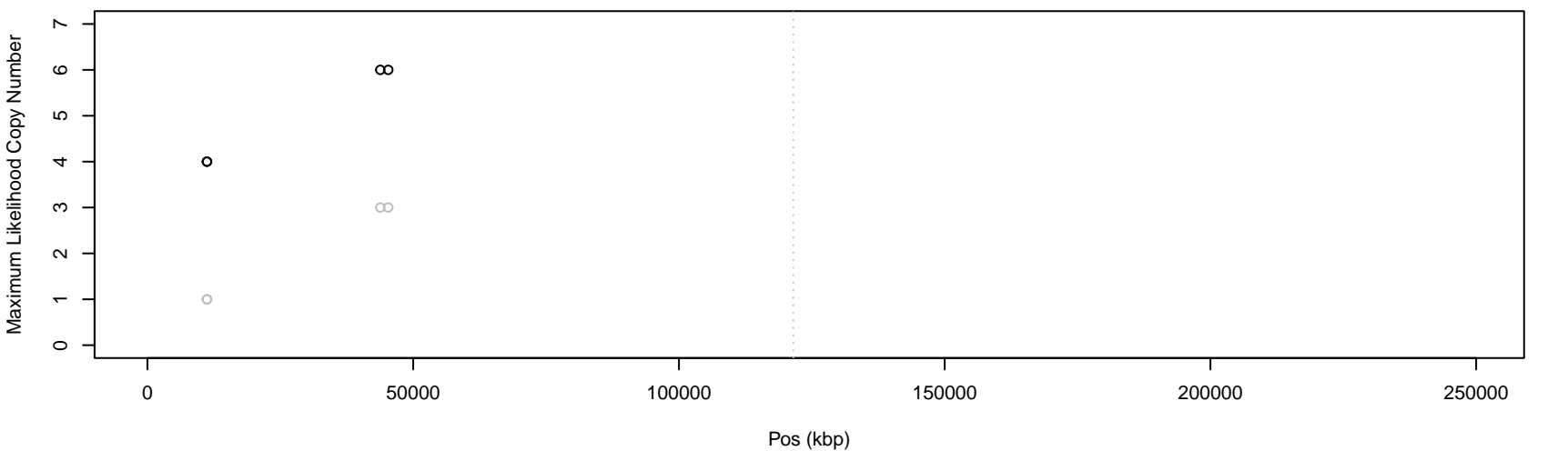
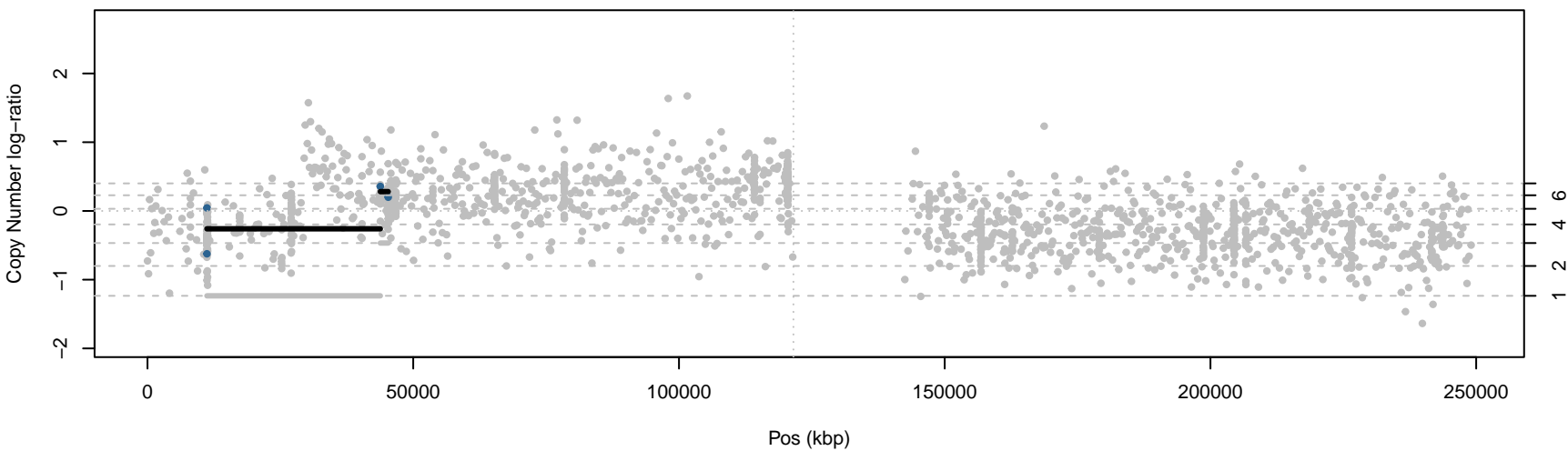
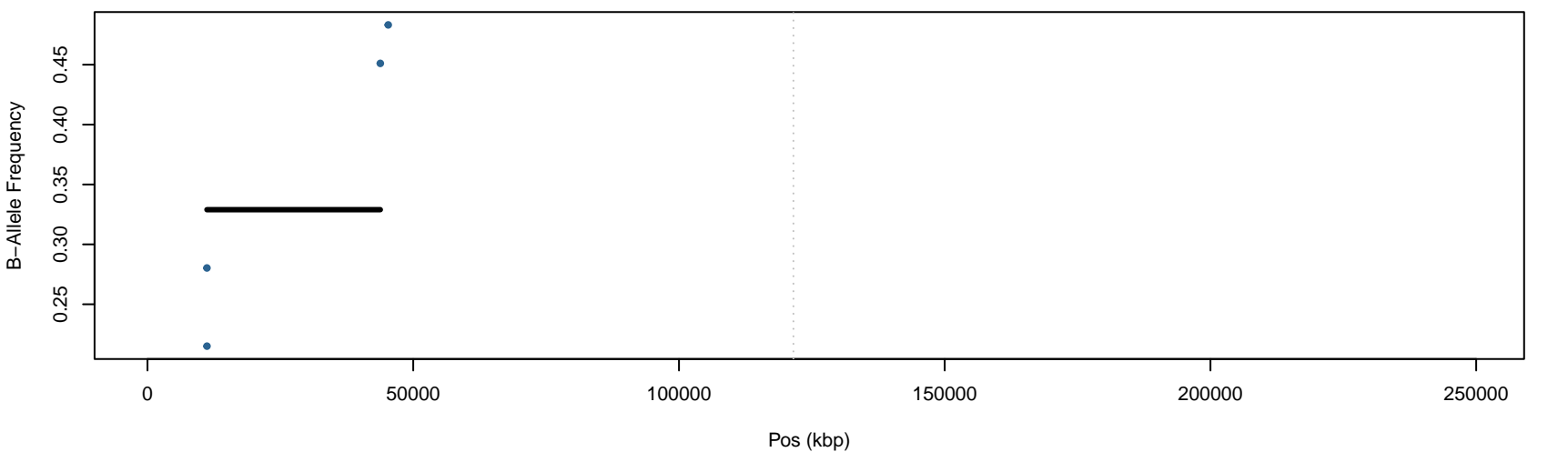
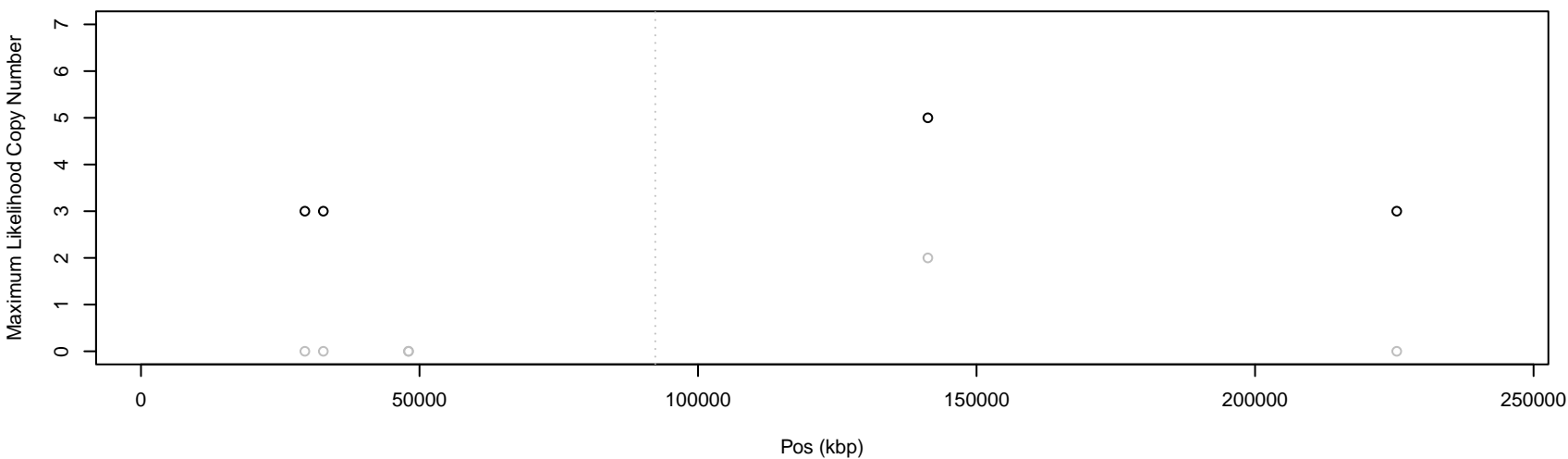
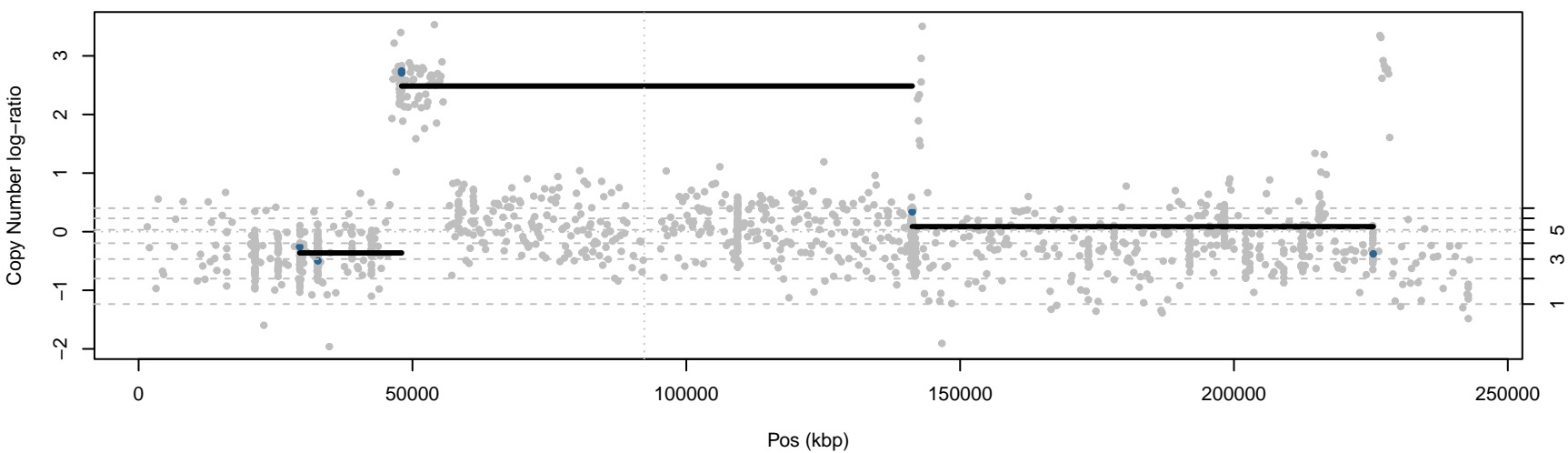
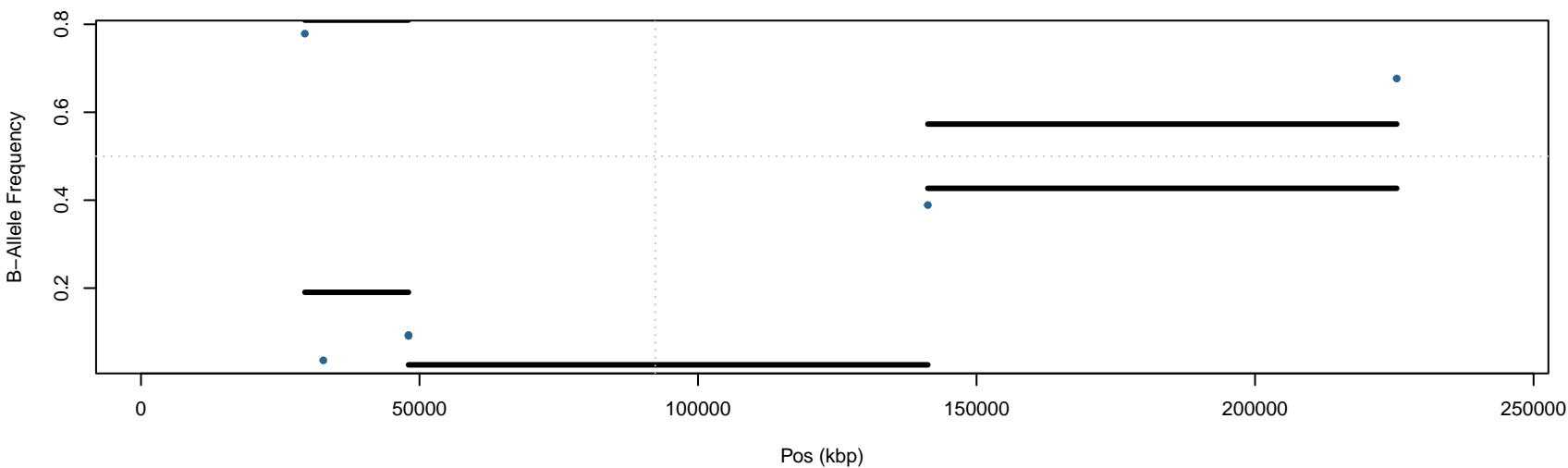


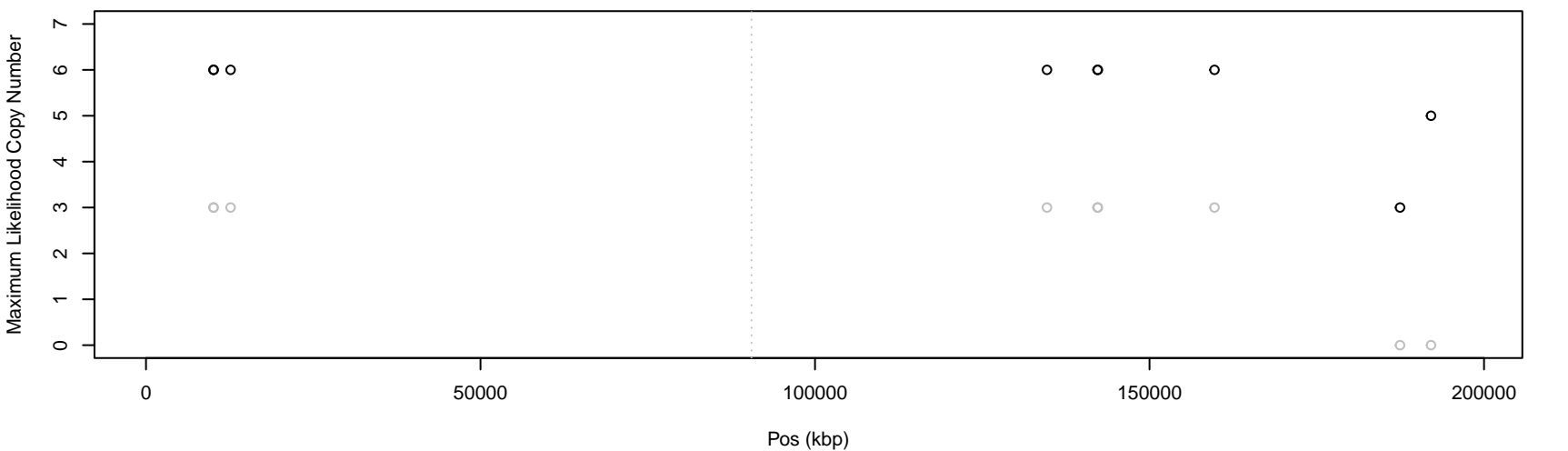
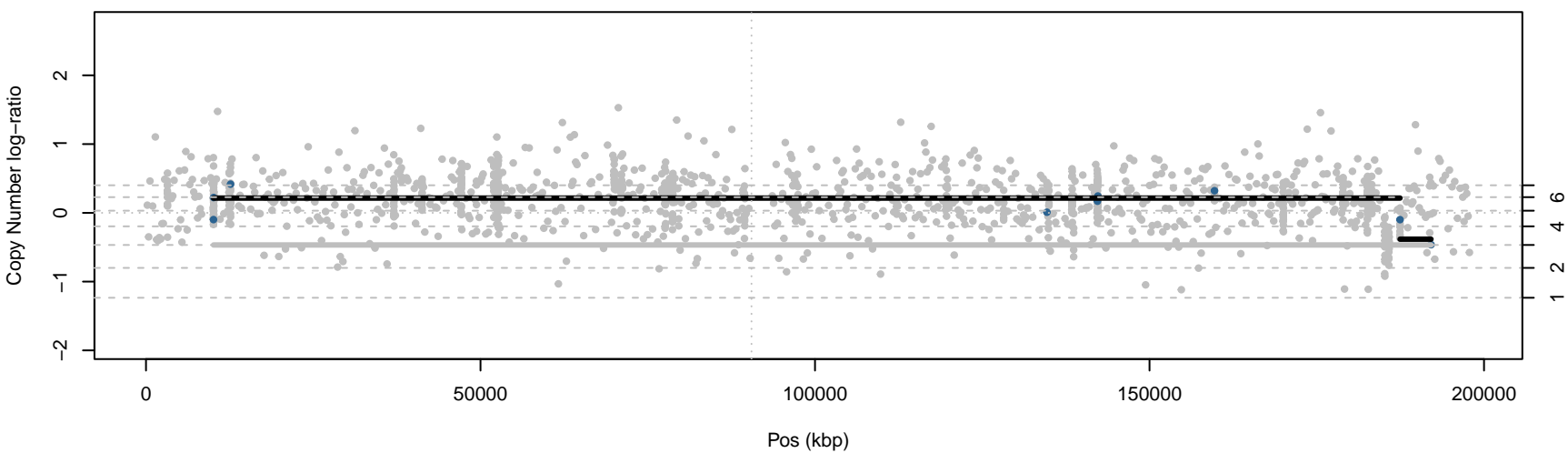
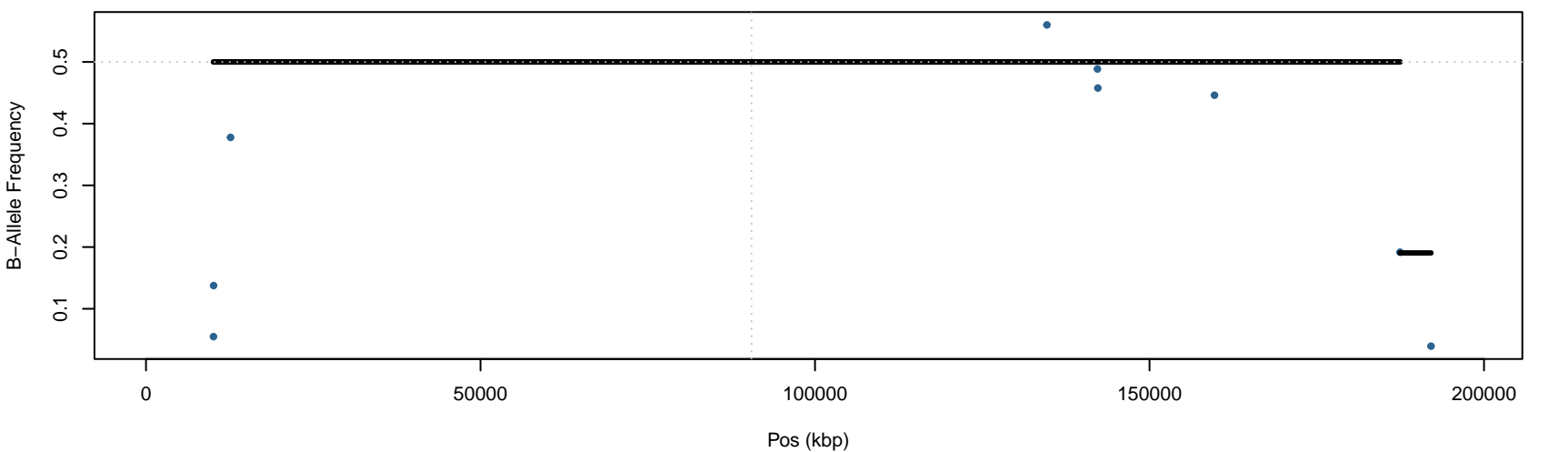
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 1



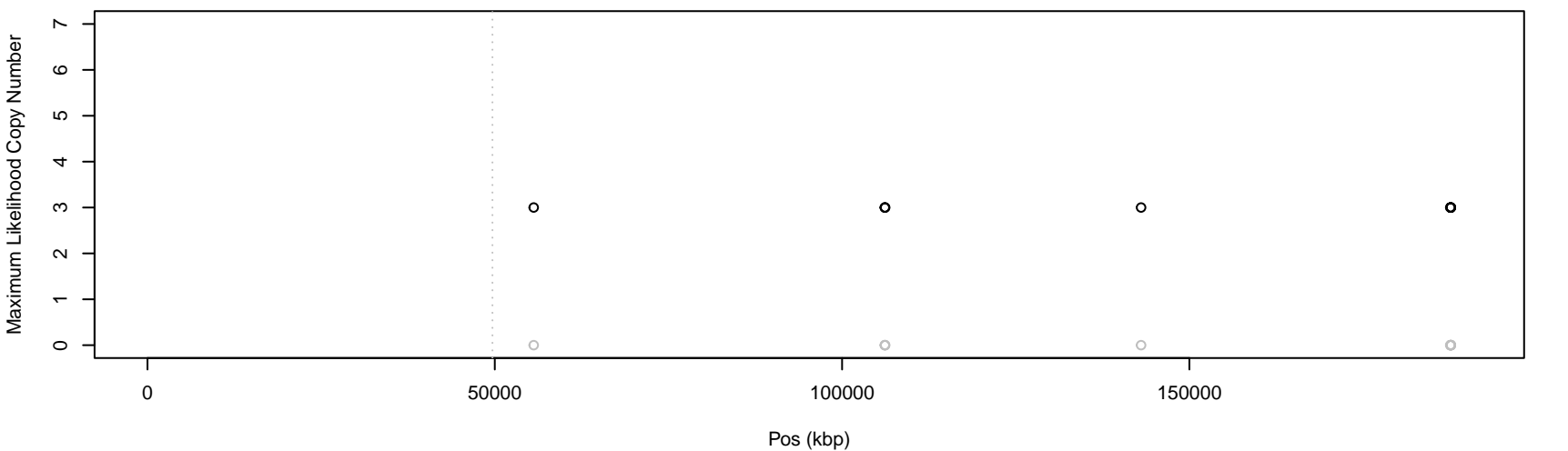
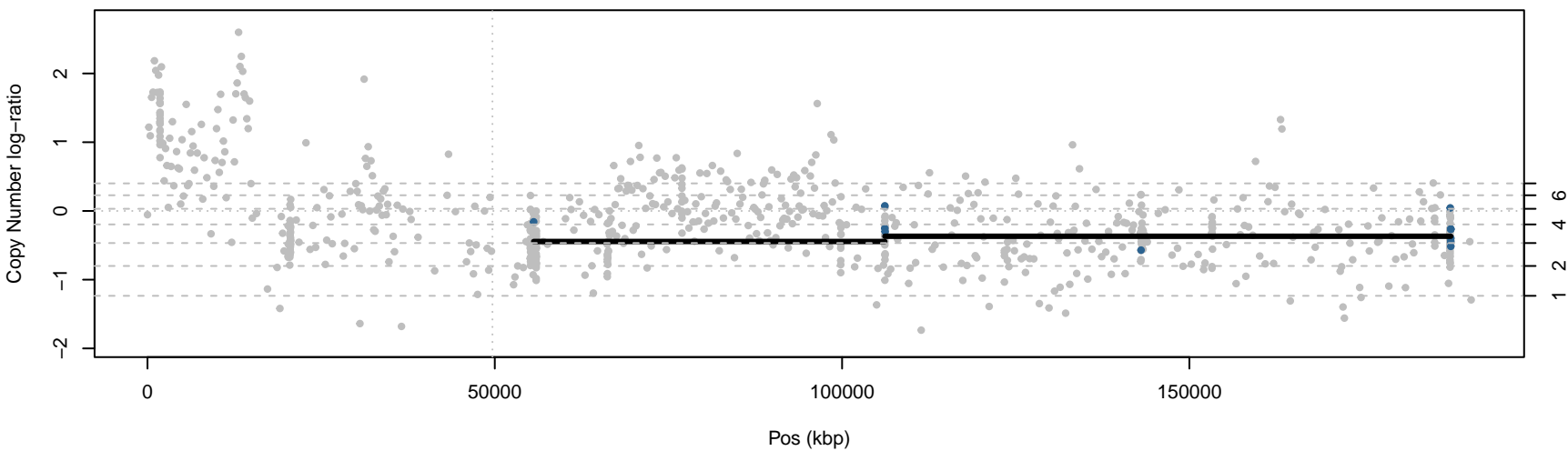
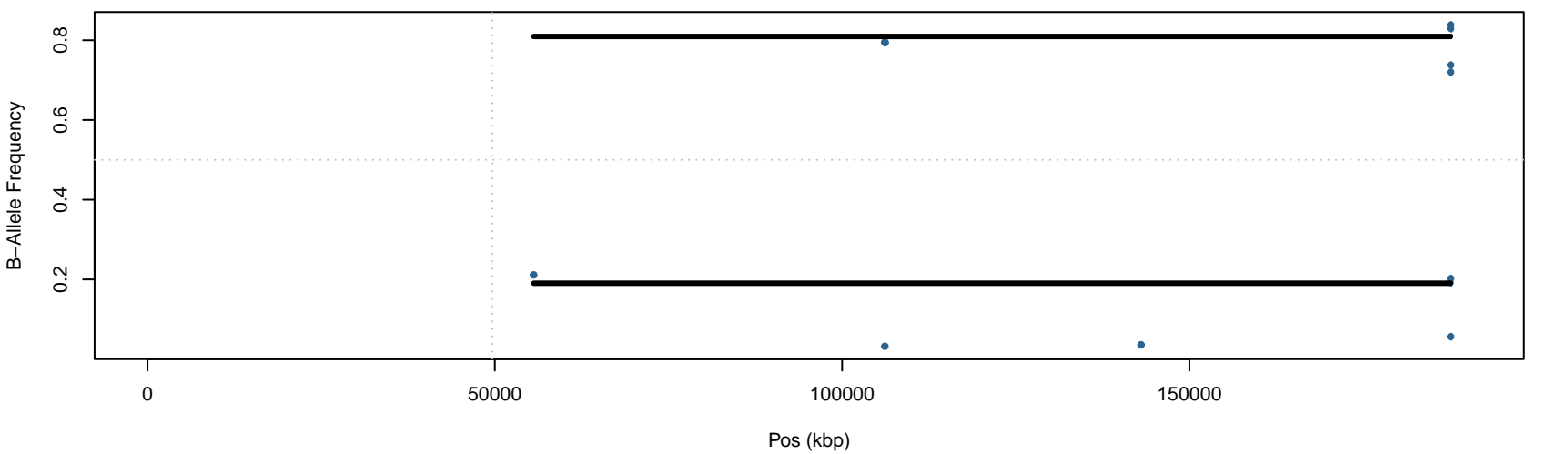
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 2



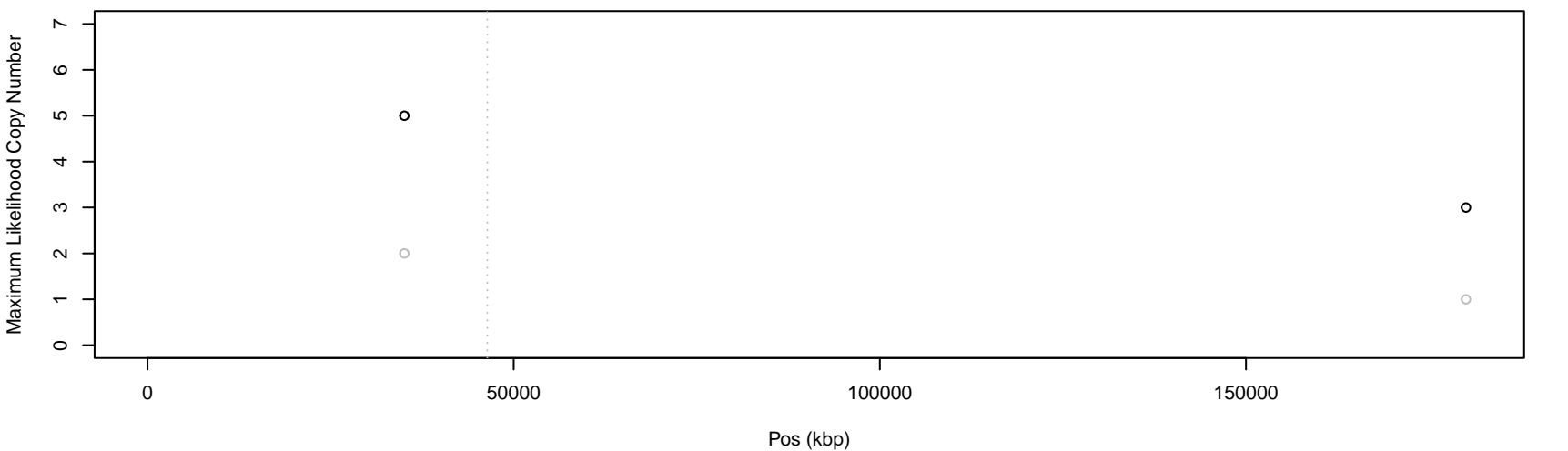
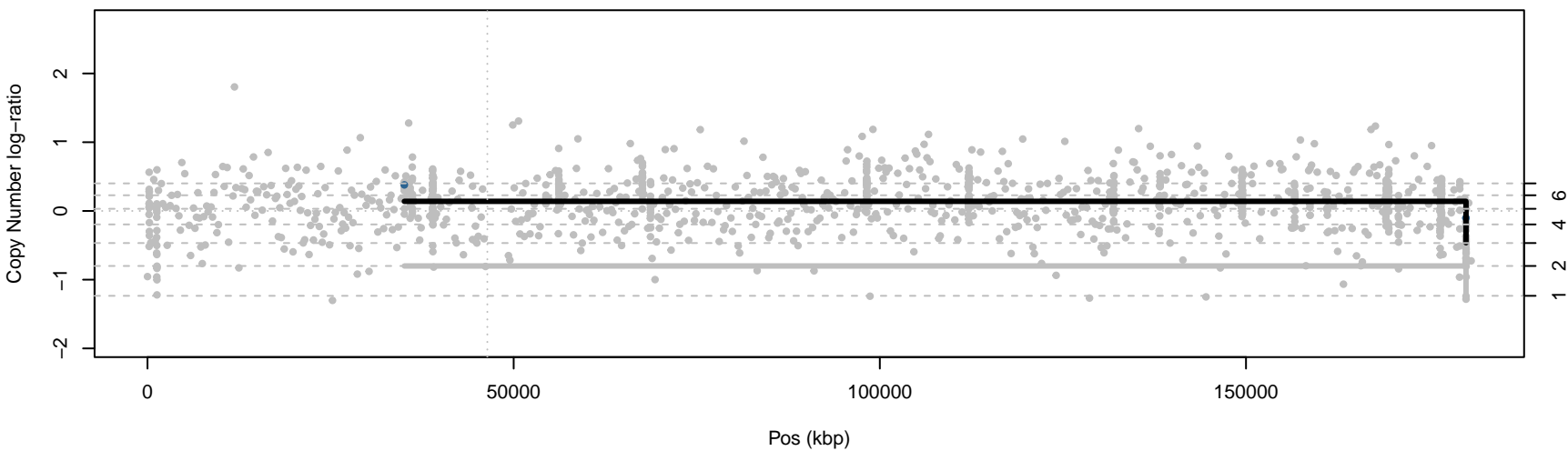
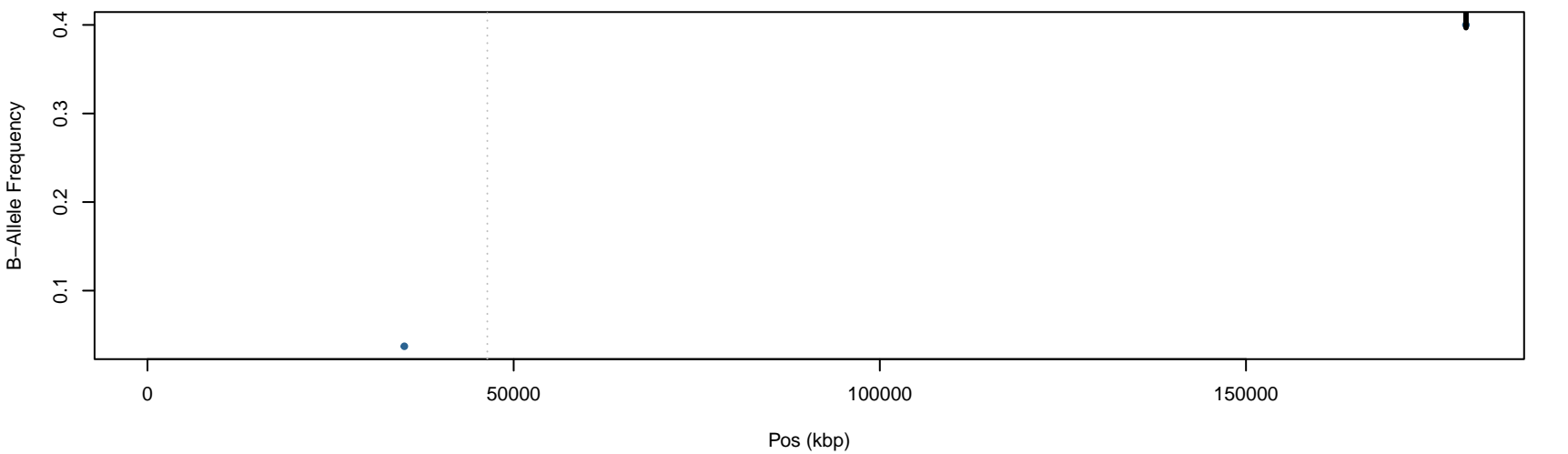
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 3



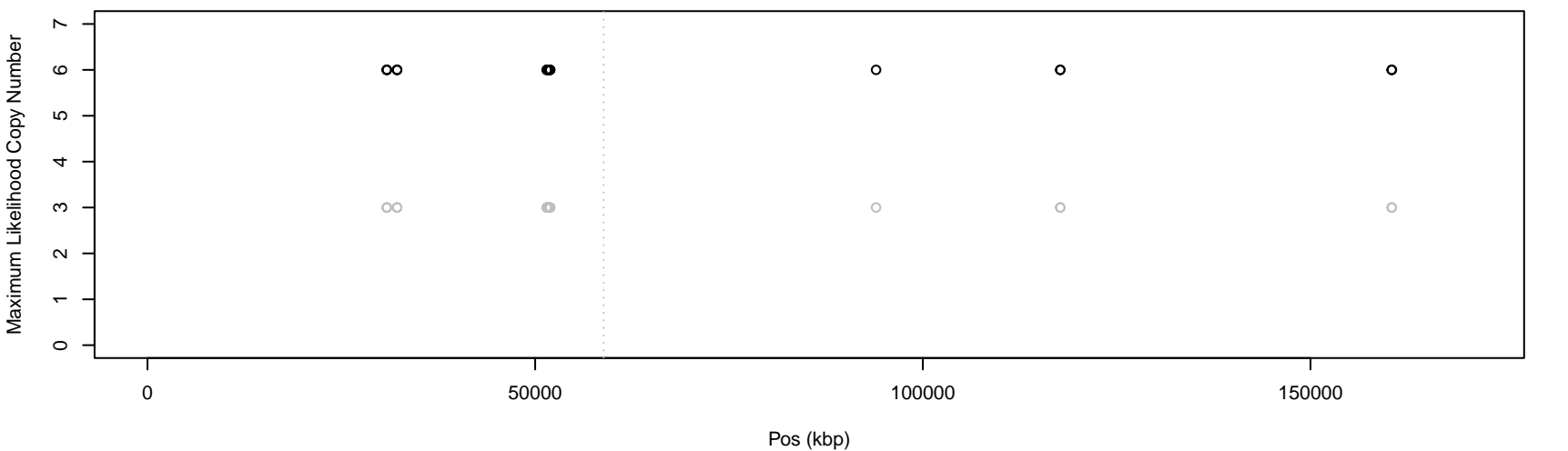
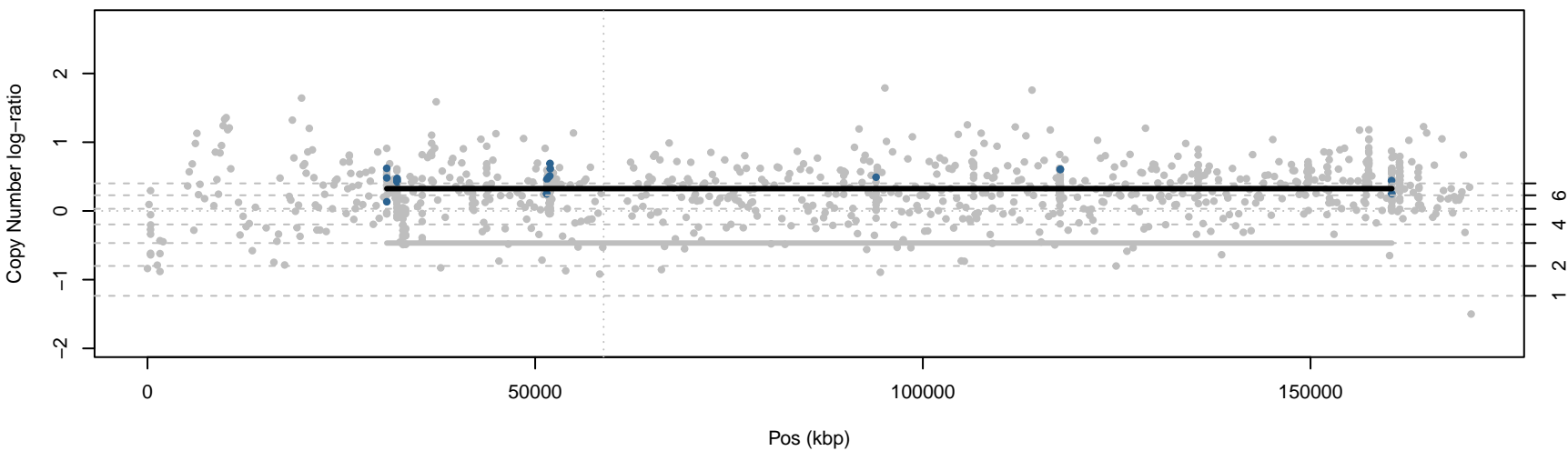
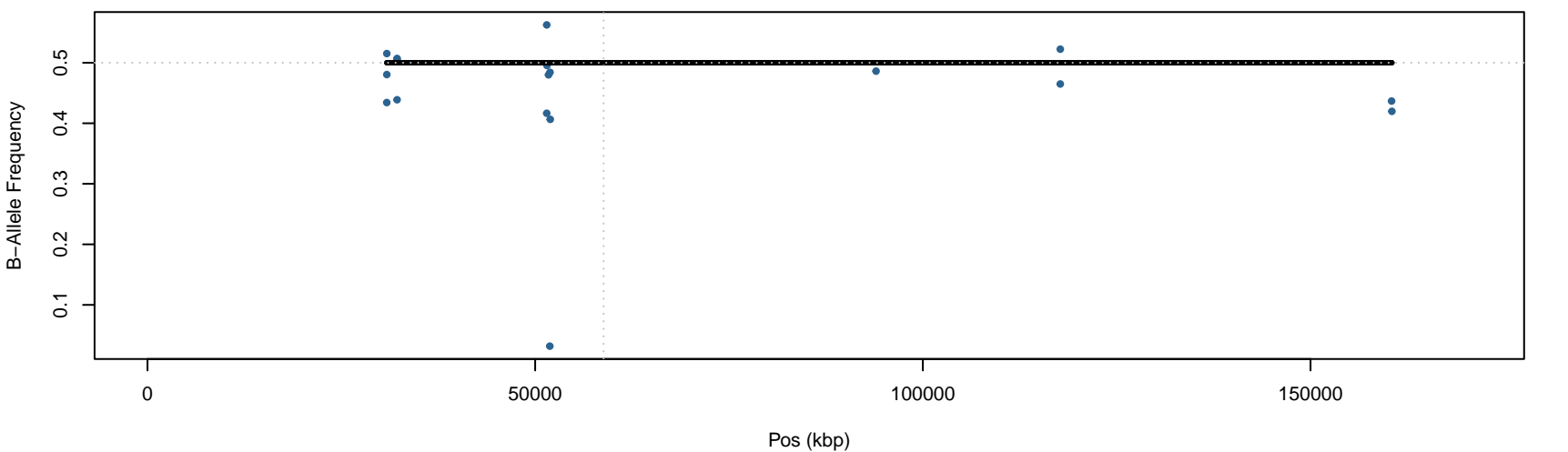
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 4



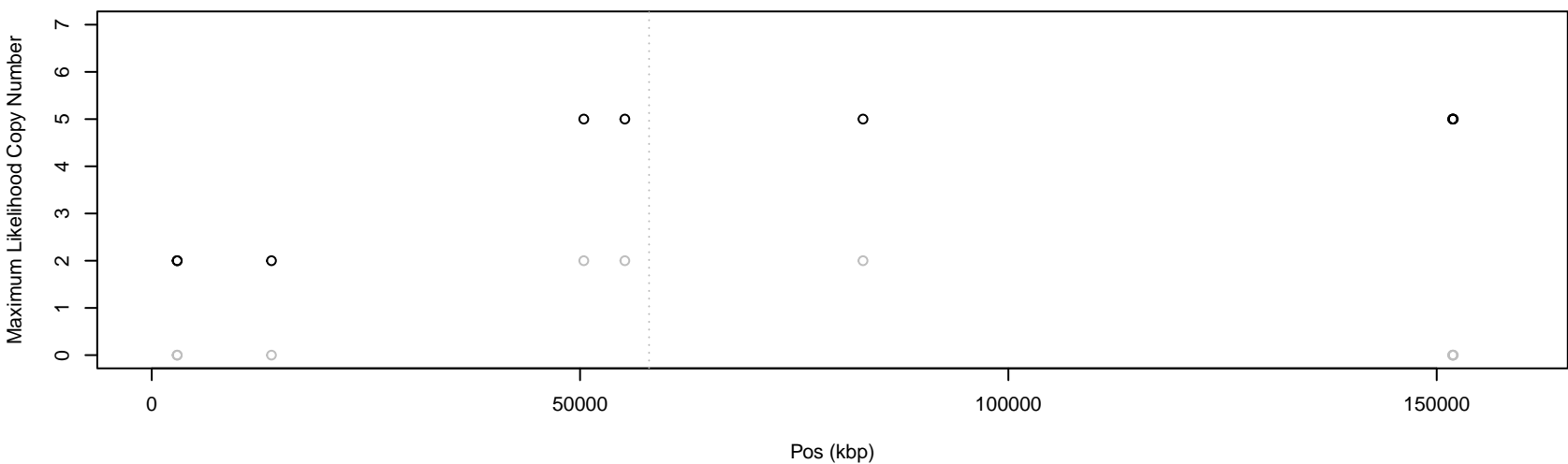
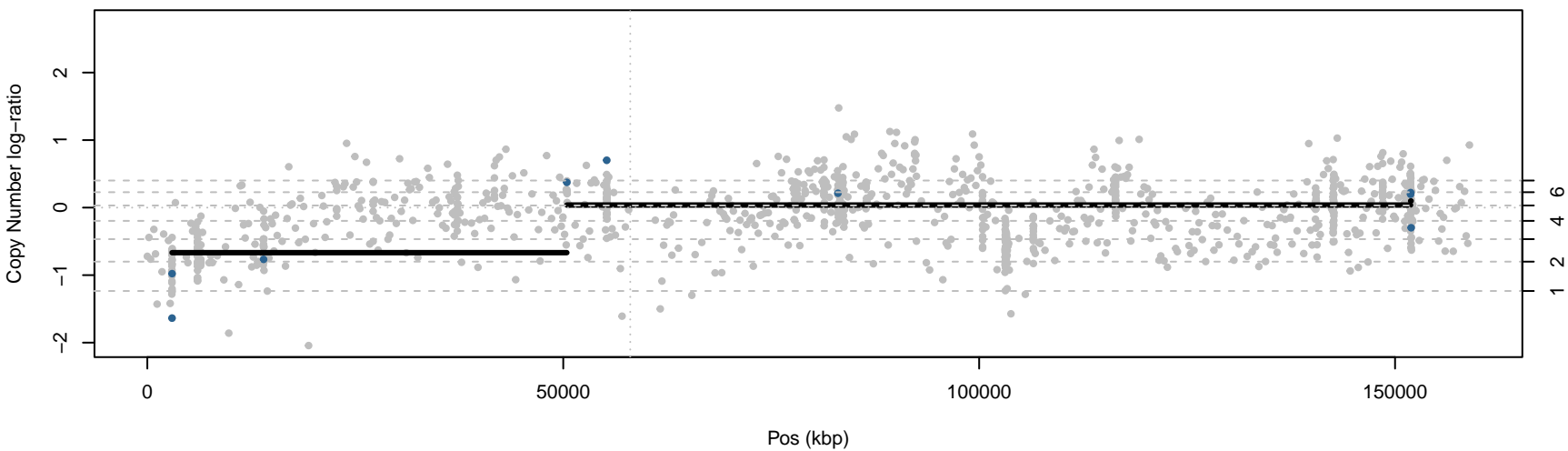
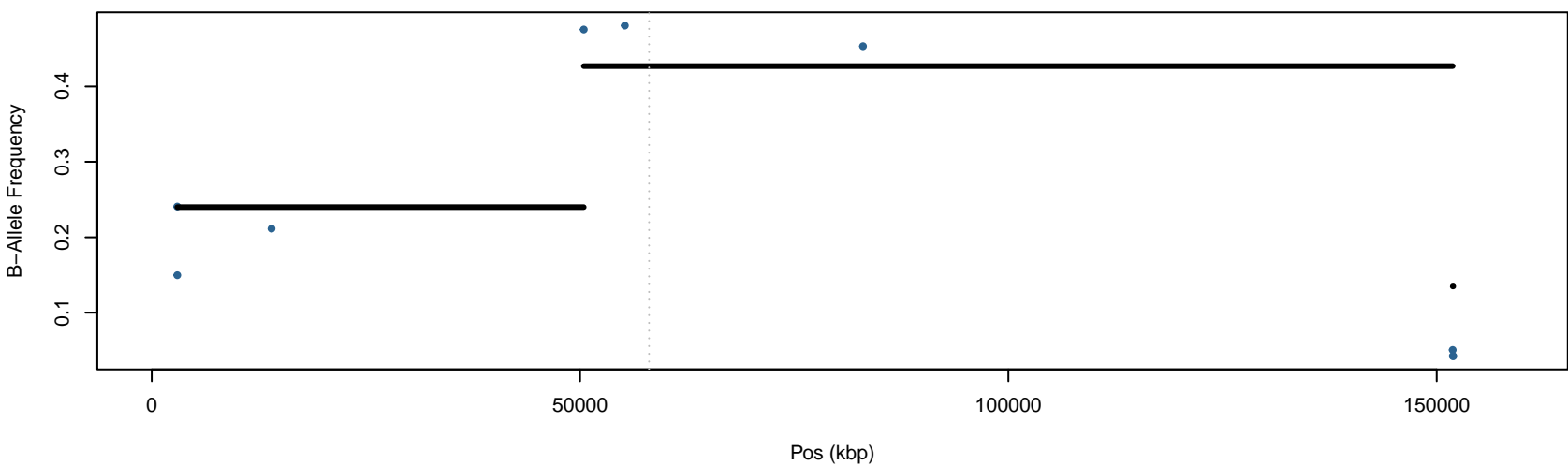
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 5



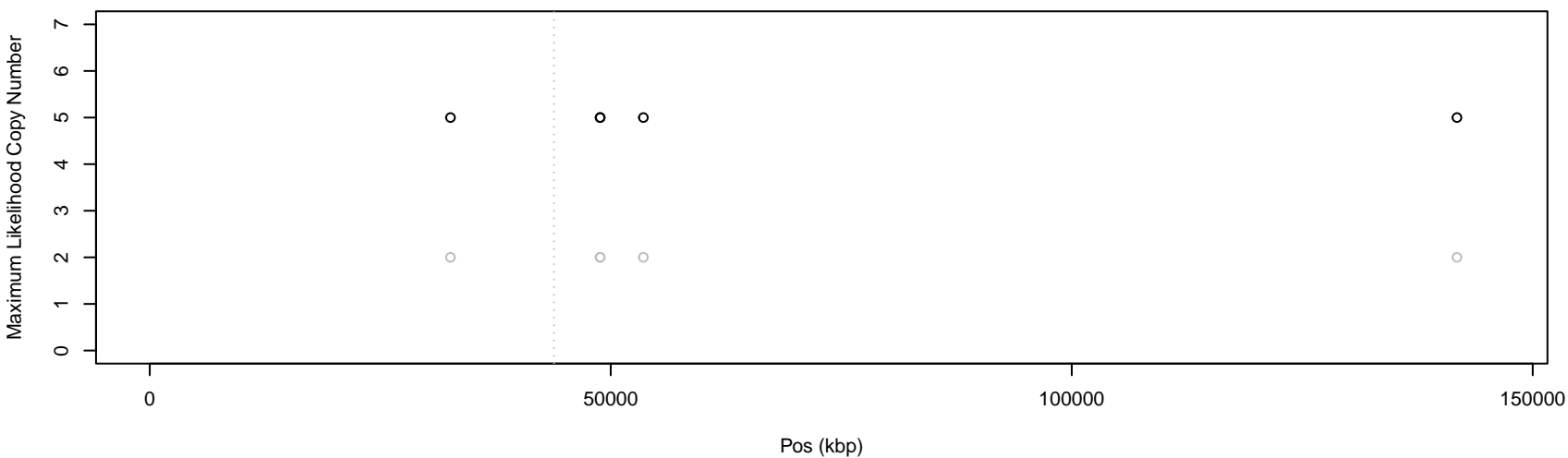
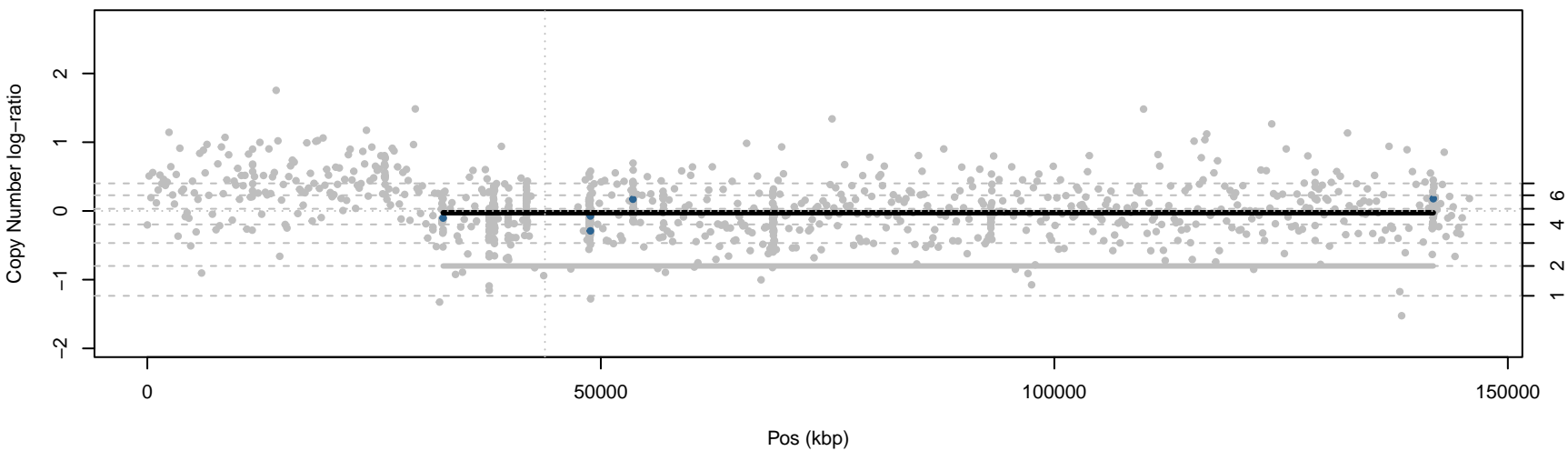
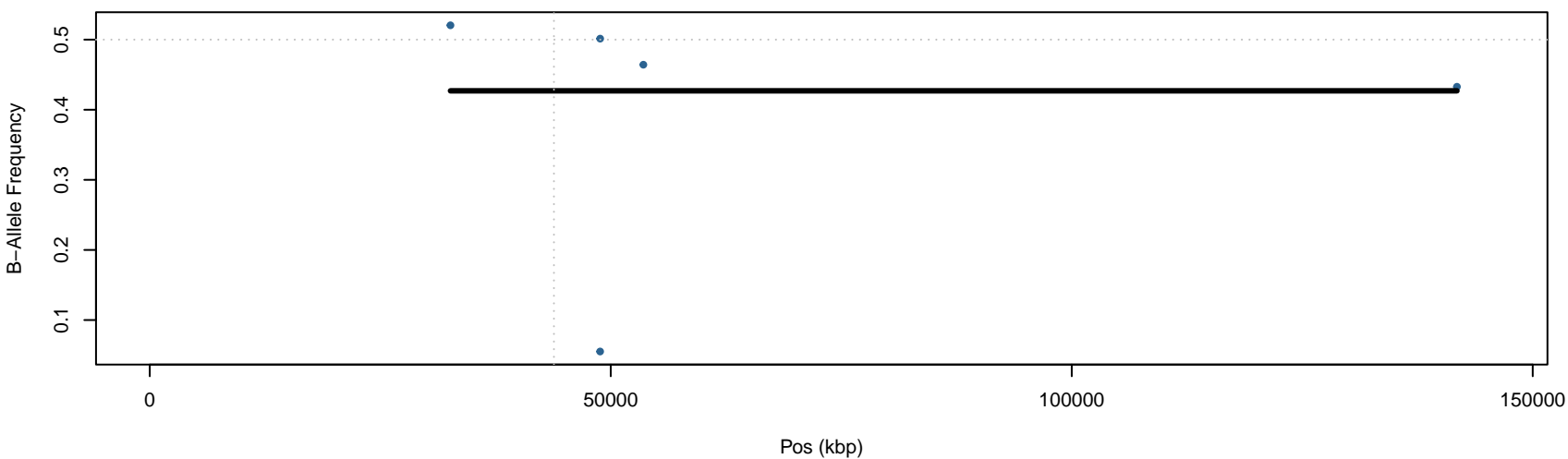
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 6



Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 7

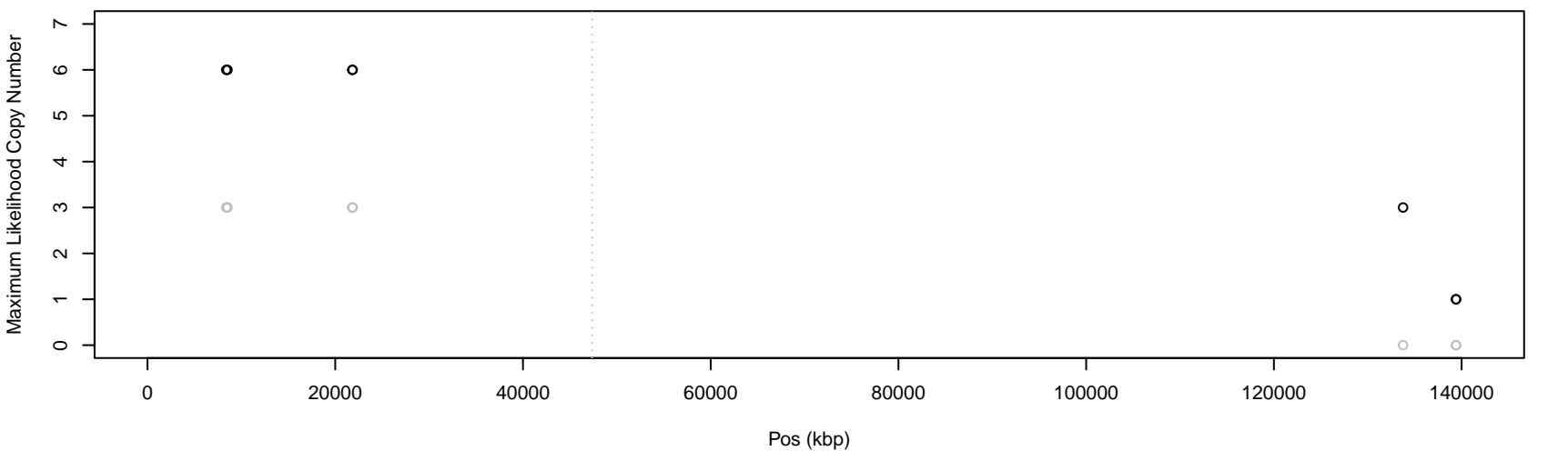
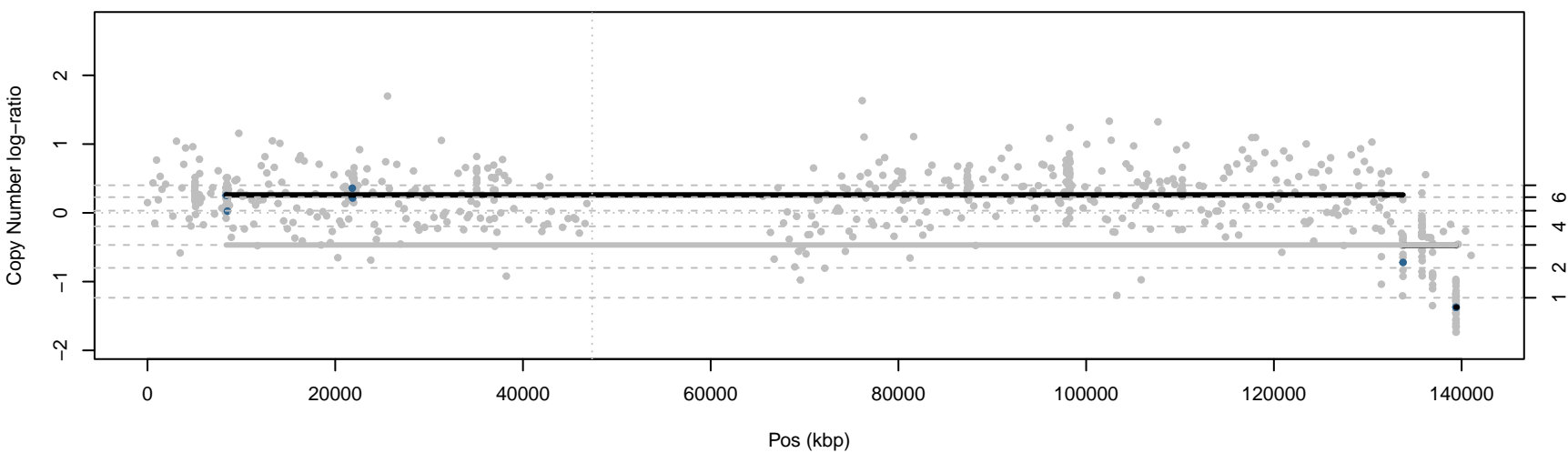
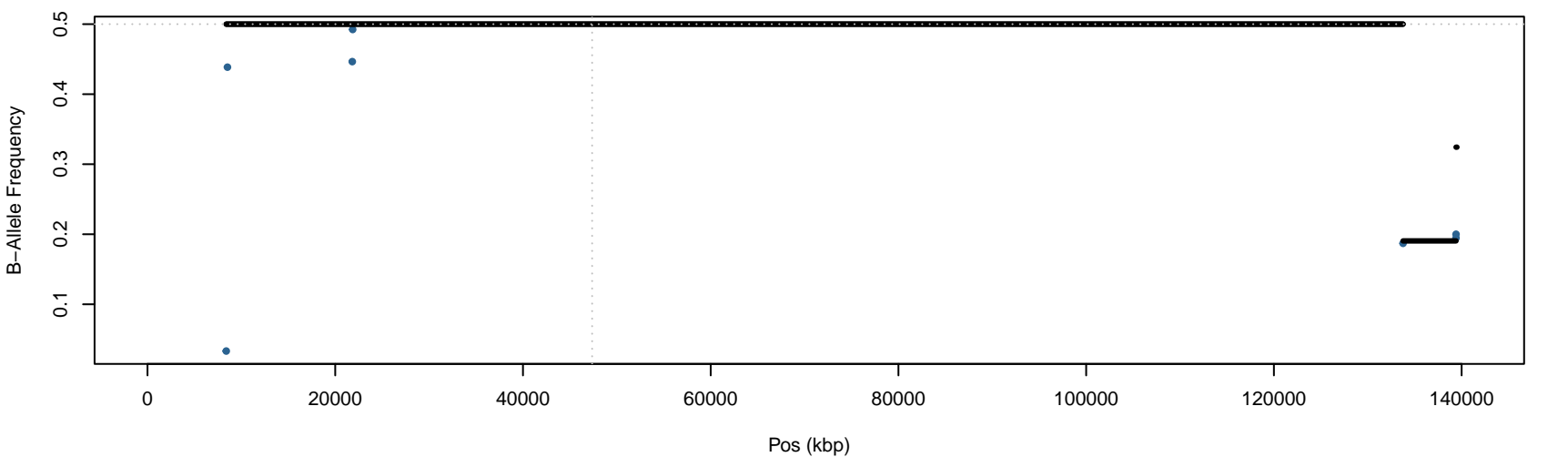


Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 8

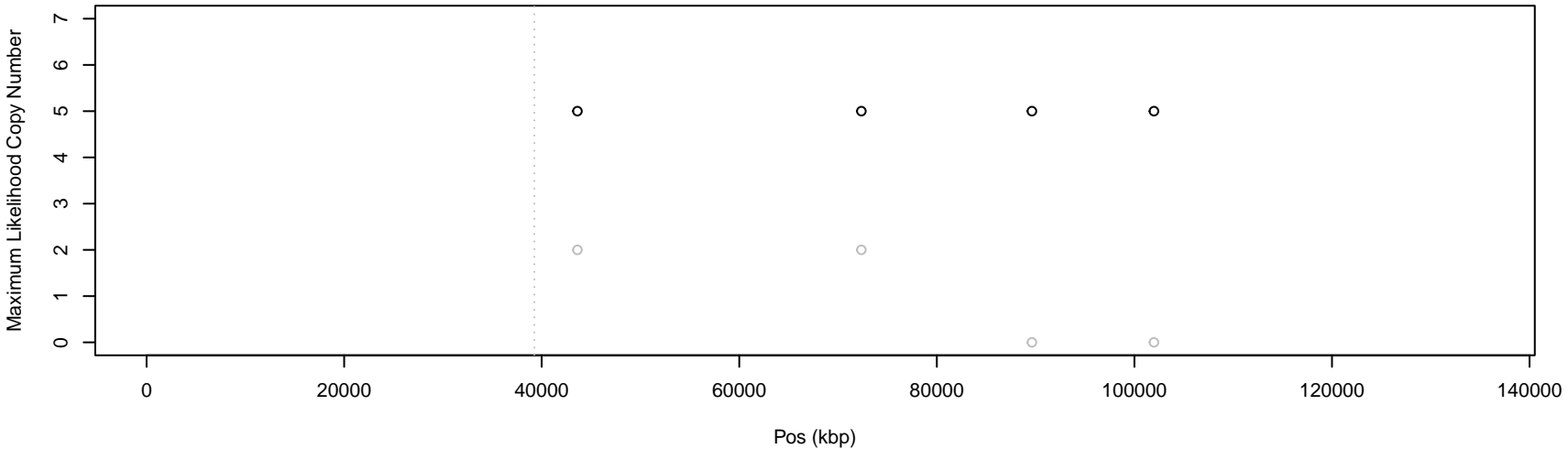
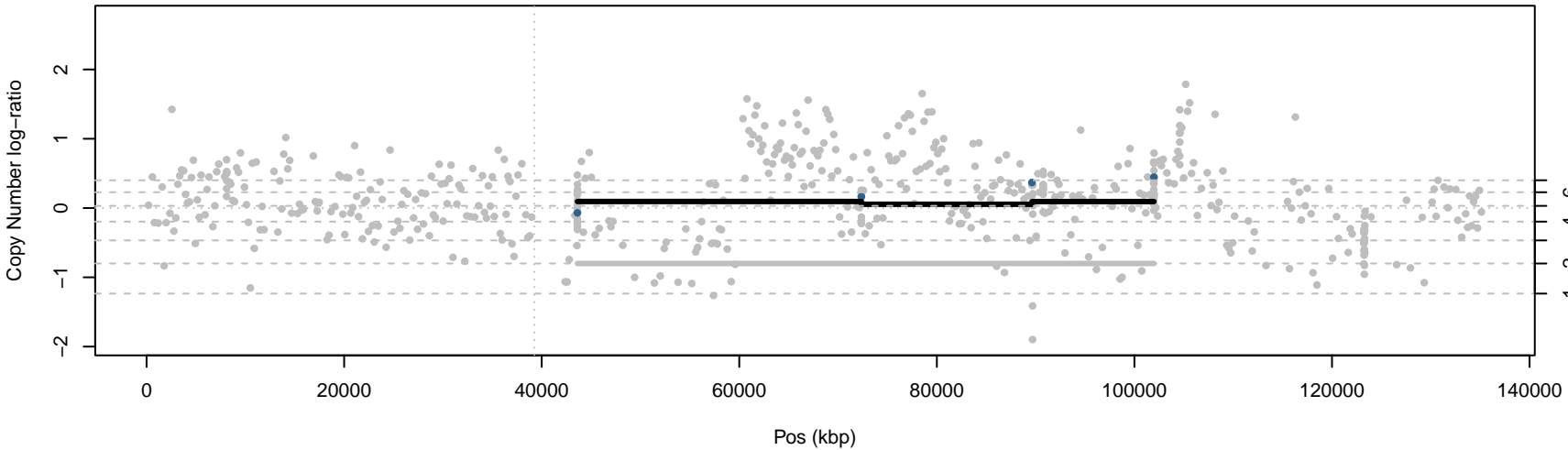
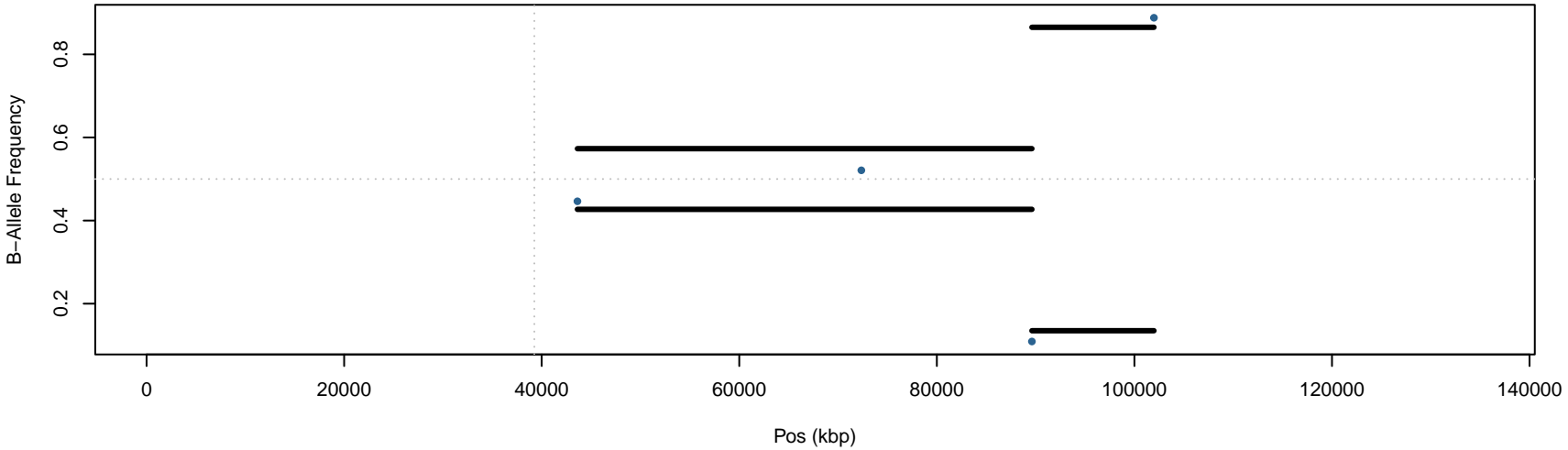




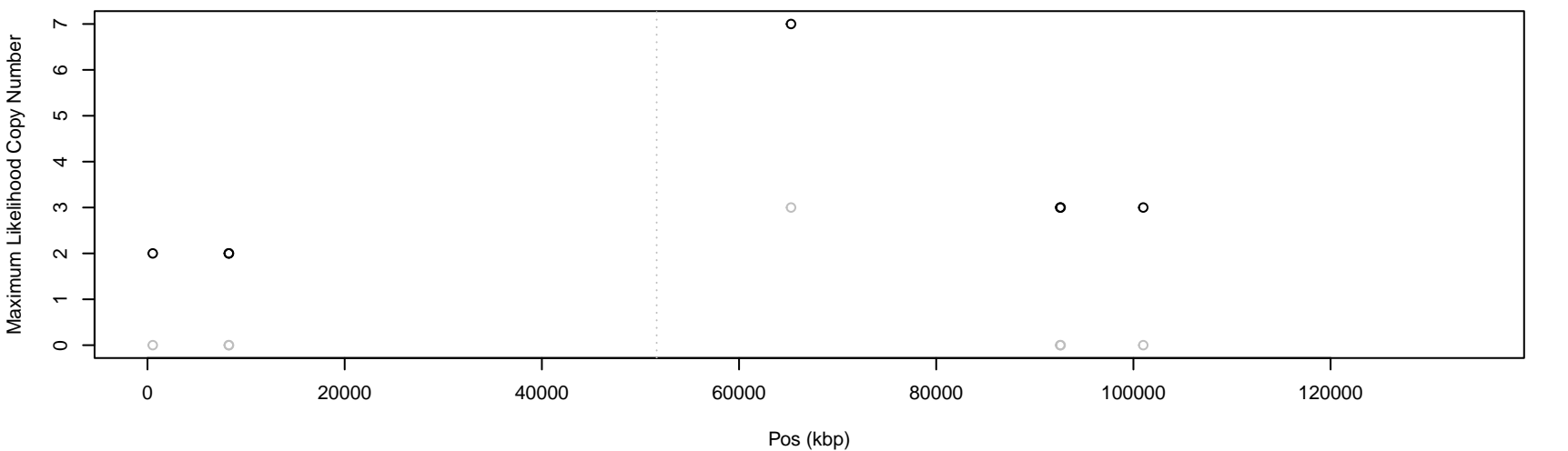
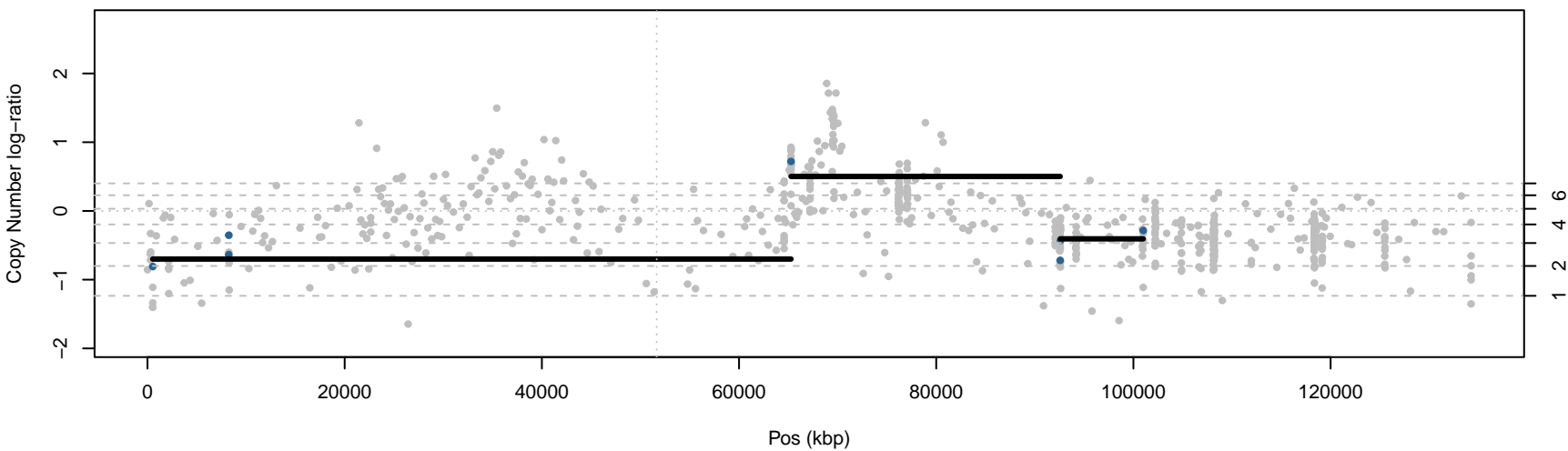
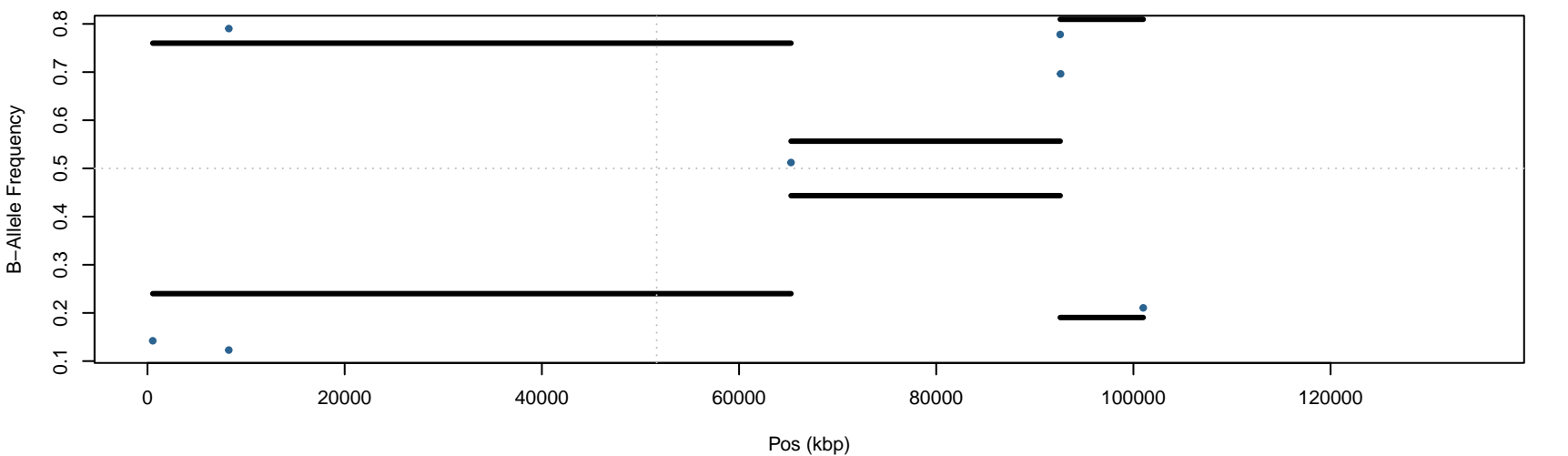
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 9



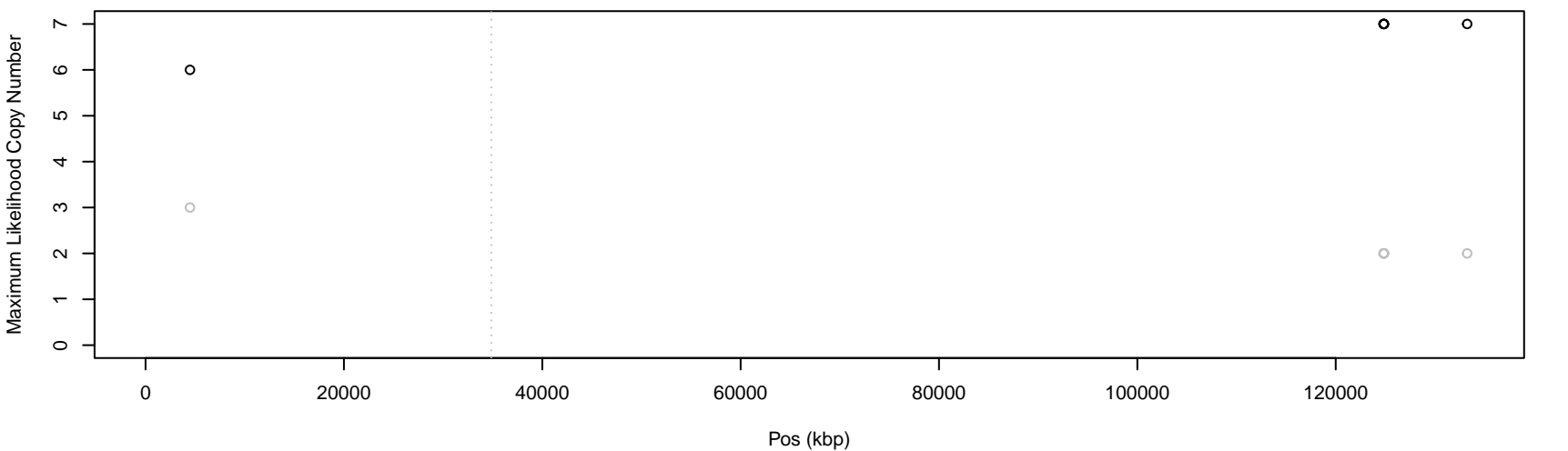
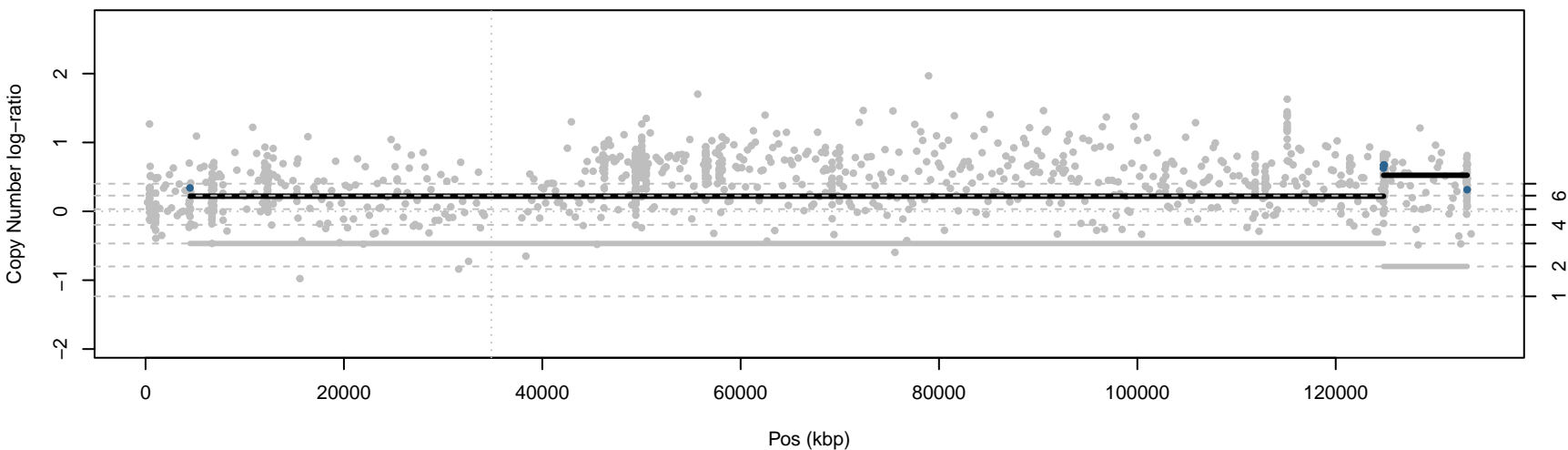
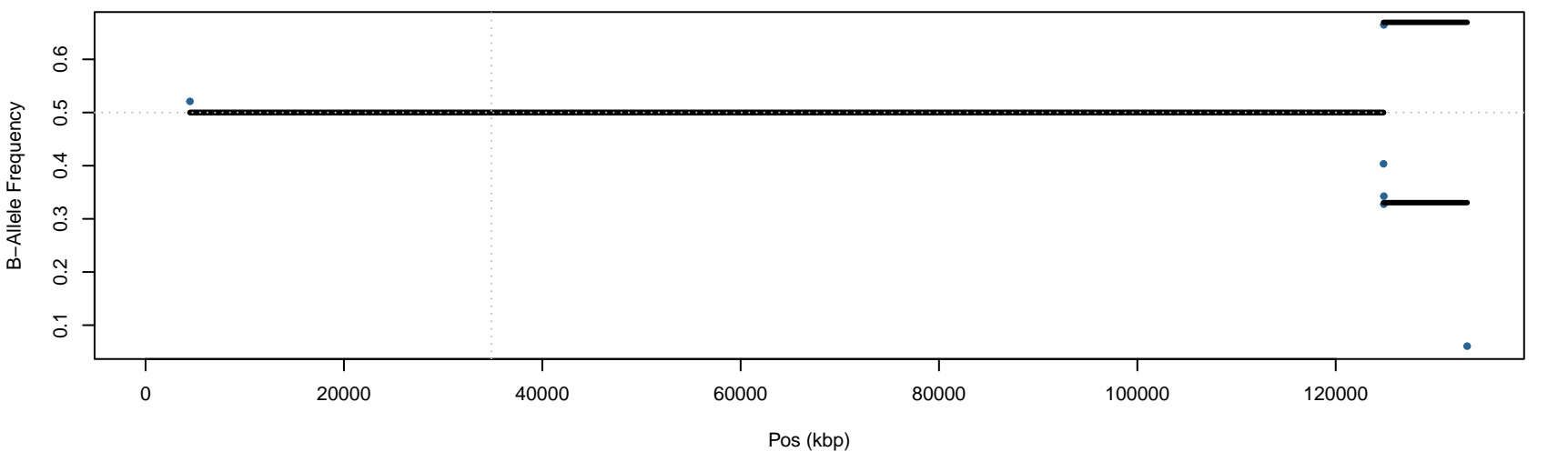
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 10



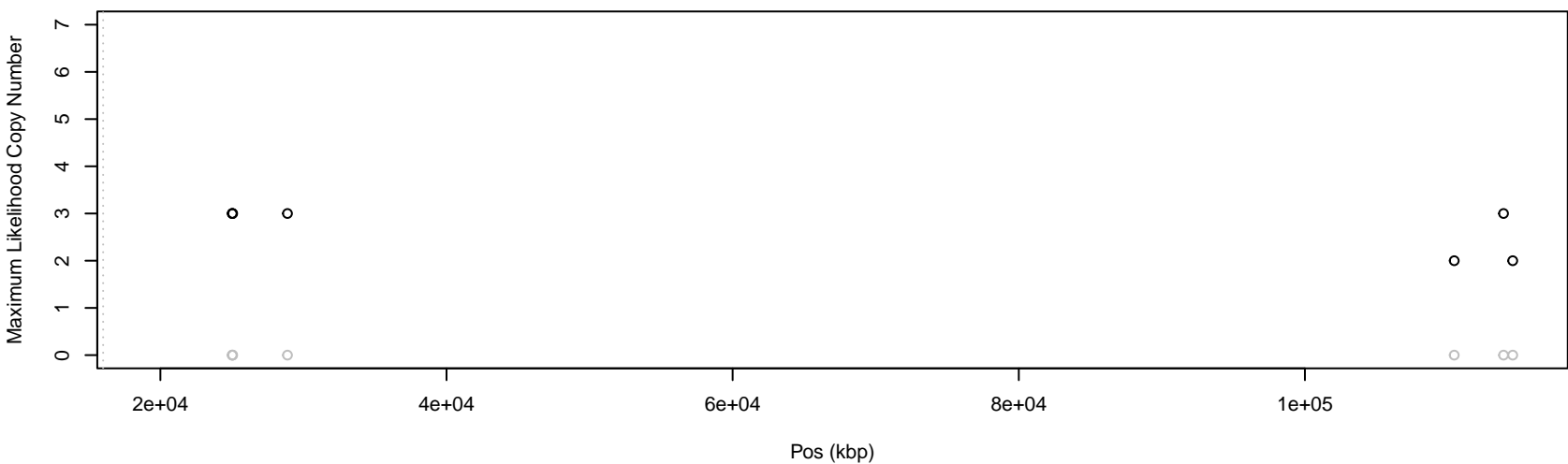
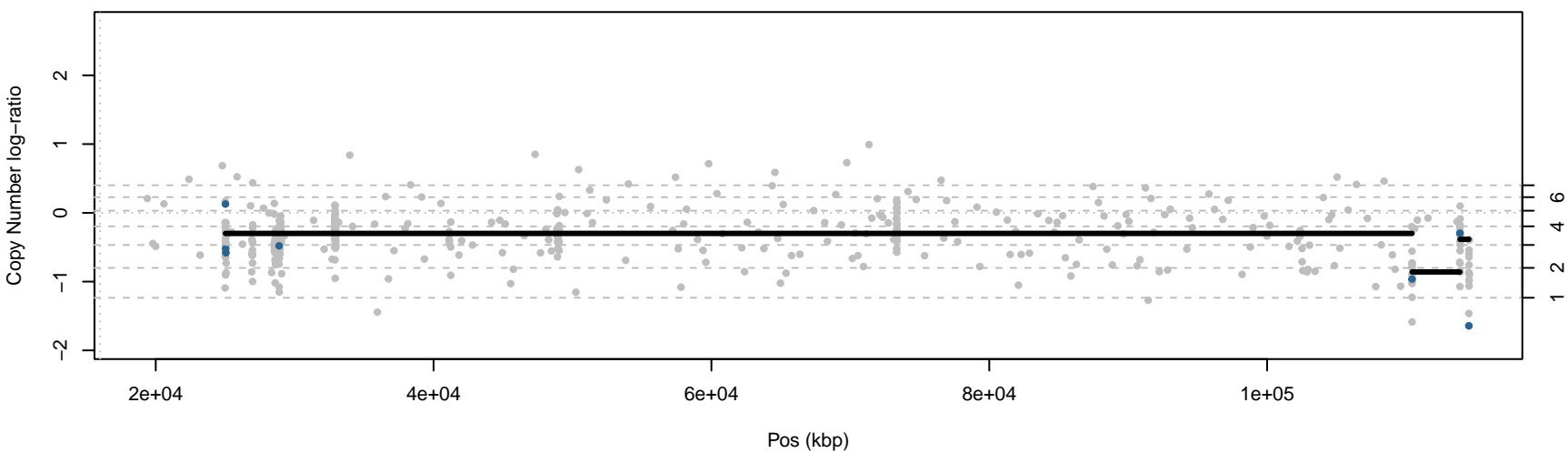
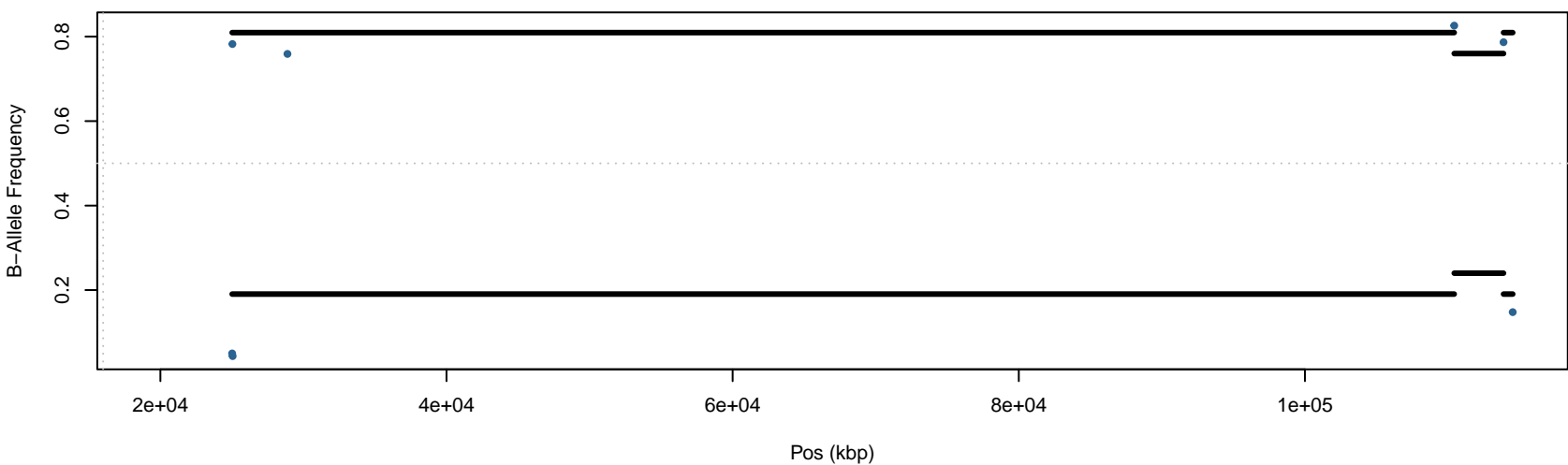
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 11

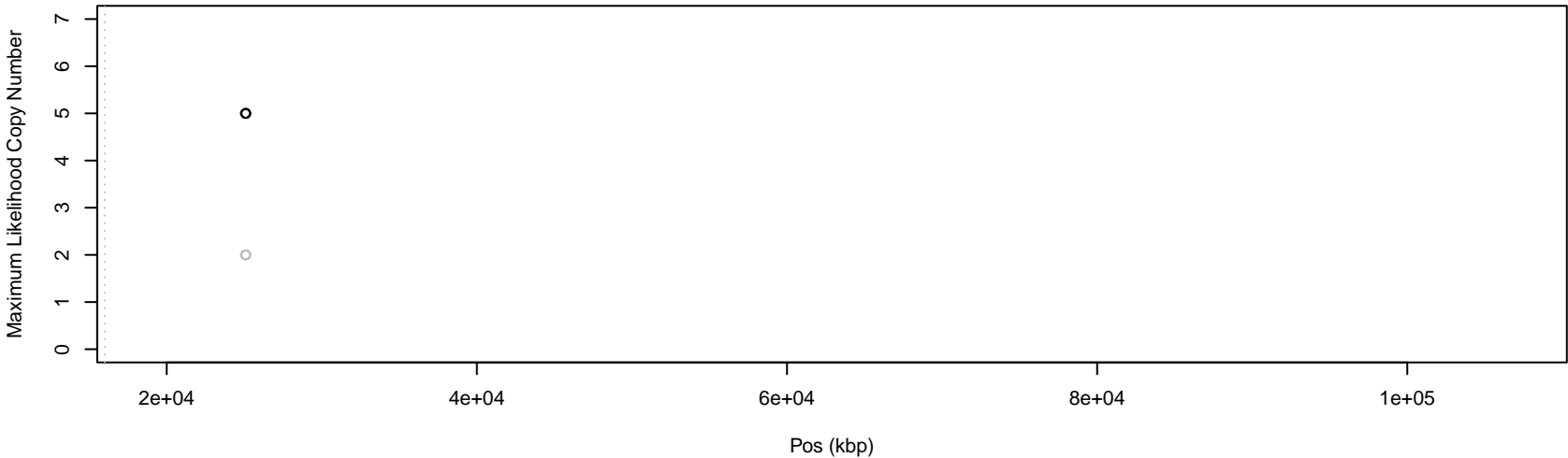
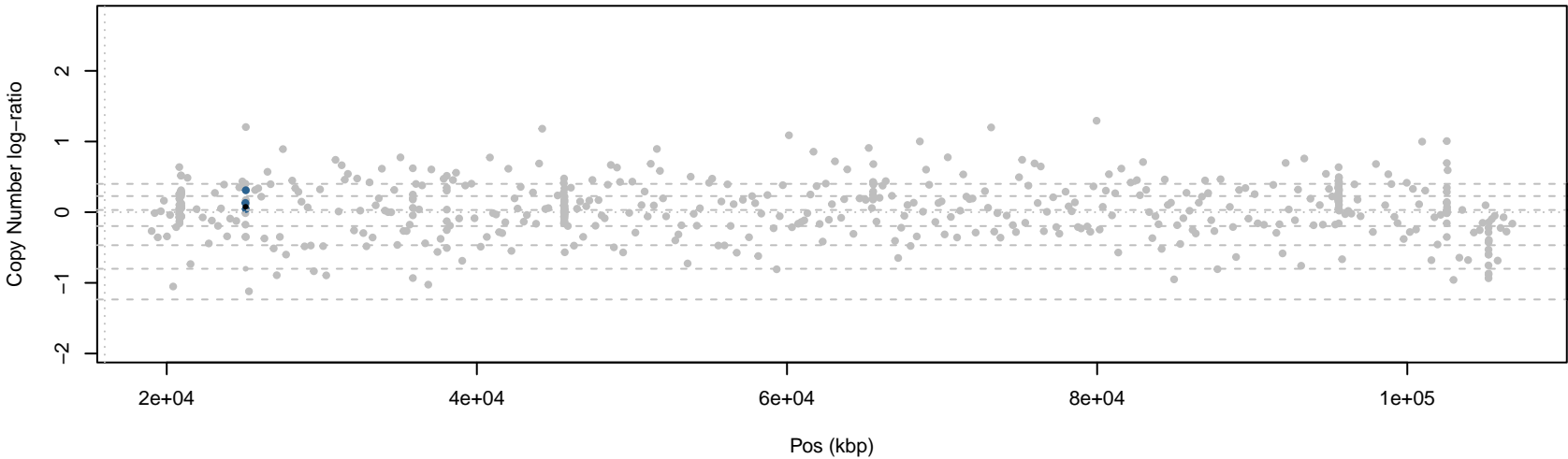
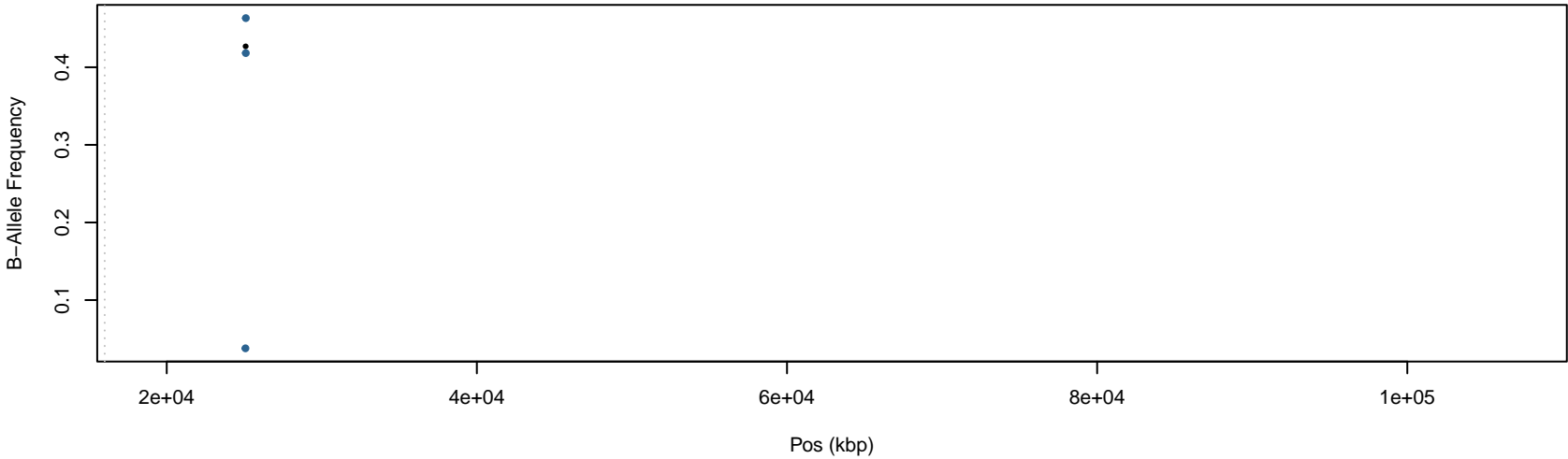


Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 12

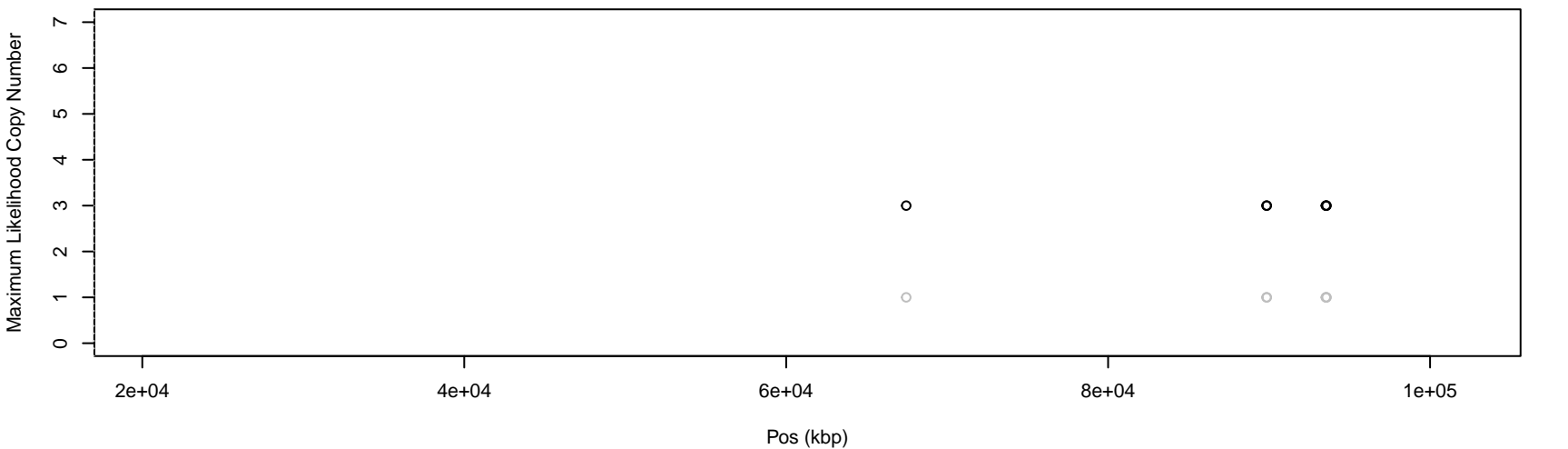
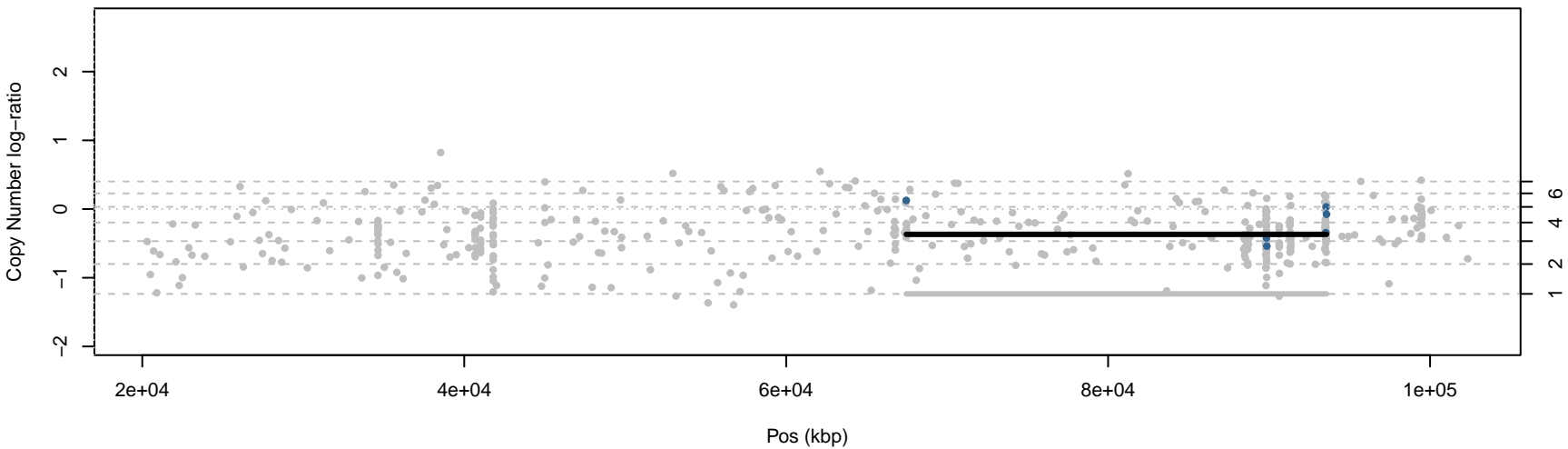
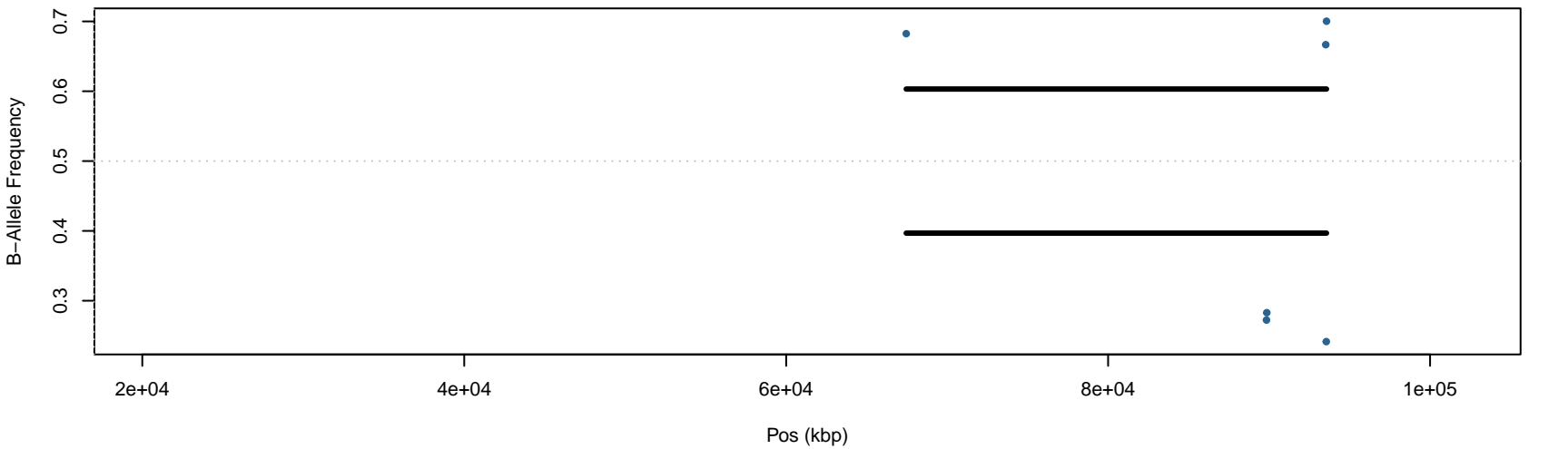


Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 13

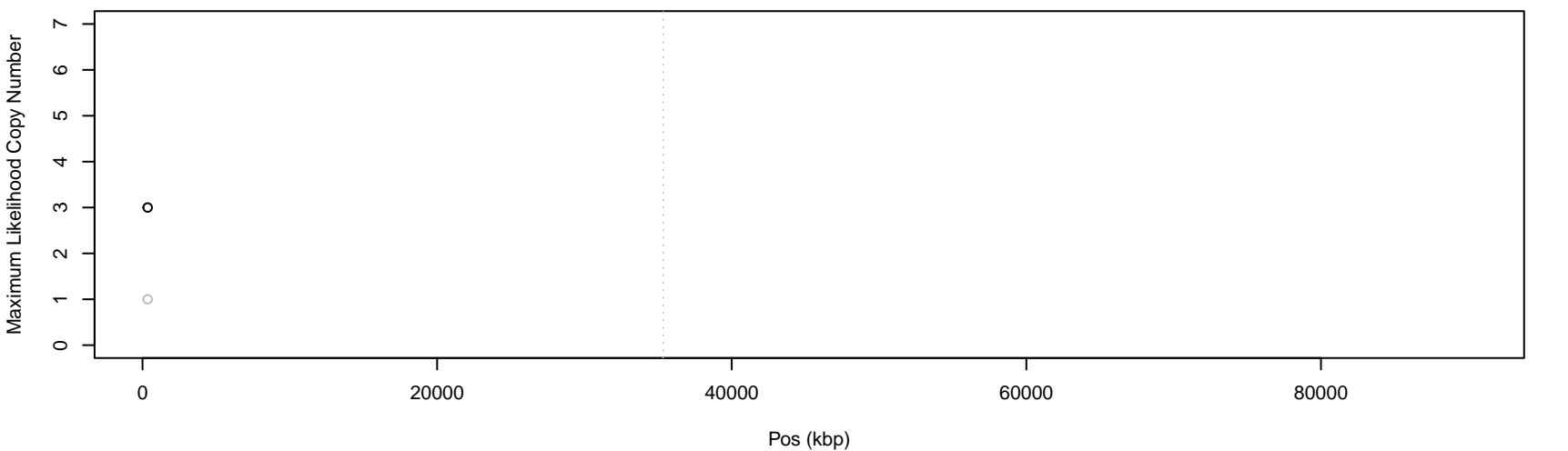
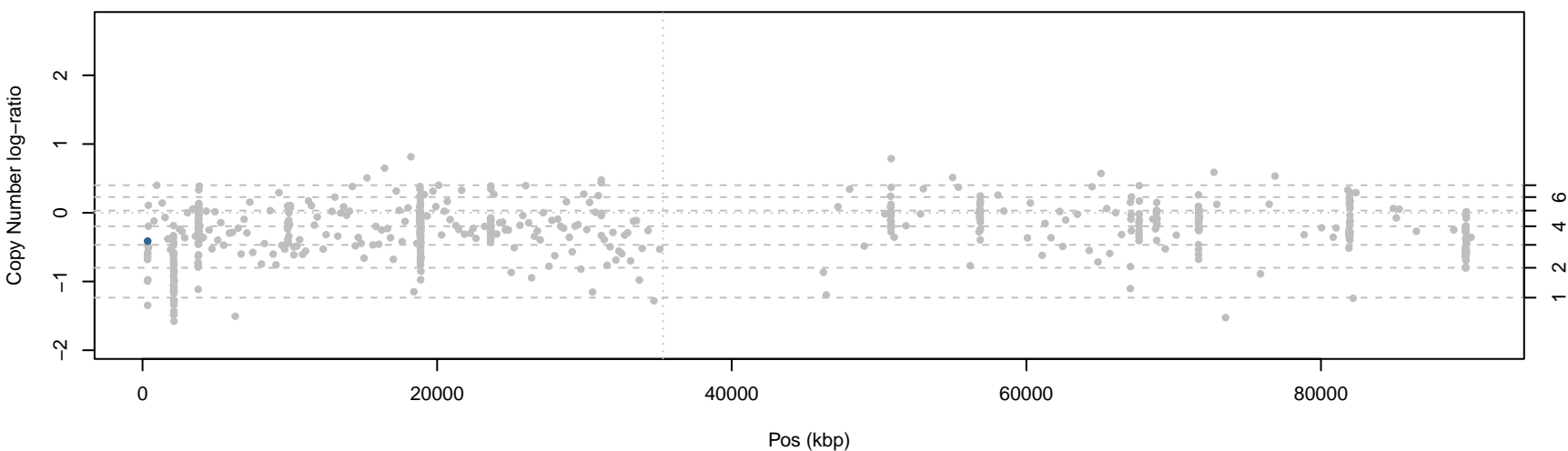
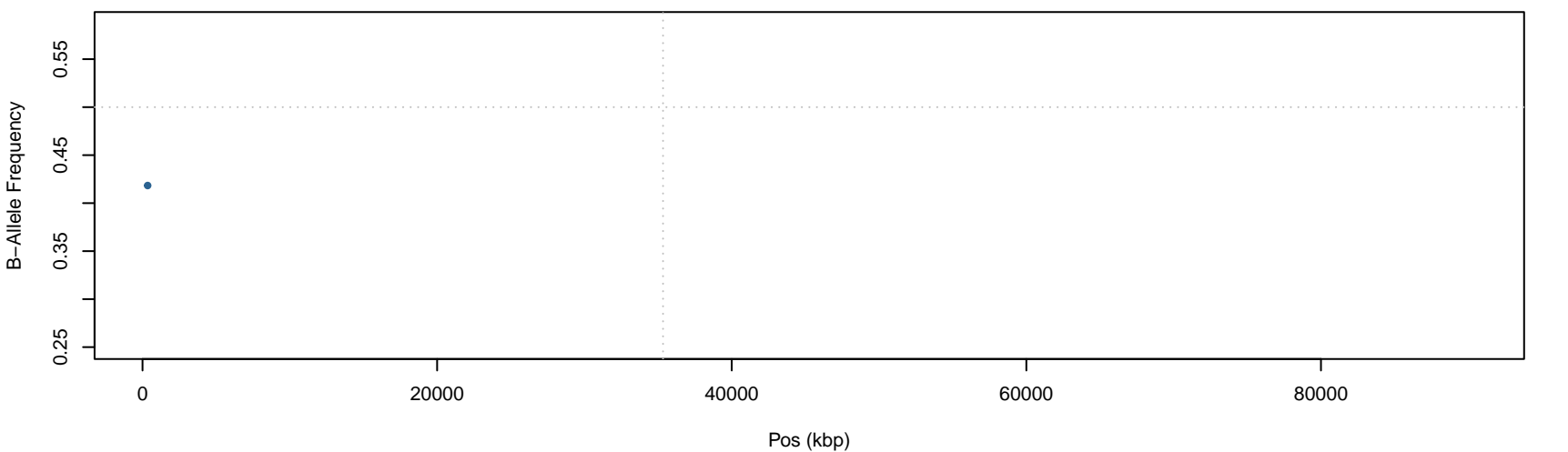




Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 15

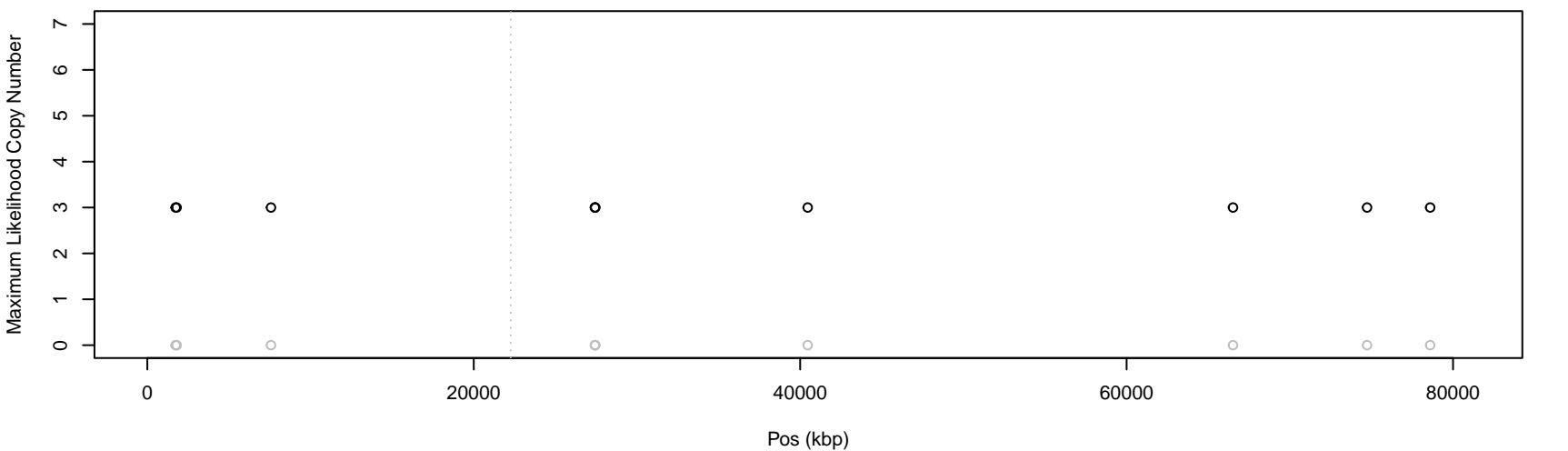
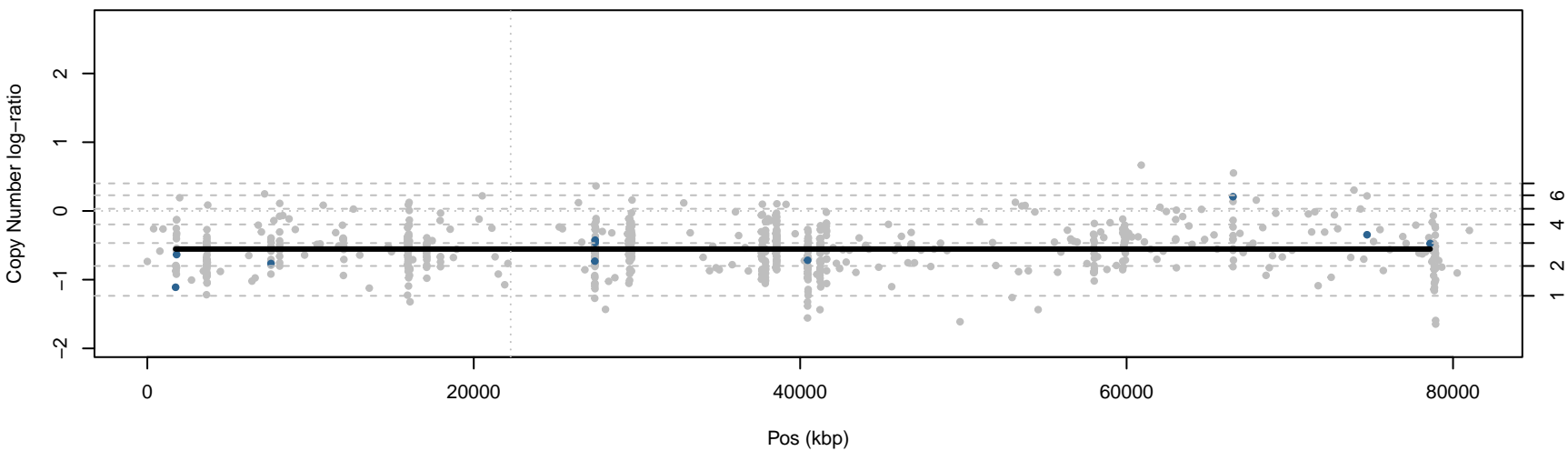
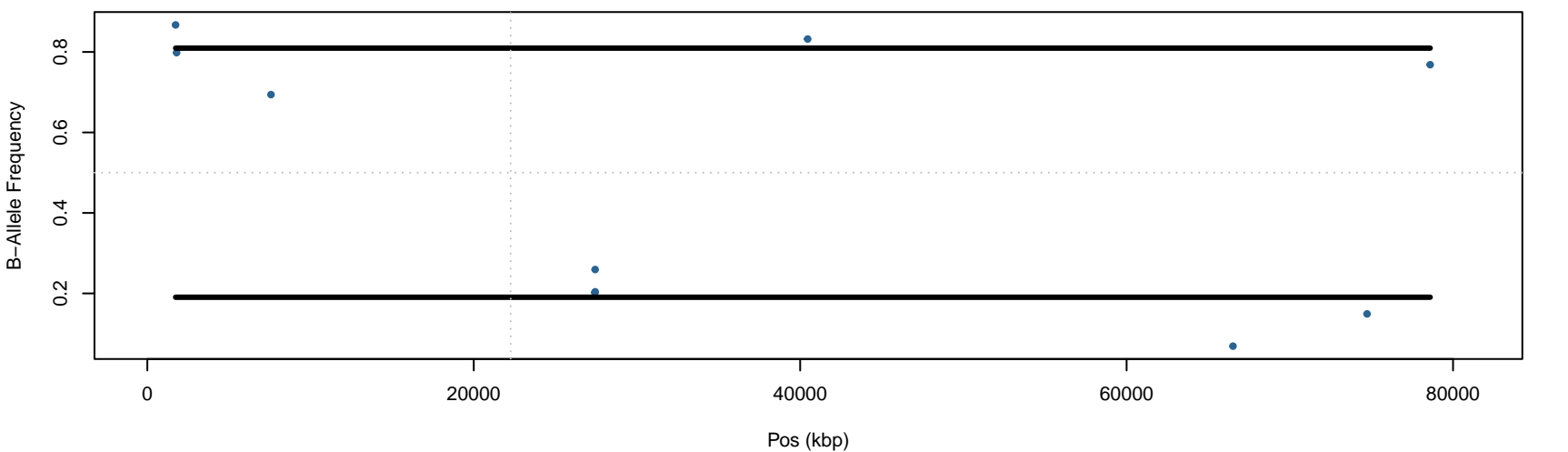


Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 16

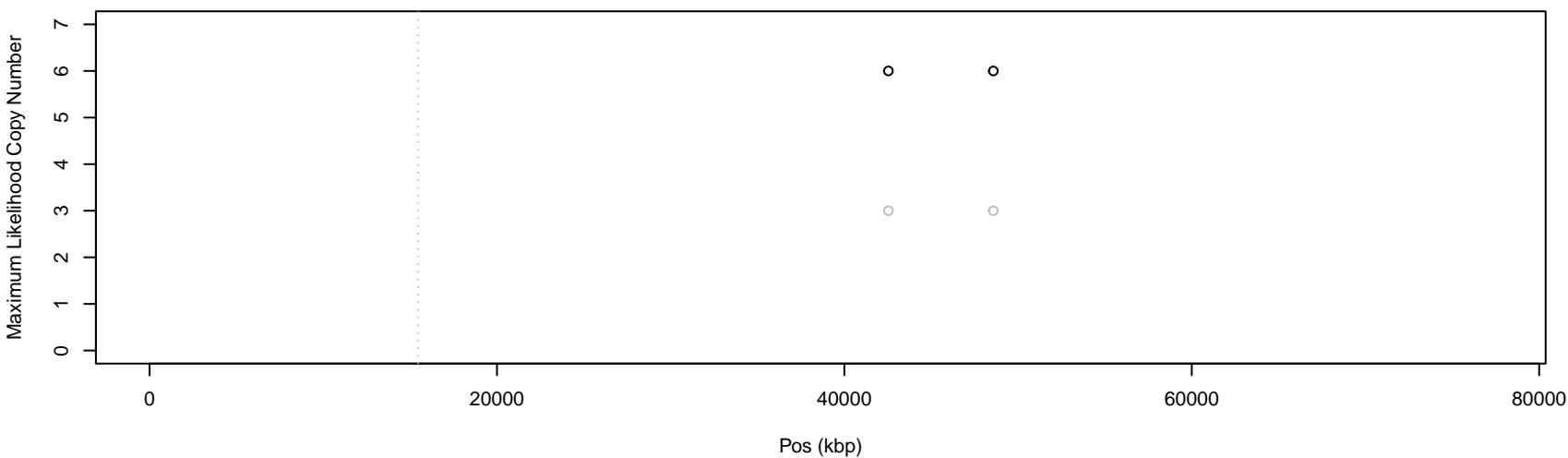
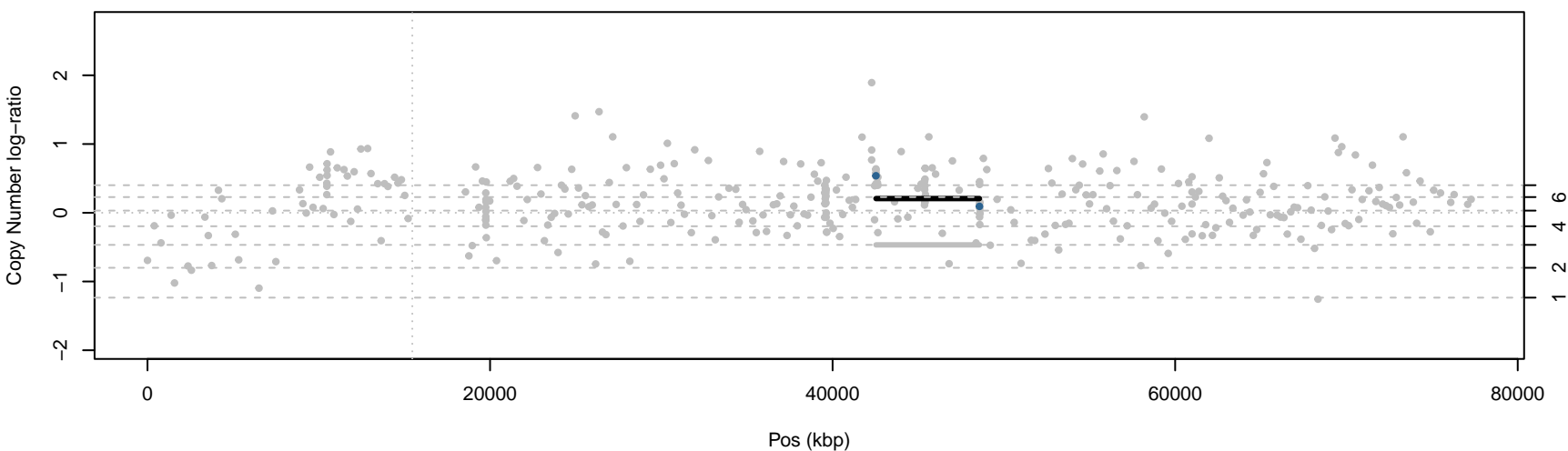
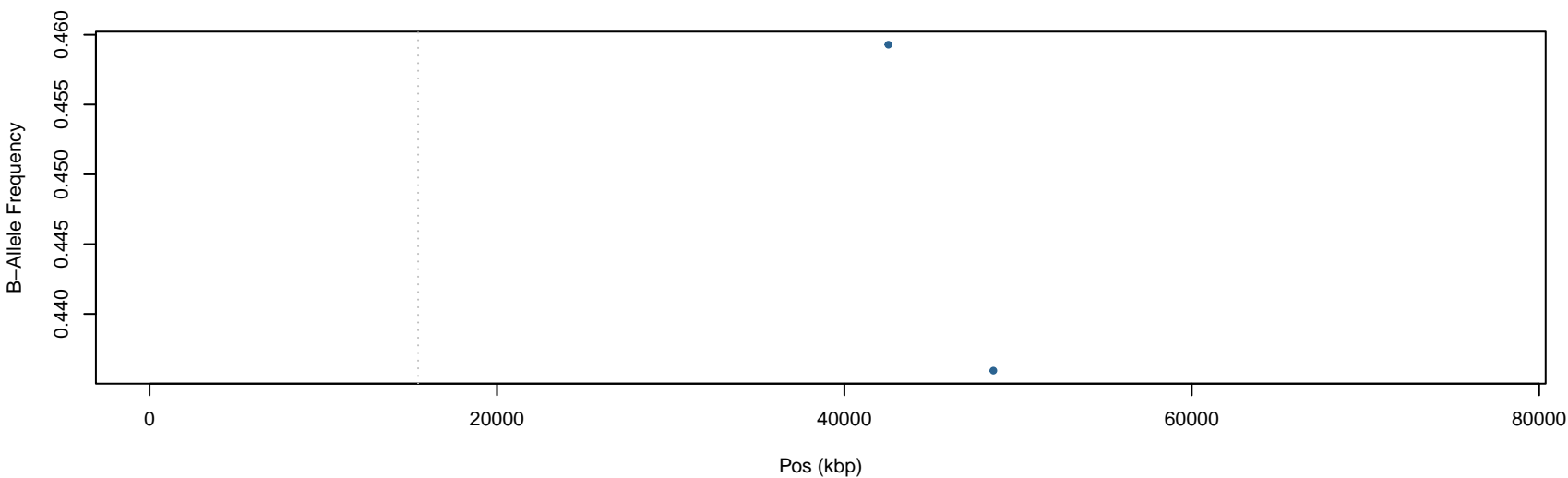




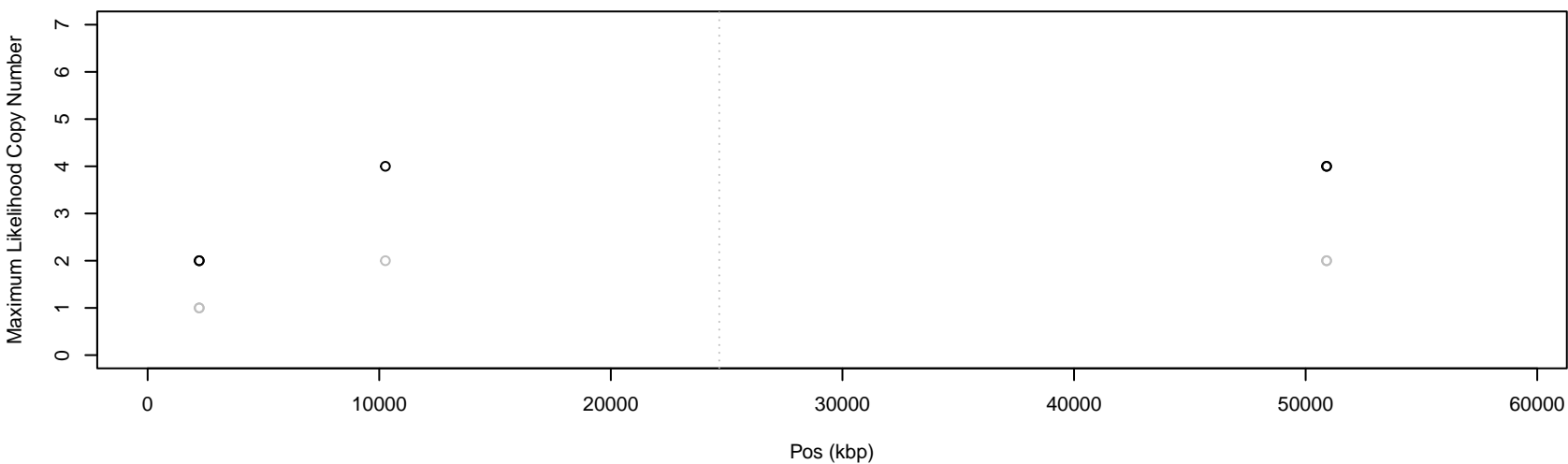
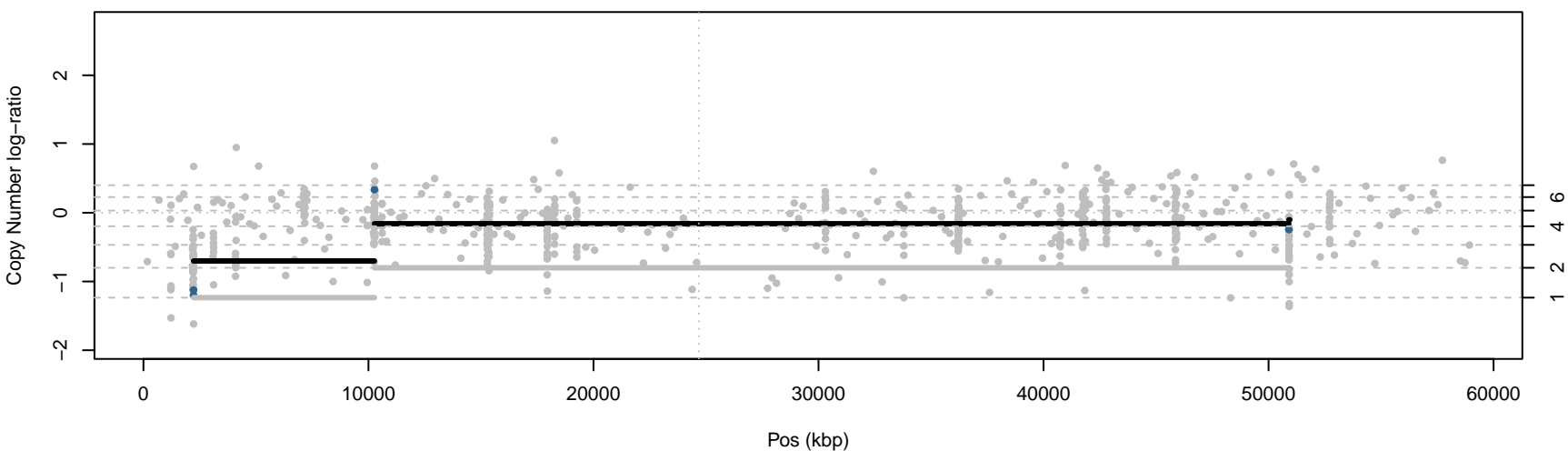
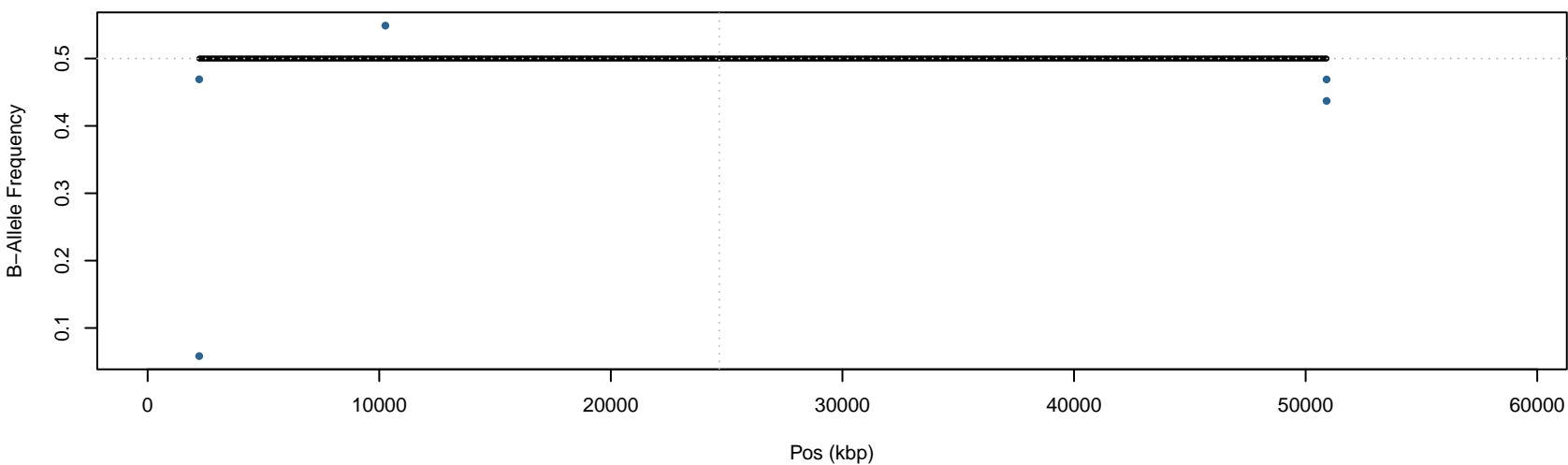
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 17



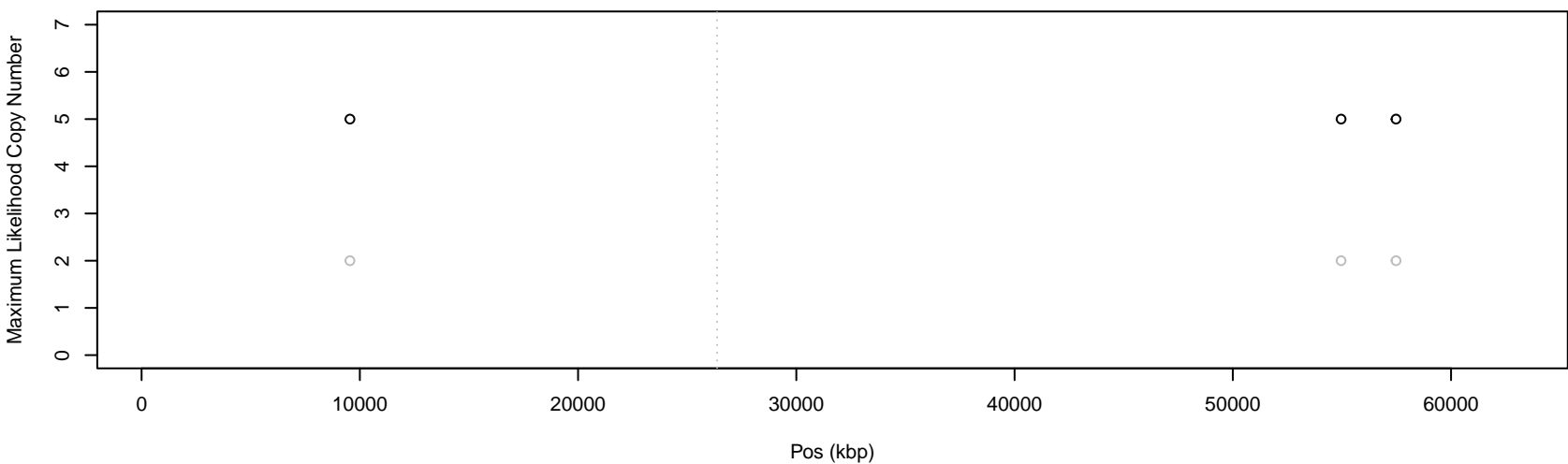
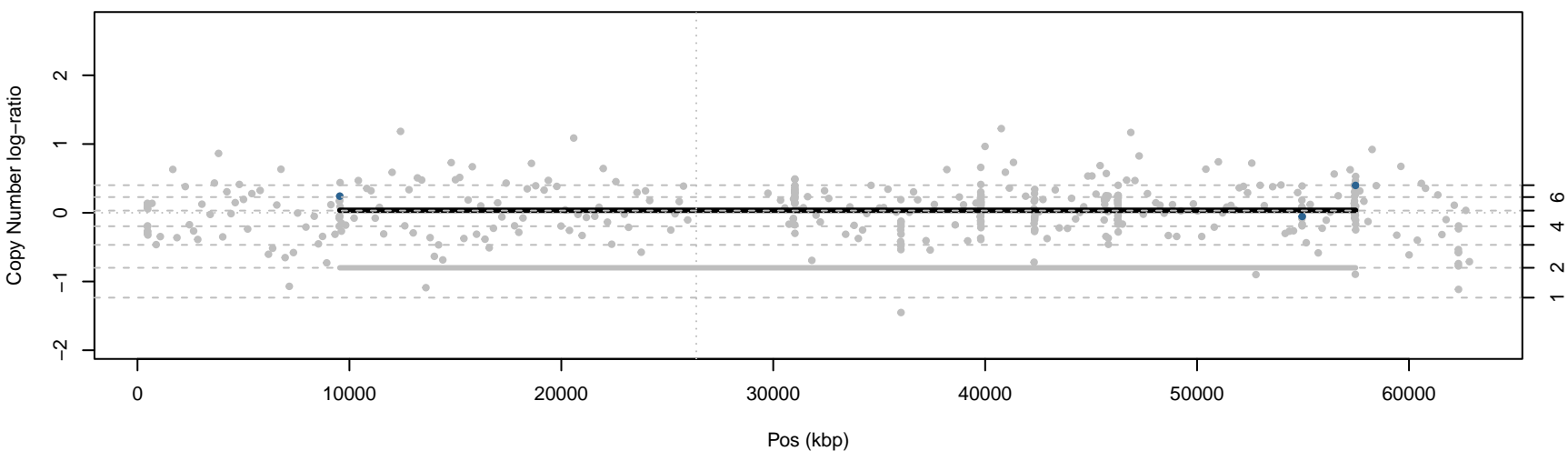
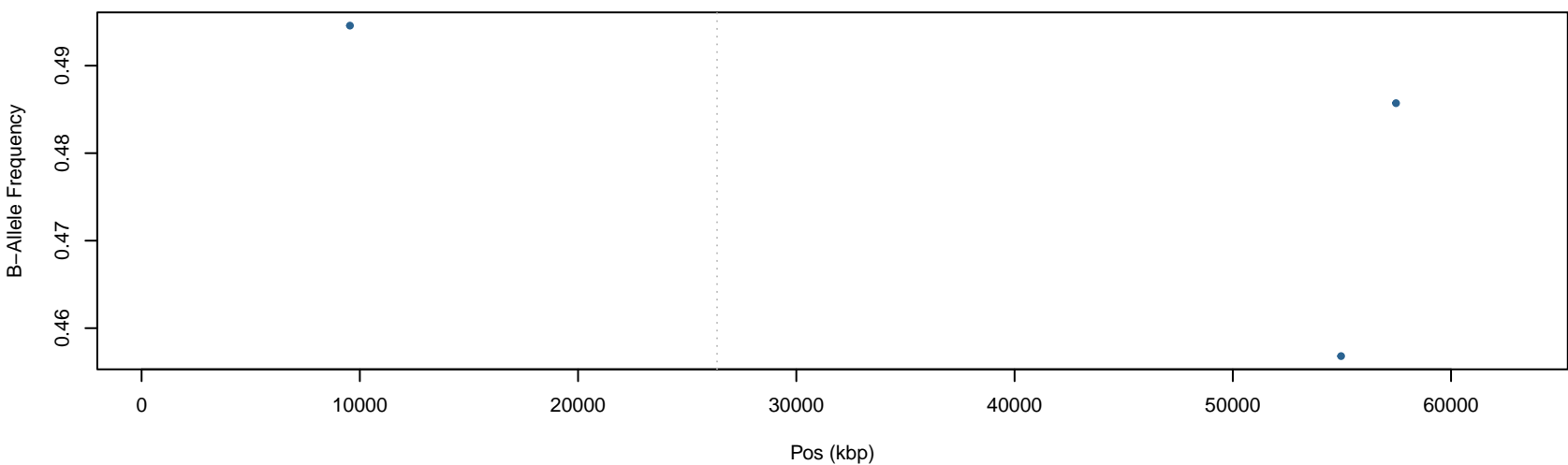
Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 18

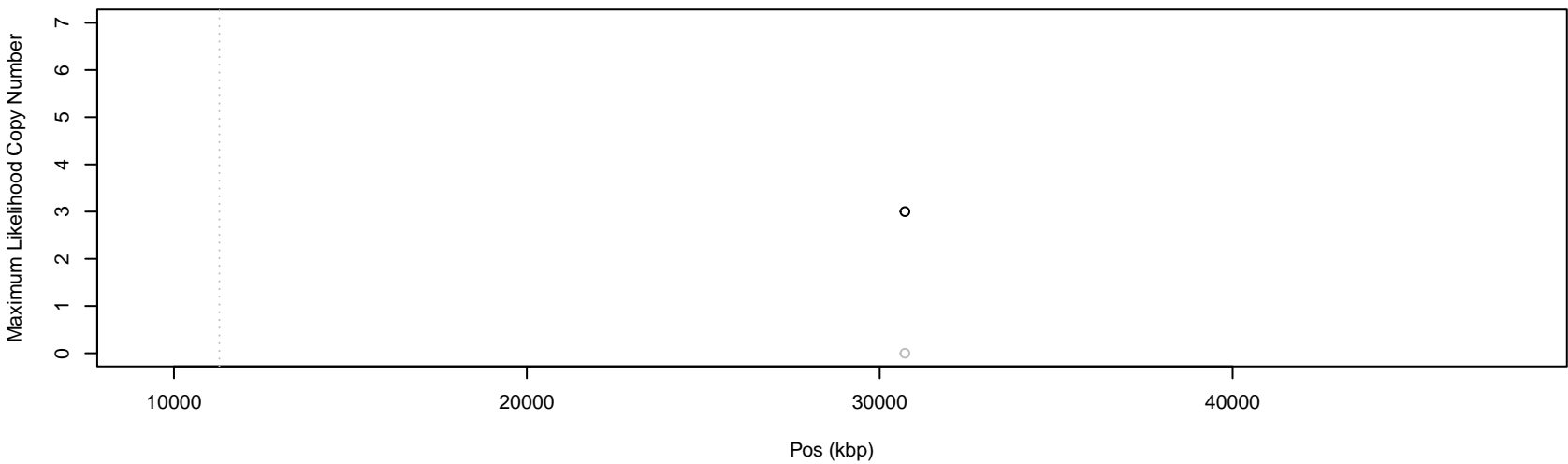
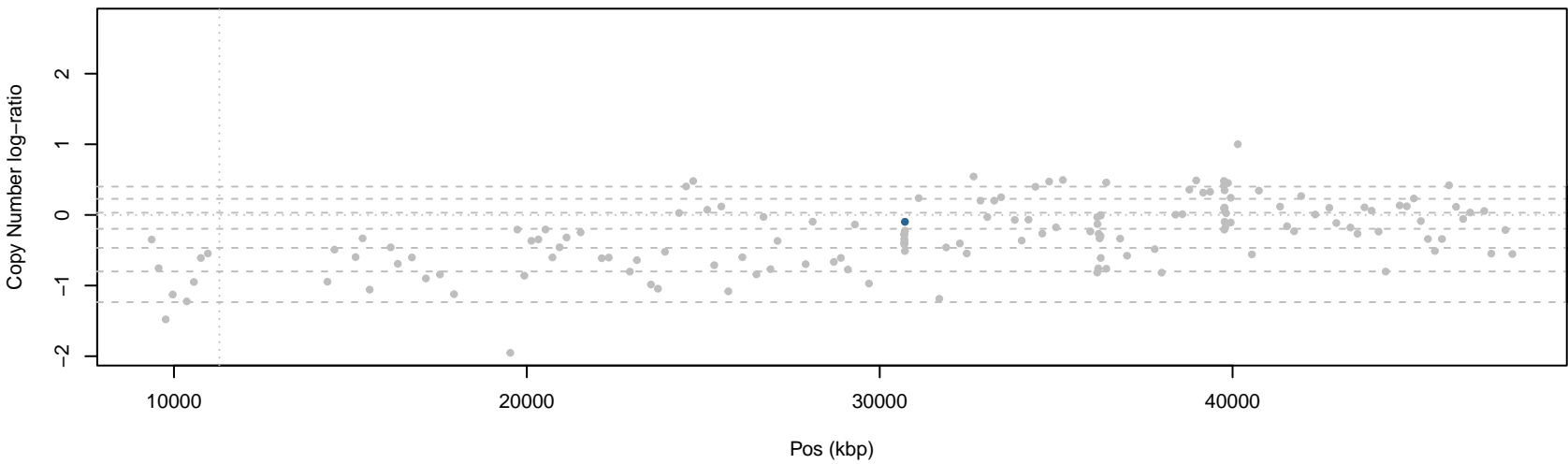
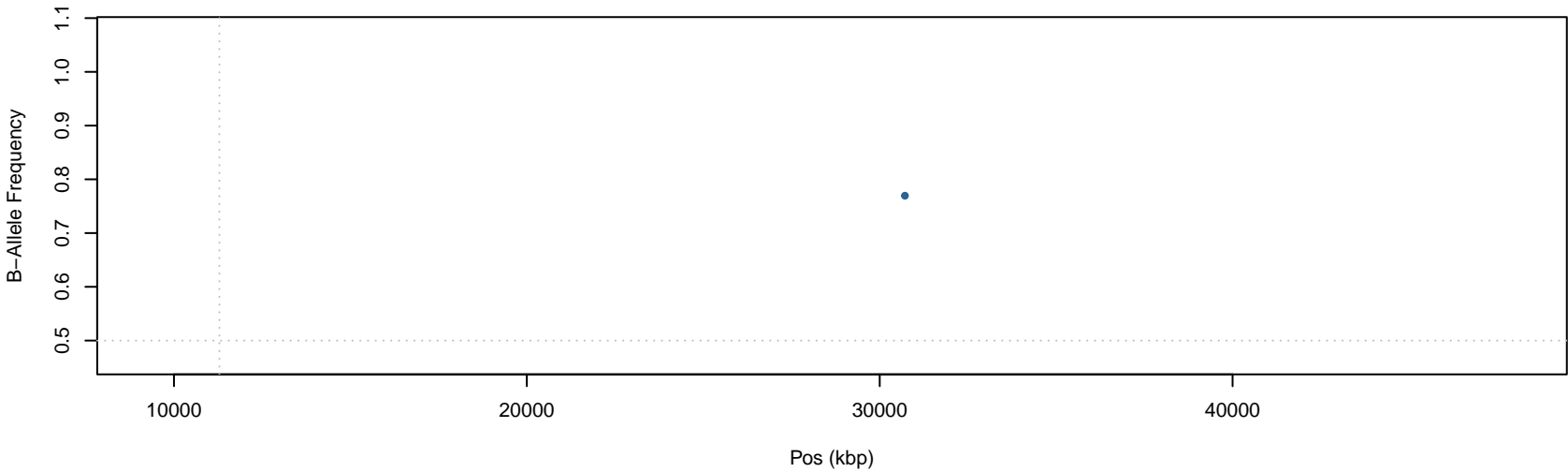


Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 19



Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 20





Purity: 0.52 Tumor ploidy: 4.853 SNV log-likelihood: -191.01 GoF: 85.9% Mean coverage: 305;536 Chromosome: 22

