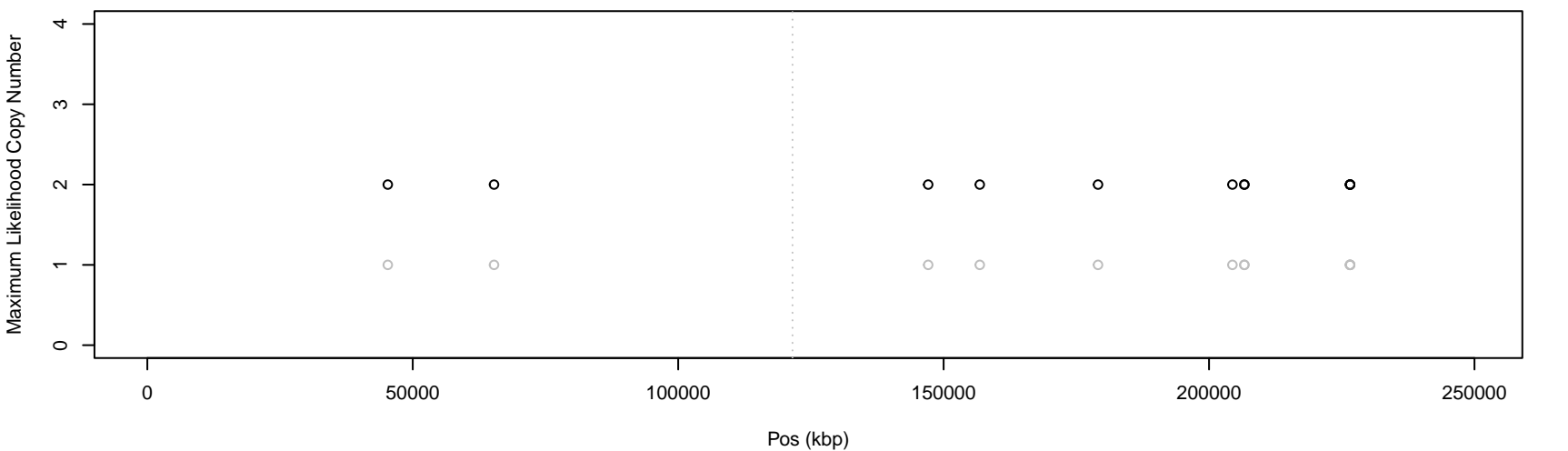
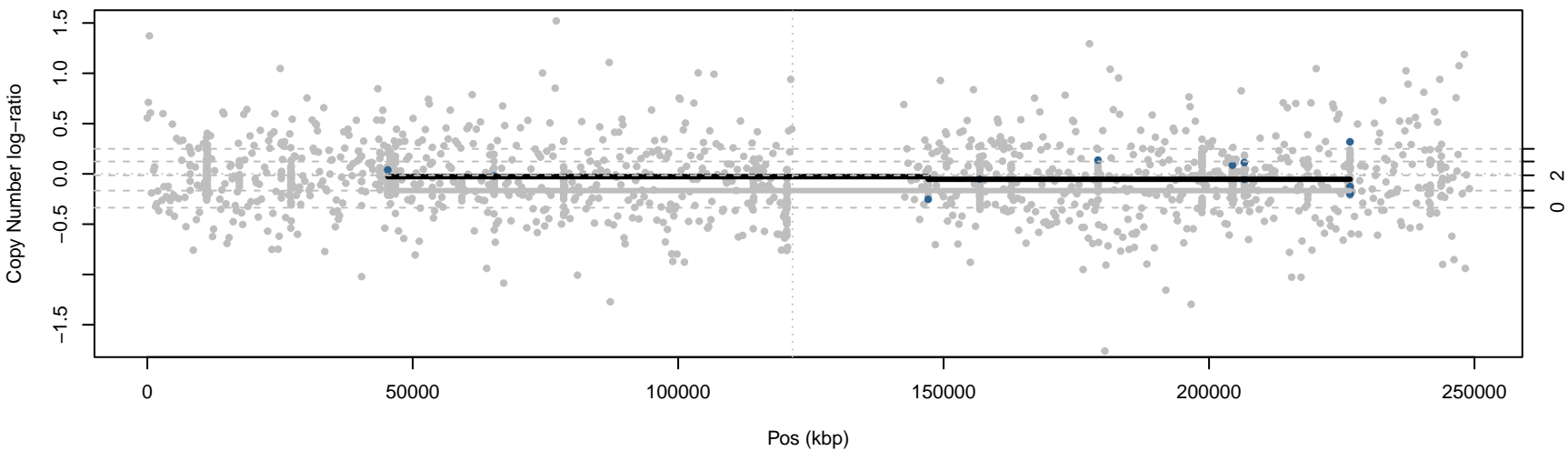
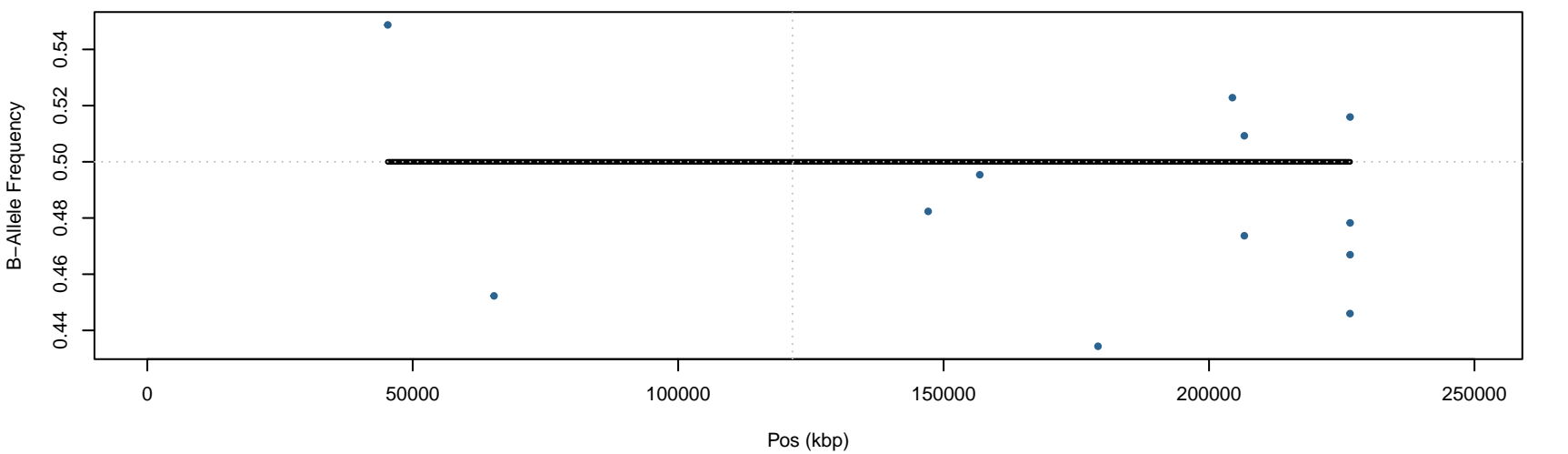
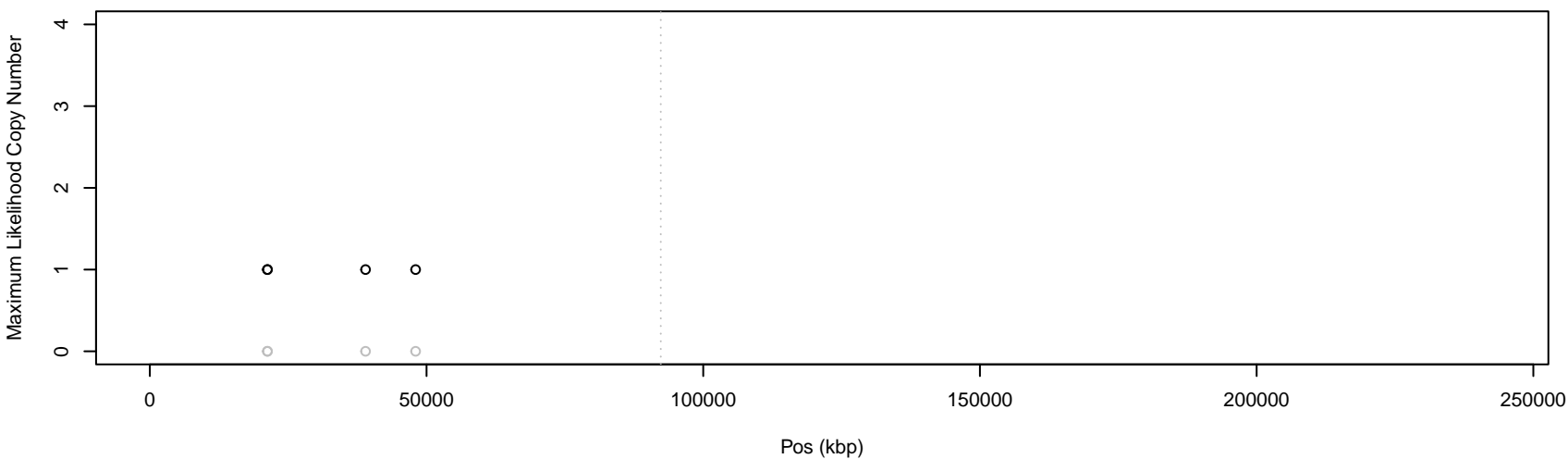
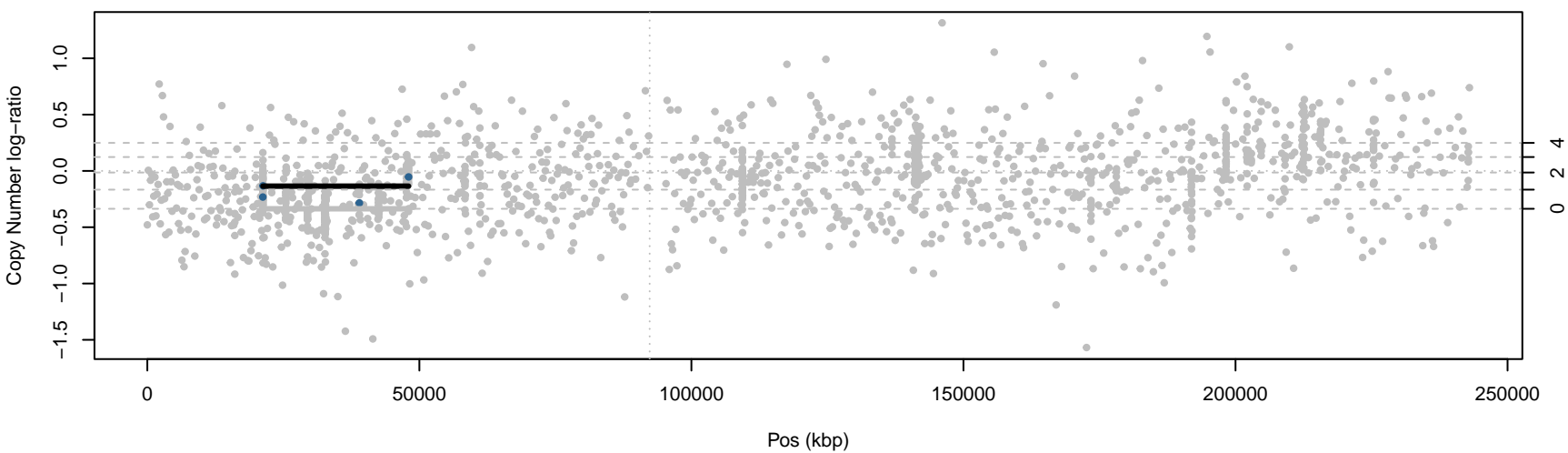
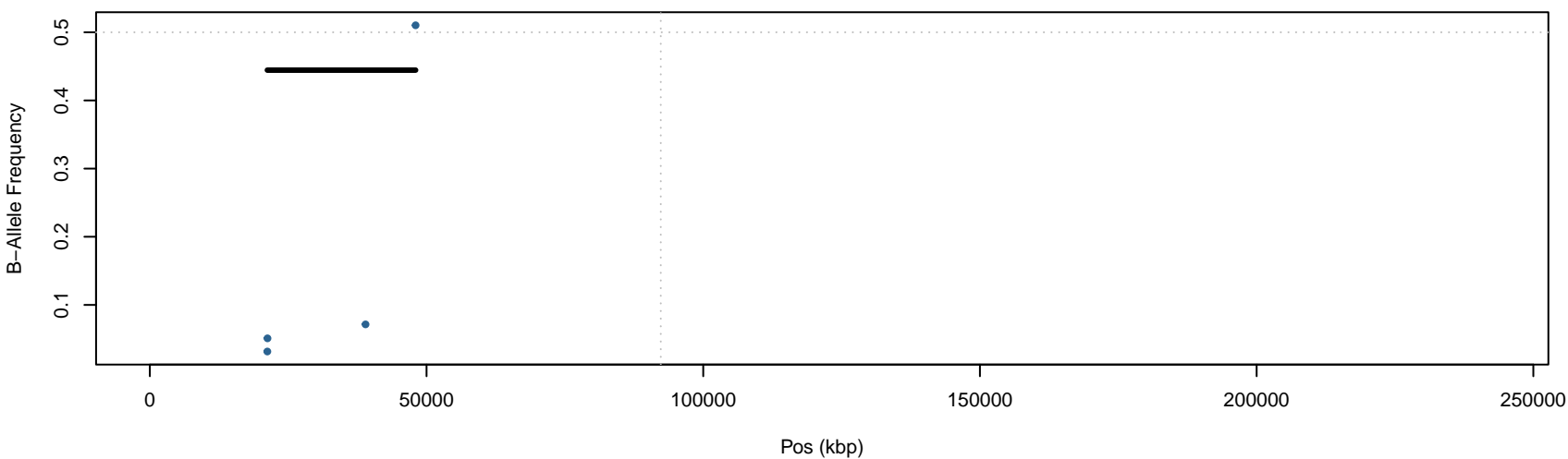


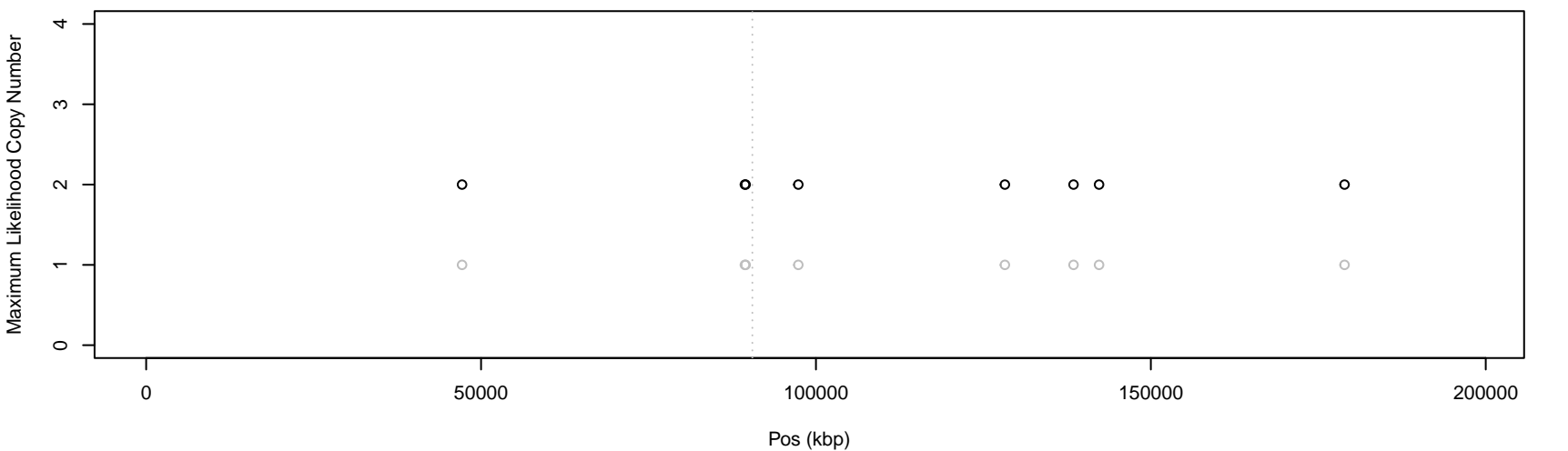
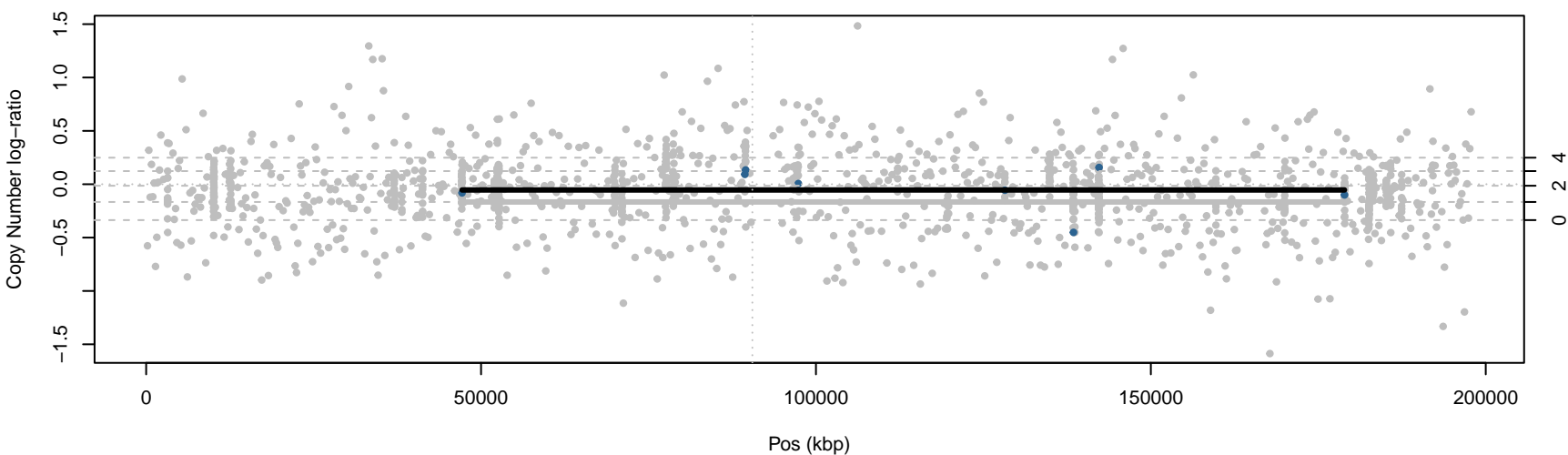
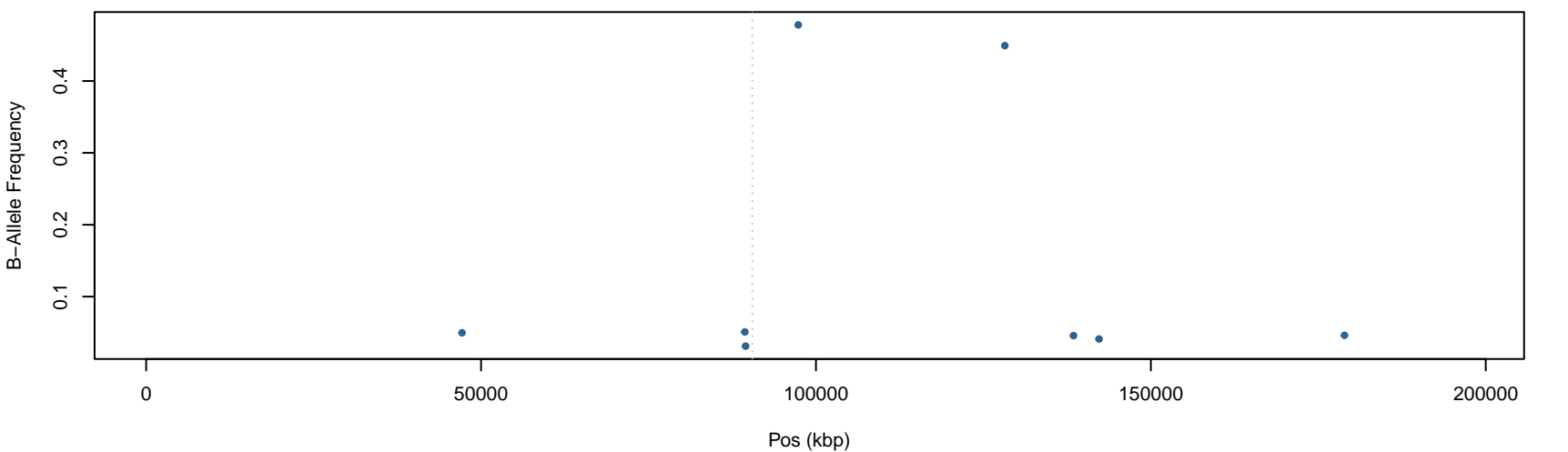
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 1



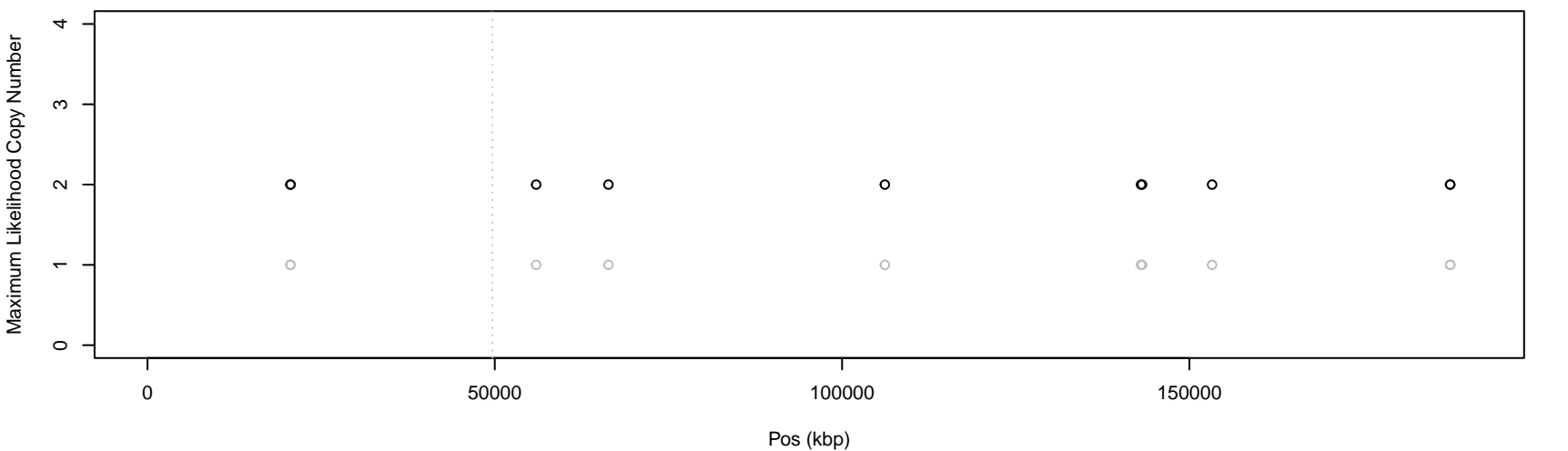
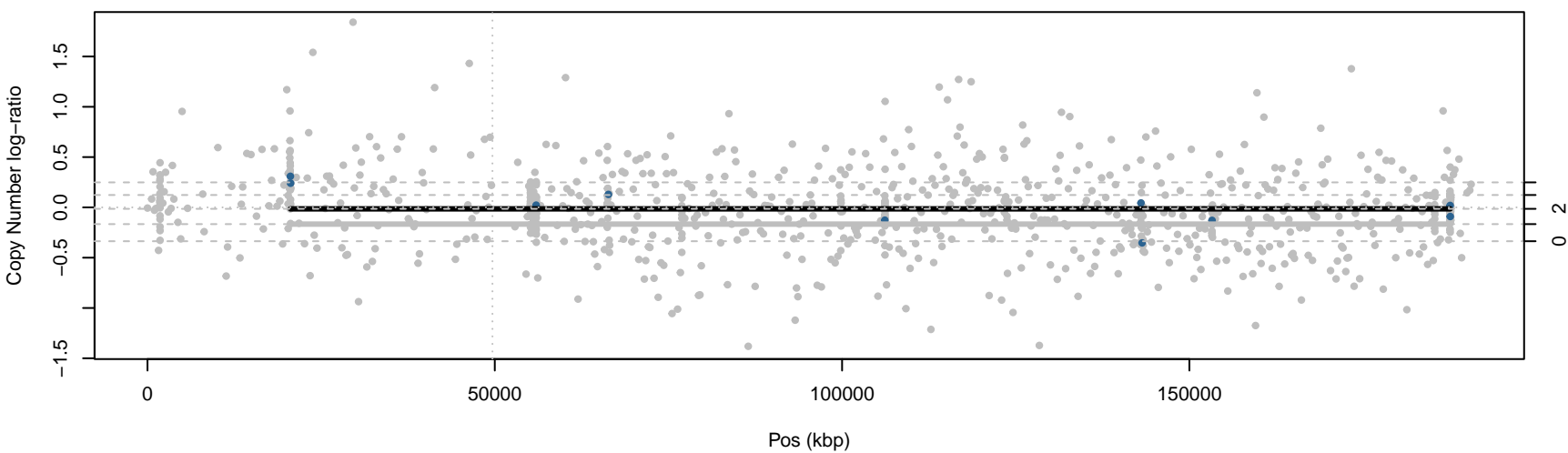
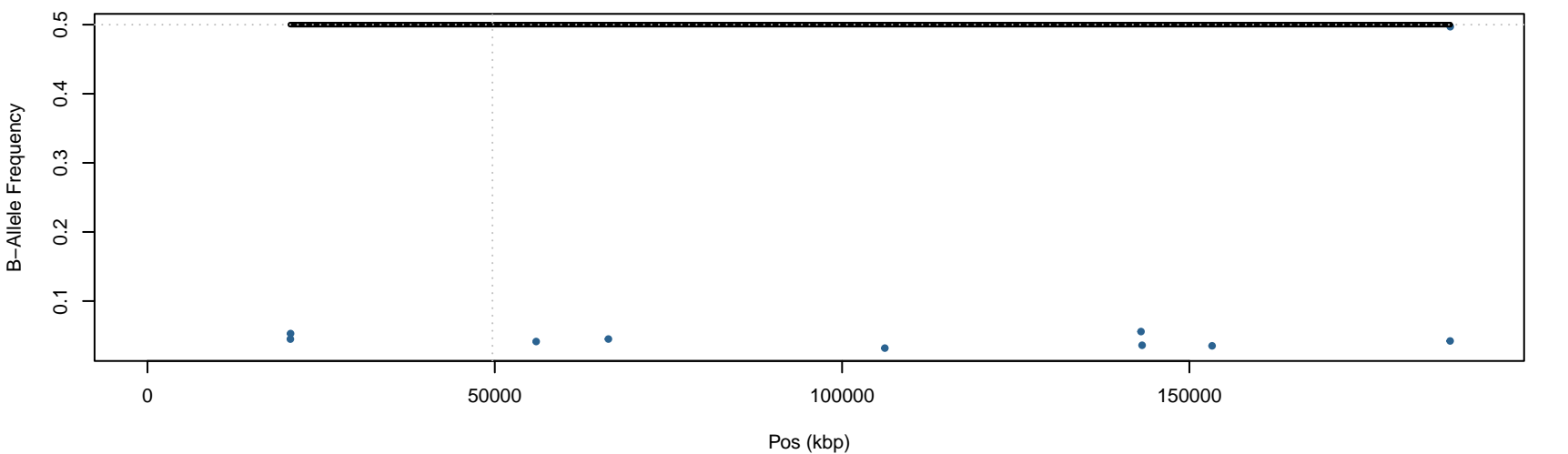
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 2



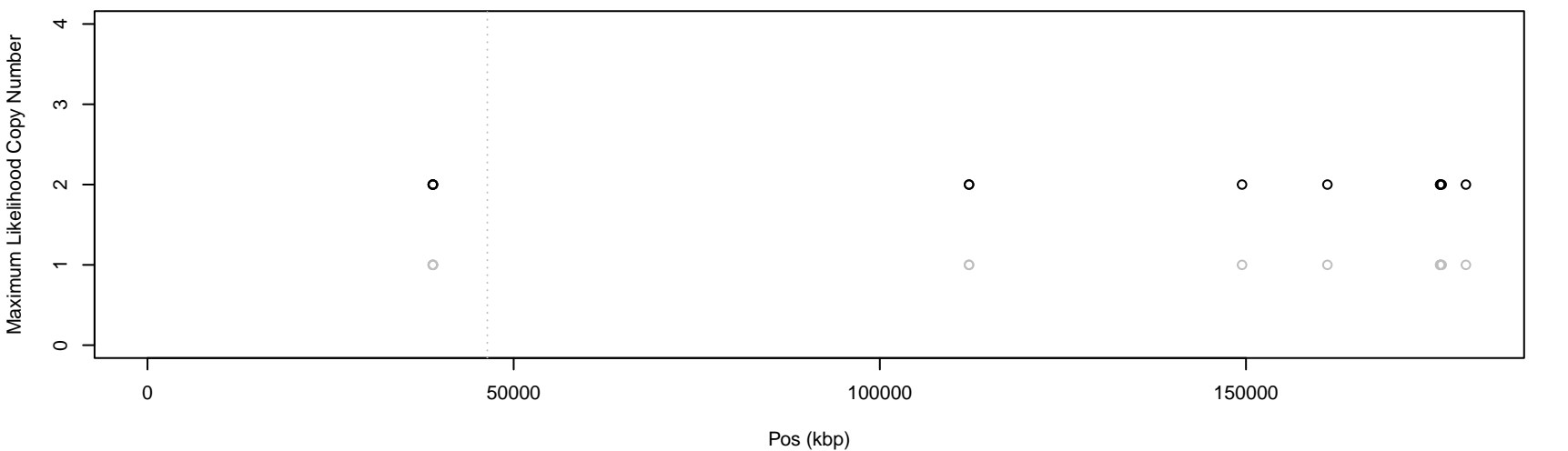
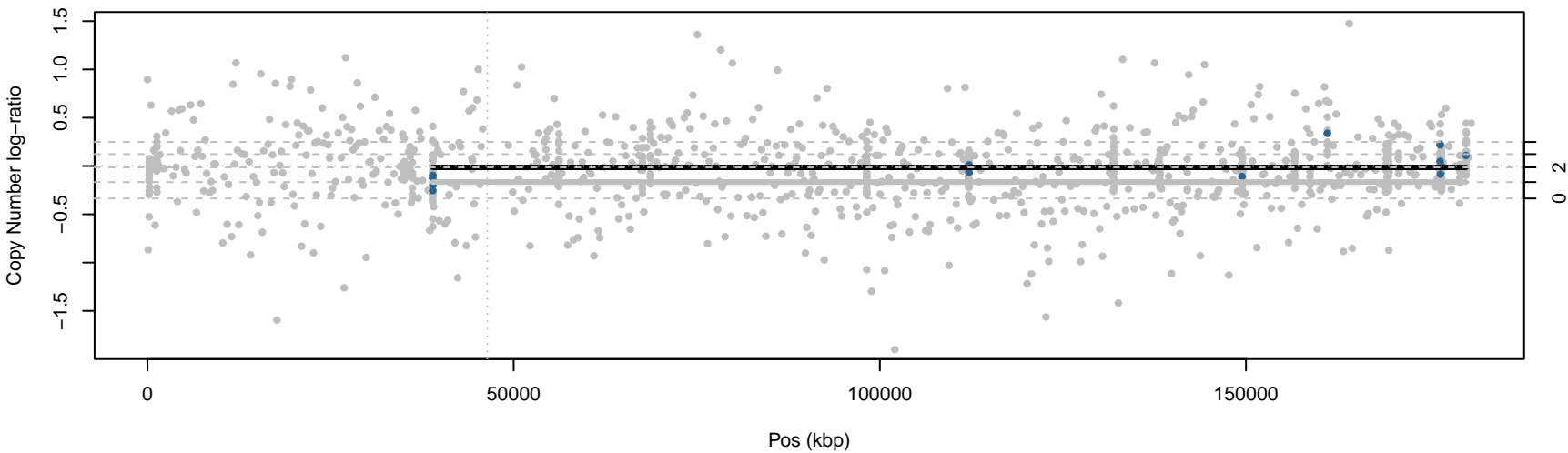
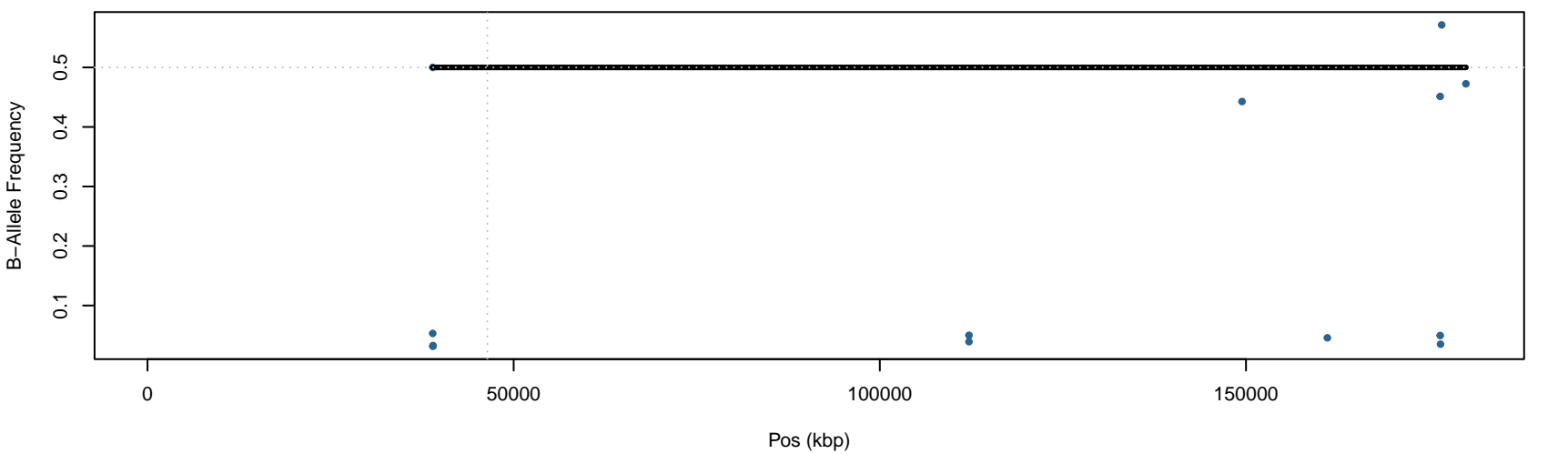
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 3



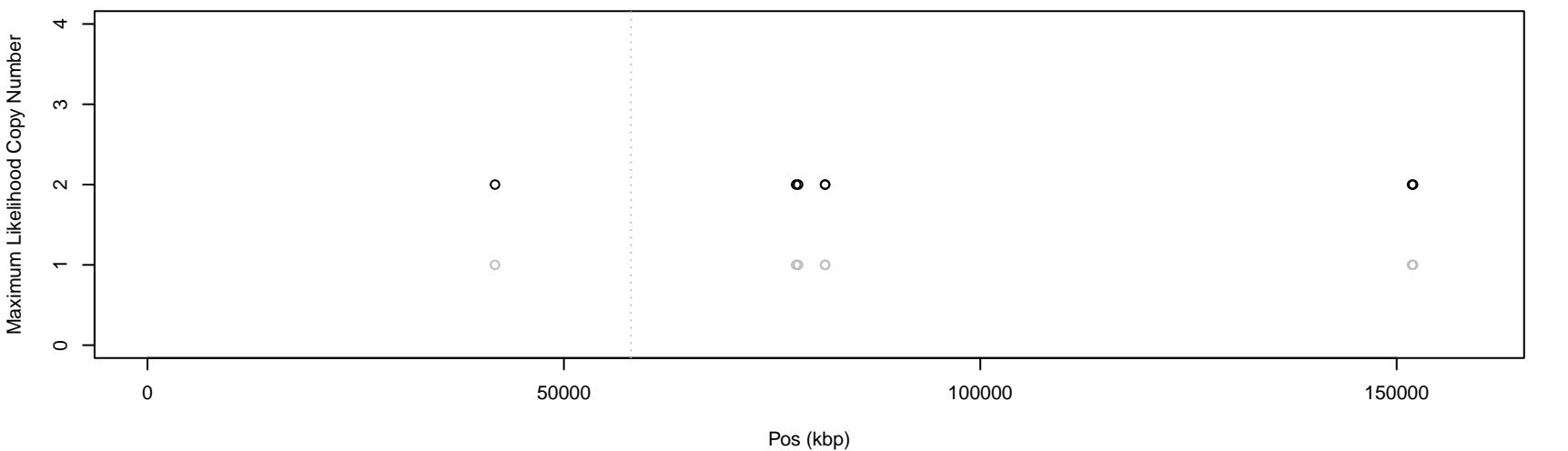
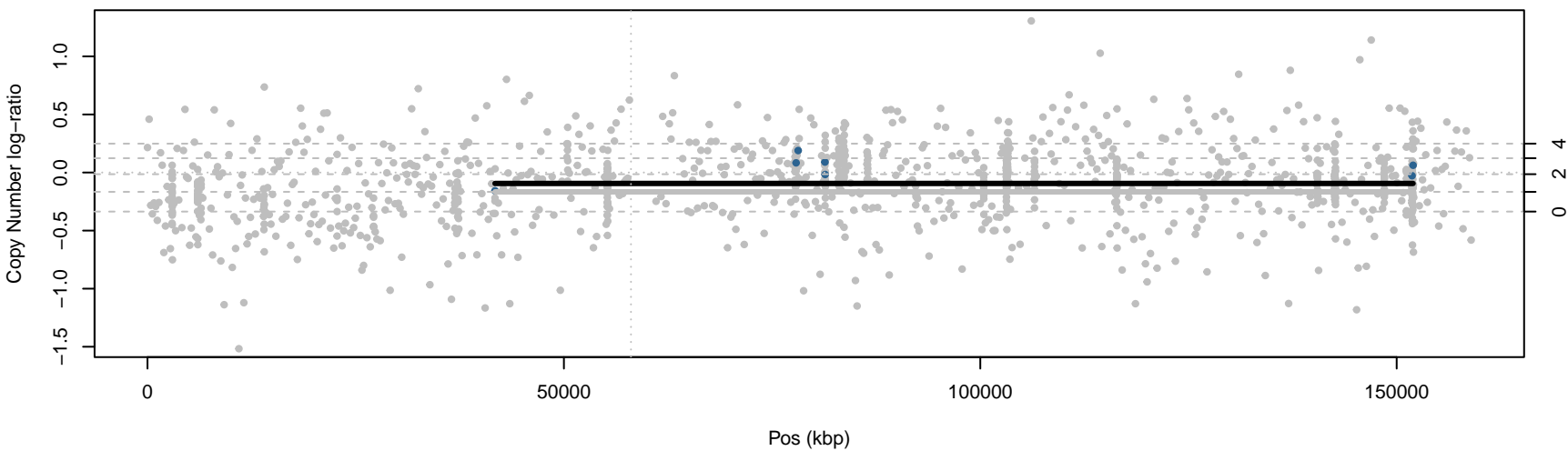
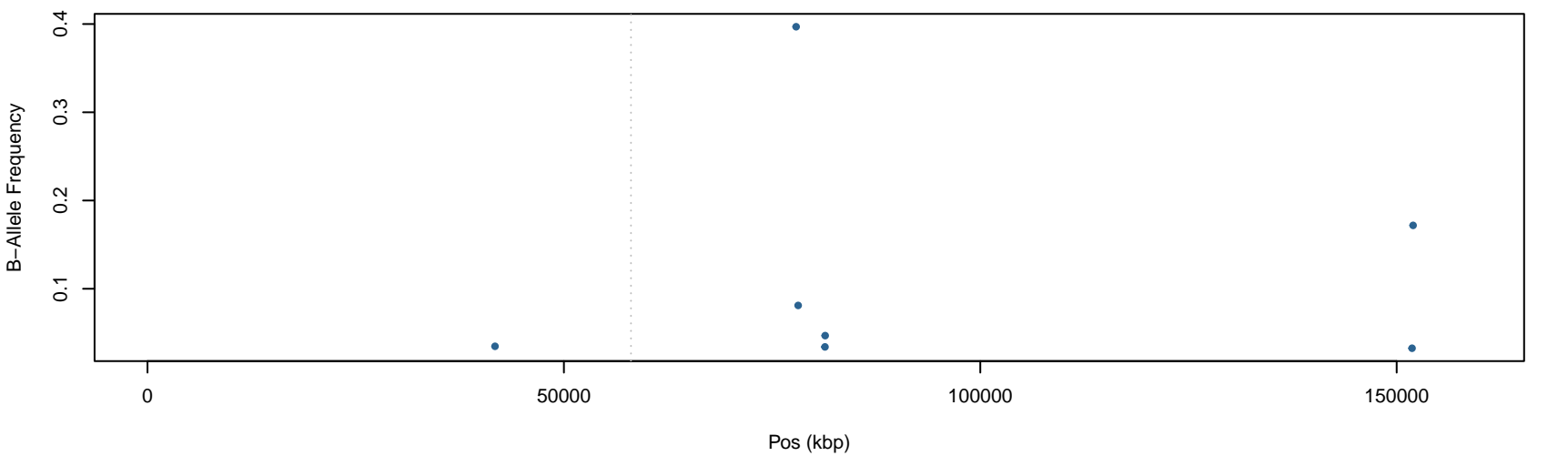
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 4



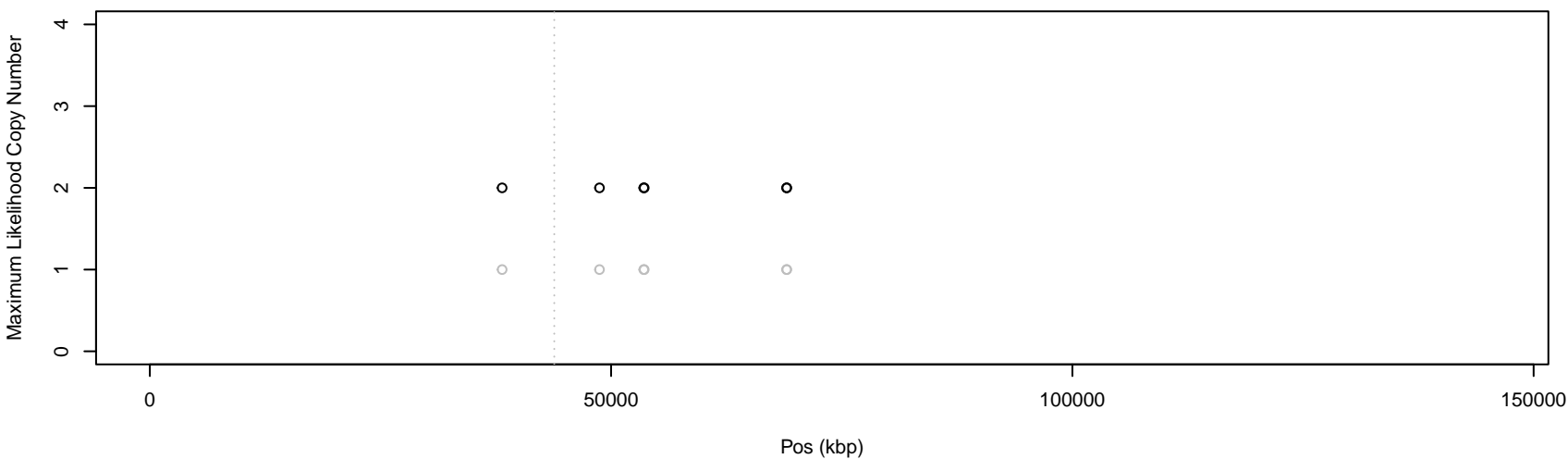
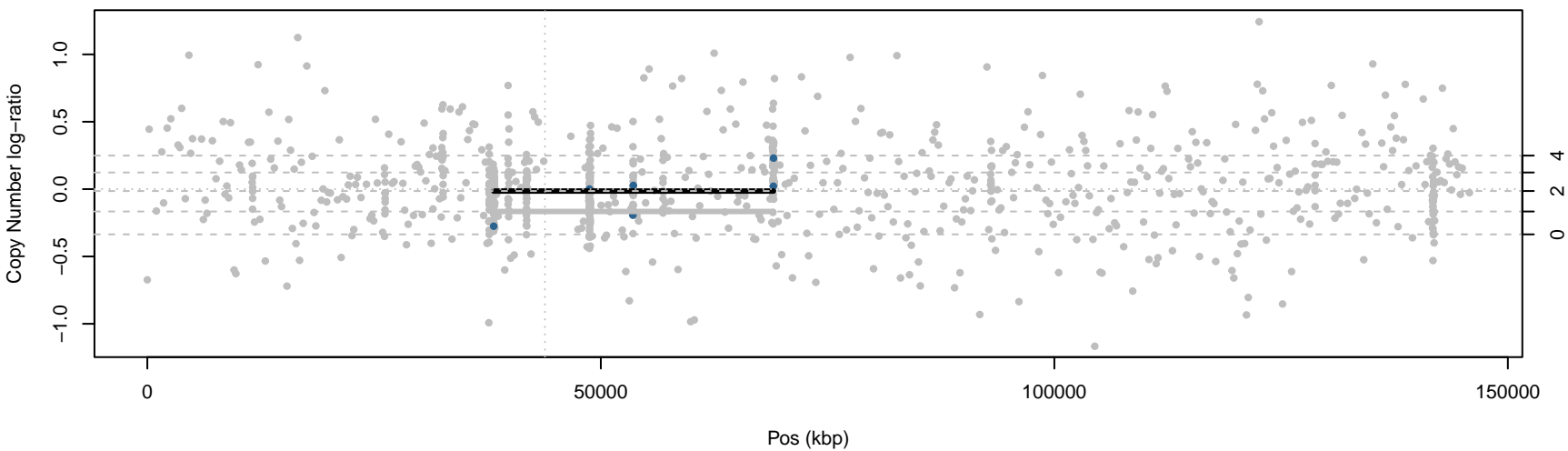
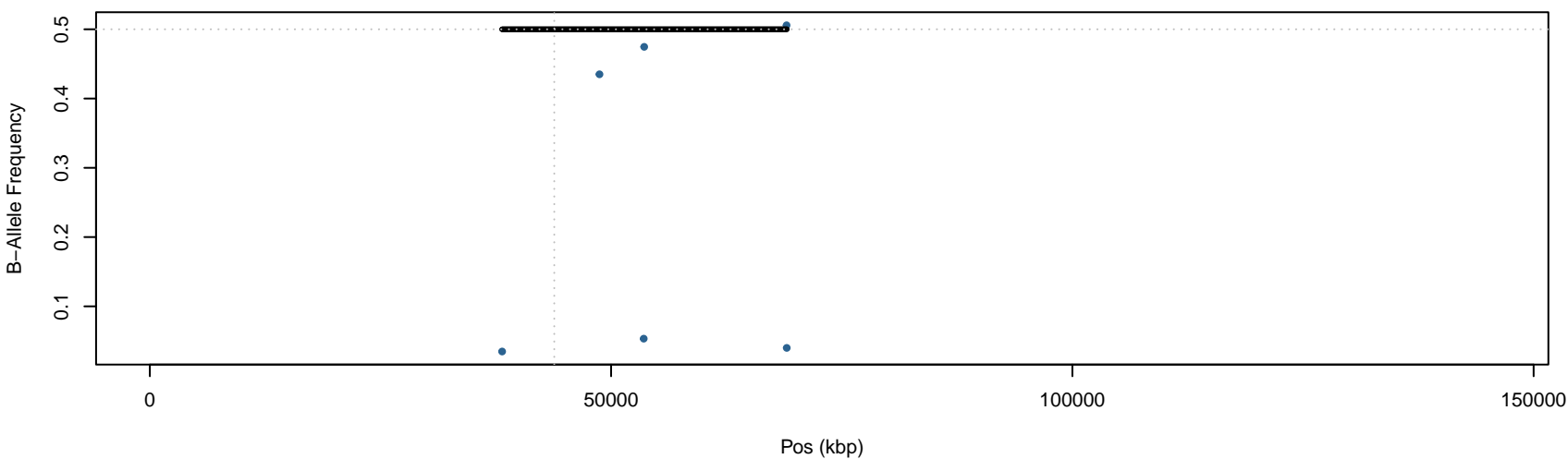
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 5



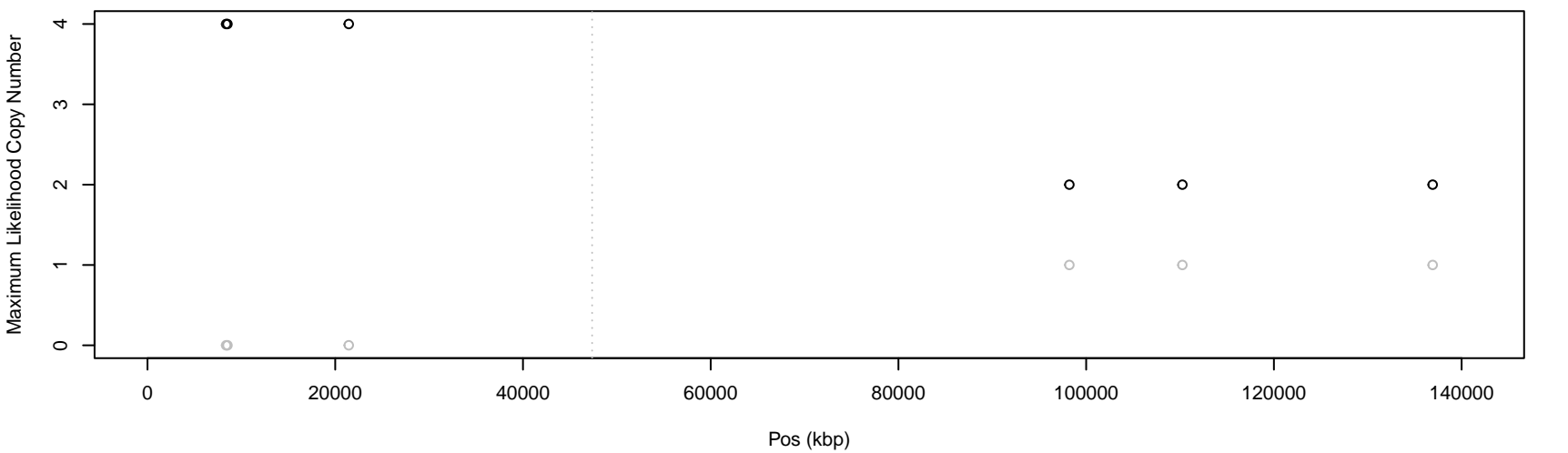
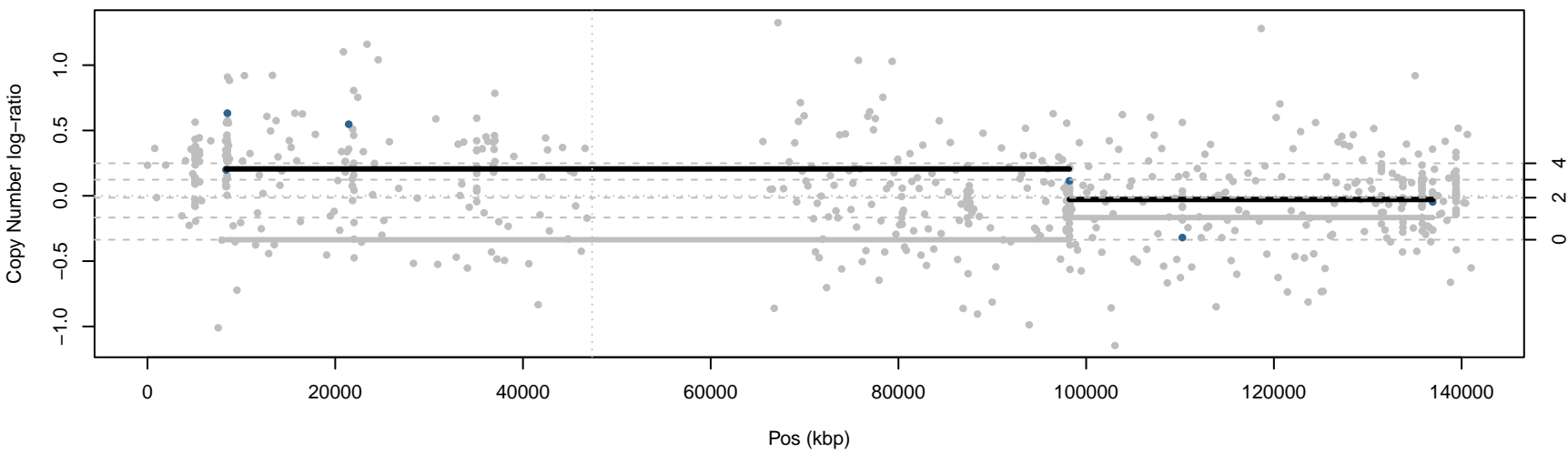
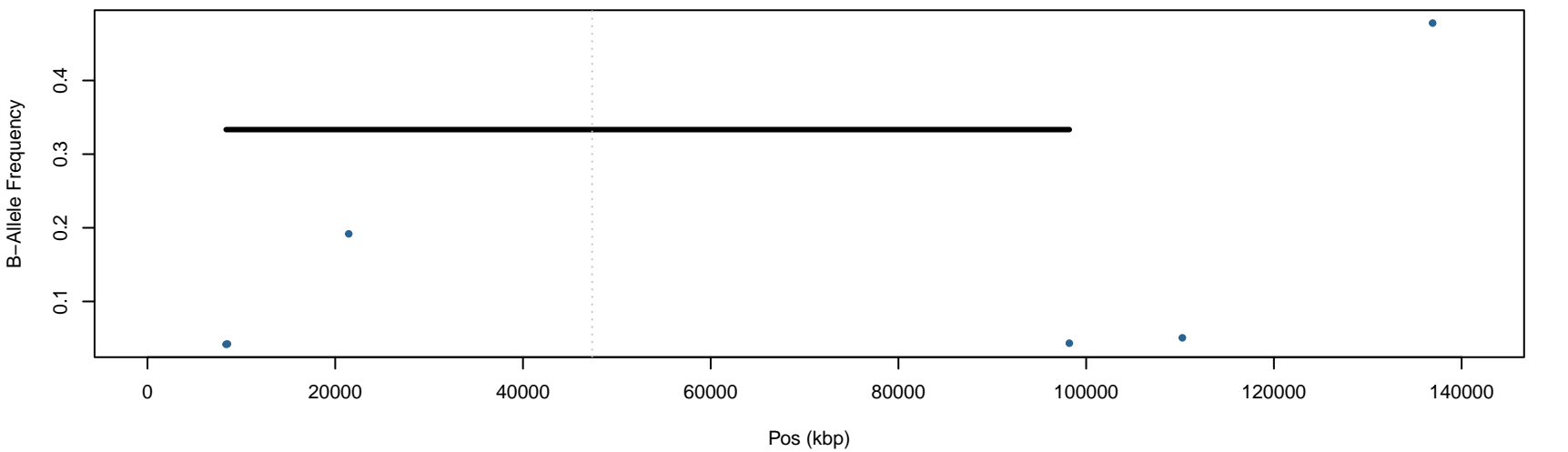
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 7



Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 8

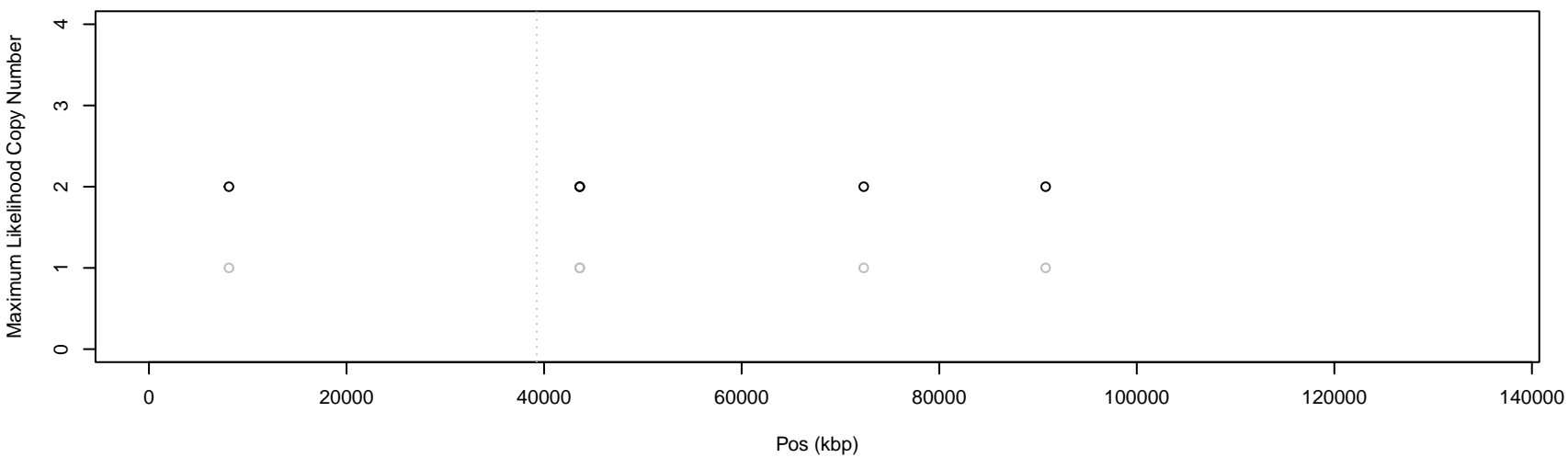
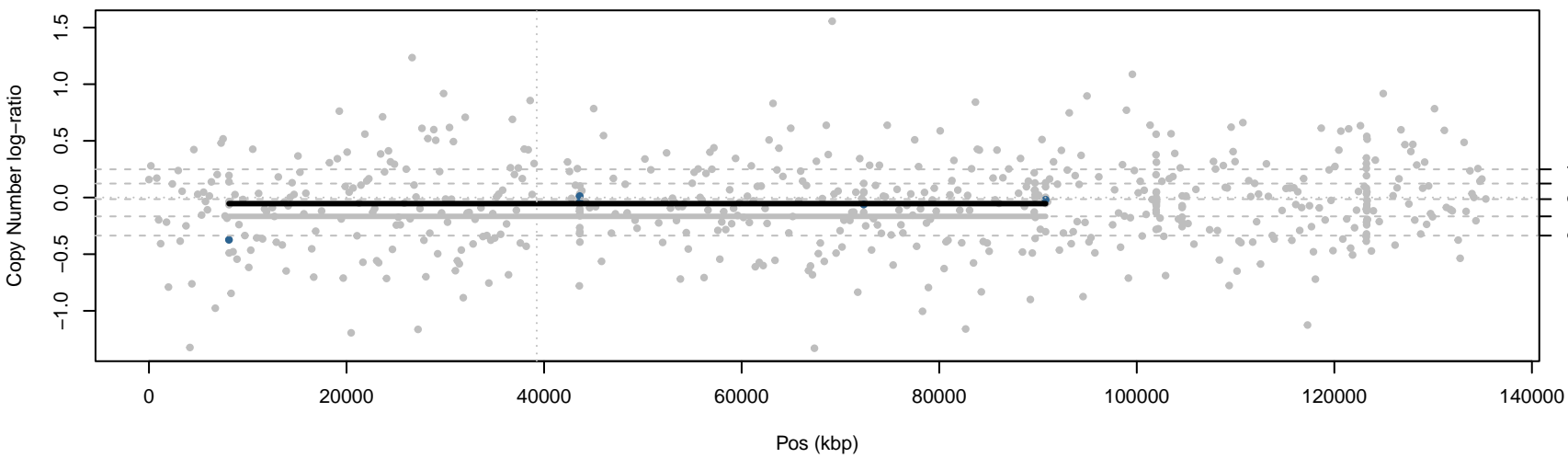
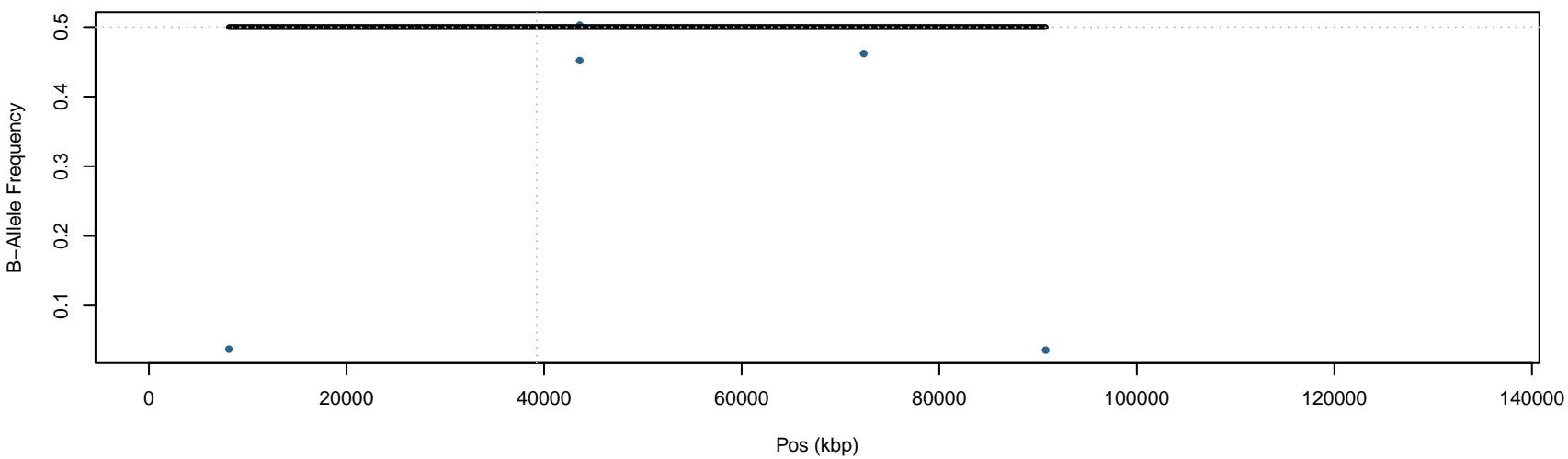


Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 9

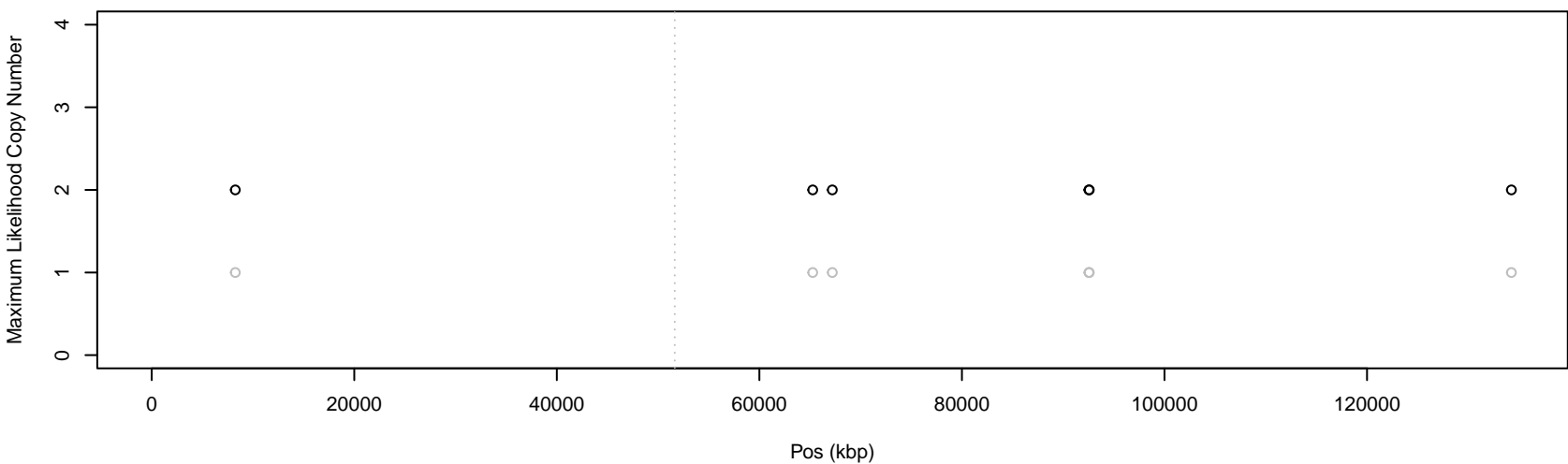
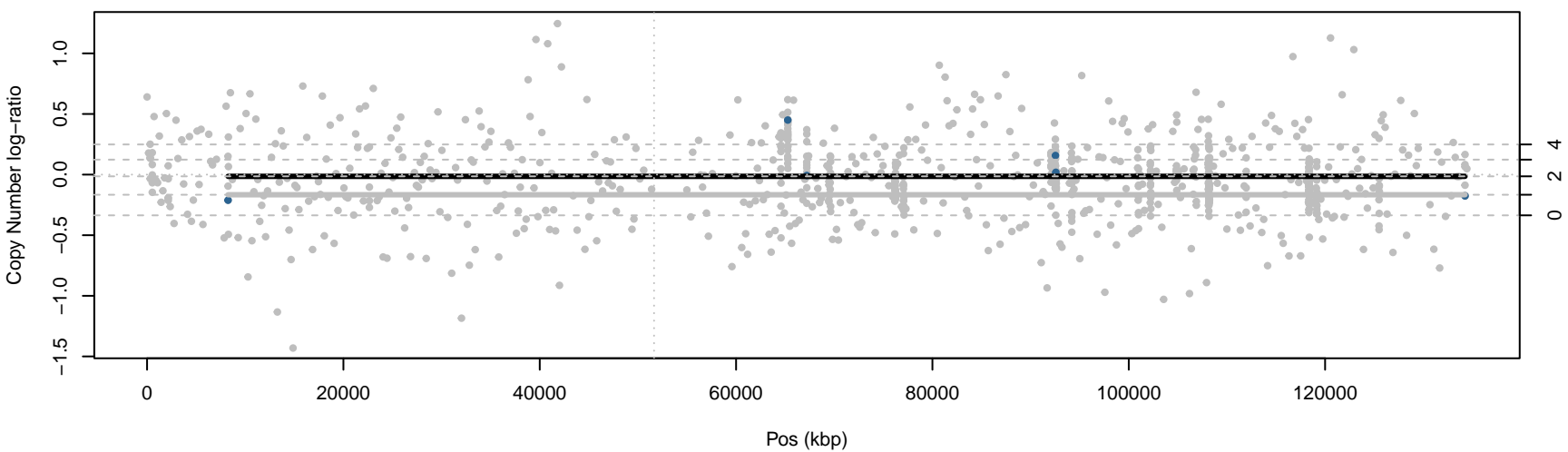
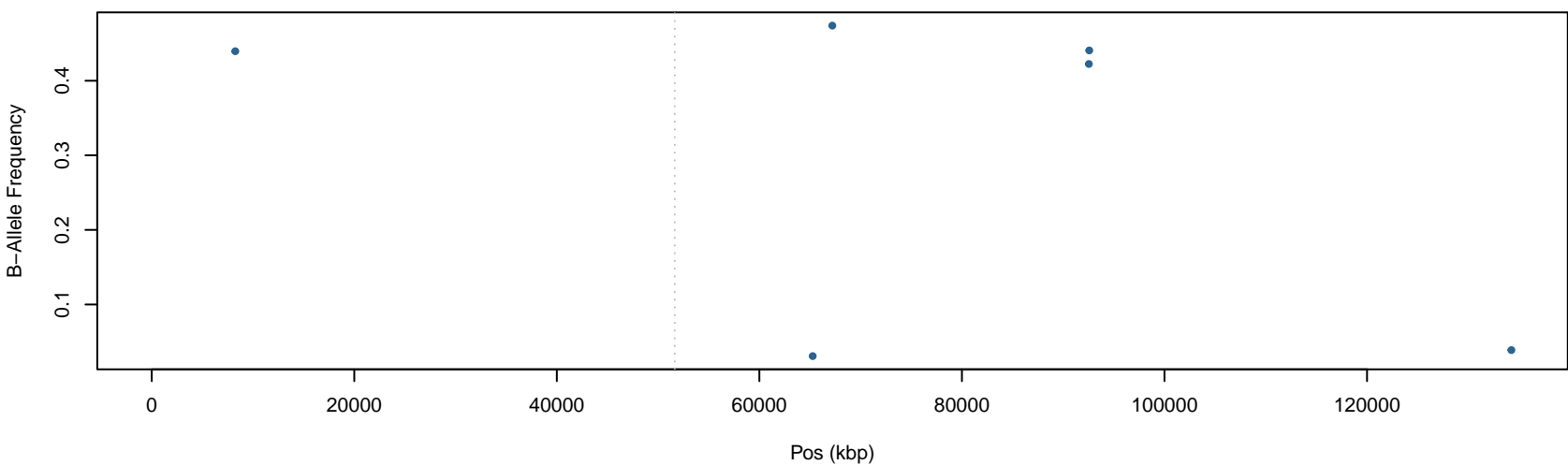




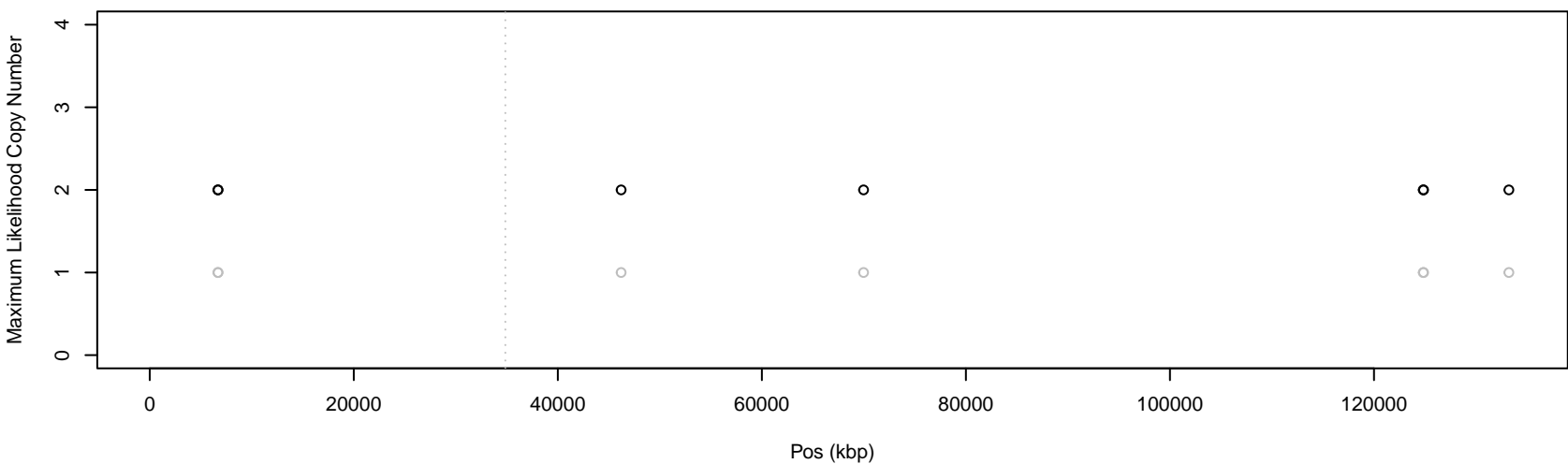
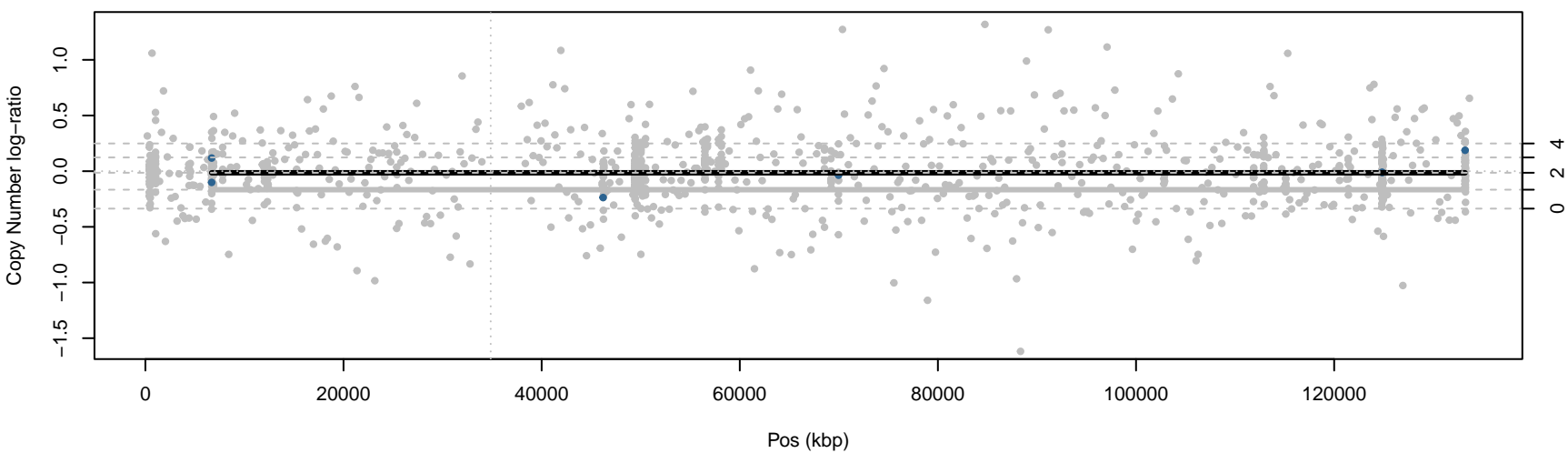
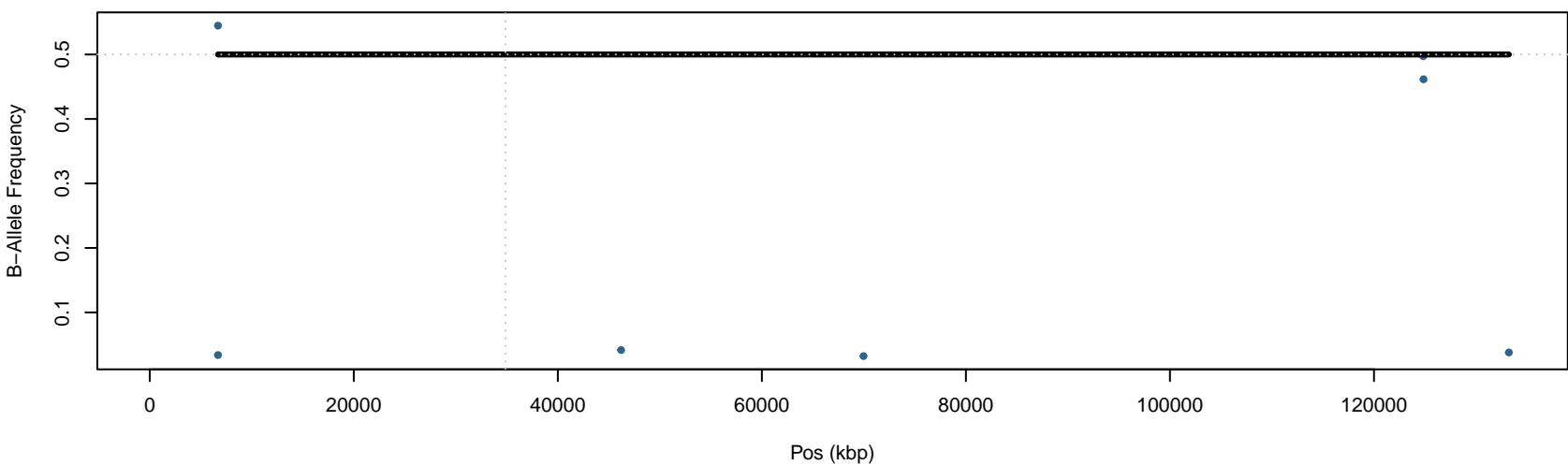
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 10



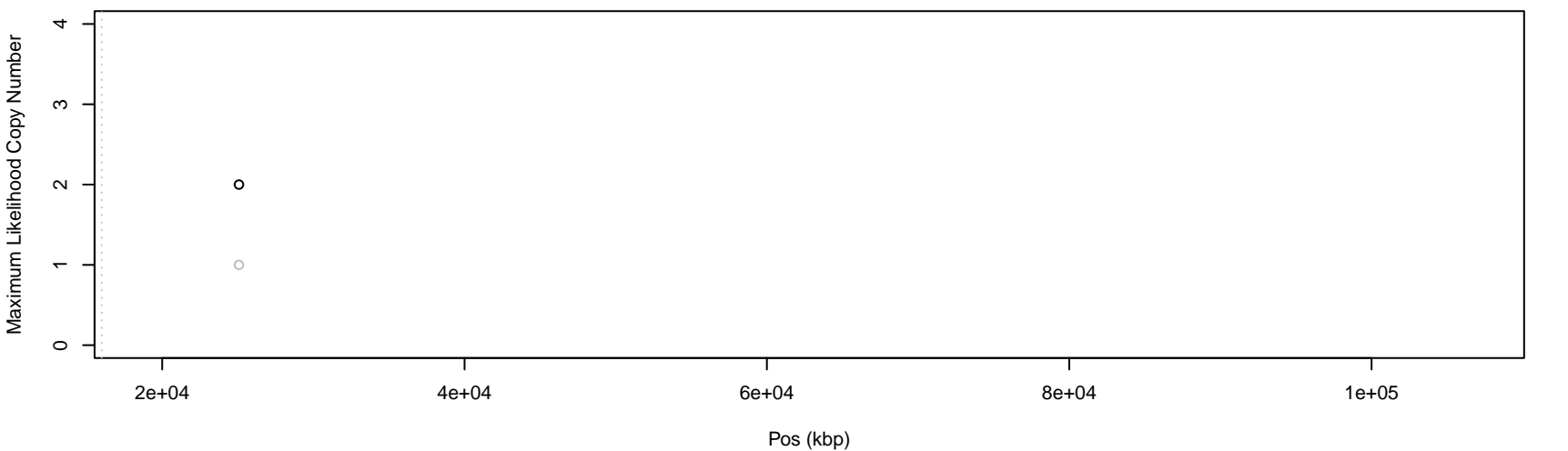
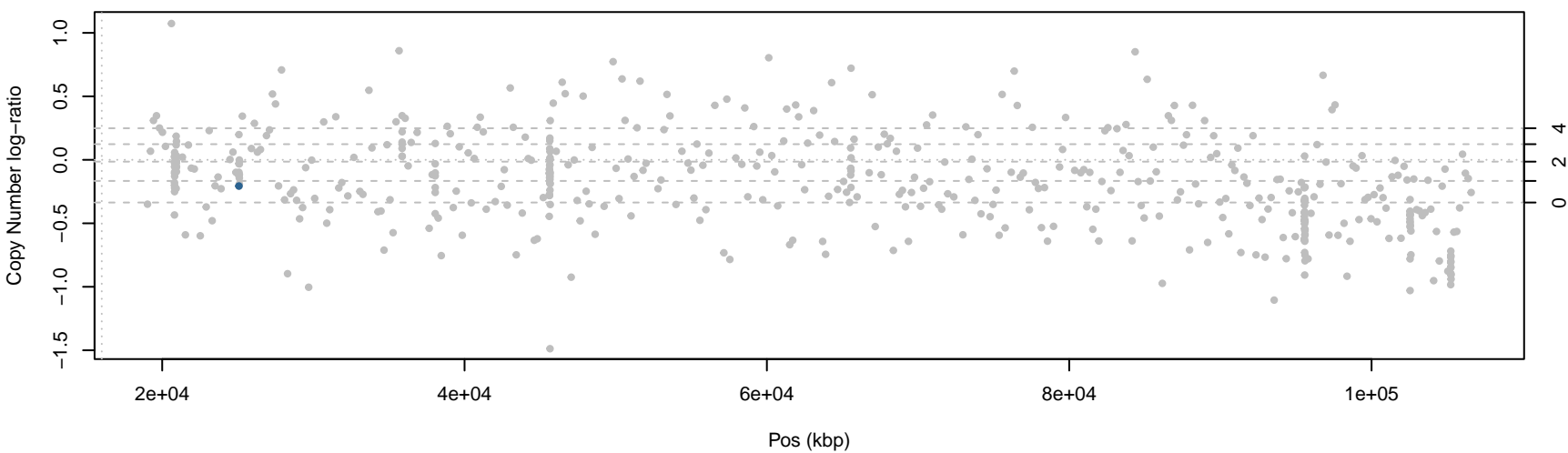
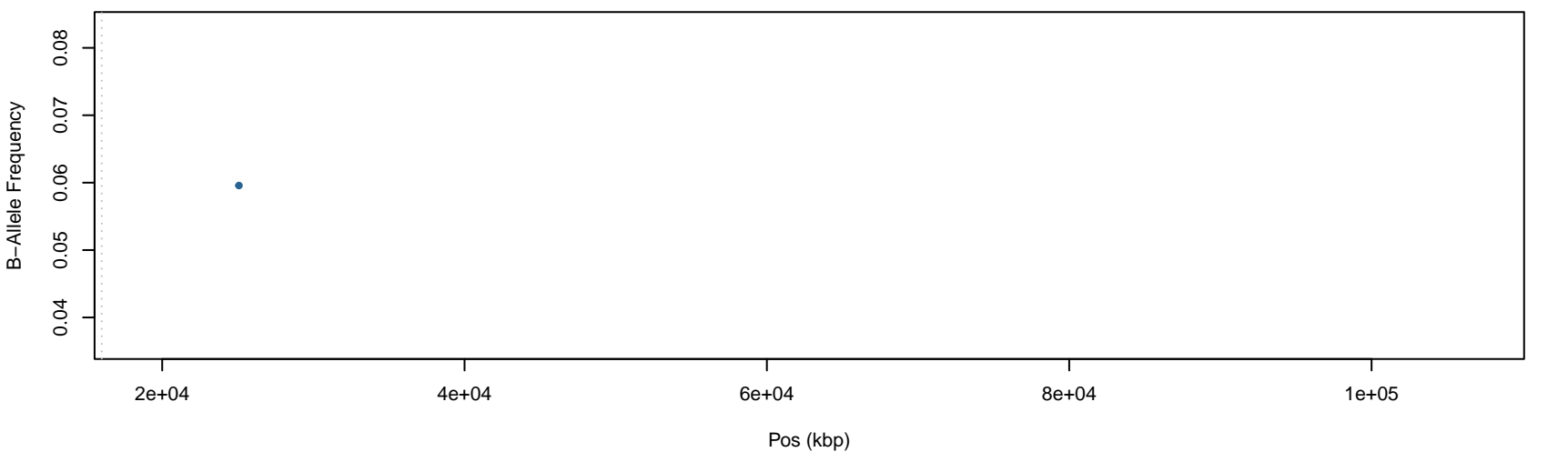
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 11



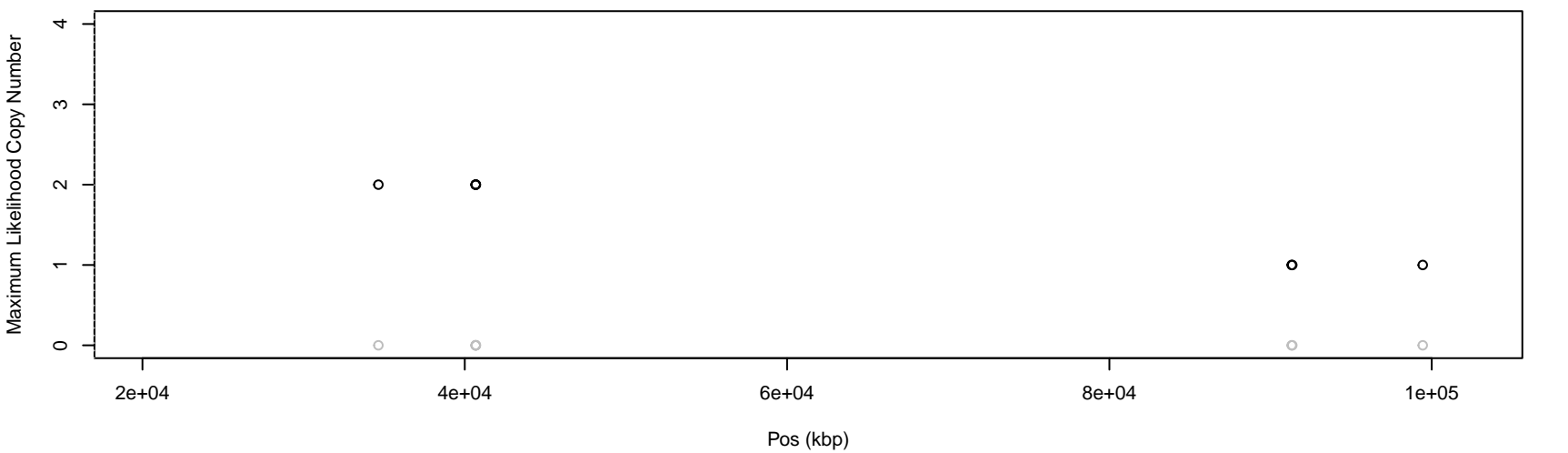
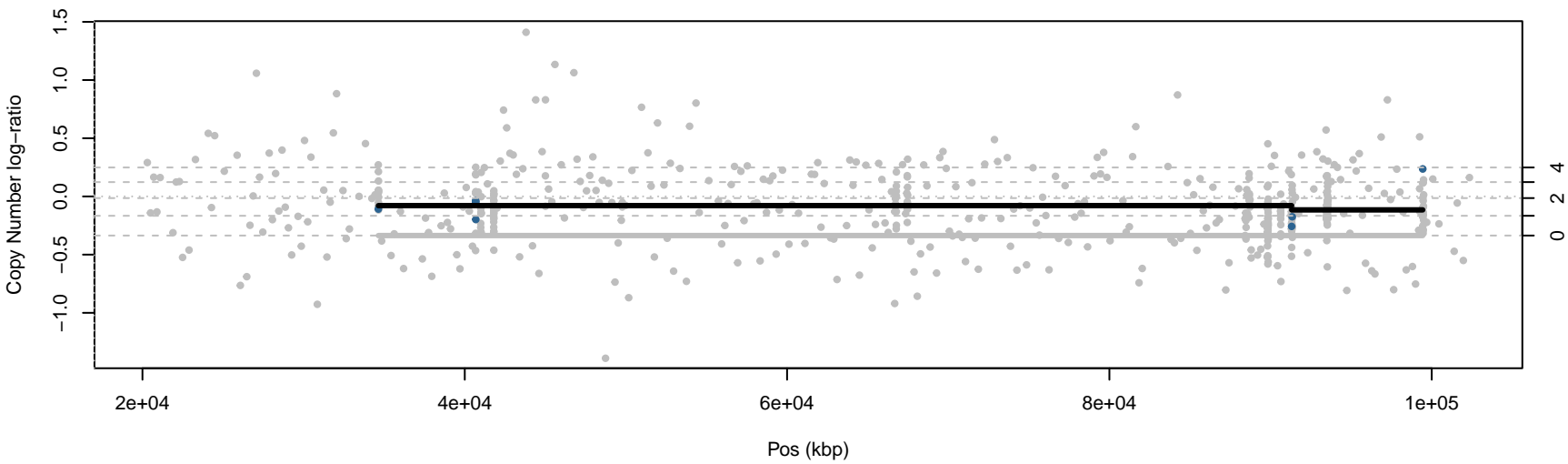
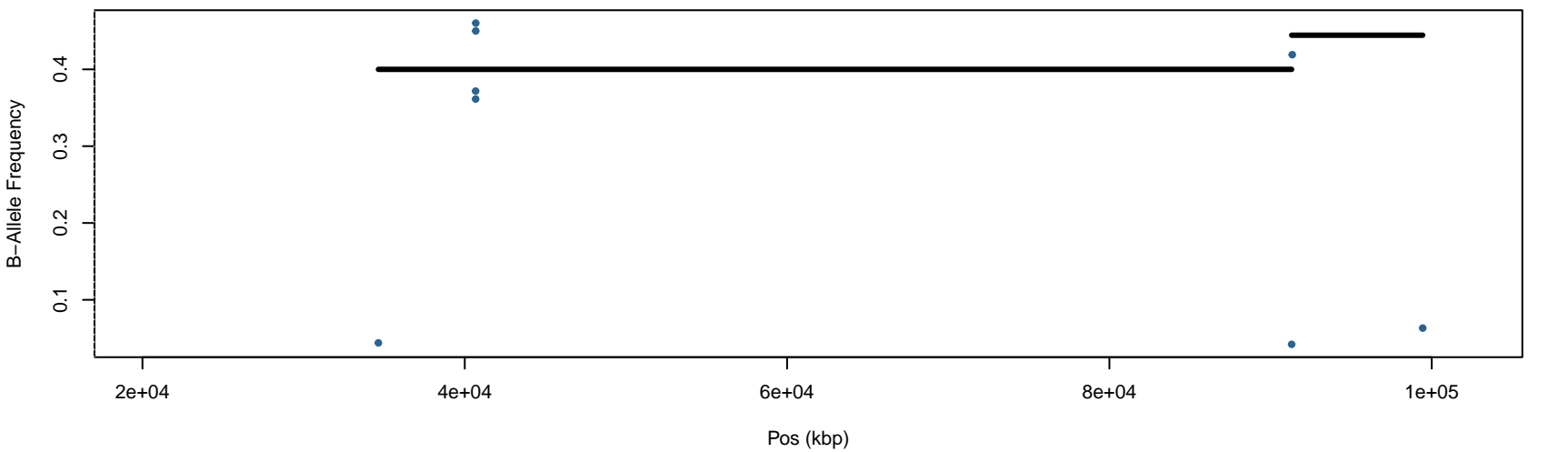
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 12



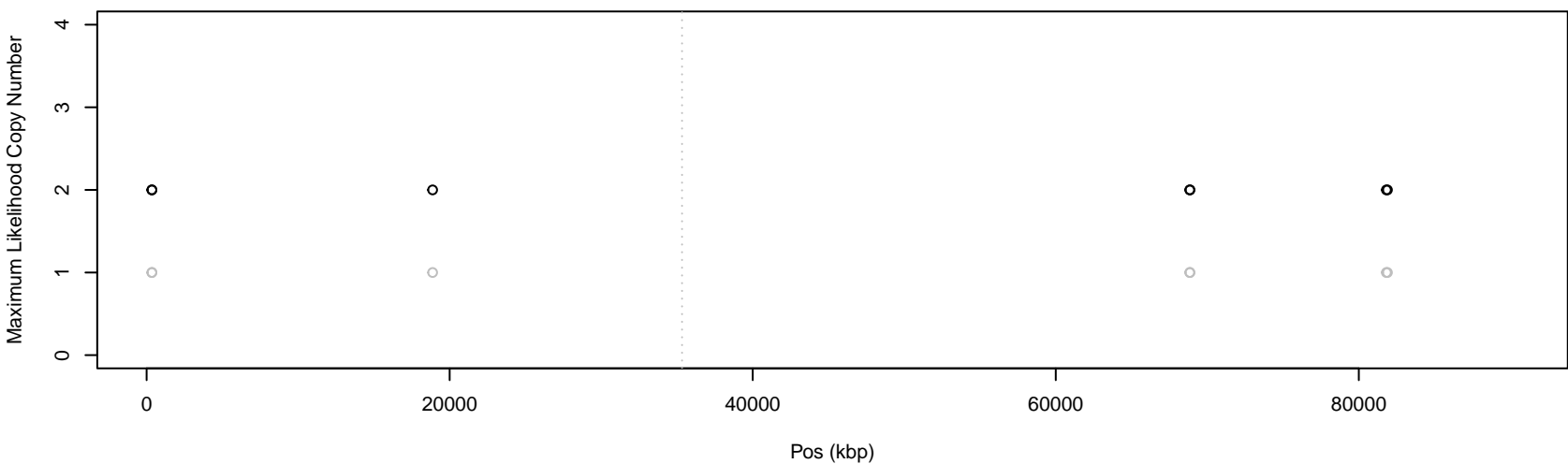
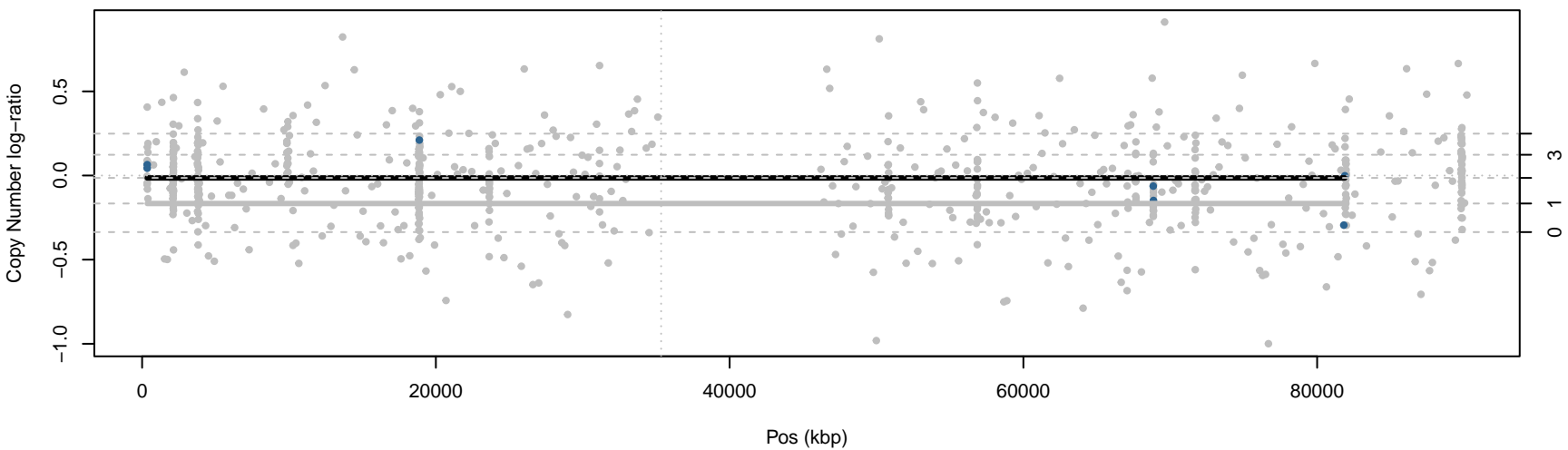
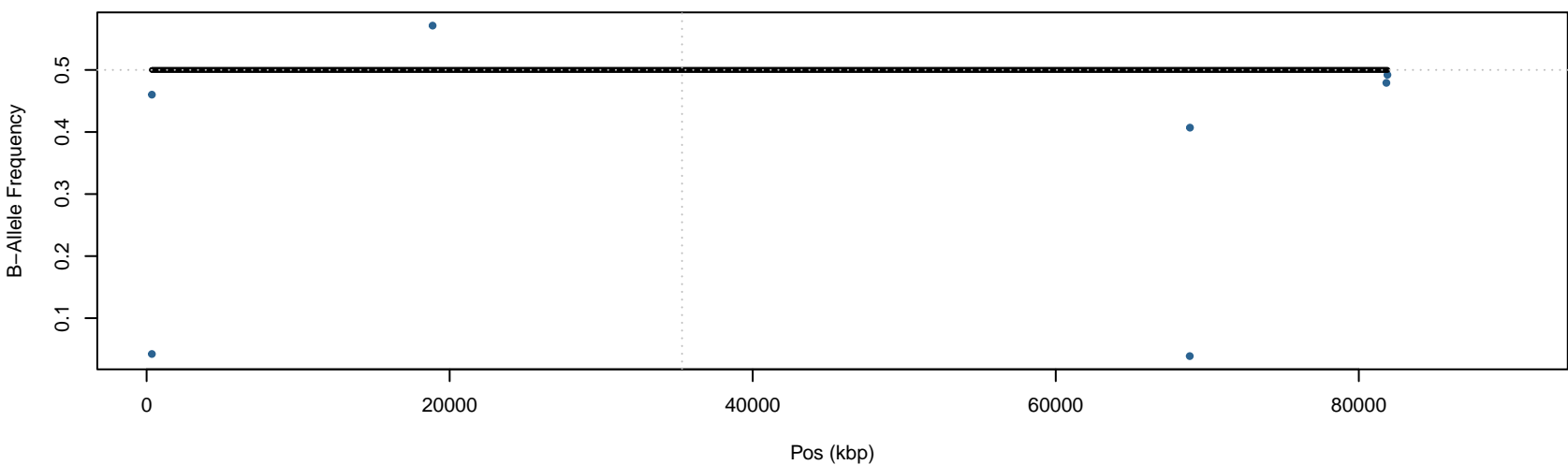
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 14



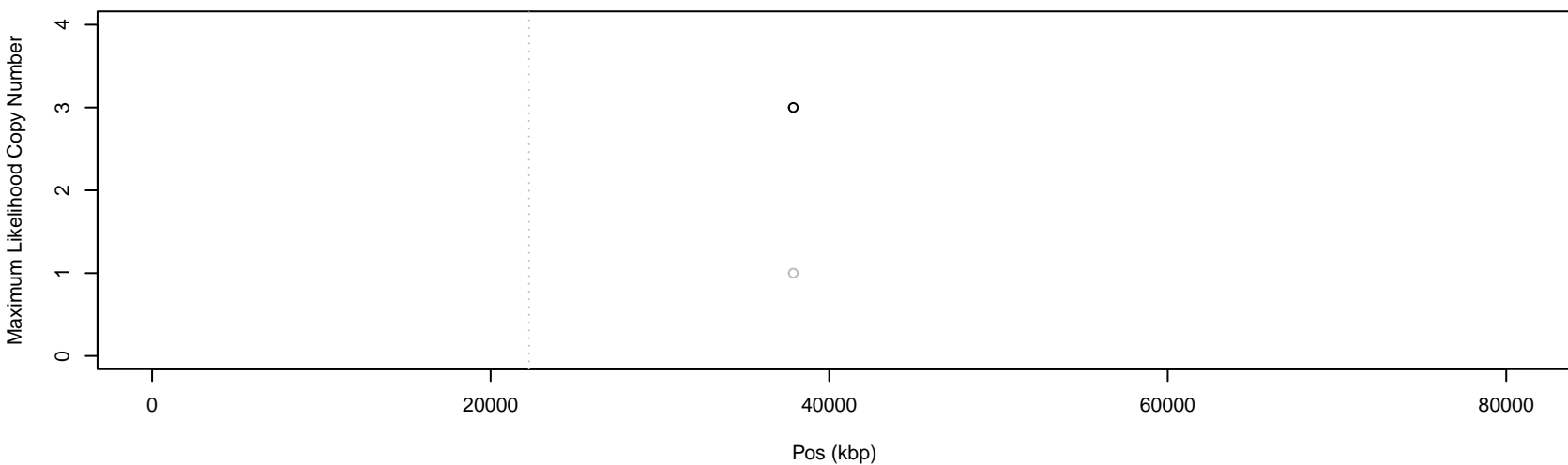
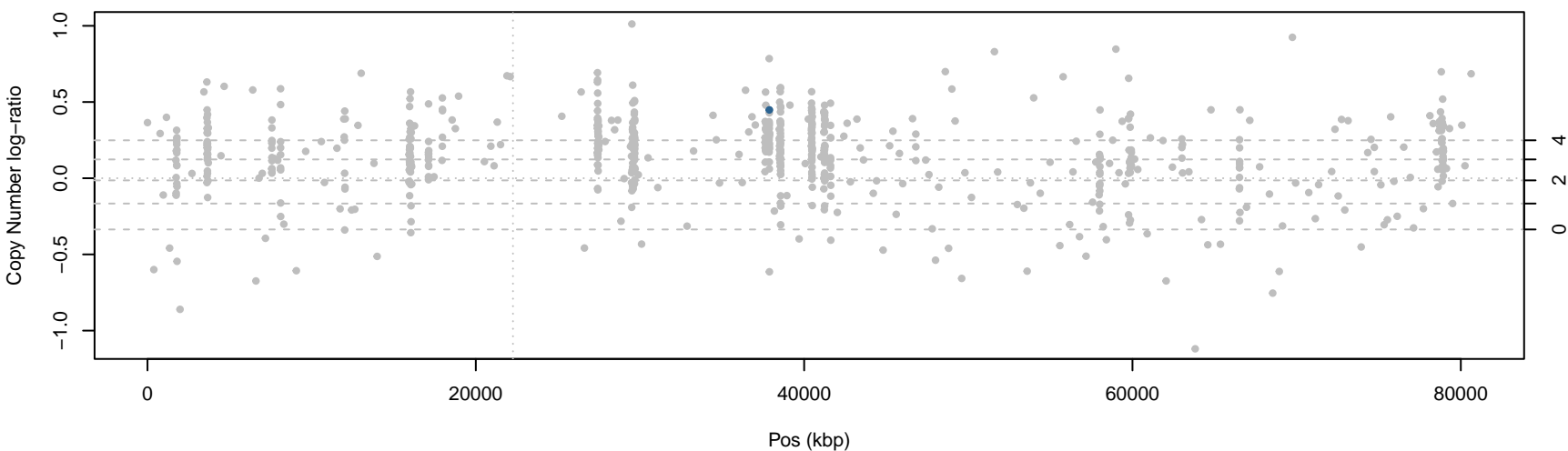
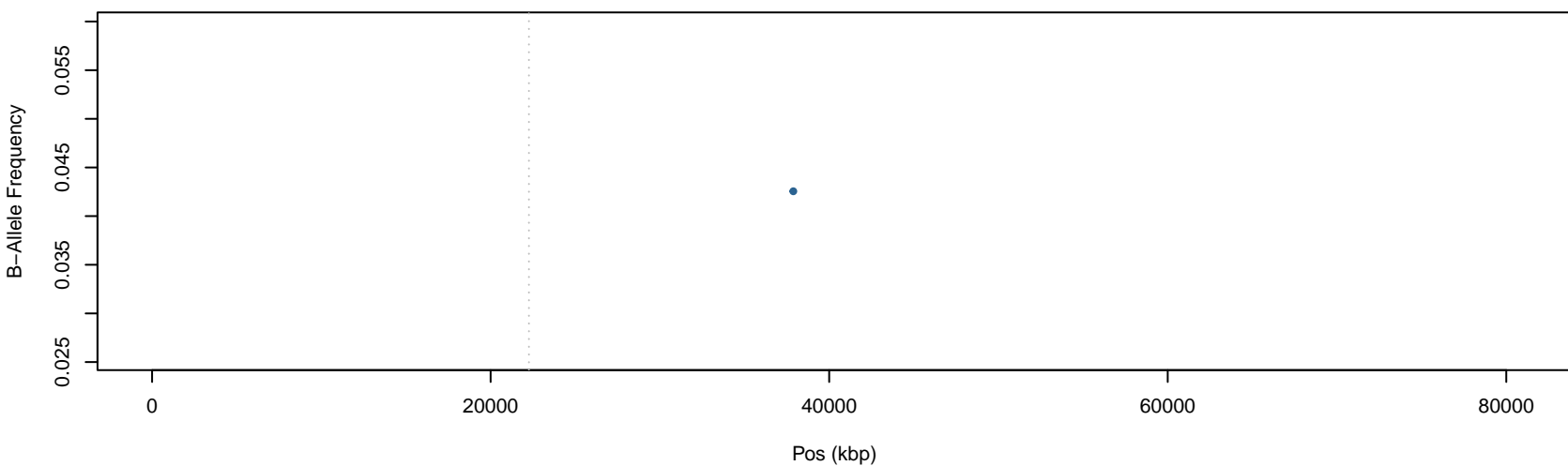
Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 15



Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 16



Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 17



Purity: 0.2 Tumor ploidy: 2.099 SNV log-likelihood: 280.05 GoF: 92.7% Mean coverage: 250;235 Chromosome: 22

