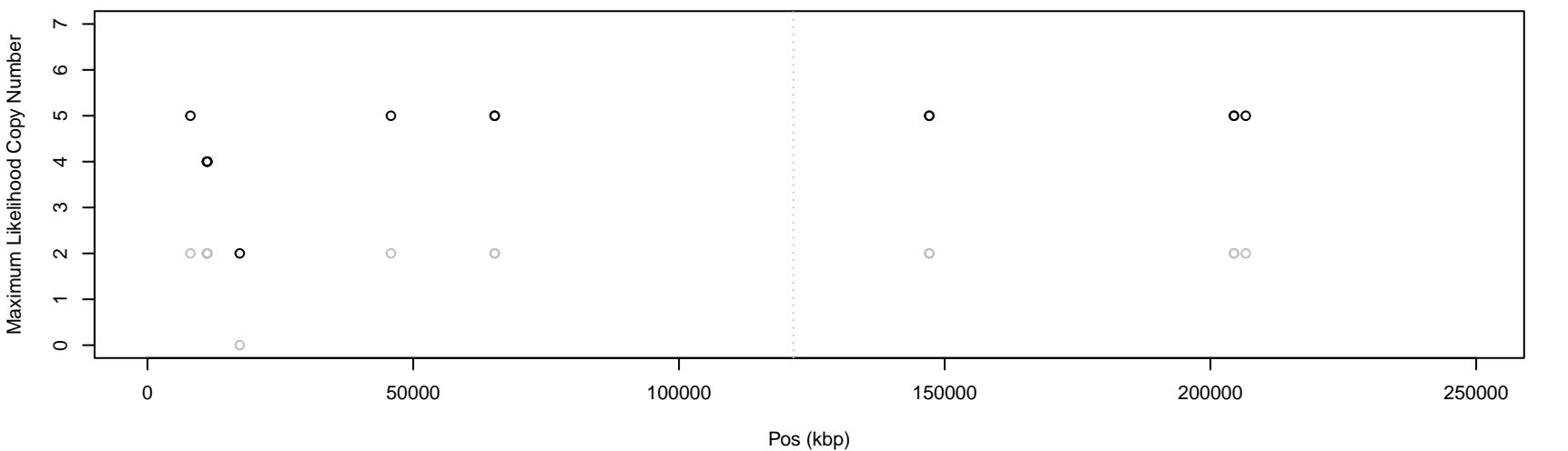
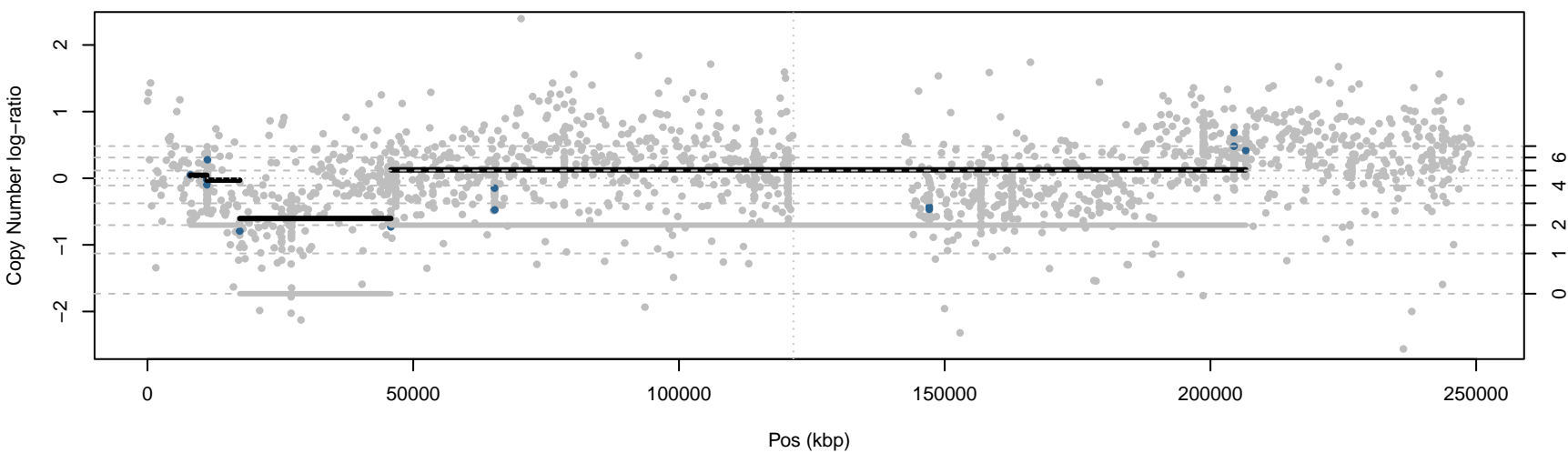
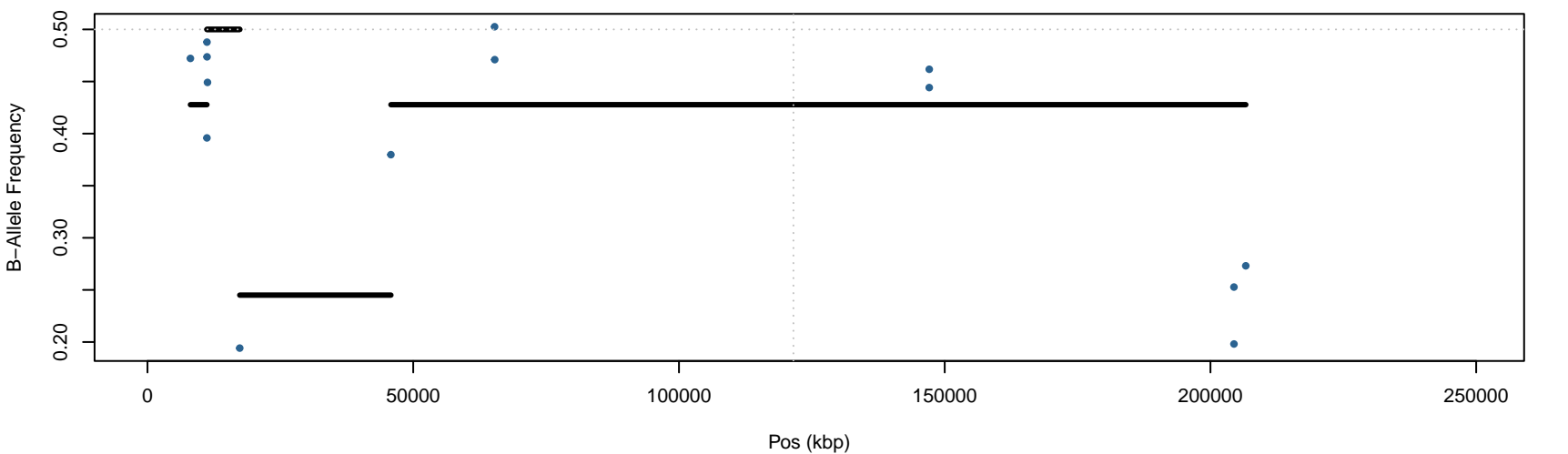
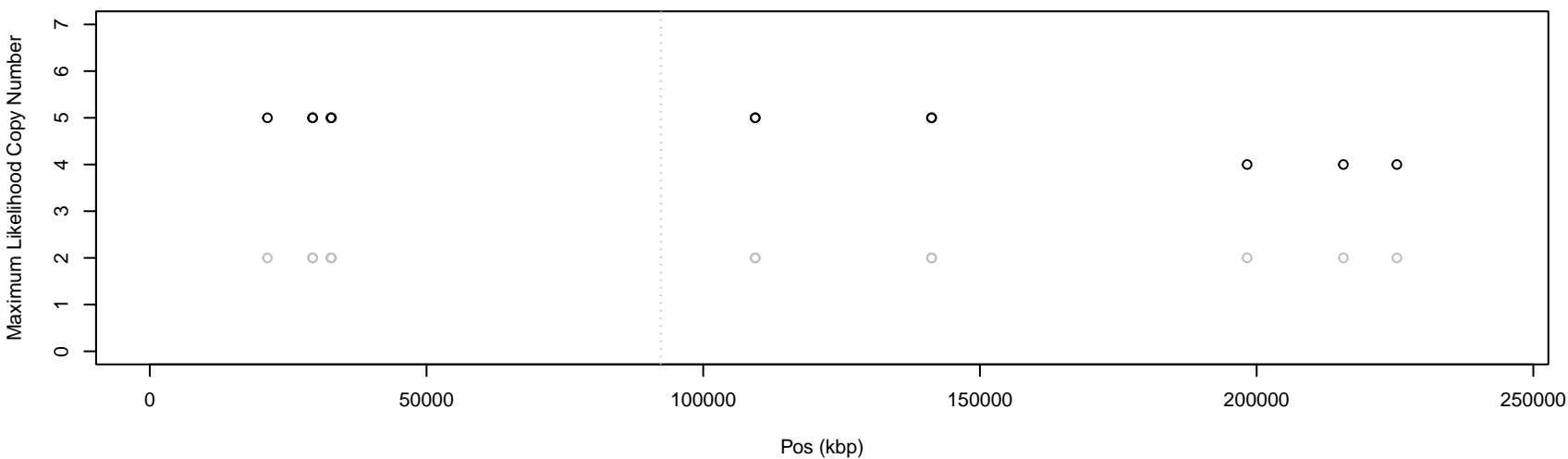
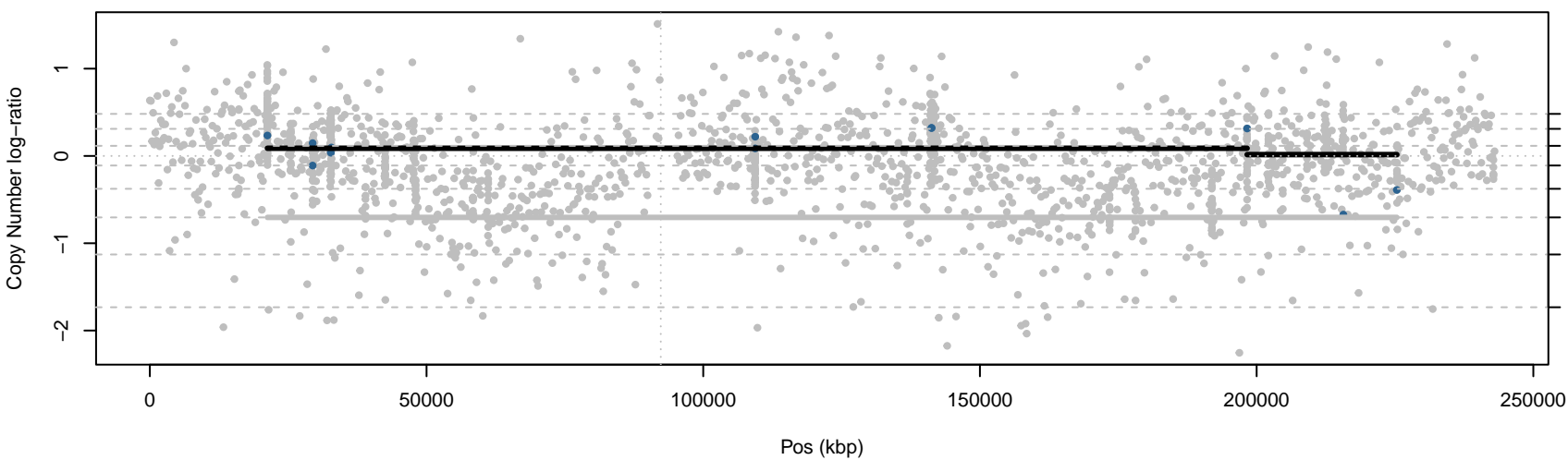
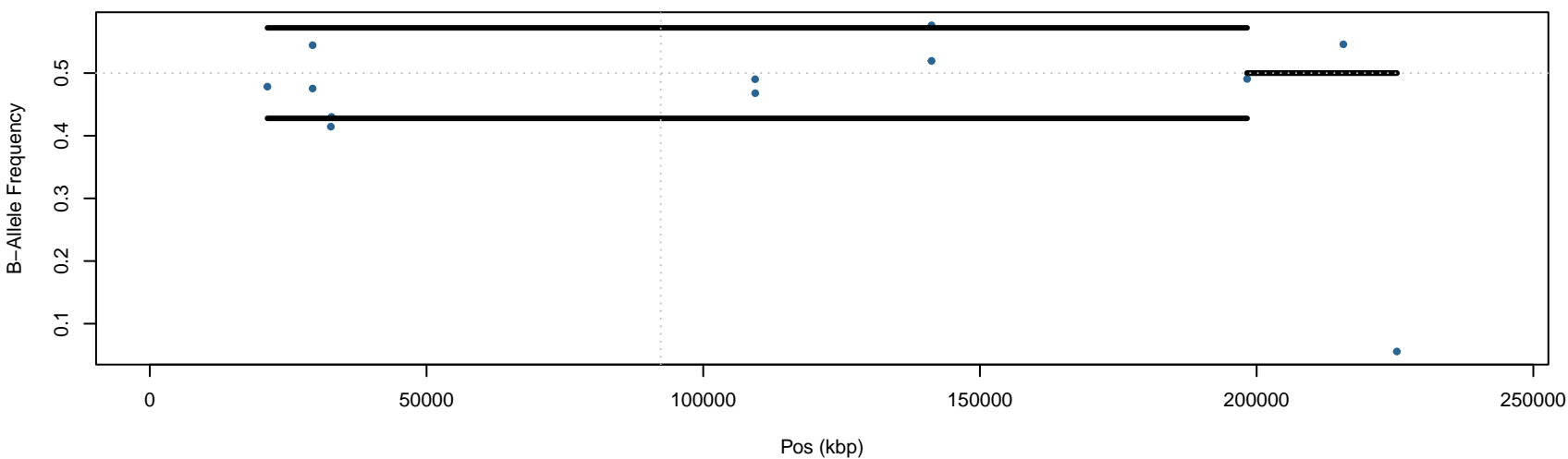


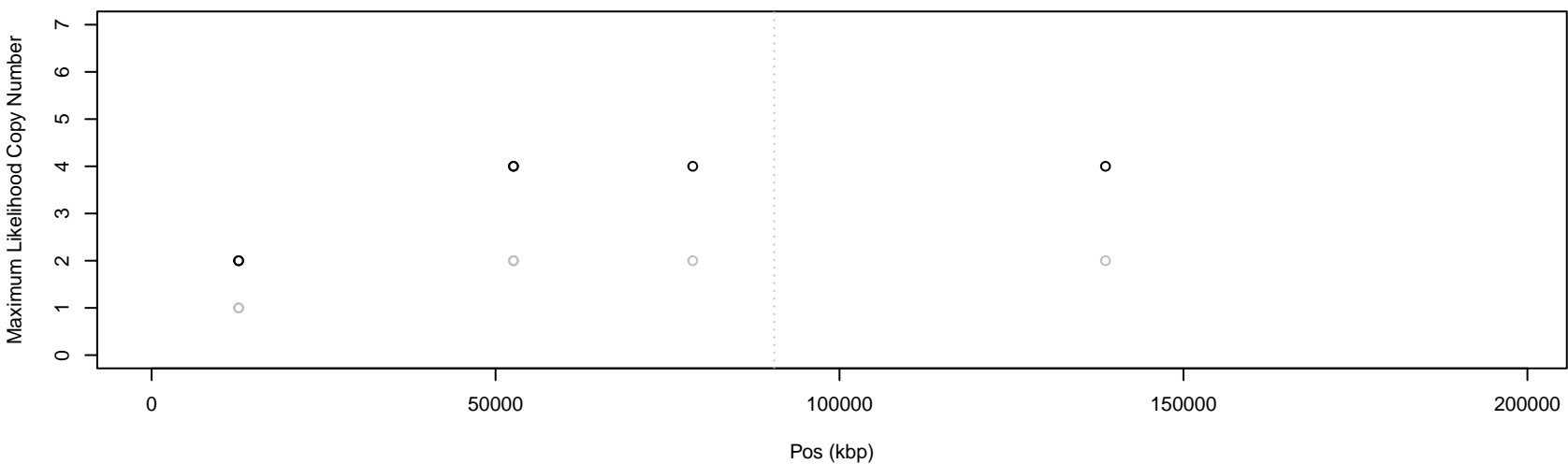
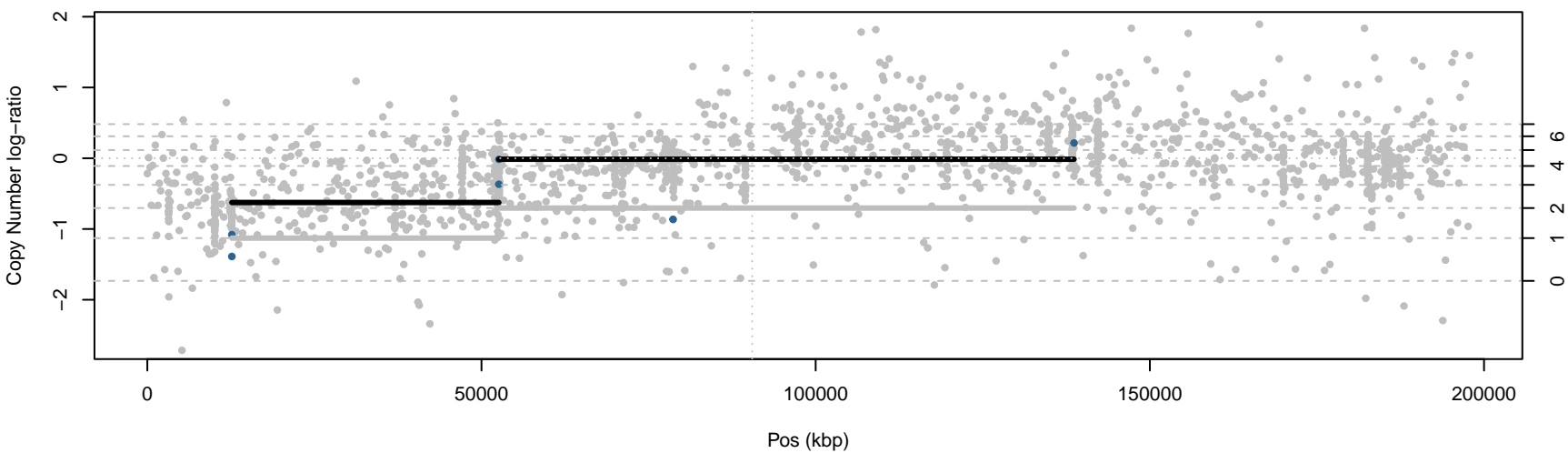
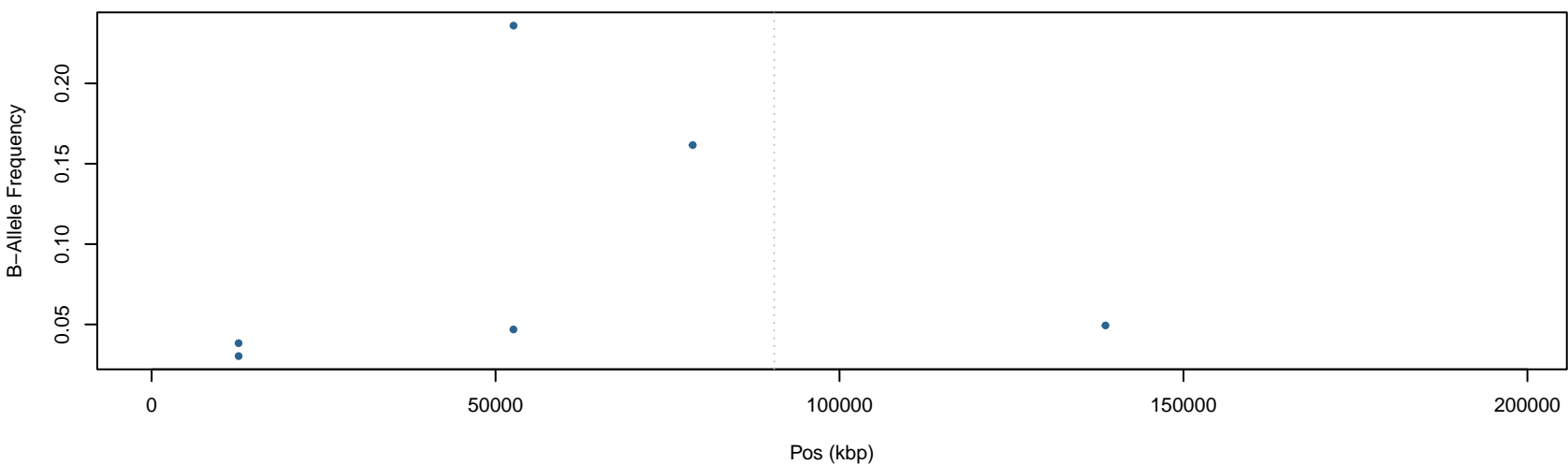
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 1



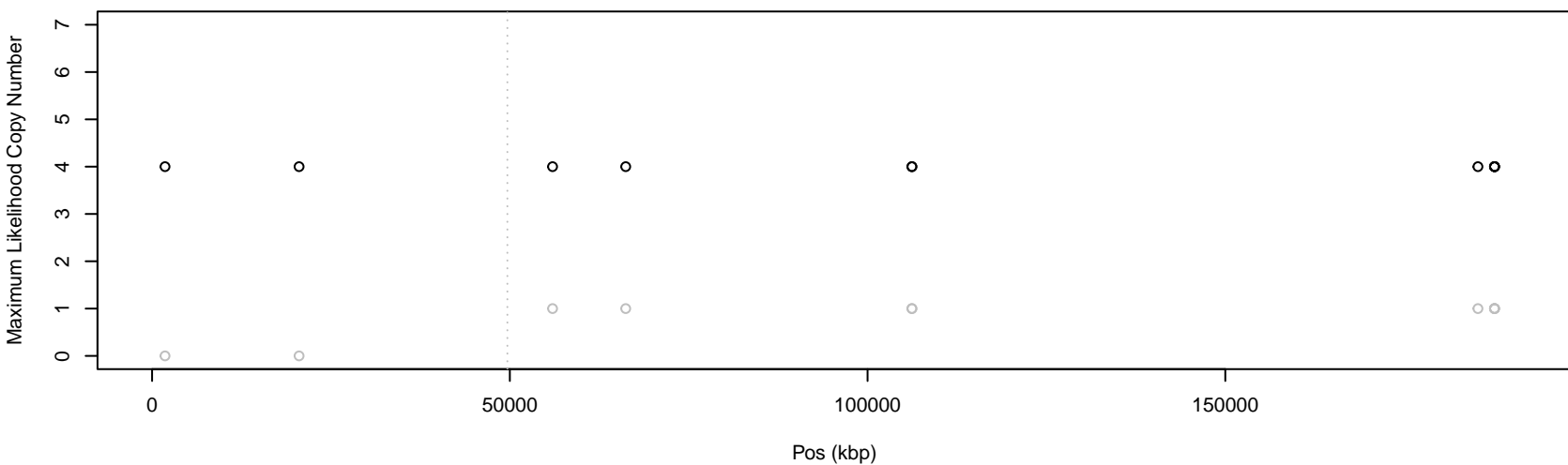
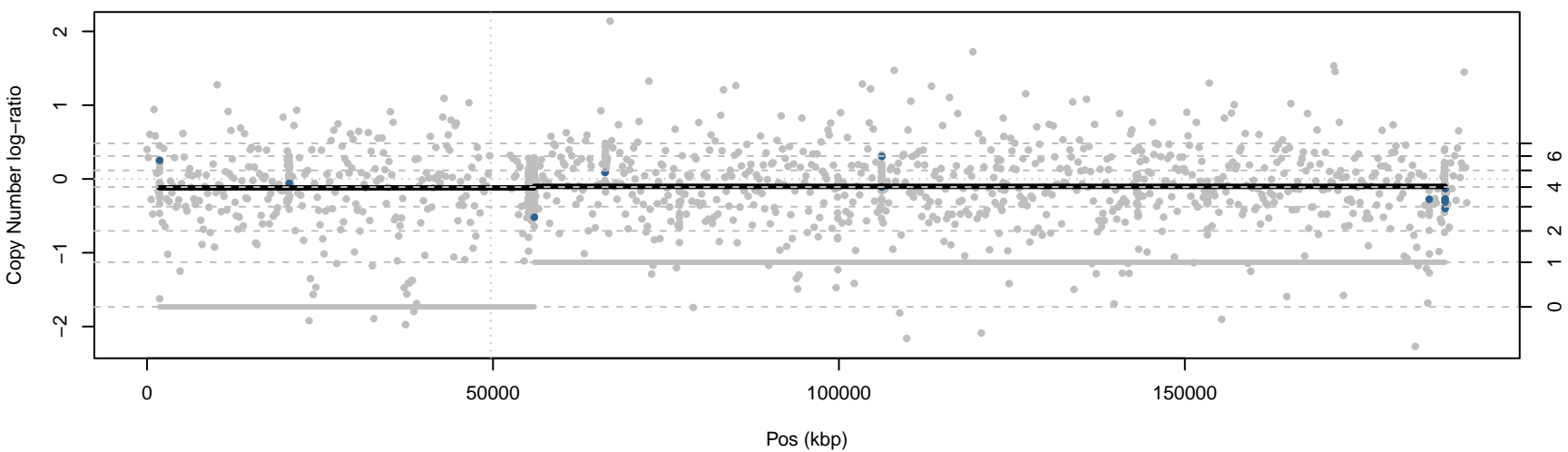
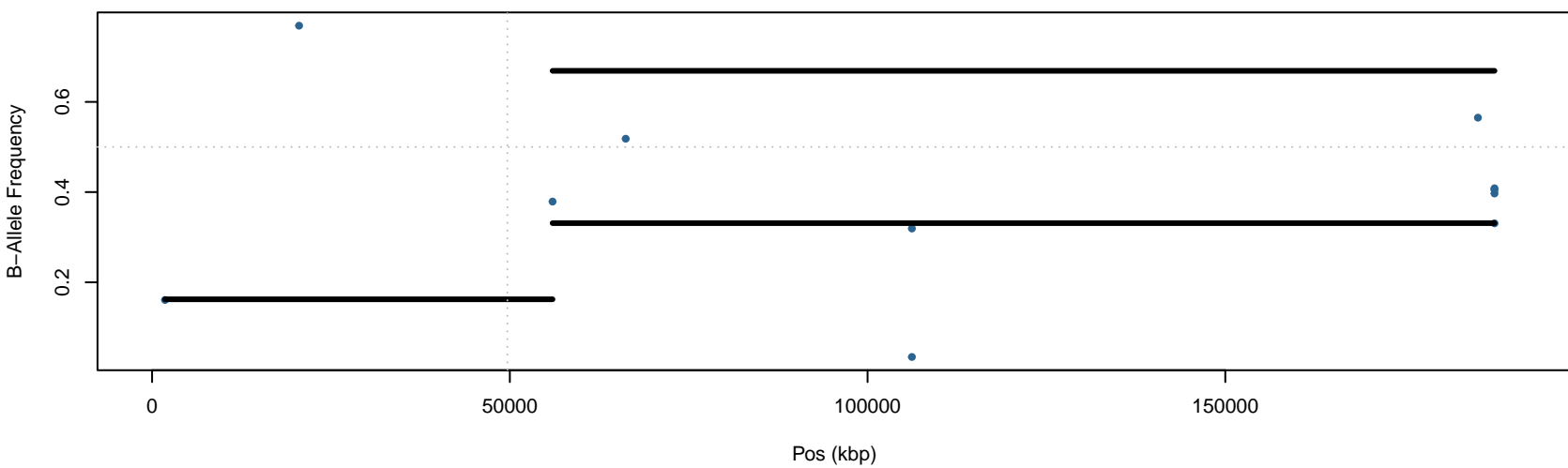
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 2



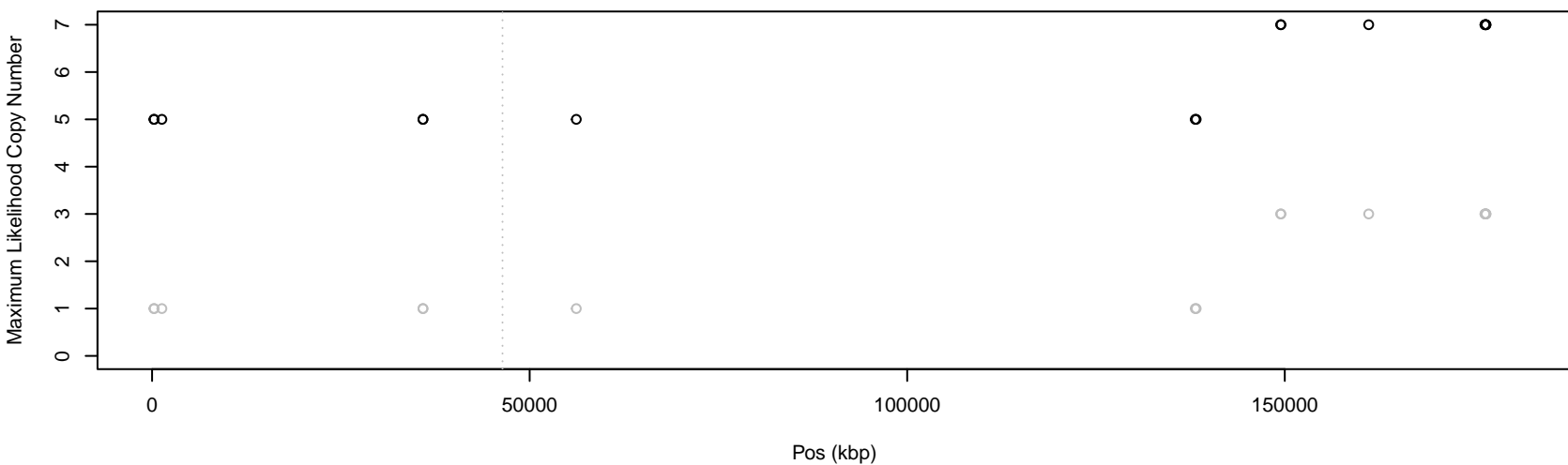
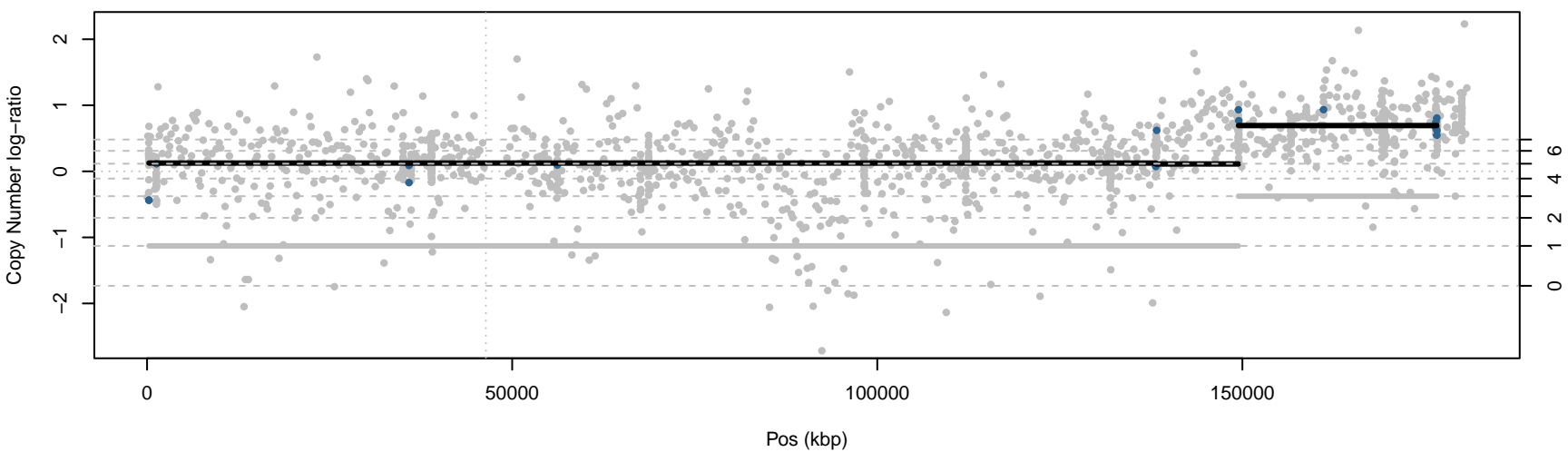
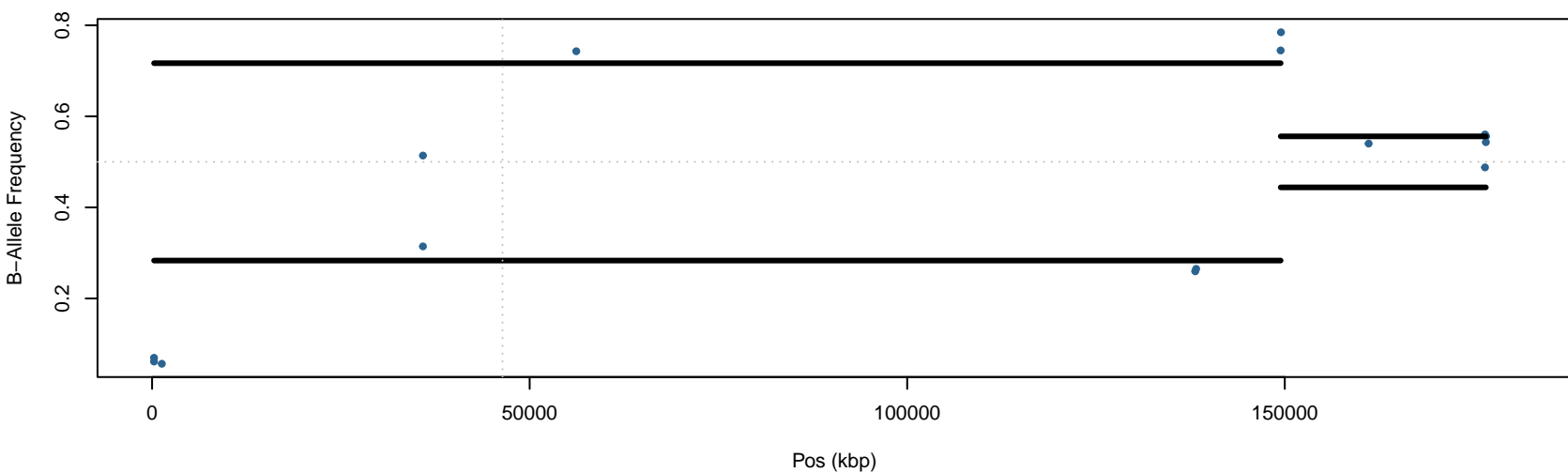
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 3



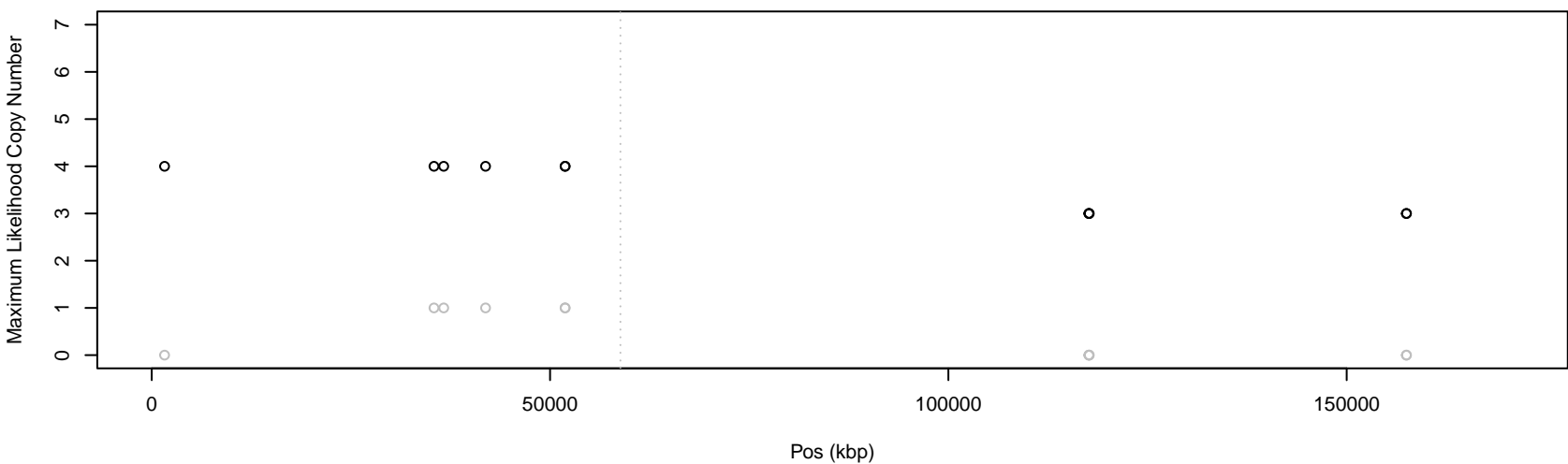
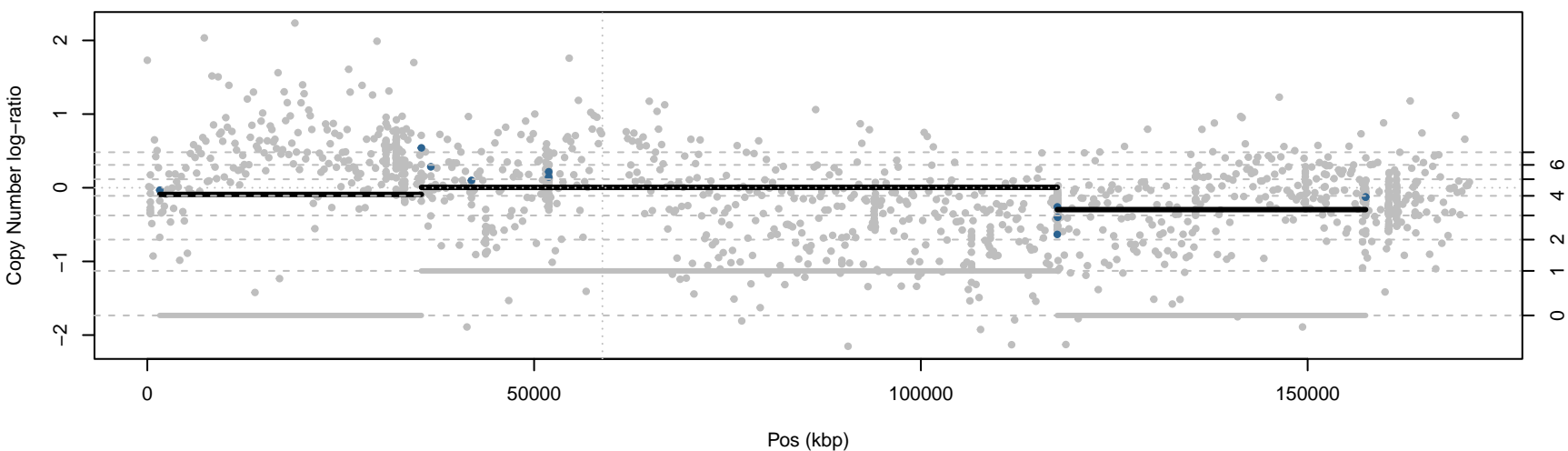
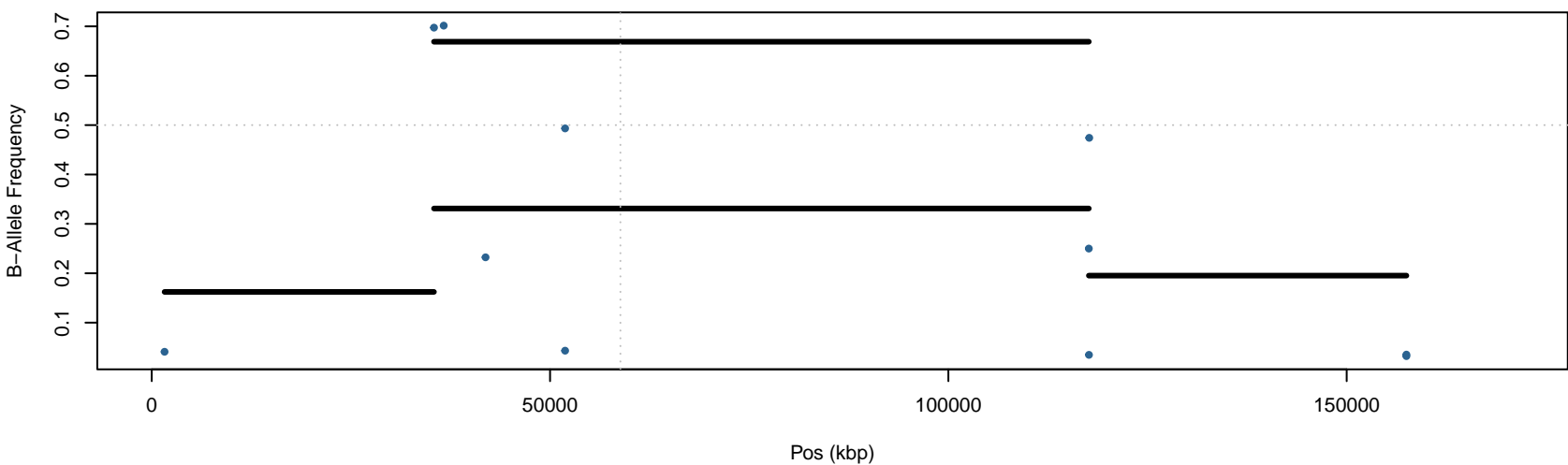
**Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 4**



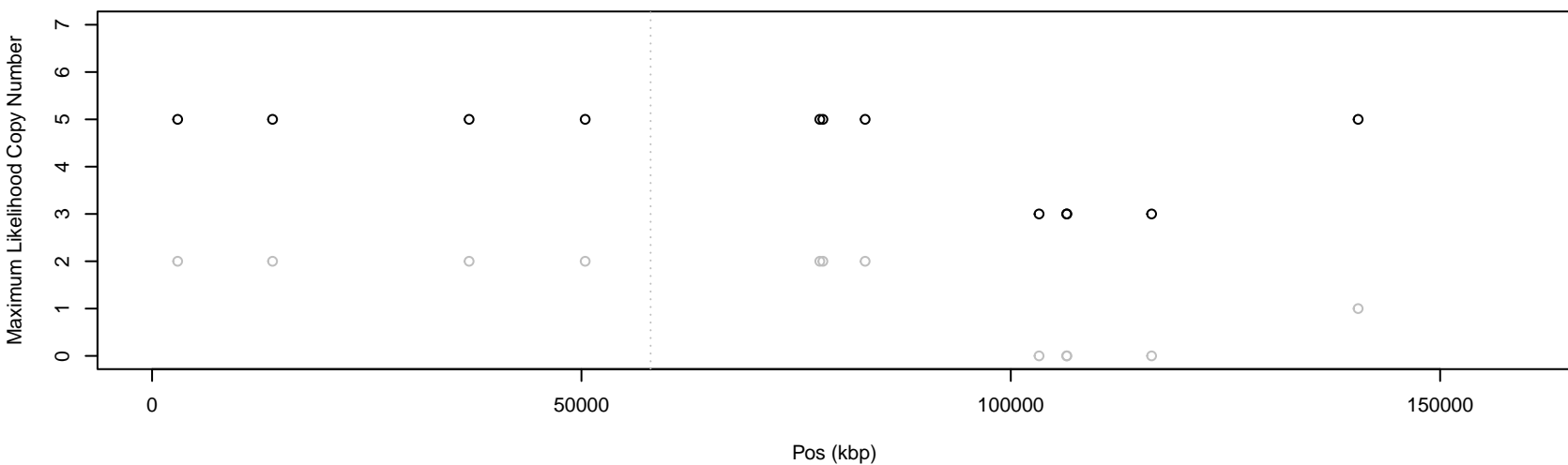
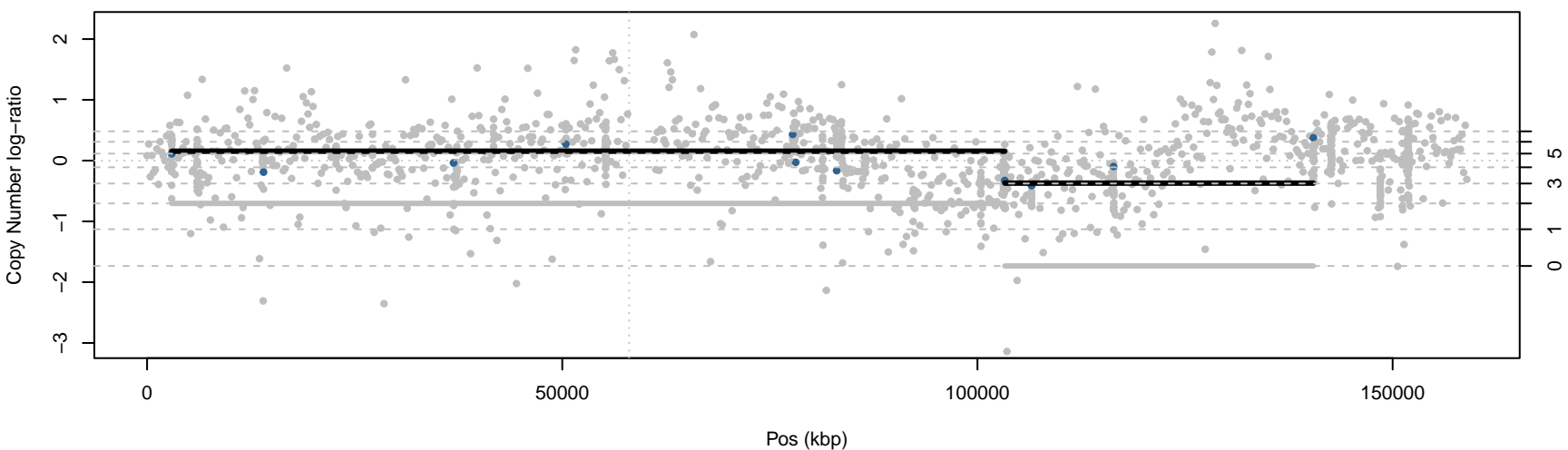
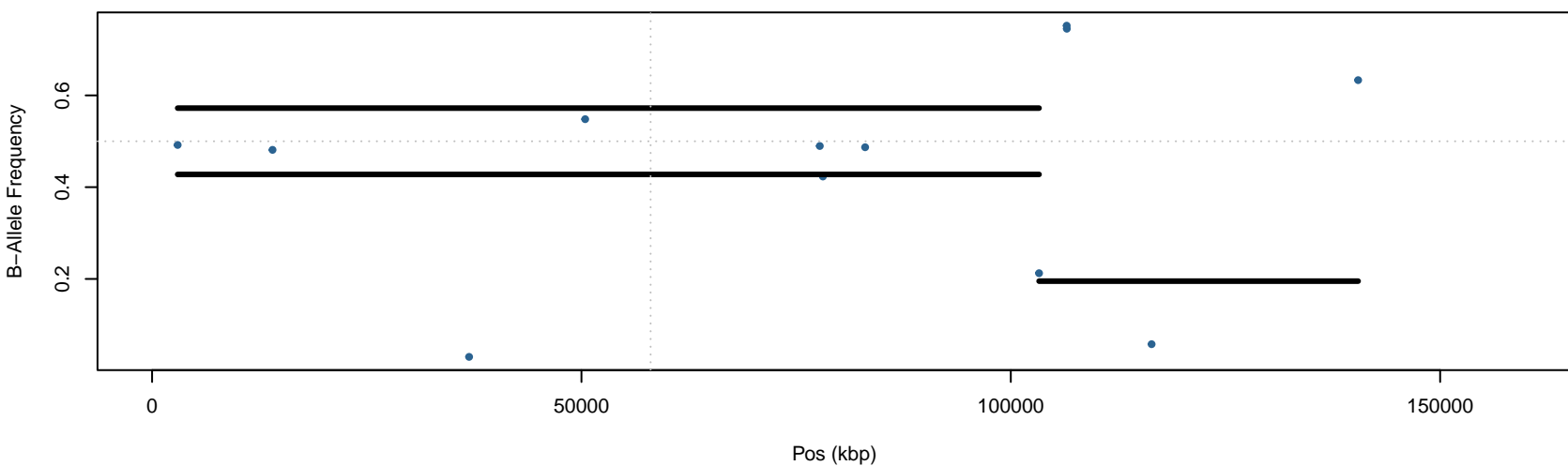
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 5



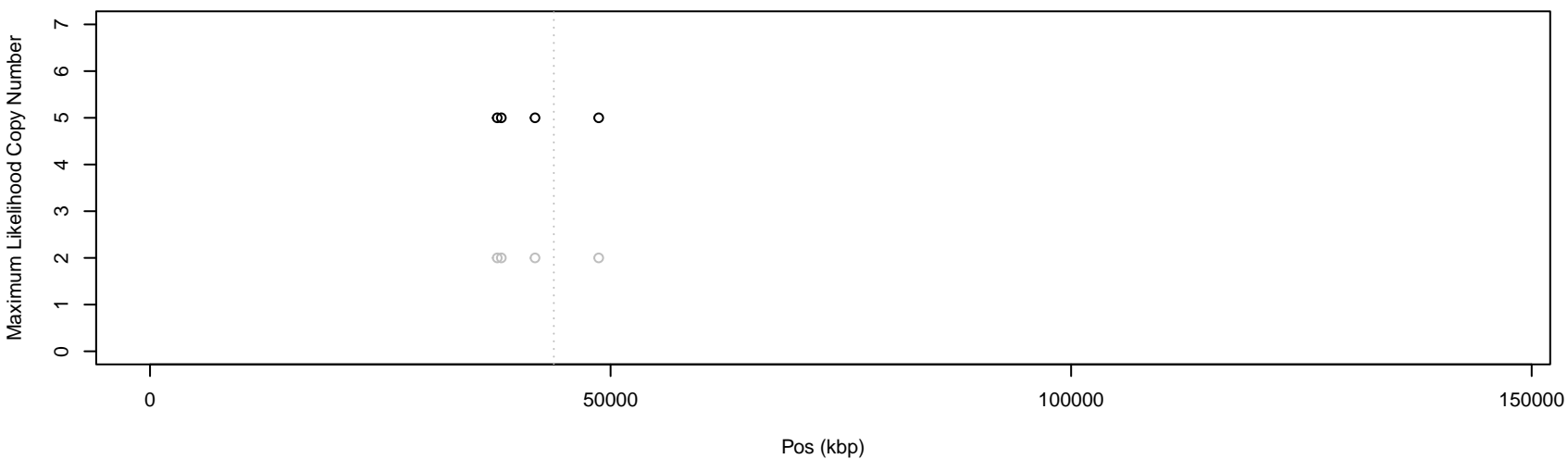
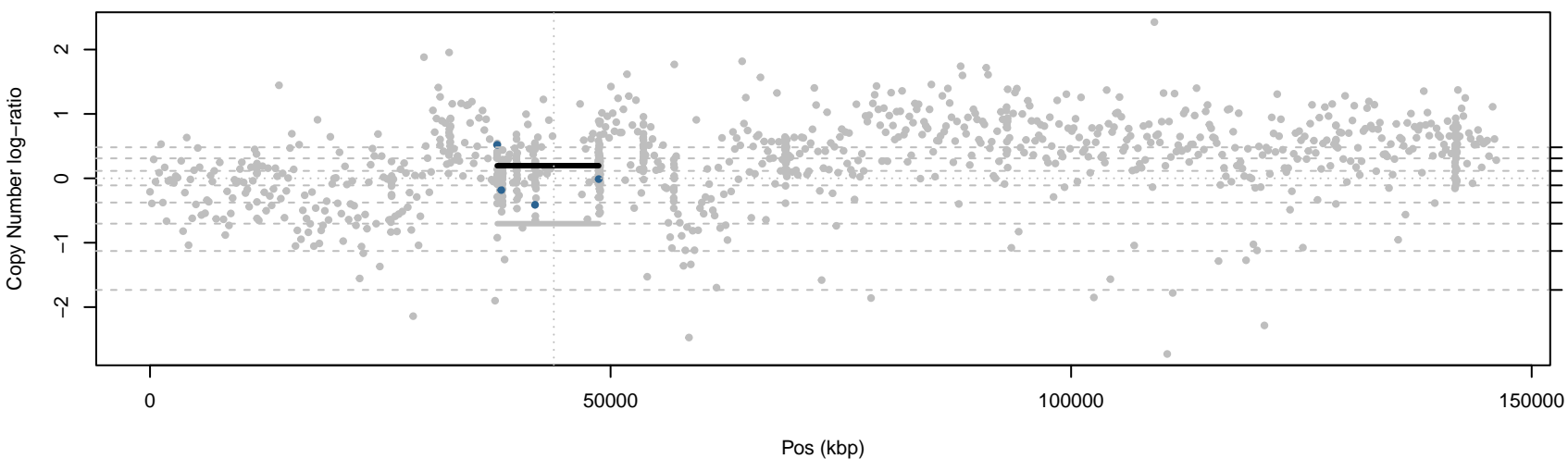
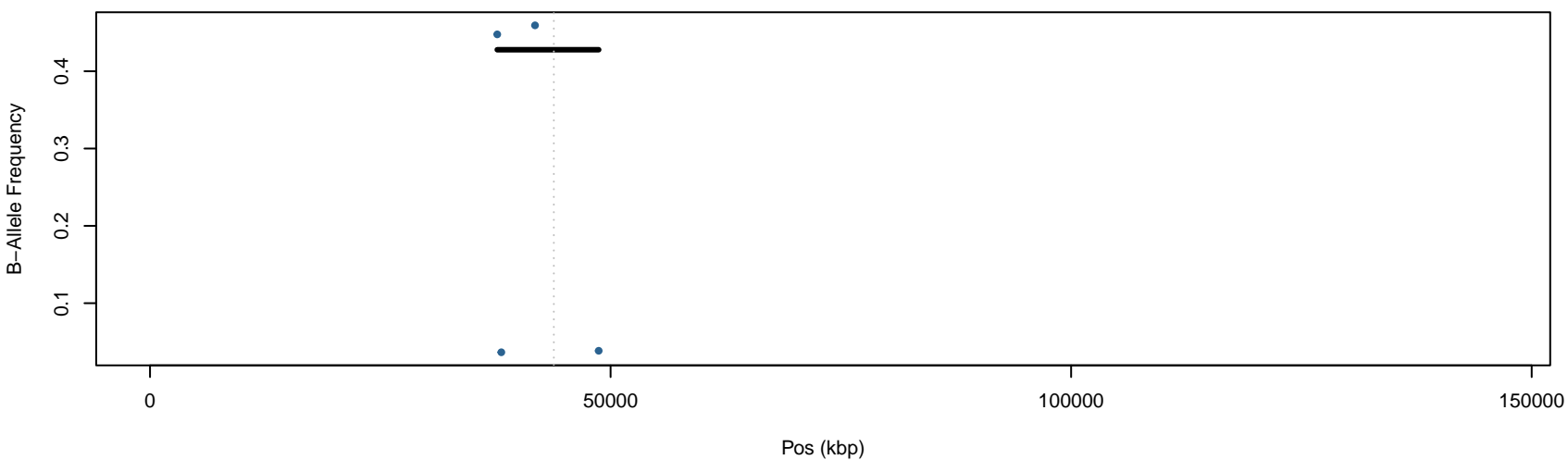
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 6



Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 7

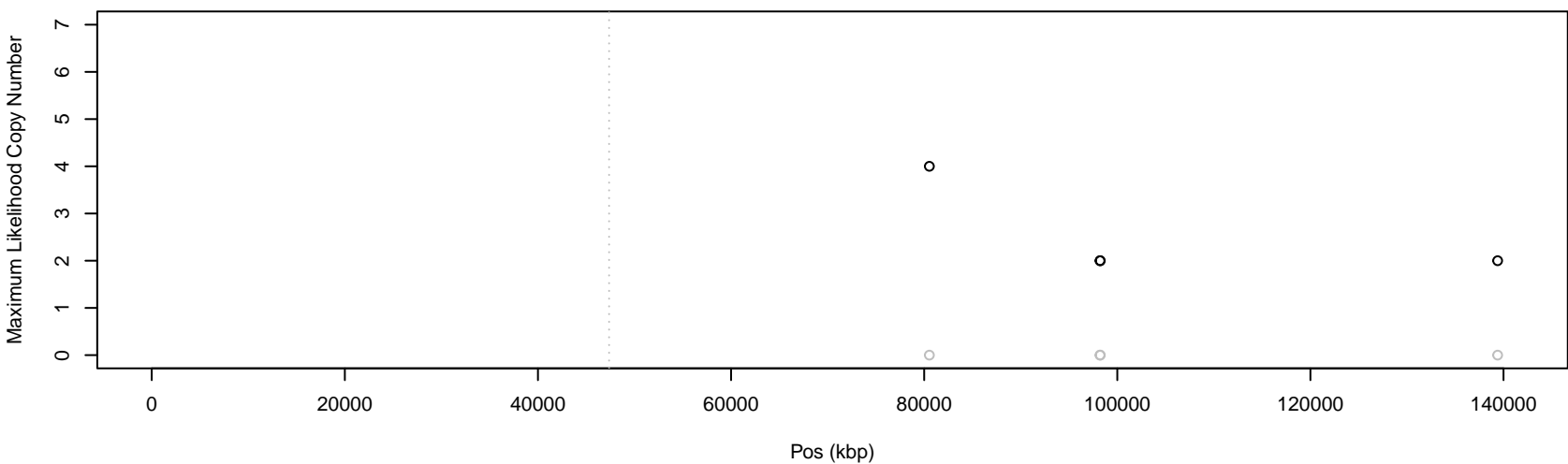
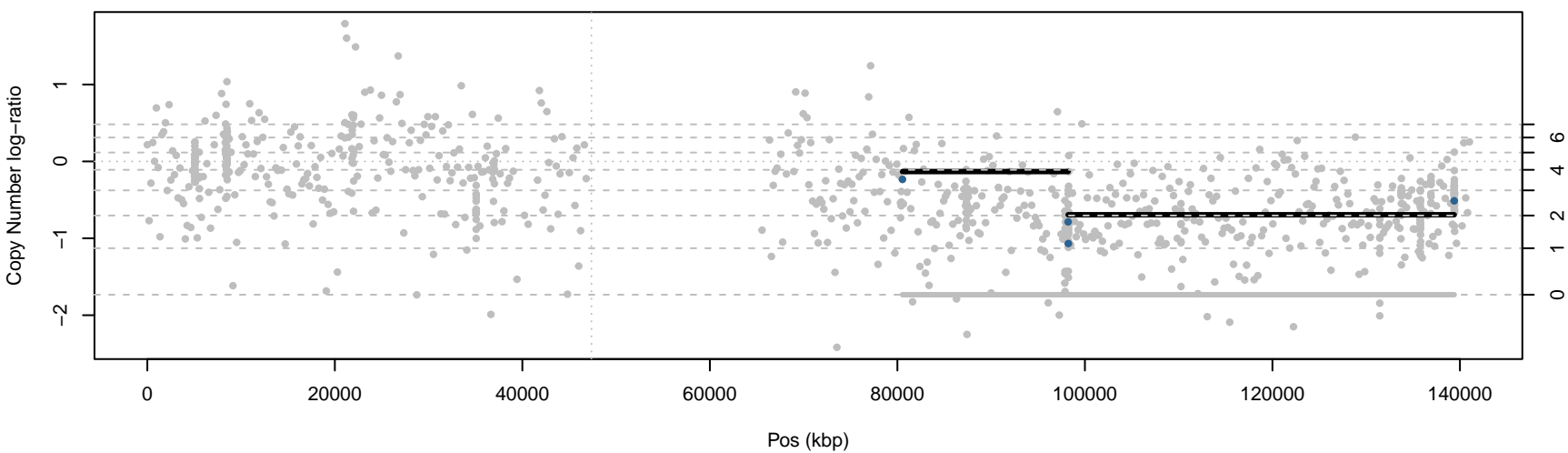
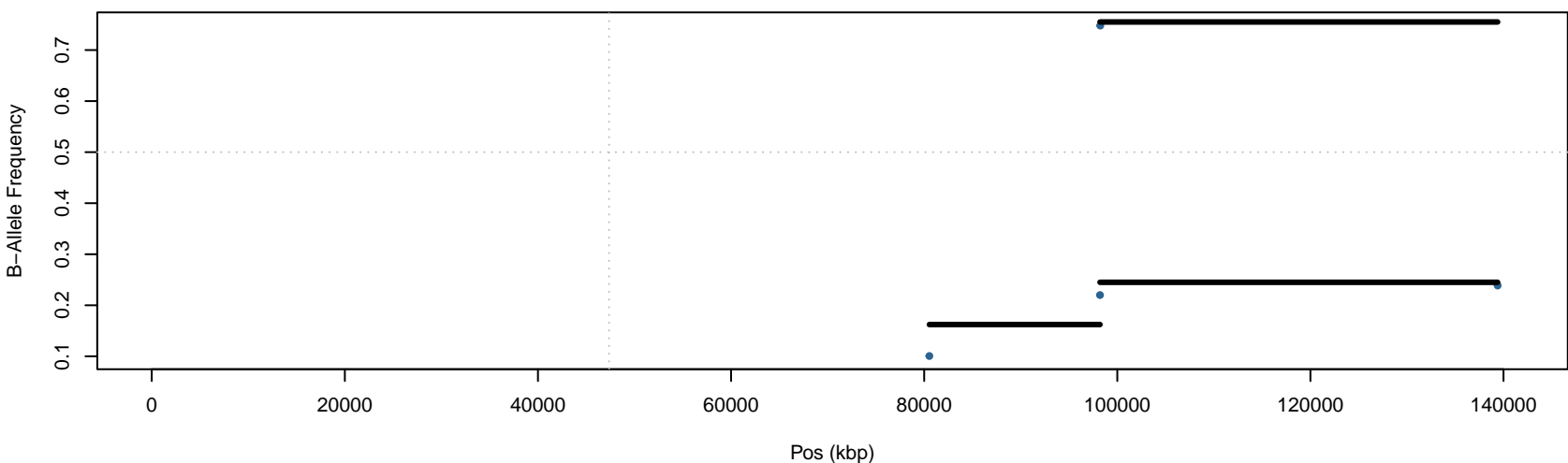


Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 8

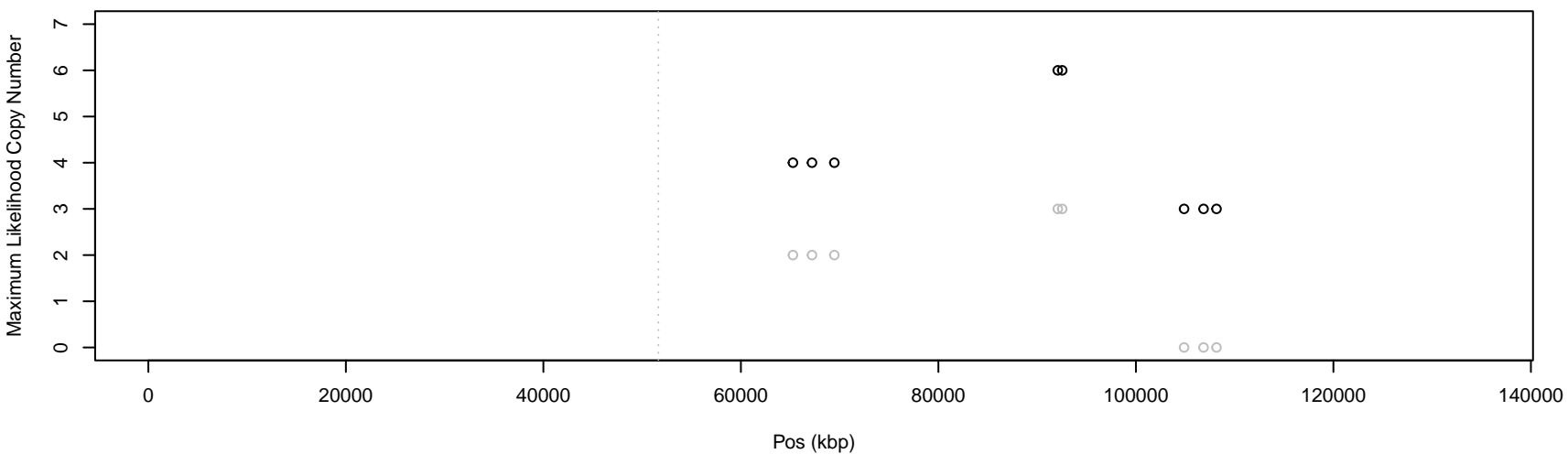
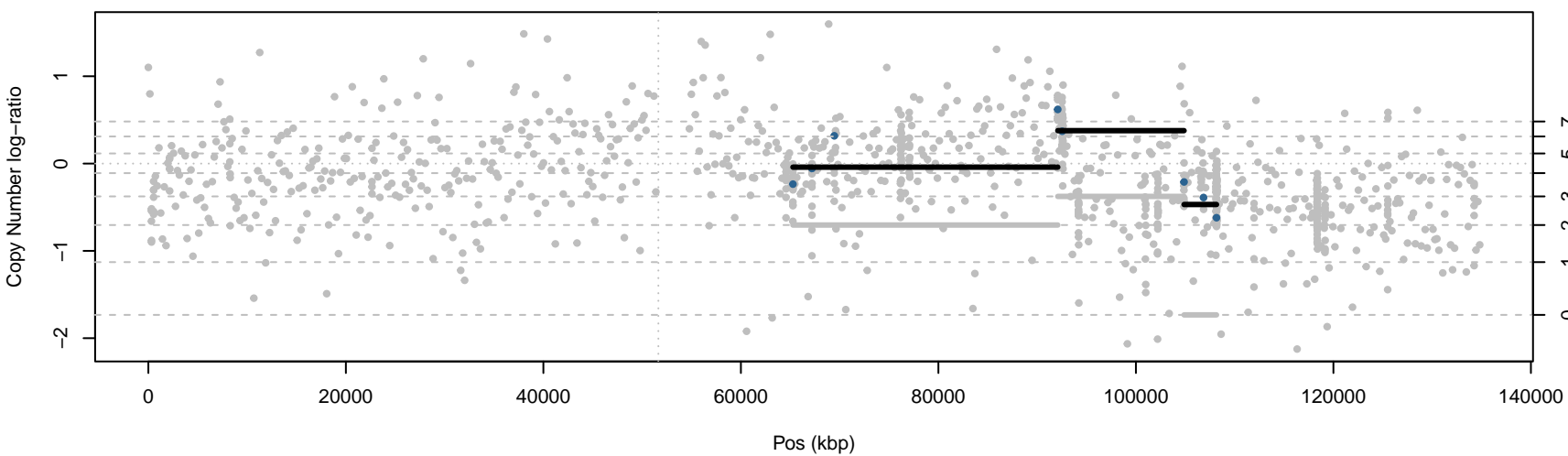
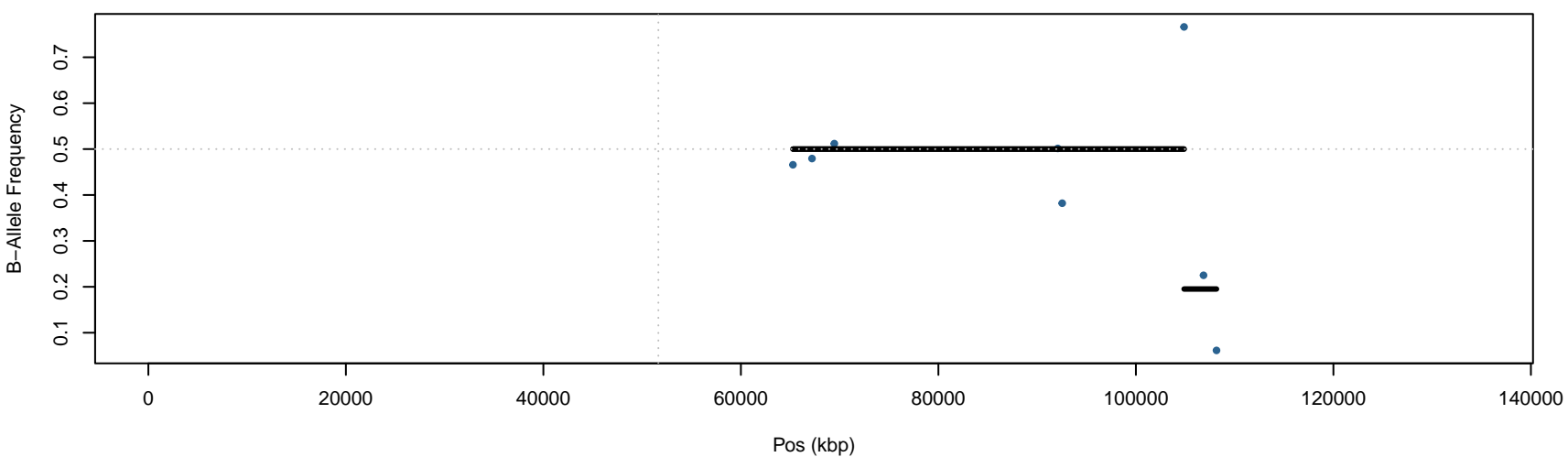




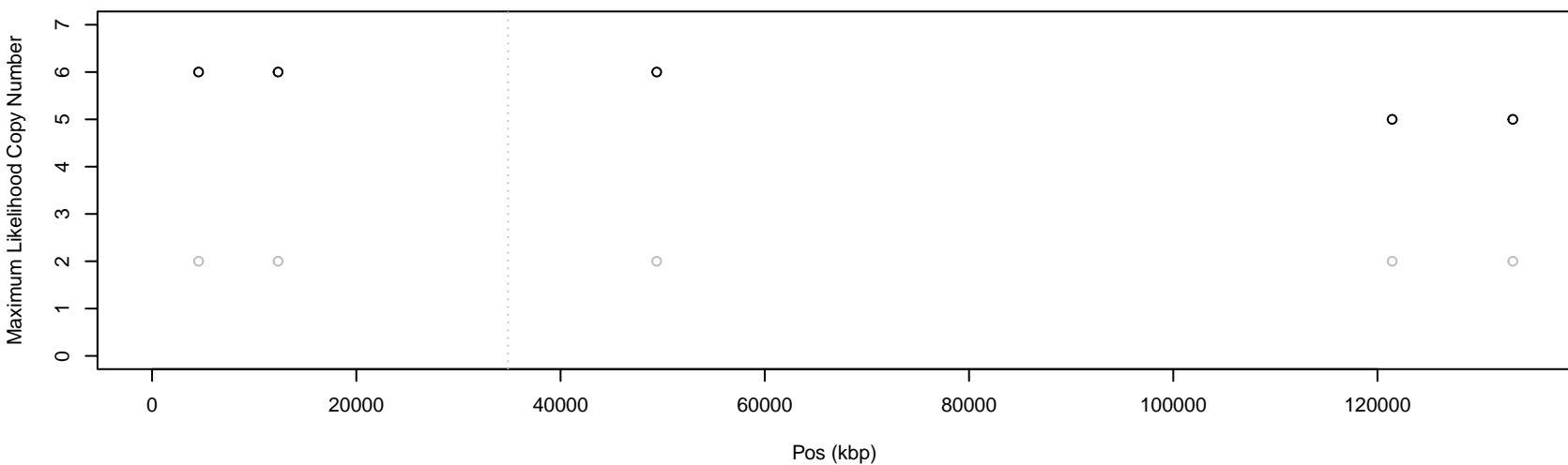
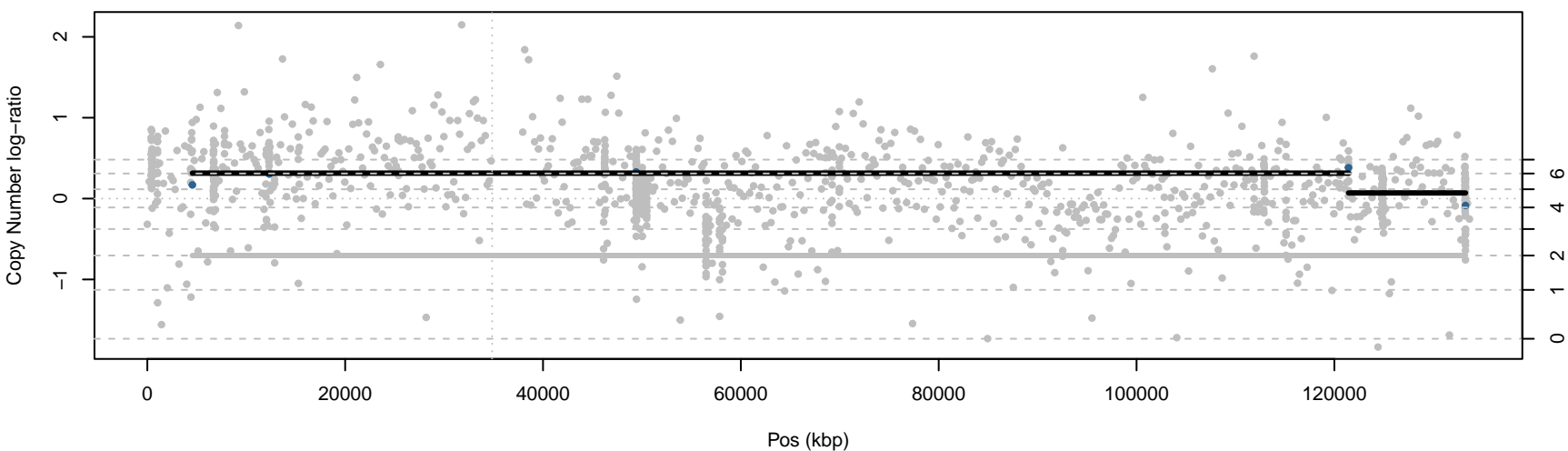
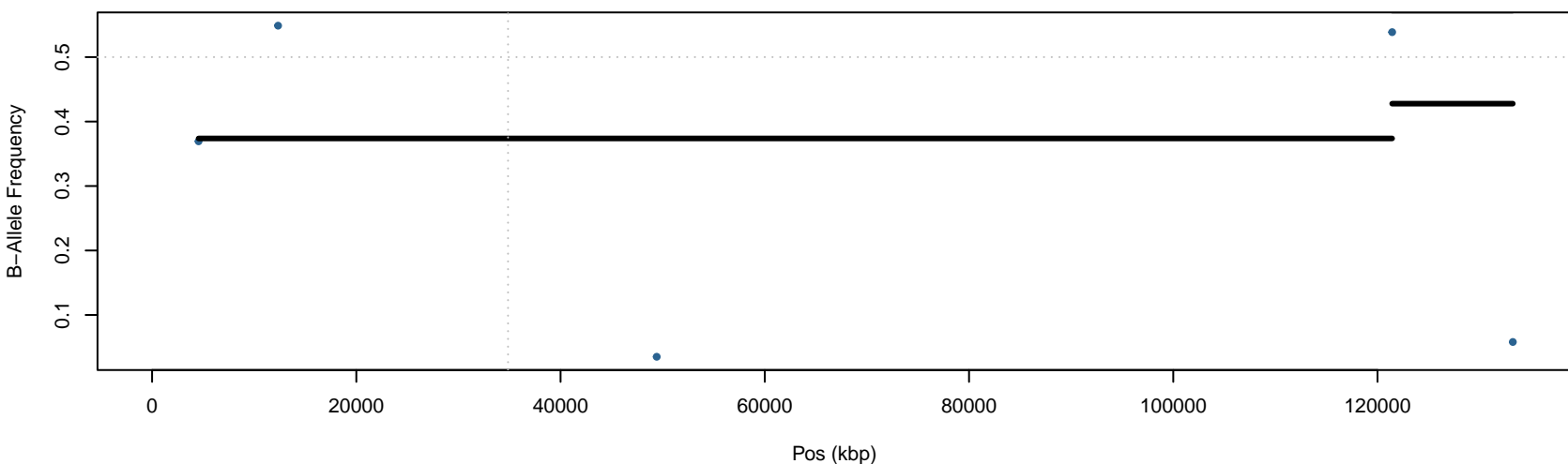
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 9



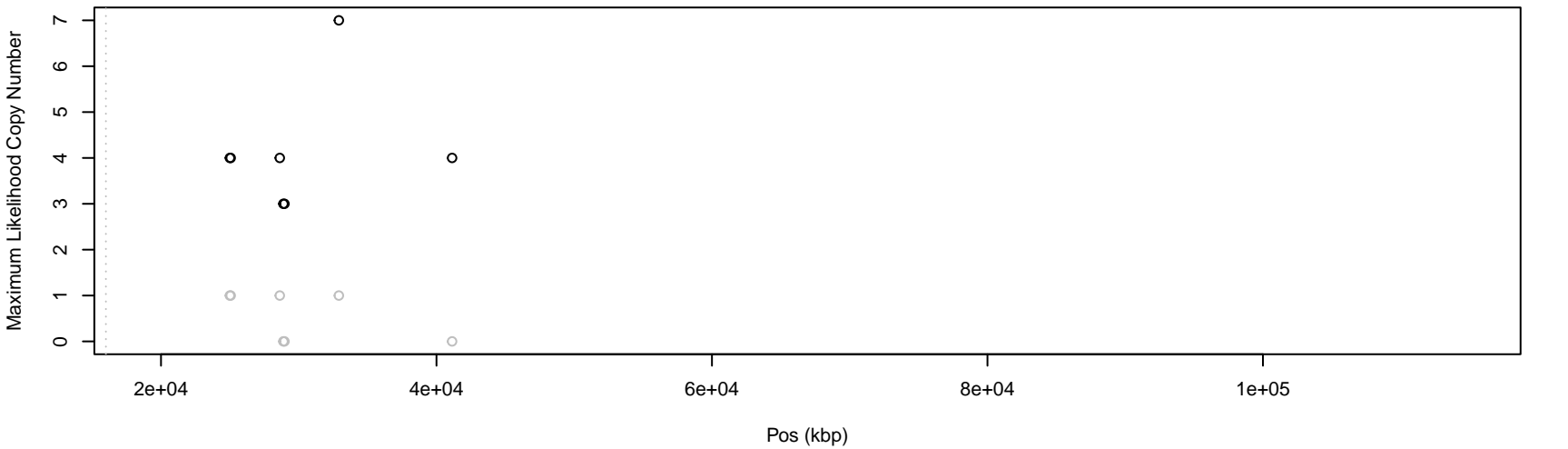
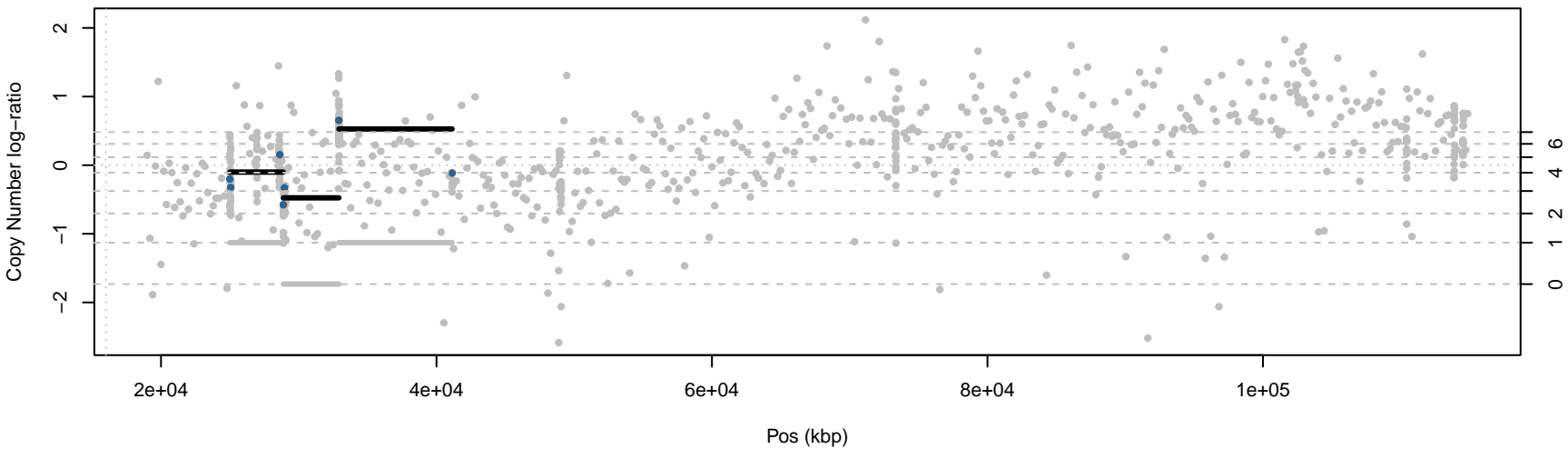
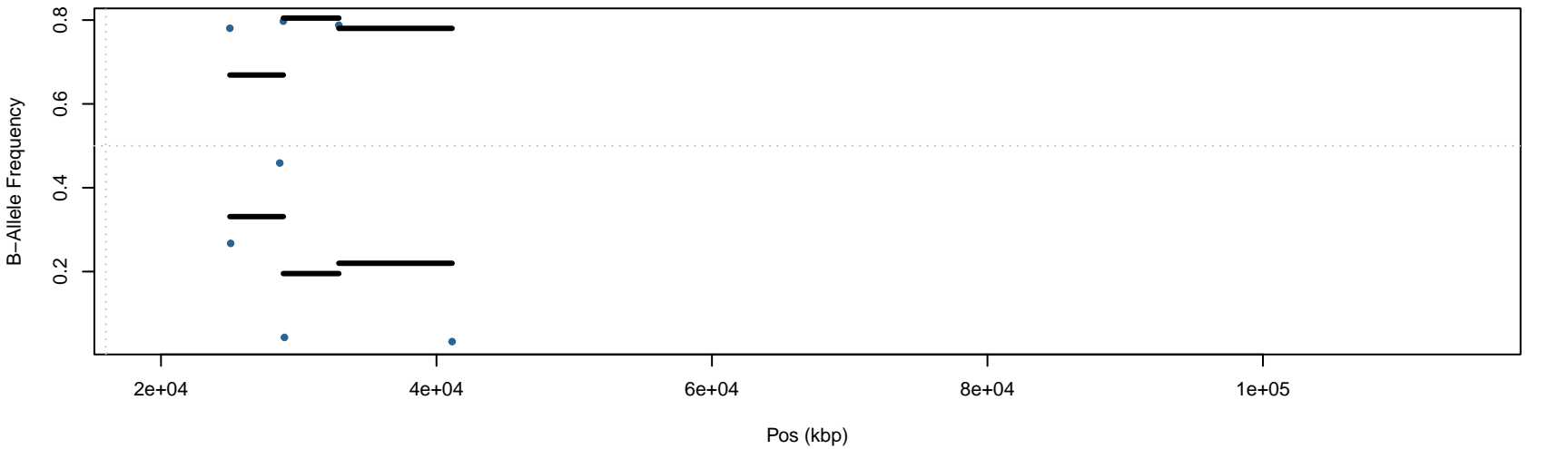
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 11



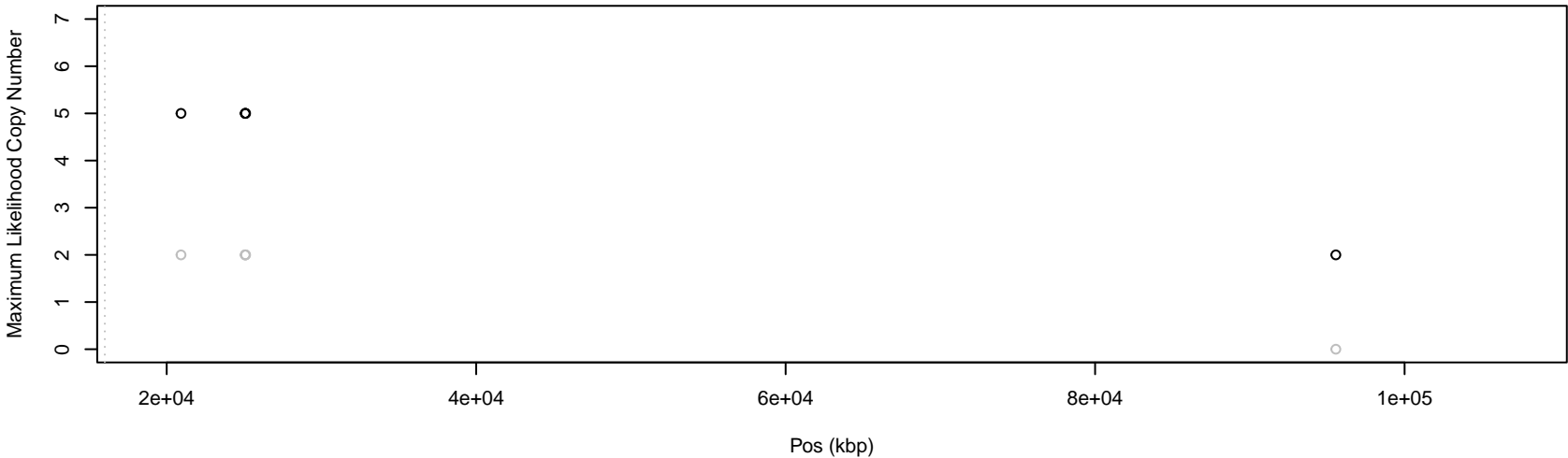
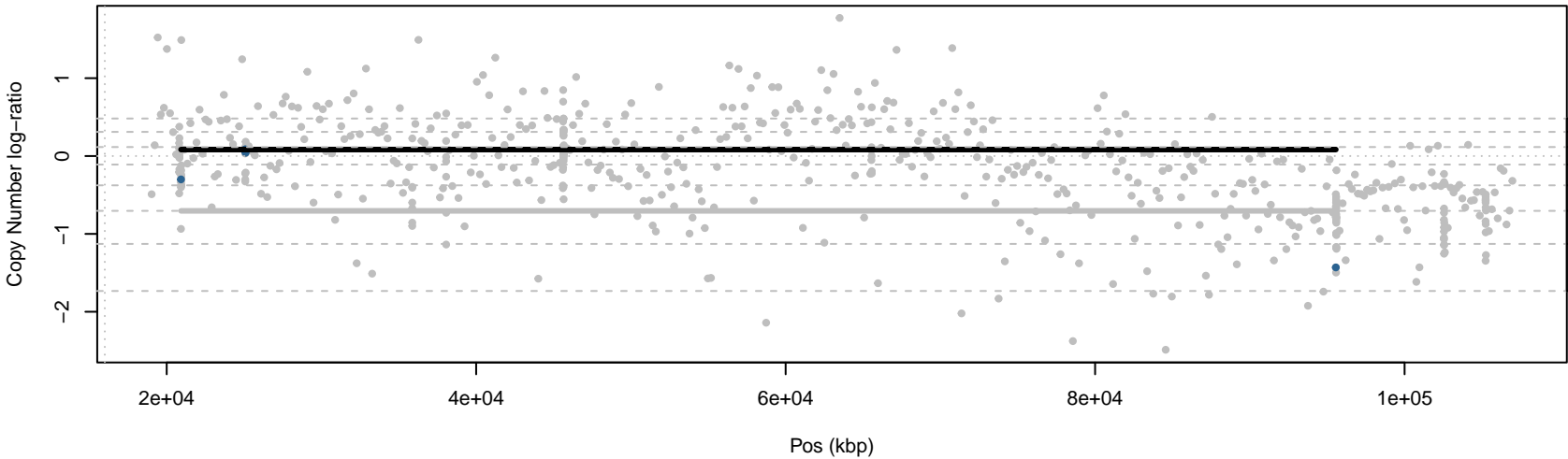
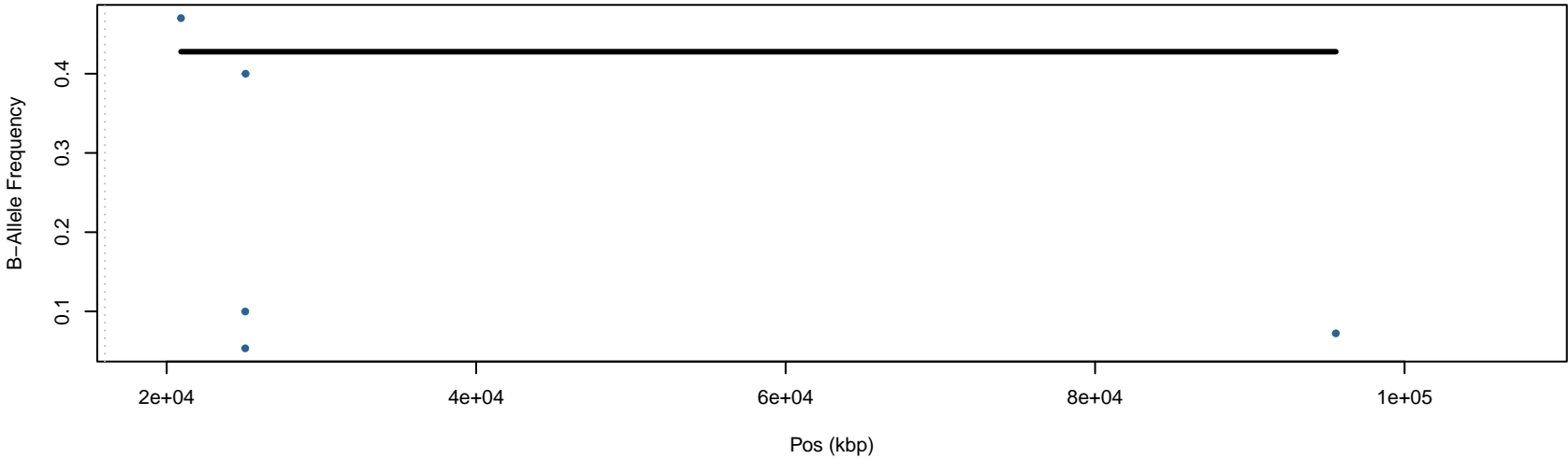
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 12



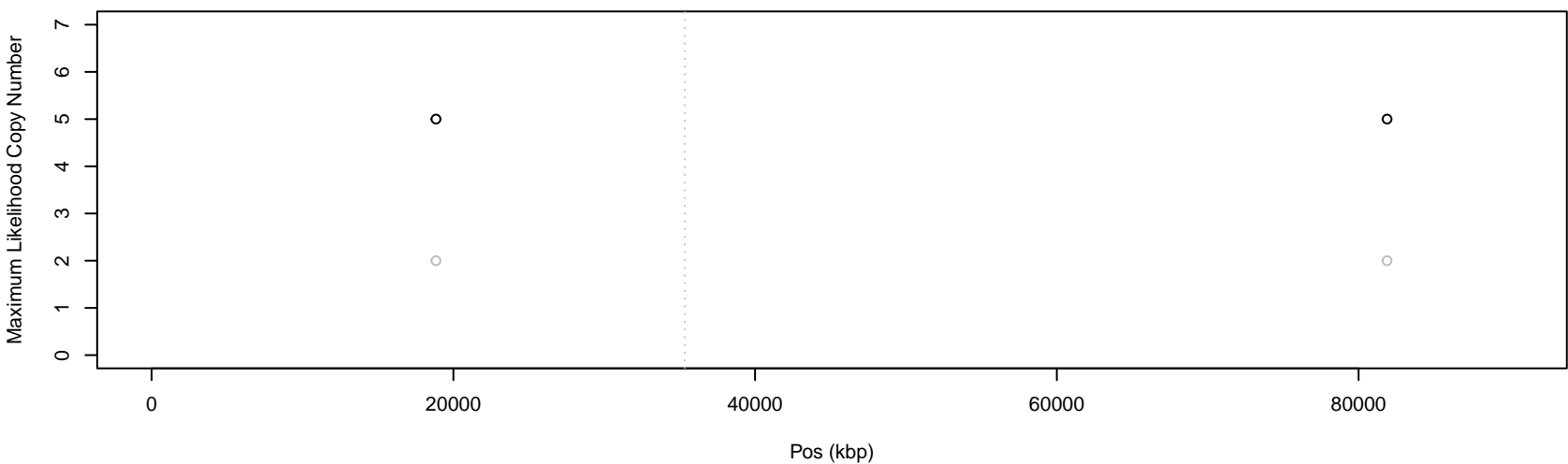
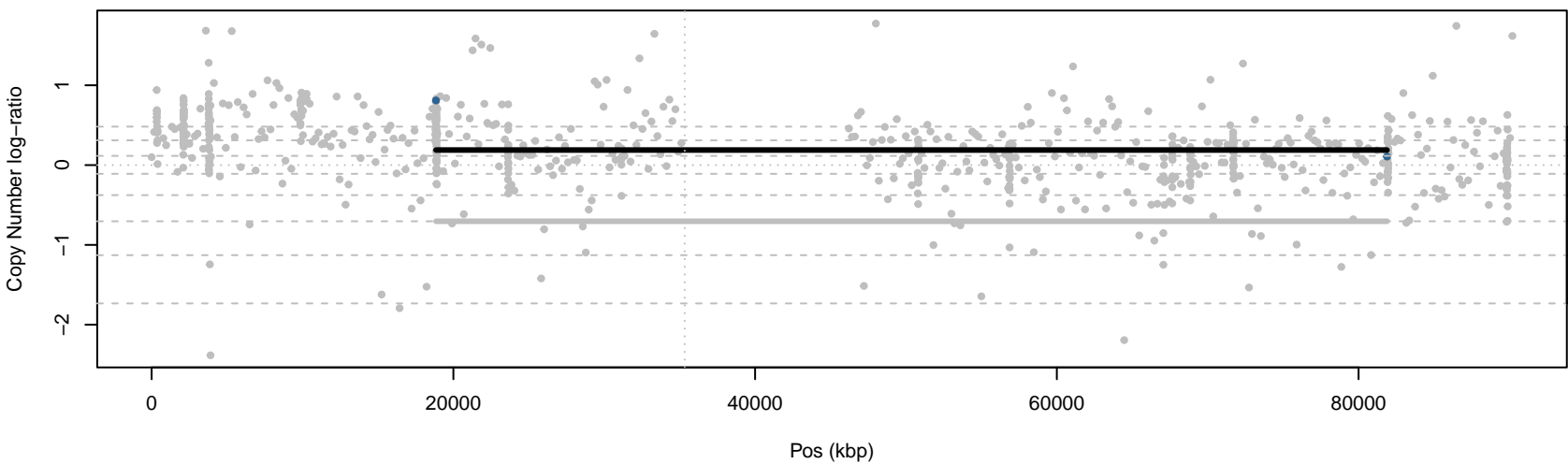
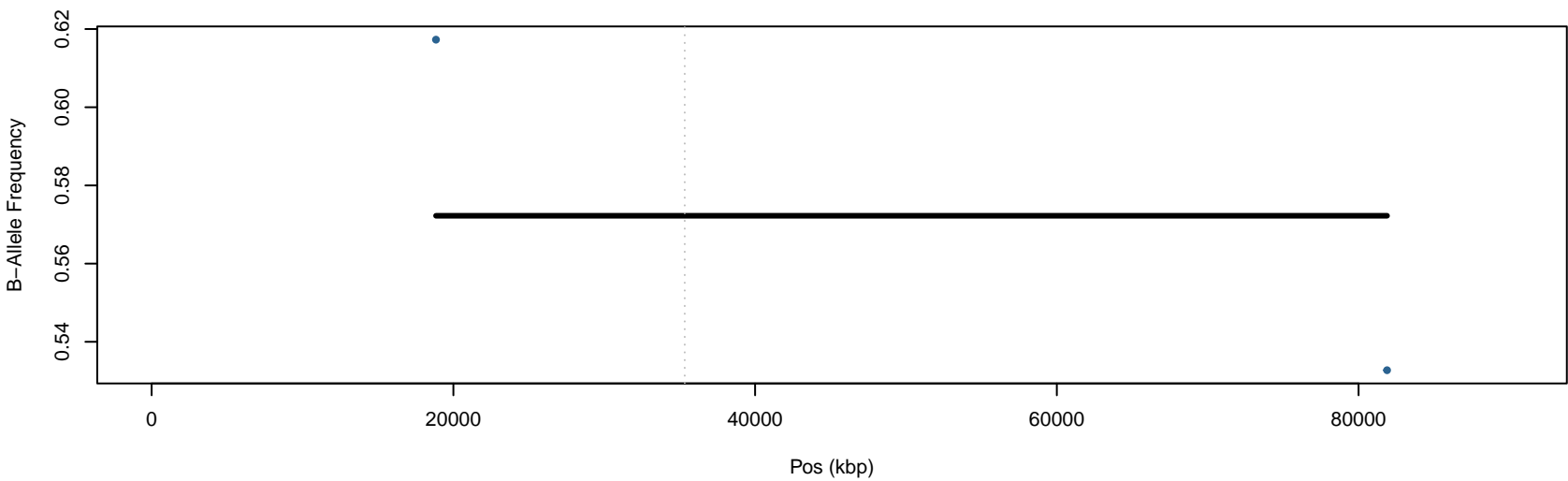
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 13



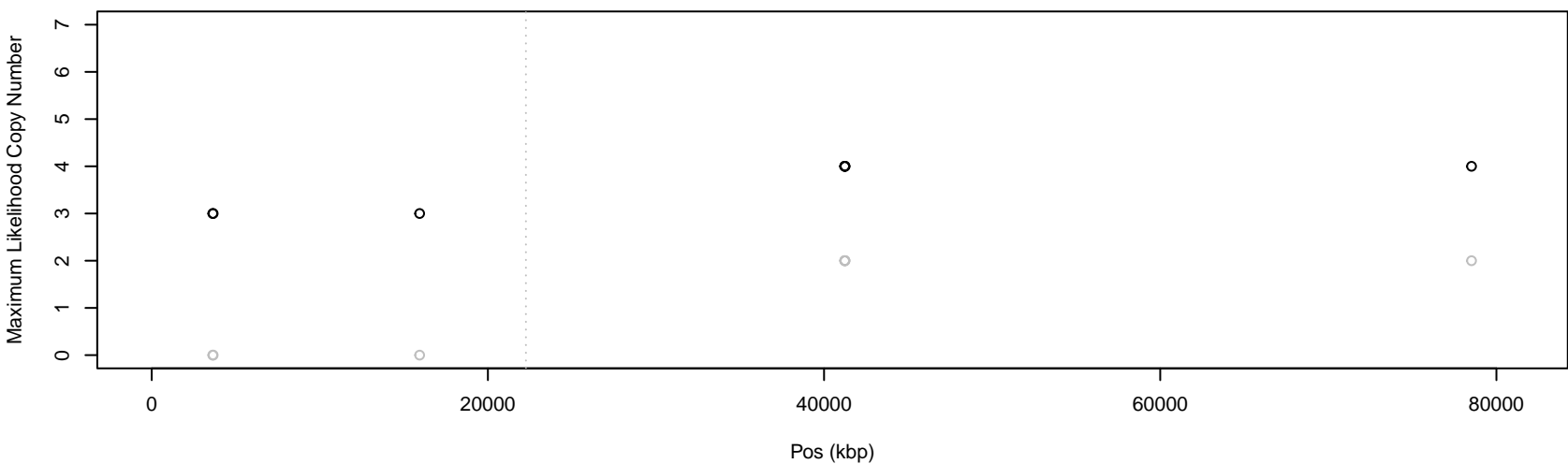
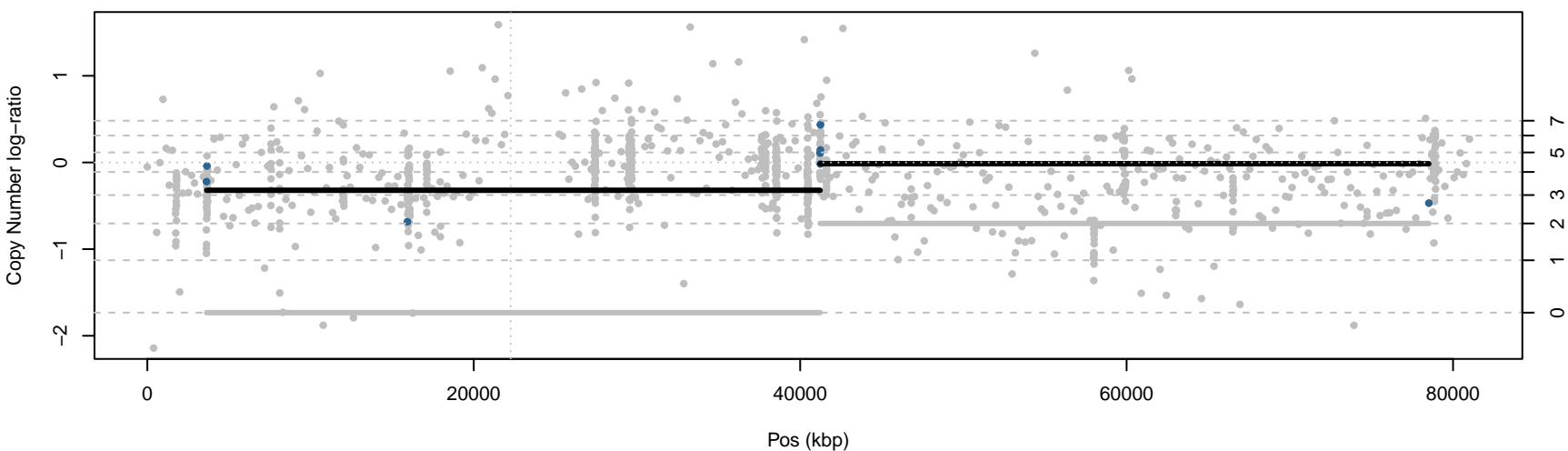
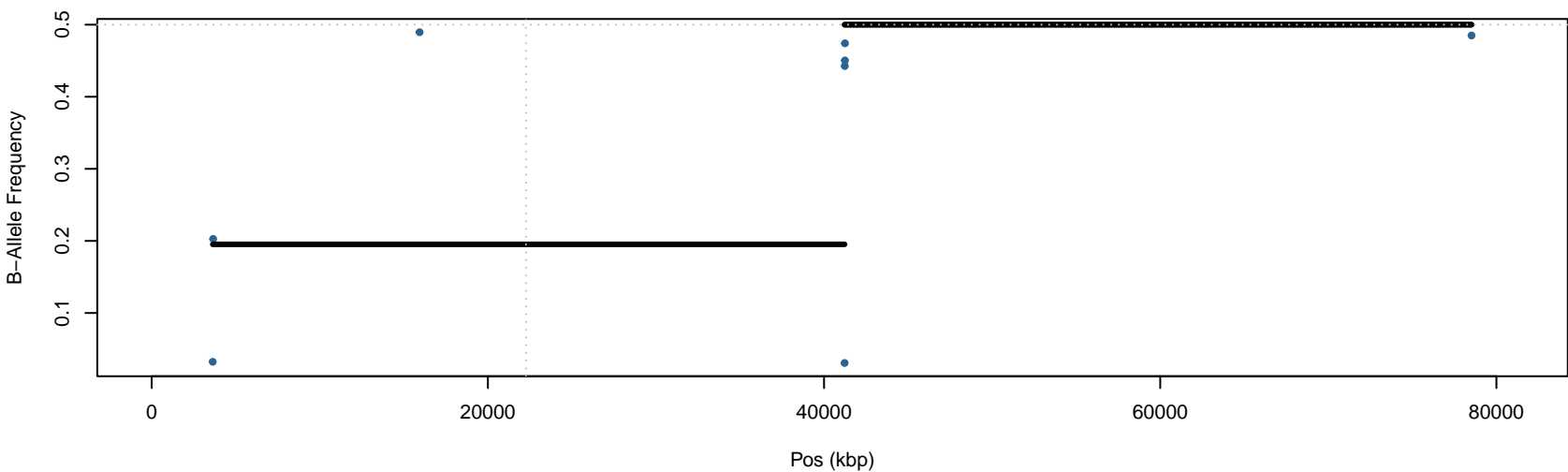
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 14



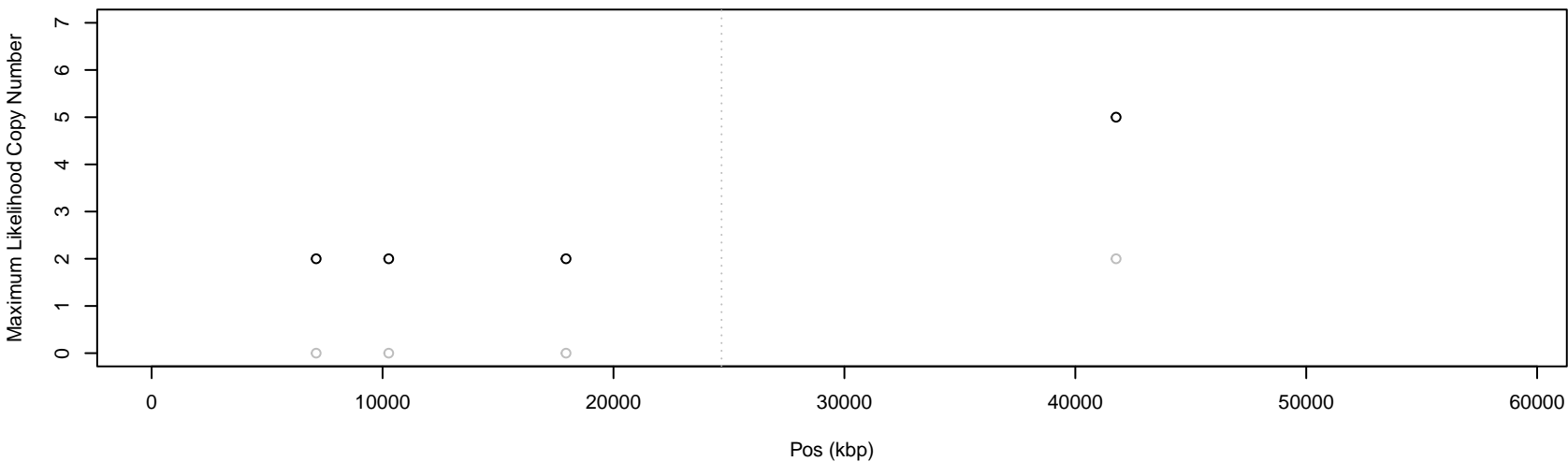
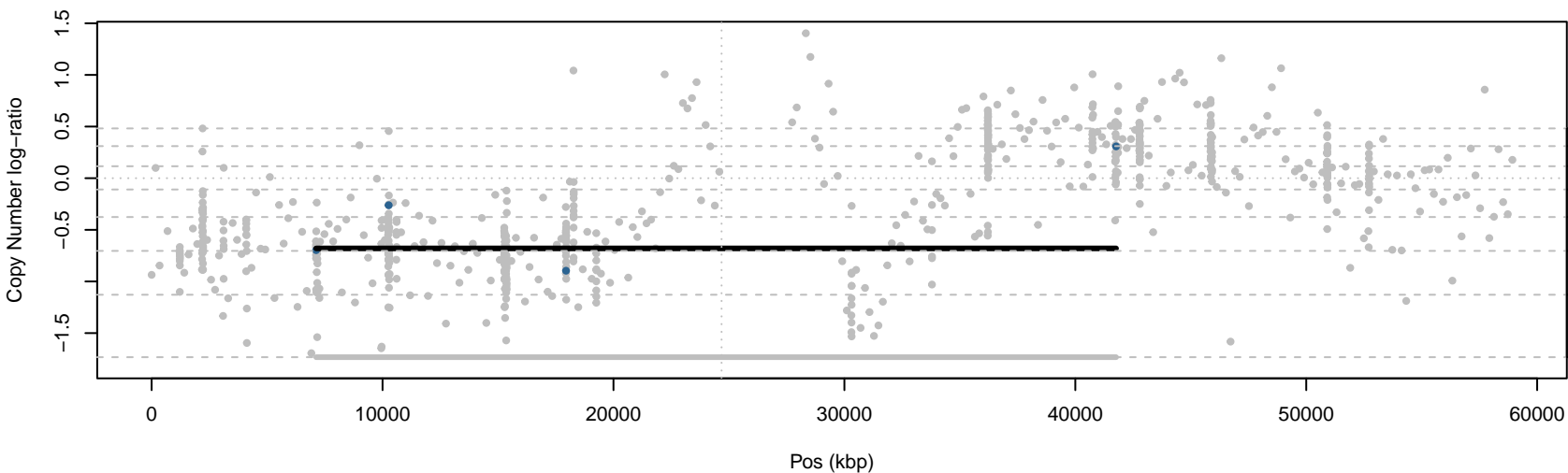
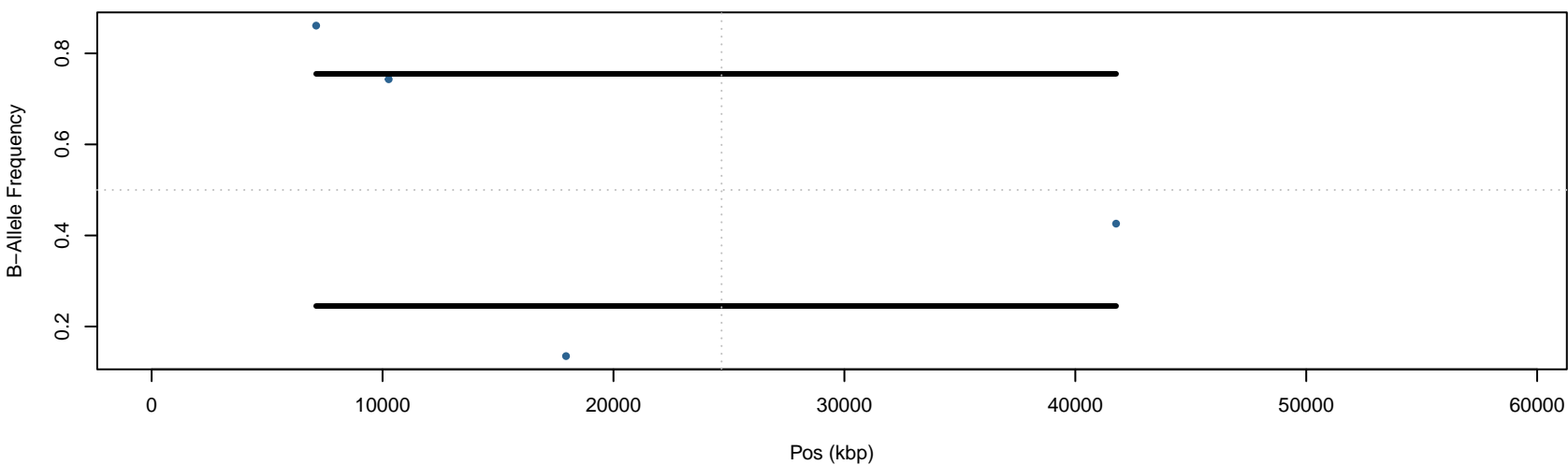
Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 16



Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 17

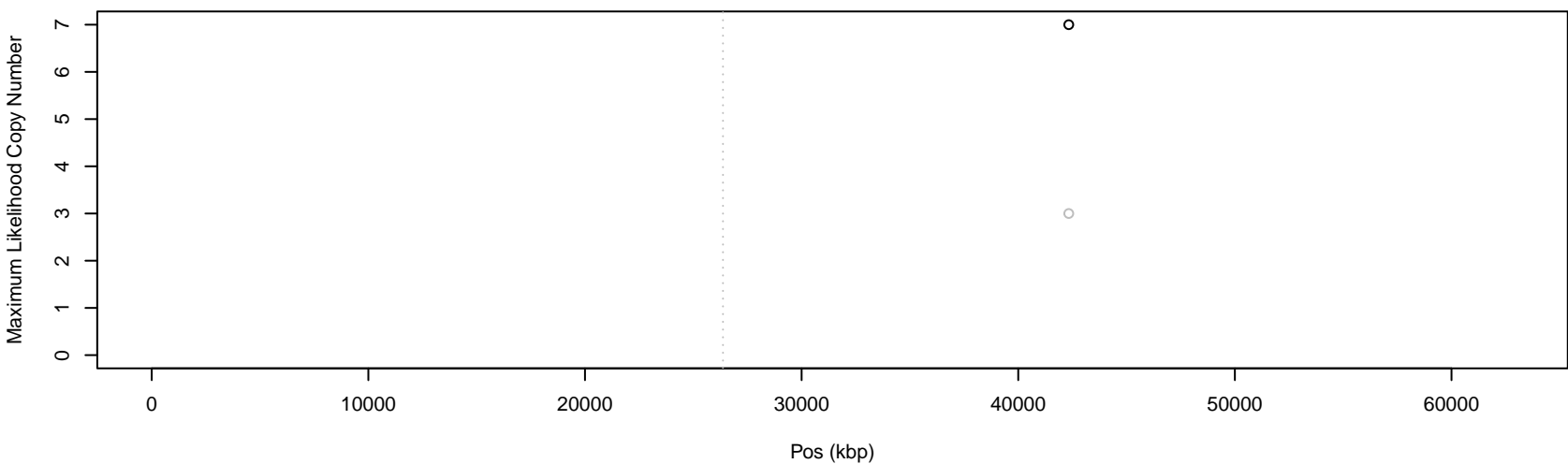
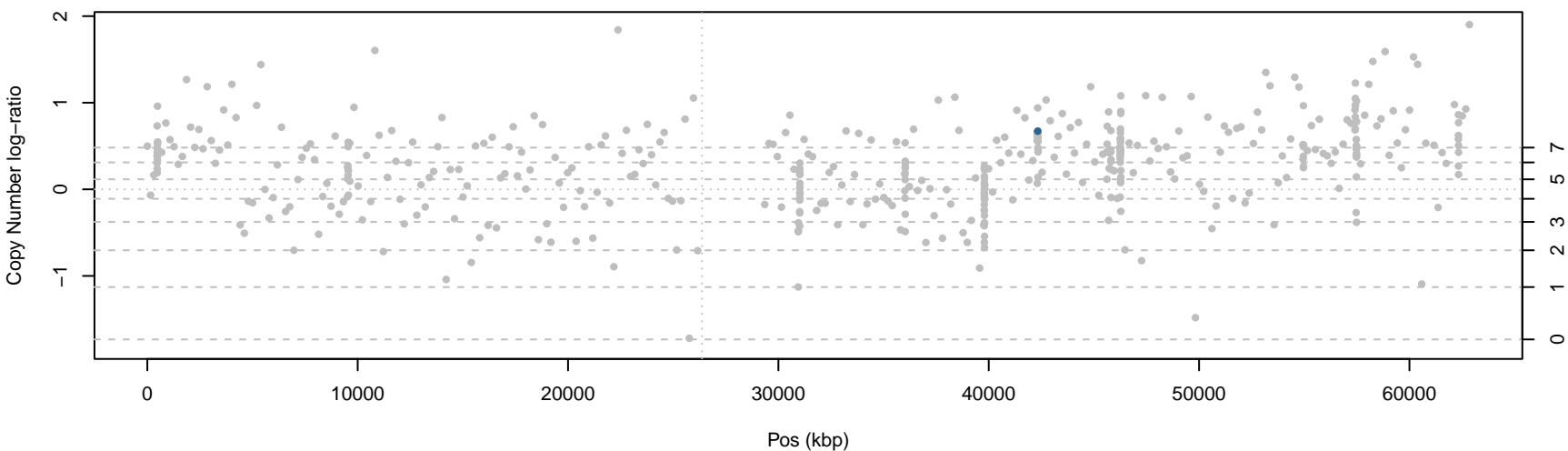
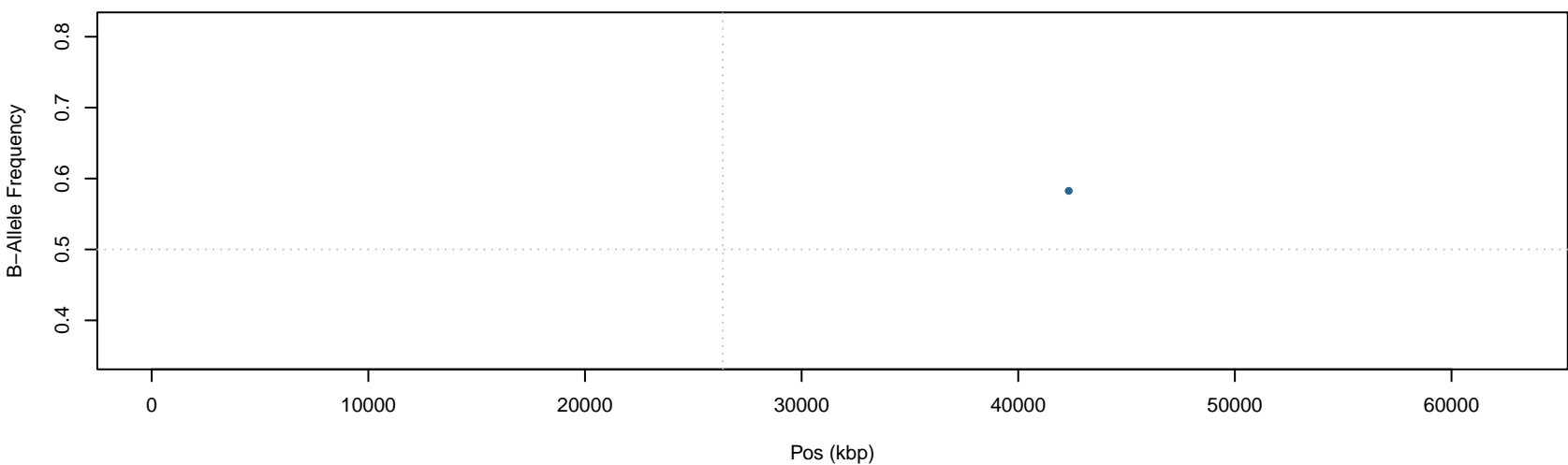


Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 19

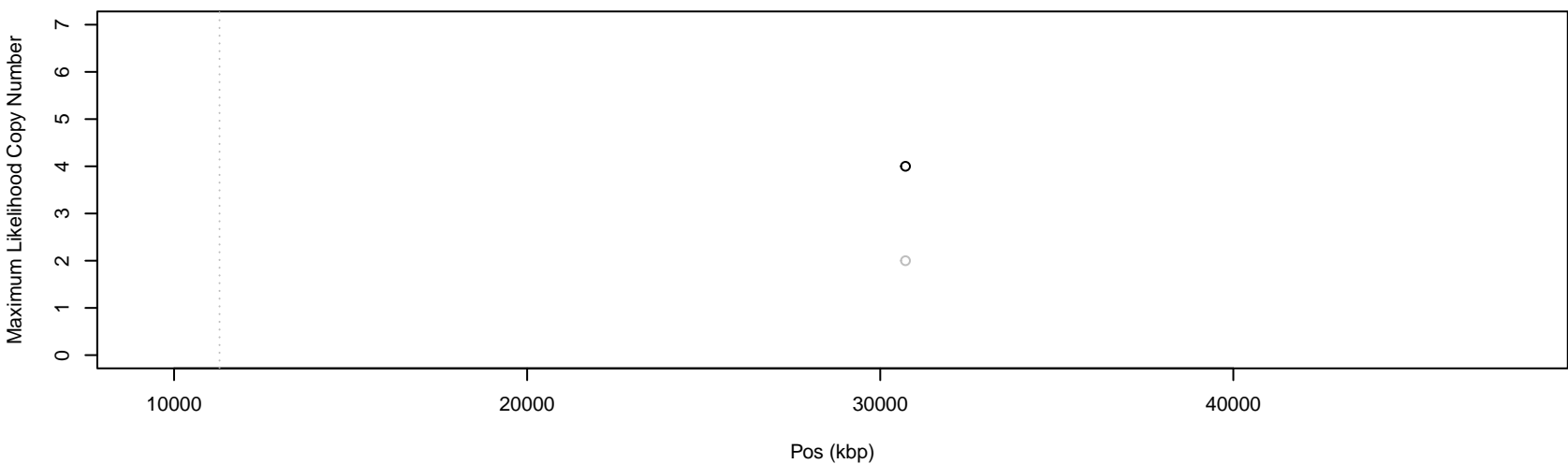
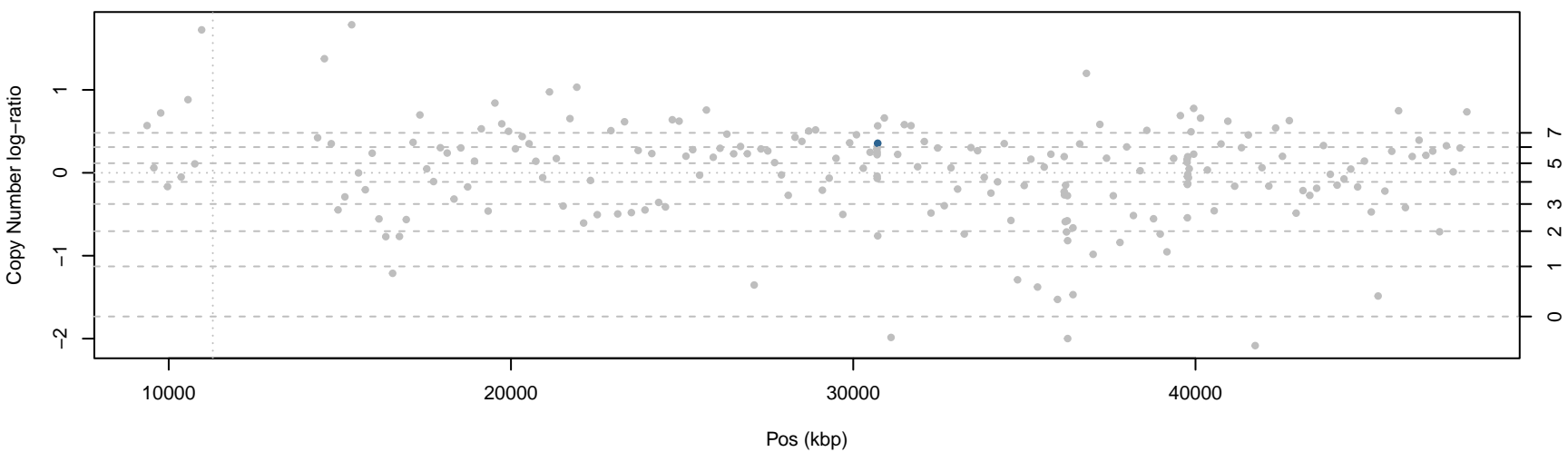
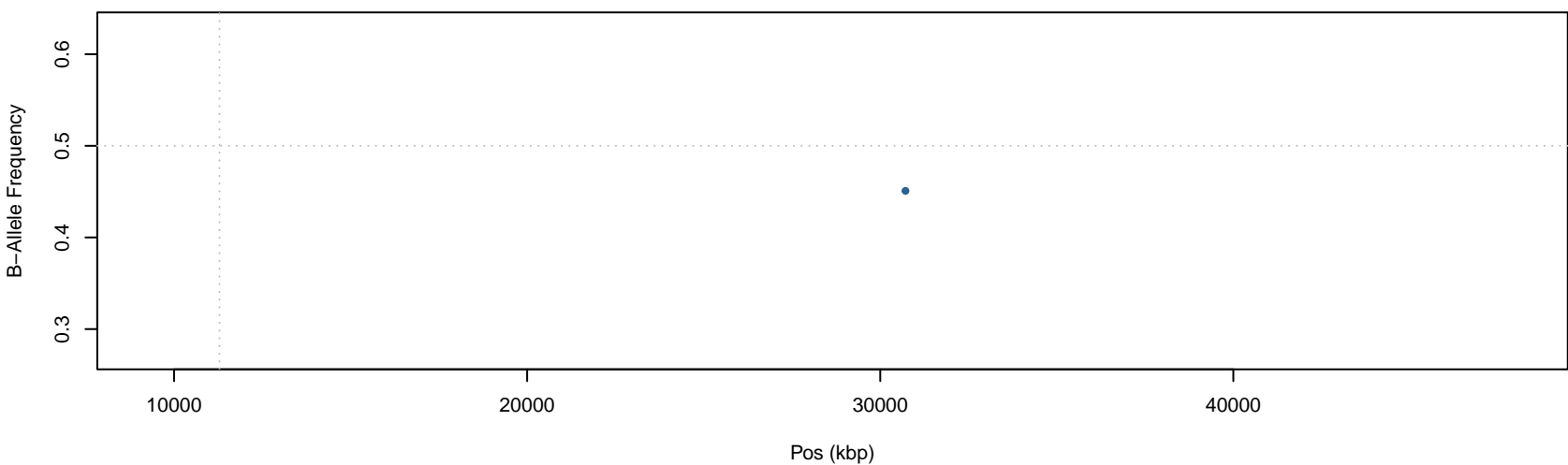




Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 20



Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 21



Purity: 0.51 Tumor ploidy: 4.467 SNV log-likelihood: -258.99 GoF: 84.8% Mean coverage: 465;225 Chromosome: 22

