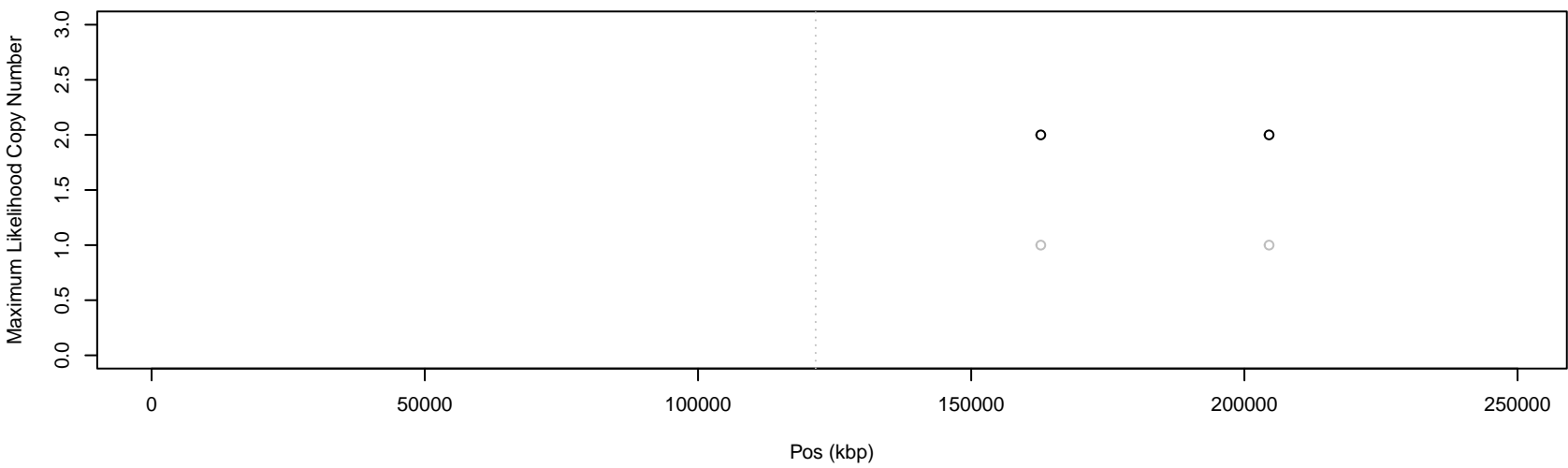
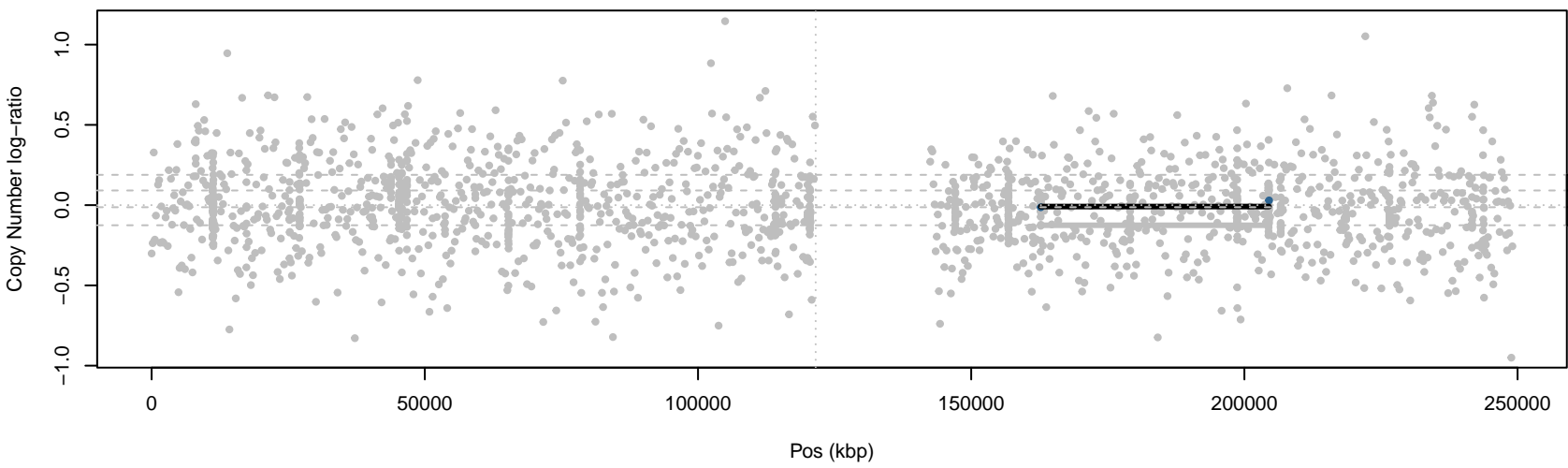
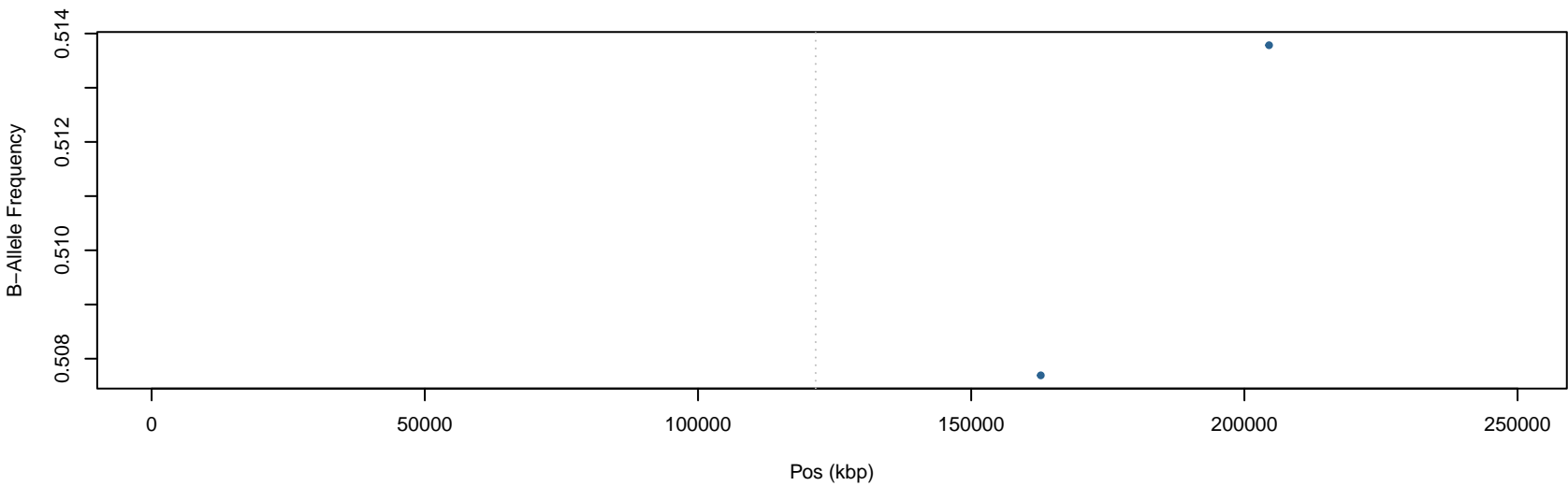
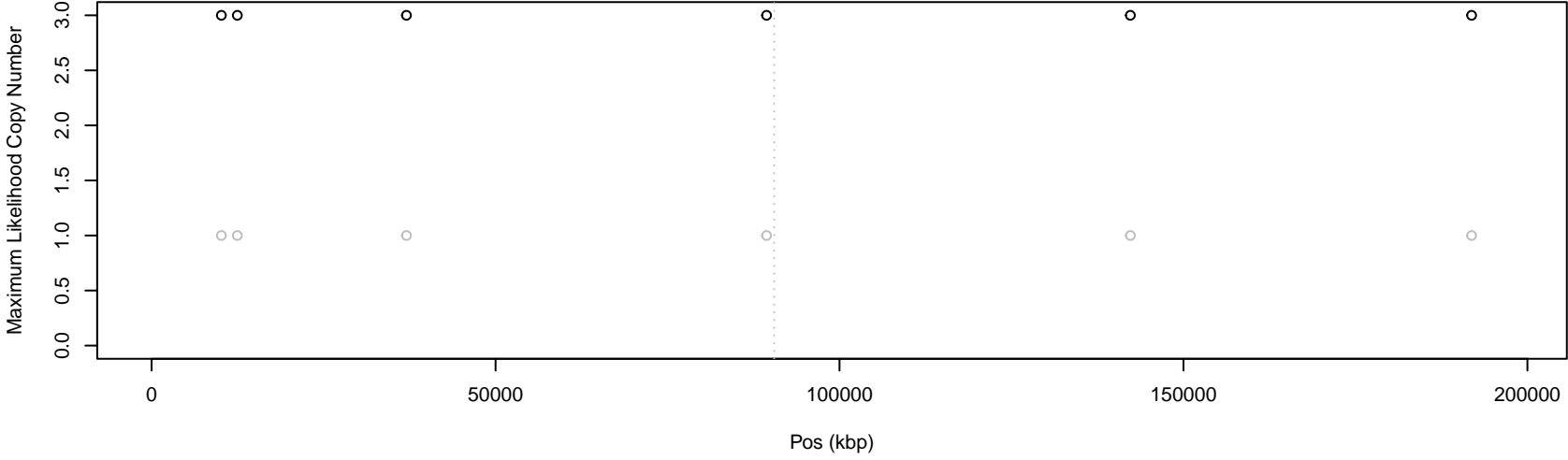
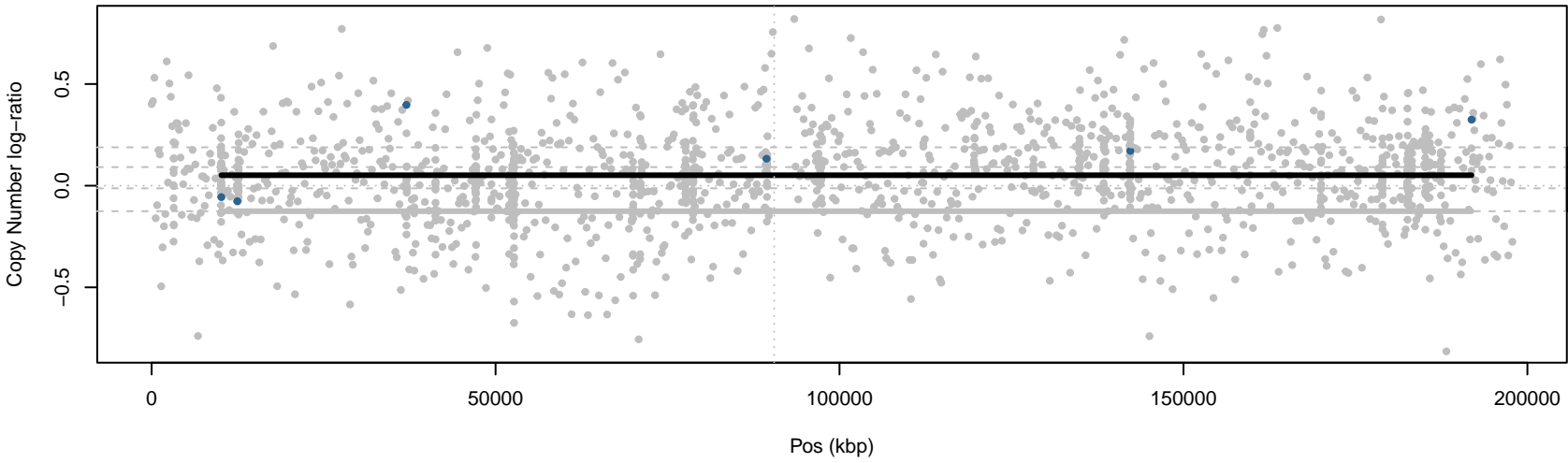
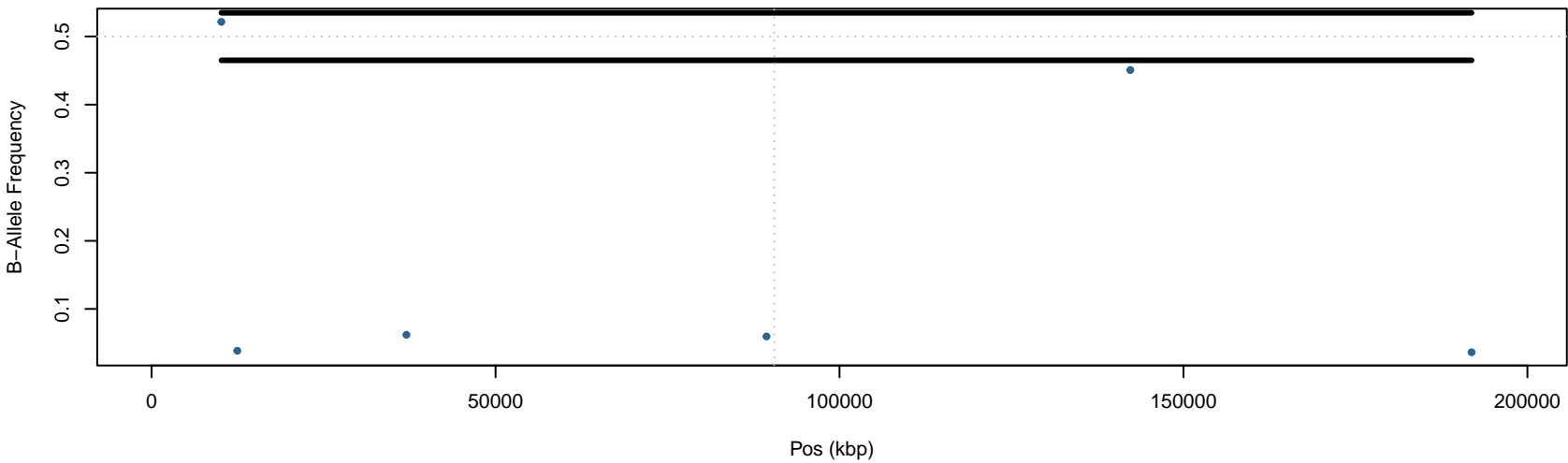


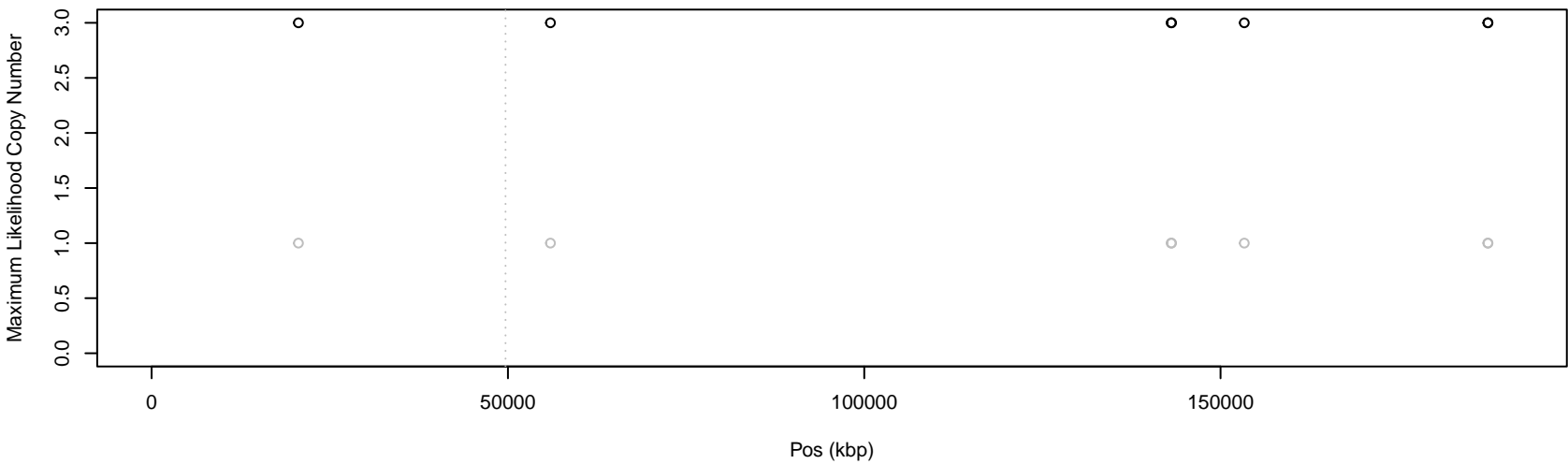
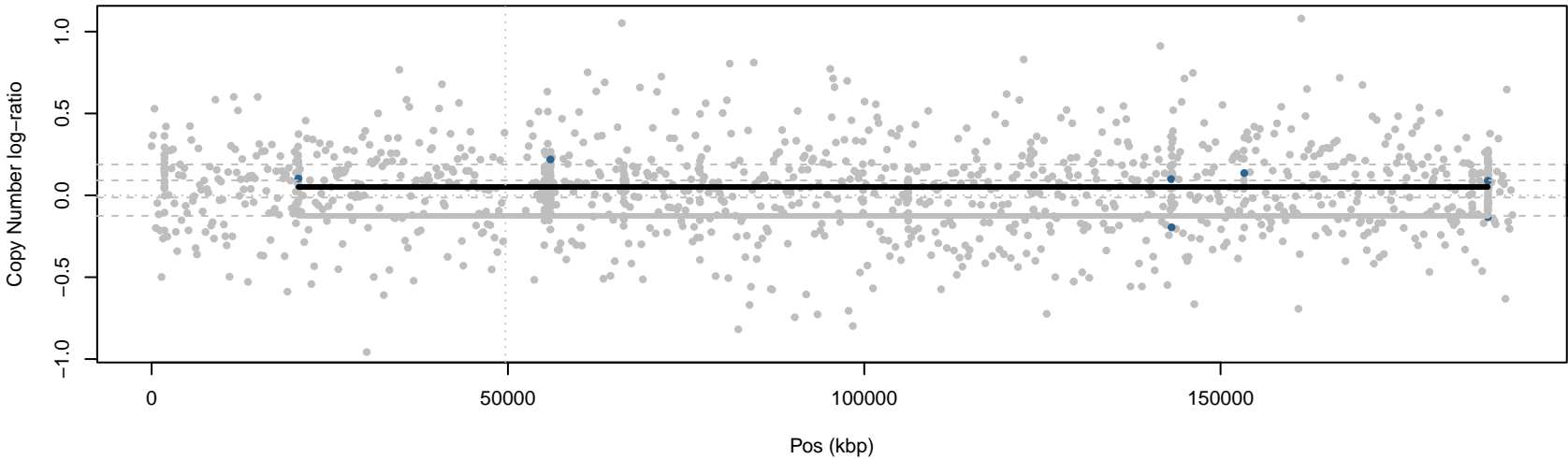
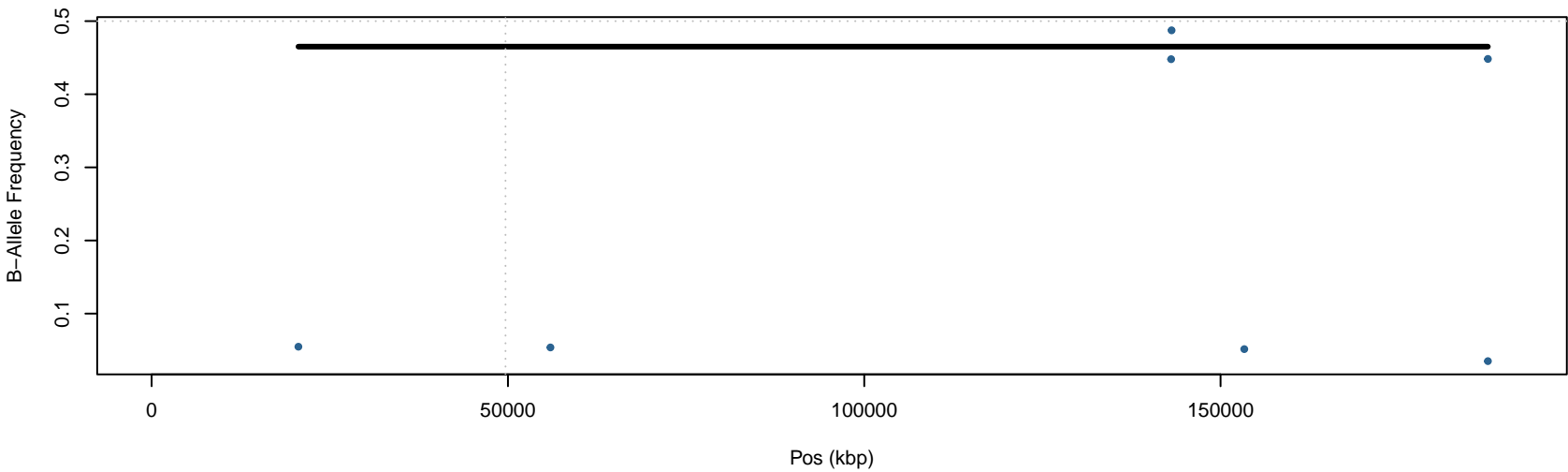
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 1



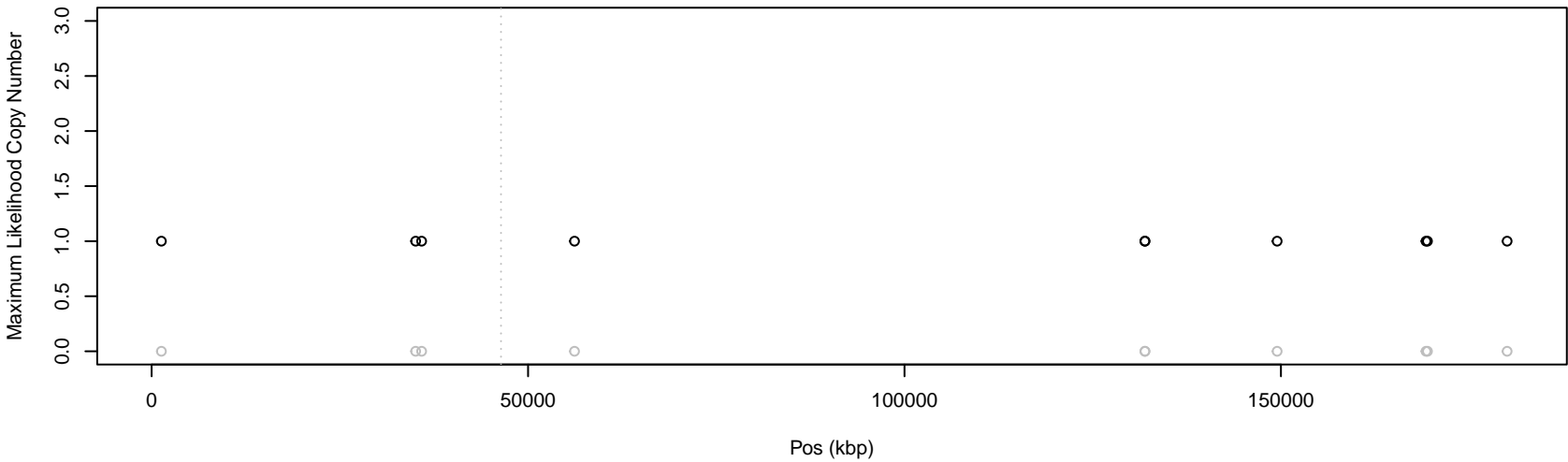
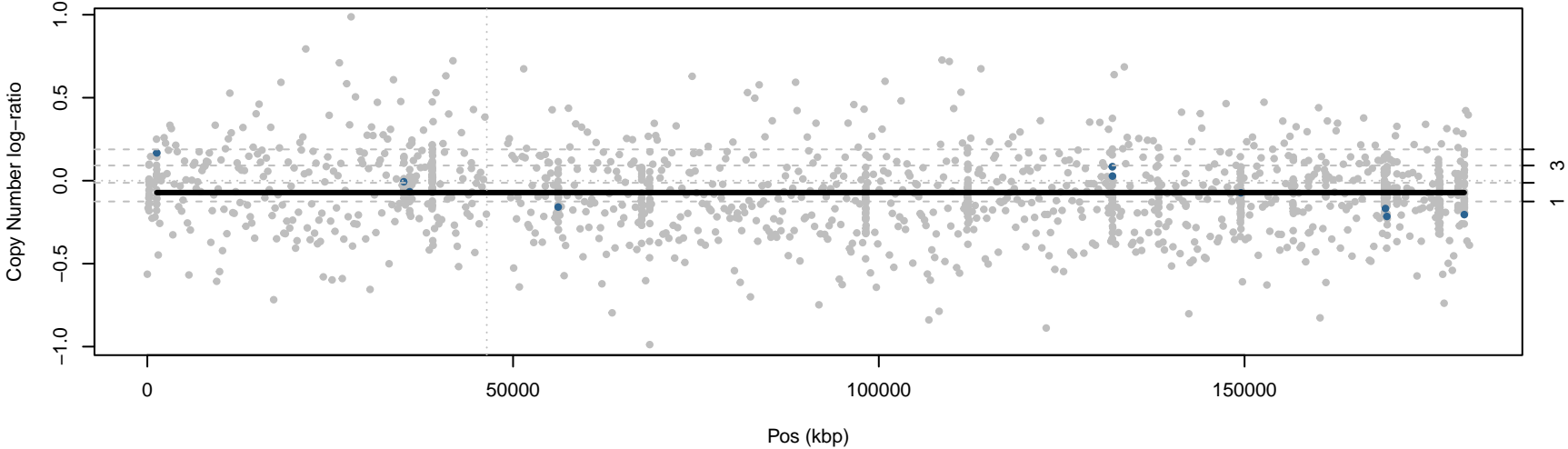
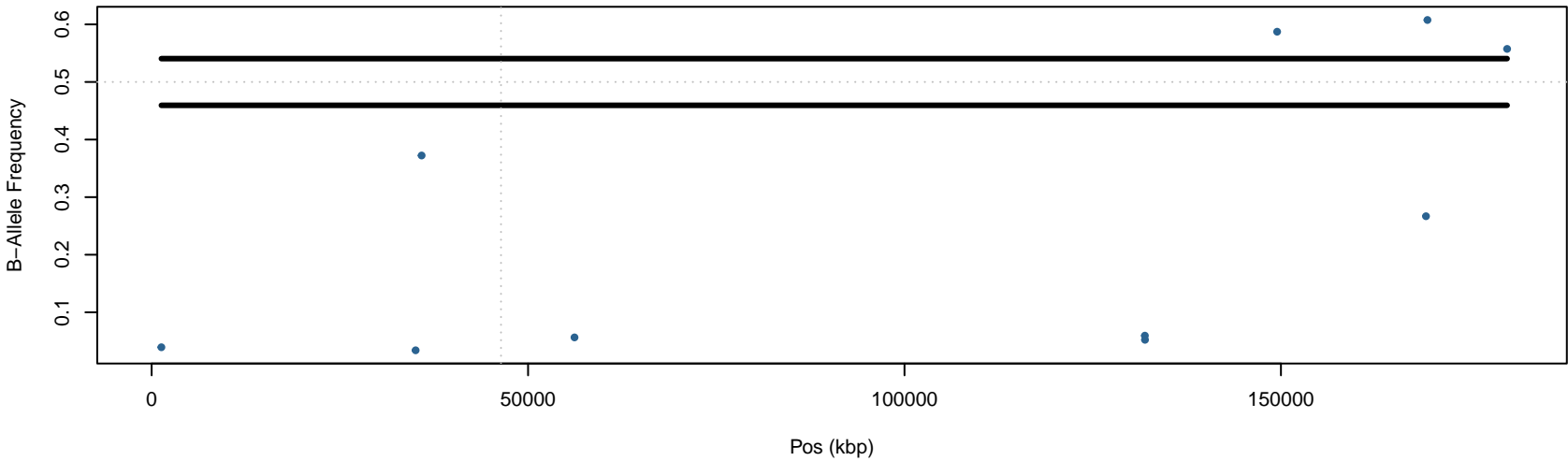
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 3



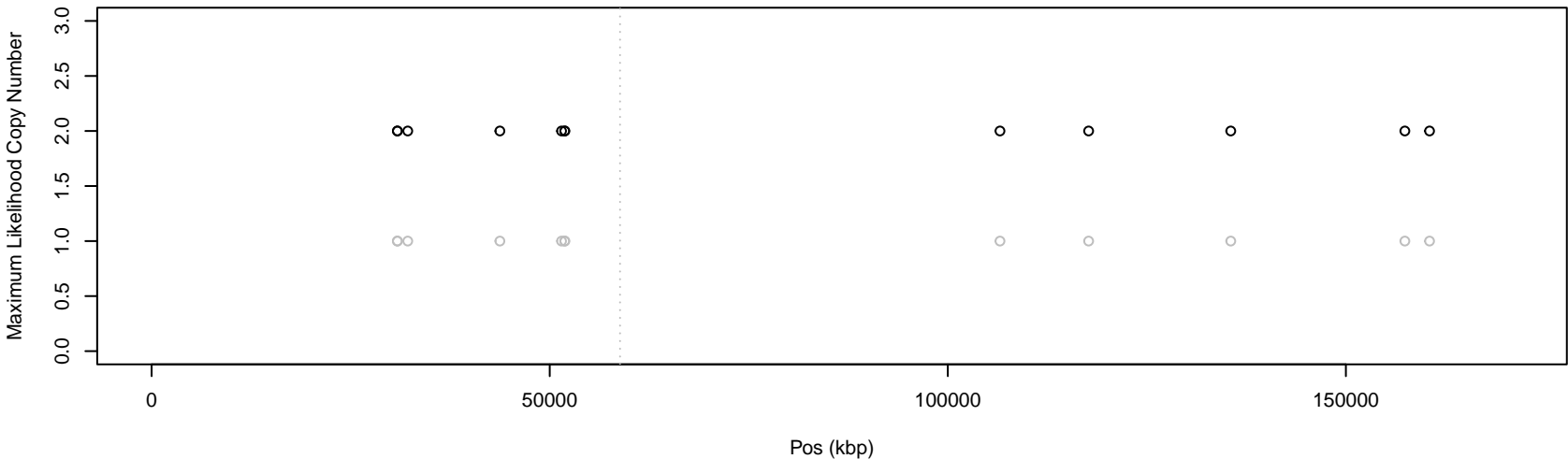
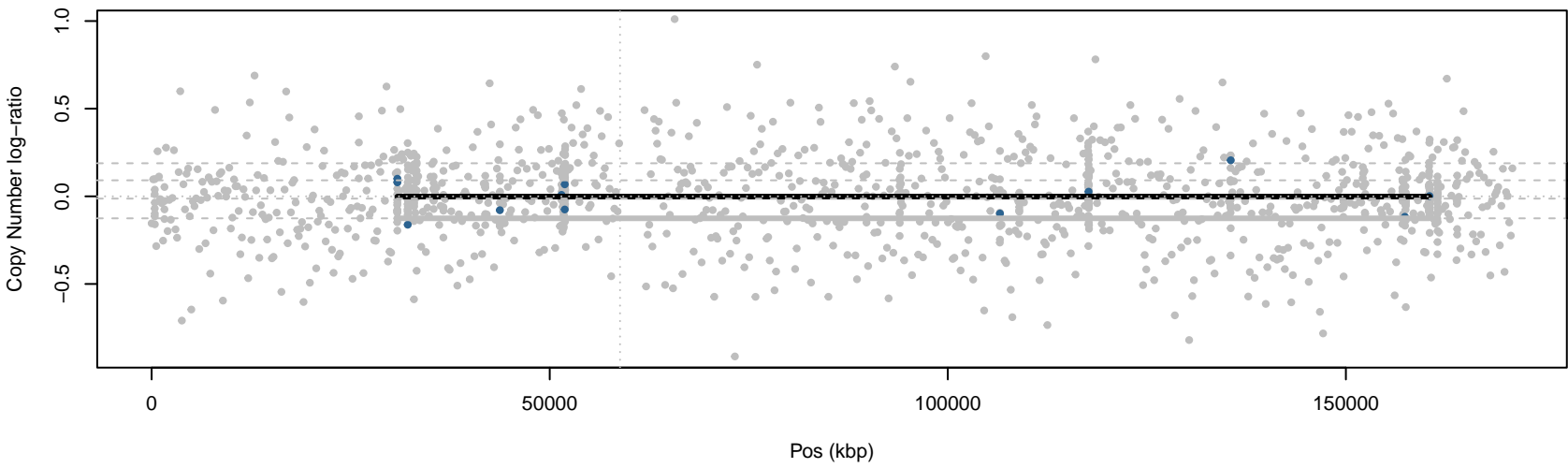
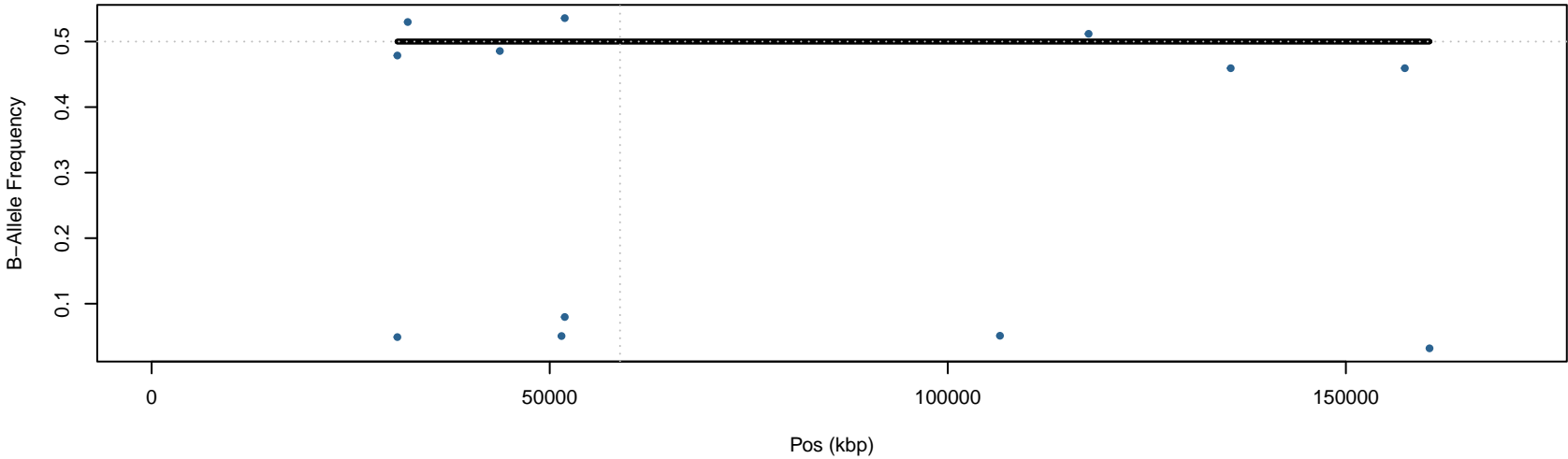
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 4



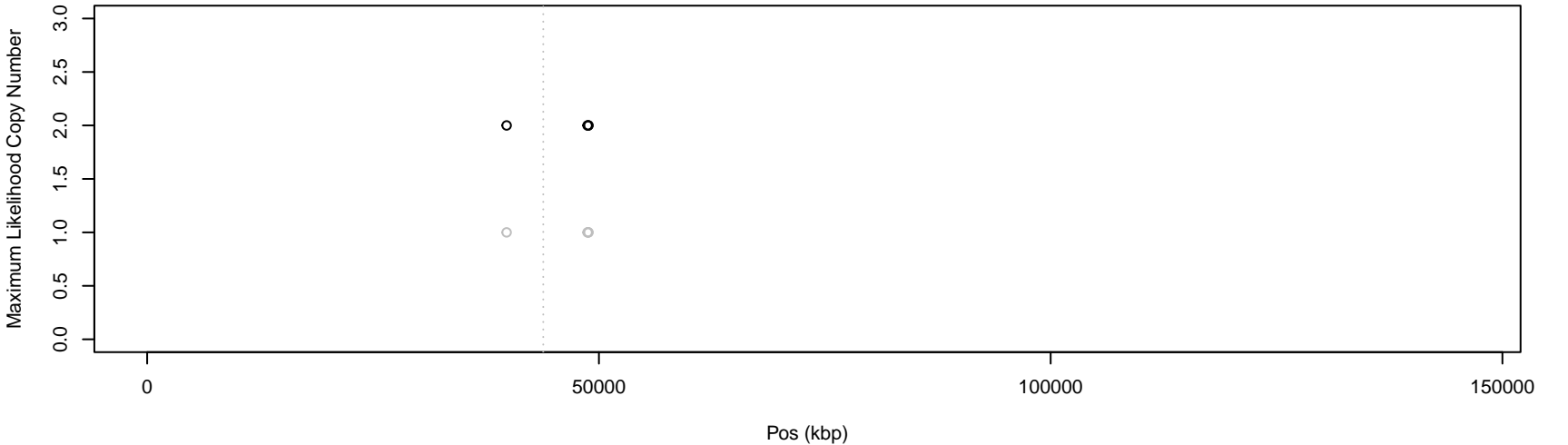
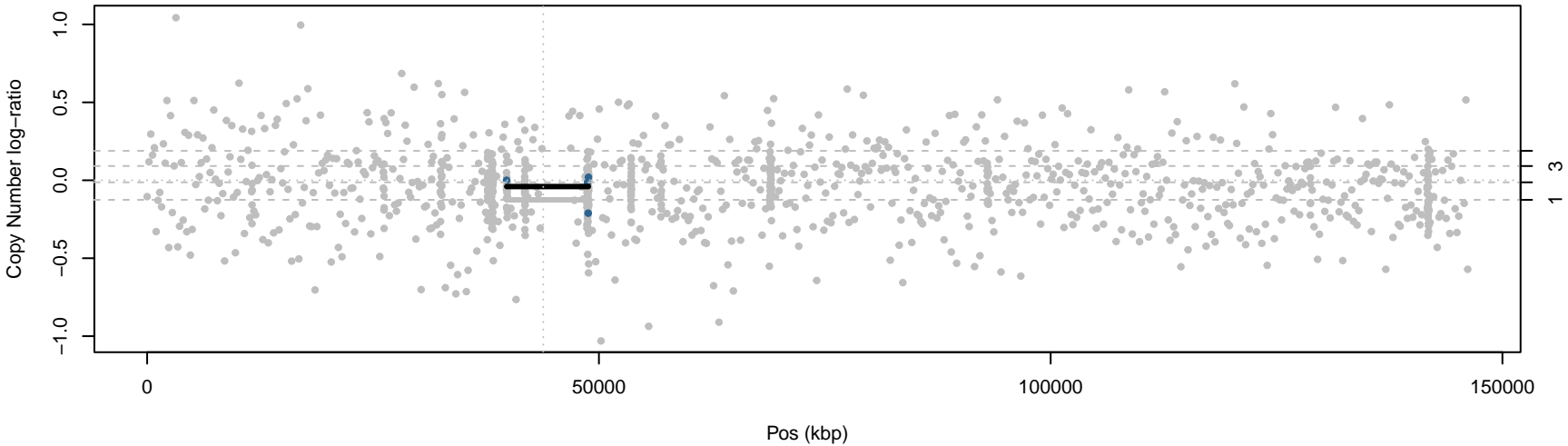
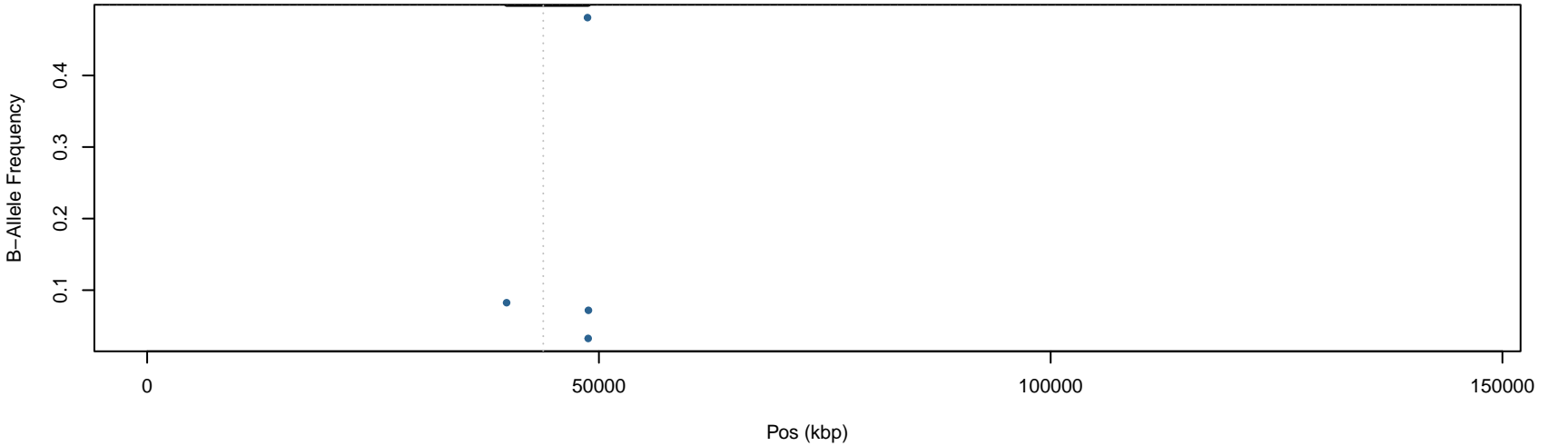
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 5



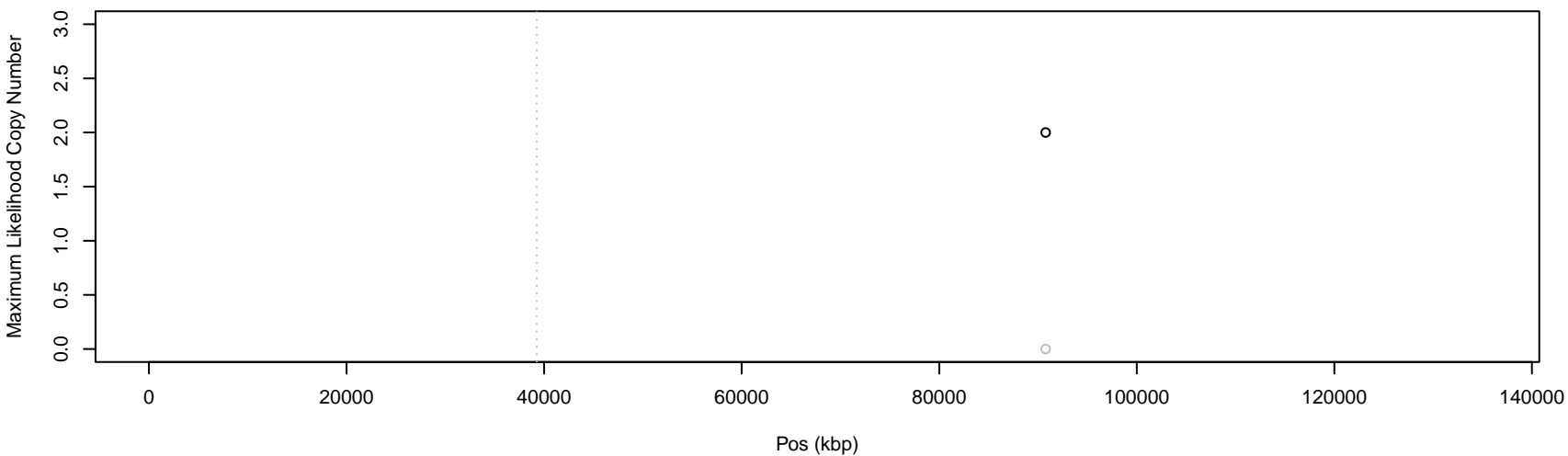
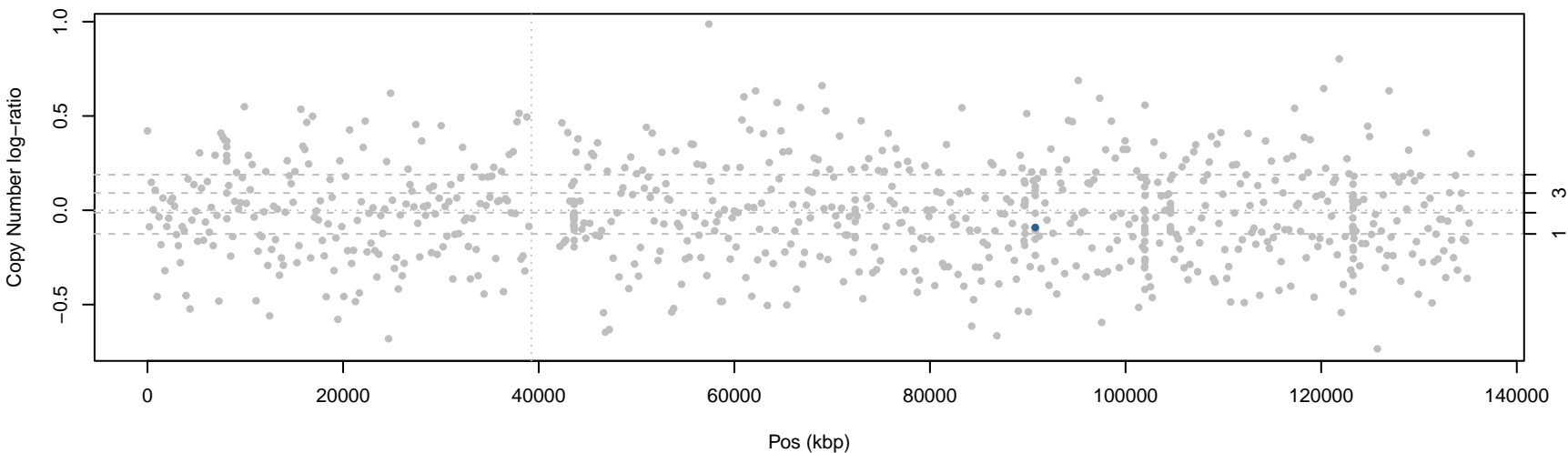
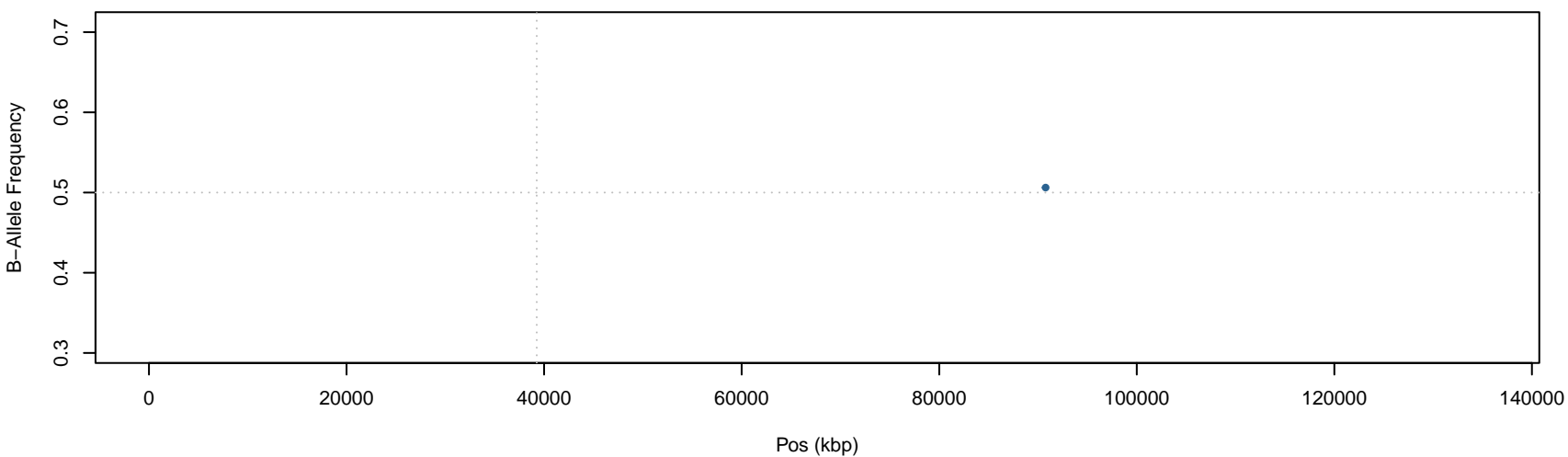
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 6



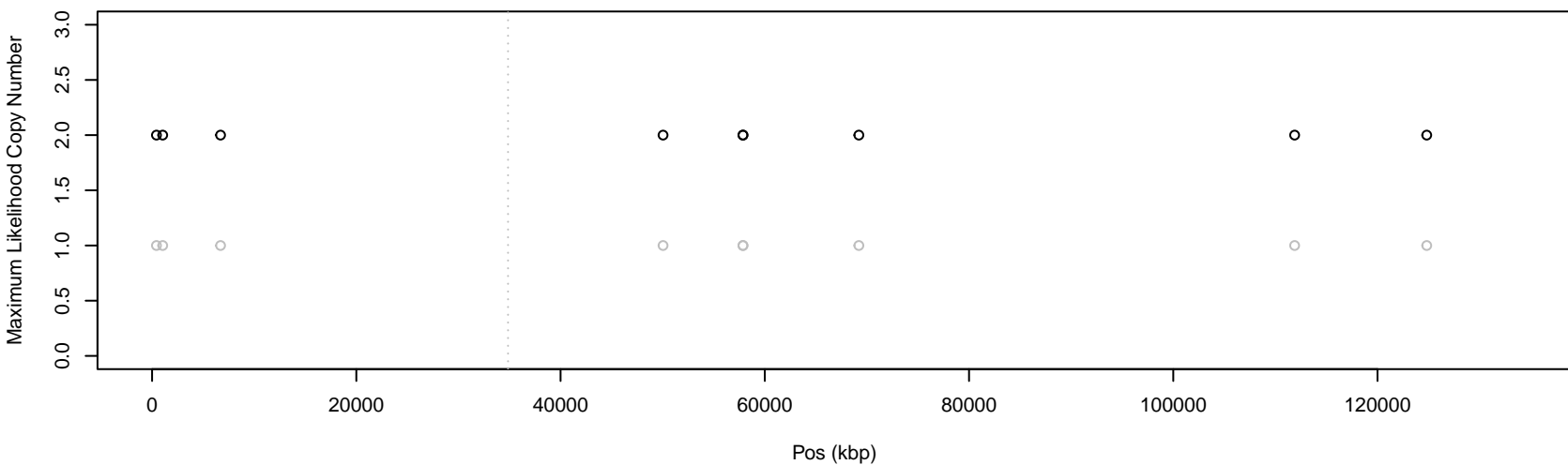
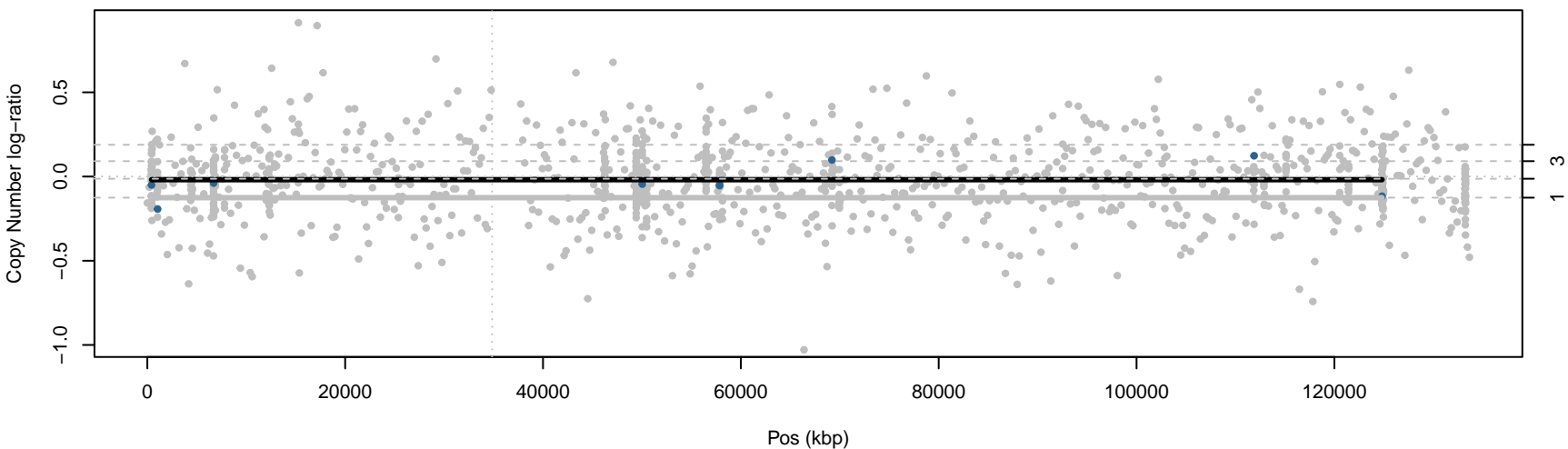
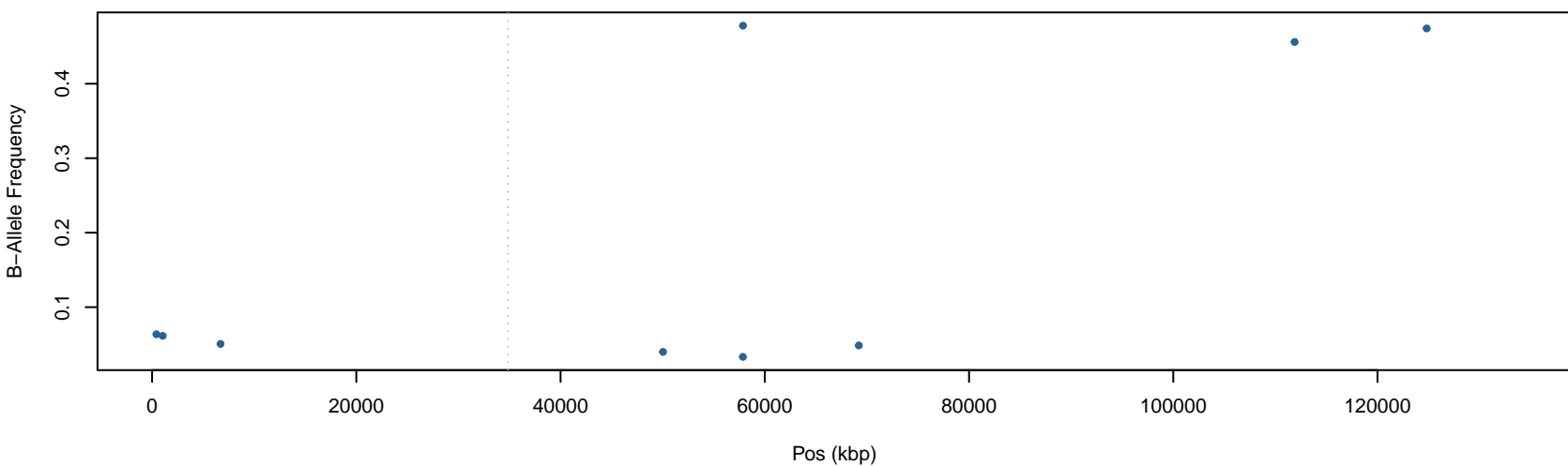
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 8



Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 10

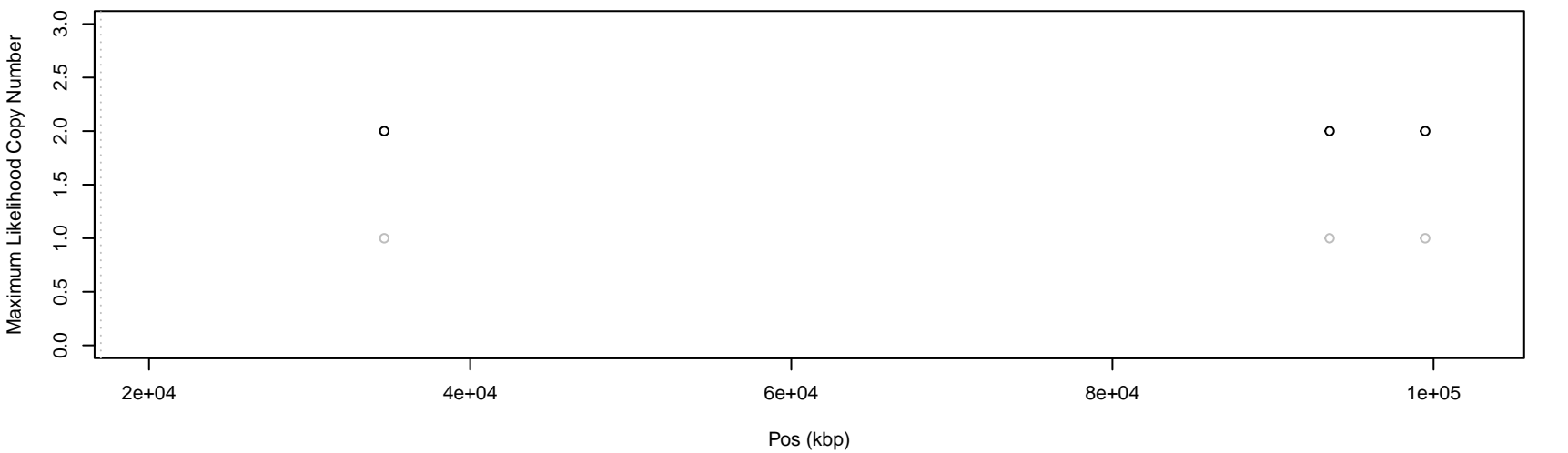
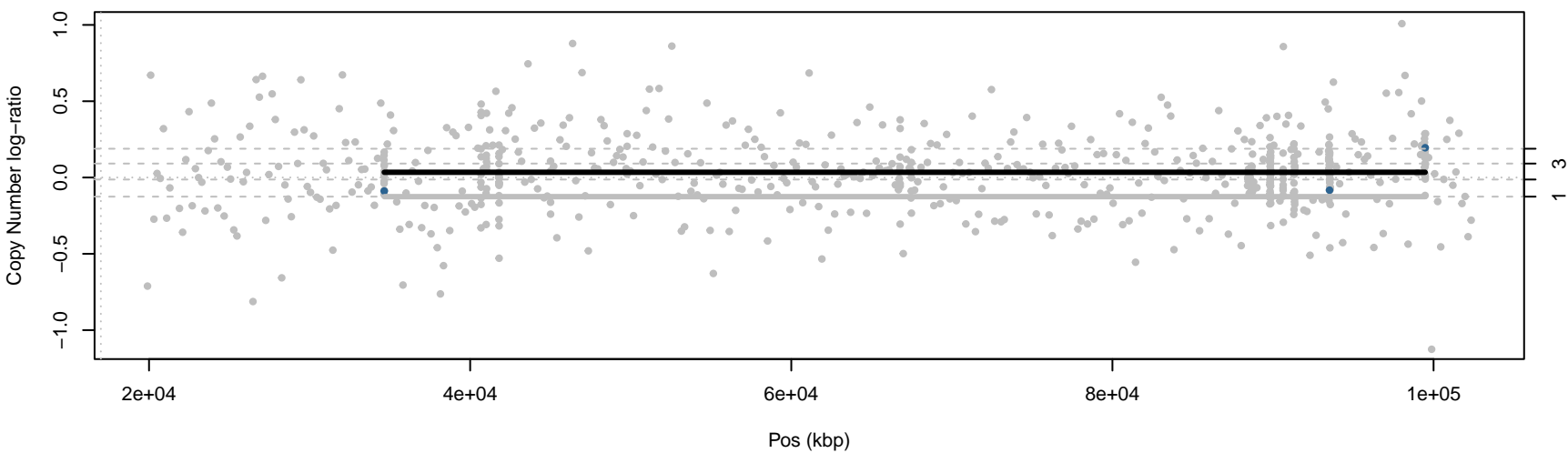
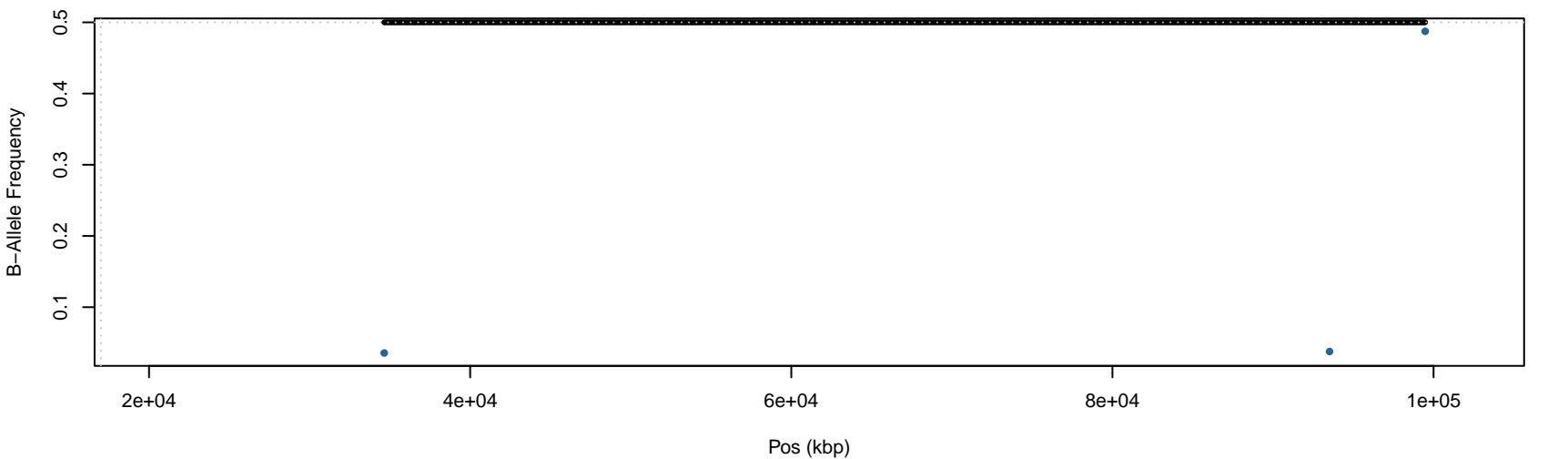


Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 12

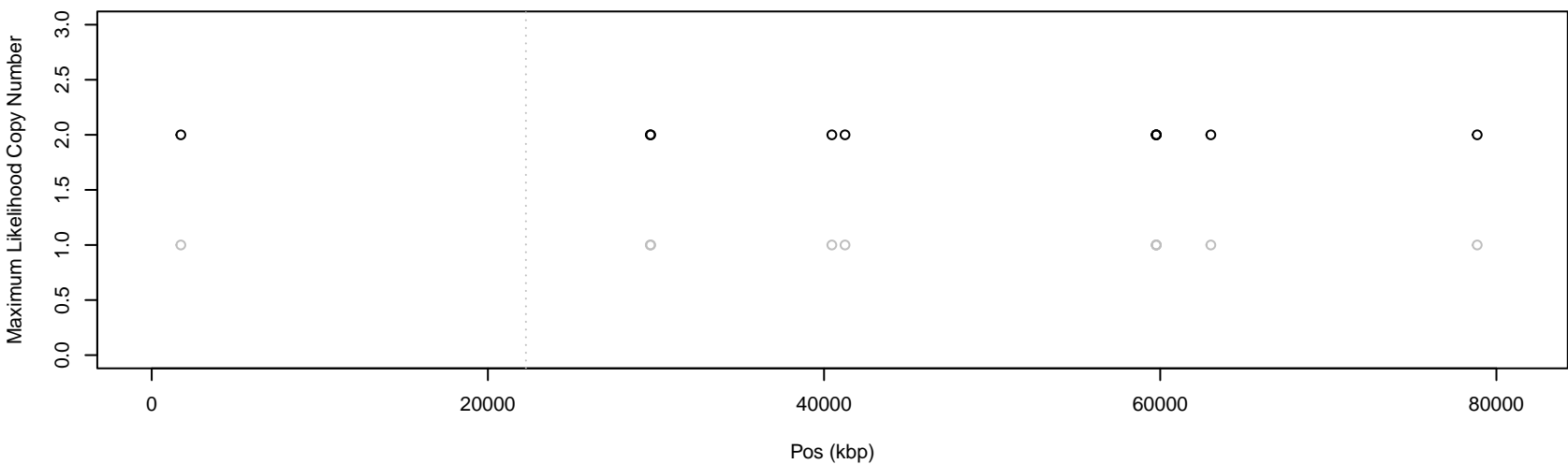
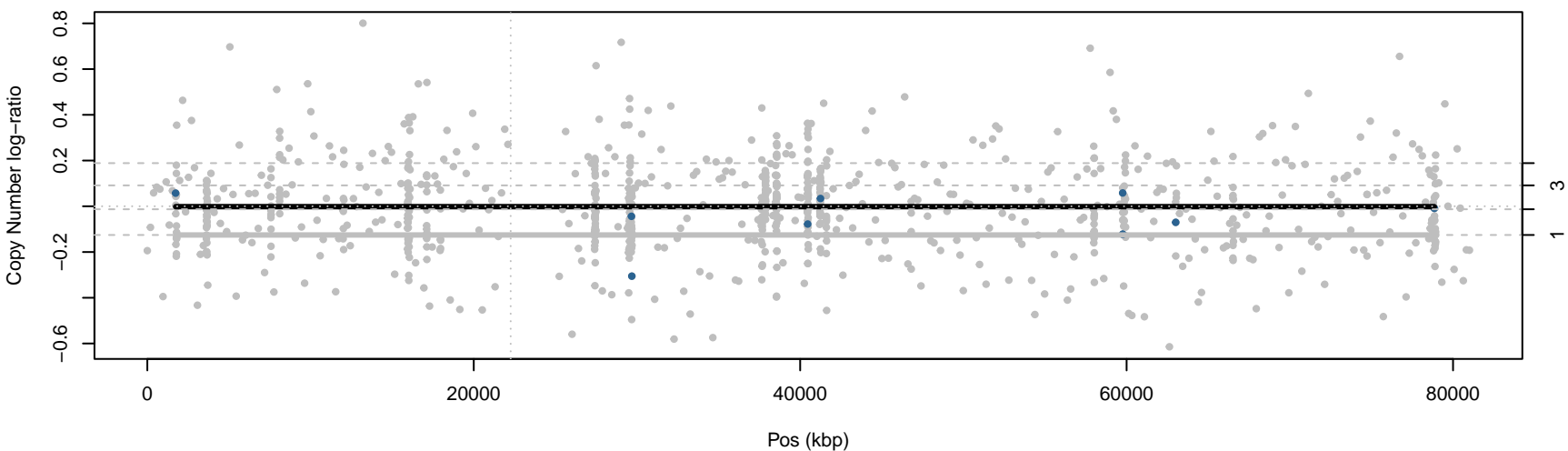
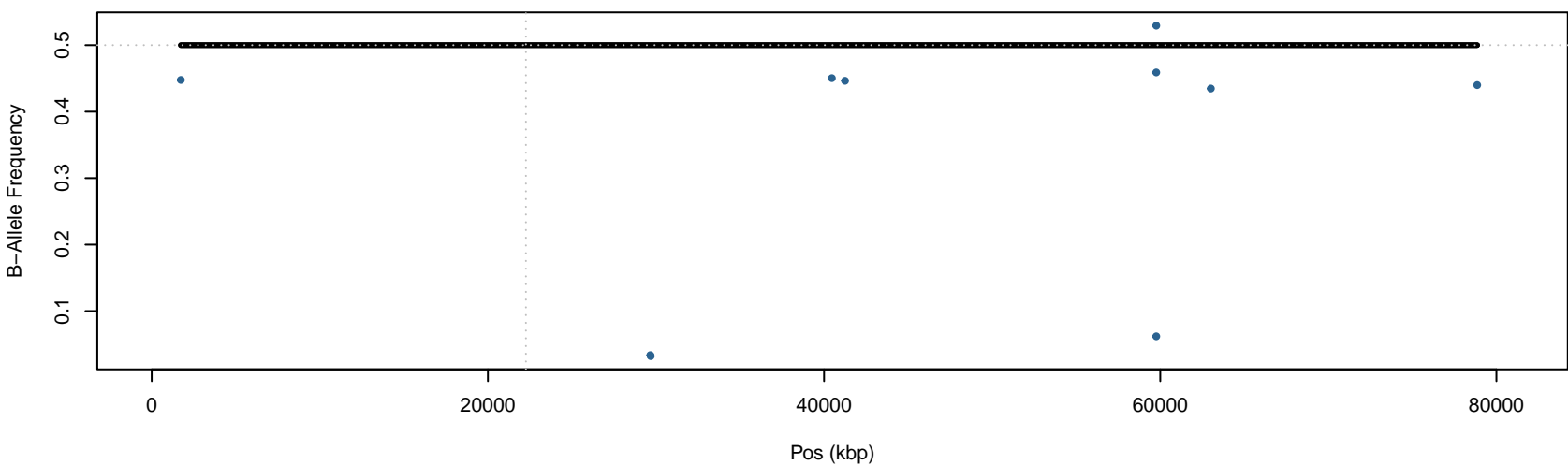




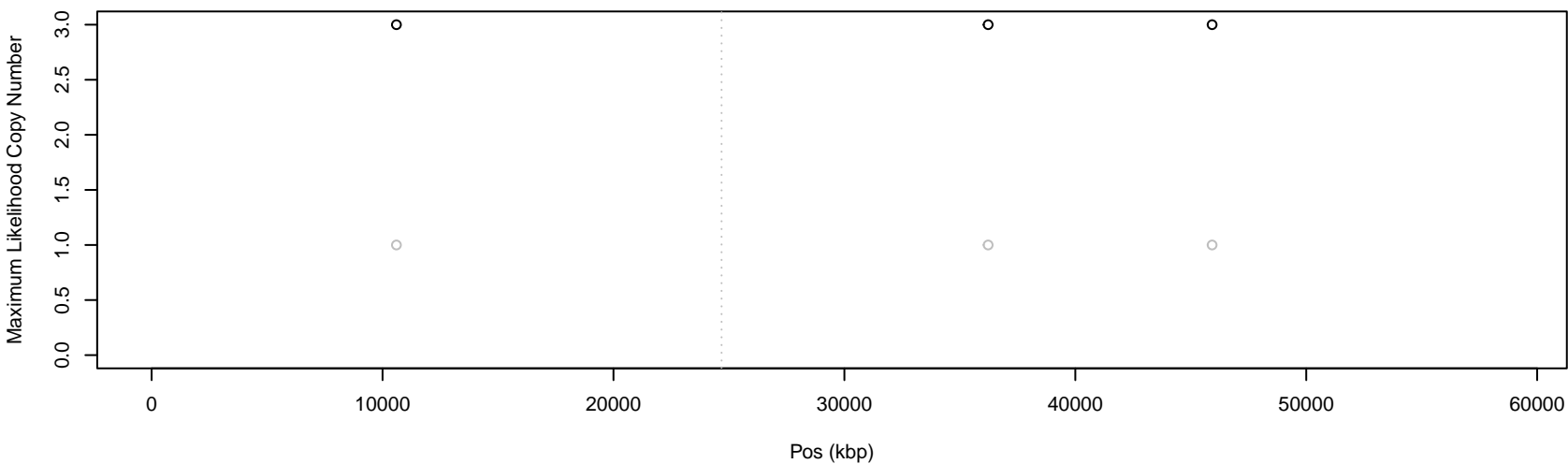
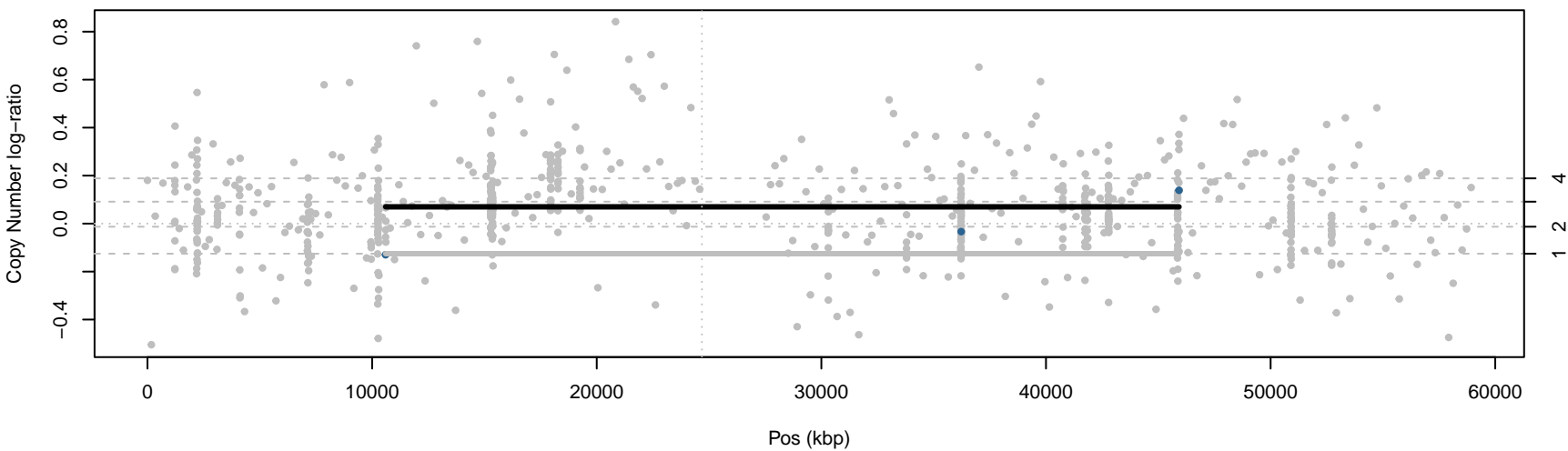
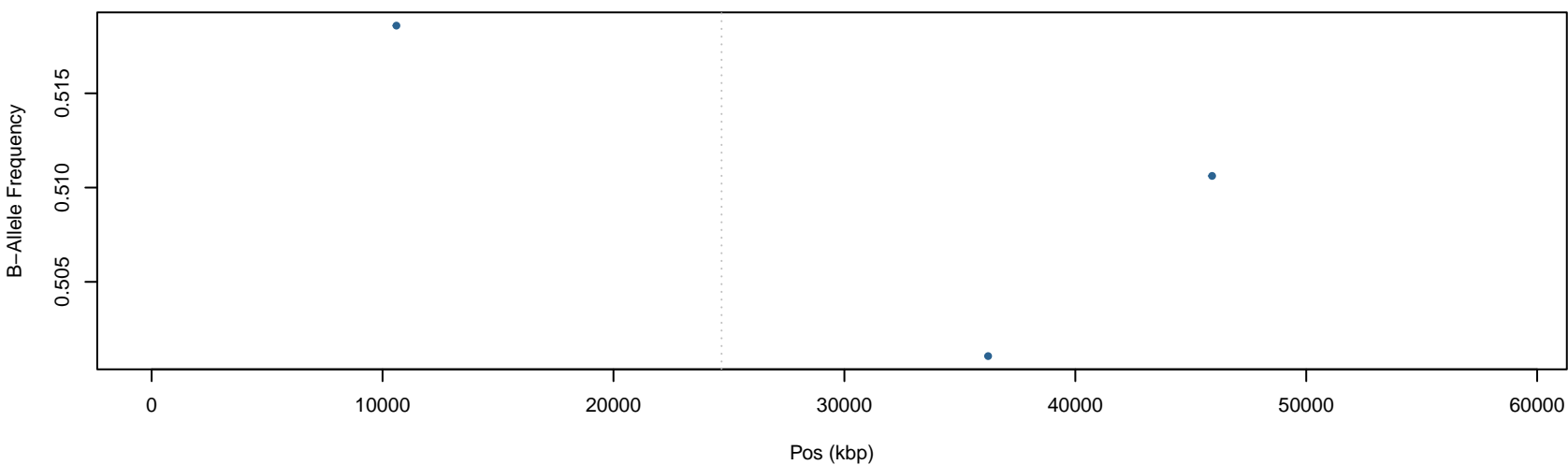
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 15



Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 17



Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 19



Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394 Chromosome: 20

