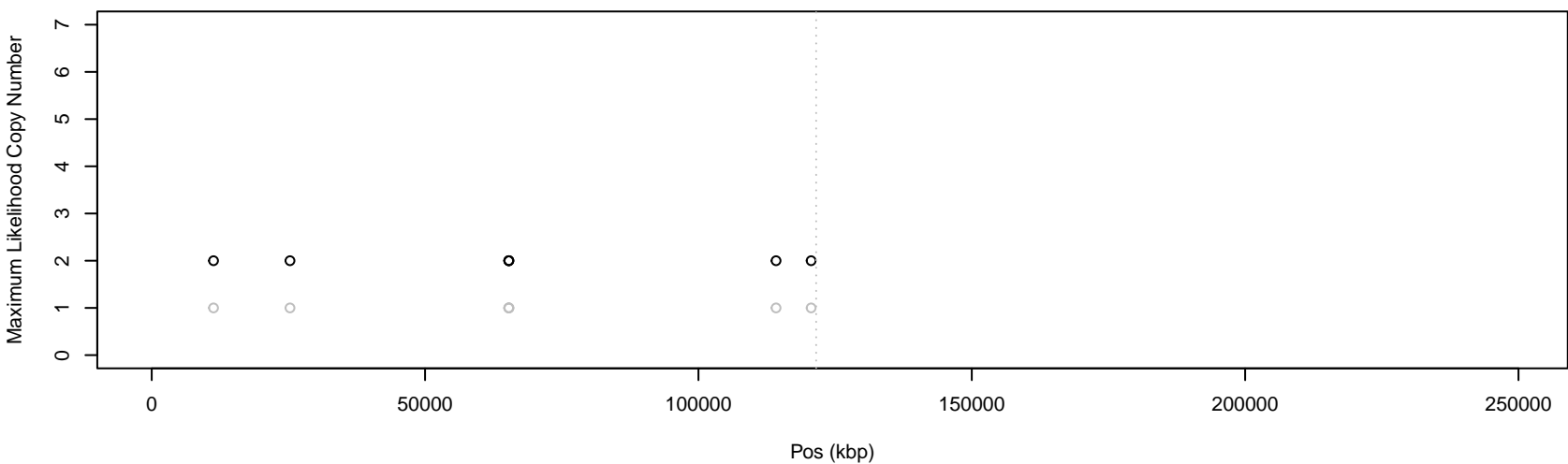
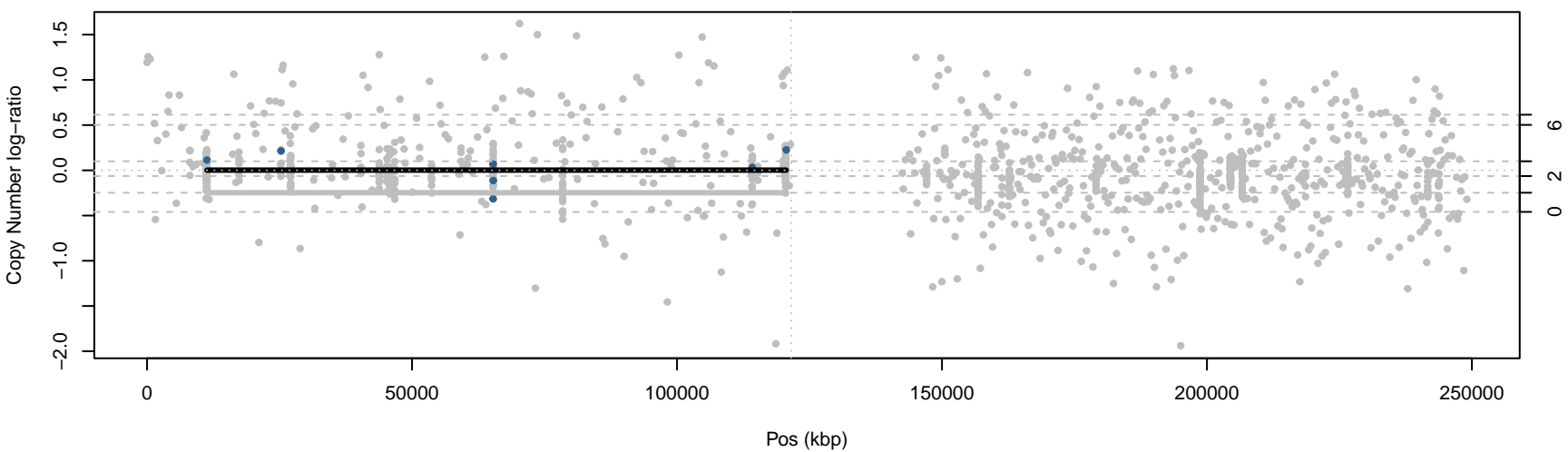
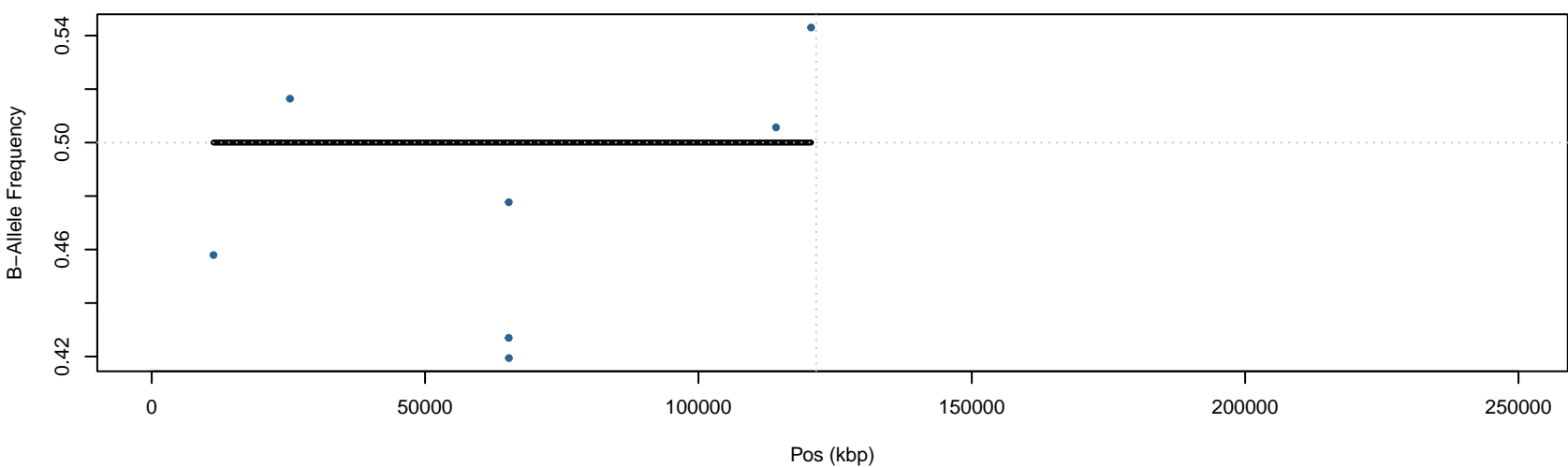
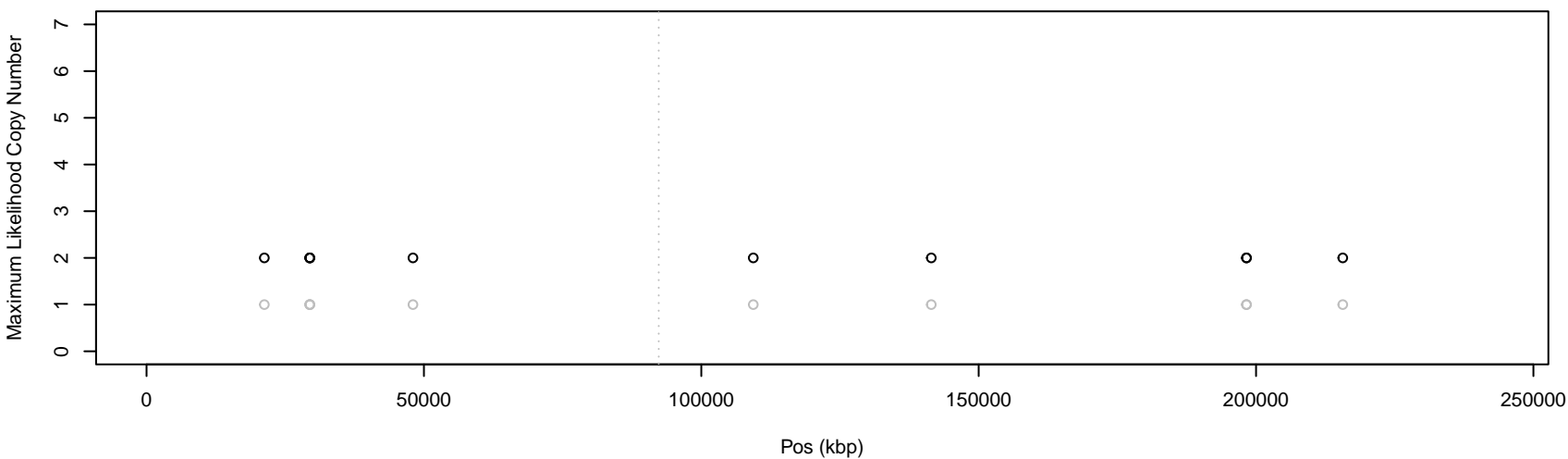
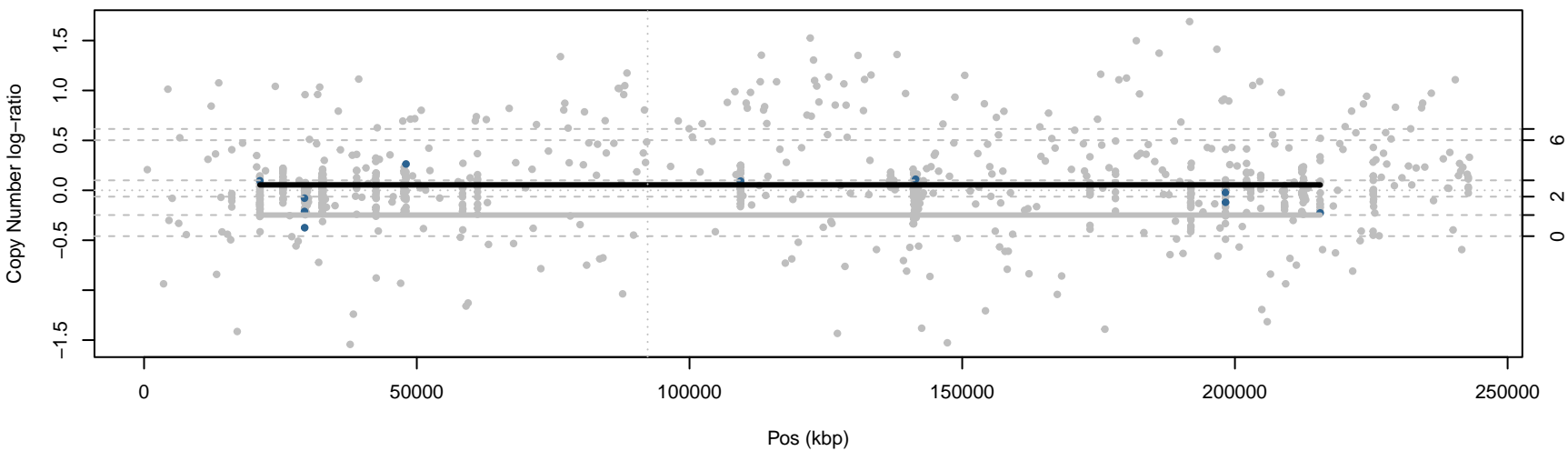
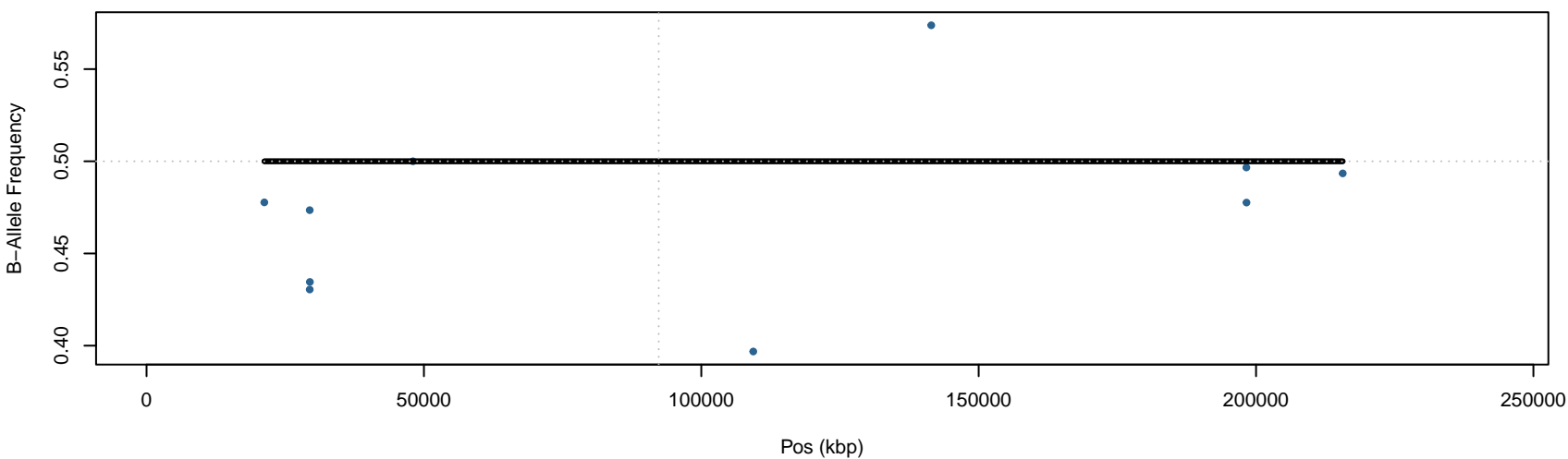


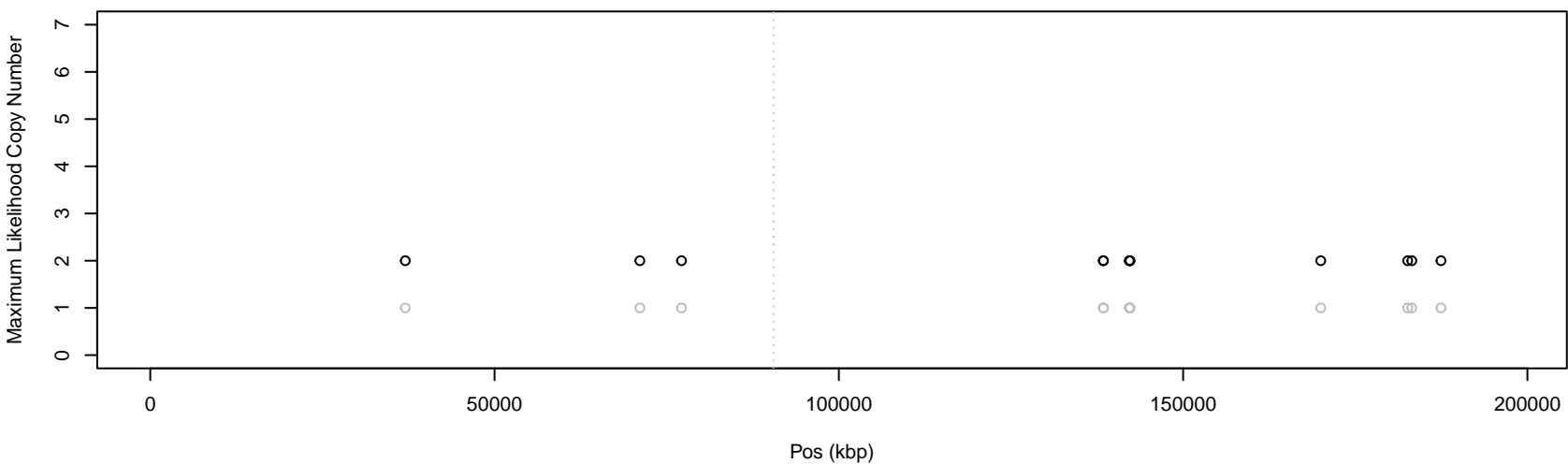
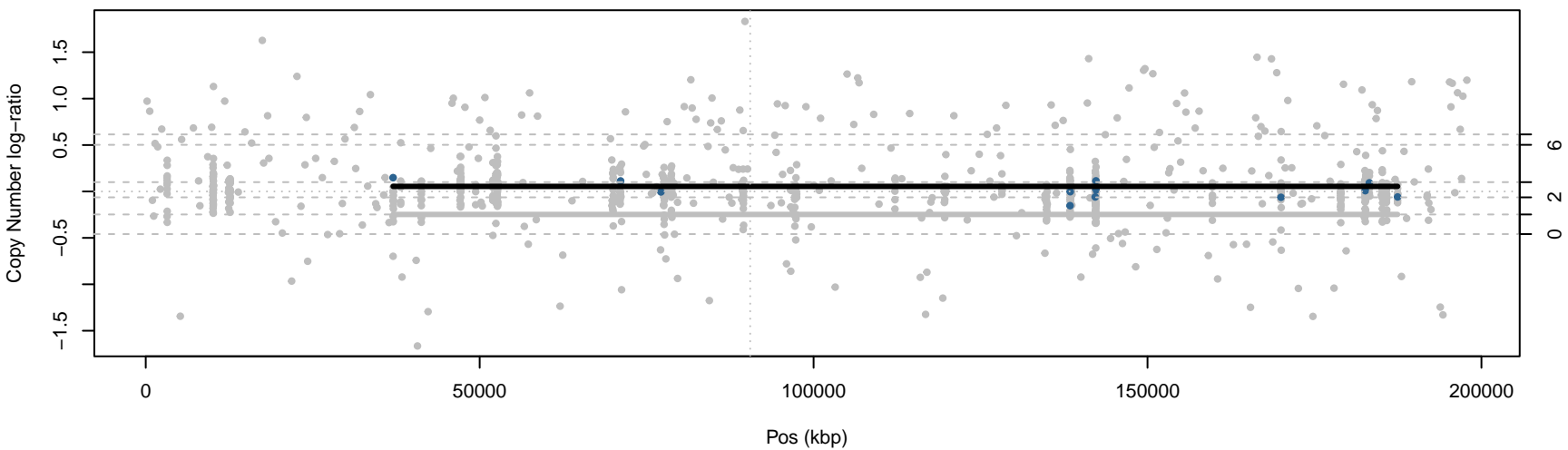
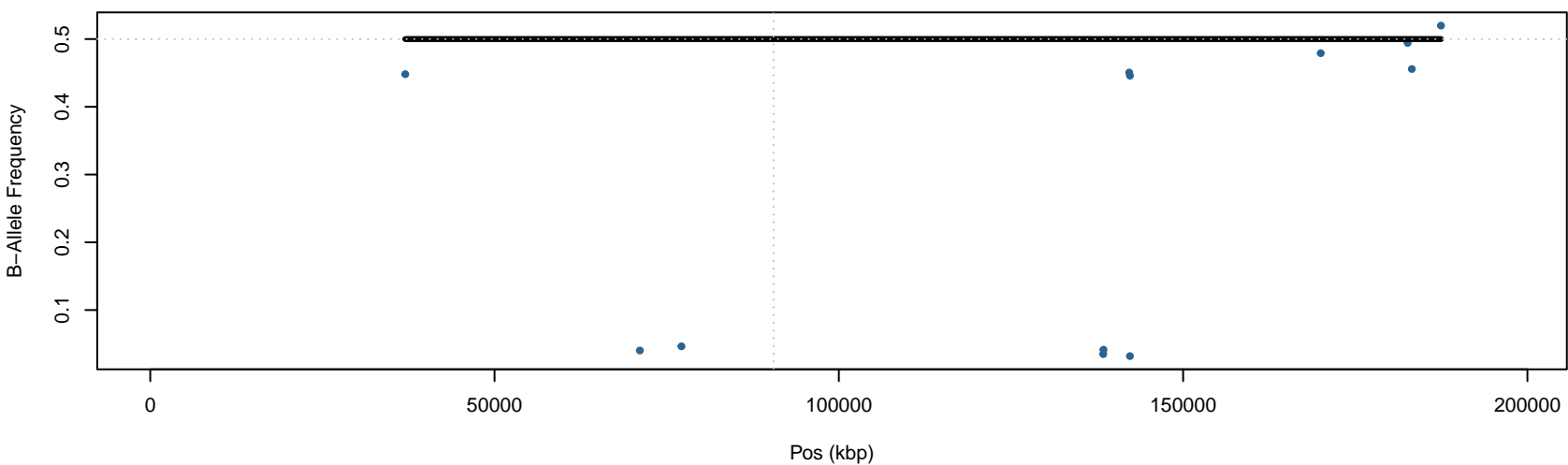
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 1



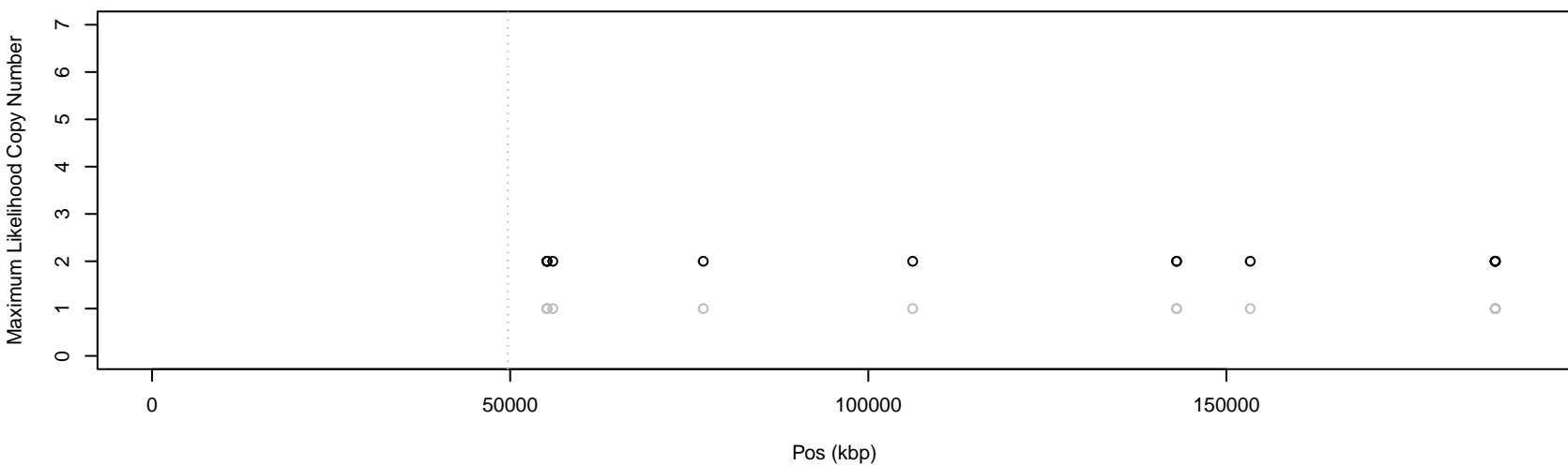
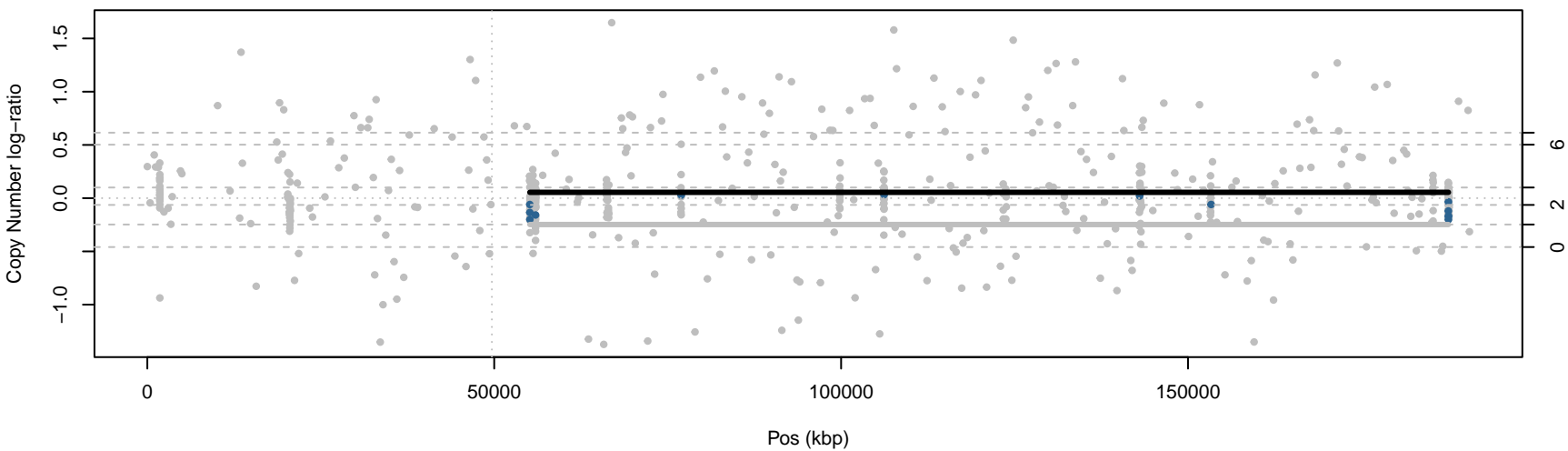
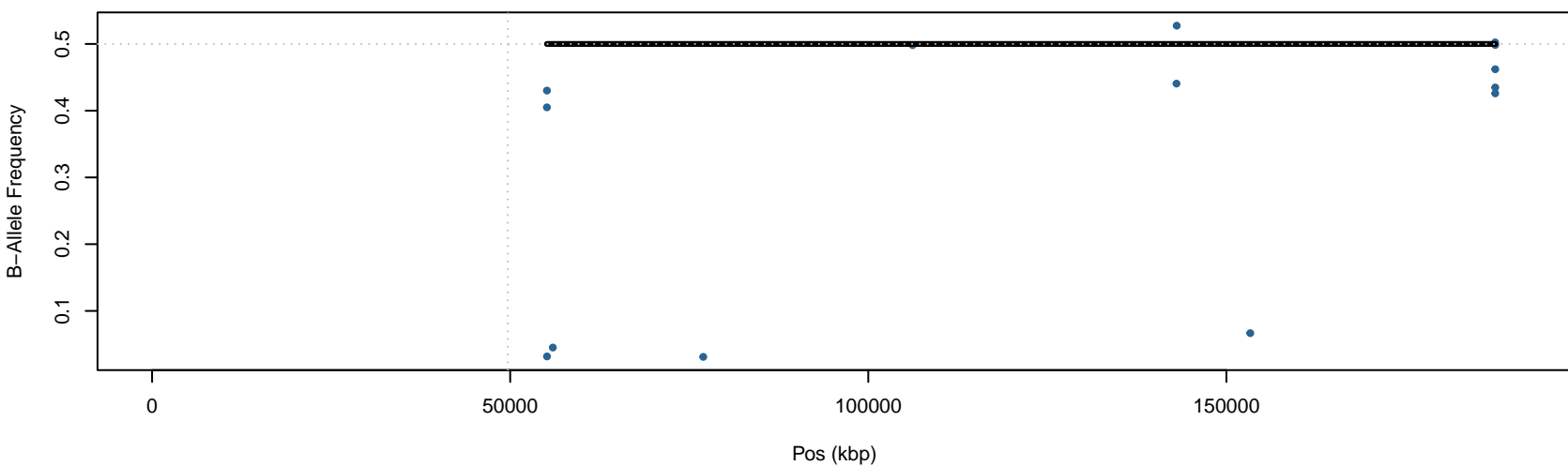
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 2



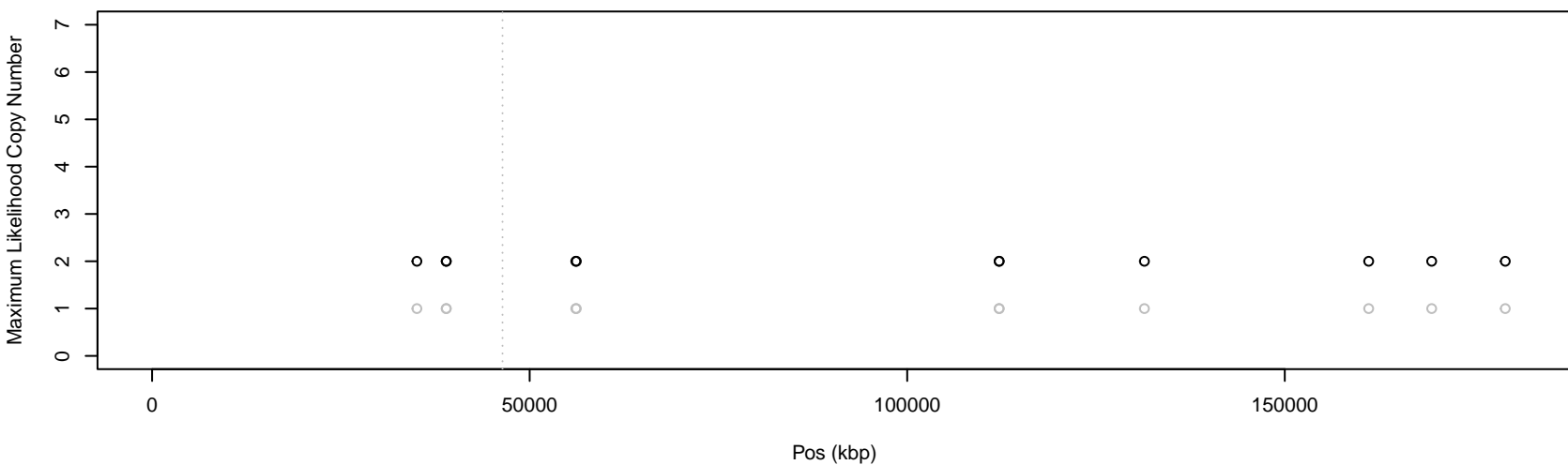
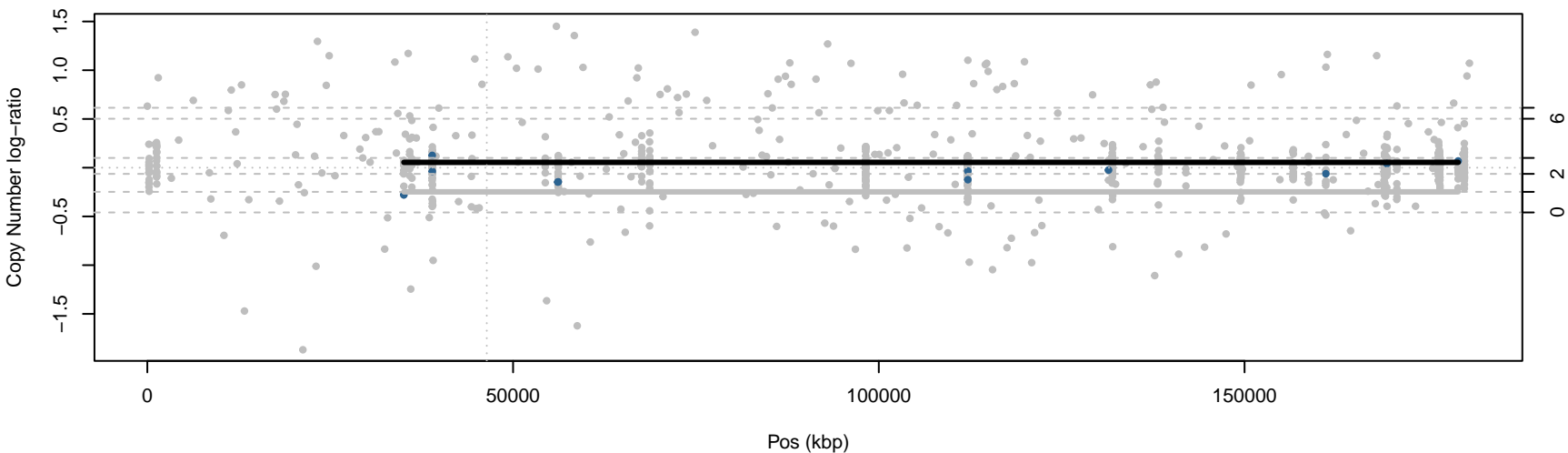
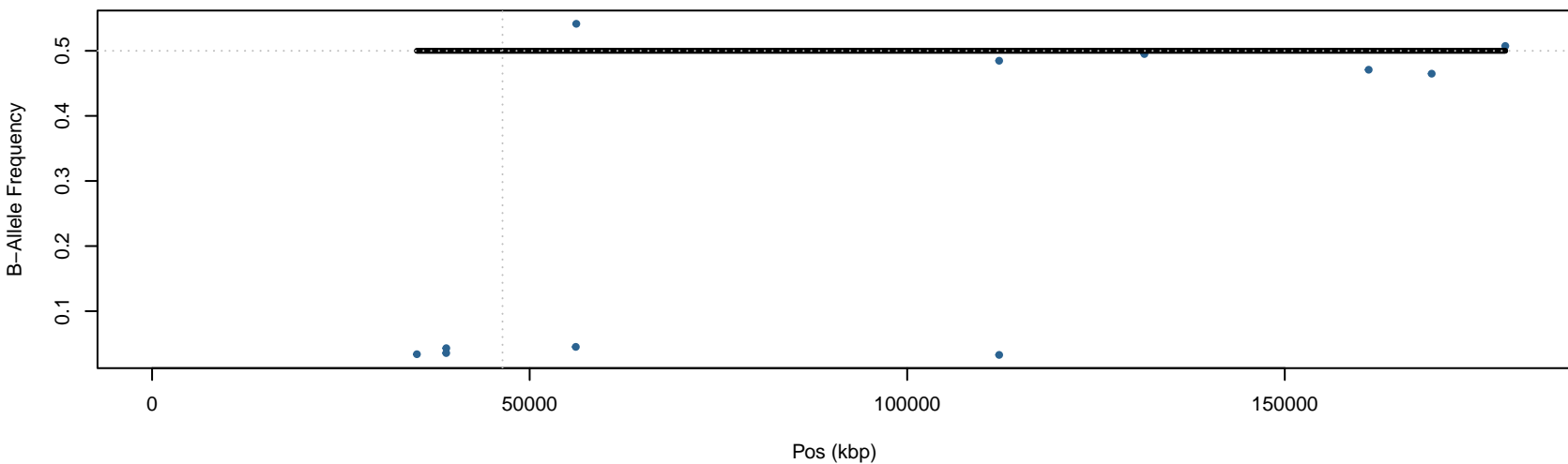
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 3



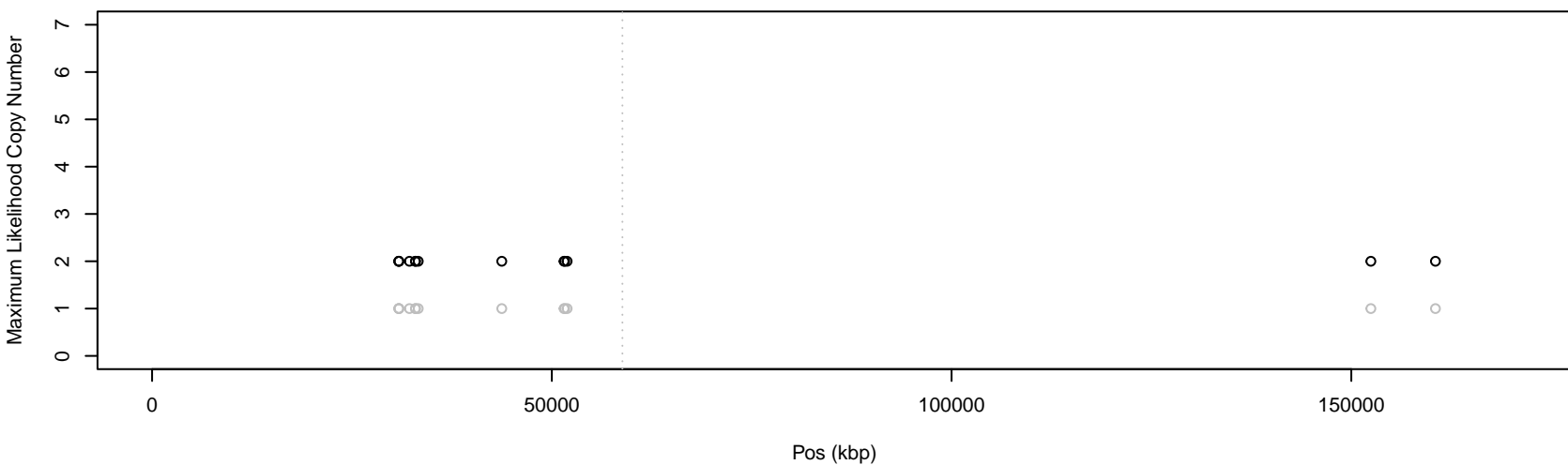
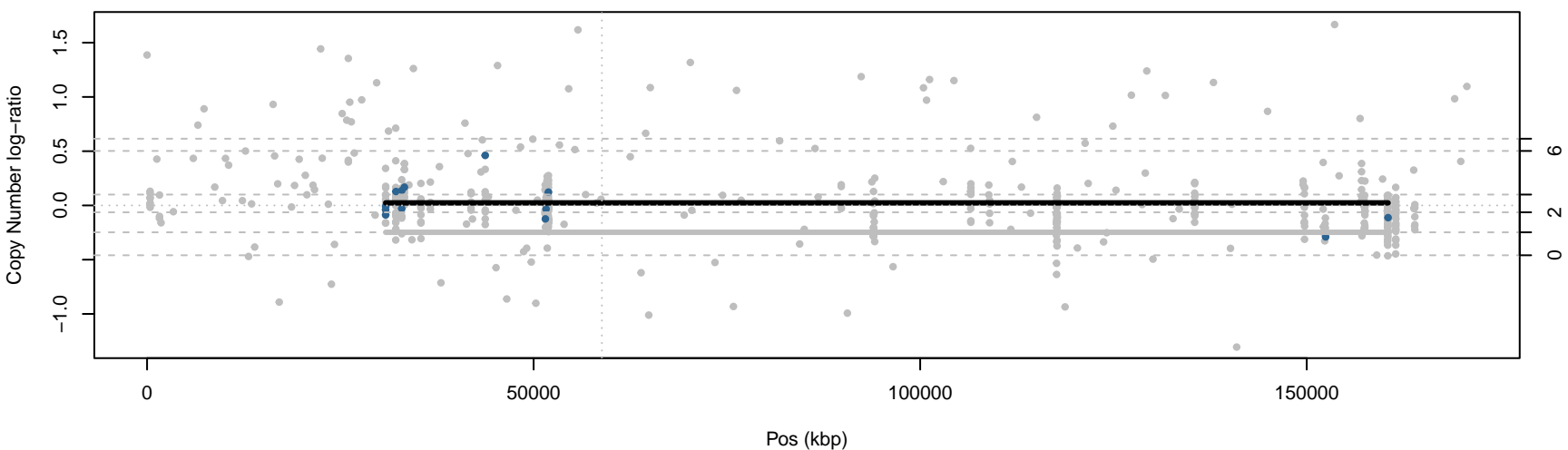
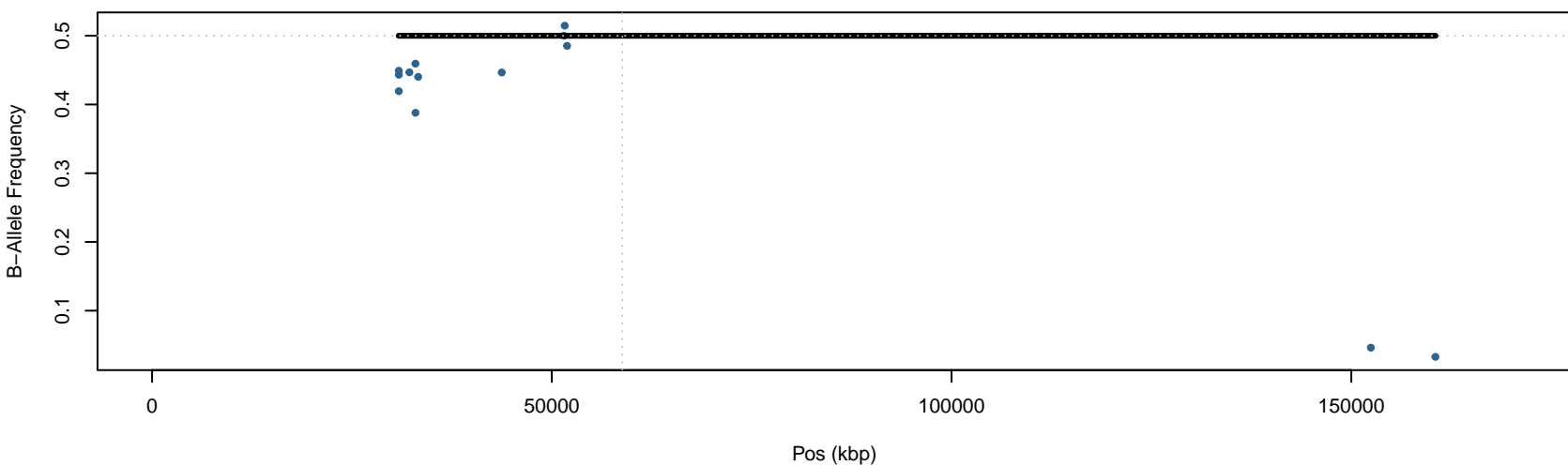
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 4



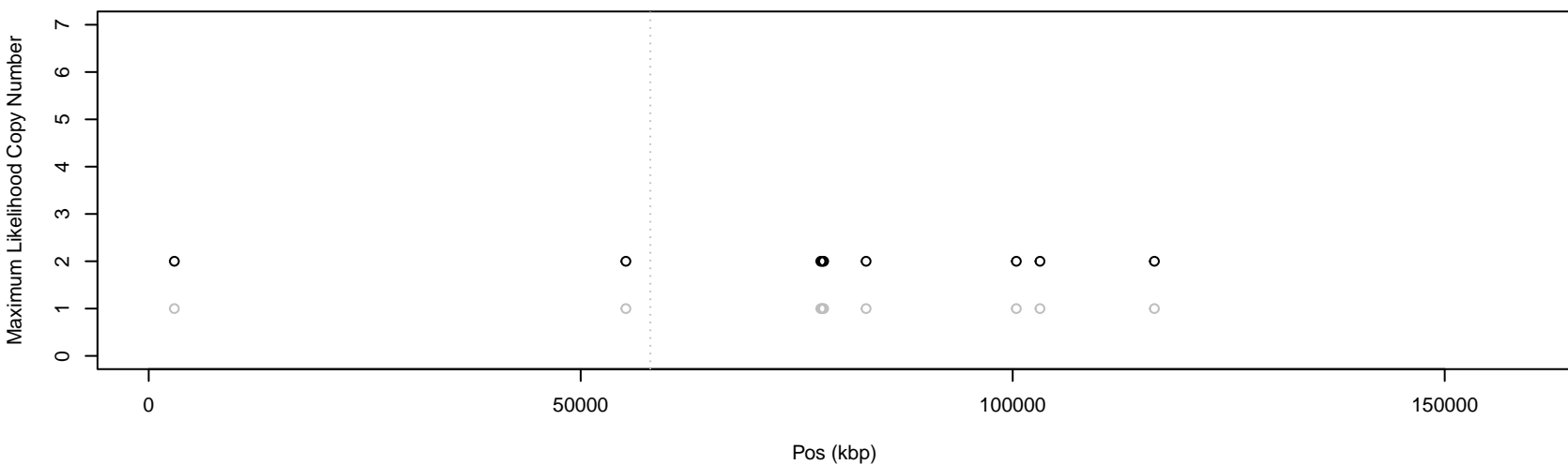
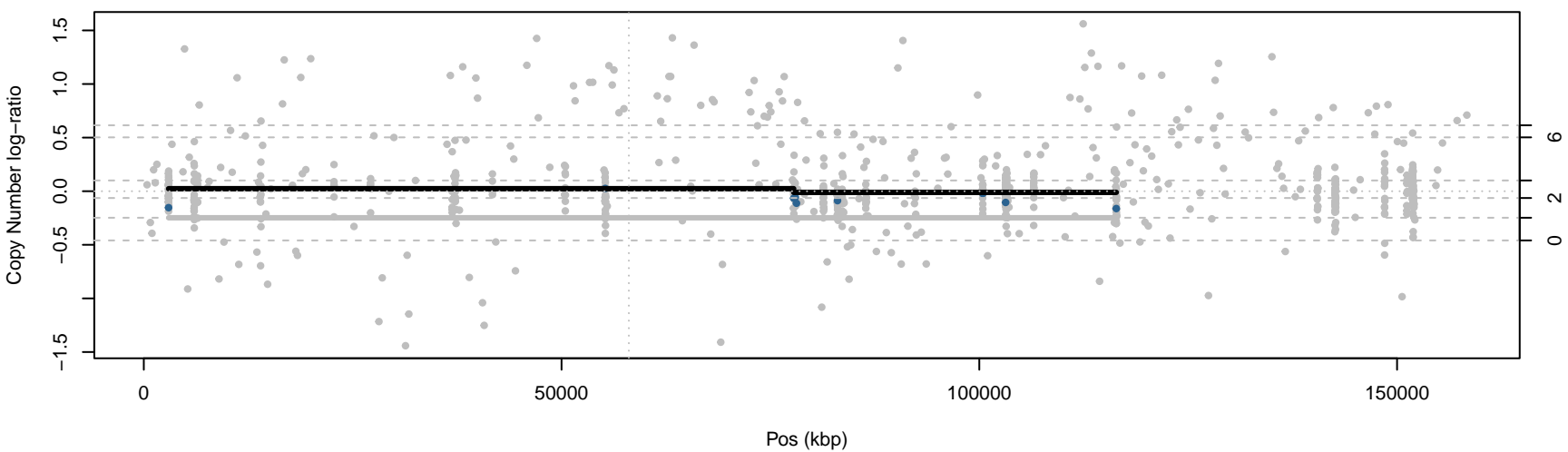
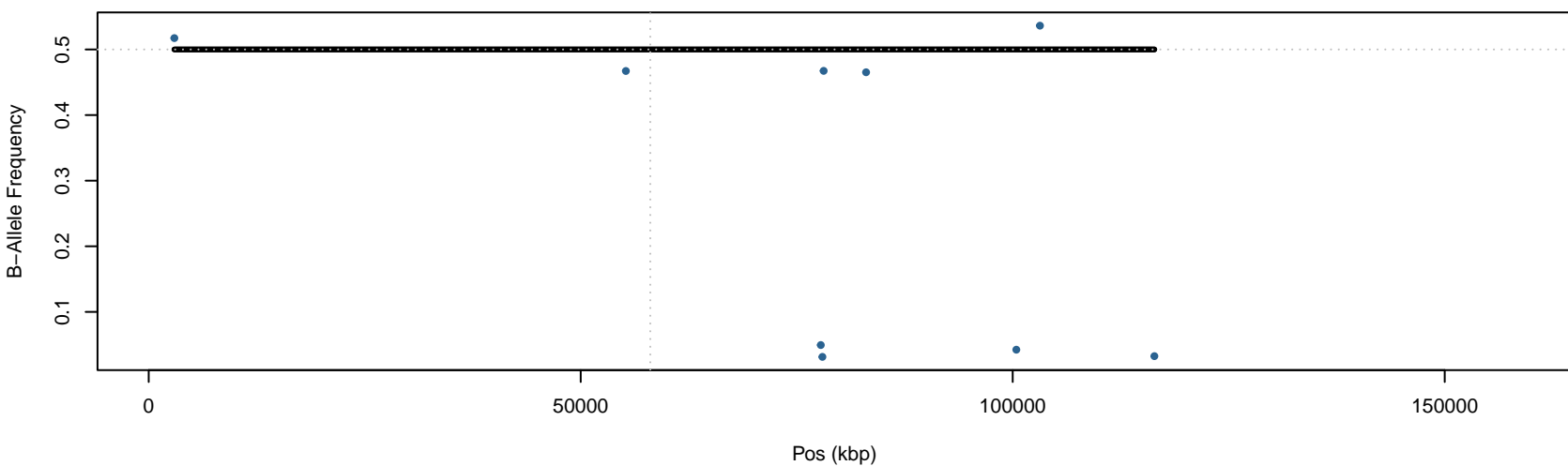
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 5



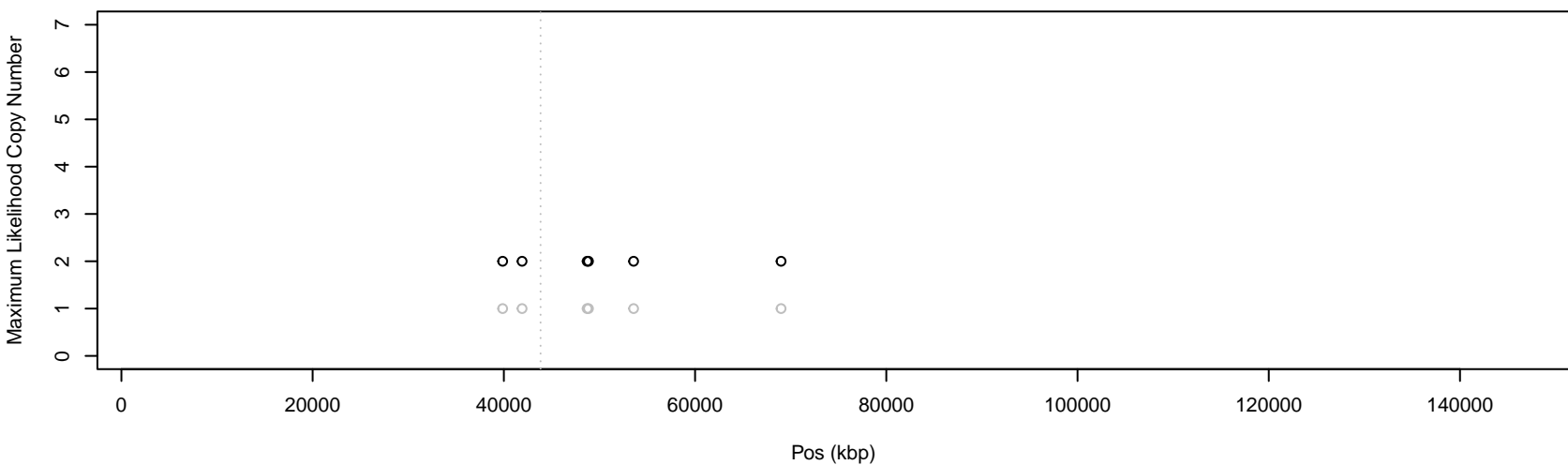
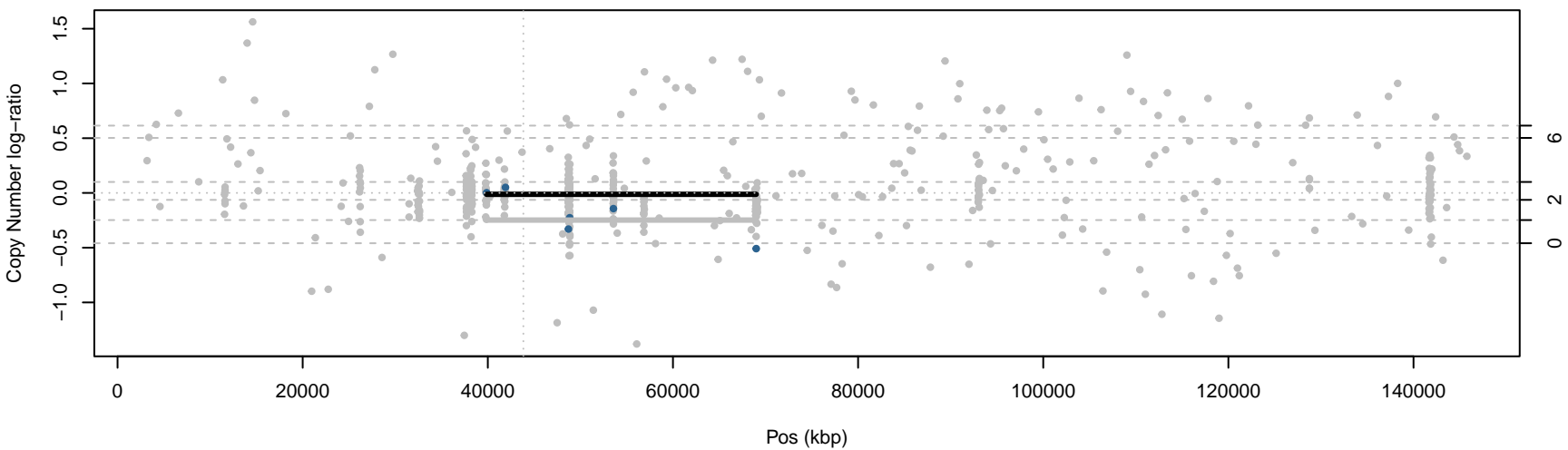
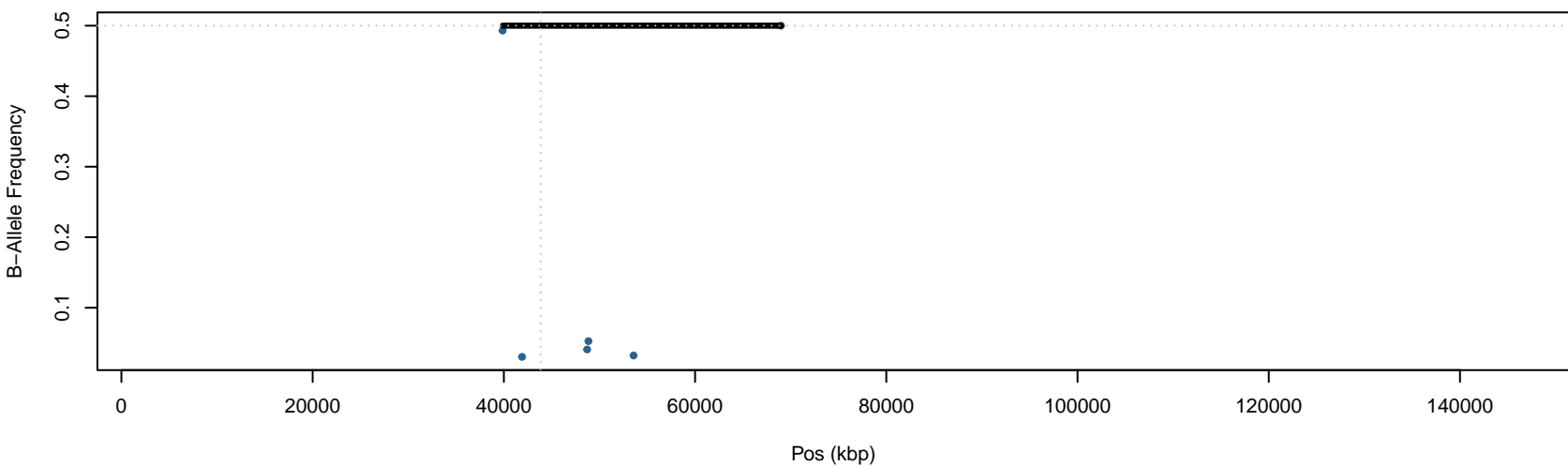
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 6



Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 7

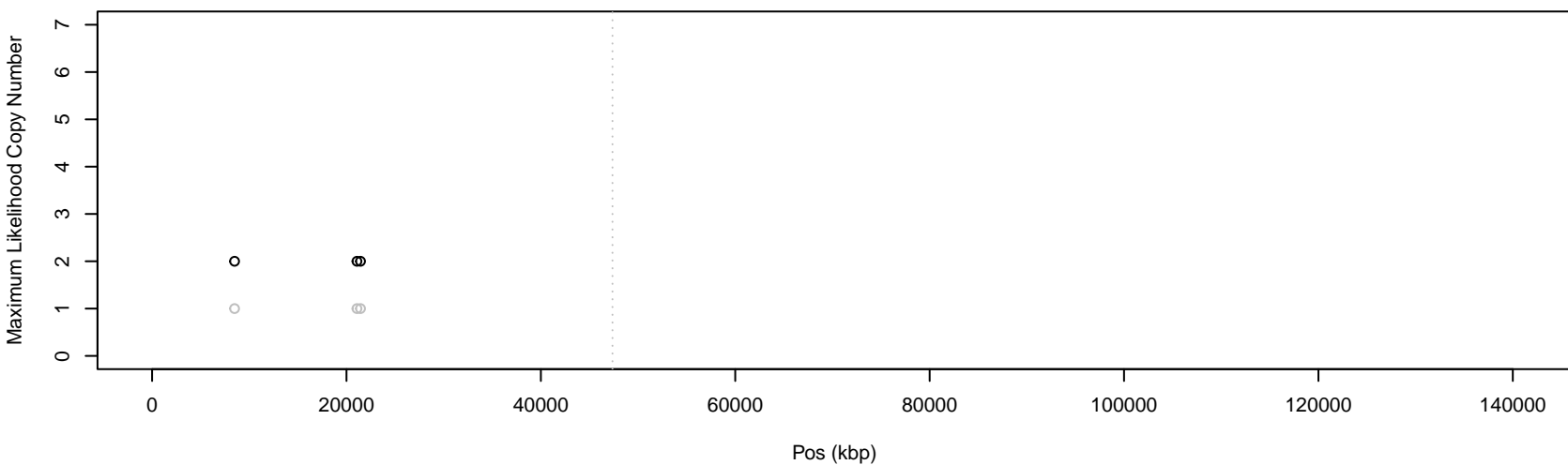
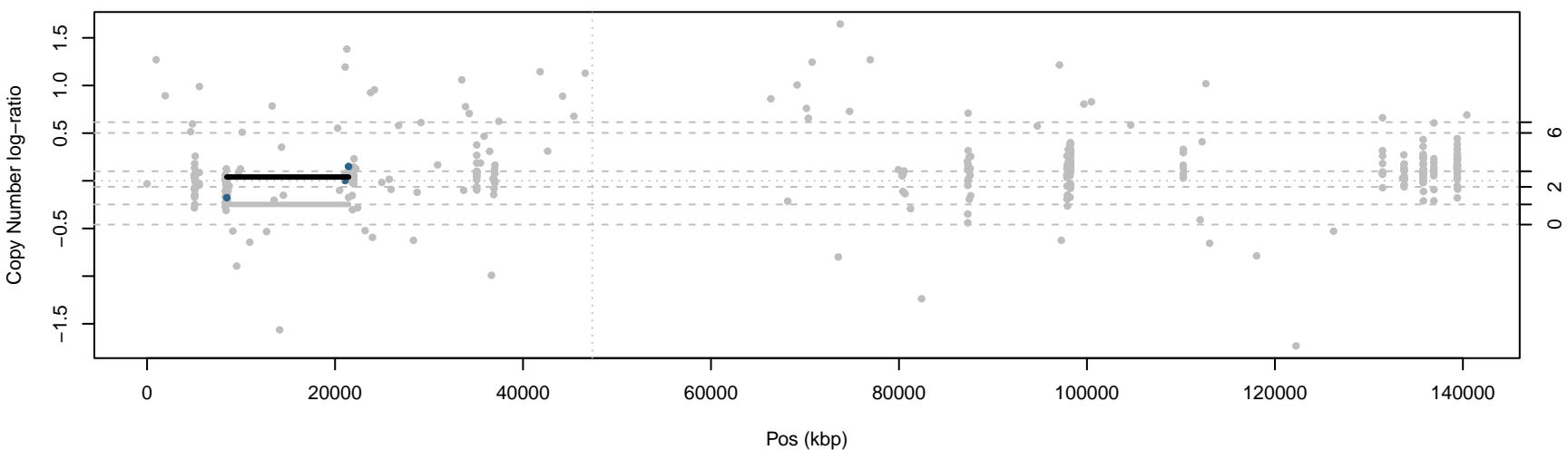
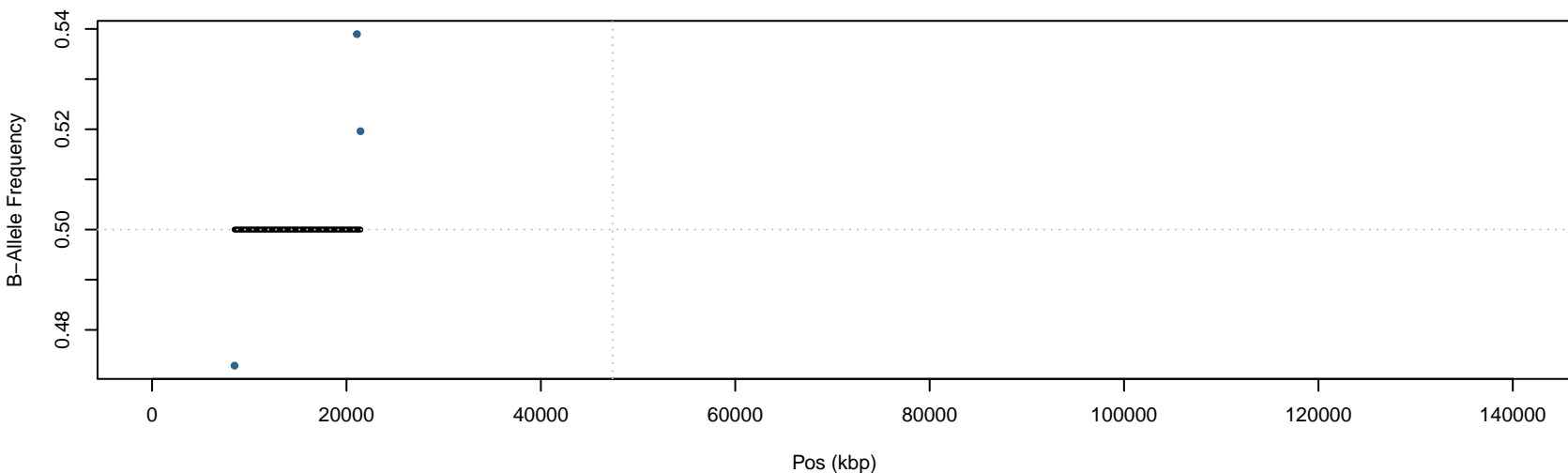


Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 8

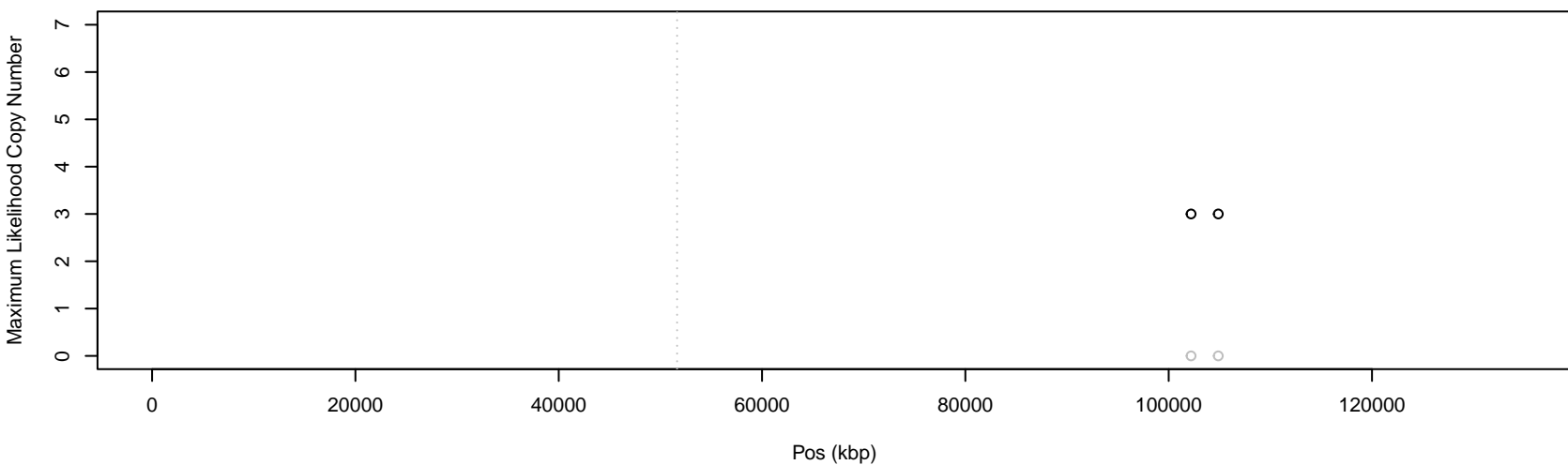
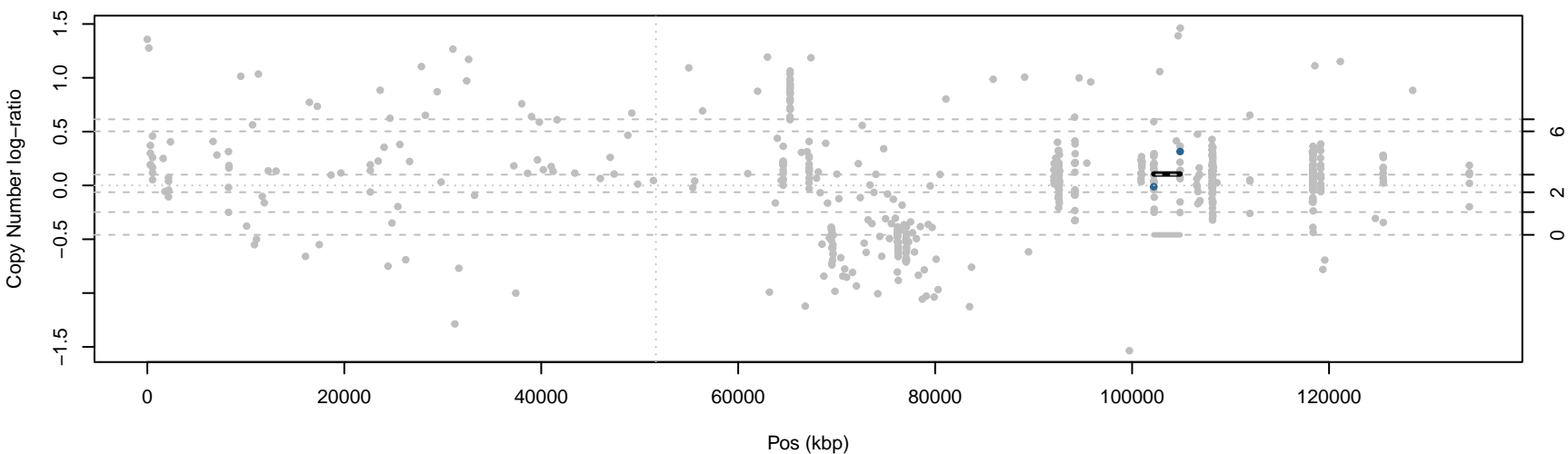
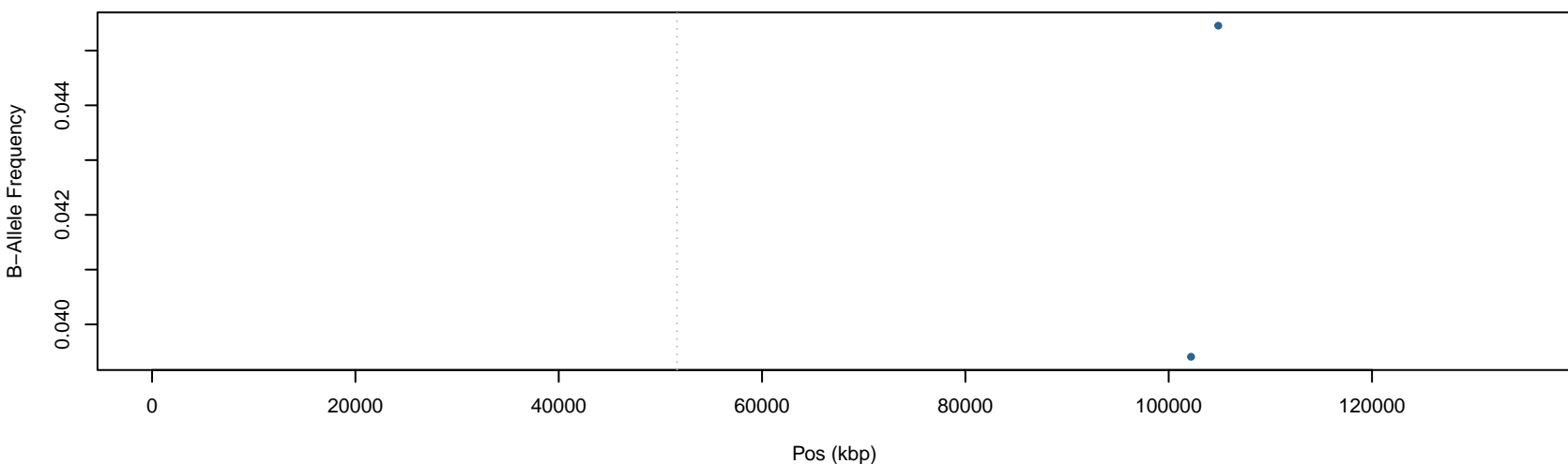




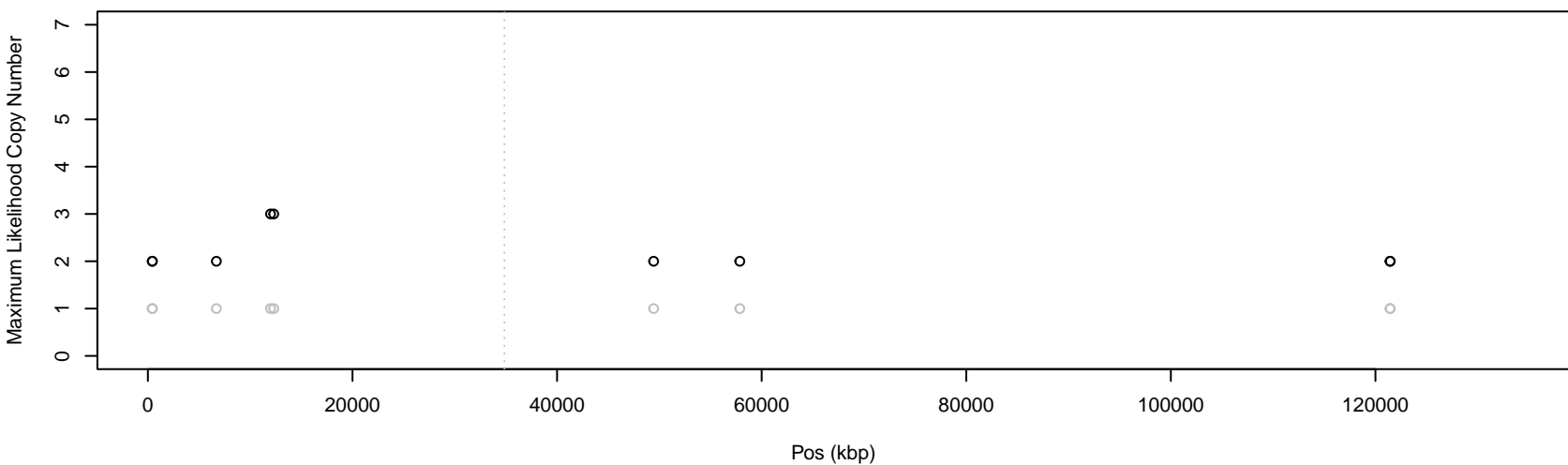
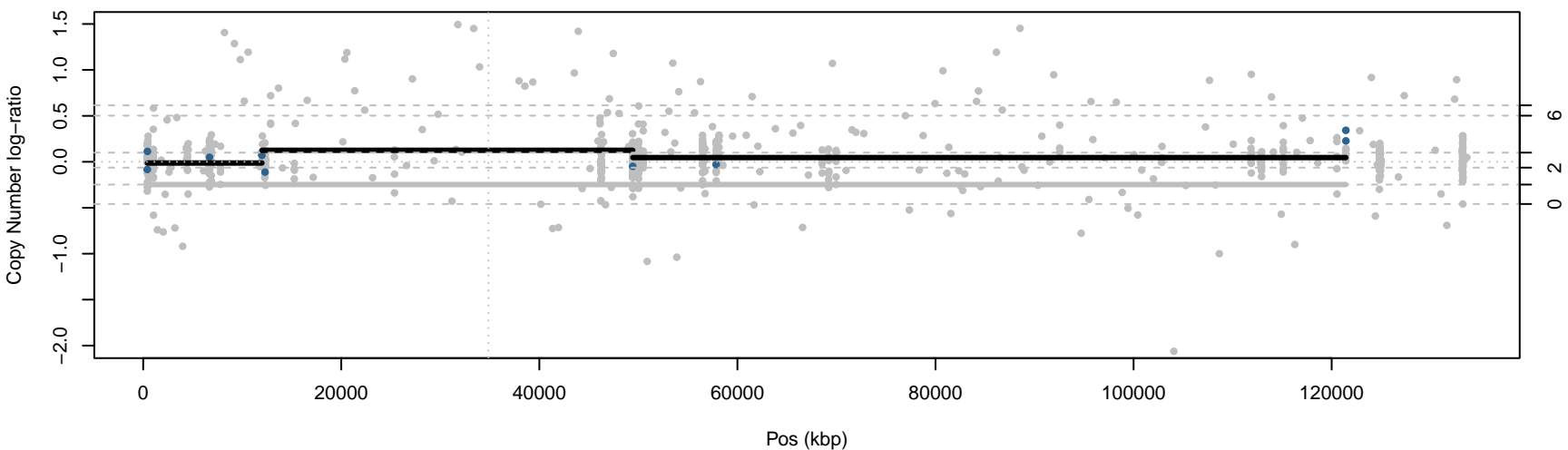
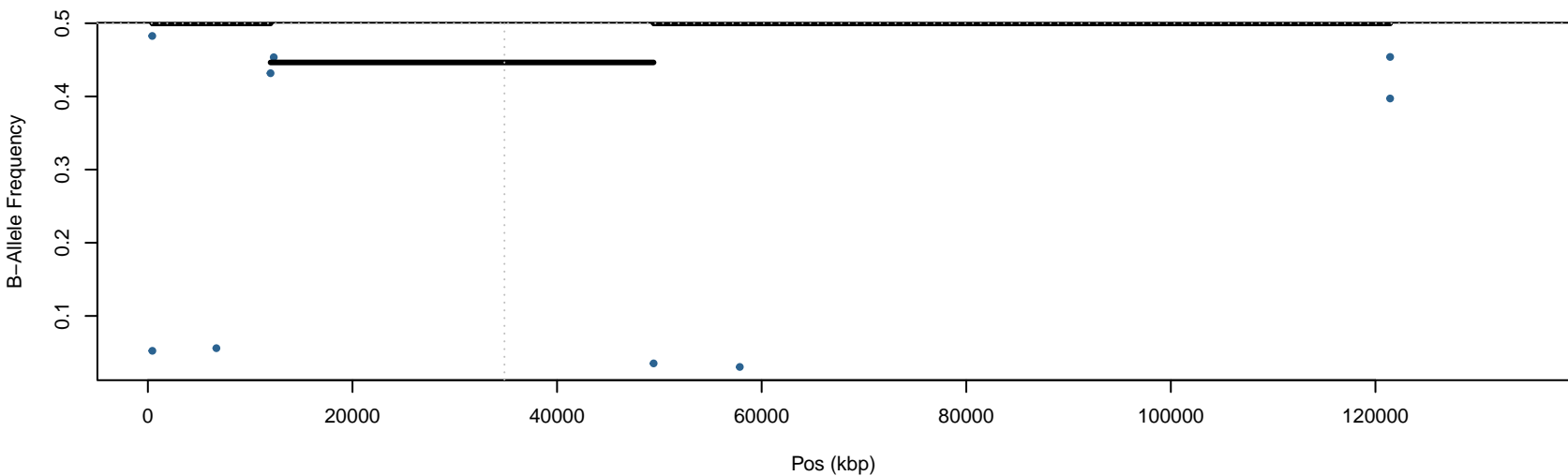
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 9



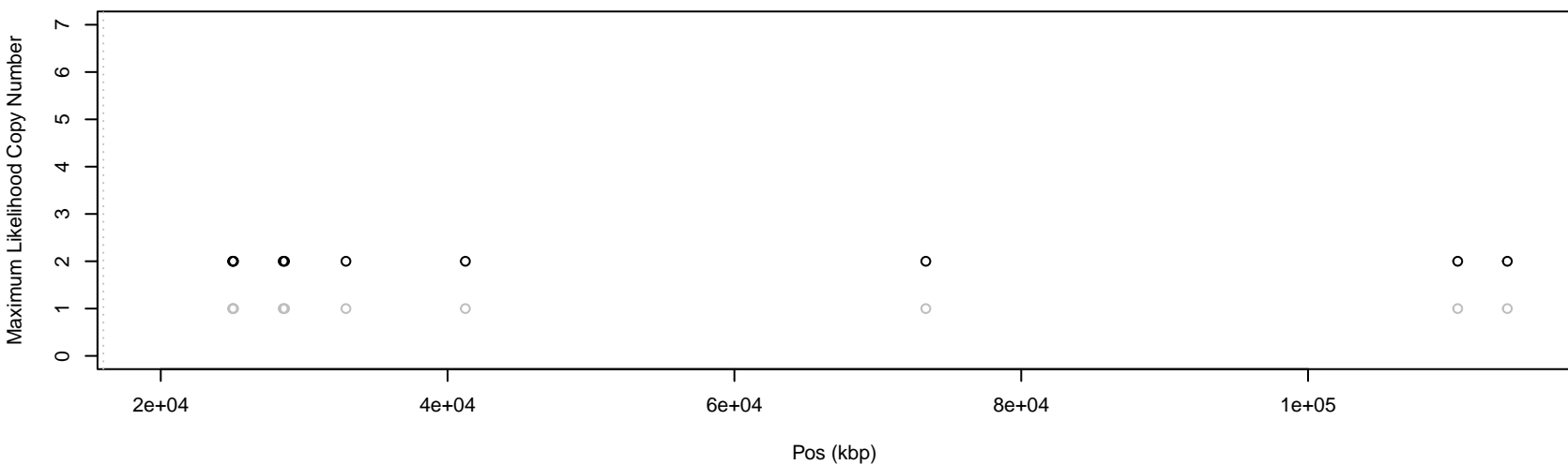
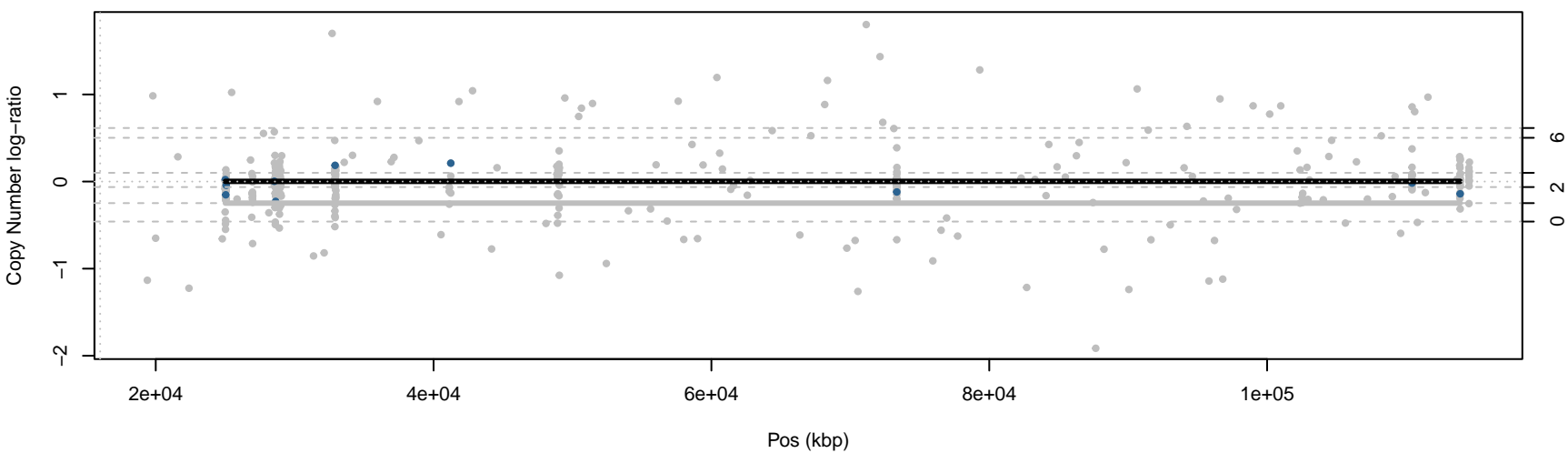
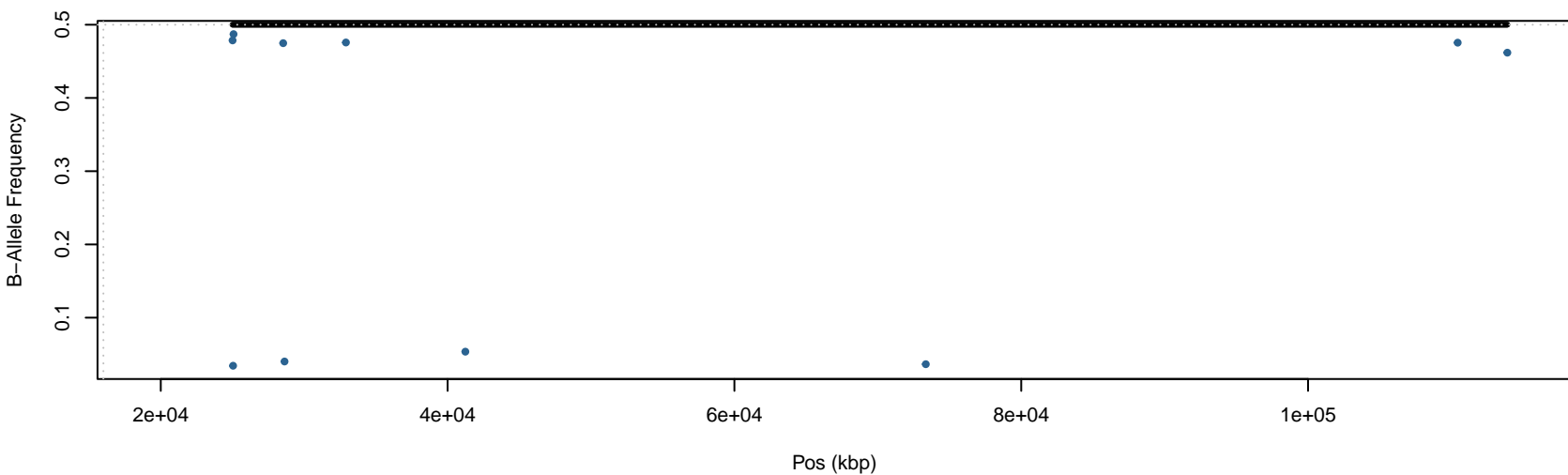
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 11



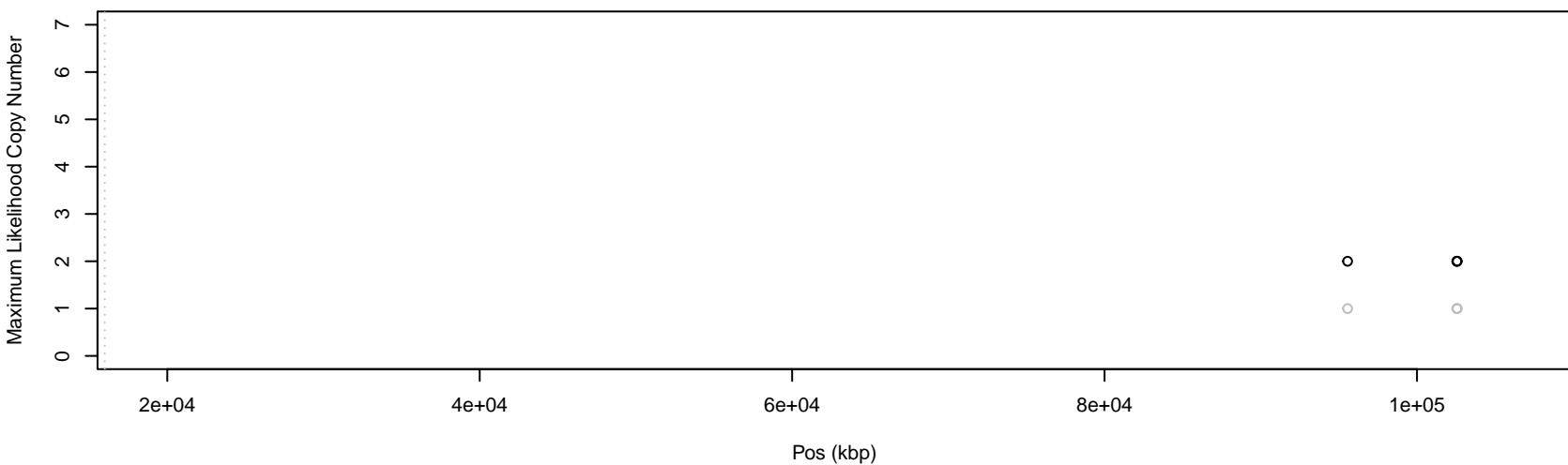
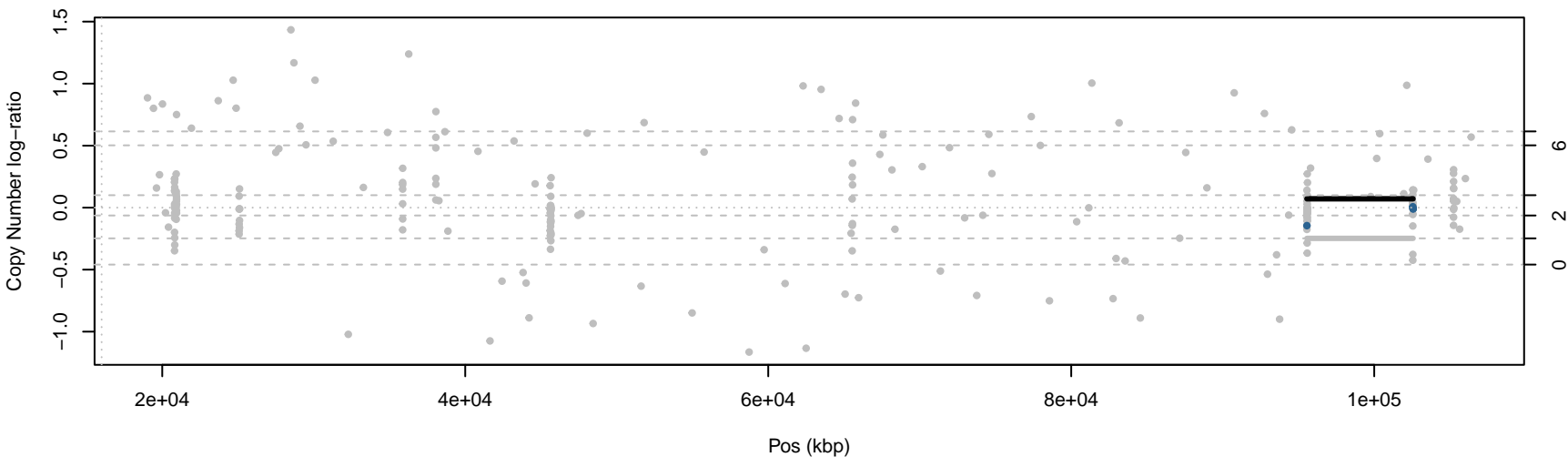
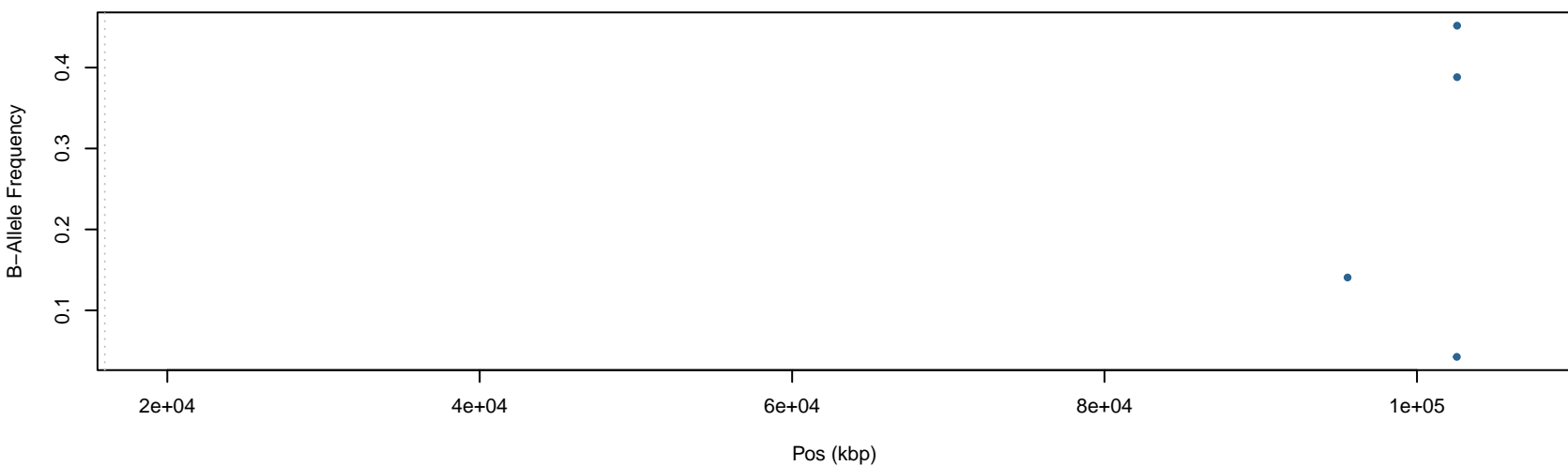
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 12



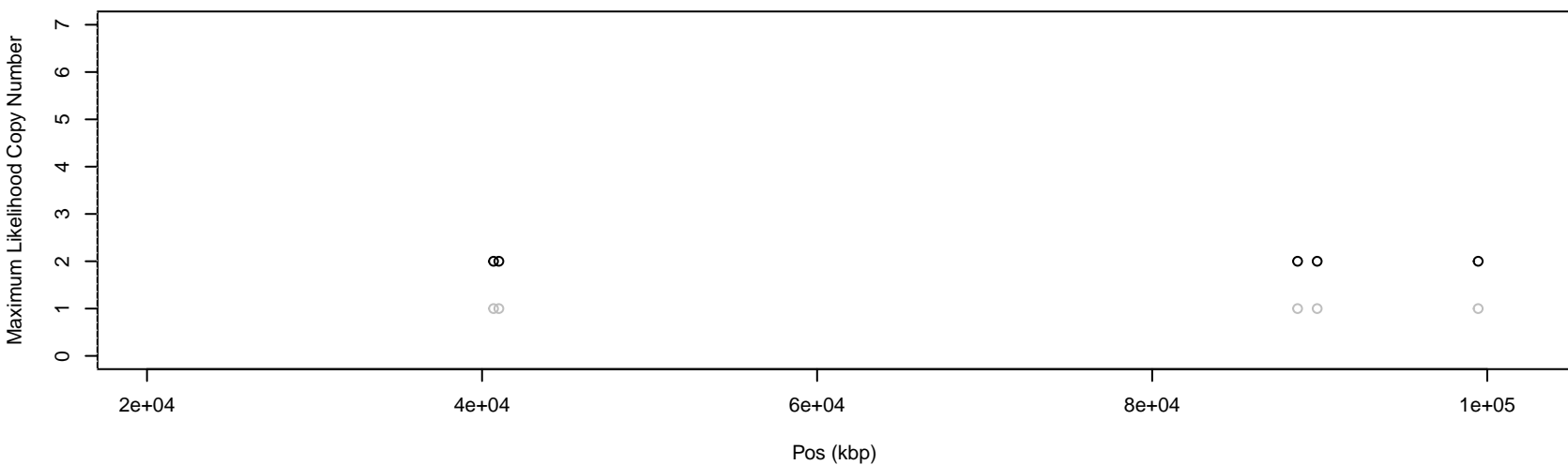
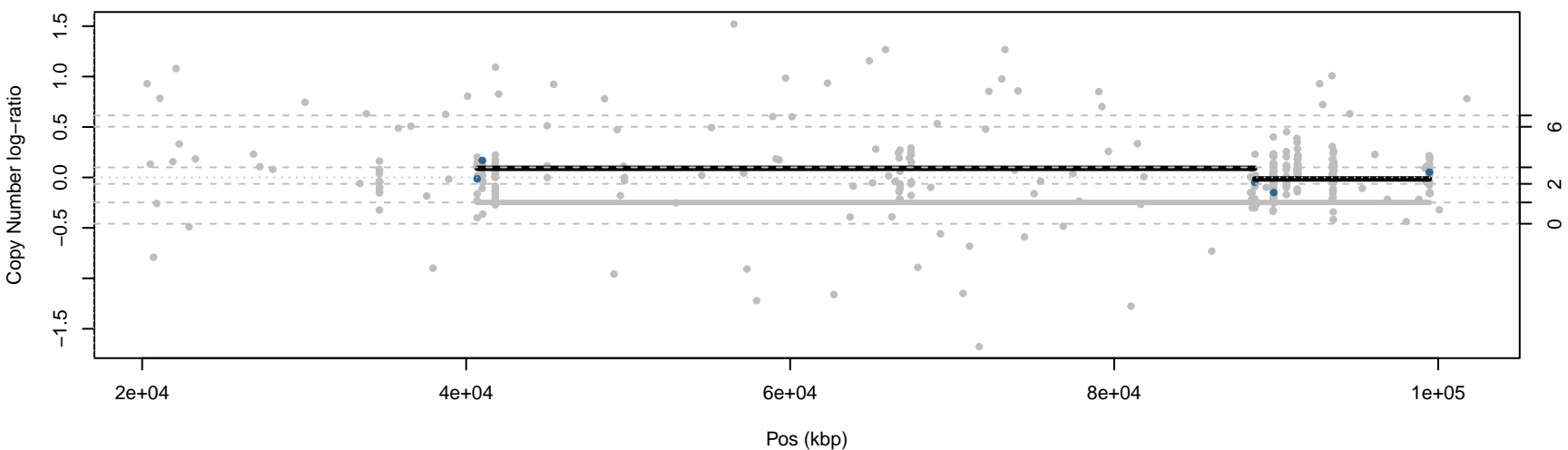
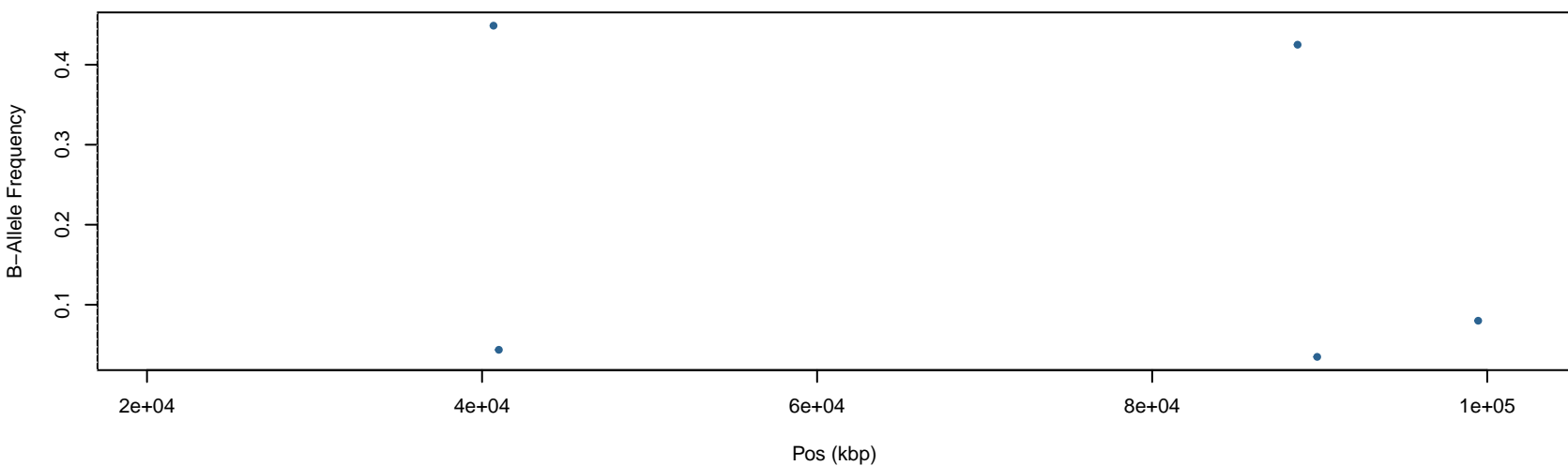
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 13



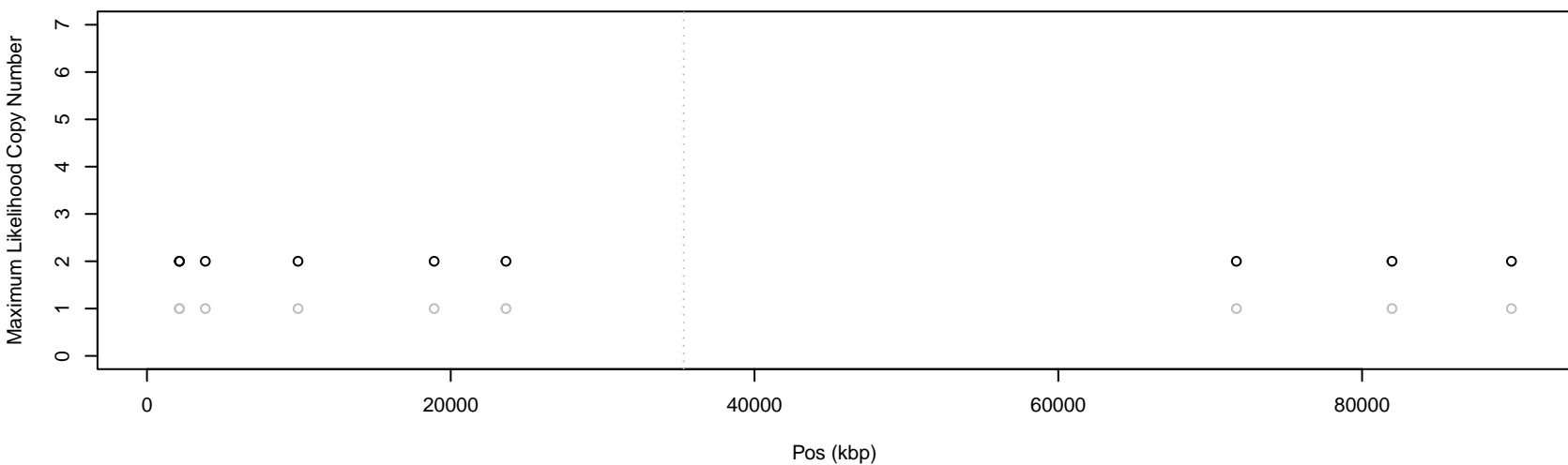
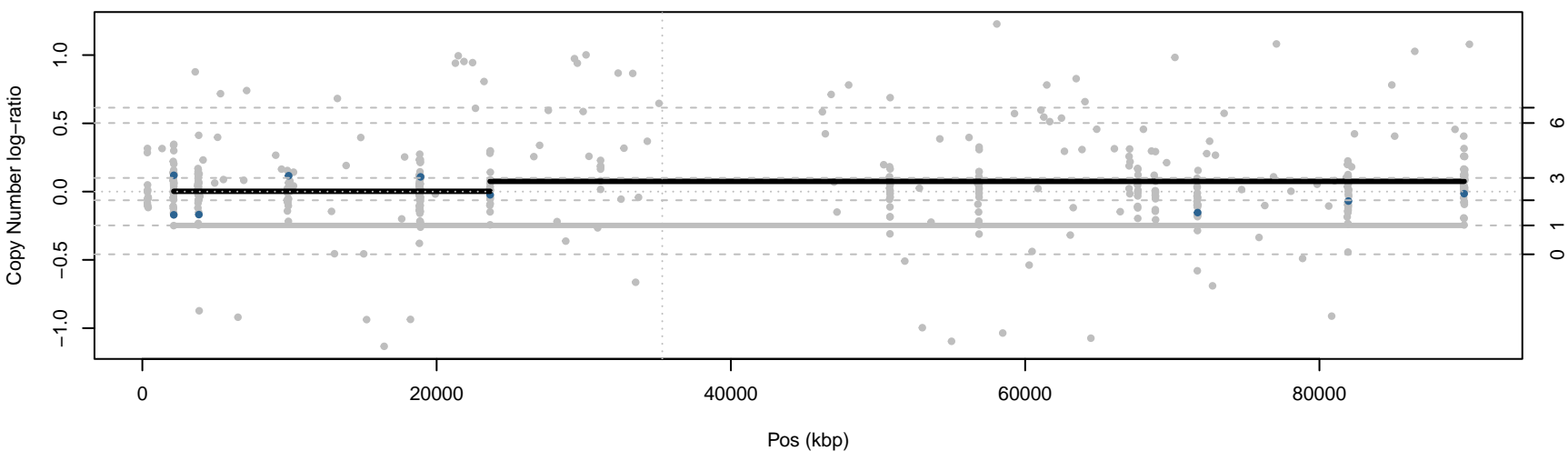
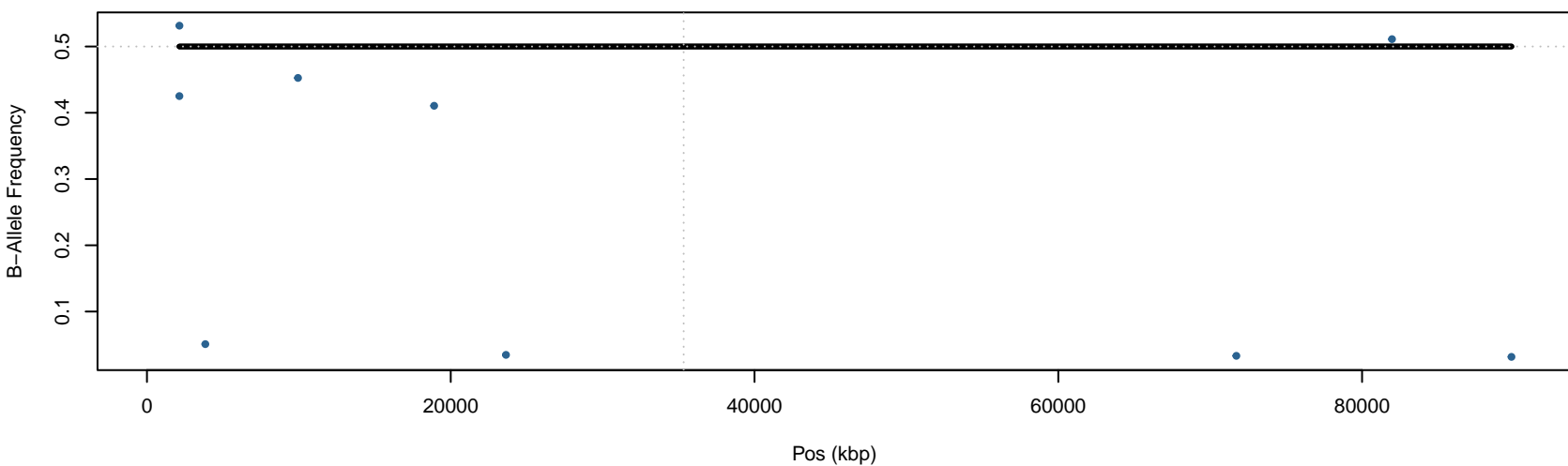
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 14



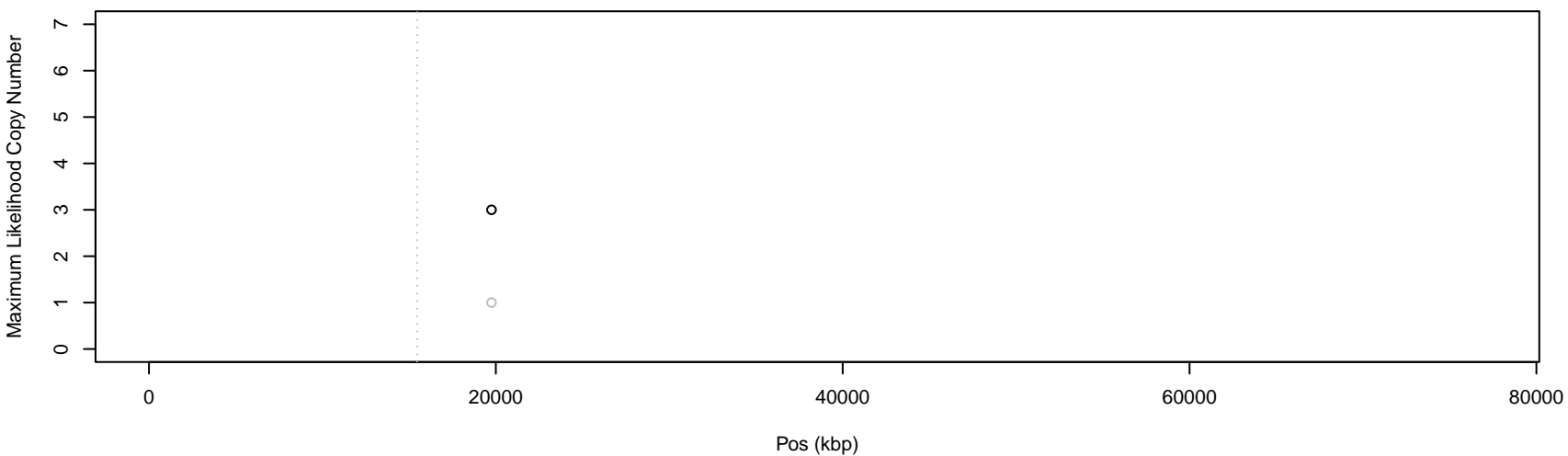
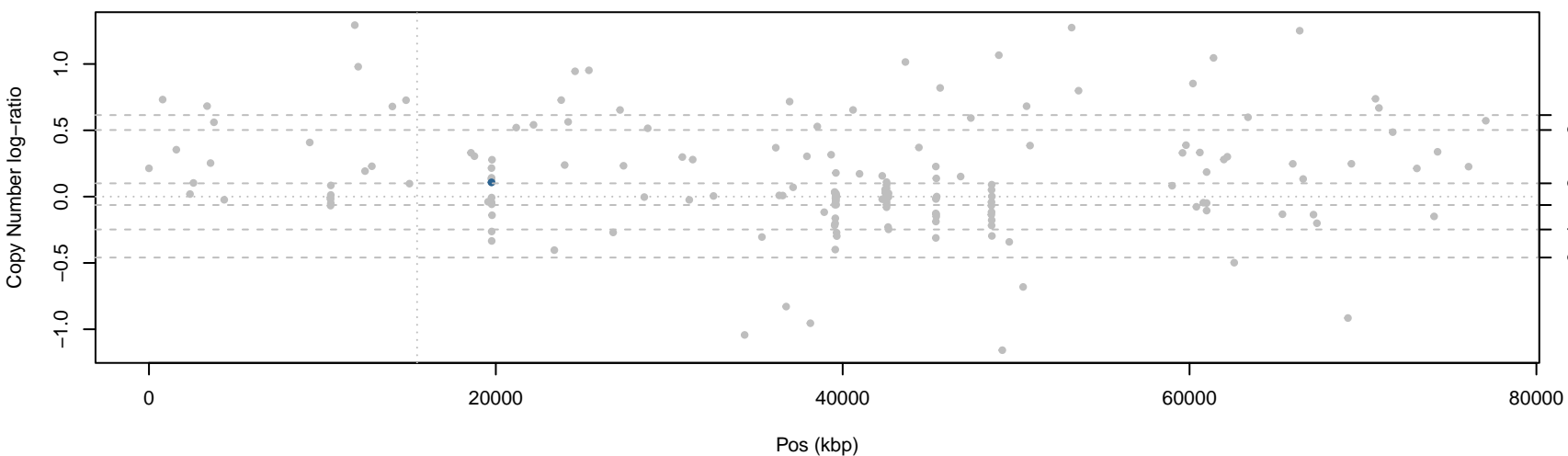
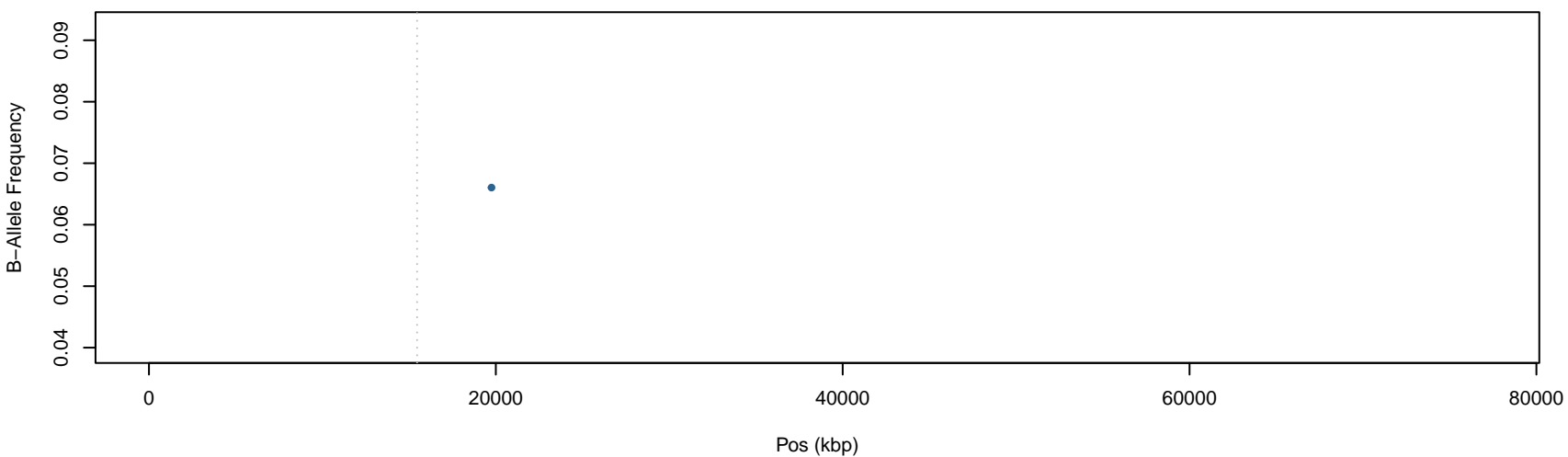
Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 15



Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 16

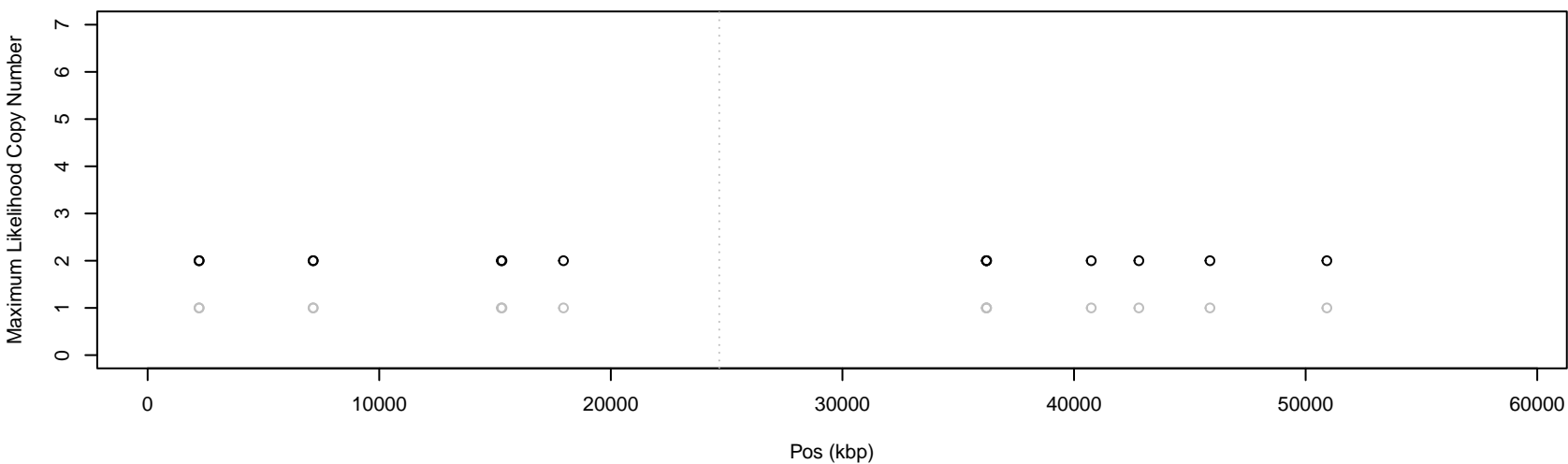
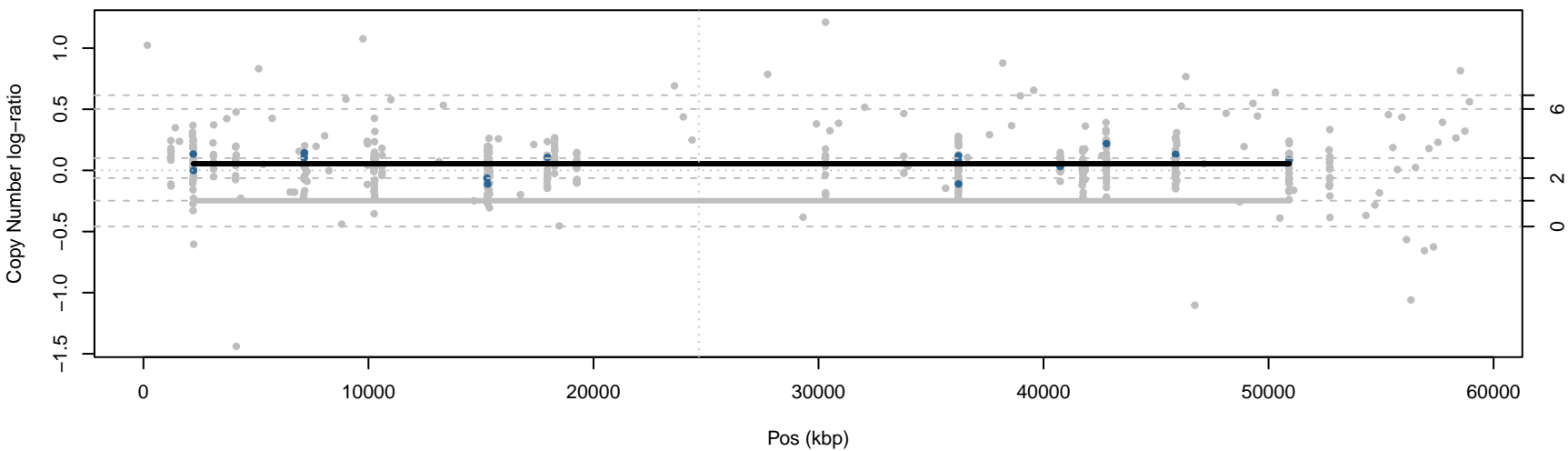
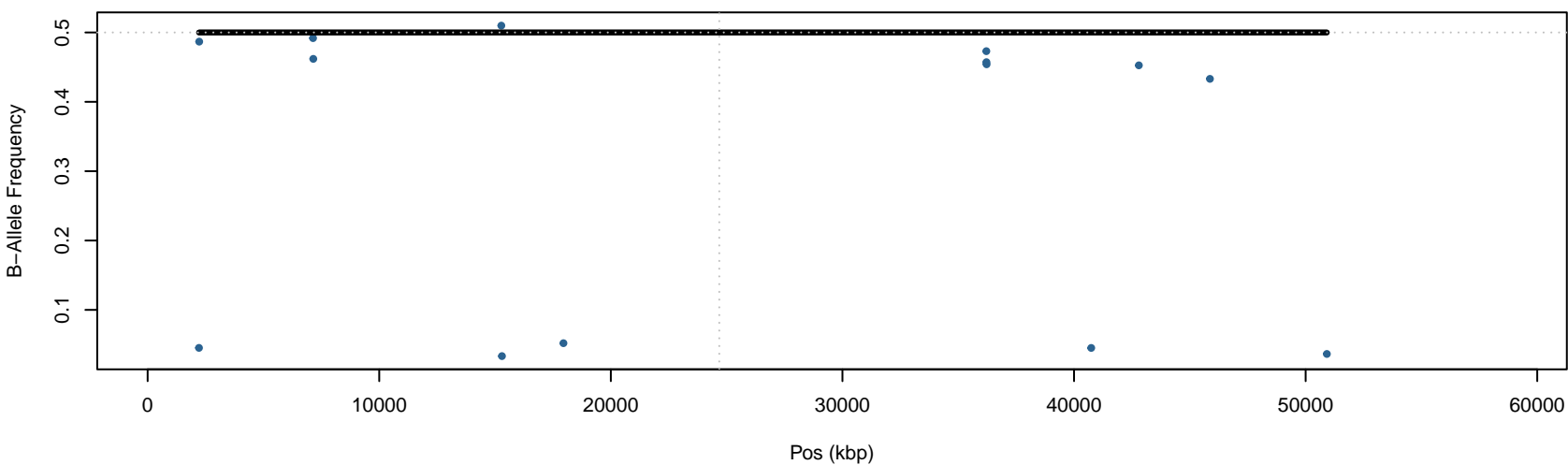


Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 18

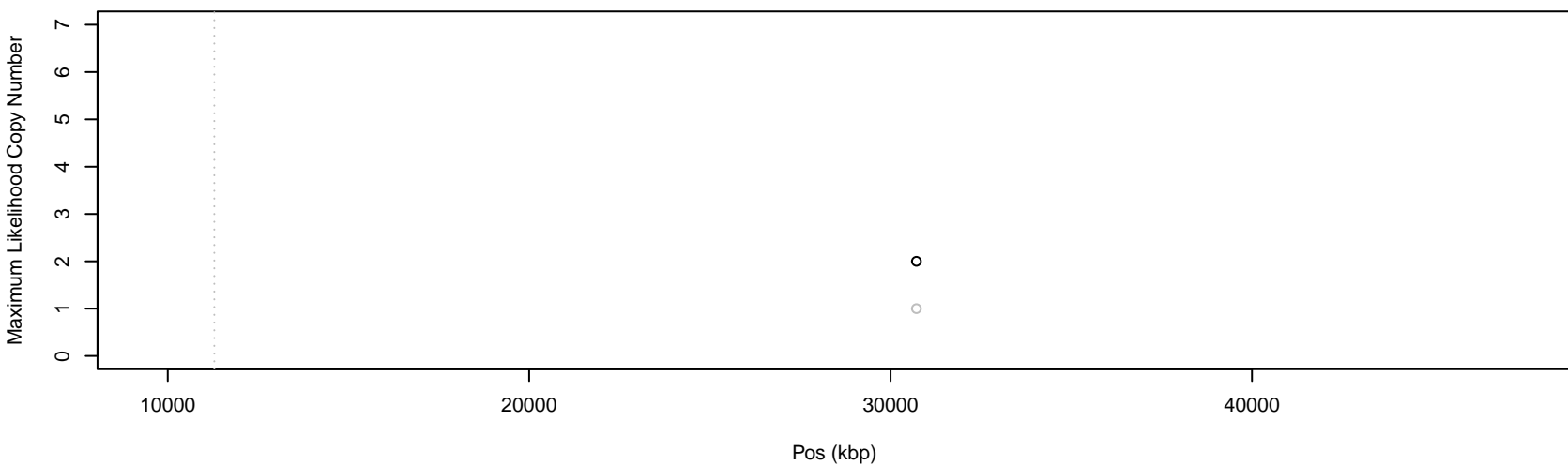
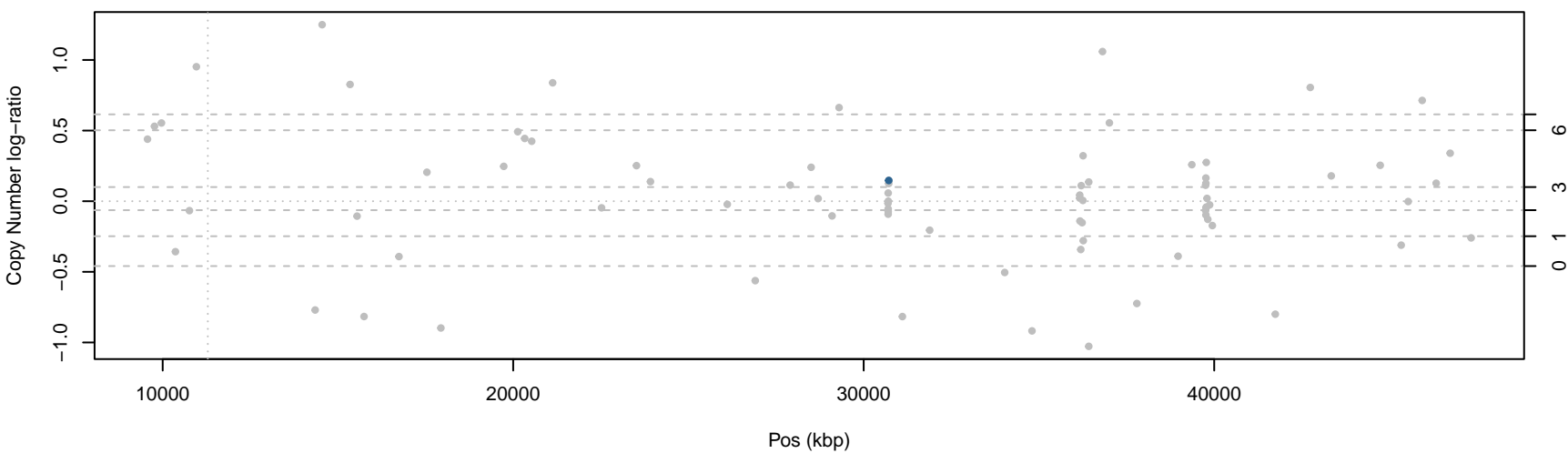
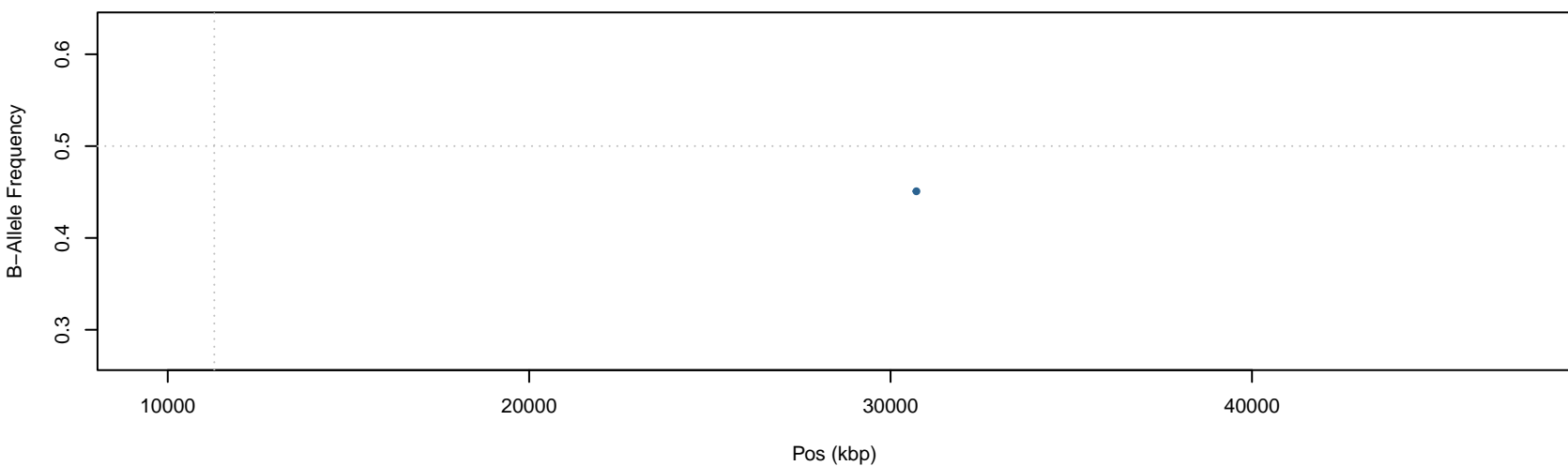




Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 19



Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 21



Purity: 0.24 Tumor ploidy: 2.374 SNV log-likelihood: 208.81 GoF: 90.8% Mean coverage: 306;377 Chromosome: 22

