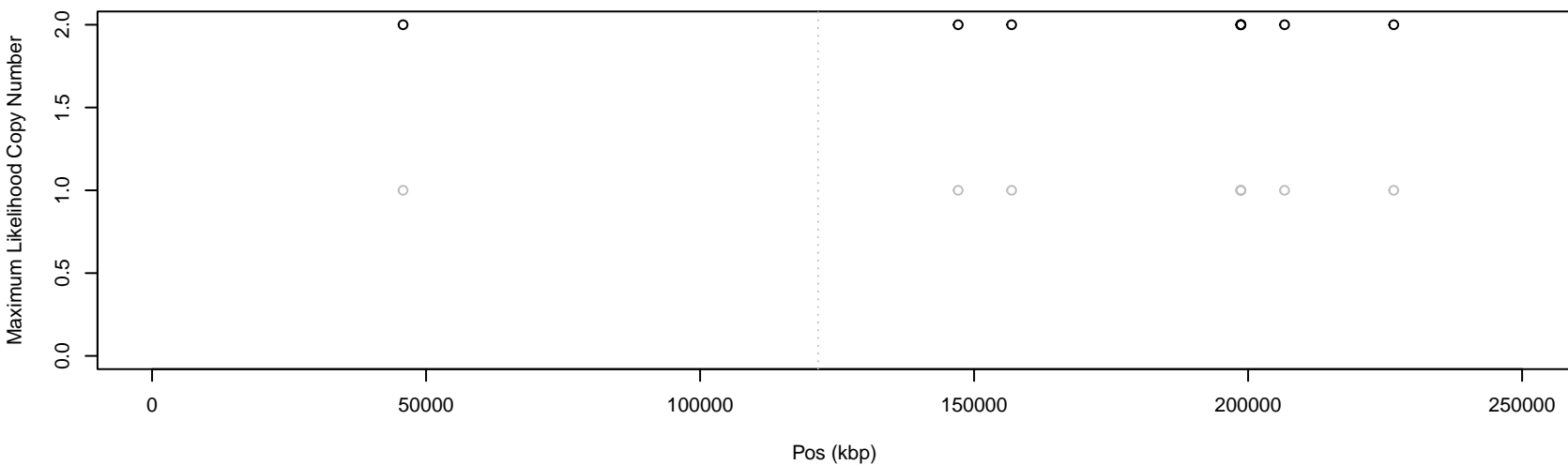
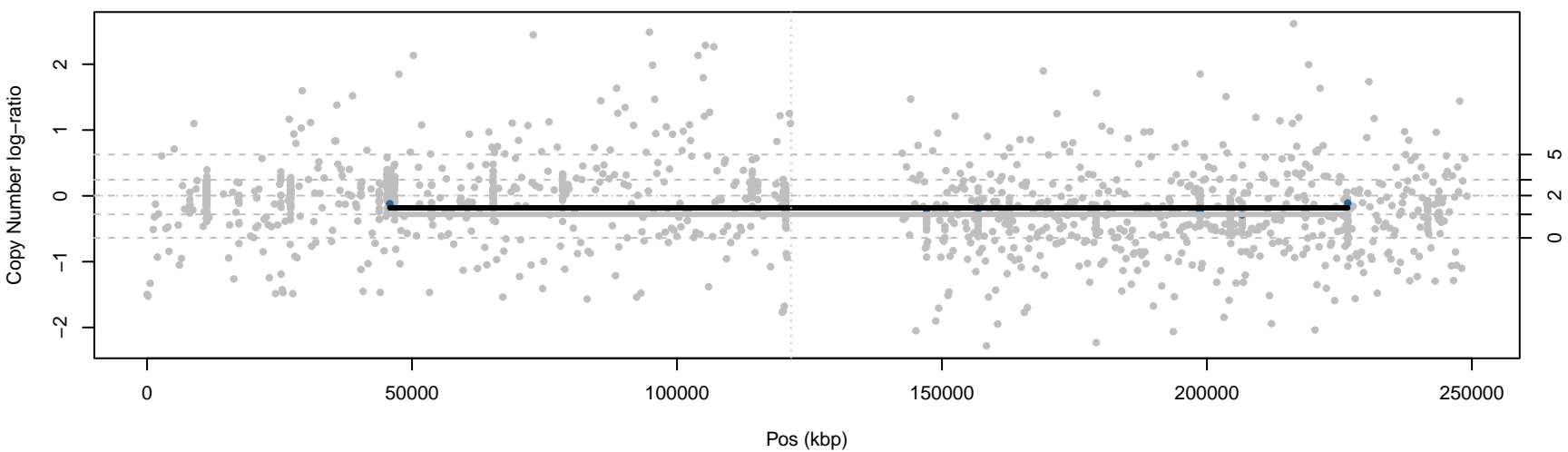
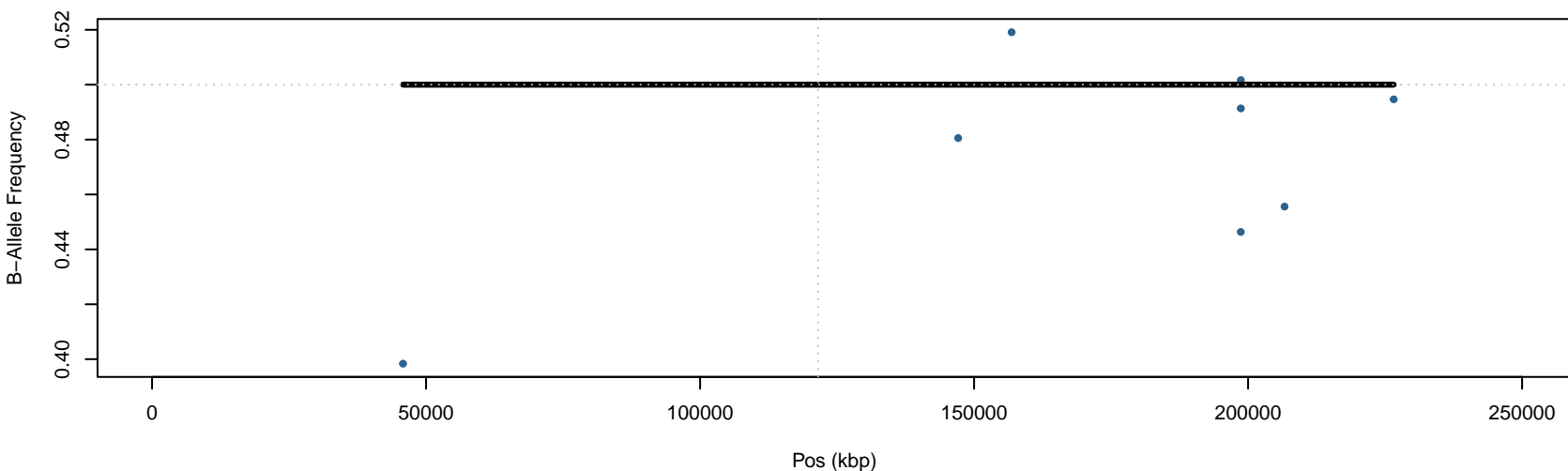
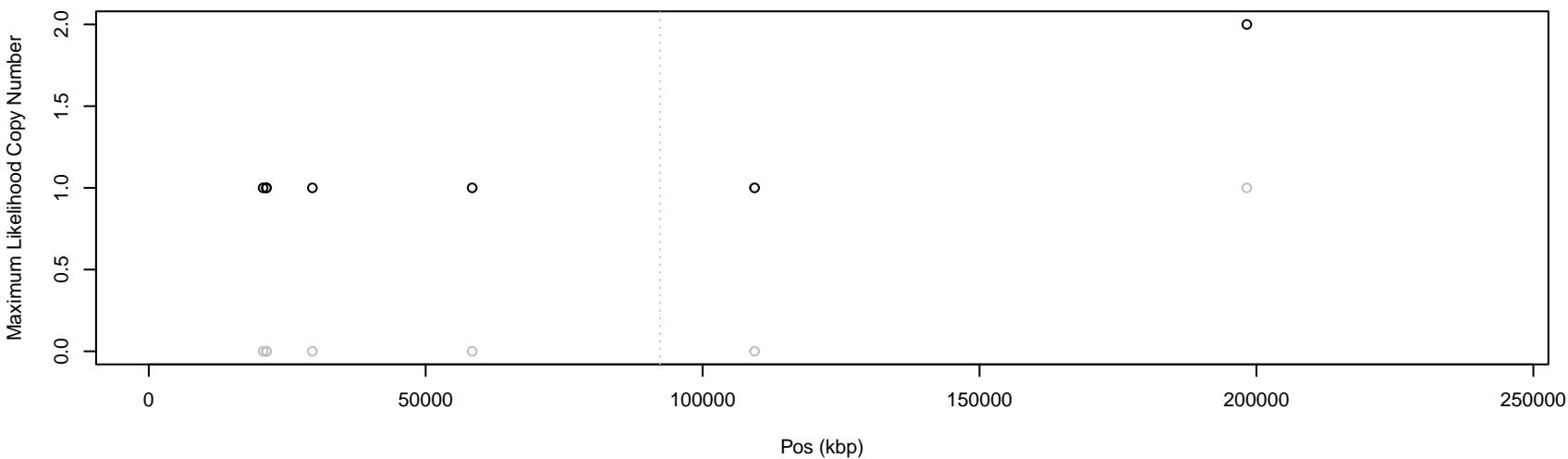
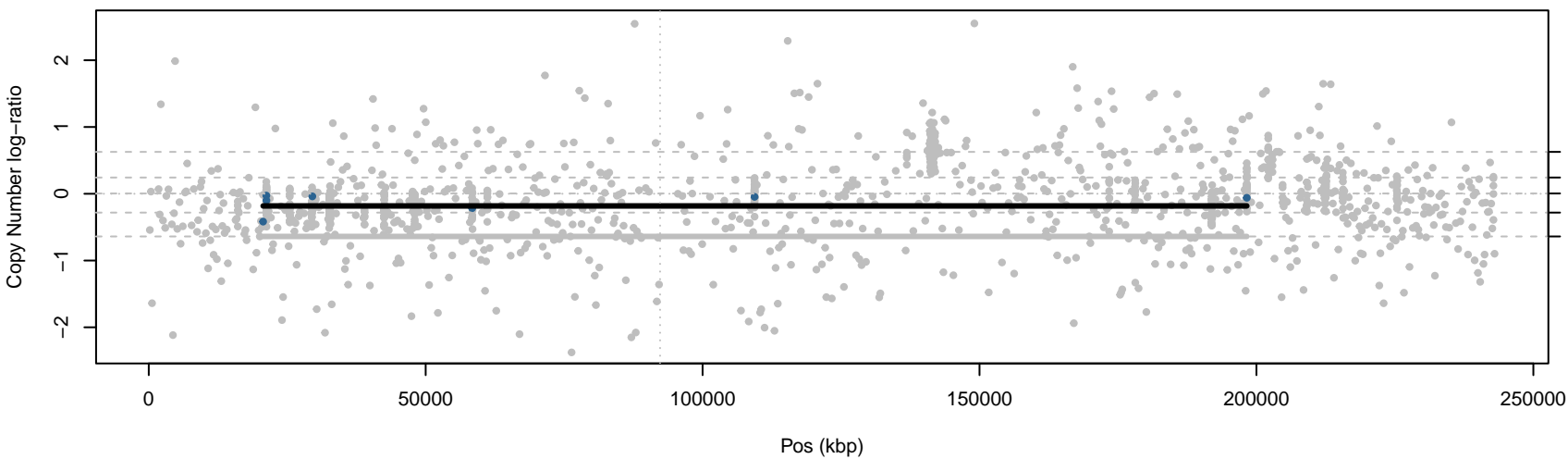
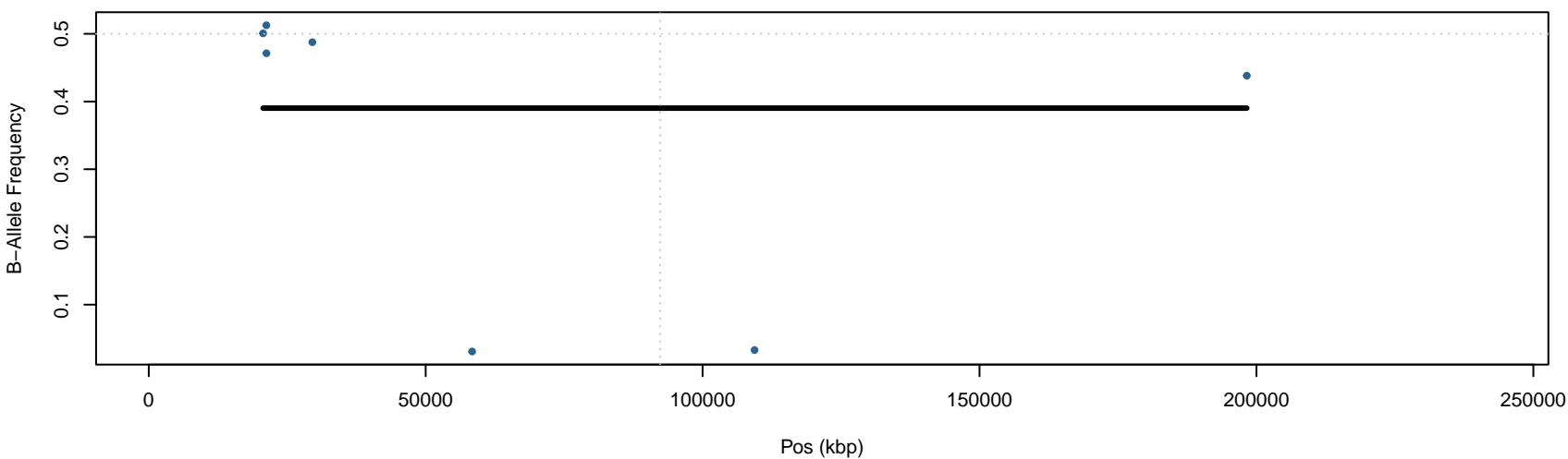


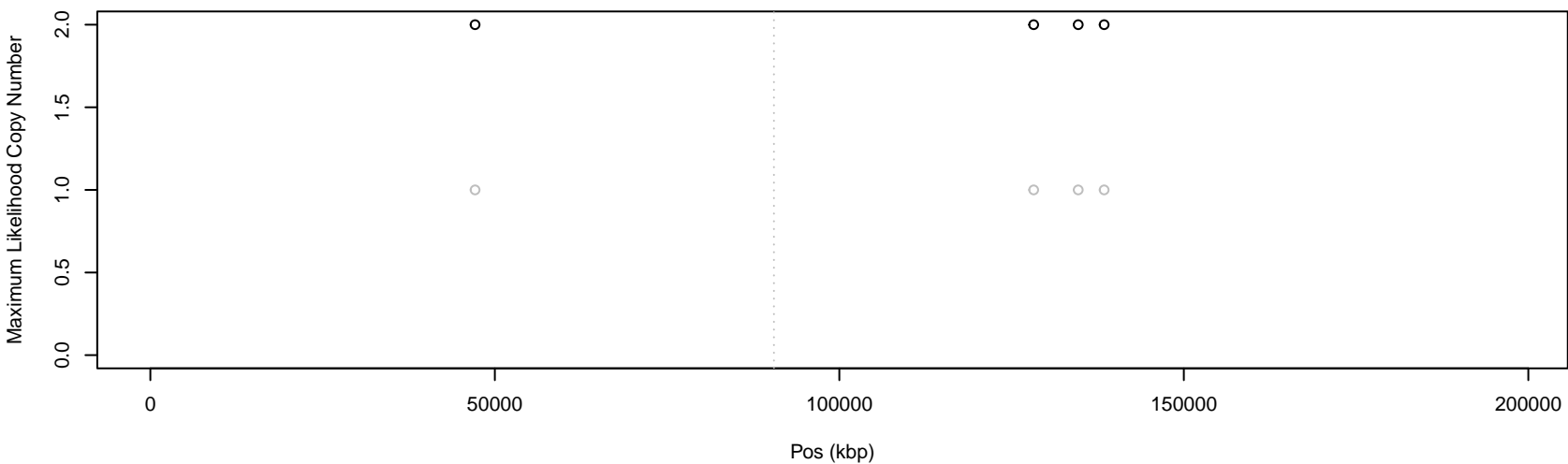
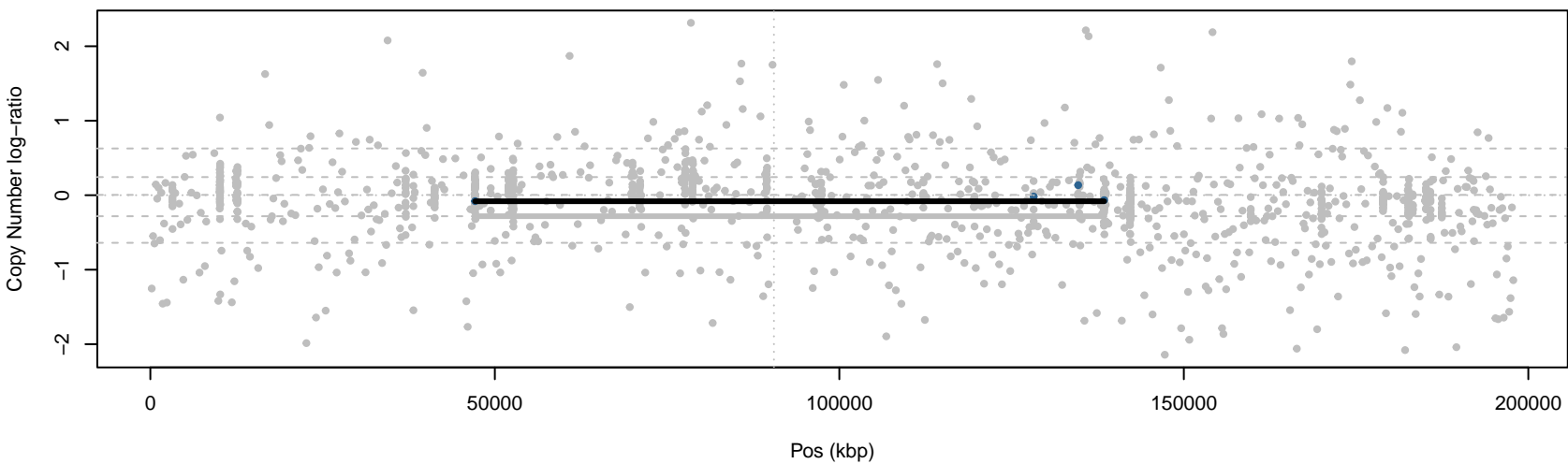
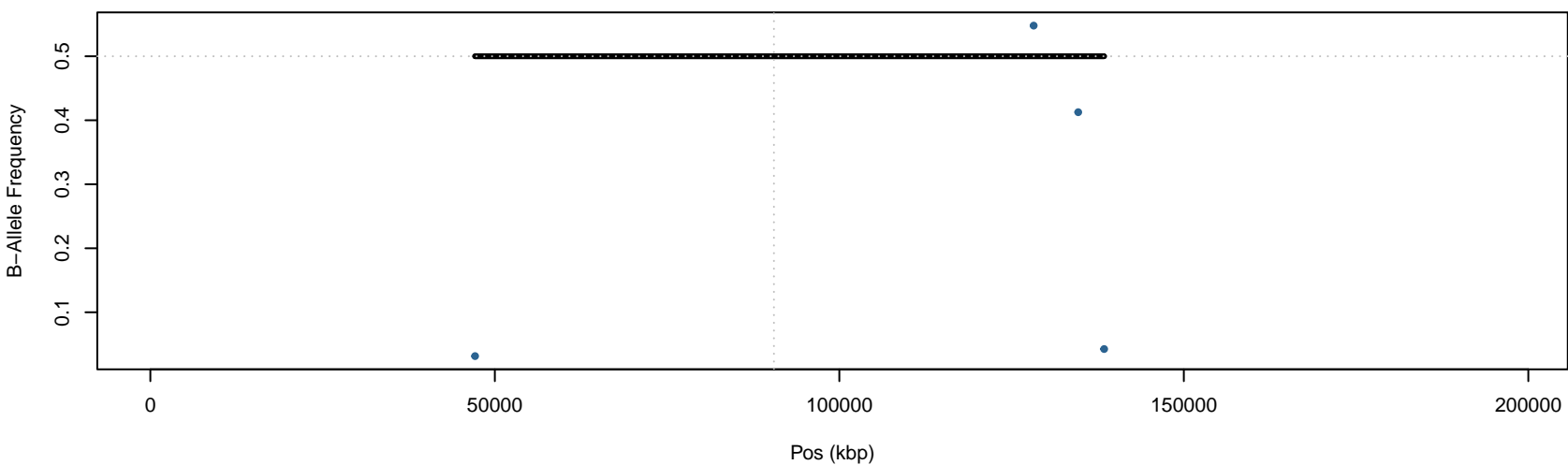
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 1



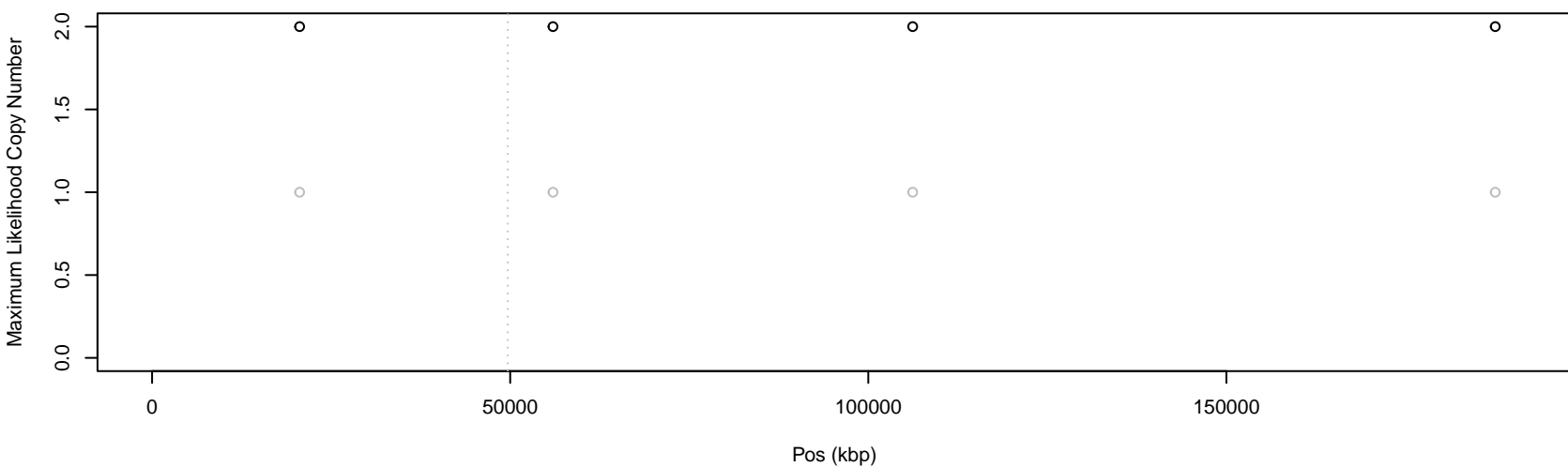
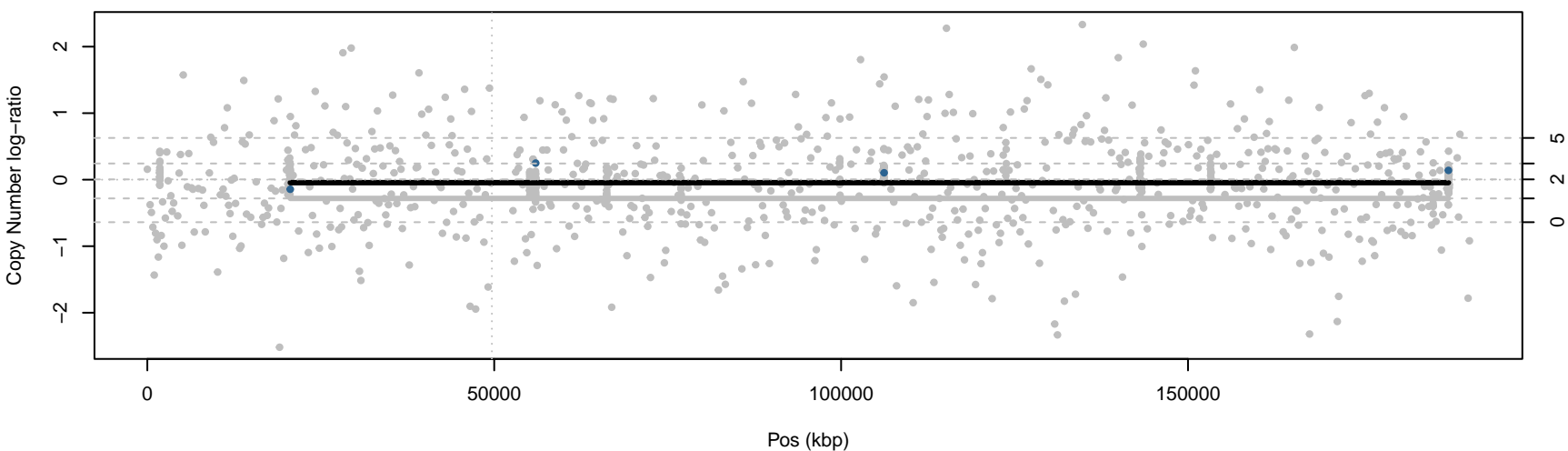
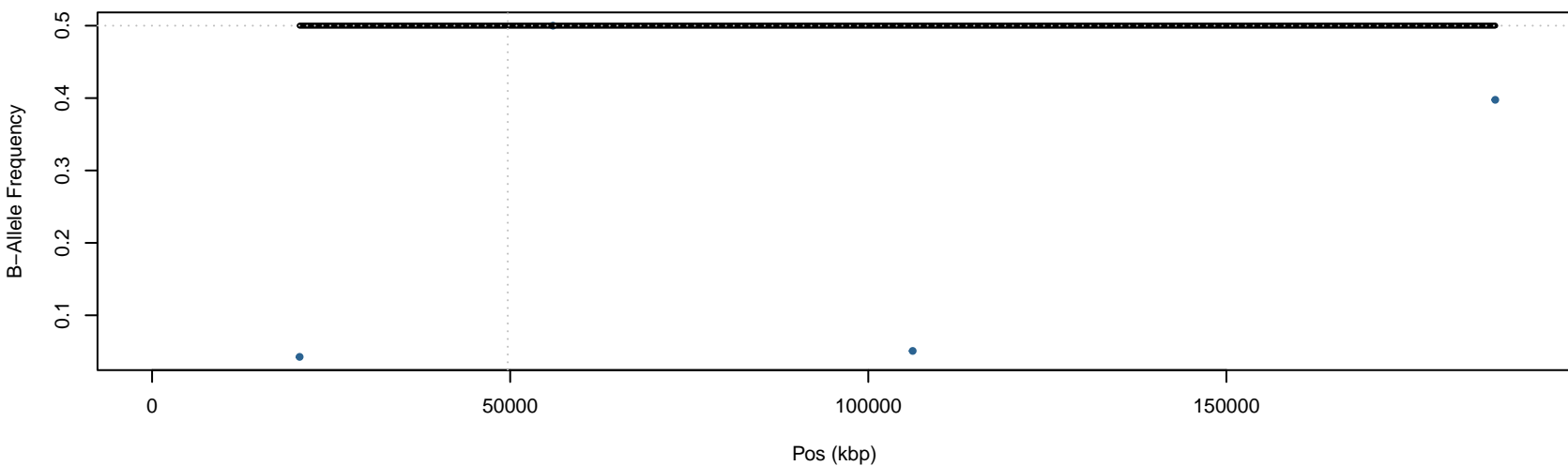
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 2



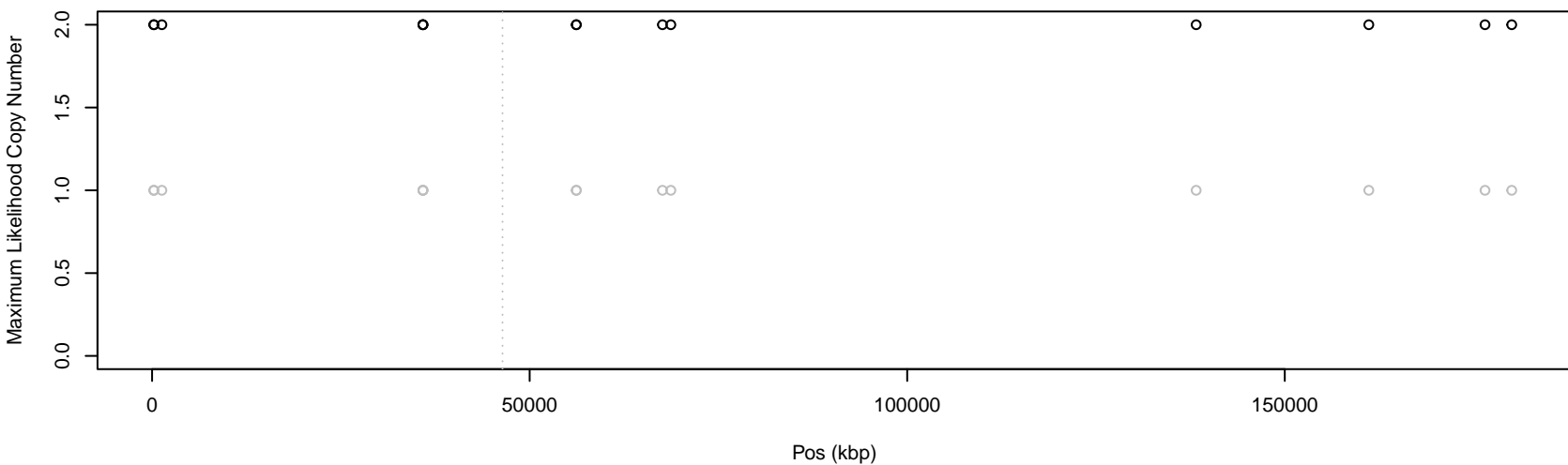
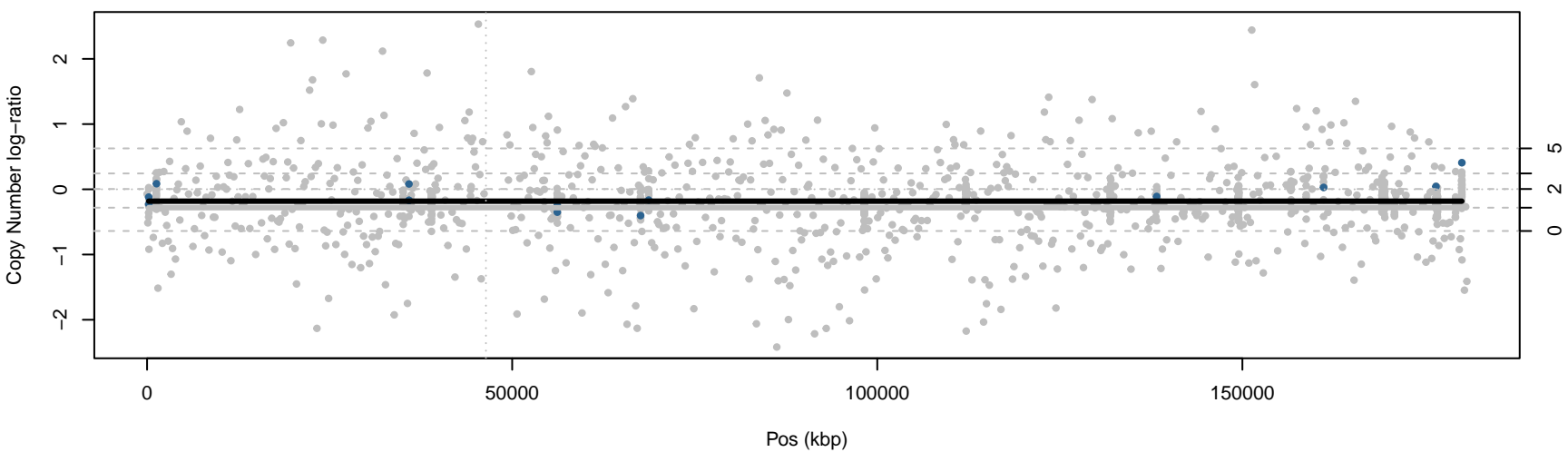
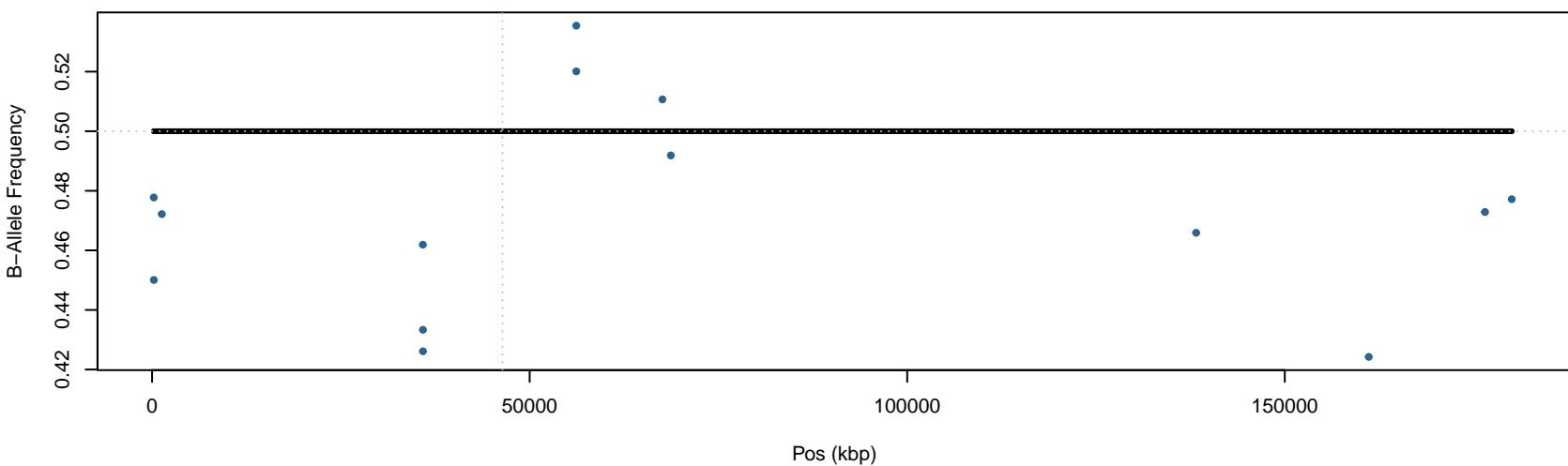
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 3



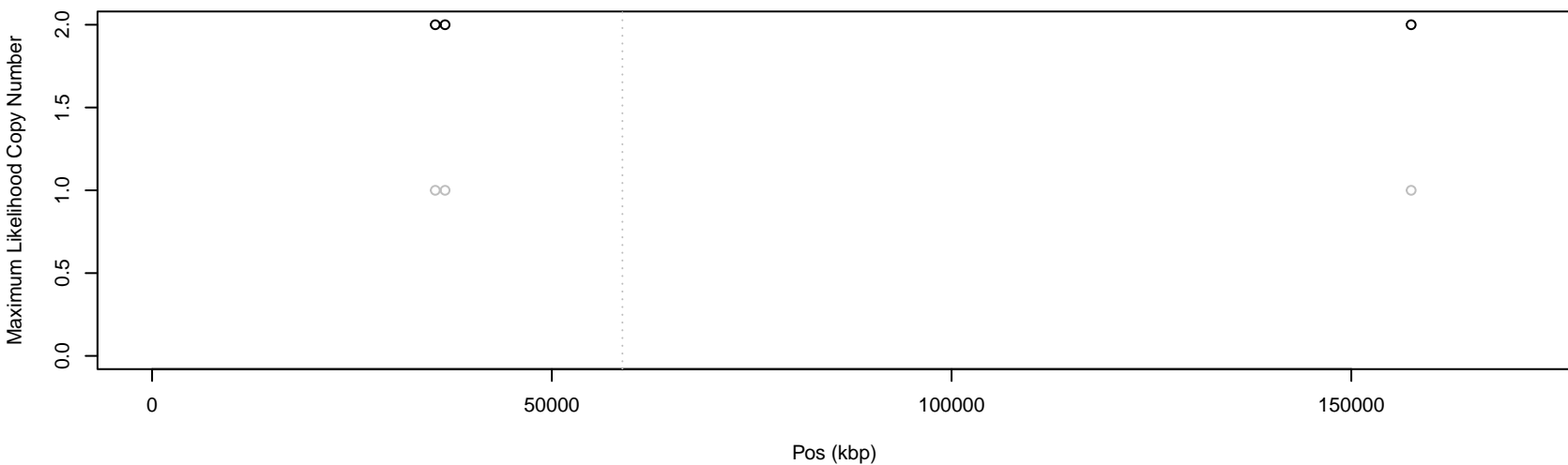
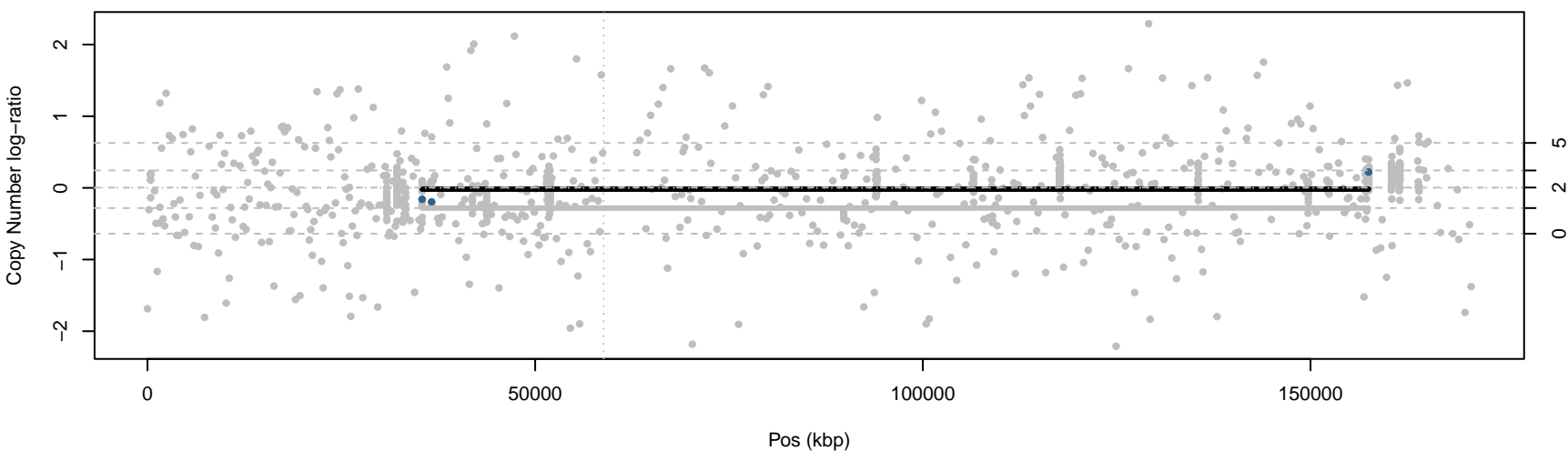
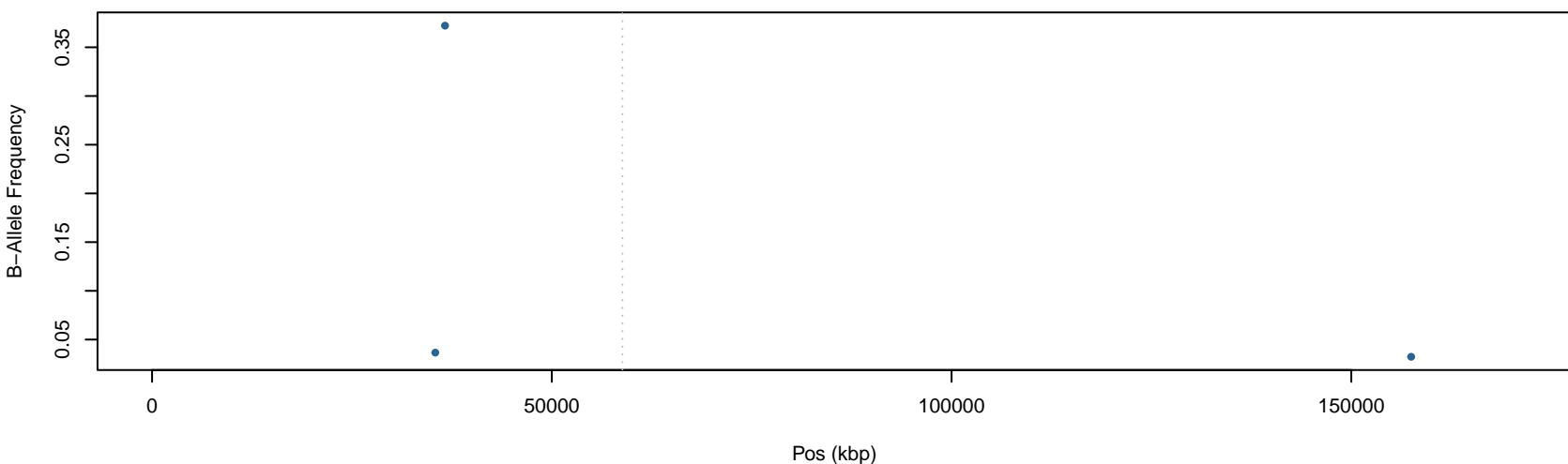
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 4



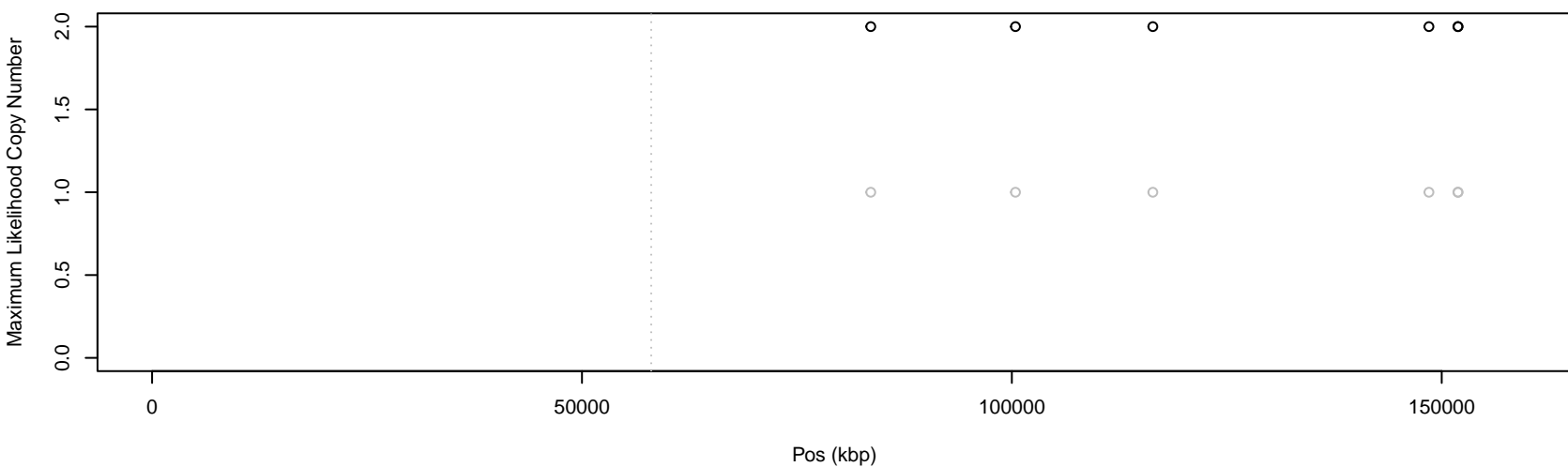
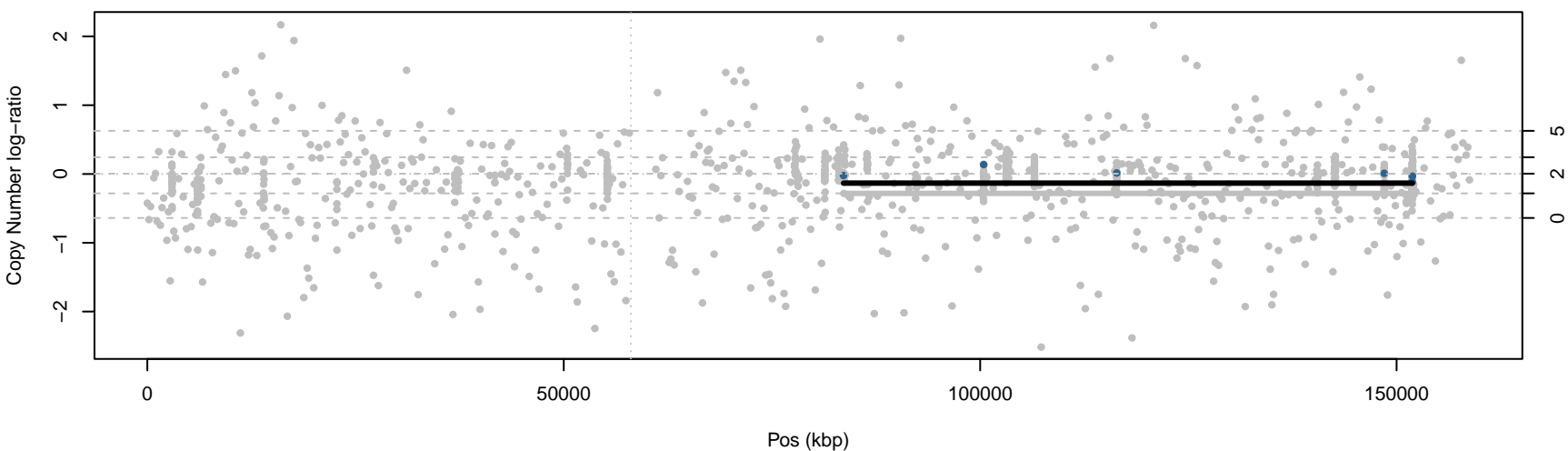
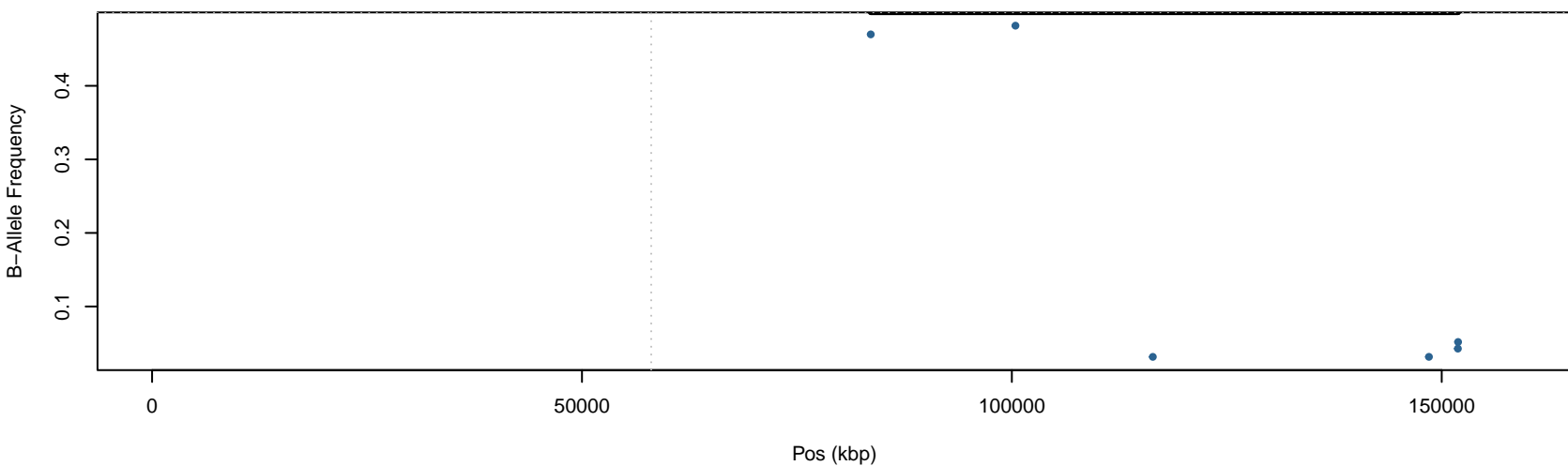
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 5



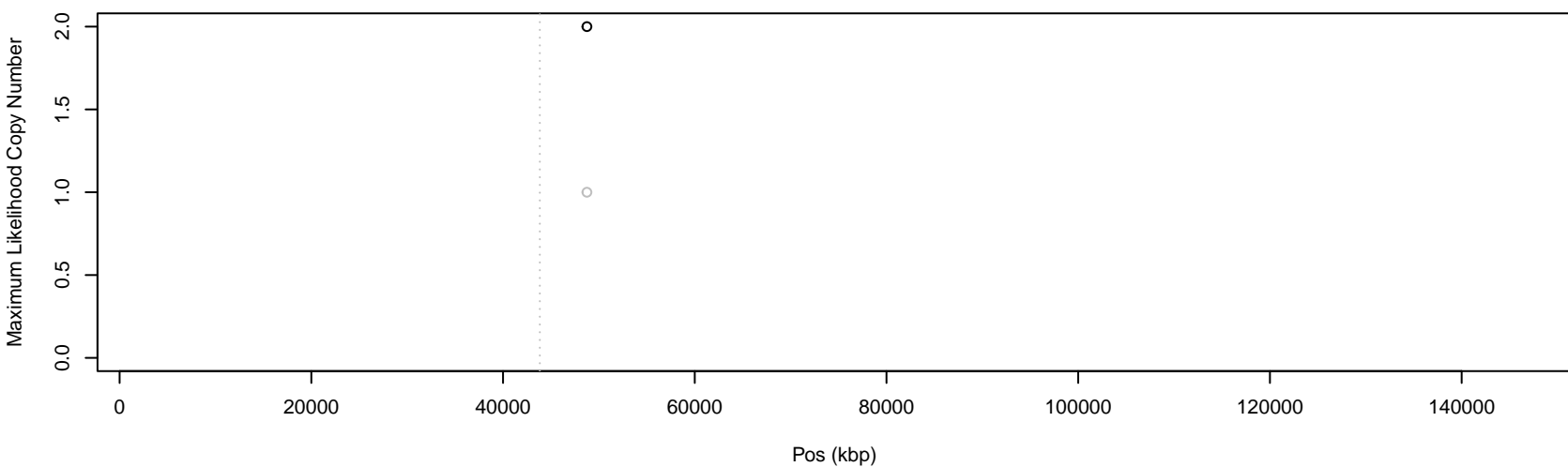
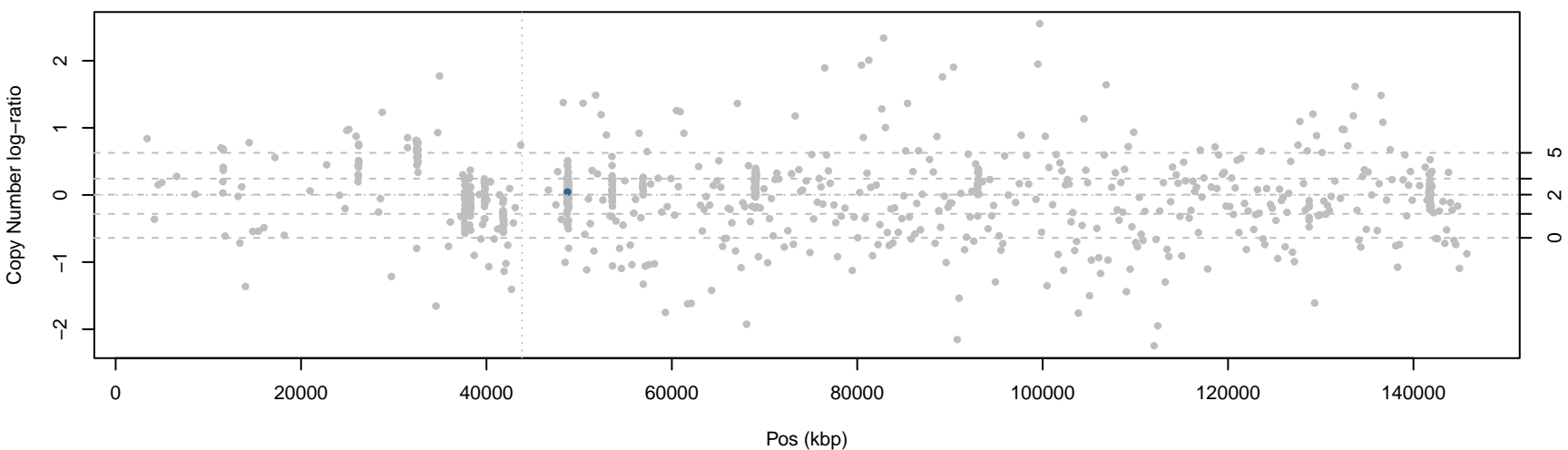
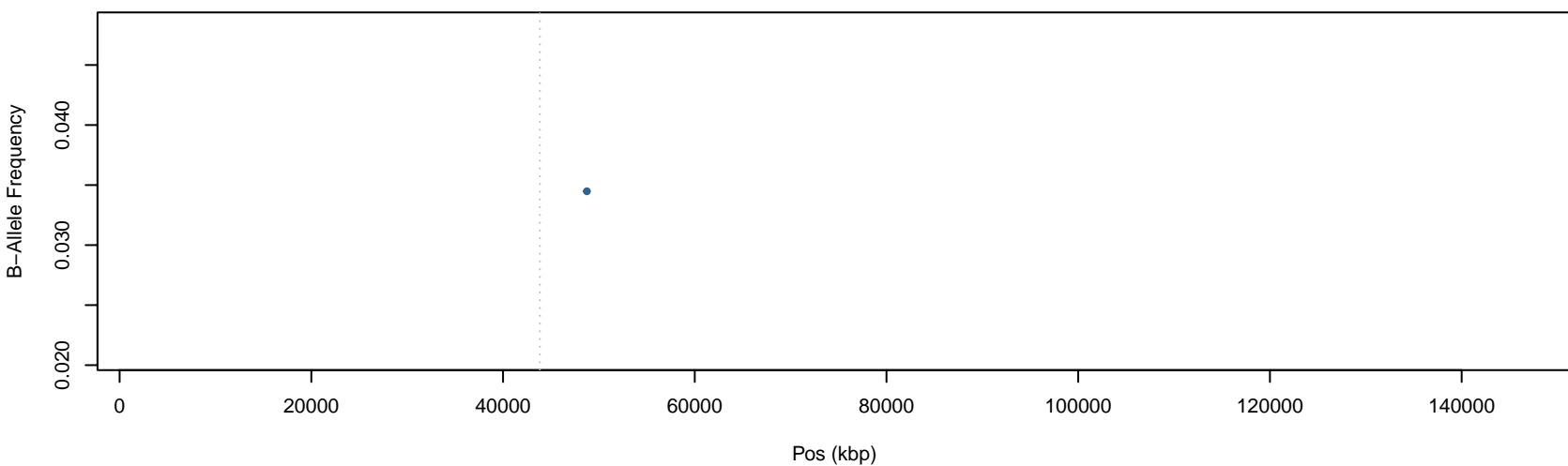
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 6



Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 7

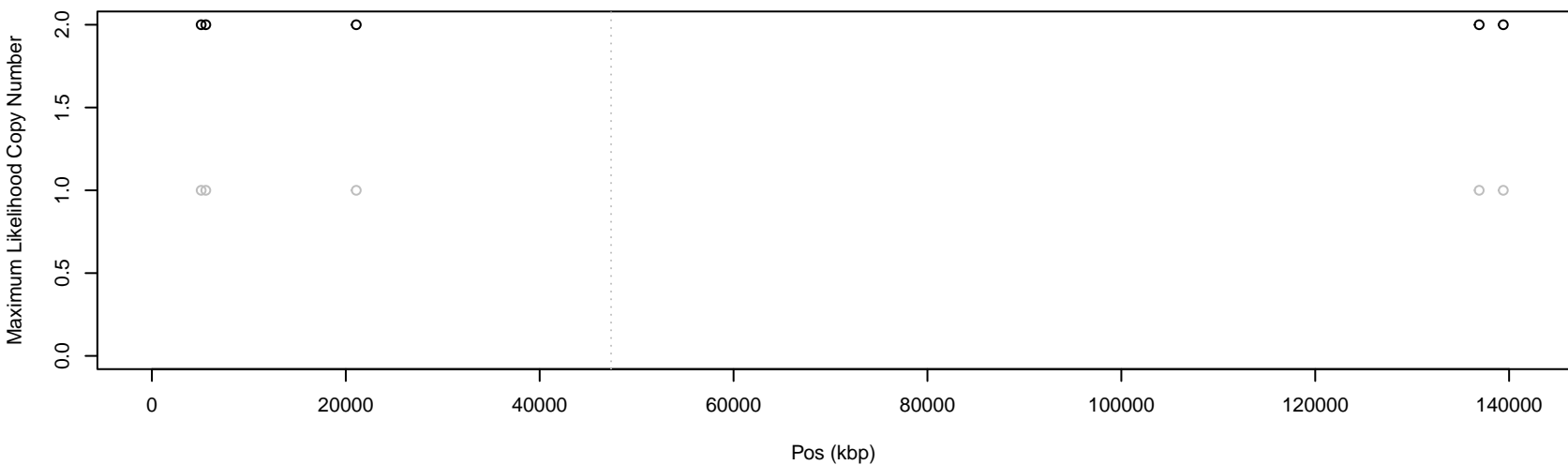
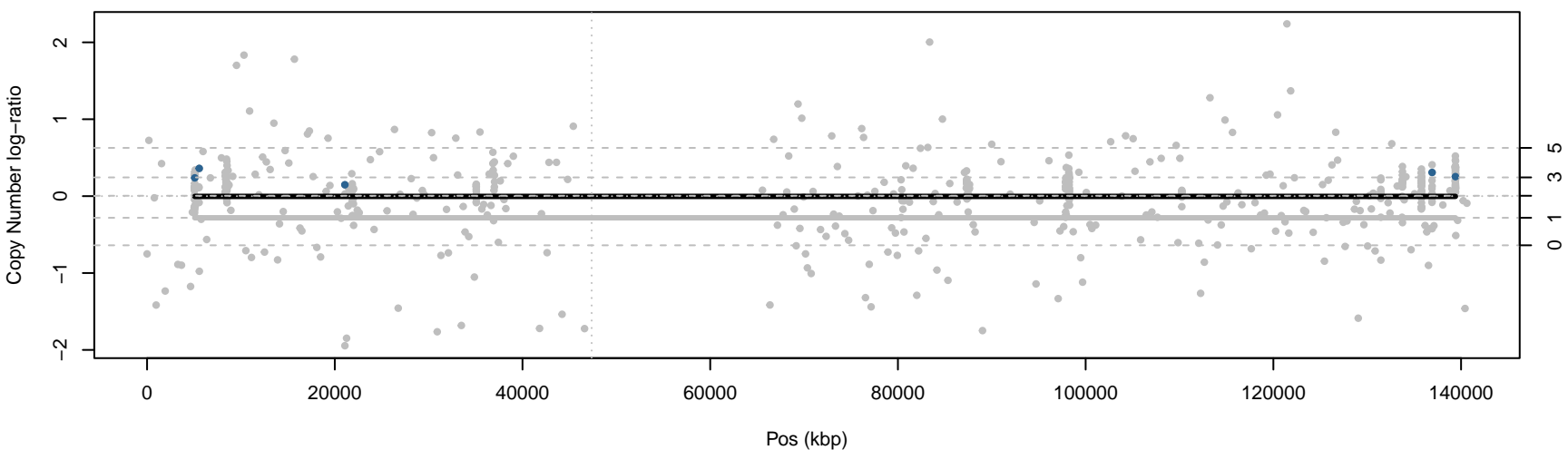
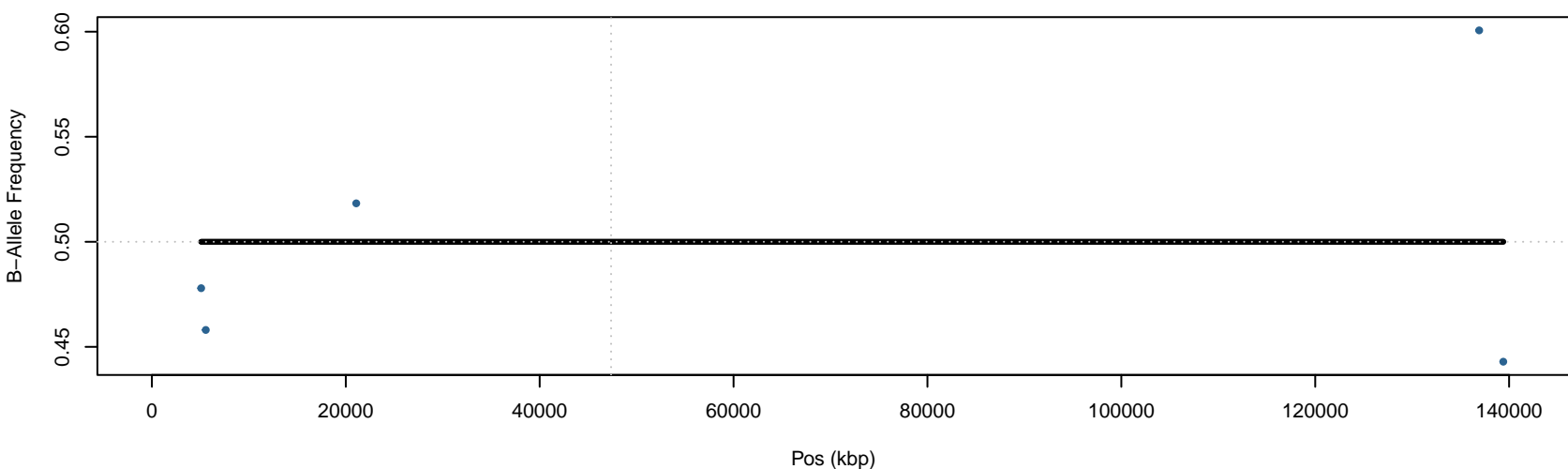


Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 8

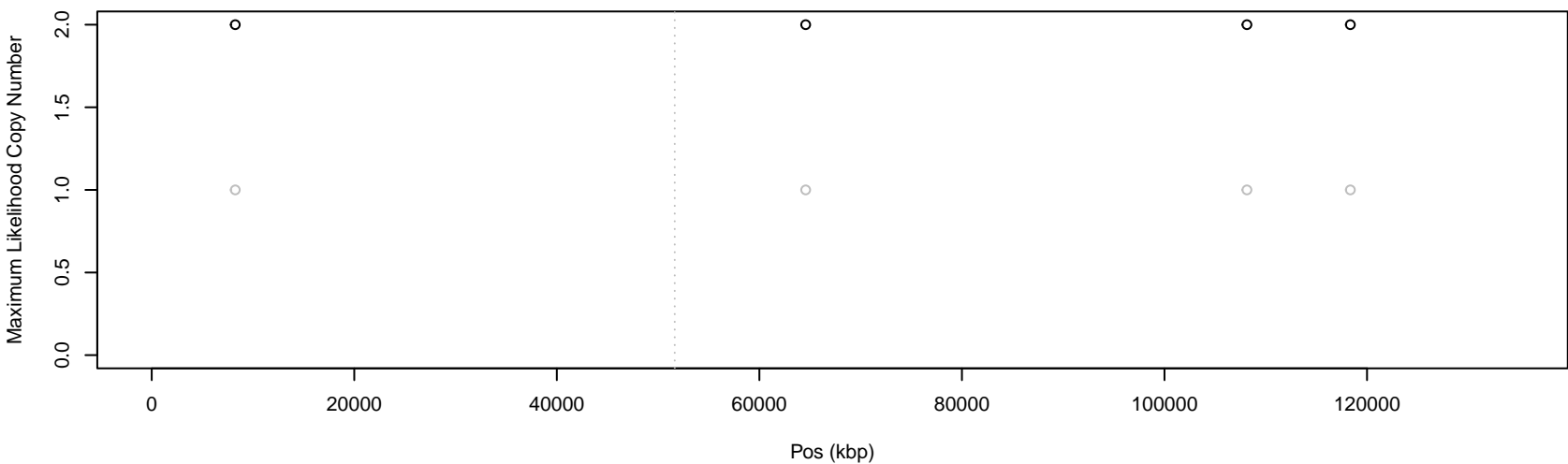
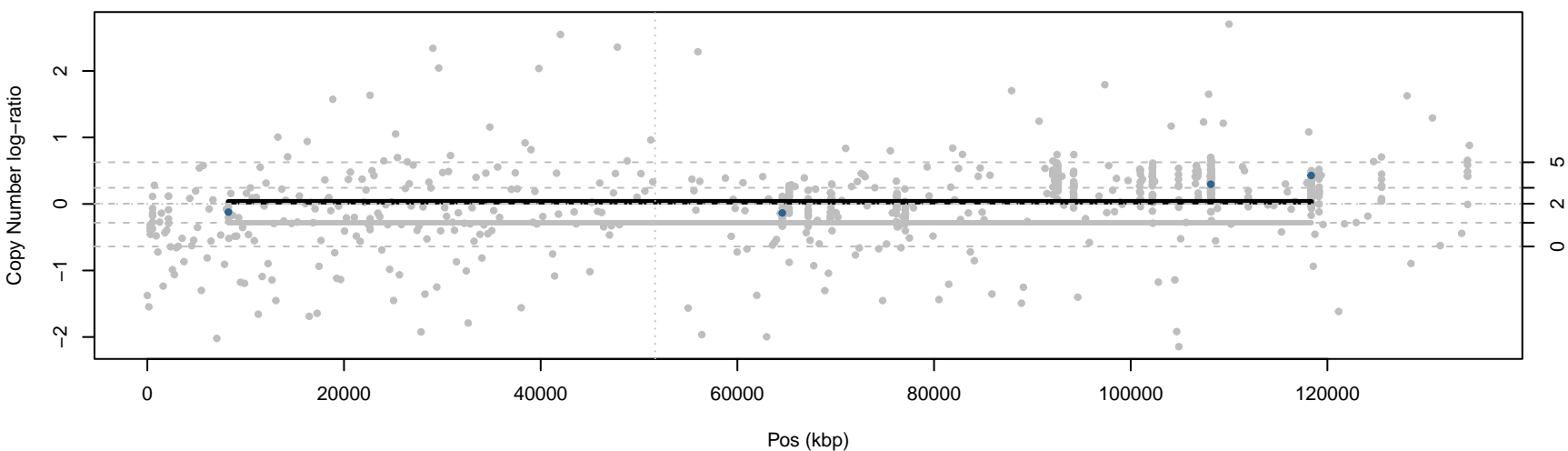
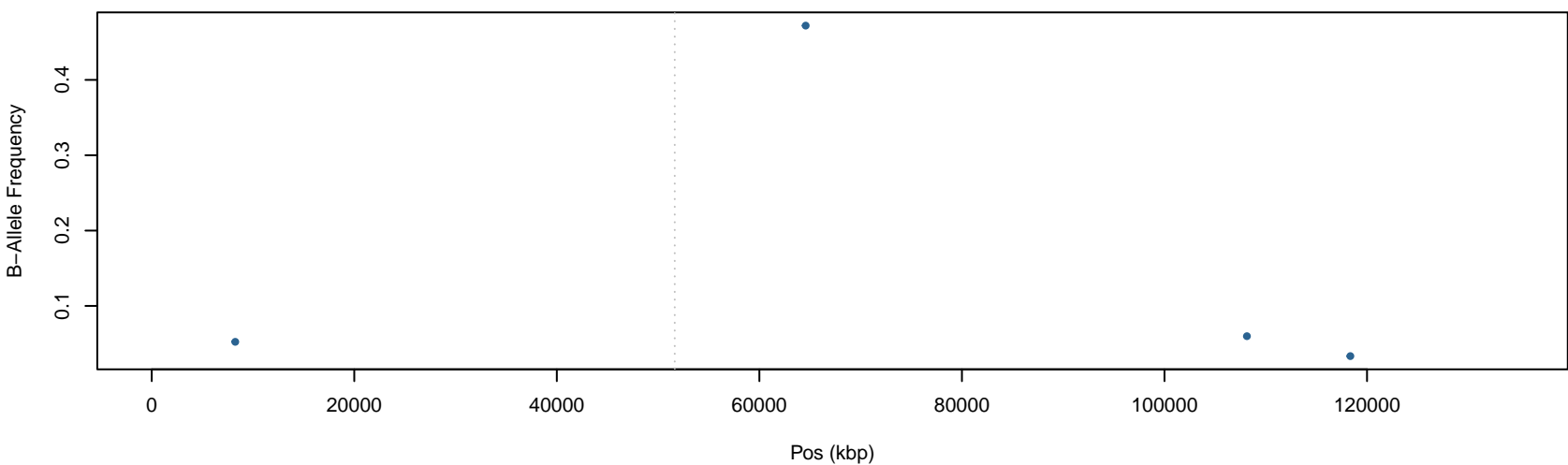




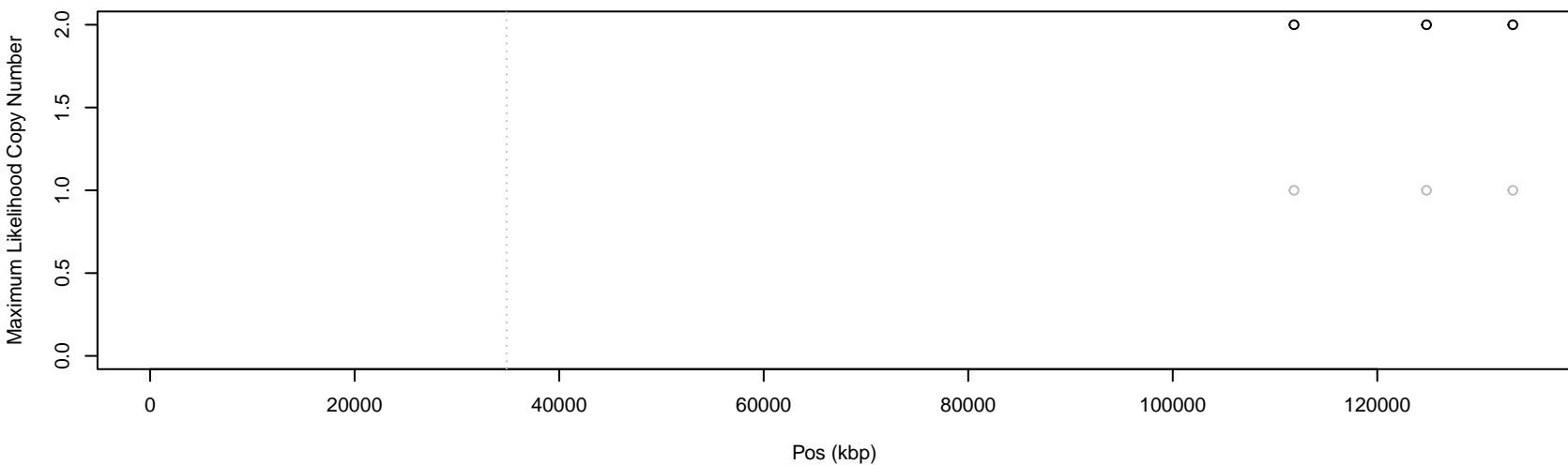
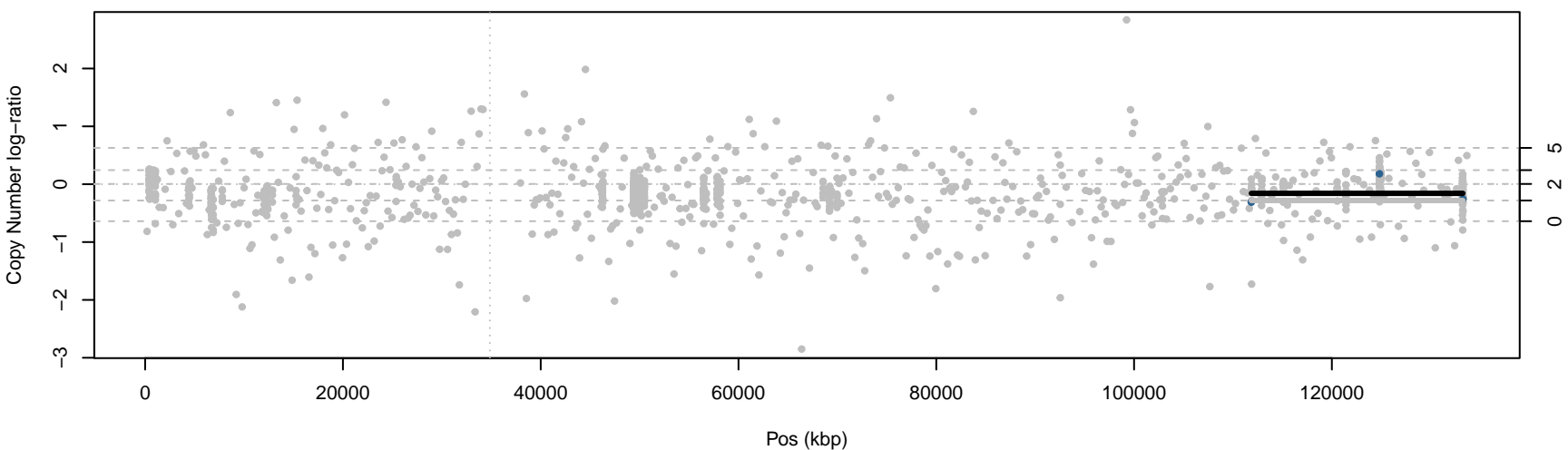
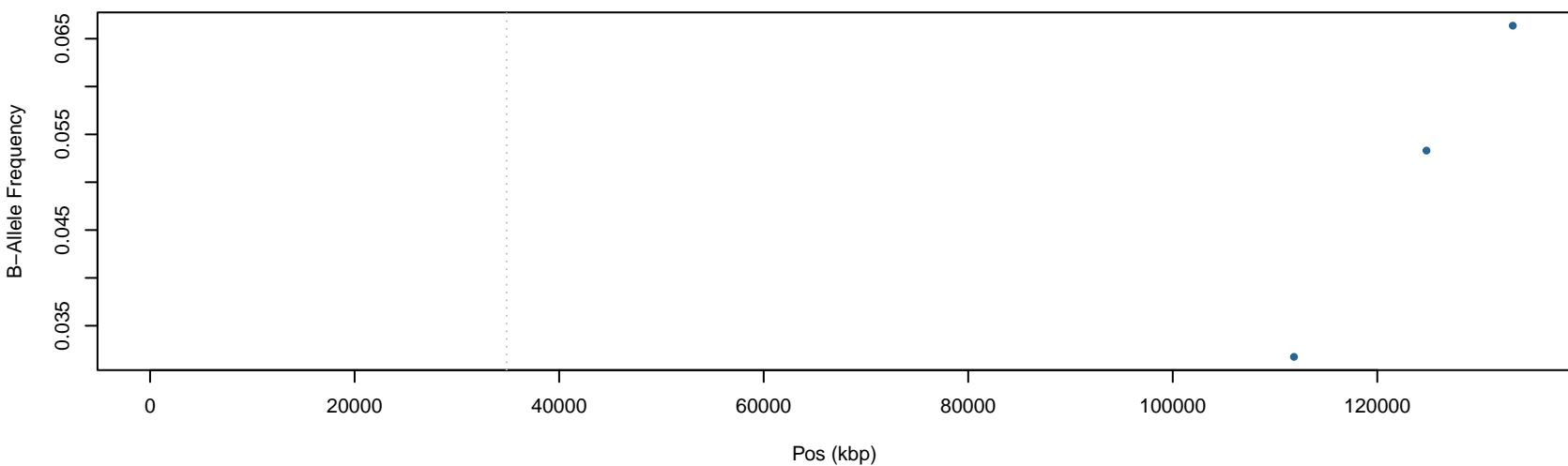
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 9



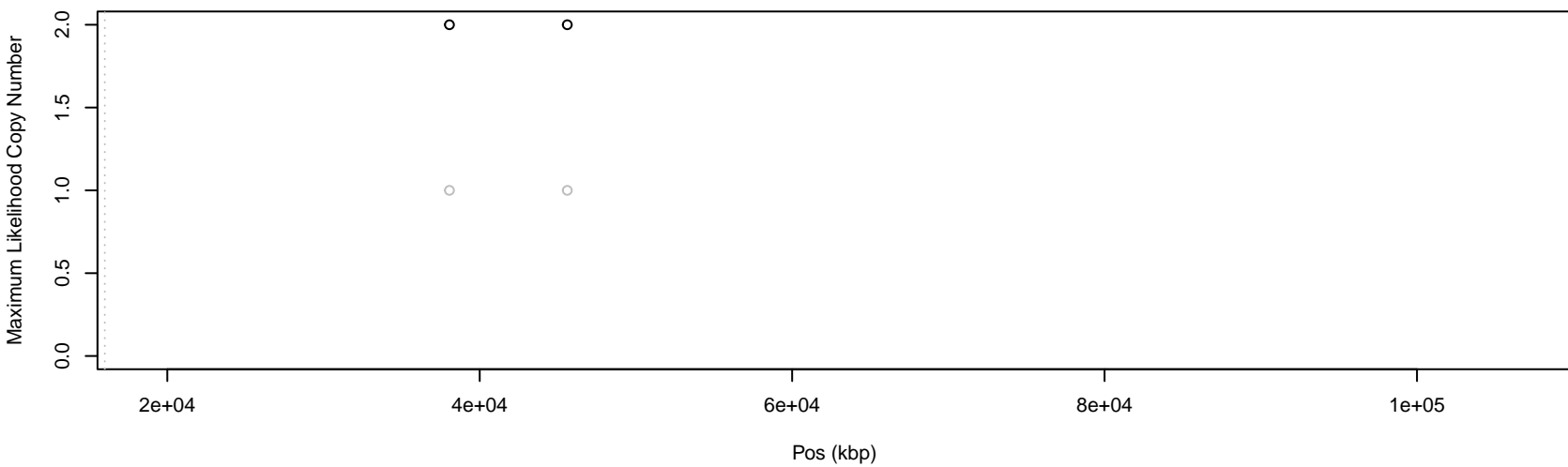
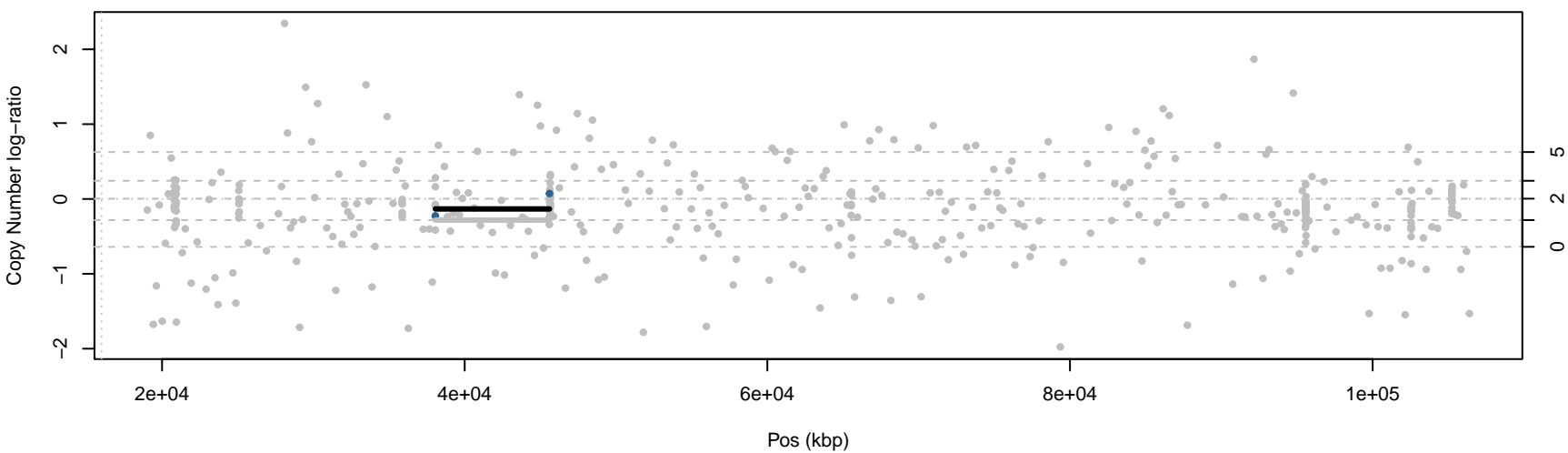
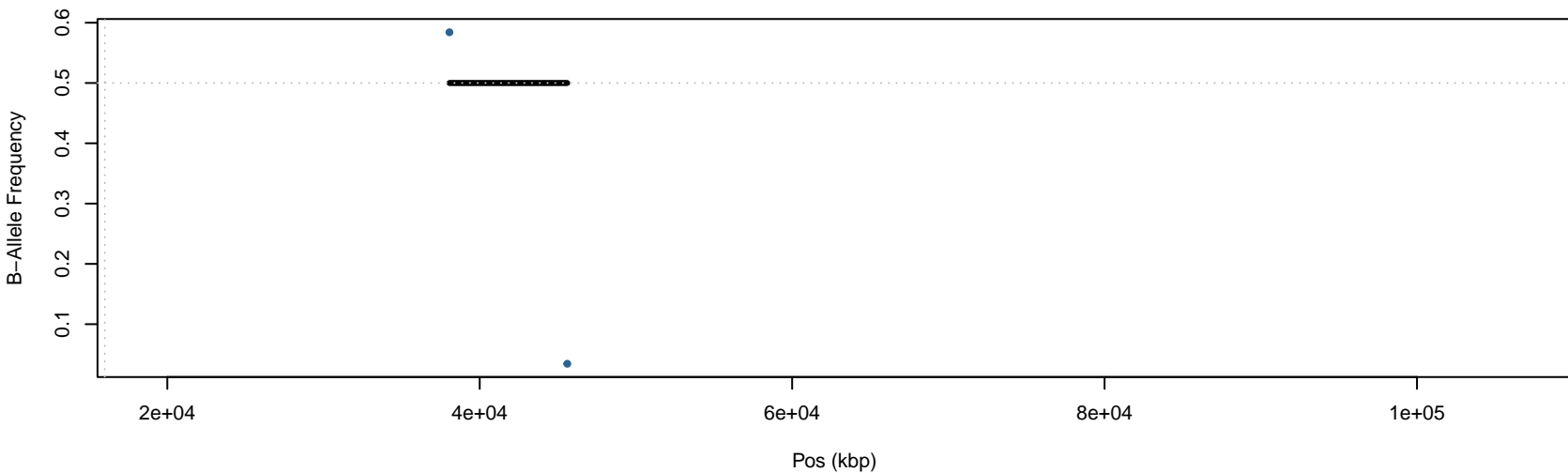
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 11



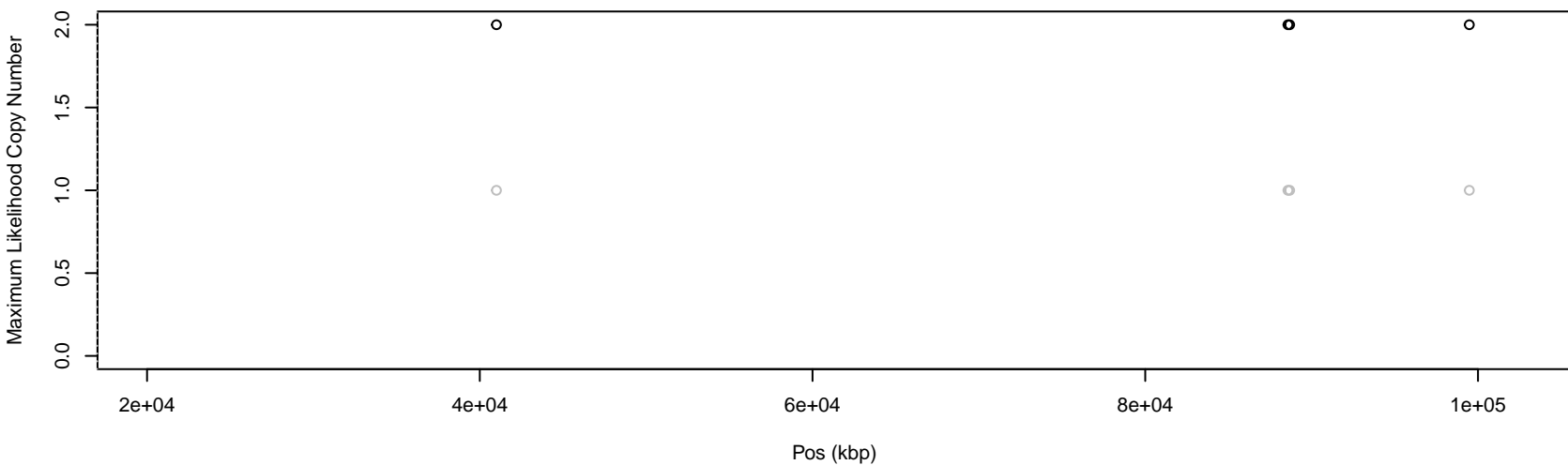
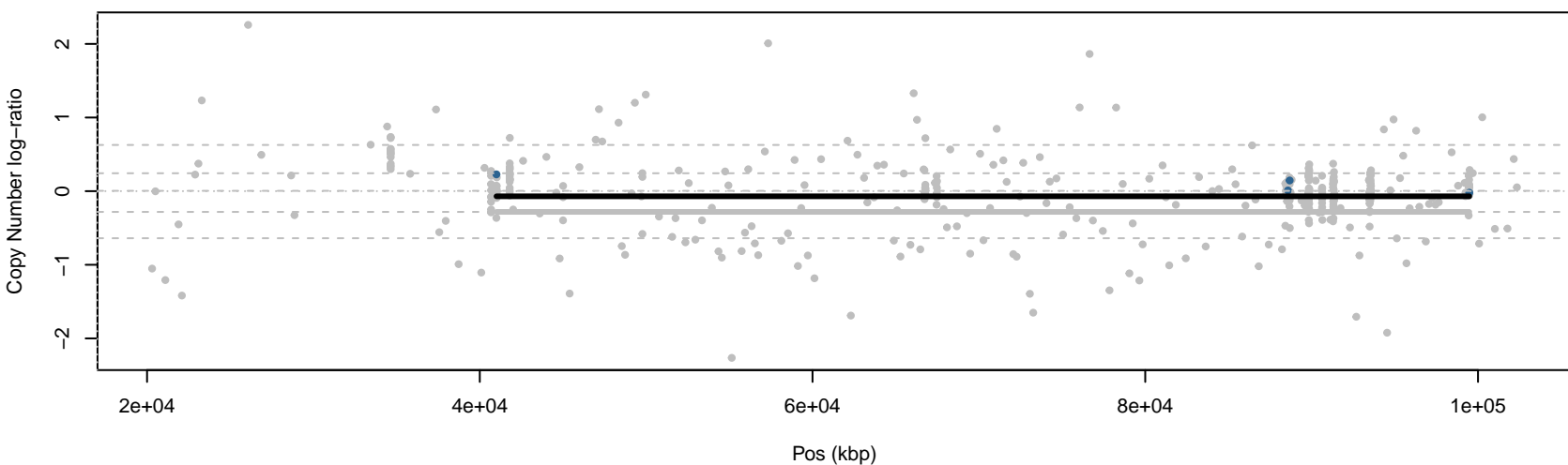
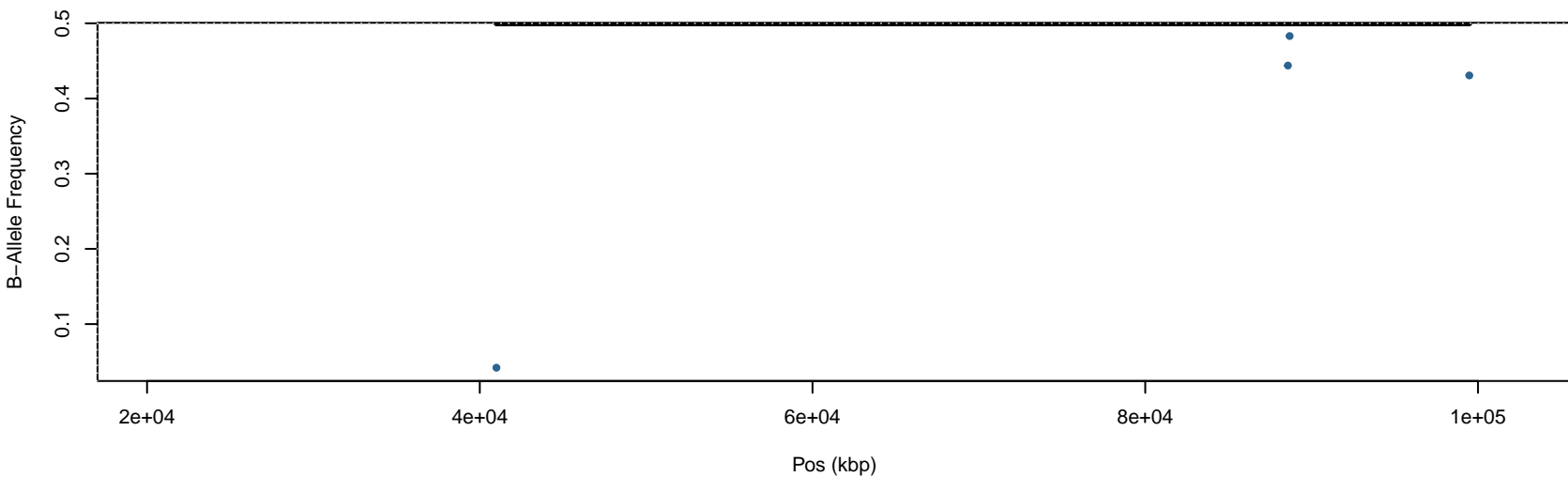
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 12



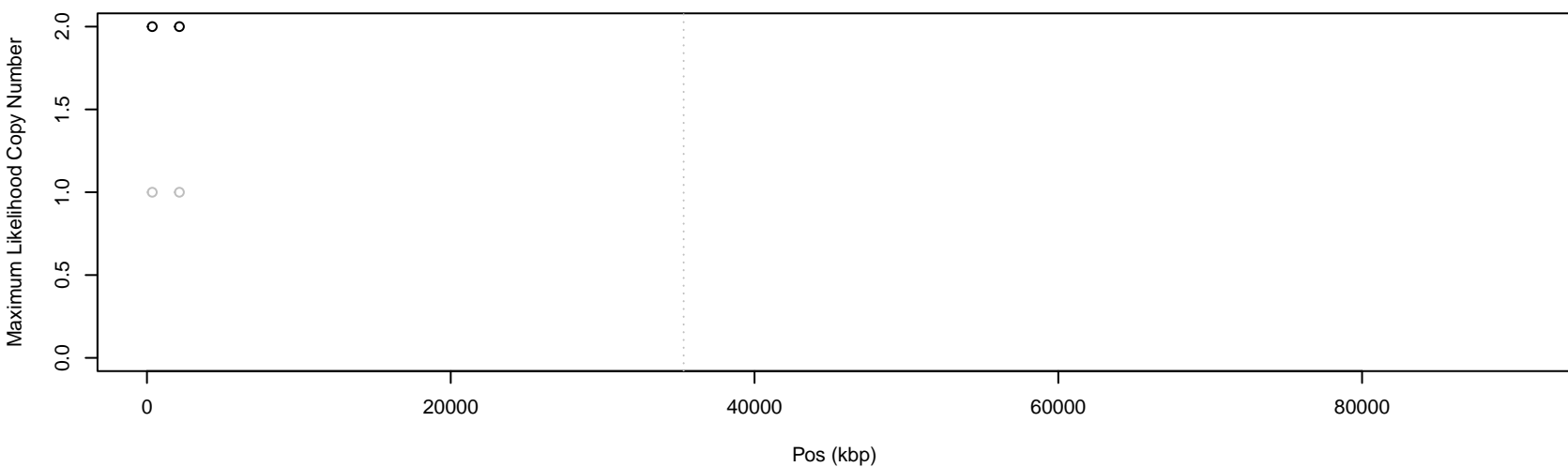
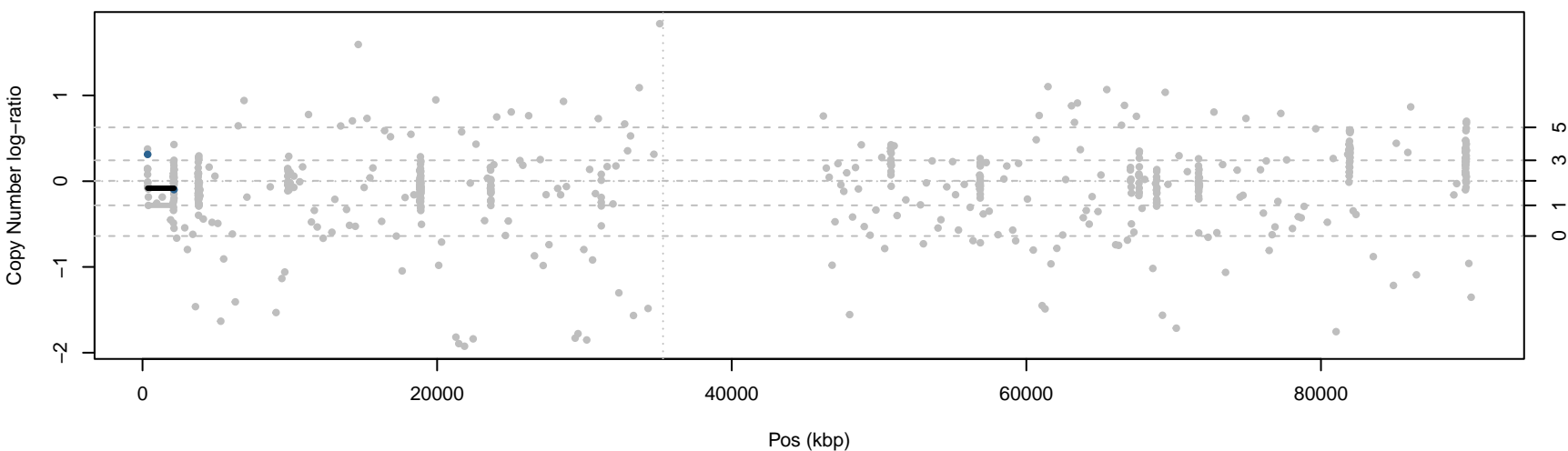
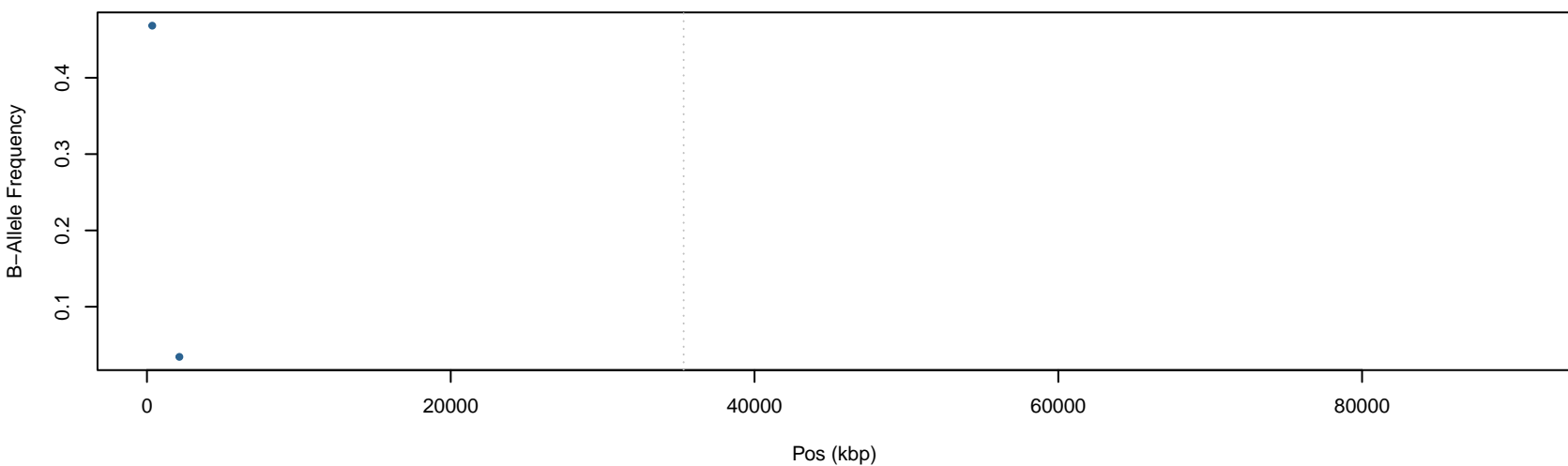
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 14



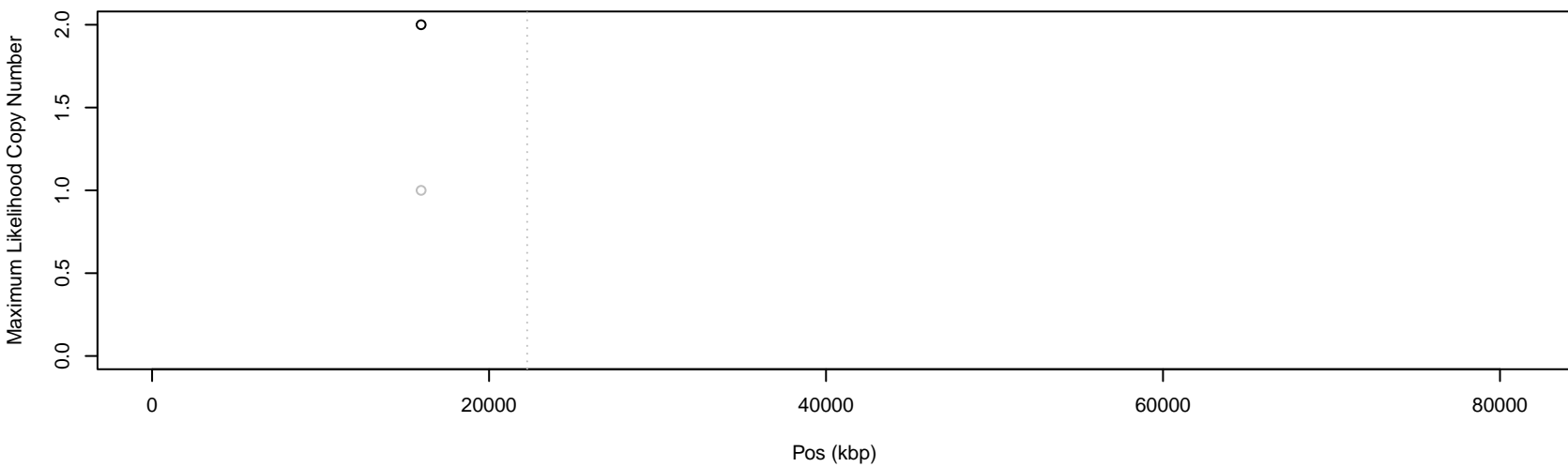
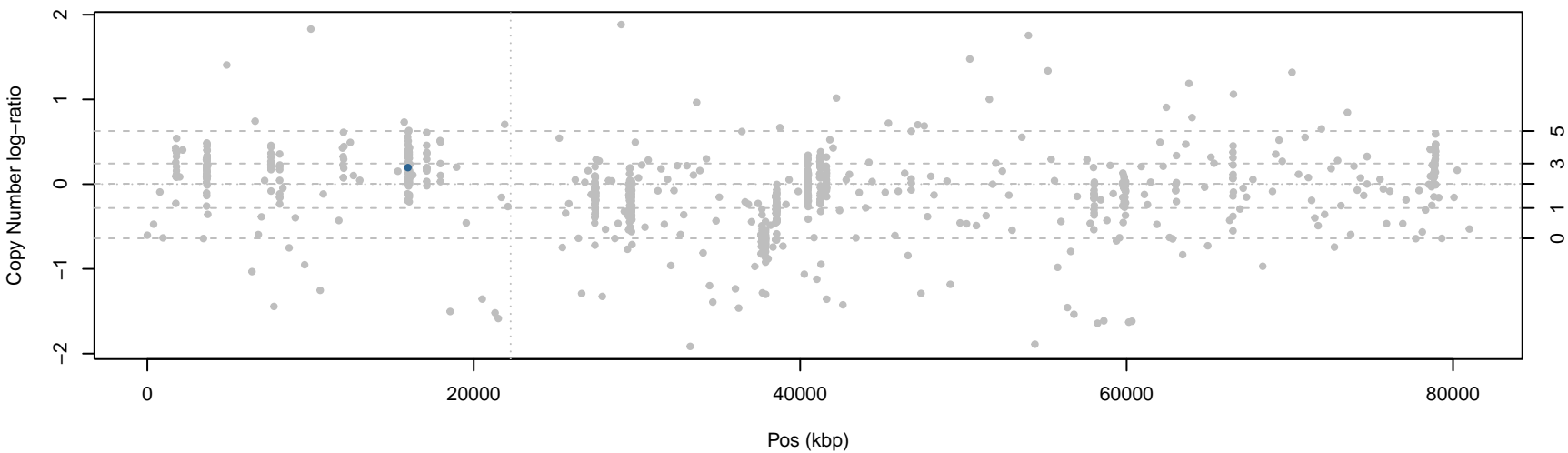
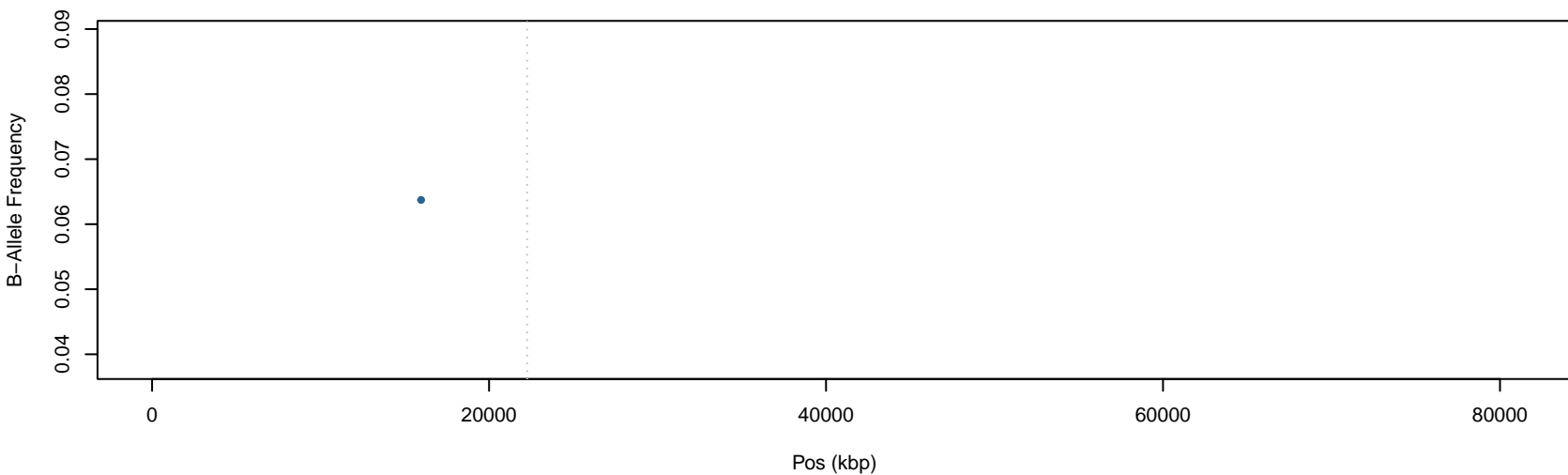
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 15



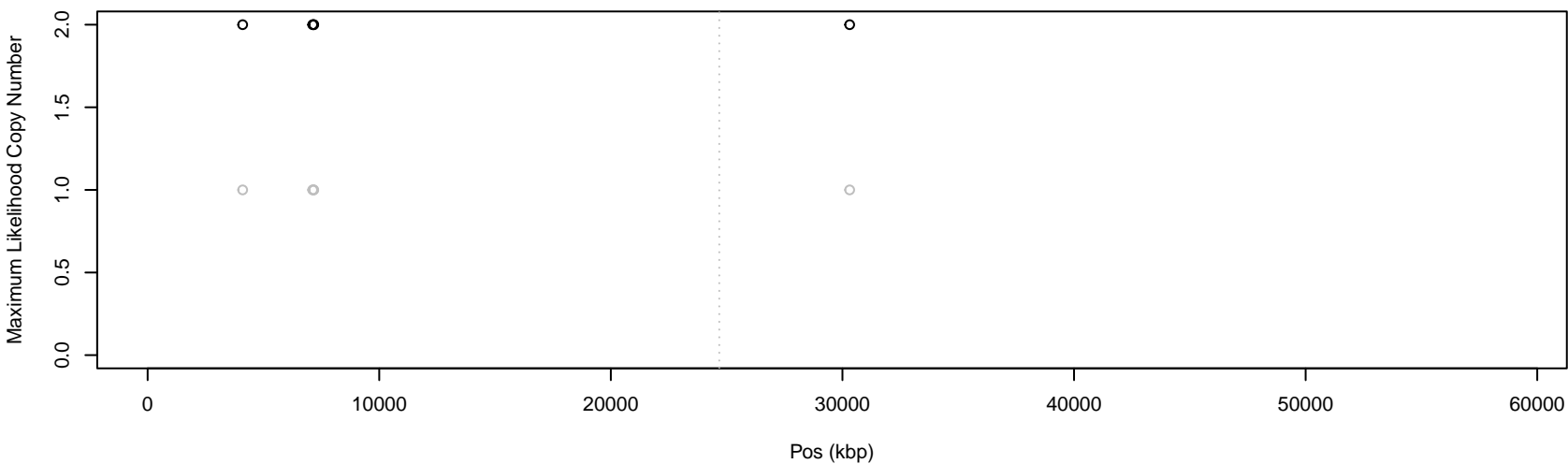
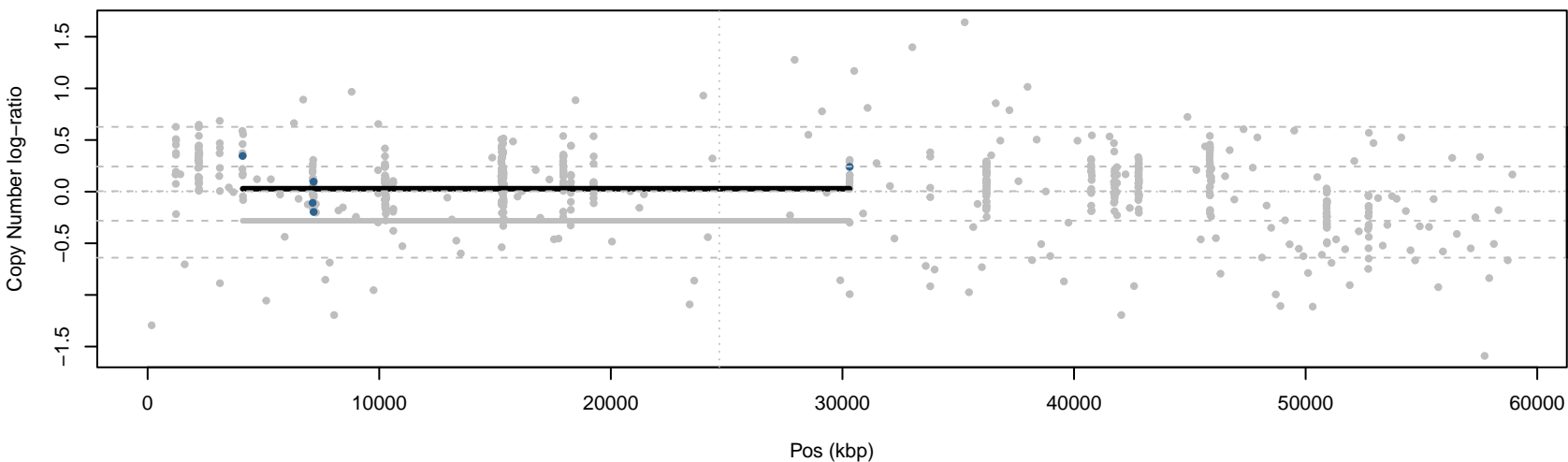
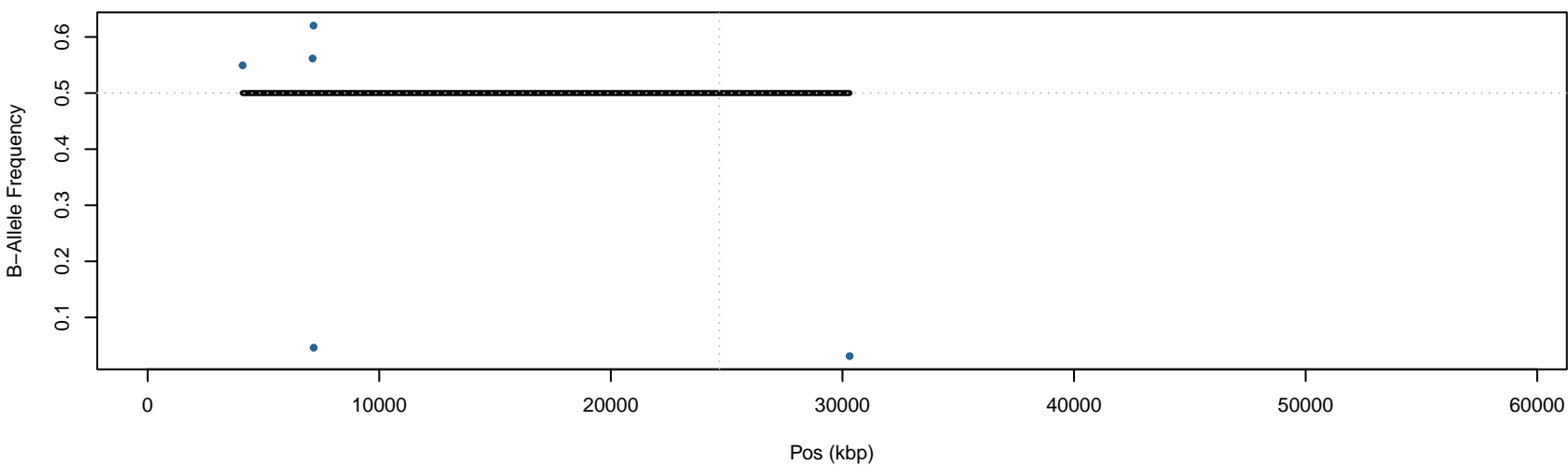
Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 16



Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 17

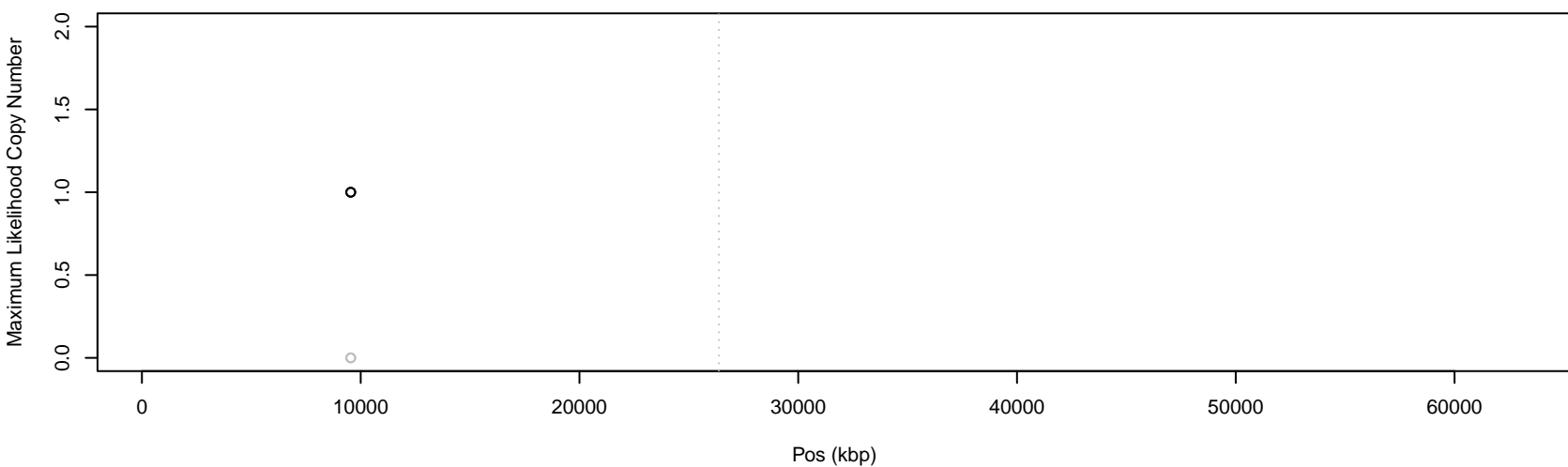
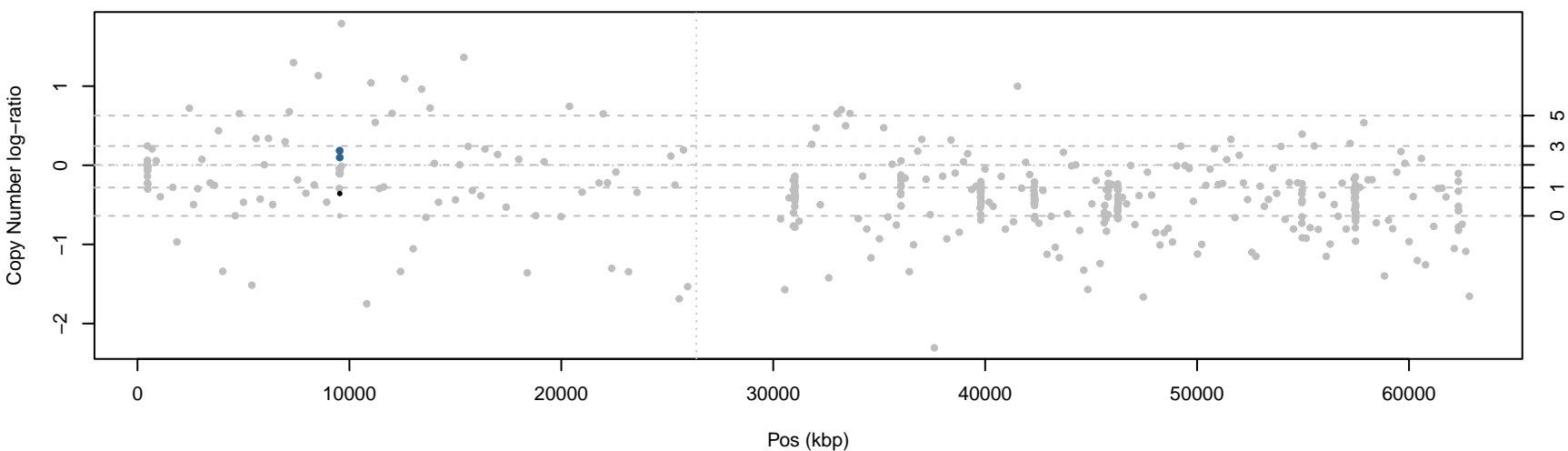
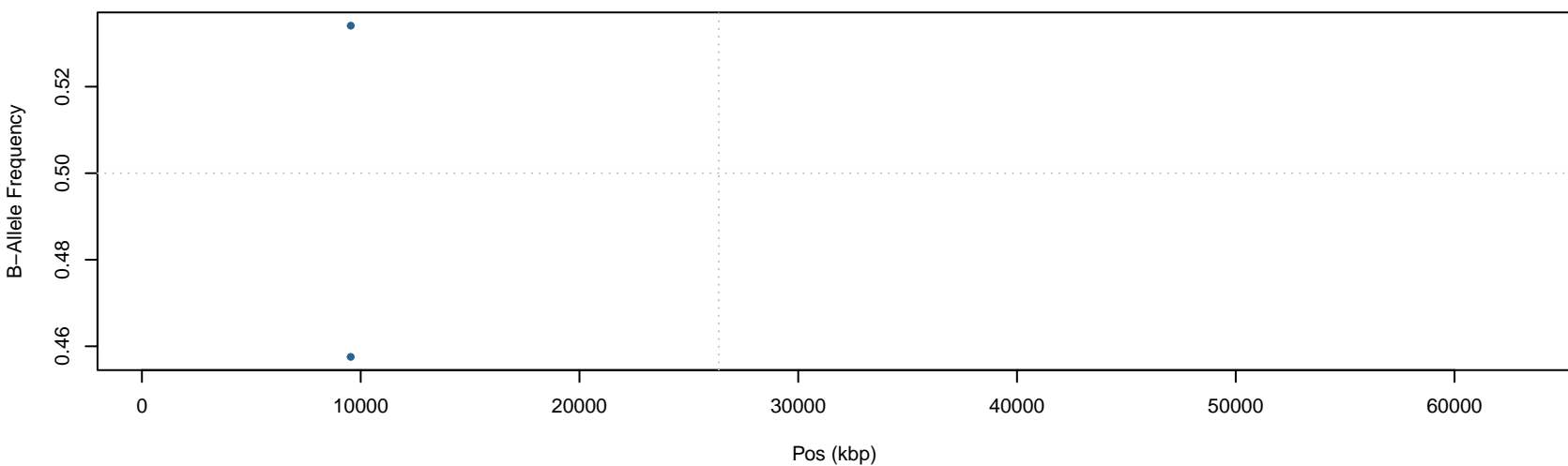


Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 19

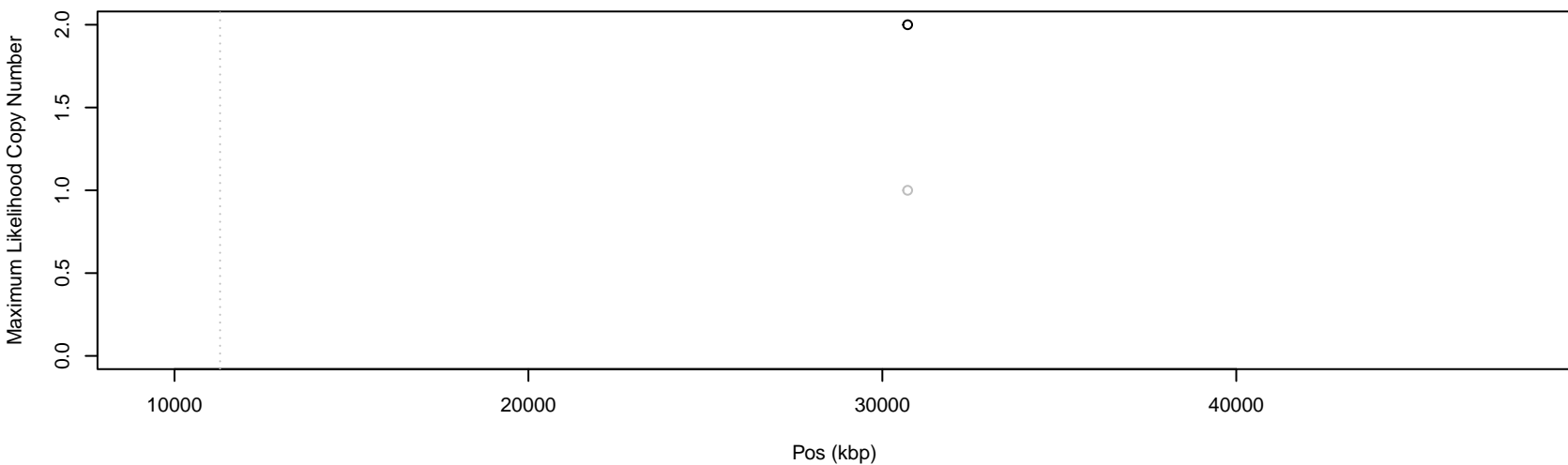
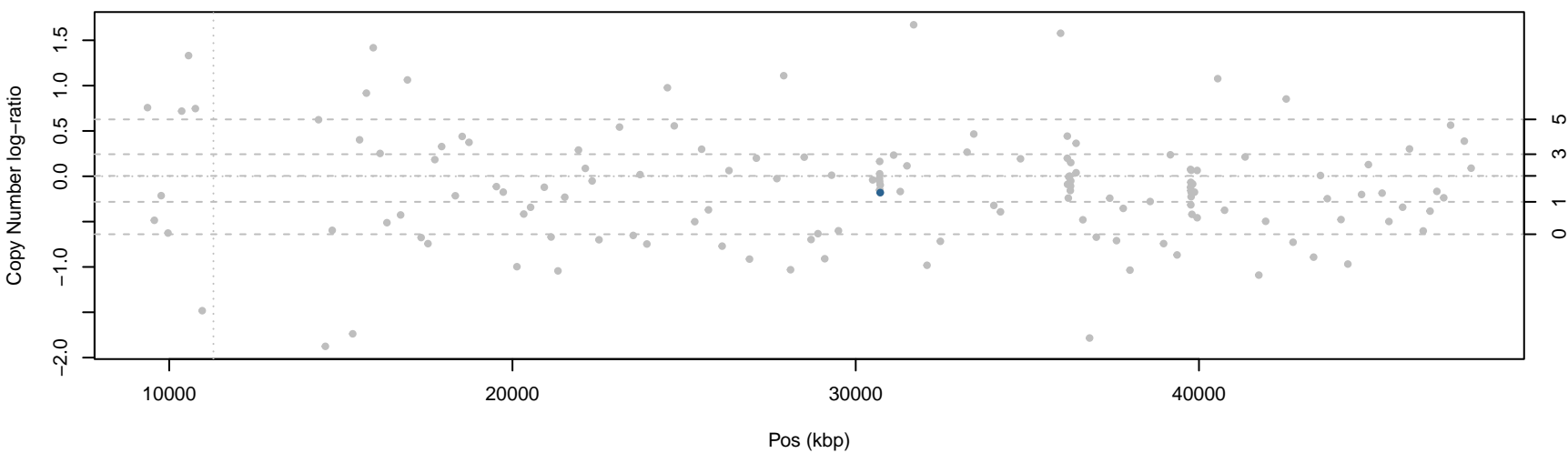
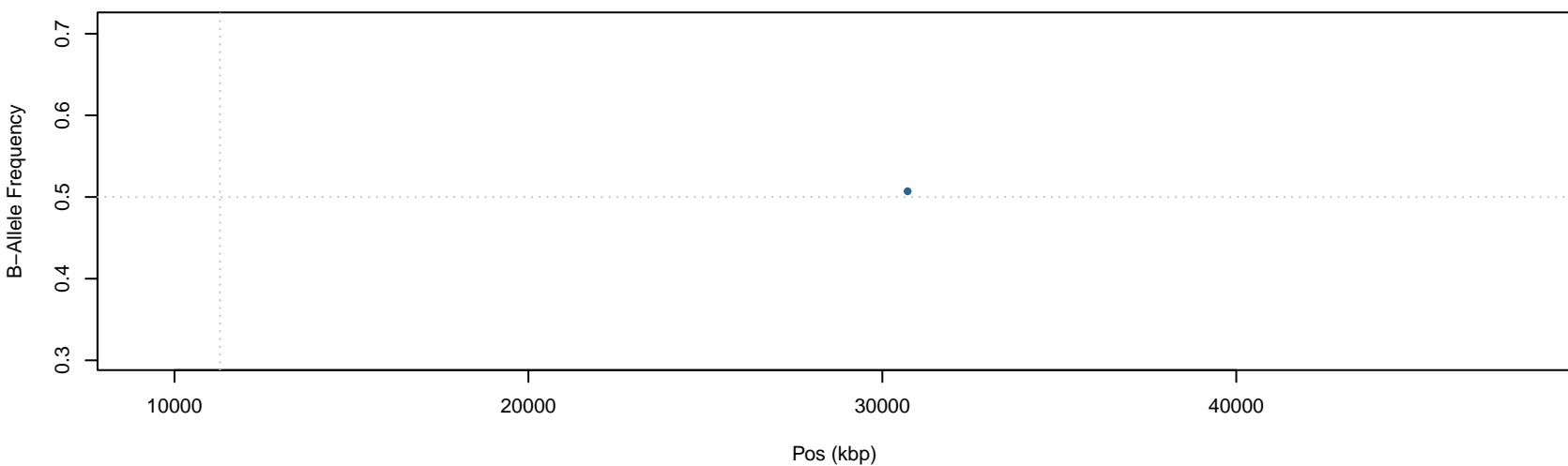




Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 20



Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 21



Purity: 0.36 Tumor ploidy: 1.983 SNV log-likelihood: -10.69 GoF: 74.1% Mean coverage: 493;307 Chromosome: 22

