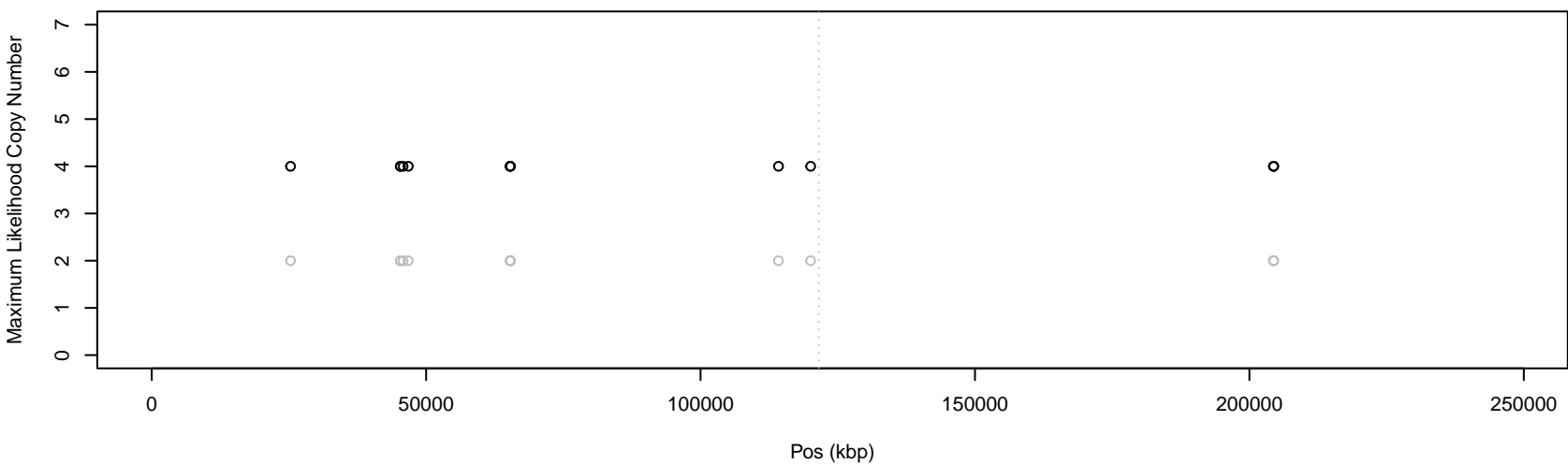
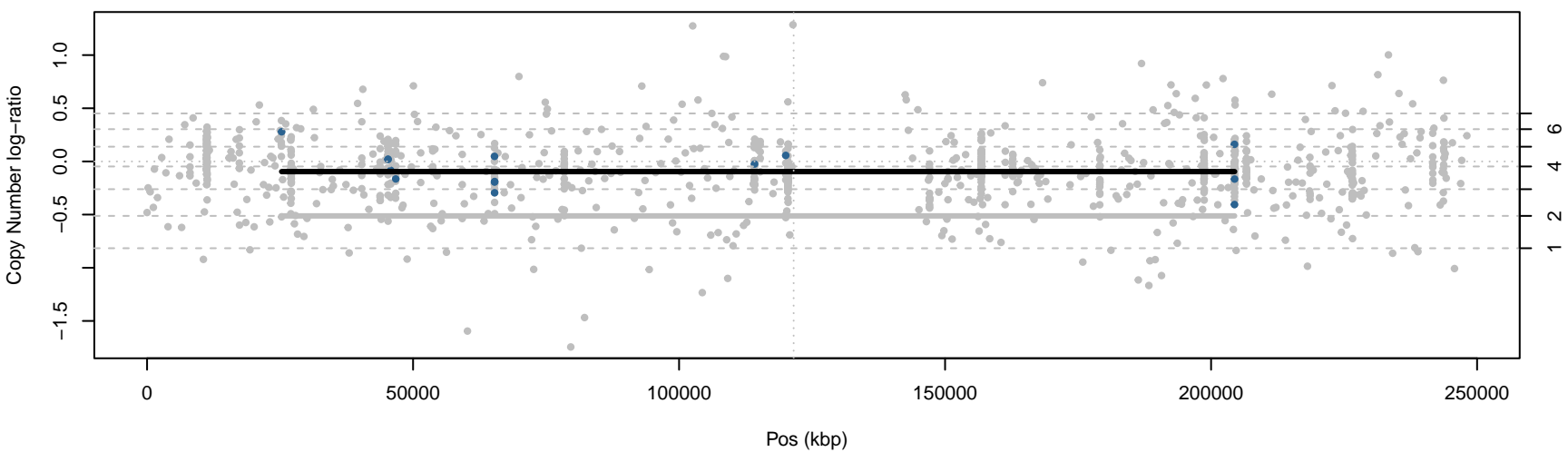
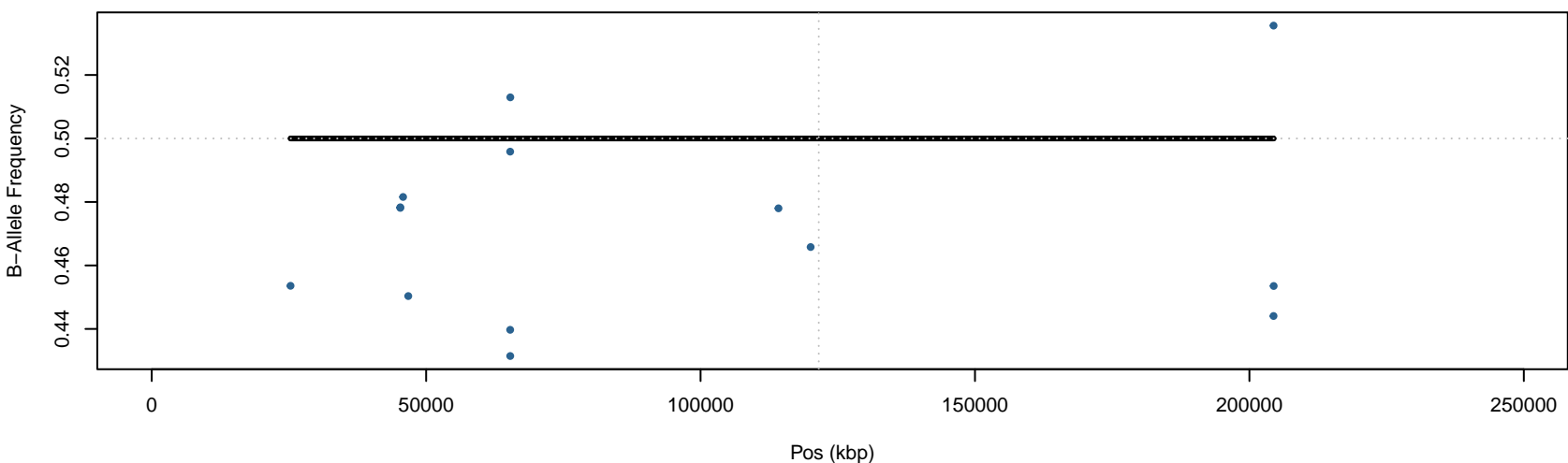
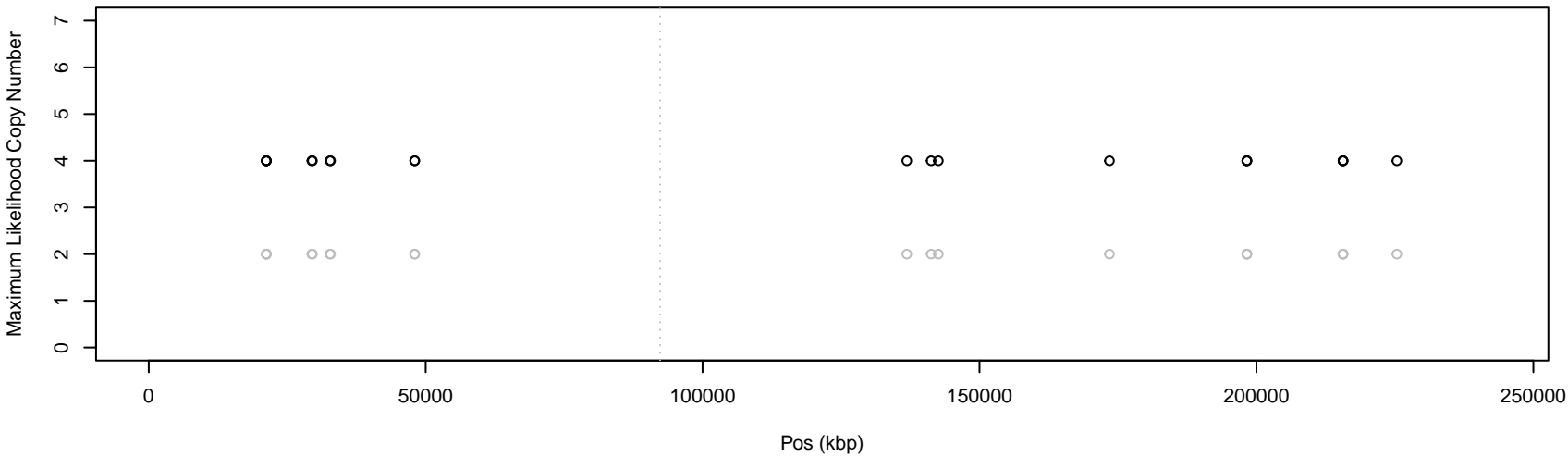
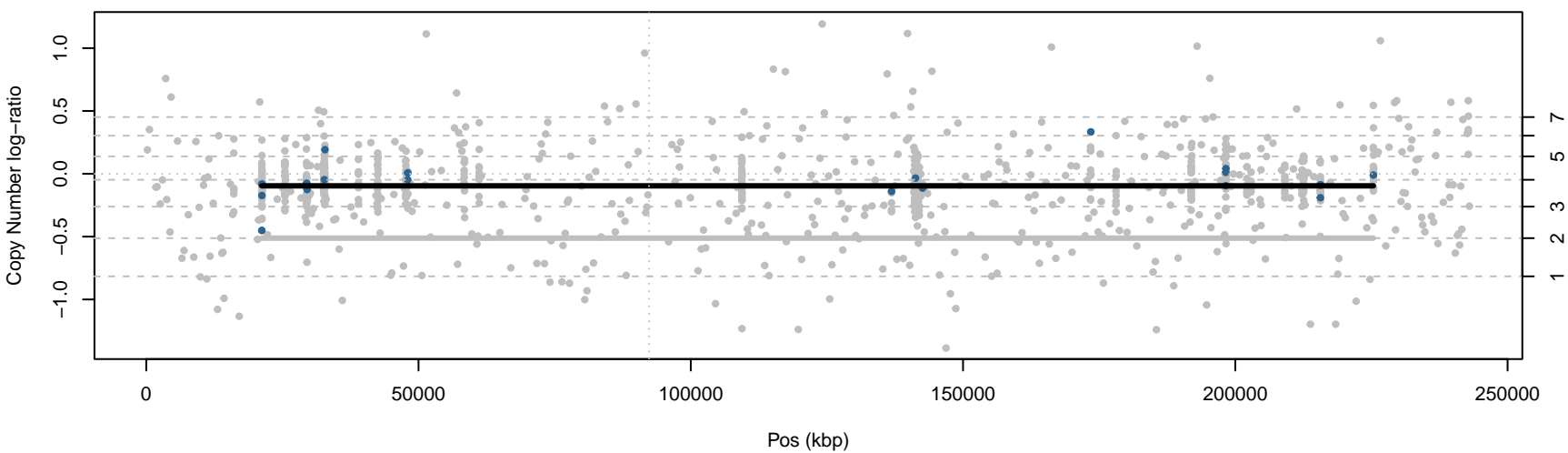
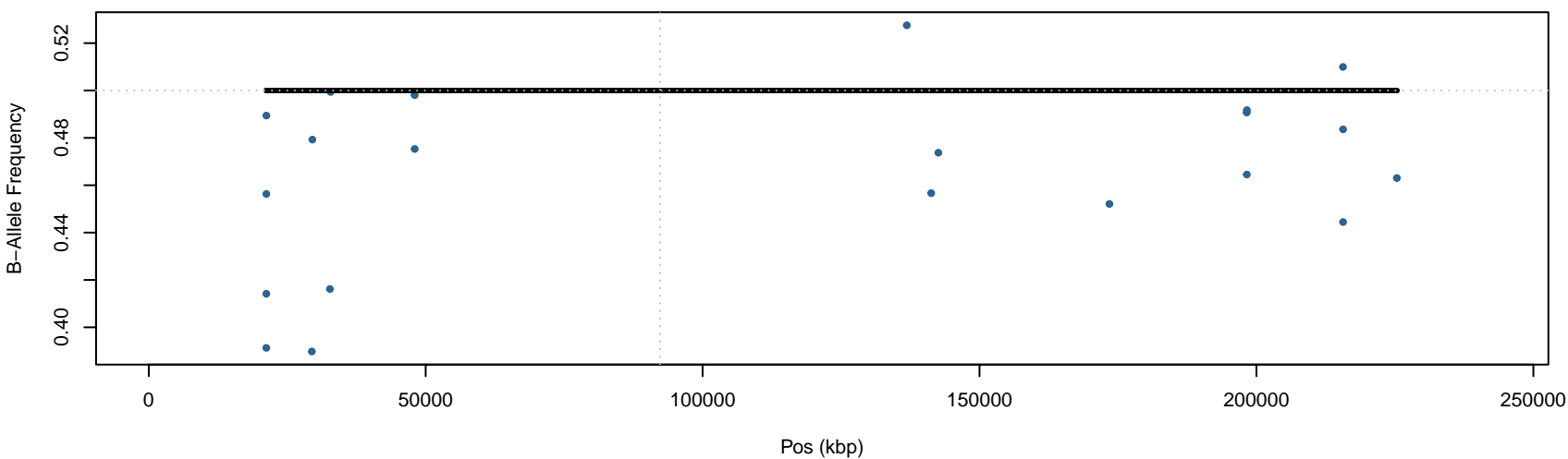


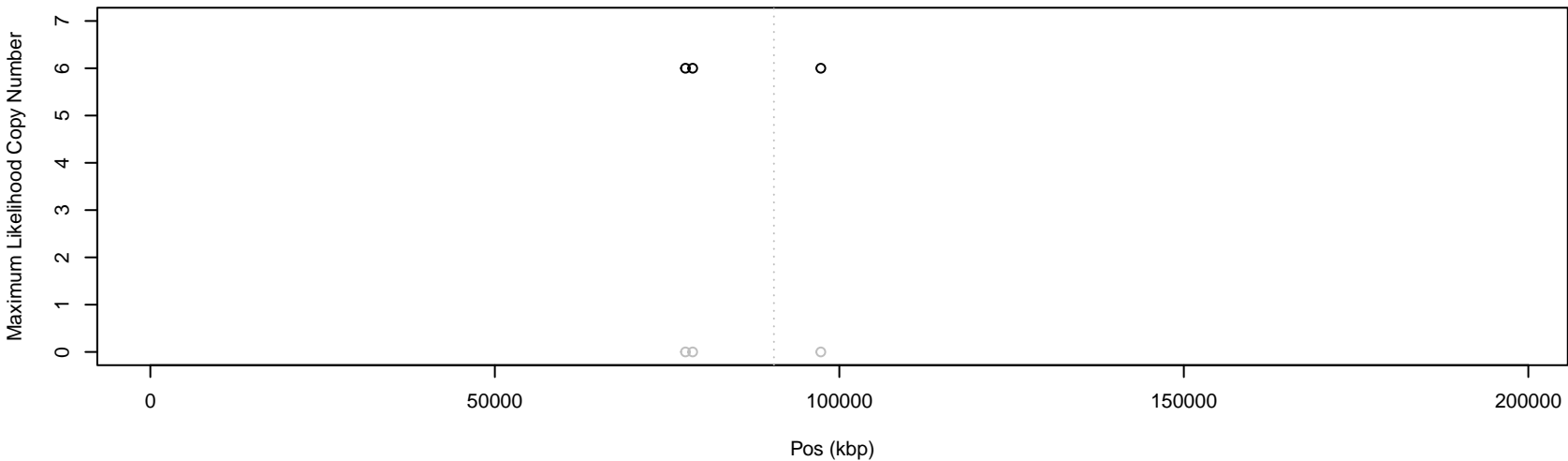
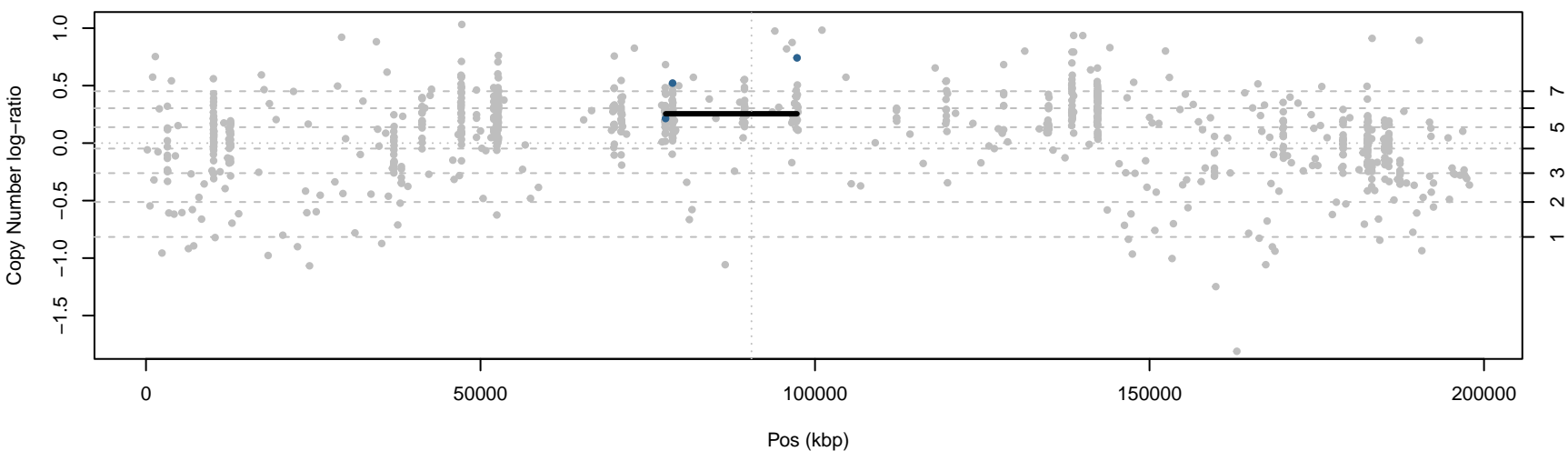
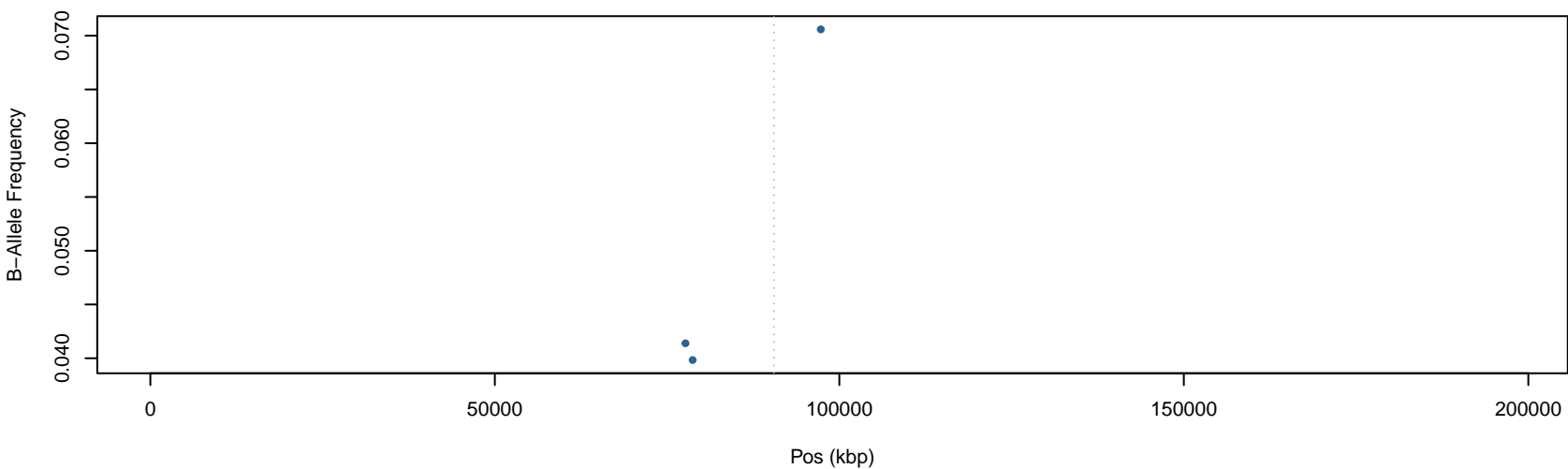
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 1



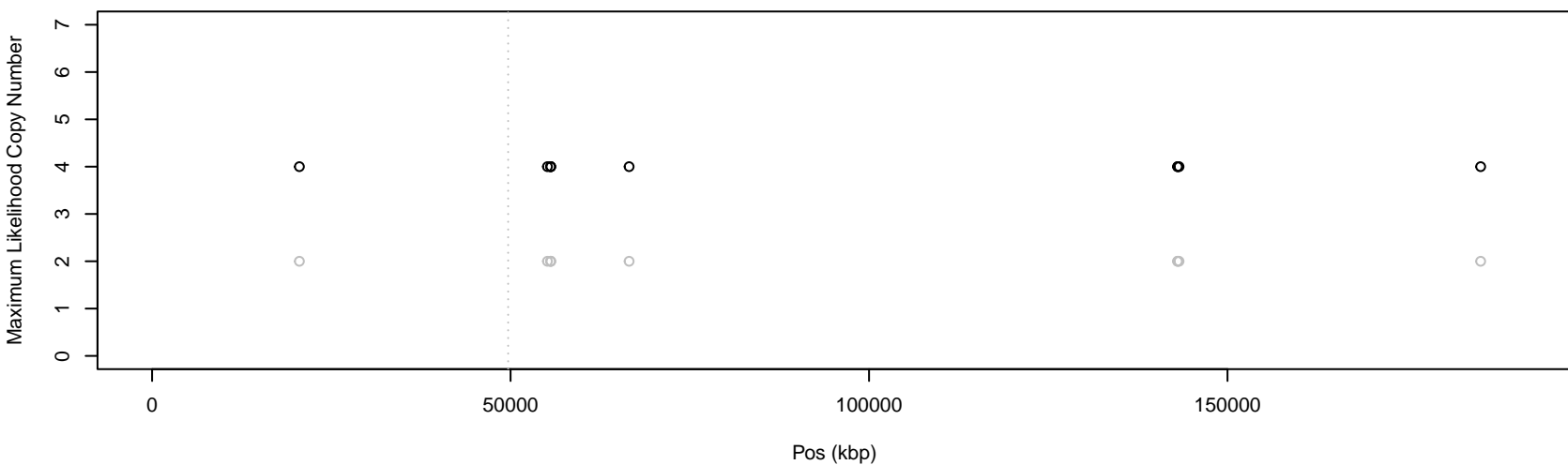
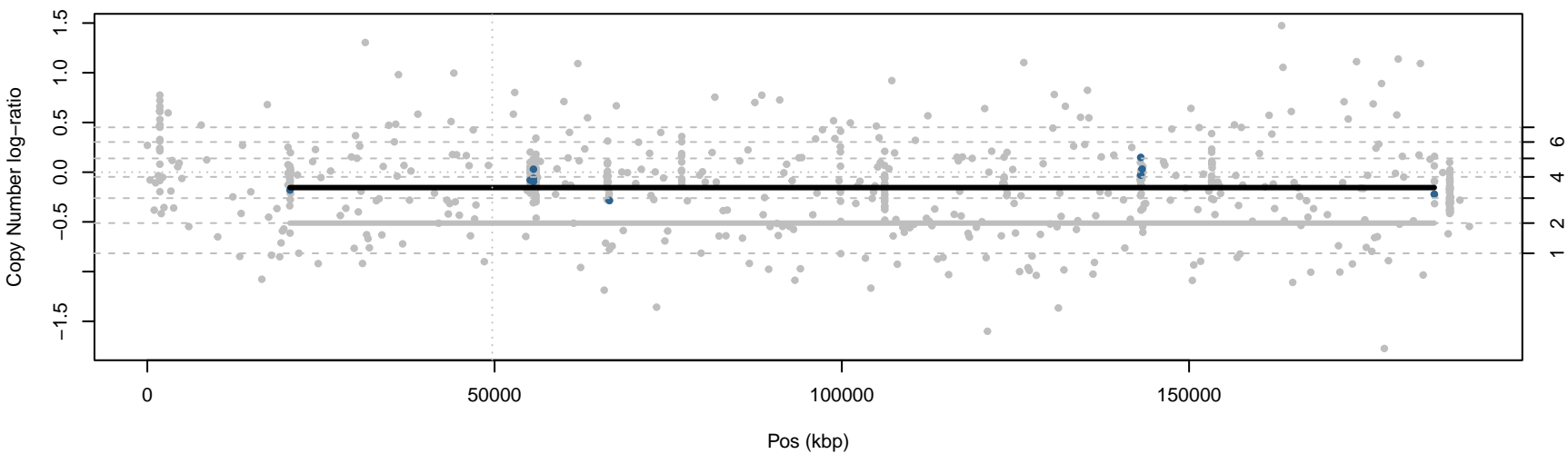
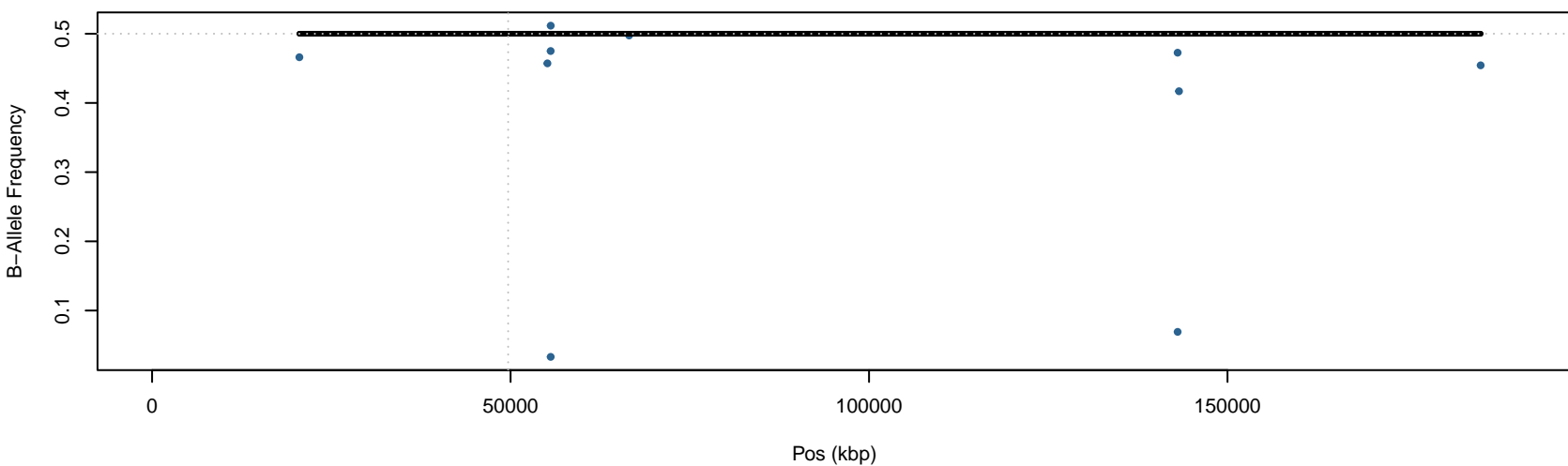
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 2



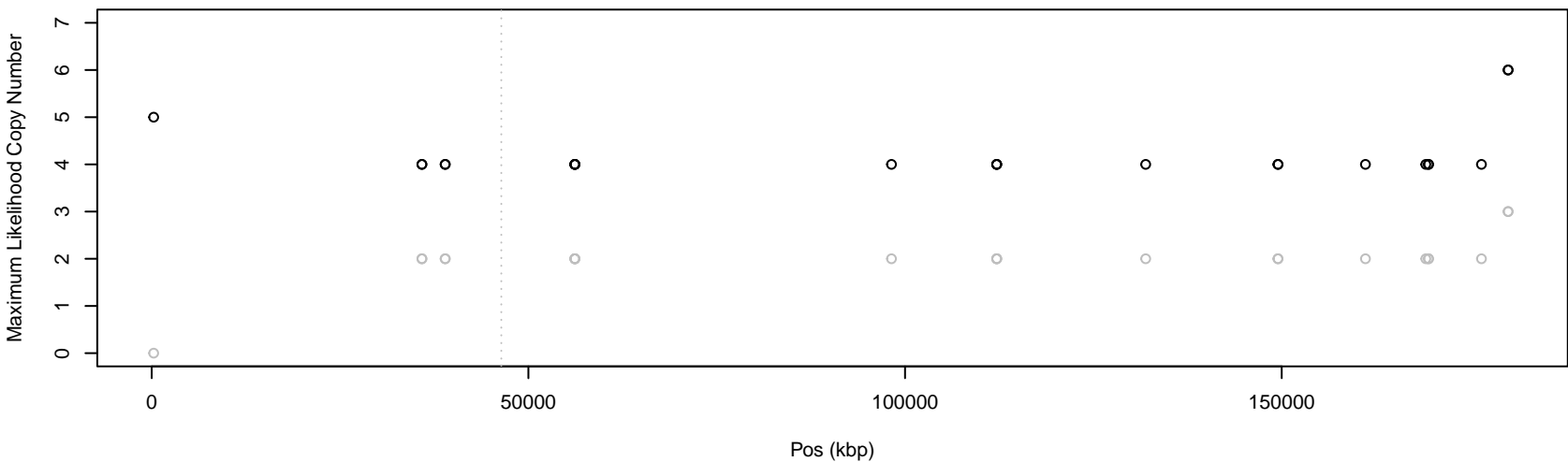
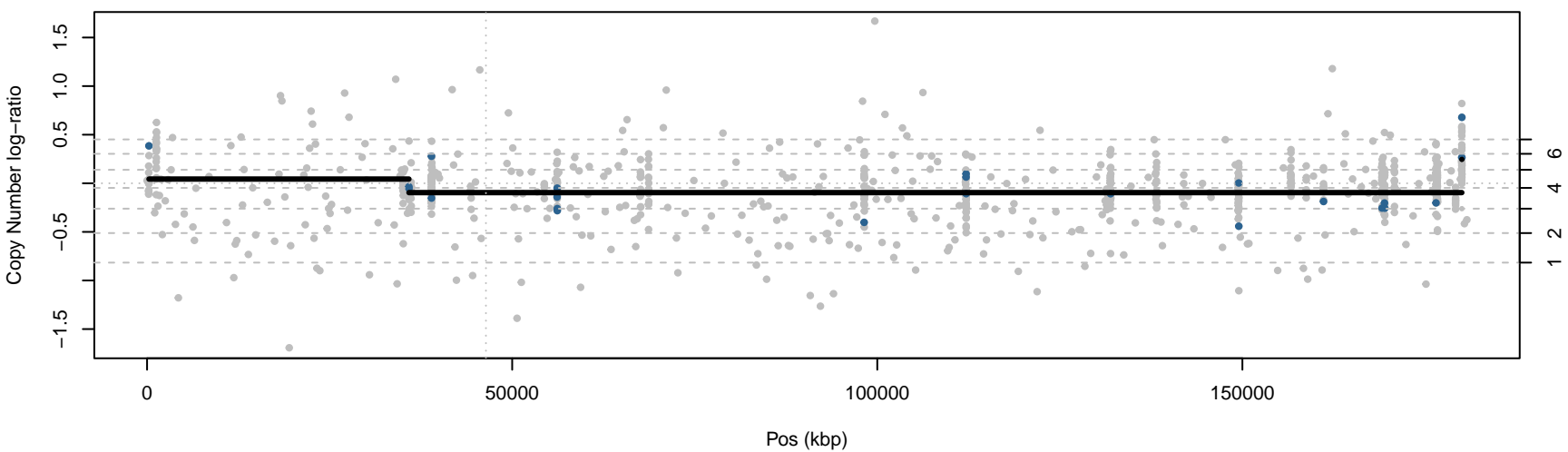
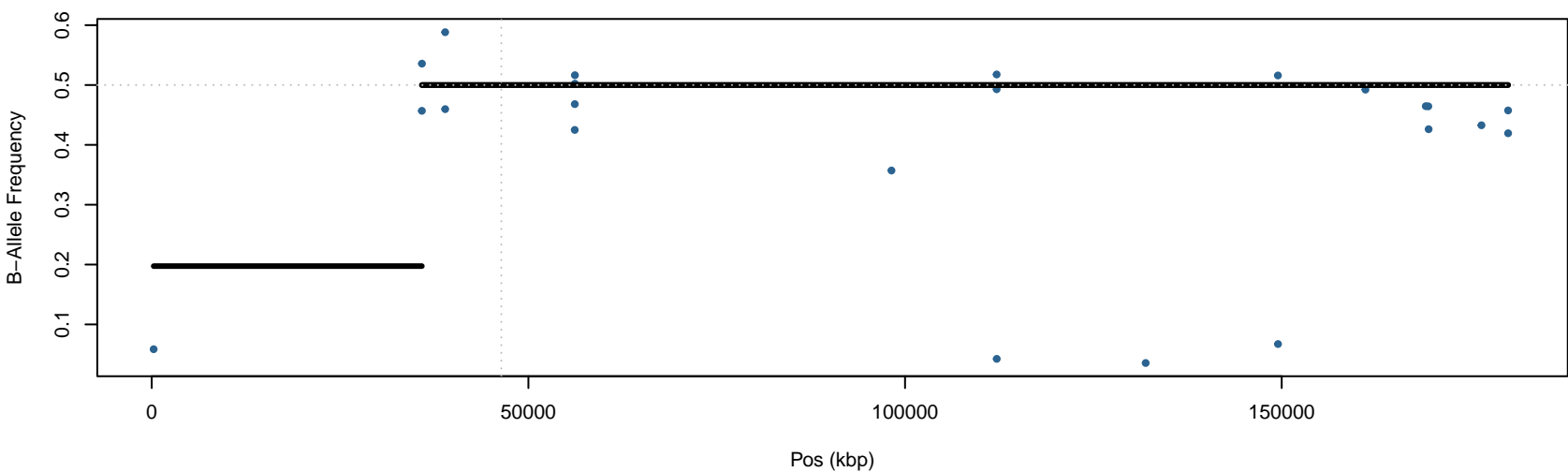
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 3



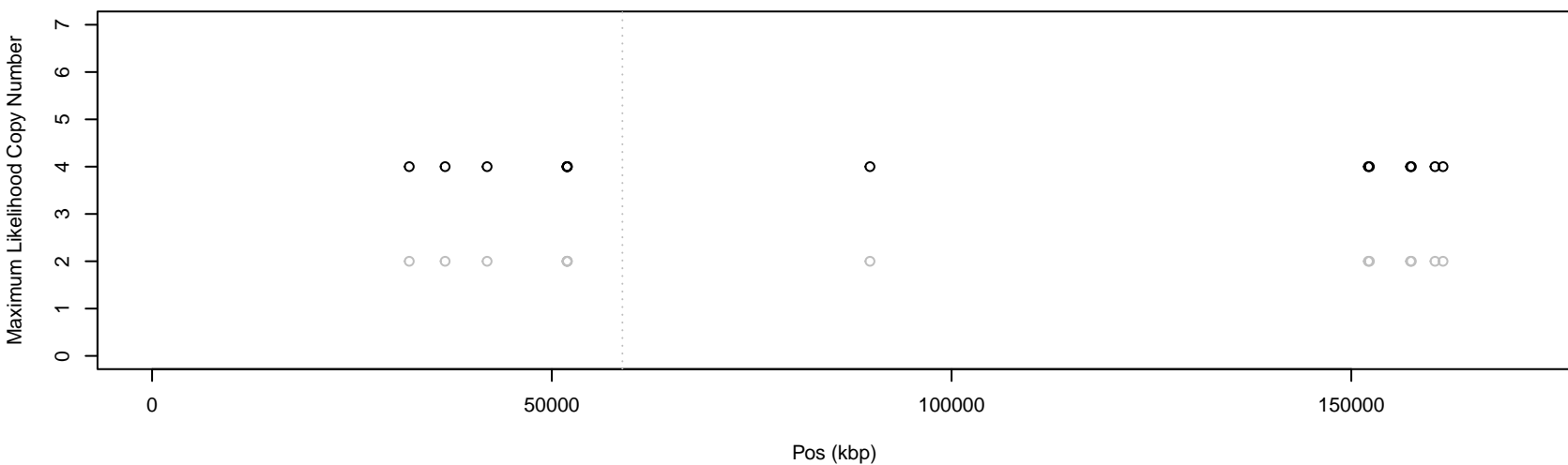
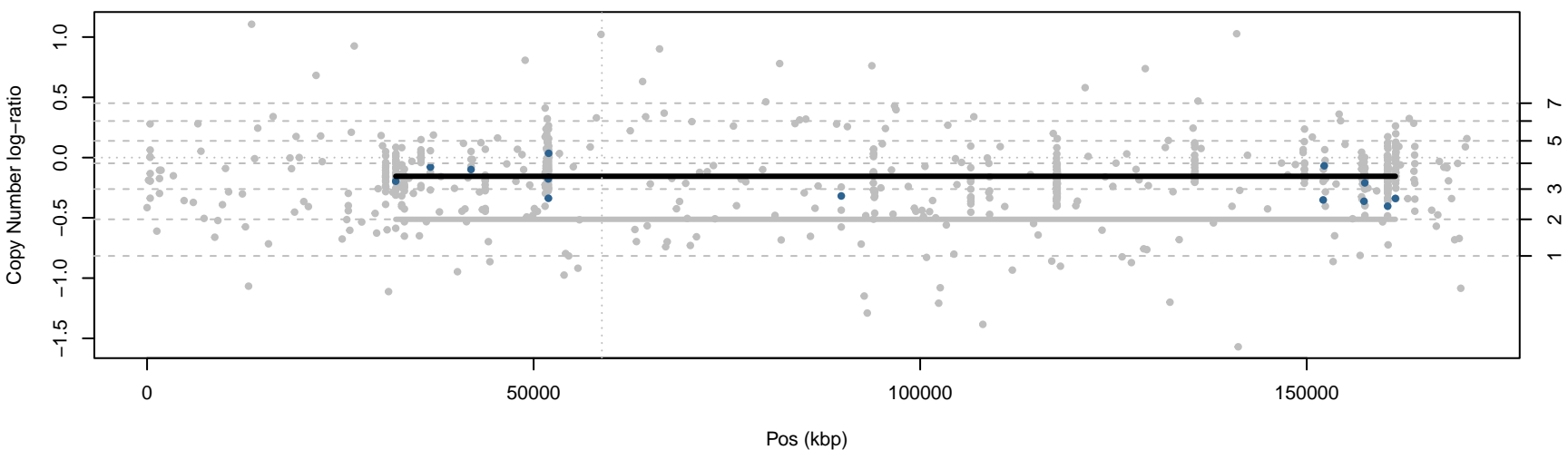
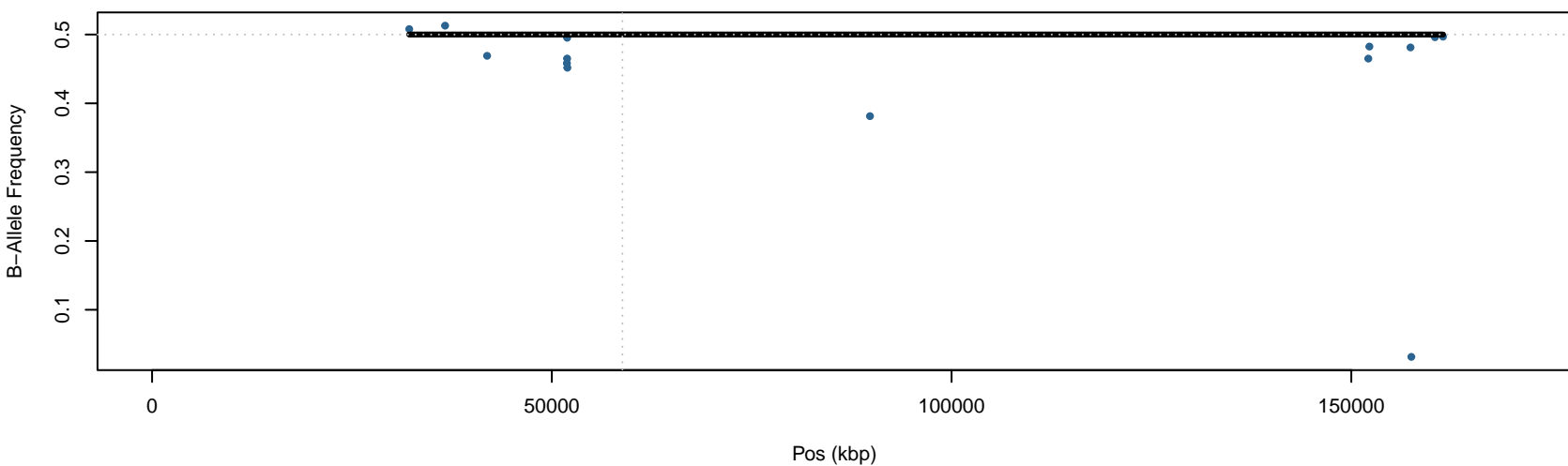
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 4



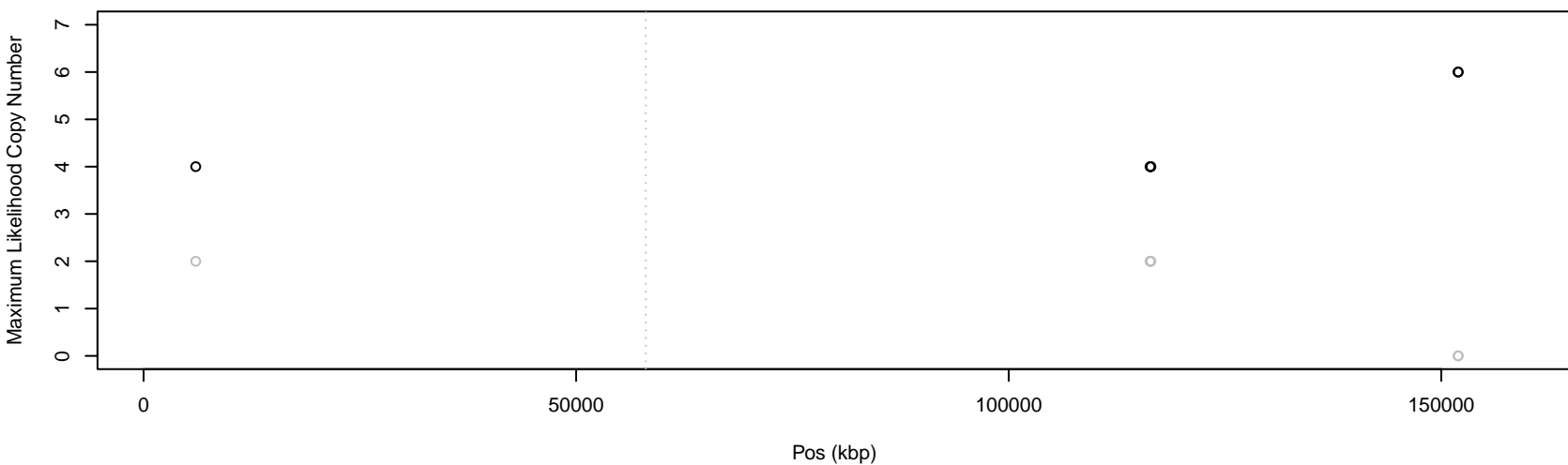
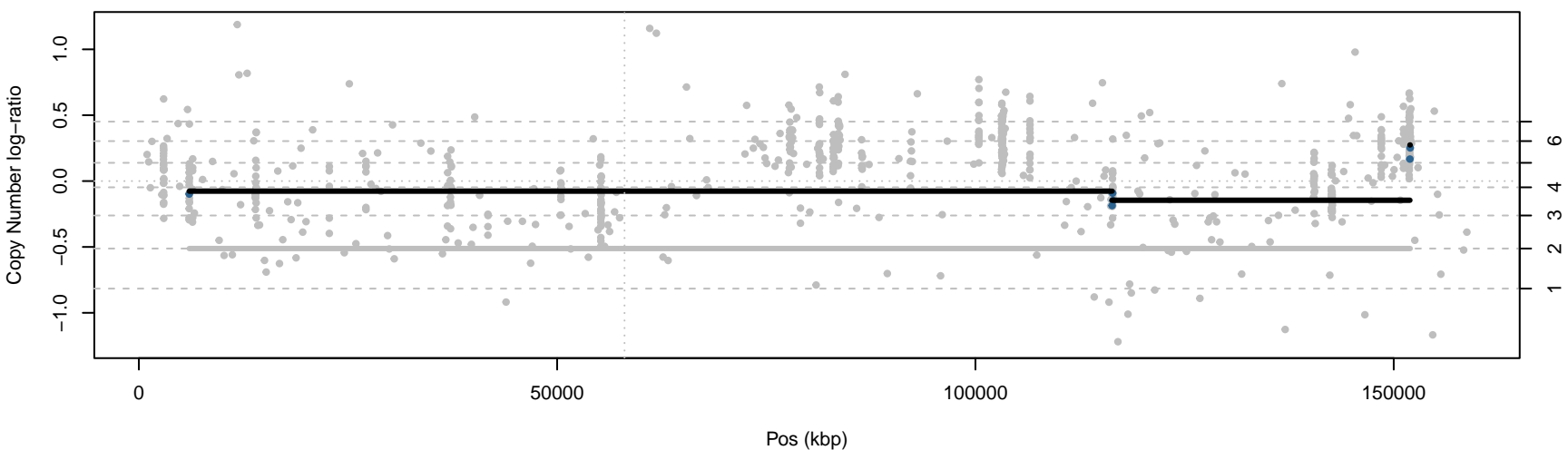
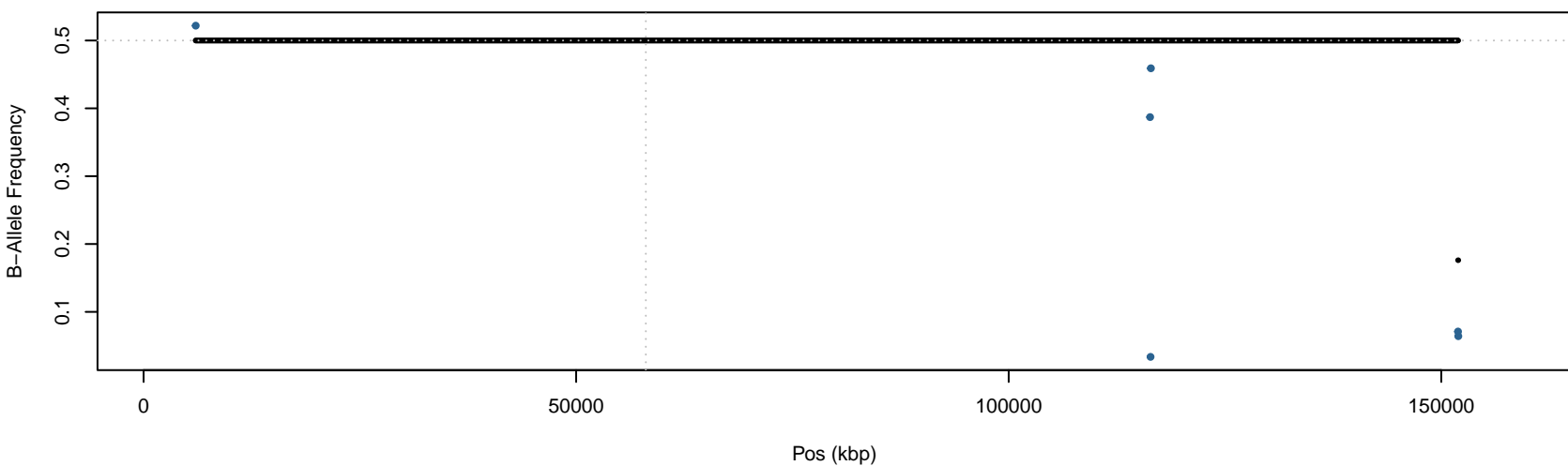
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 5



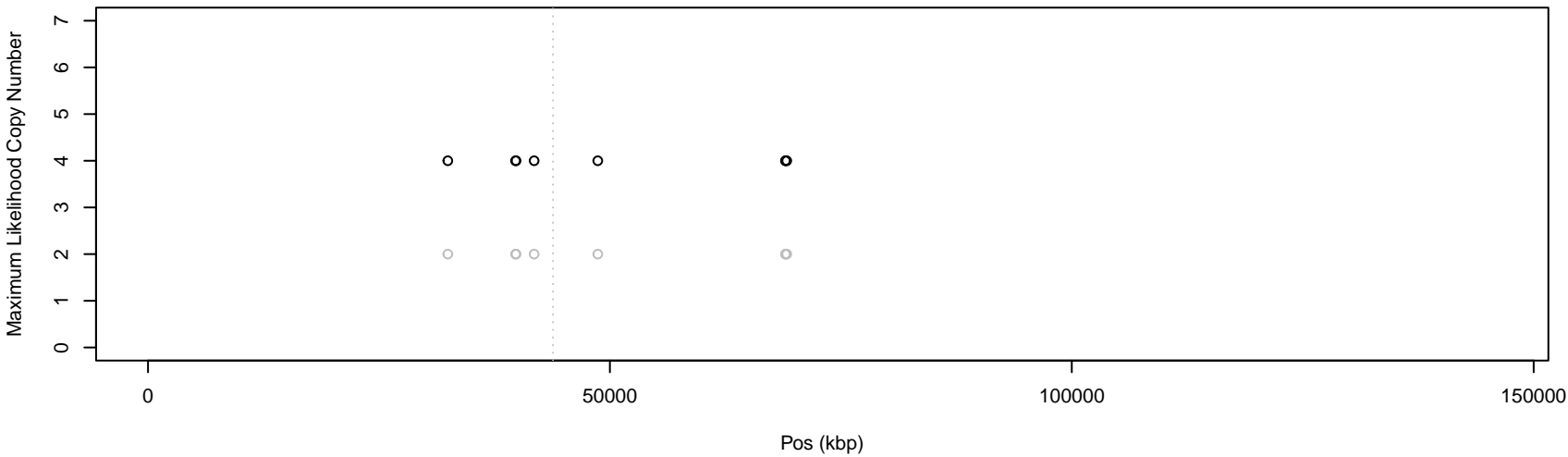
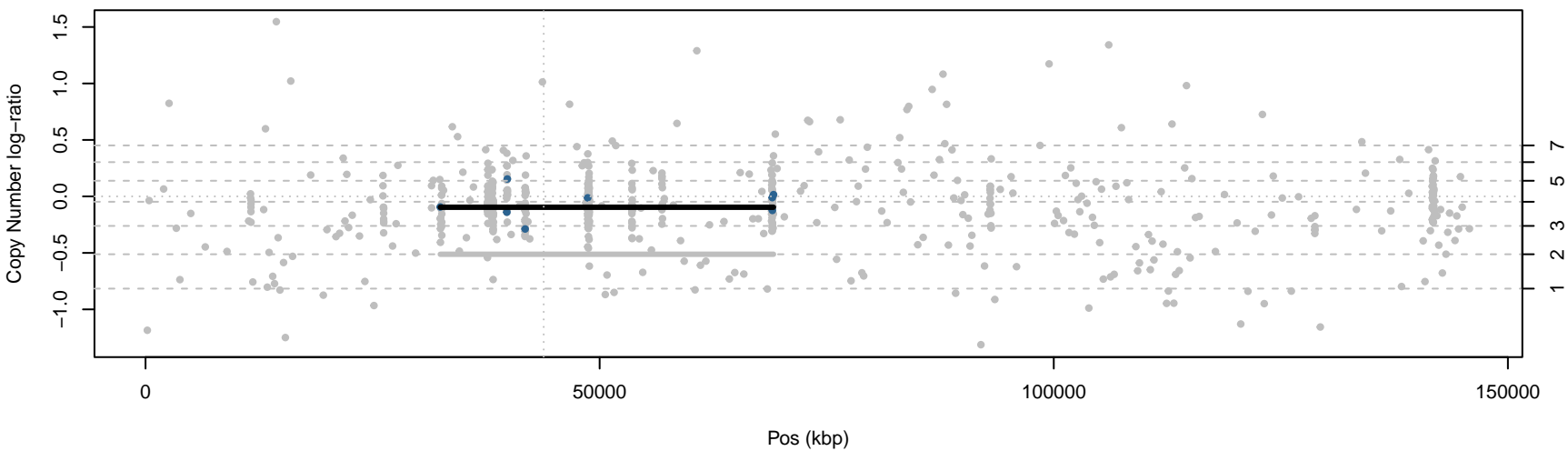
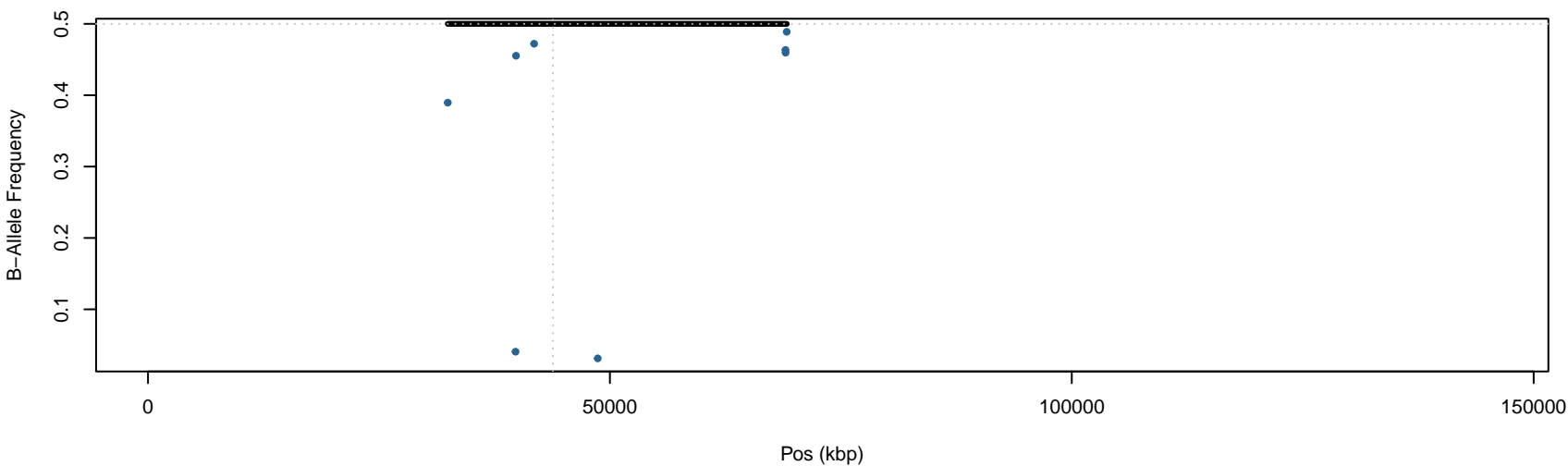
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 6



Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 7

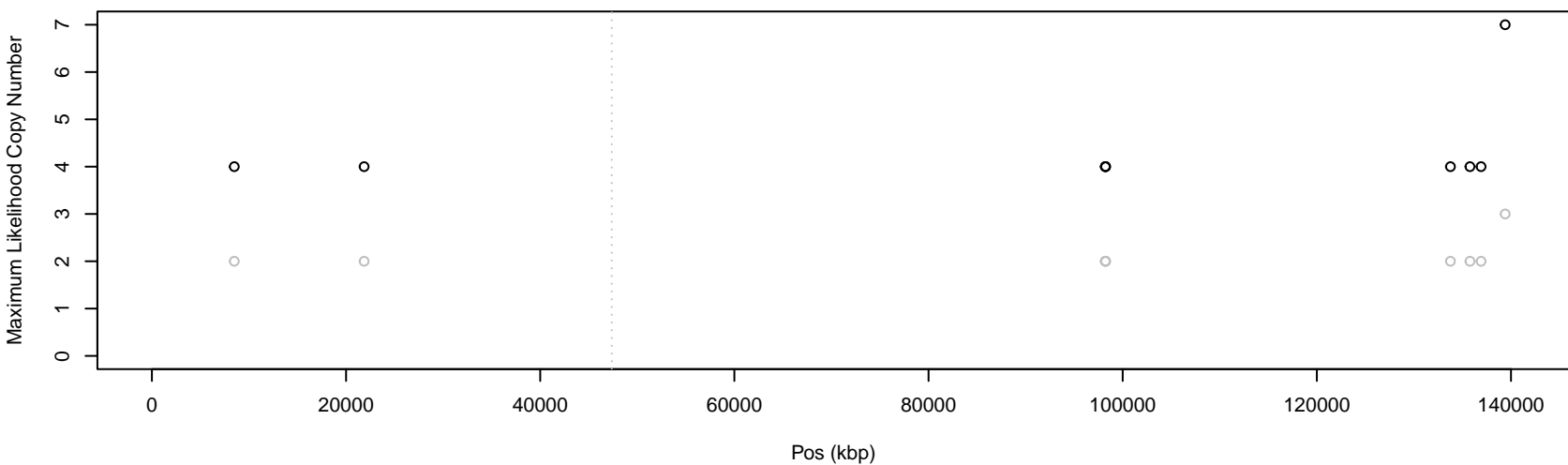
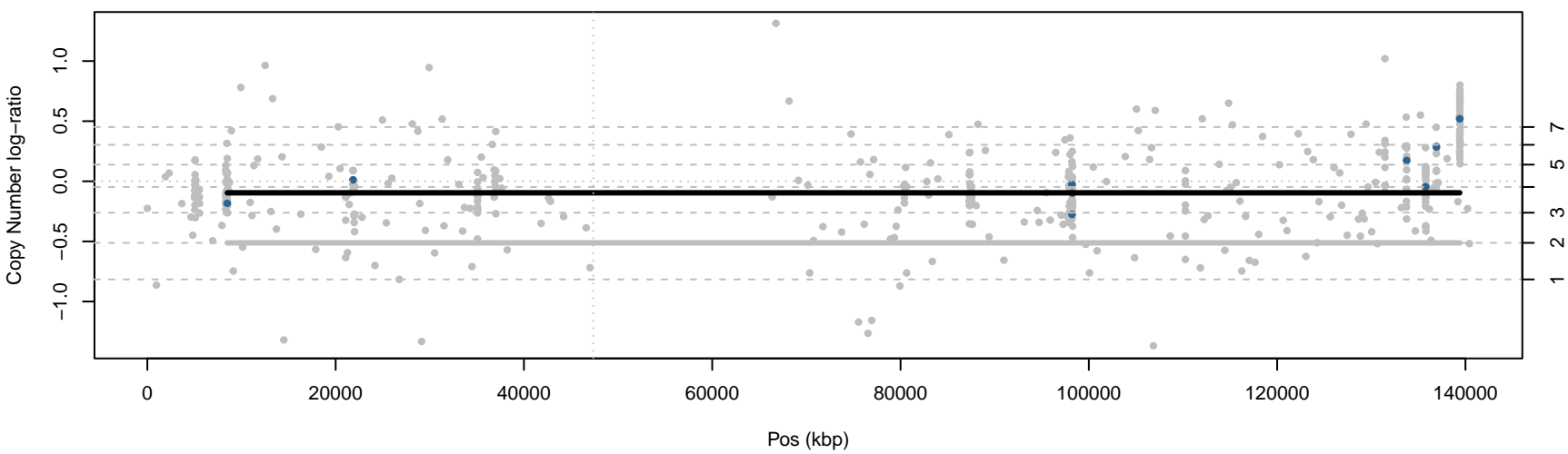
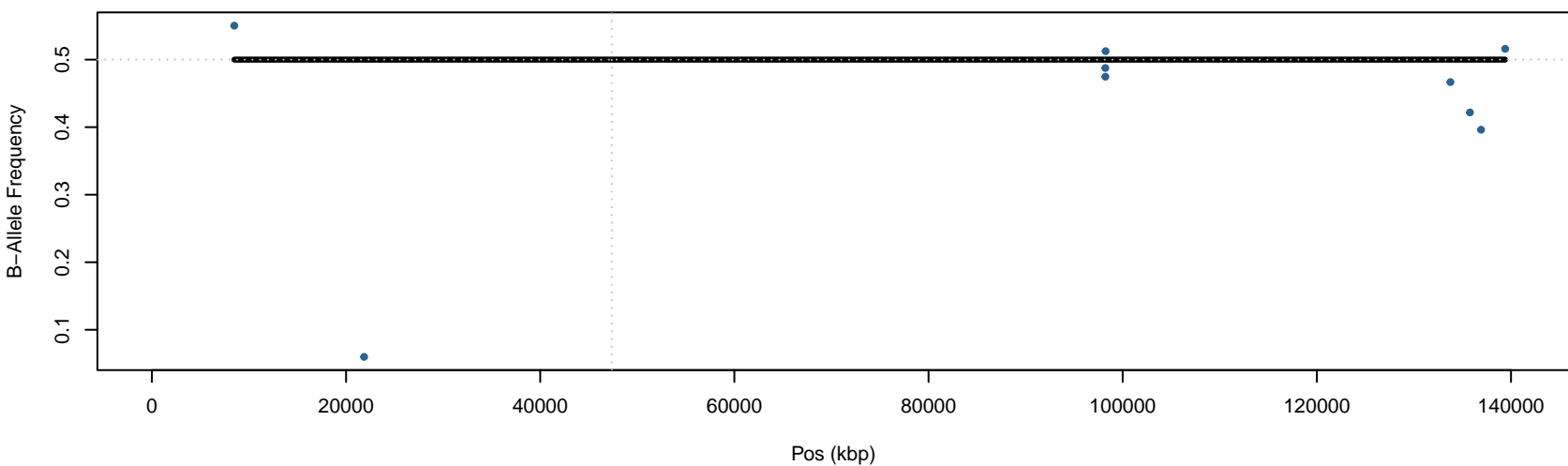


Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 8

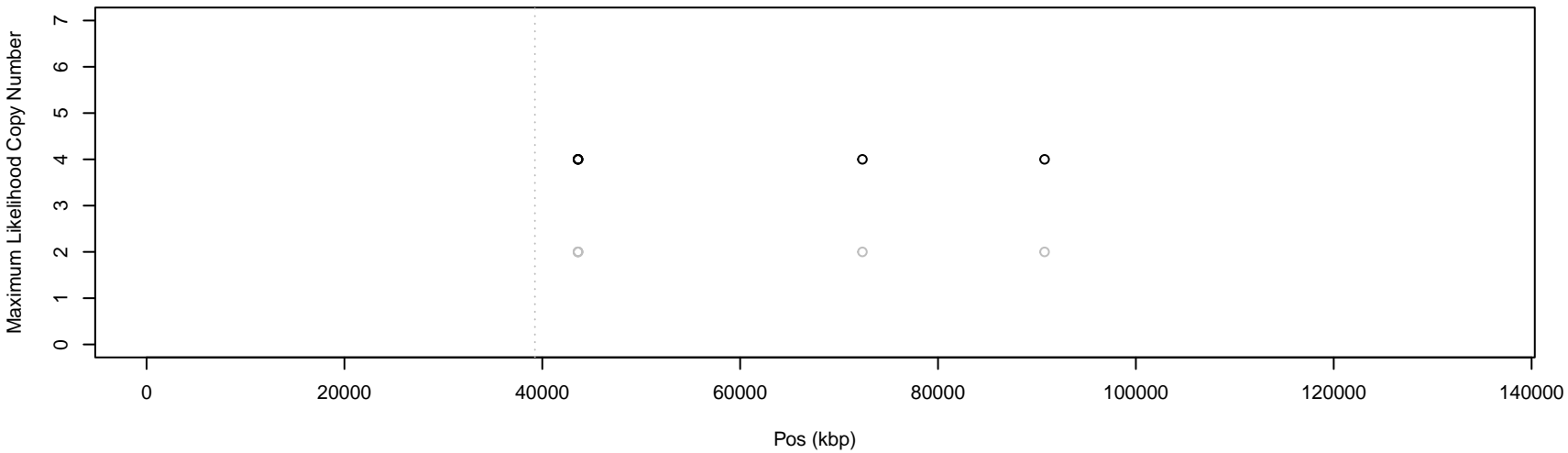
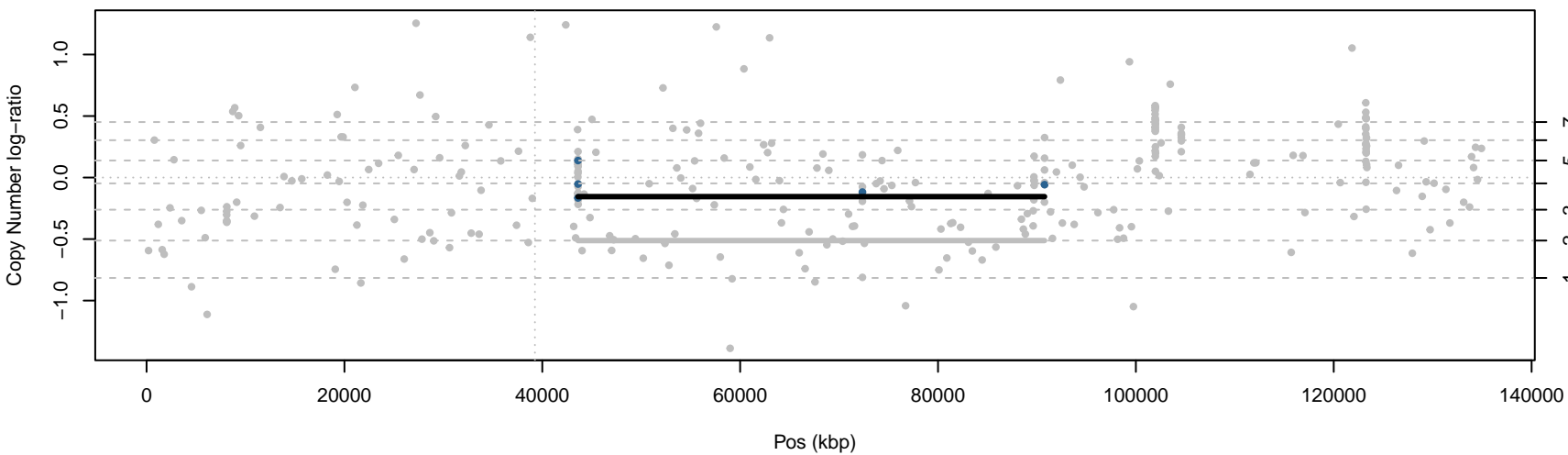
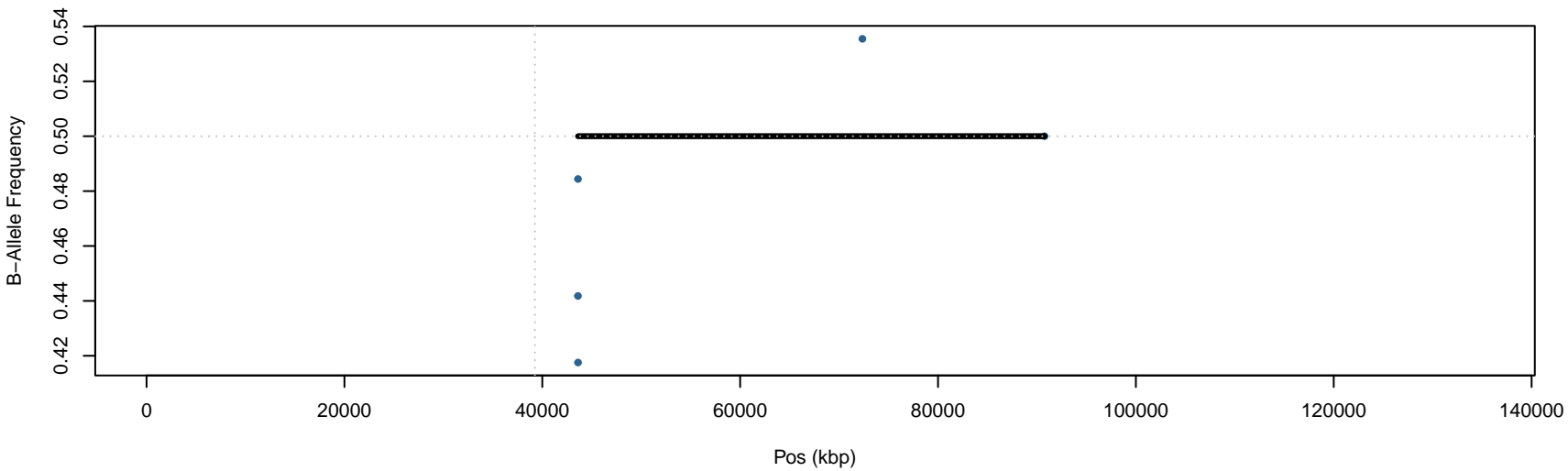




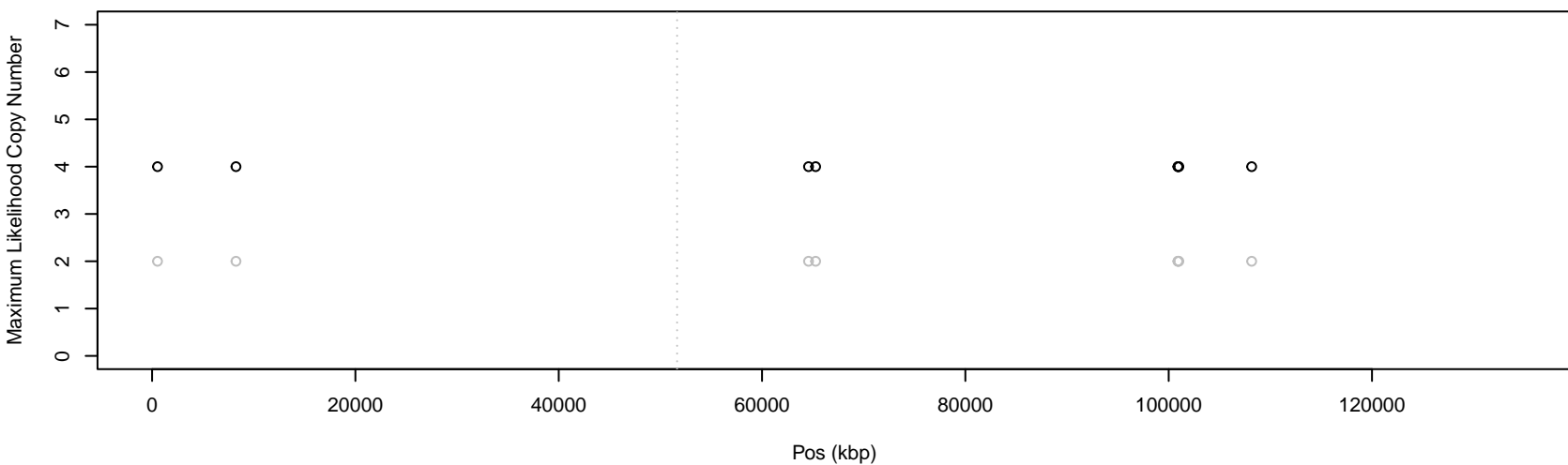
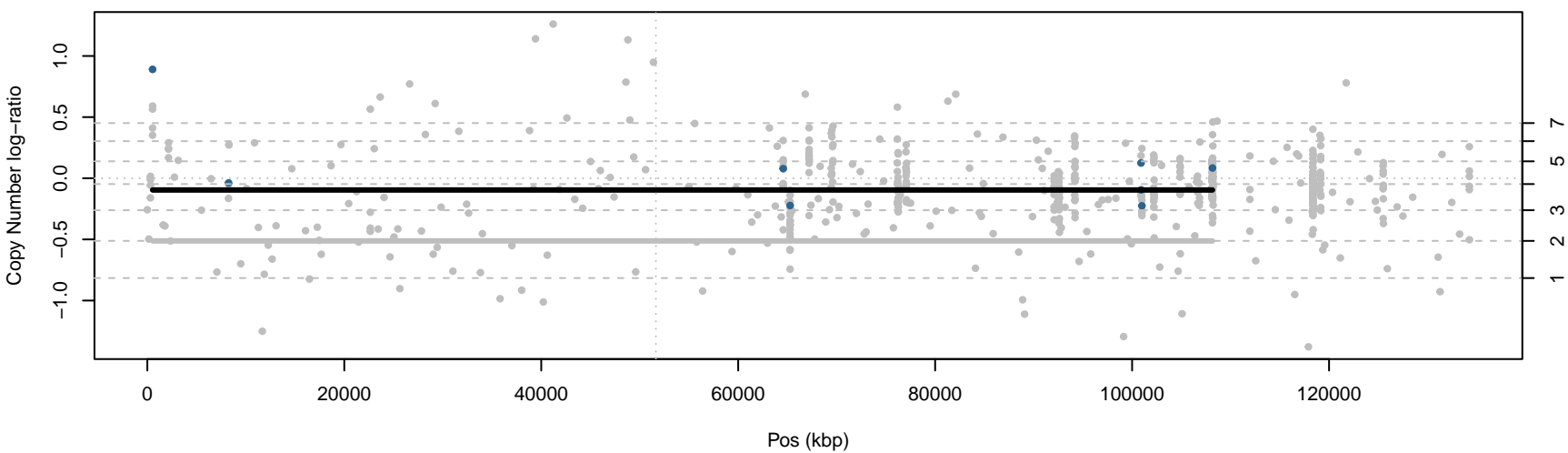
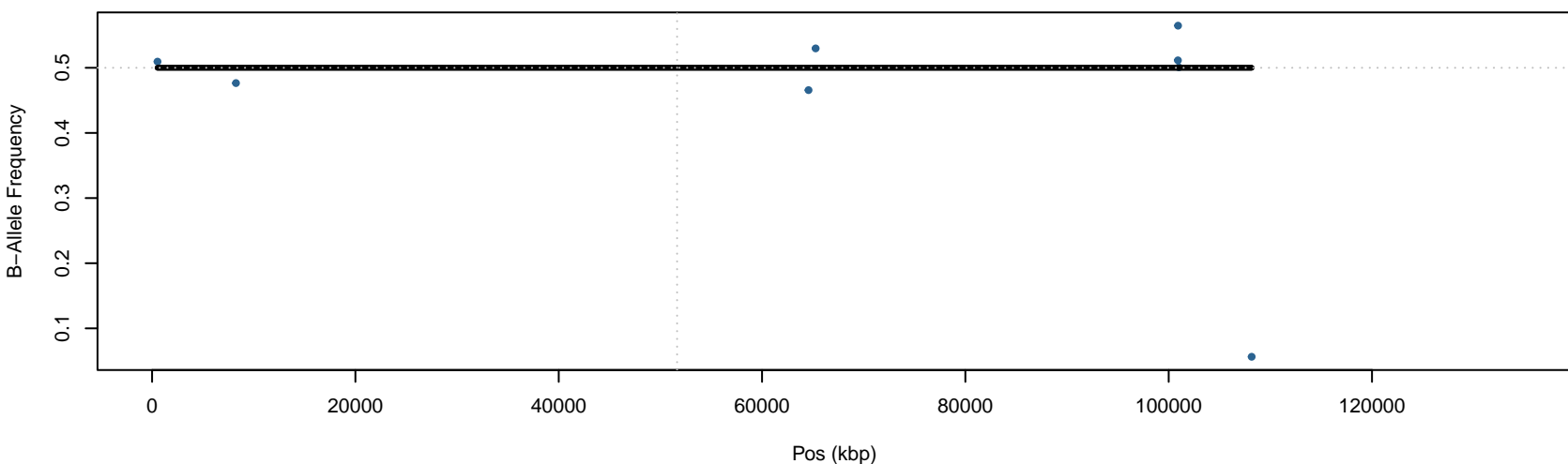
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 9



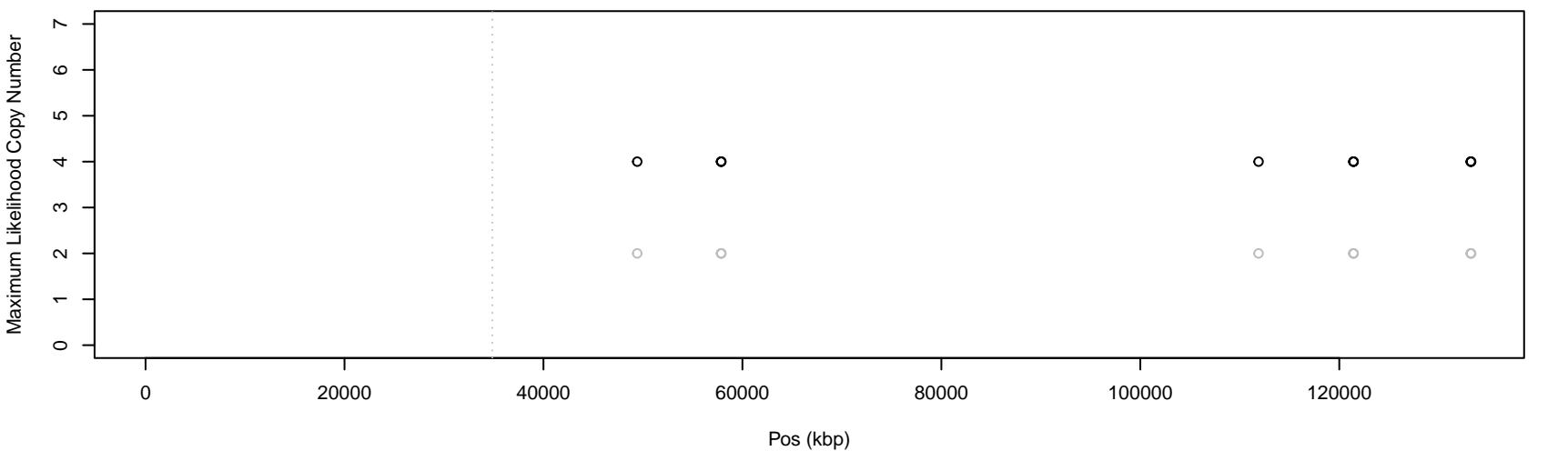
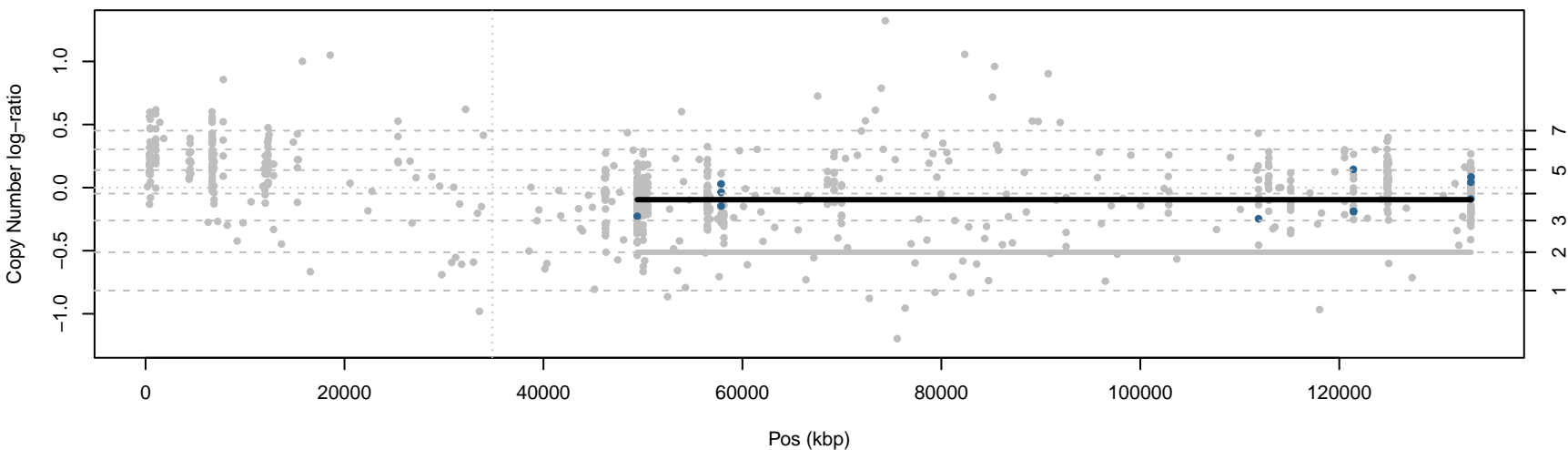
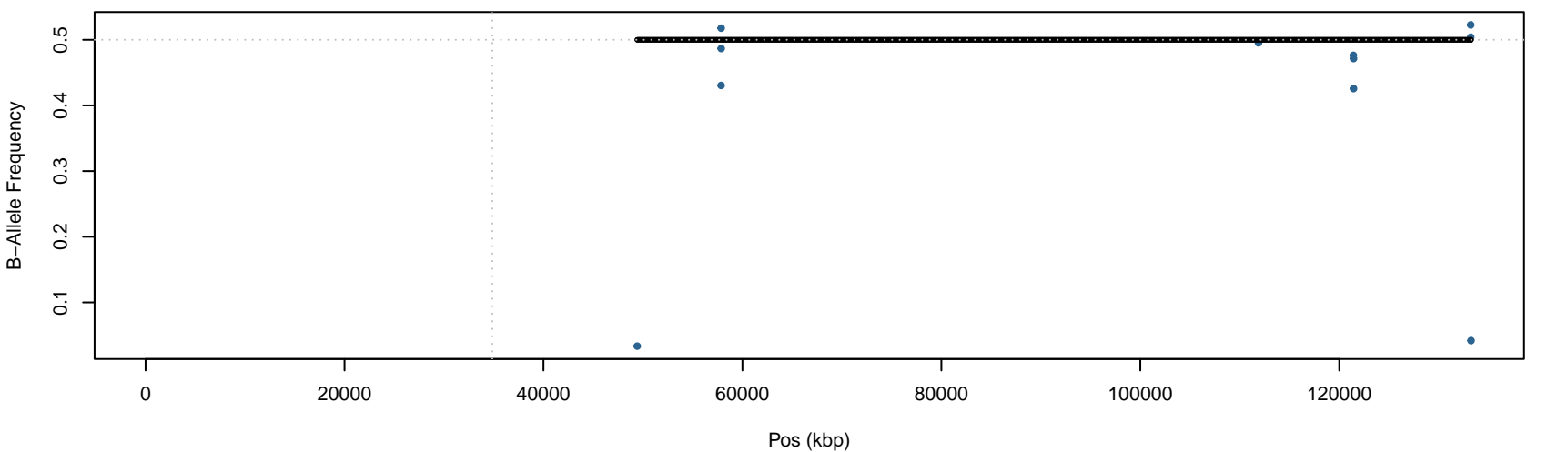
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 10



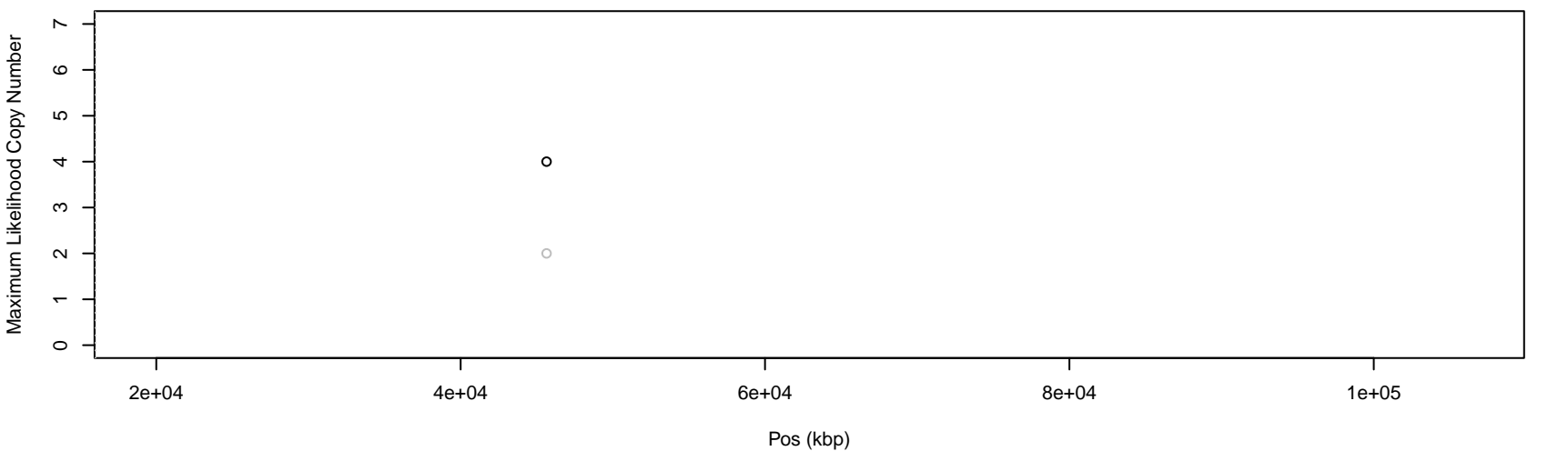
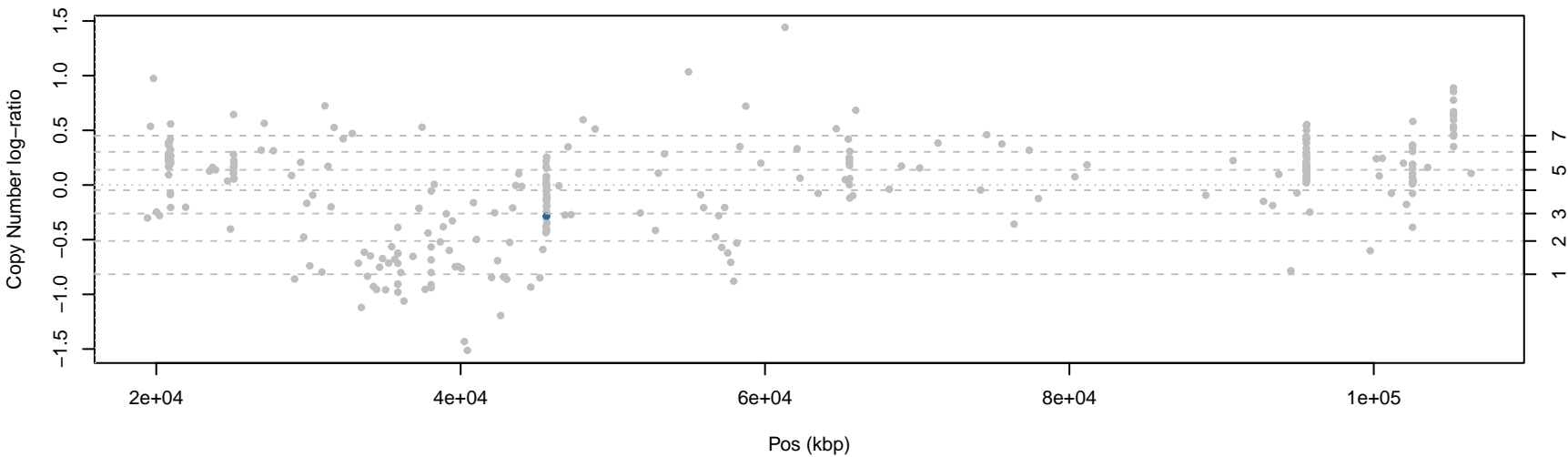
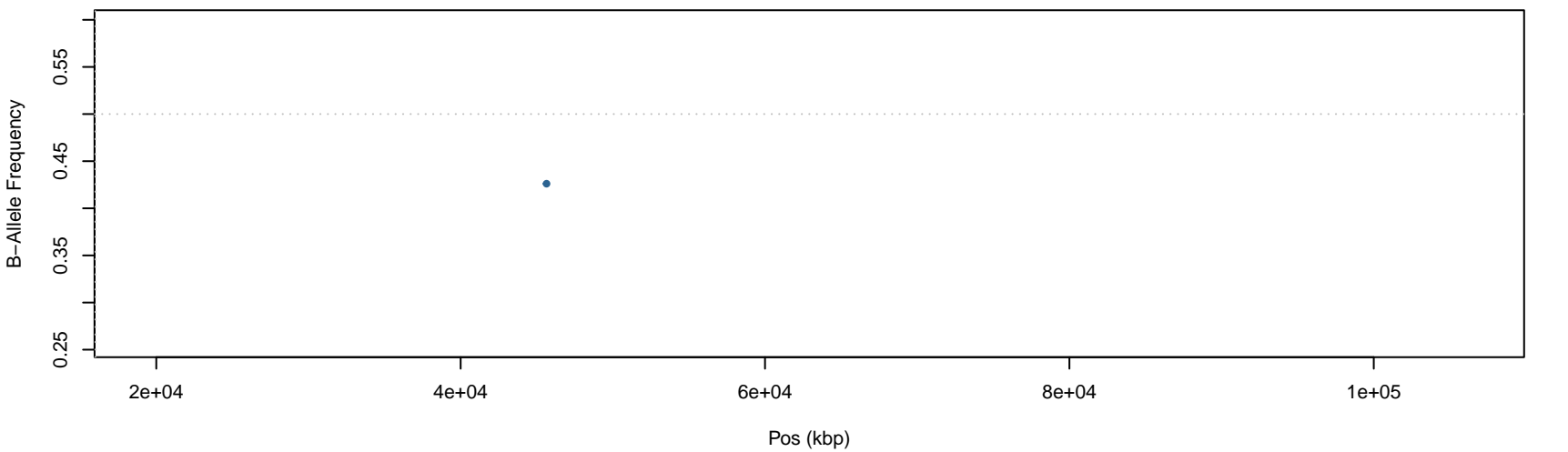
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 11



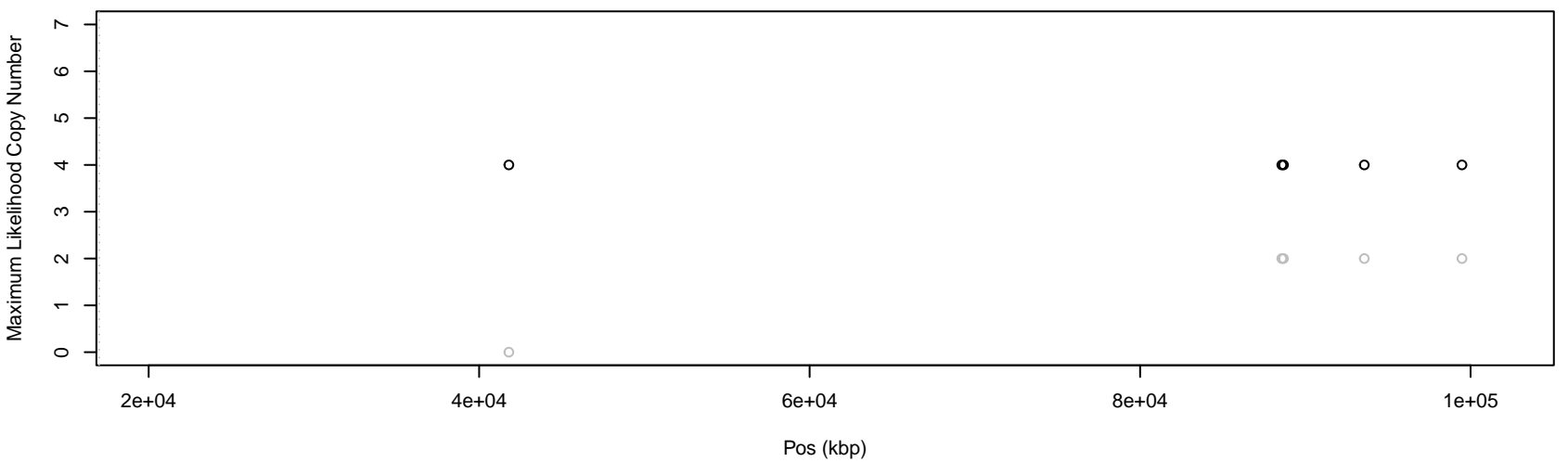
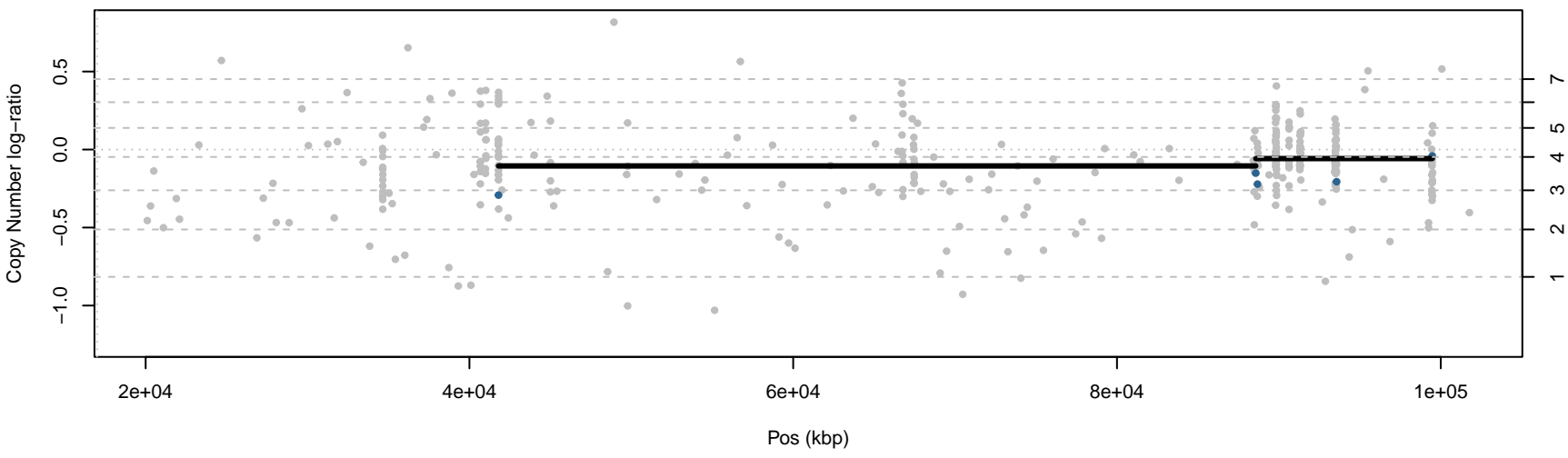
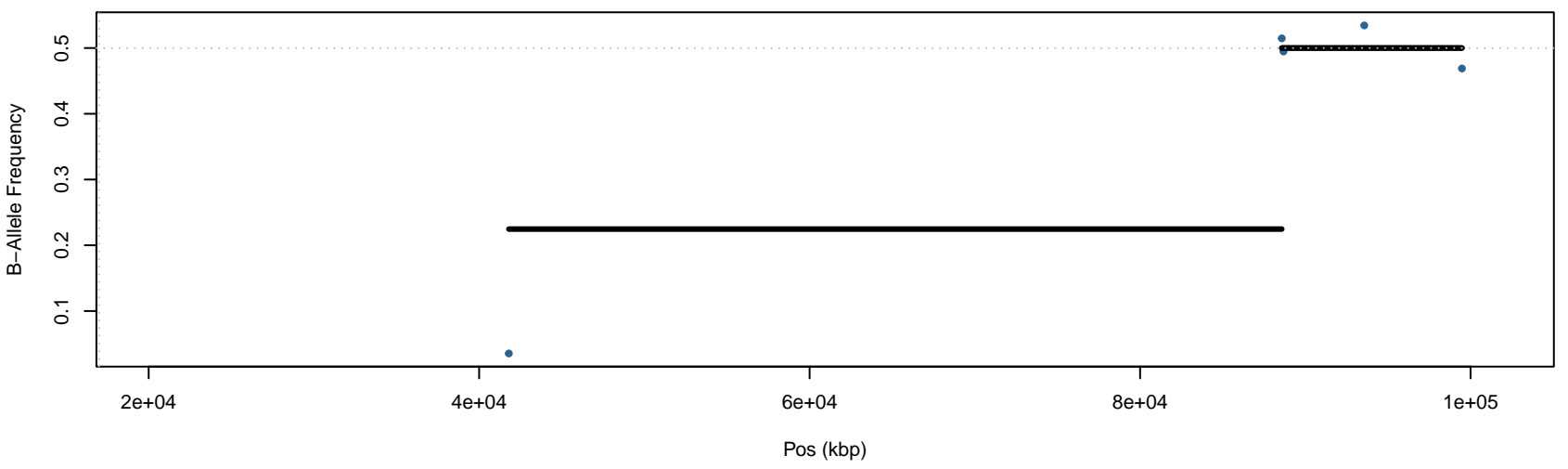
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 12



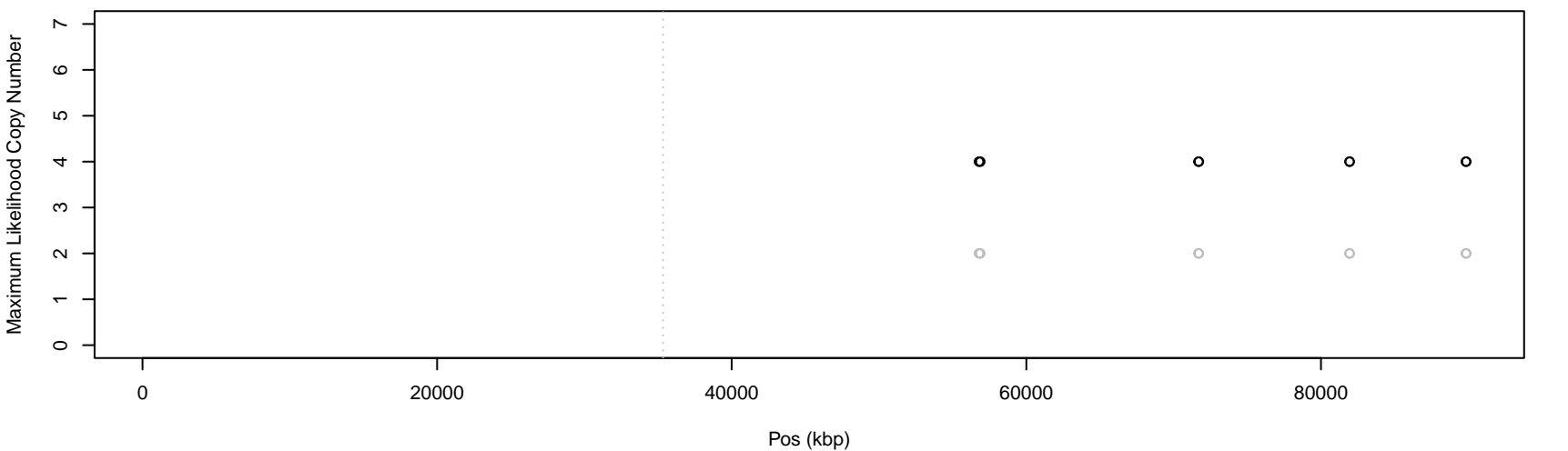
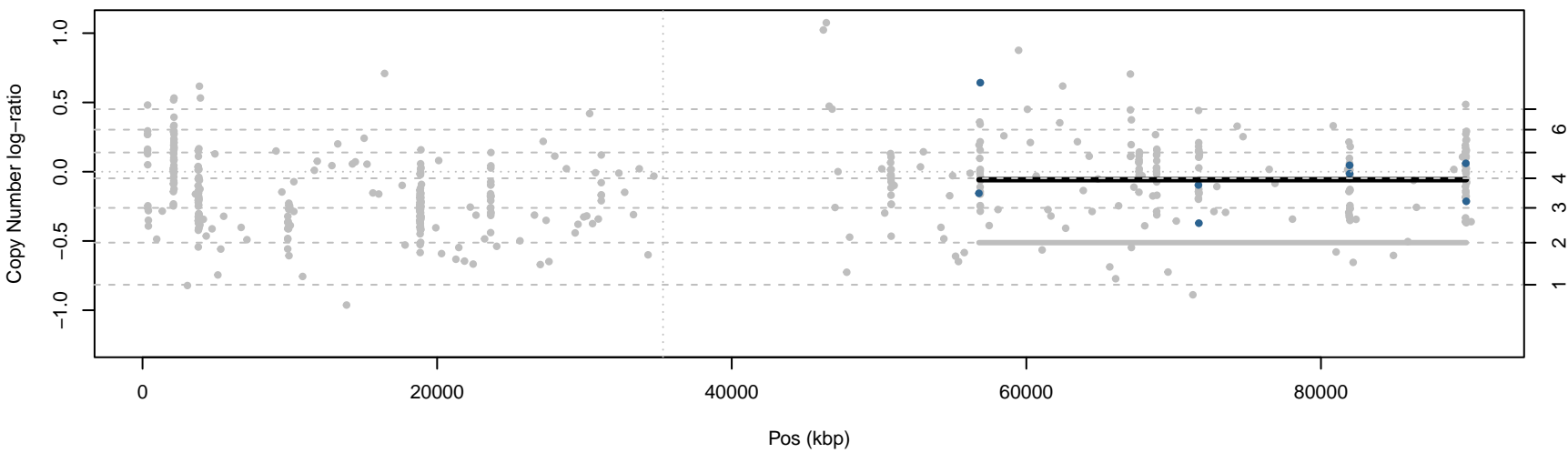
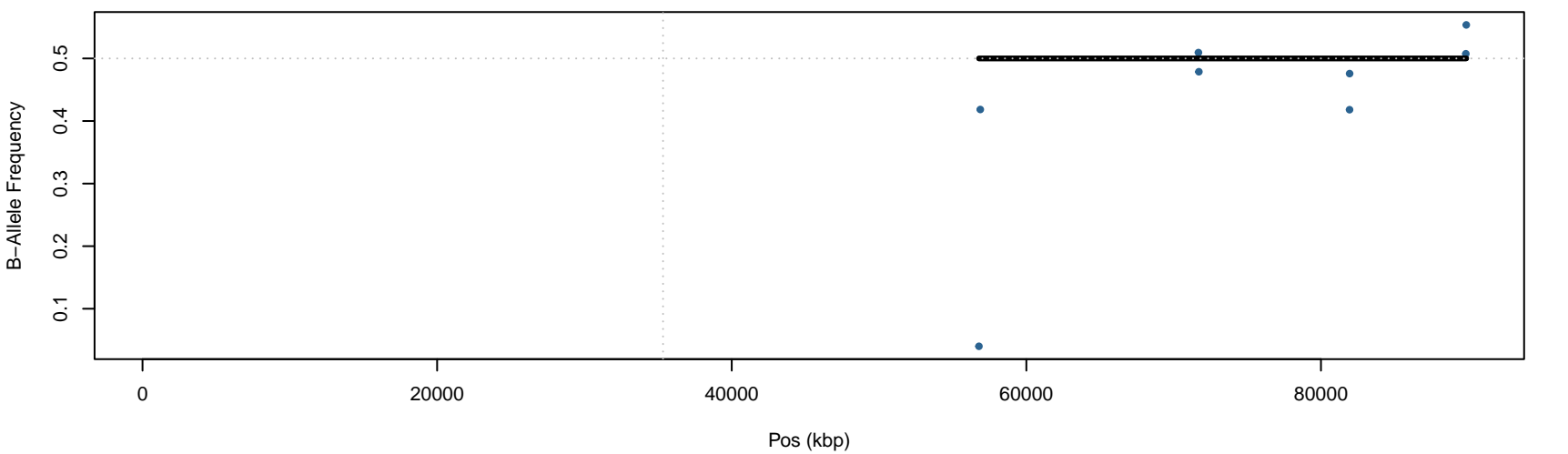
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 14



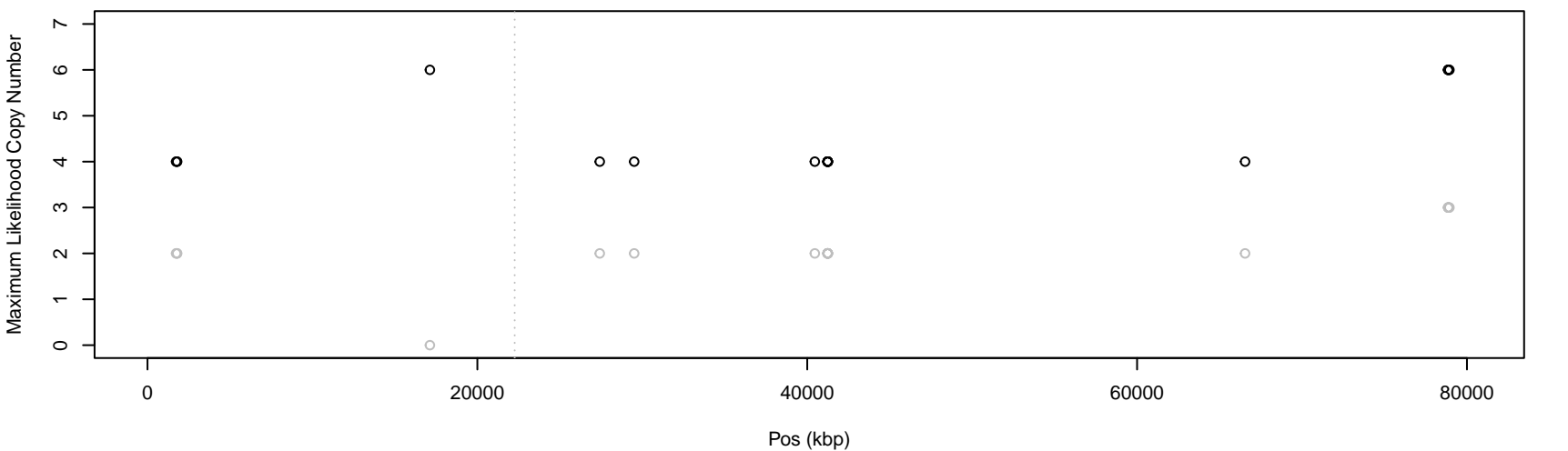
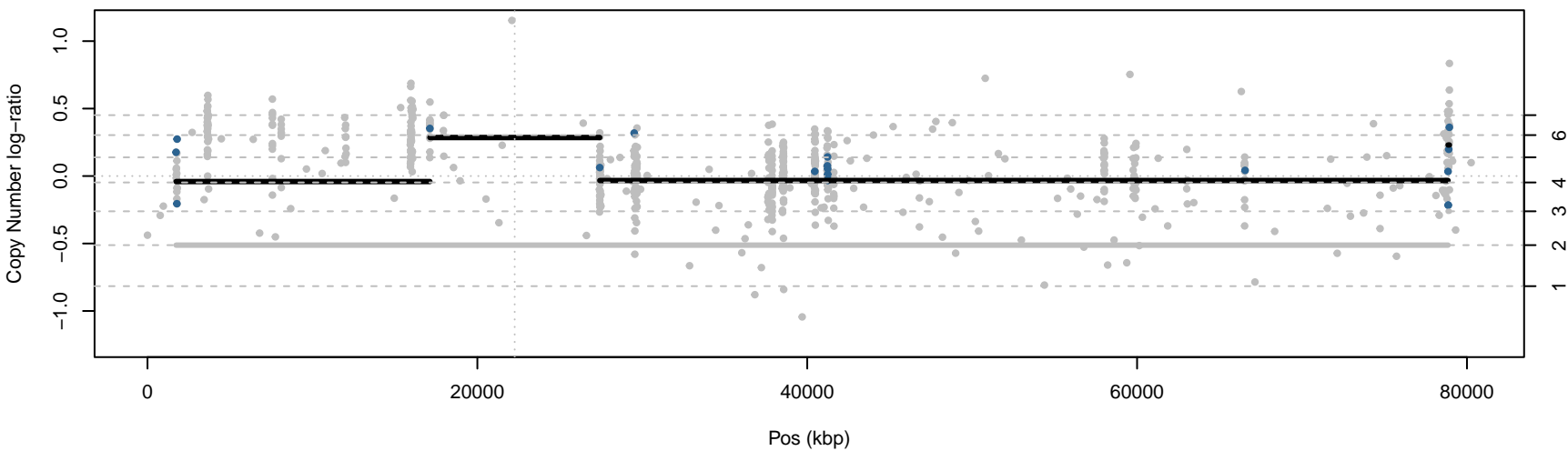
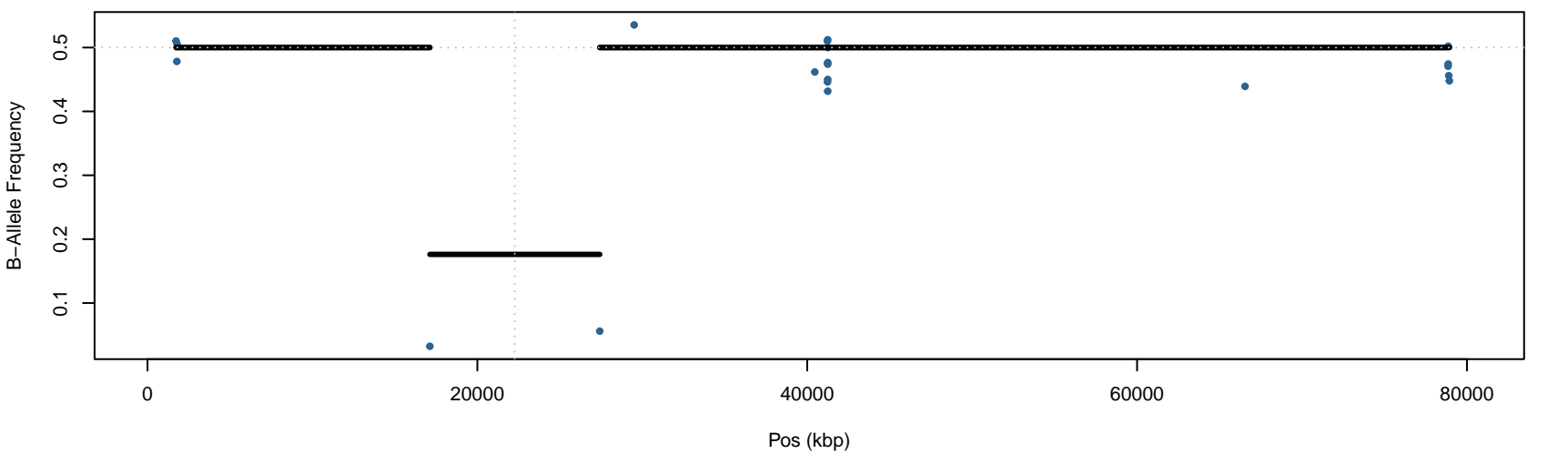
Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 15



Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 16

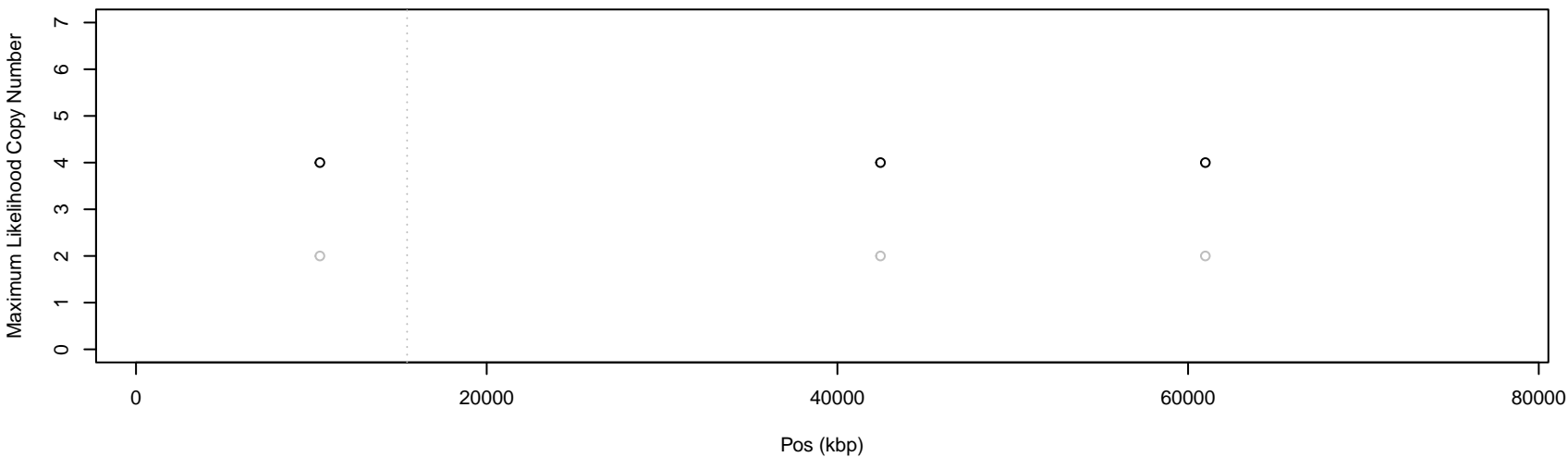
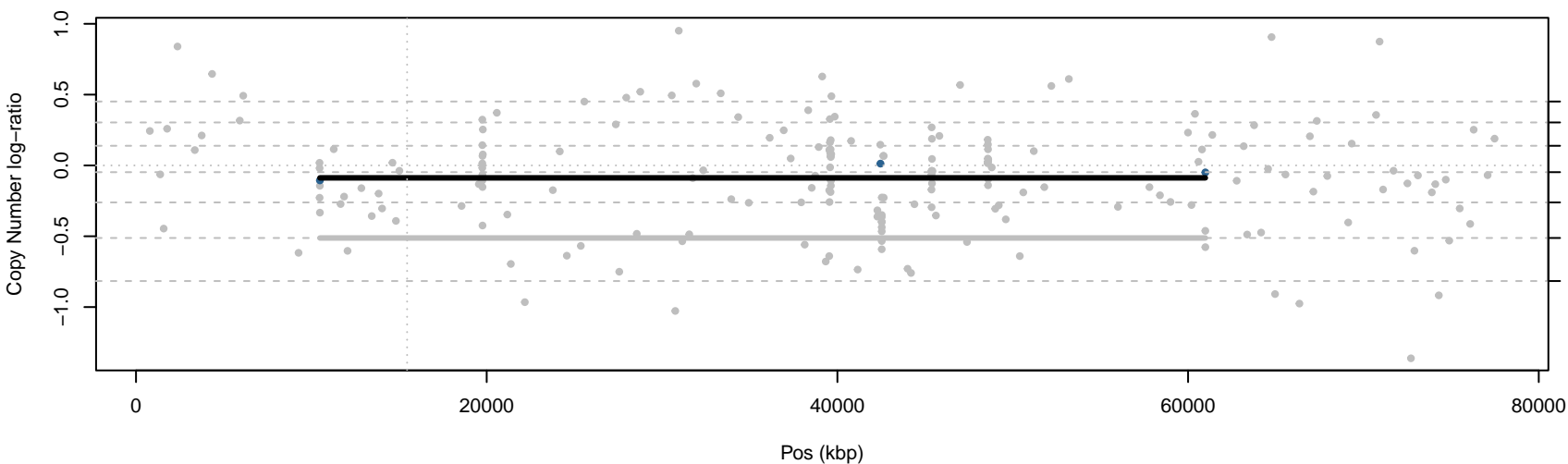
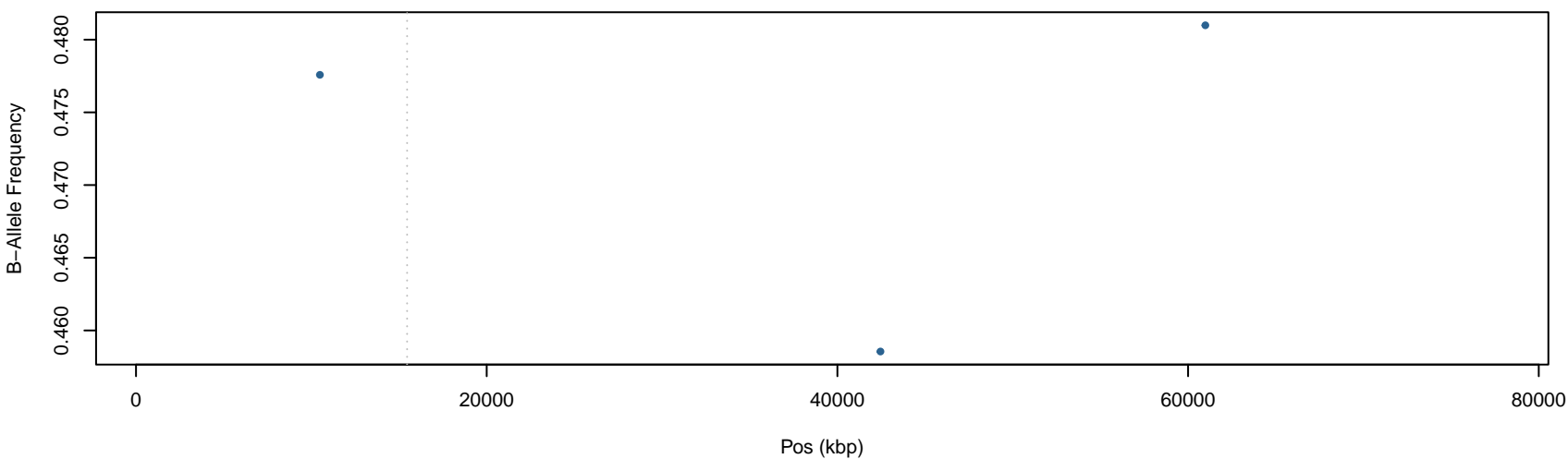


Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 17

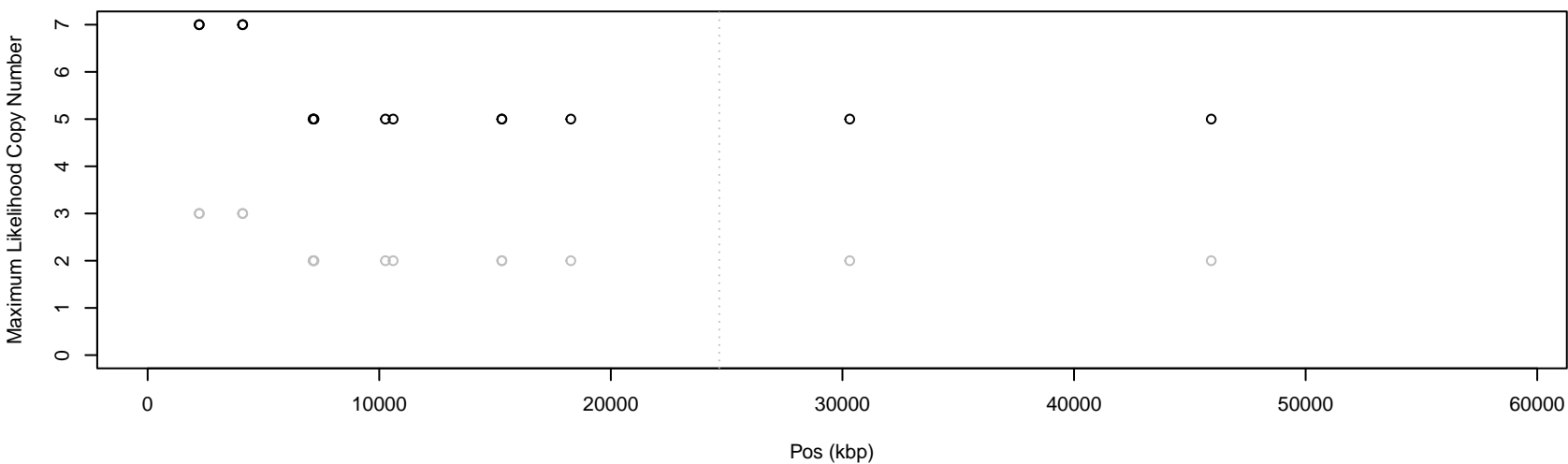
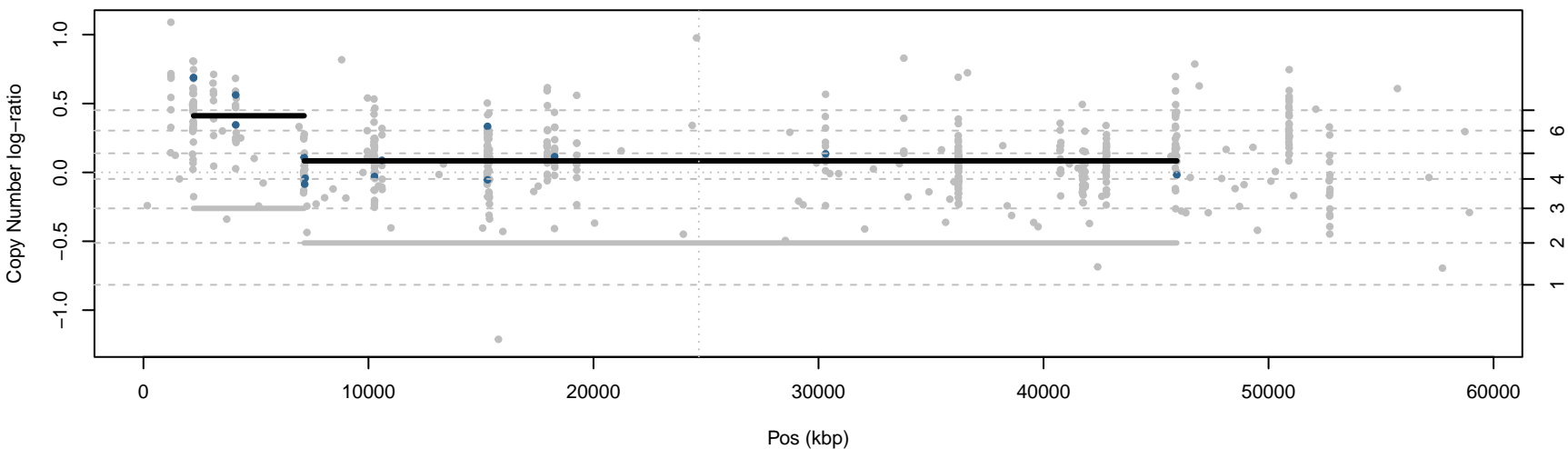
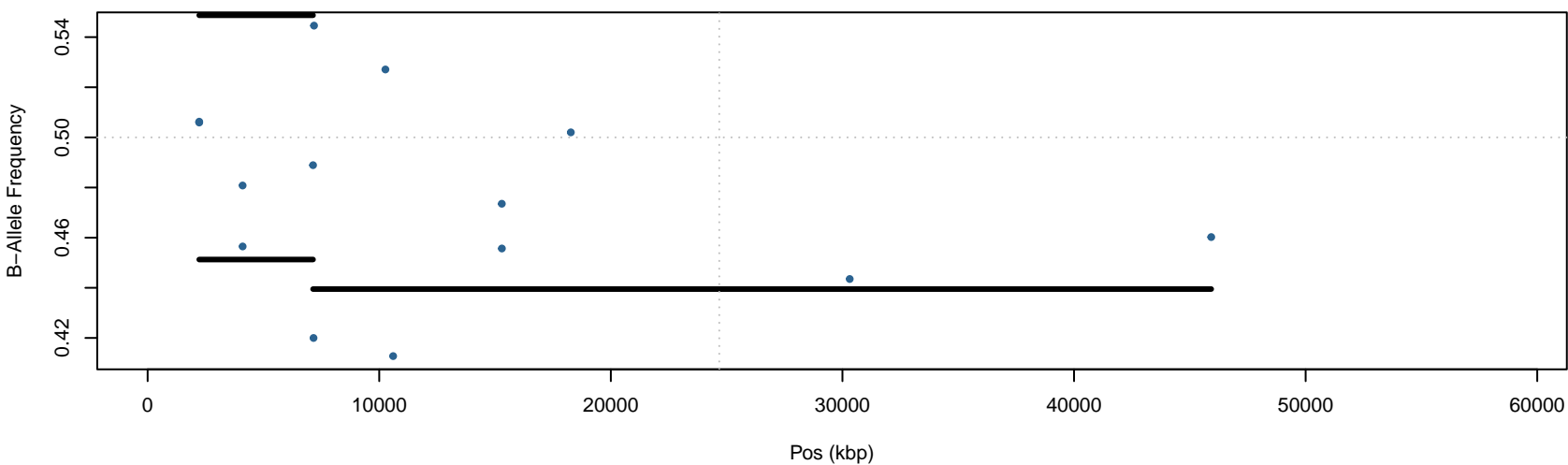




Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 18



Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 19



Purity: 0.38 Tumor ploidy: 4.243 SNV log-likelihood: 2.82 GoF: 93.7% Mean coverage: 889;627 Chromosome: 21

