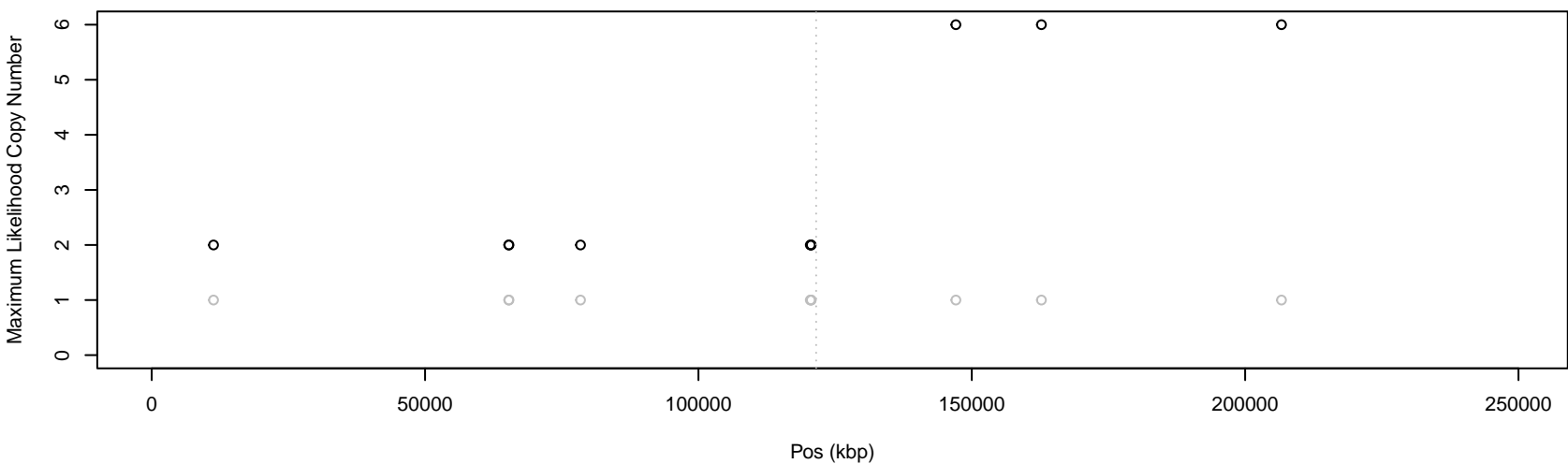
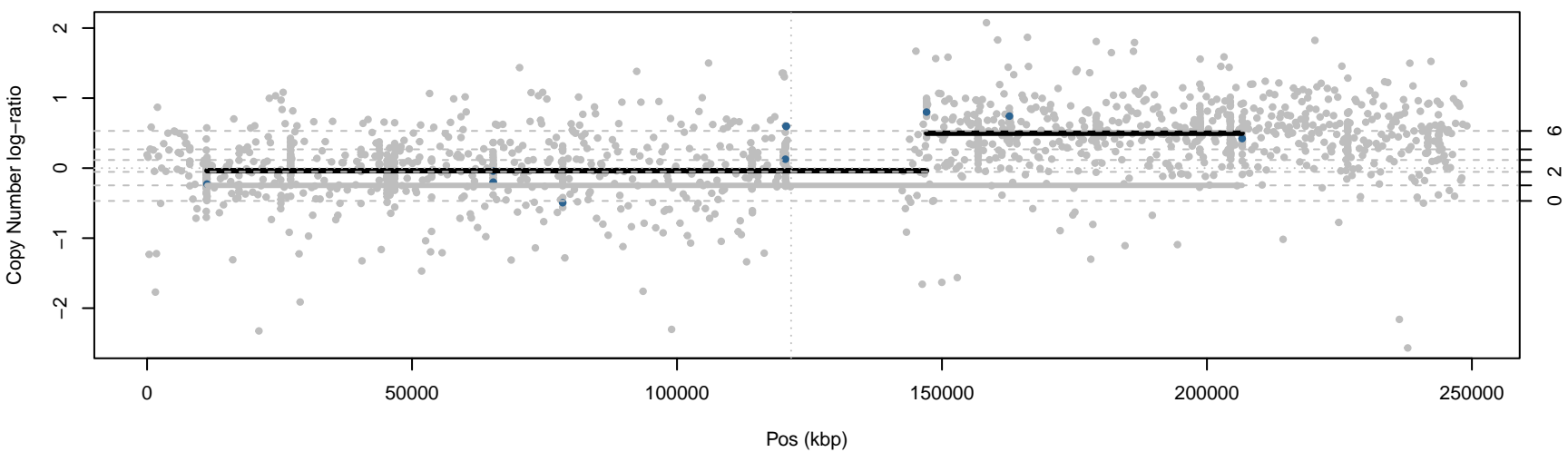
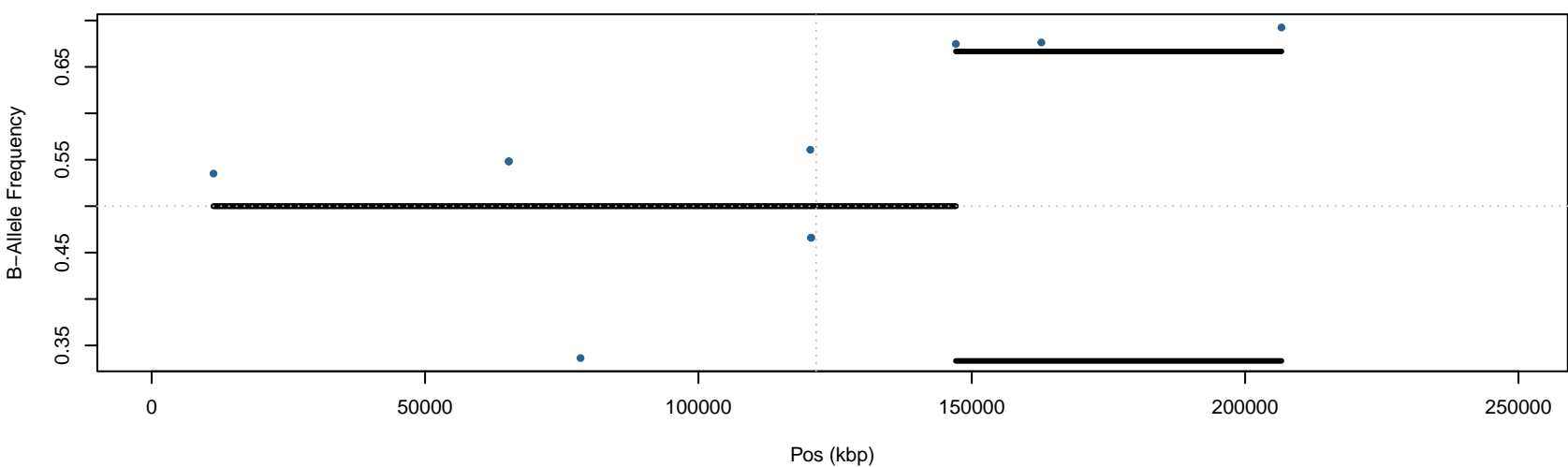
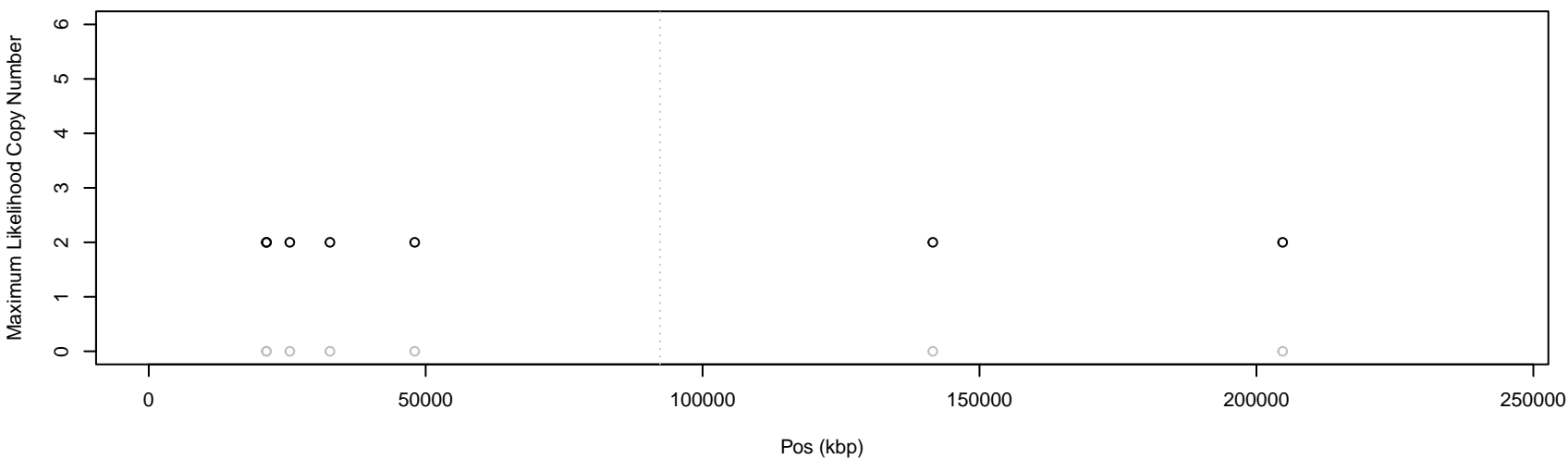
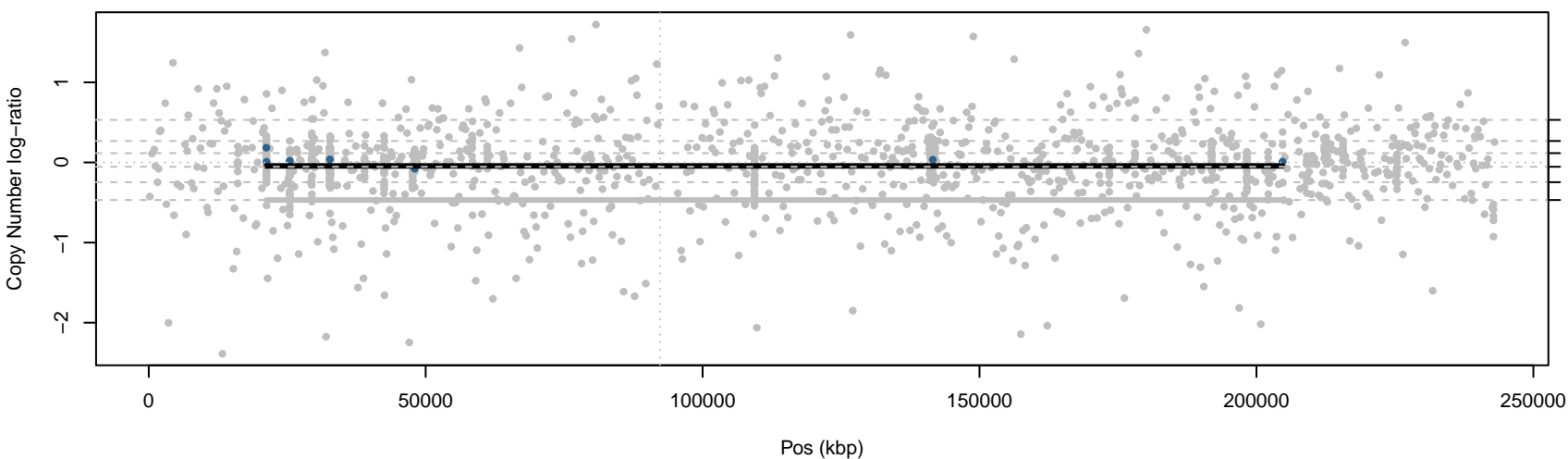
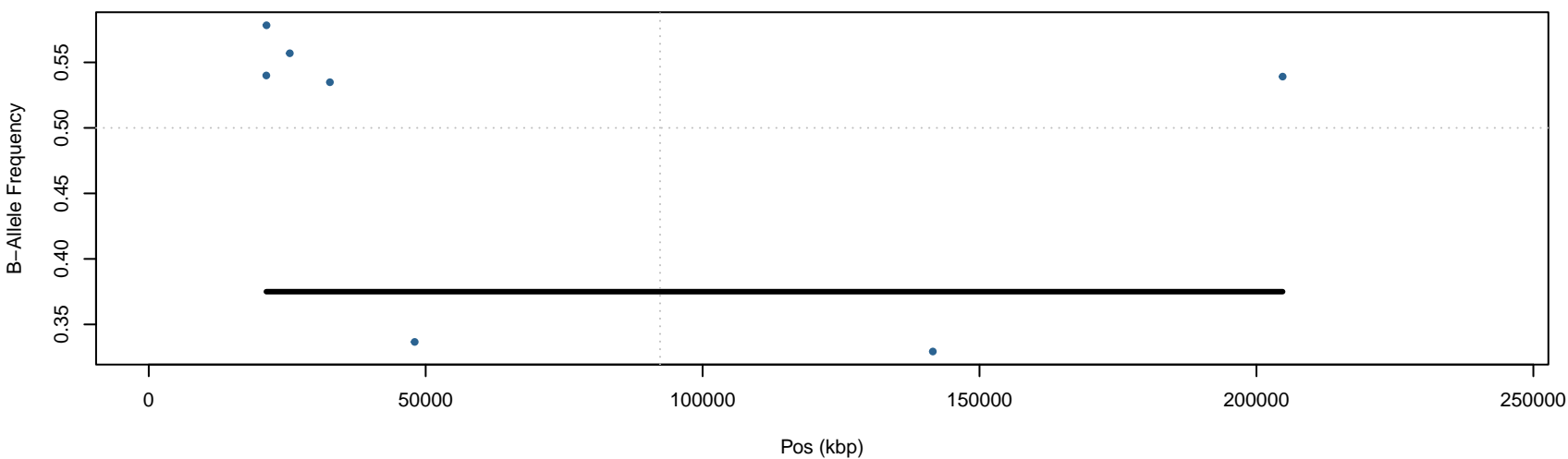


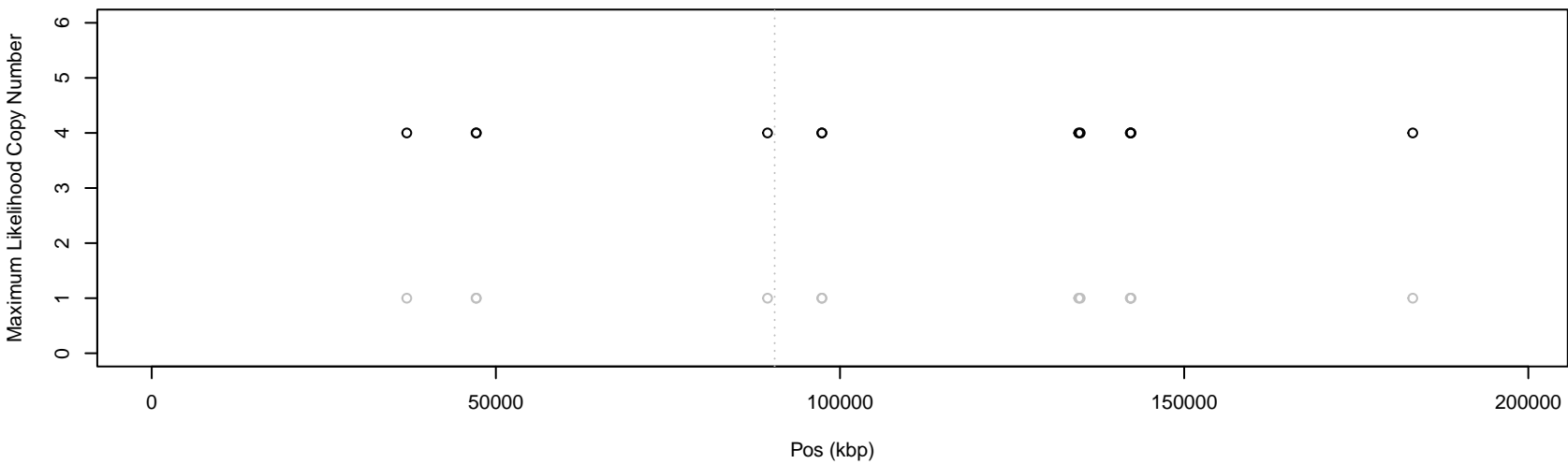
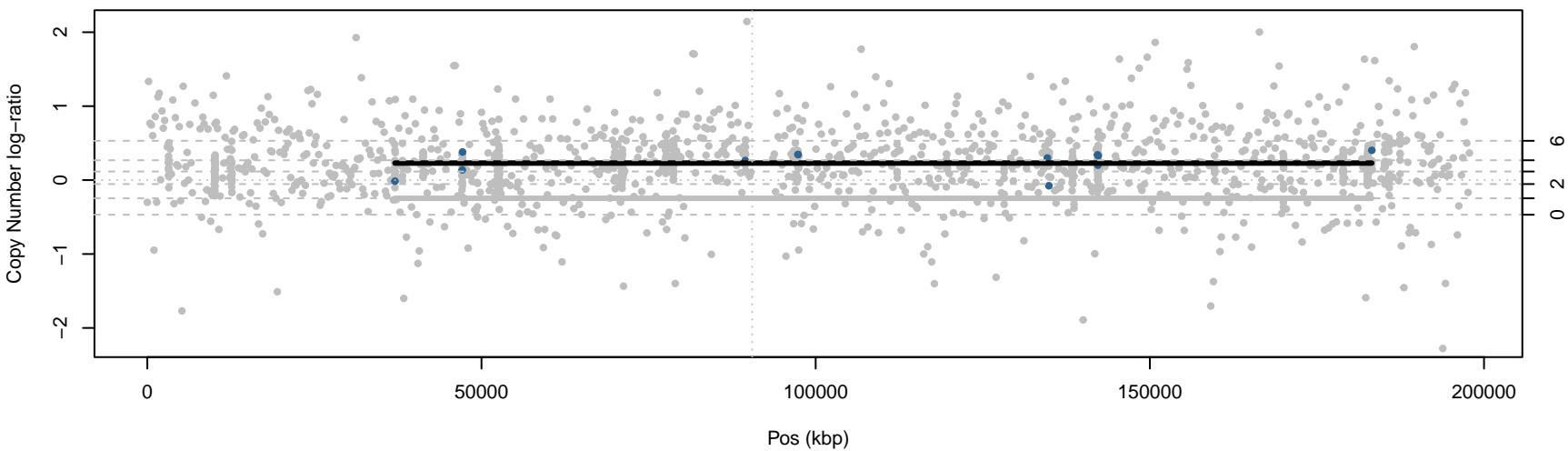
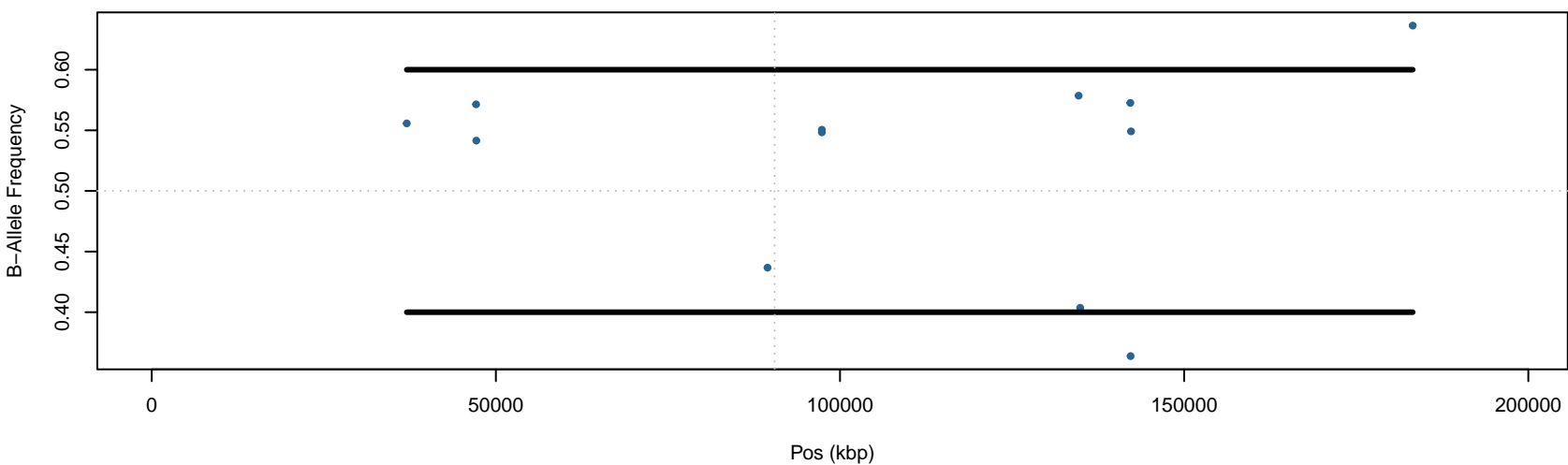
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 1



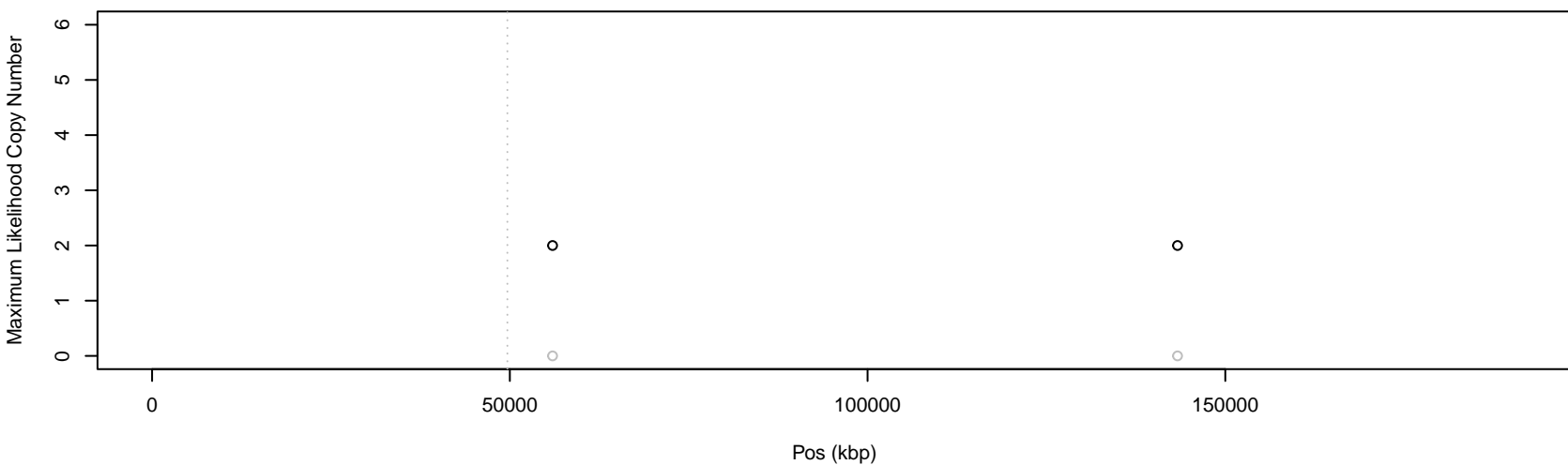
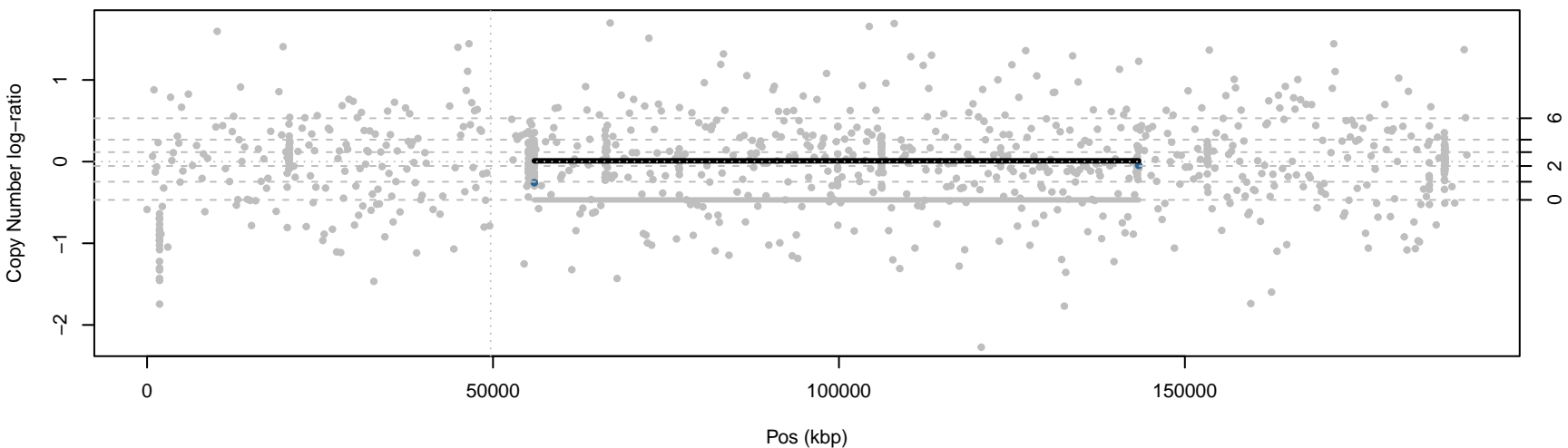
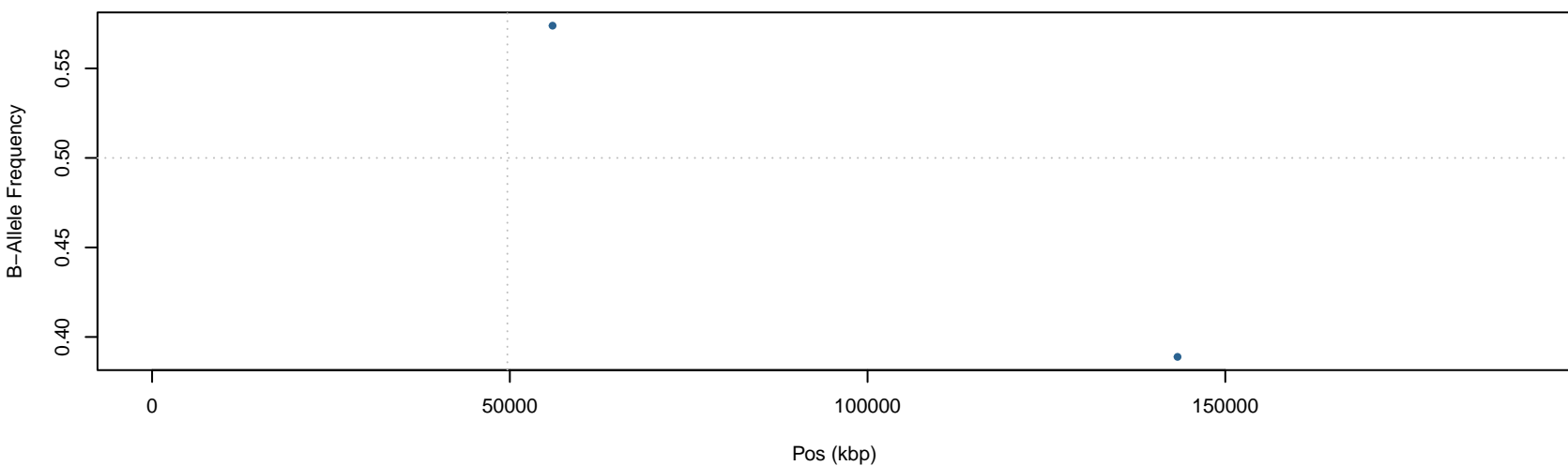
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 2



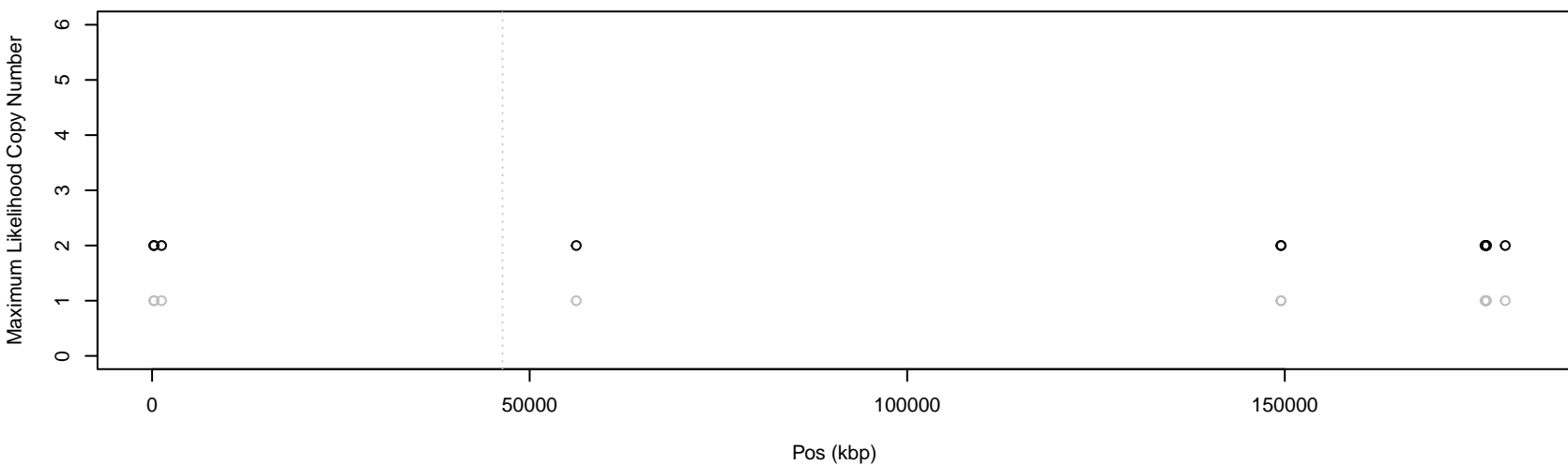
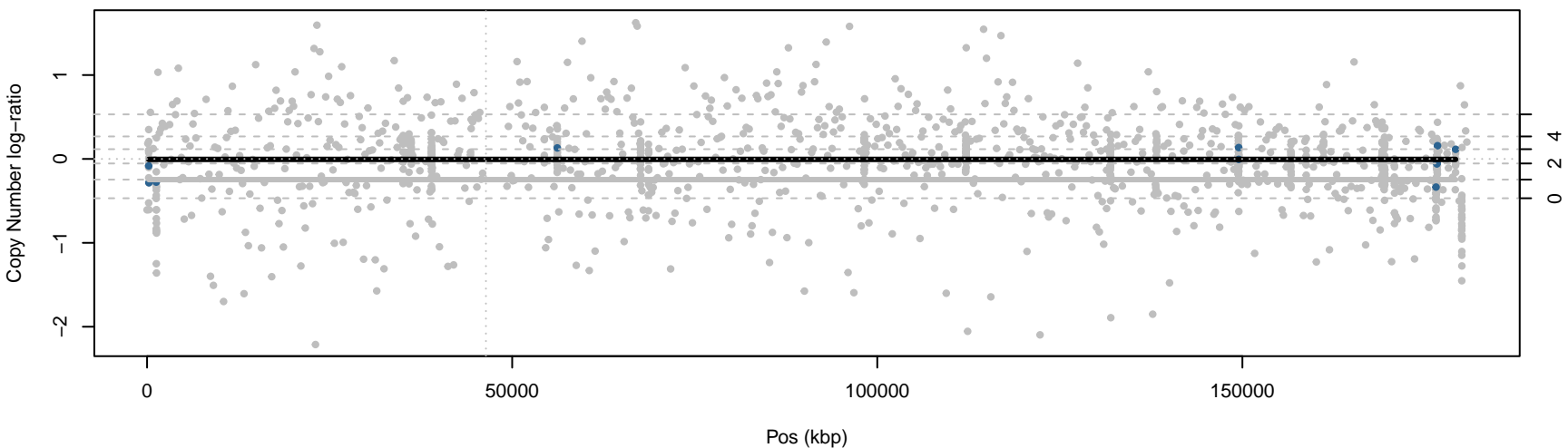
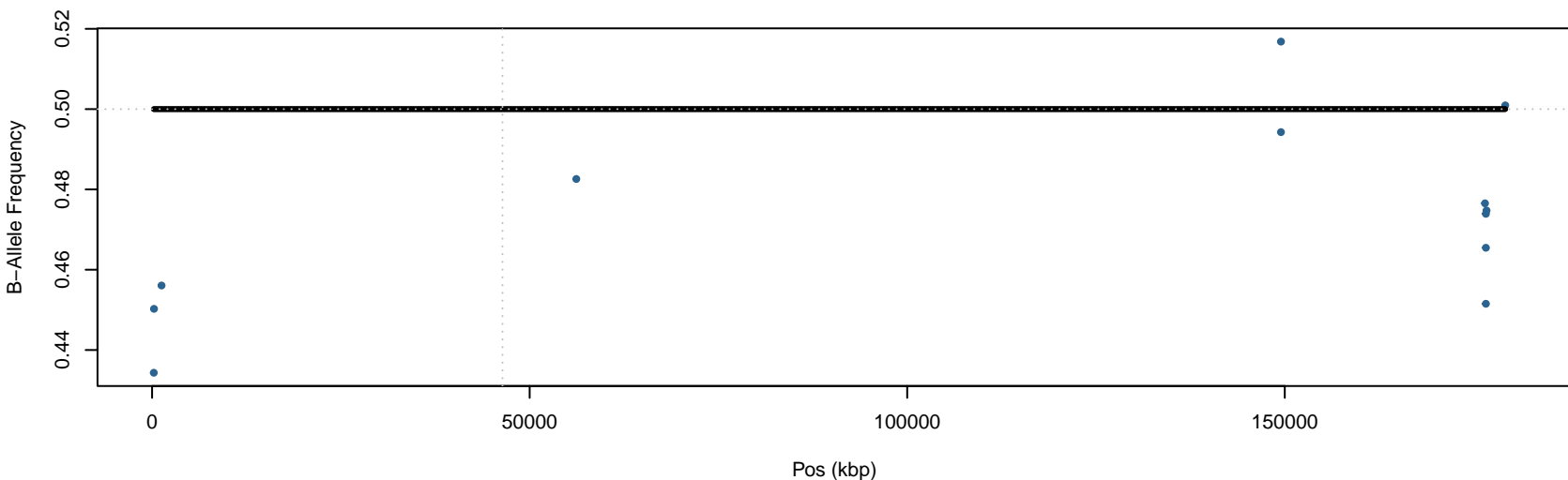
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 3



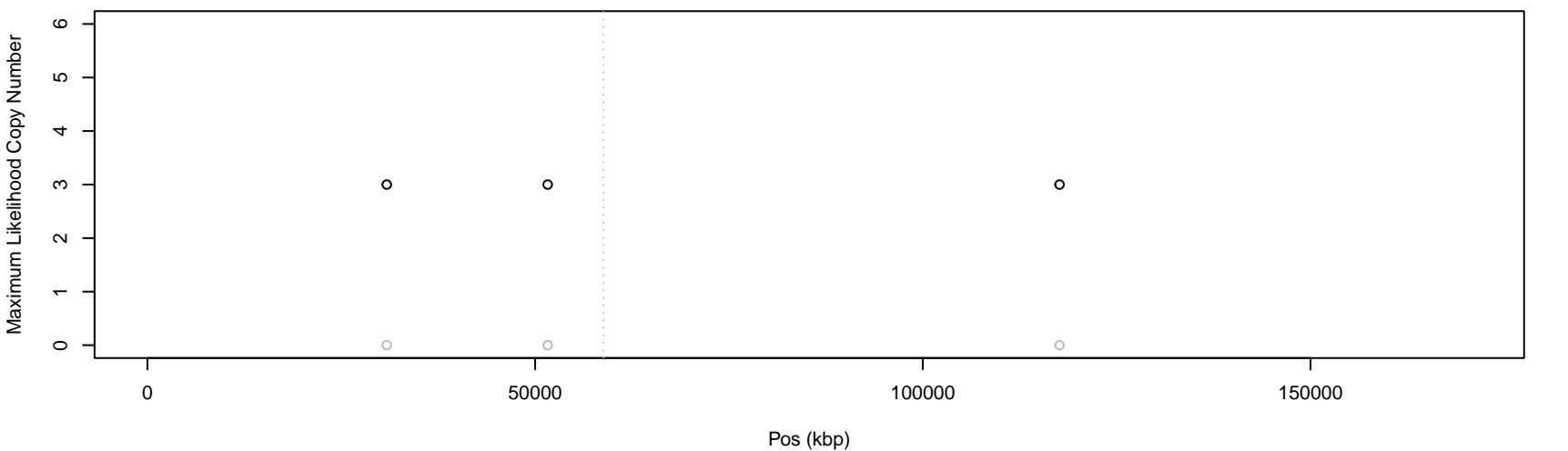
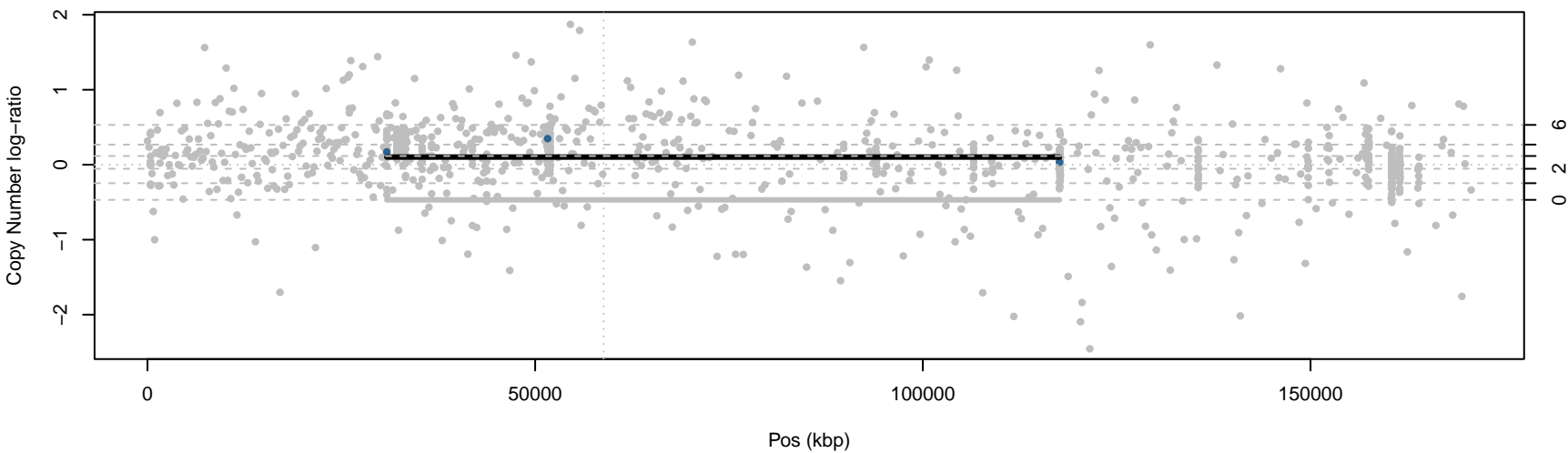
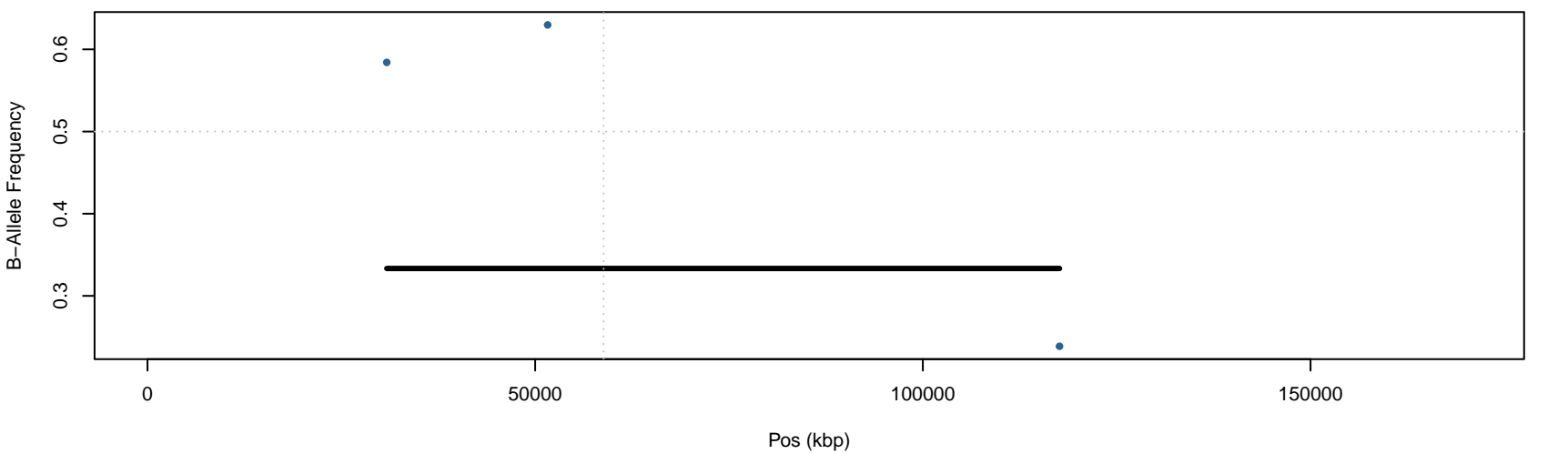
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 4



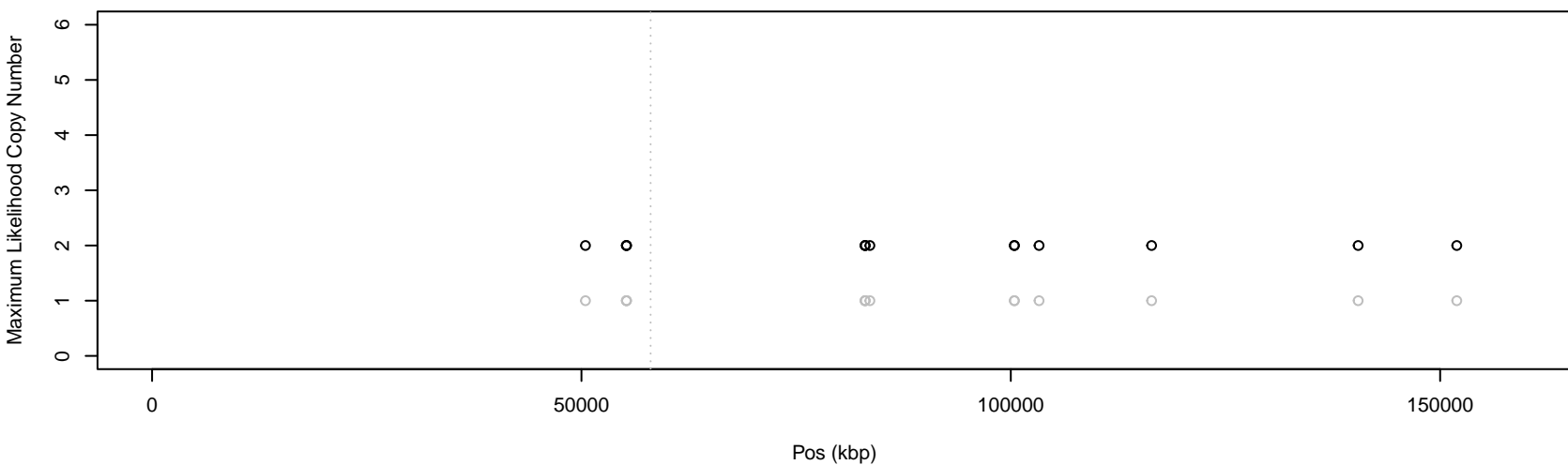
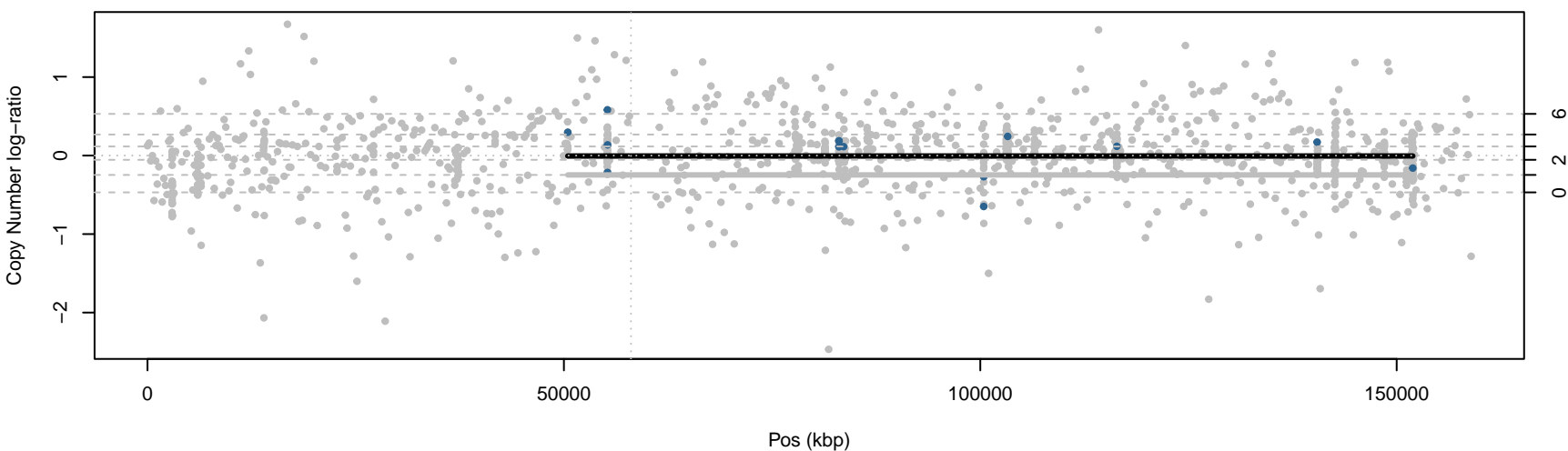
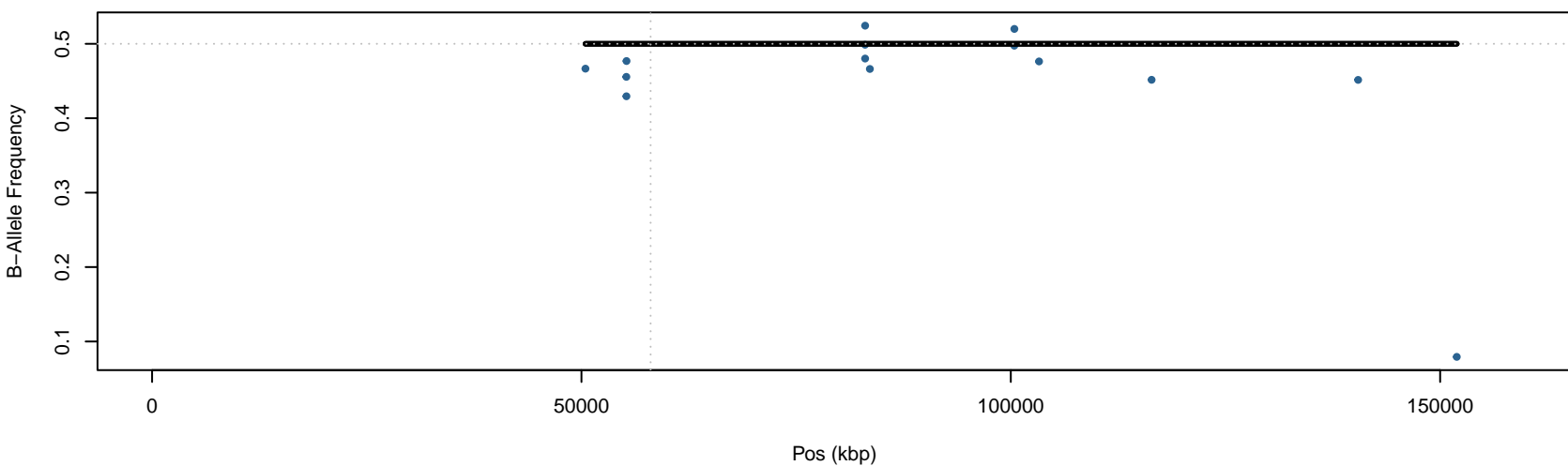
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 5



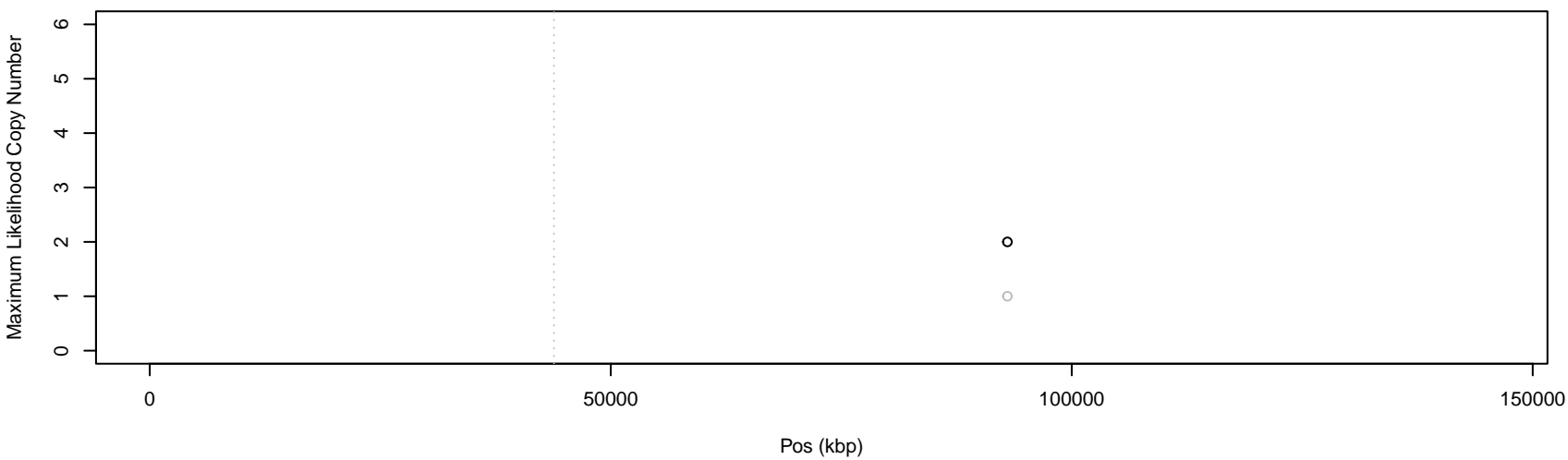
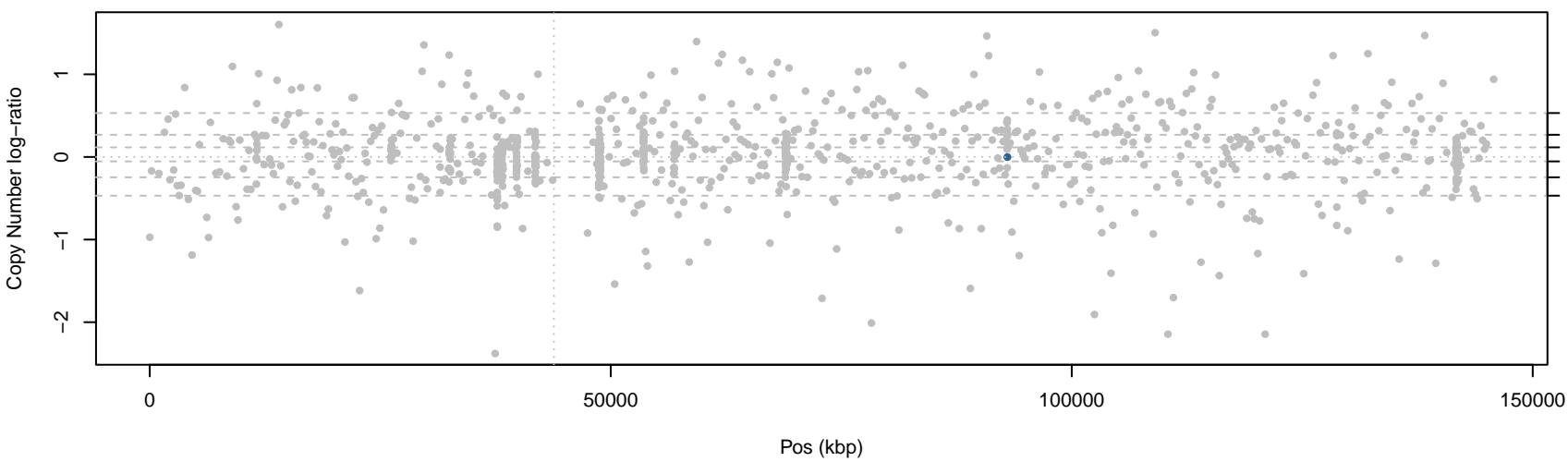
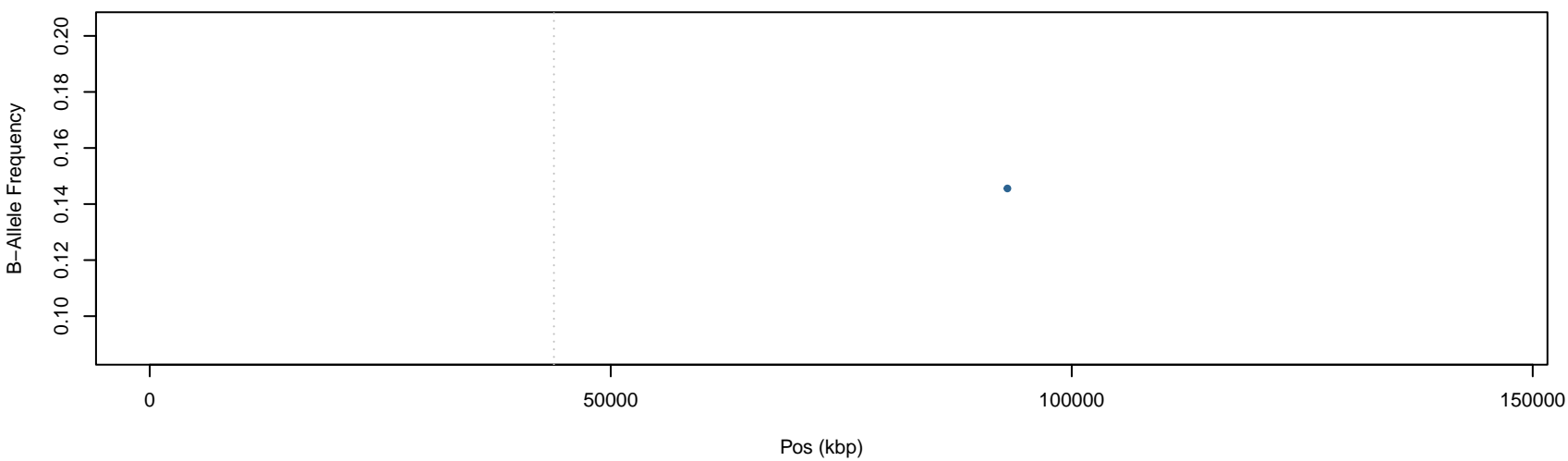
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 6



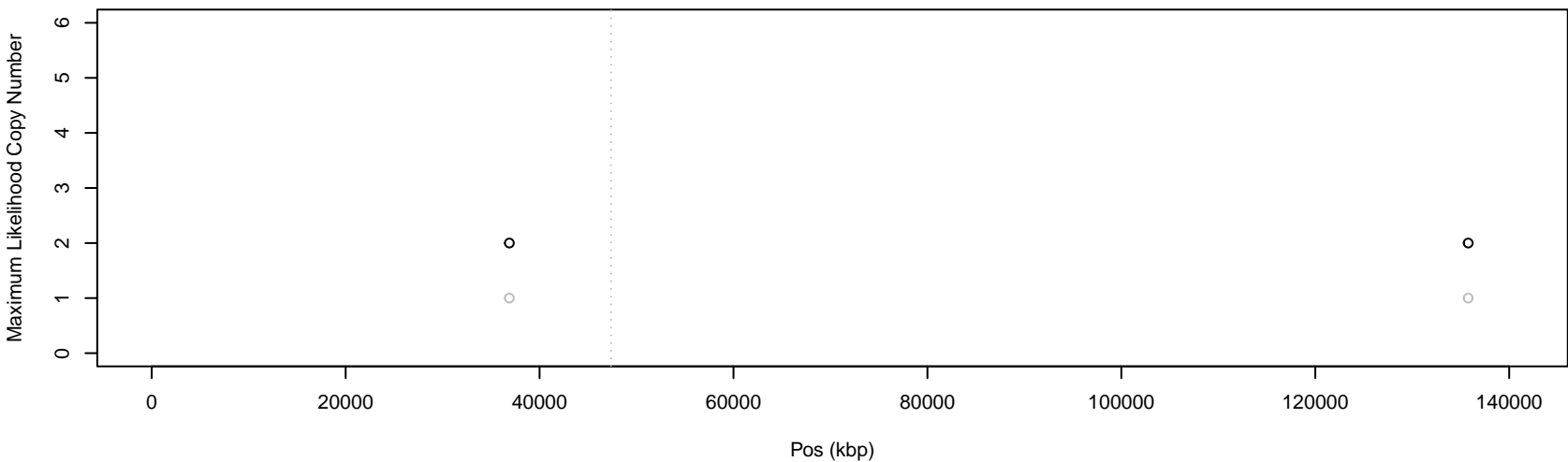
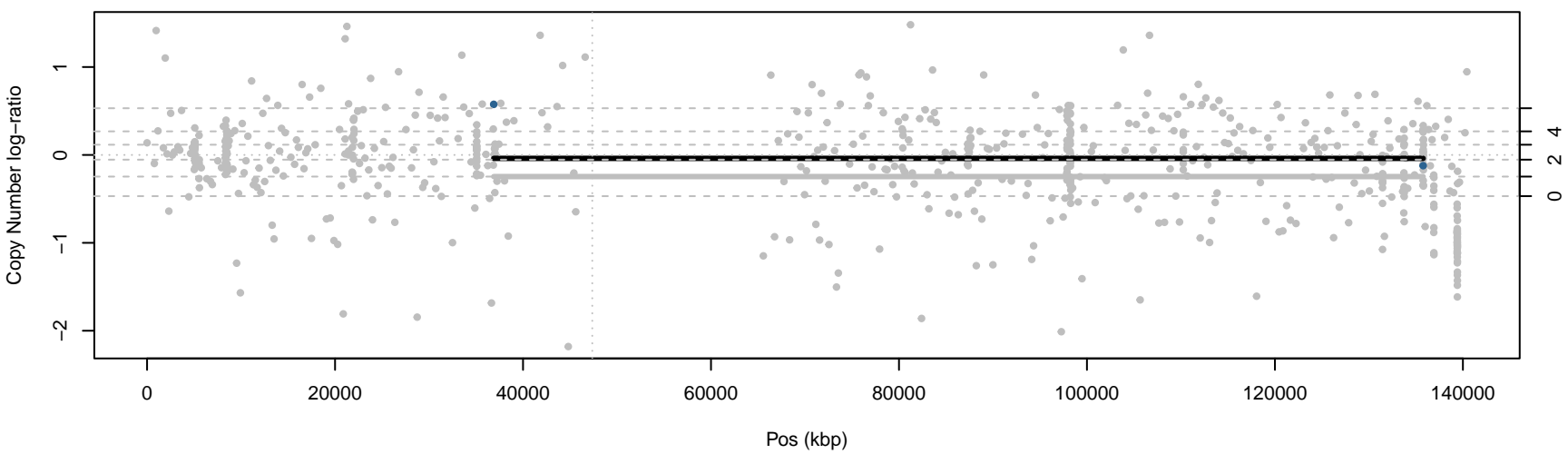
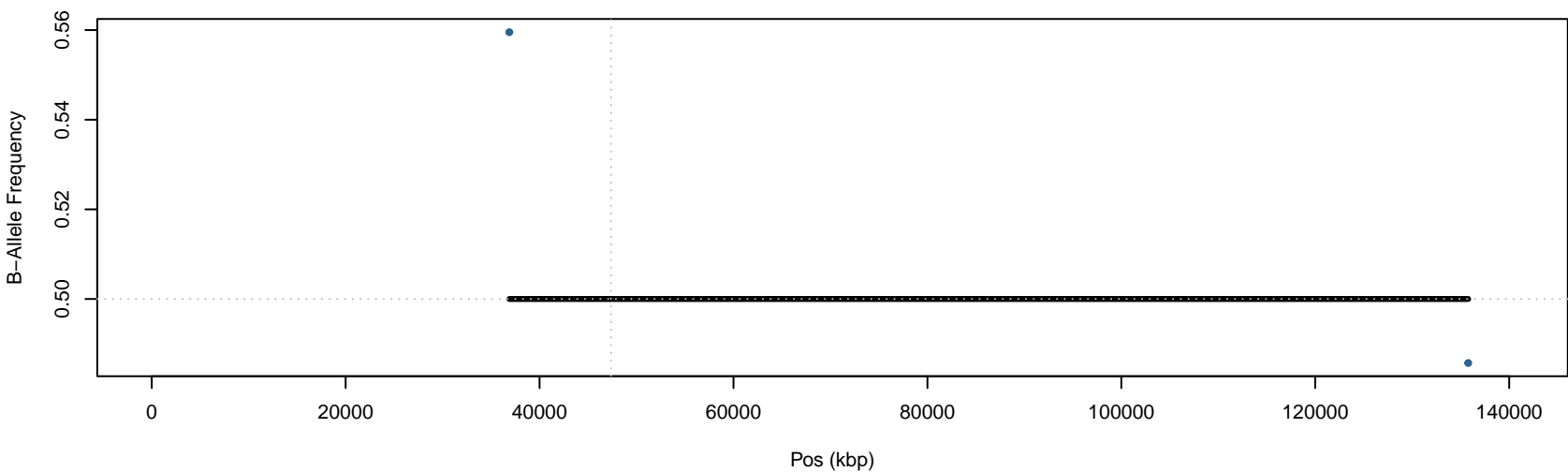
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 7



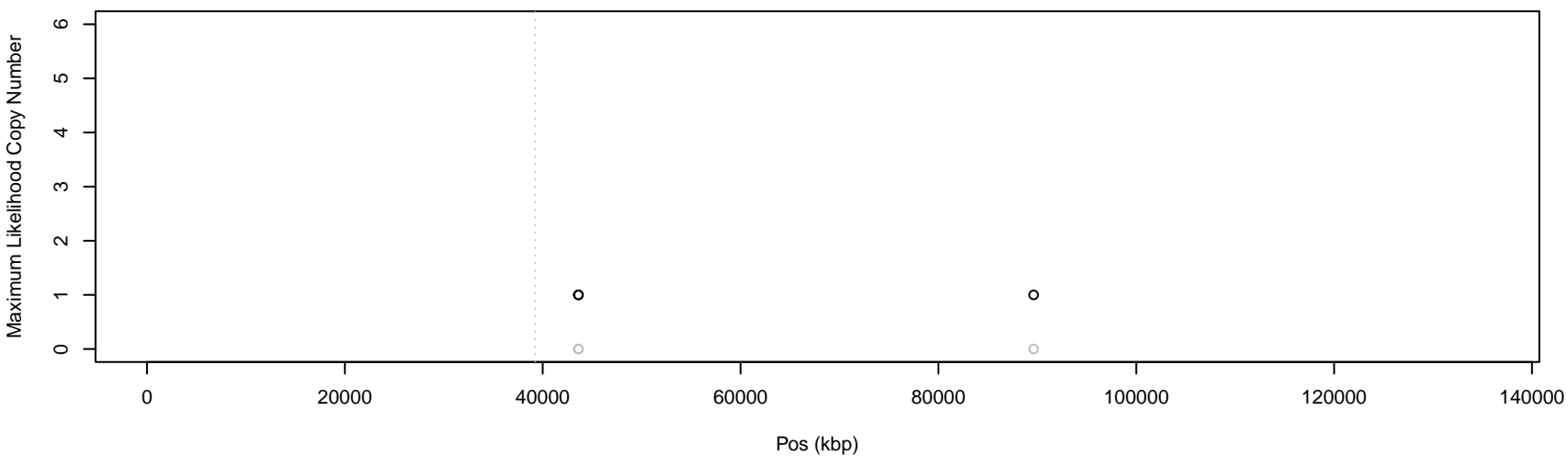
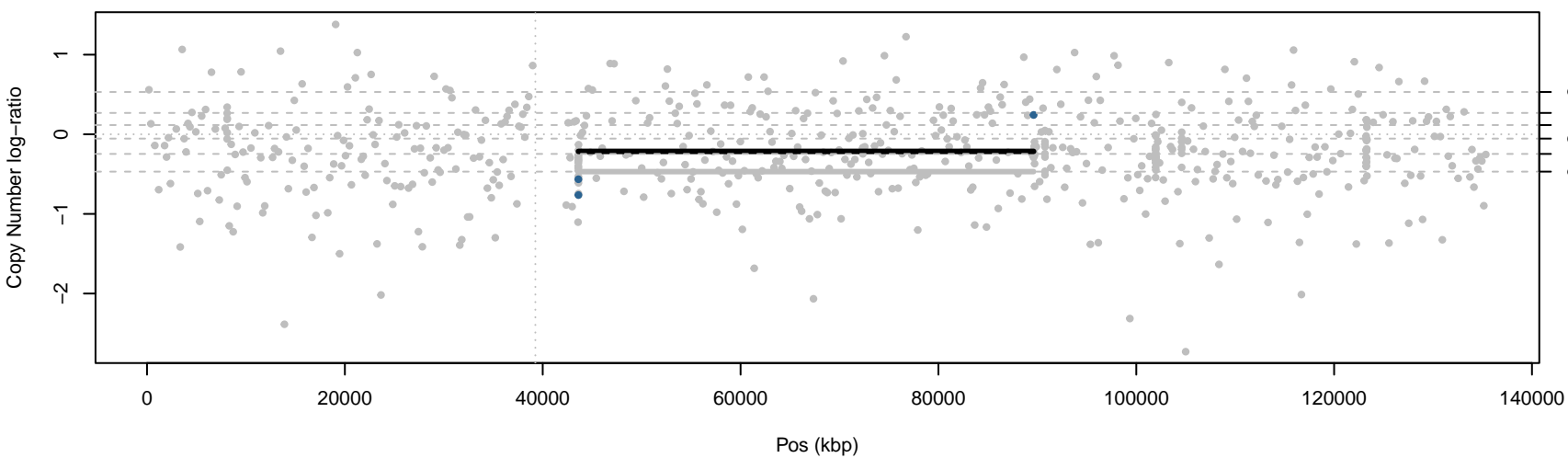
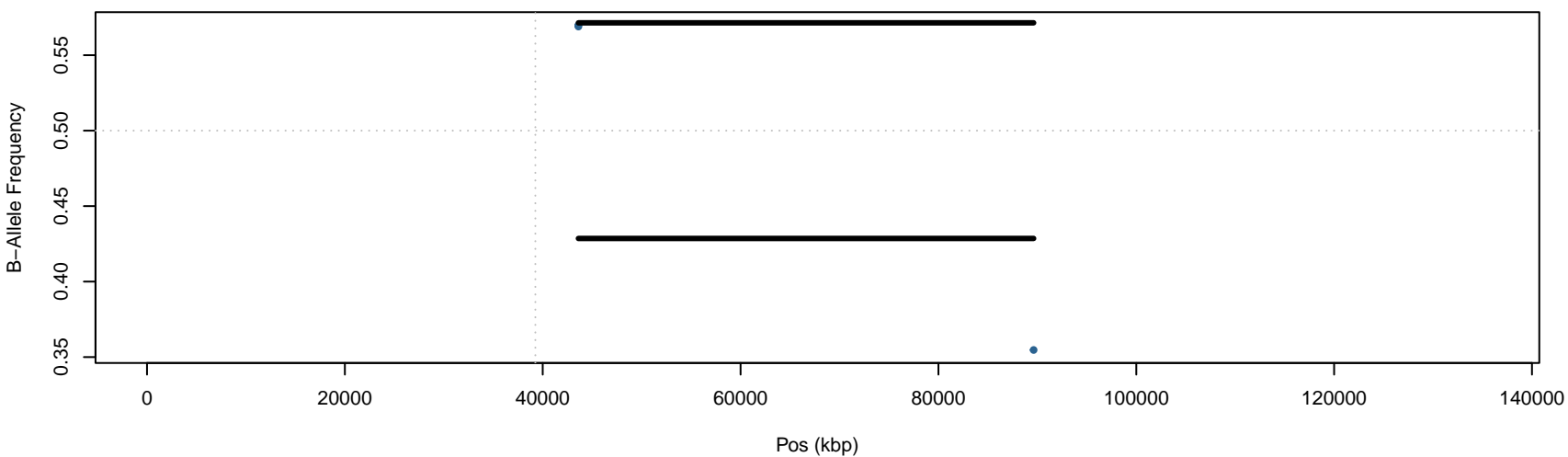
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 8



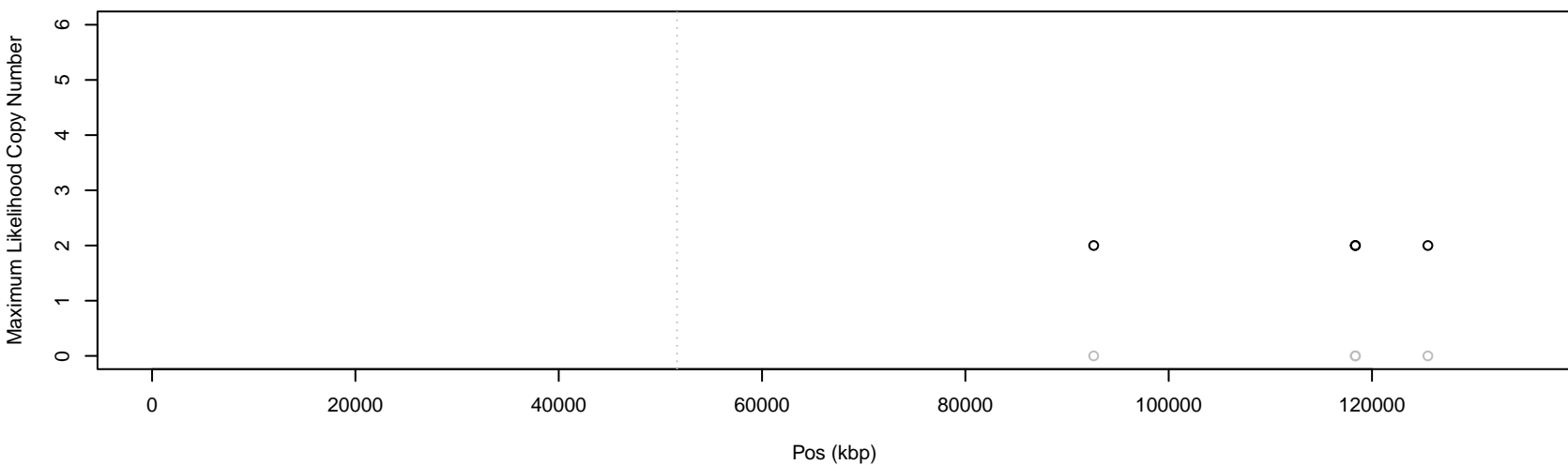
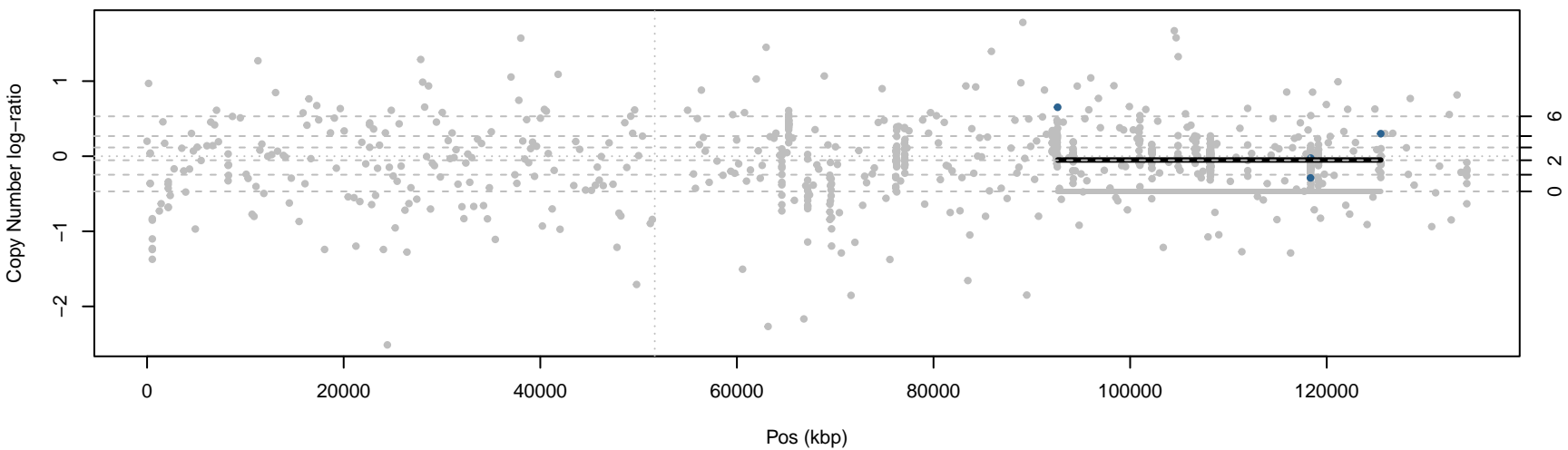
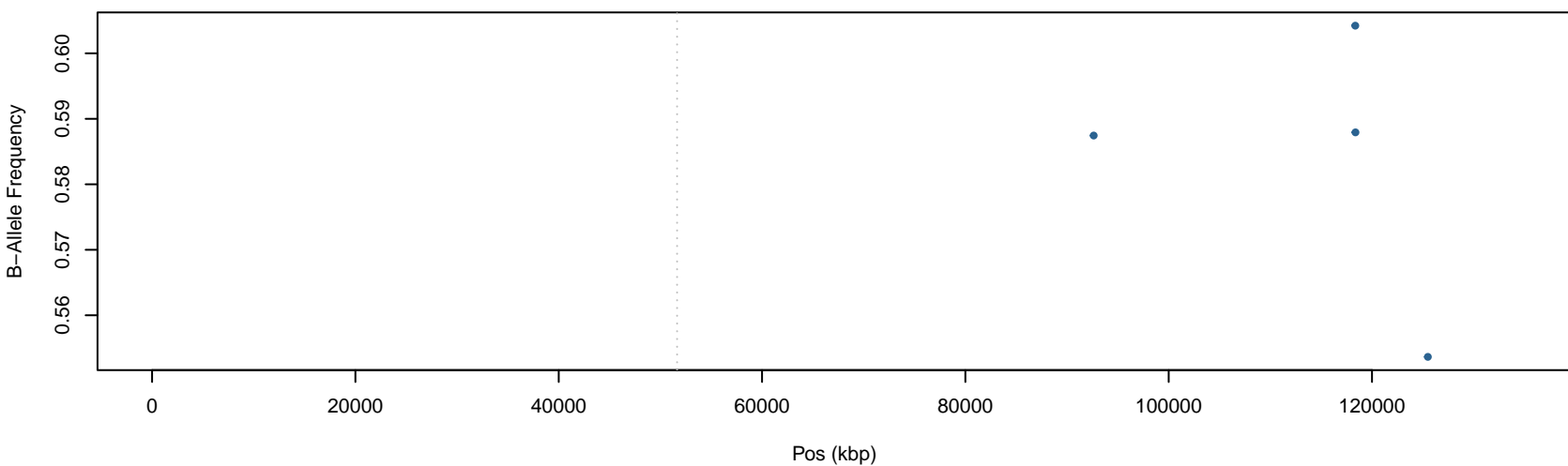
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 9



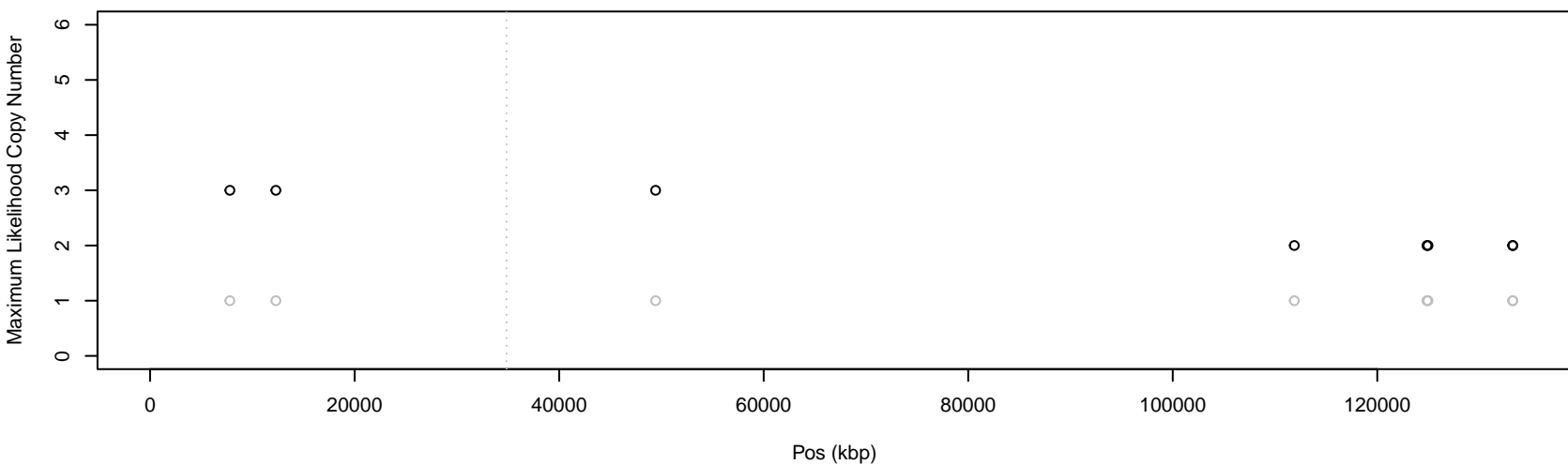
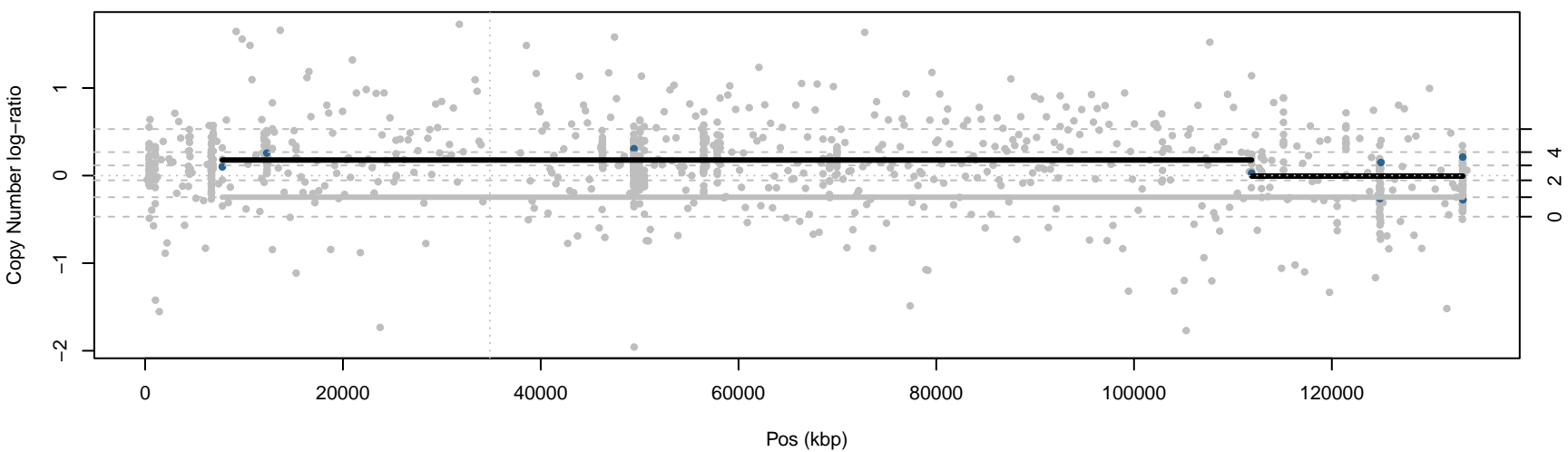
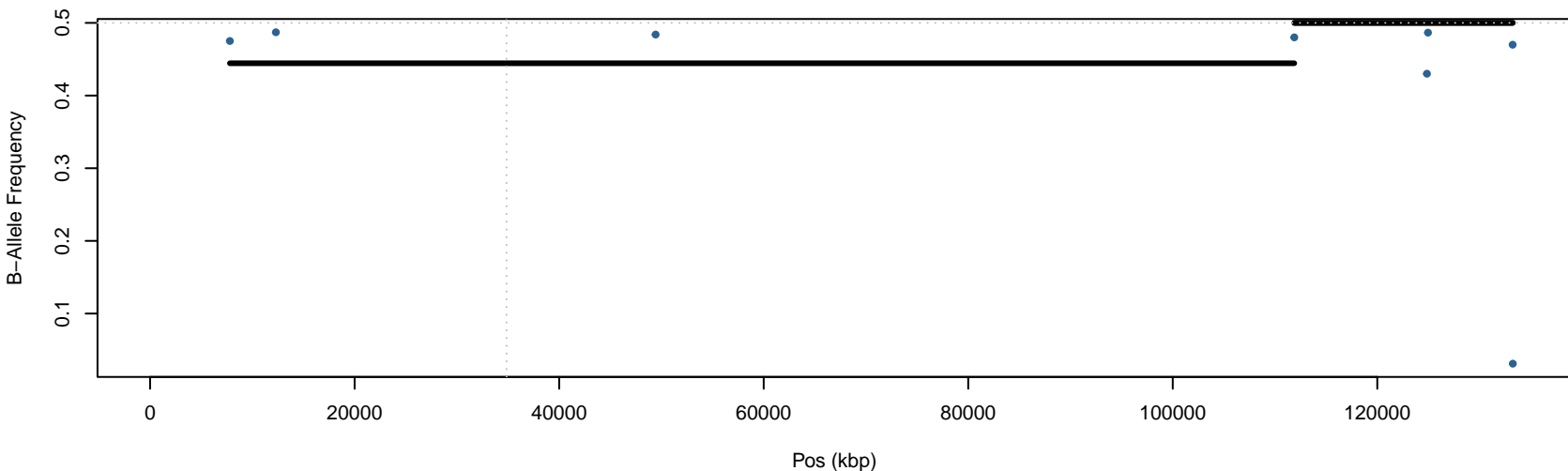
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 10



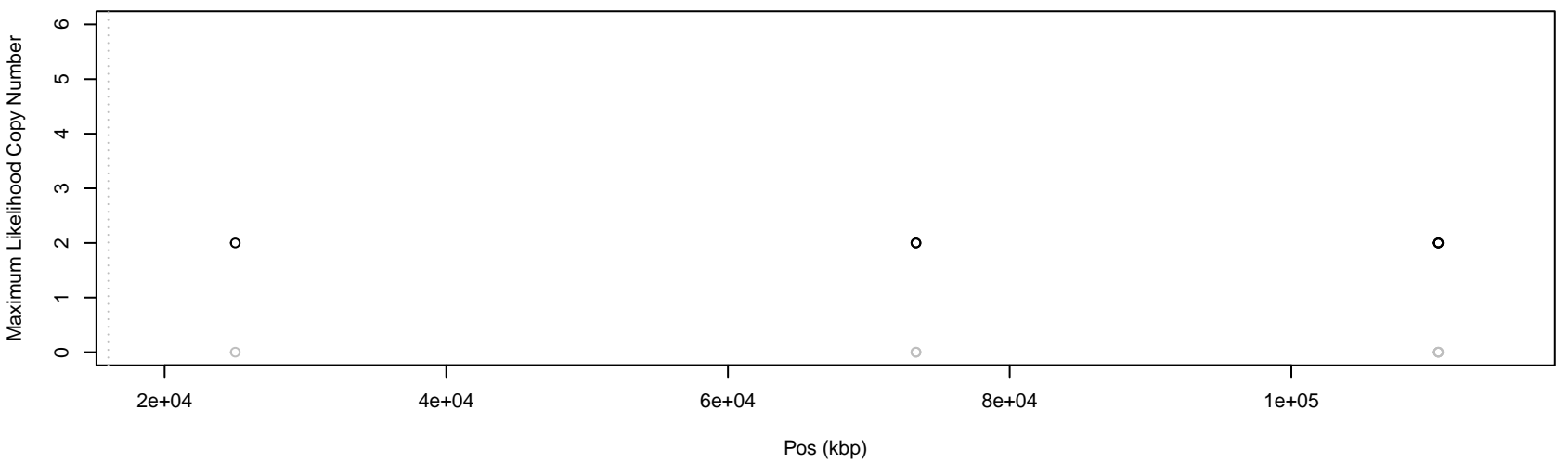
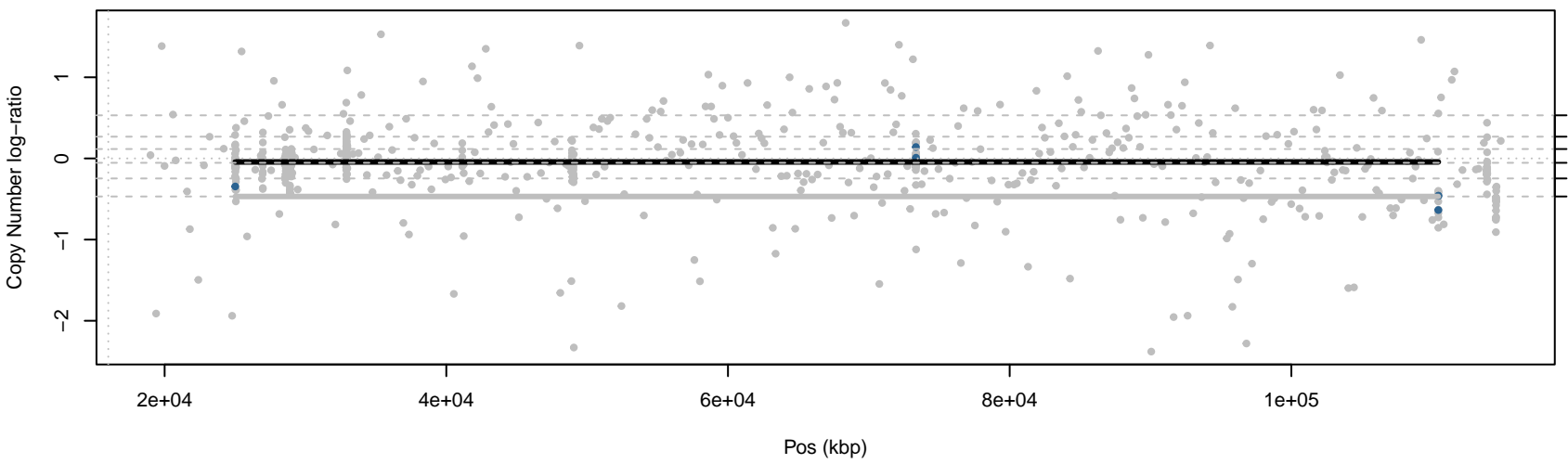
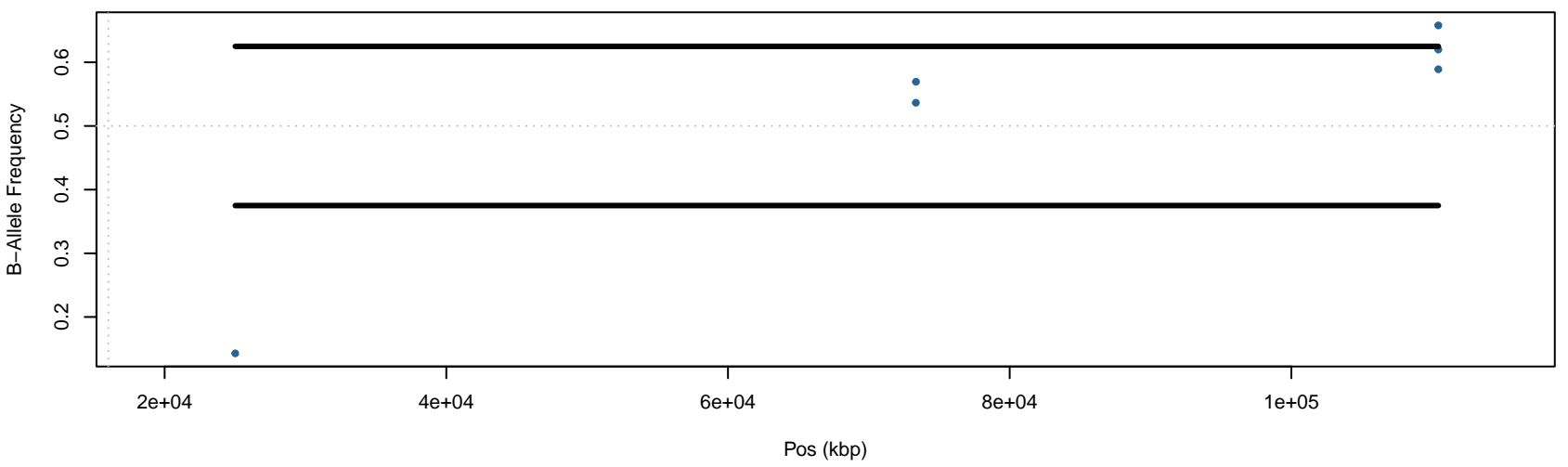
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 11



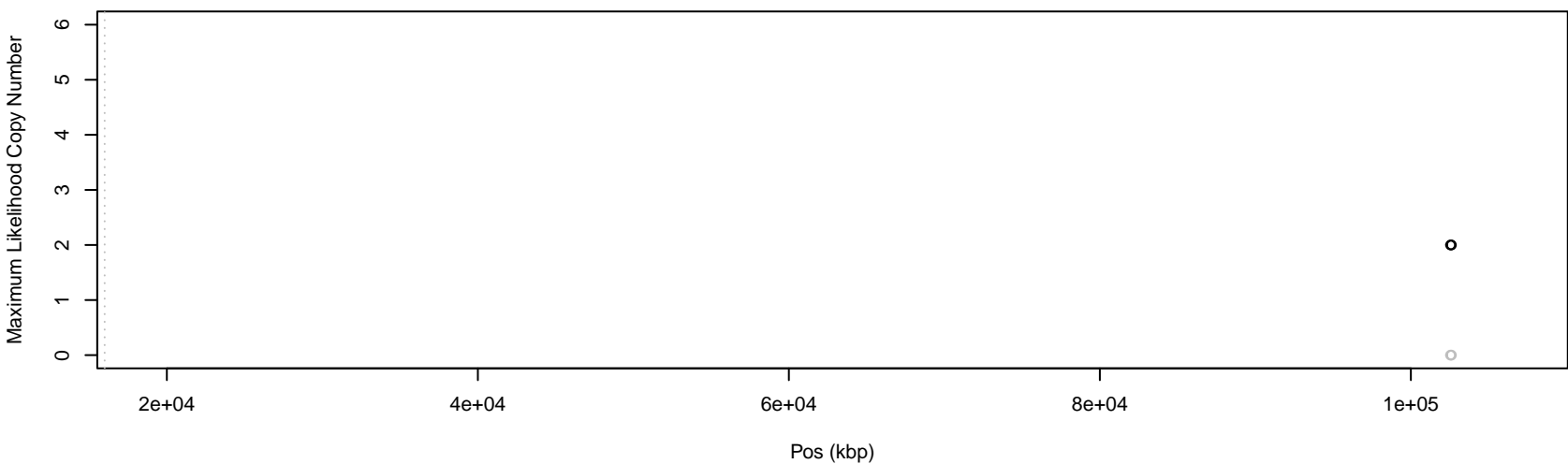
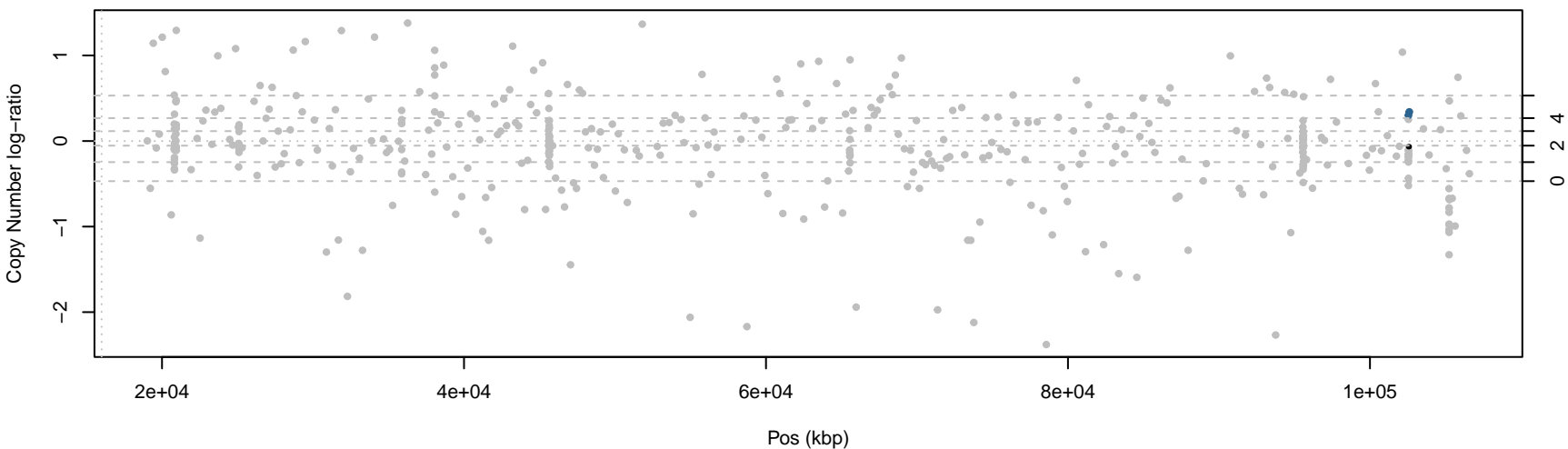
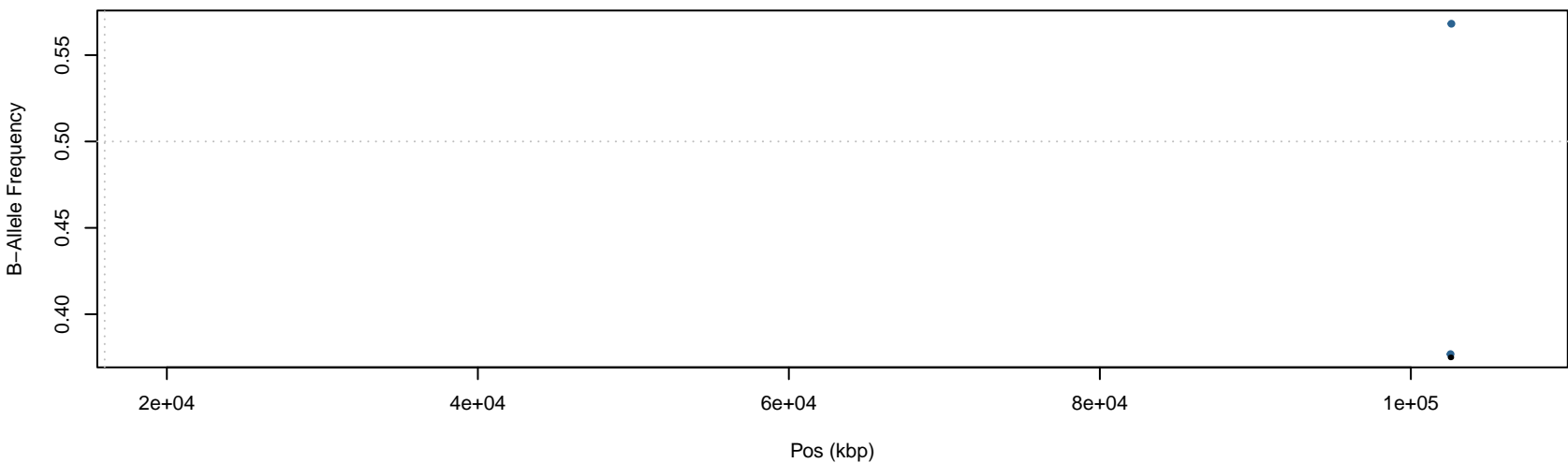
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 12



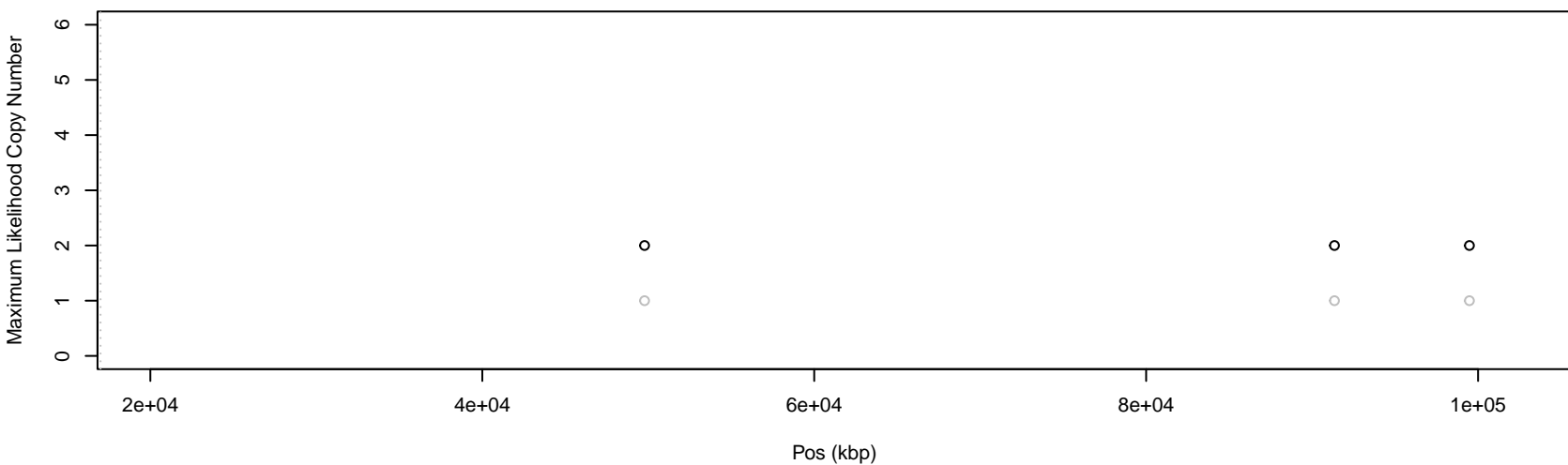
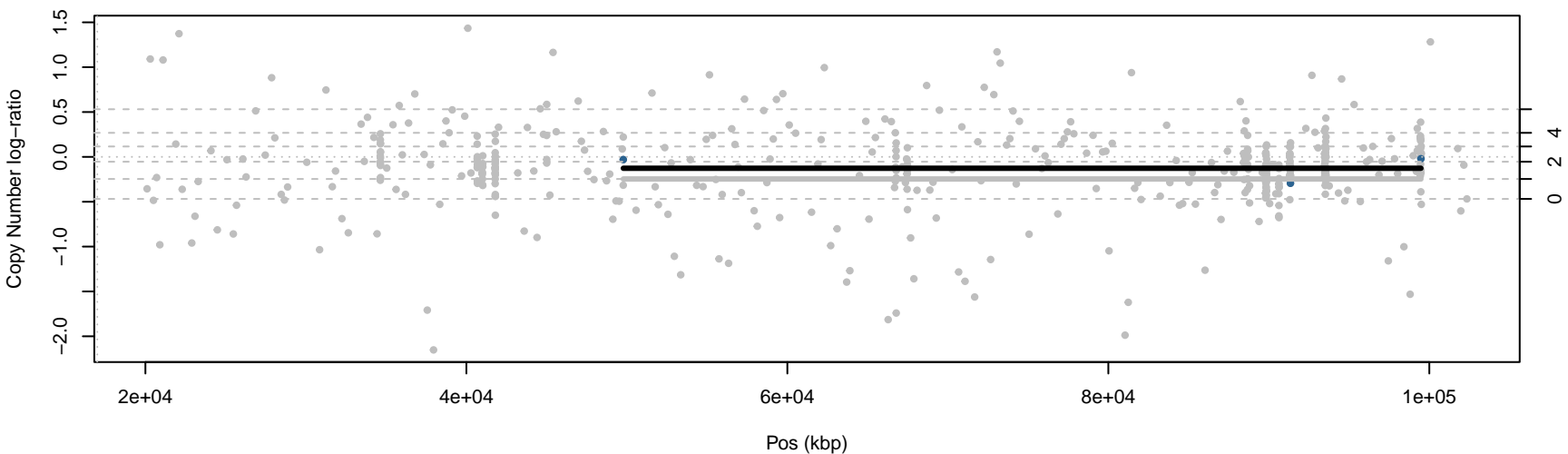
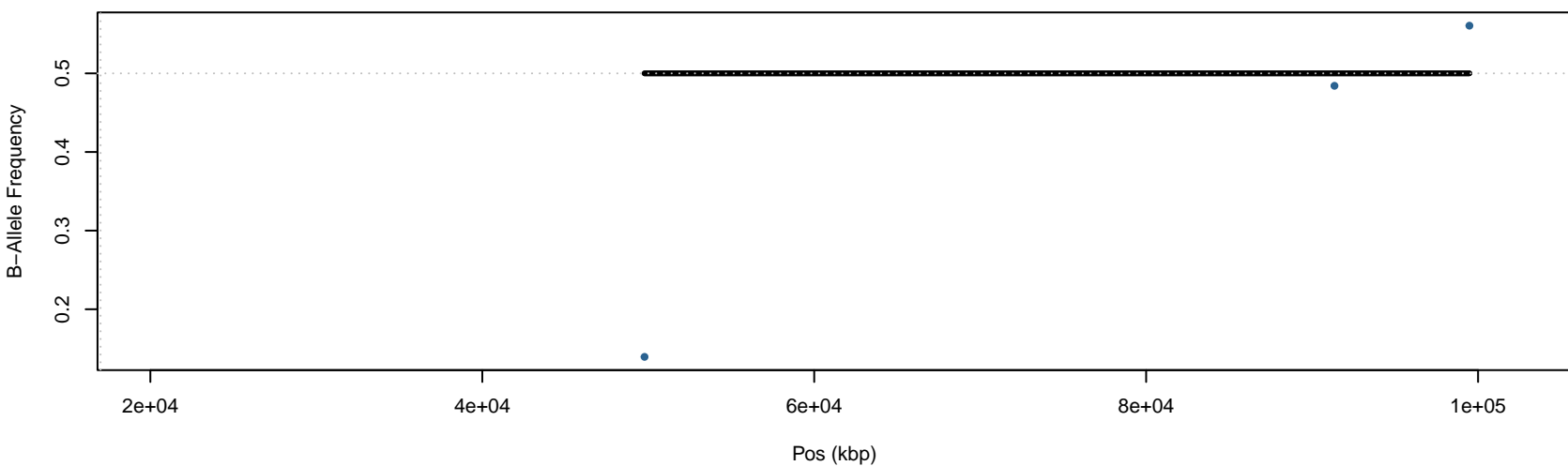
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 13



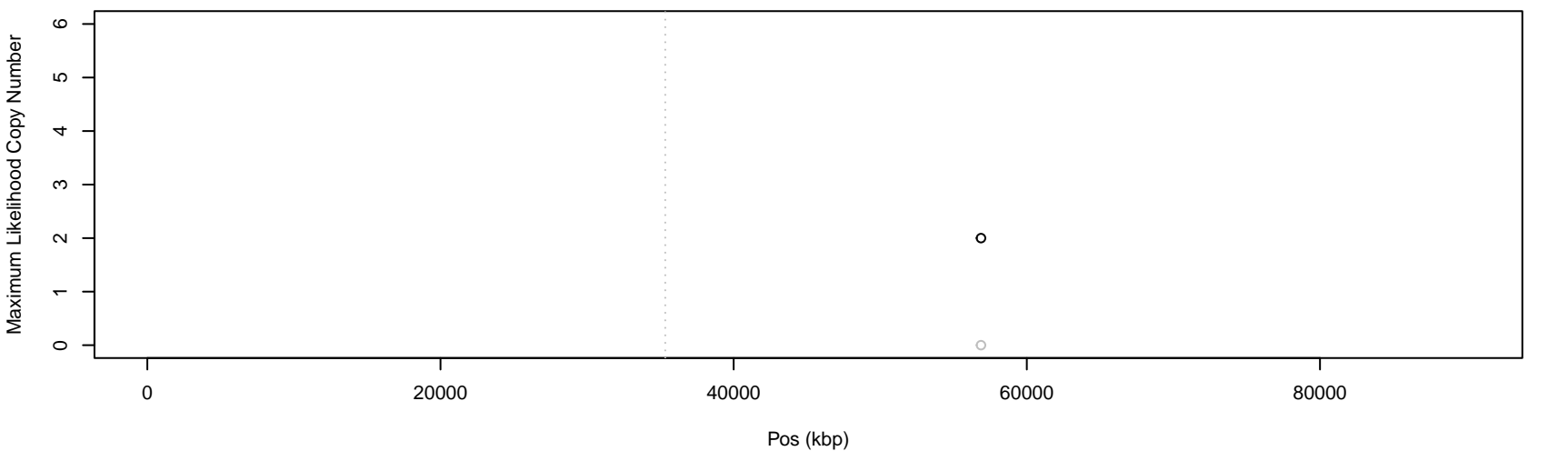
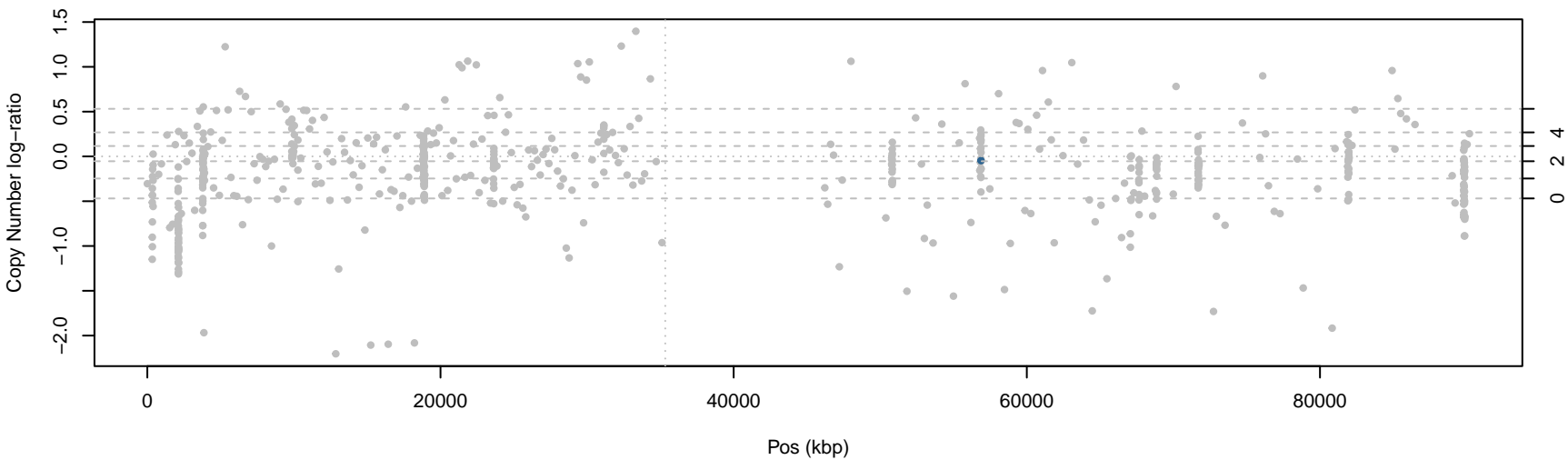
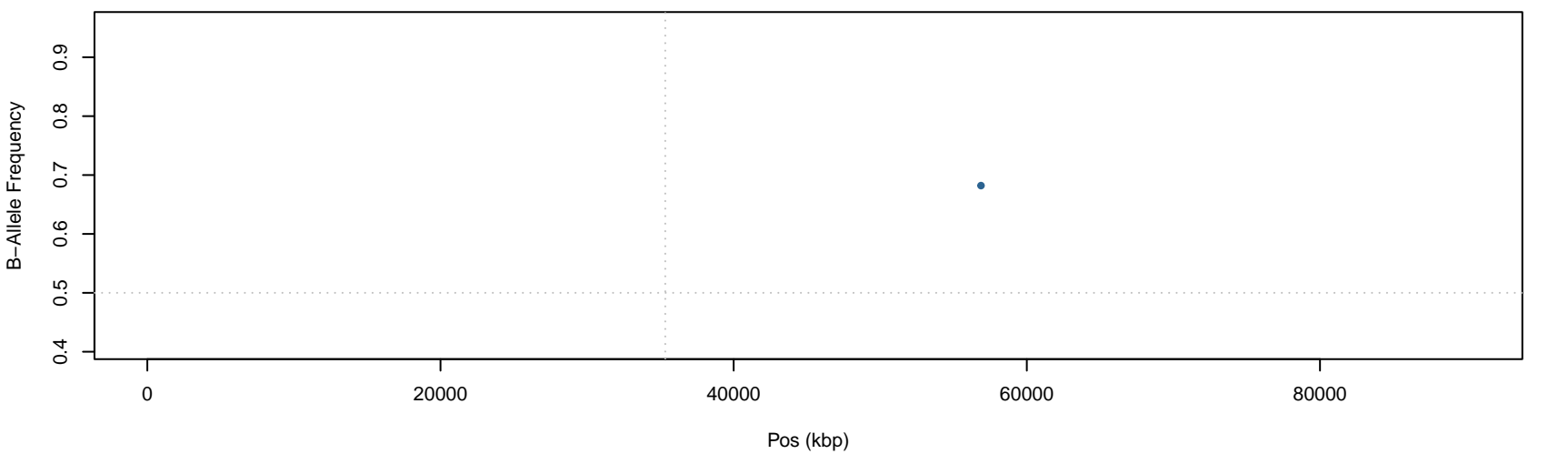
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 14



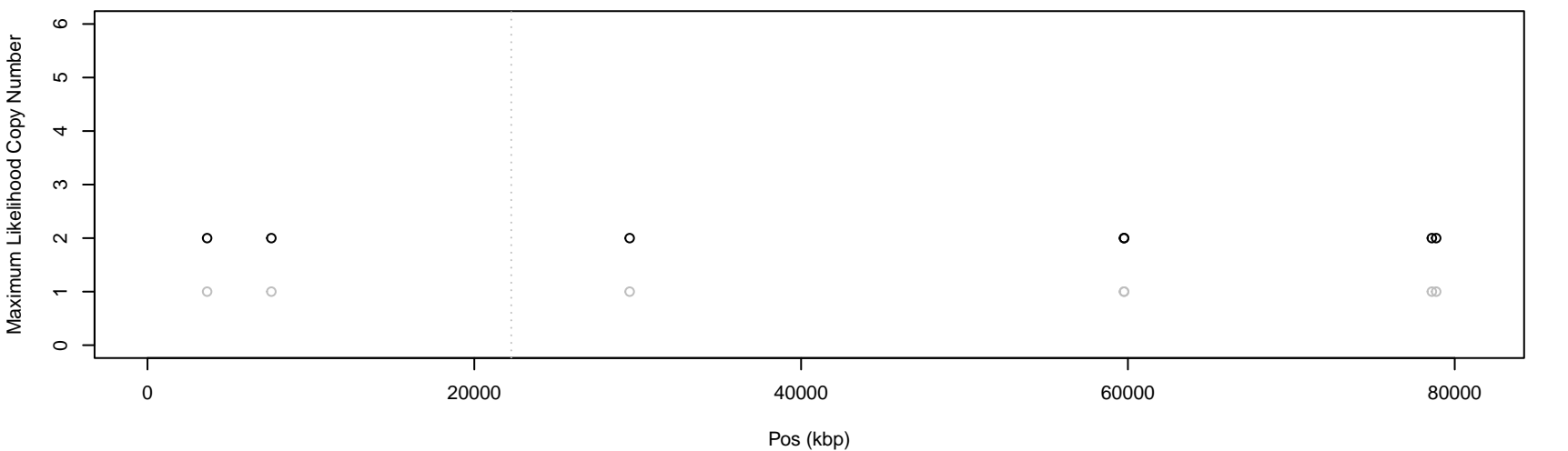
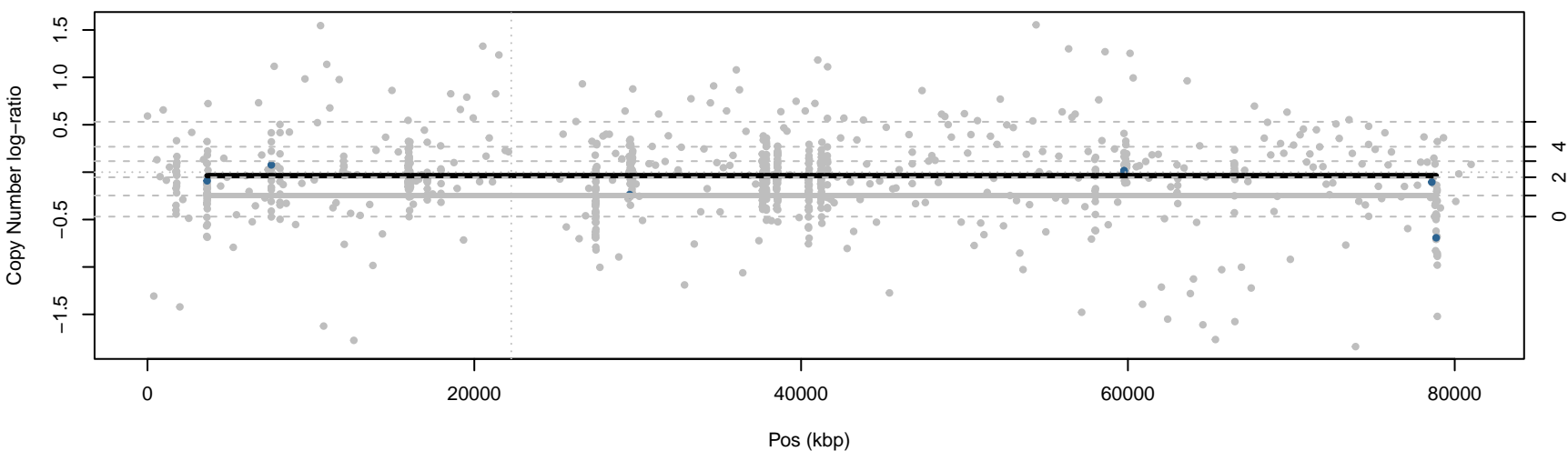
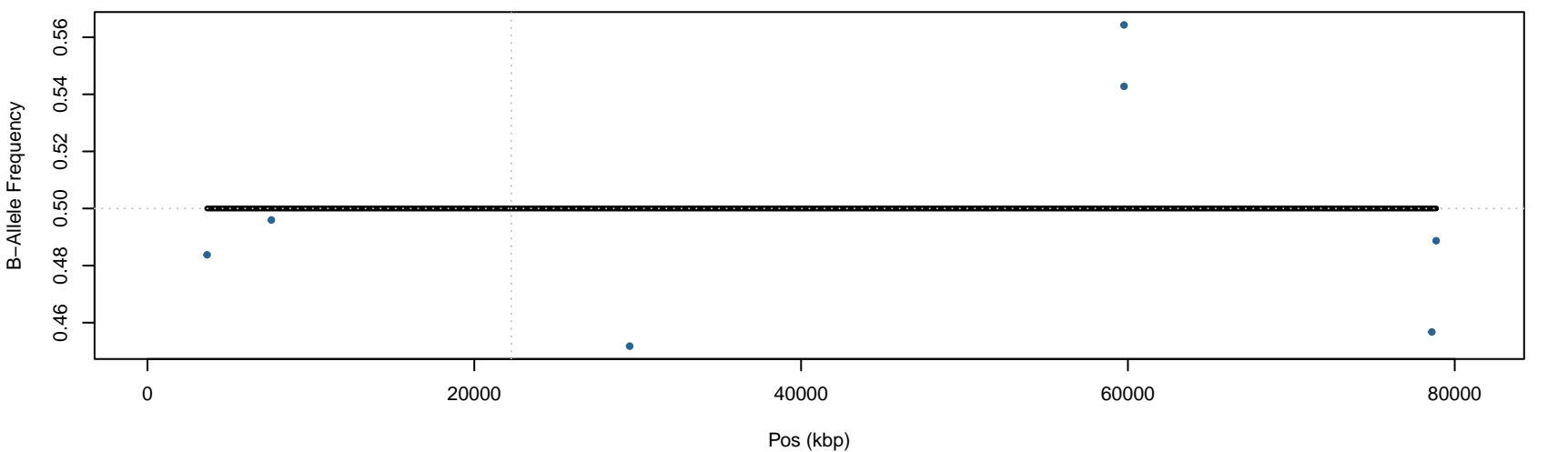
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 15



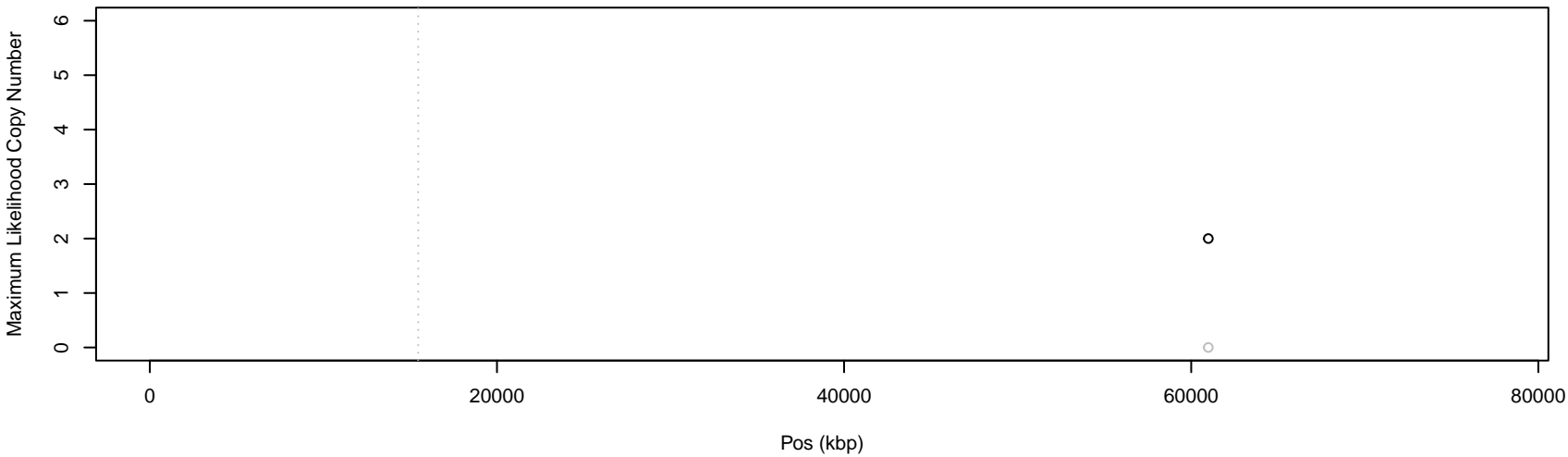
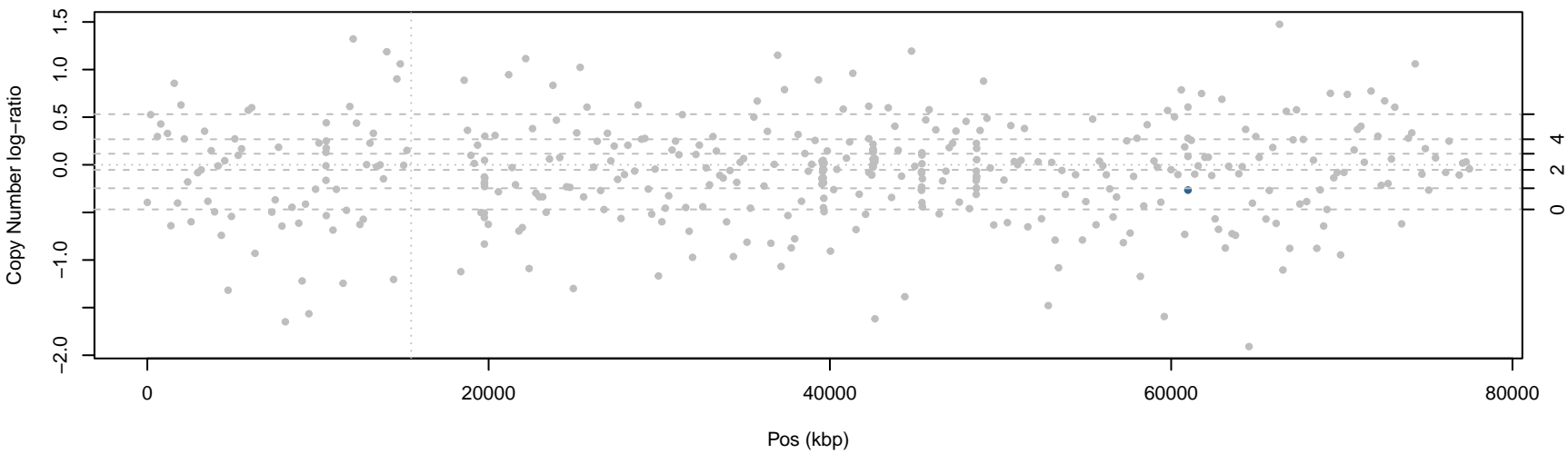
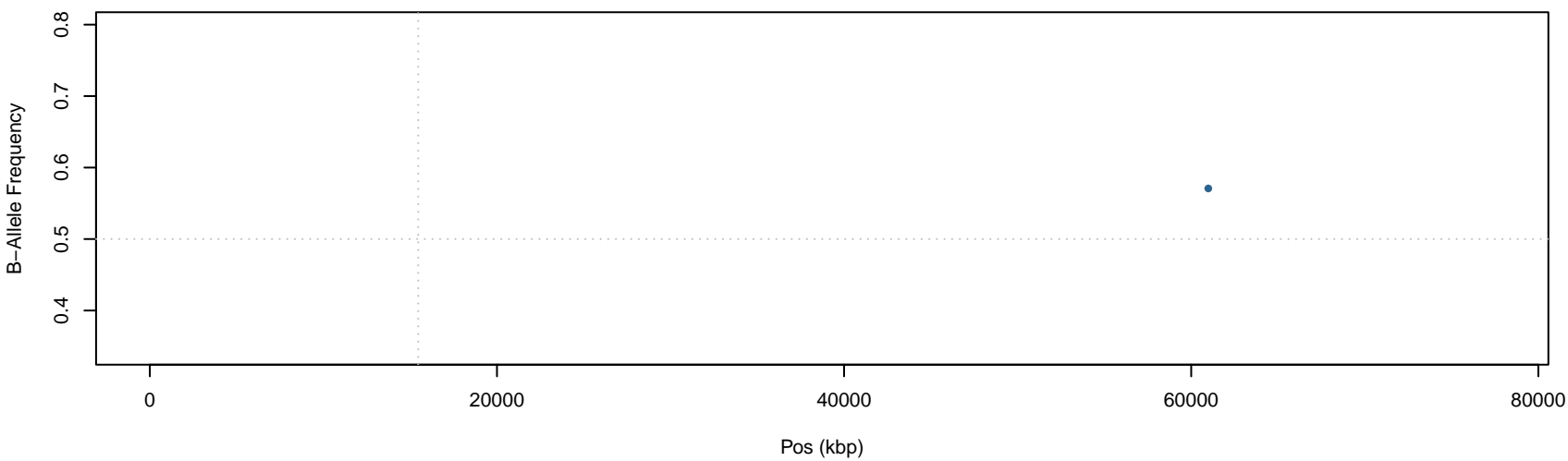
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 16



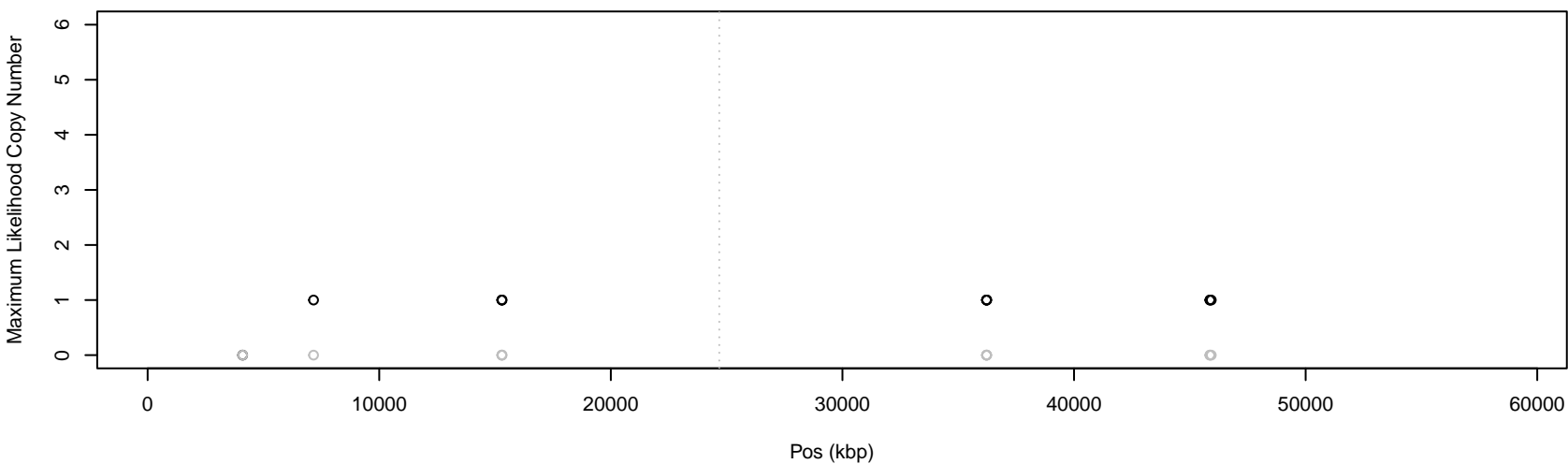
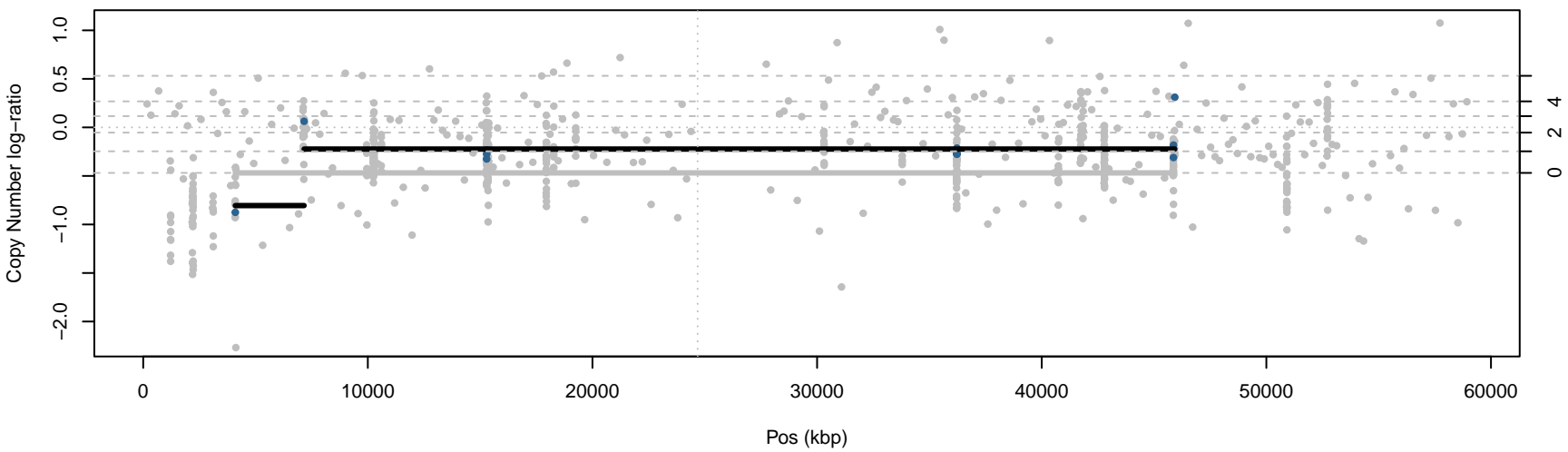
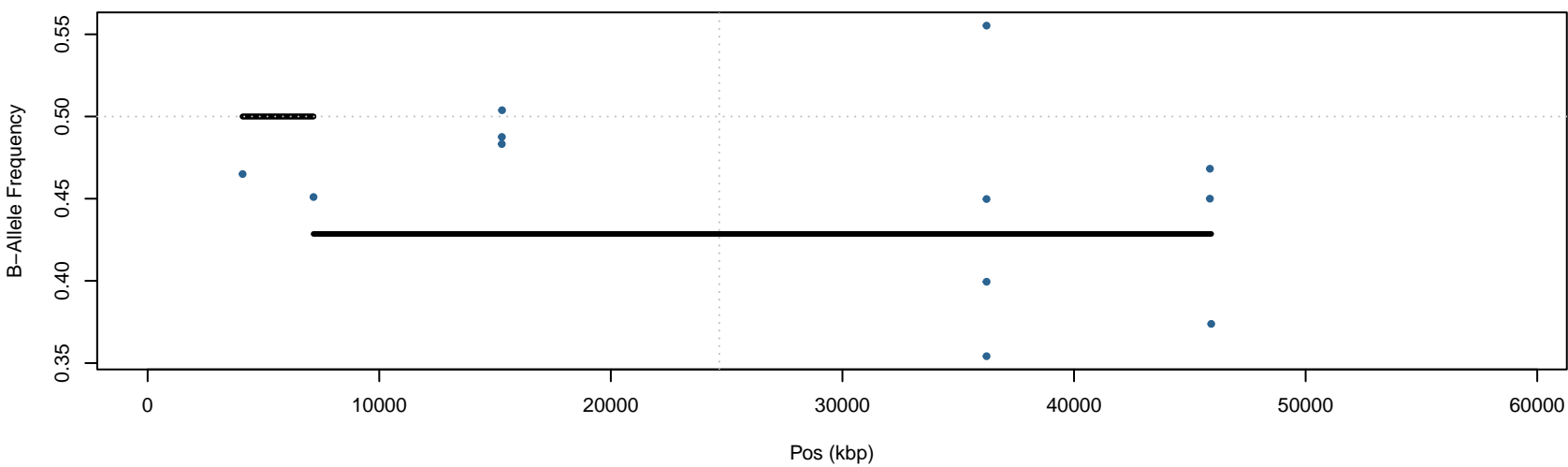
Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 17



Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 18



Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 19



Purity: 0.25 Tumor ploidy: 2.304 SNV log-likelihood: 16.1 GoF: 95.6% Mean coverage: 397;595 Chromosome: 20

