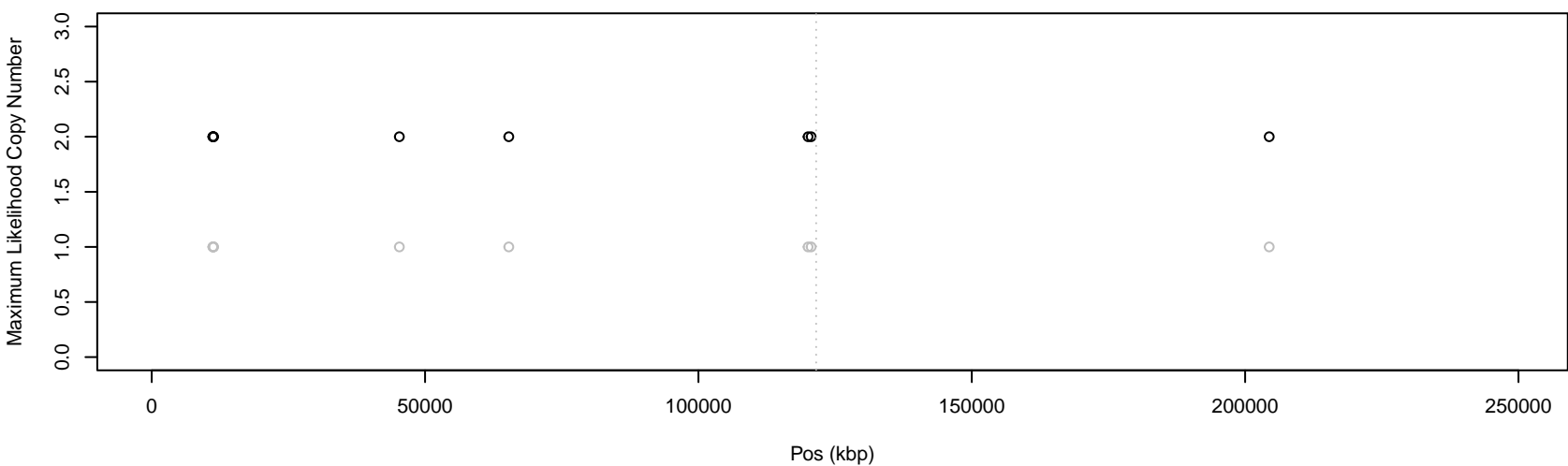
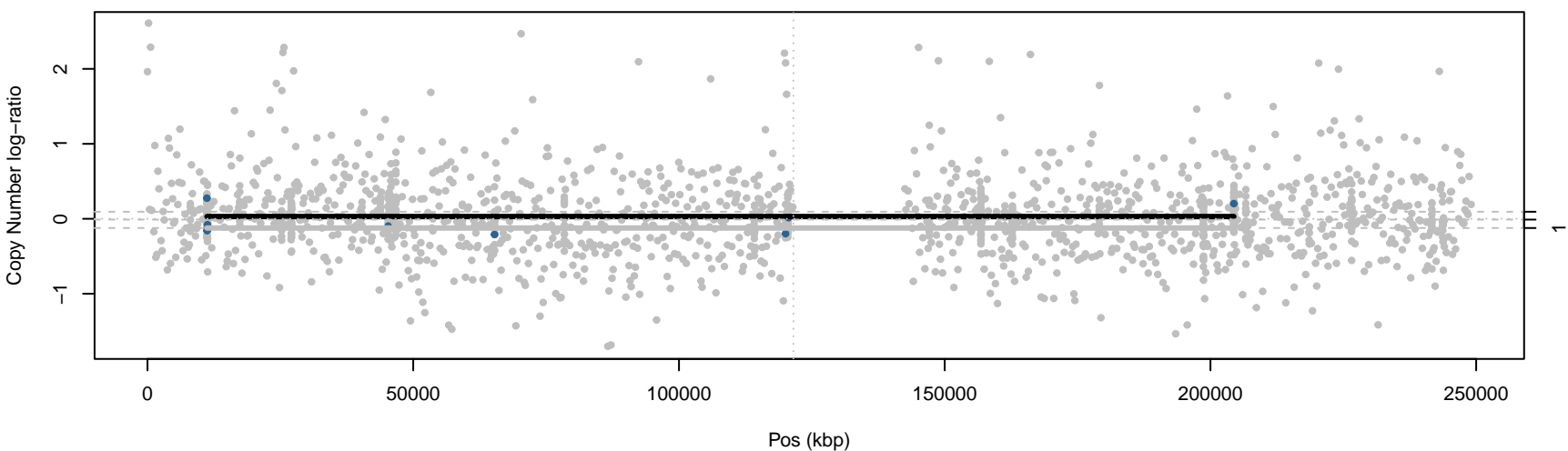
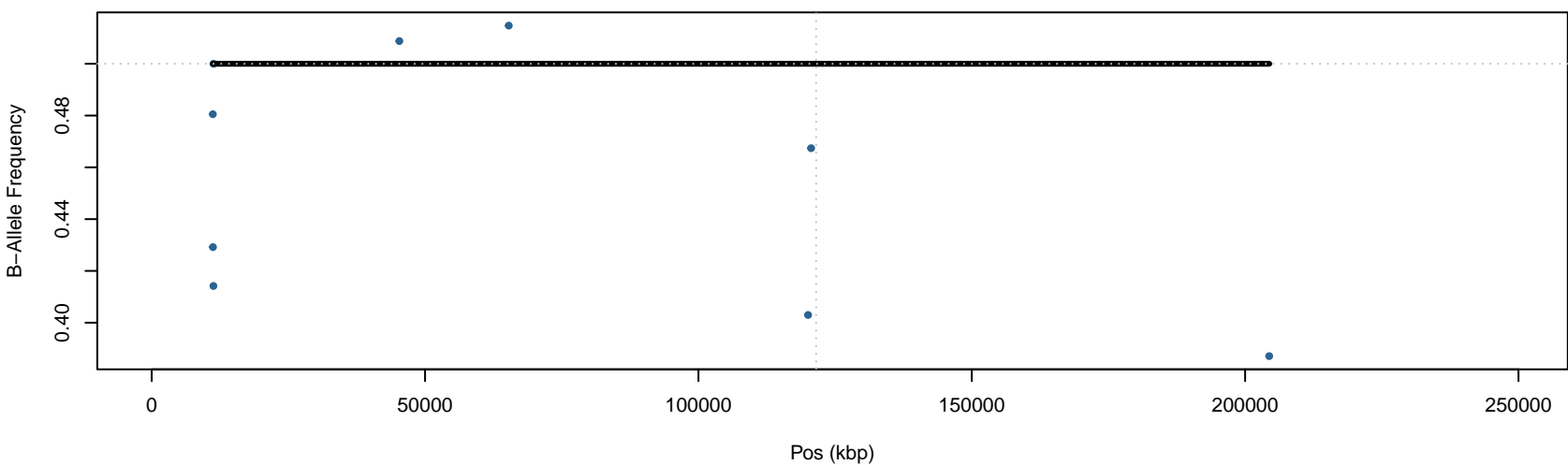
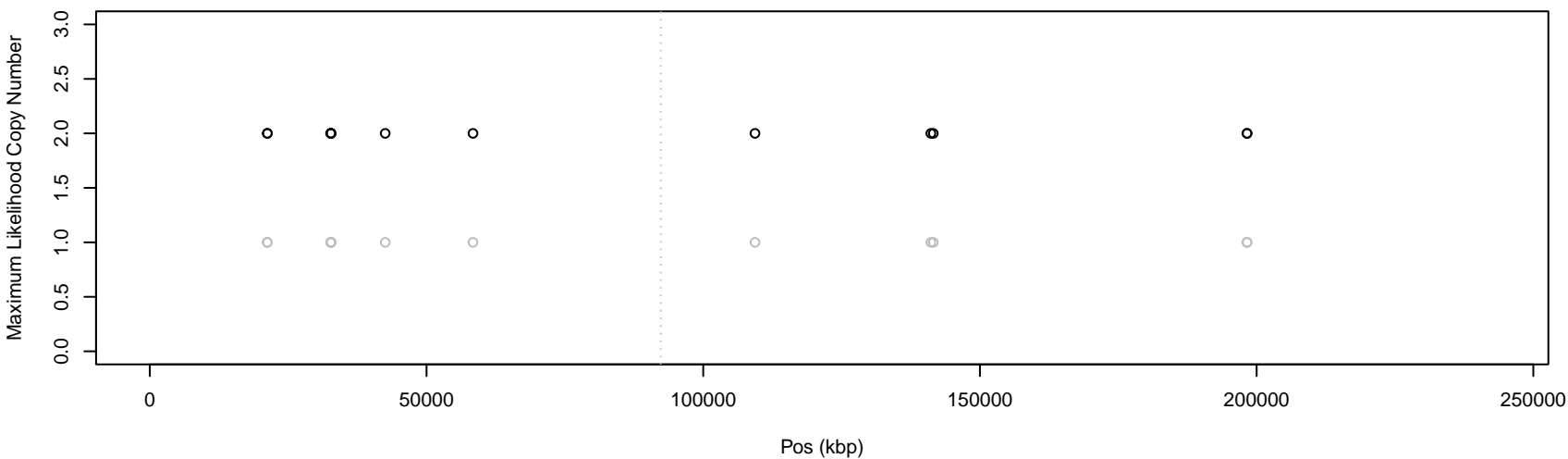
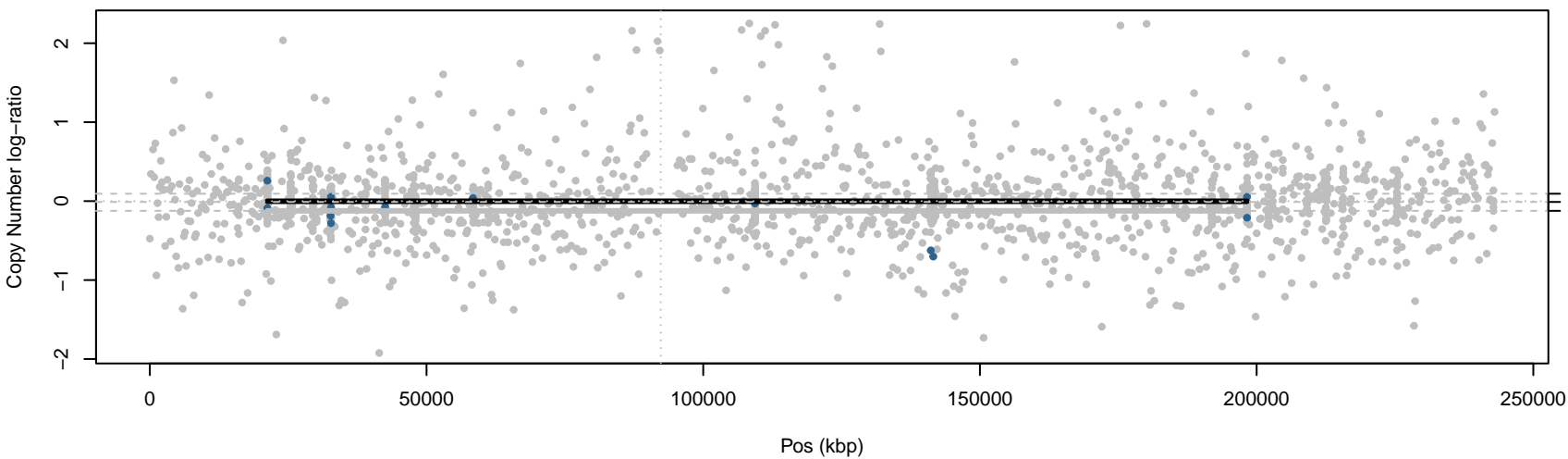
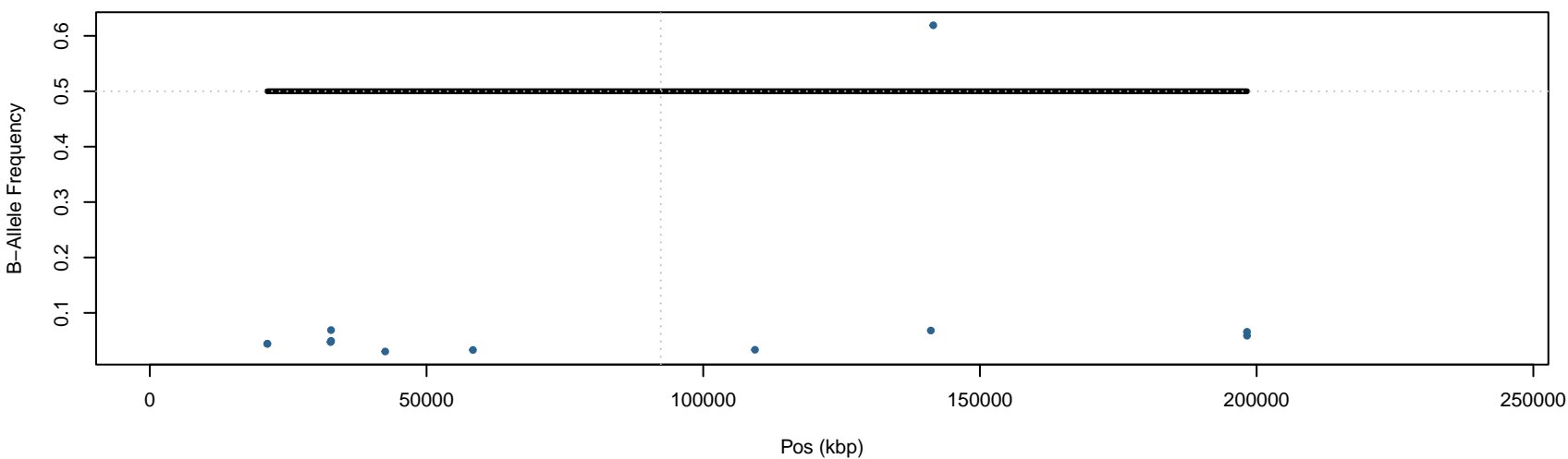


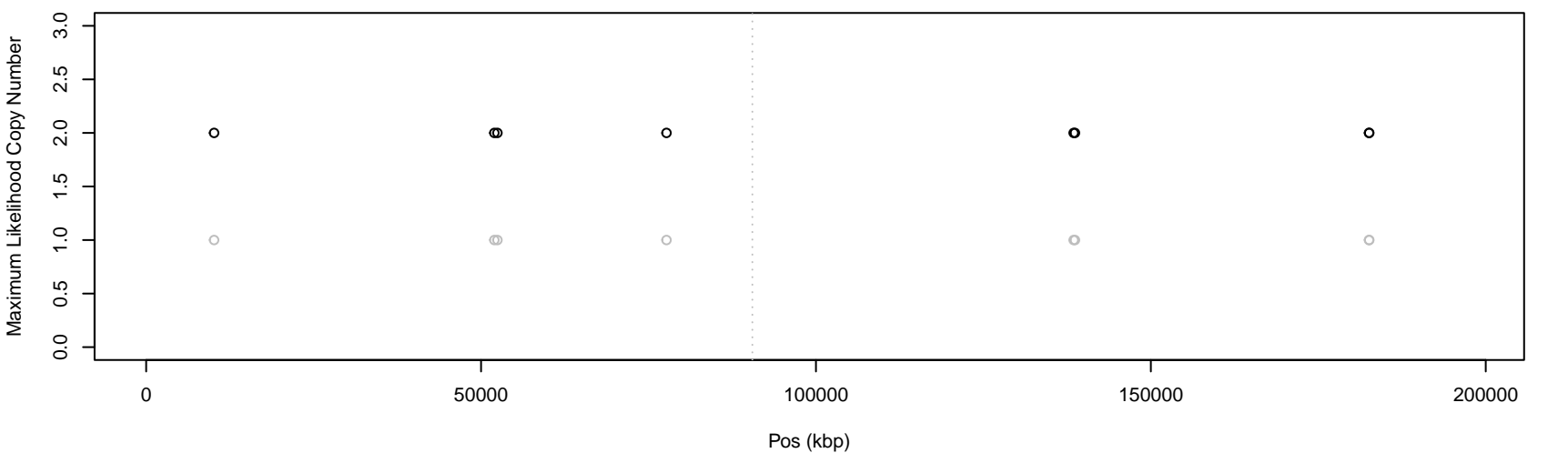
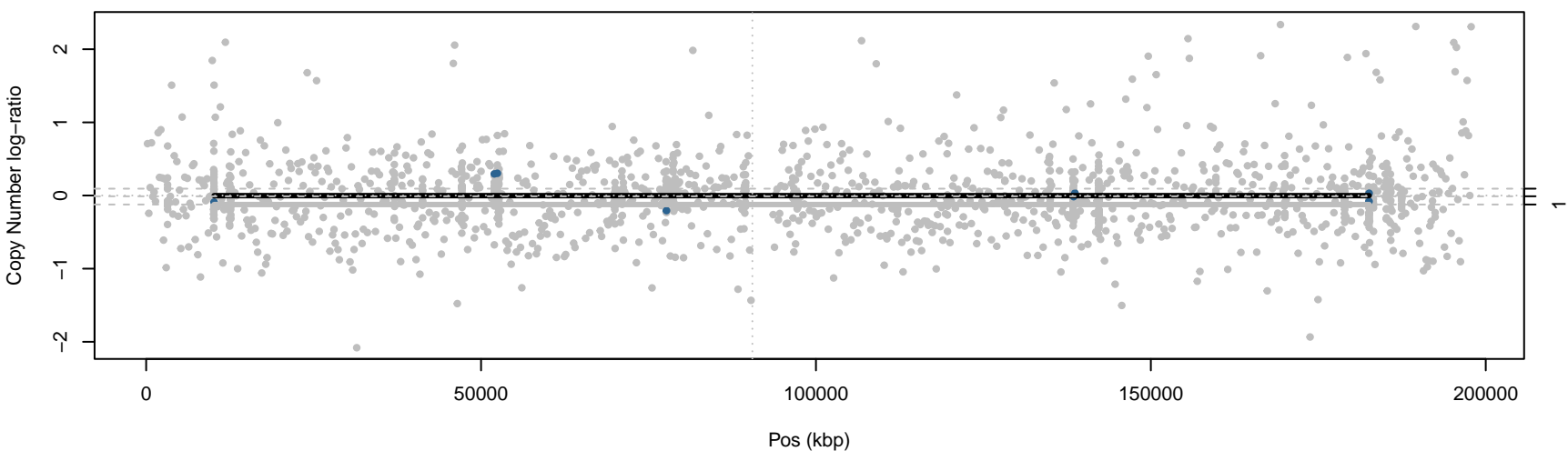
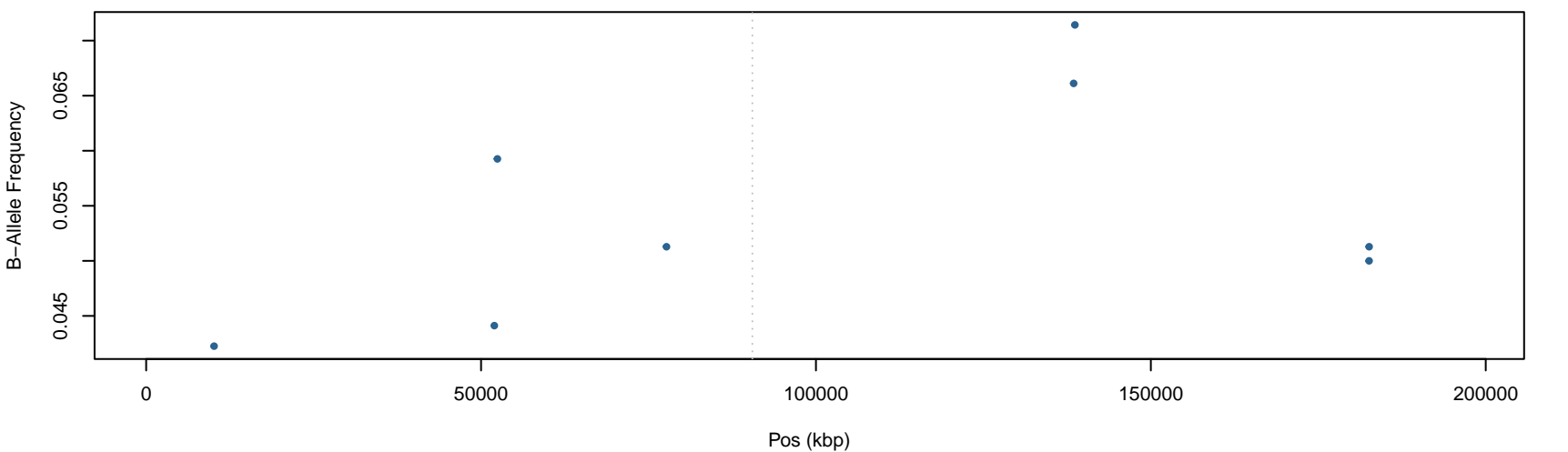
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 1



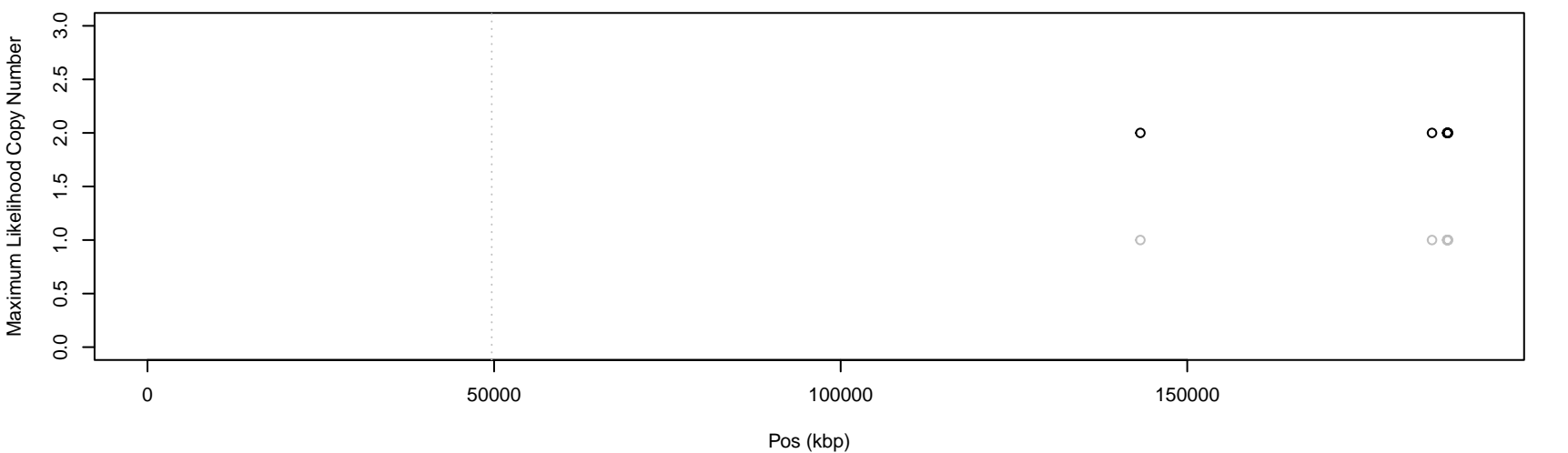
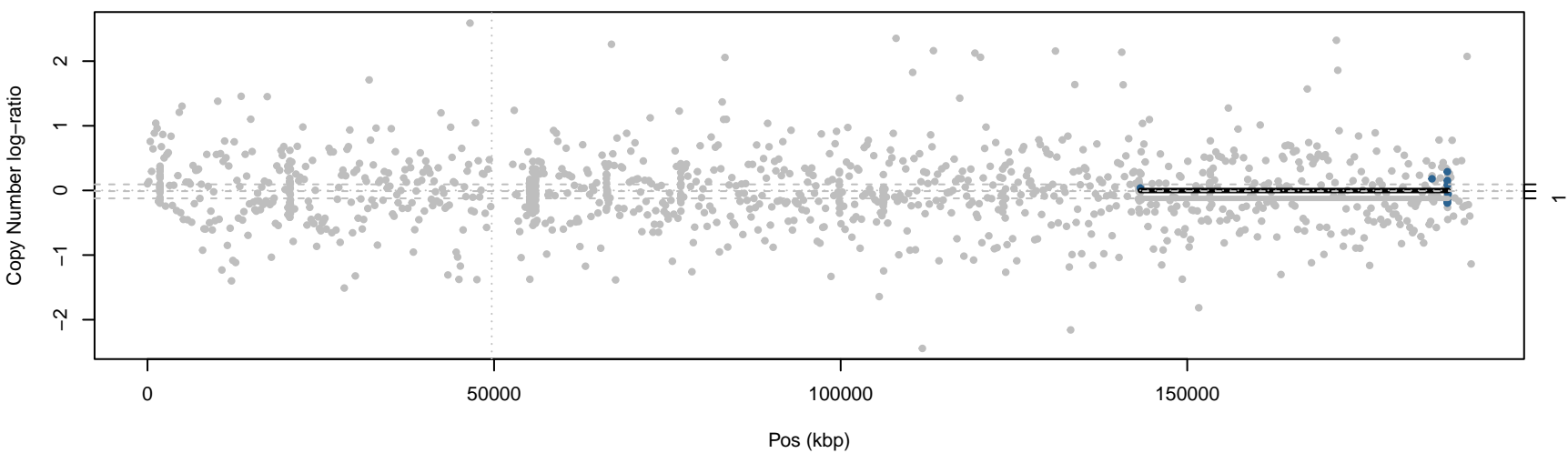
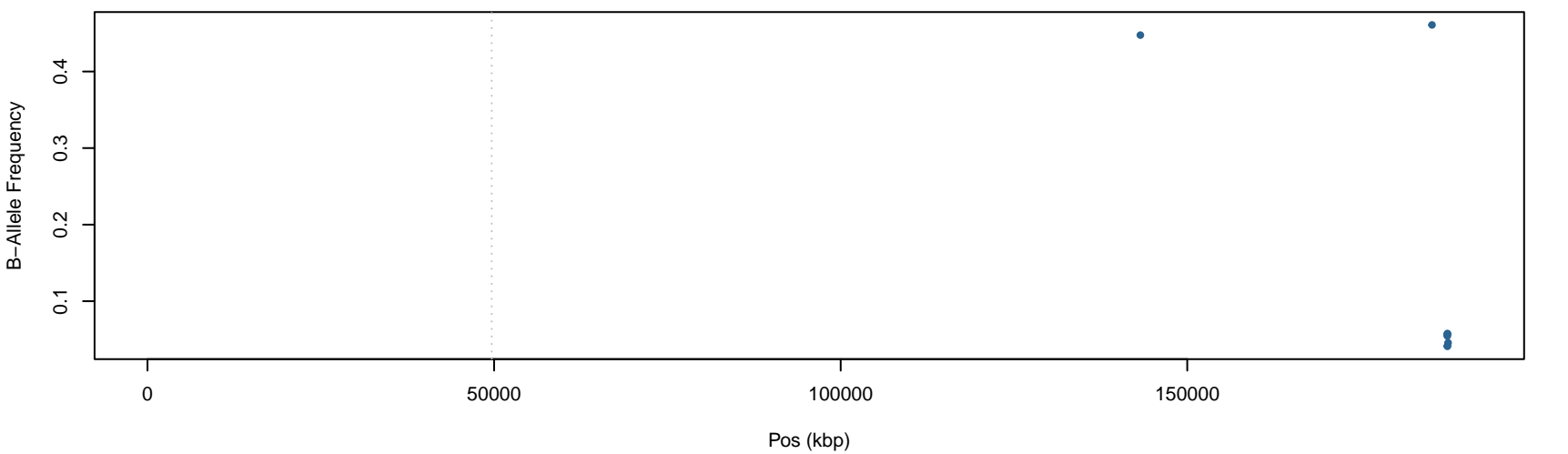
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 2



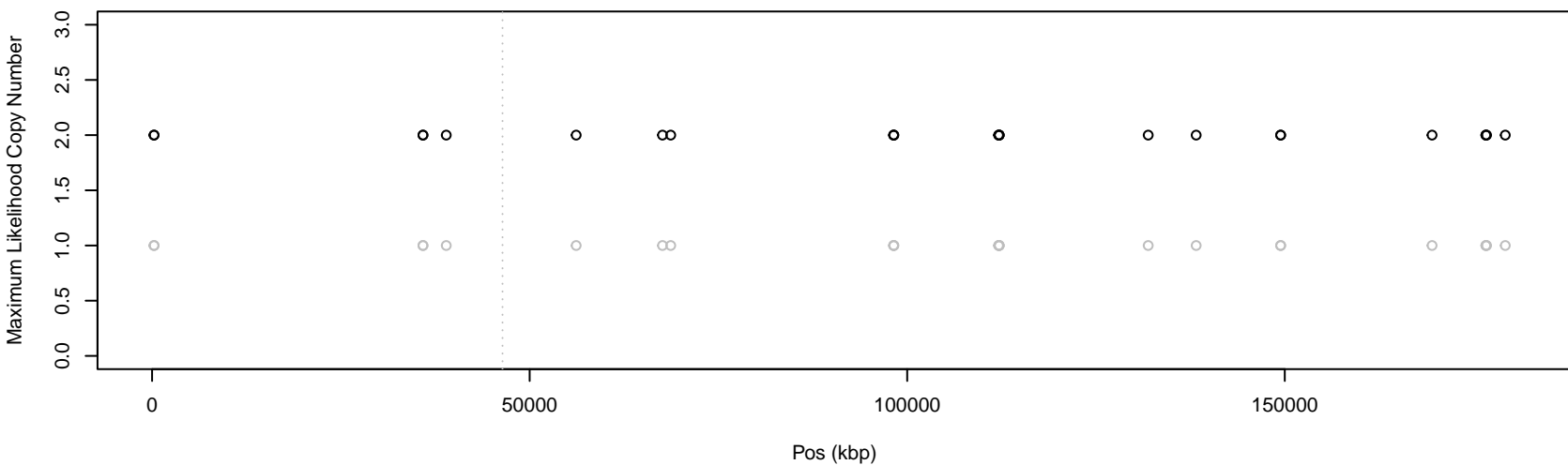
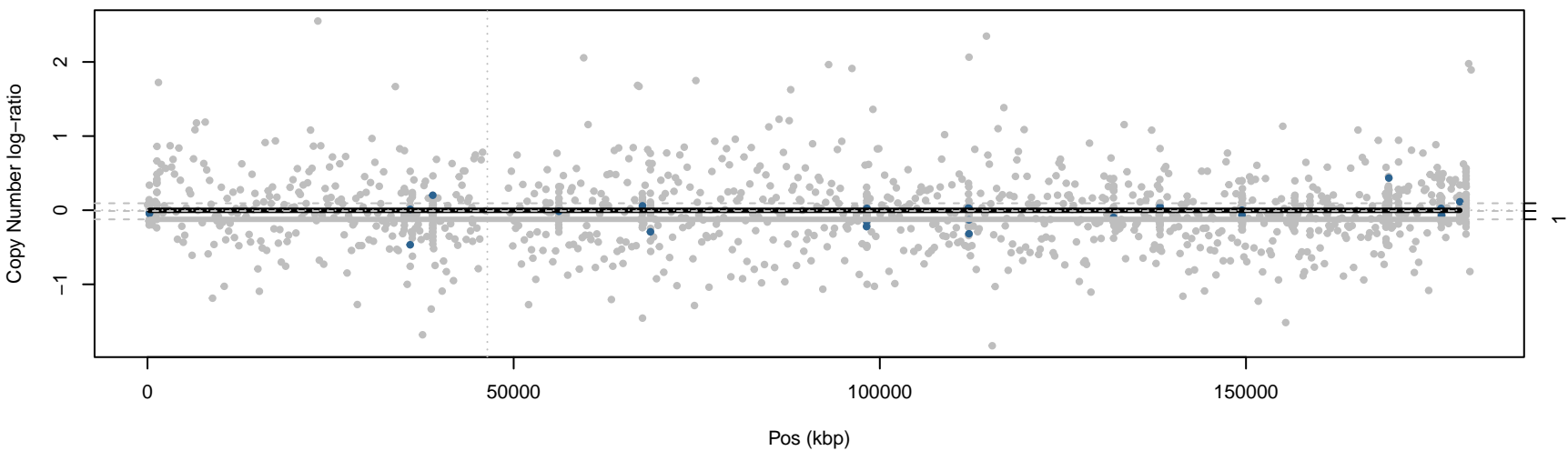
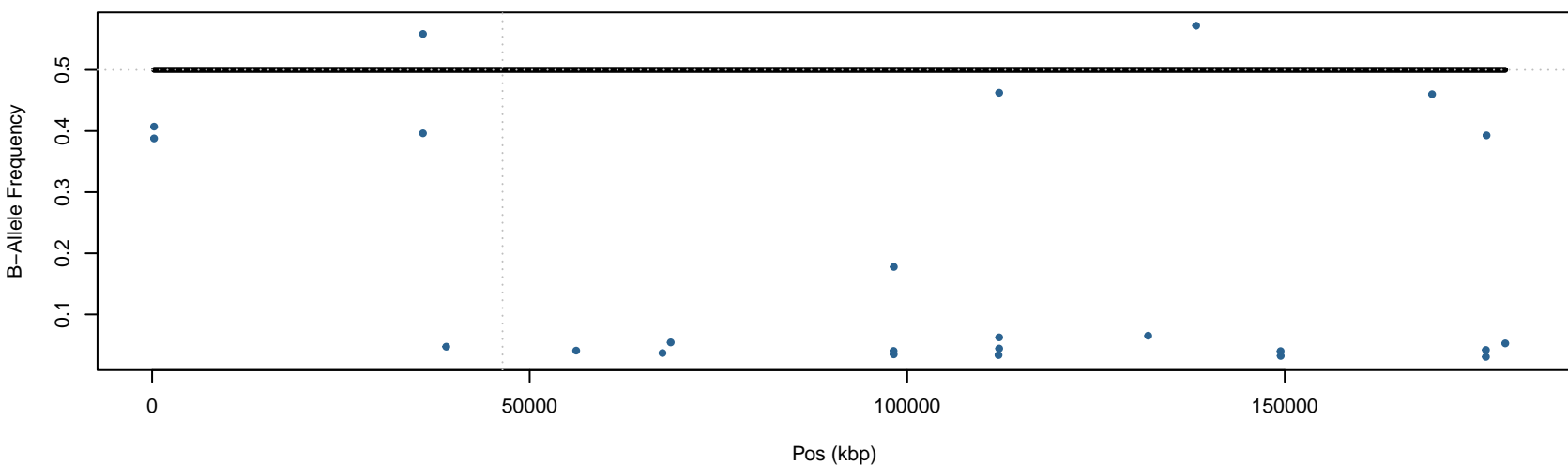
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 3



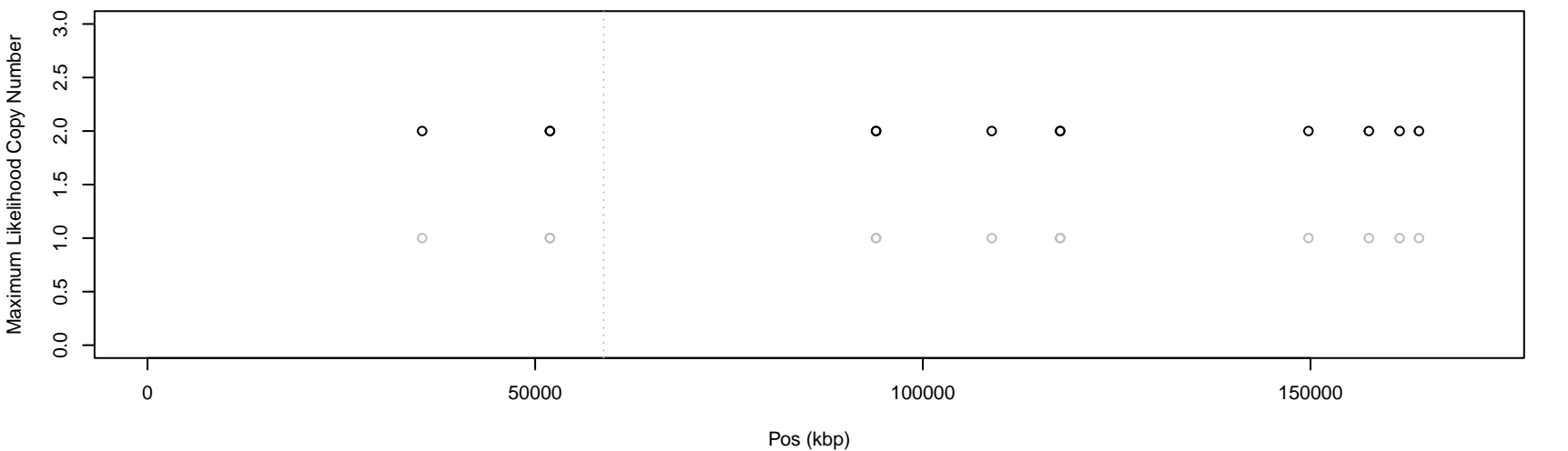
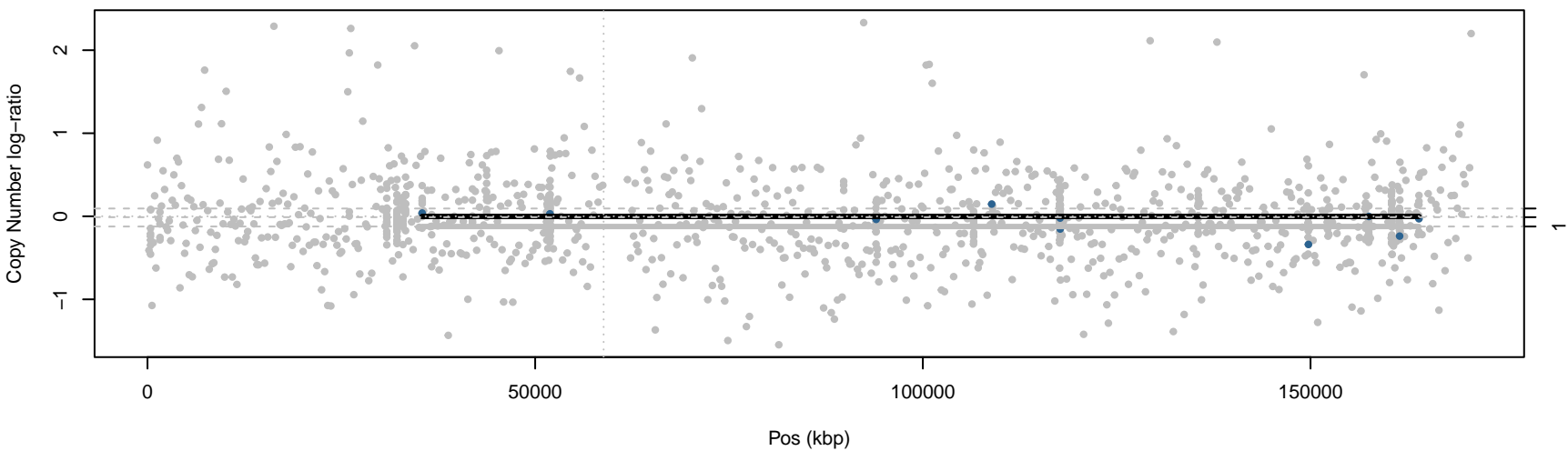
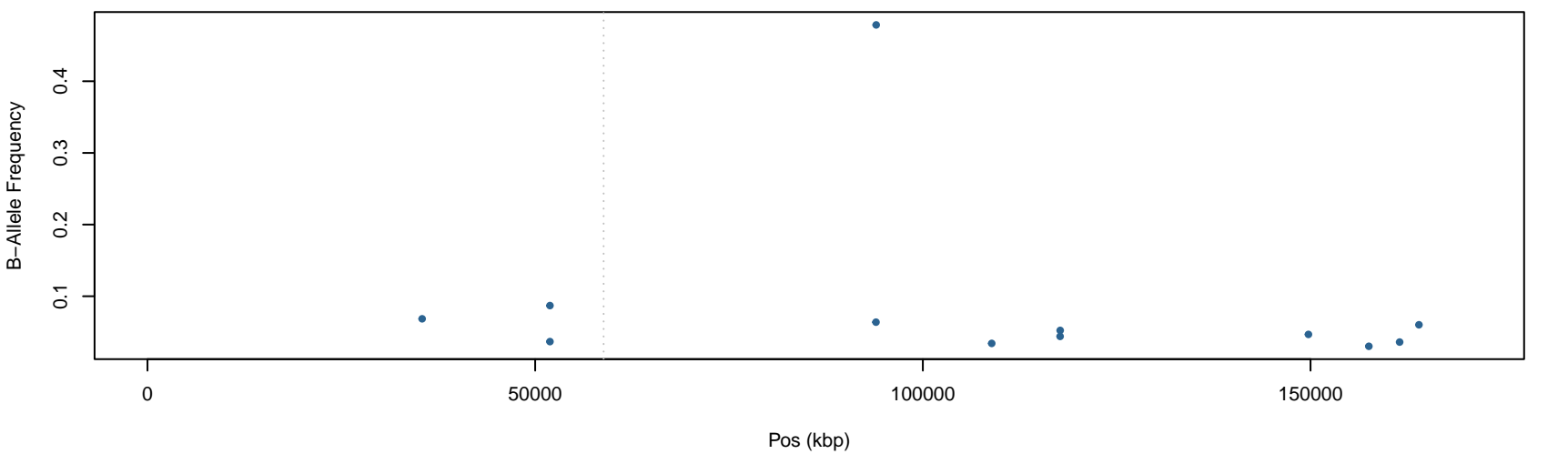
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 4



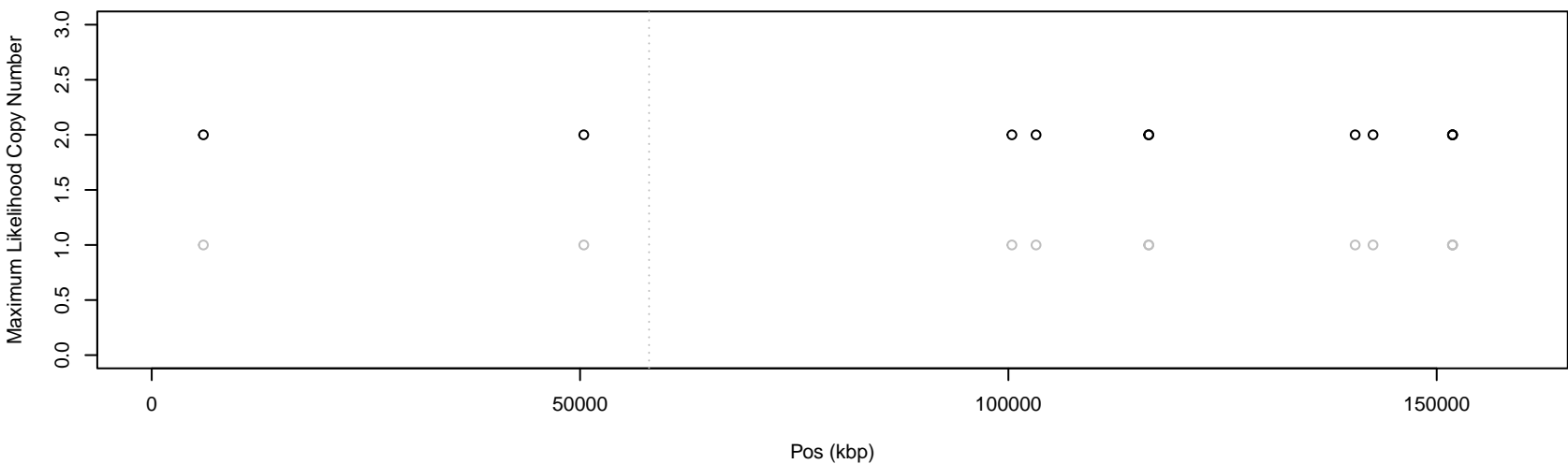
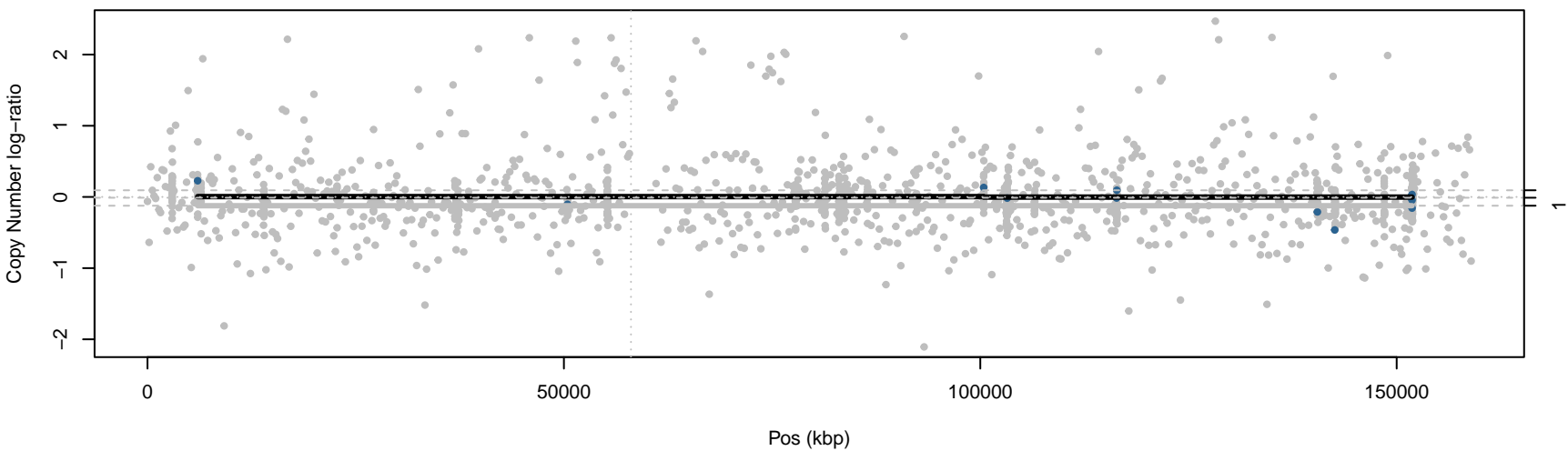
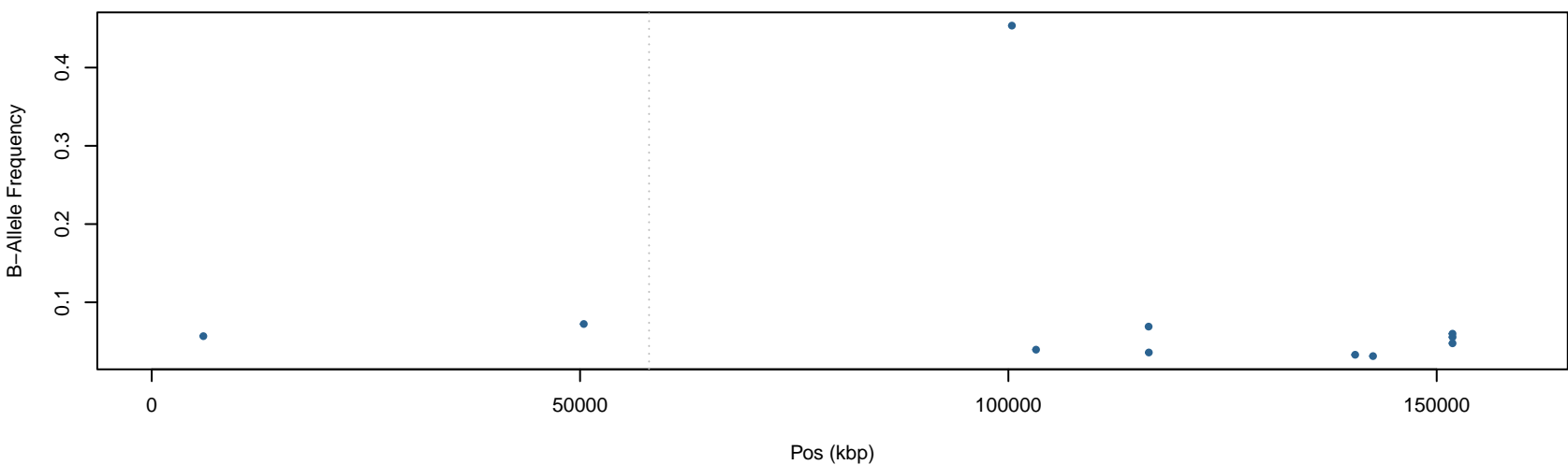
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 5



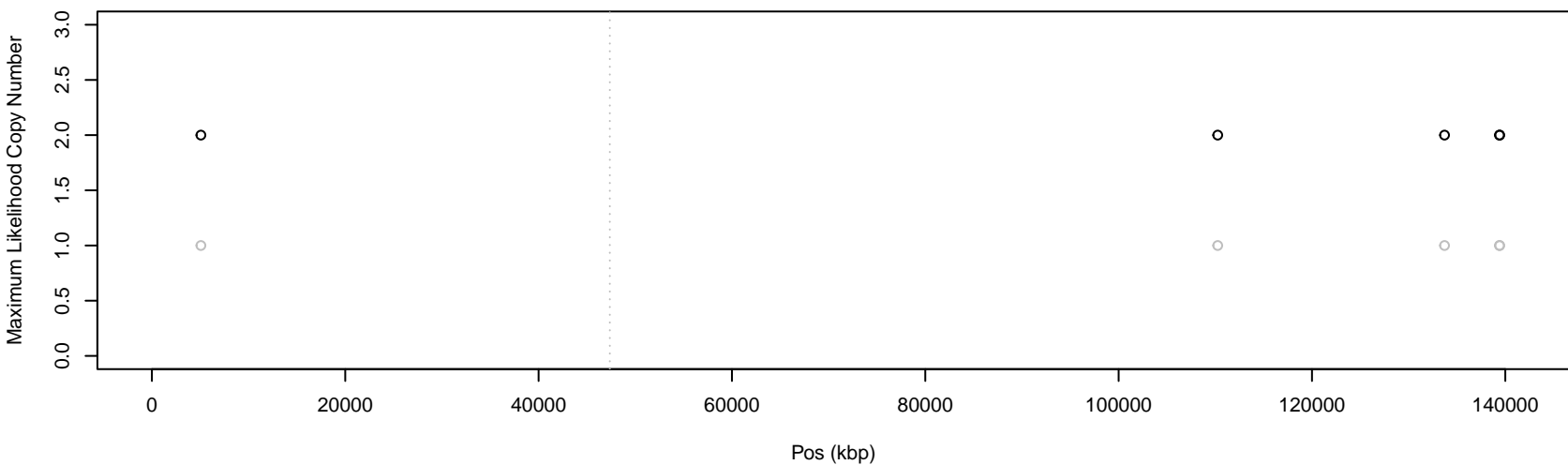
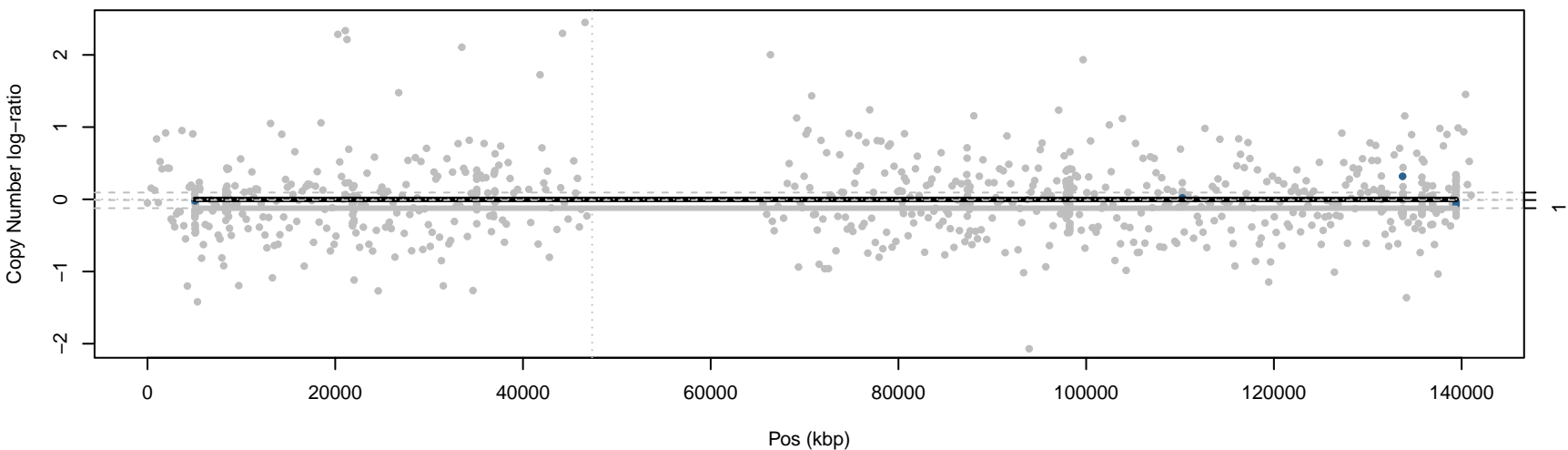
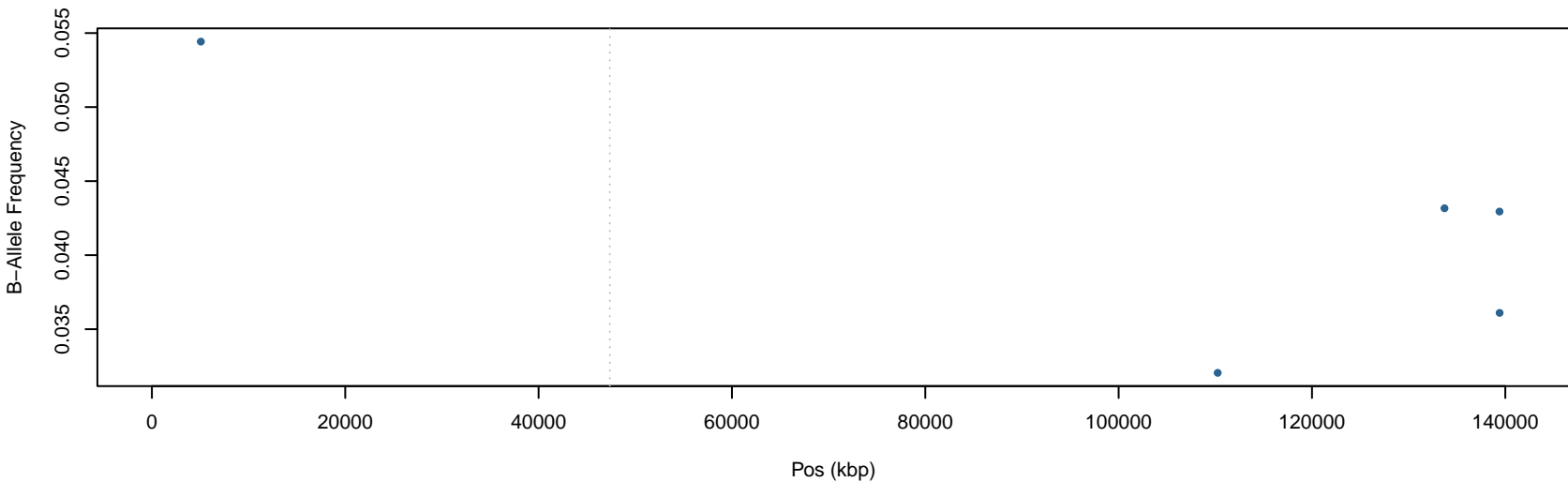
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 6



Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 7

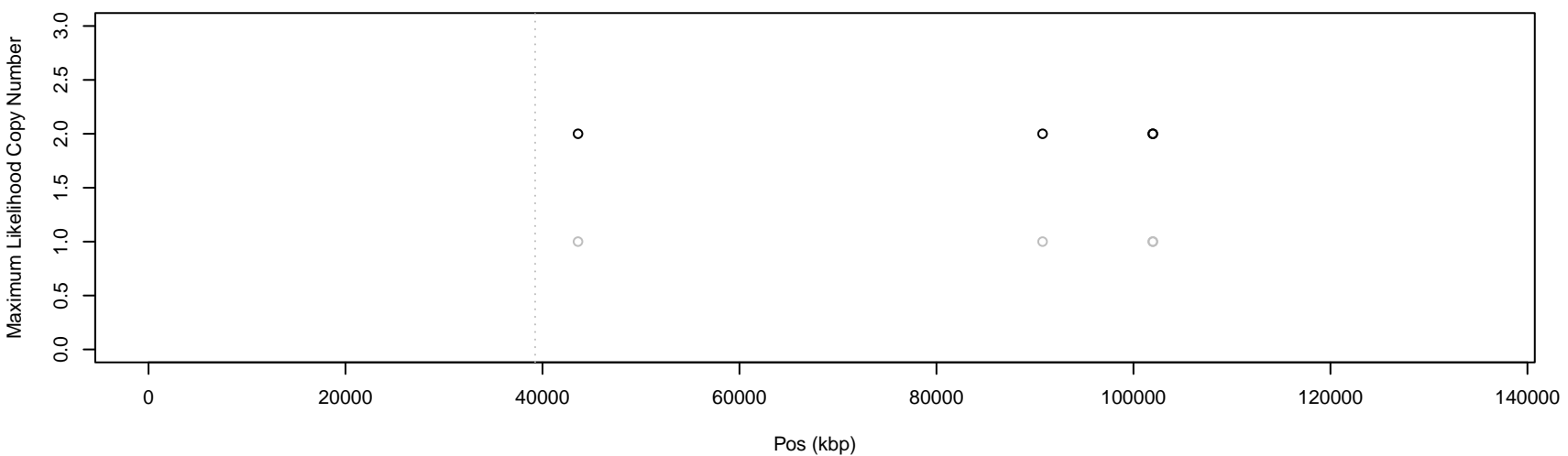
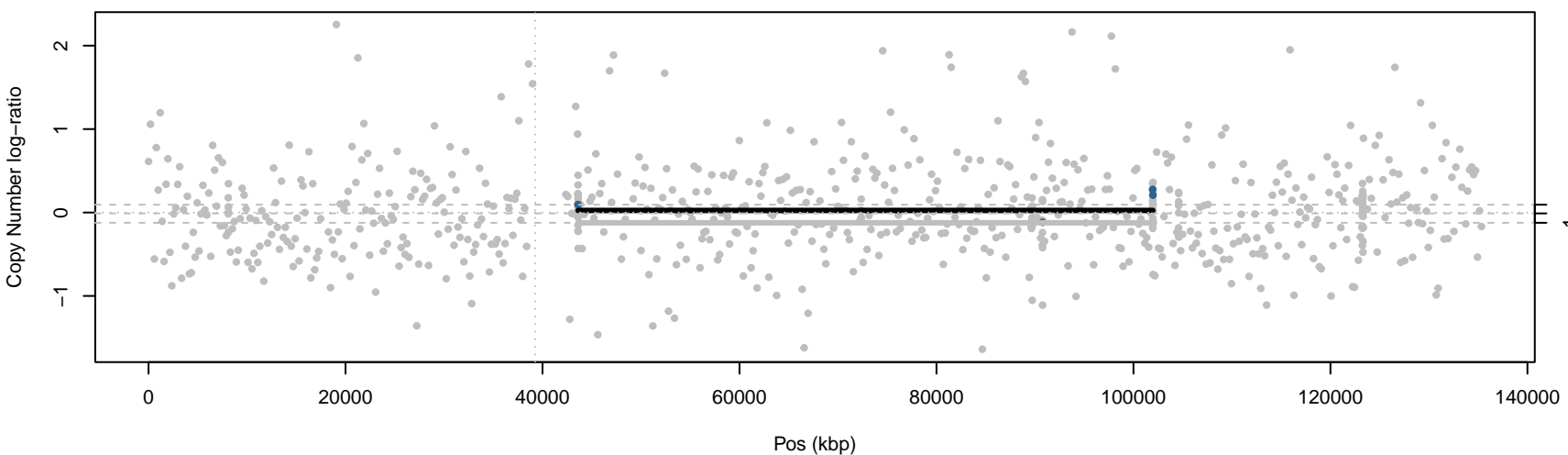
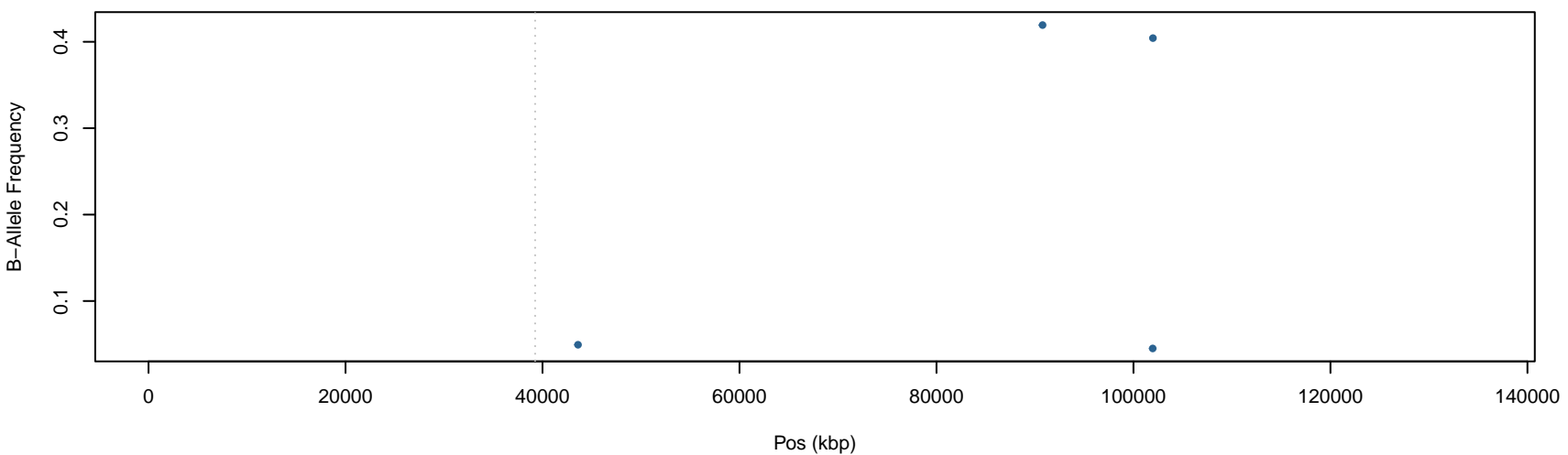


Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 9

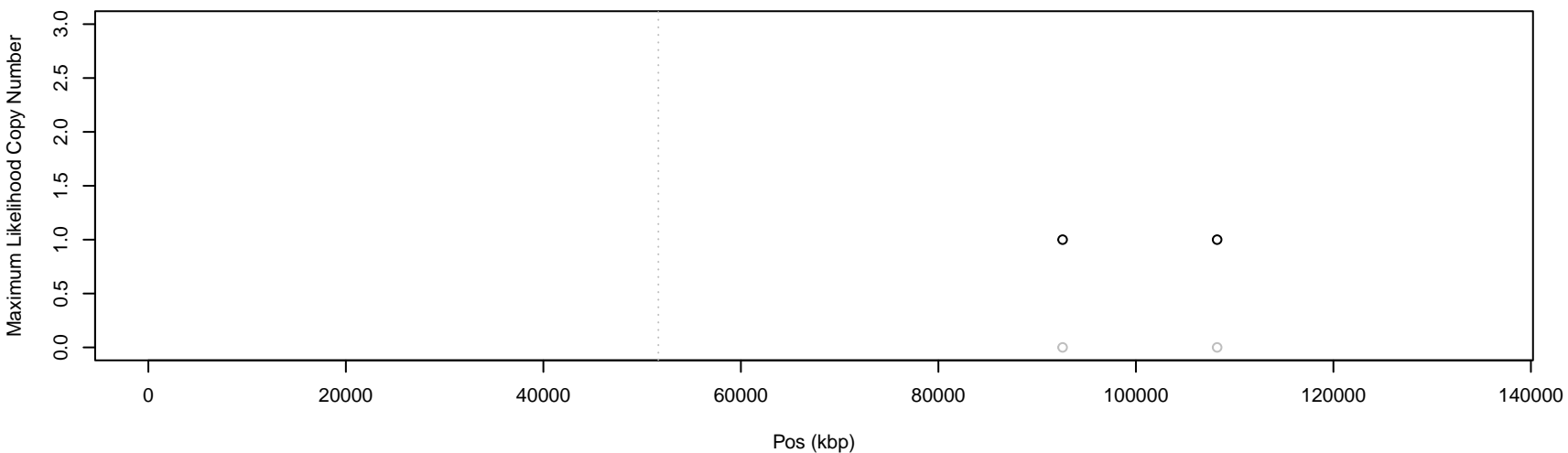
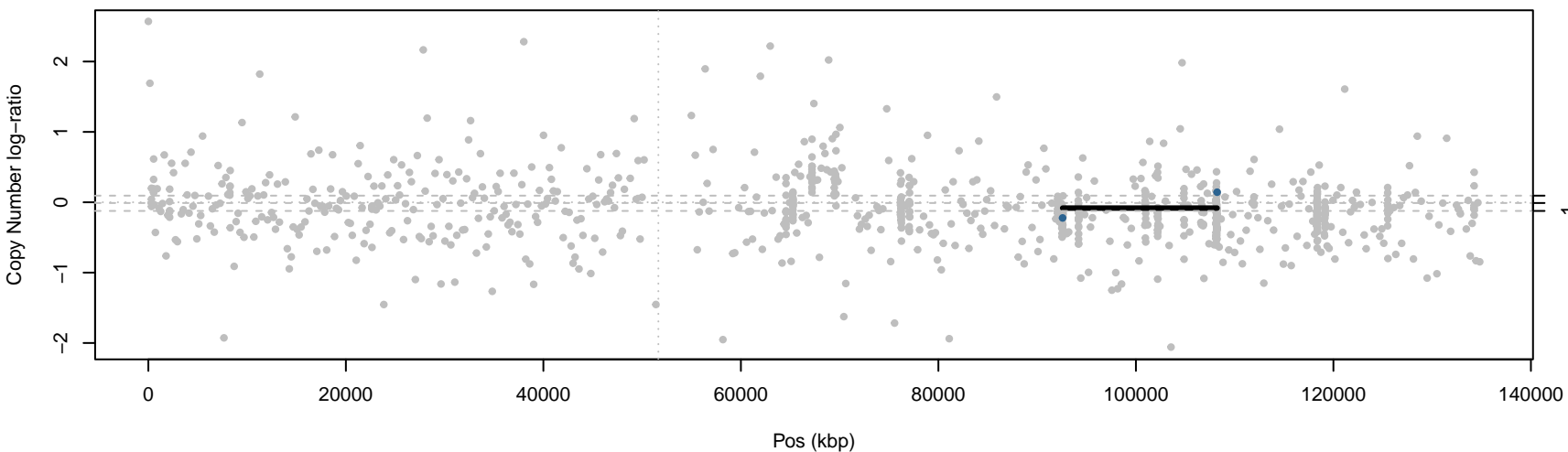
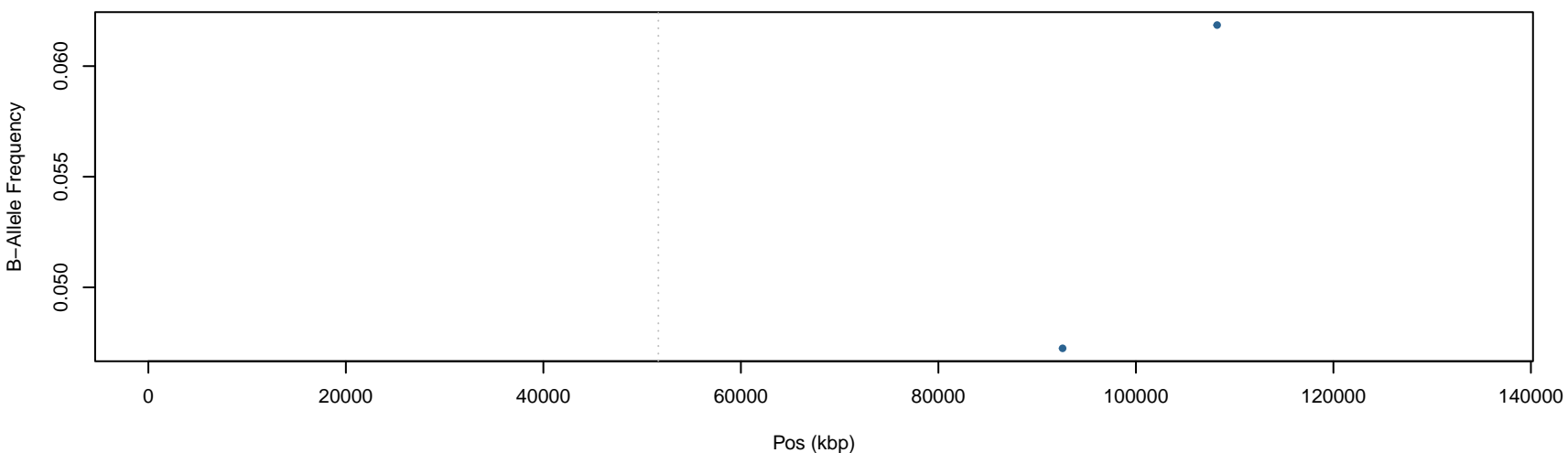




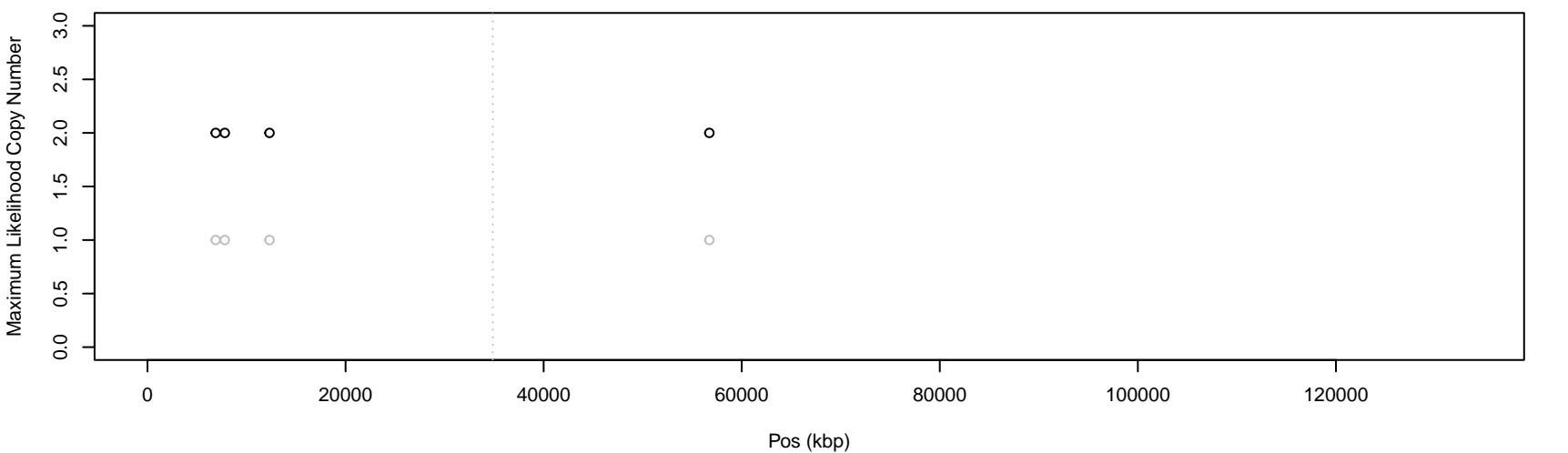
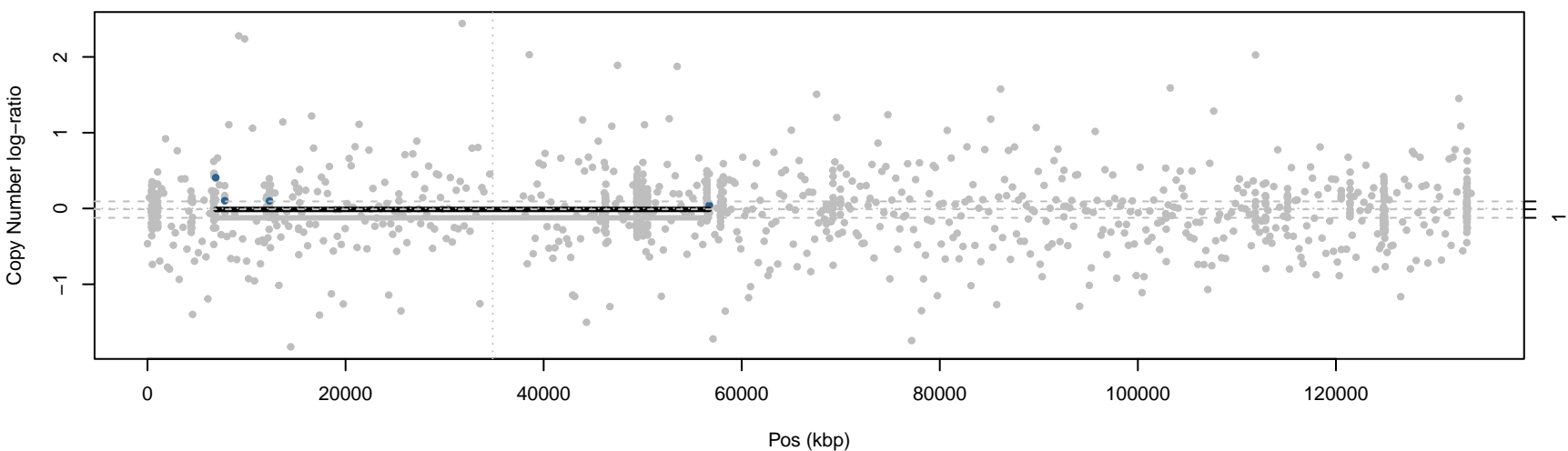
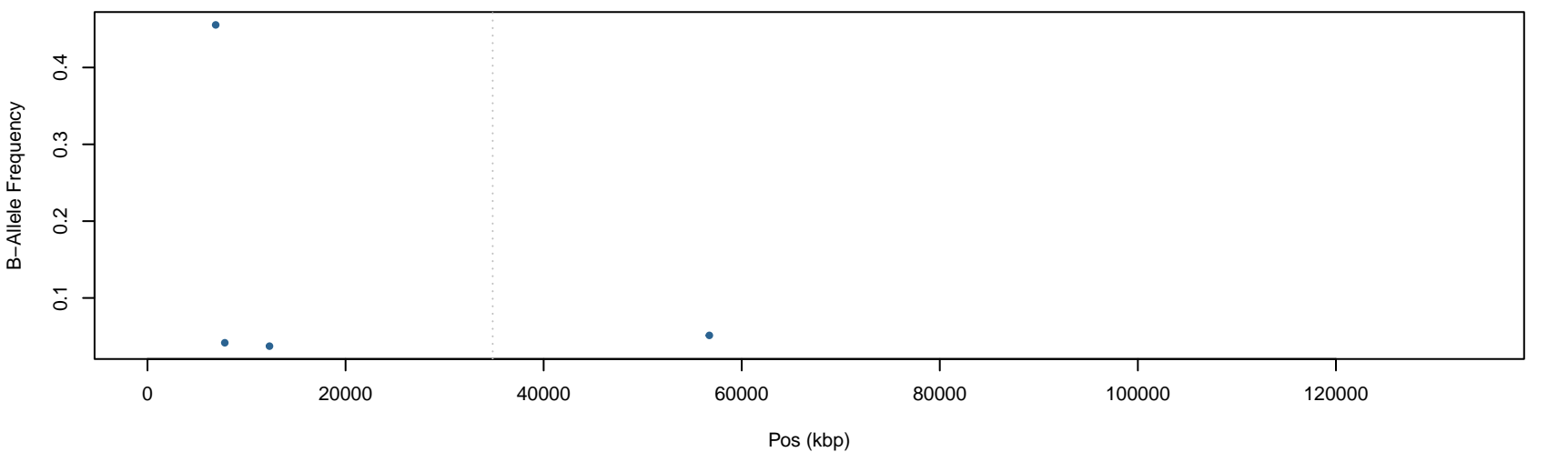
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 10



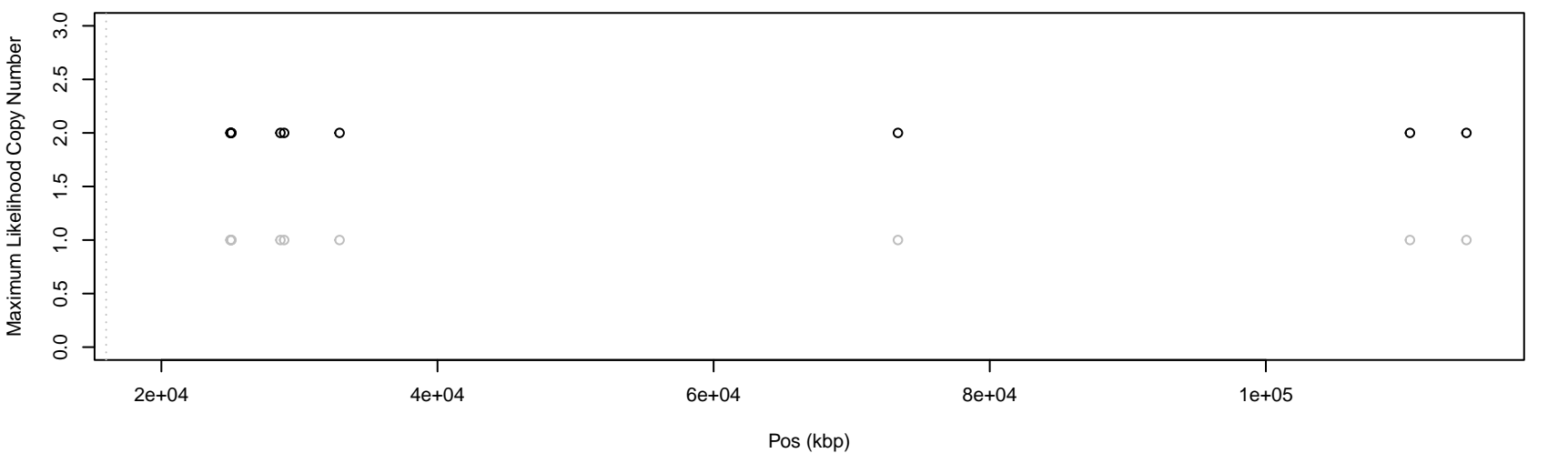
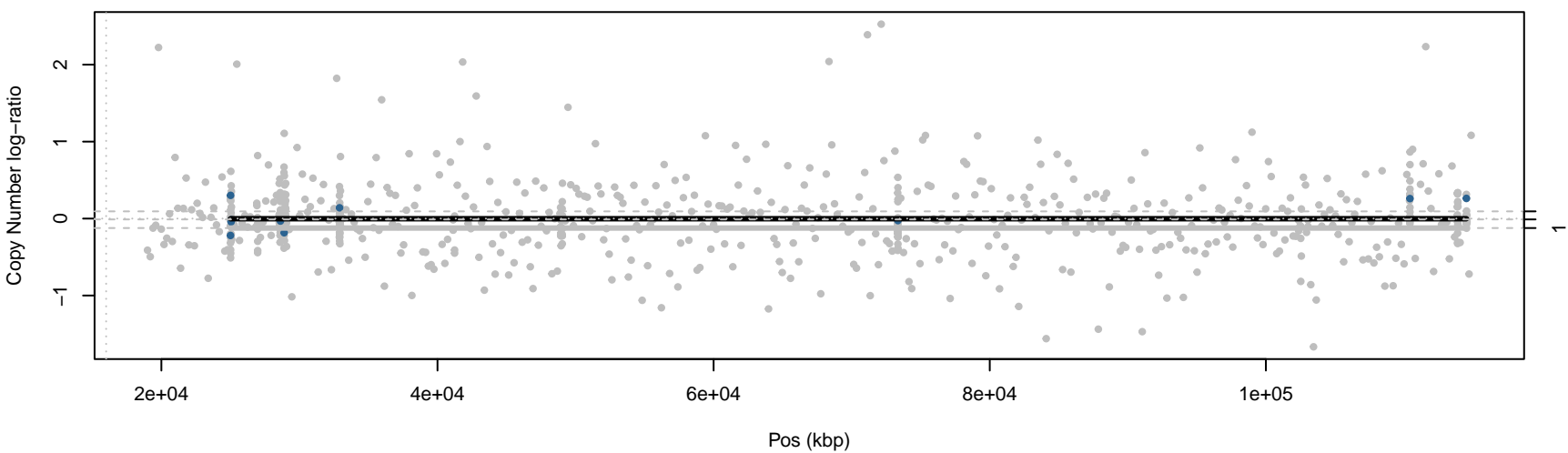
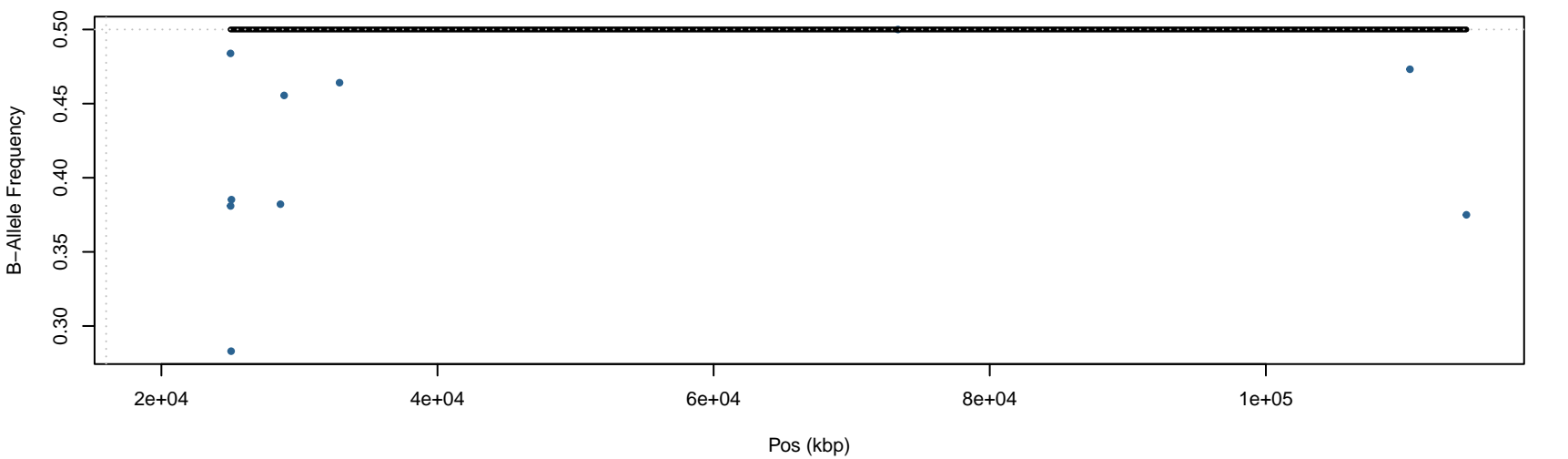
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 11



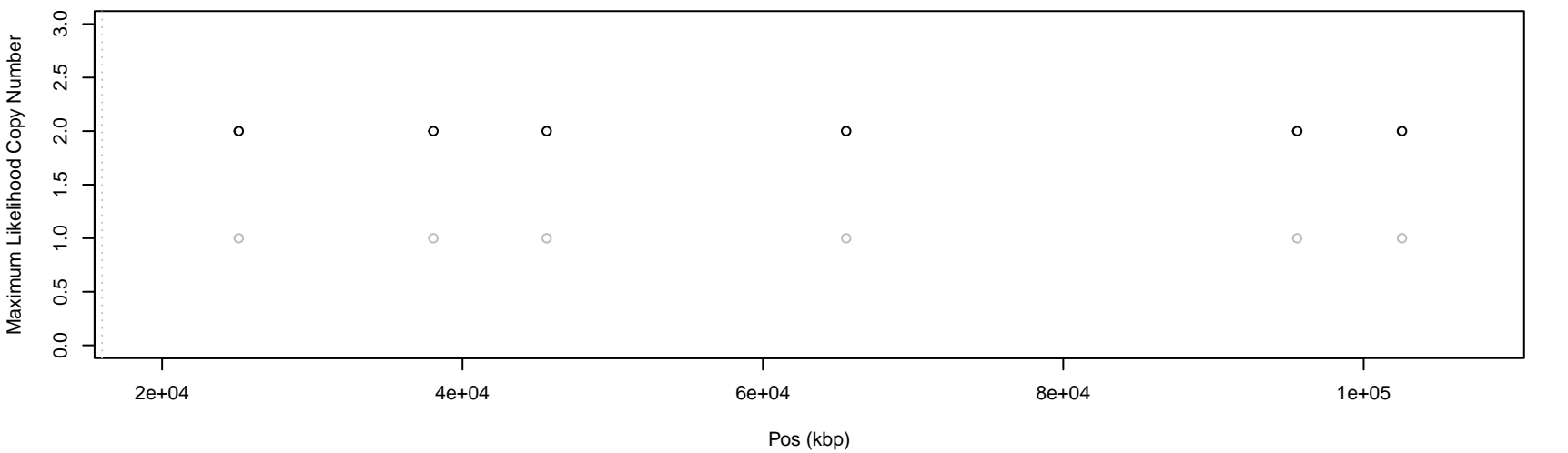
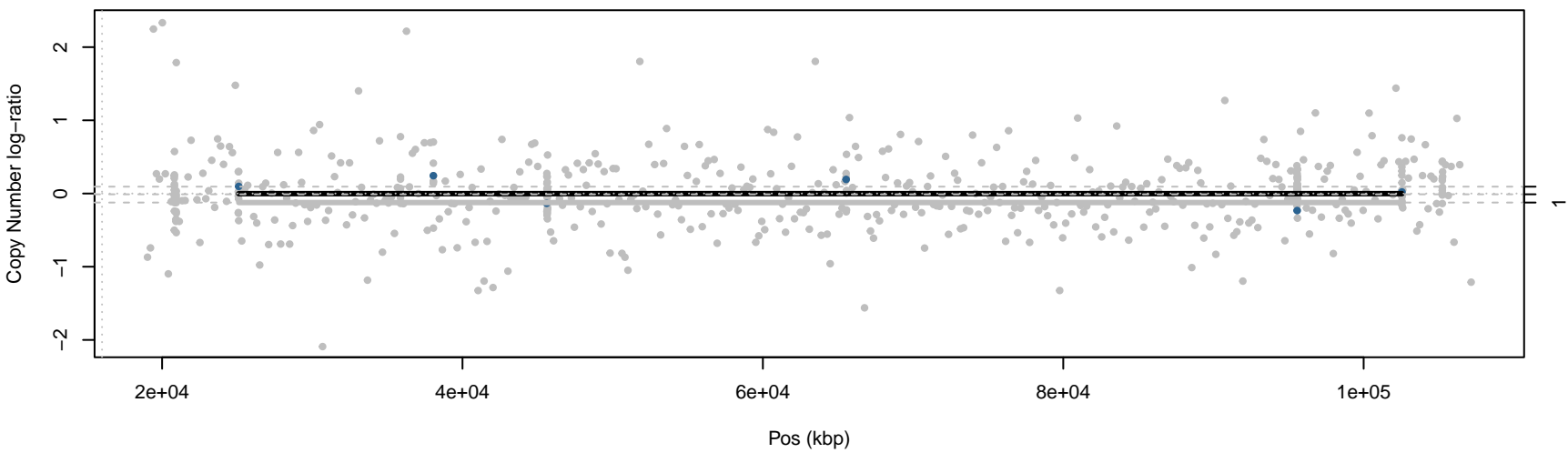
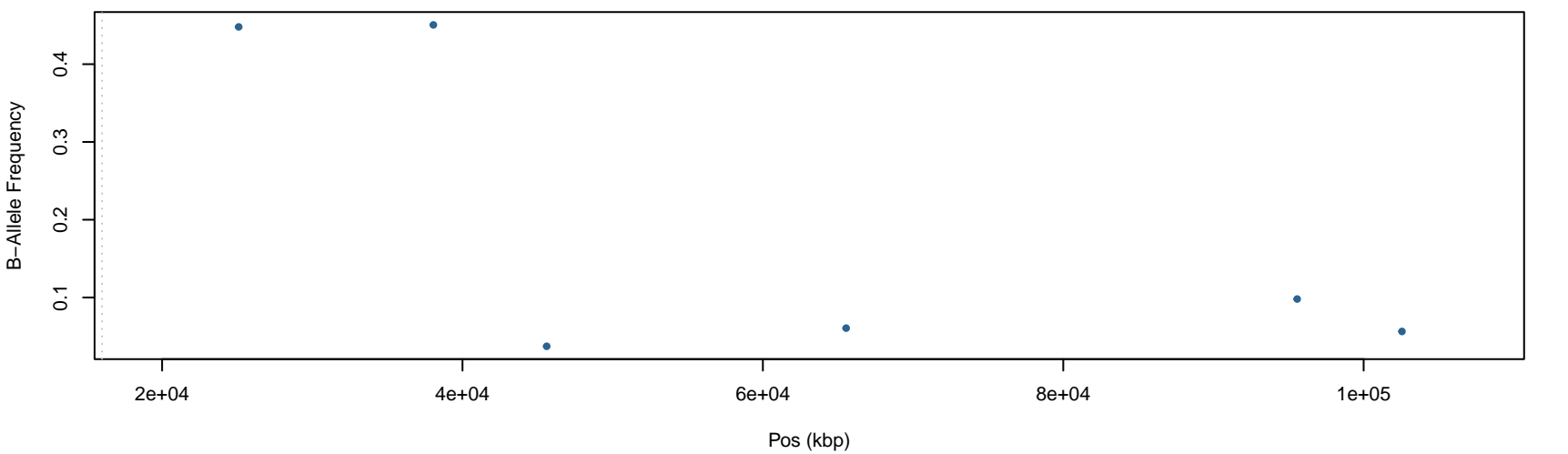
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 12



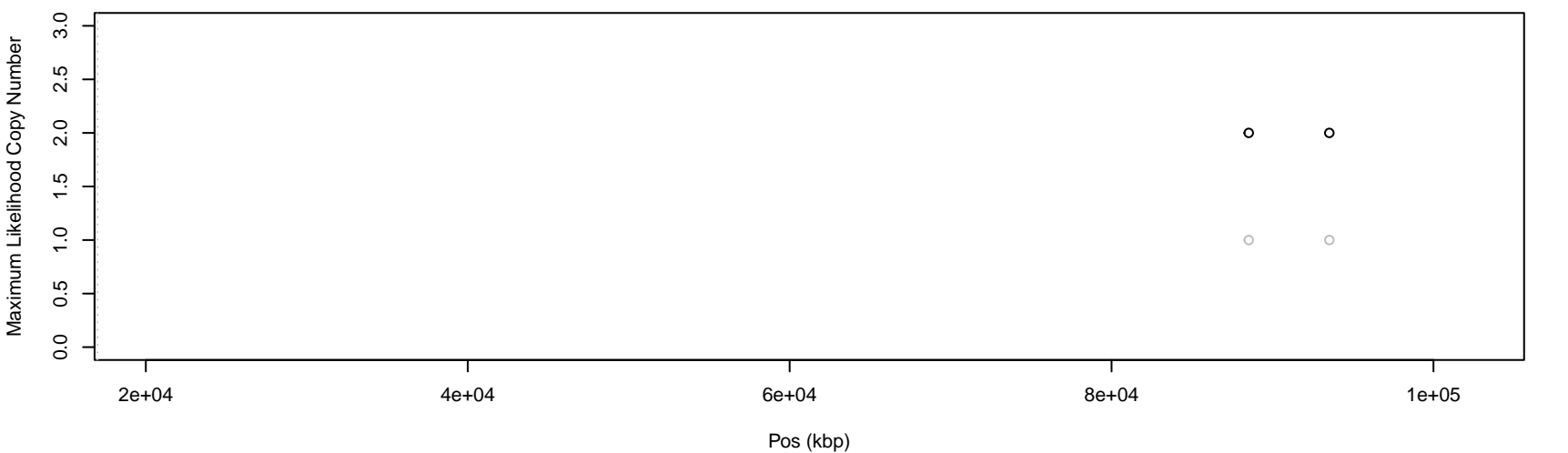
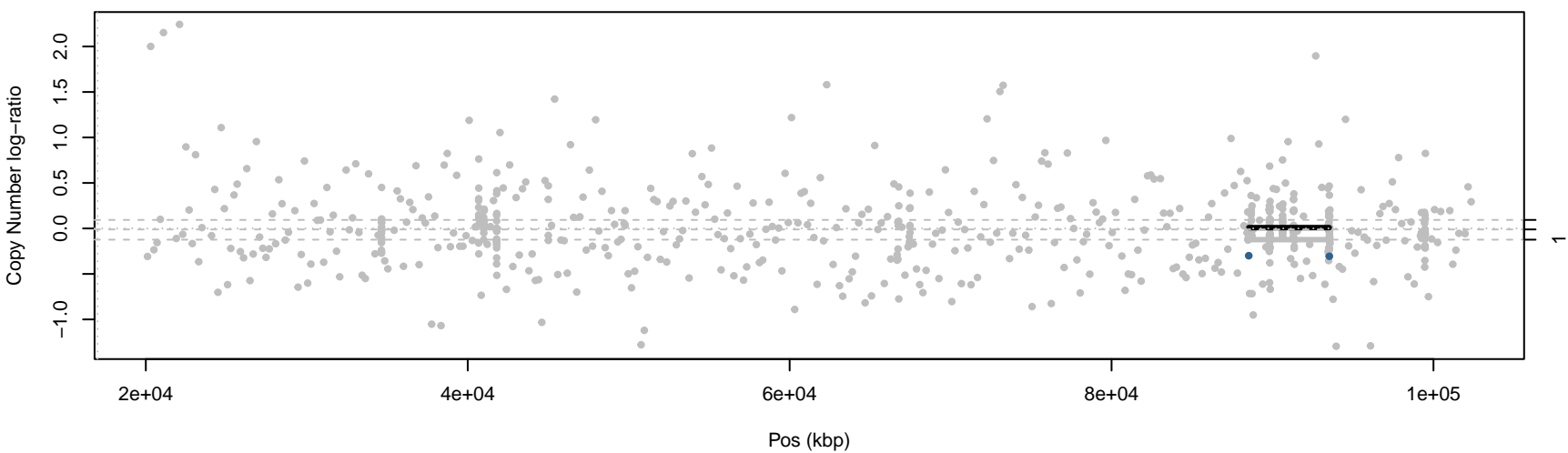
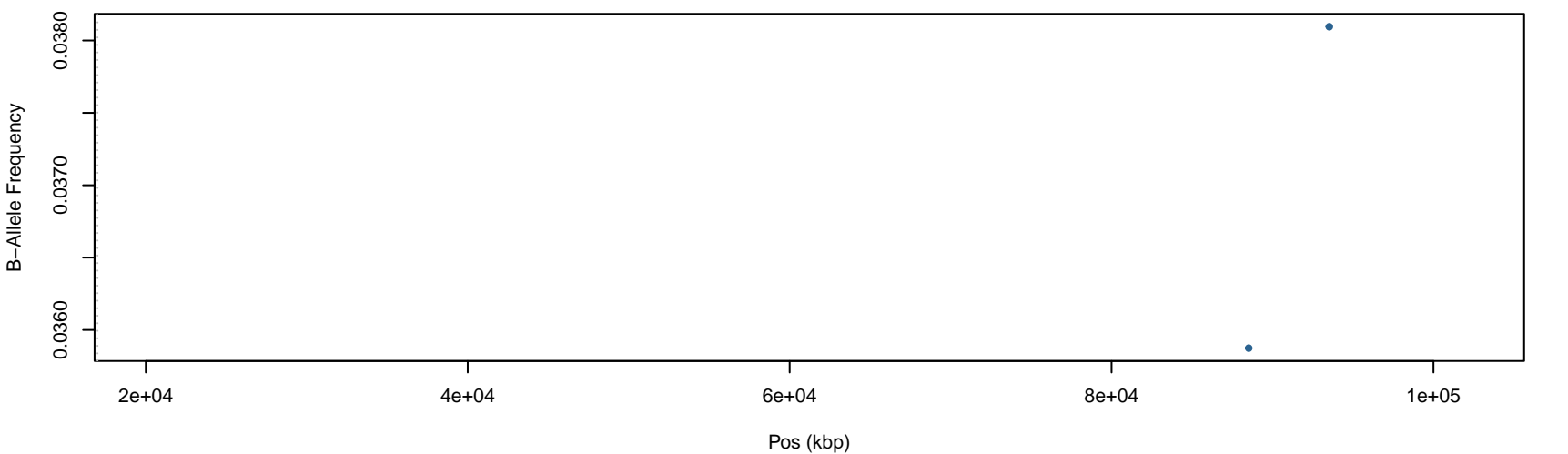
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 13



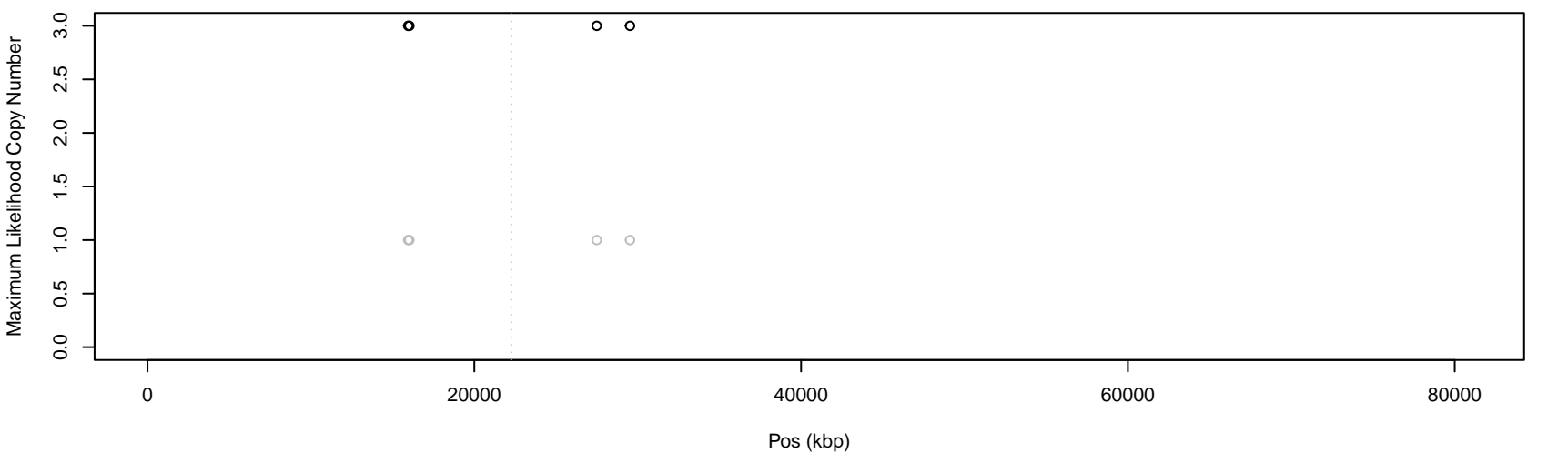
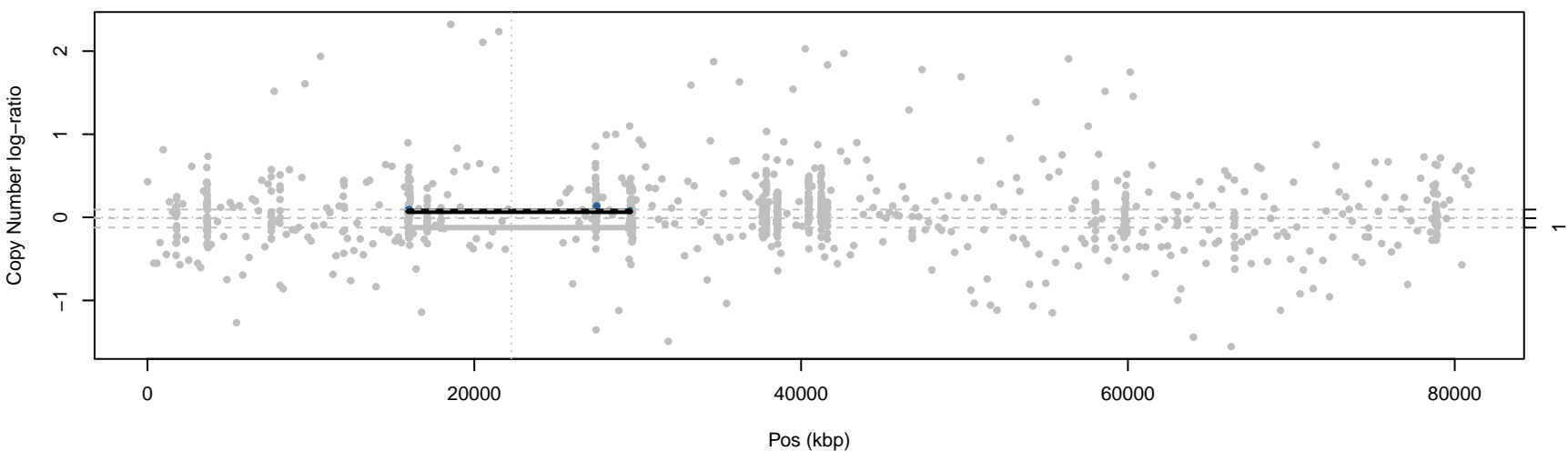
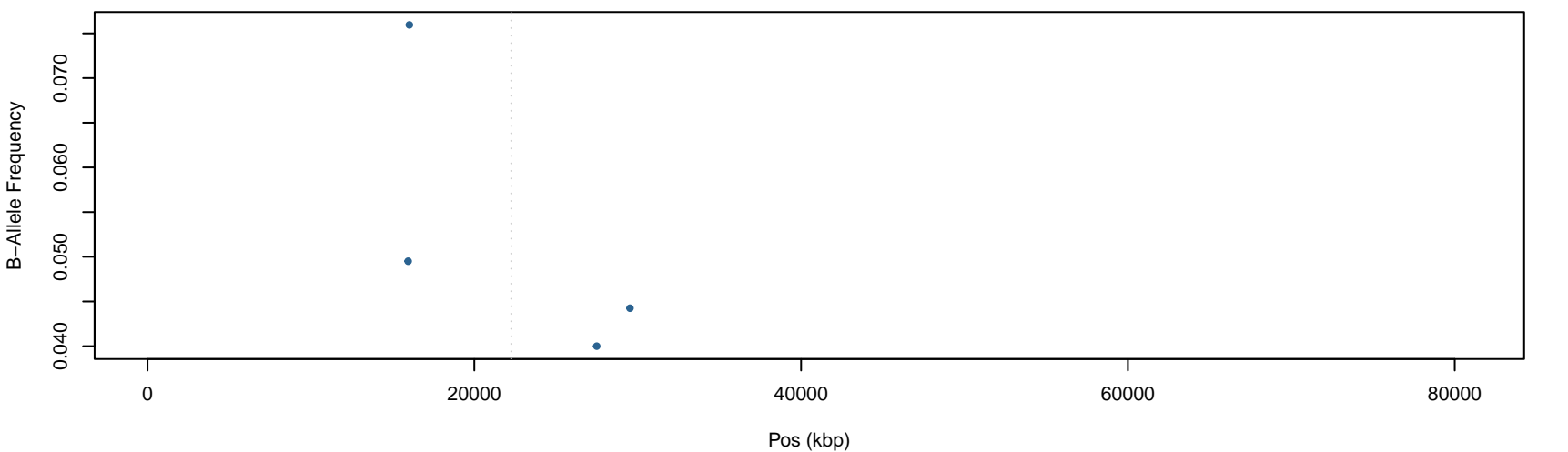
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 14



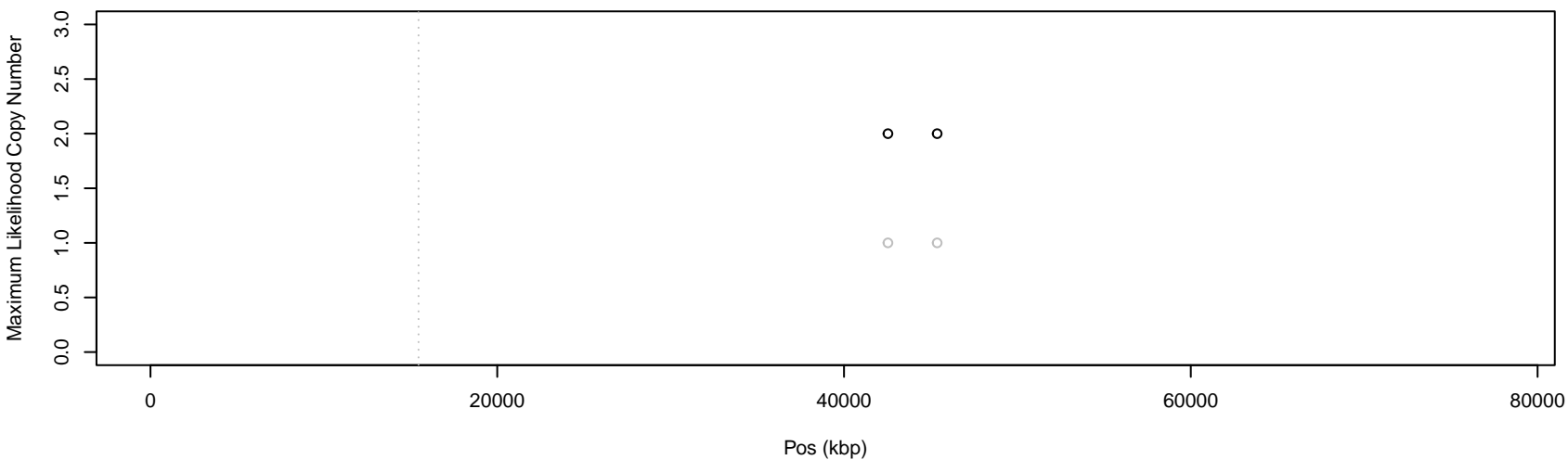
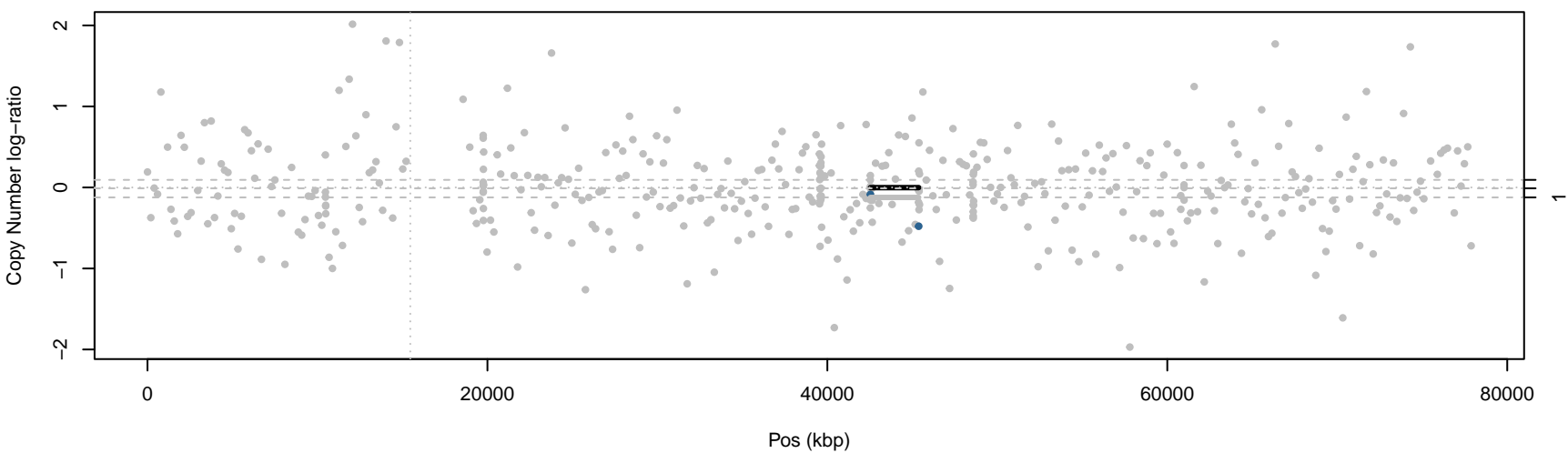
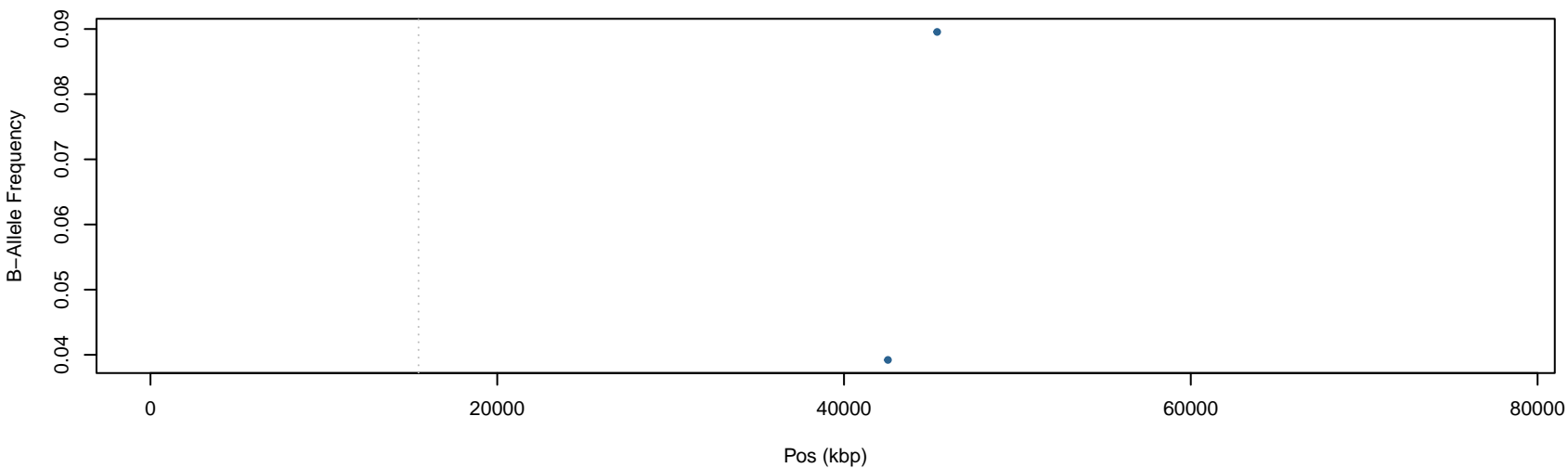
Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 15



Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 17

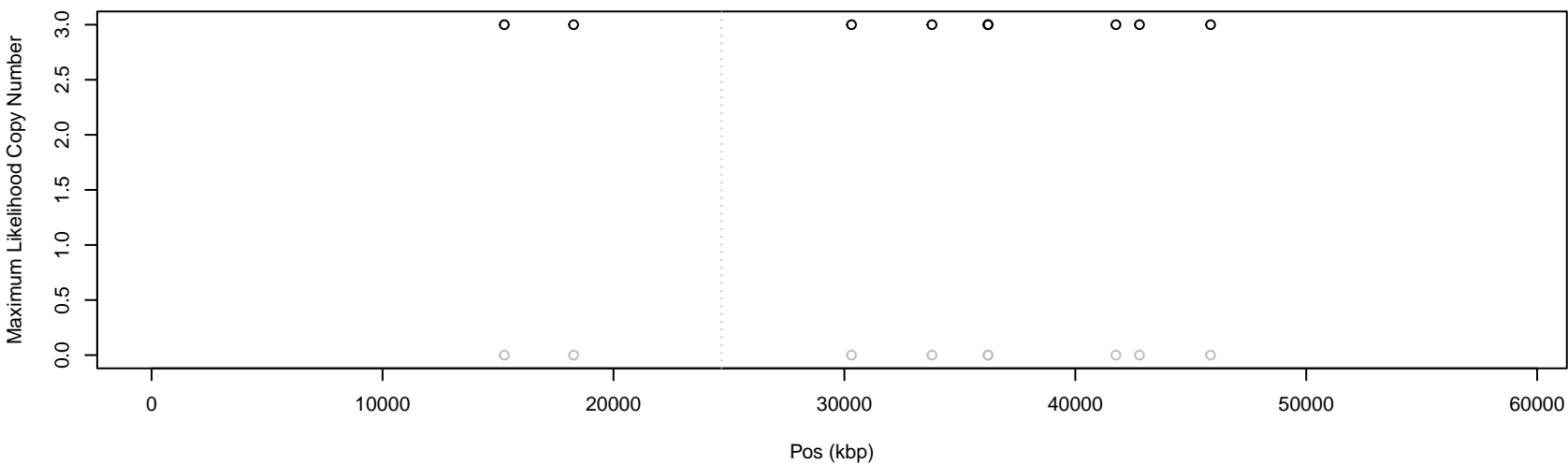
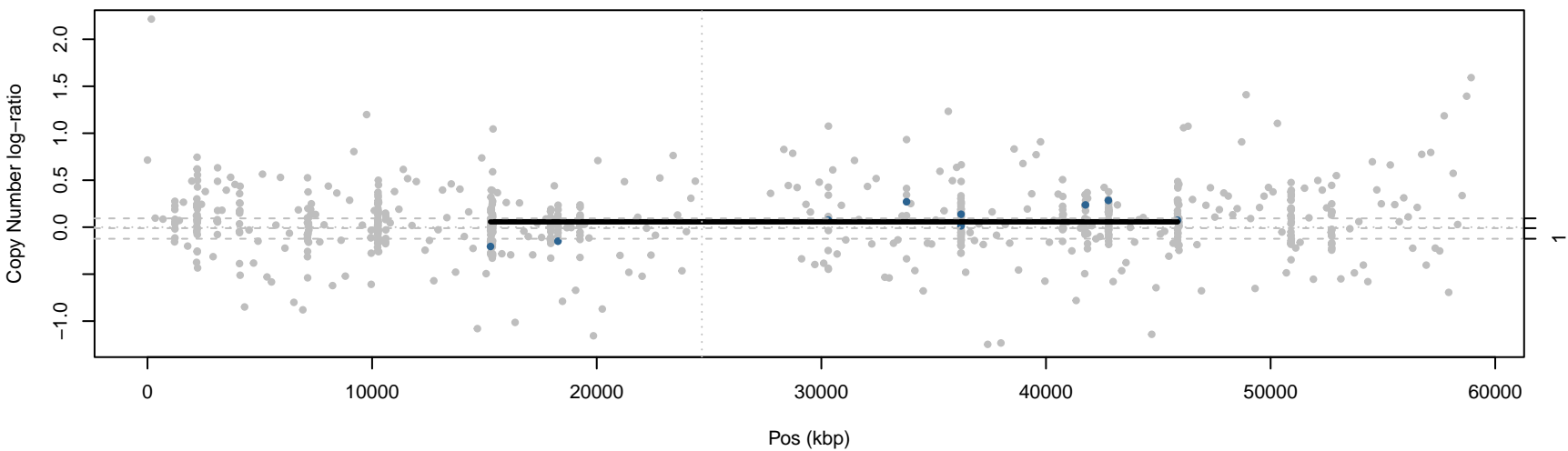
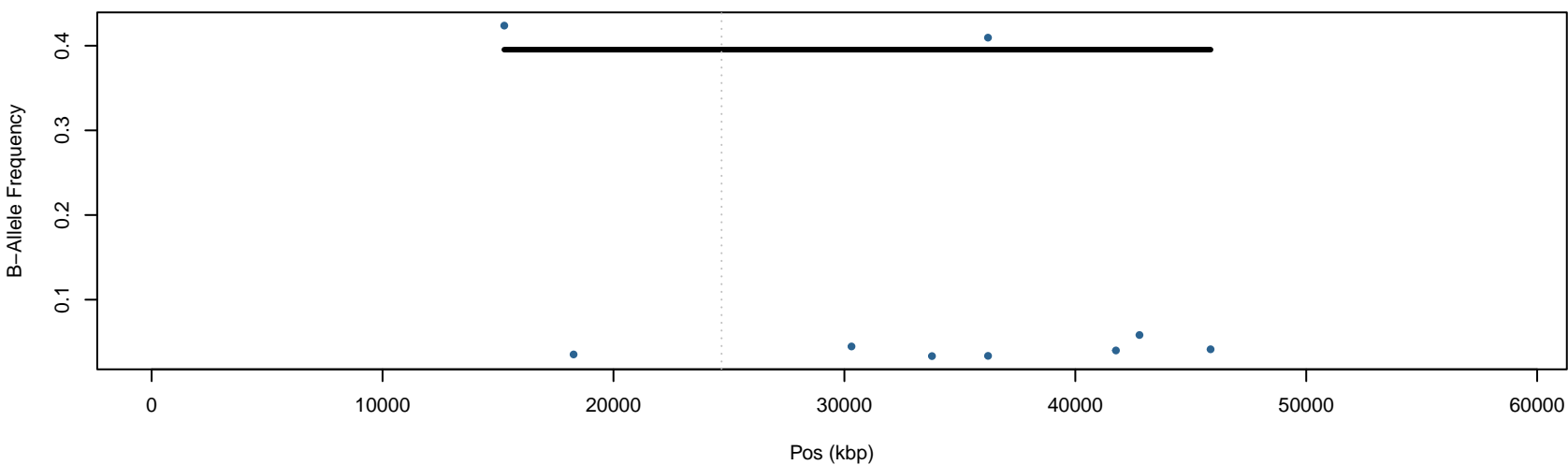


Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 18





Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 19



Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149 Chromosome: 20

