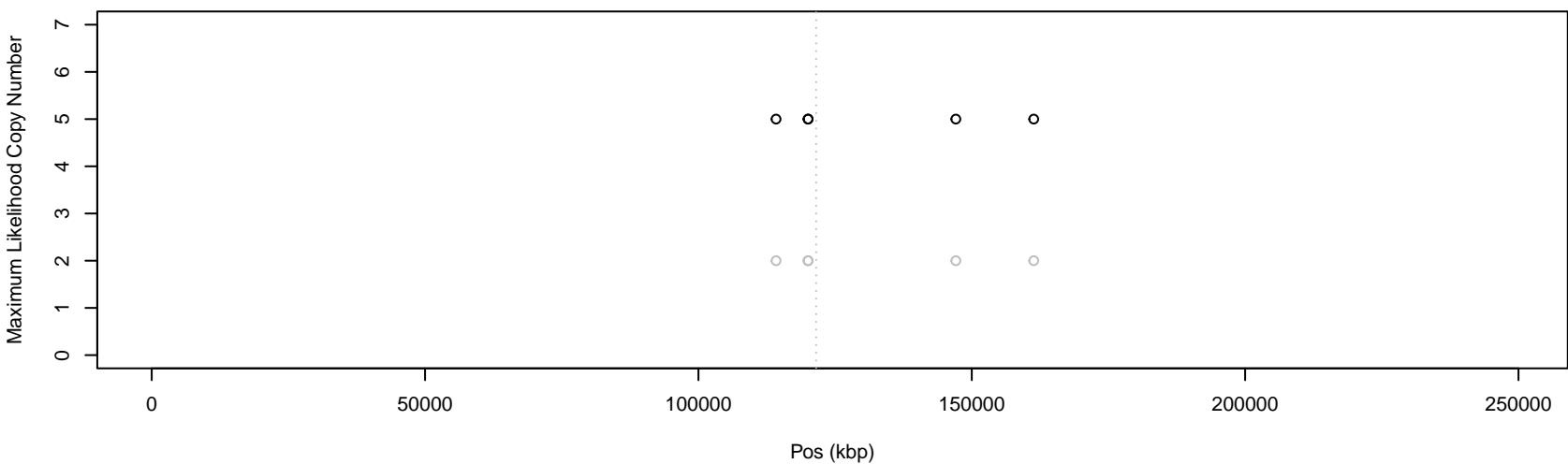
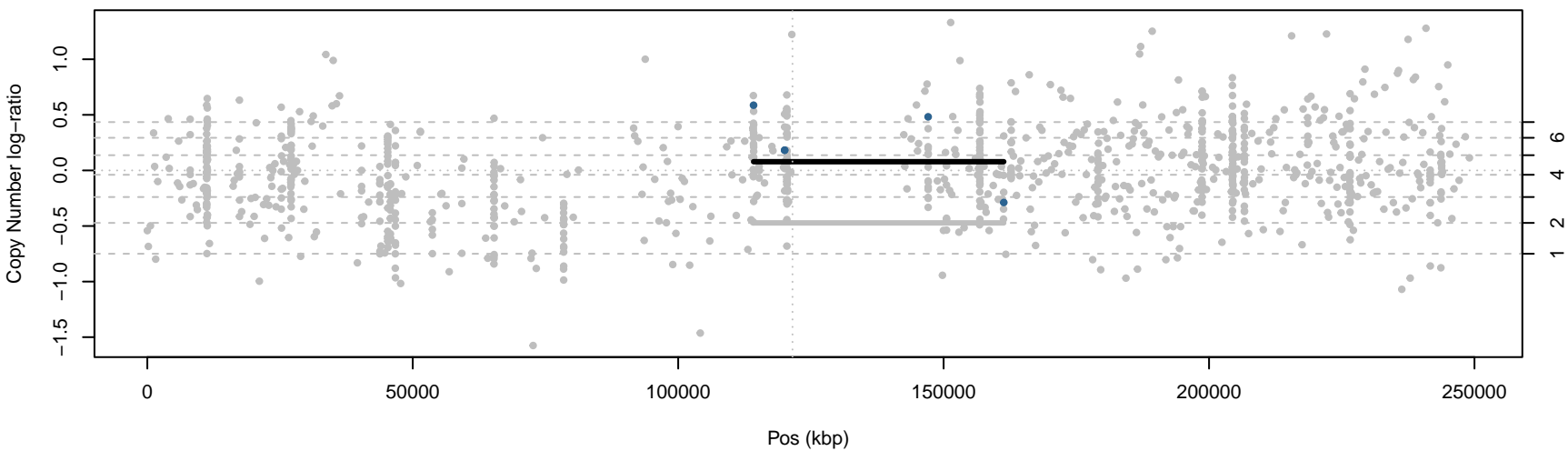
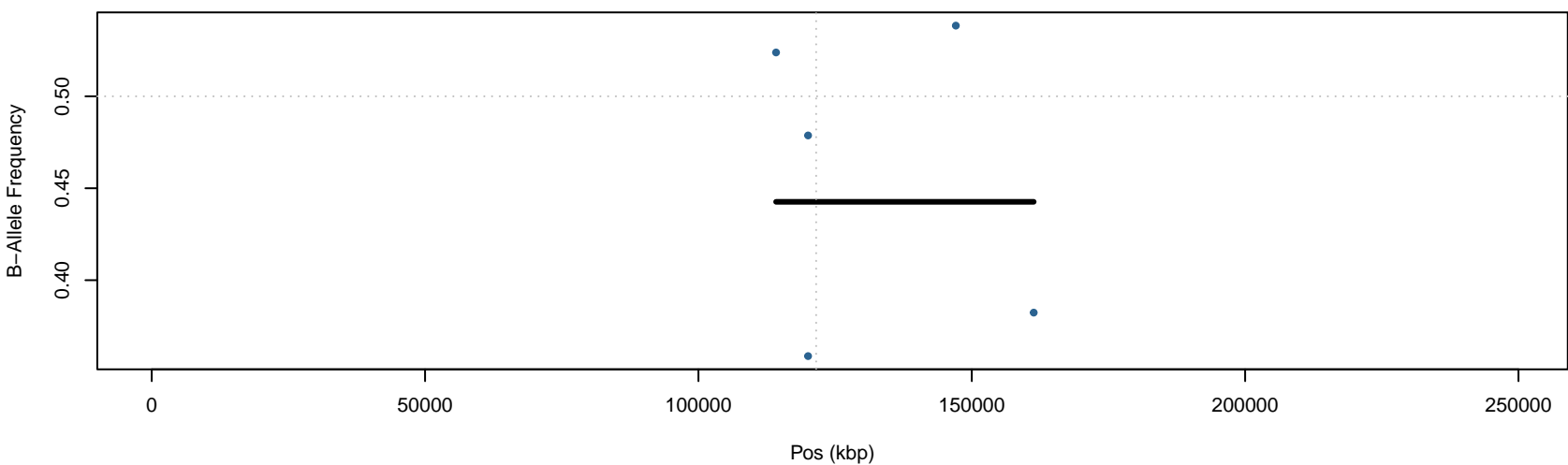
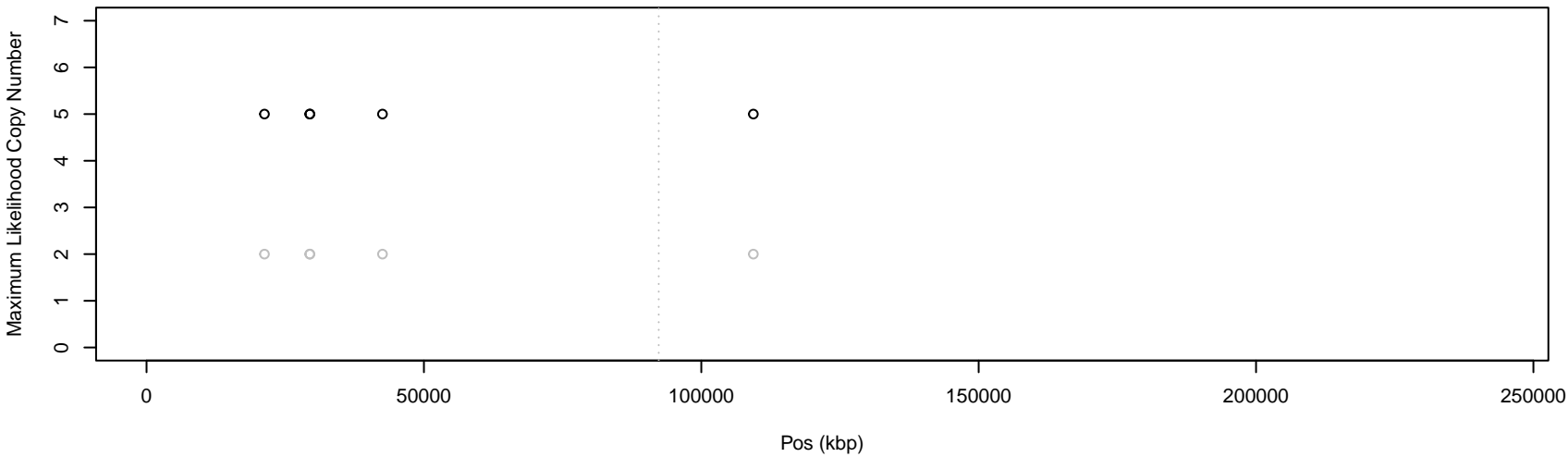
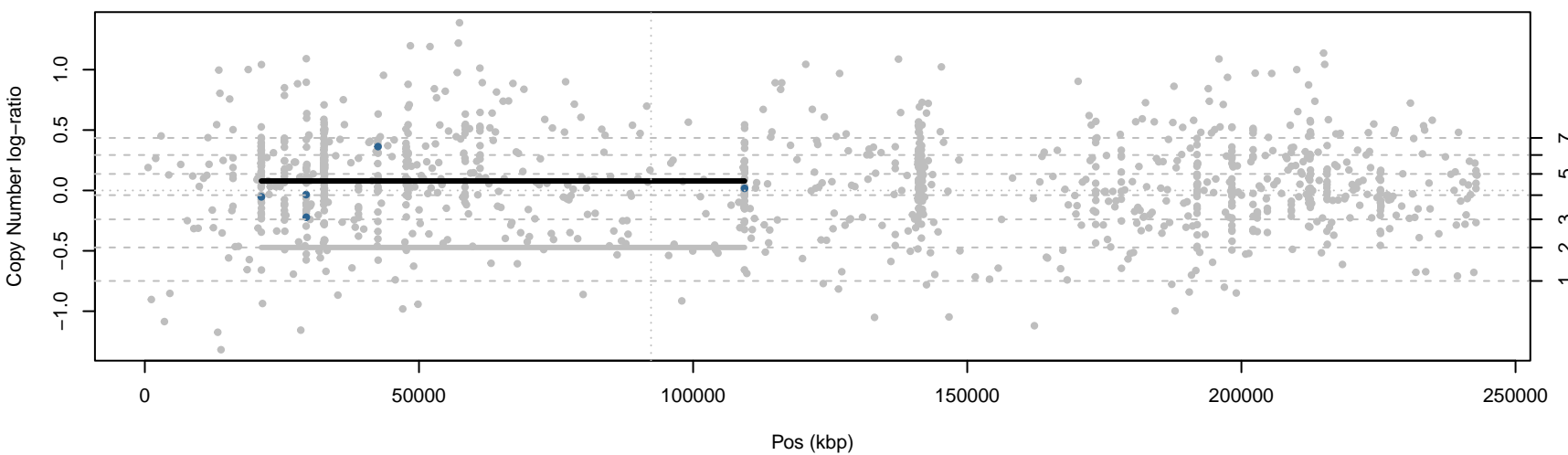
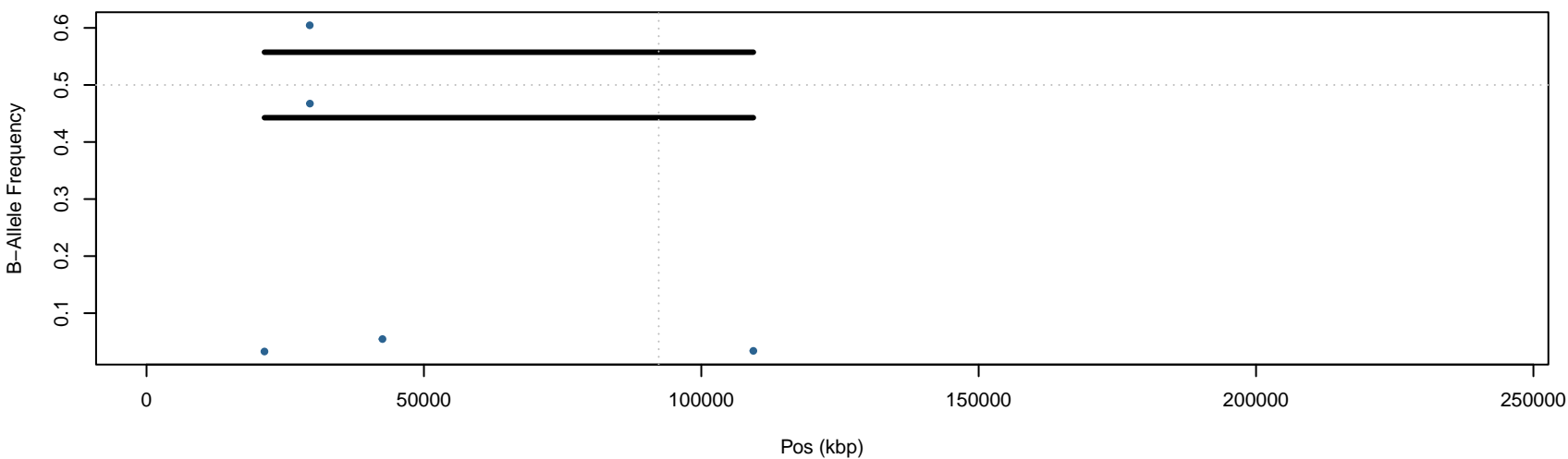


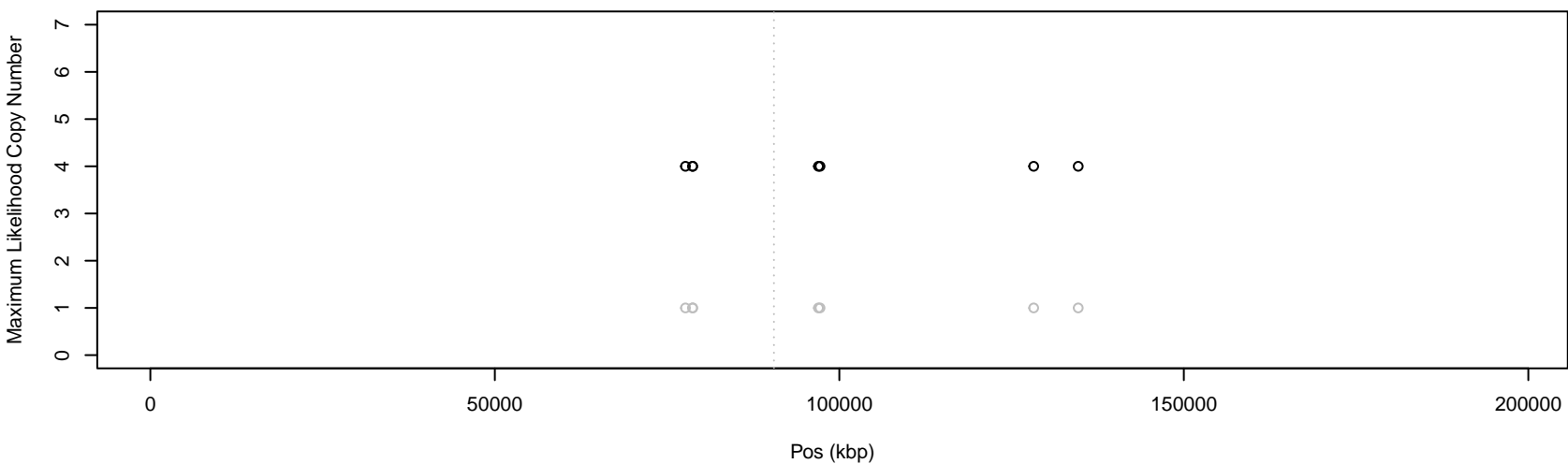
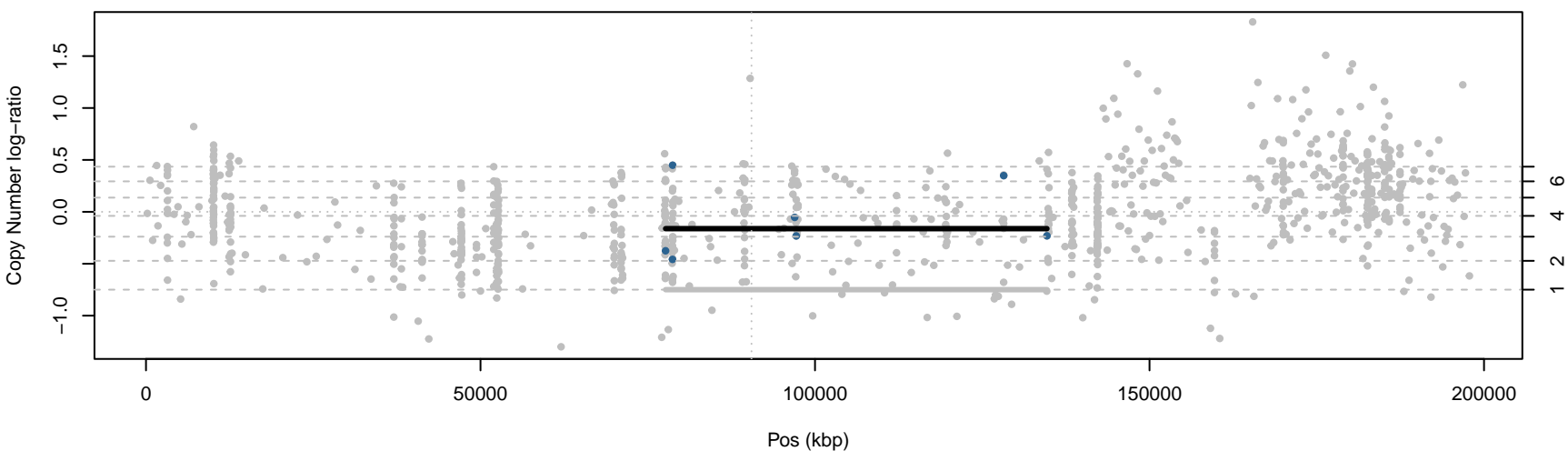
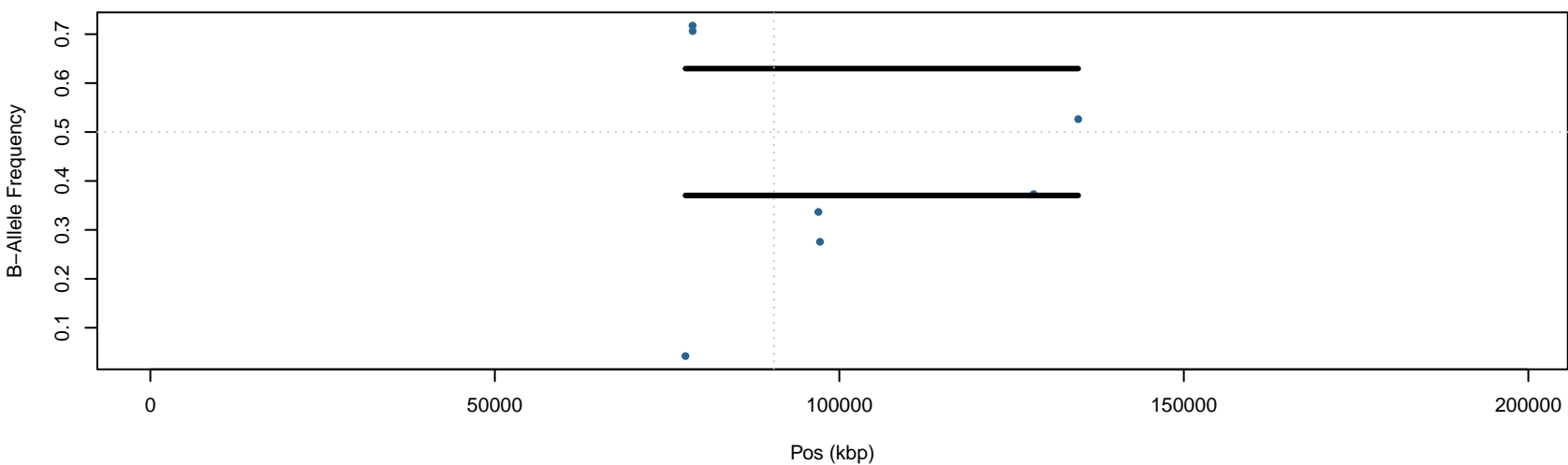
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 1



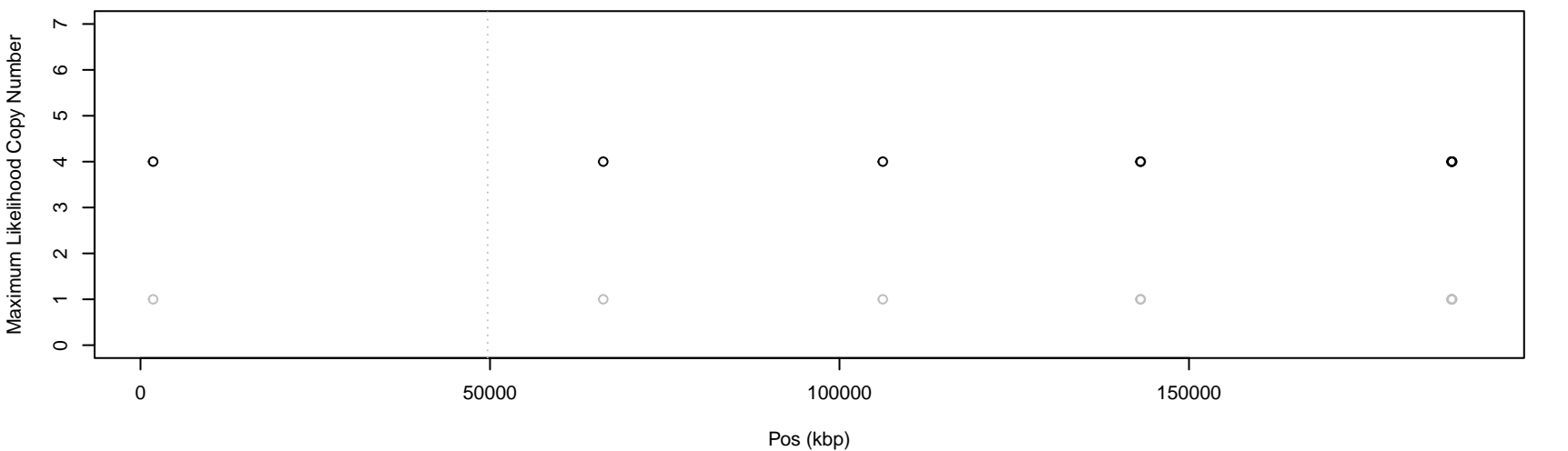
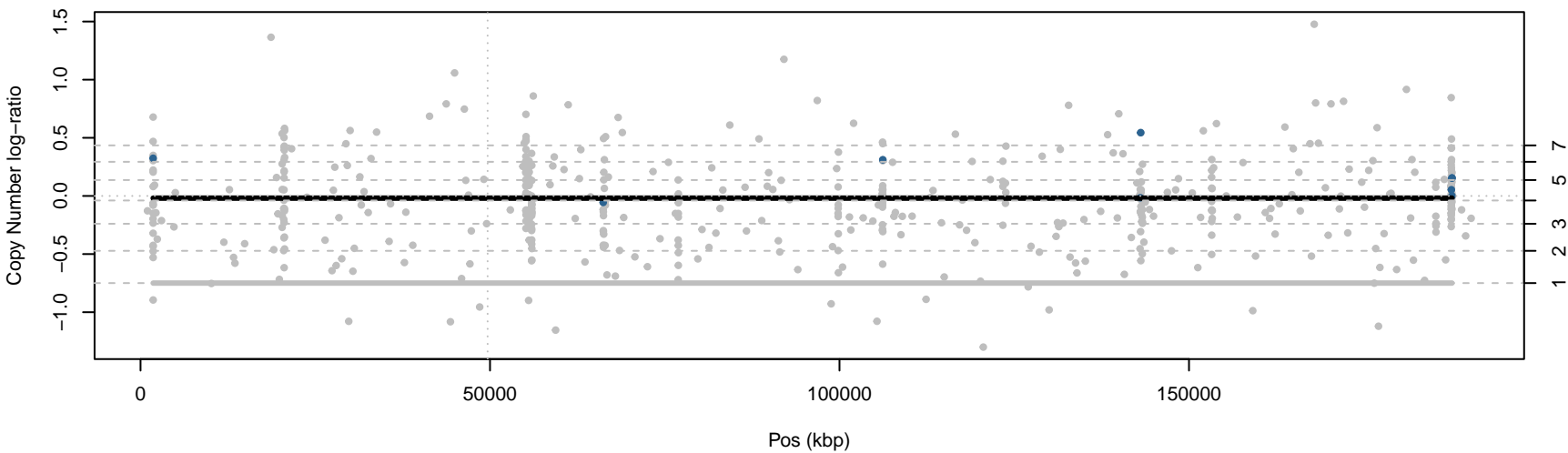
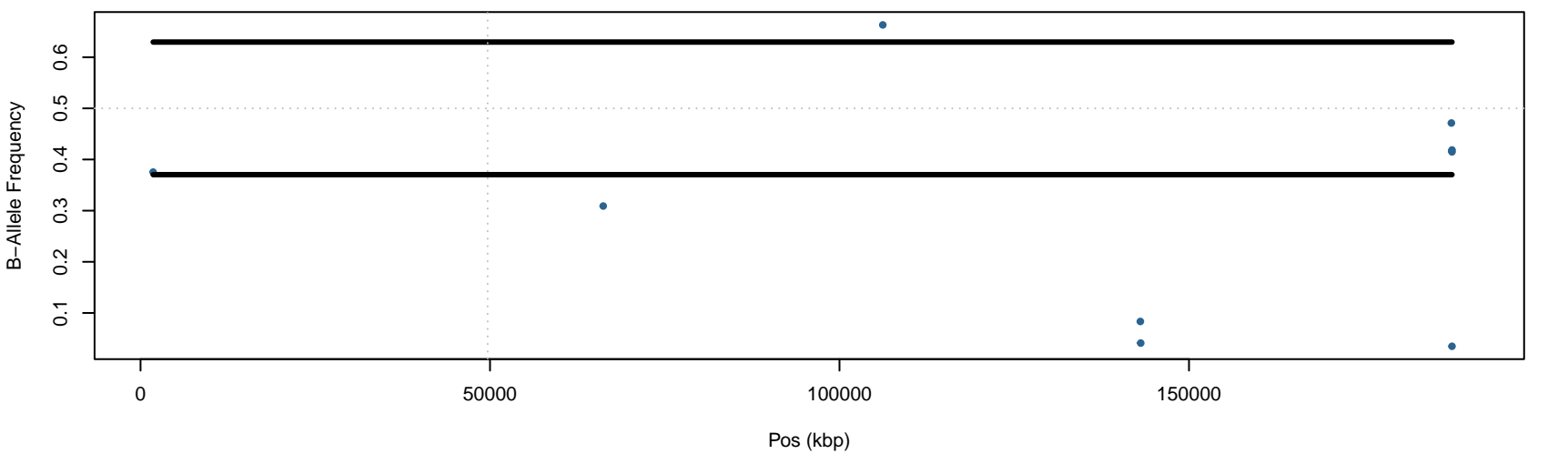
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 2



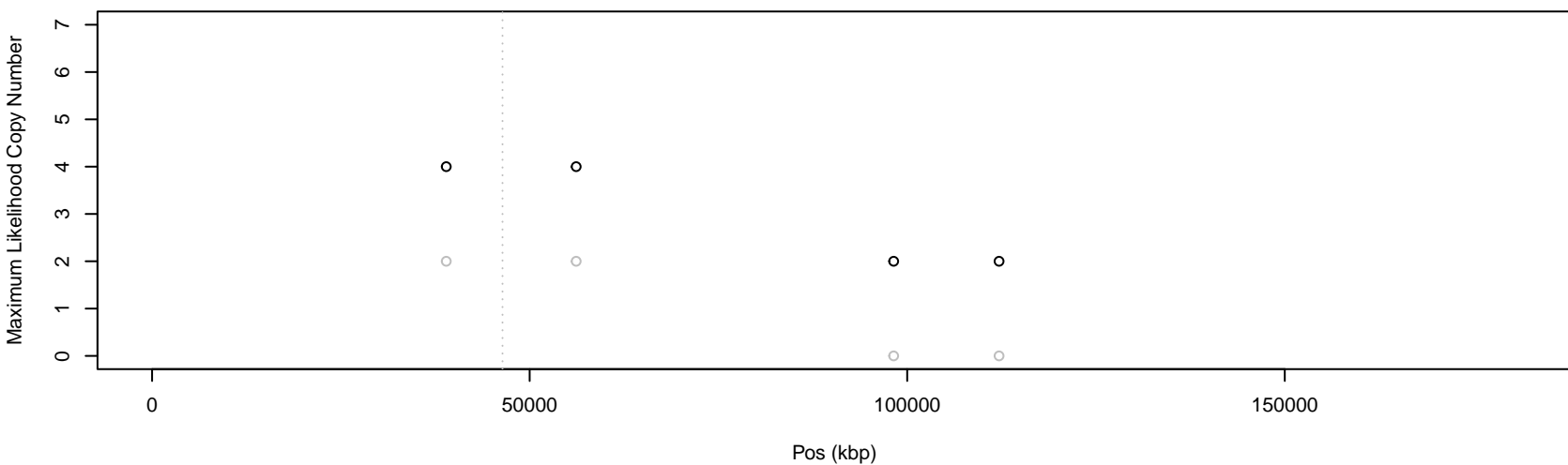
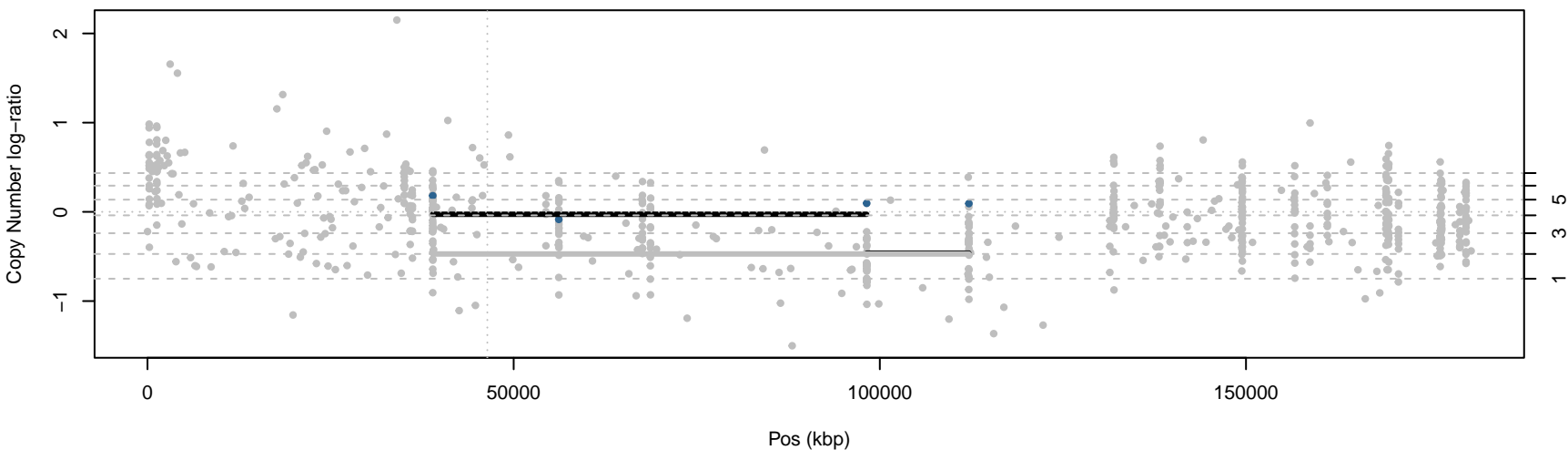
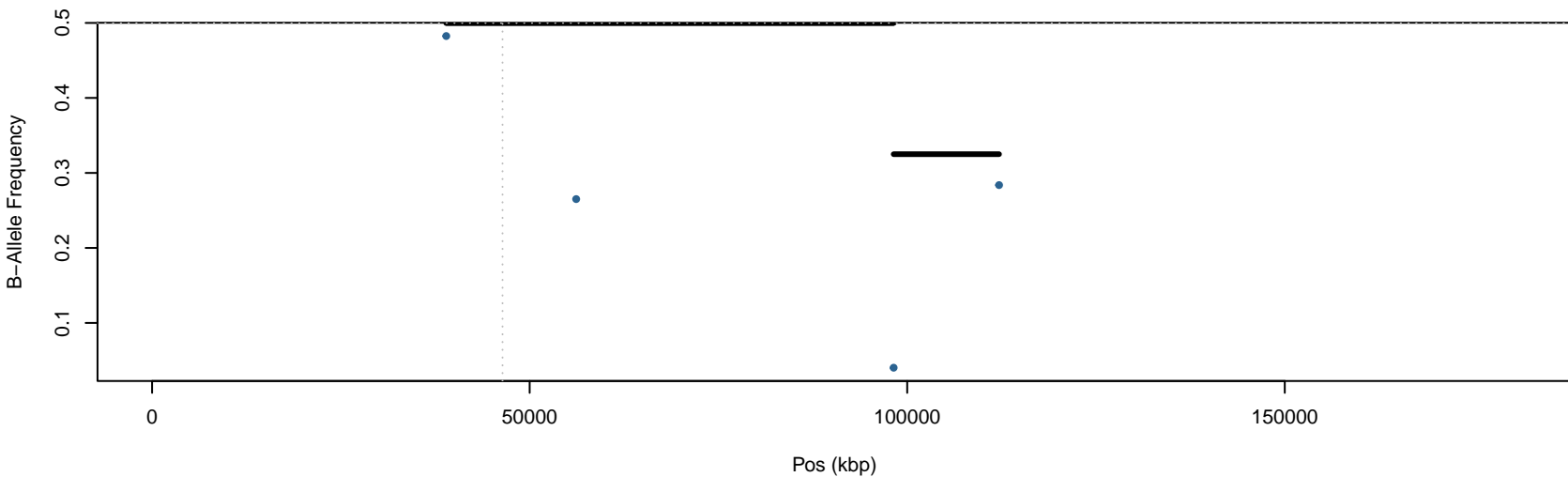
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 3



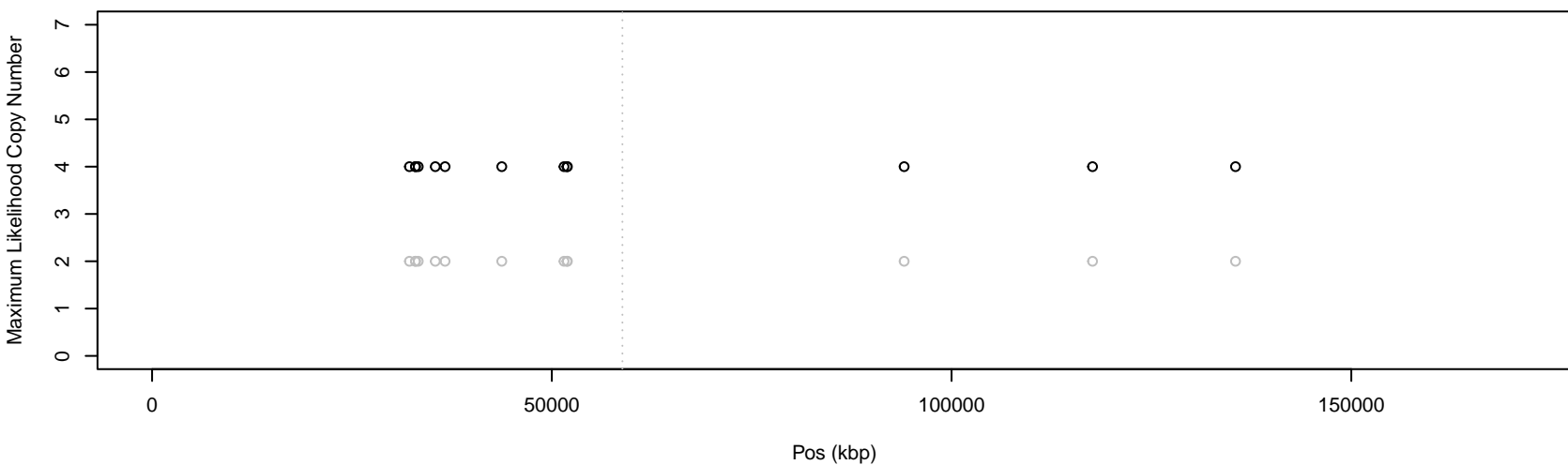
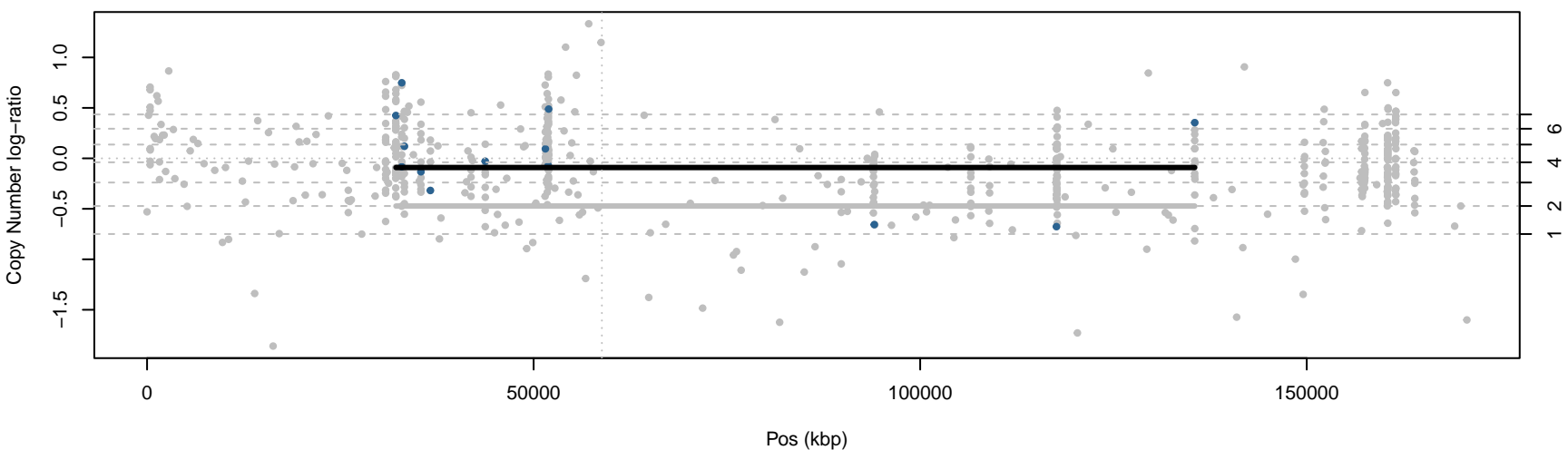
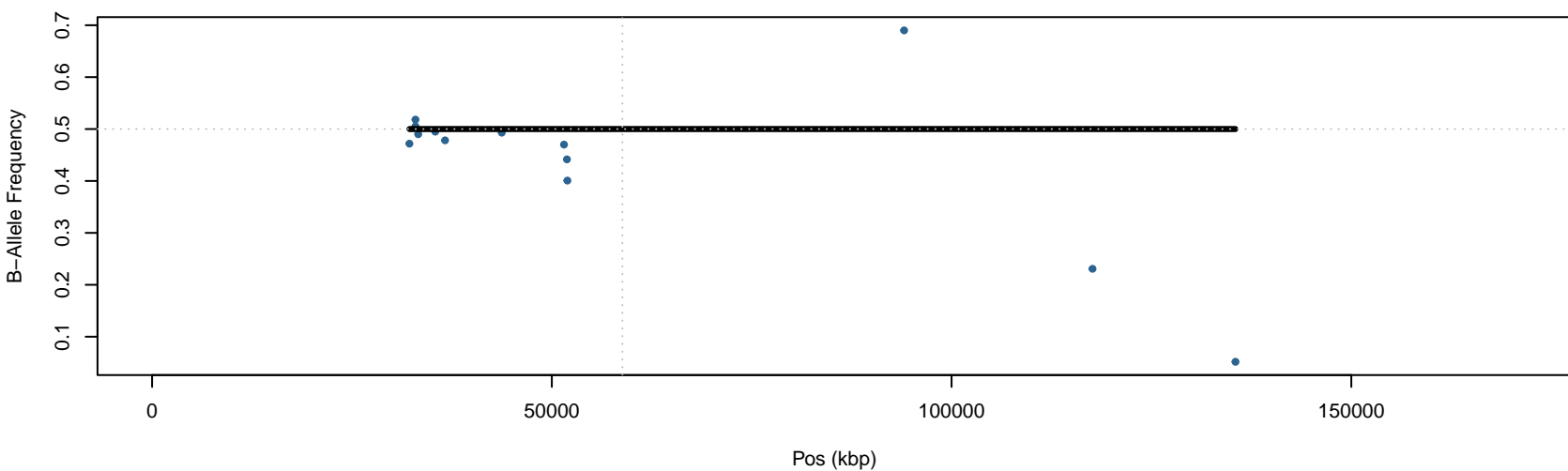
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 4



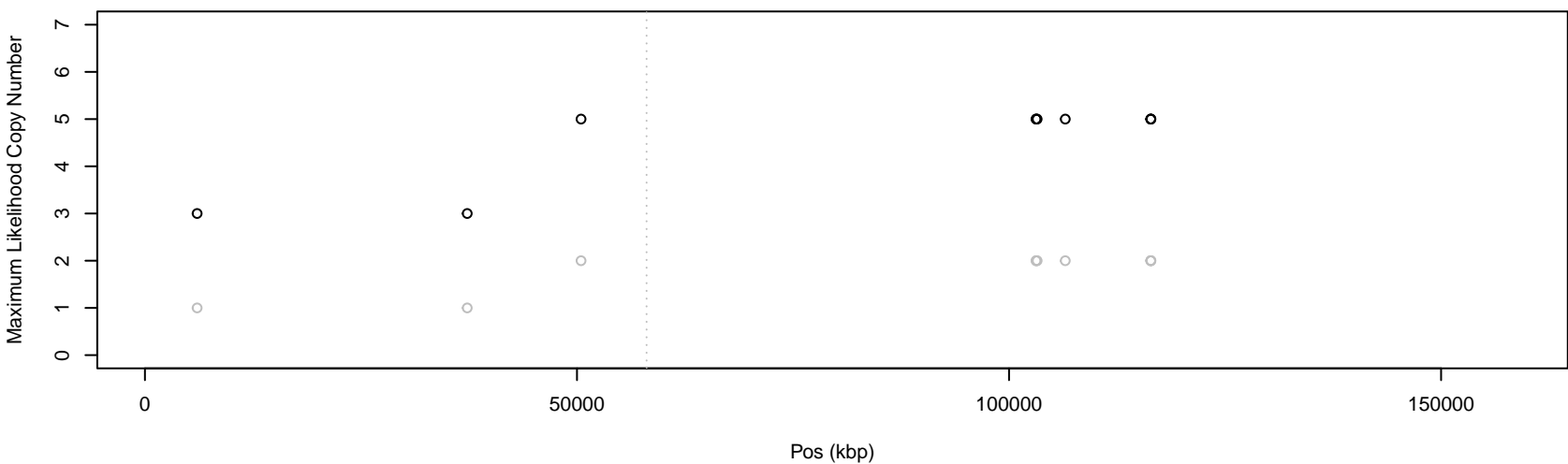
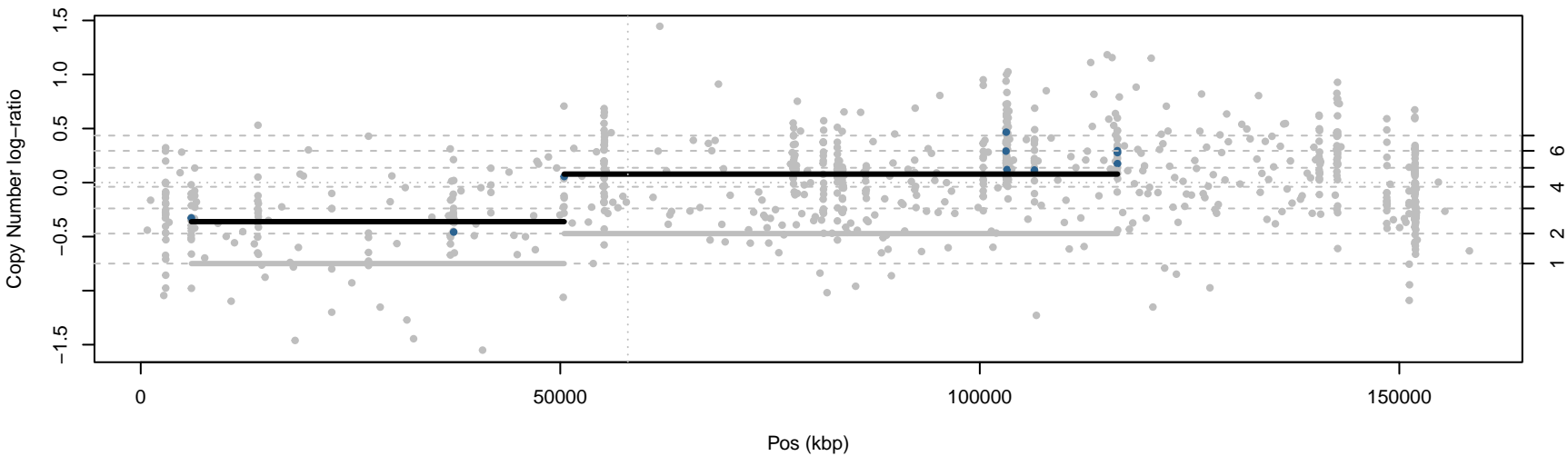
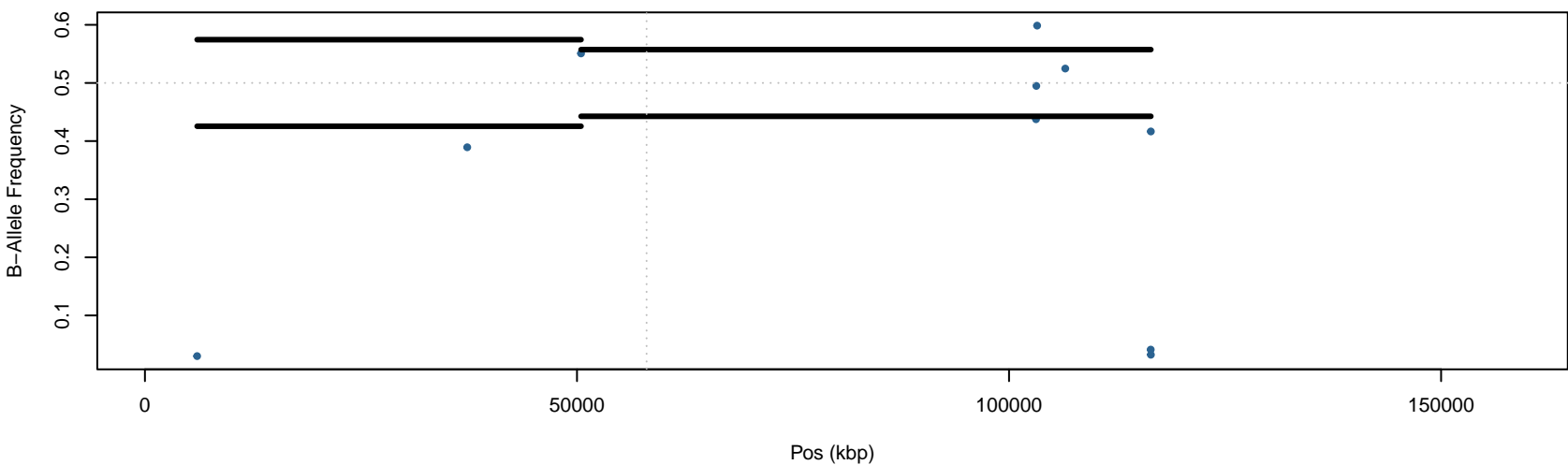
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 5



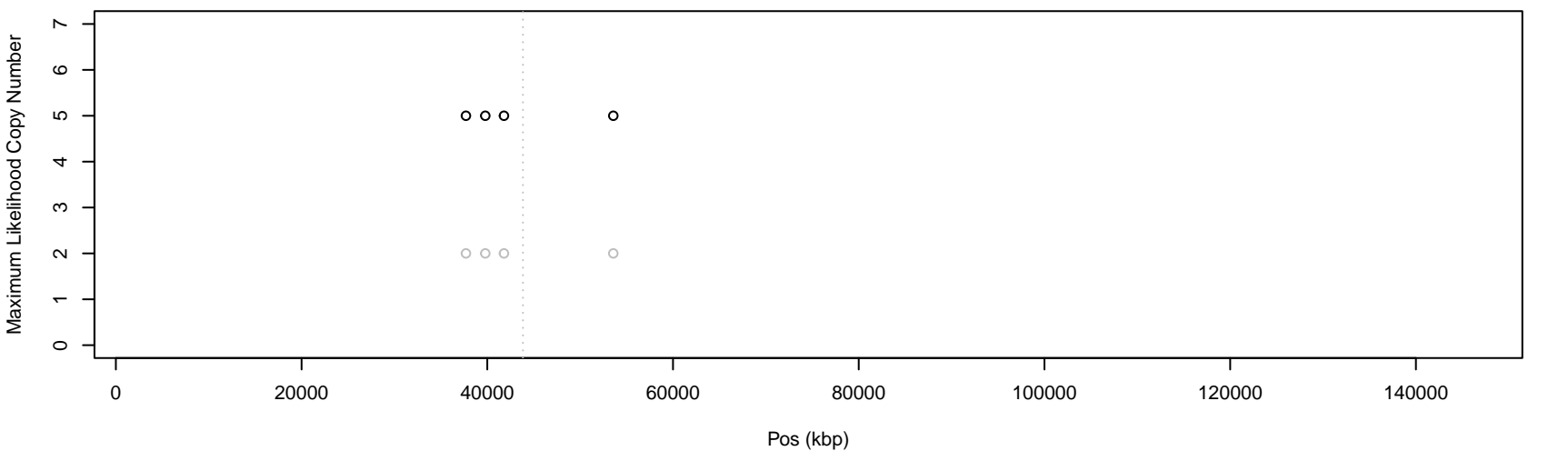
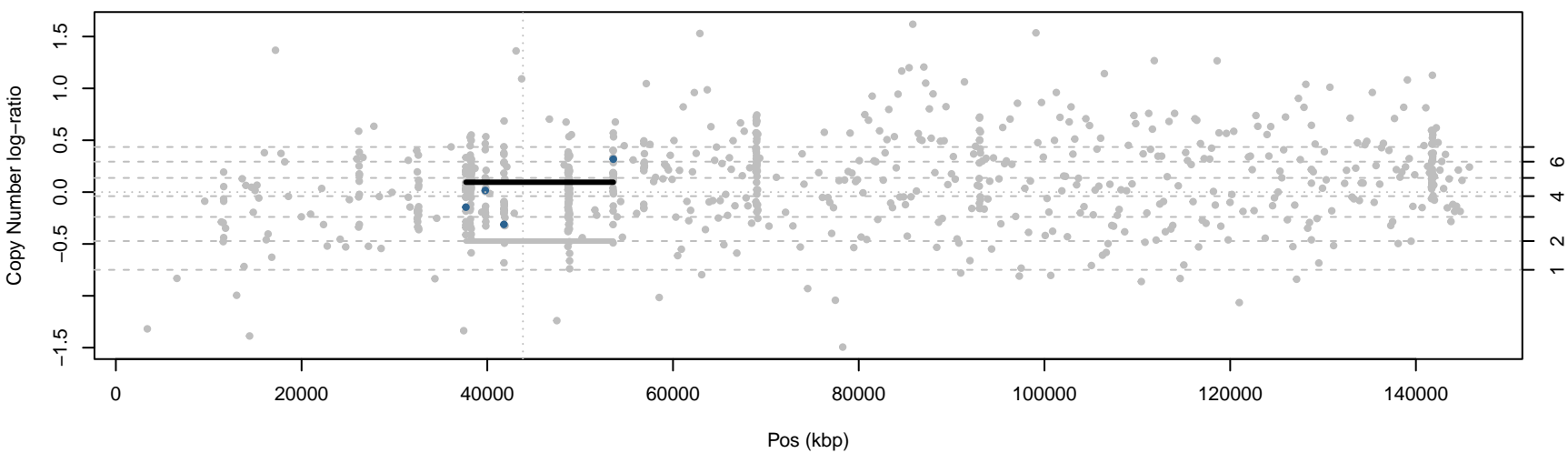
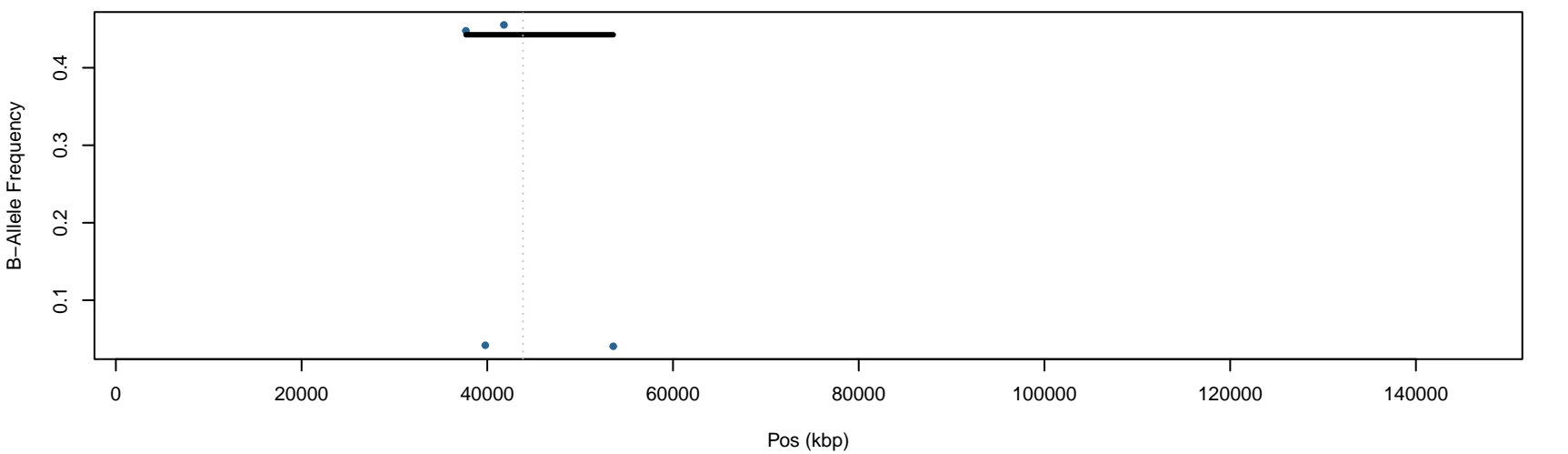
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 6



Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 7

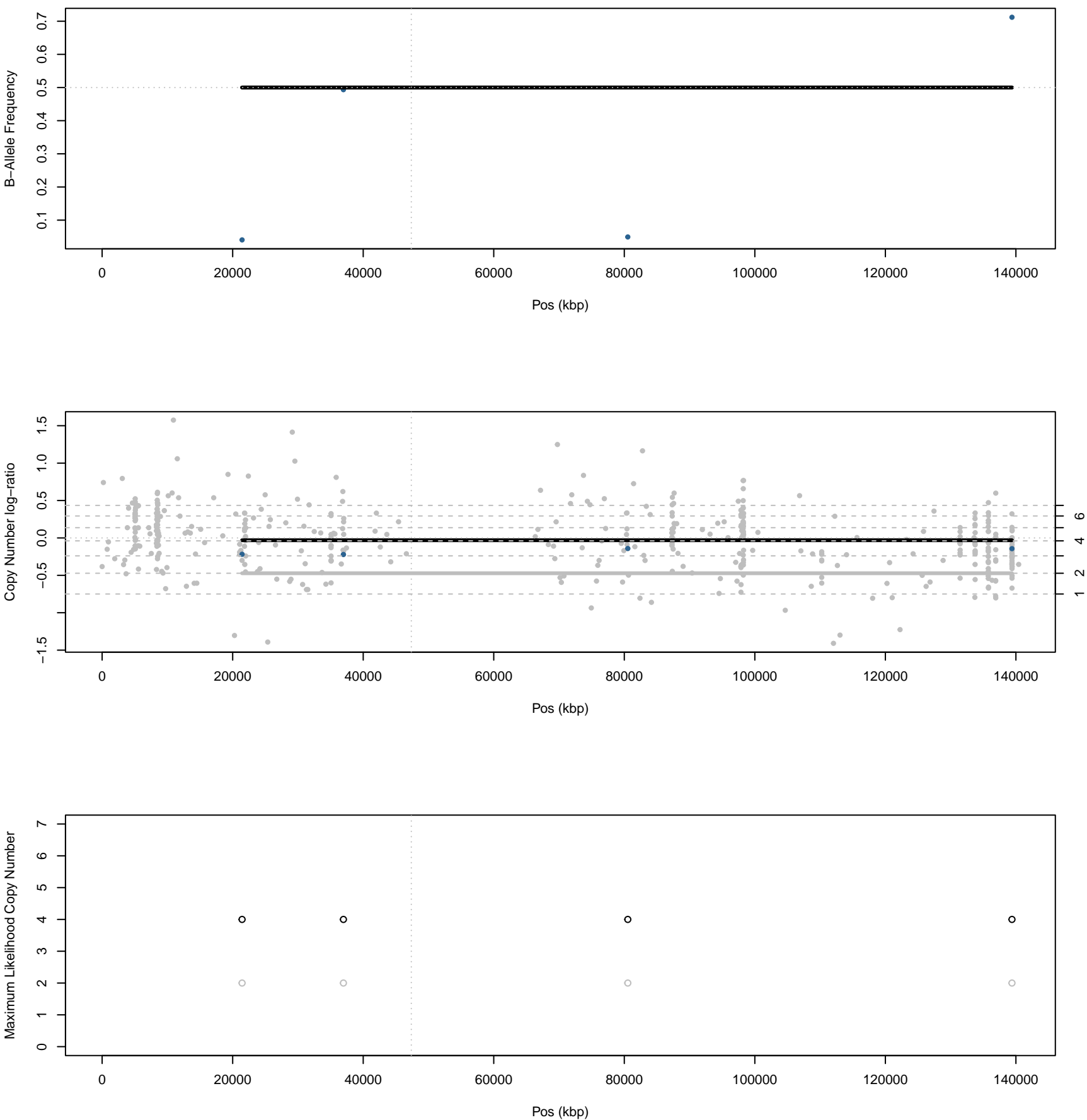


Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 8

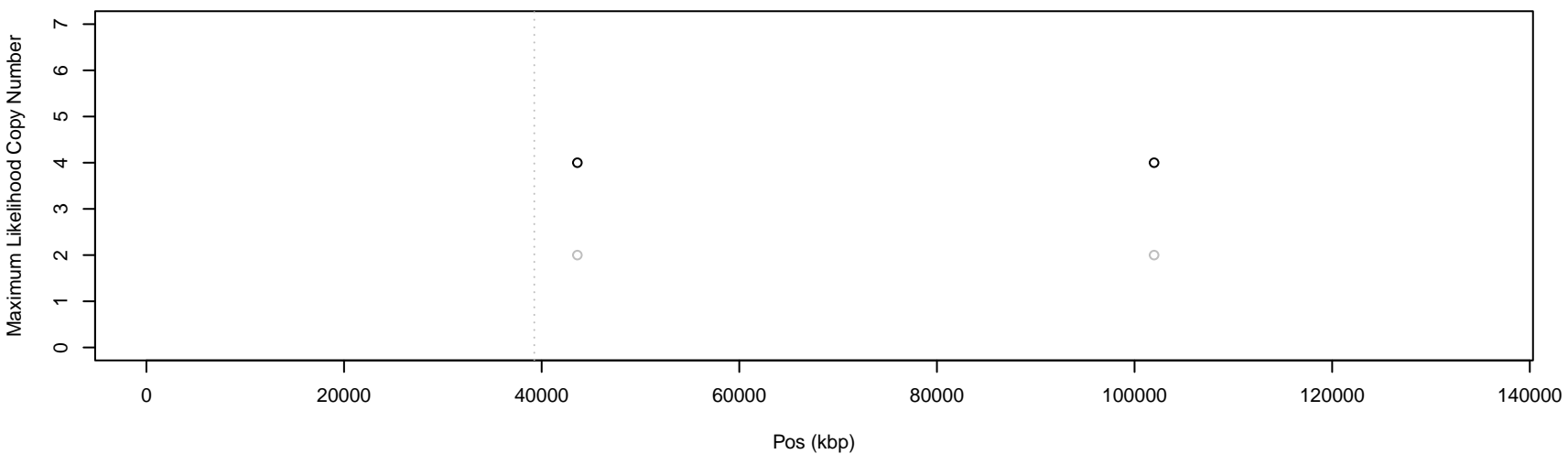
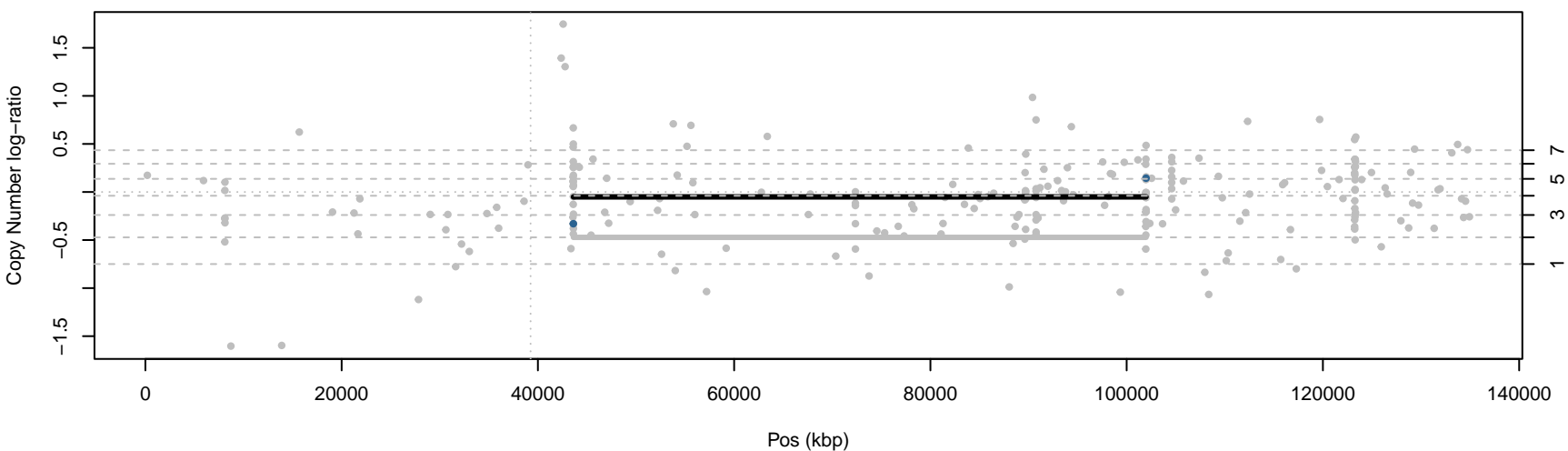
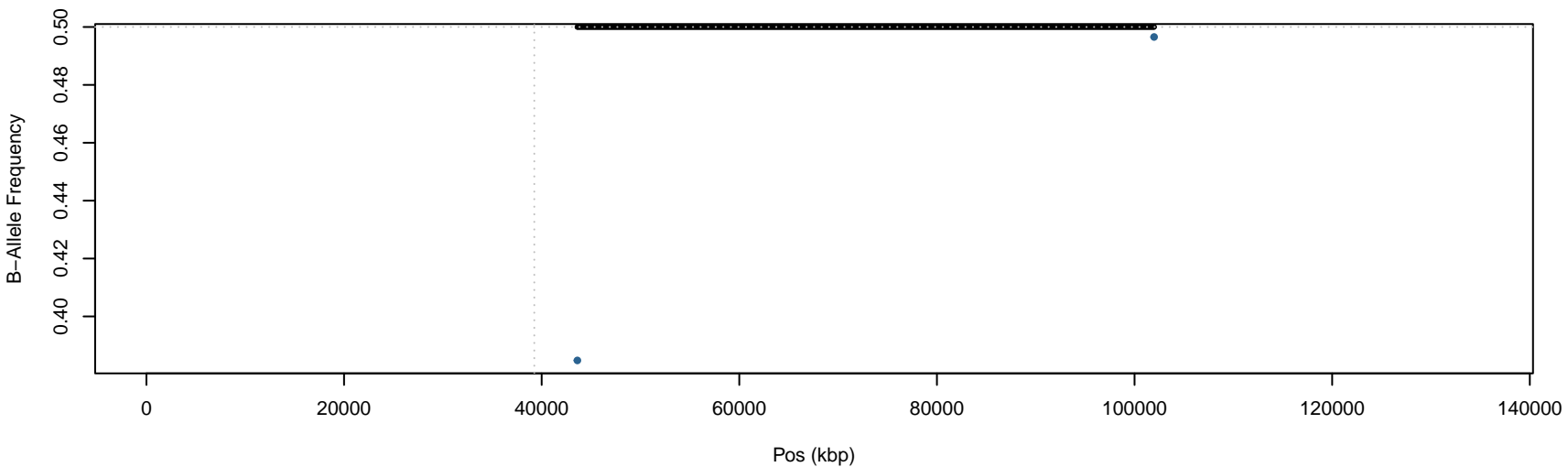




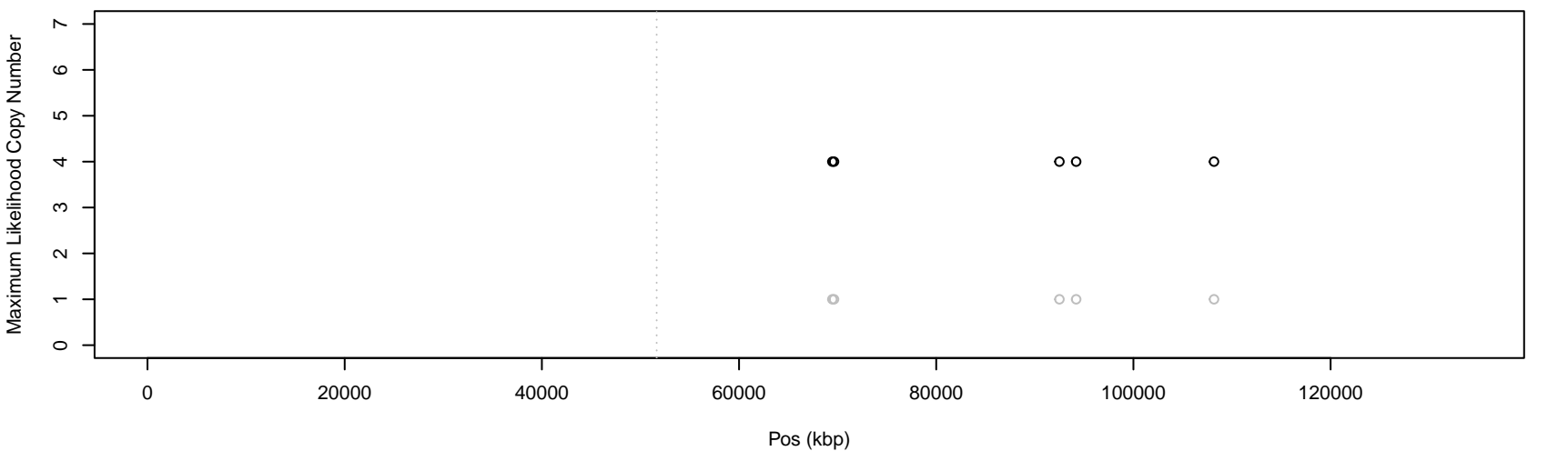
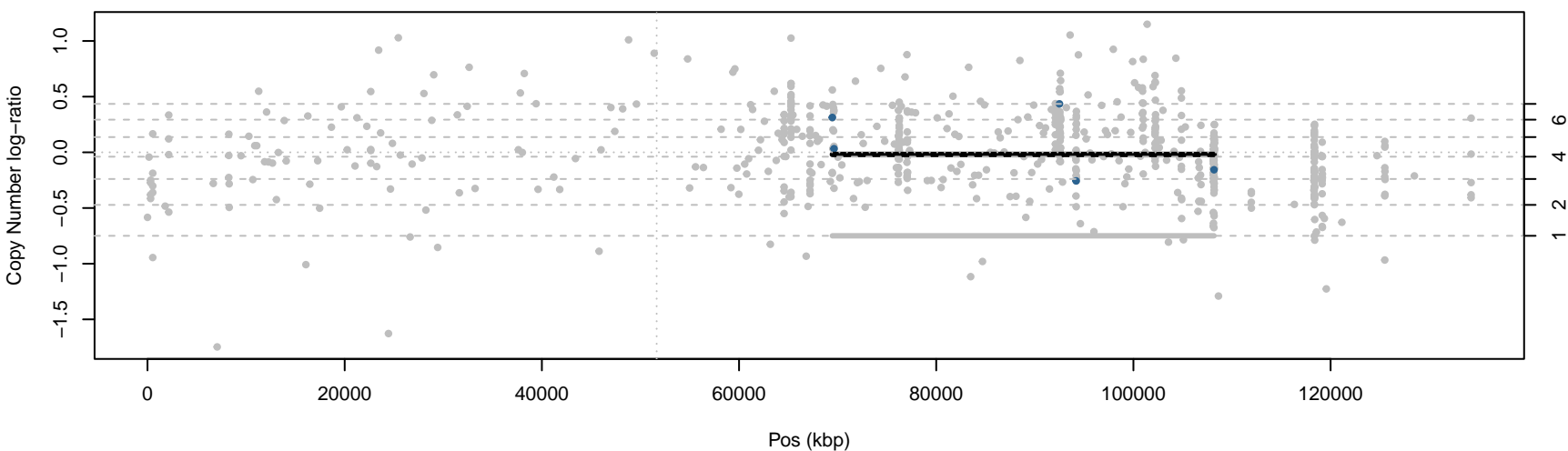
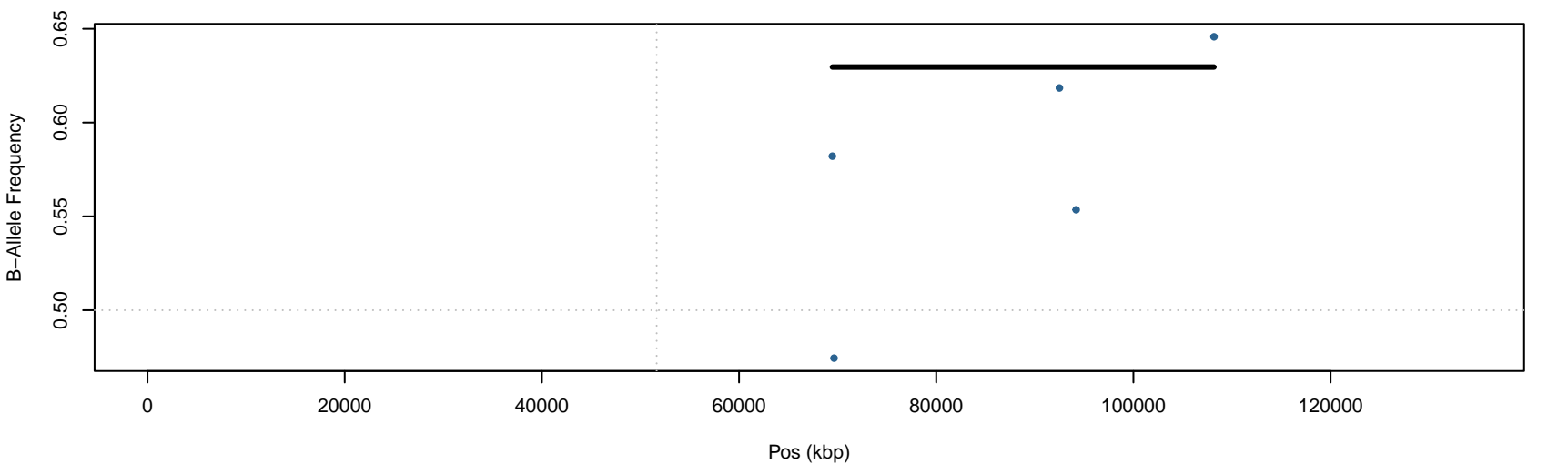
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 9



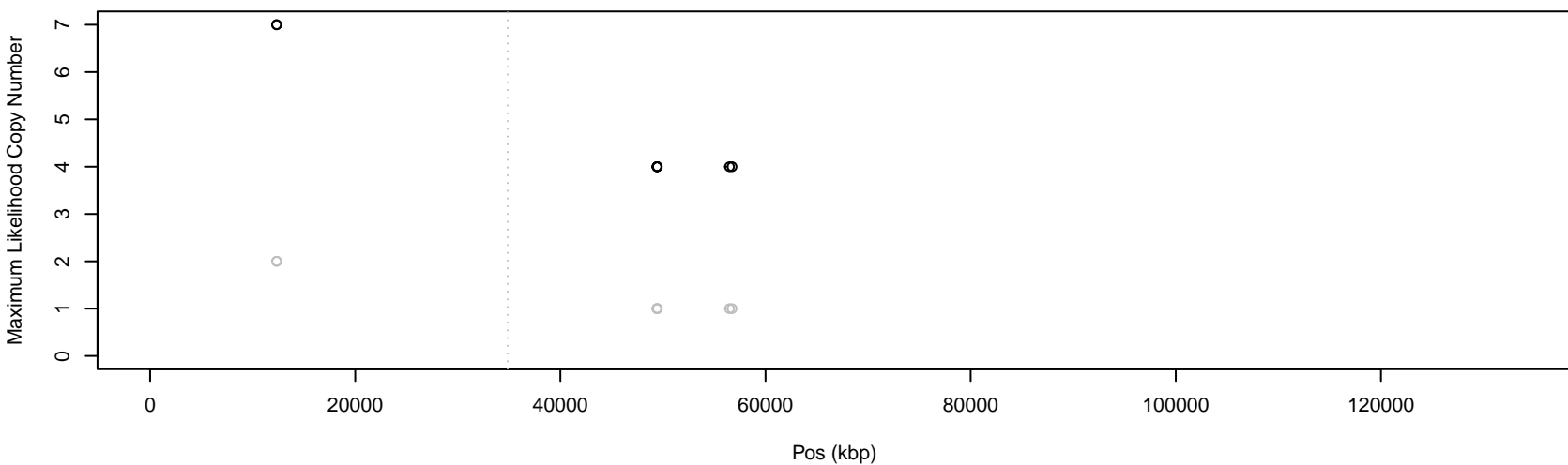
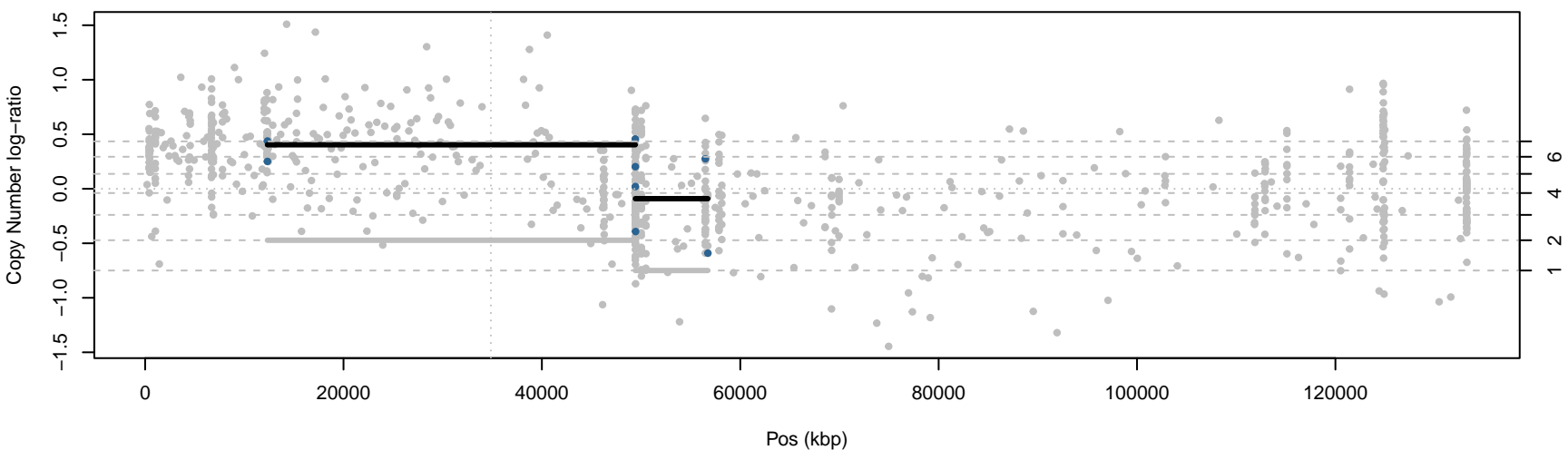
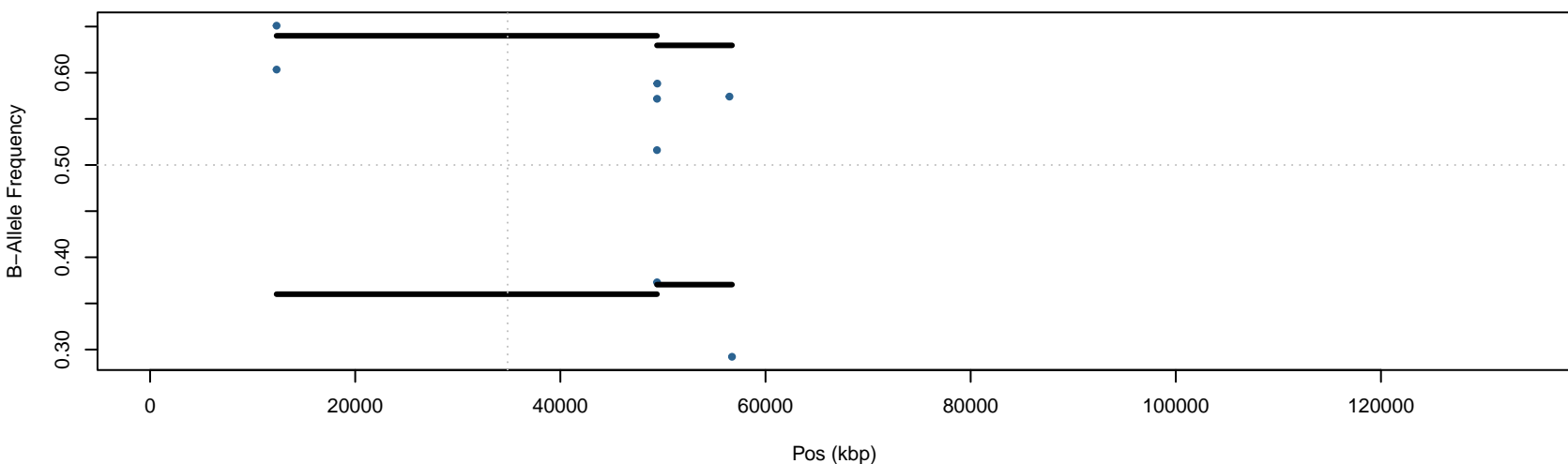
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 10



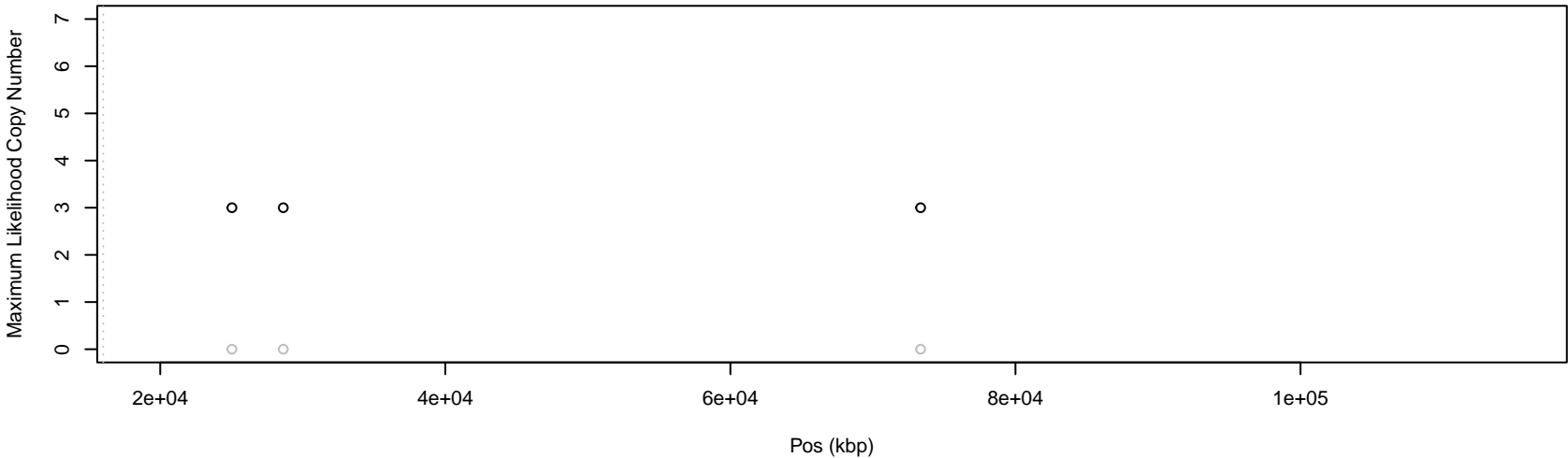
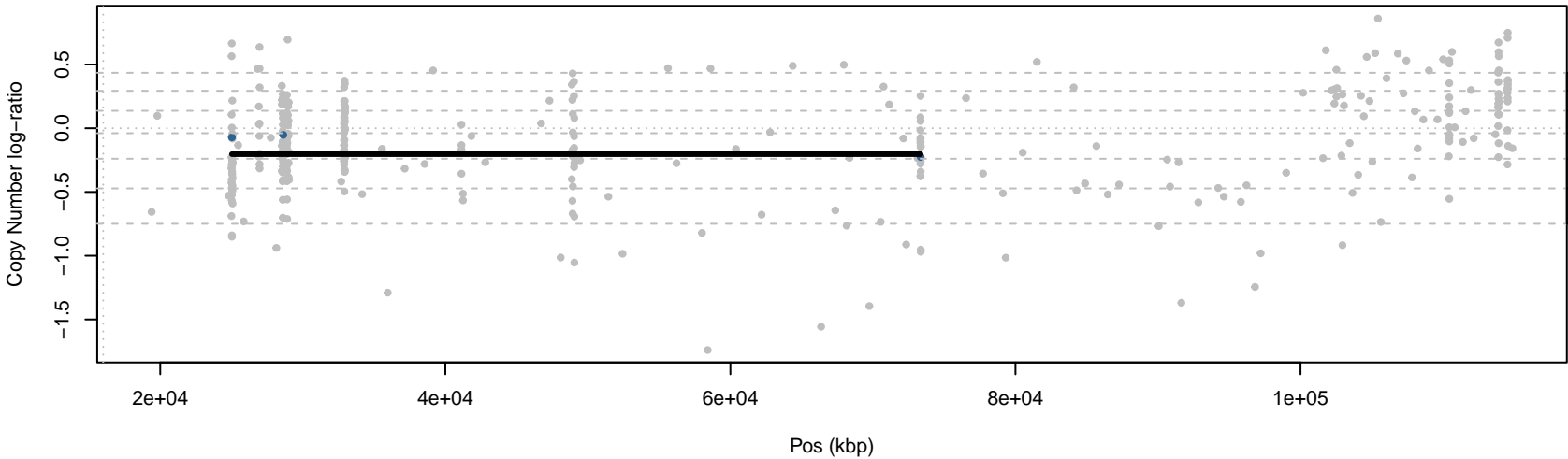
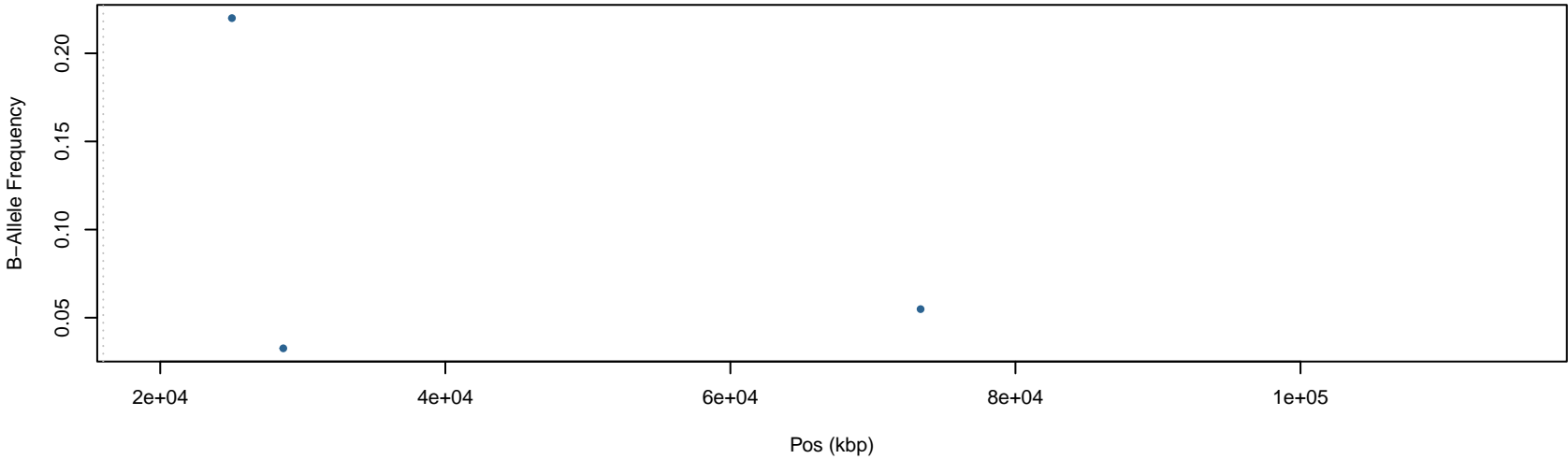
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 11



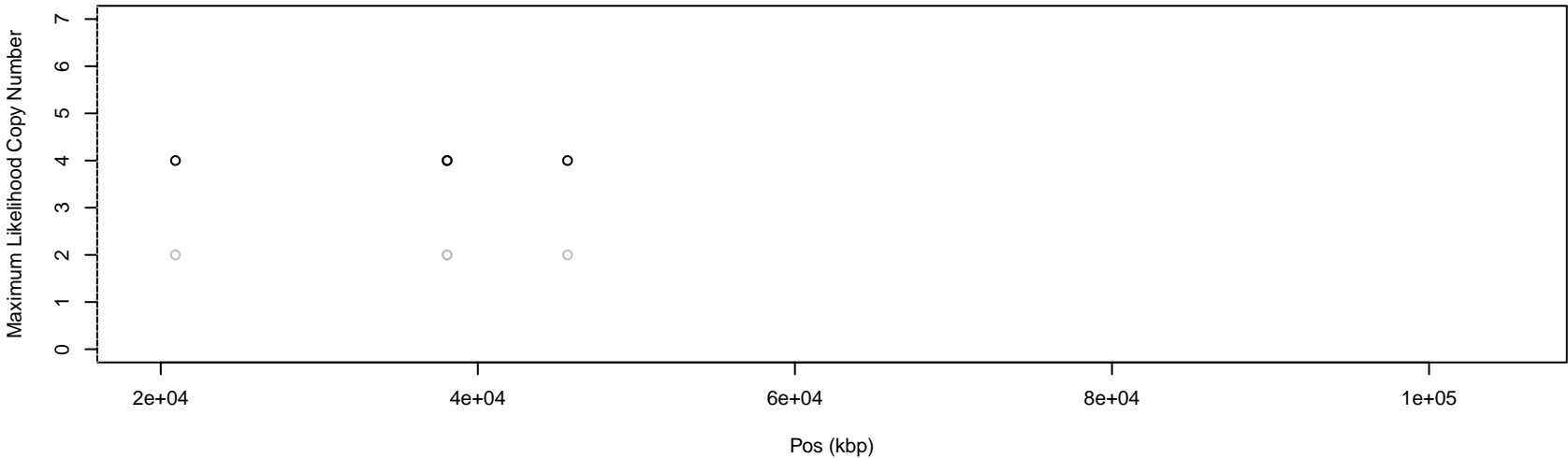
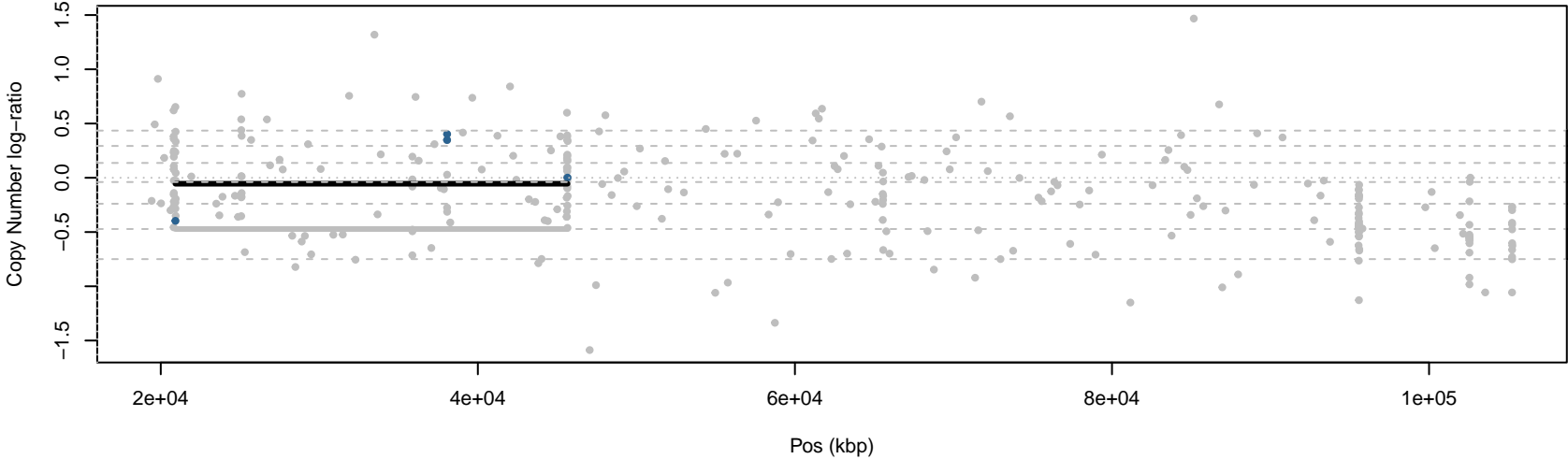
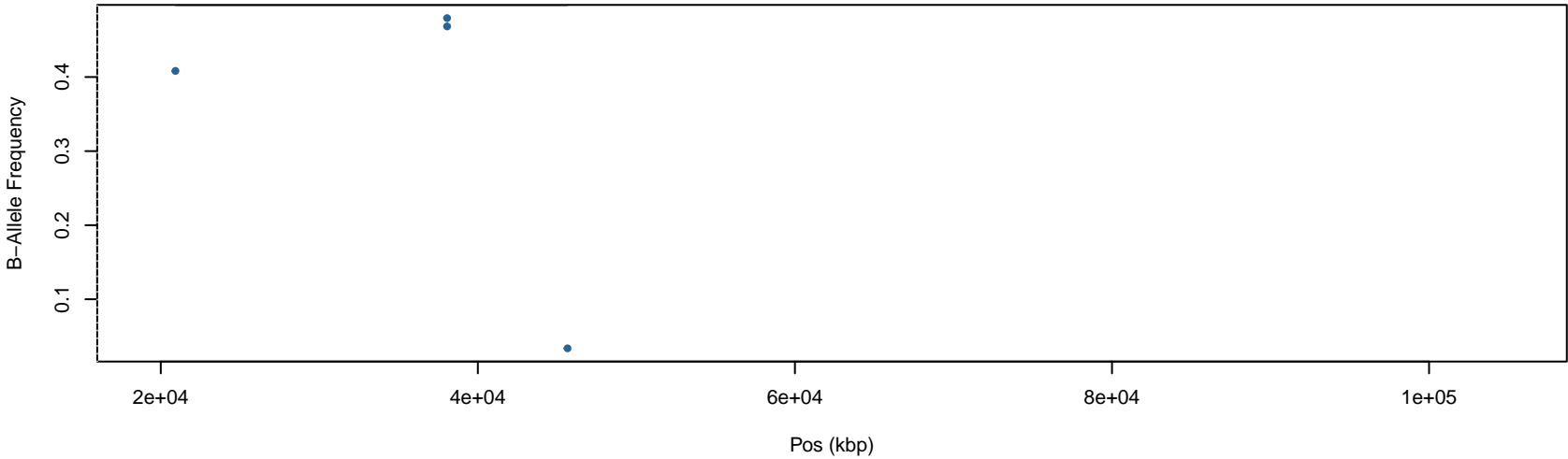
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 12



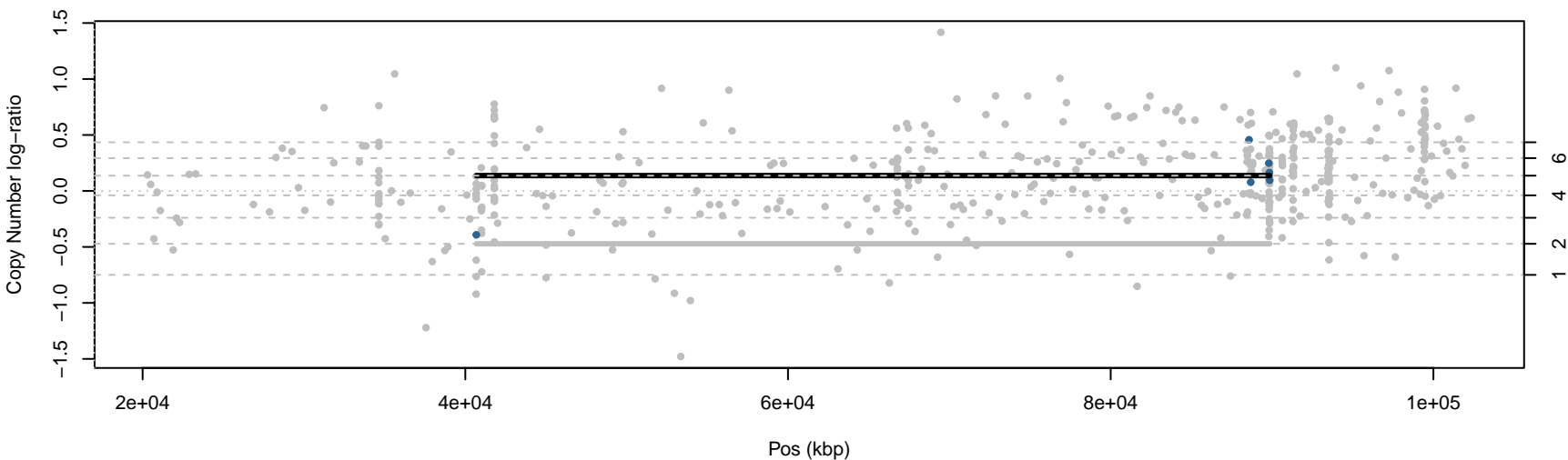
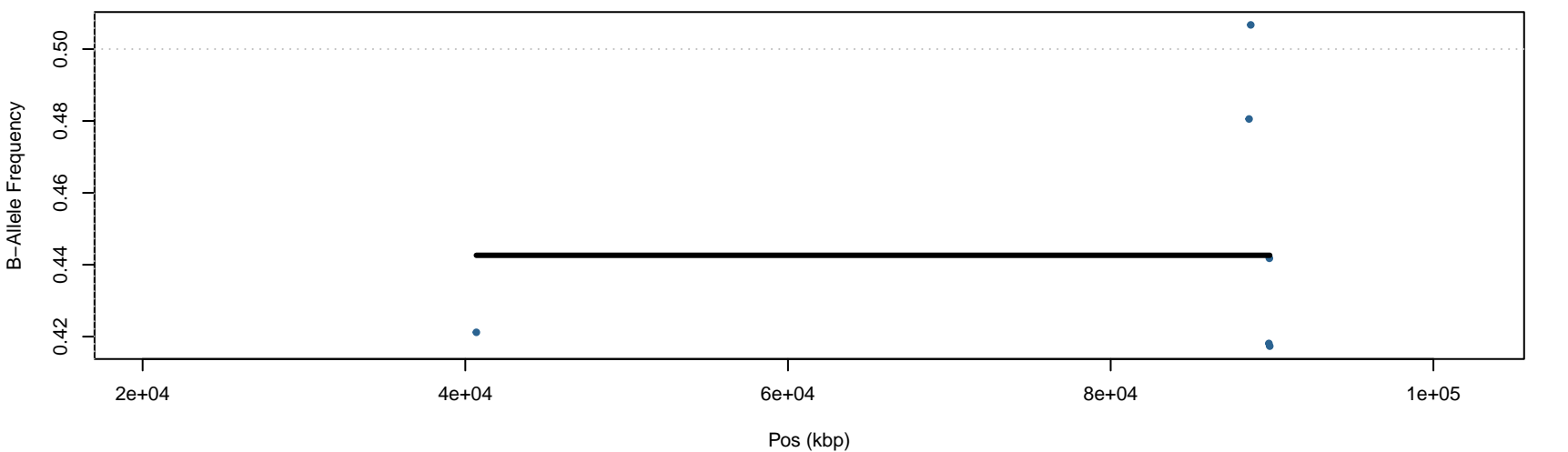
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 13



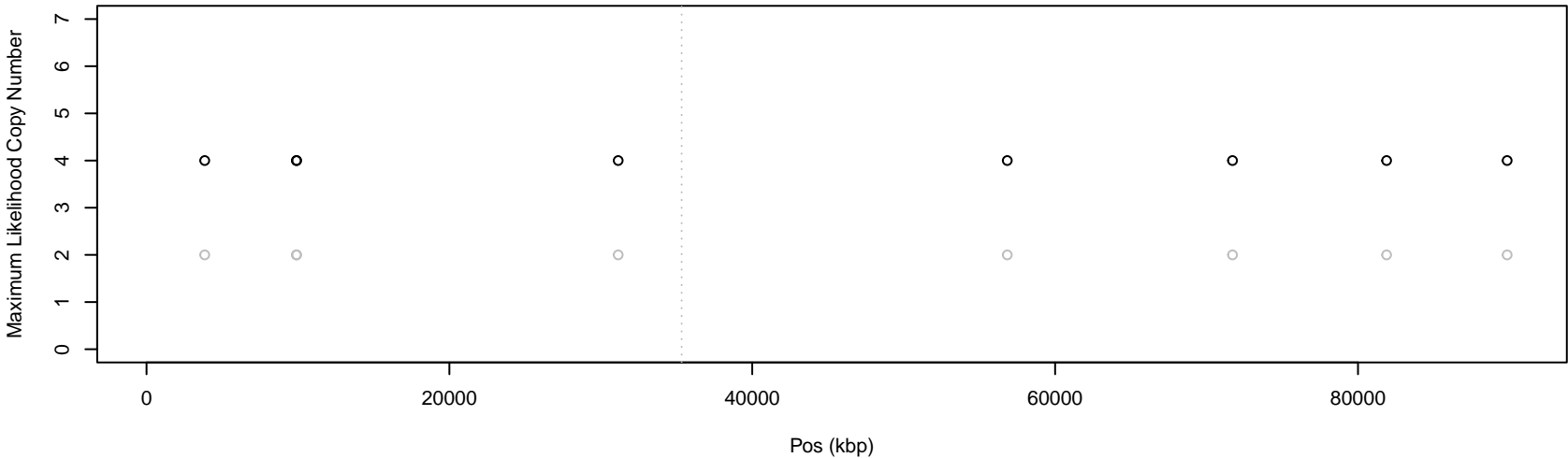
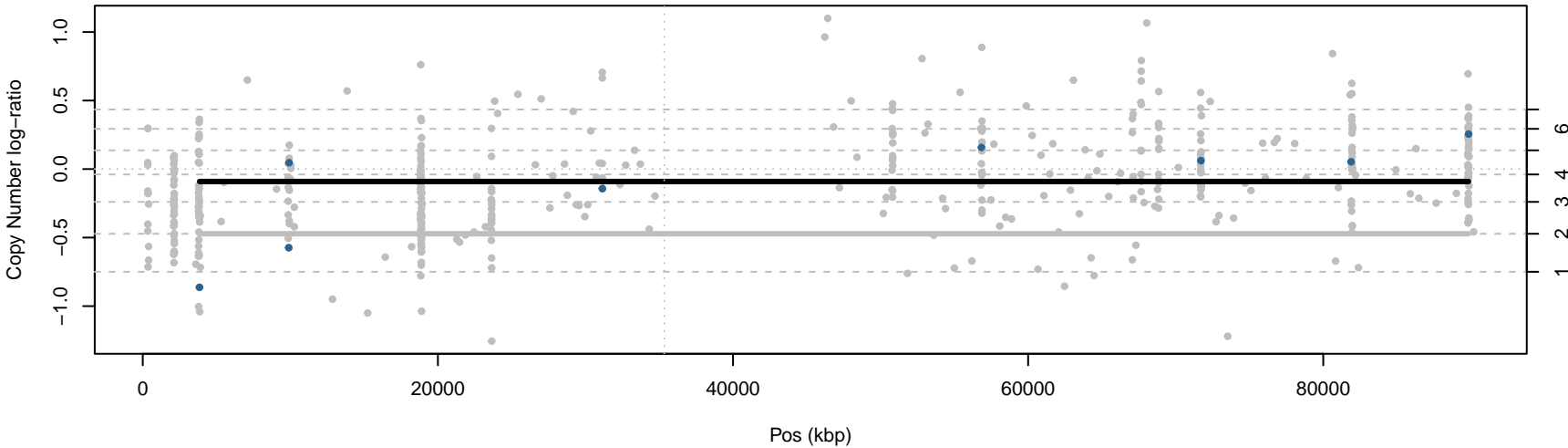
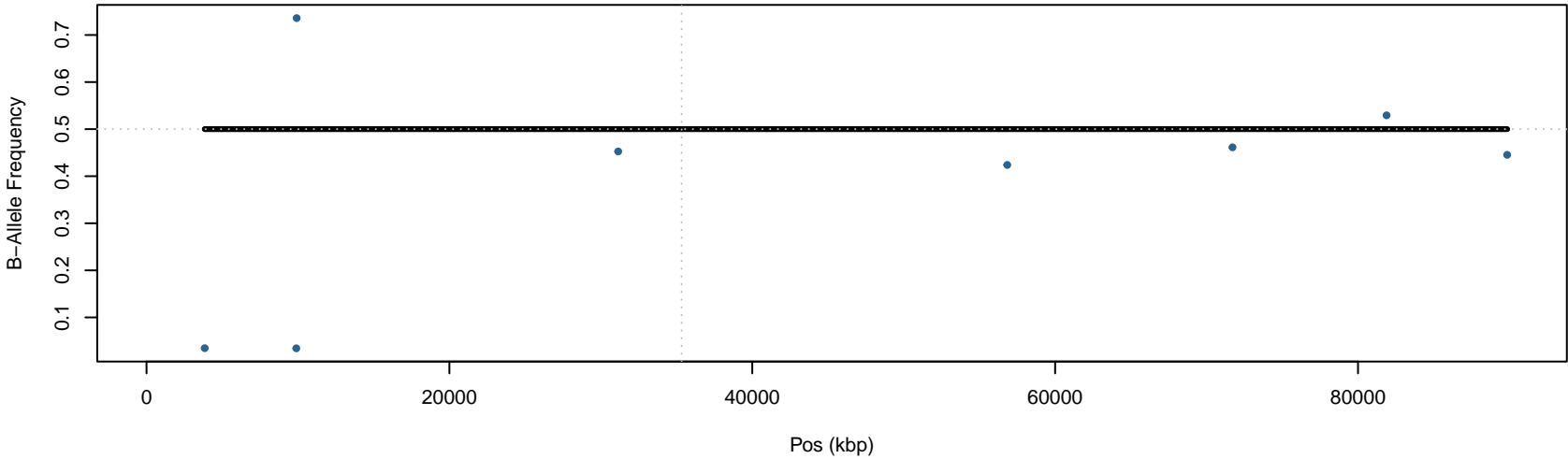
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 14



Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 15

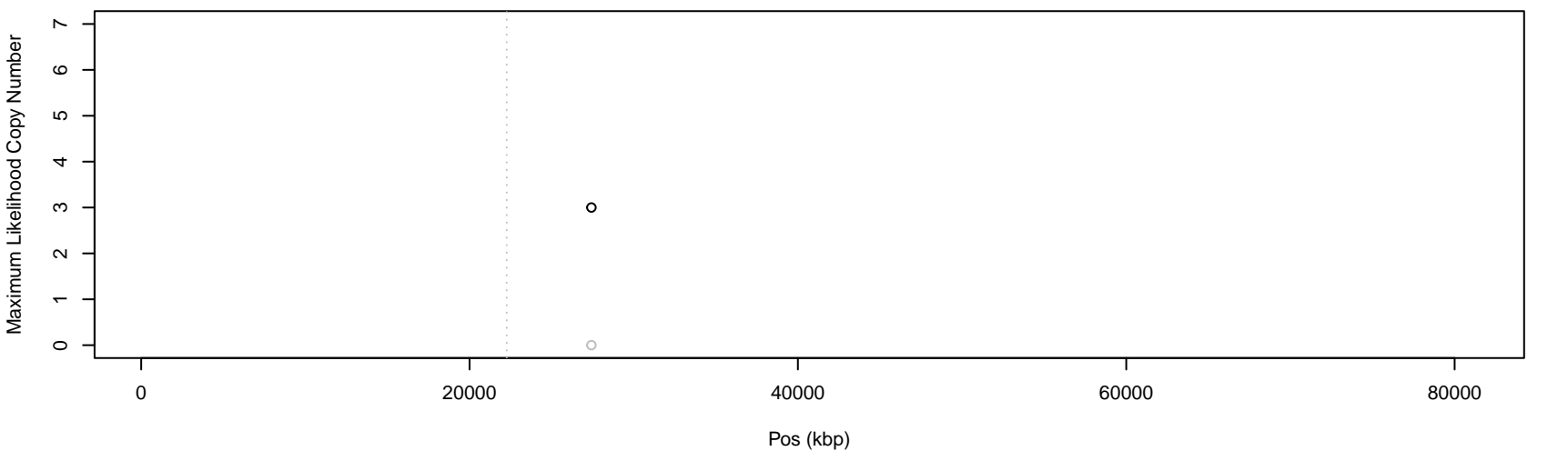
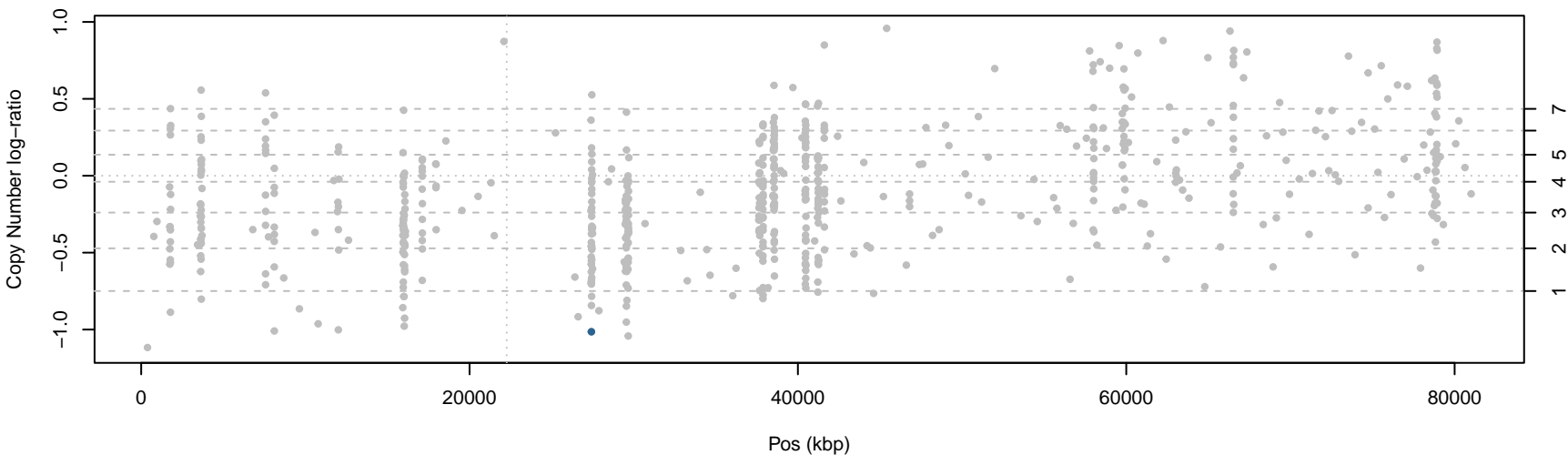
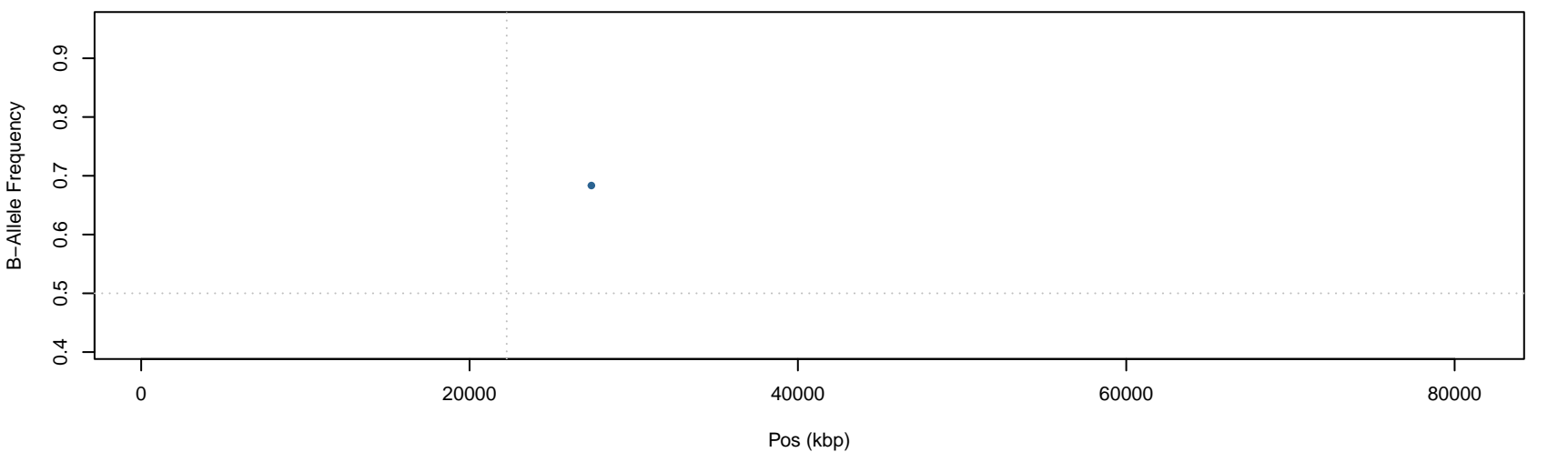


Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 16

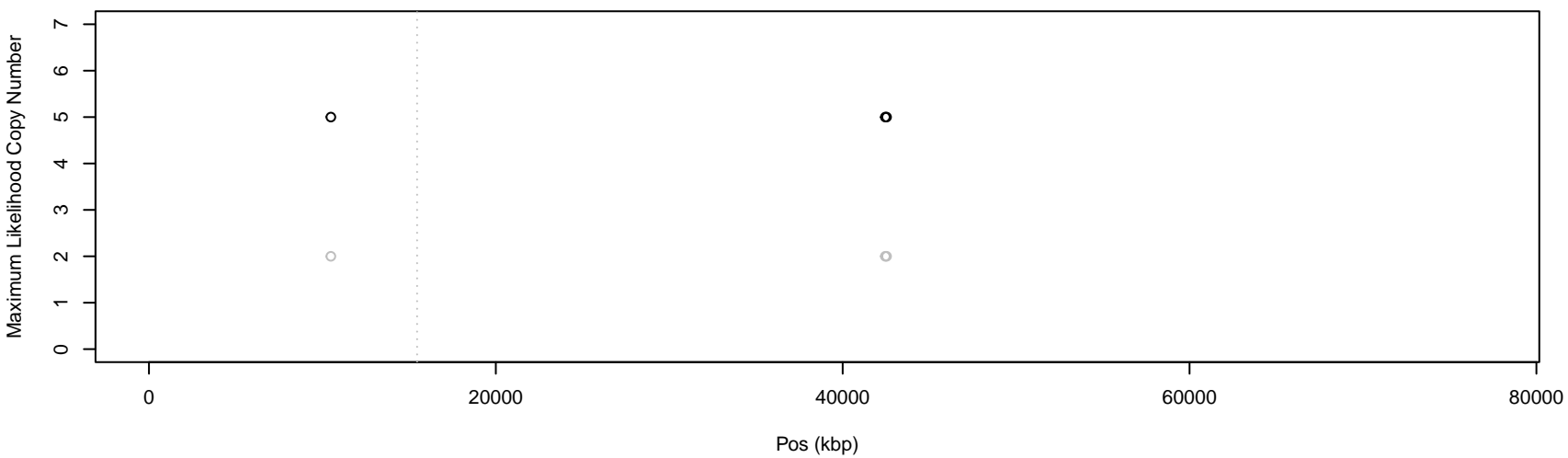
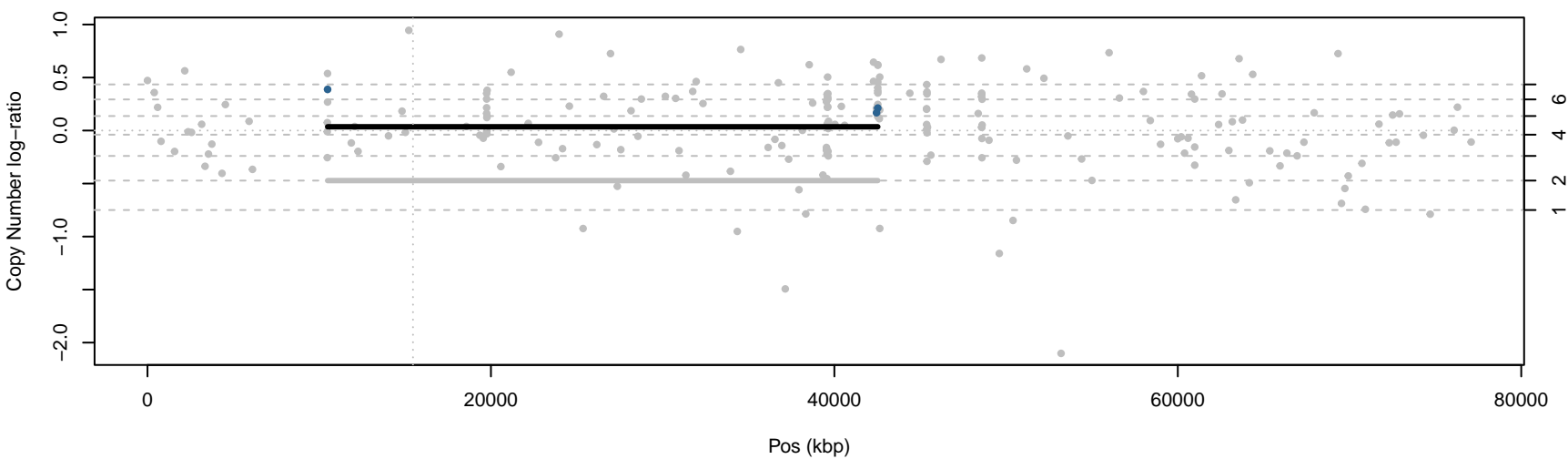
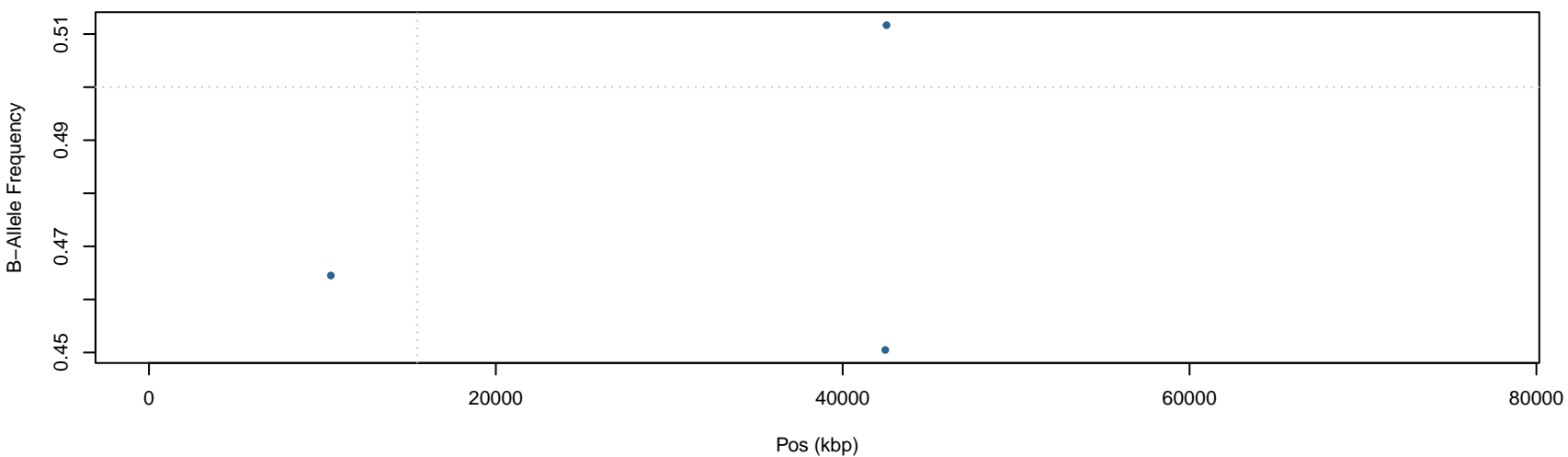




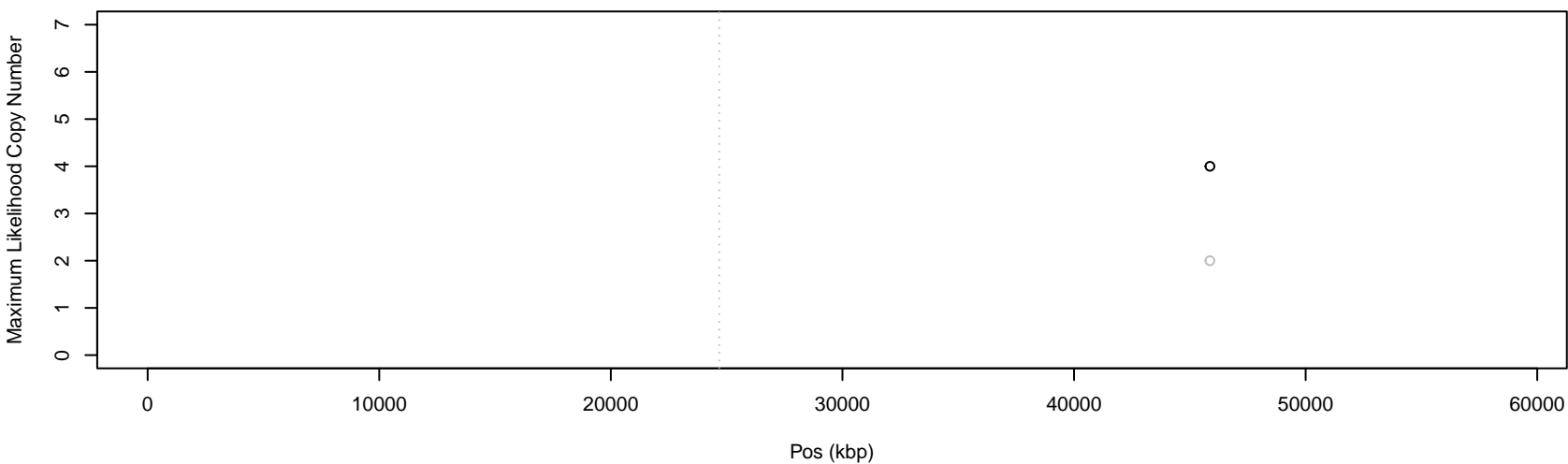
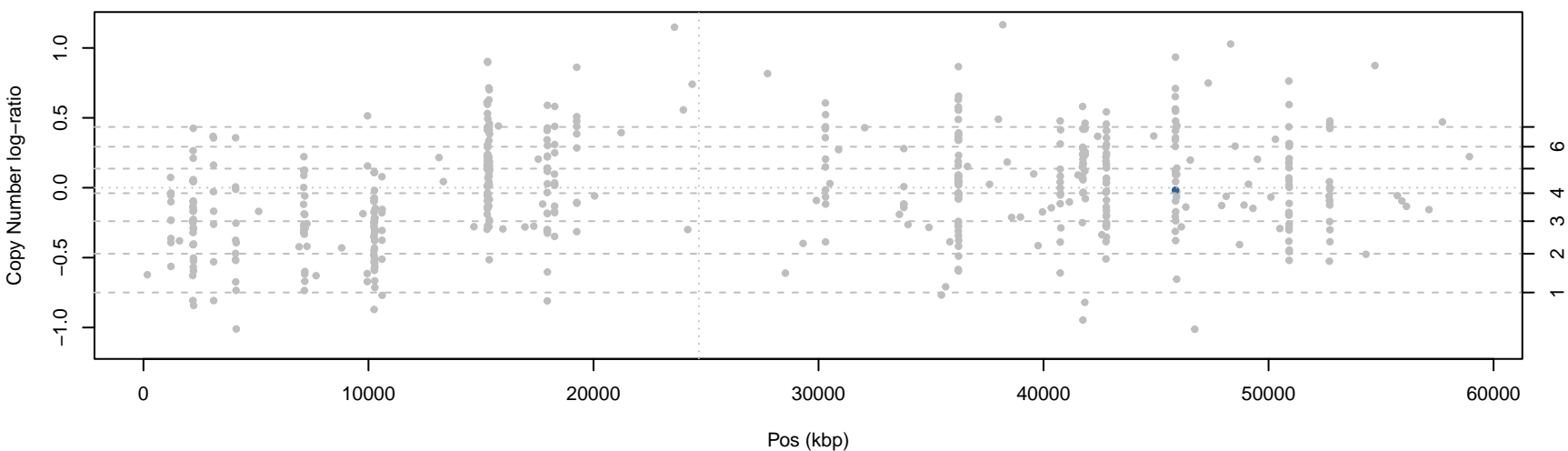
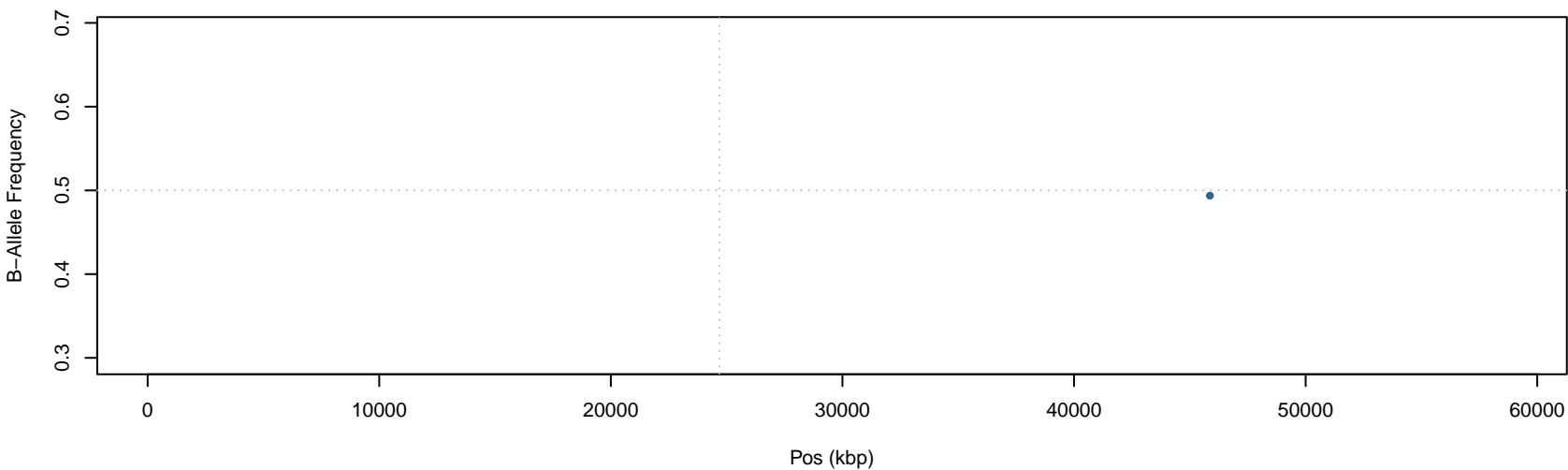
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 17



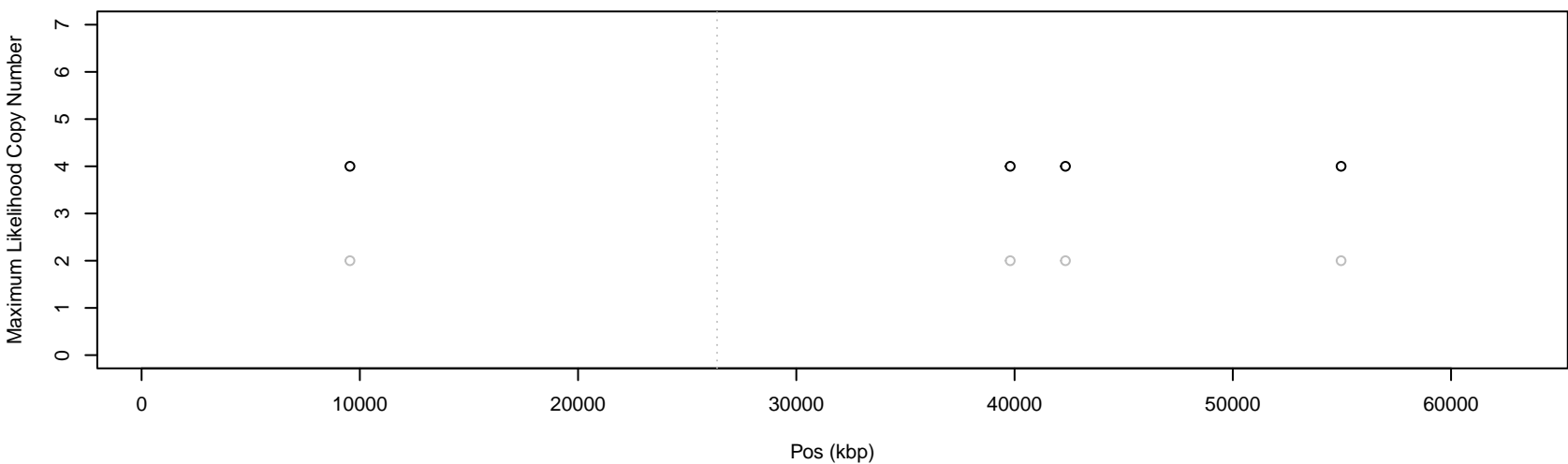
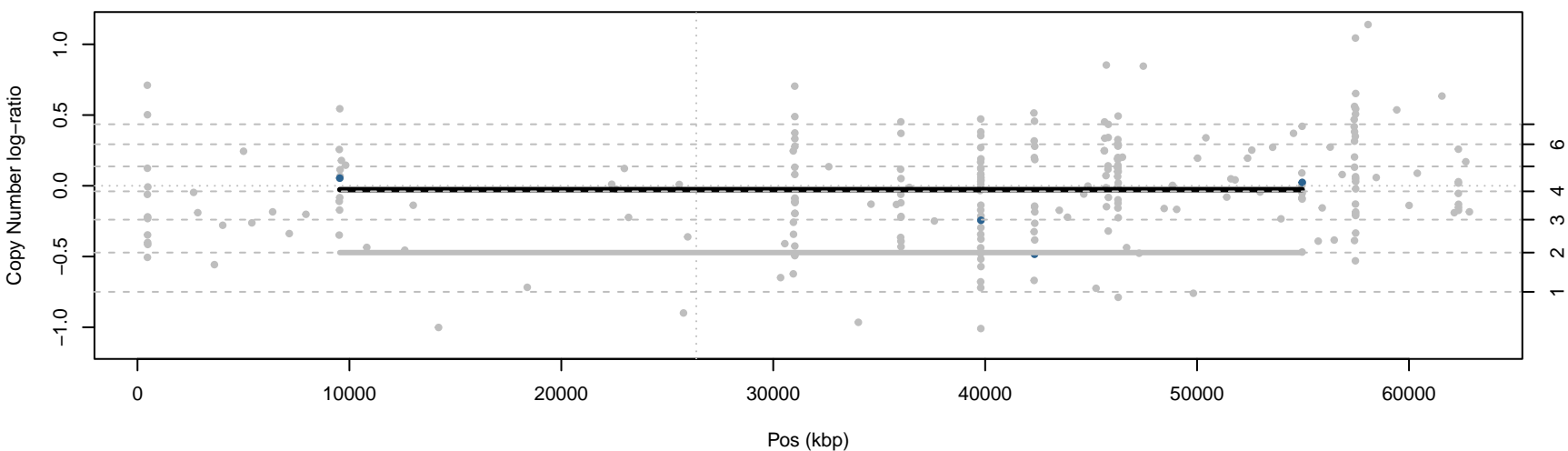
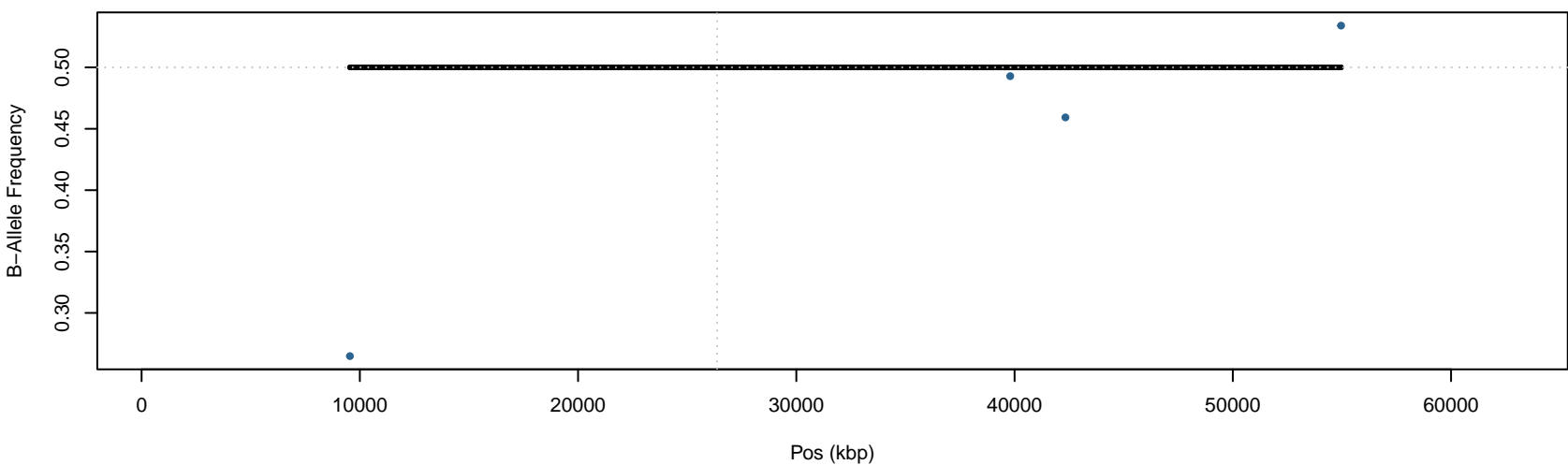
Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 18



Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 19



Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 20



Purity: 0.35 Tumor ploidy: 4.212 SNV log-likelihood: -122.56 GoF: 89.6% Mean coverage: 296;515 Chromosome: 22

