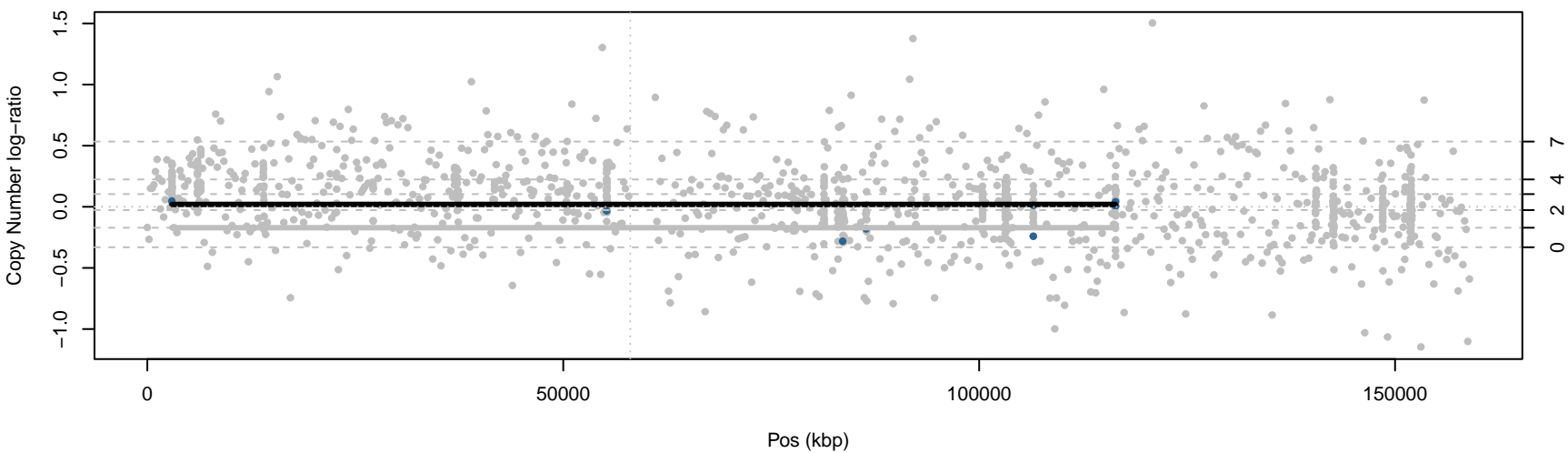
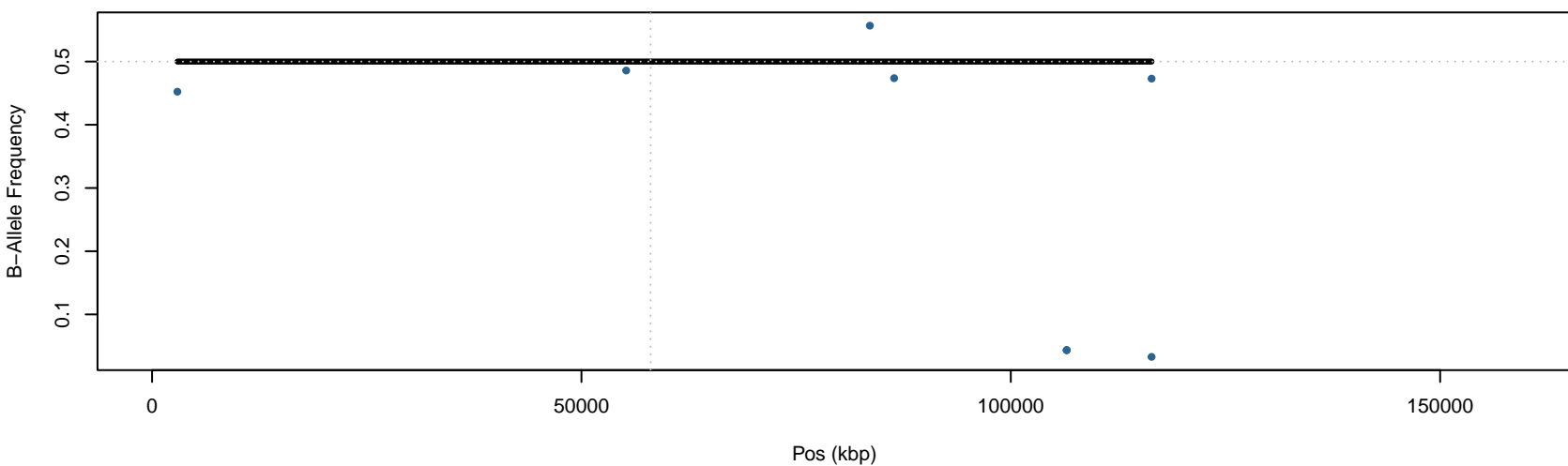
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 5



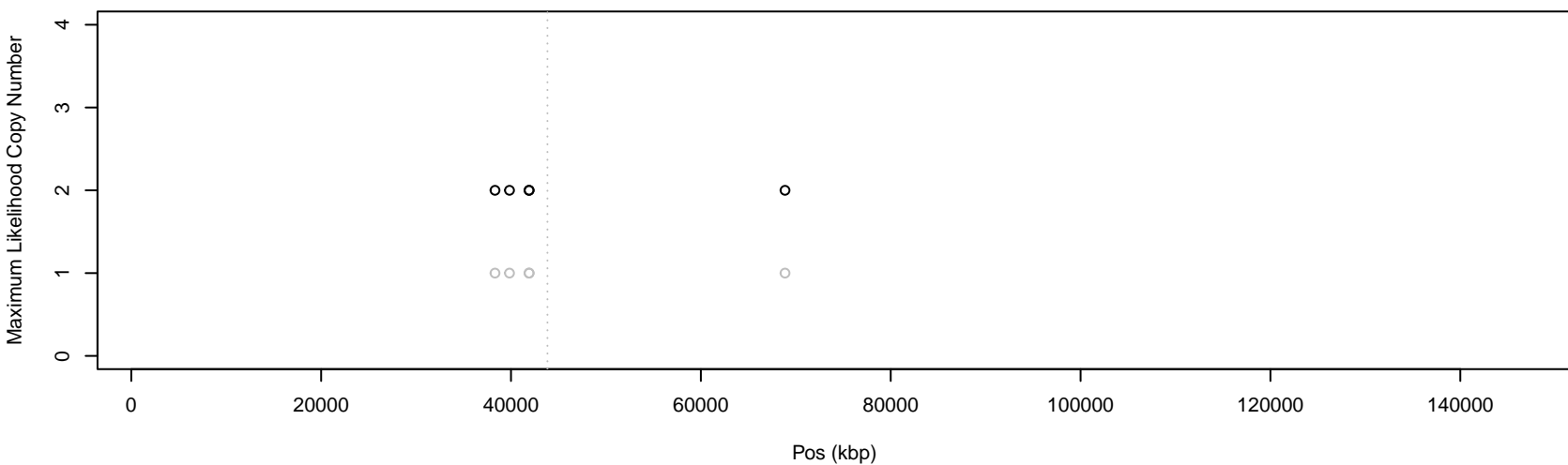
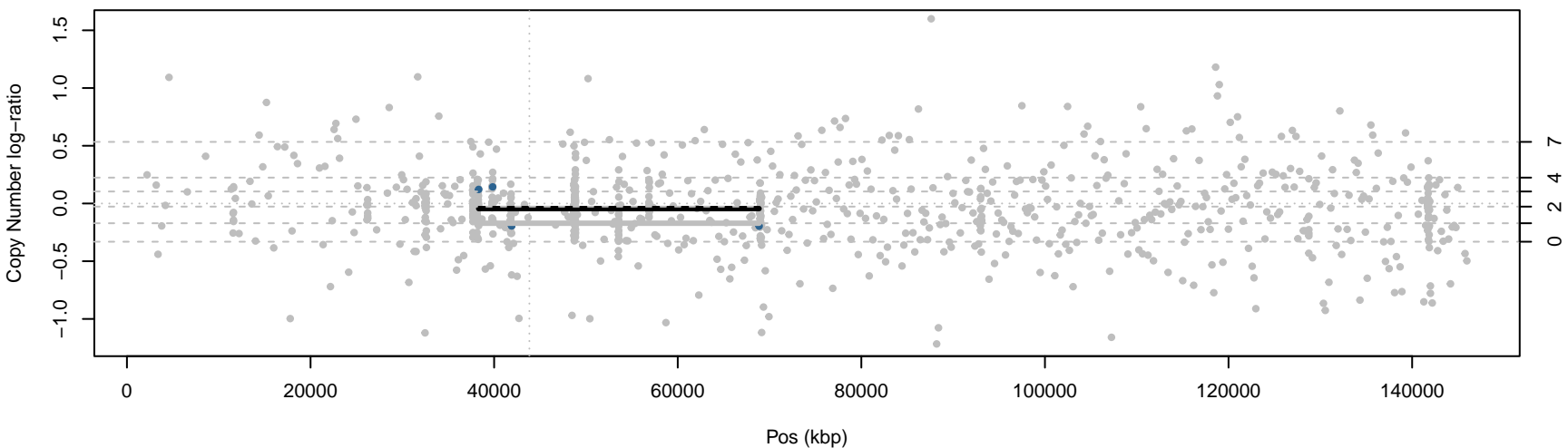
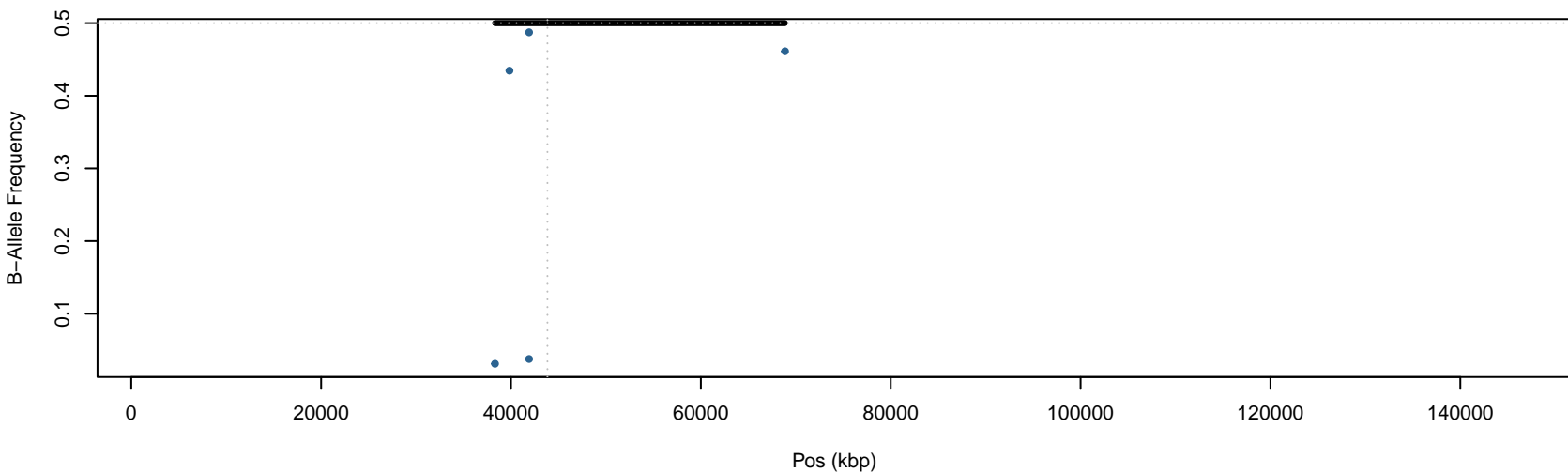
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 6



Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 7

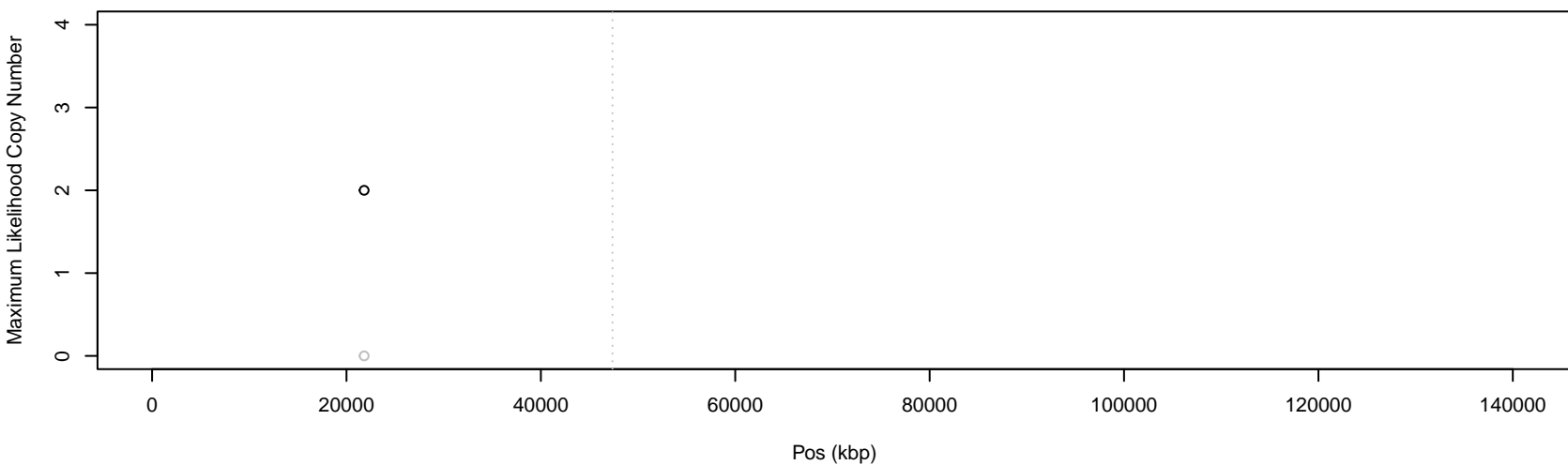
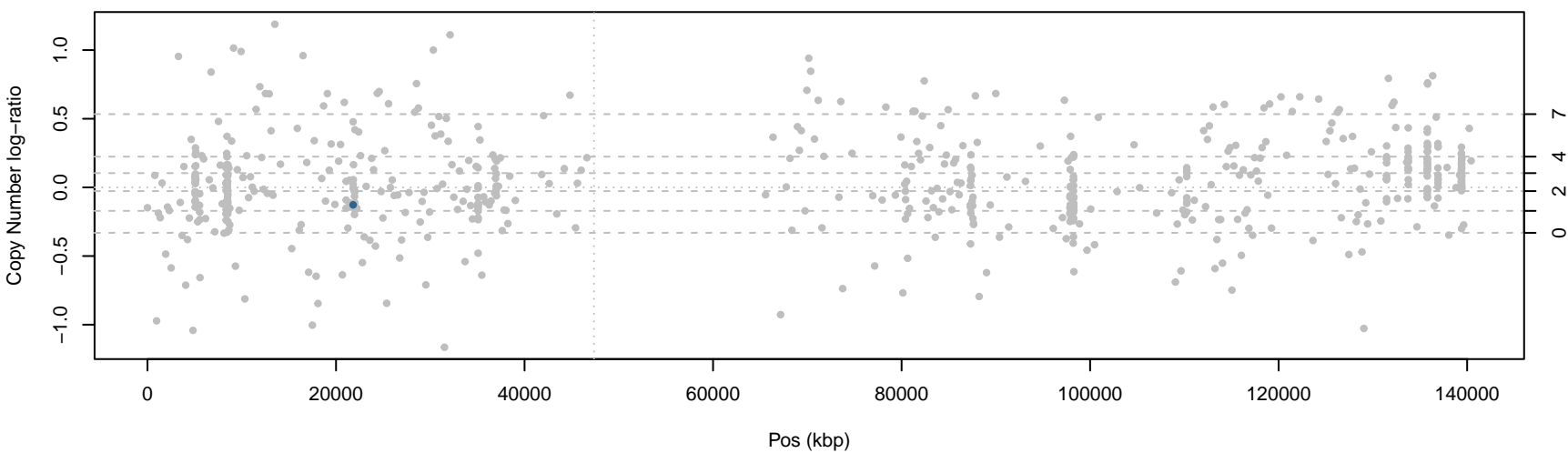
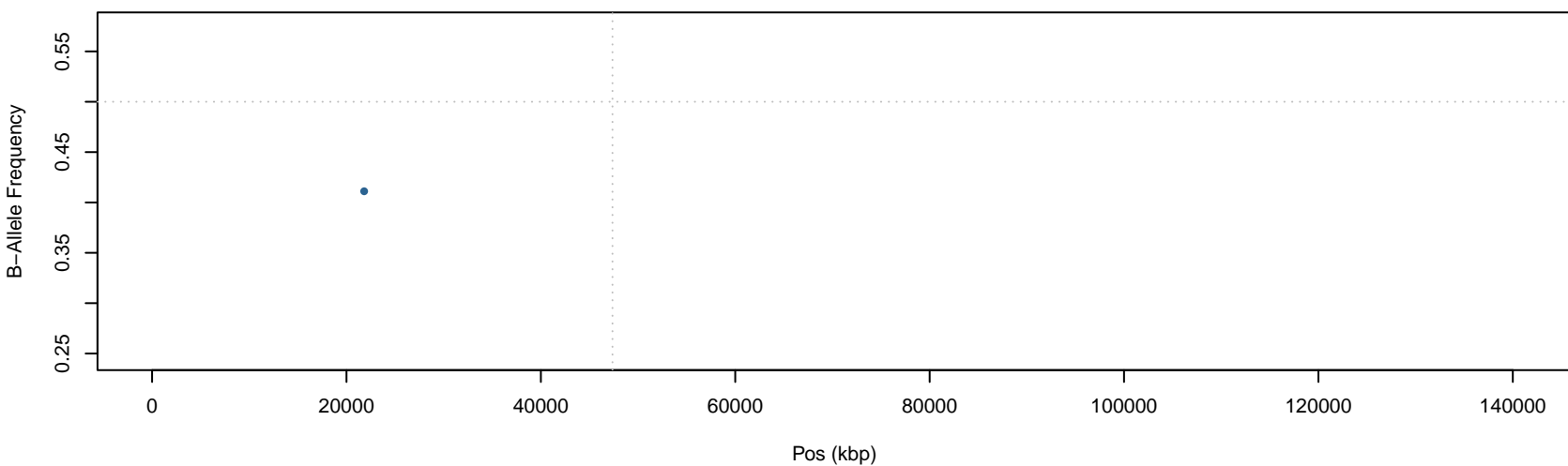


Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 8

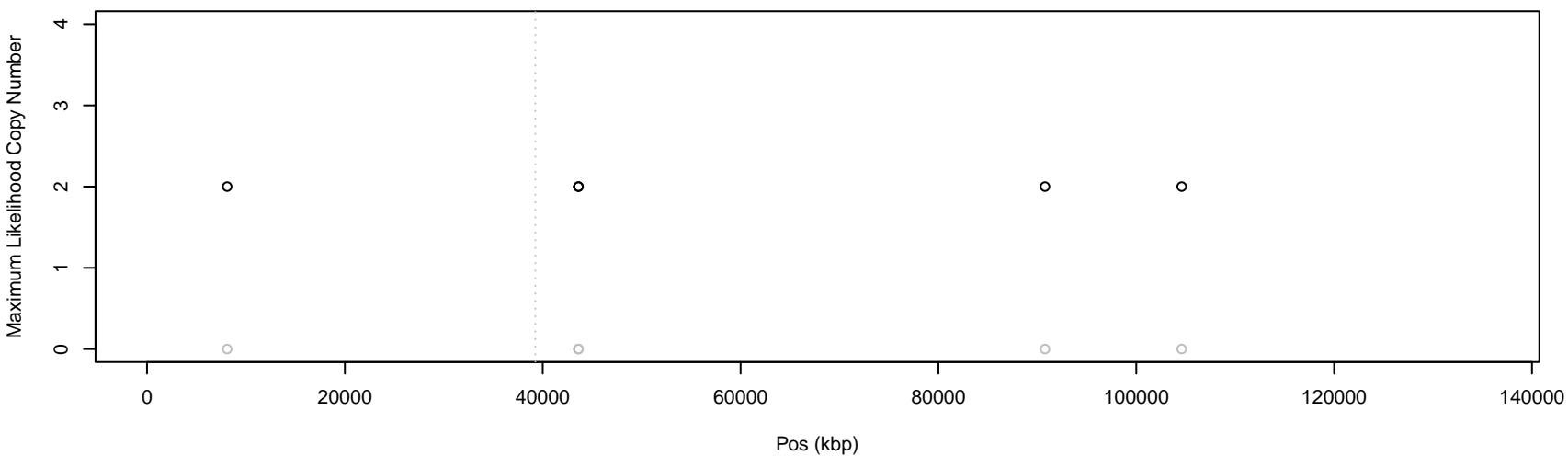
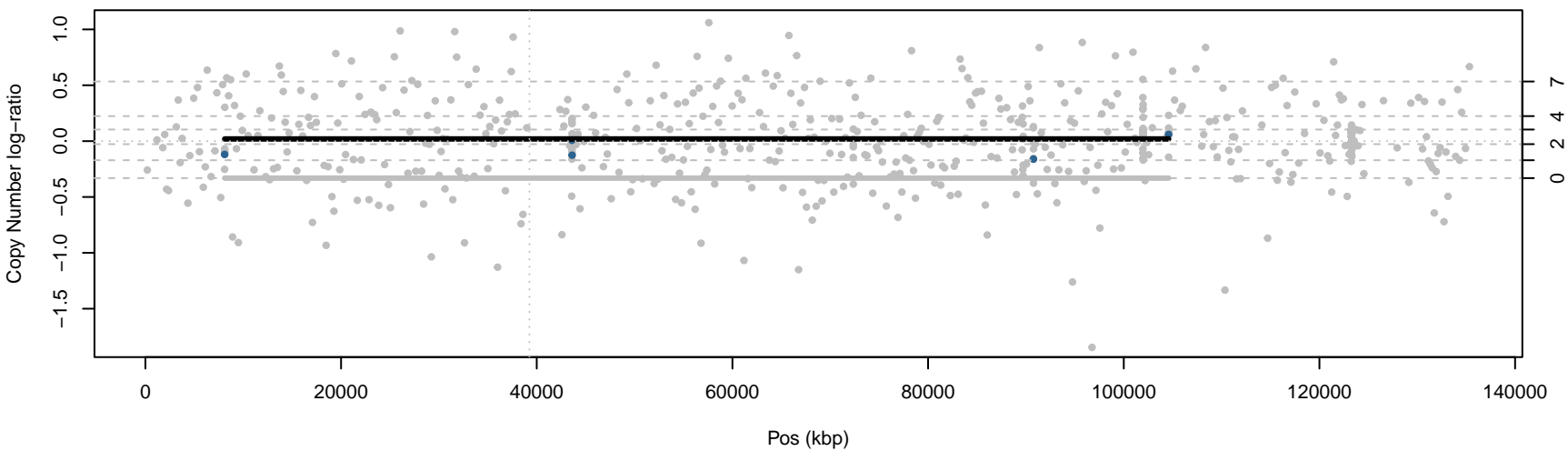
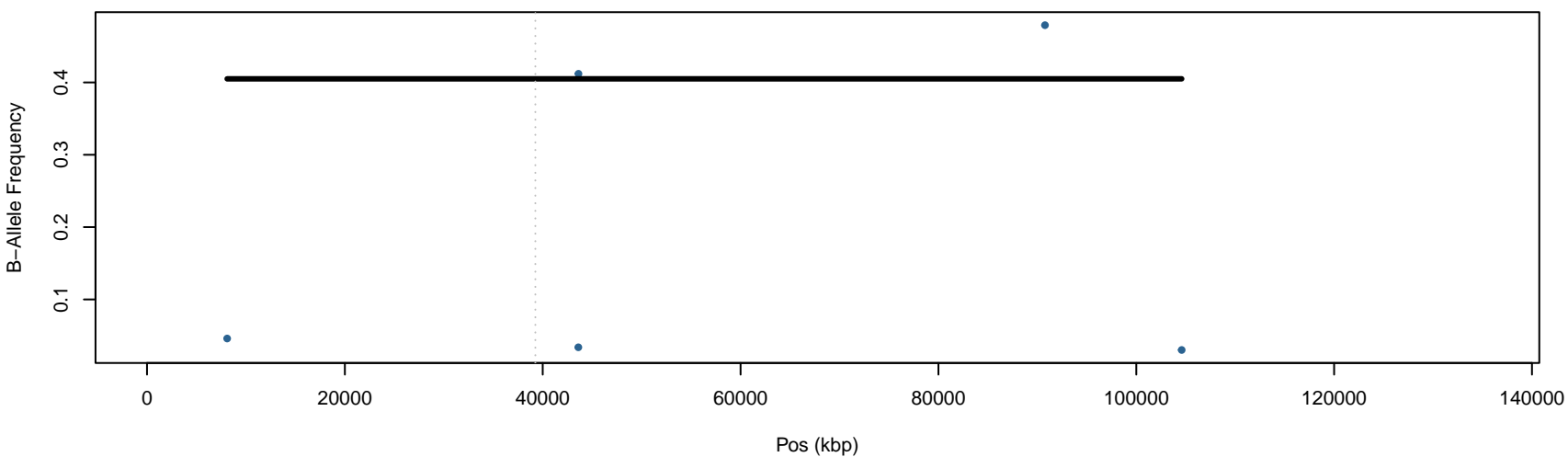




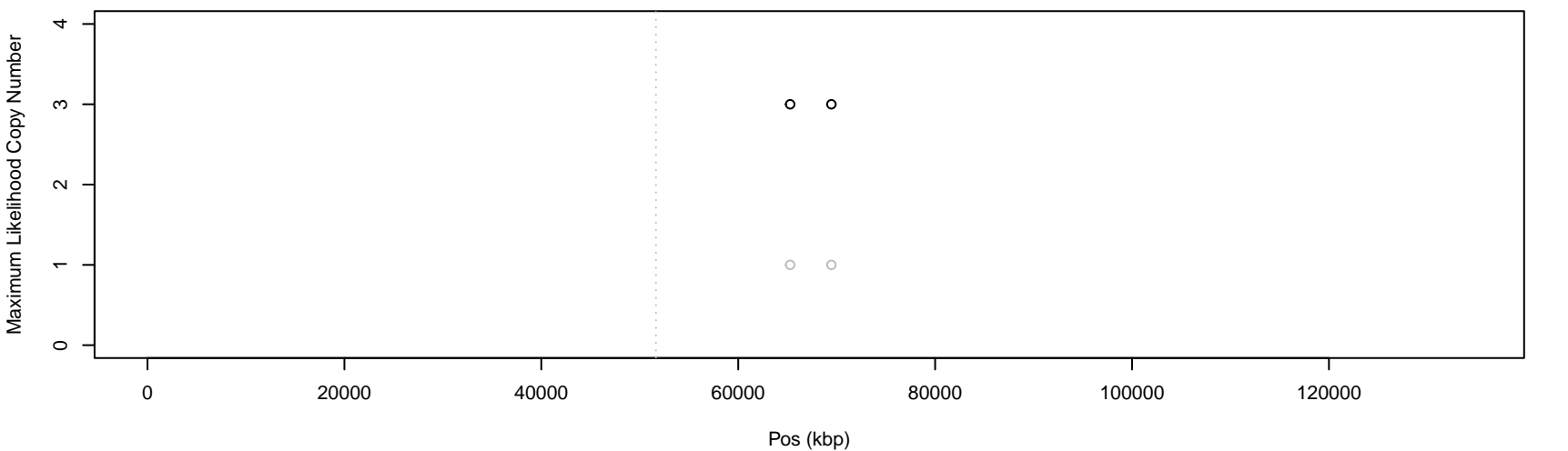
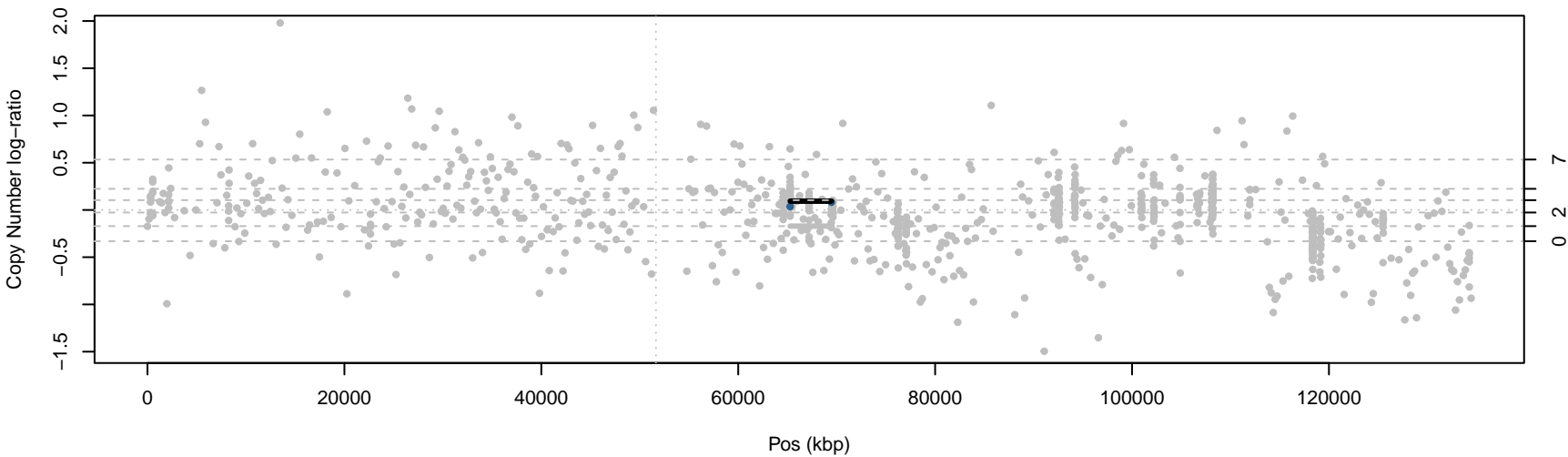
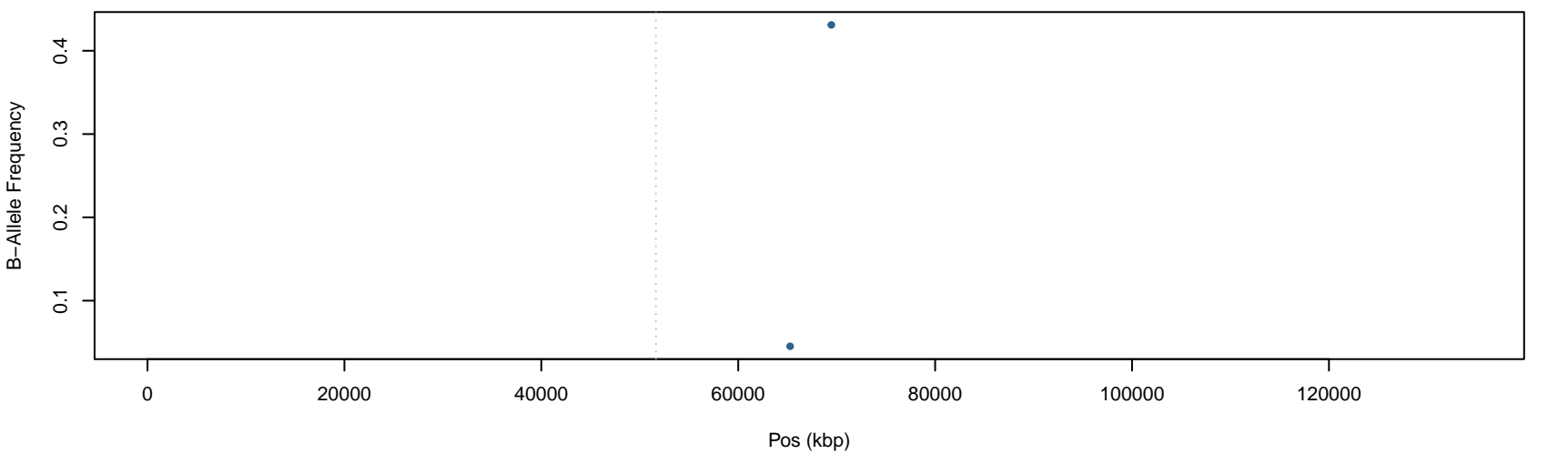
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 9



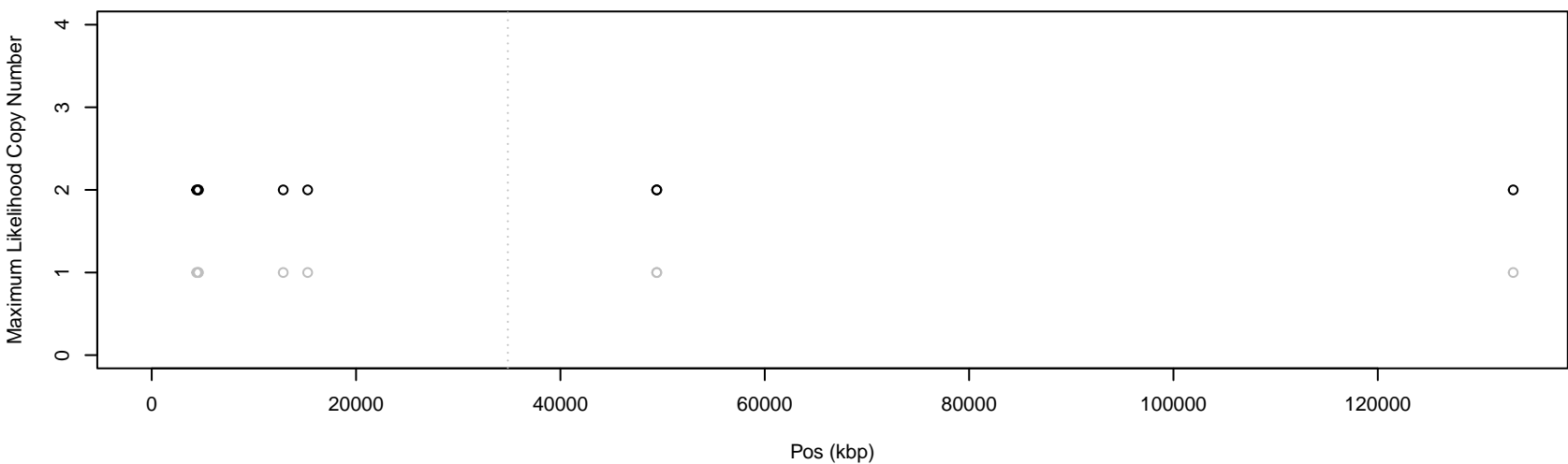
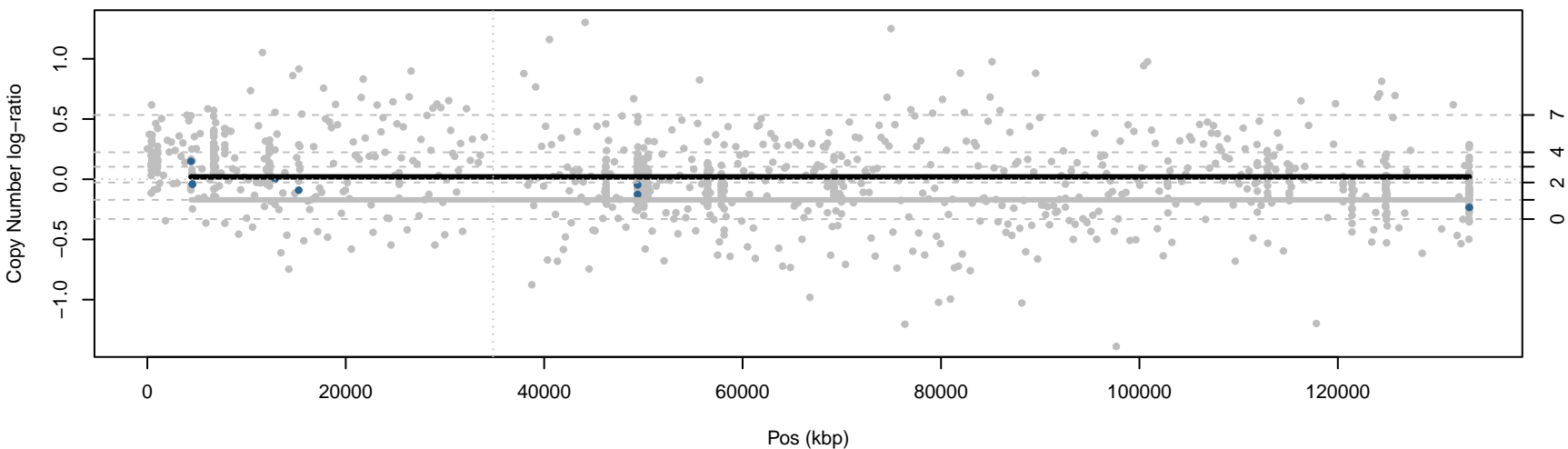
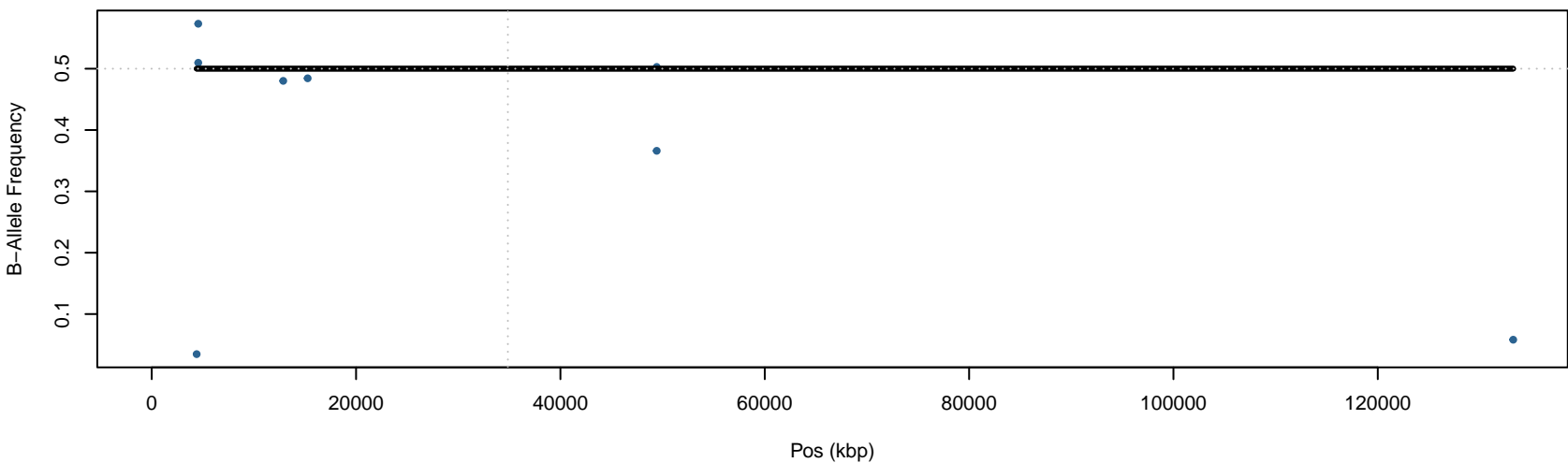
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 10



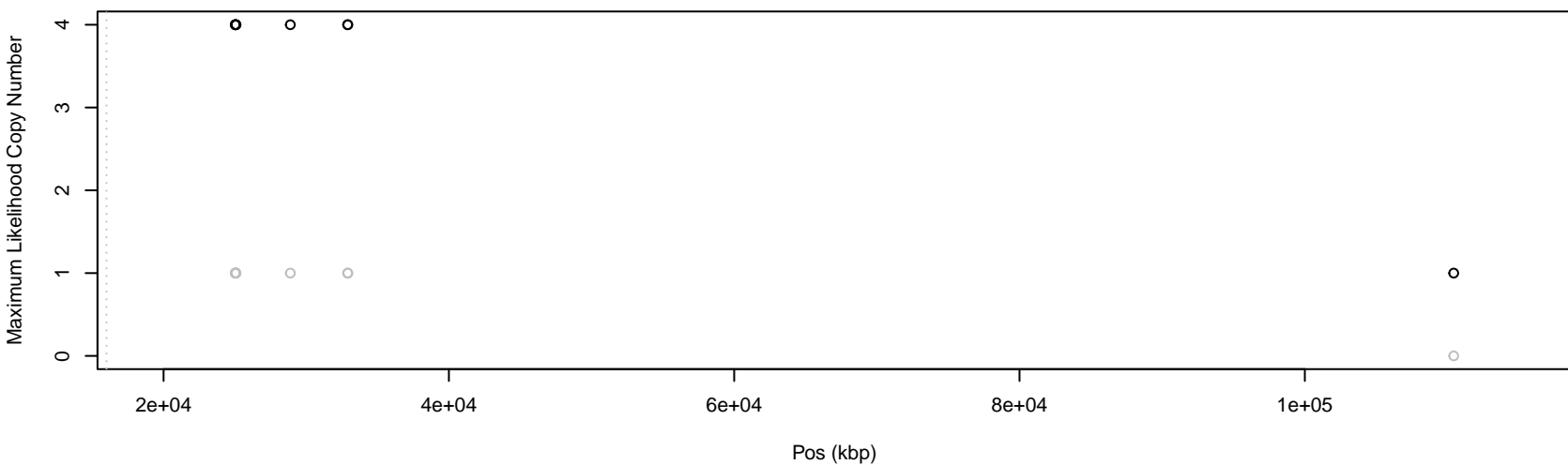
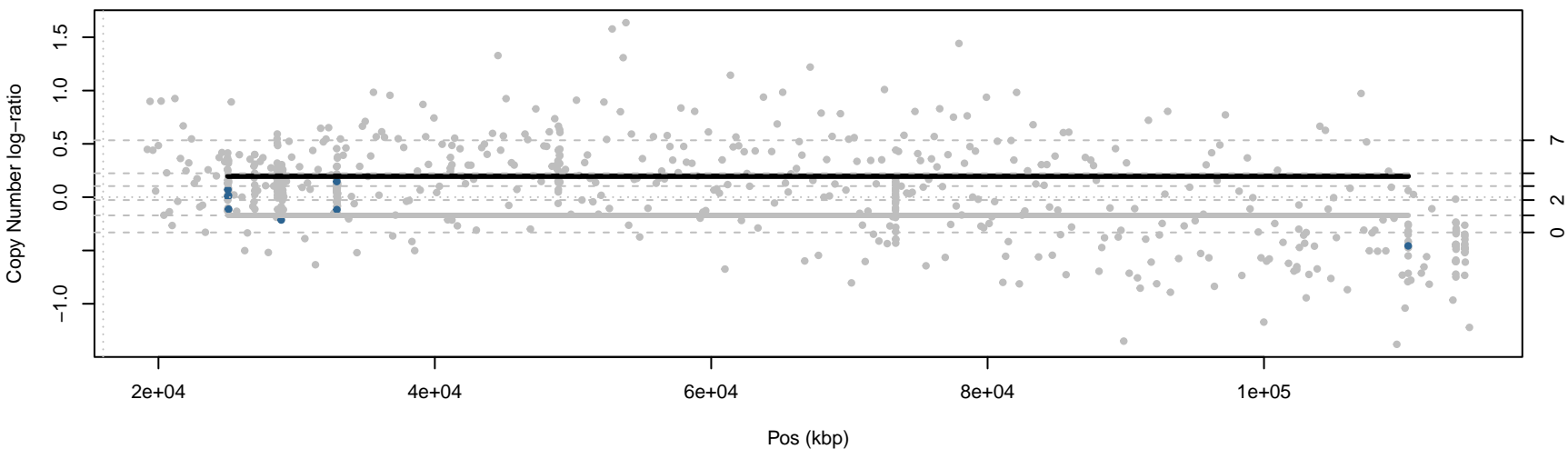
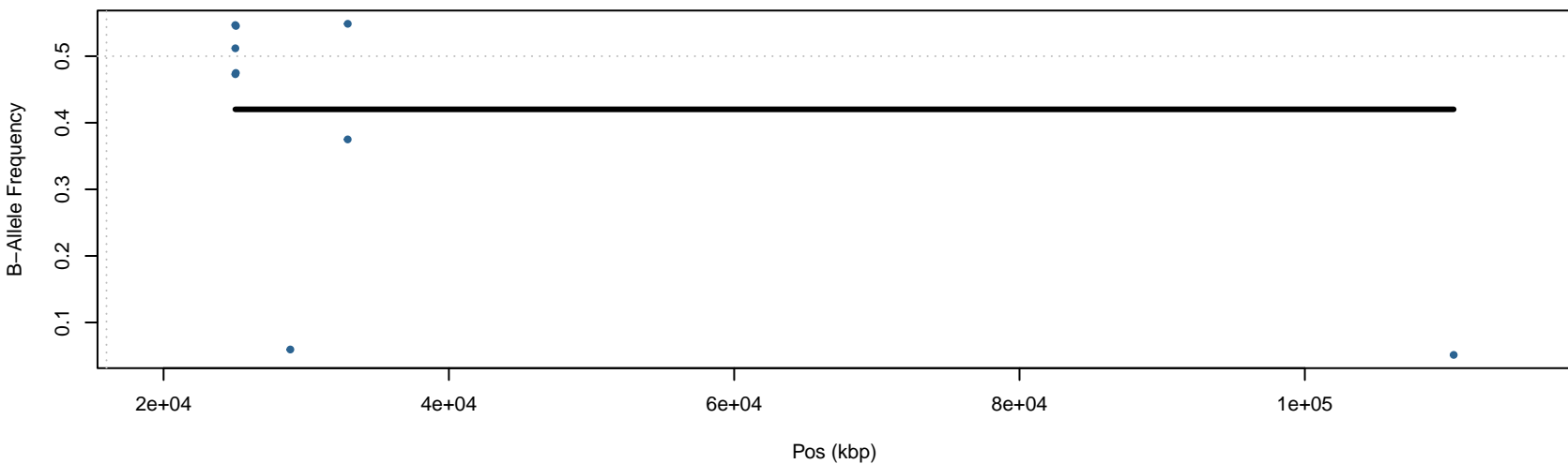
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 11



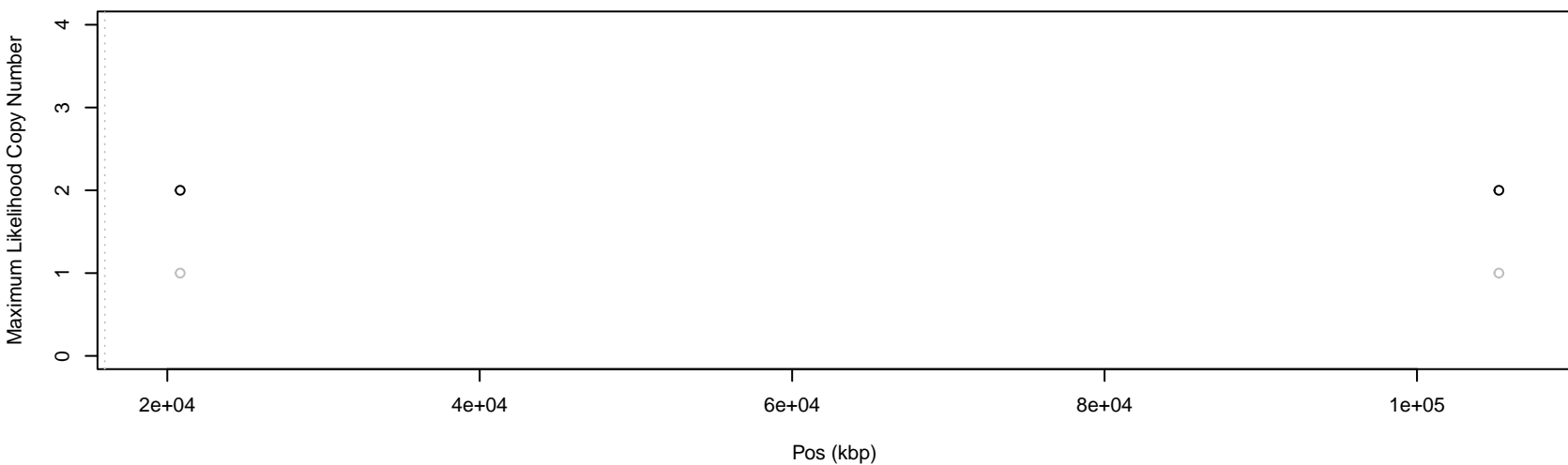
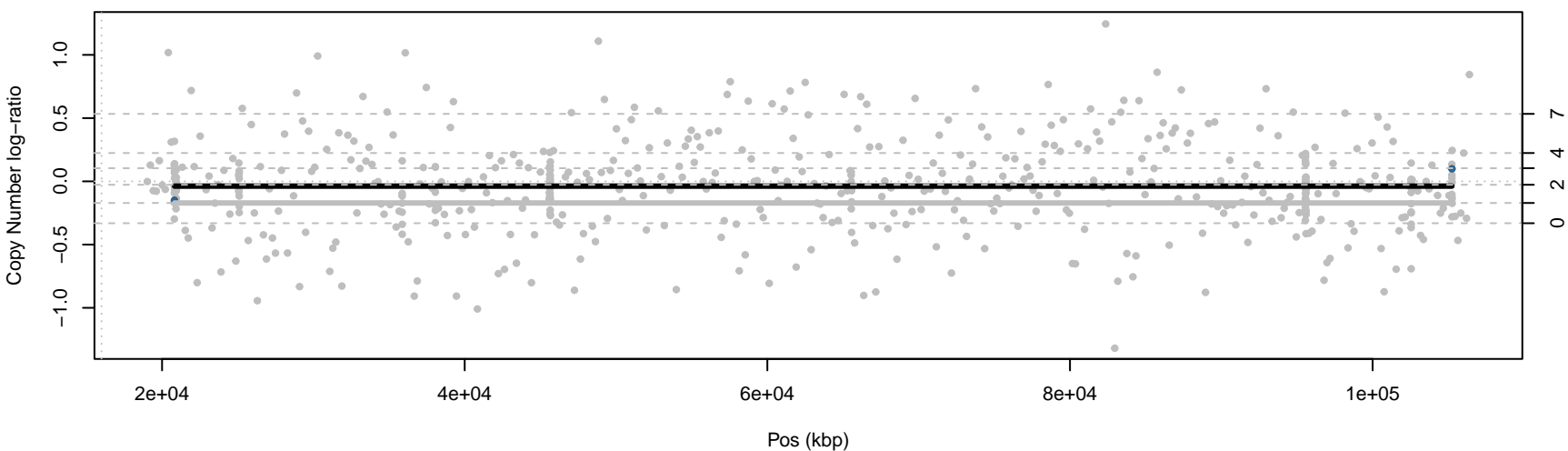
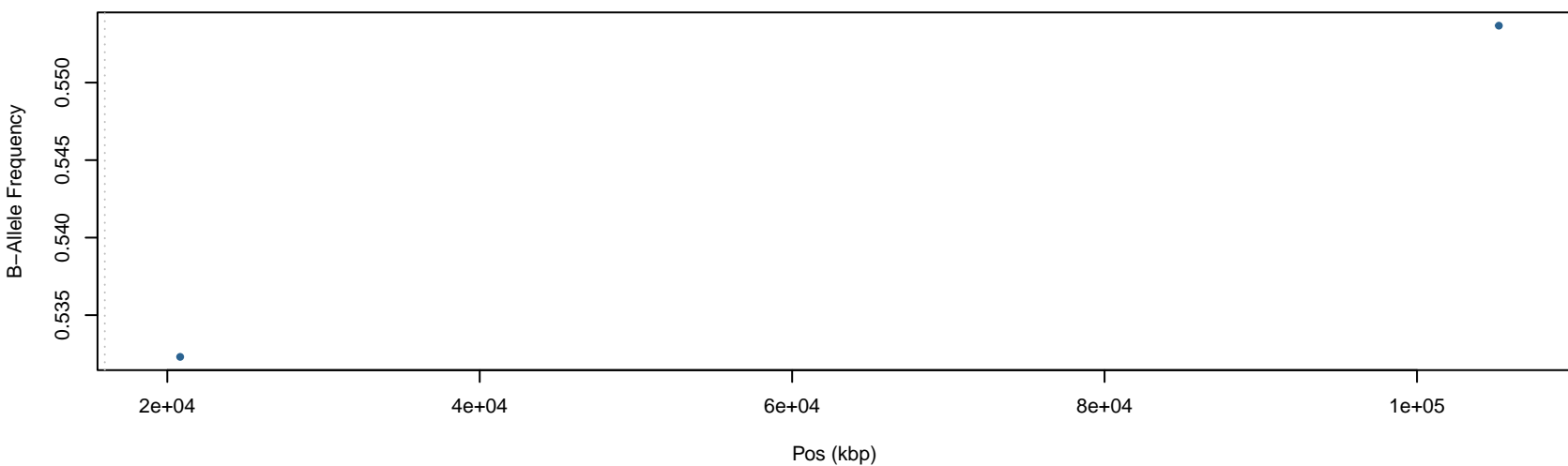
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 12



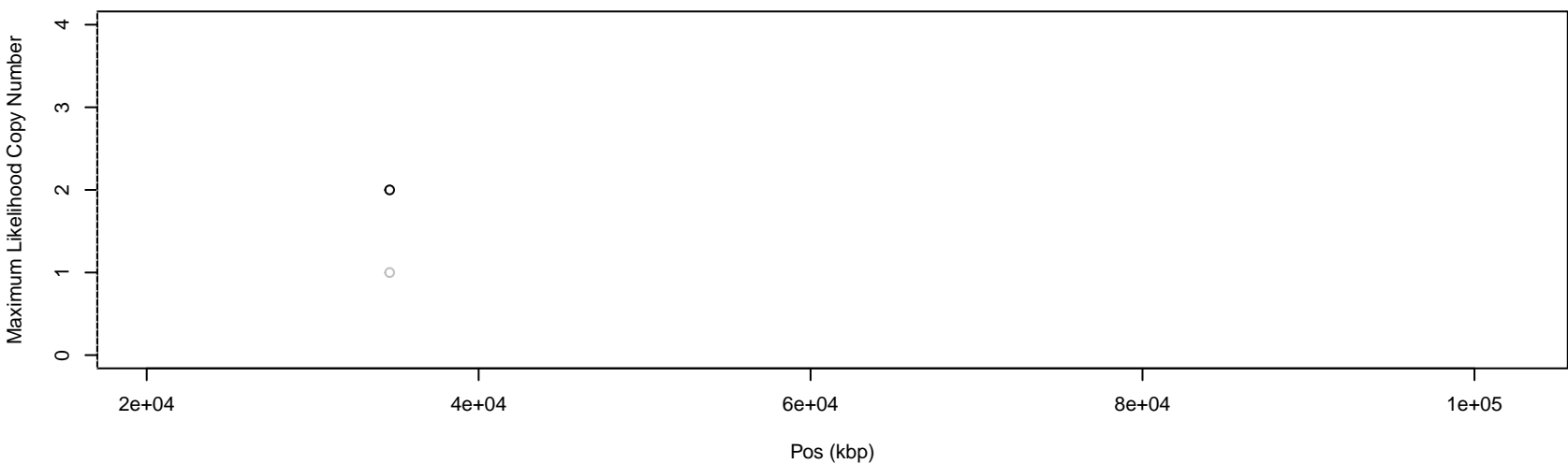
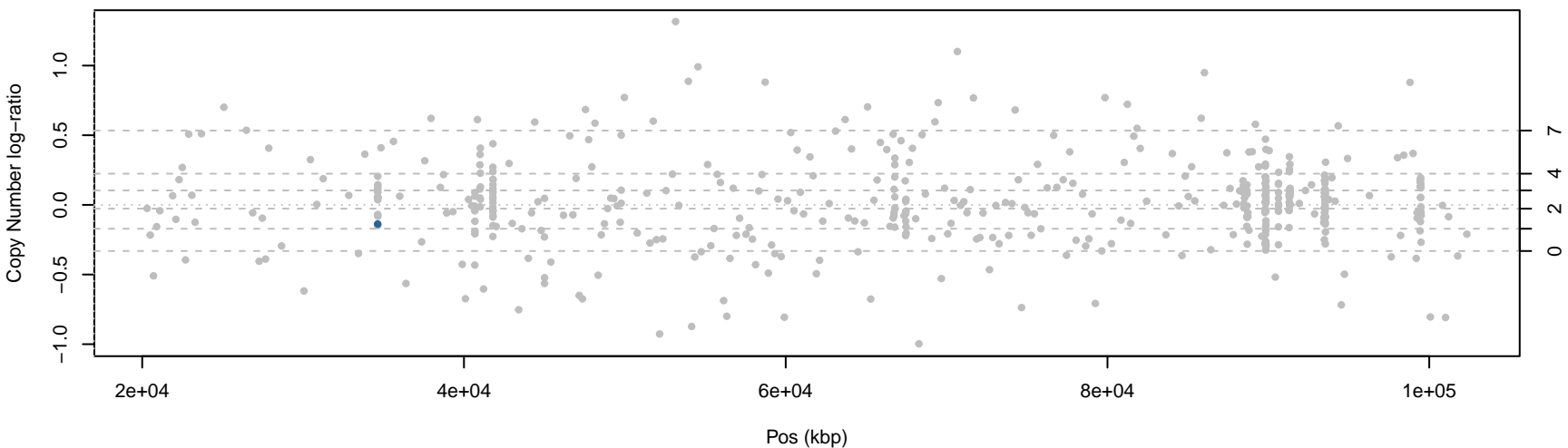
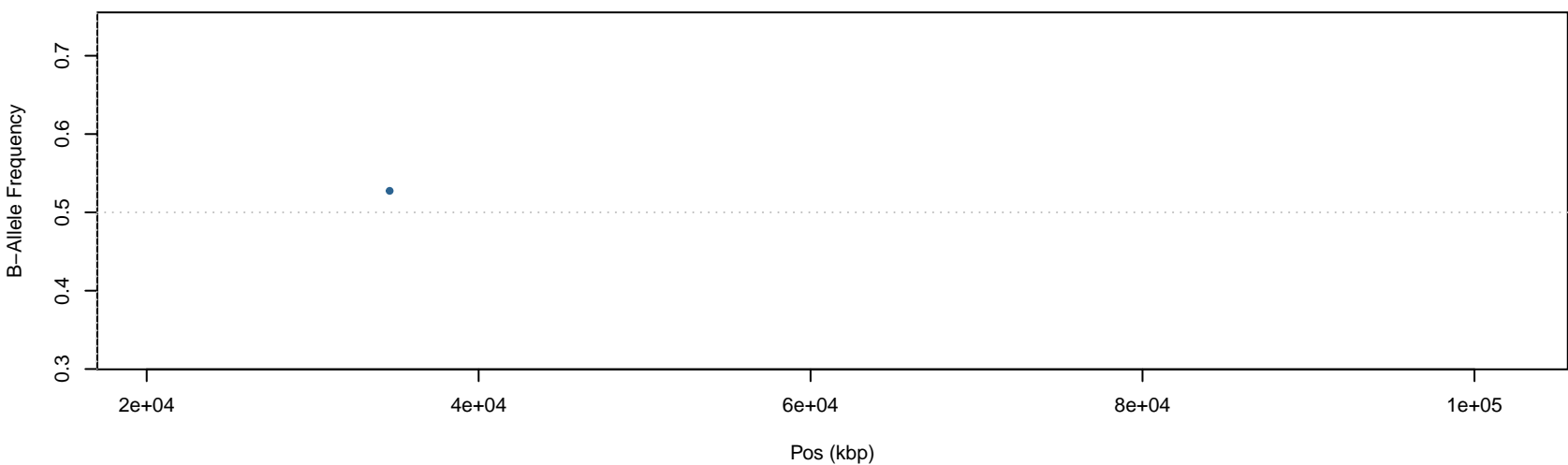
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 13



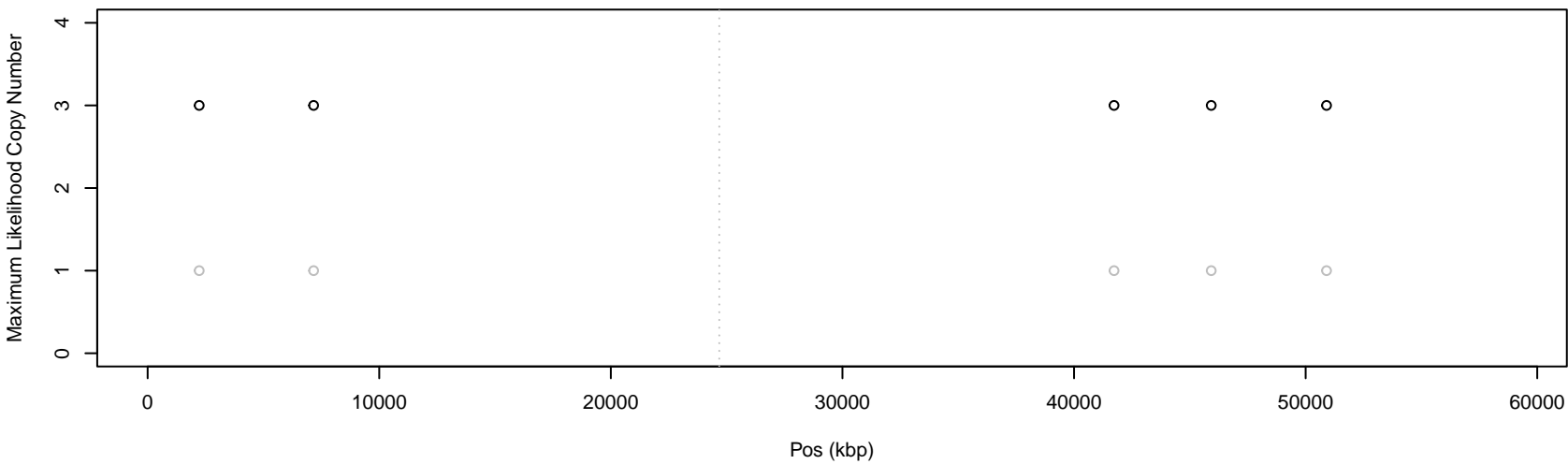
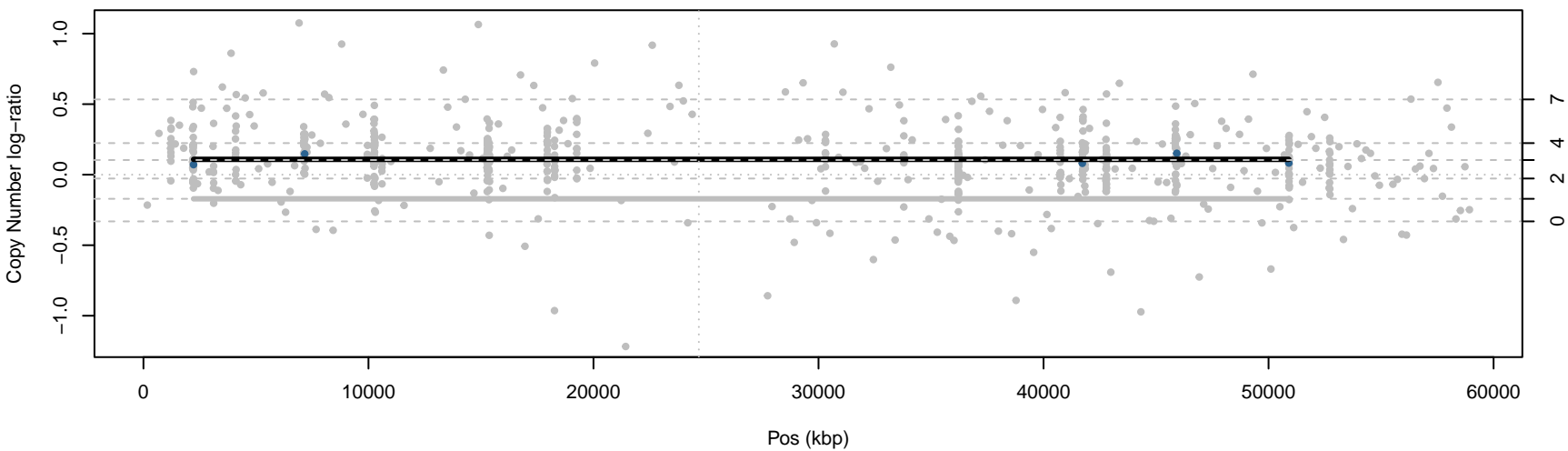
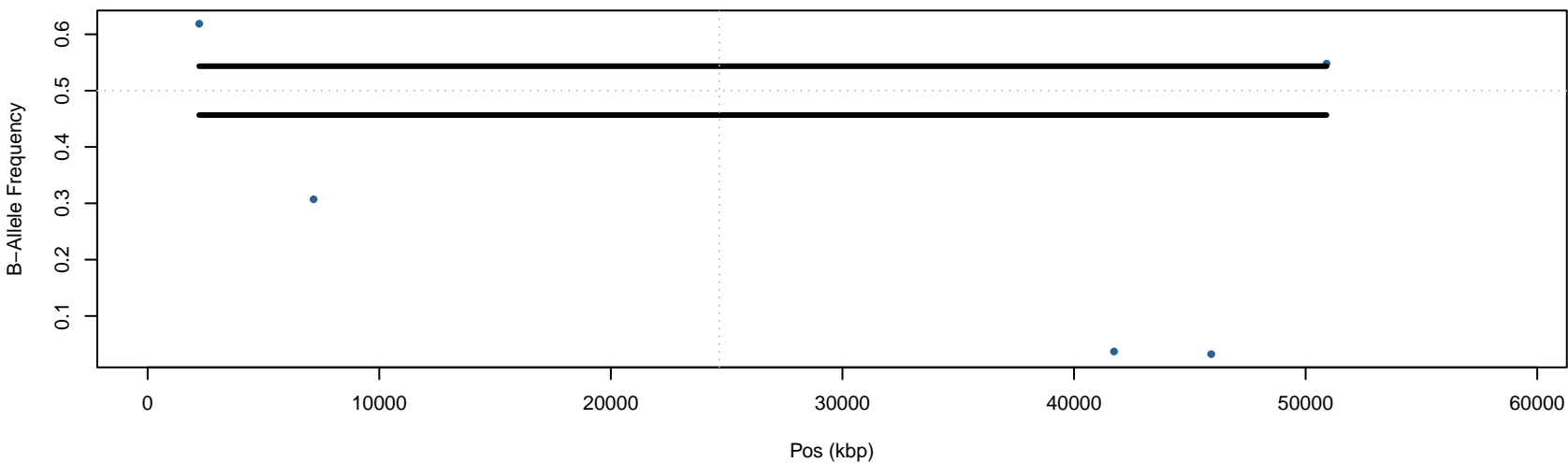
Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 14



Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 15



Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 19





Purity: 0.19 Tumor ploidy: 2.2 SNV log-likelihood: -77.41 GoF: 90.9% Mean coverage: 427;361 Chromosome: 20

