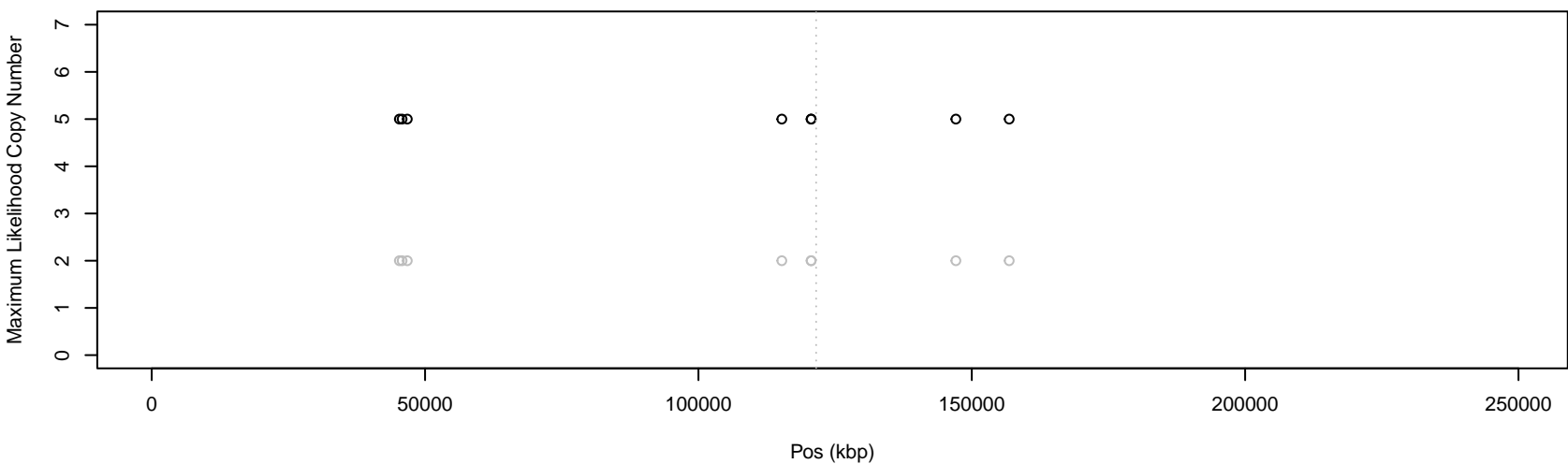
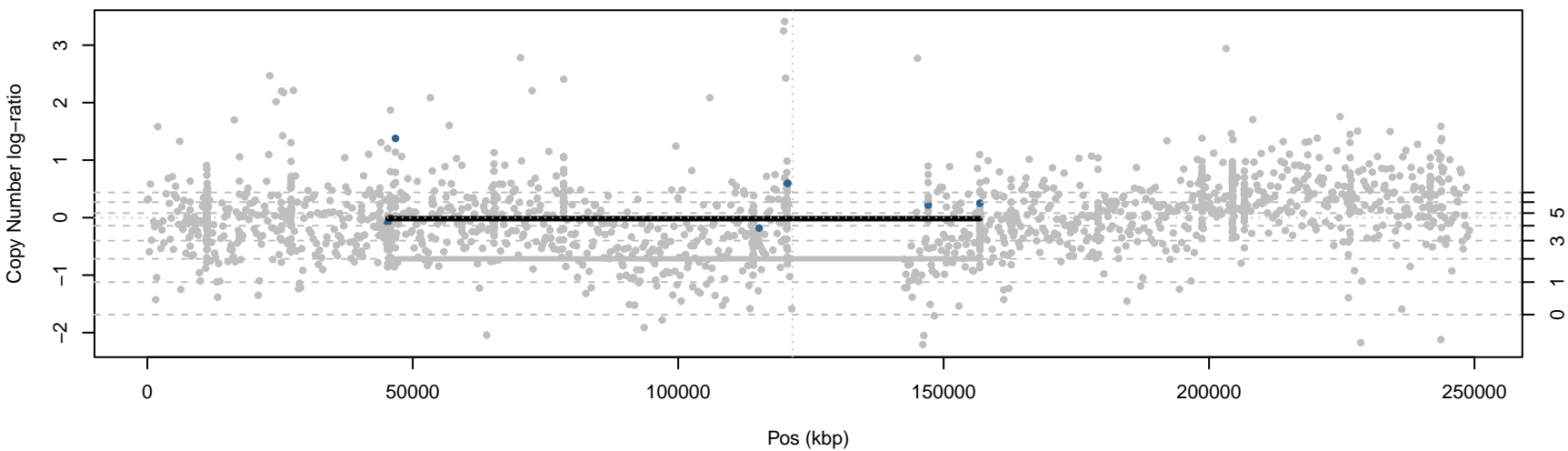
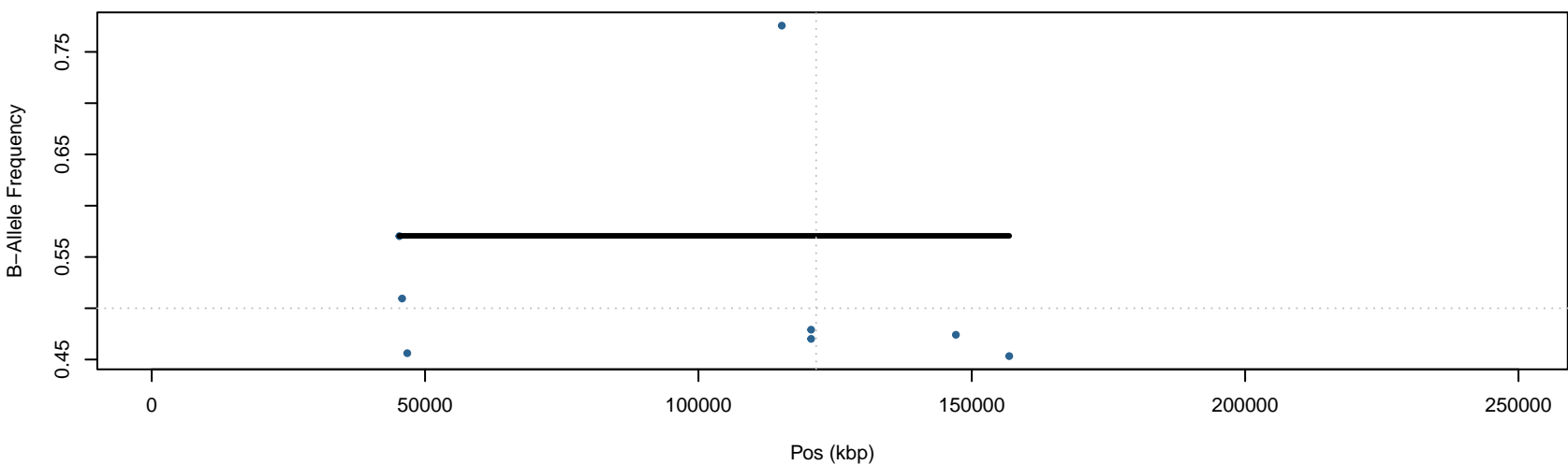
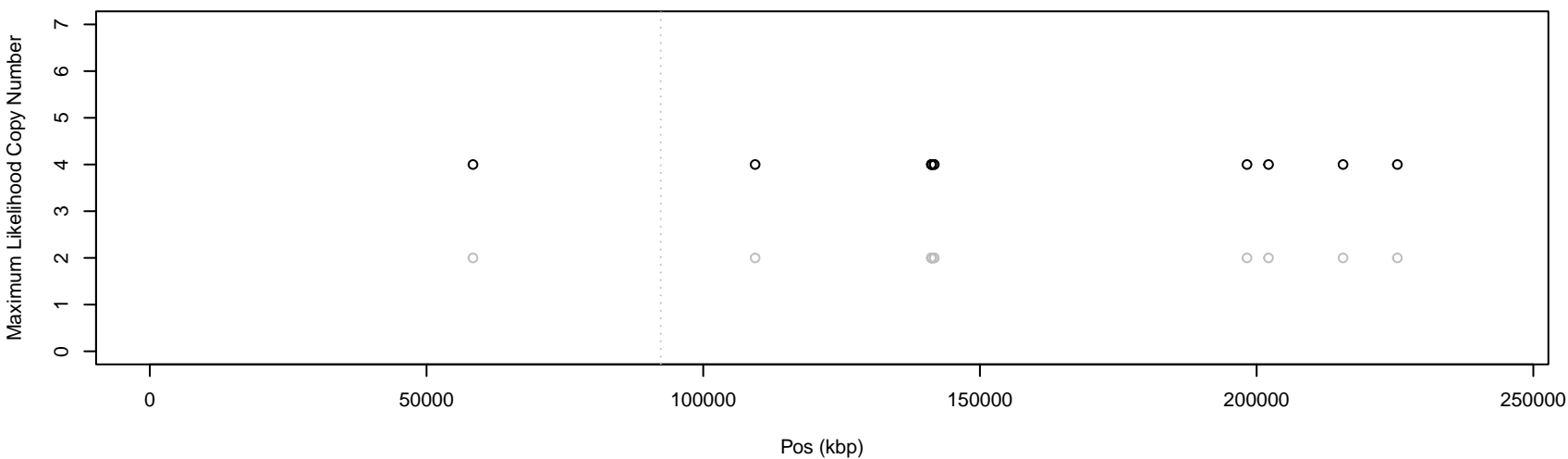
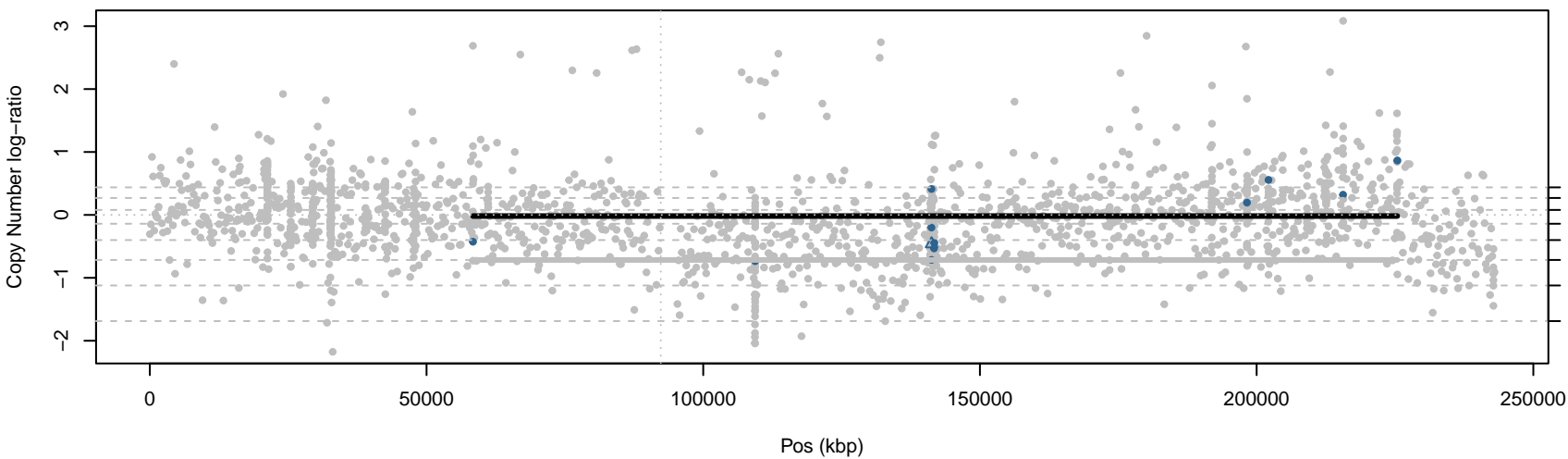
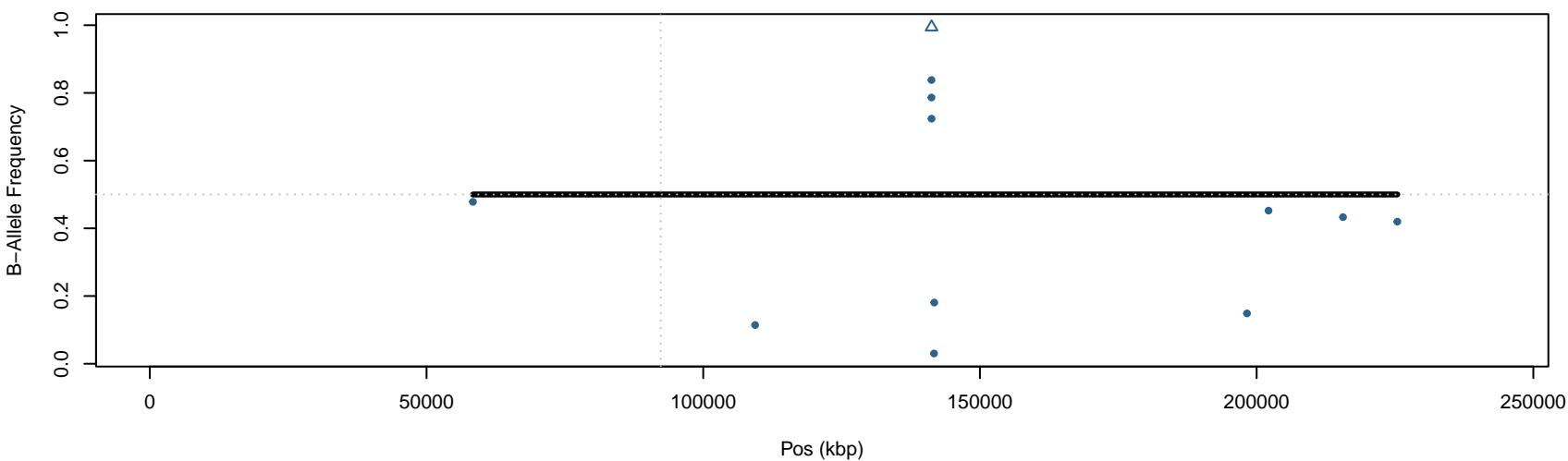


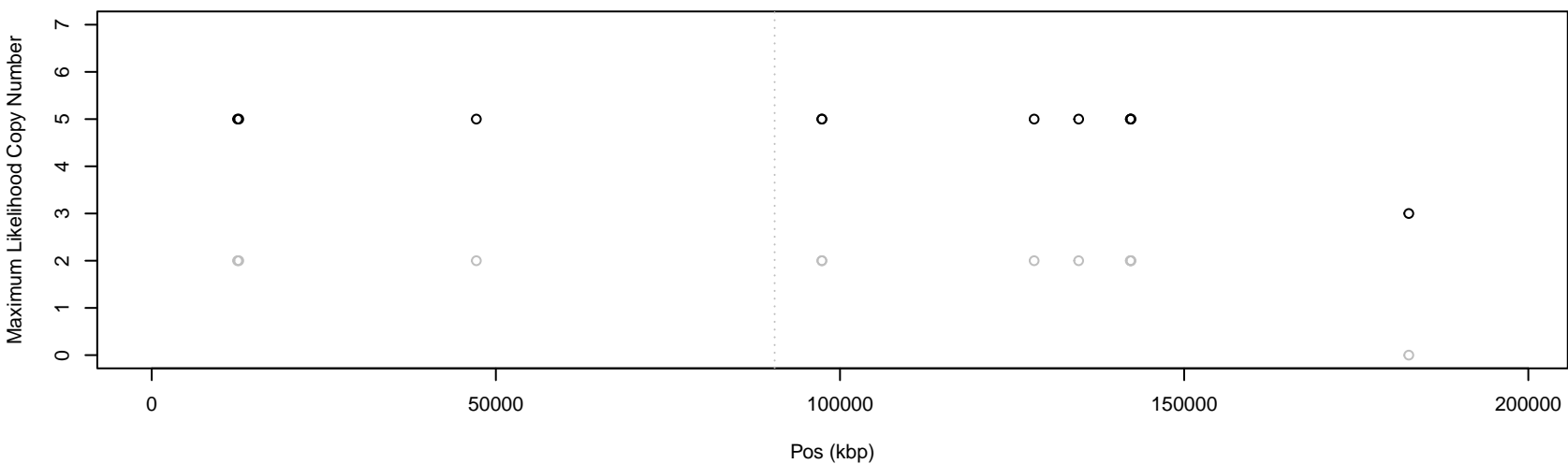
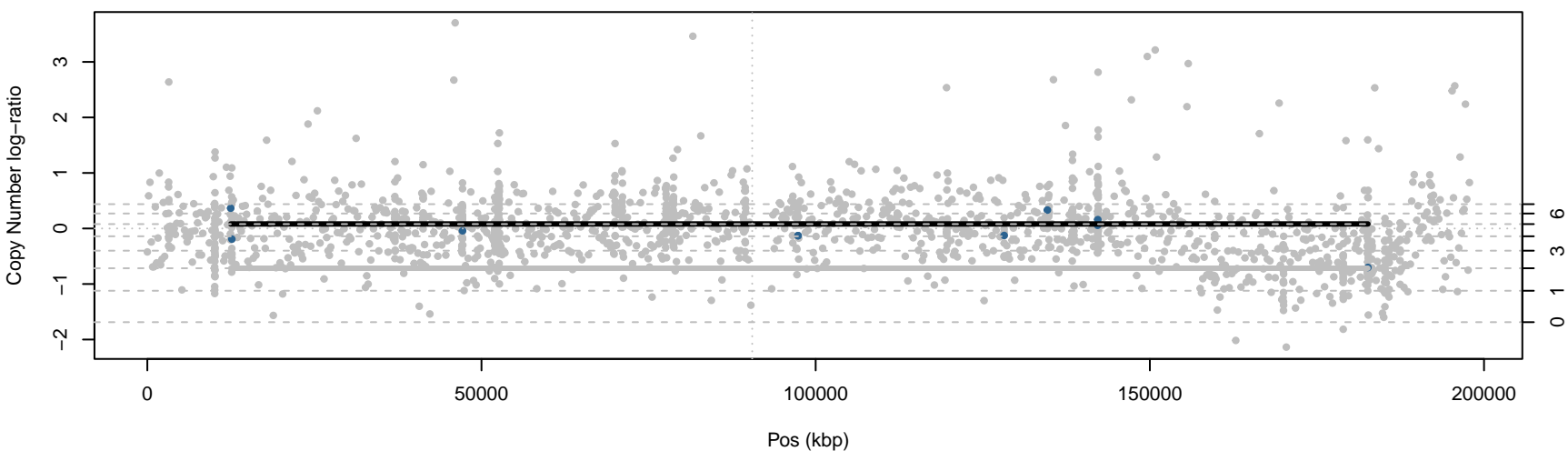
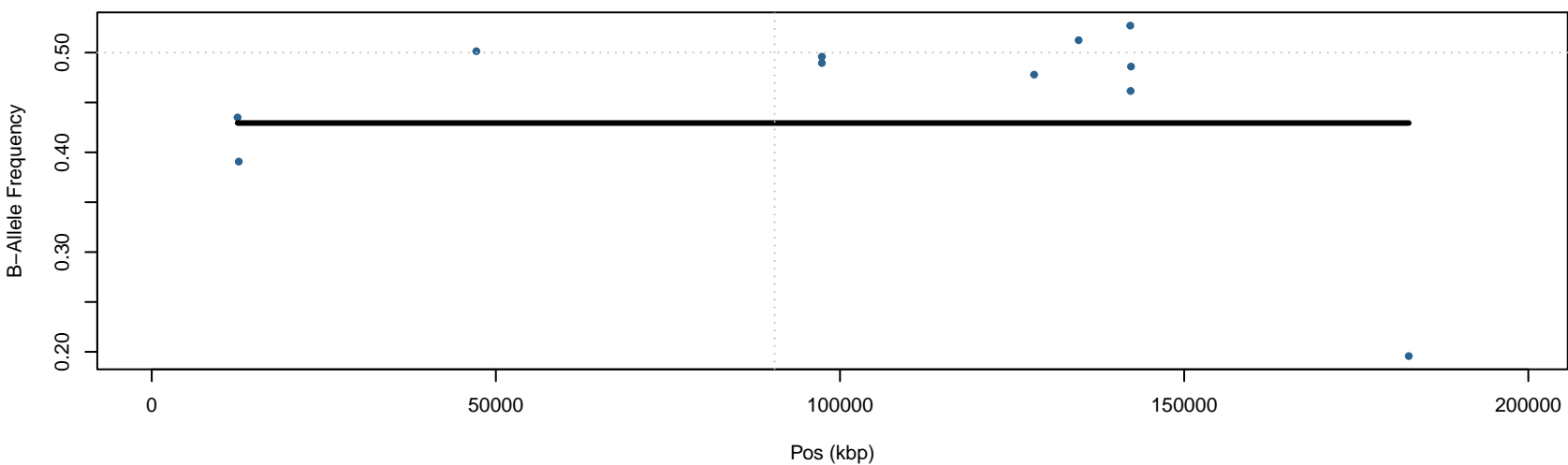
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 1



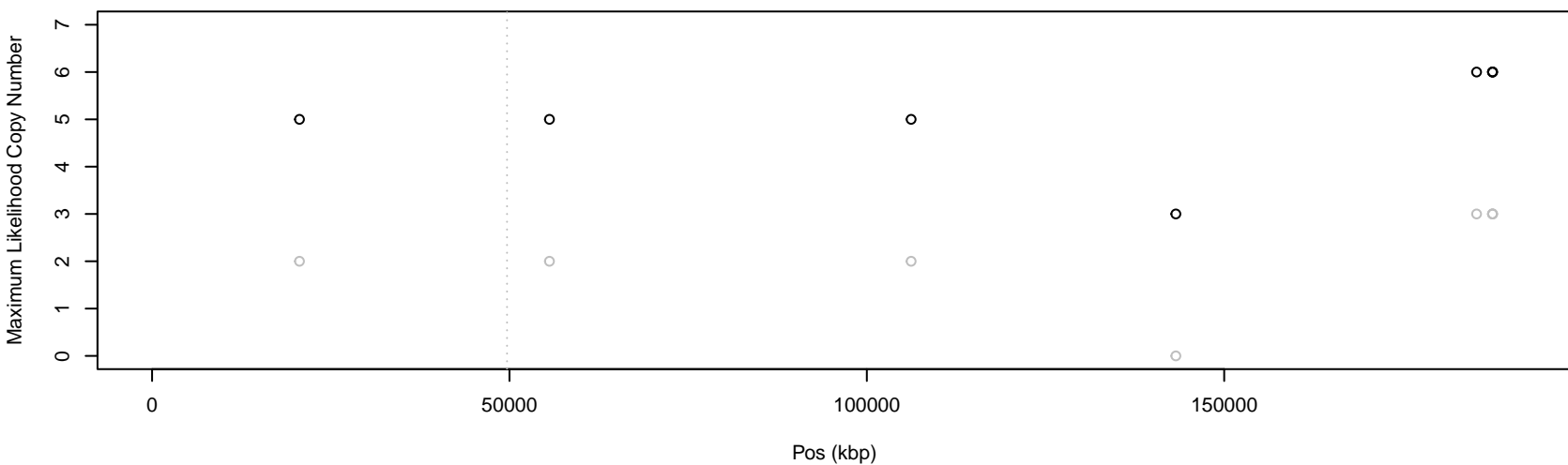
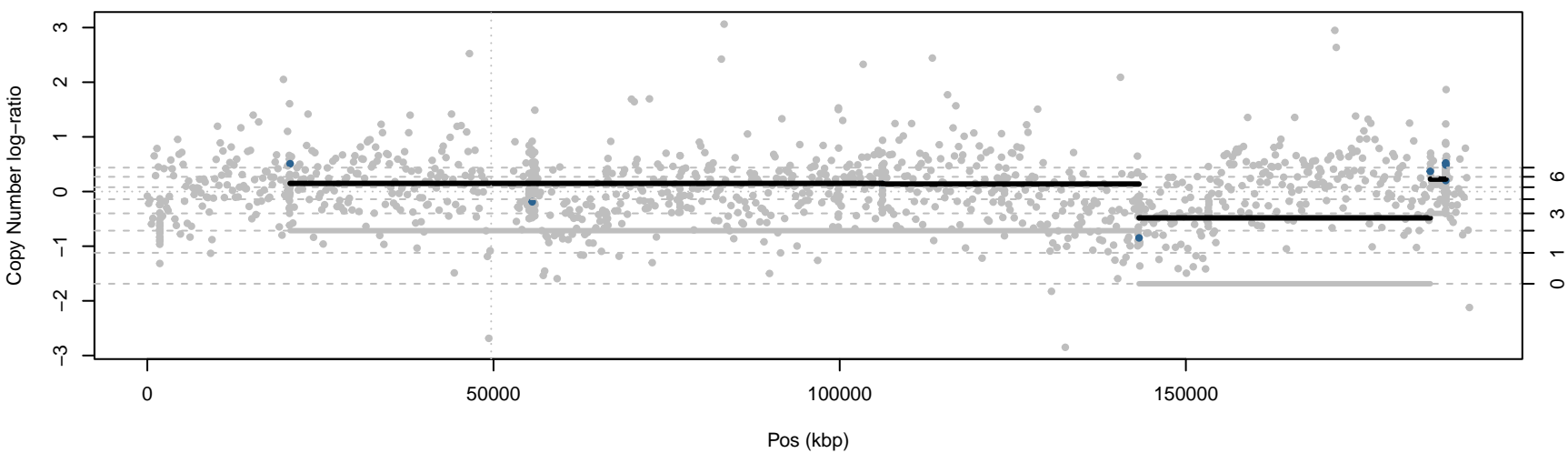
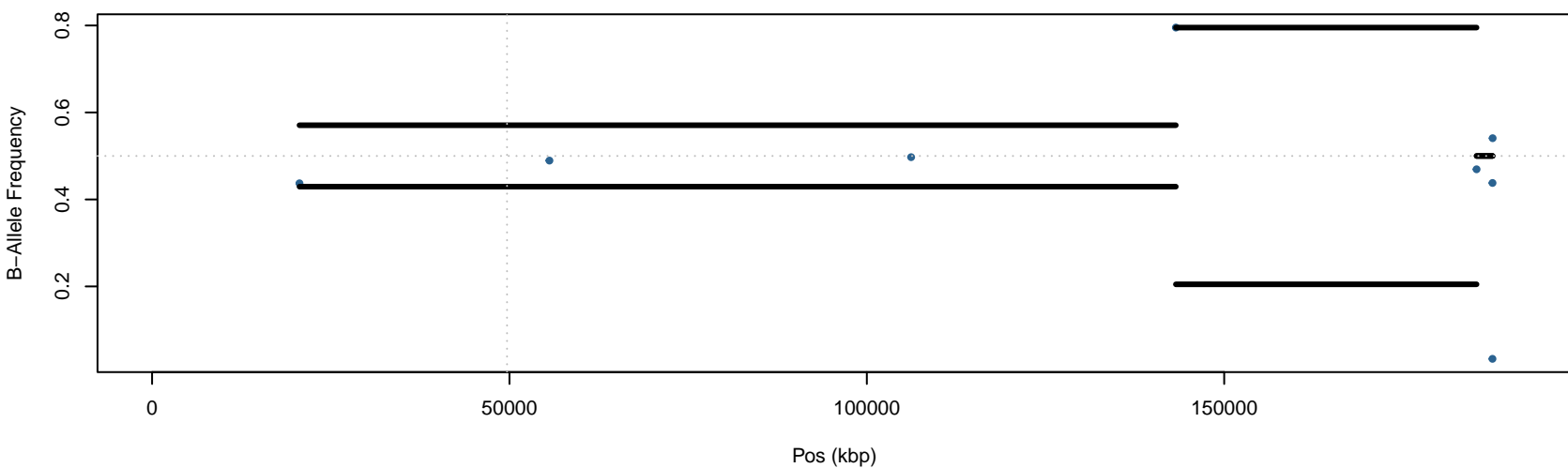
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 2



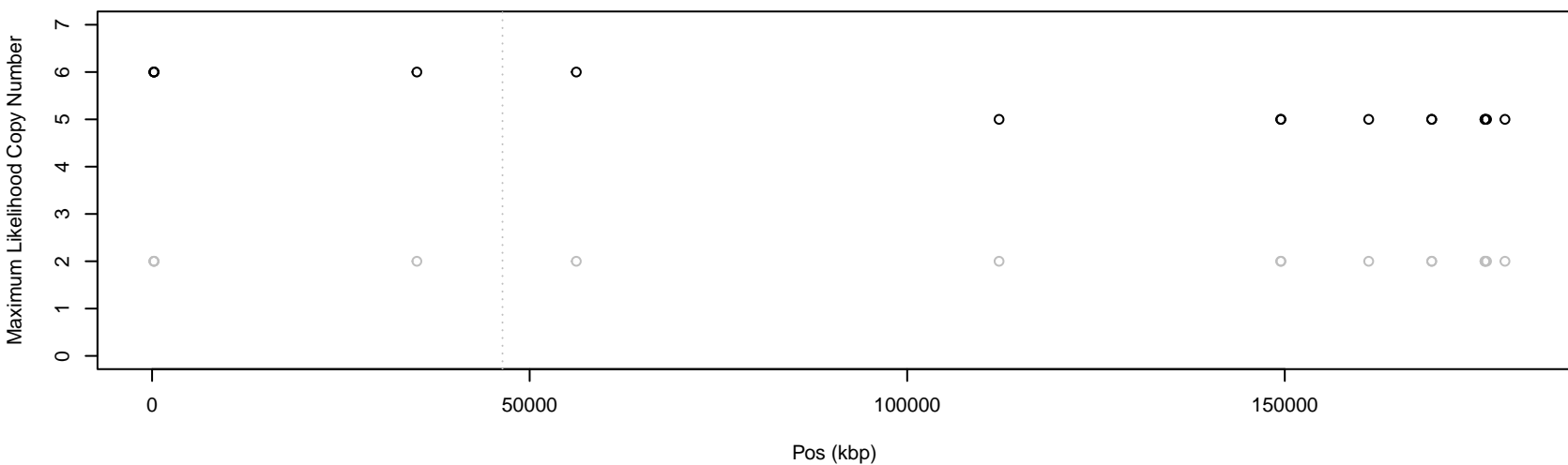
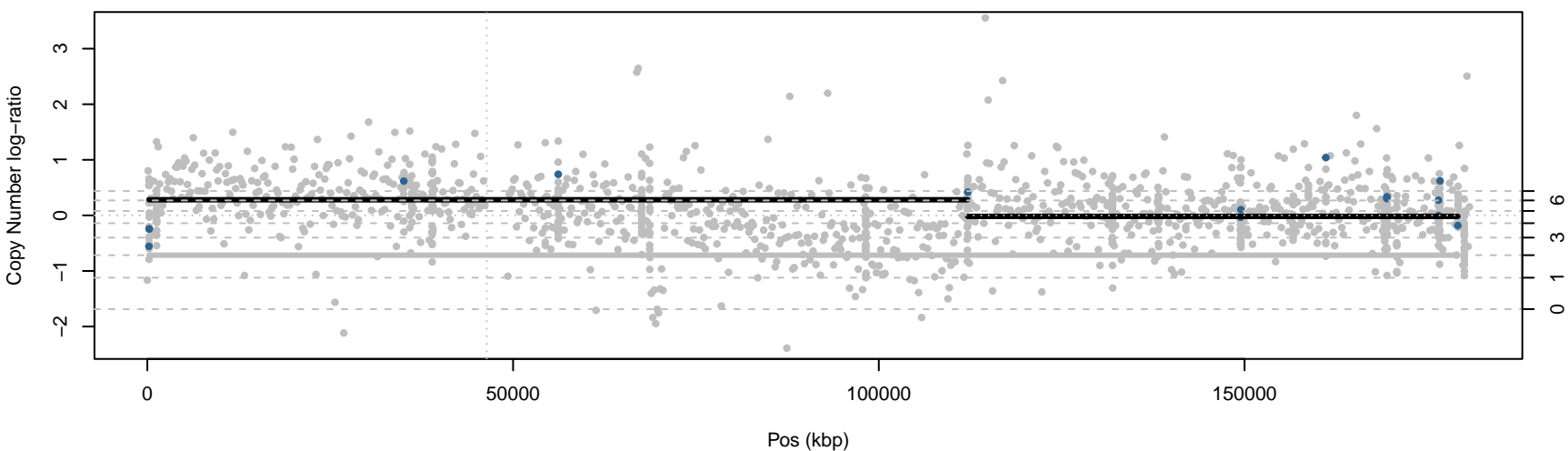
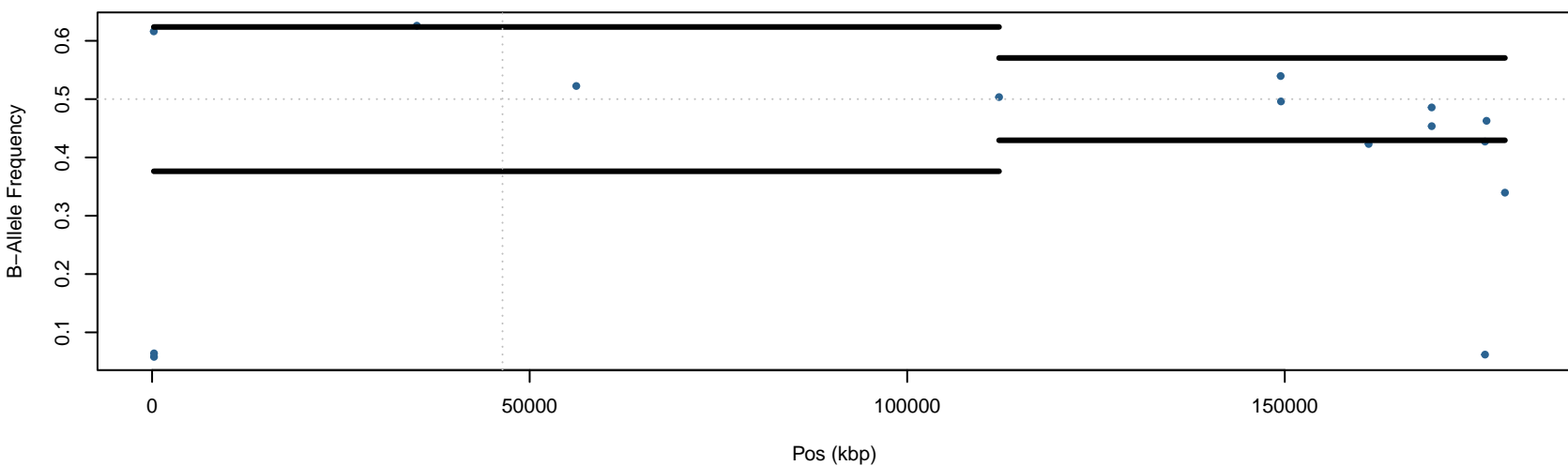
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 3



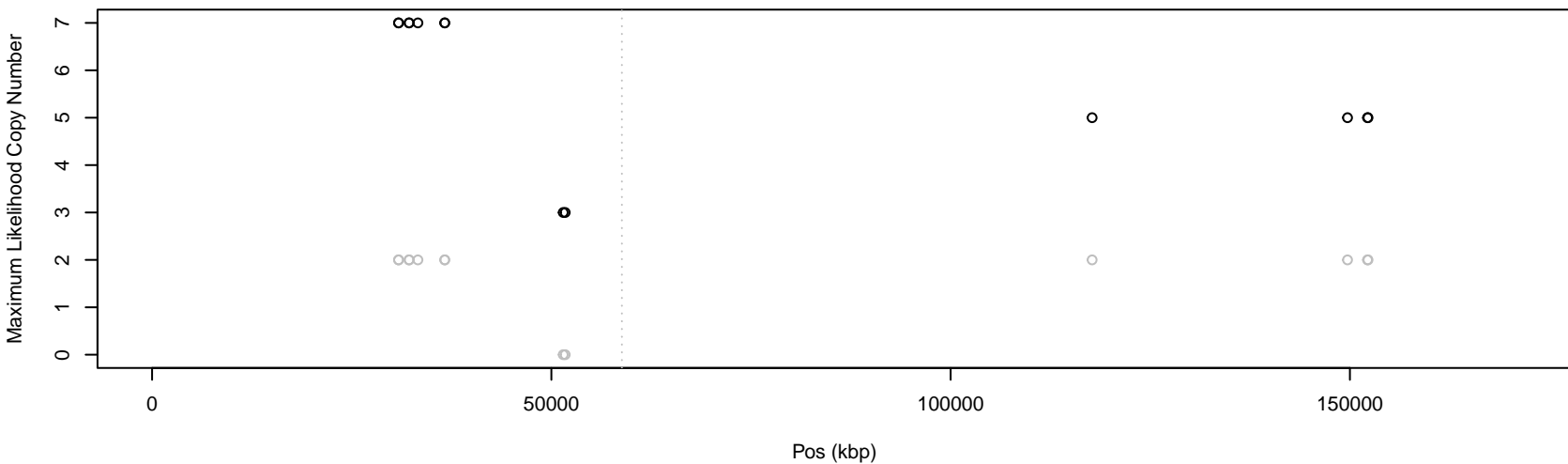
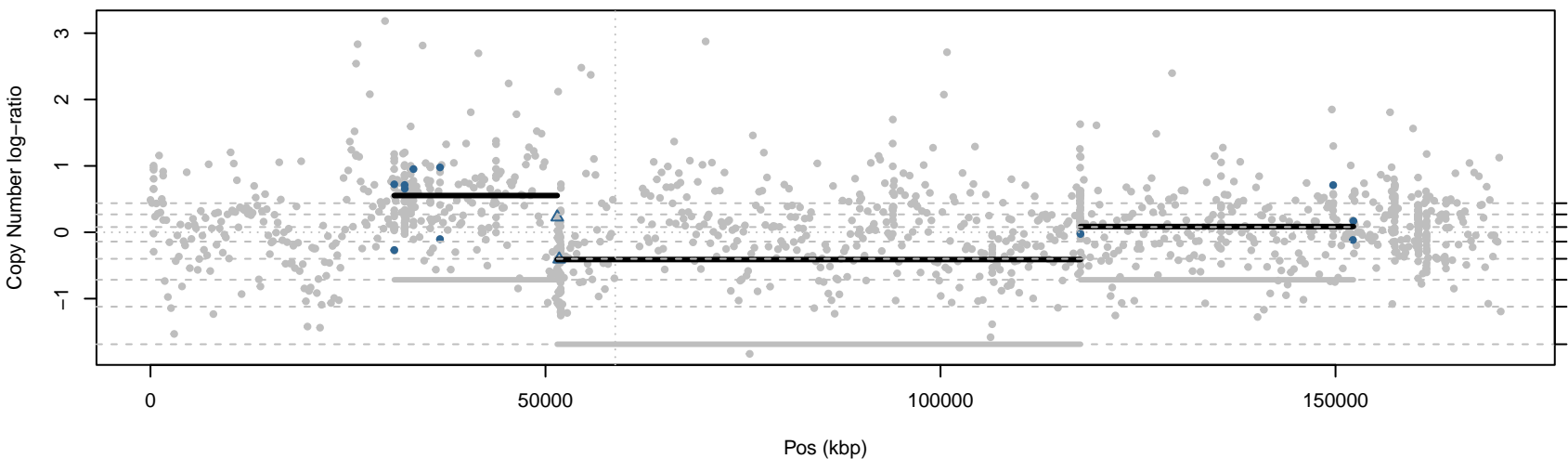
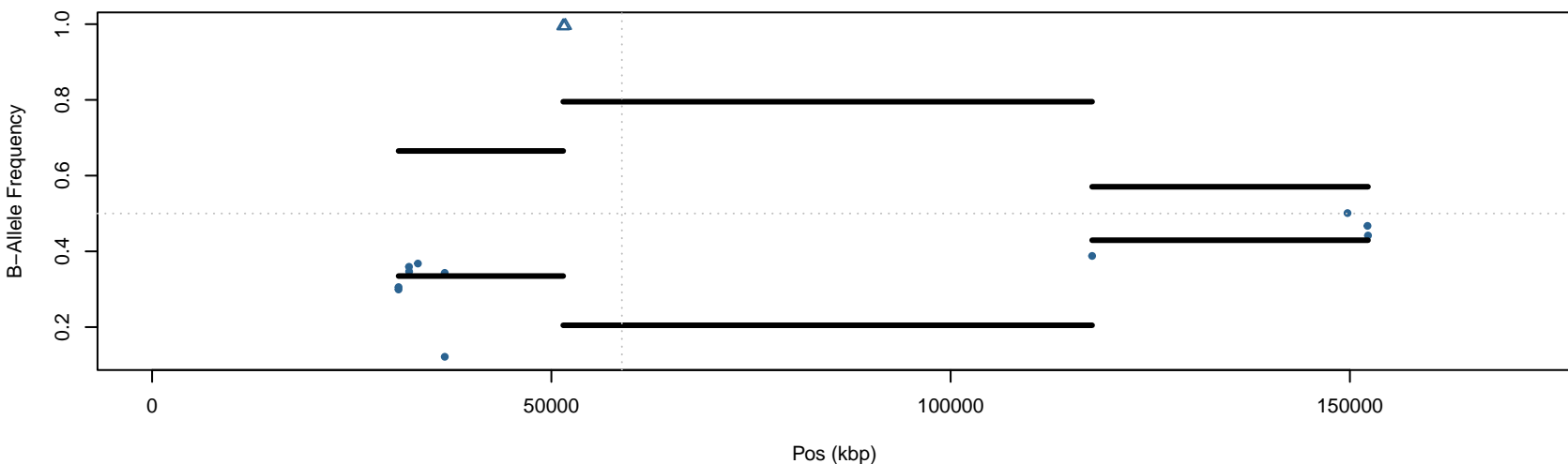
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 4



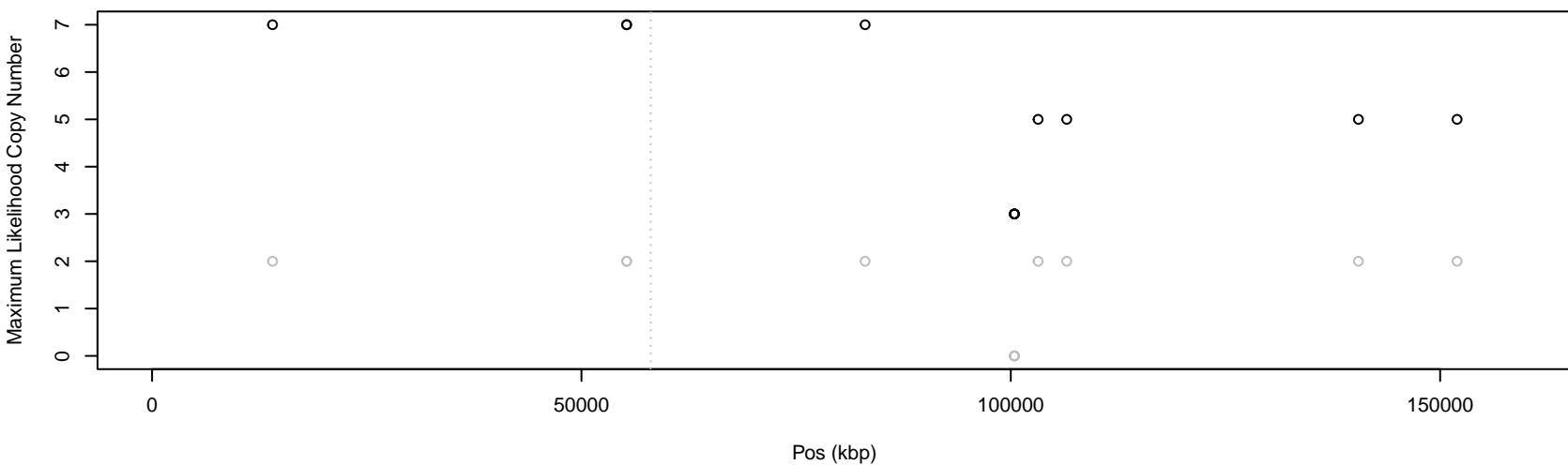
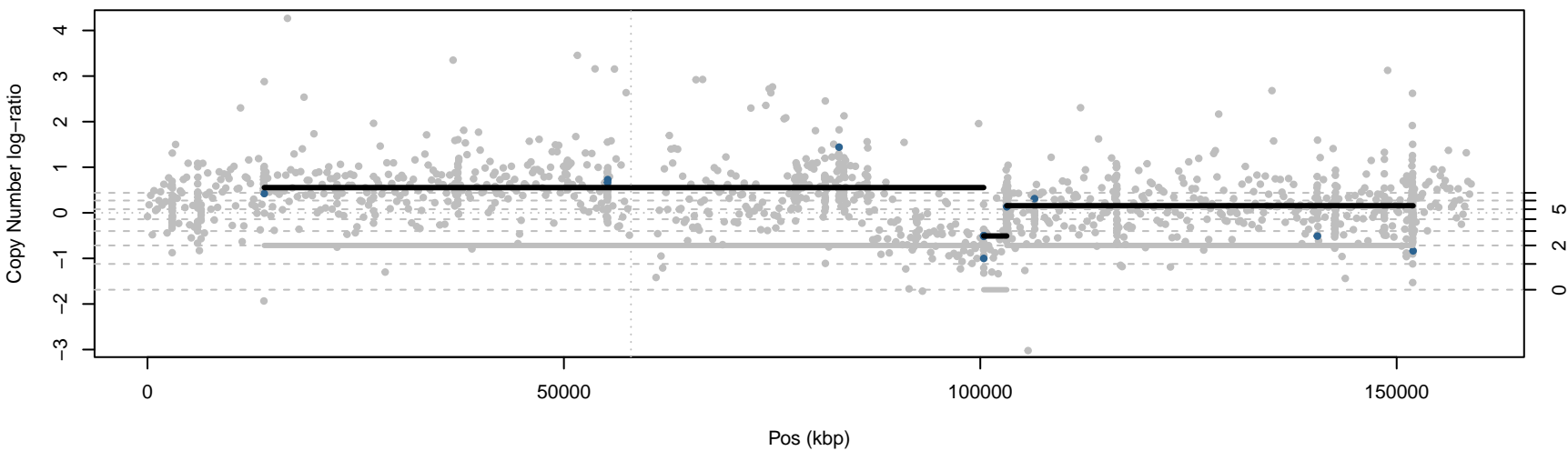
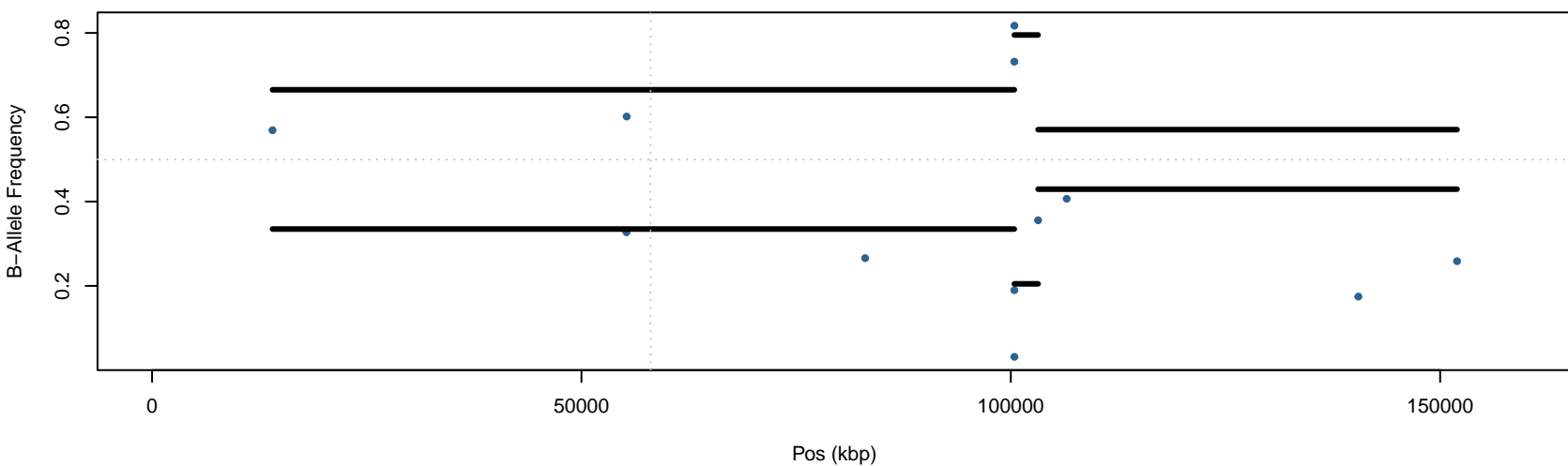
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 5



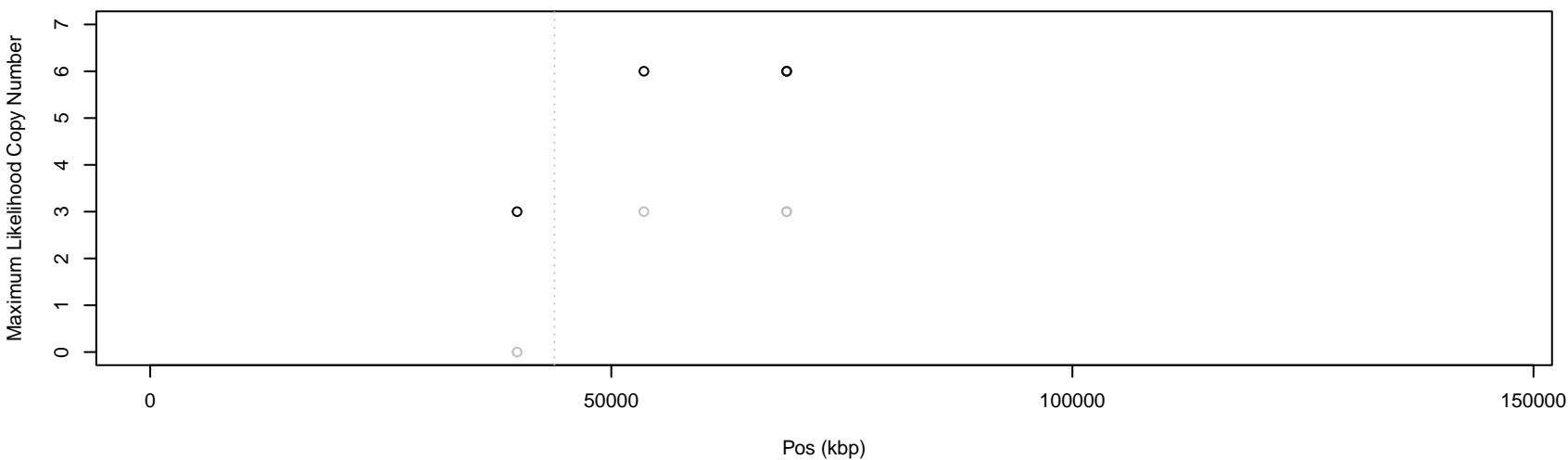
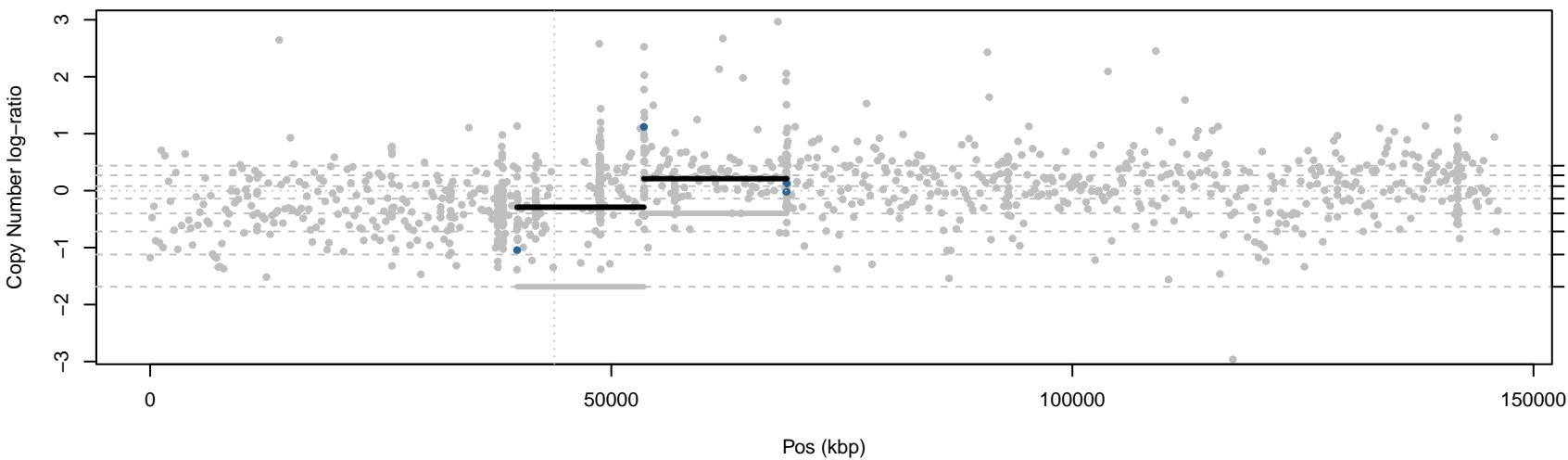
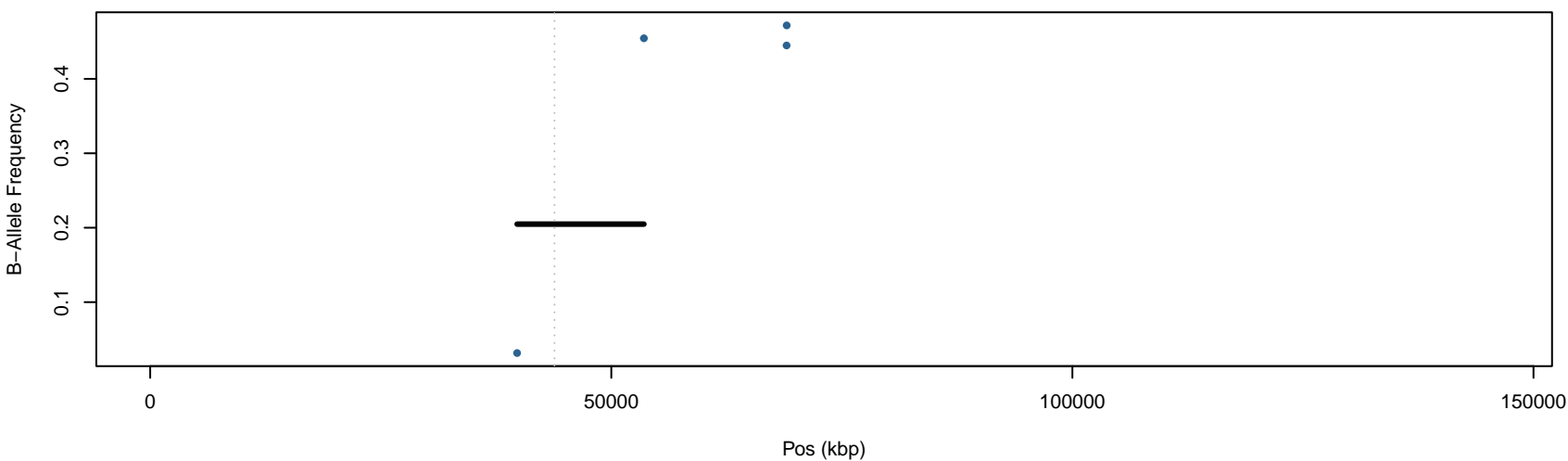
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 6



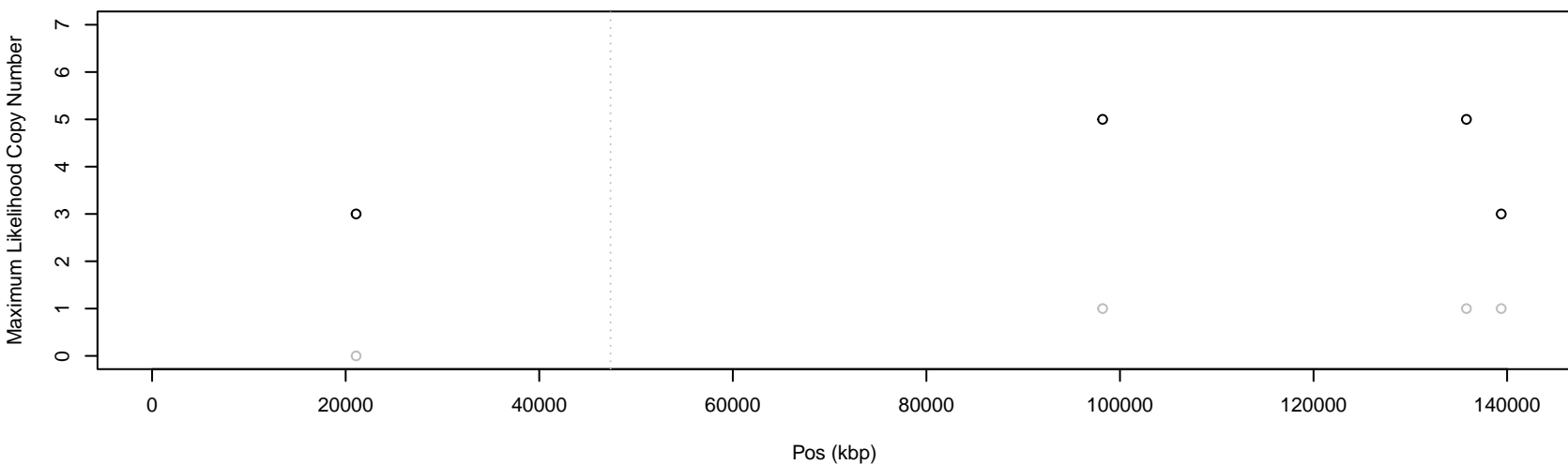
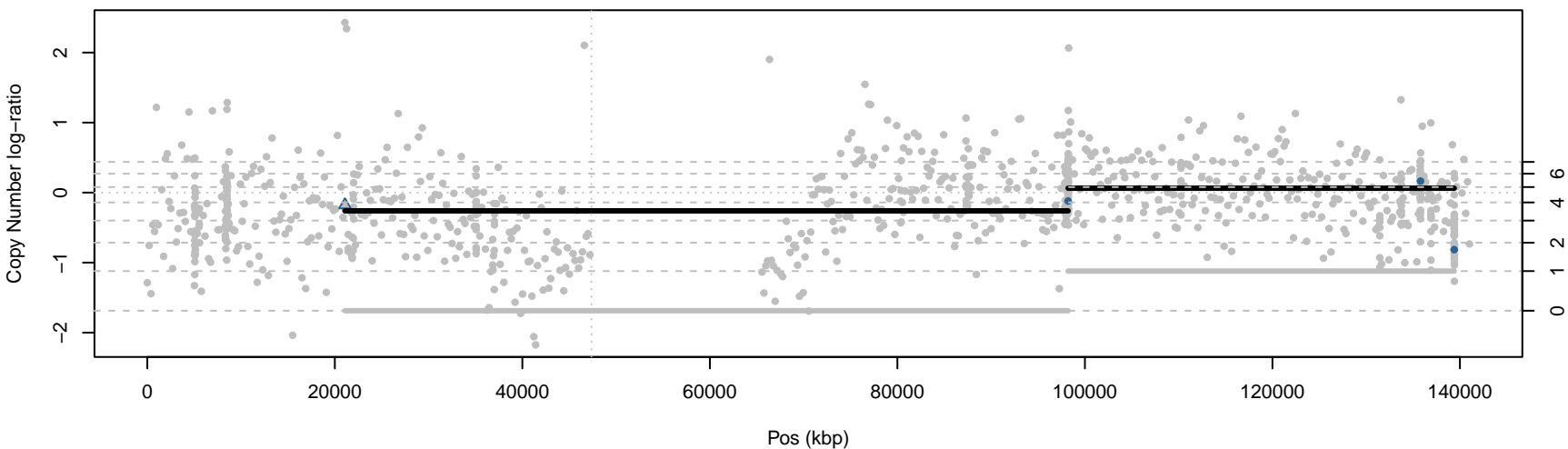
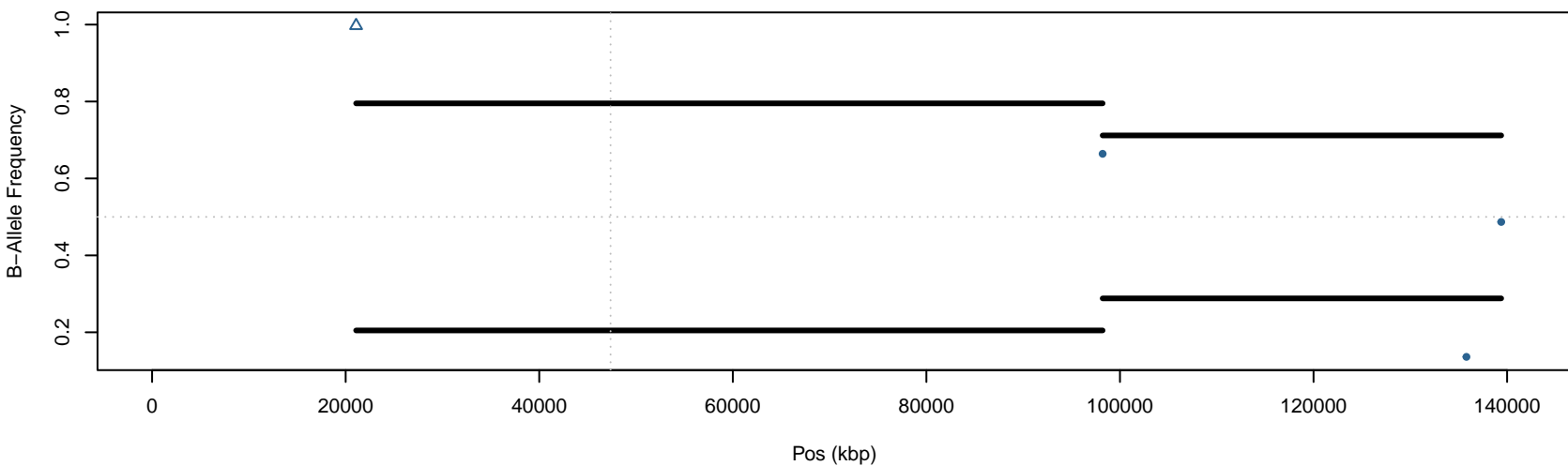
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 7



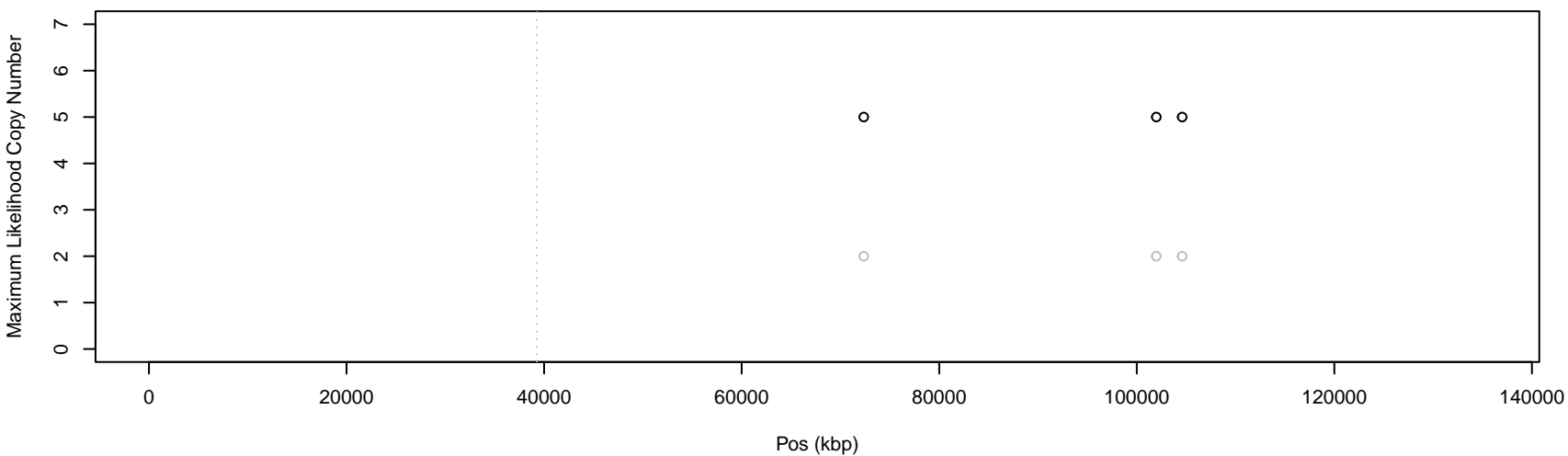
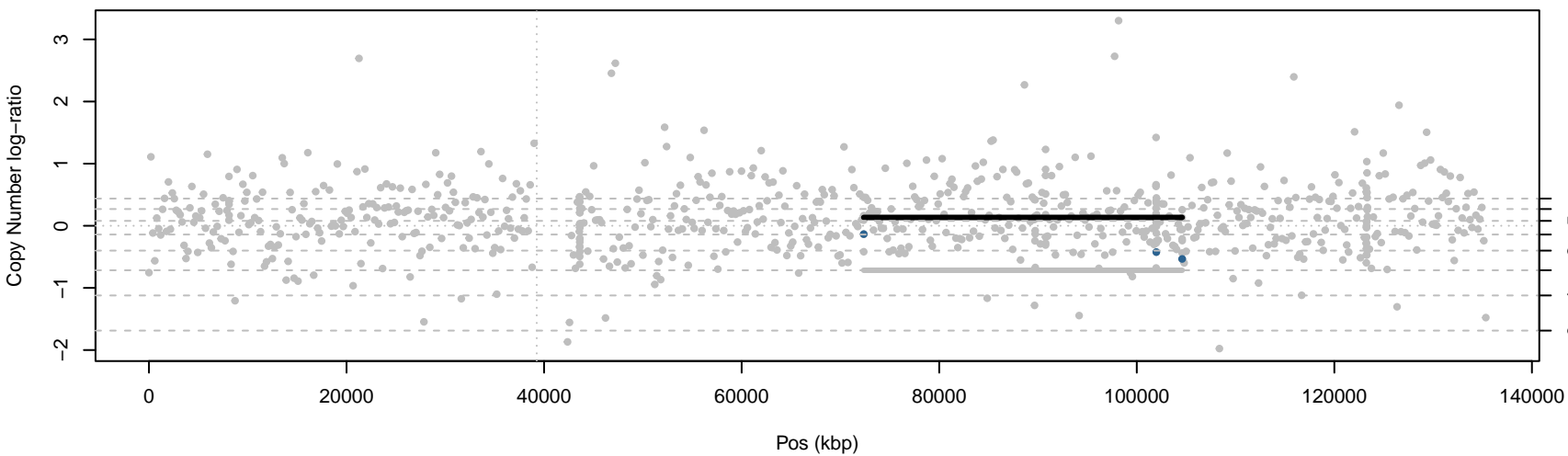
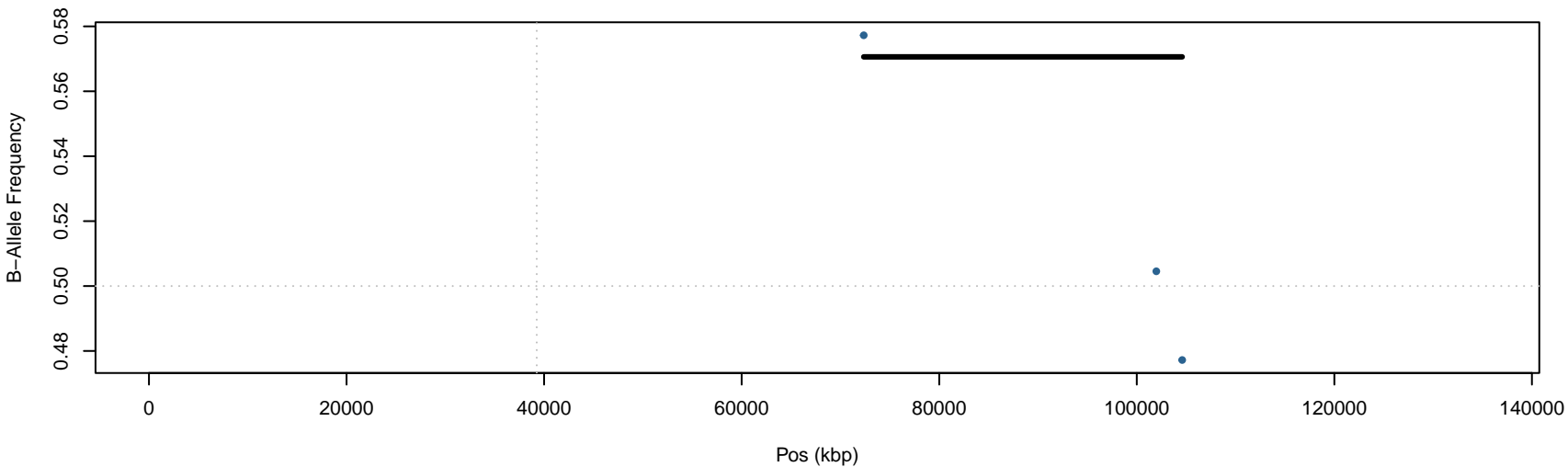
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 8



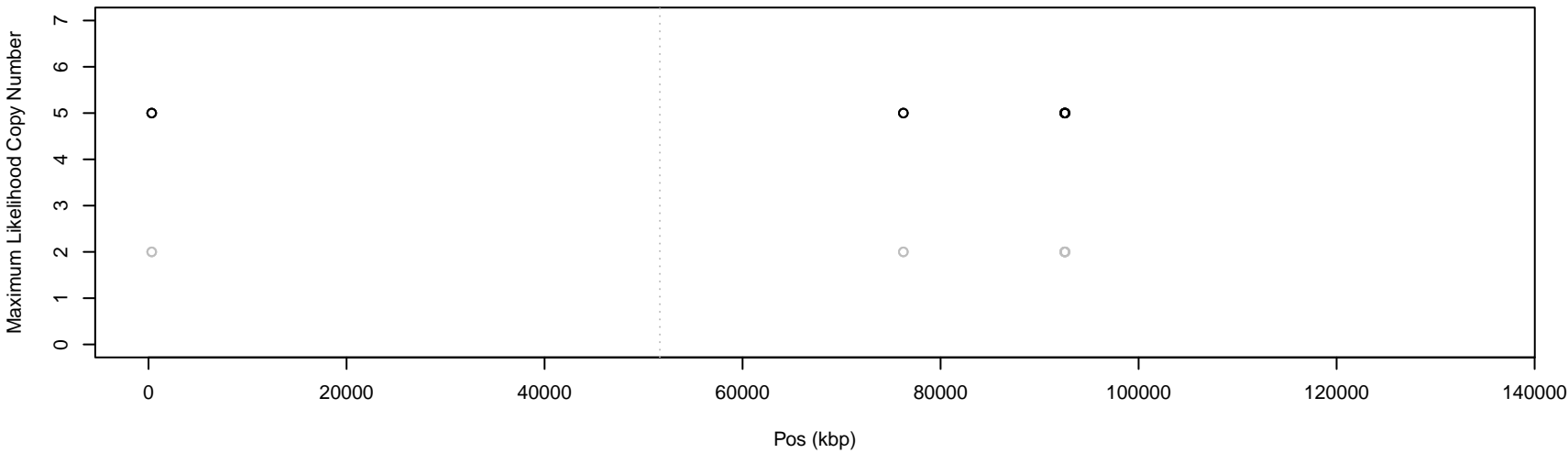
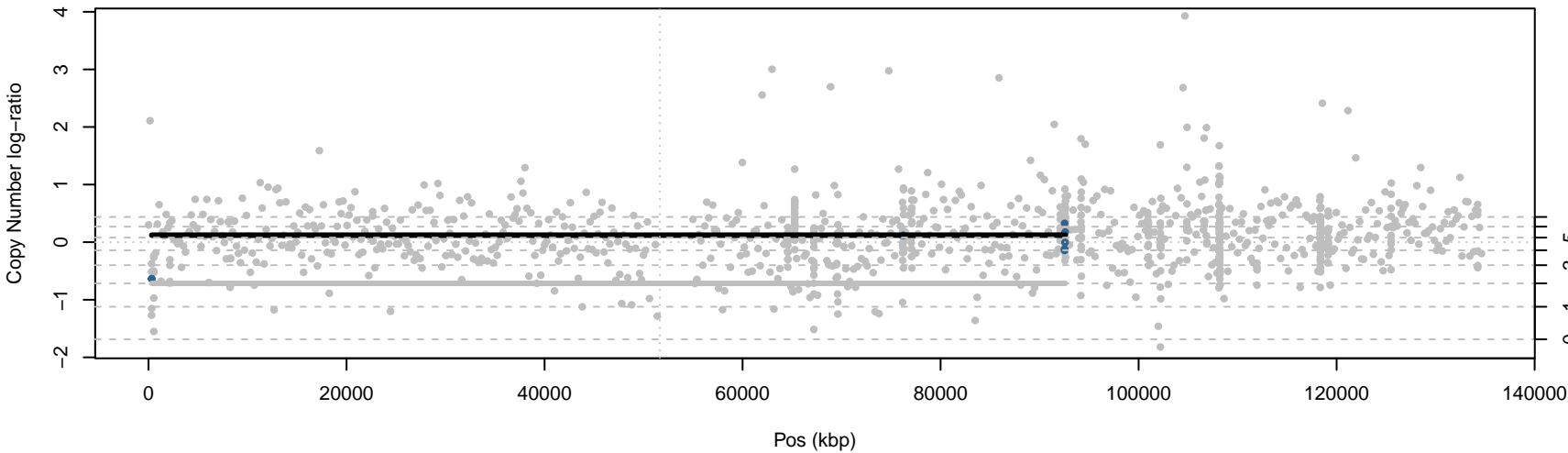
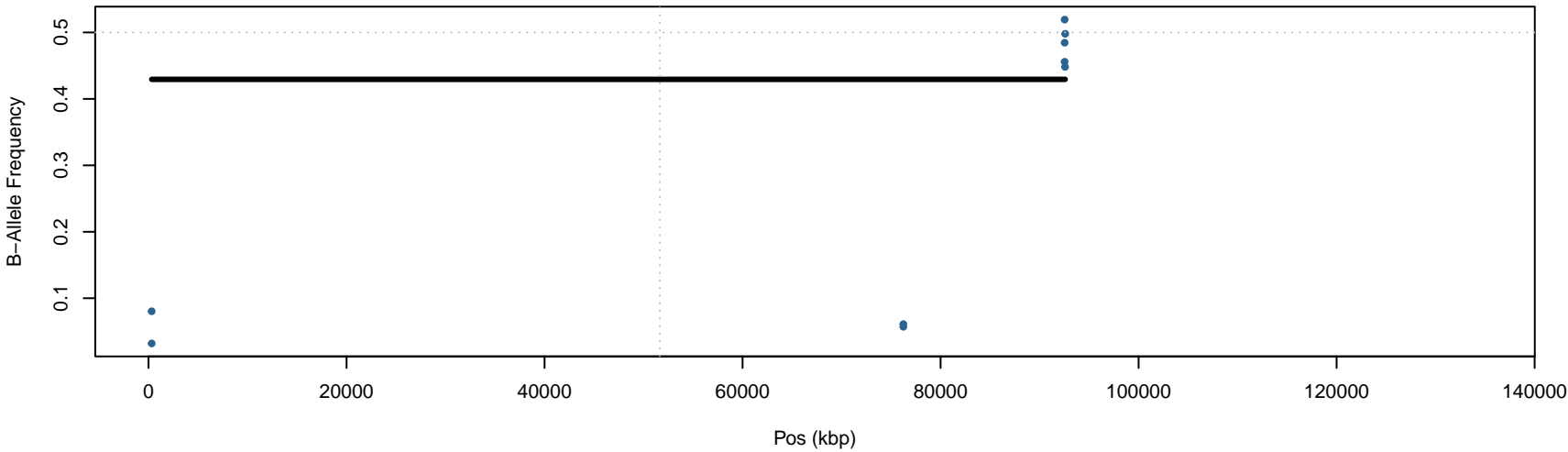
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 9



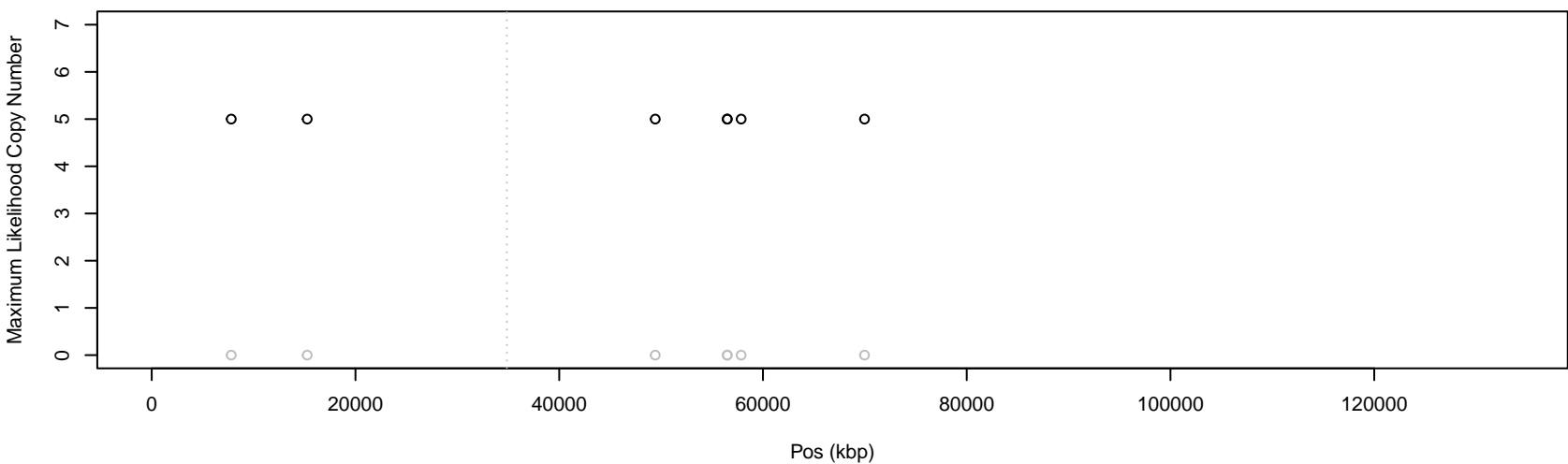
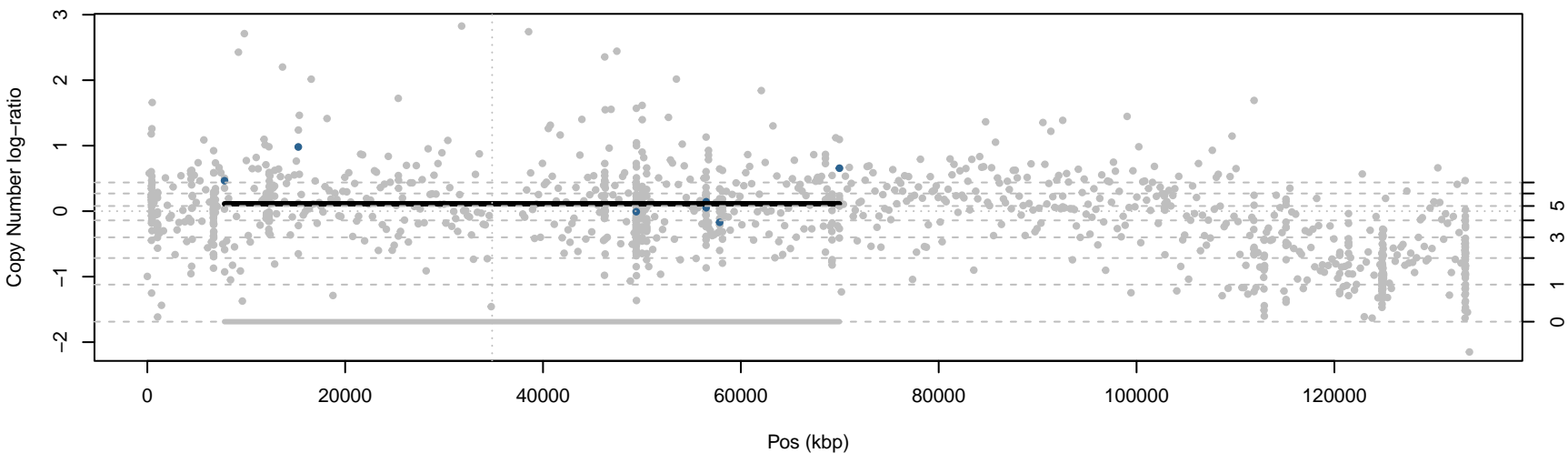
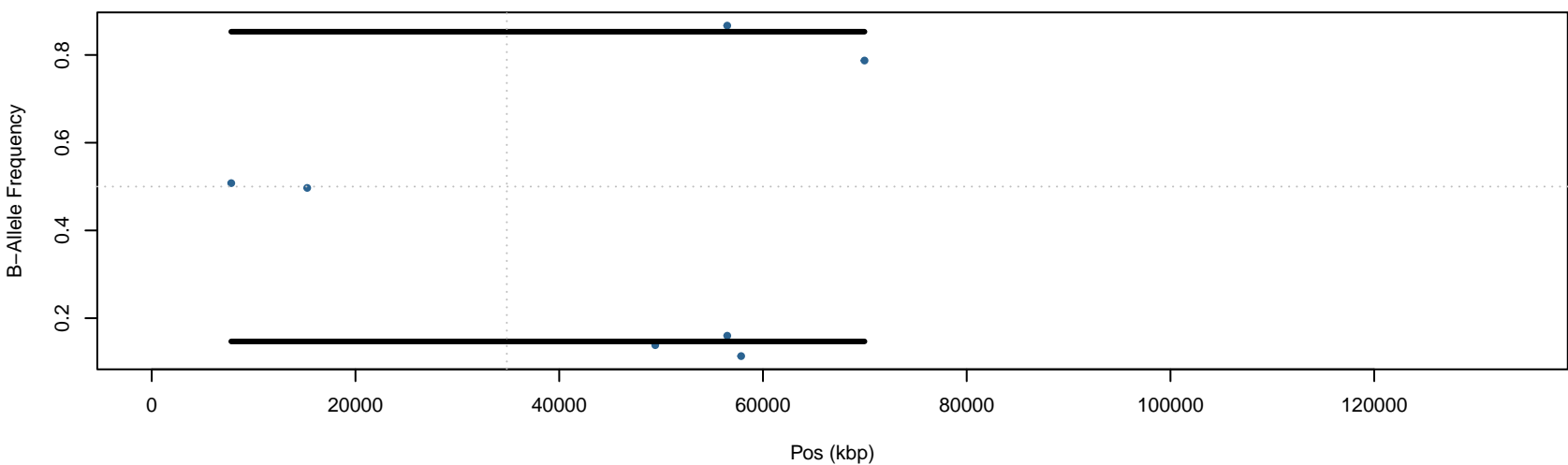
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 10



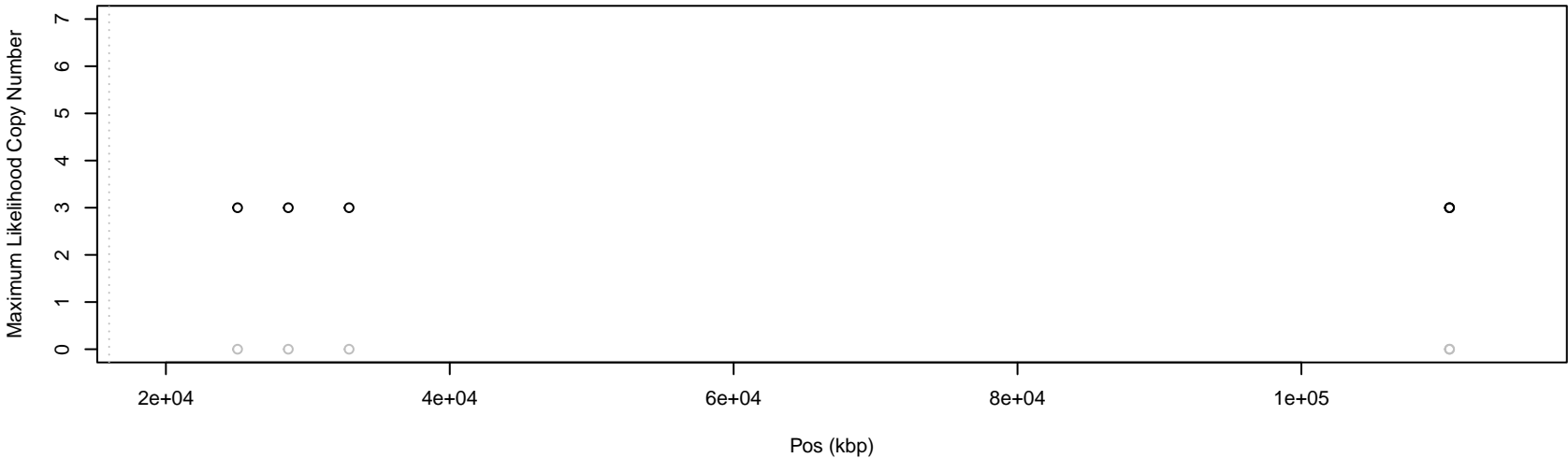
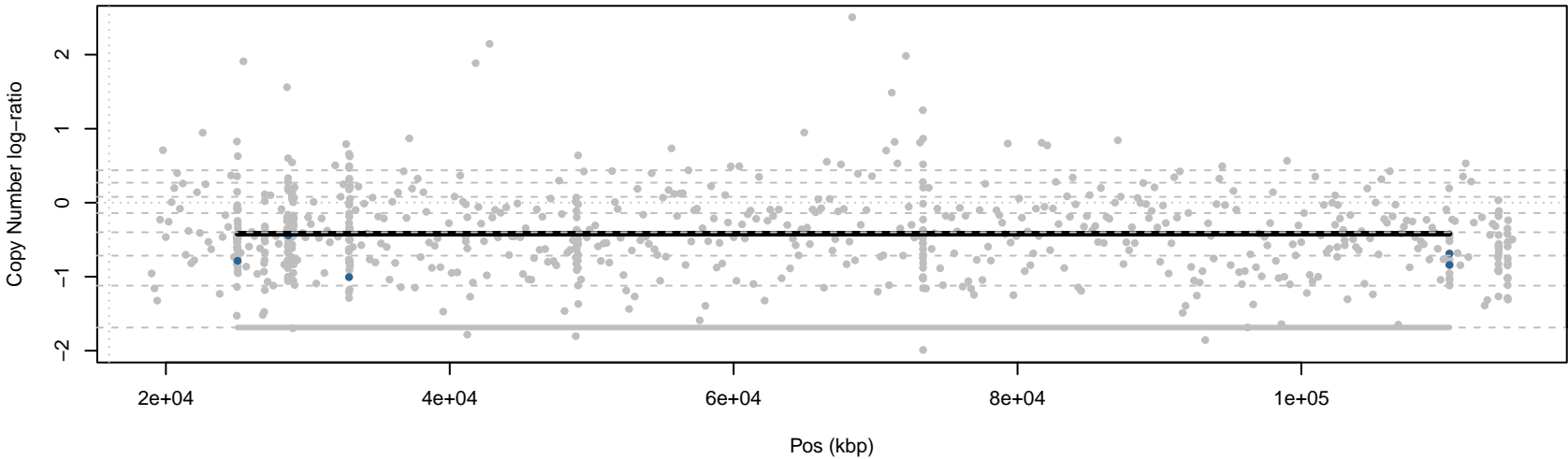
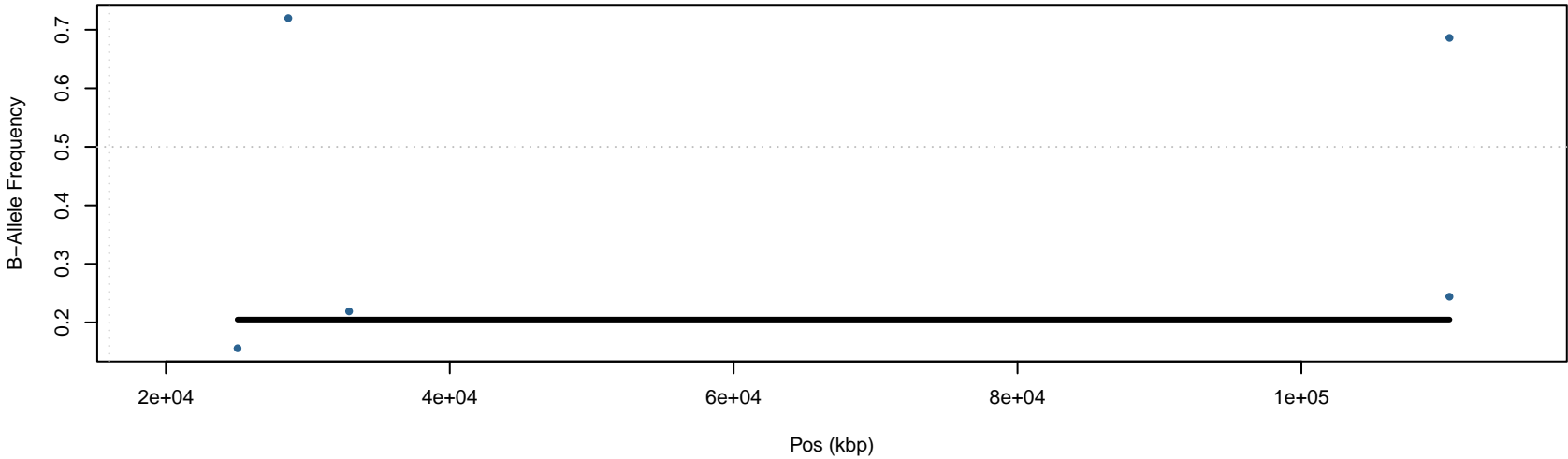
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 11



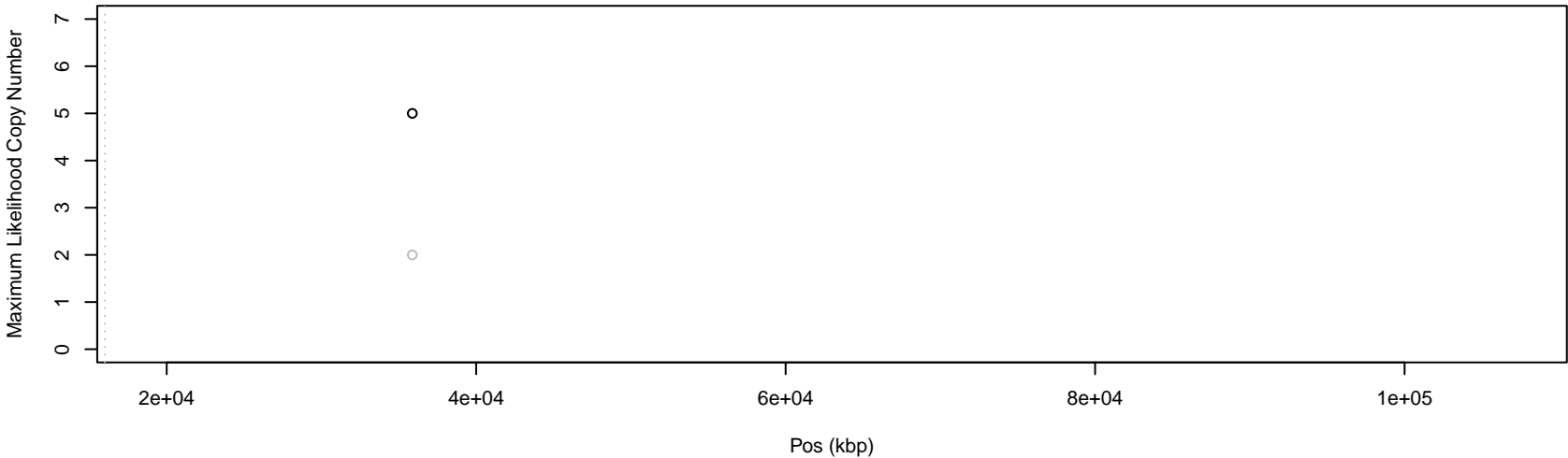
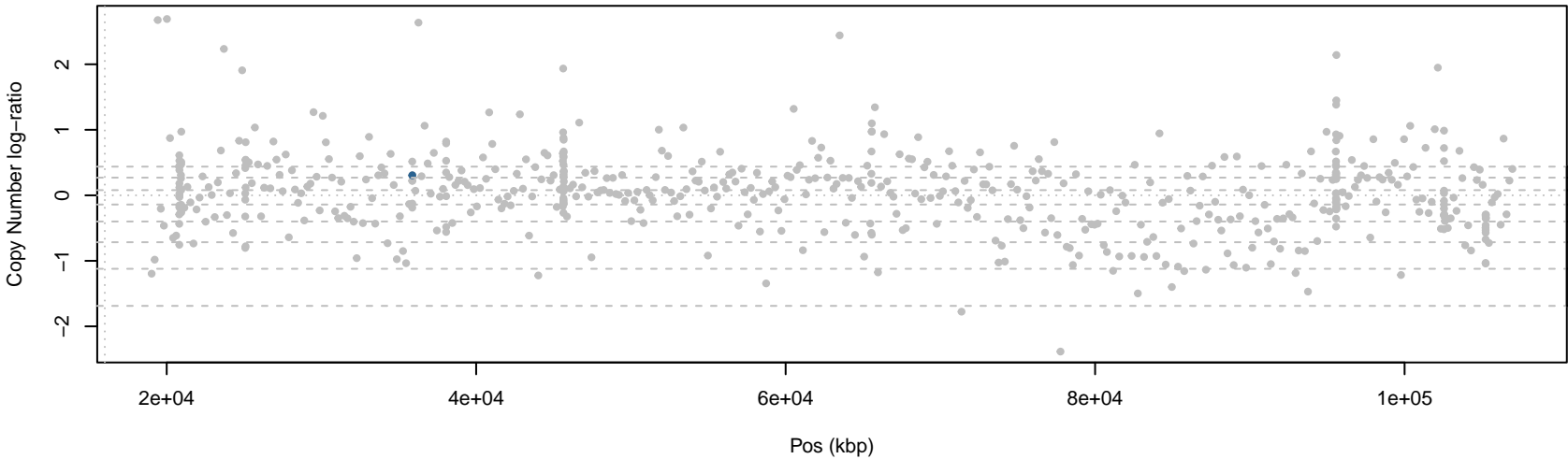
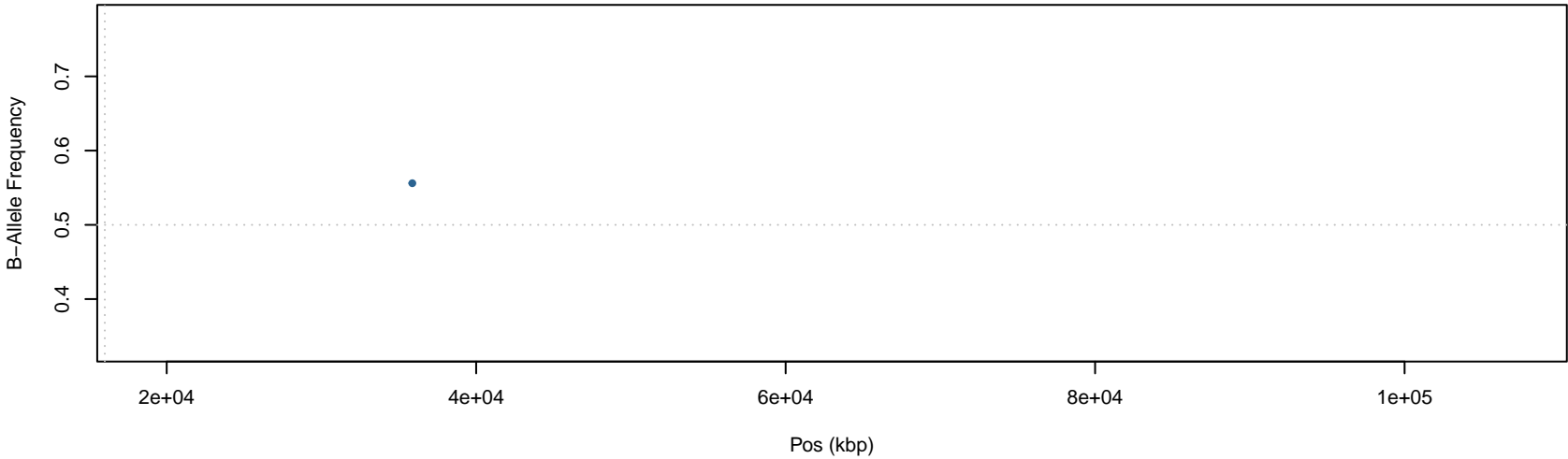
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 12



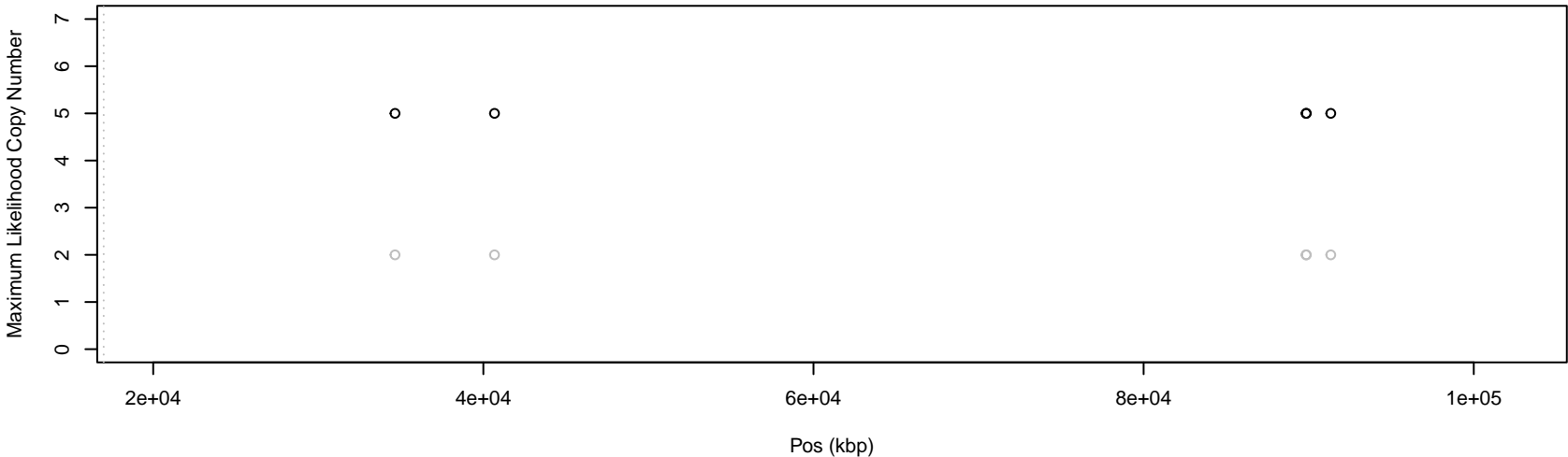
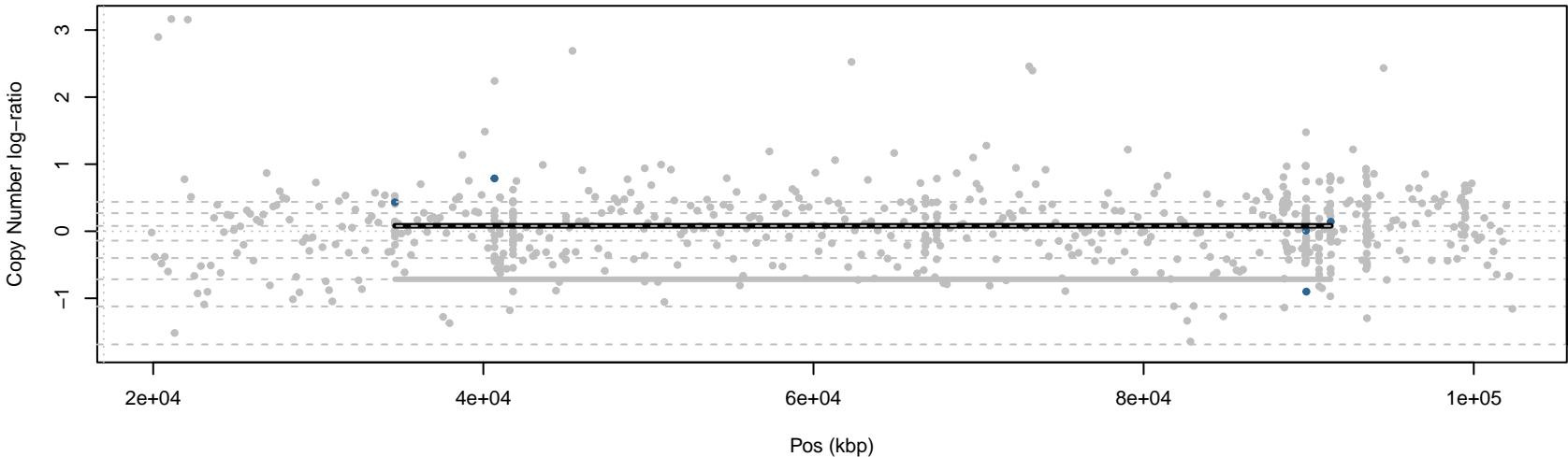
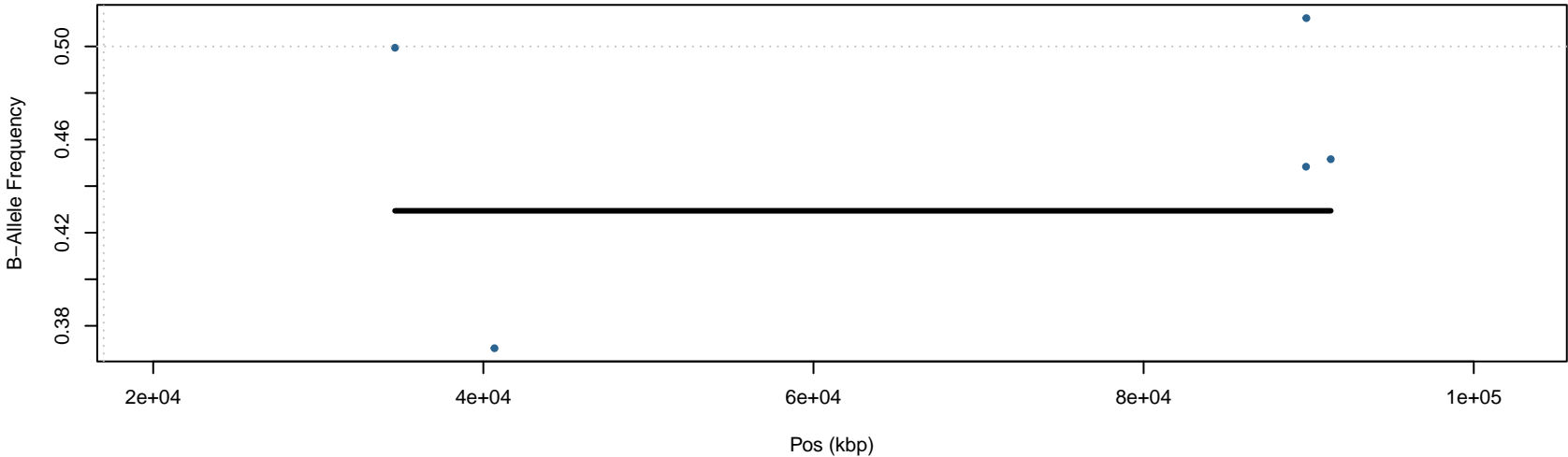
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 13



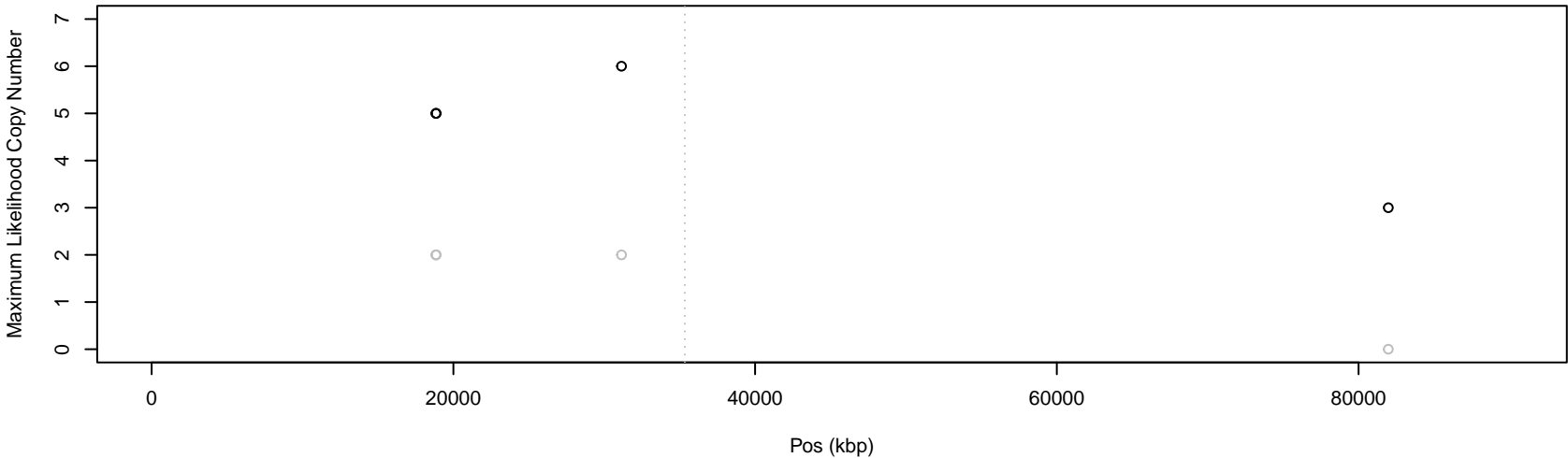
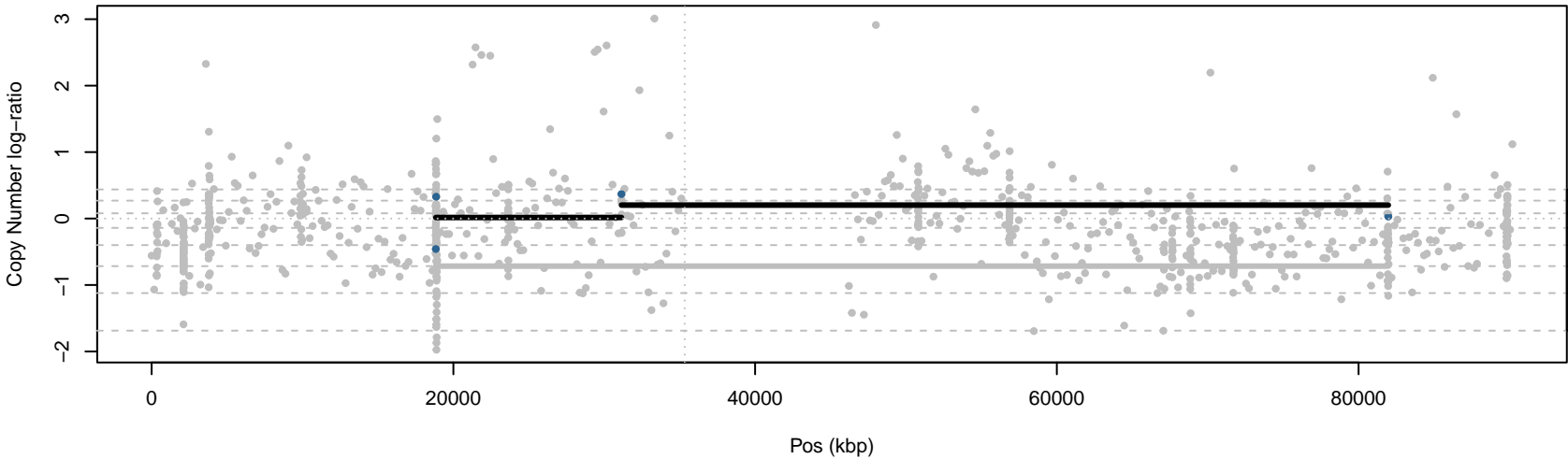
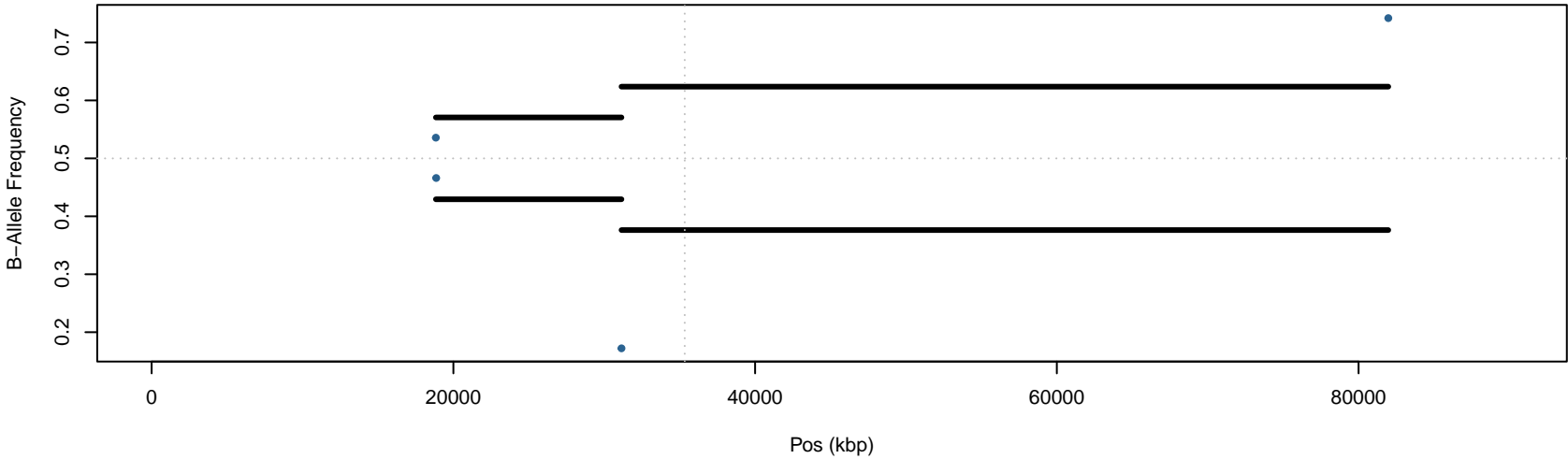
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 14



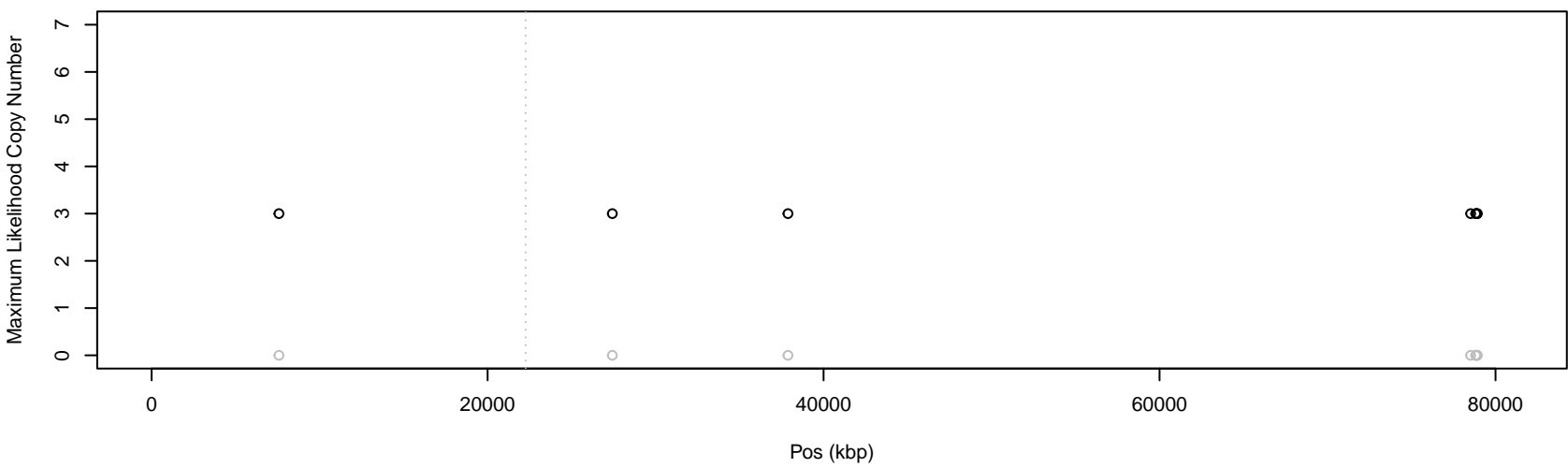
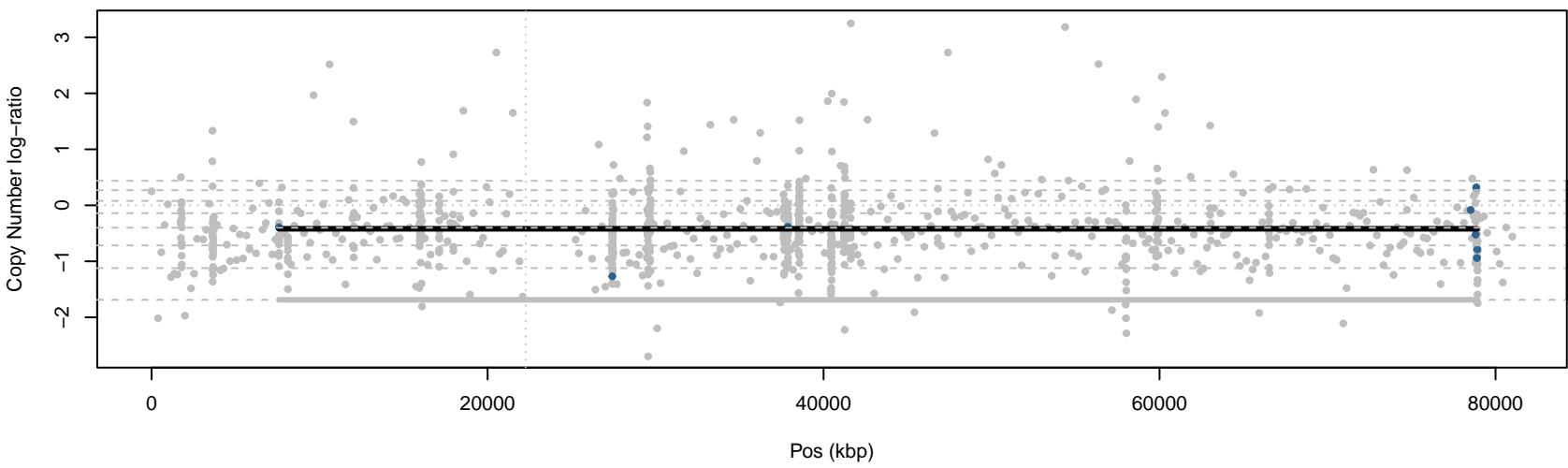
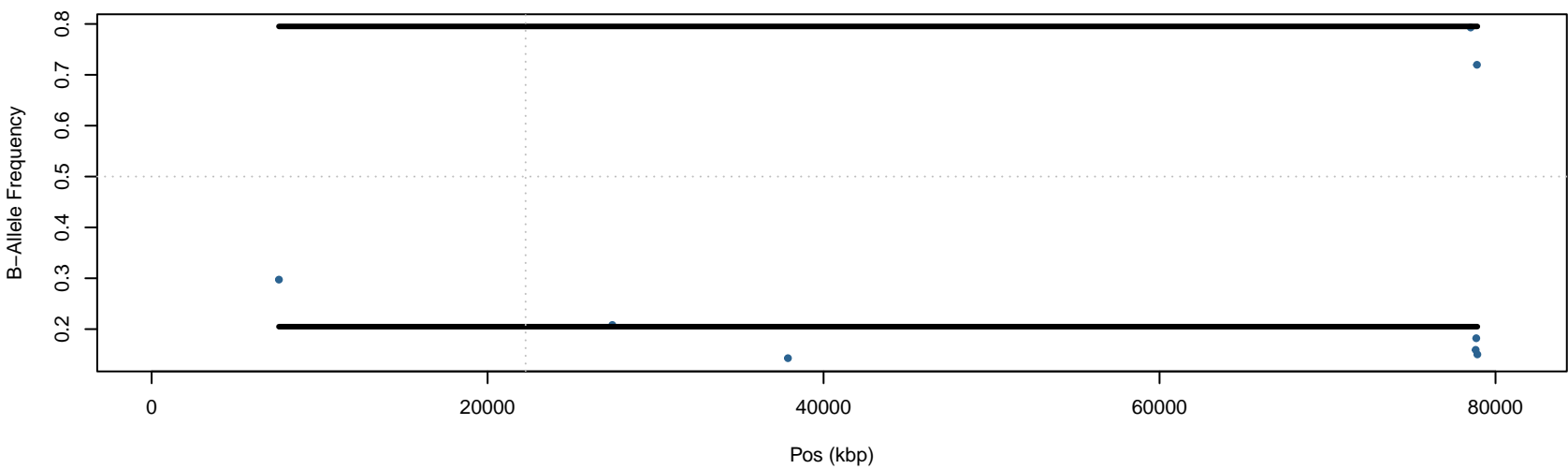
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 15



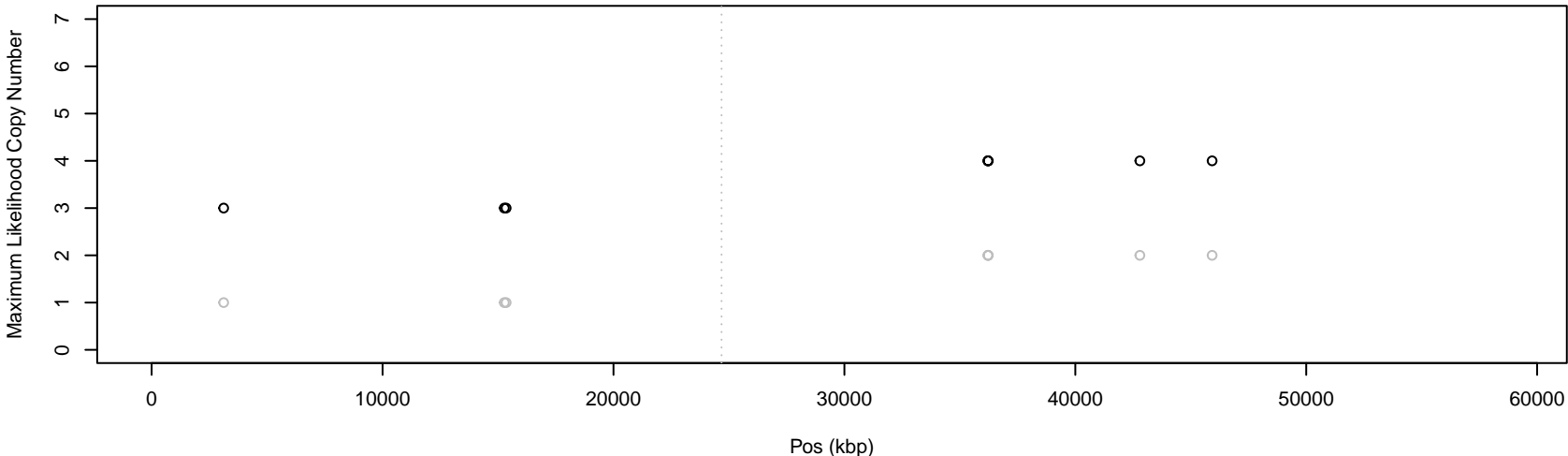
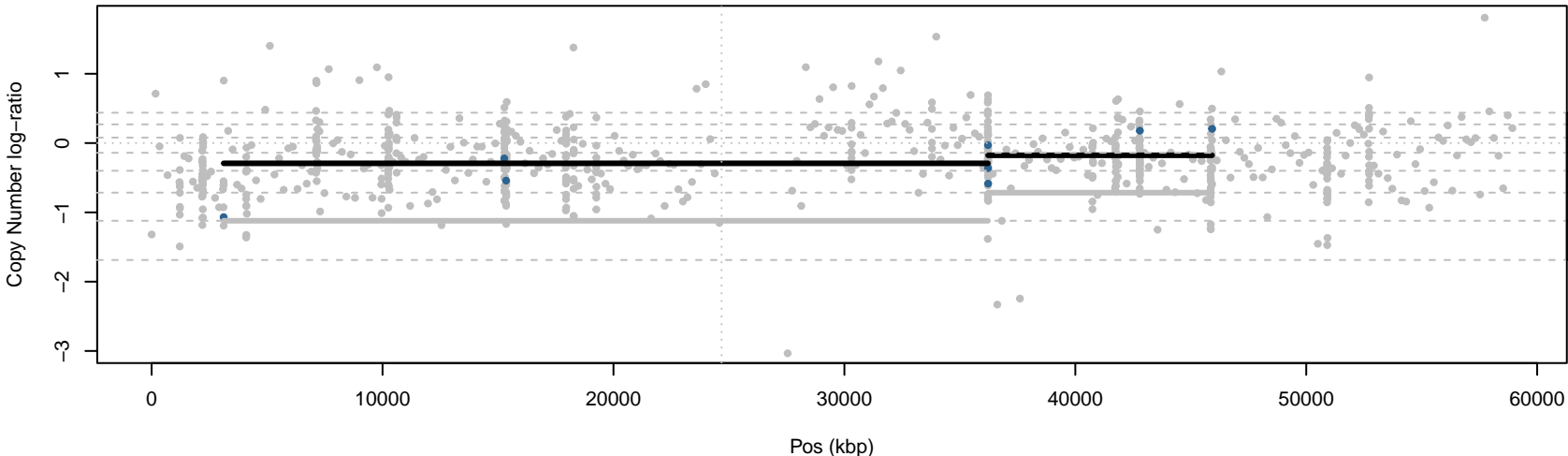
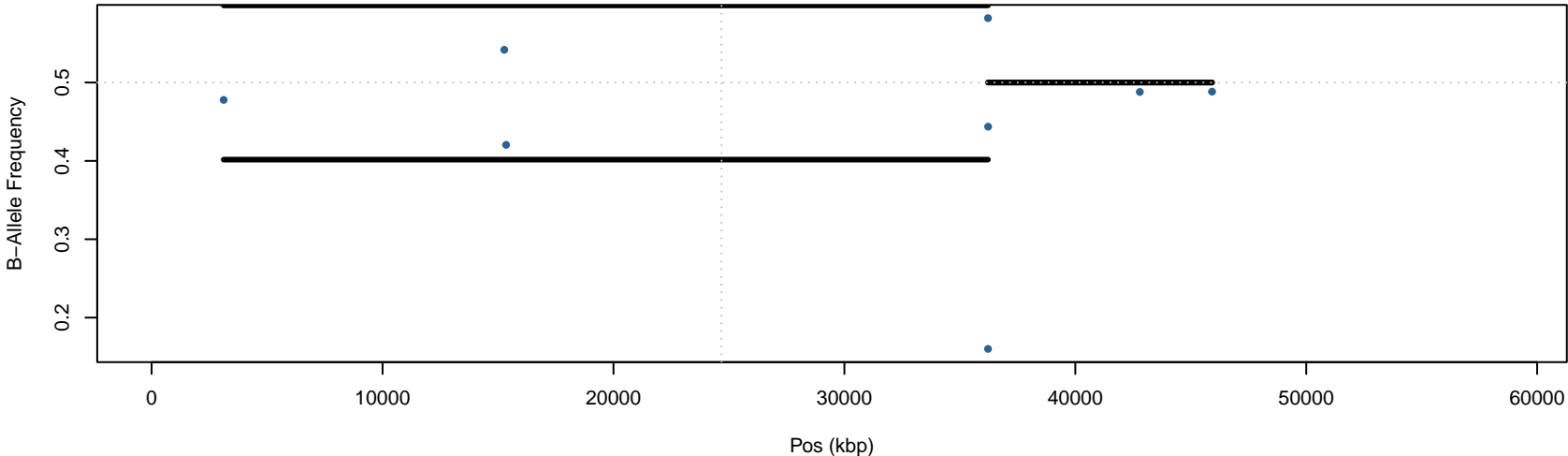
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 16



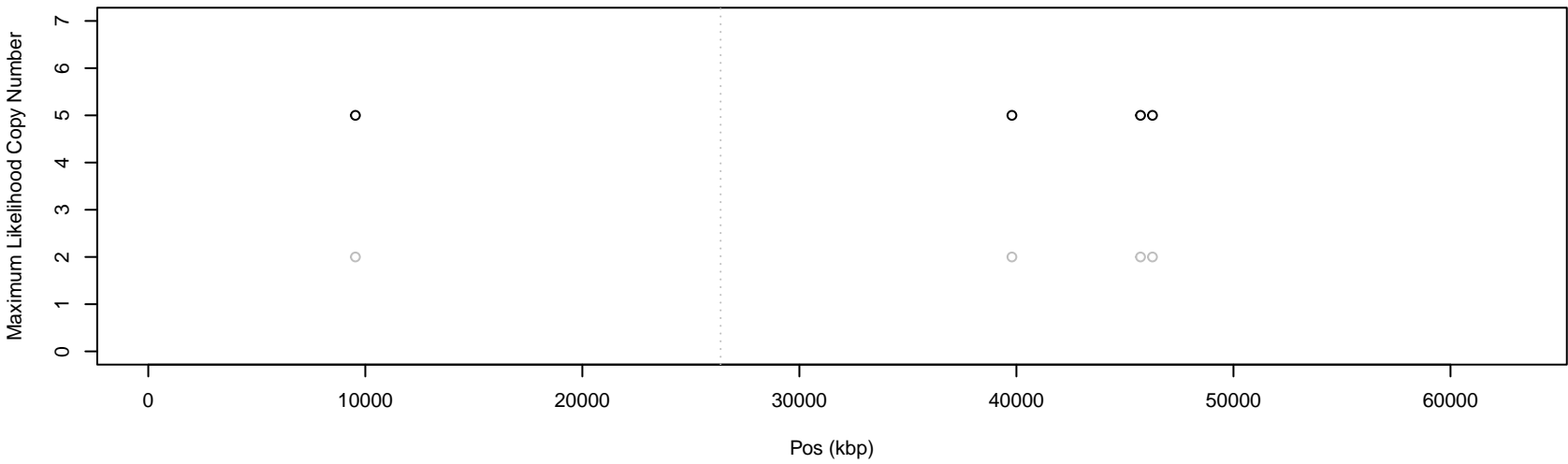
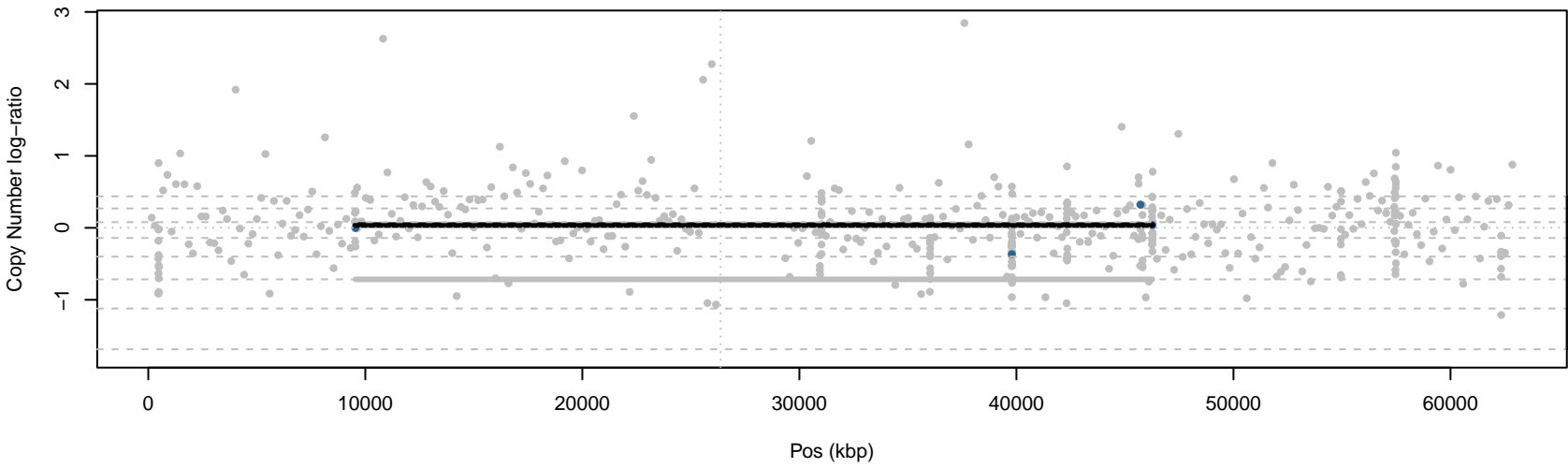
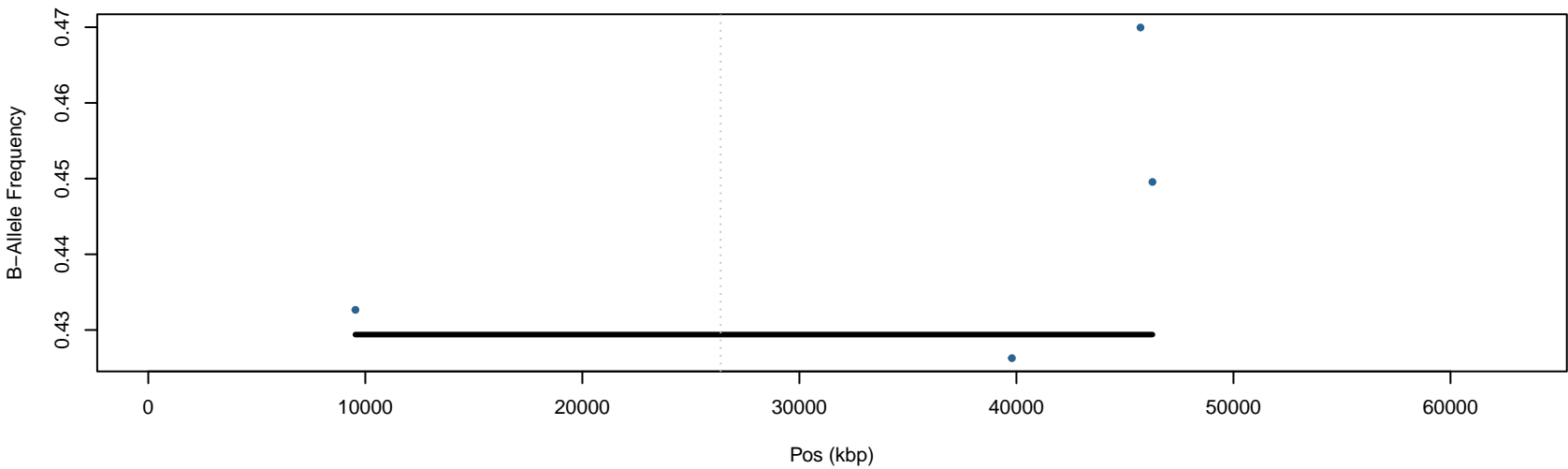
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 17



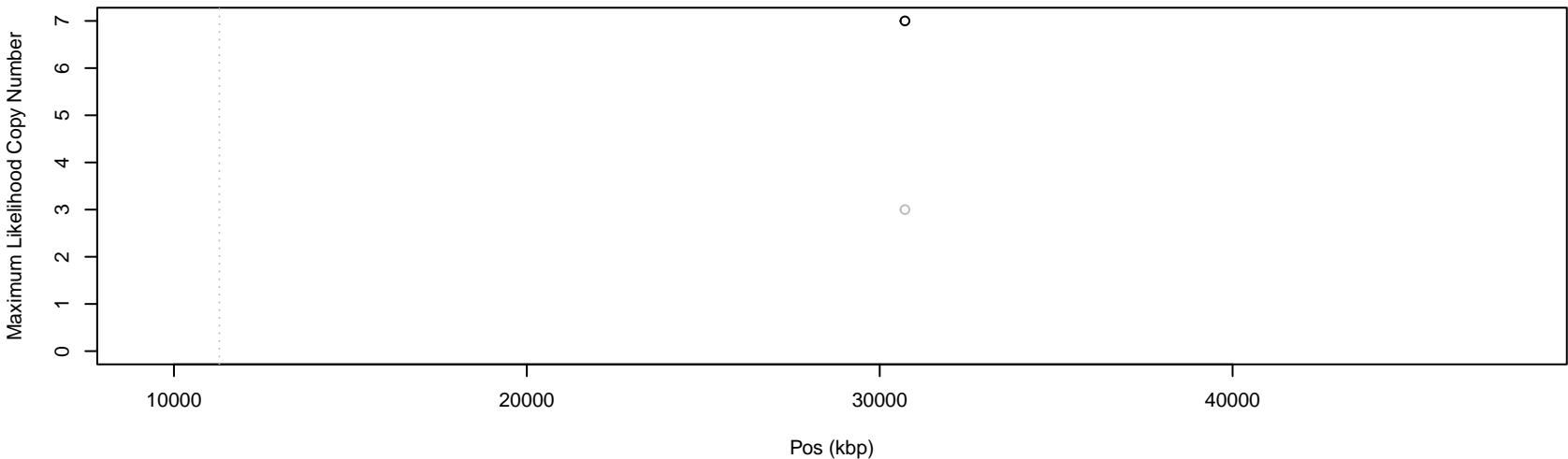
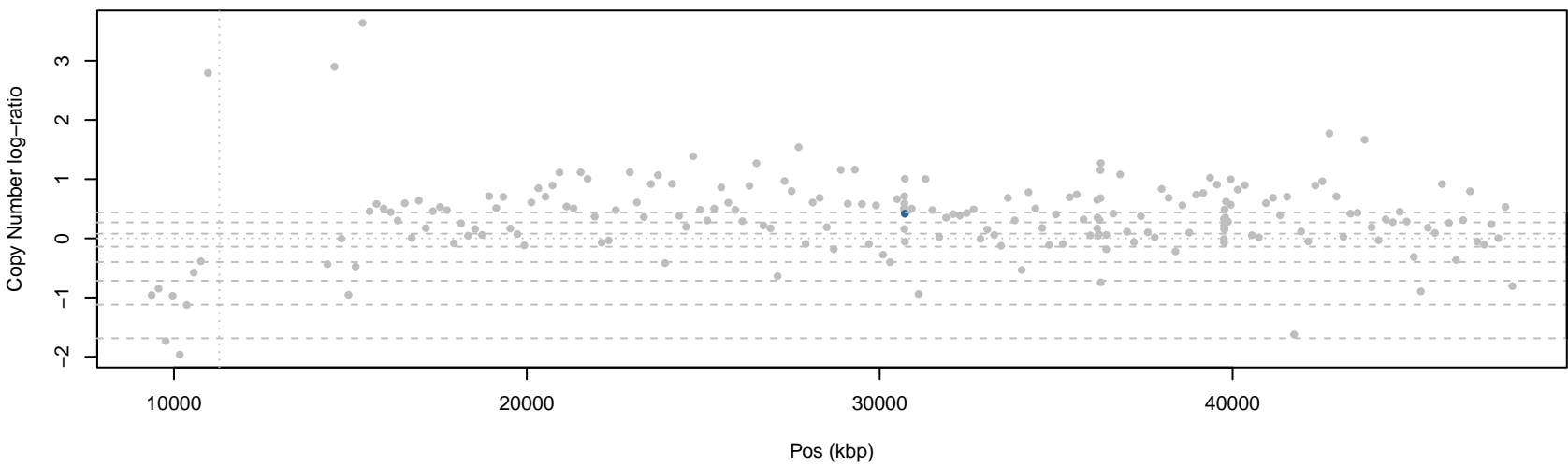
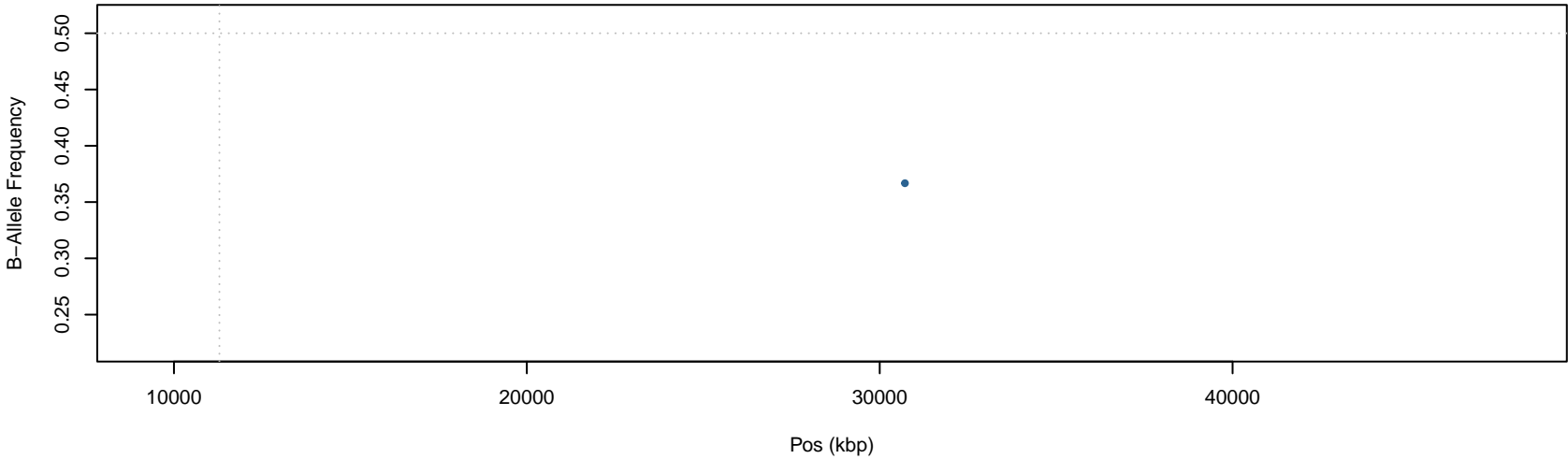
Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 19



Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 20



Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 21



Purity: 0.49 Tumor ploidy: 4.623 SNV log-likelihood: -379.67 GoF: 93% Mean coverage: 639;28 Chromosome: 22

