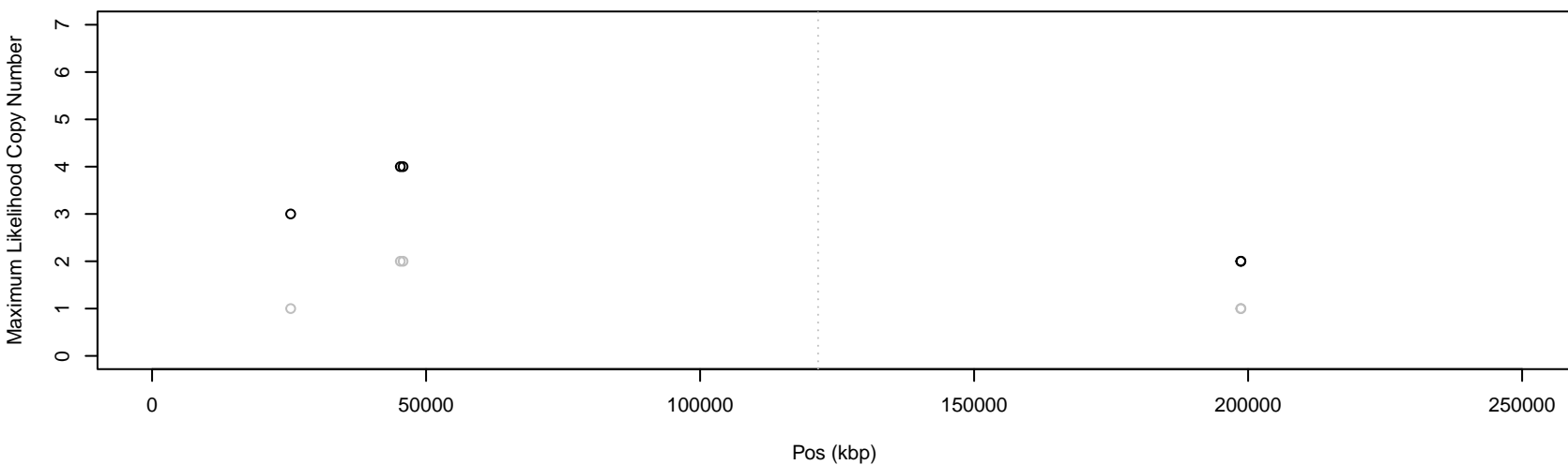
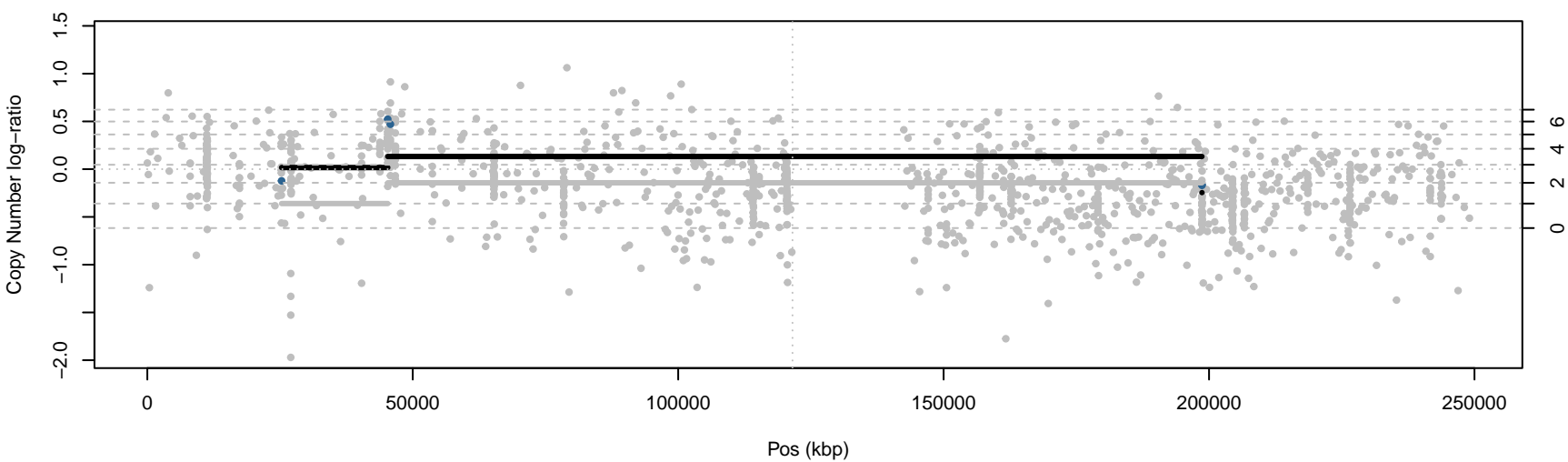
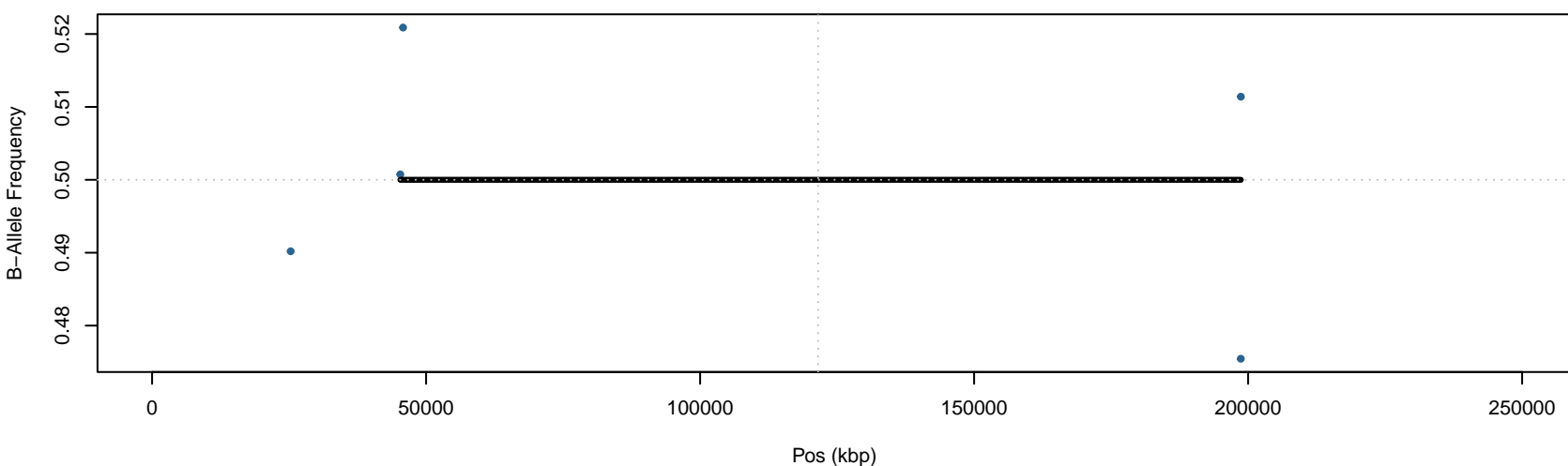
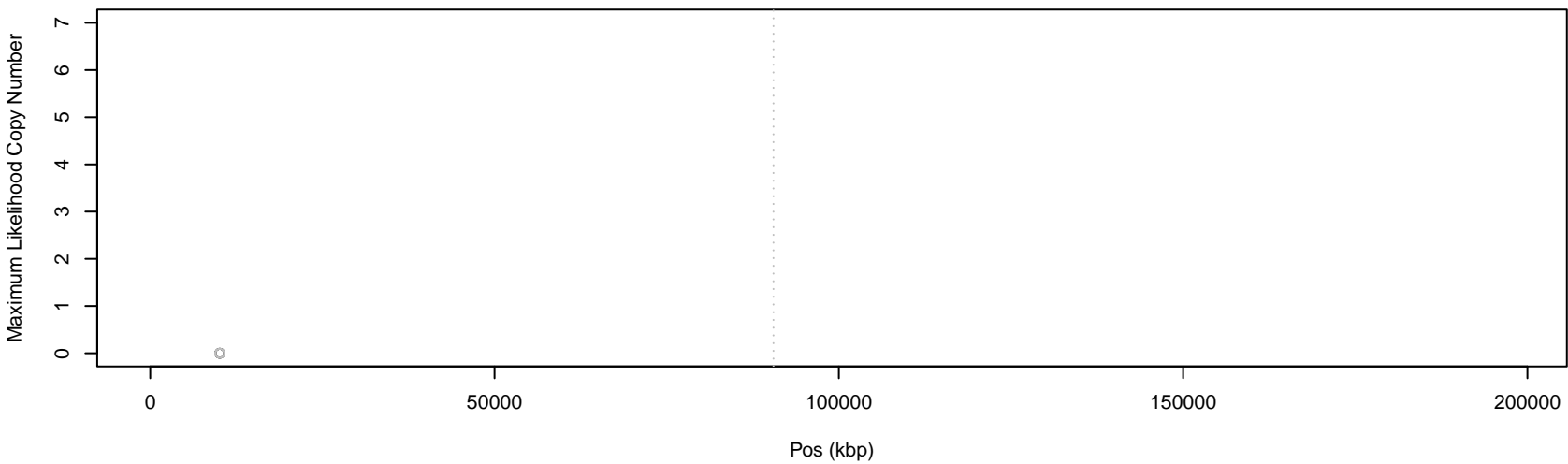
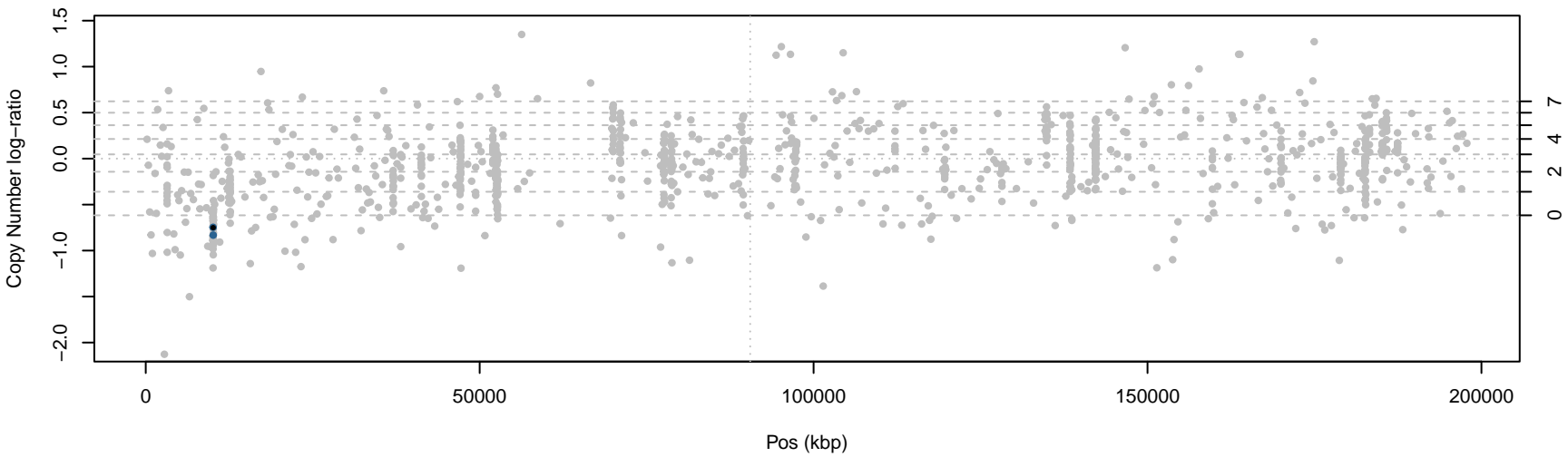
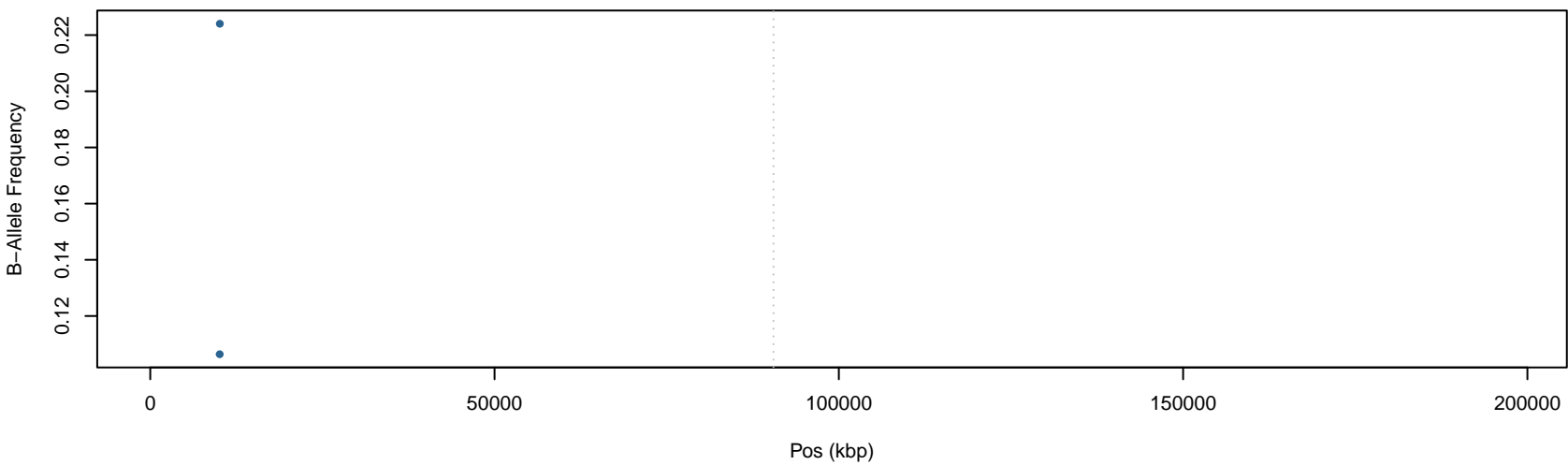


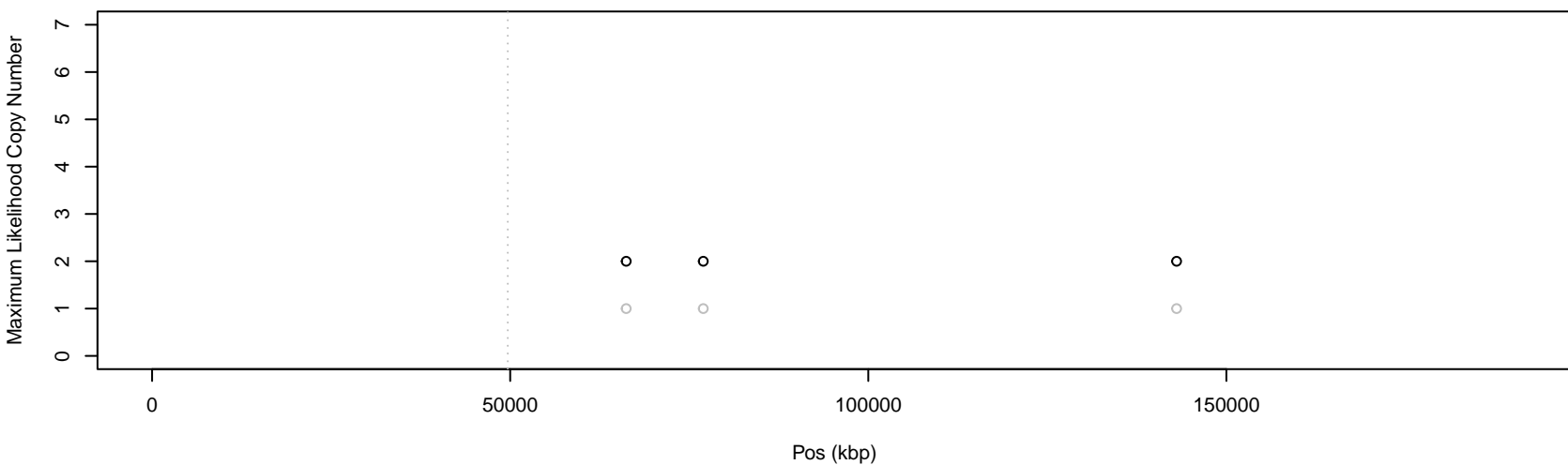
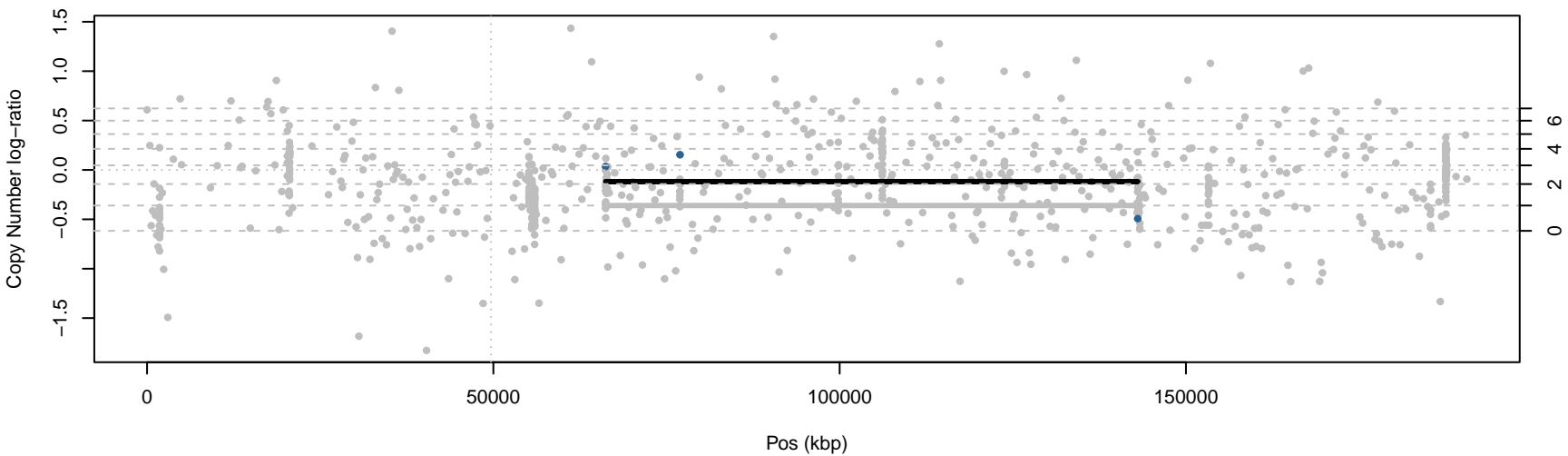
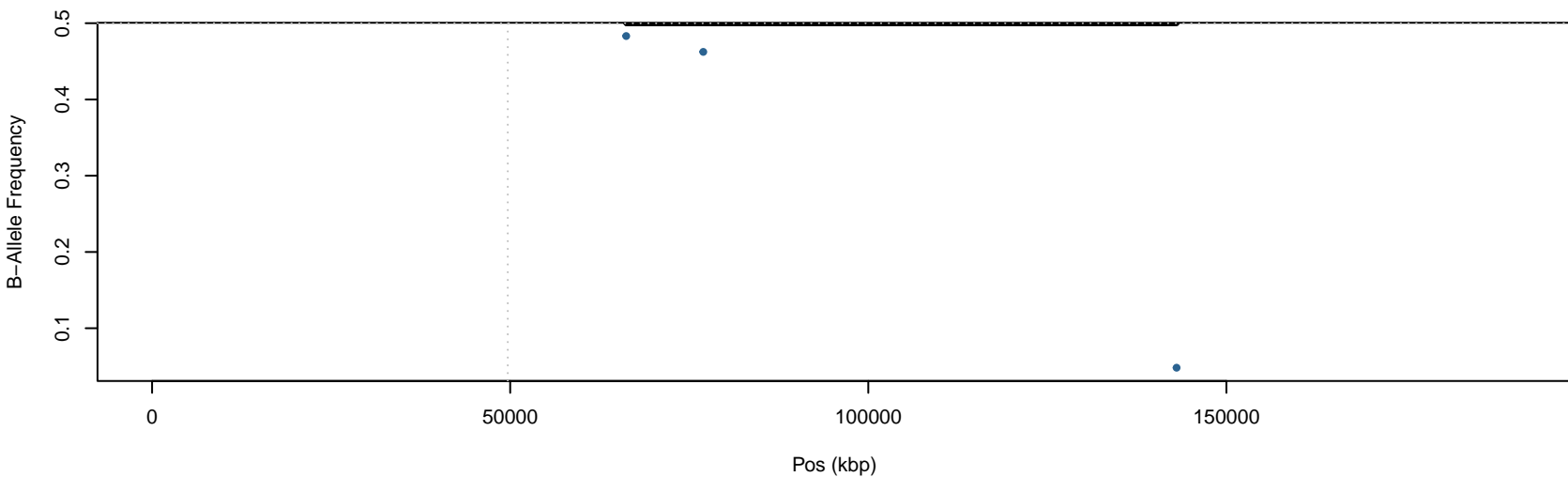
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 1



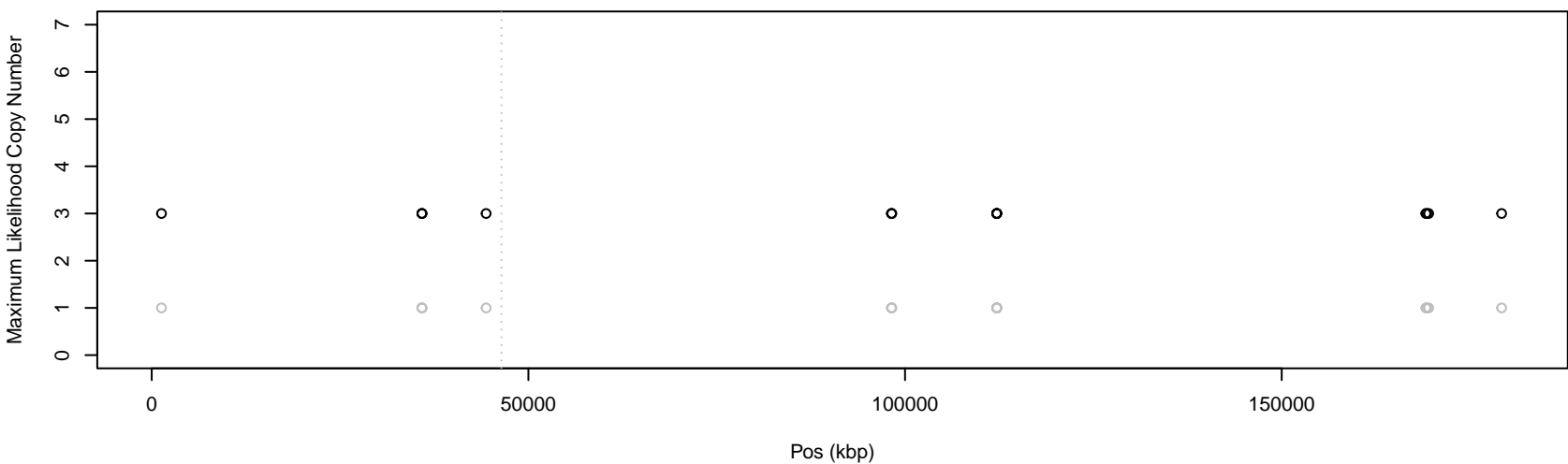
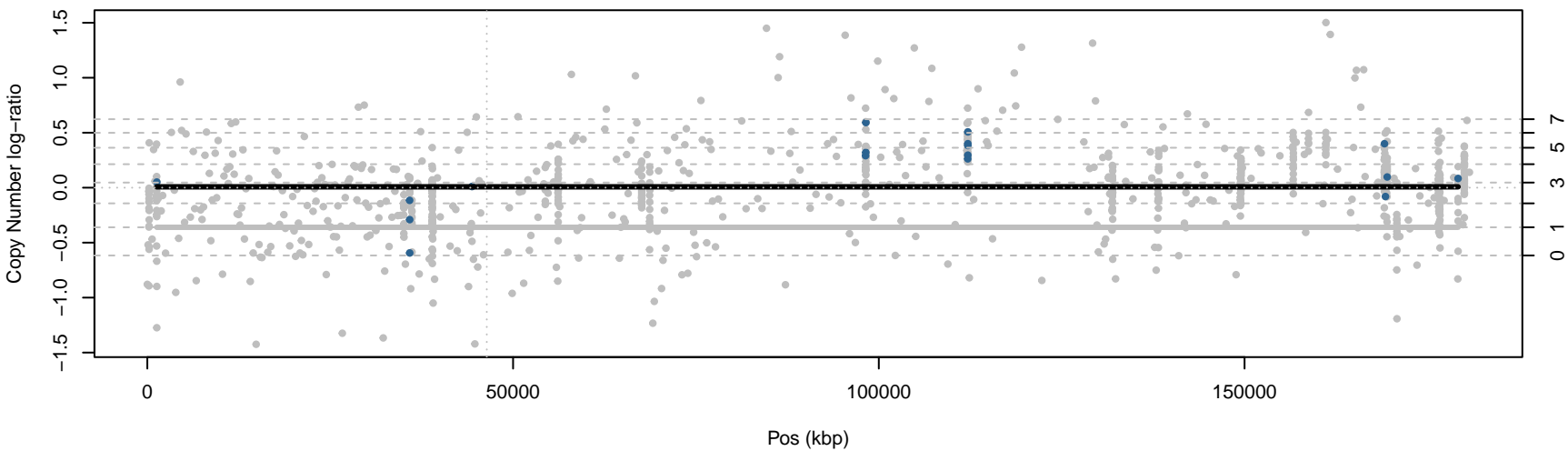
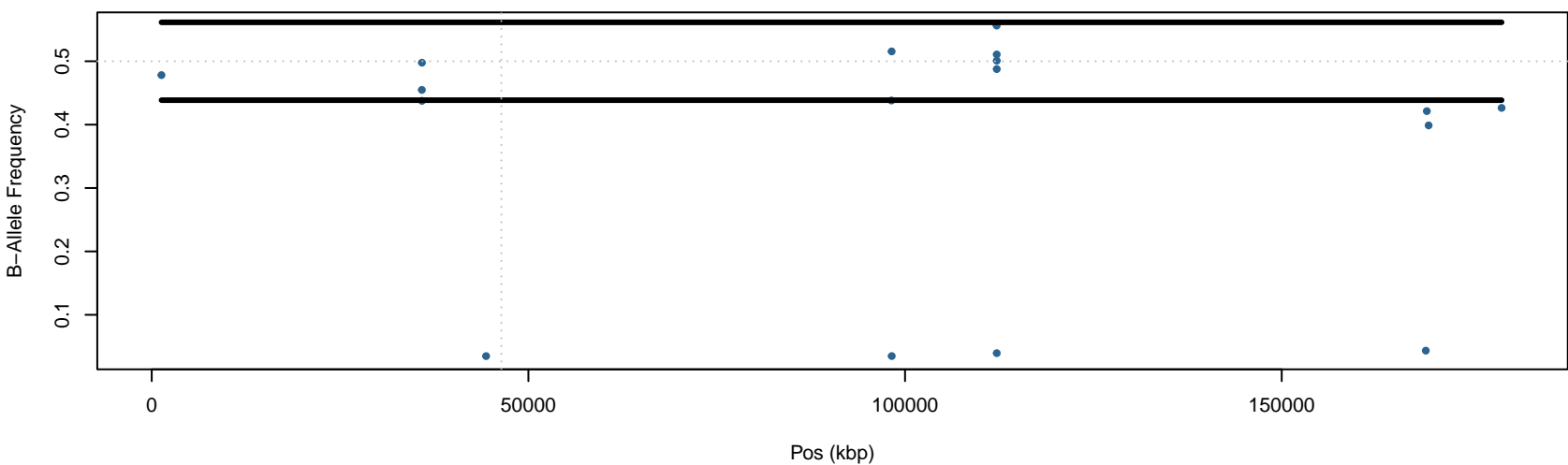
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 3



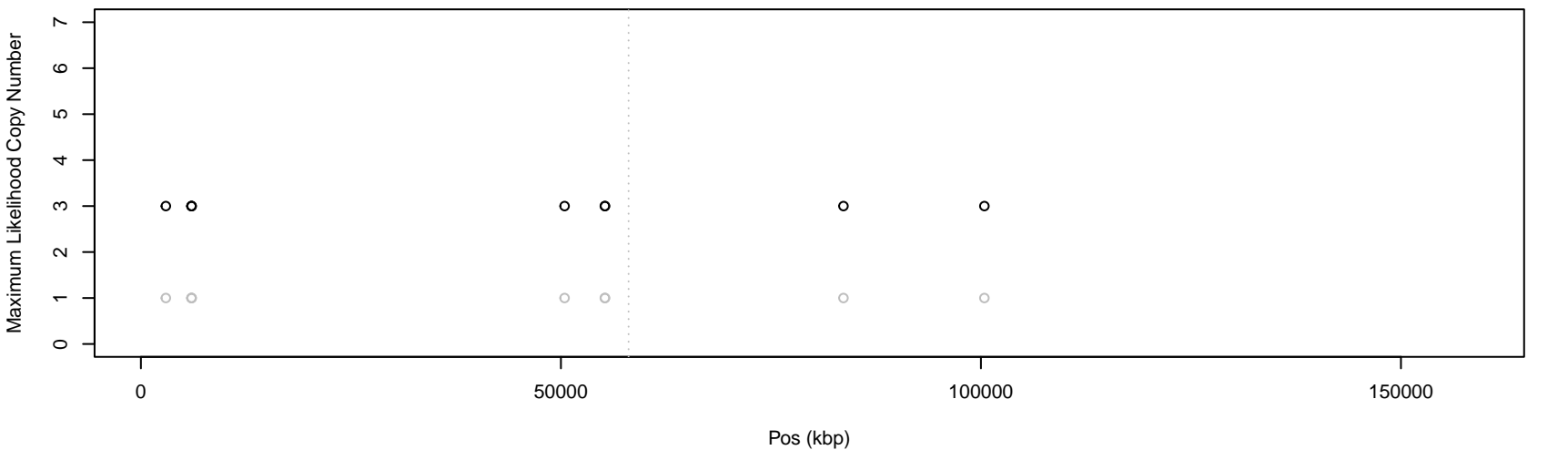
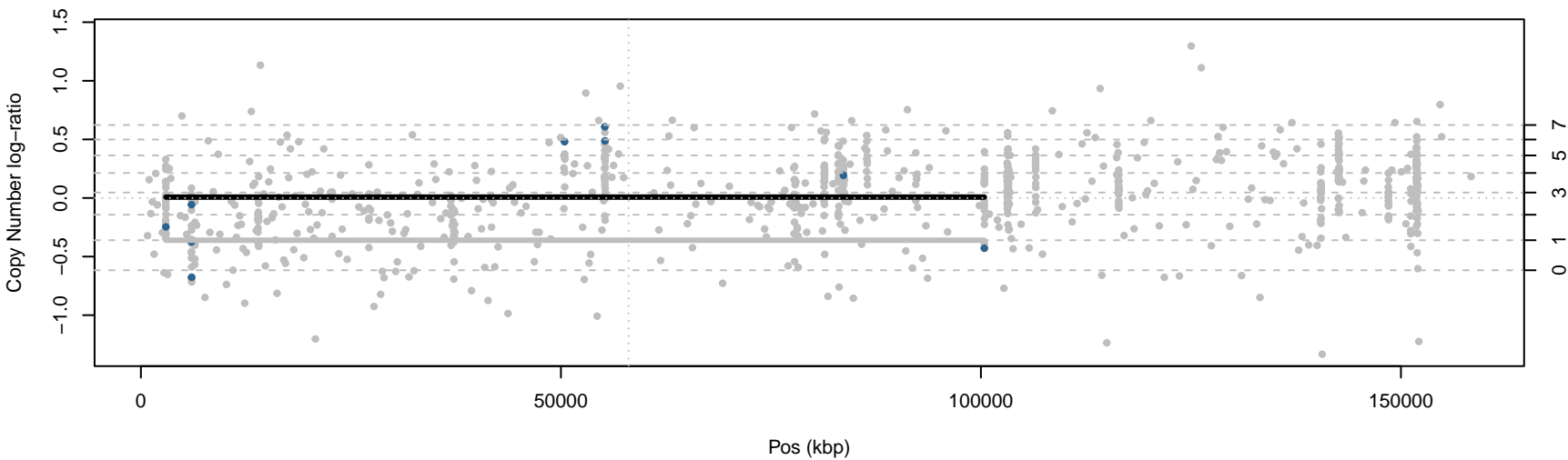
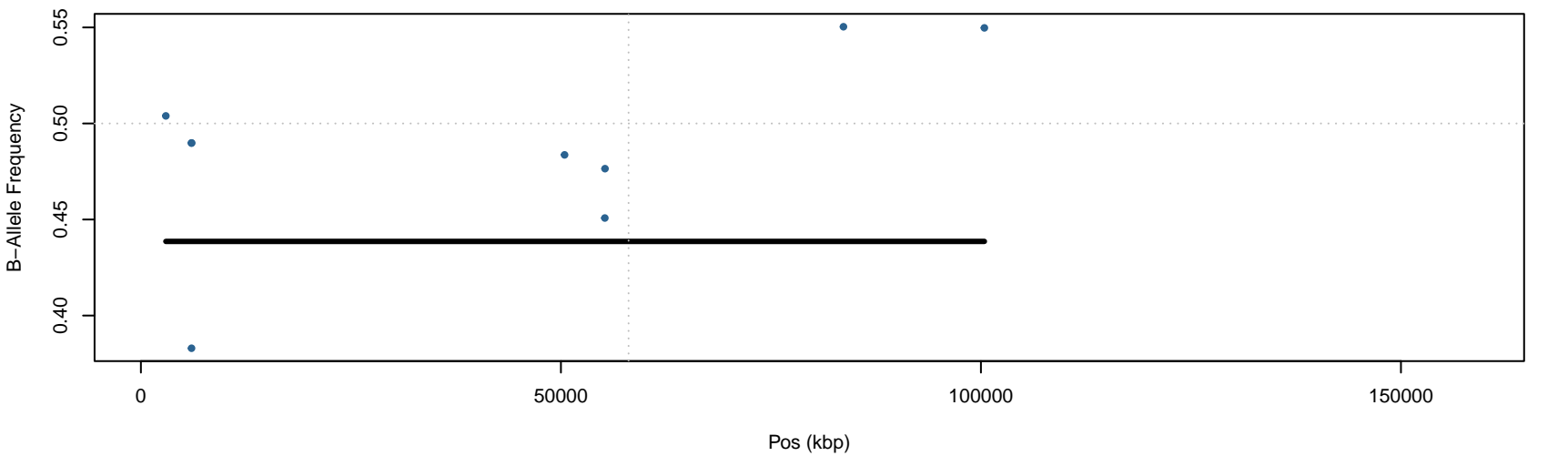
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 4



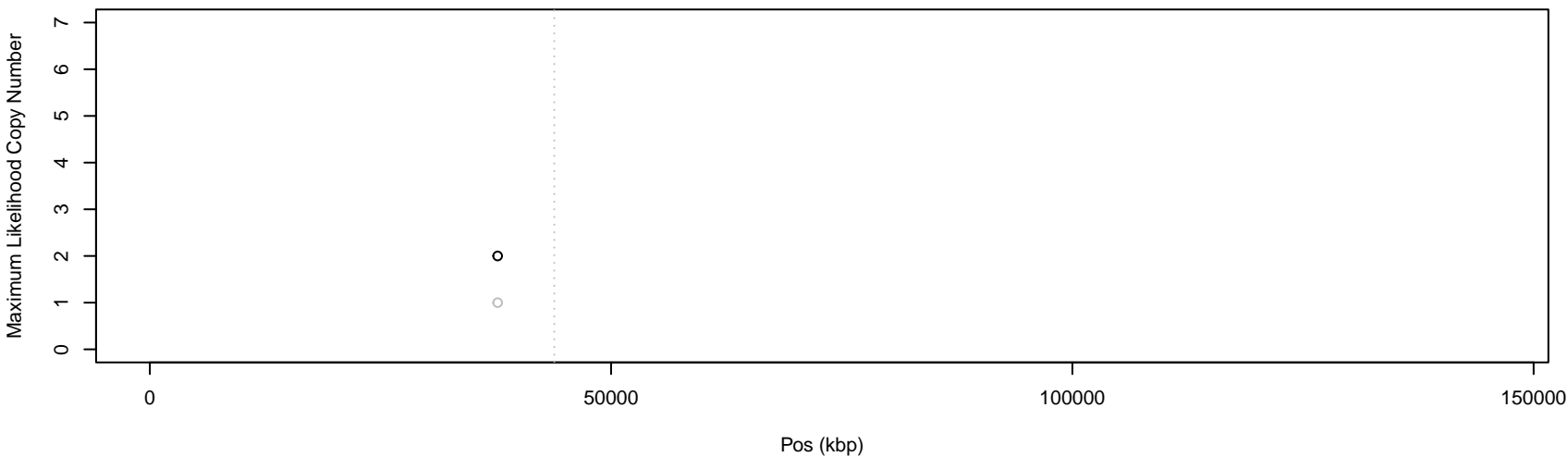
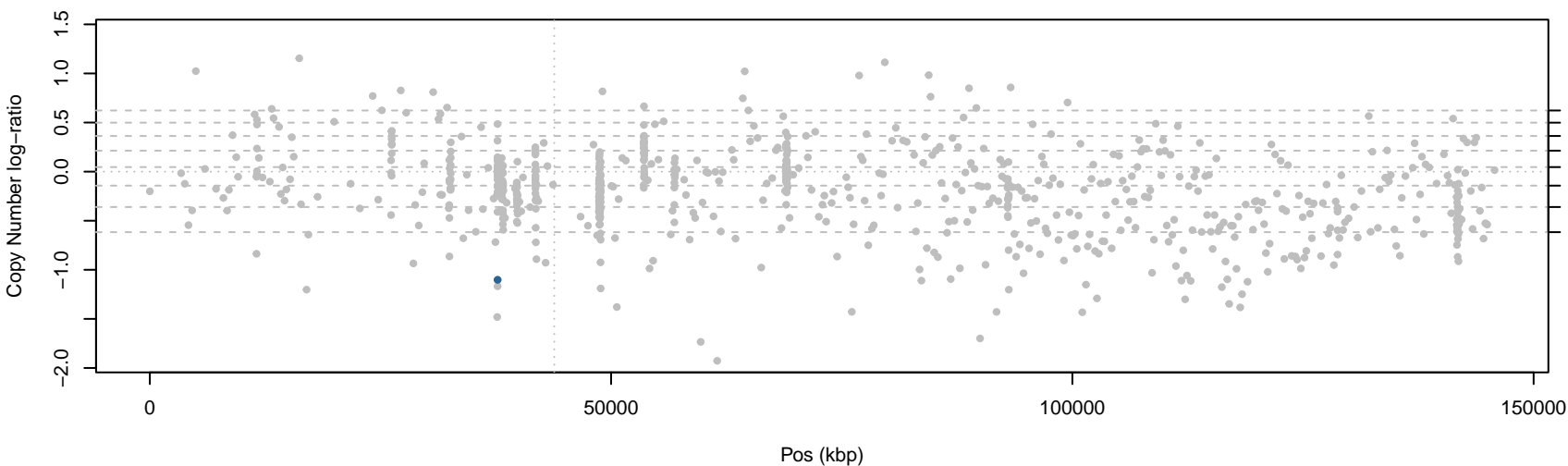
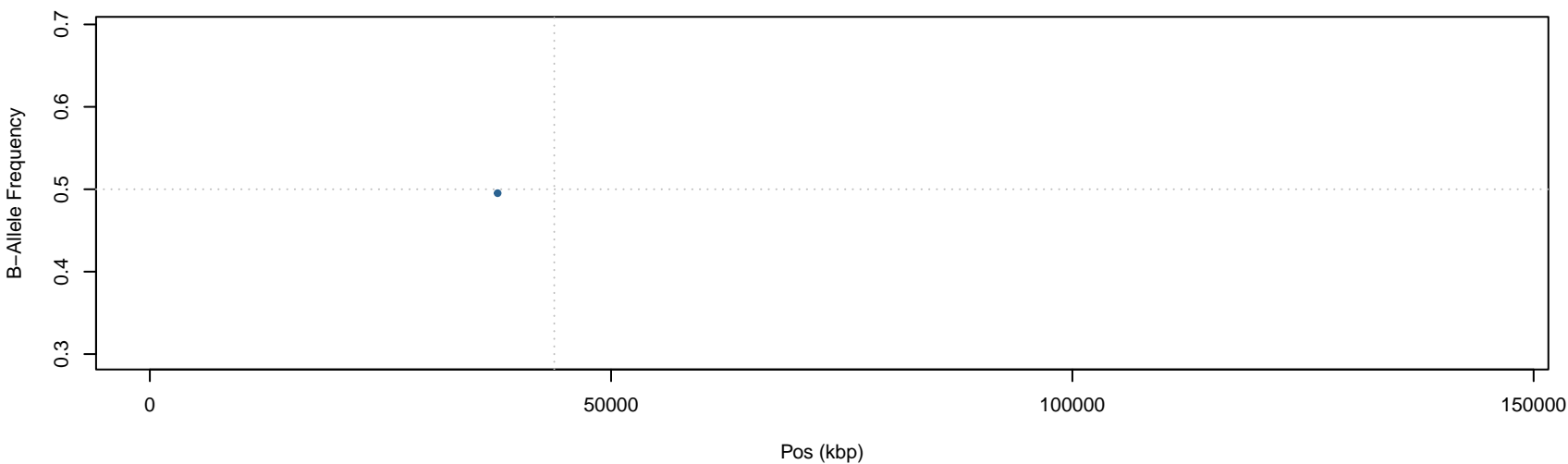
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 5



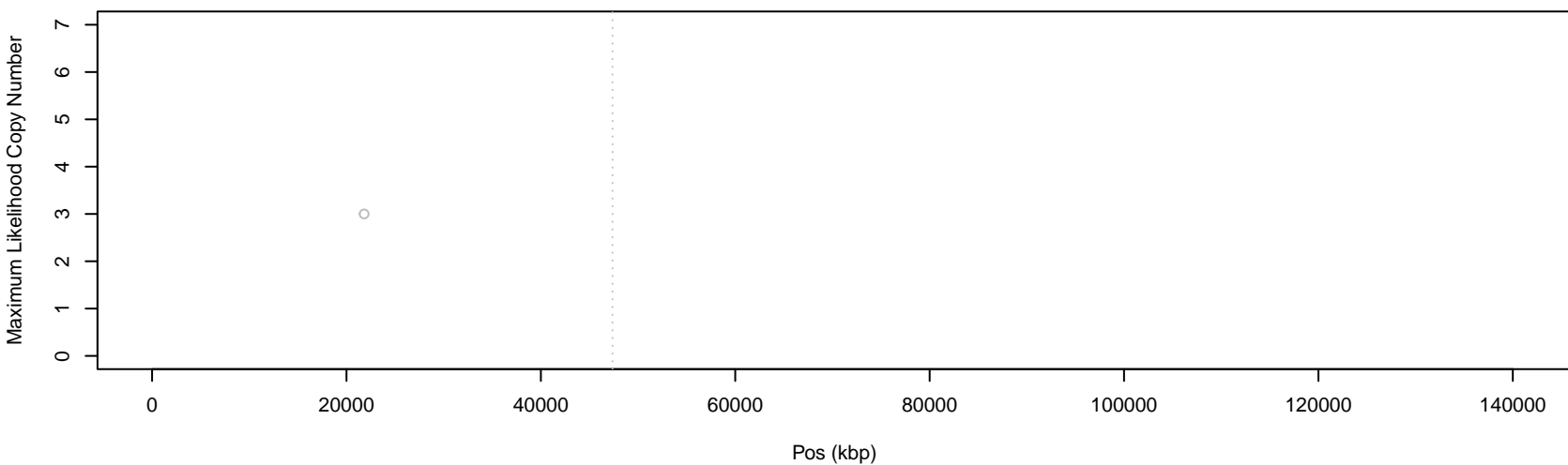
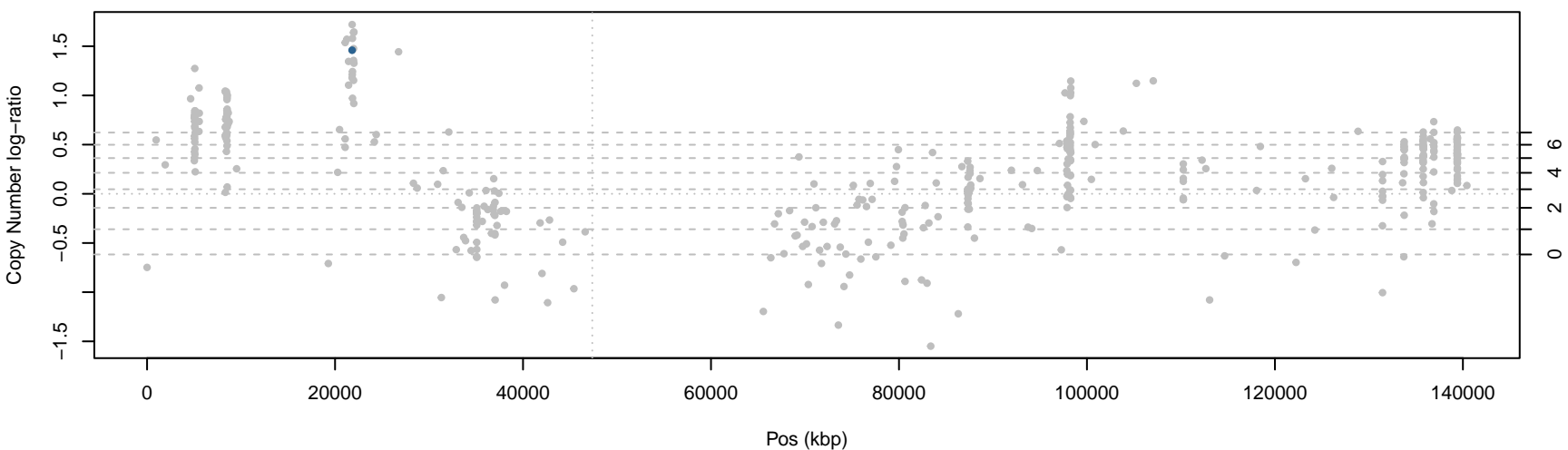
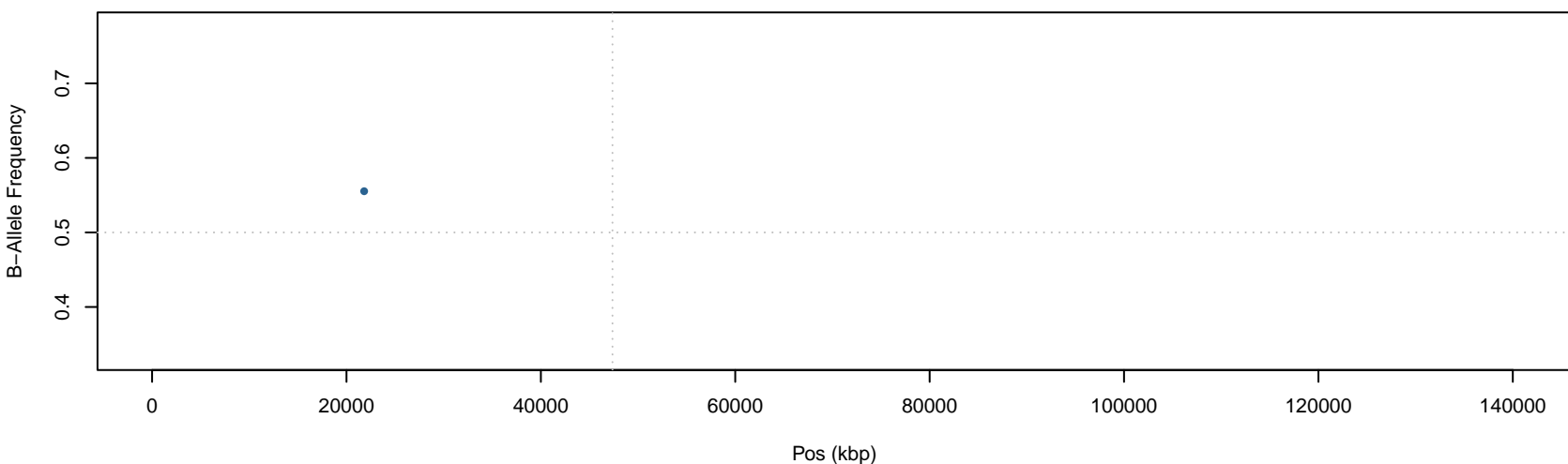
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 7



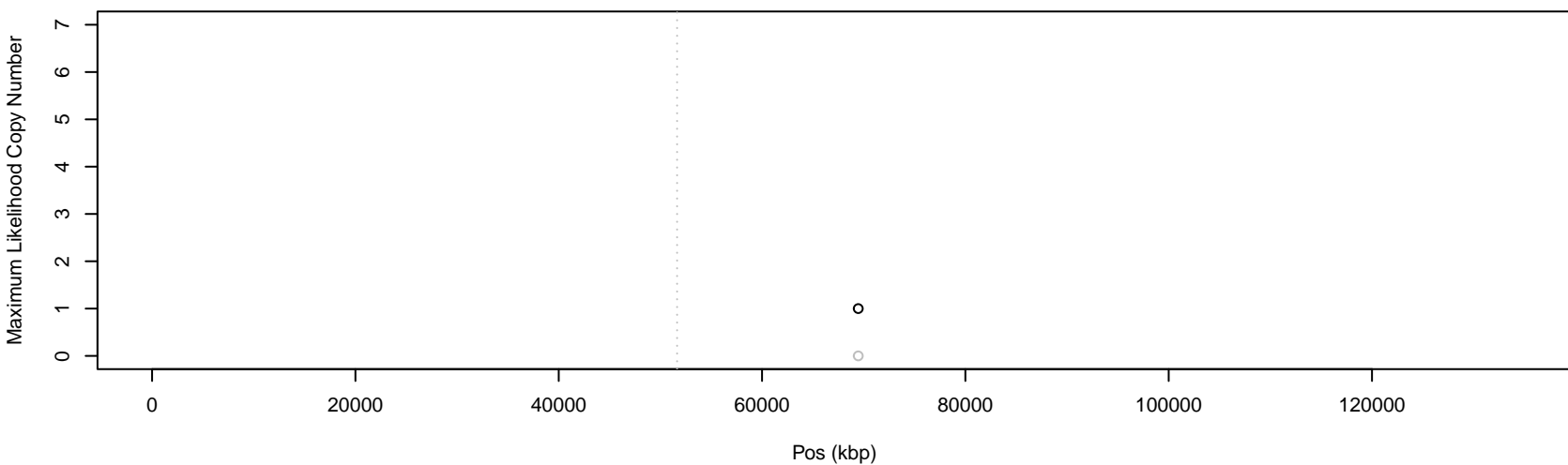
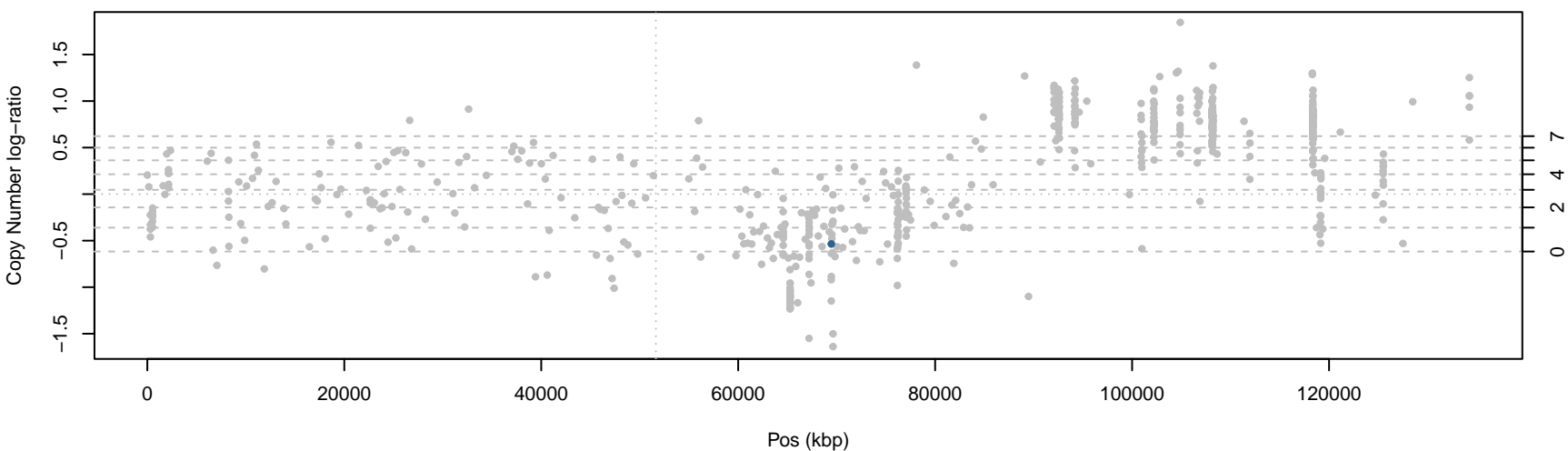
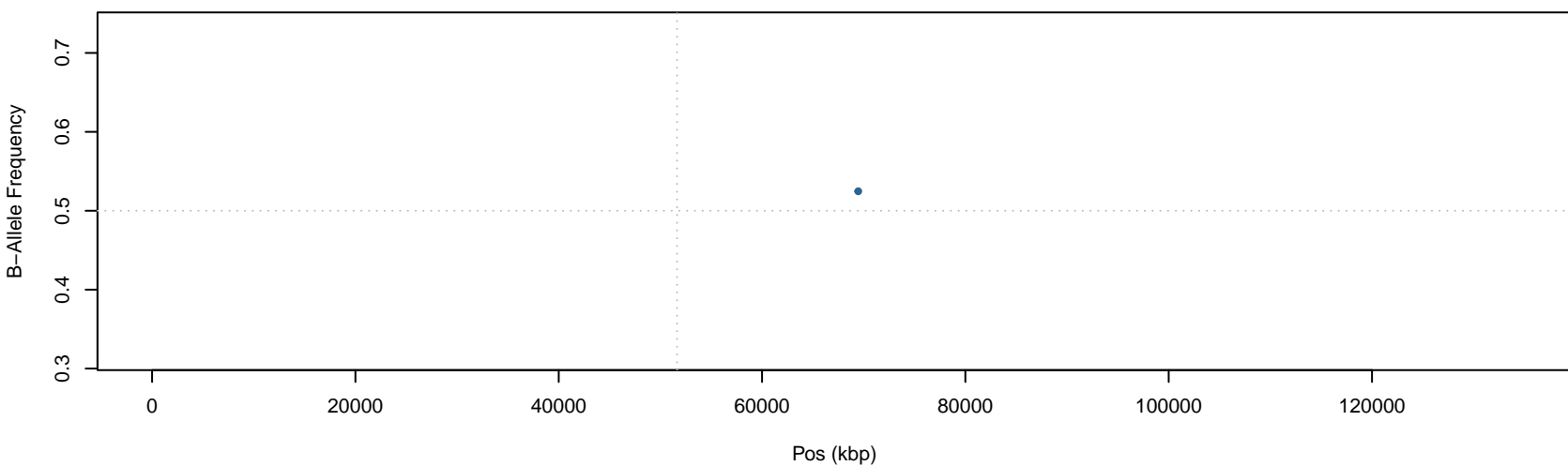
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 8



Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 9

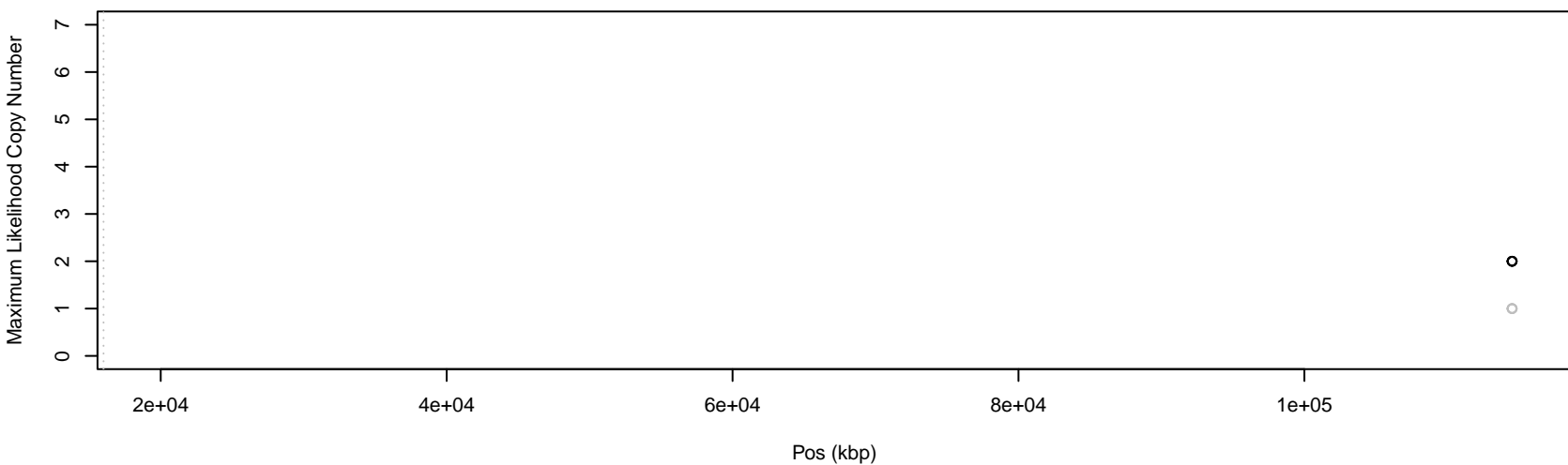
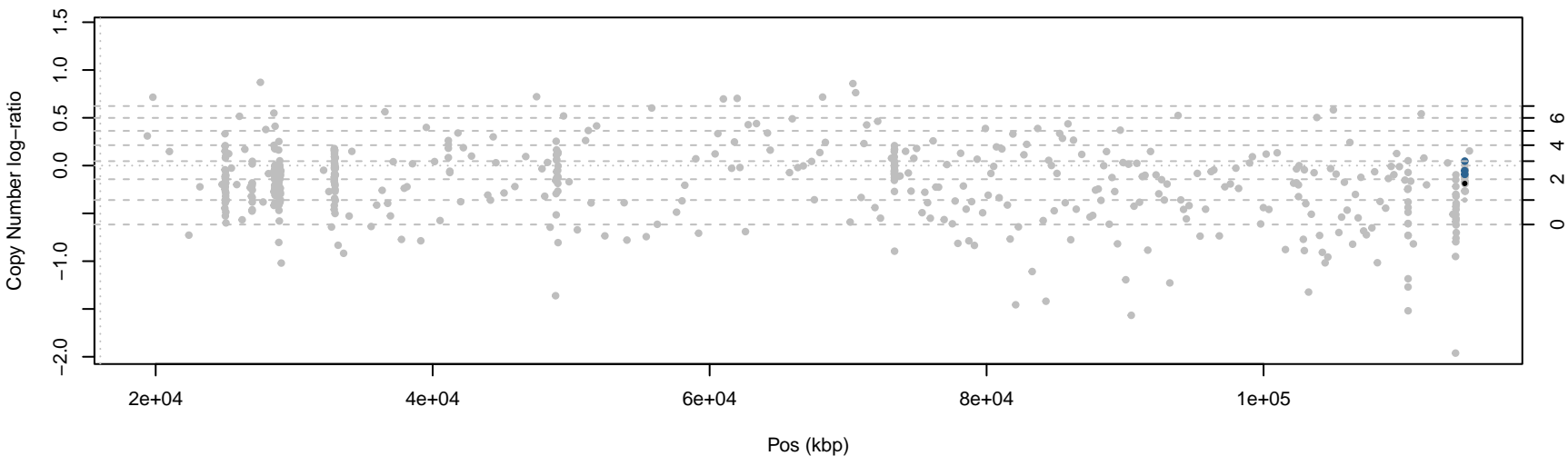
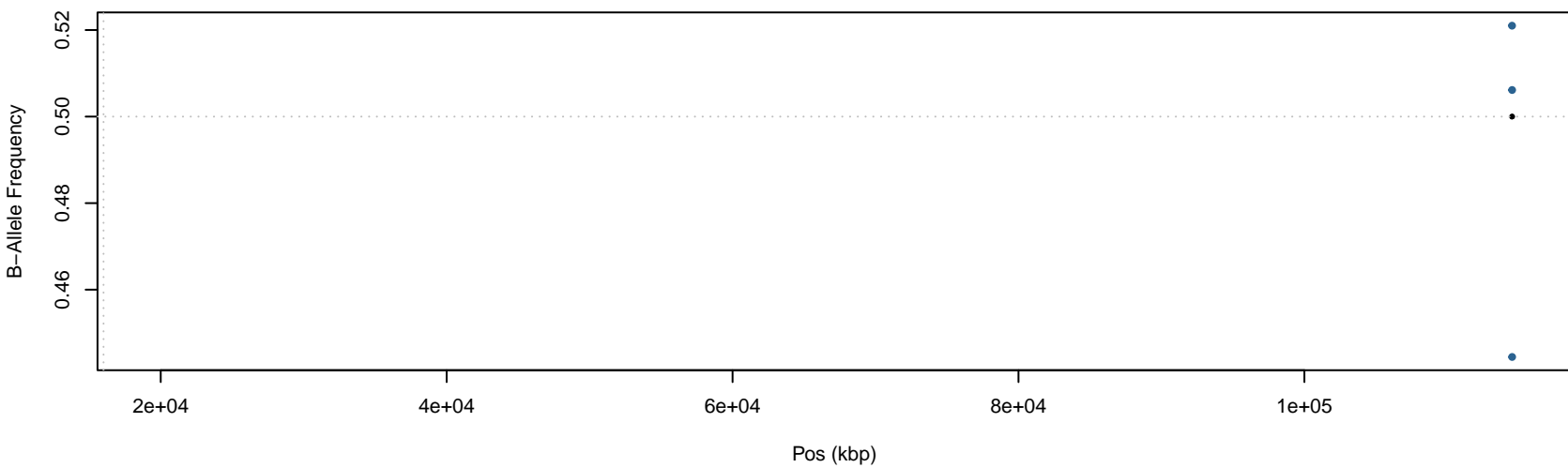


Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 11

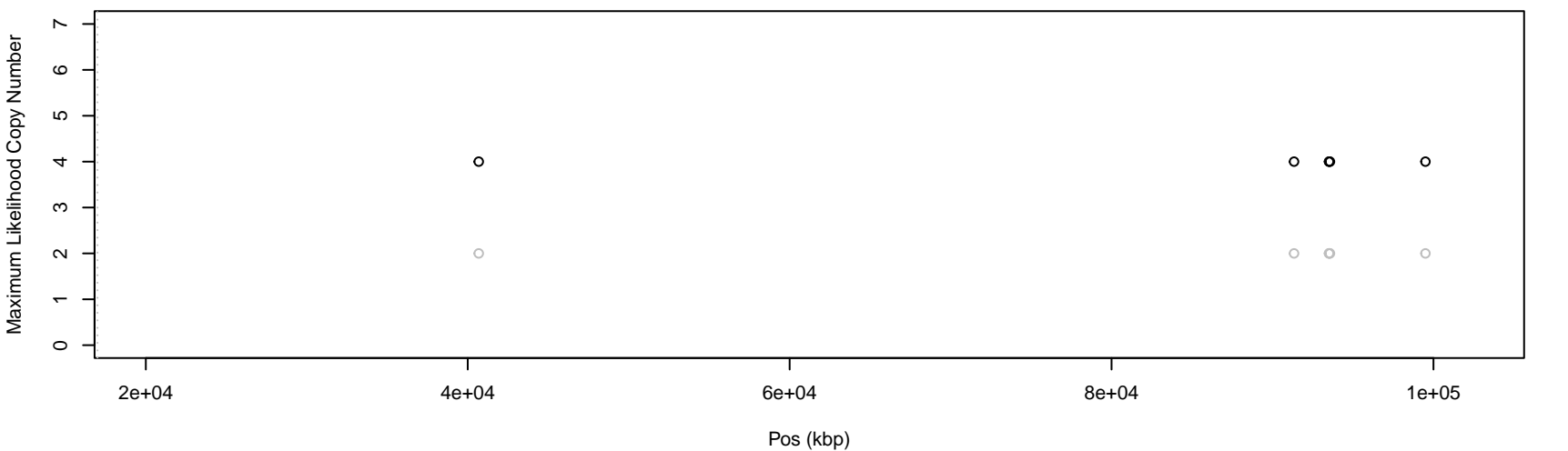
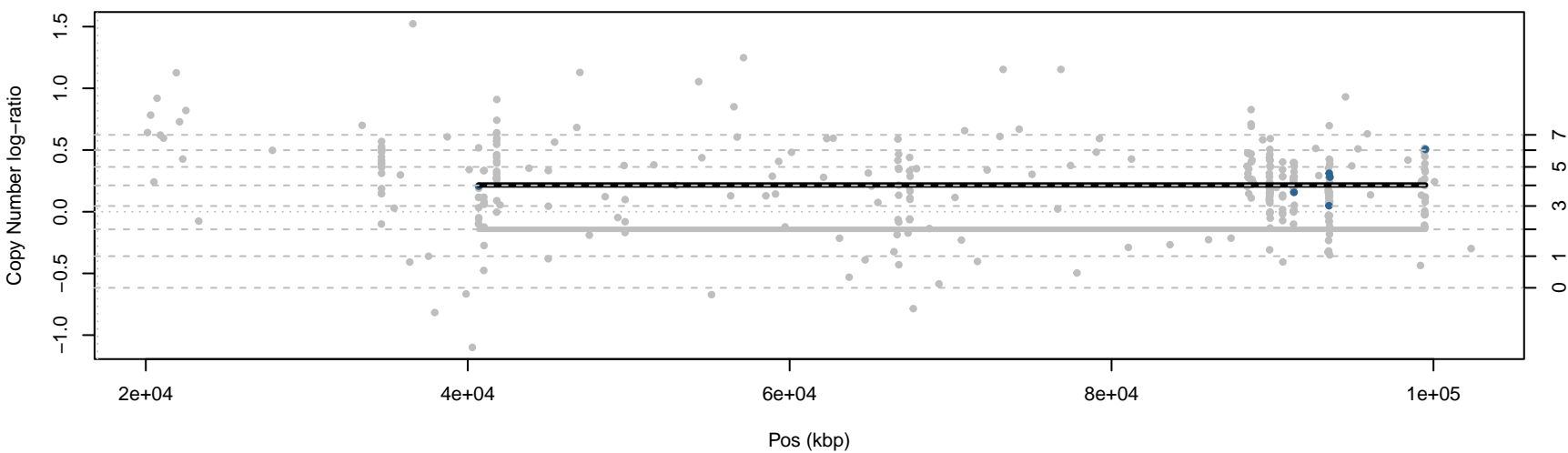
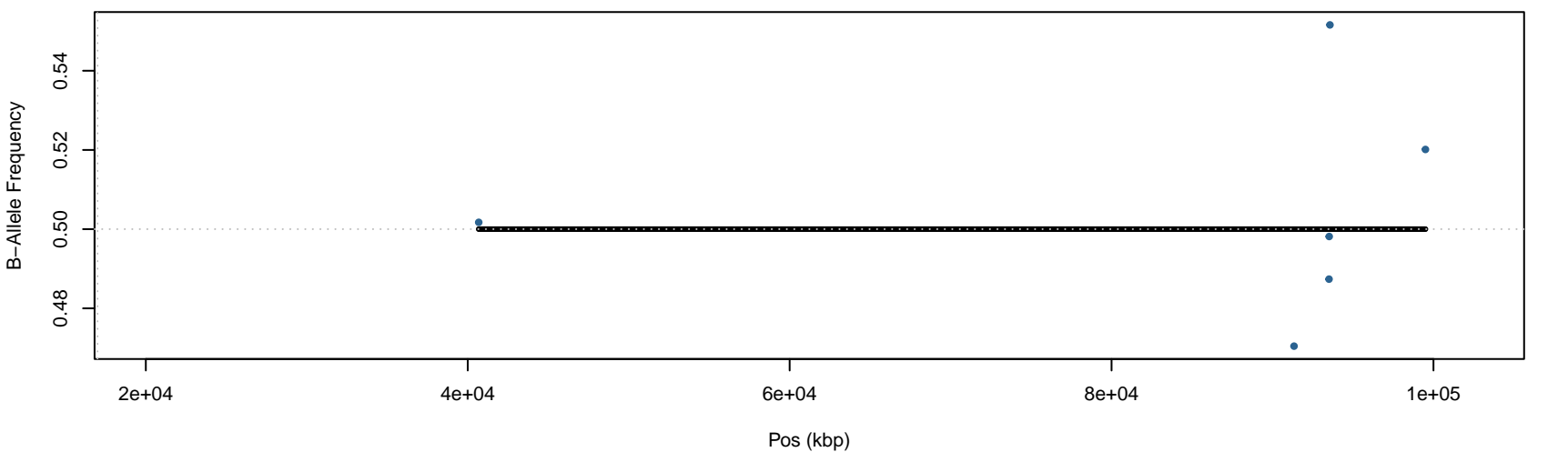




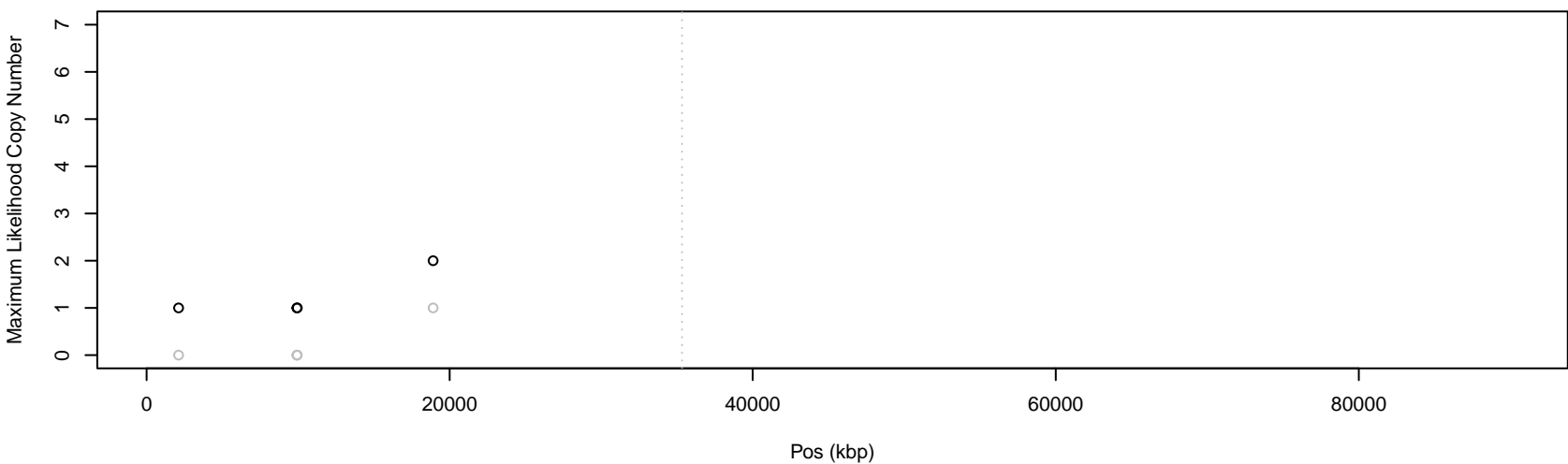
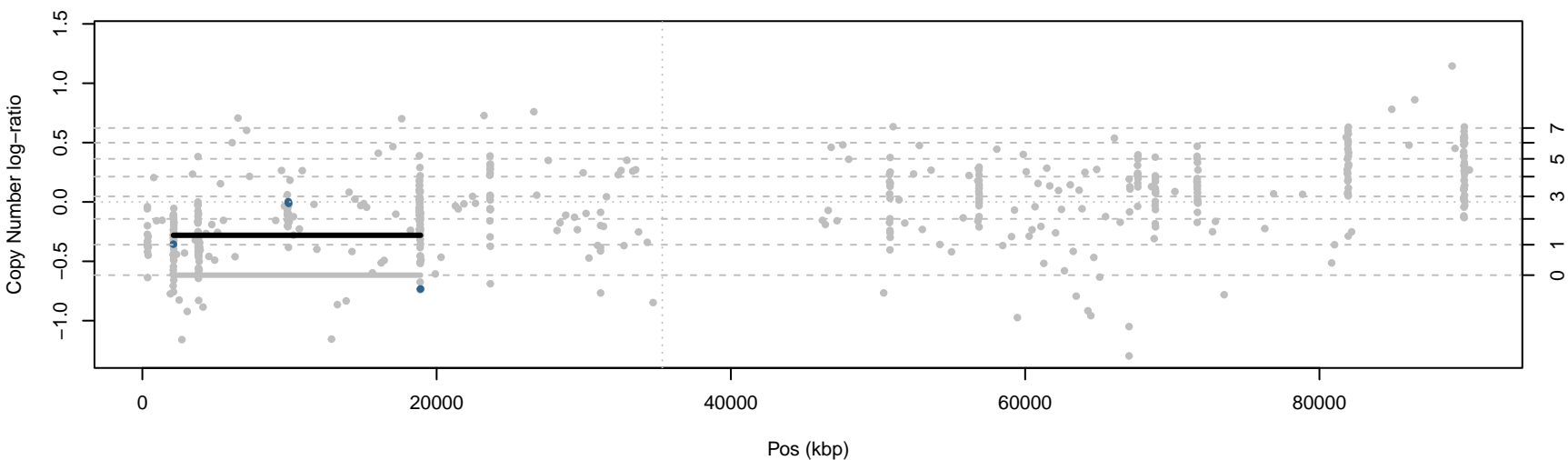
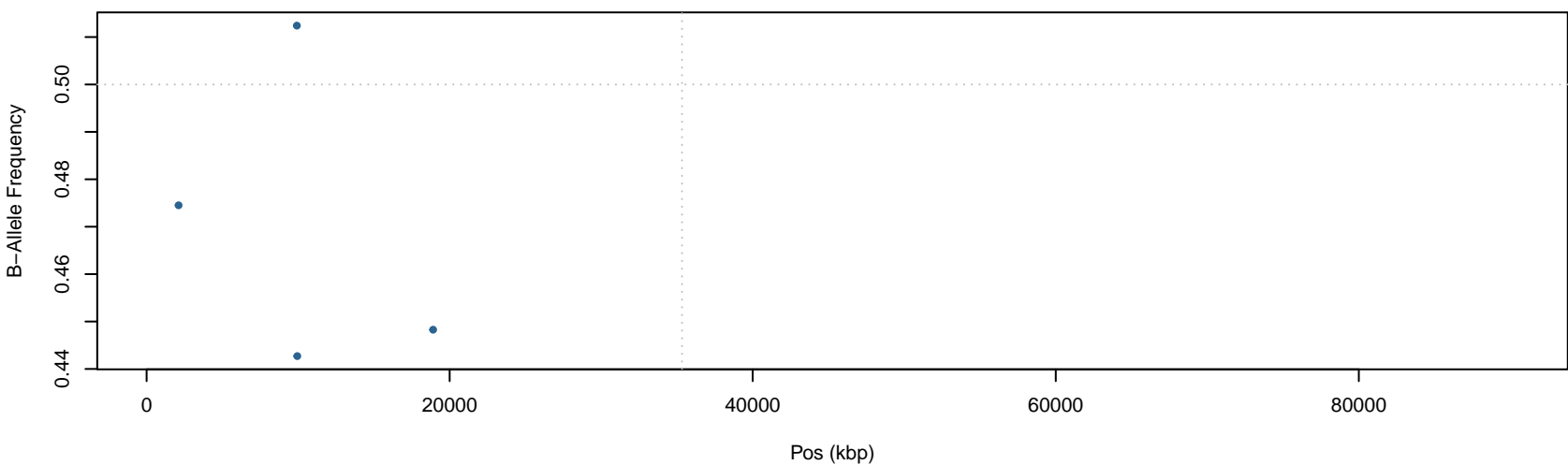
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 13



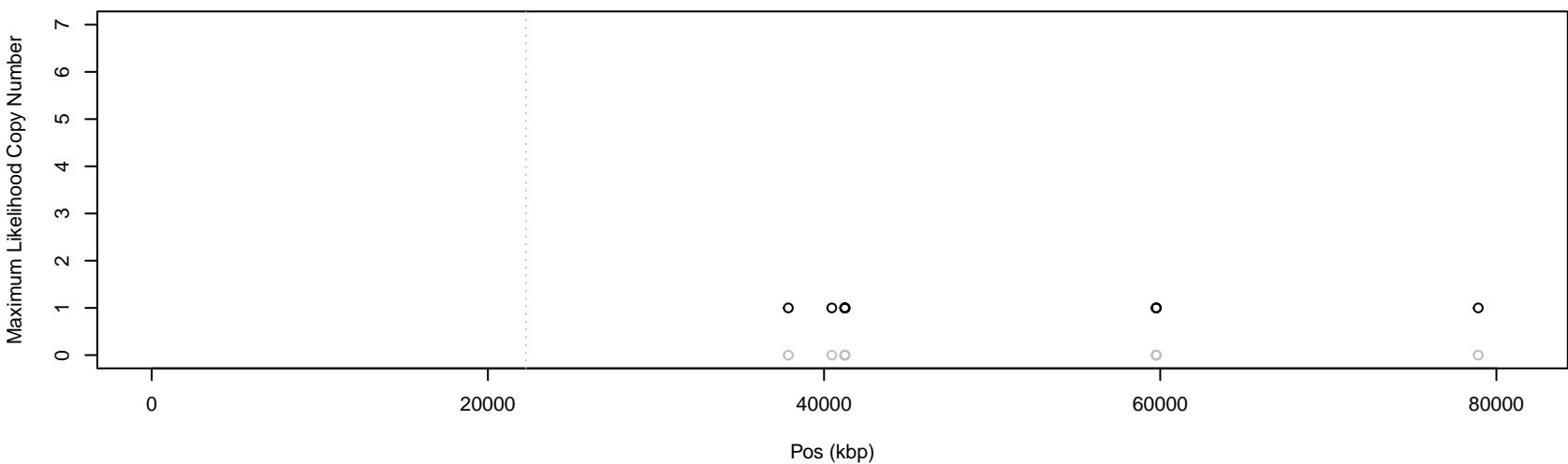
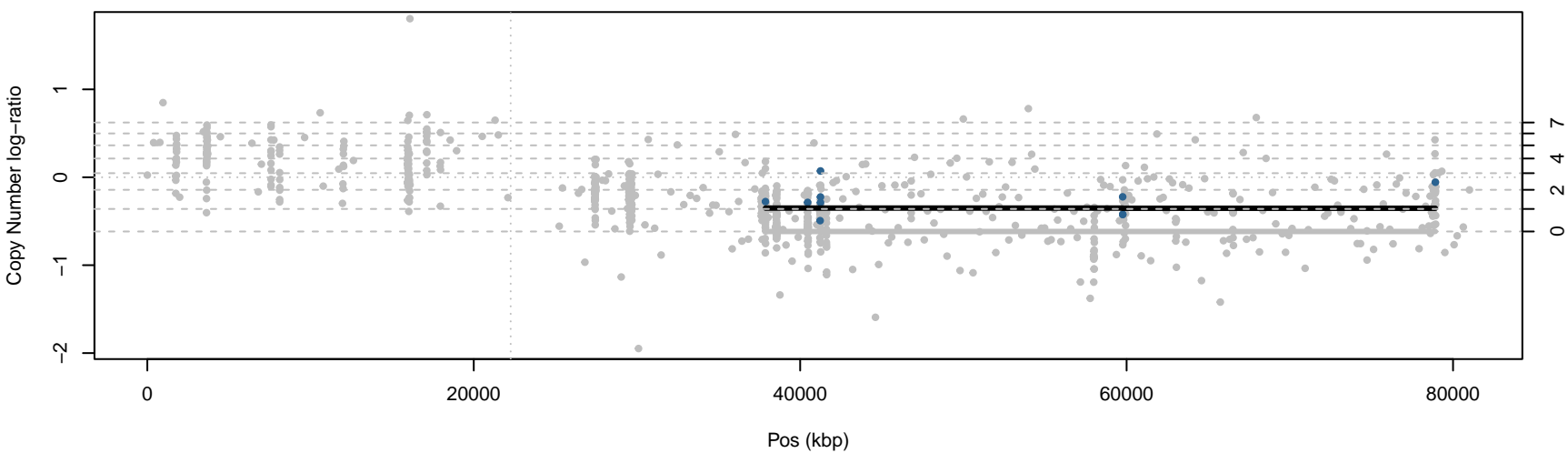
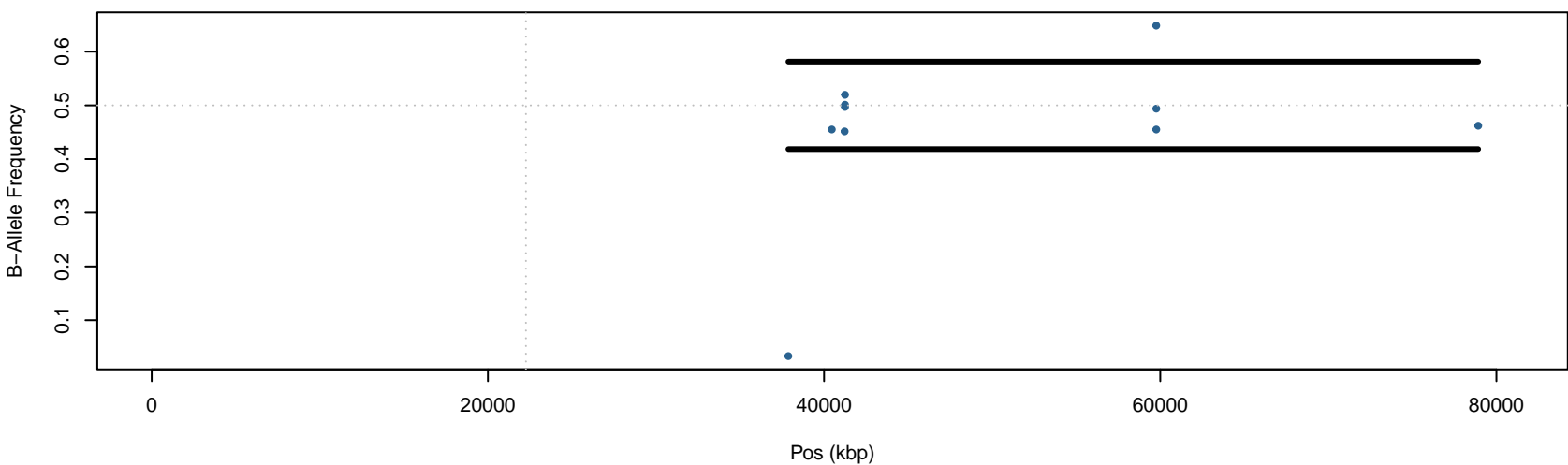
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 15



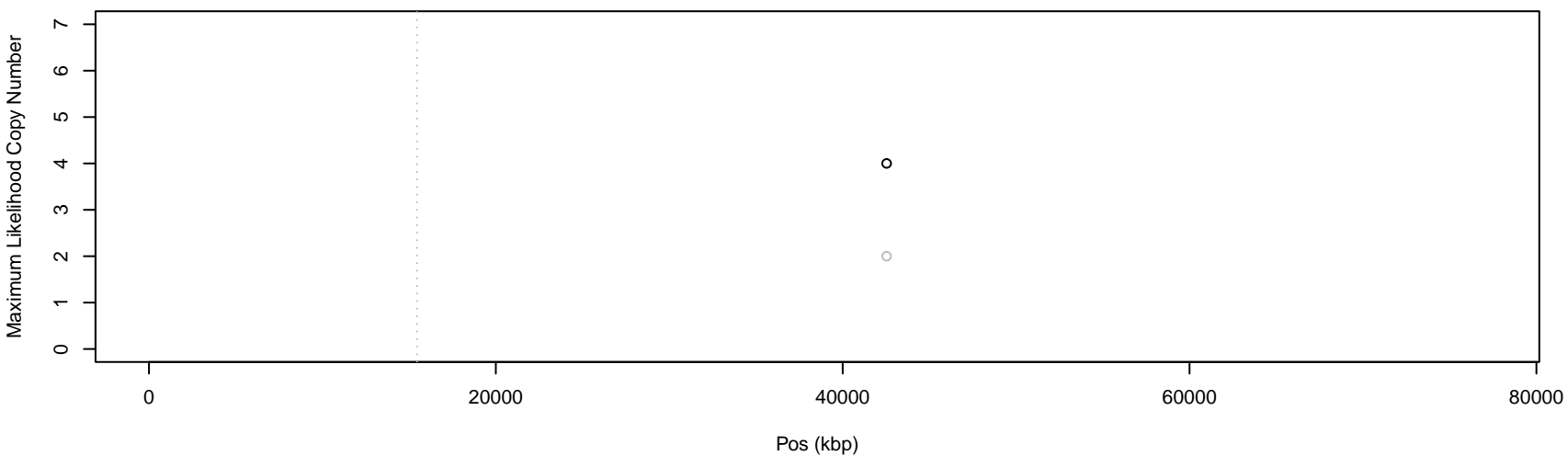
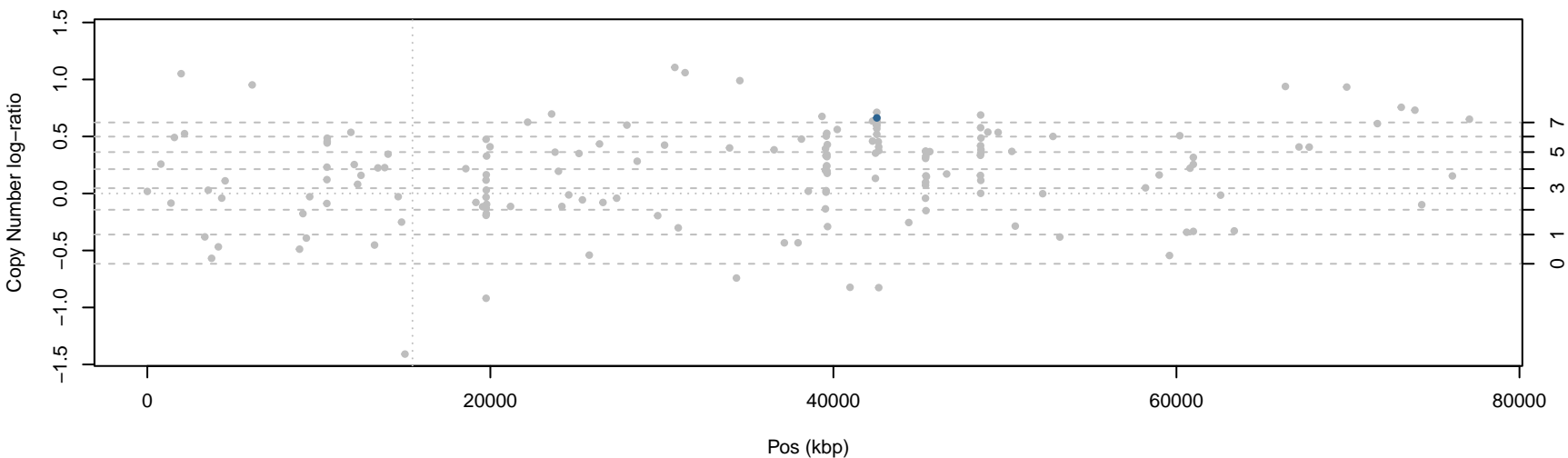
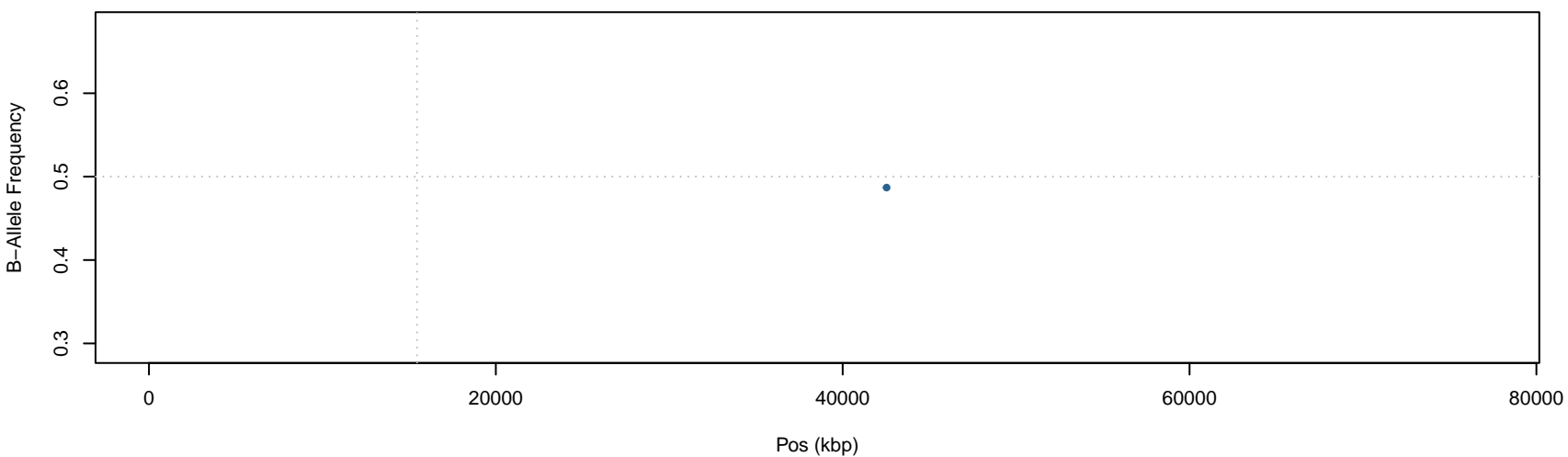
Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 16



Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 17



Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 18



Purity: 0.28 Tumor ploidy: 2.743 SNV log-likelihood: -39.28 GoF: 91.4% Mean coverage: 290;505 Chromosome: 19

