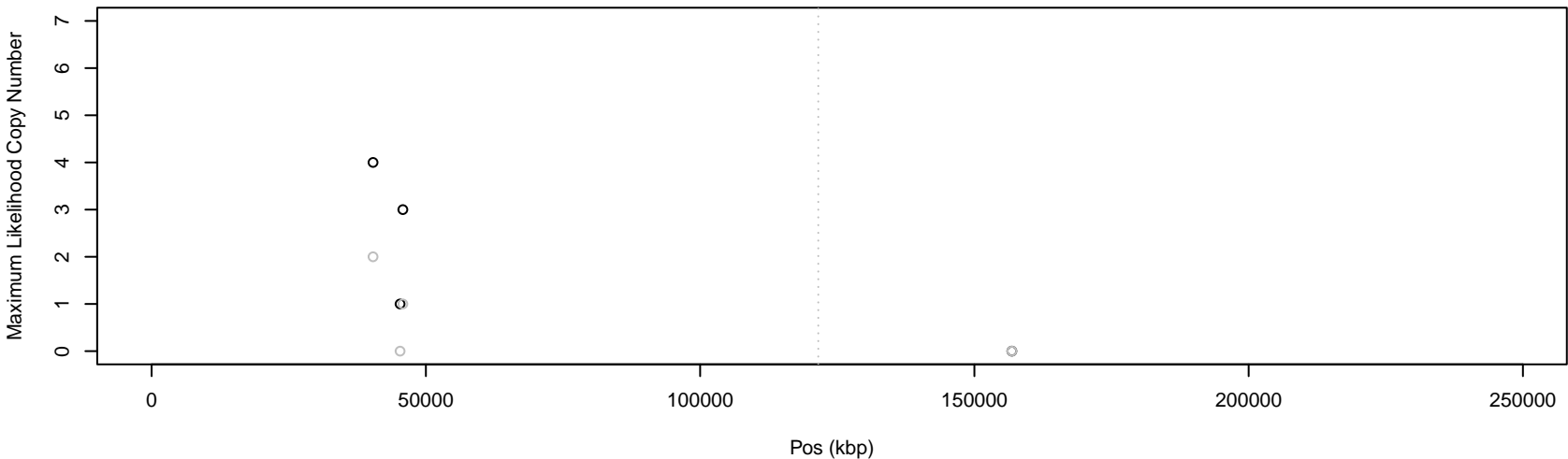
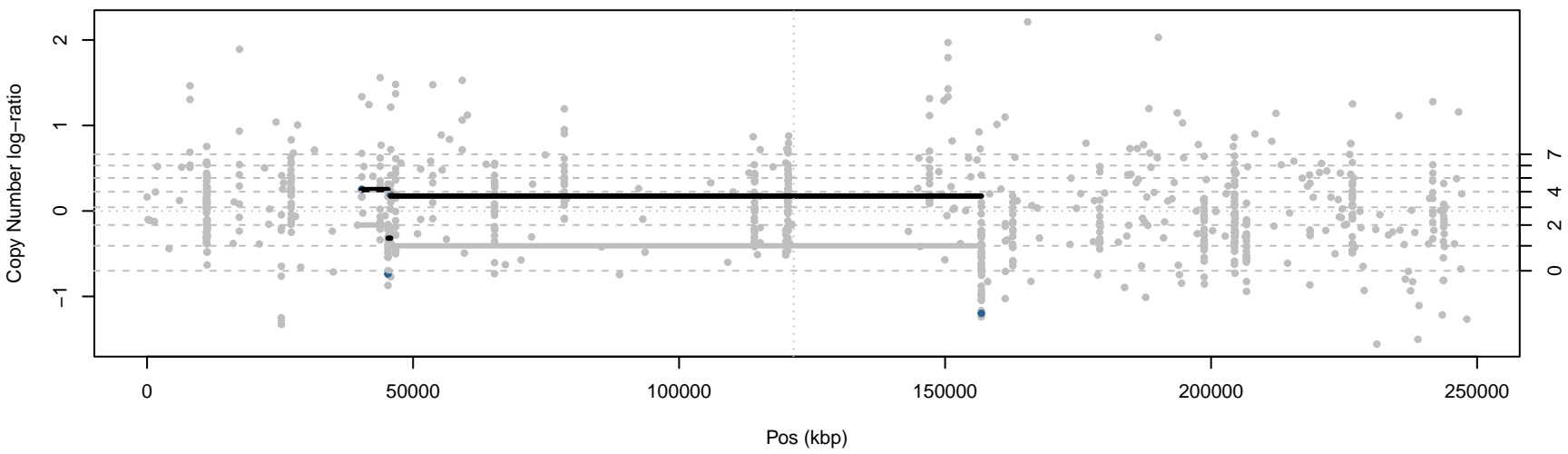
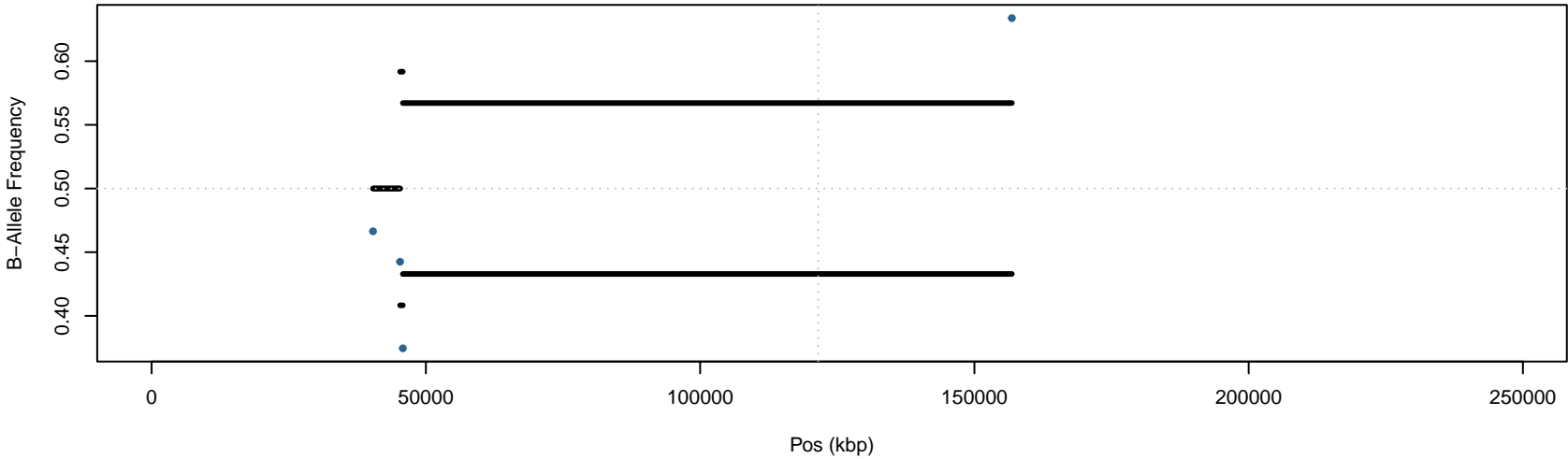
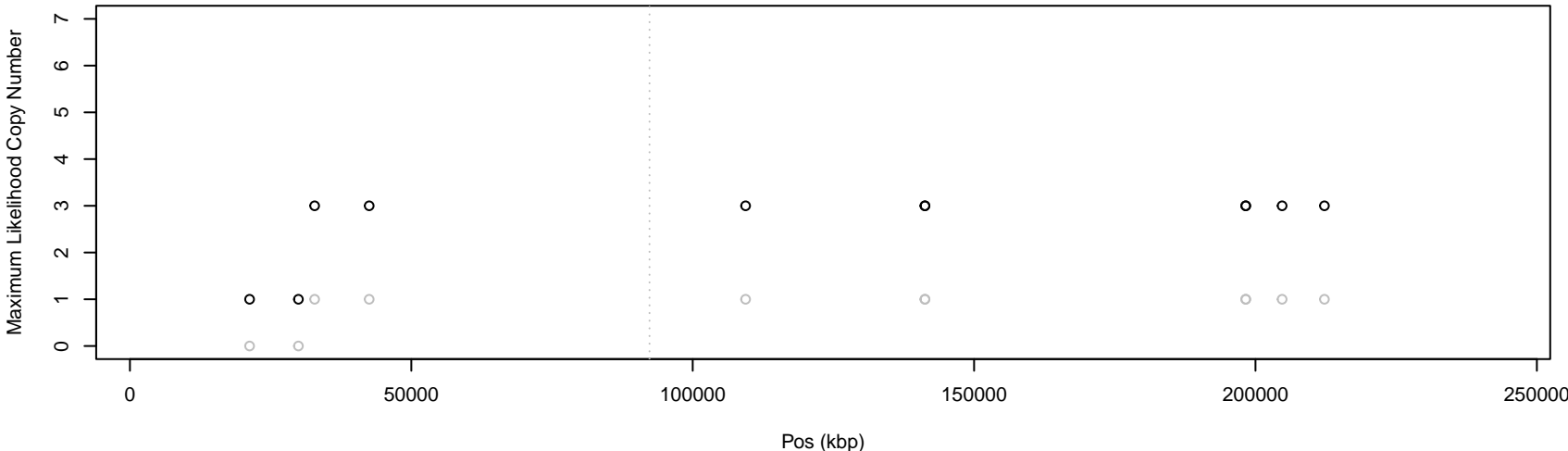
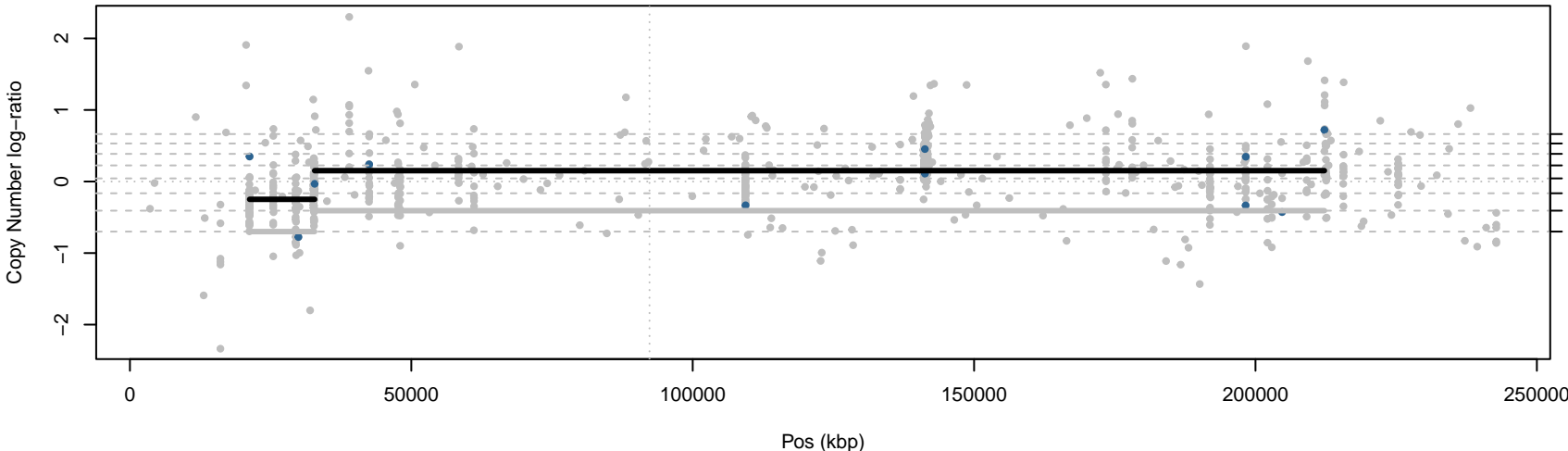
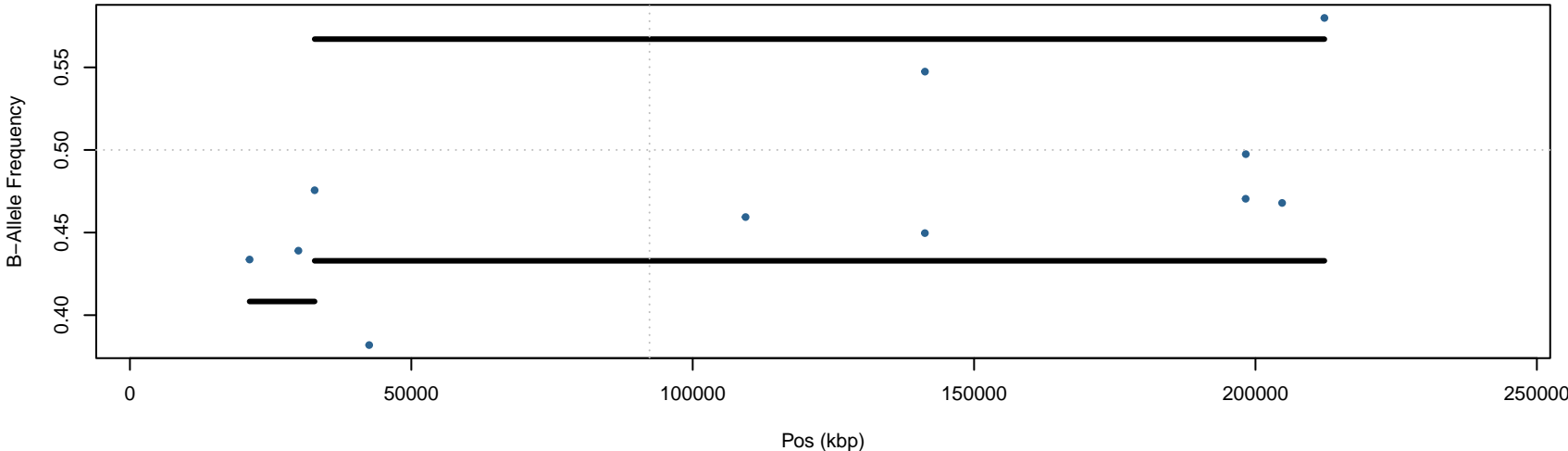
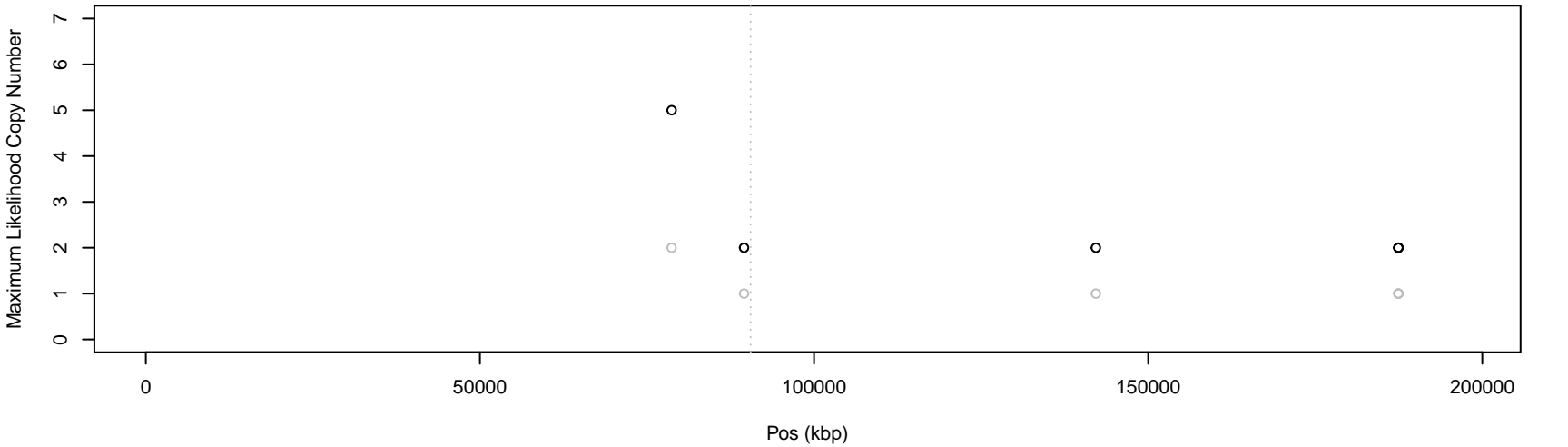
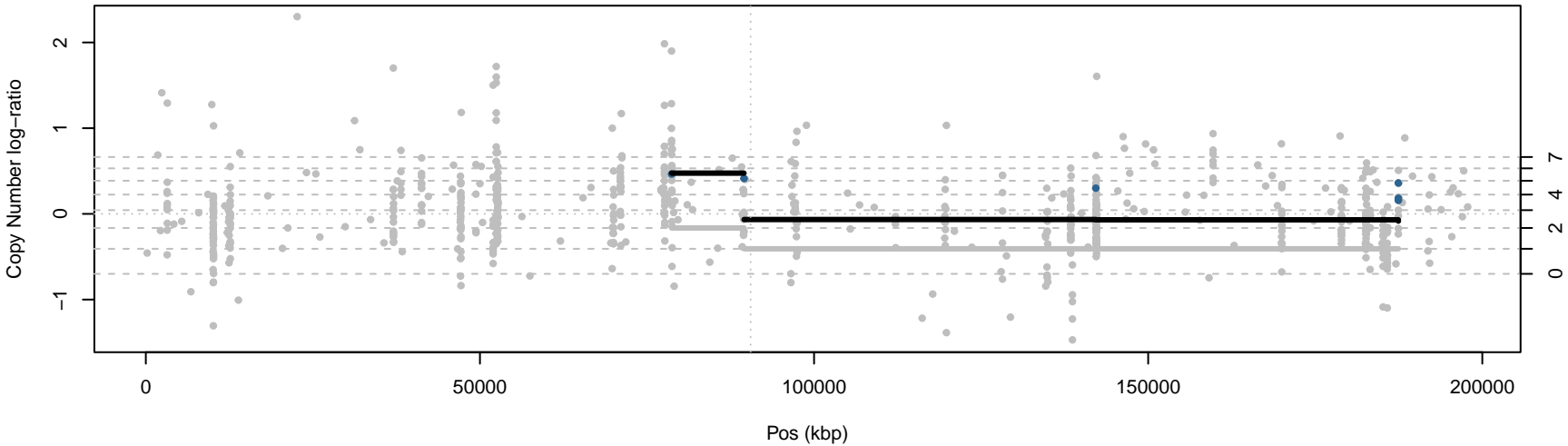
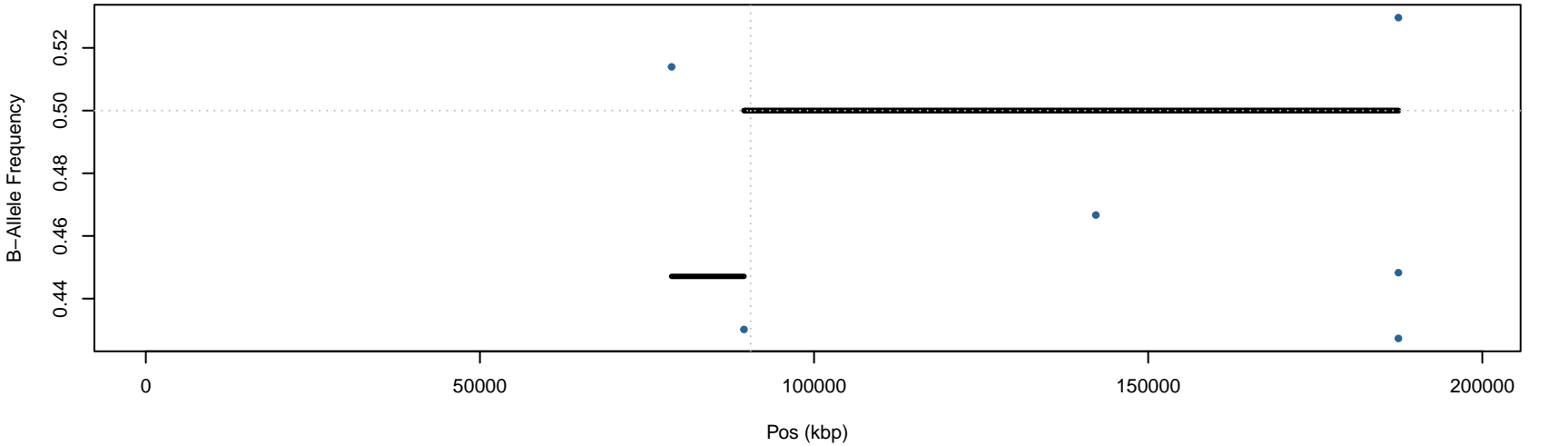


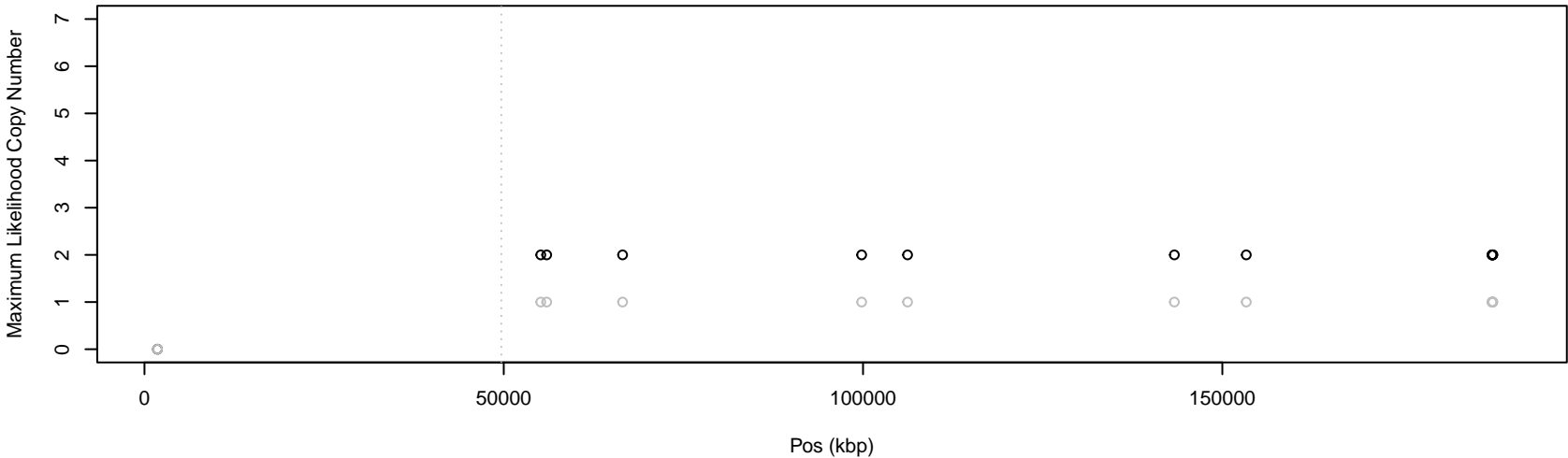
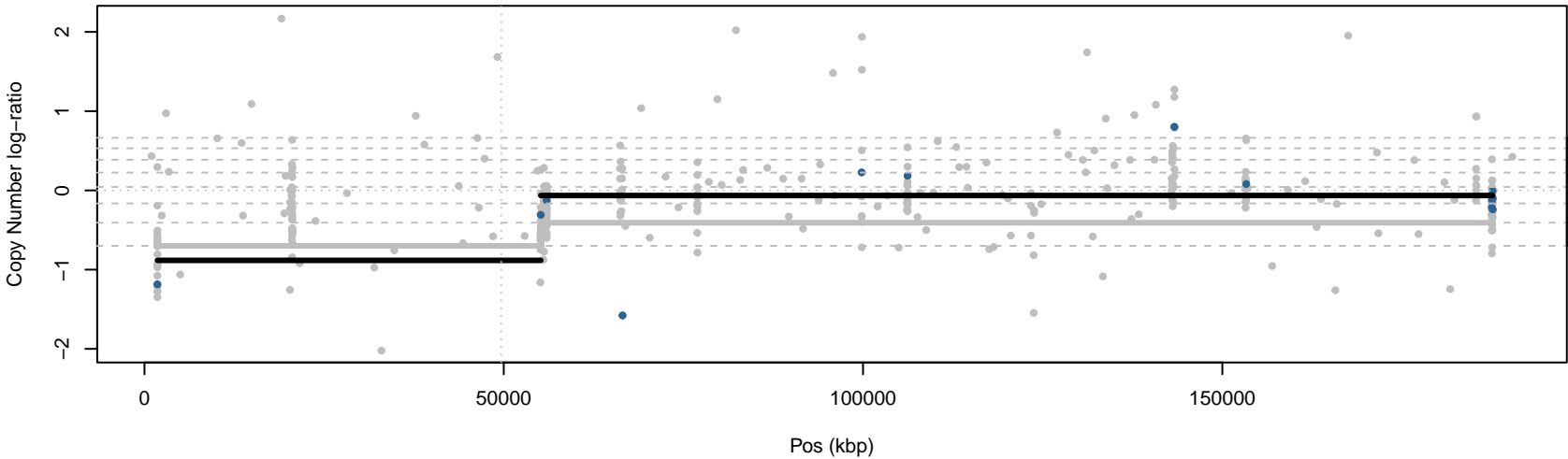
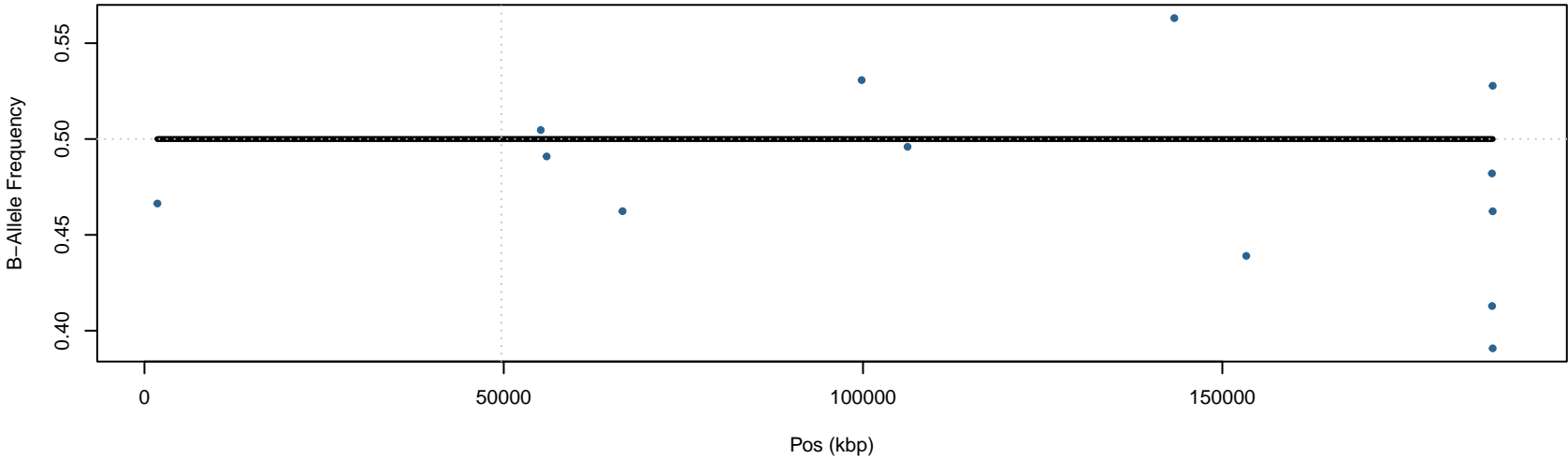
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 1

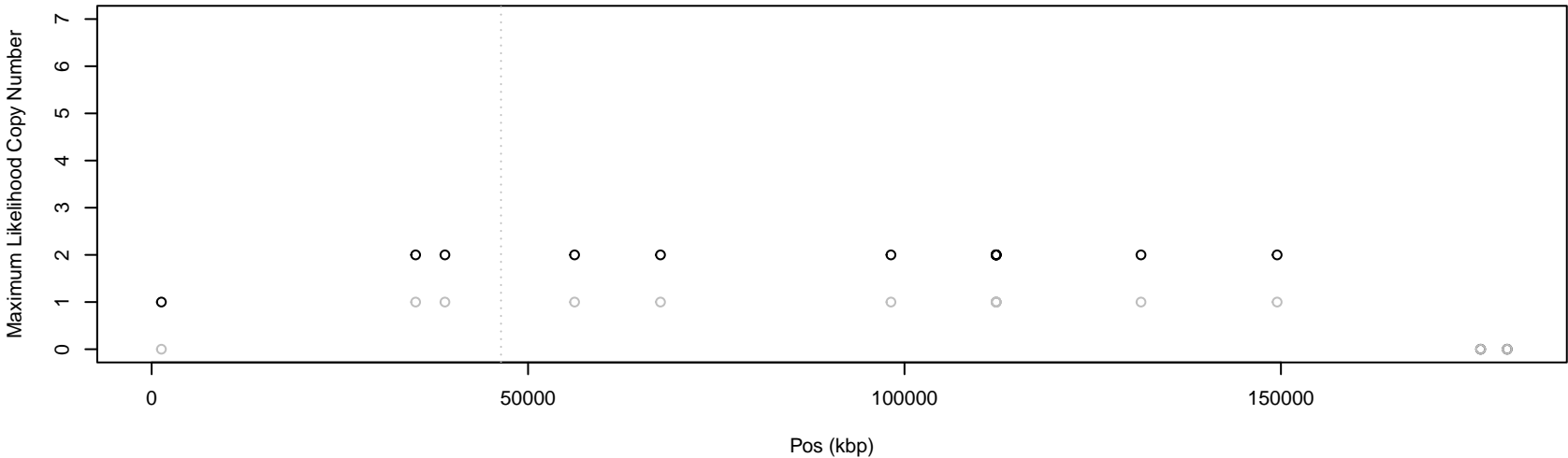
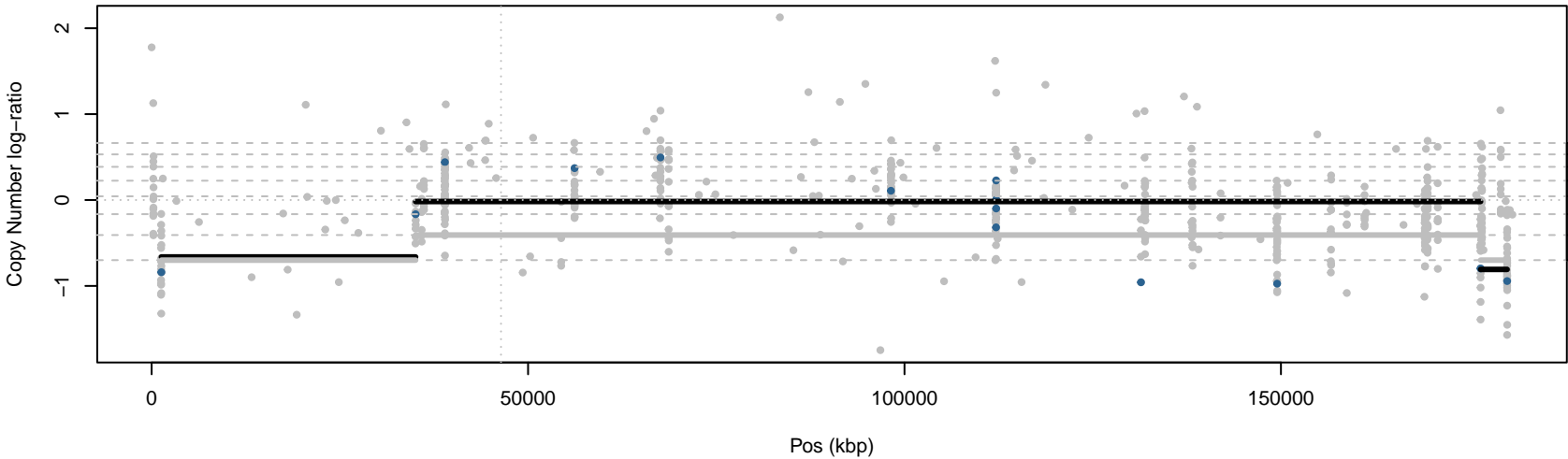
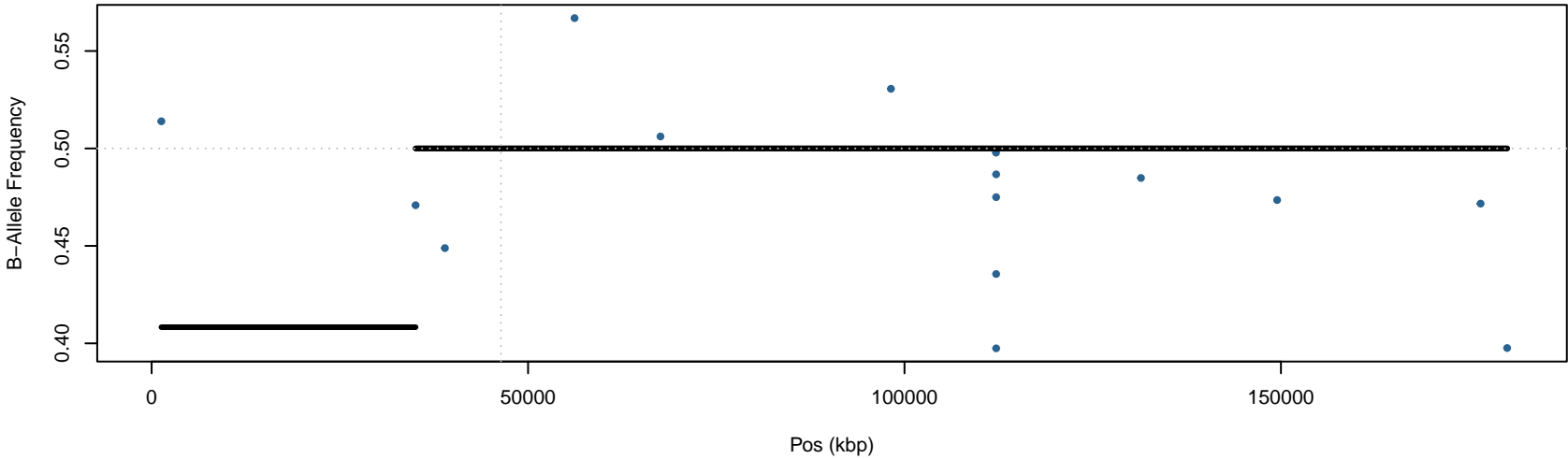


Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 2

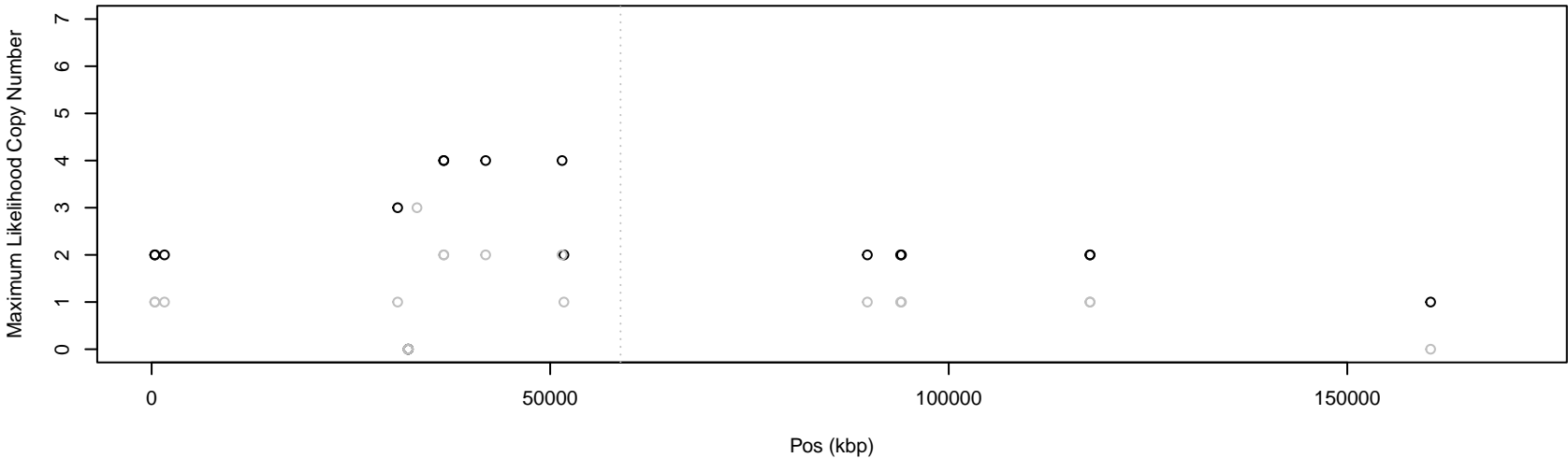
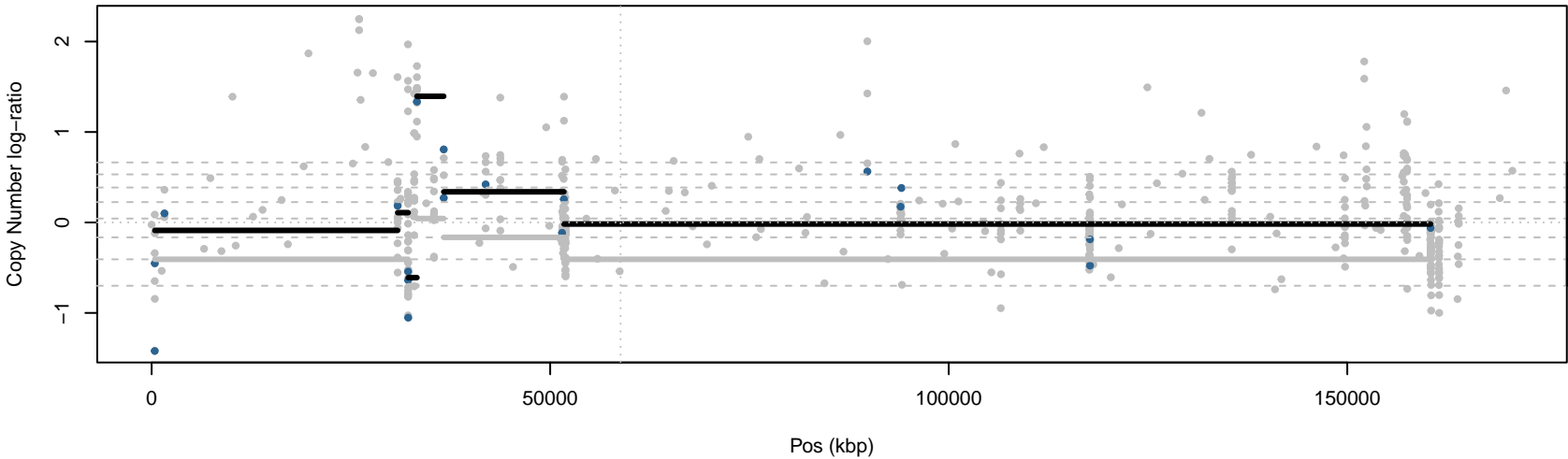
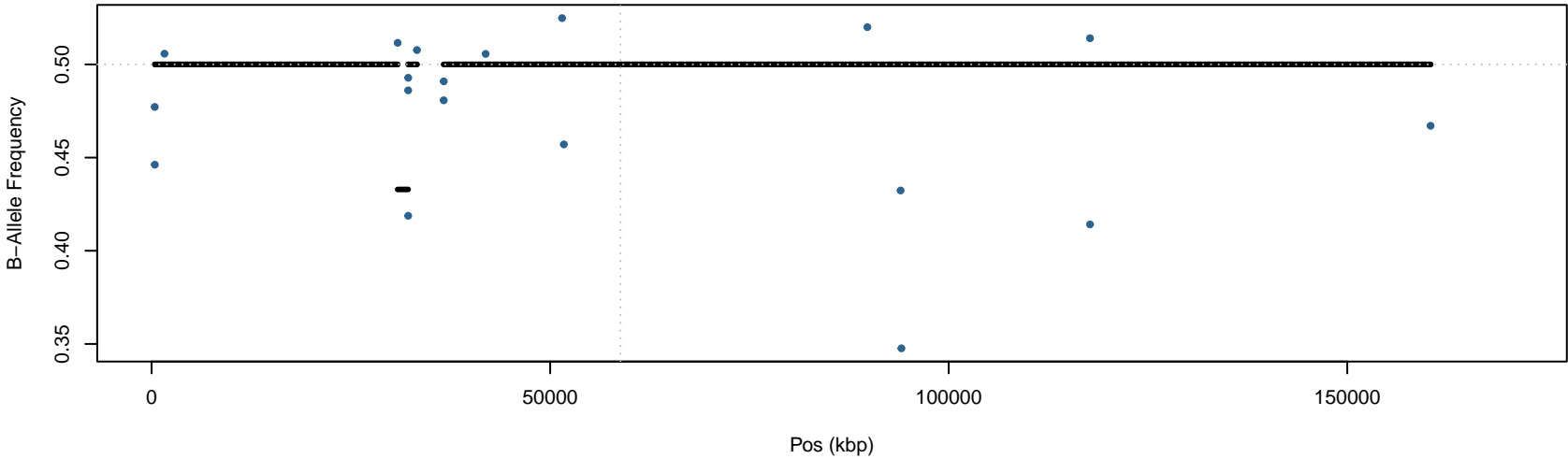


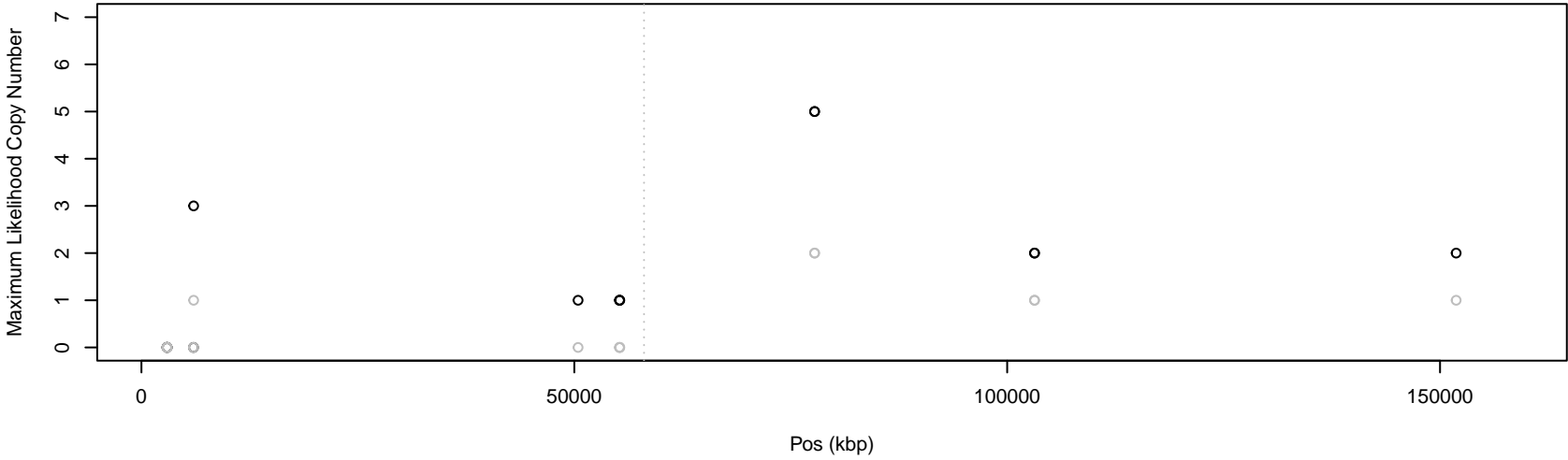
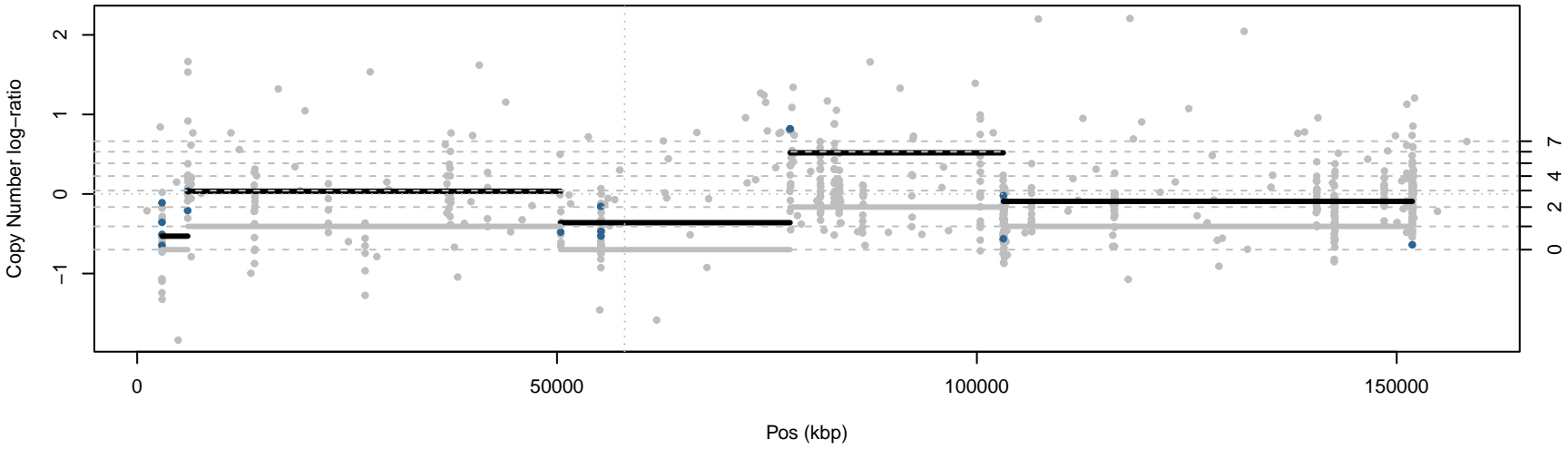
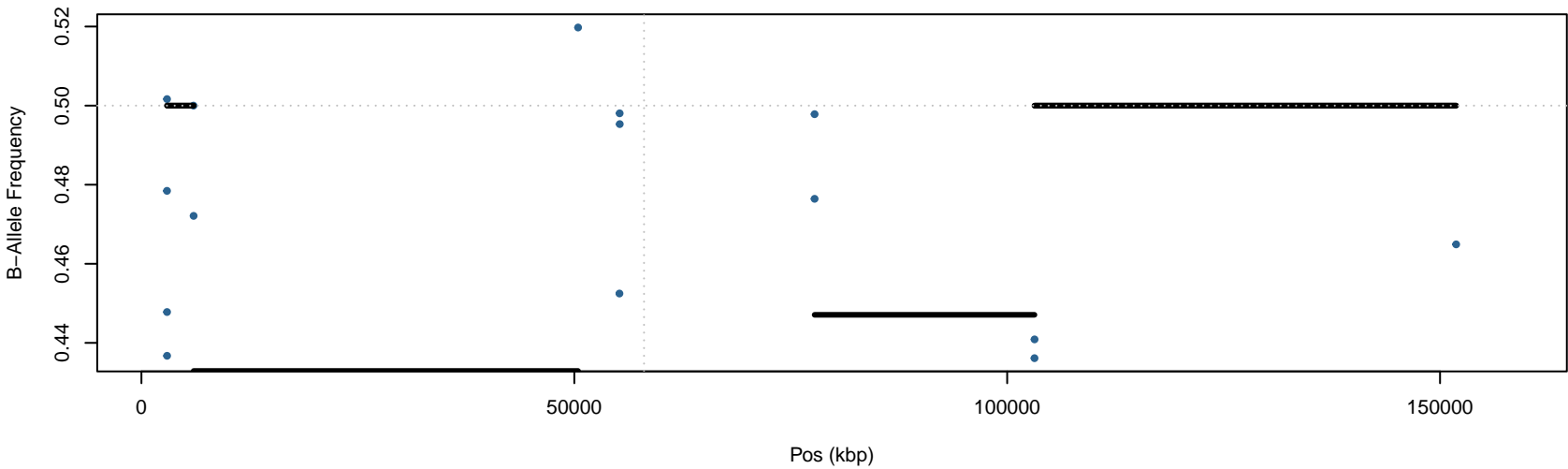


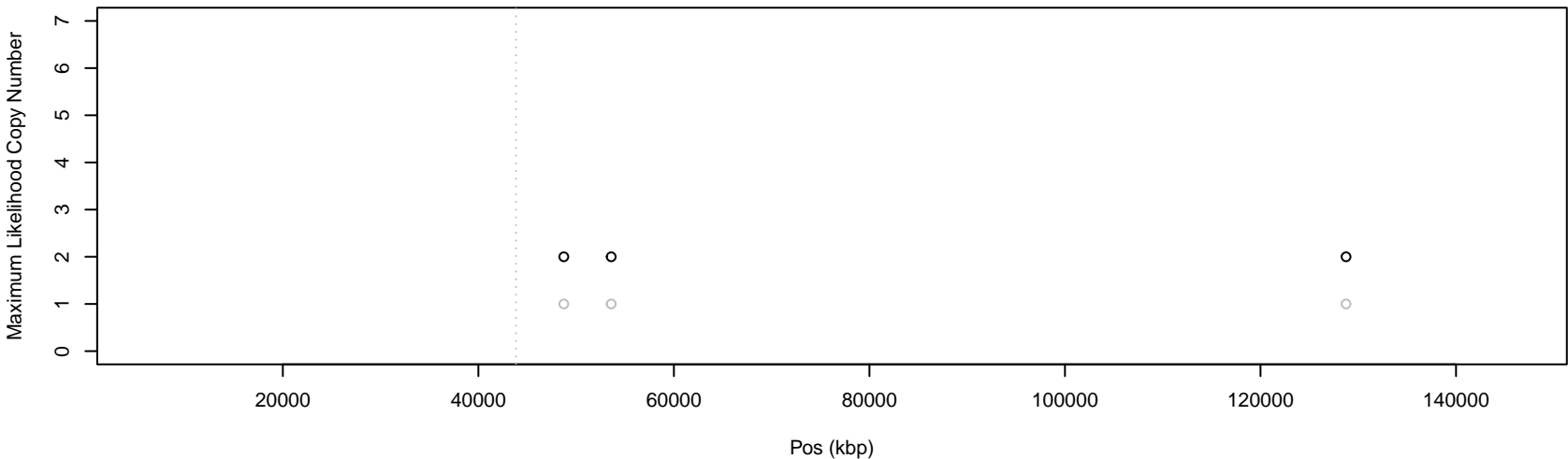
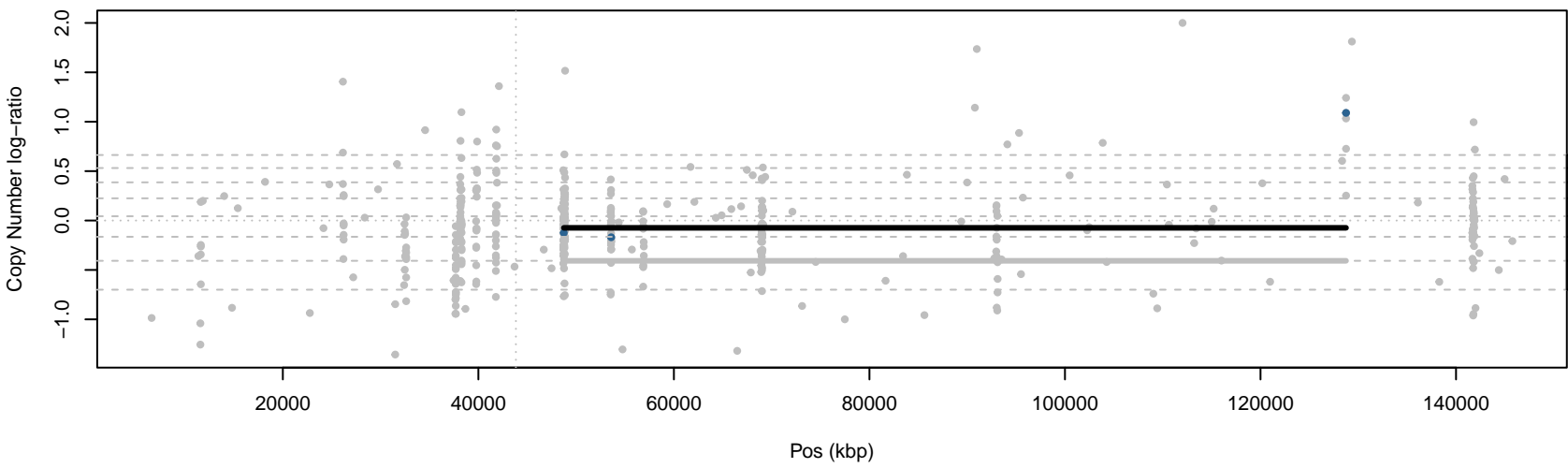
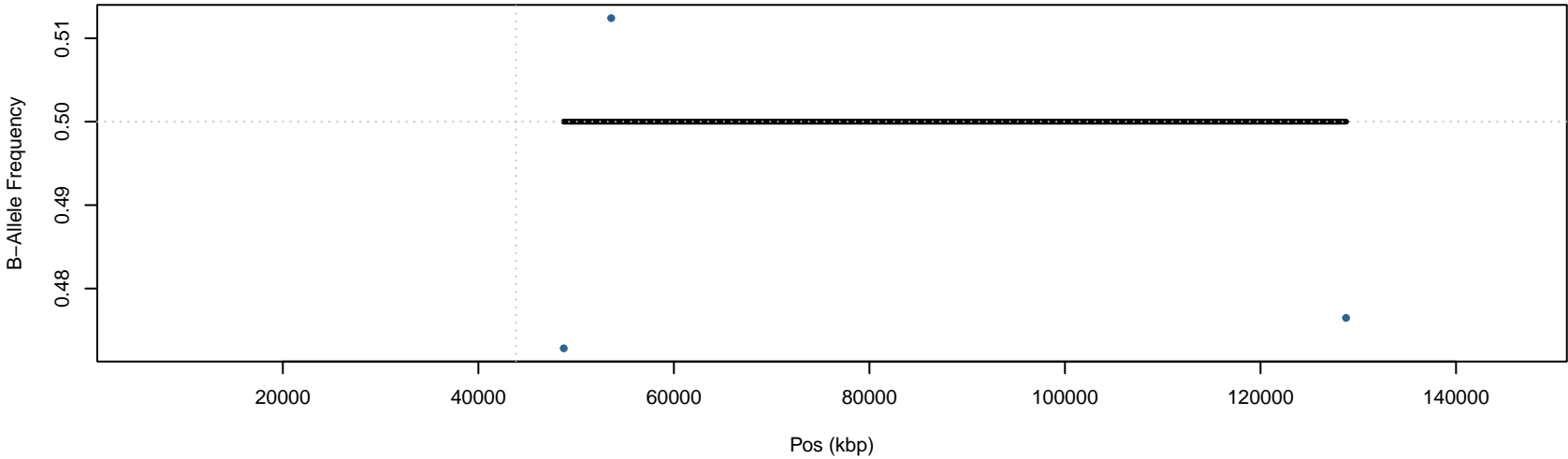




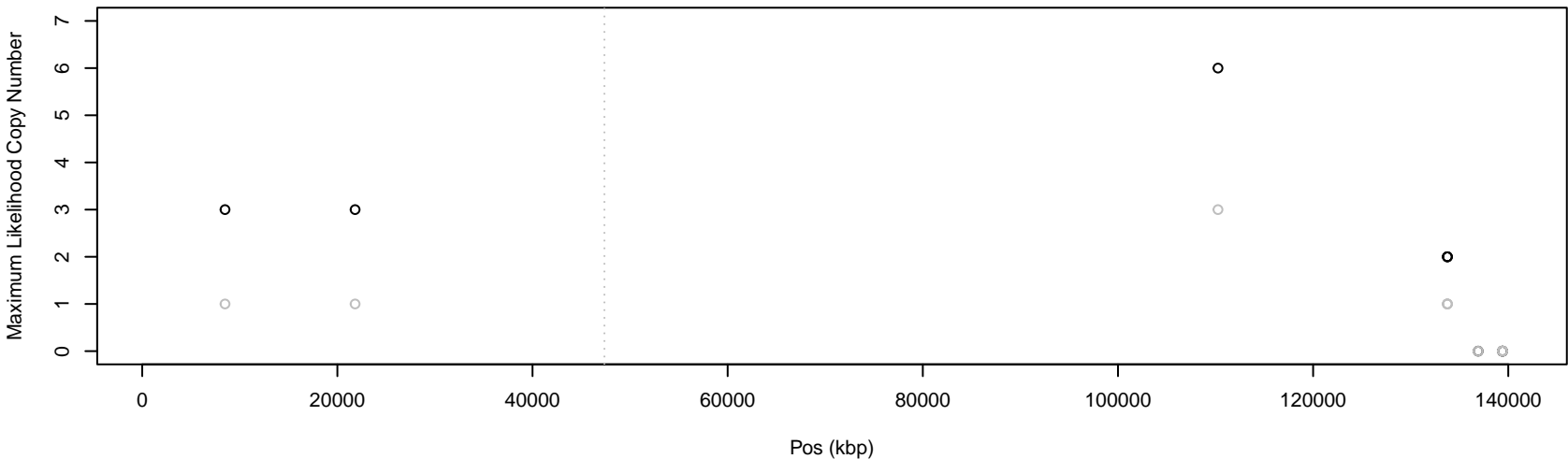
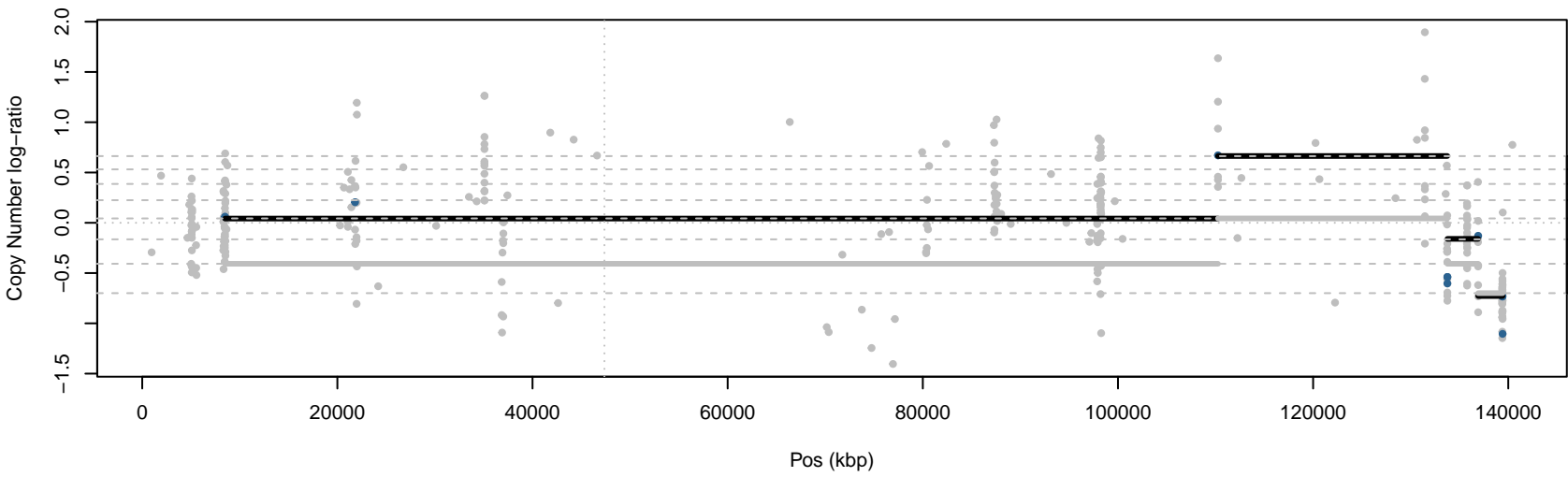
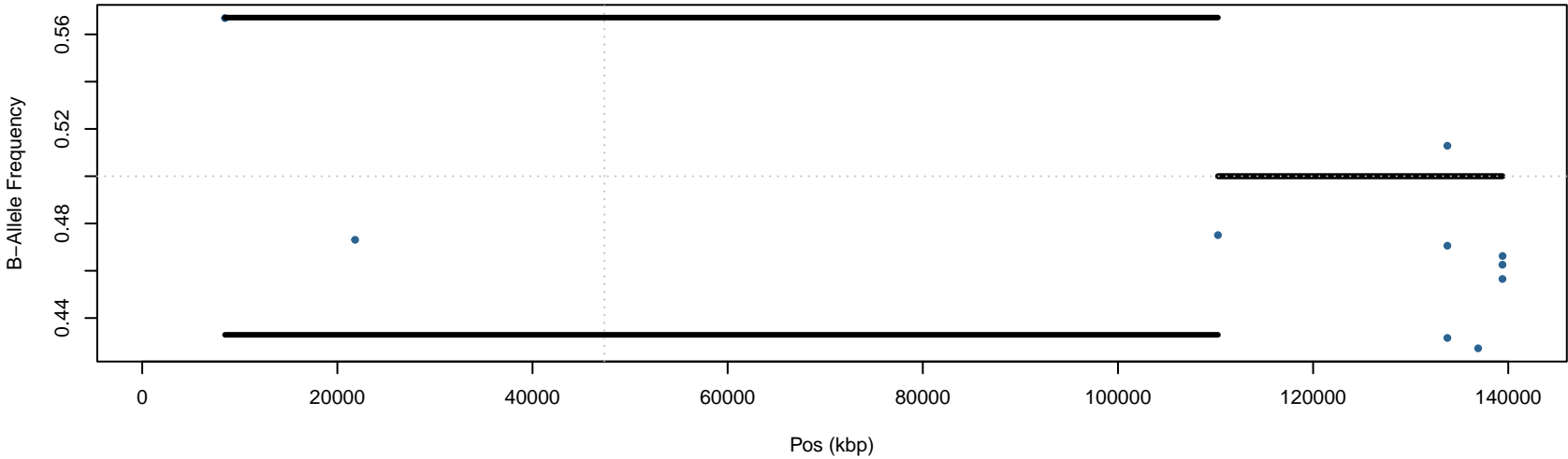
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 6



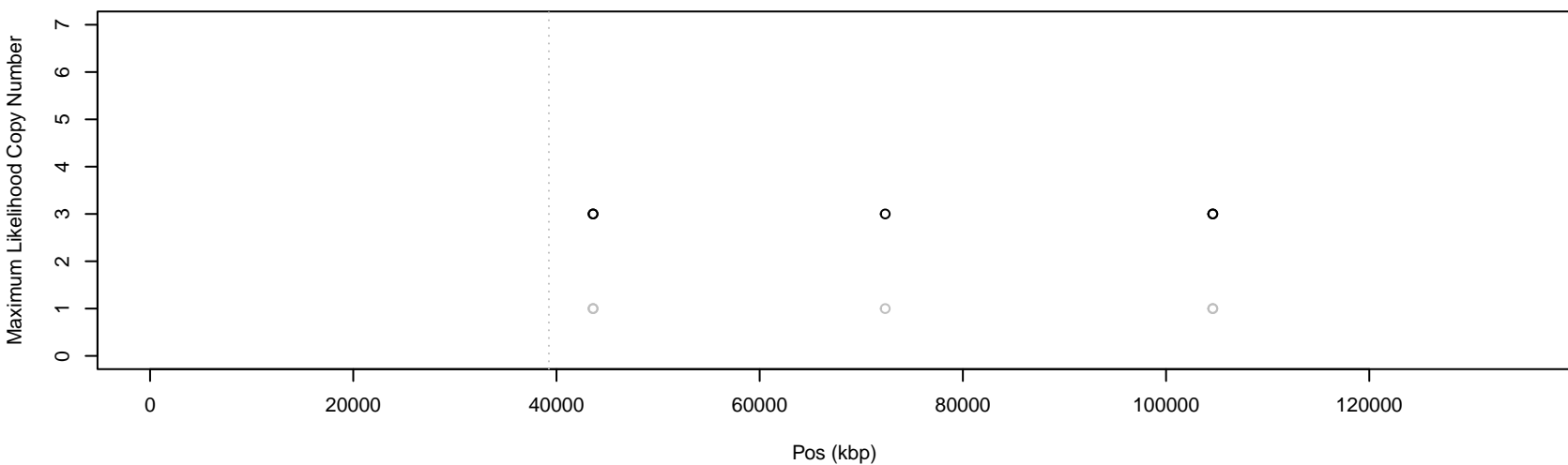
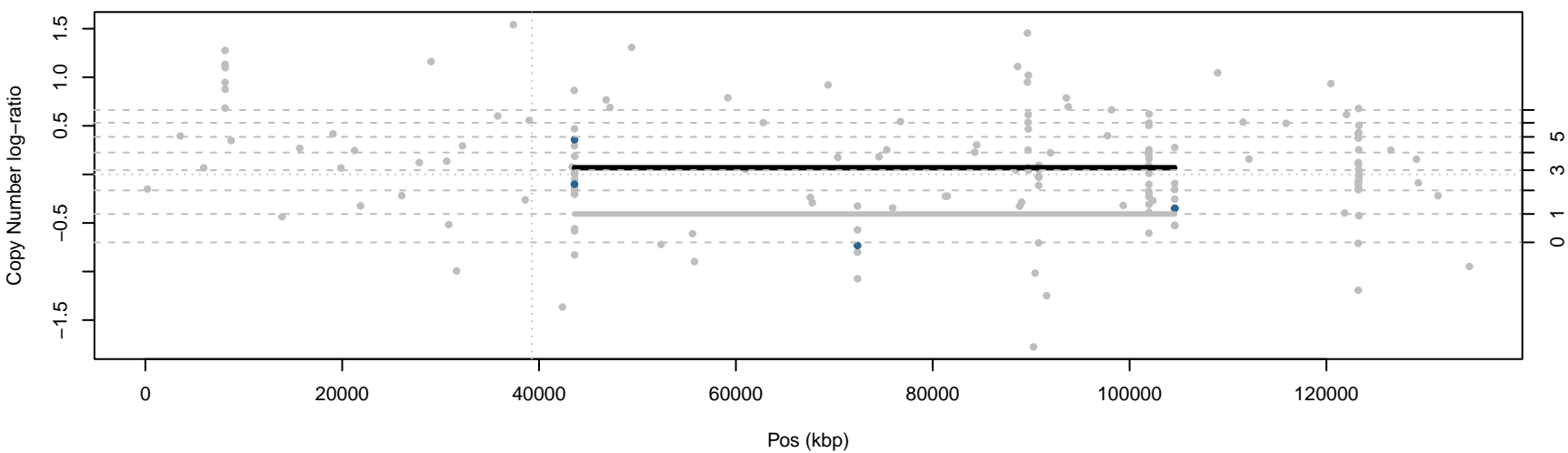
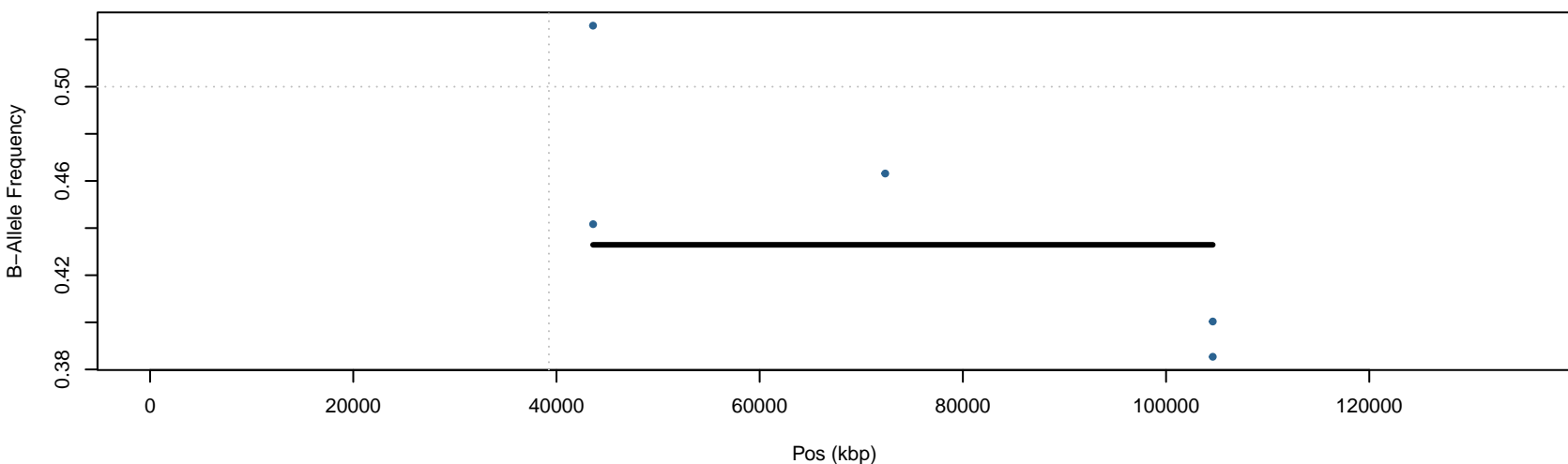




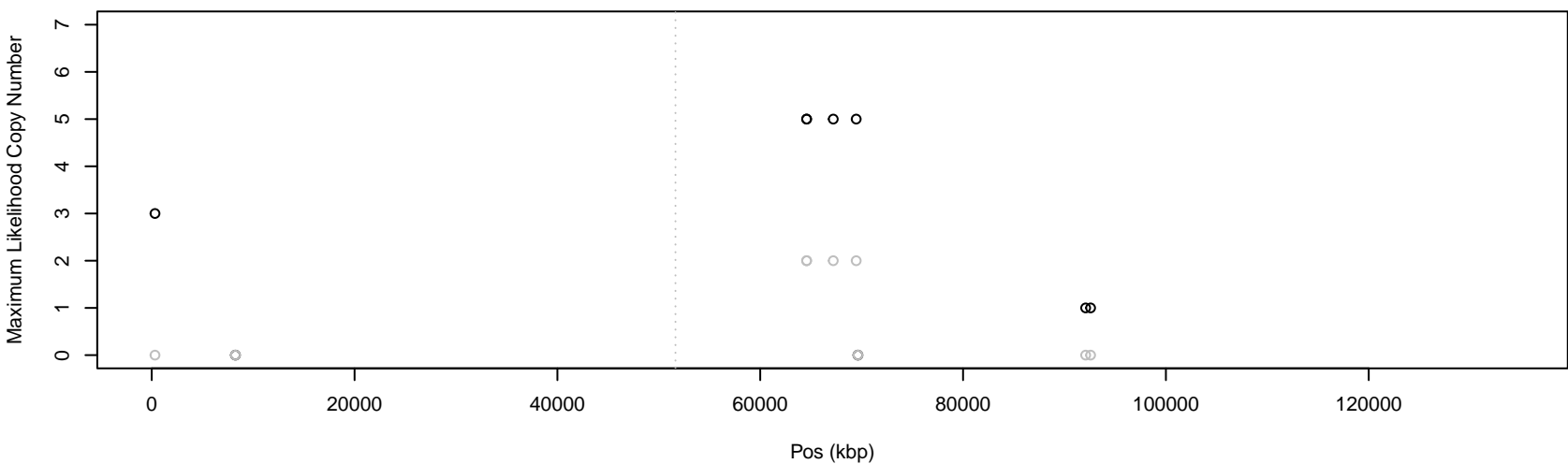
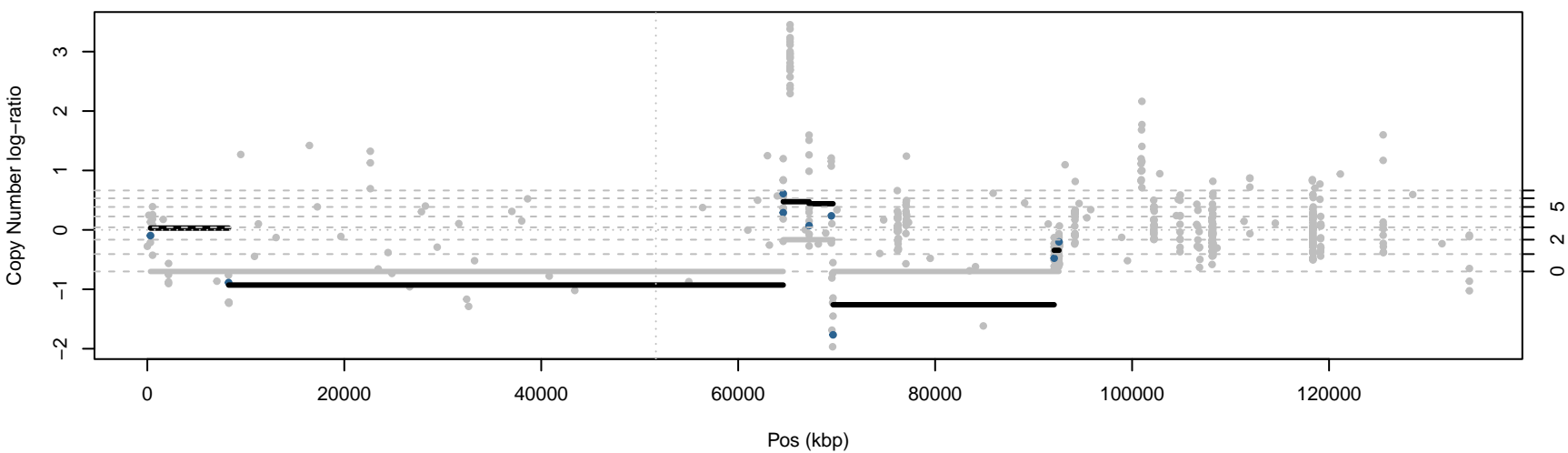
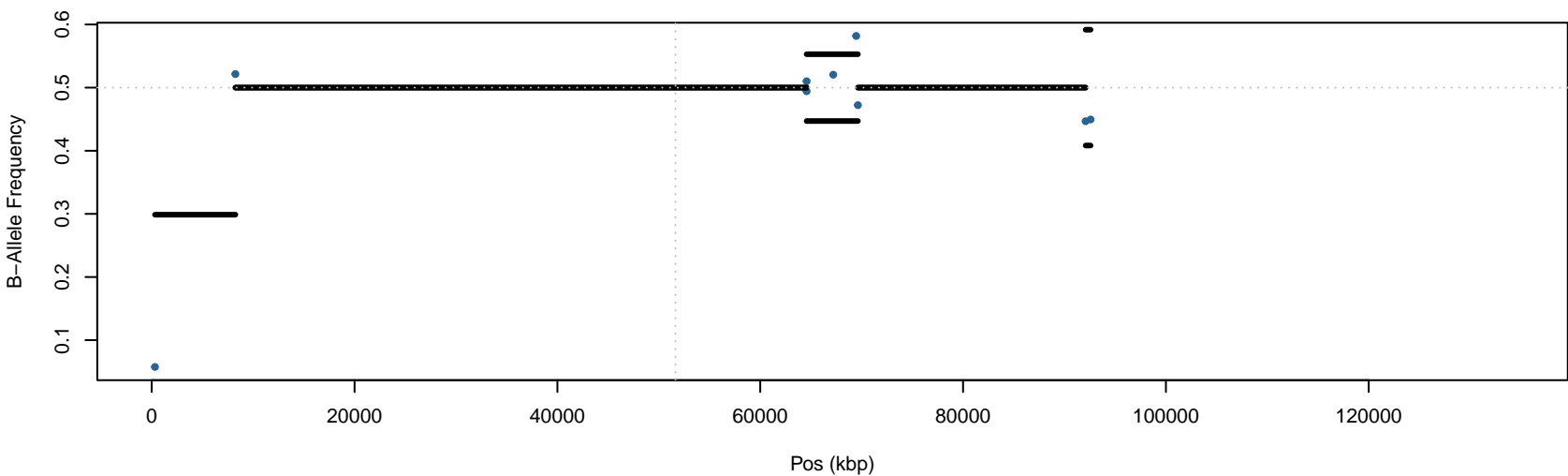




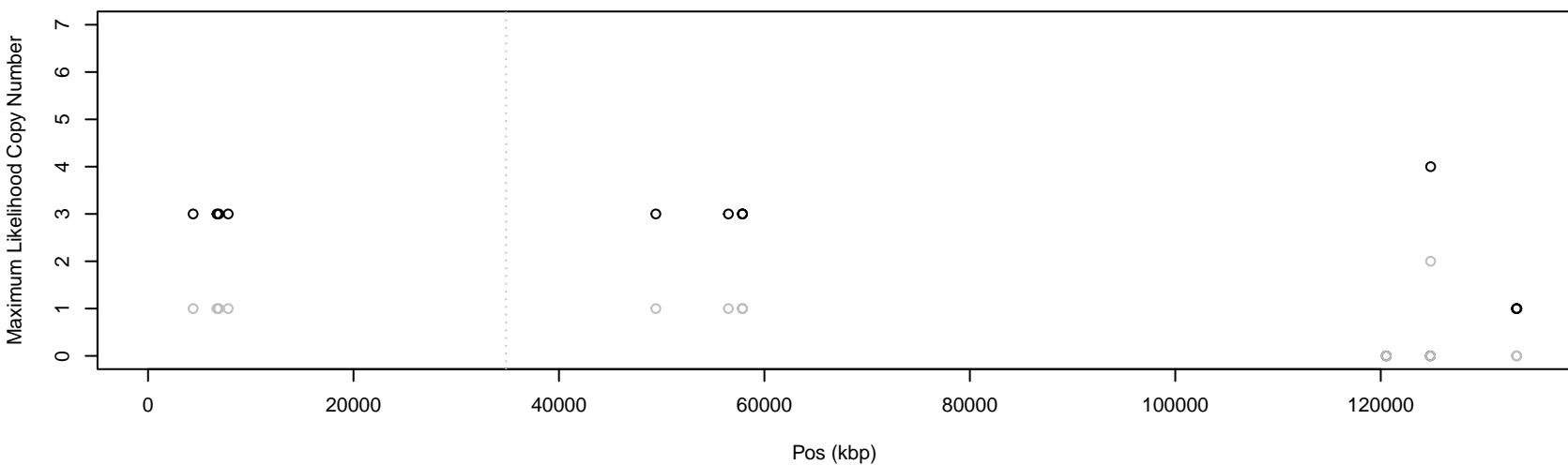
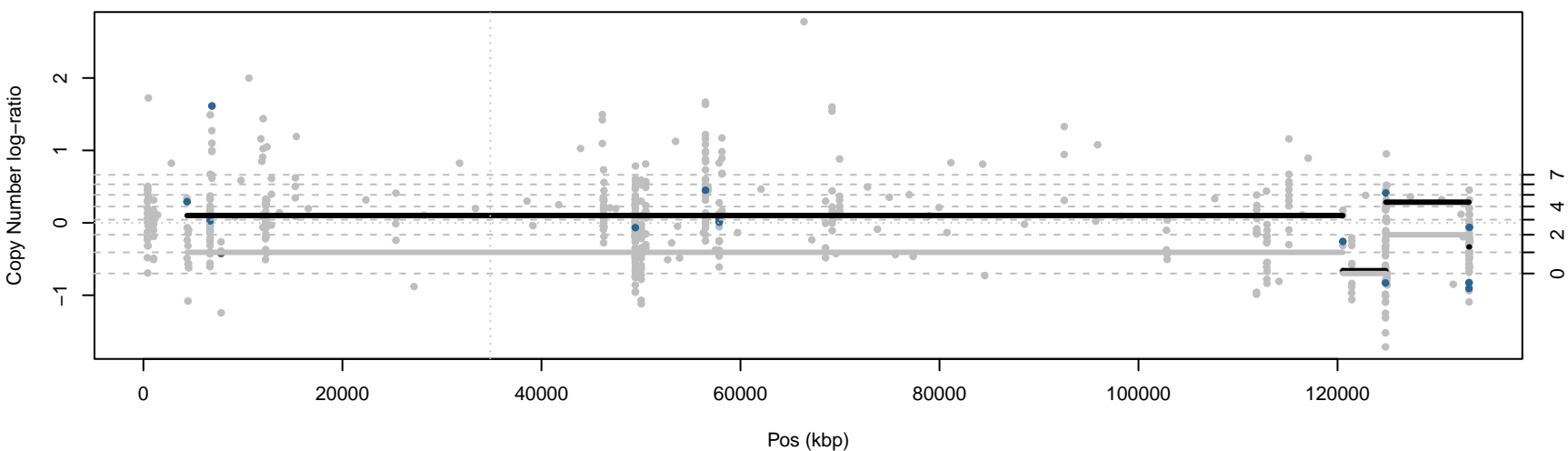
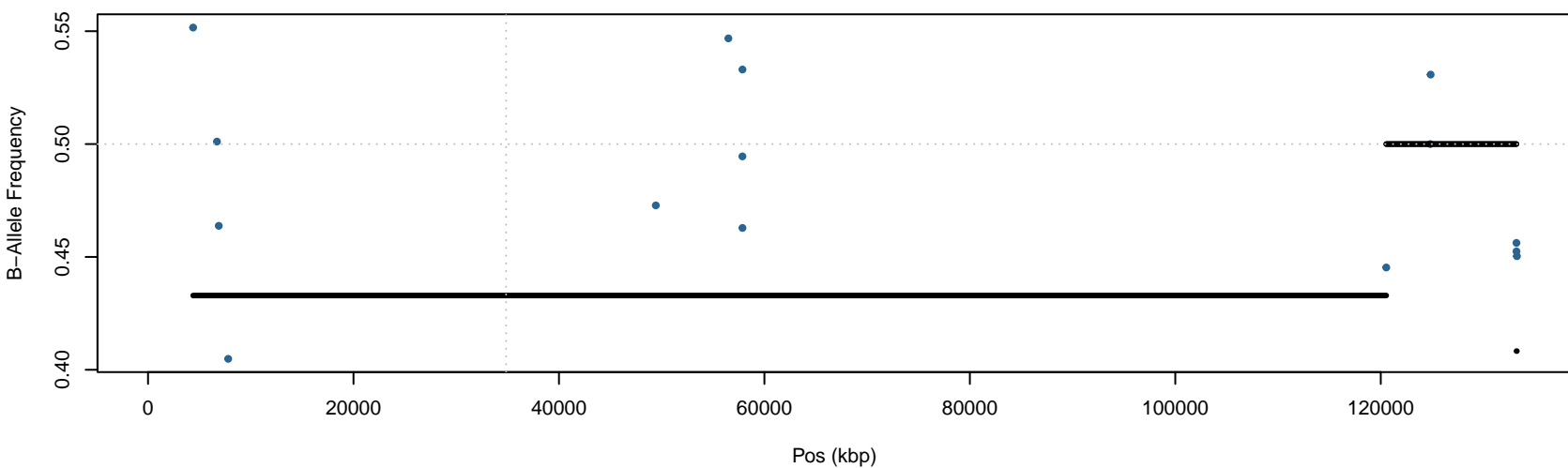
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 10



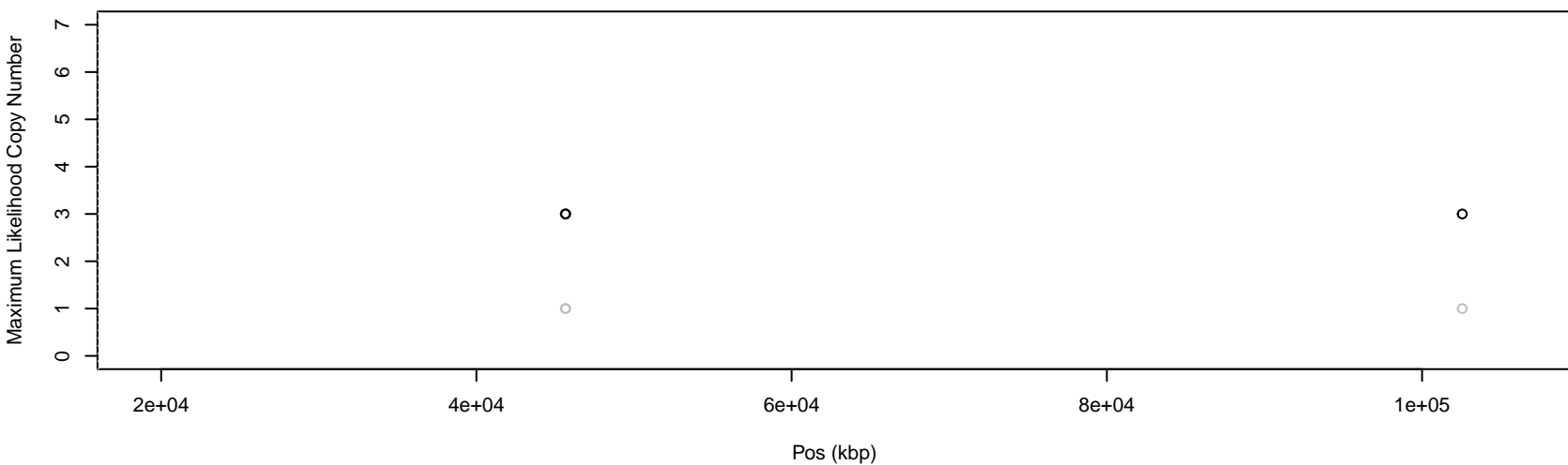
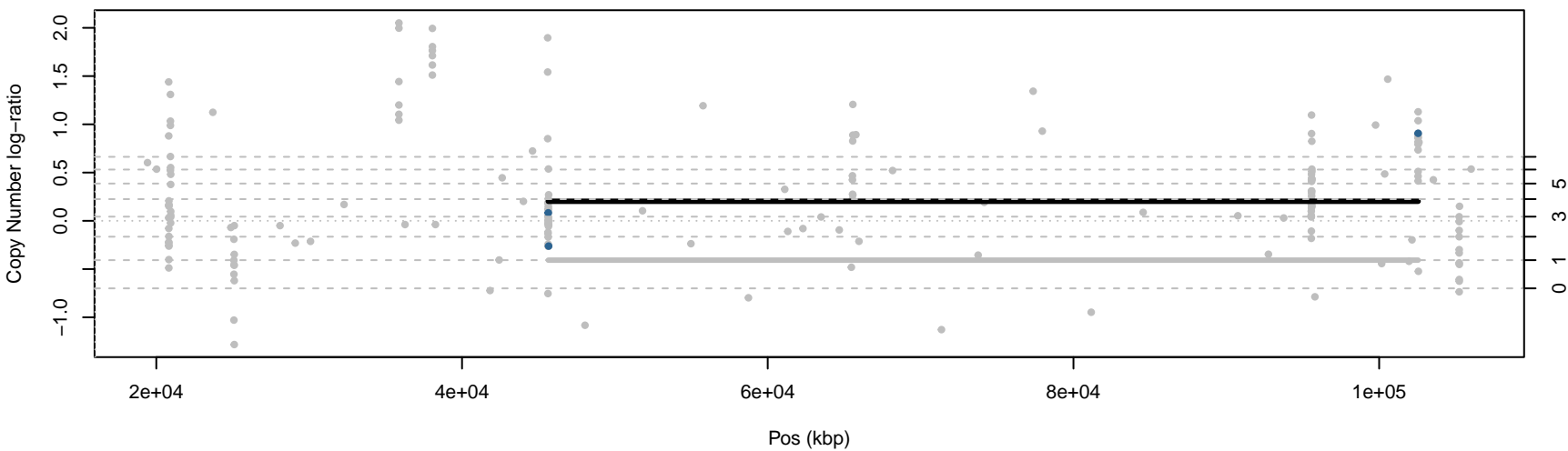
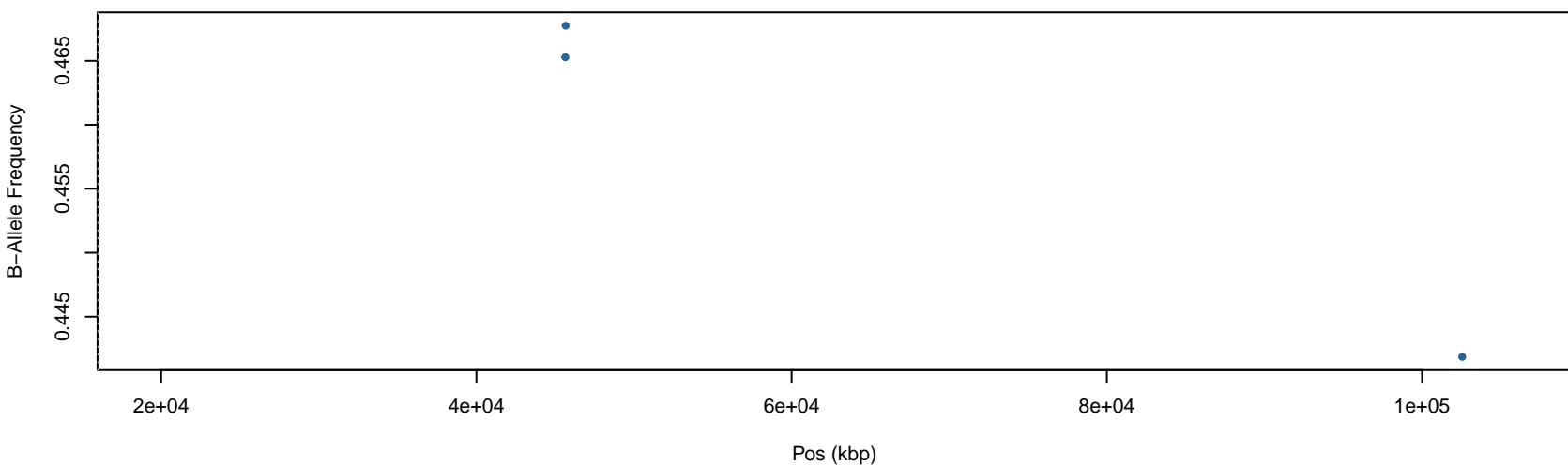
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 11



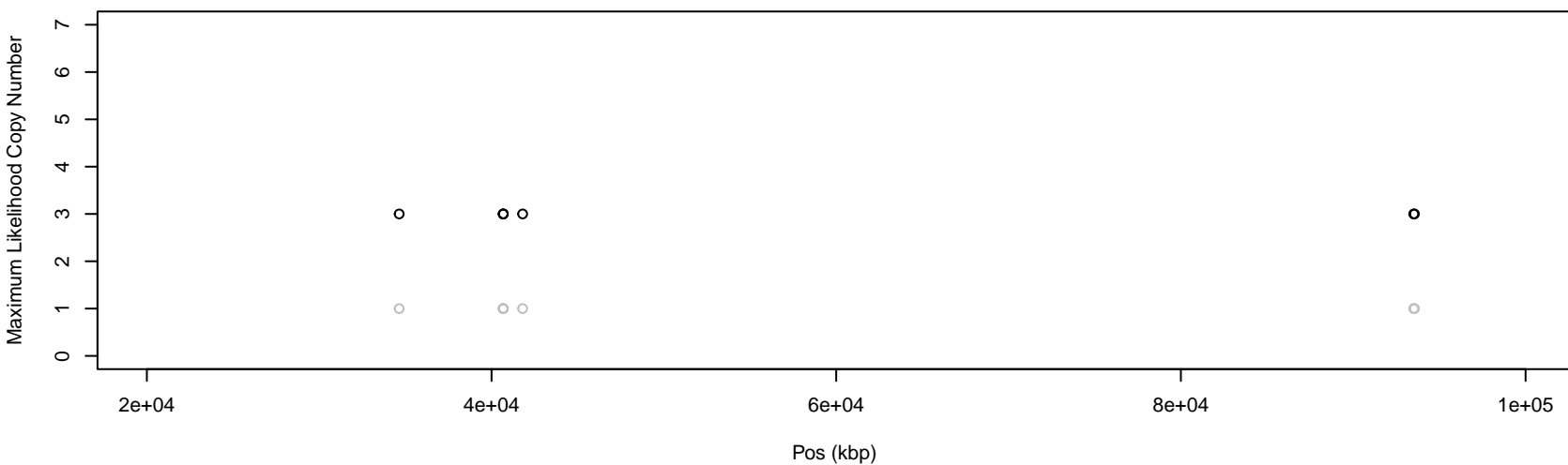
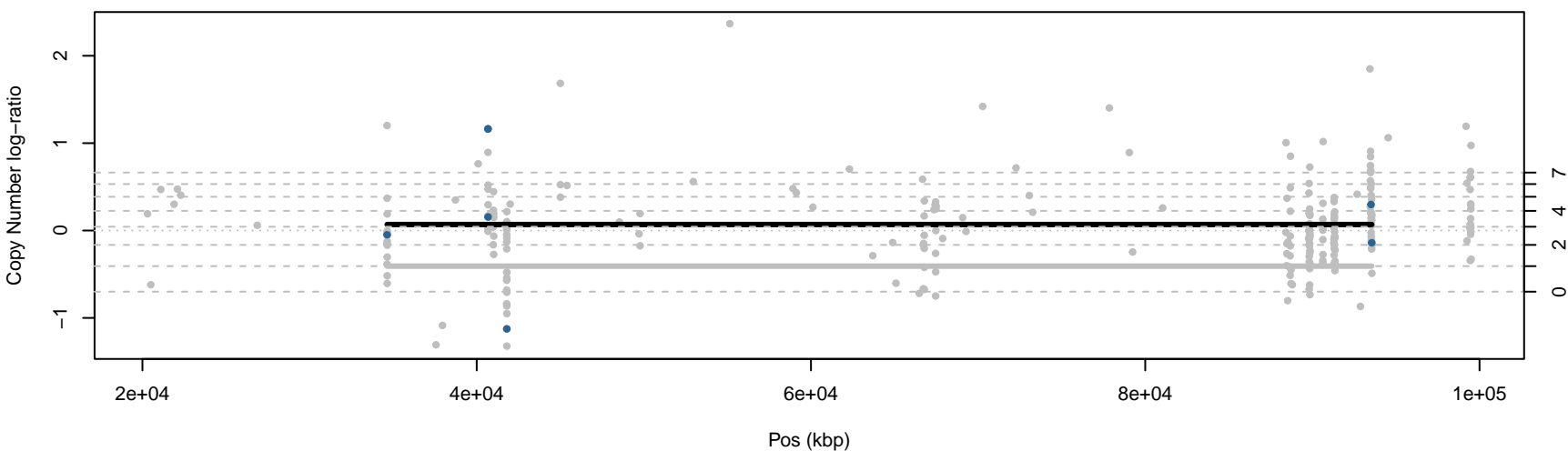
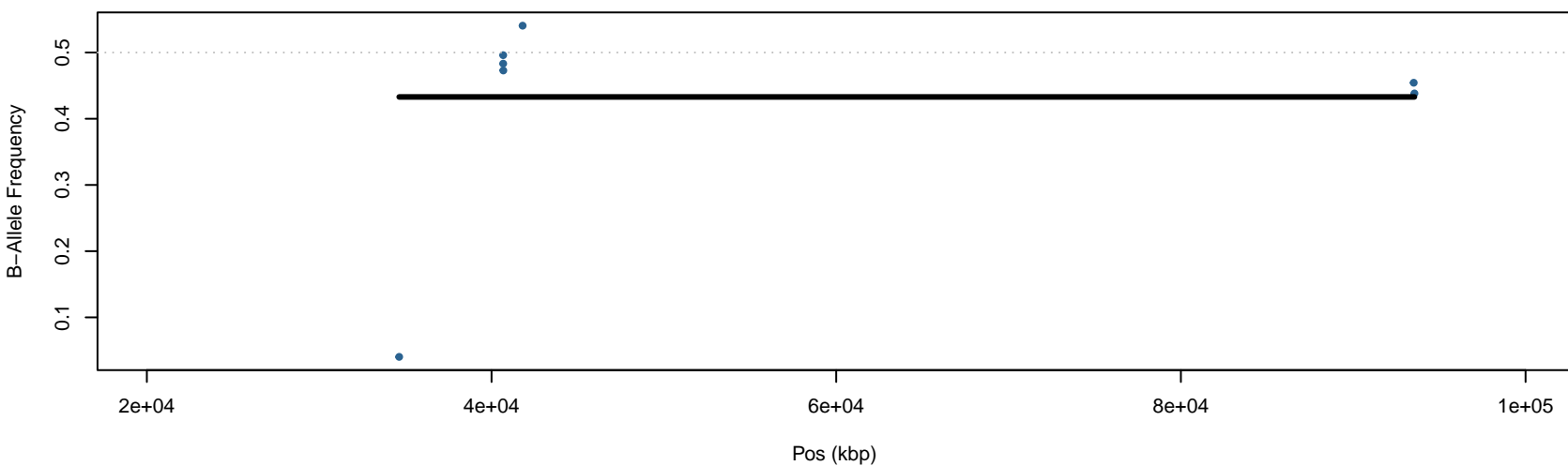
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 12



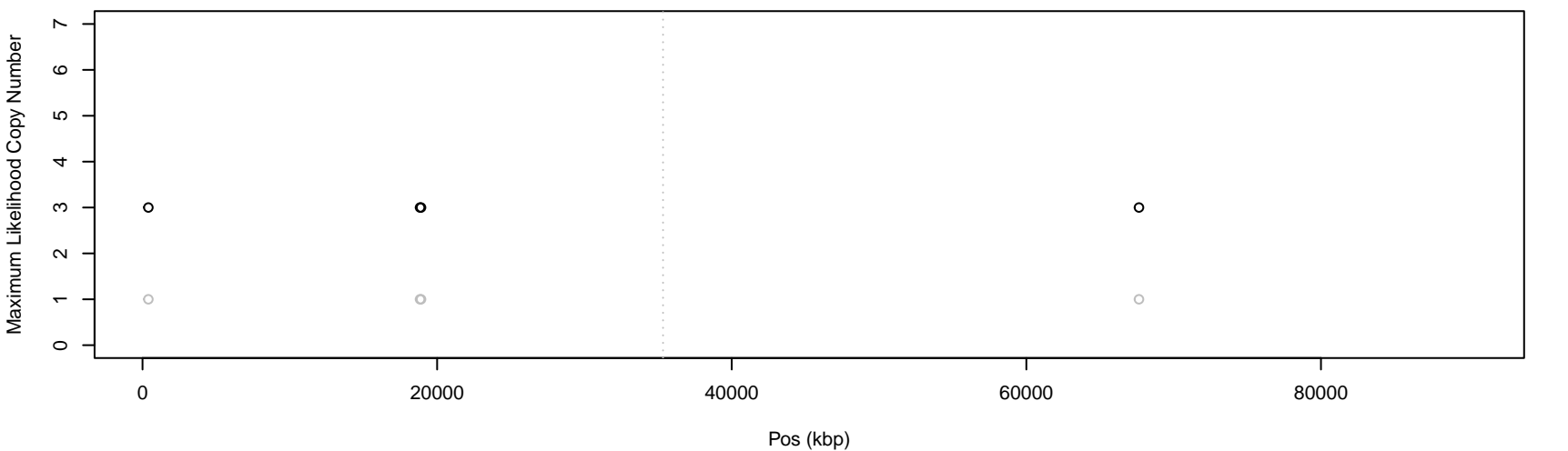
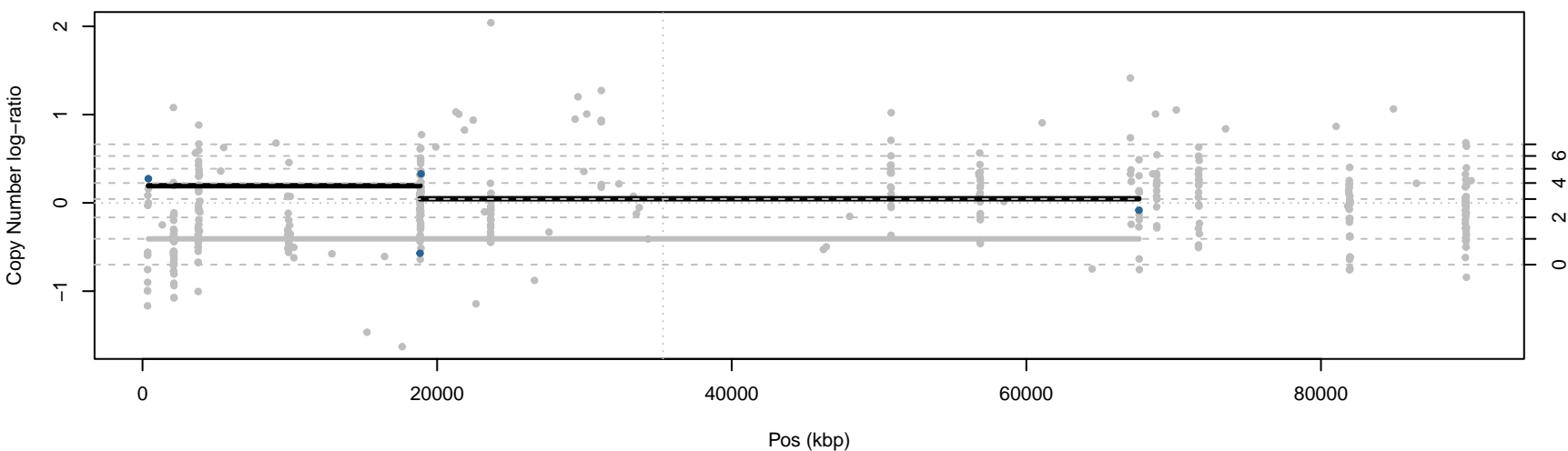
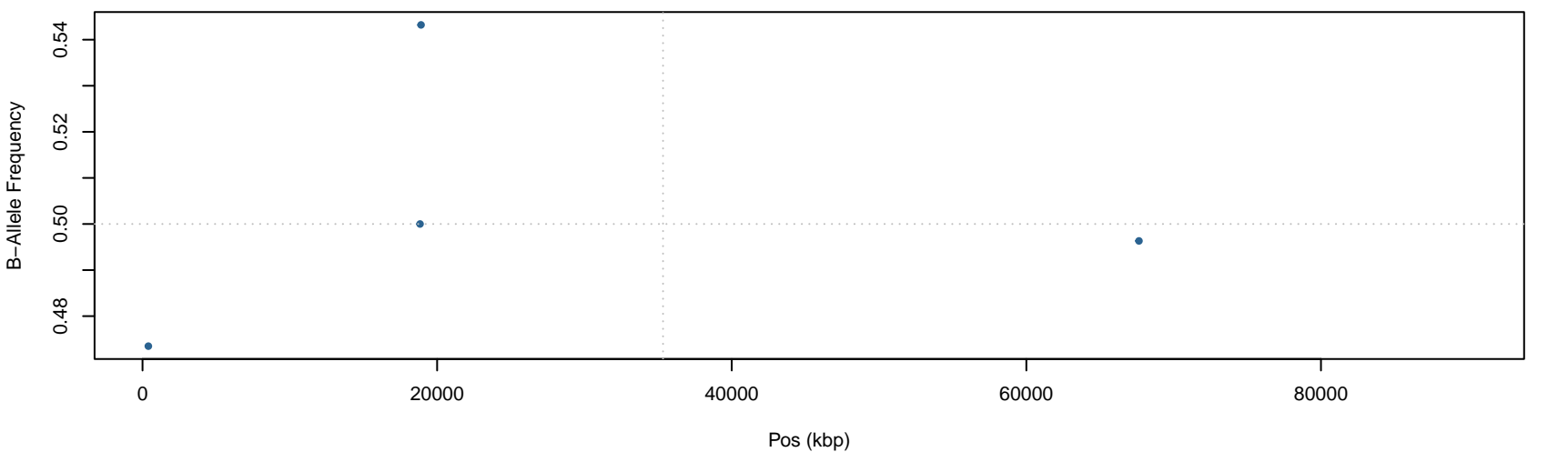
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 14



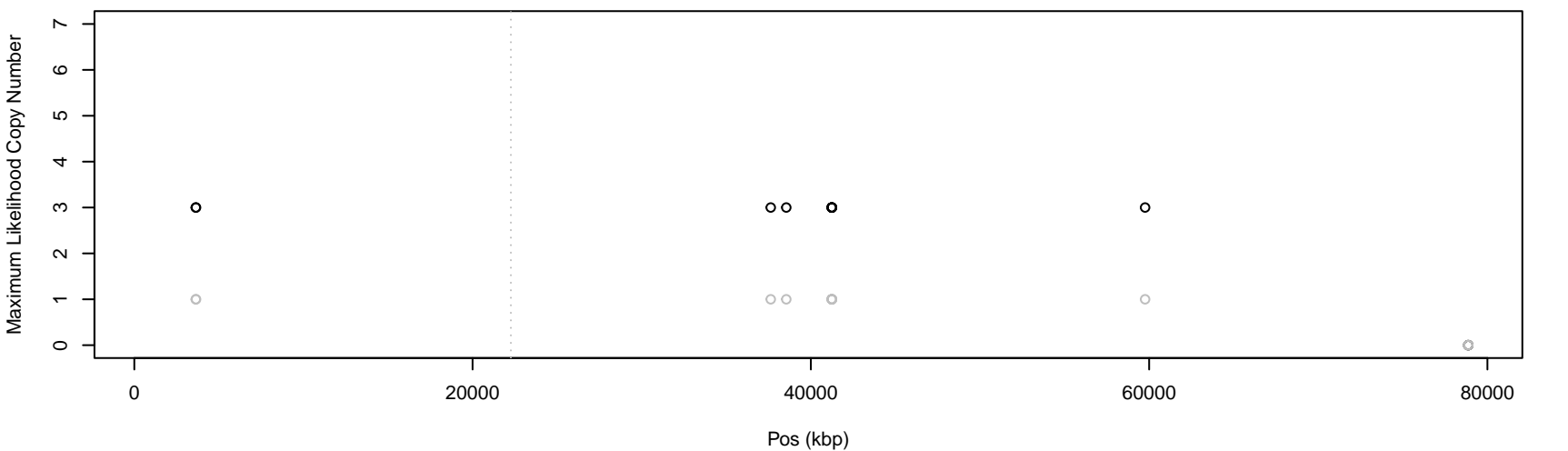
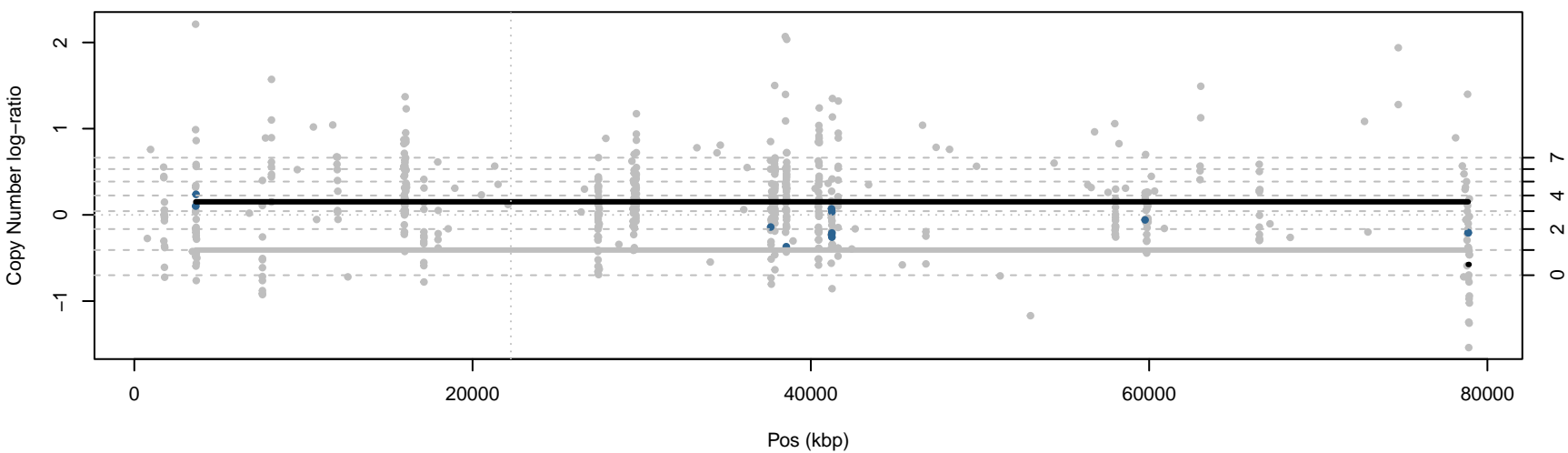
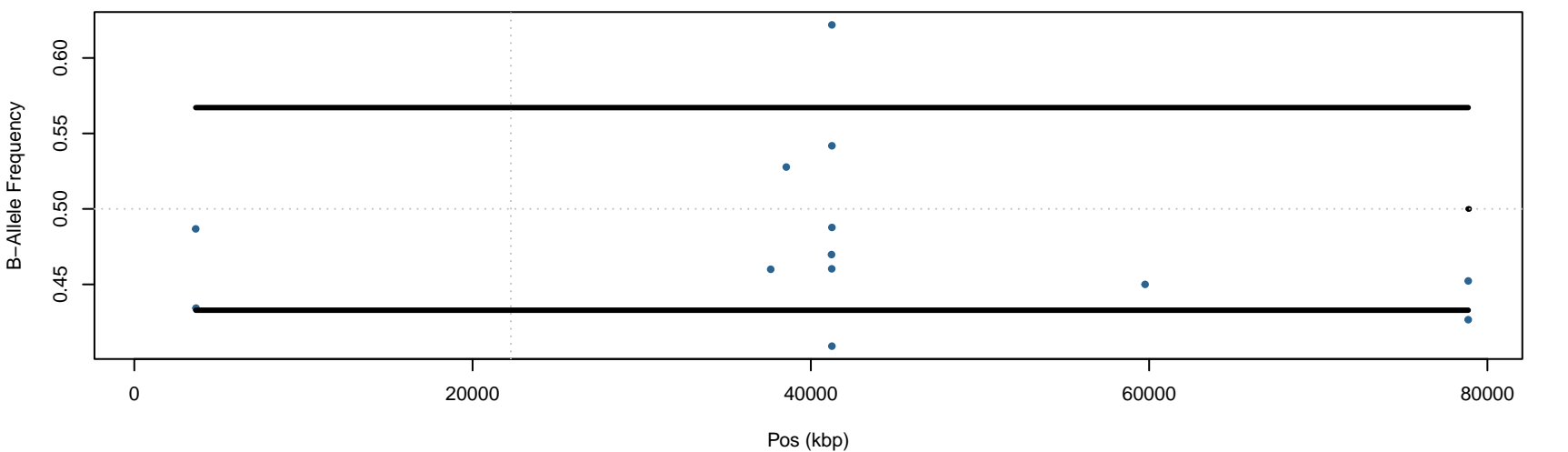
Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 15



Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 16

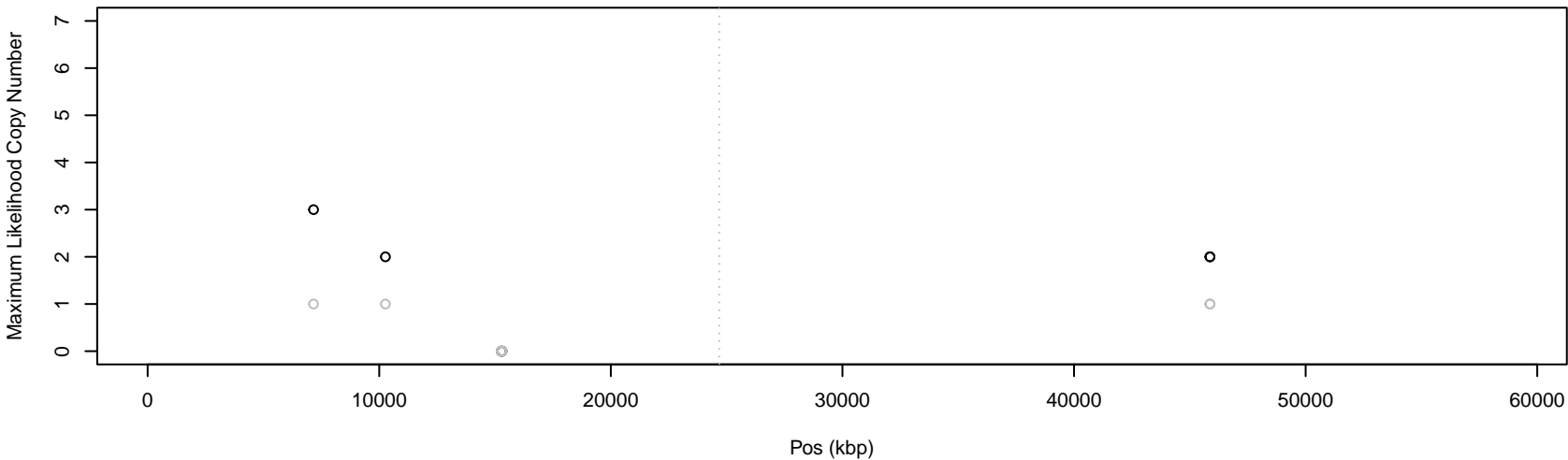
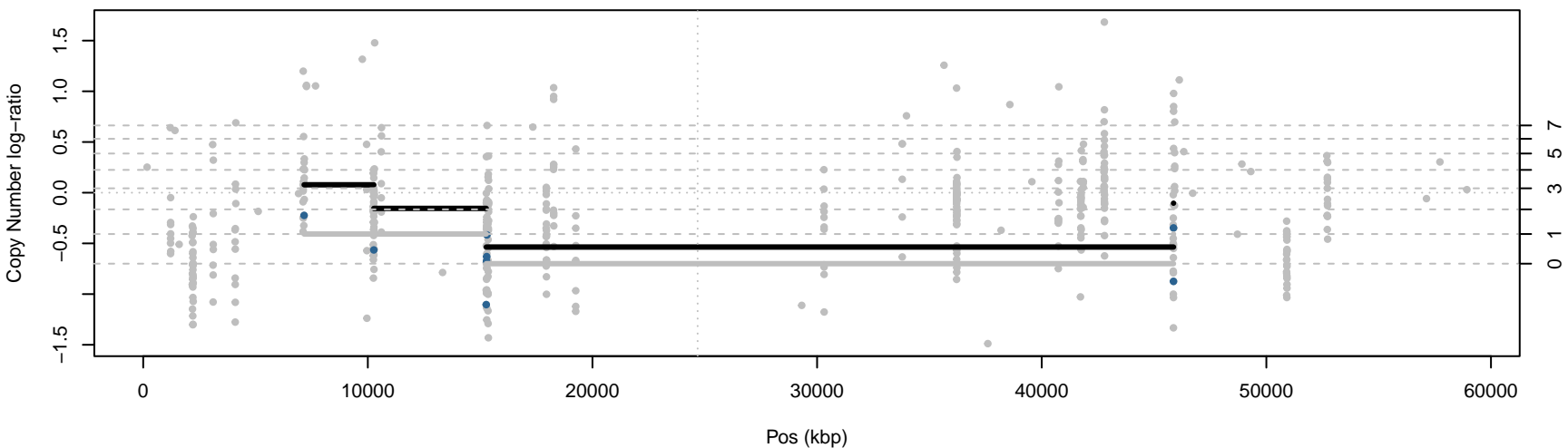
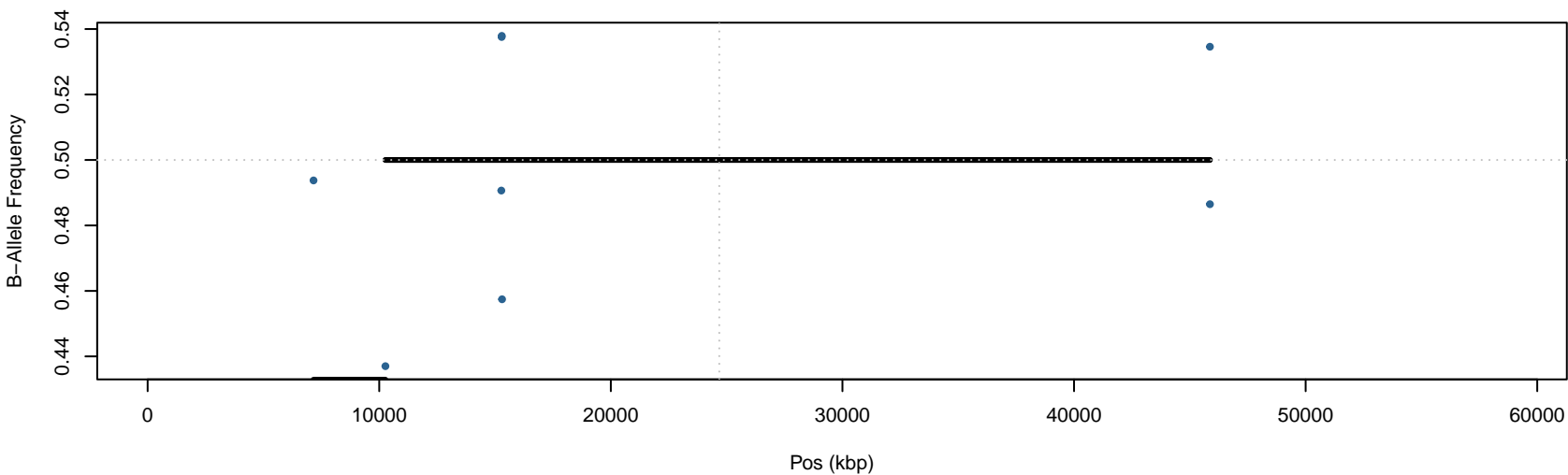


Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 17

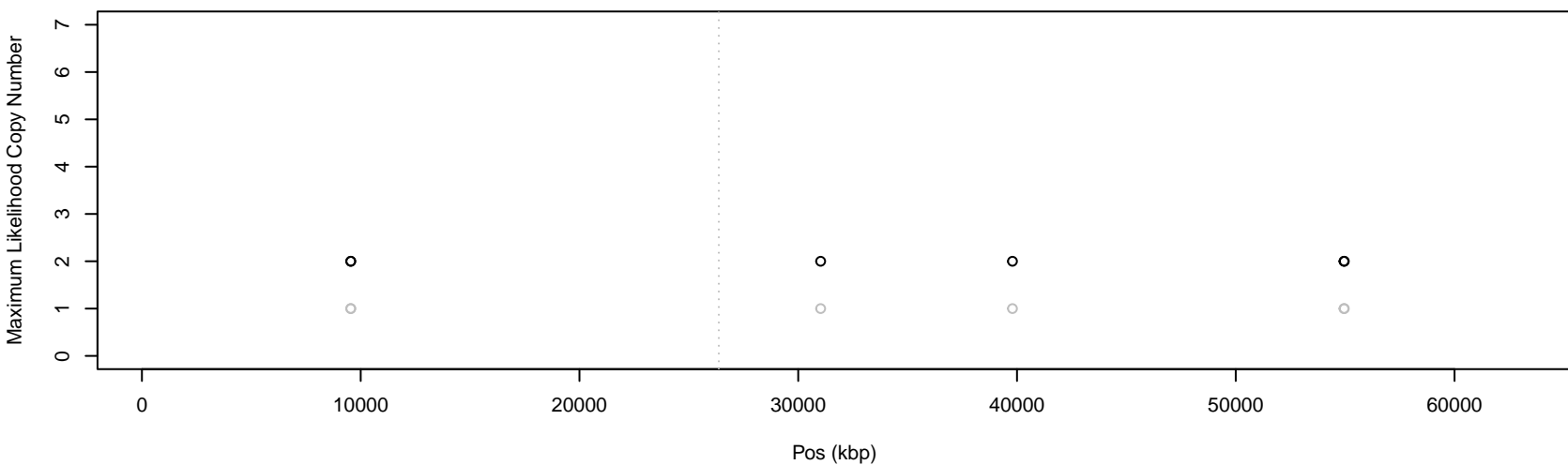
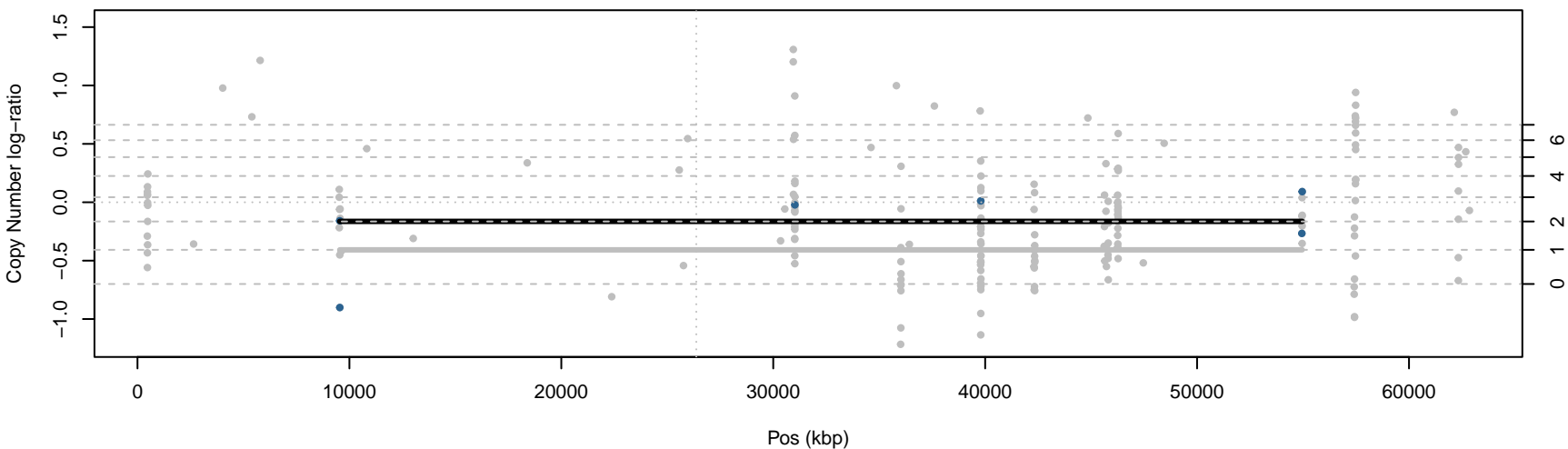
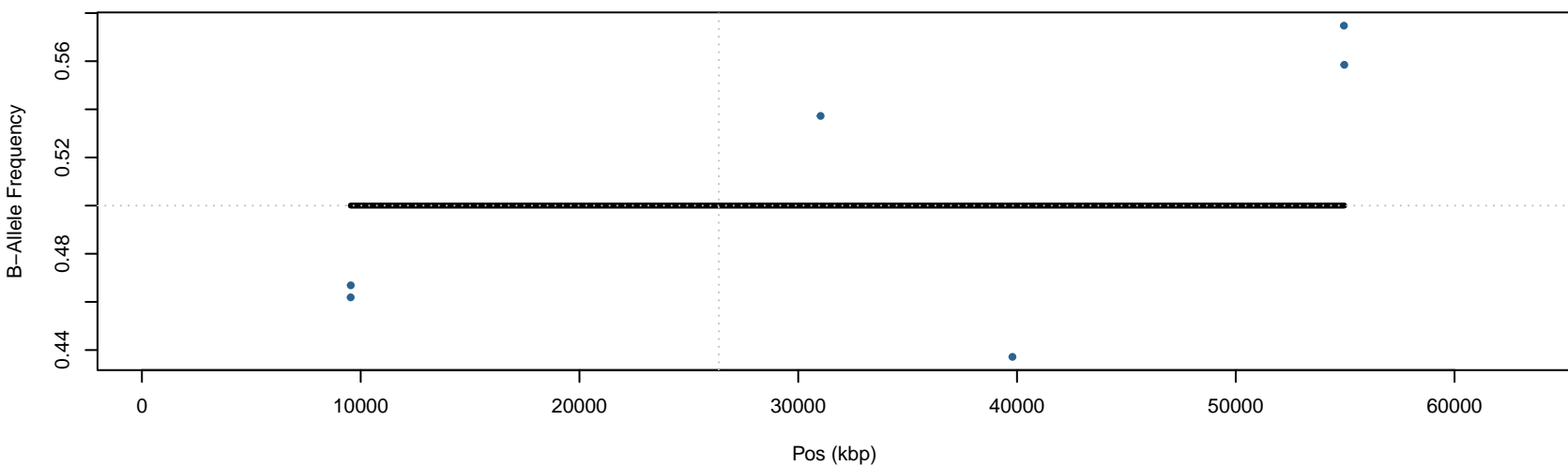




Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367,591 Chromosome: 19



Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367;591 Chromosome: 20



Purity: 0.31 Tumor ploidy: 2.78 SNV log-likelihood: 8.22 GoF: 93.8% Mean coverage: 367,591 Chromosome: 22

