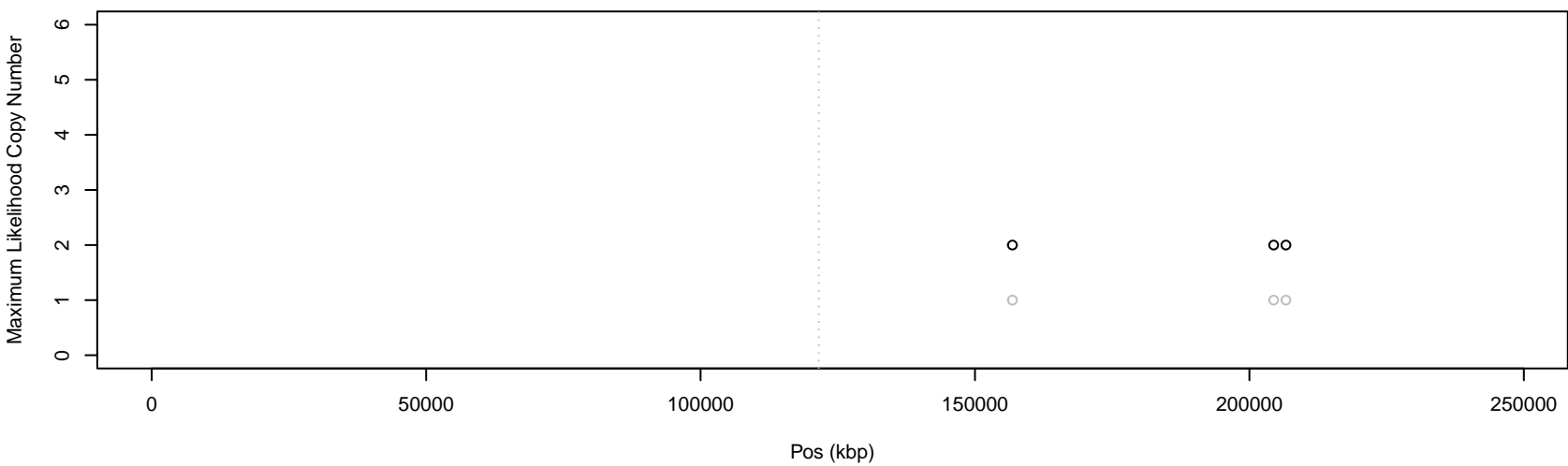
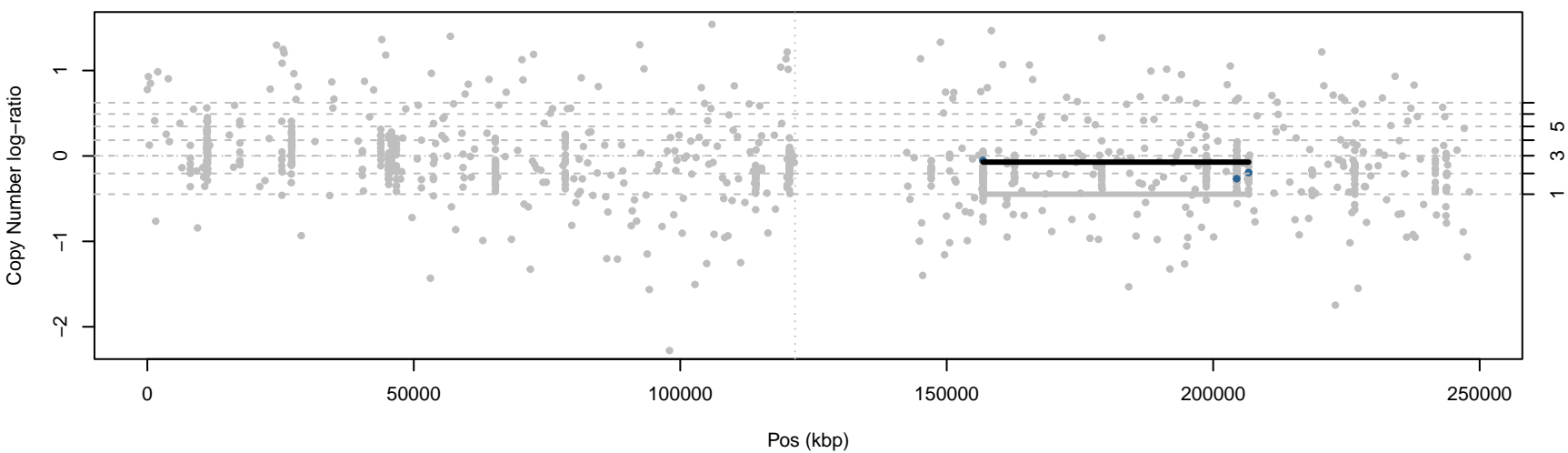
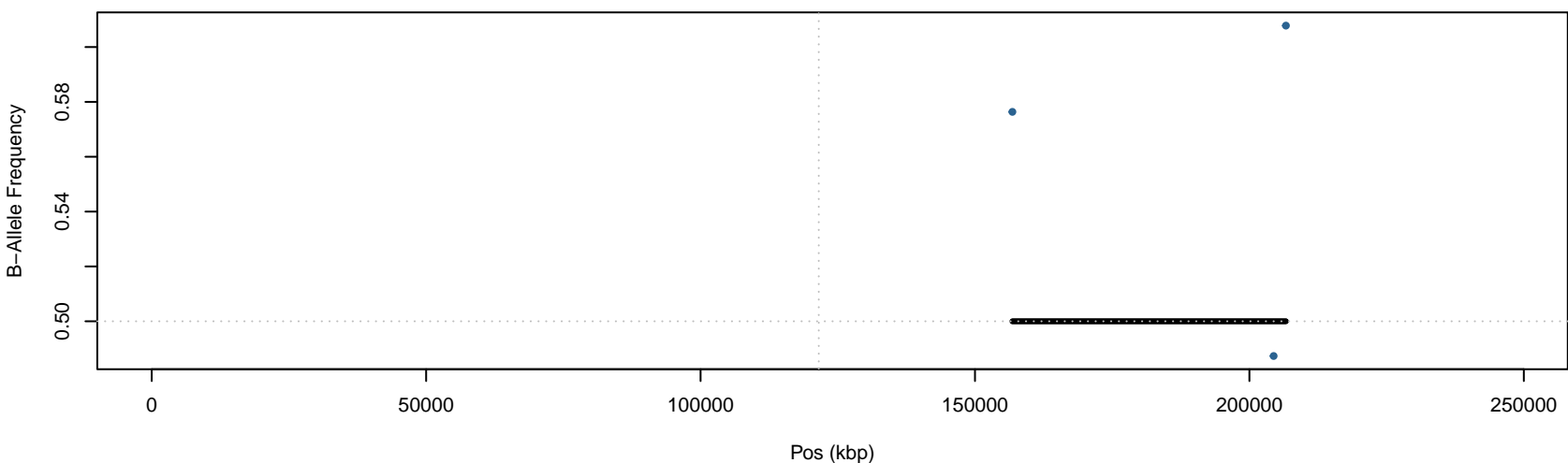
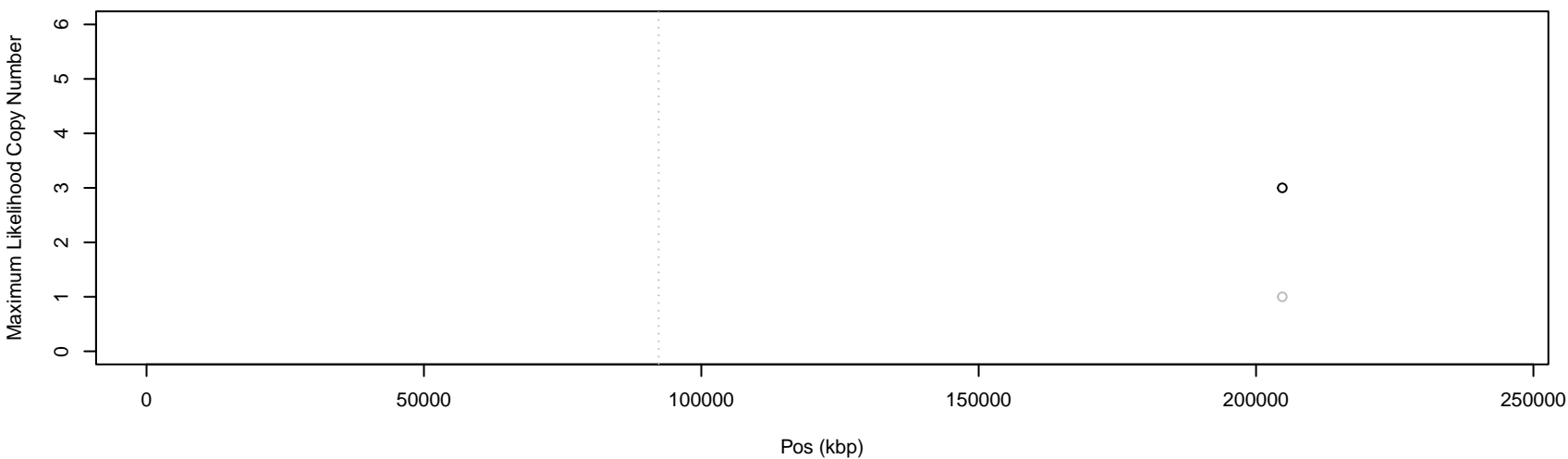
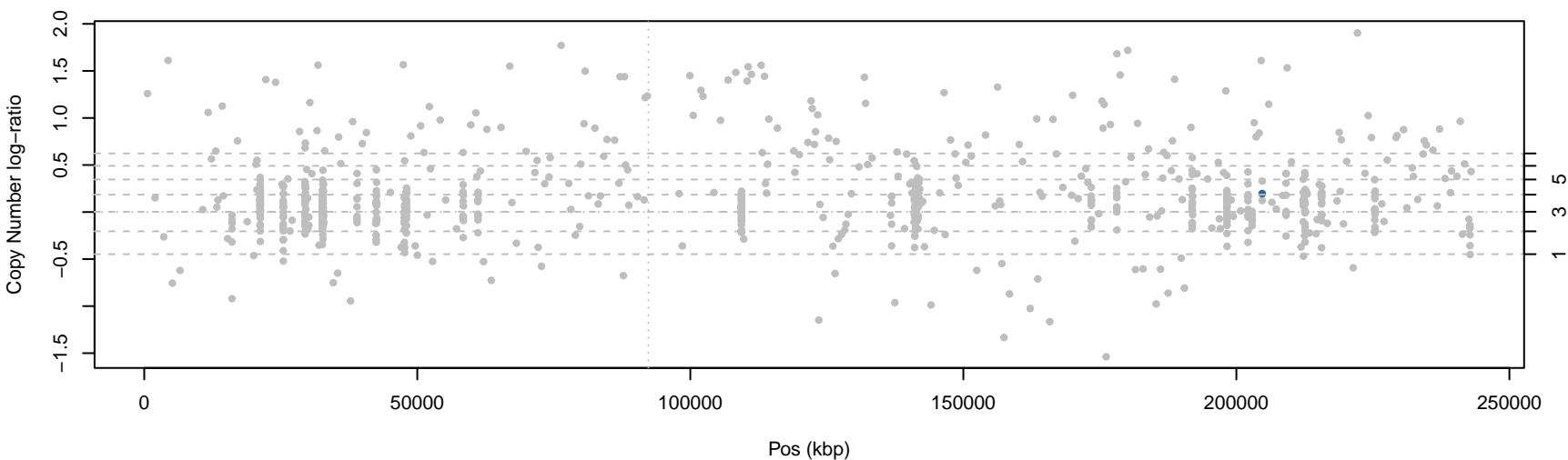
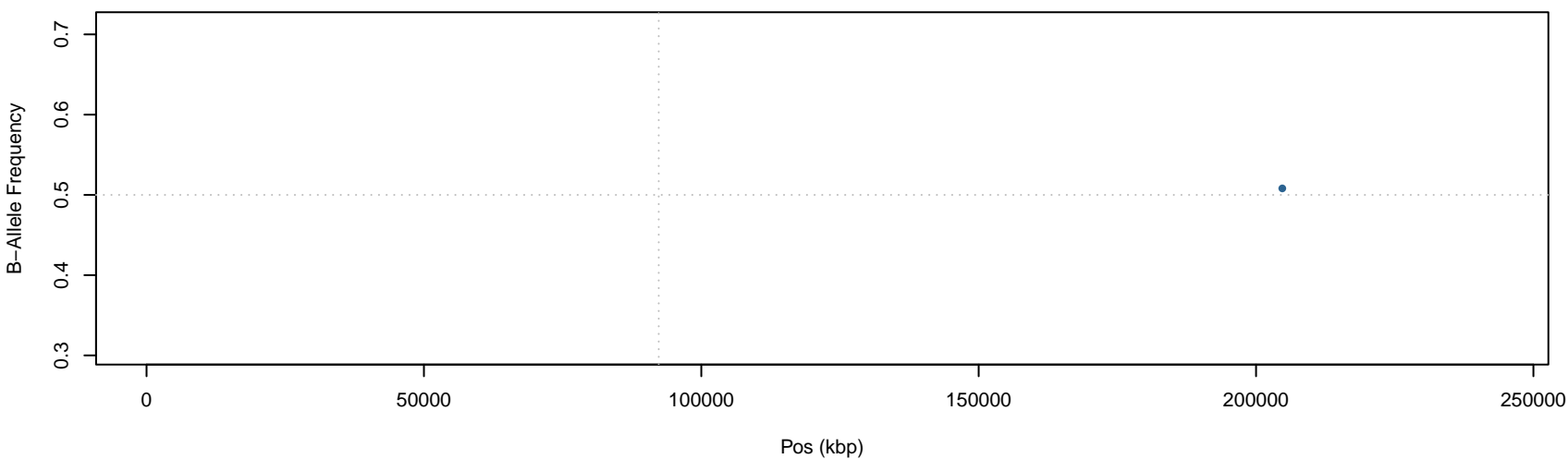


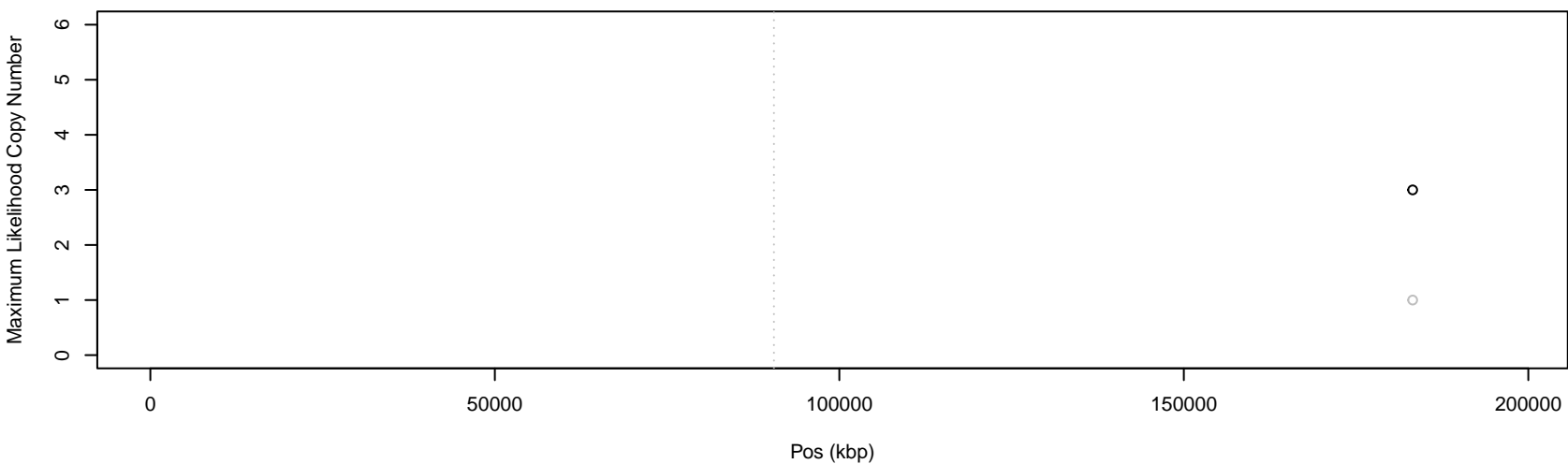
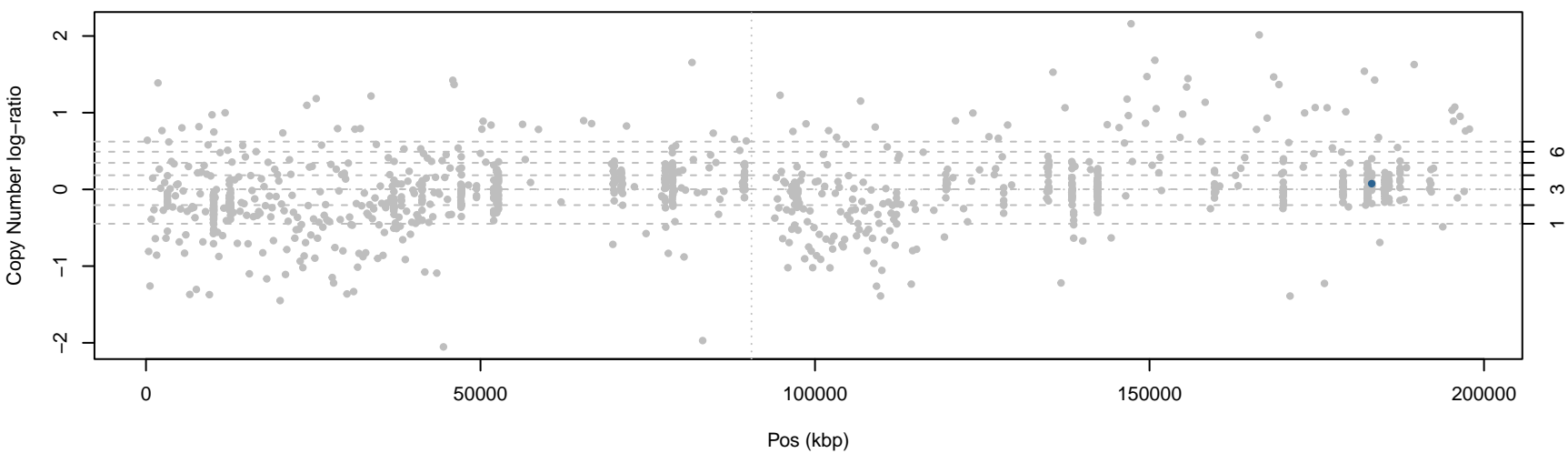
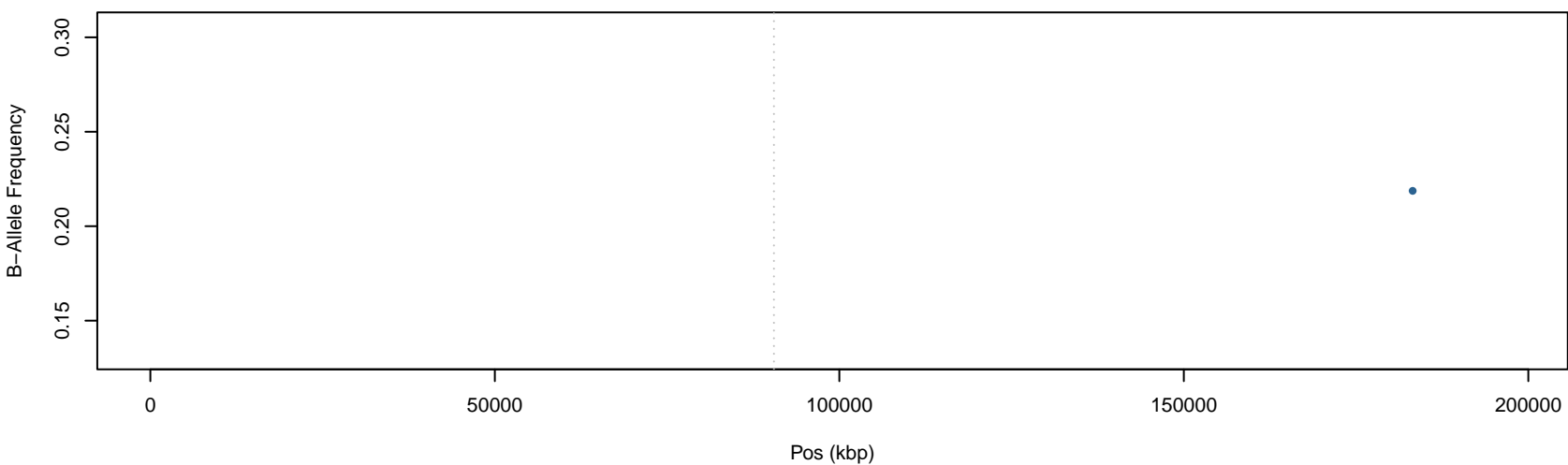
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 1



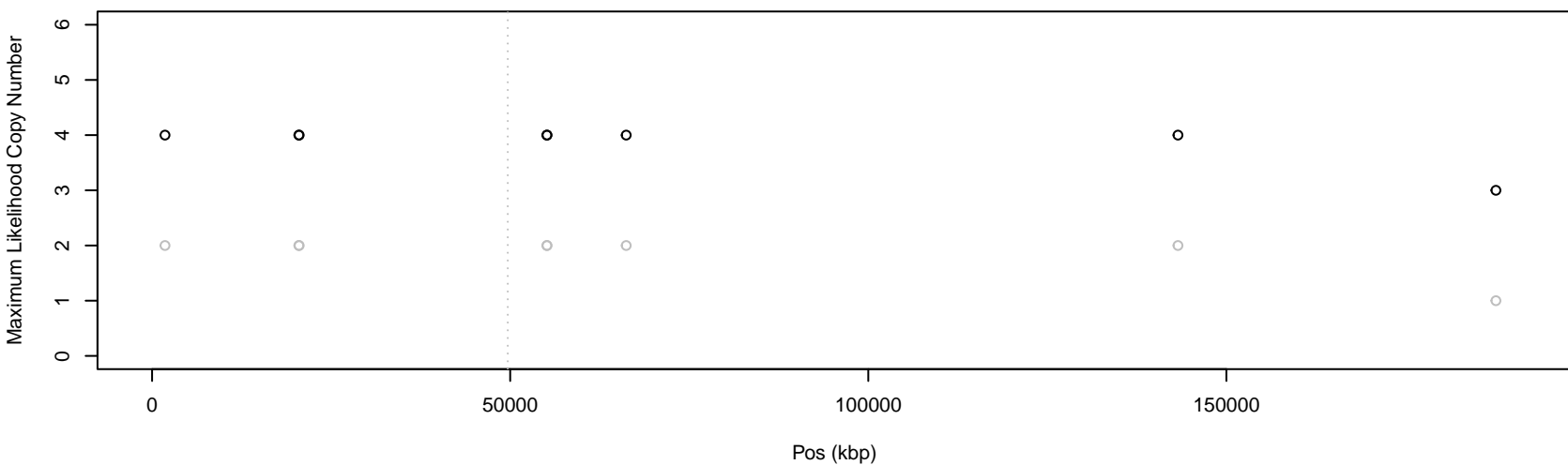
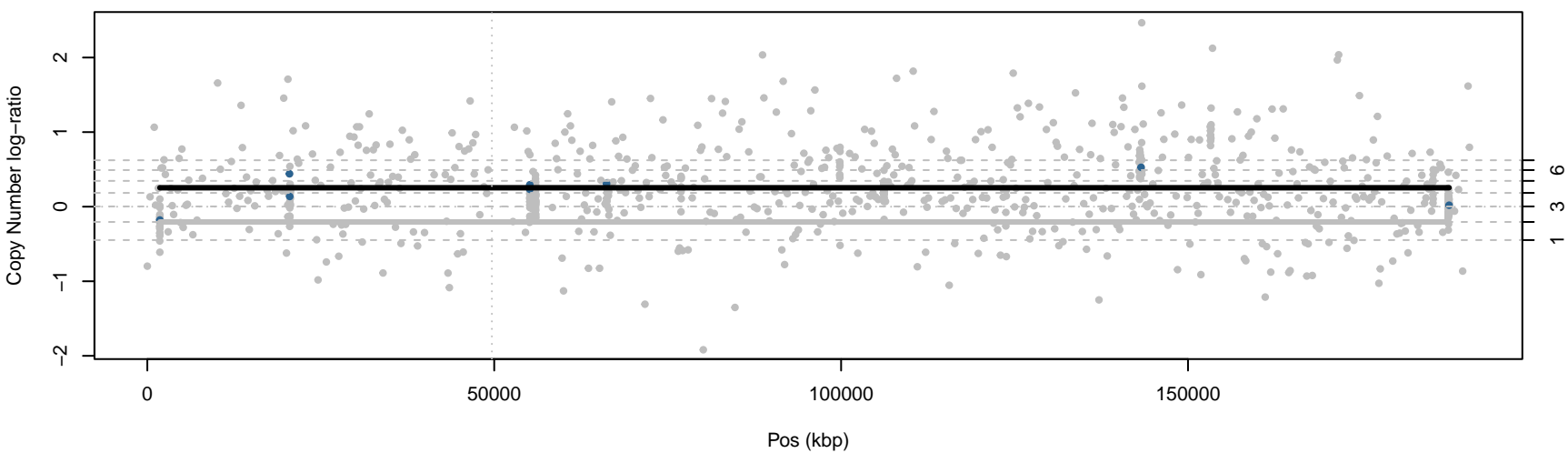
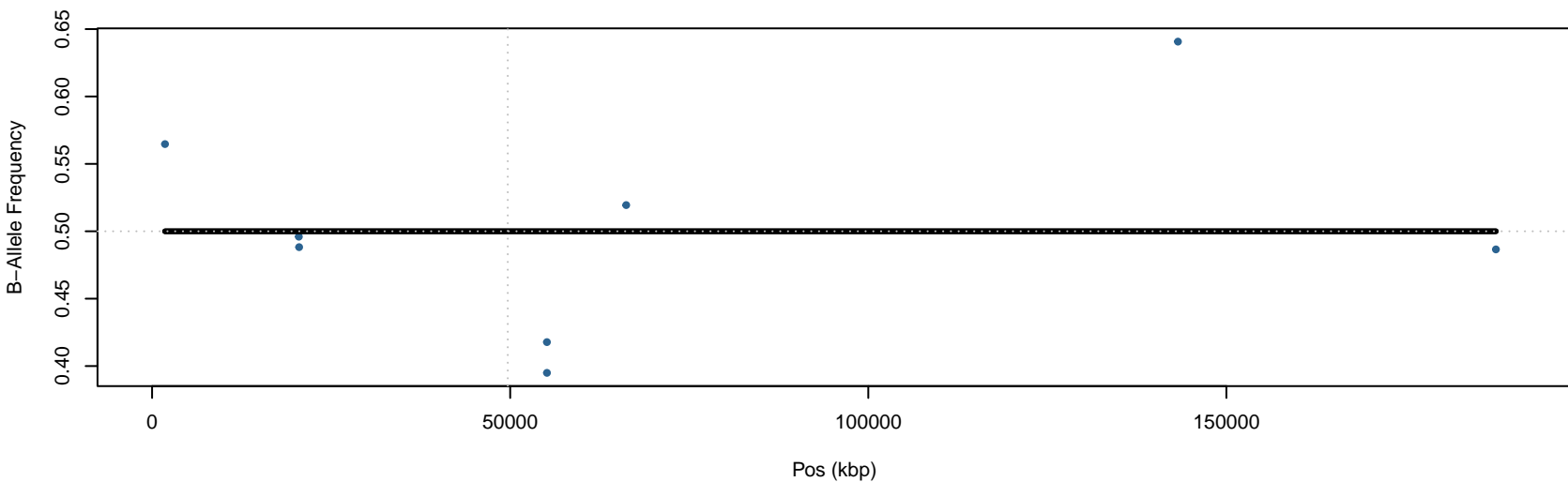
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 2



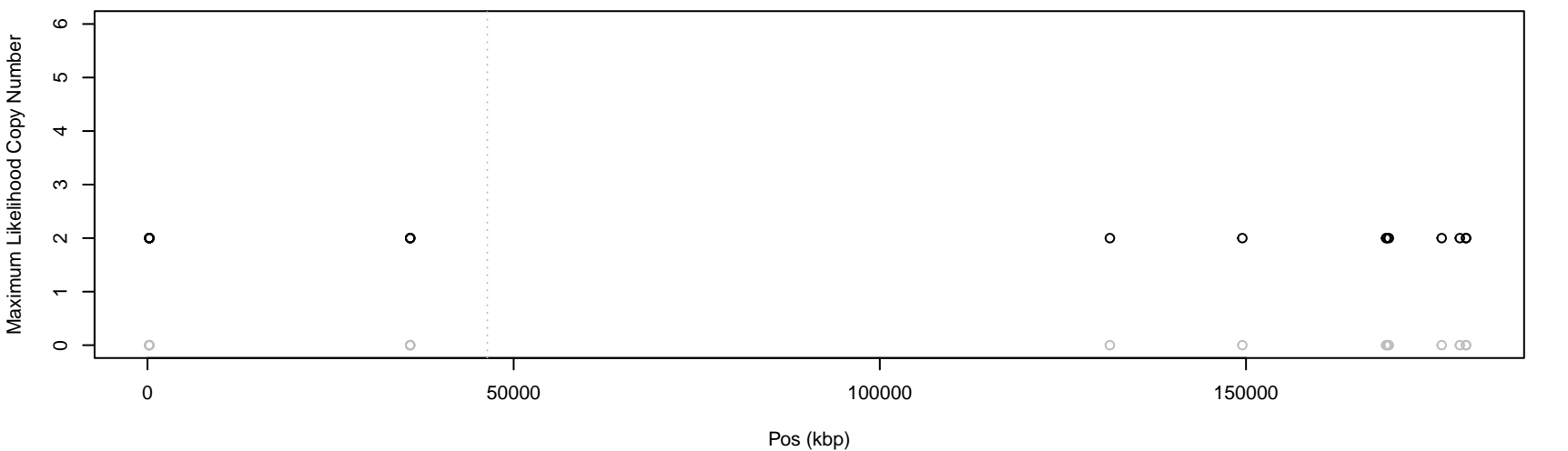
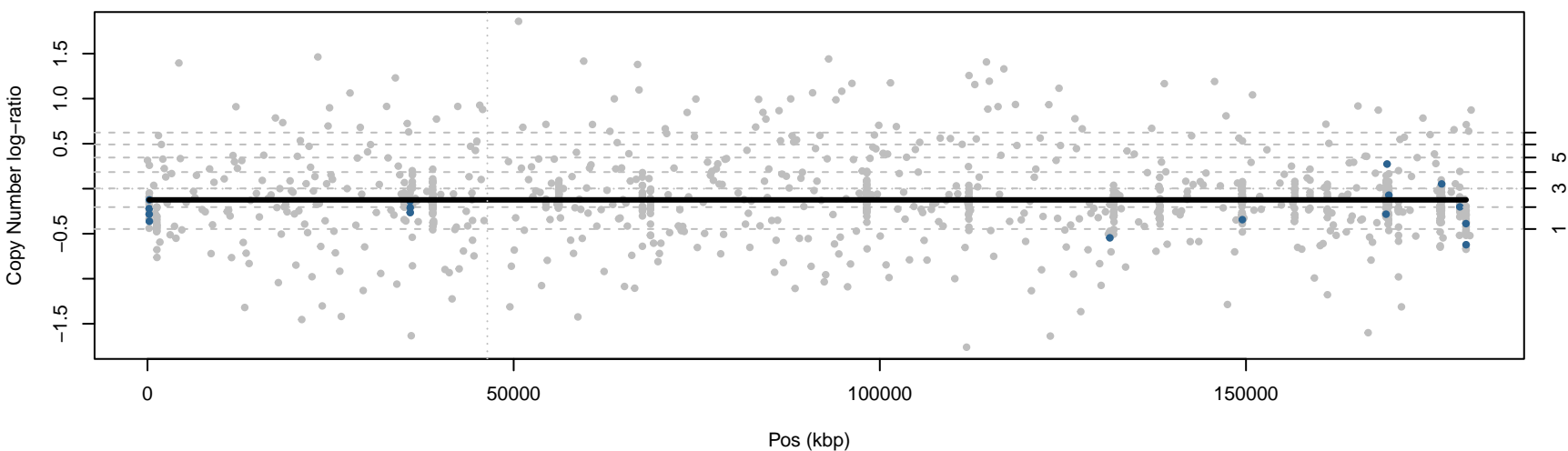
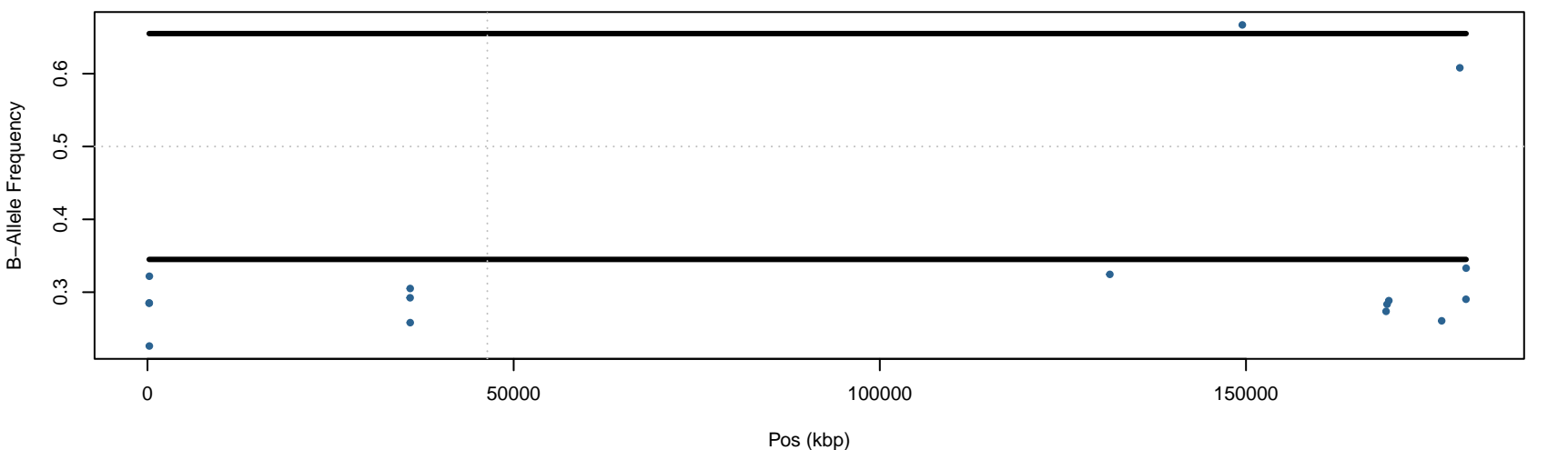
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 3



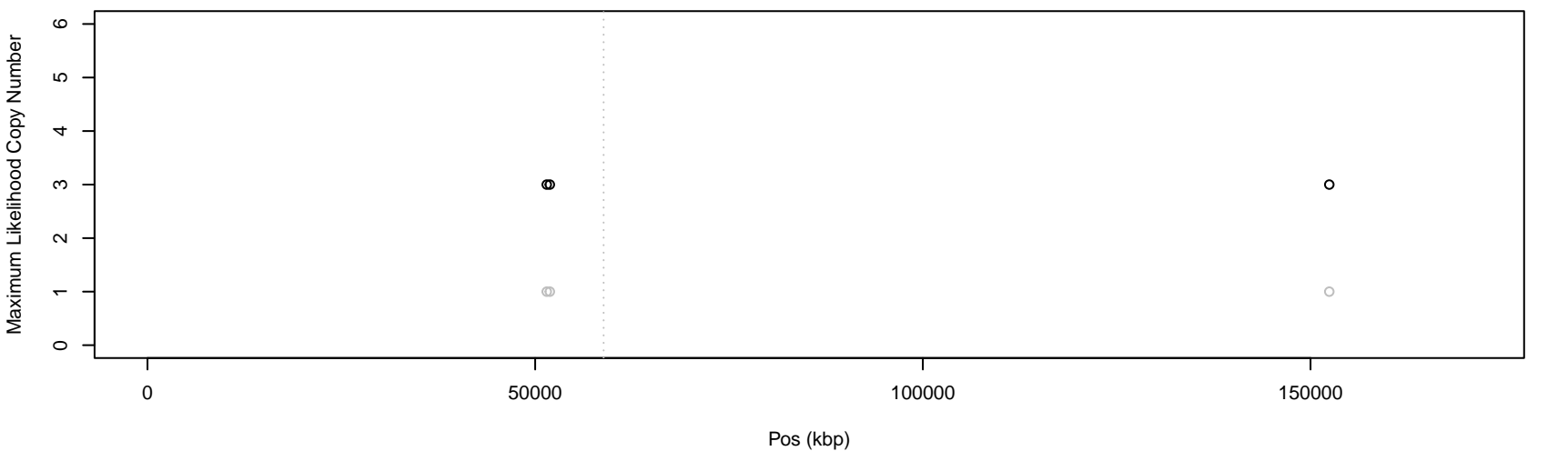
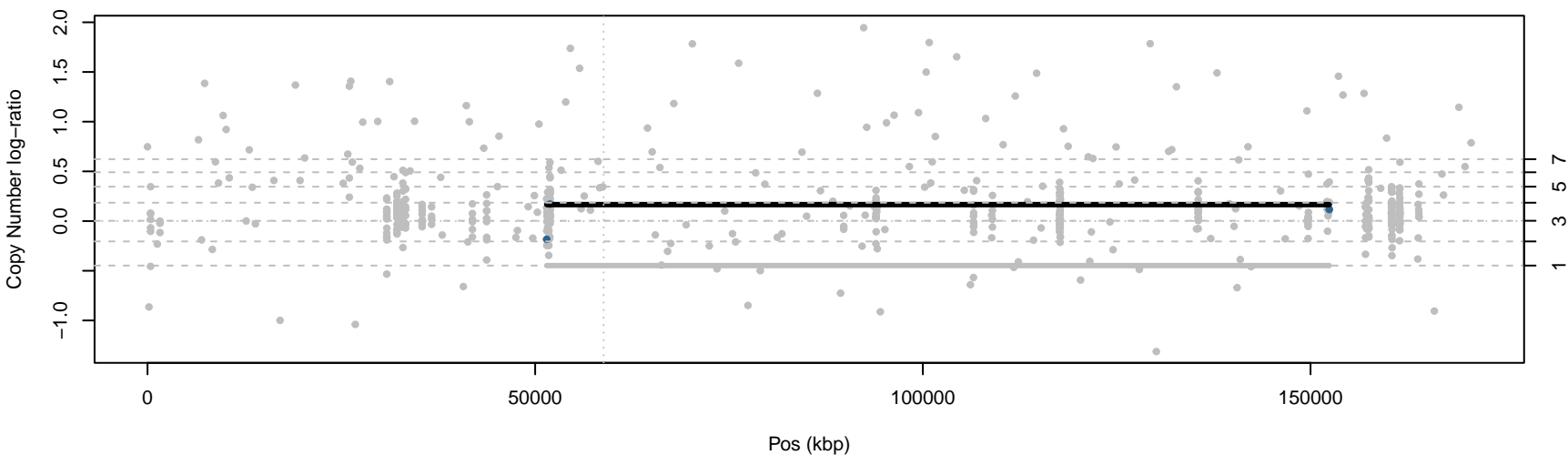
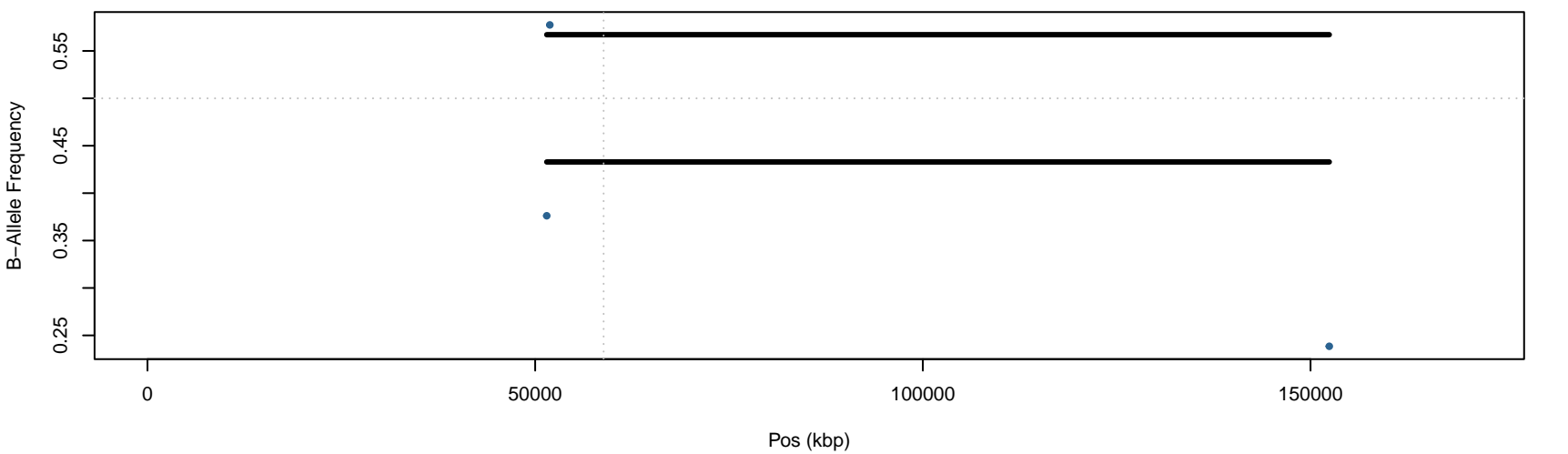
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 4



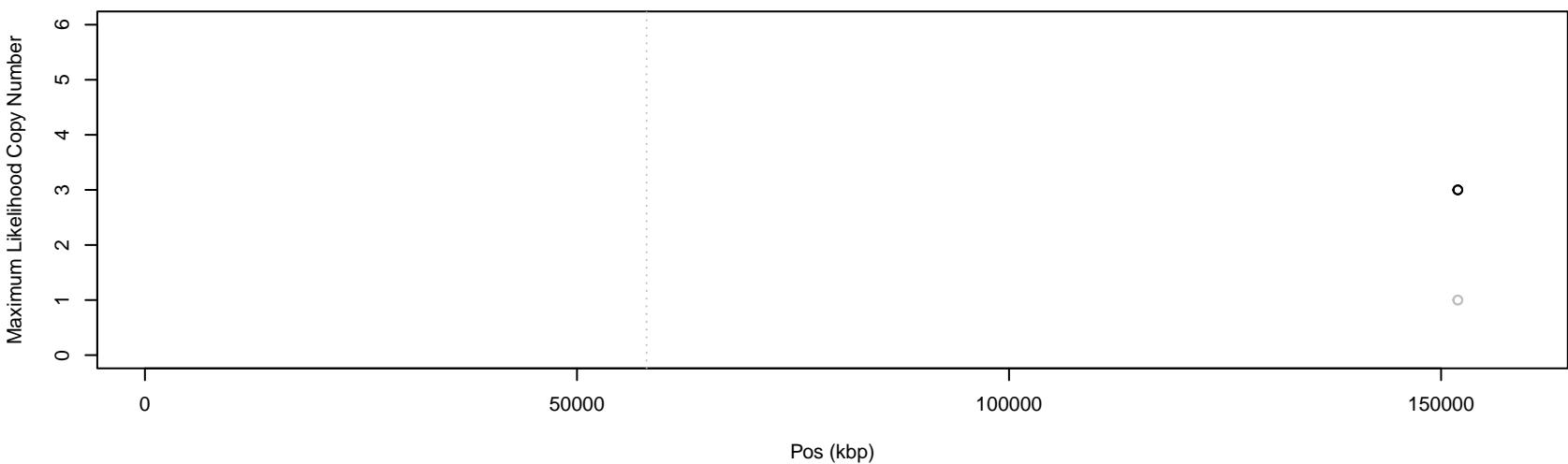
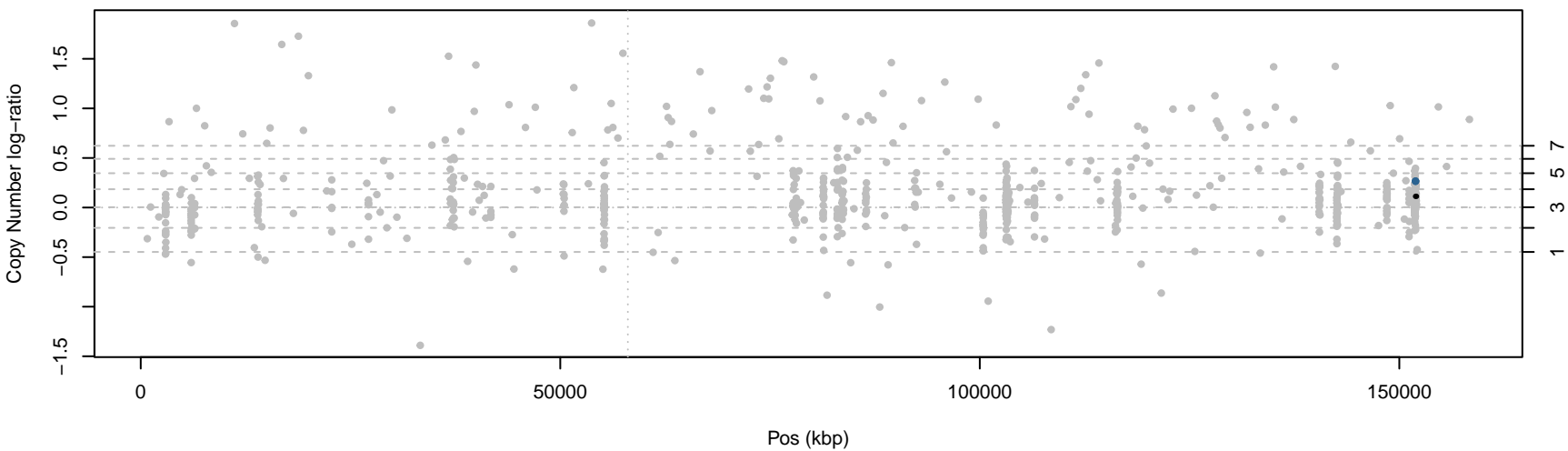
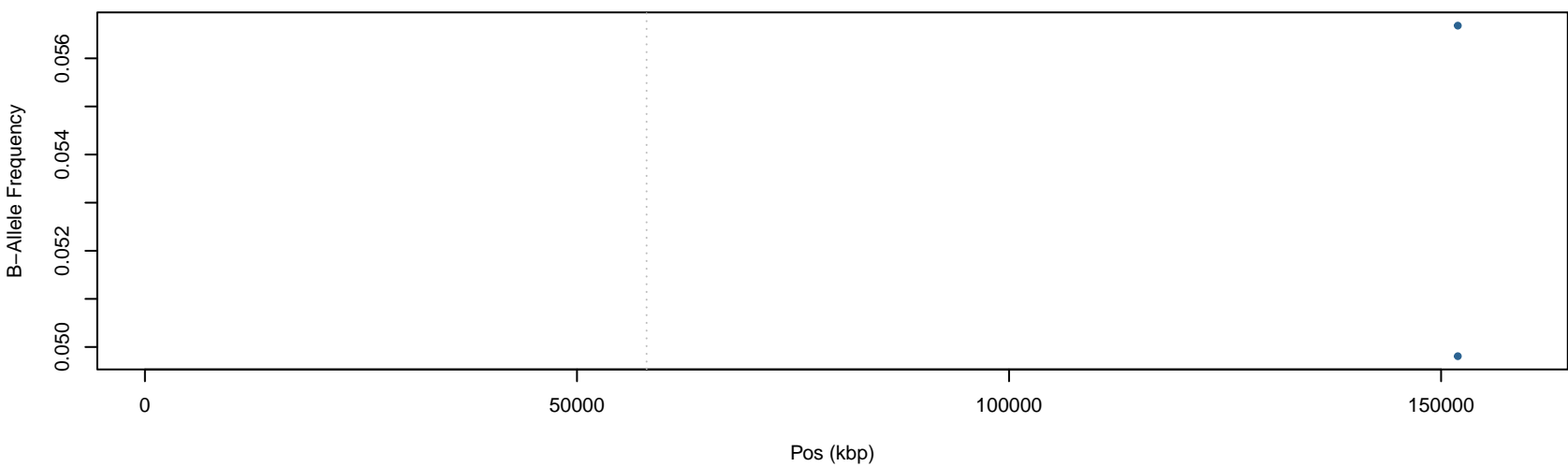
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 5



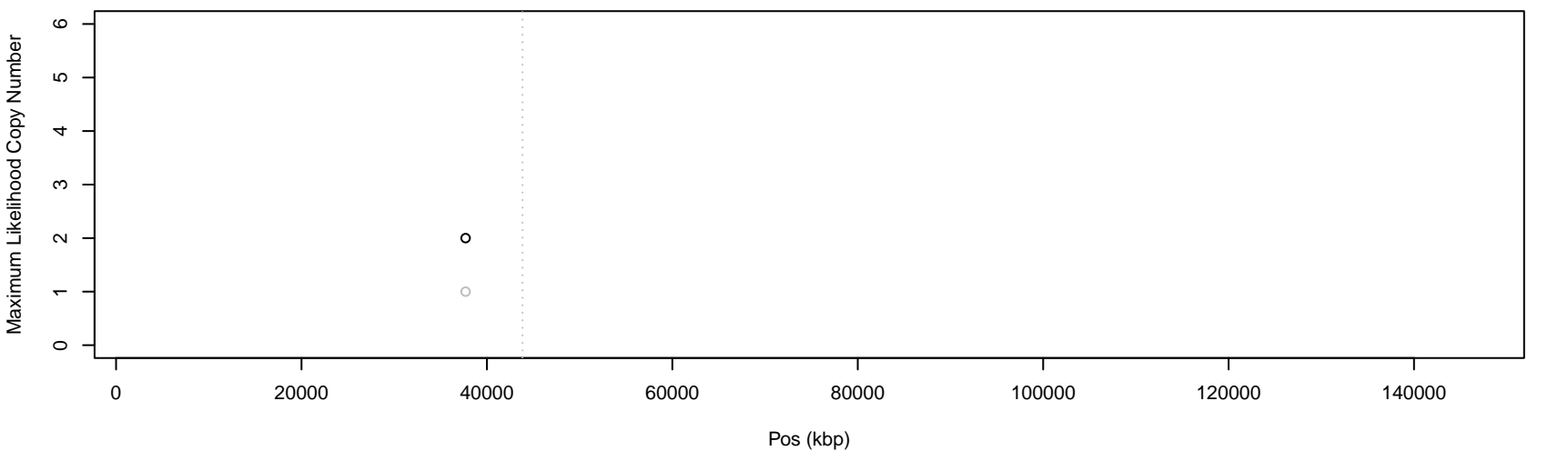
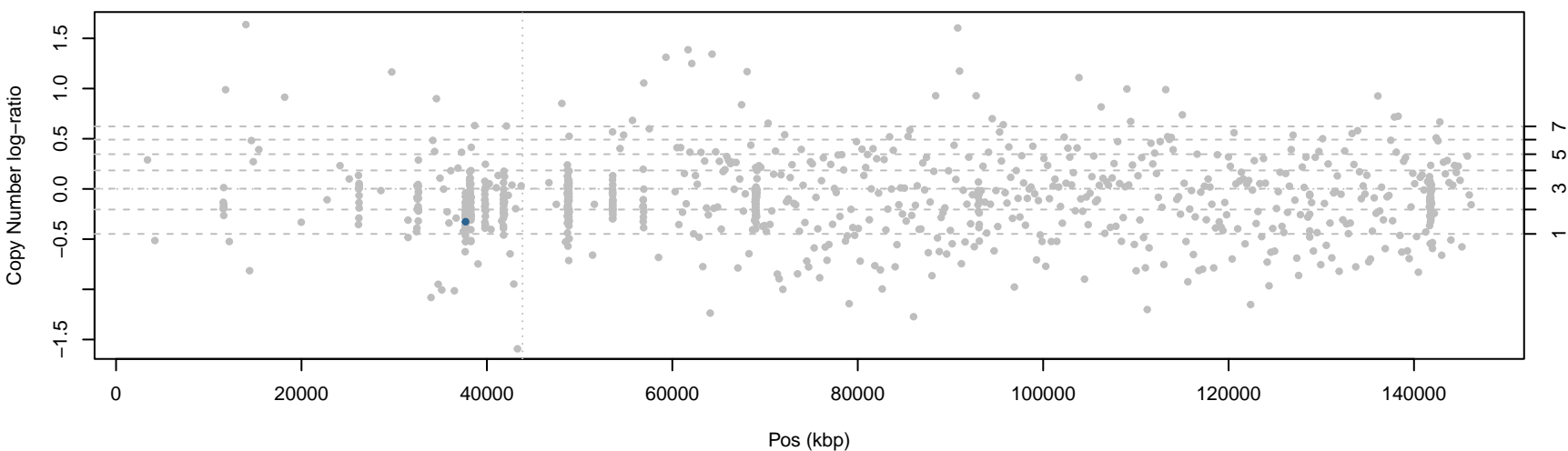
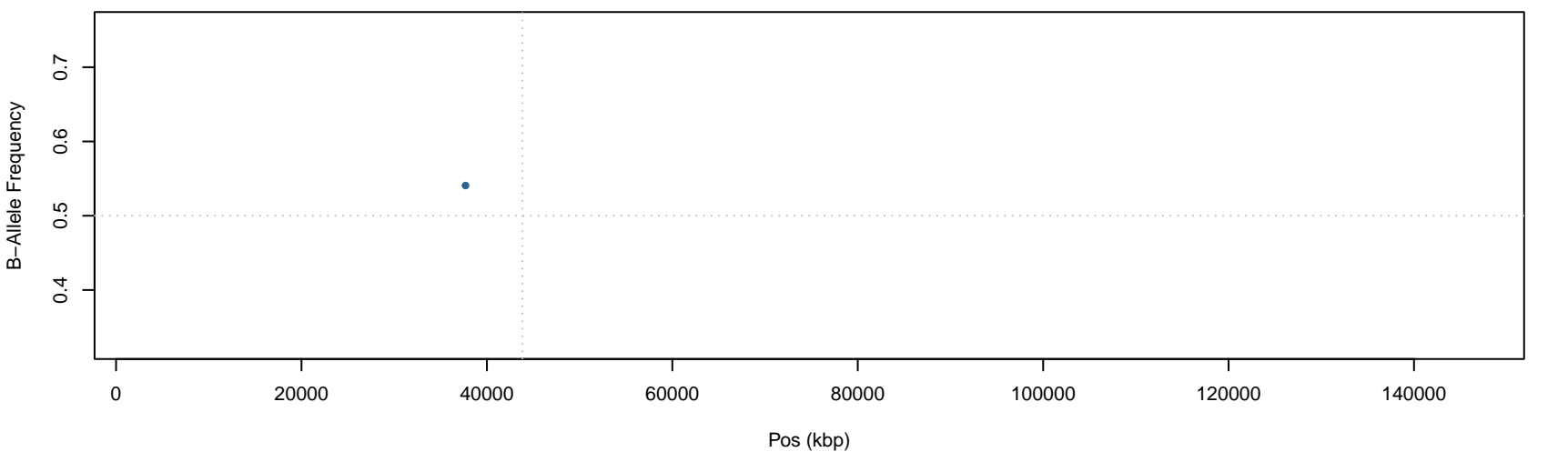
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 6



Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 7

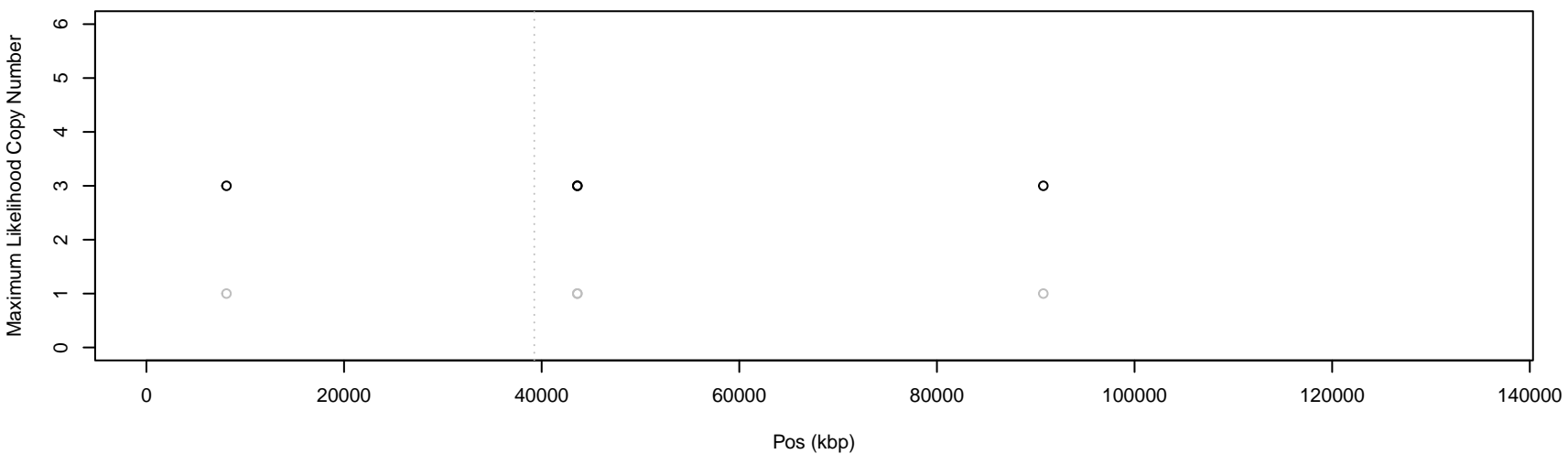
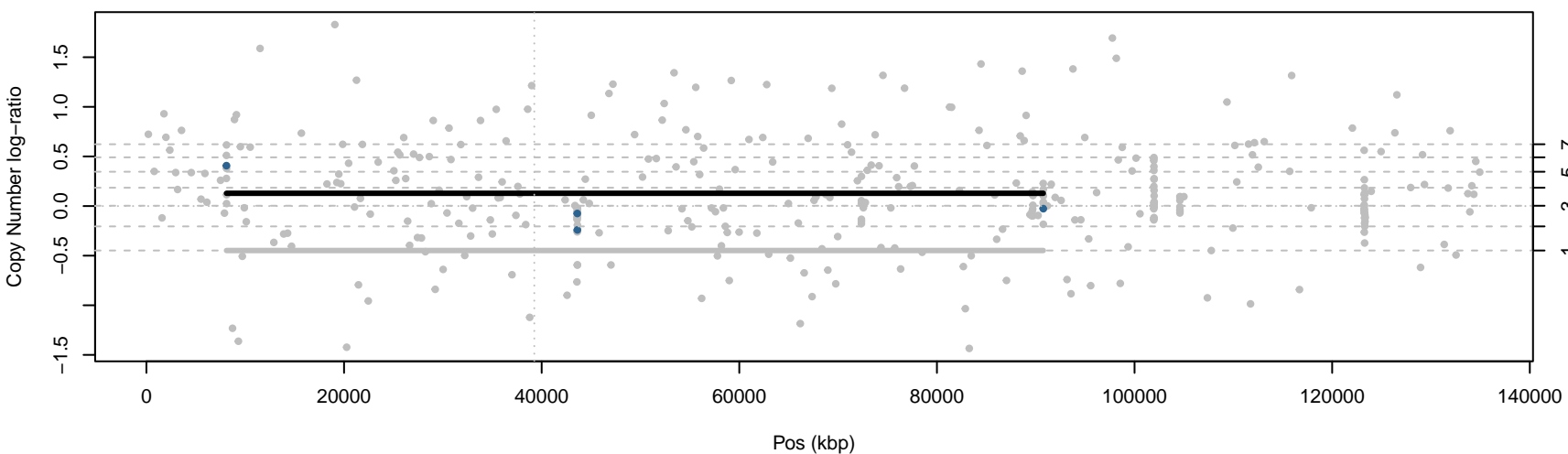
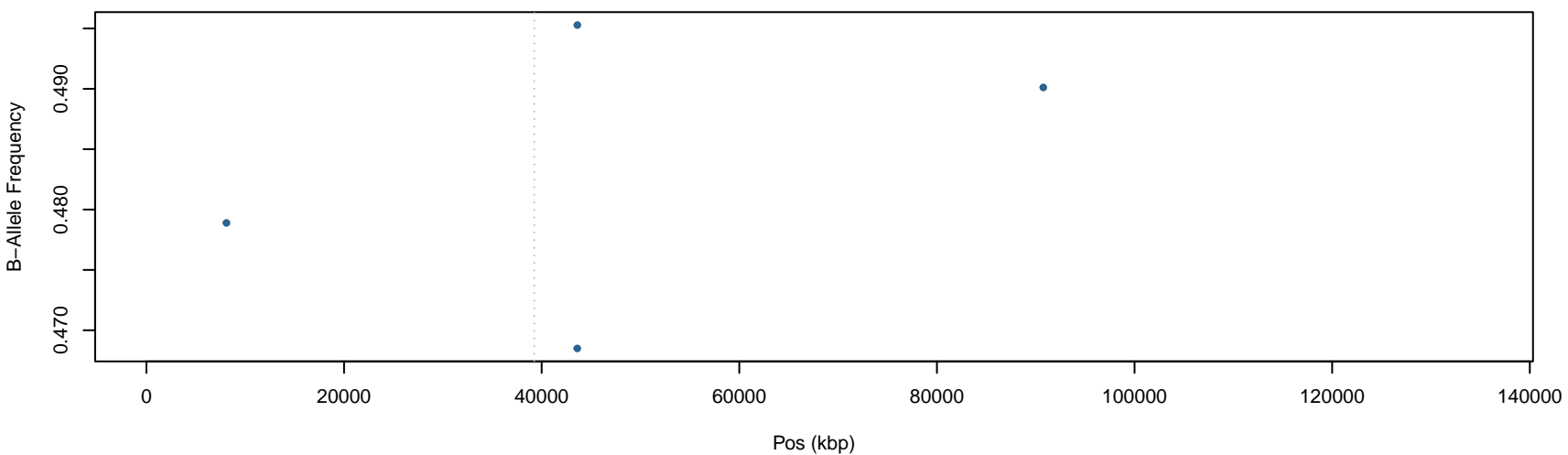


Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 8

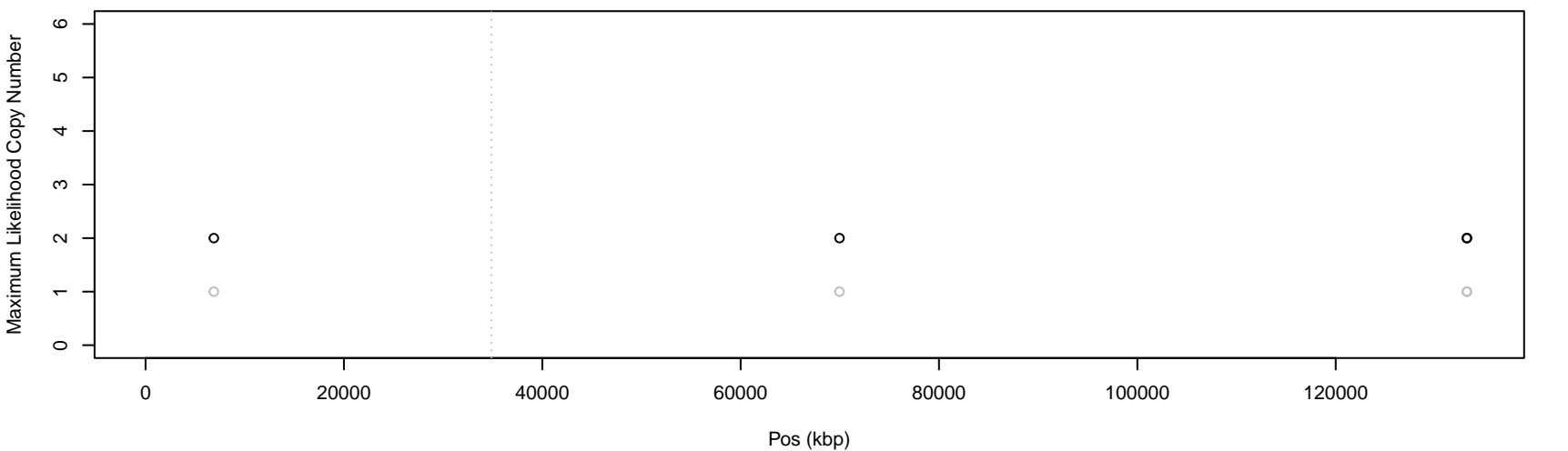
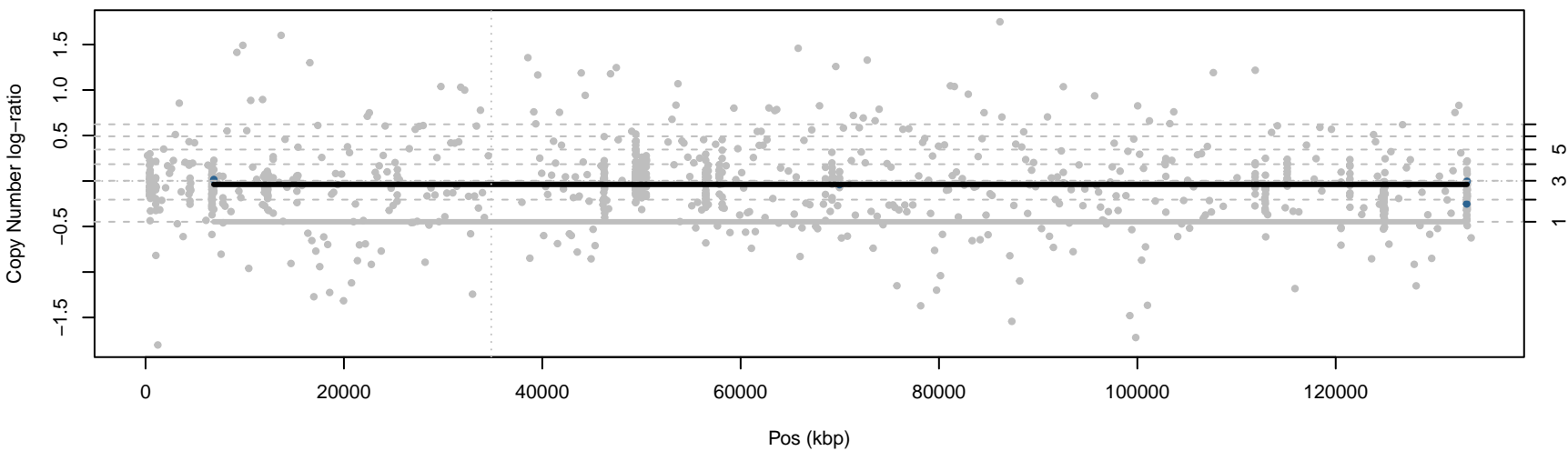
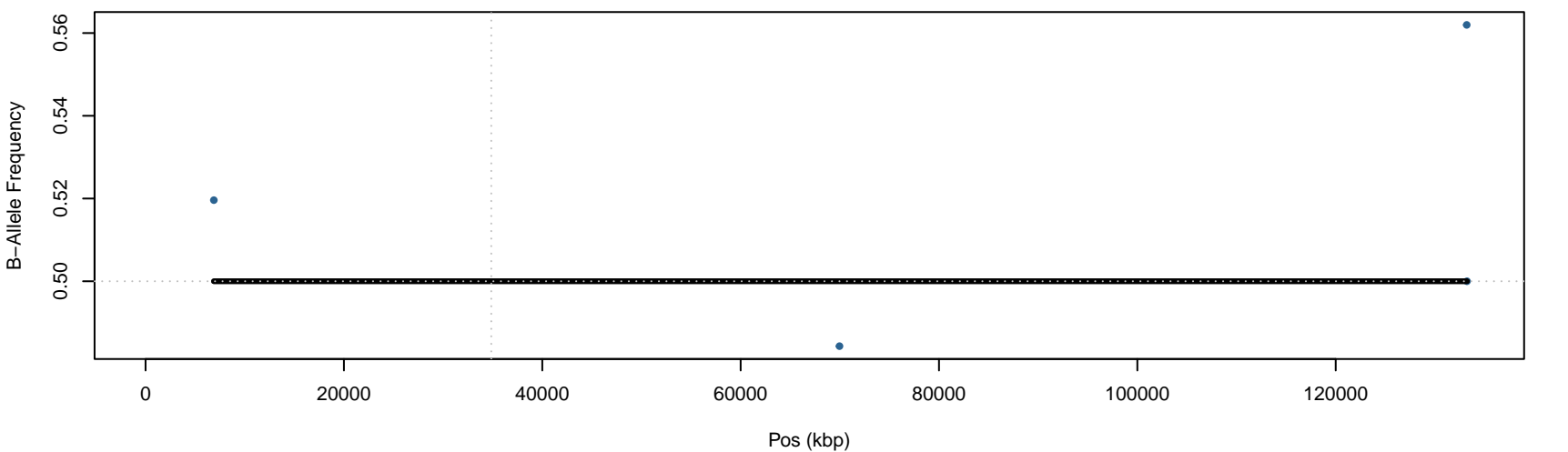




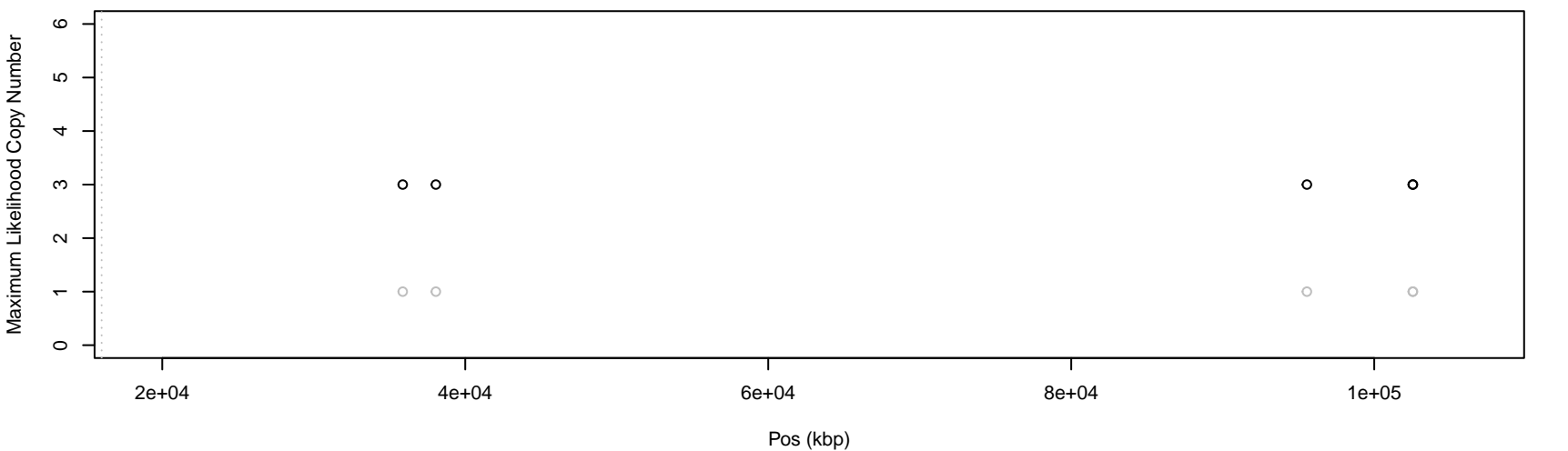
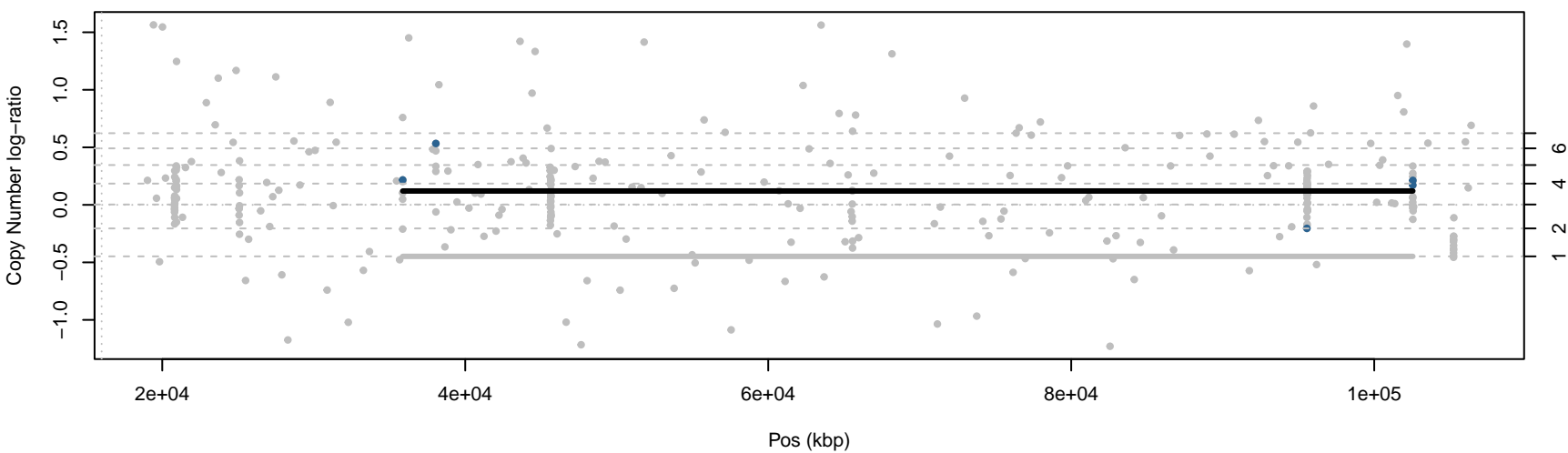
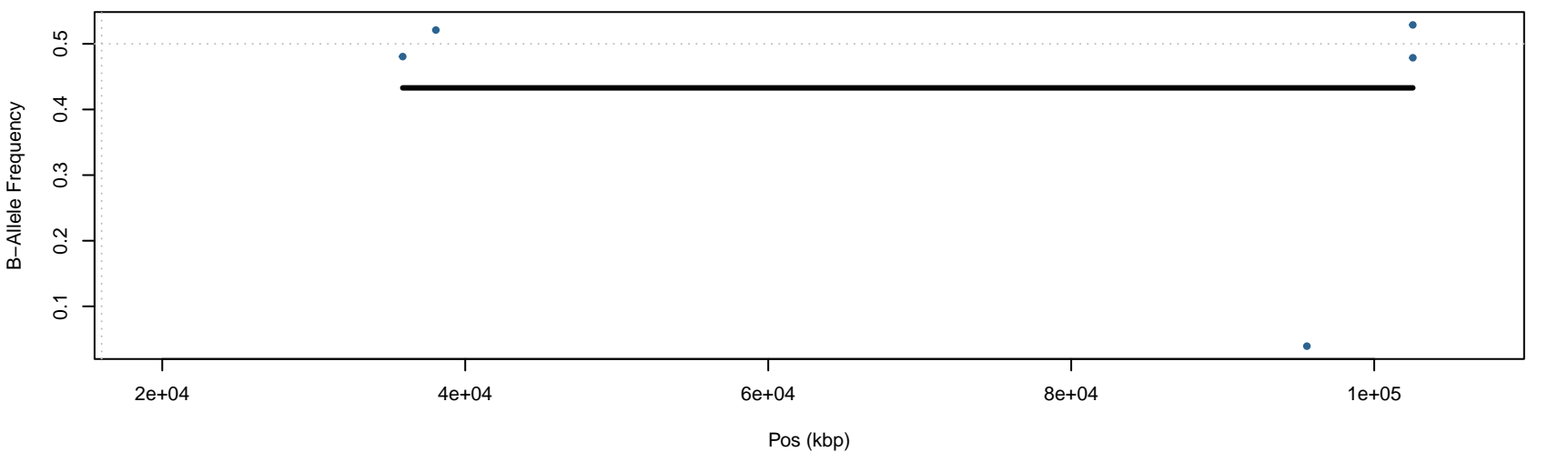
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 10



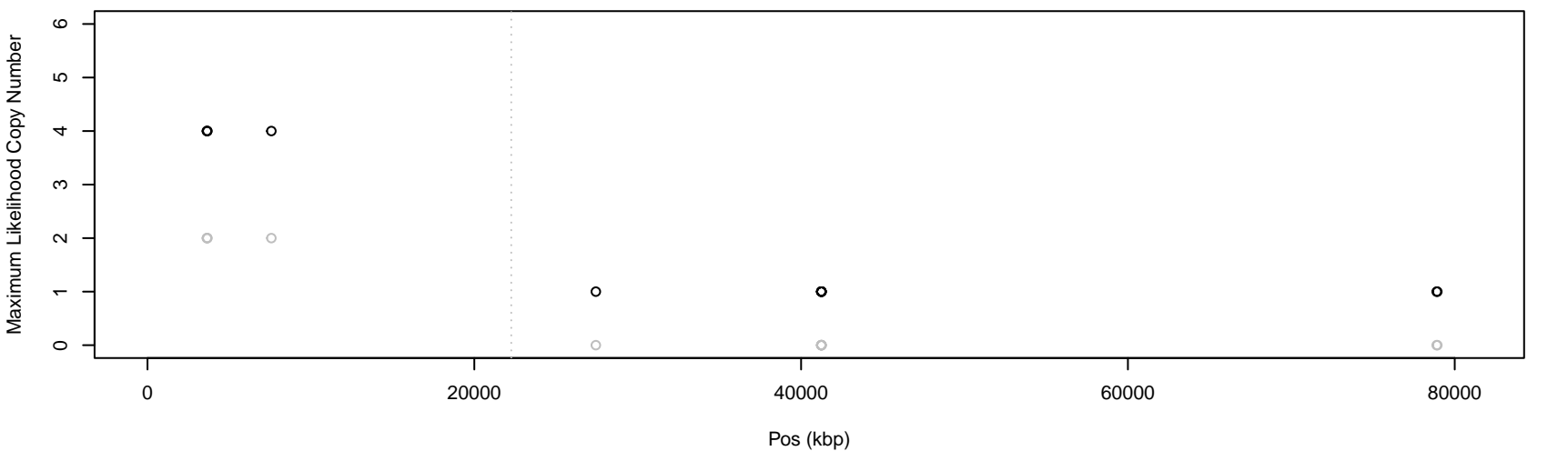
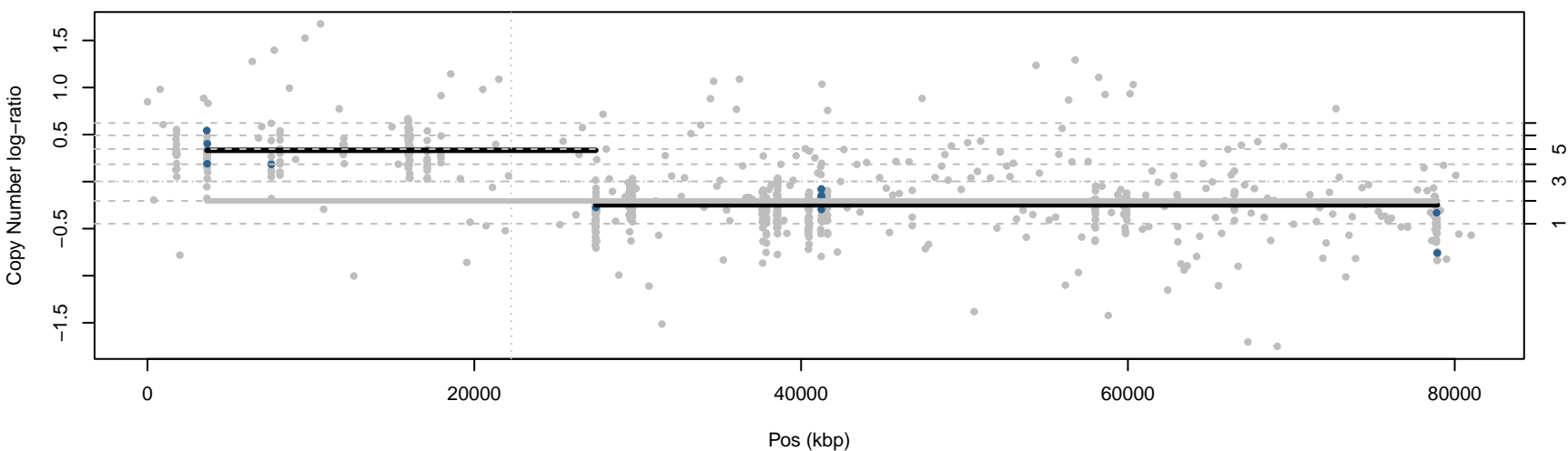
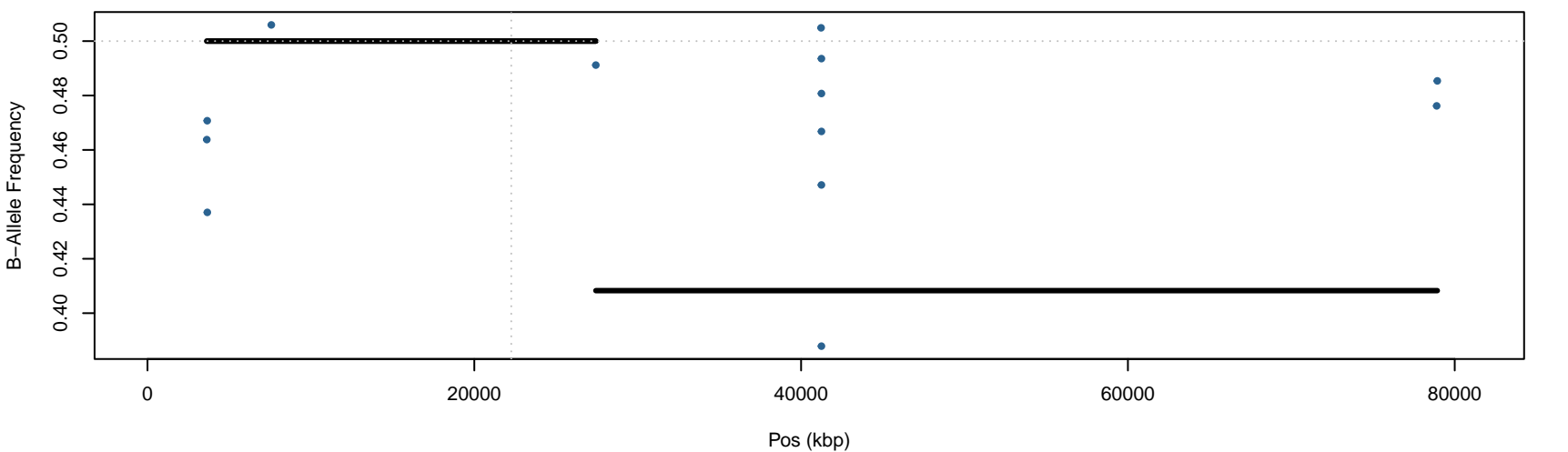
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 12



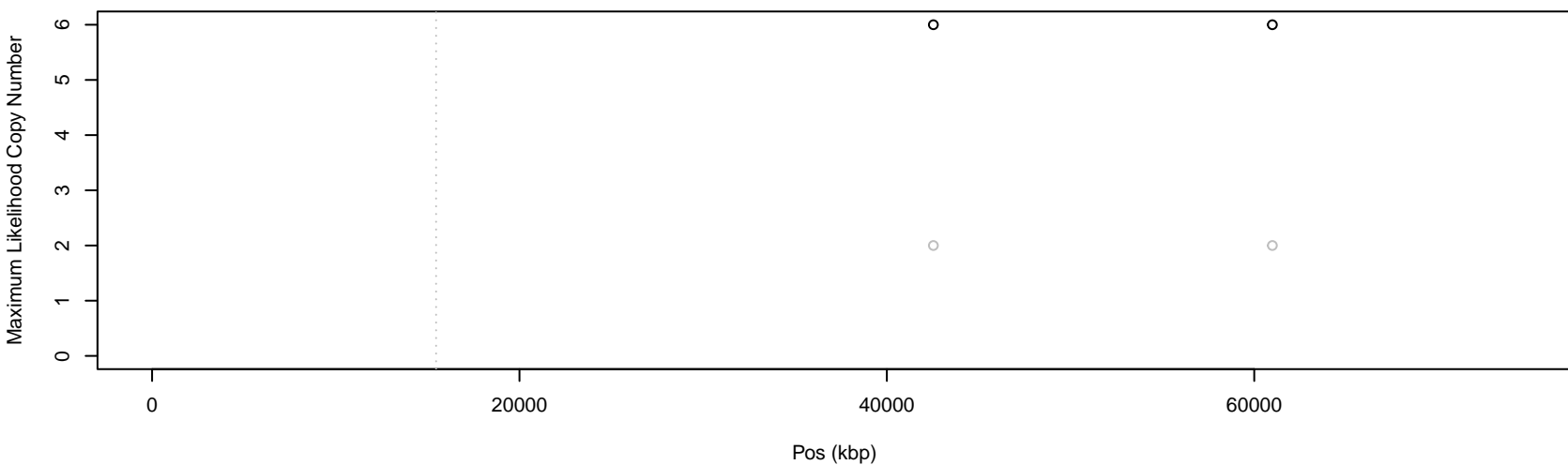
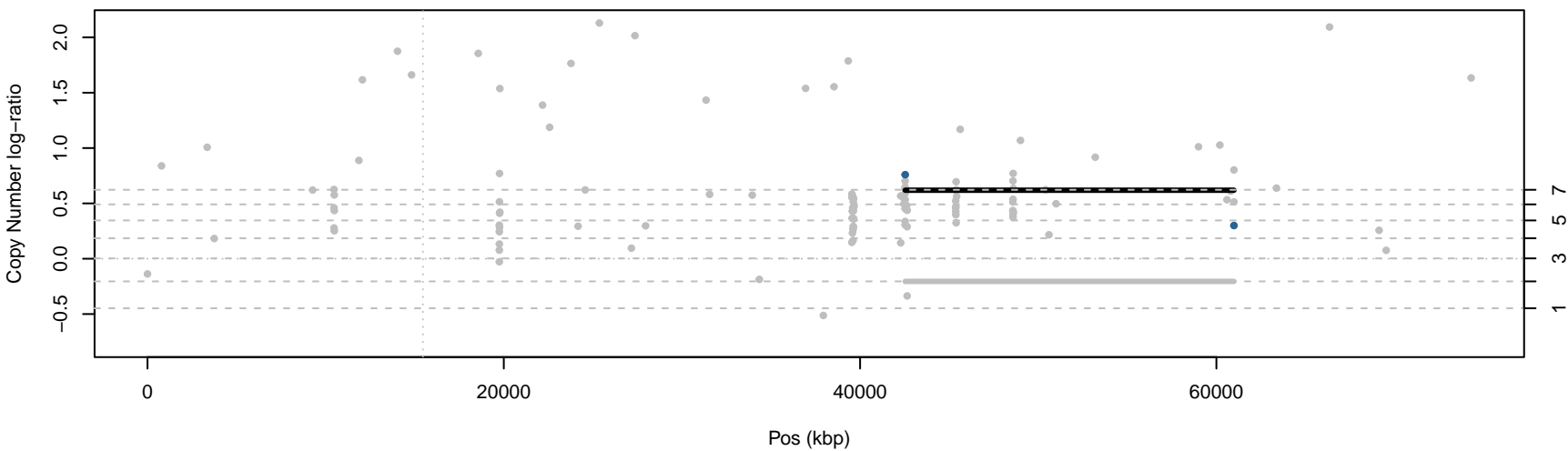
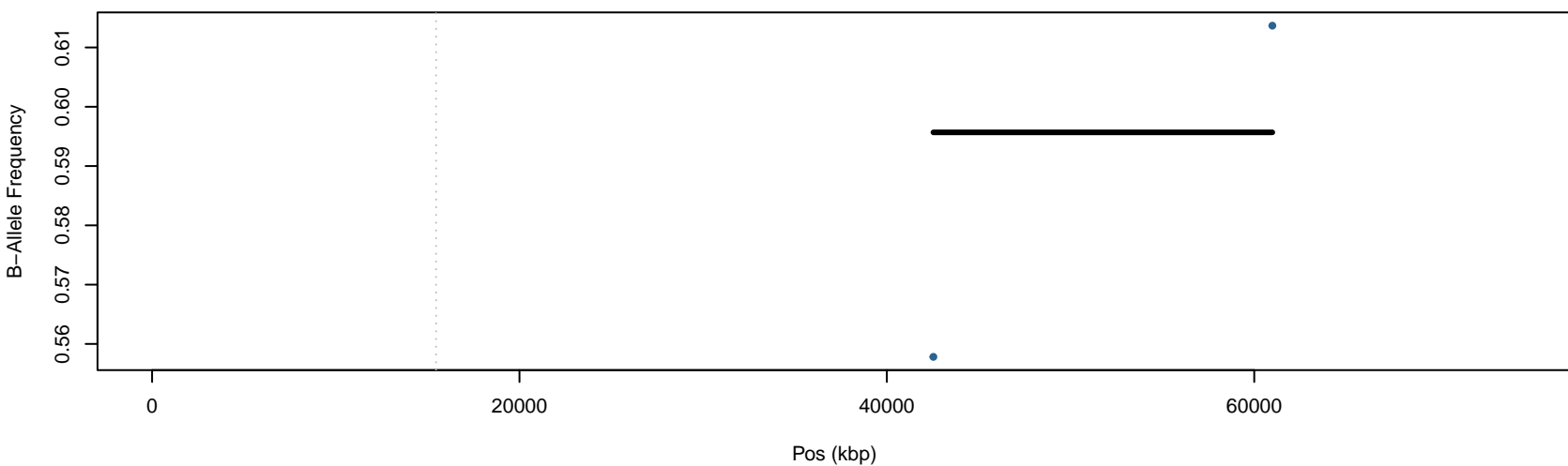
Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 14



Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 17



Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 18



Purity: 0.31 Tumor ploidy: 2.985 SNV log-likelihood: -71.89 GoF: 91.7% Mean coverage: 586;641 Chromosome: 19

