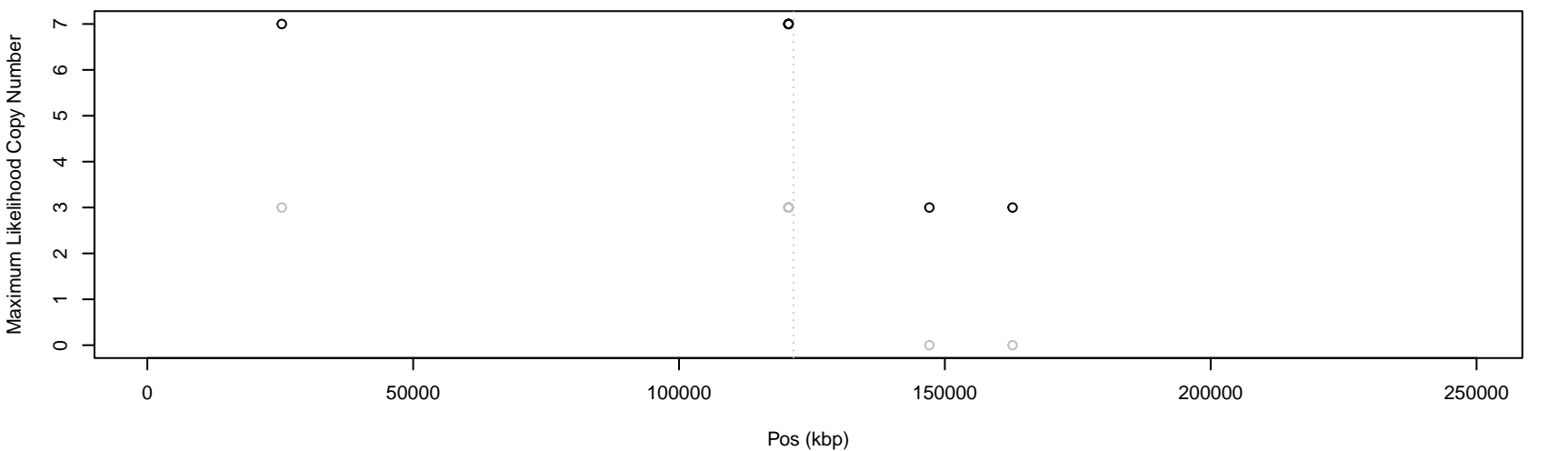
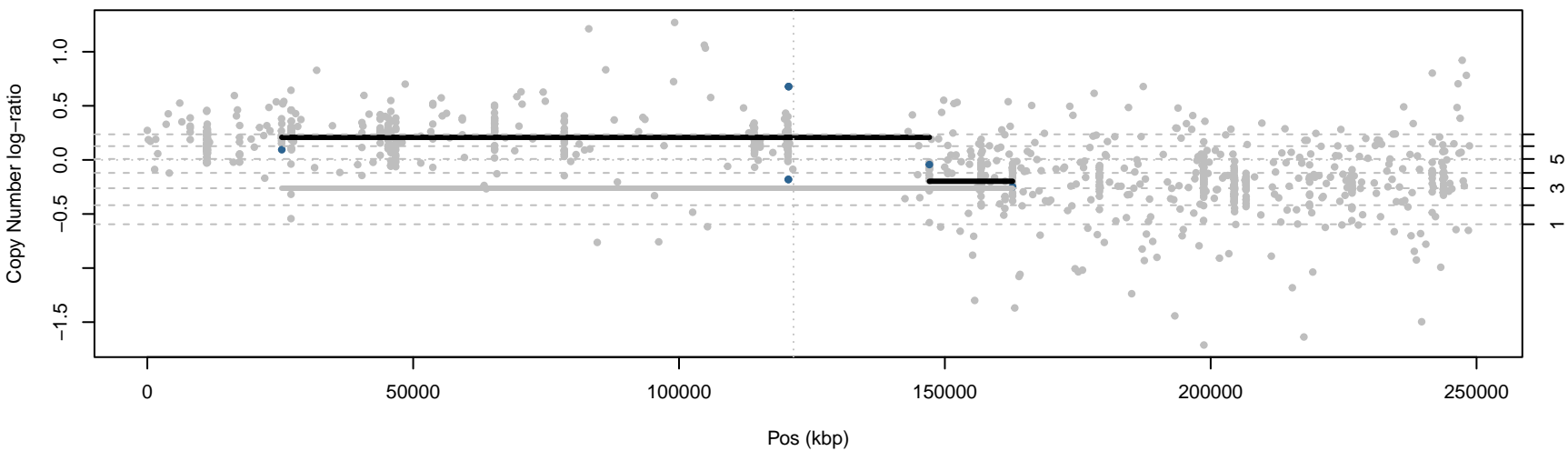
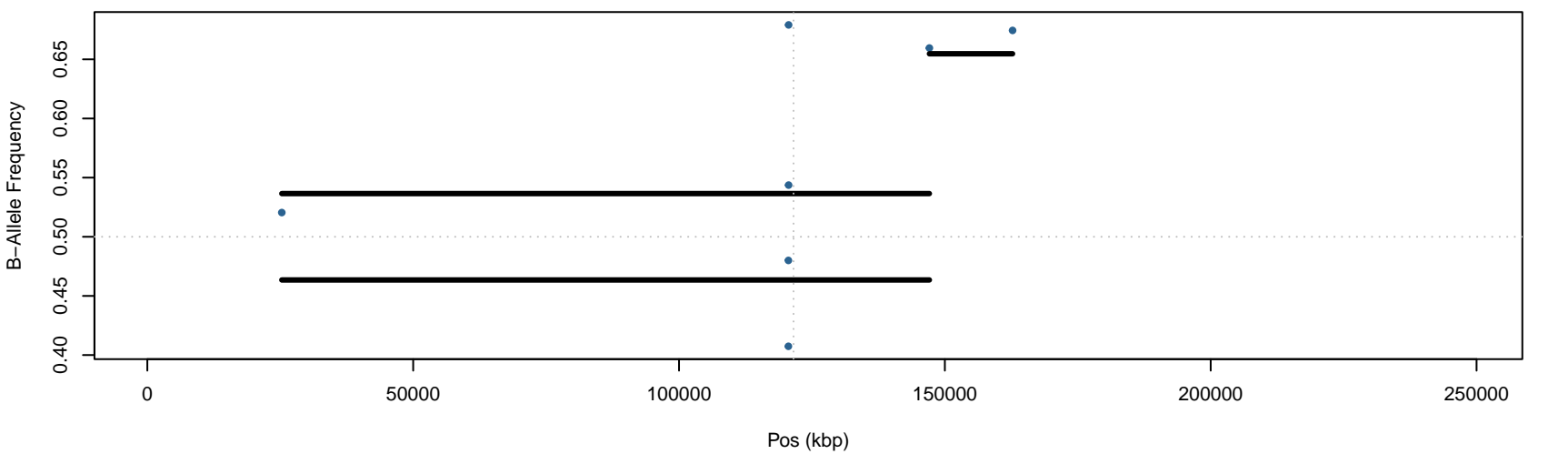
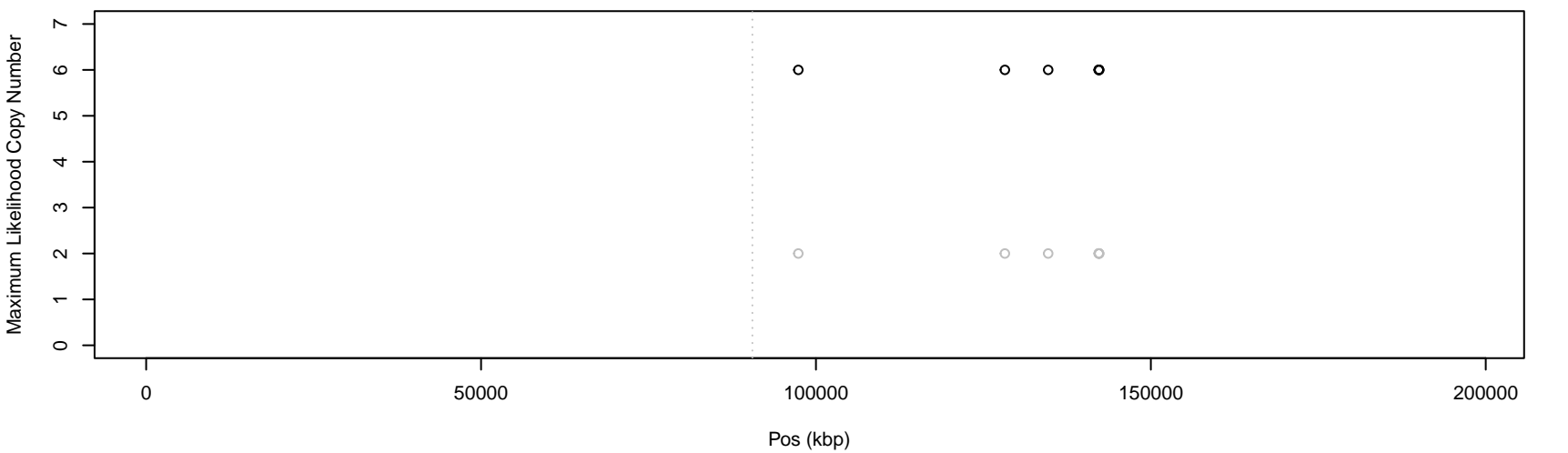
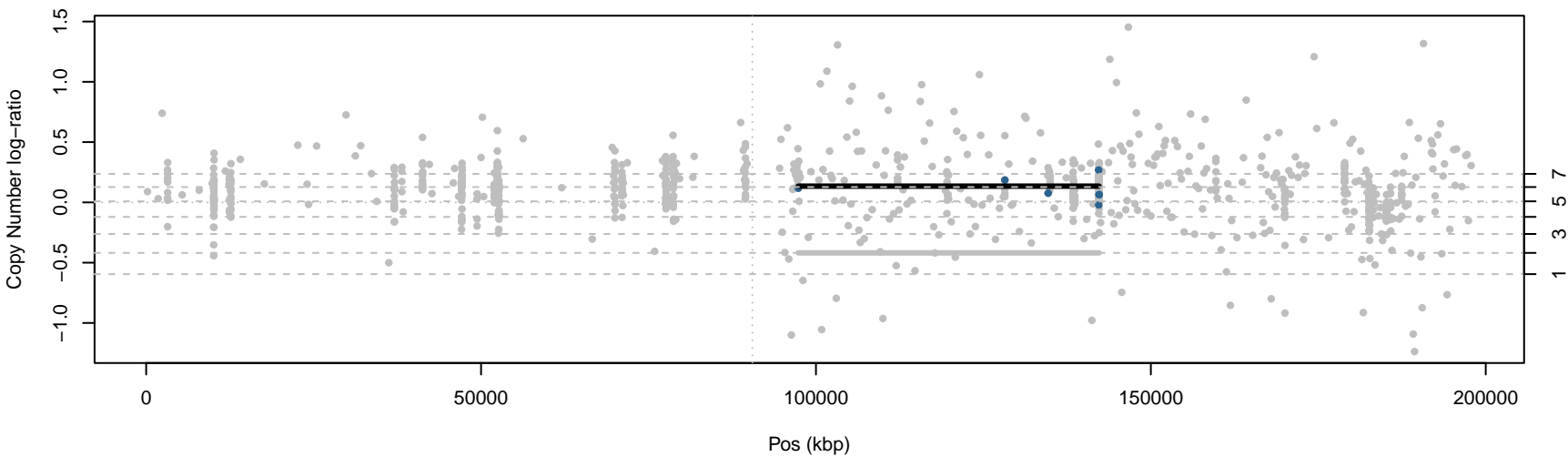
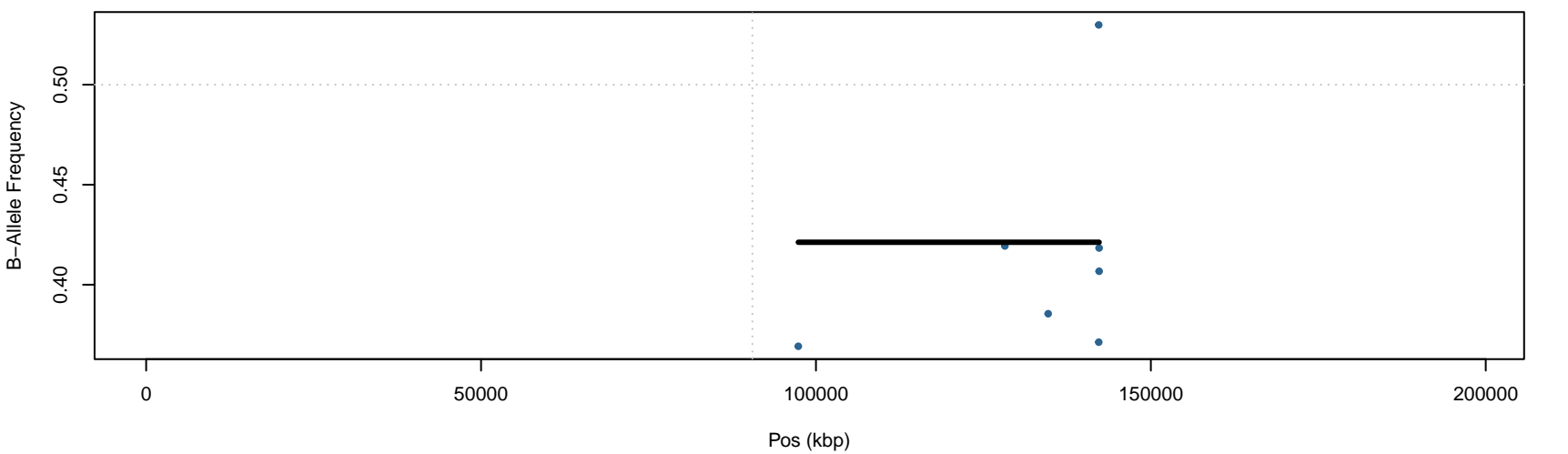


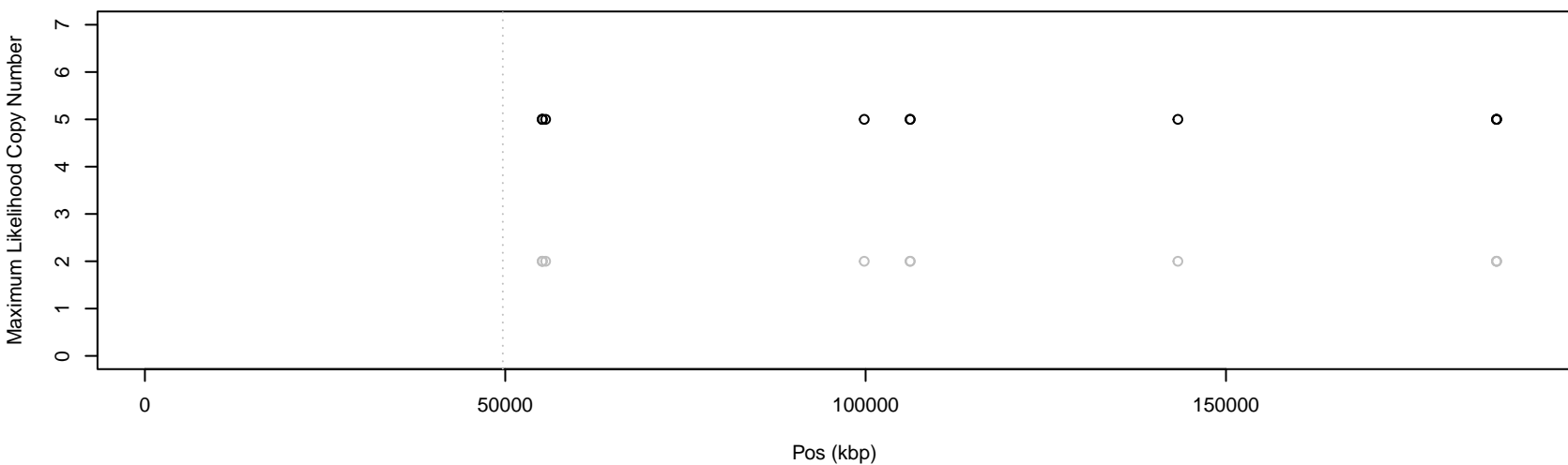
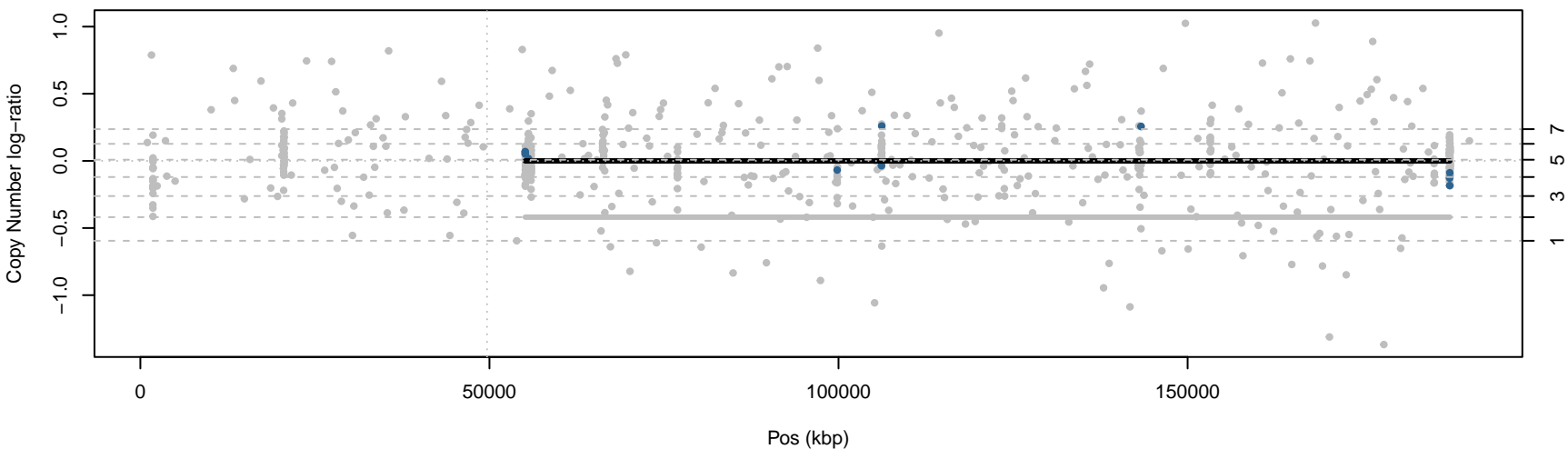
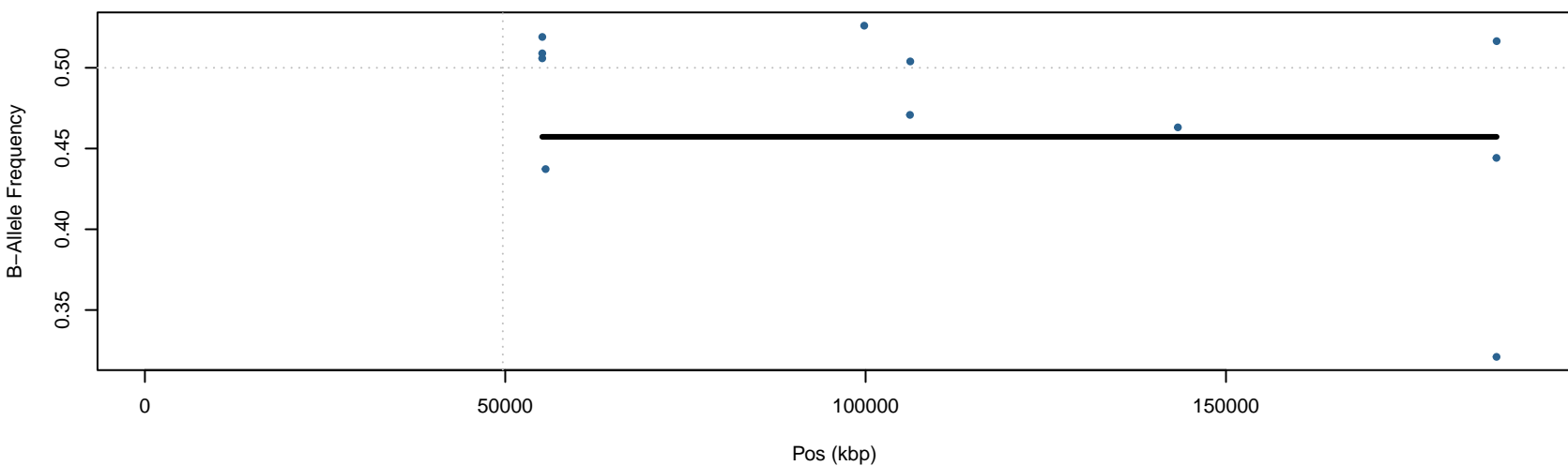
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 1



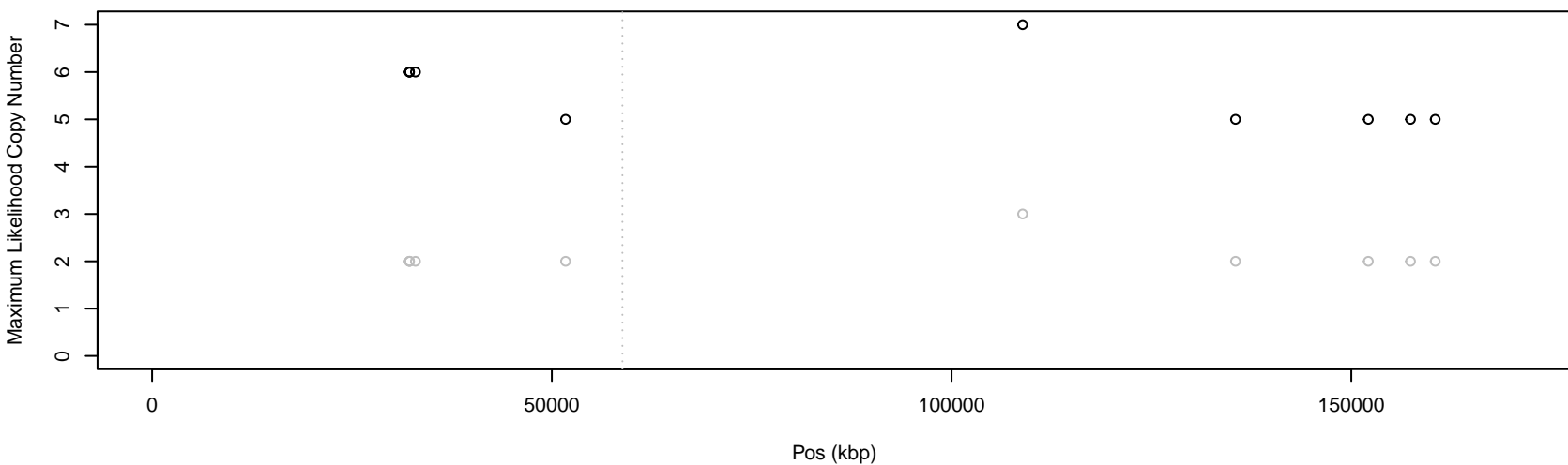
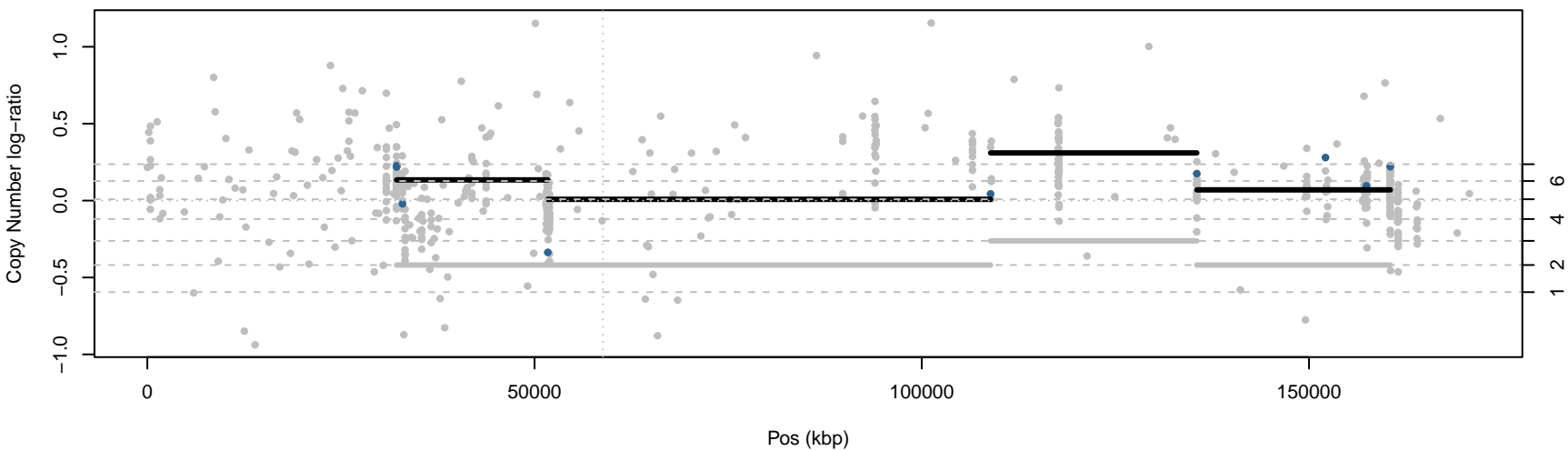
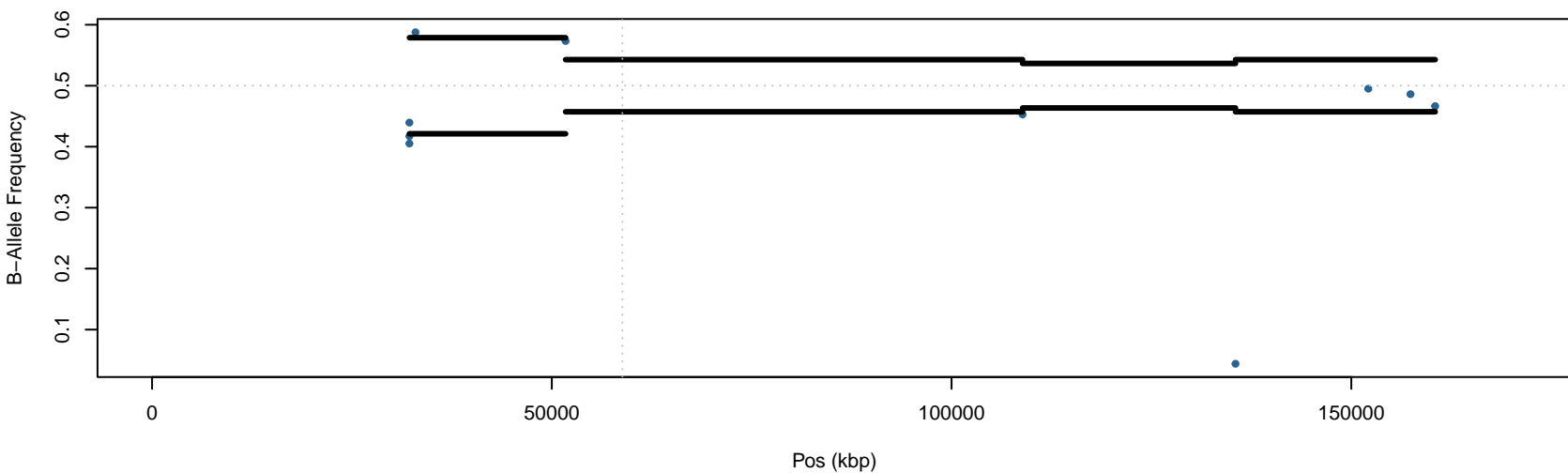
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 3



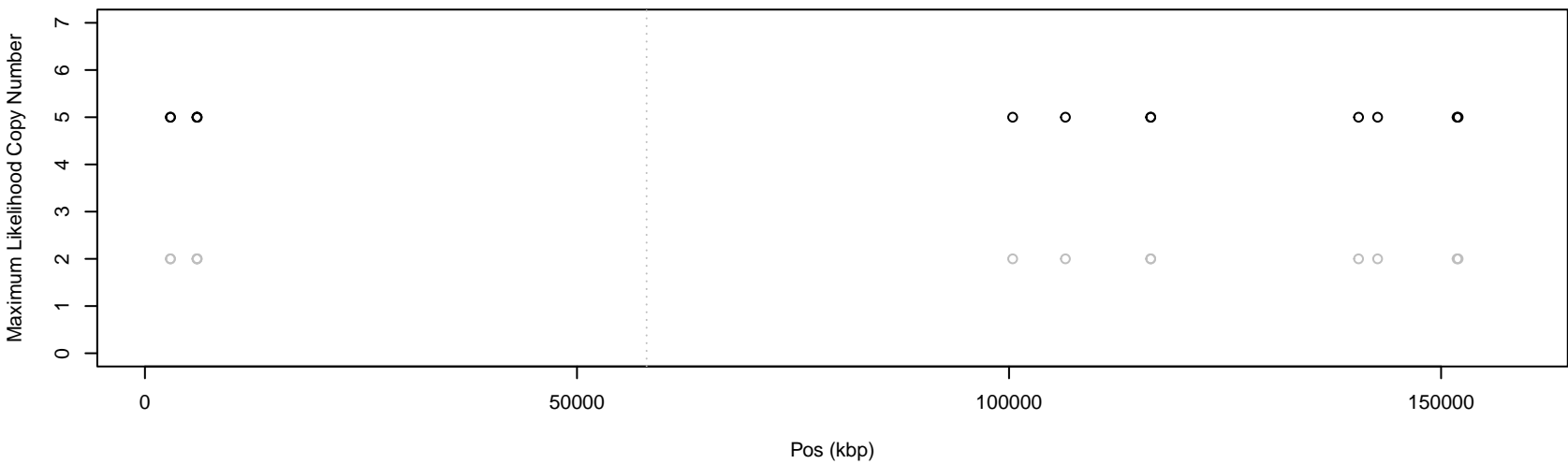
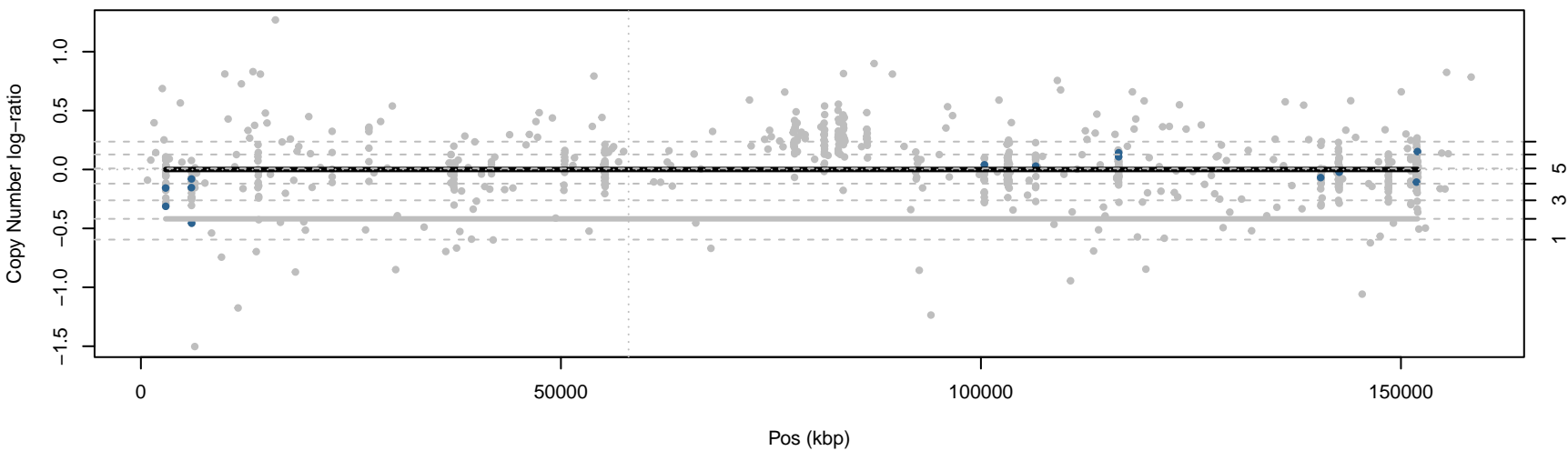
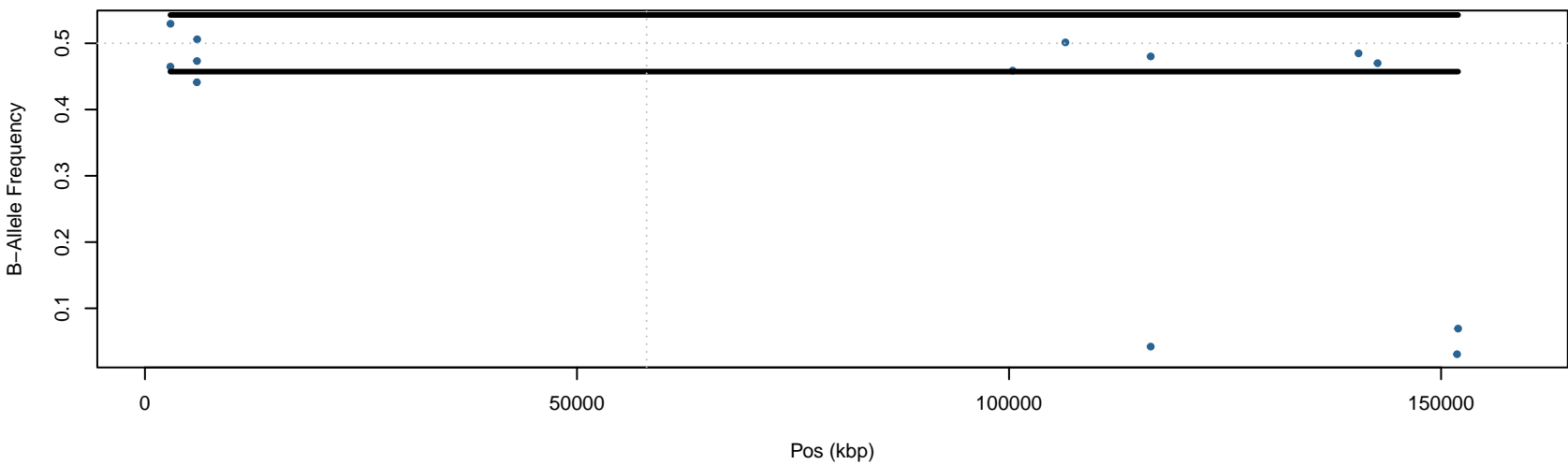
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 4



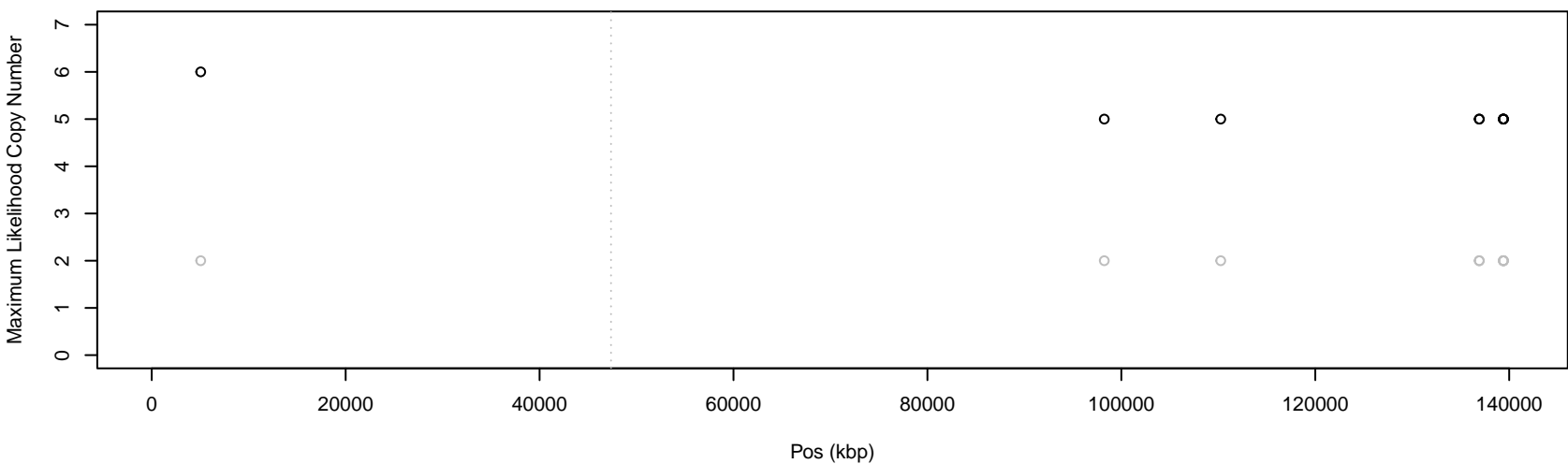
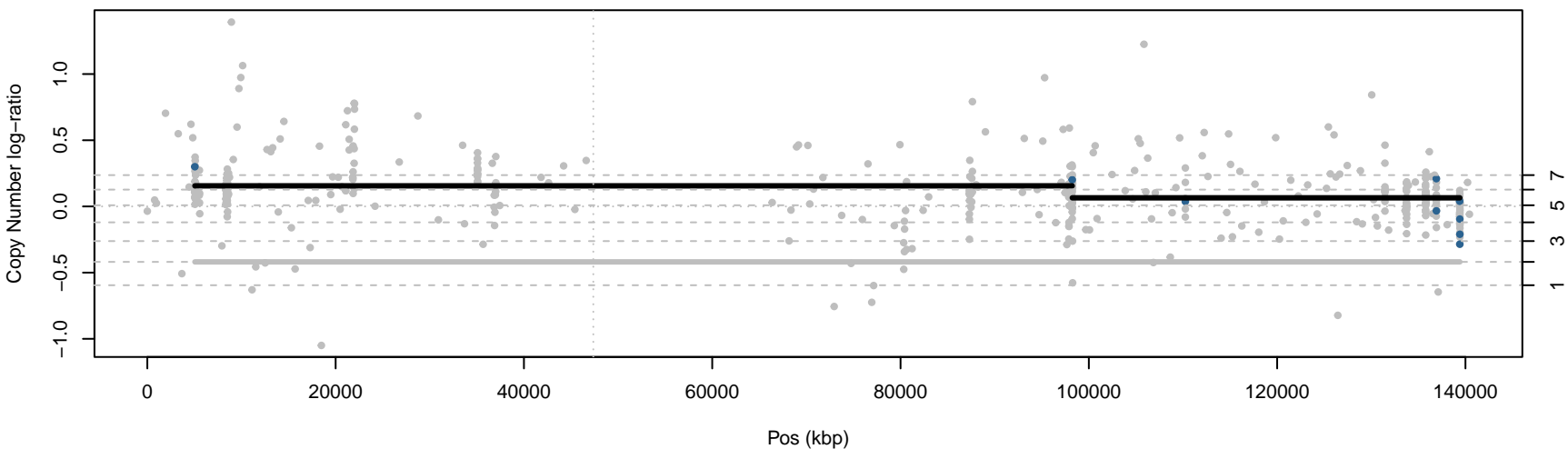
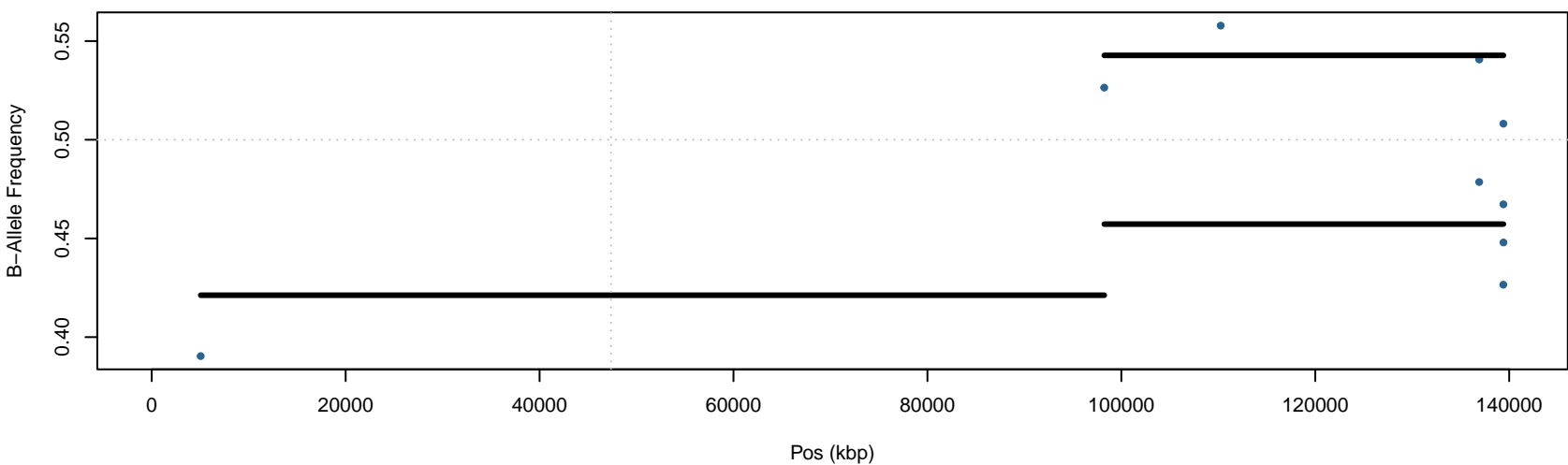
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 6



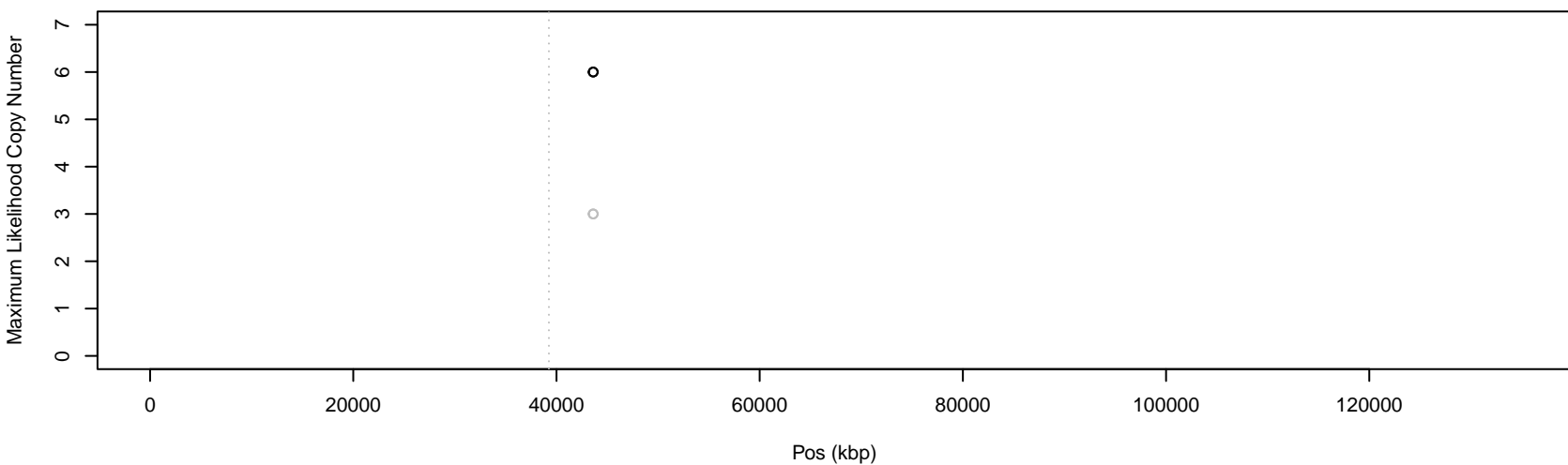
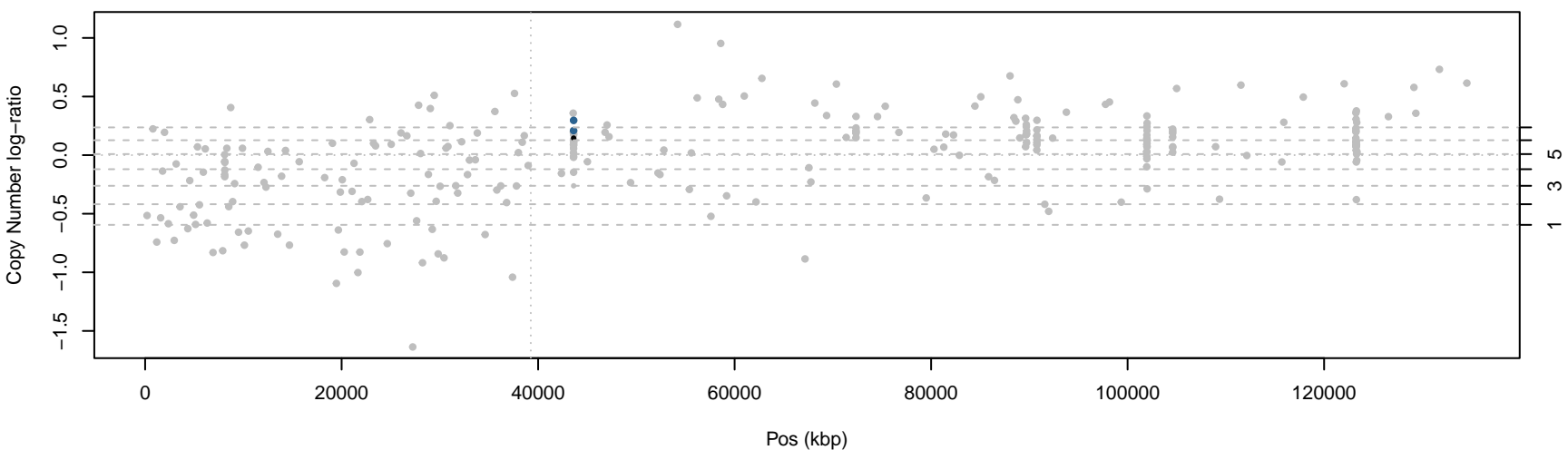
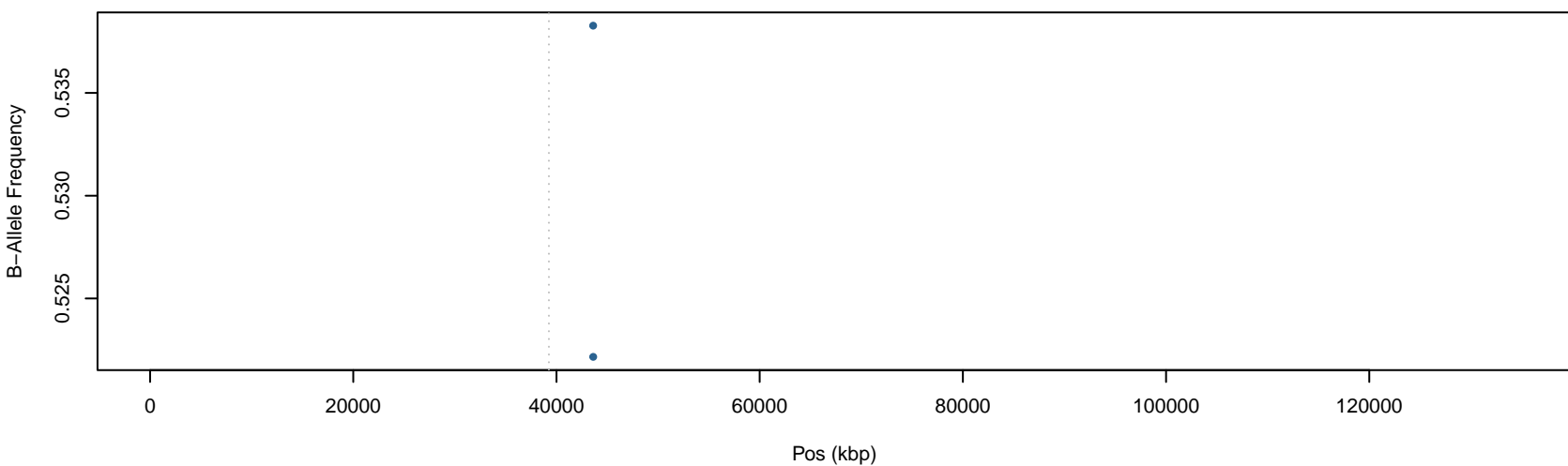
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 7



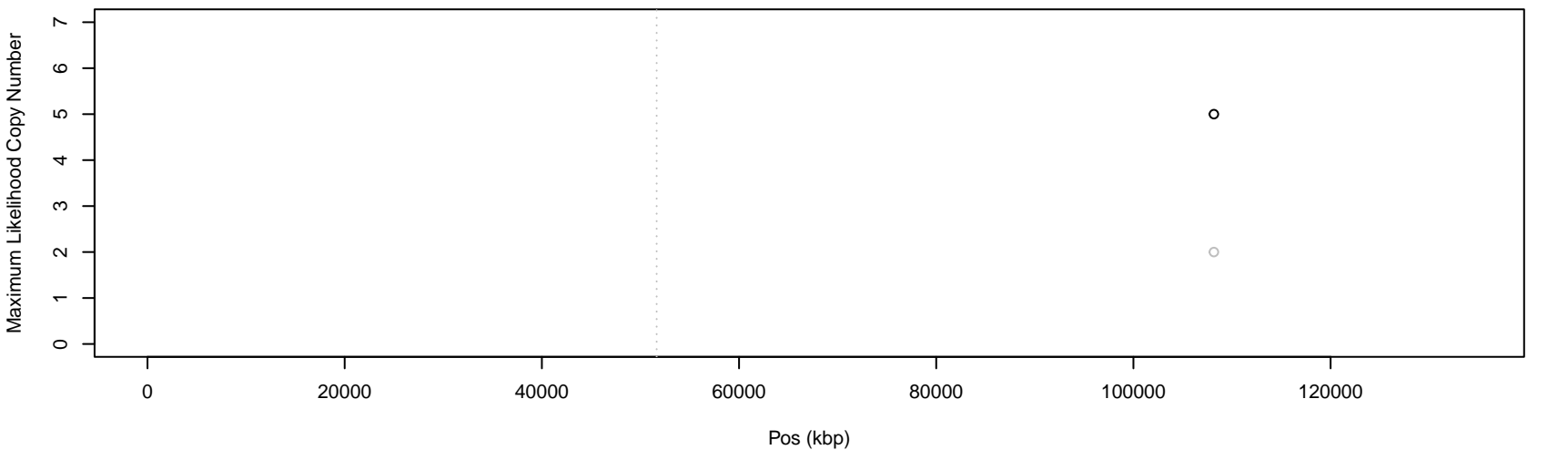
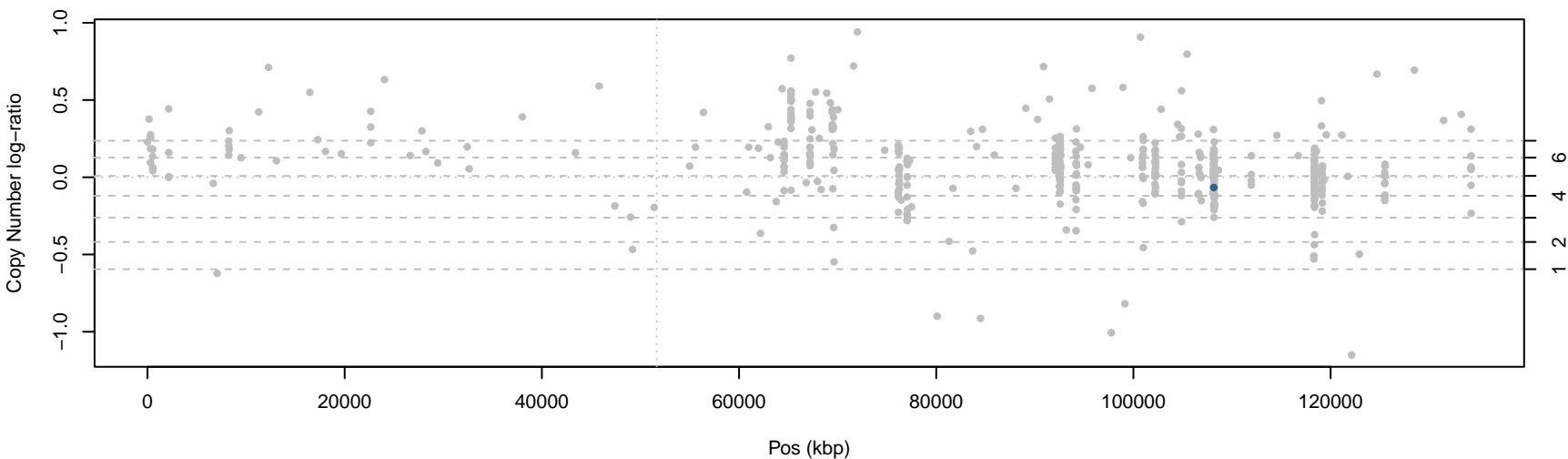
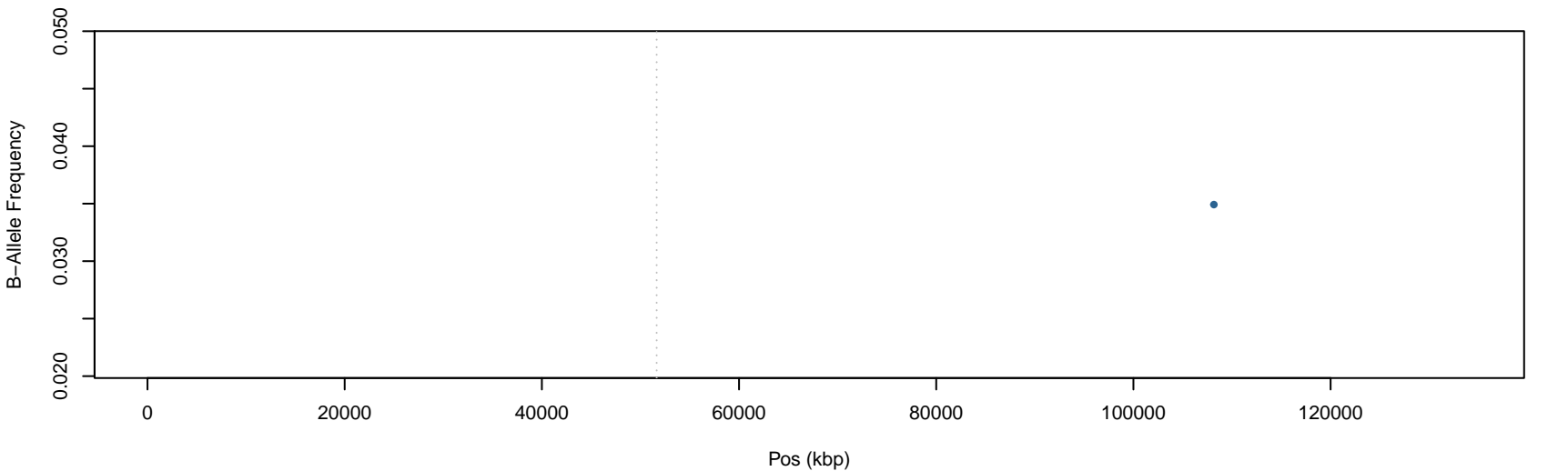
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 9



Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 10

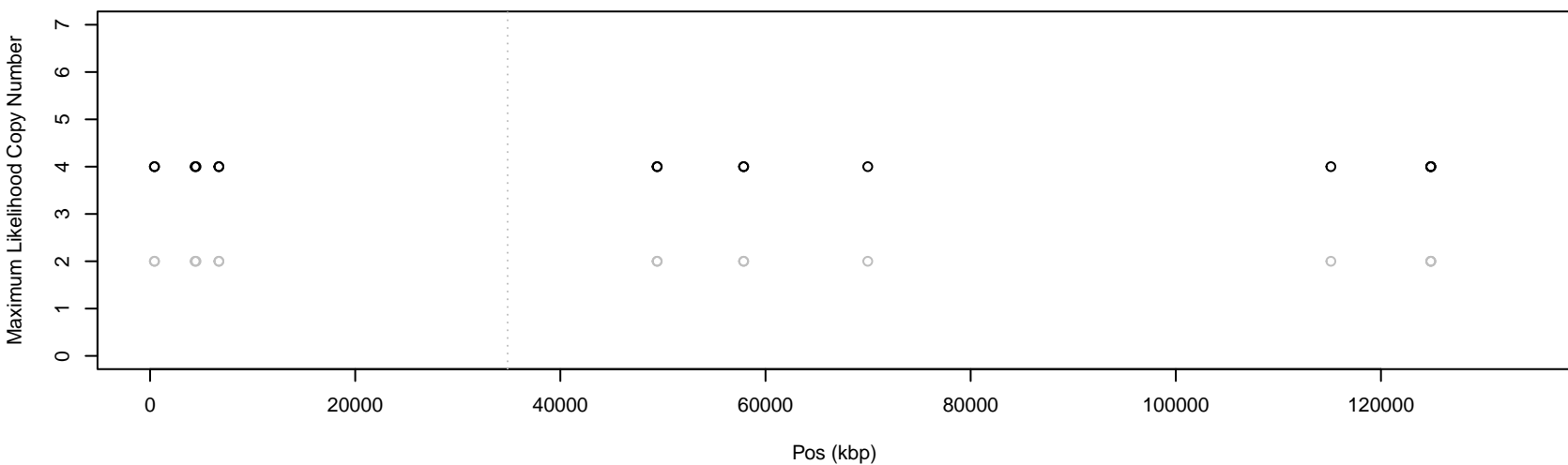
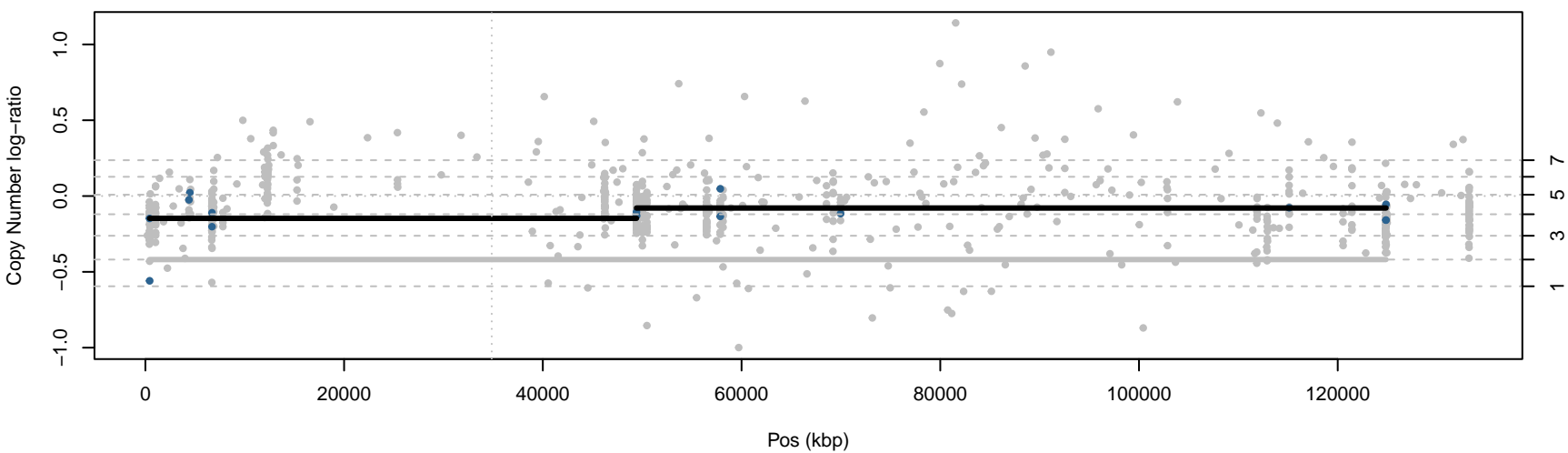
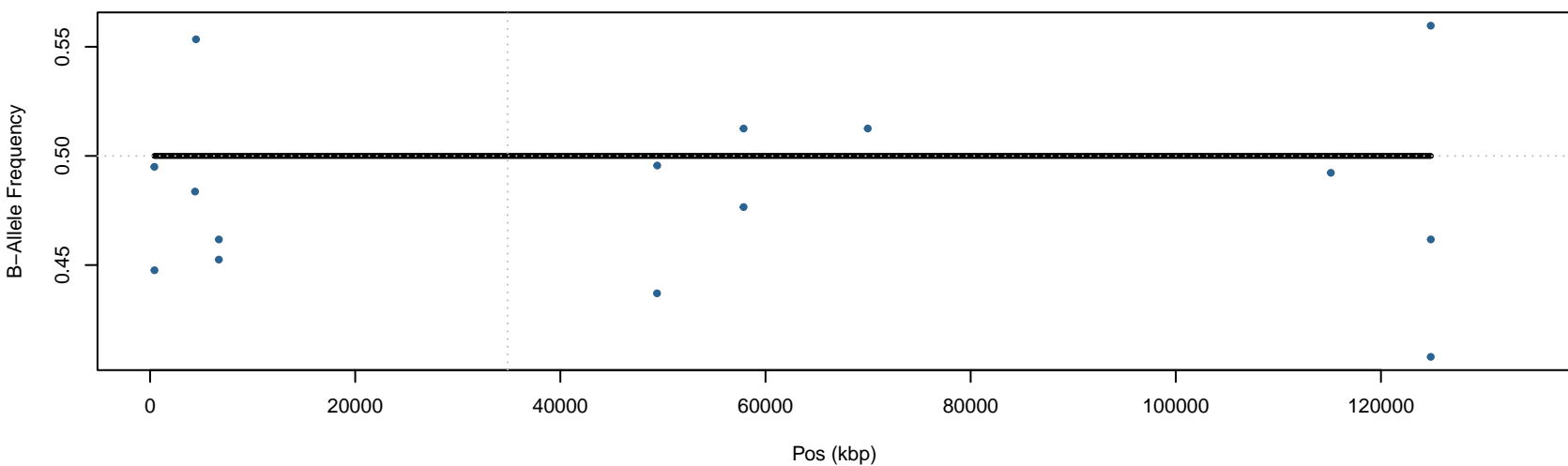


Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 11

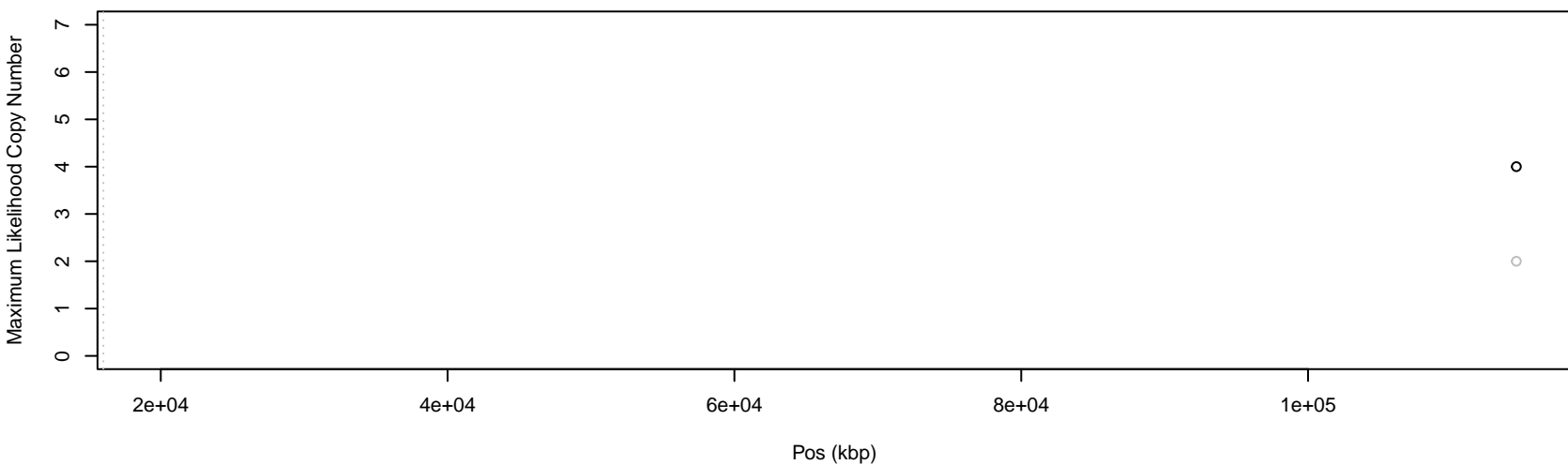
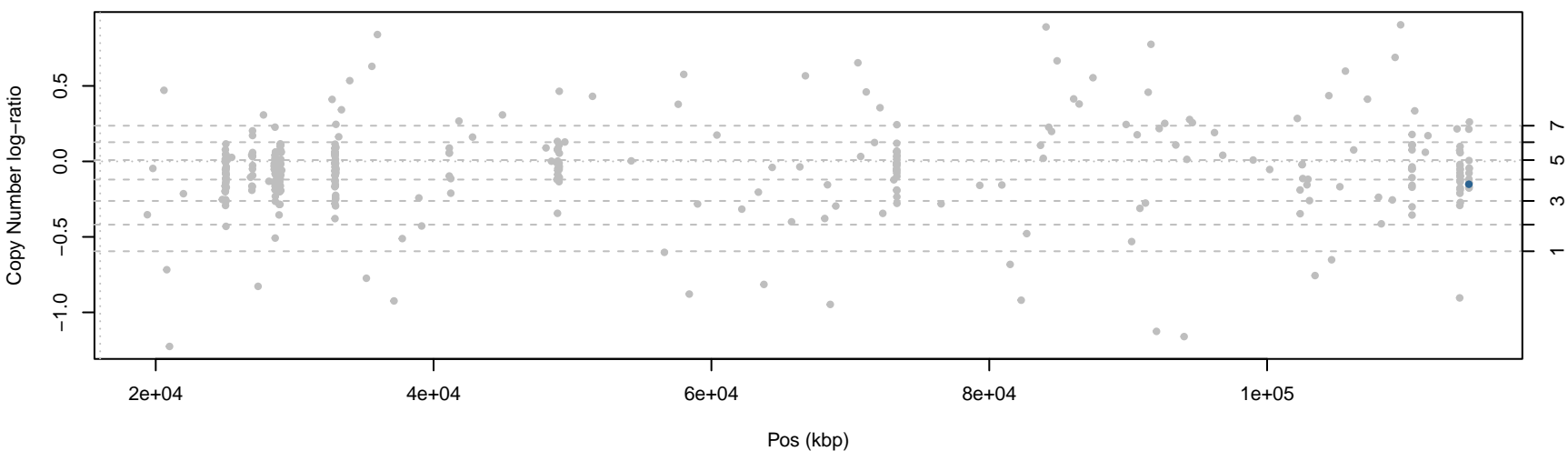
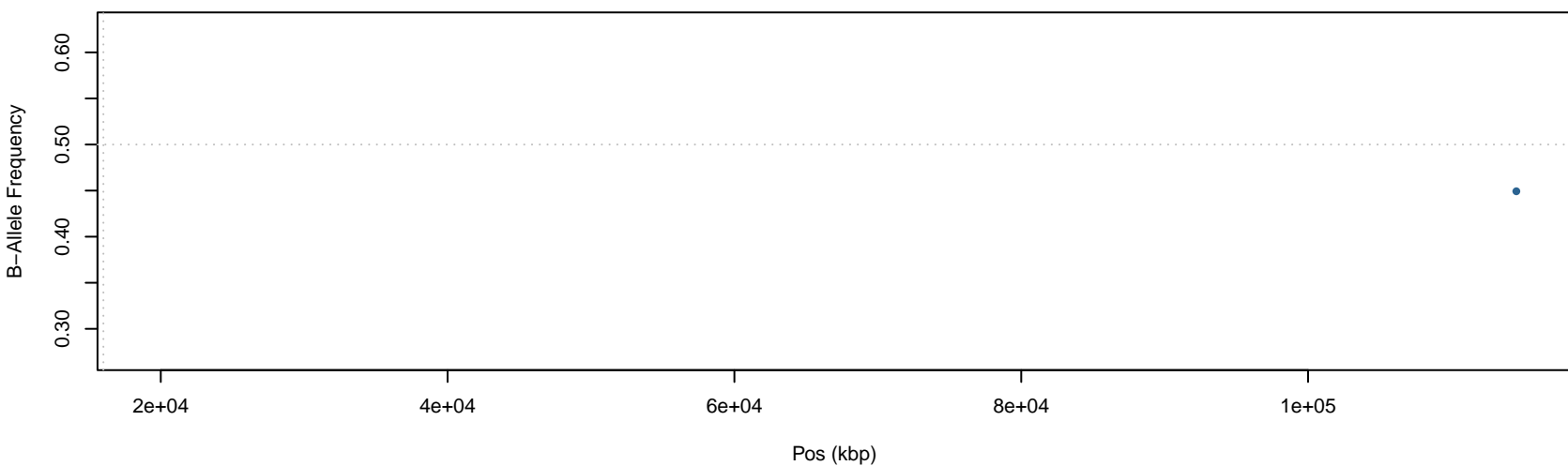




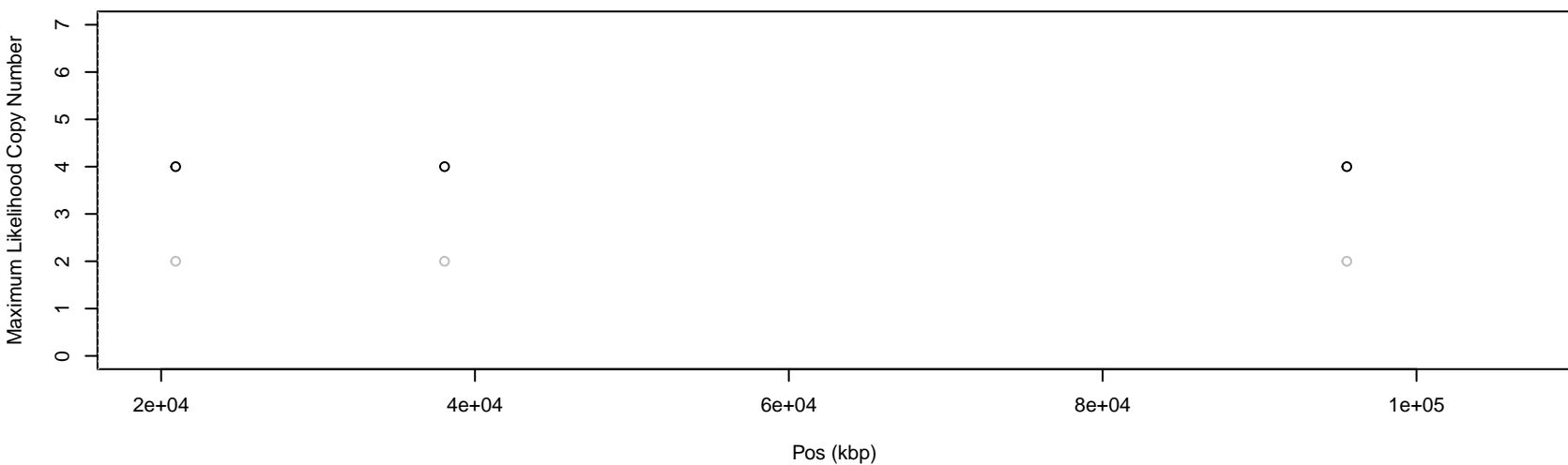
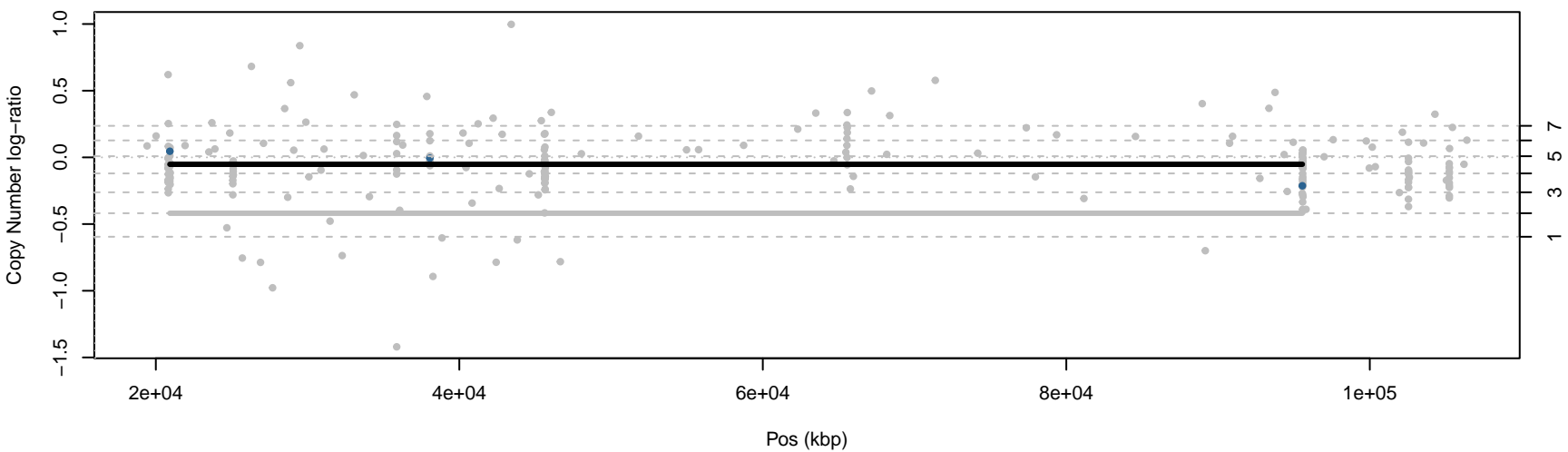
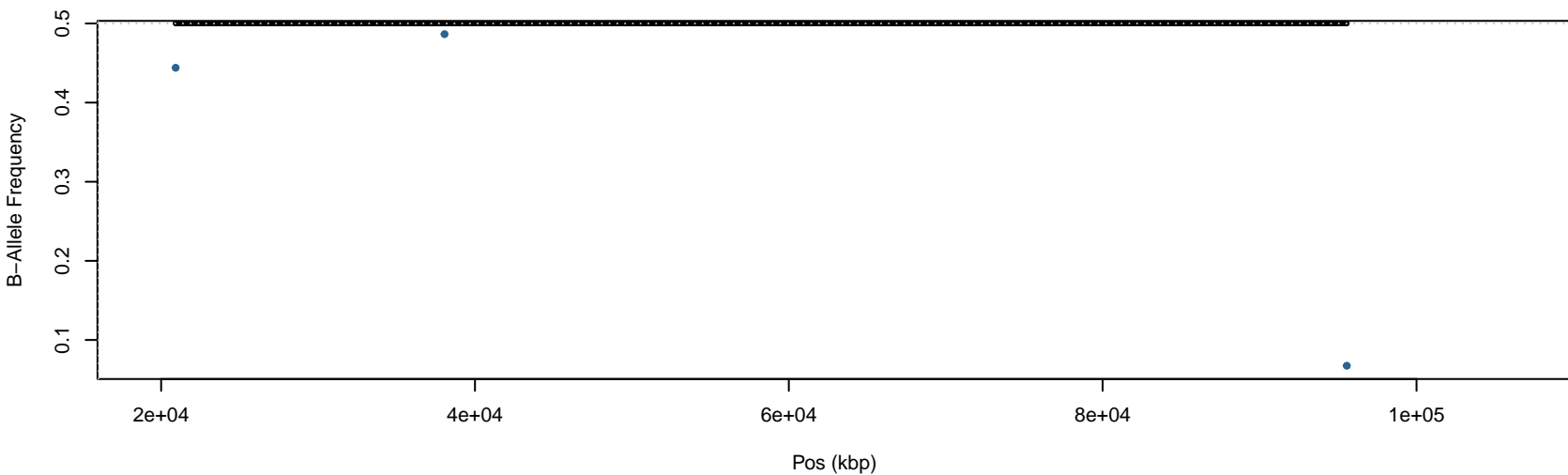
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 12



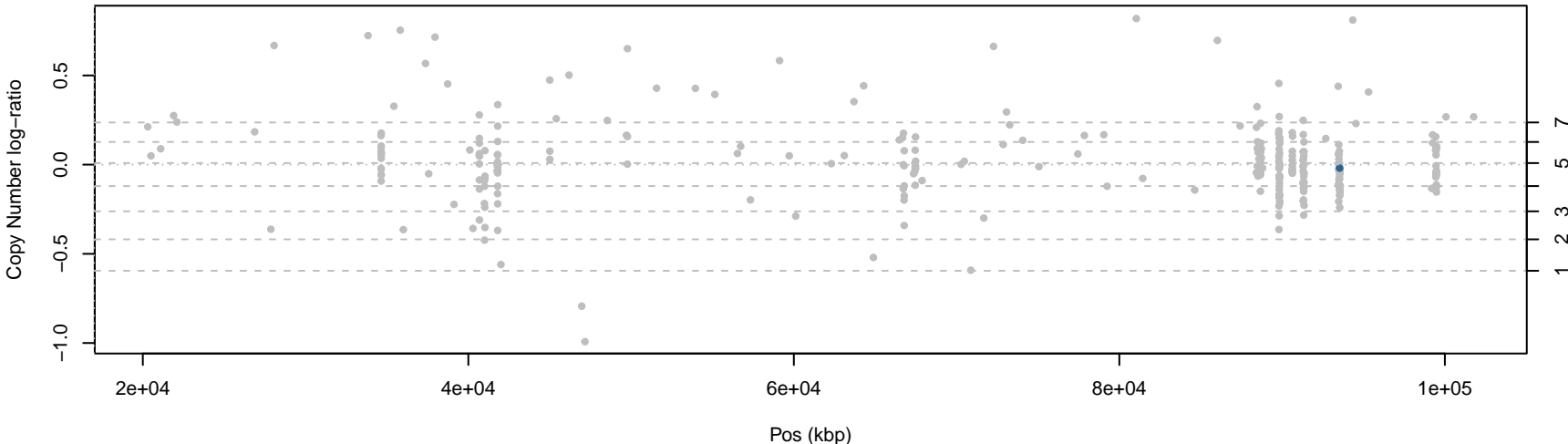
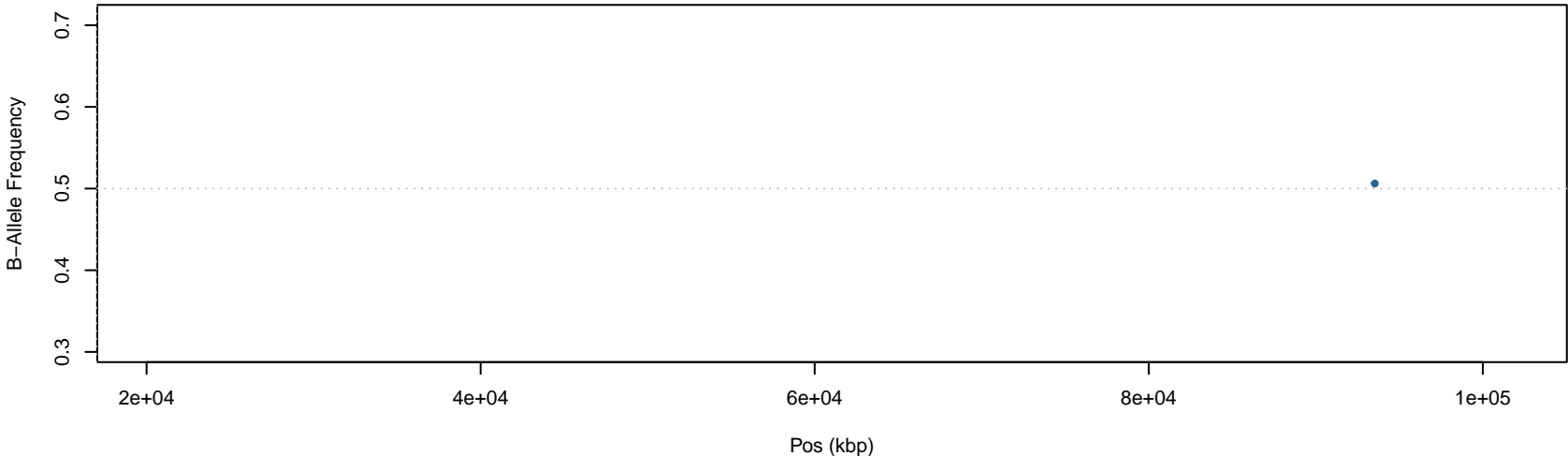
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 13



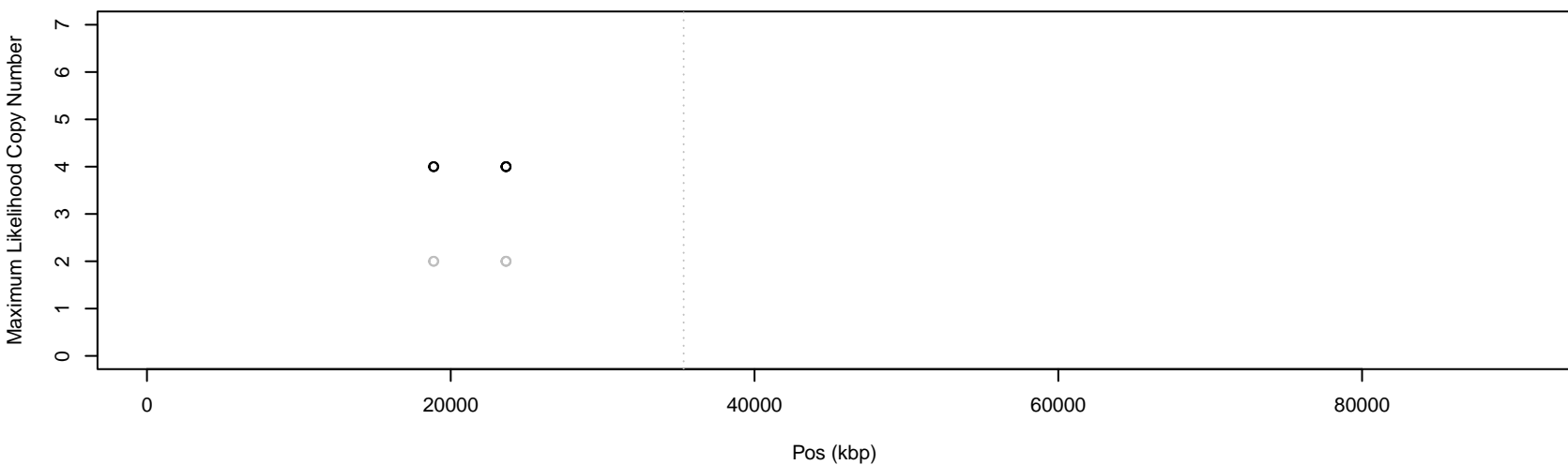
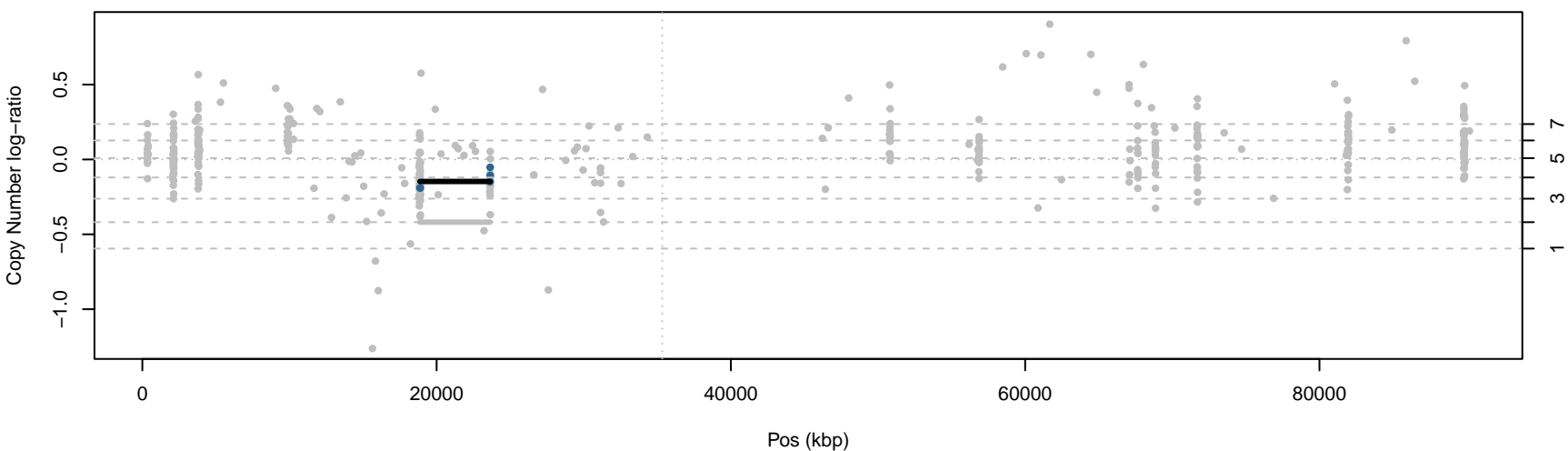
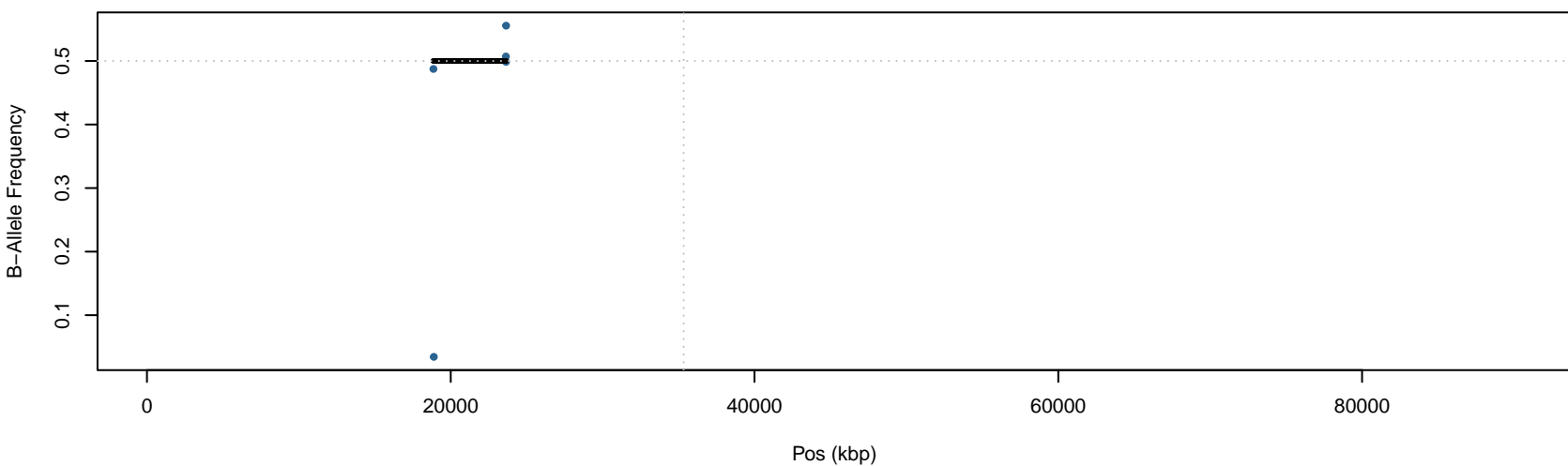
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 14



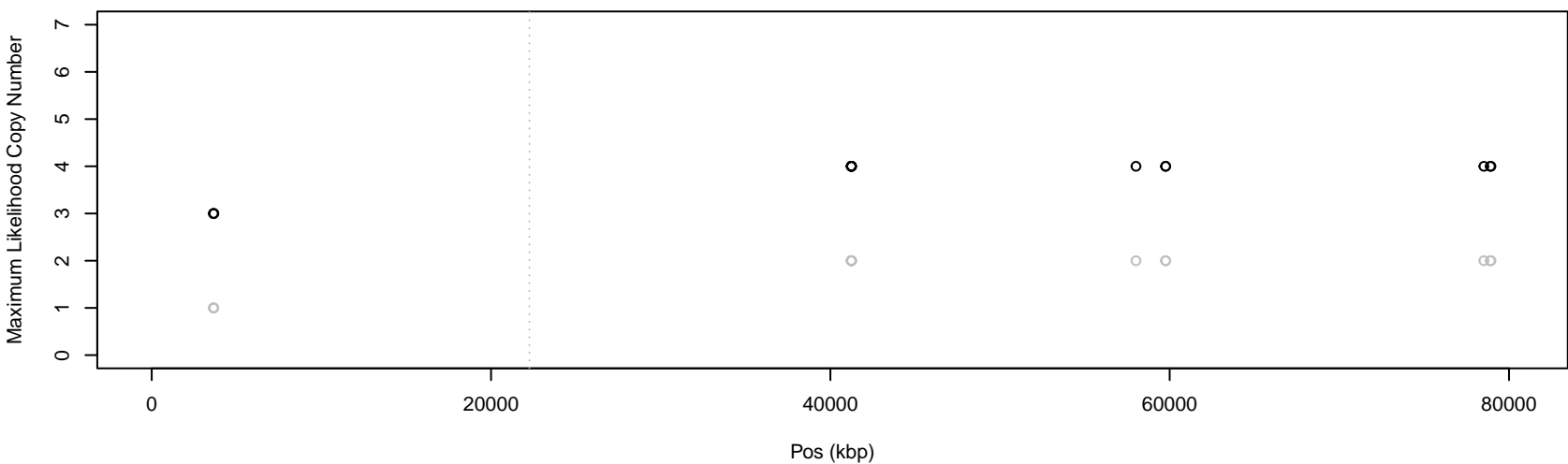
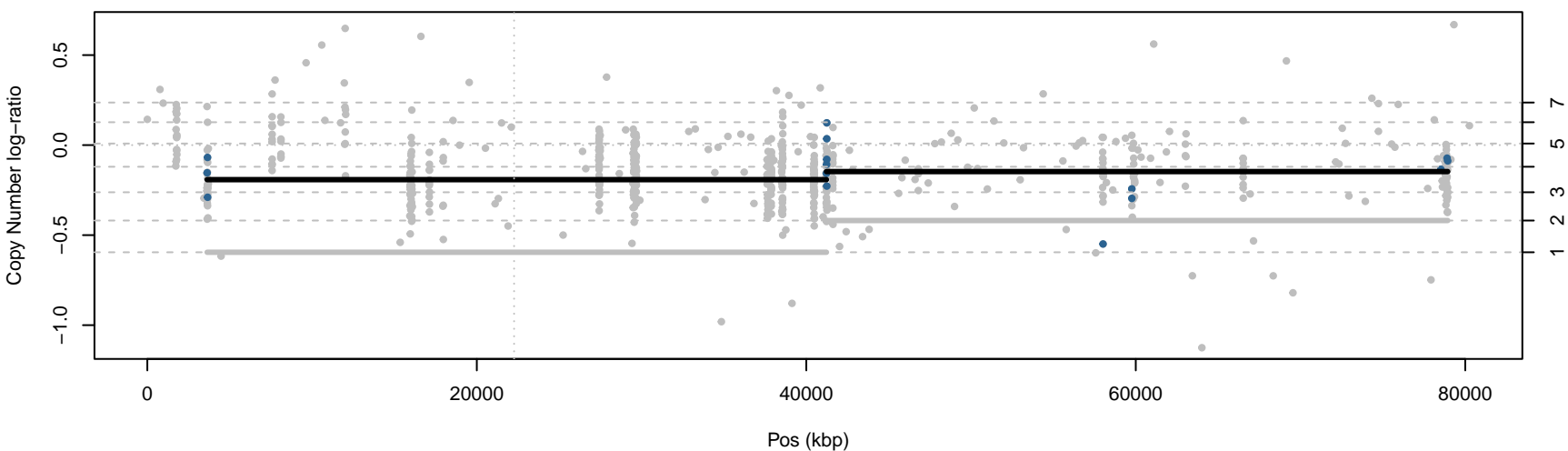
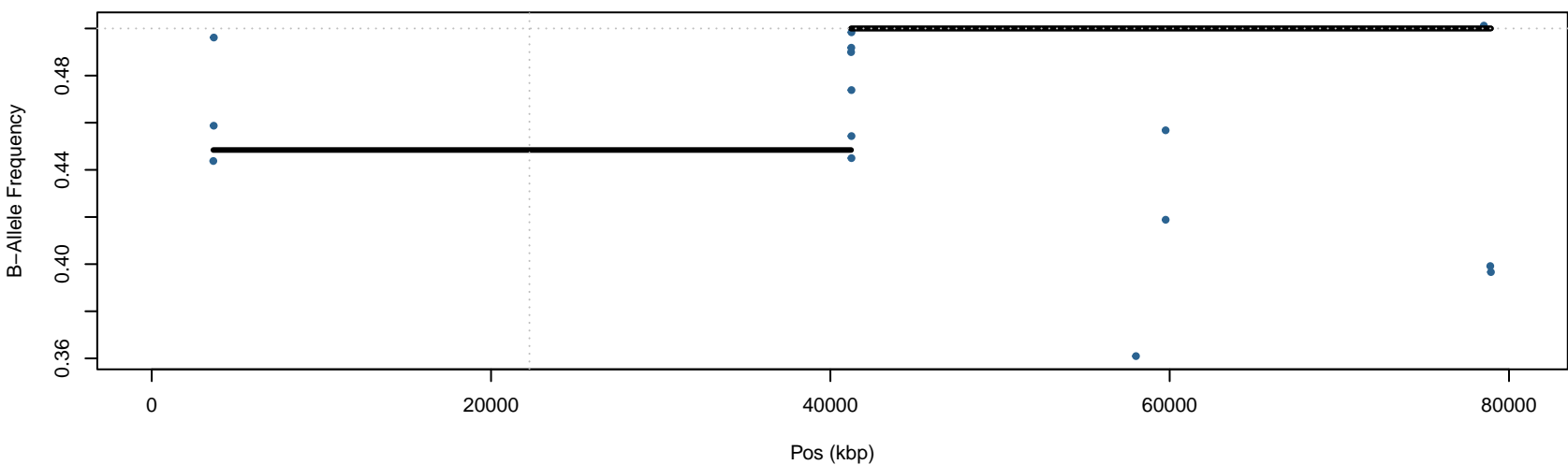
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 15



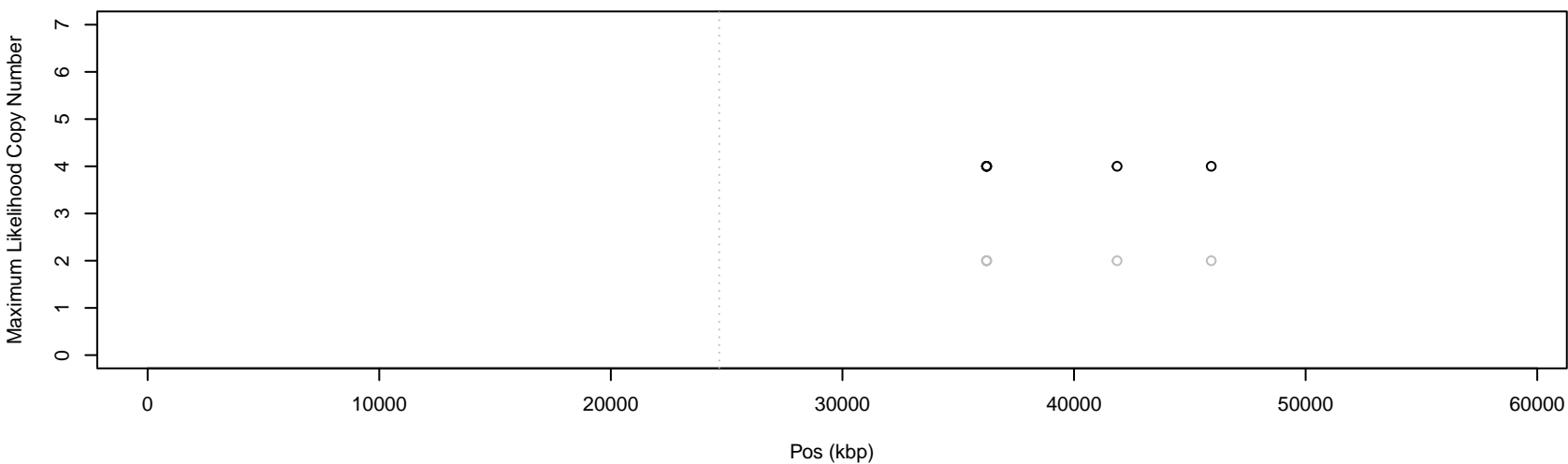
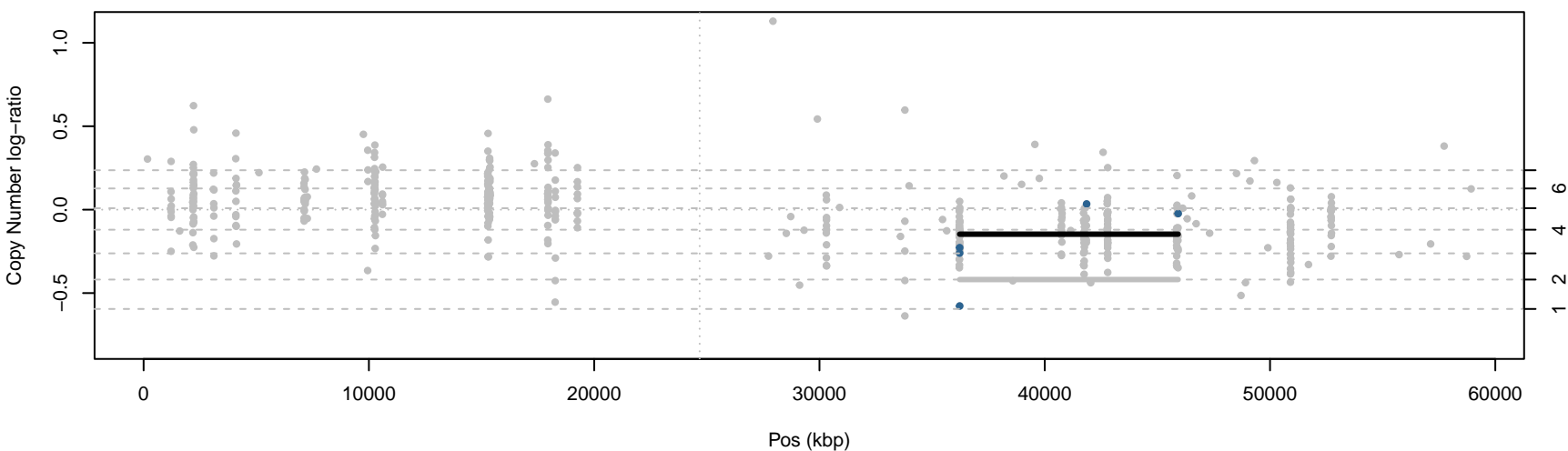
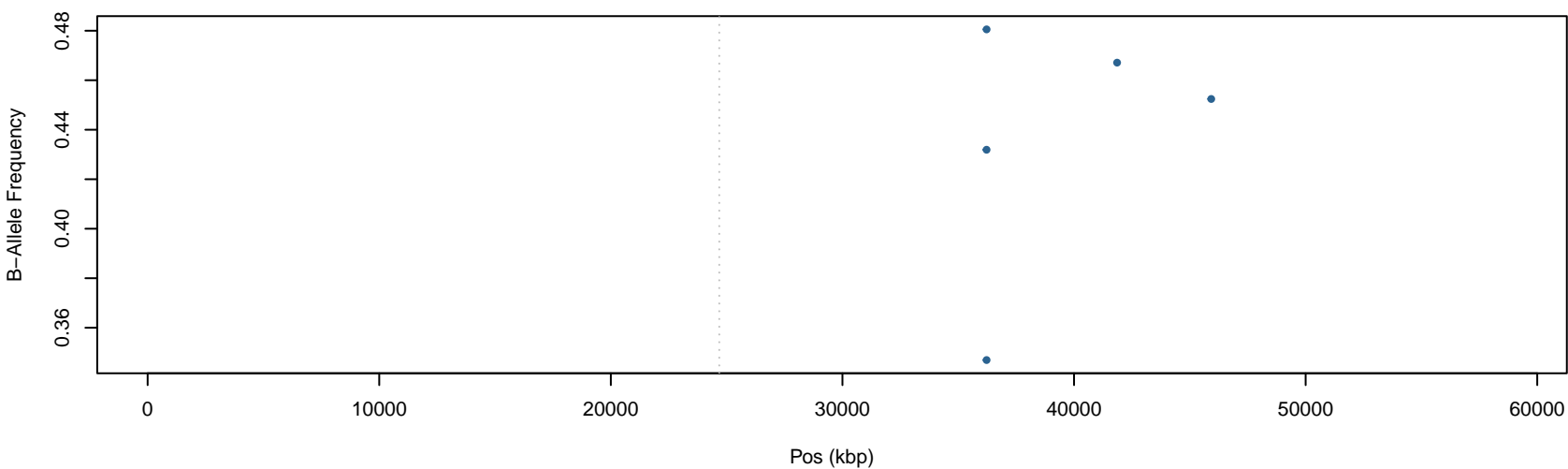
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 16



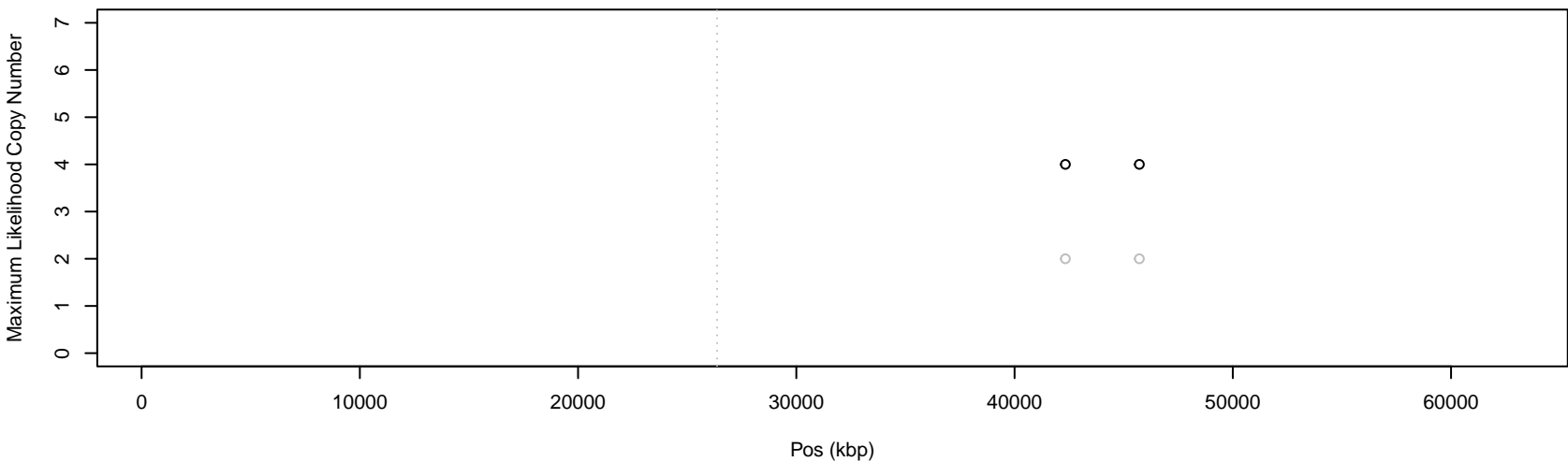
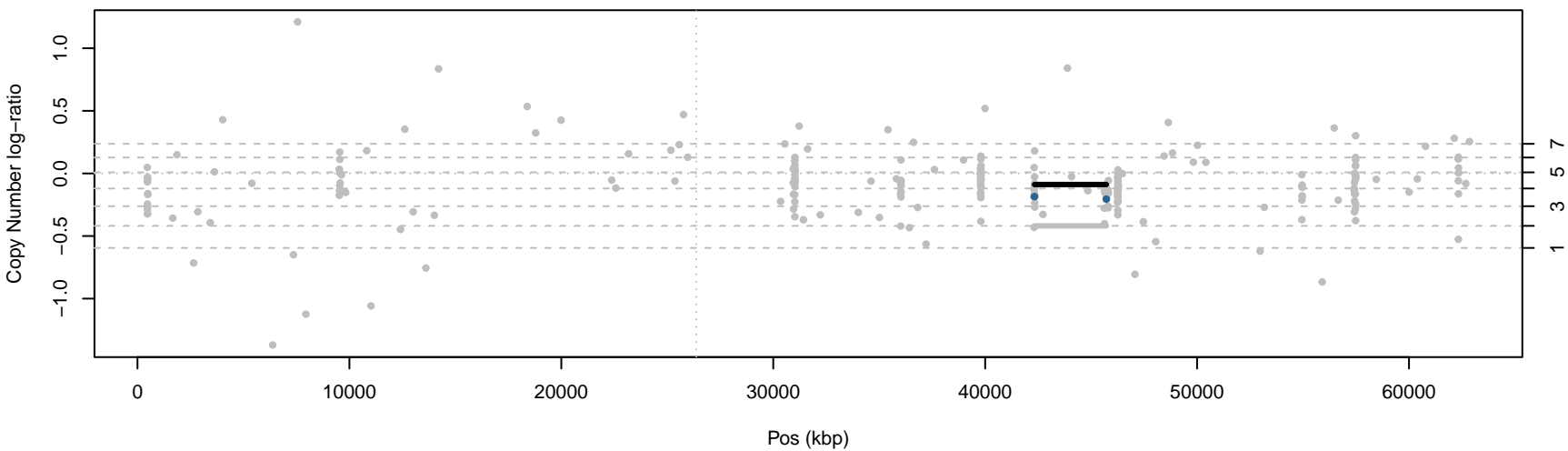
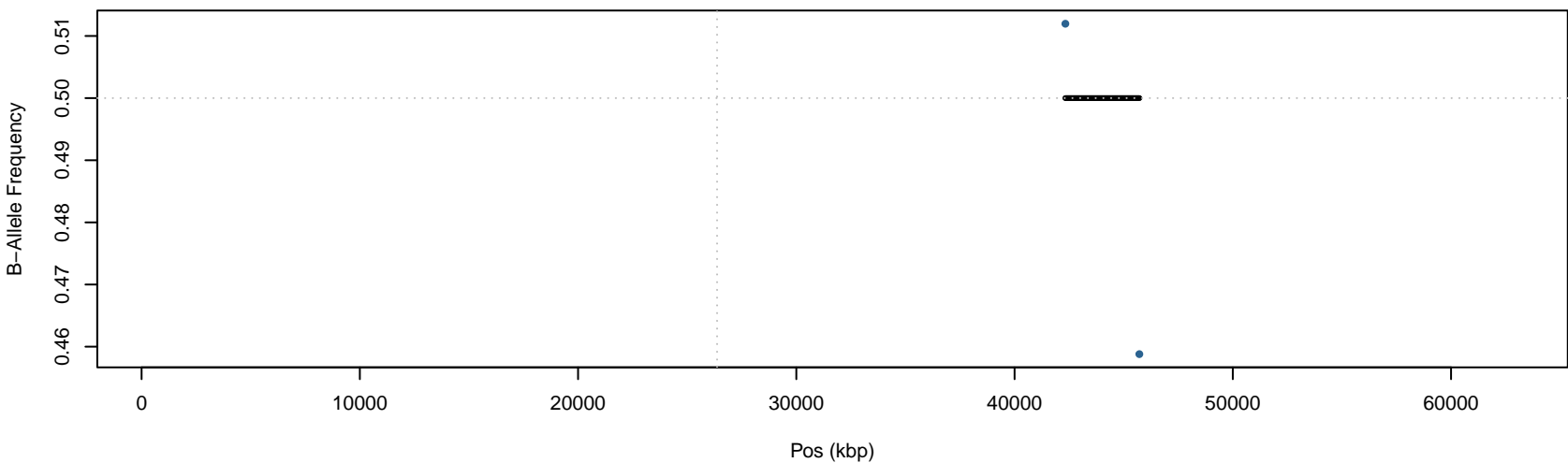
Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 17



Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 19



Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 20





Purity: 0.23 Tumor ploidy: 4.93 SNV log-likelihood: -40.11 GoF: 95.4% Mean coverage: 573;766 Chromosome: 21

