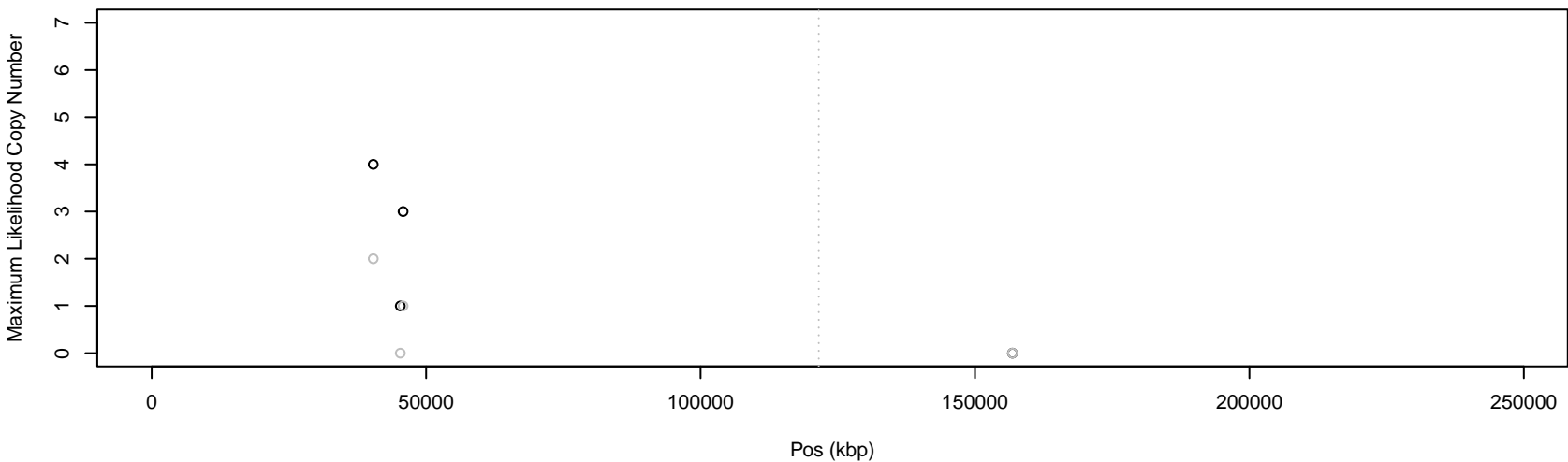
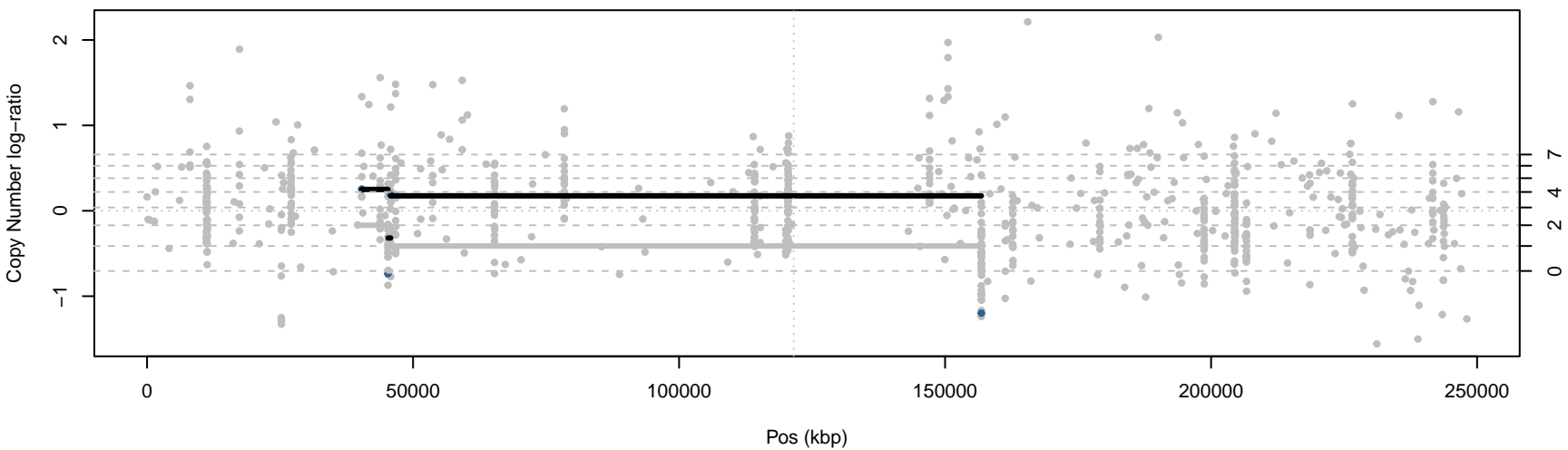
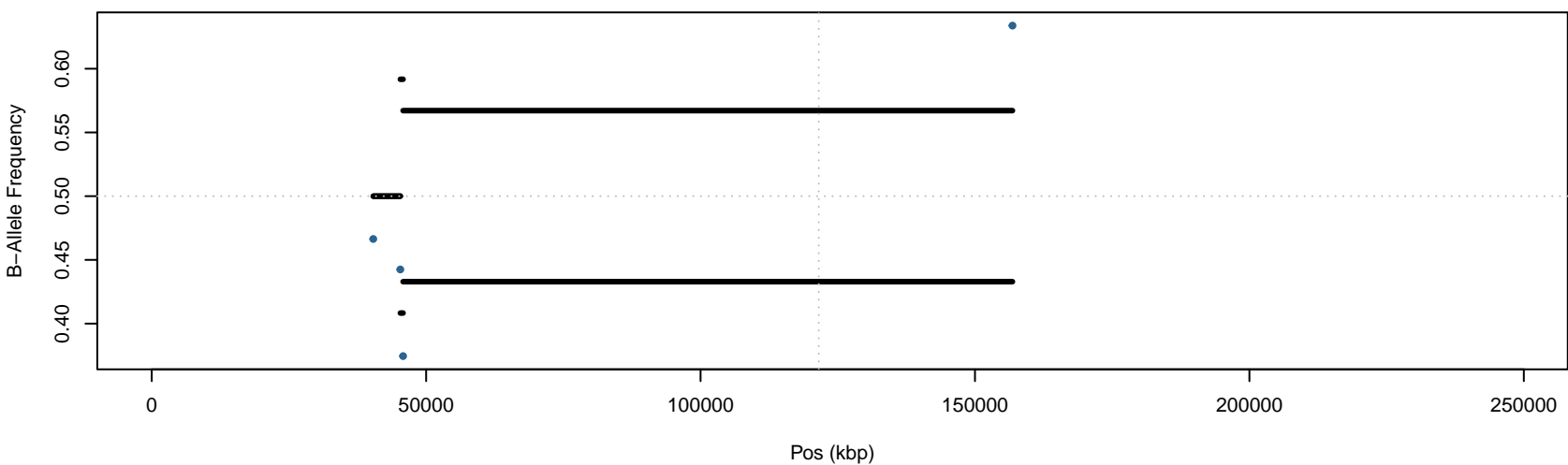
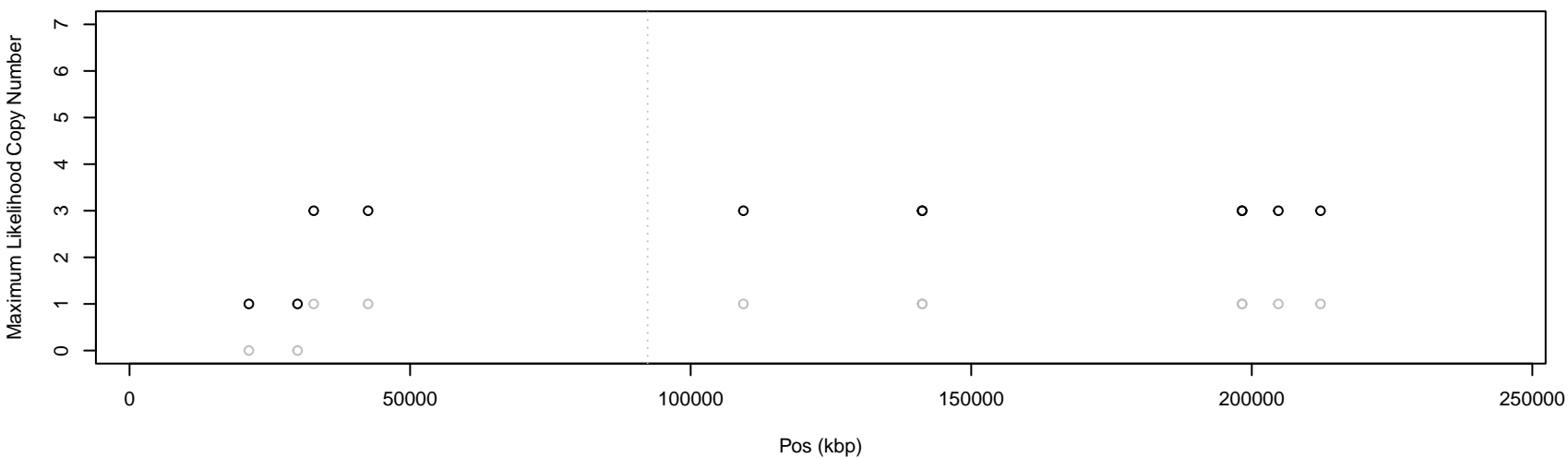
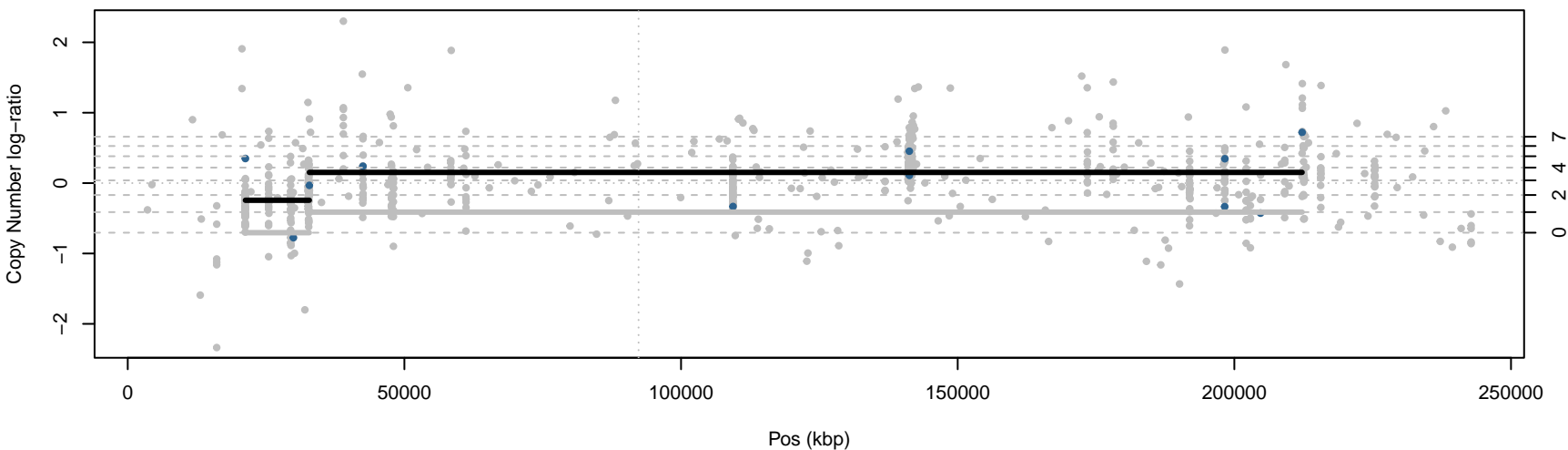
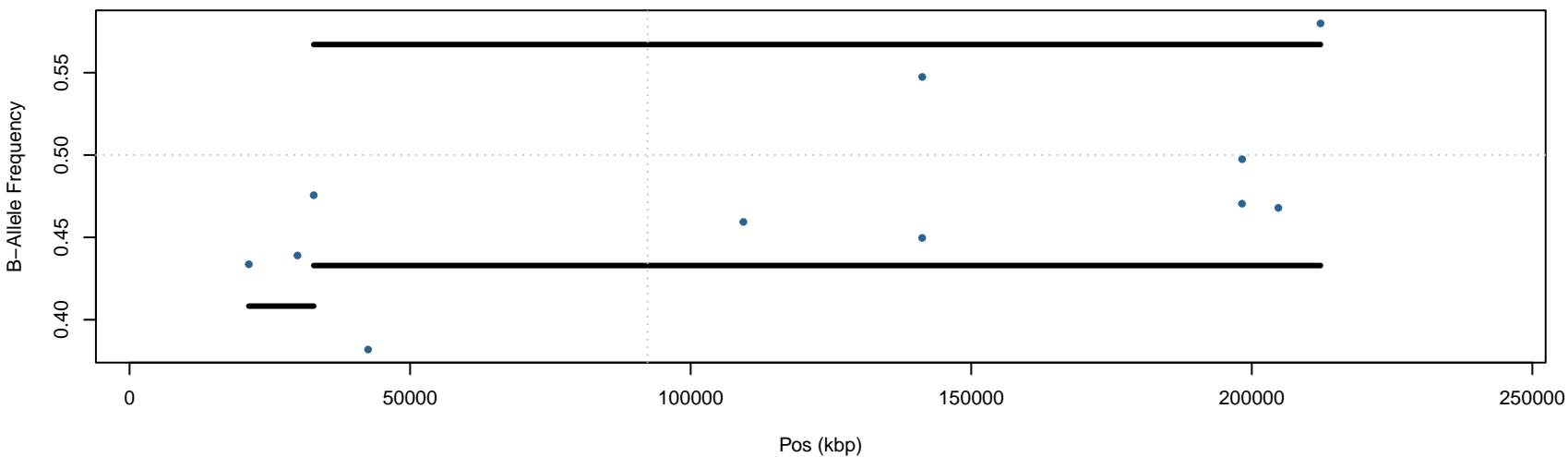


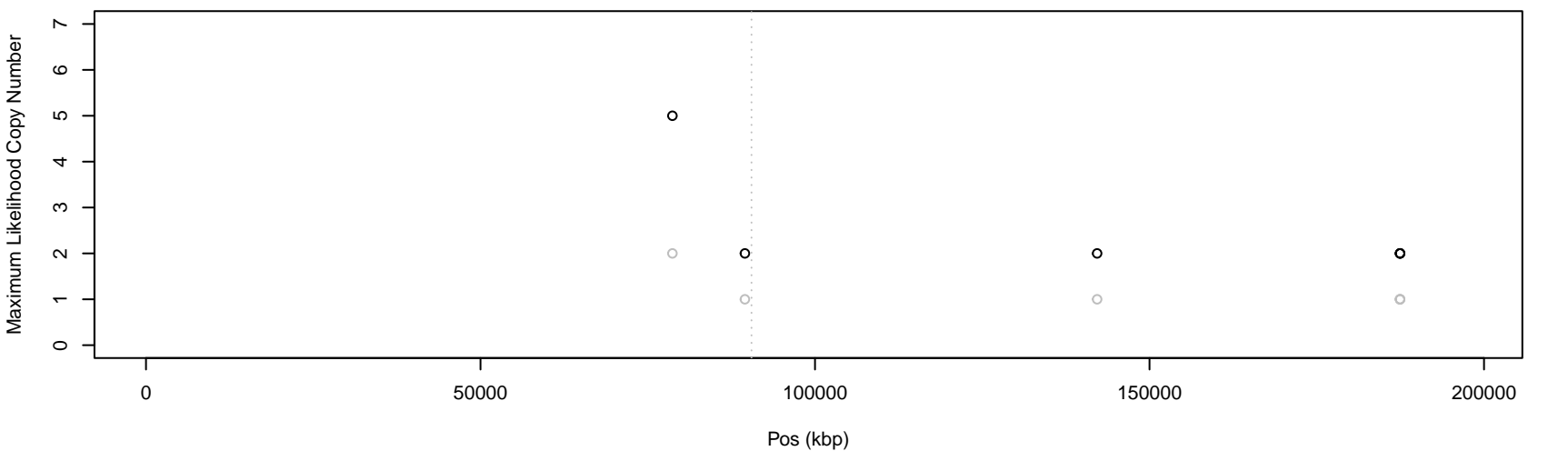
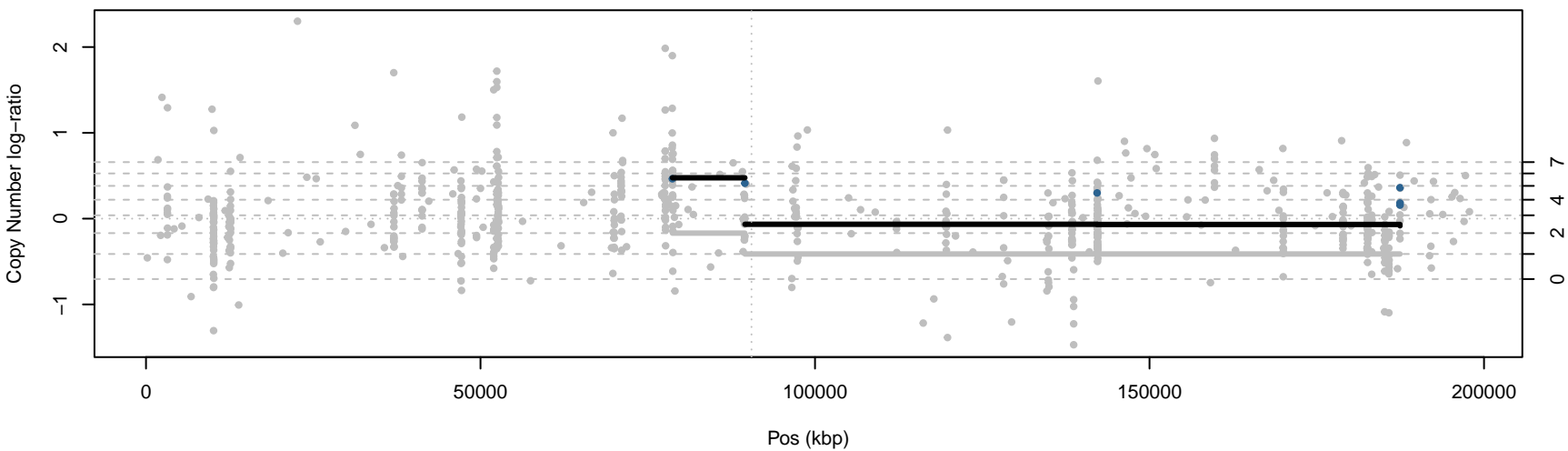
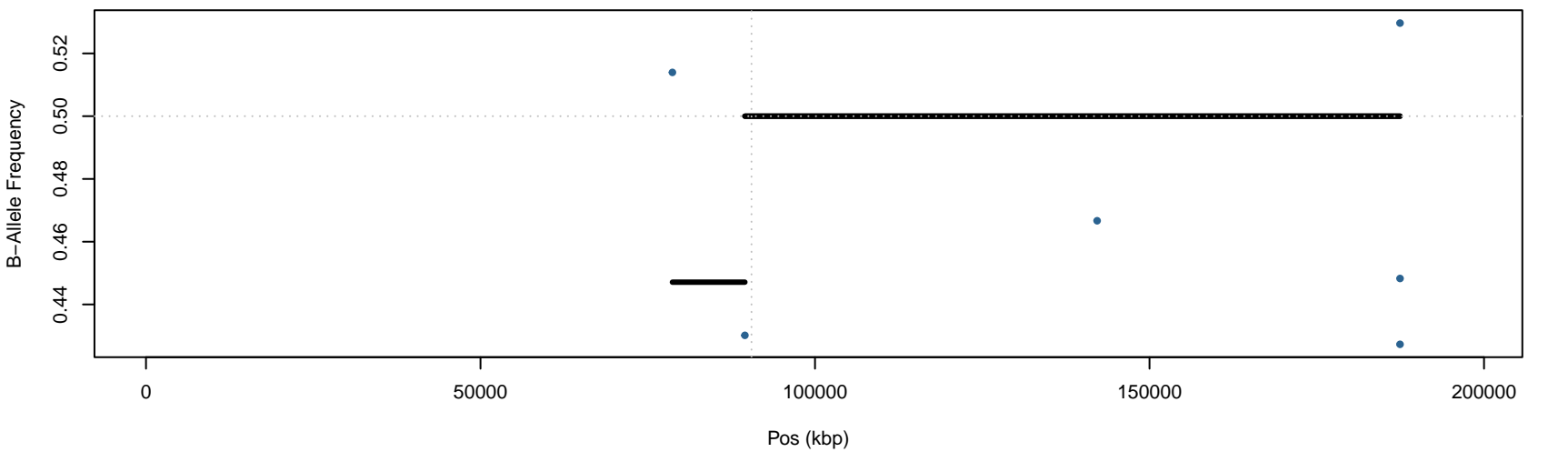
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 1



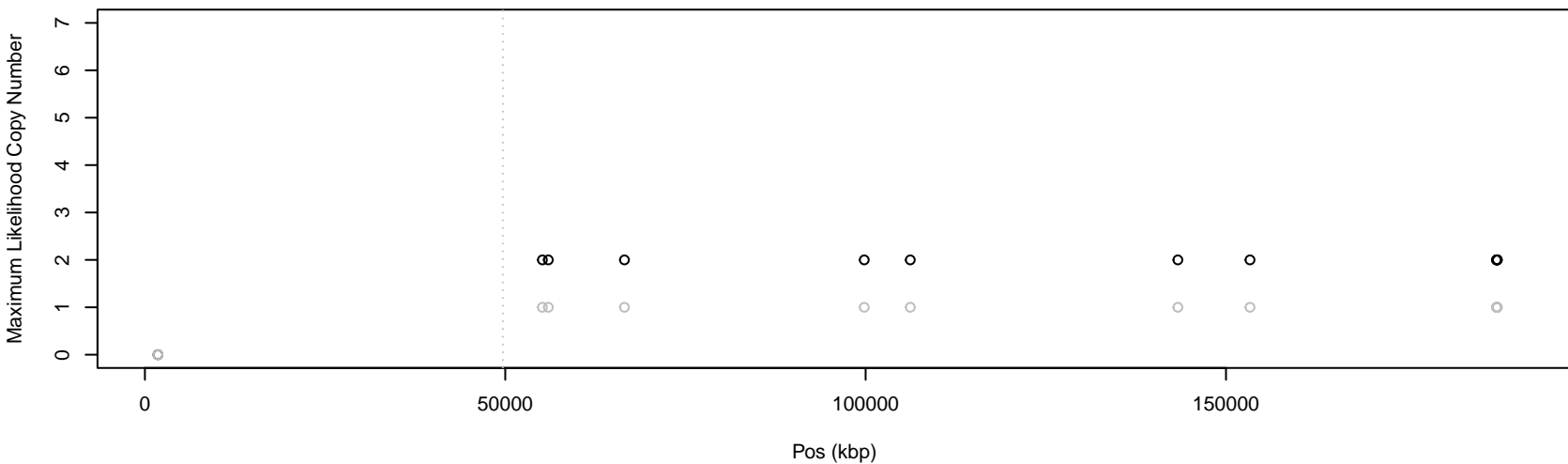
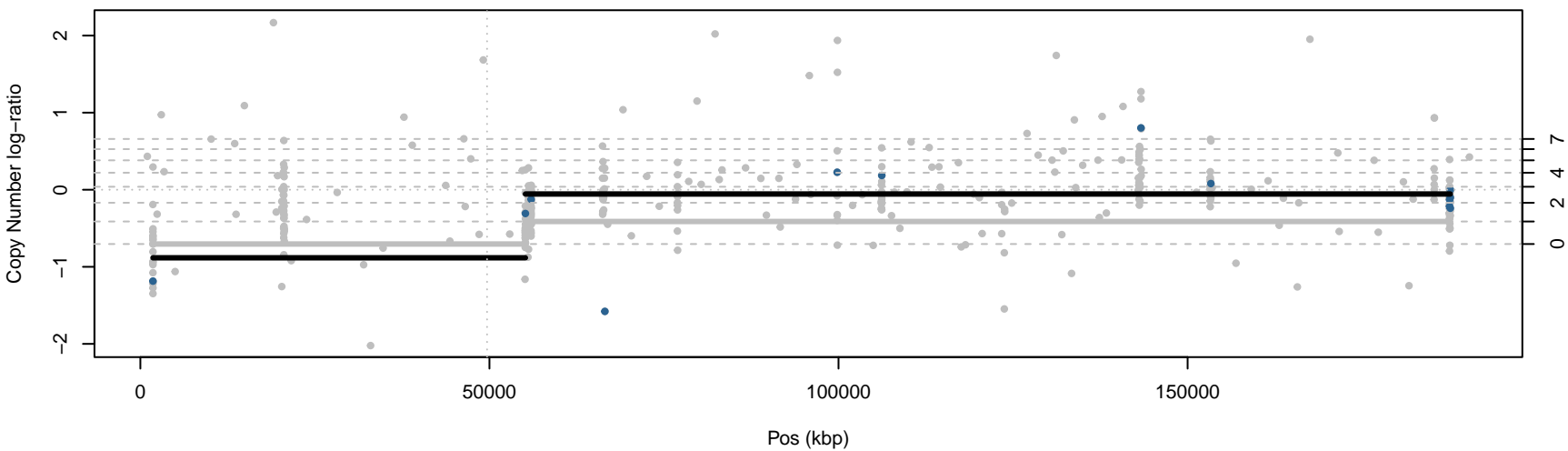
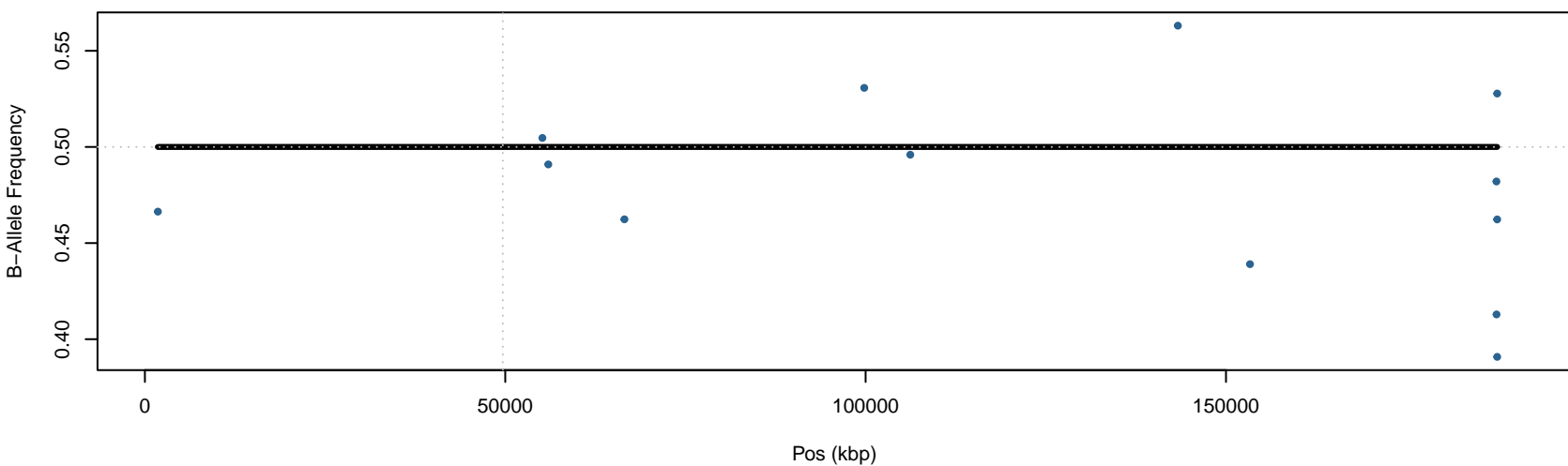
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 2



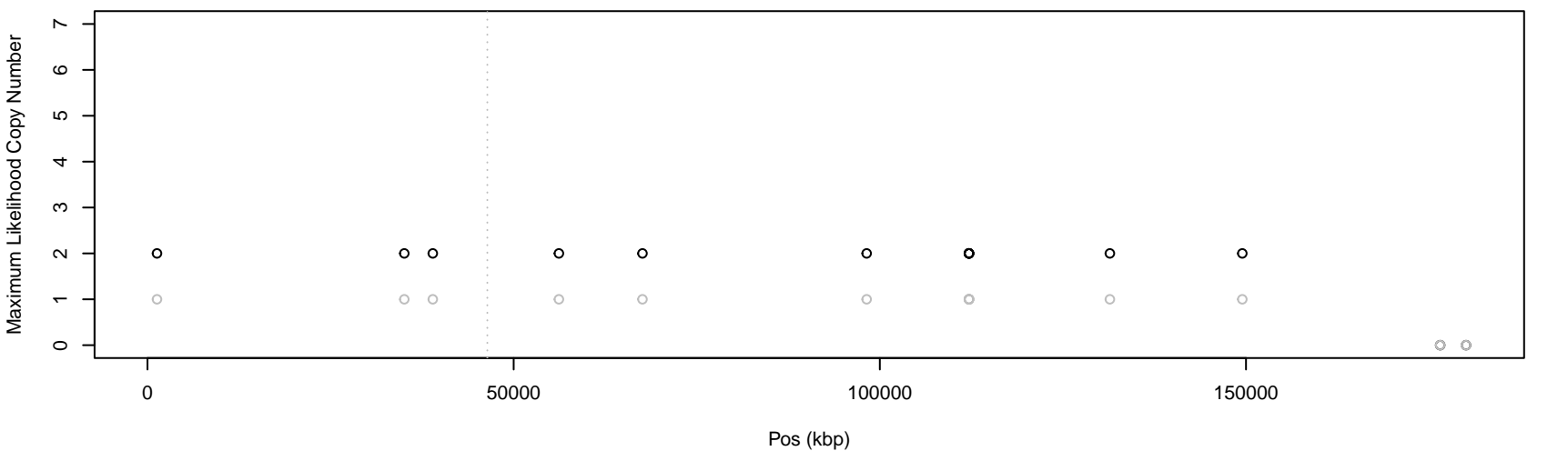
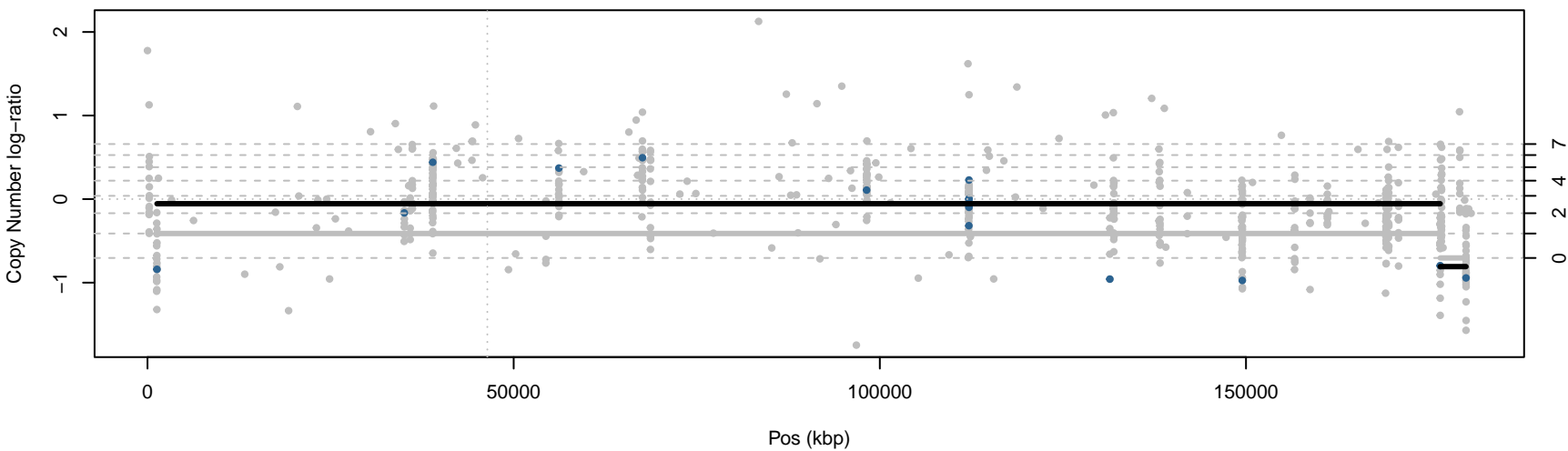
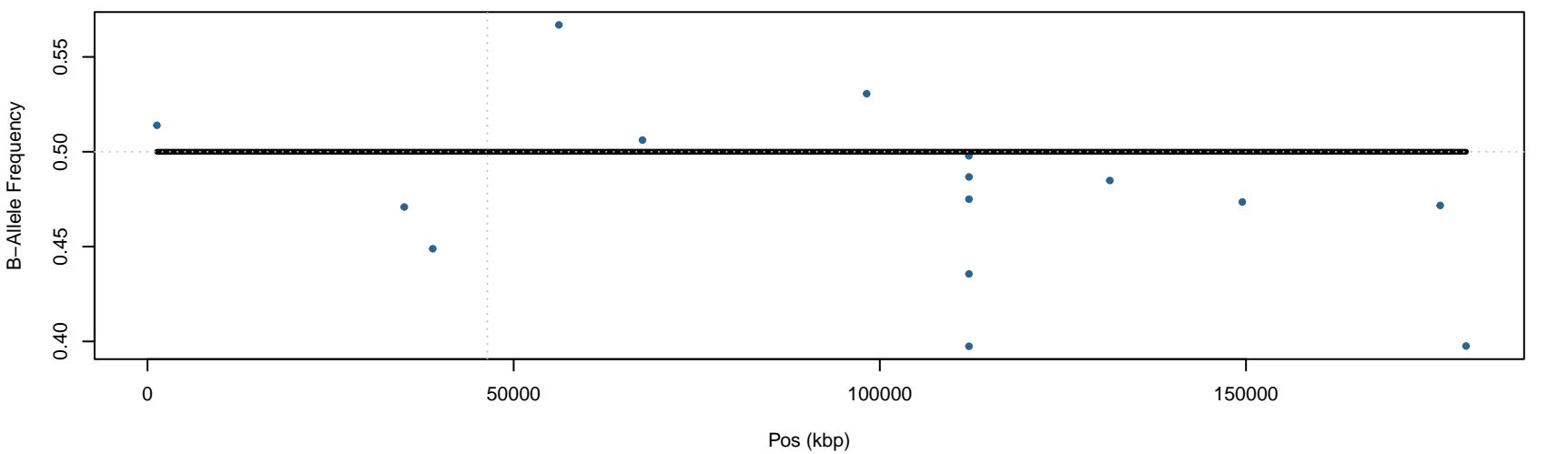
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 3



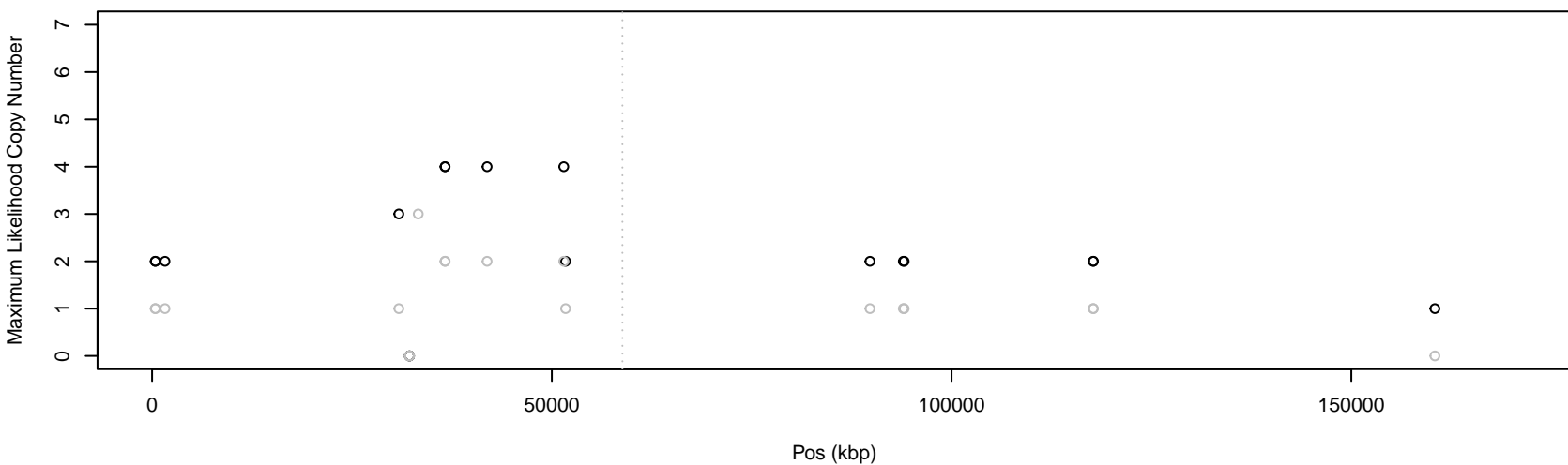
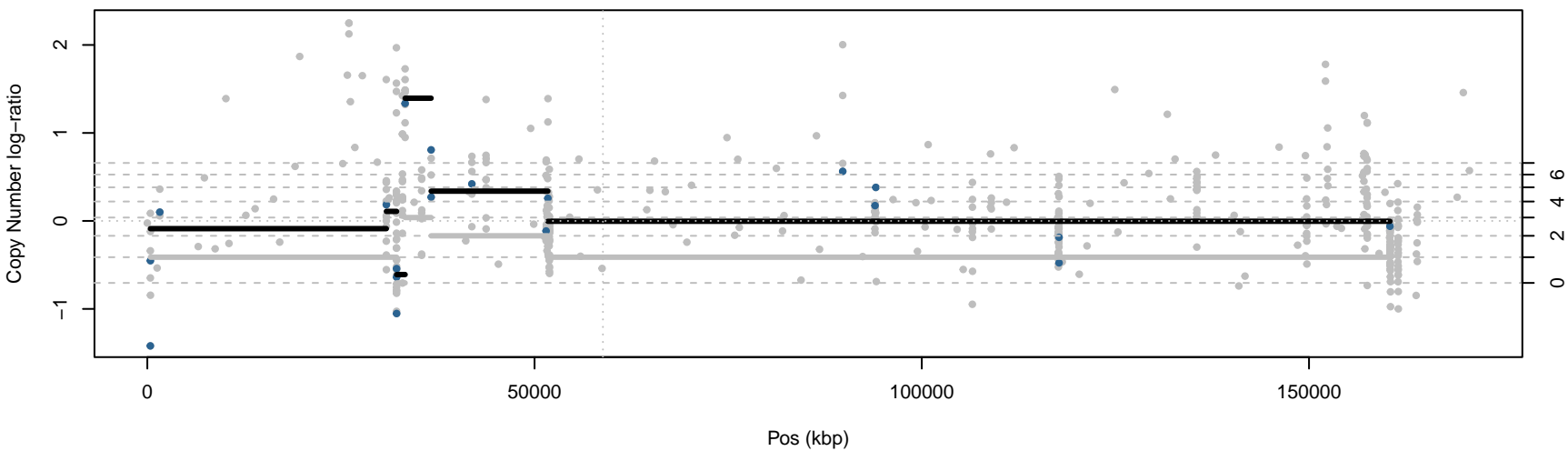
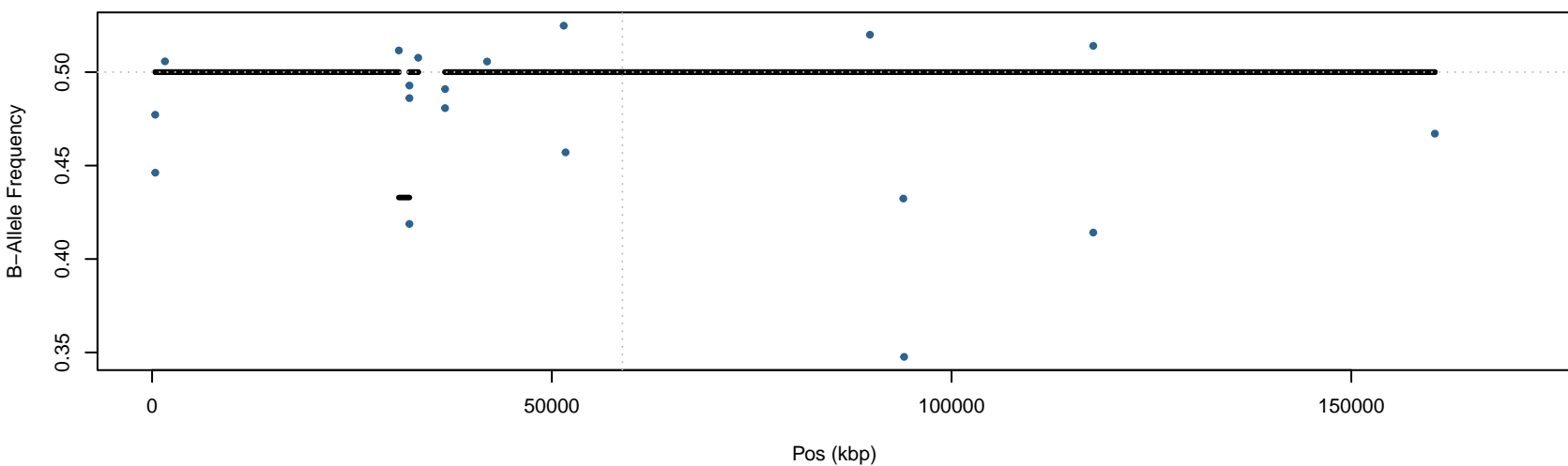
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 4



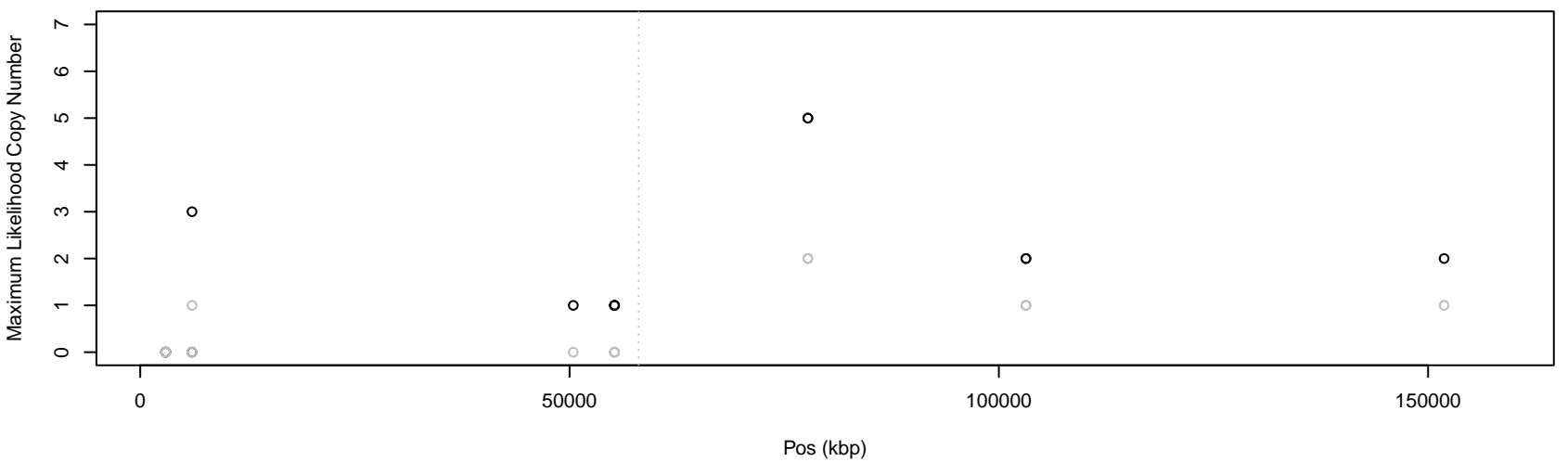
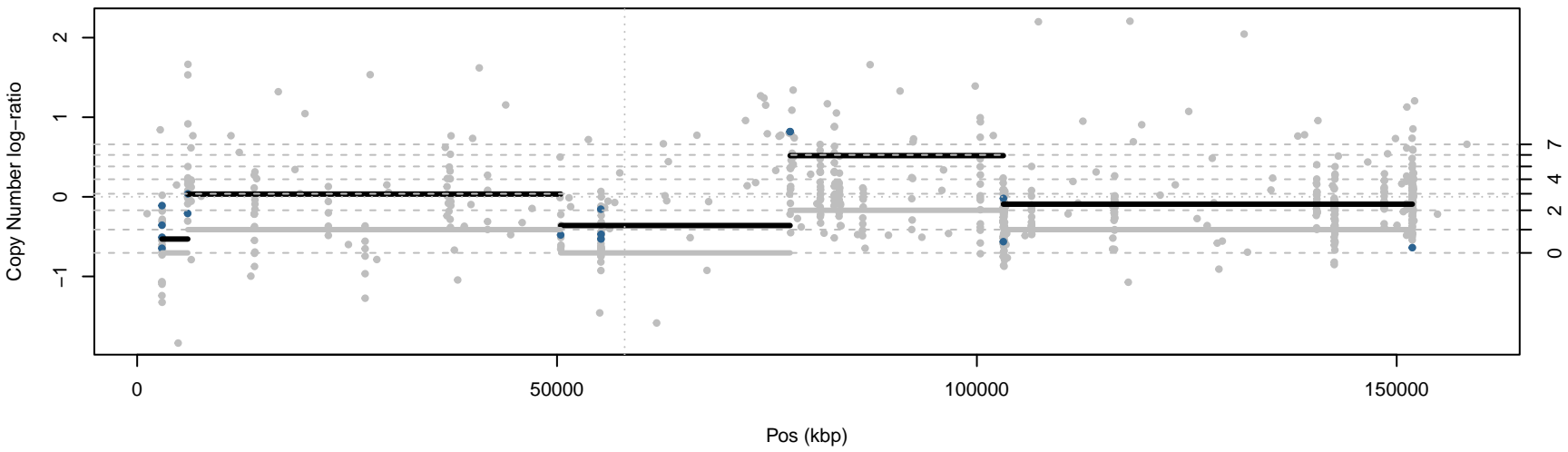
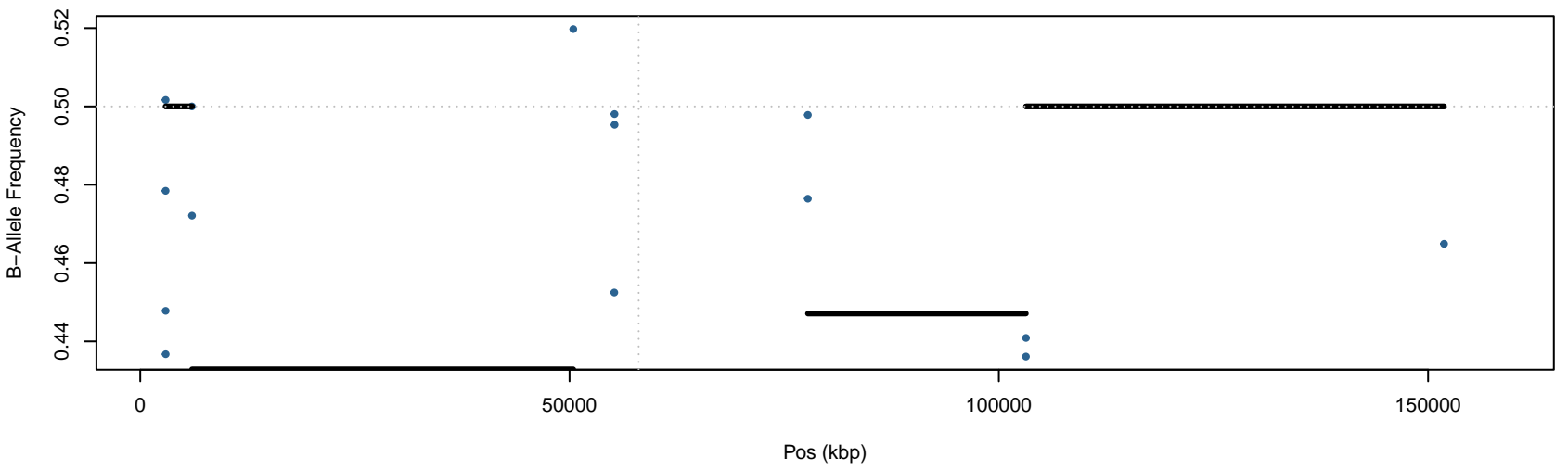
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 5



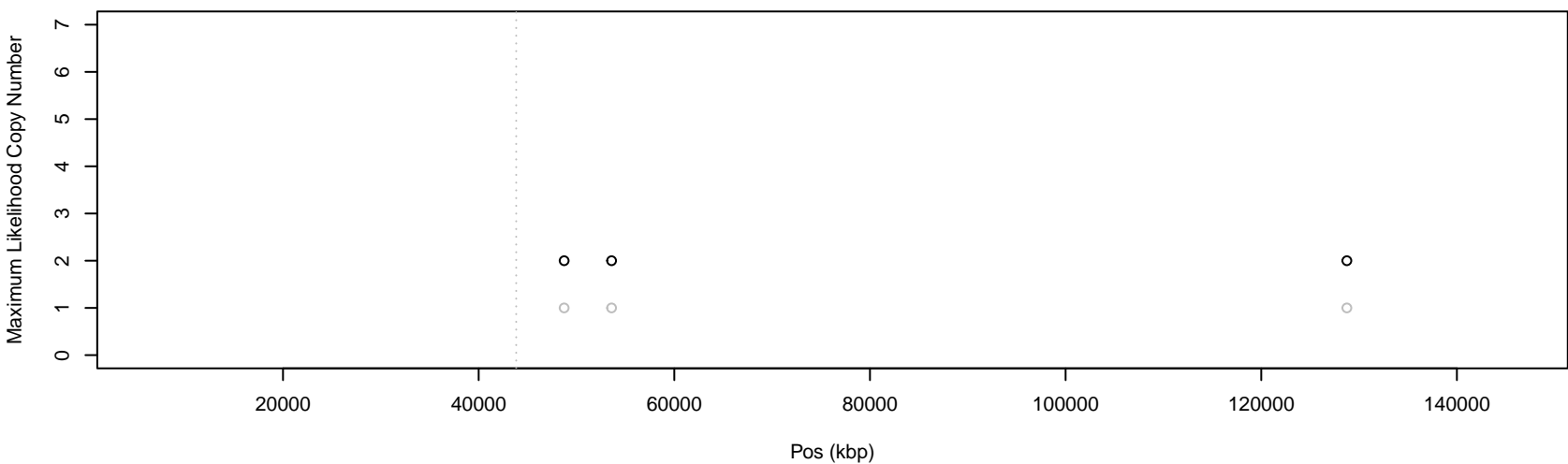
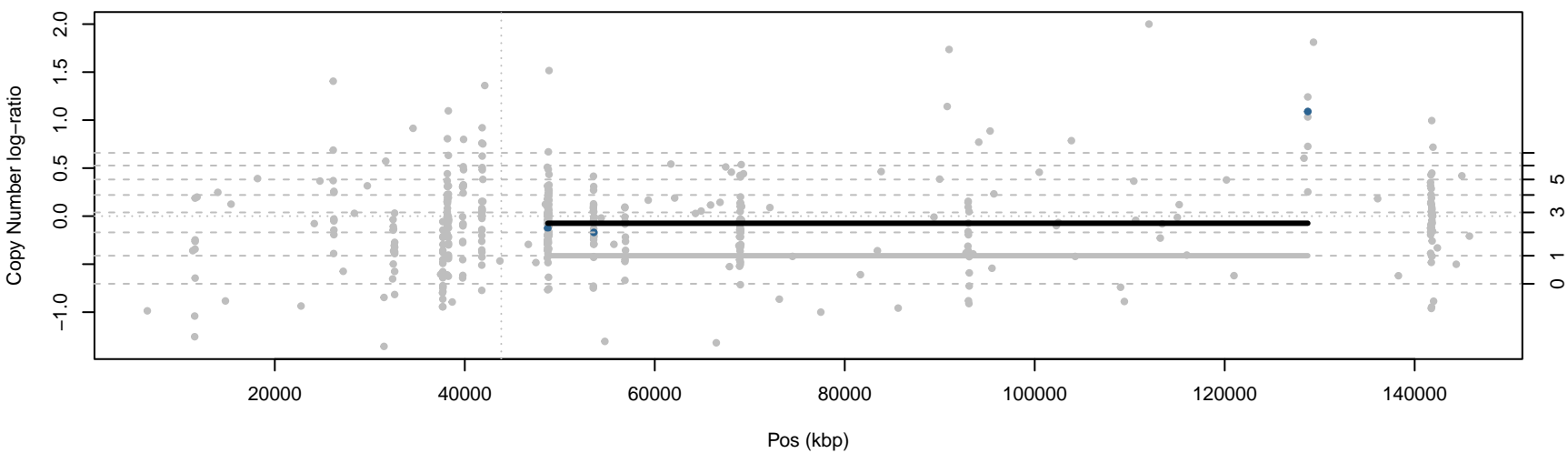
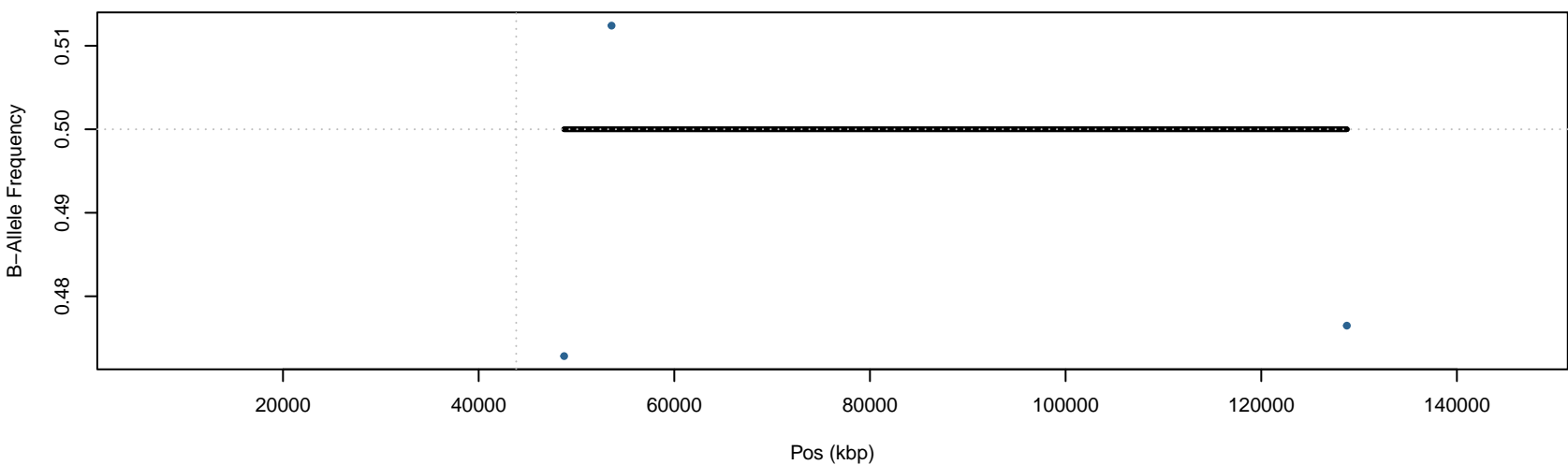
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 6



Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 7

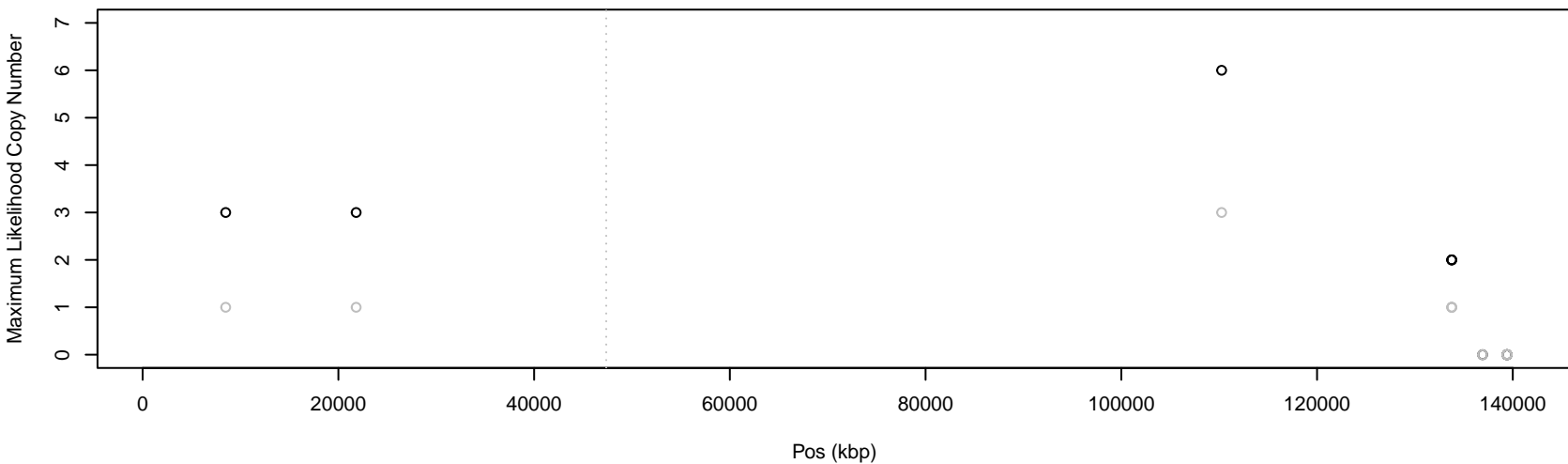
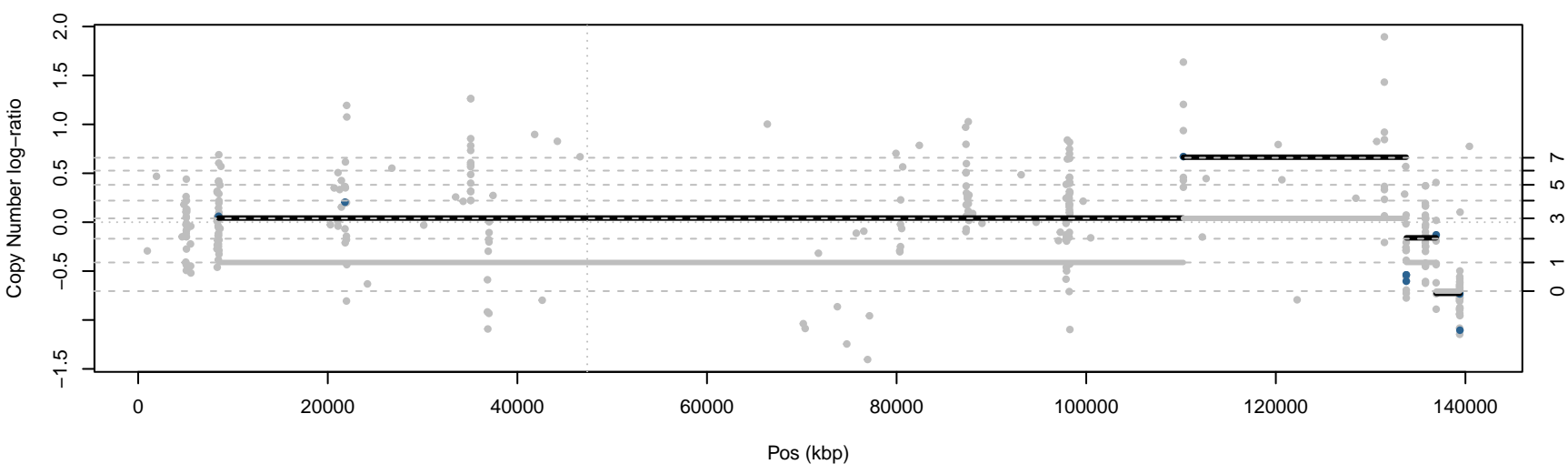
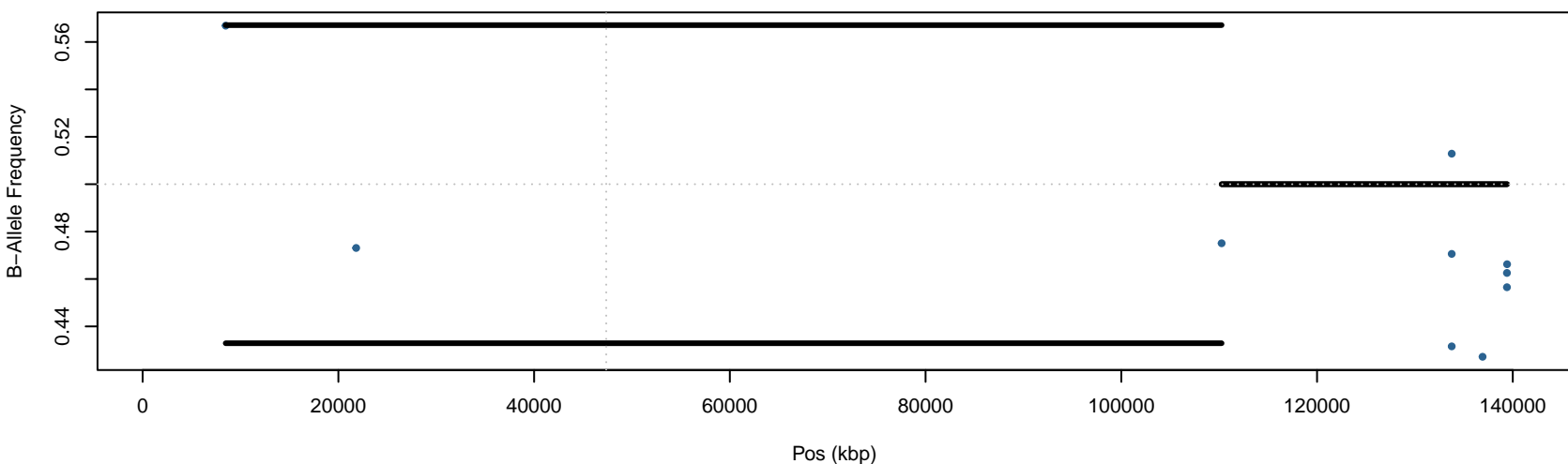


Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 8

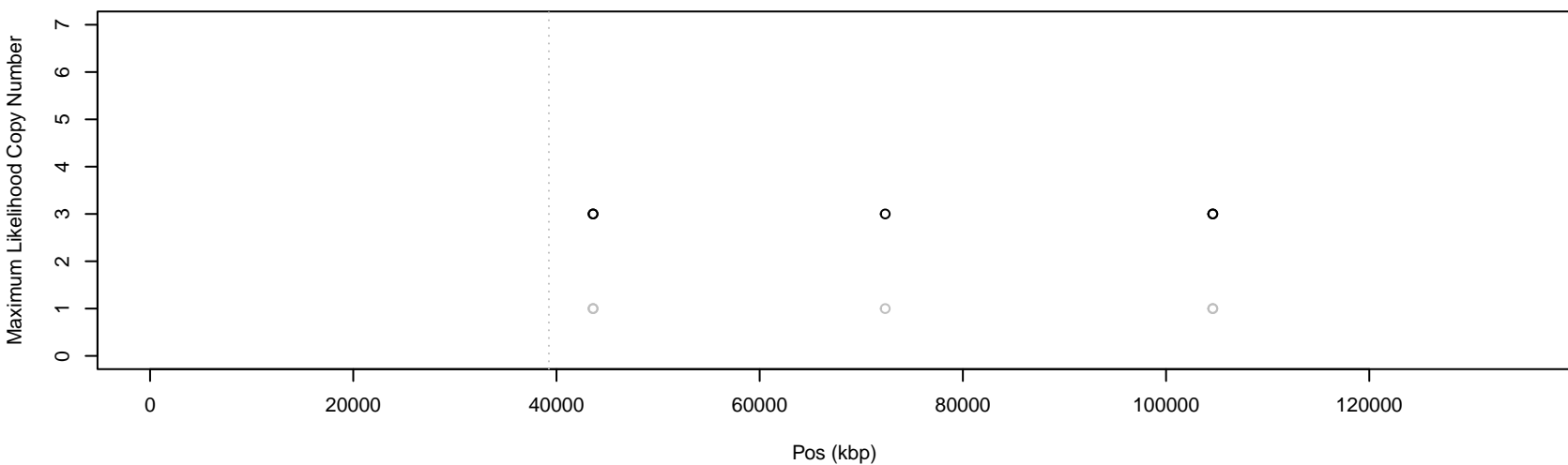
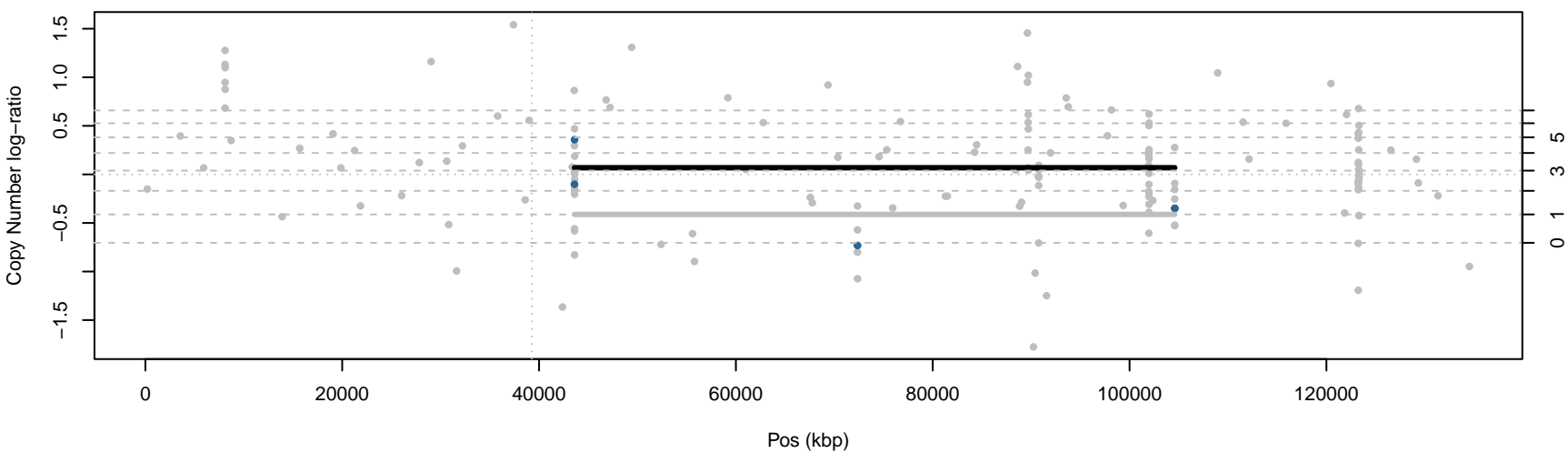
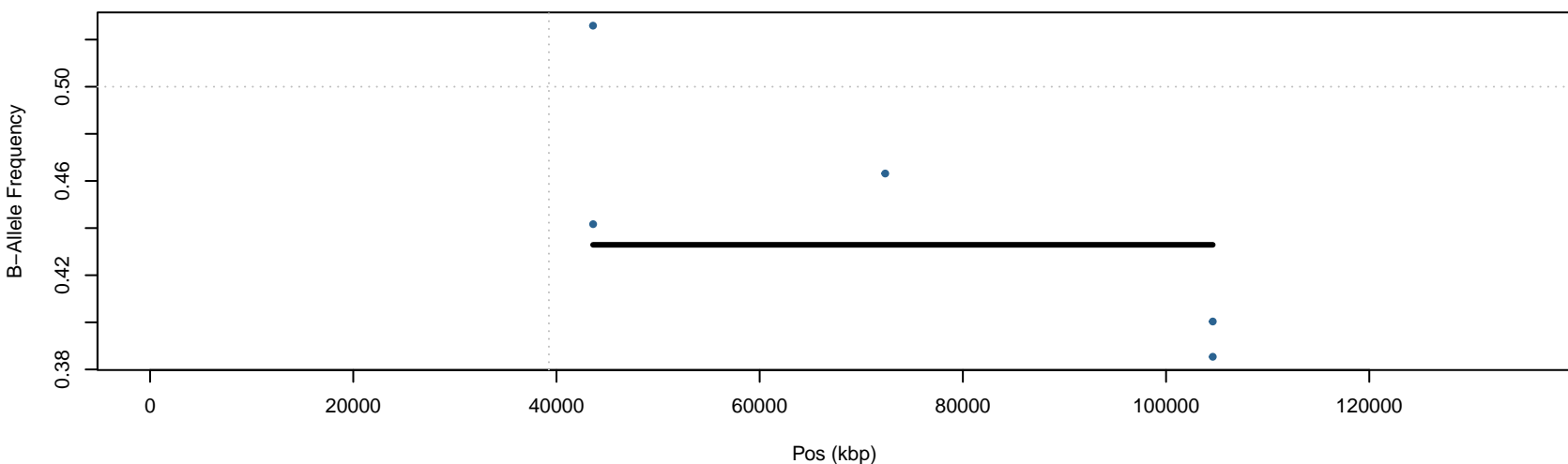




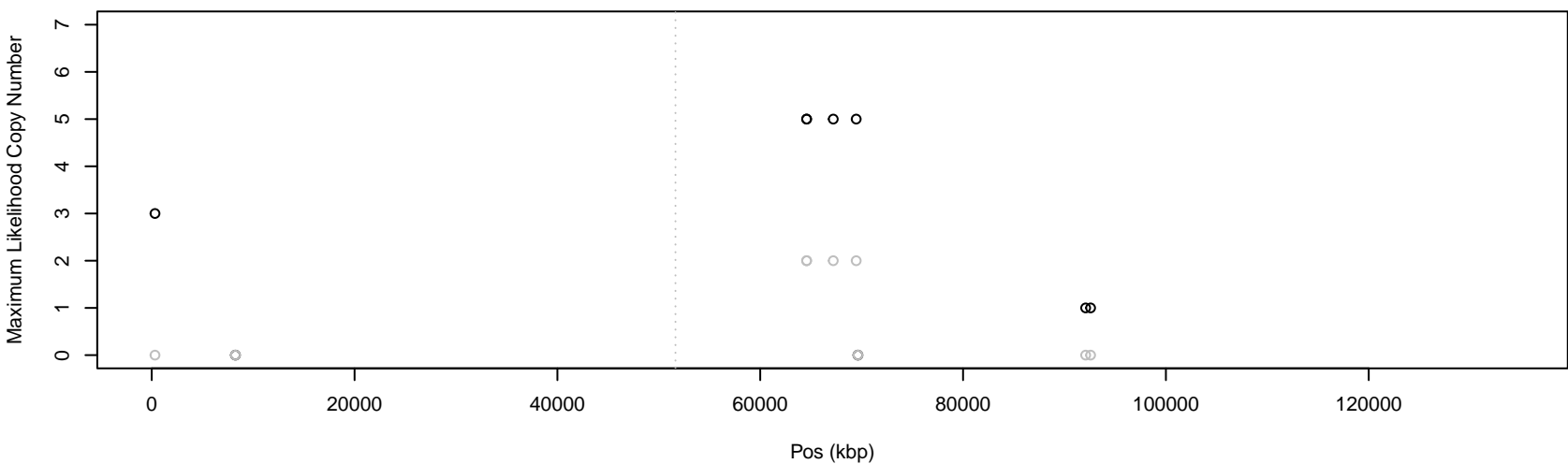
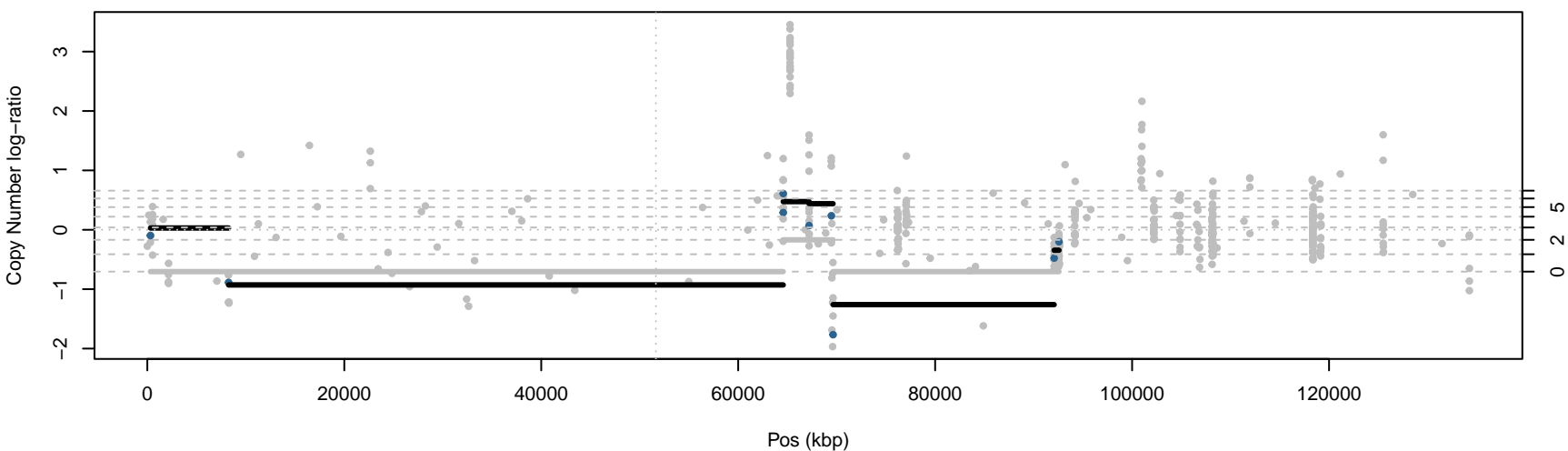
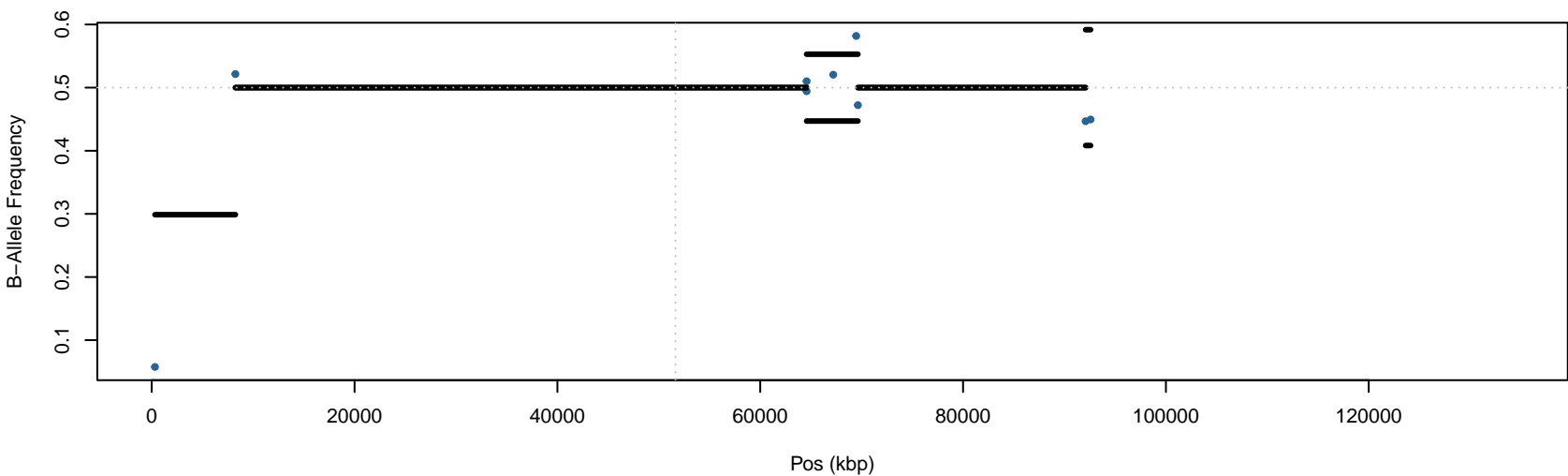
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 9



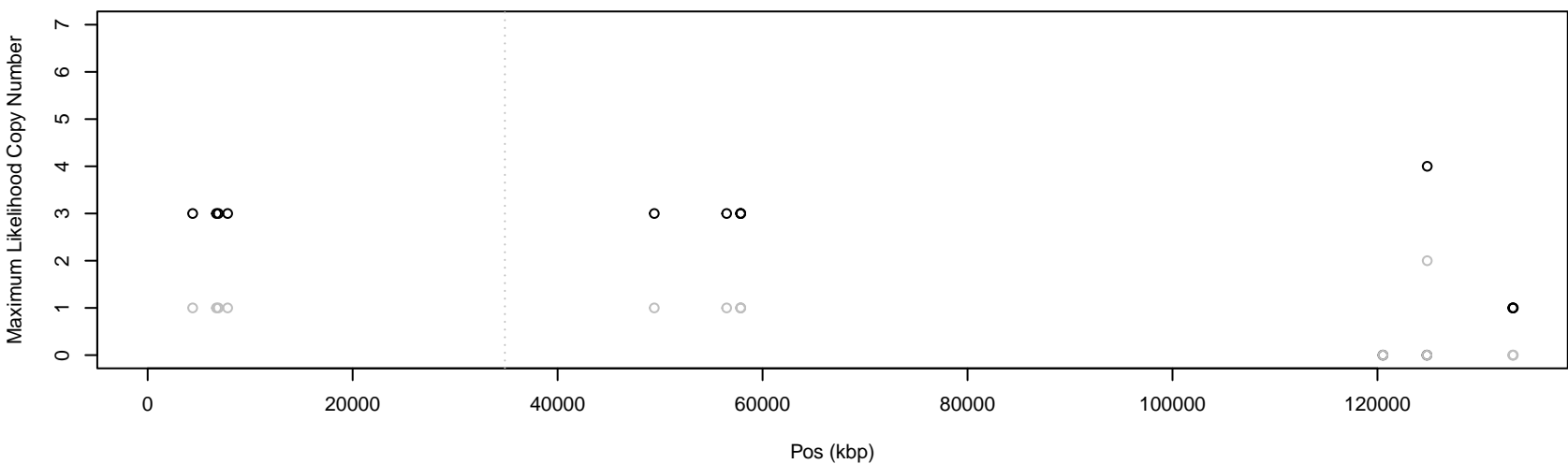
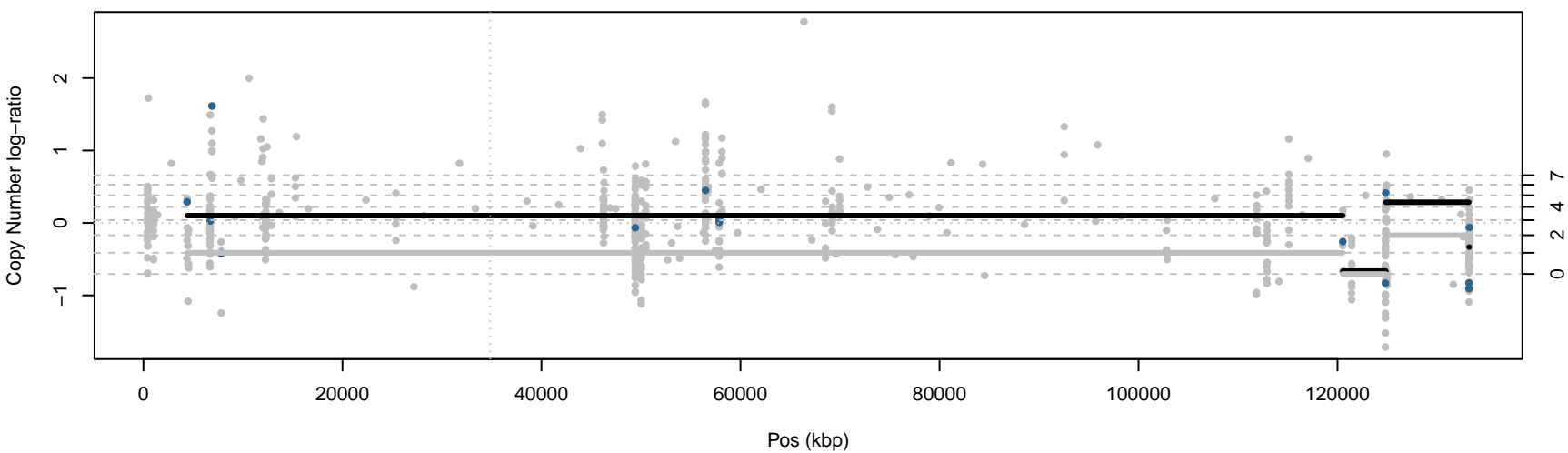
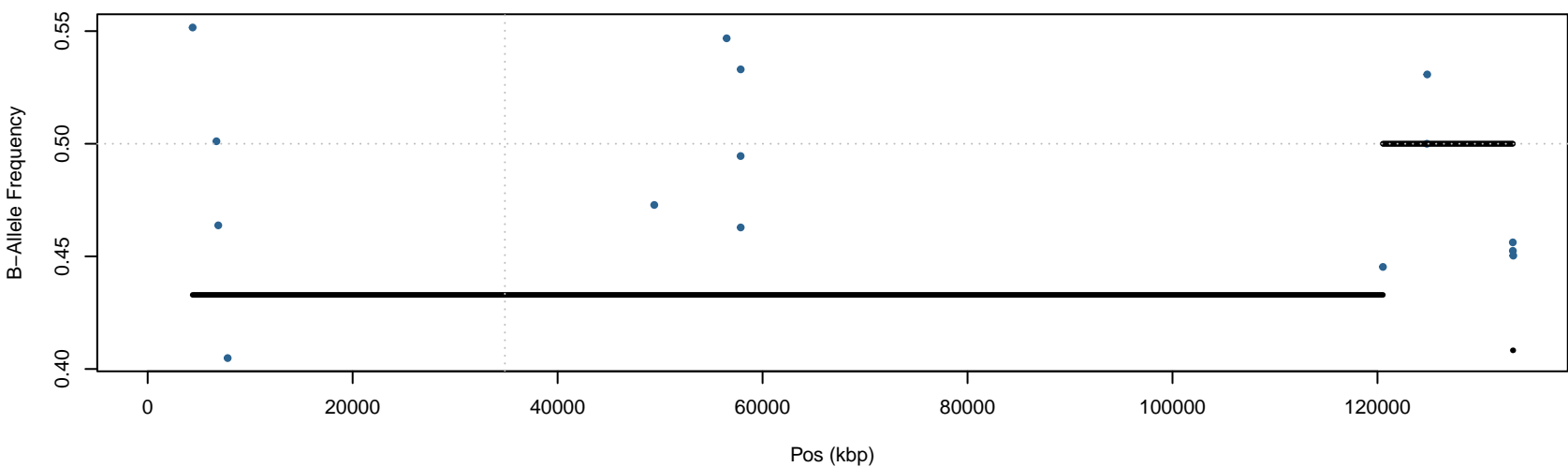
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 10



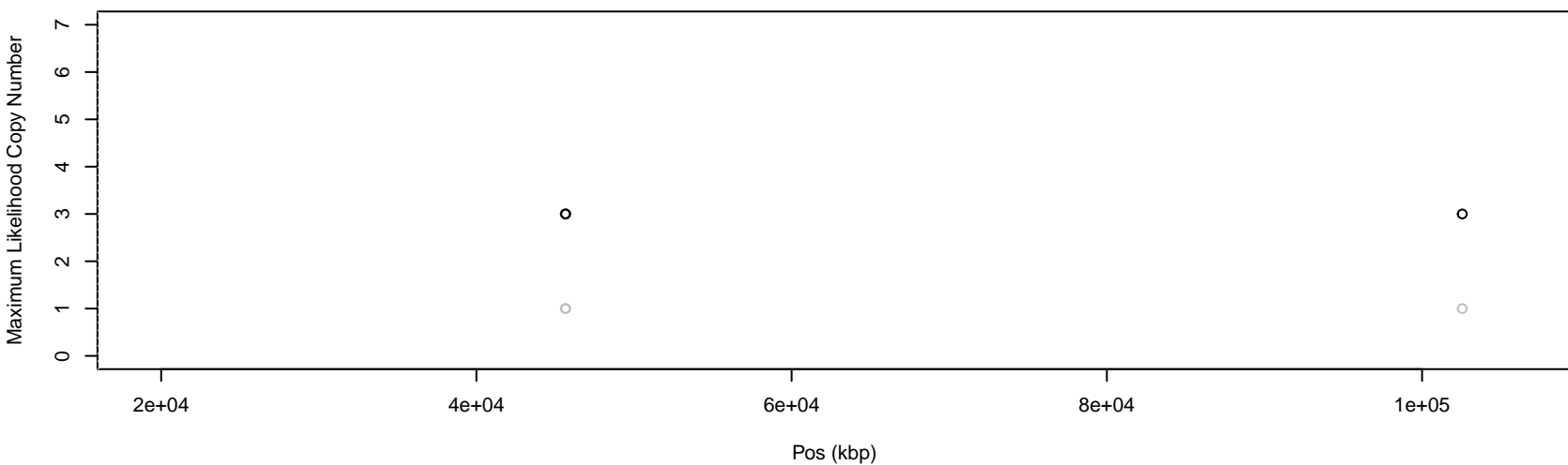
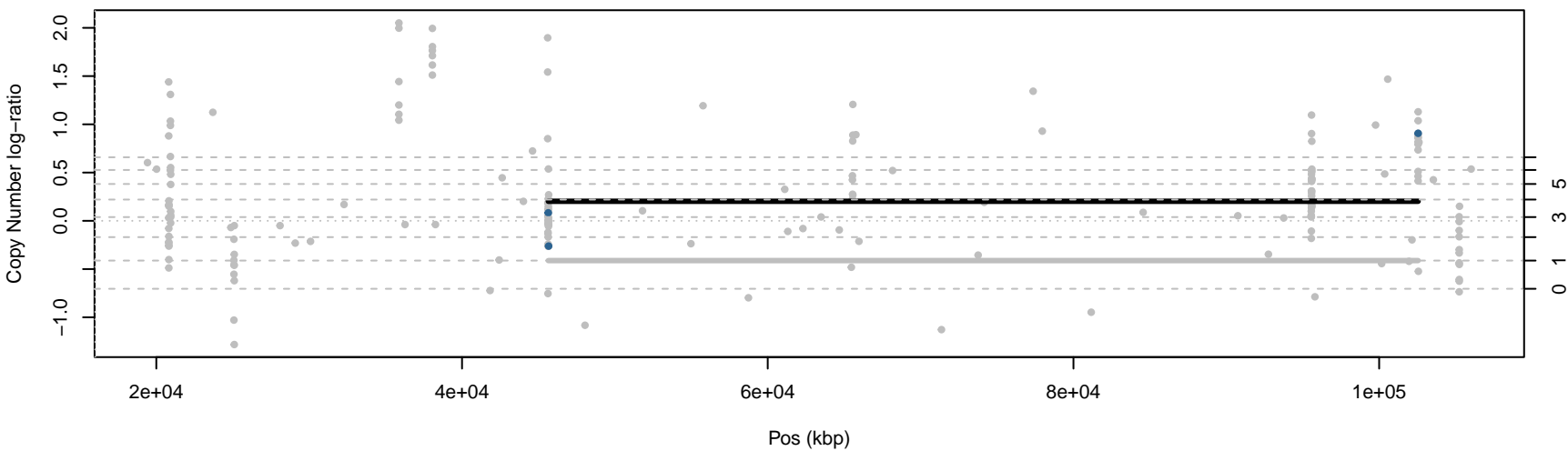
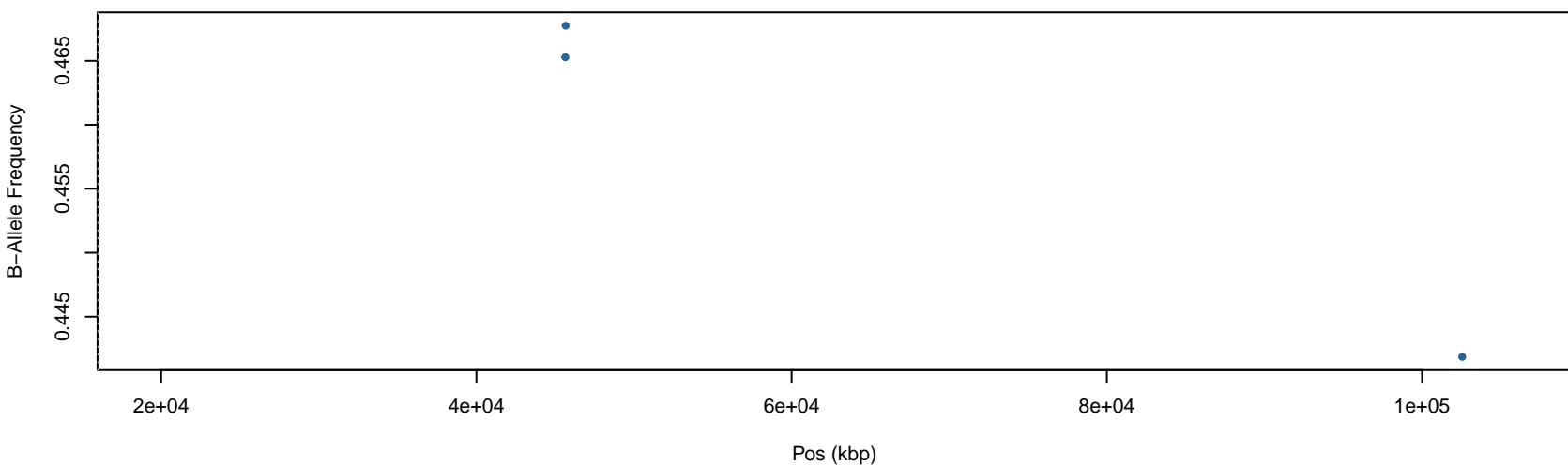
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 11



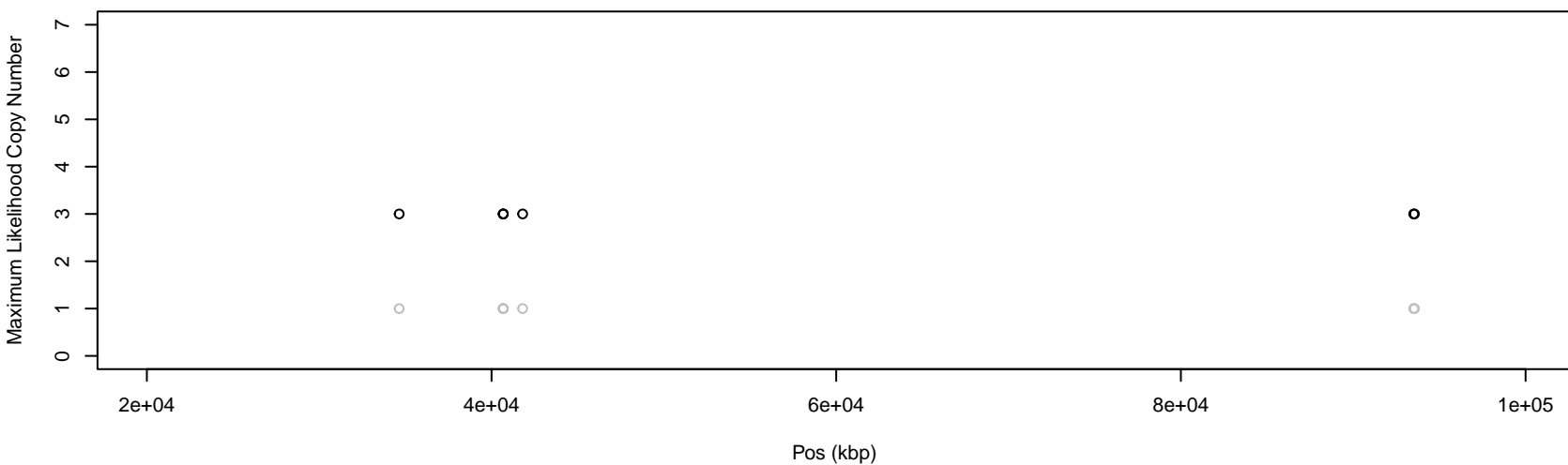
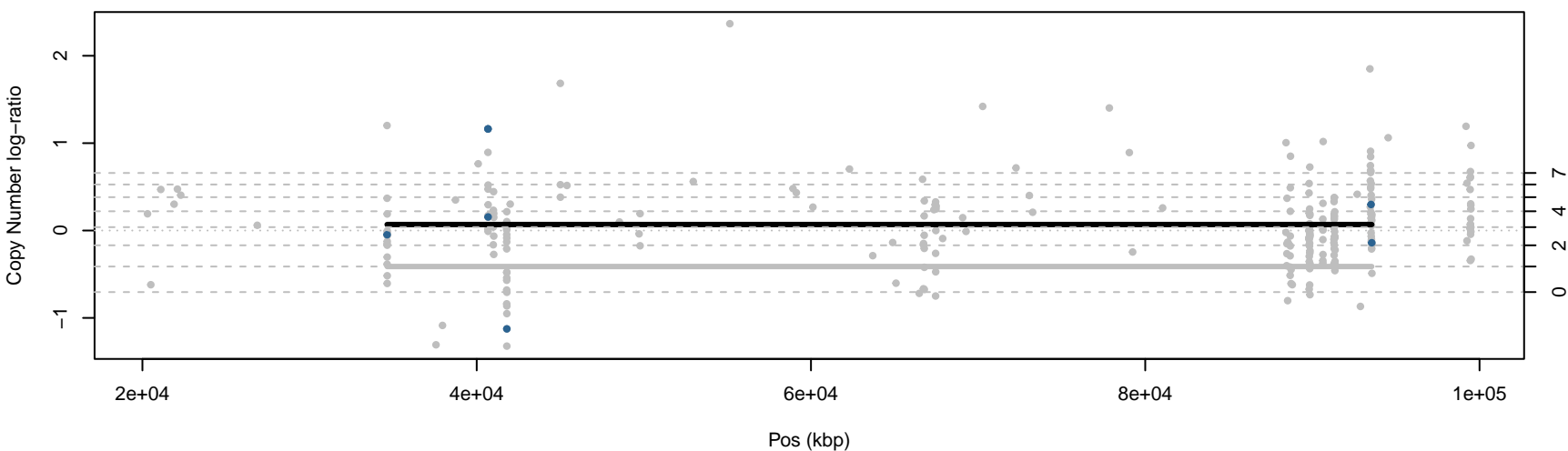
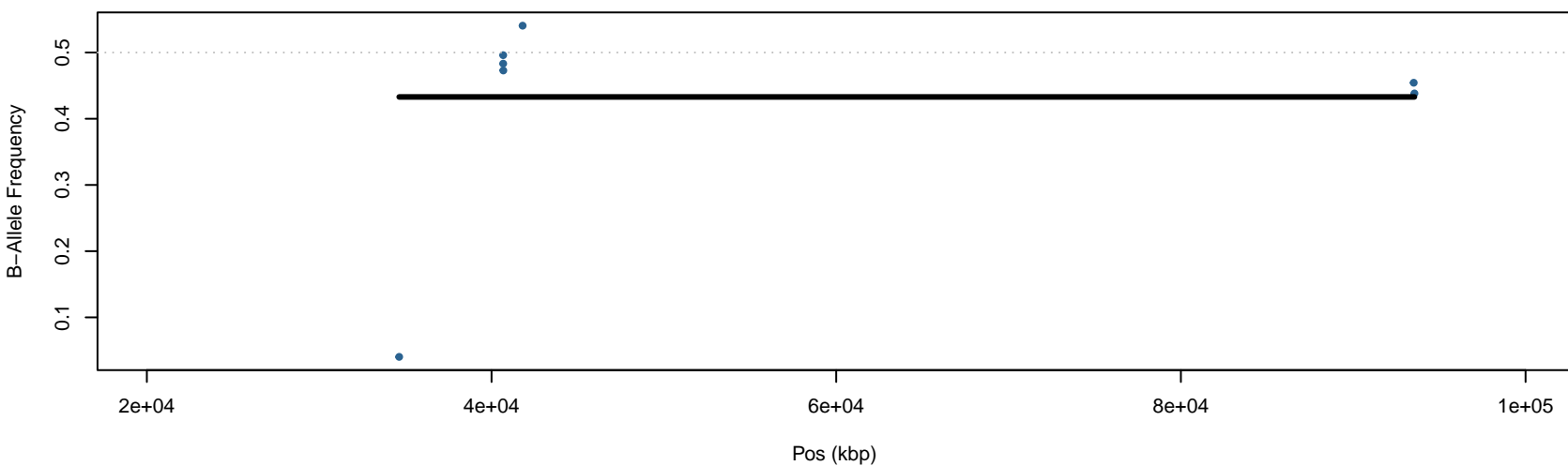
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 12



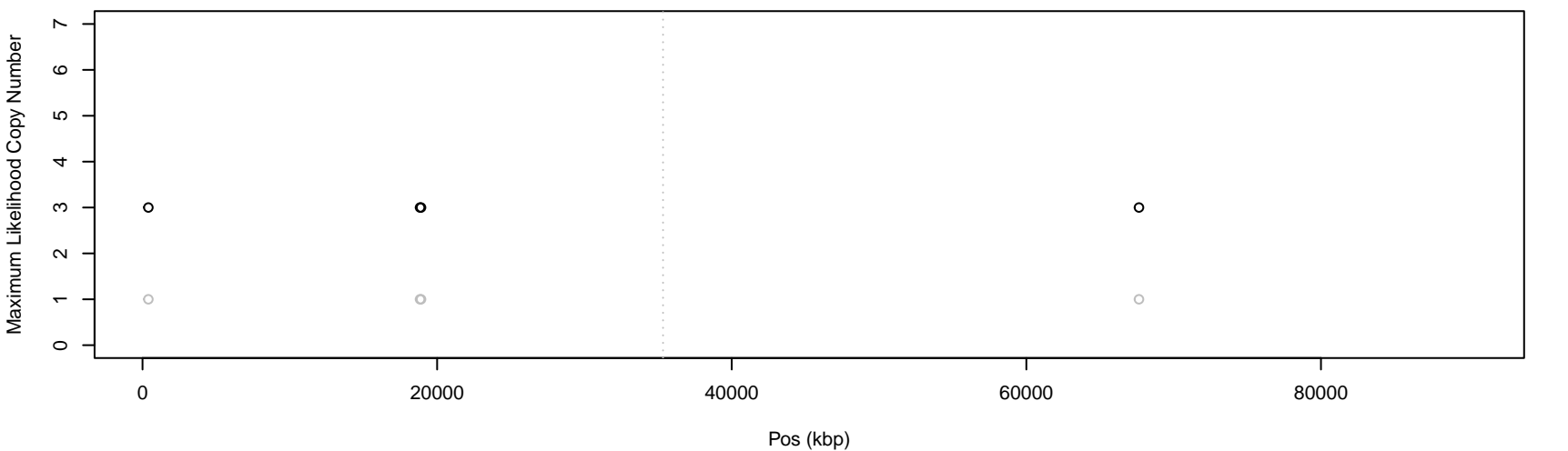
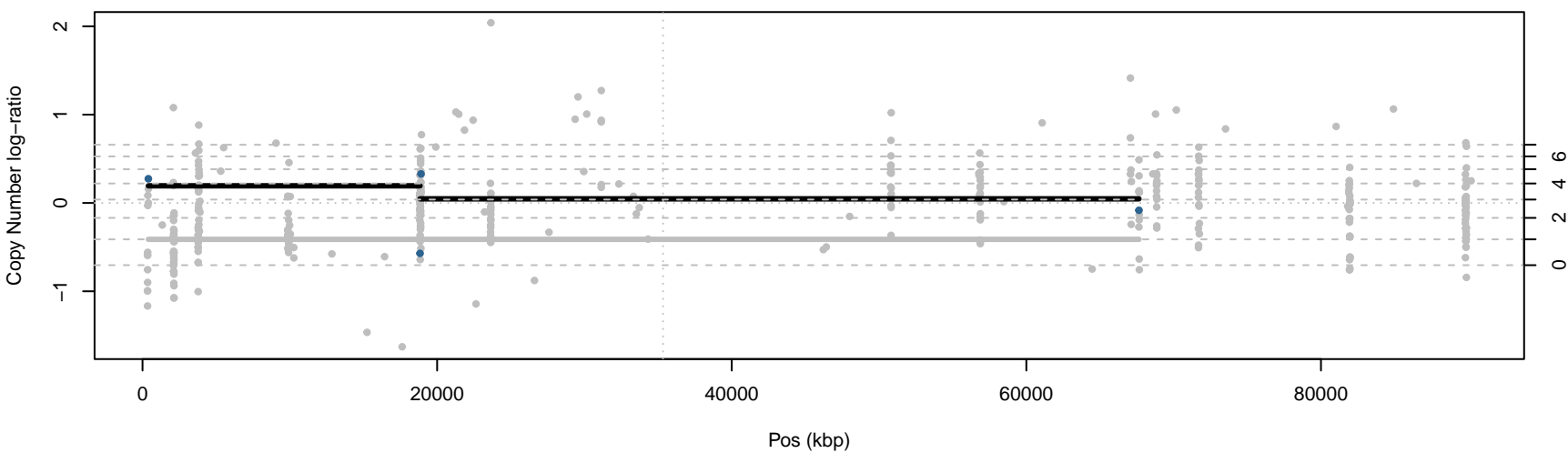
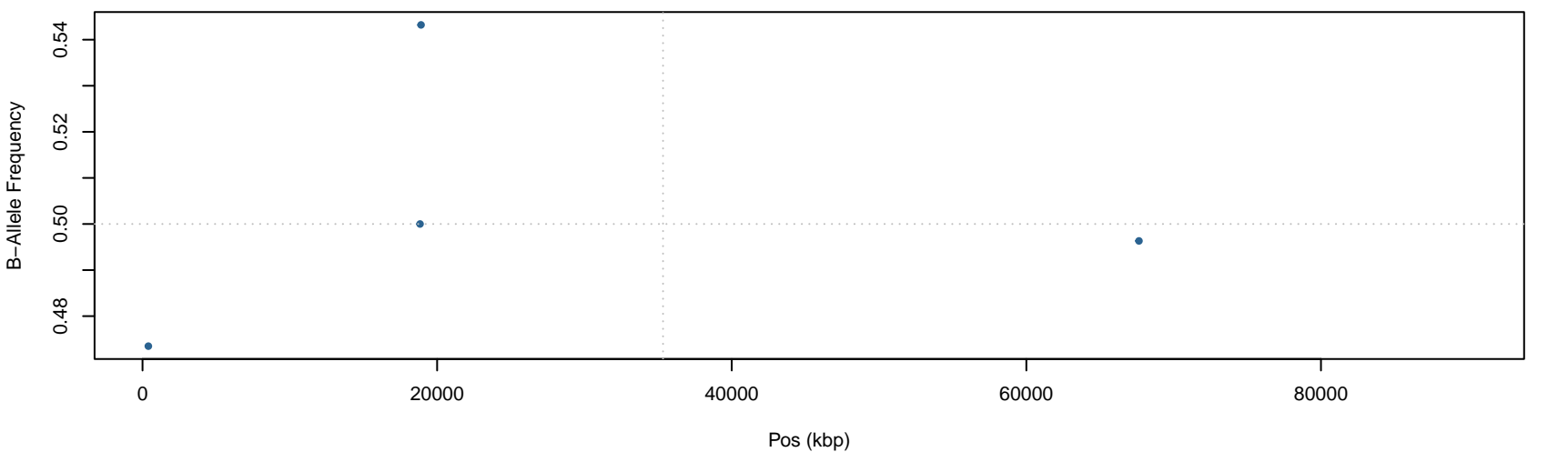
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 14



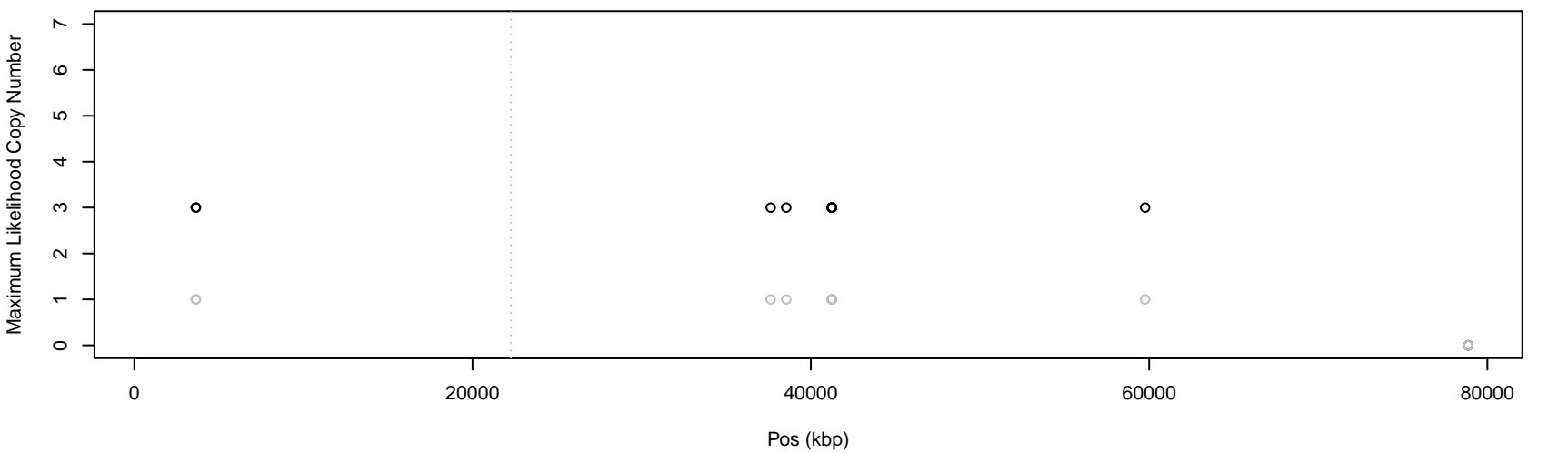
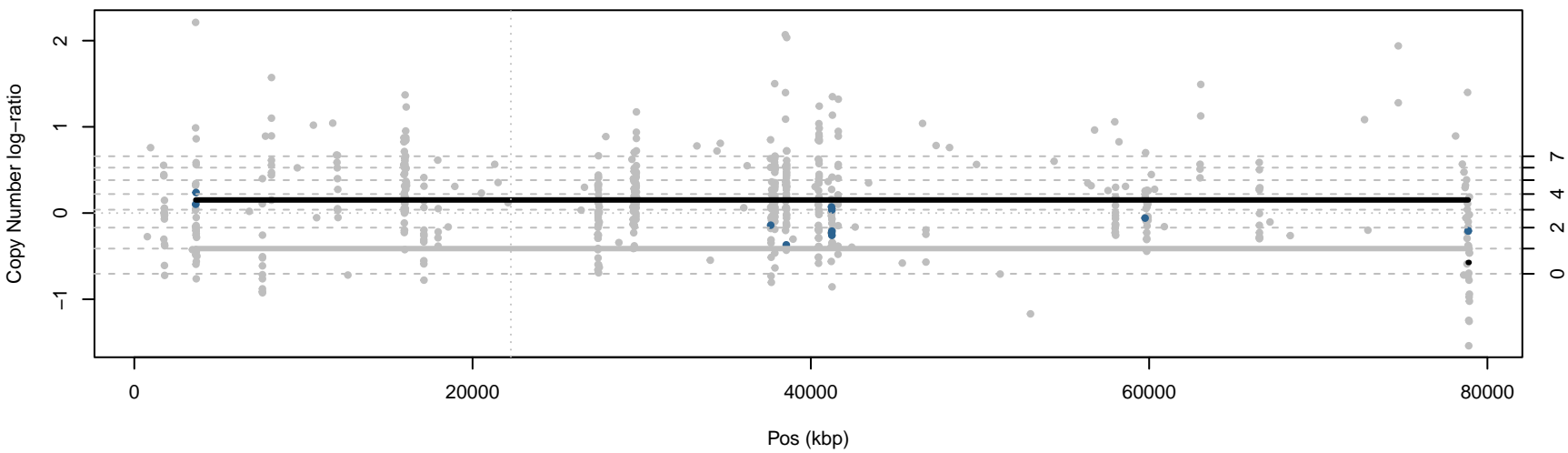
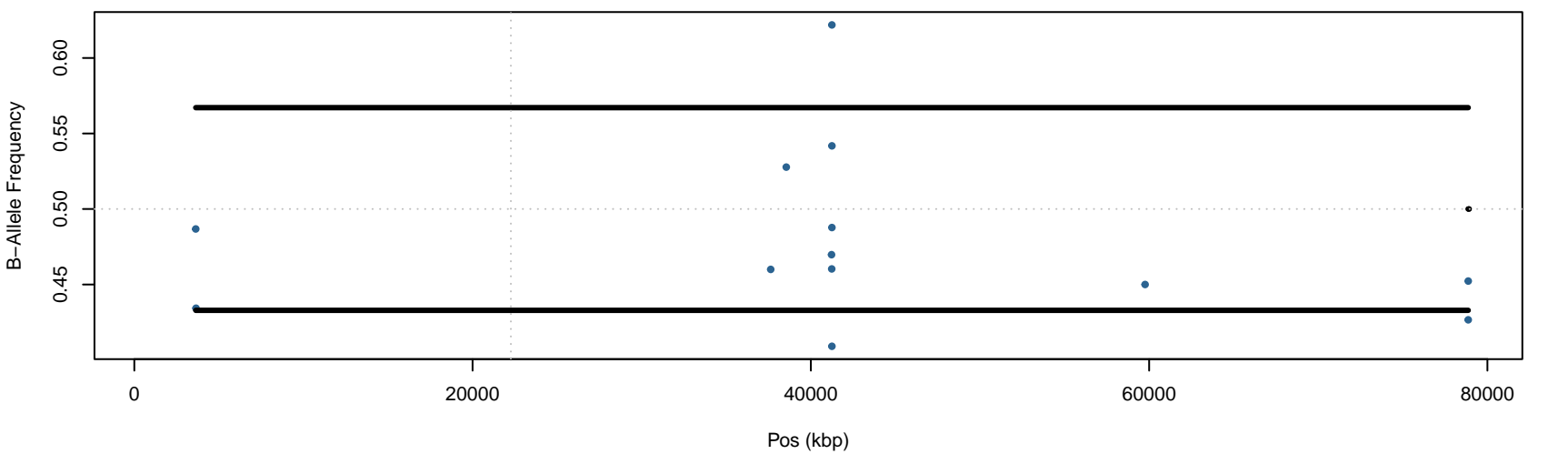
Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 15



Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 16

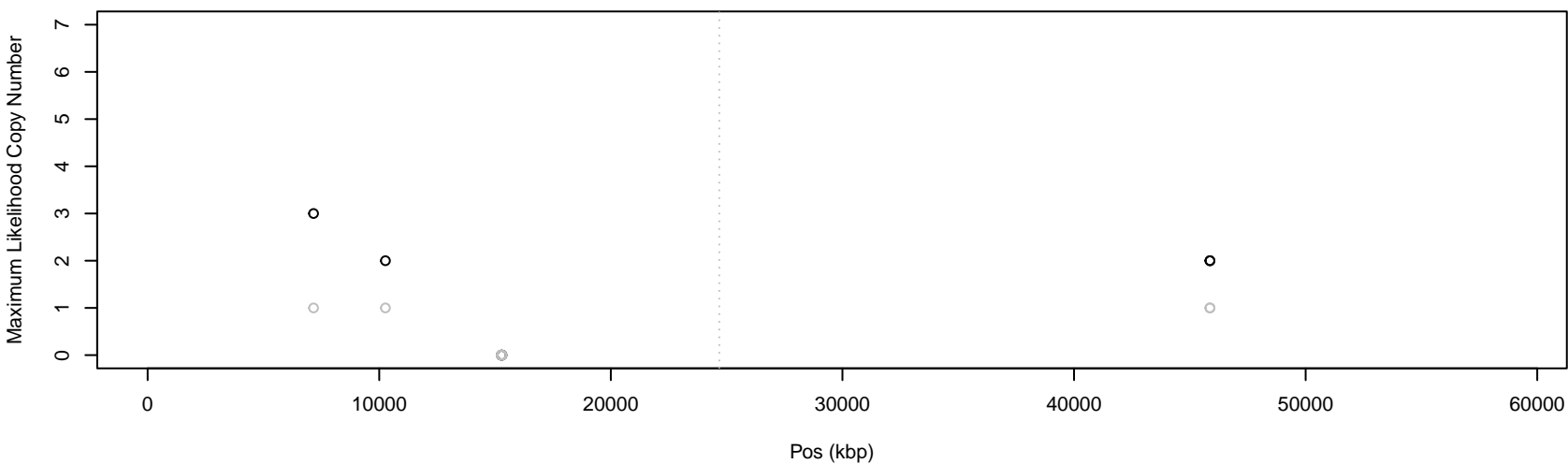
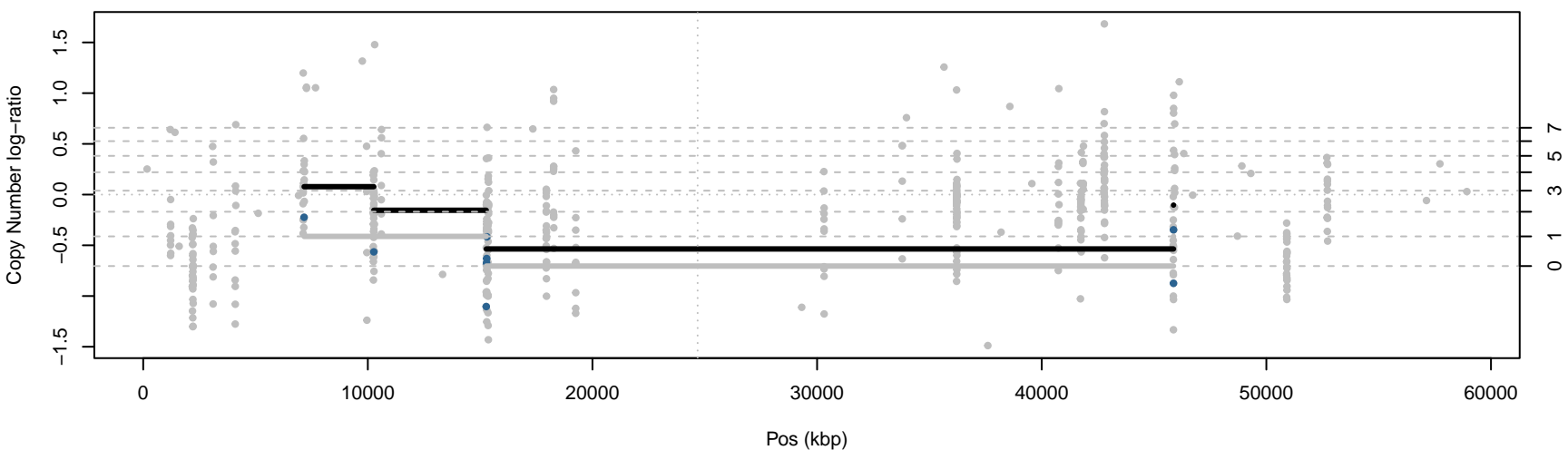
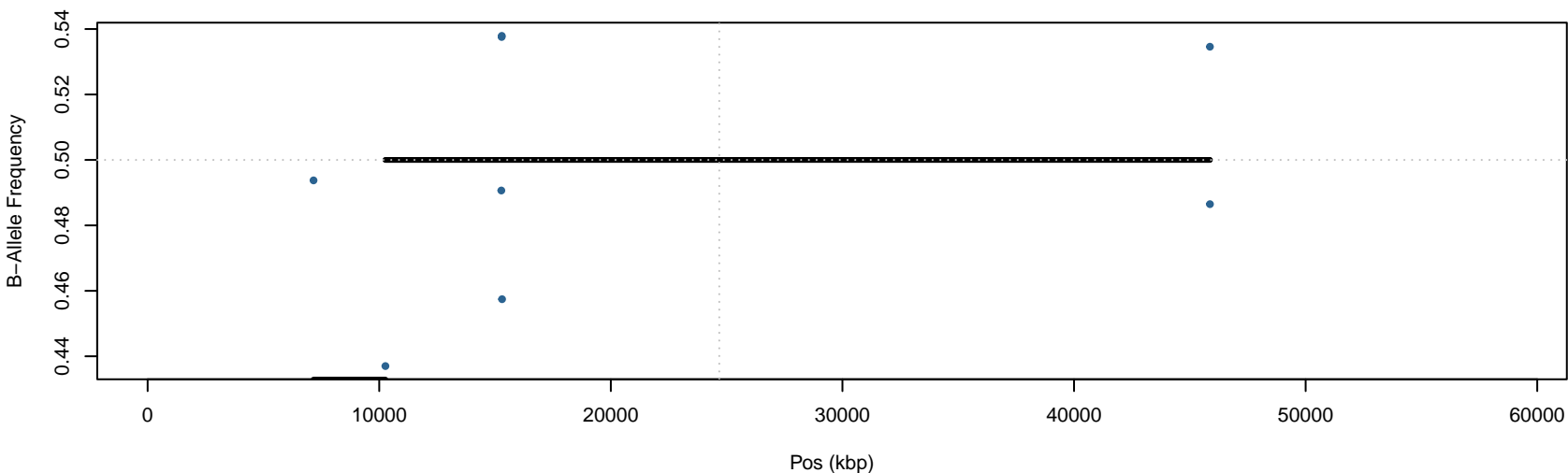


Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 17

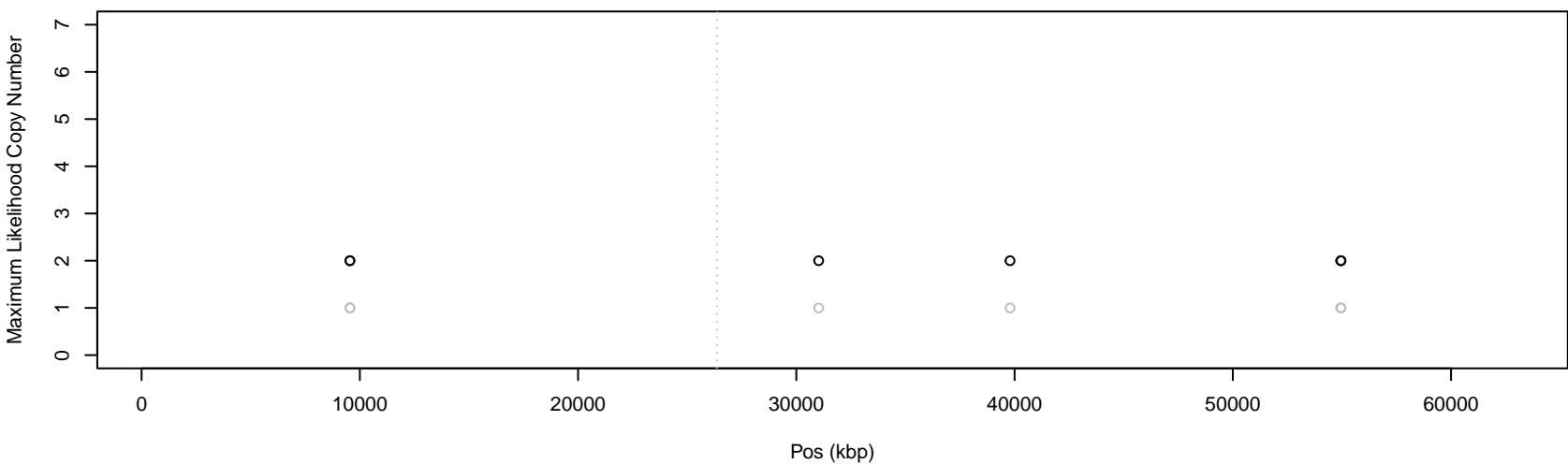
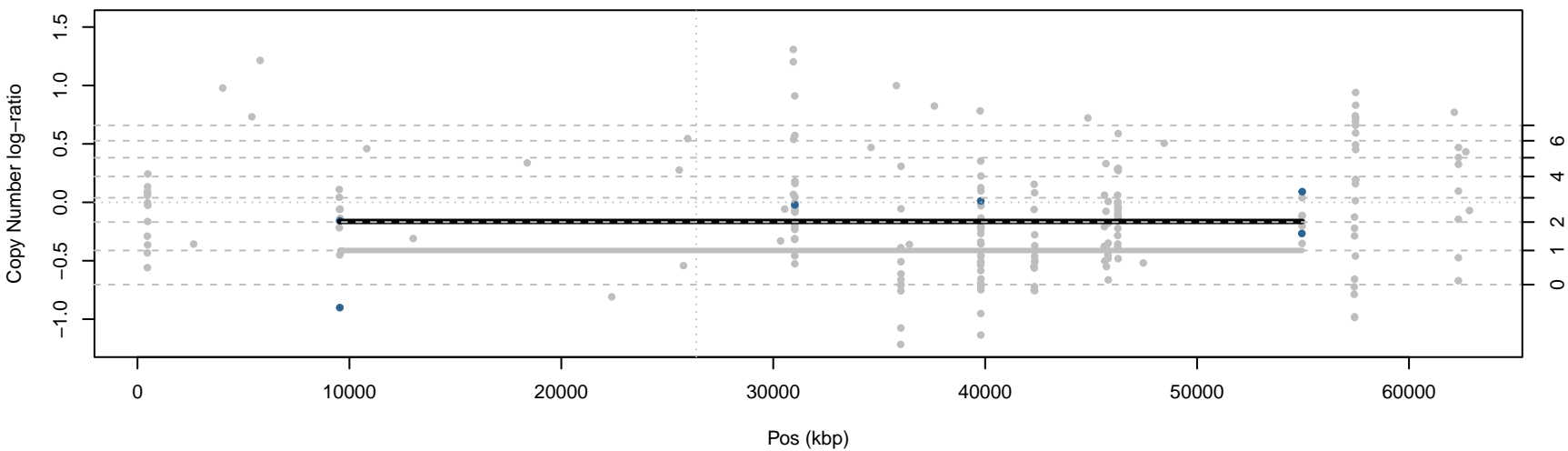
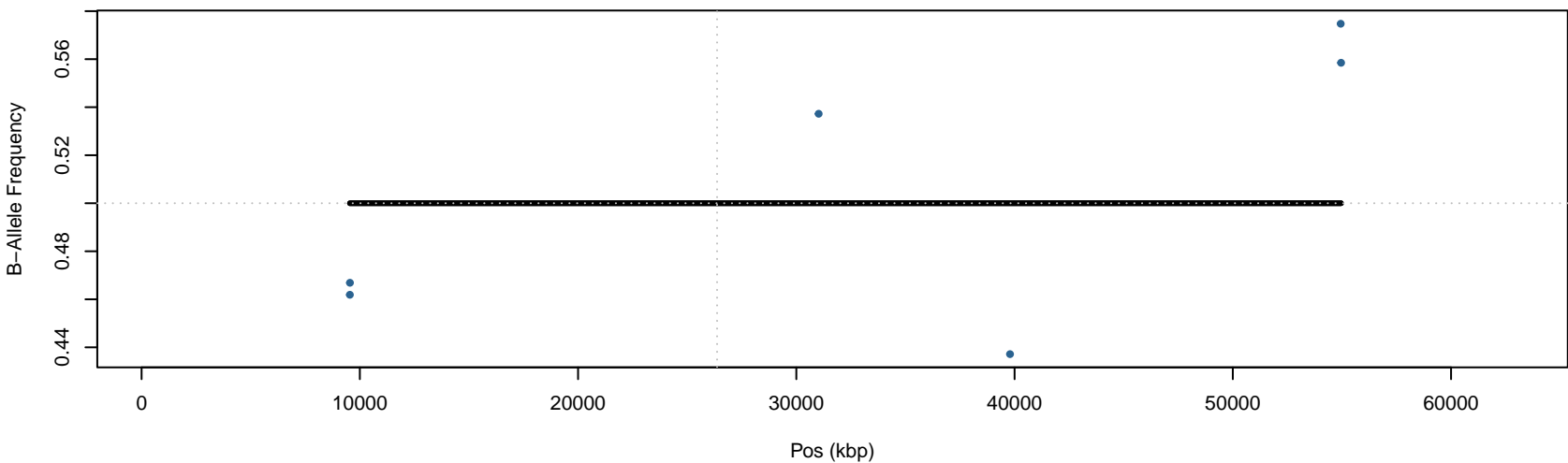




Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 19



Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 20



Purity: 0.31 Tumor ploidy: 2.802 SNV log-likelihood: 10.24 GoF: 93.5% Mean coverage: 367;591 Chromosome: 22

