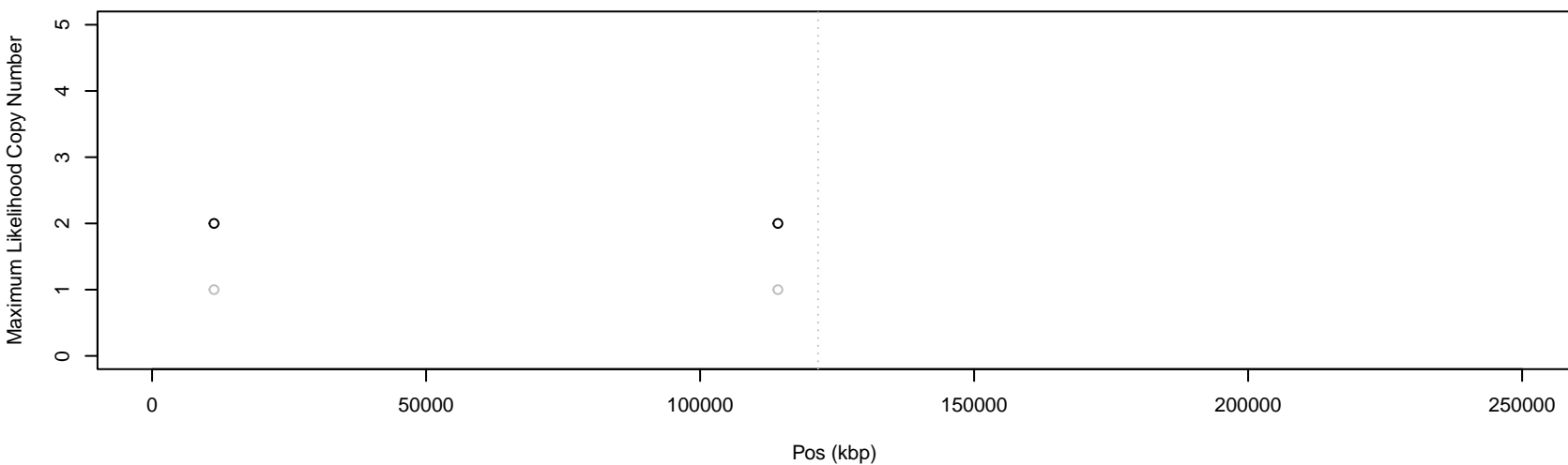
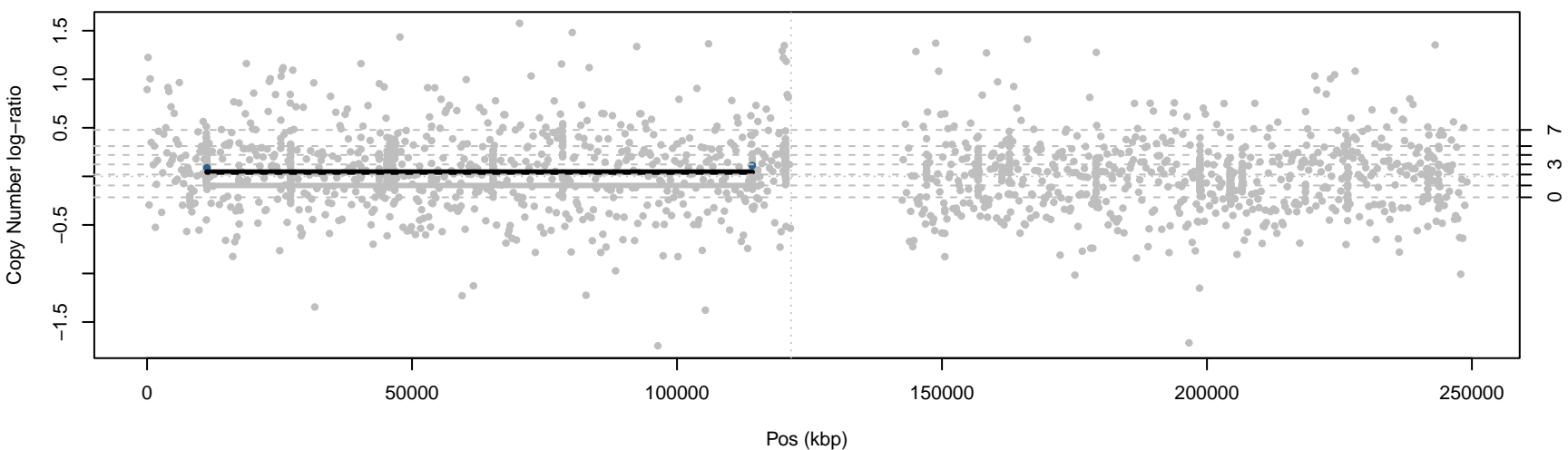
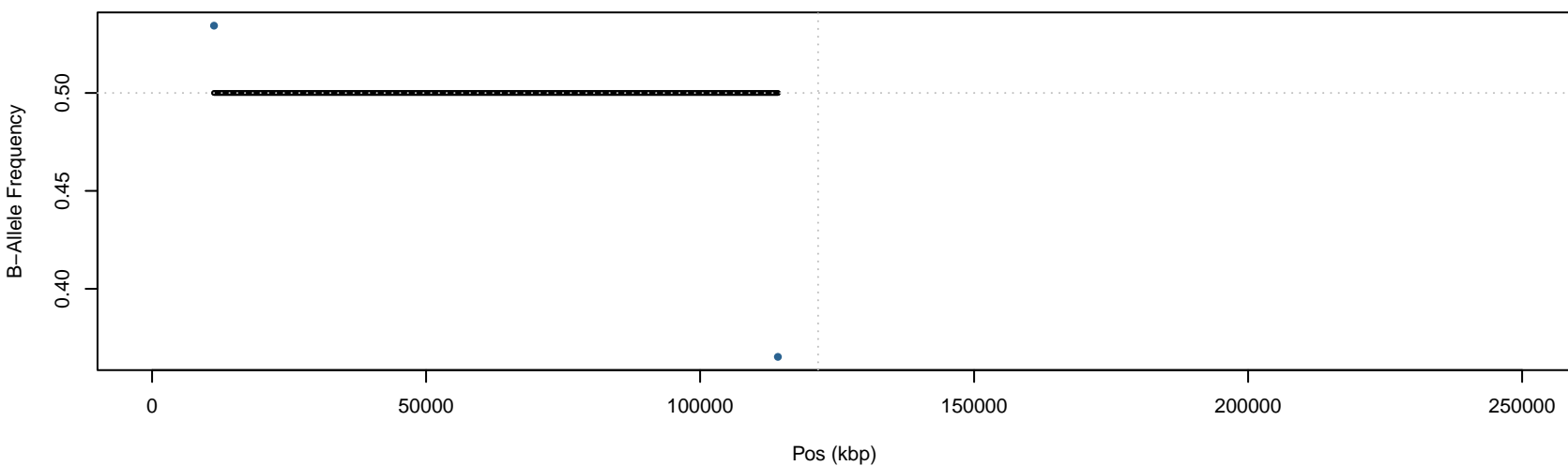
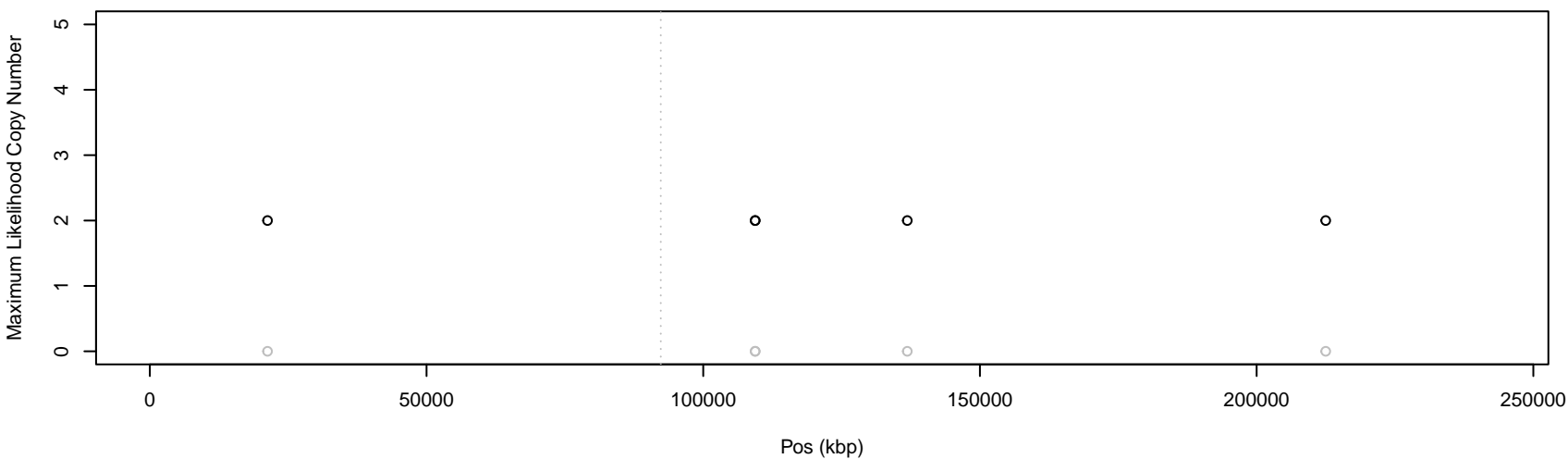
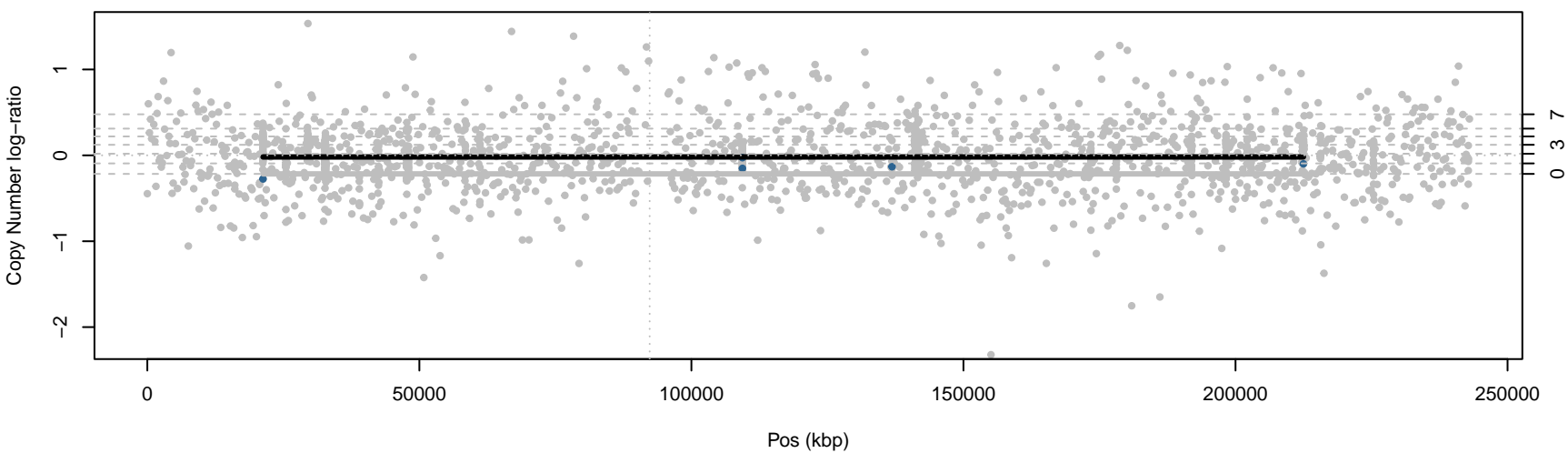
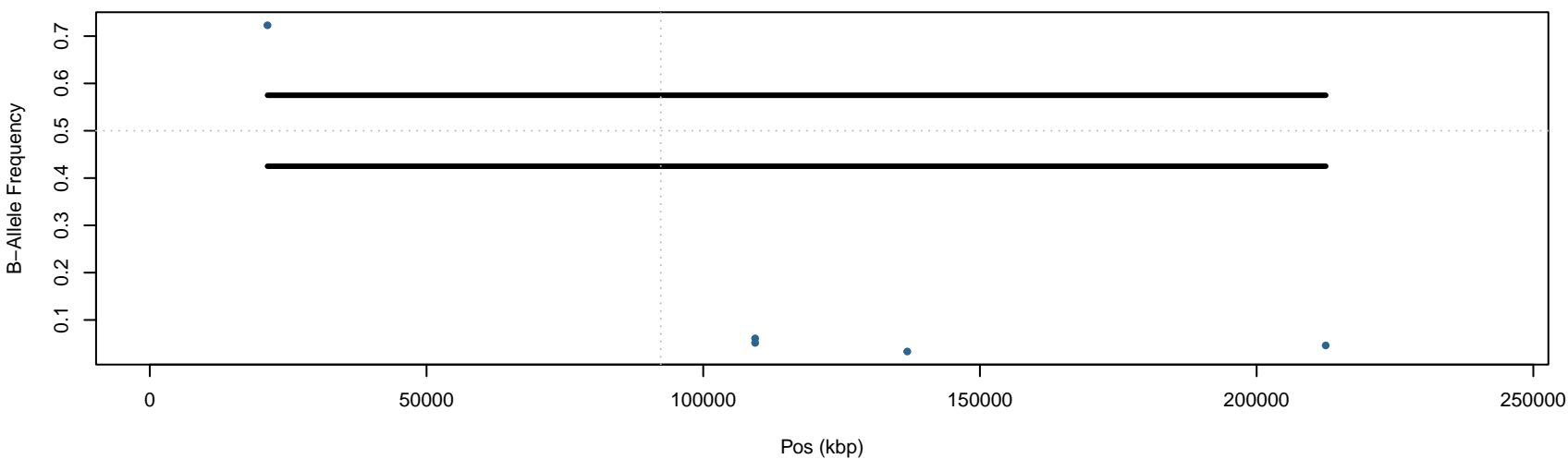


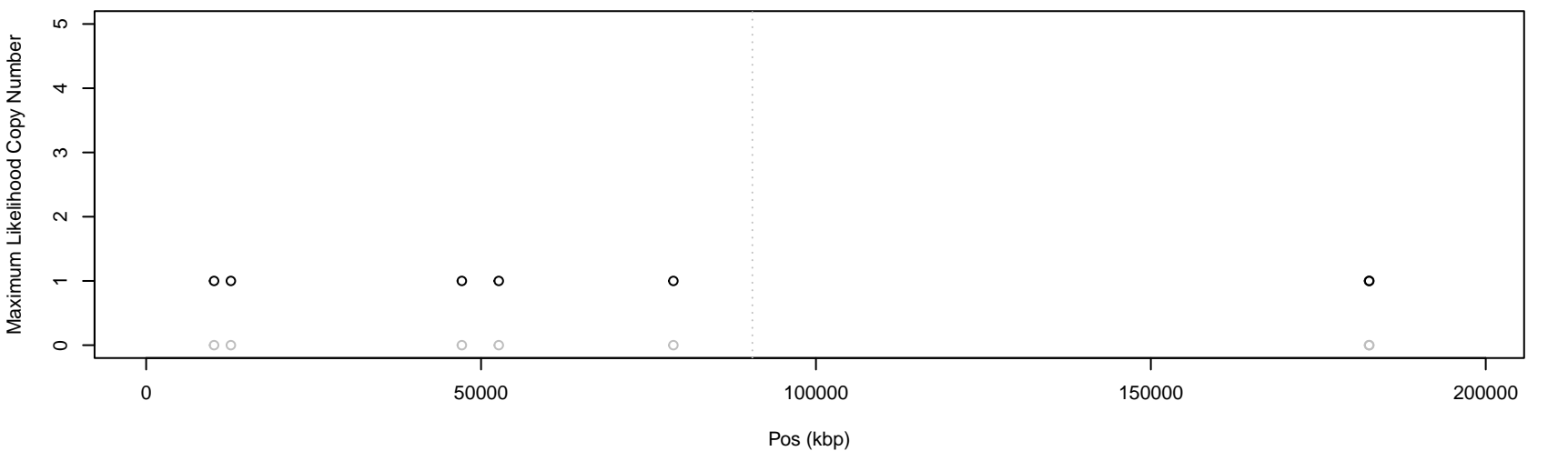
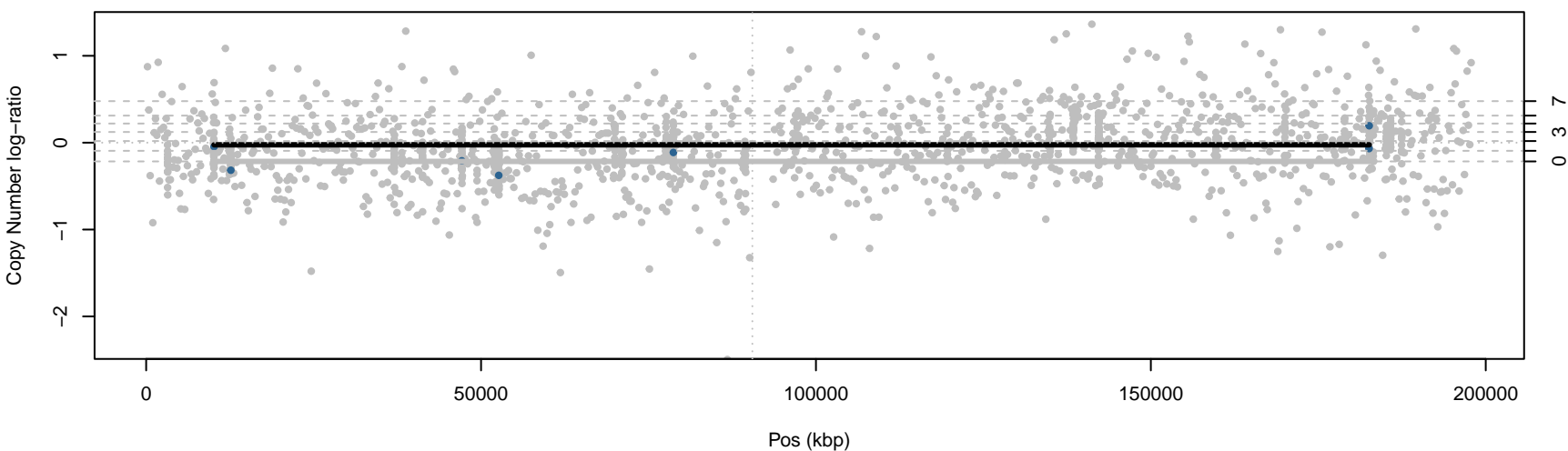
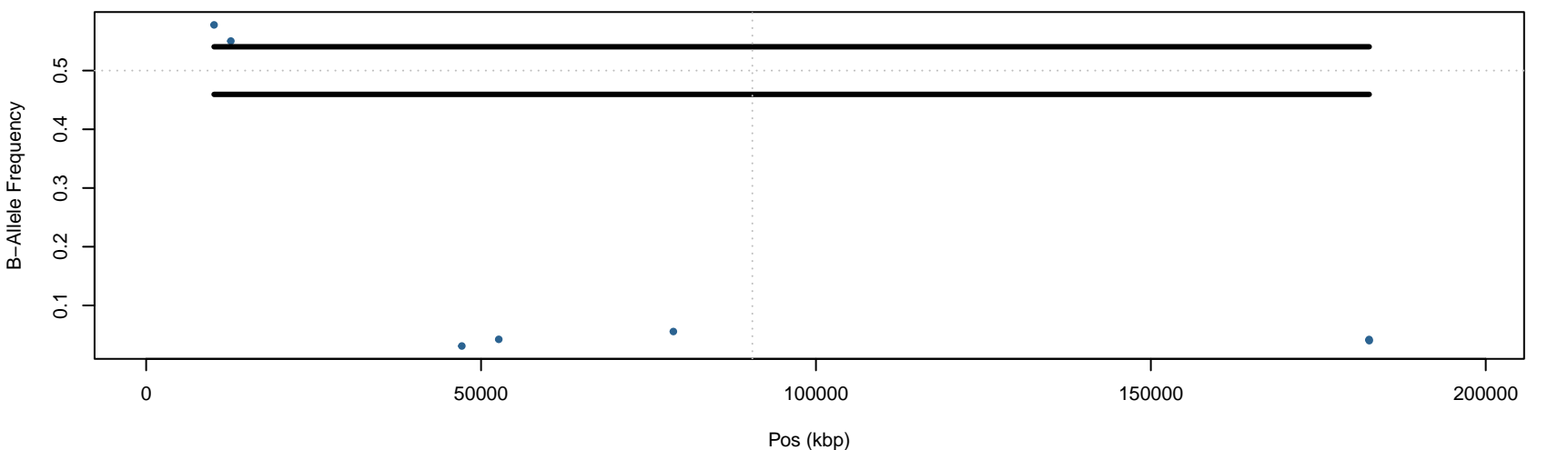
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 1



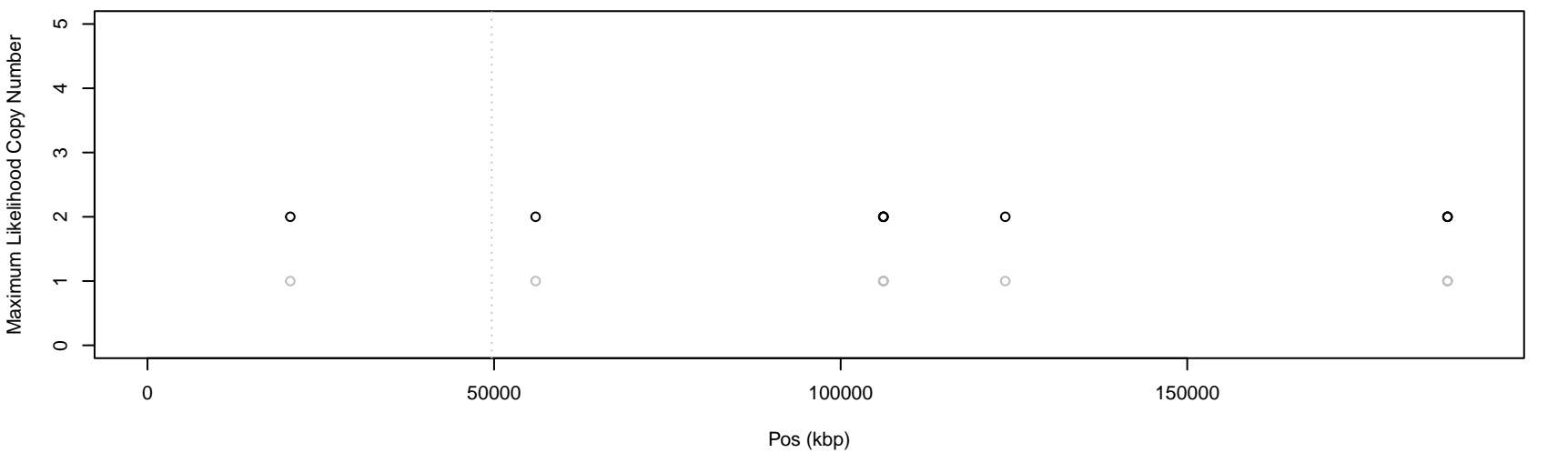
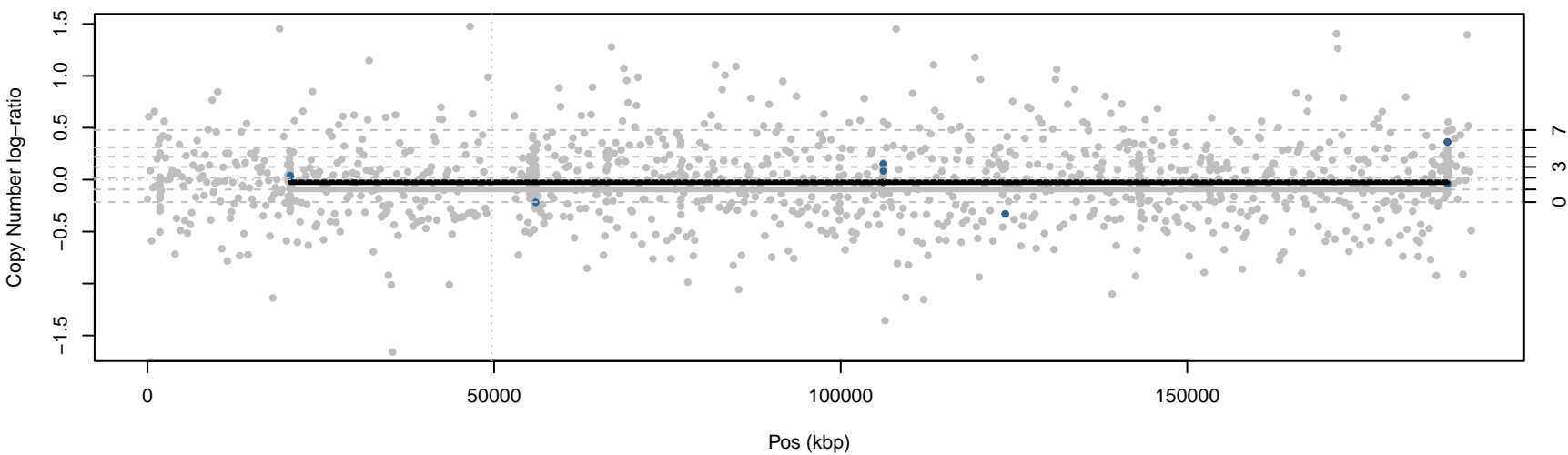
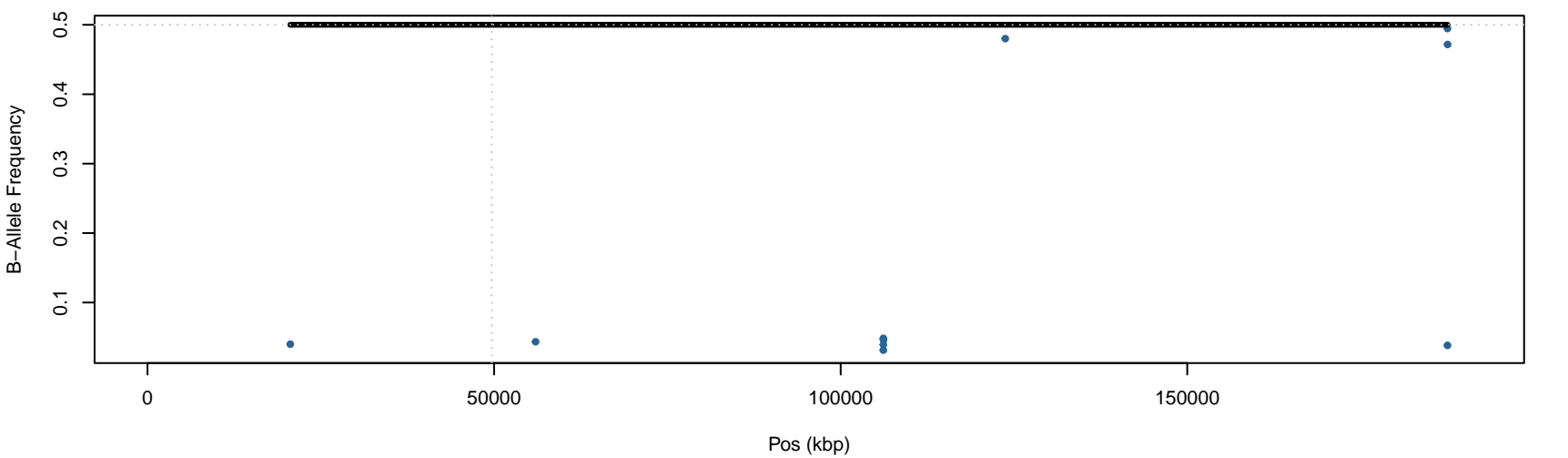
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 2



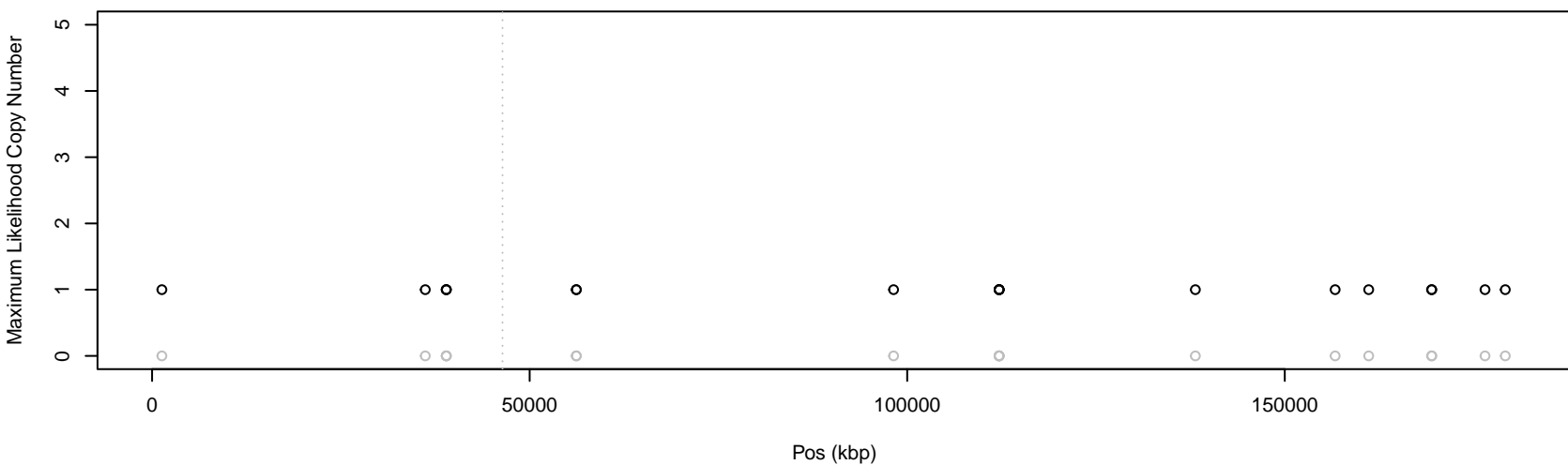
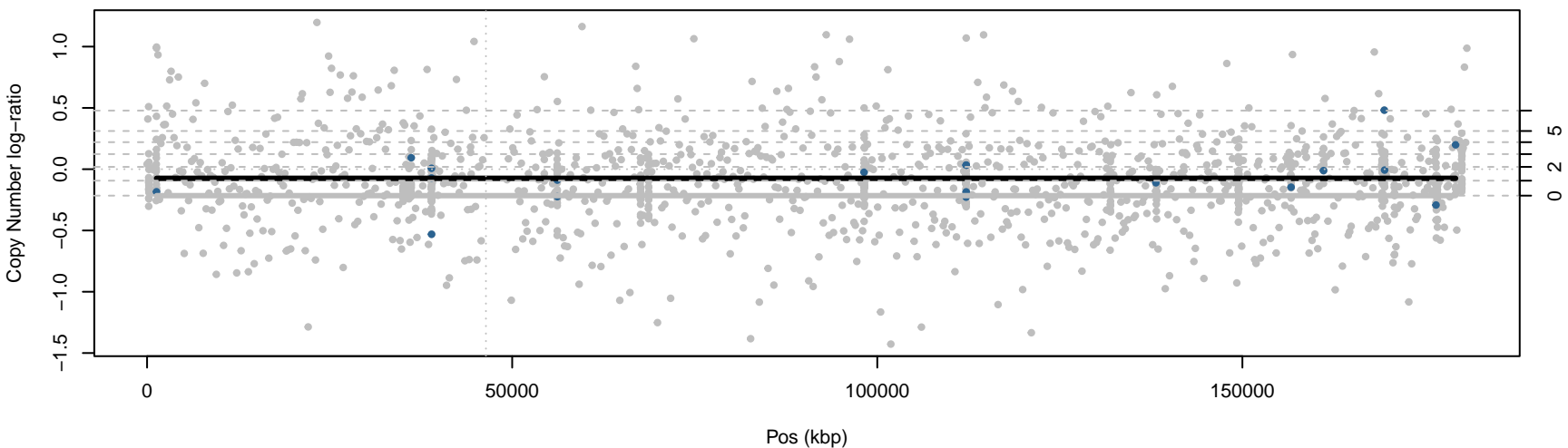
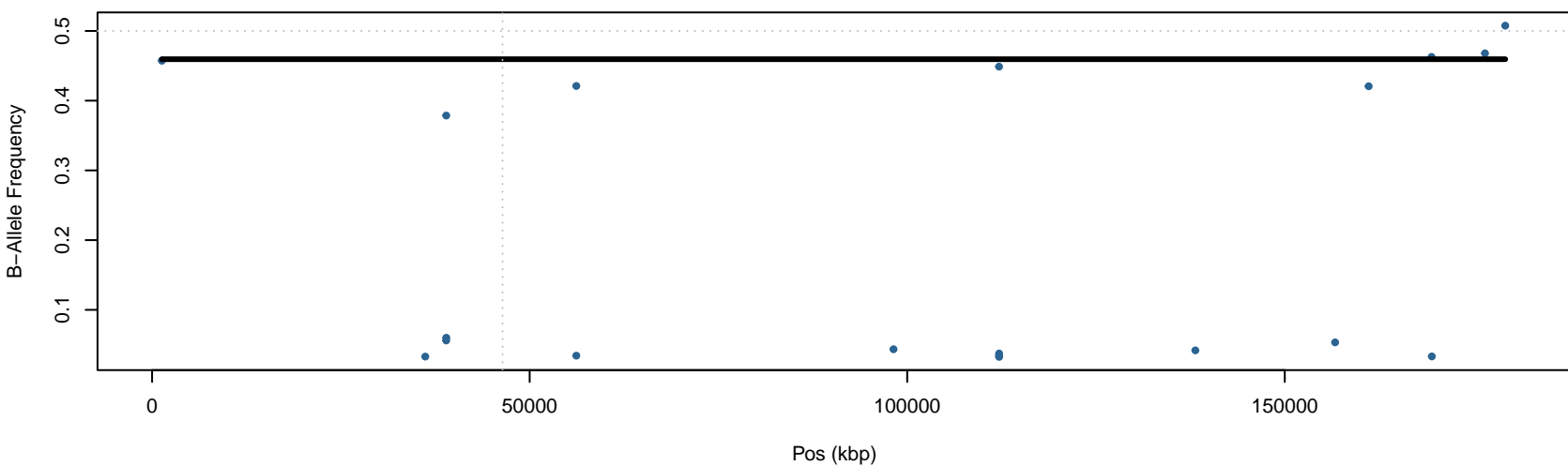
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 3



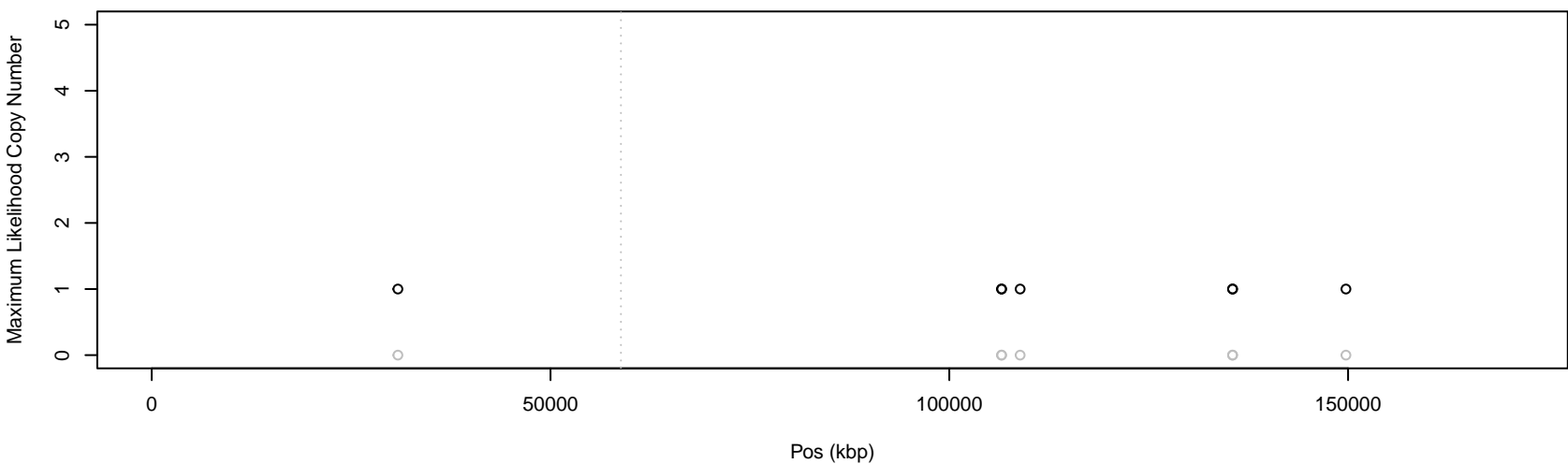
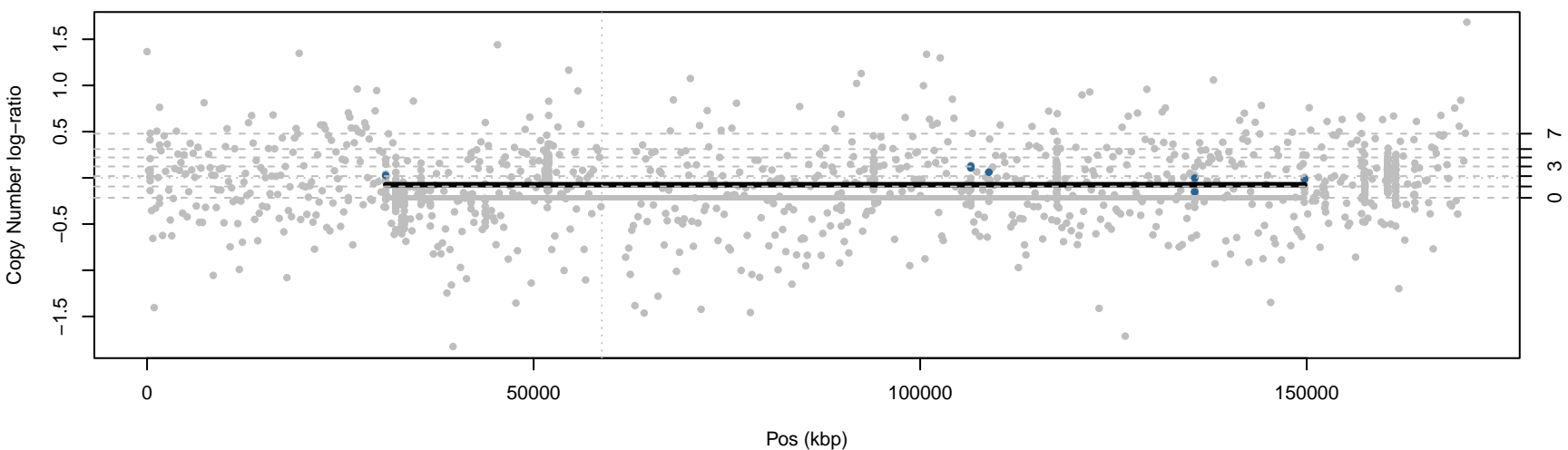
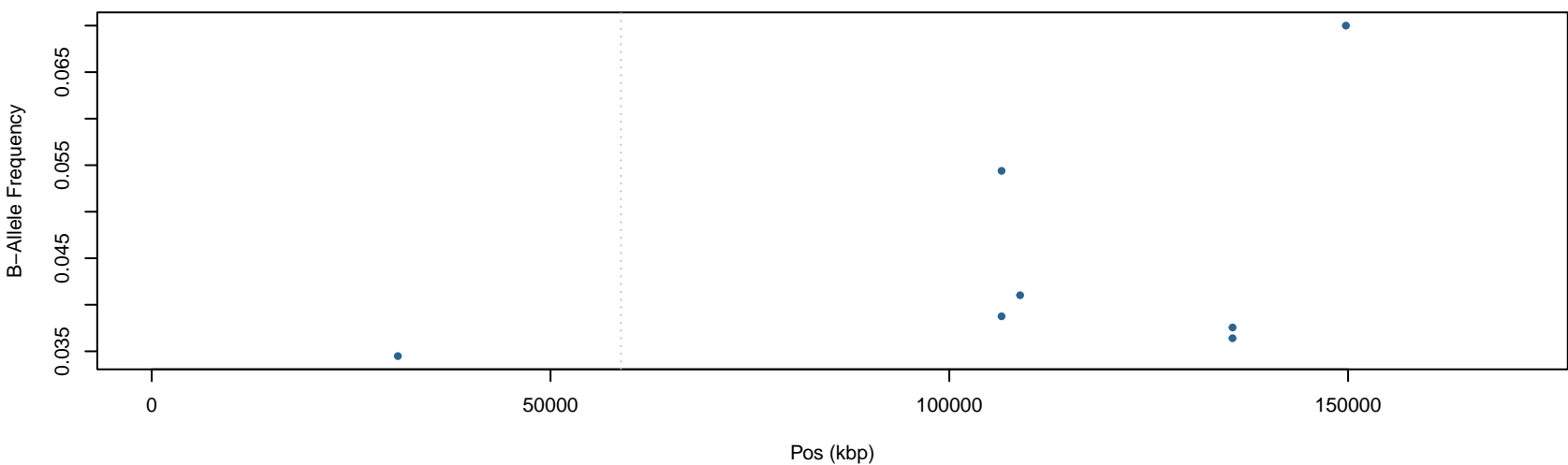
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 4



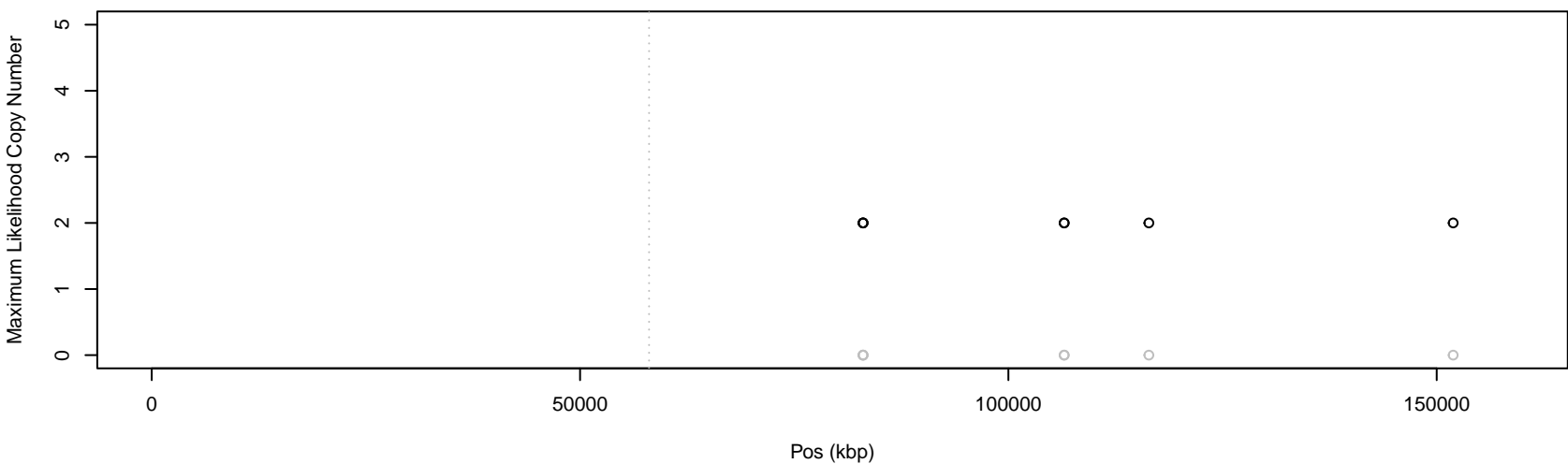
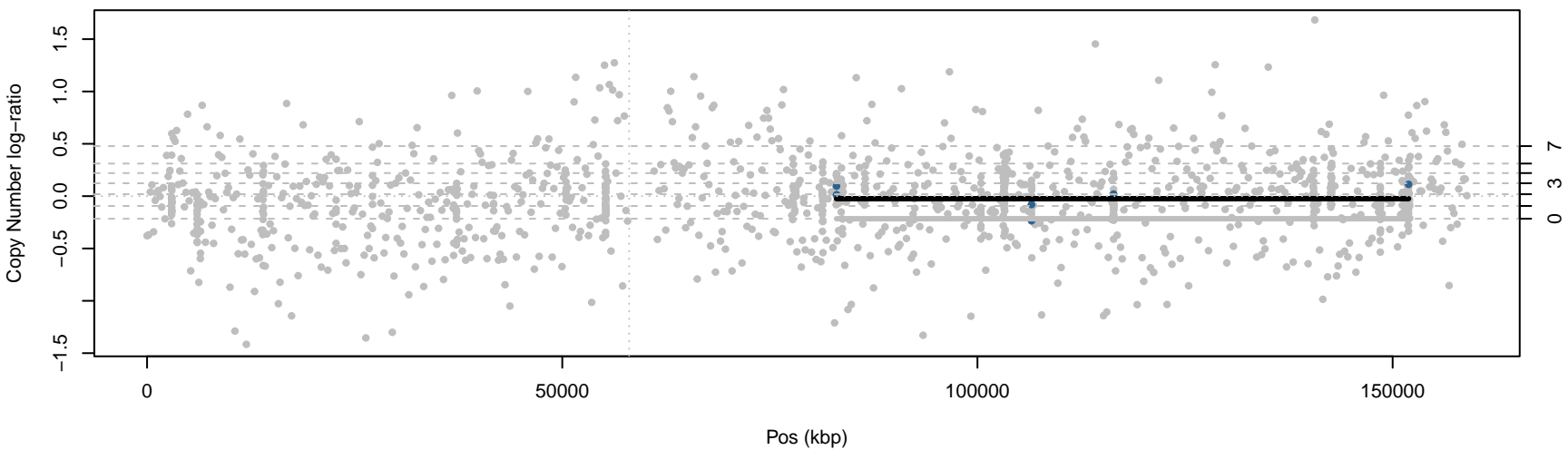
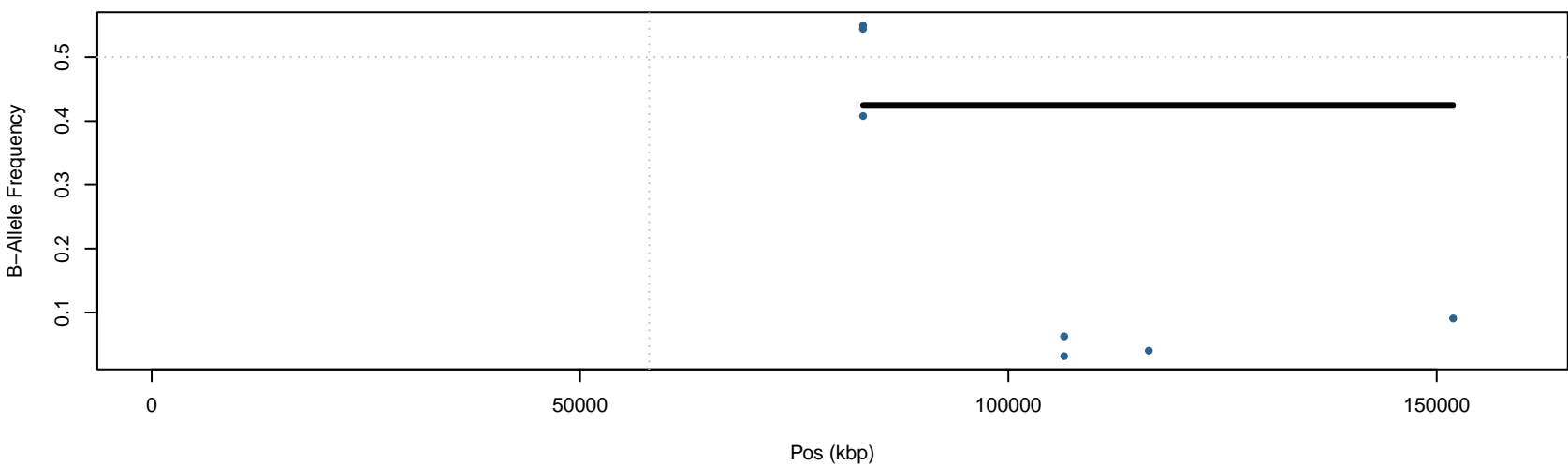
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 5



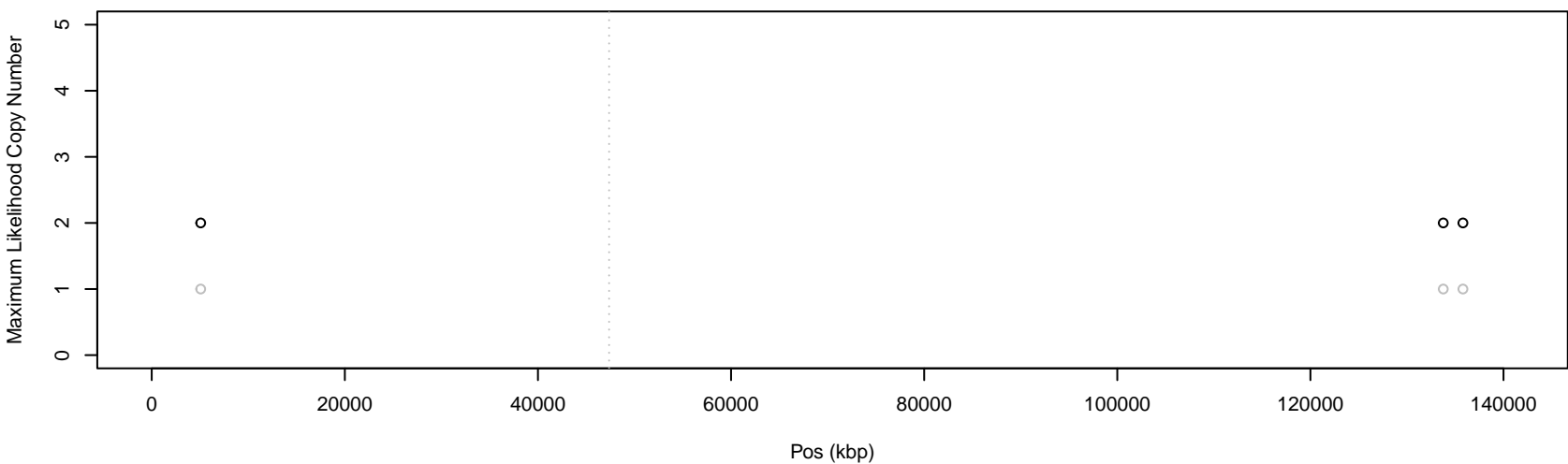
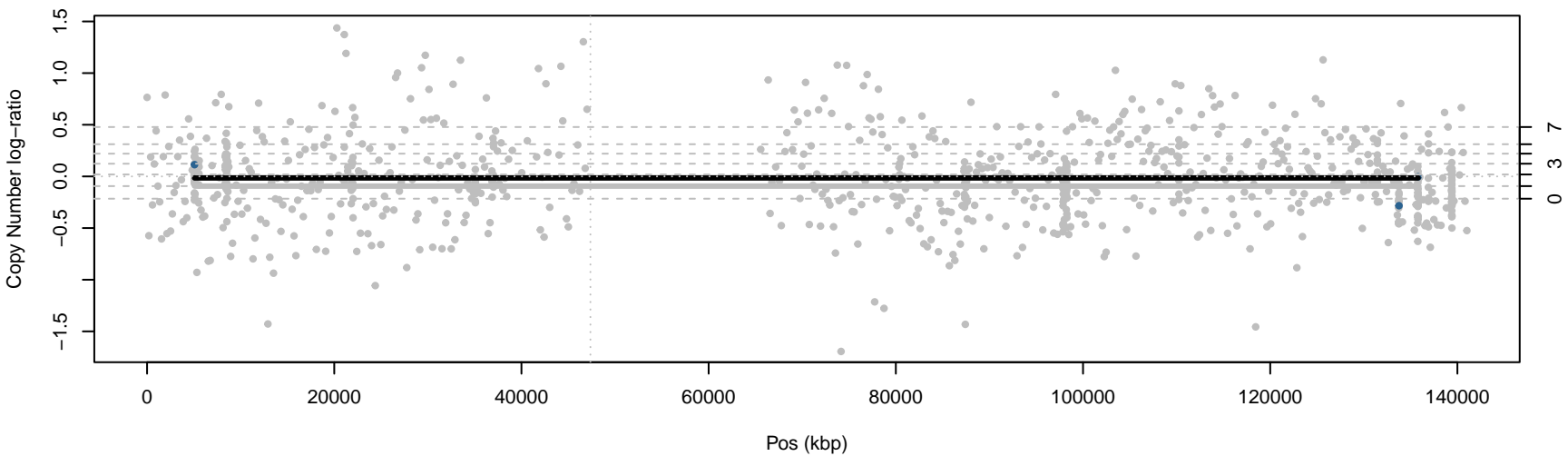
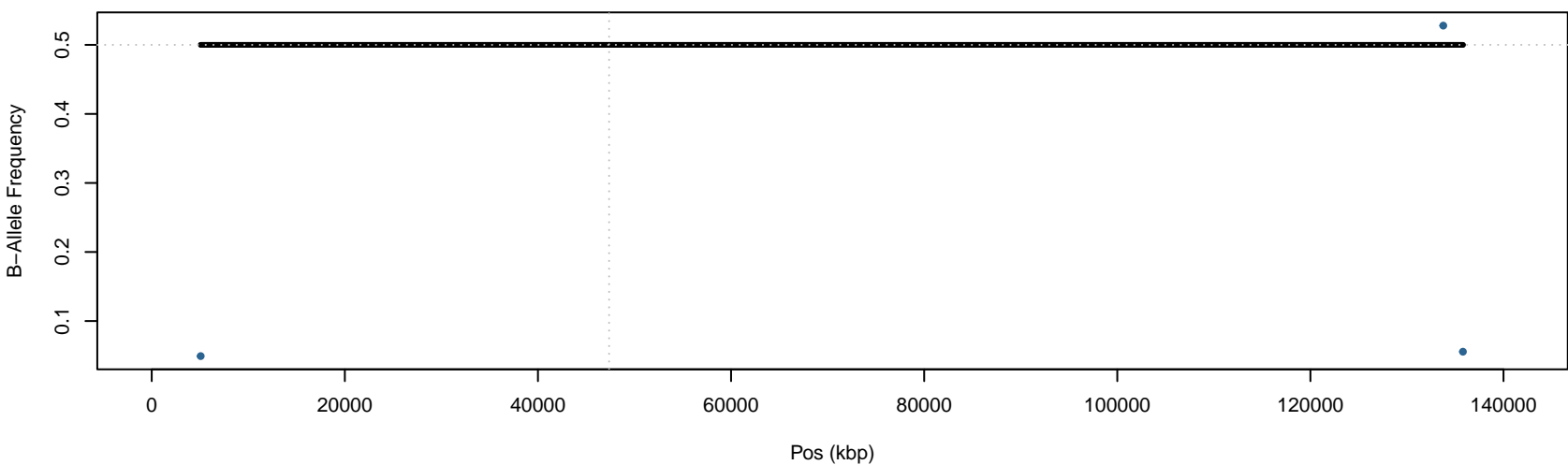
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 6



Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 7

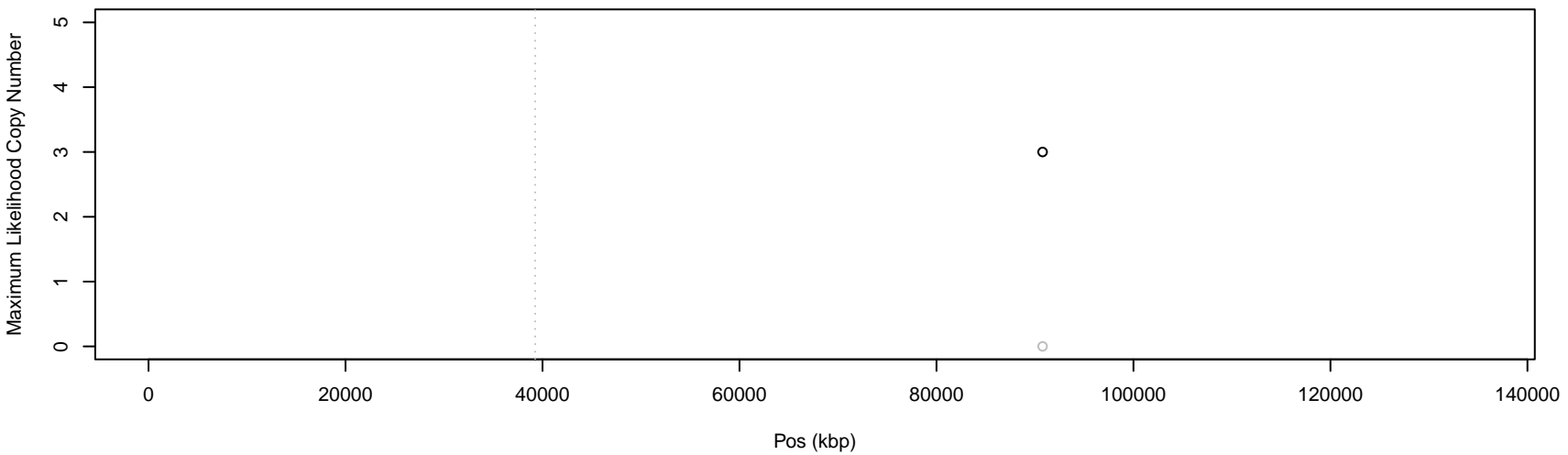
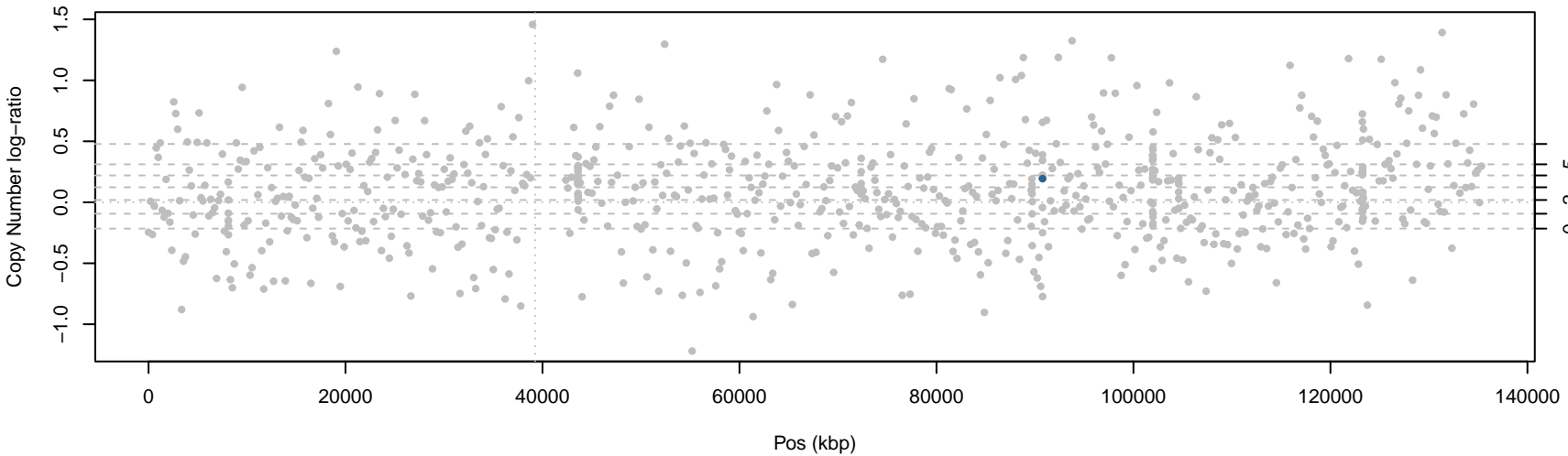
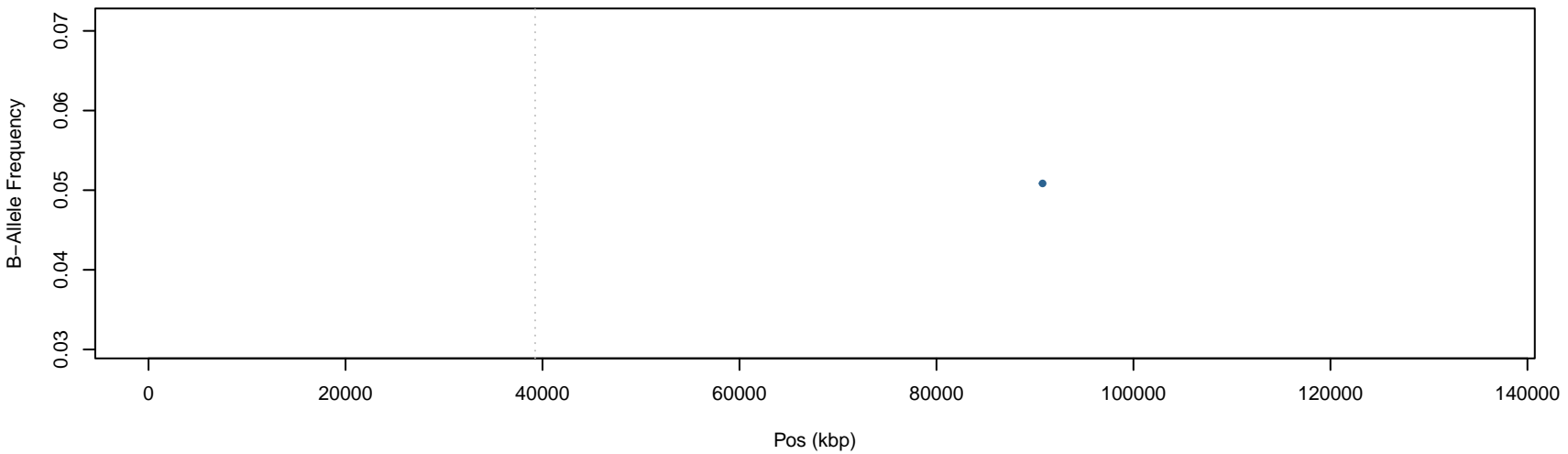


Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 9

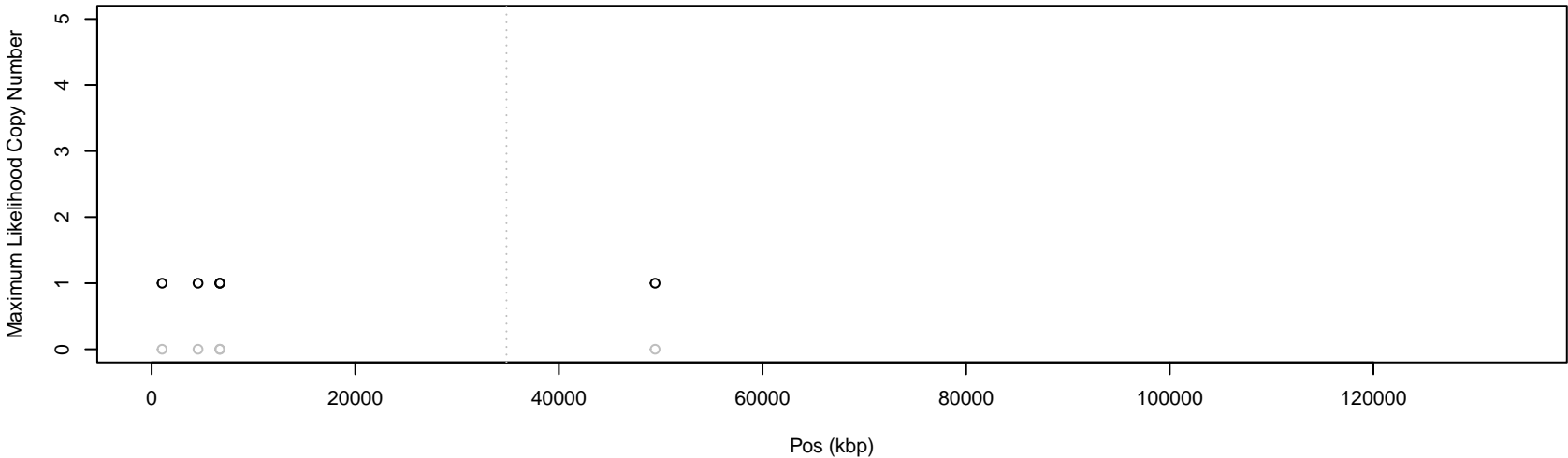
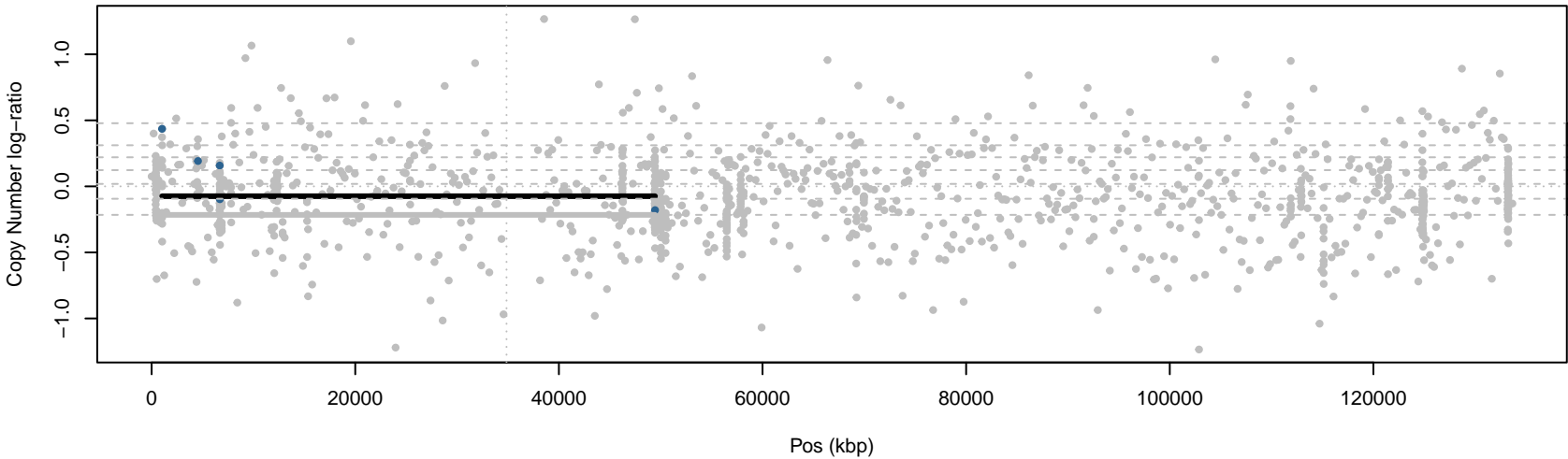
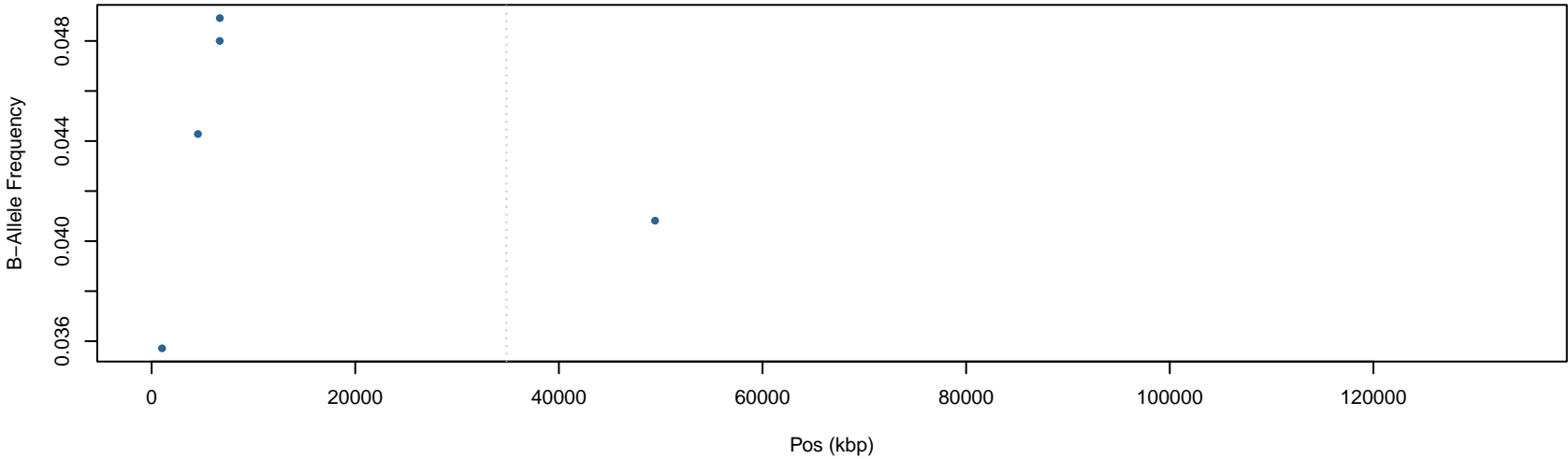




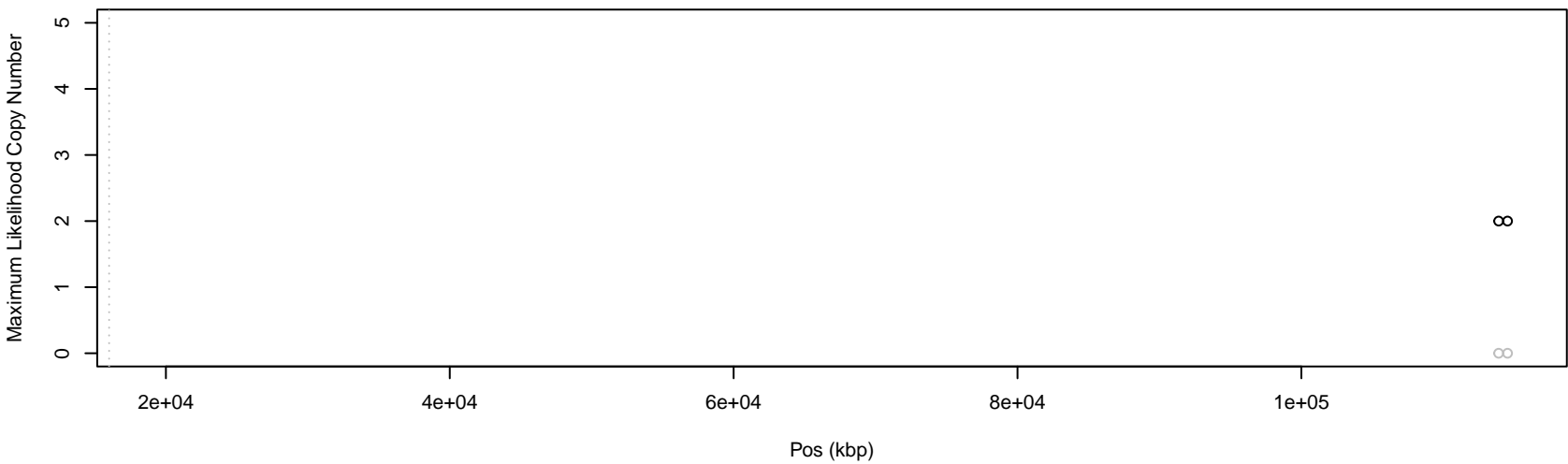
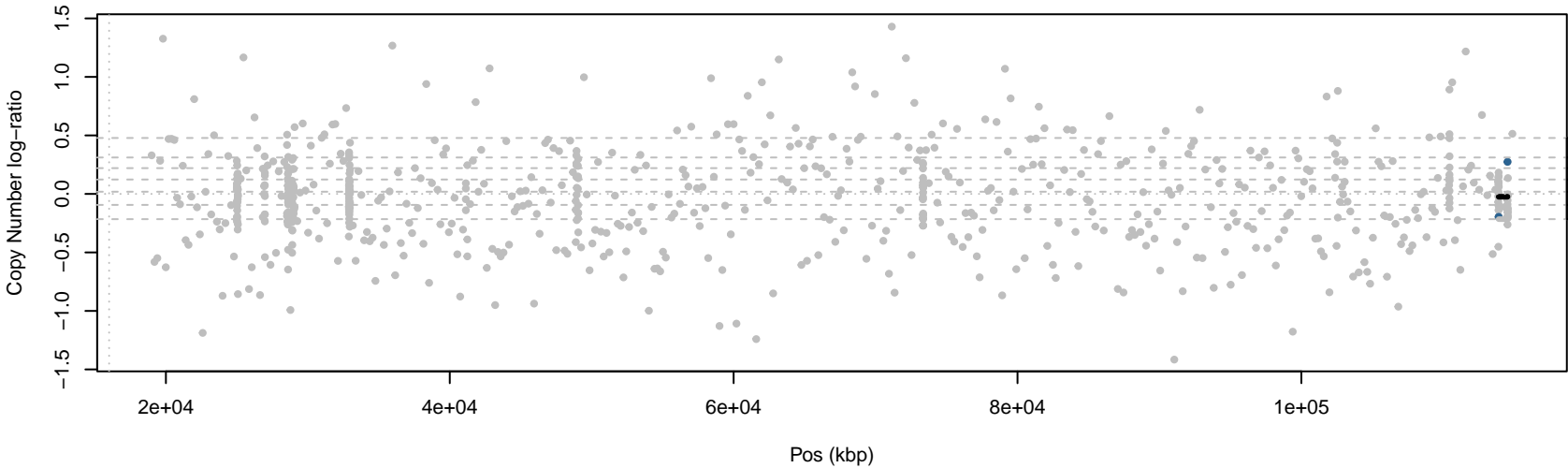
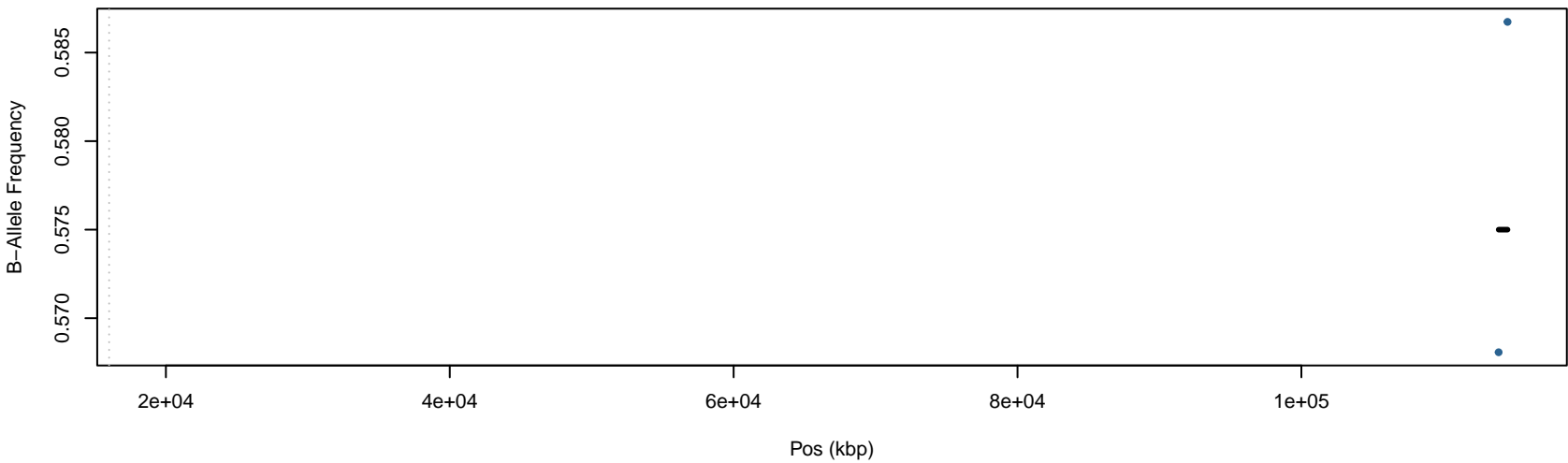
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 10



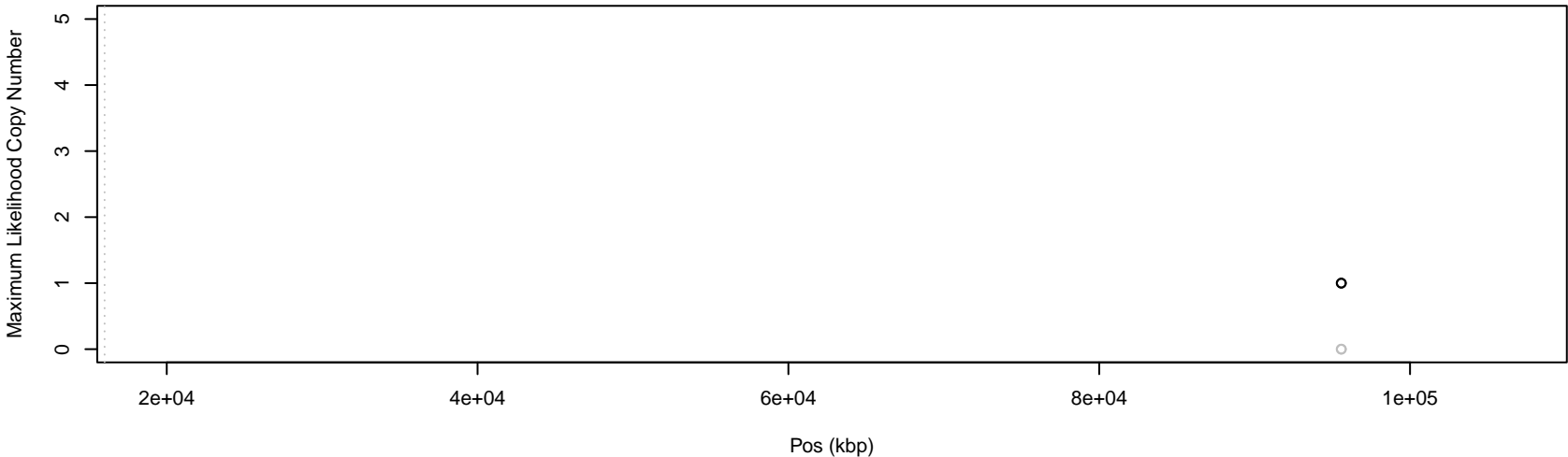
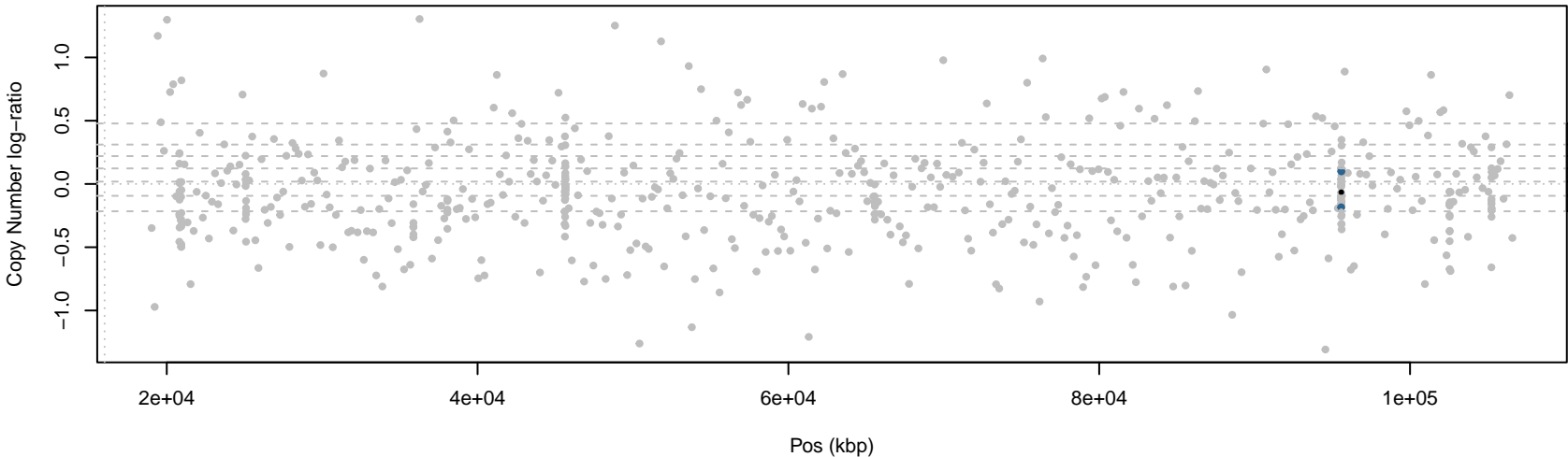
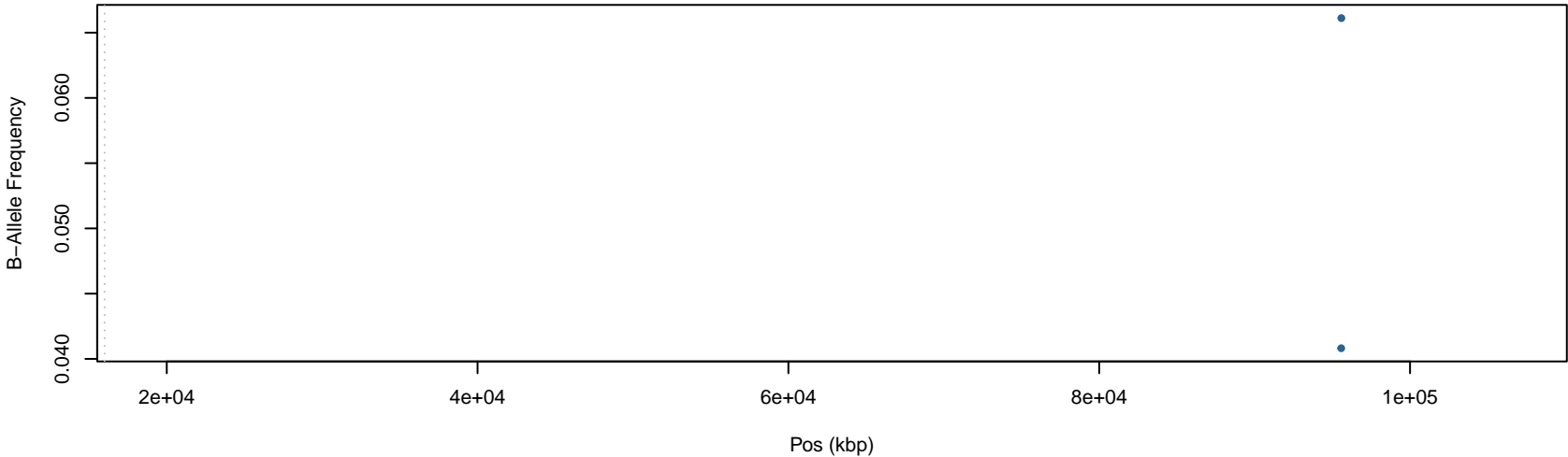
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 12



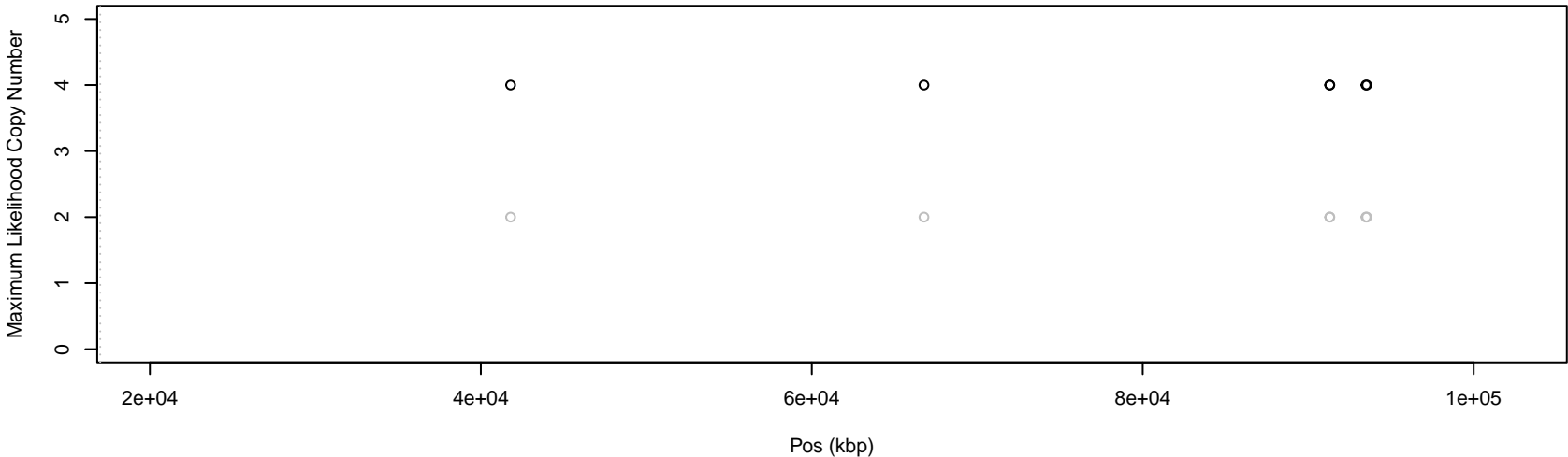
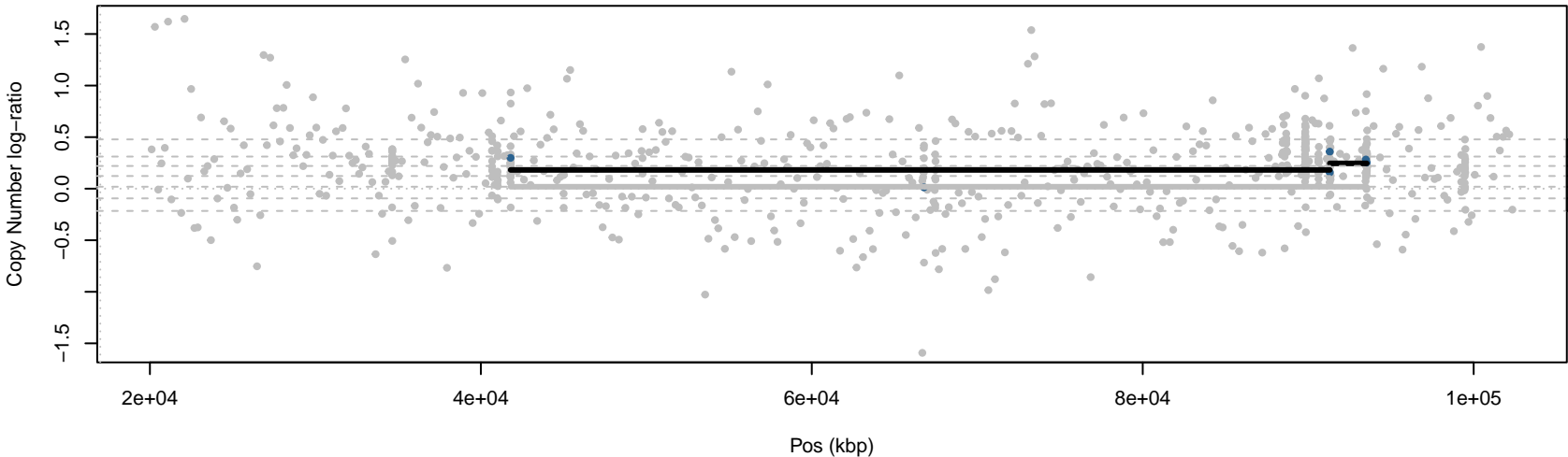
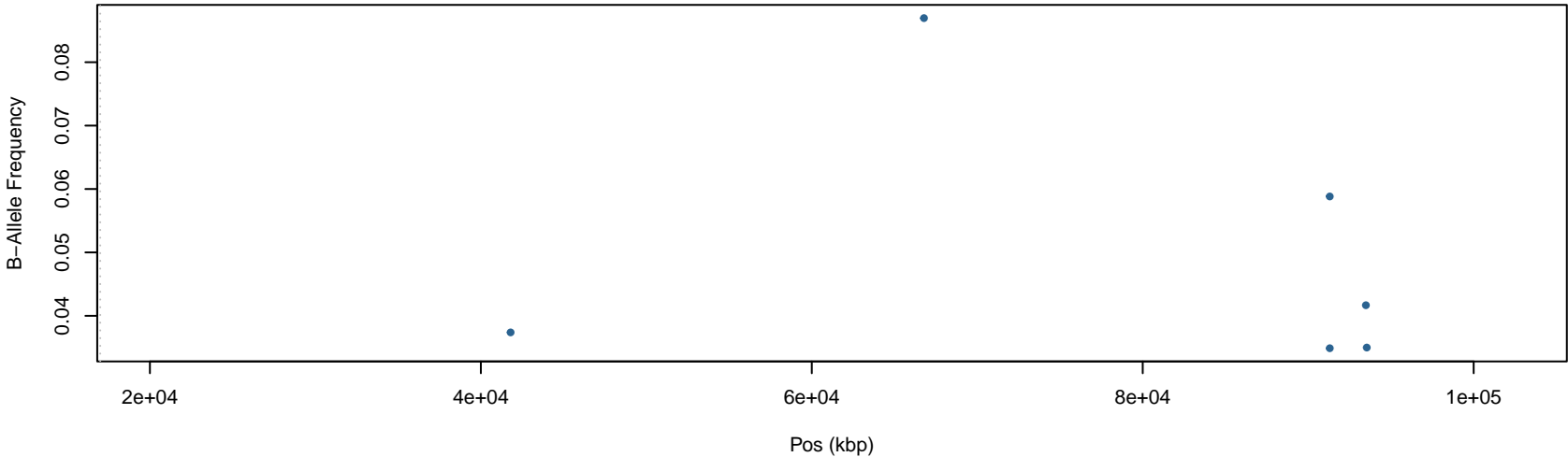
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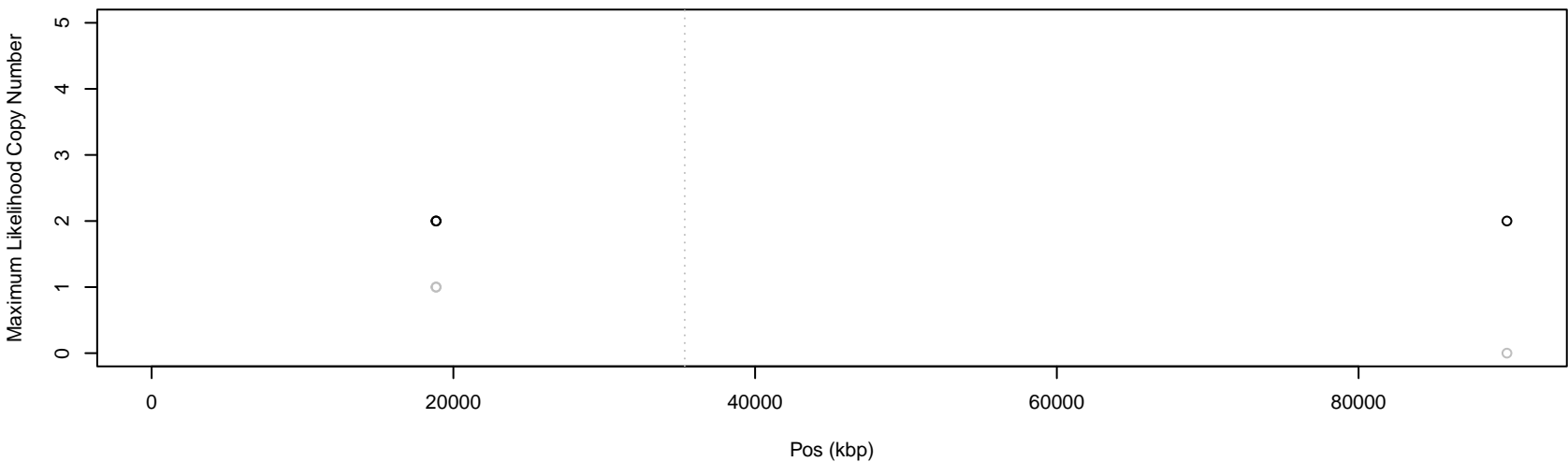
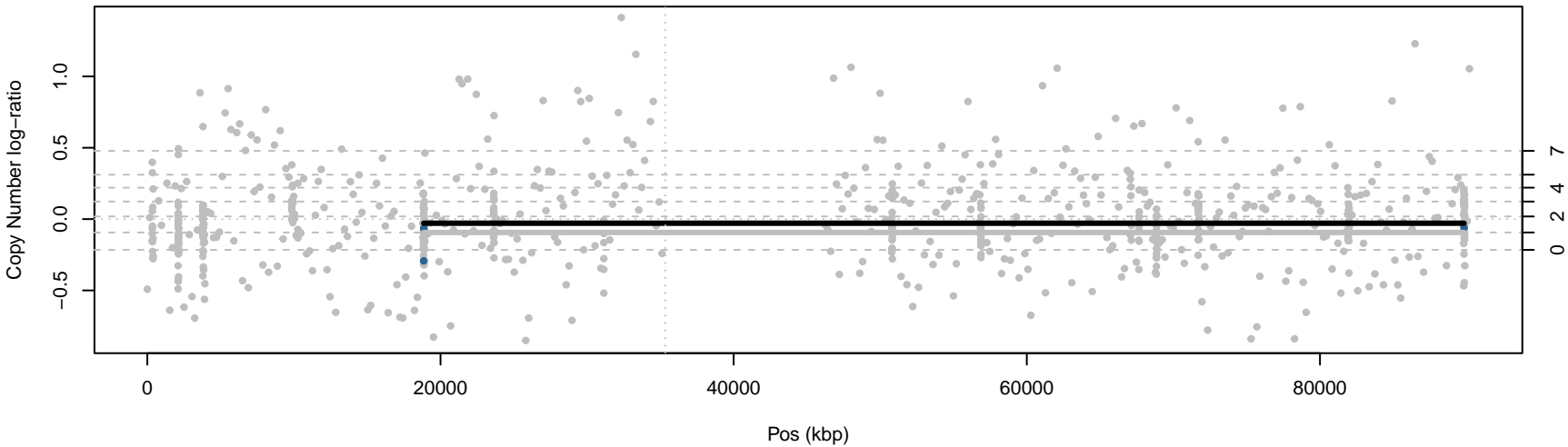
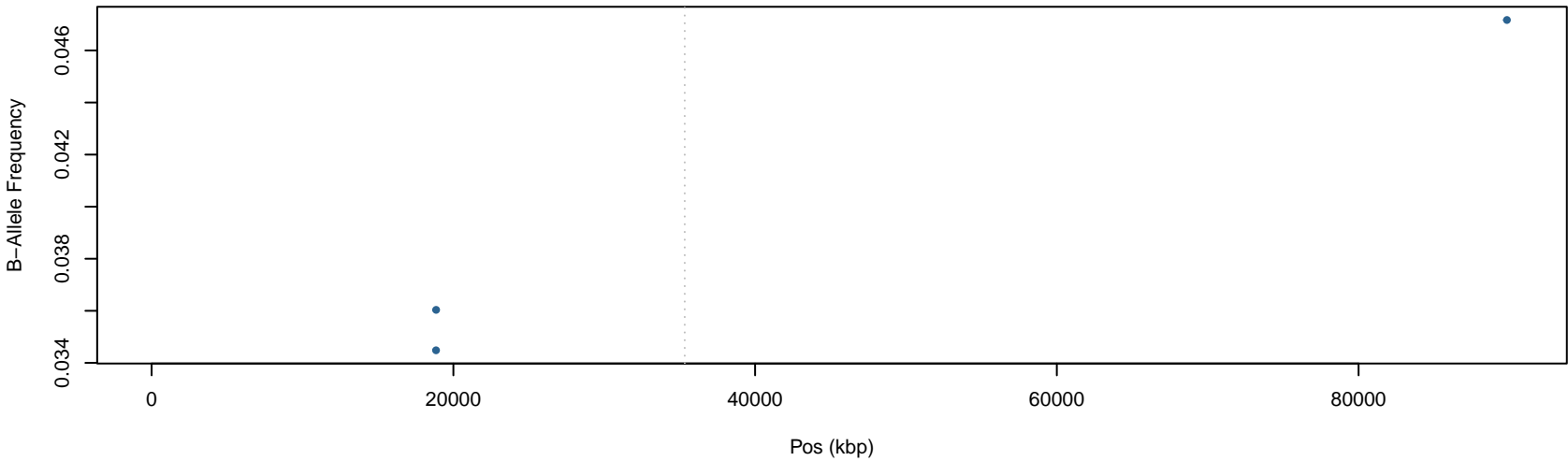
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 14



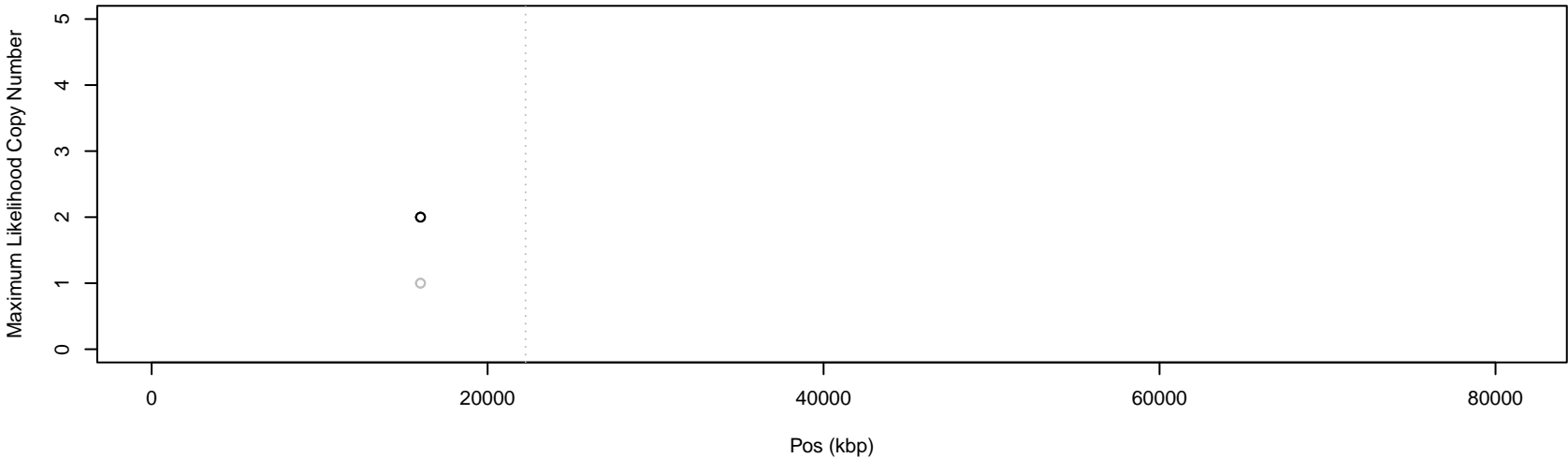
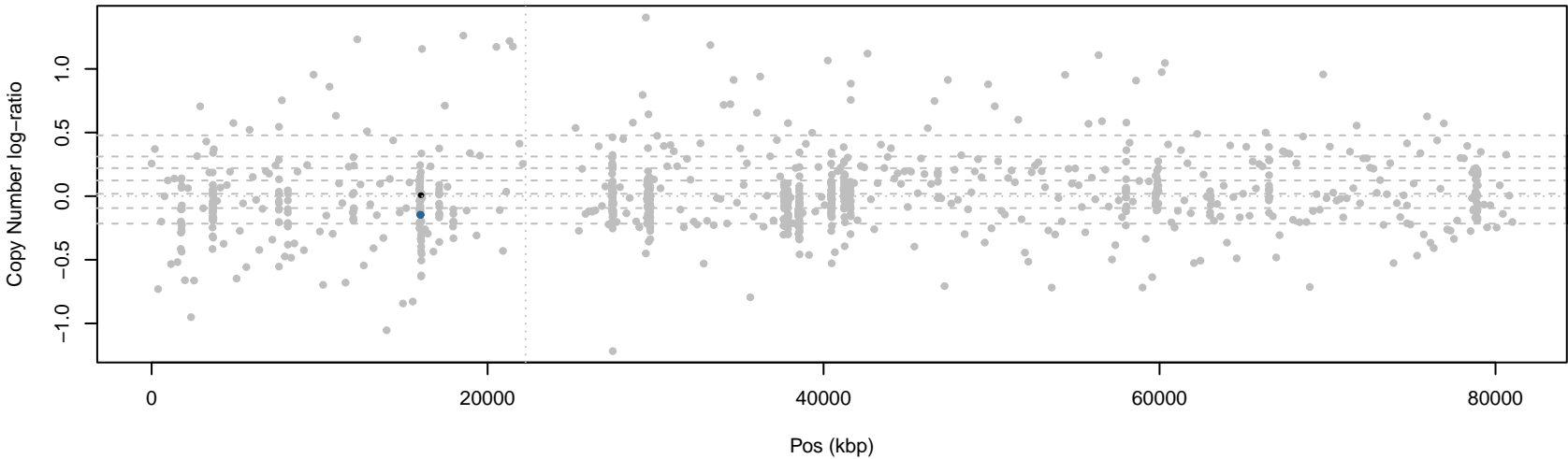
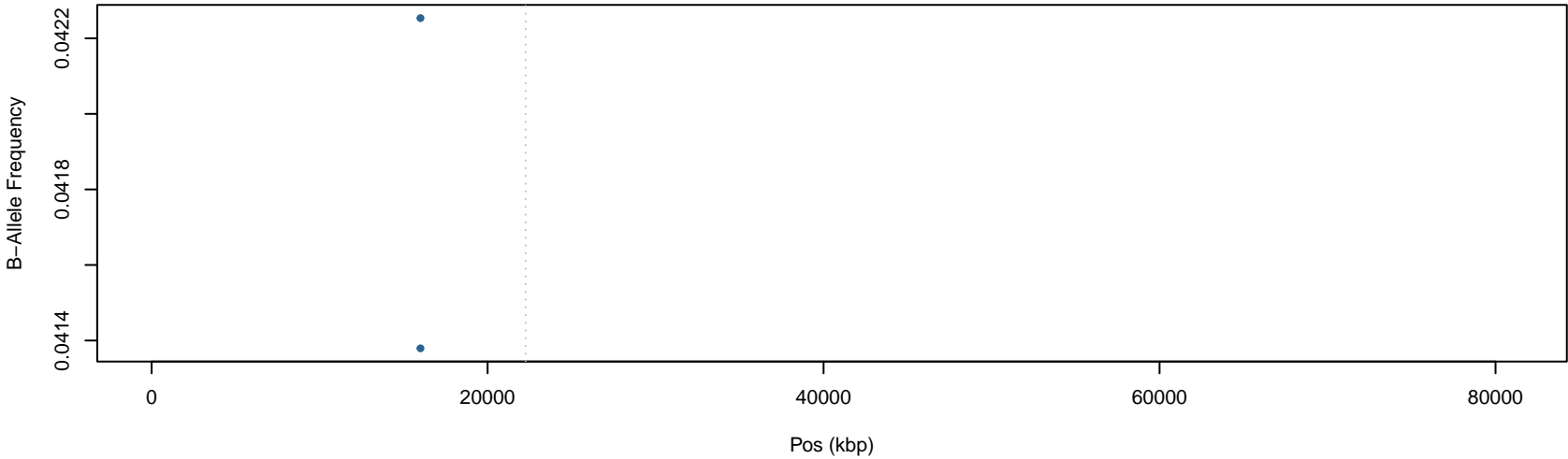
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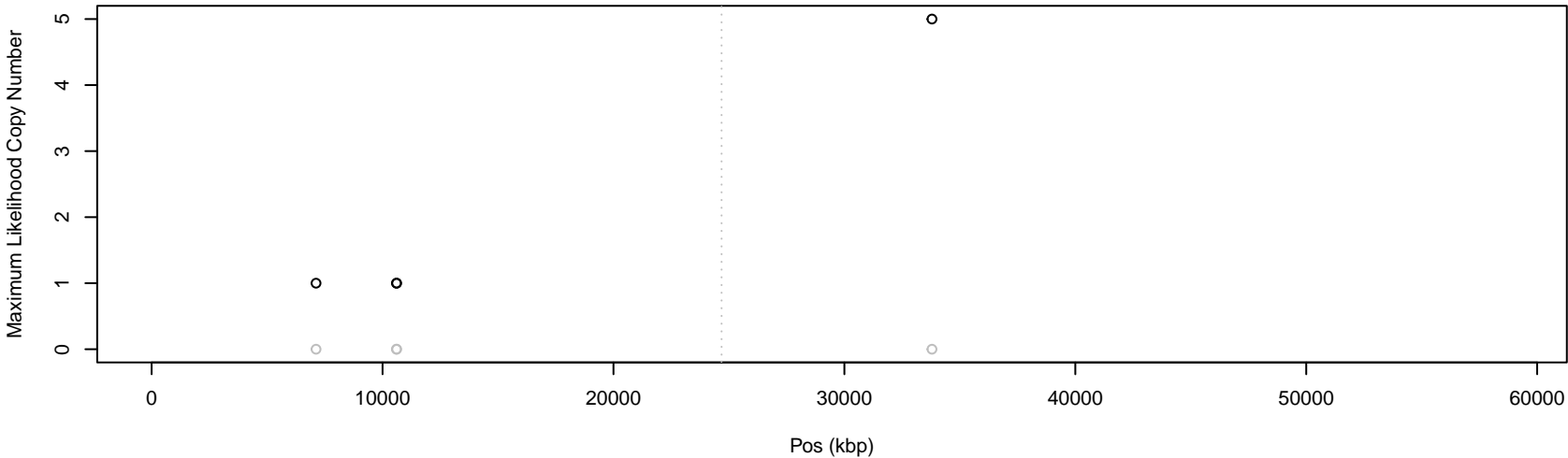
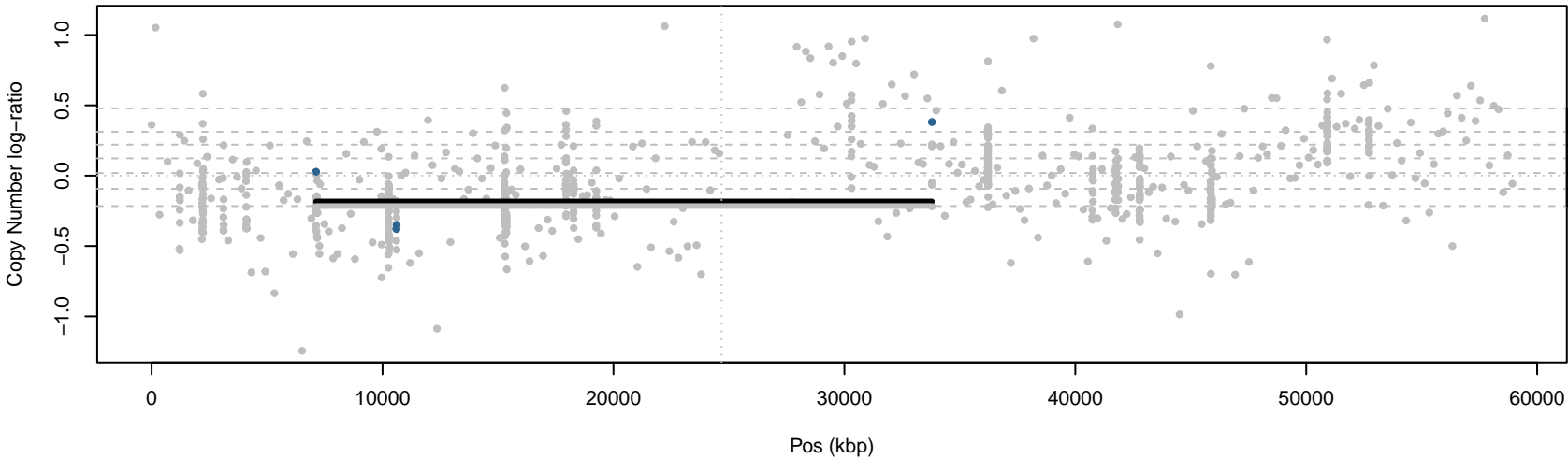
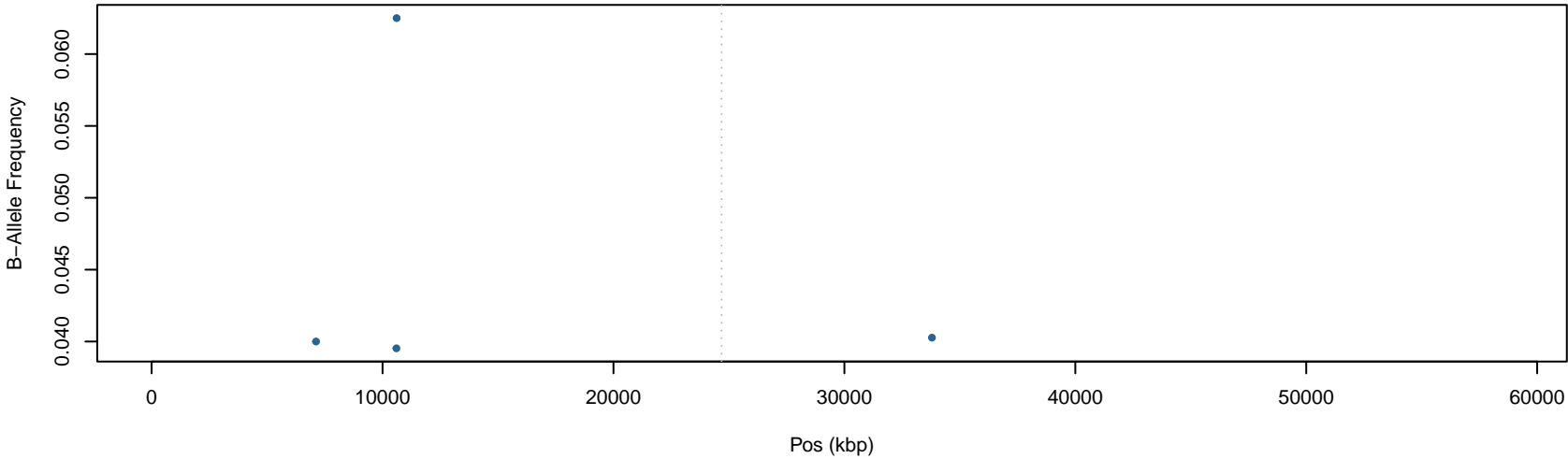
Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 16



Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 17



Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 19





Purity: 0.15 Tumor ploidy: 1.829 SNV log-likelihood: 310.32 GoF: 96% Mean coverage: 219;199 Chromosome: 20

