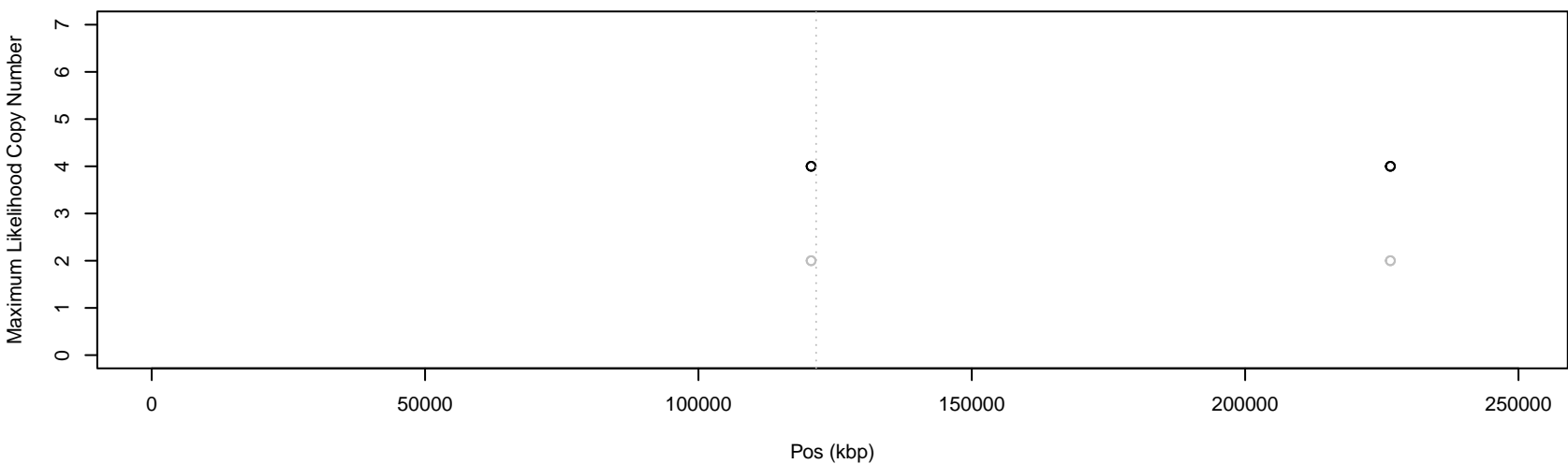
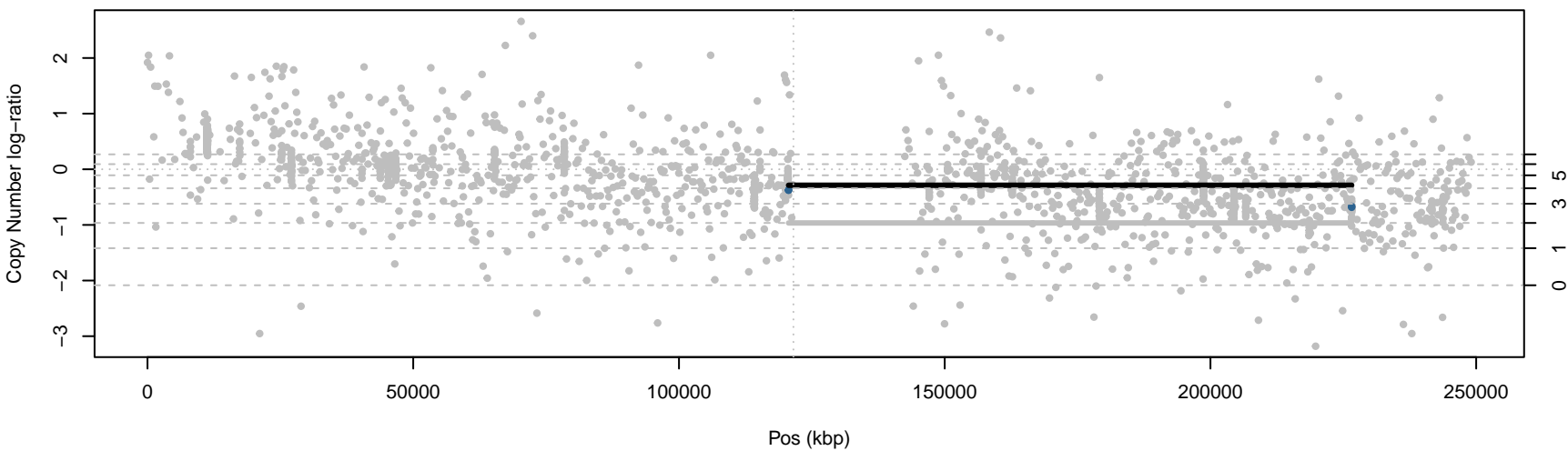
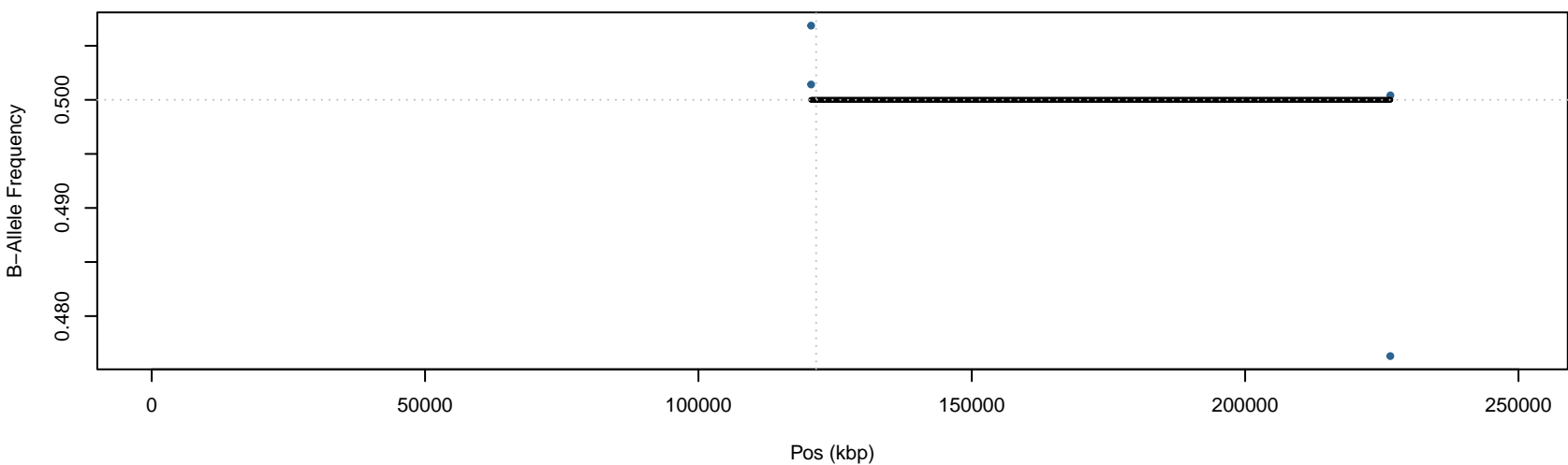
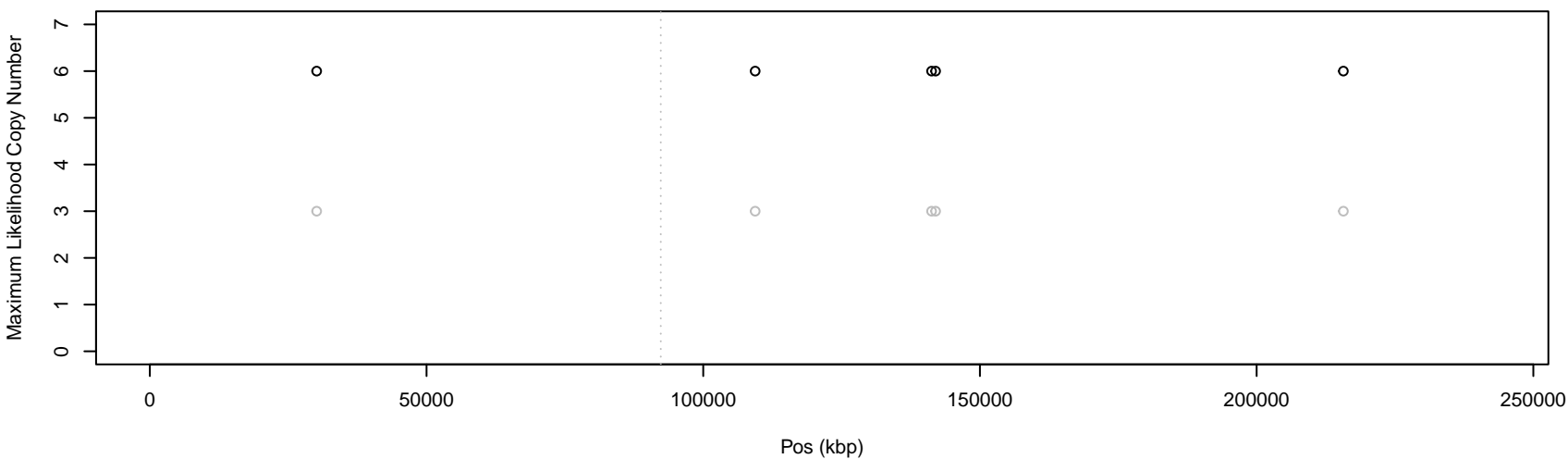
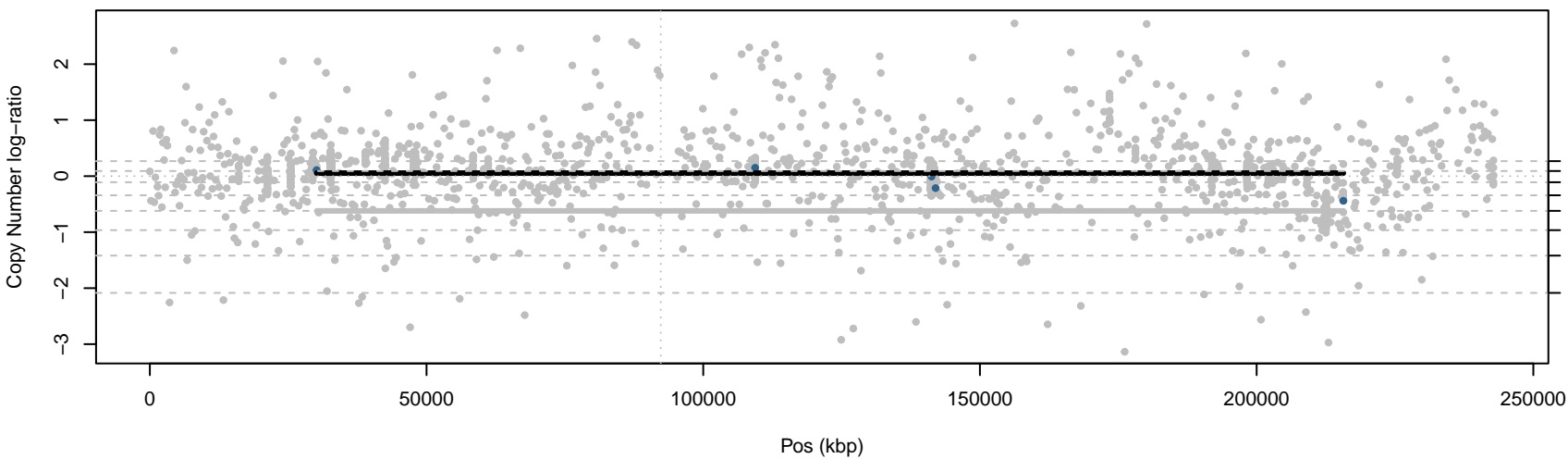
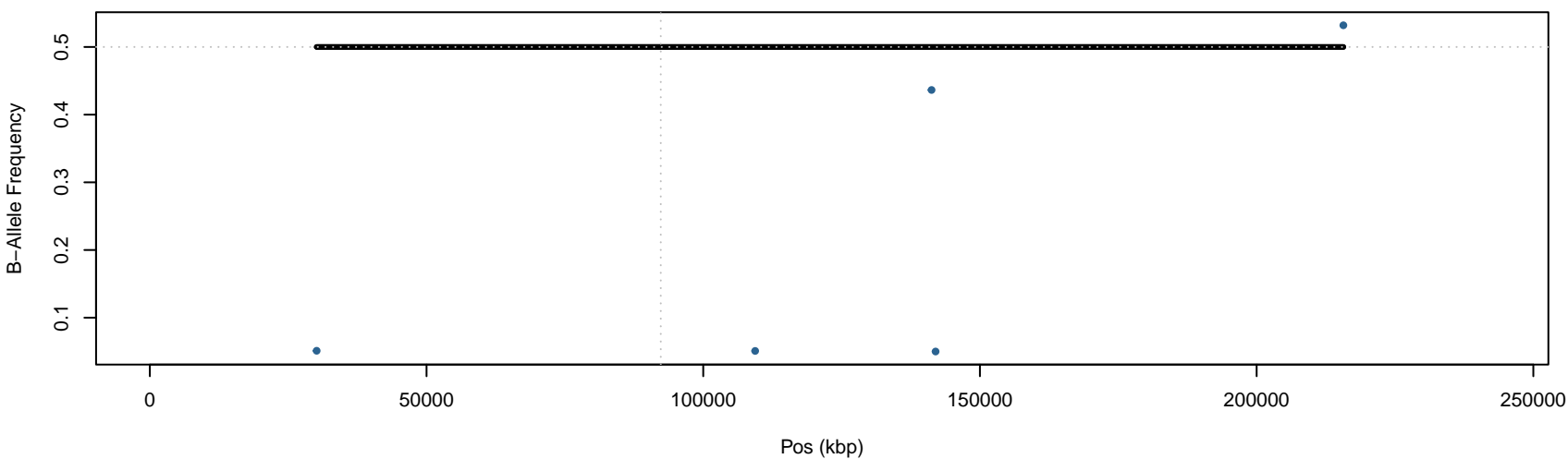


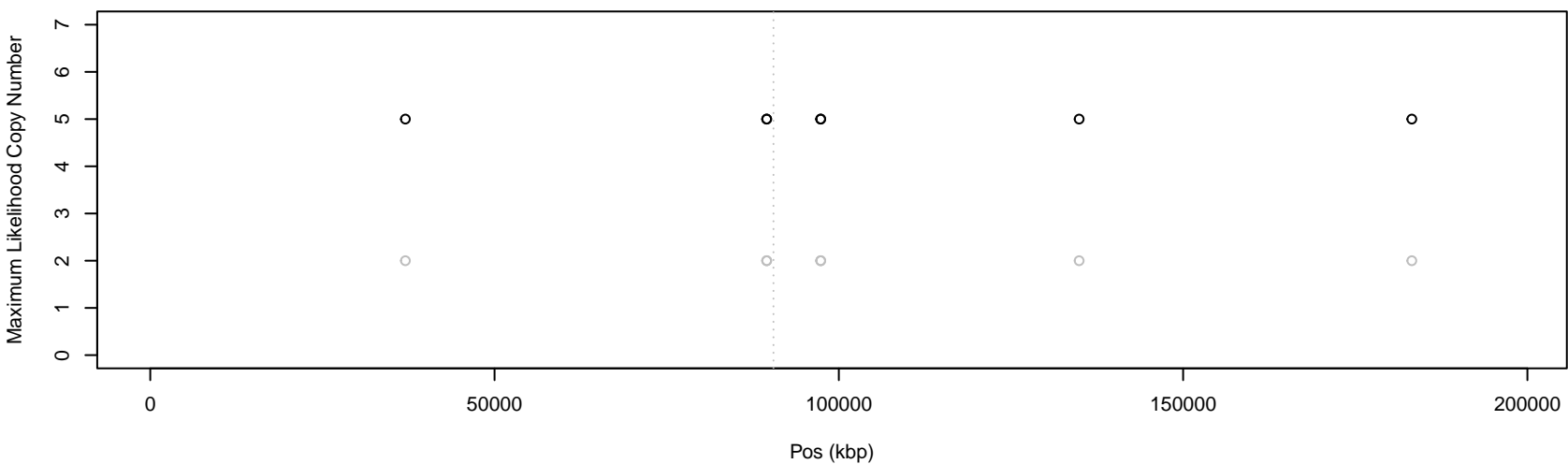
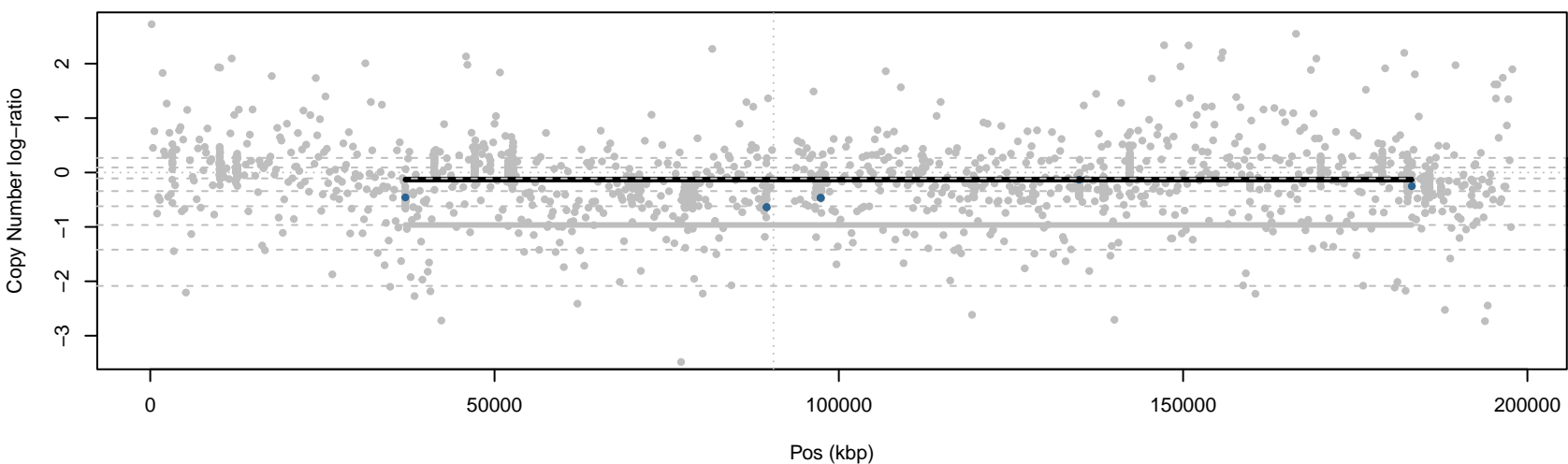
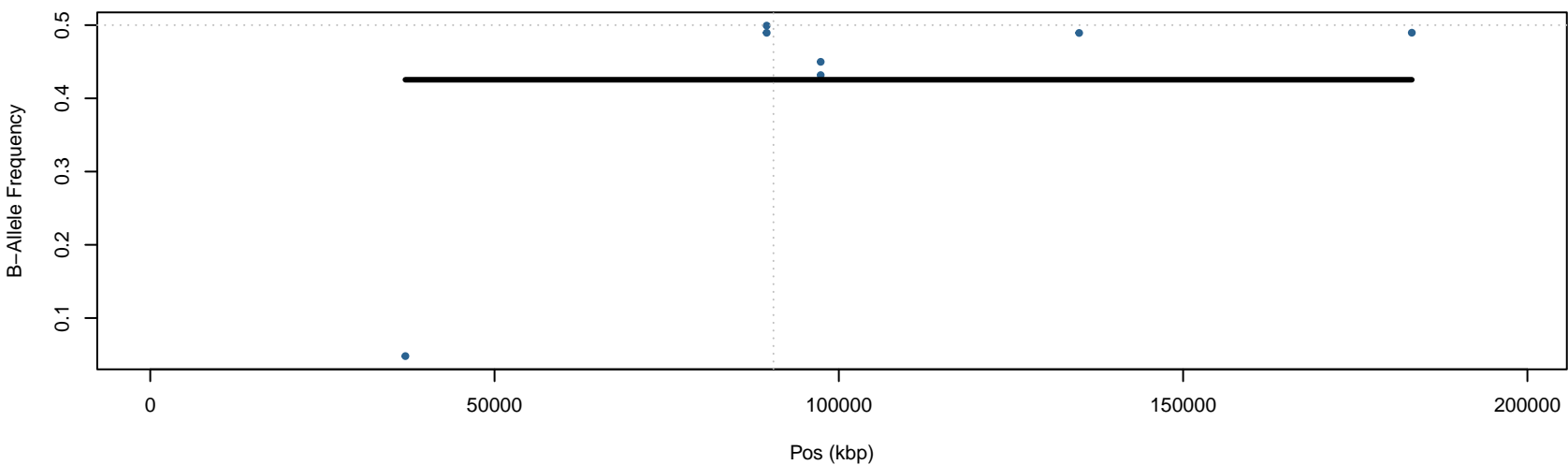
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 1



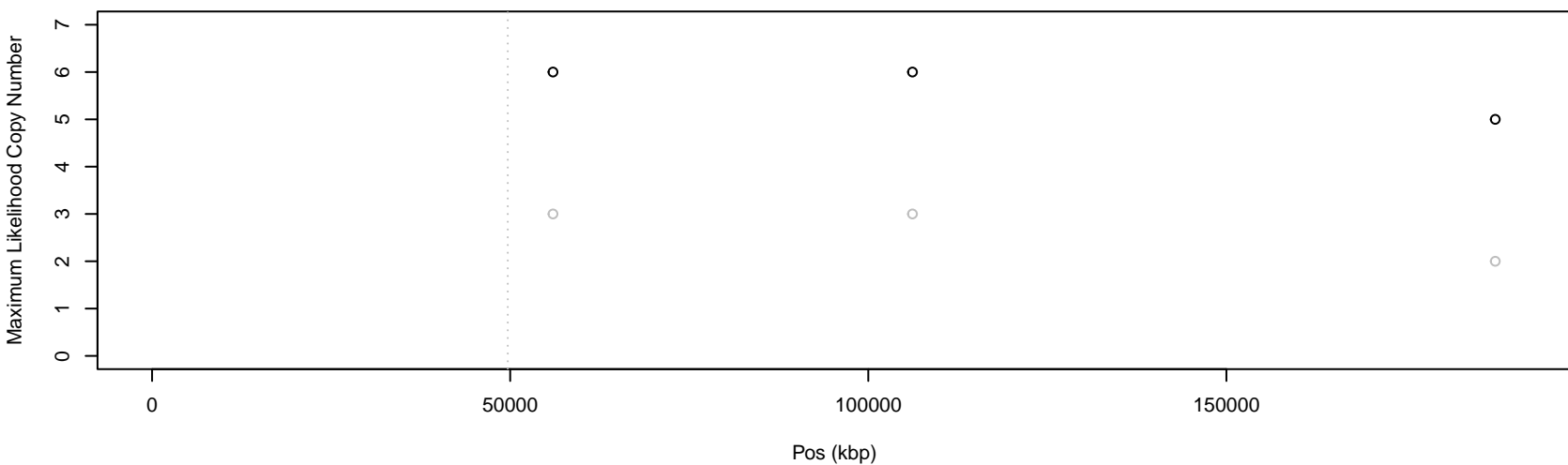
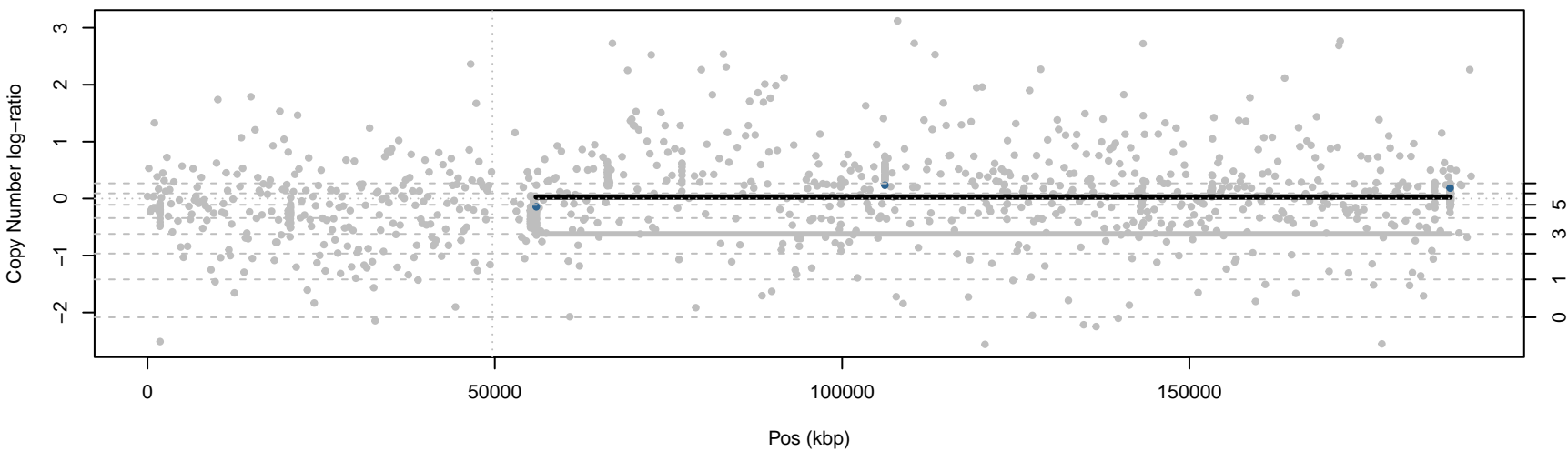
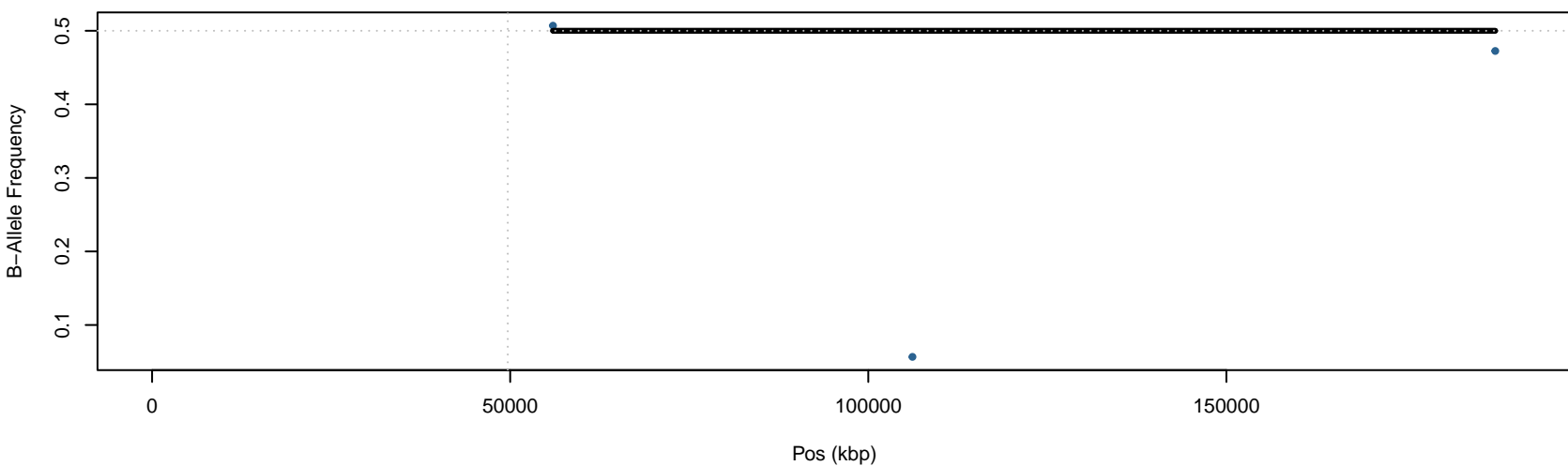
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 2



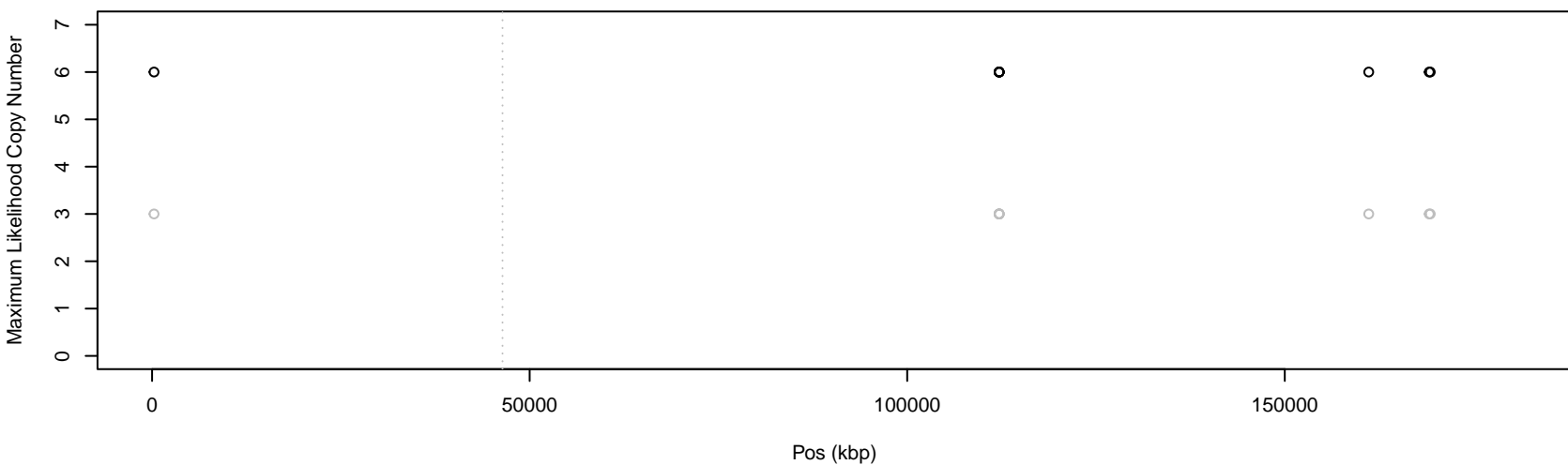
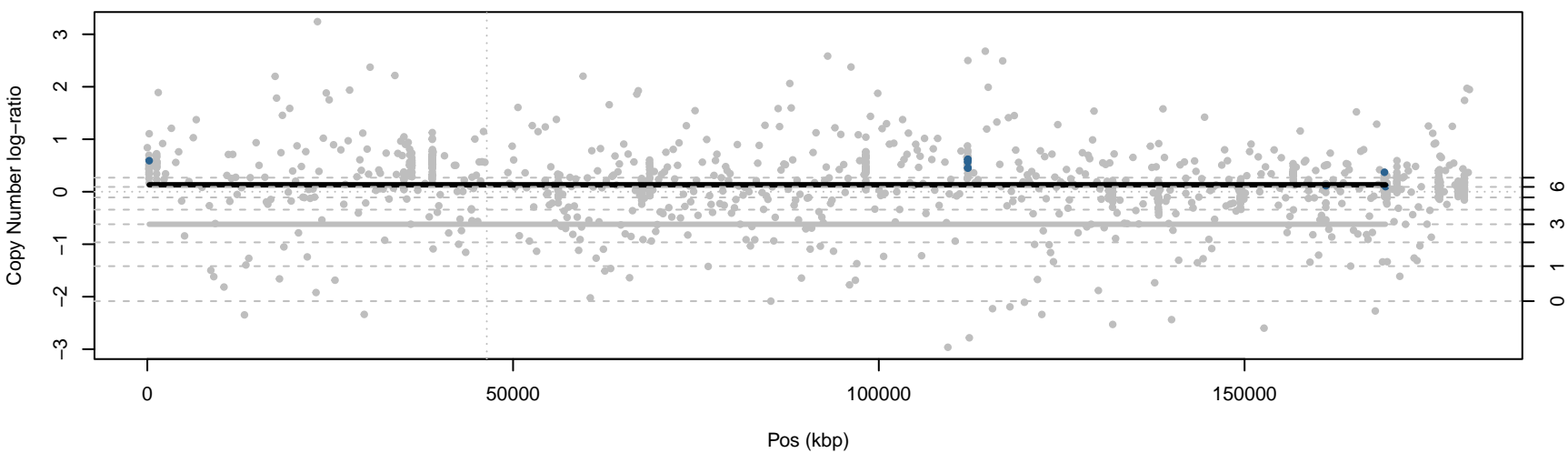
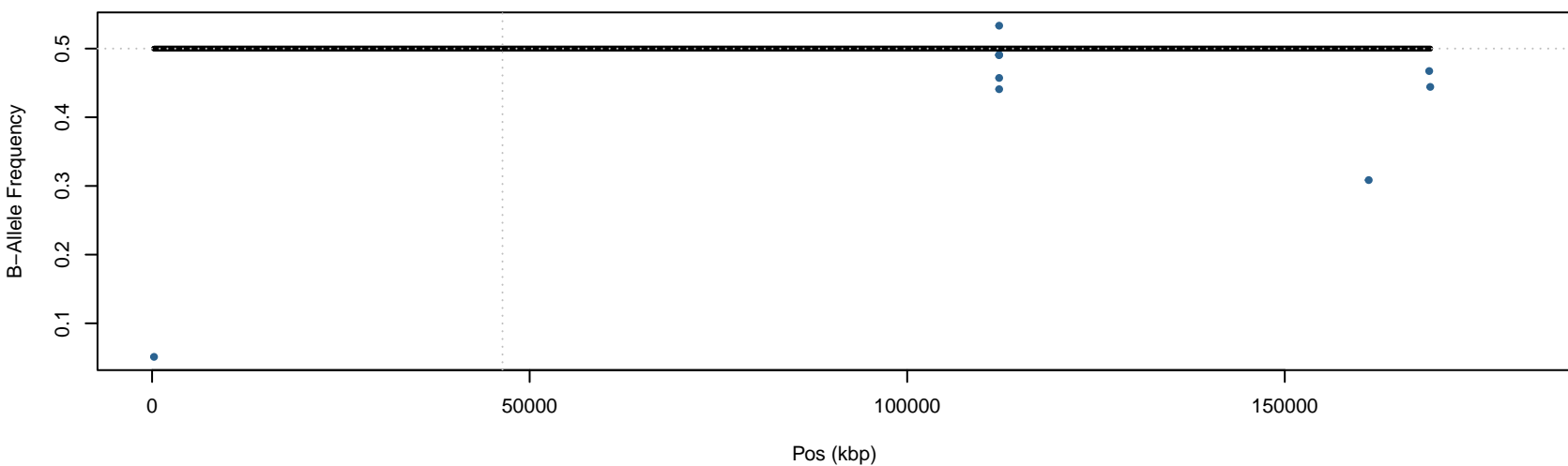
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 3



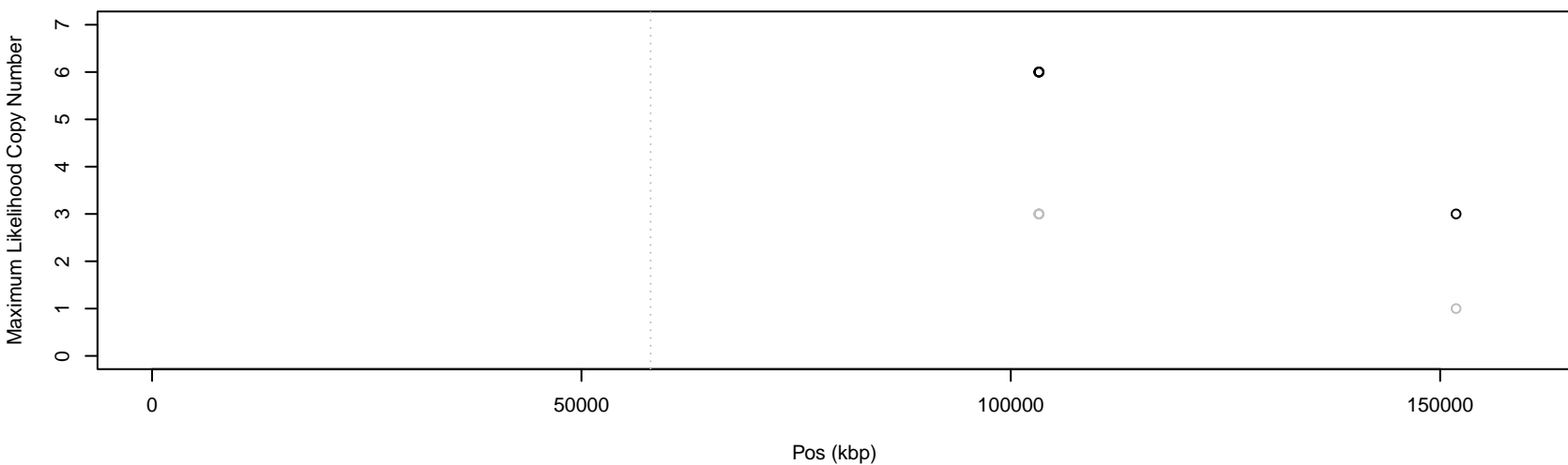
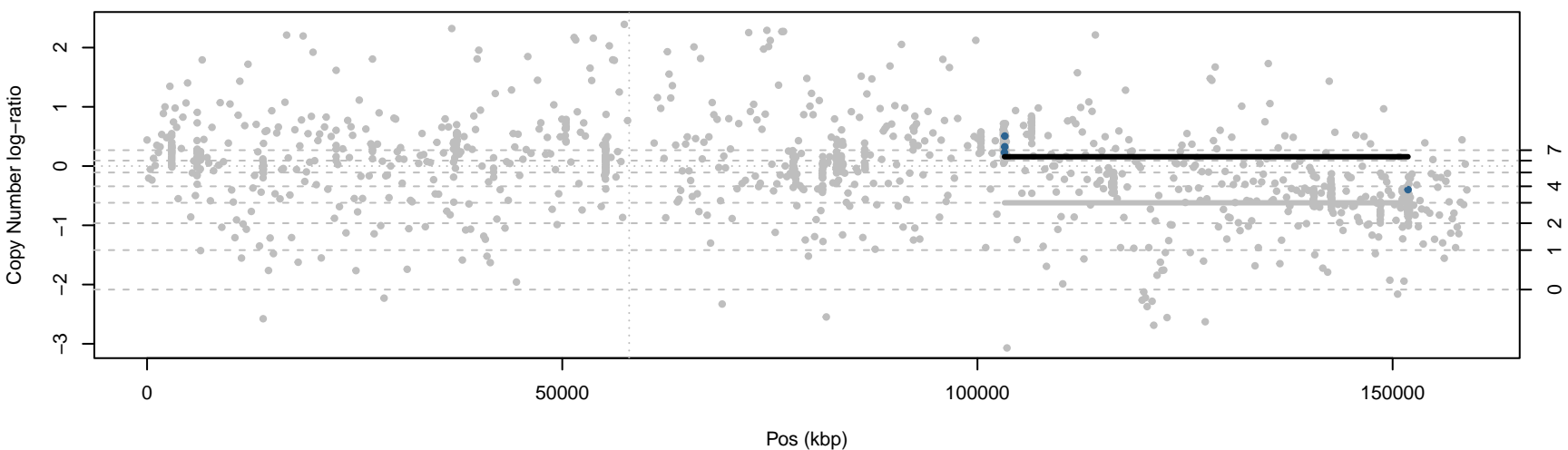
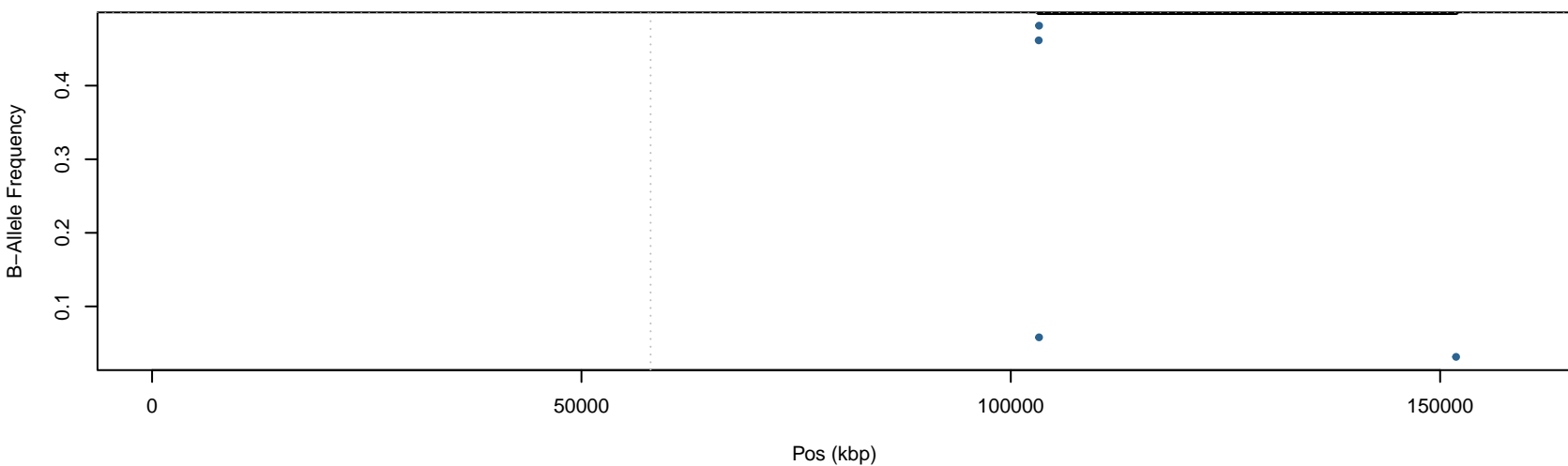
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 4



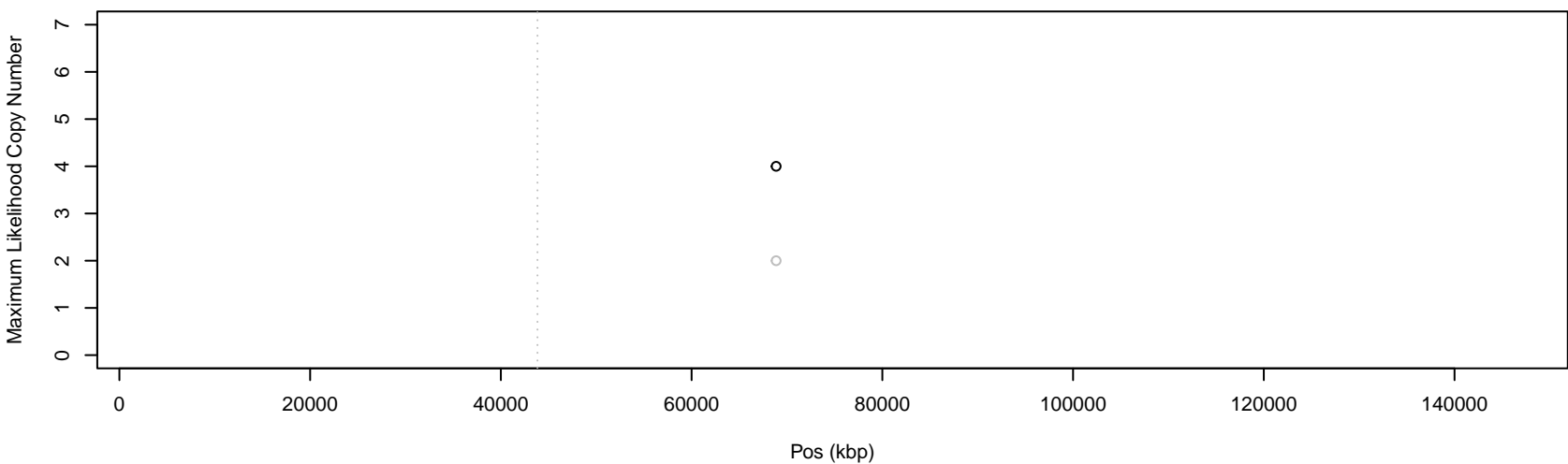
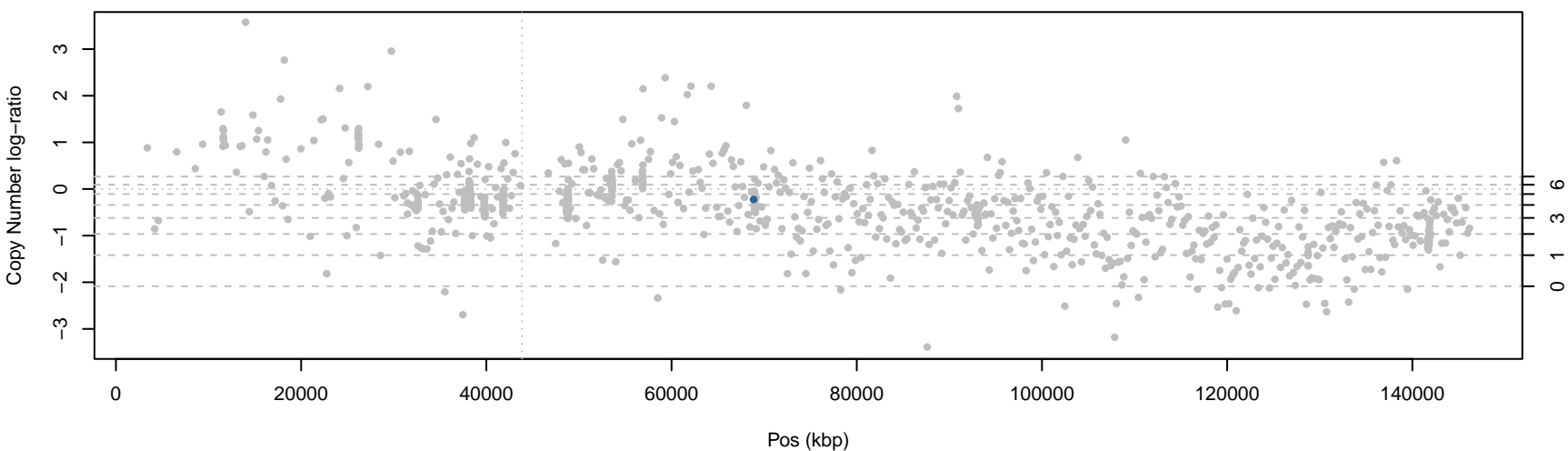
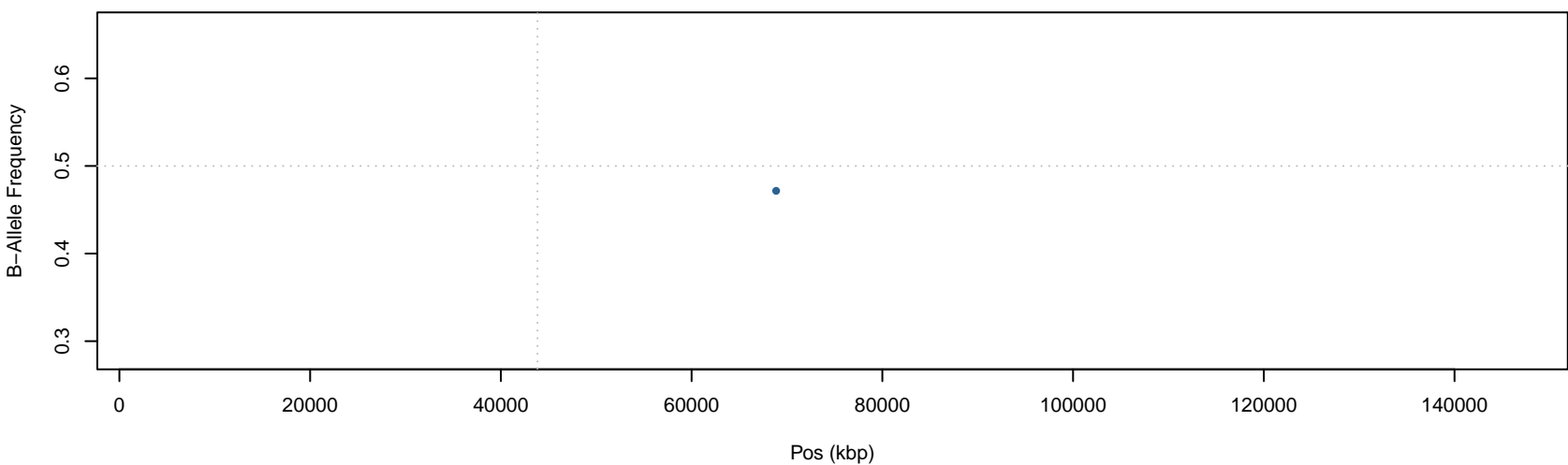
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 5



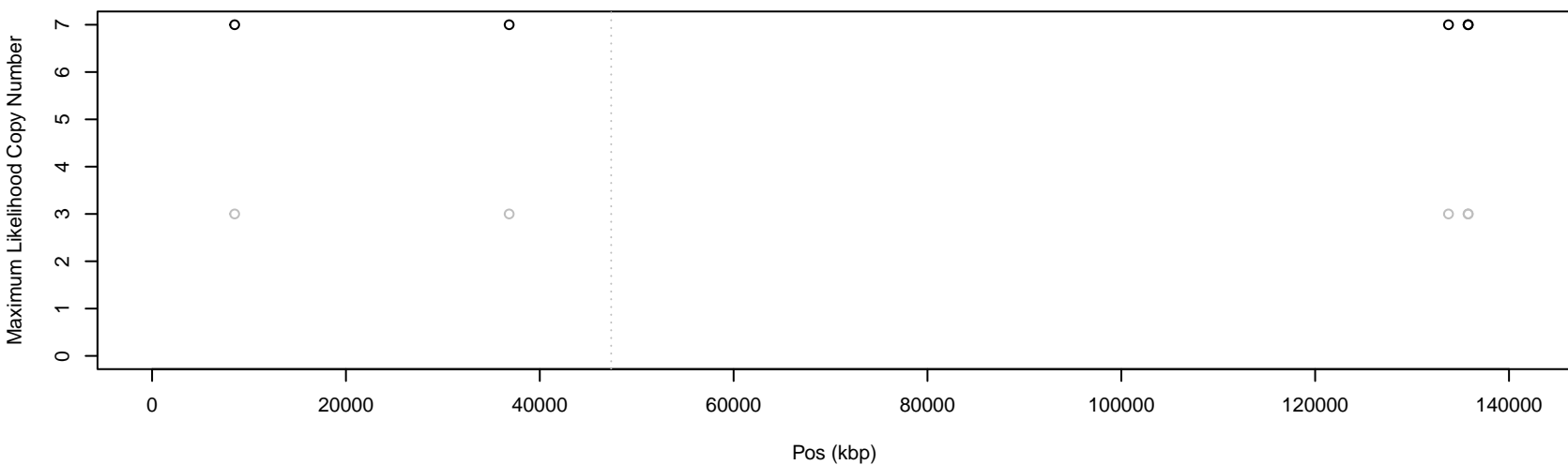
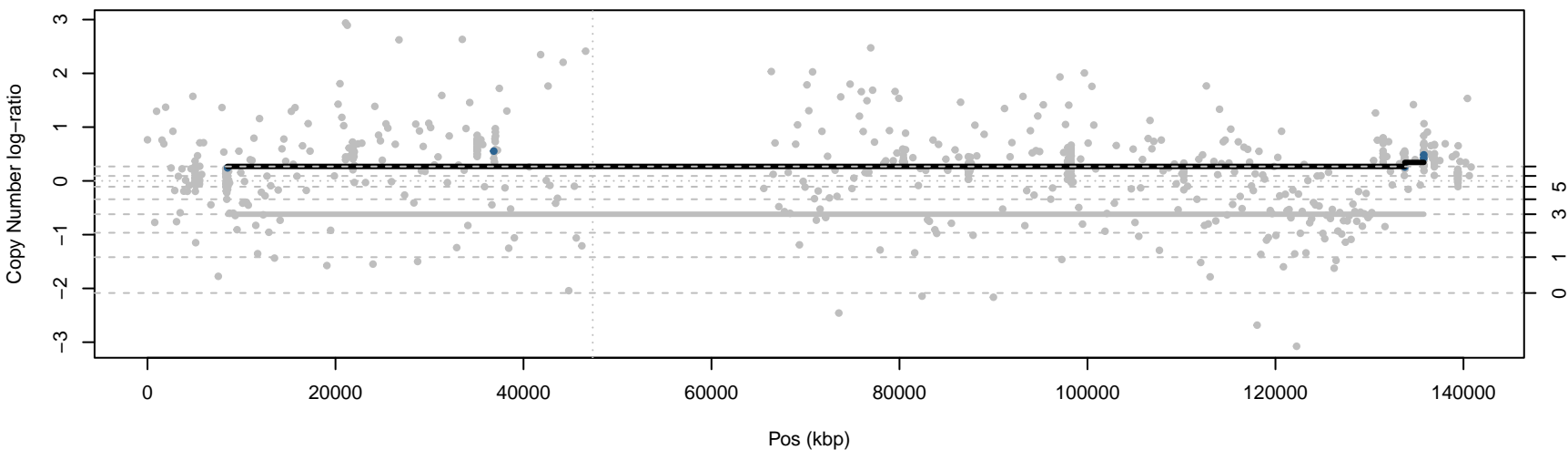
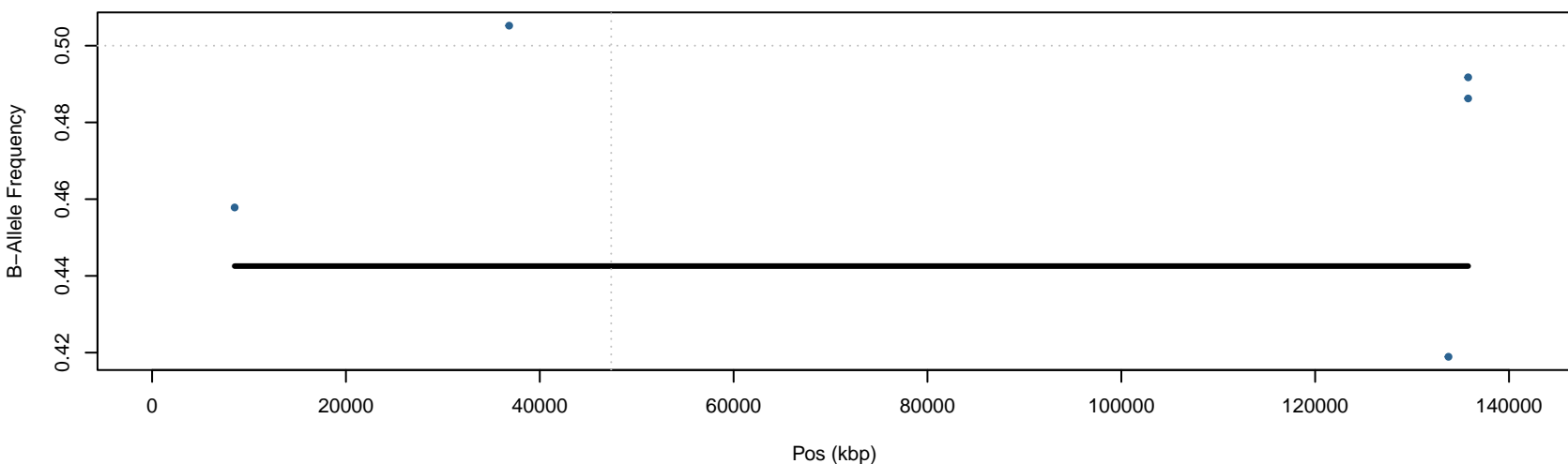
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 7



Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 8

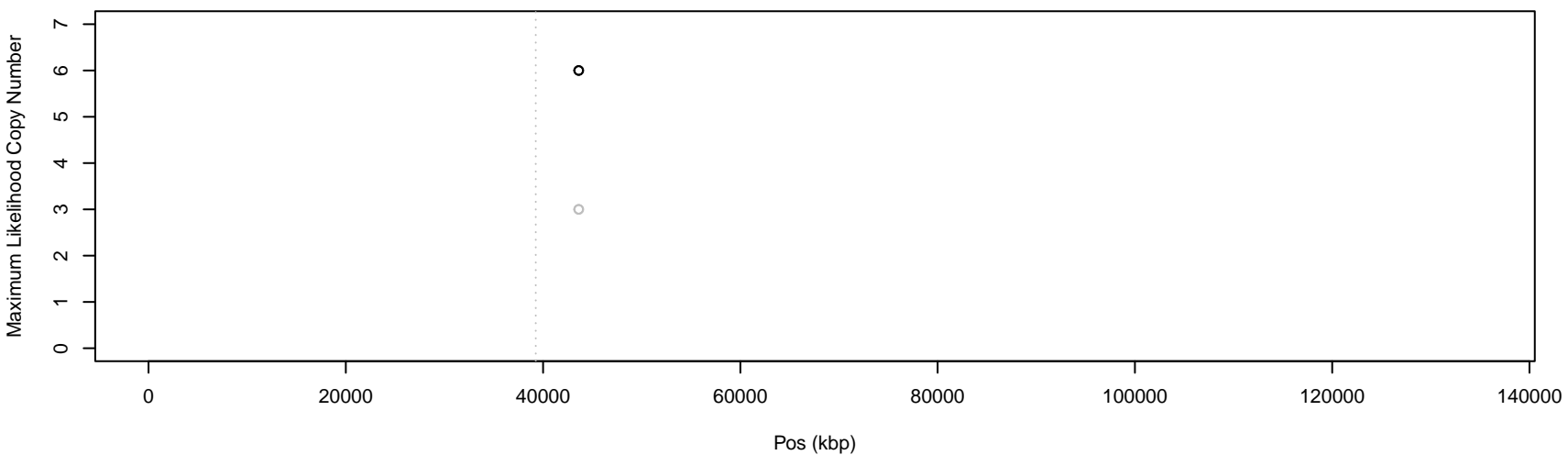
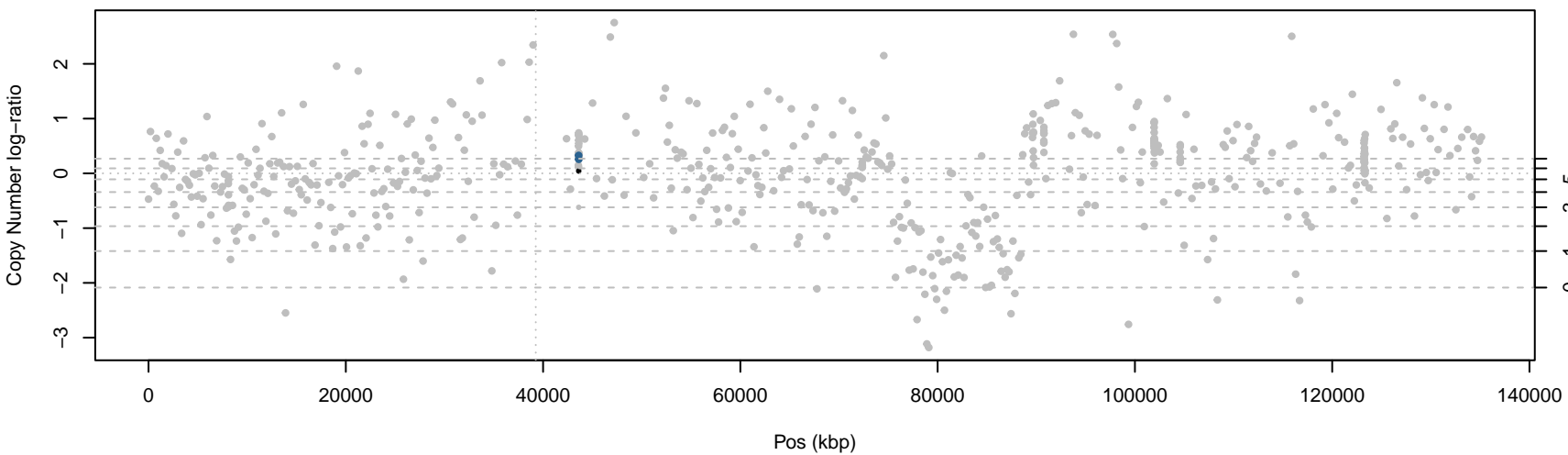
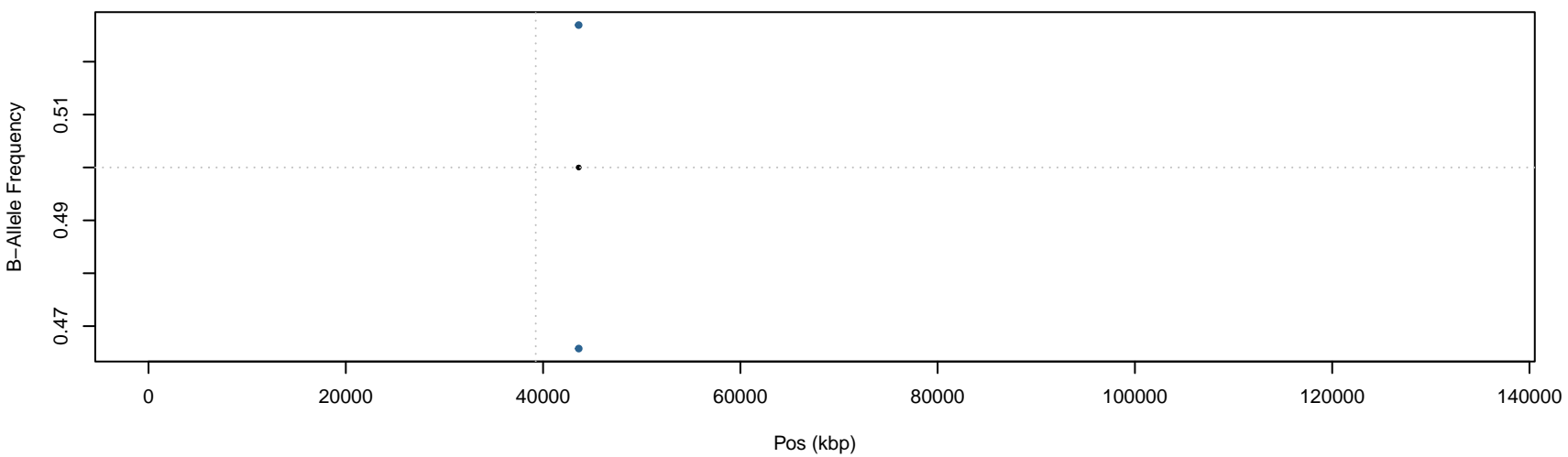


Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 9

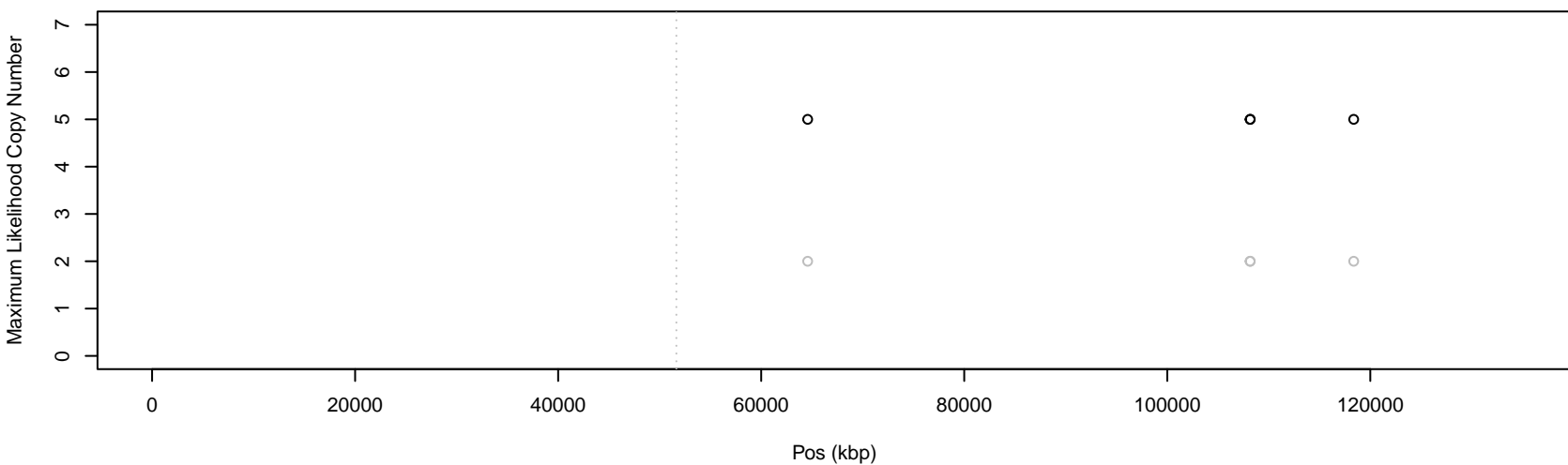
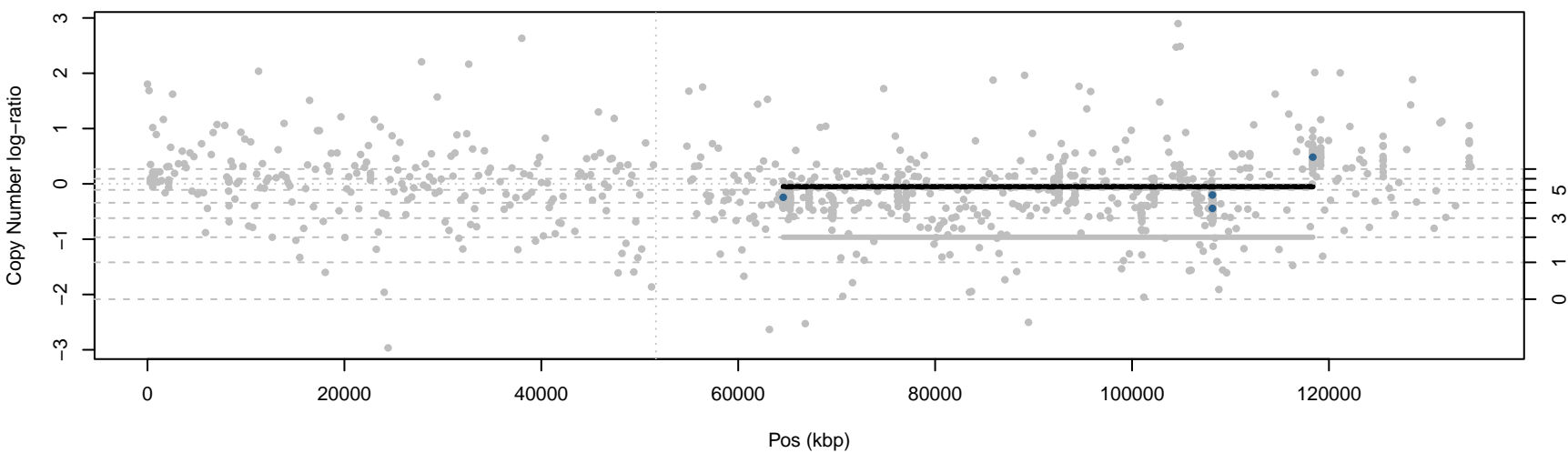
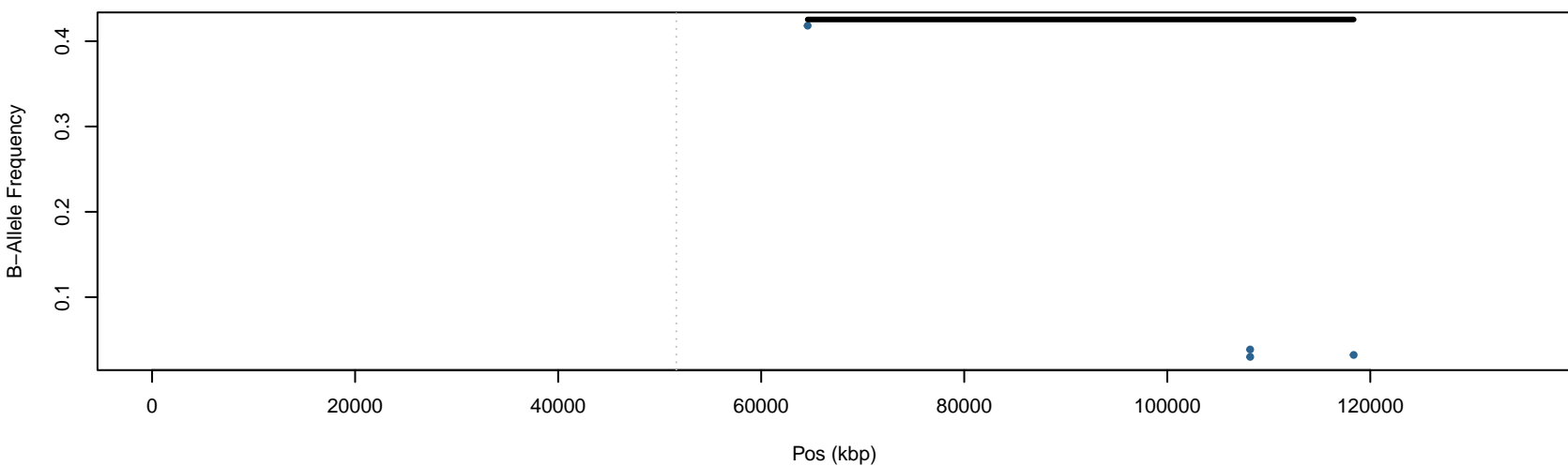




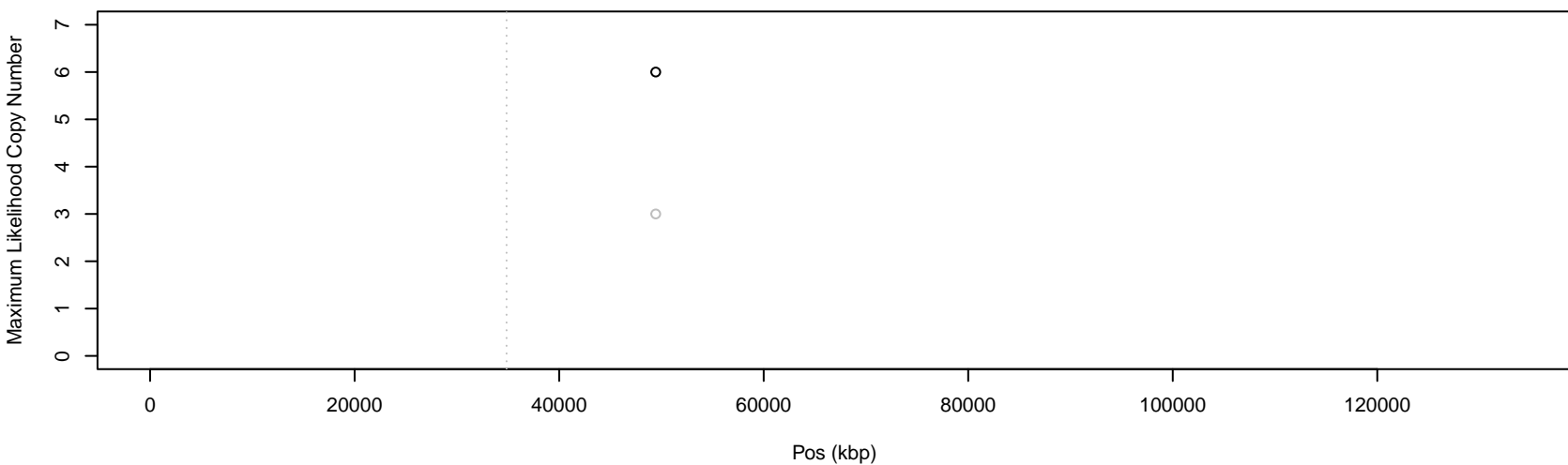
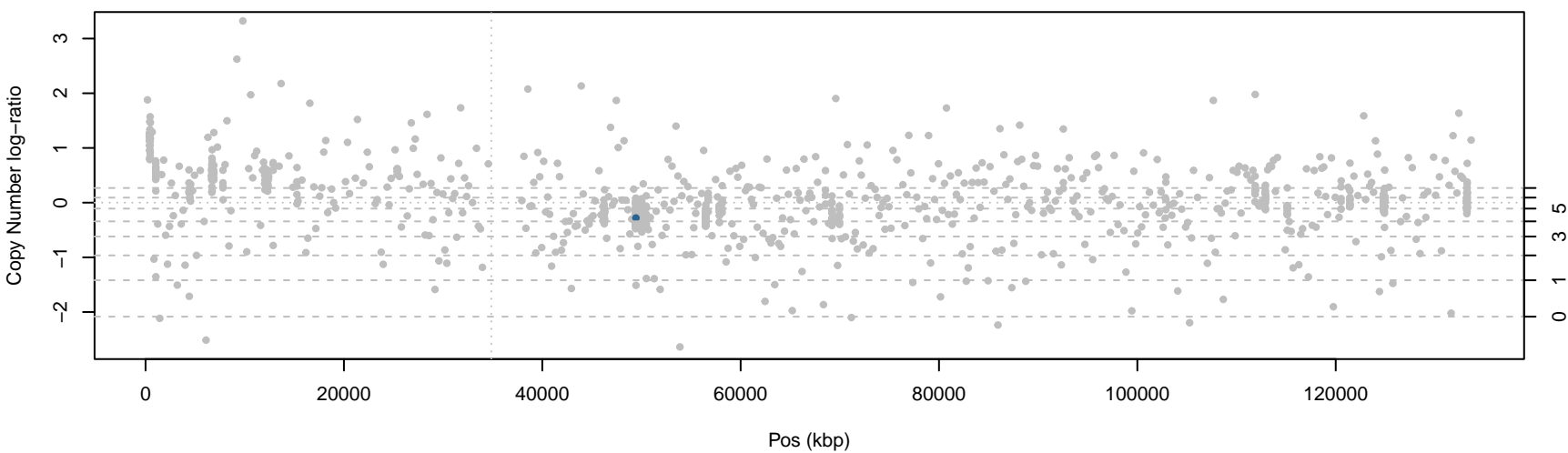
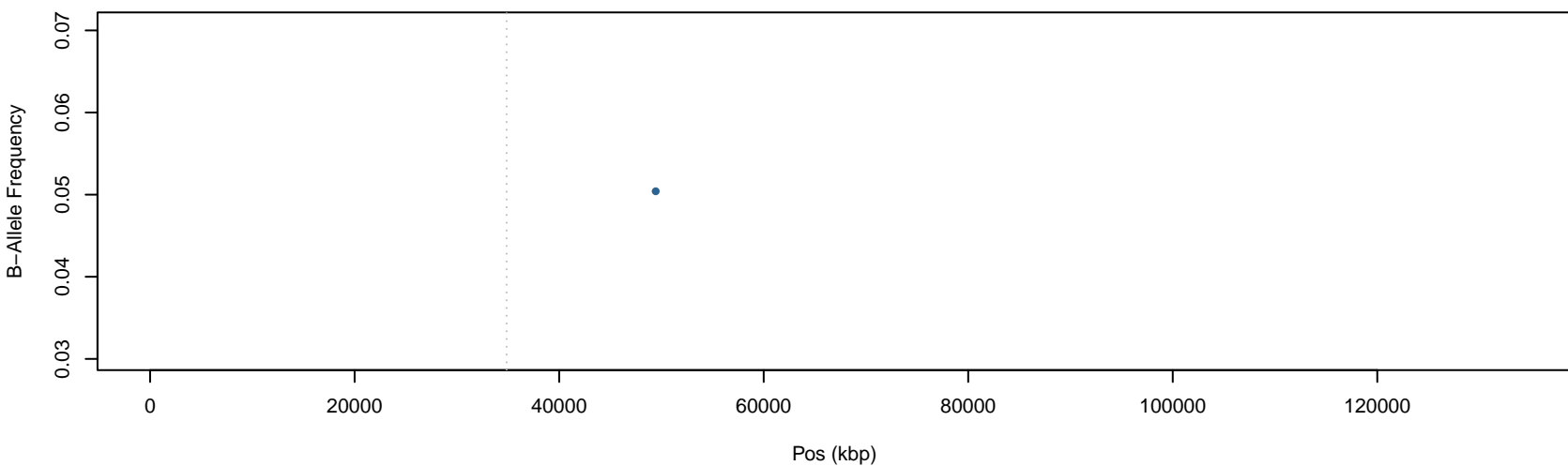
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 10



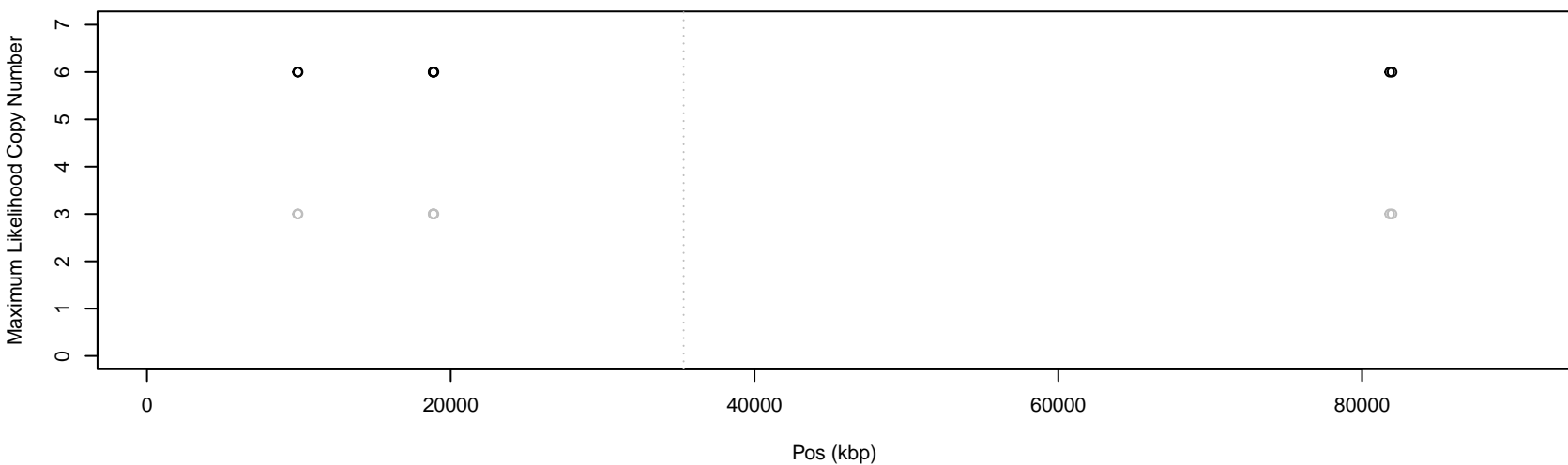
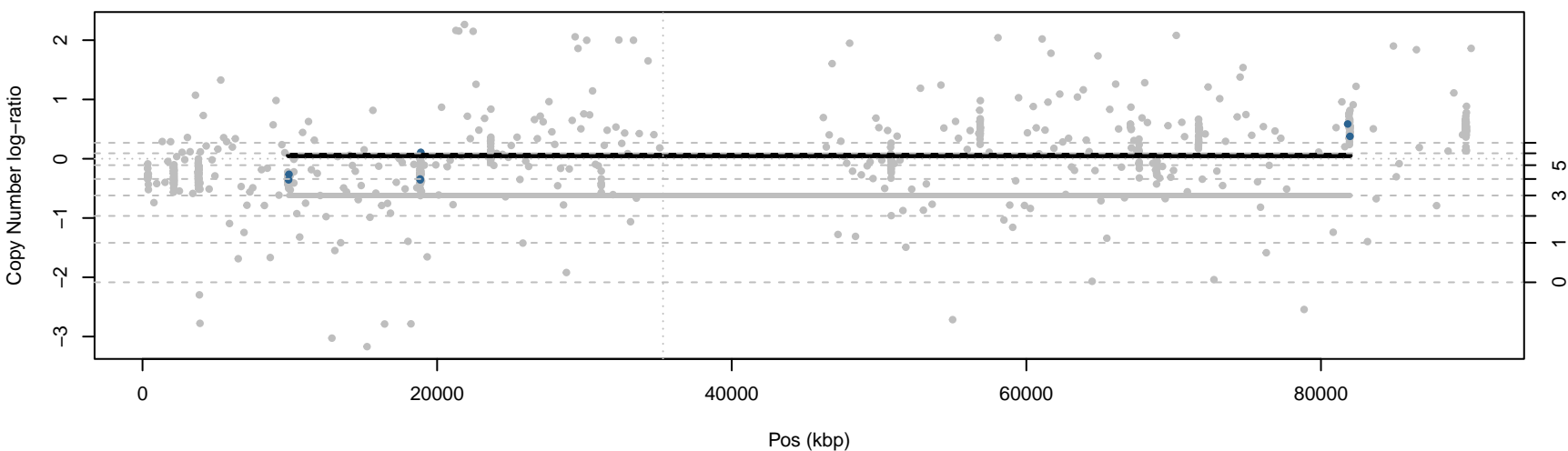
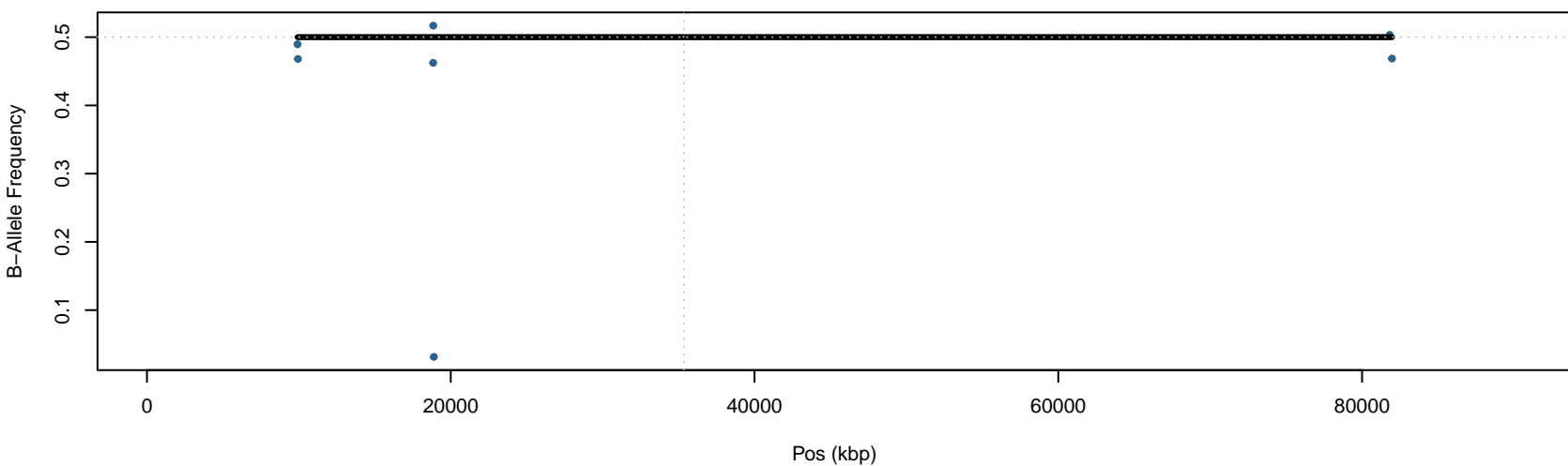
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 11



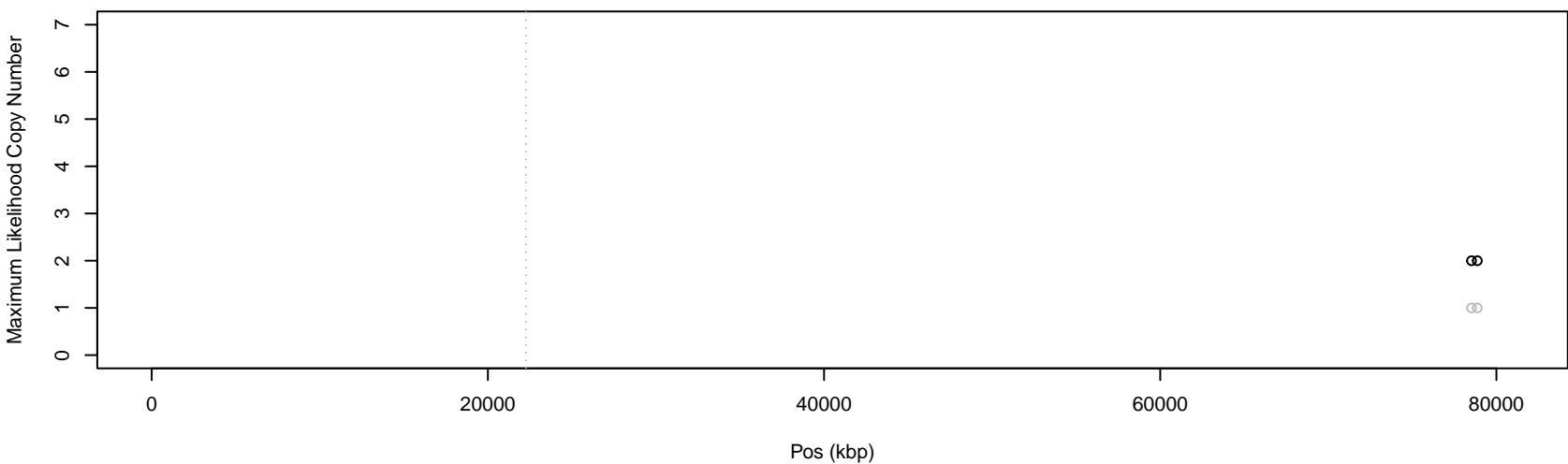
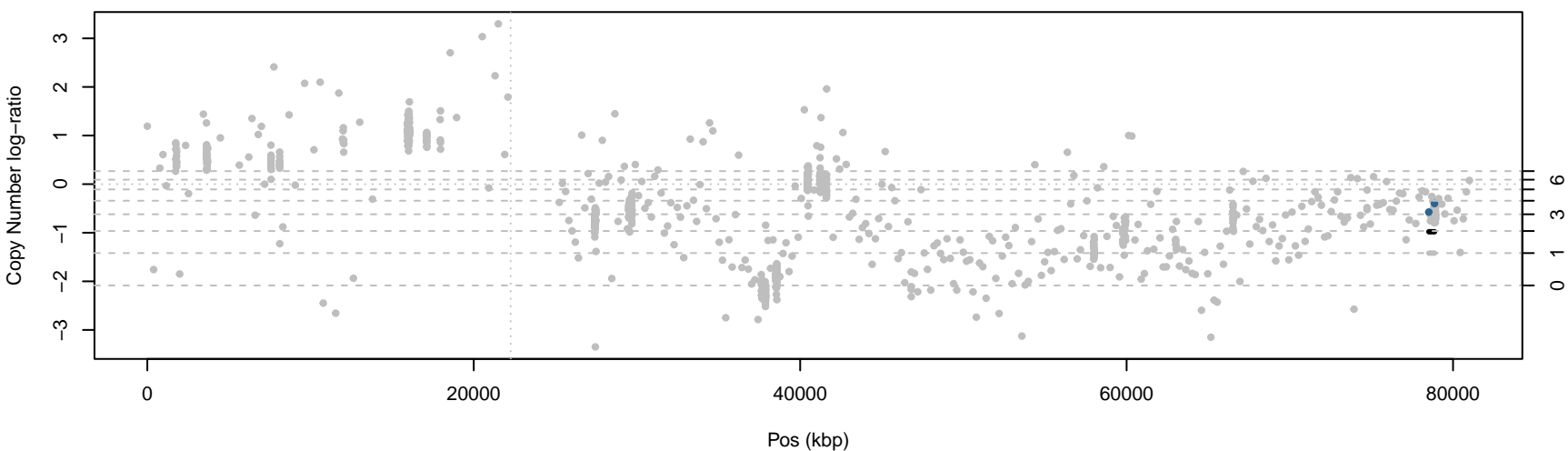
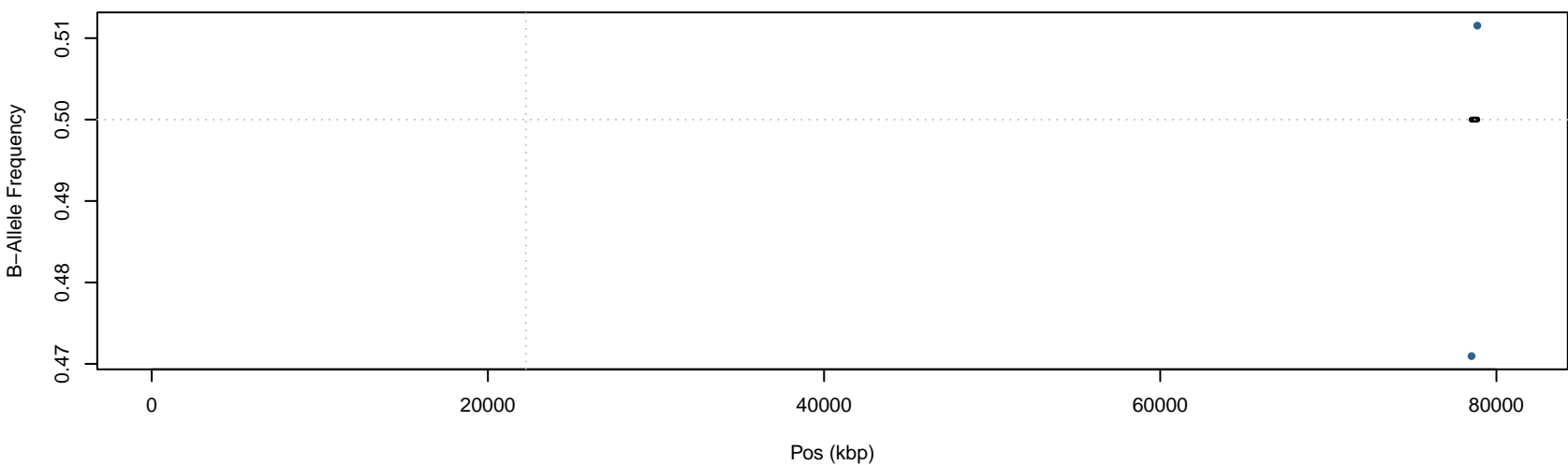
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 12



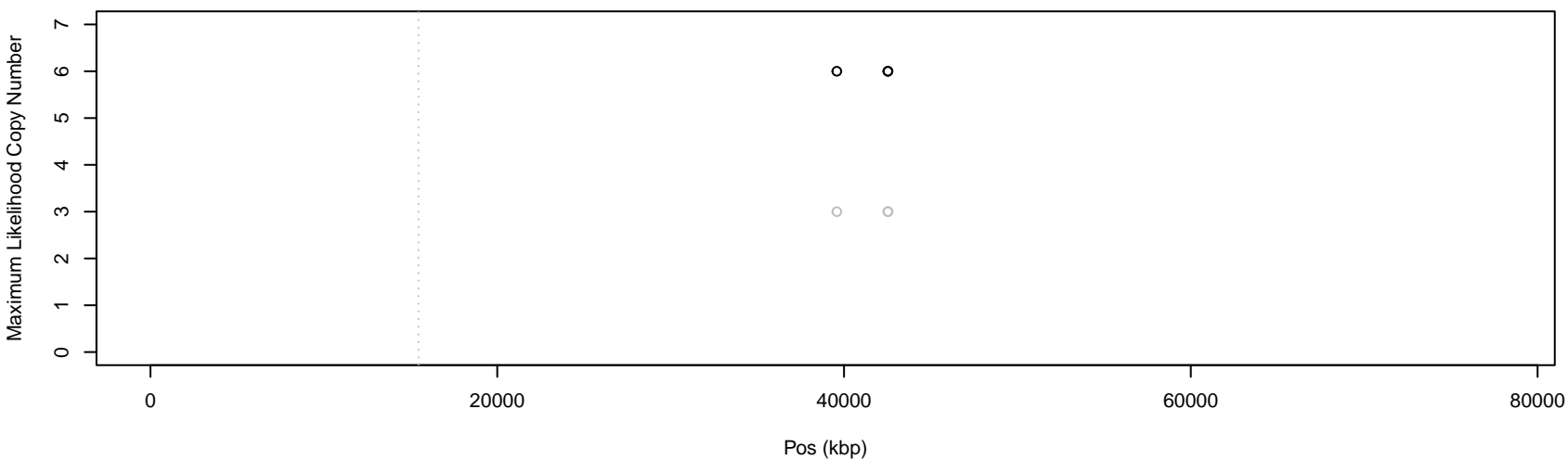
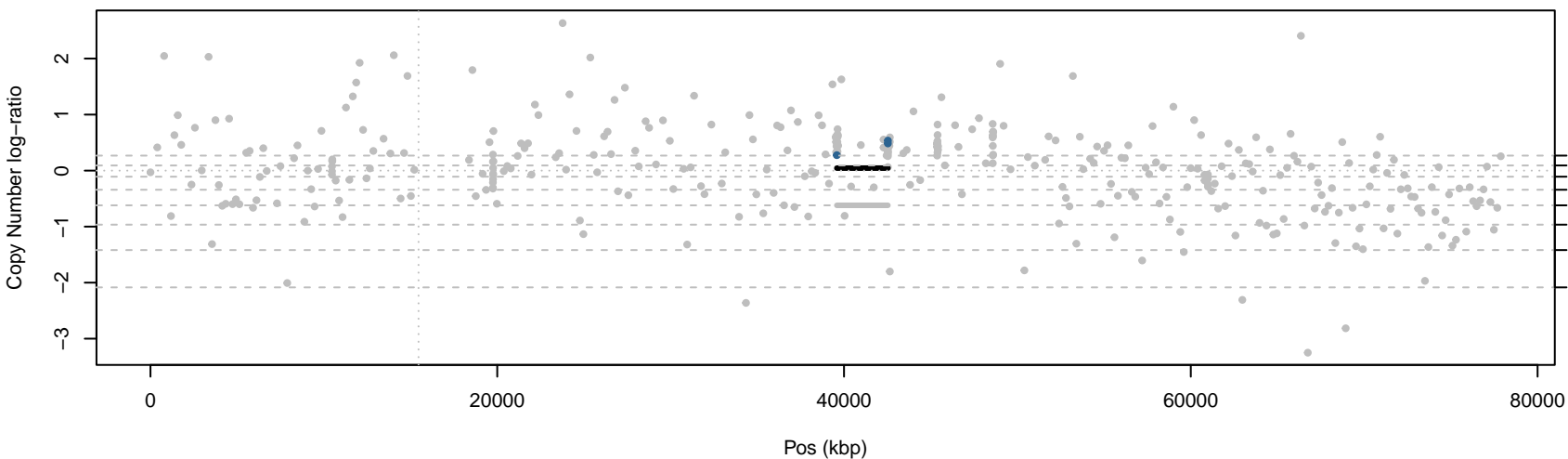
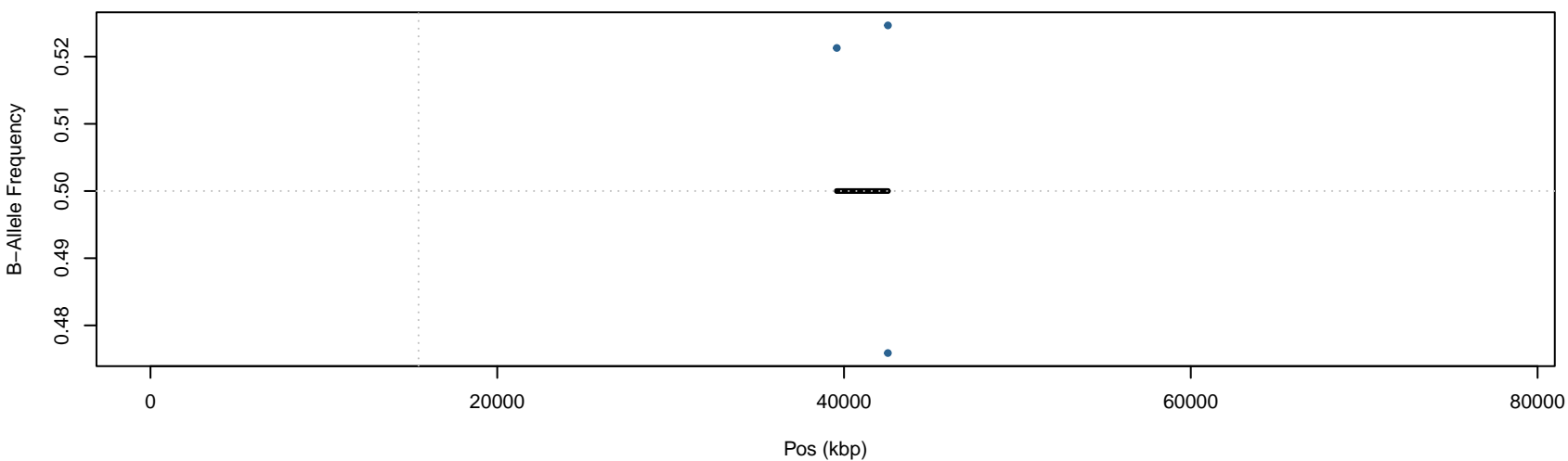
Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 16



Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 17



Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 18



Purity: 0.54 Tumor ploidy: 5.525 SNV log-likelihood: -36.34 GoF: 89.1% Mean coverage: 505;329 Chromosome: 20

