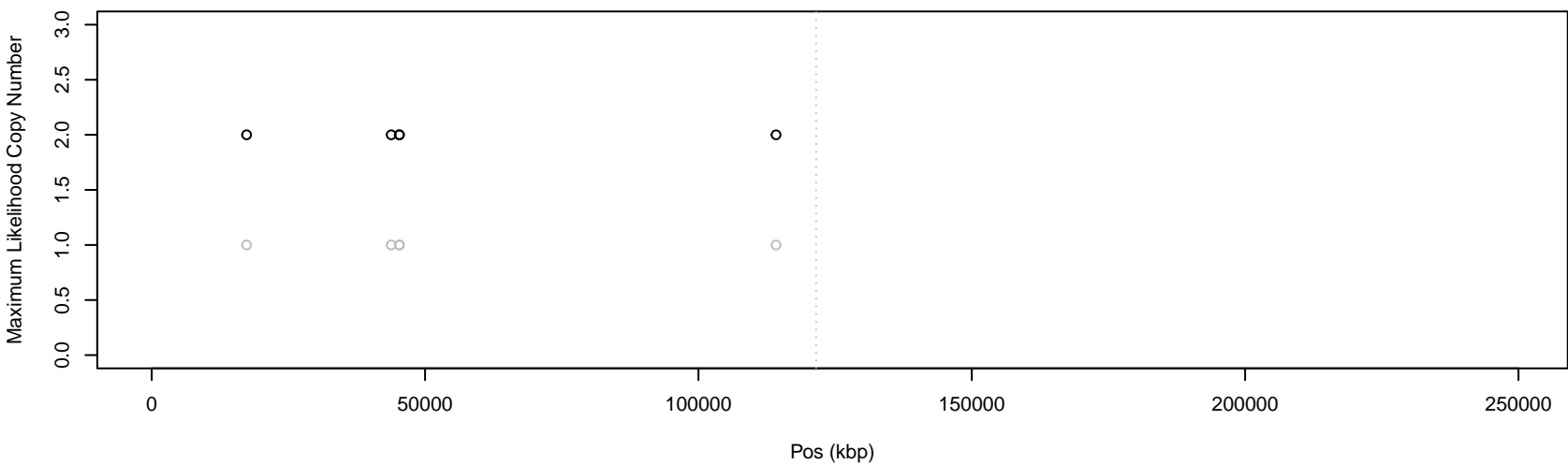
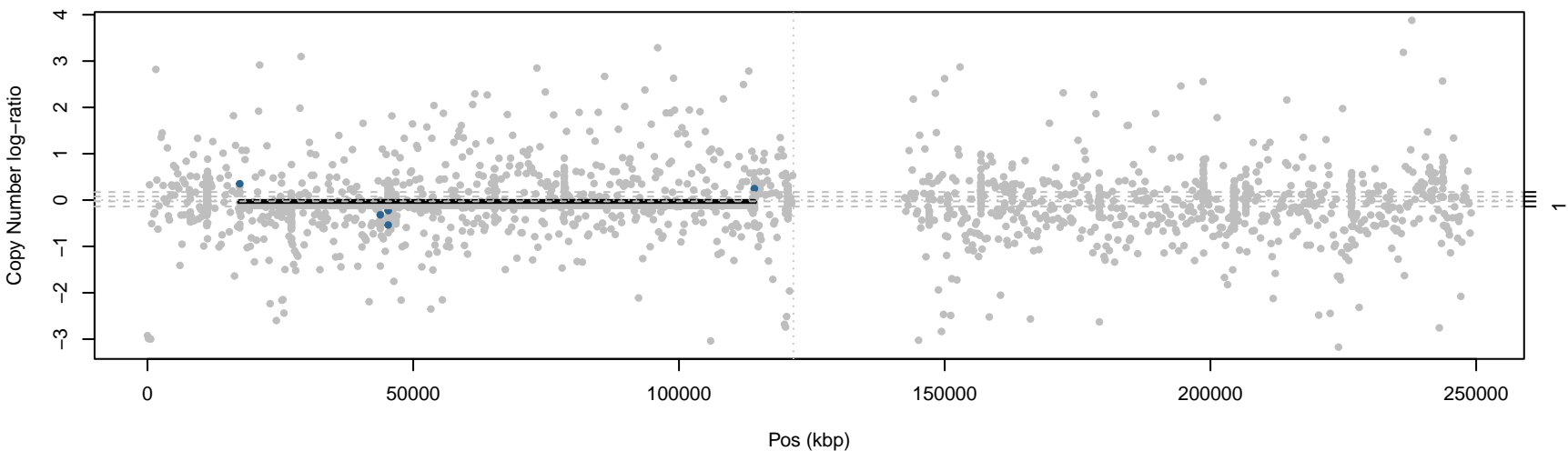
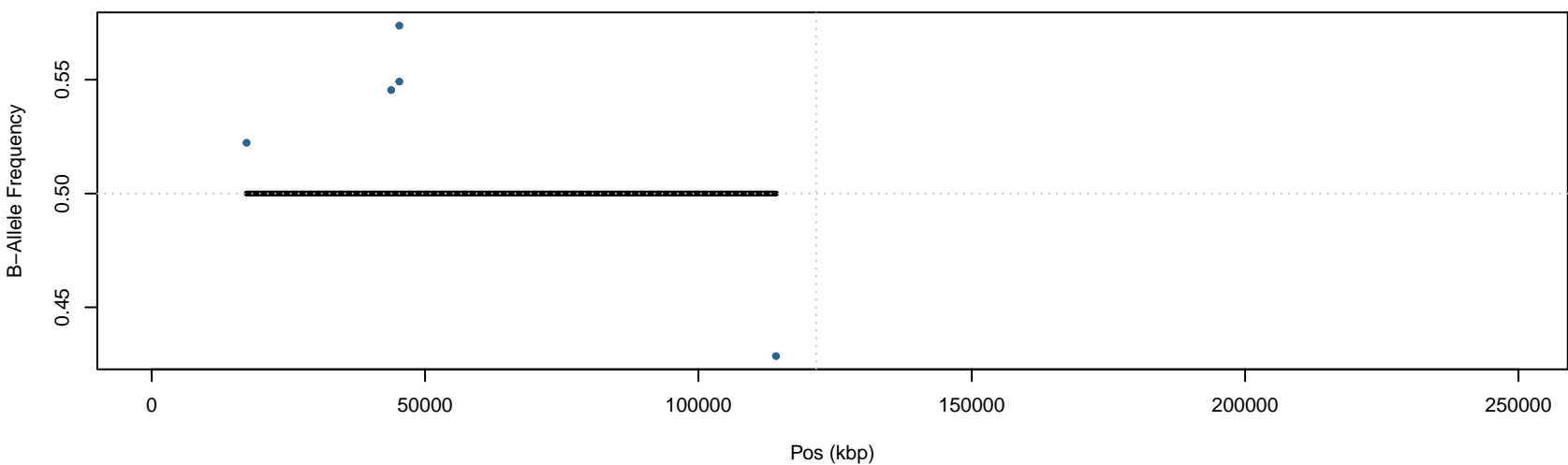
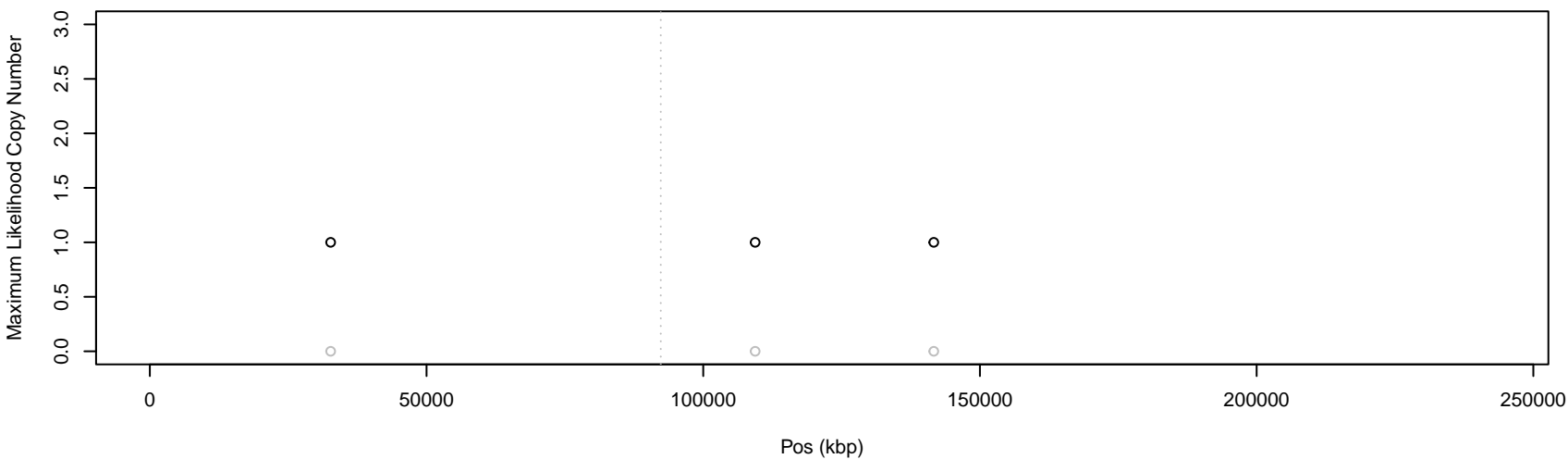
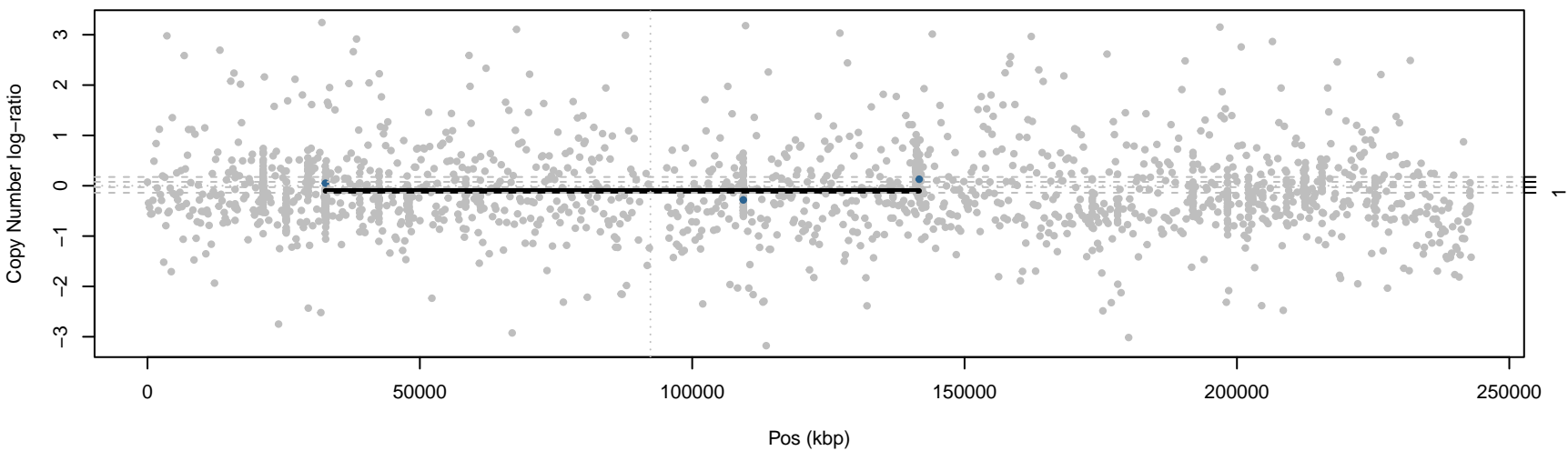
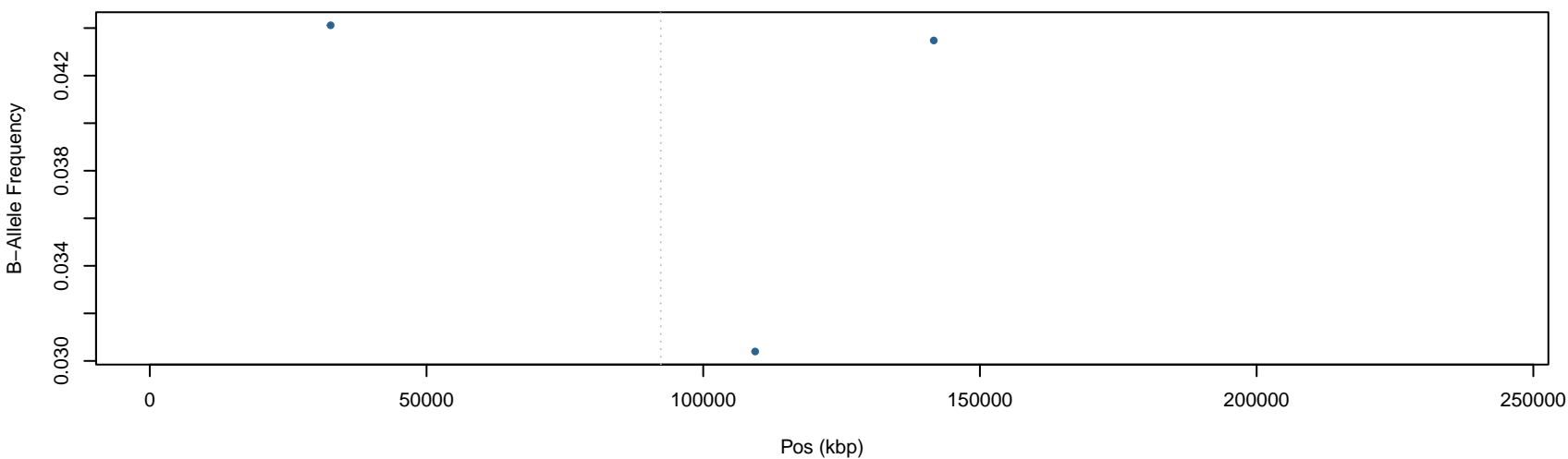


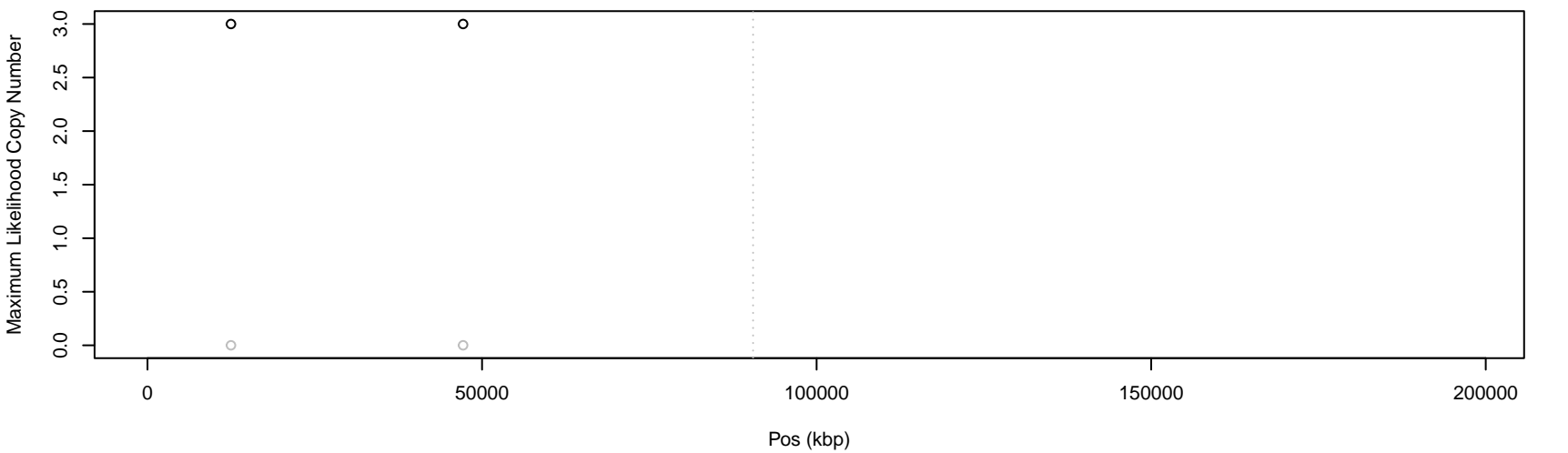
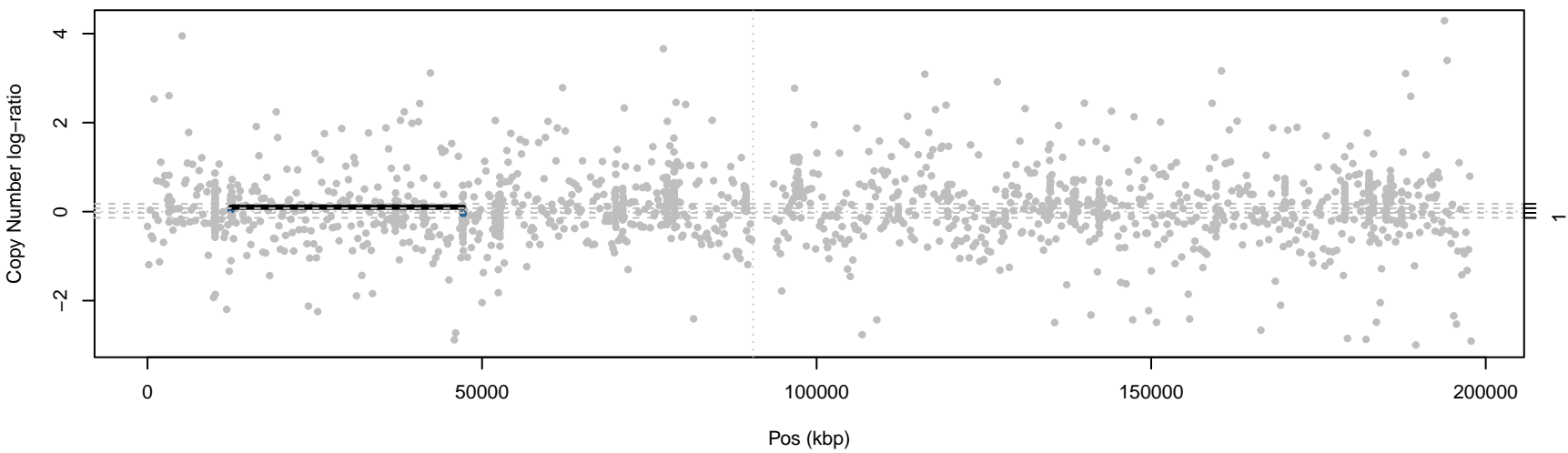
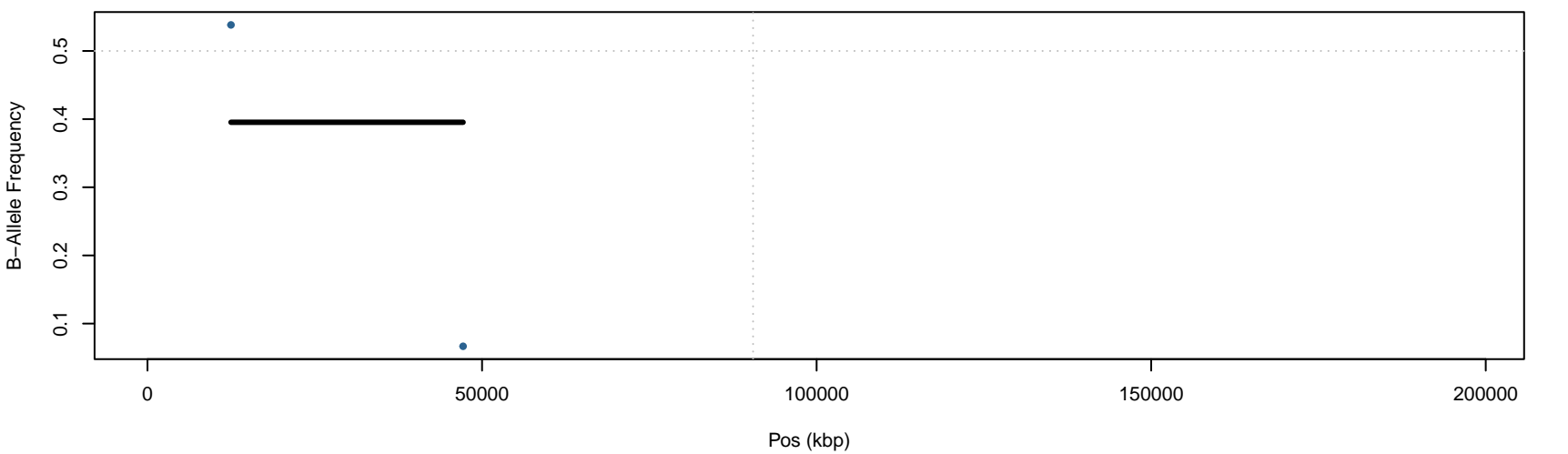
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 1



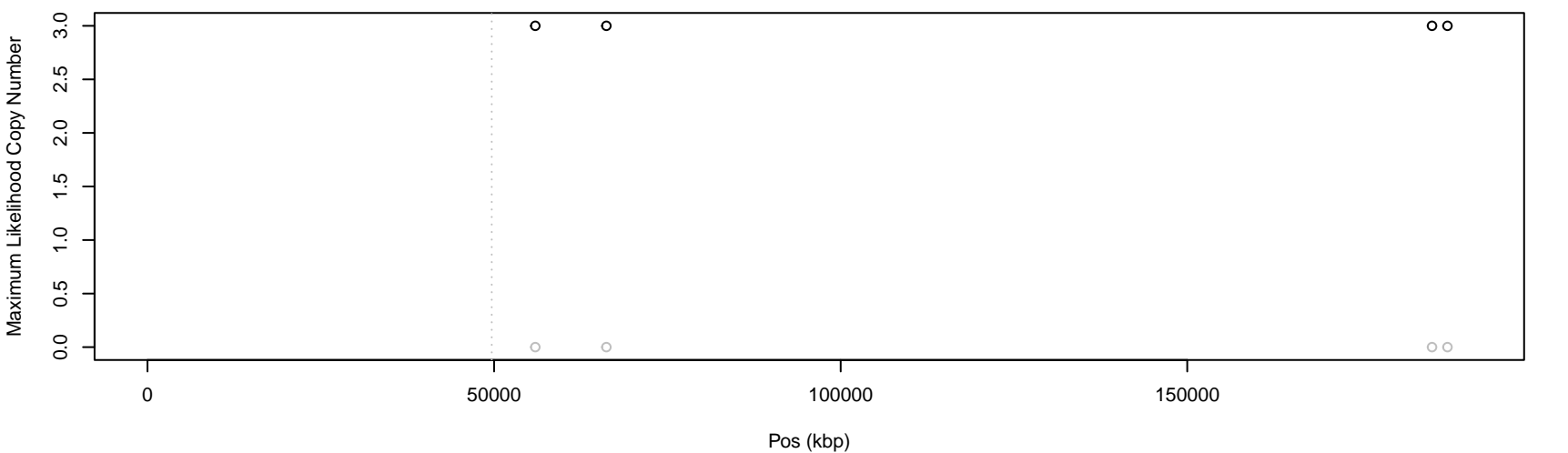
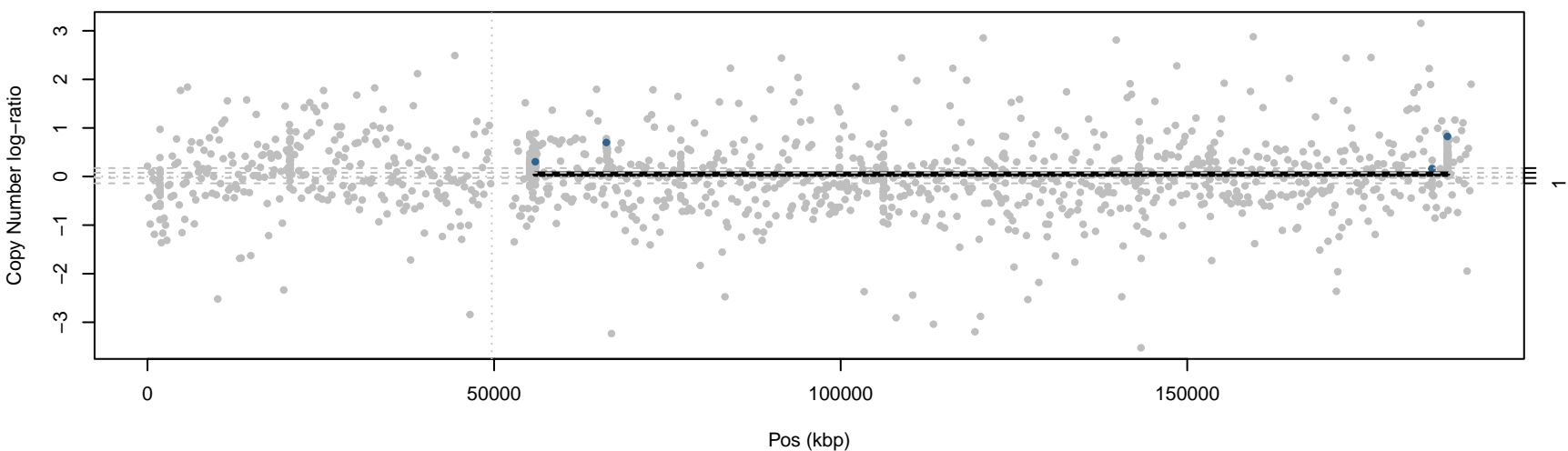
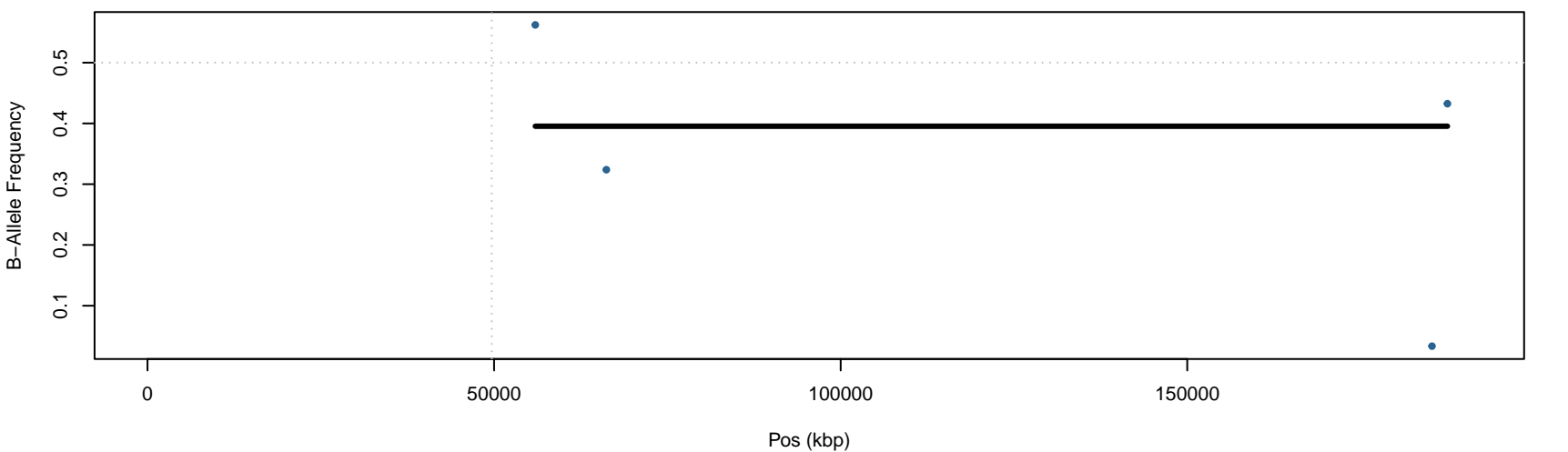
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 2



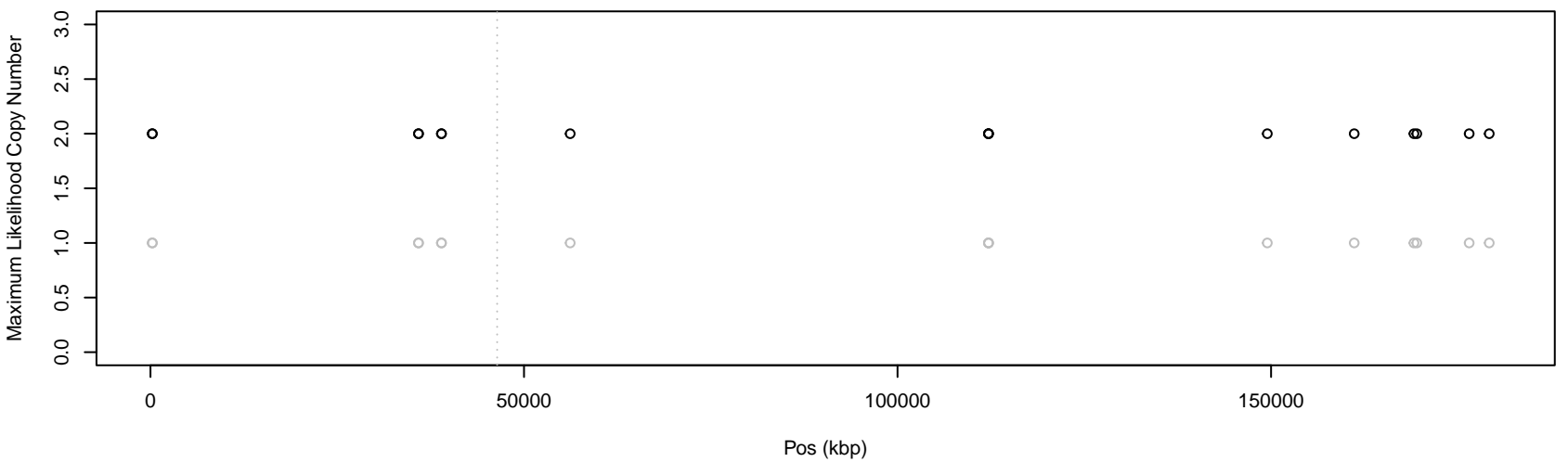
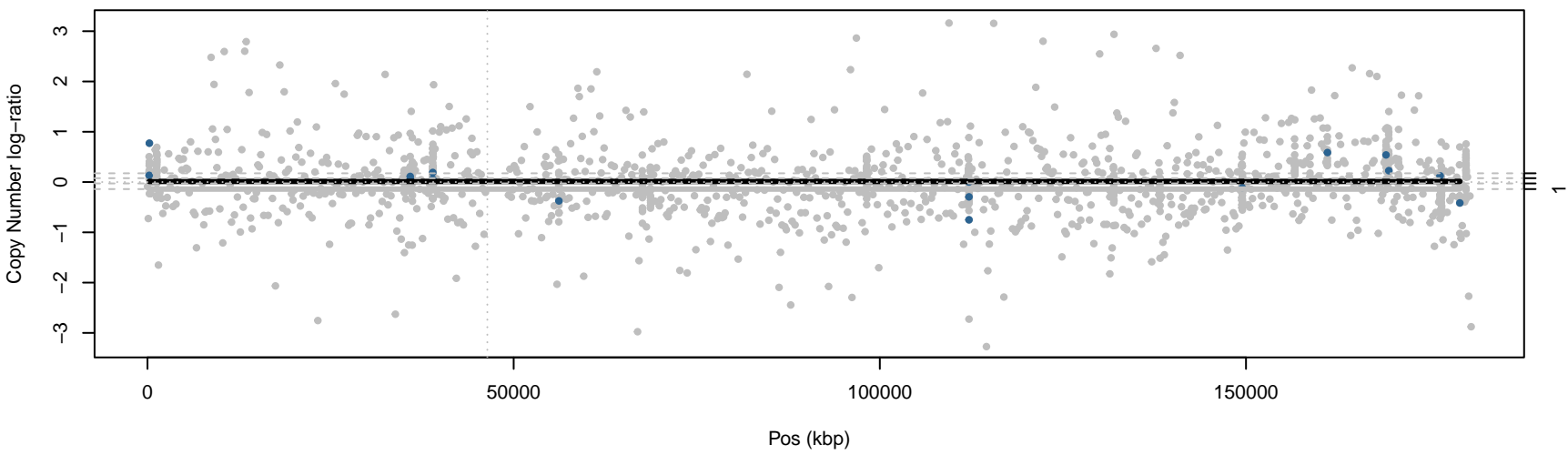
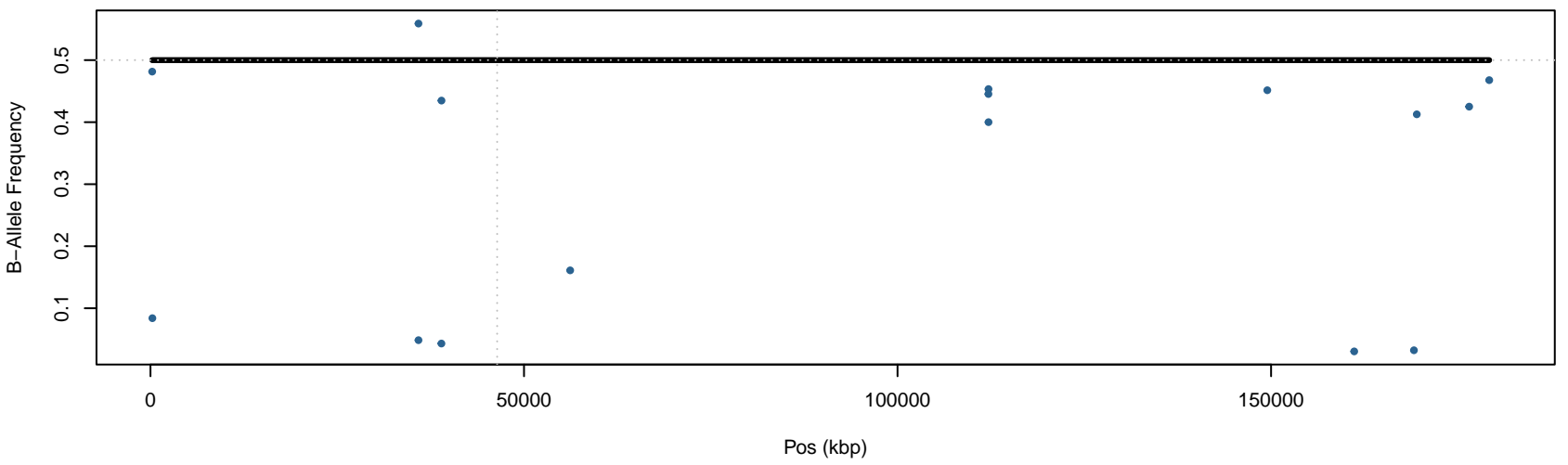
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 3



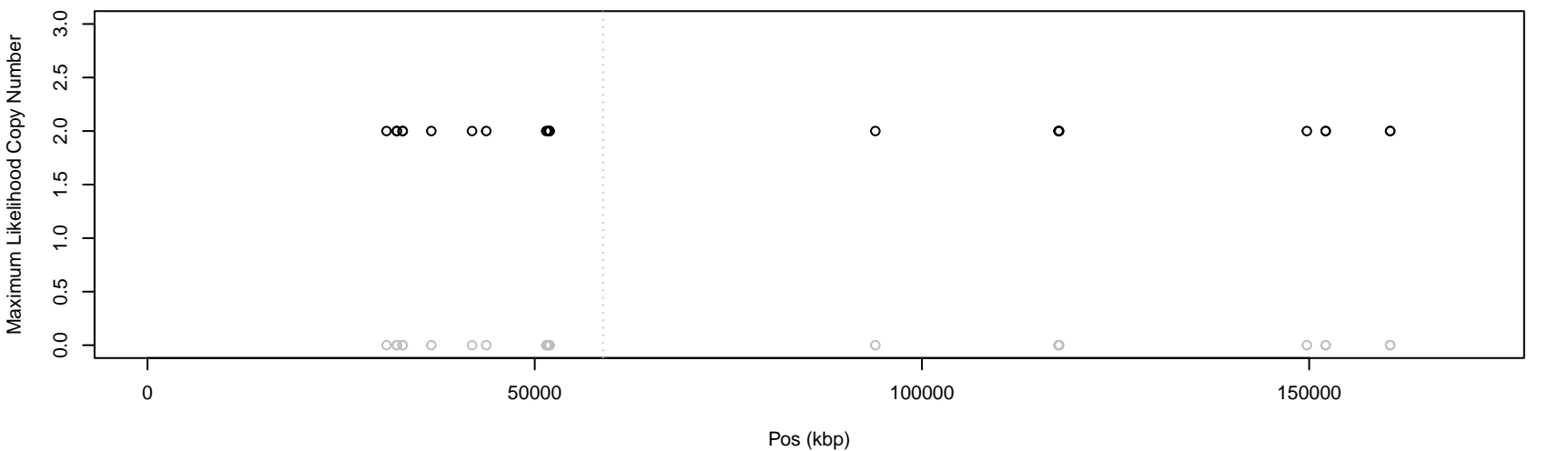
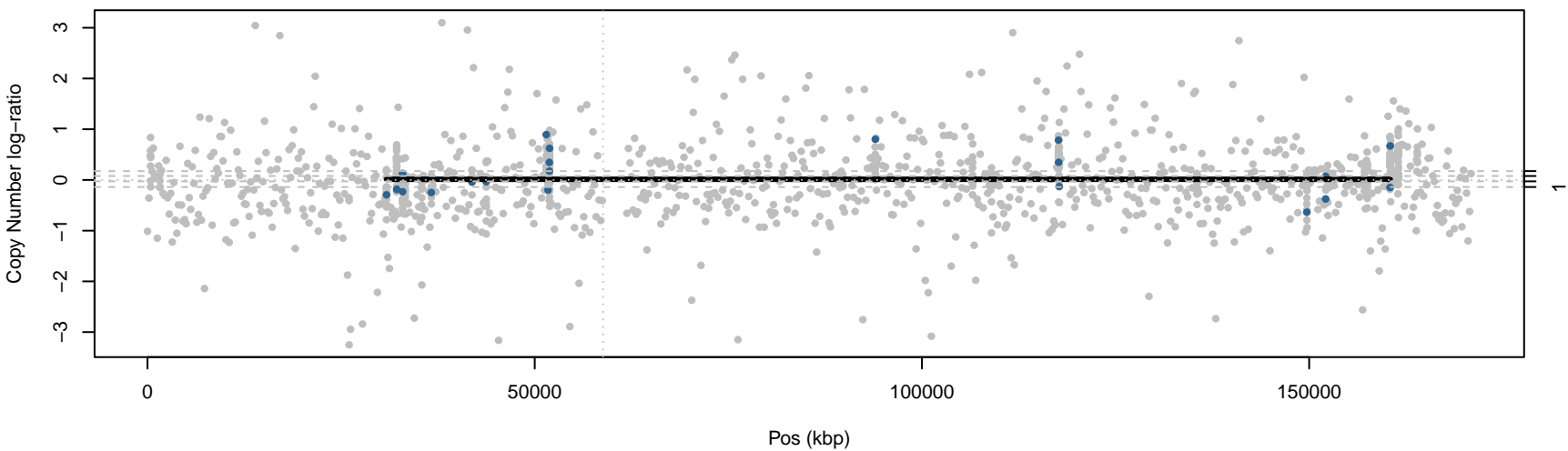
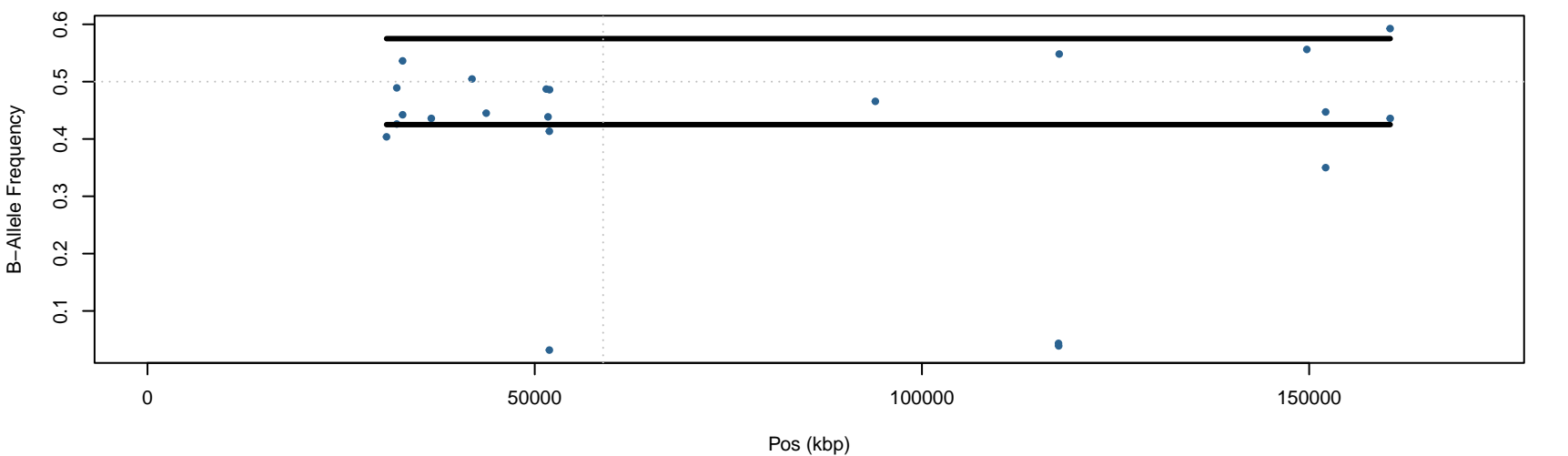
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 4



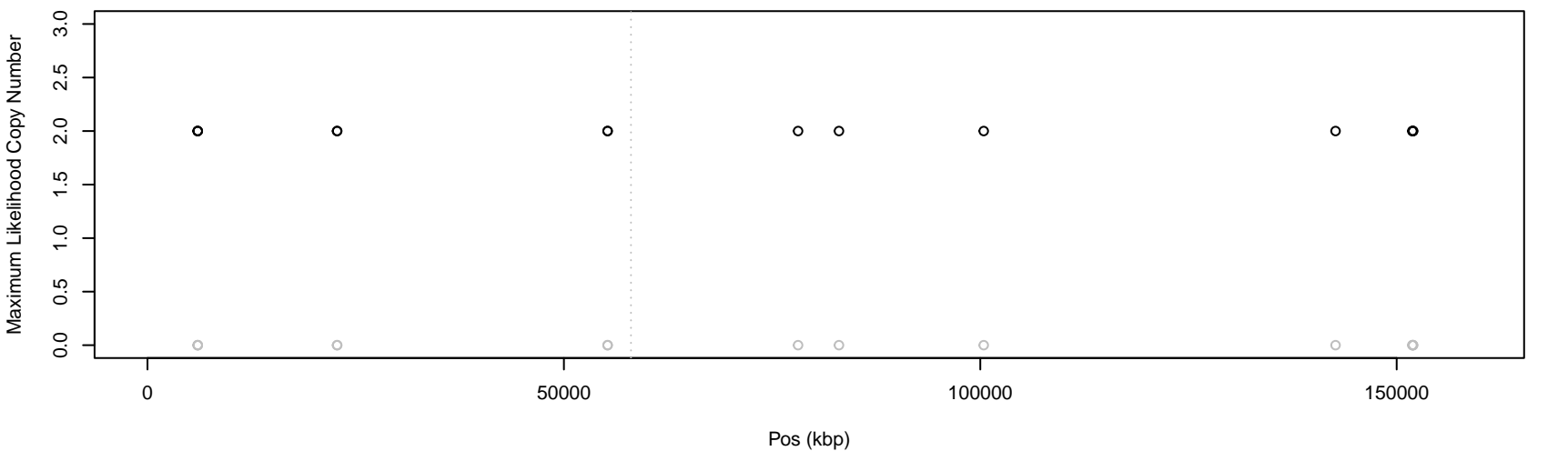
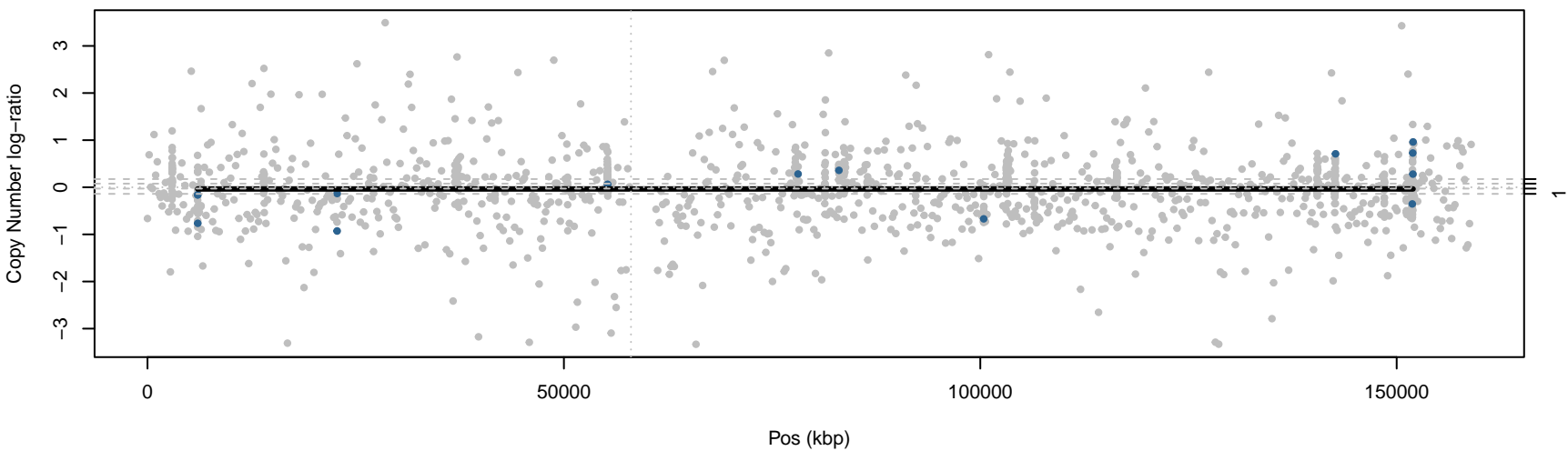
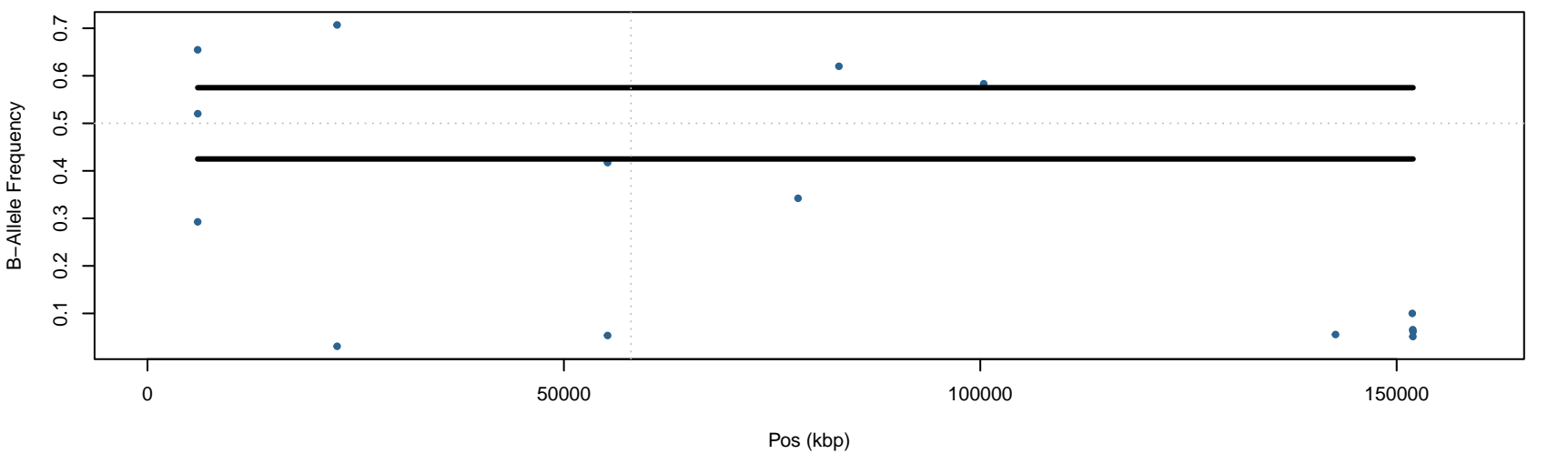
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 5



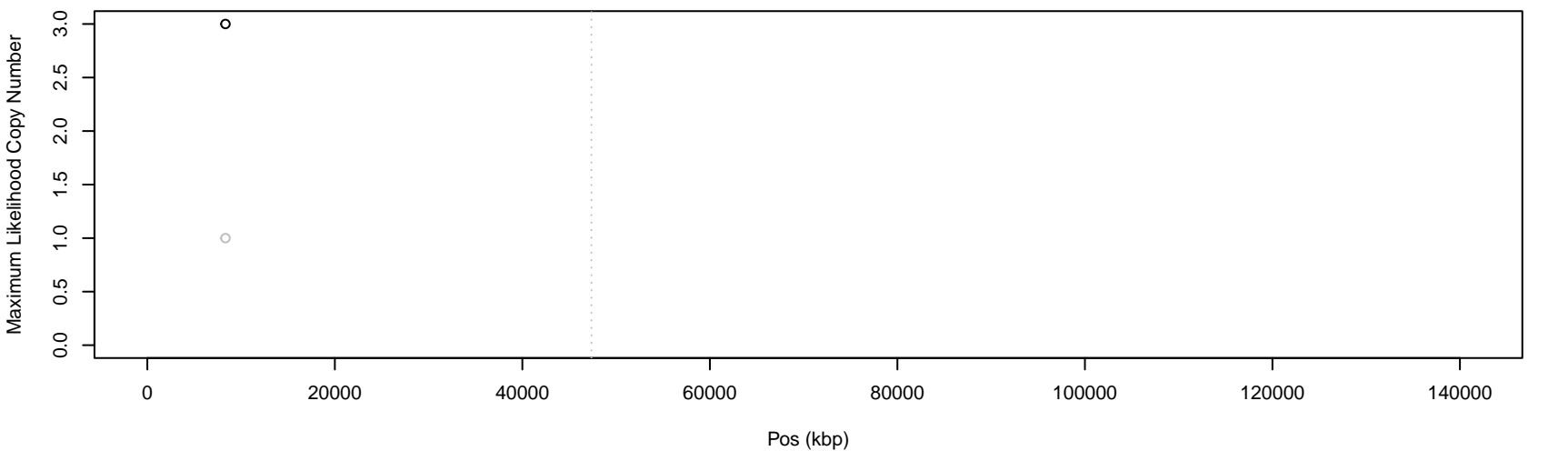
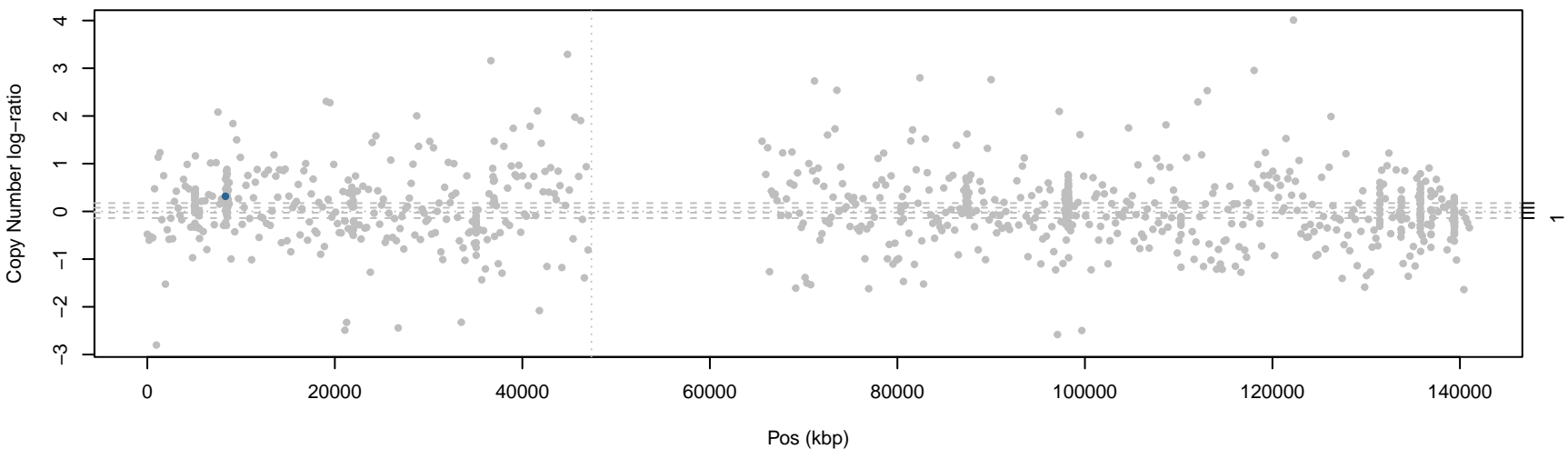
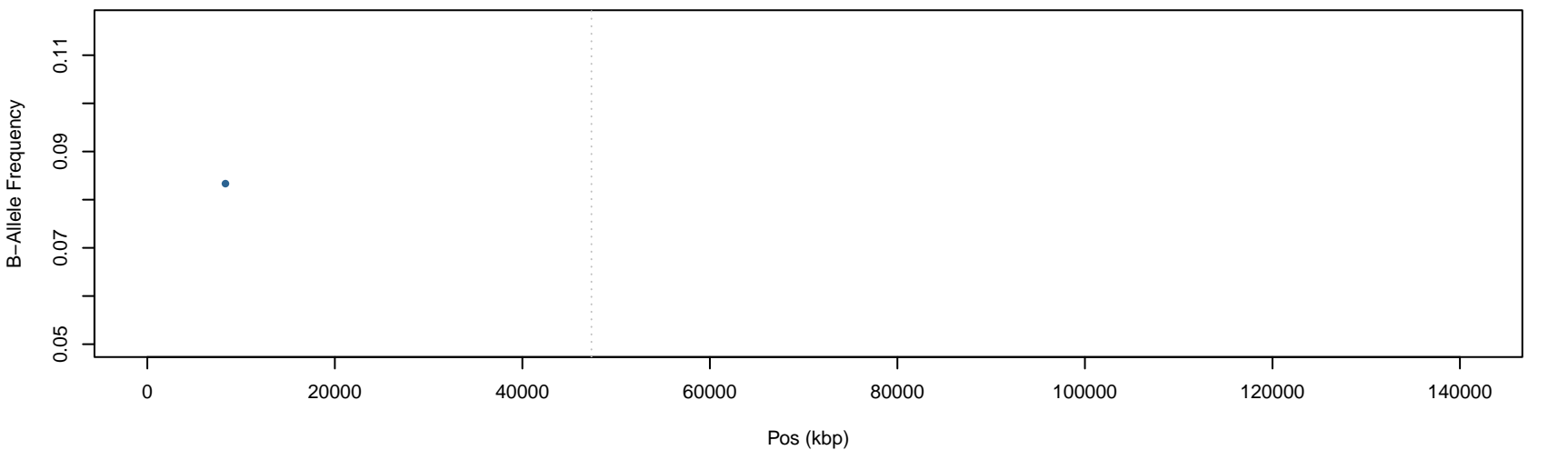
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 6



Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 7

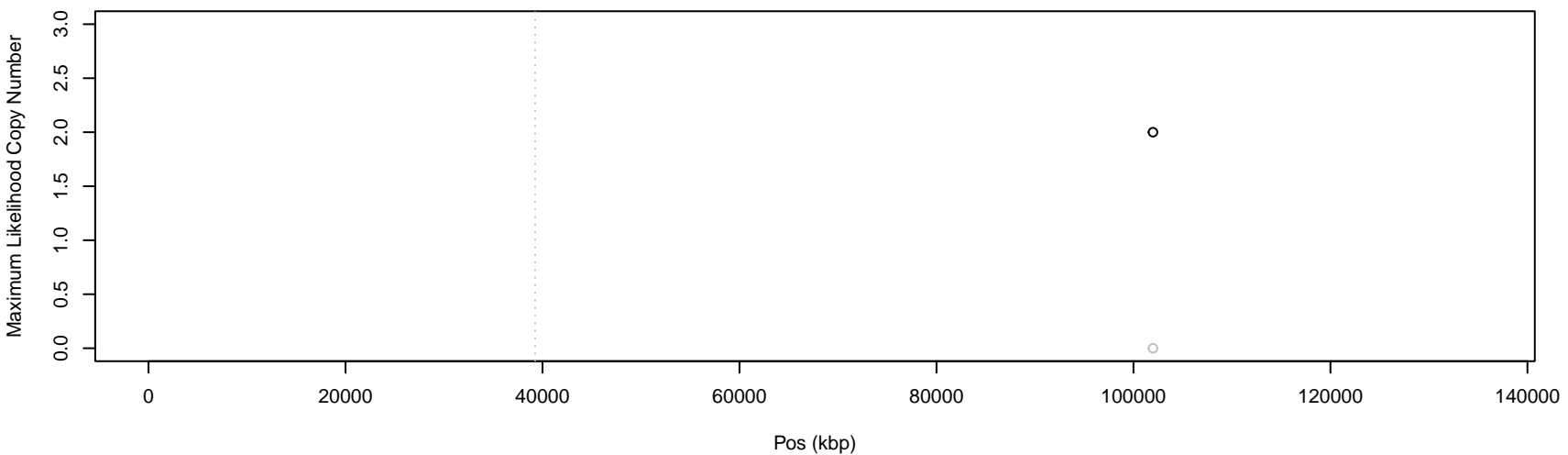
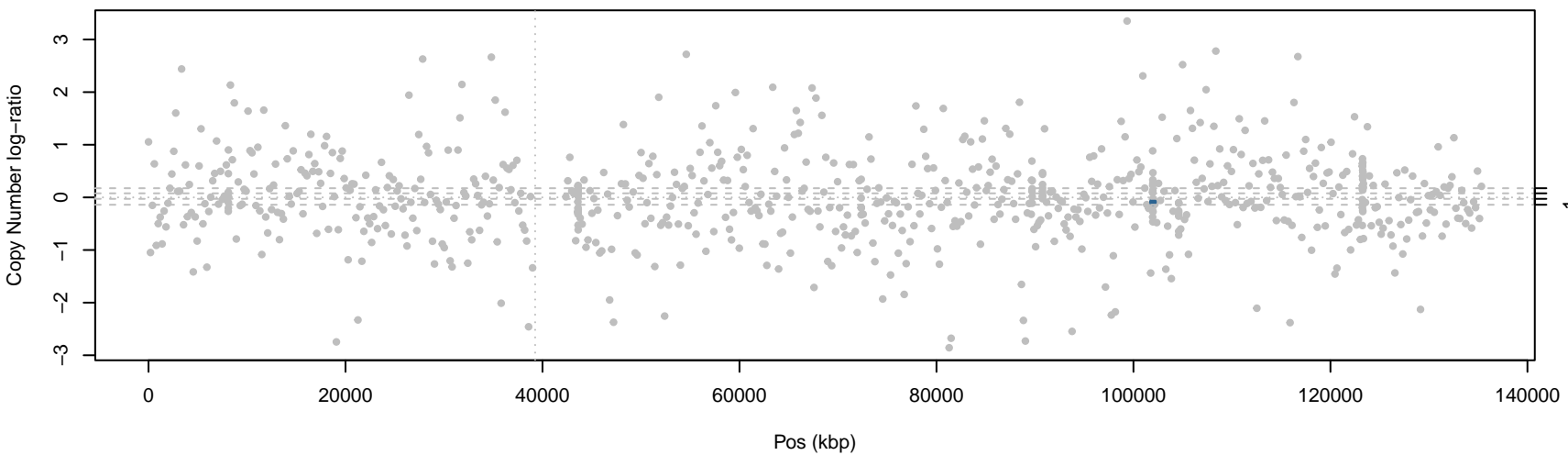
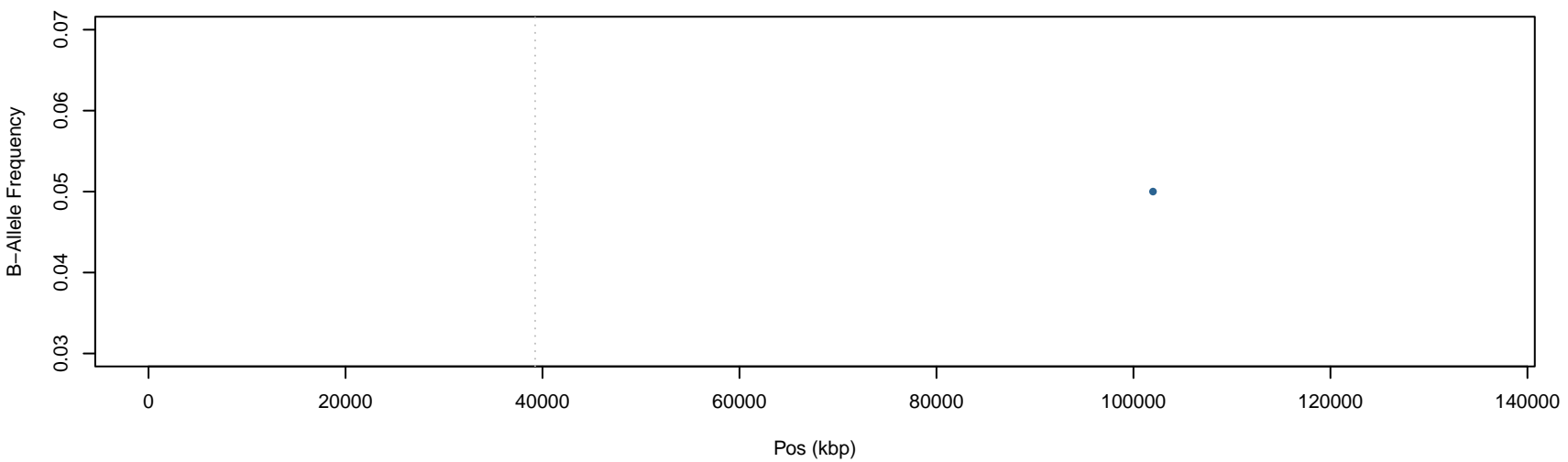


Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 9

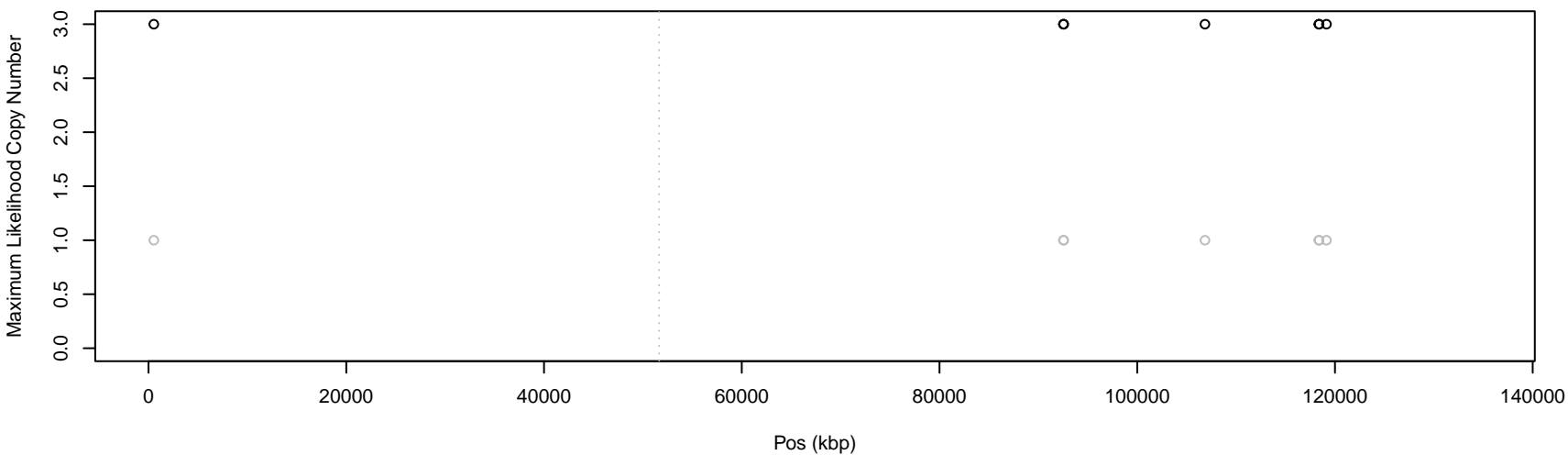
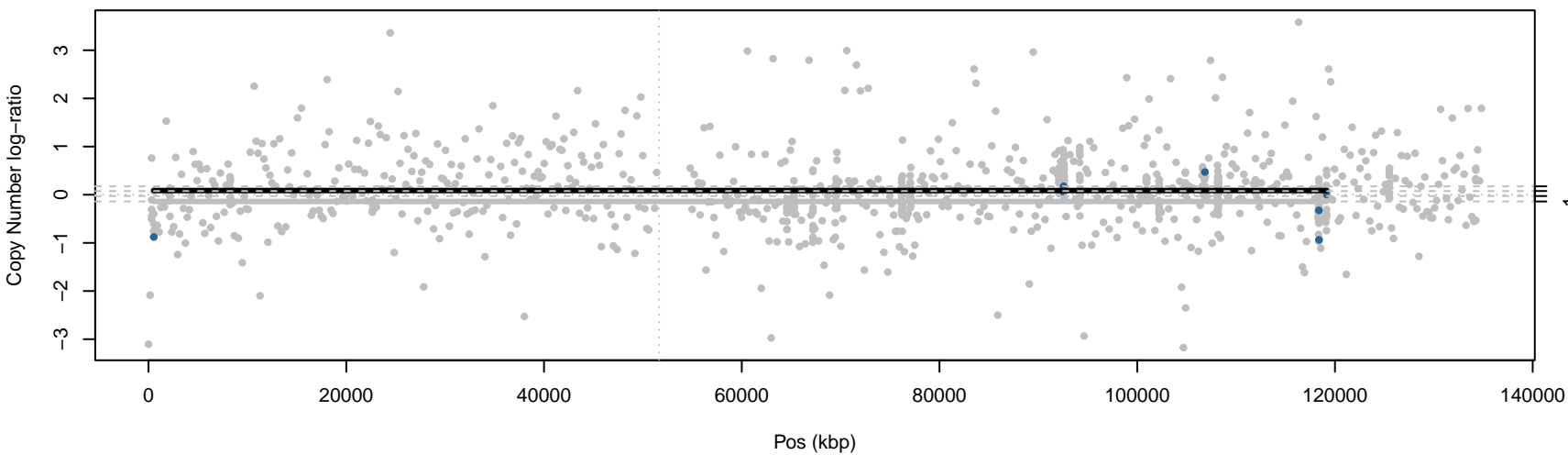
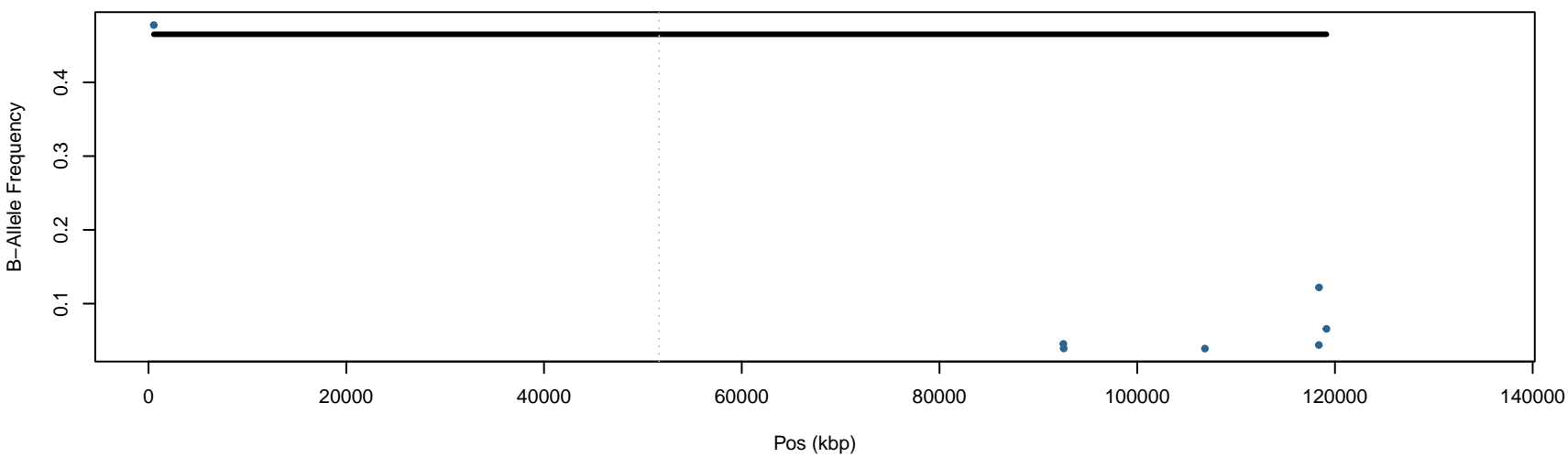




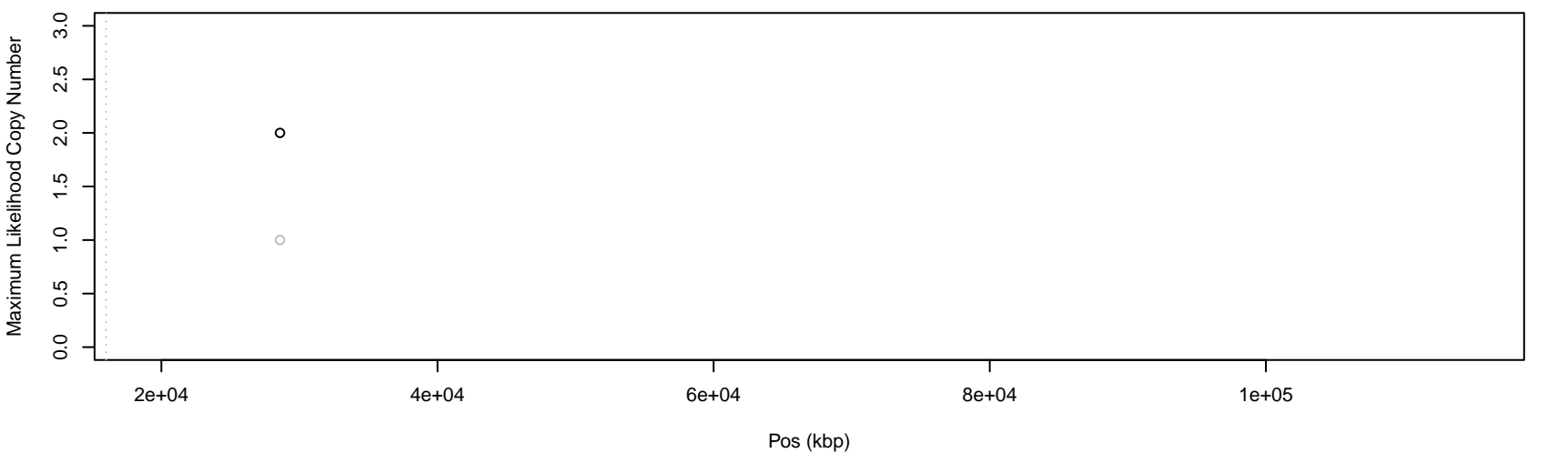
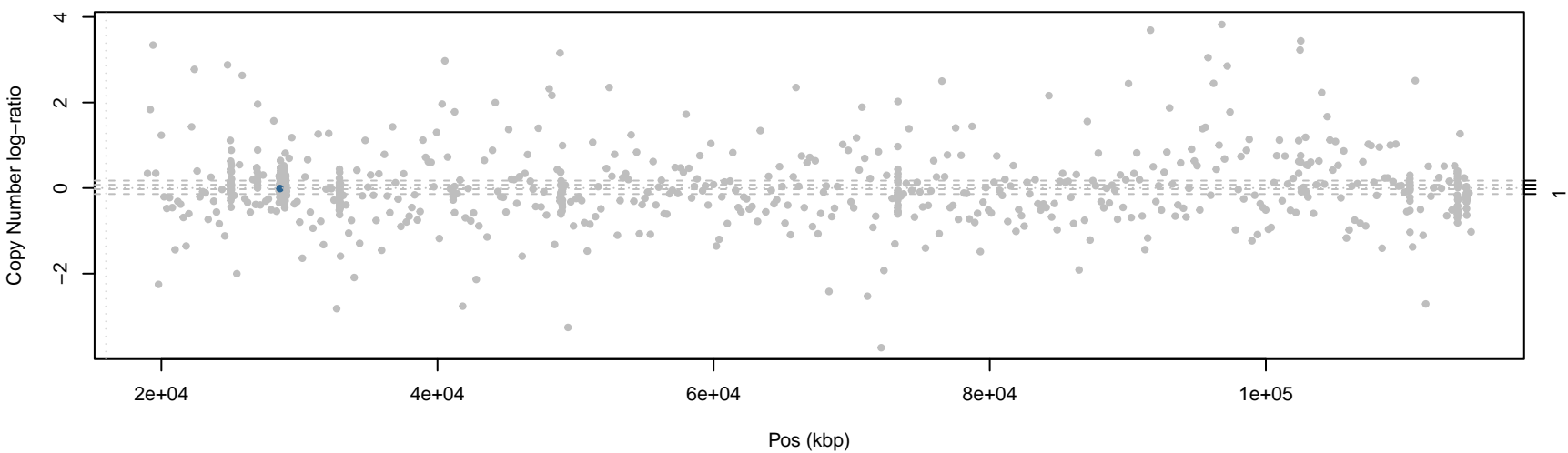
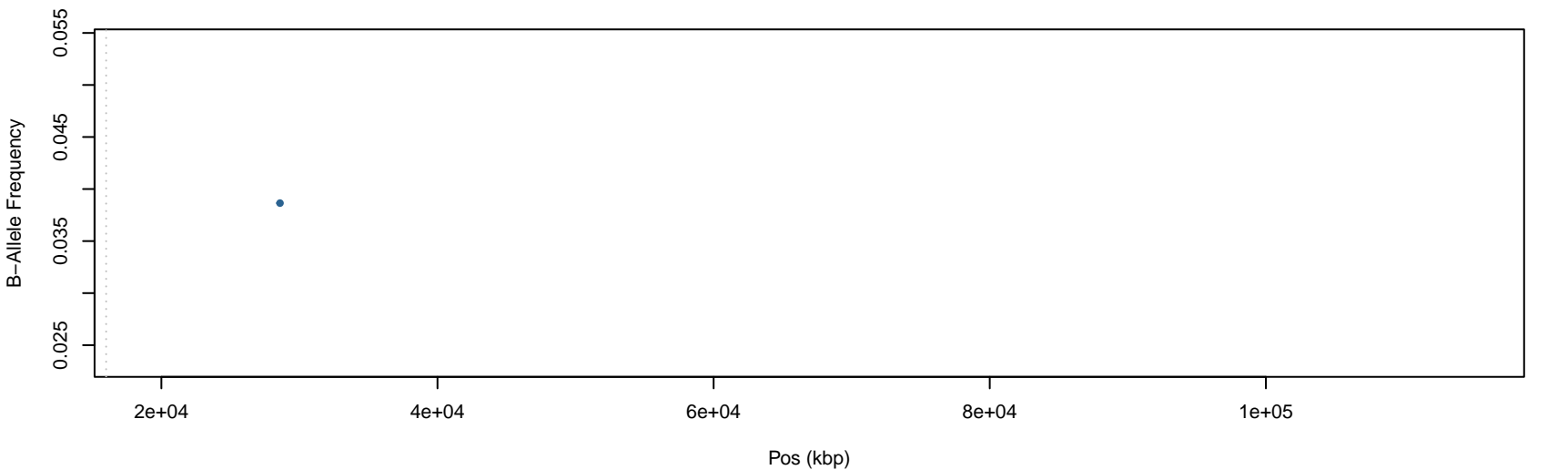
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 10



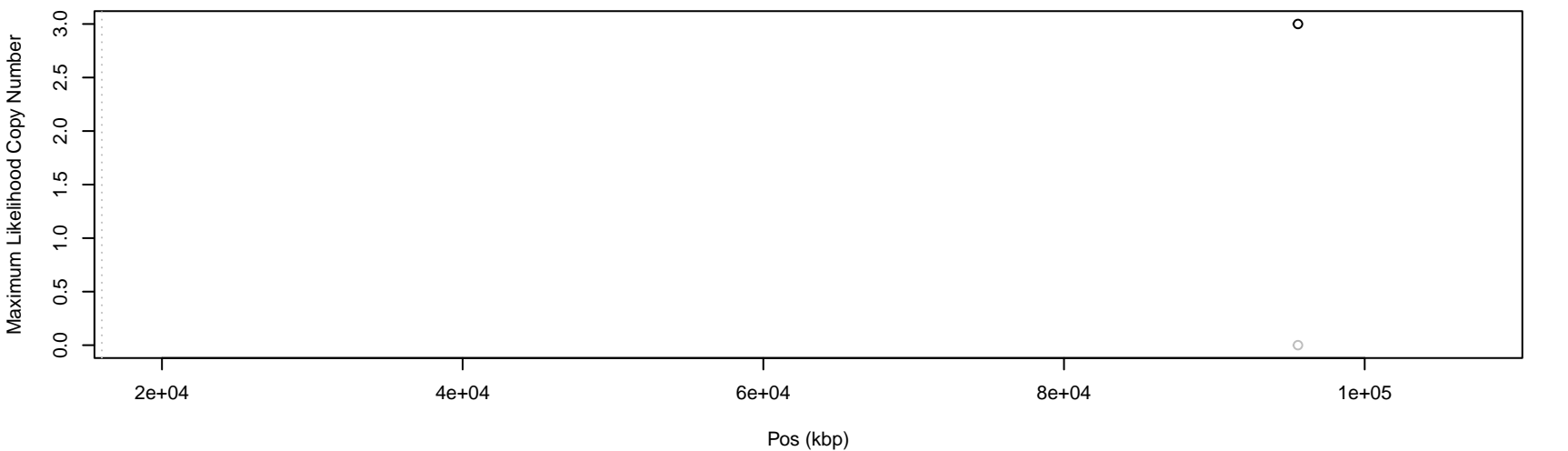
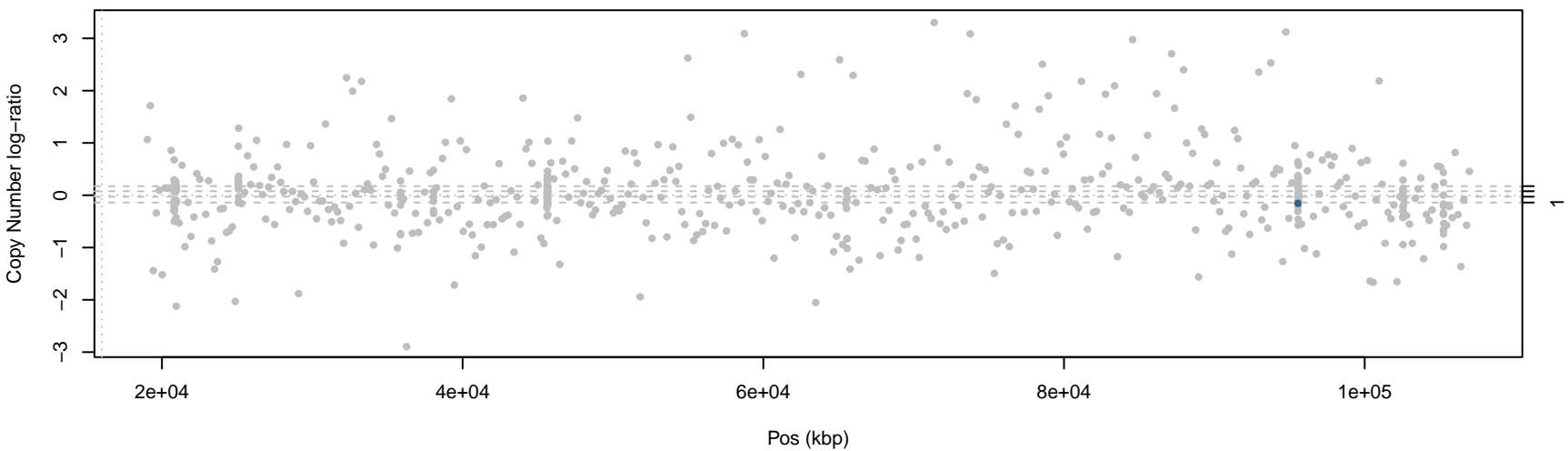
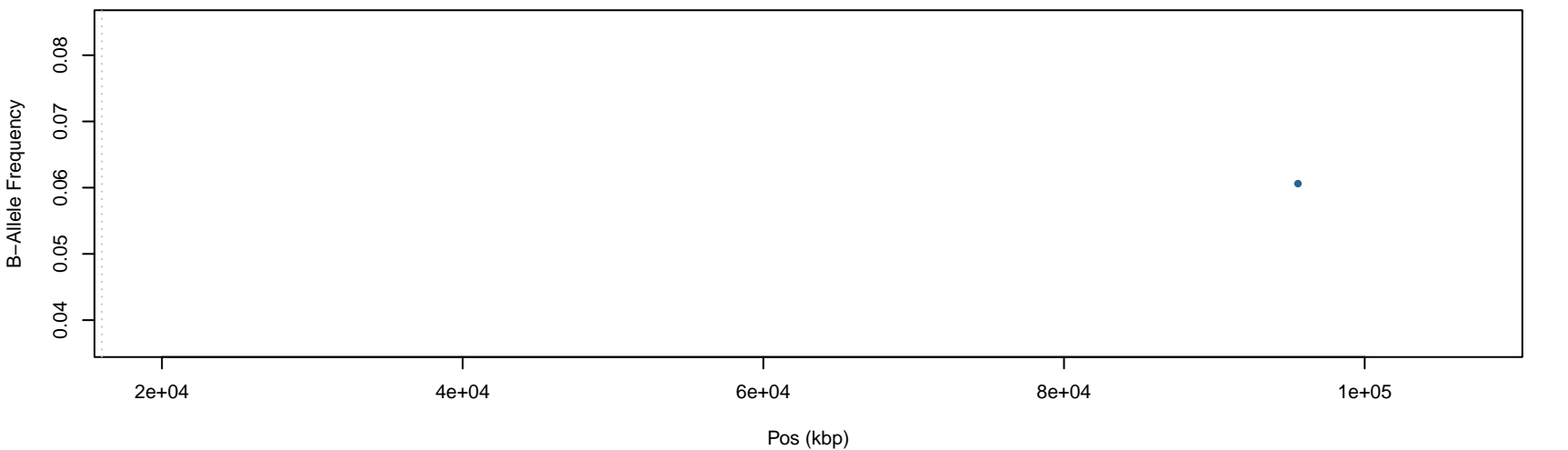
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 11



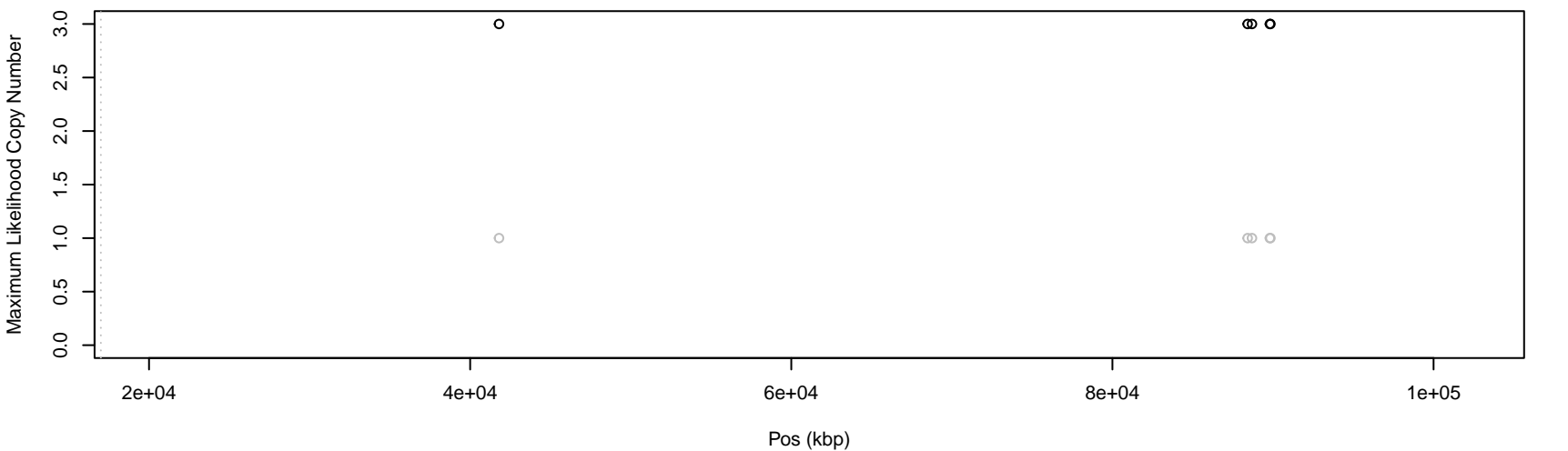
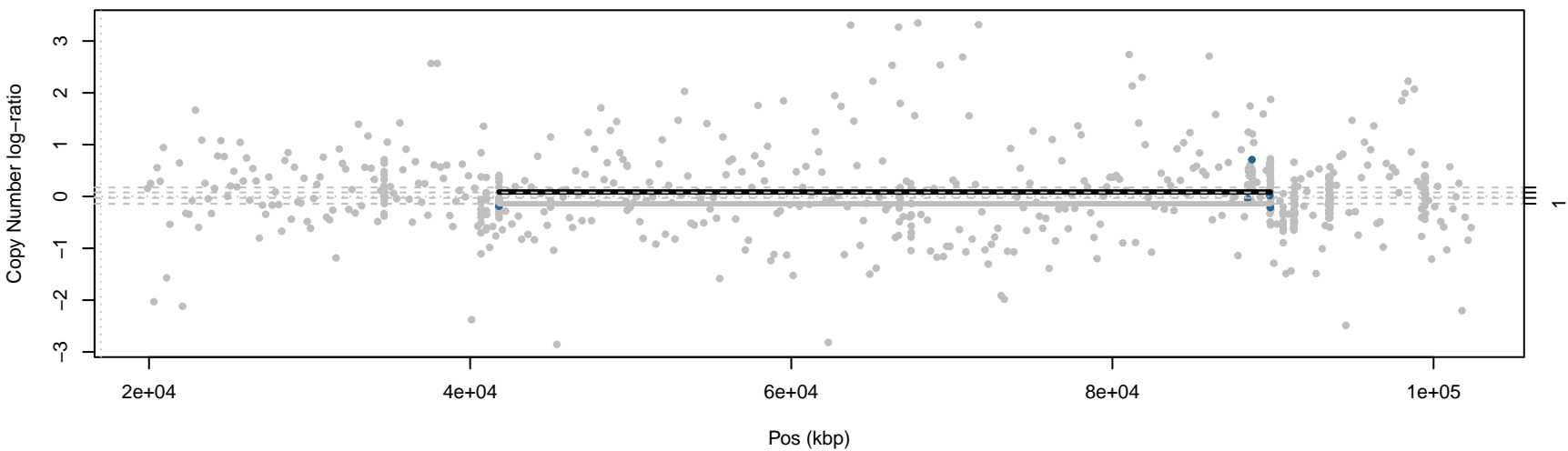
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 13



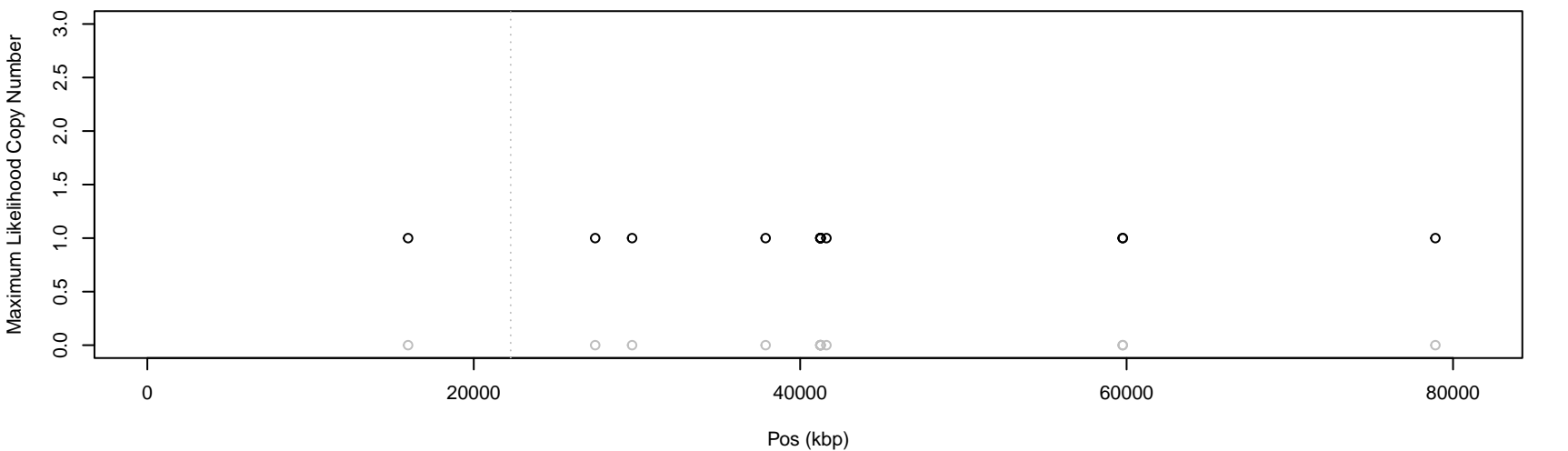
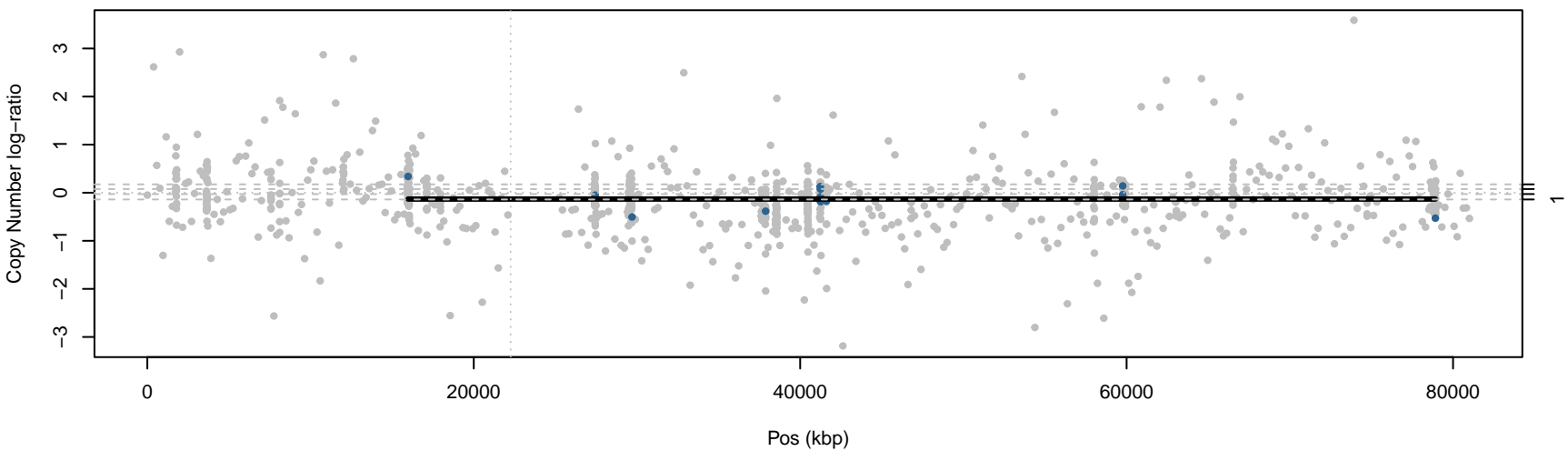
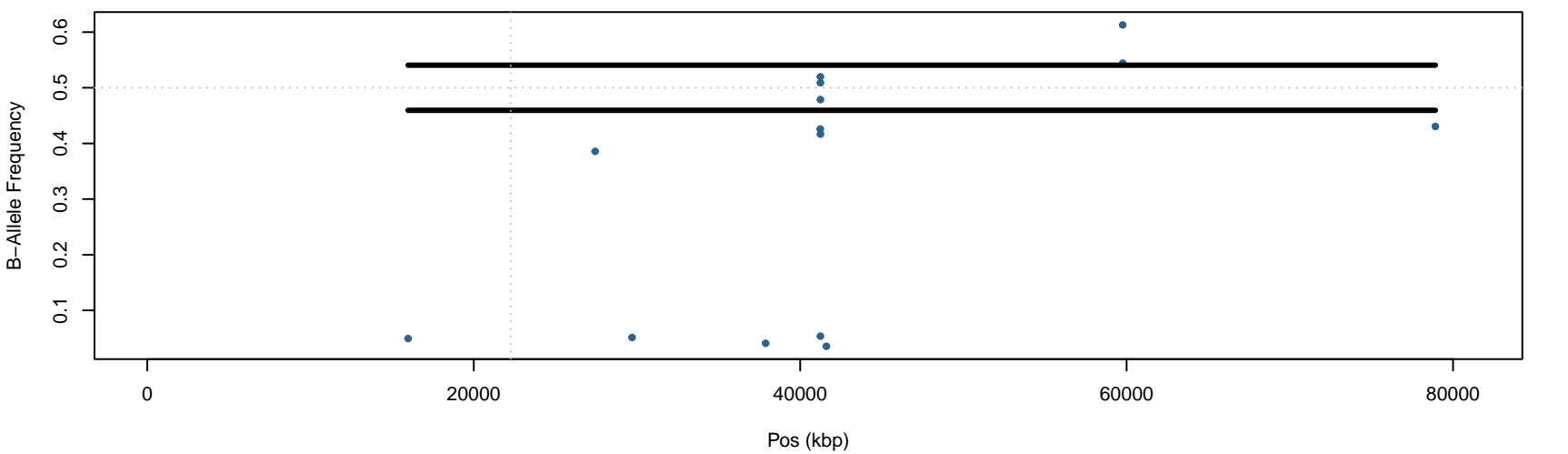
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 14



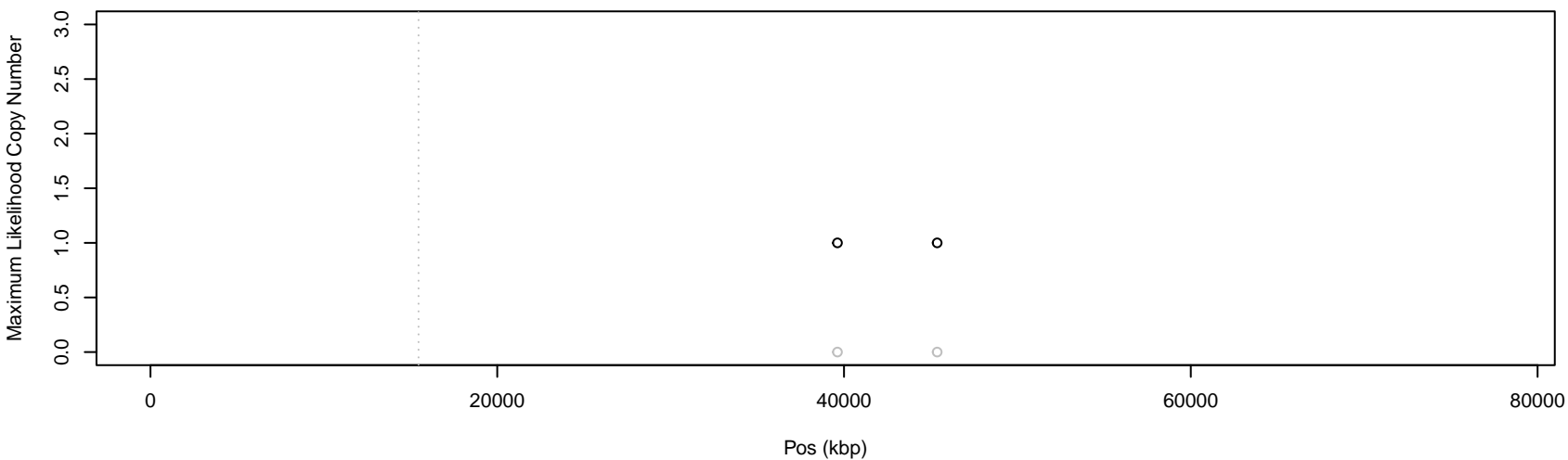
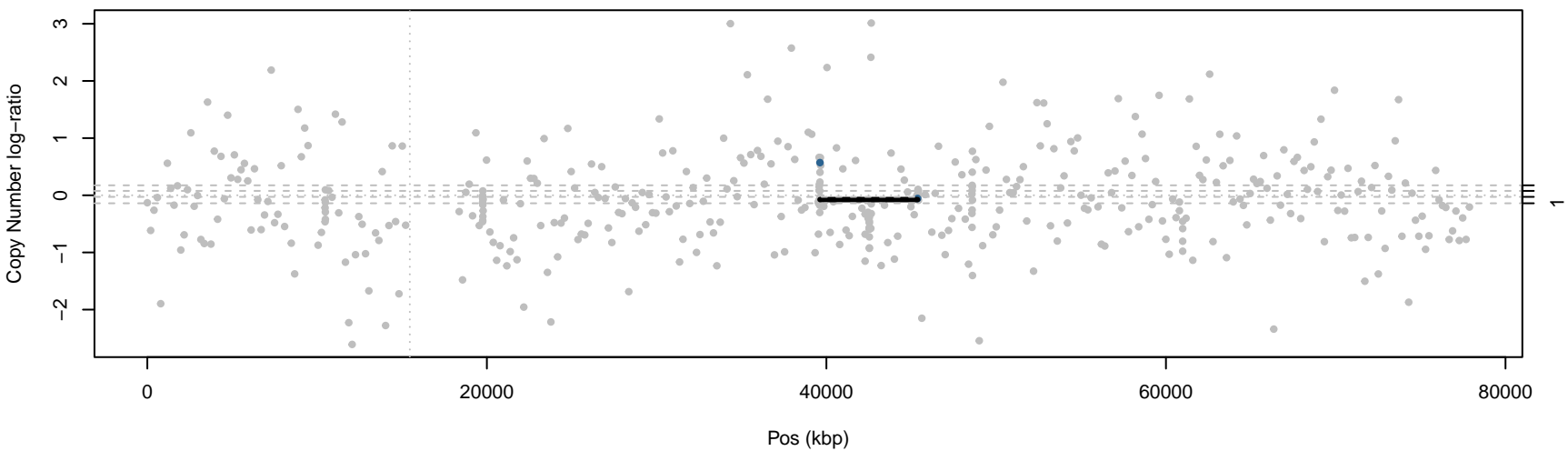
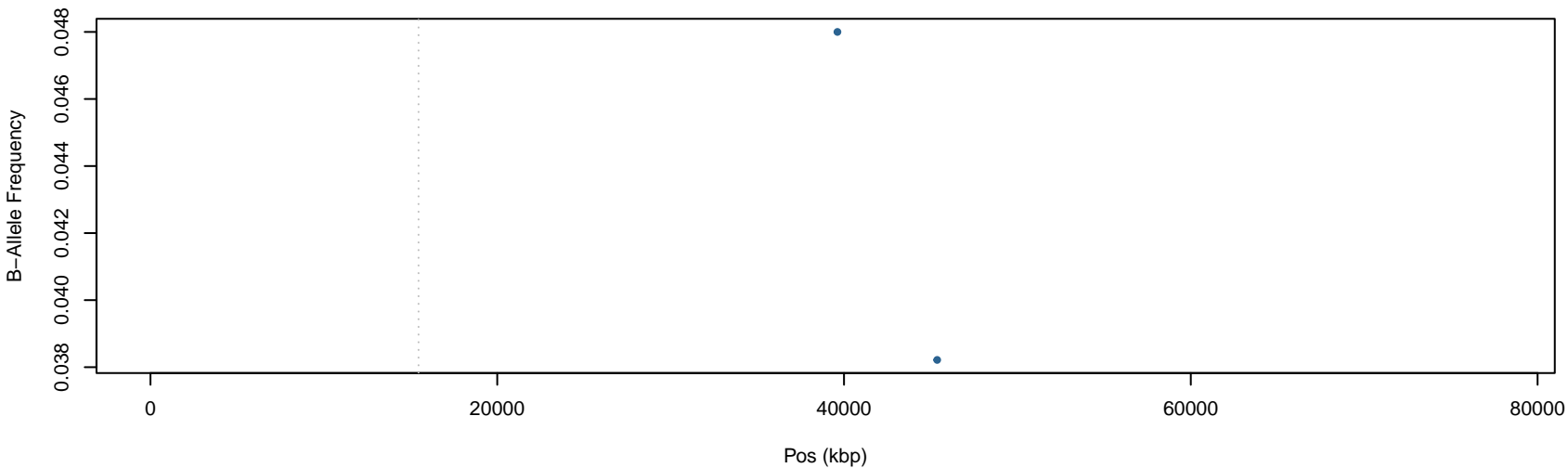
Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 15



Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 17



Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 18



Purity: 0.15 Tumor ploidy: 2.256 SNV log-likelihood: 51.53 GoF: 95.6% Mean coverage: 187;674 Chromosome: 19

