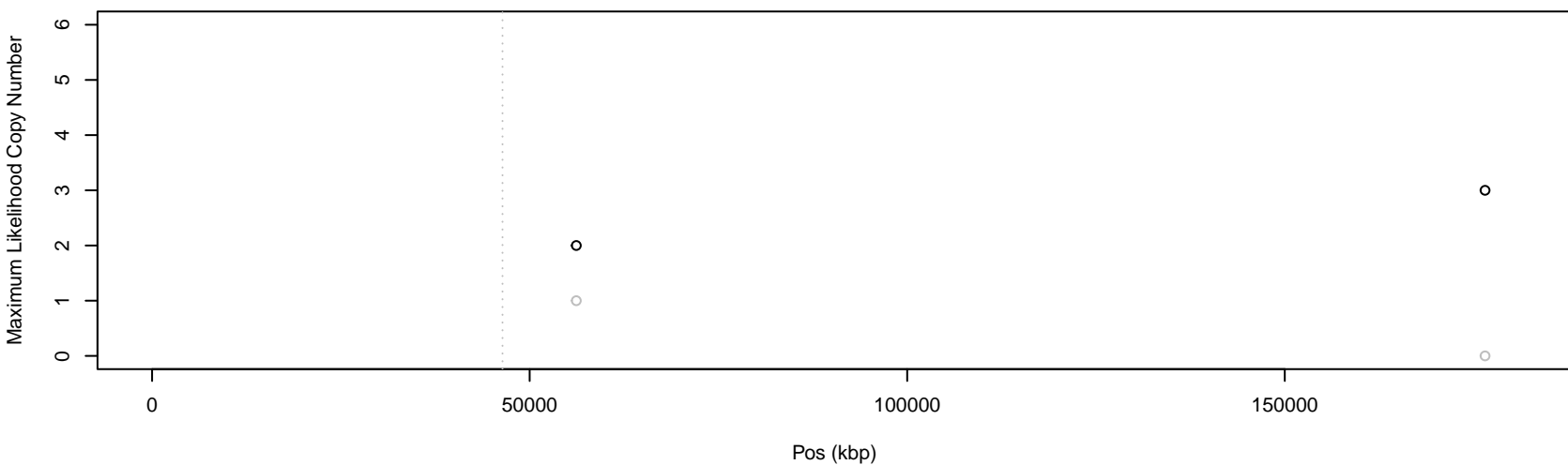
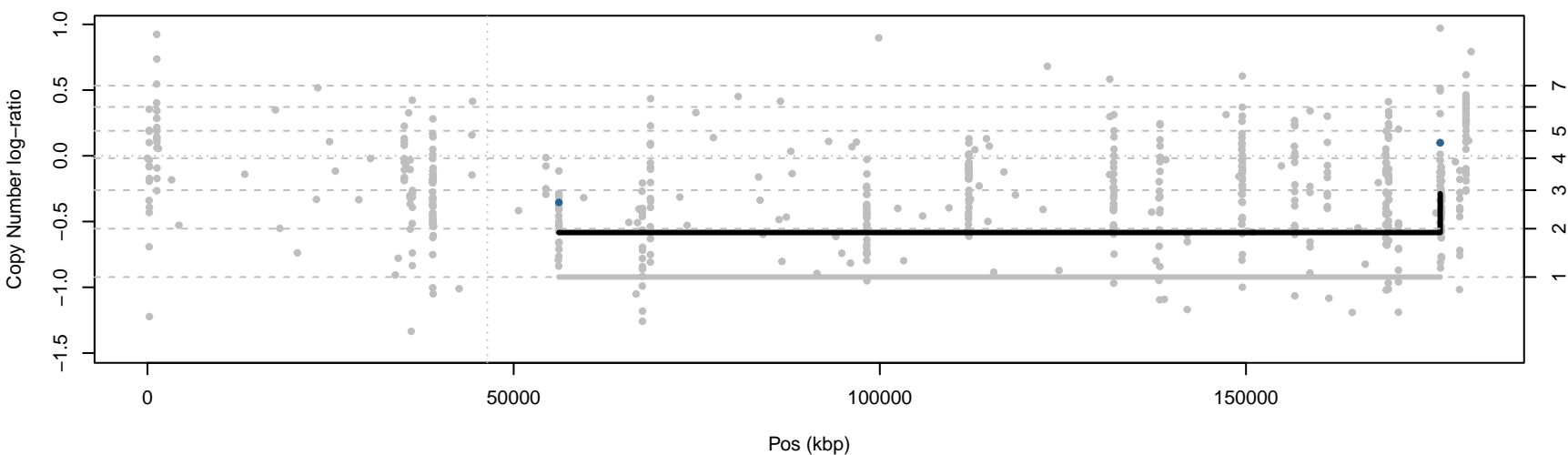
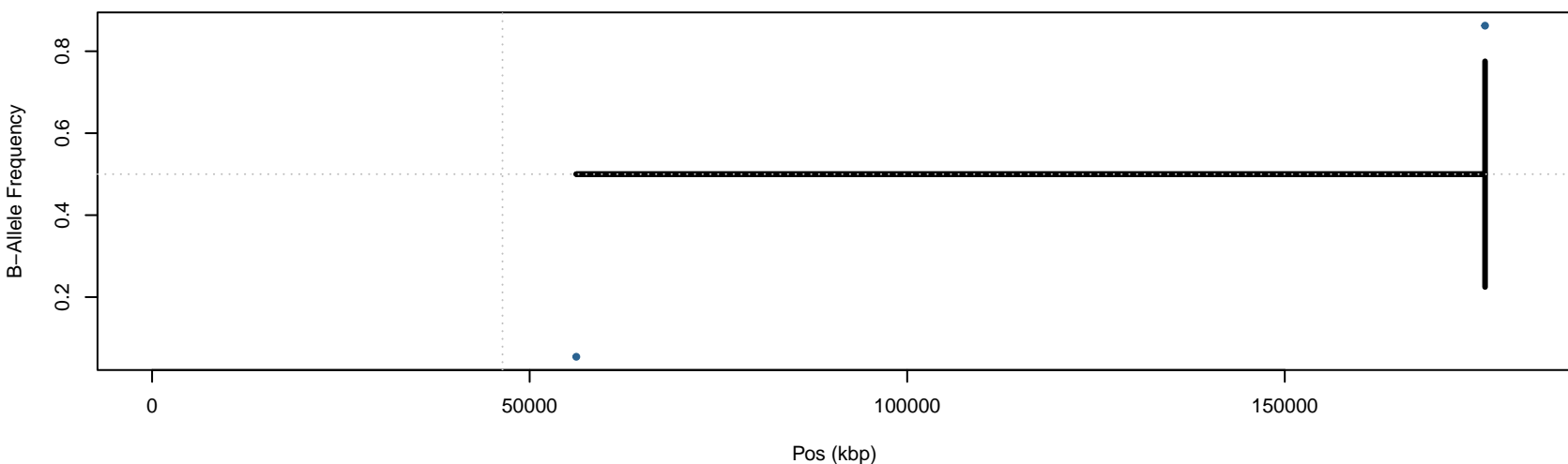
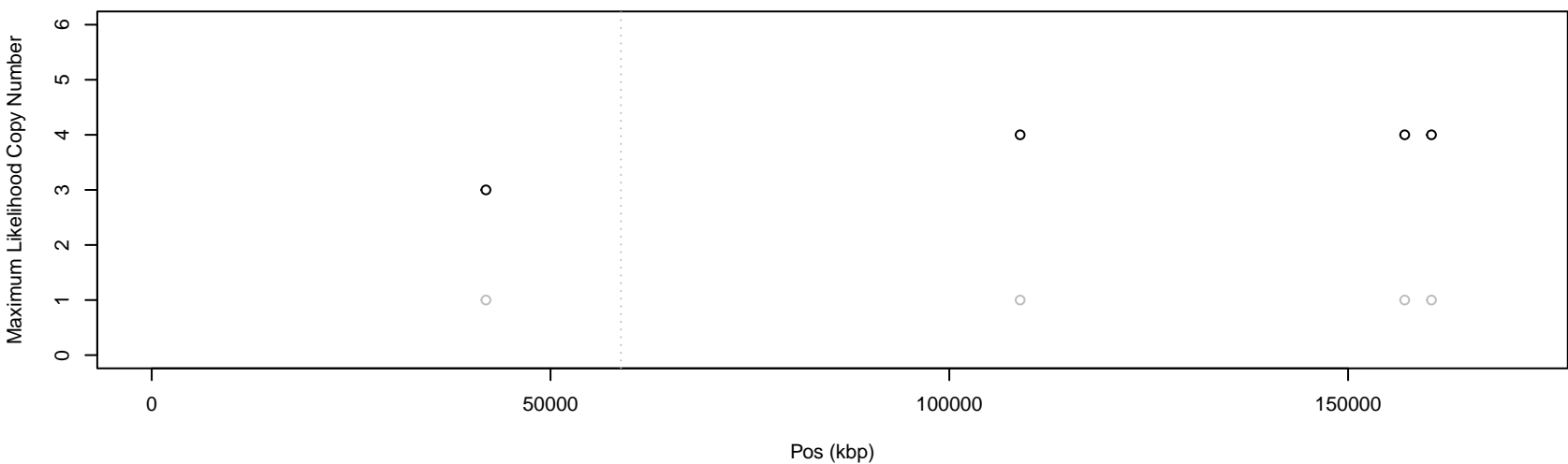
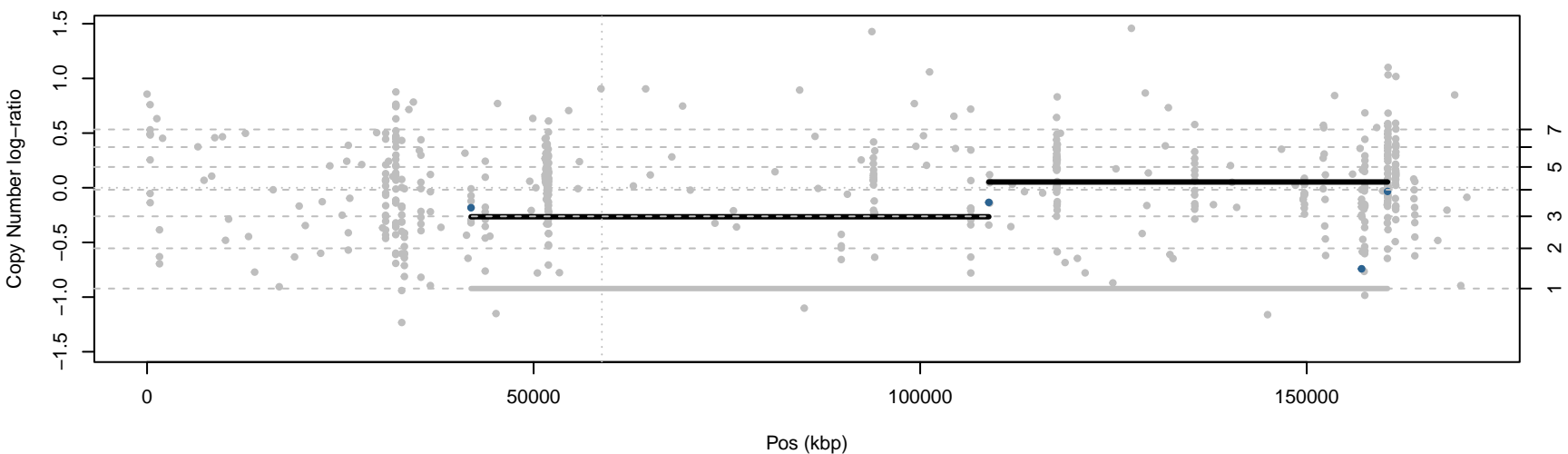
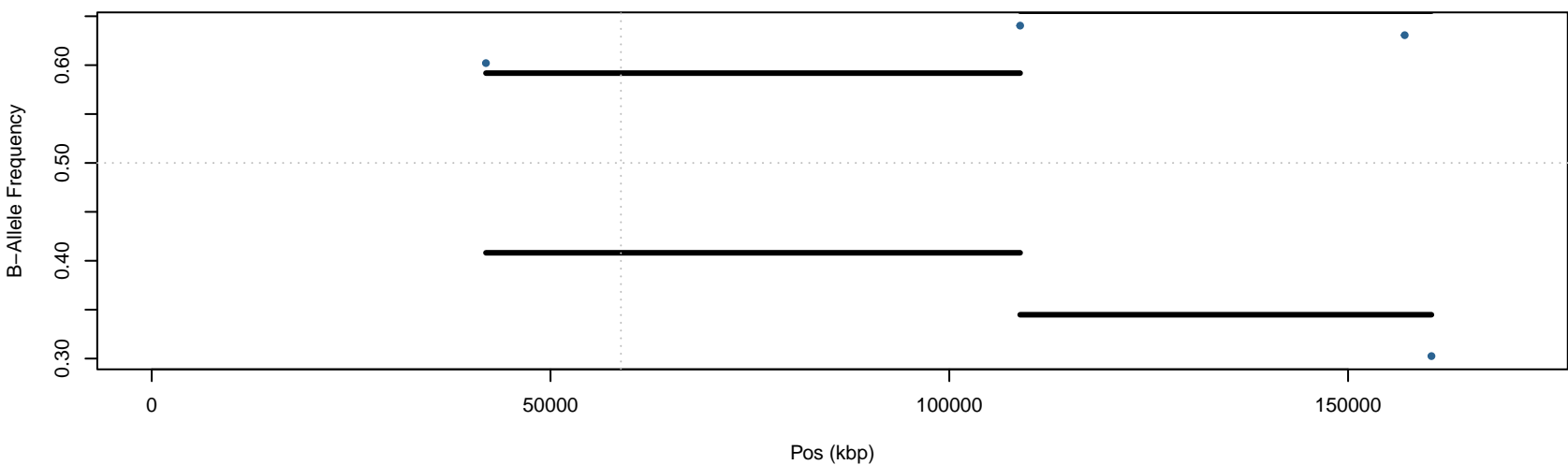


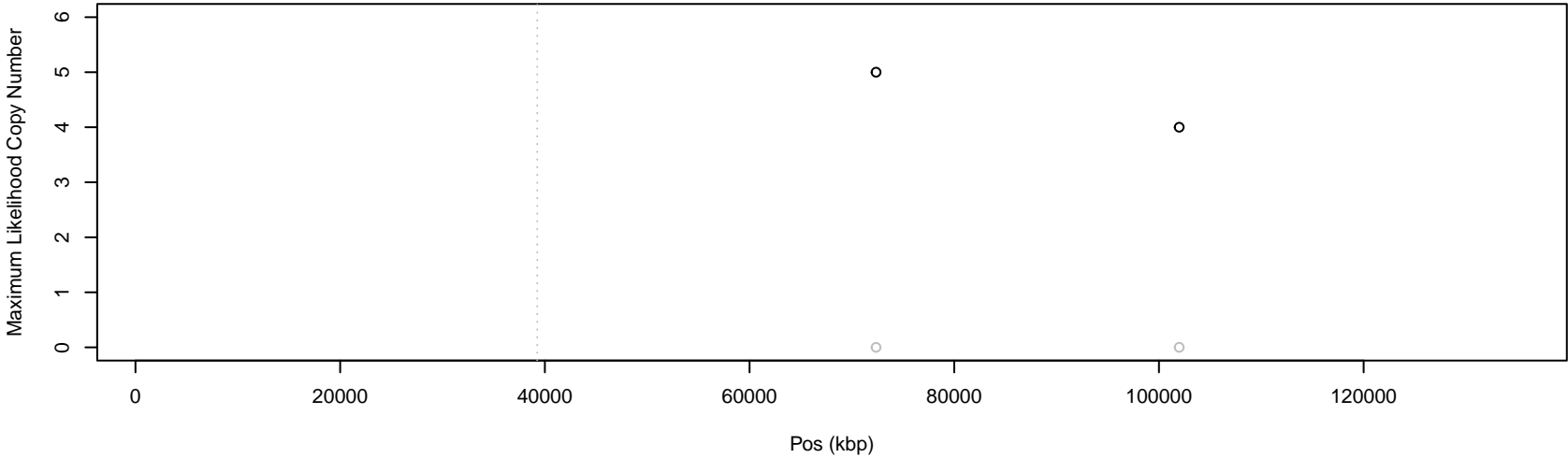
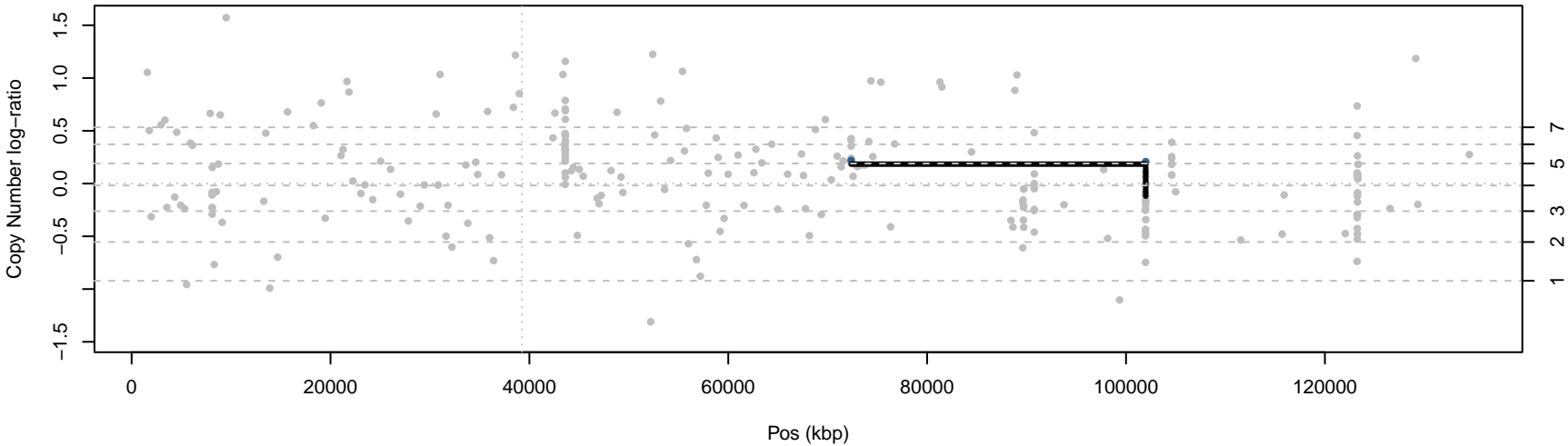
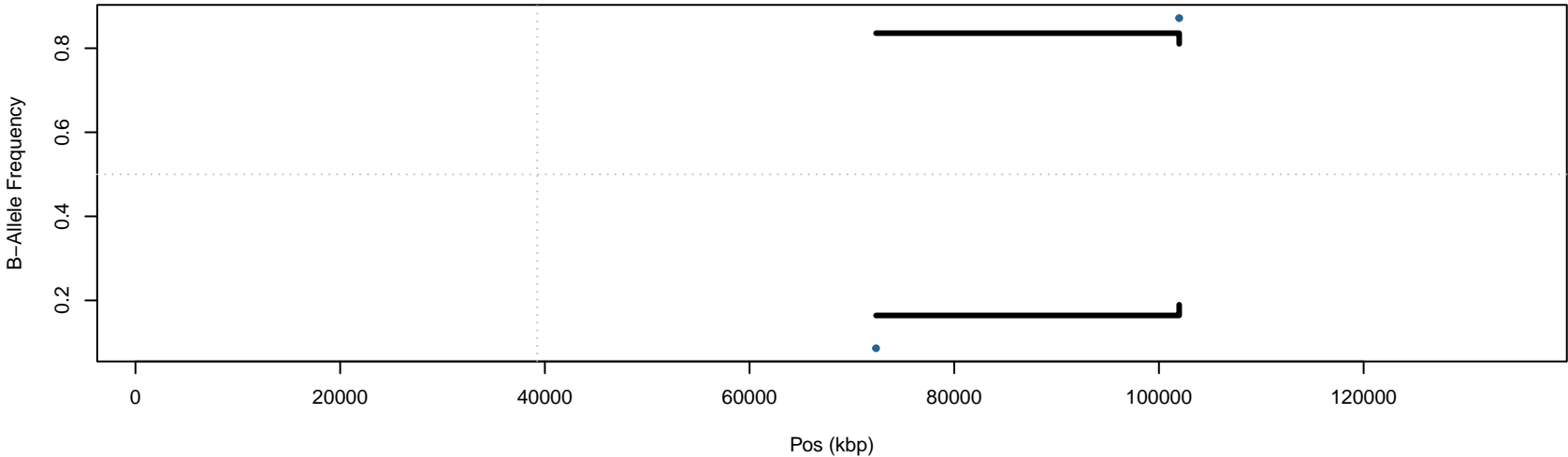
Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 5



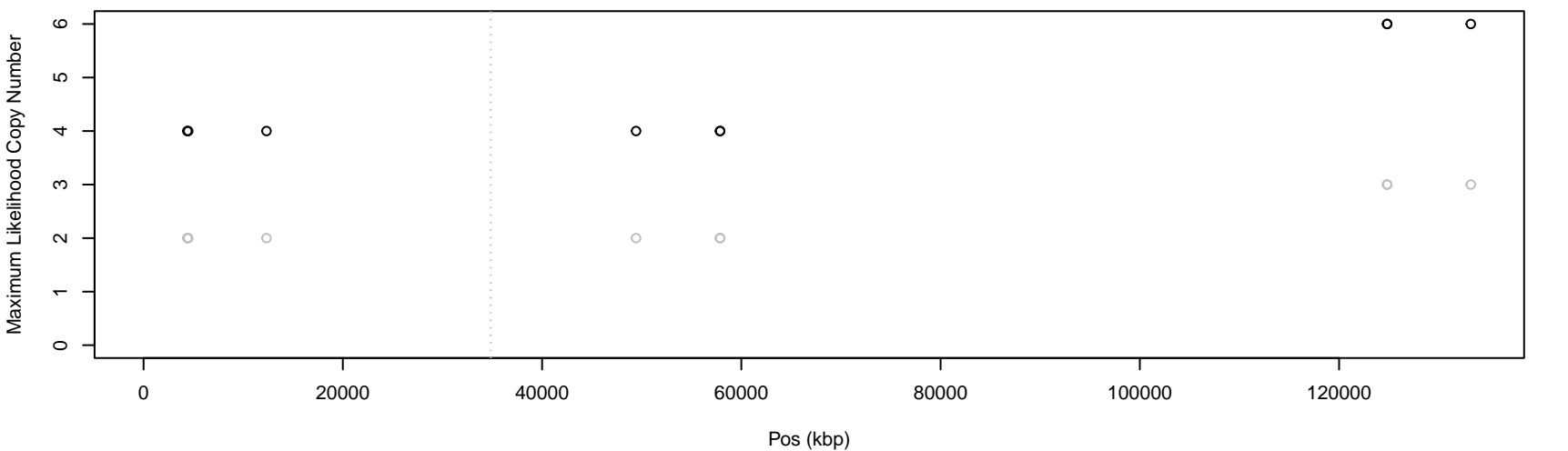
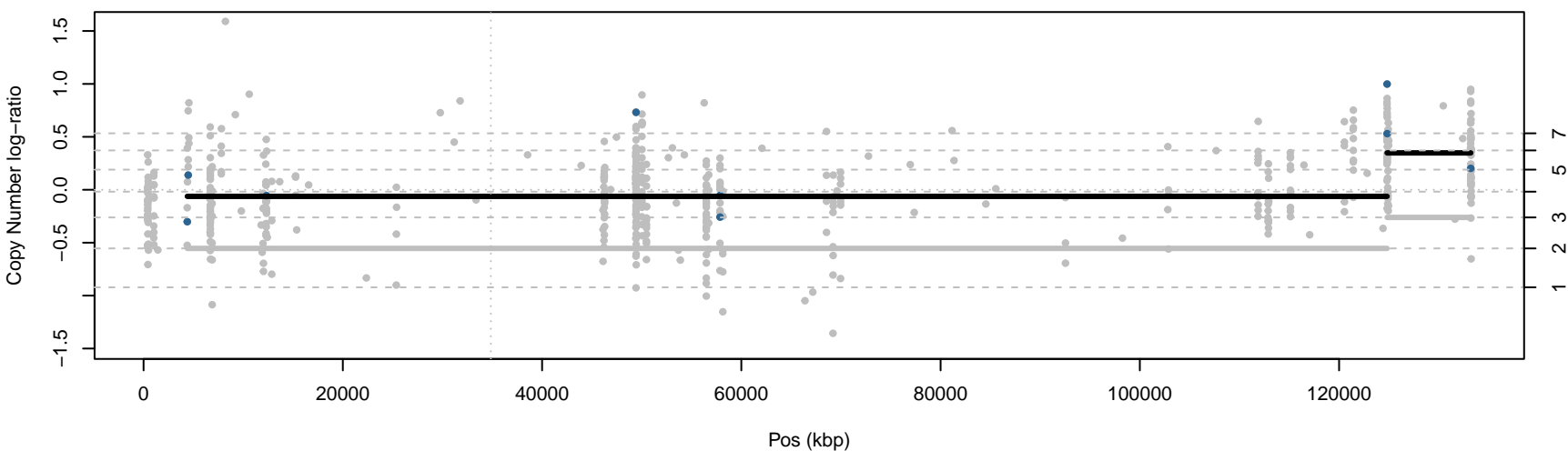
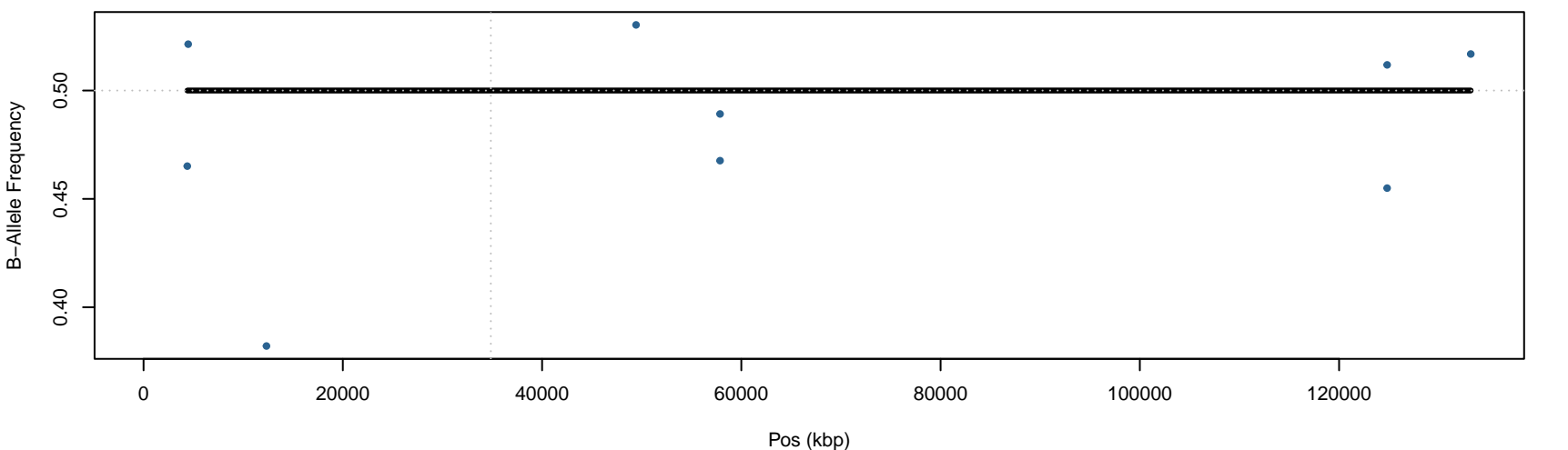
Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 6



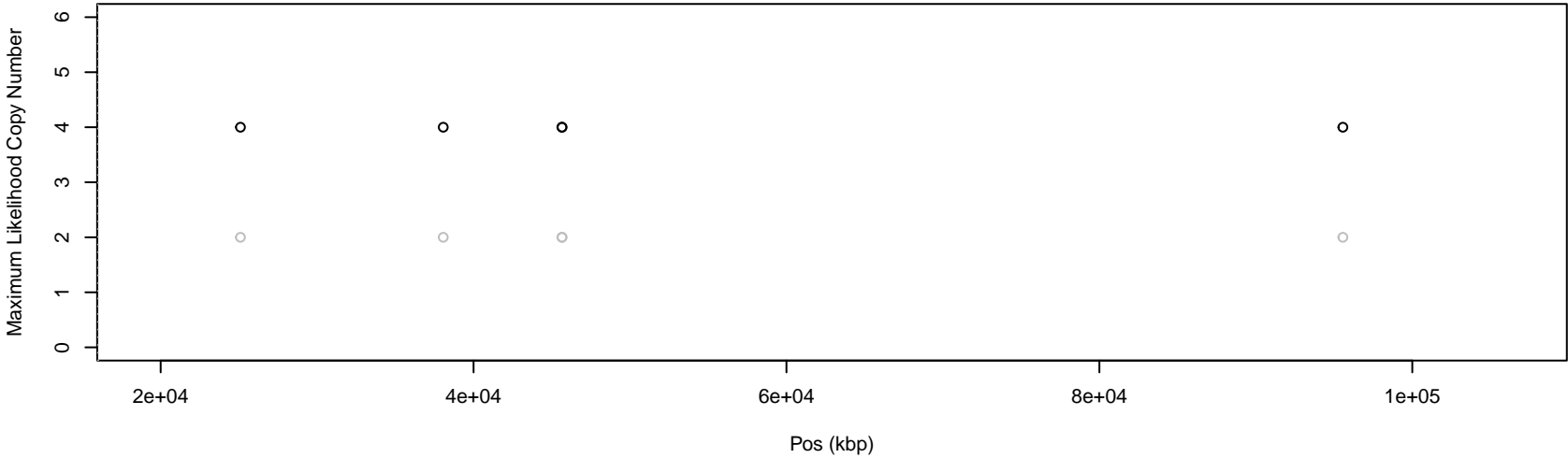
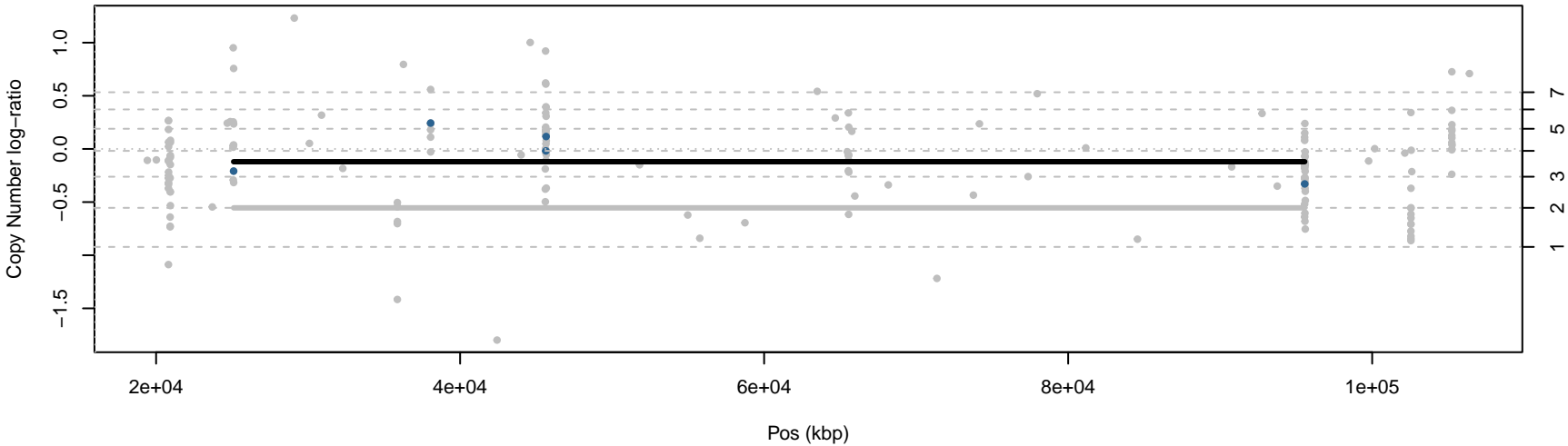
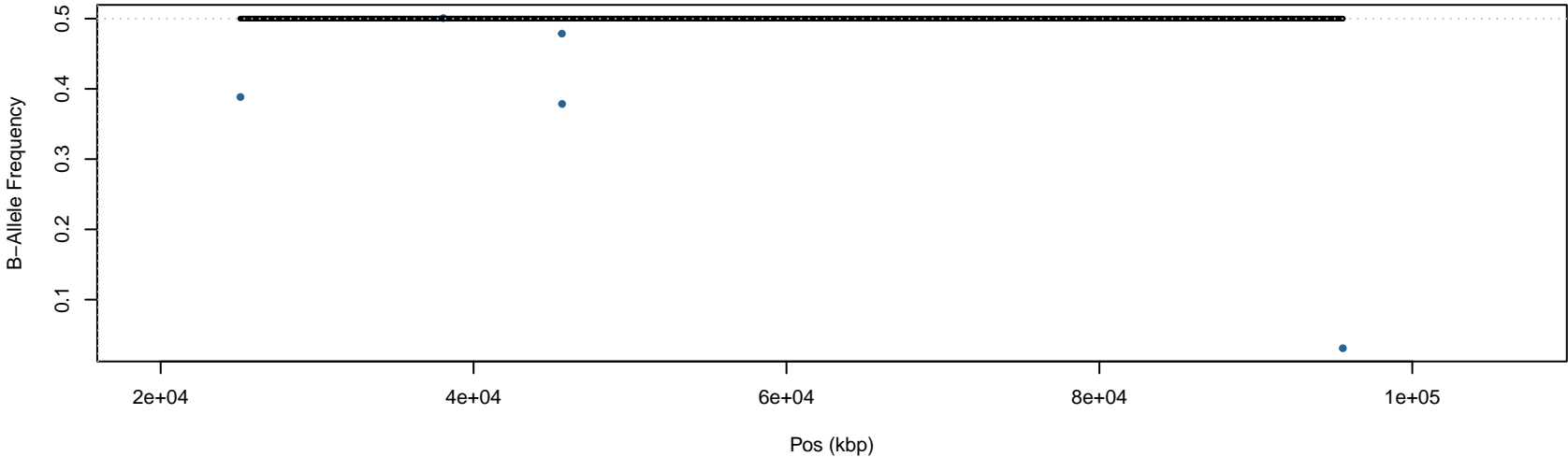
Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 10



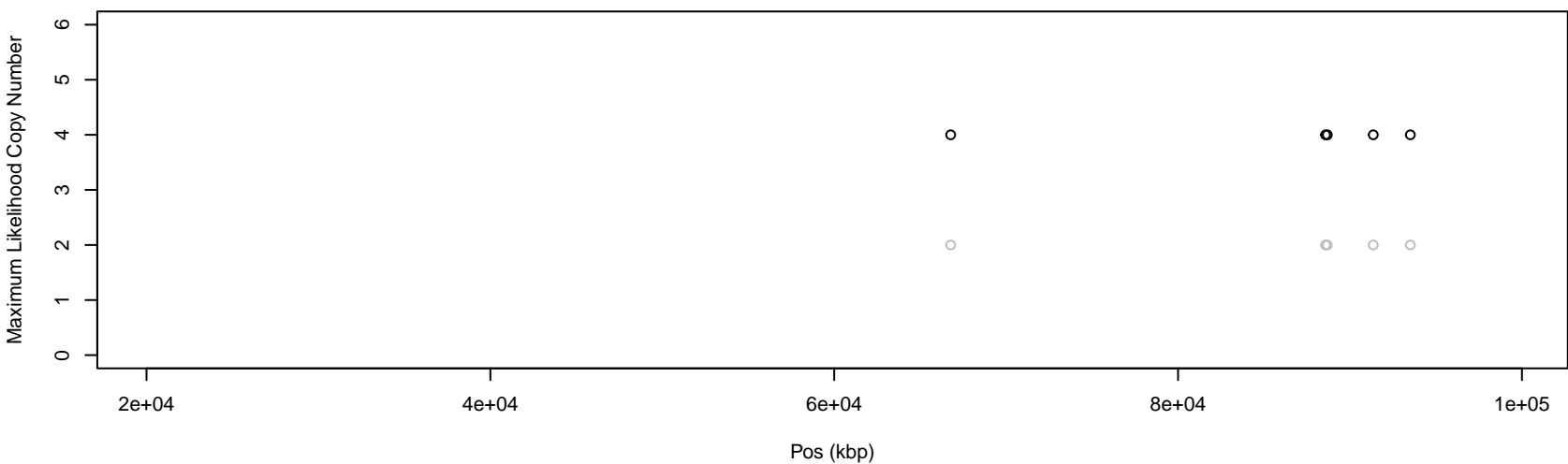
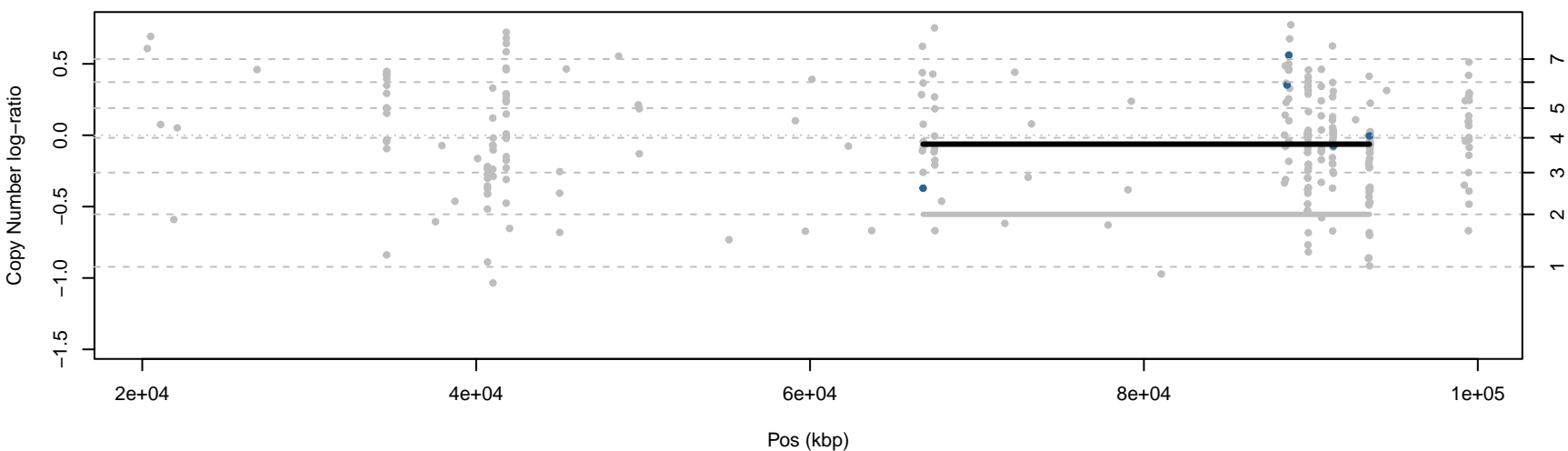
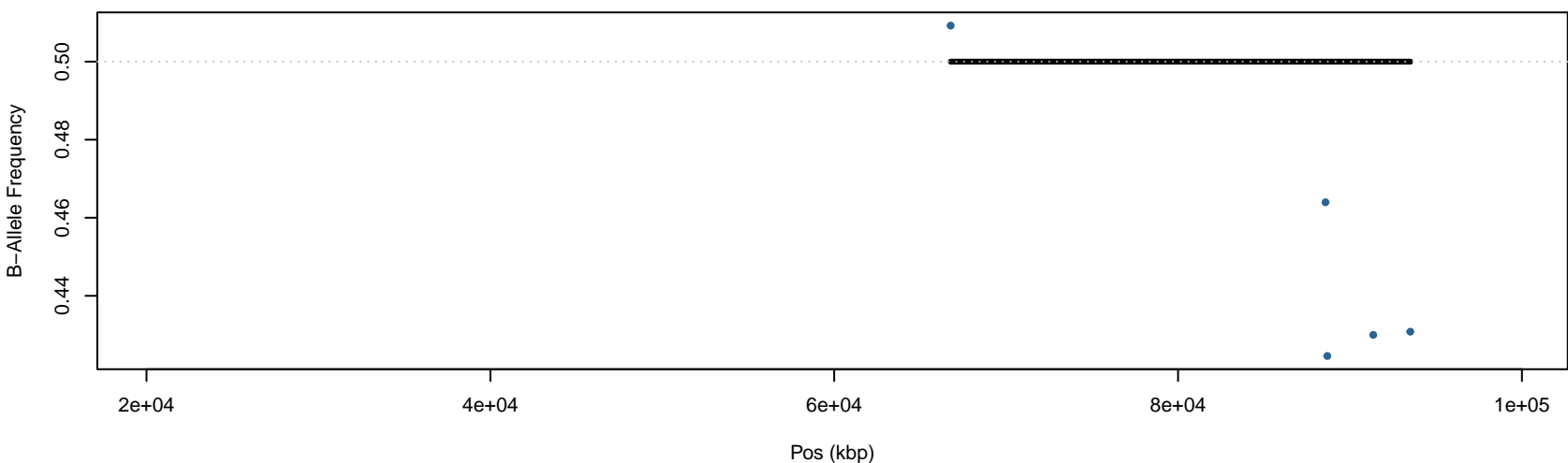
Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 12



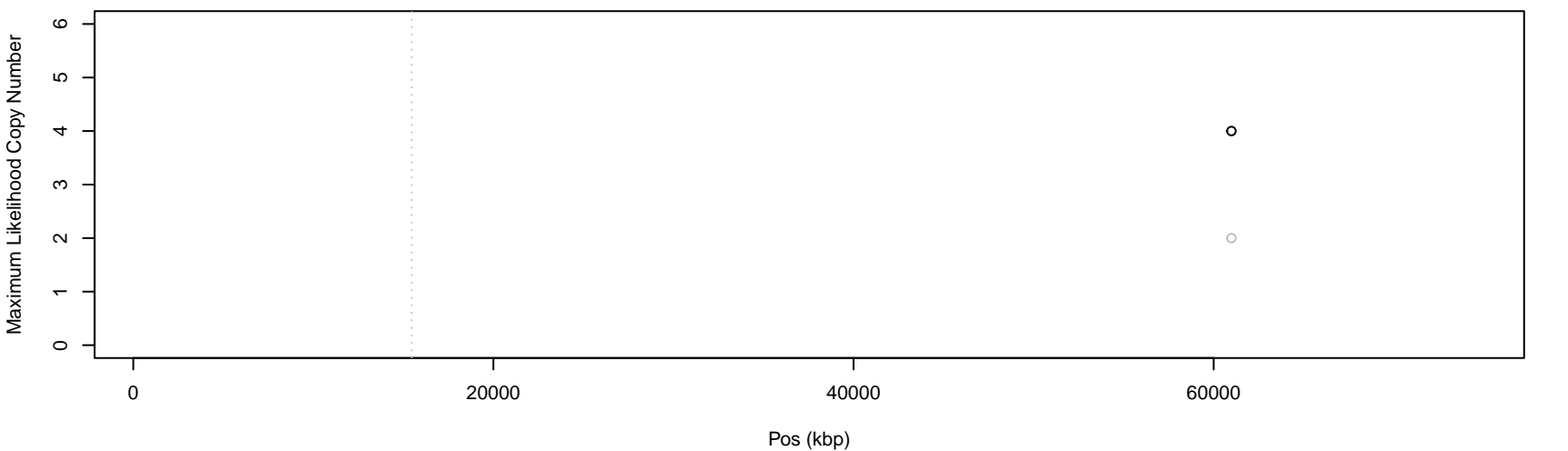
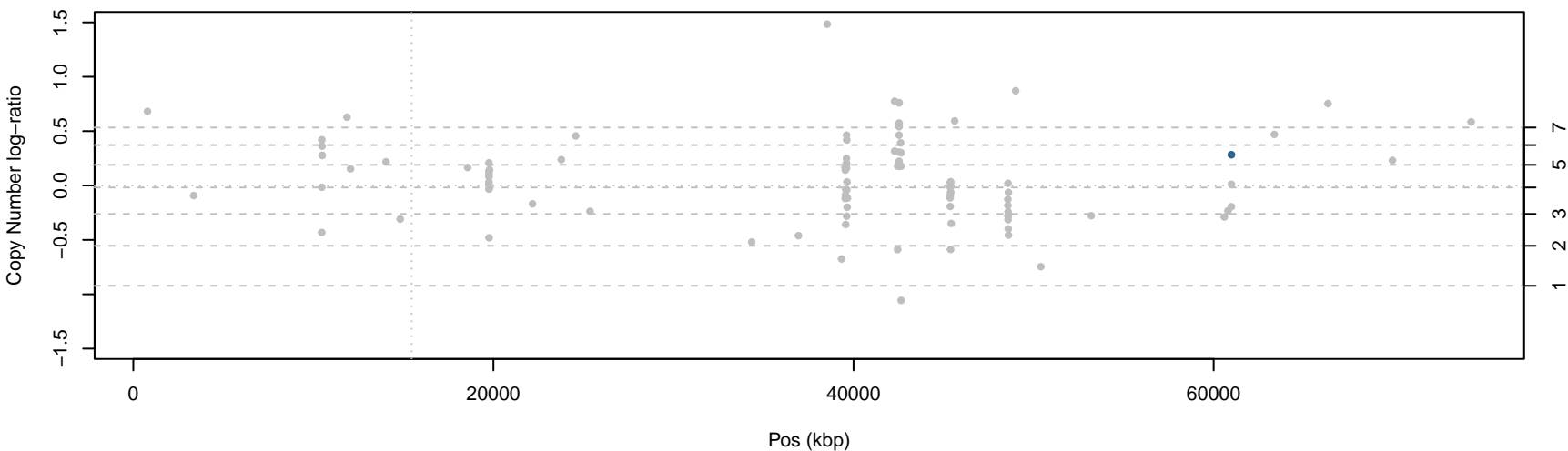
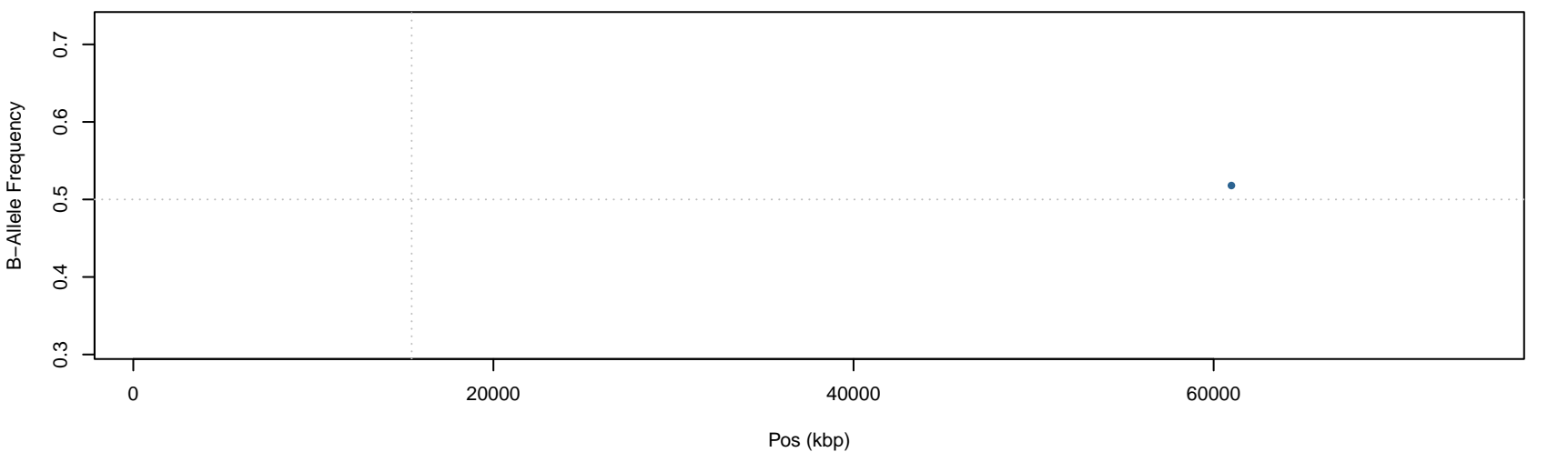
Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 14

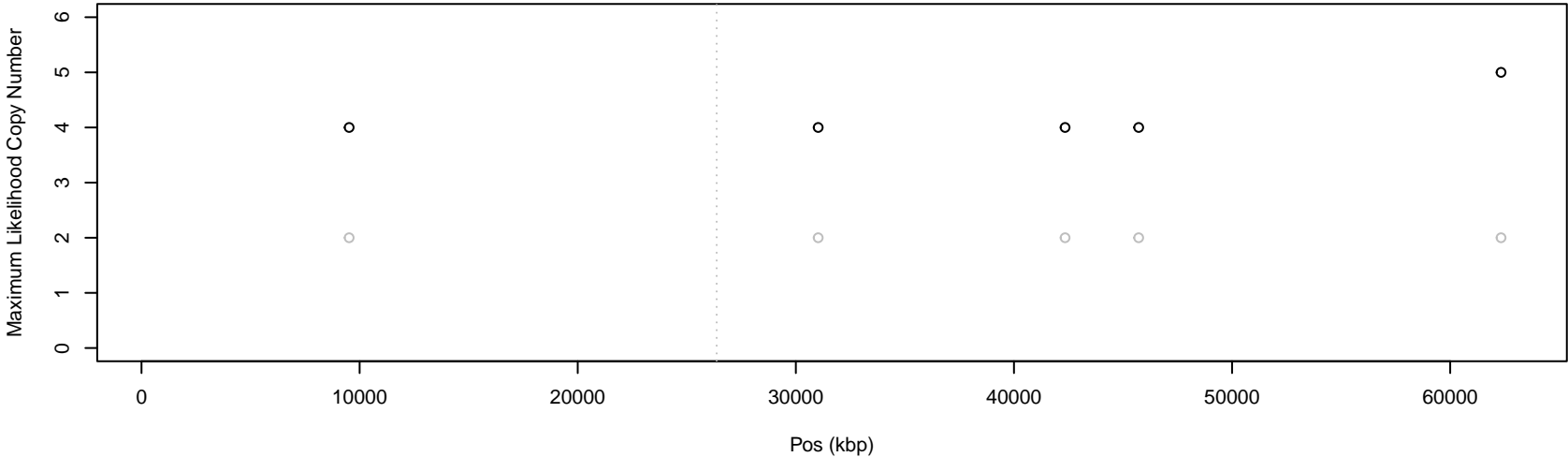
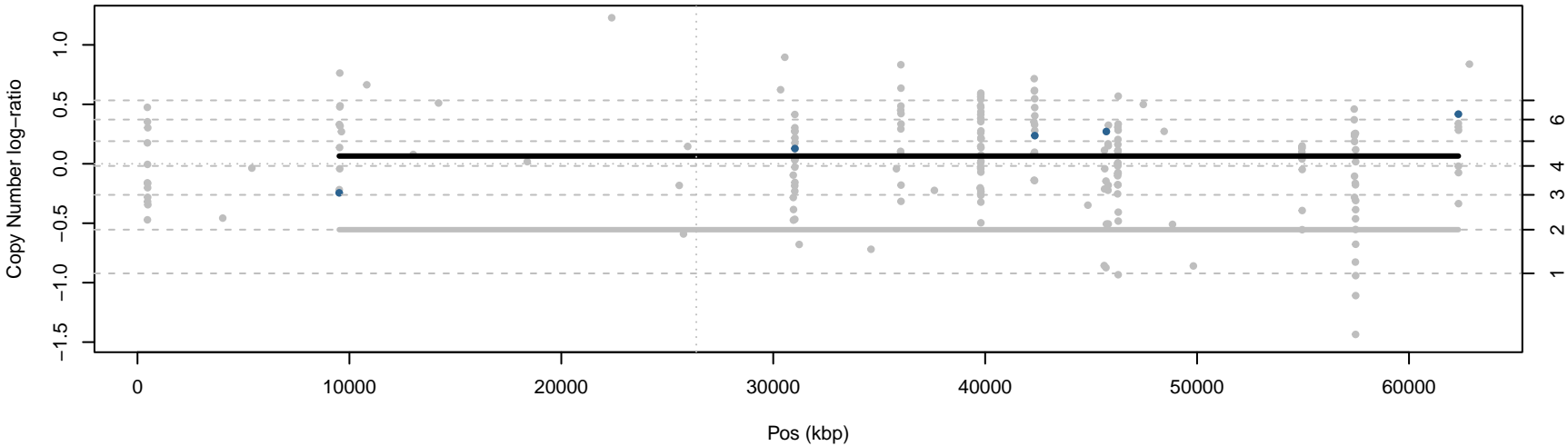
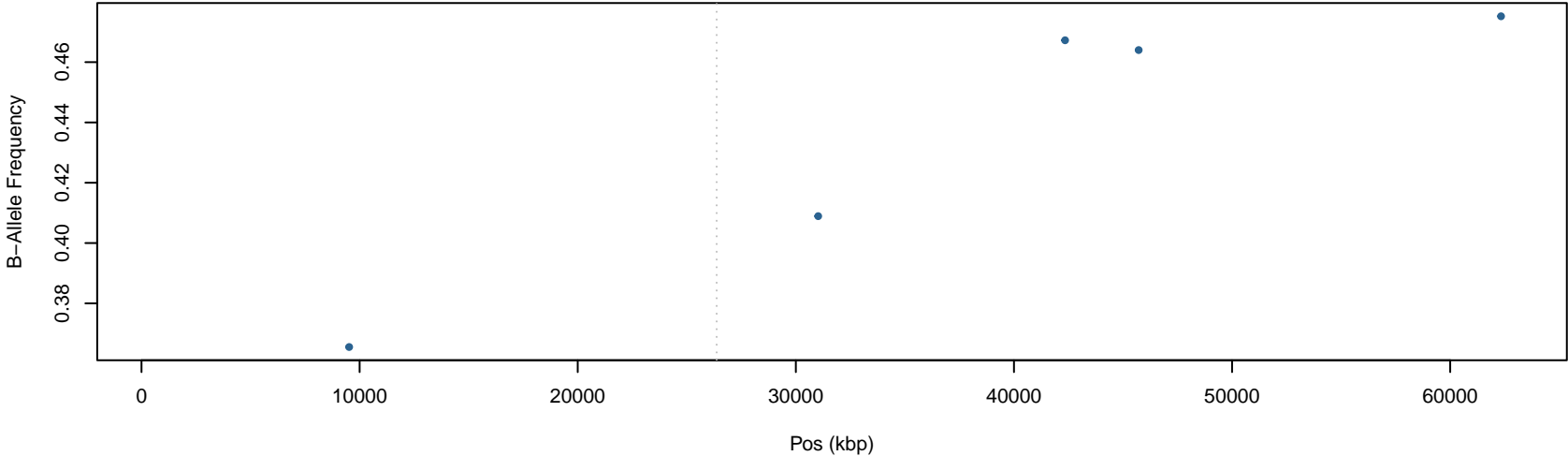


Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 15



Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 18







Purity: 0.45 Tumor ploidy: 4.081 SNV log-likelihood: -122.67 GoF: 89.6% Mean coverage: 350;709 Chromosome: 22

