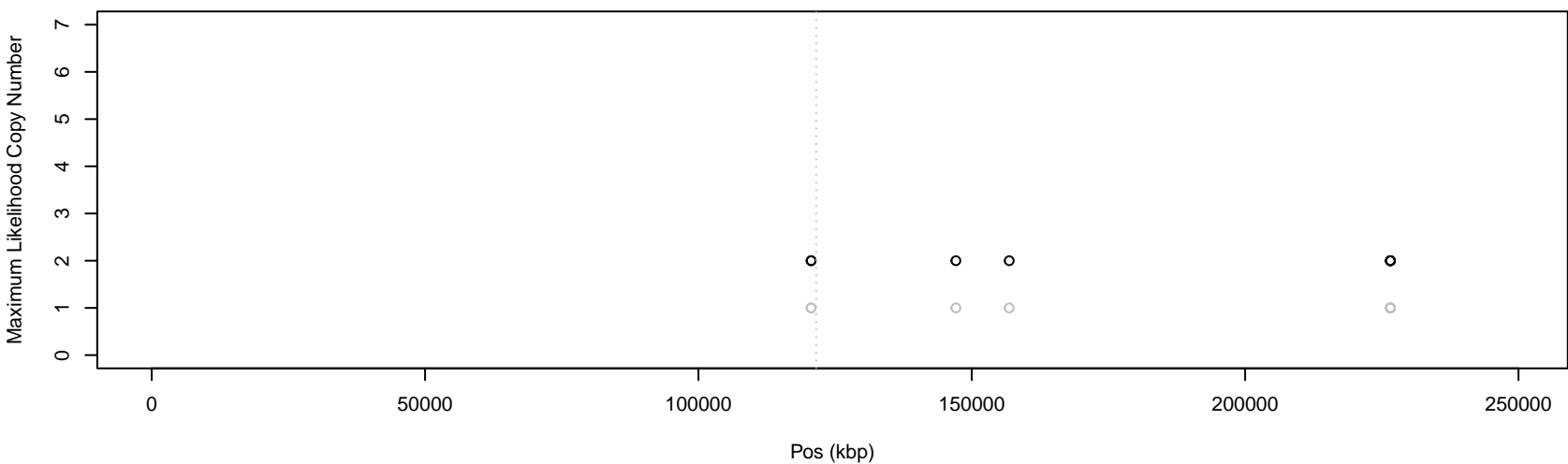
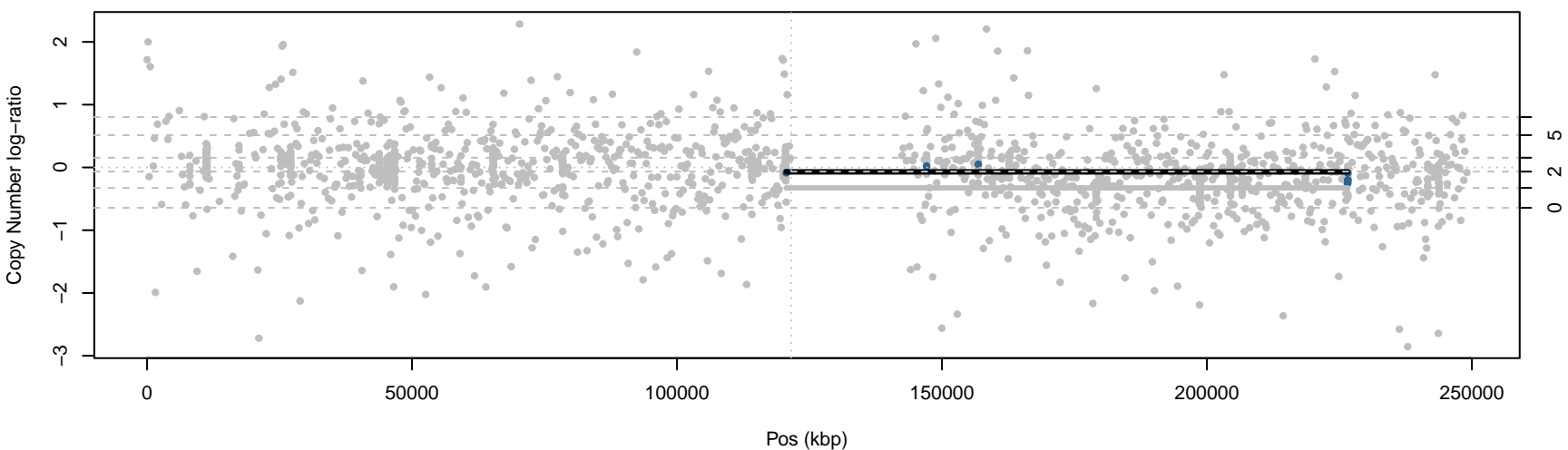
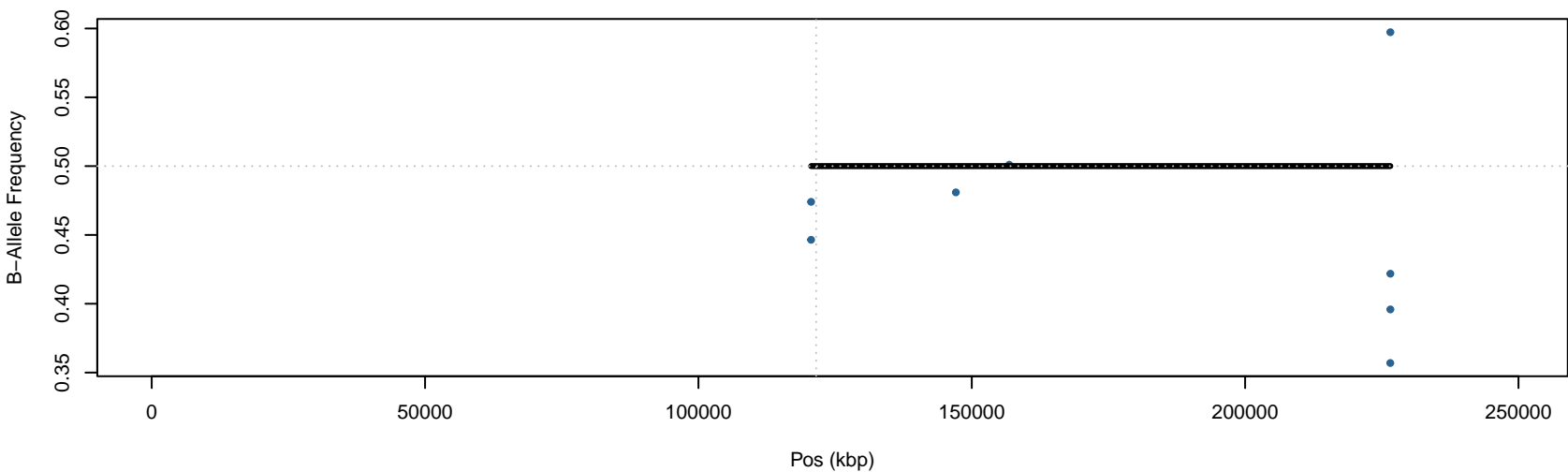
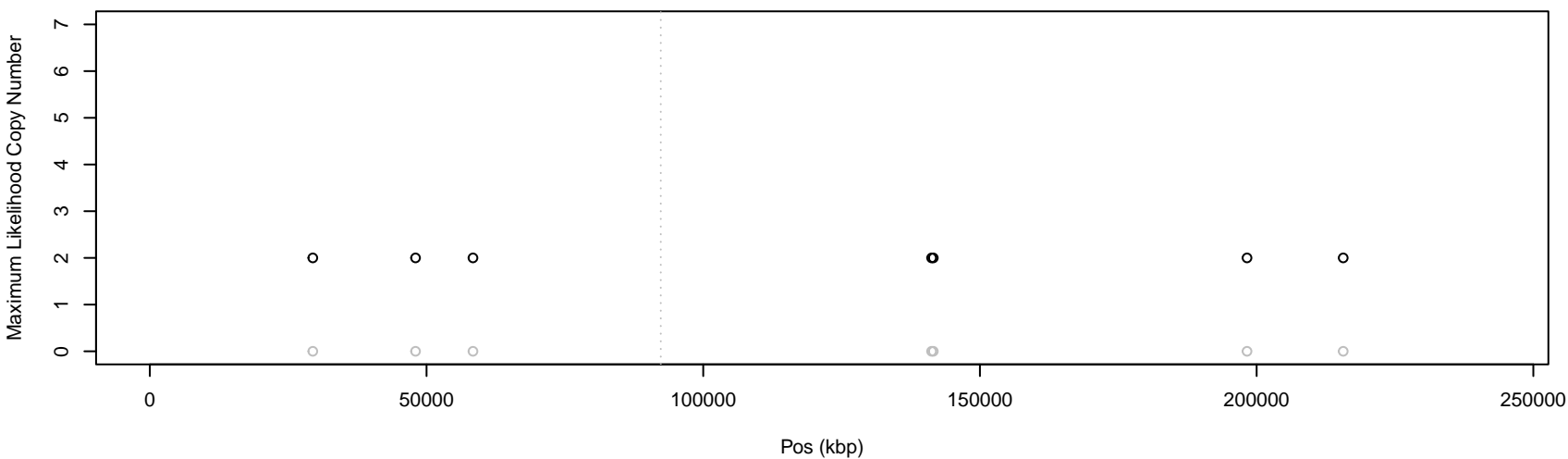
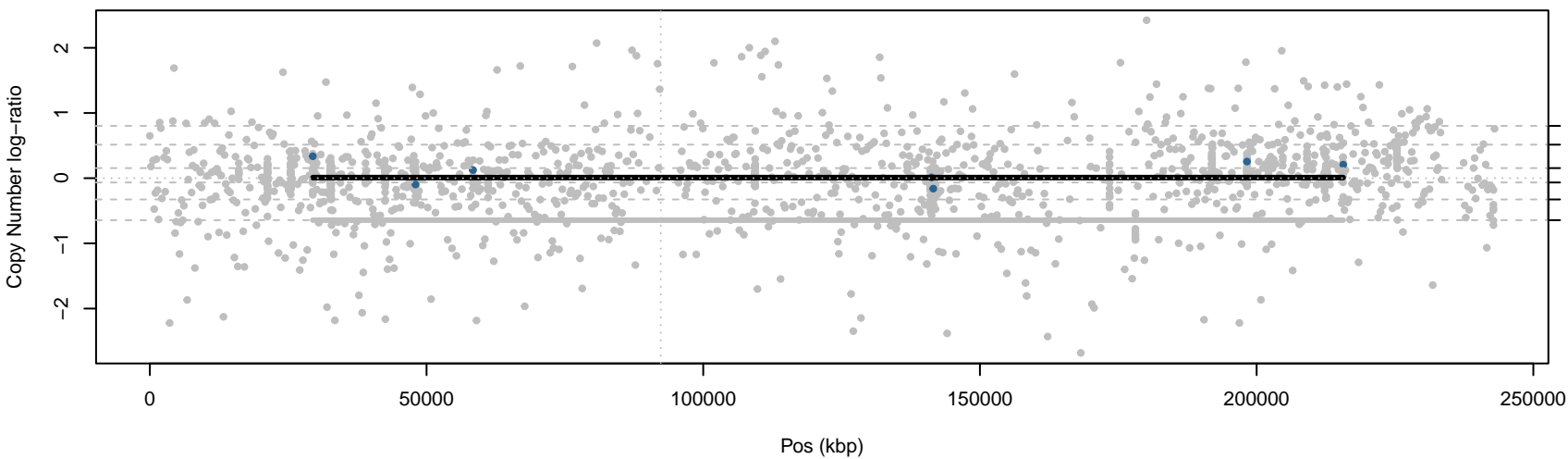
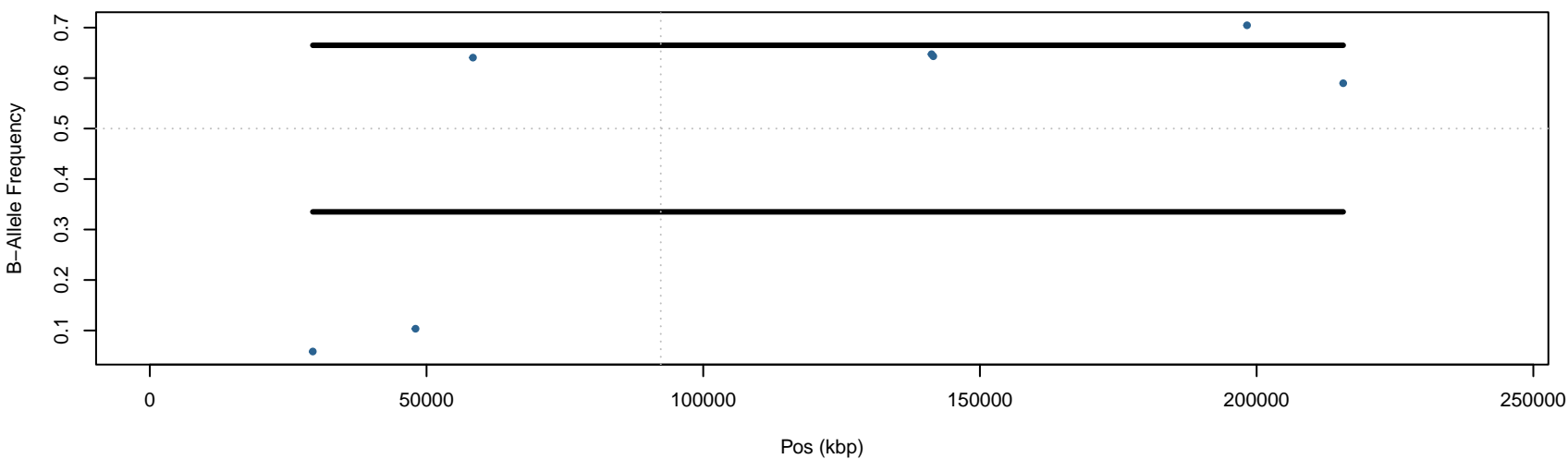


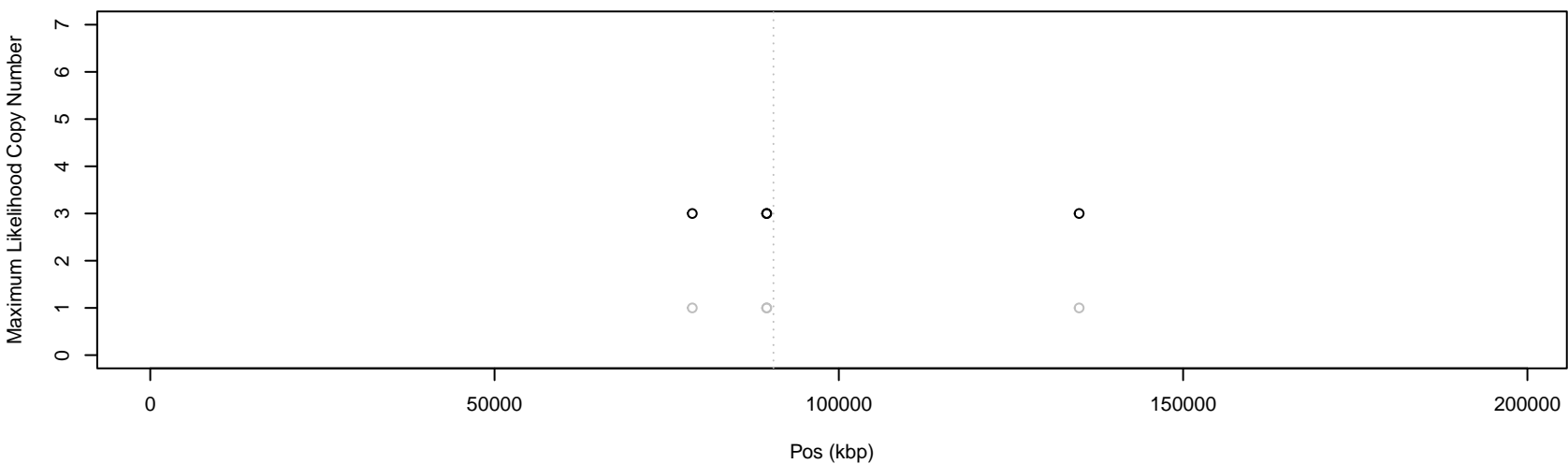
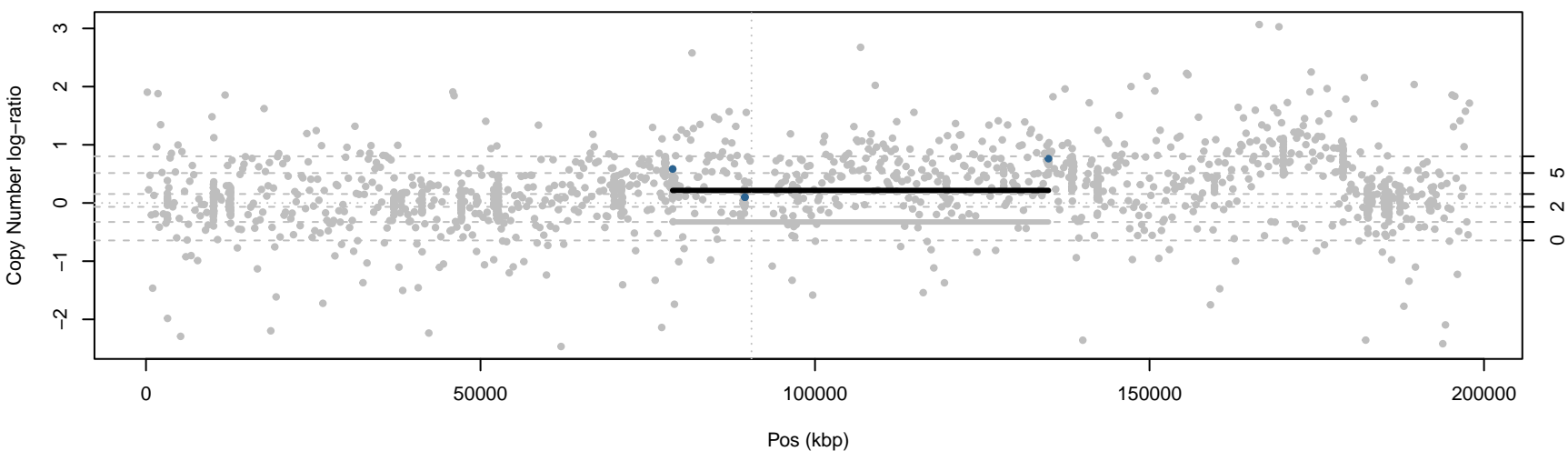
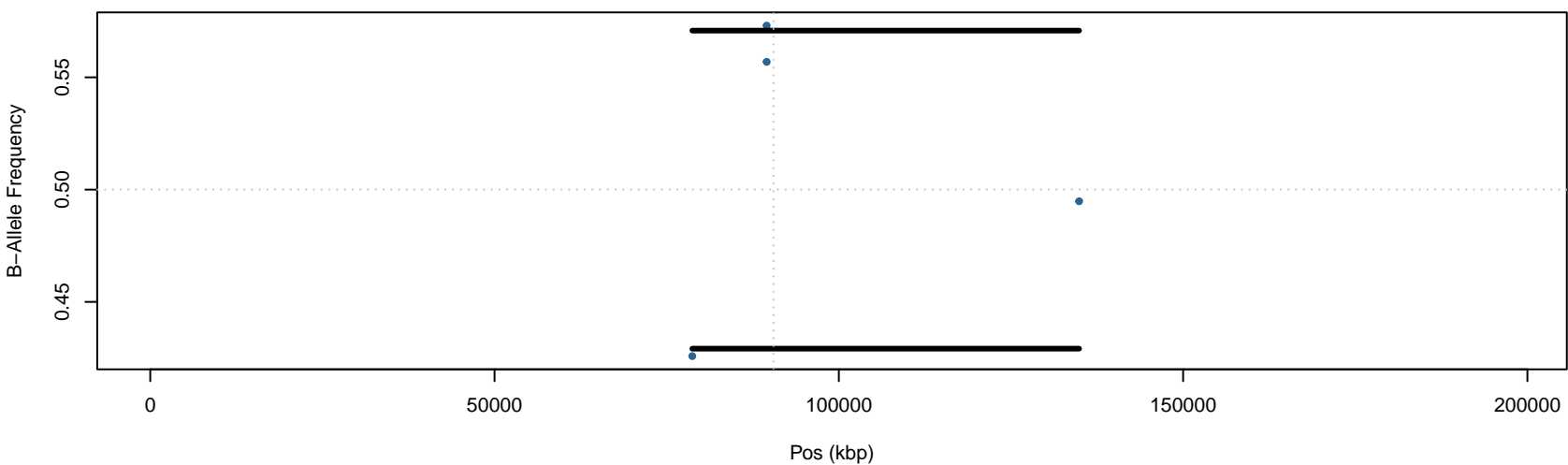
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 1



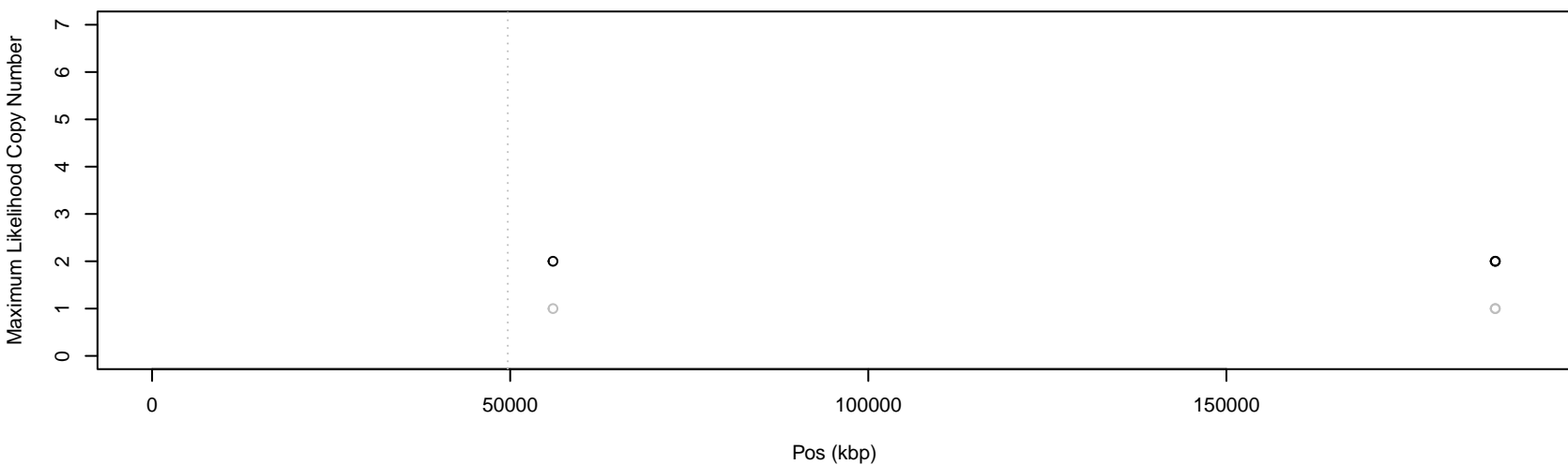
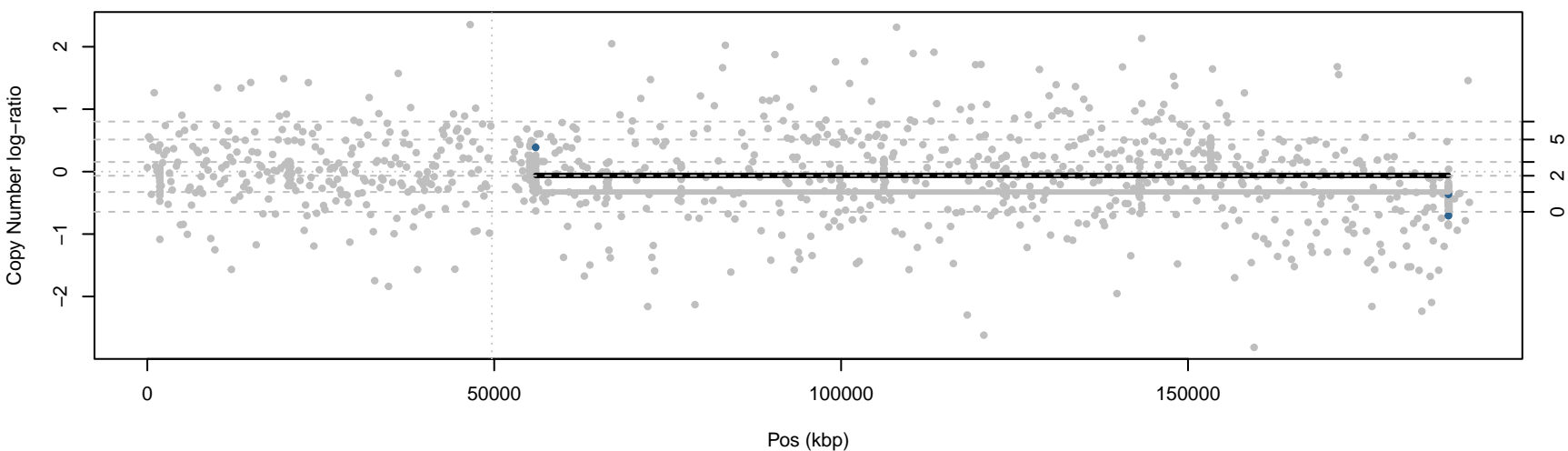
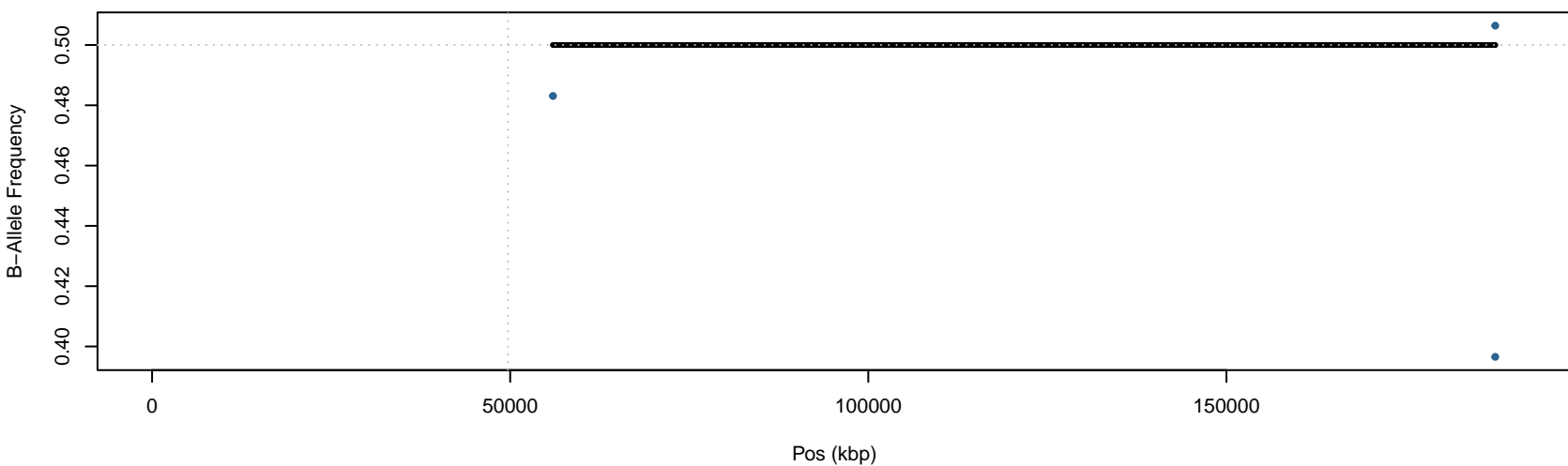
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 2



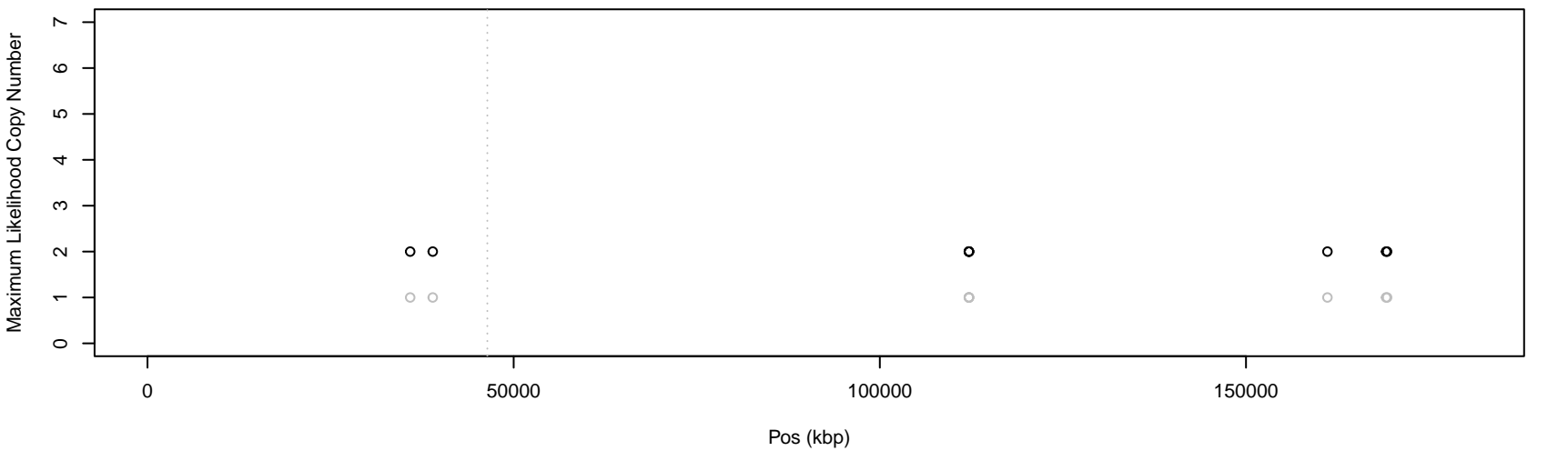
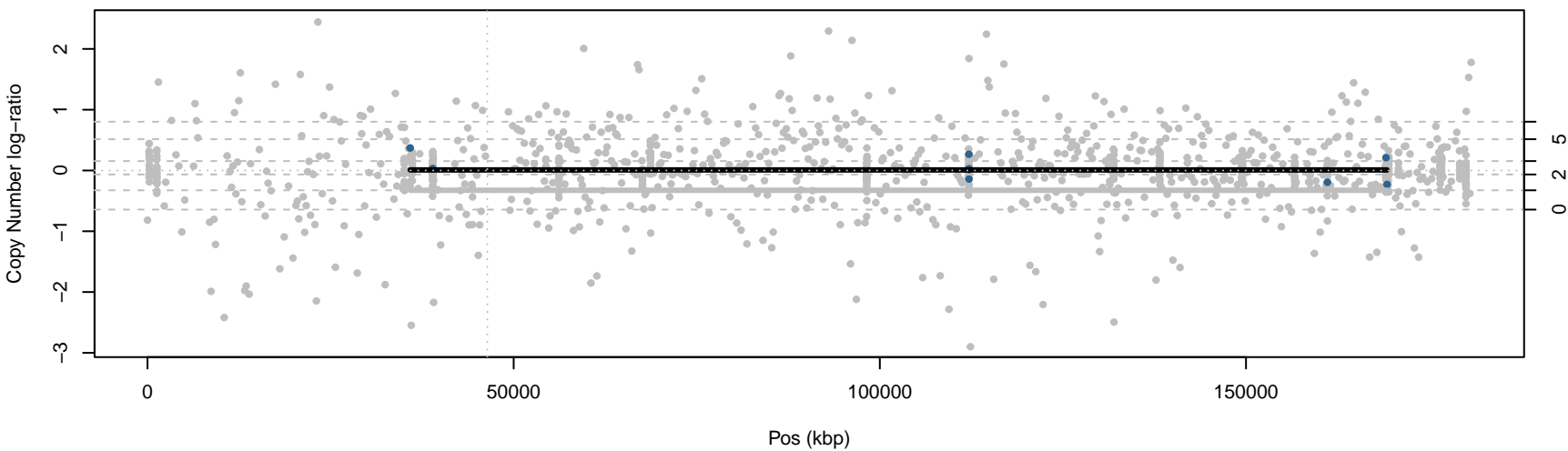
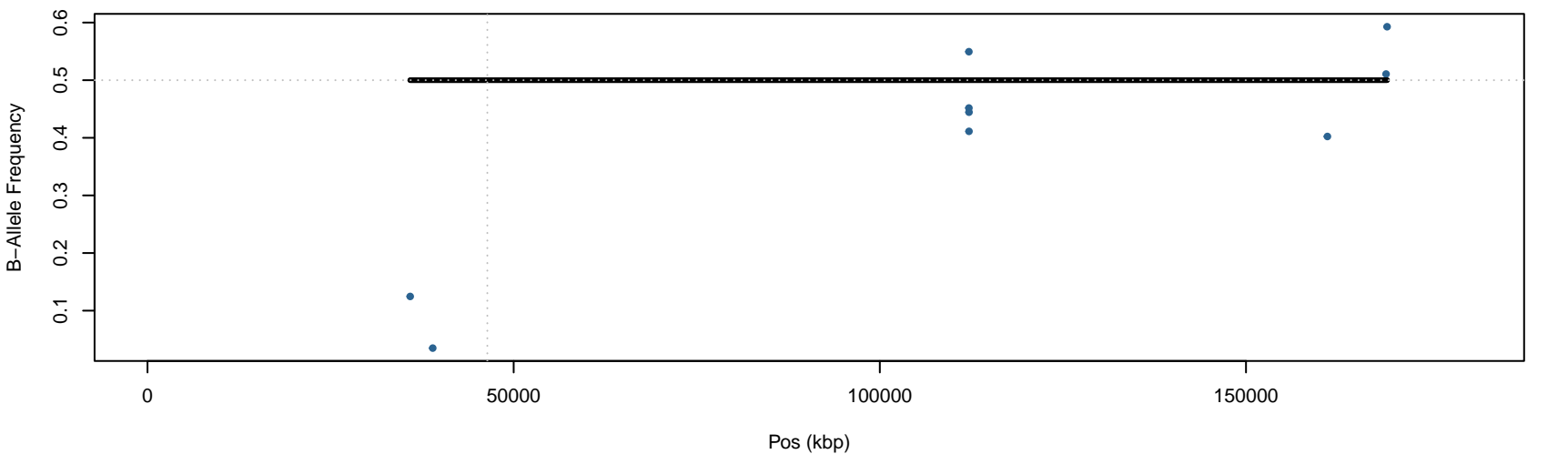
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 3



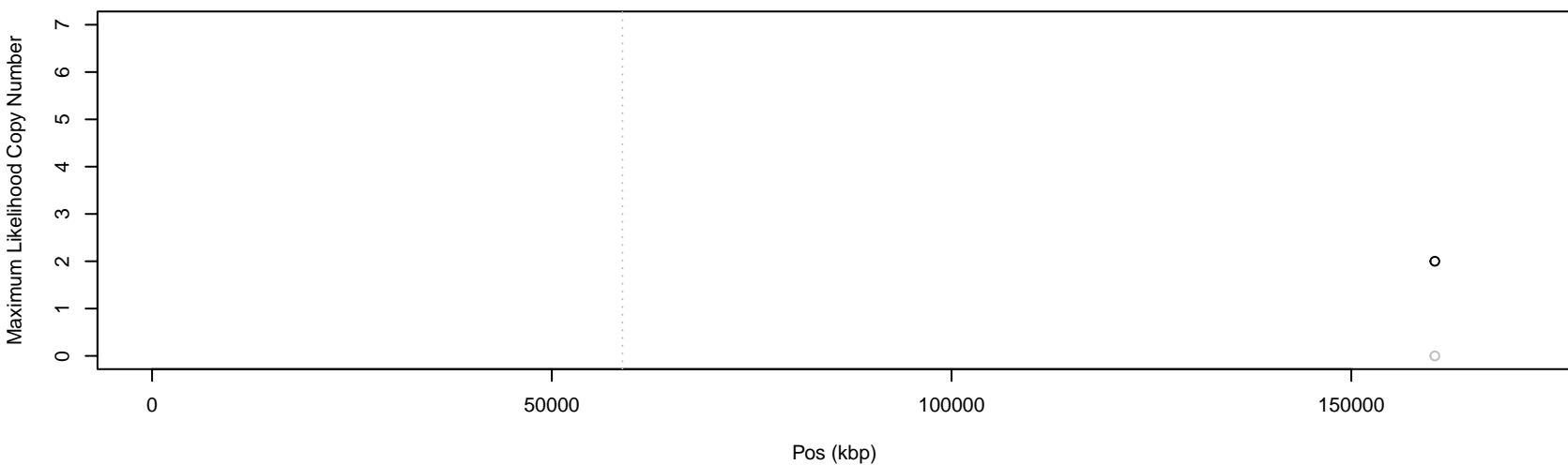
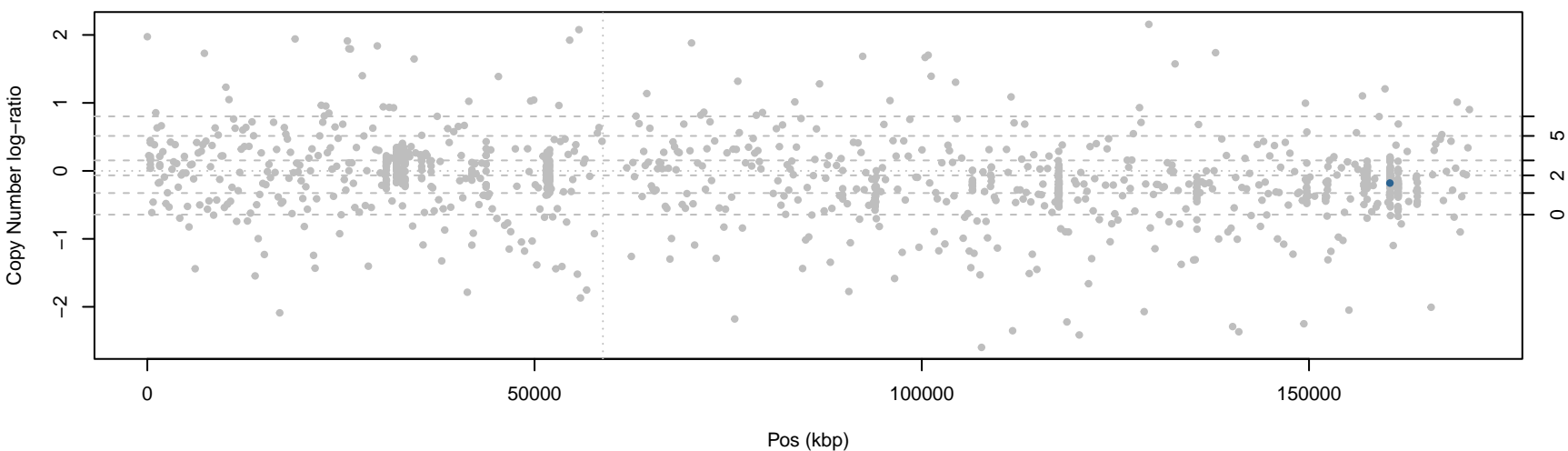
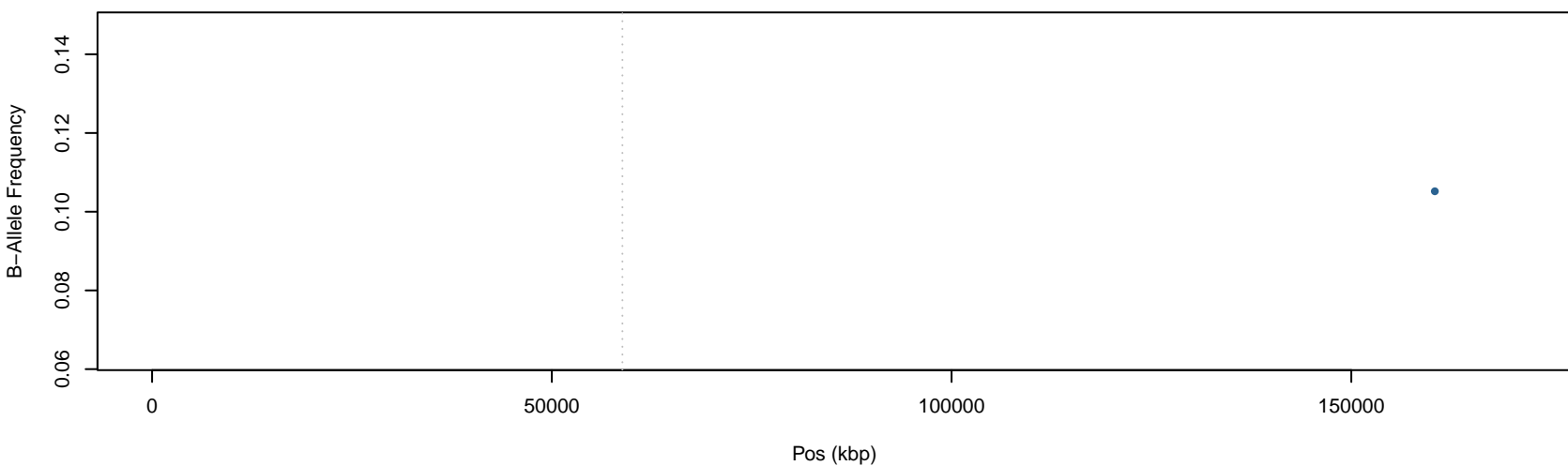
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 4



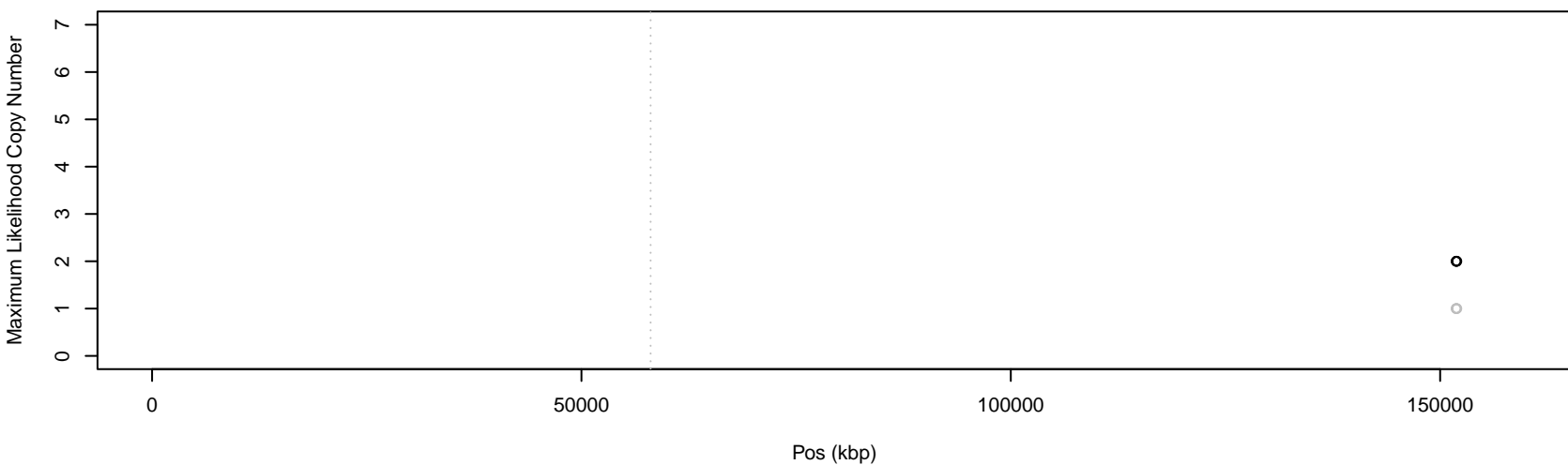
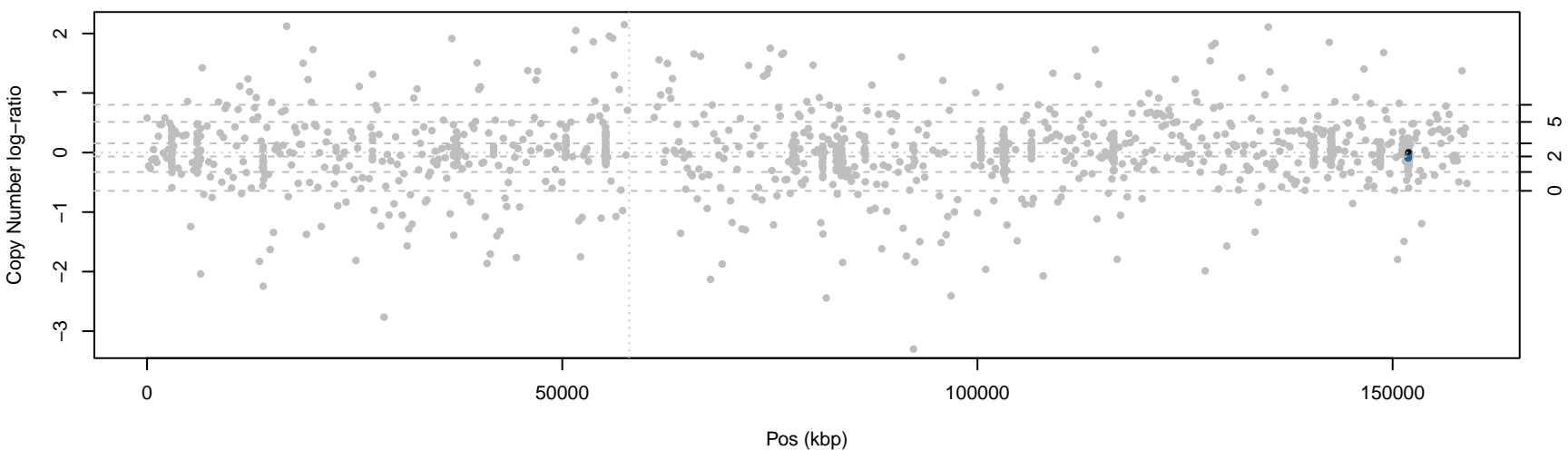
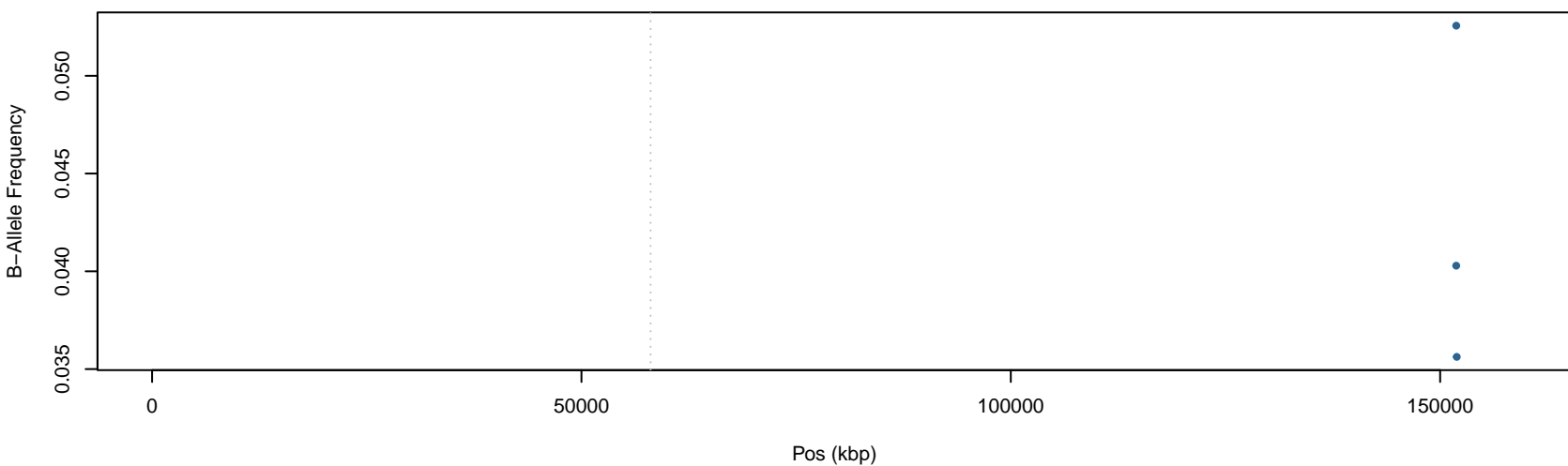
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 5



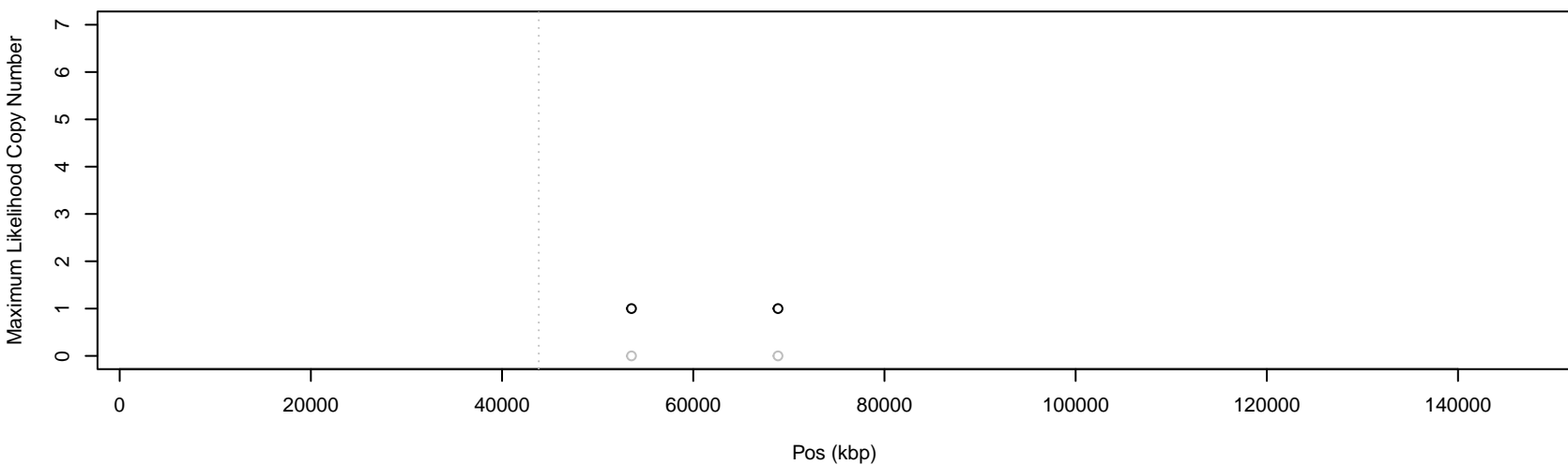
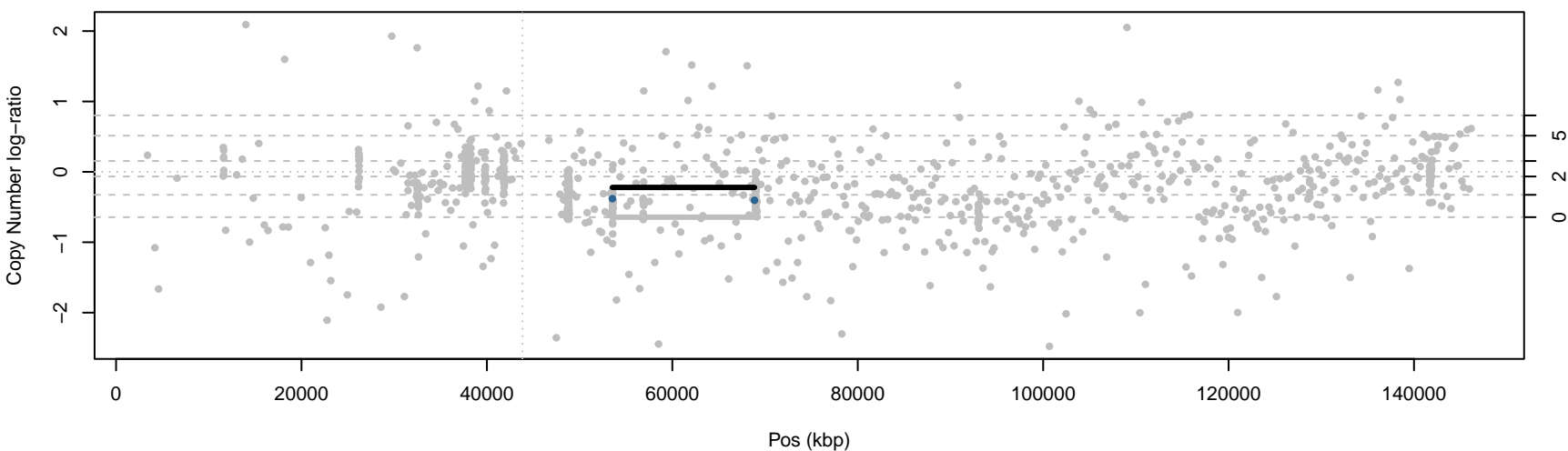
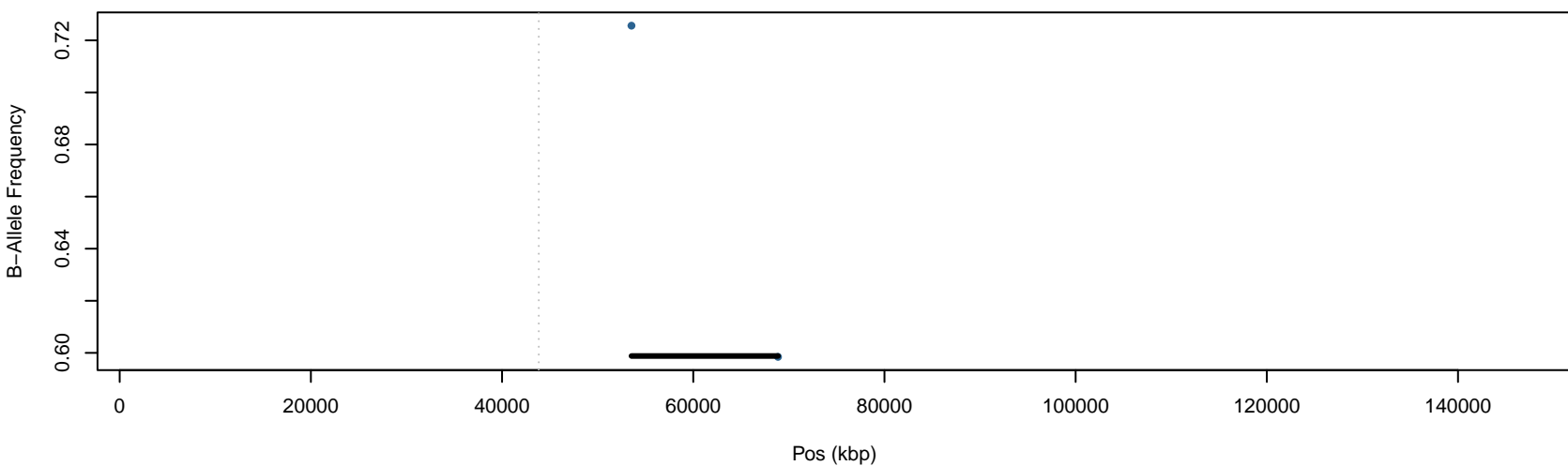
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 6



Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 7

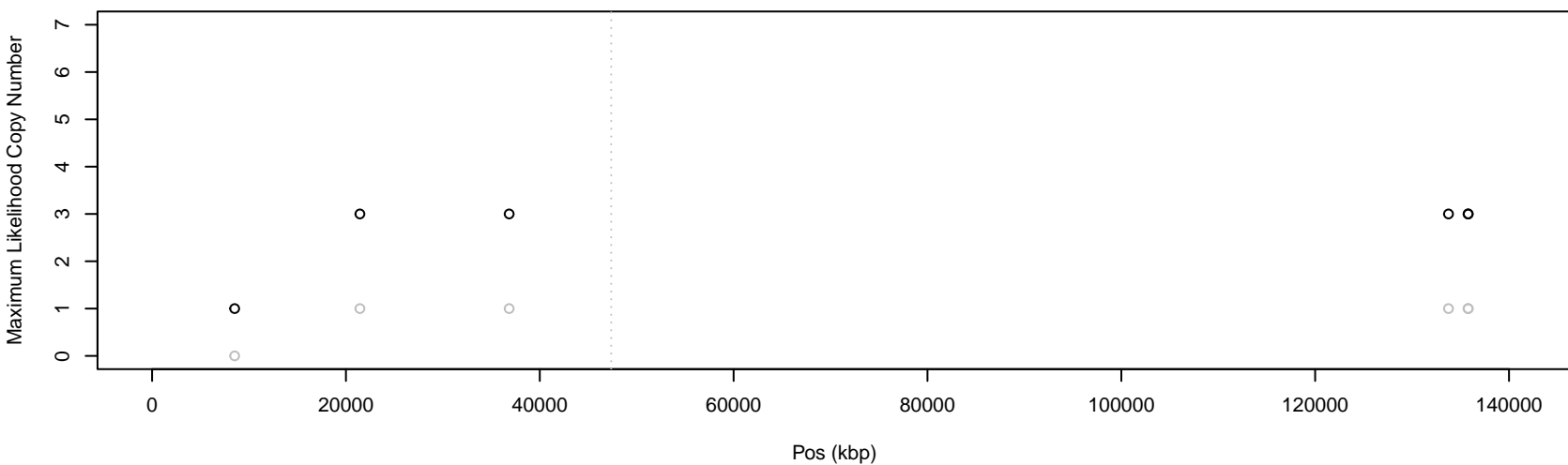
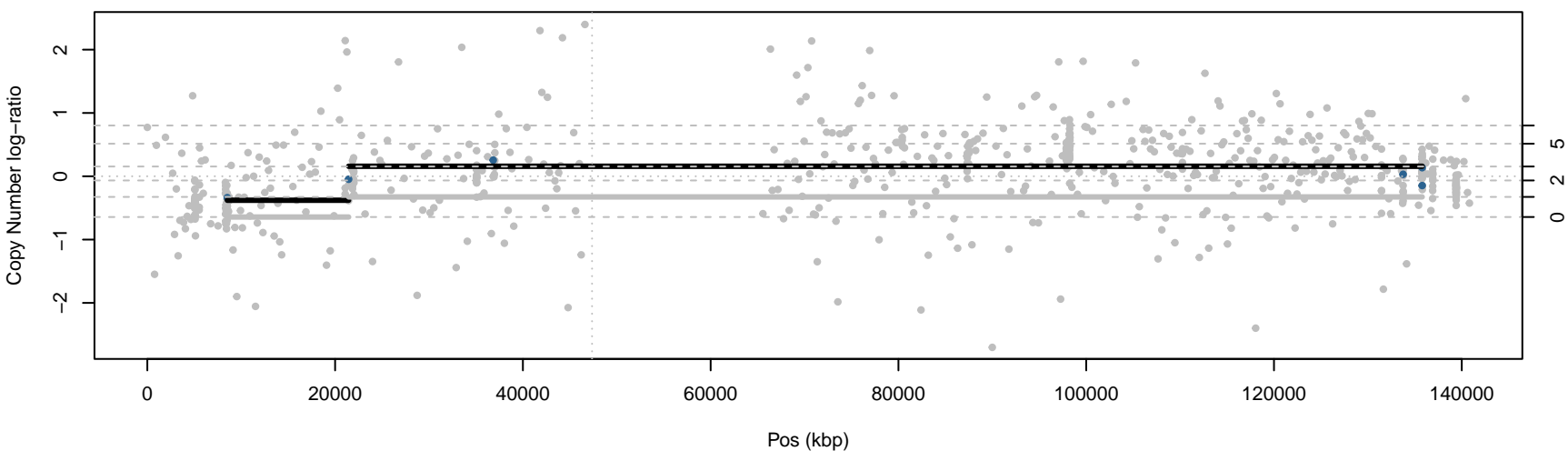
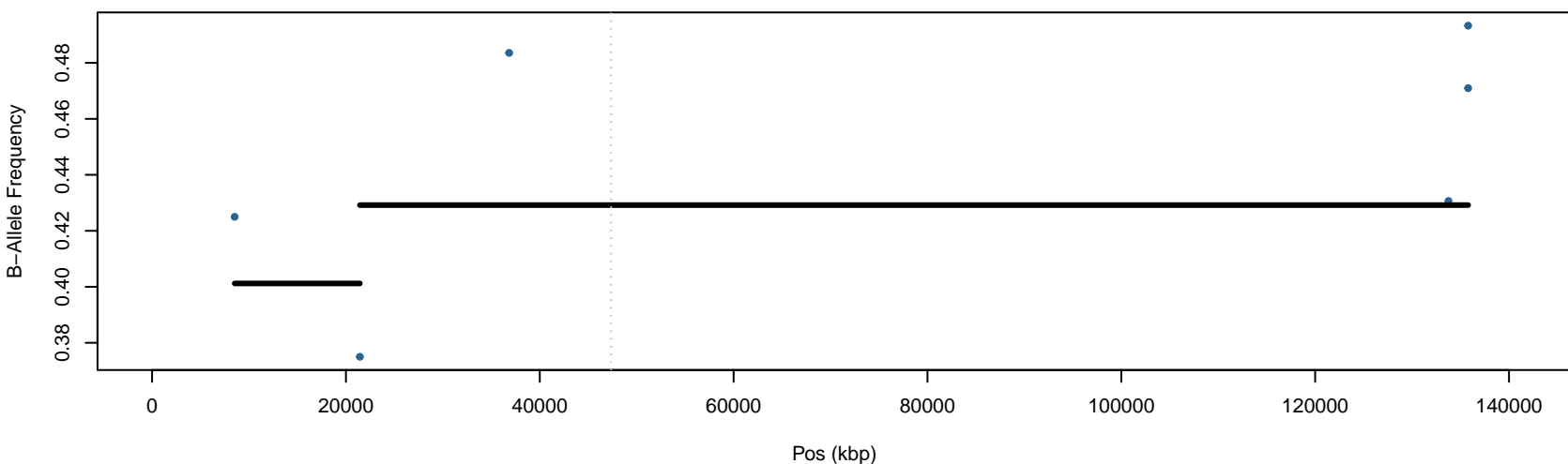


Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 8

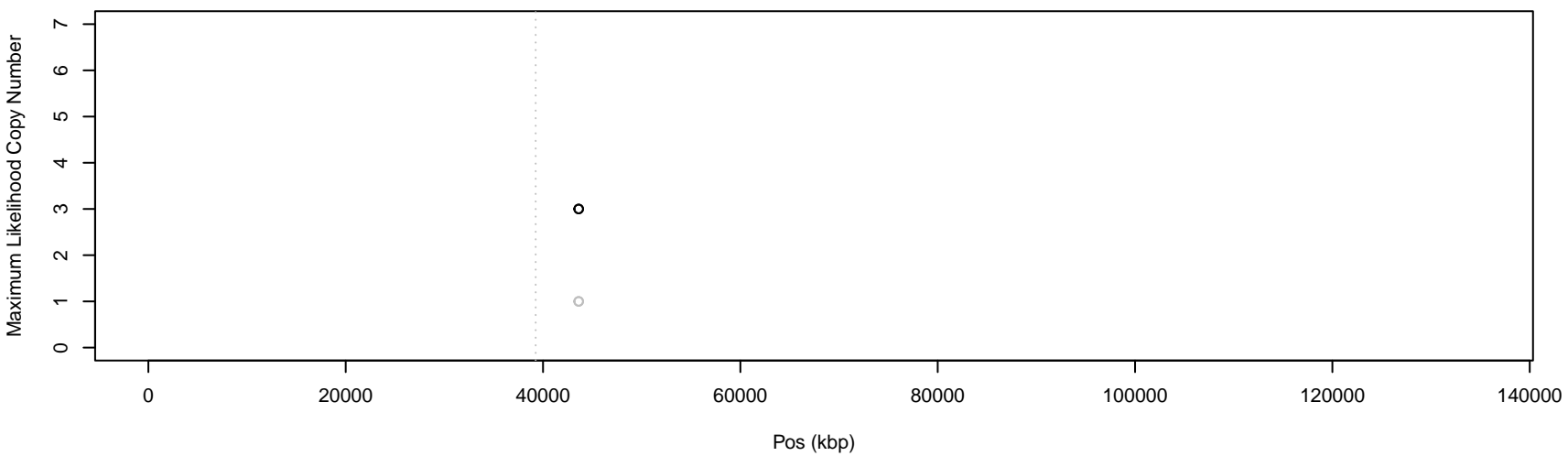
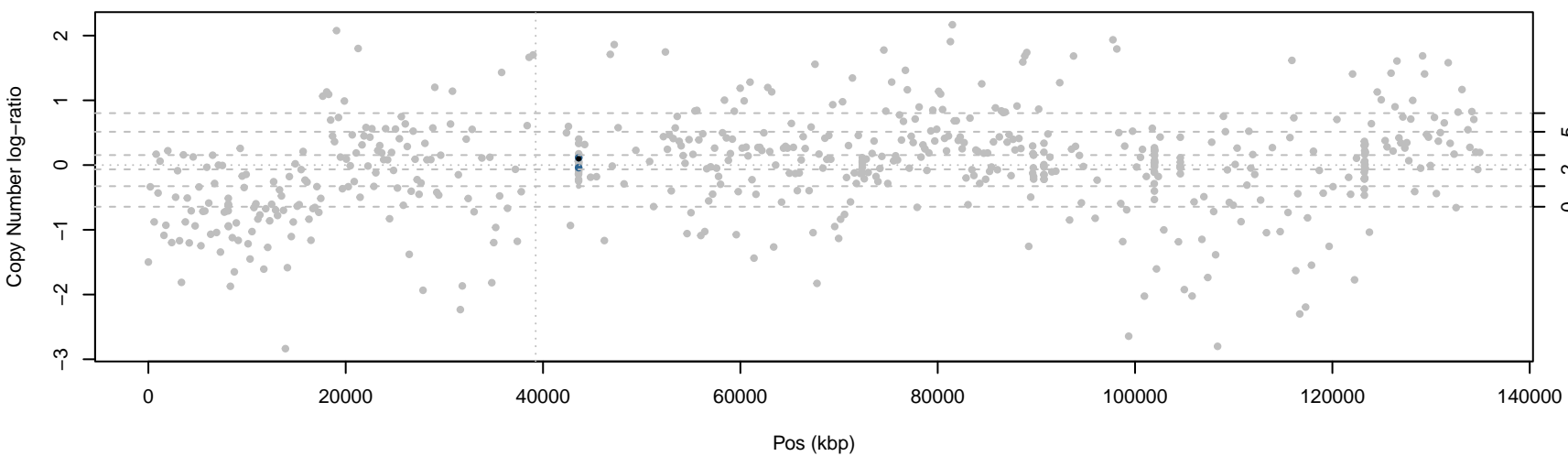
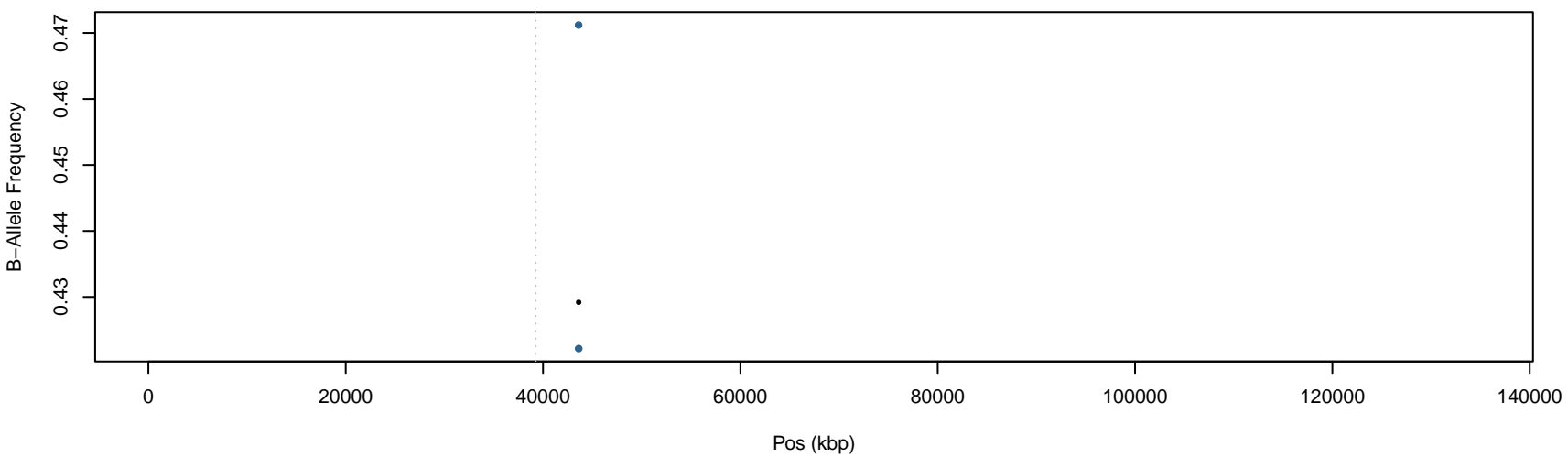




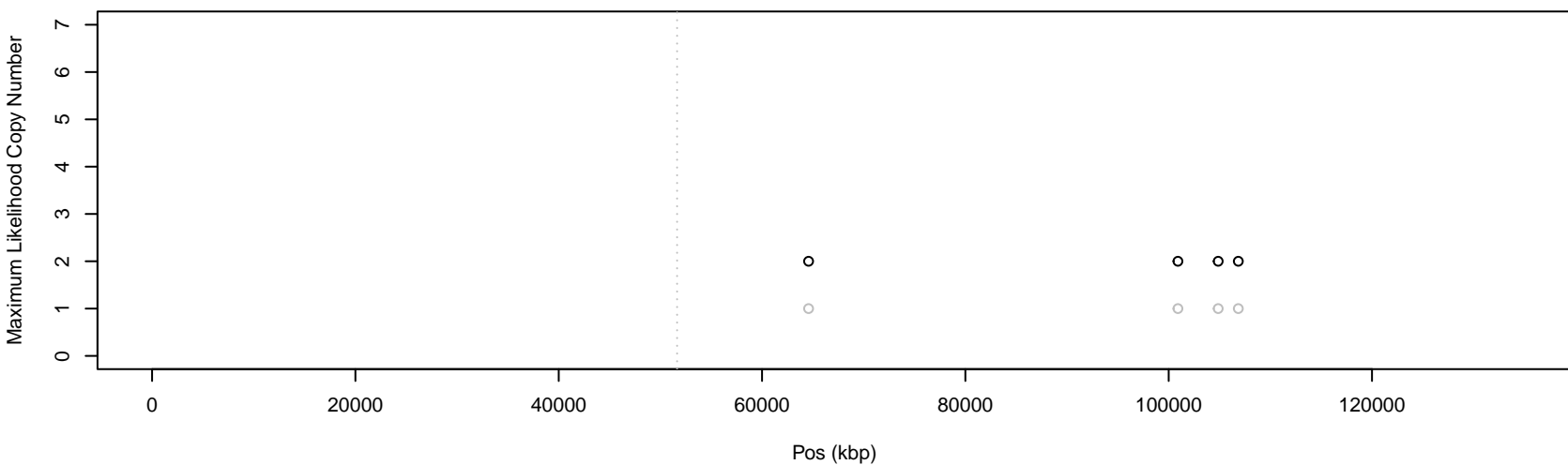
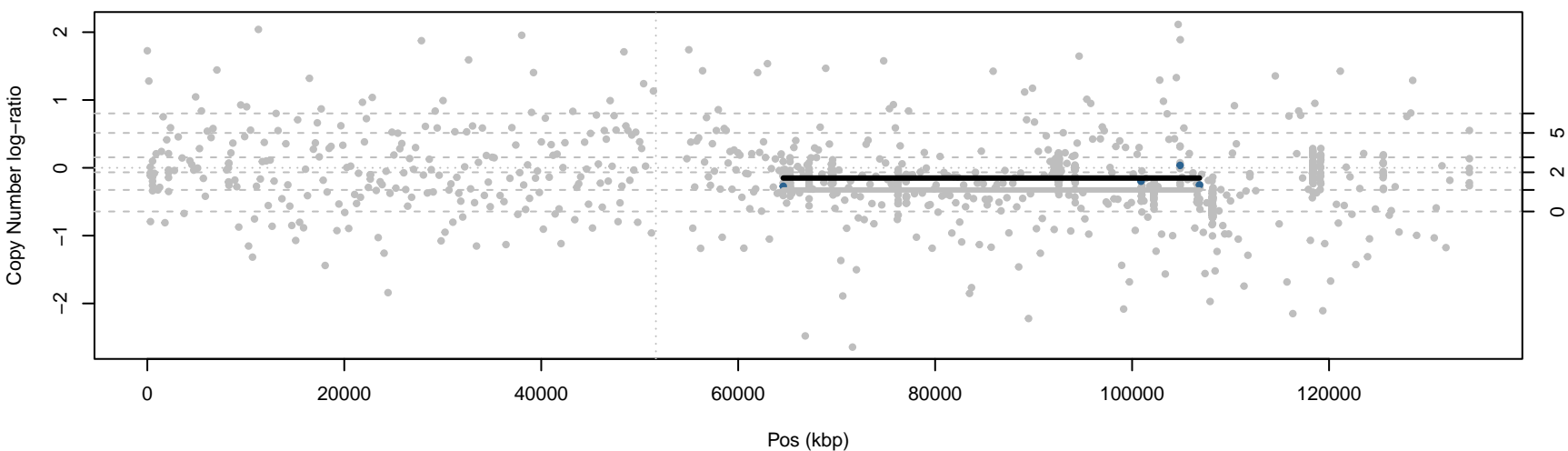
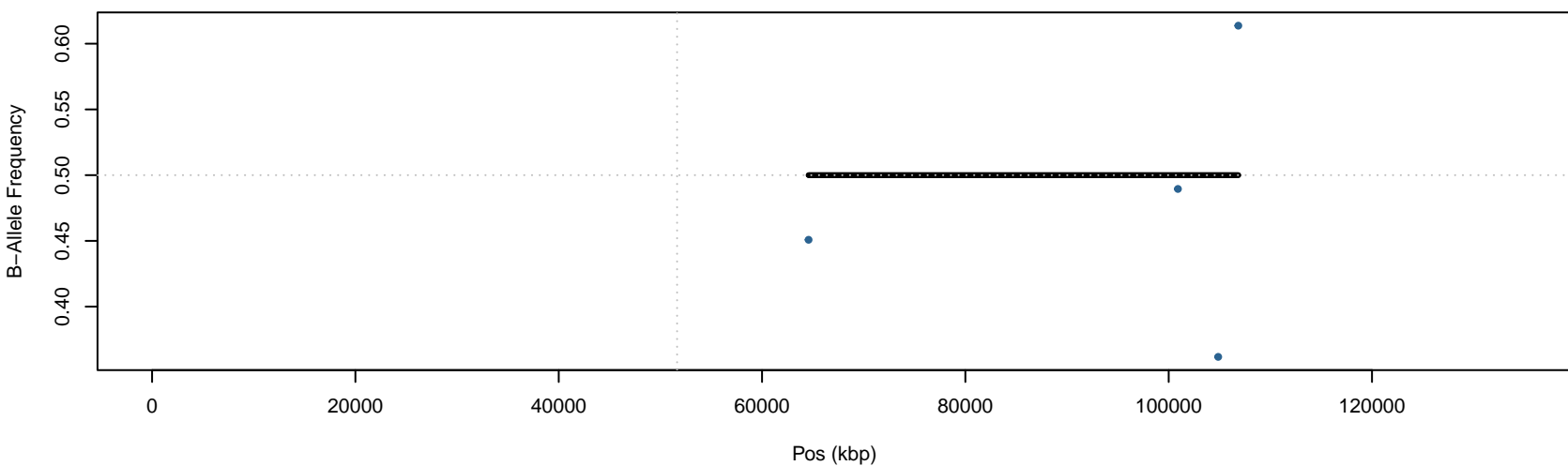
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 9



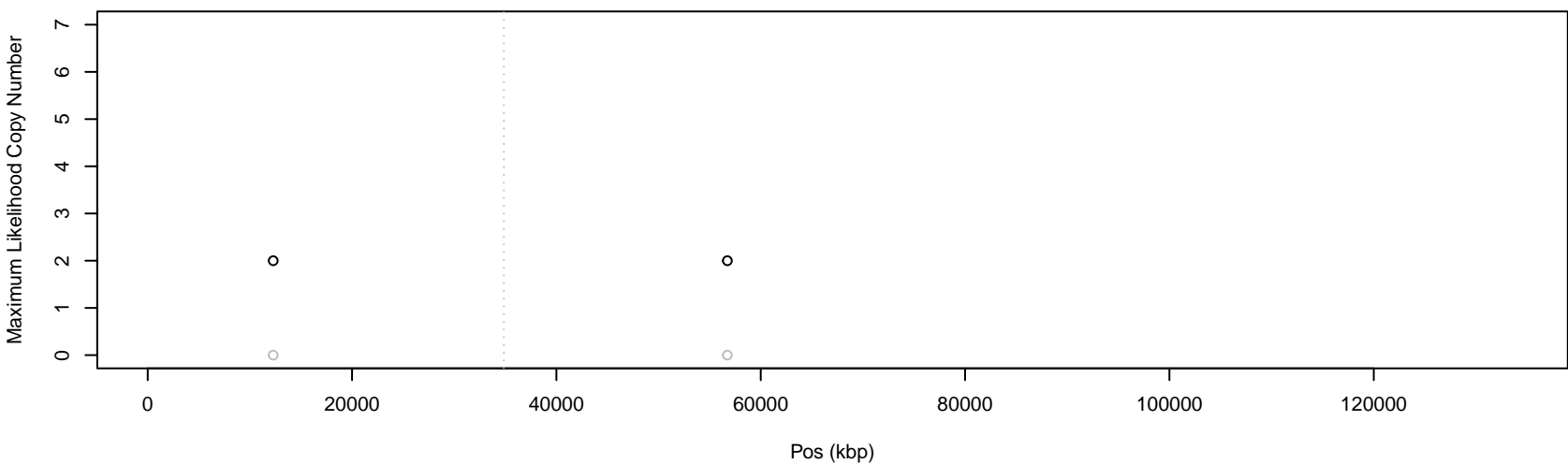
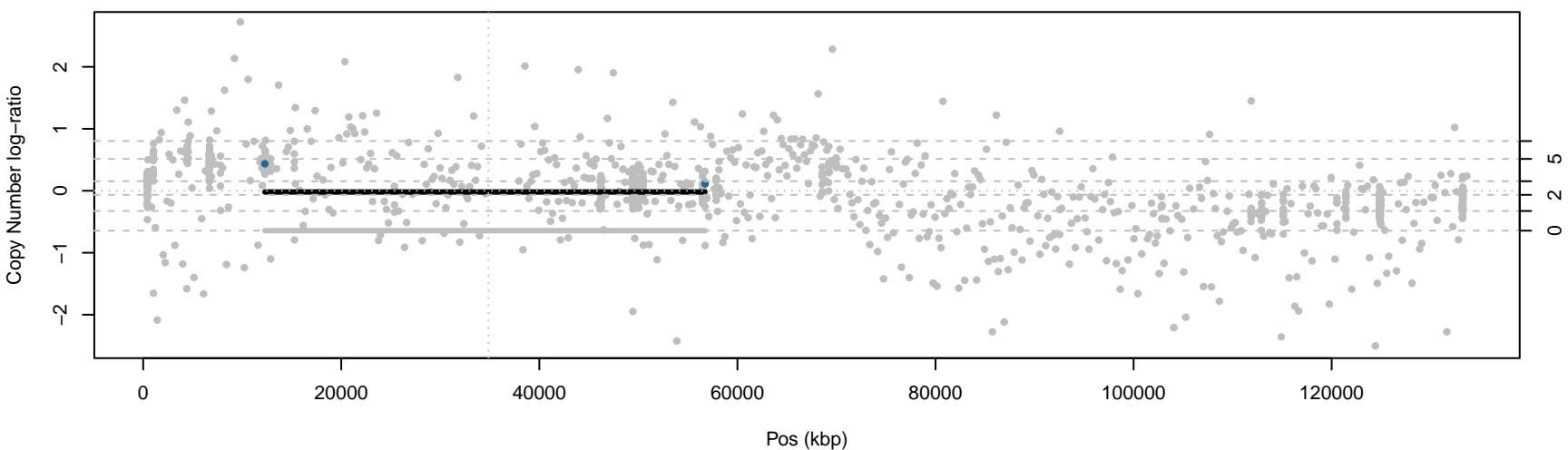
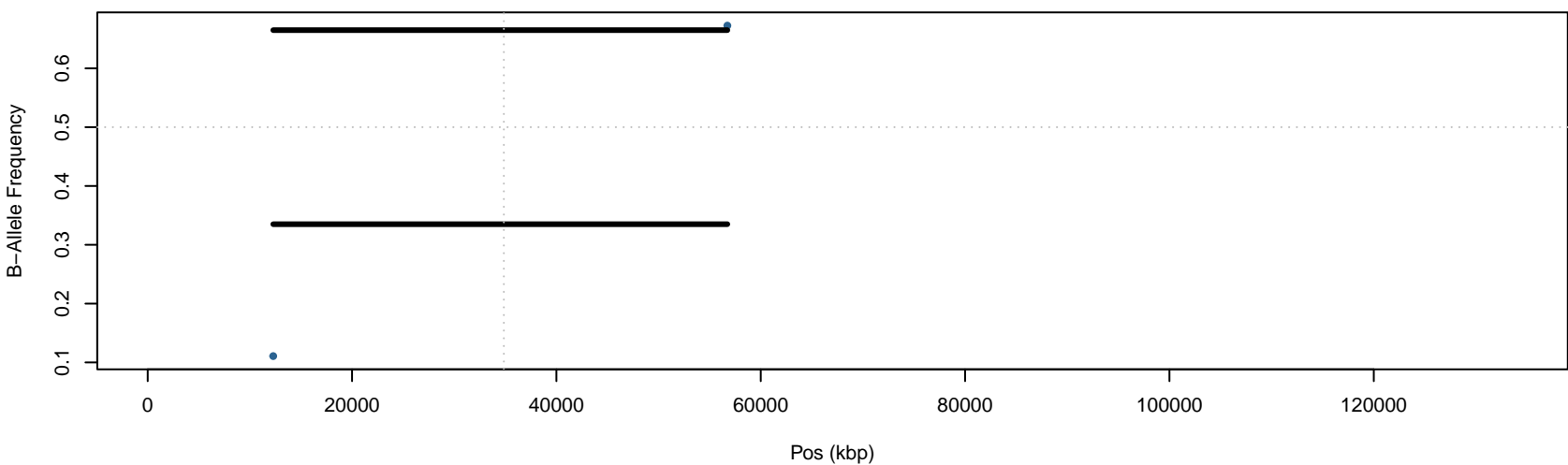
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 10



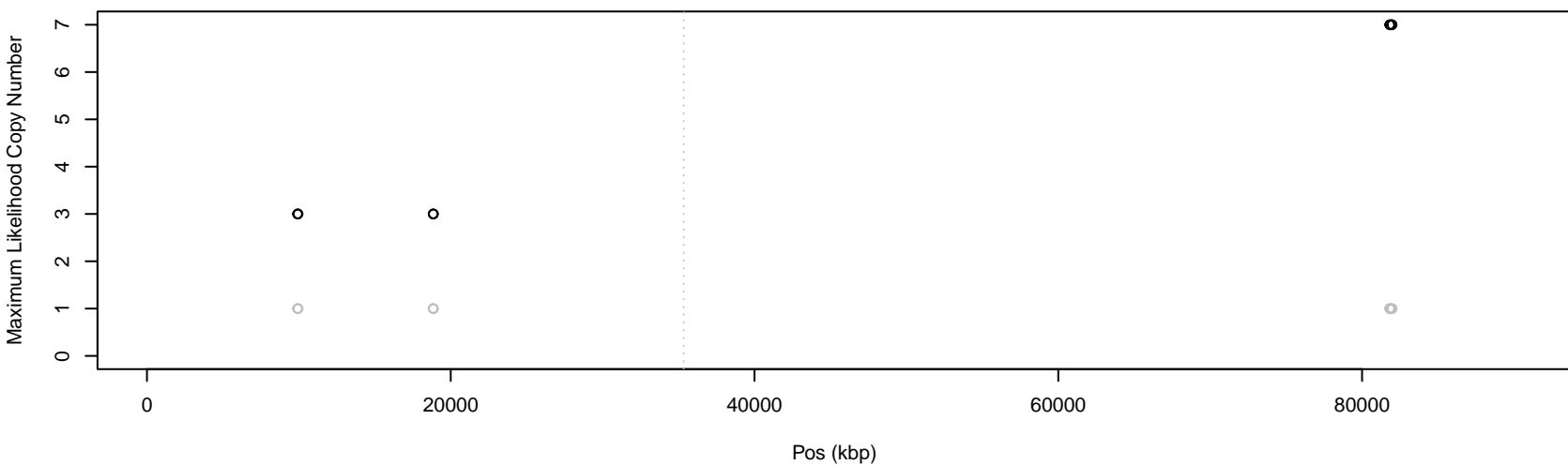
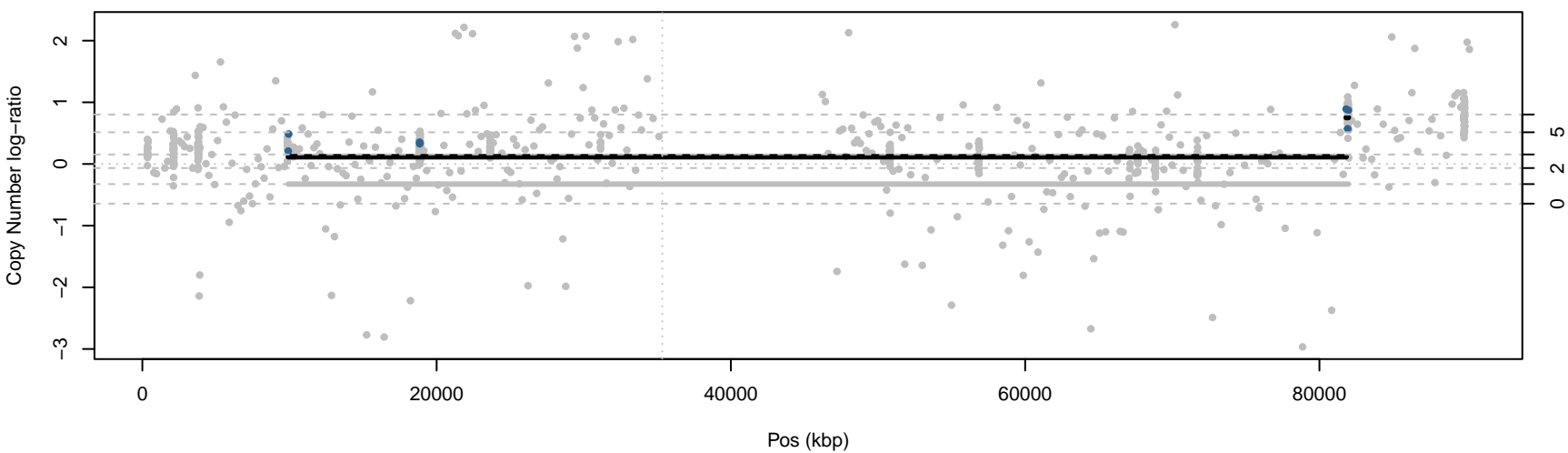
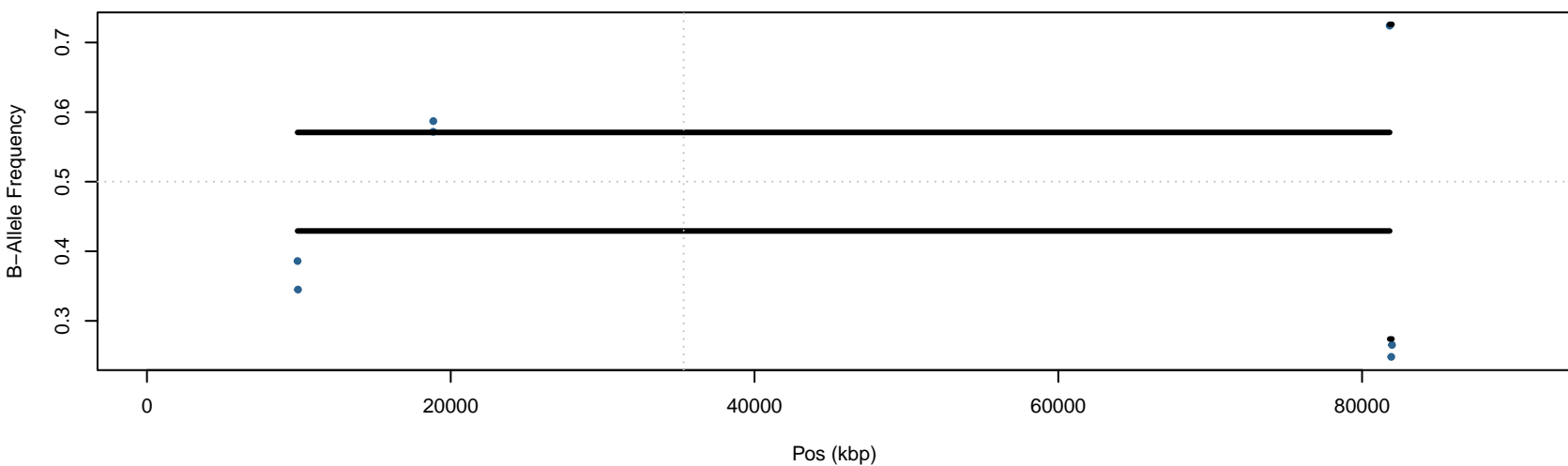
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 11



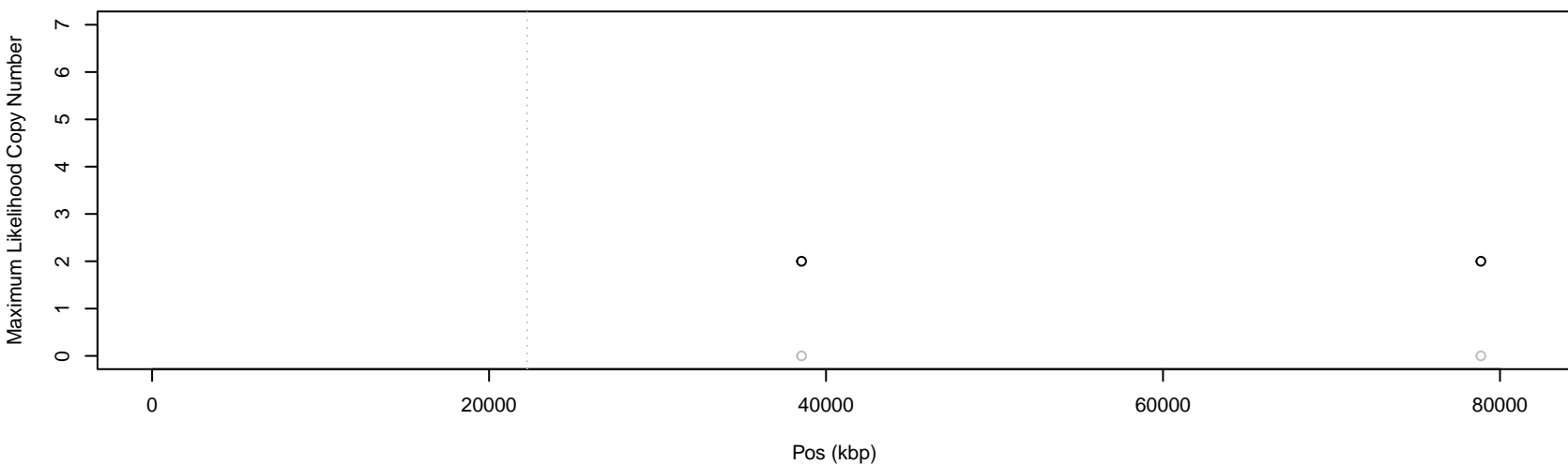
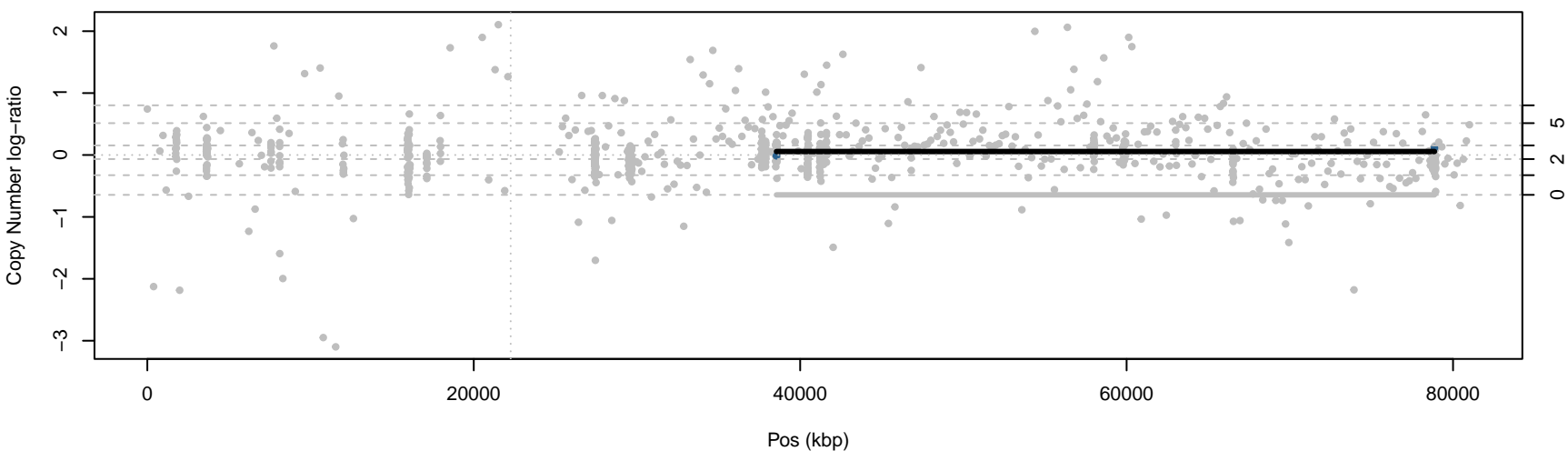
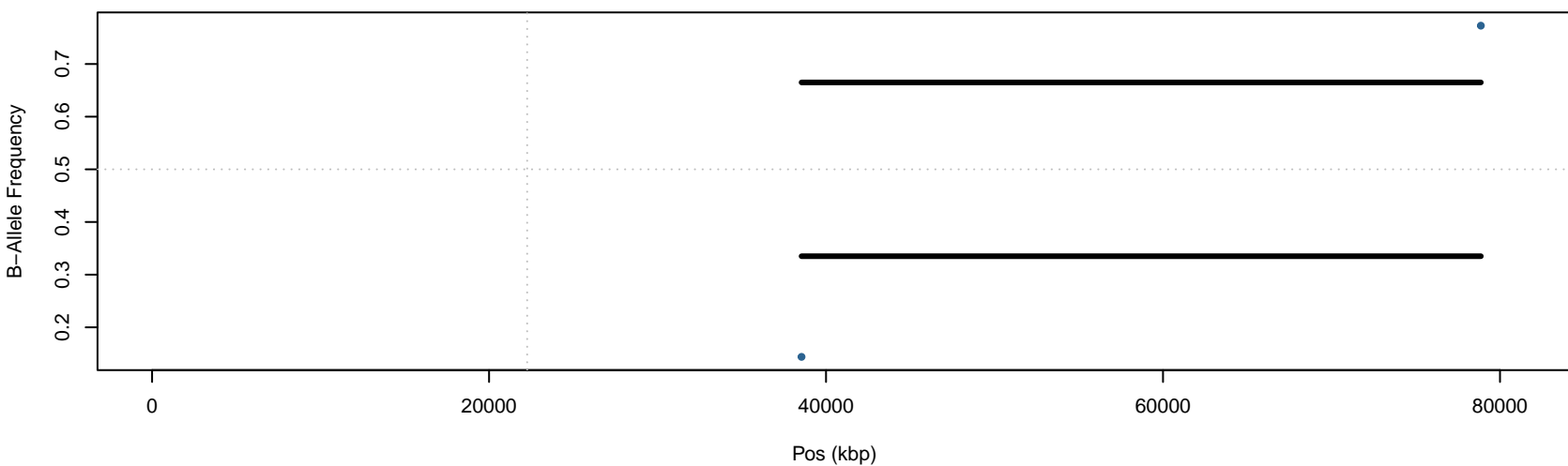
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 12



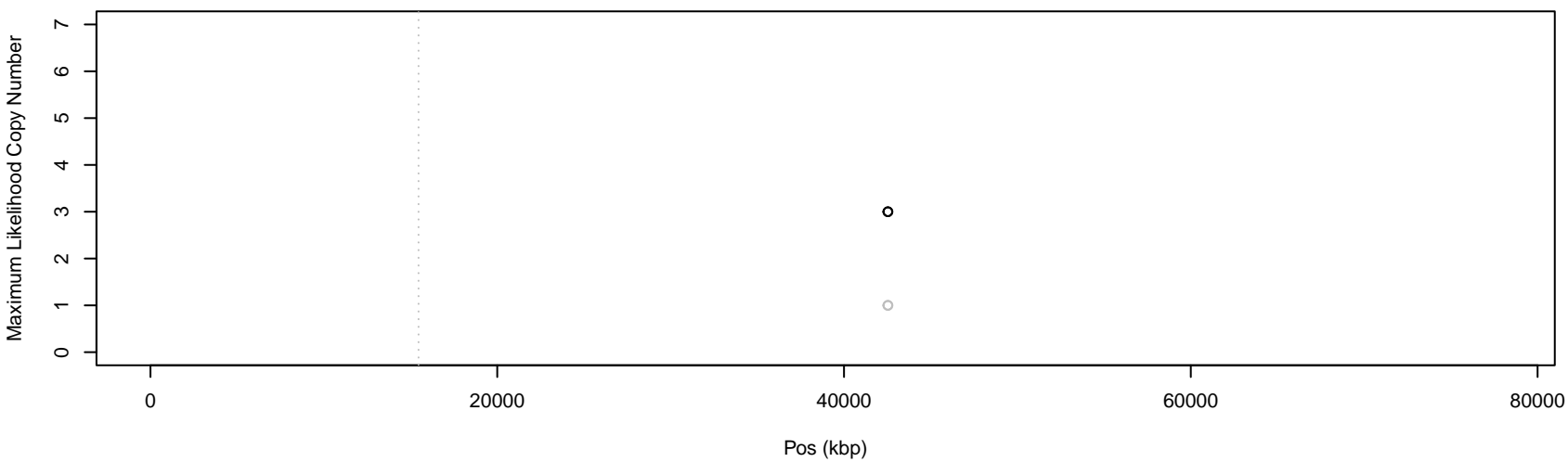
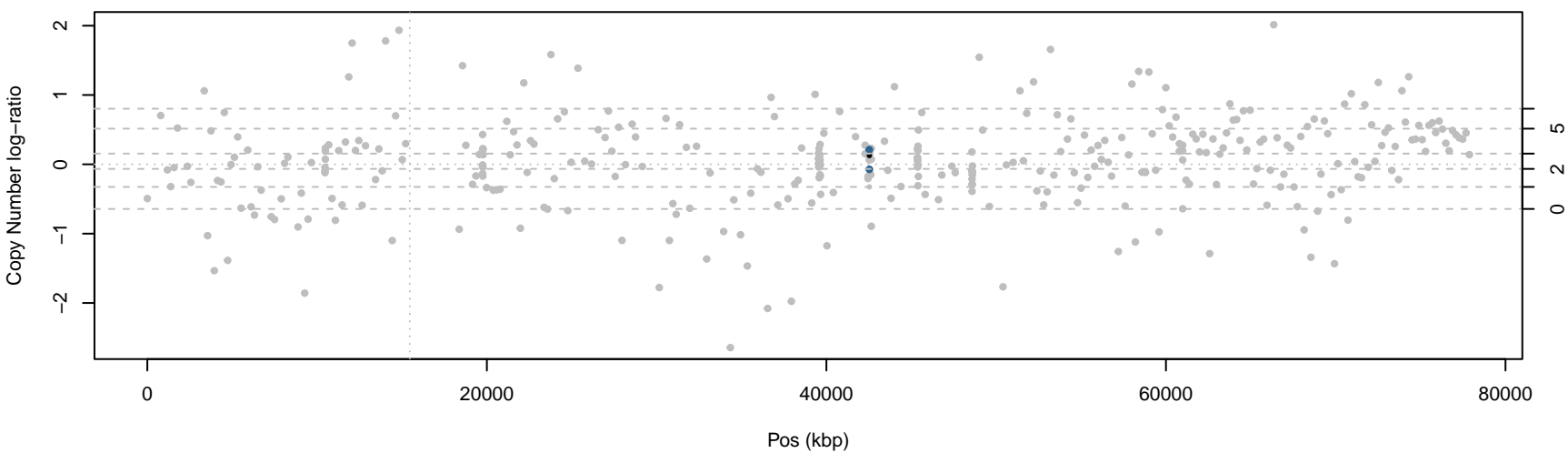
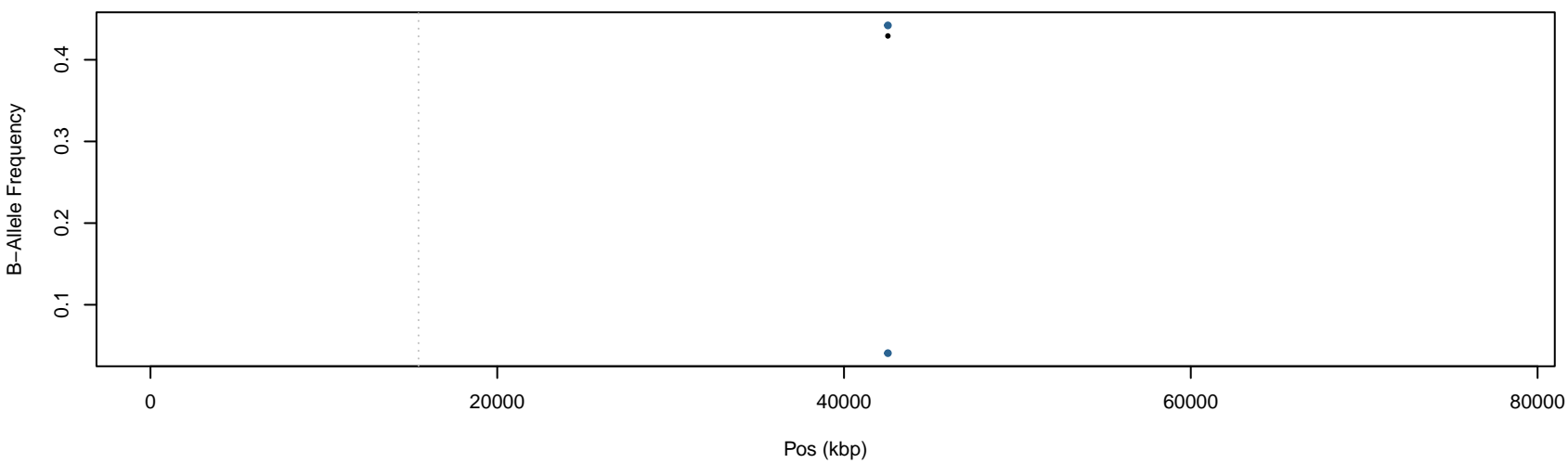
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 16



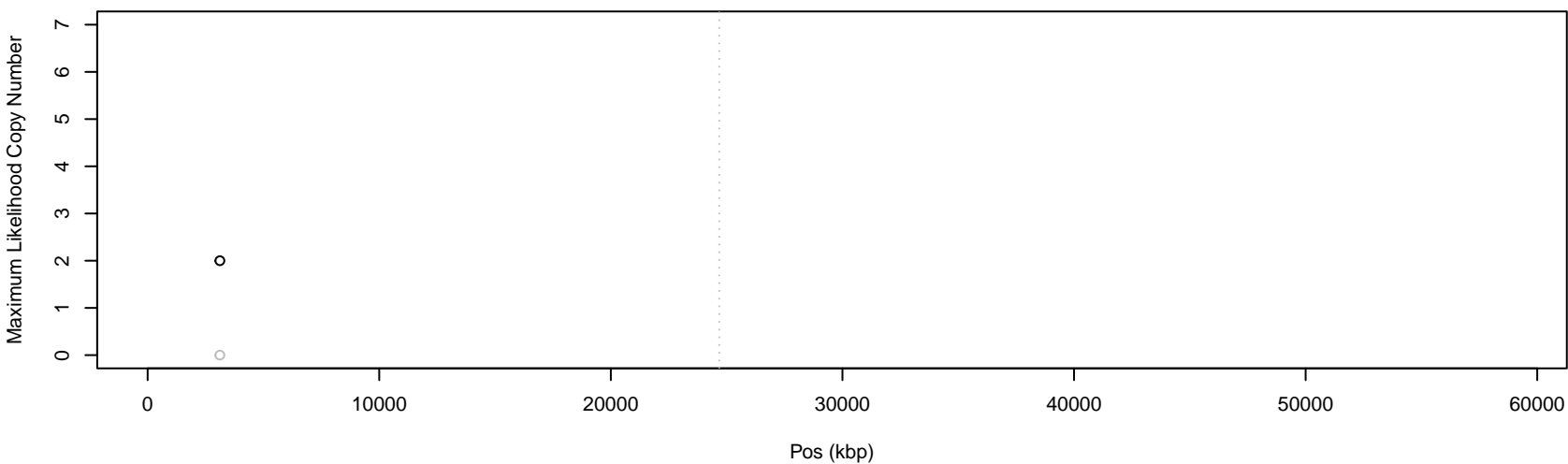
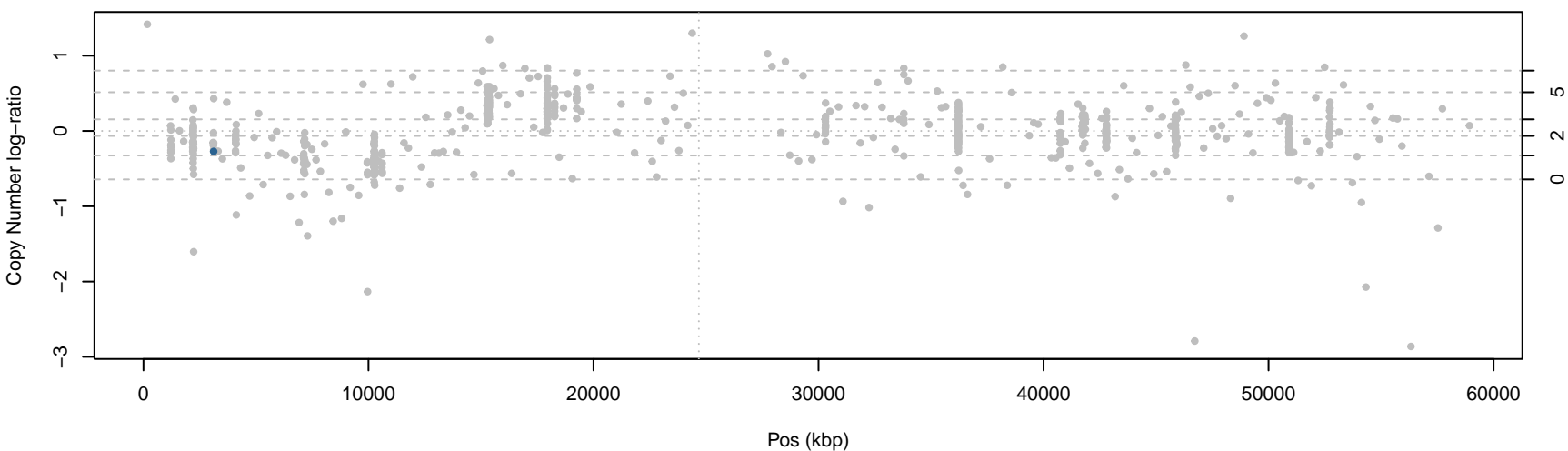
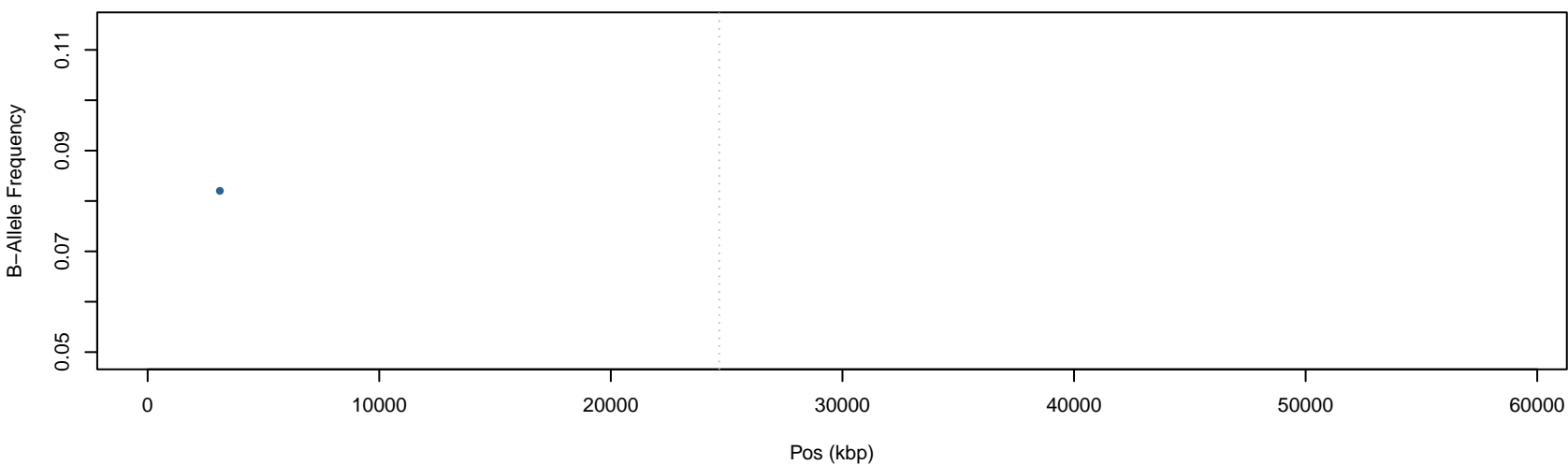
Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 17



Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 18

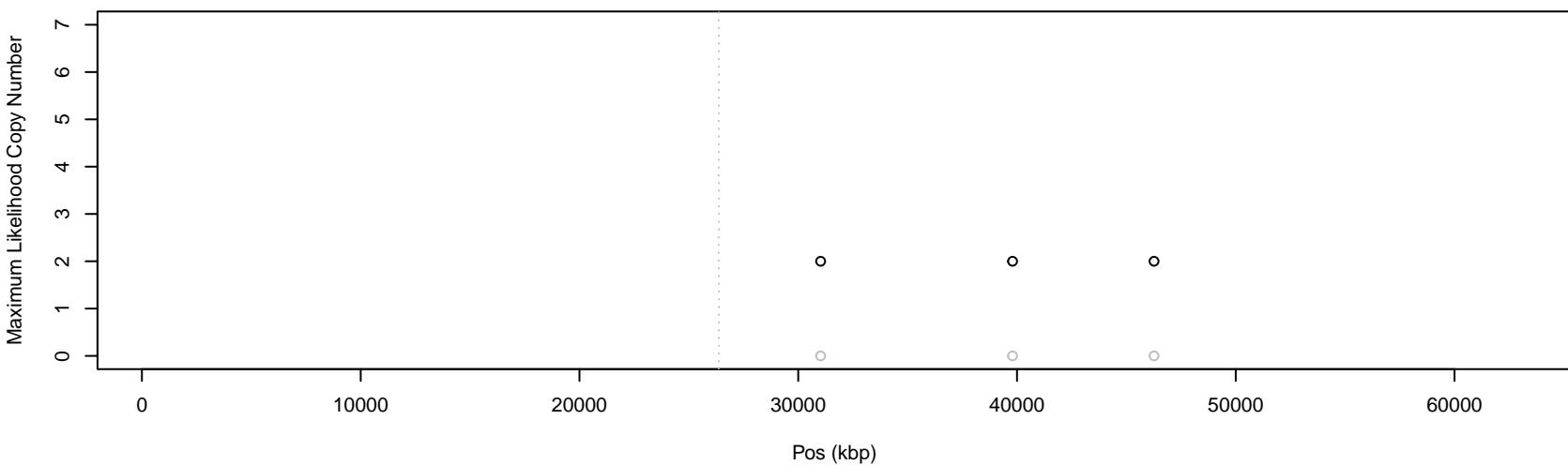
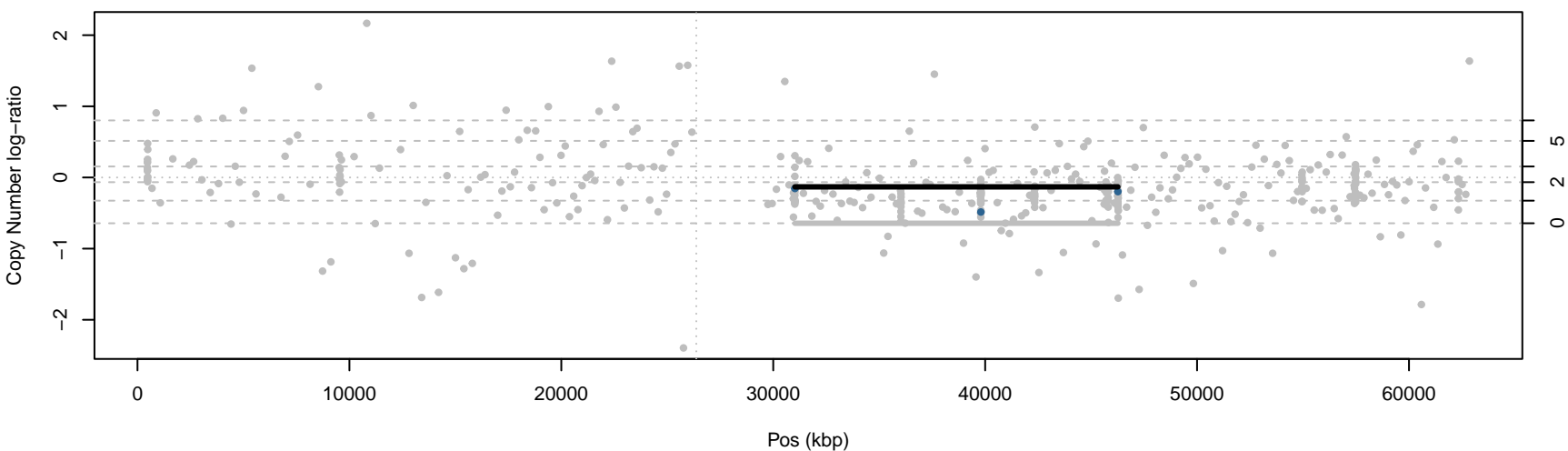
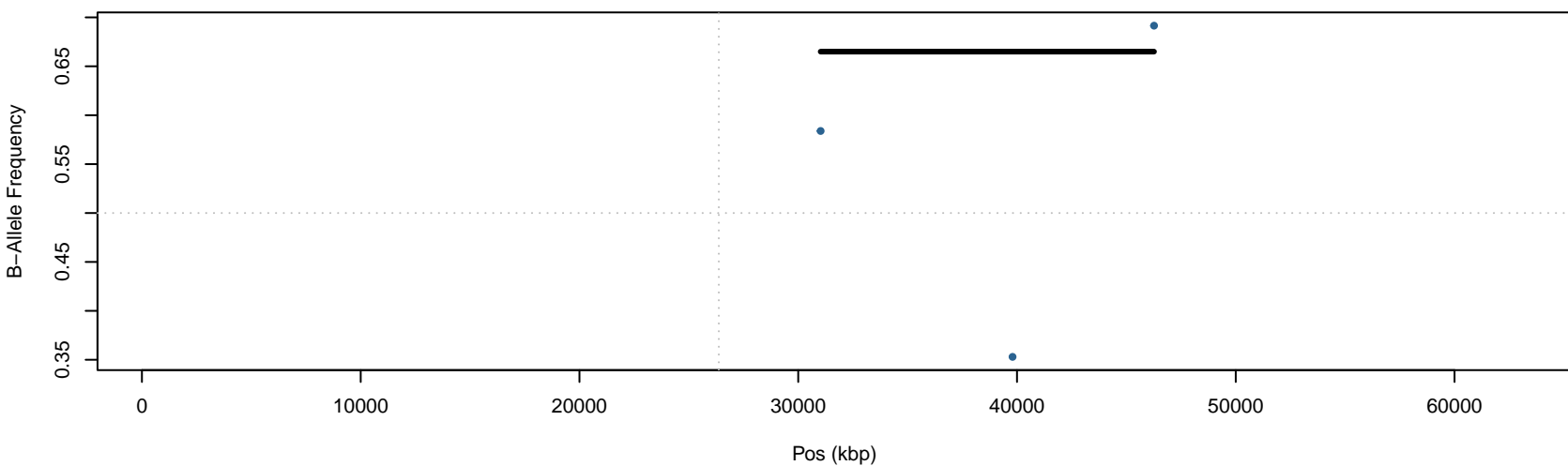


Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 19





Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 20



Purity: 0.33 Tumor ploidy: 2.283 SNV log-likelihood: -92.82 GoF: 88.7% Mean coverage: 385;708 Chromosome: 22

