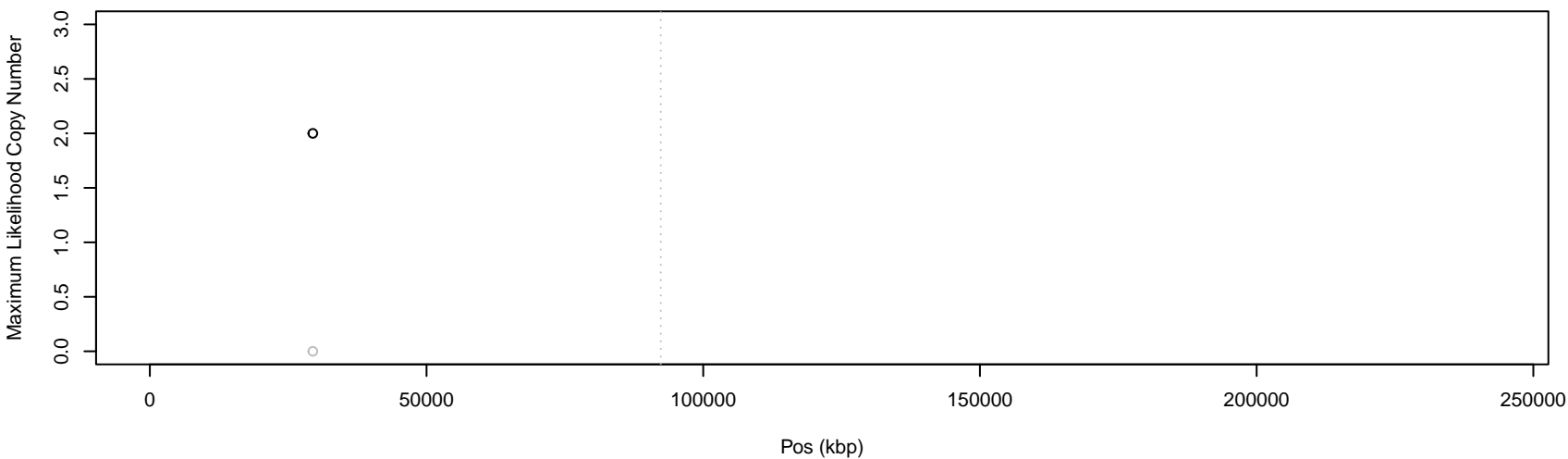
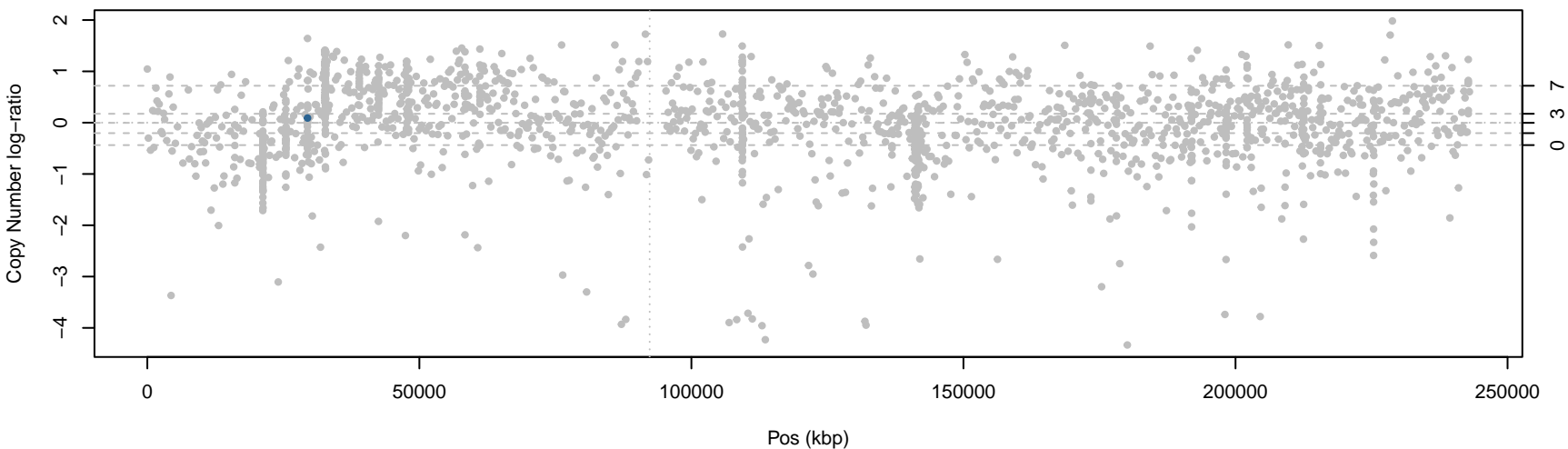
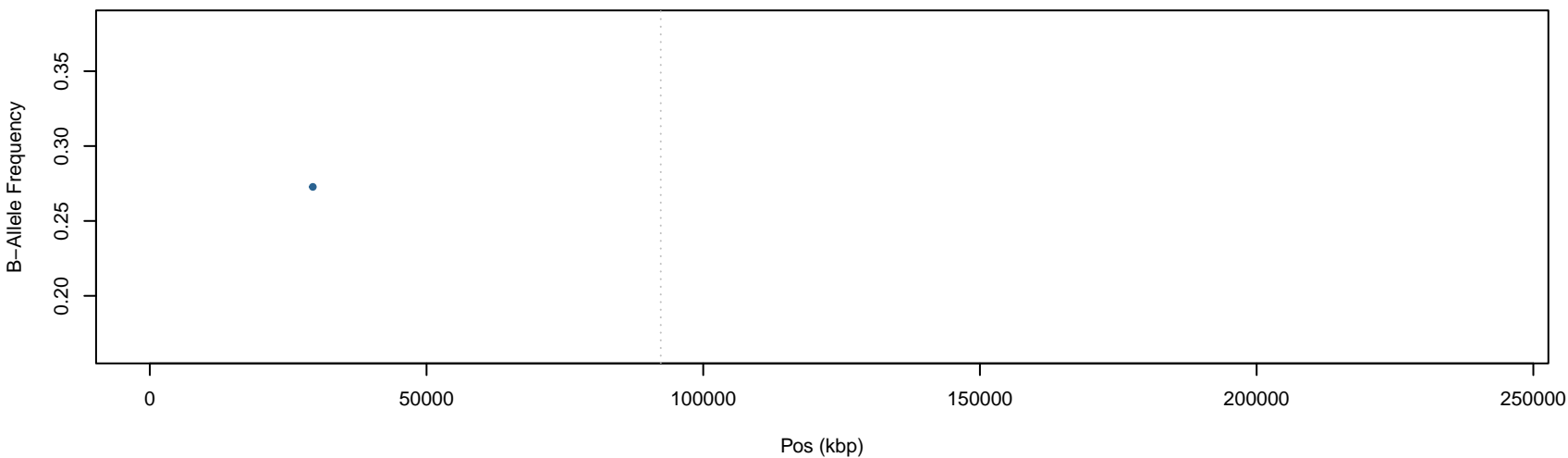
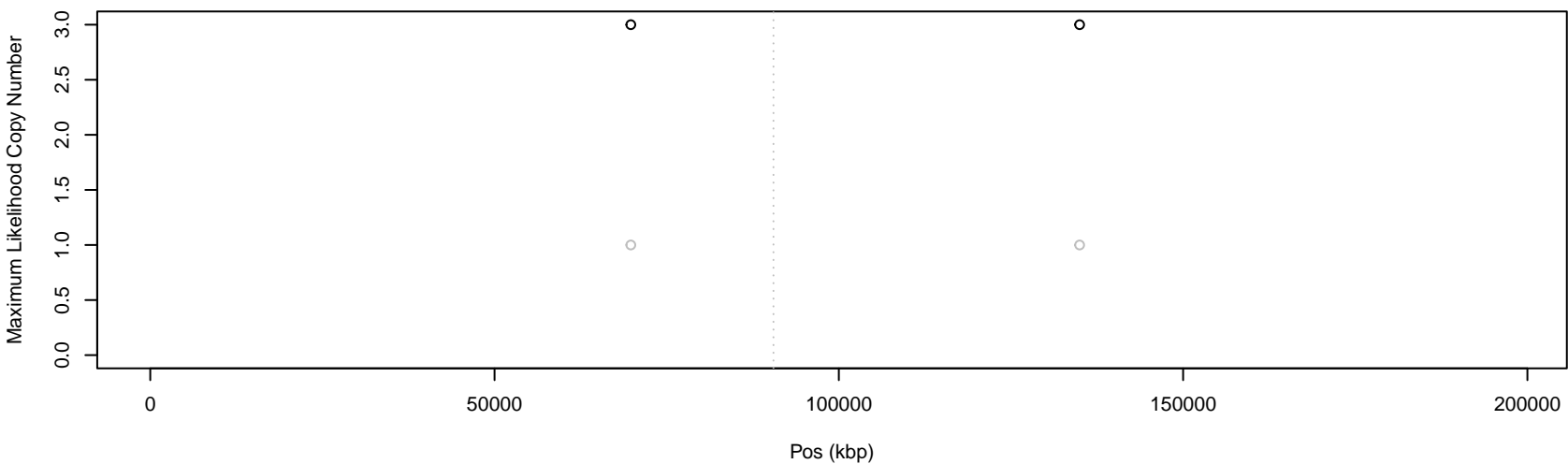
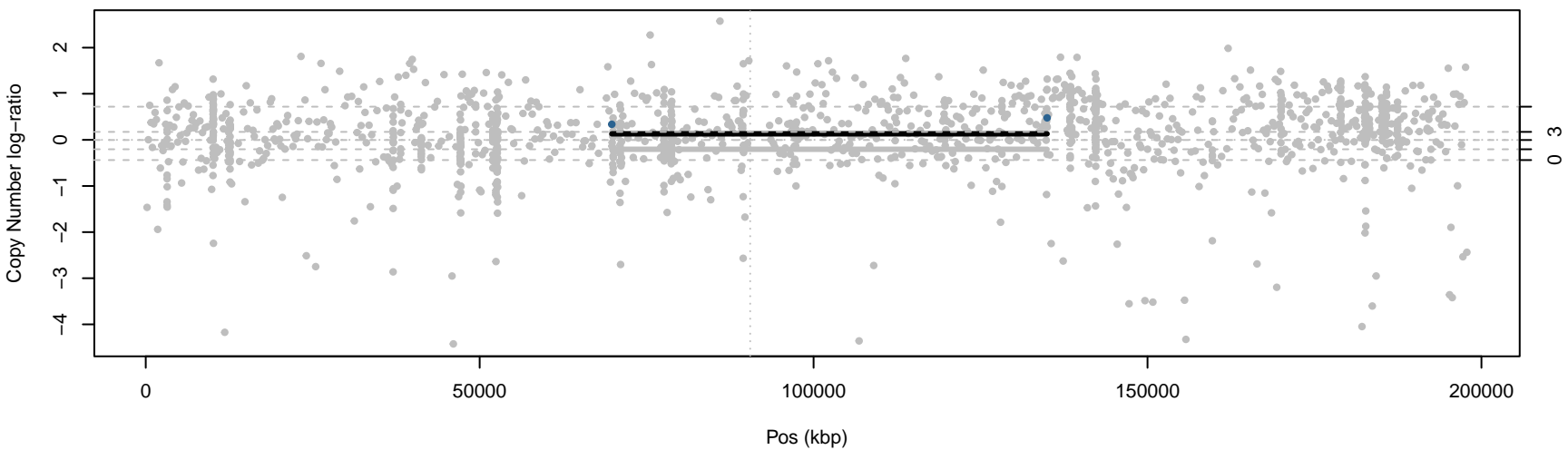
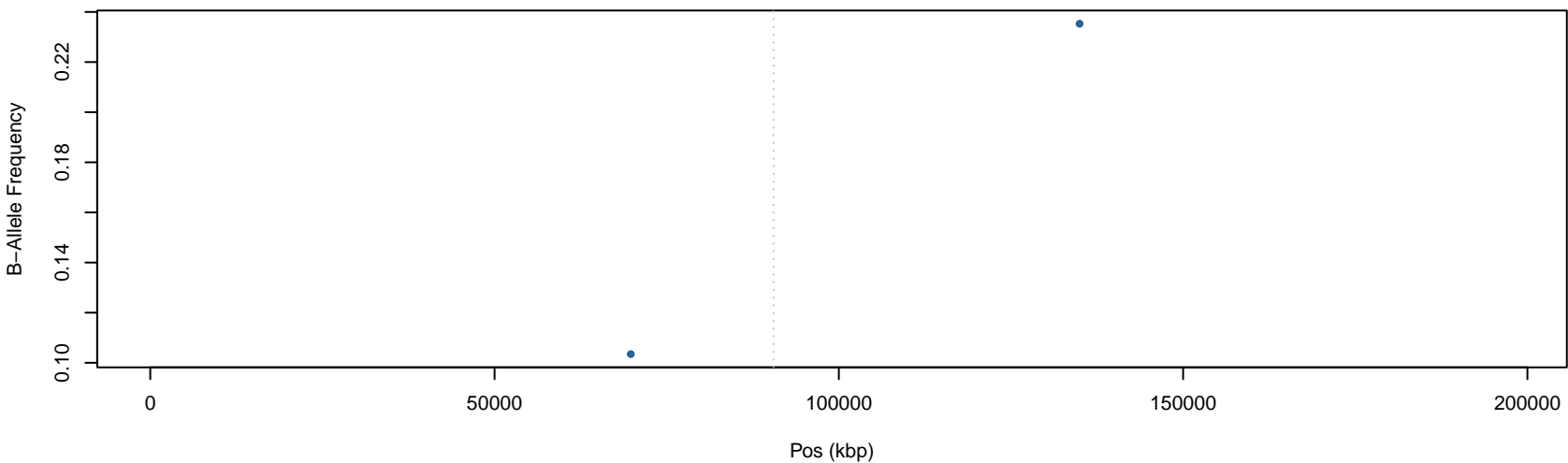


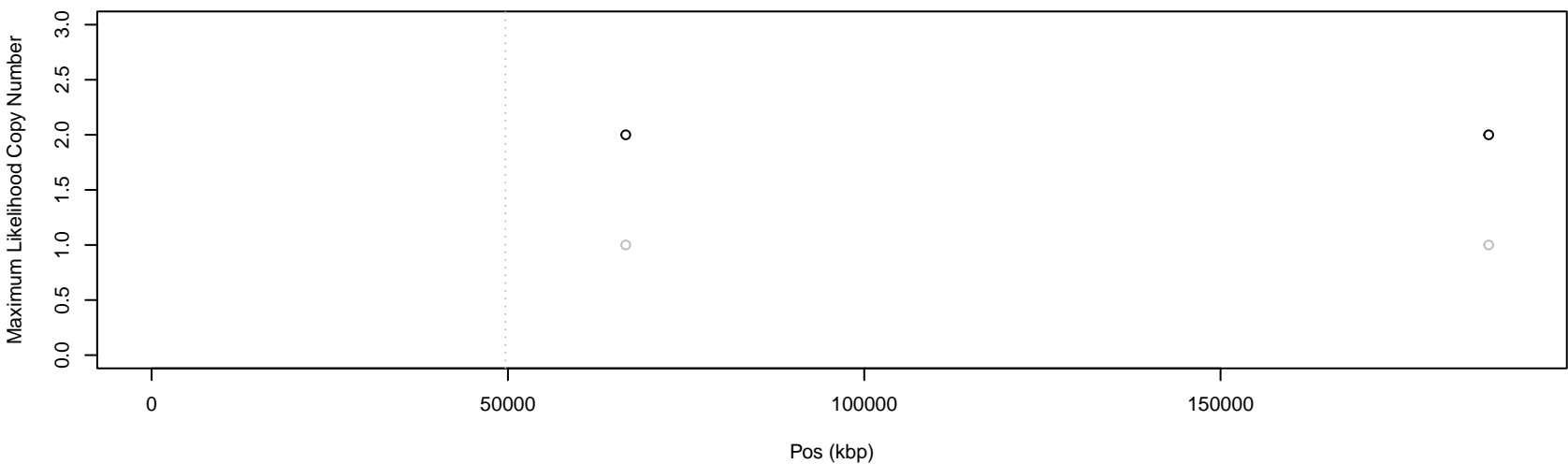
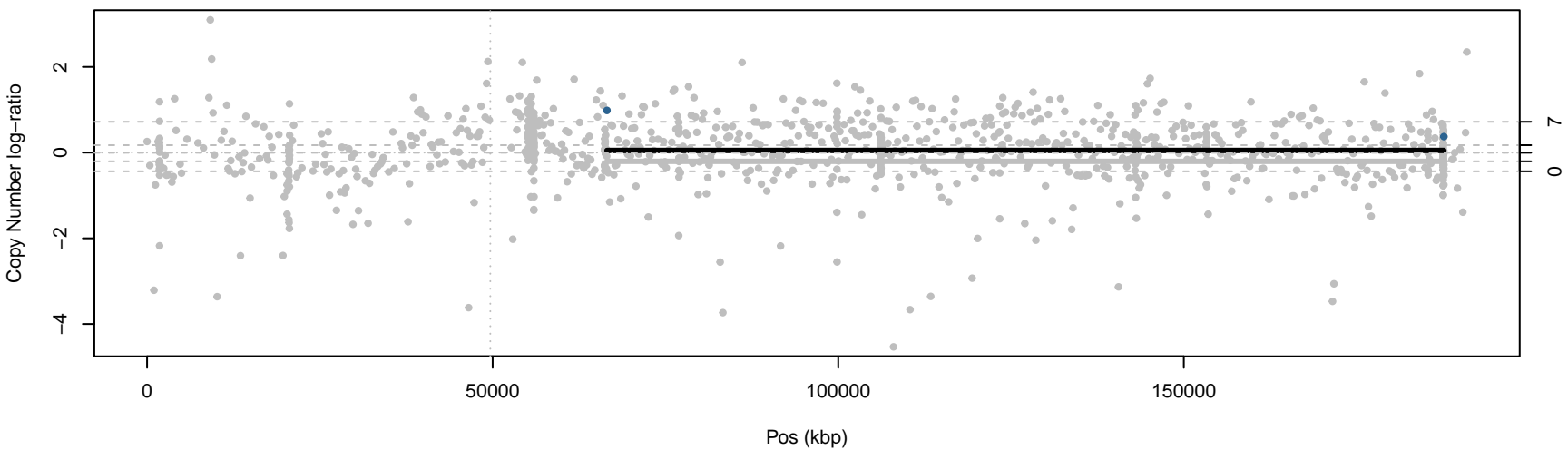
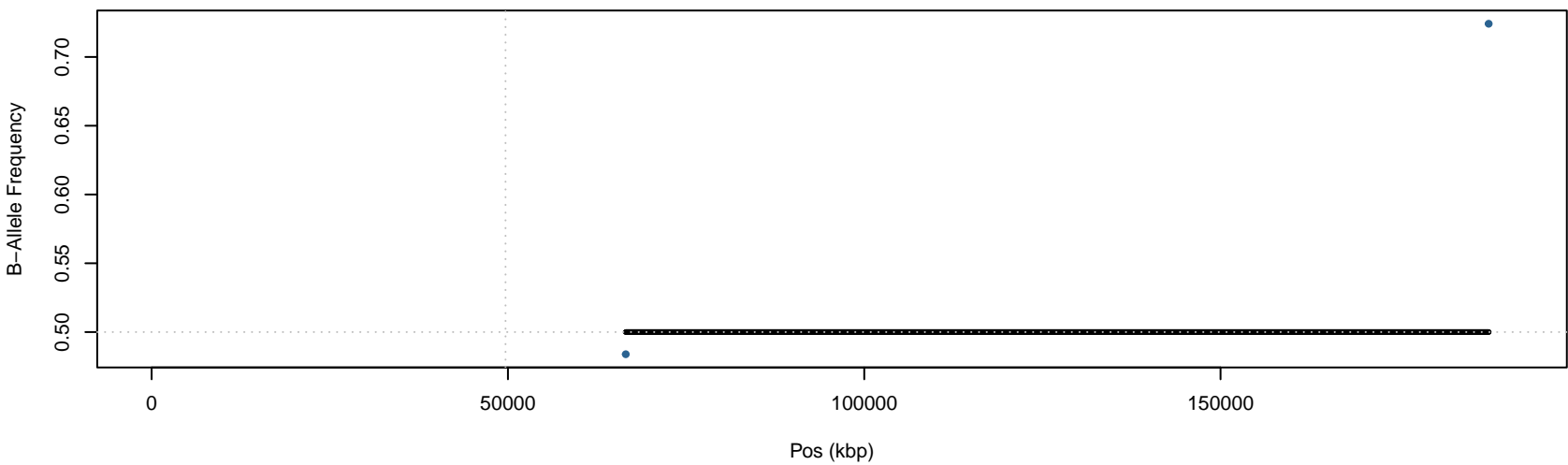
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 2



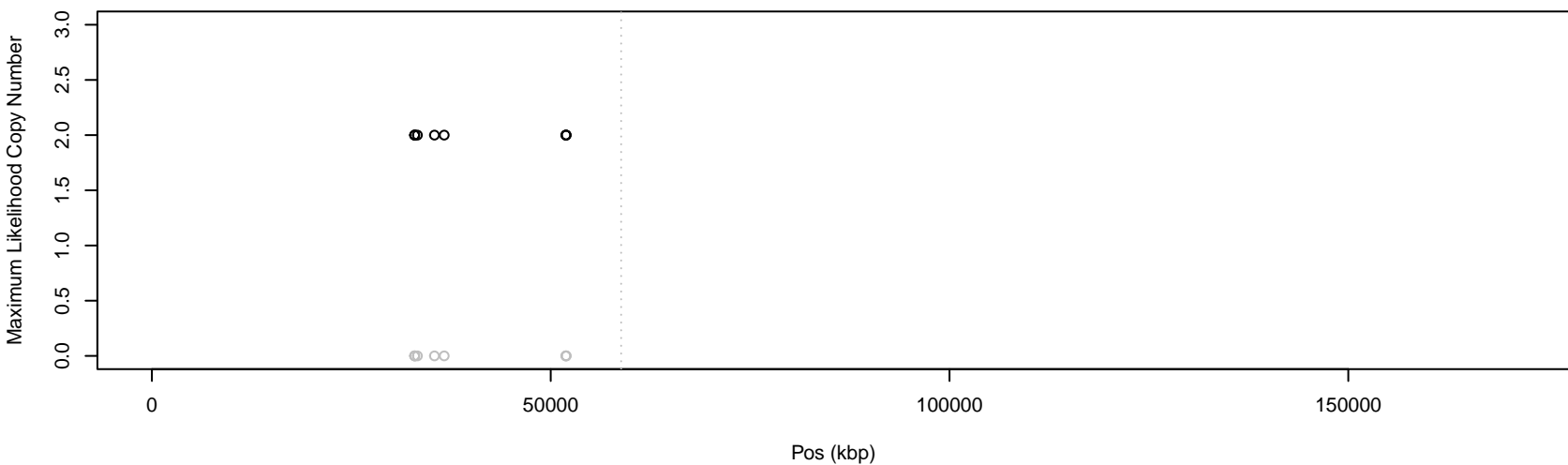
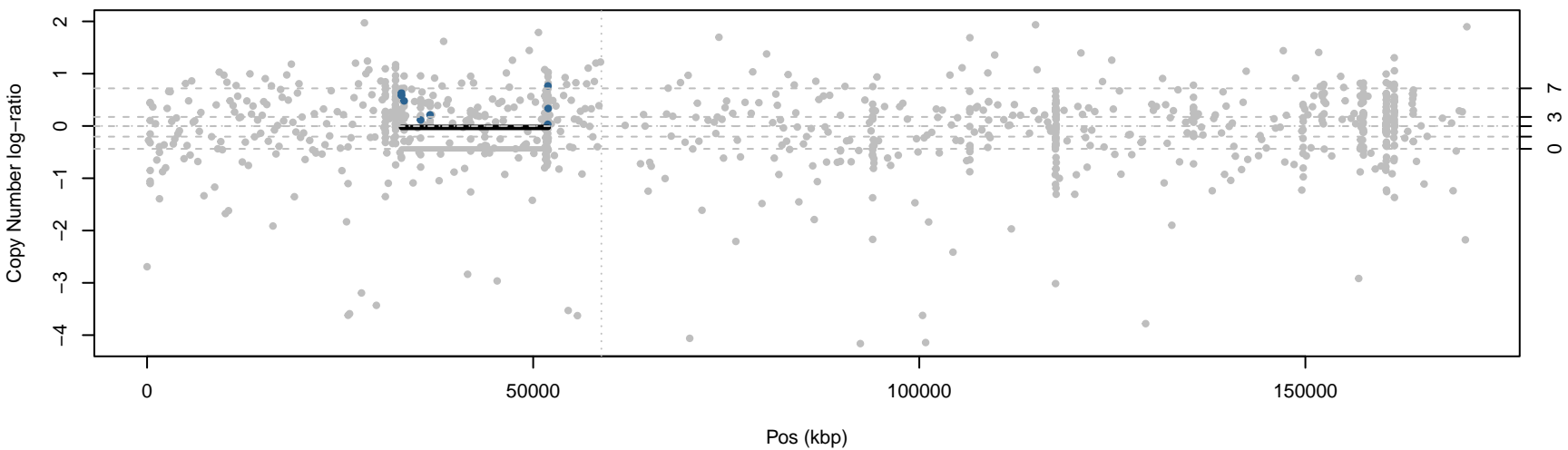
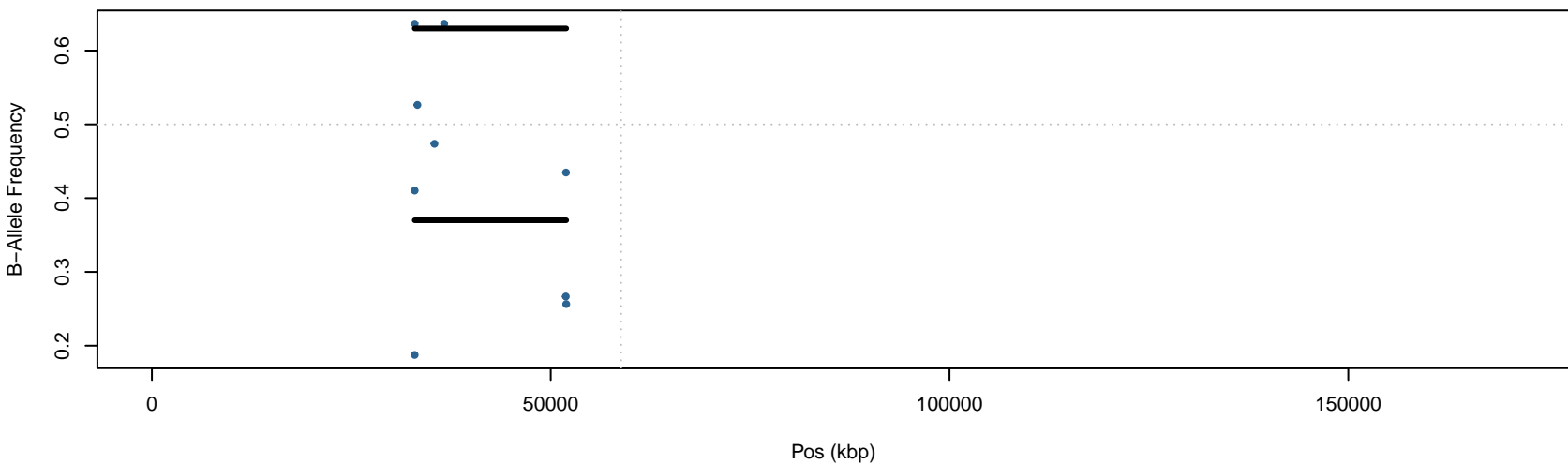
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 3



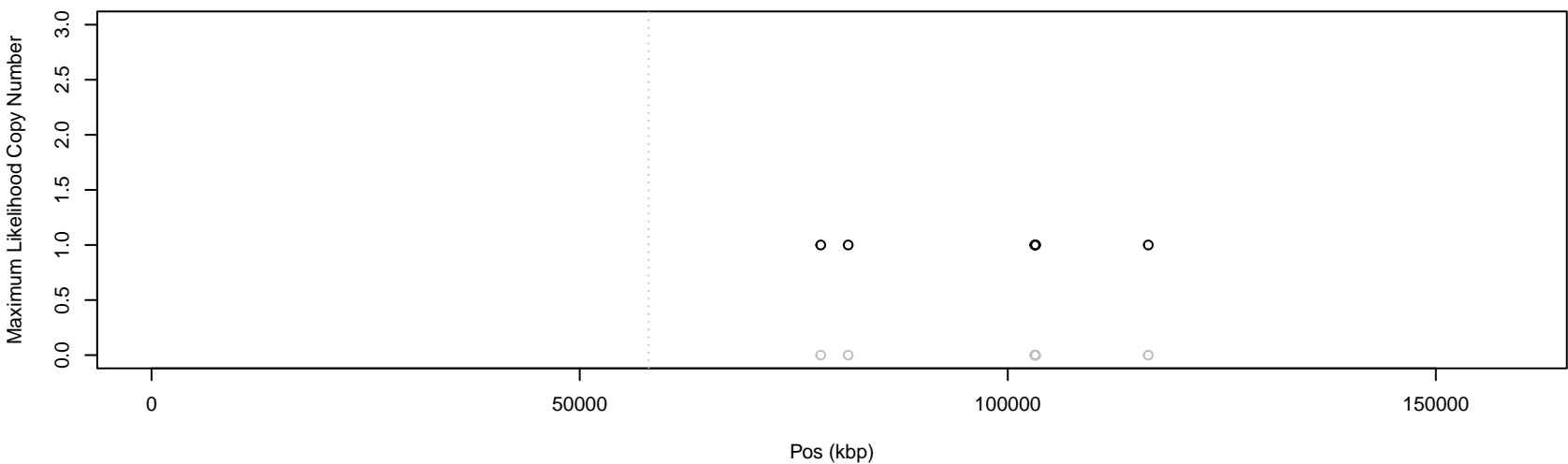
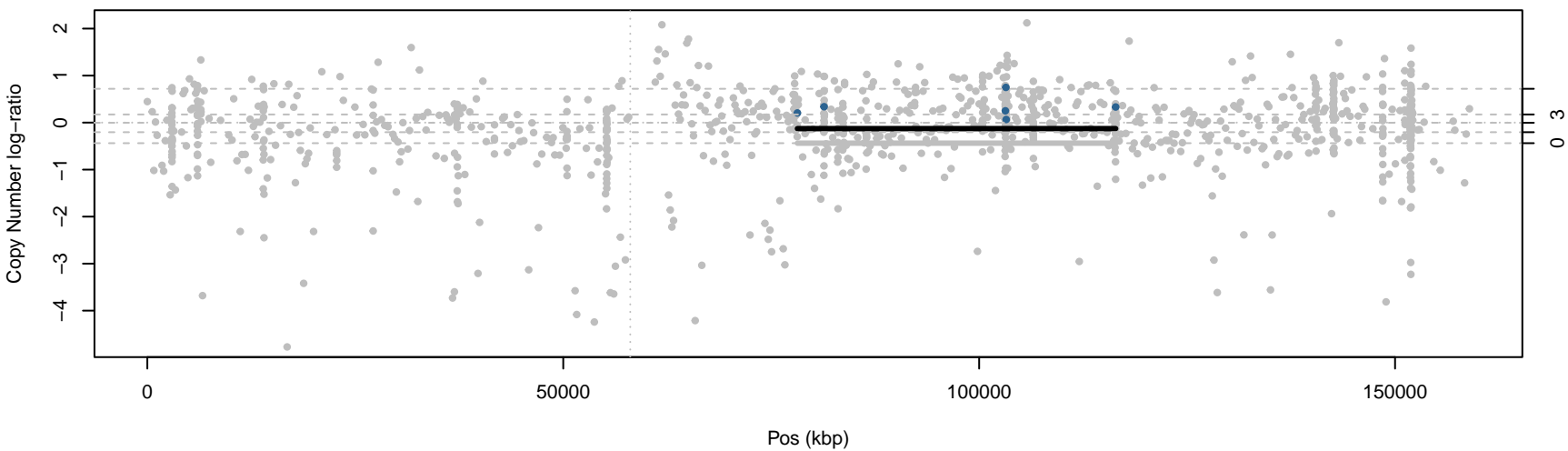
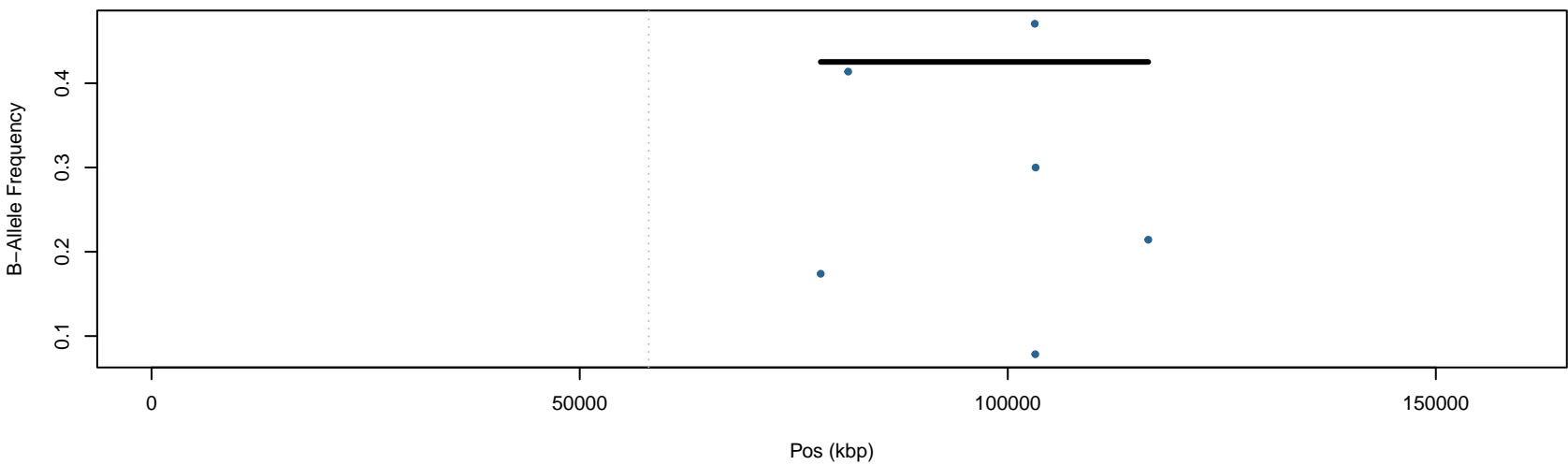
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 4



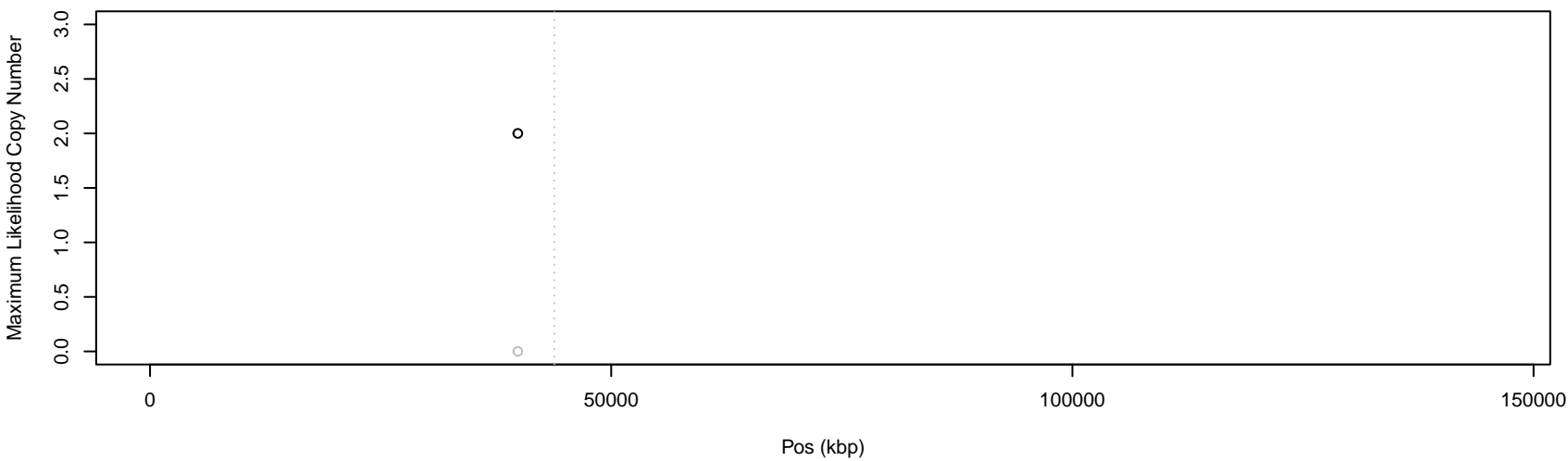
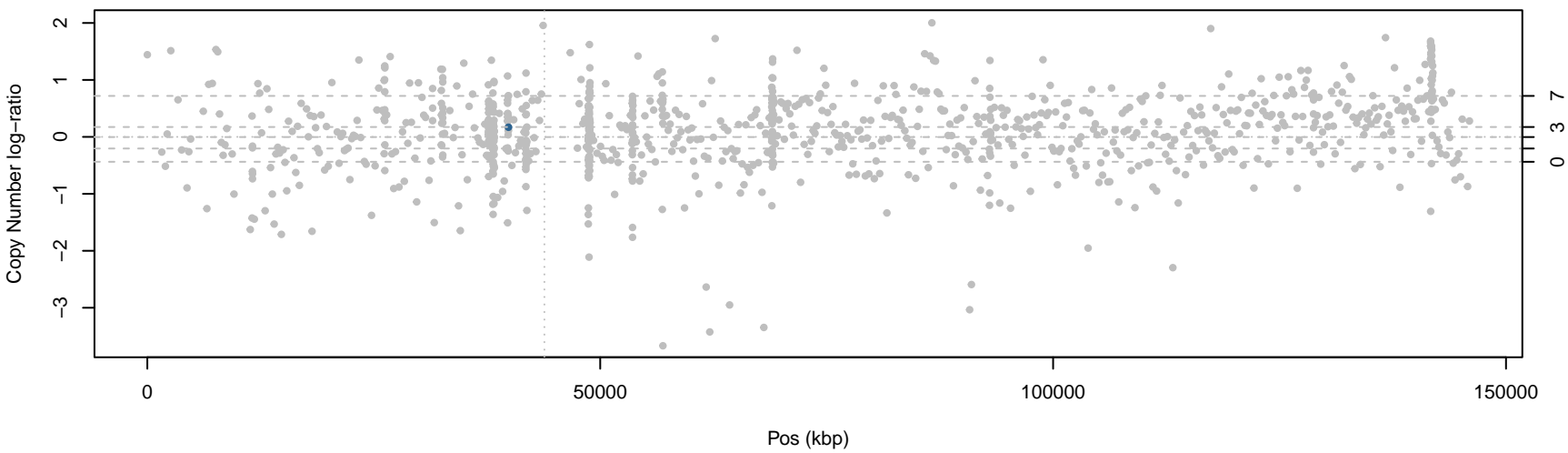
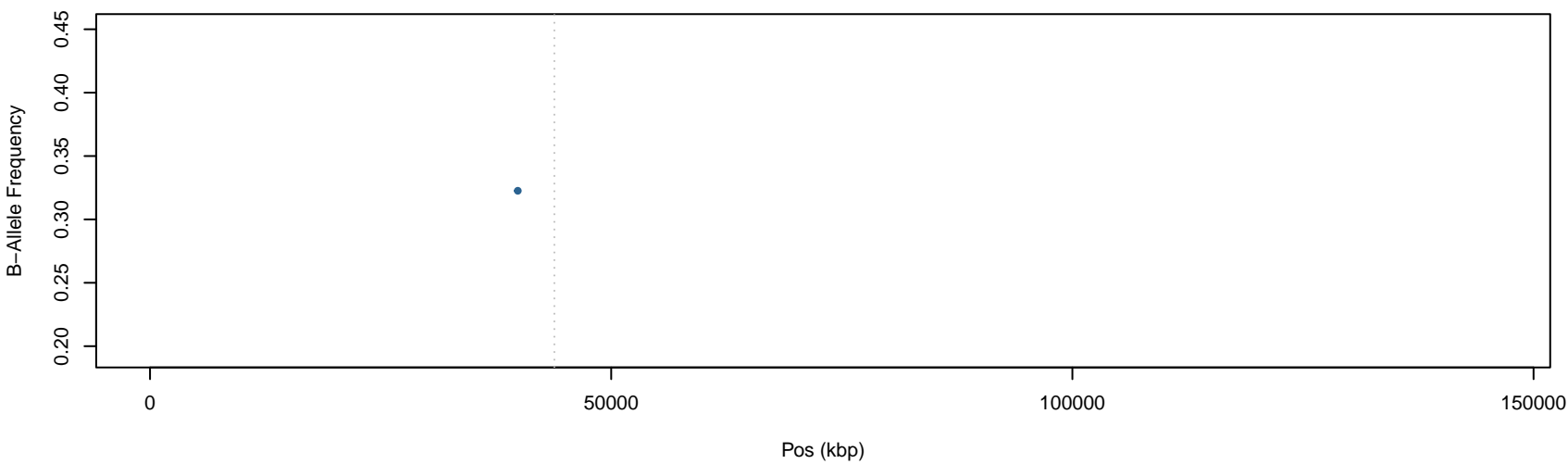
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 6



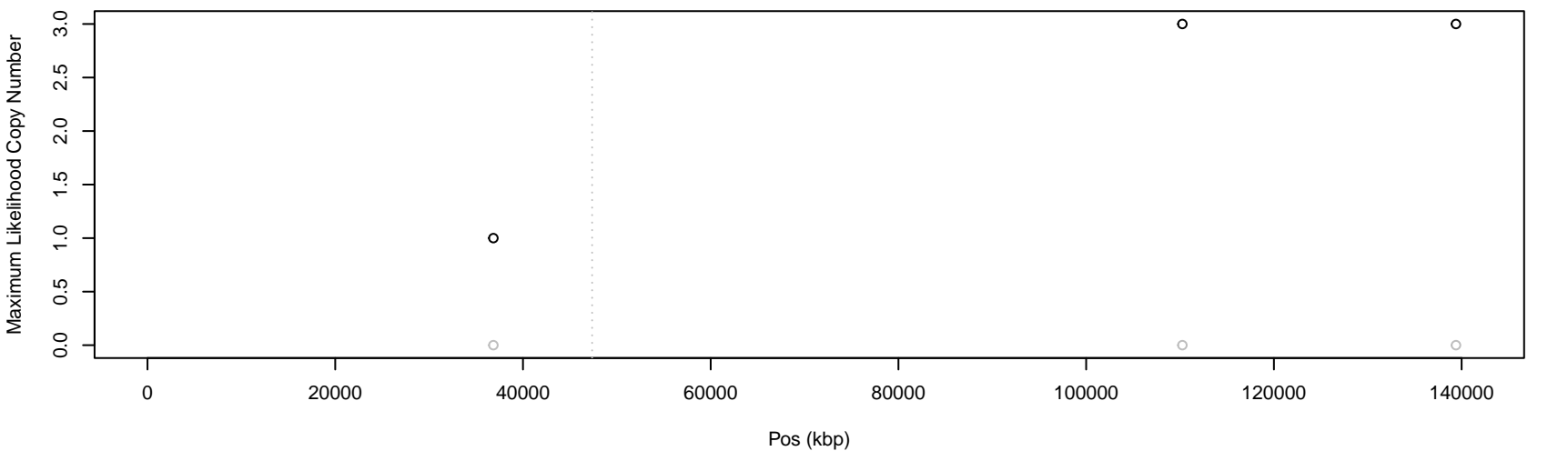
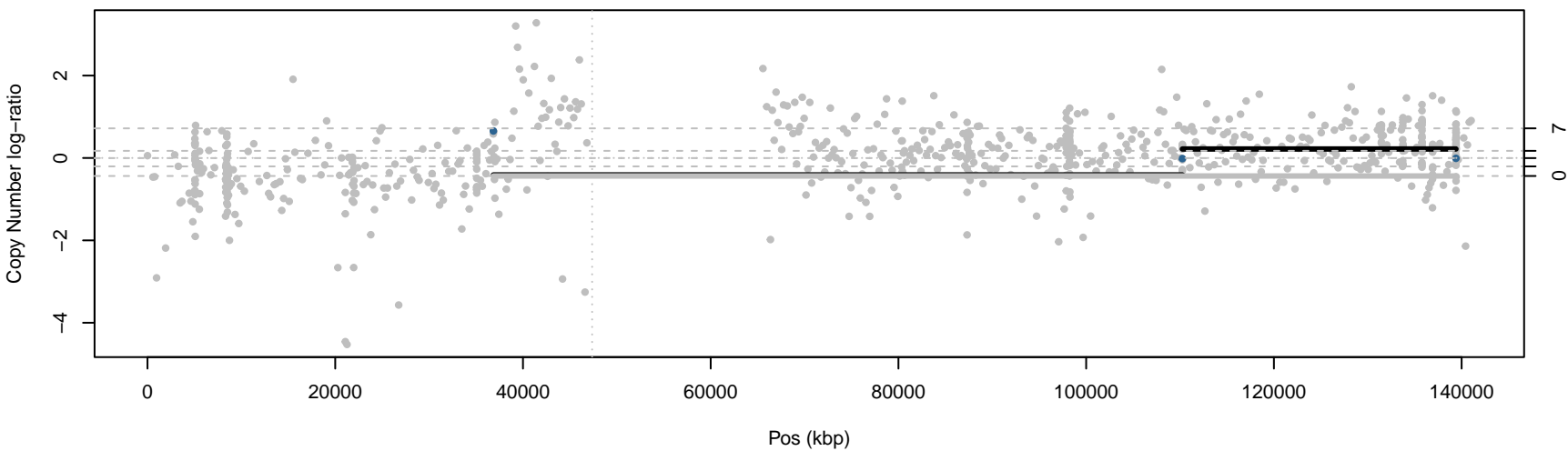
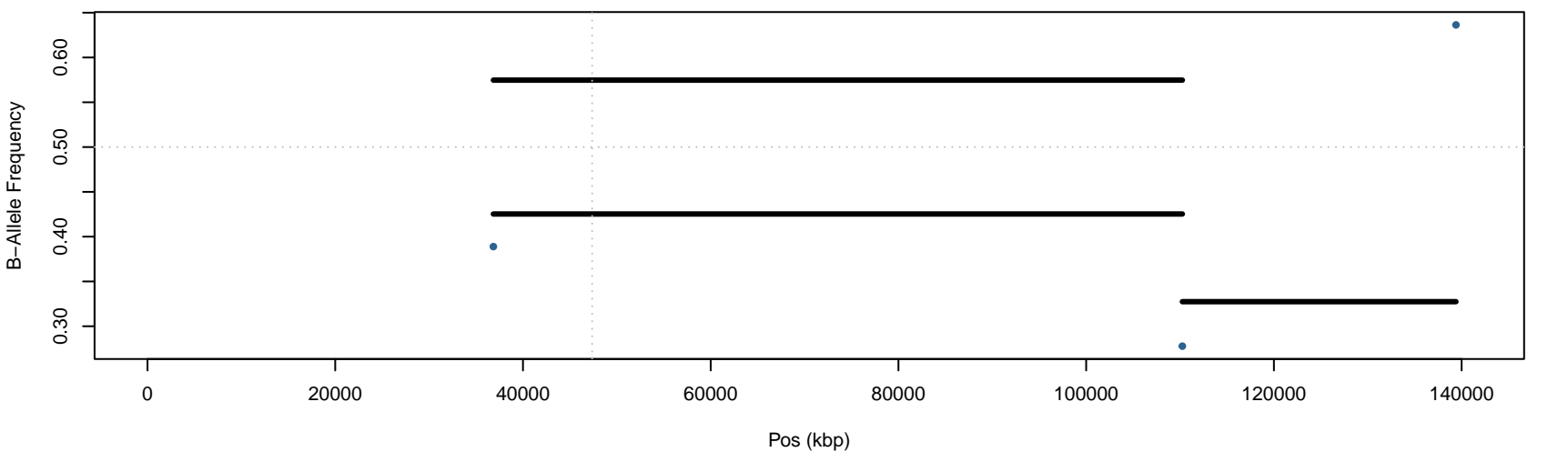
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 7



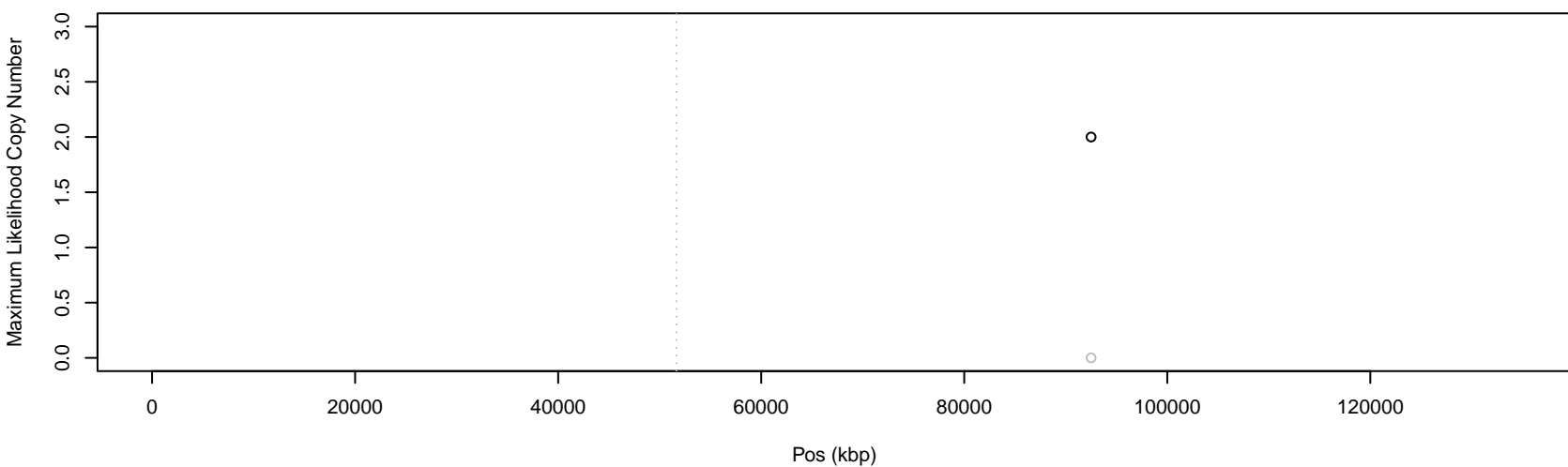
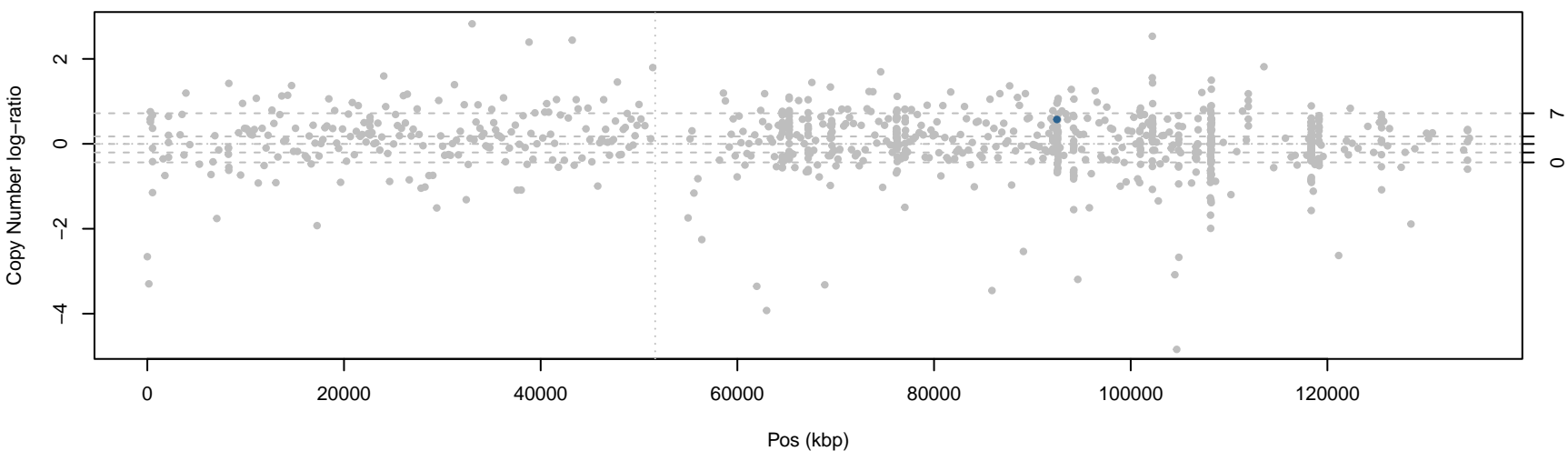
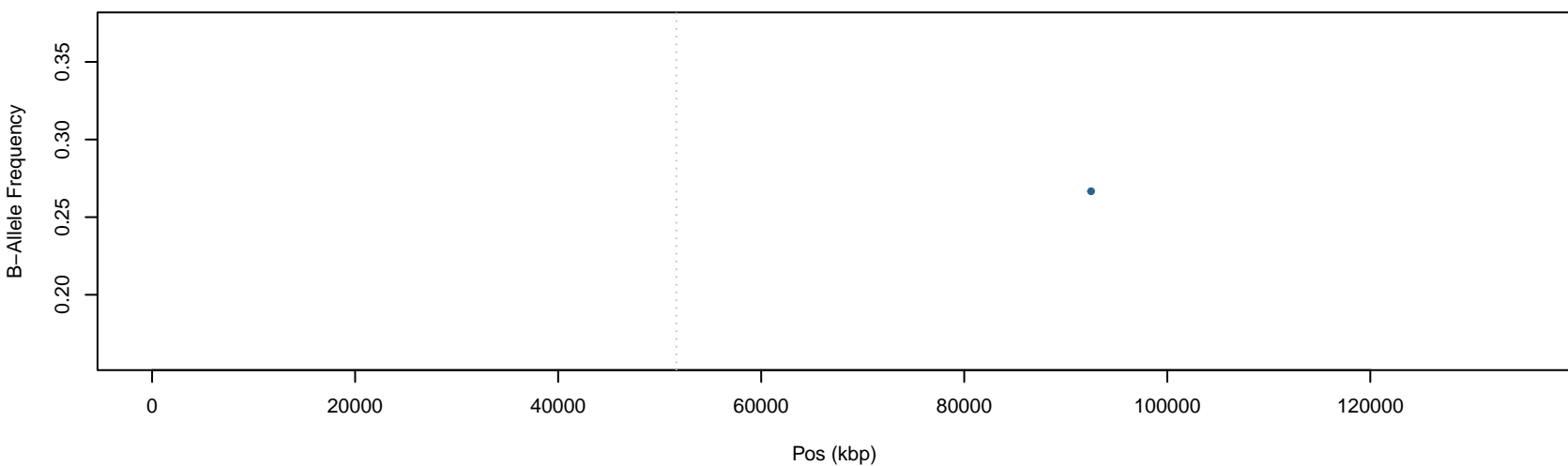
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 8



Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 9

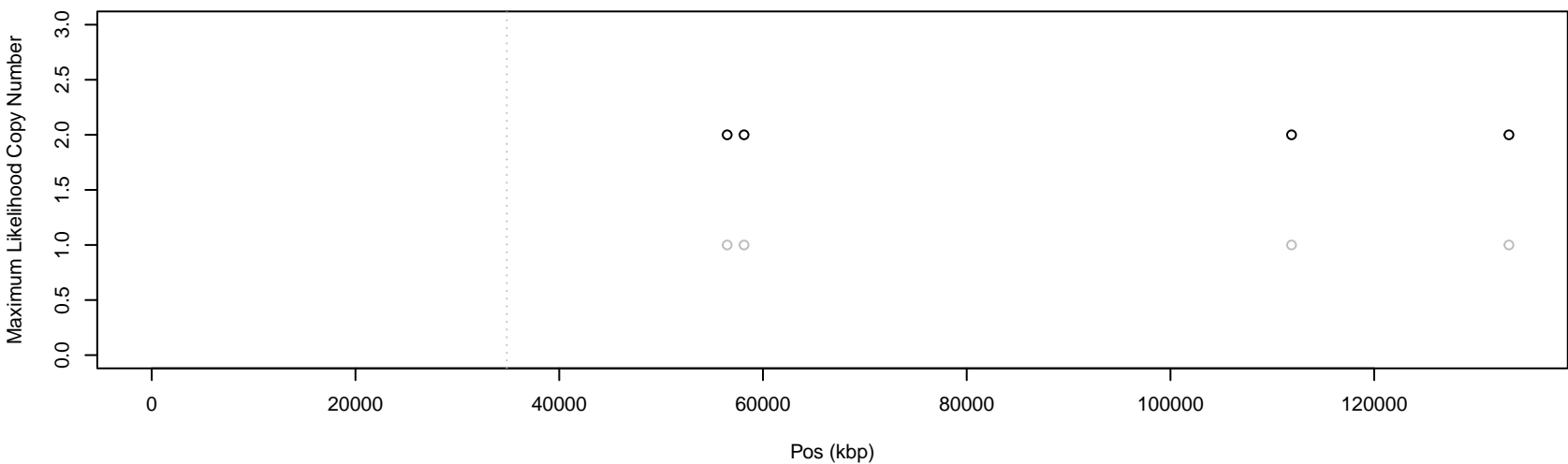
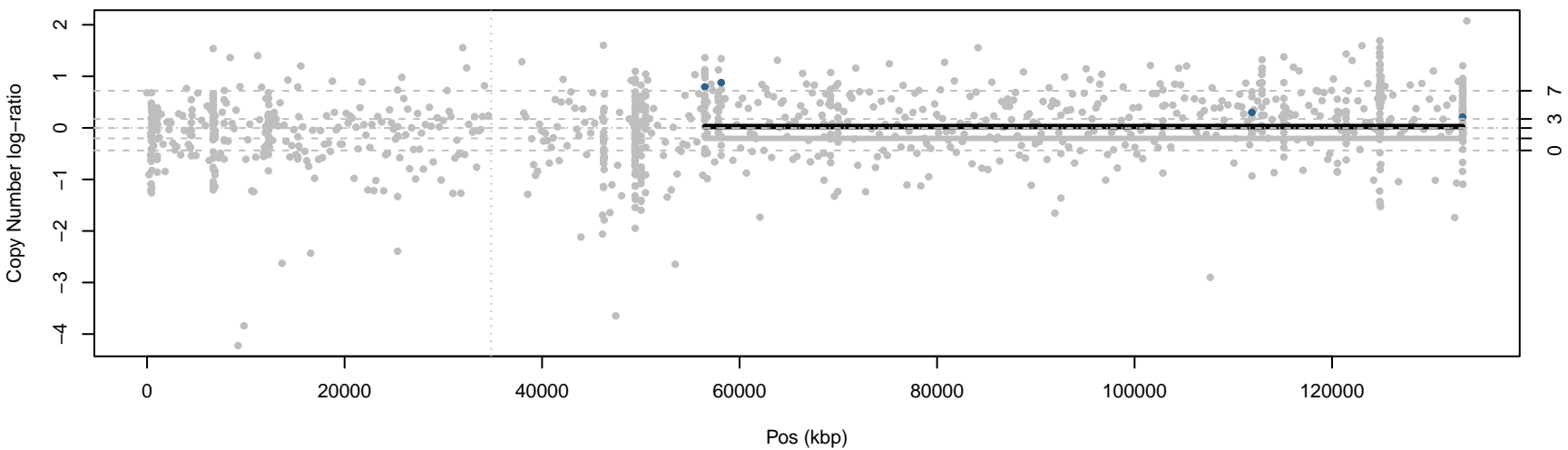
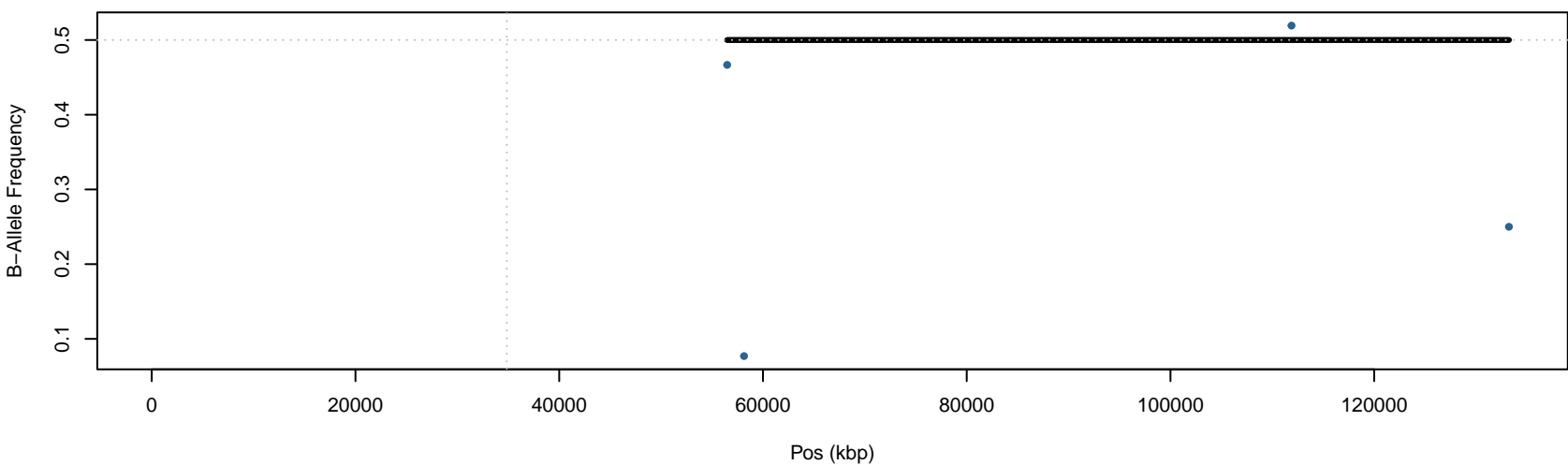


Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 11

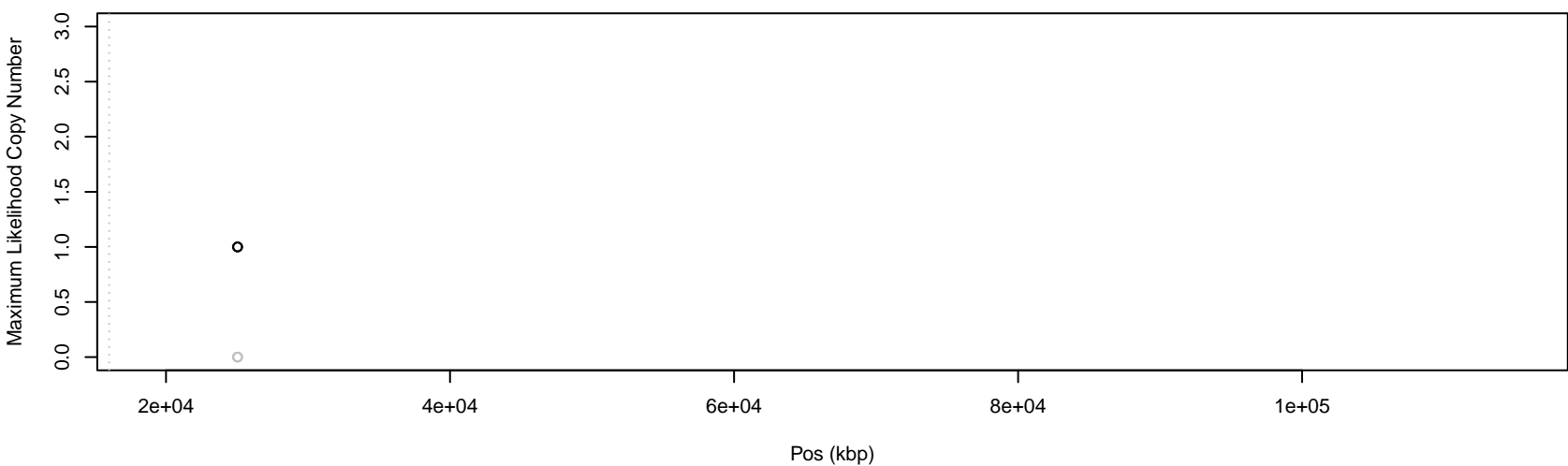
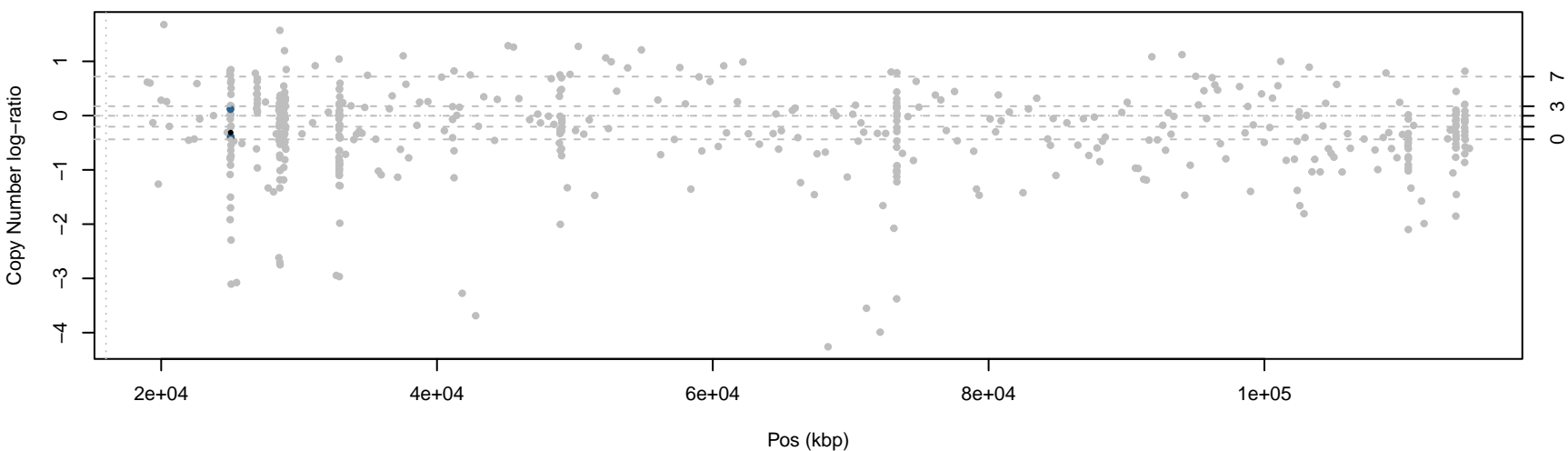
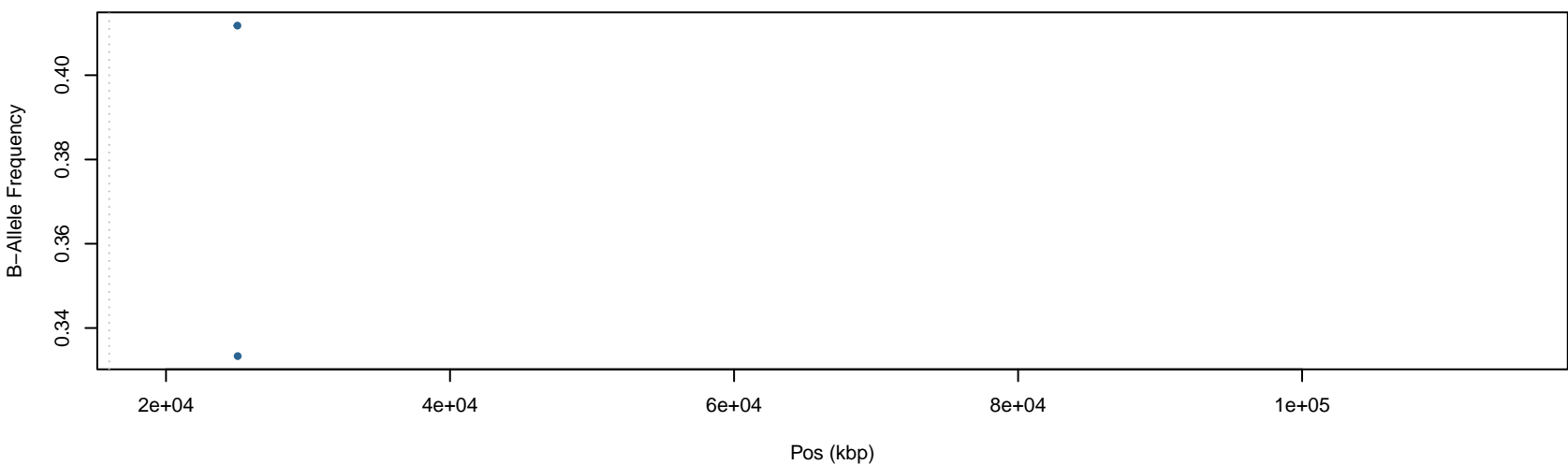




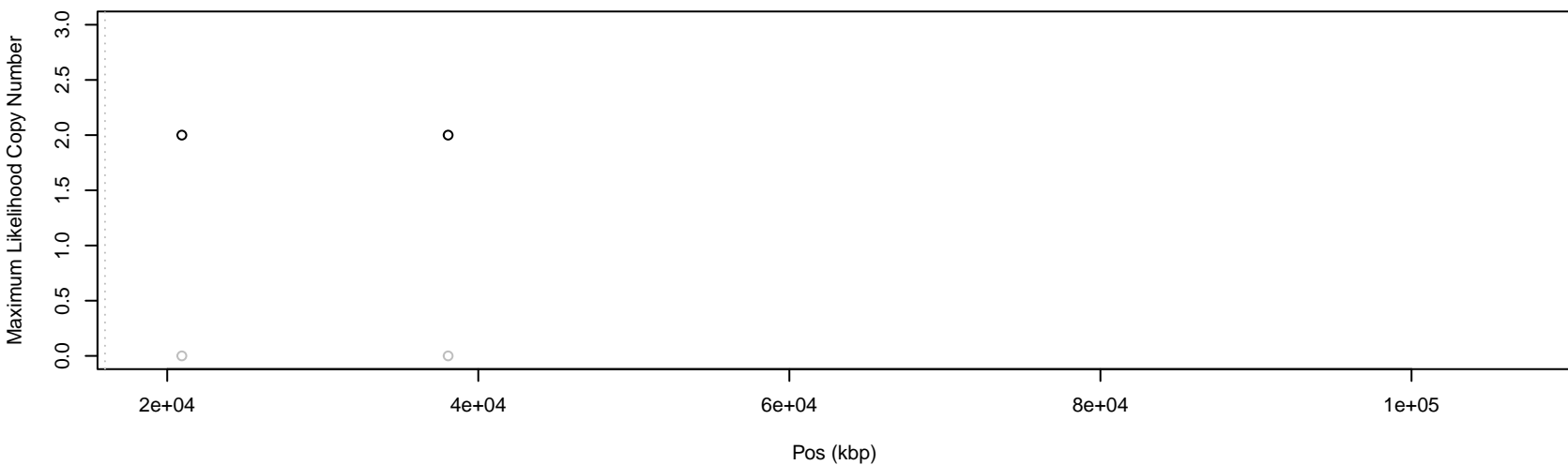
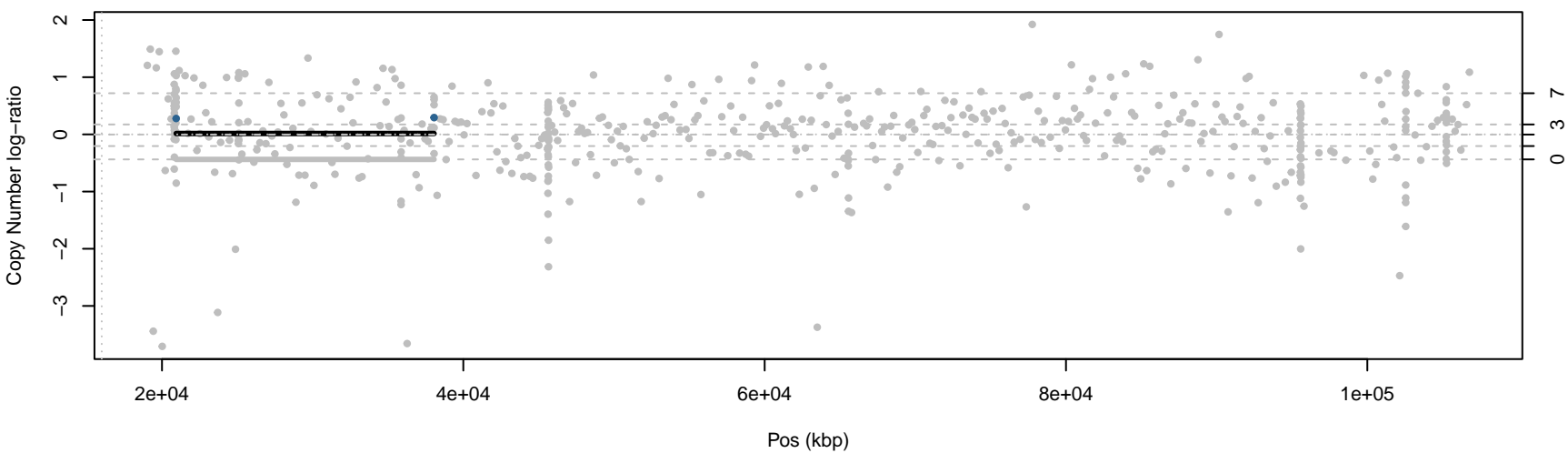
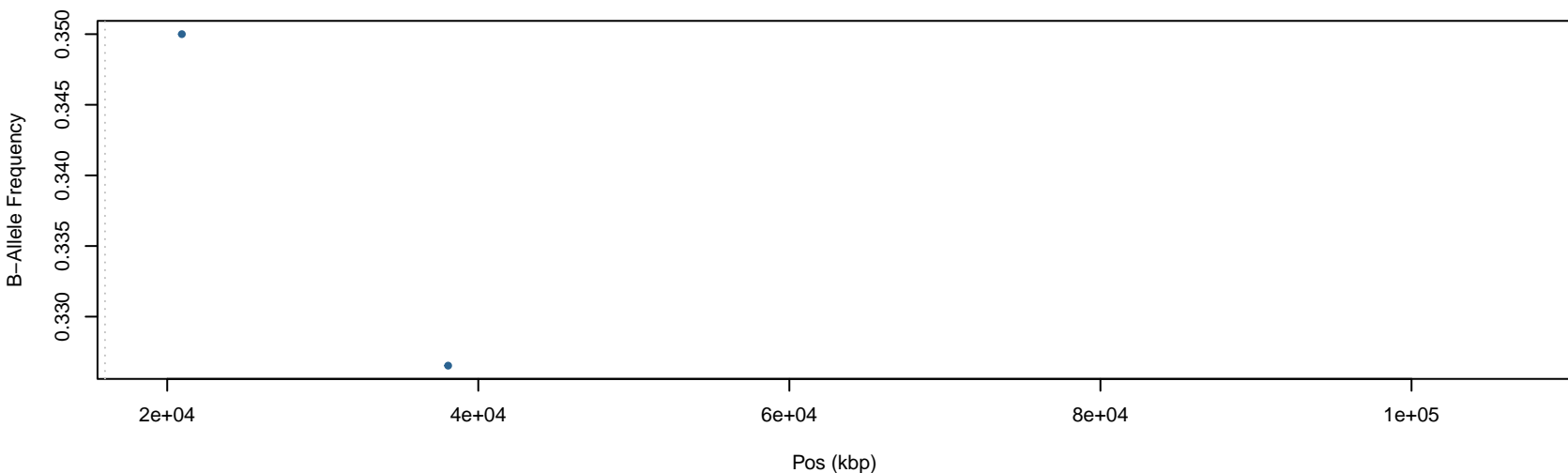
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 12



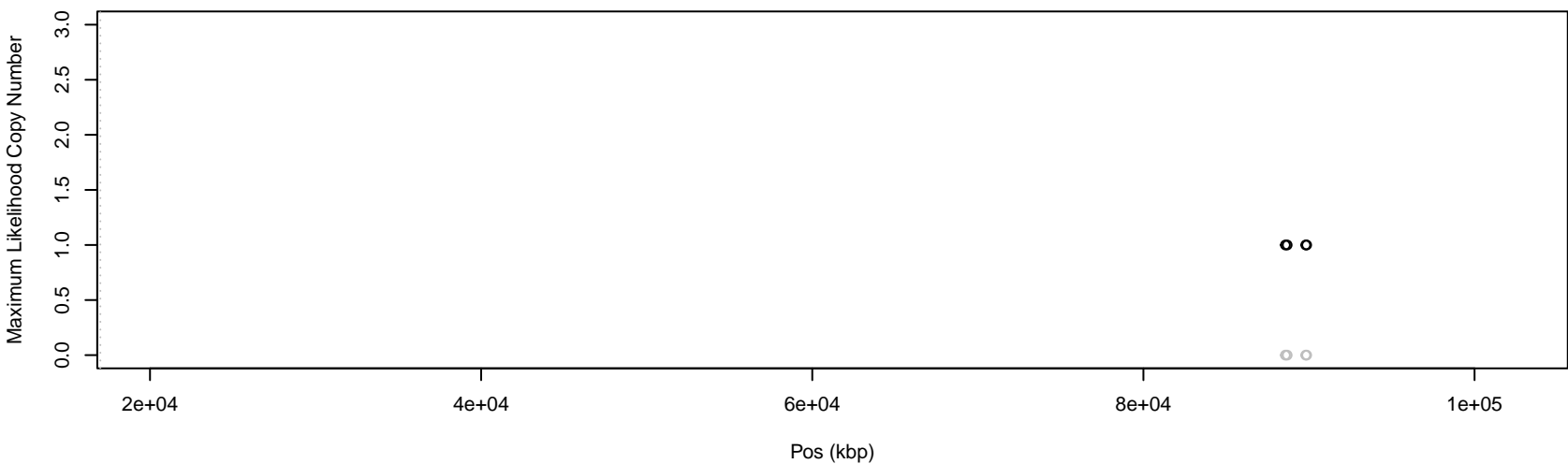
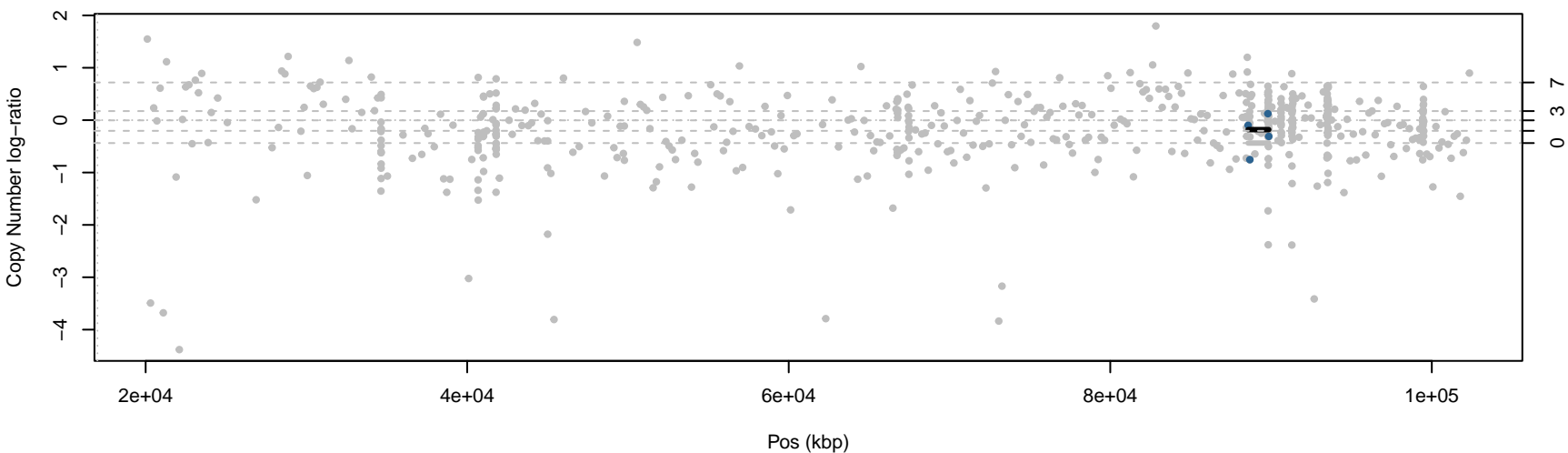
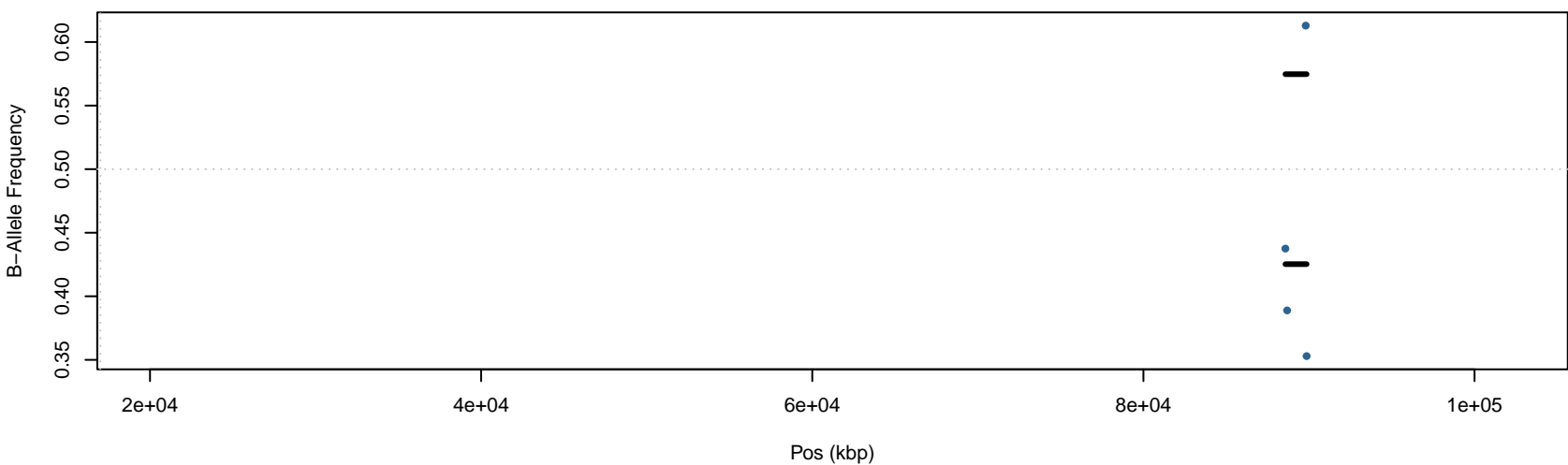
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 13



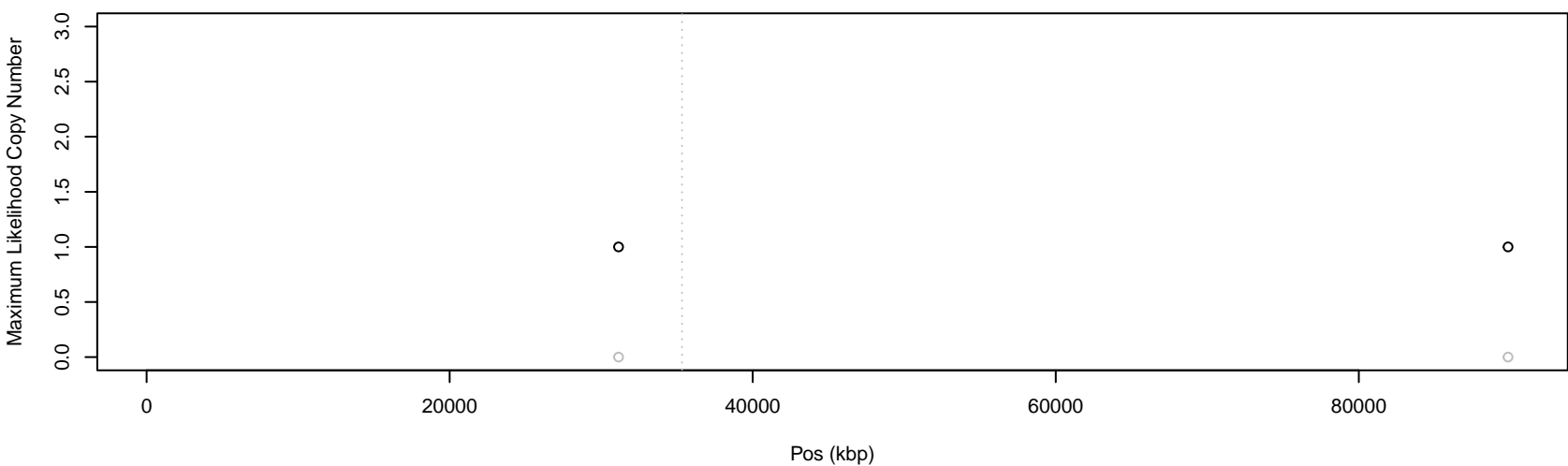
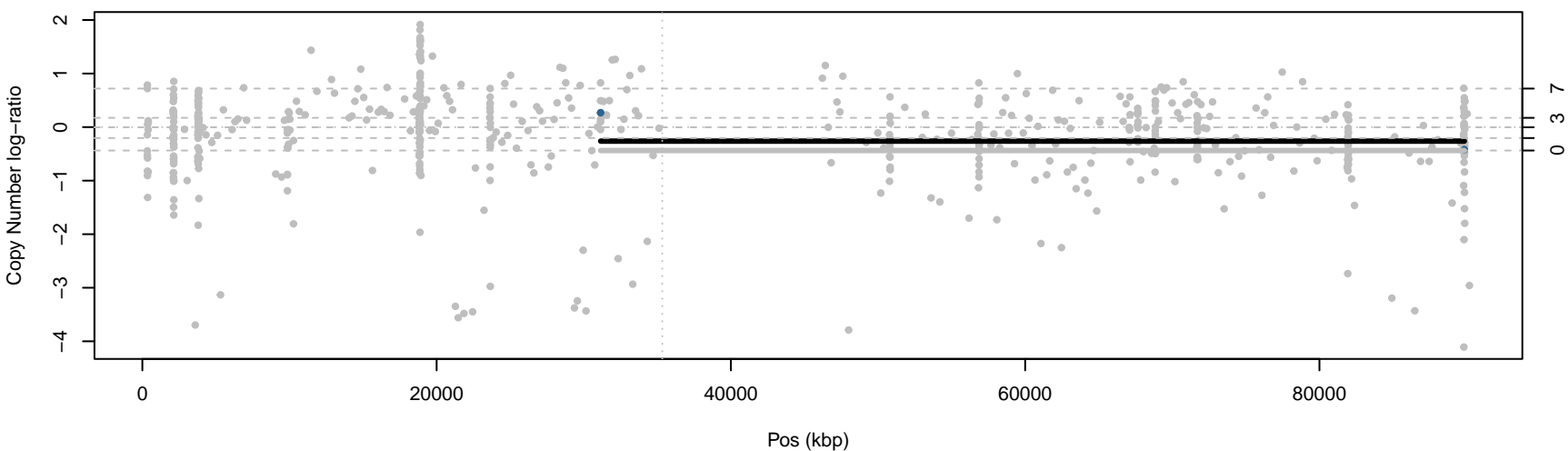
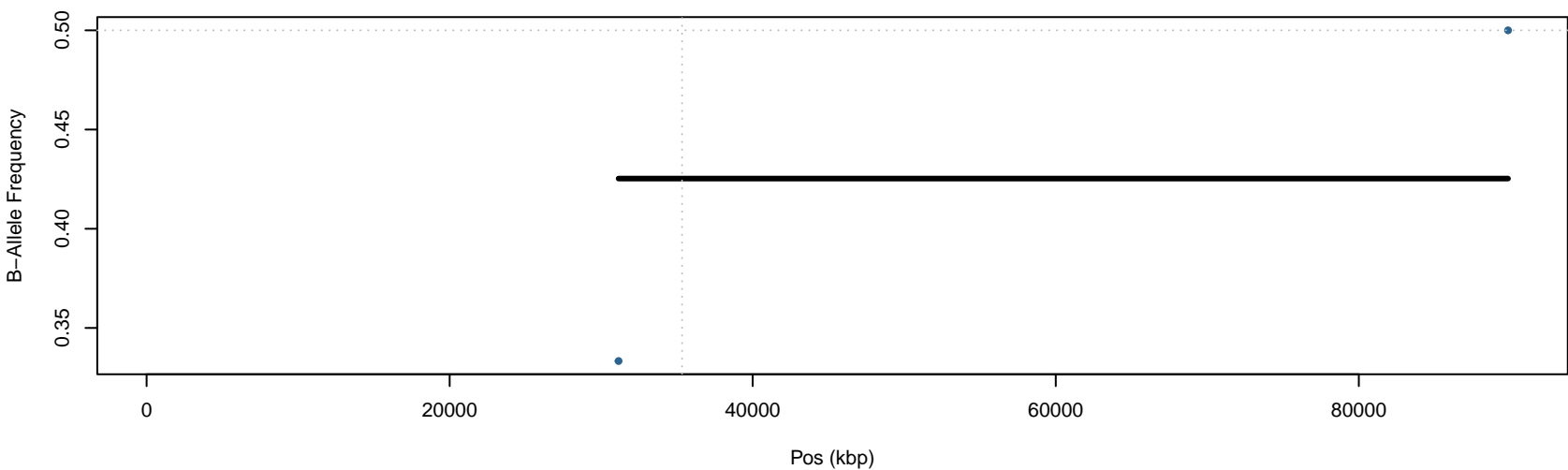
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 14



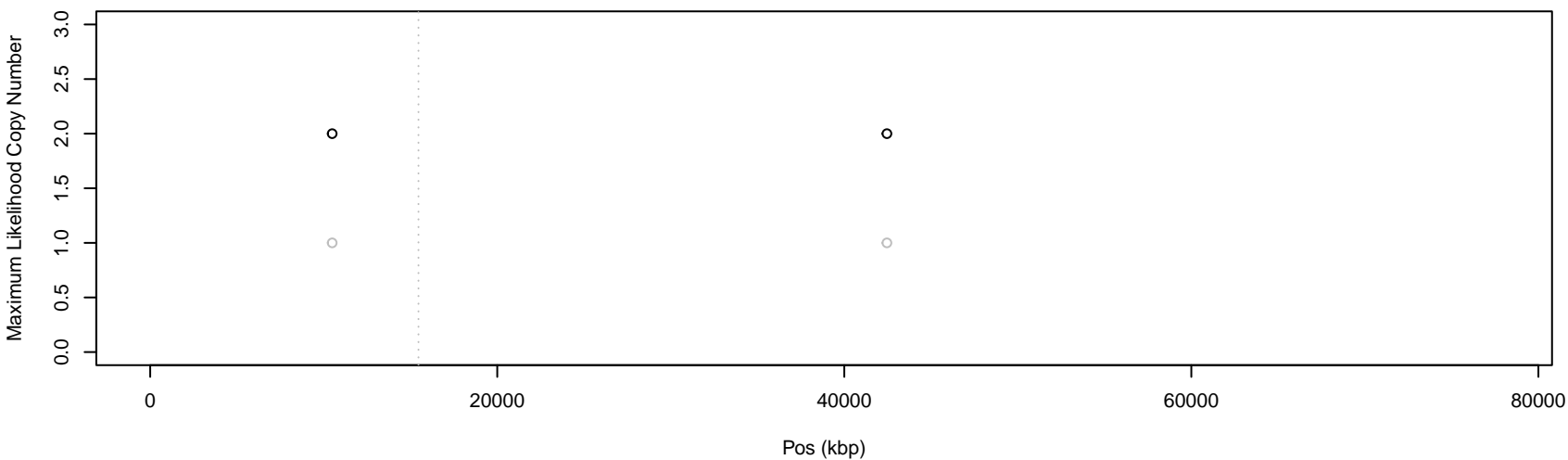
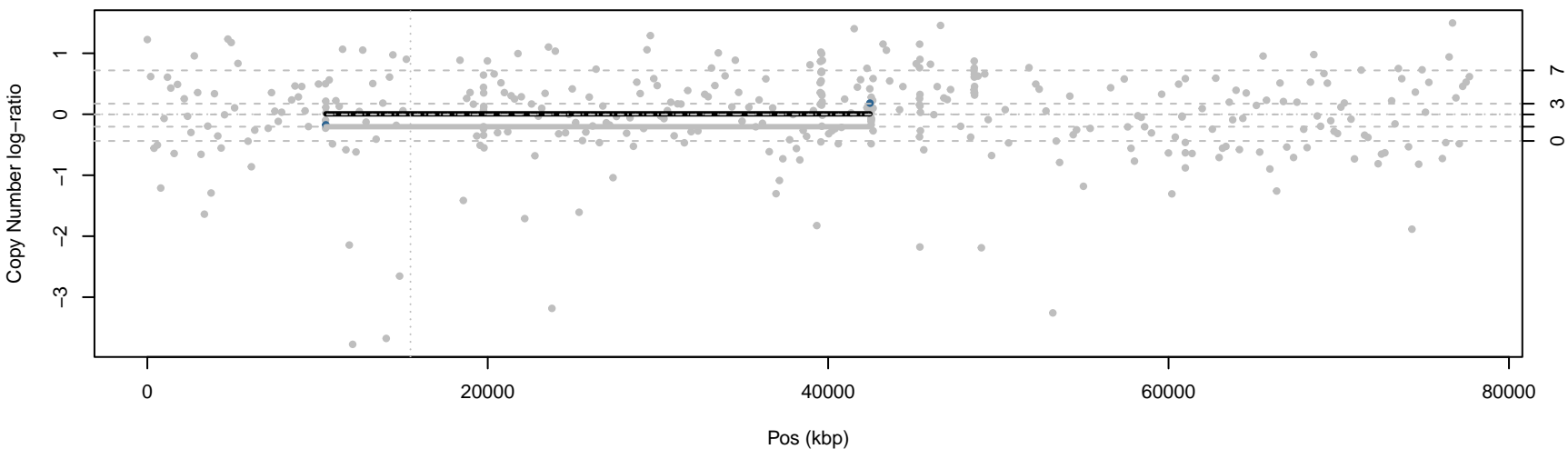
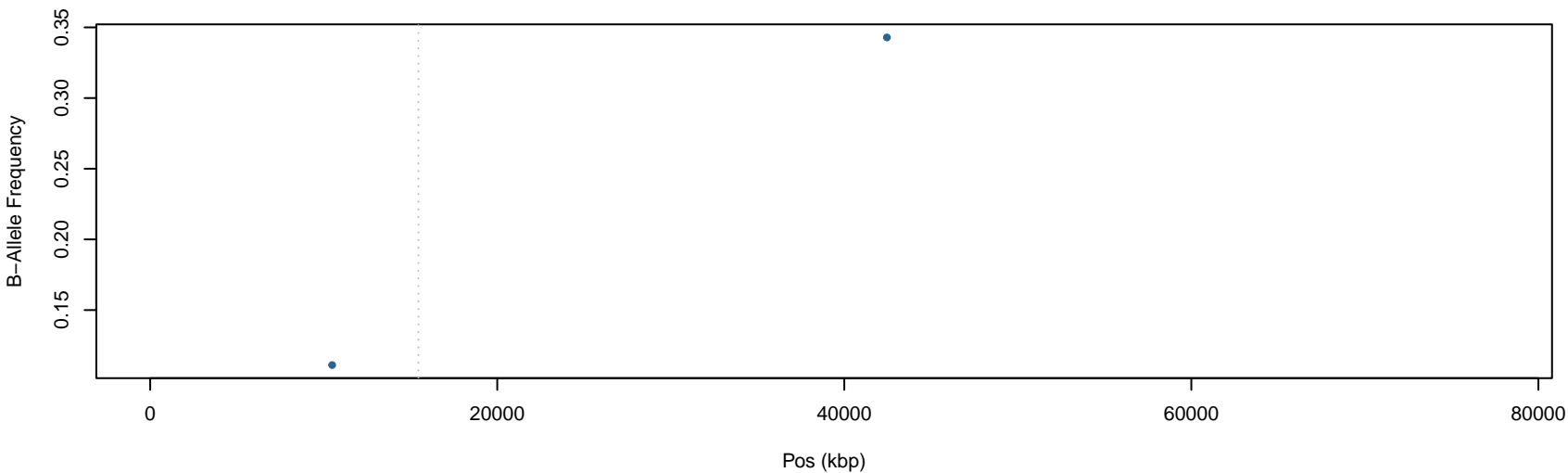
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 15



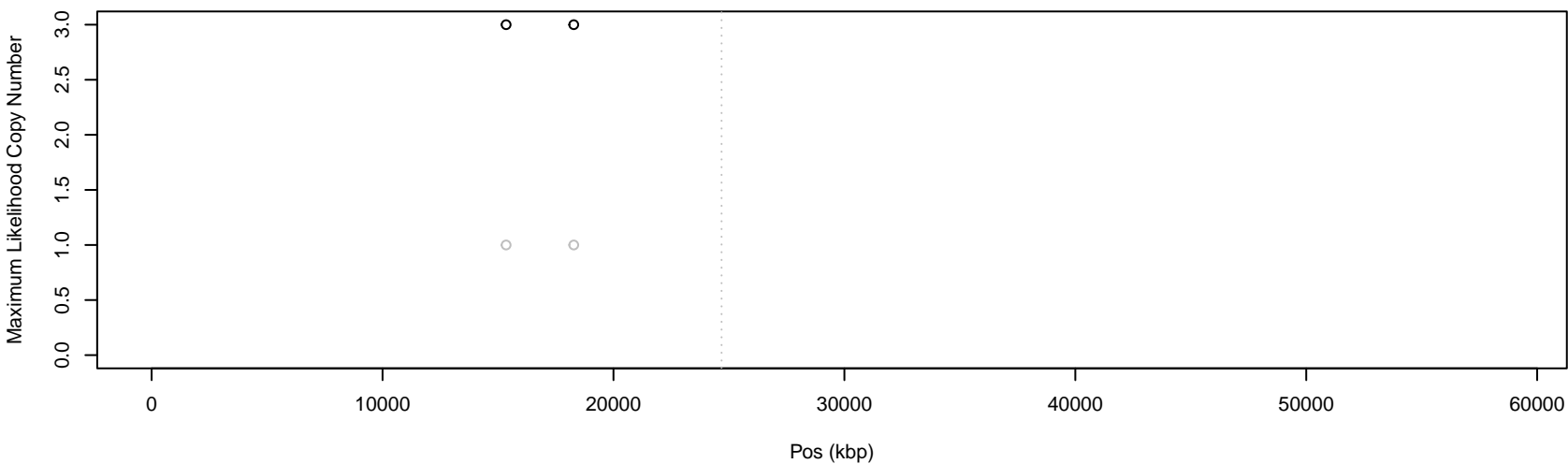
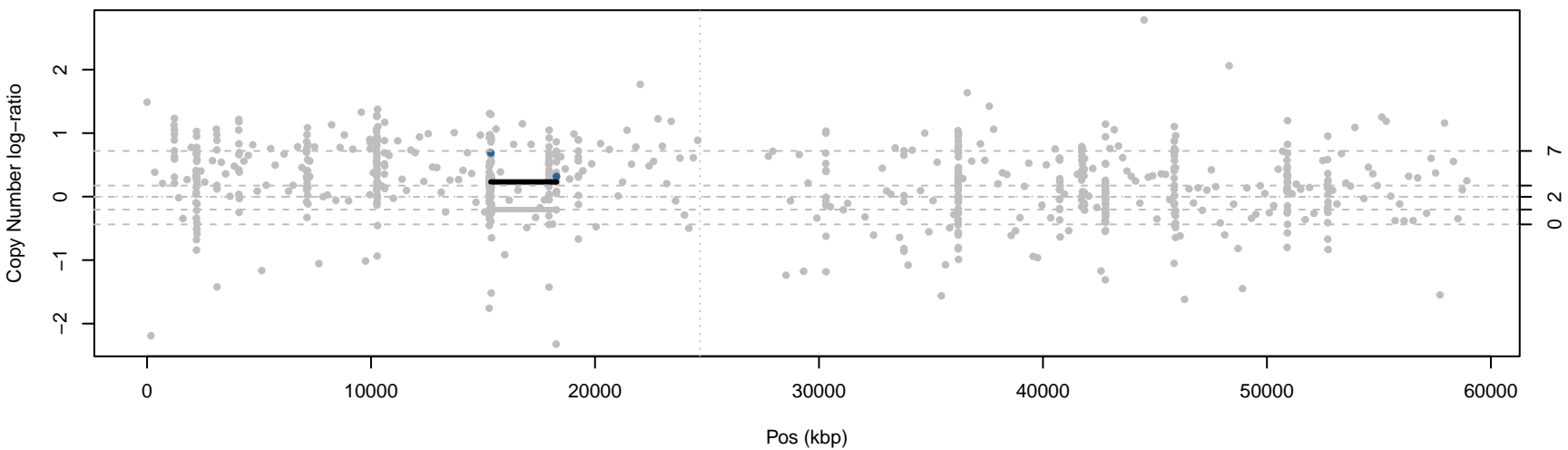
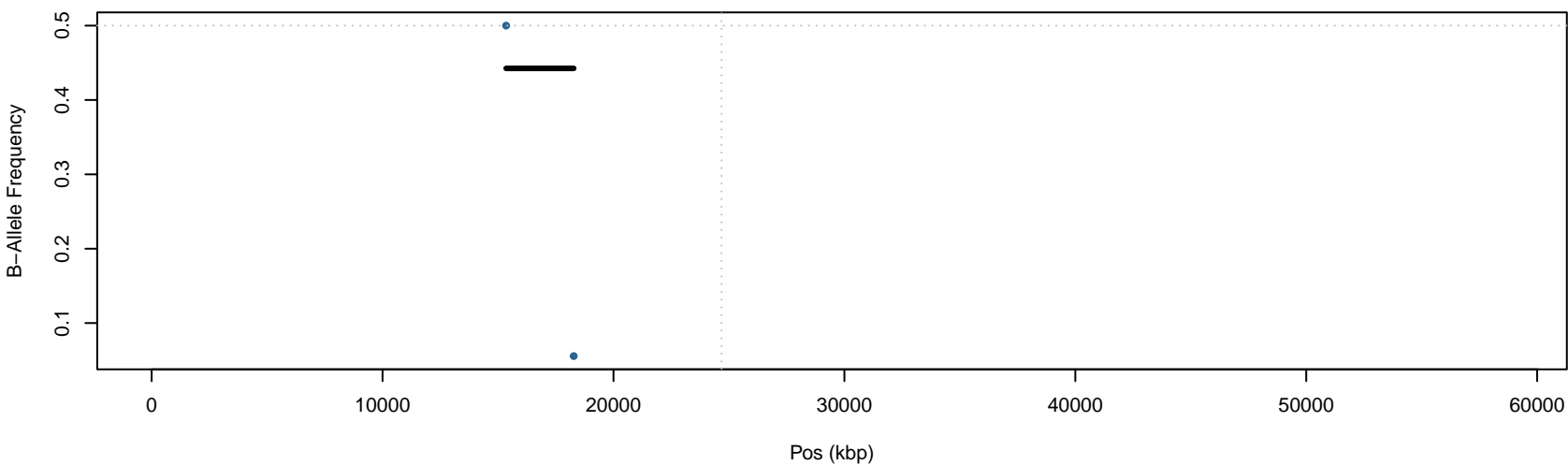
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 16



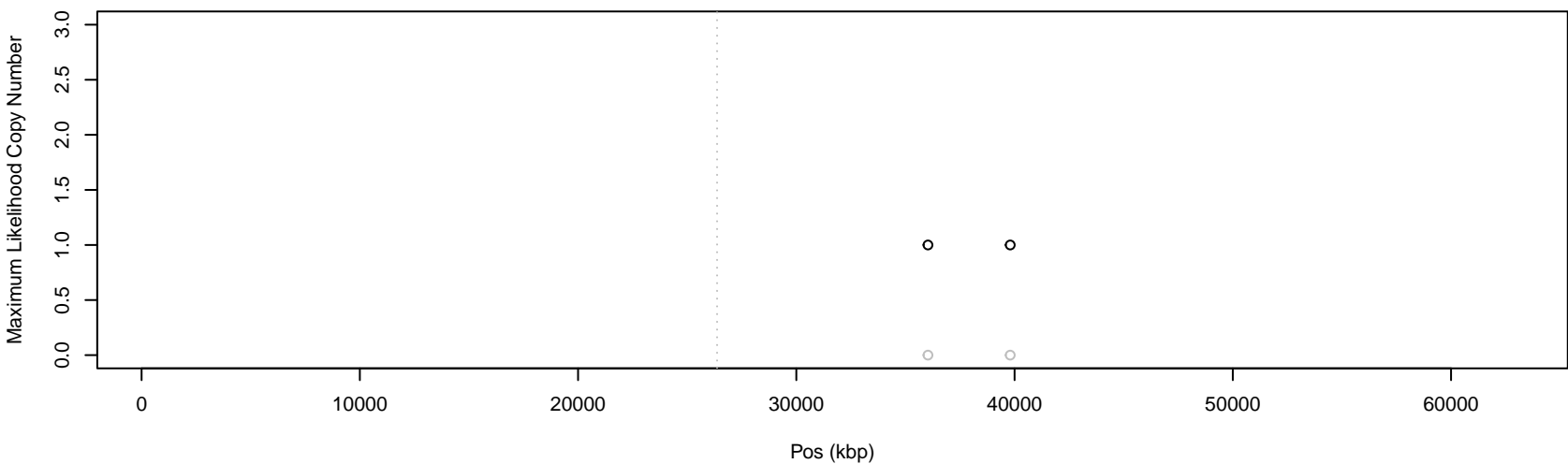
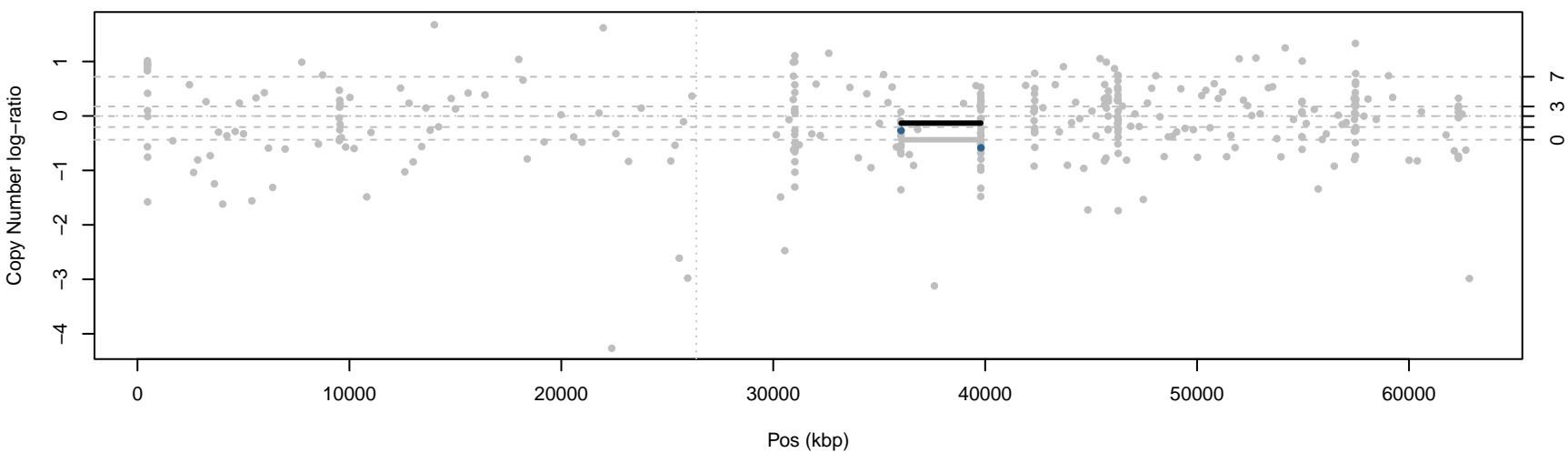
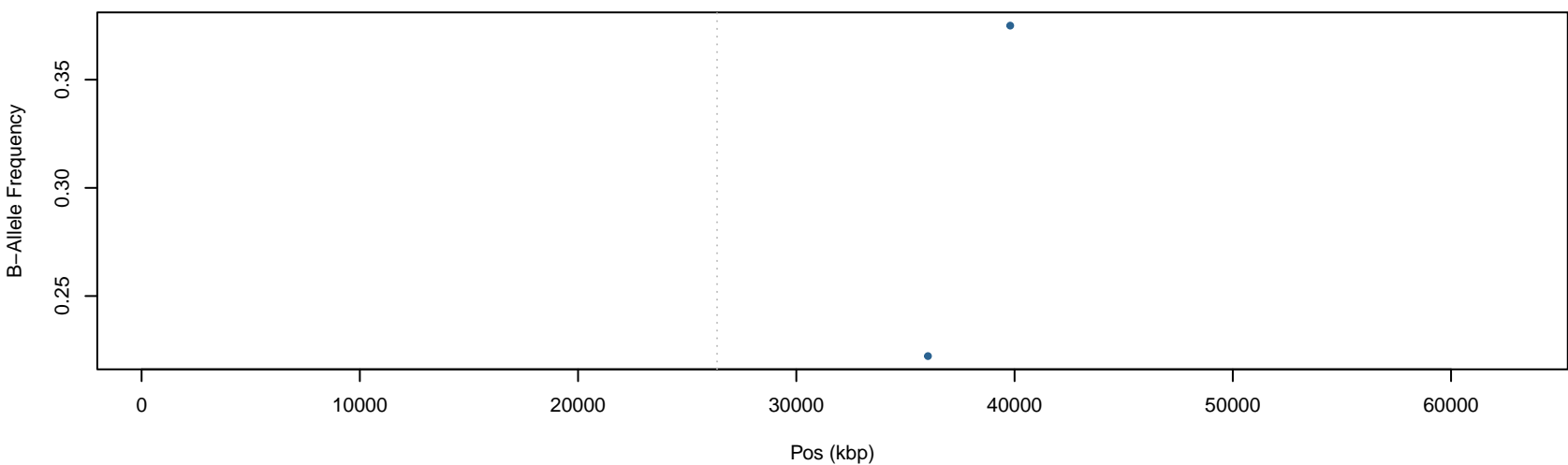
Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 18



Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 19



Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 20





Purity: 0.26 Tumor ploidy: 2.017 SNV log-likelihood: -25.29 GoF: 85.6% Mean coverage: 441;34 Chromosome: 22

