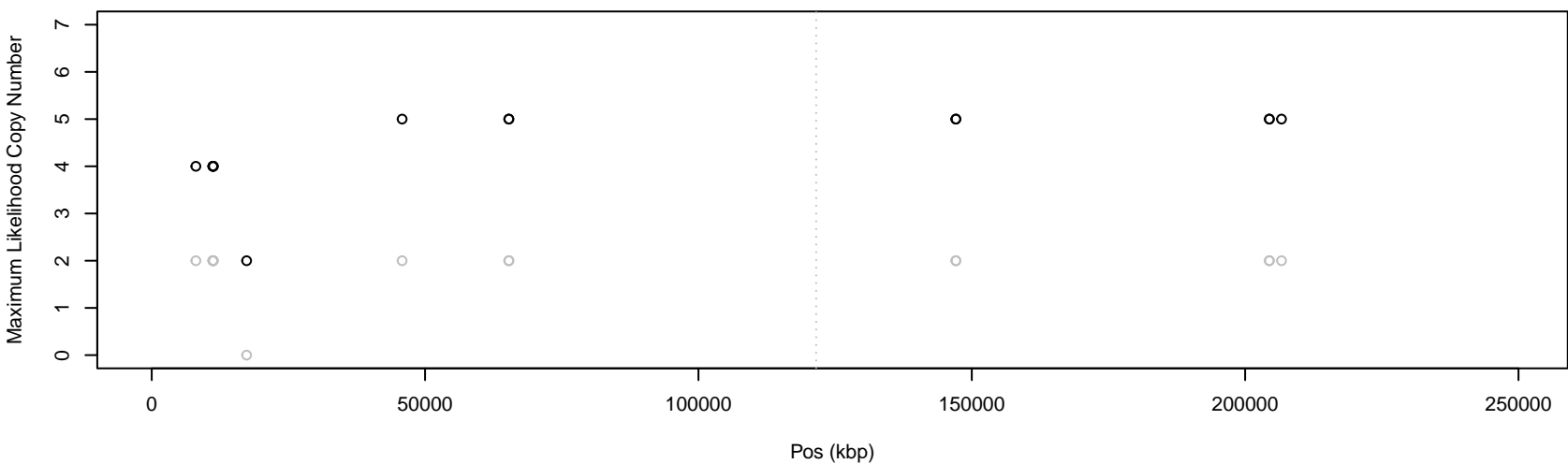
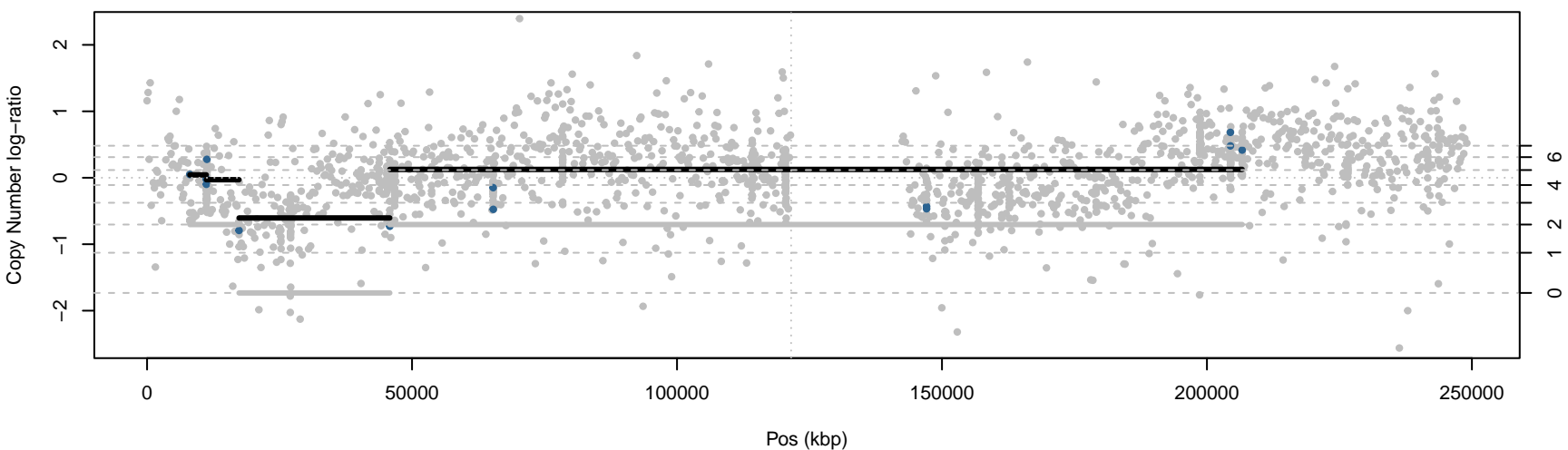
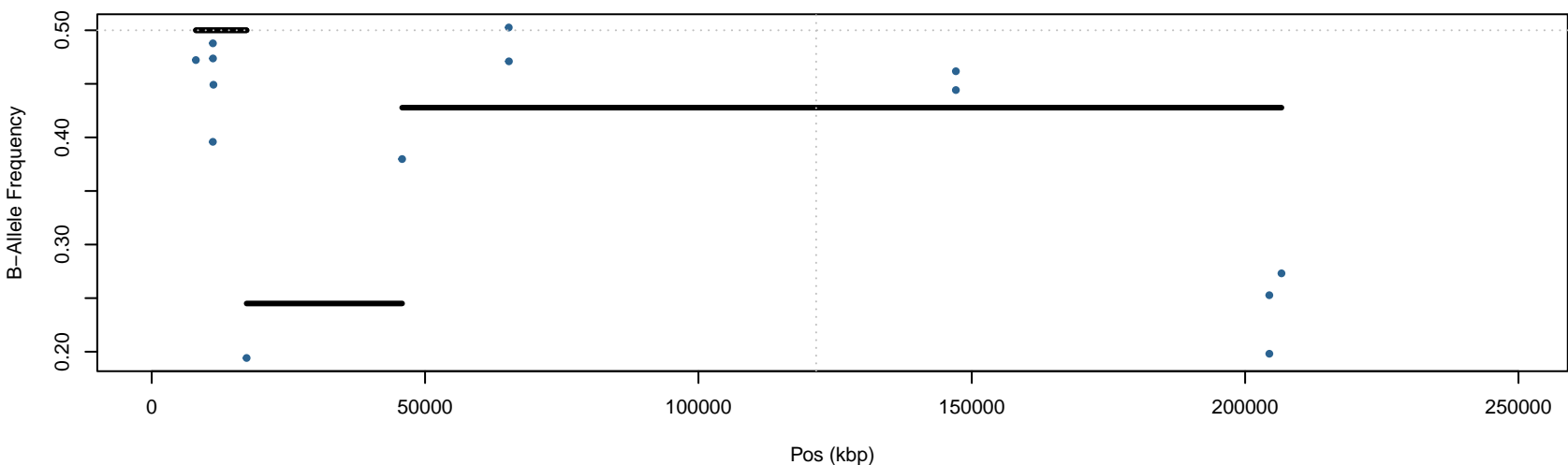
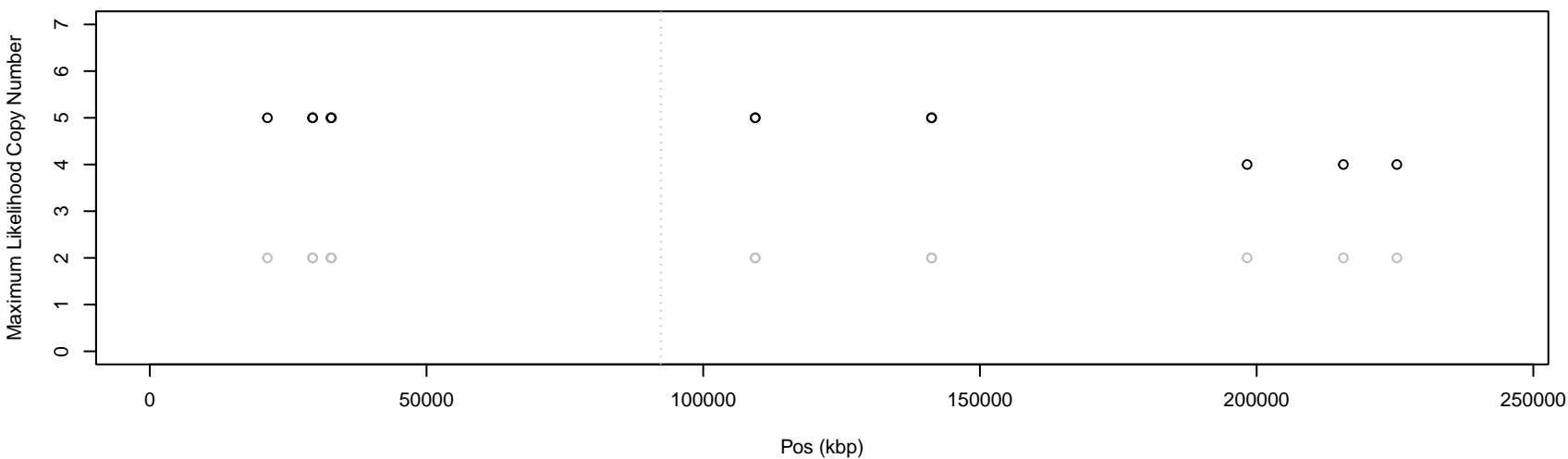
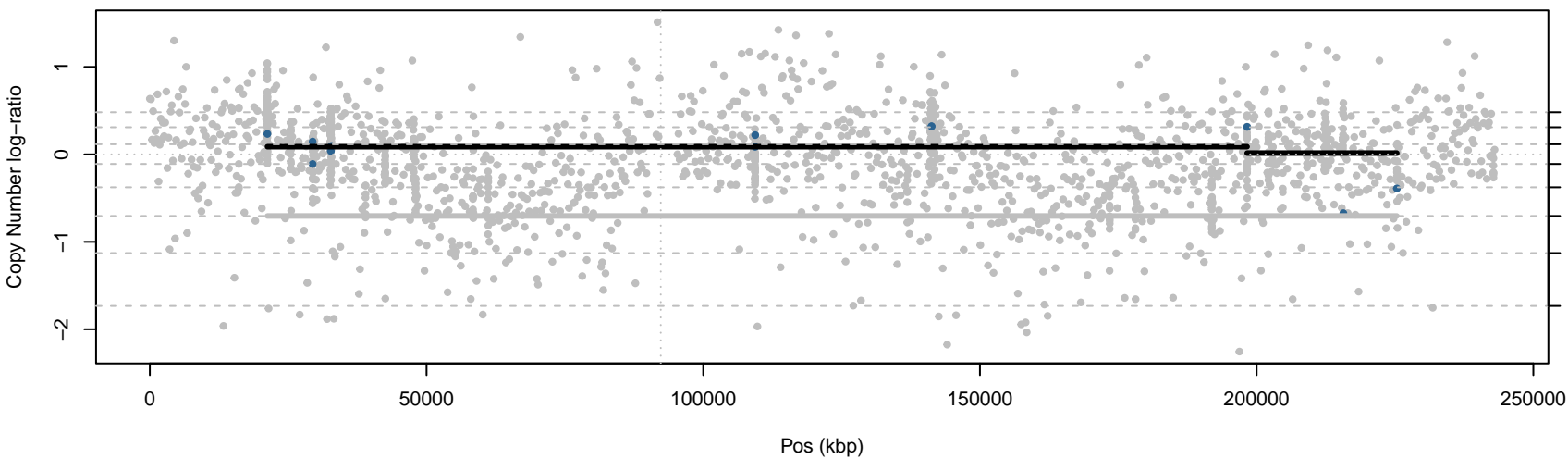
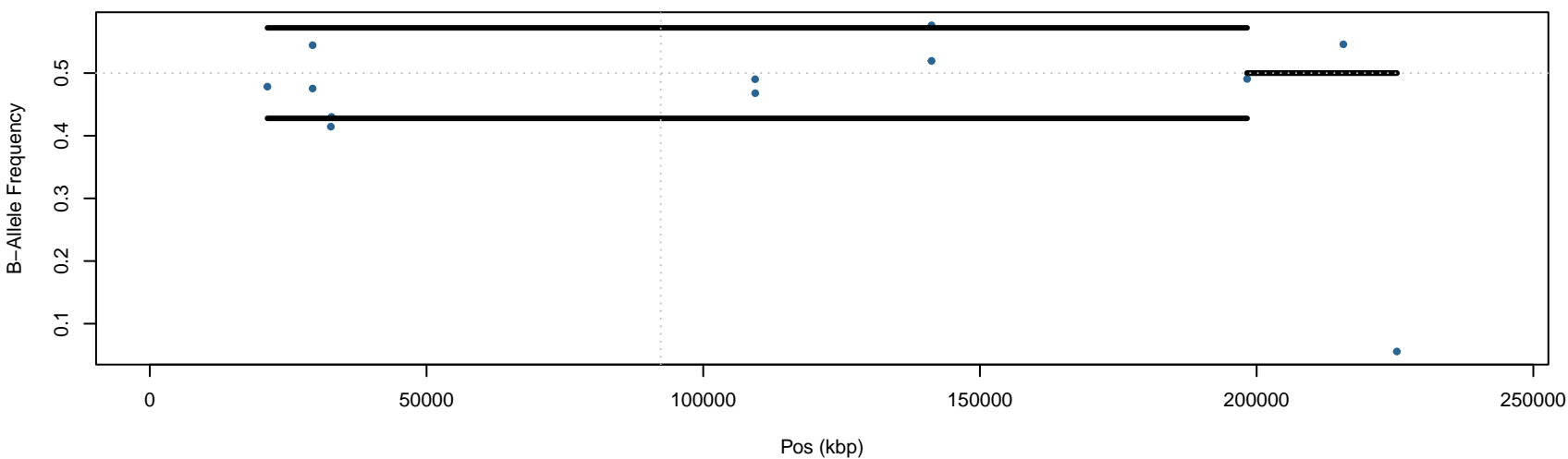


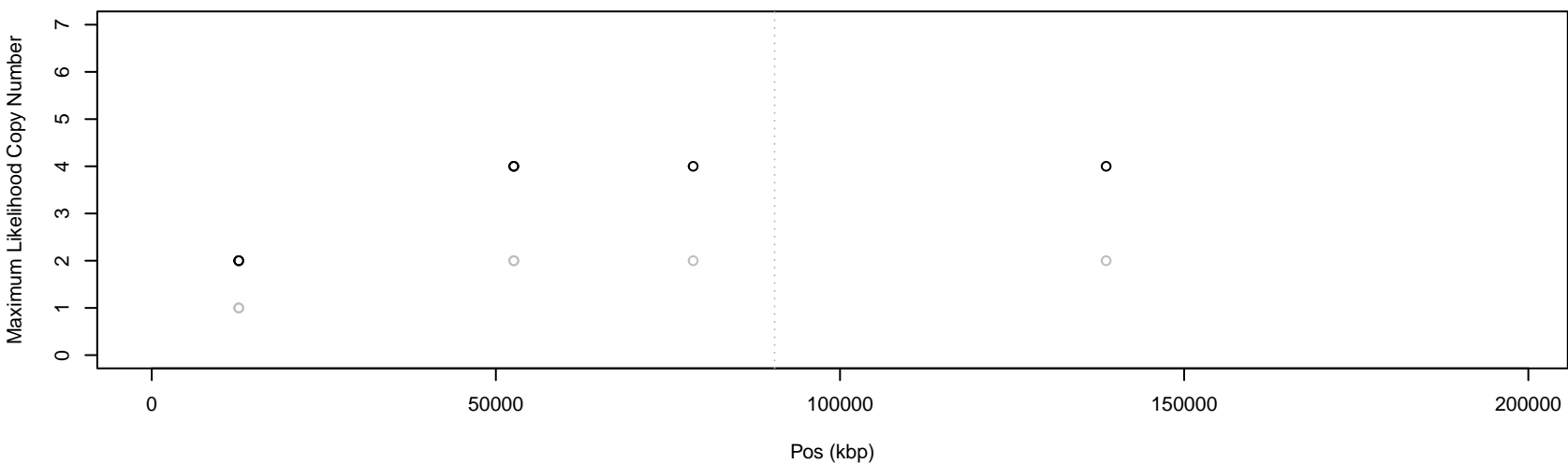
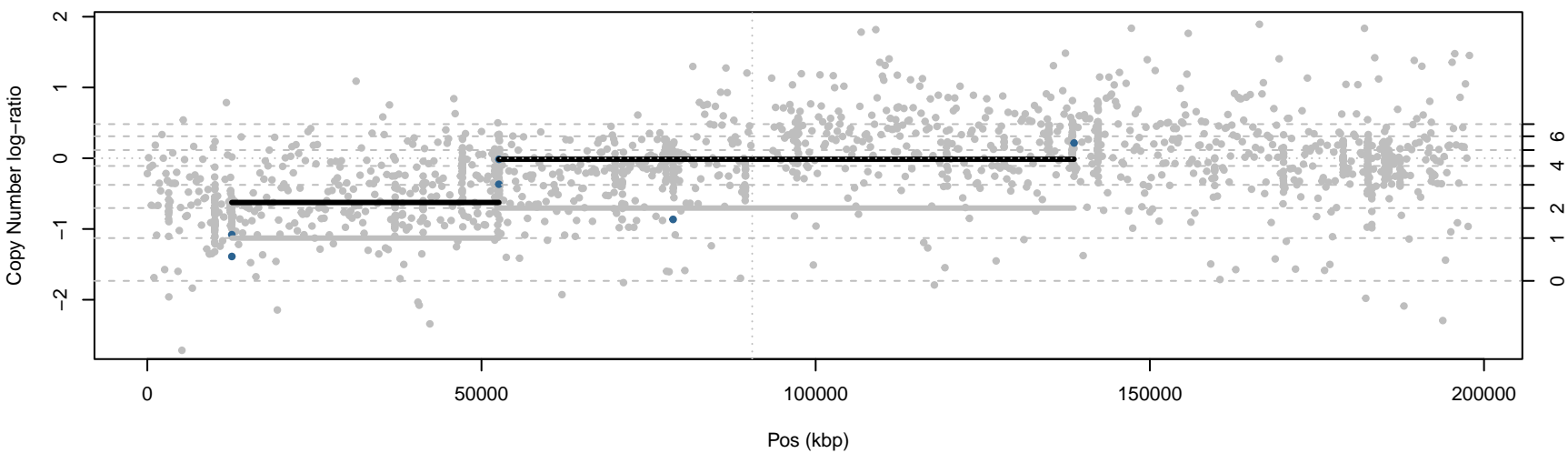
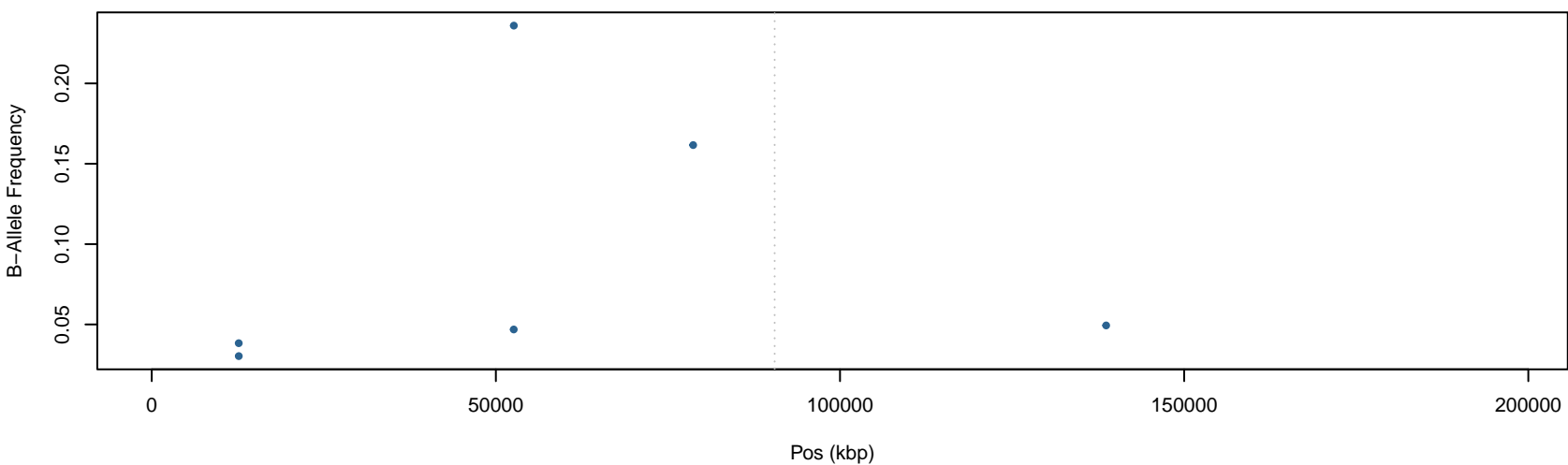
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 1



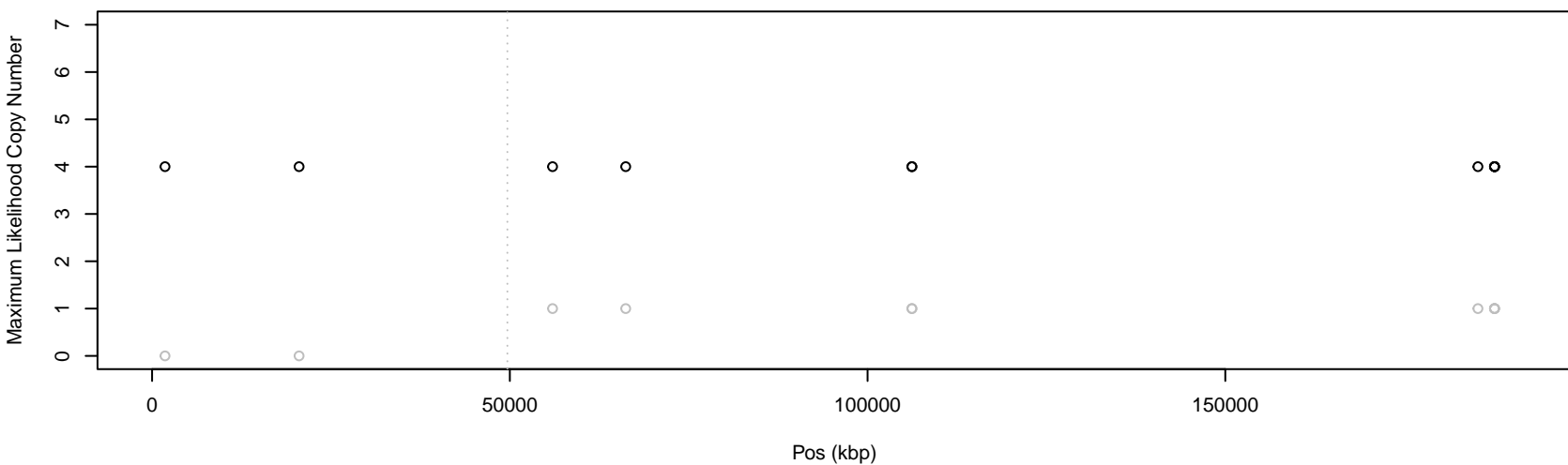
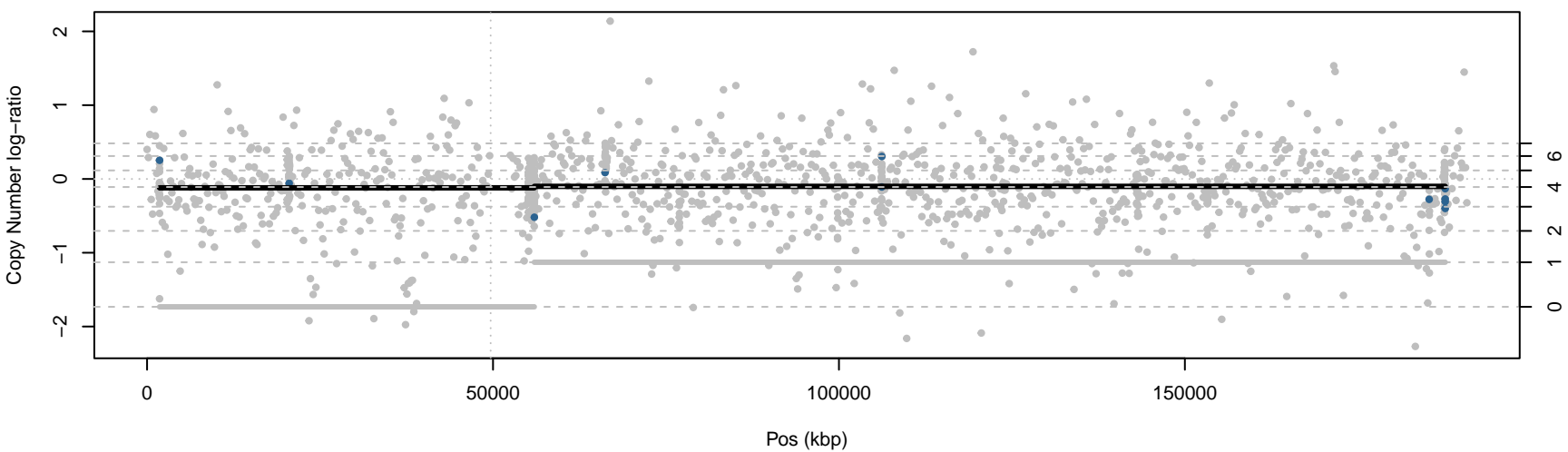
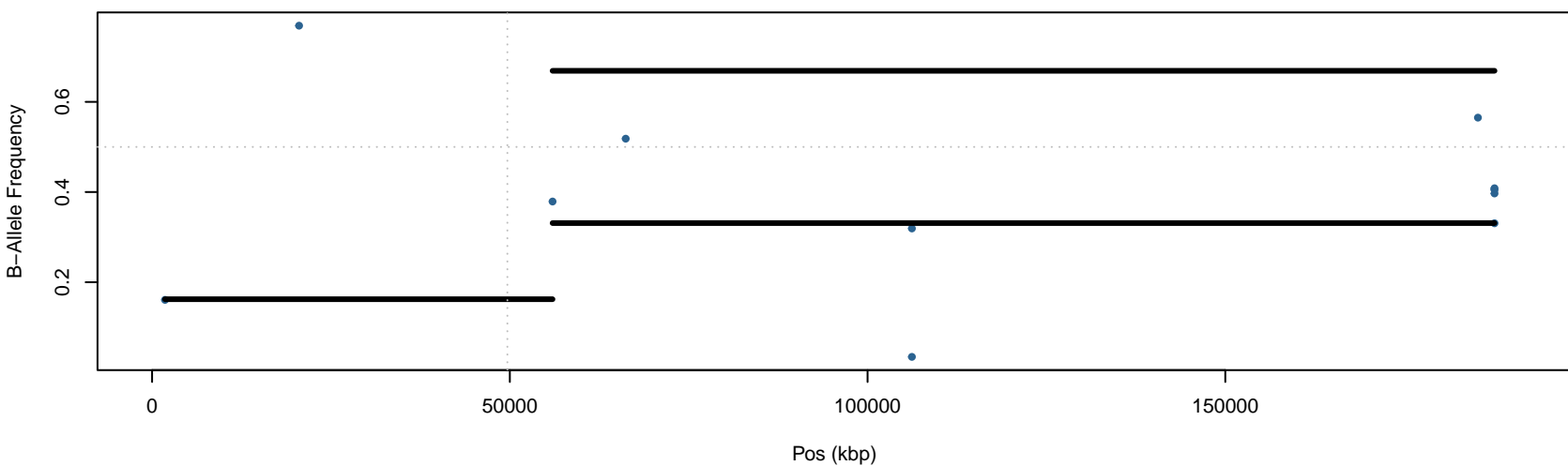
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 2



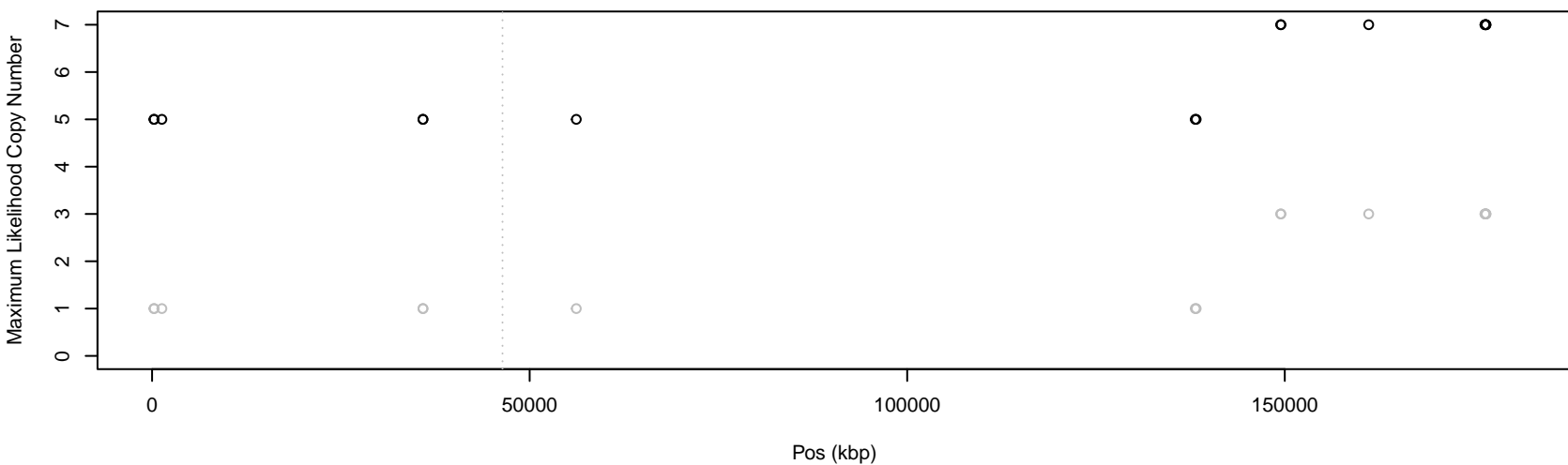
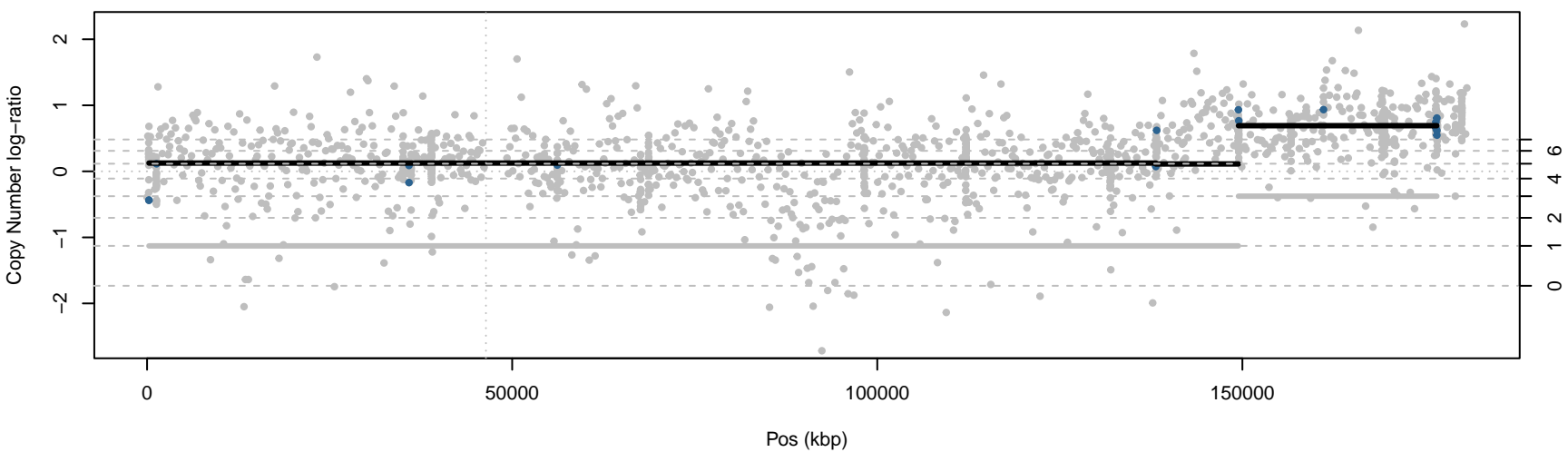
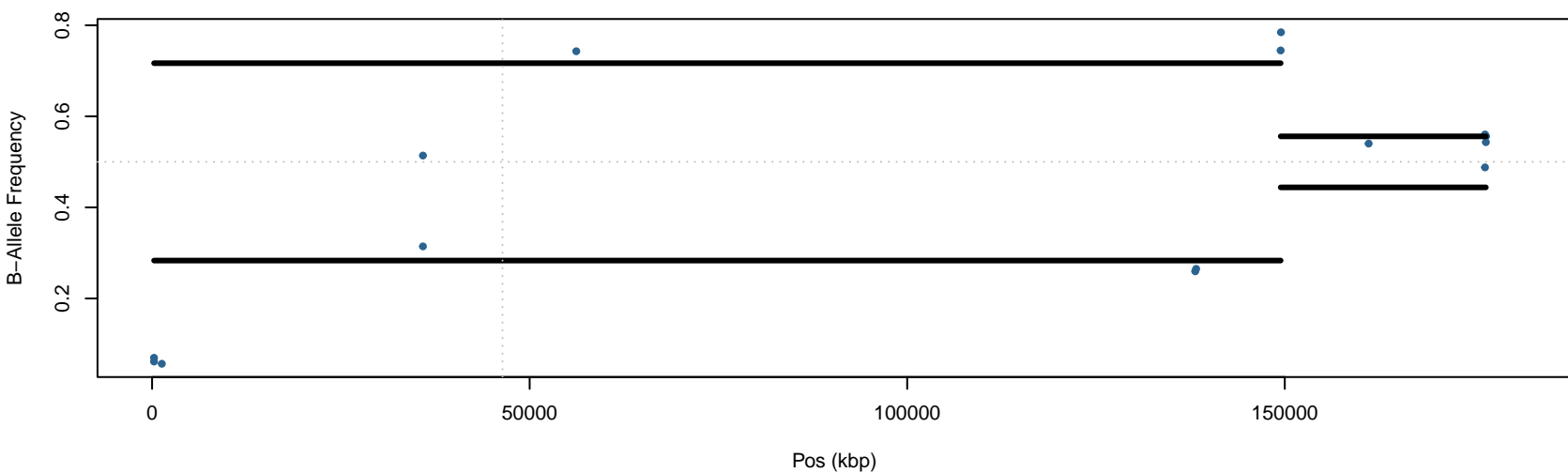
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 3



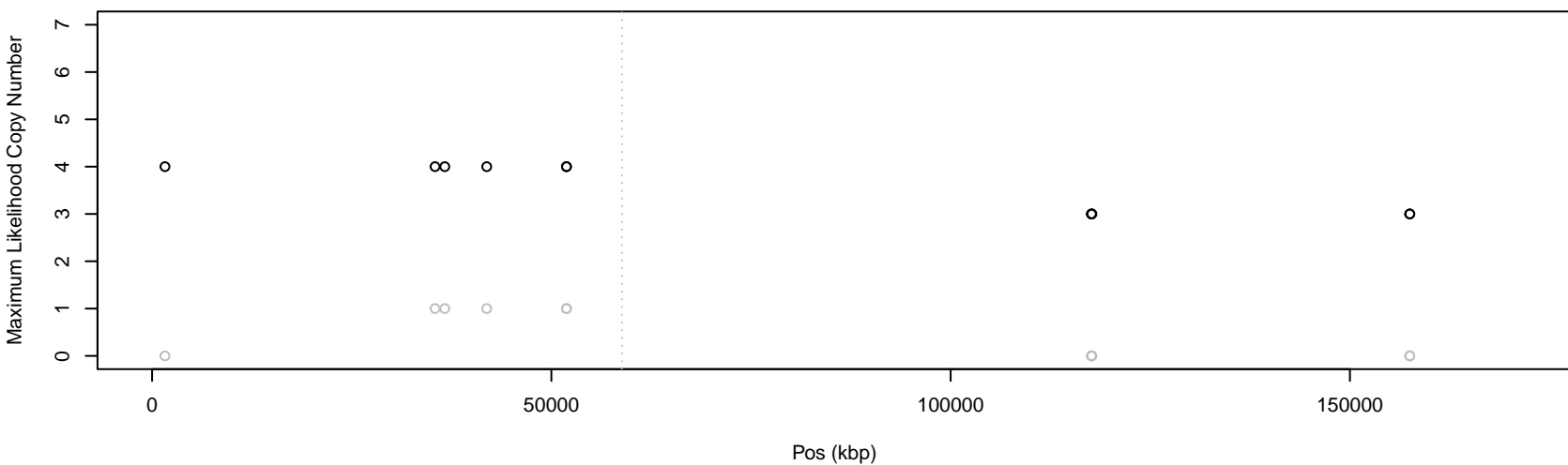
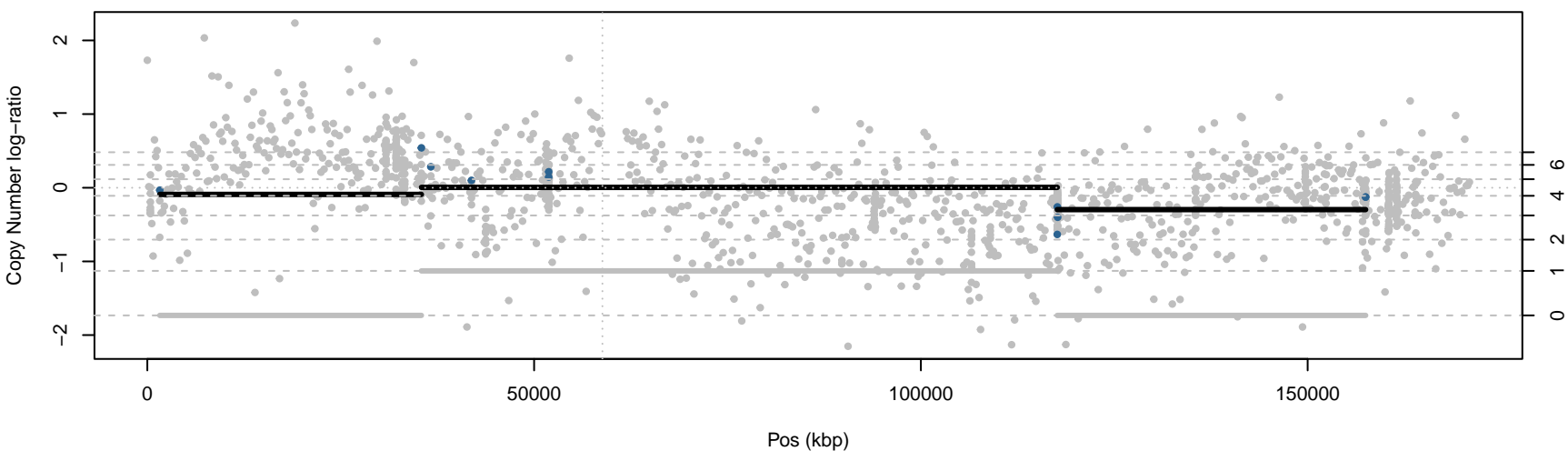
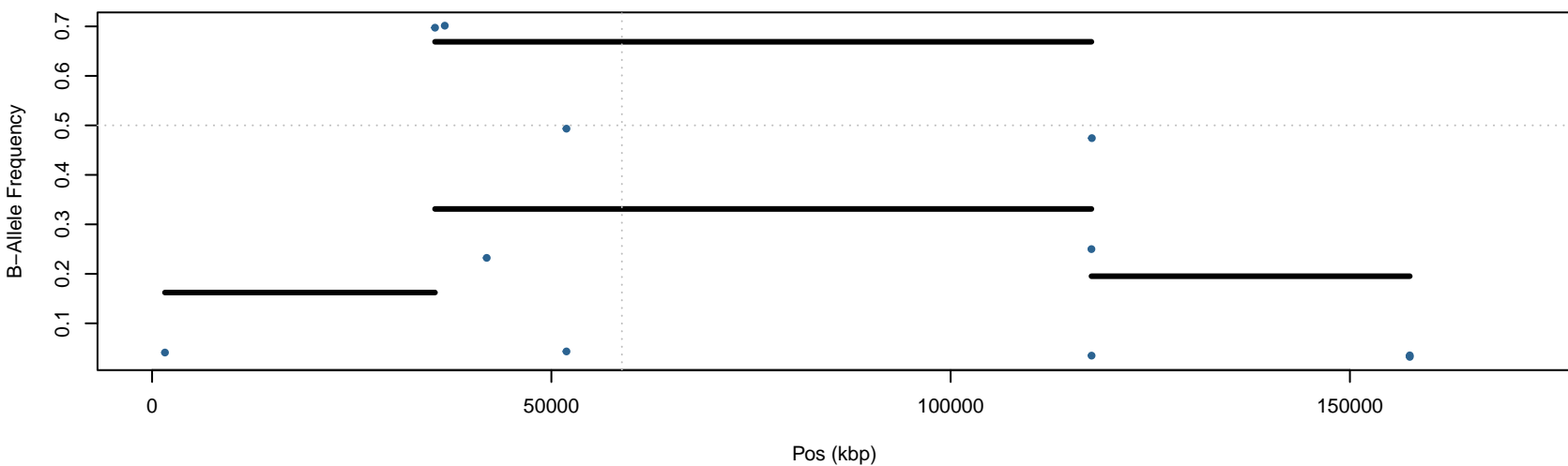
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 4



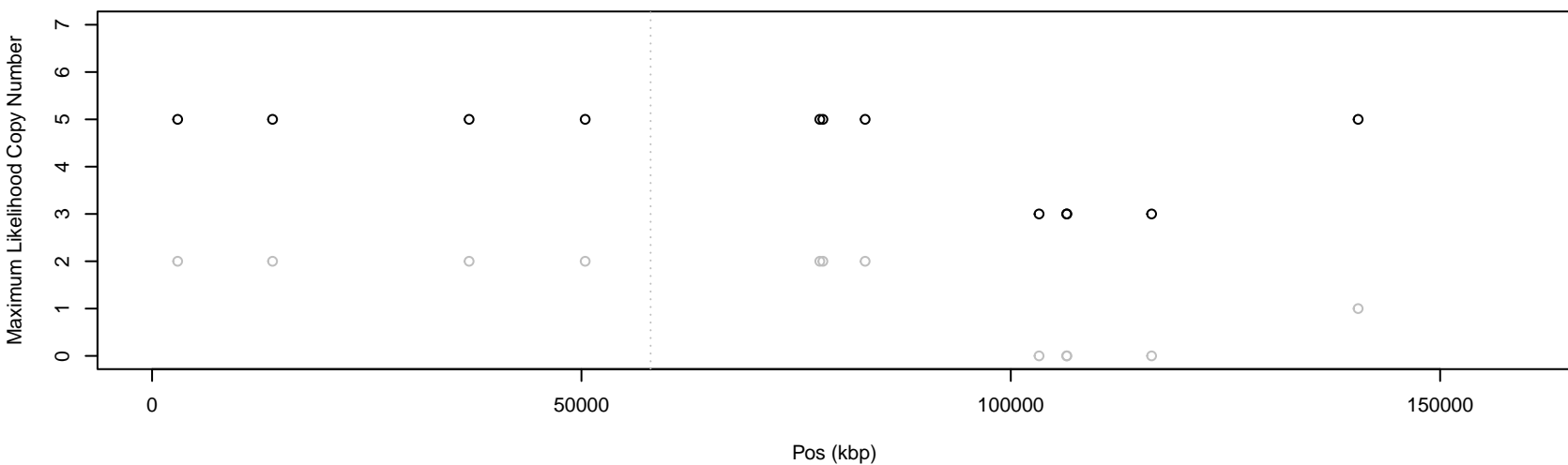
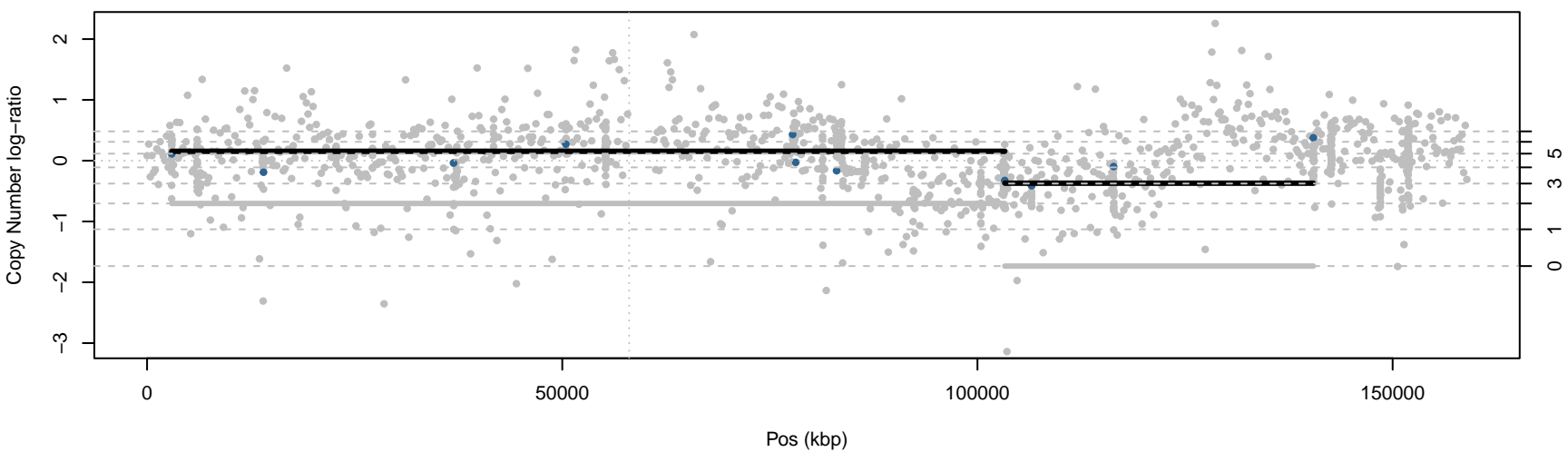
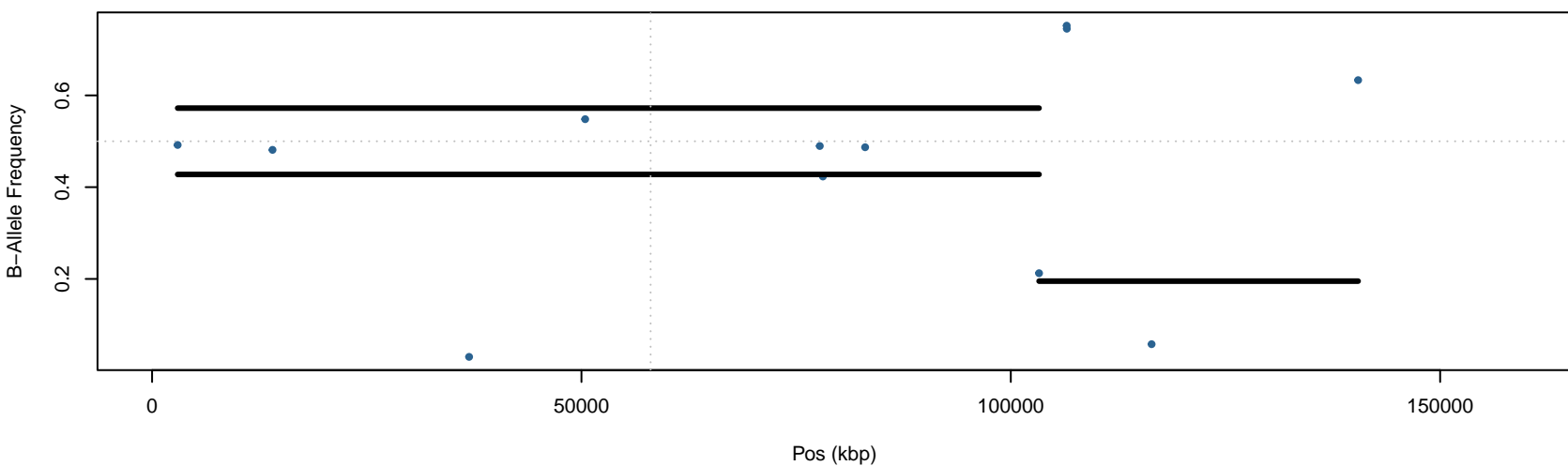
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 5



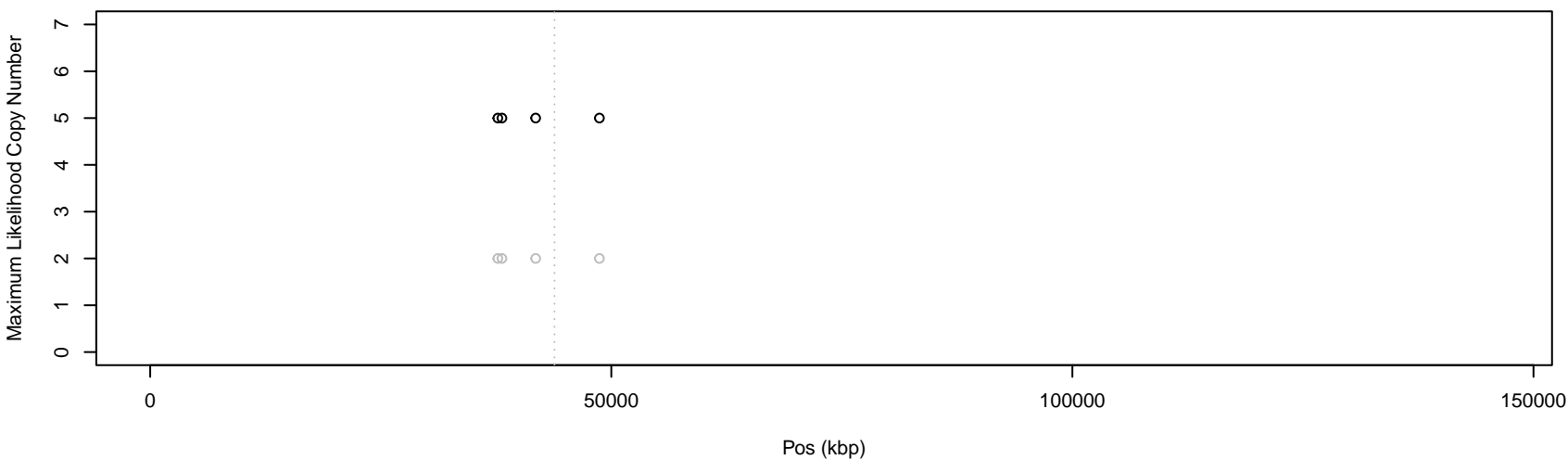
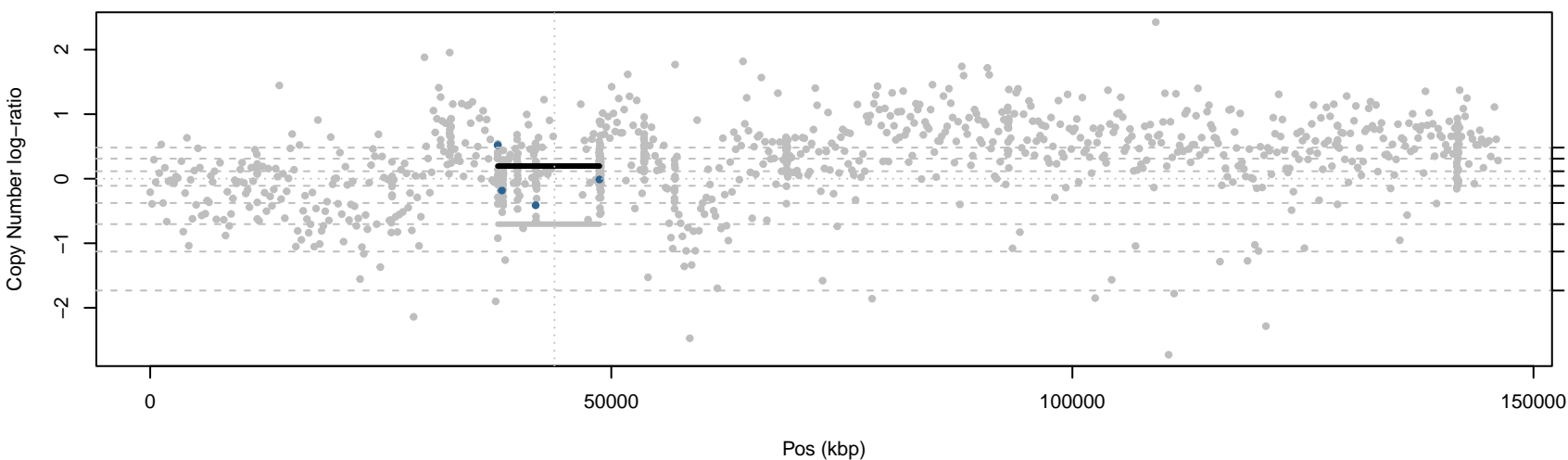
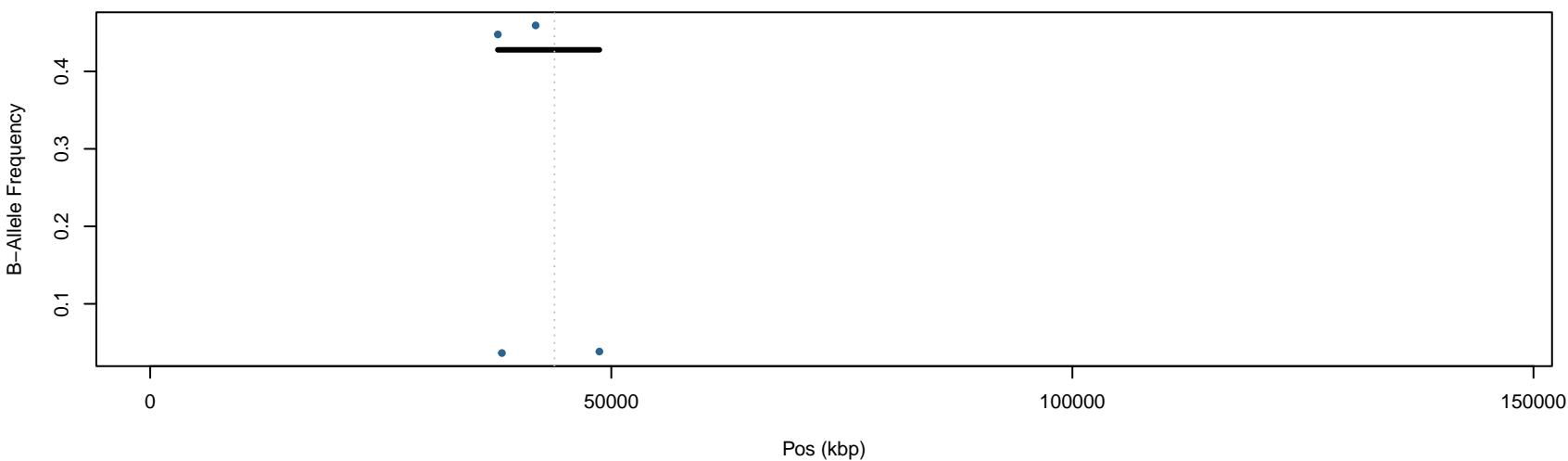
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 6



Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 7

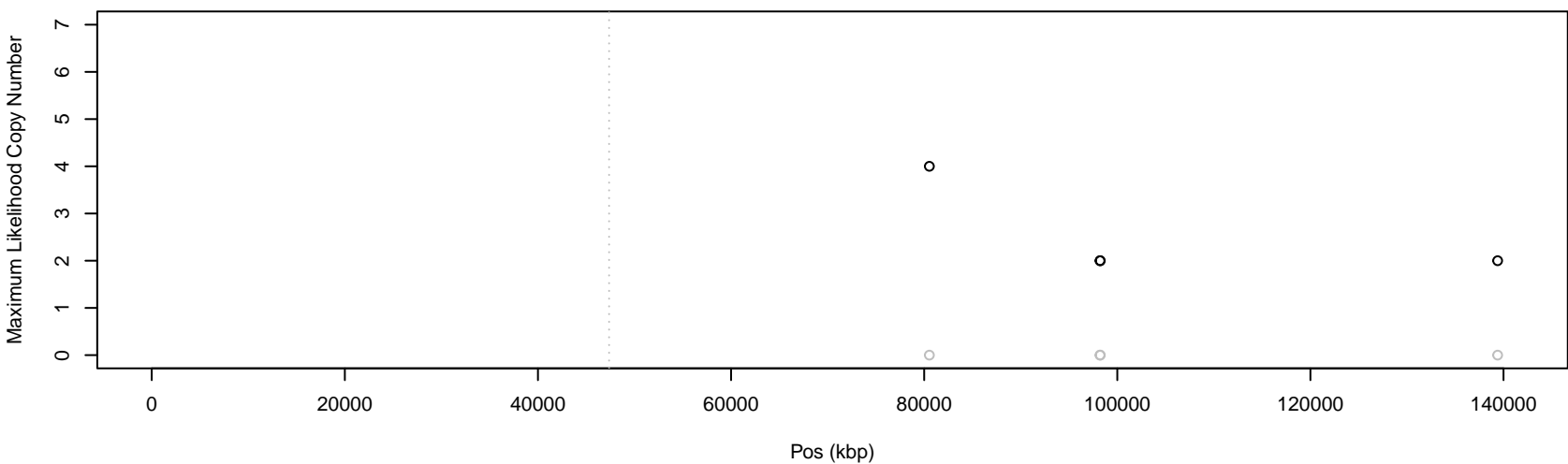
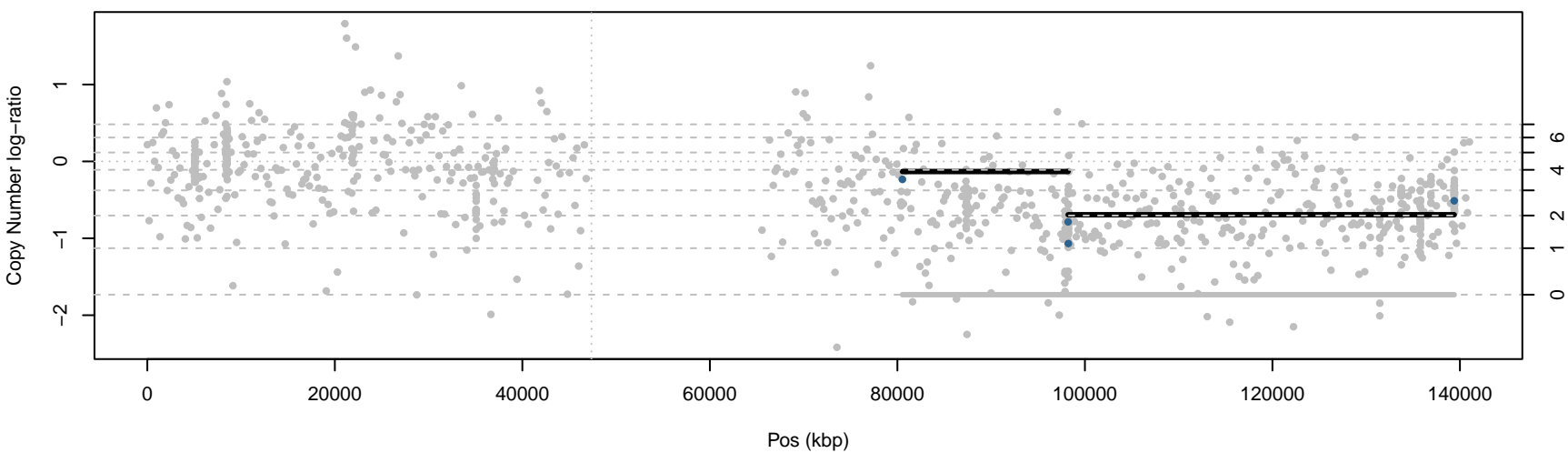
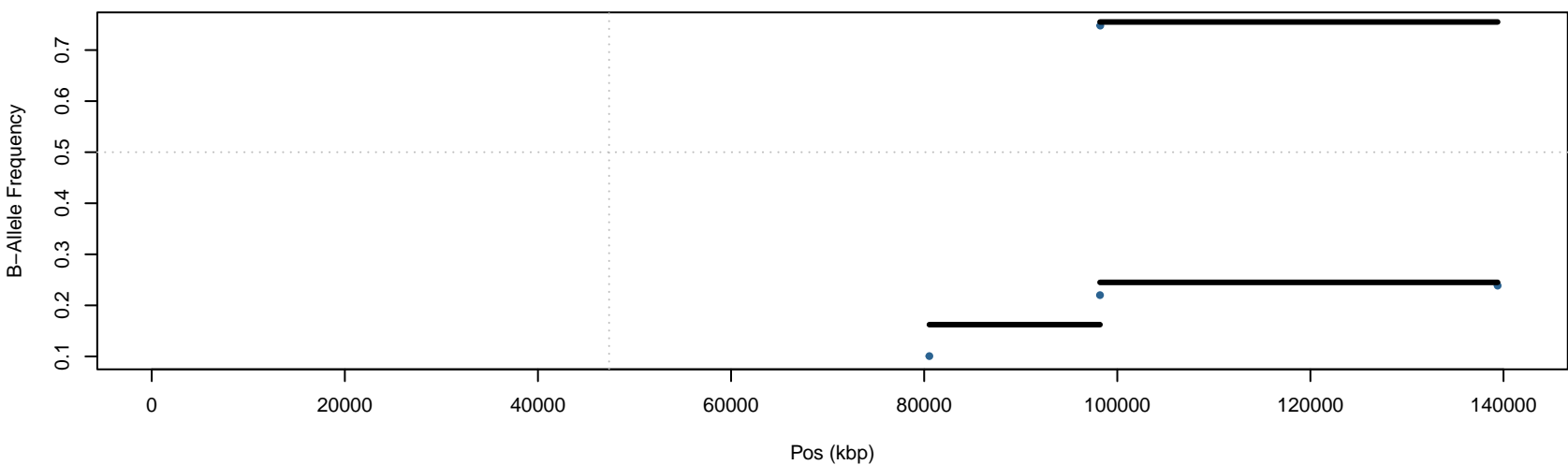


Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 8

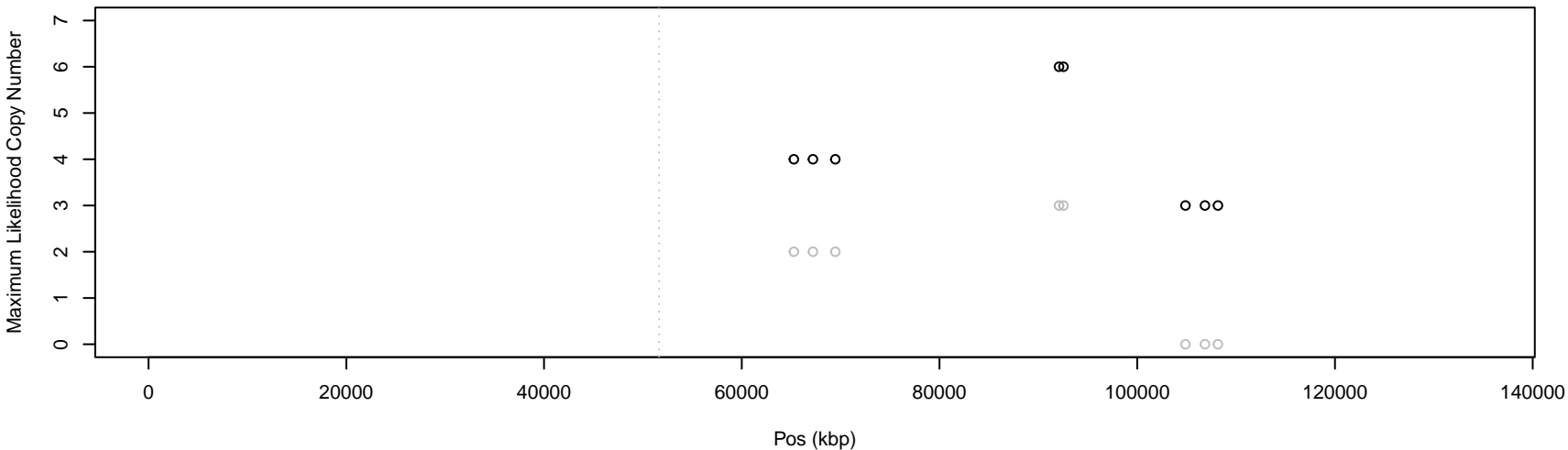
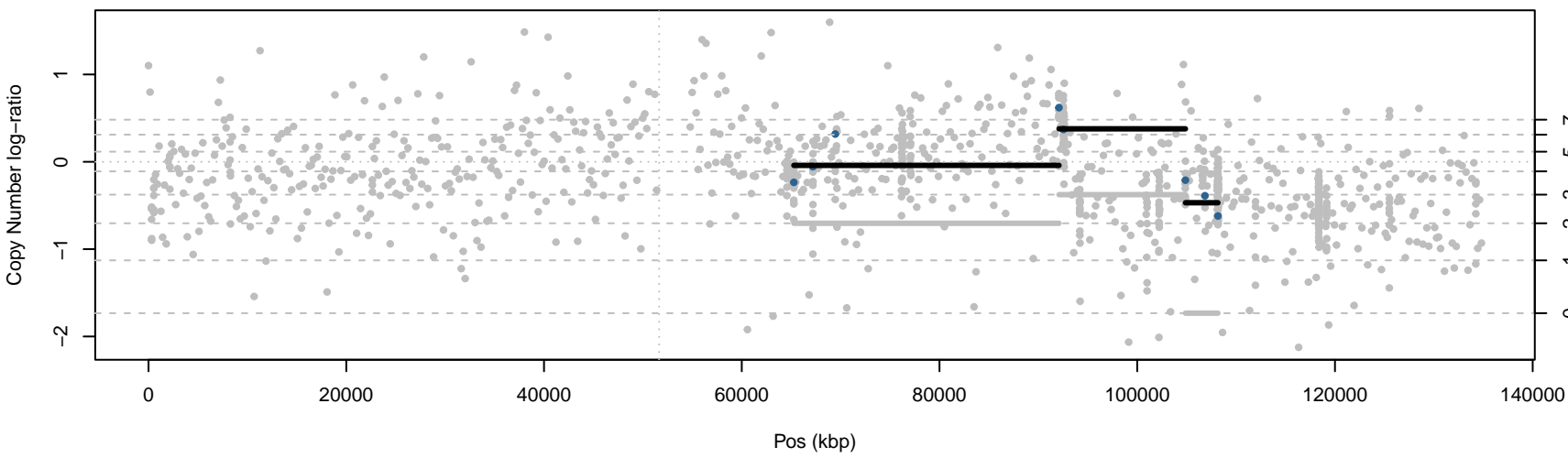
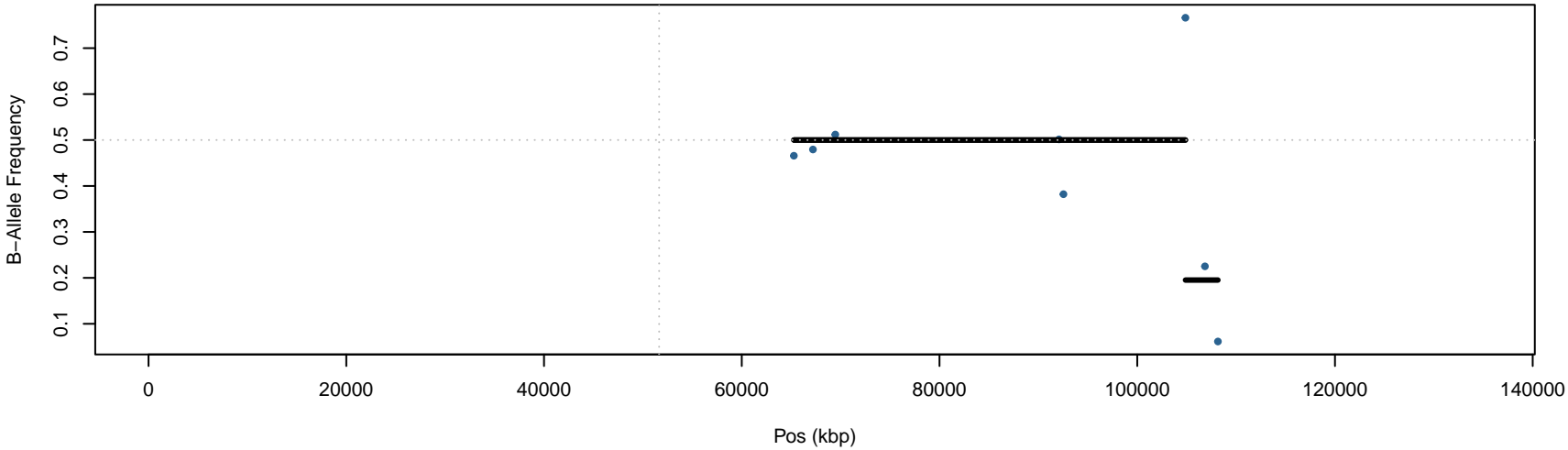




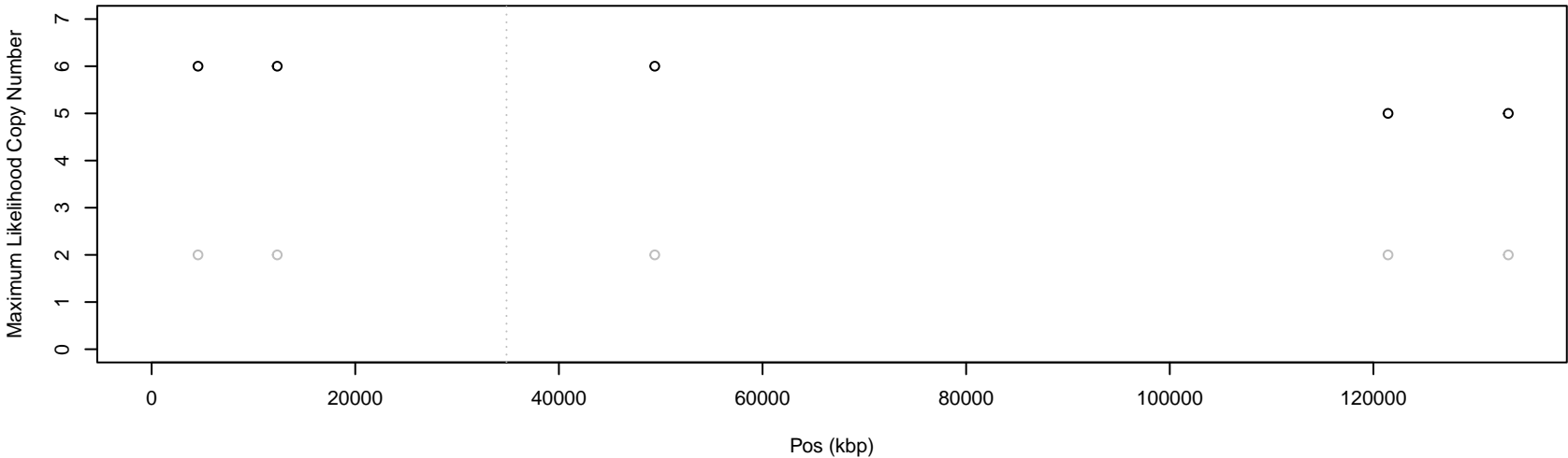
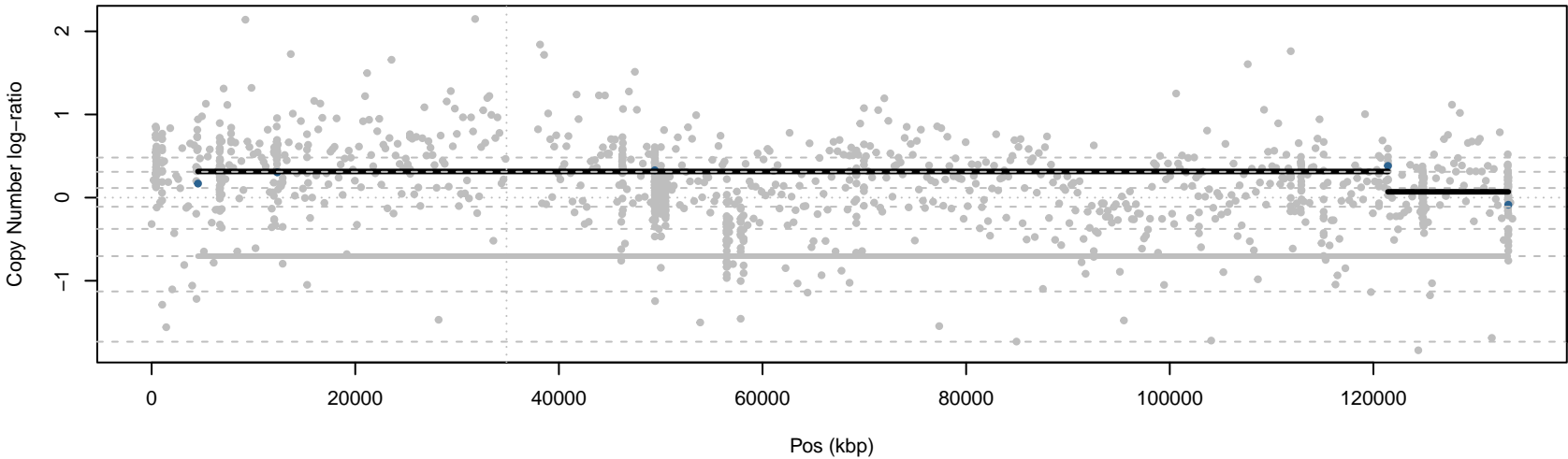
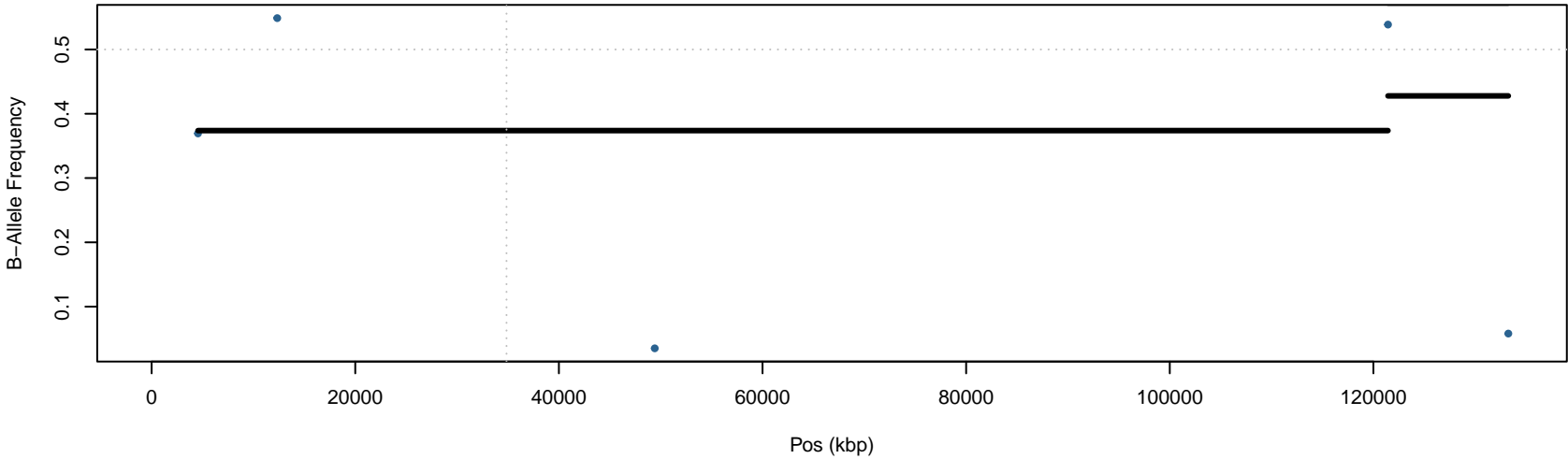
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 9

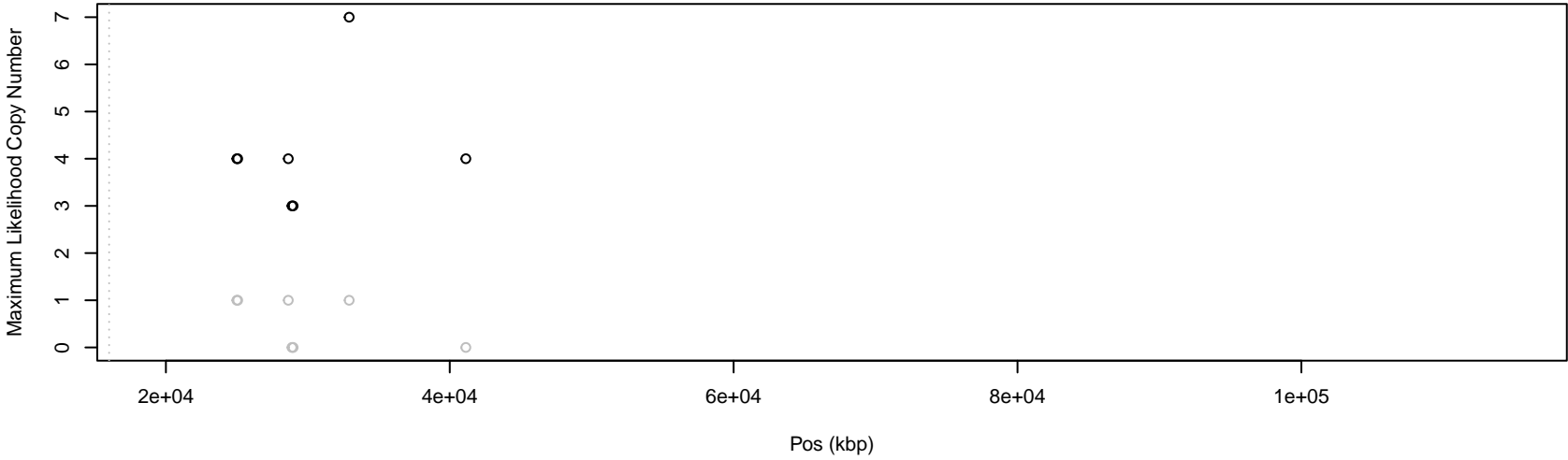
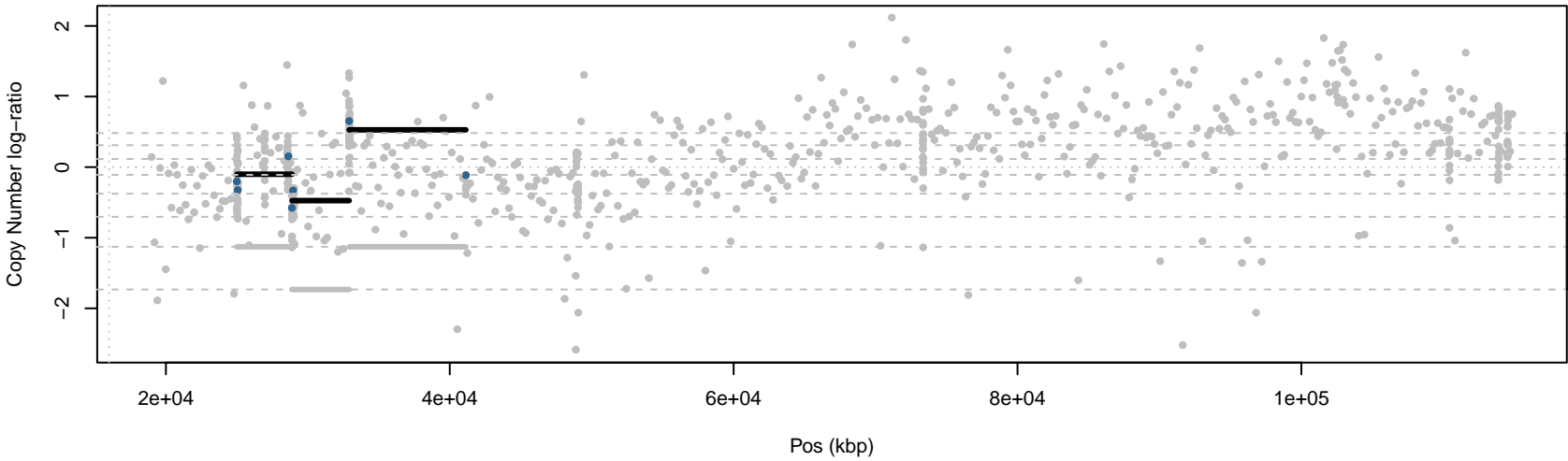
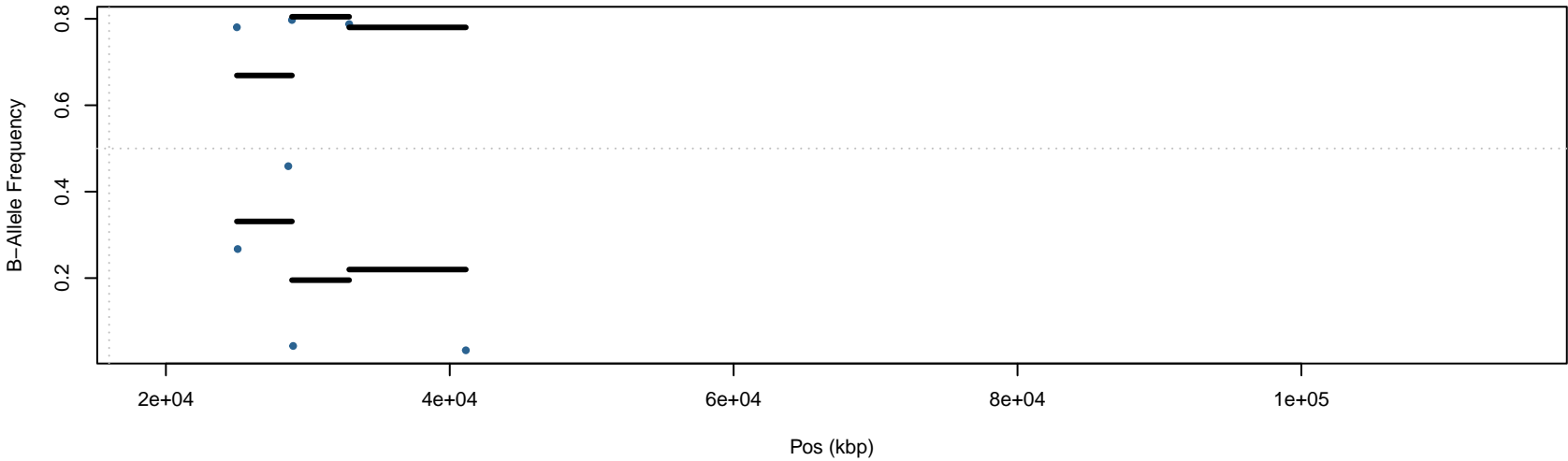


Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 11

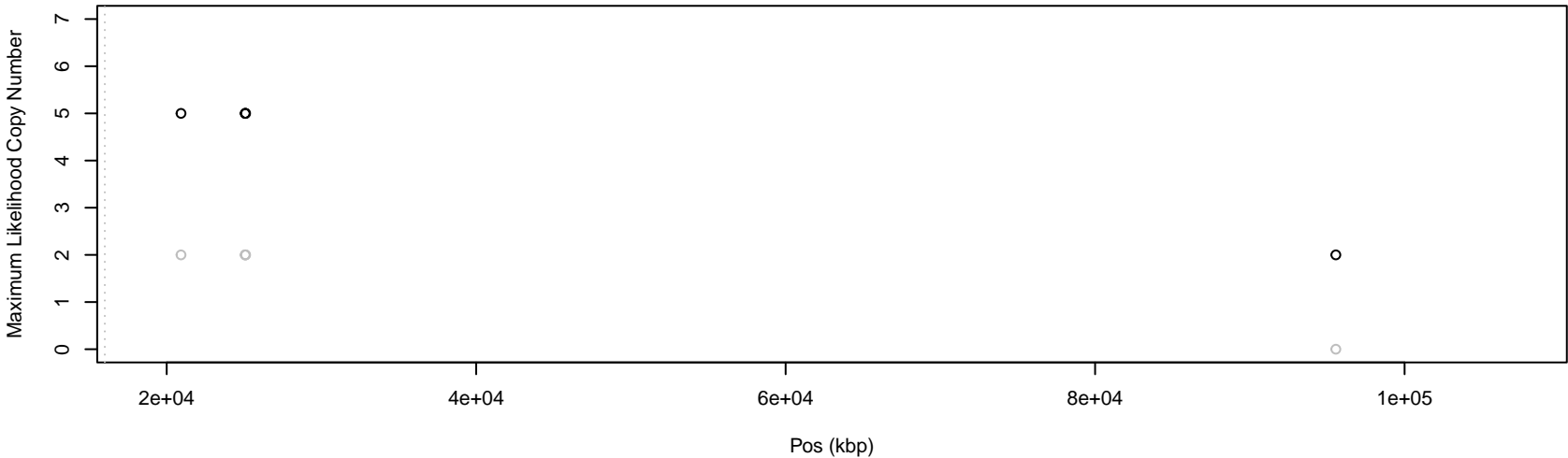
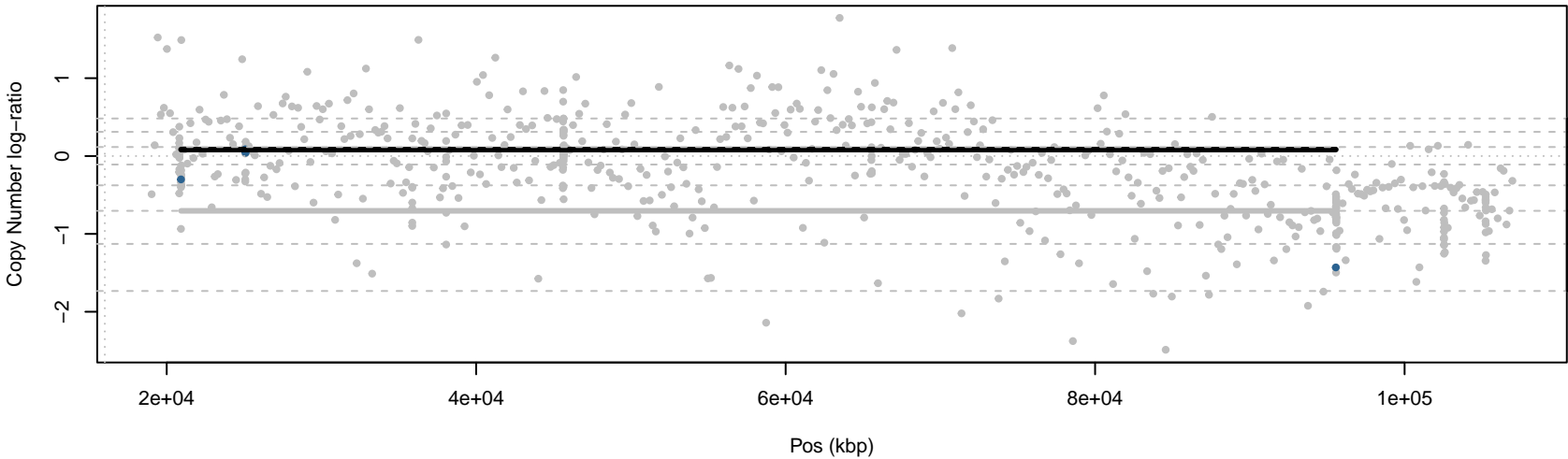
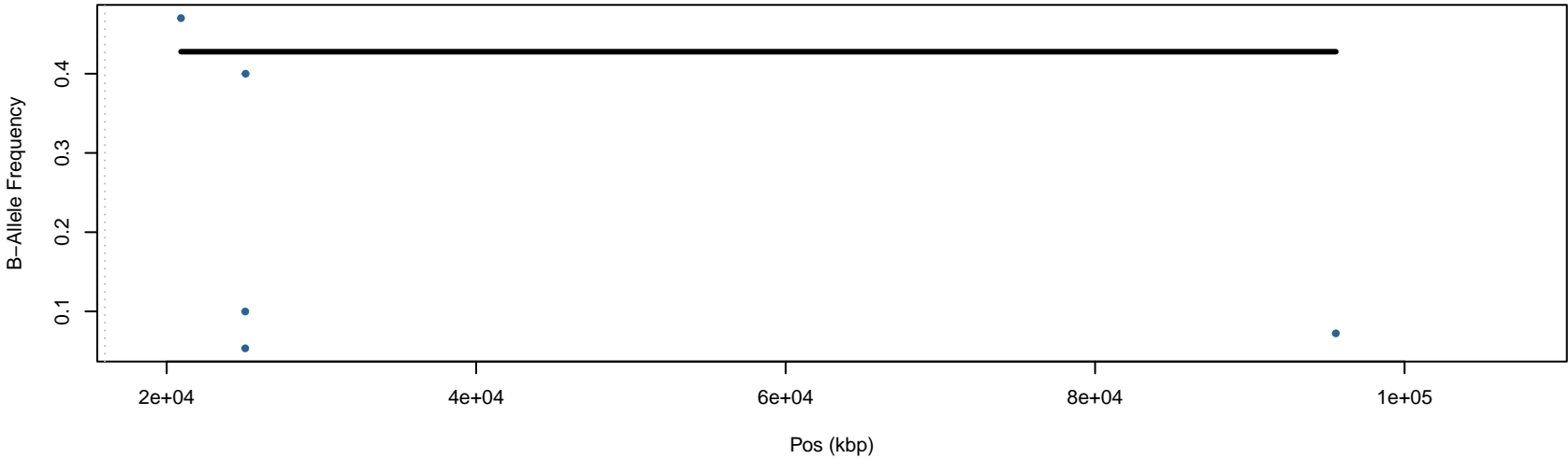


Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 12

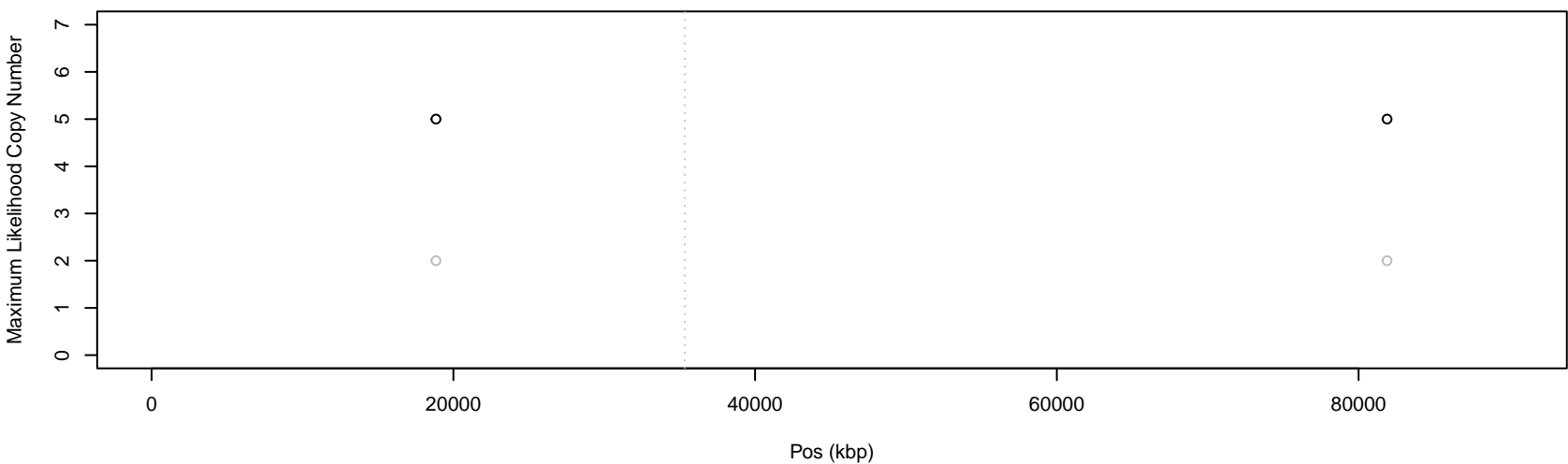
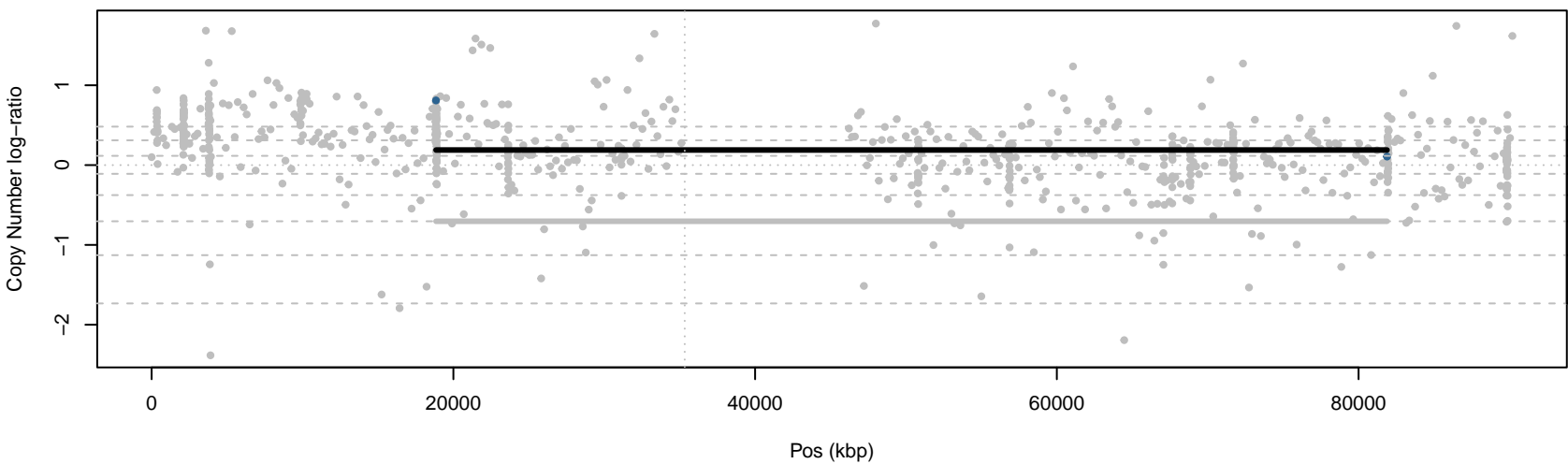
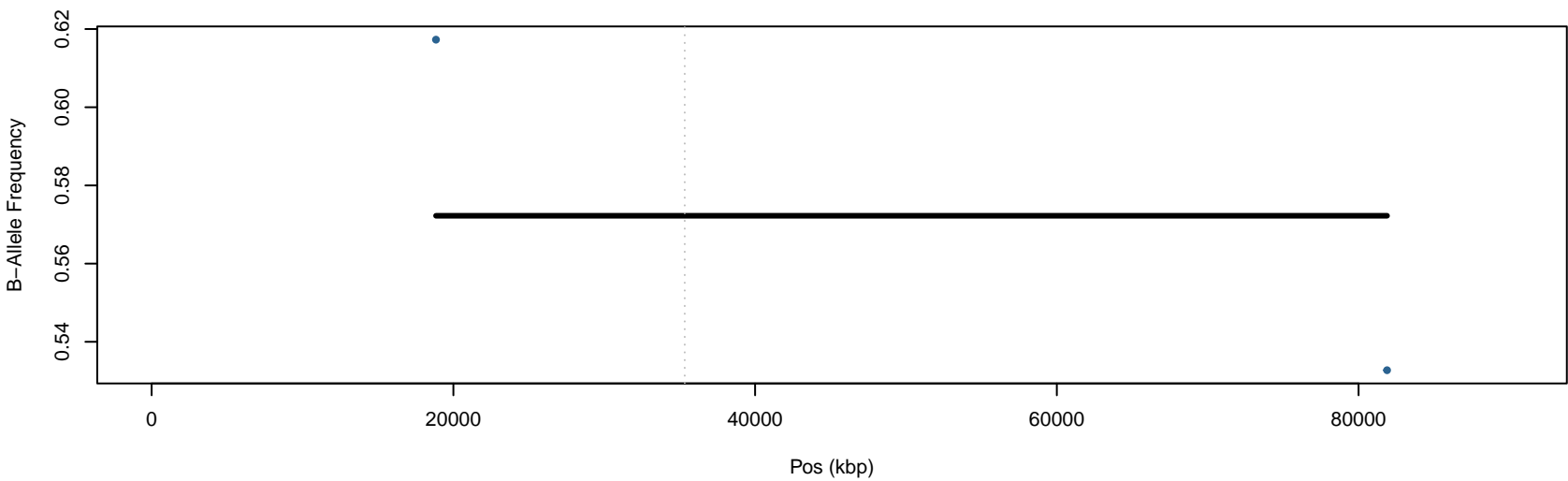


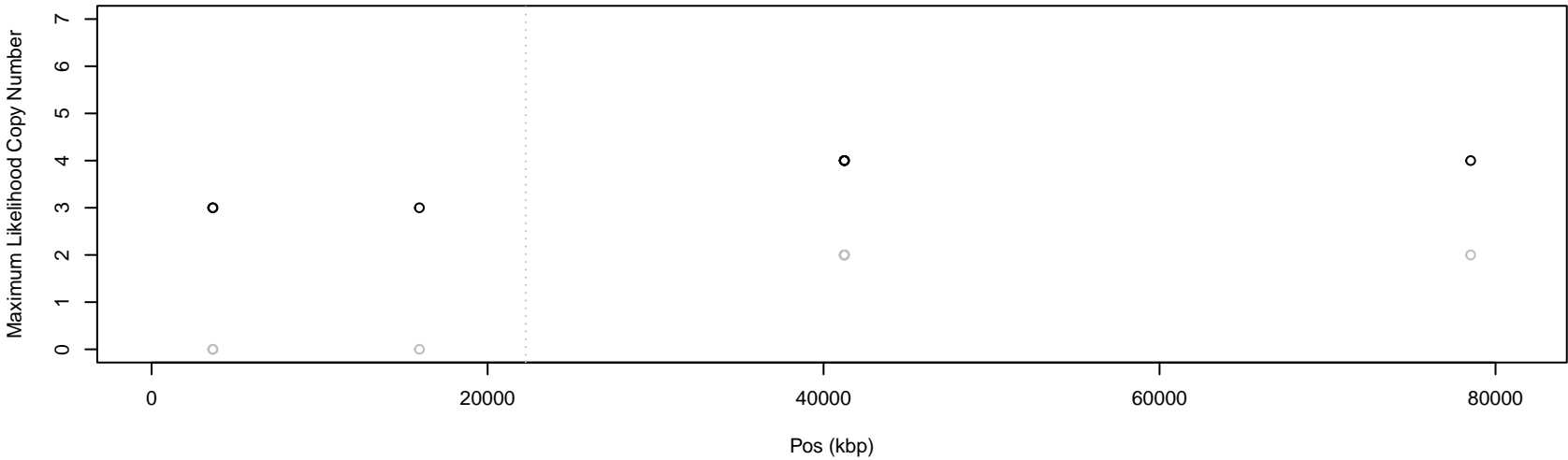
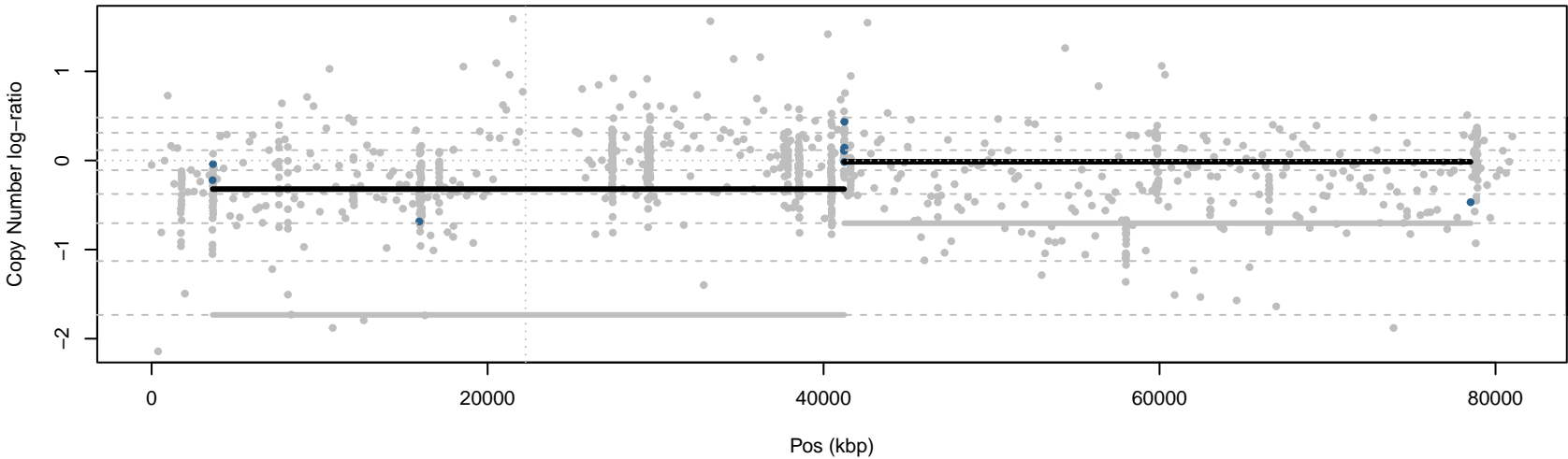
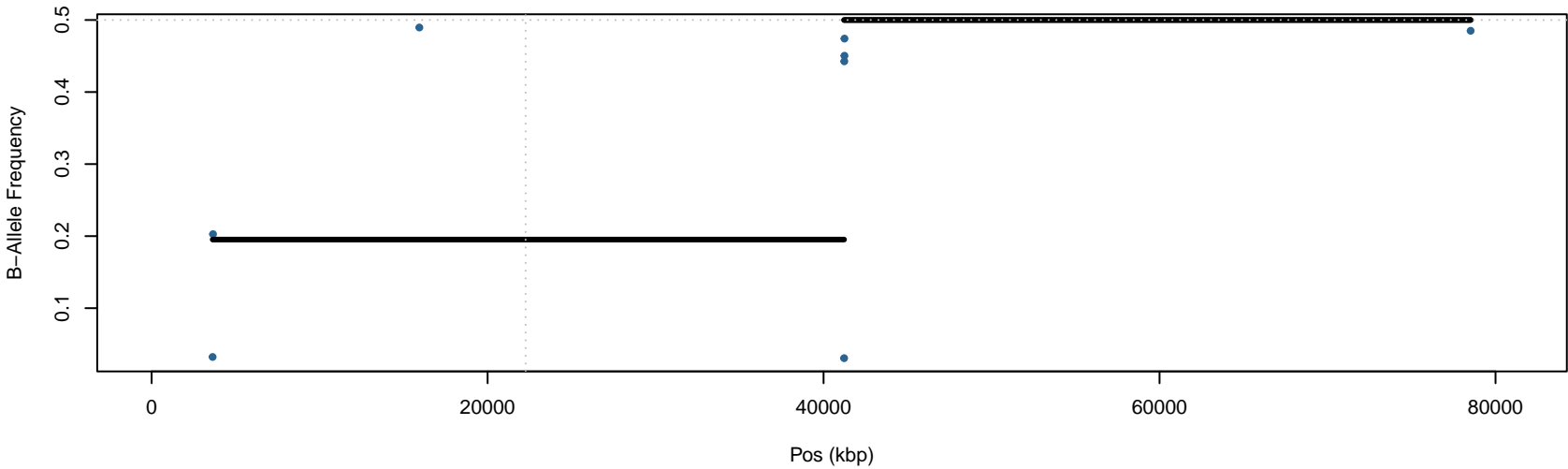


Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 14

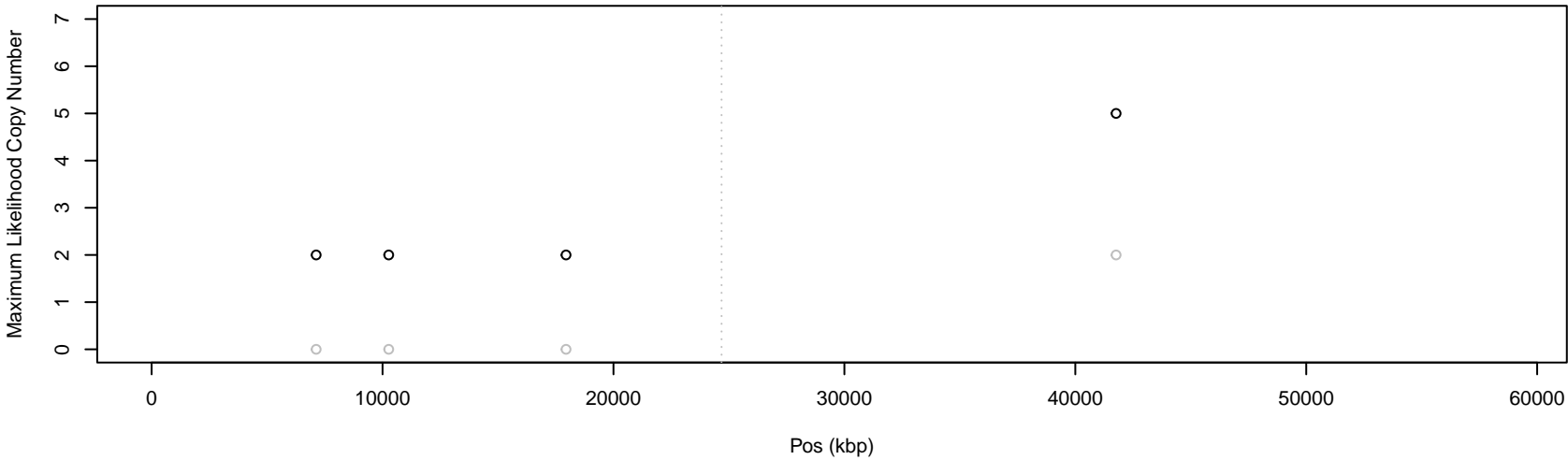
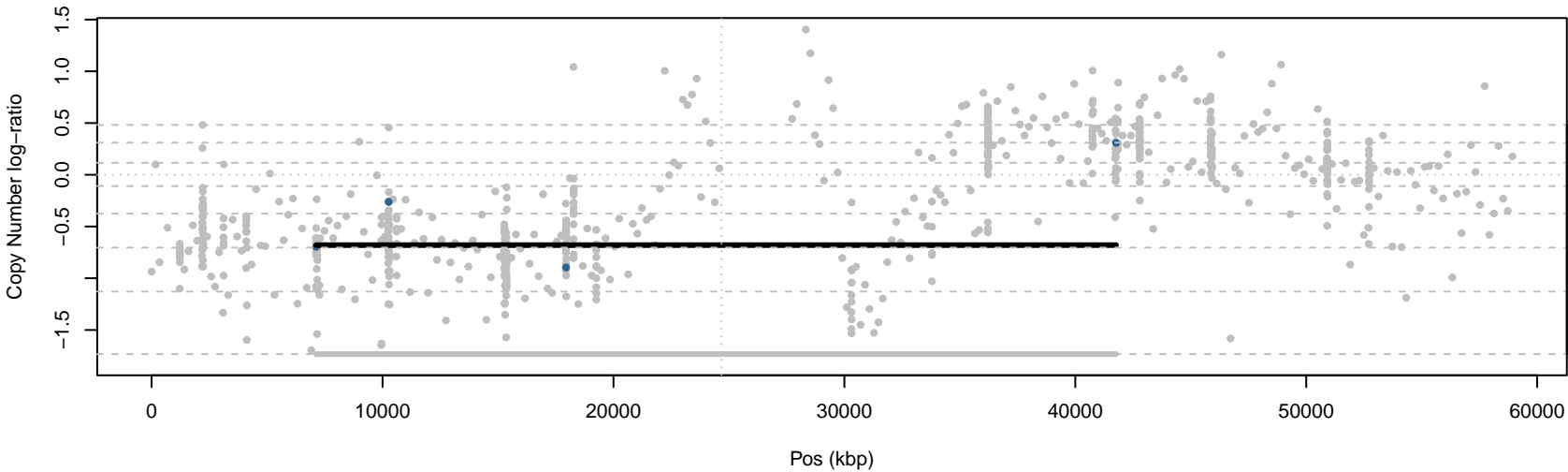
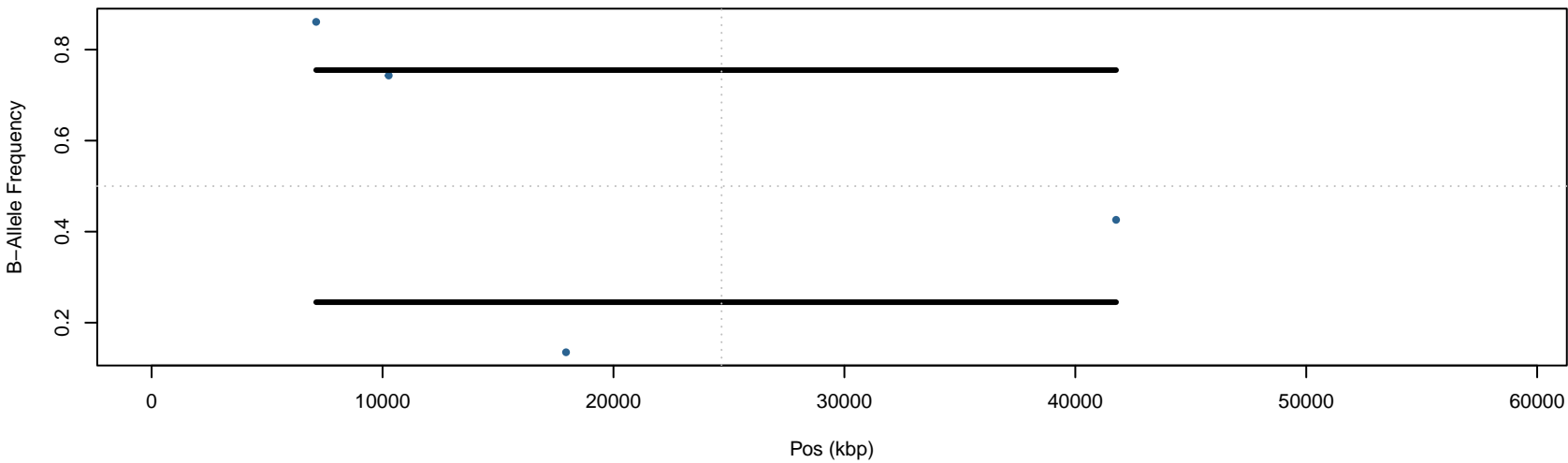


Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 16



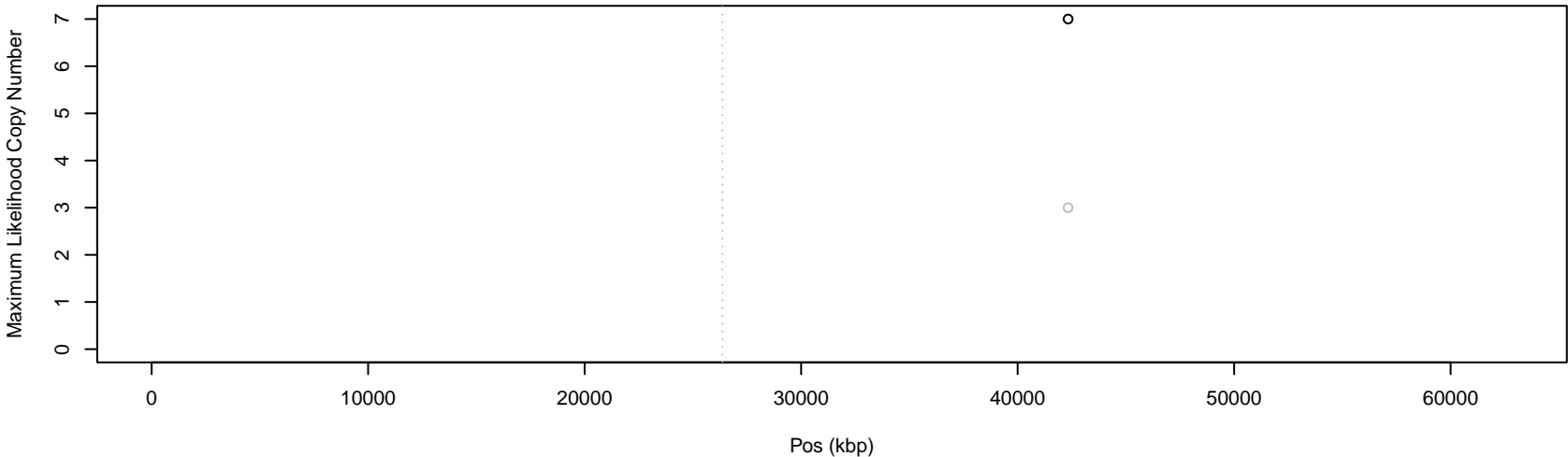
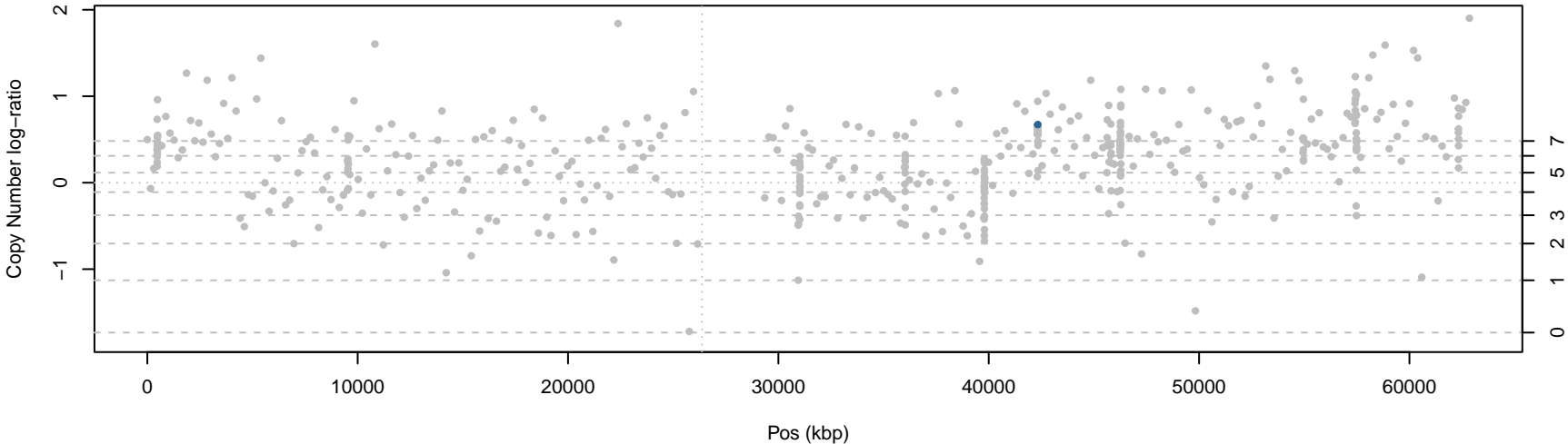
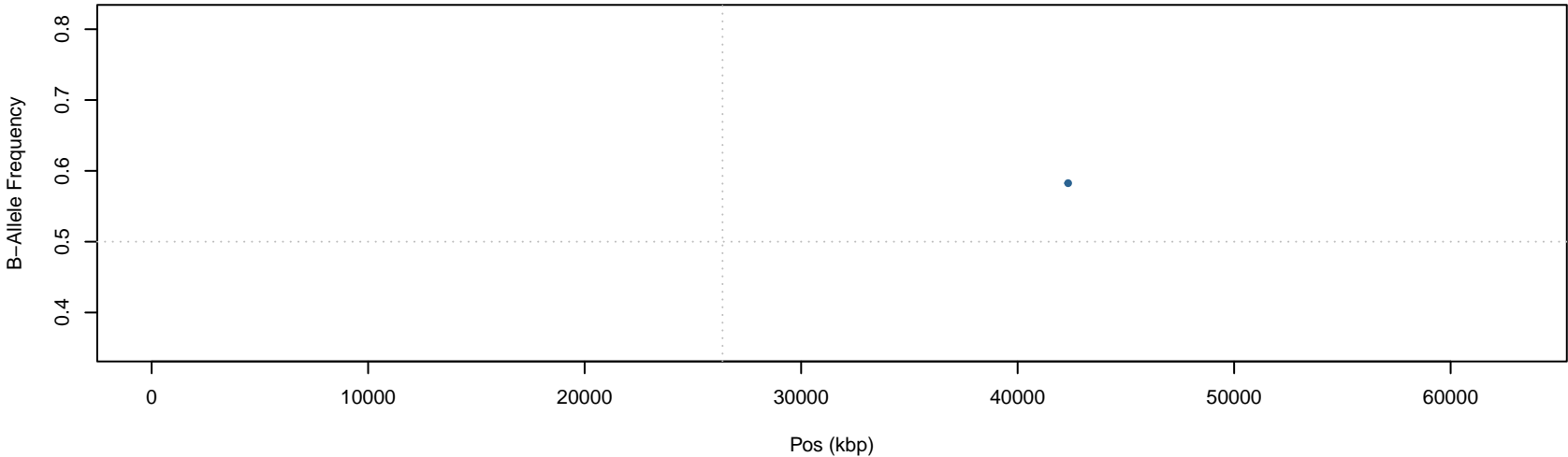


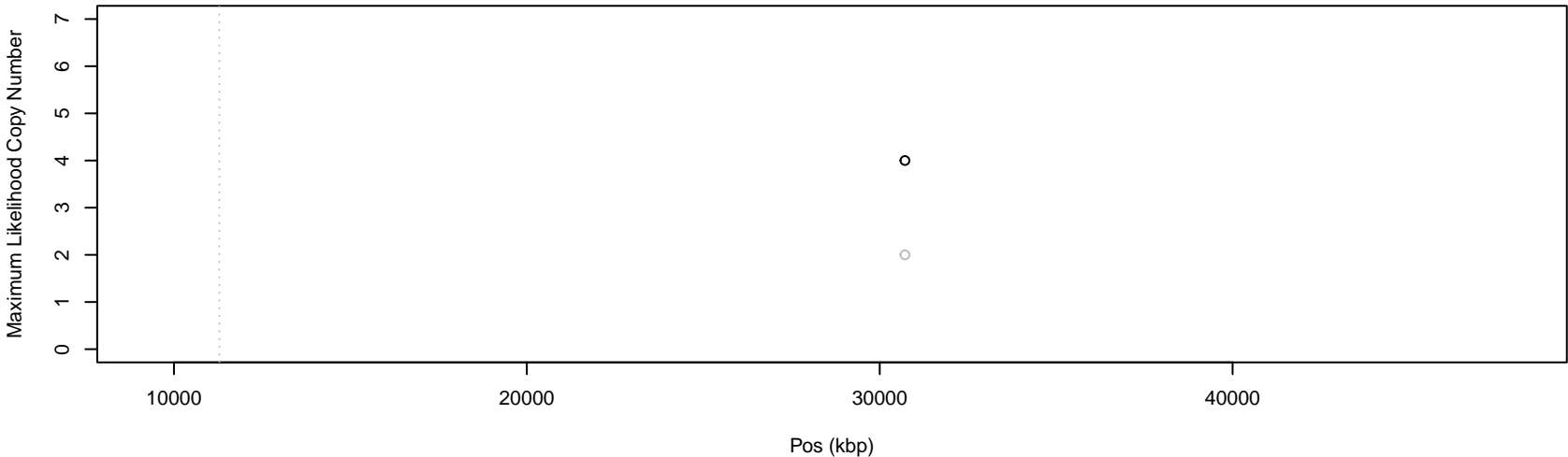
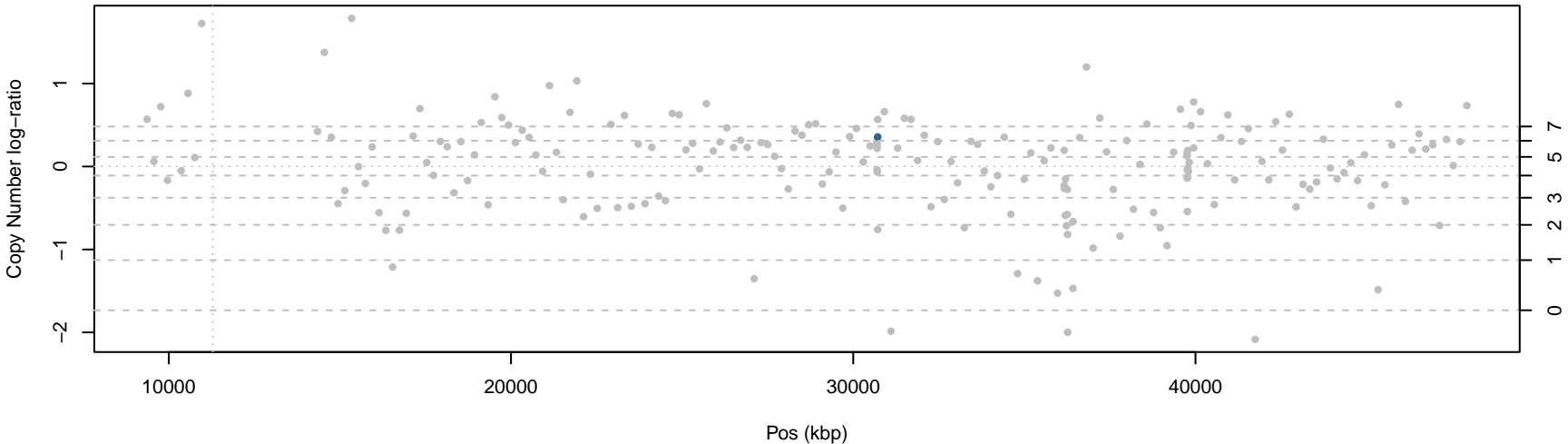
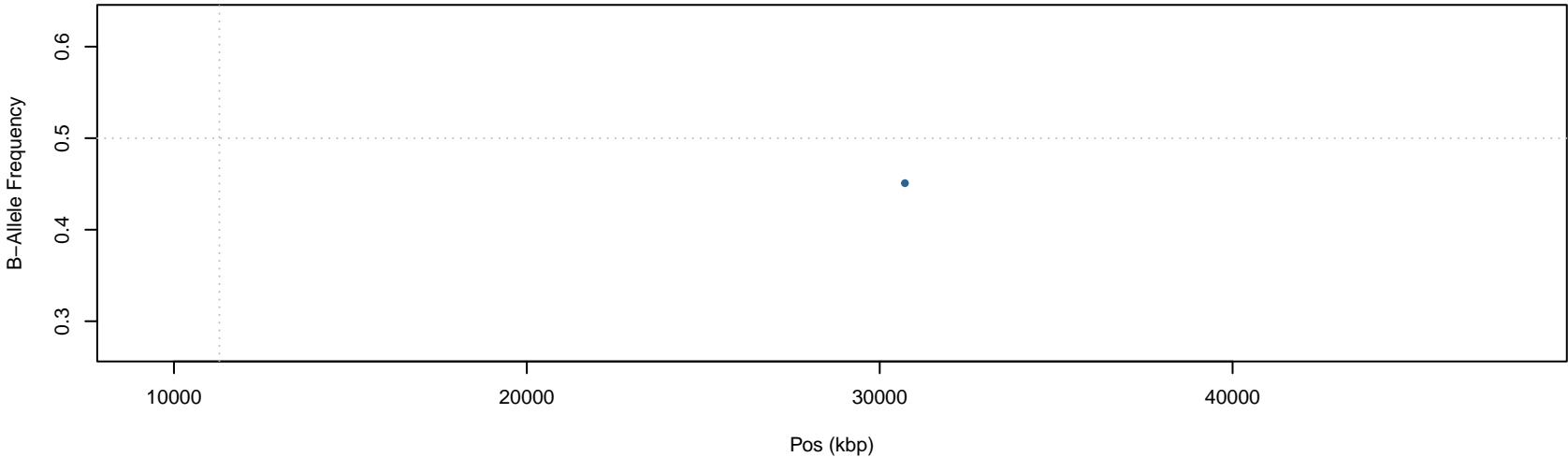
Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 19





Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 20





Purity: 0.51 Tumor ploidy: 4.466 SNV log-likelihood: -259 GoF: 84.8% Mean coverage: 465;225 Chromosome: 22

