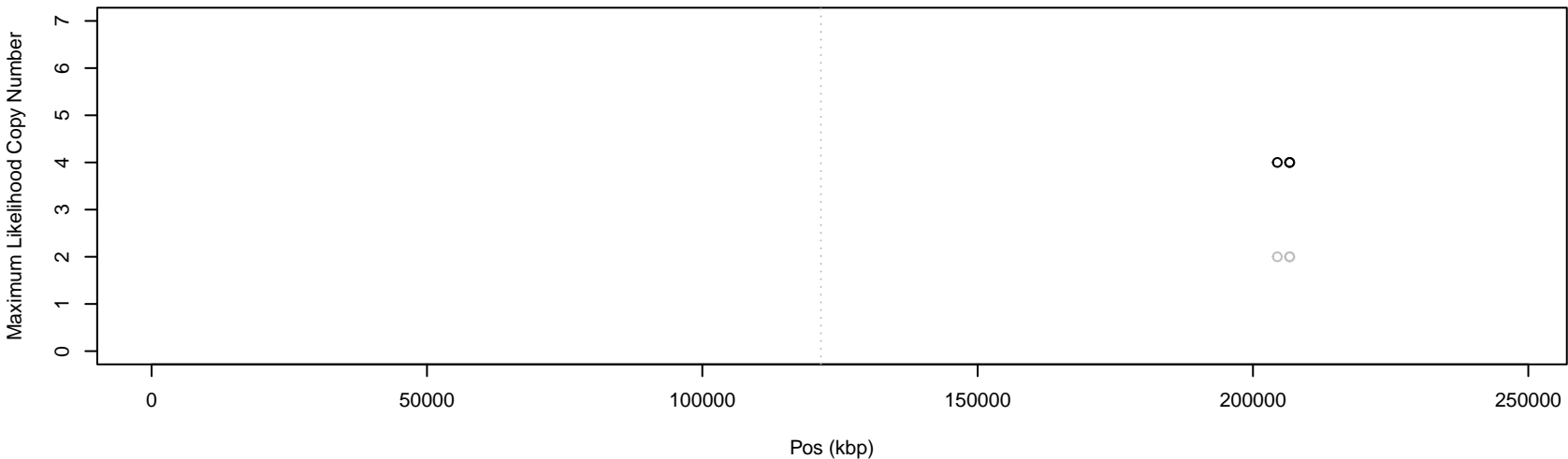
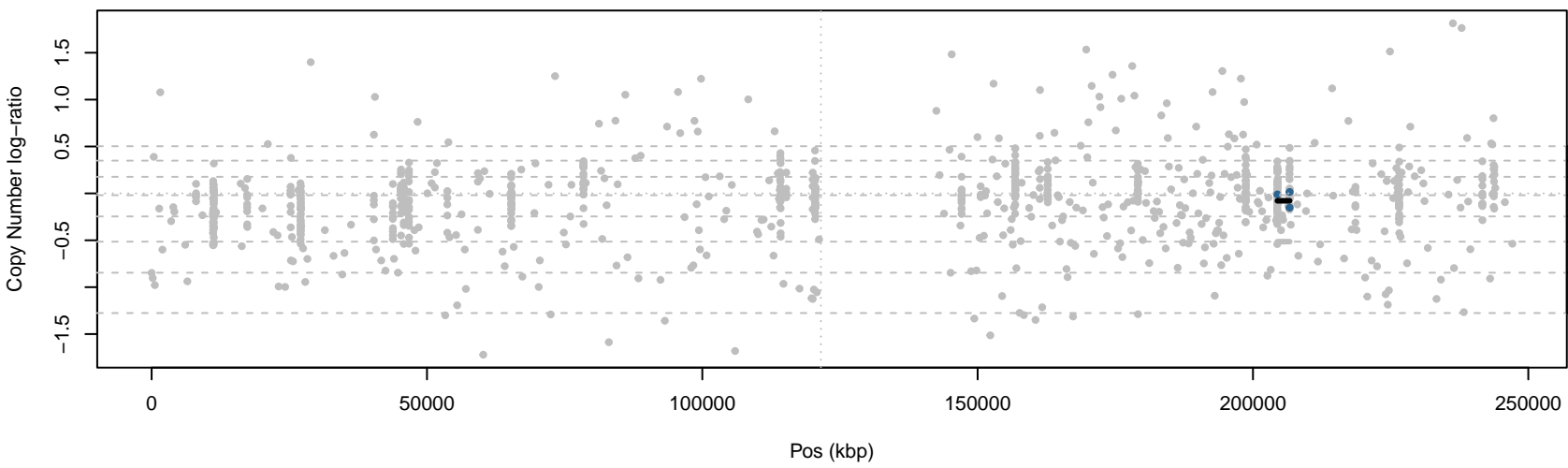
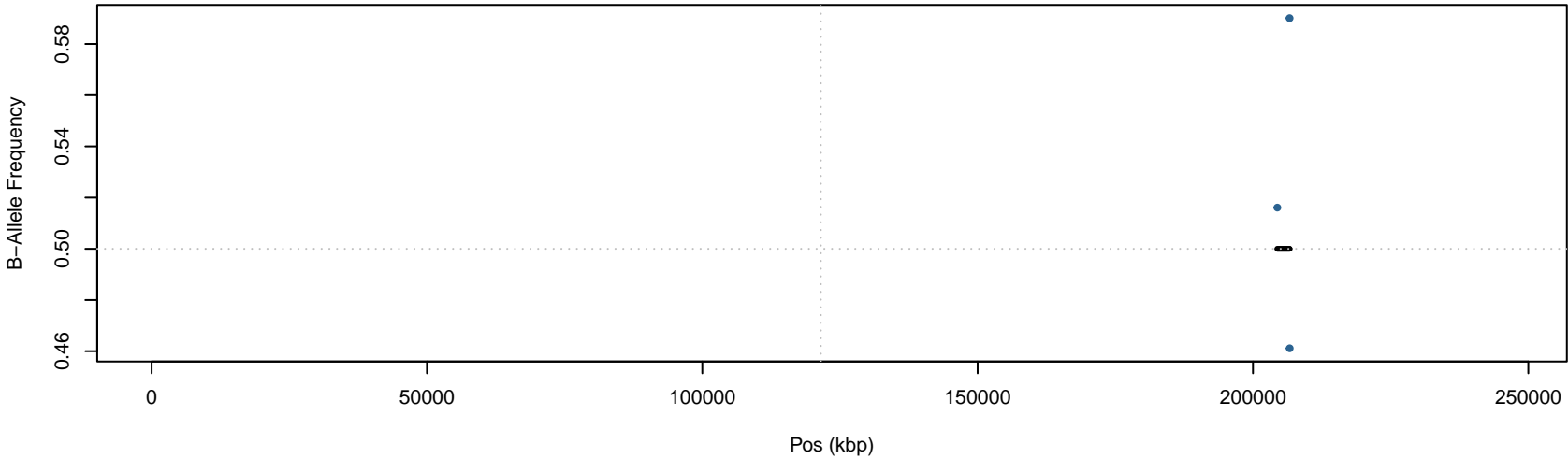
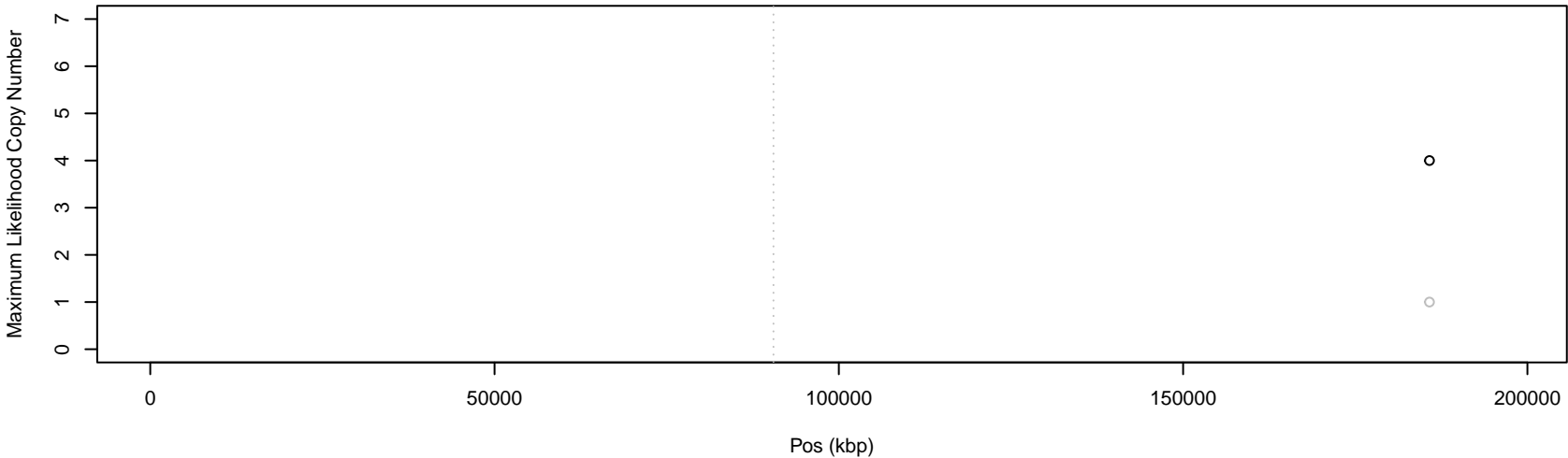
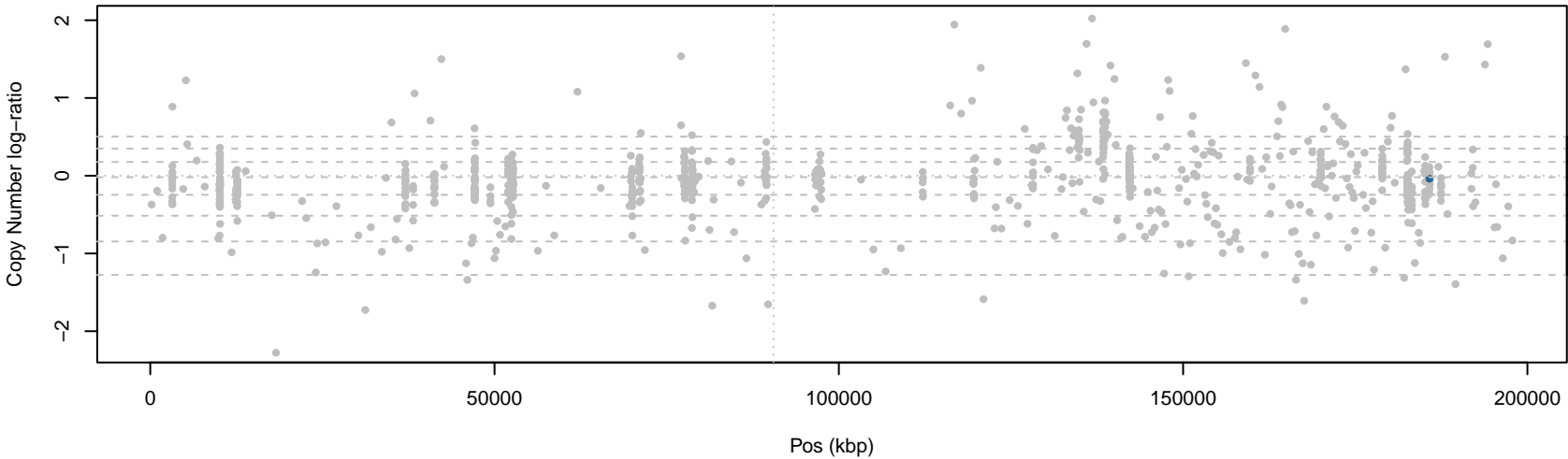
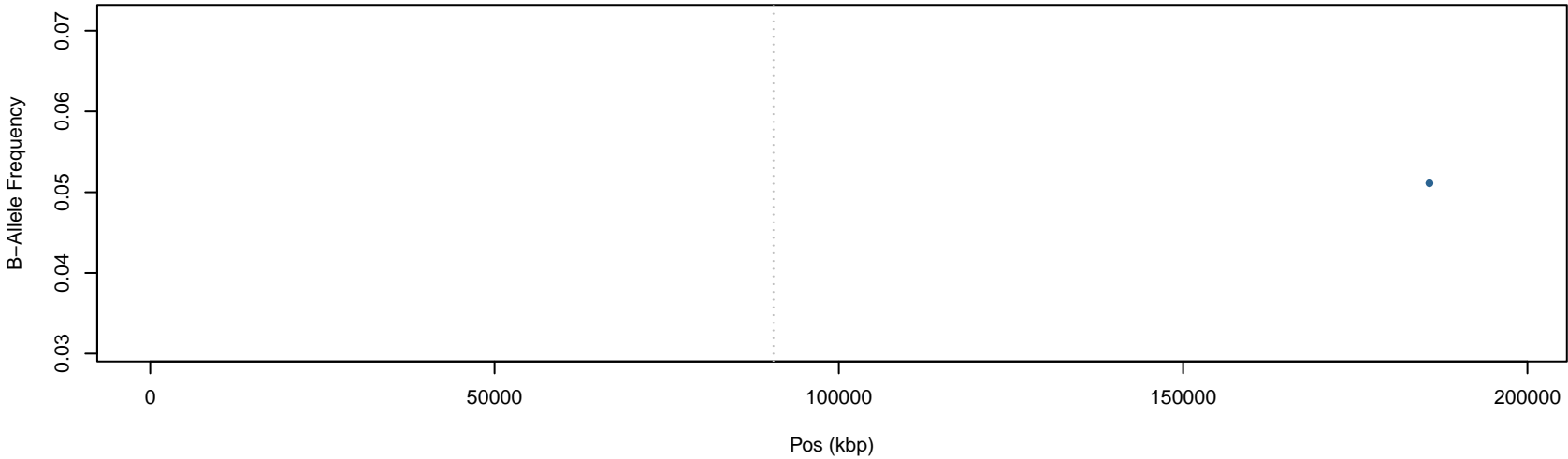


Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 1

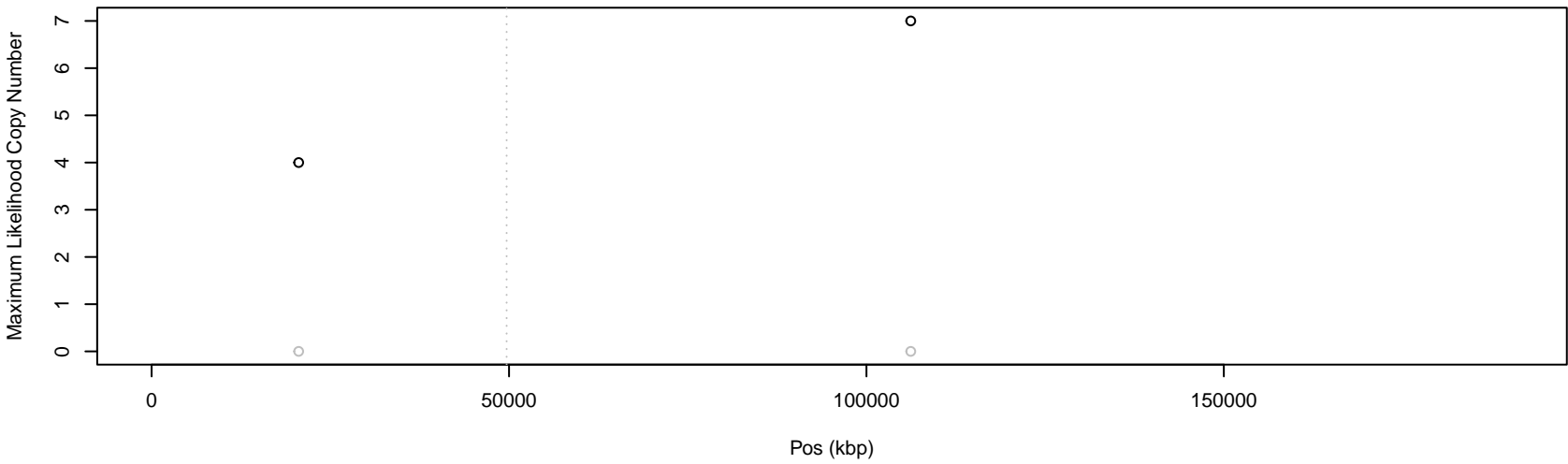
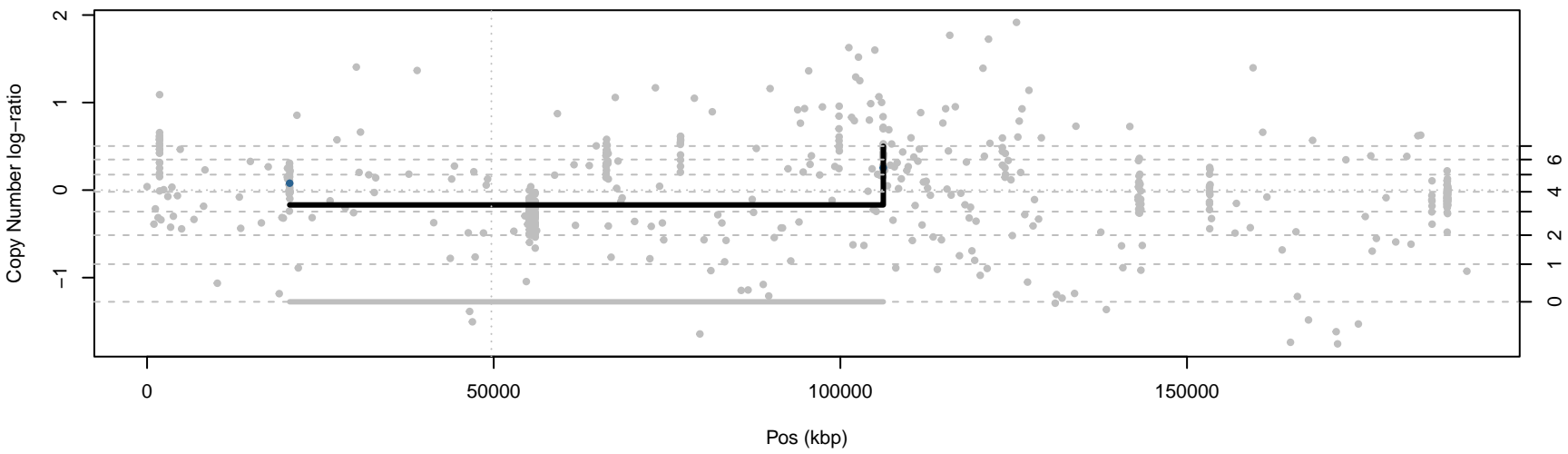
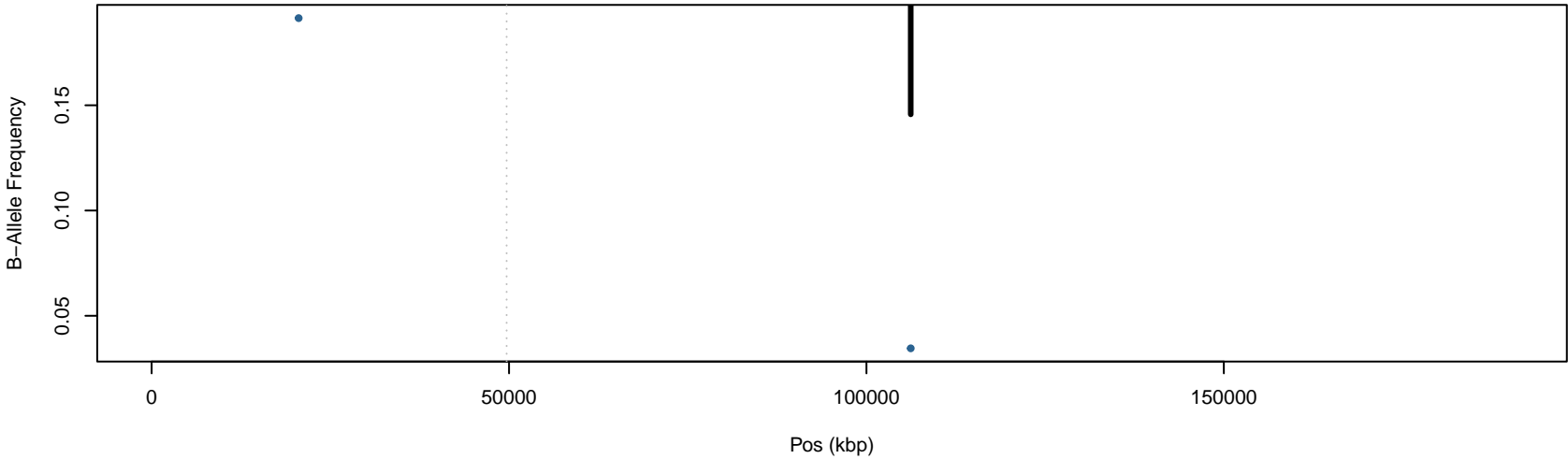




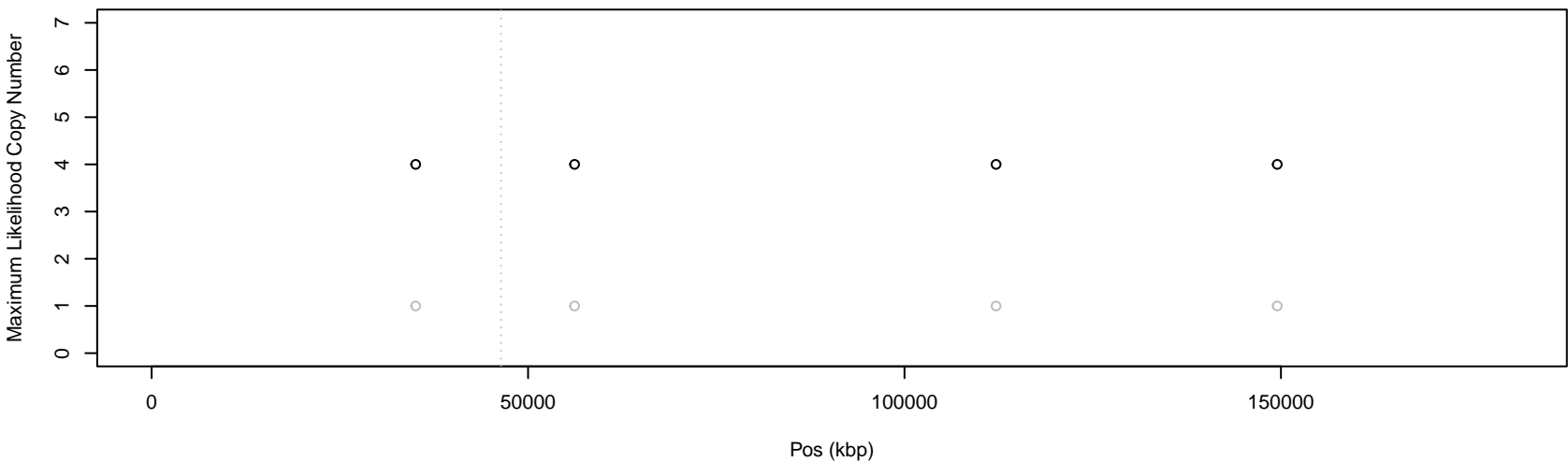
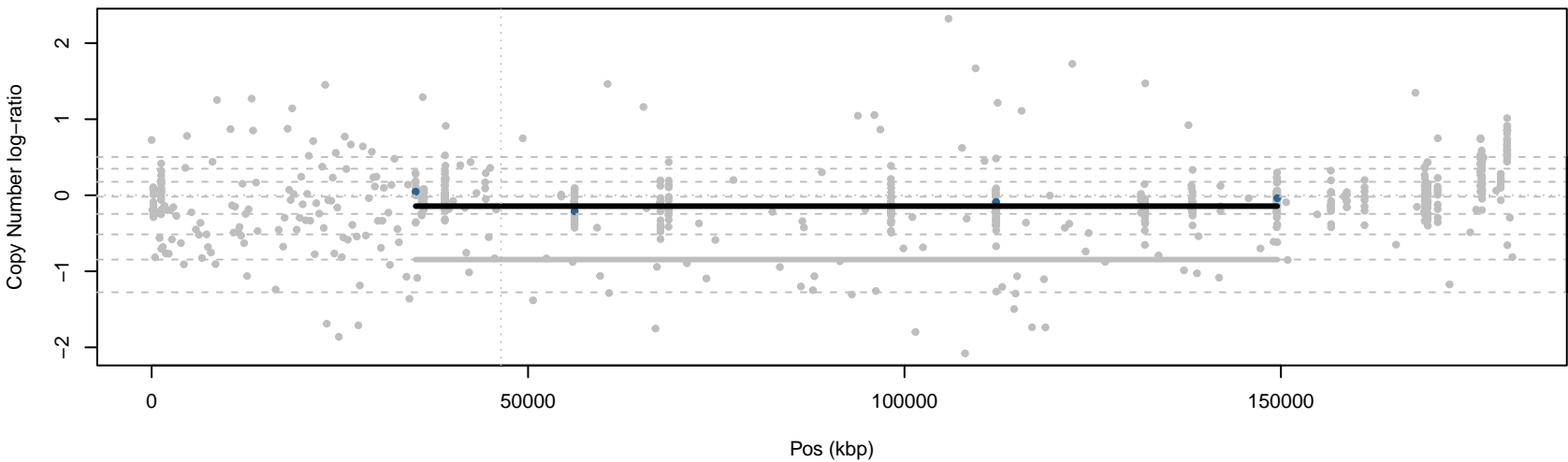
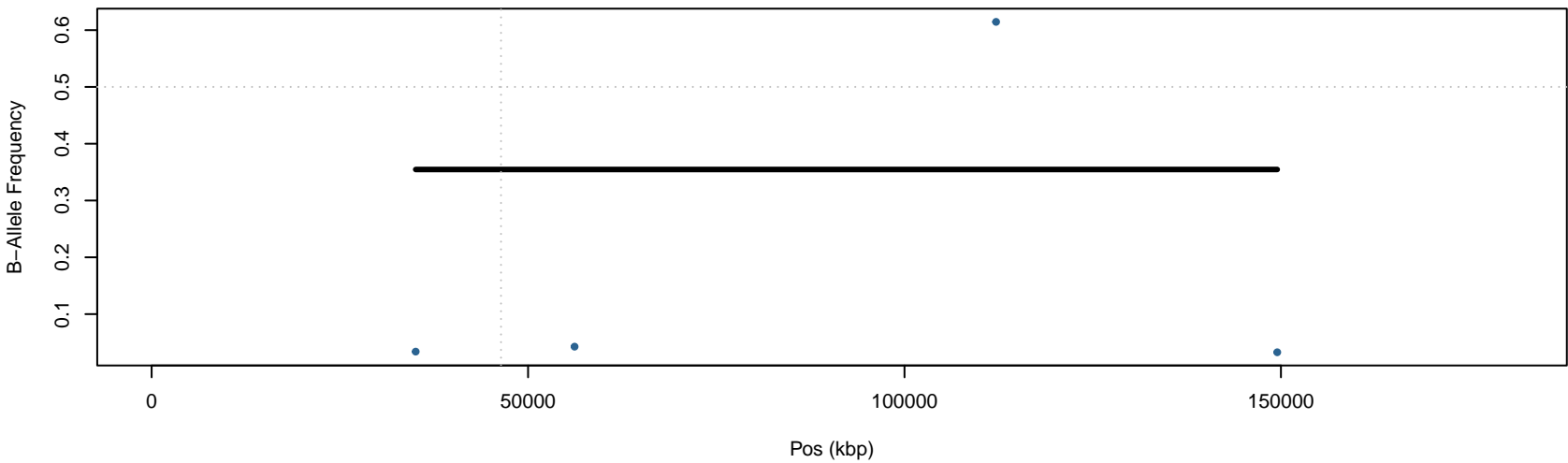
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 3



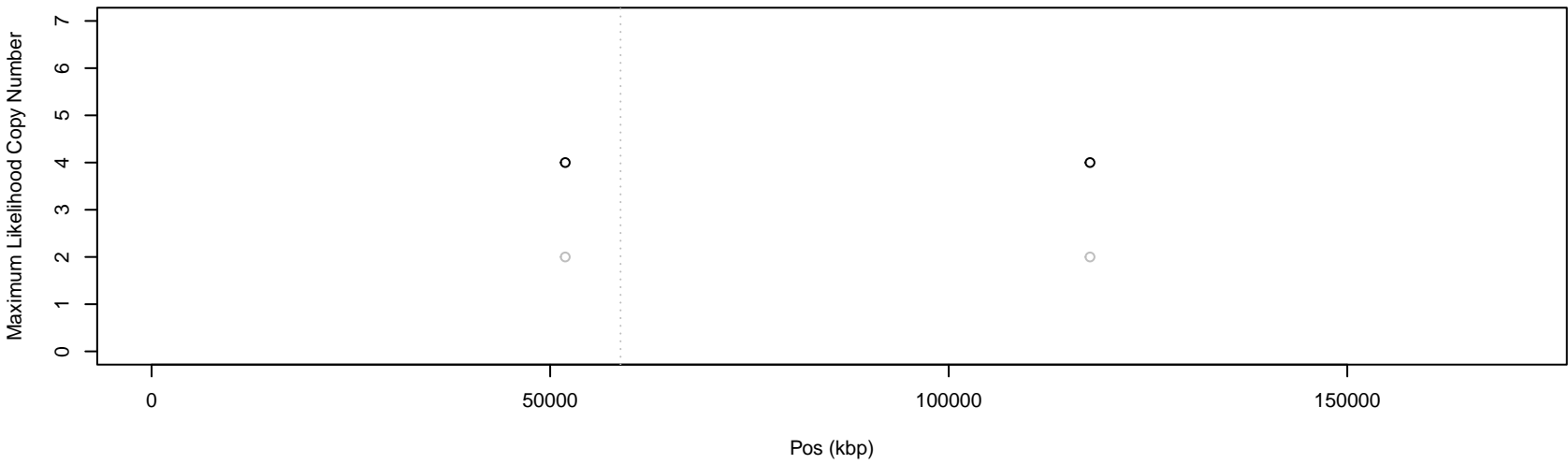
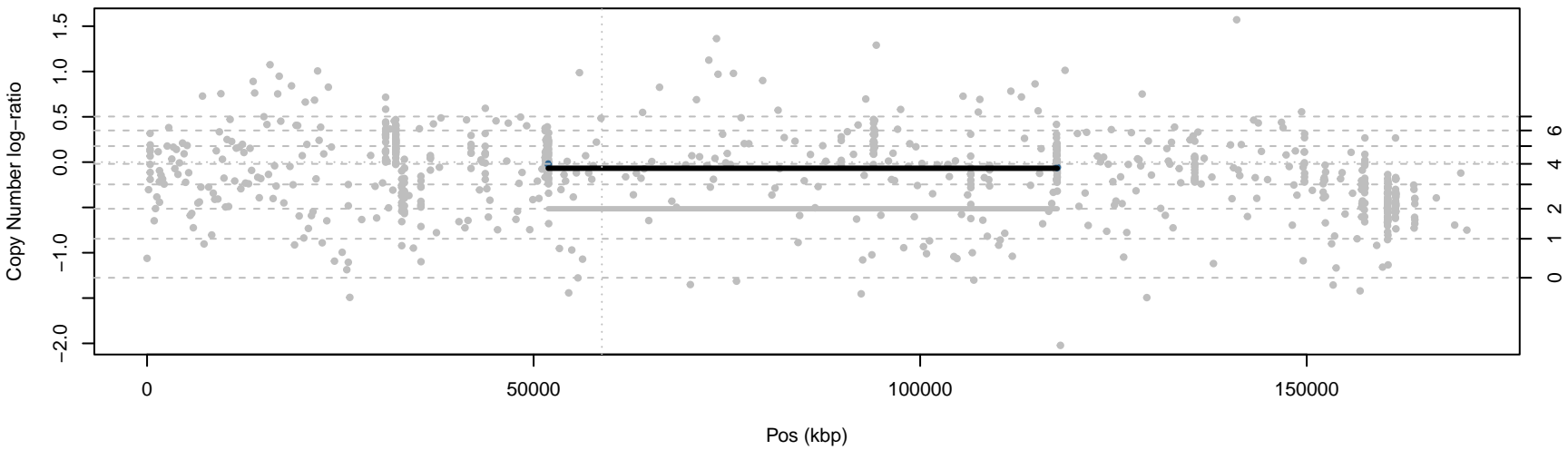
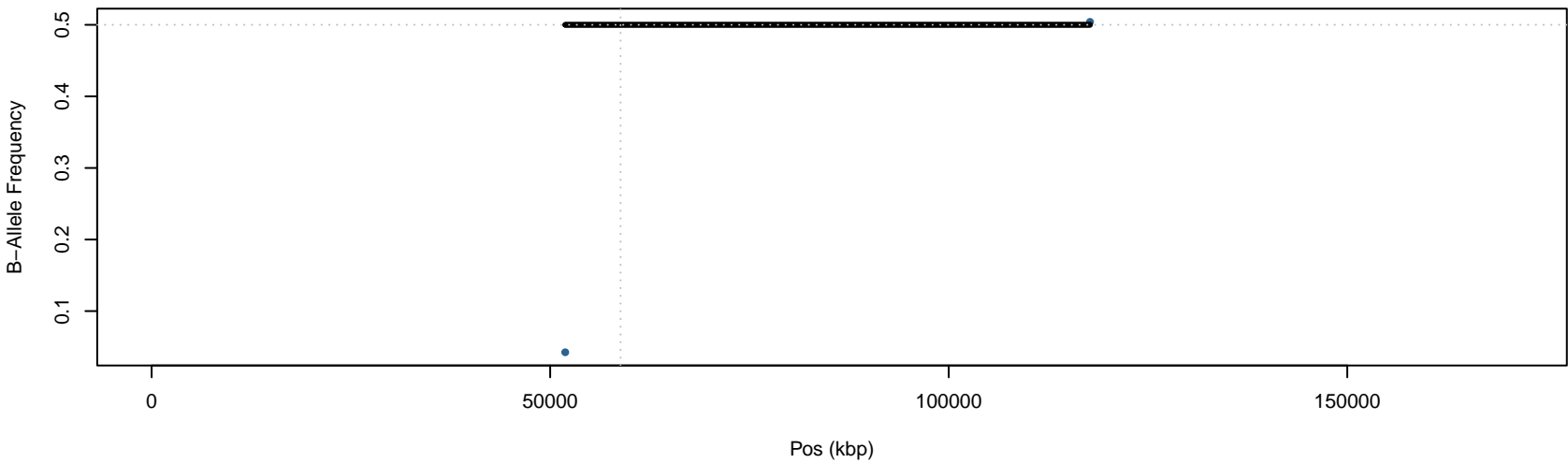
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 4



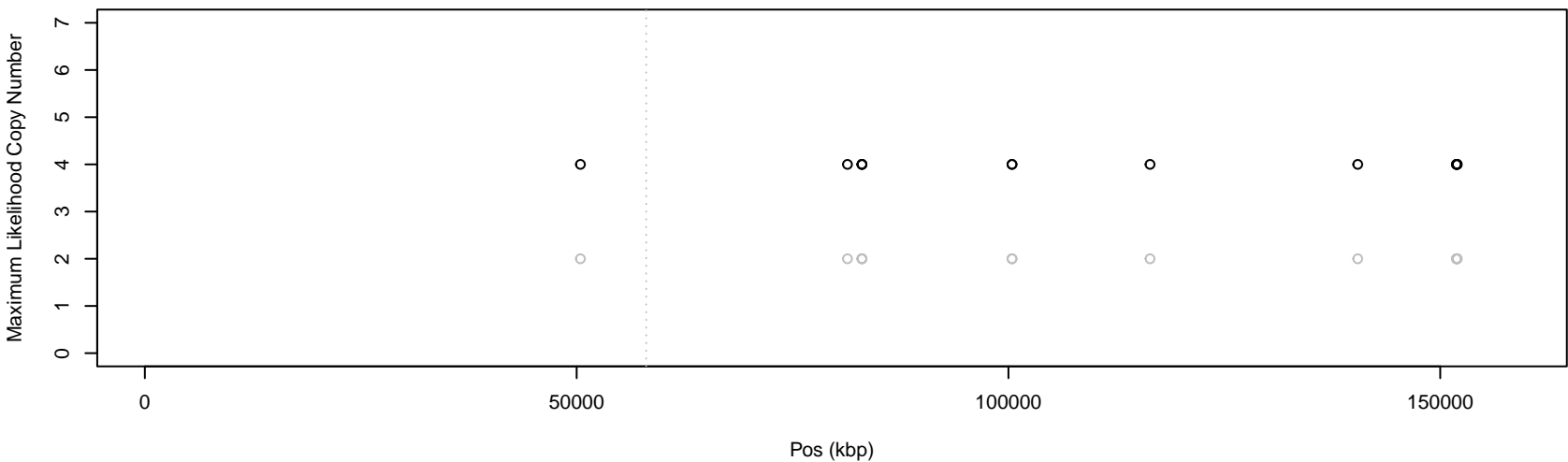
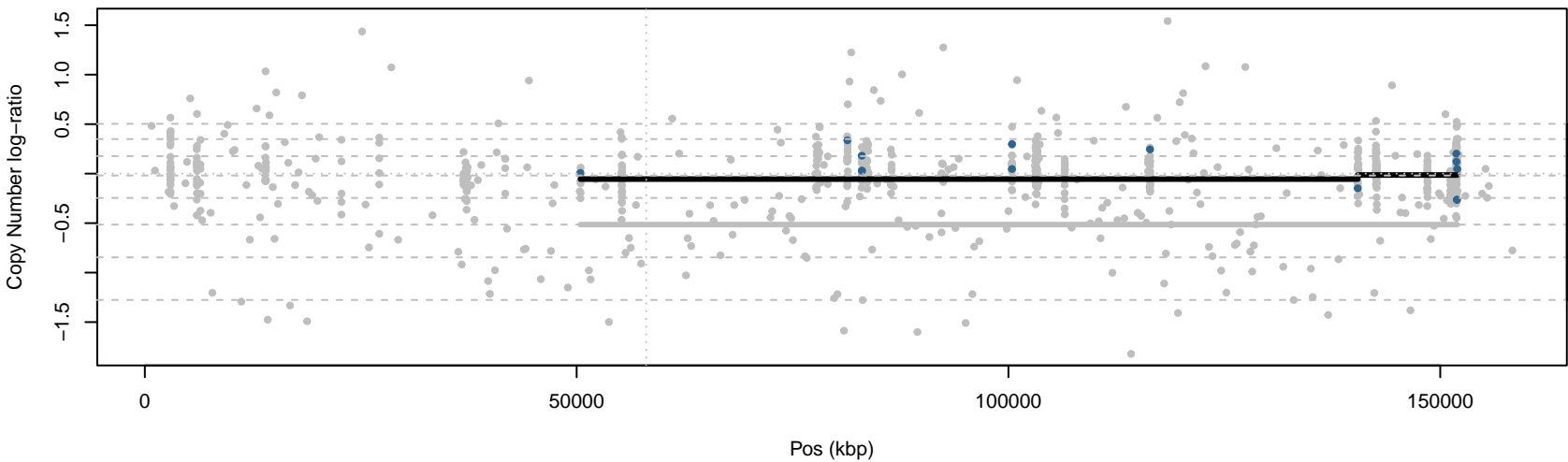
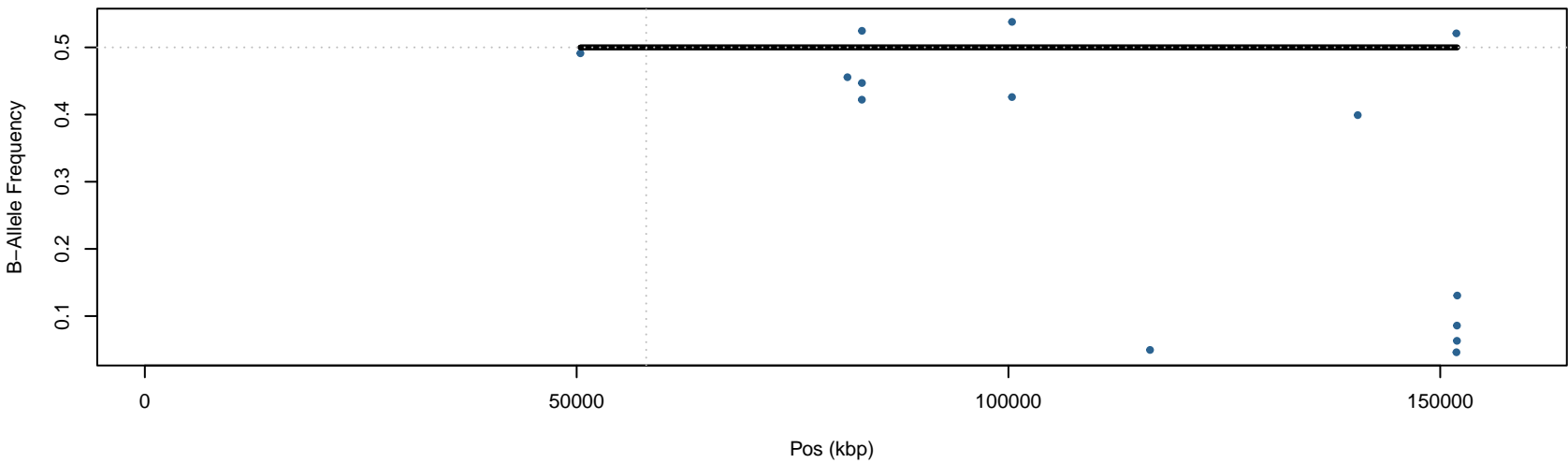
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 5



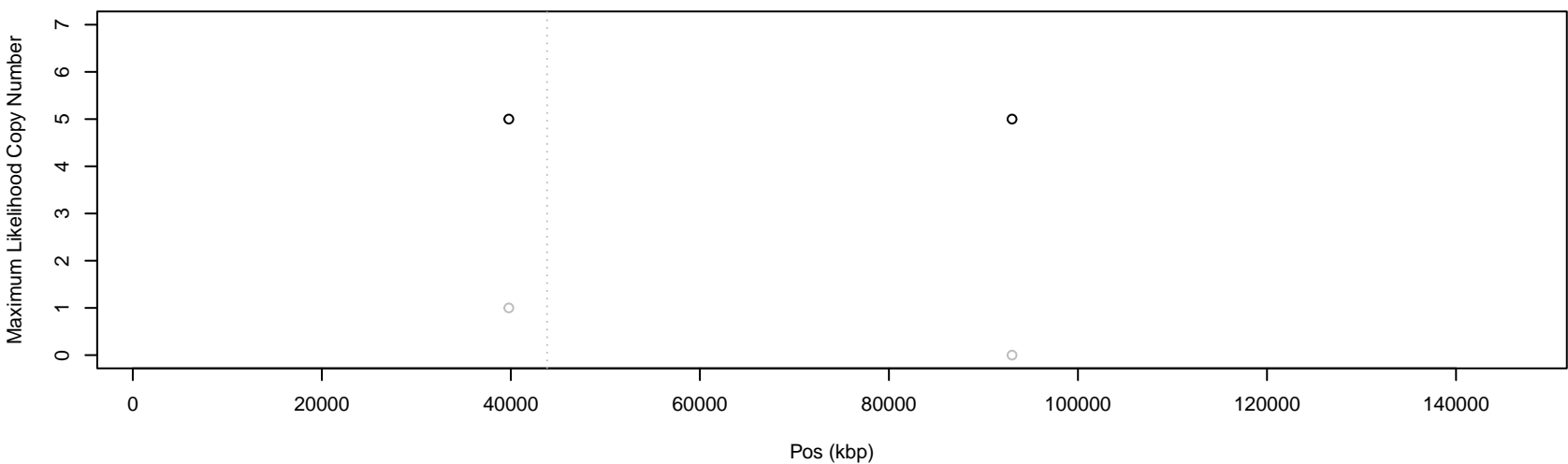
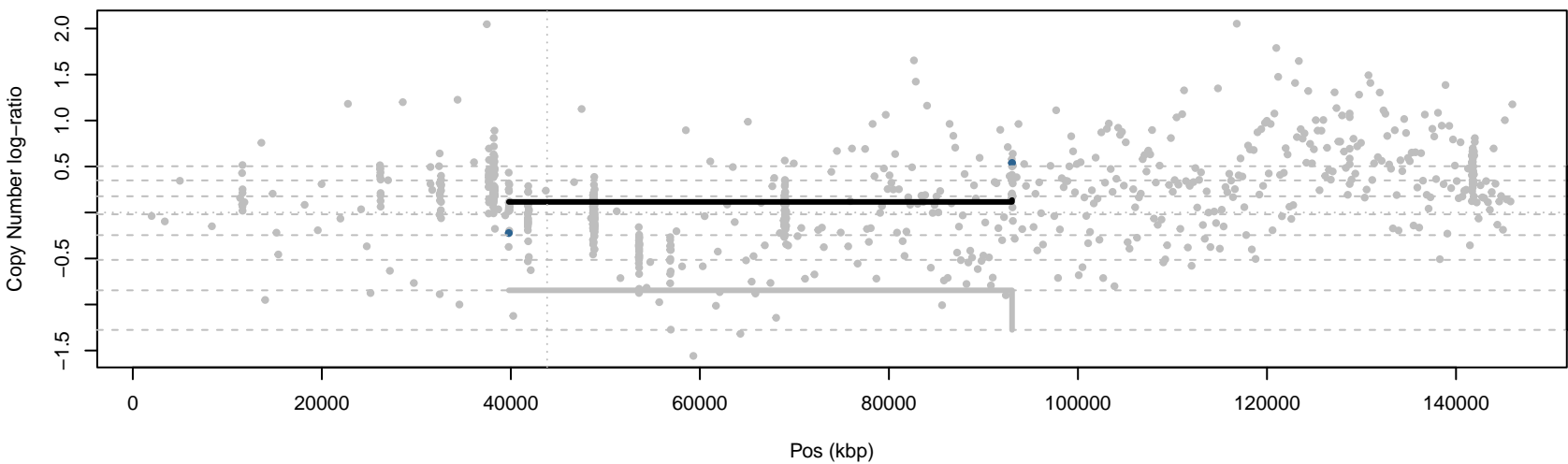
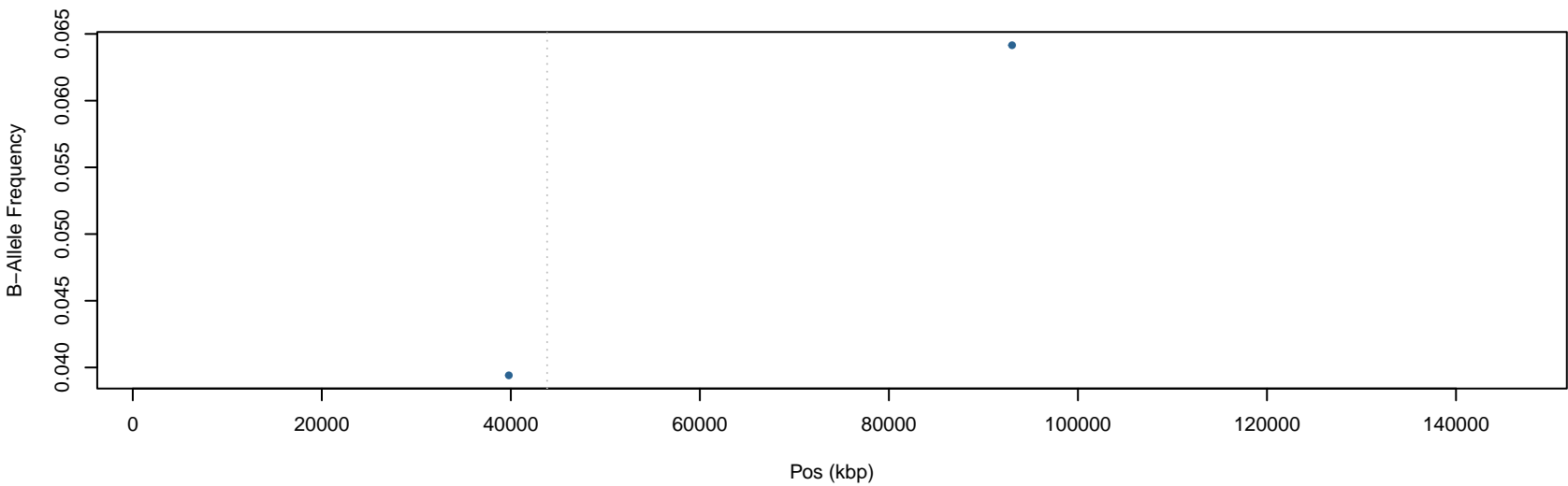
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 6



Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 7

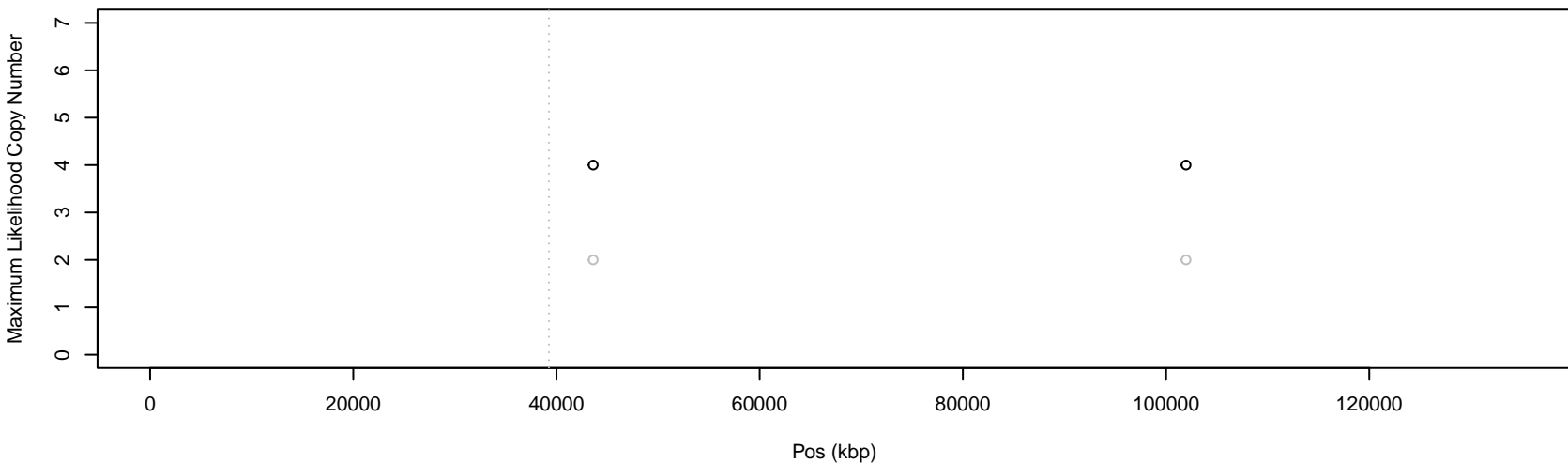
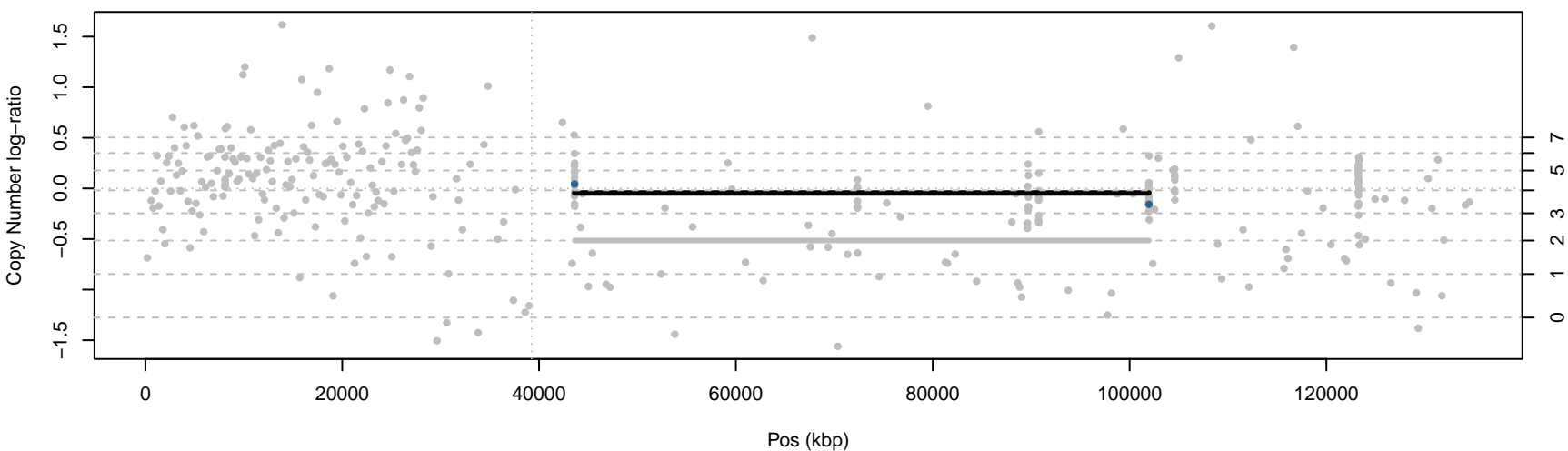
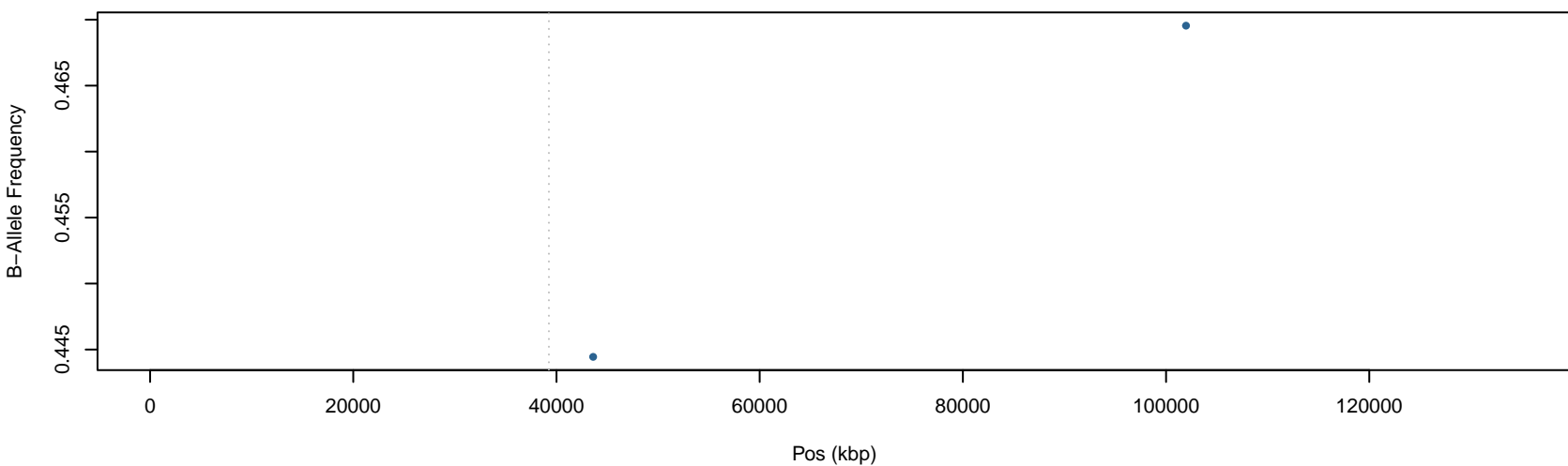


Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710;524 Chromosome: 8

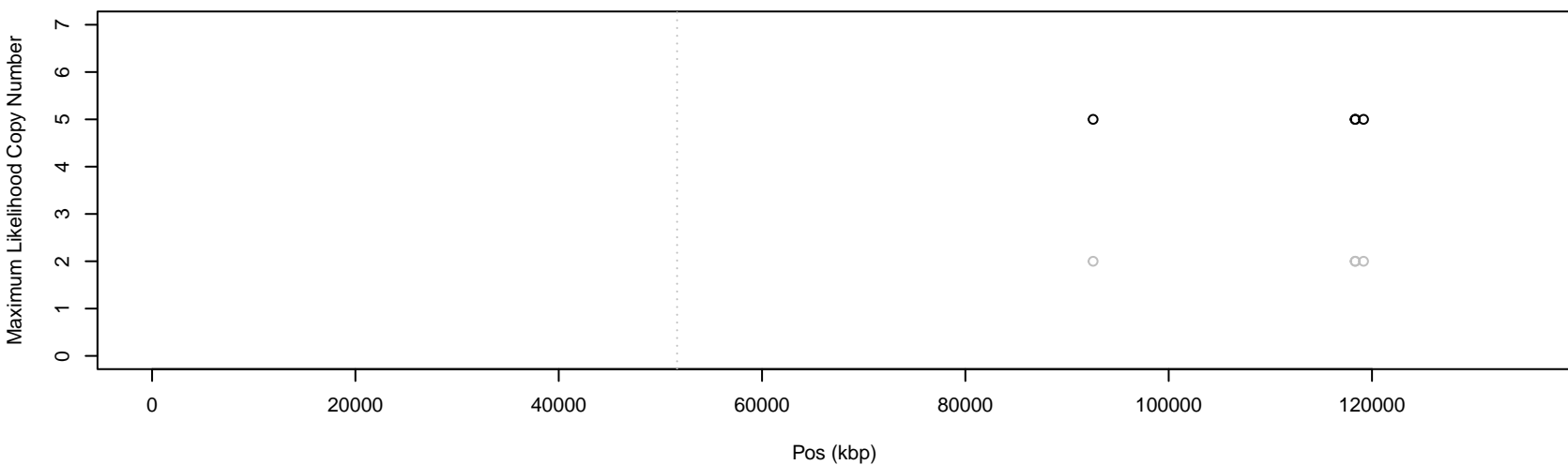
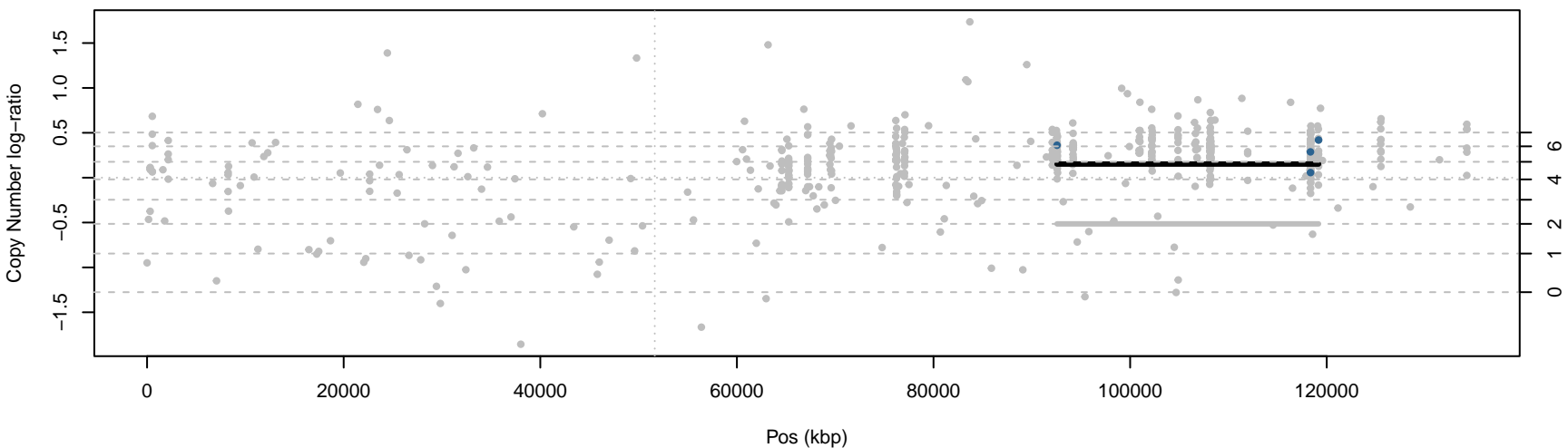
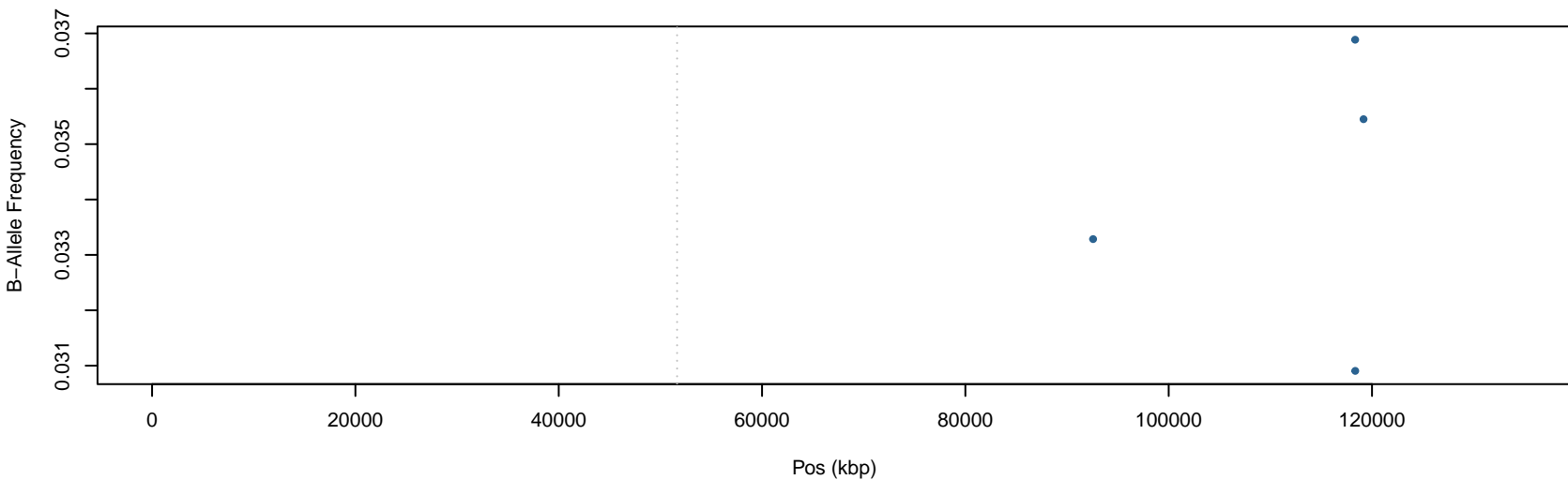




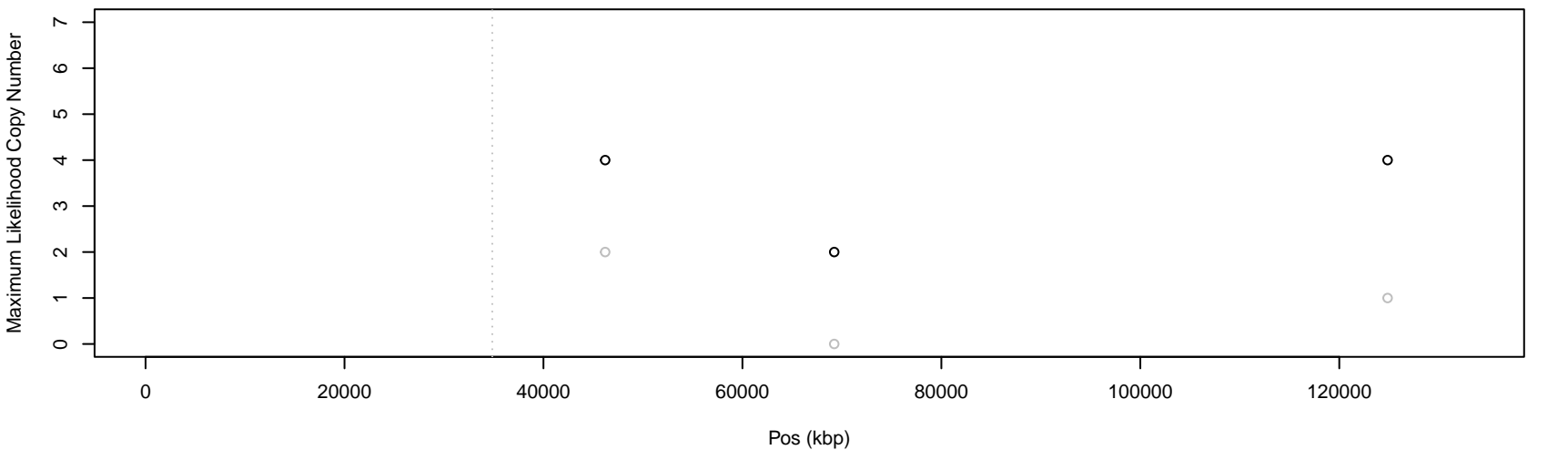
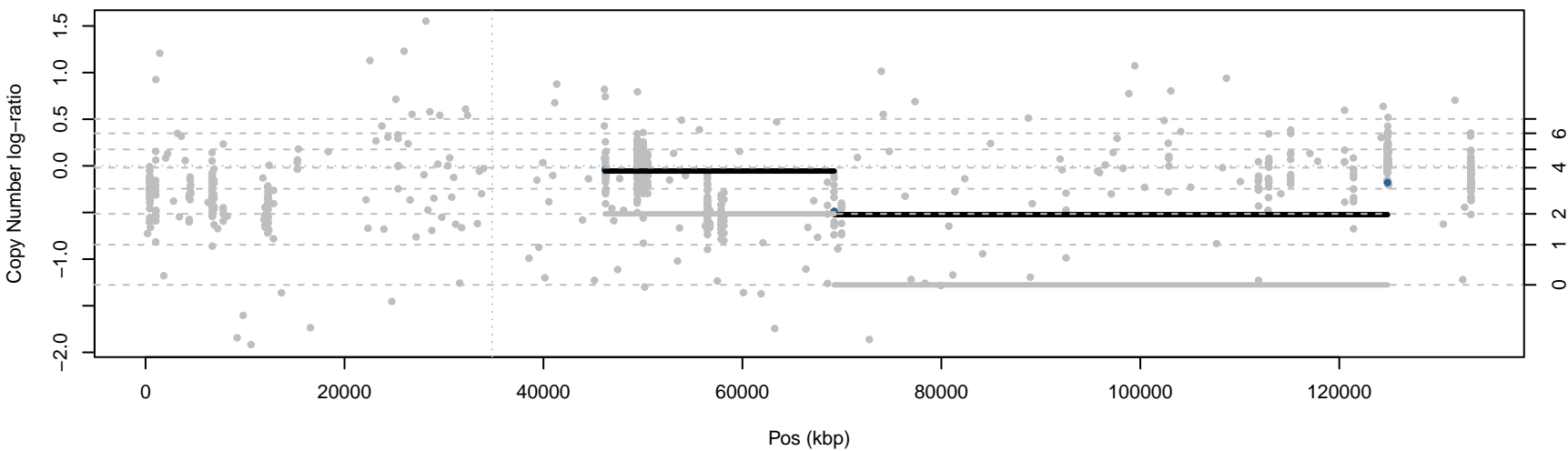
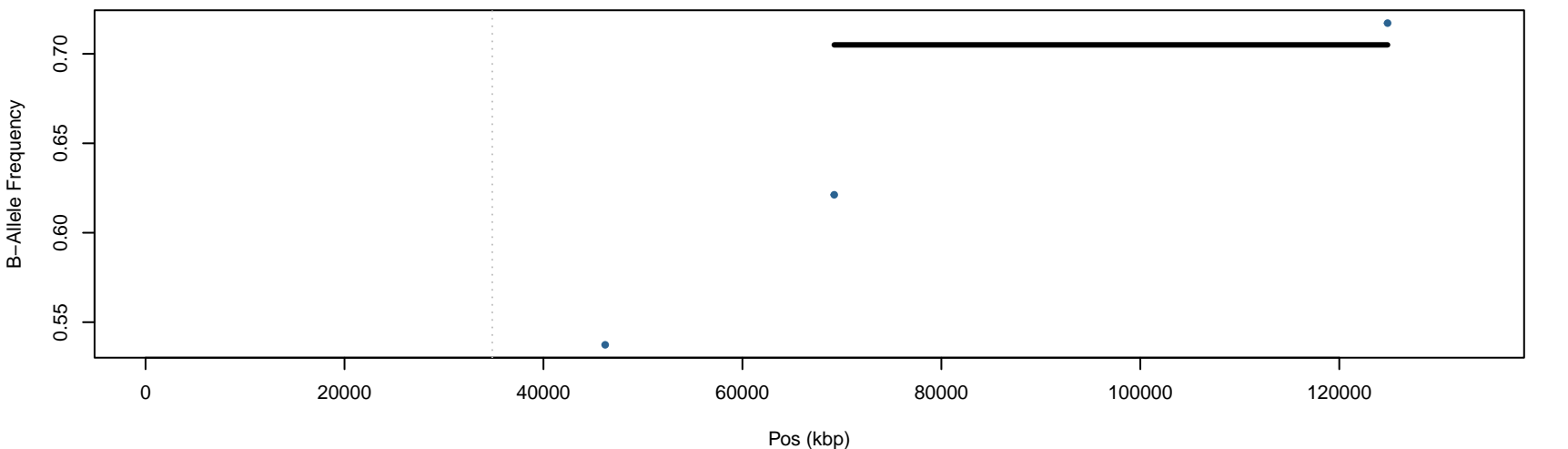
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 10



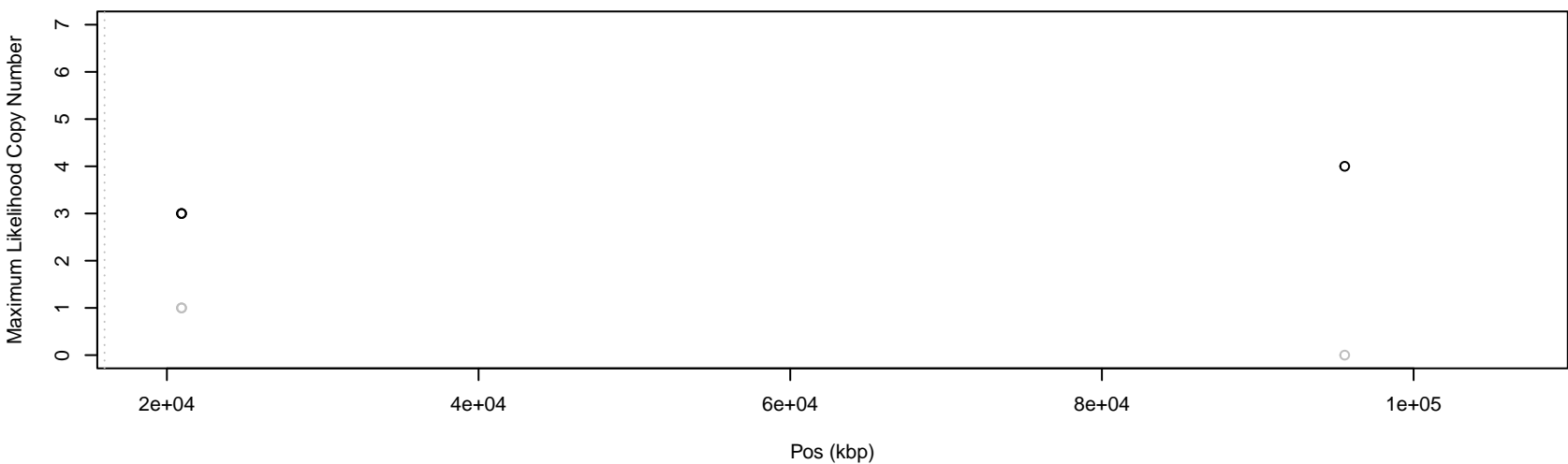
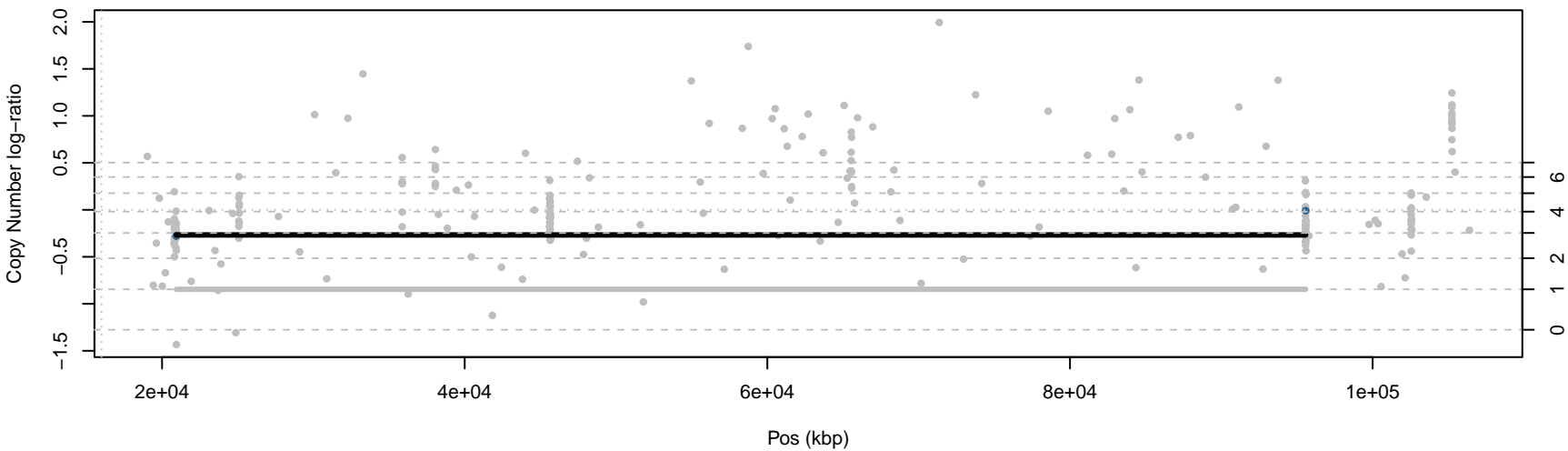
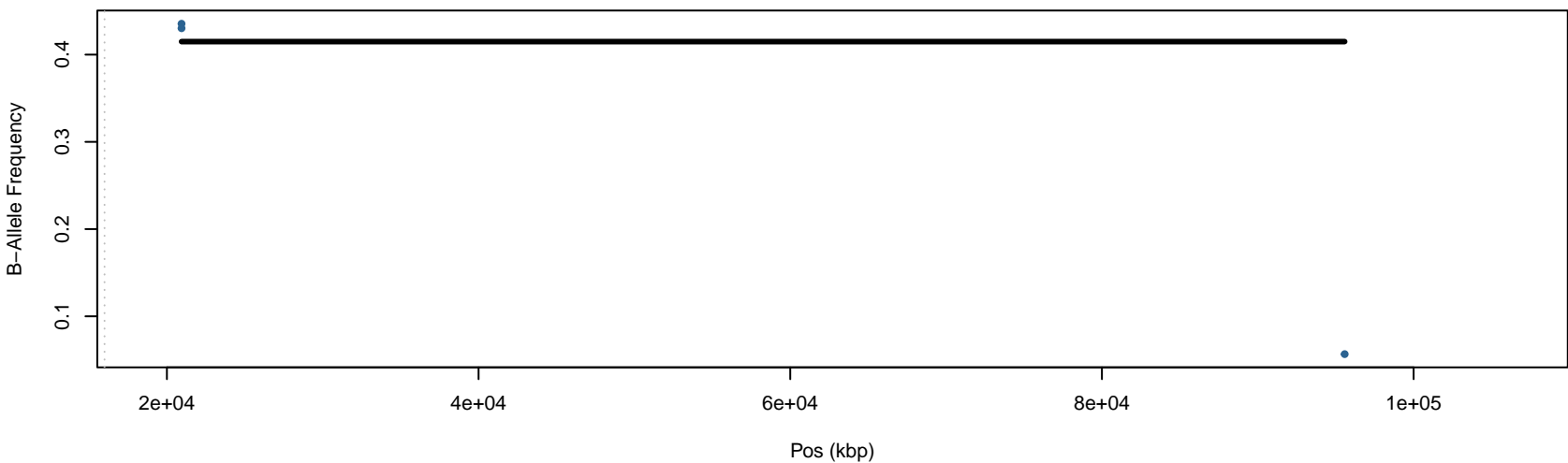
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 11



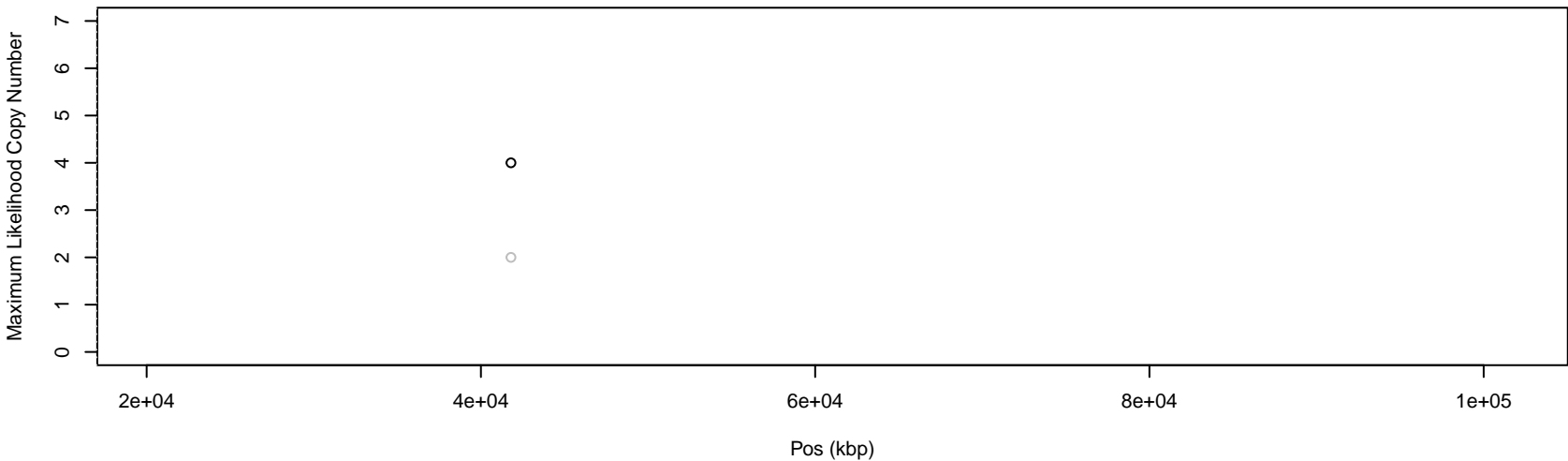
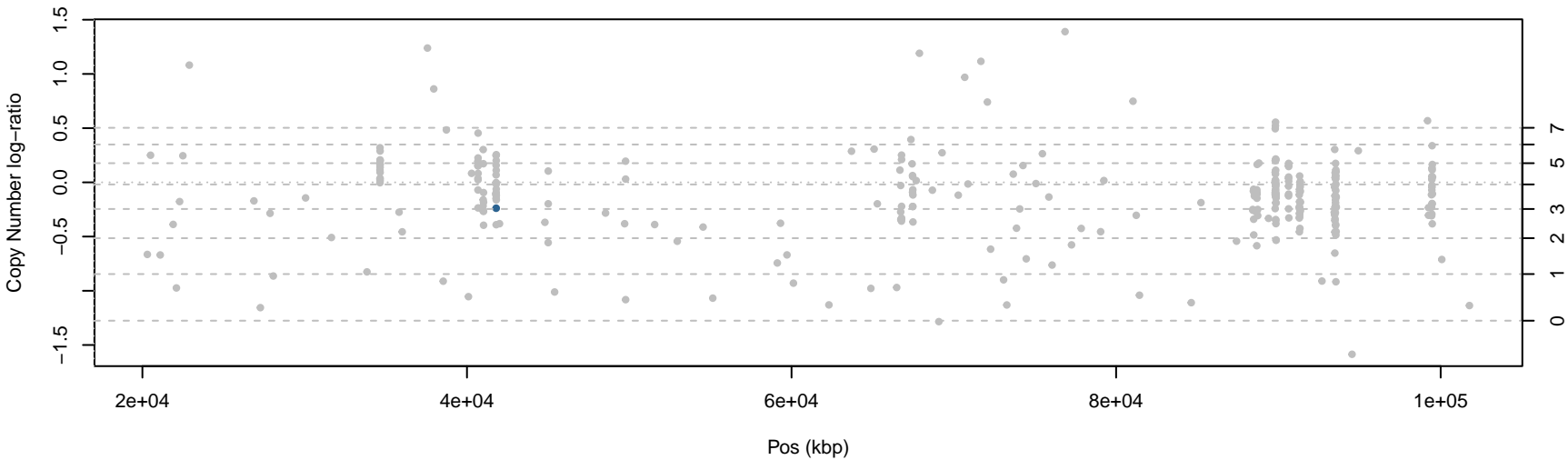
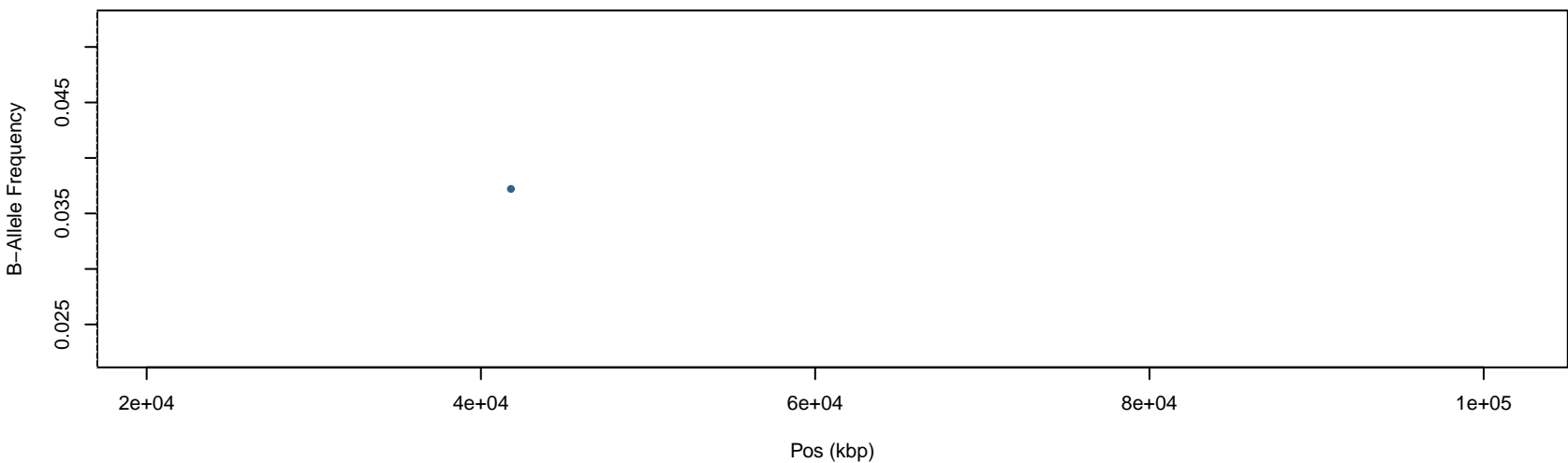
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 12



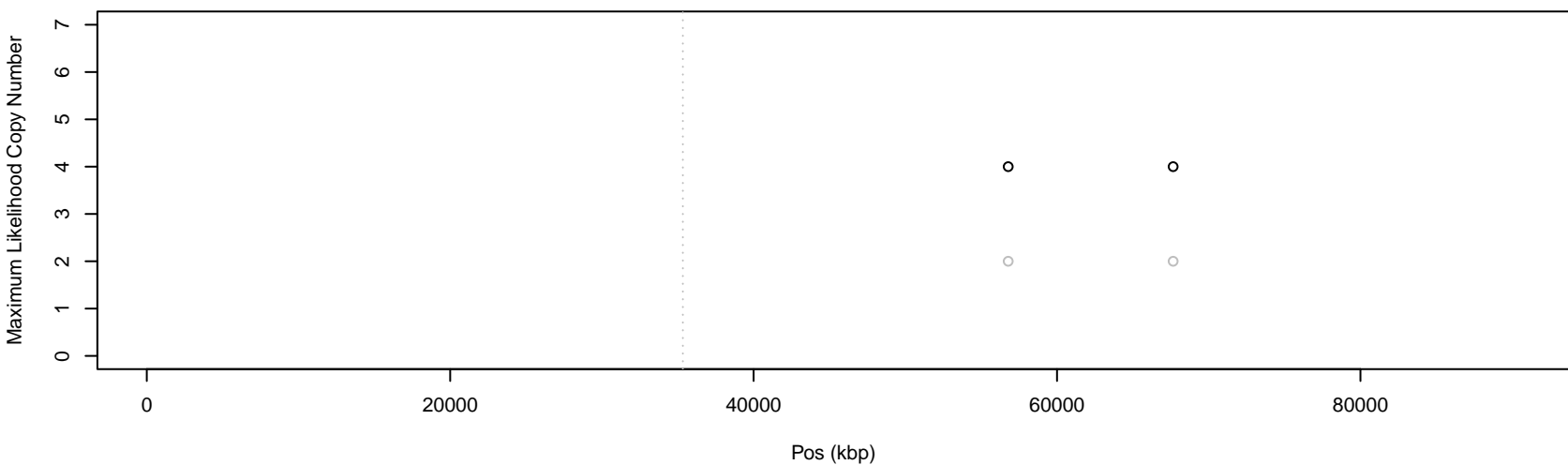
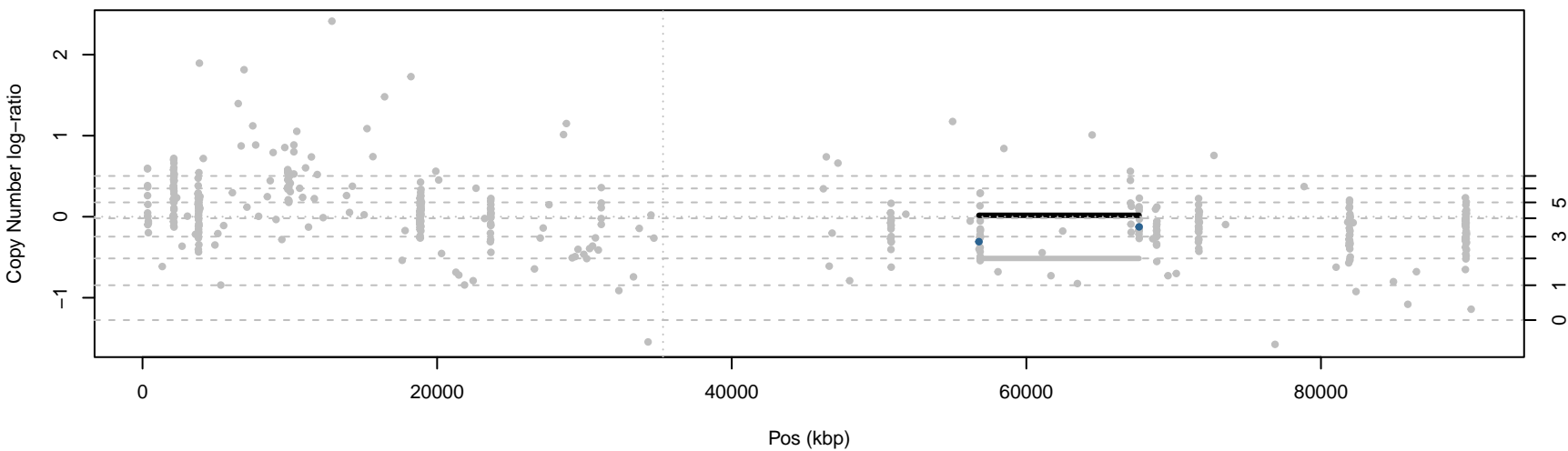
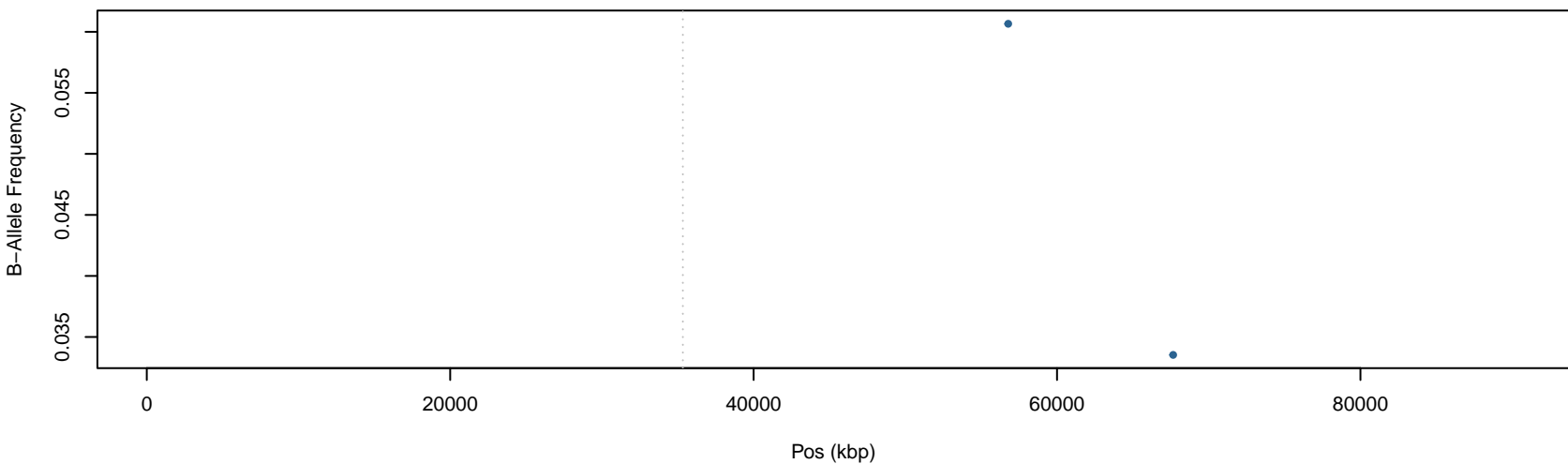
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 14



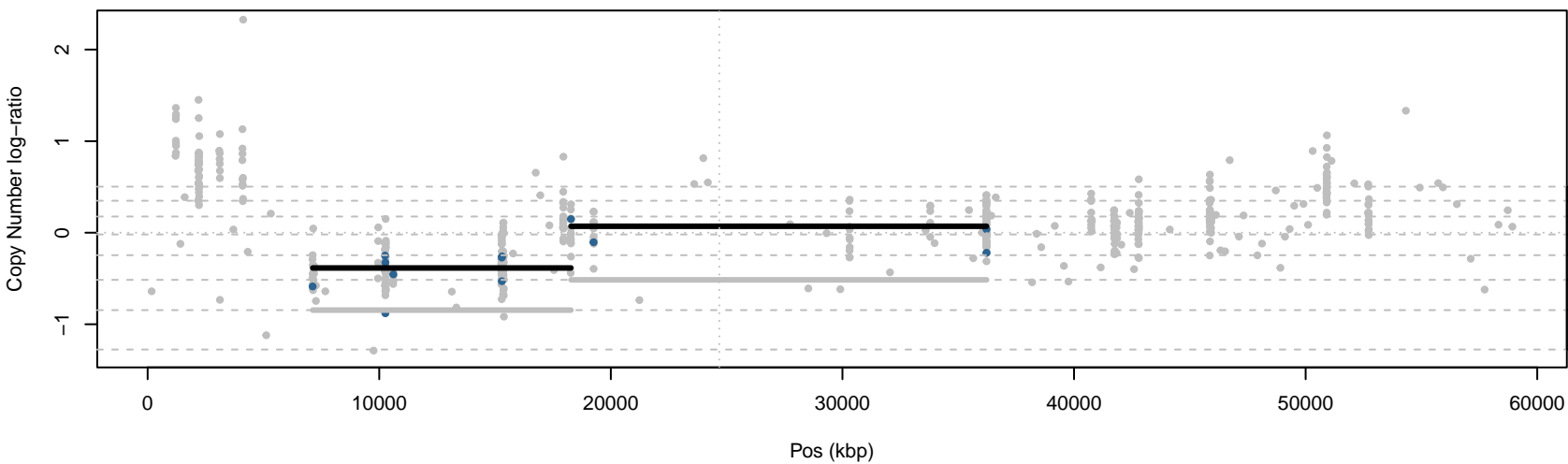
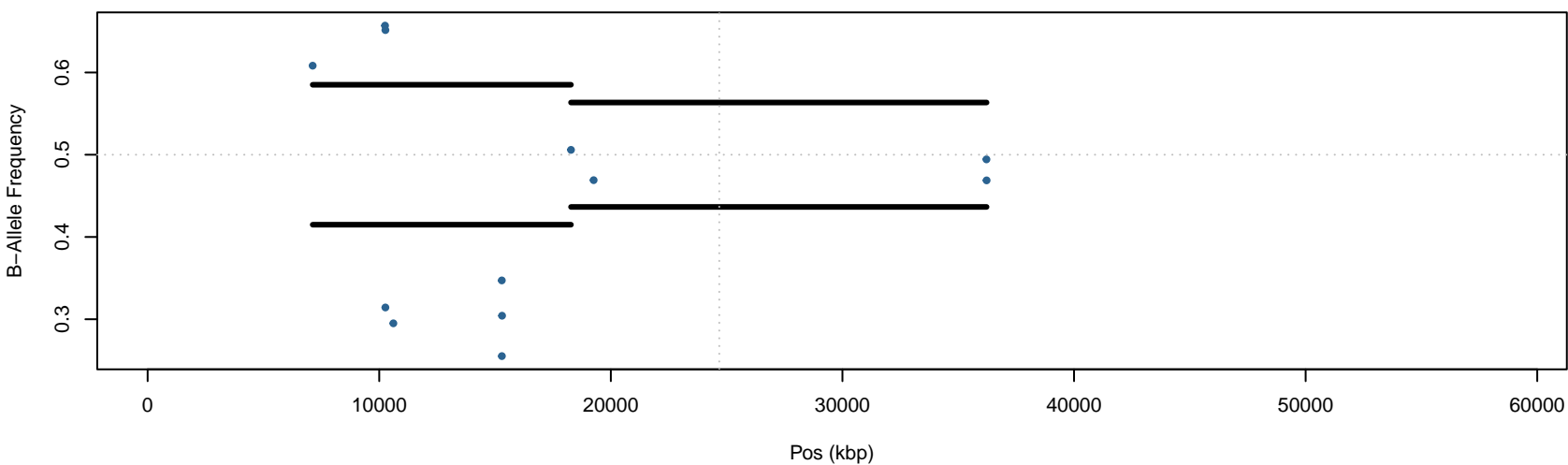
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 15



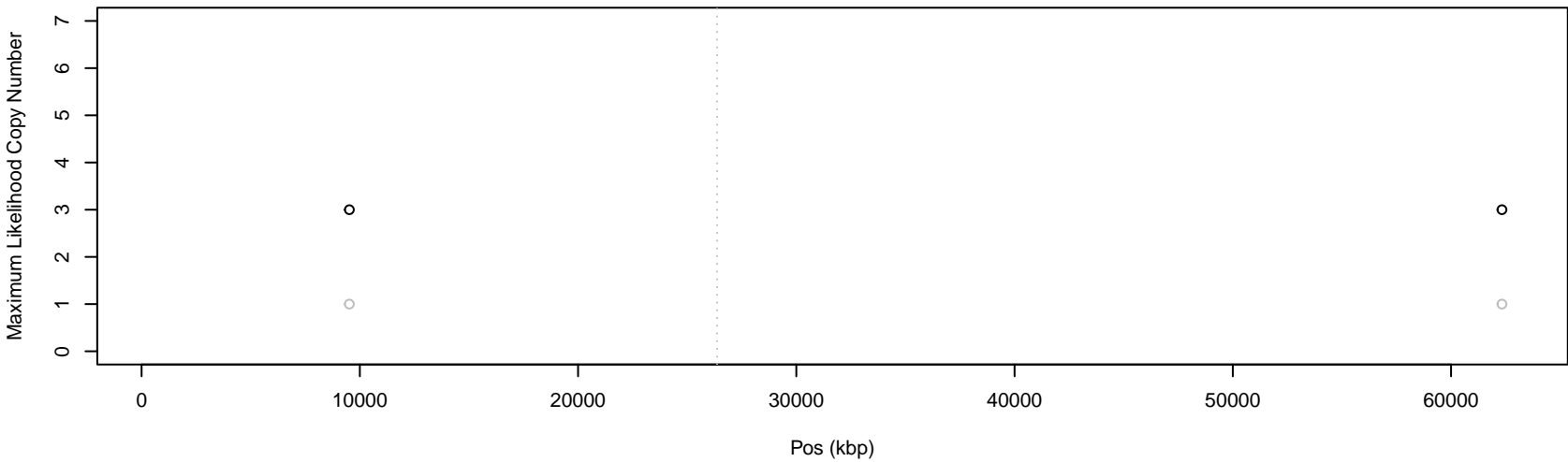
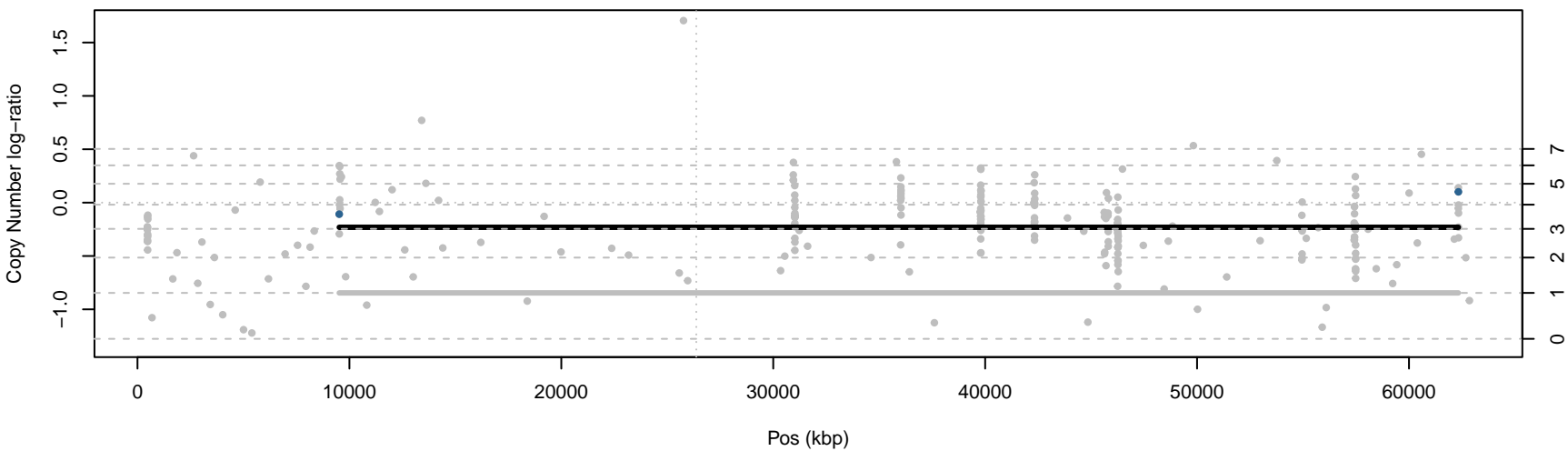
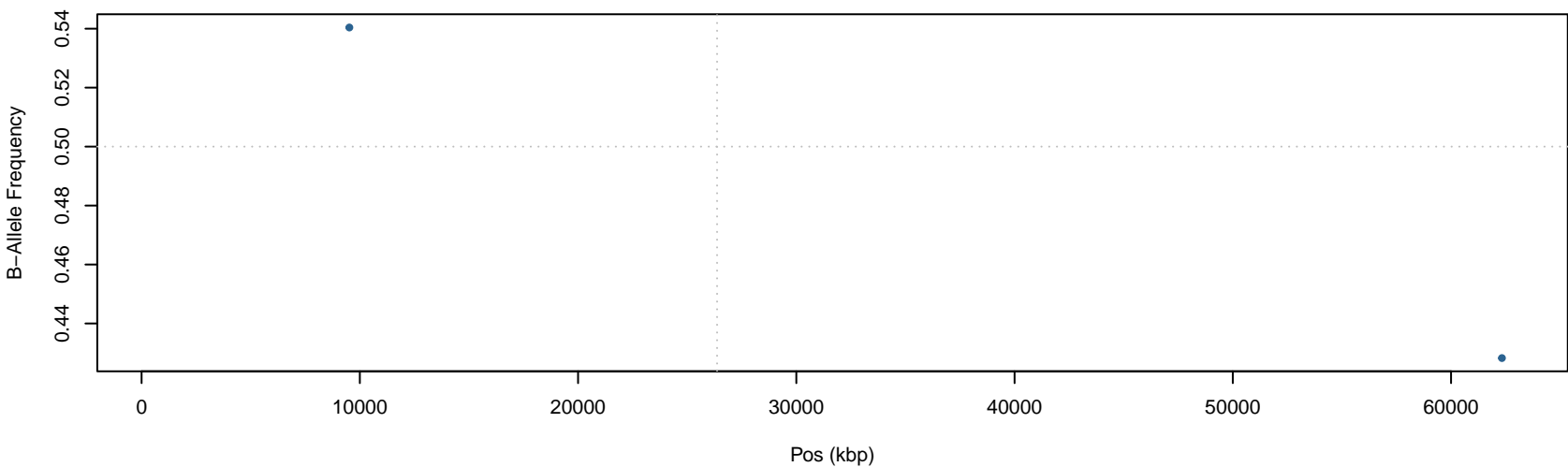
Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 16



Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 19

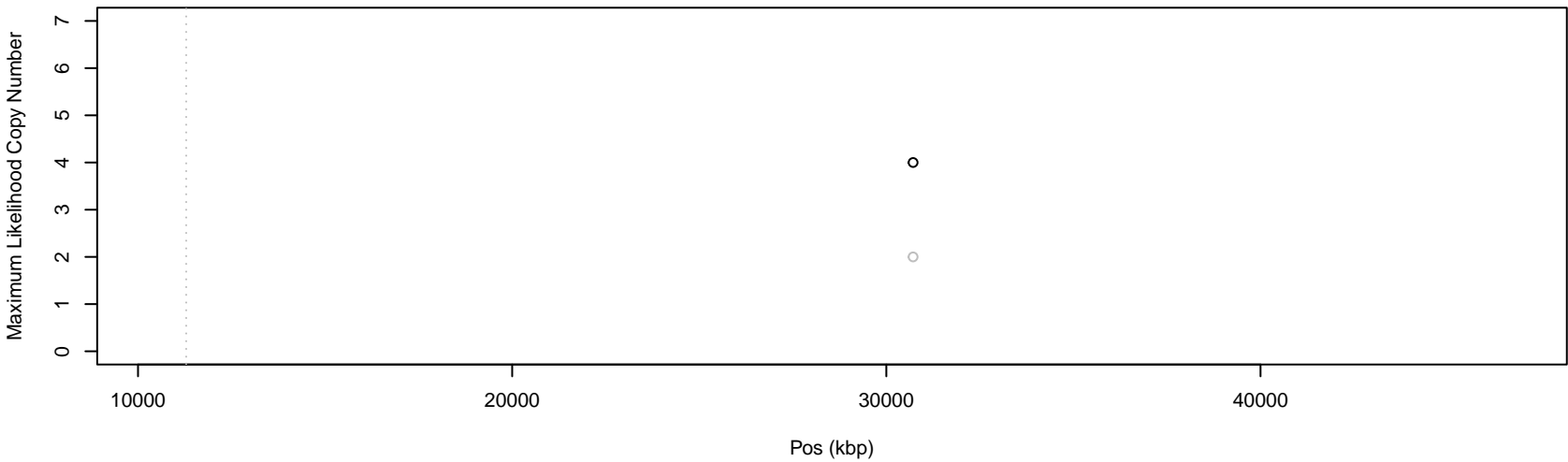
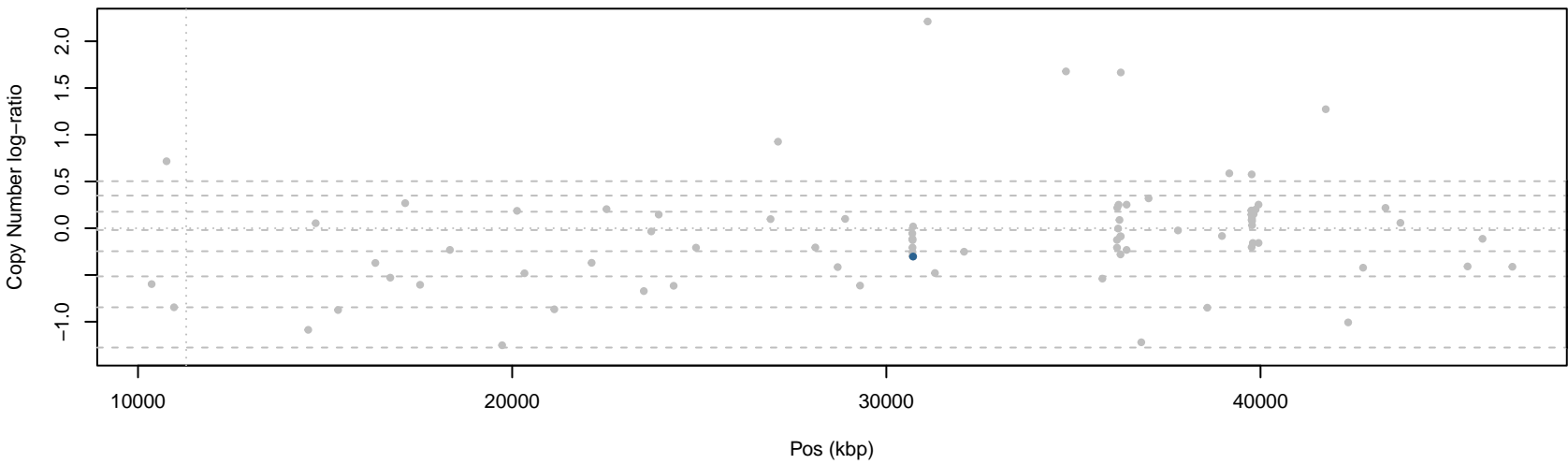
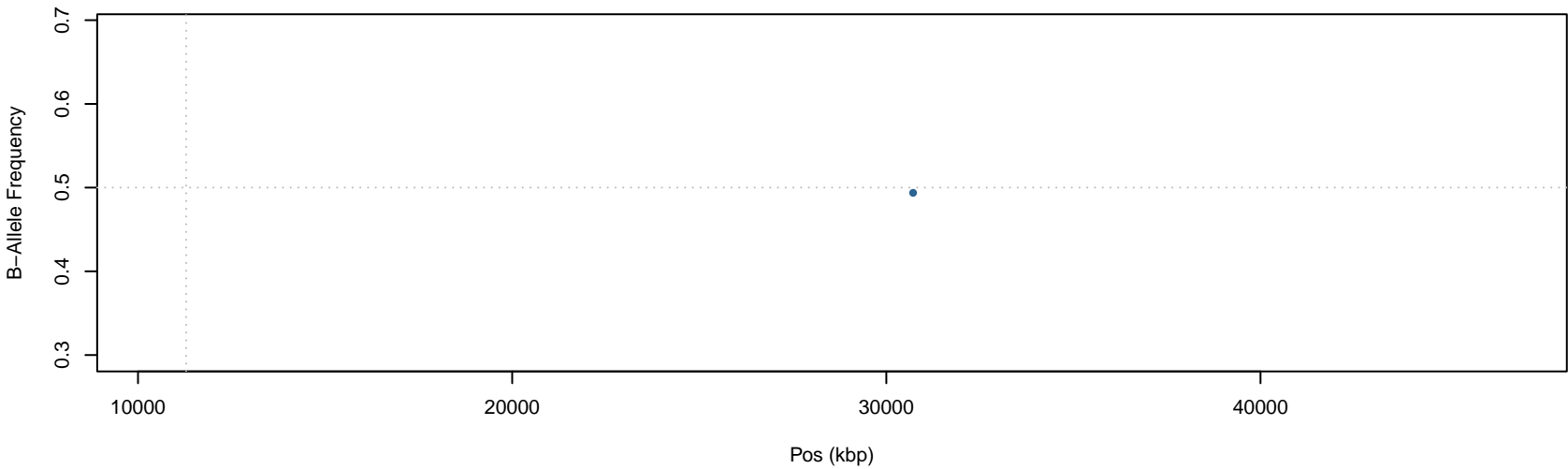


Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 20





Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 21



Purity: 0.41 Tumor ploidy: 4.092 SNV log-likelihood: -102.22 GoF: 86% Mean coverage: 710,524 Chromosome: 22

