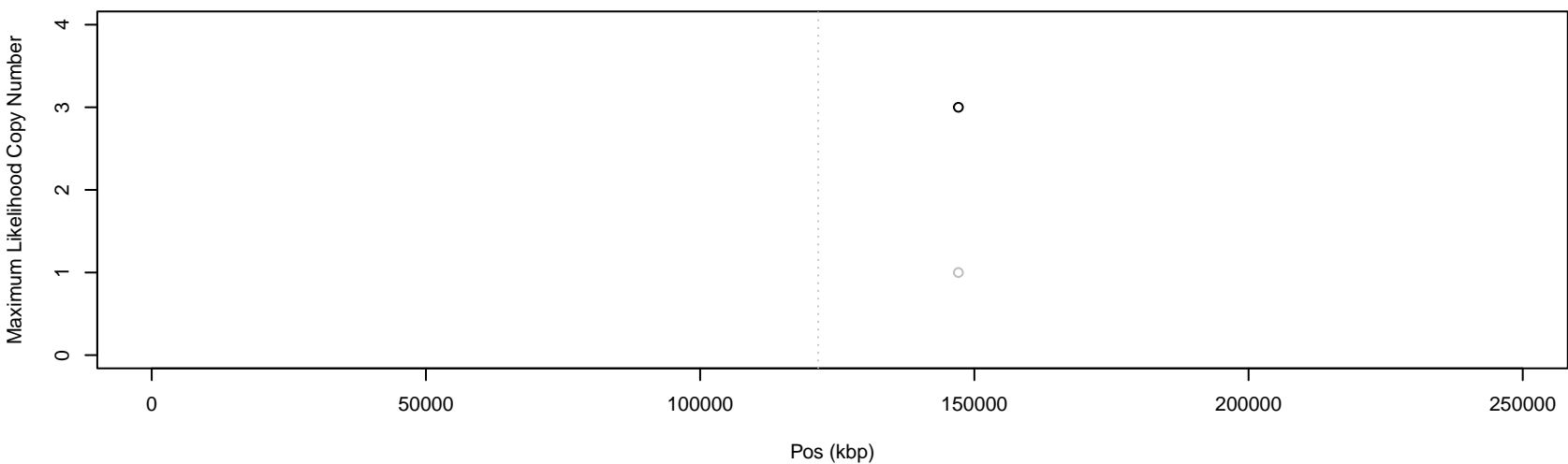
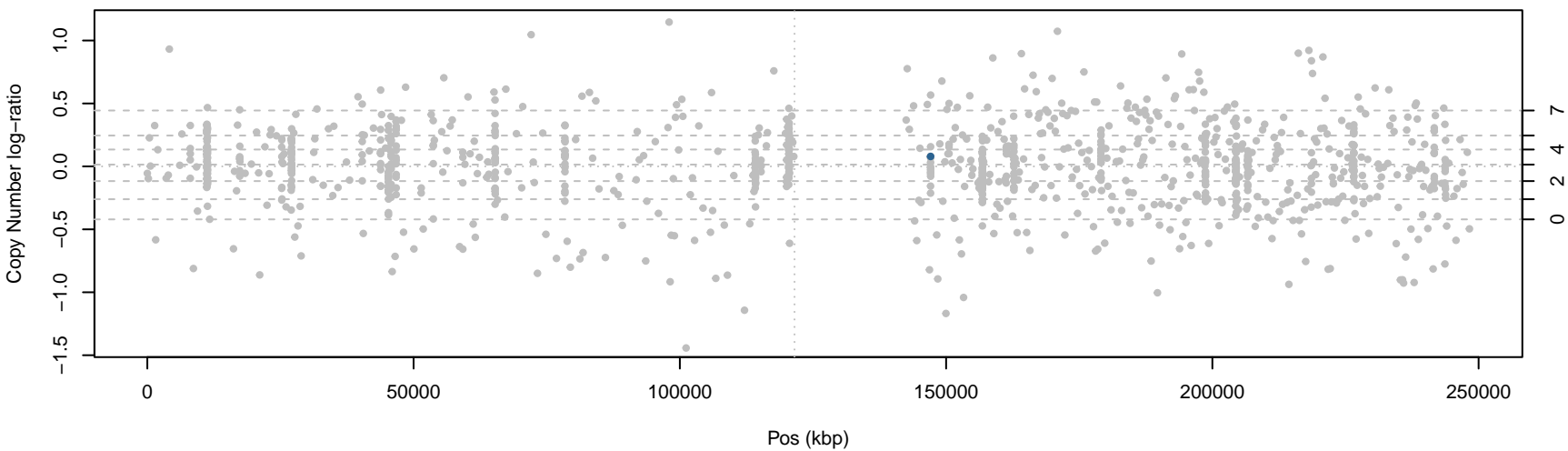
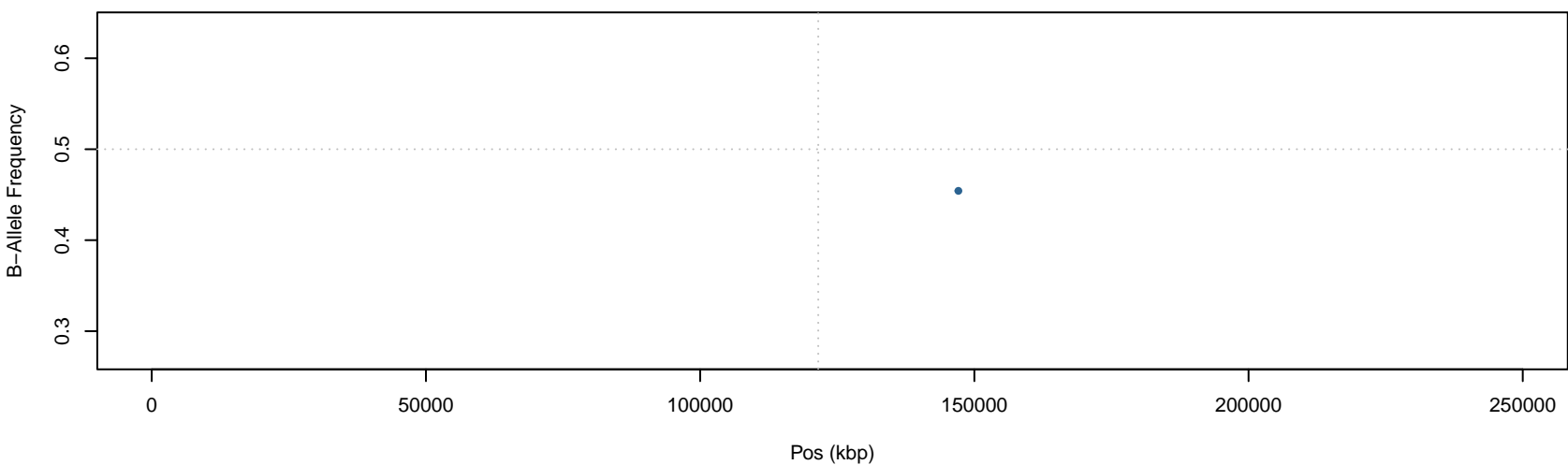
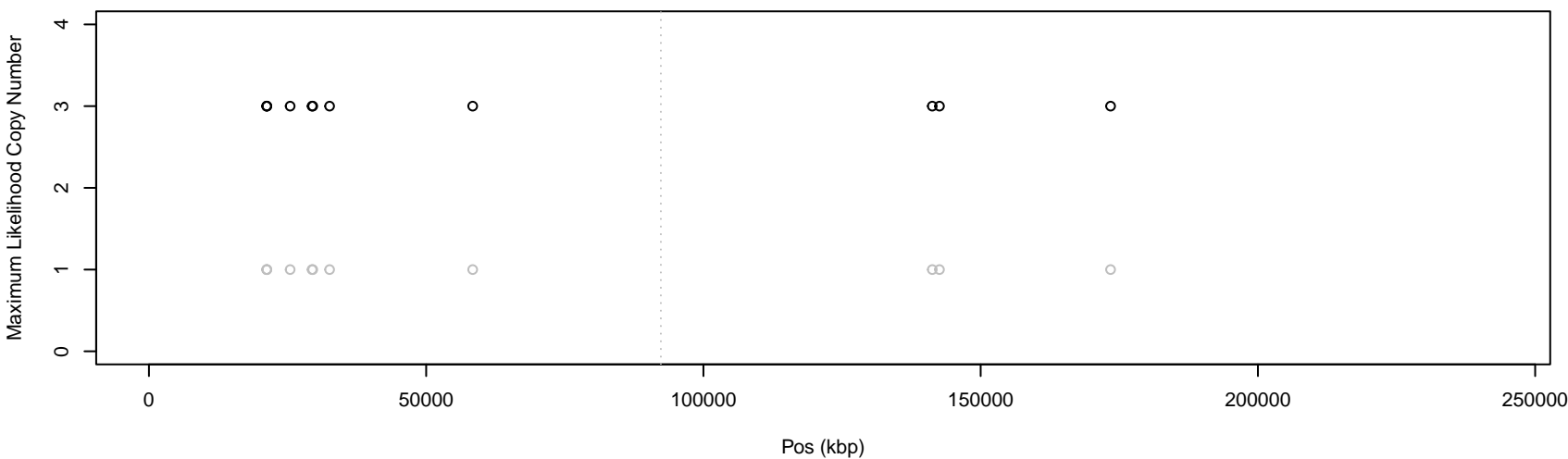
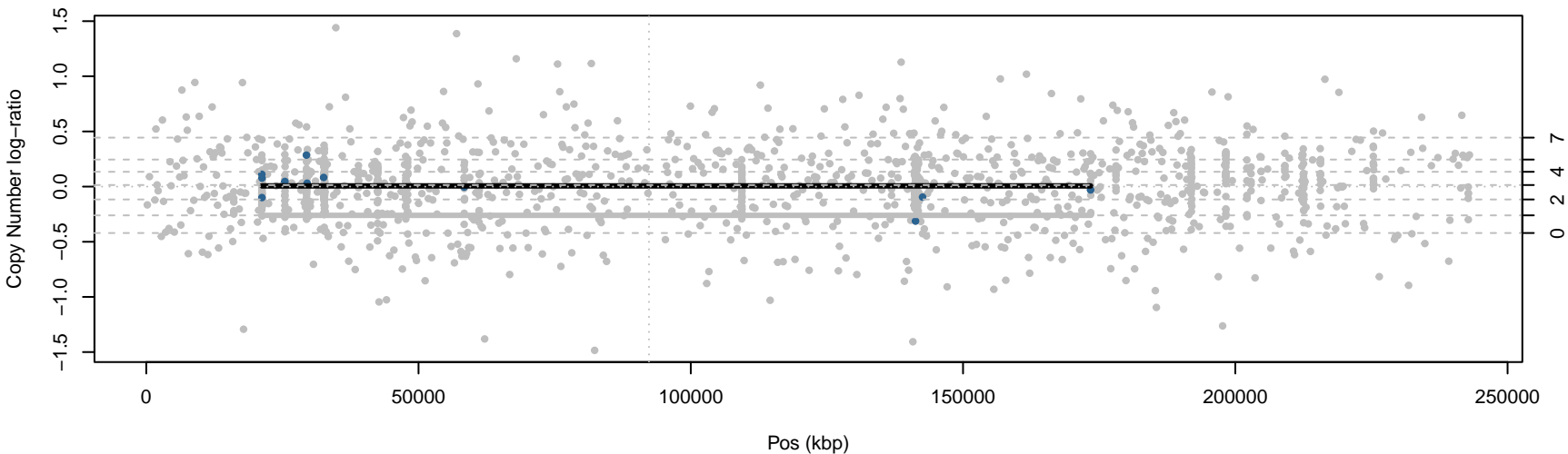
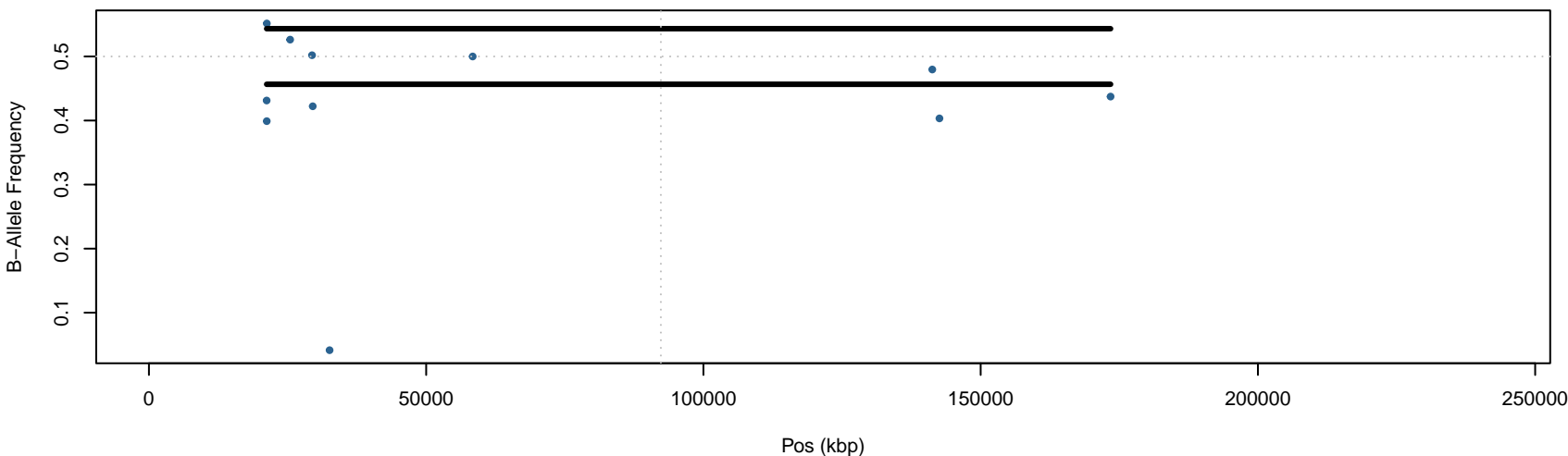


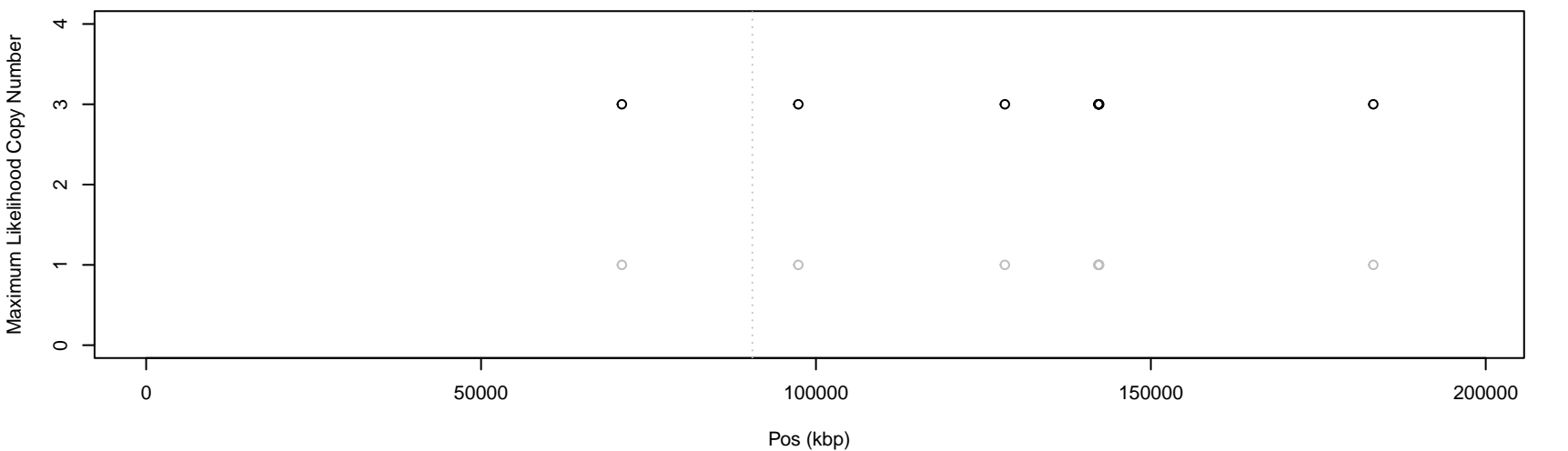
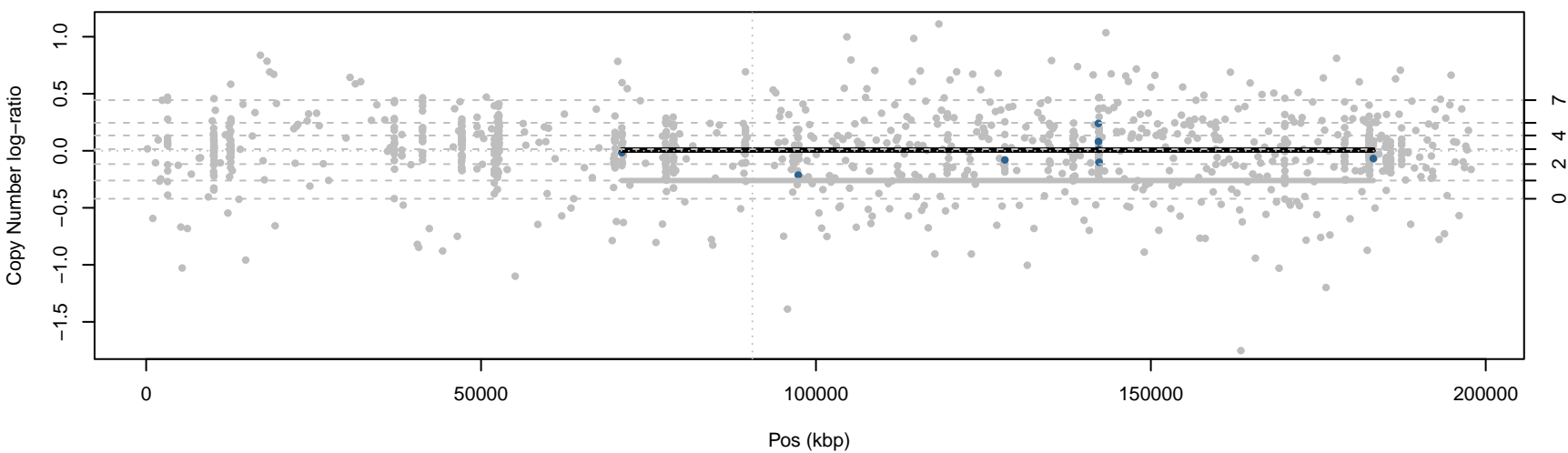
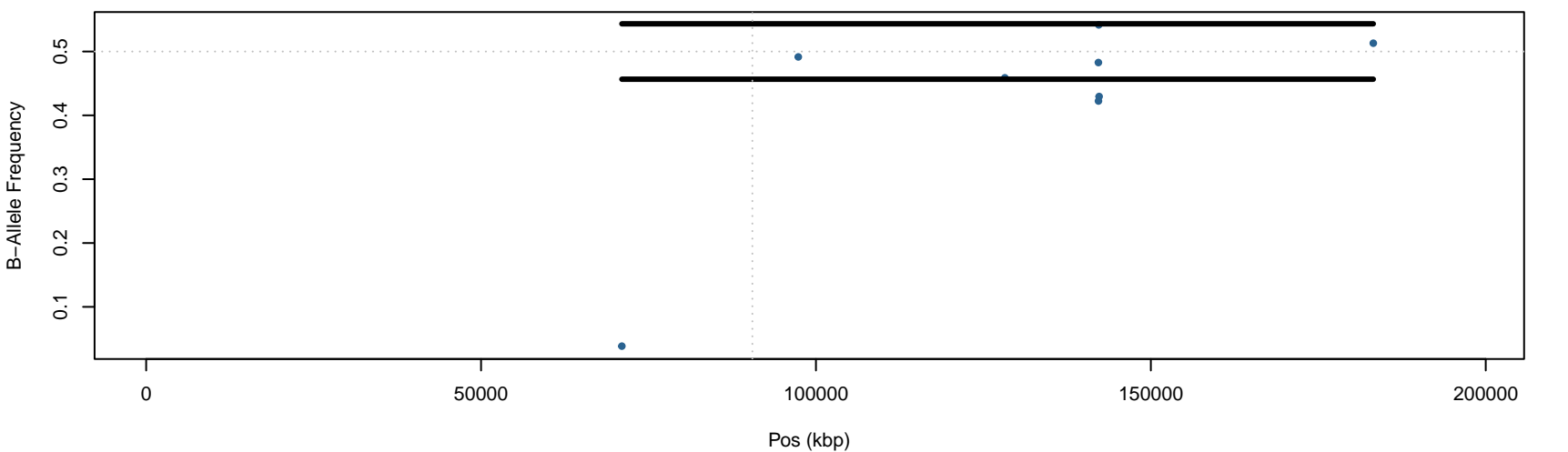
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 1



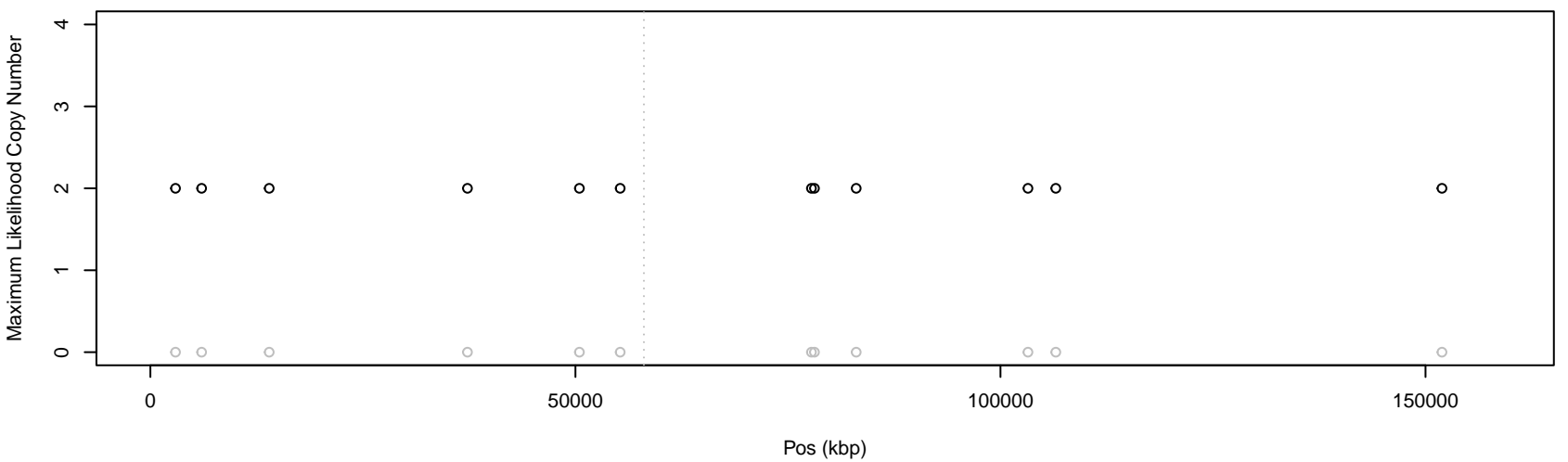
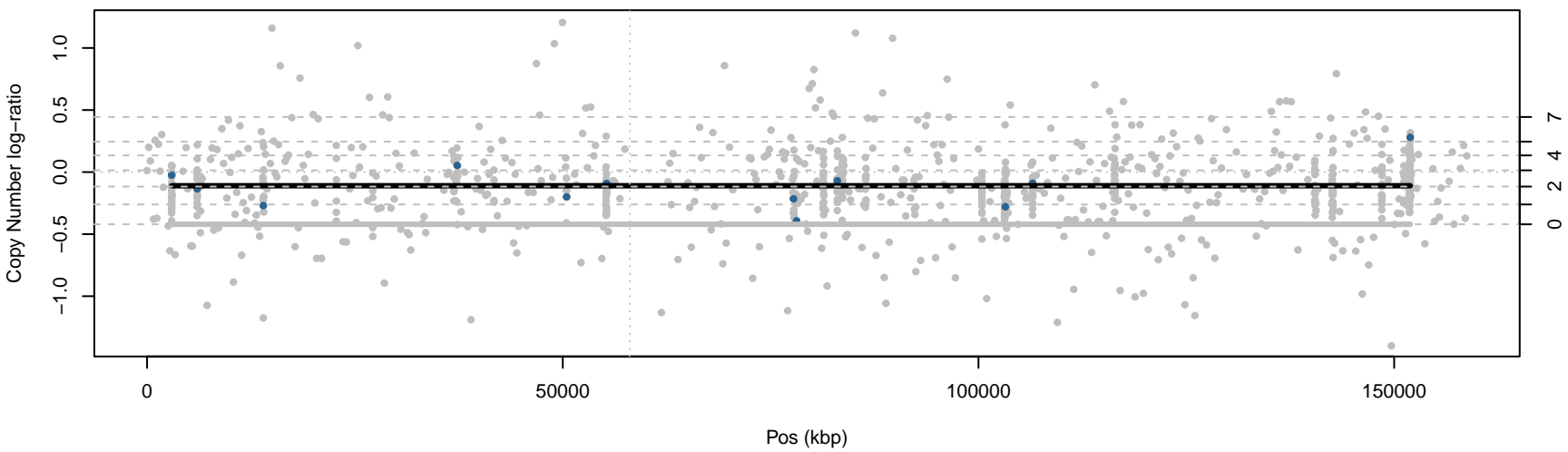
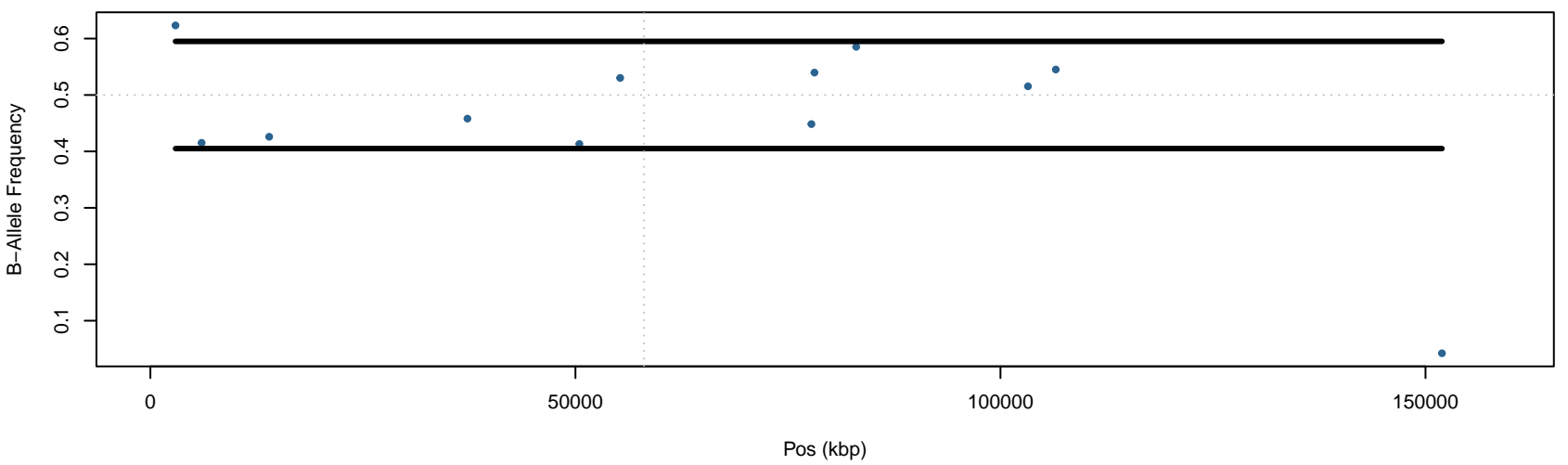
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 2



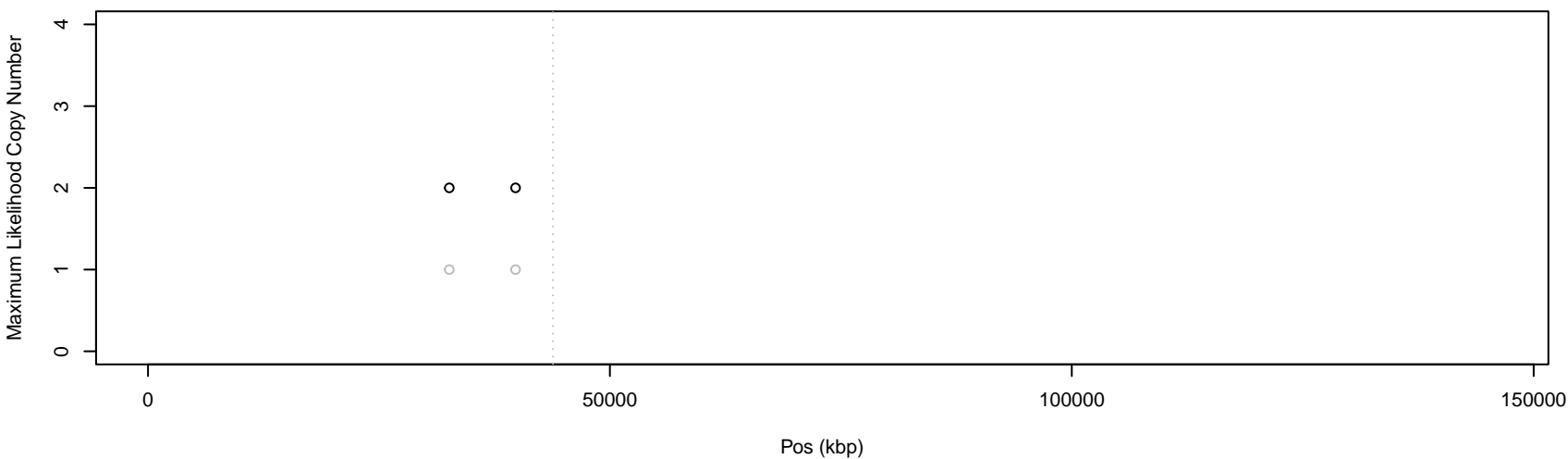
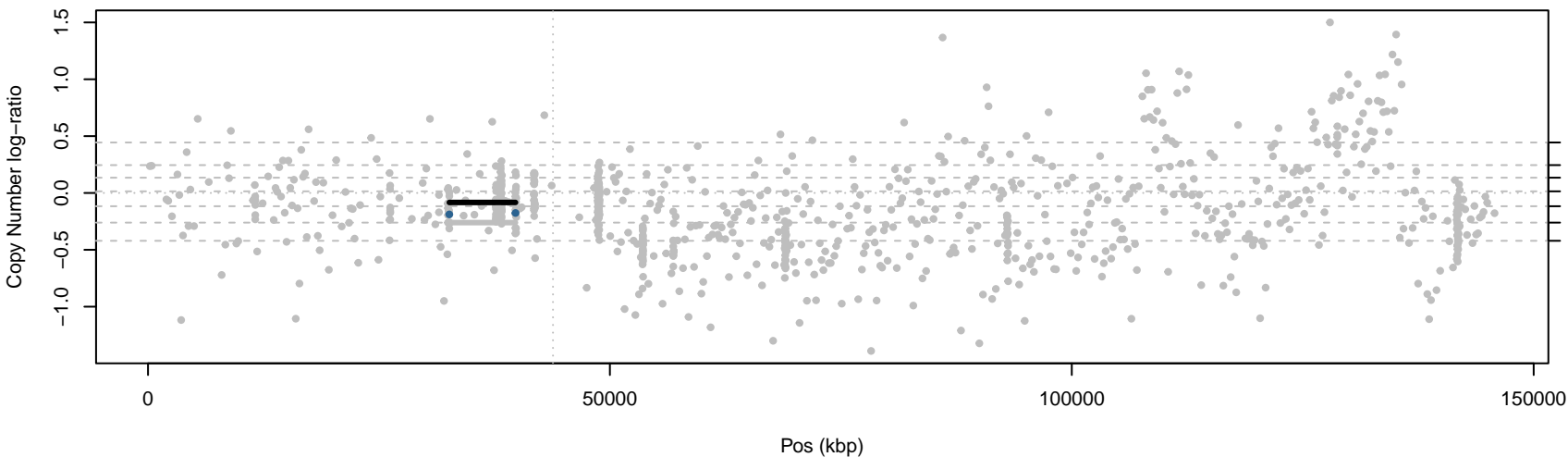
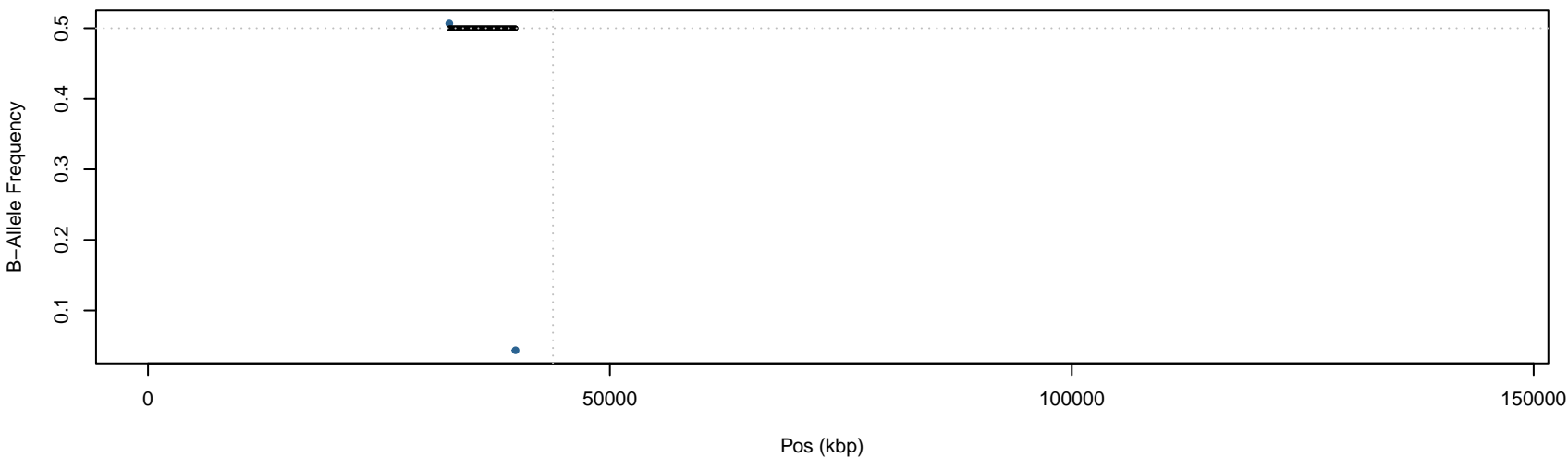
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 3



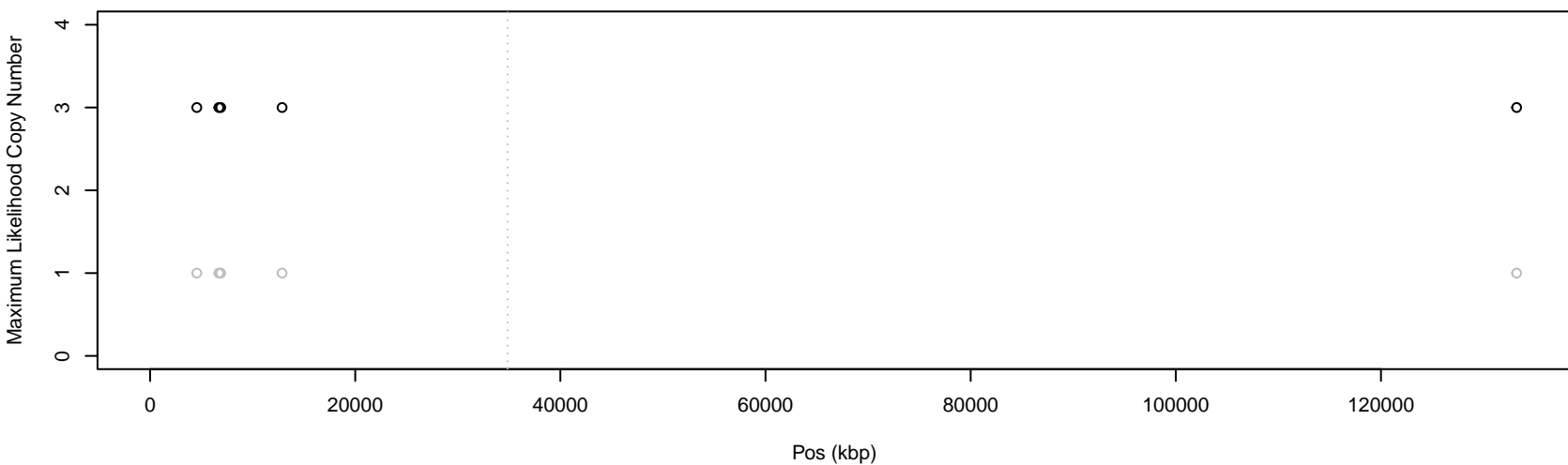
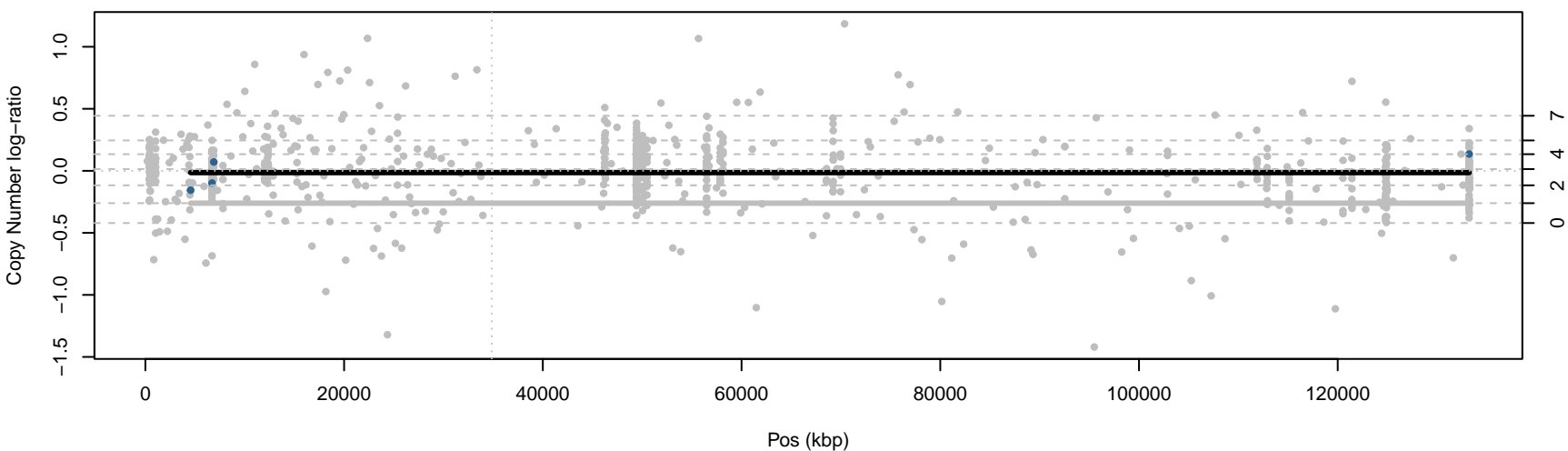
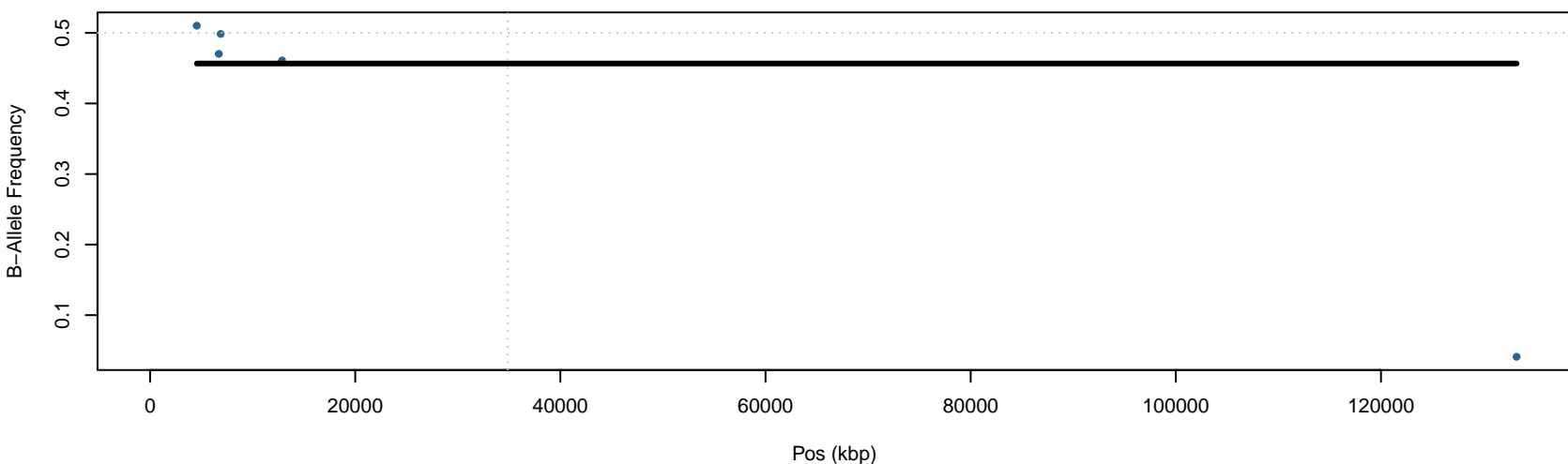
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 7



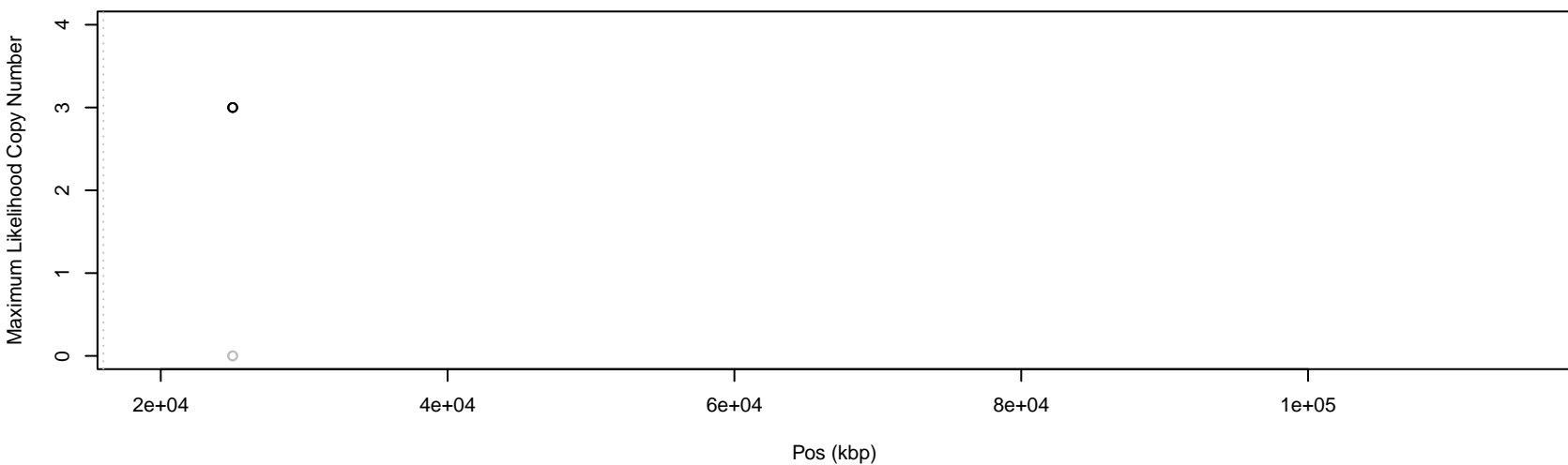
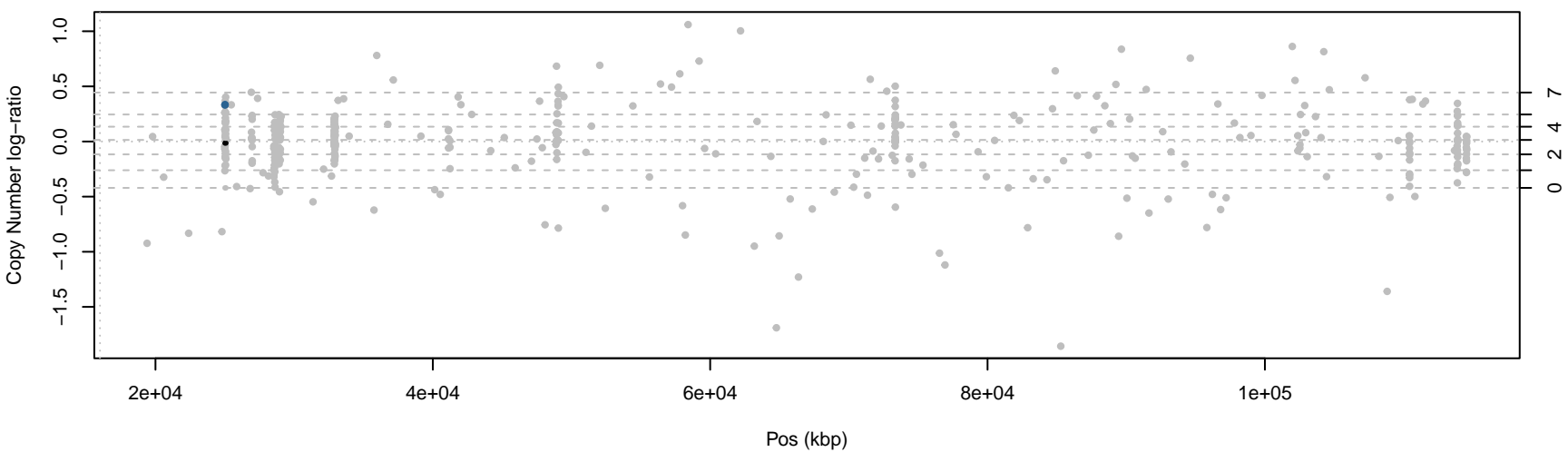
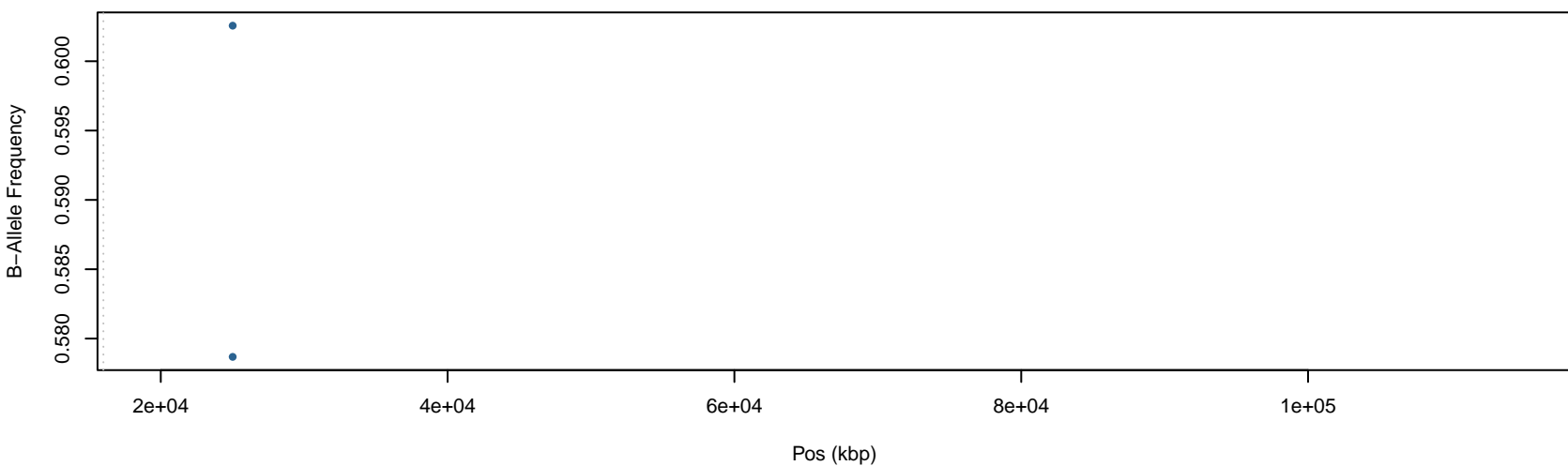
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 8



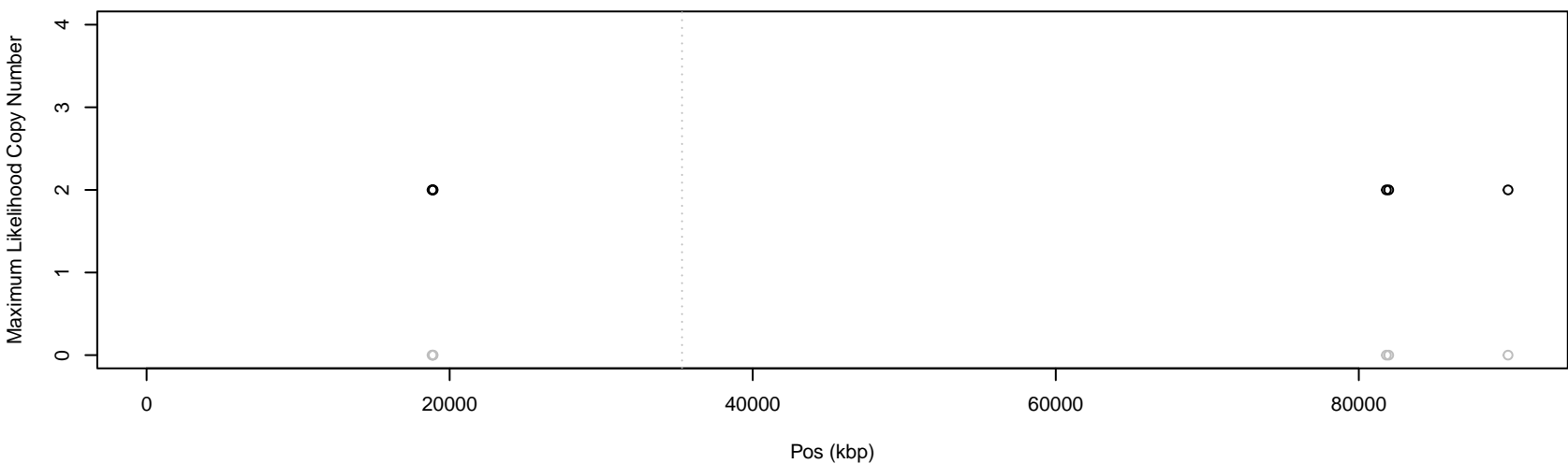
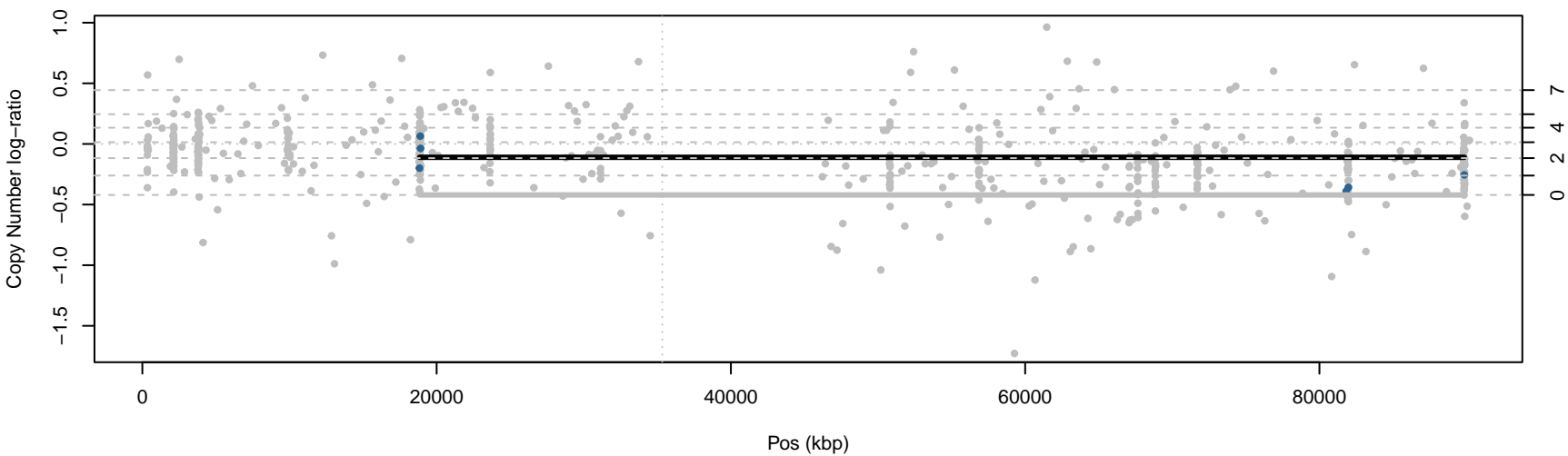
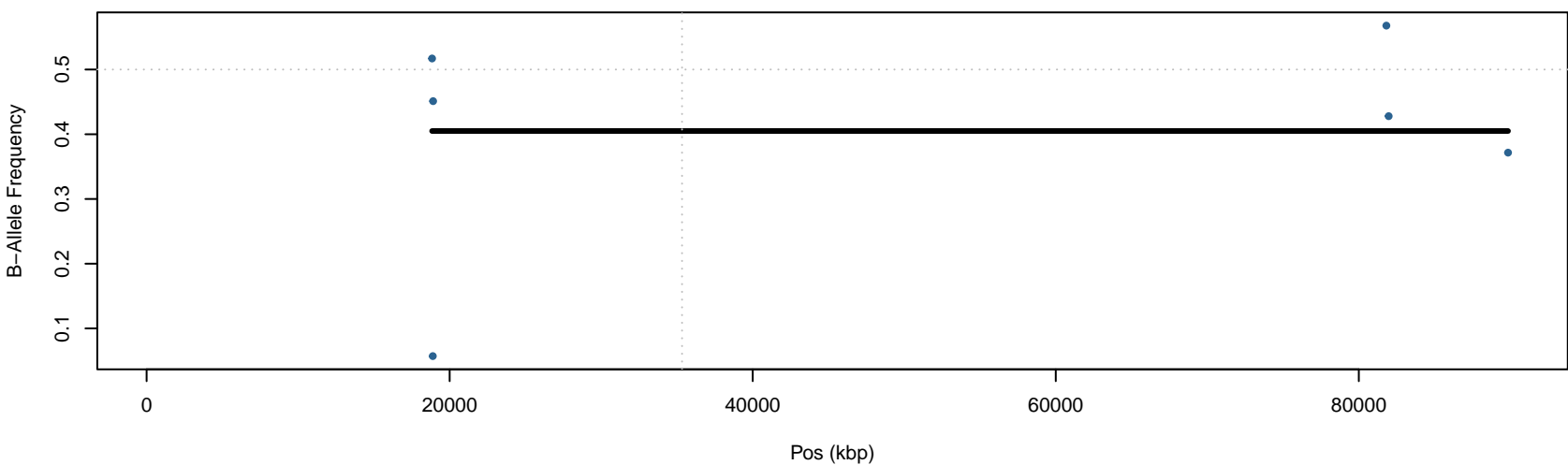
Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 12



Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 13

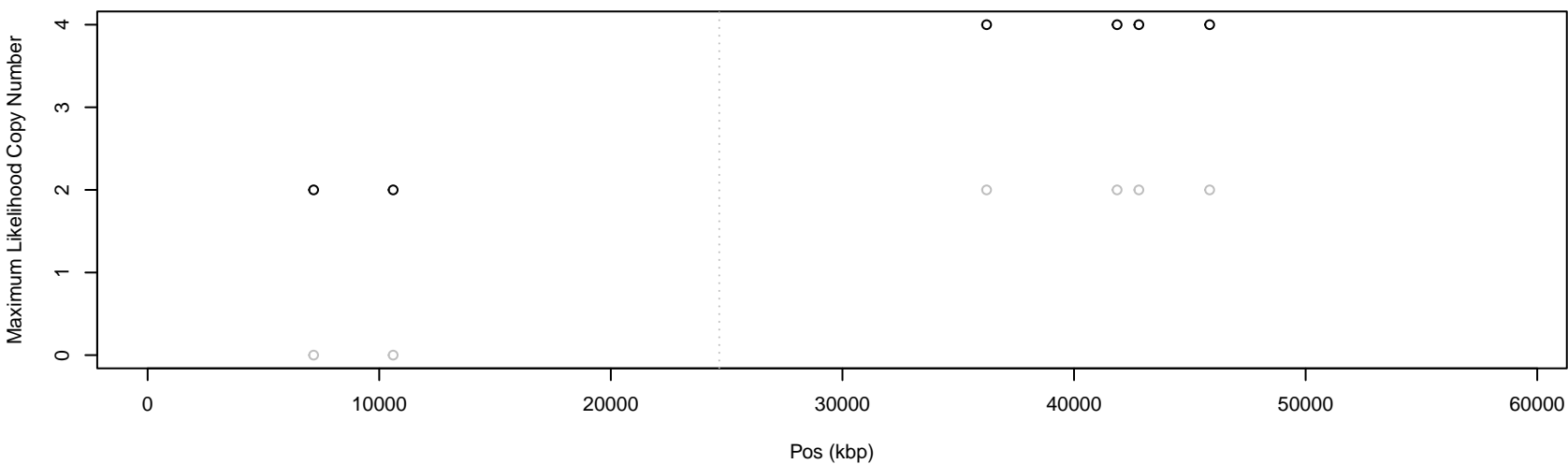
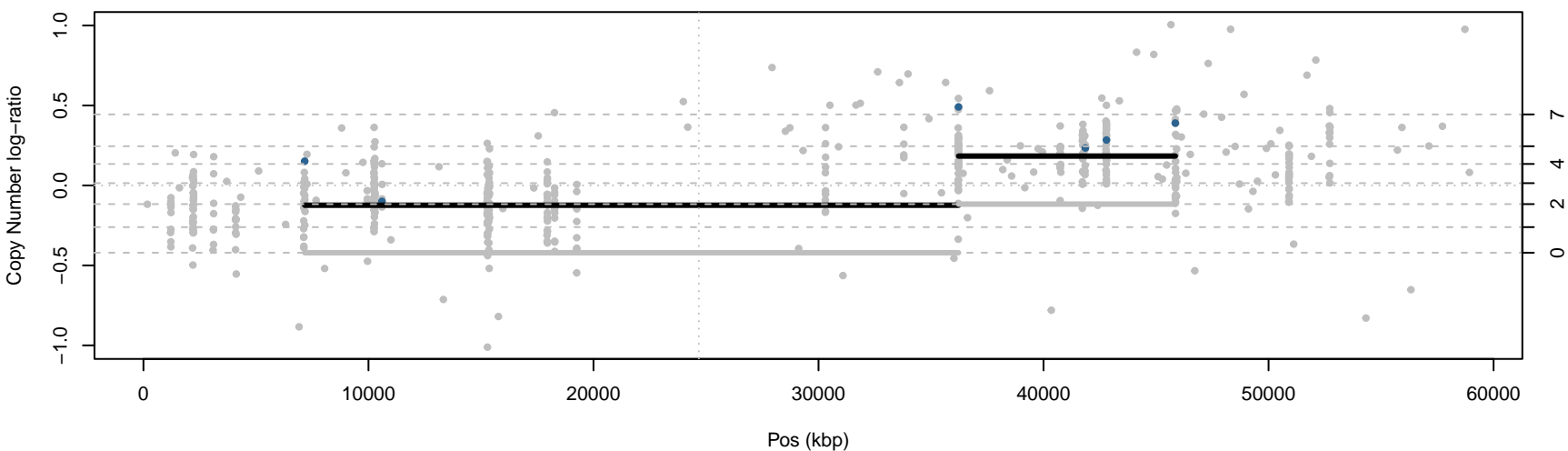
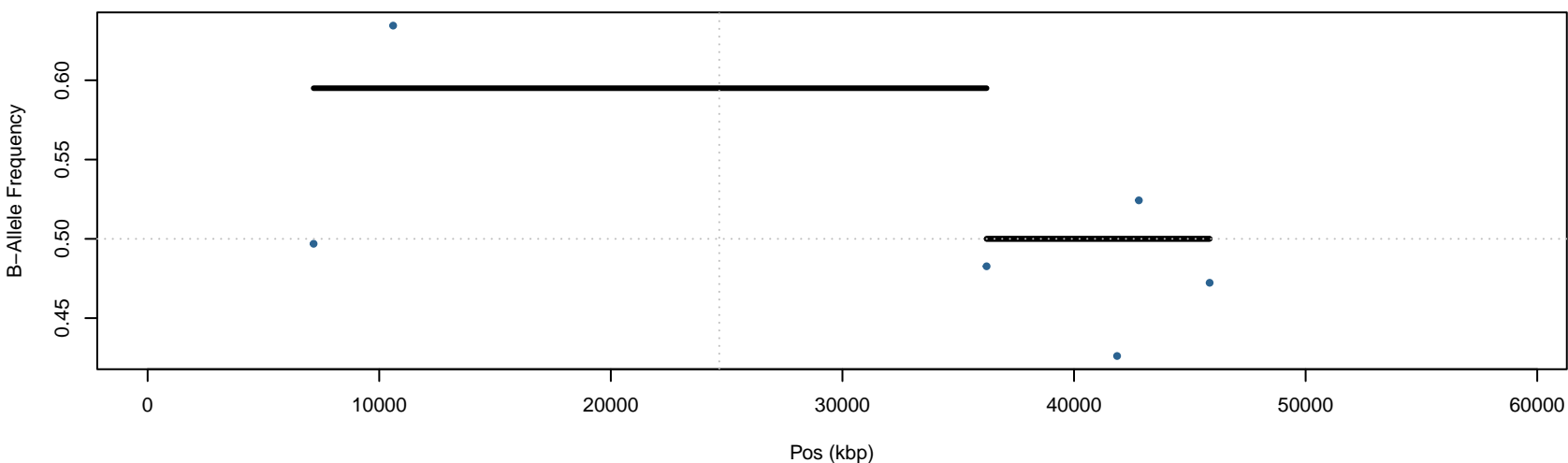


Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 16

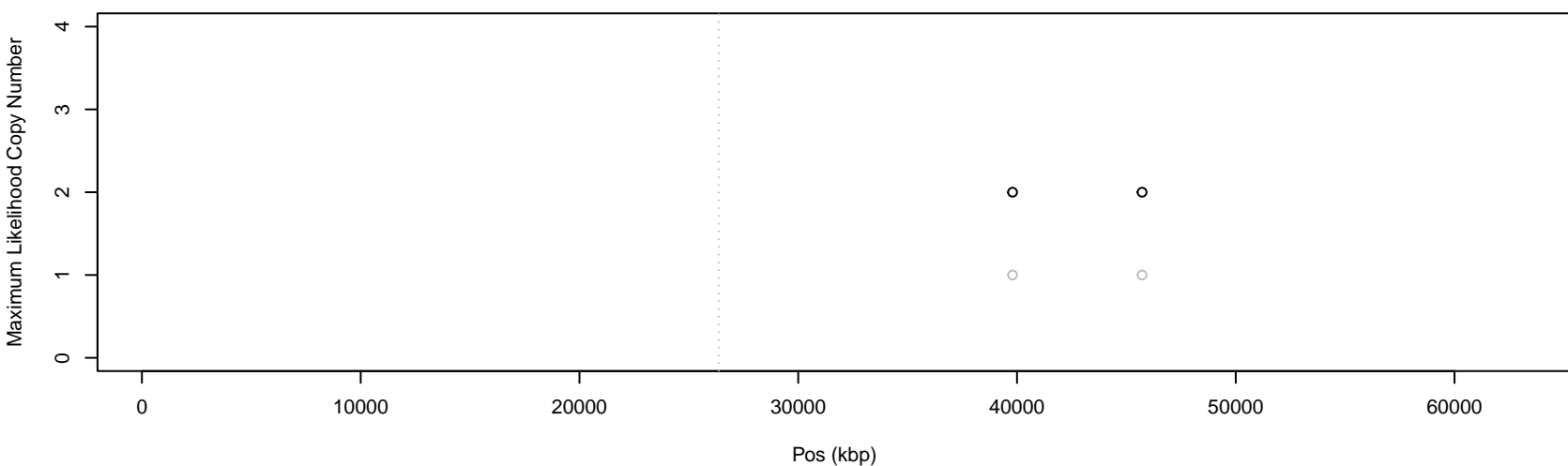
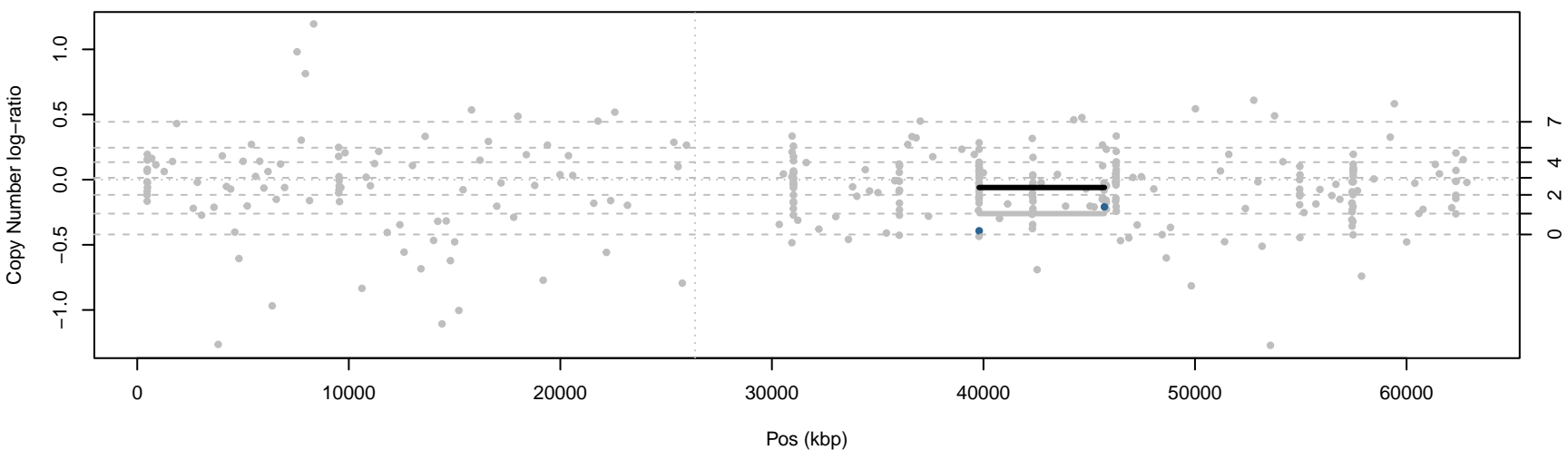
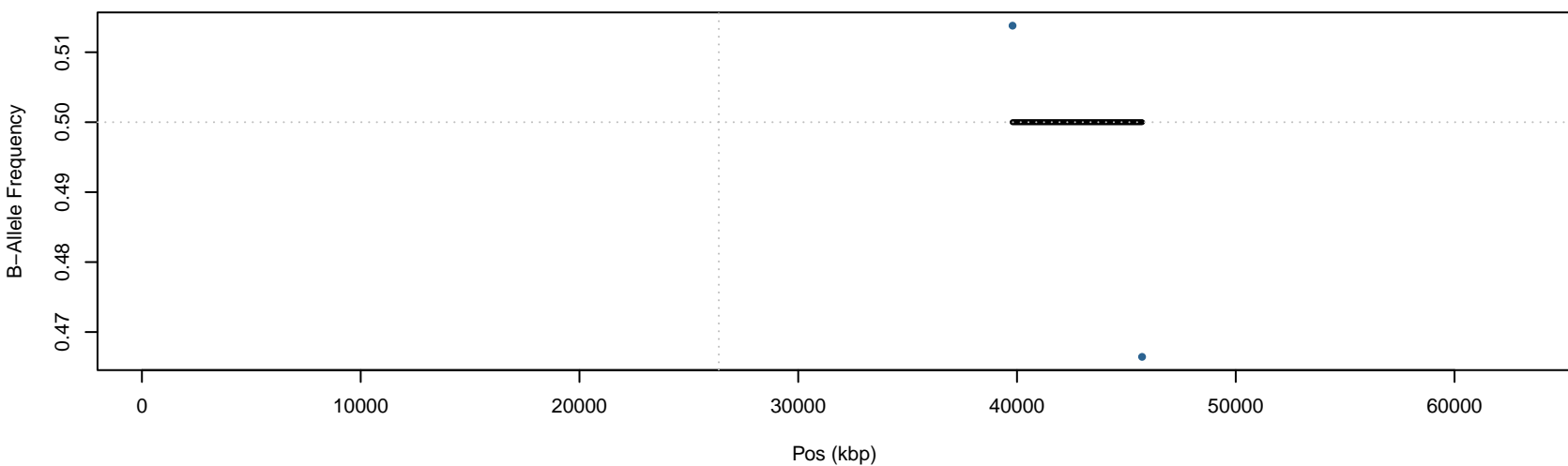




Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 19



Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 20



Purity: 0.19 Tumor ploidy: 2.887 SNV log-likelihood: -0.71 GoF: 96.3% Mean coverage: 545;642 Chromosome: 22

