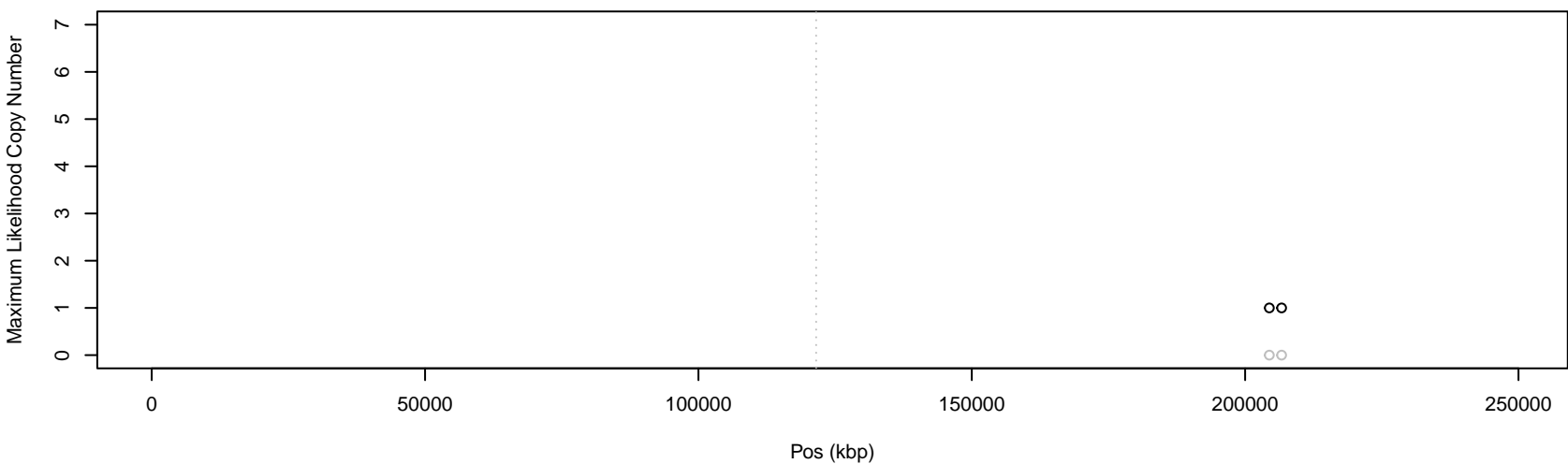
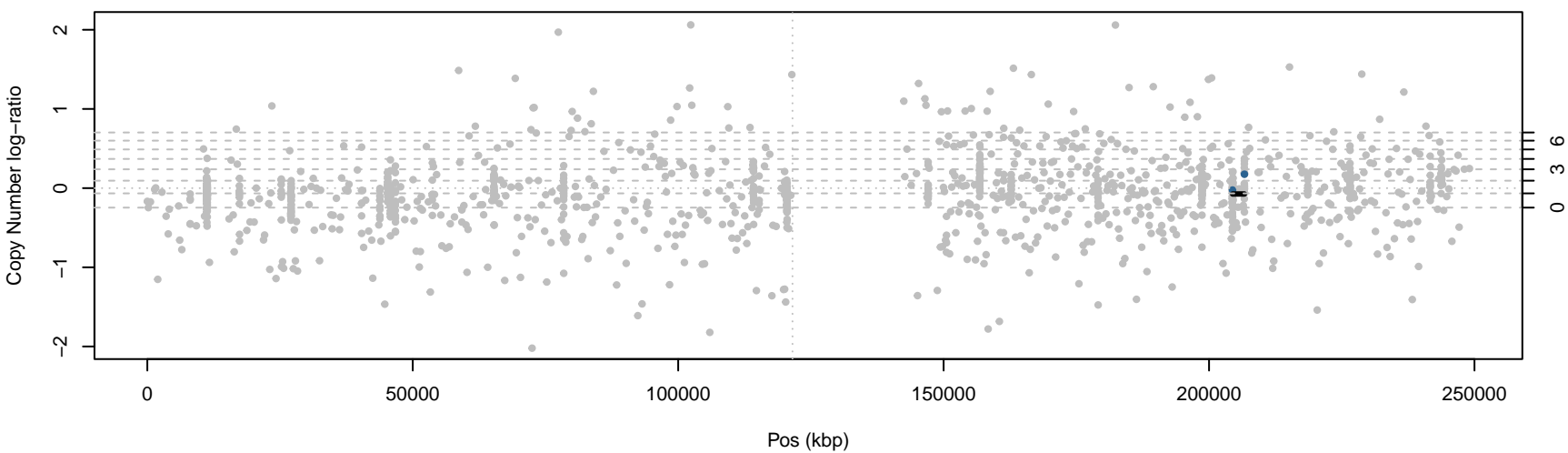
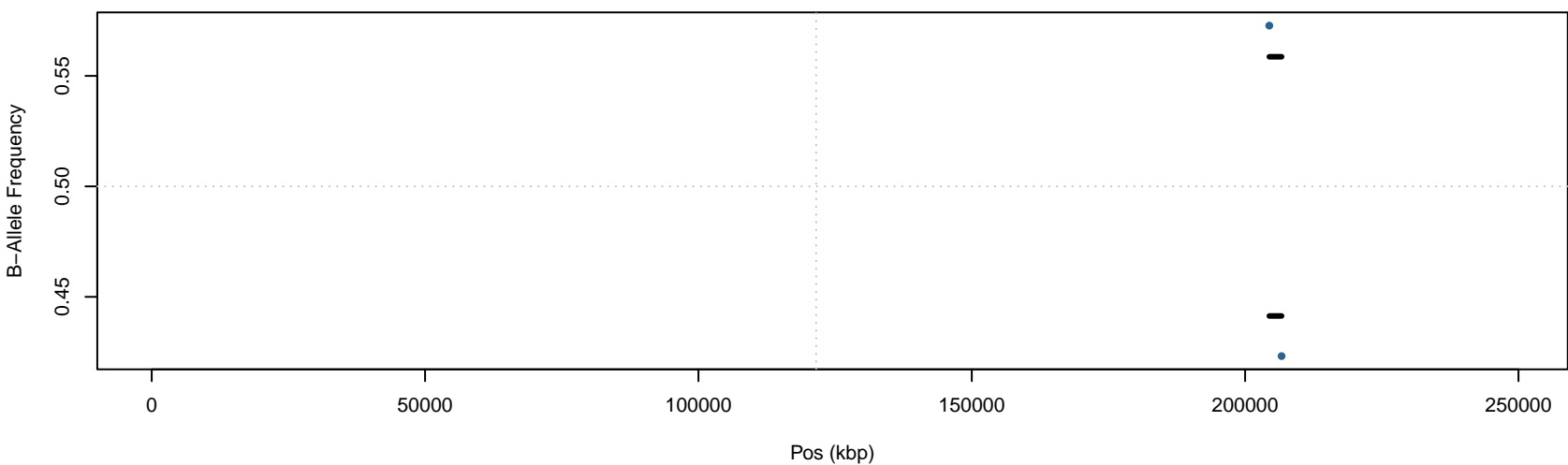
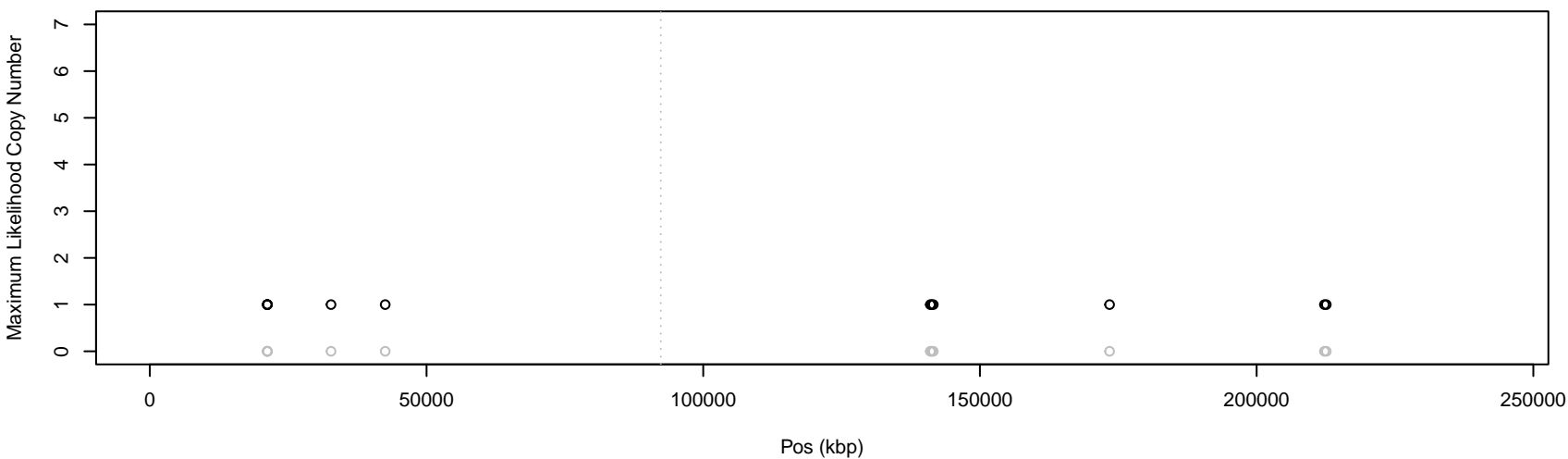
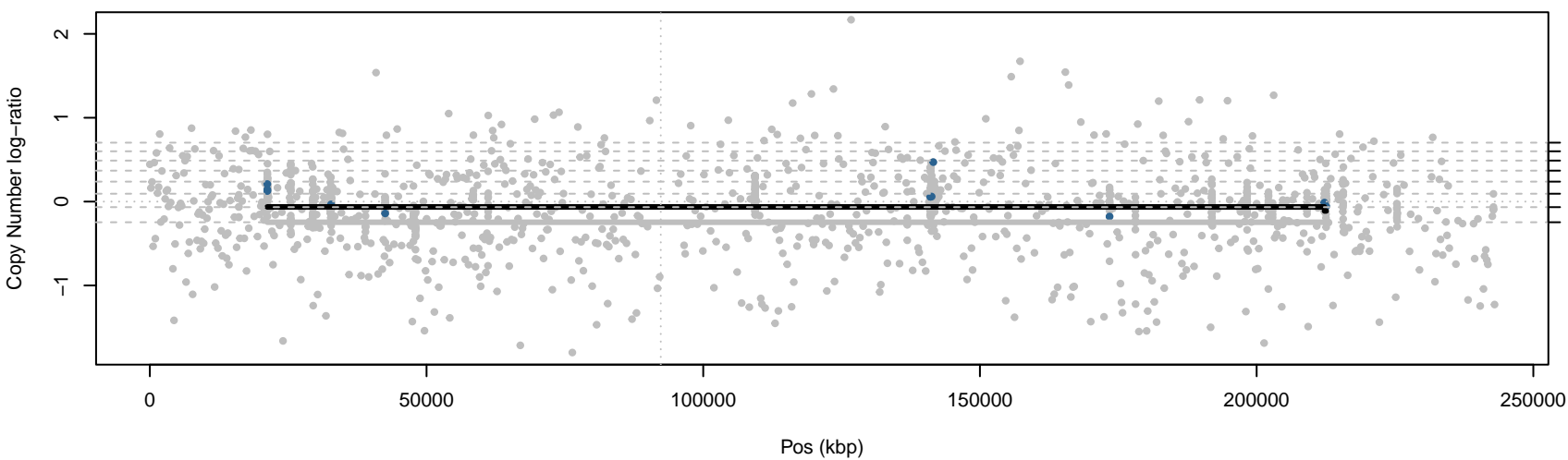
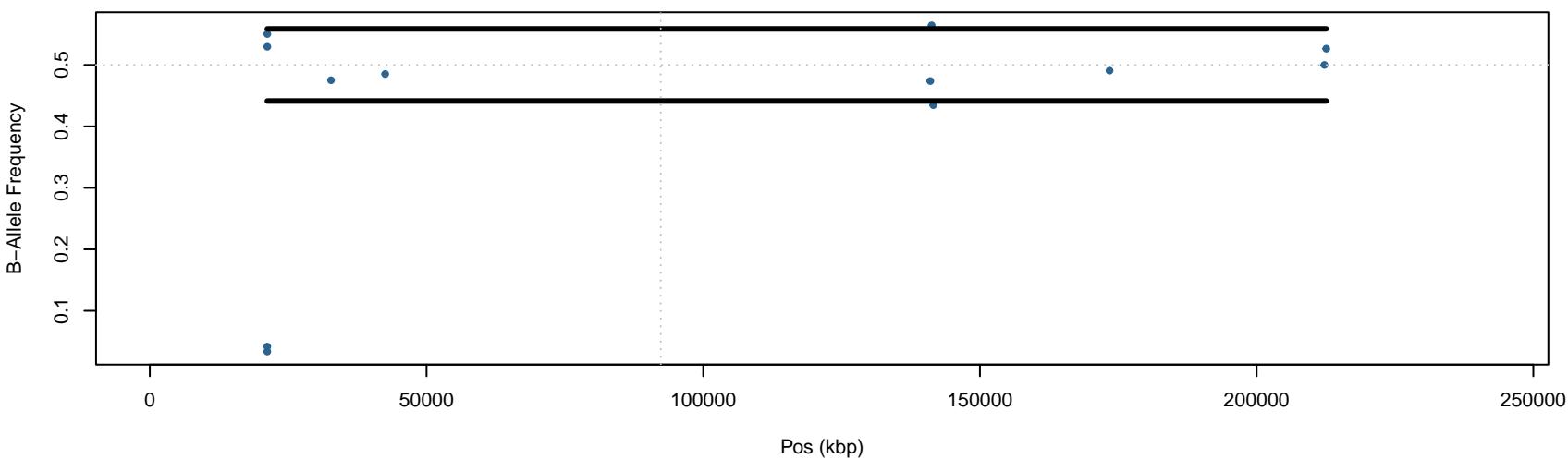


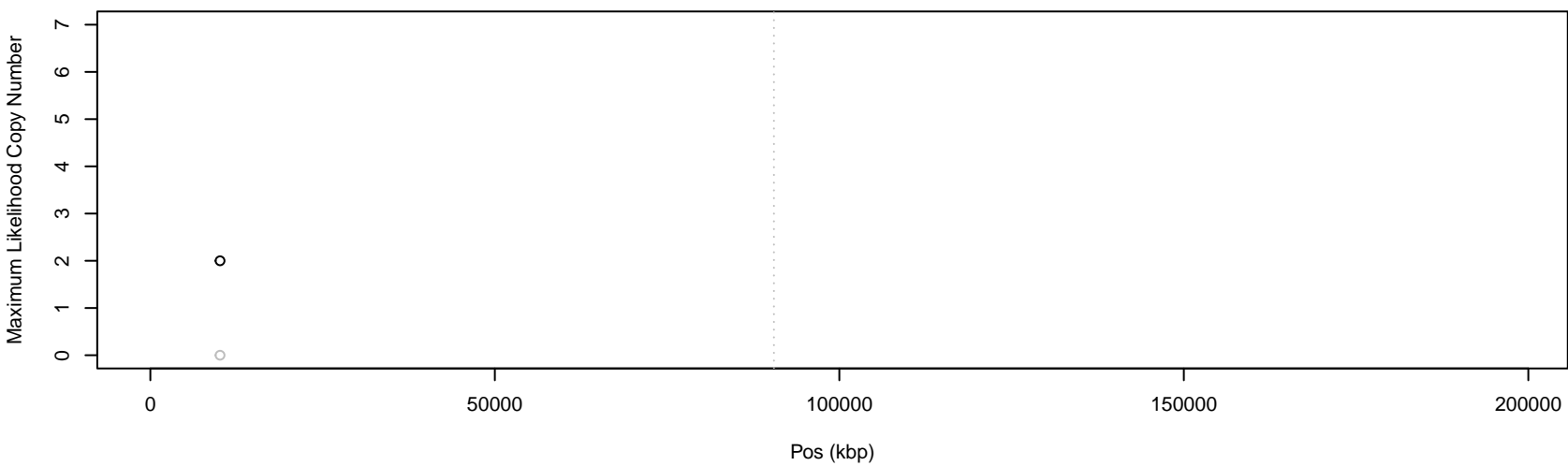
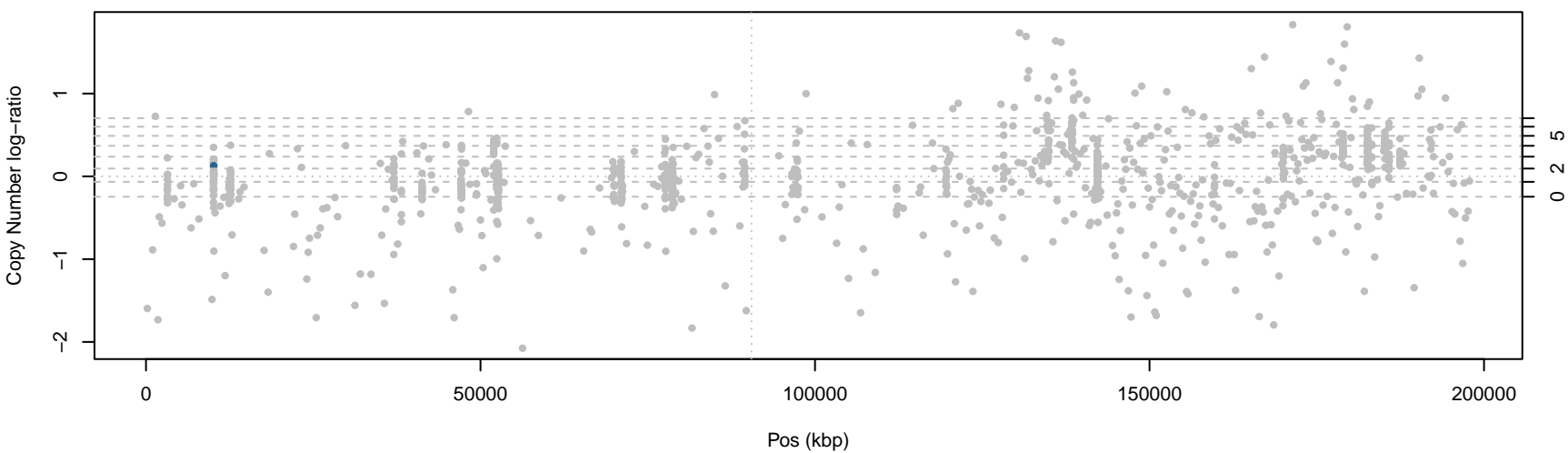
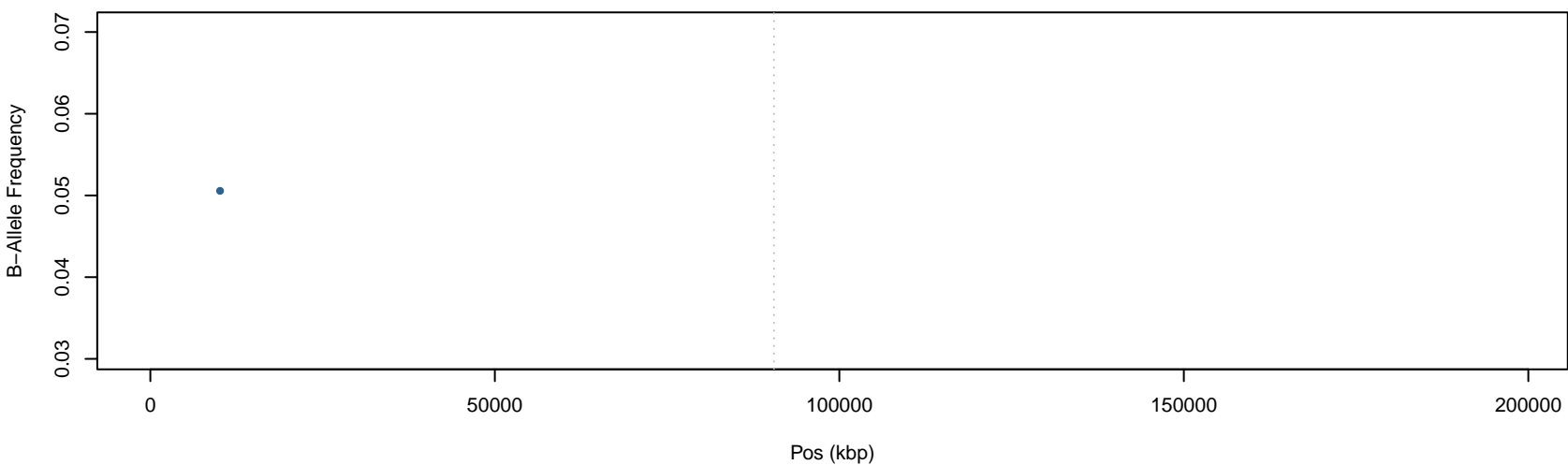
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 1



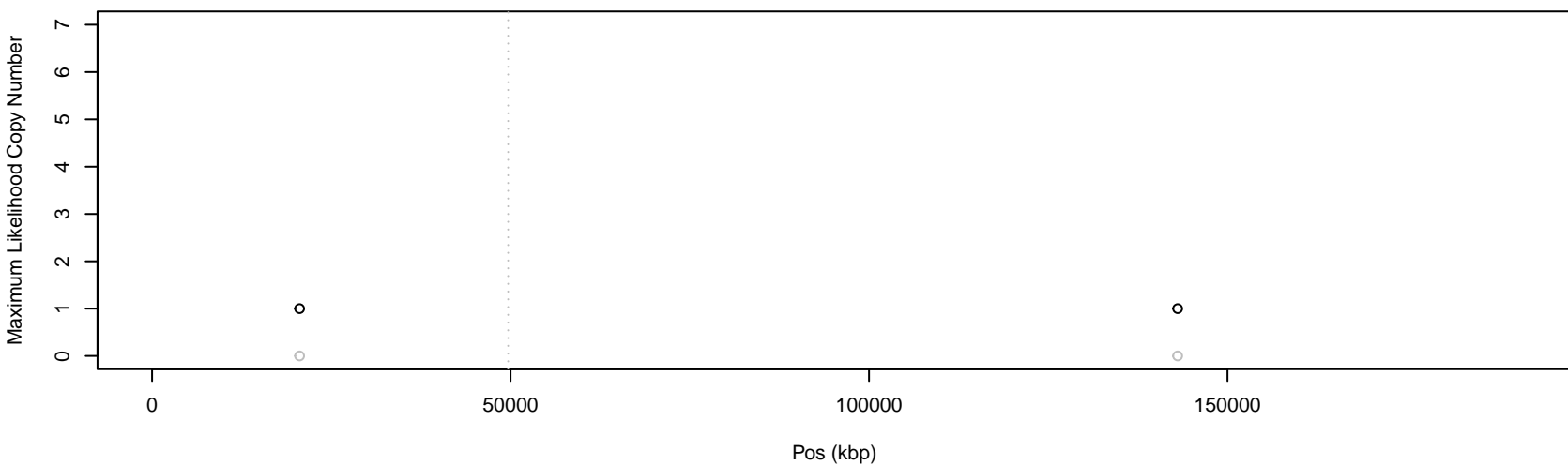
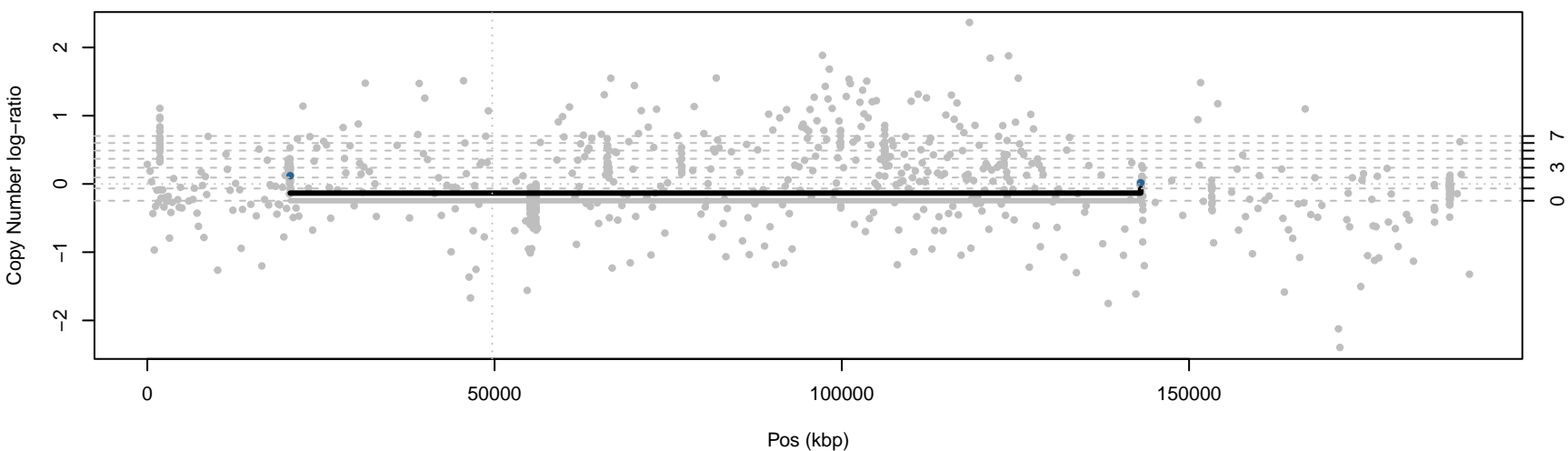
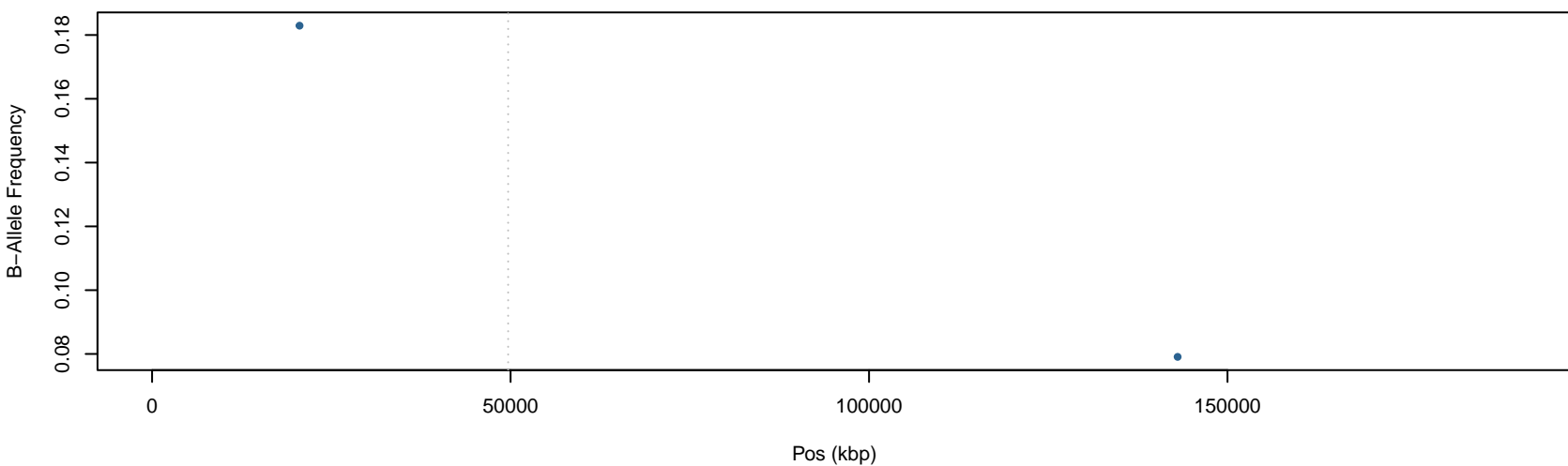
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 2



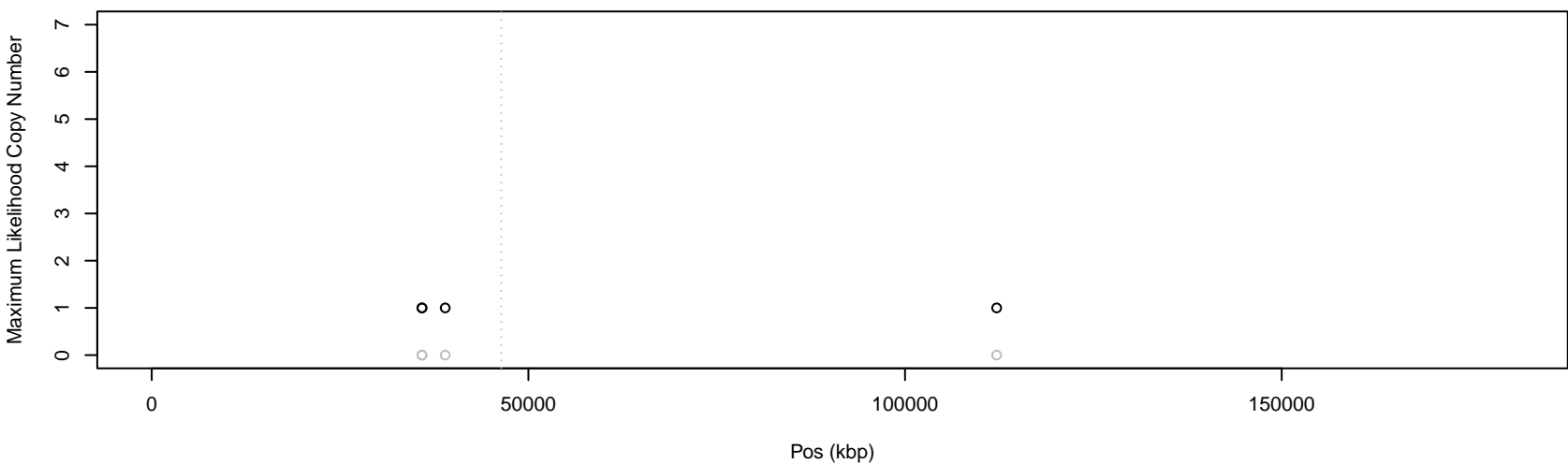
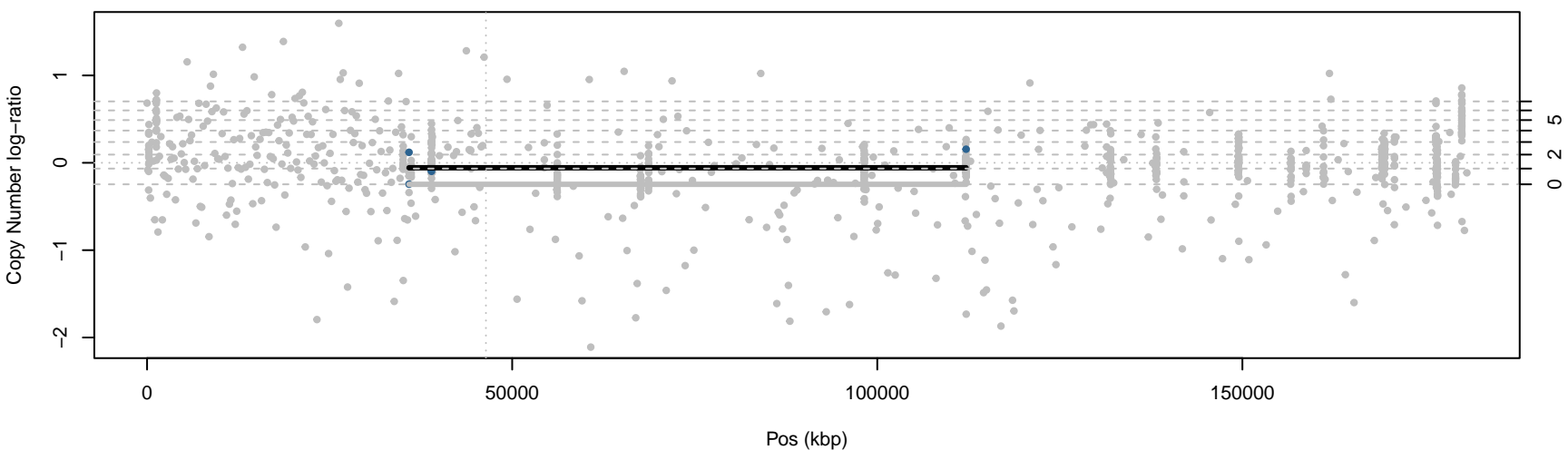
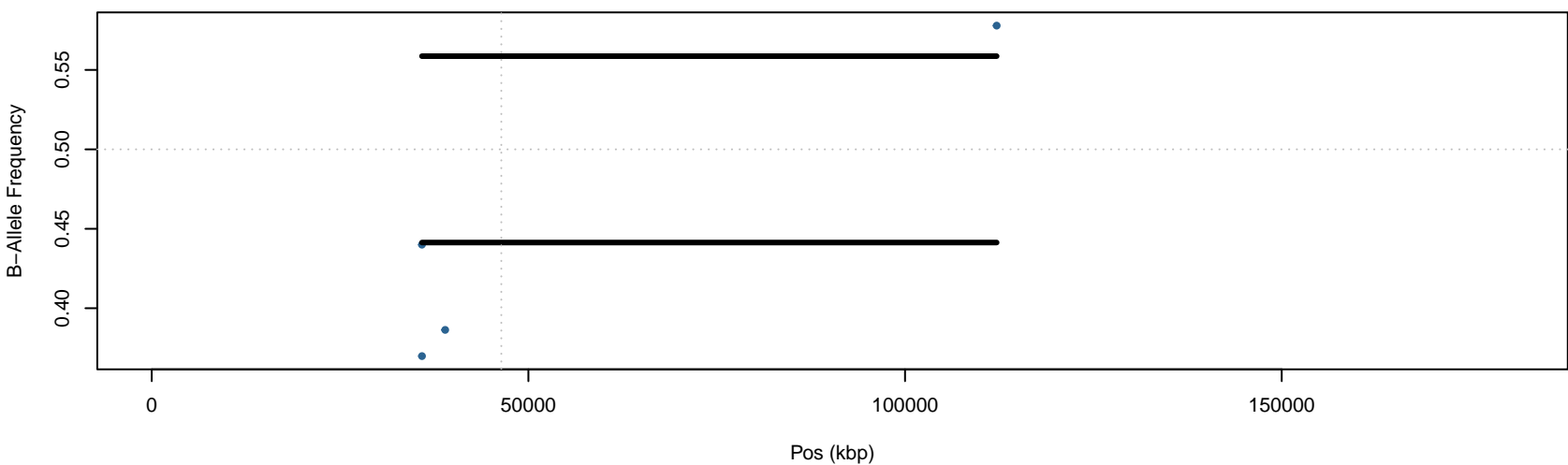
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 3



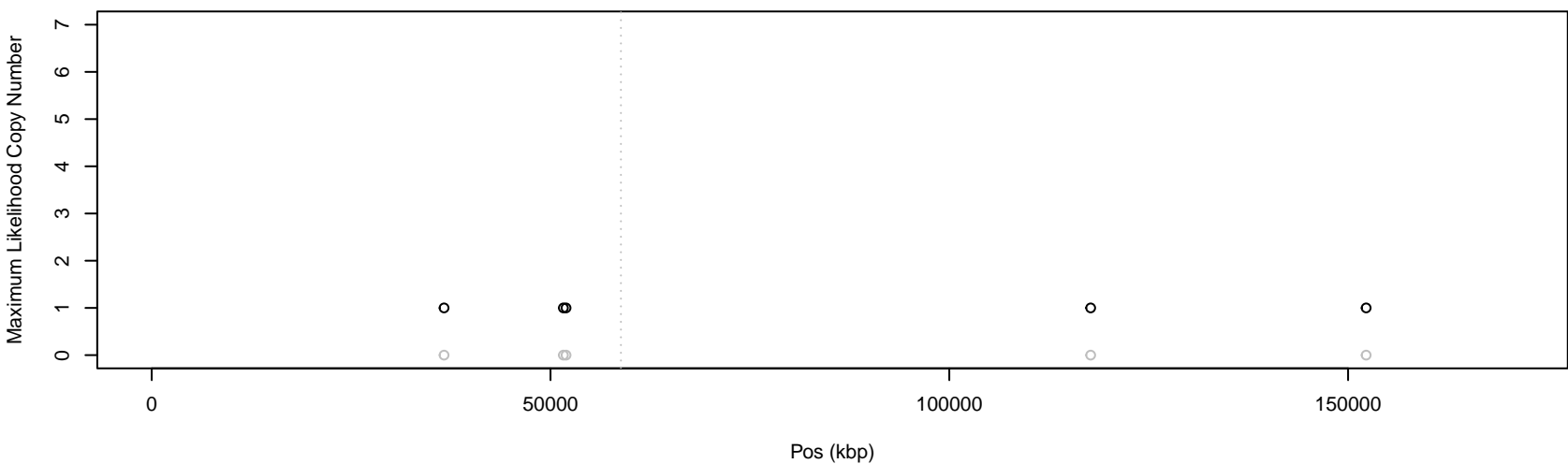
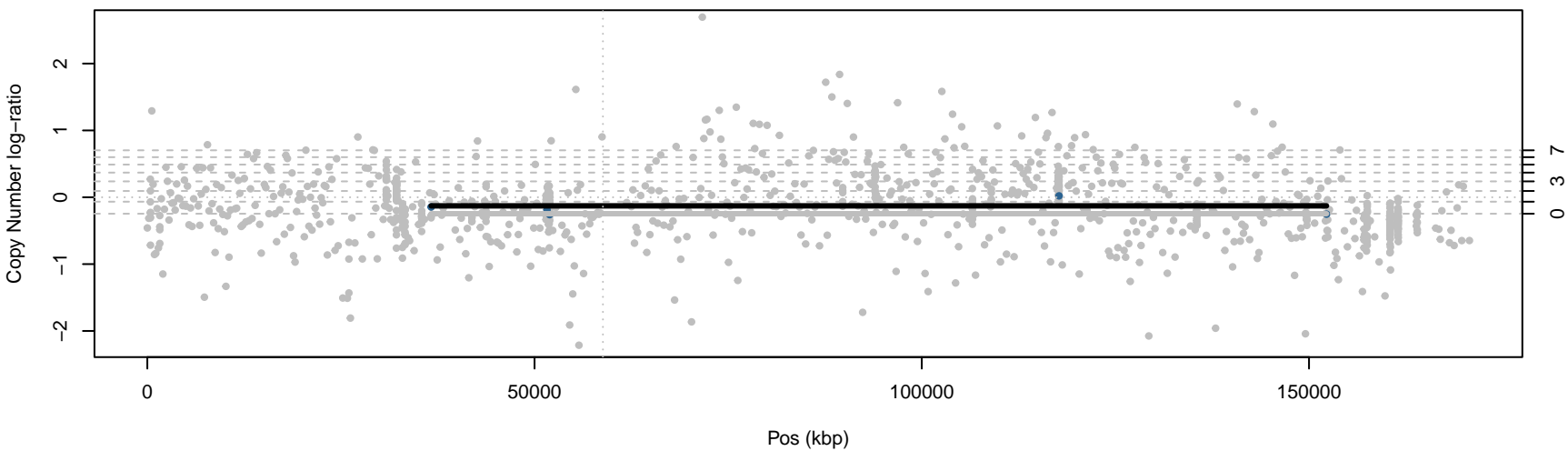
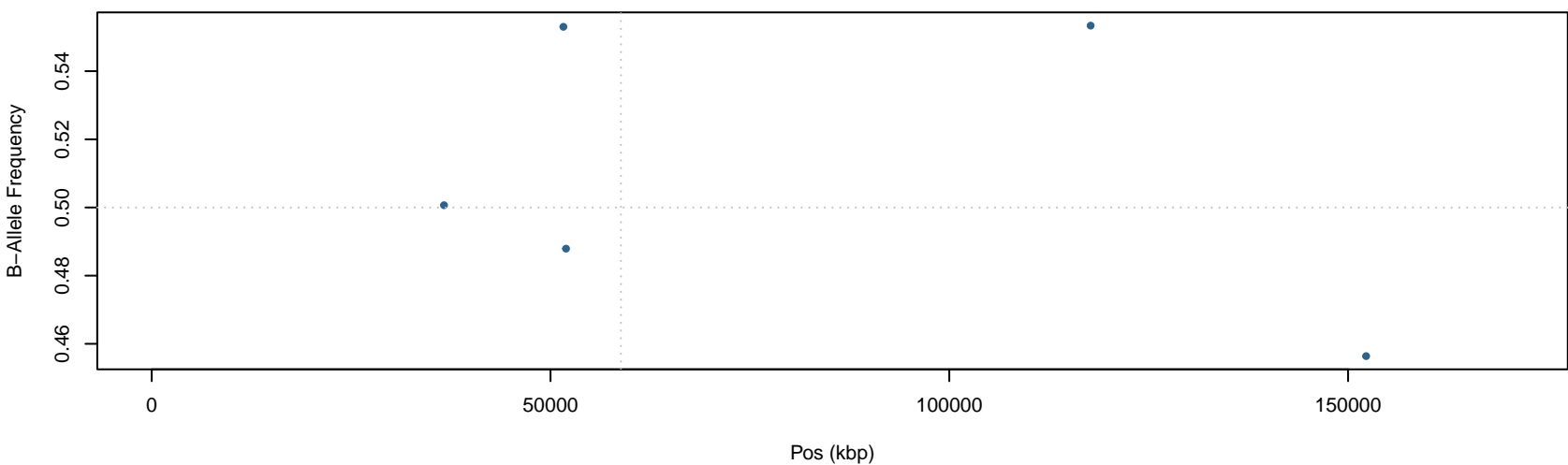
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 4



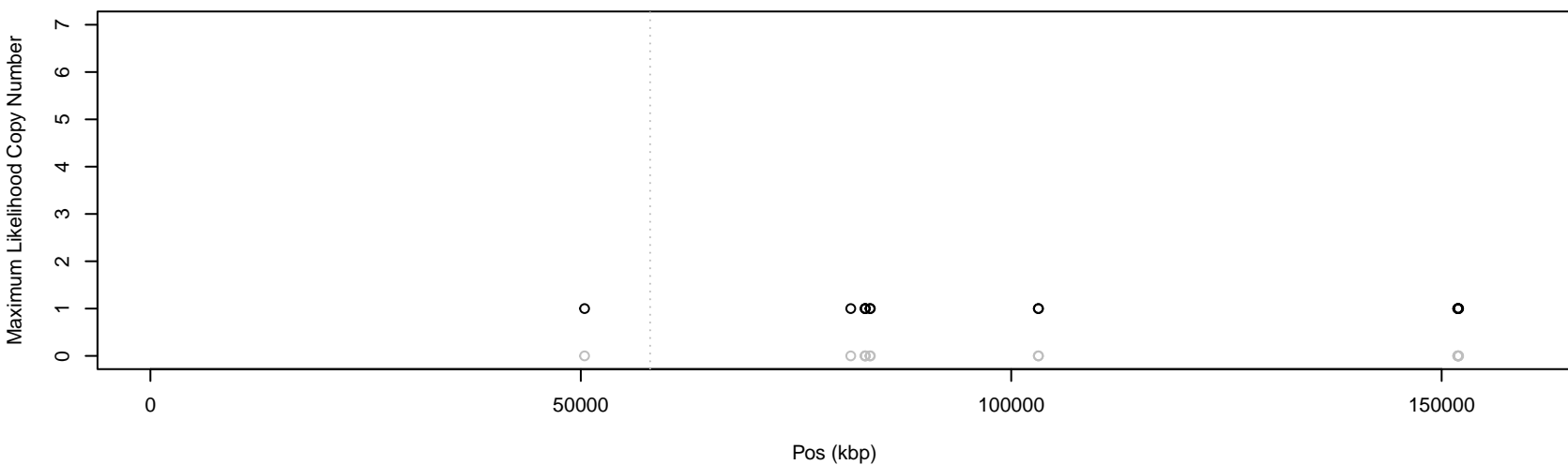
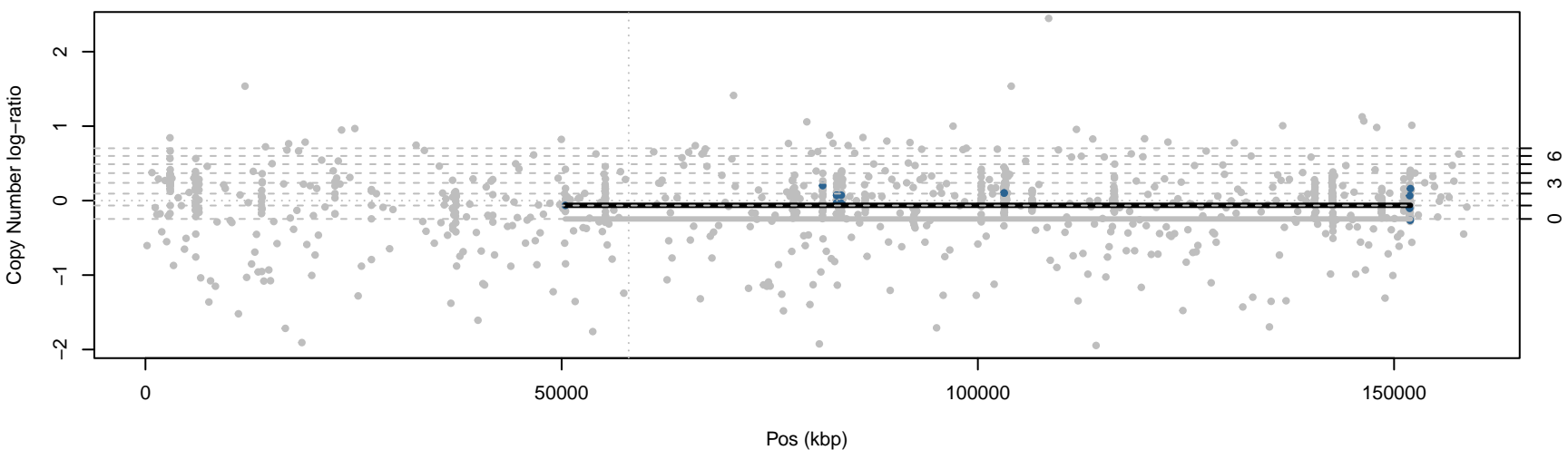
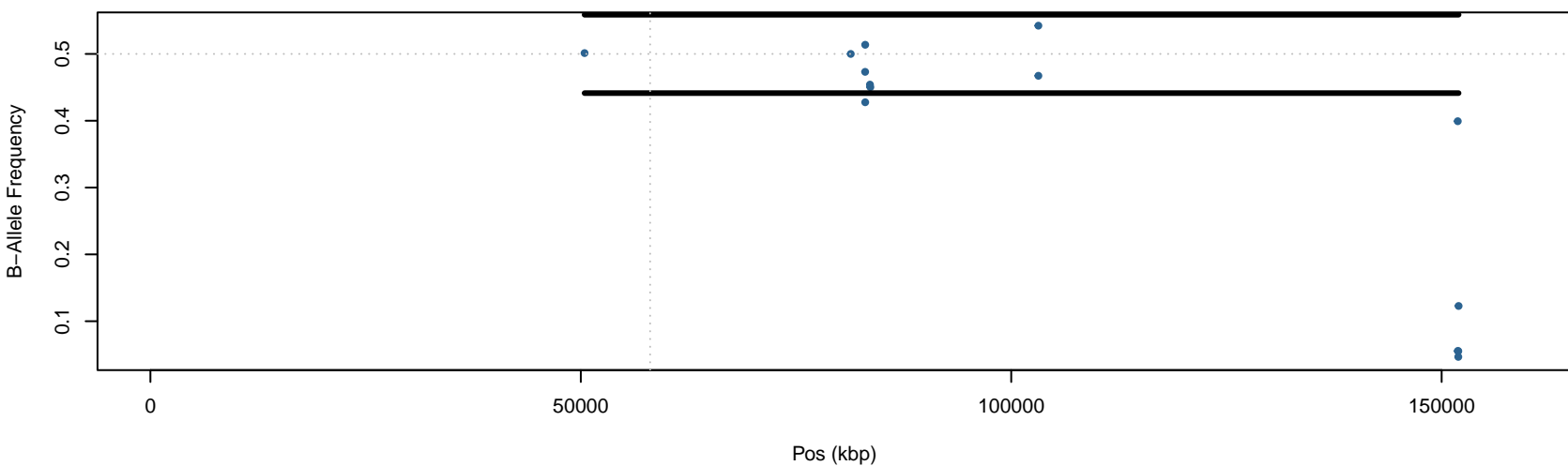
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 5



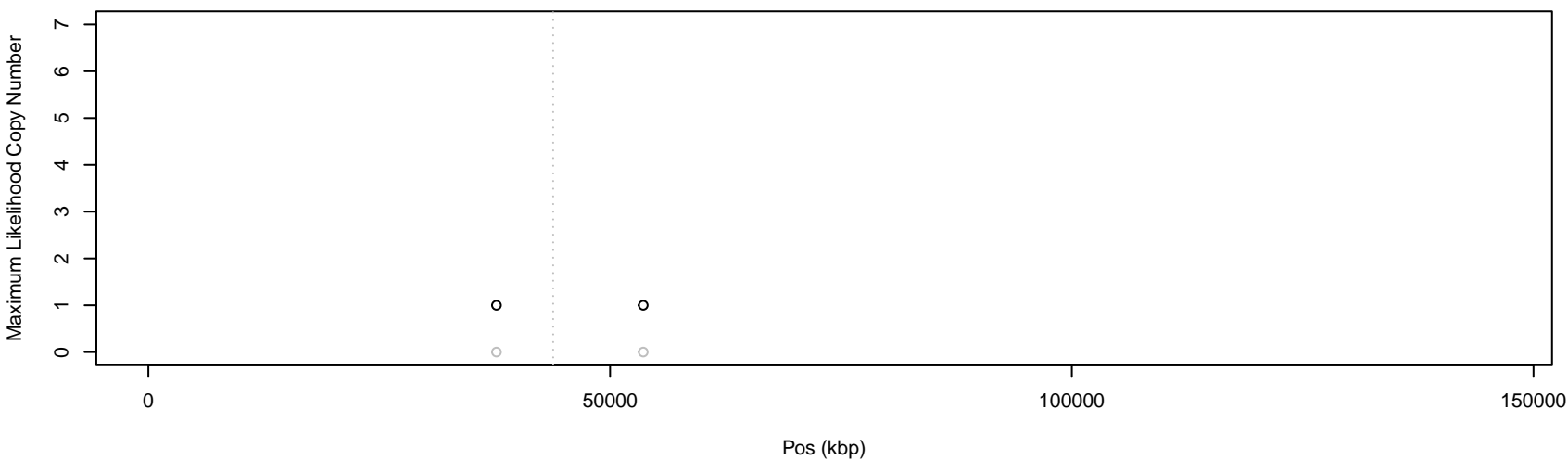
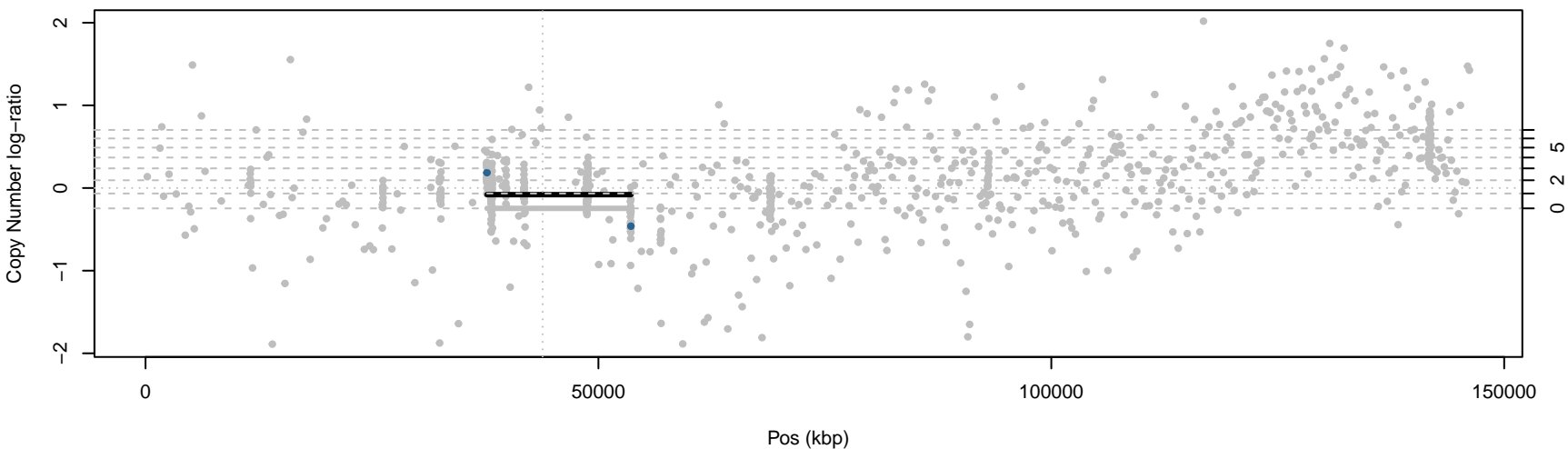
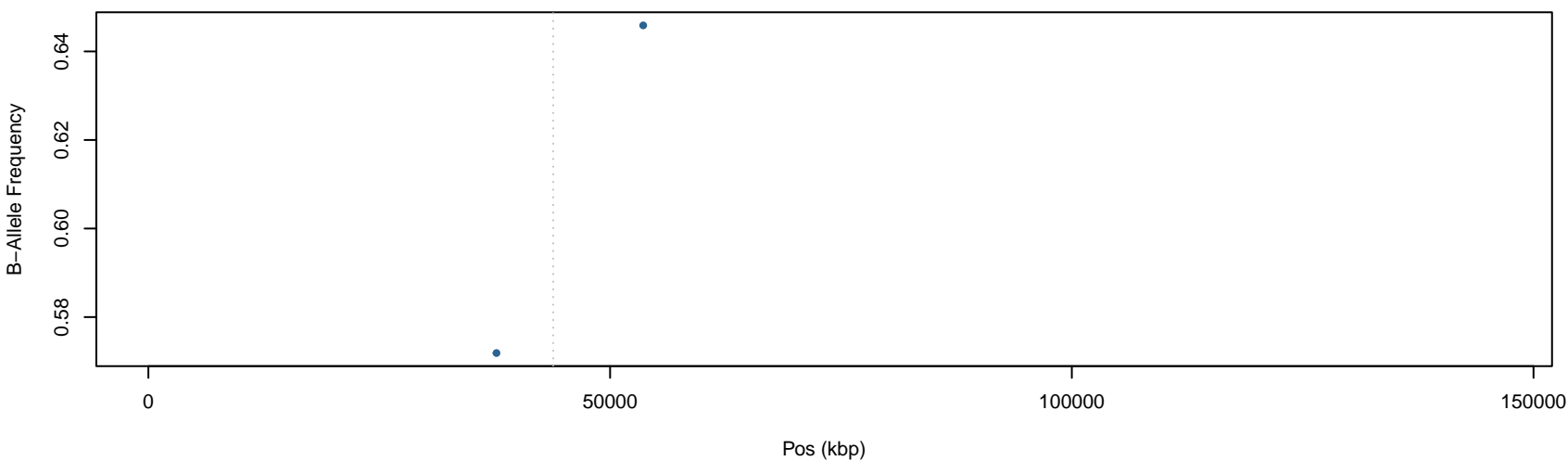
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 6



Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 7

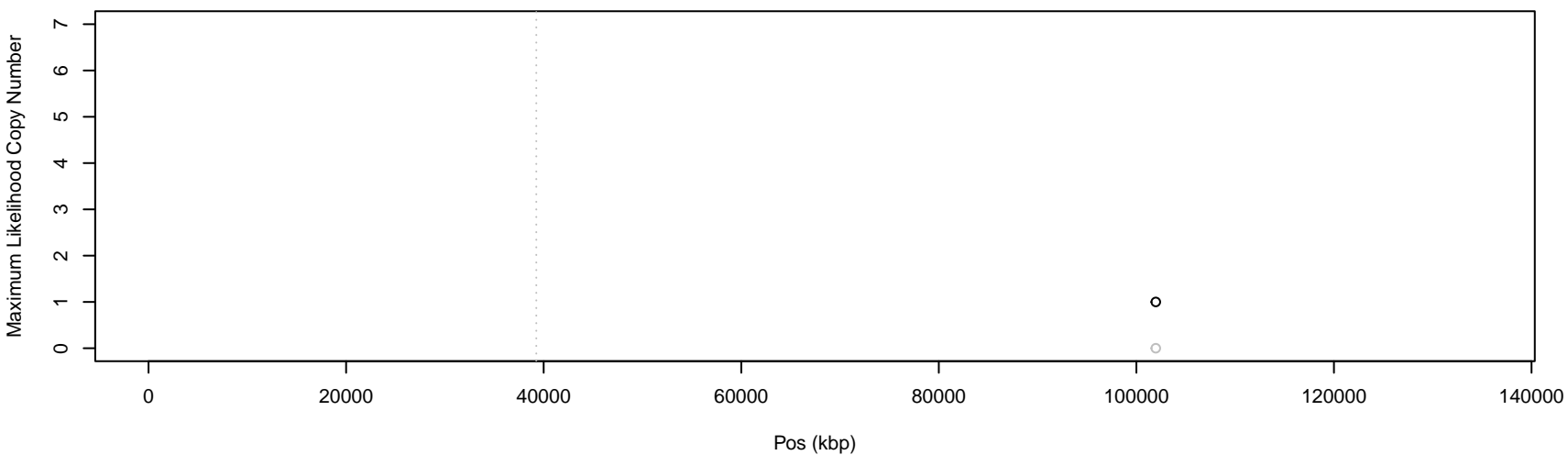
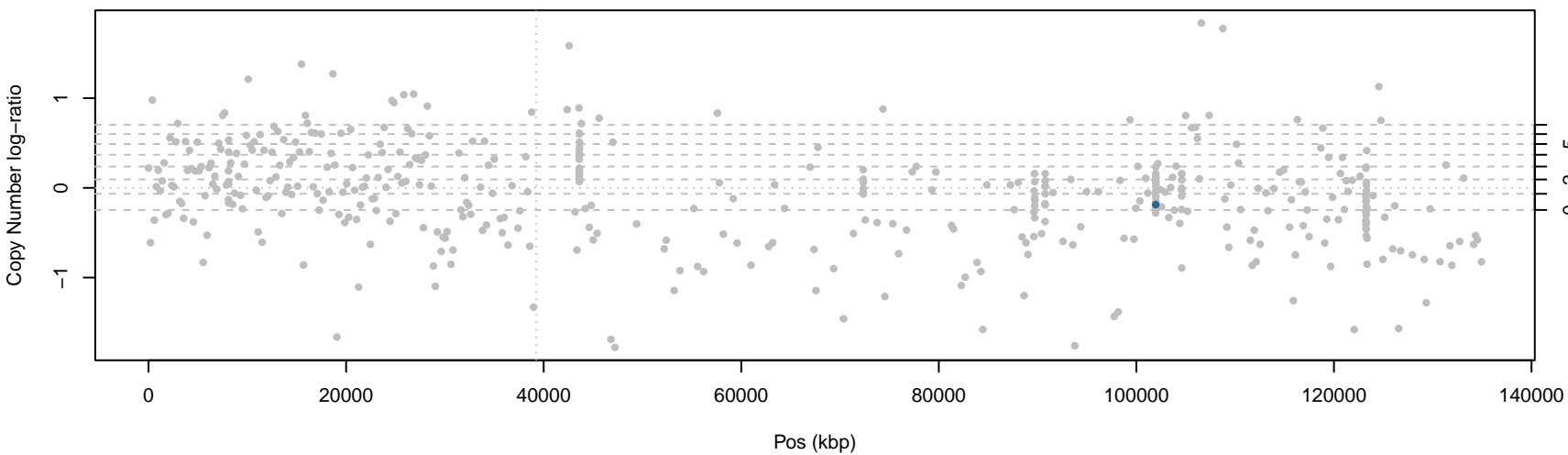
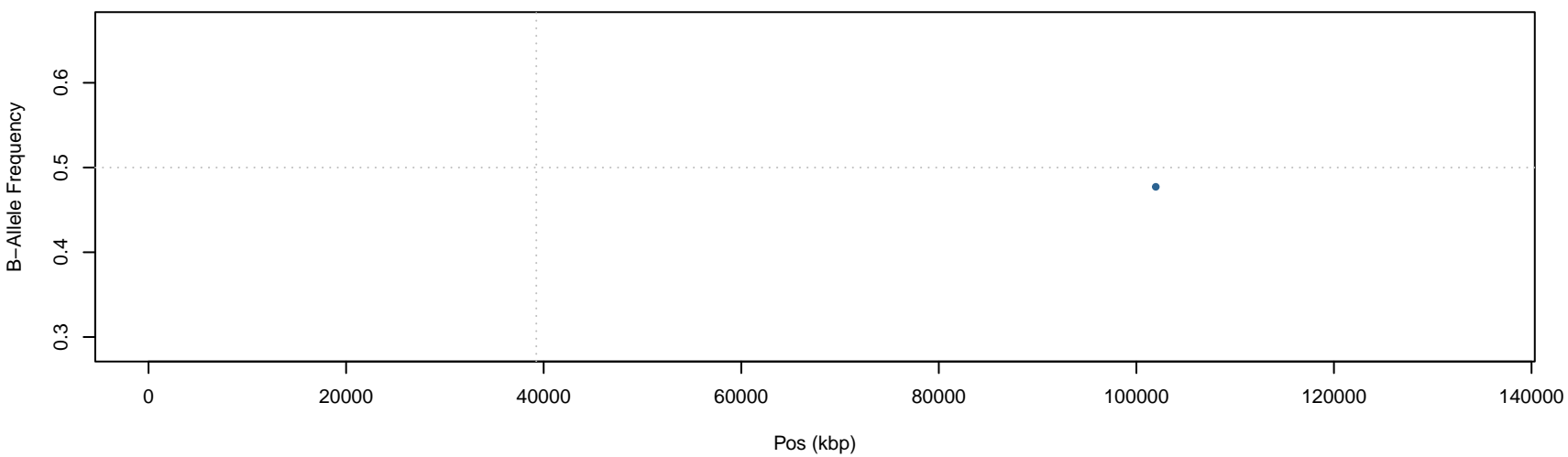


Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 8

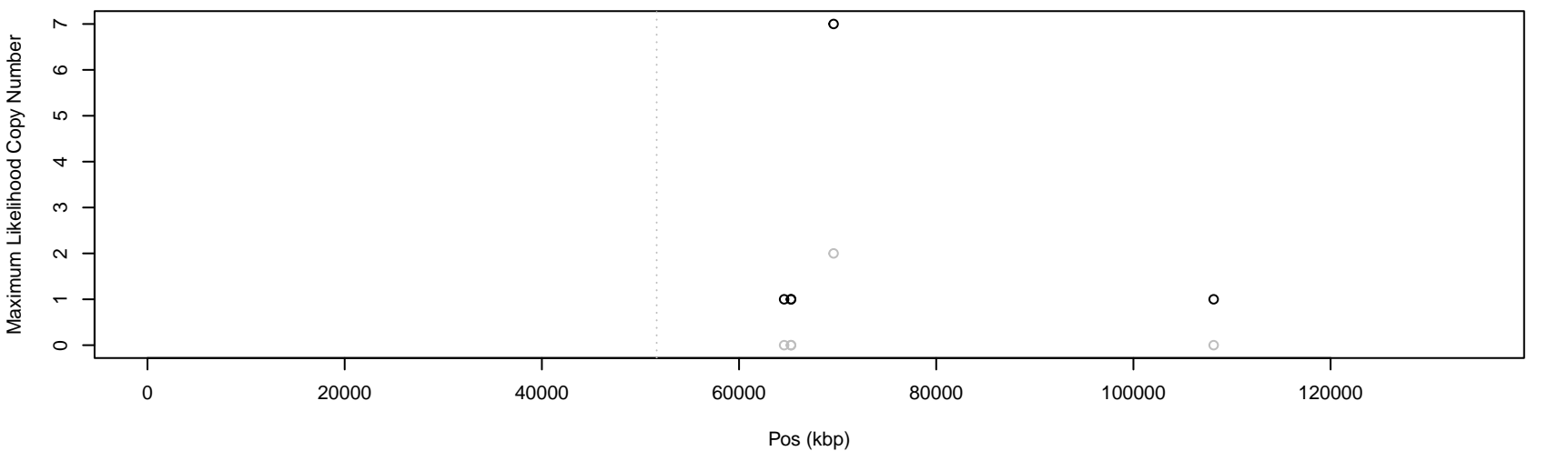
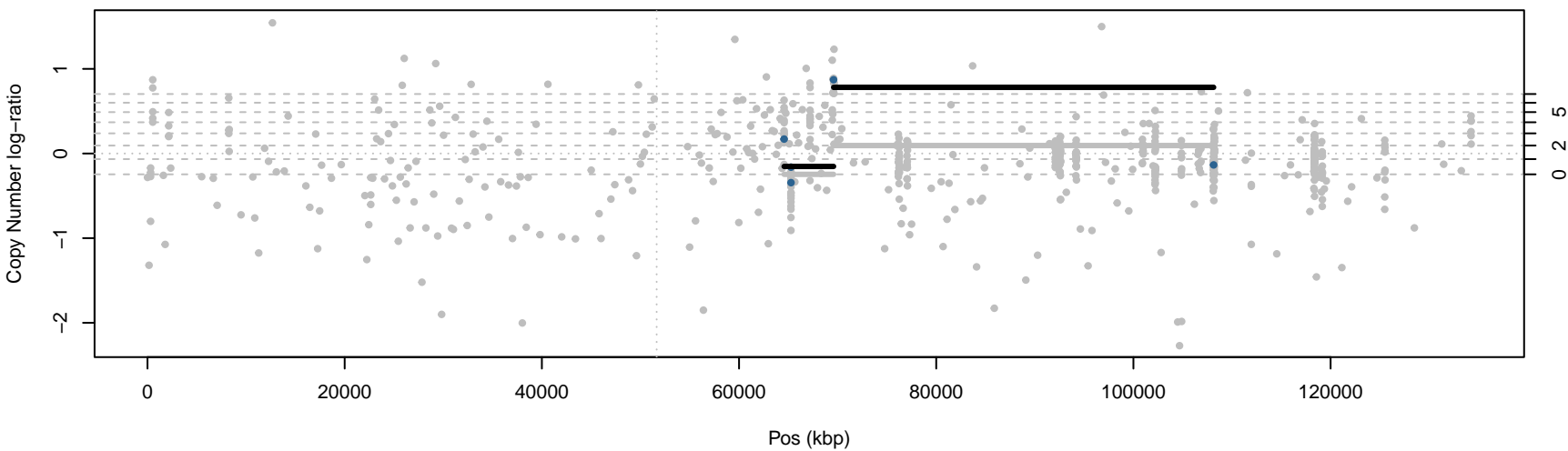
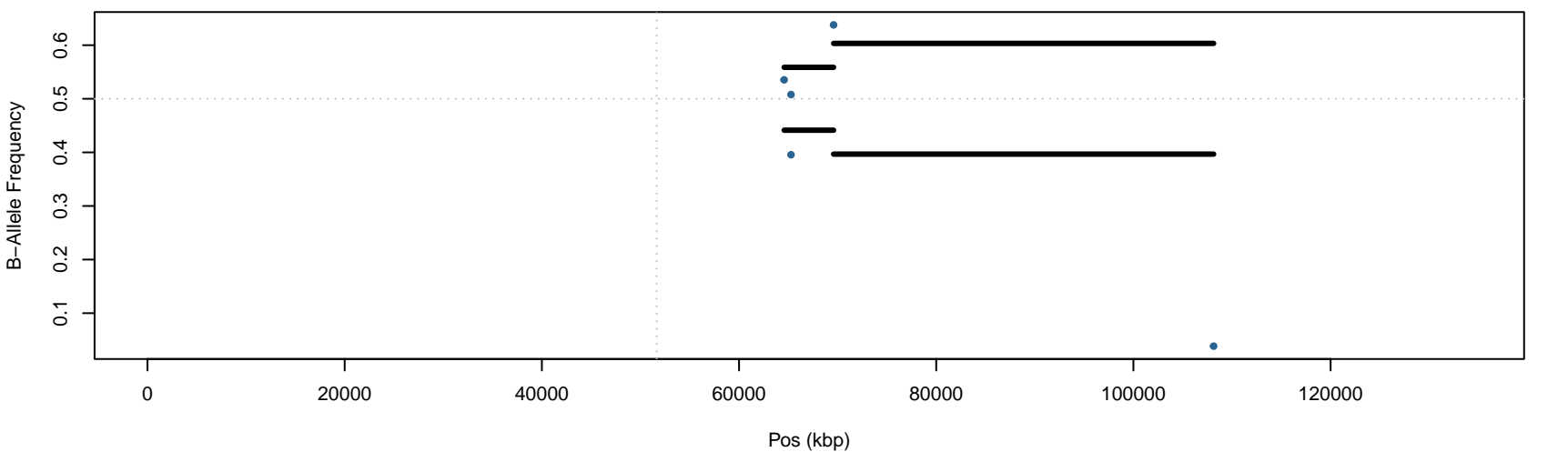




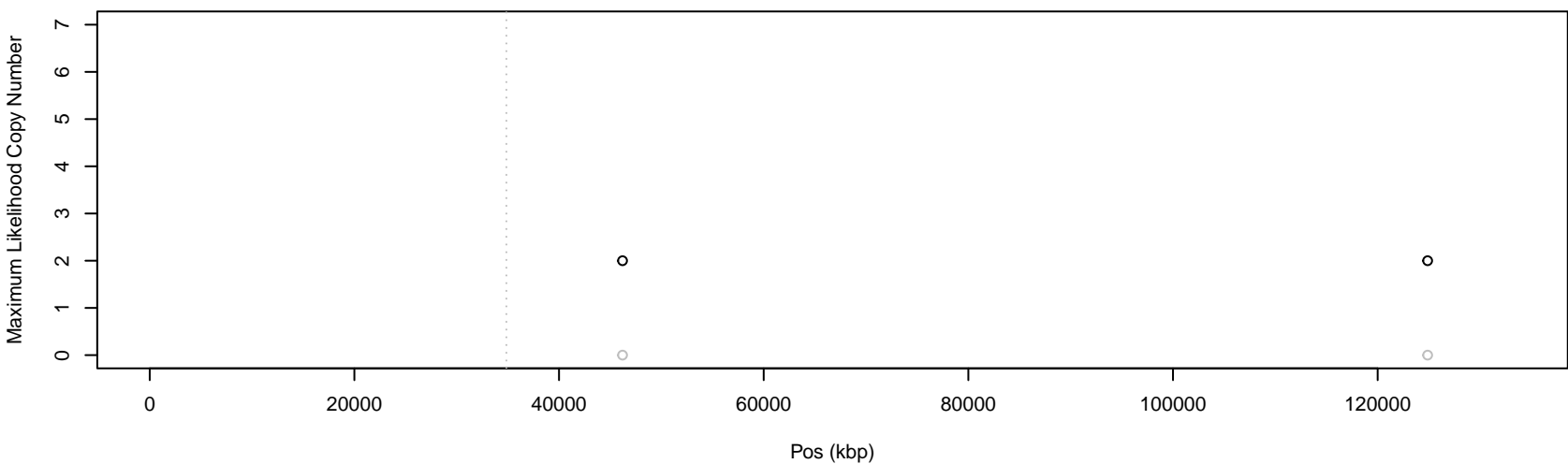
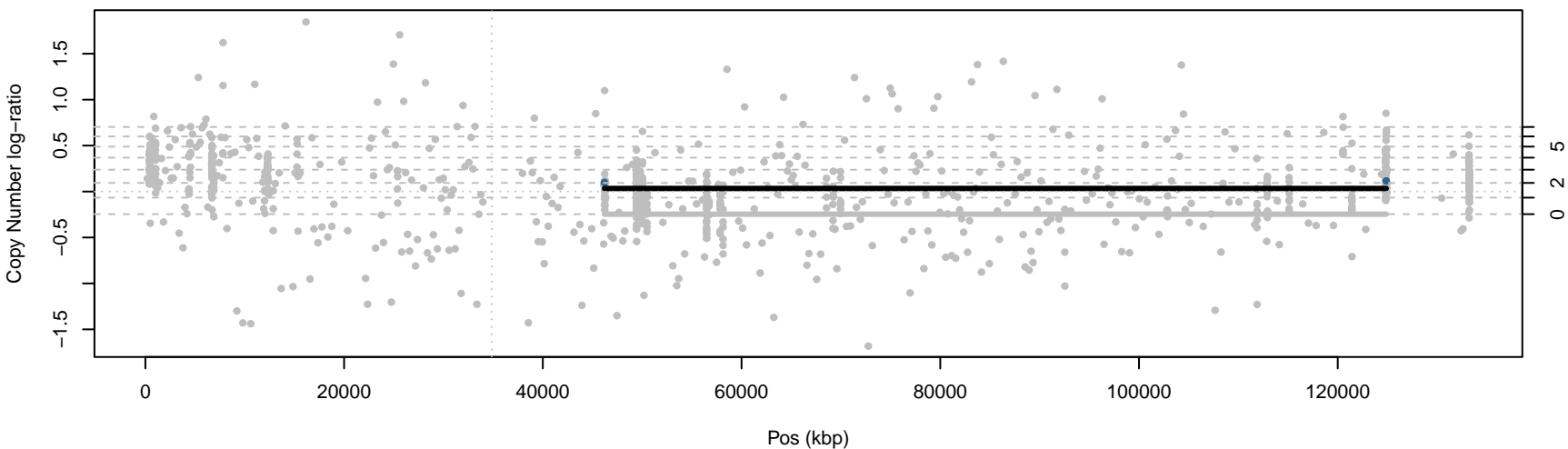
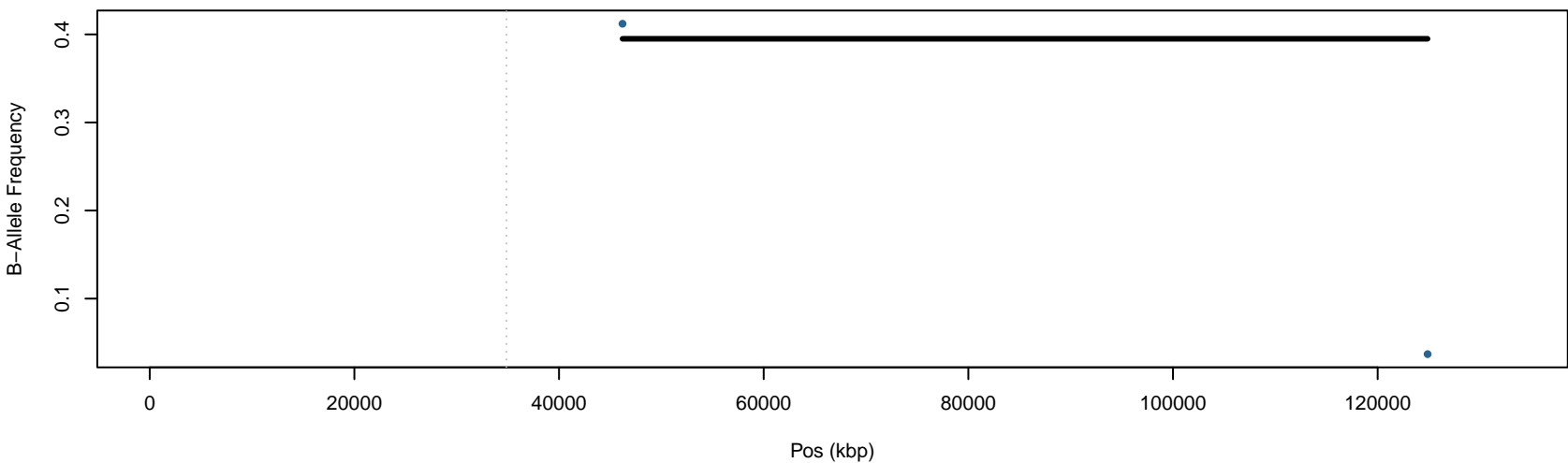
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 10



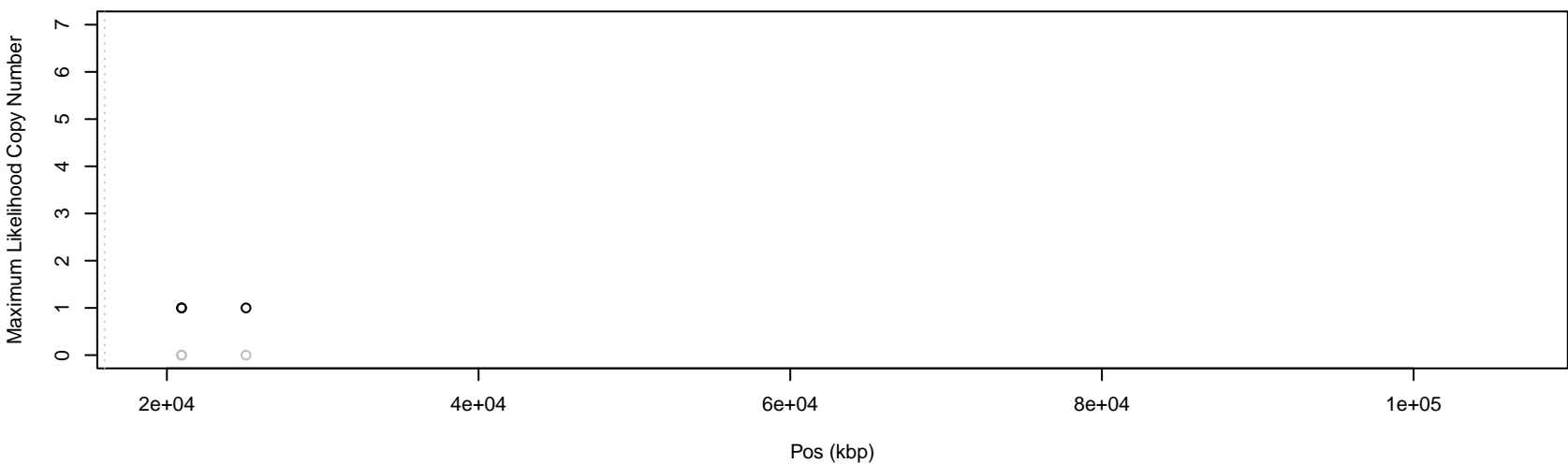
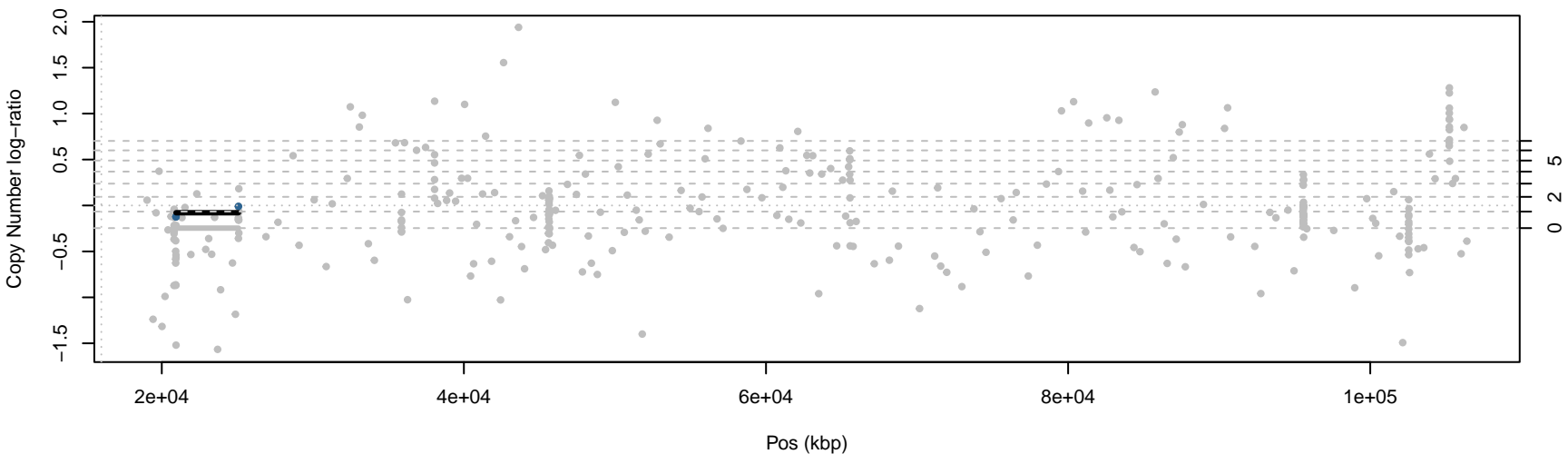
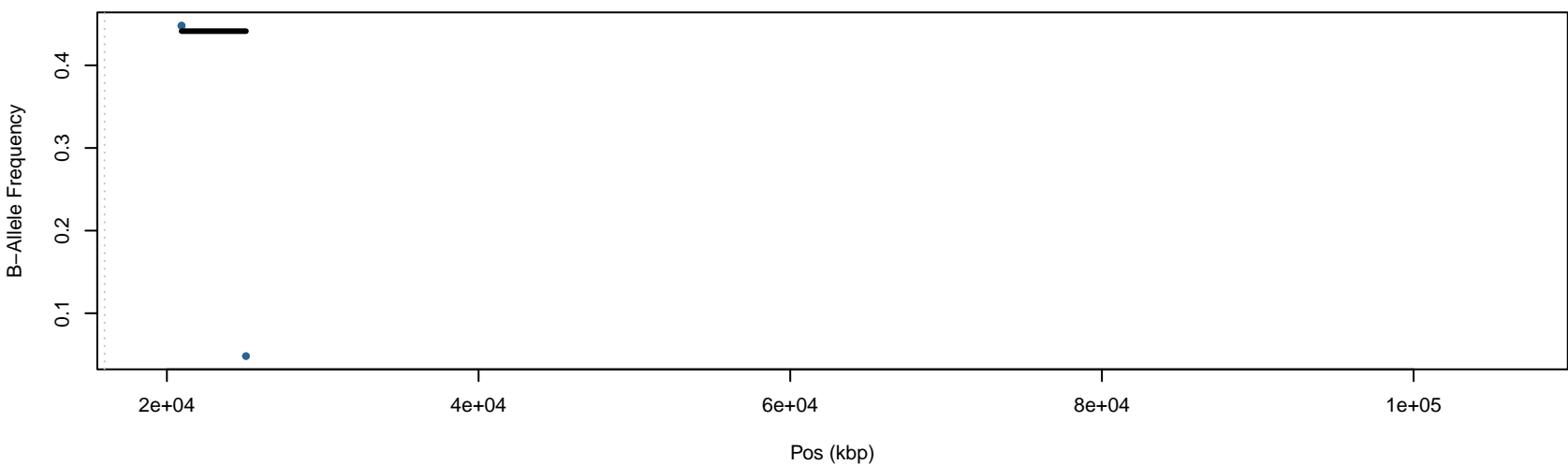
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 11



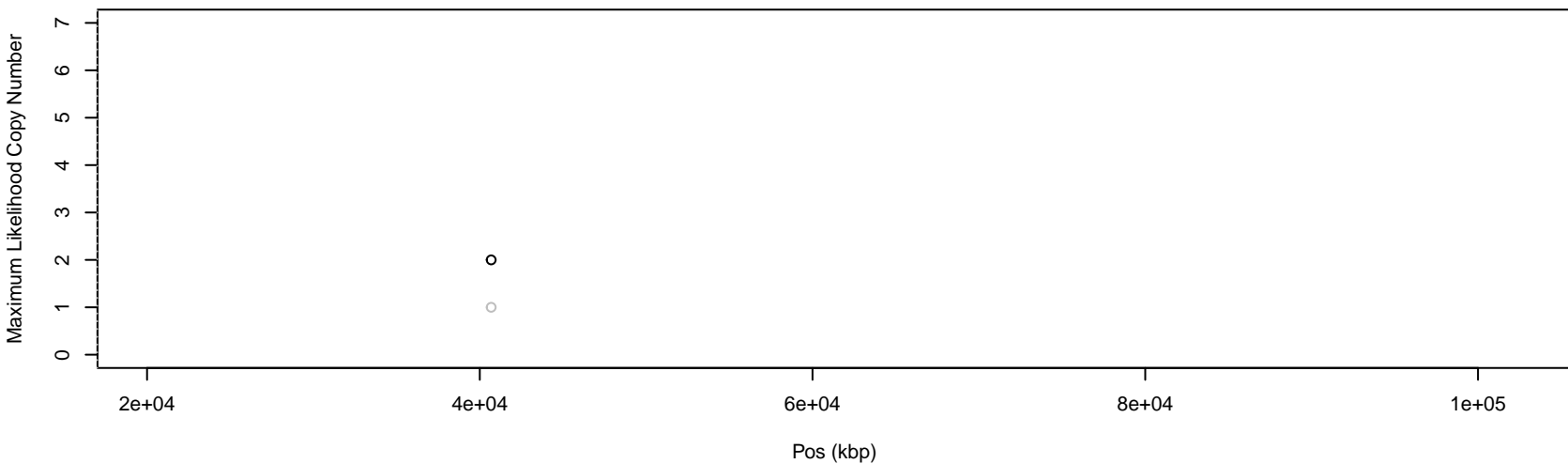
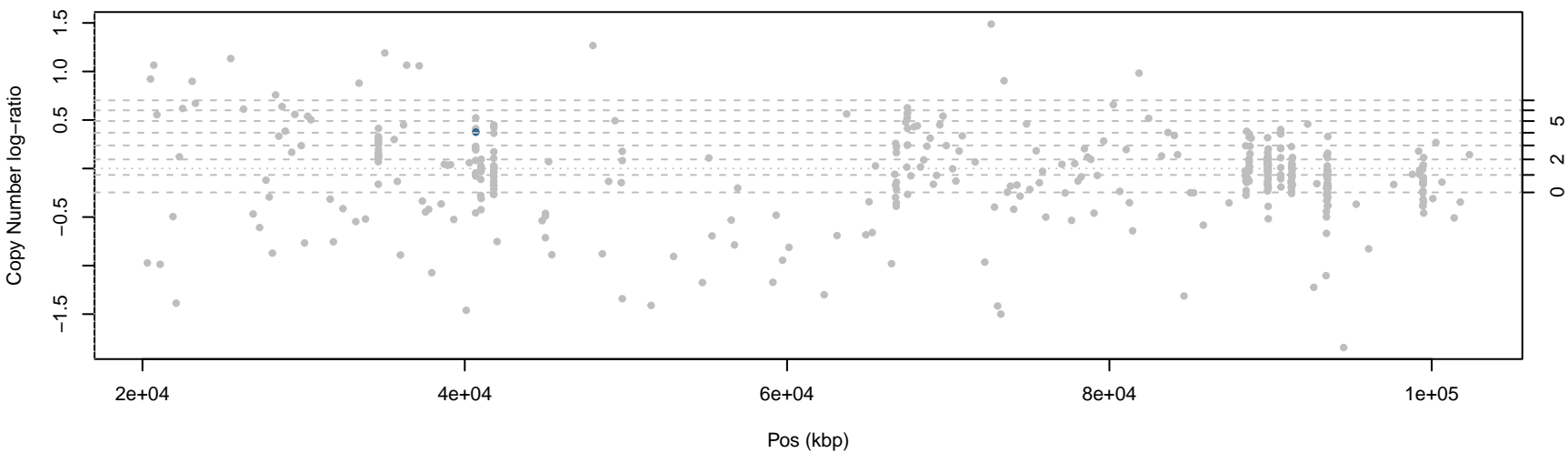
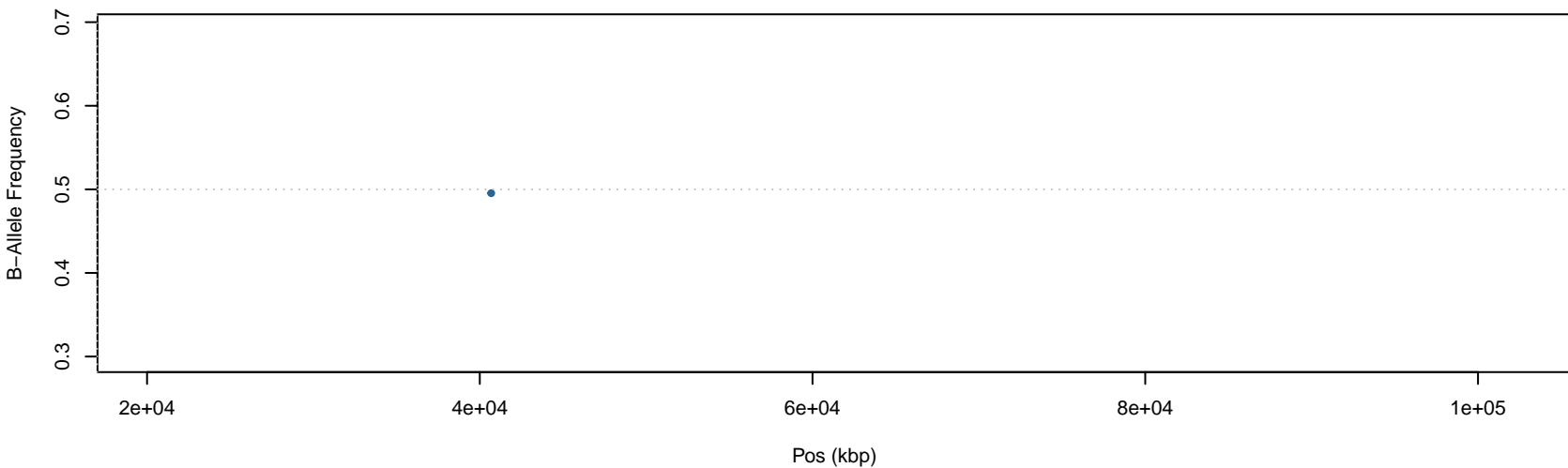
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 12



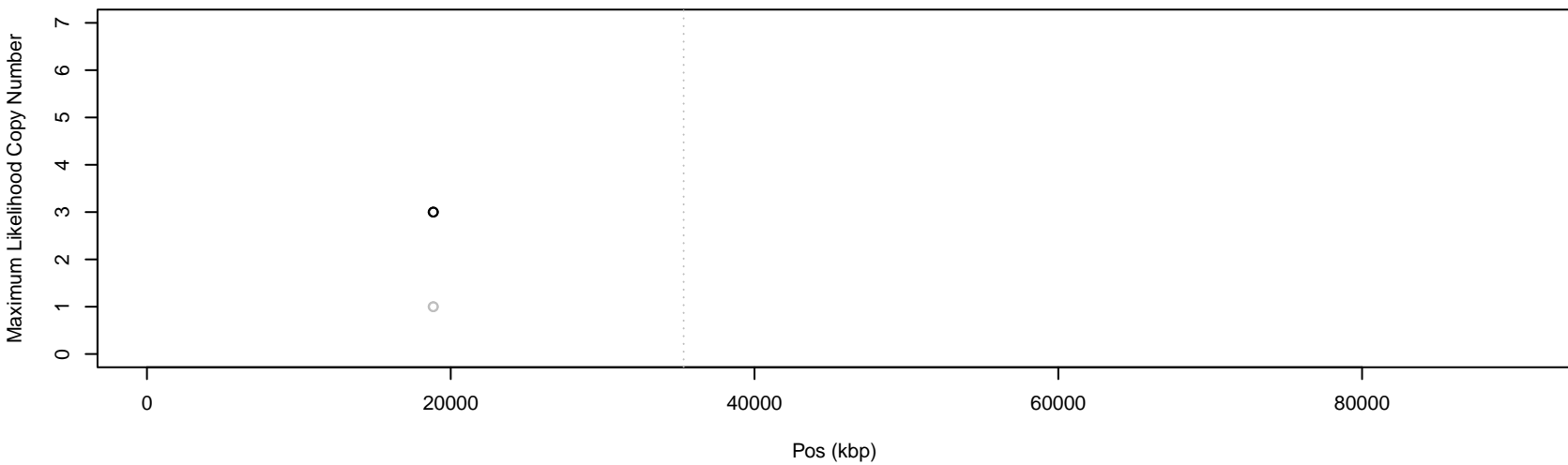
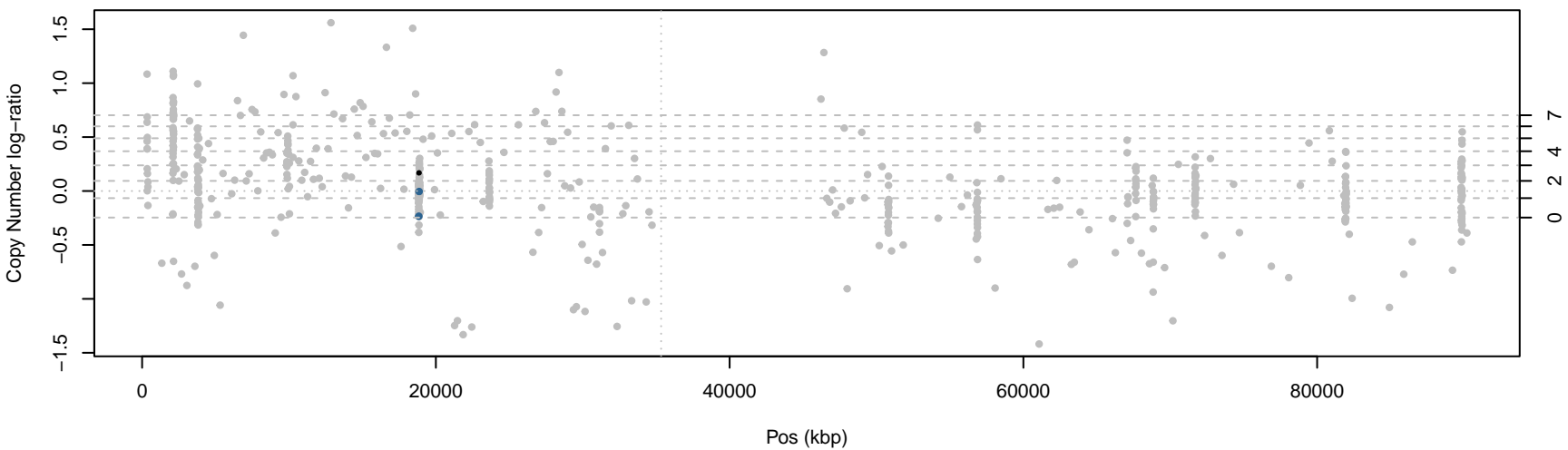
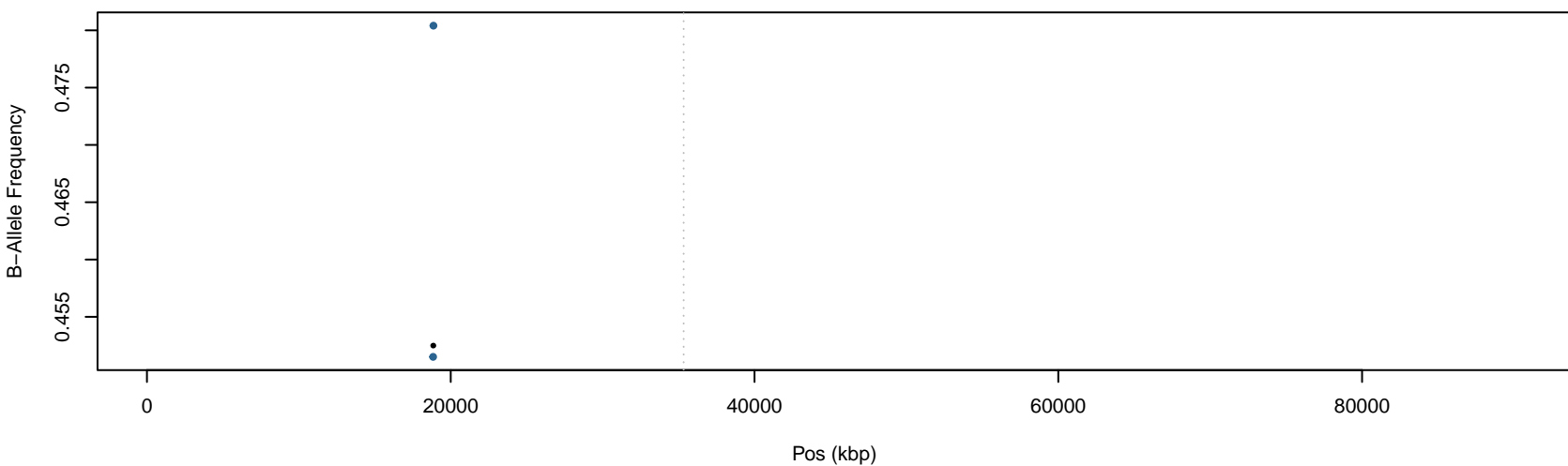
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 14



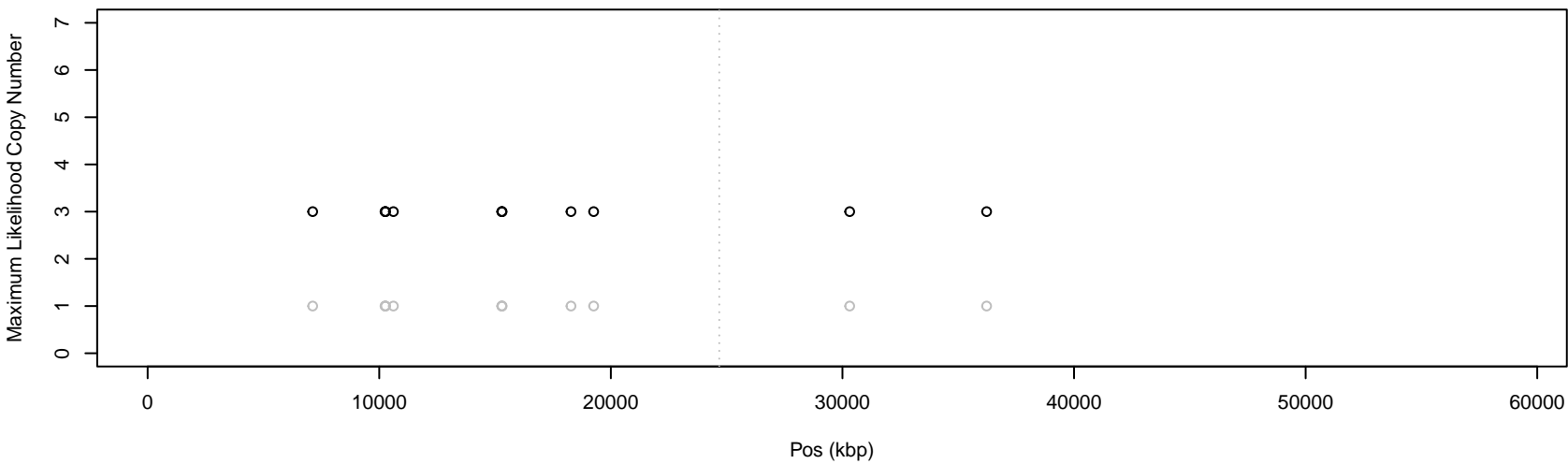
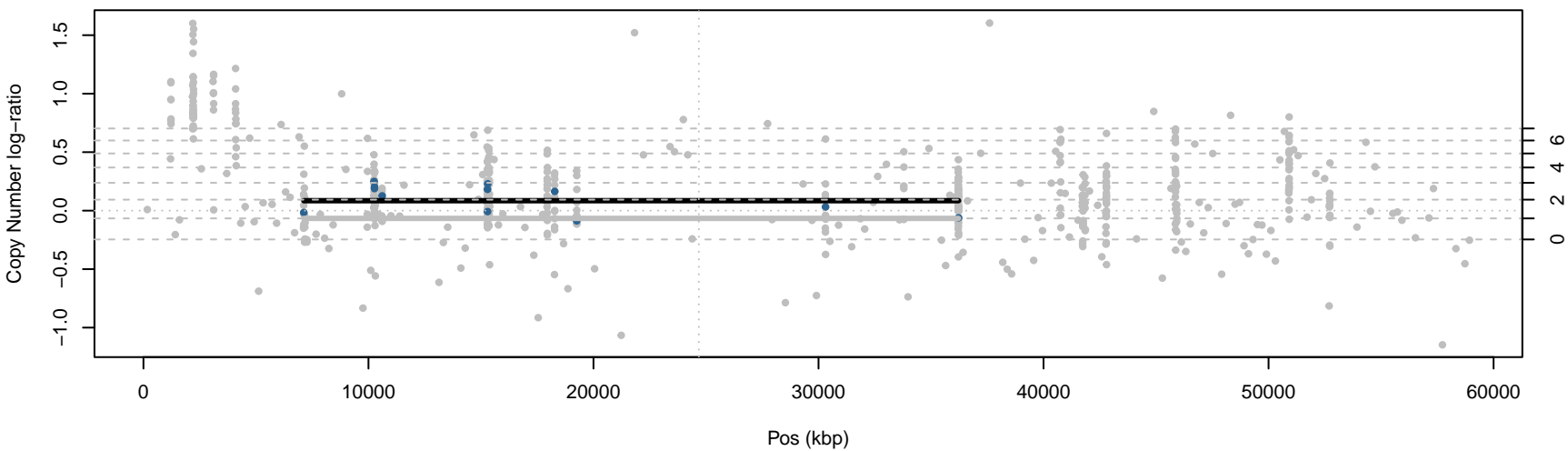
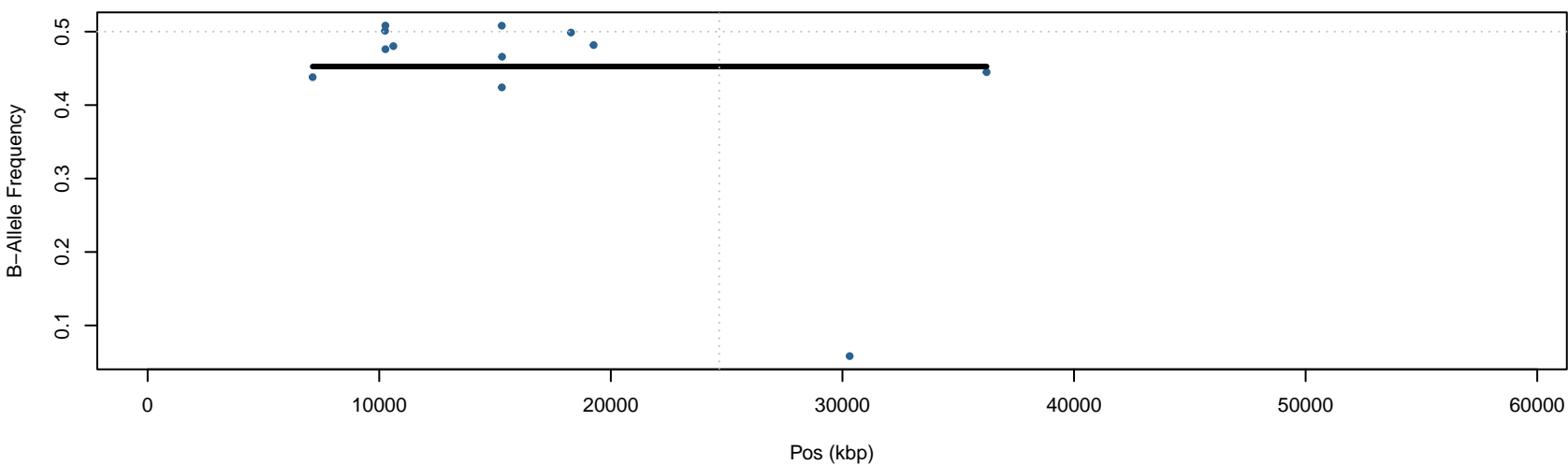
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 15



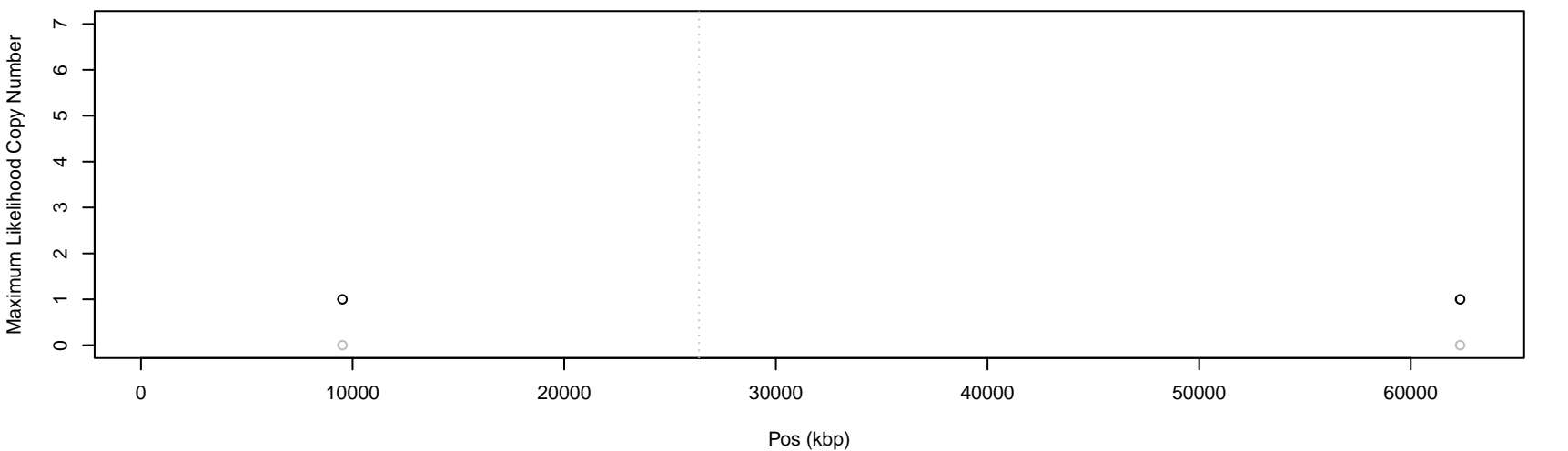
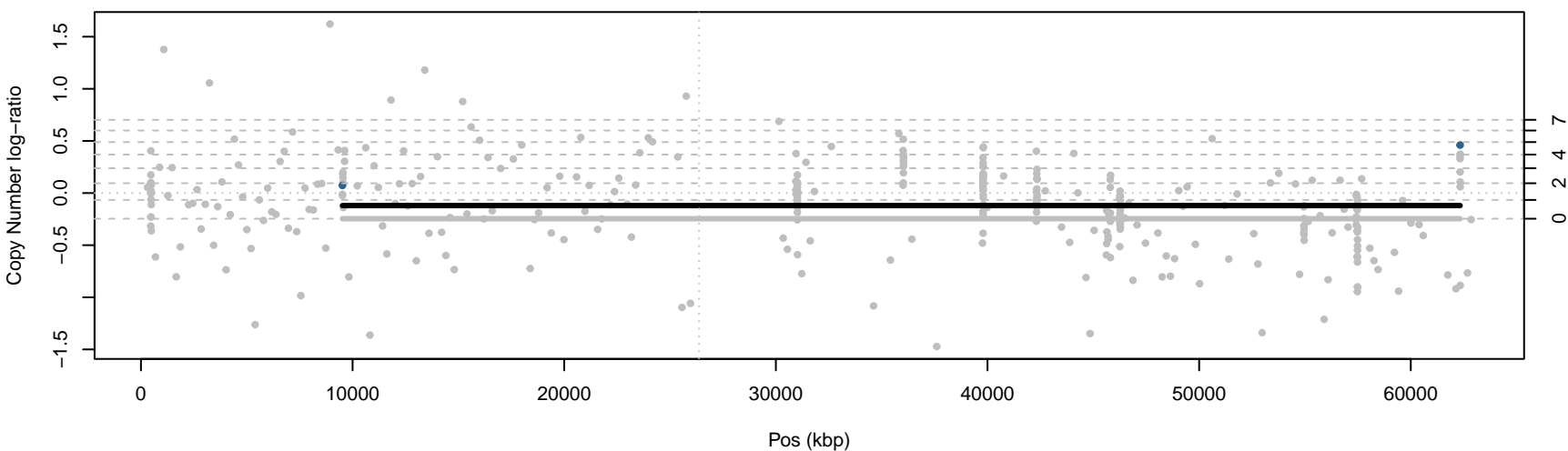
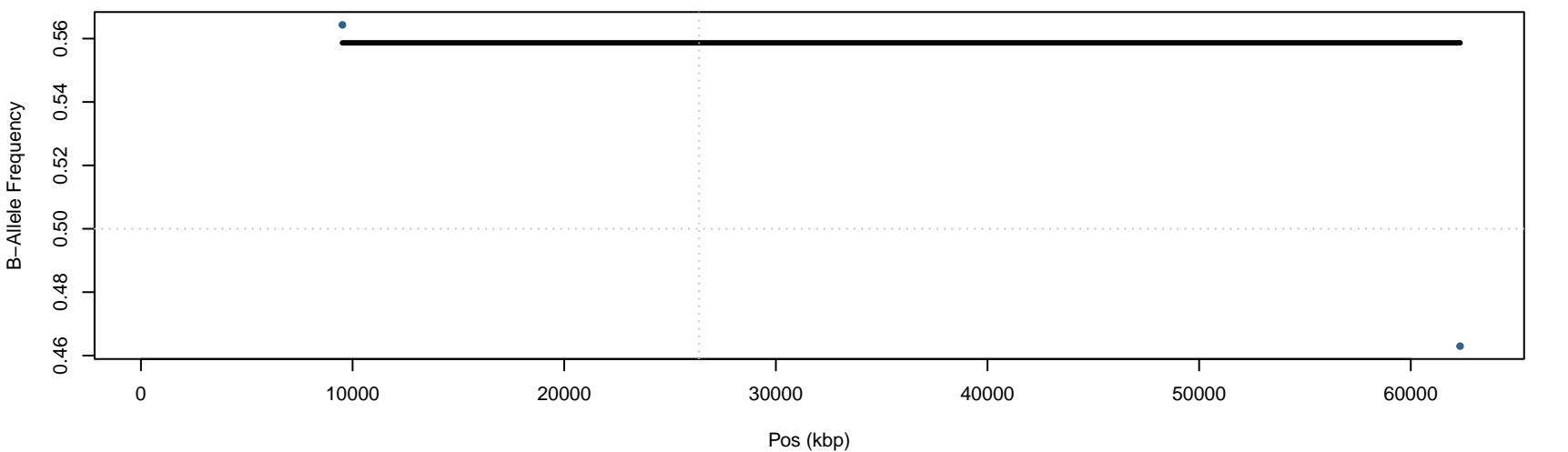
Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 16



Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 19

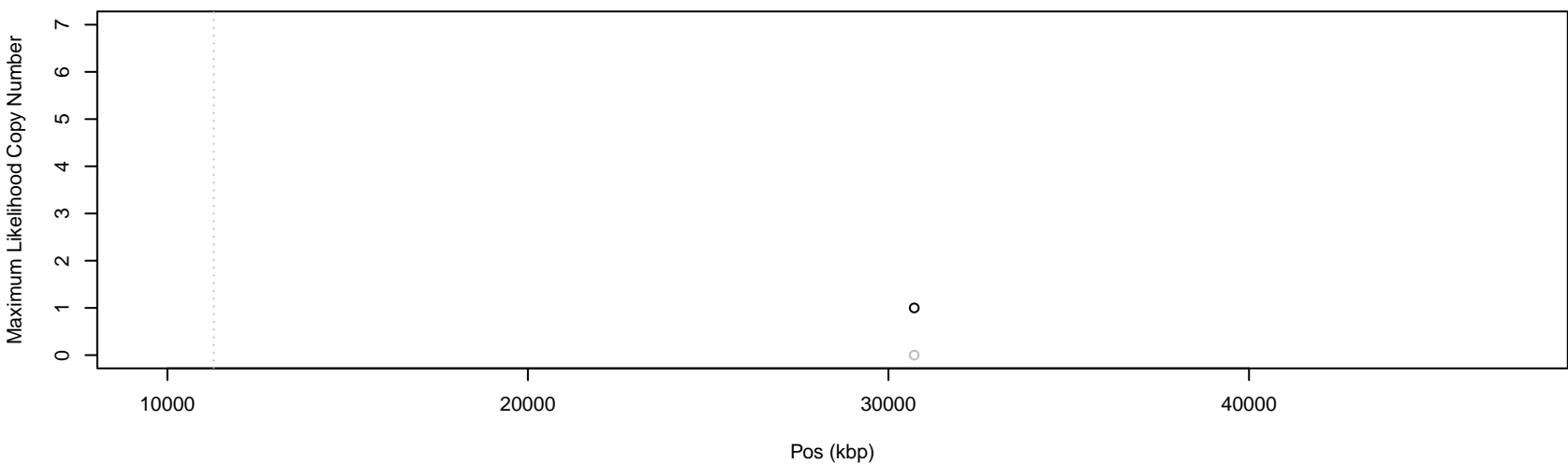
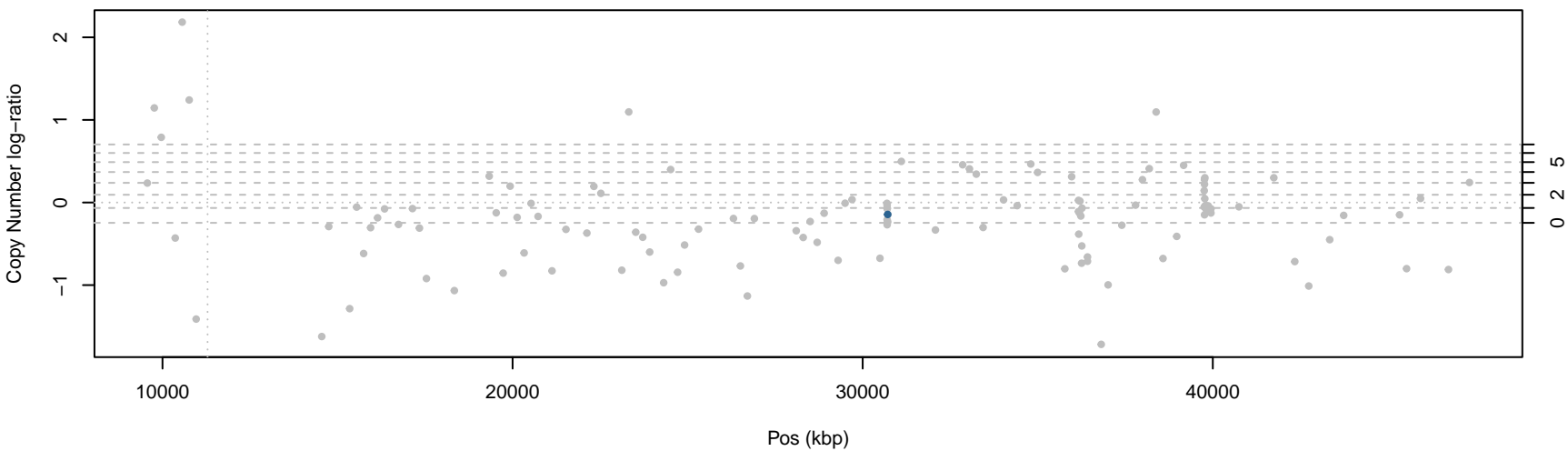
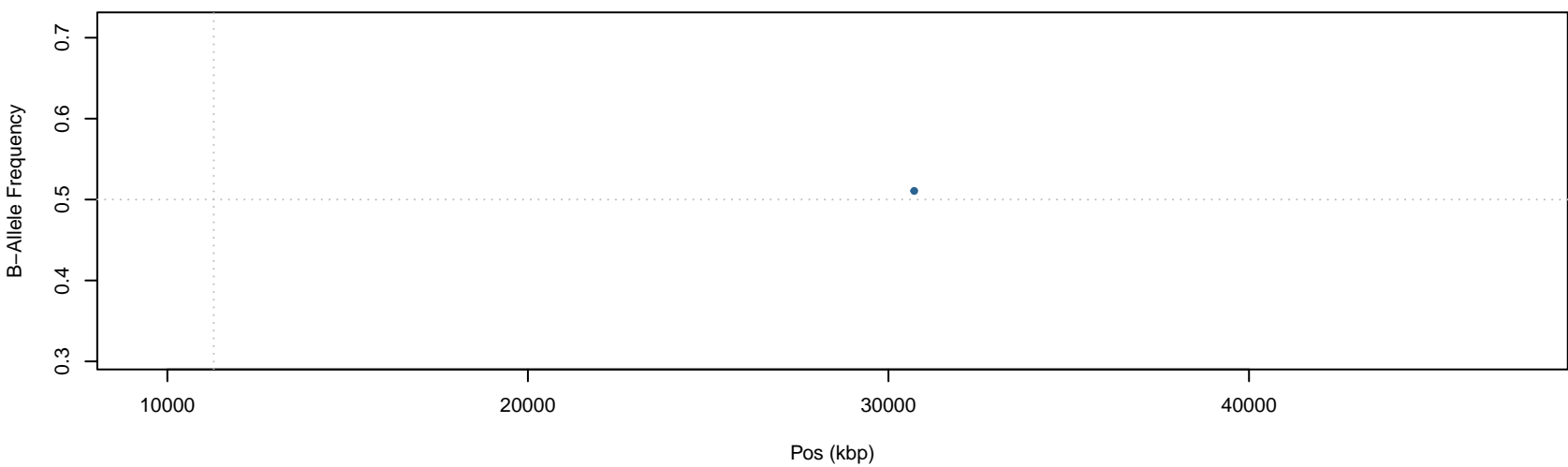


Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 20





Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 21



Purity: 0.21 Tumor ploidy: 1.4 SNV log-likelihood: -15.85 GoF: 93.4% Mean coverage: 520;712 Chromosome: 22

