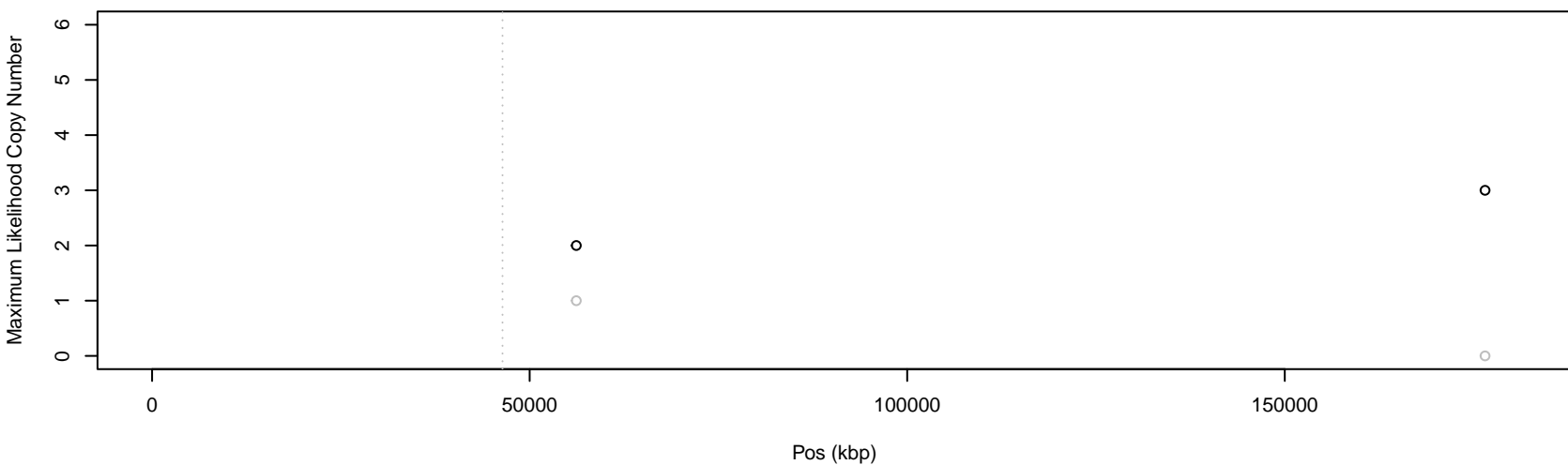
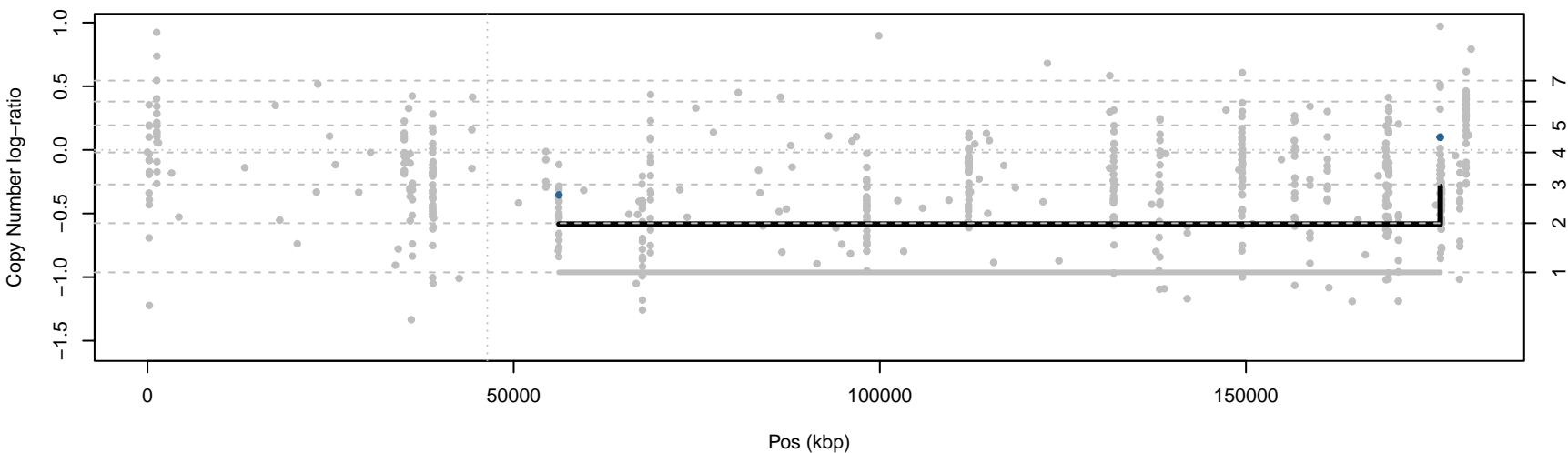
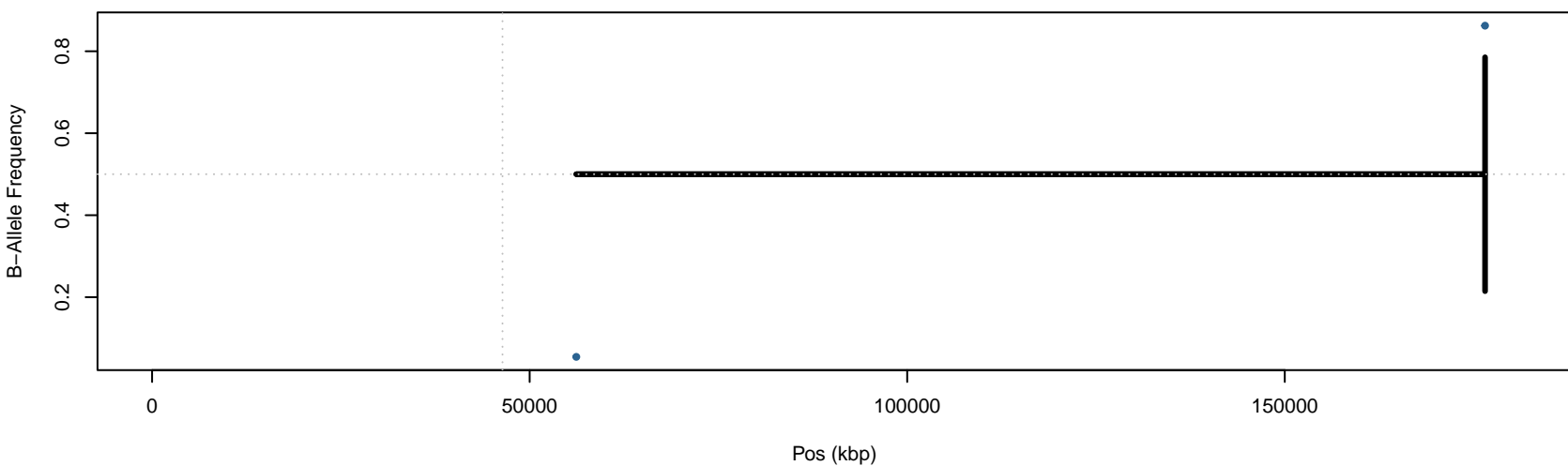
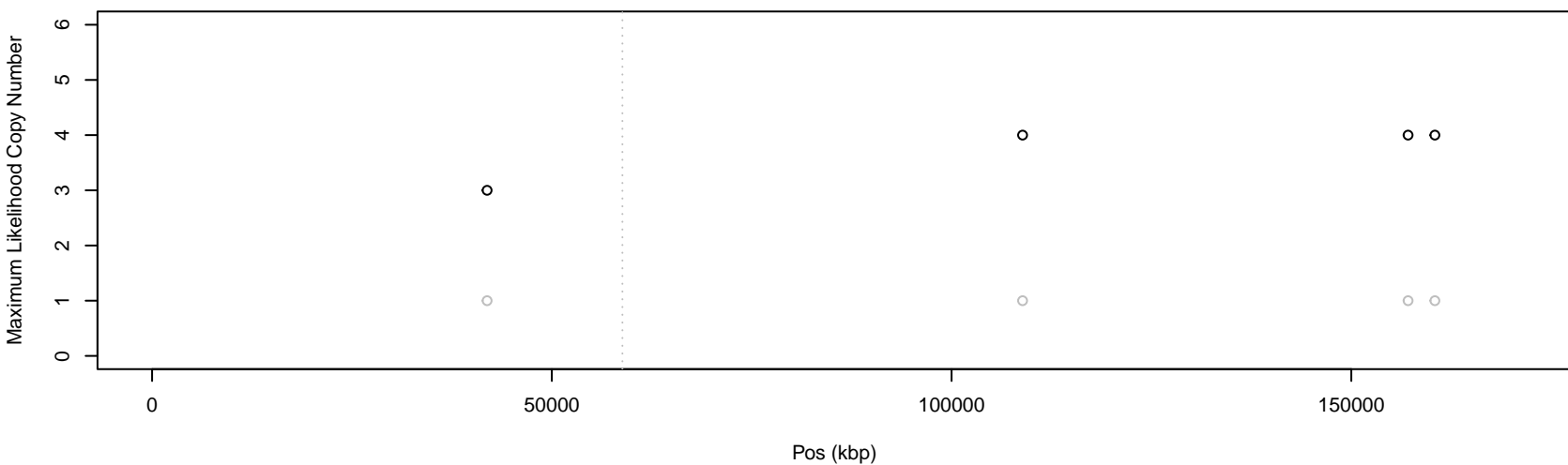
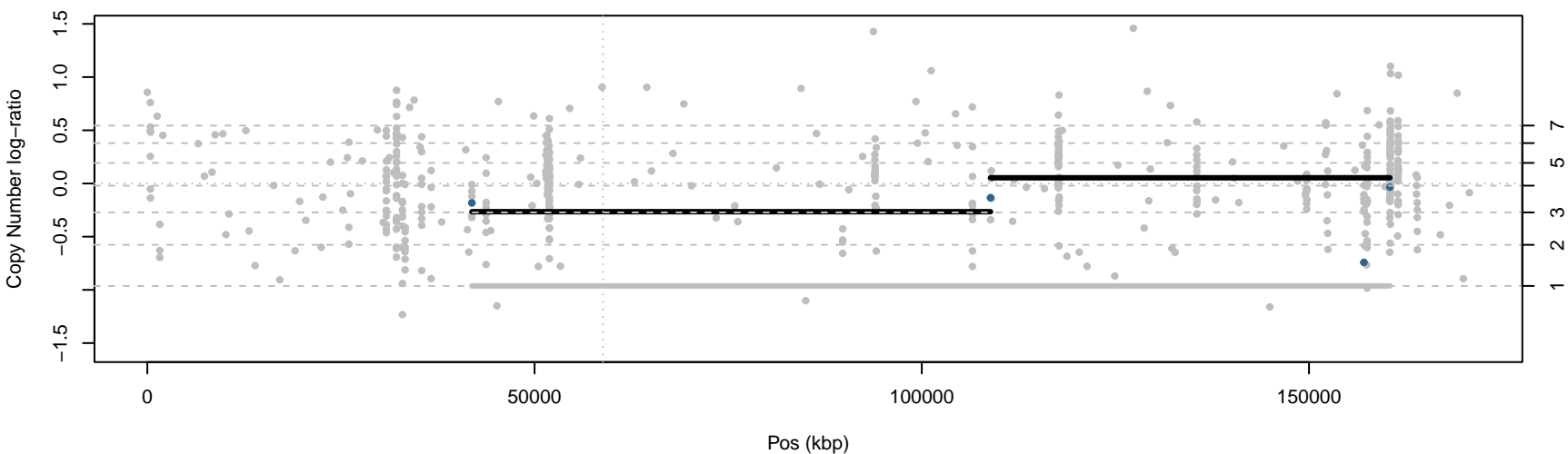
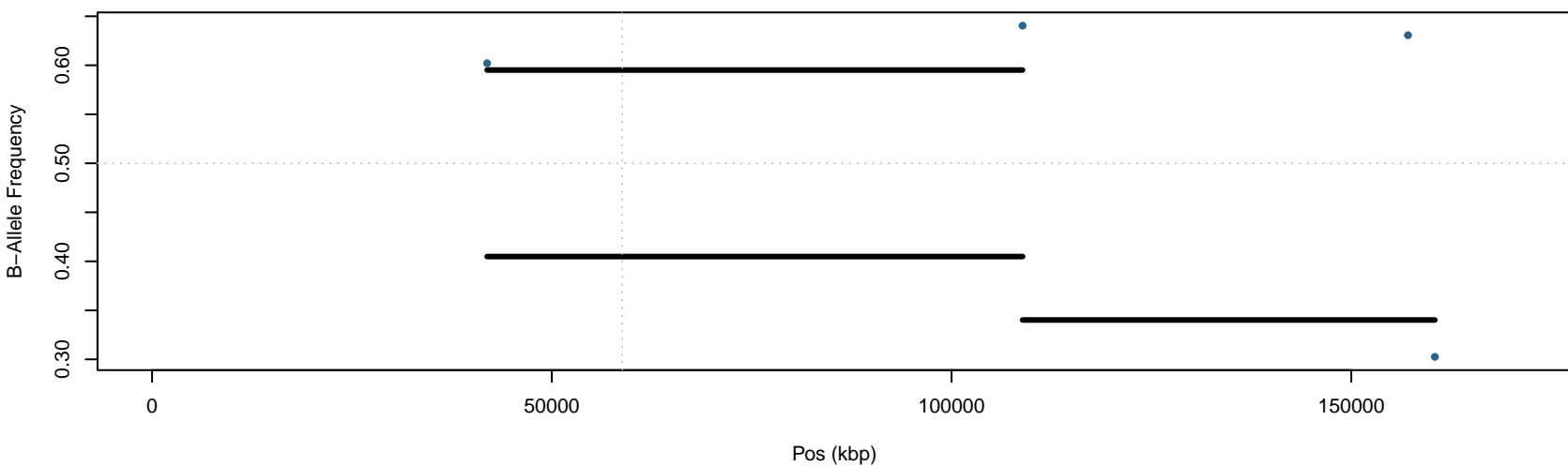


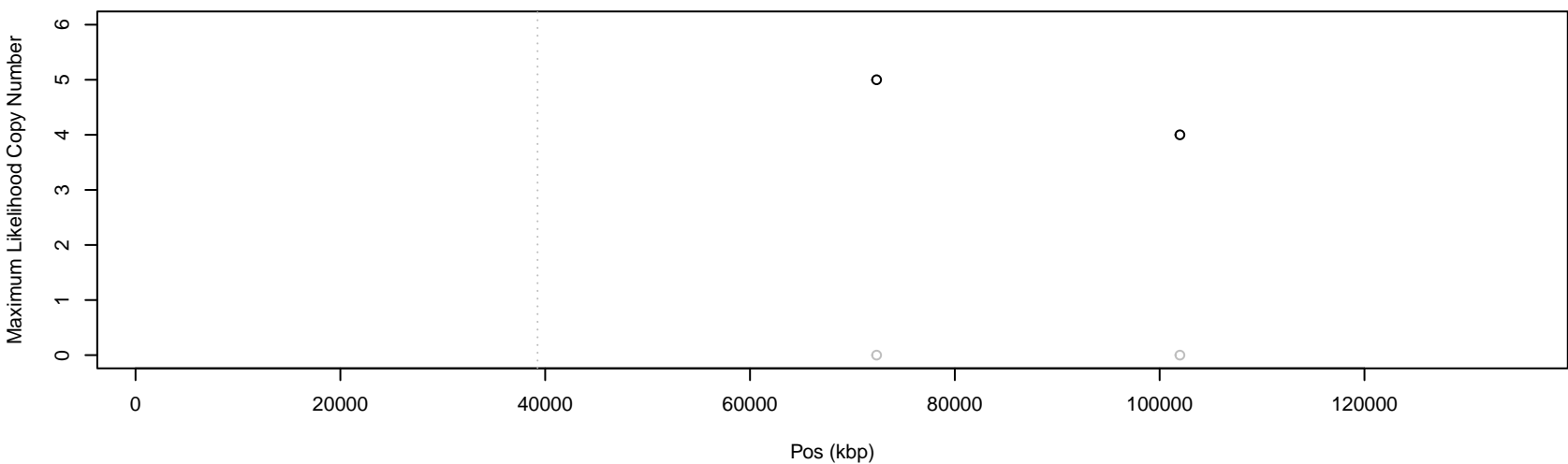
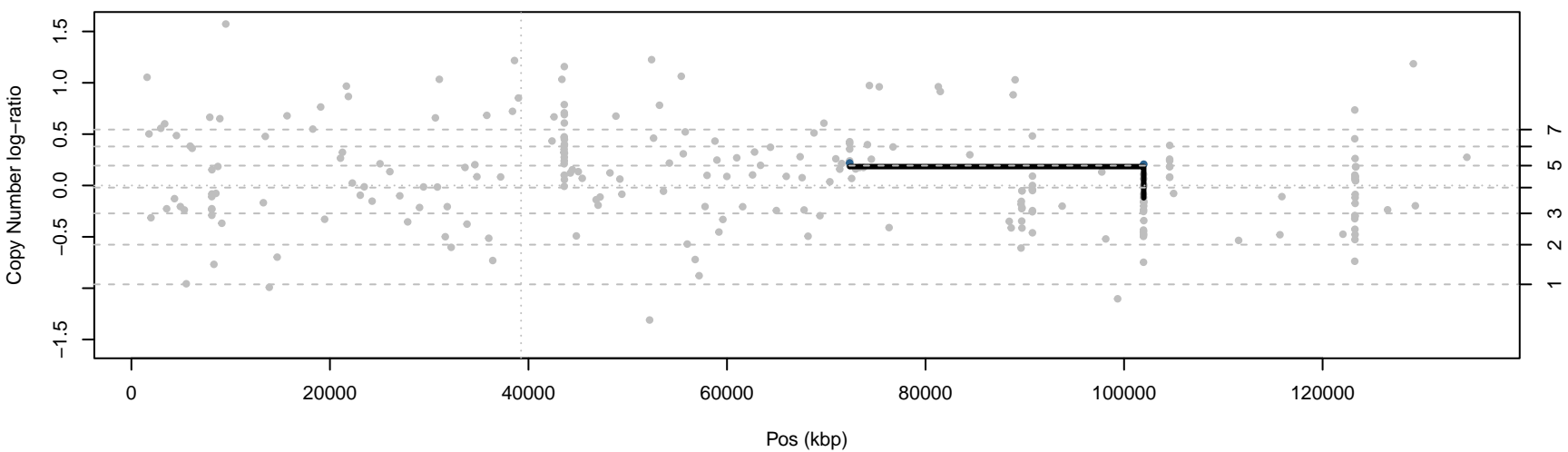
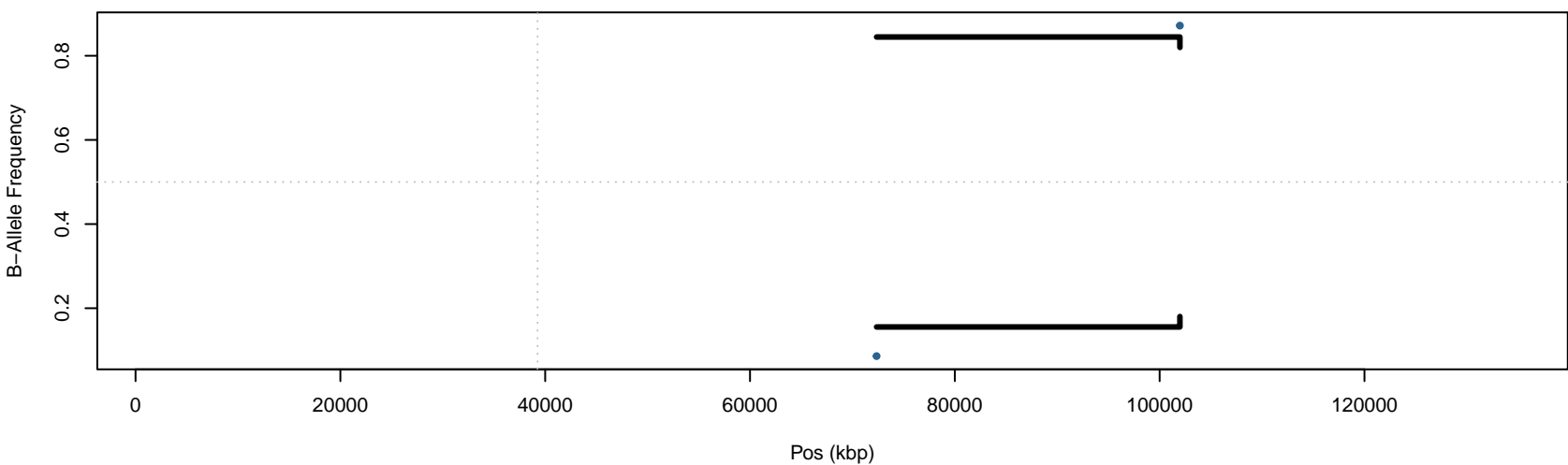
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 5



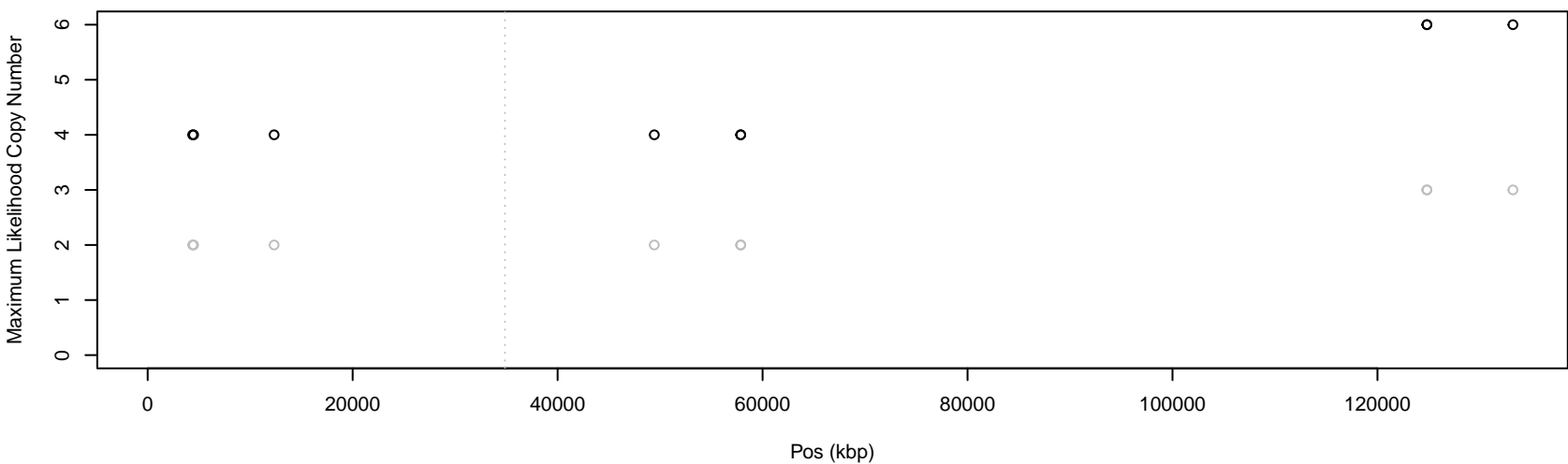
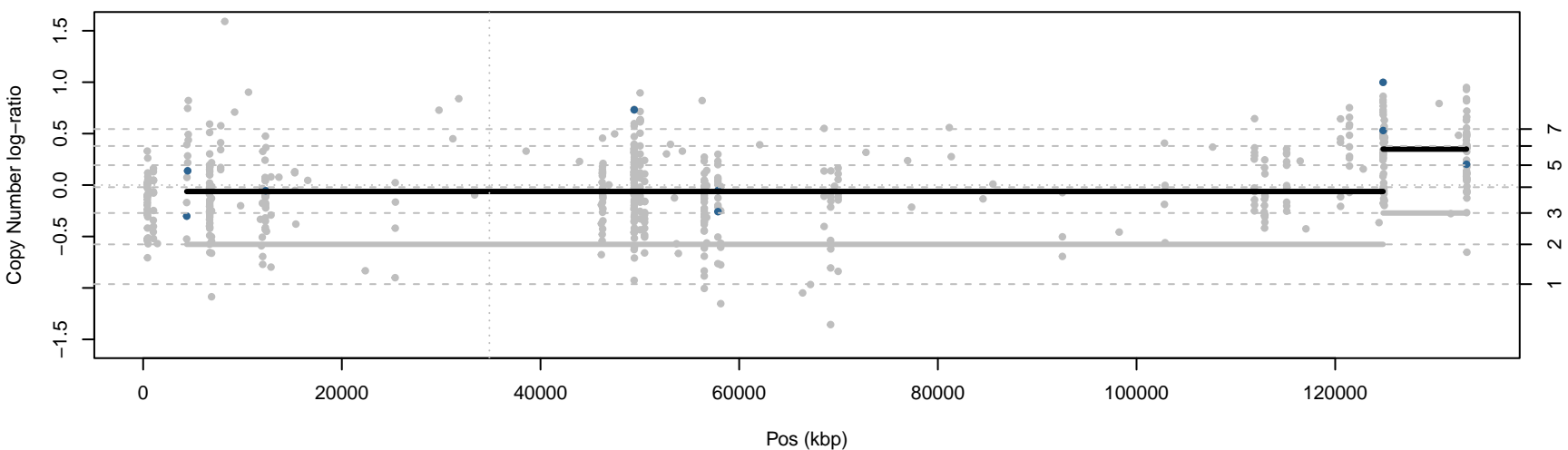
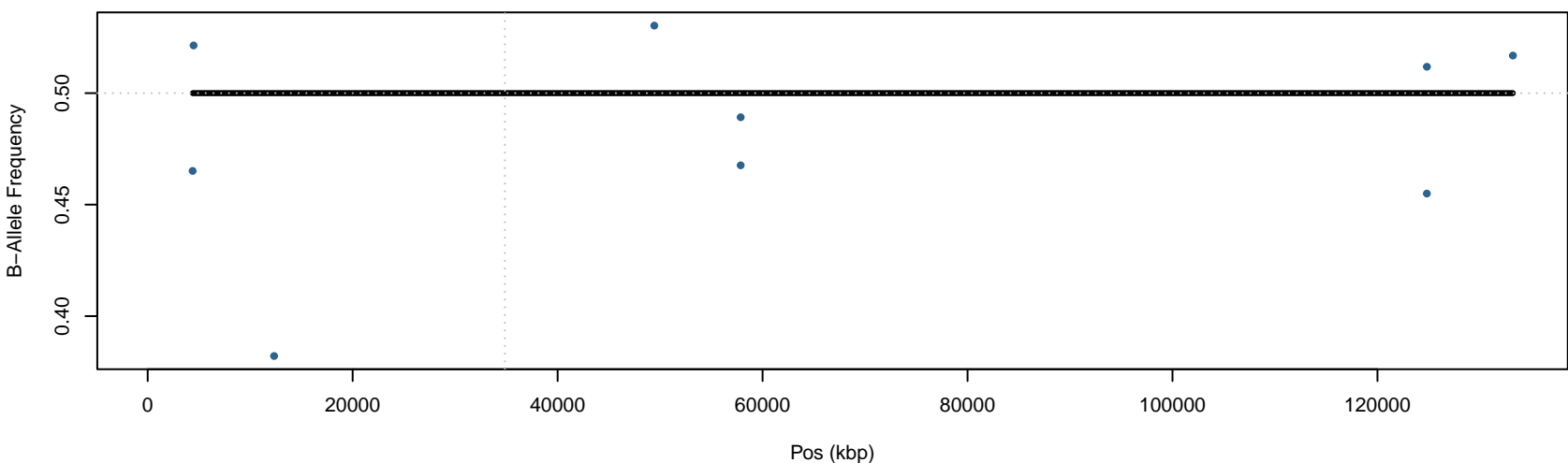
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 6



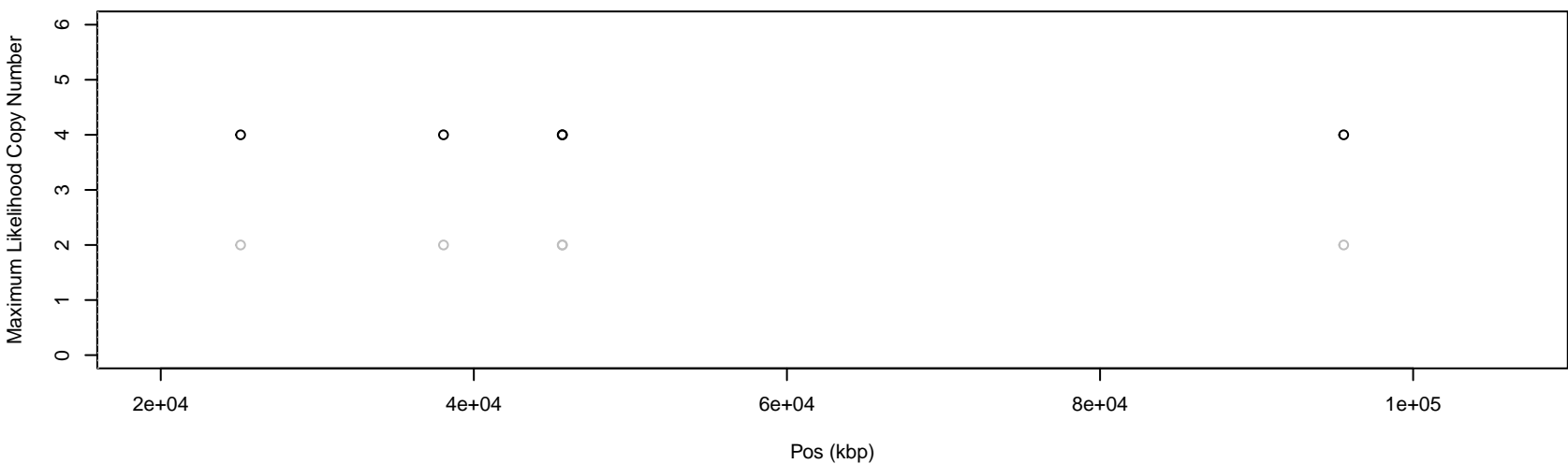
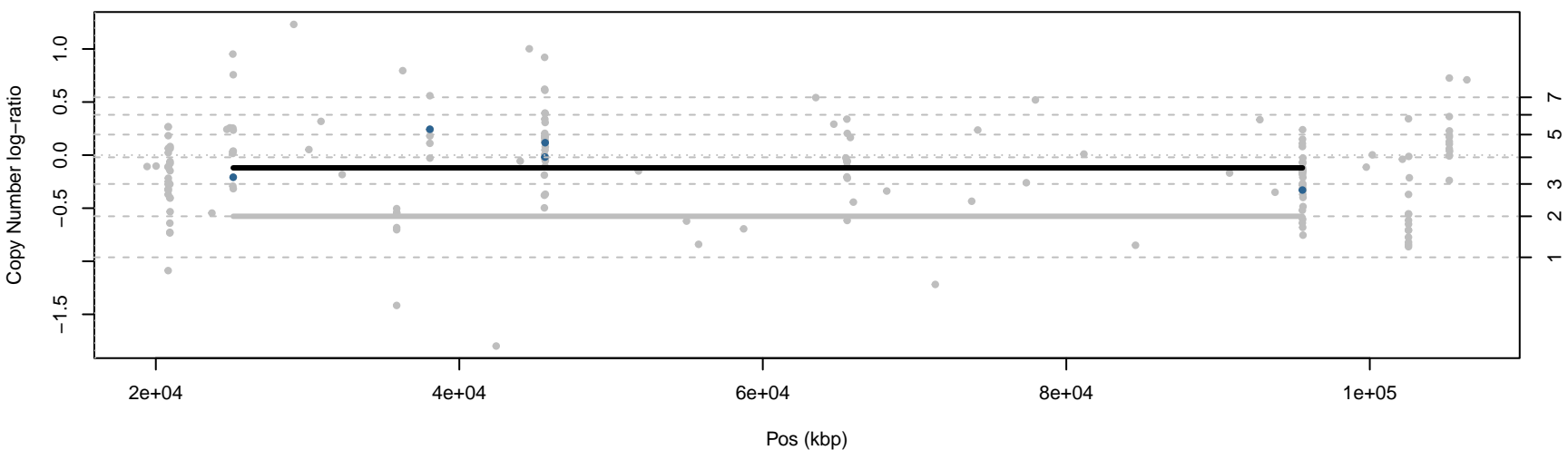
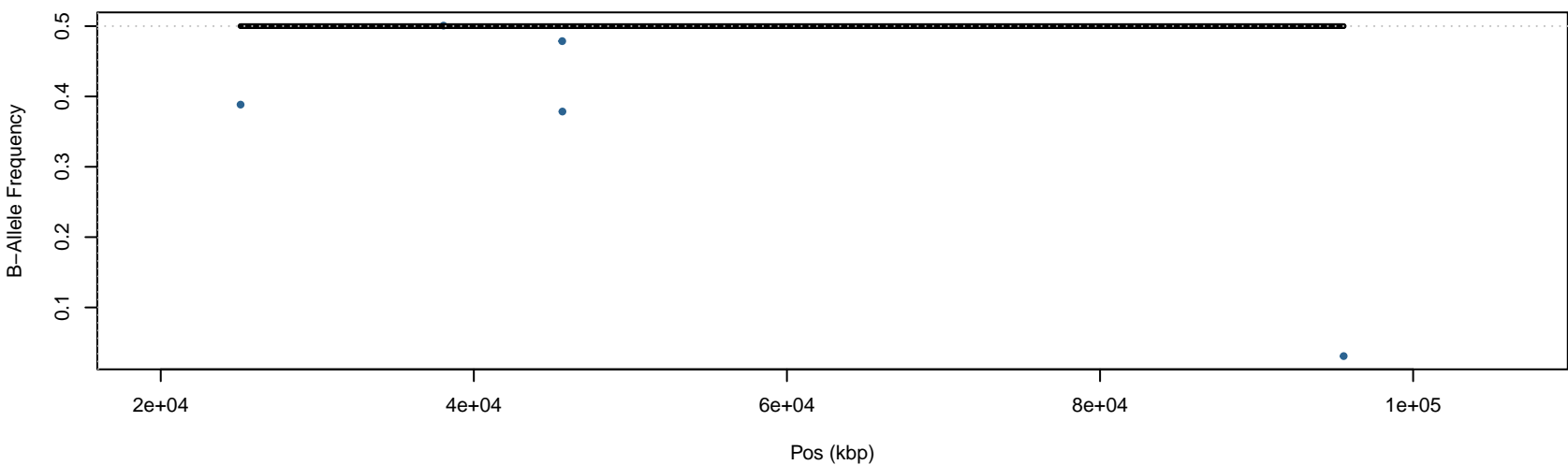
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 10



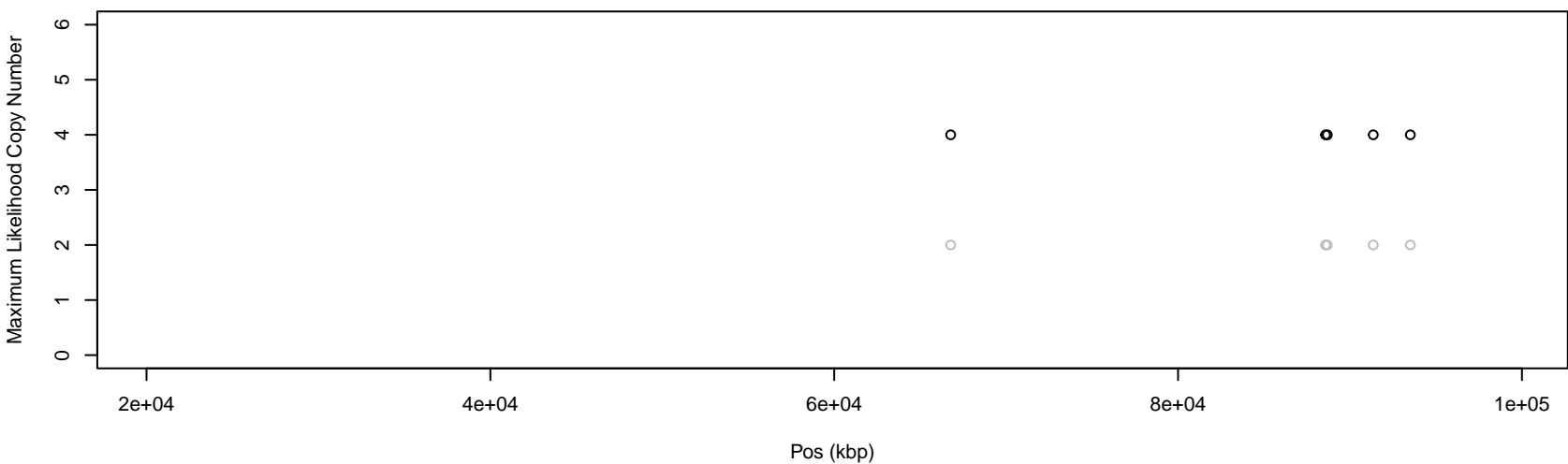
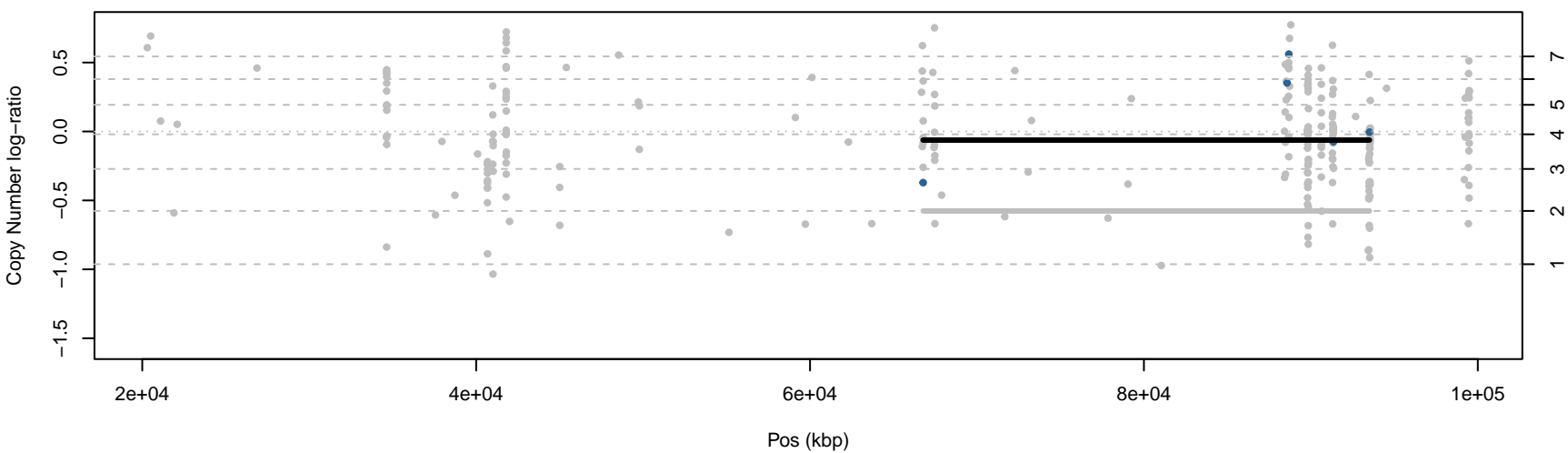
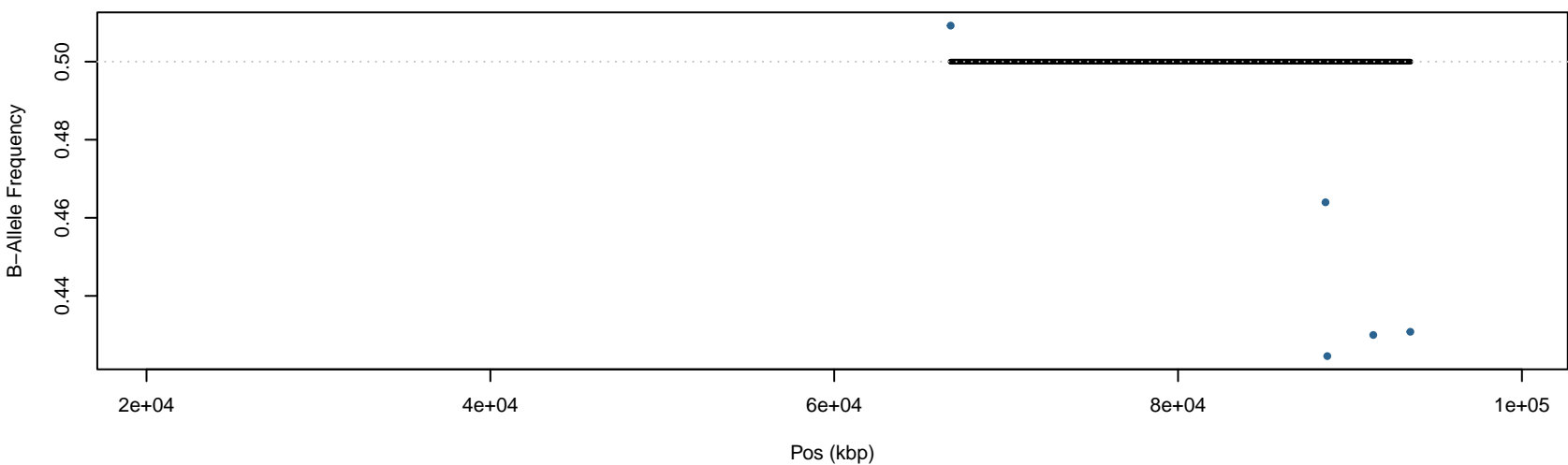
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 12



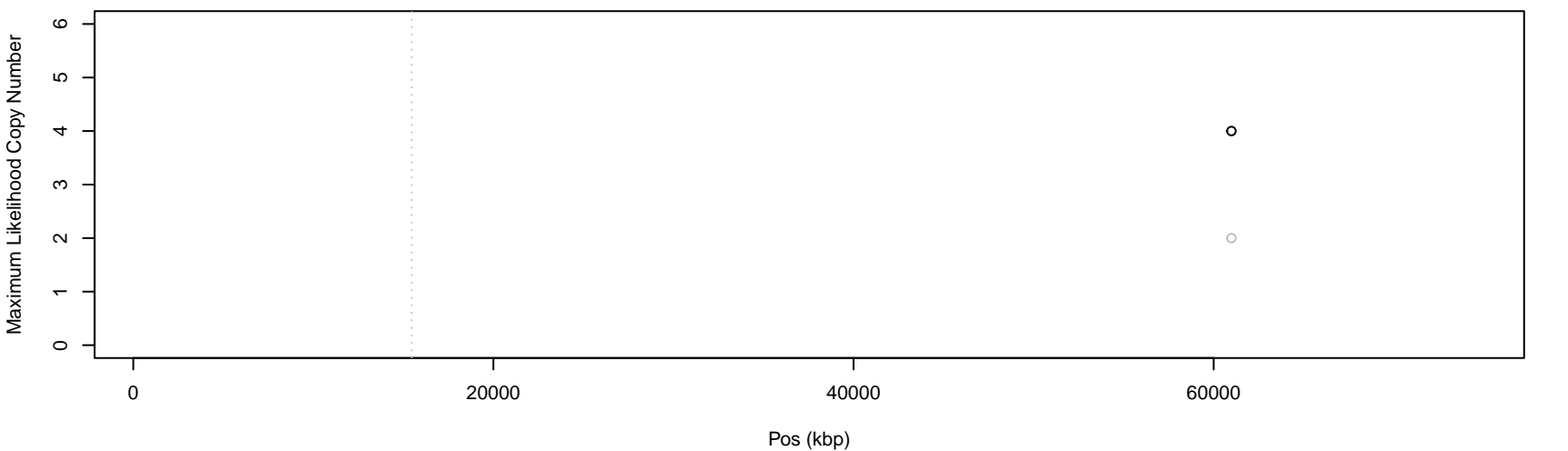
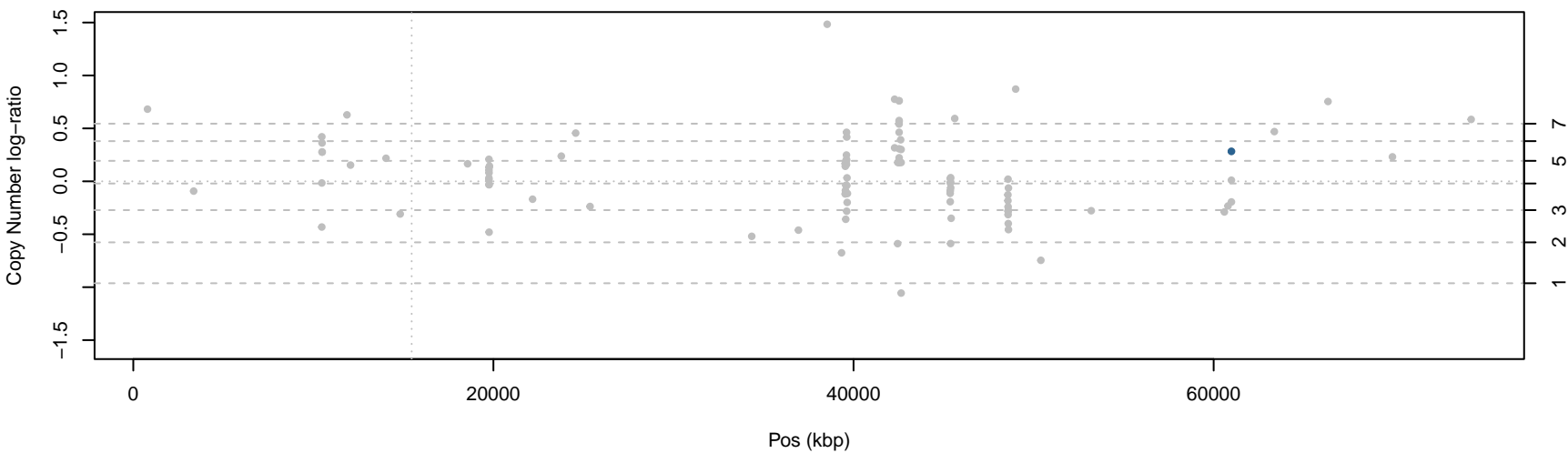
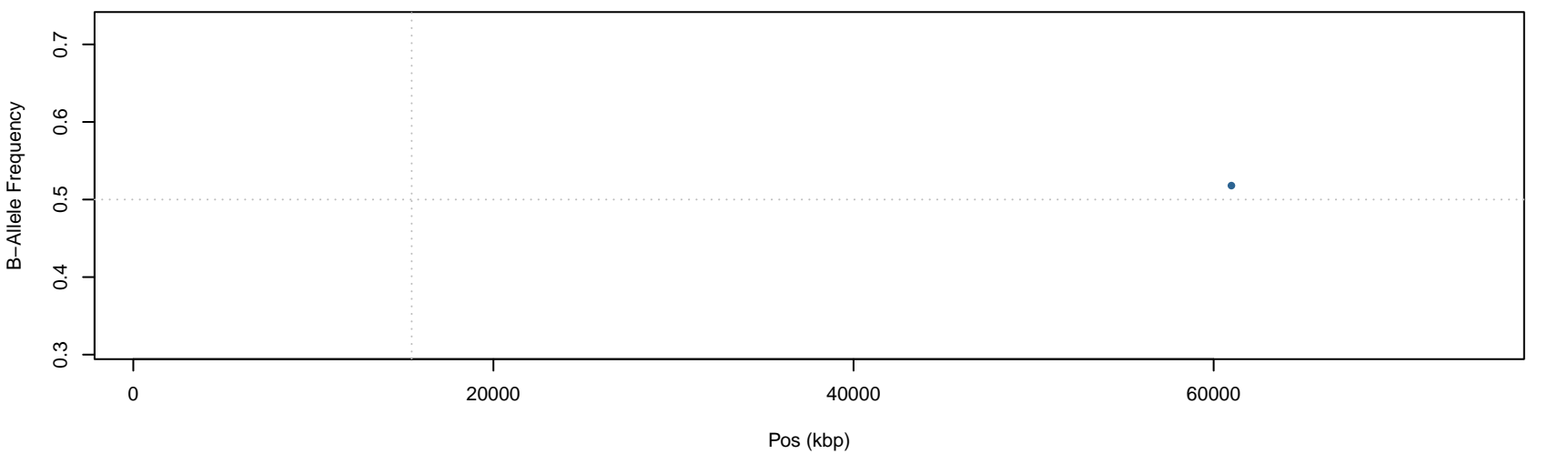
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 14



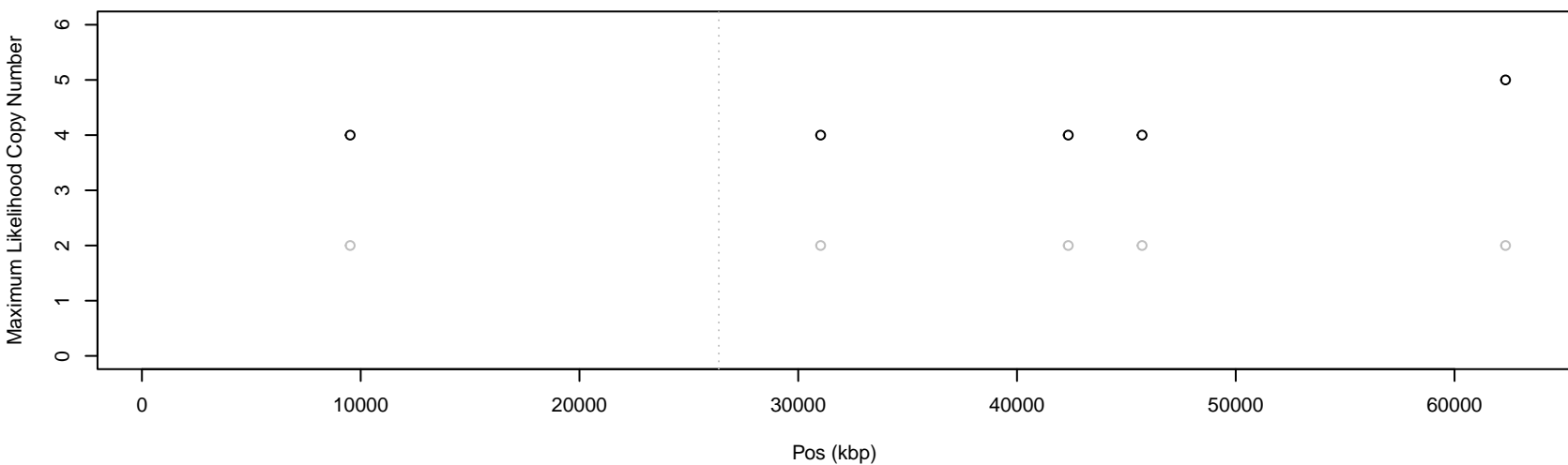
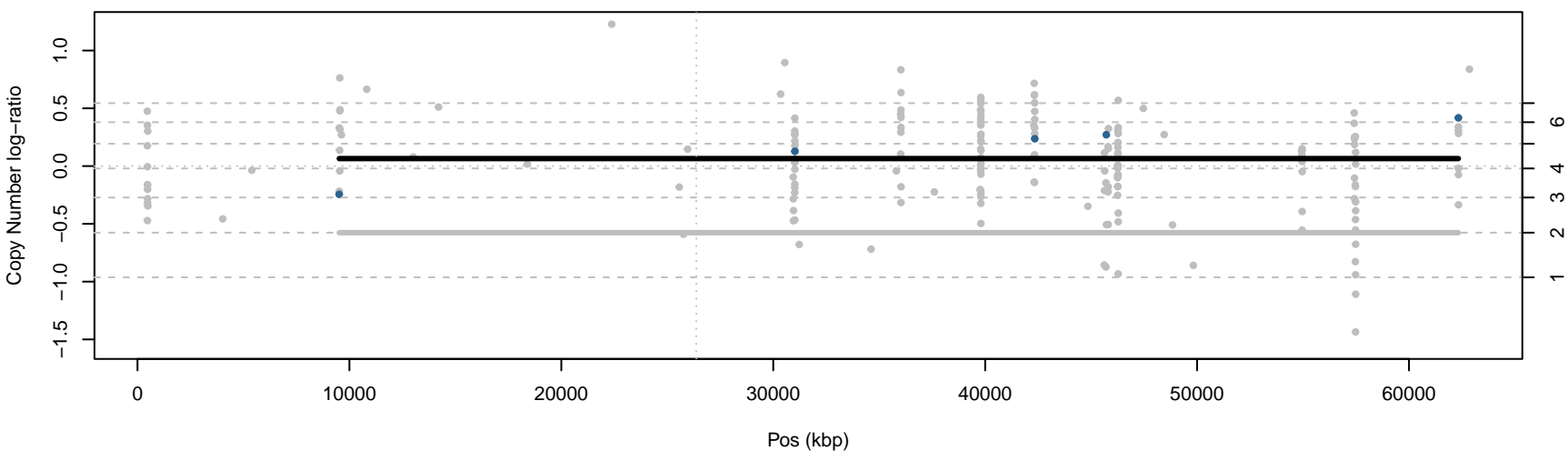
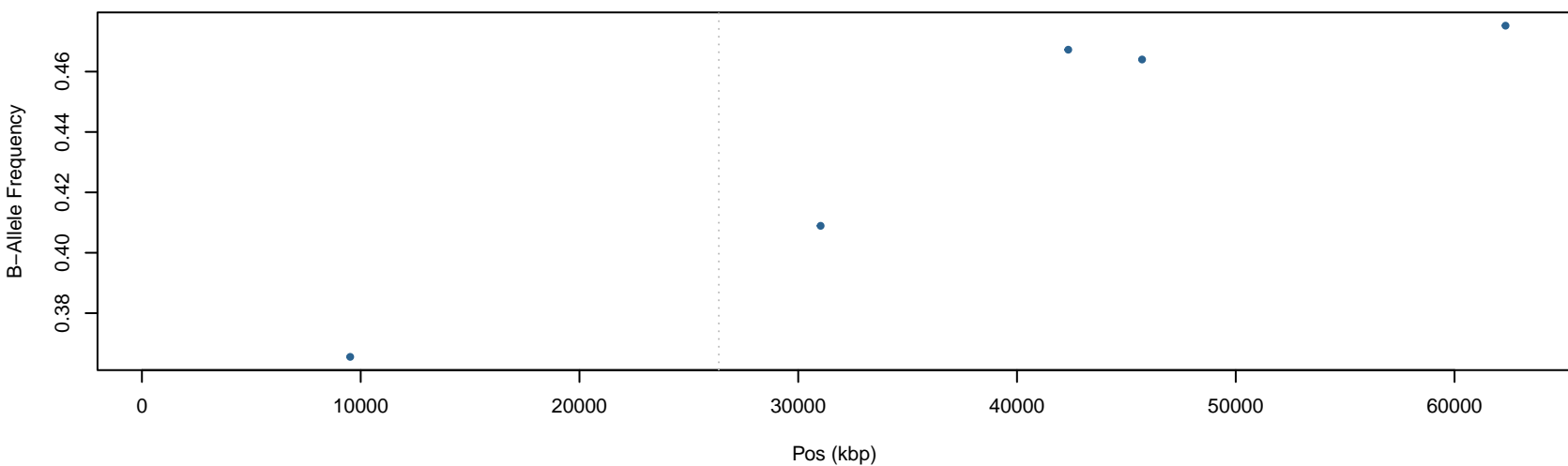
Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 15



Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 18



Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 20





Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709 Chromosome: 22

