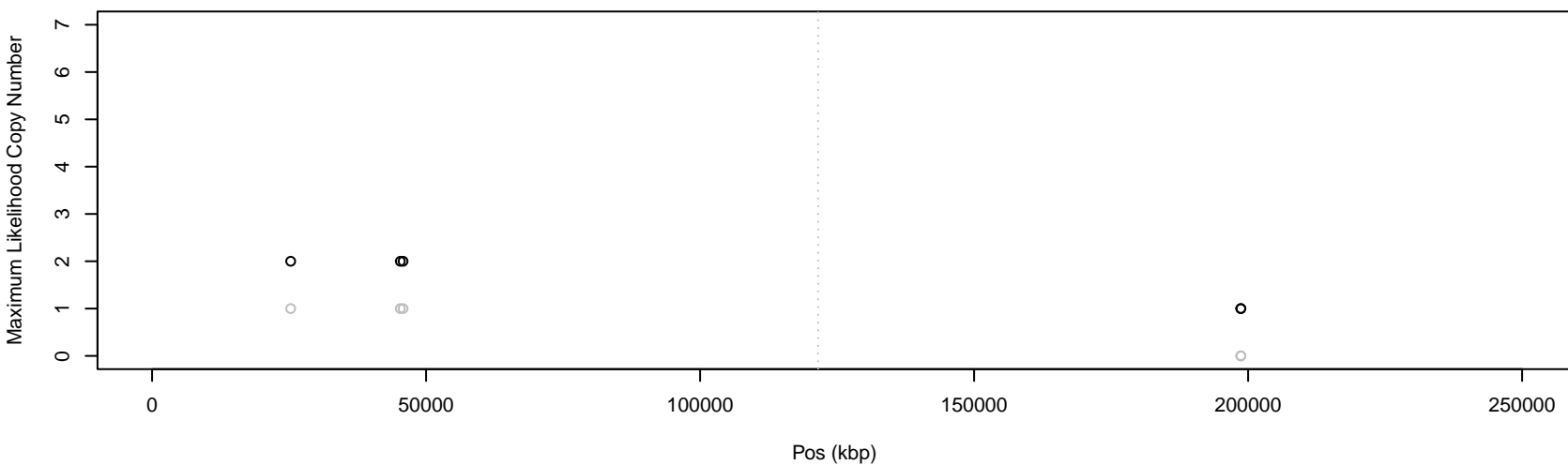
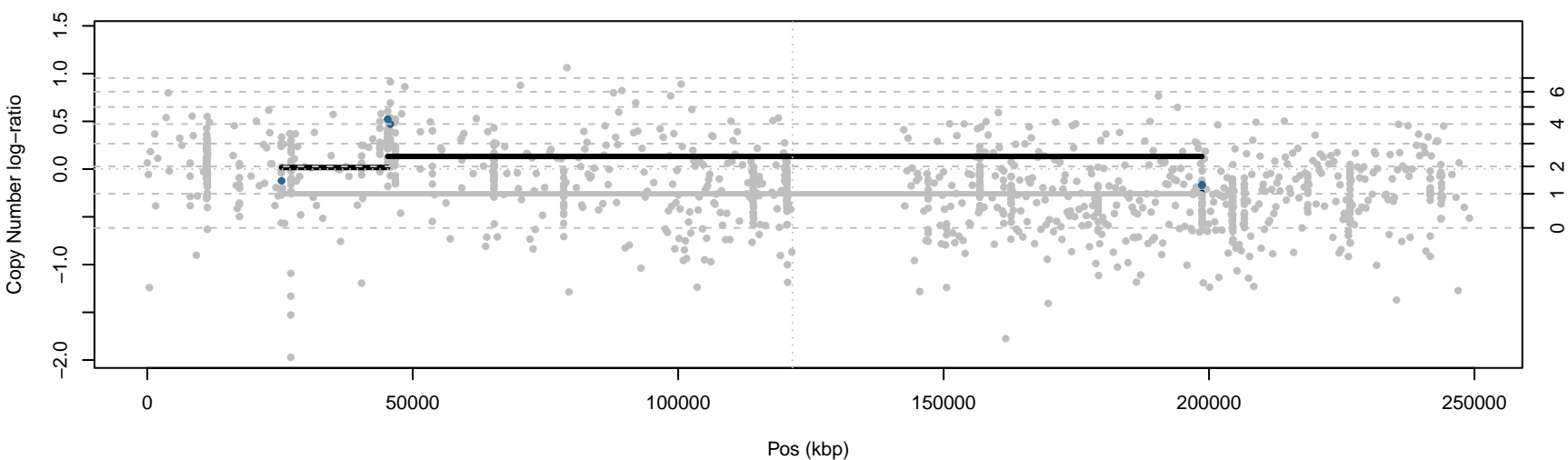
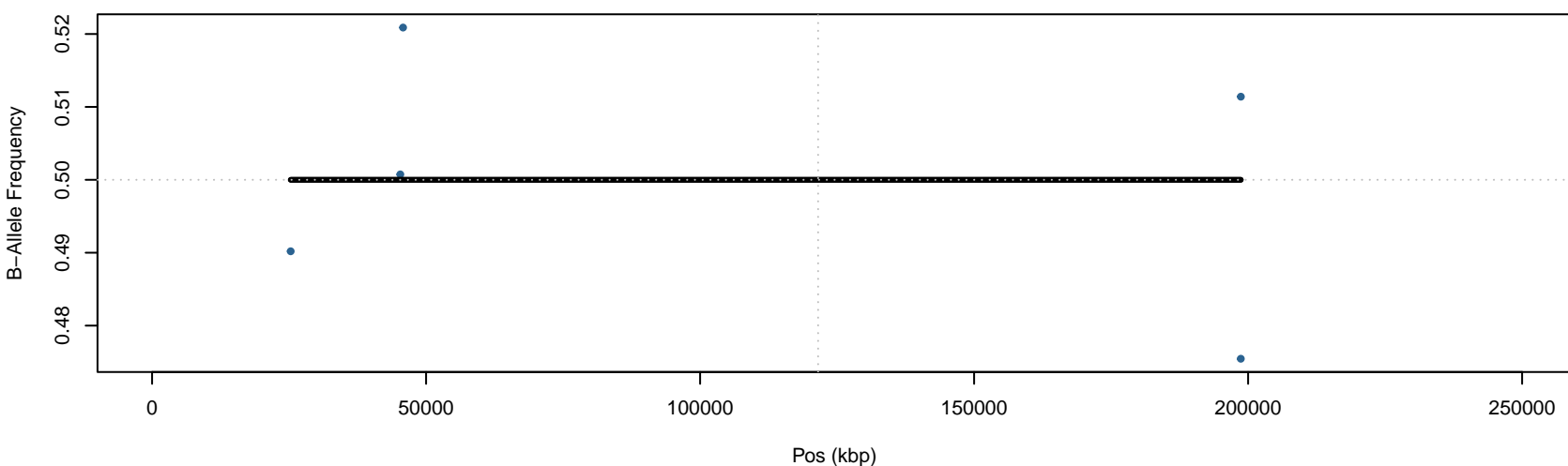
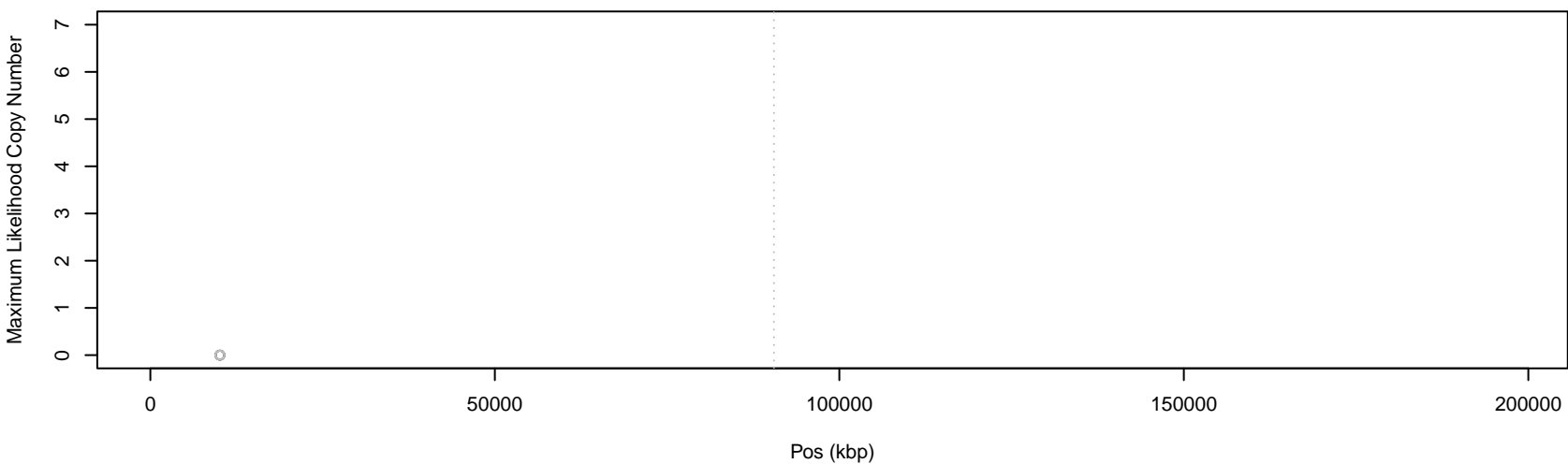
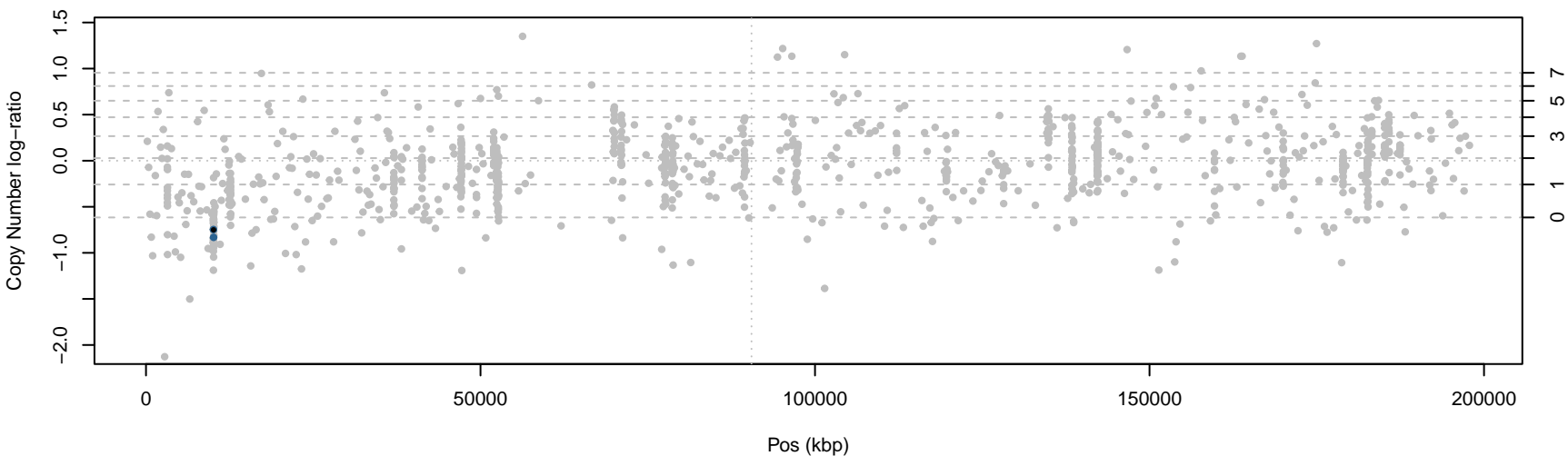
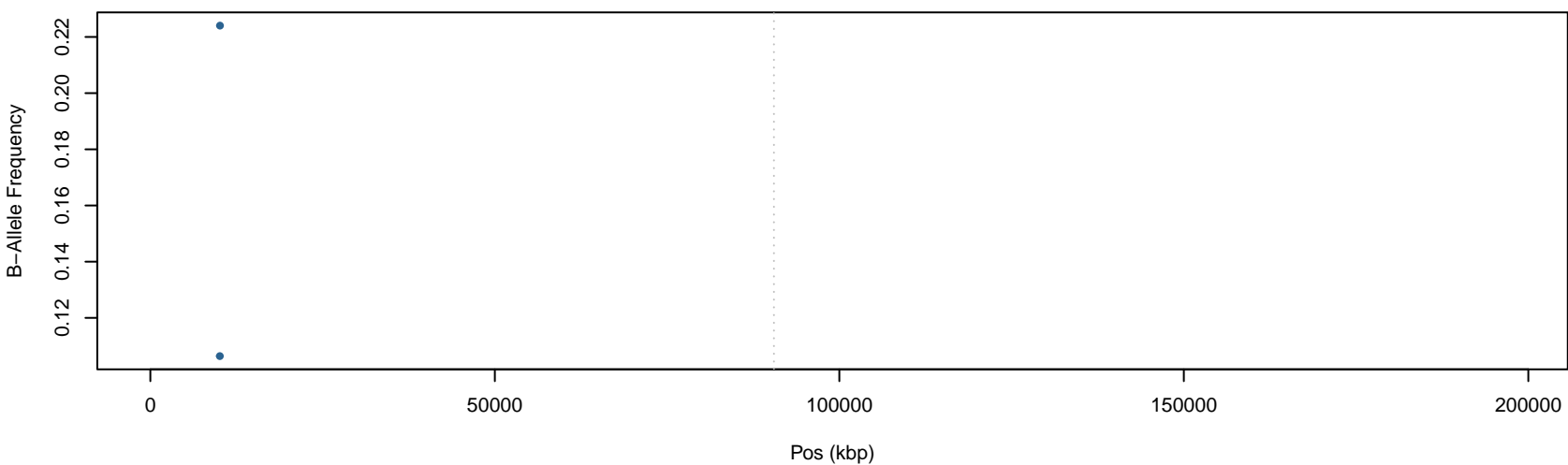


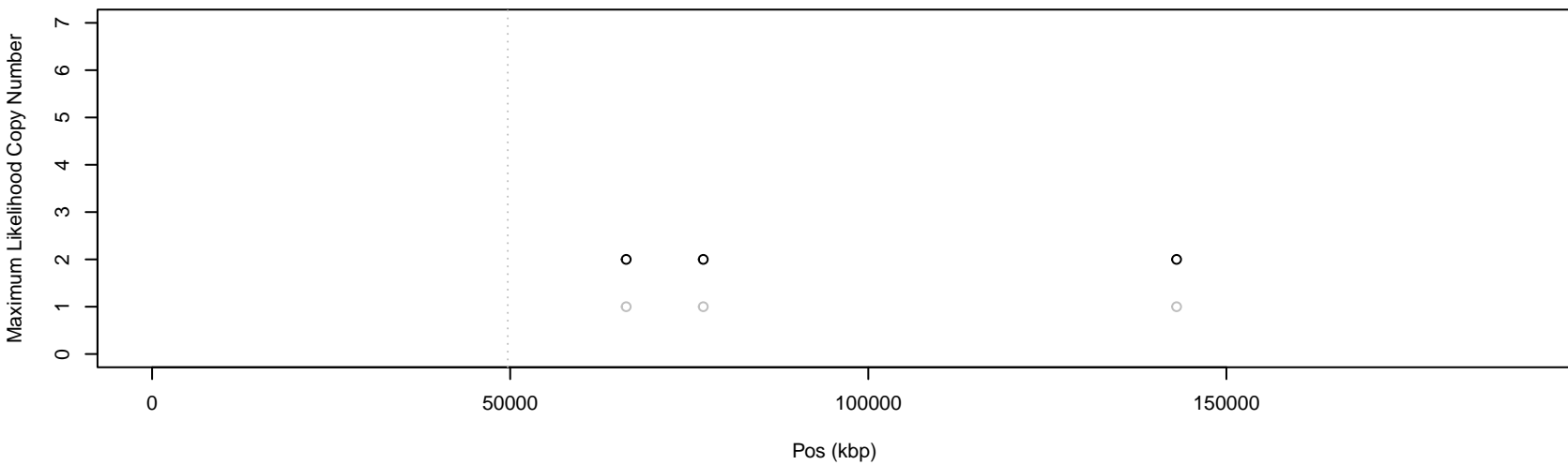
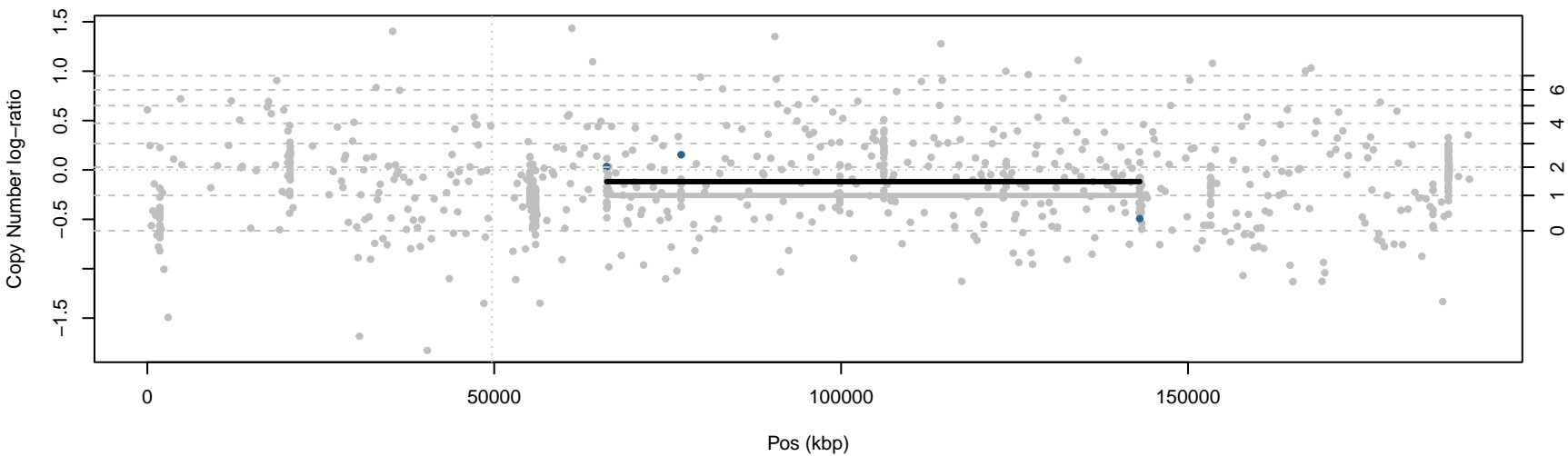
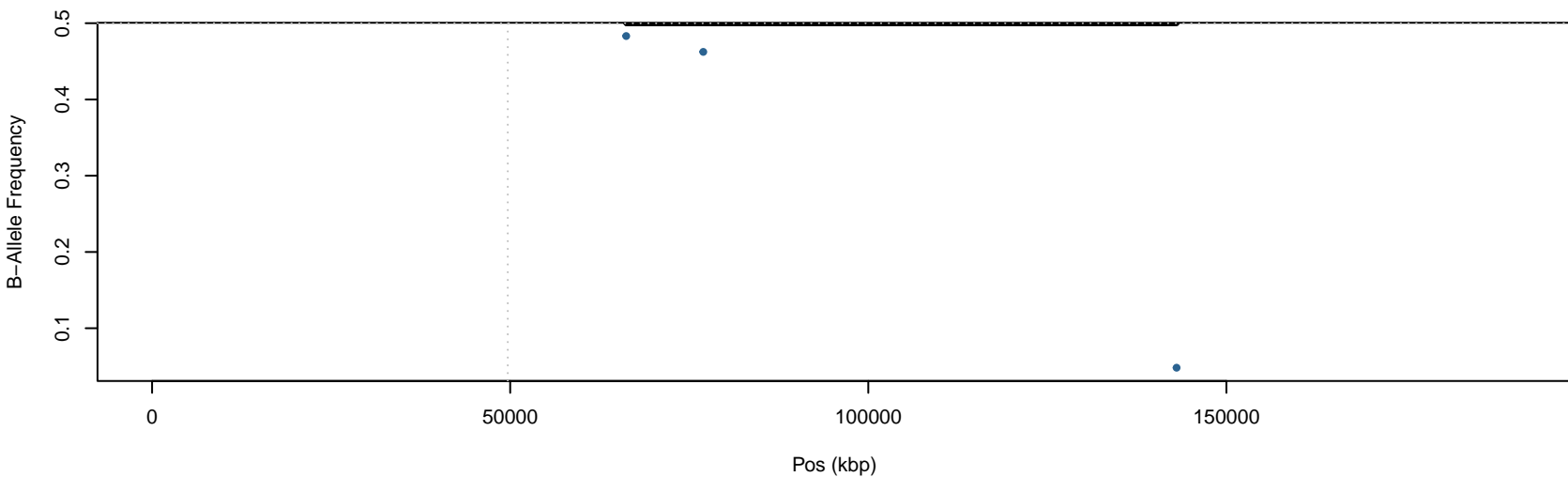
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 1



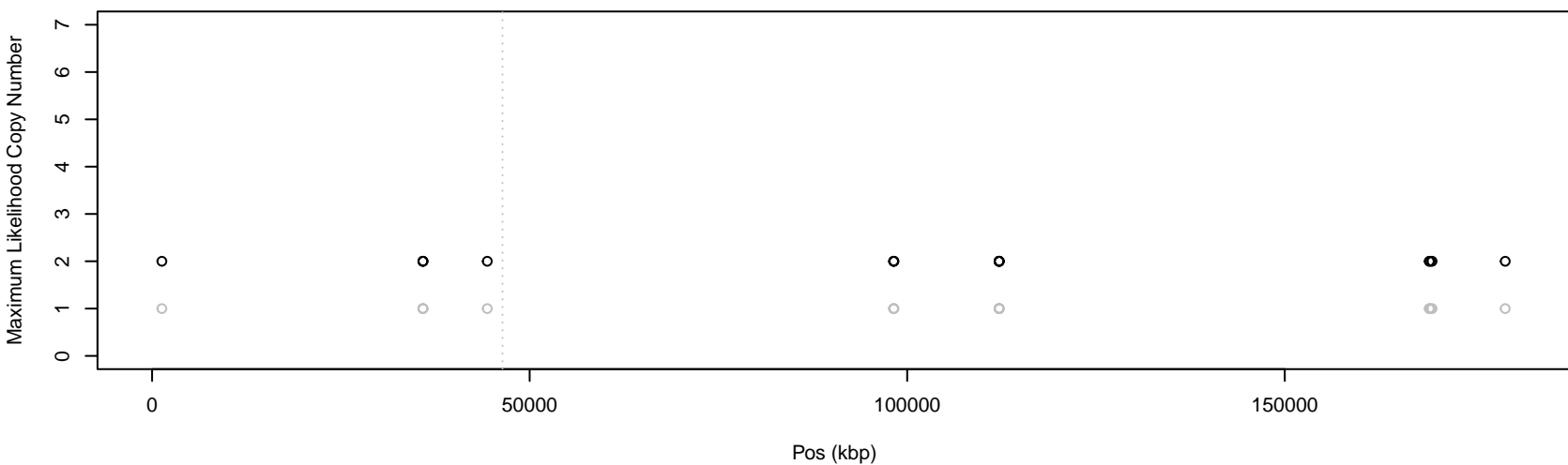
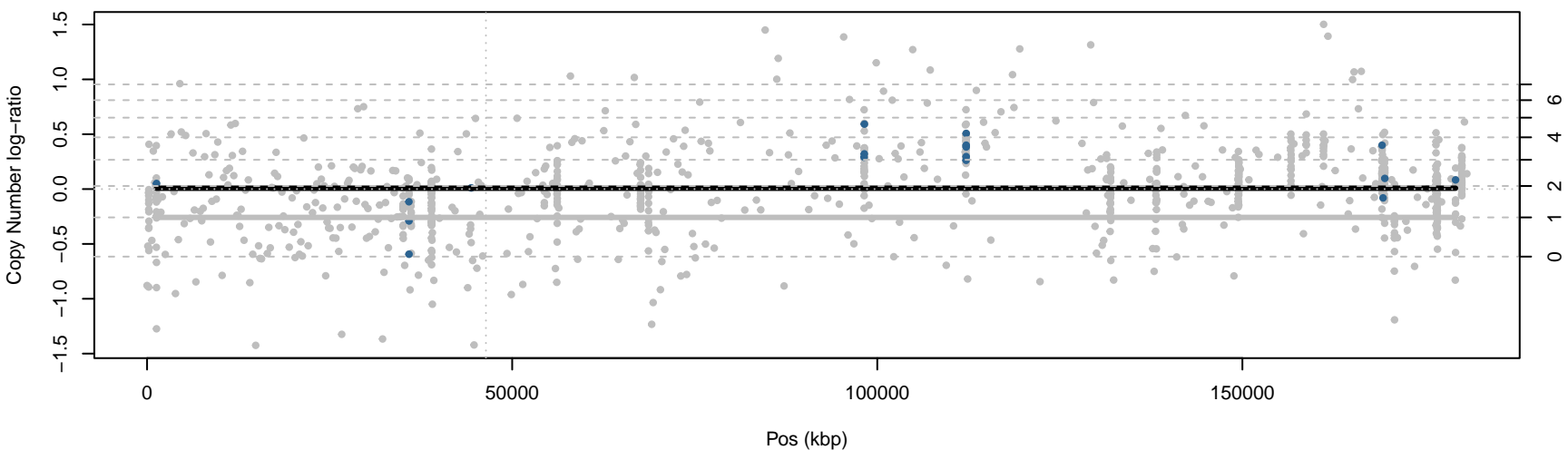
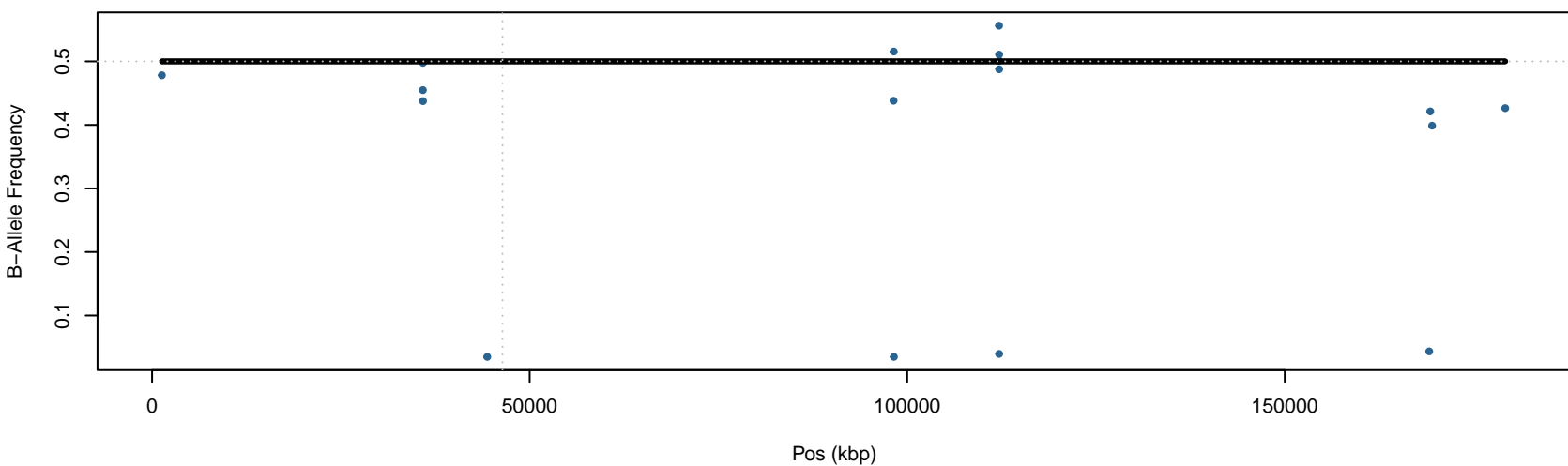
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 3



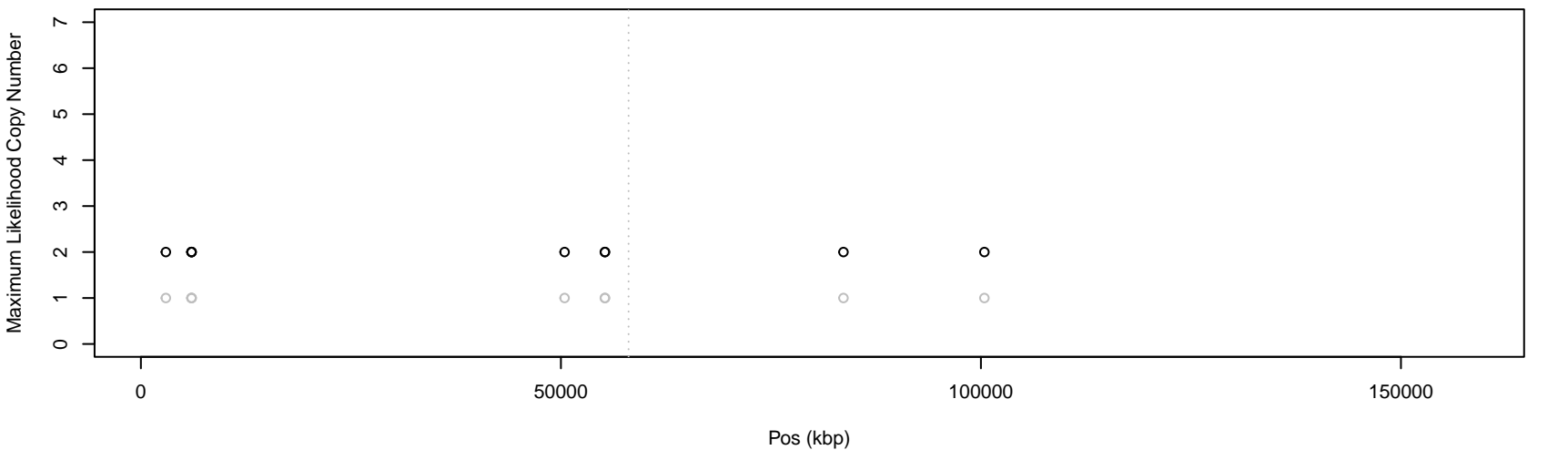
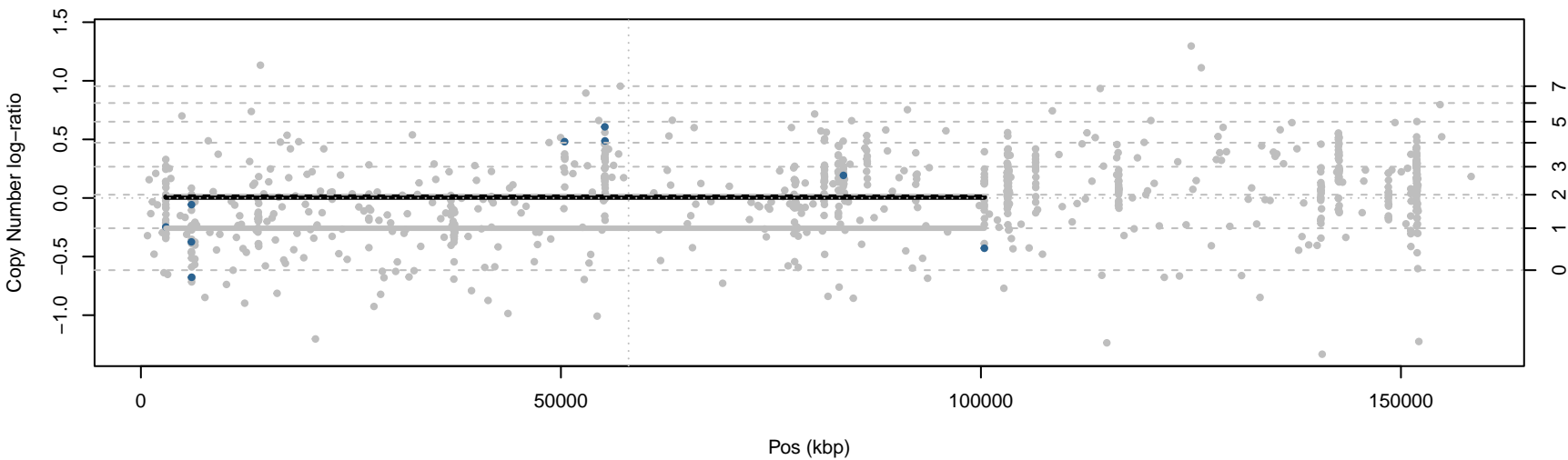
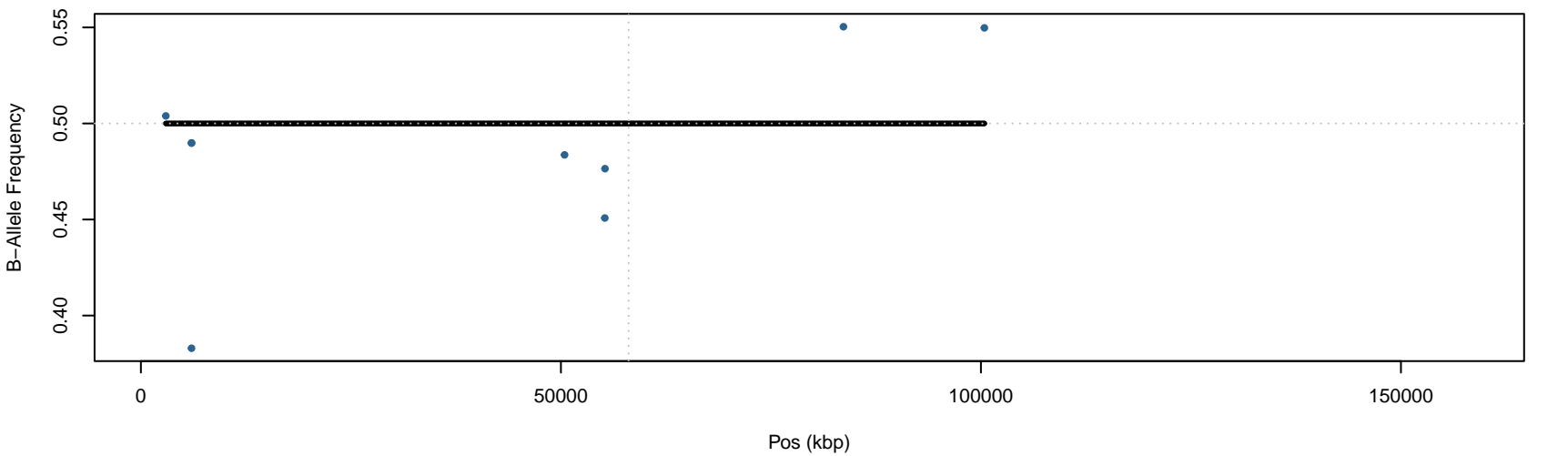
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 4



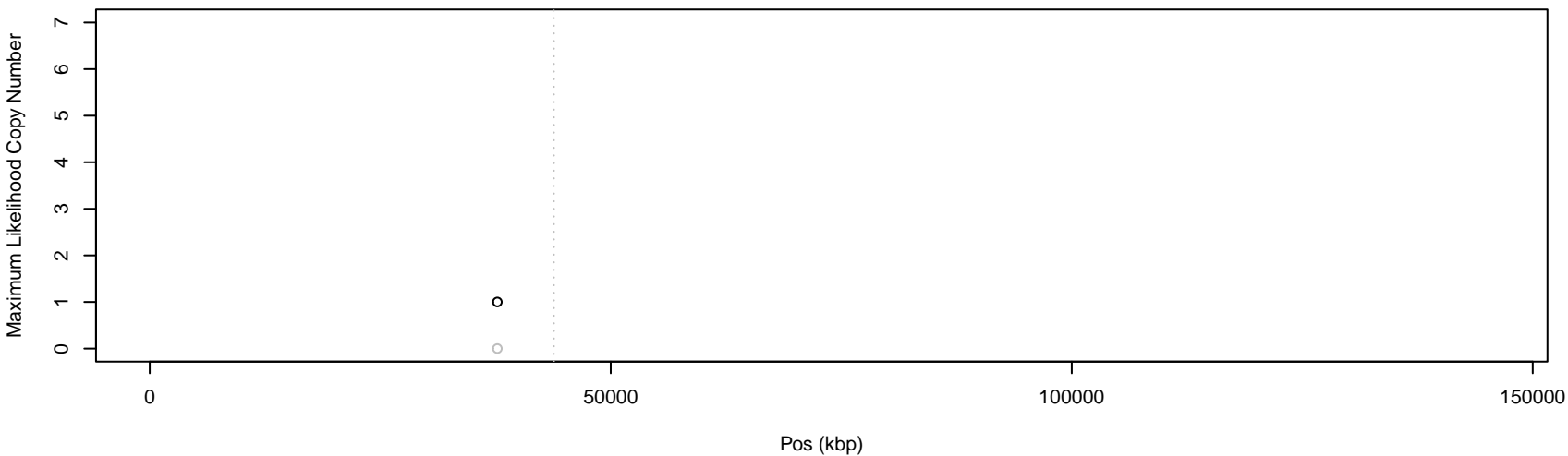
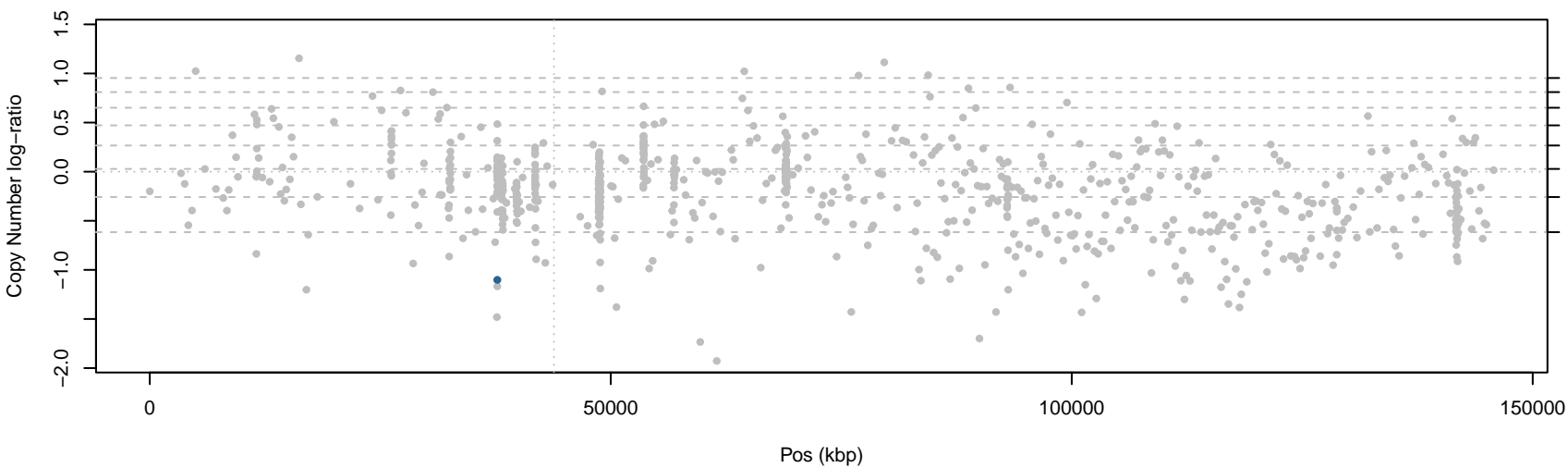
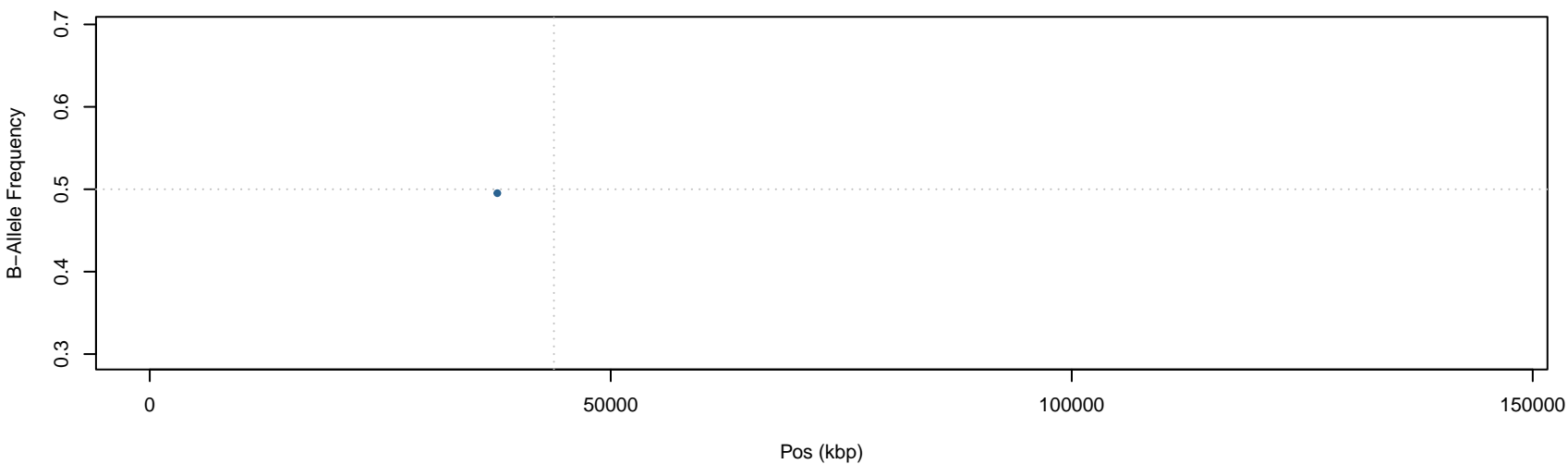
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 5



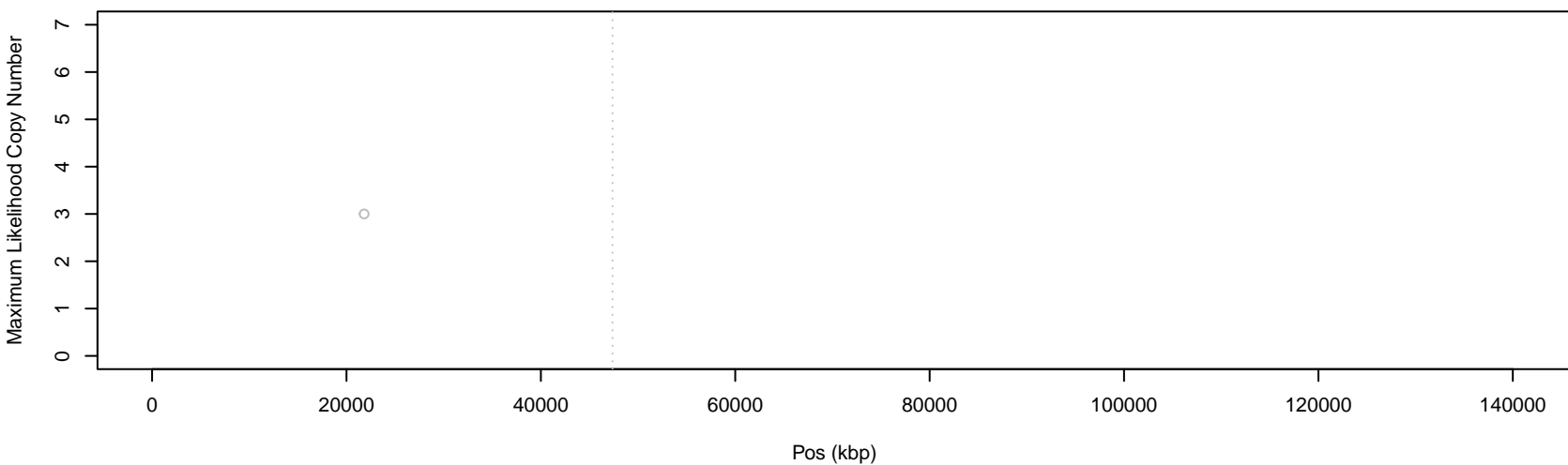
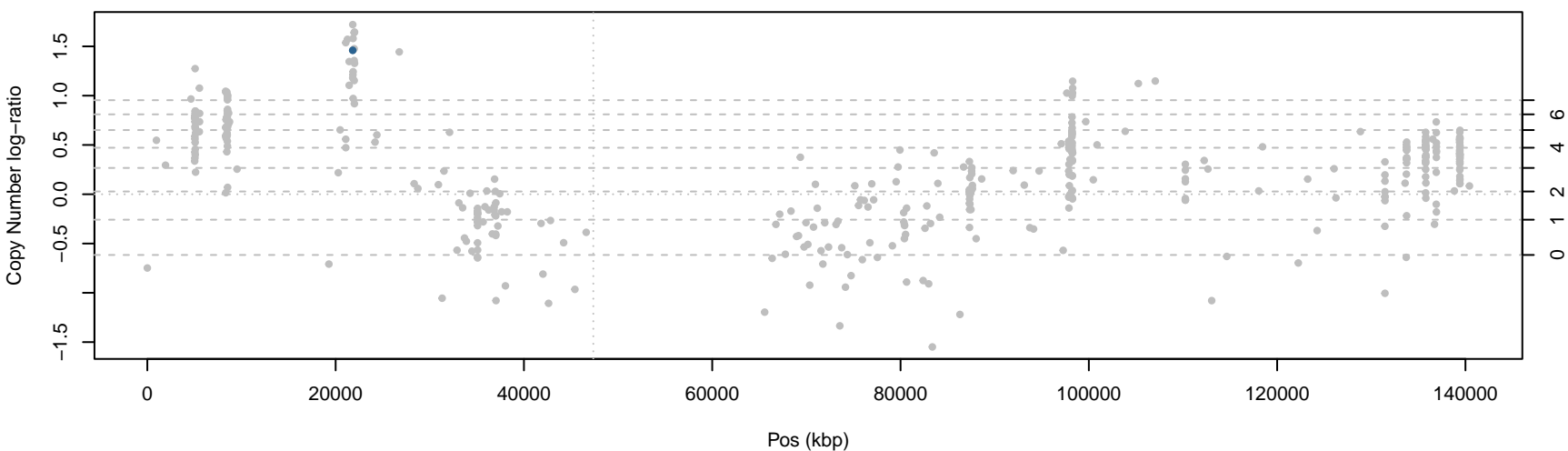
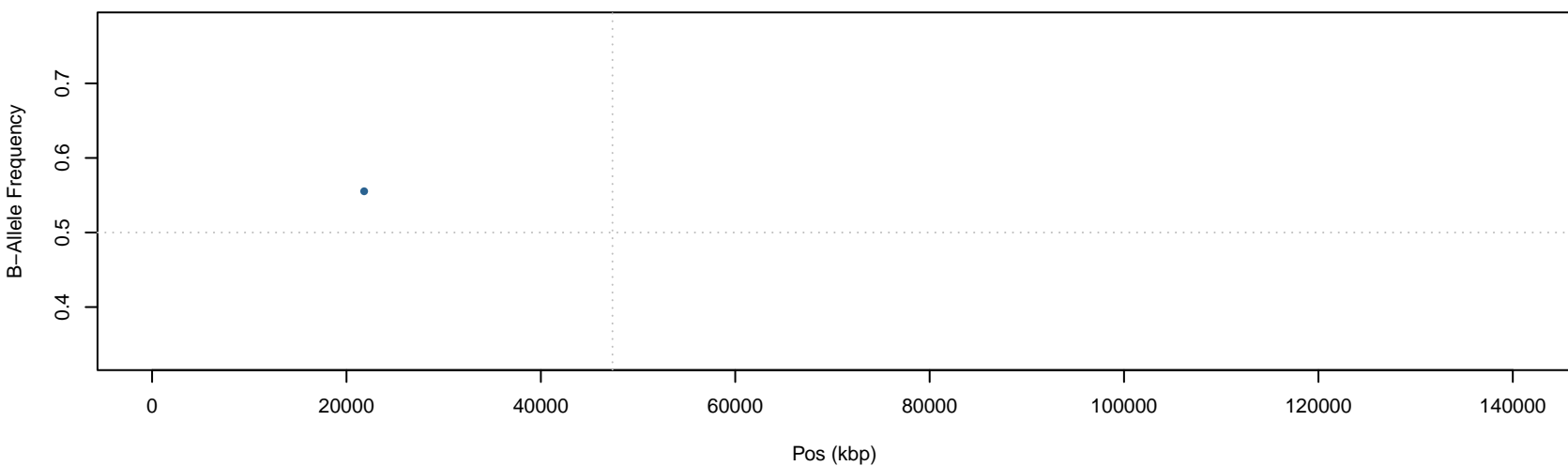
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 7



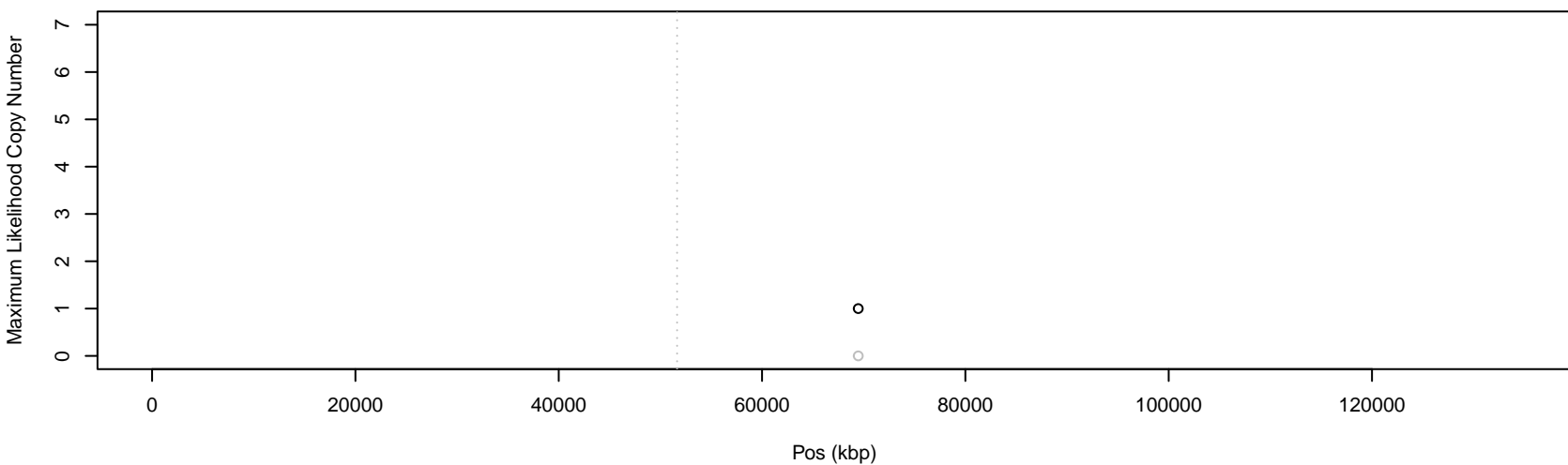
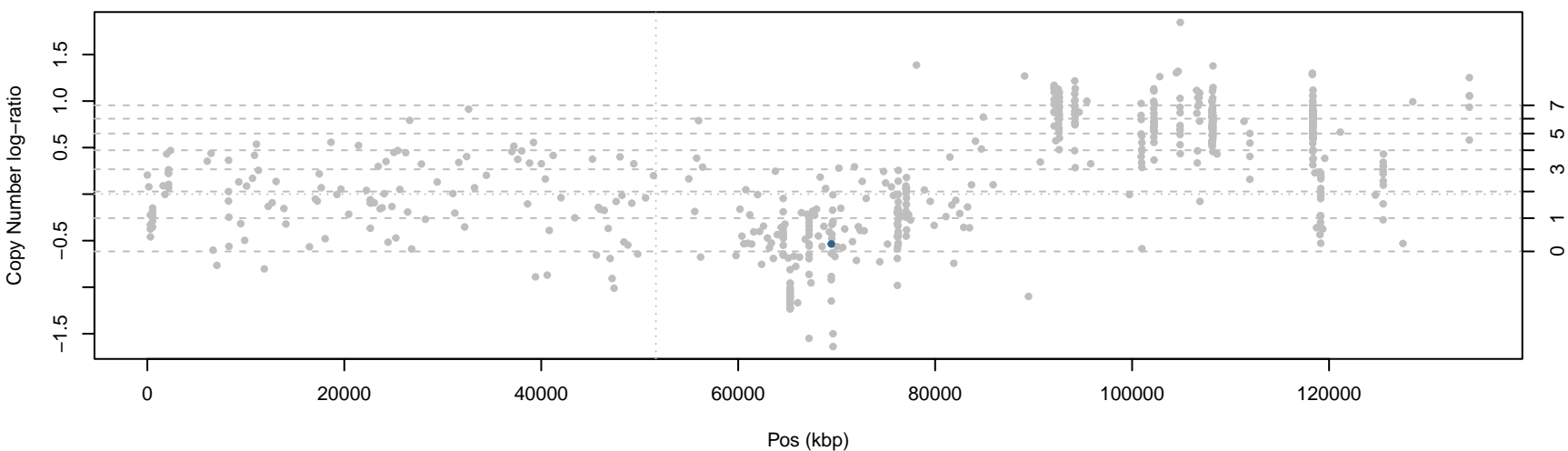
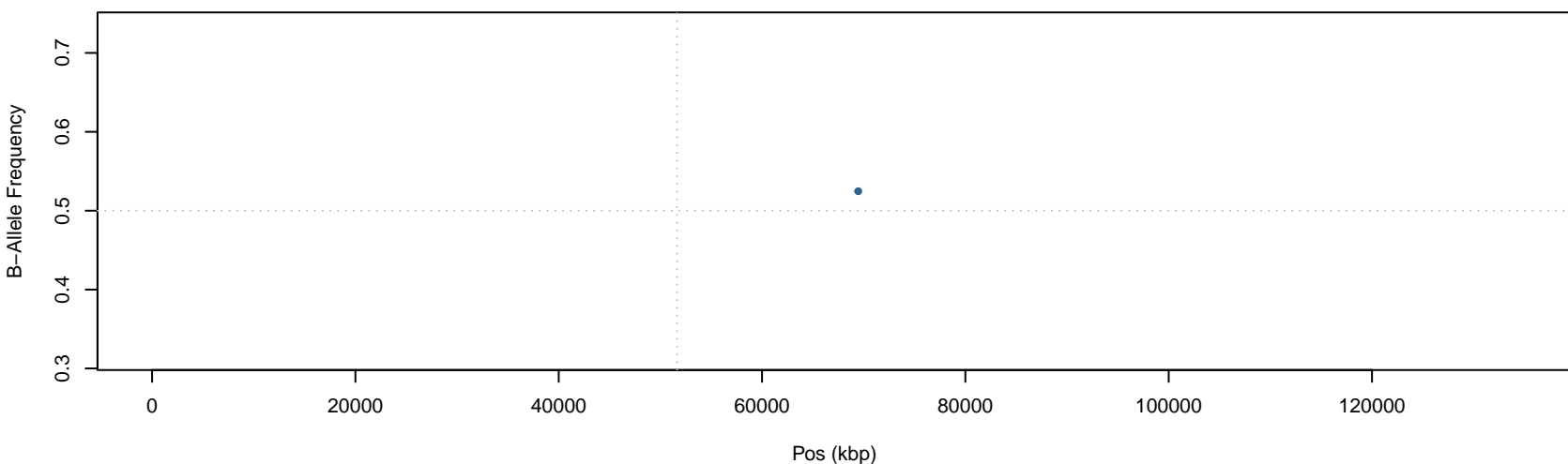
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 8



Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 9

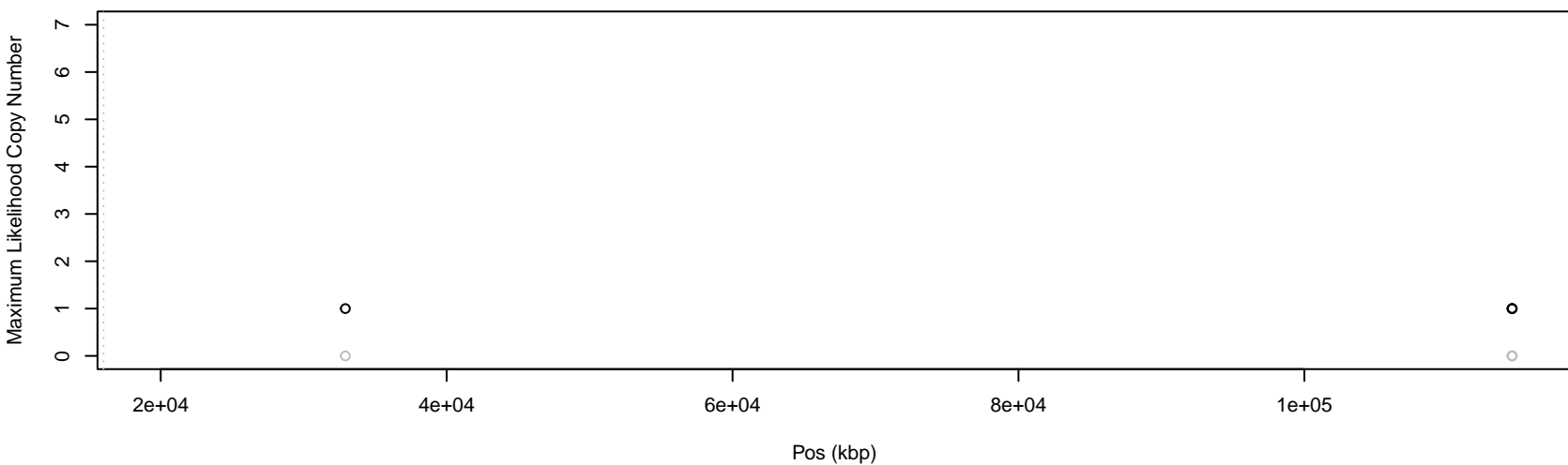
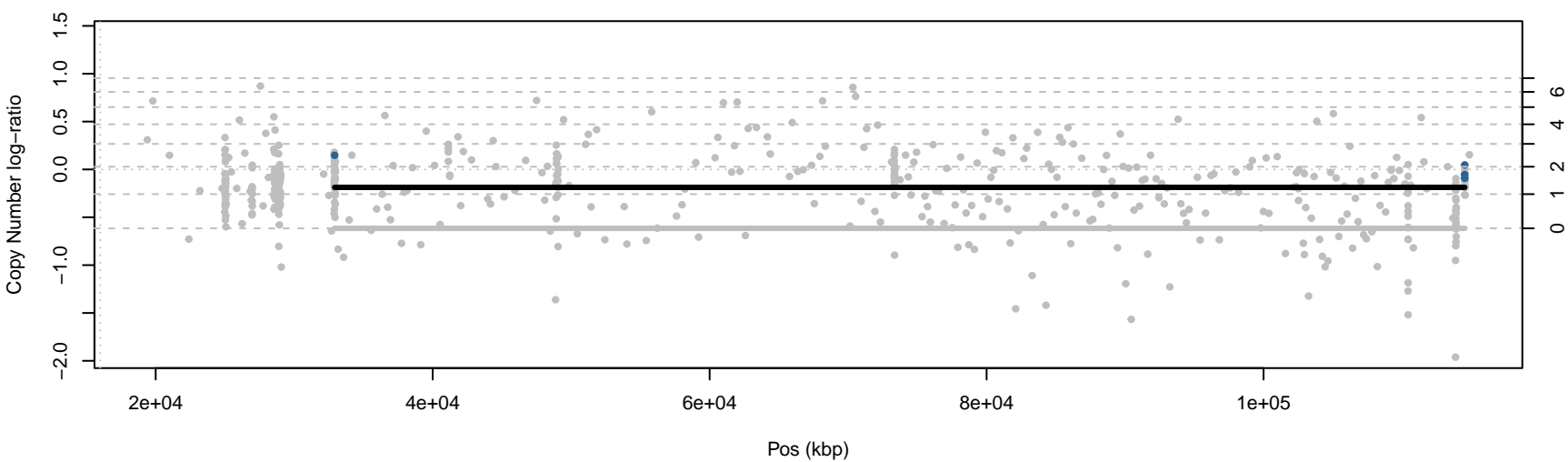
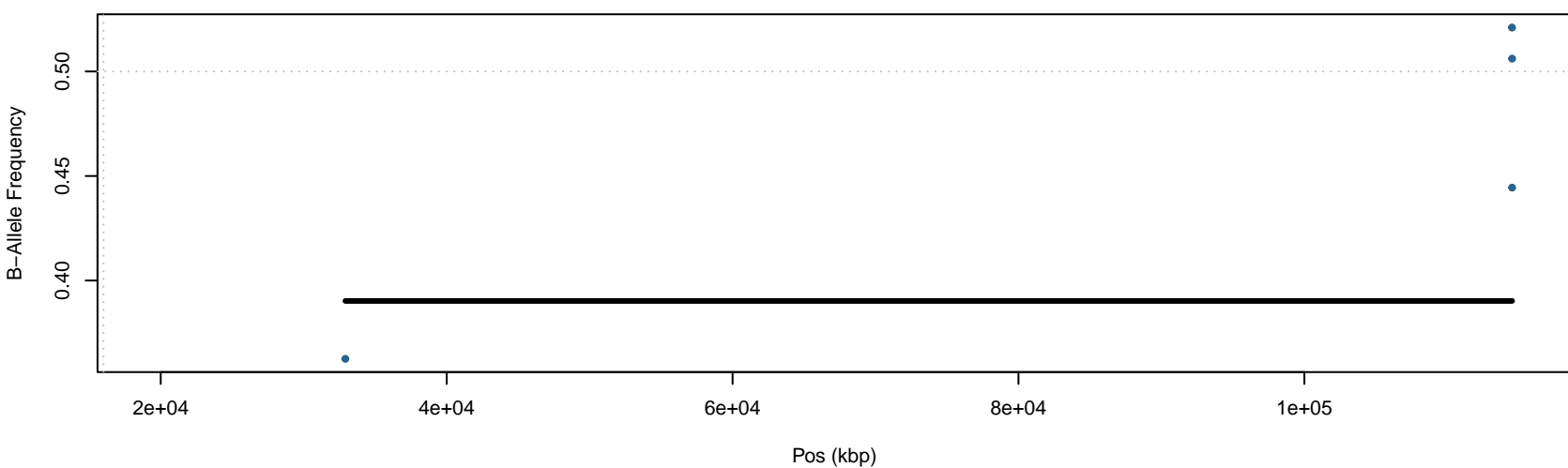


Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 11

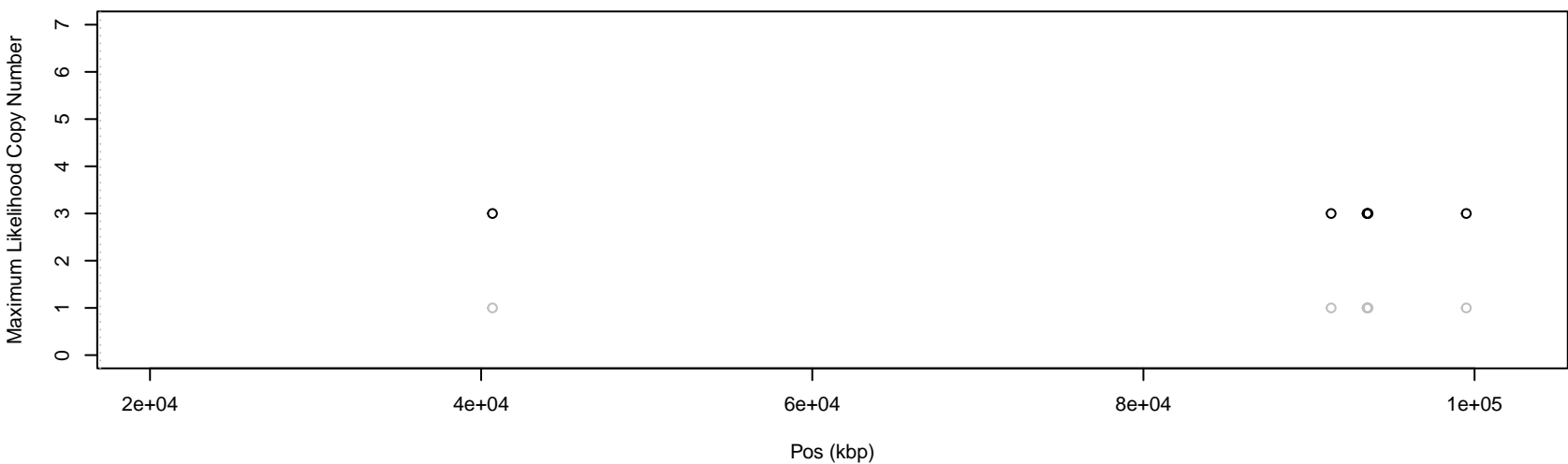
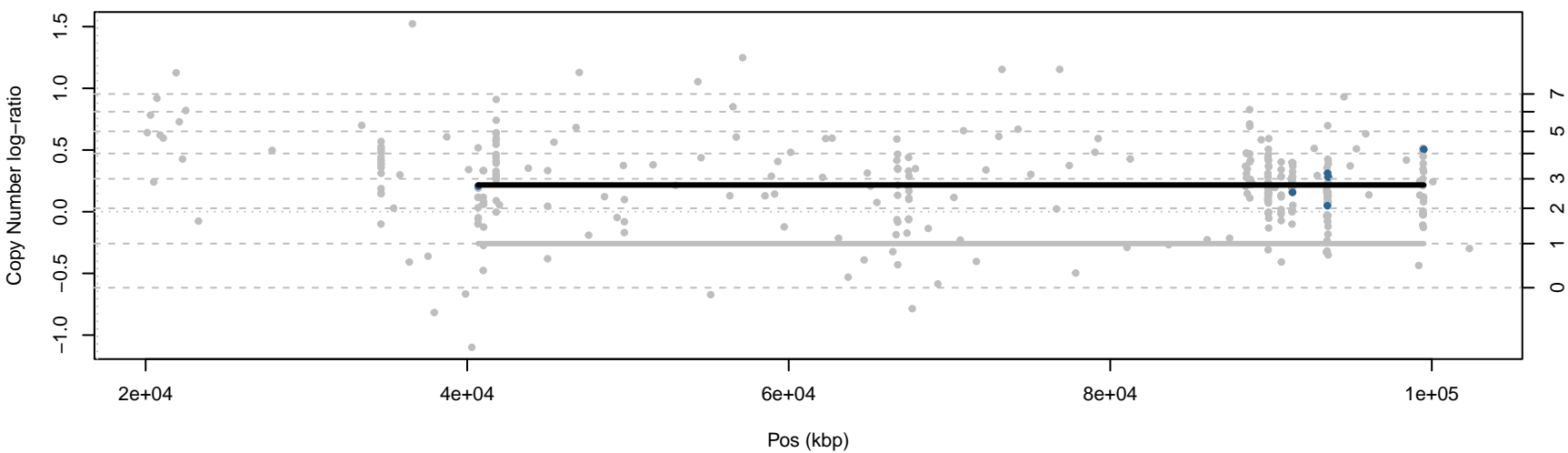
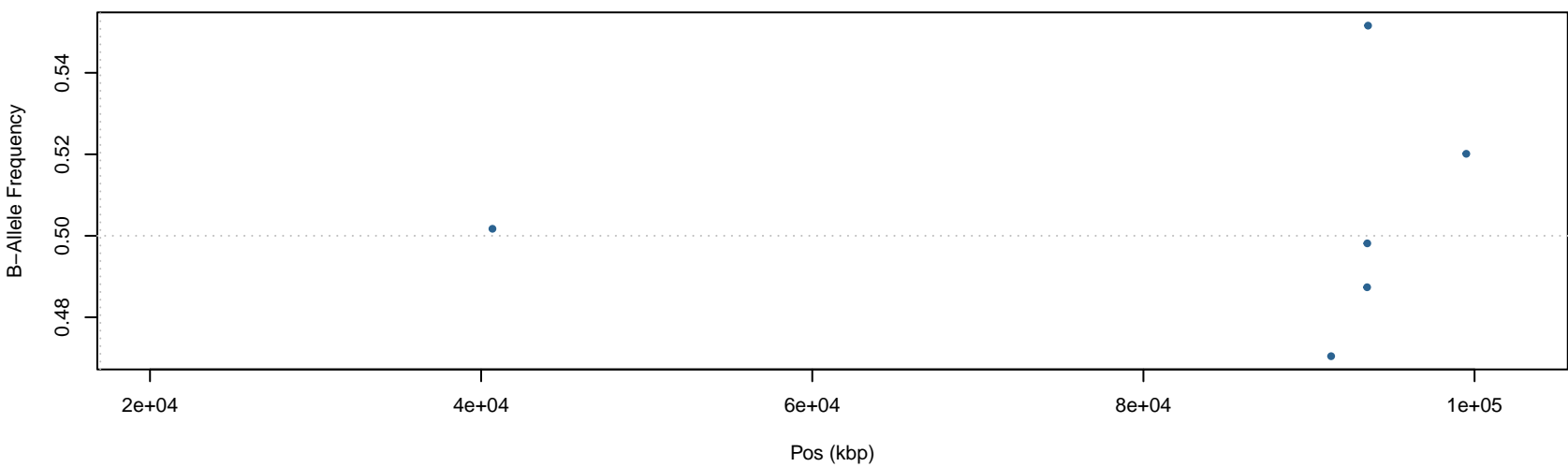




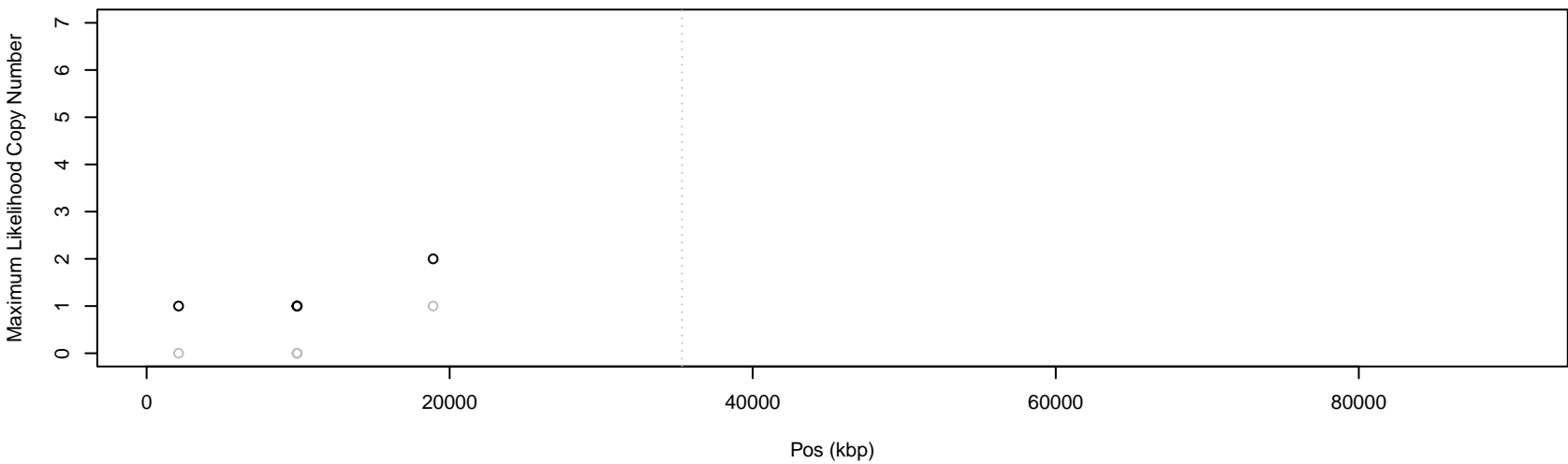
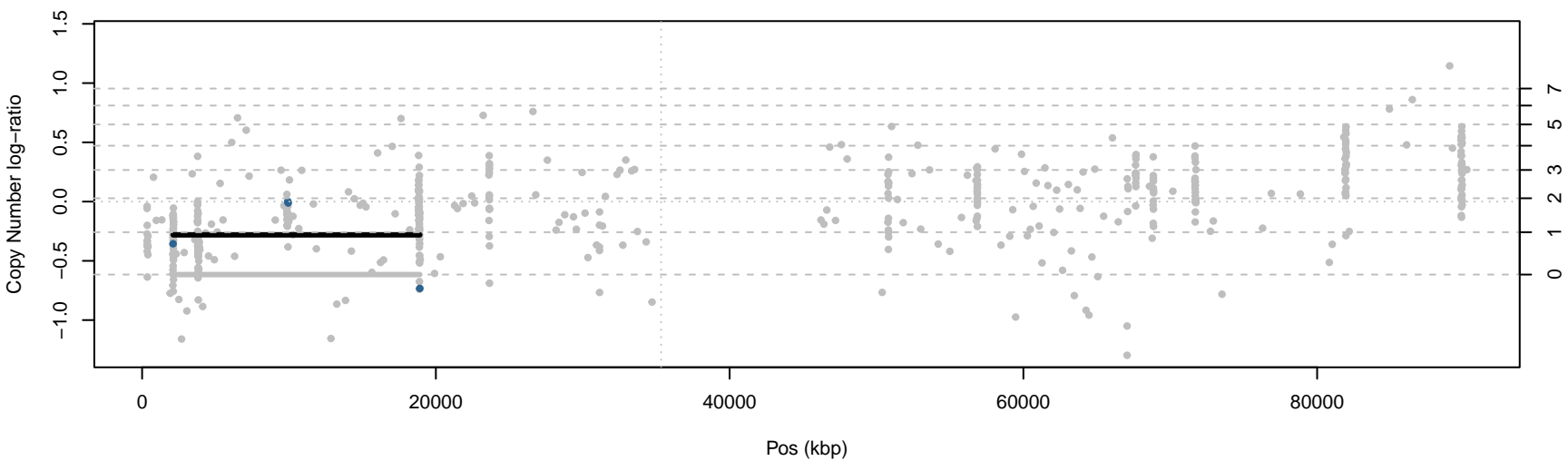
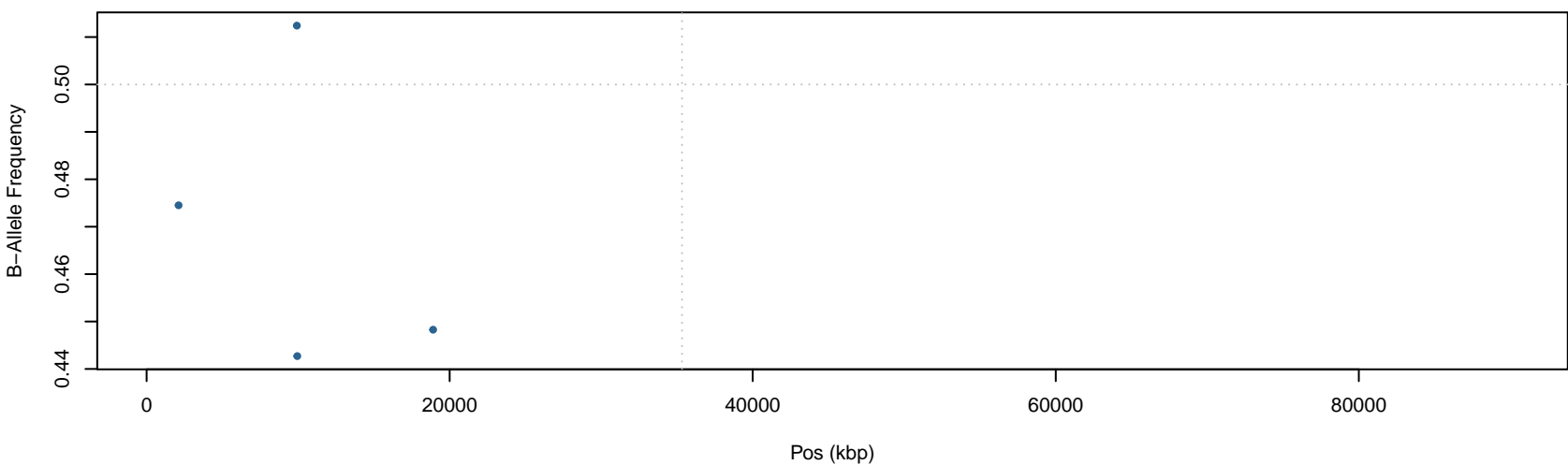
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 13



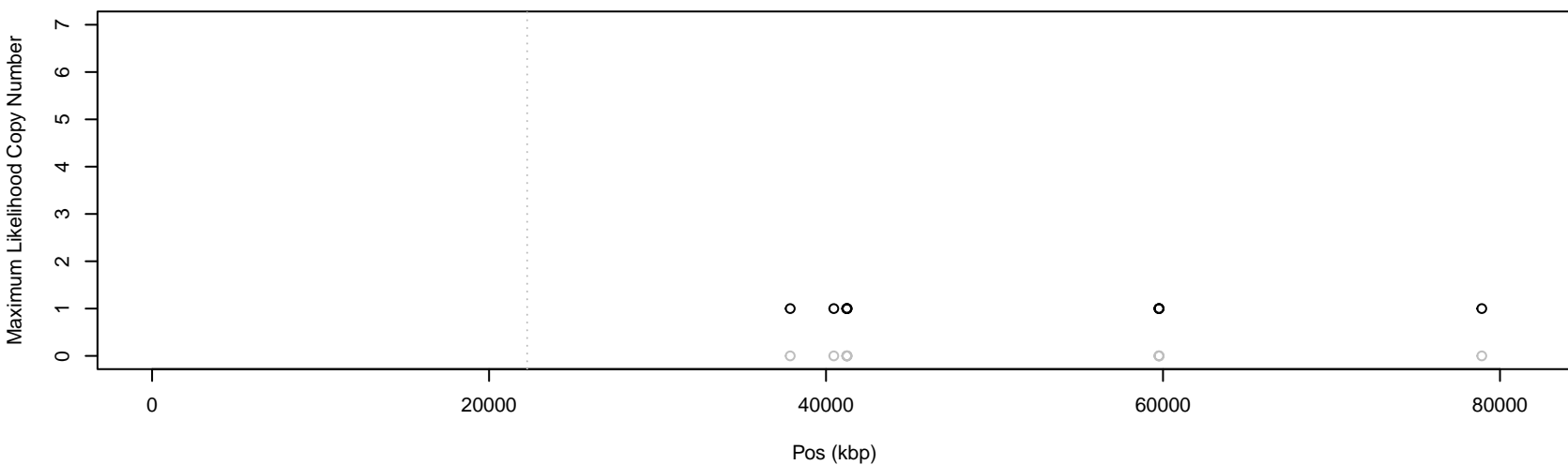
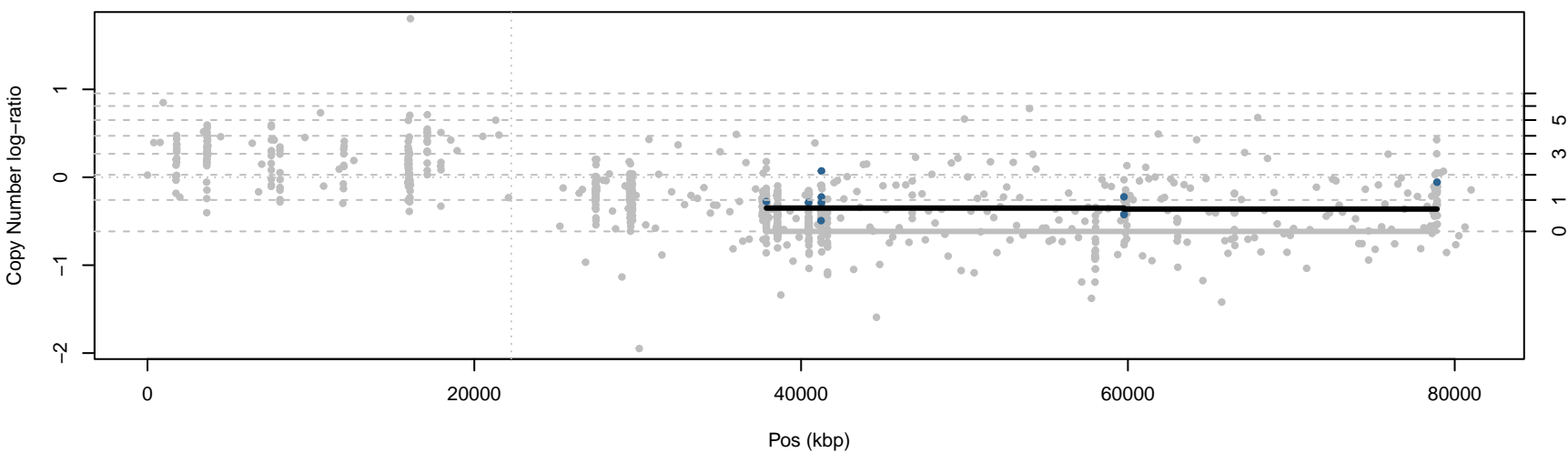
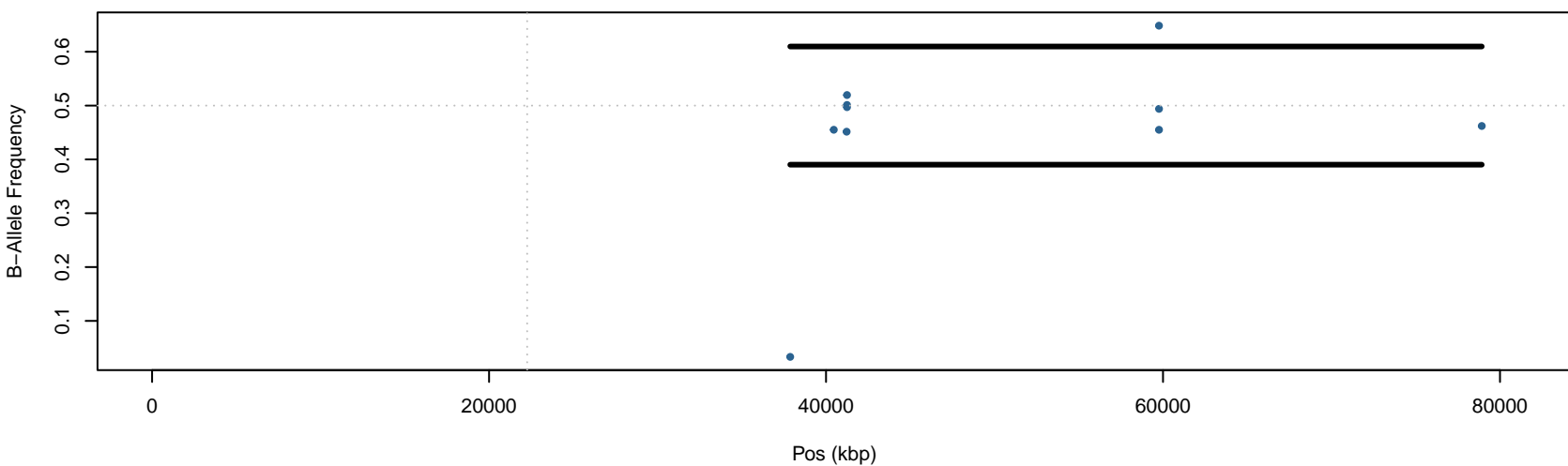
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 15



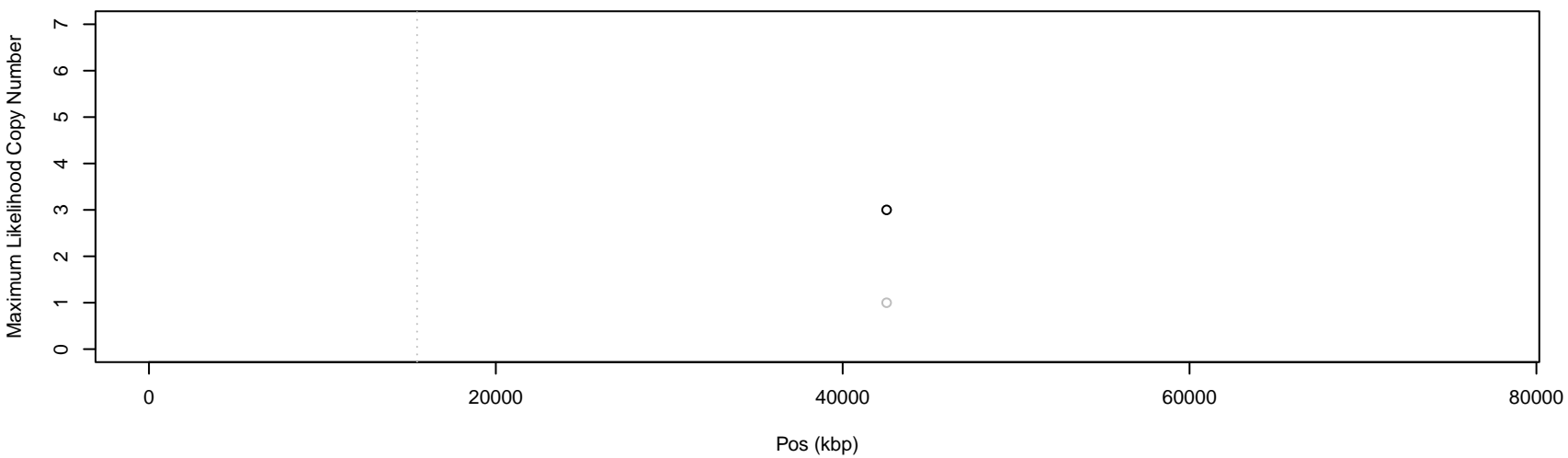
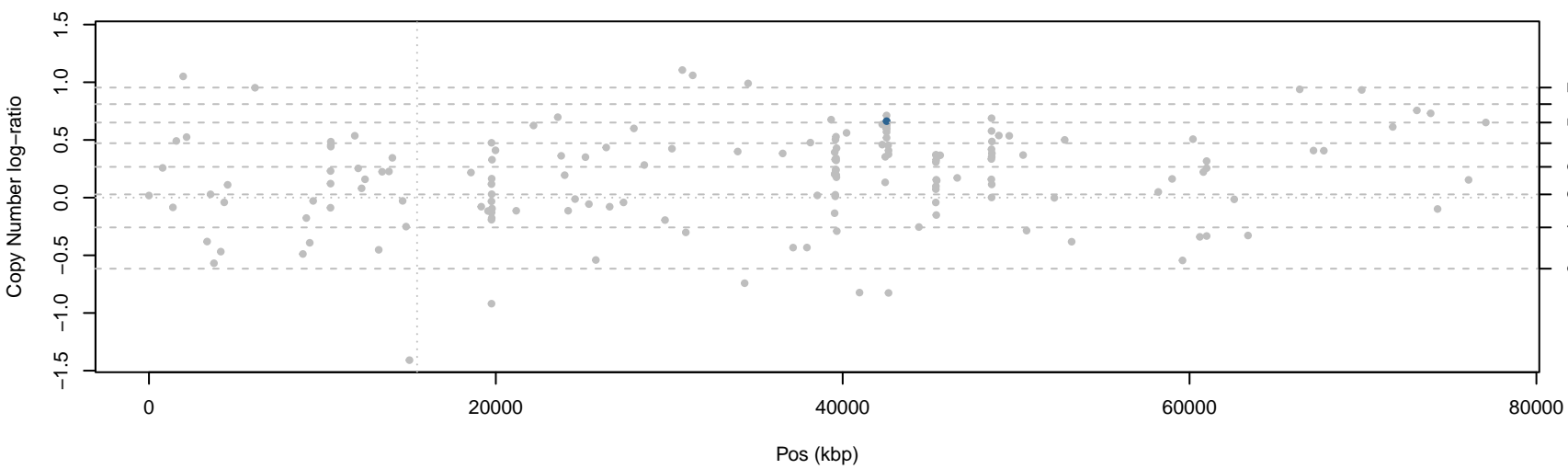
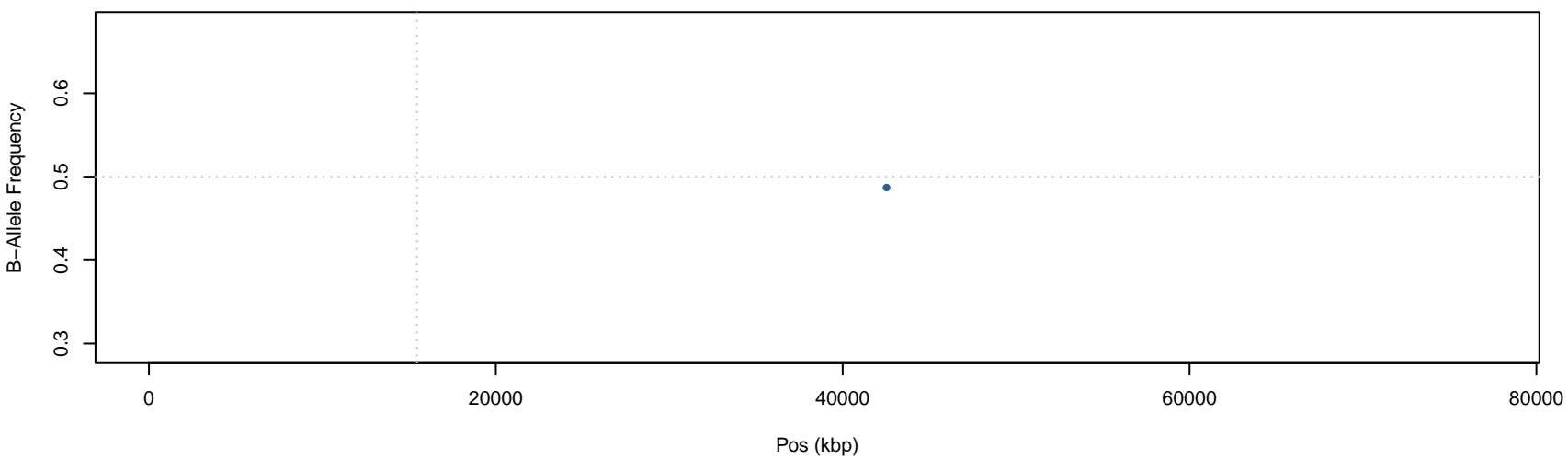
Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 16



Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 17



Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 18



Purity: 0.36 Tumor ploidy: 1.893 SNV log-likelihood: -99.21 GoF: 76.7% Mean coverage: 290;505 Chromosome: 19

