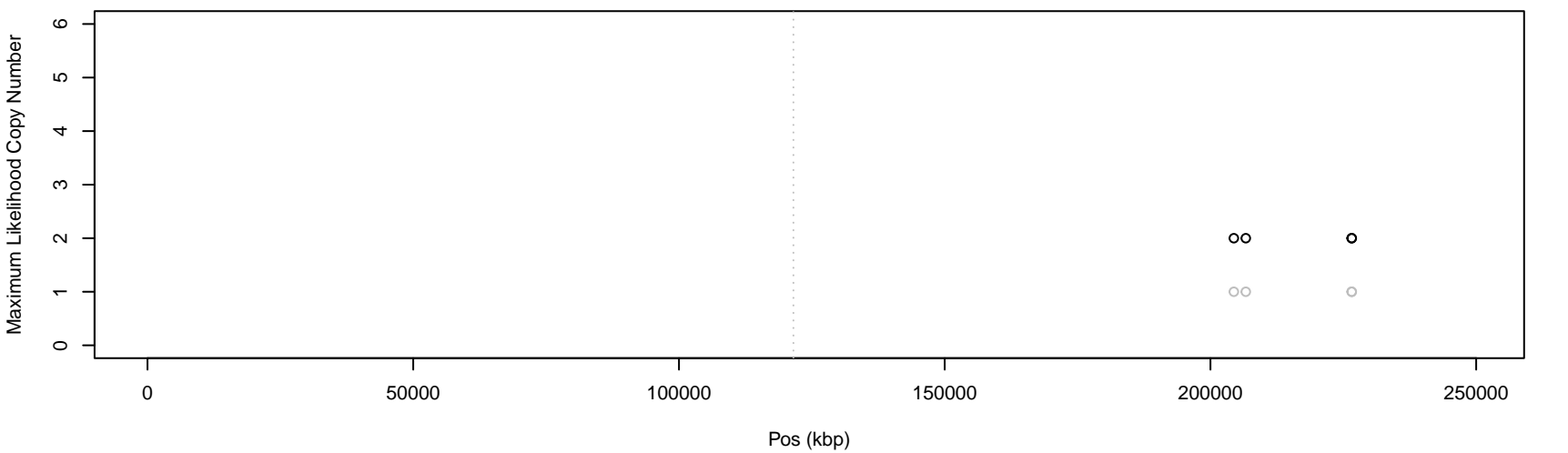
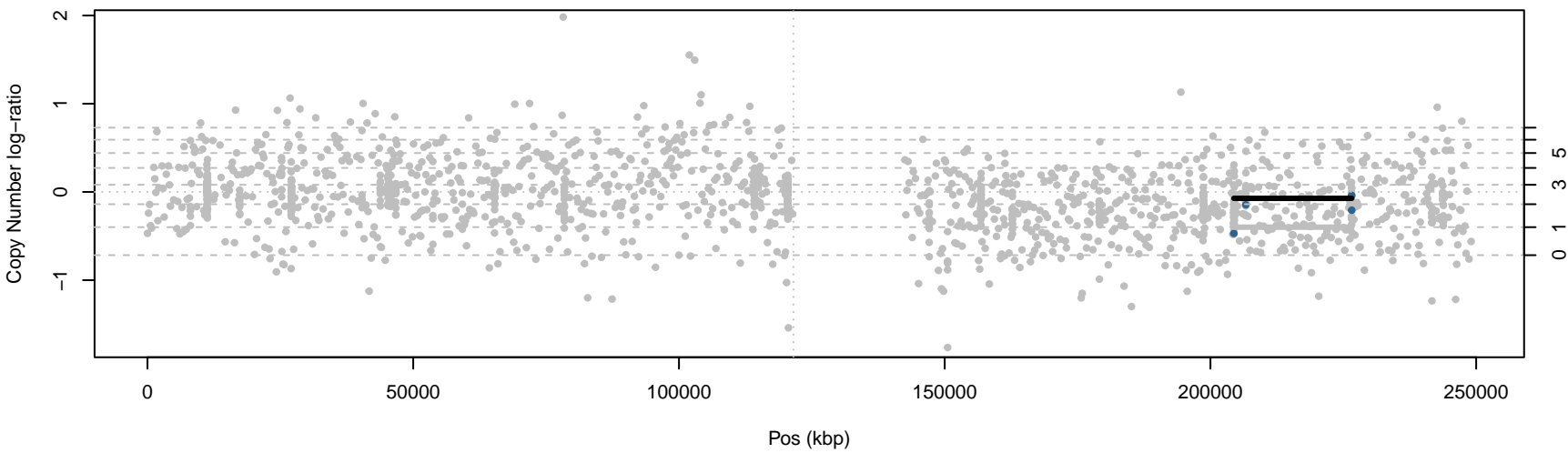
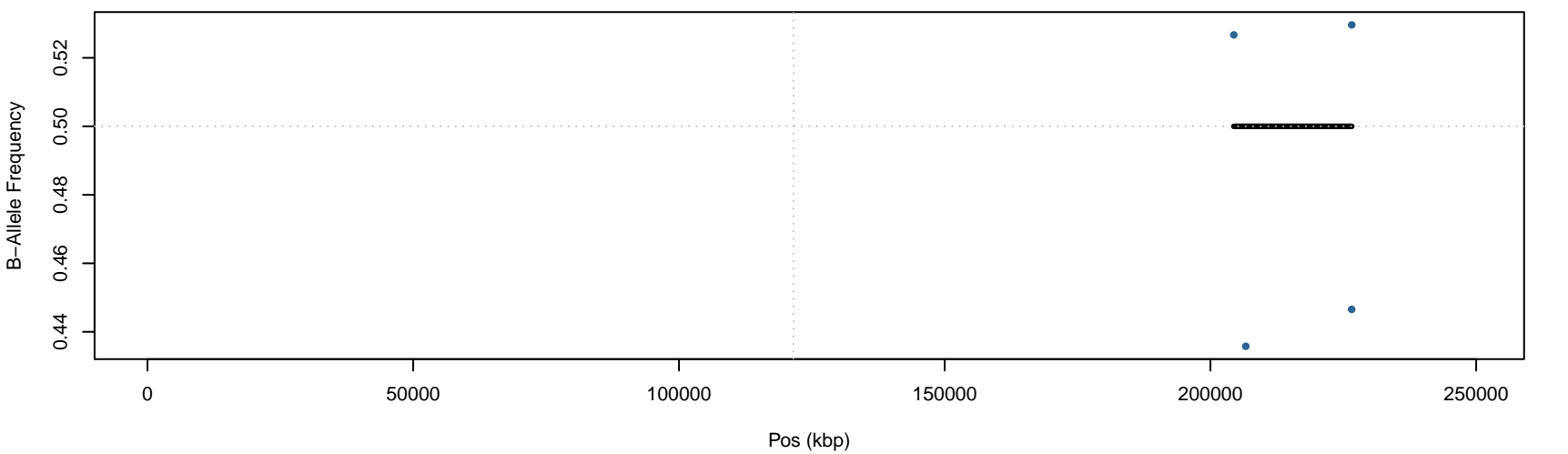
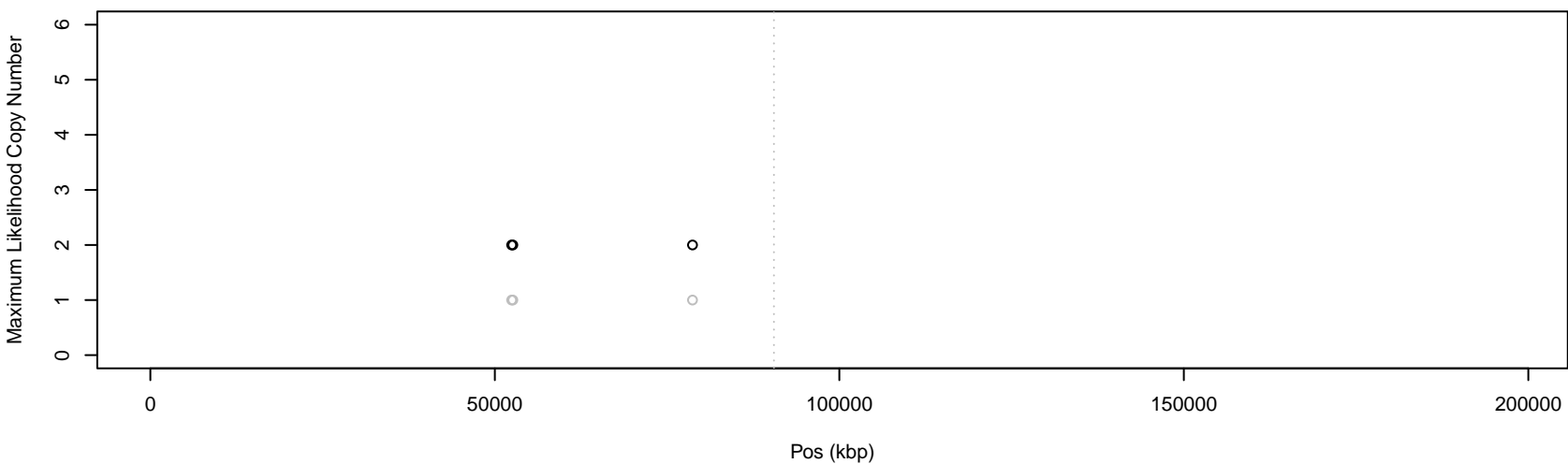
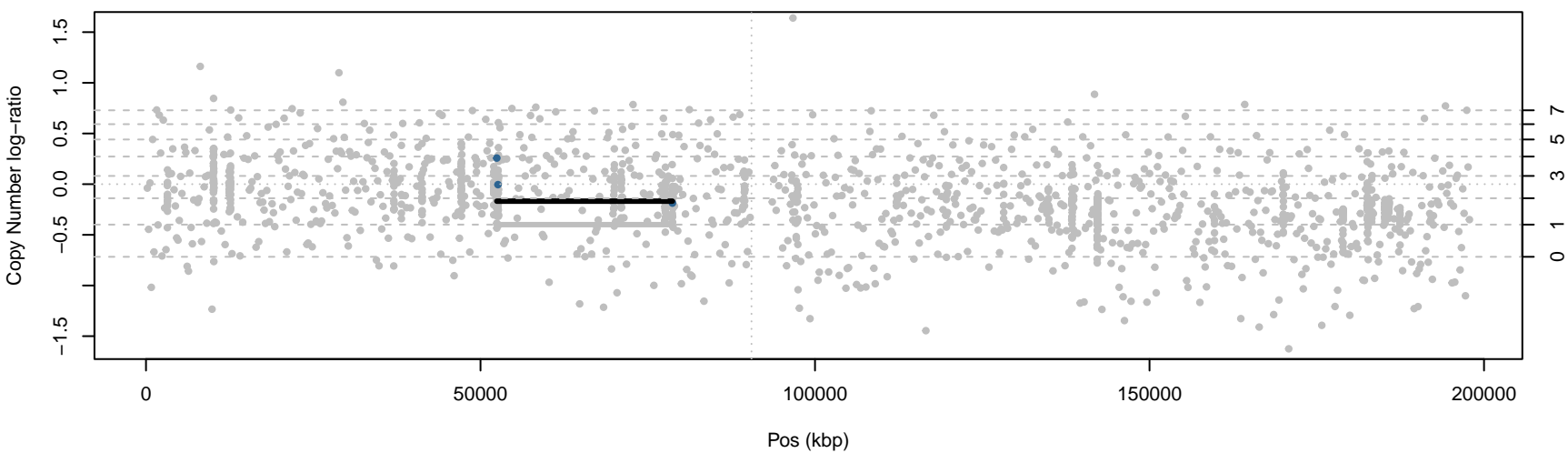
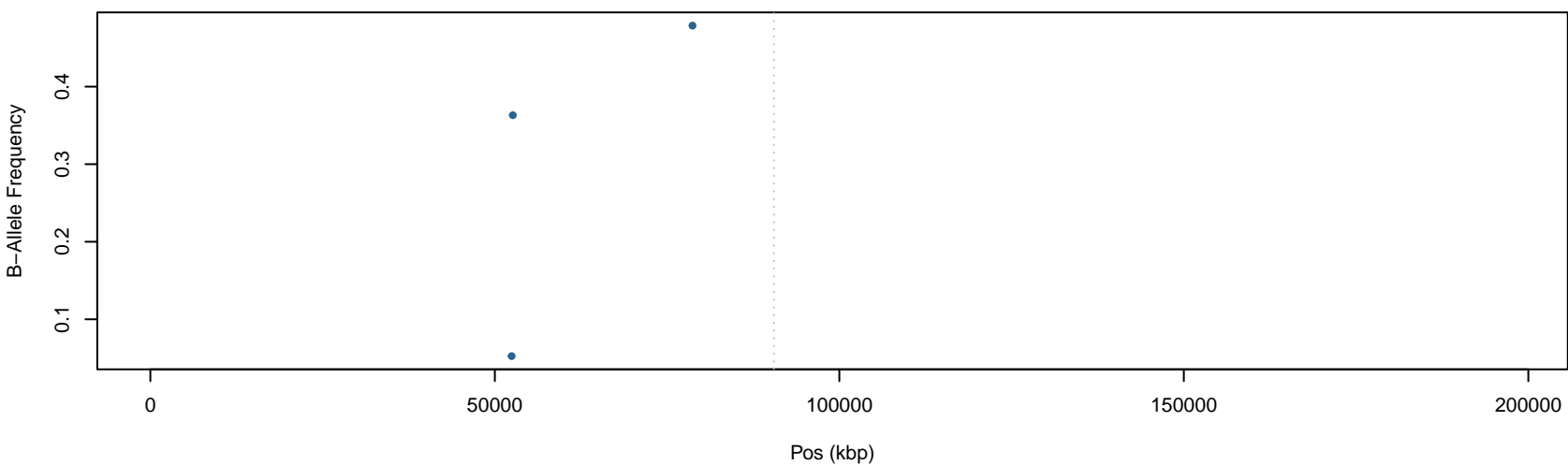


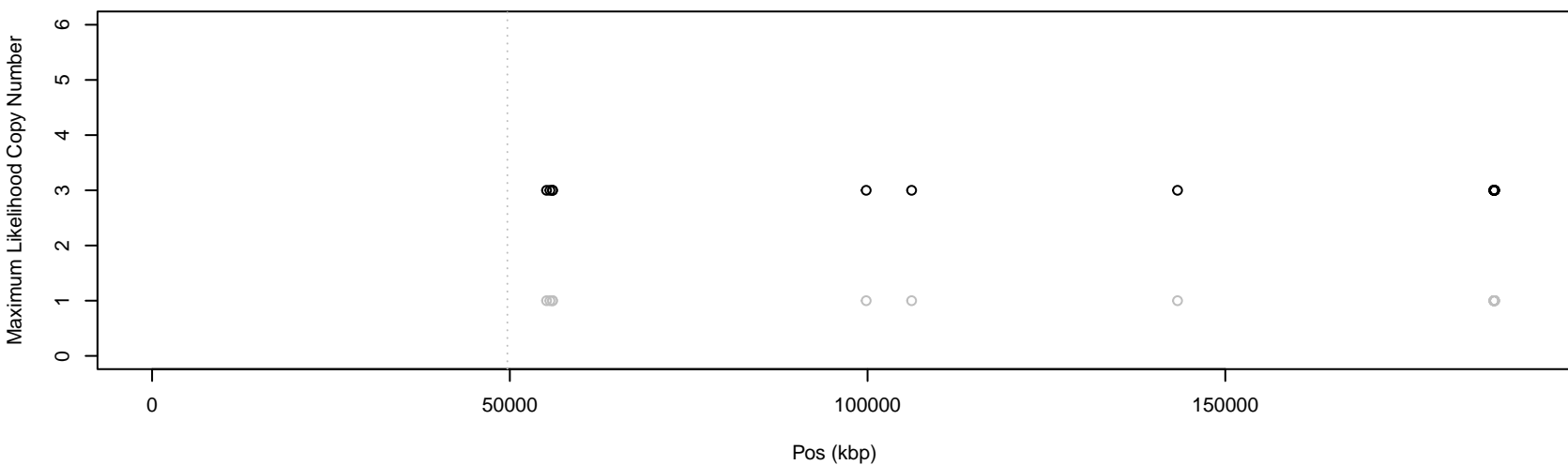
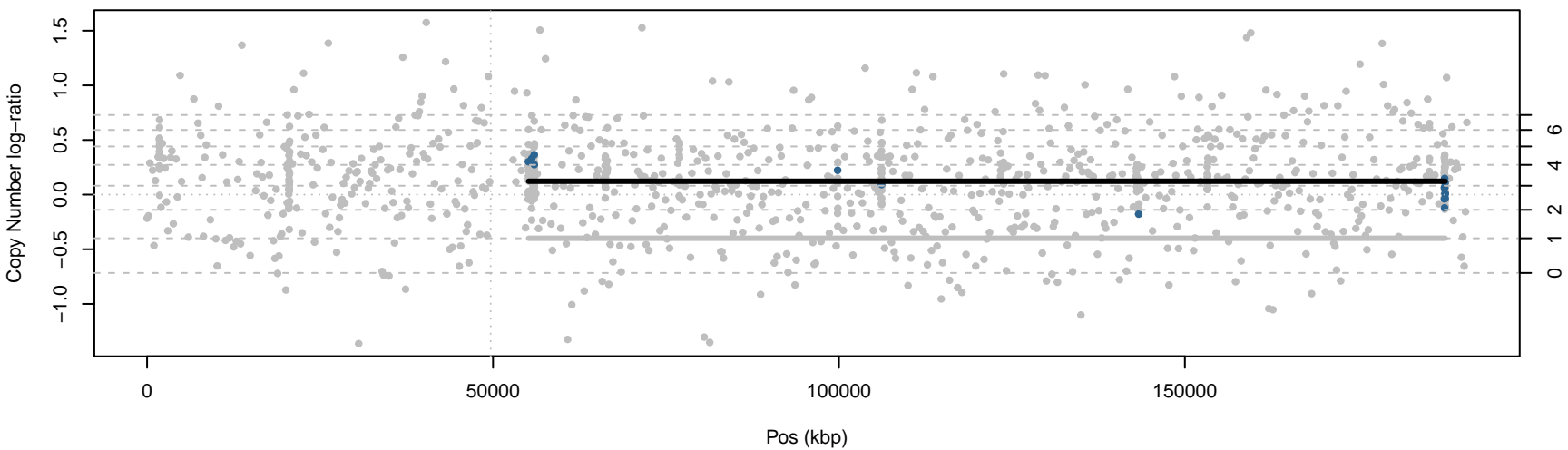
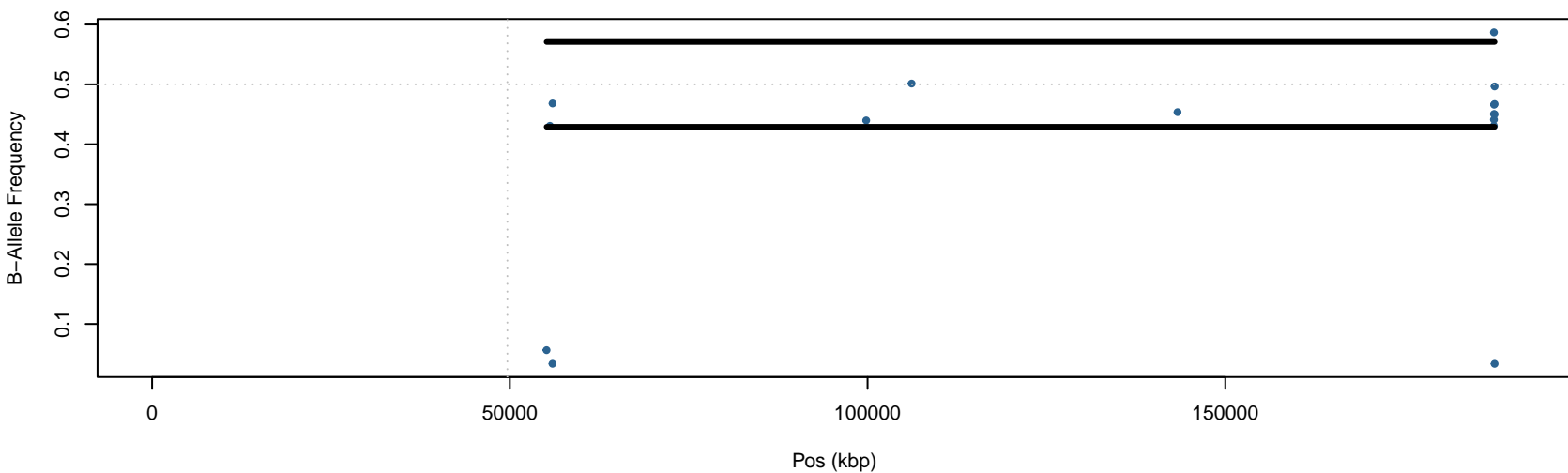
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 1



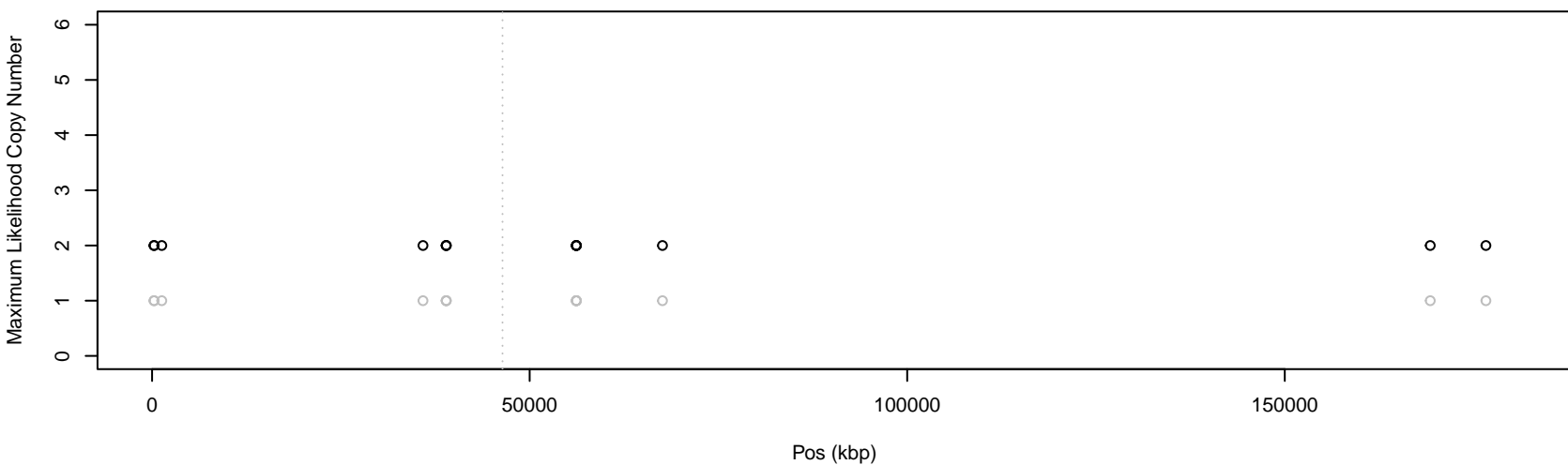
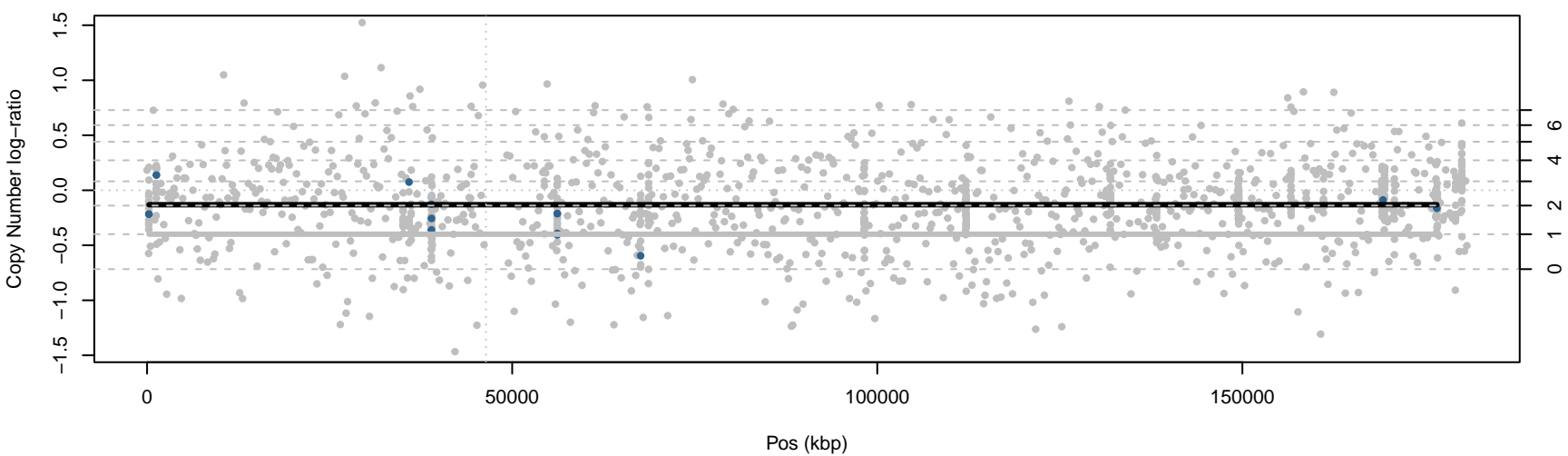
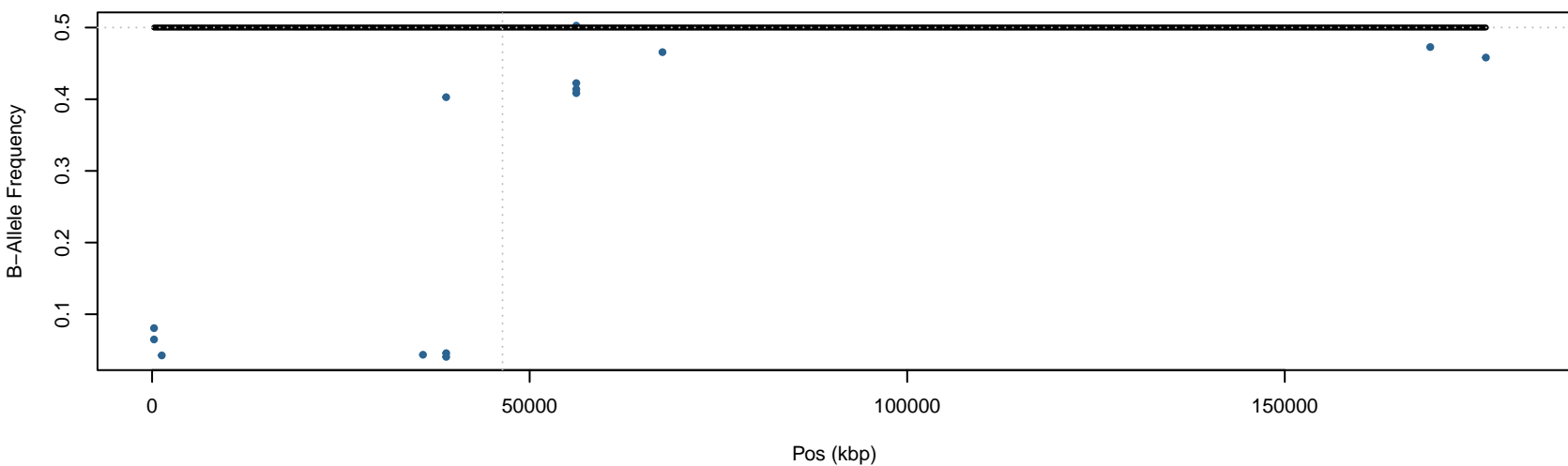
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 3



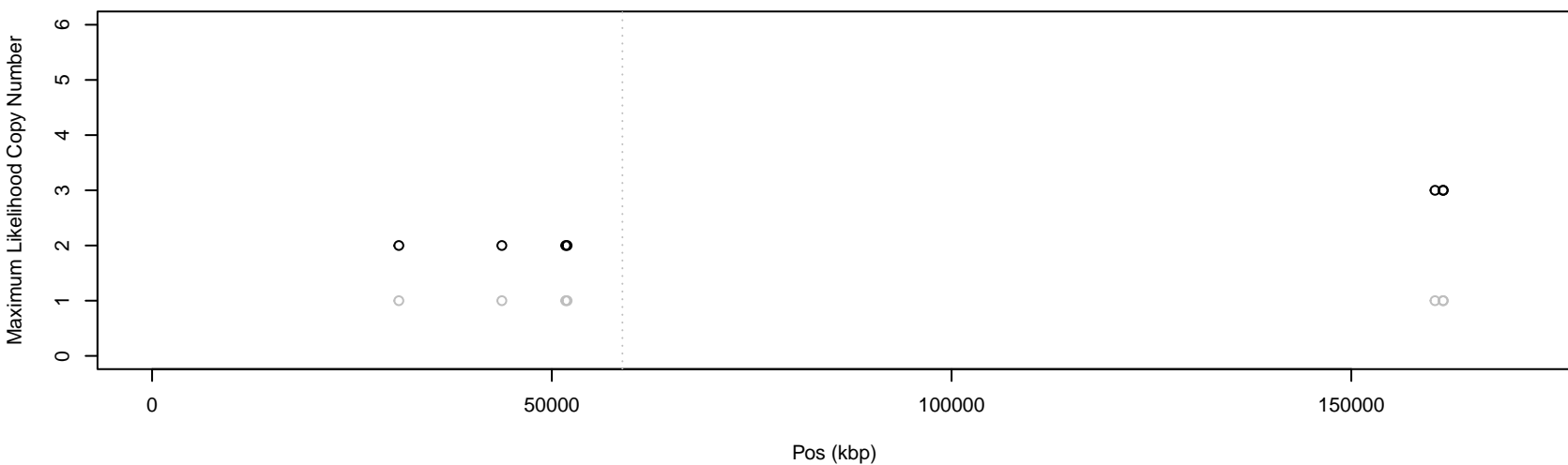
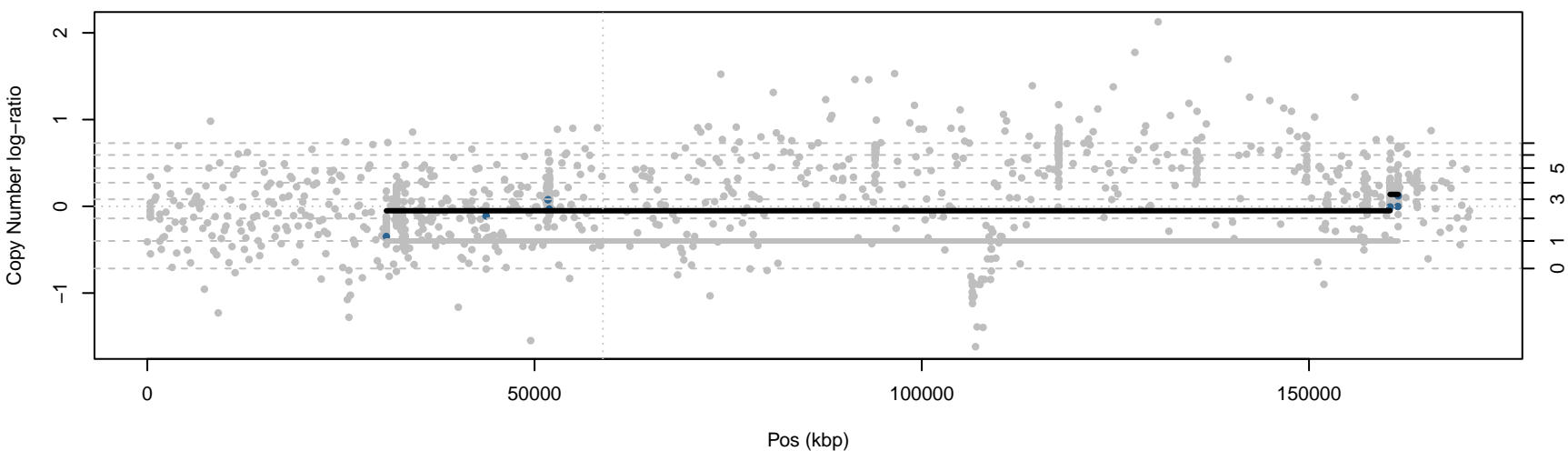
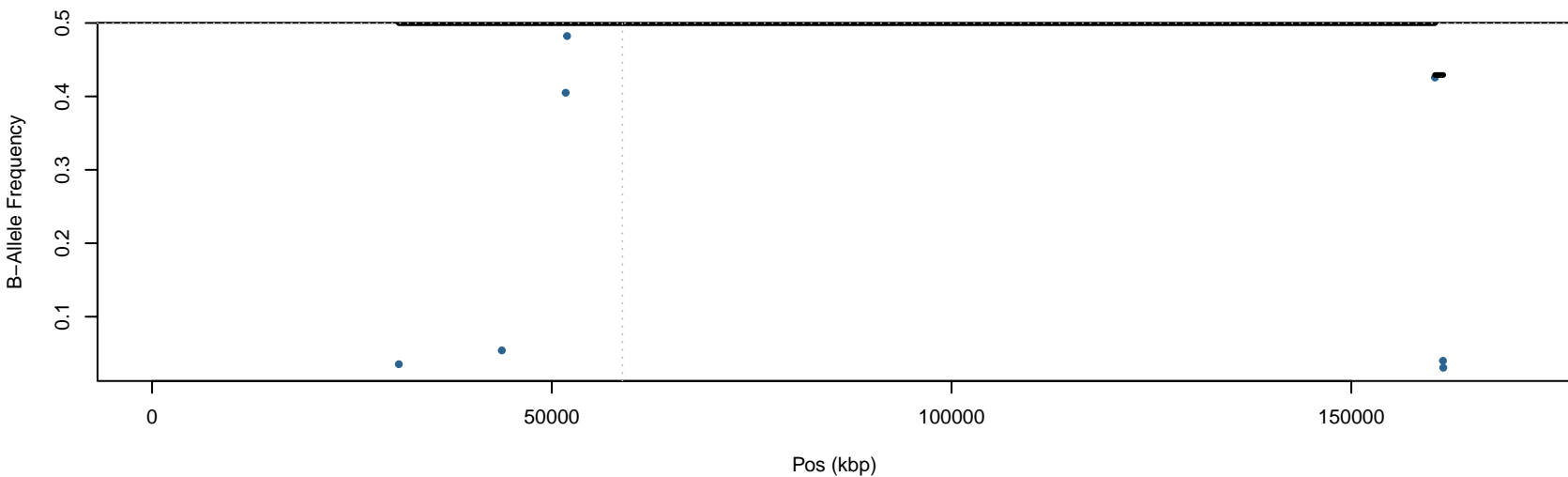
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 4



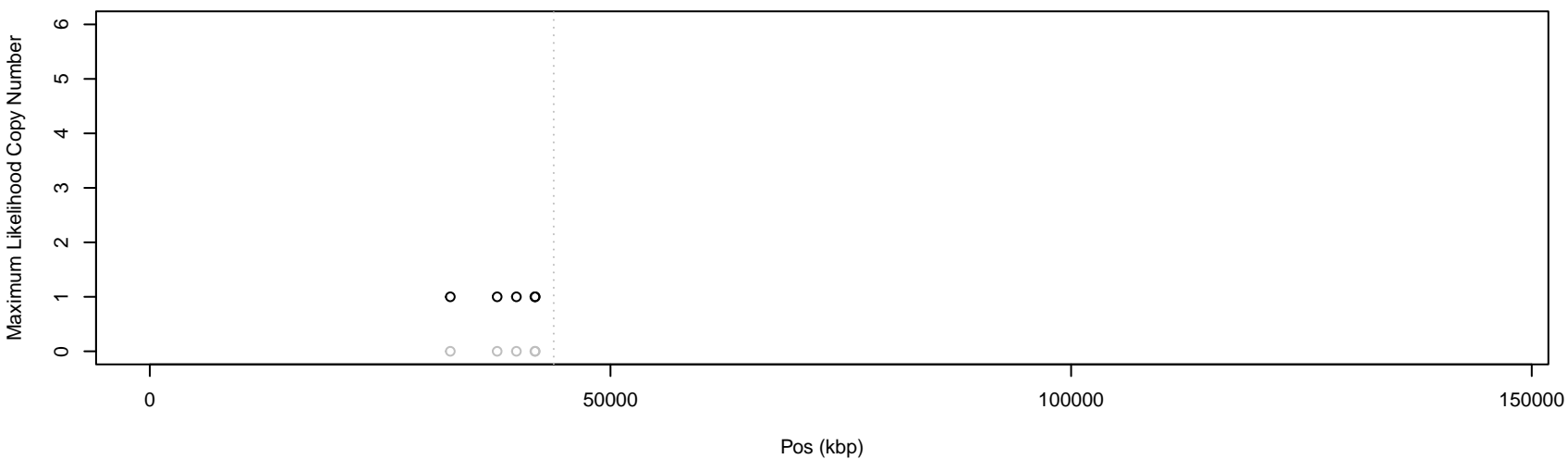
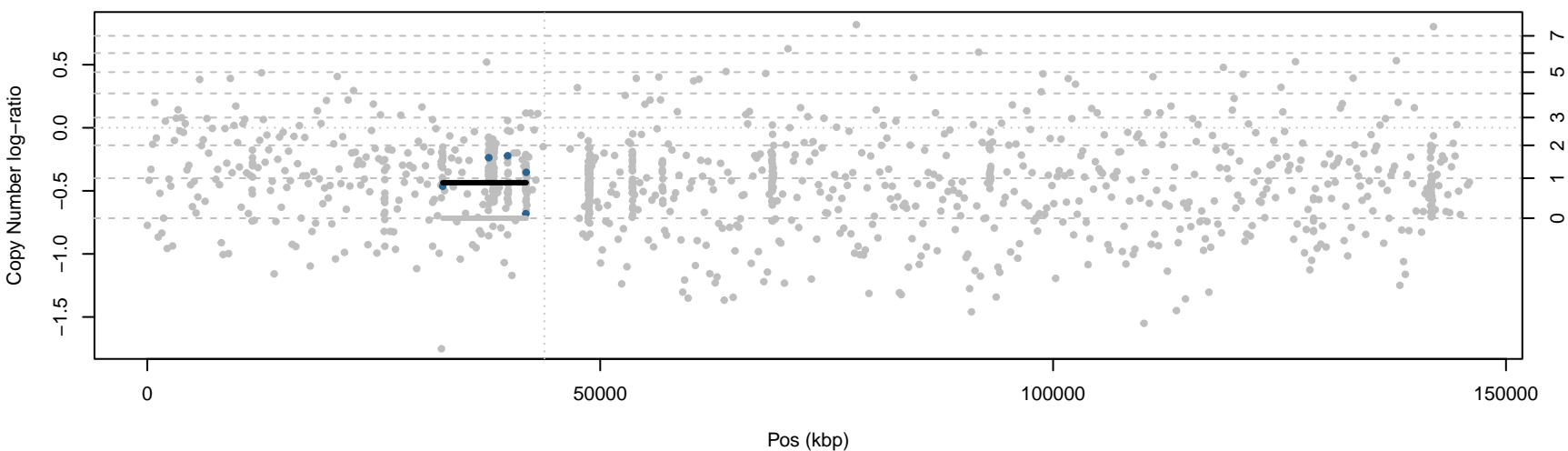
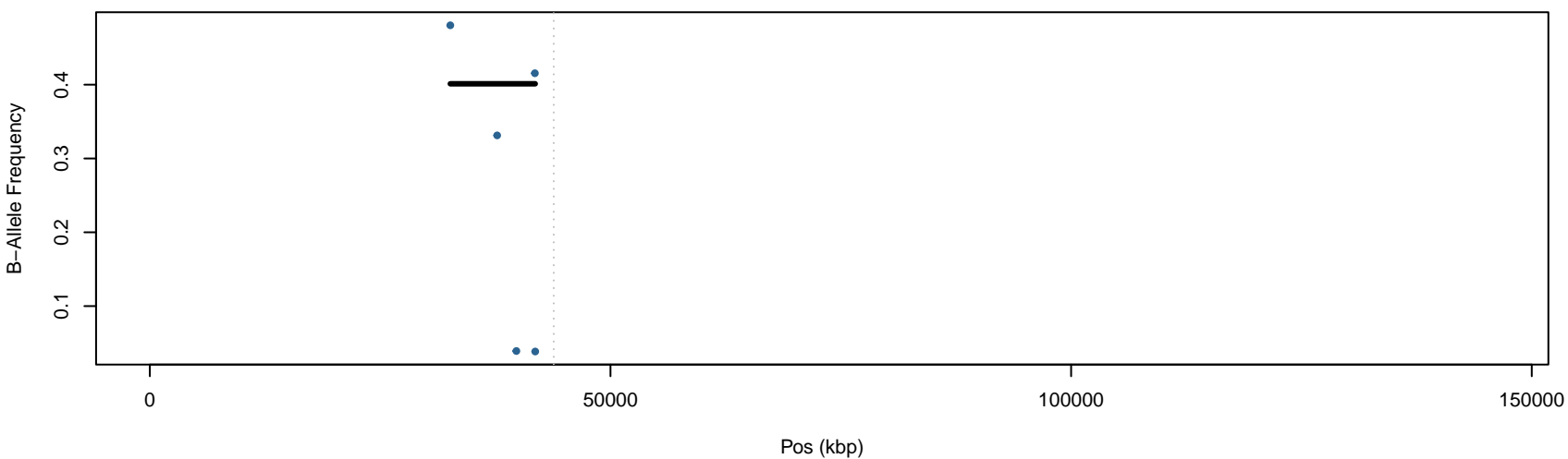
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 5



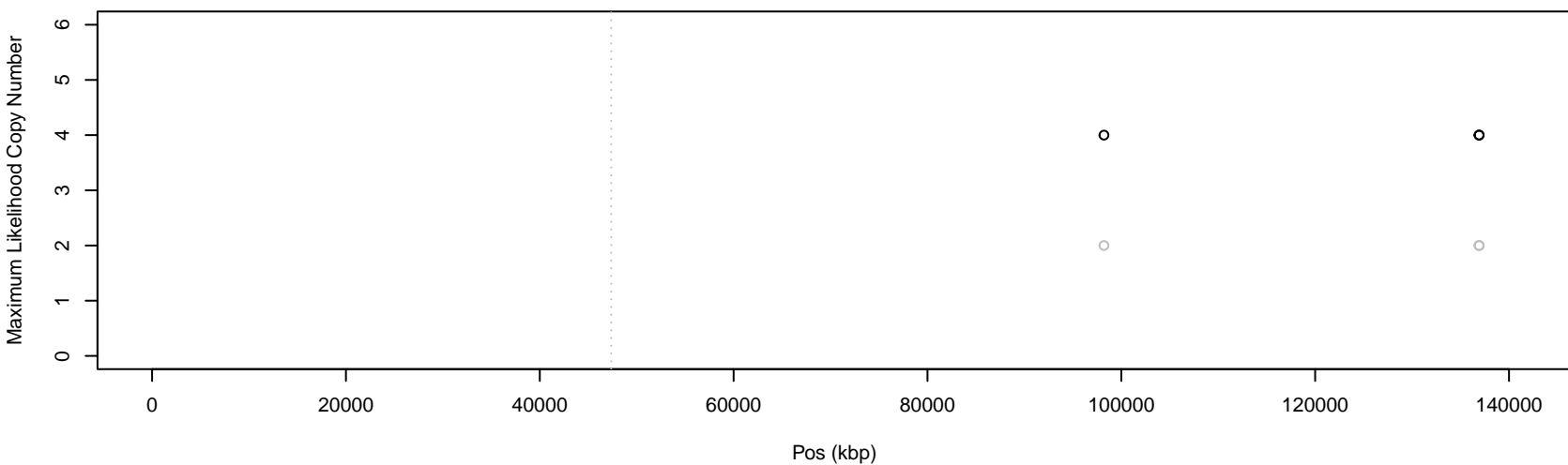
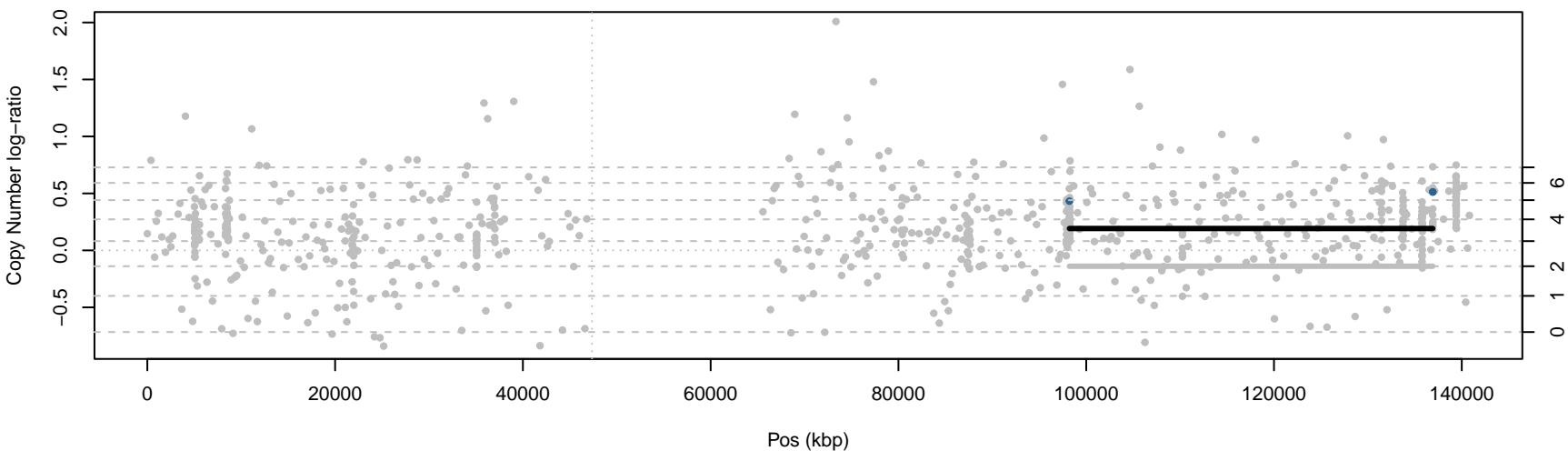
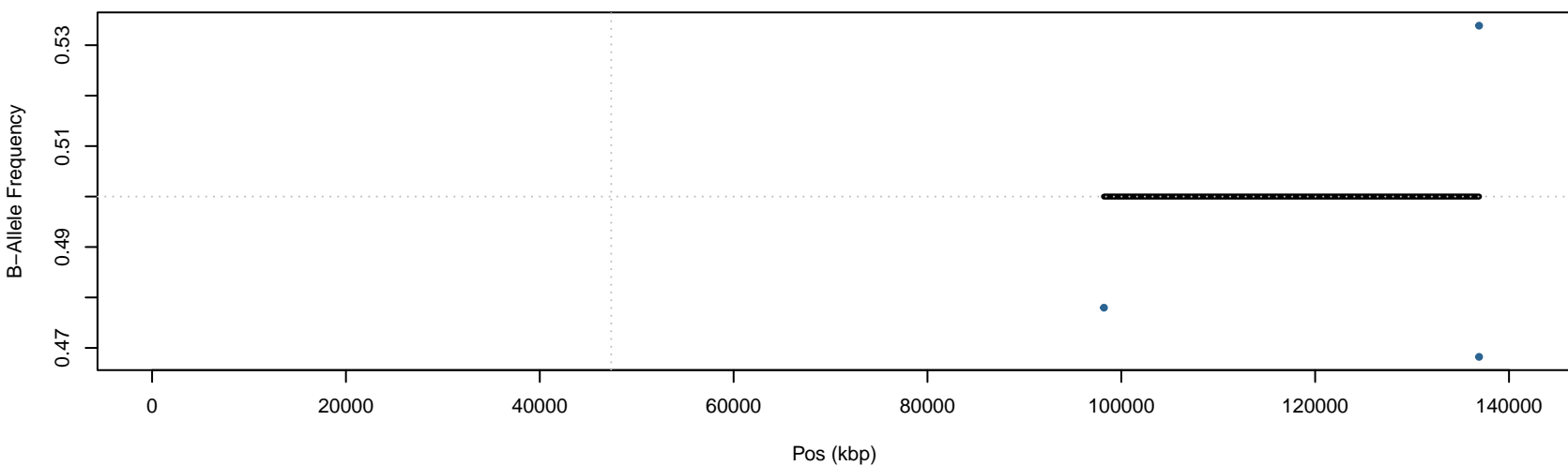
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 6



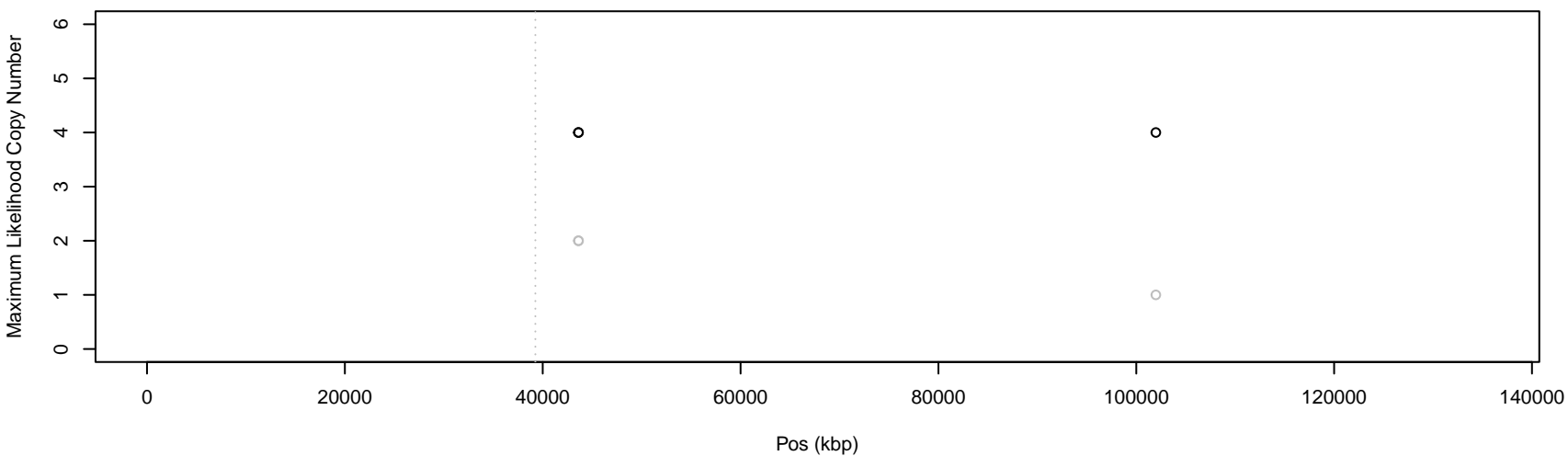
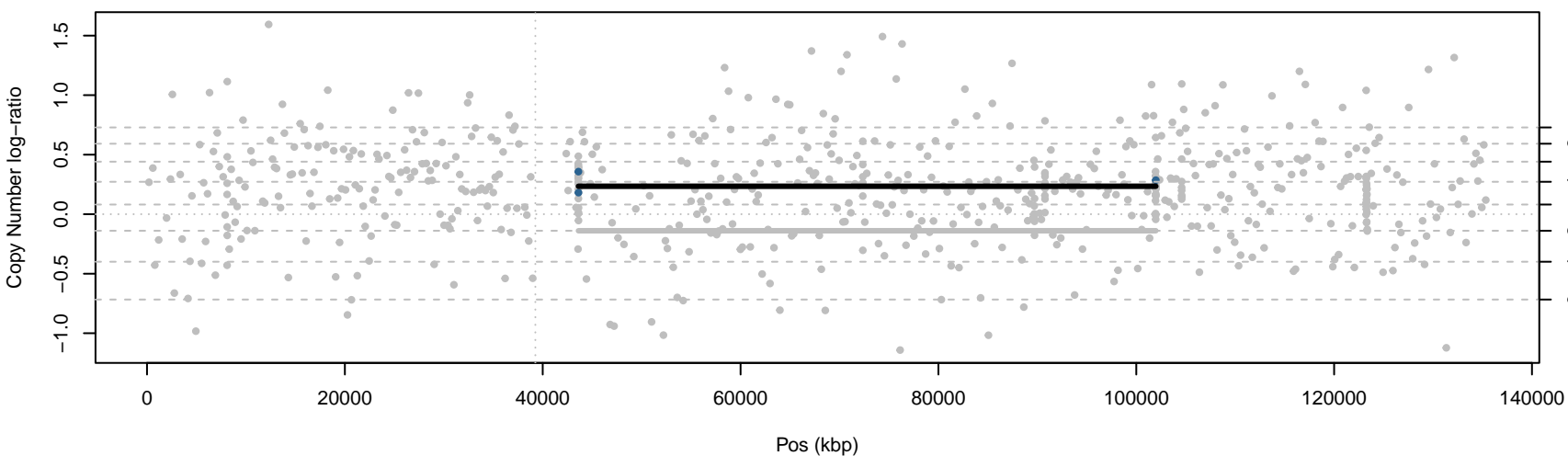
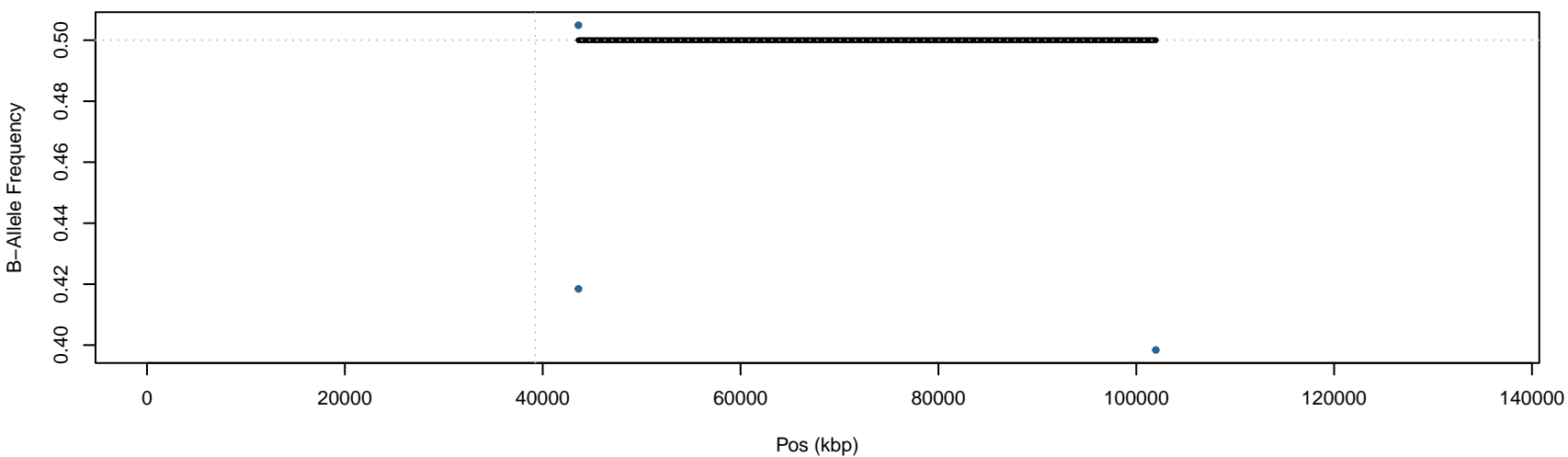
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 8



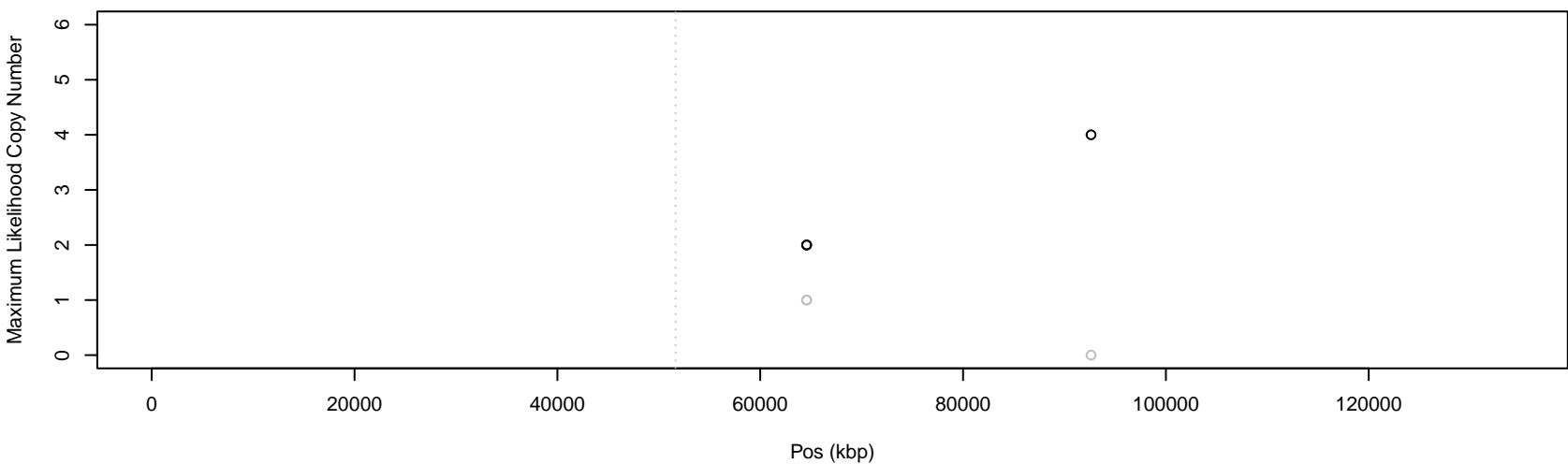
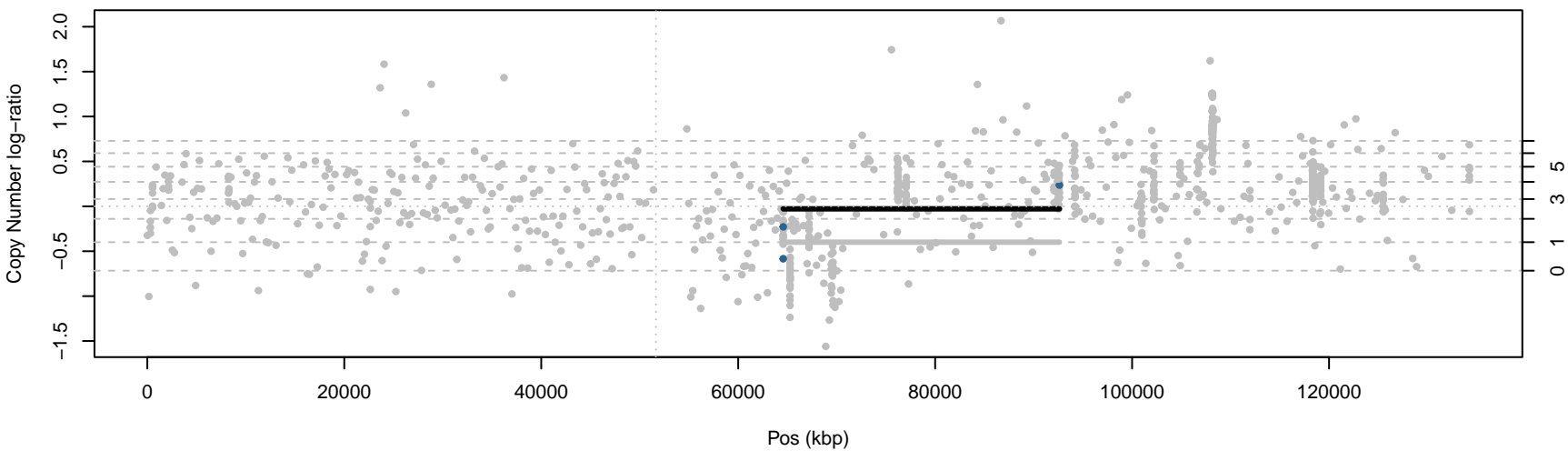
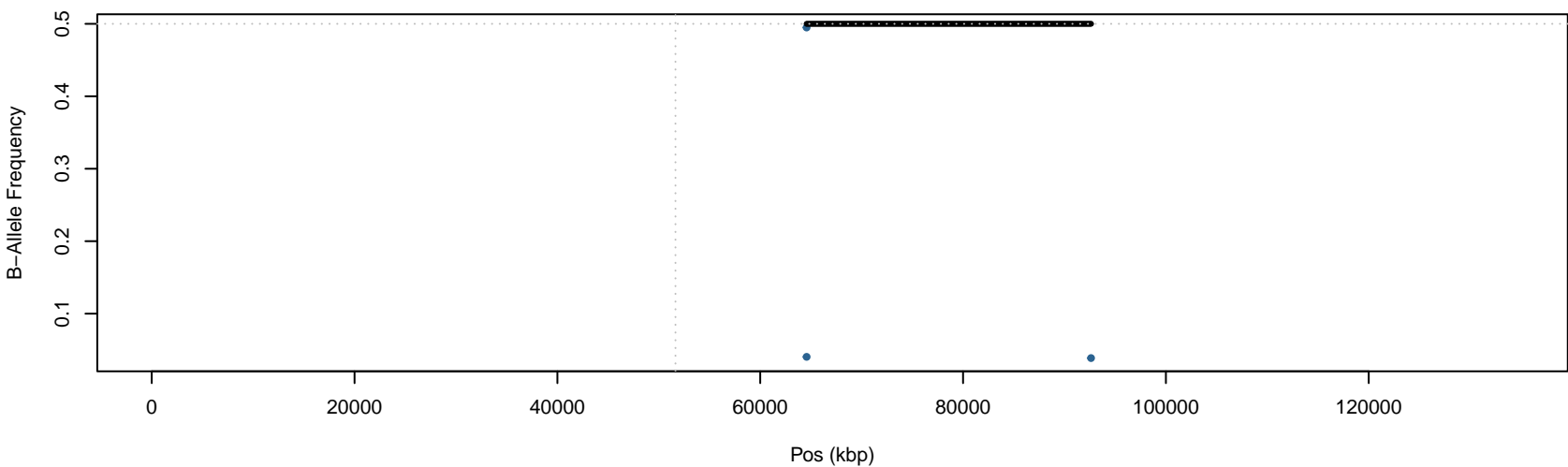
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 9



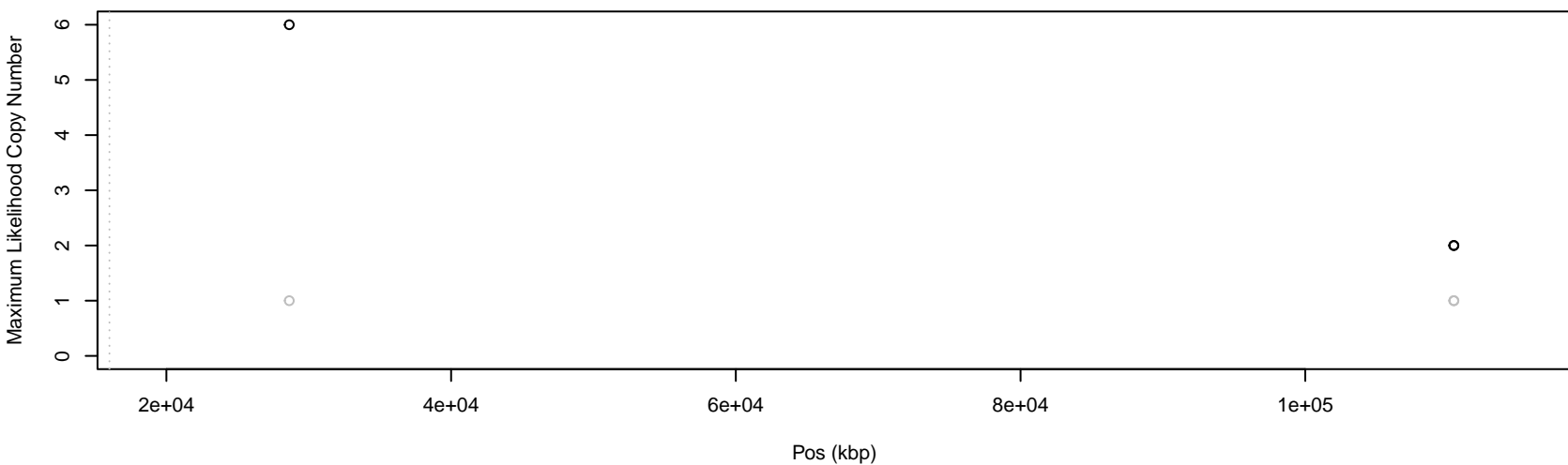
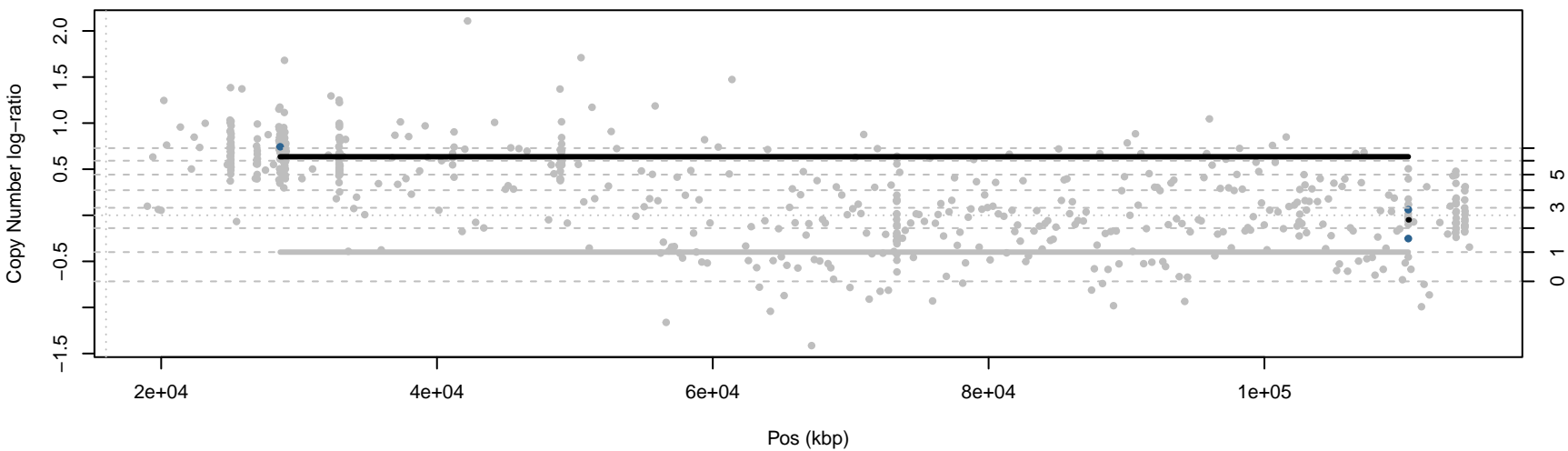
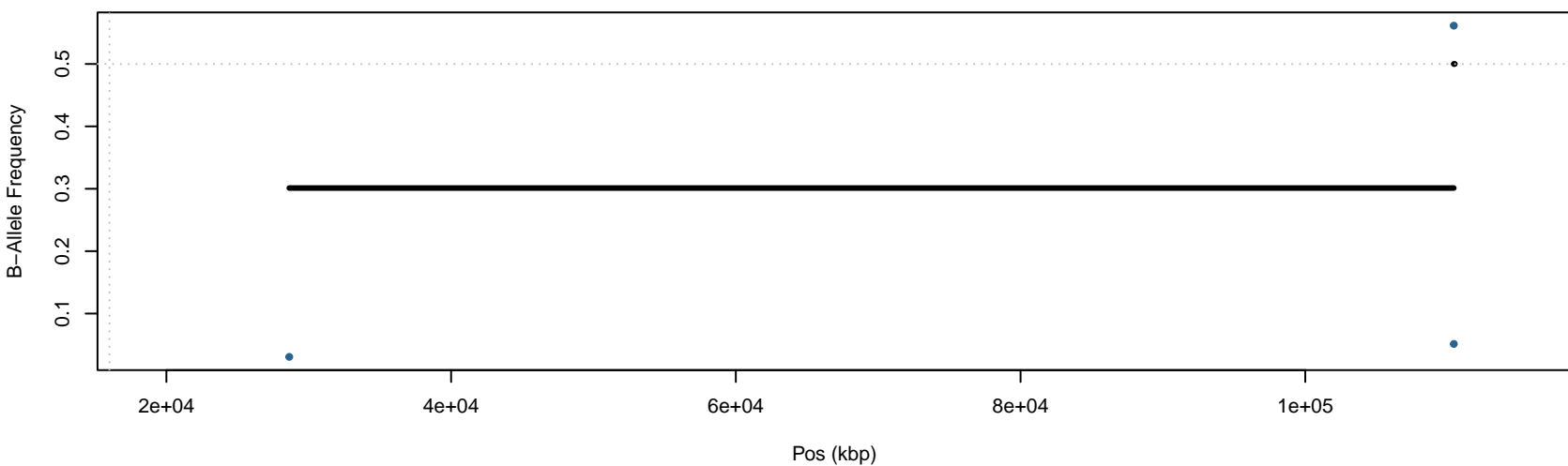
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 10



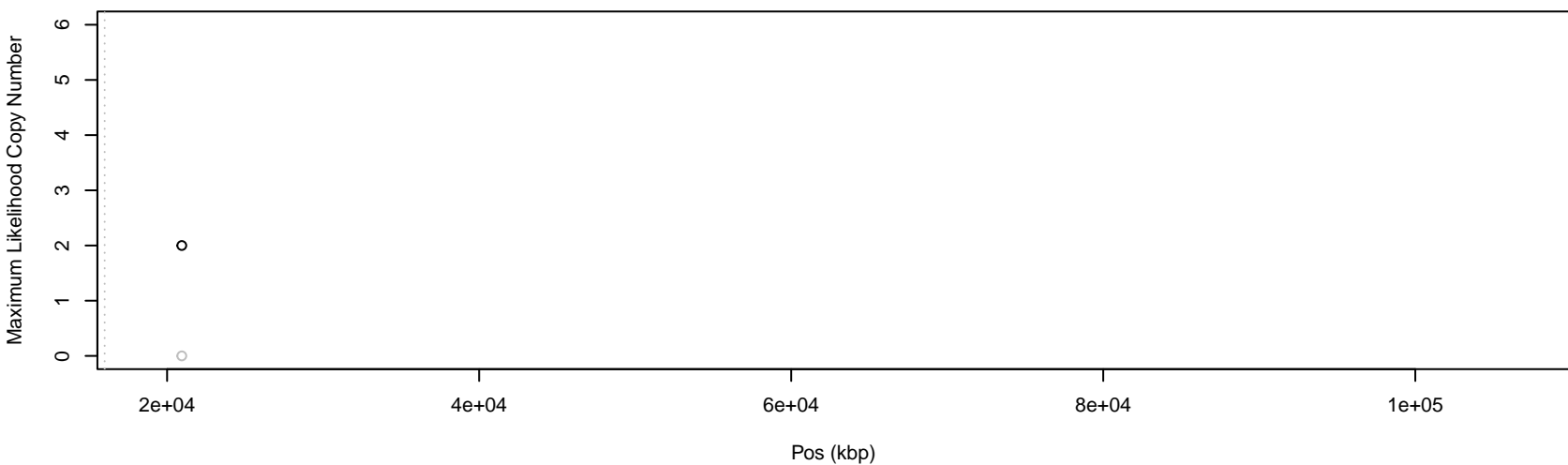
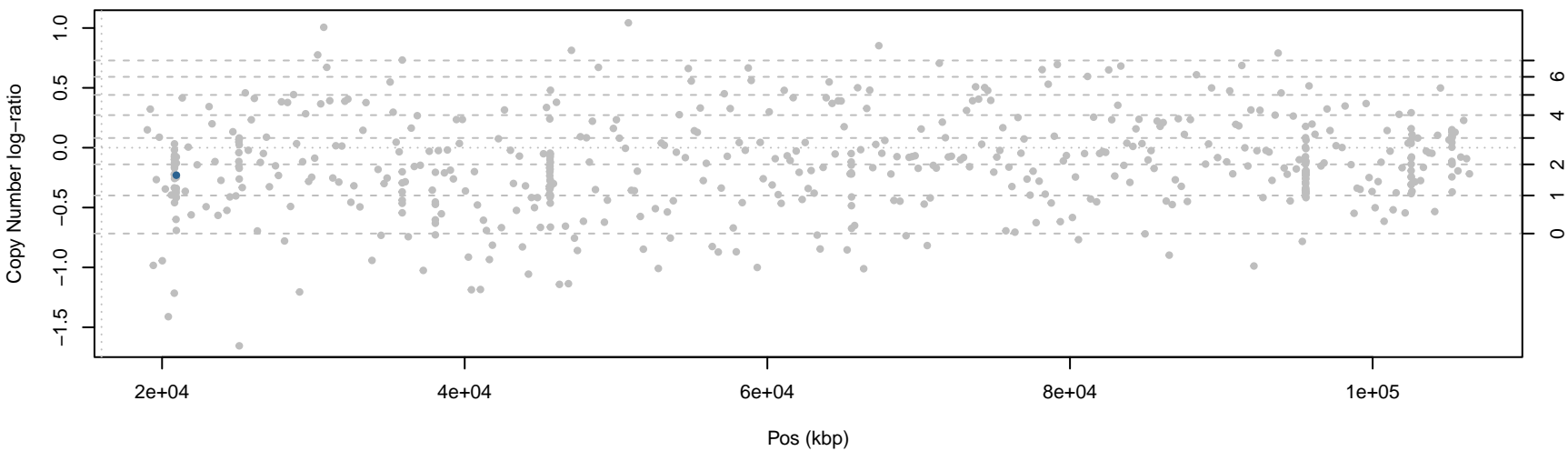
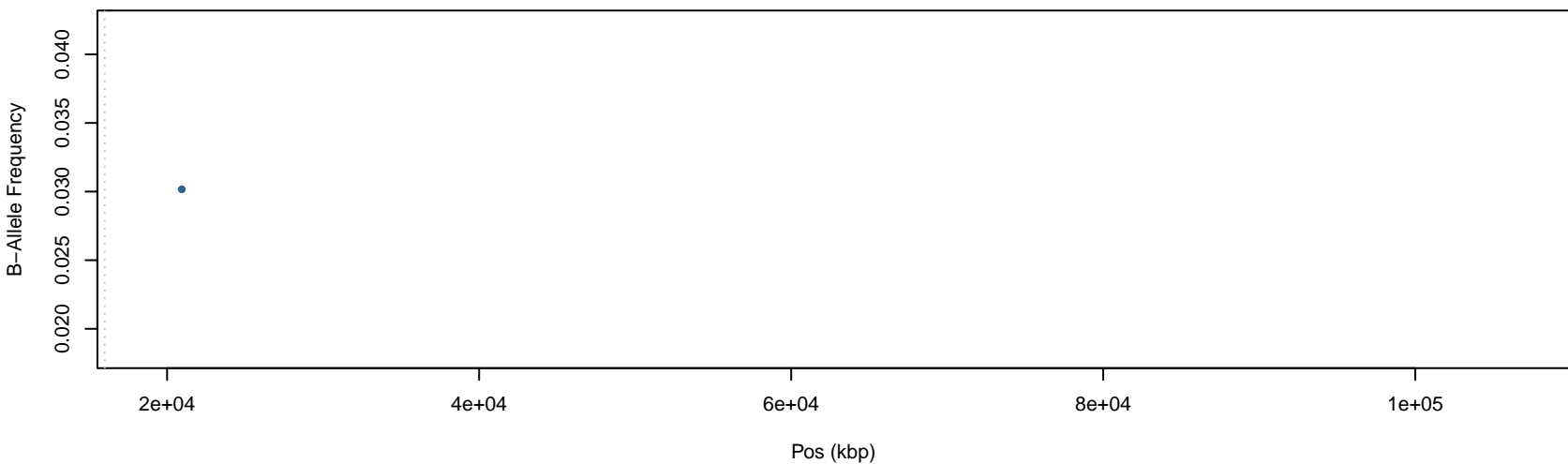
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 11



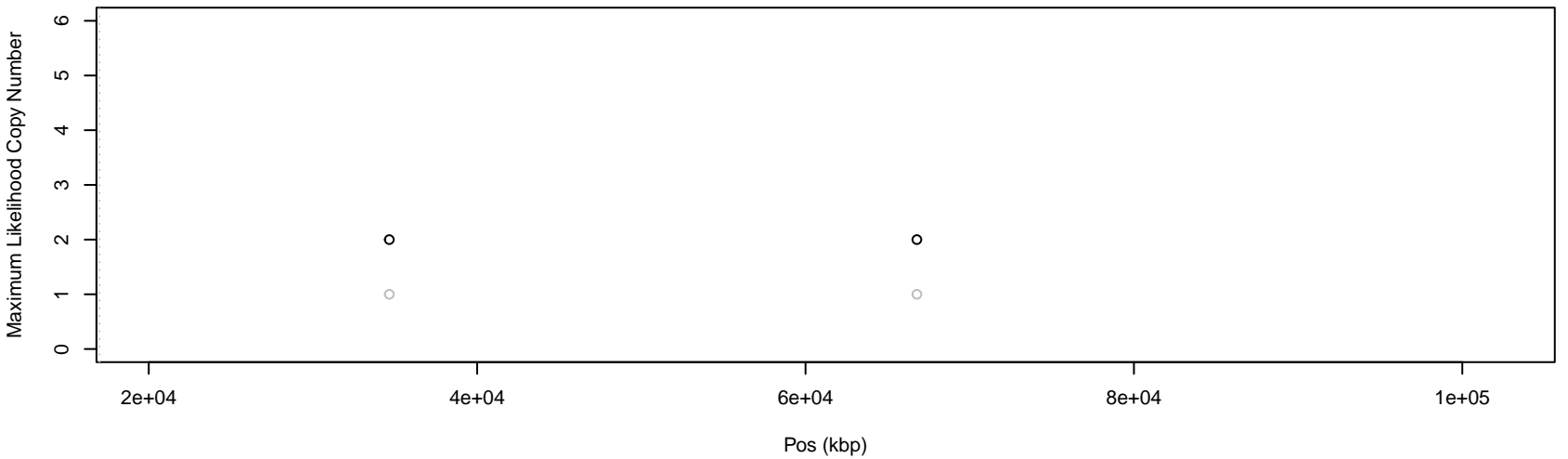
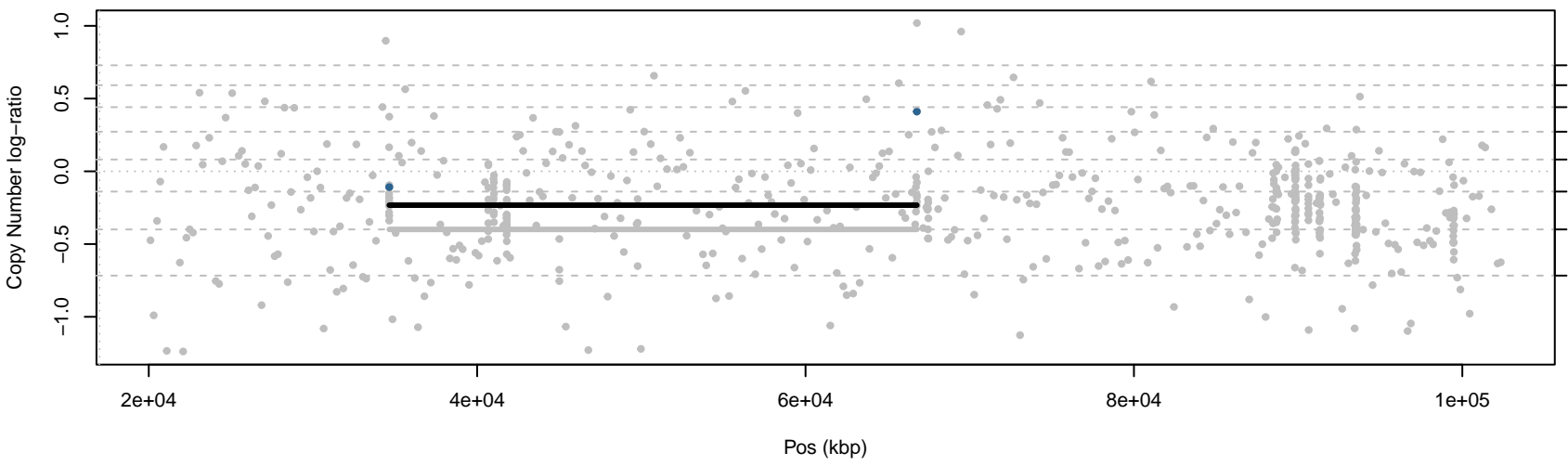
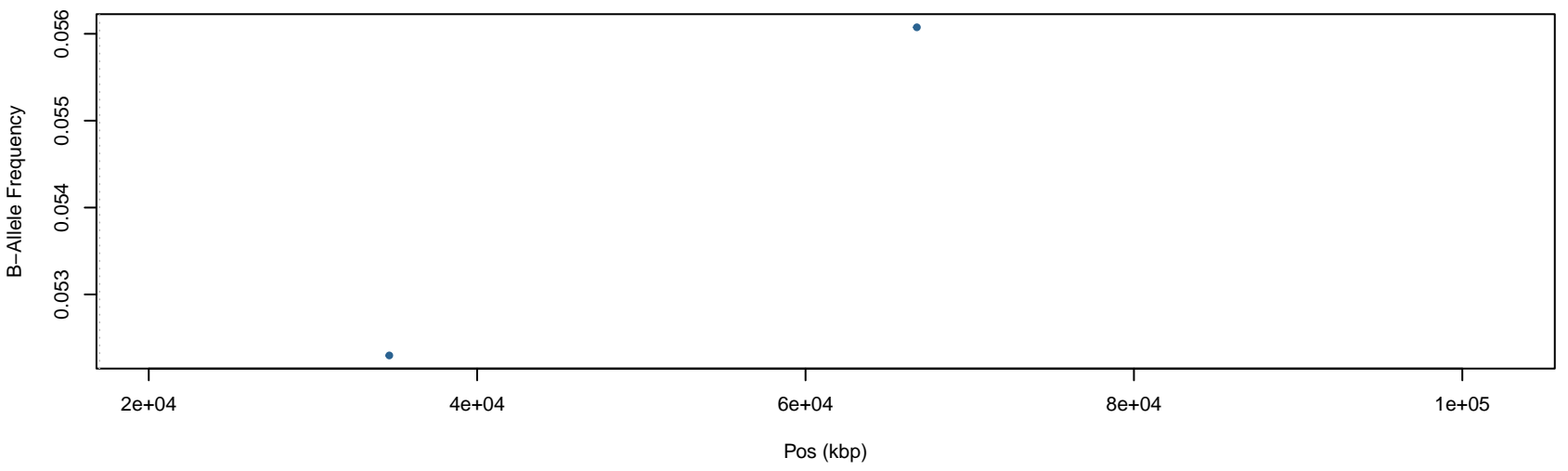
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 13



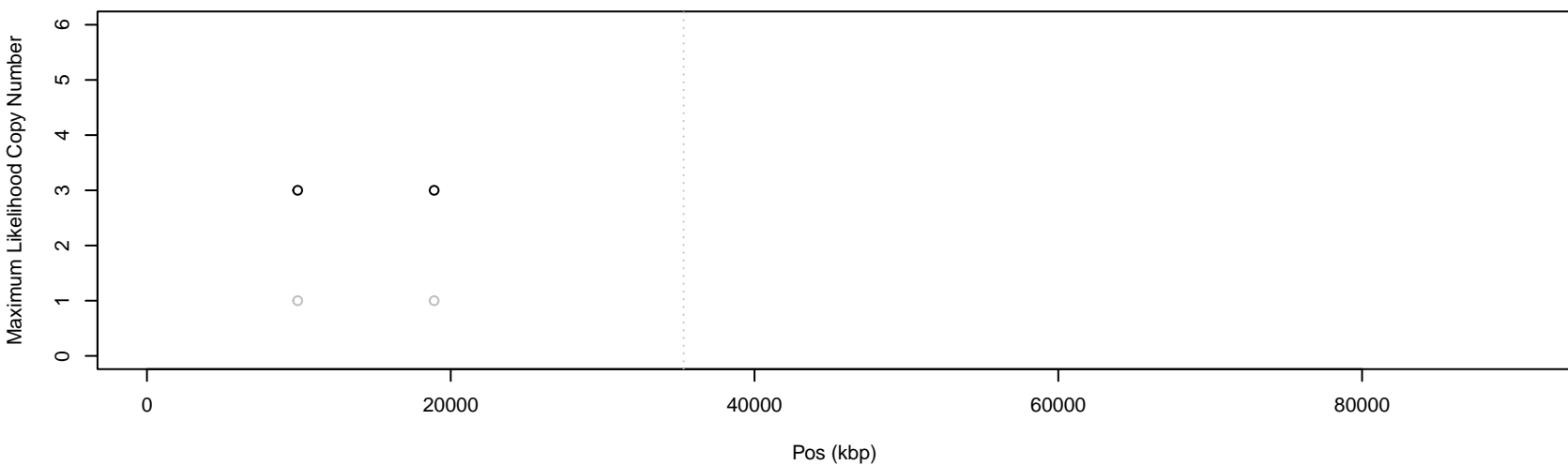
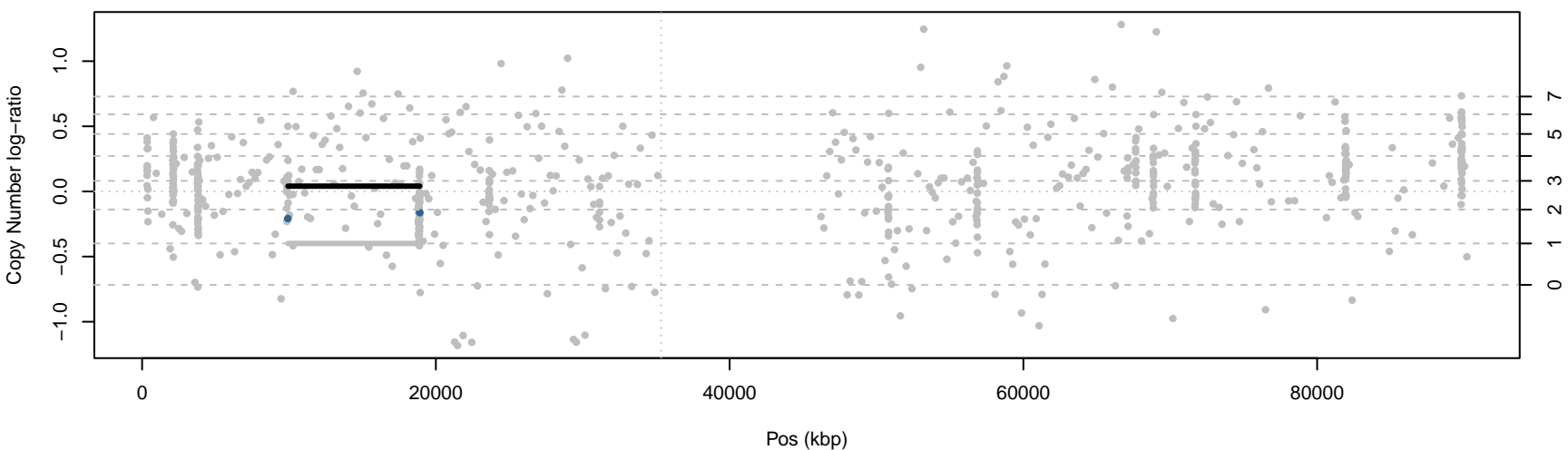
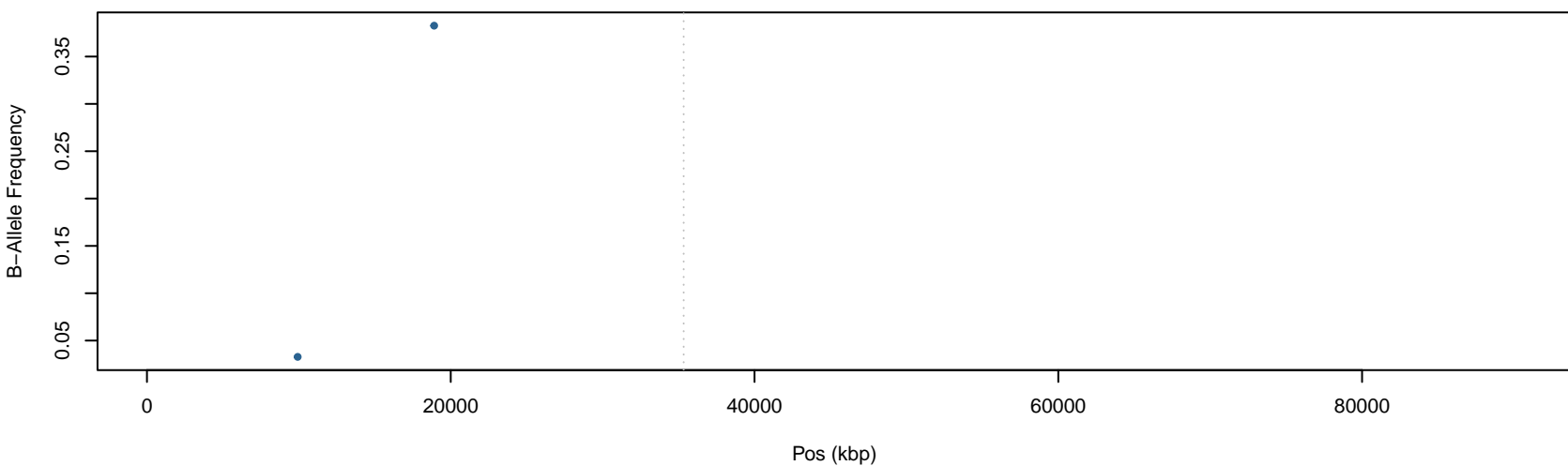
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 14



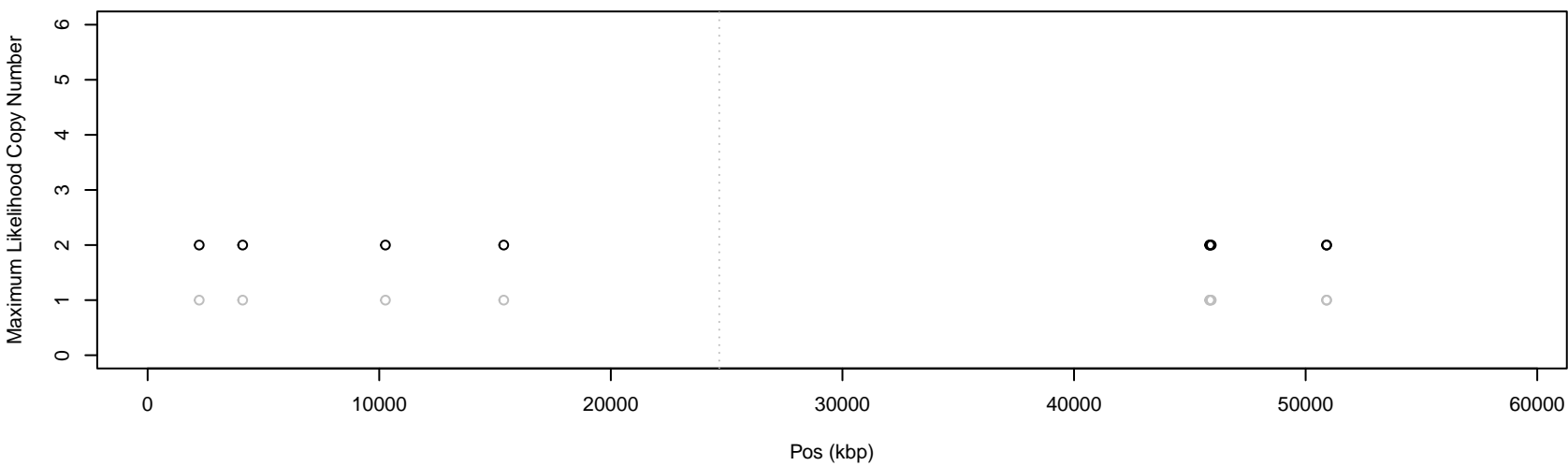
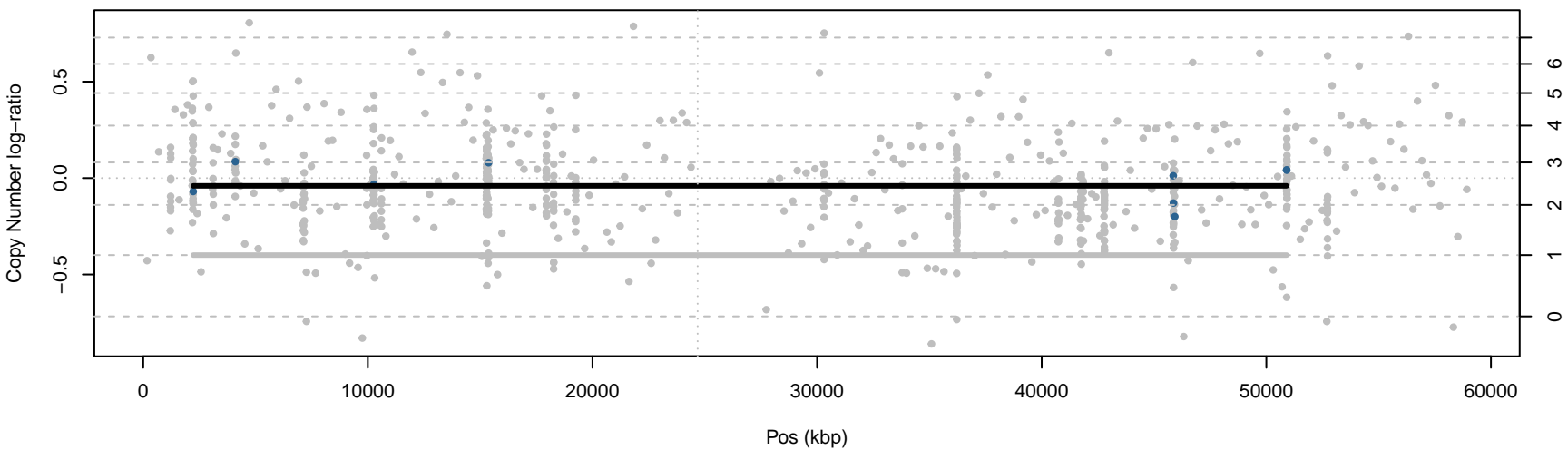
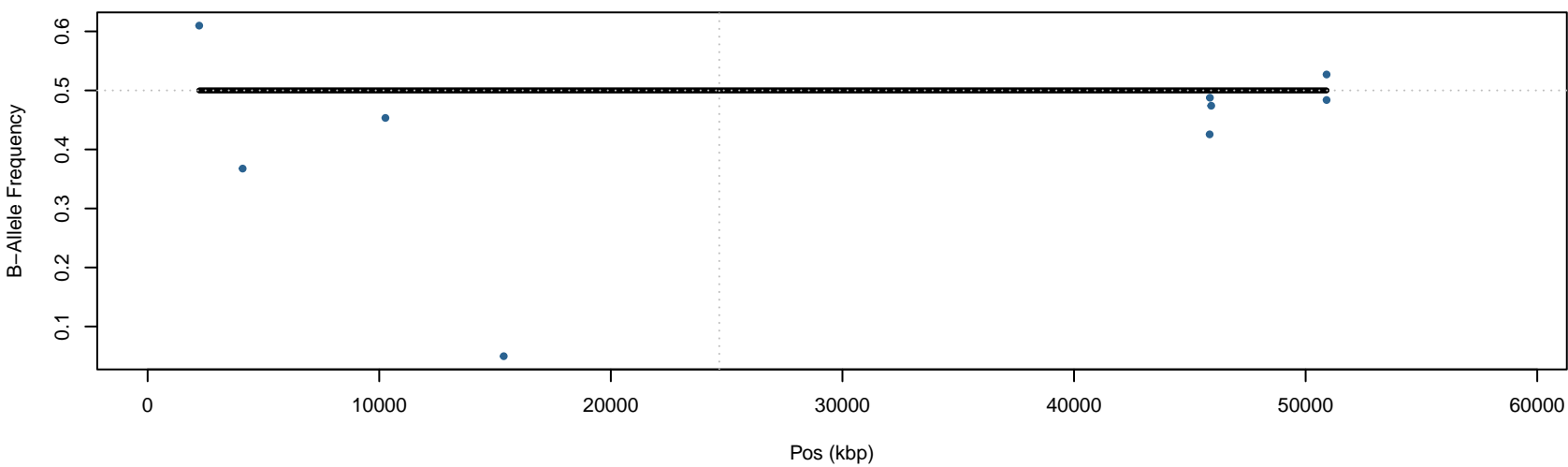
Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 15



Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 16



Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 19



Purity: 0.33 Tumor ploidy: 2.614 SNV log-likelihood: 8.06 GoF: 81.8% Mean coverage: 555;408 Chromosome: 20

