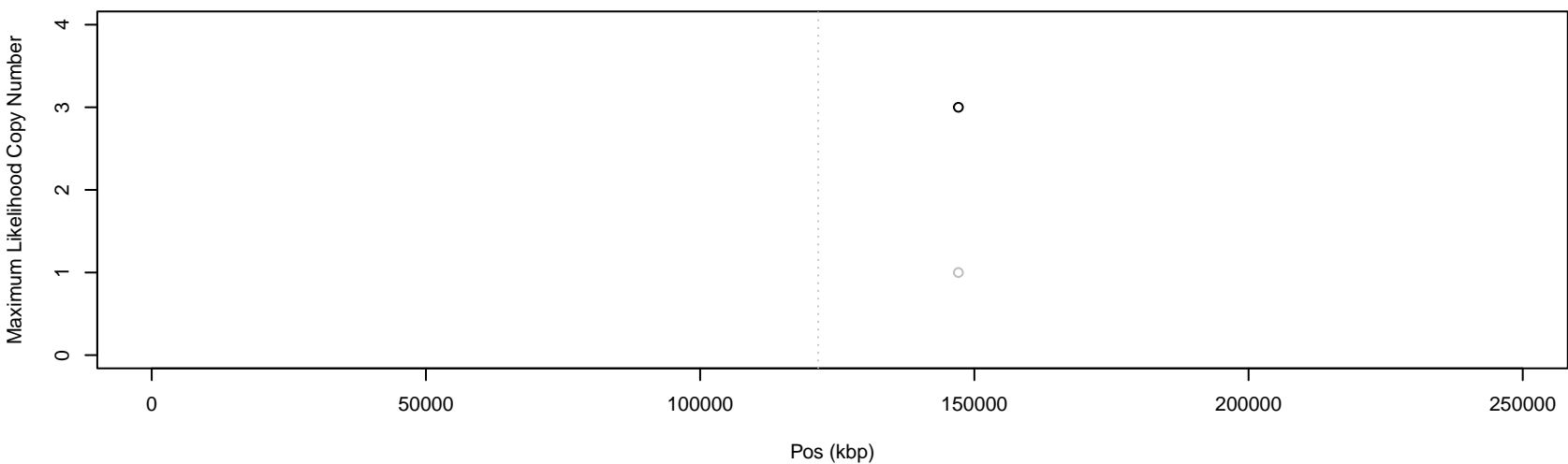
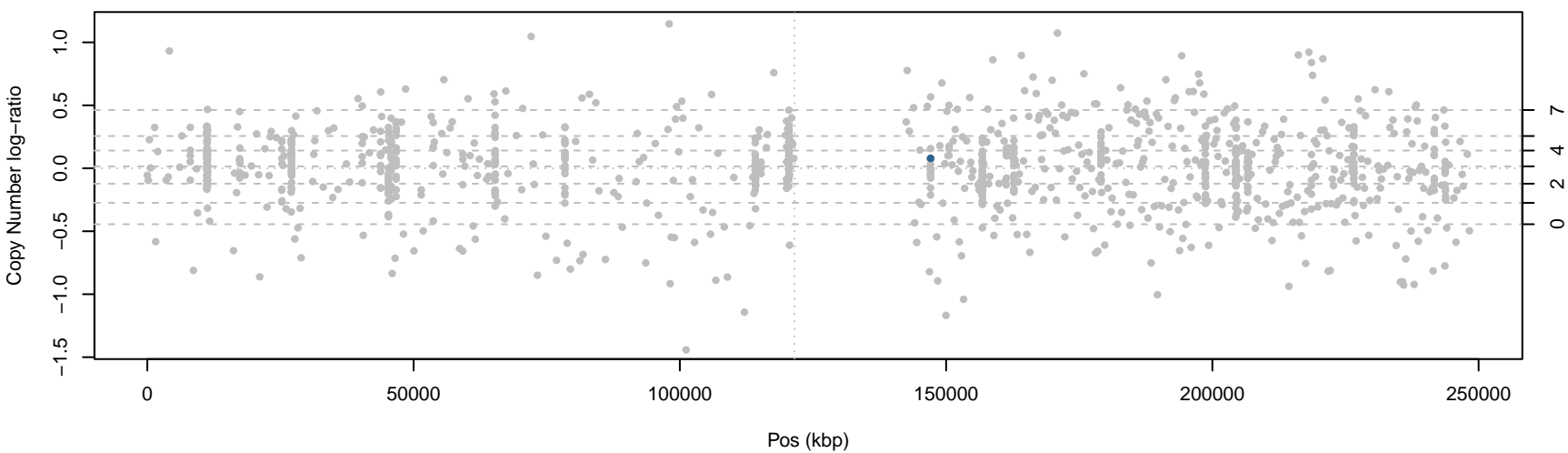
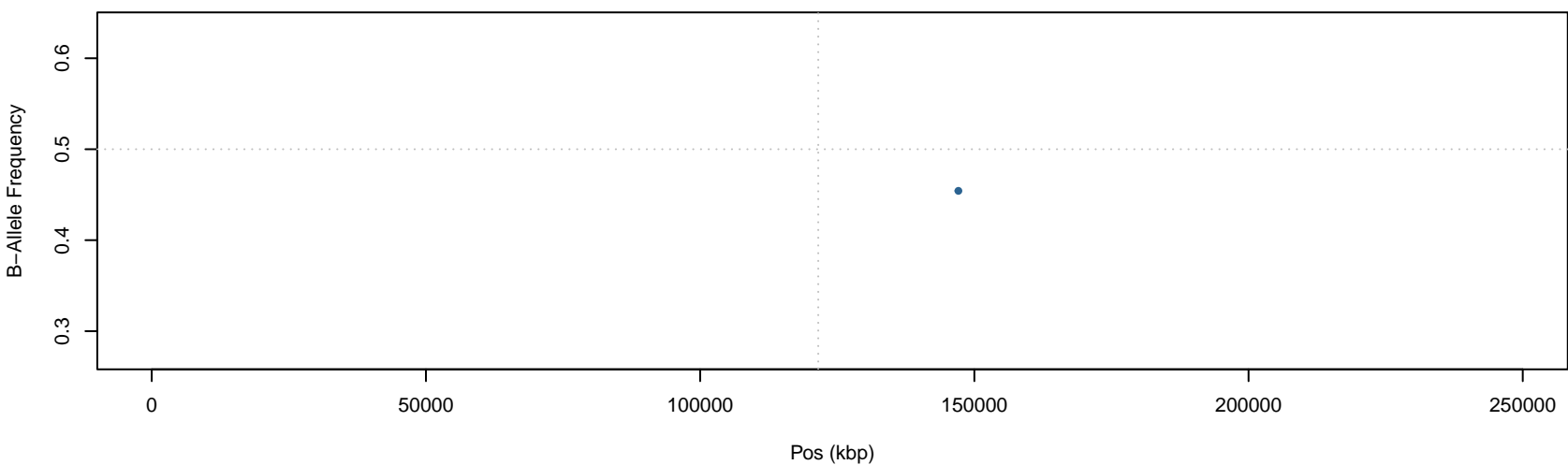
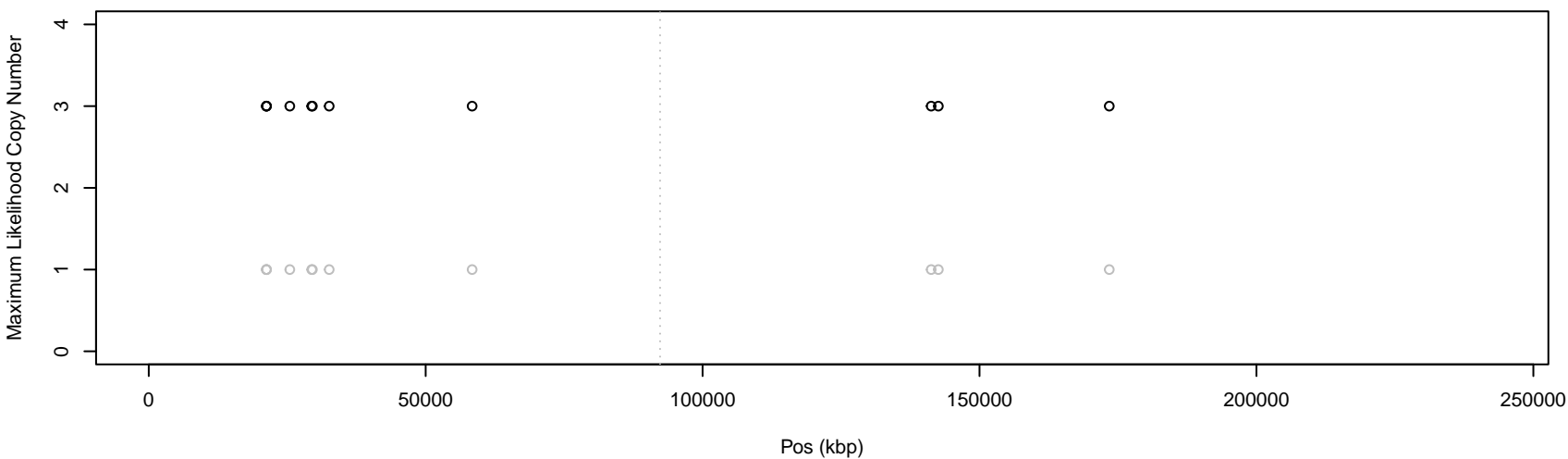
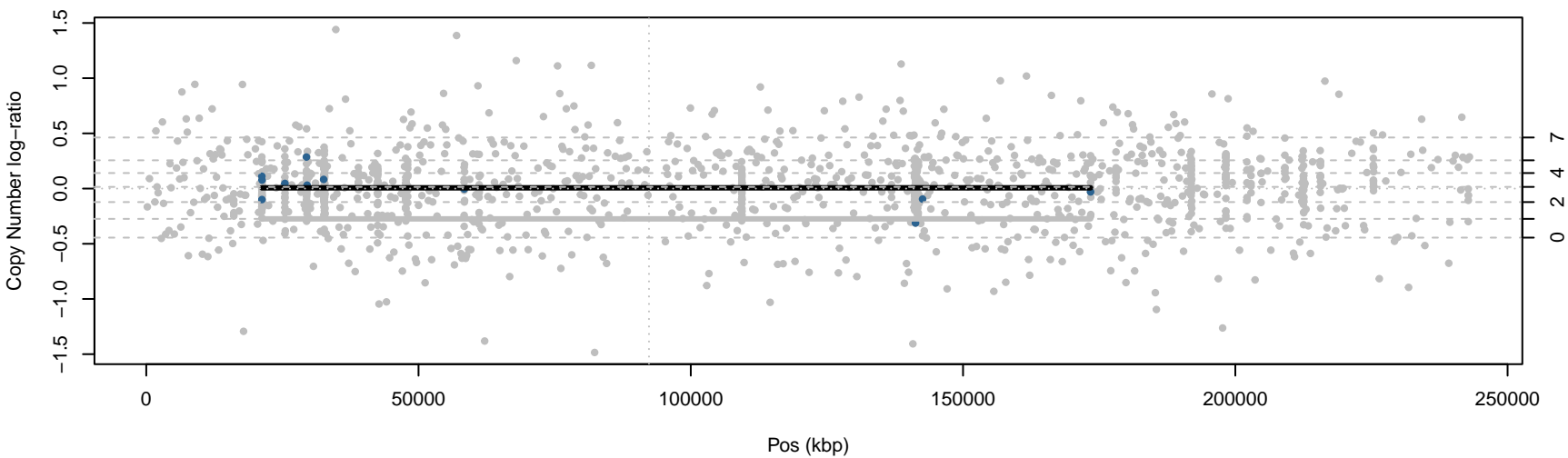
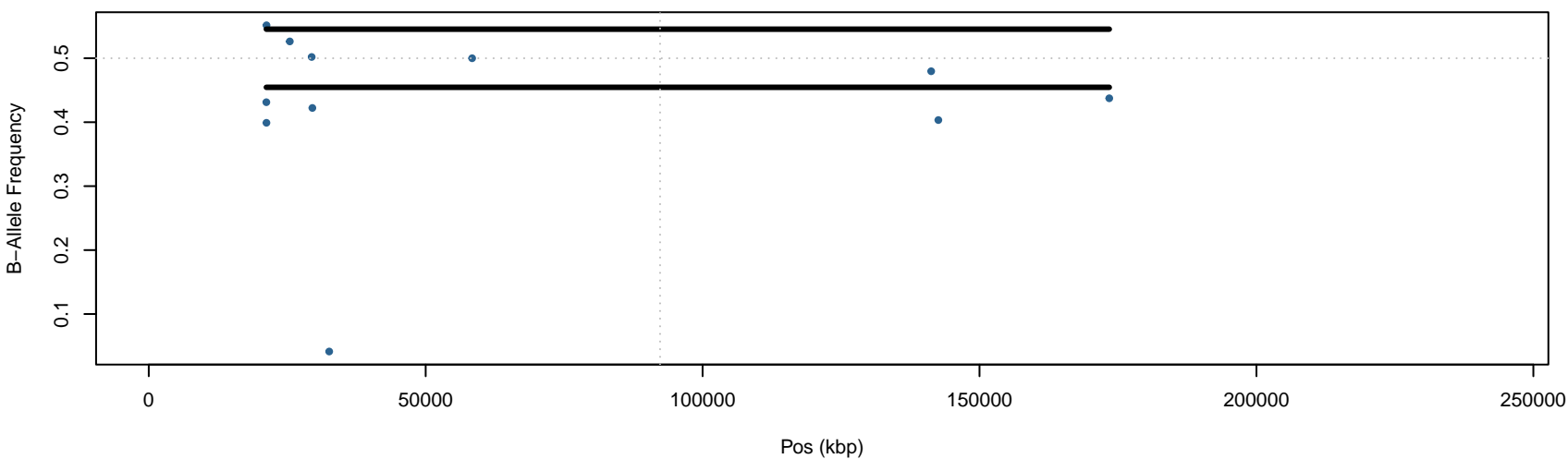


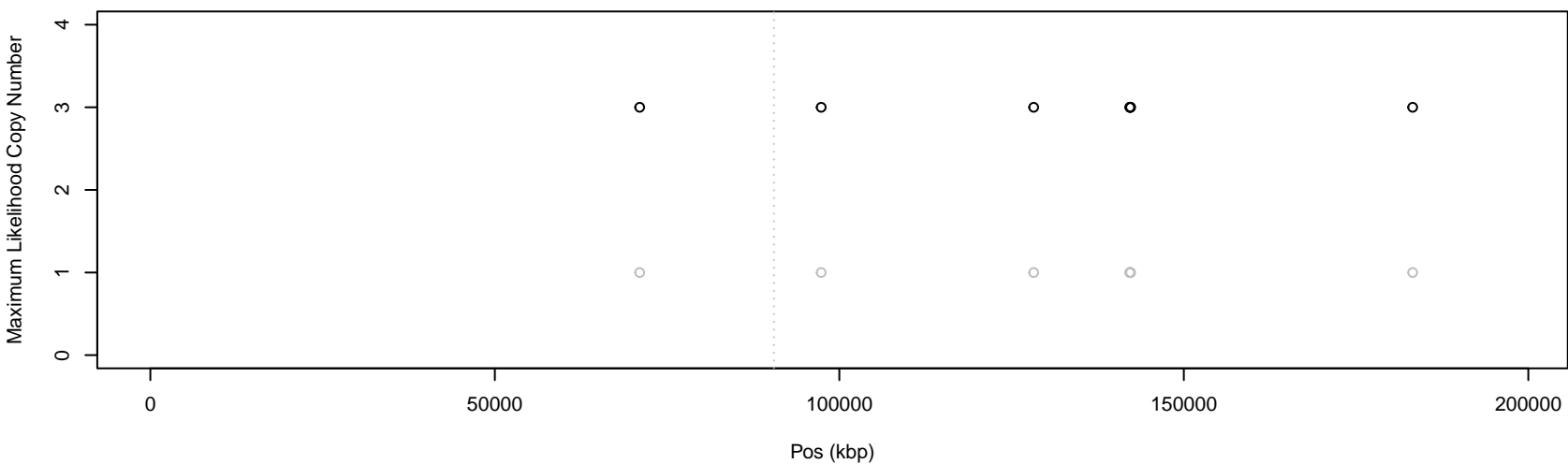
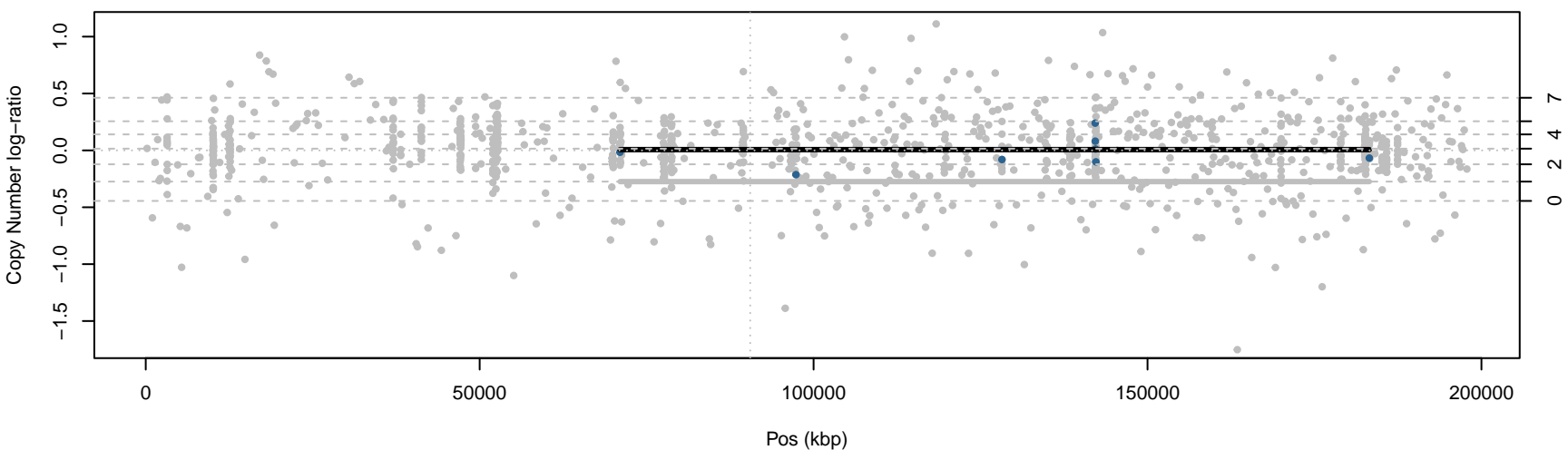
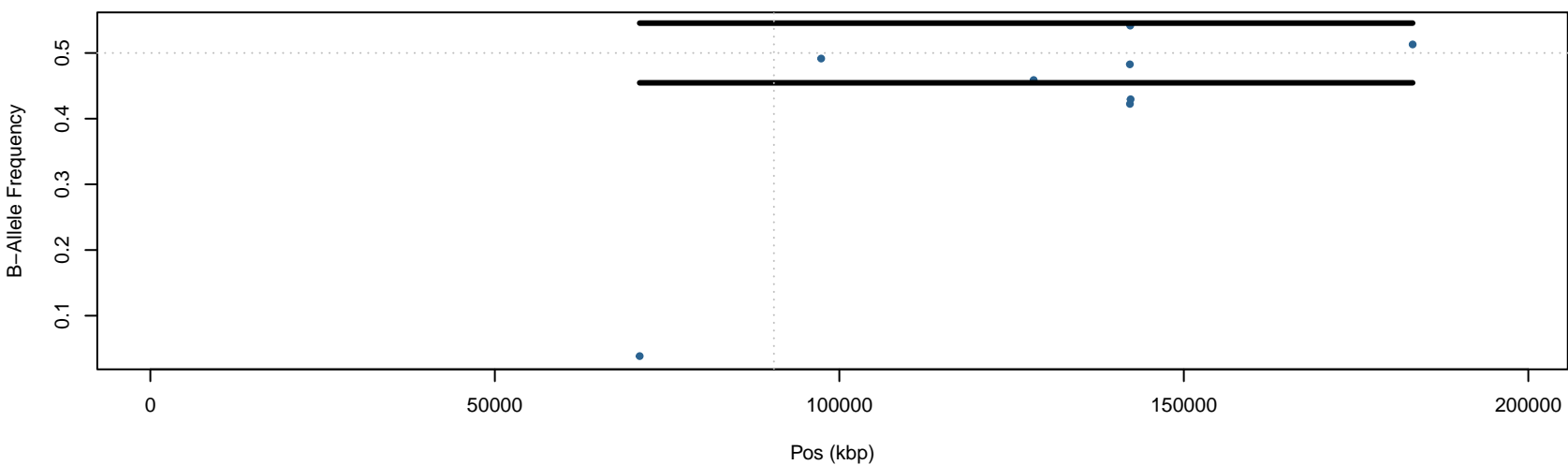
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 1



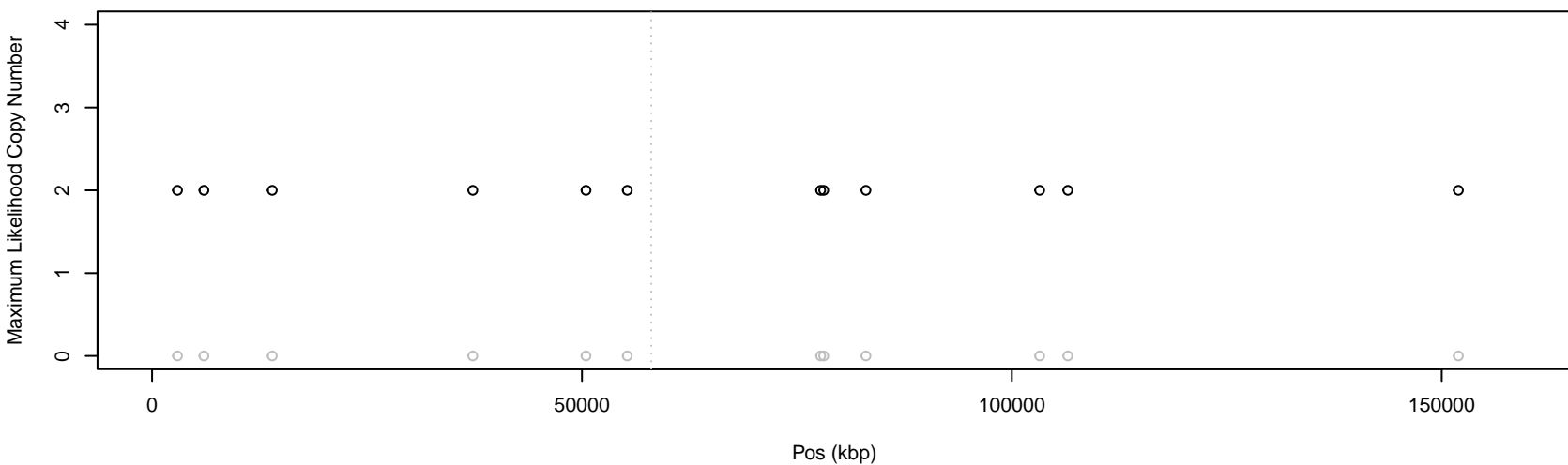
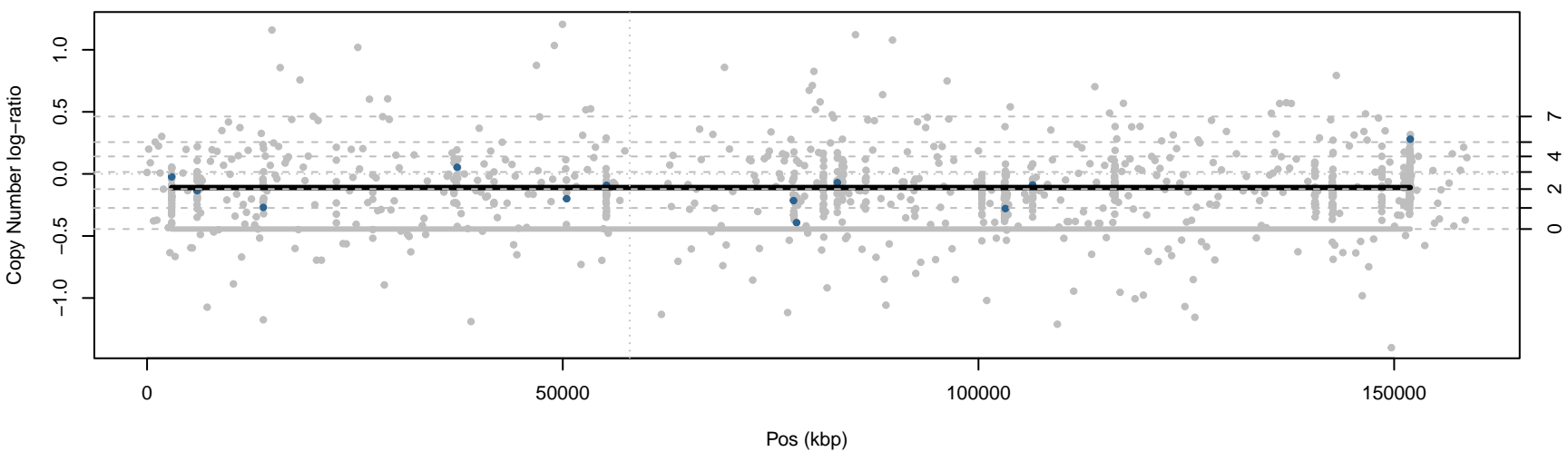
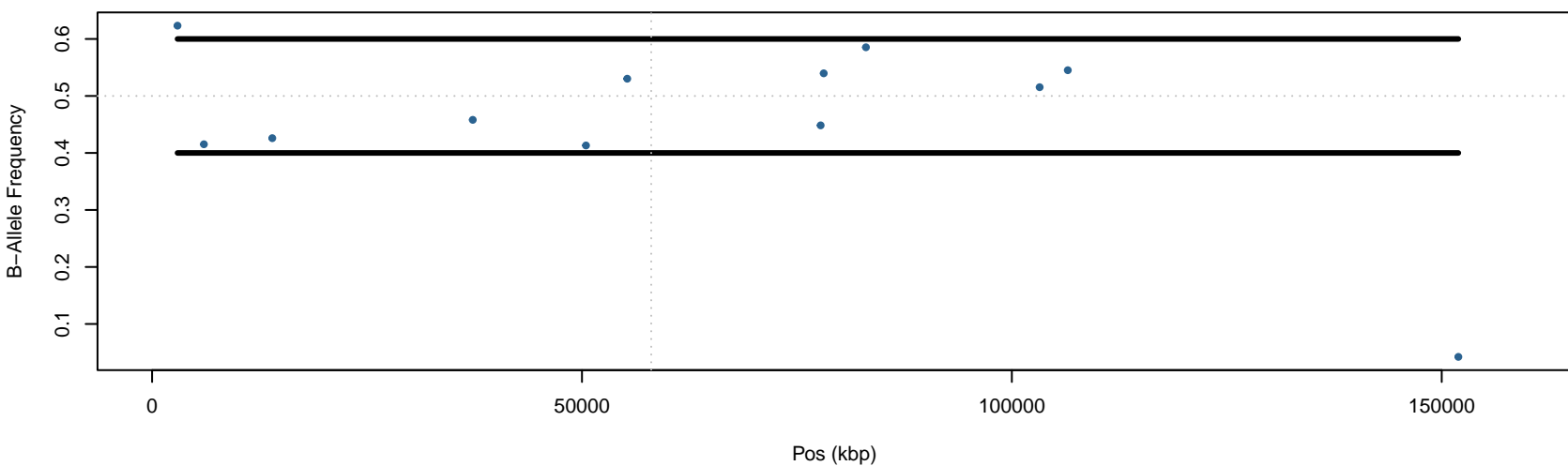
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 2



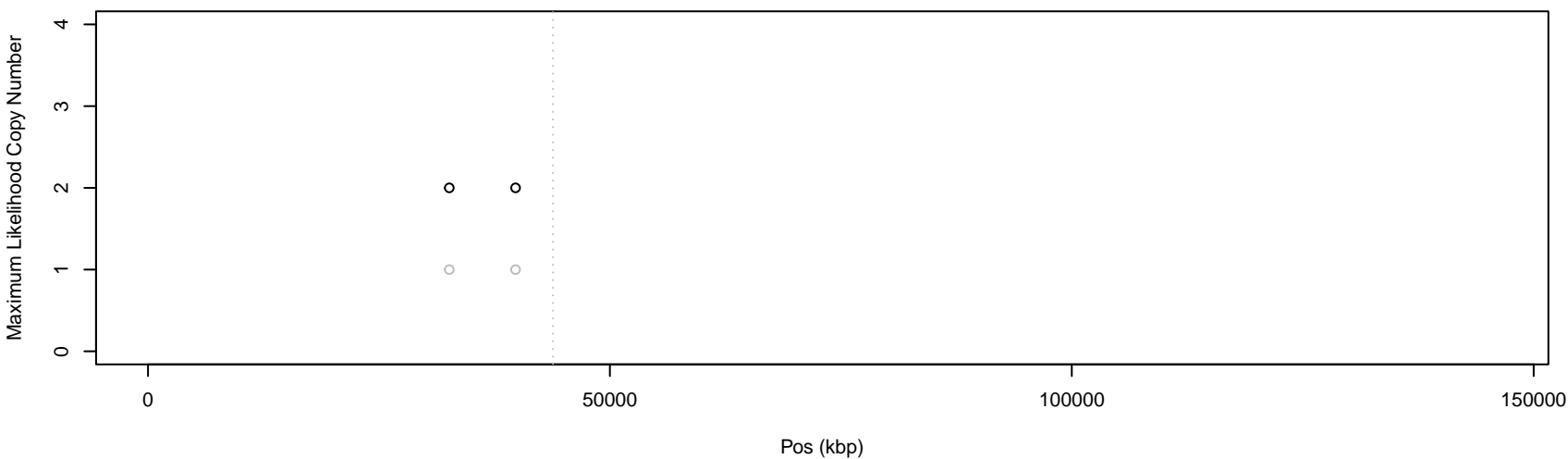
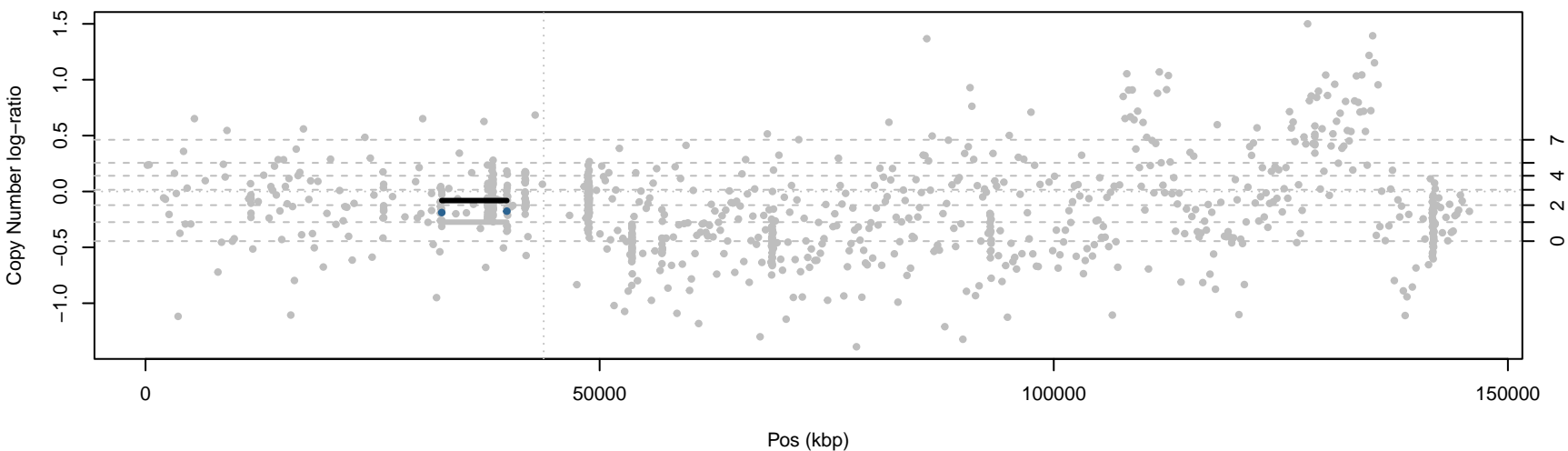
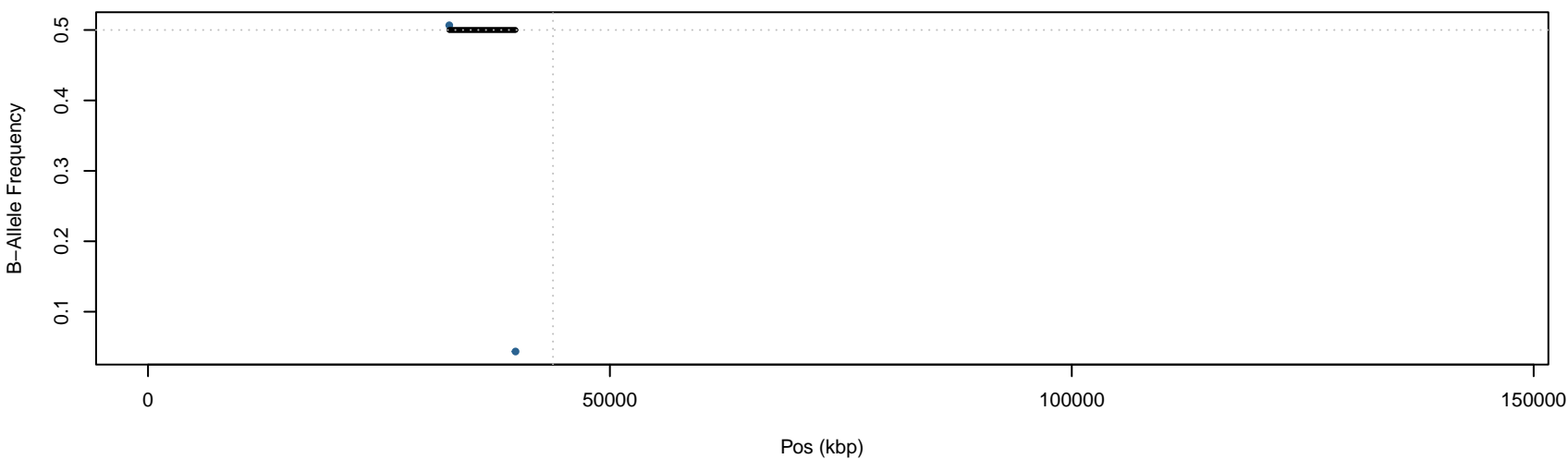
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 3



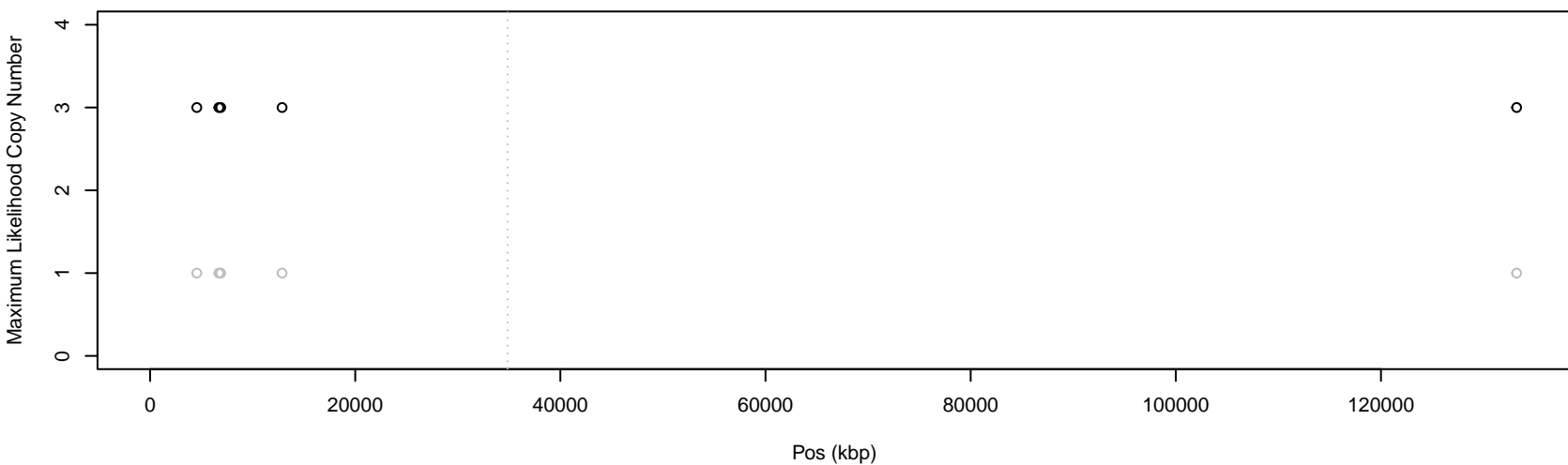
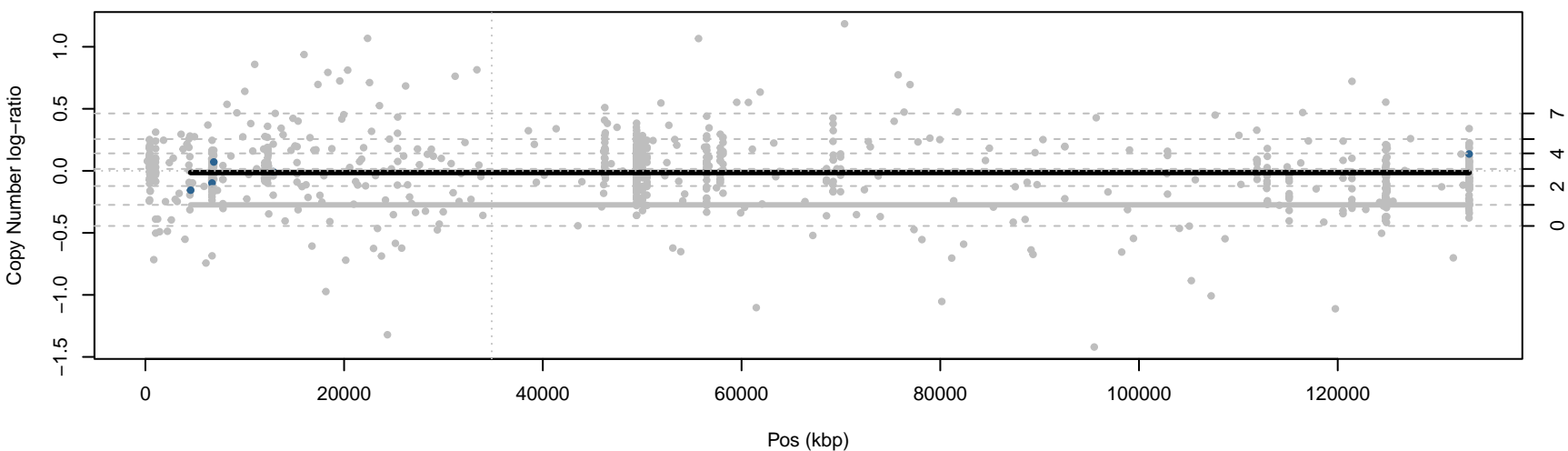
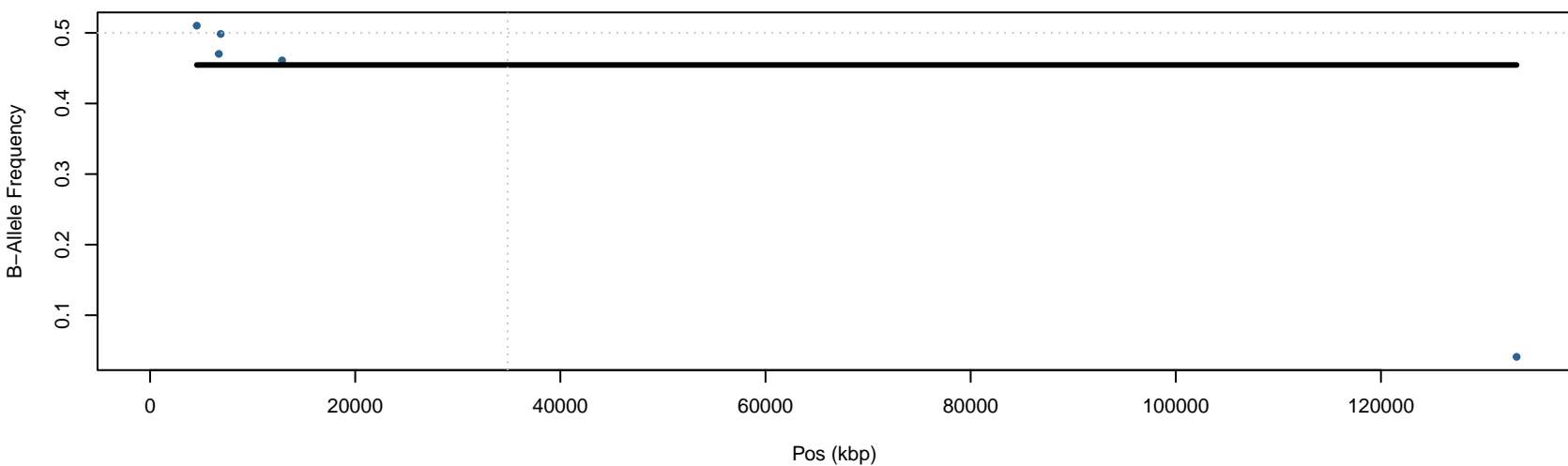
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 7



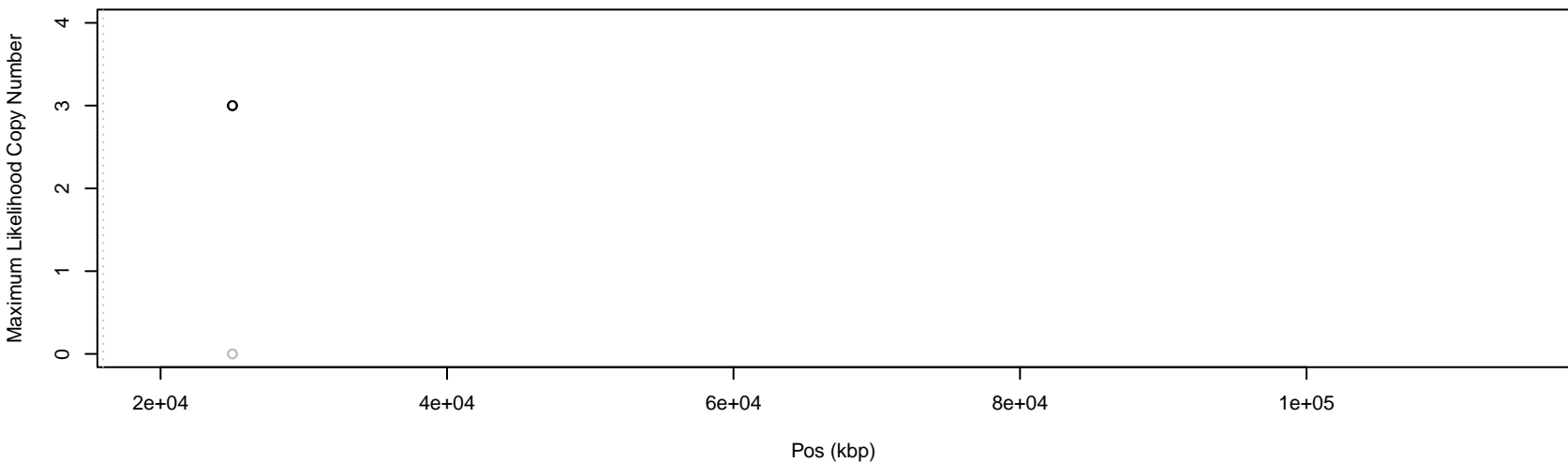
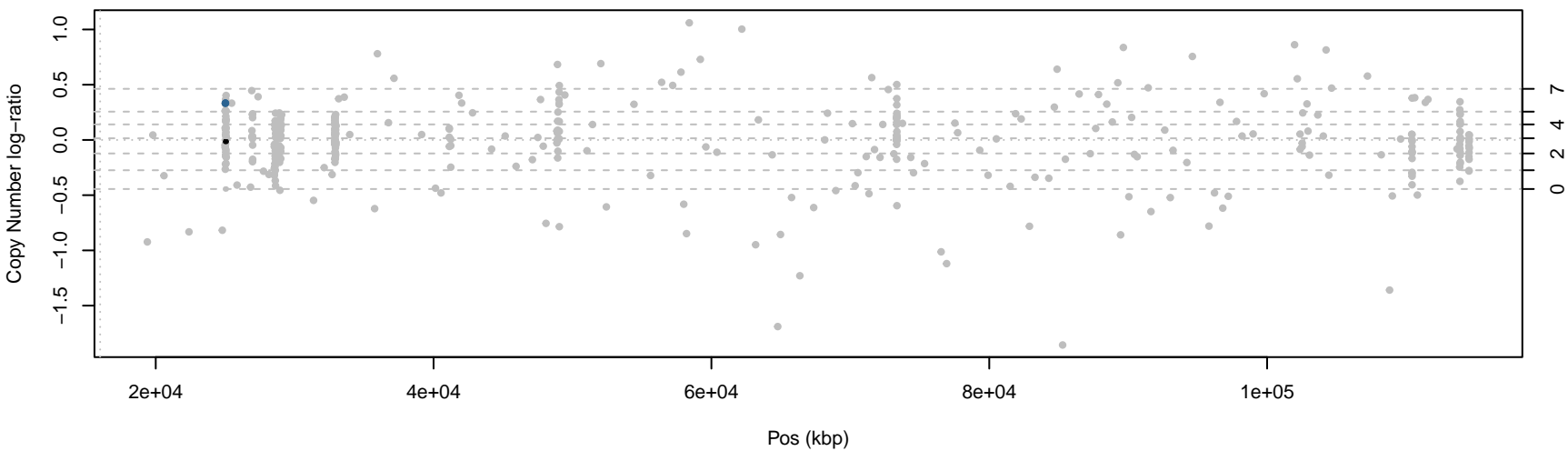
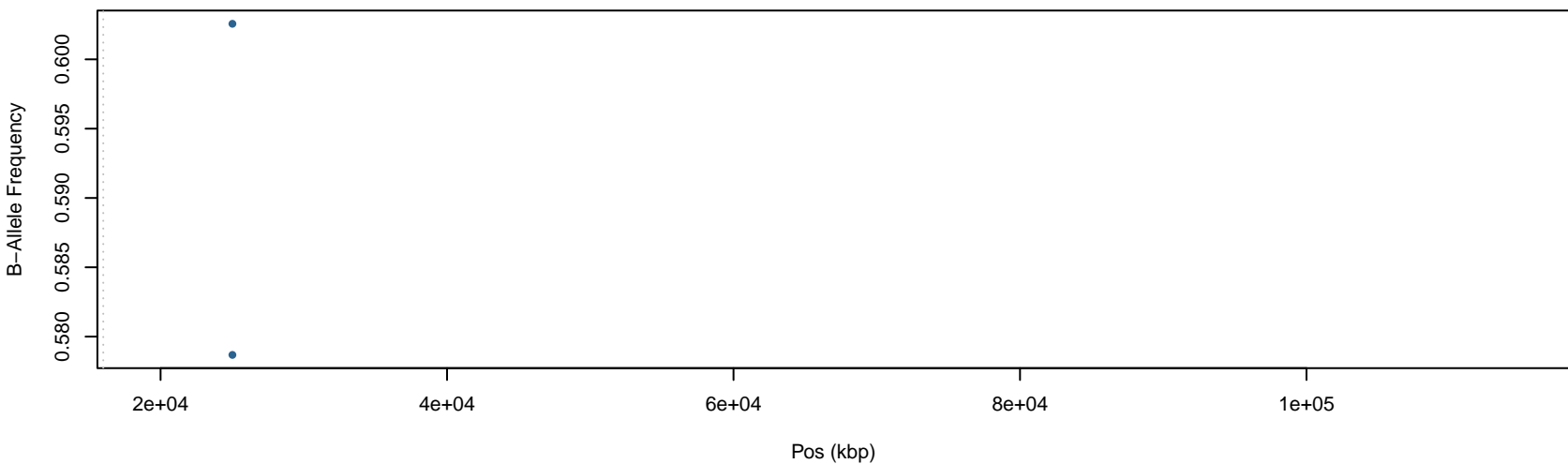
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 8



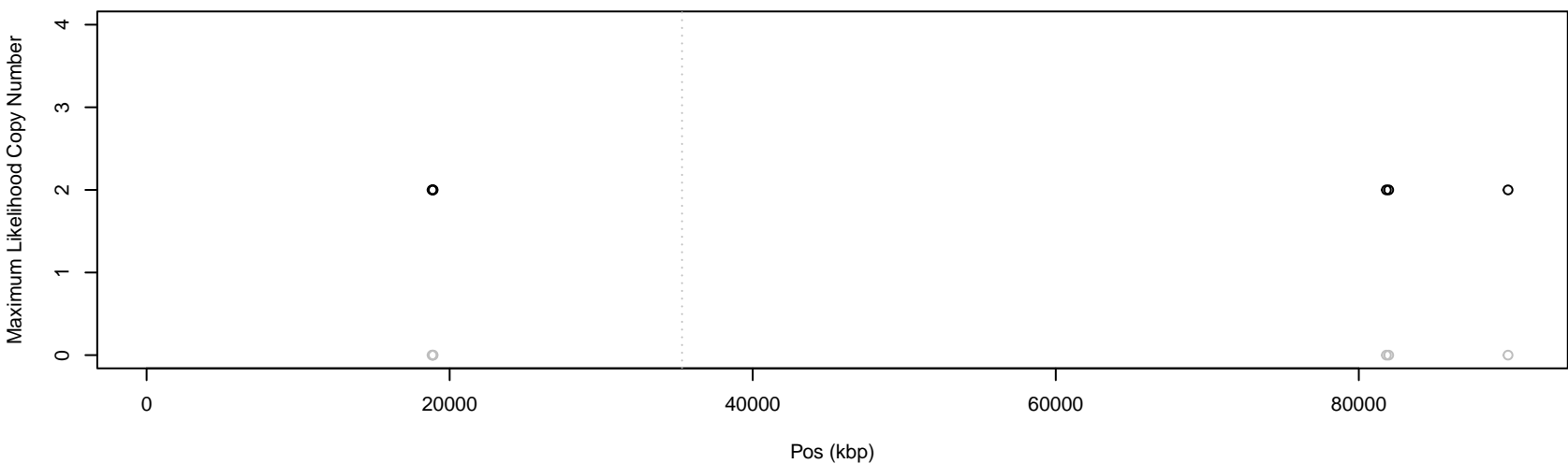
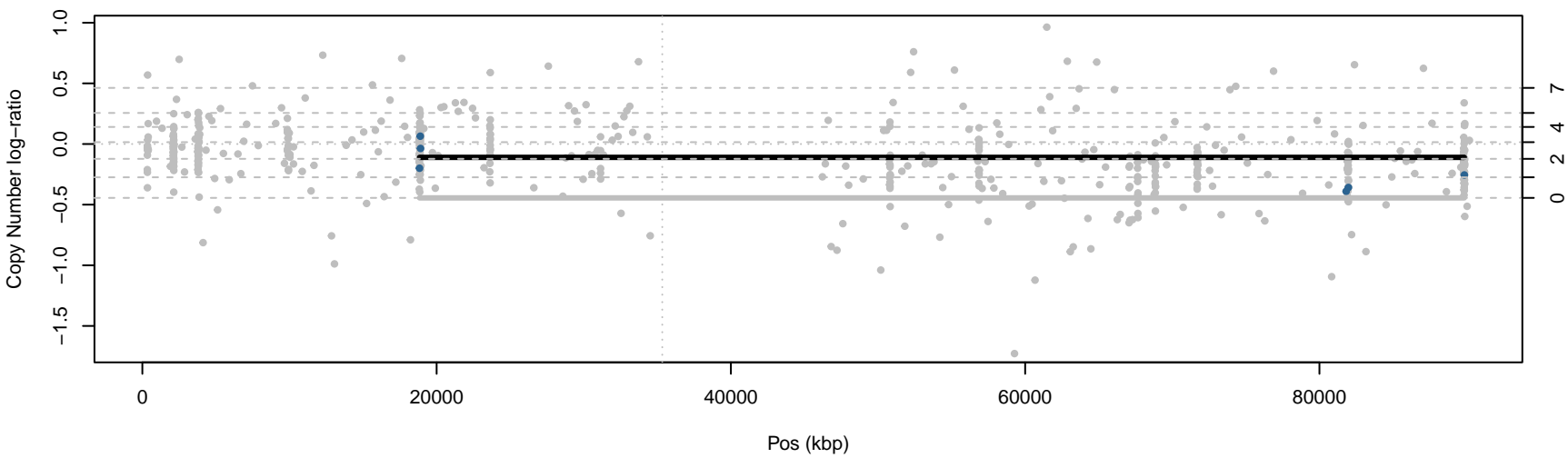
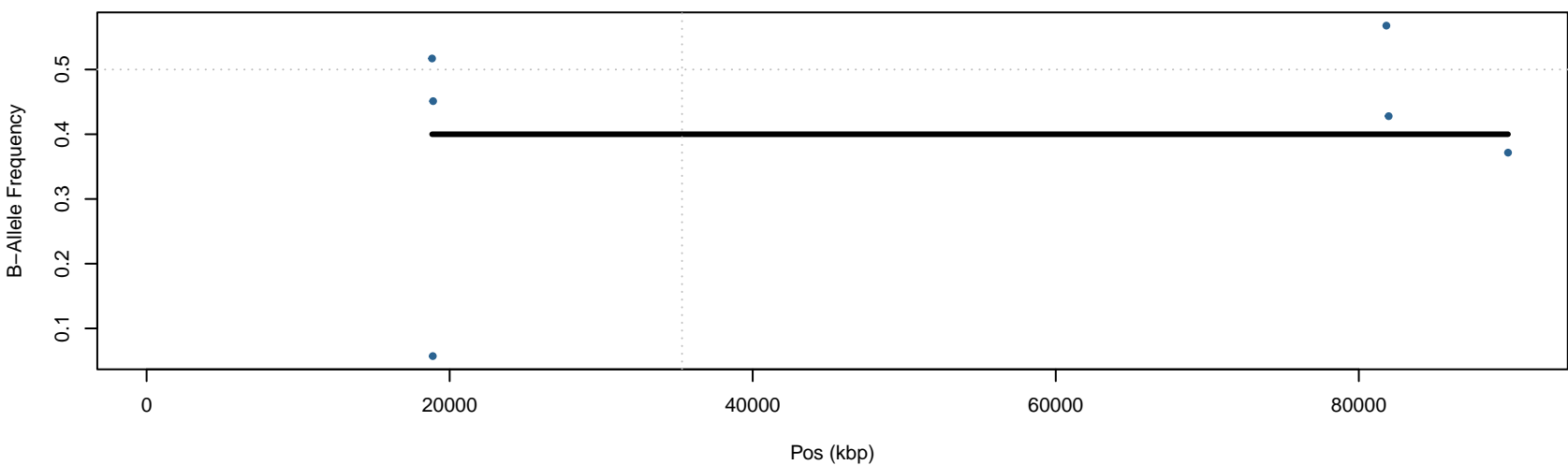
Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 12



Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 13

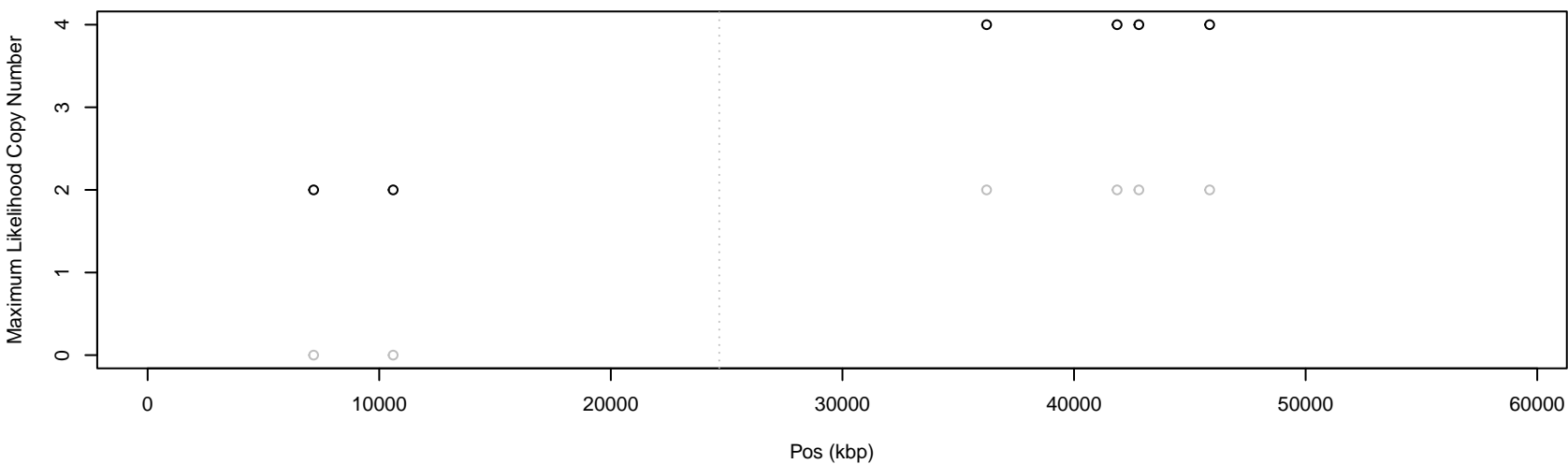
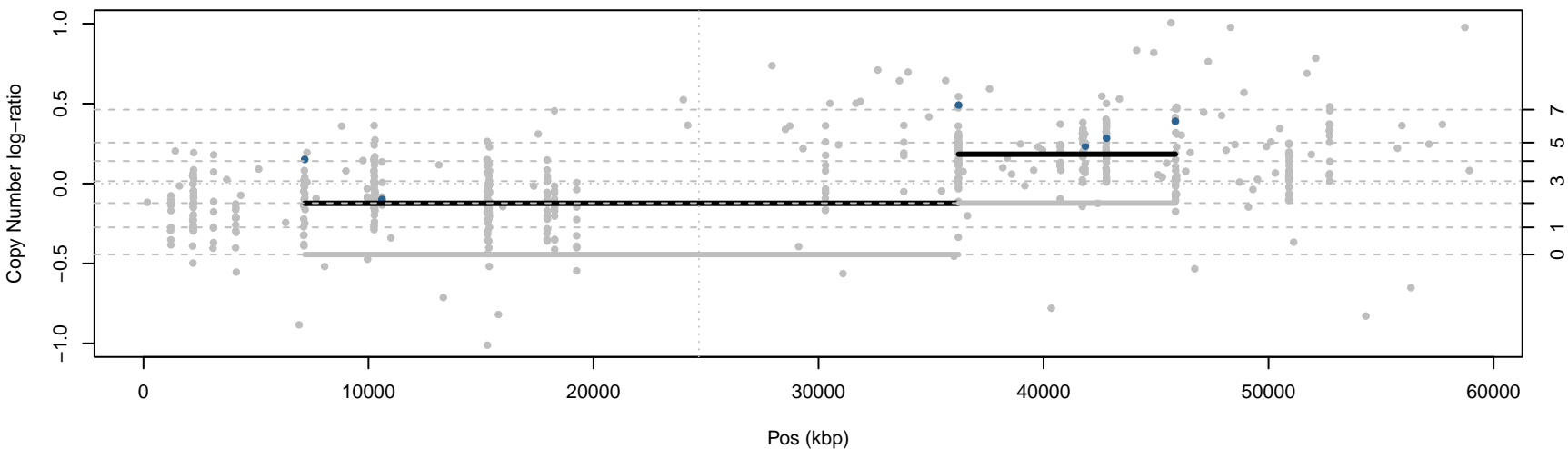
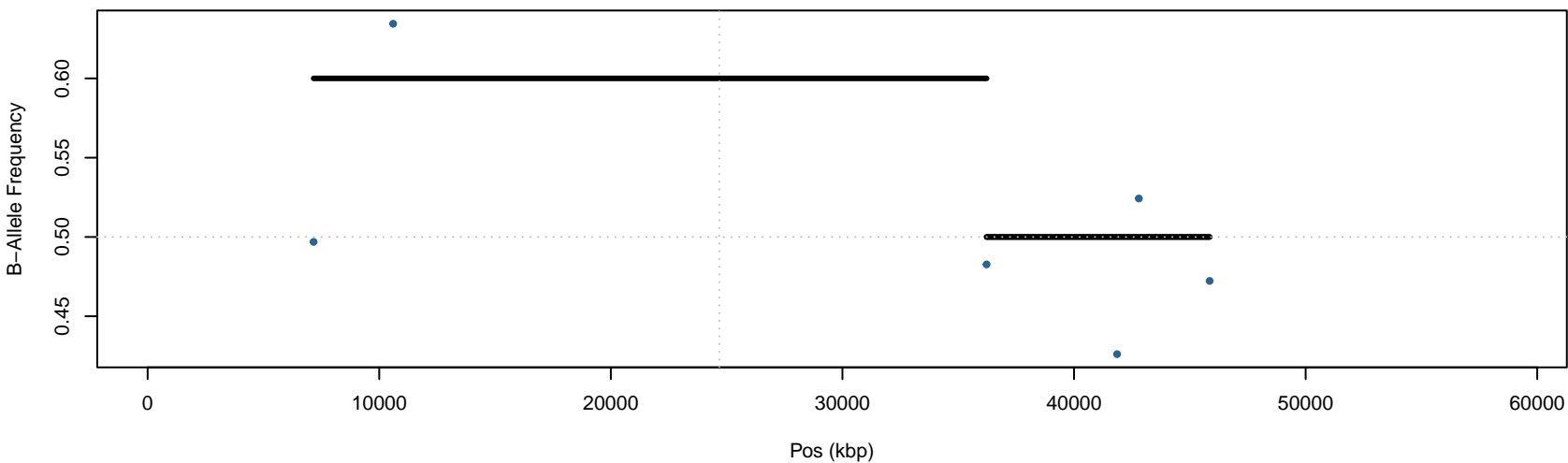


Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 16

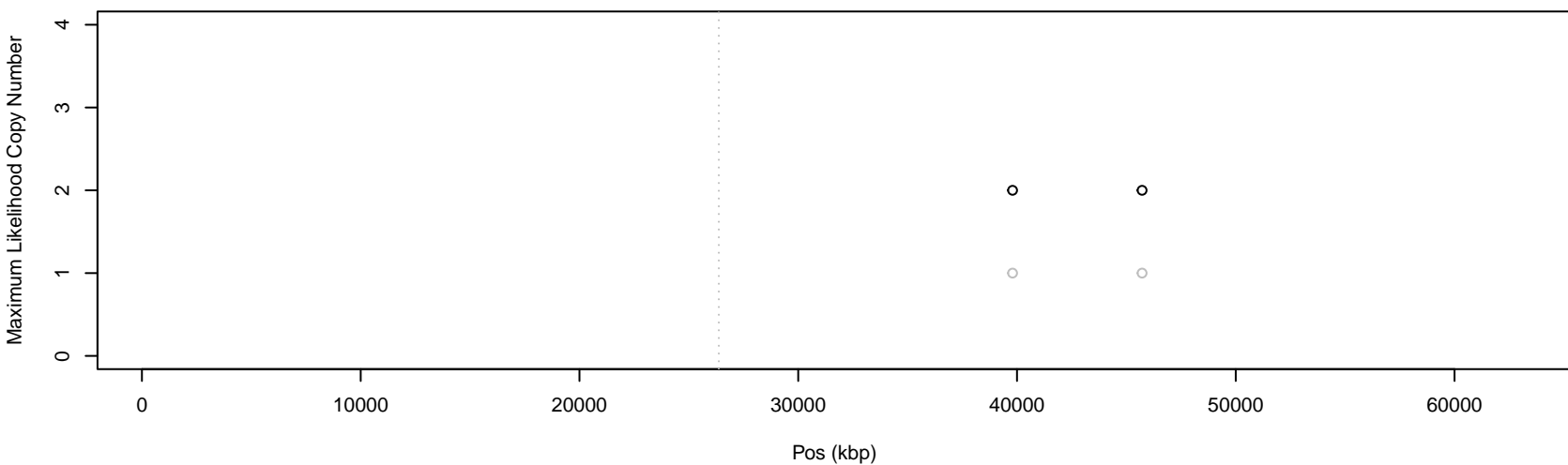
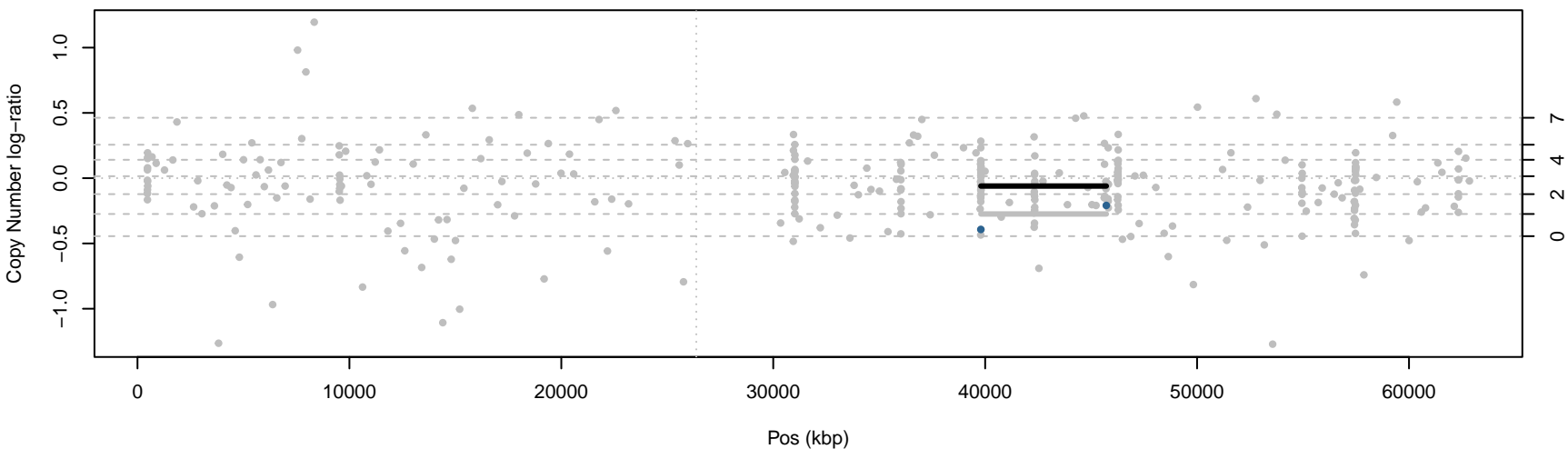
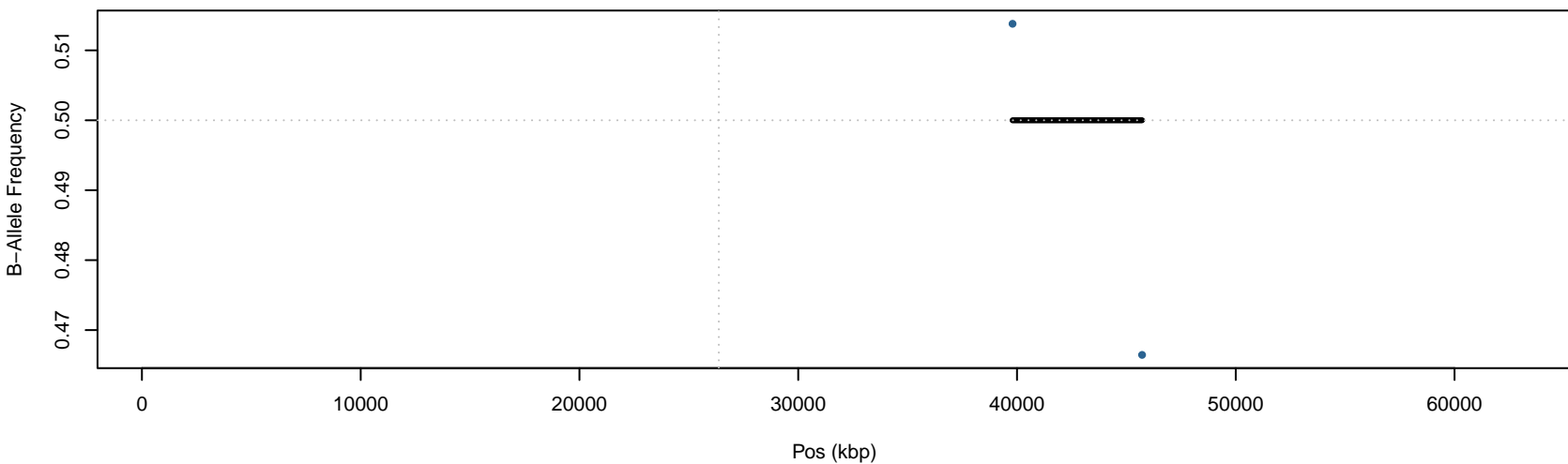




Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 19



Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 20



Purity: 0.2 Tumor ploidy: 2.885 SNV log-likelihood: -5.49 GoF: 95.8% Mean coverage: 545;642 Chromosome: 22

