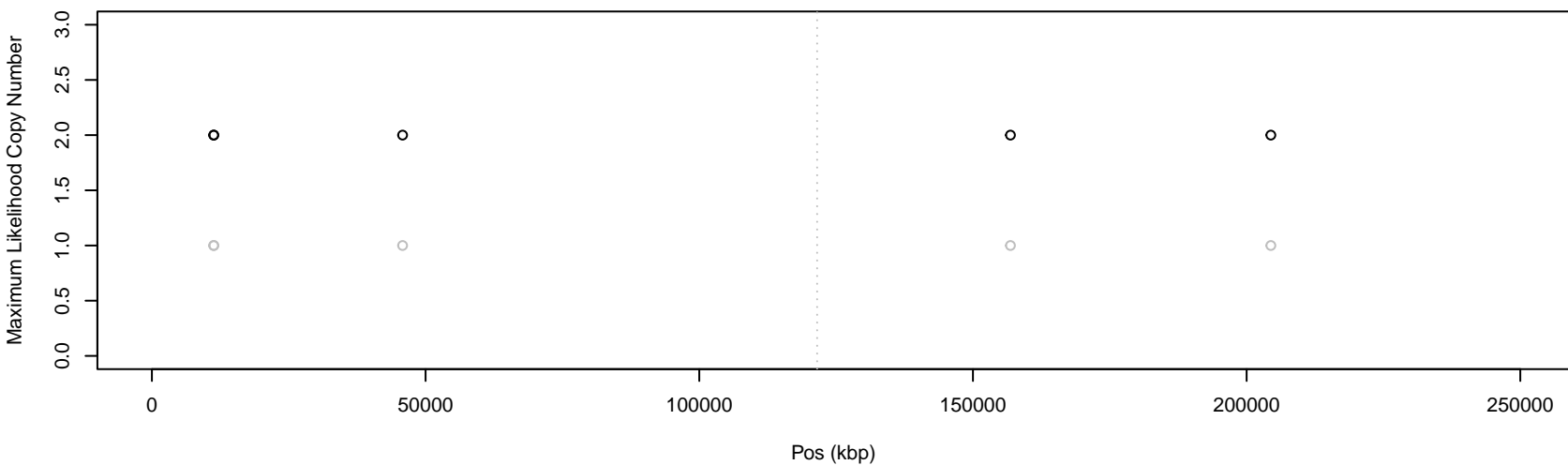
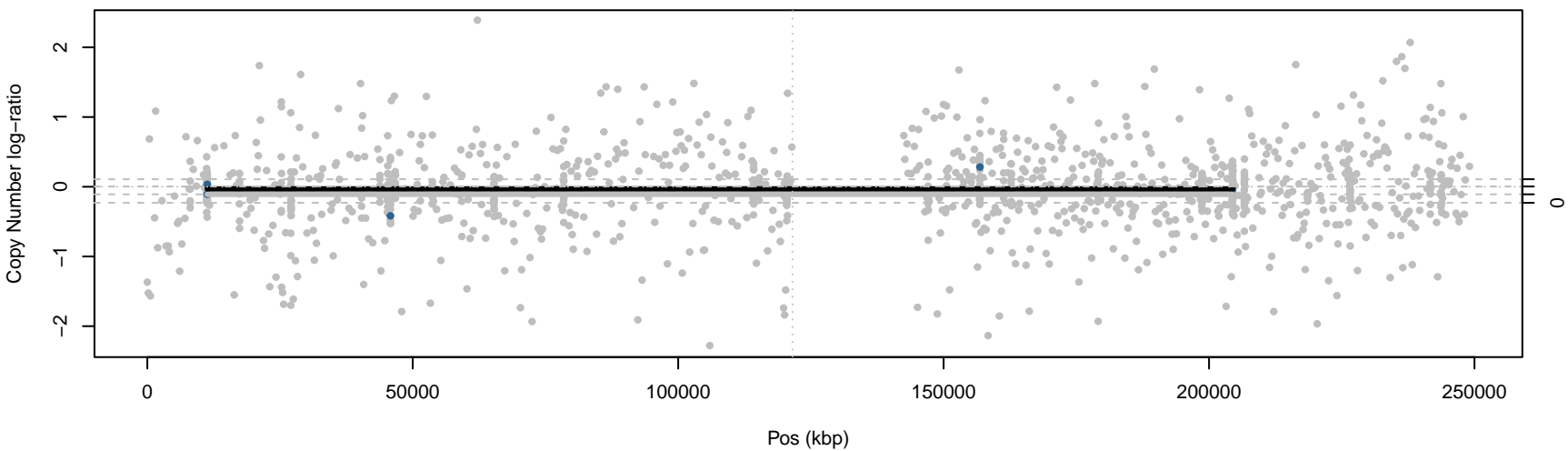
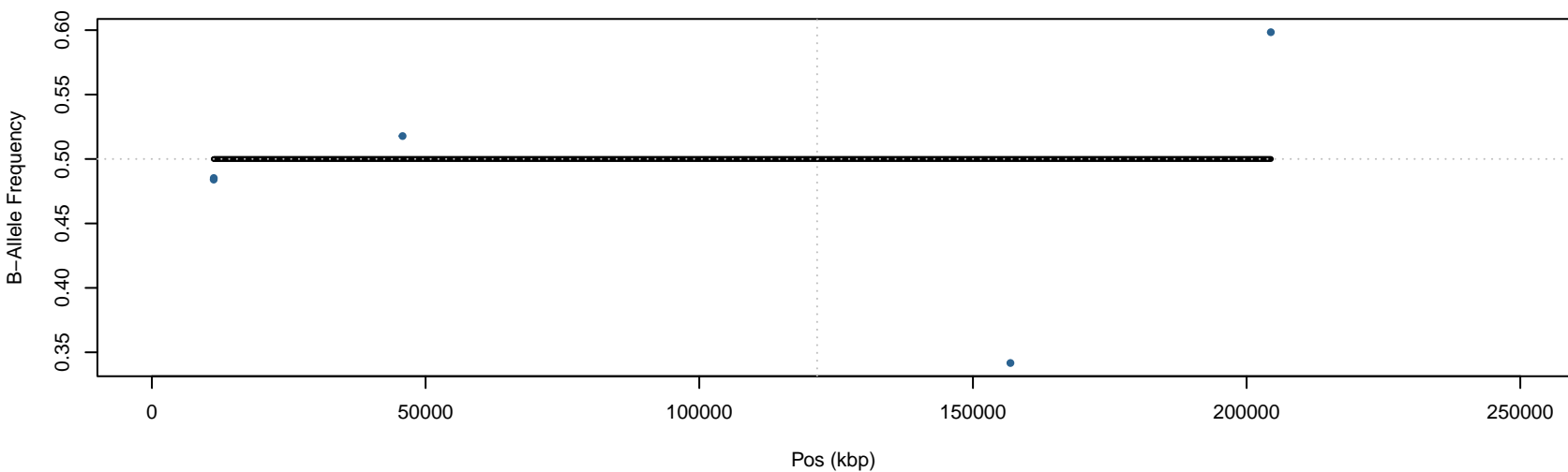
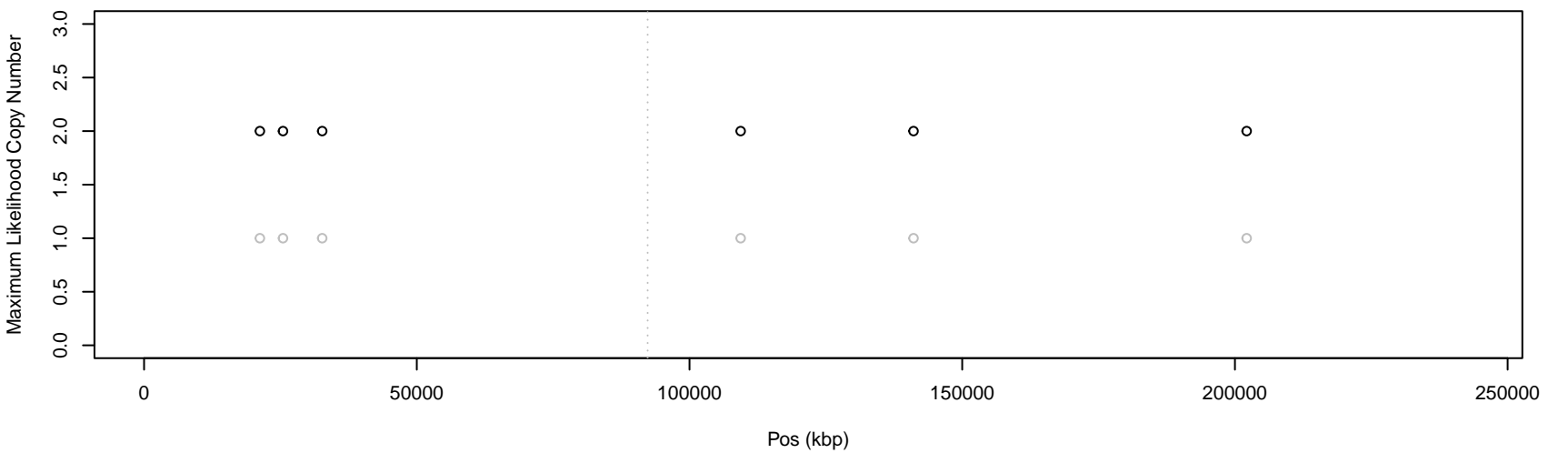
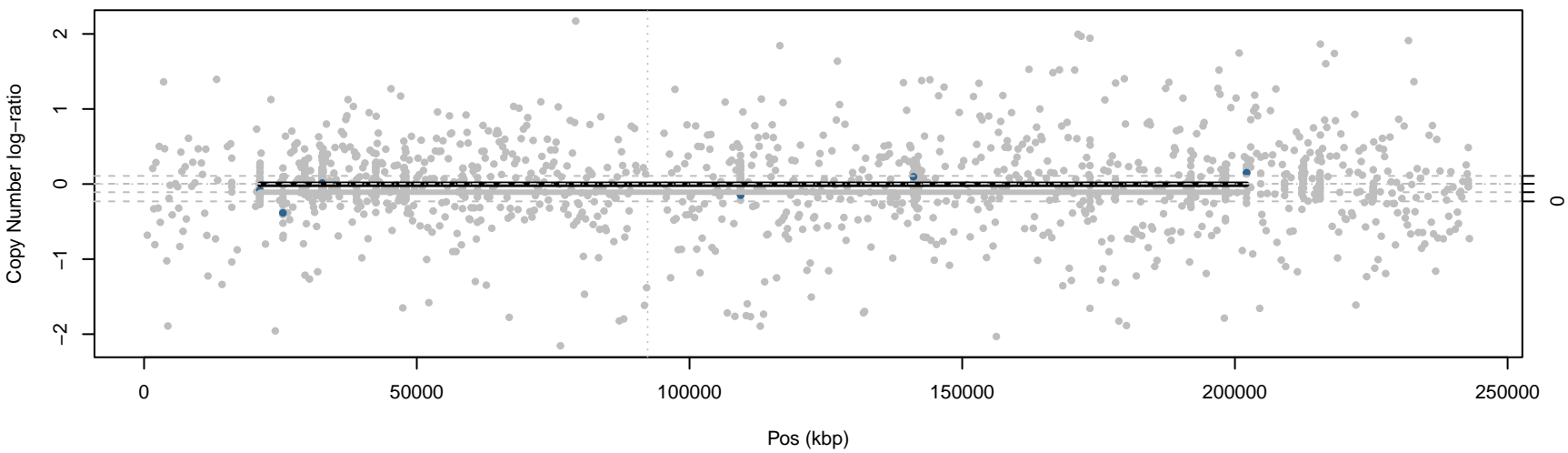
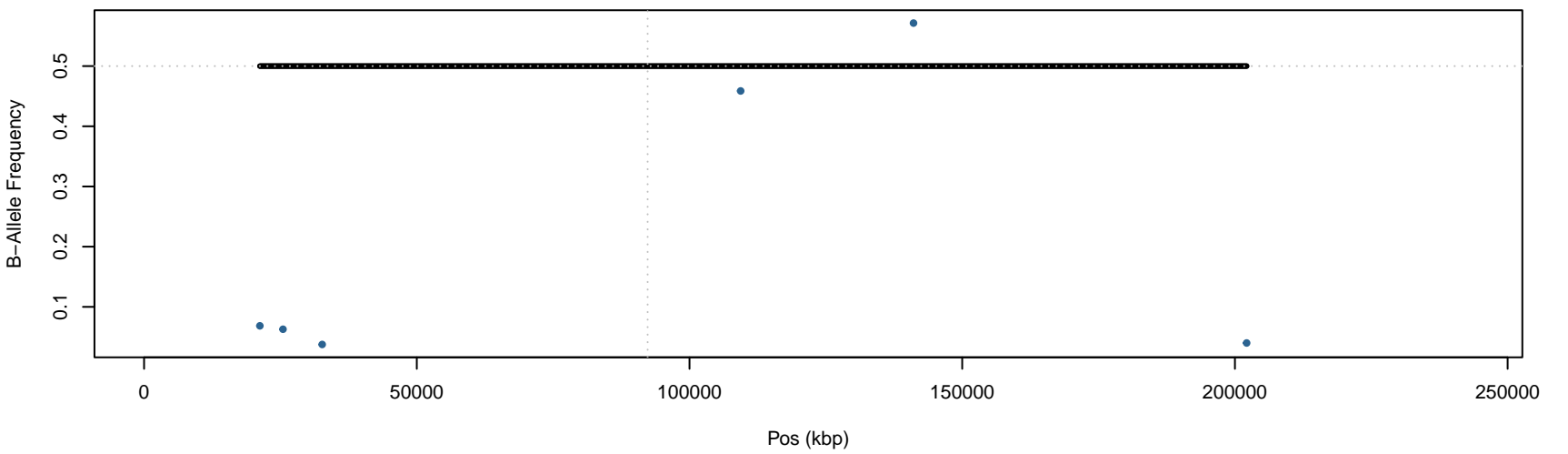


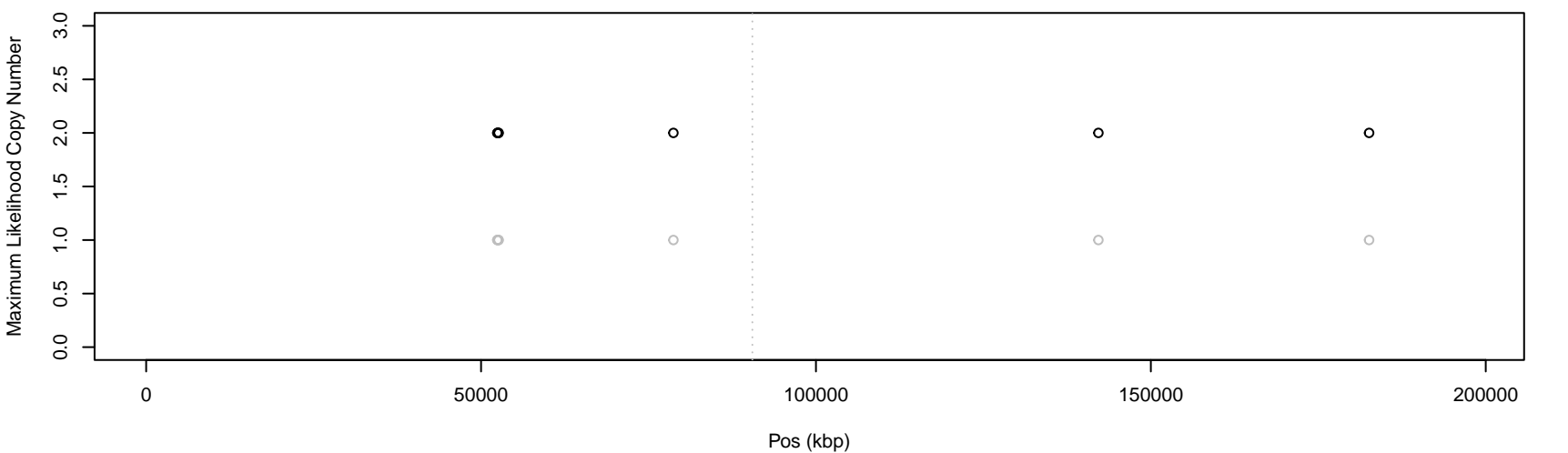
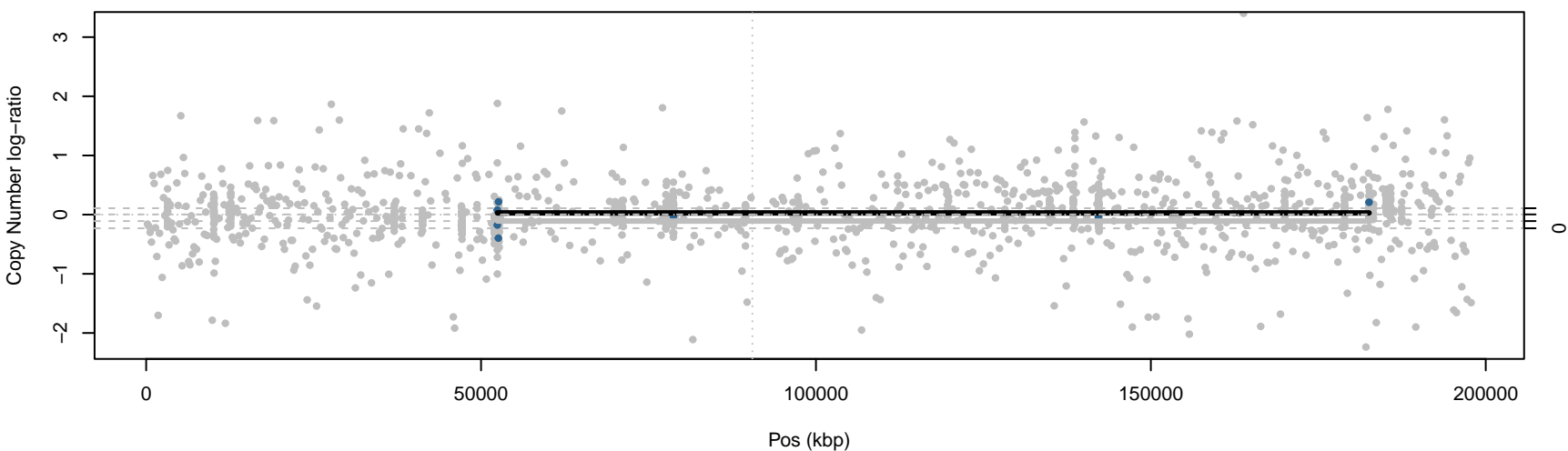
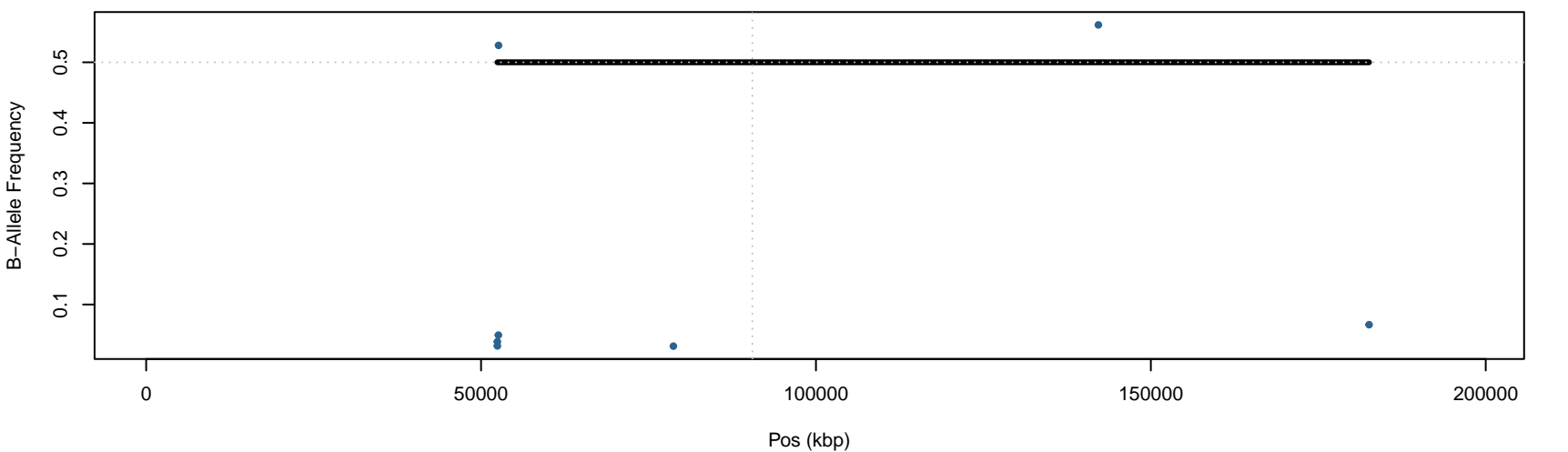
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 1



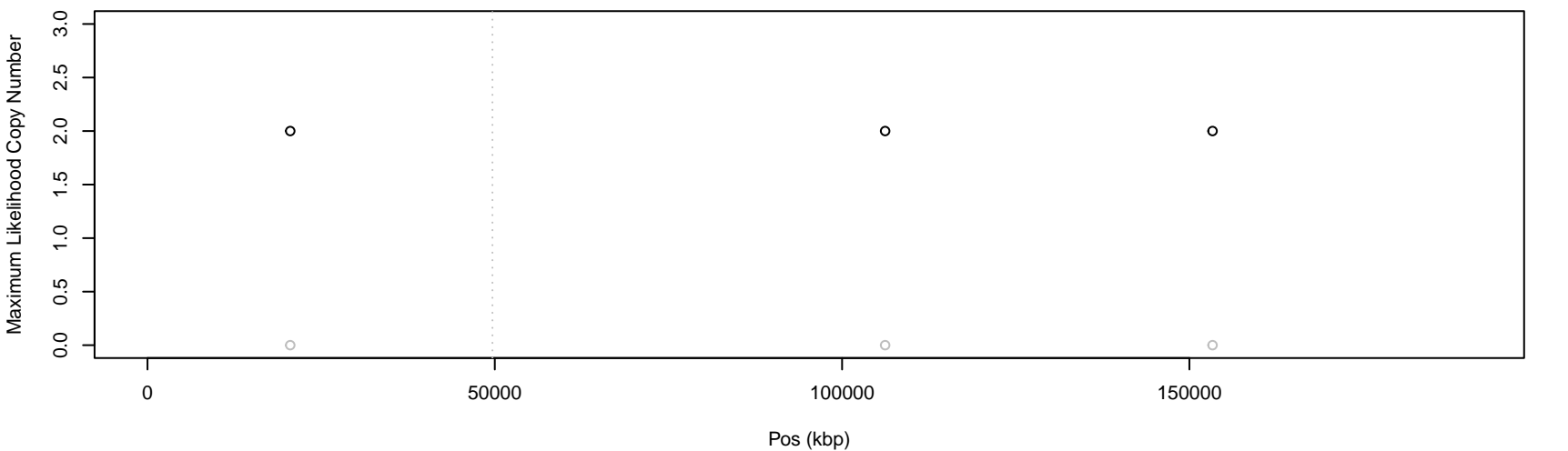
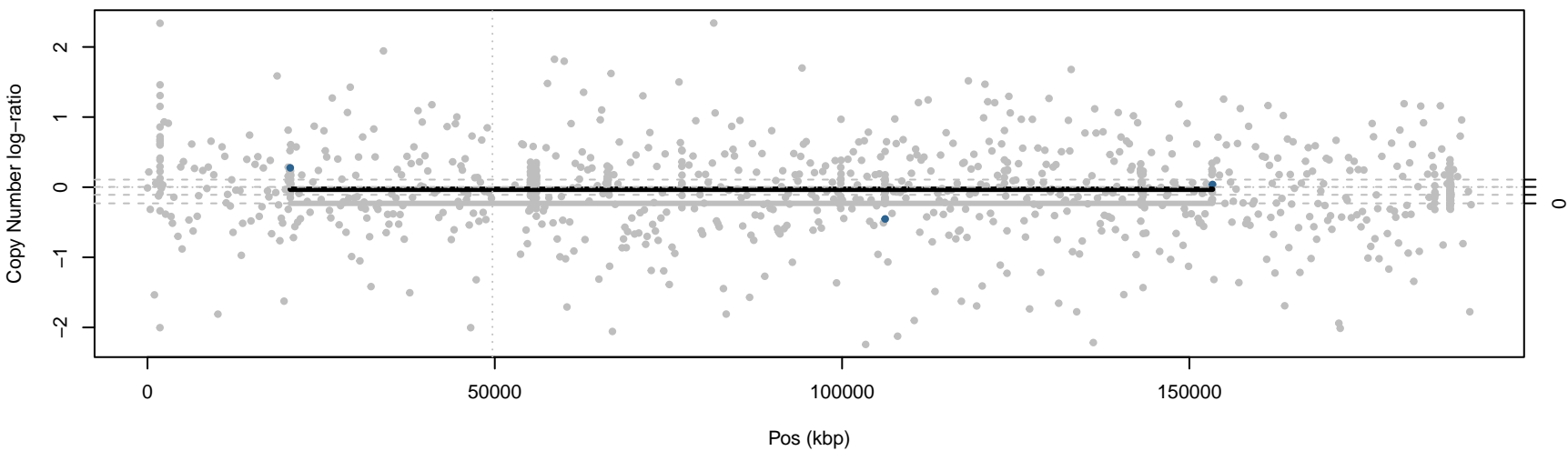
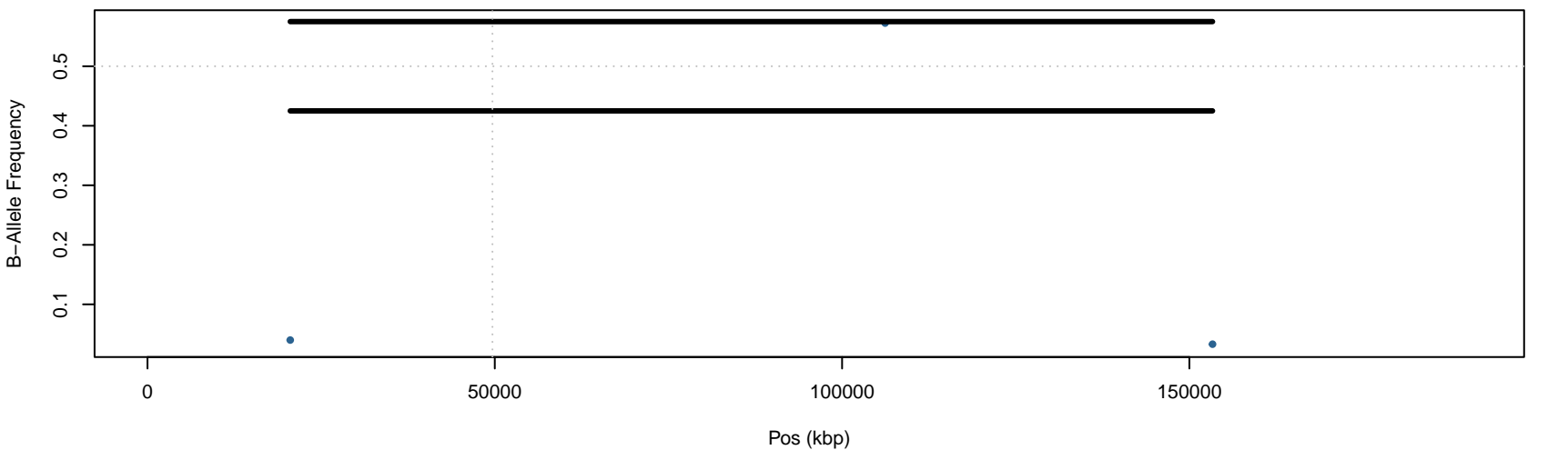
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 2



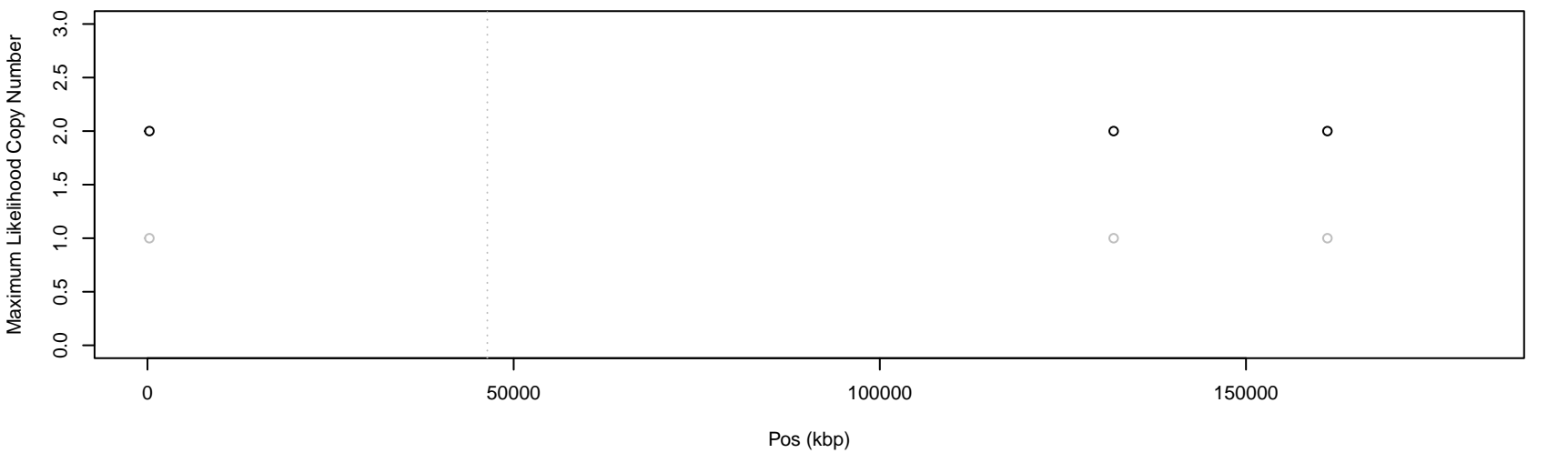
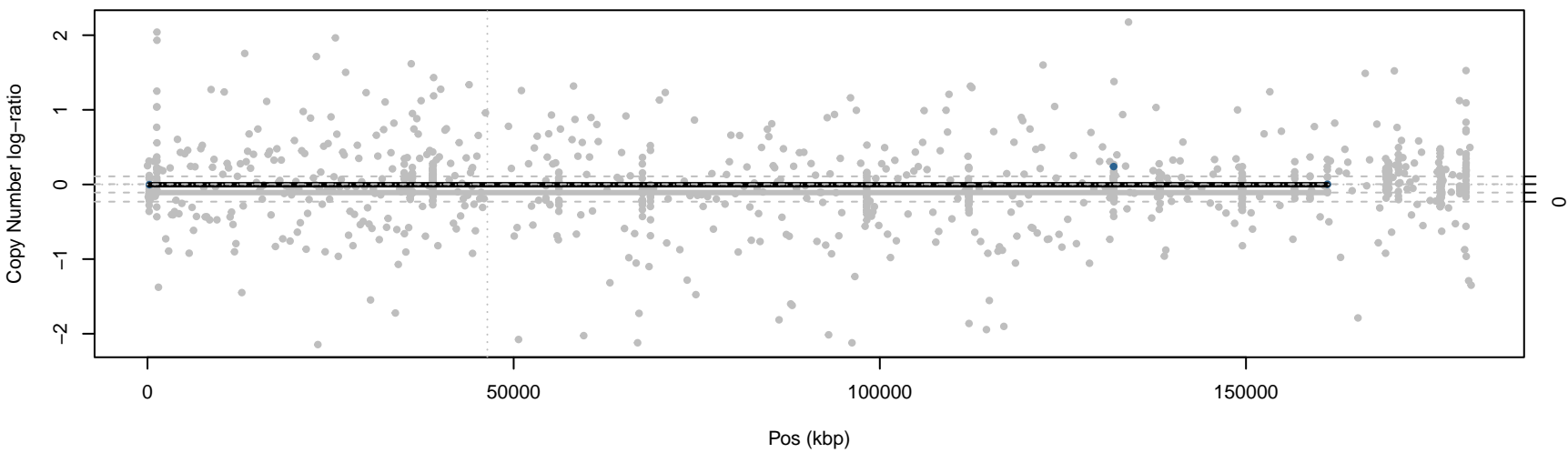
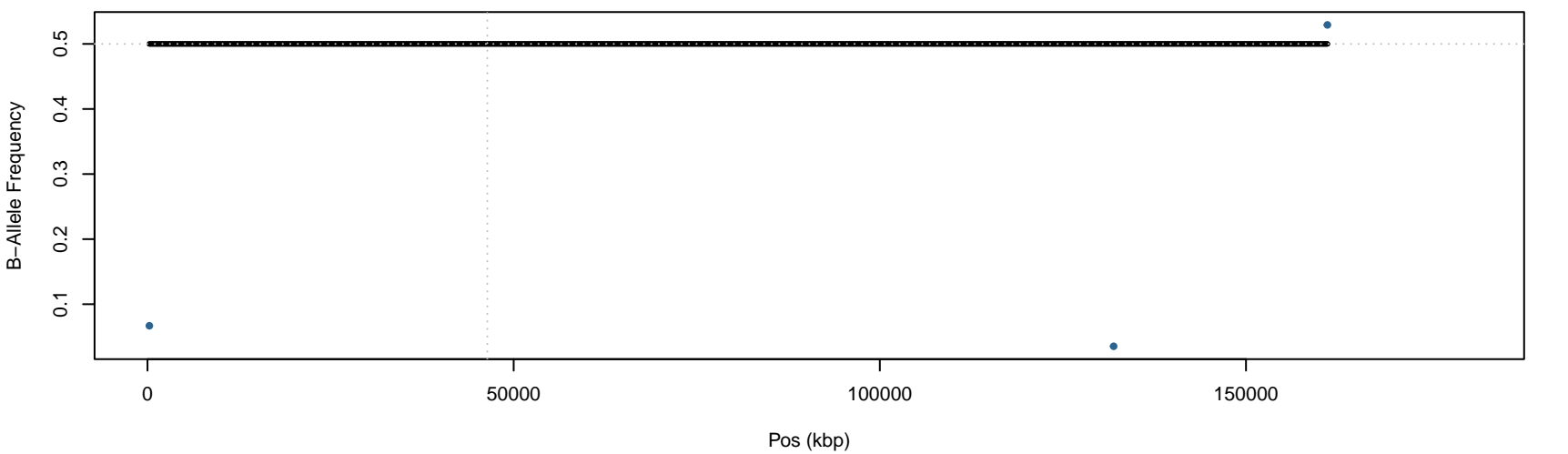
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 3



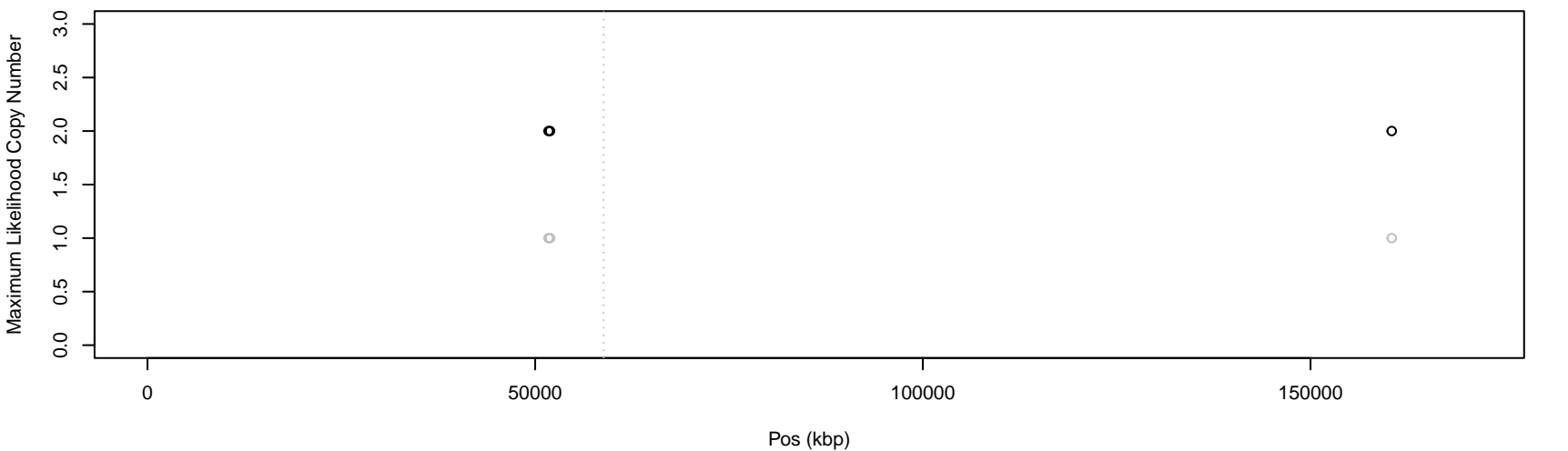
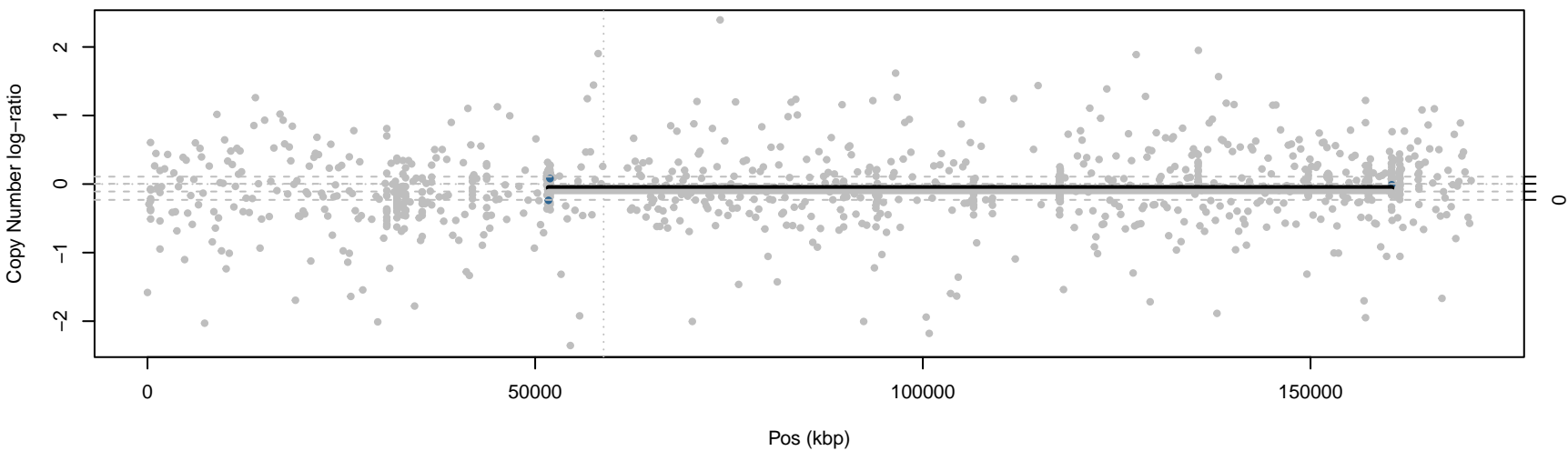
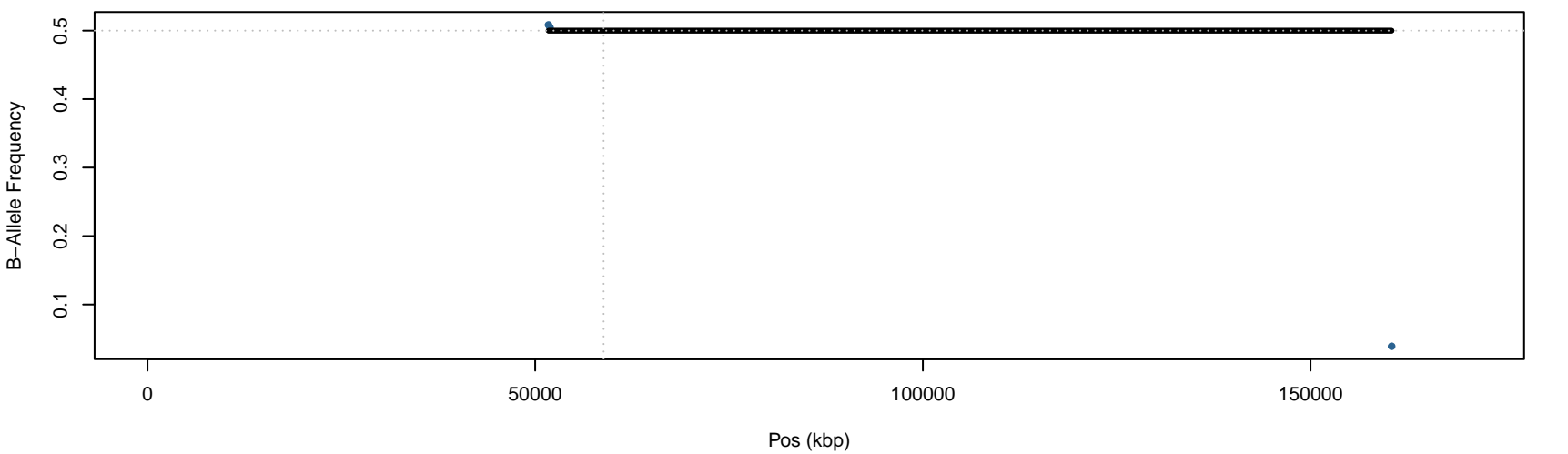
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 4



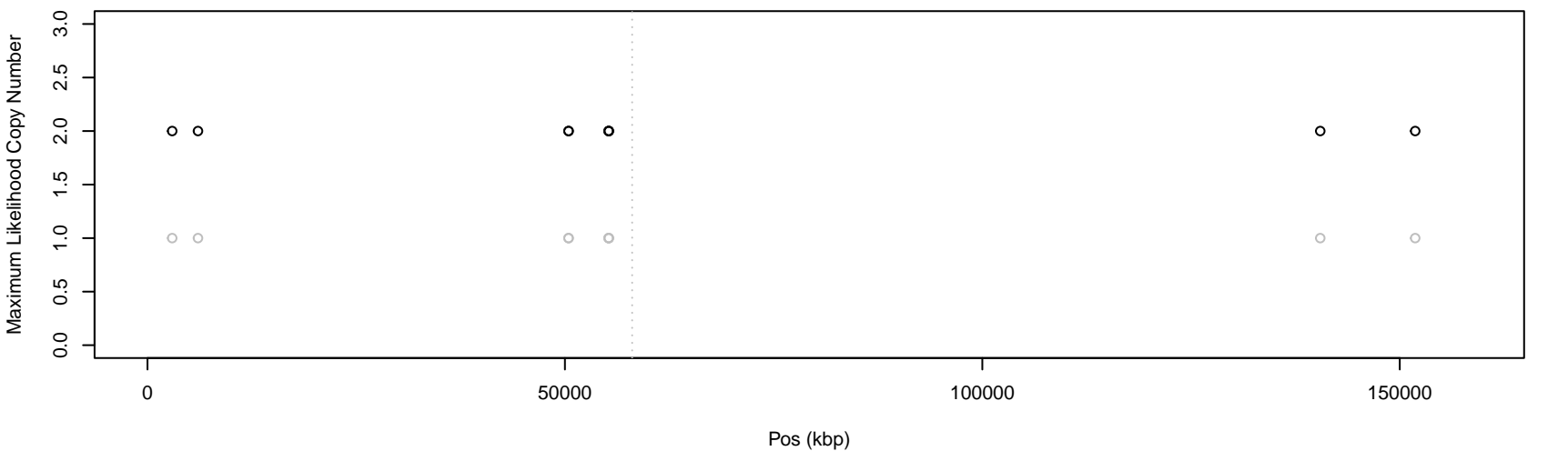
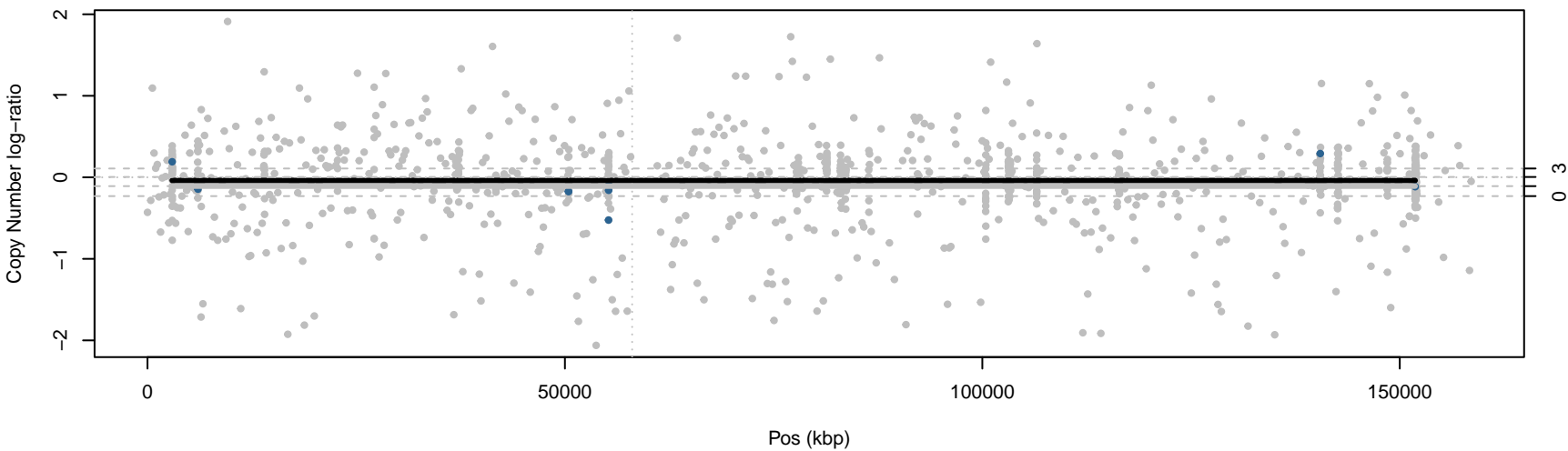
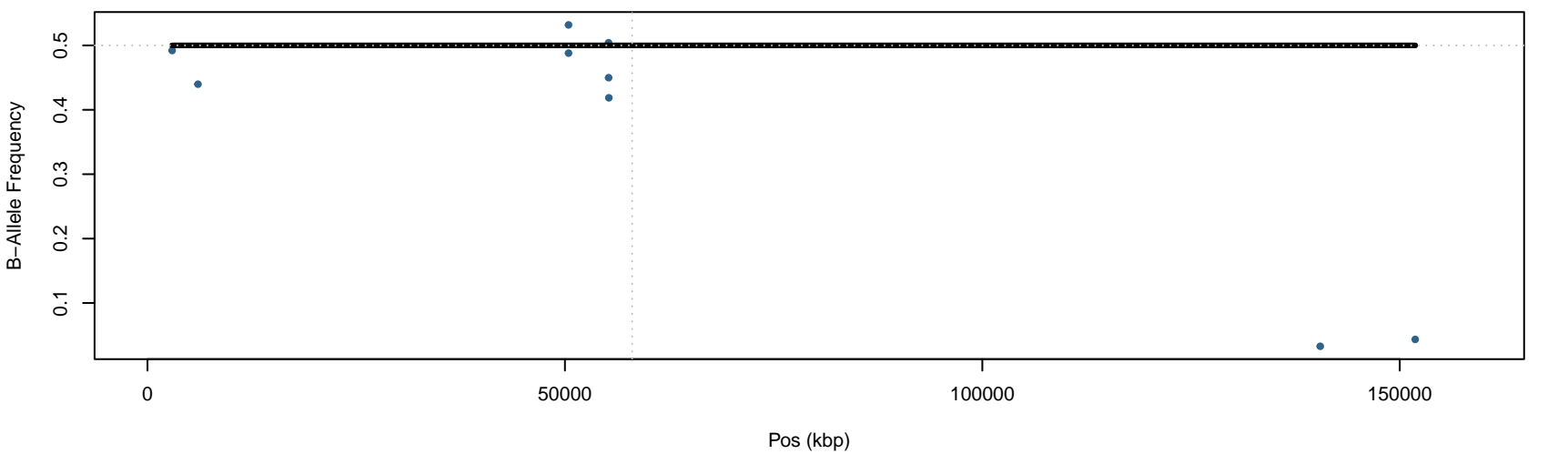
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 5



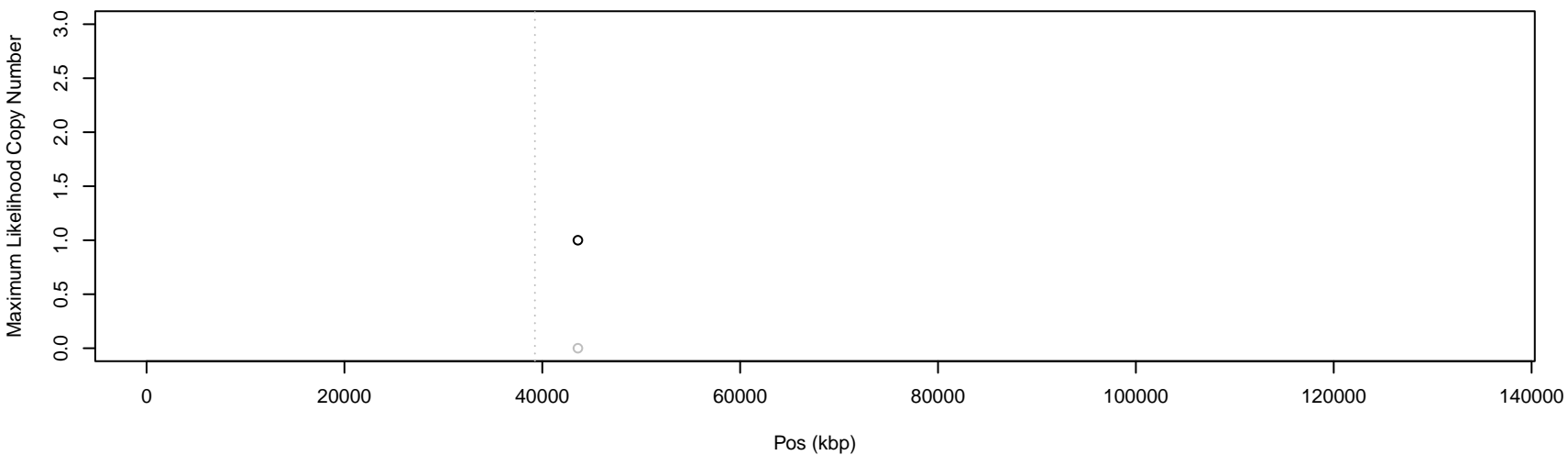
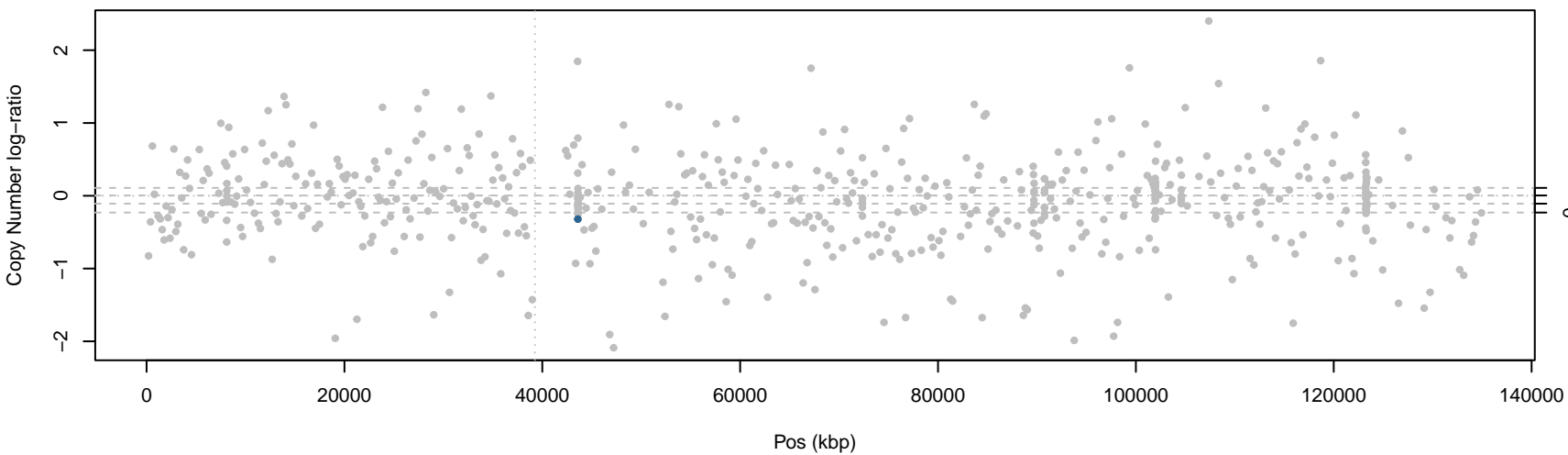
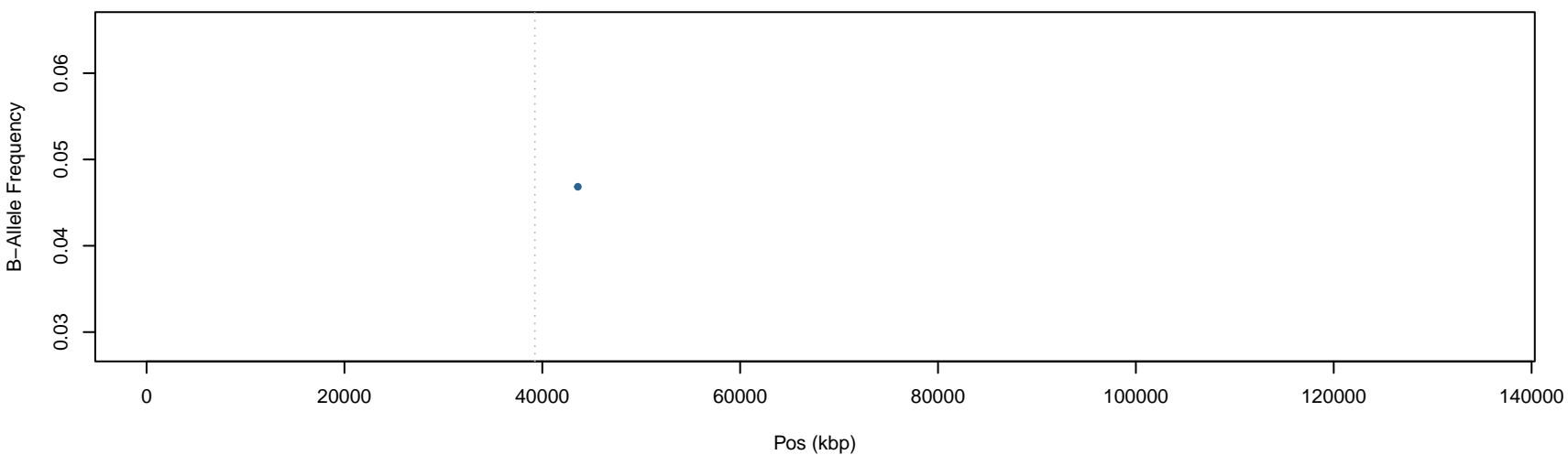
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 6



Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 7

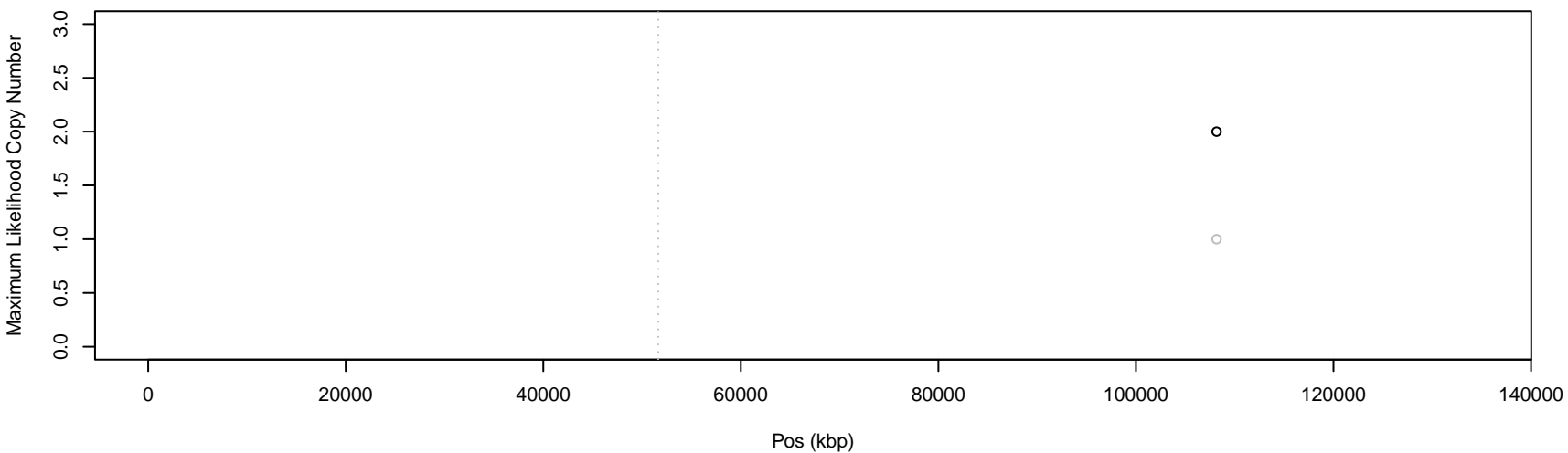
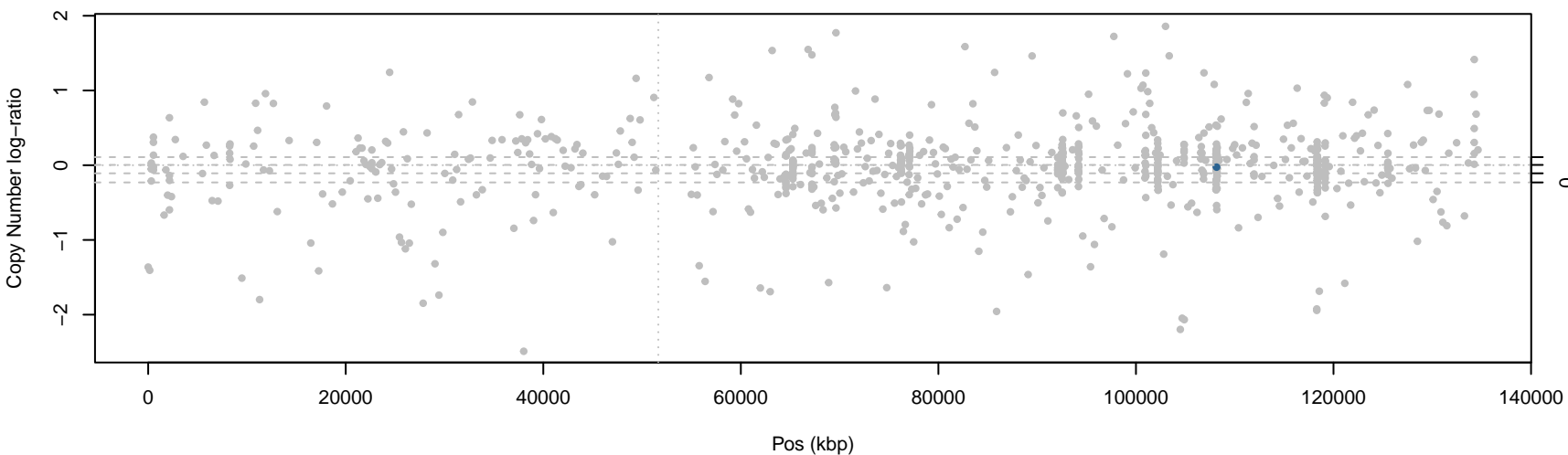
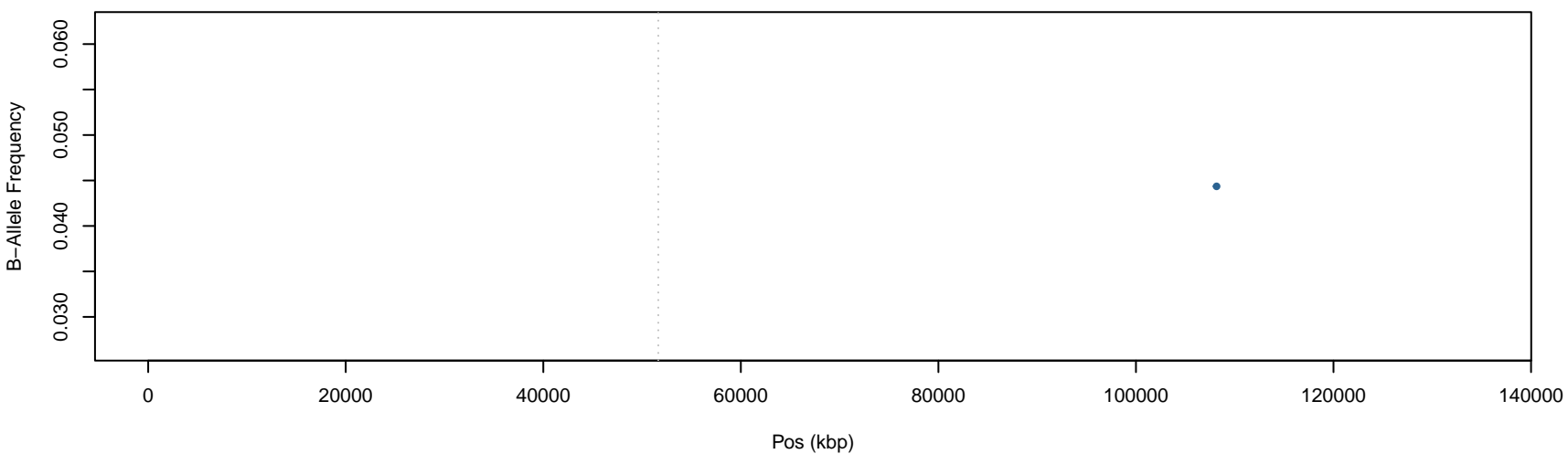


Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 10

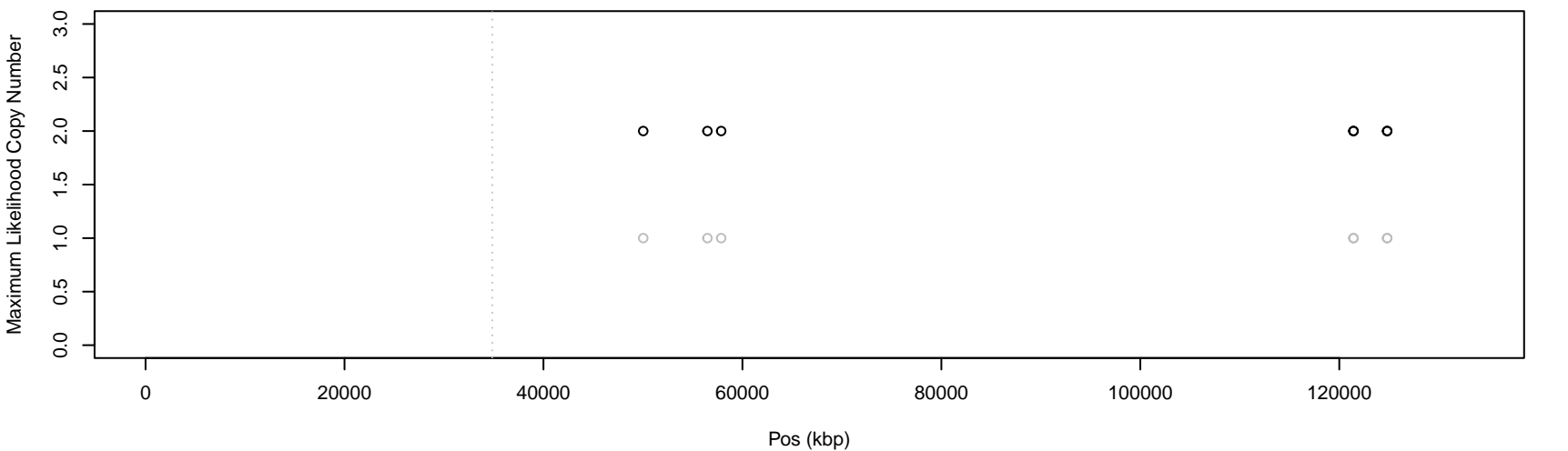
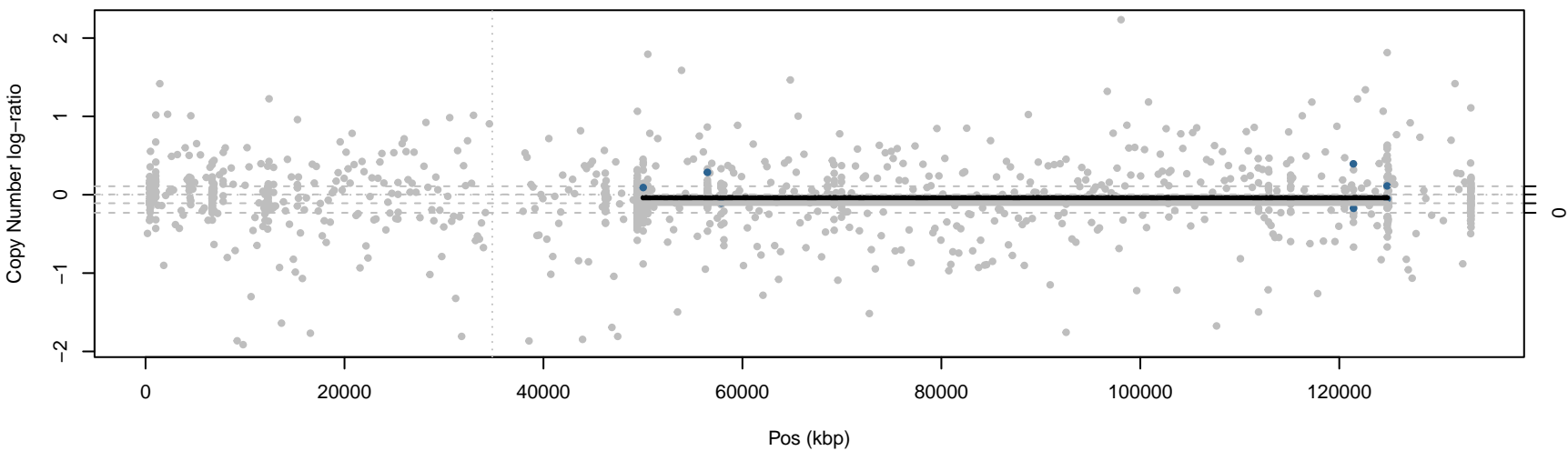
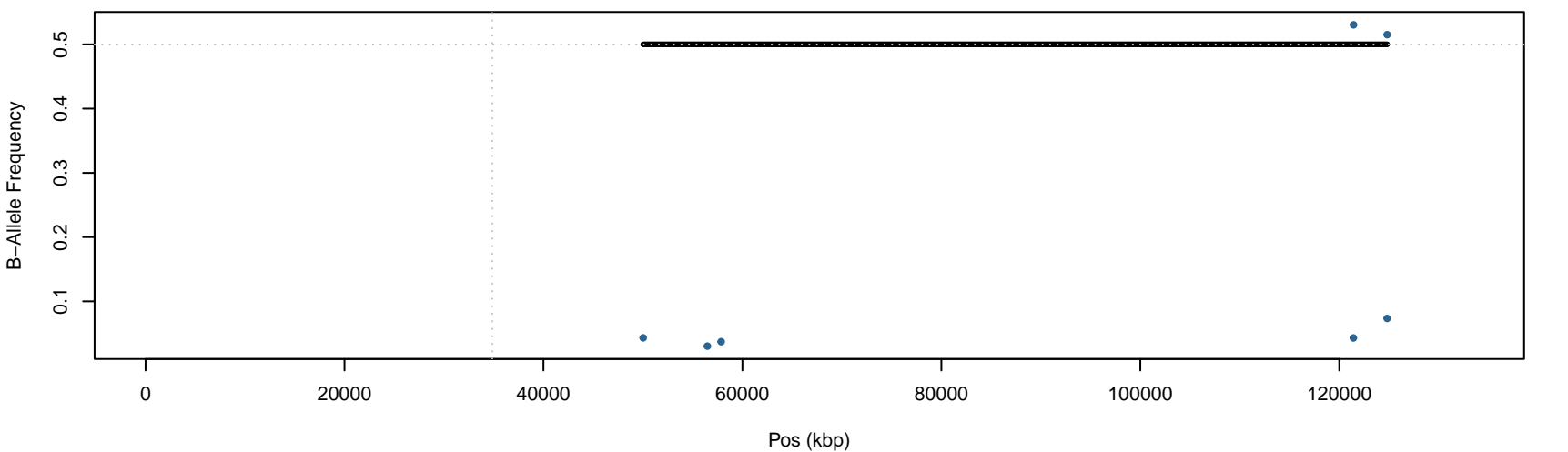




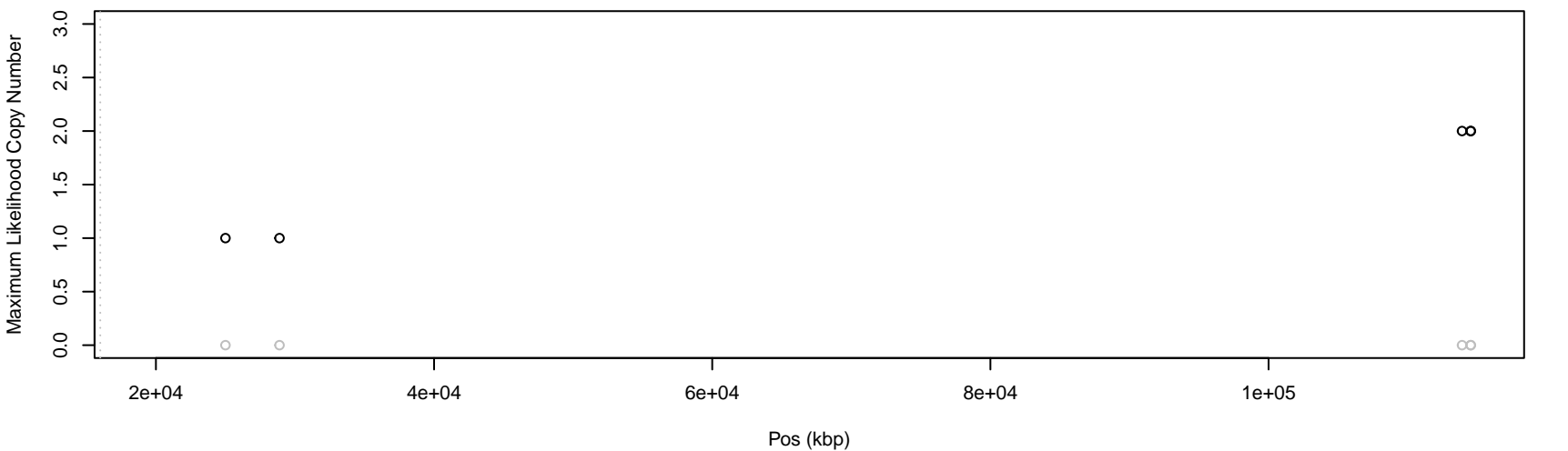
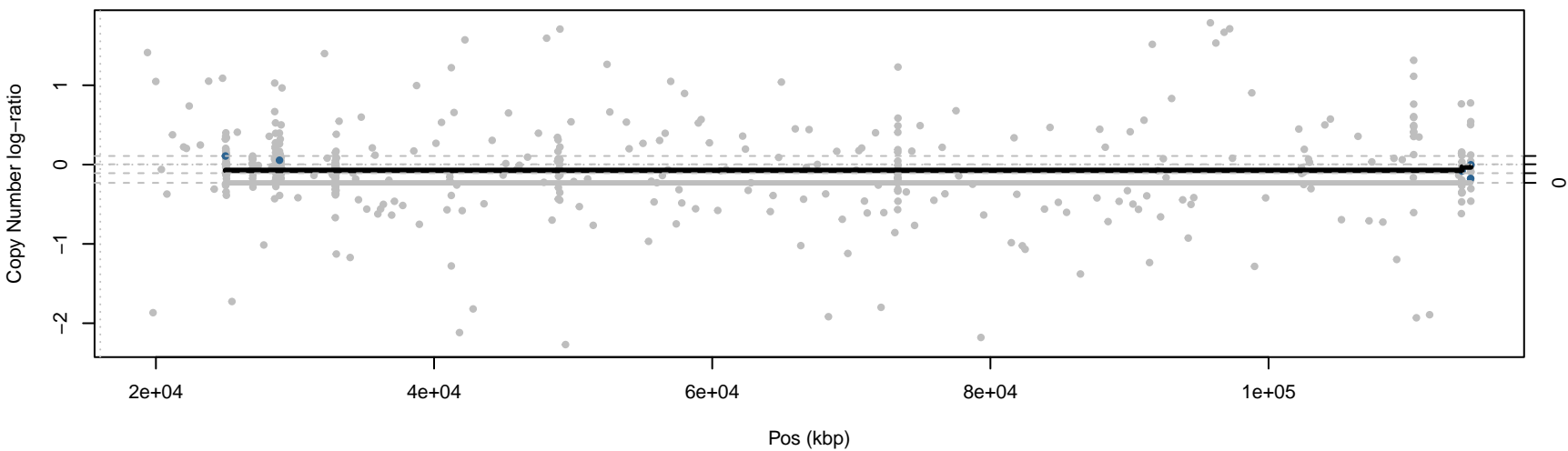
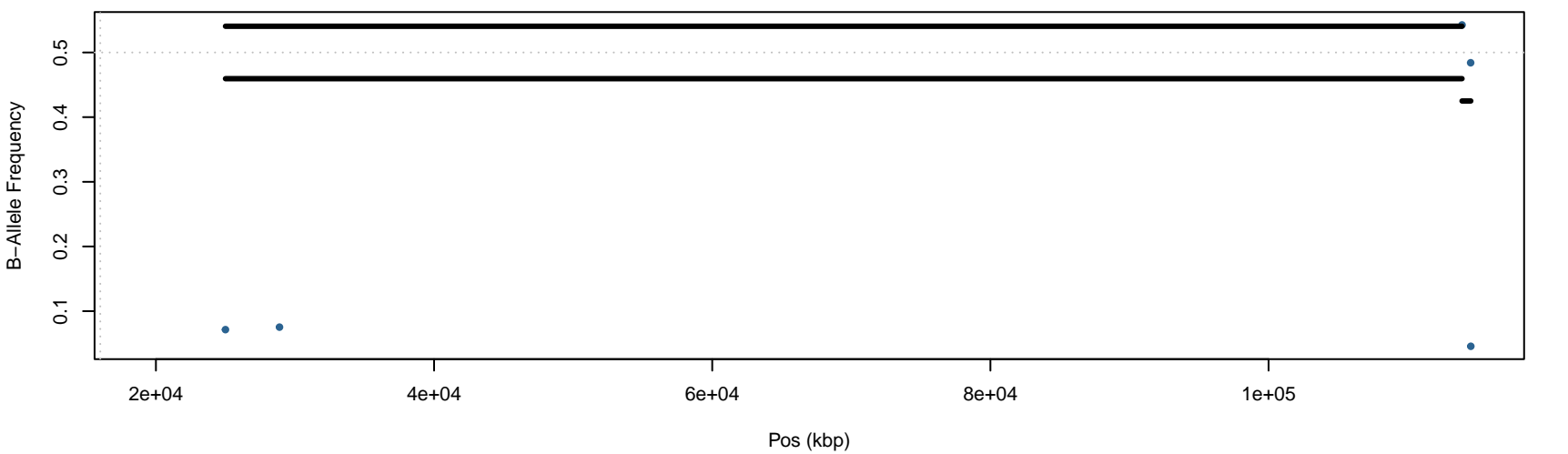
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 11



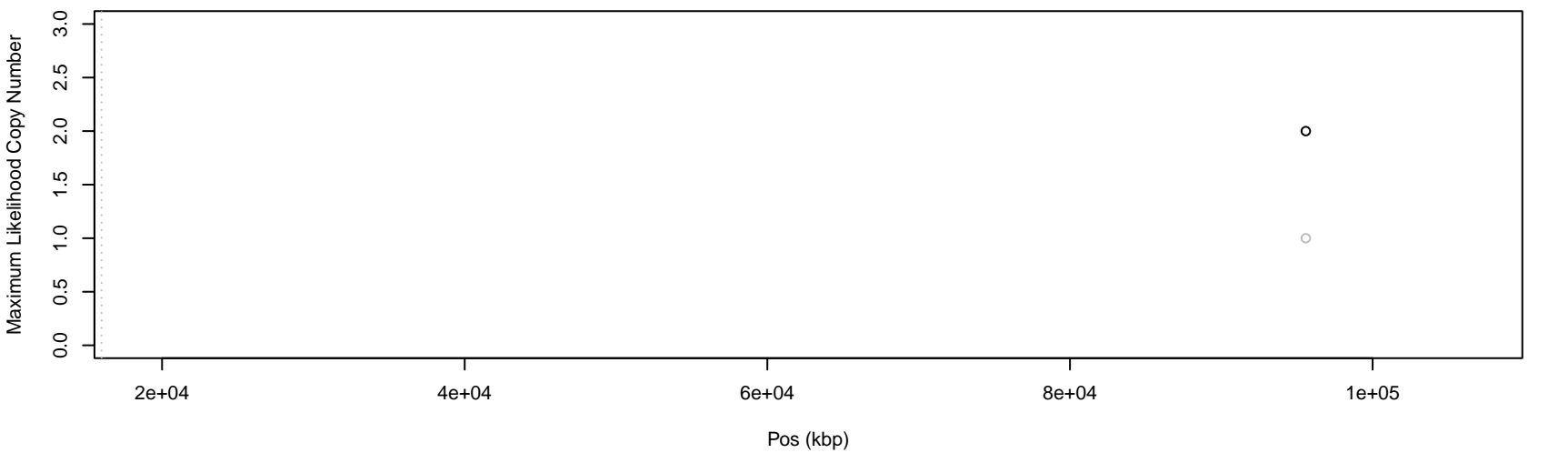
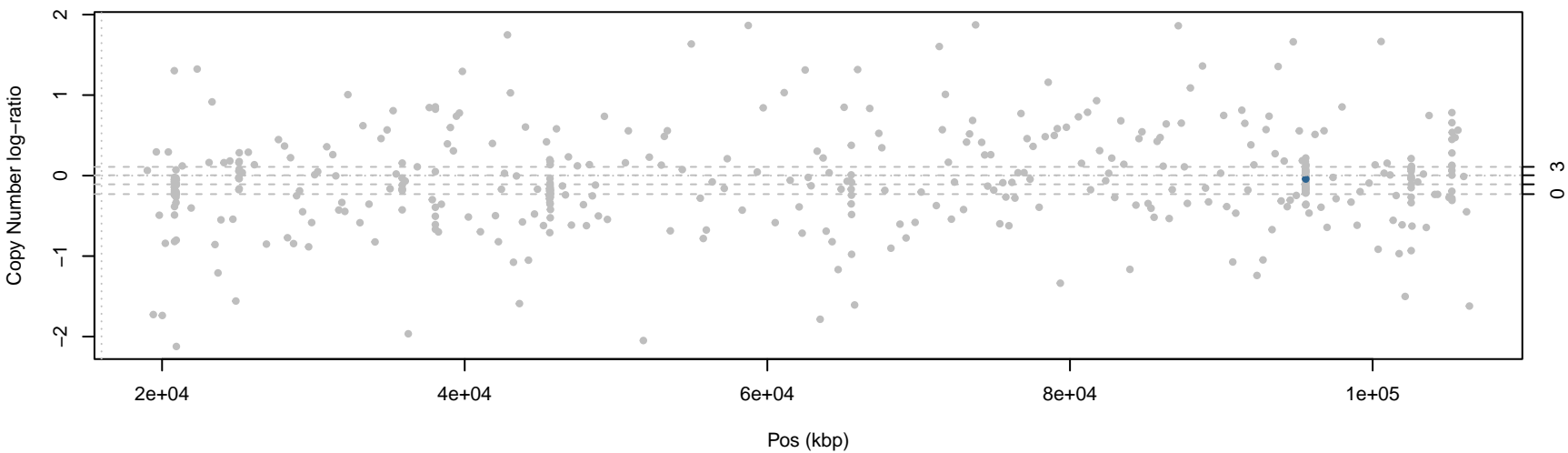
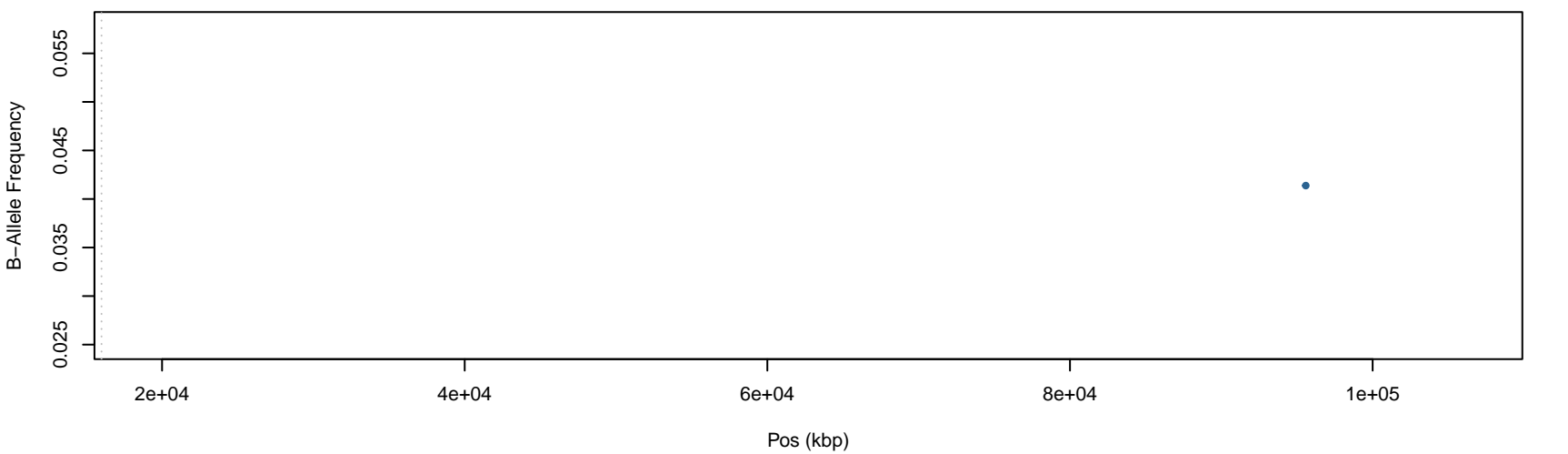
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 12



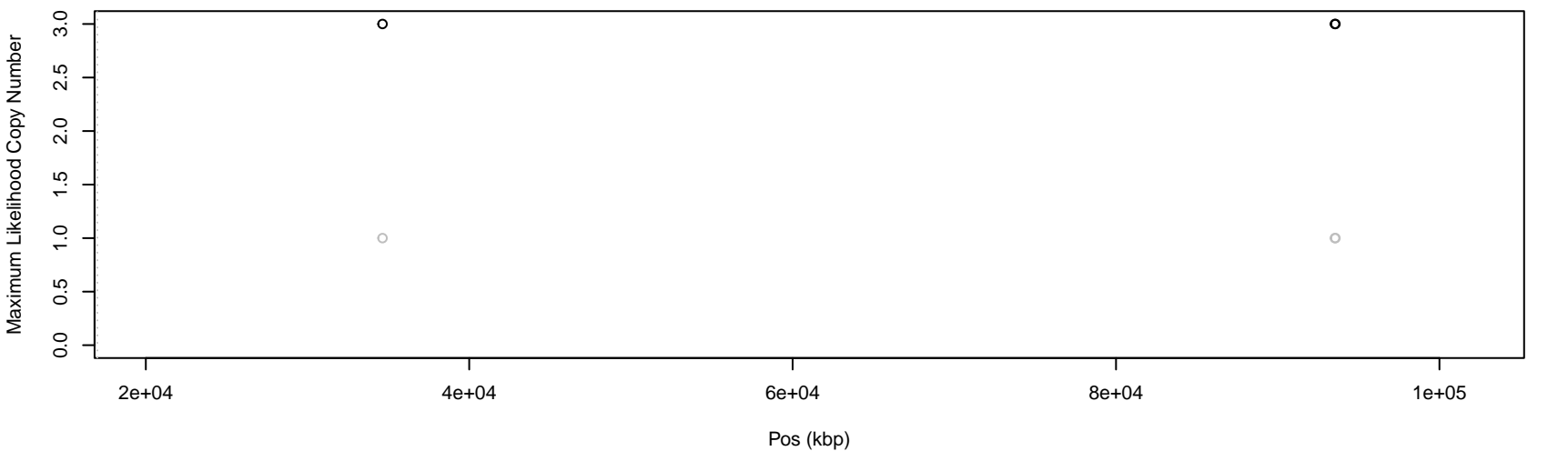
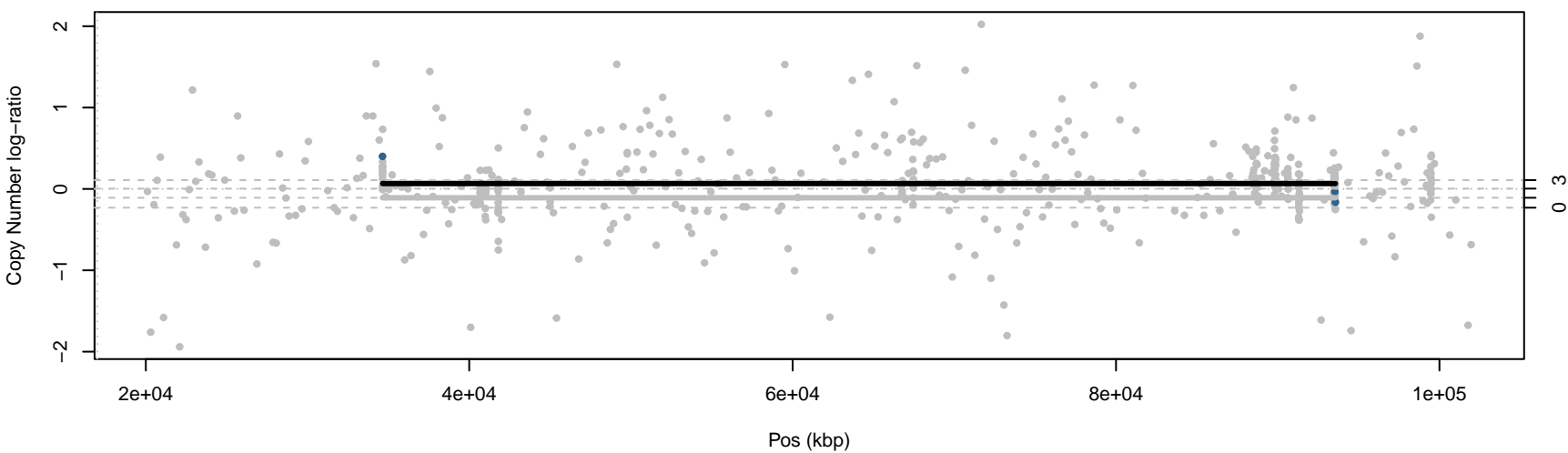
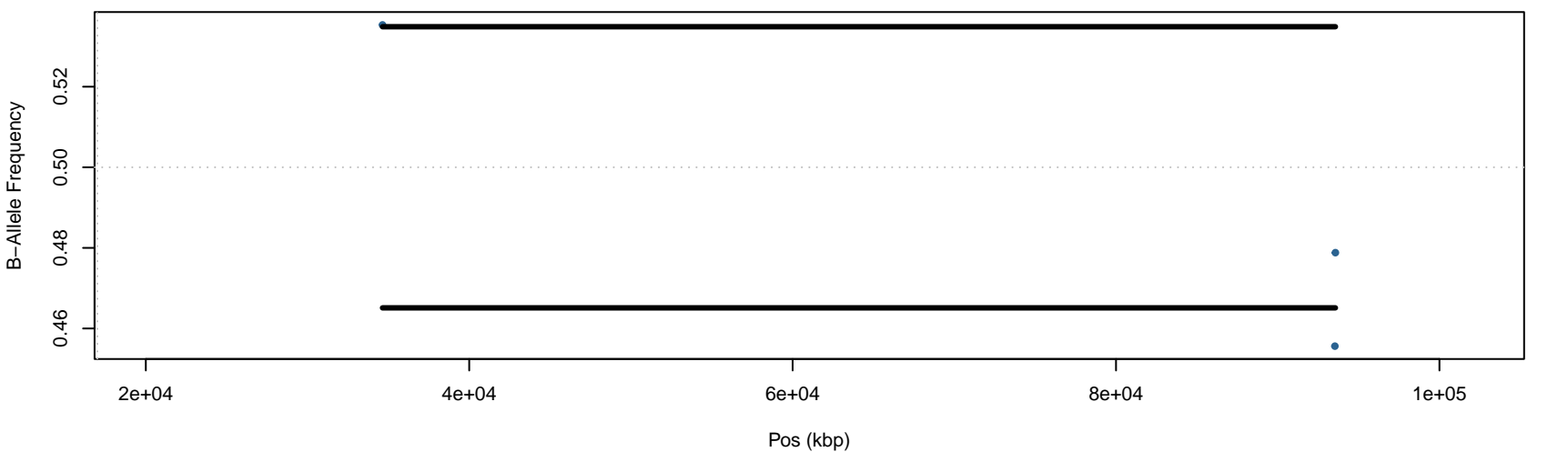
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 13



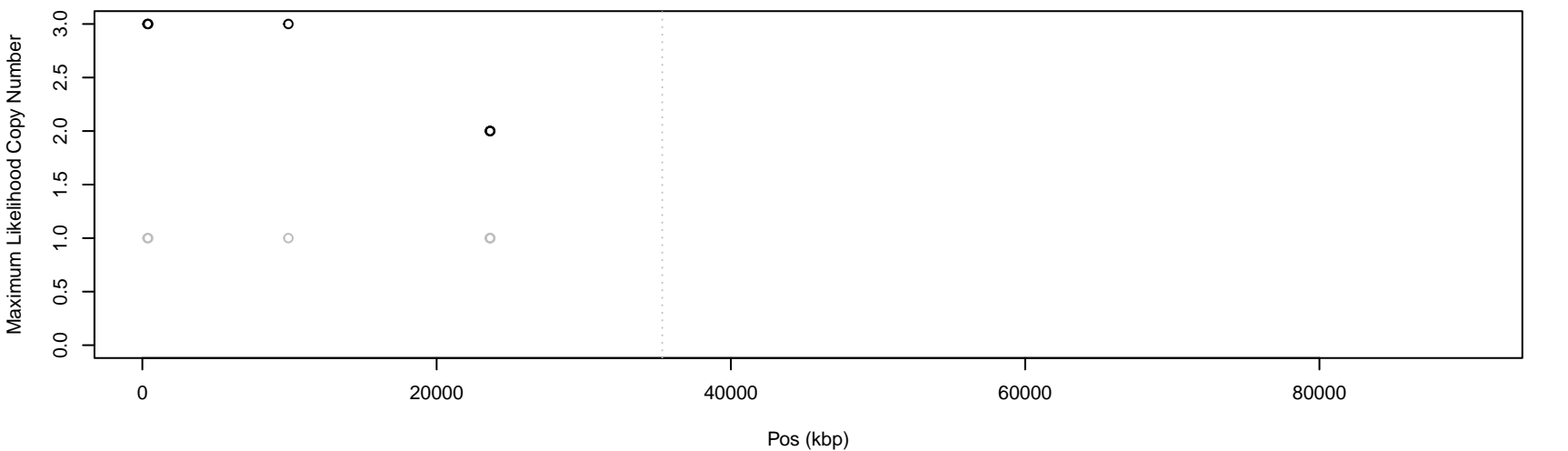
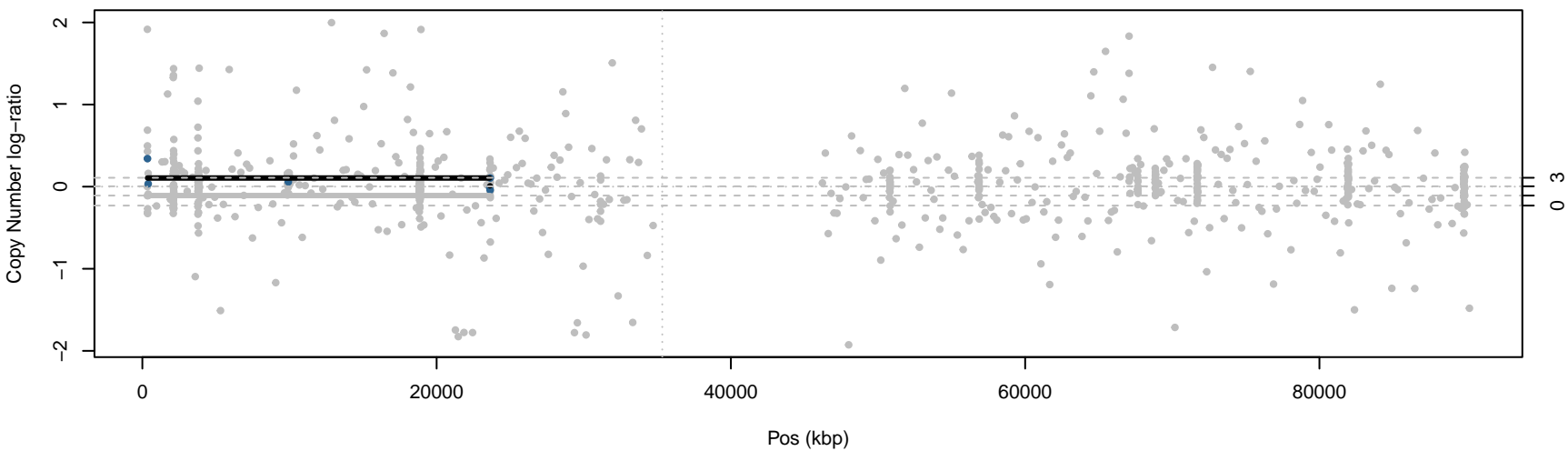
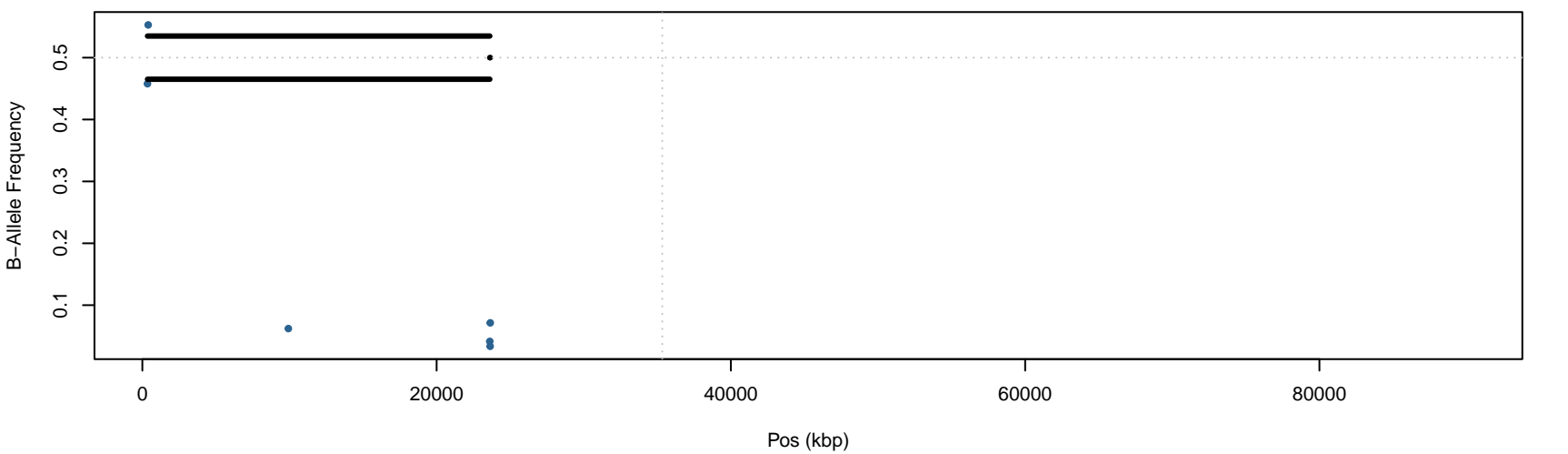
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 14



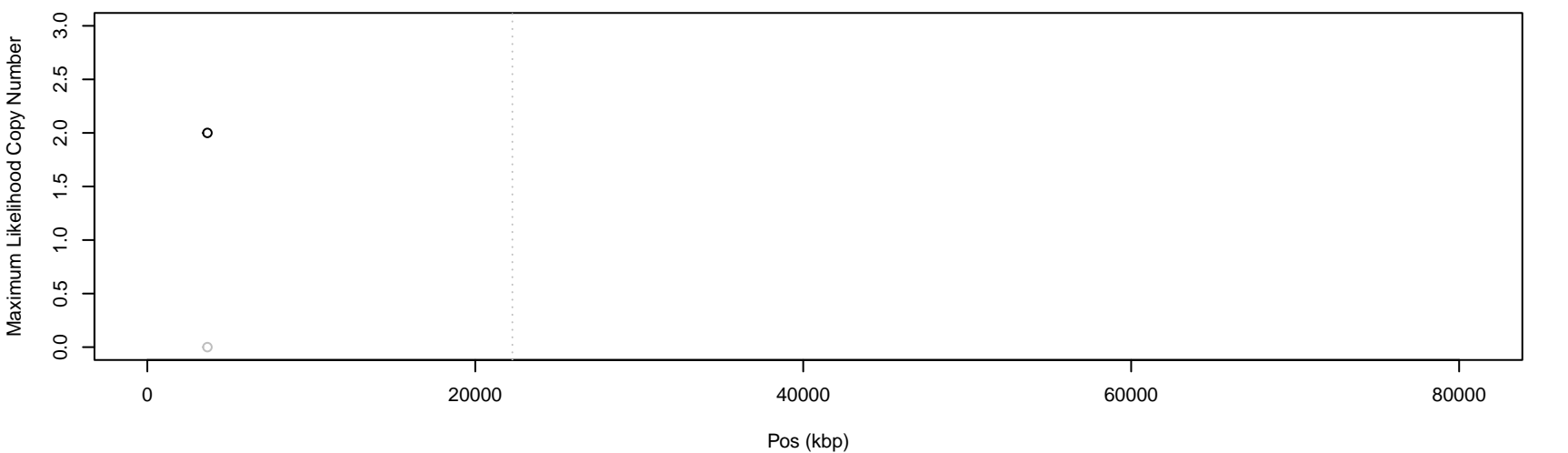
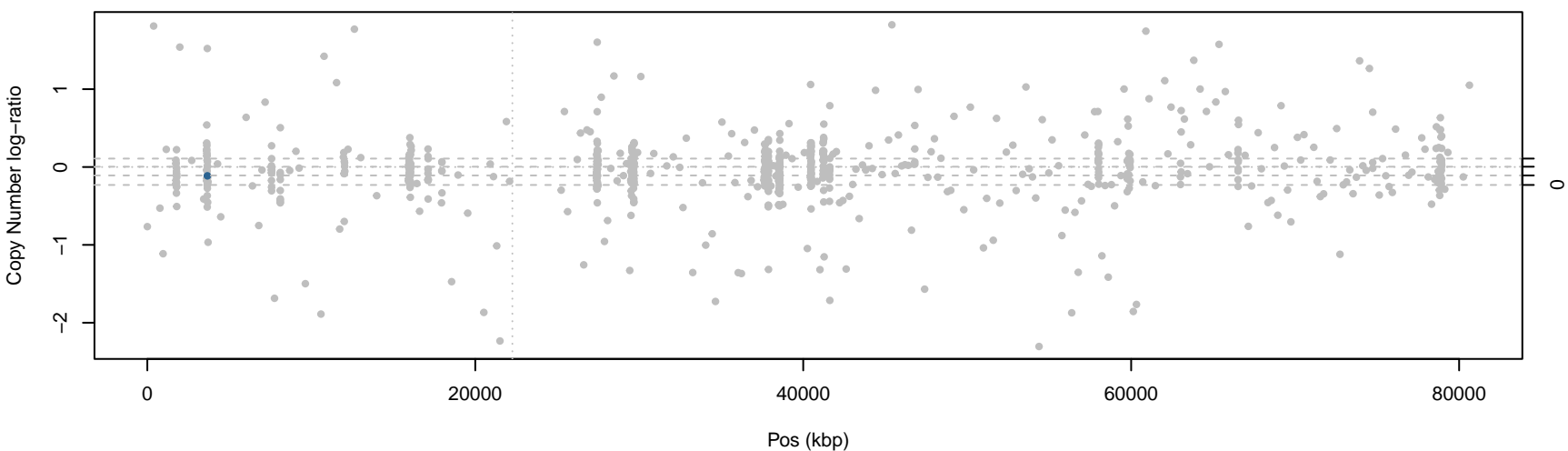
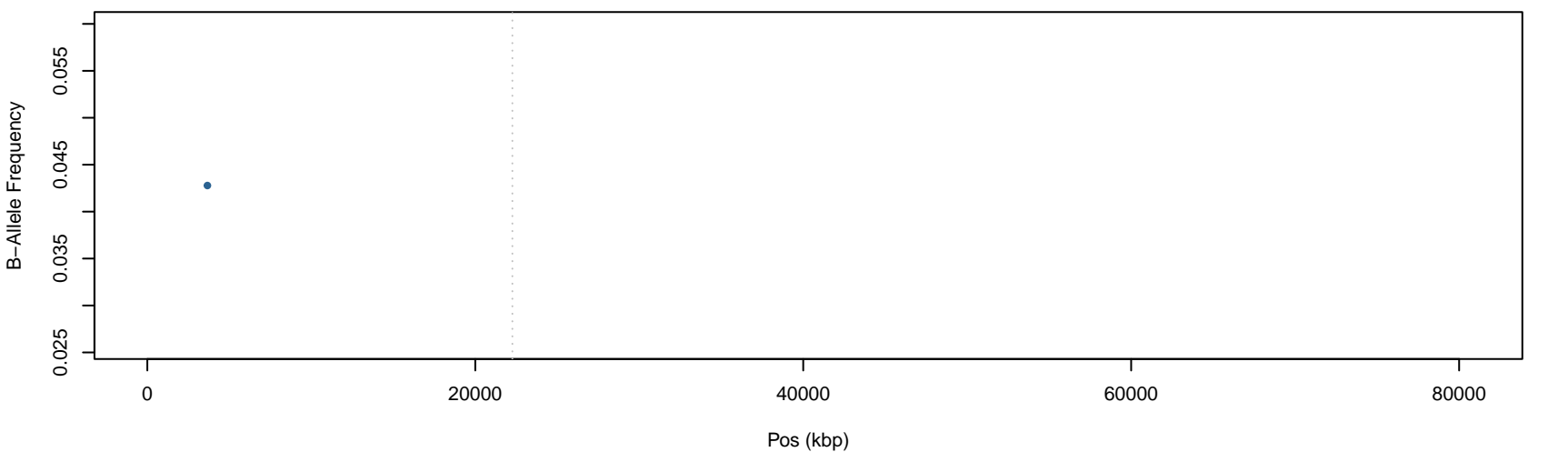
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 15



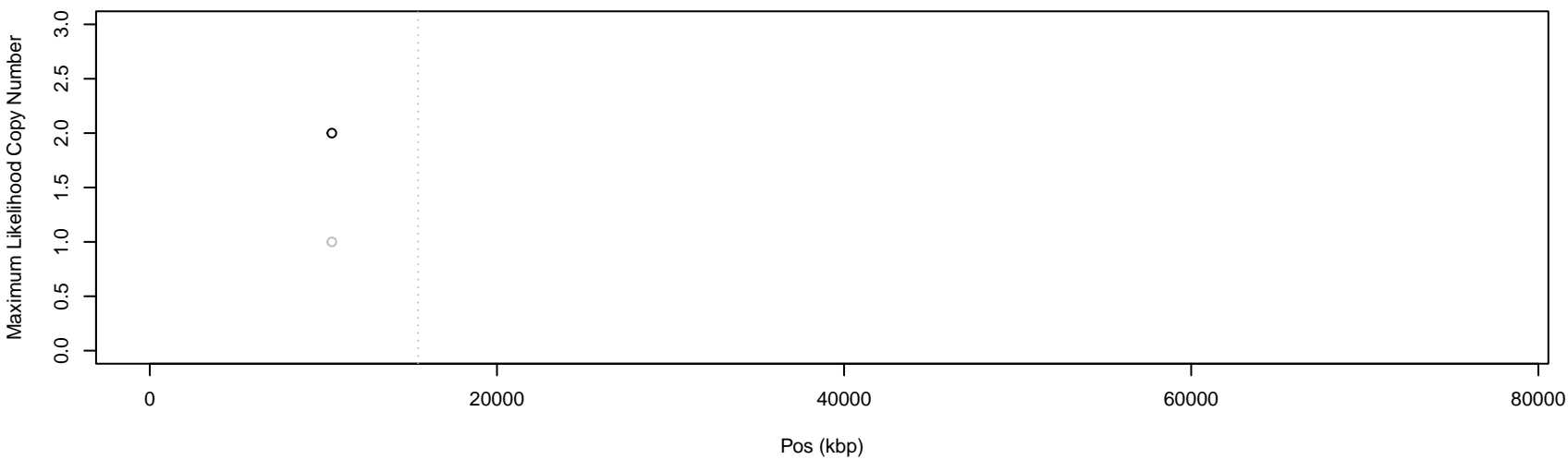
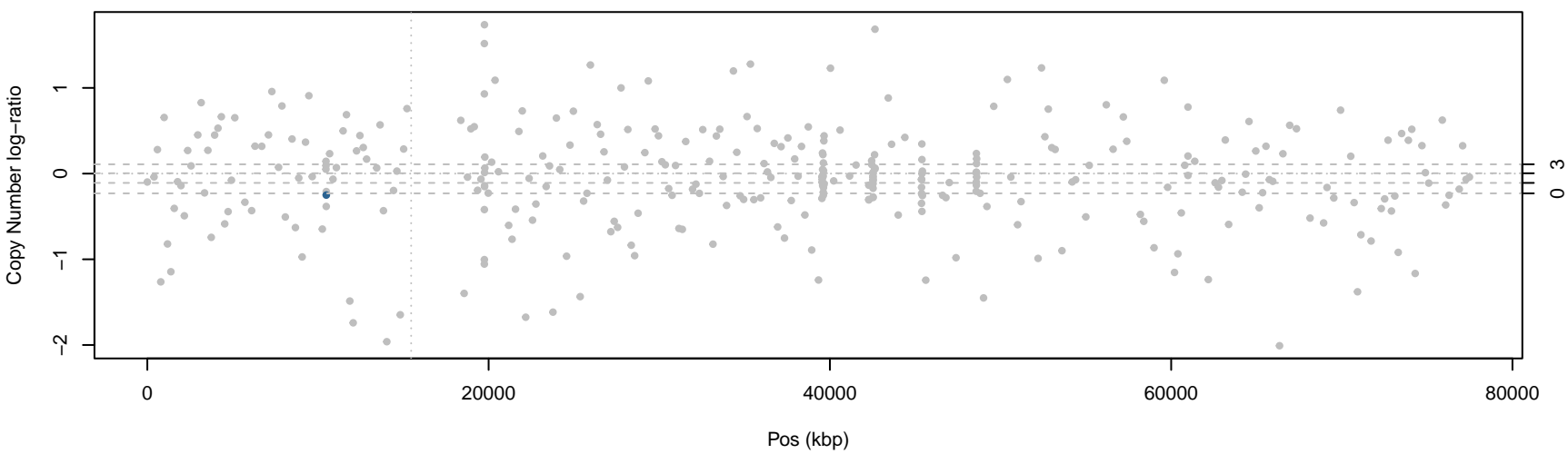
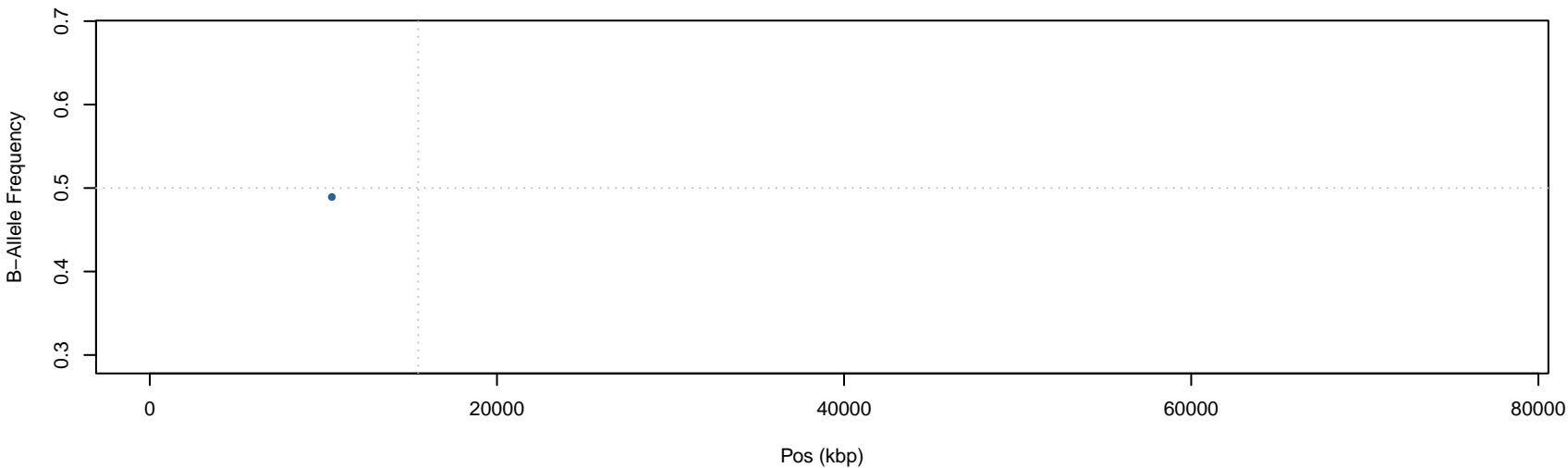
Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 16



Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 17

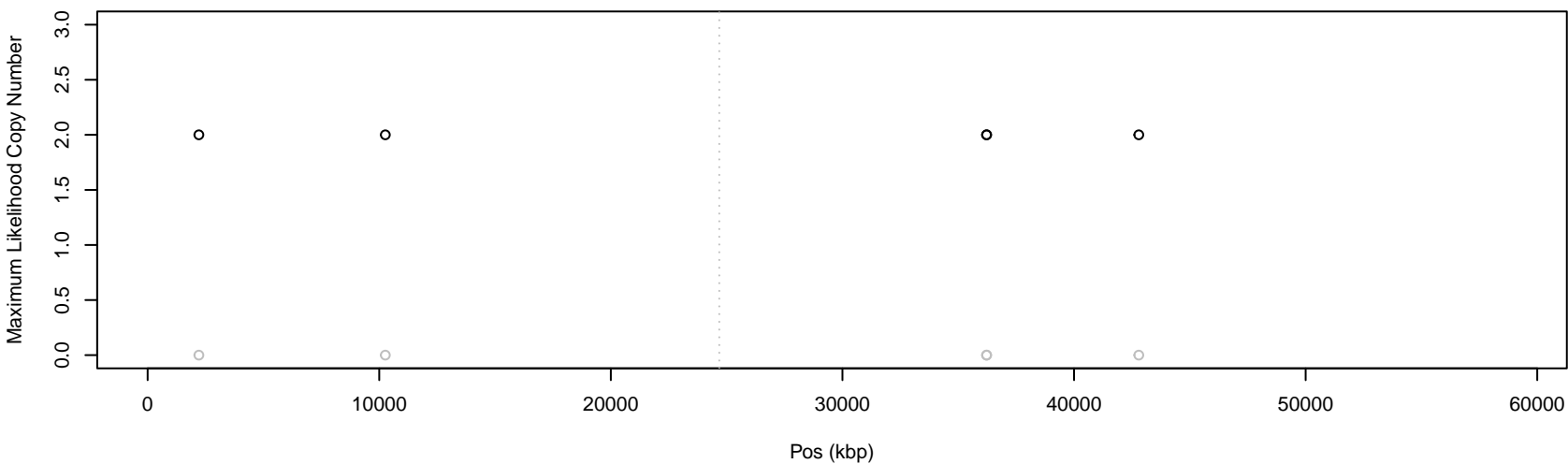
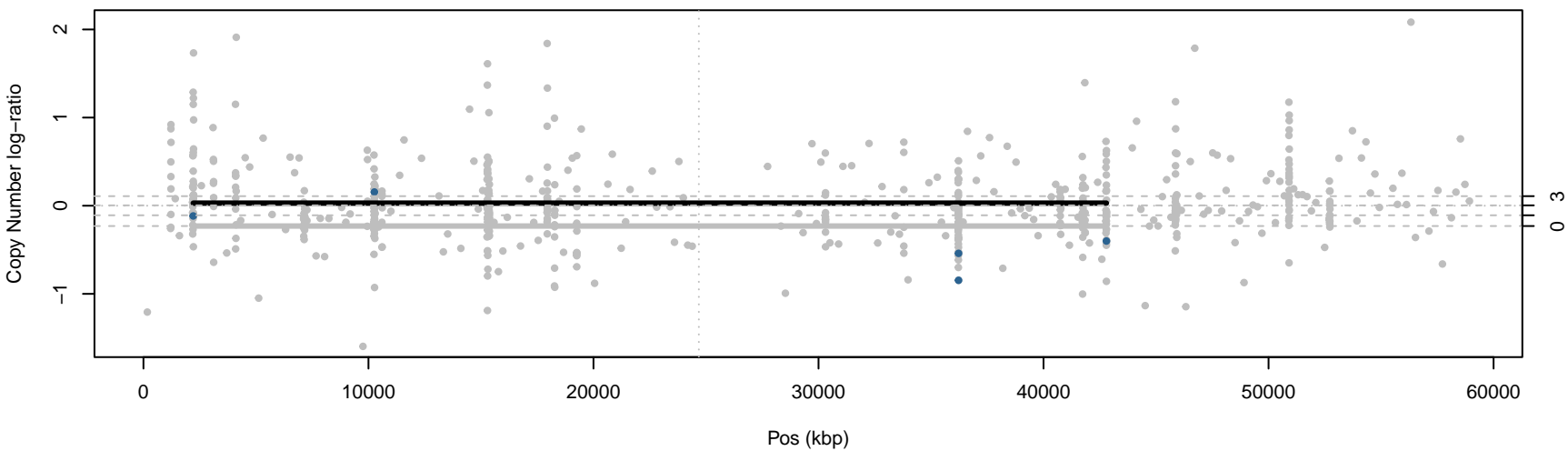
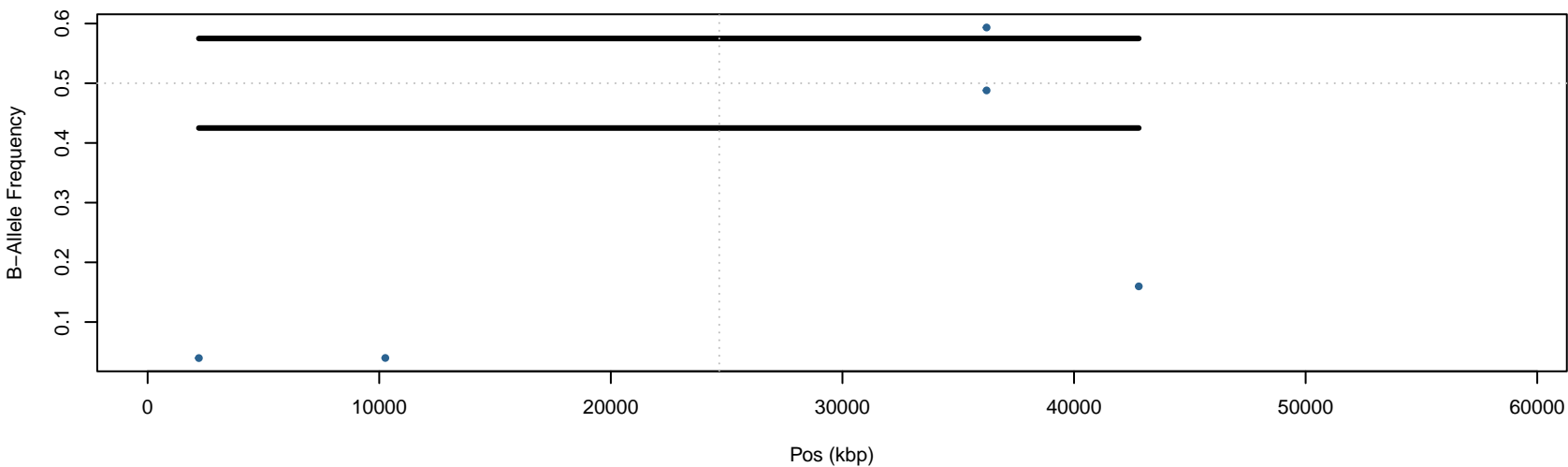


Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 18





Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 19



Purity: 0.15 Tumor ploidy: 1.963 SNV log-likelihood: 86.56 GoF: 95.8% Mean coverage: 355;510 Chromosome: 20

