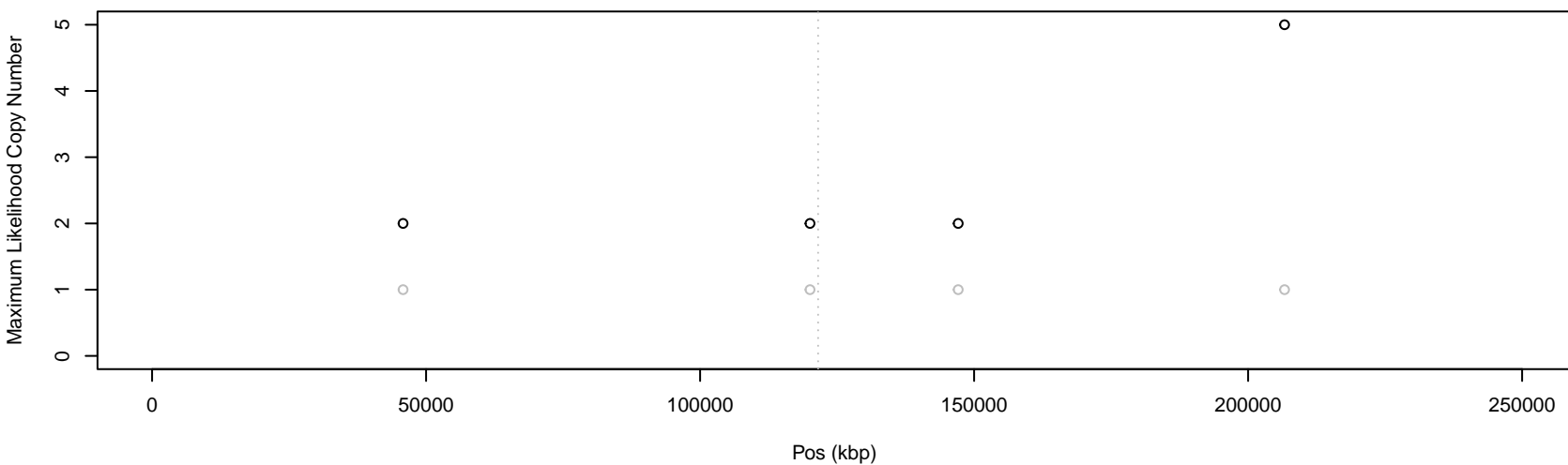
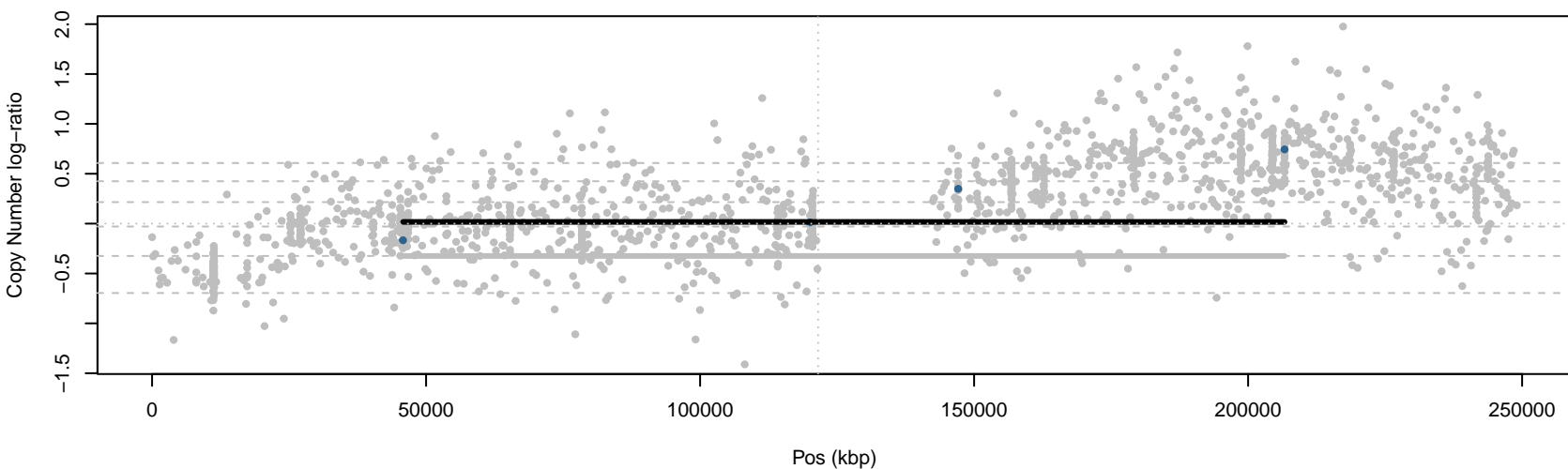
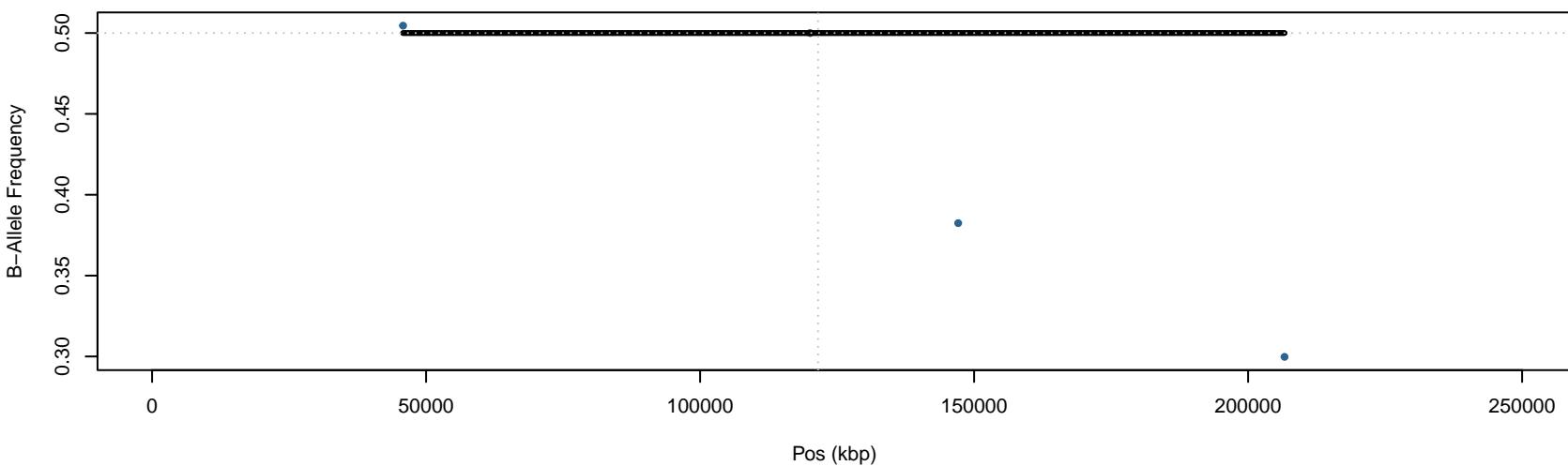
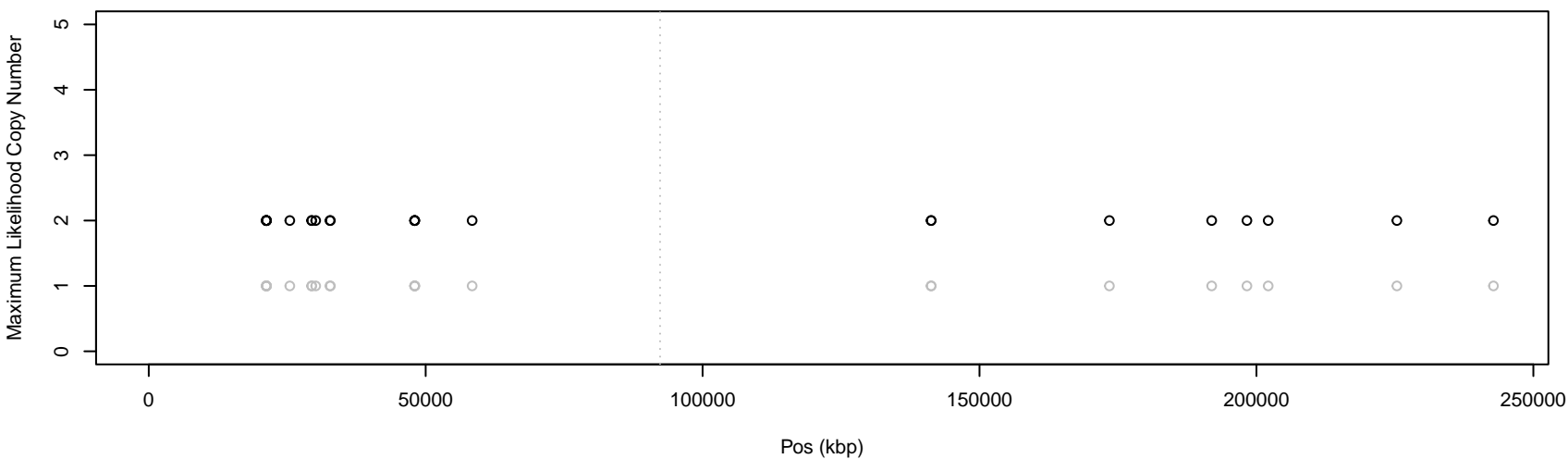
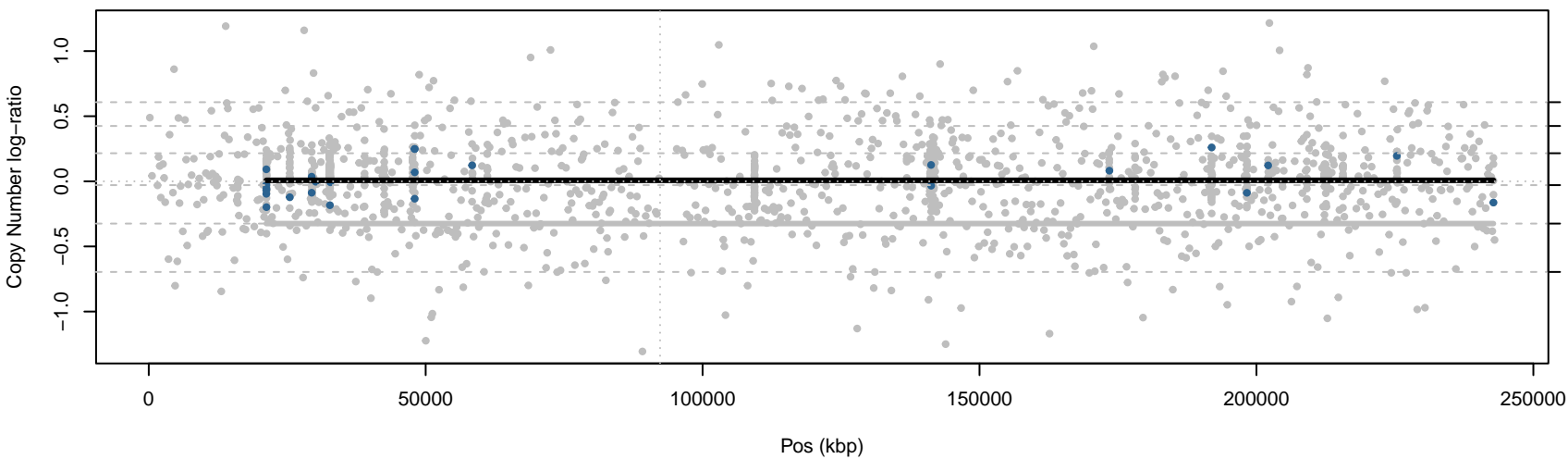
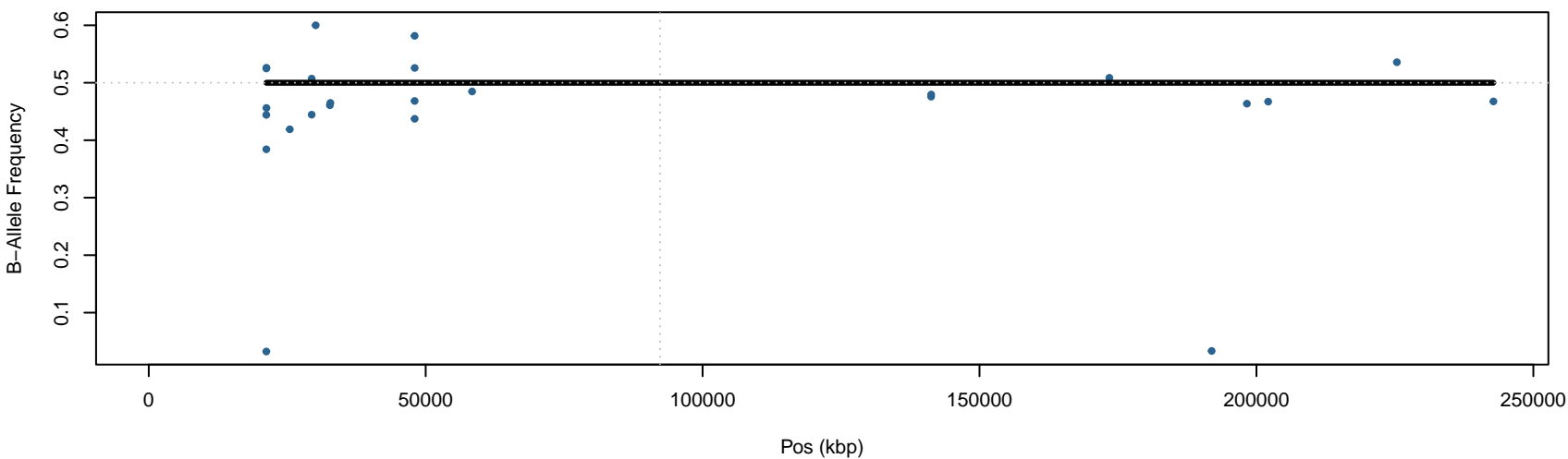


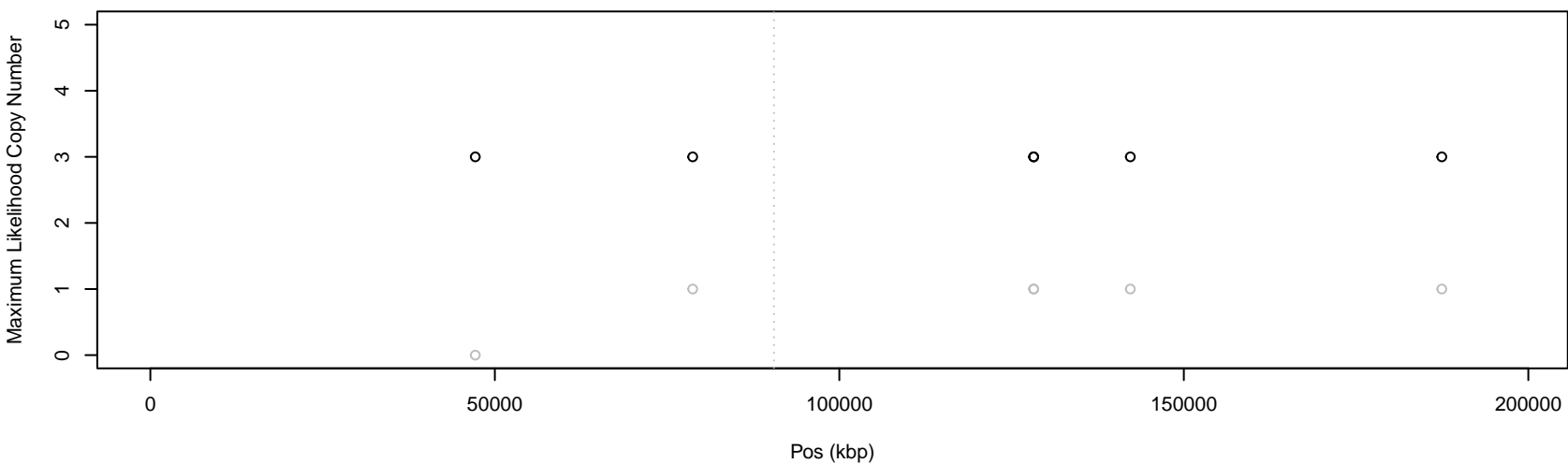
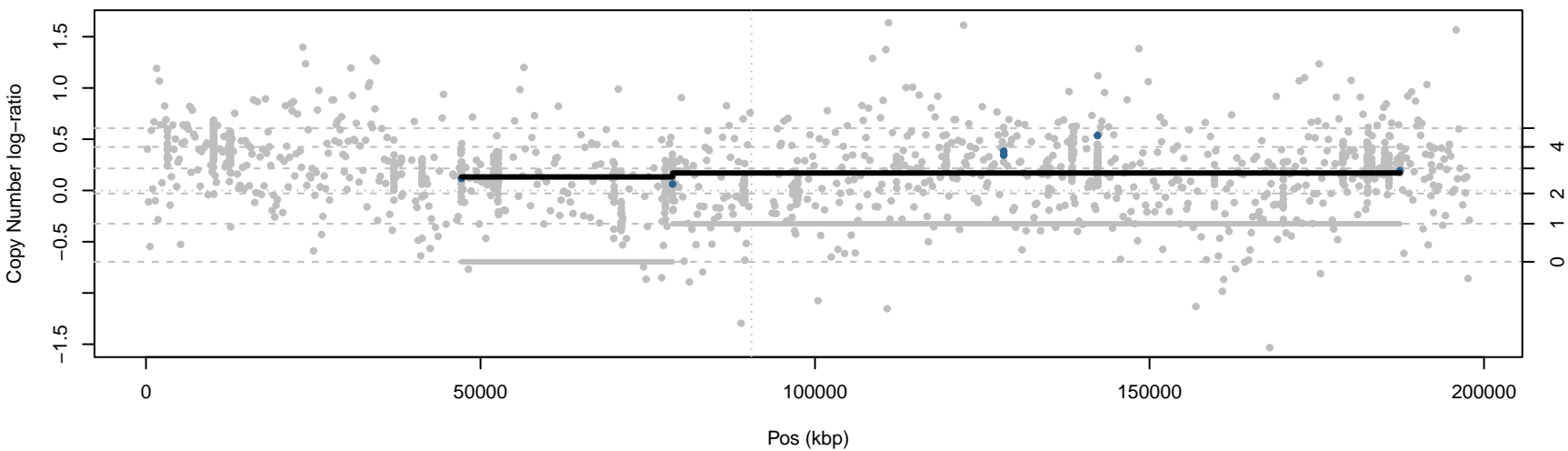
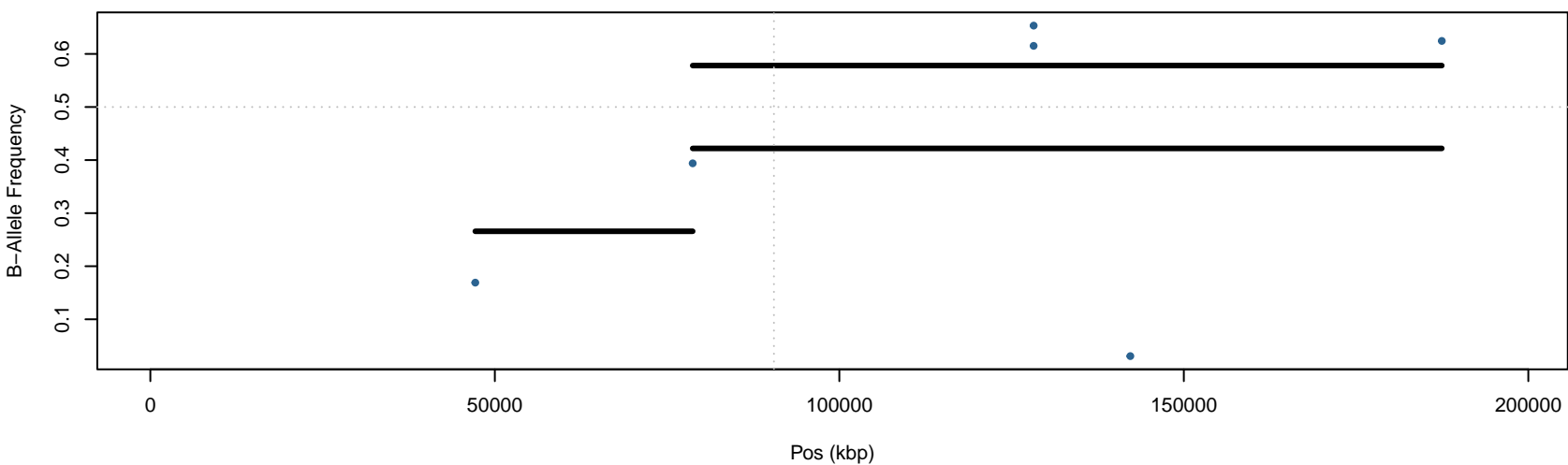
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 1



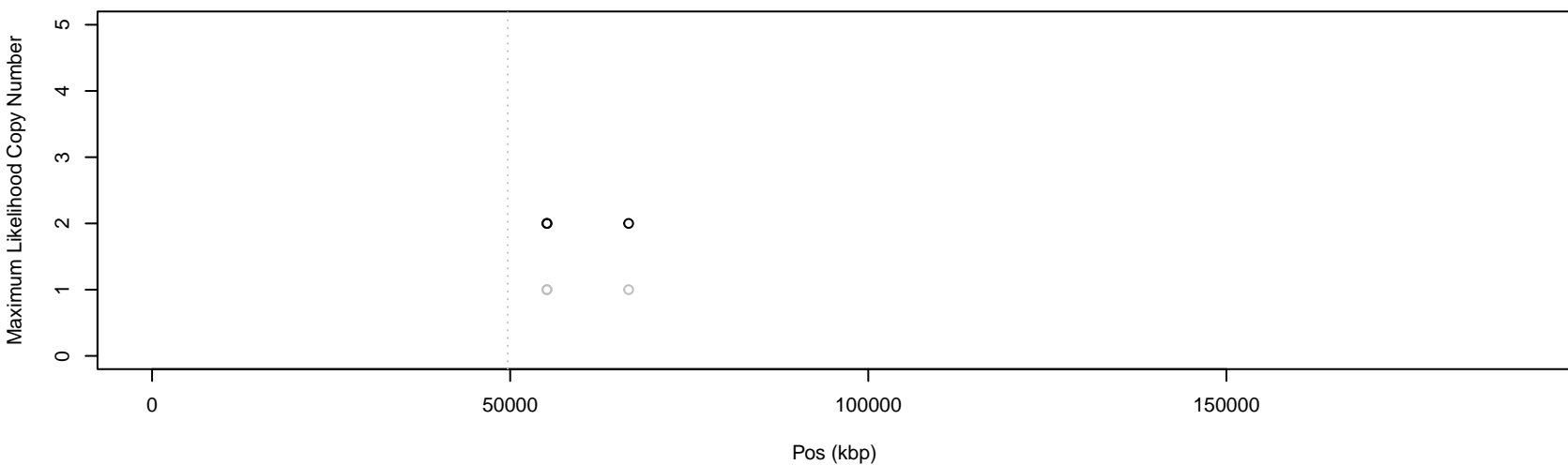
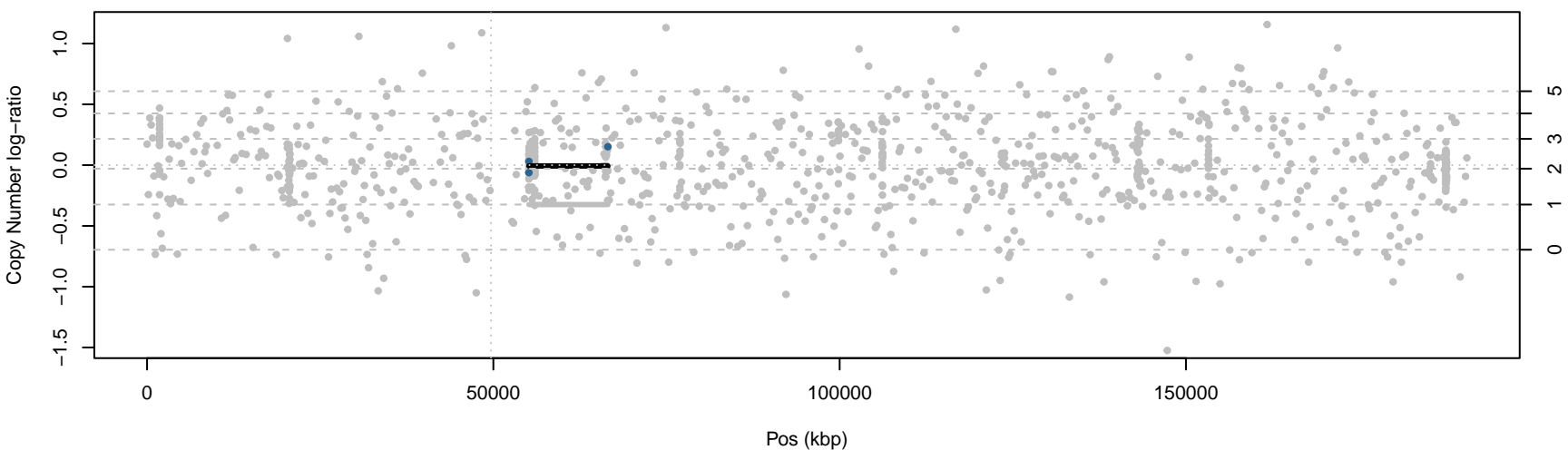
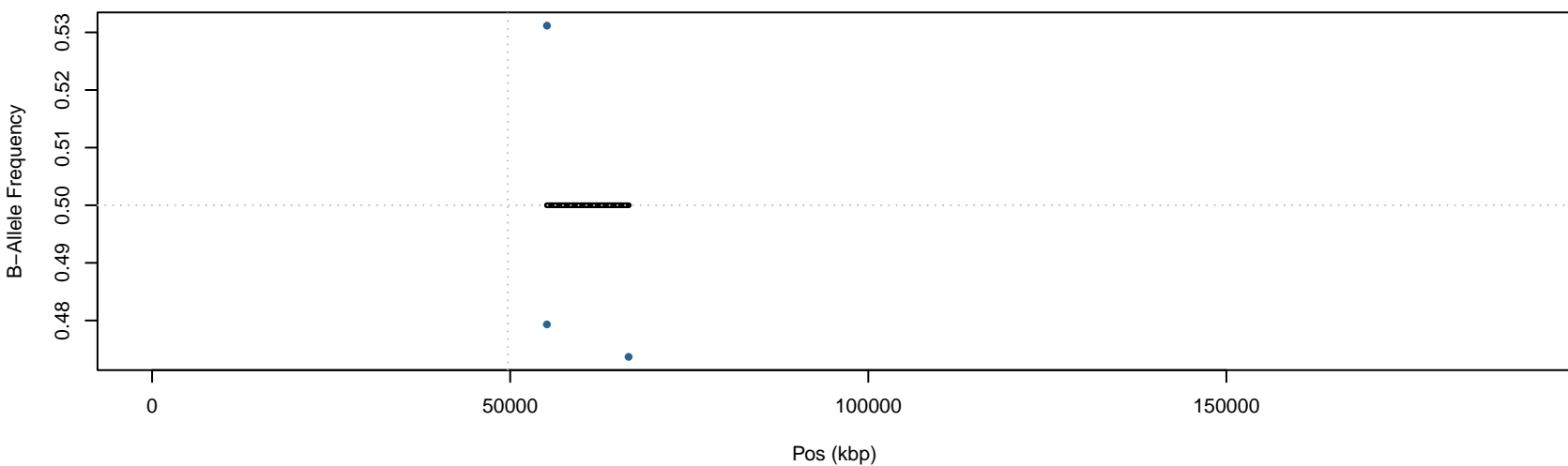
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 2



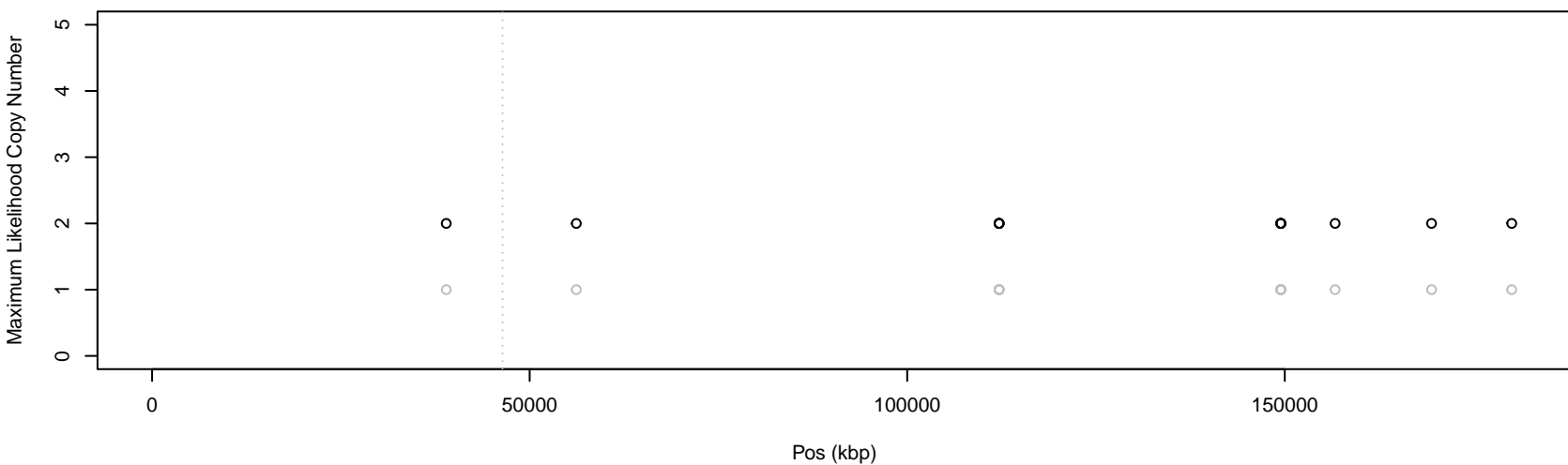
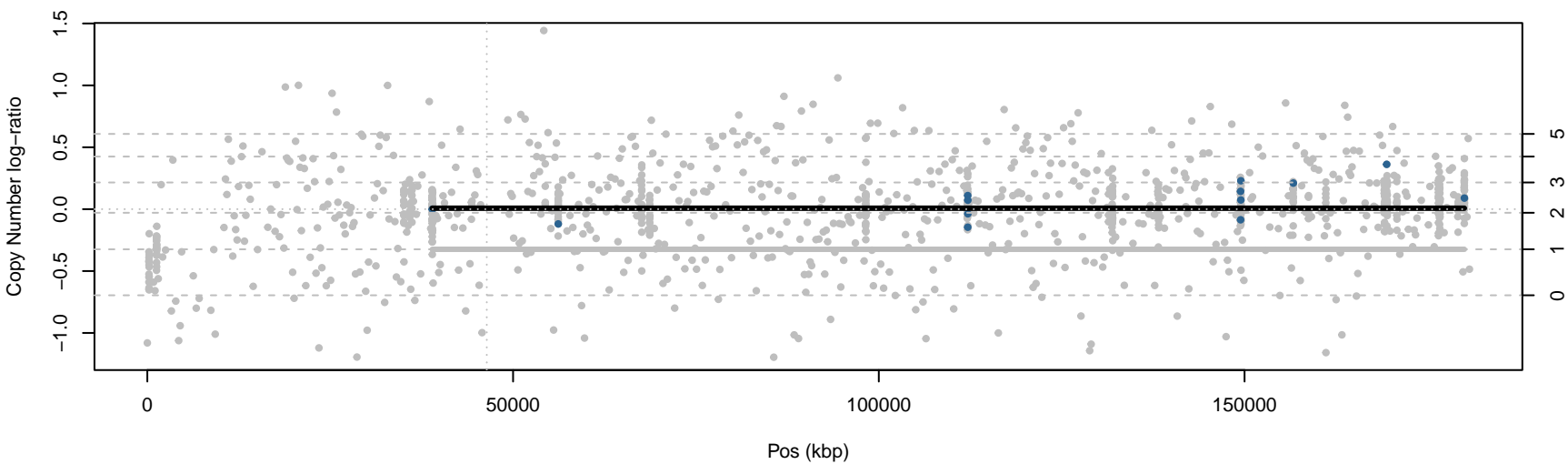
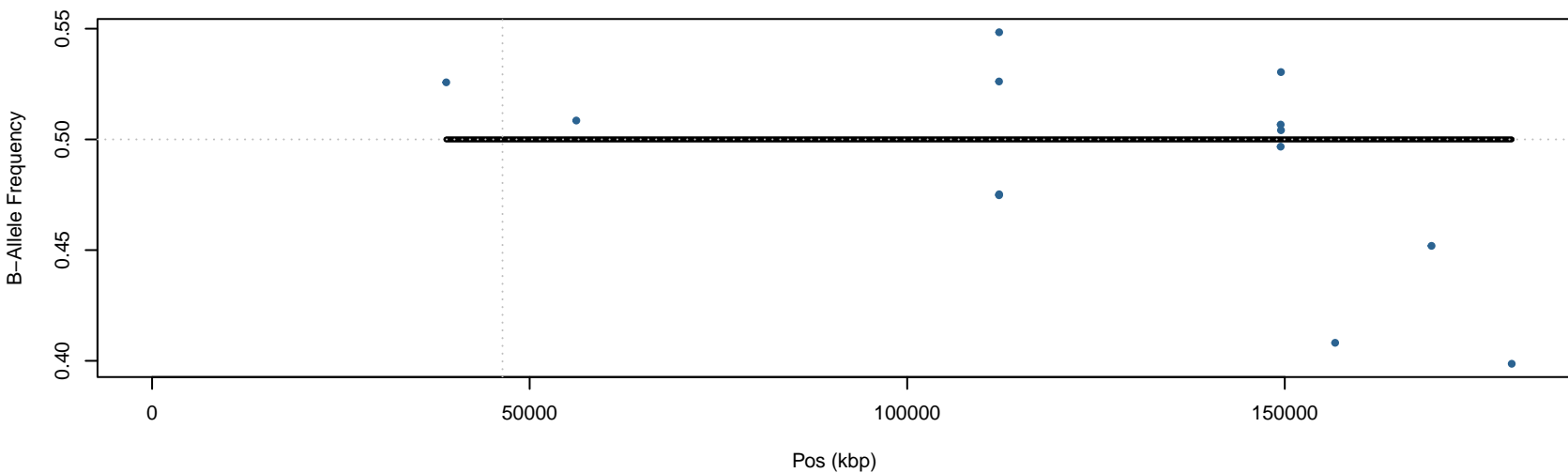
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 3



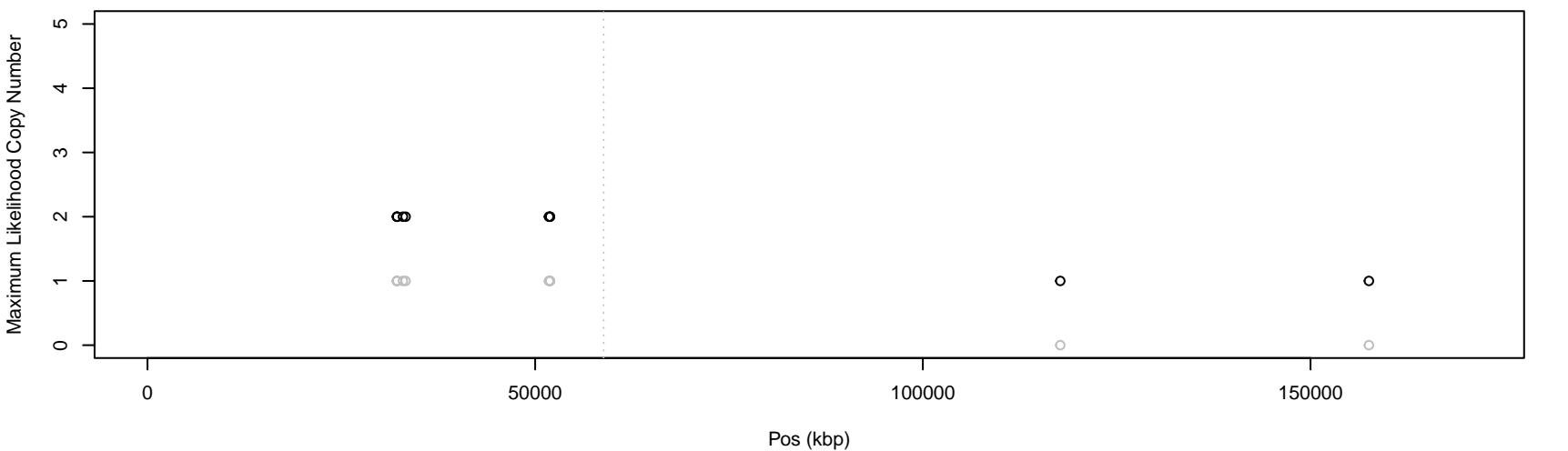
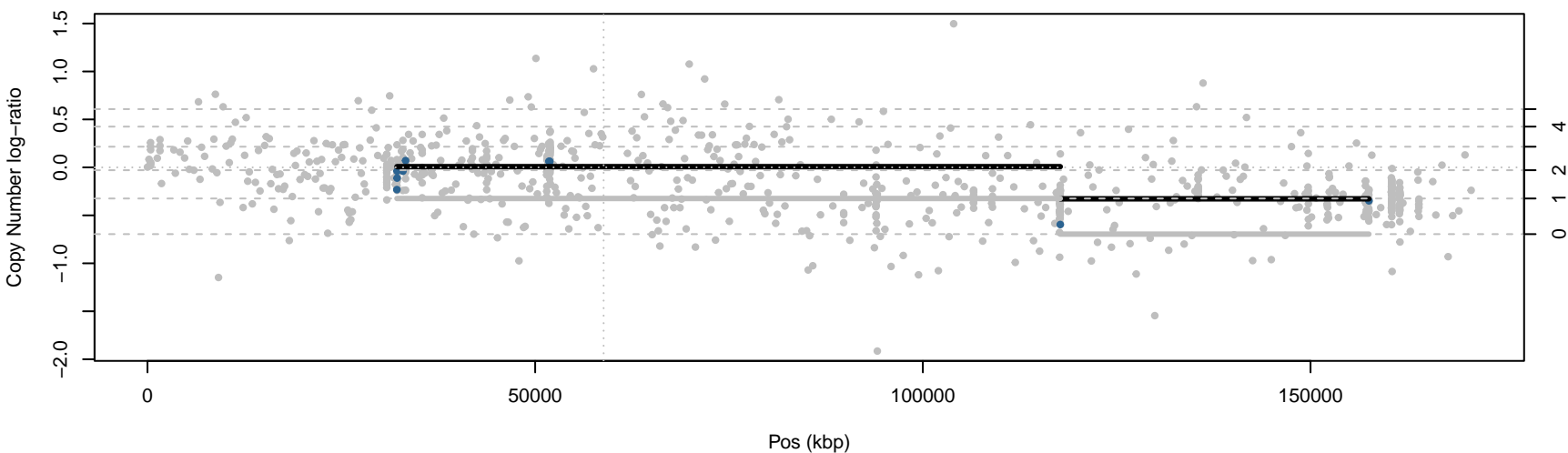
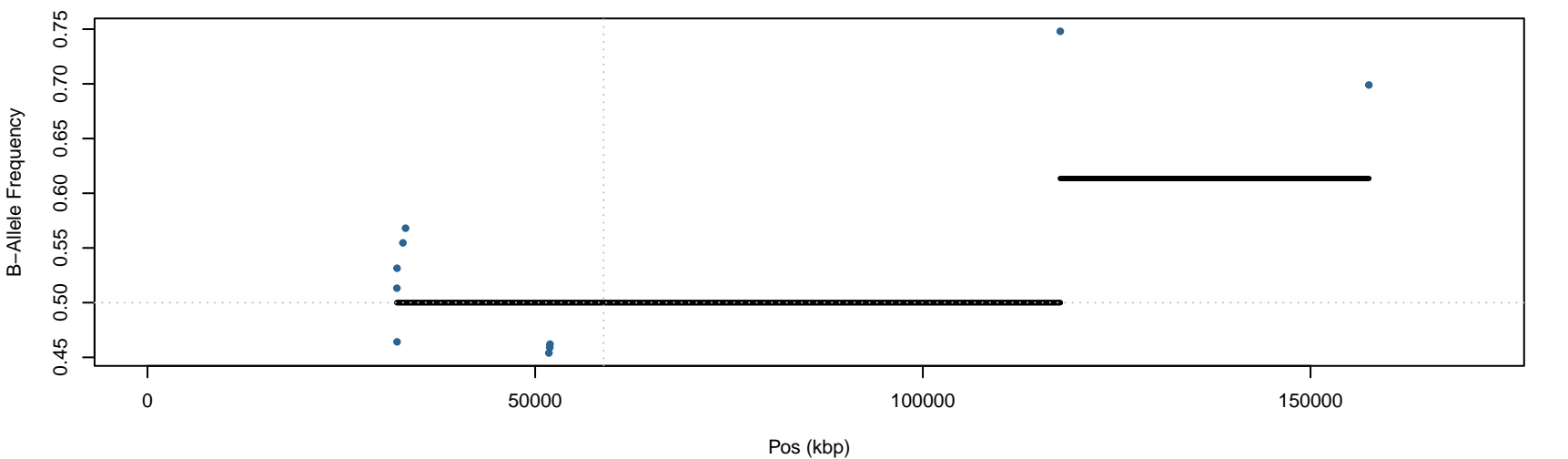
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 4



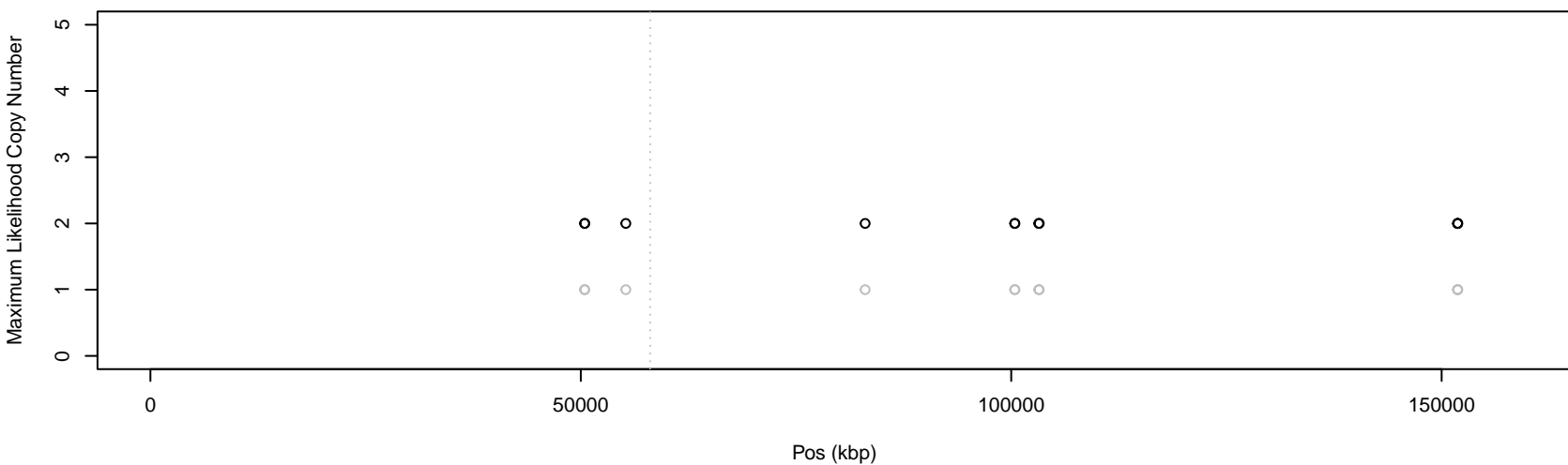
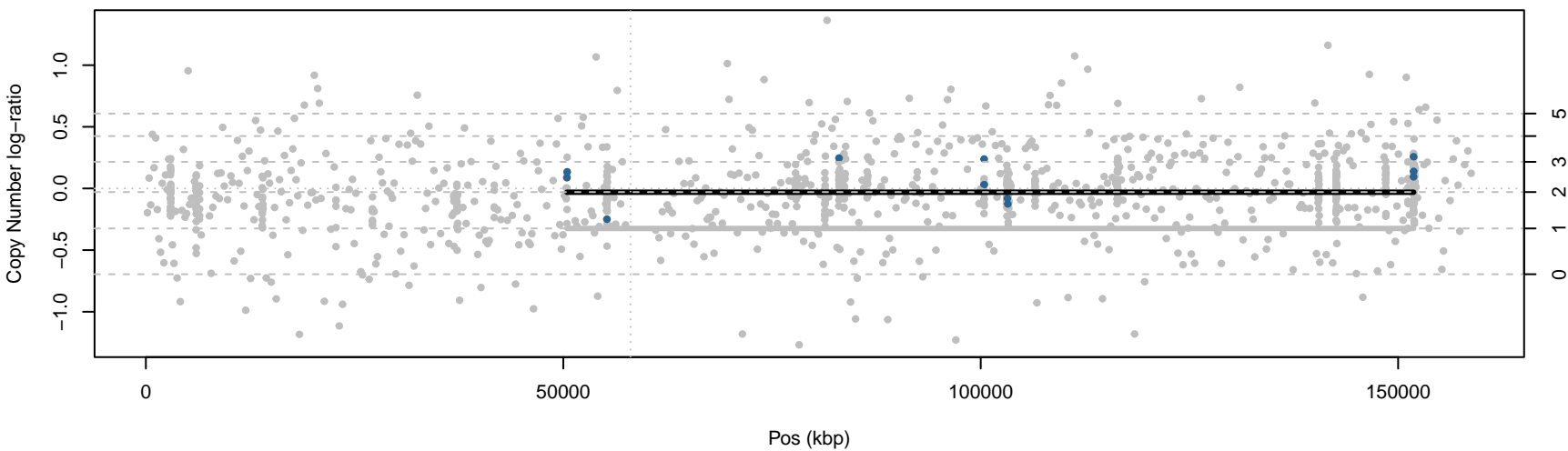
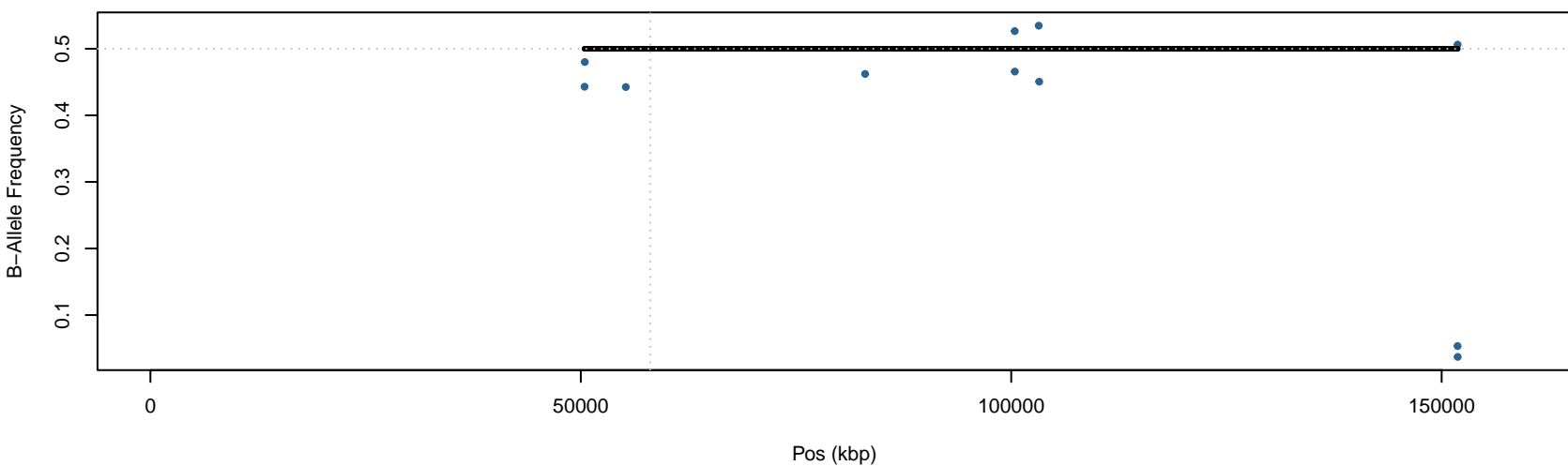
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 5



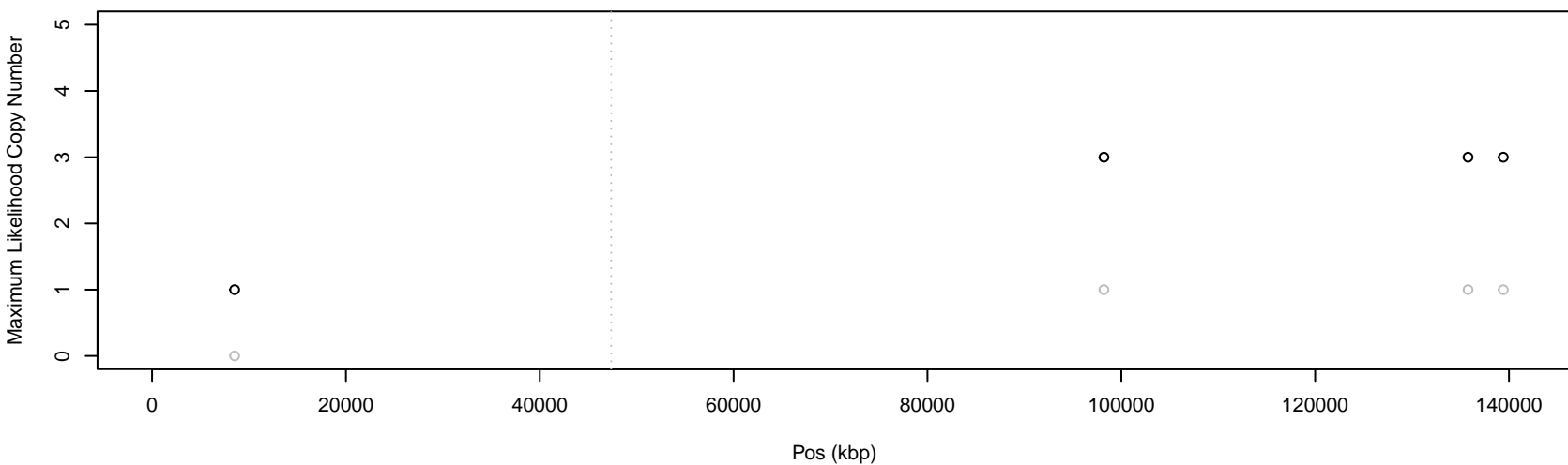
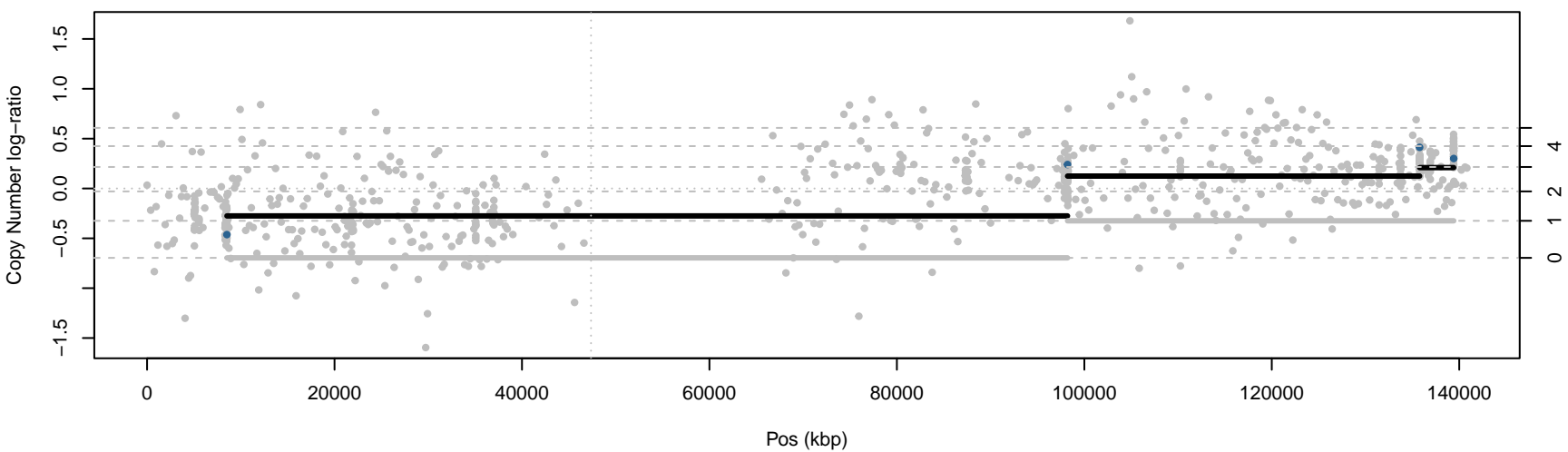
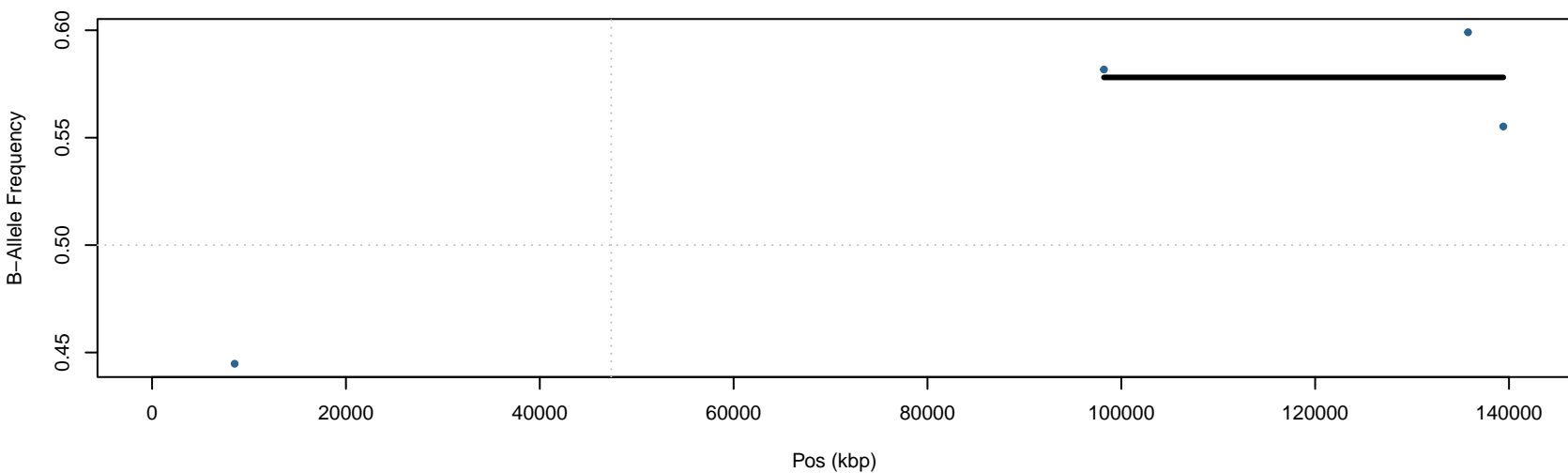
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 6



Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 7

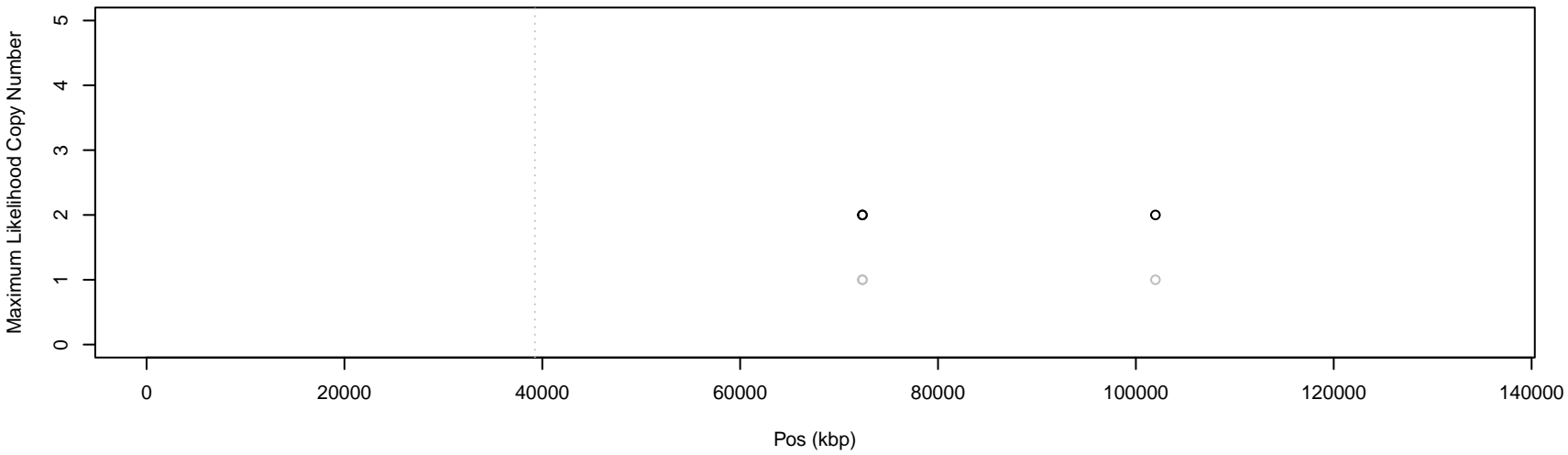
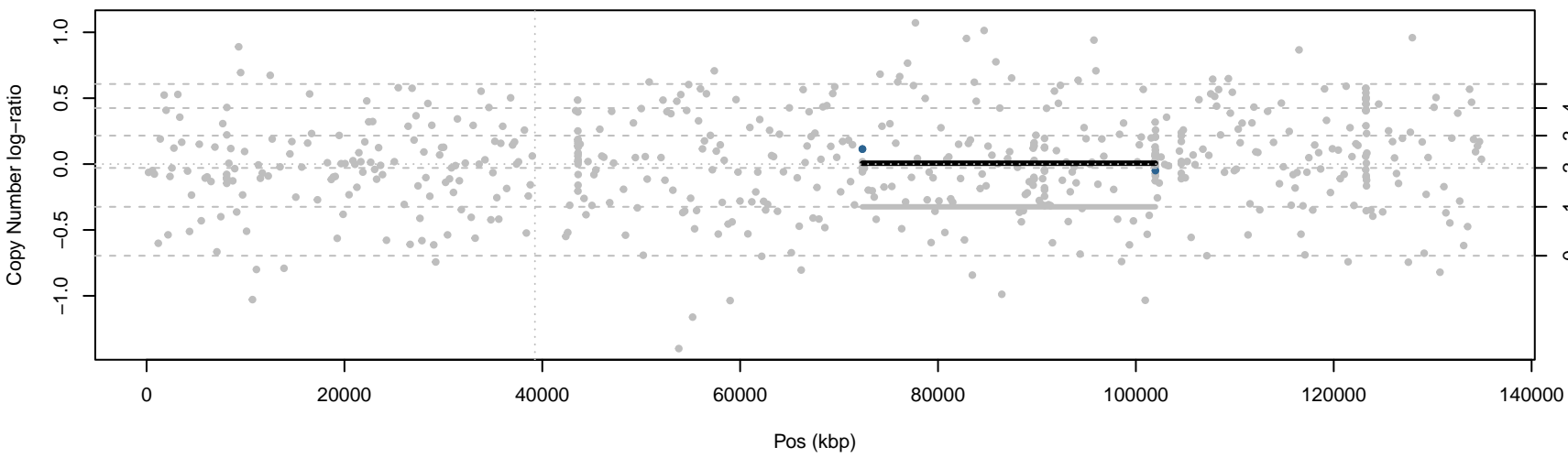
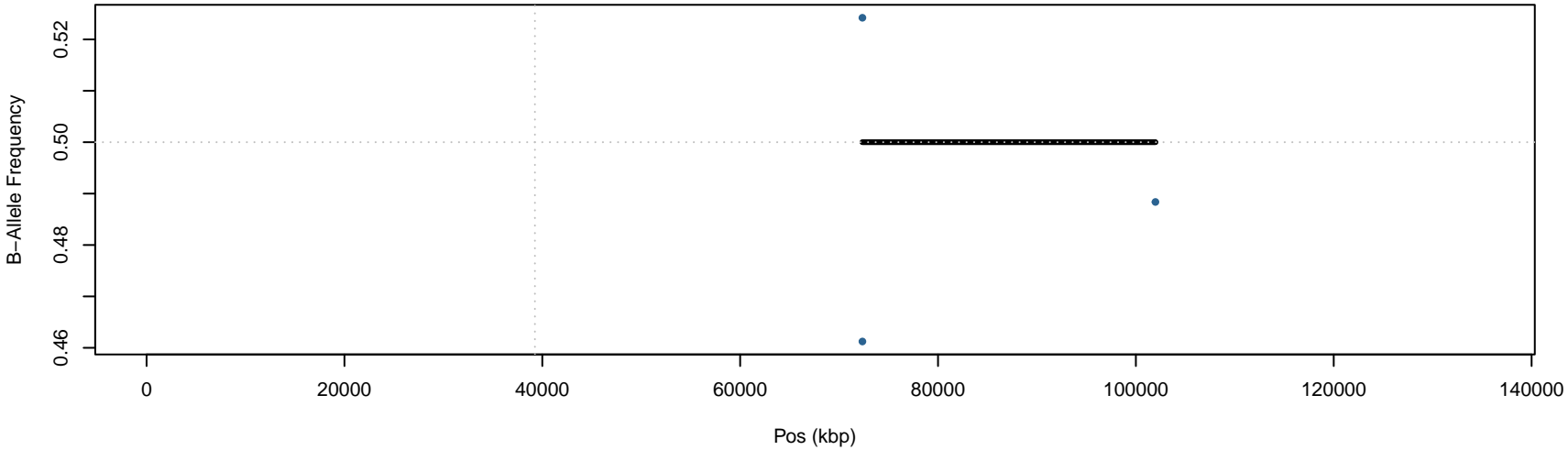


Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527,500 Chromosome: 9

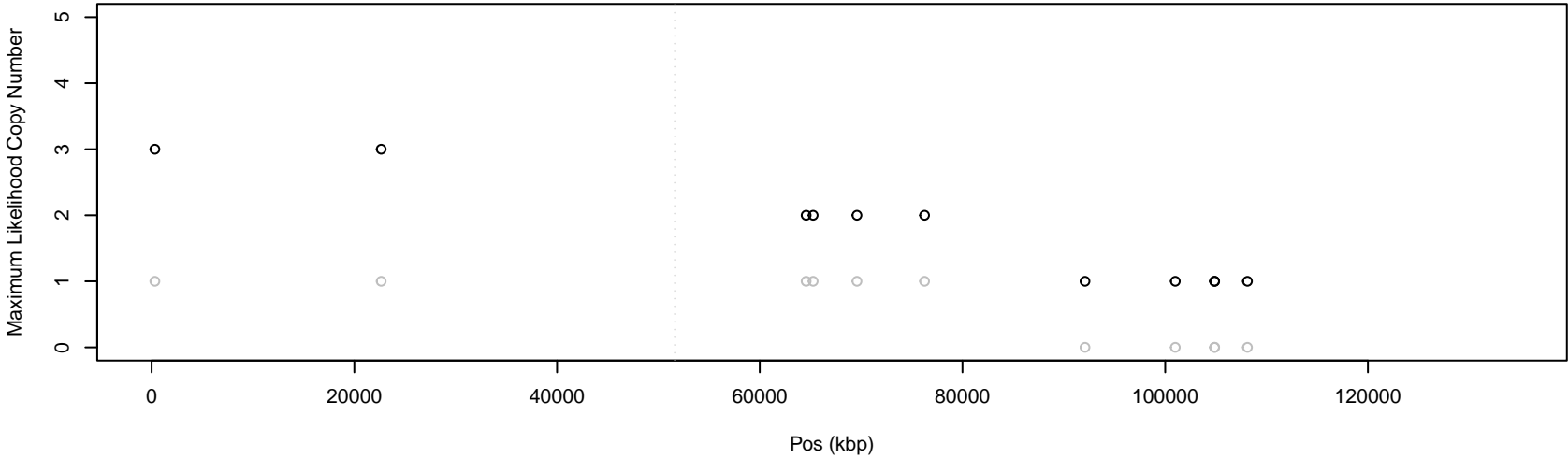
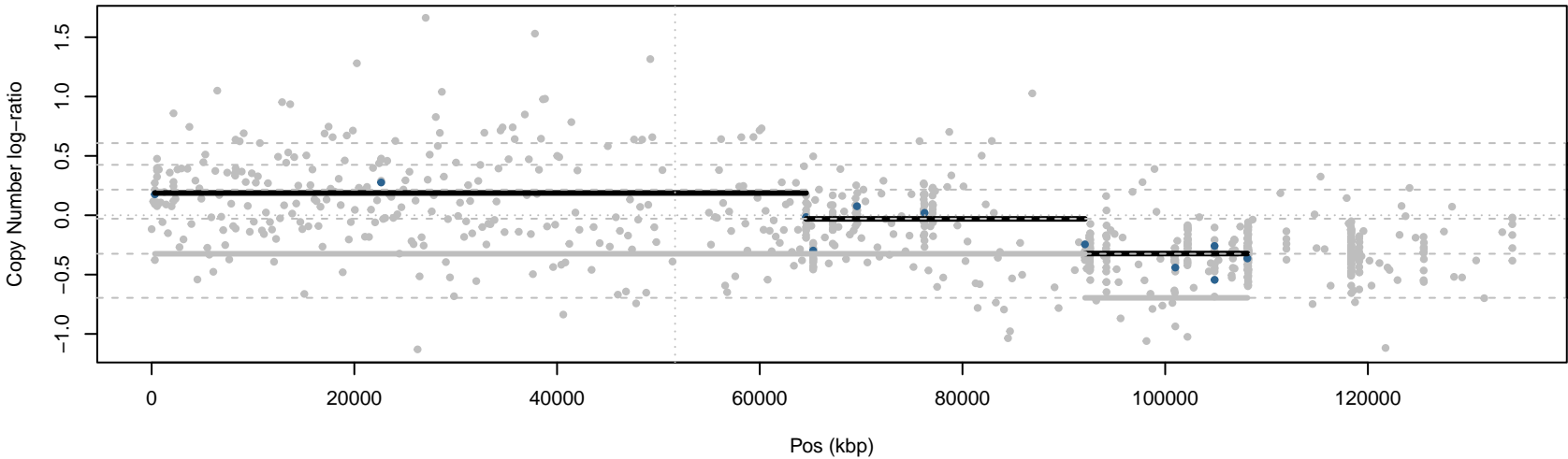
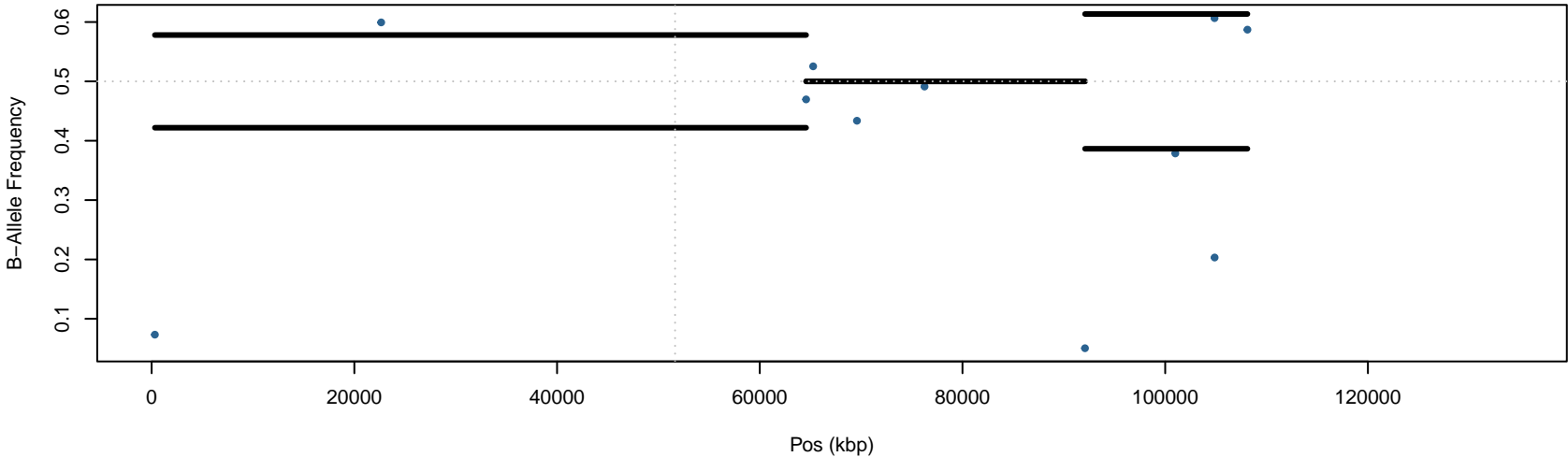




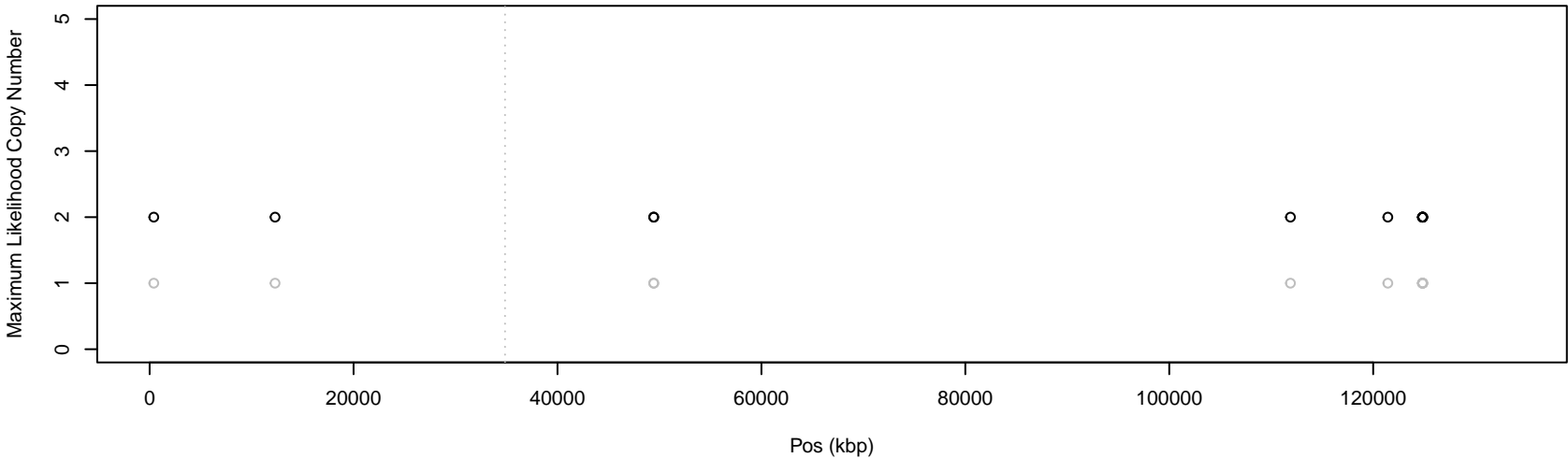
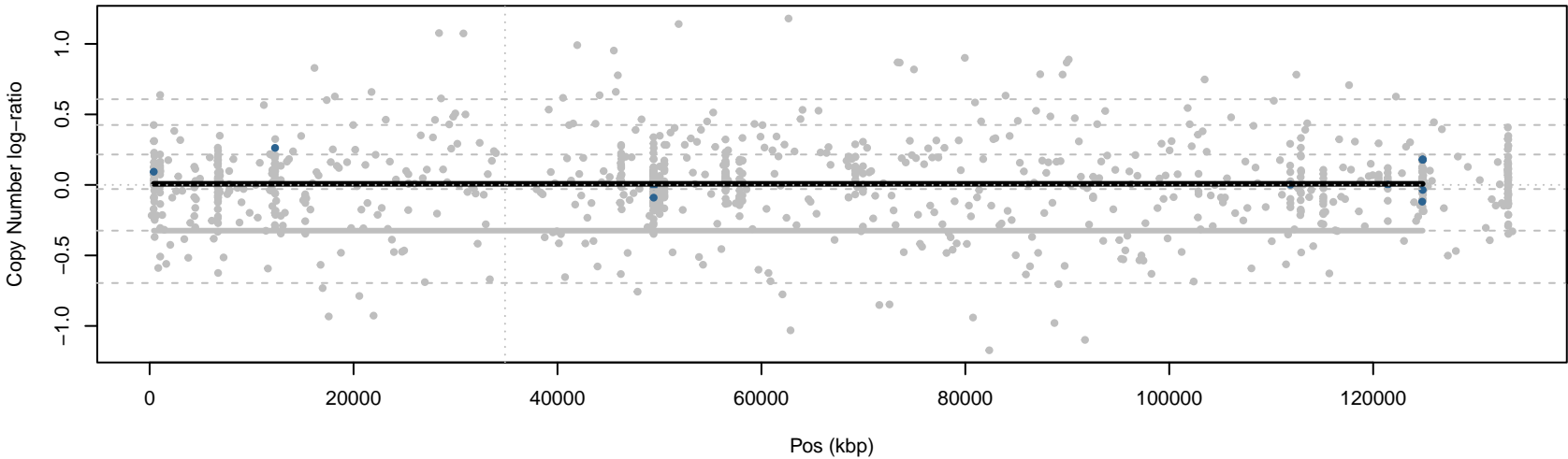
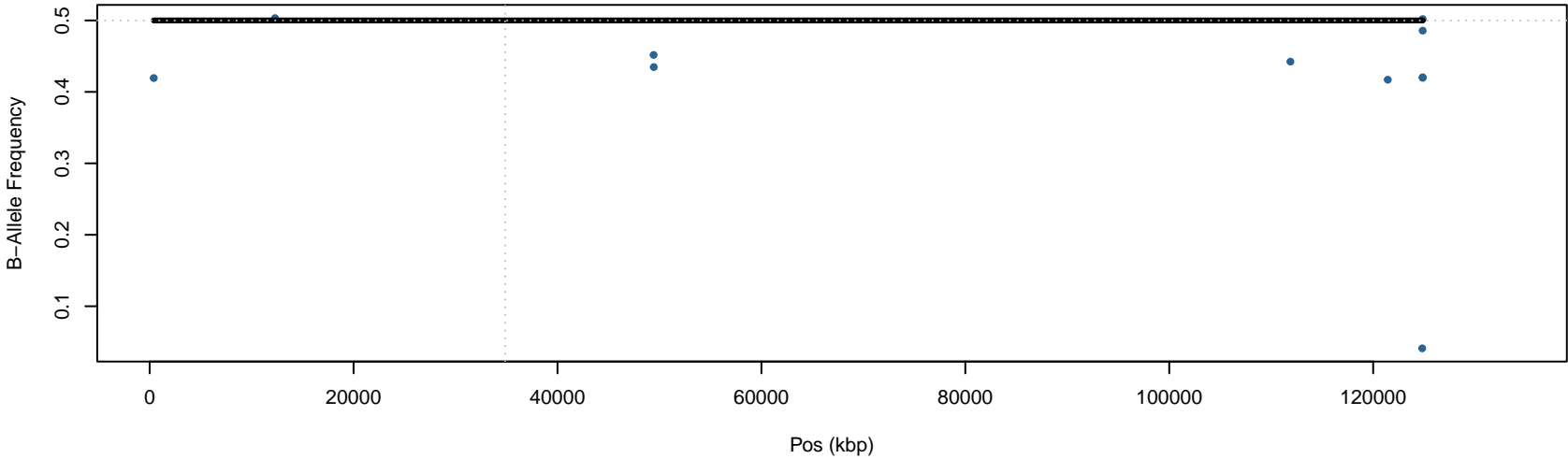
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 10



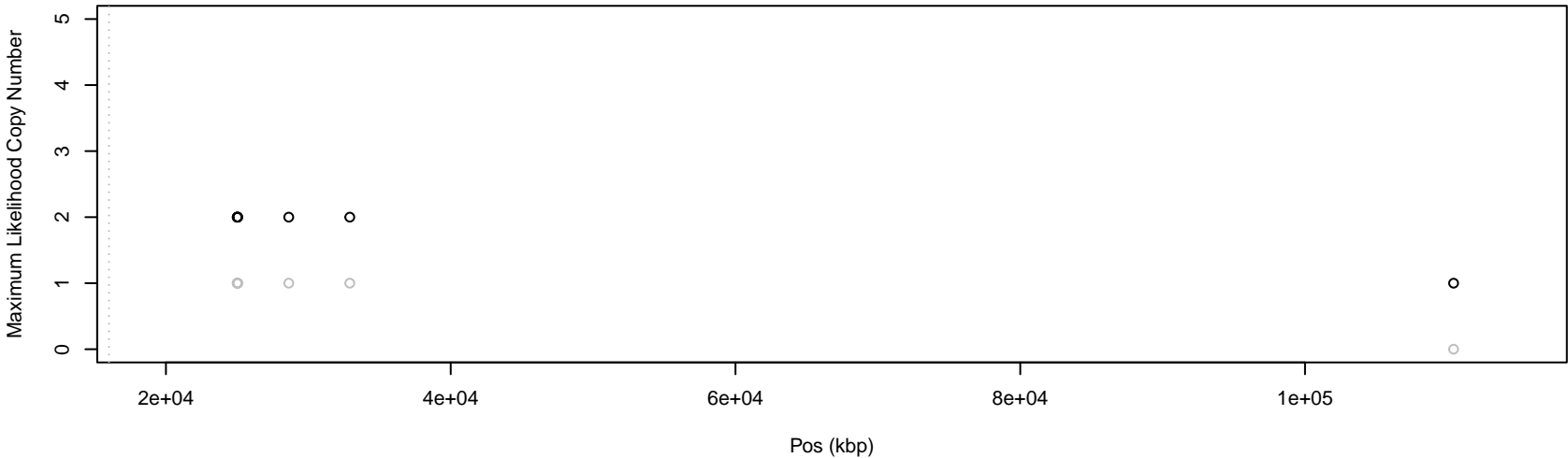
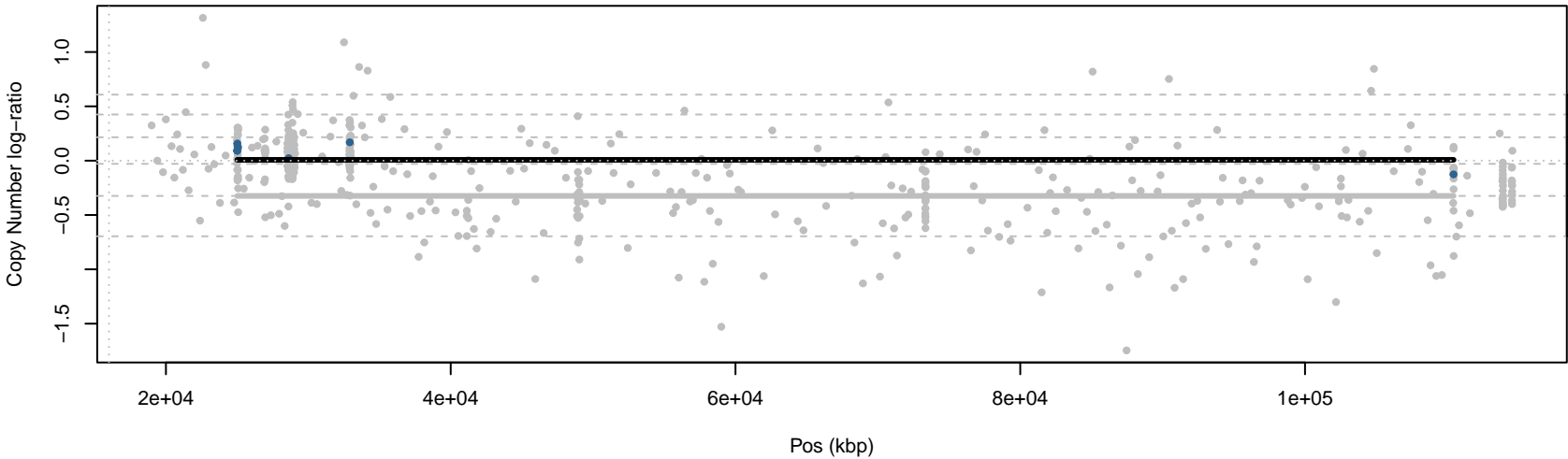
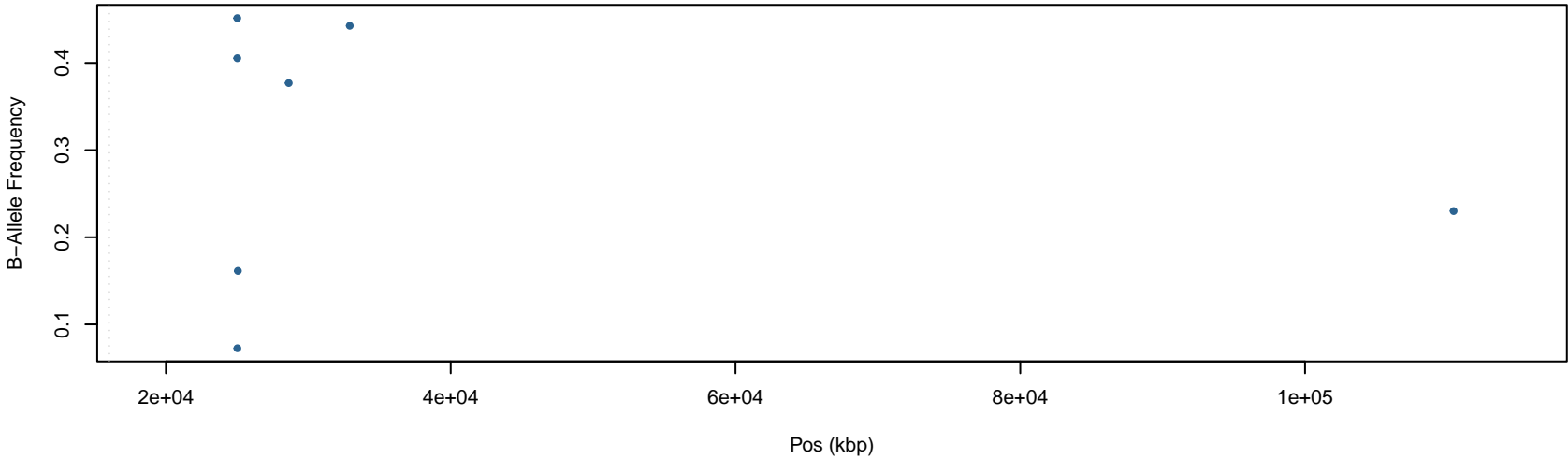
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 11



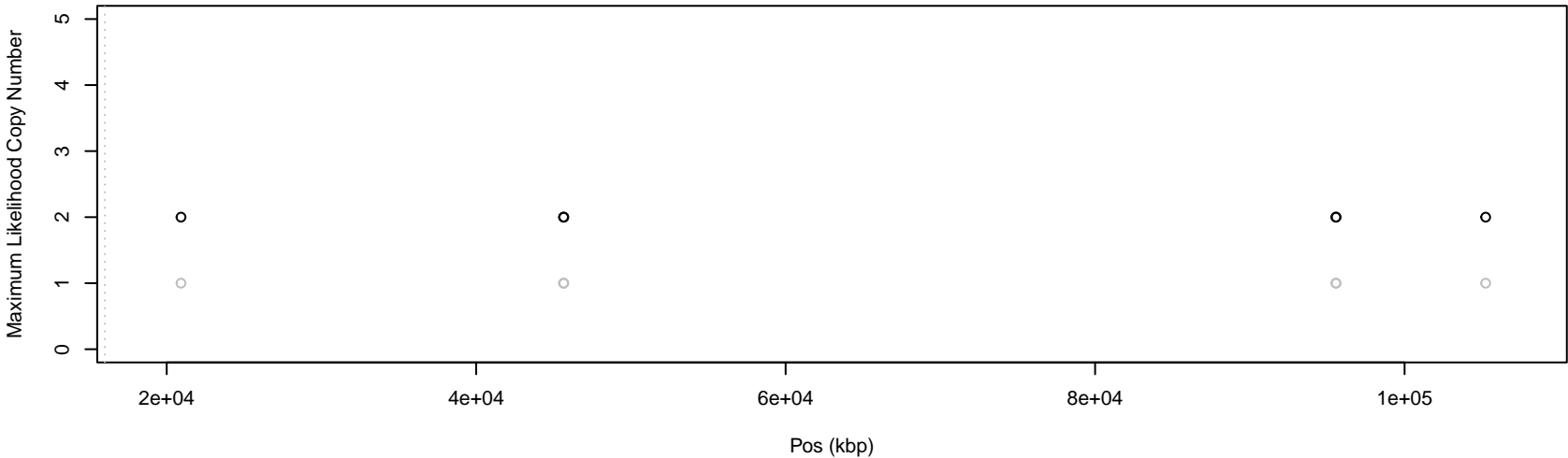
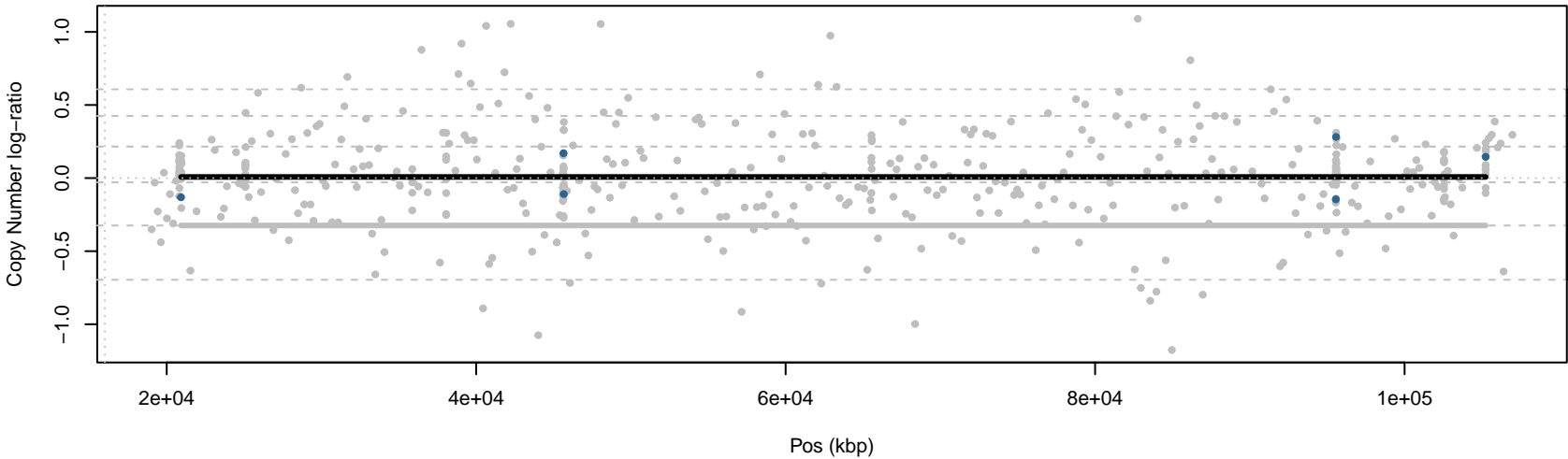
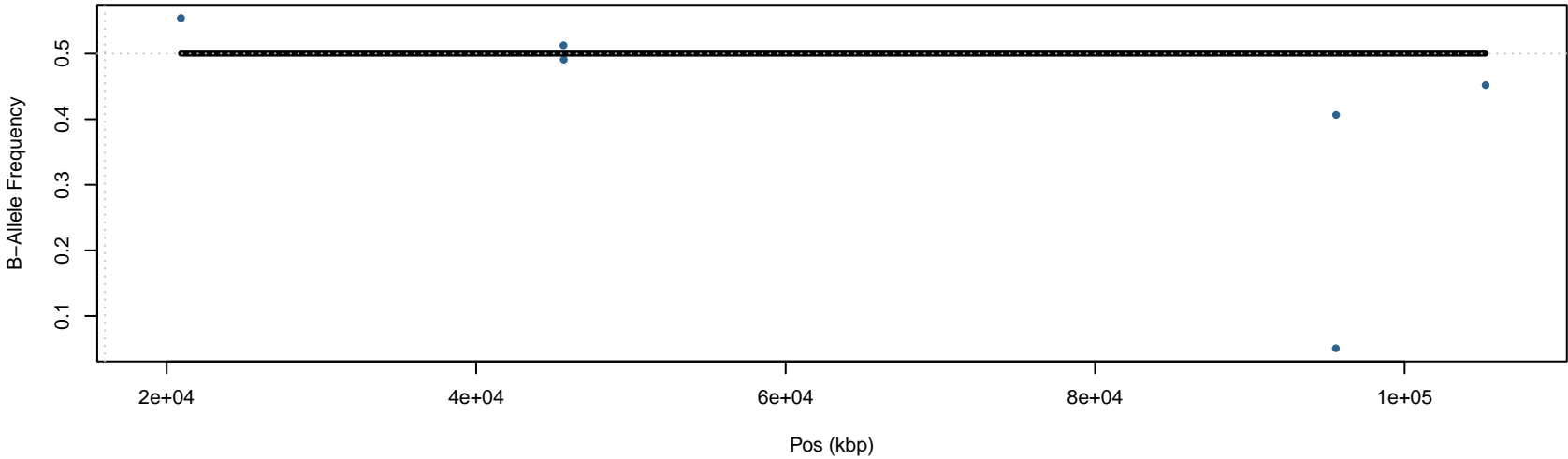
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 12



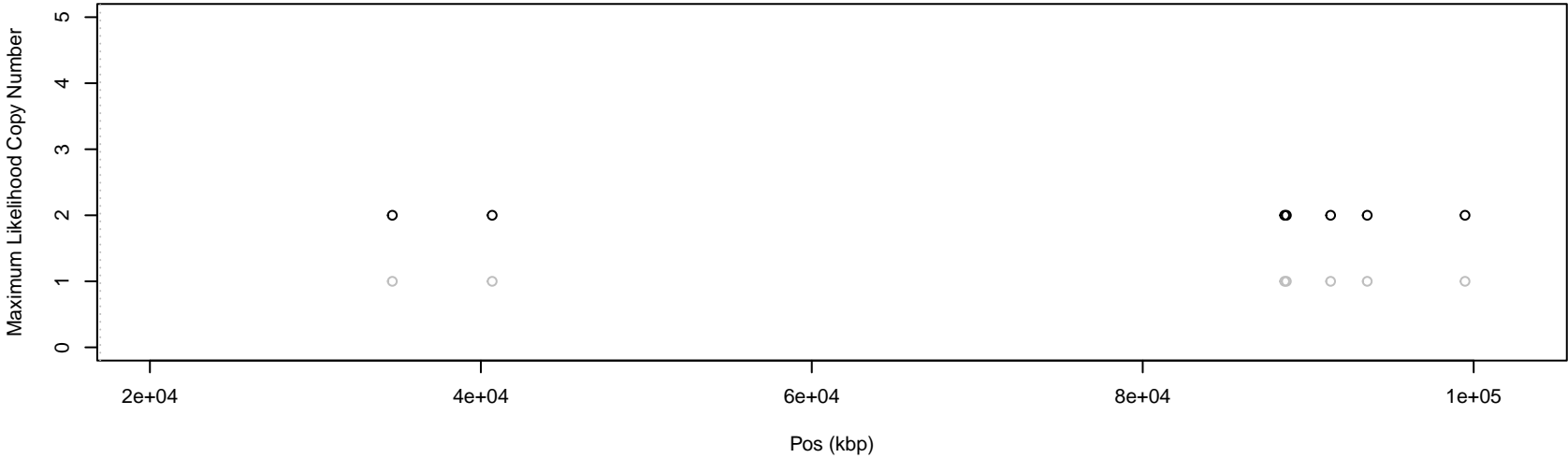
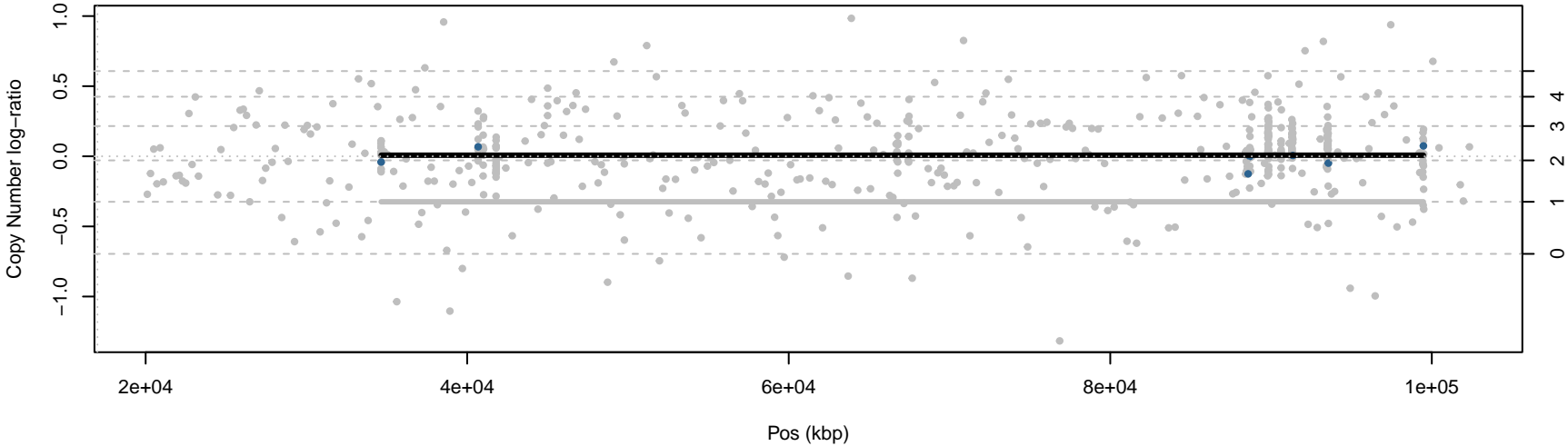
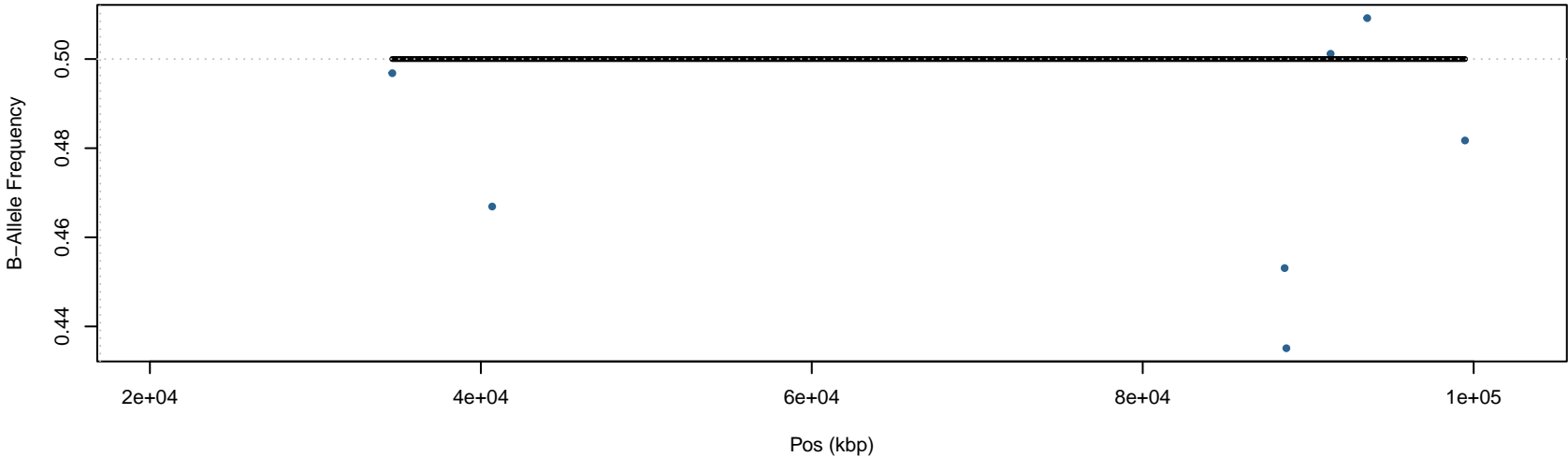
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 13



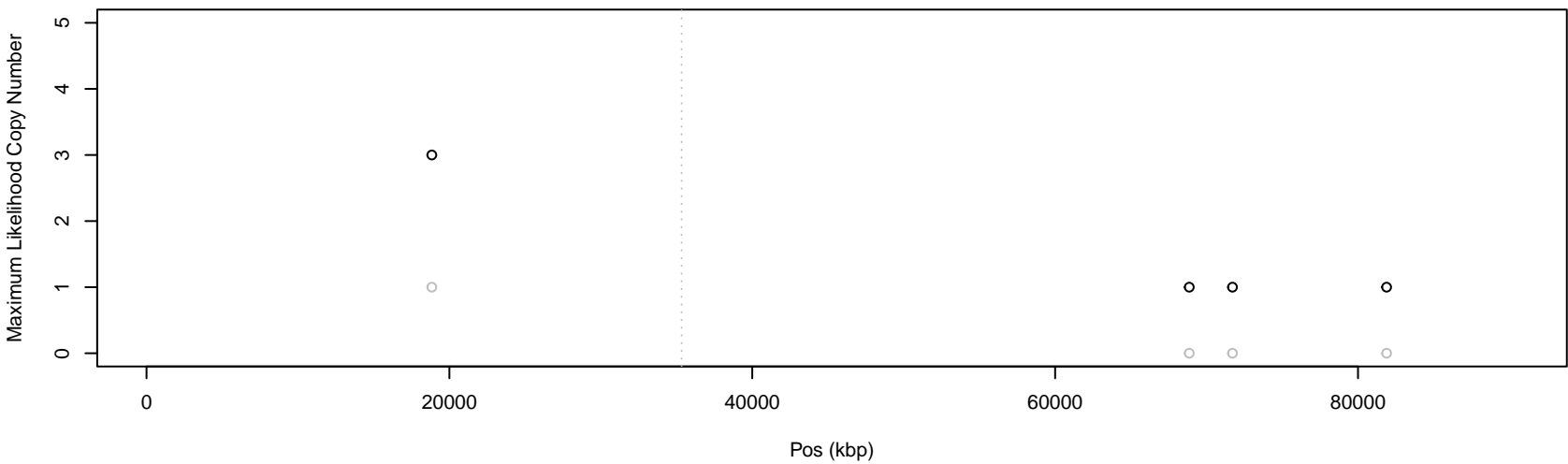
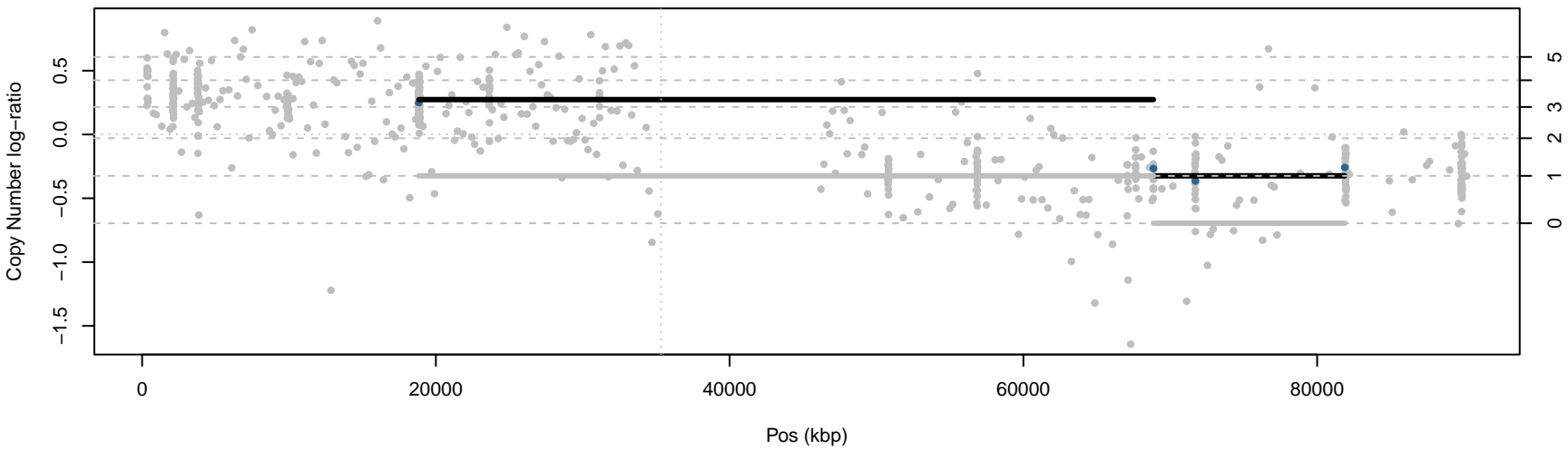
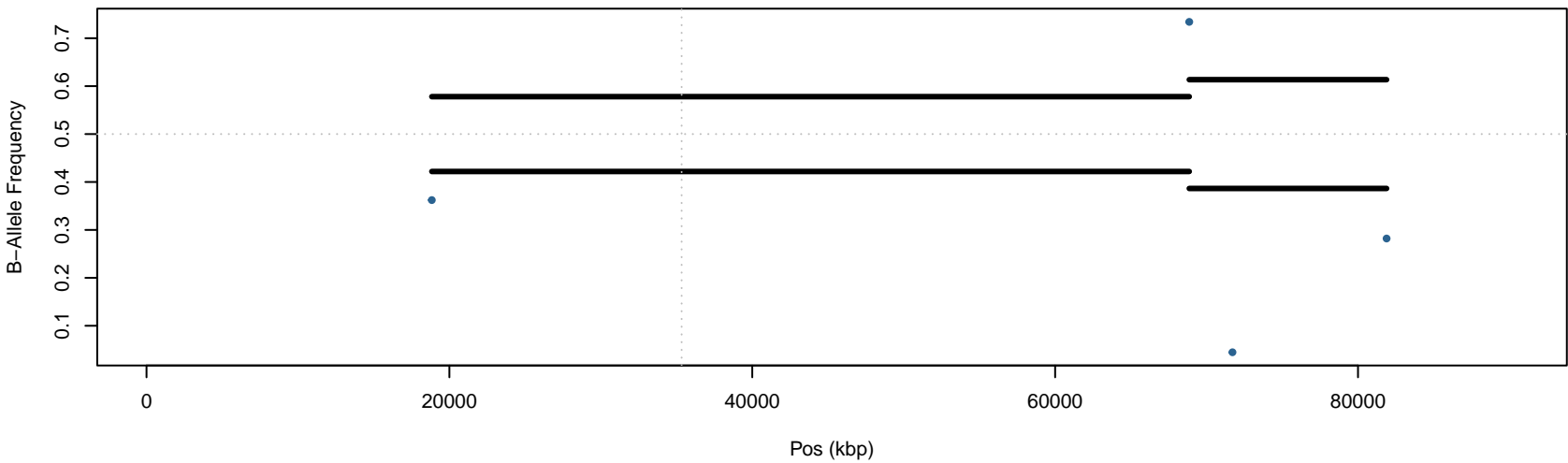
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 14



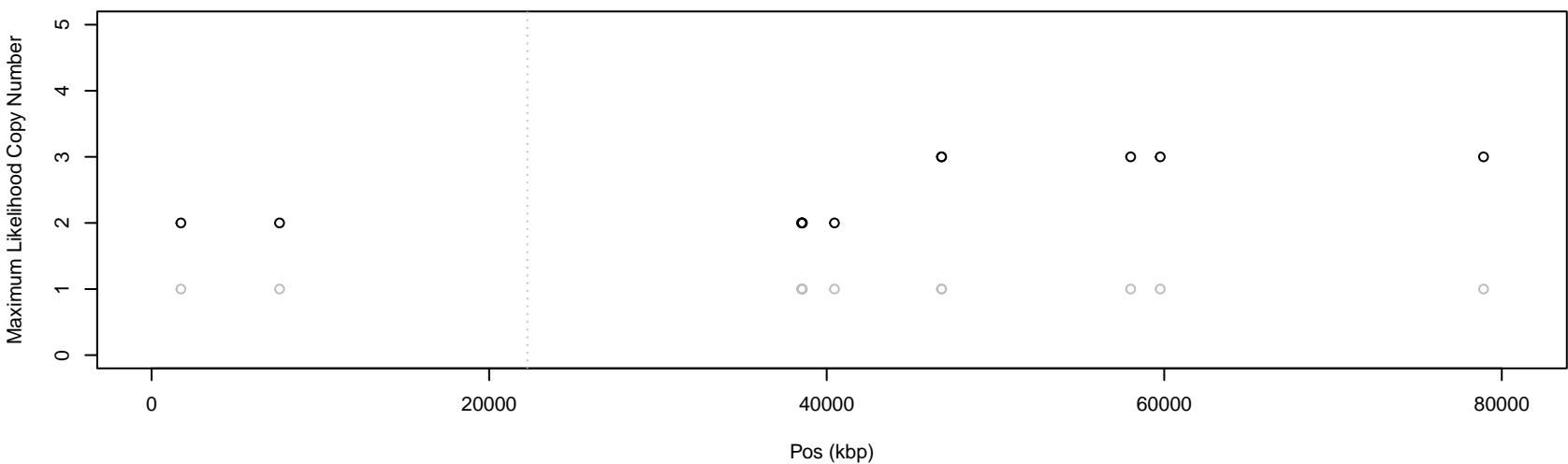
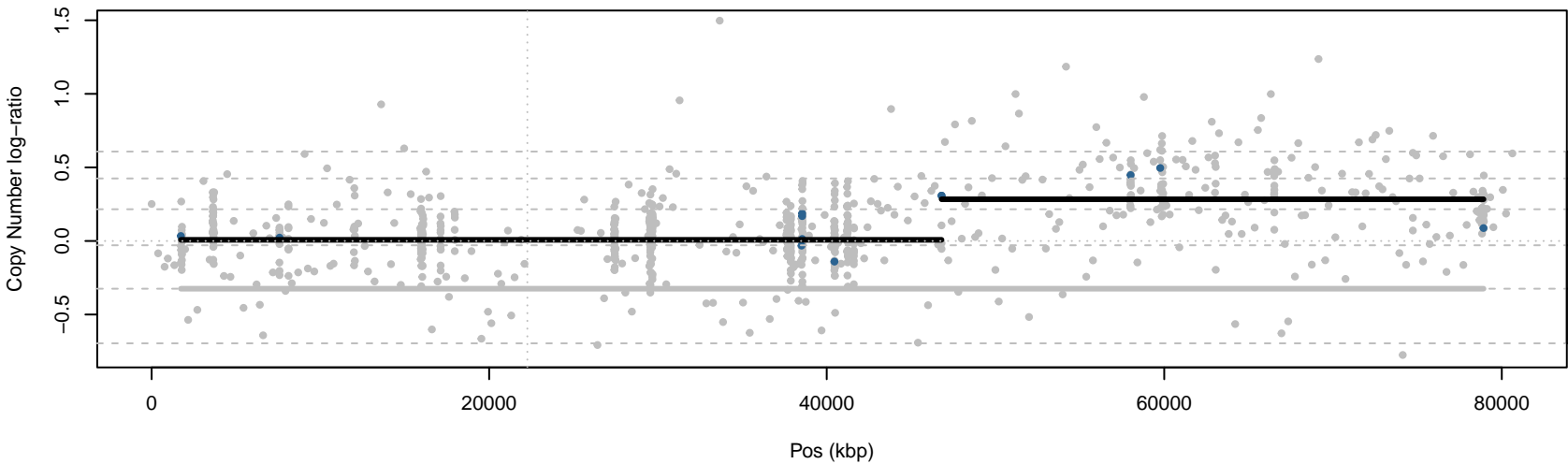
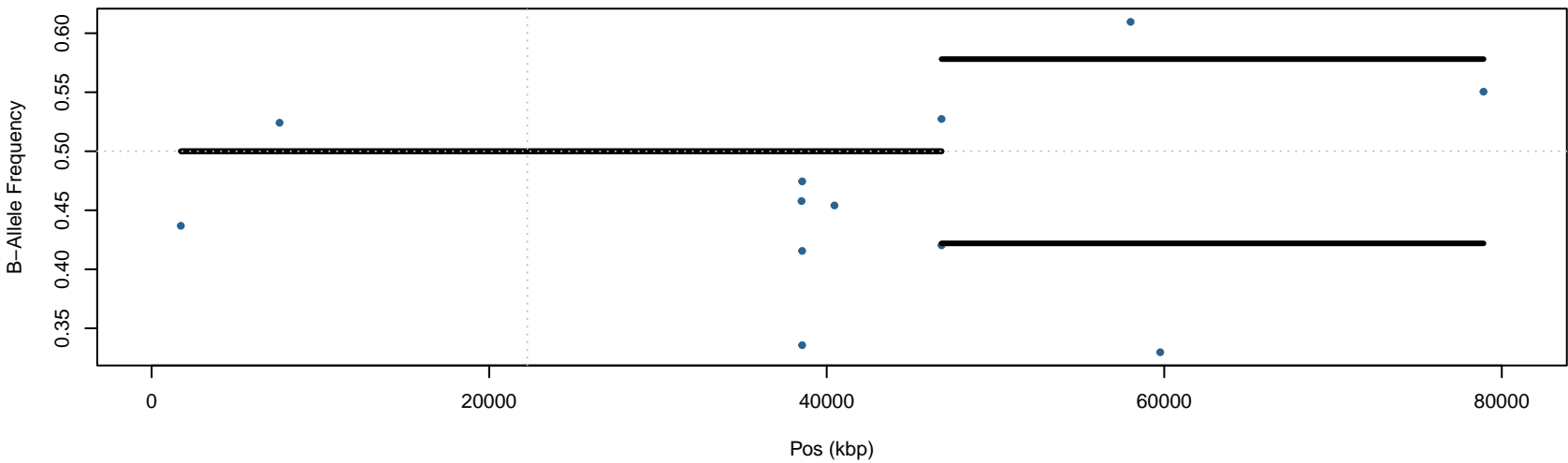
Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 15



Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 16

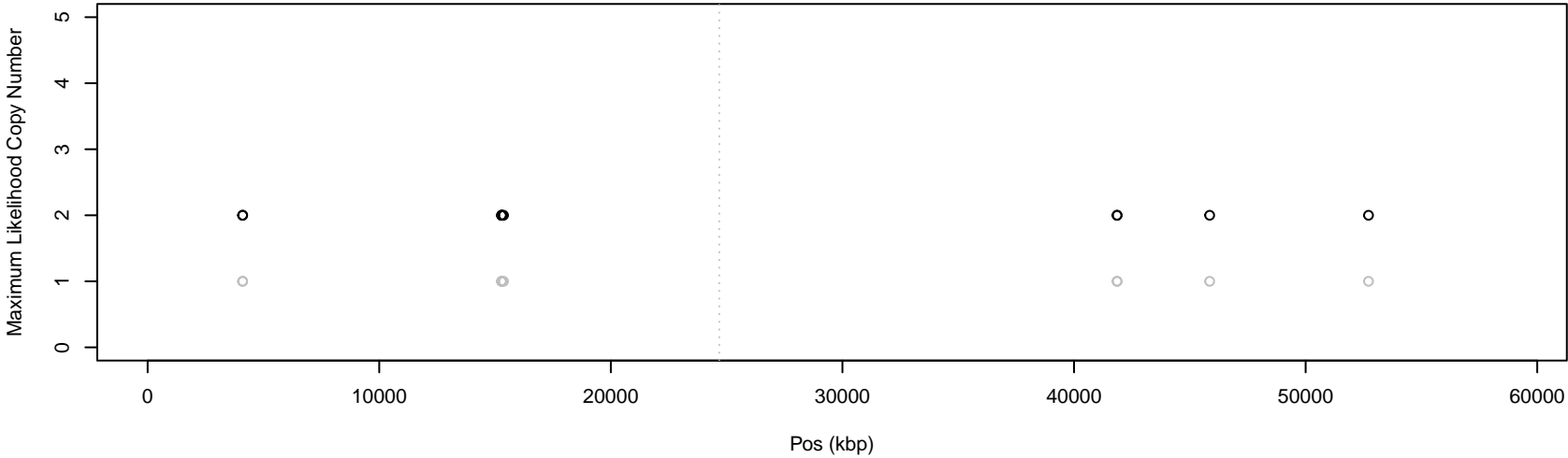
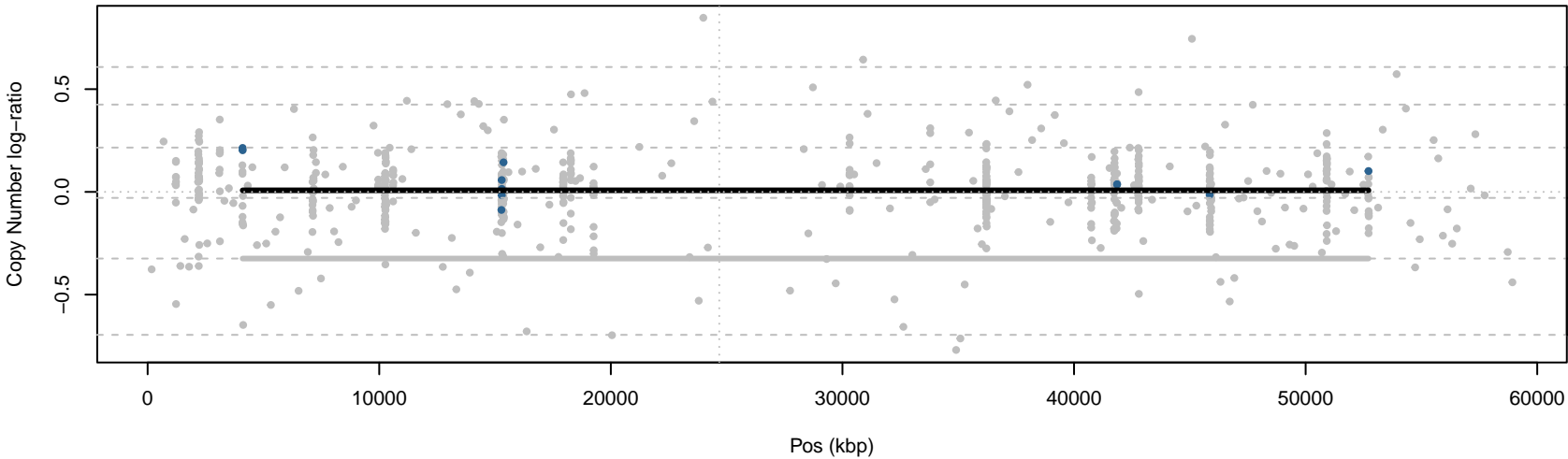
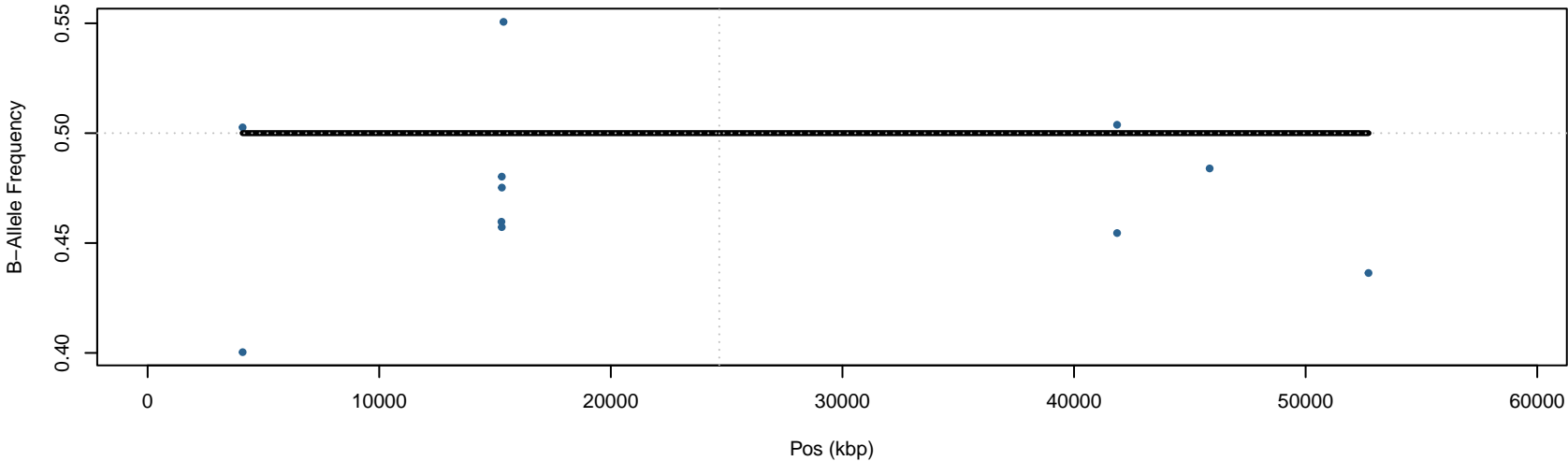


Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 17

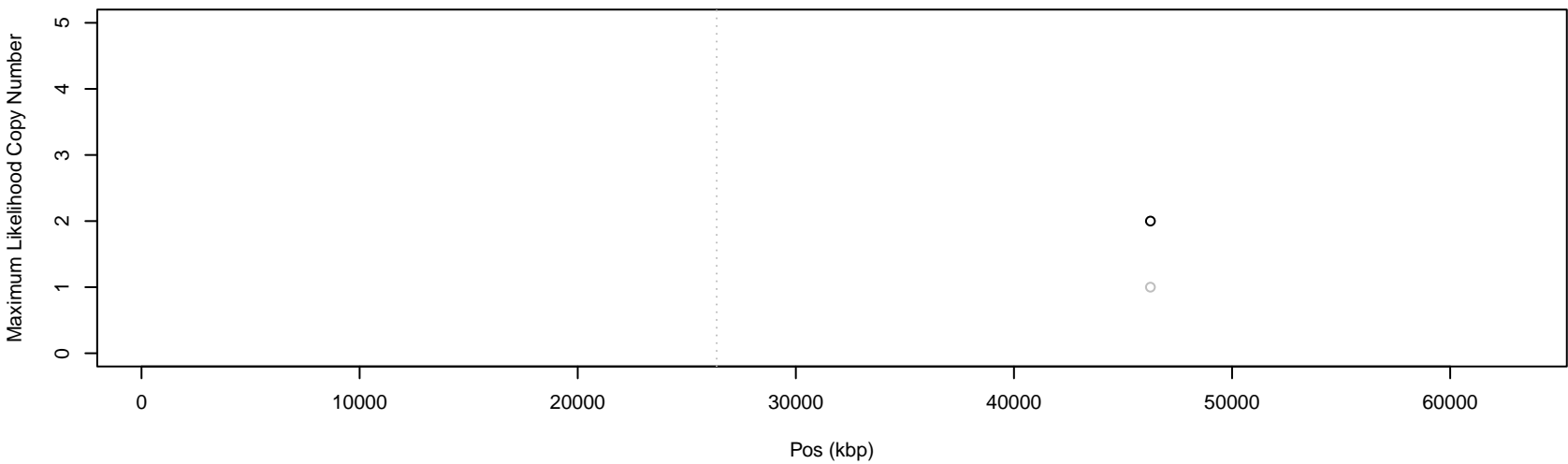
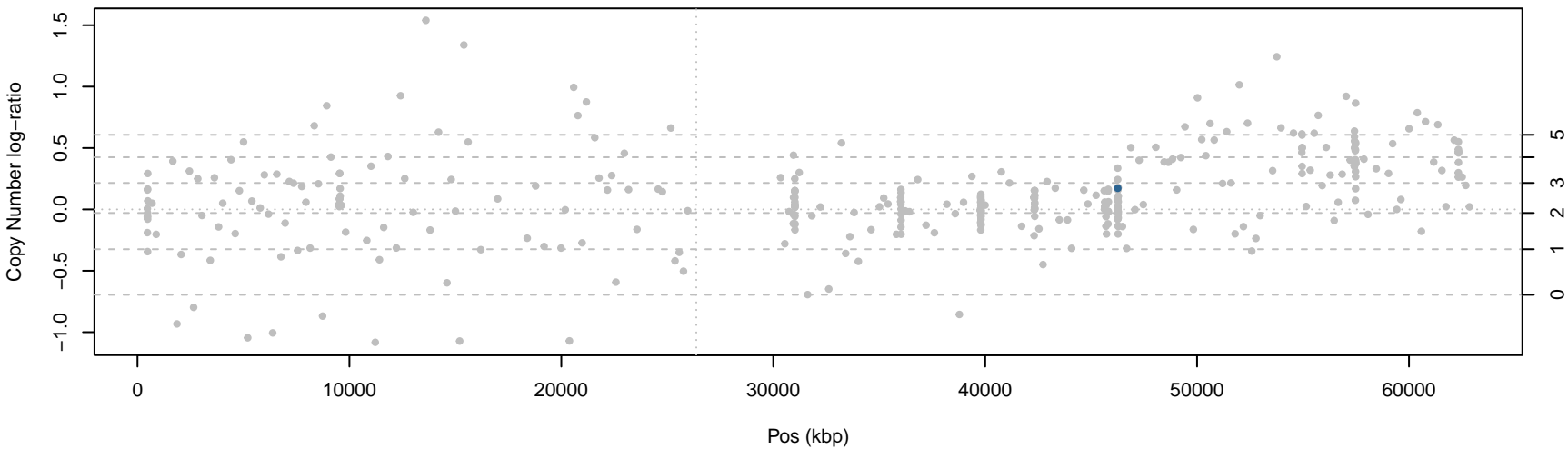
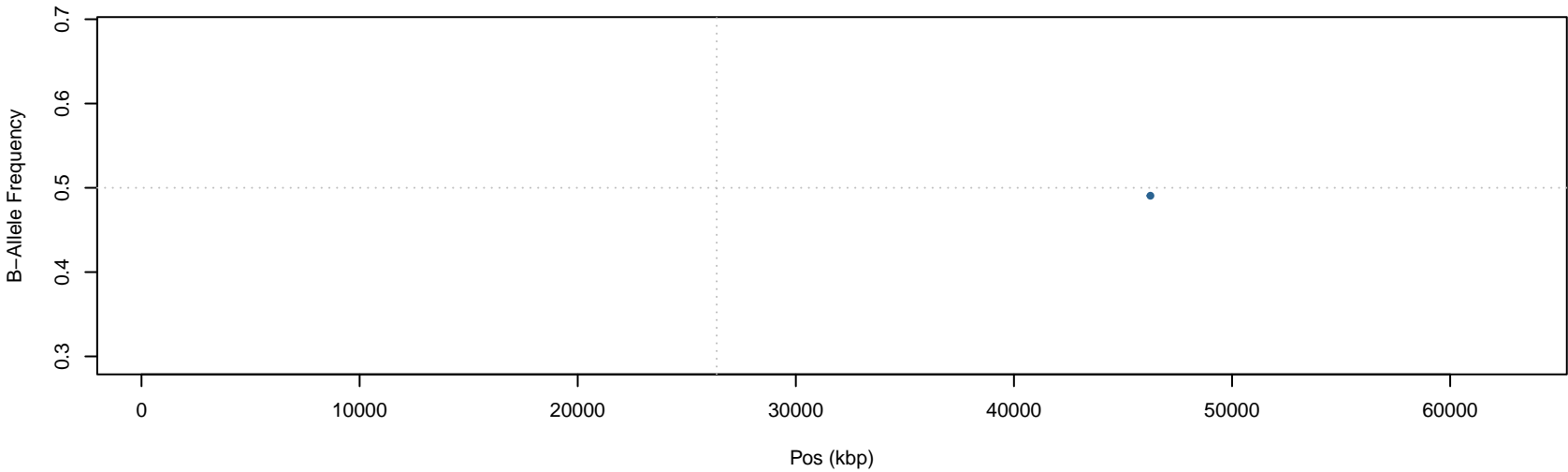




Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 19



Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 20



Purity: 0.37 Tumor ploidy: 2.111 SNV log-likelihood: -63.57 GoF: 87% Mean coverage: 527;500 Chromosome: 22

