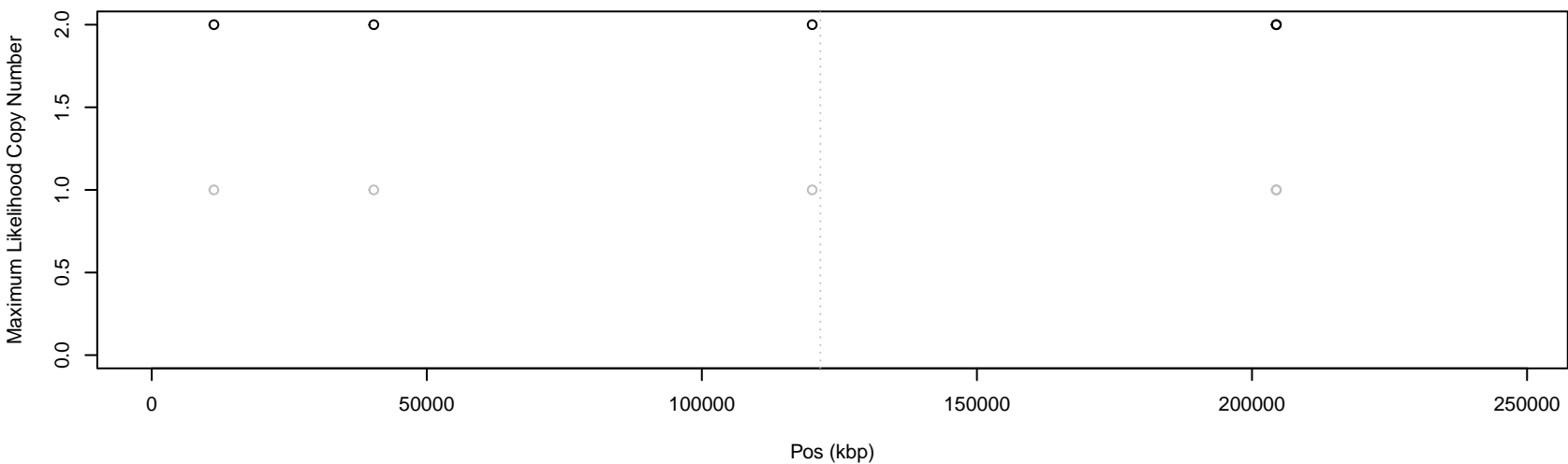
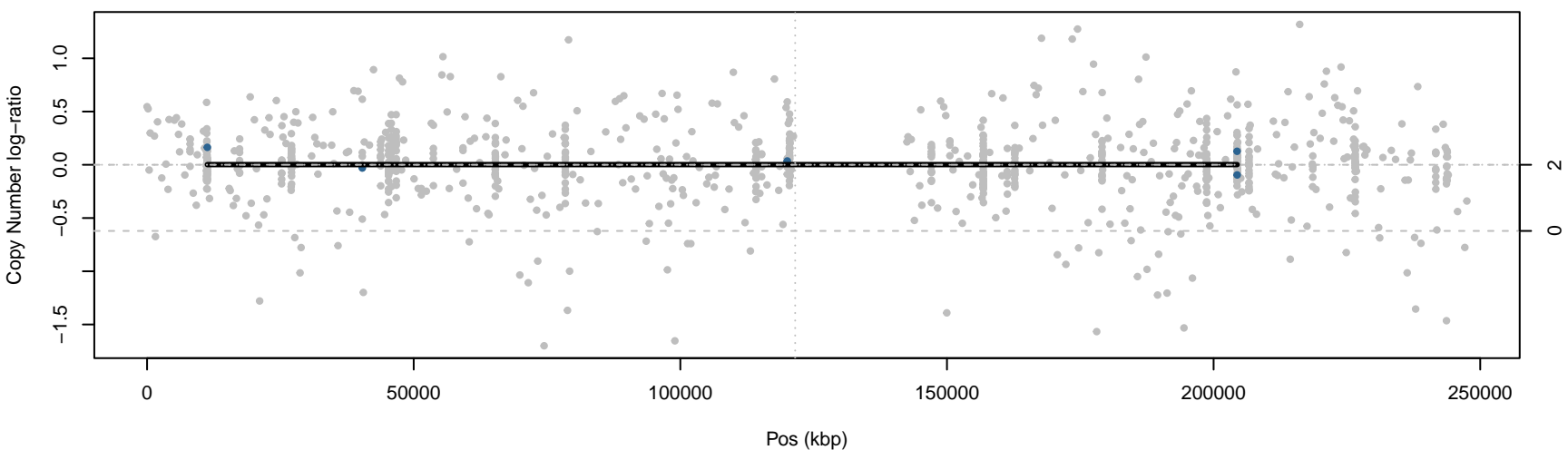
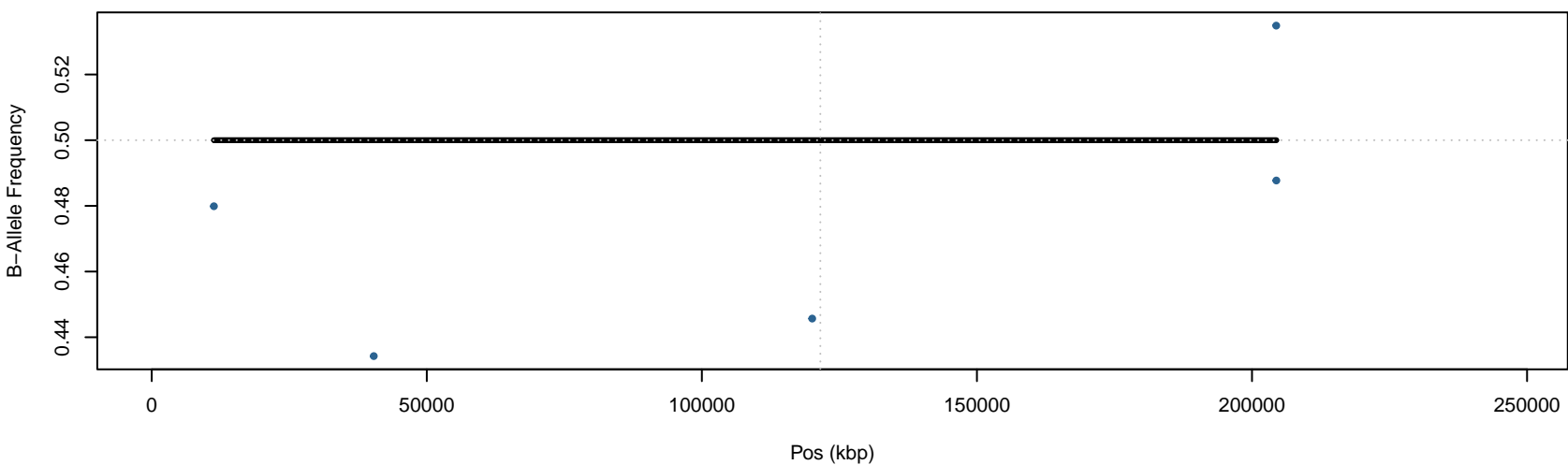
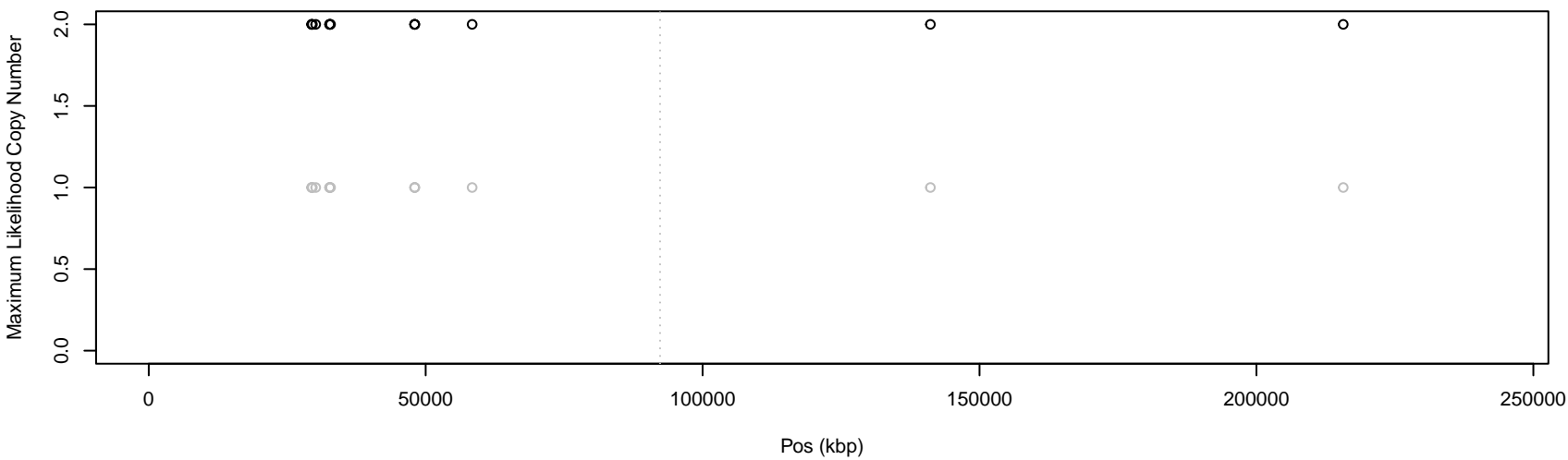
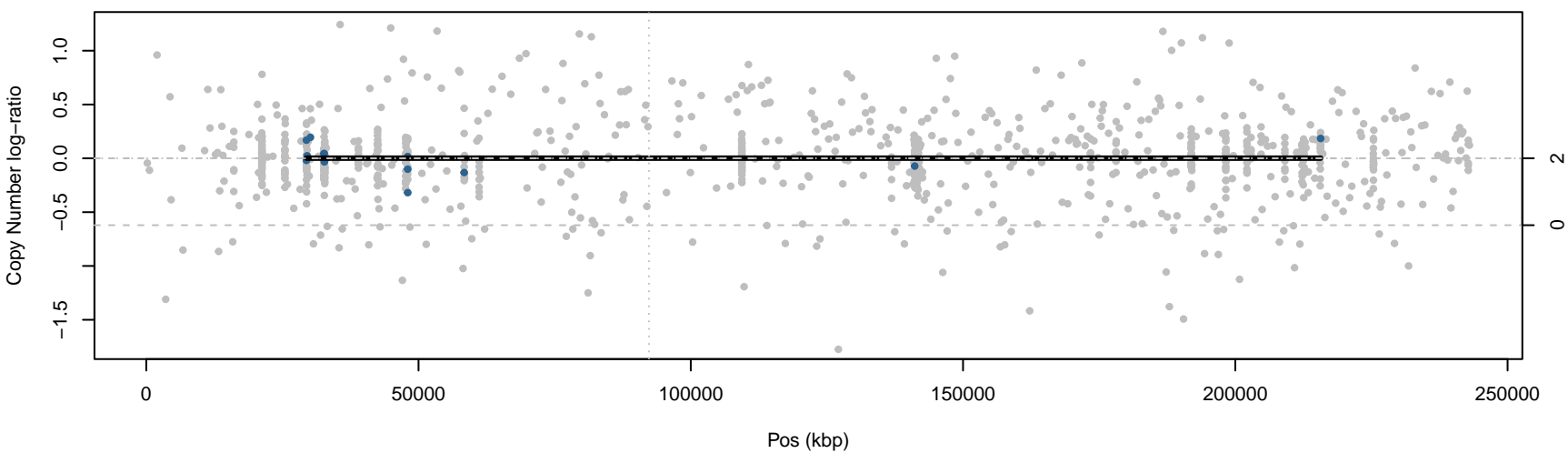
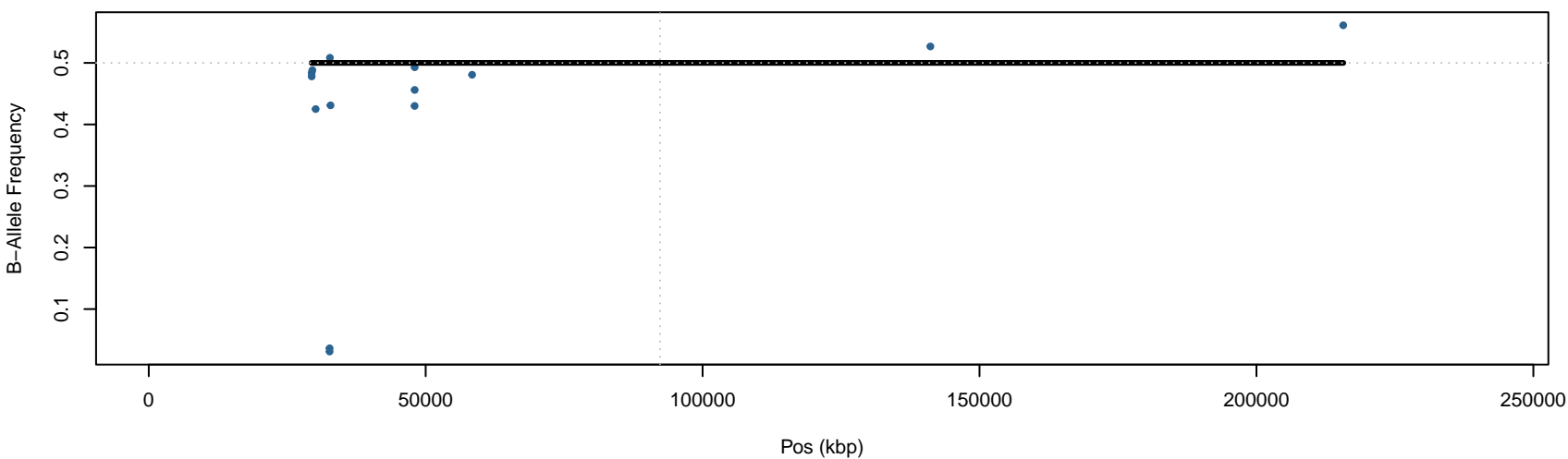


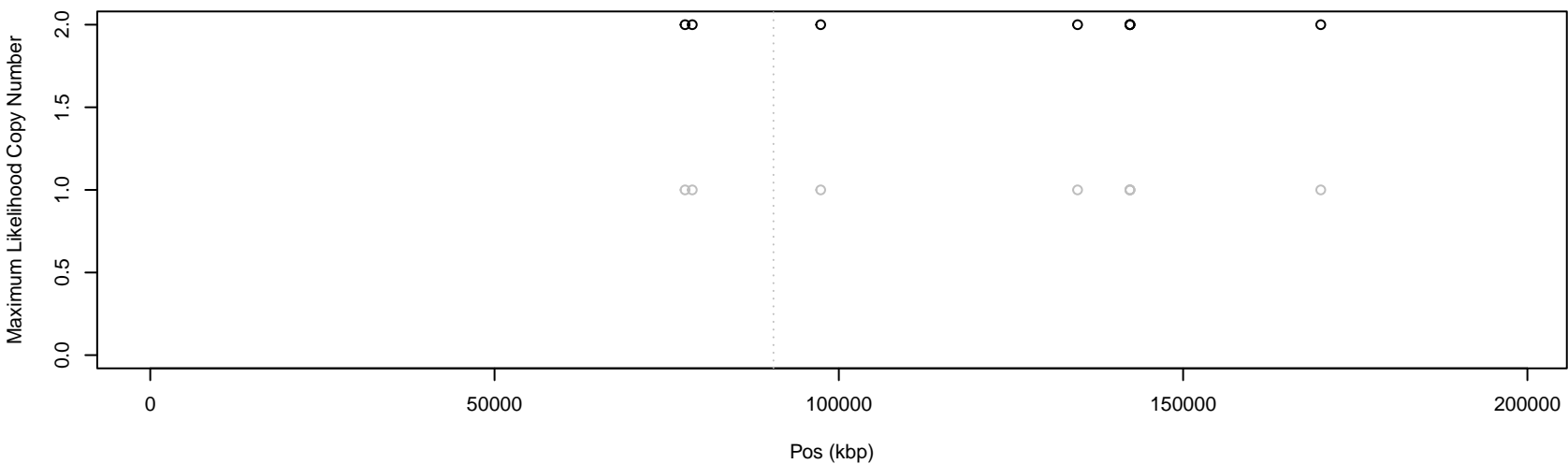
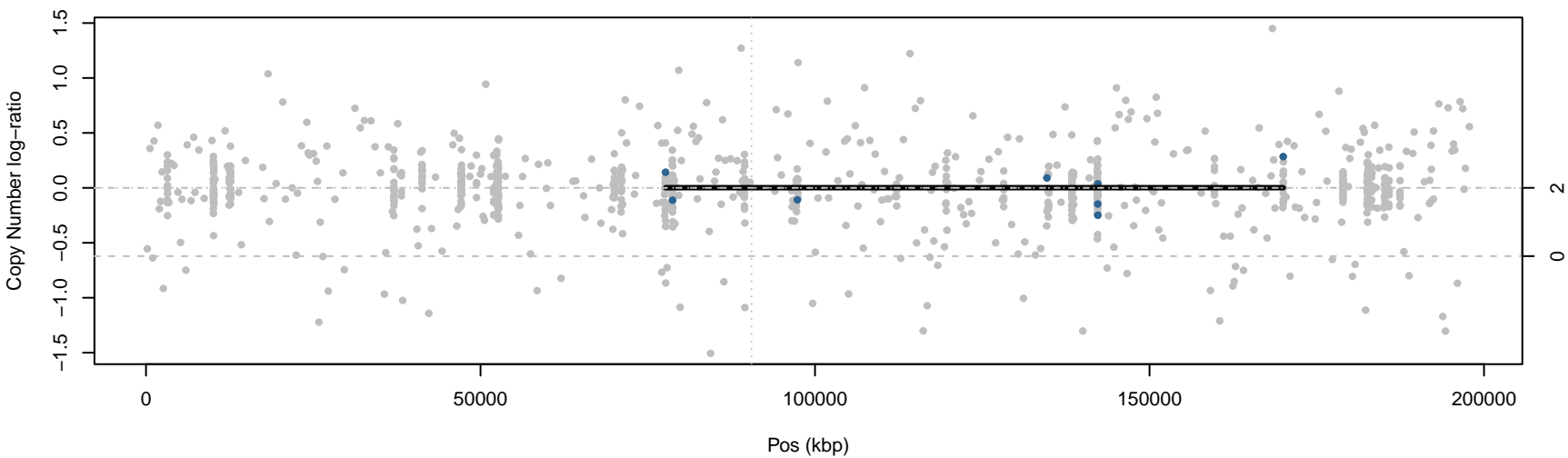
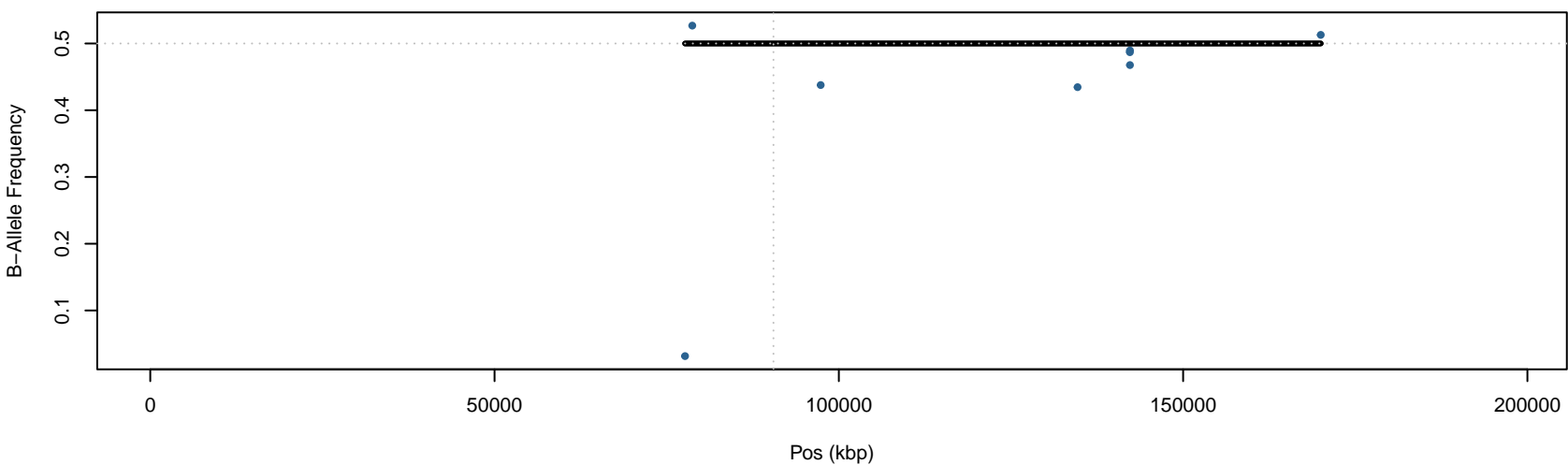
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 1



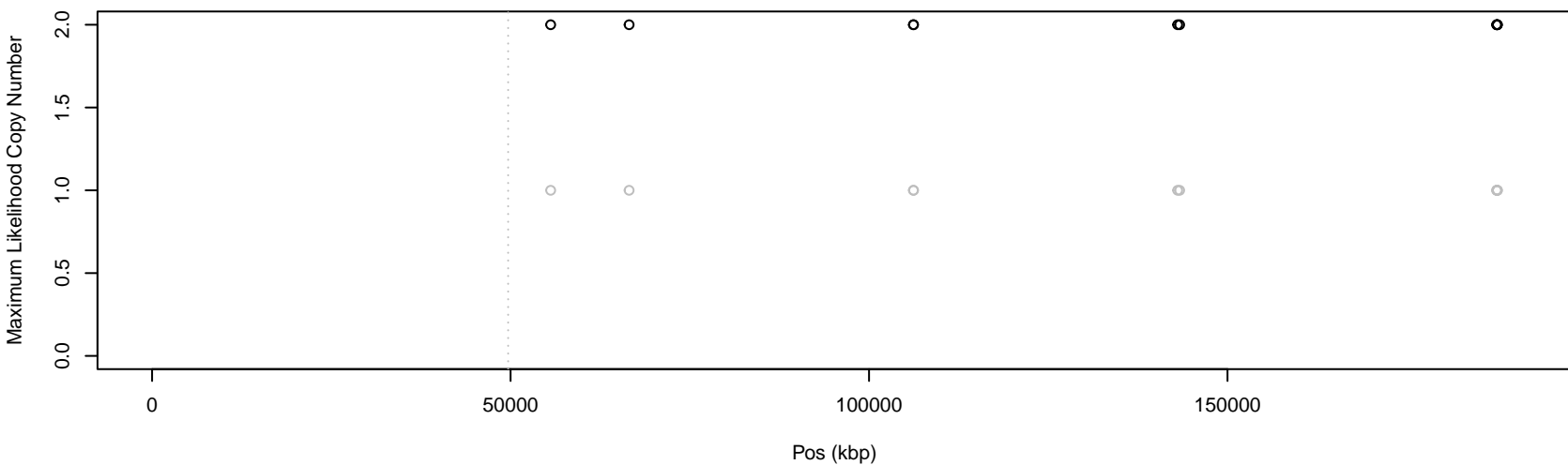
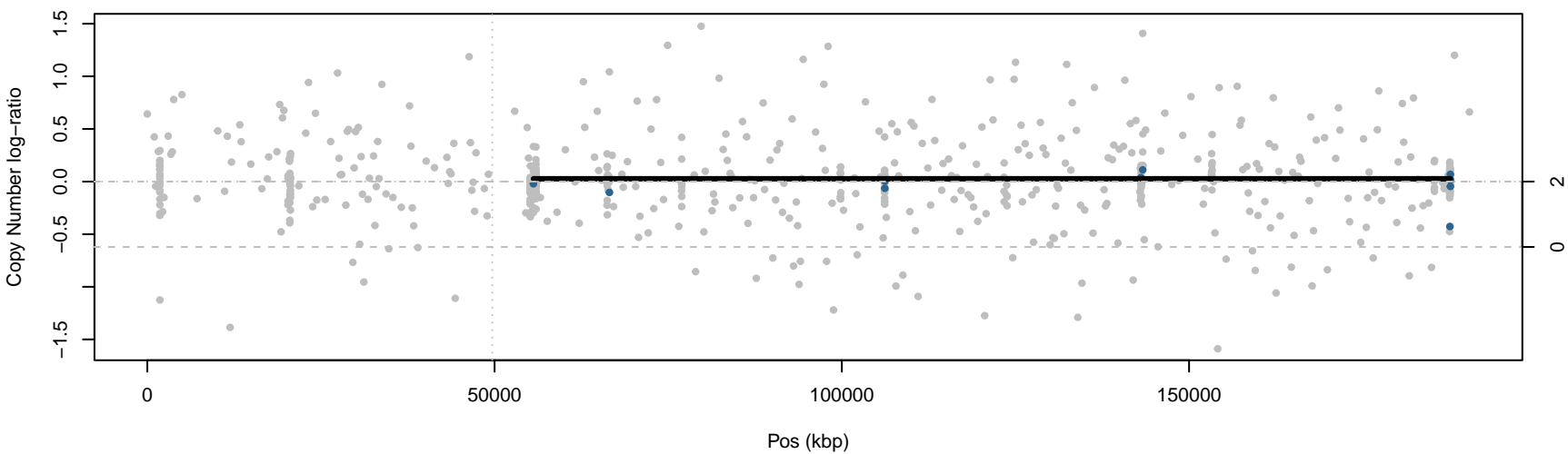
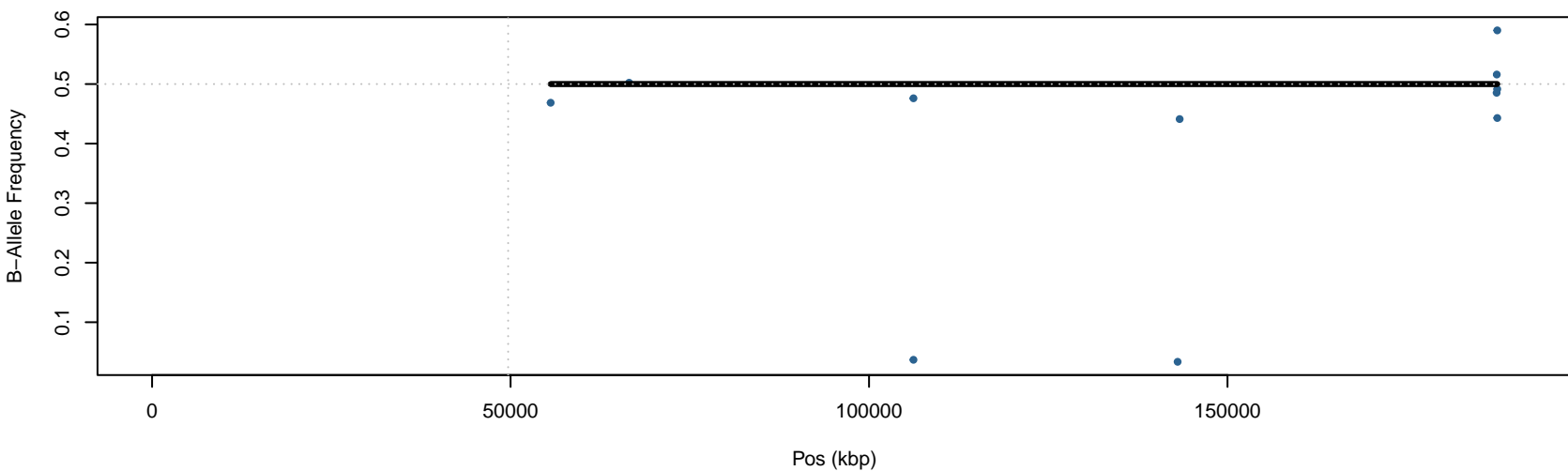
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 2



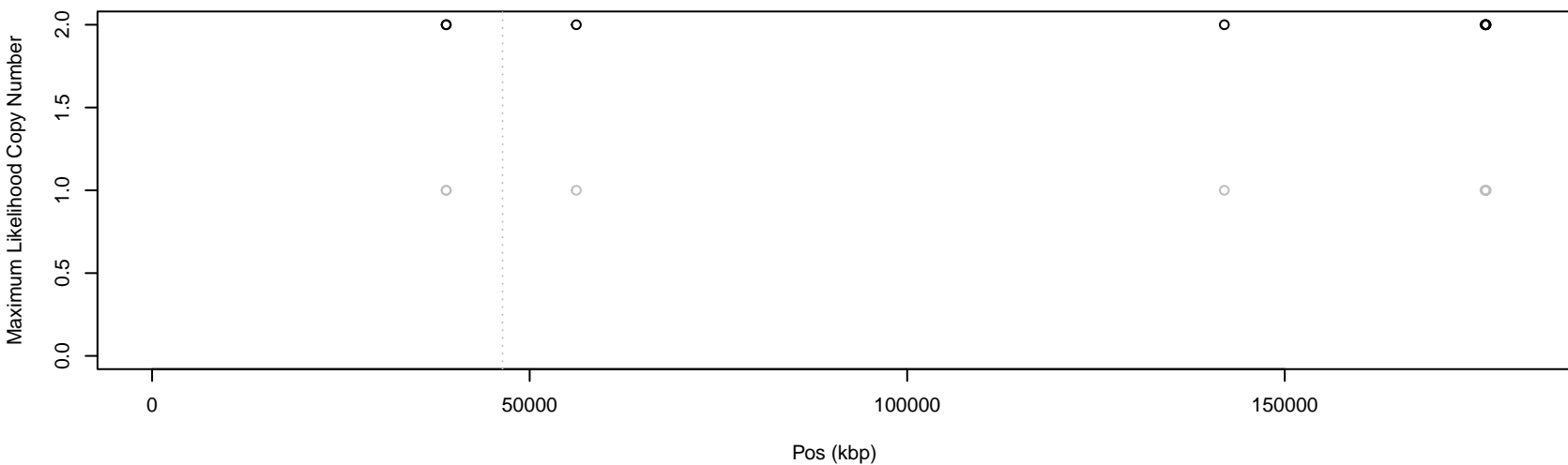
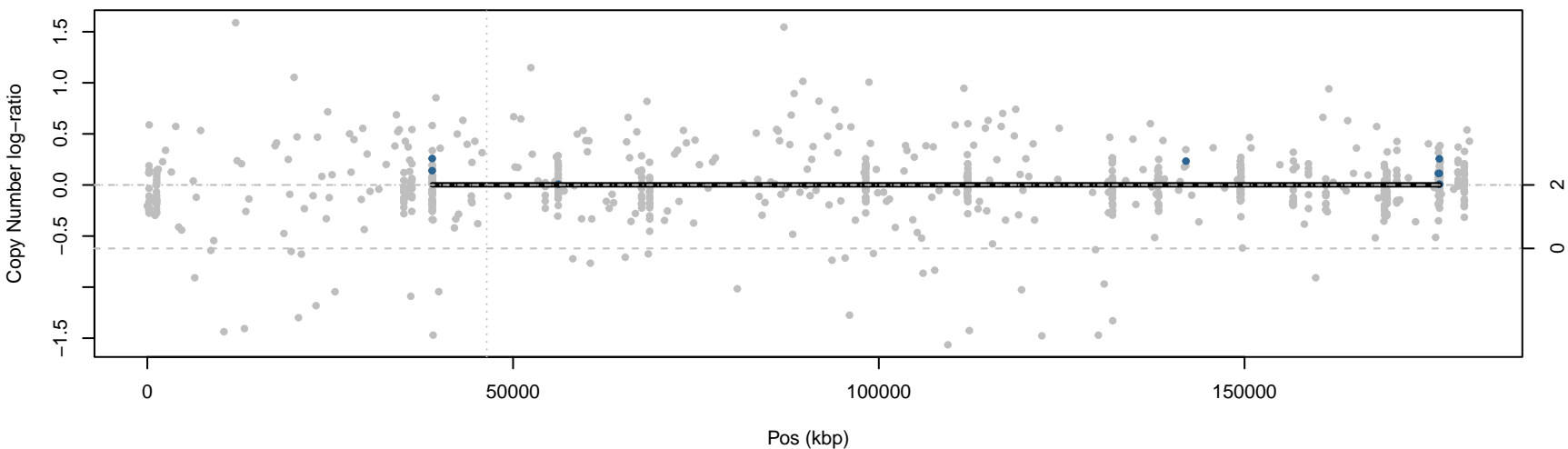
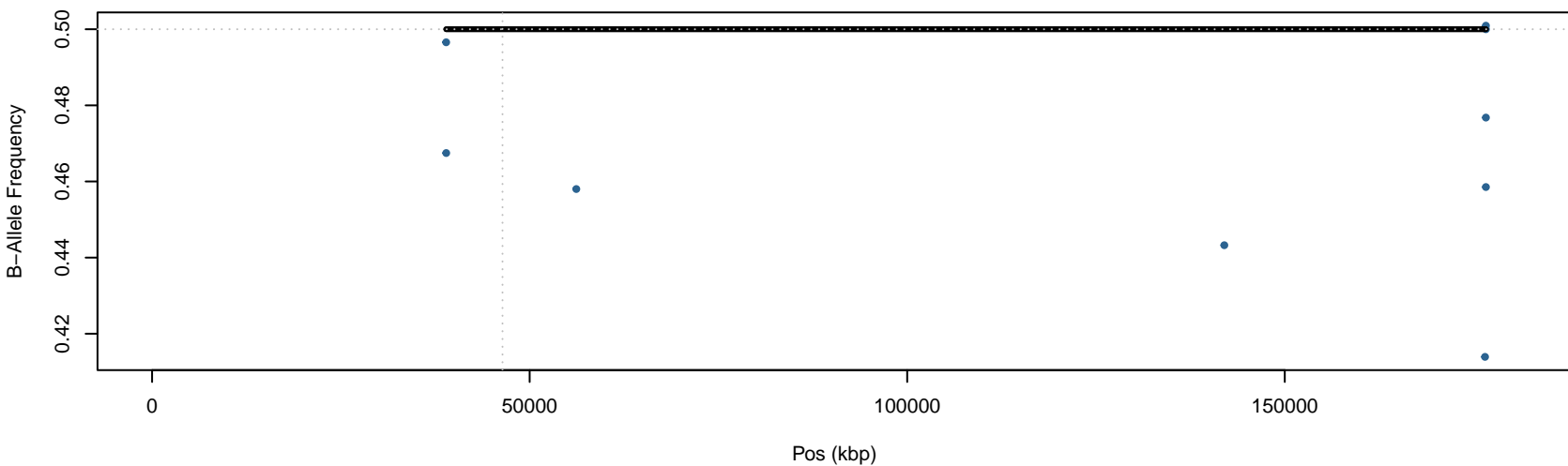
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 3



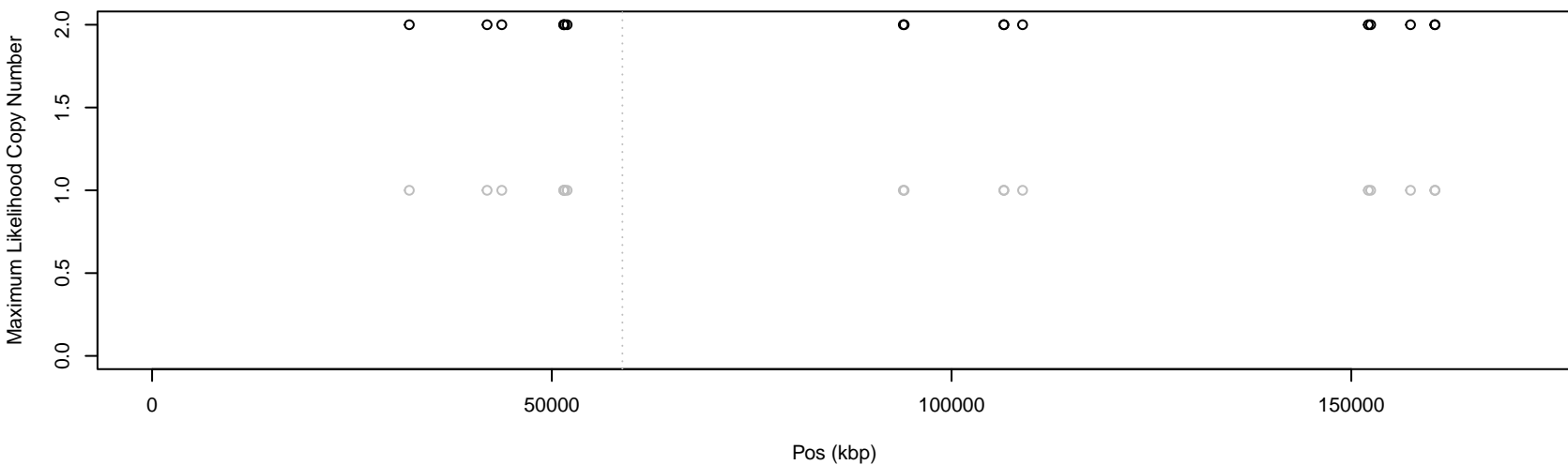
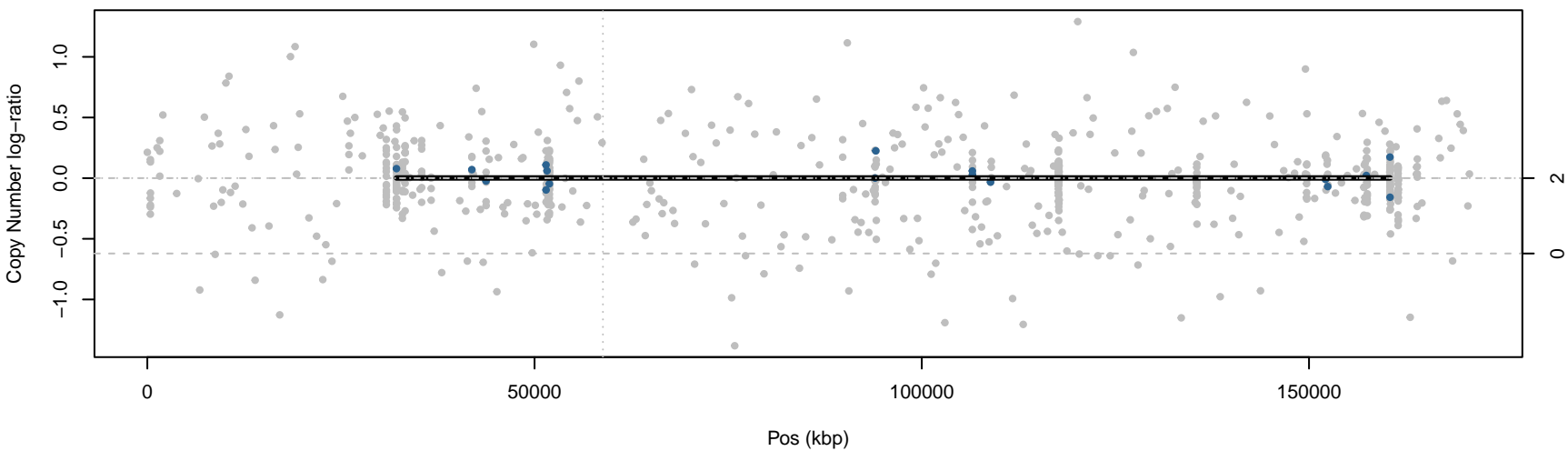
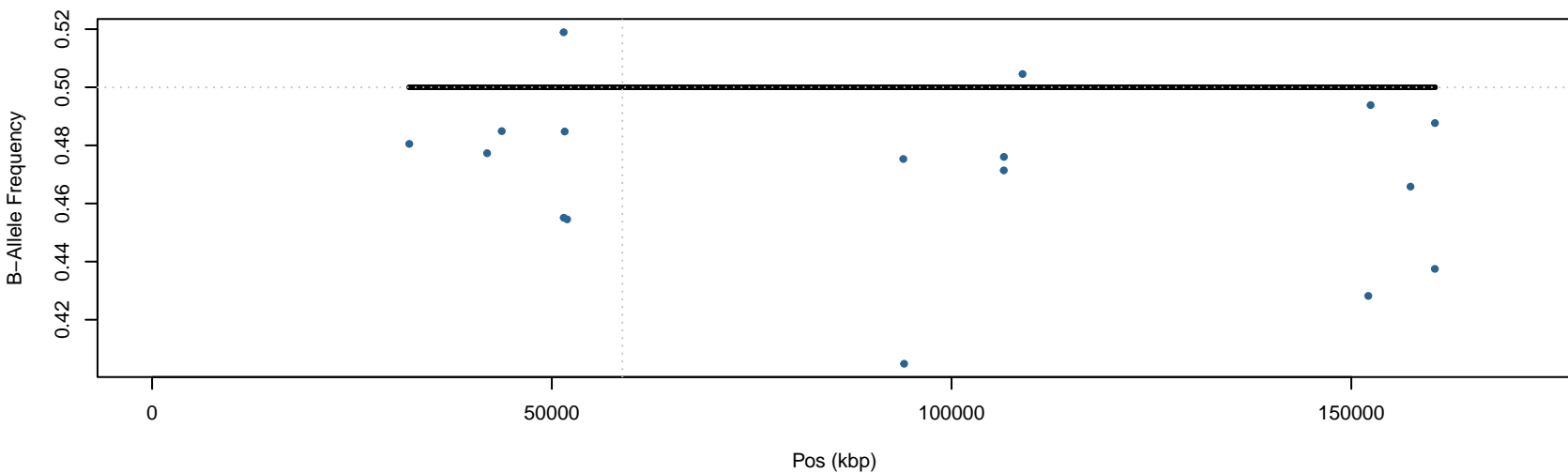
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 4



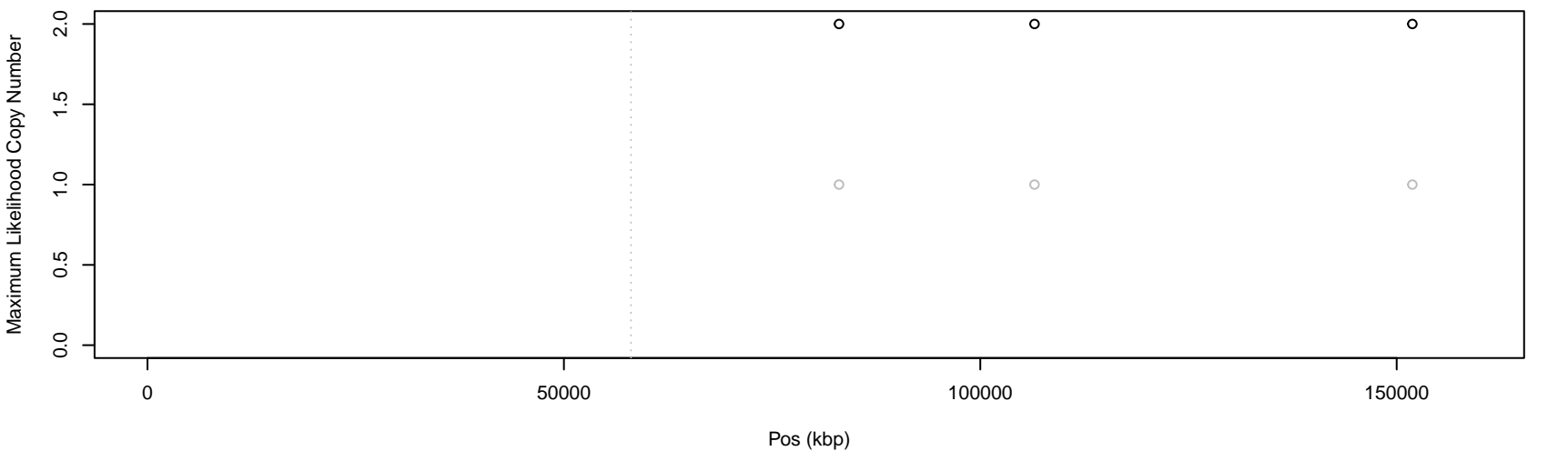
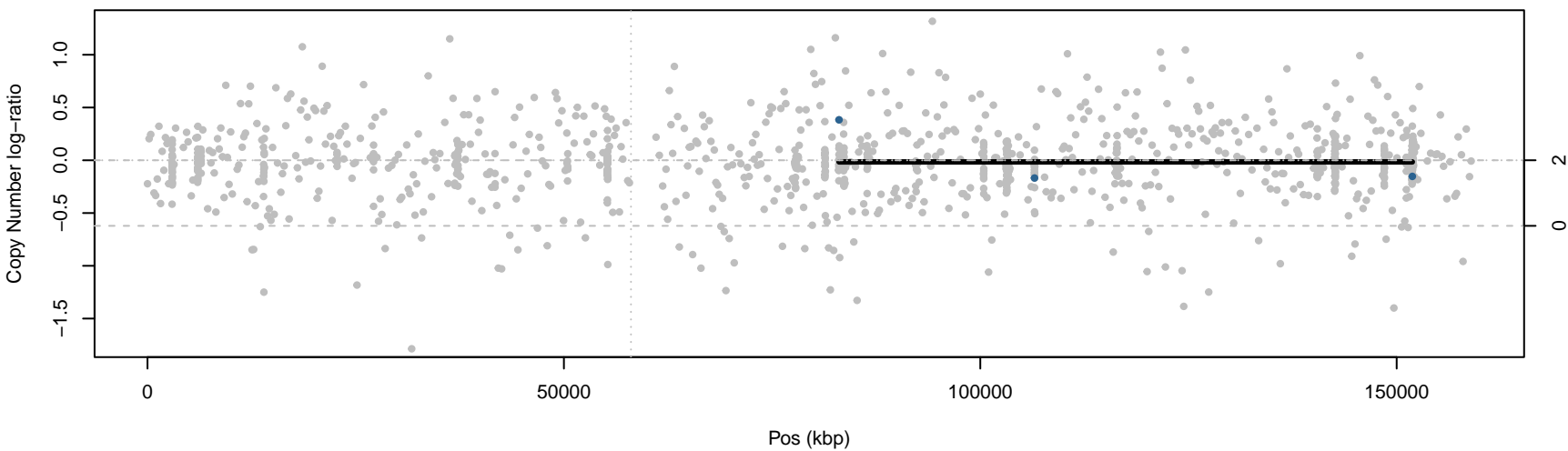
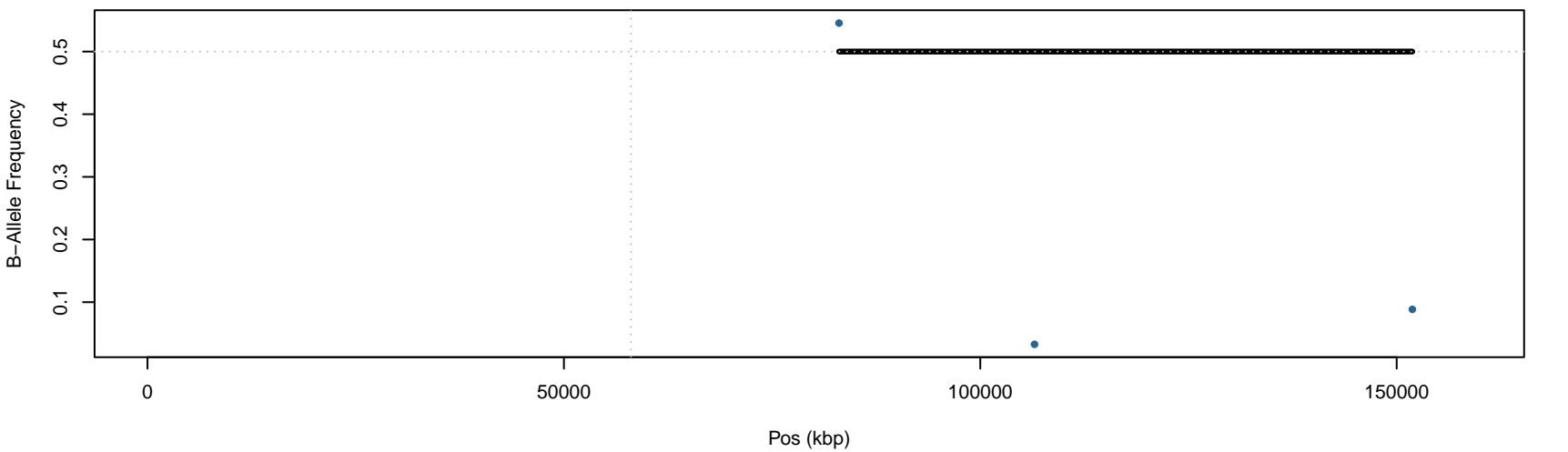
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 5



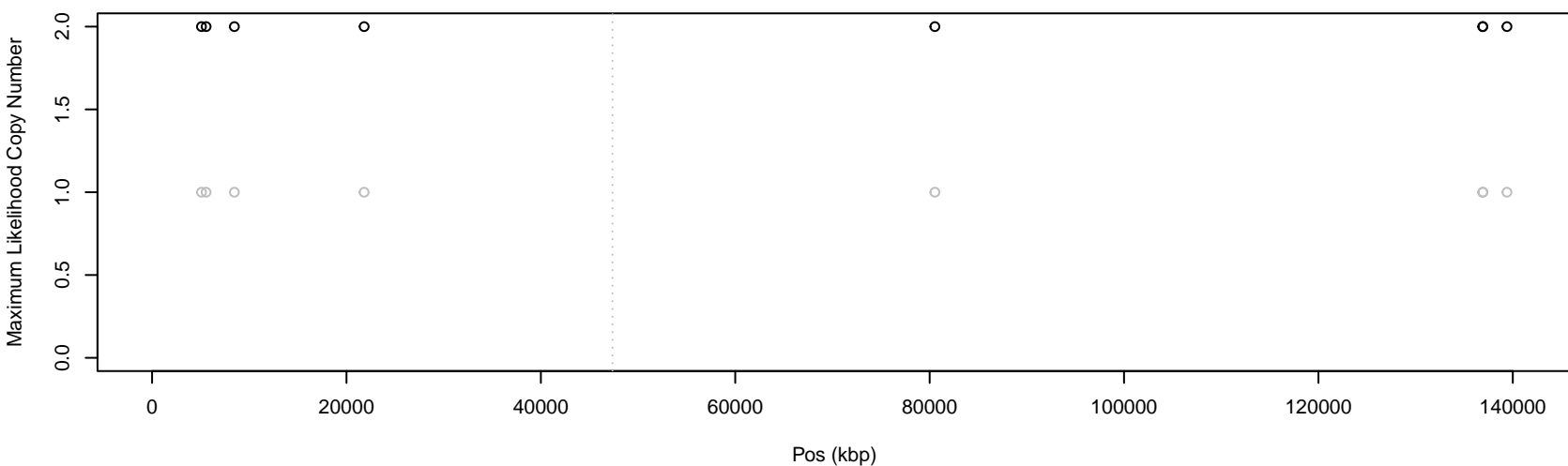
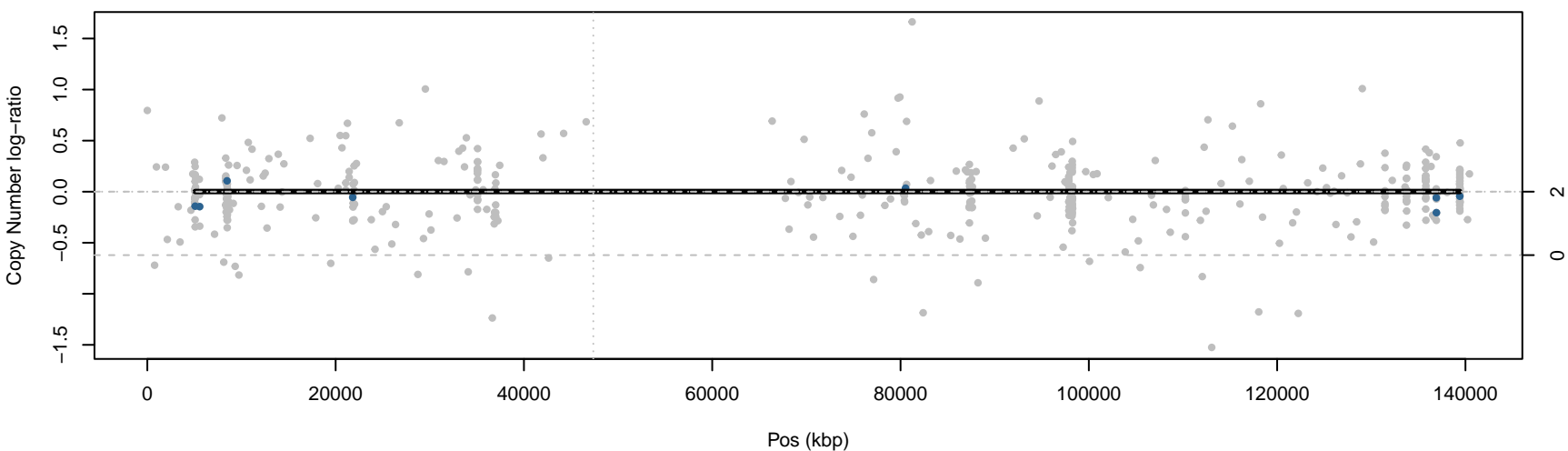
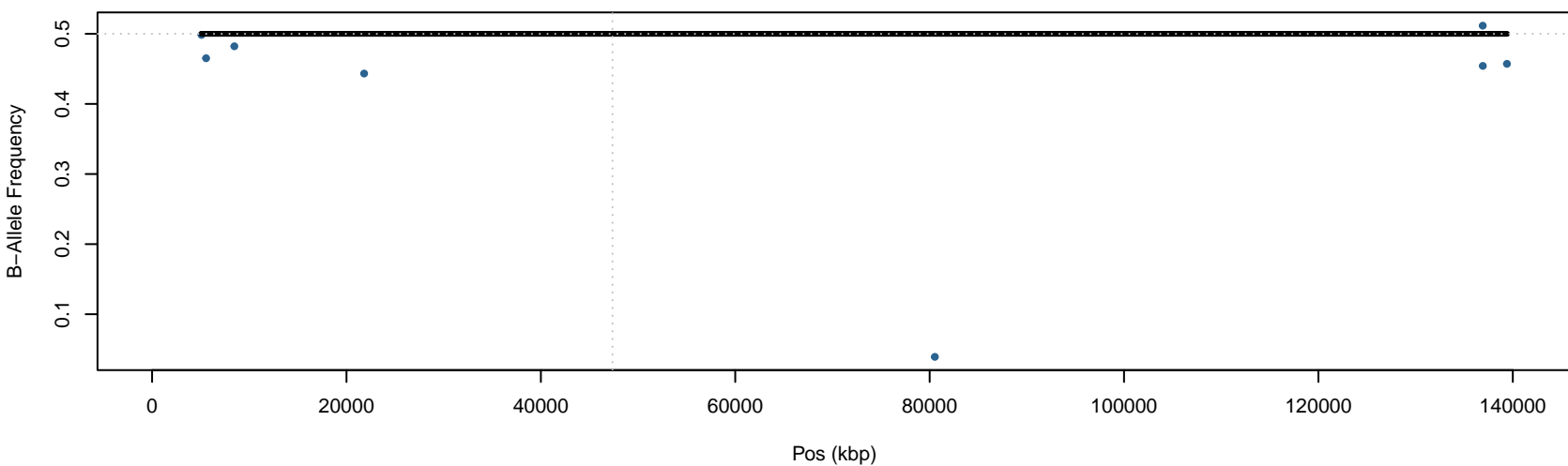
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 6



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 7

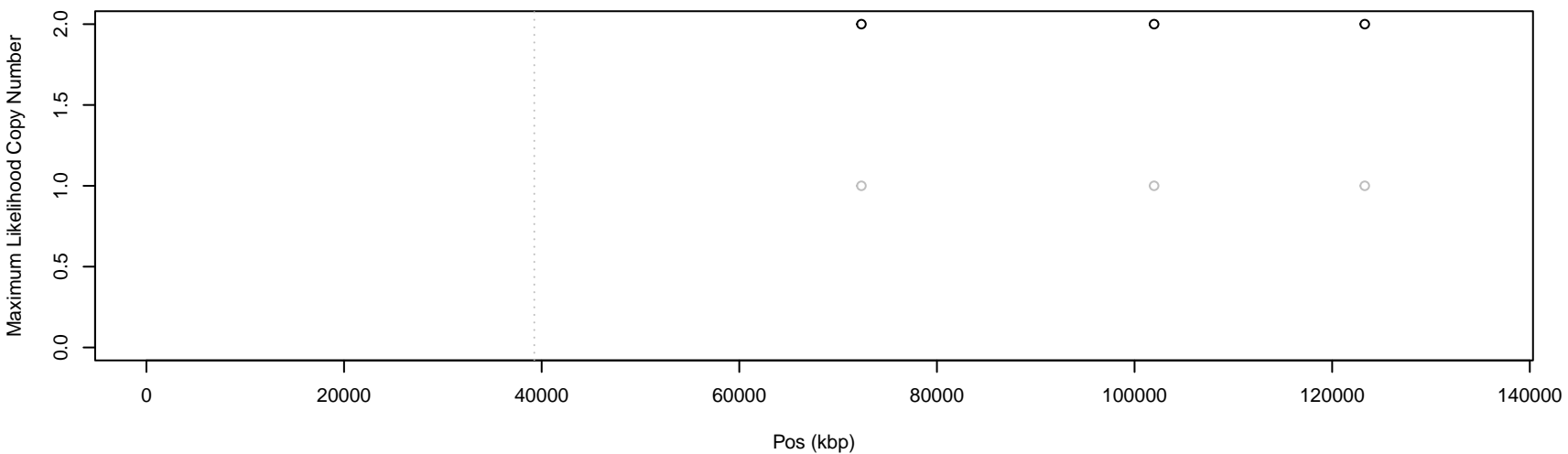
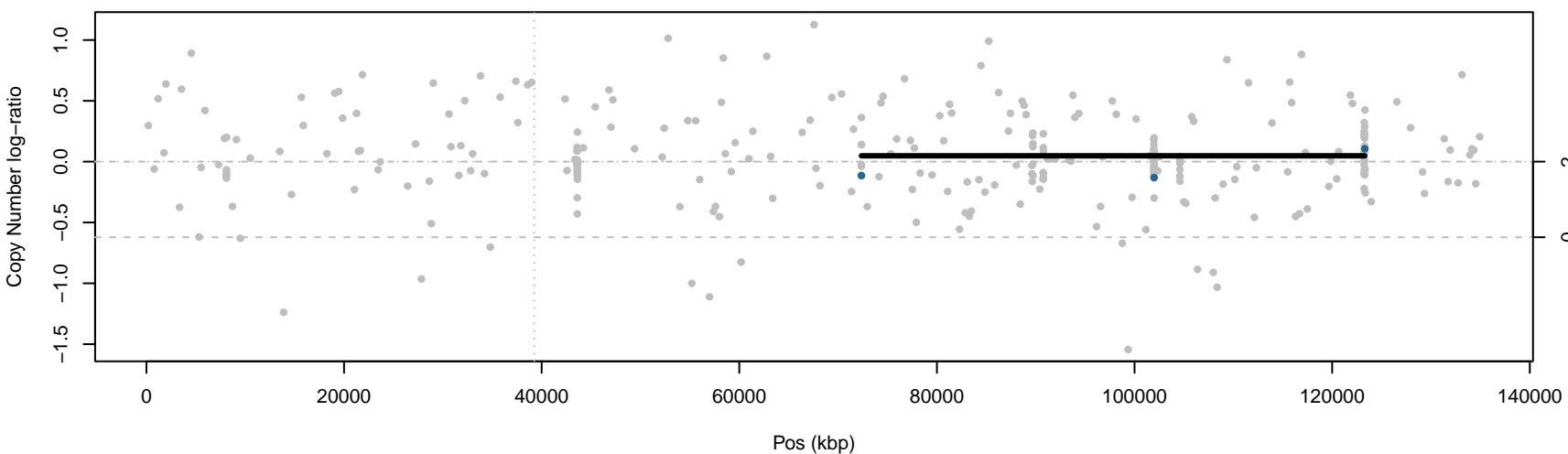
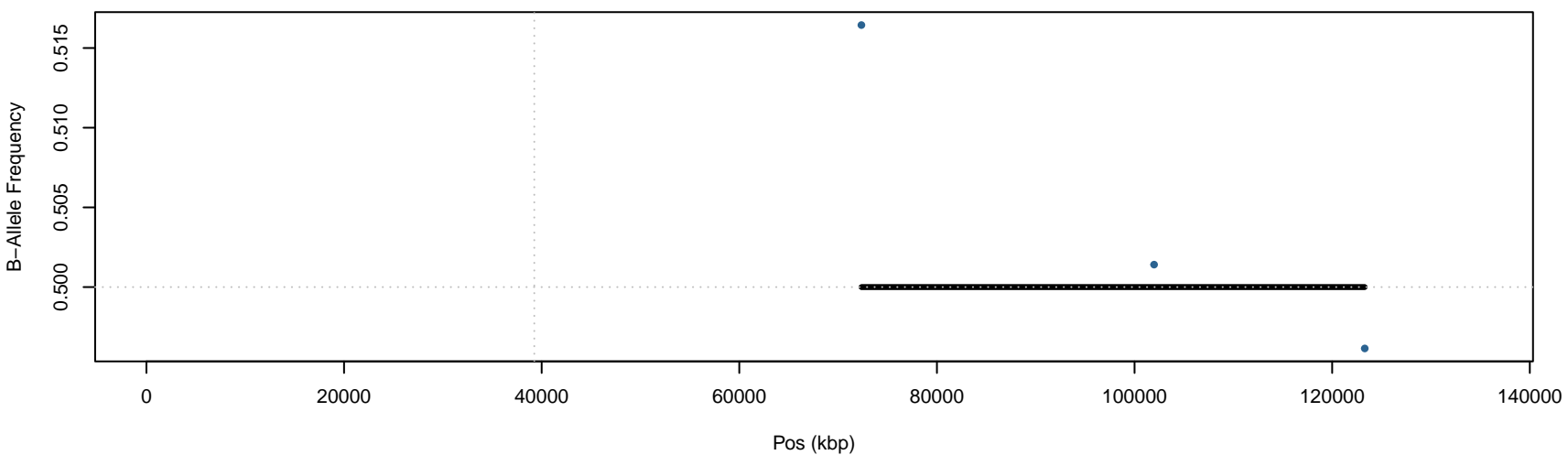


Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 9

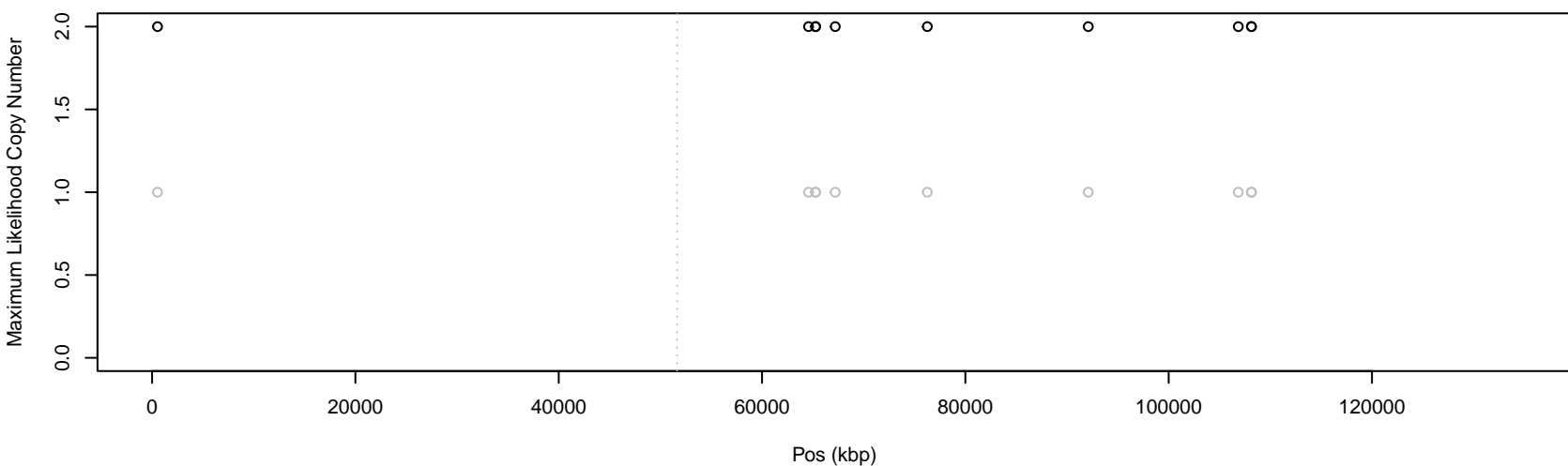
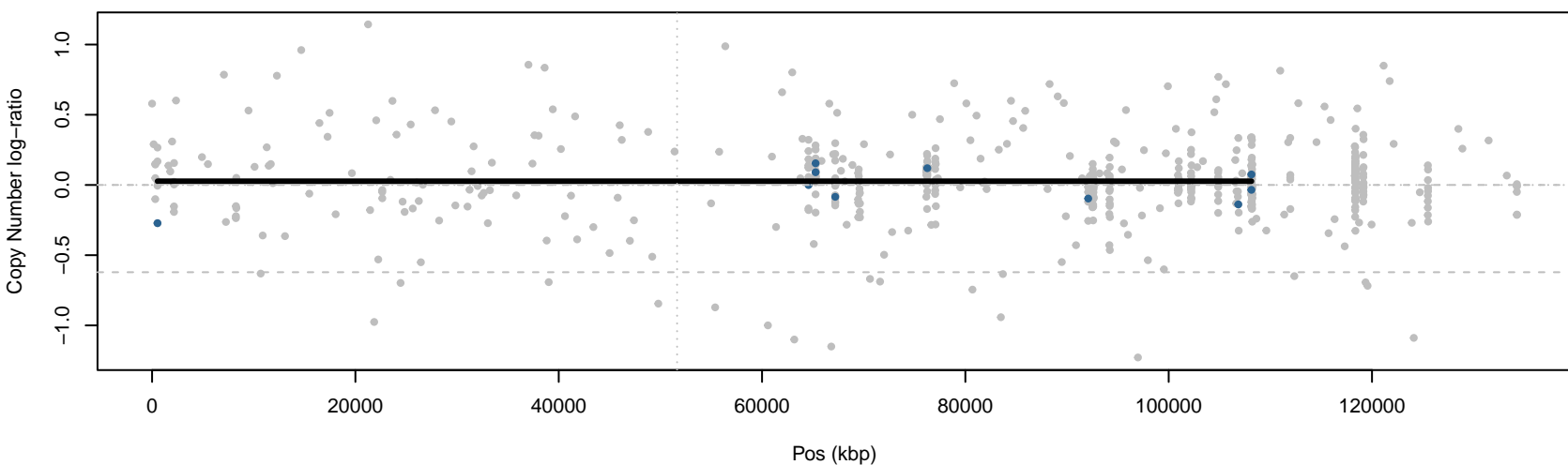
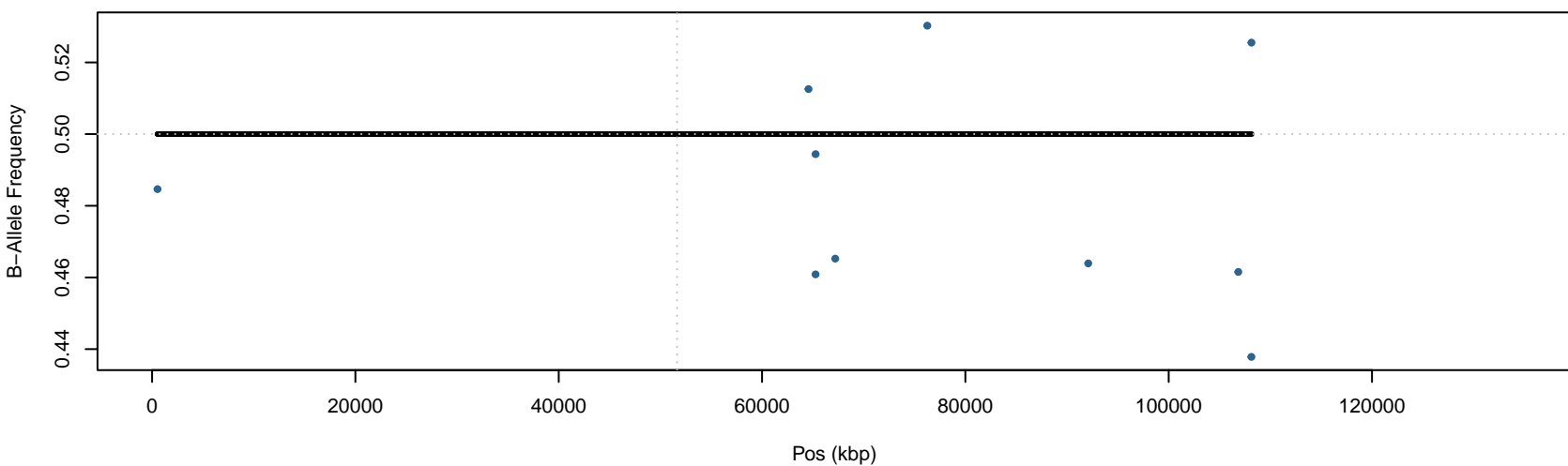




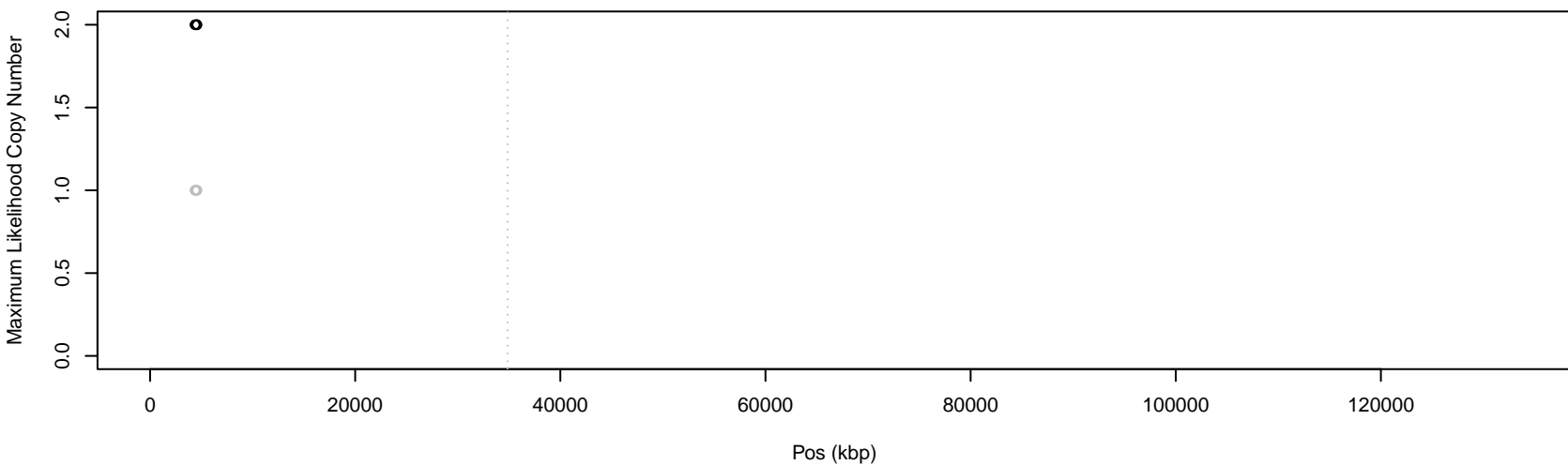
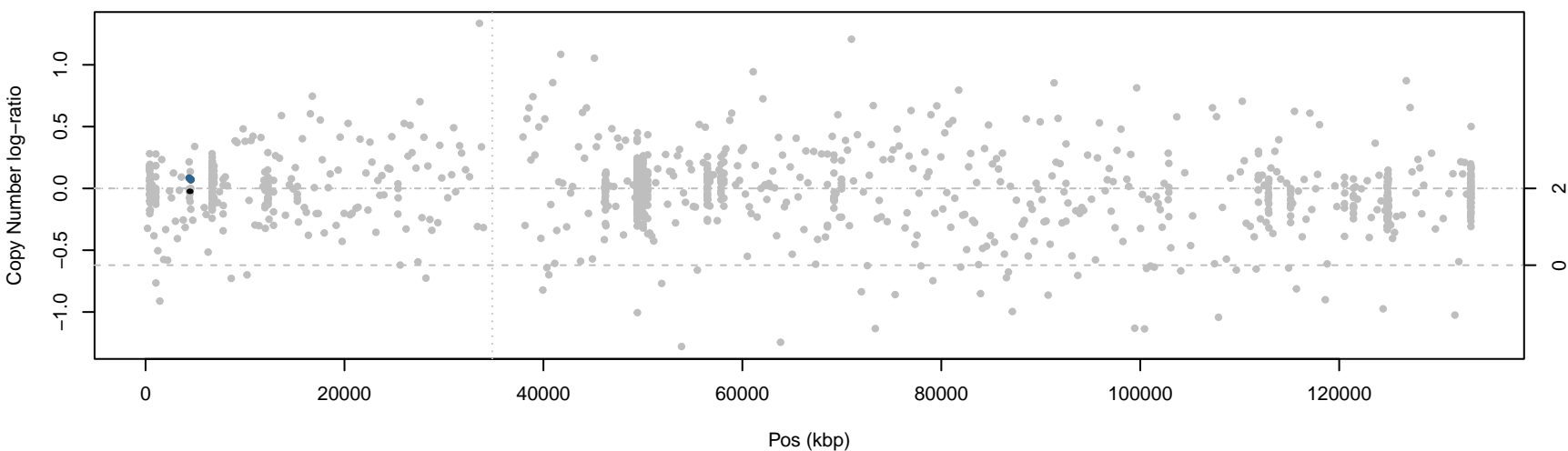
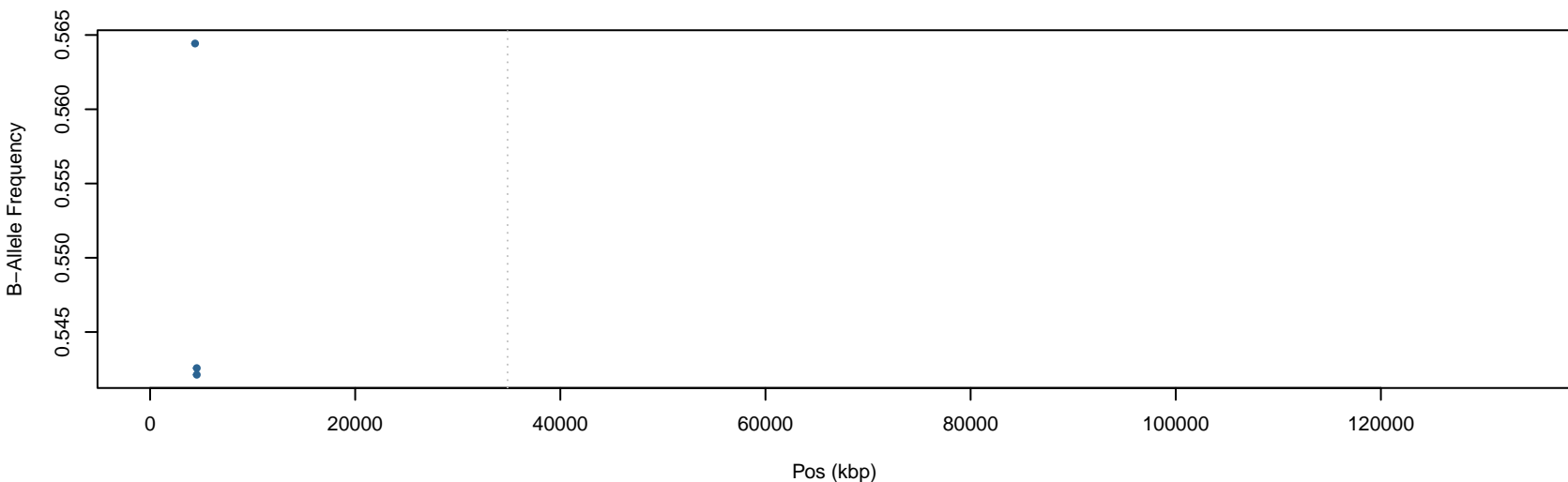
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 10



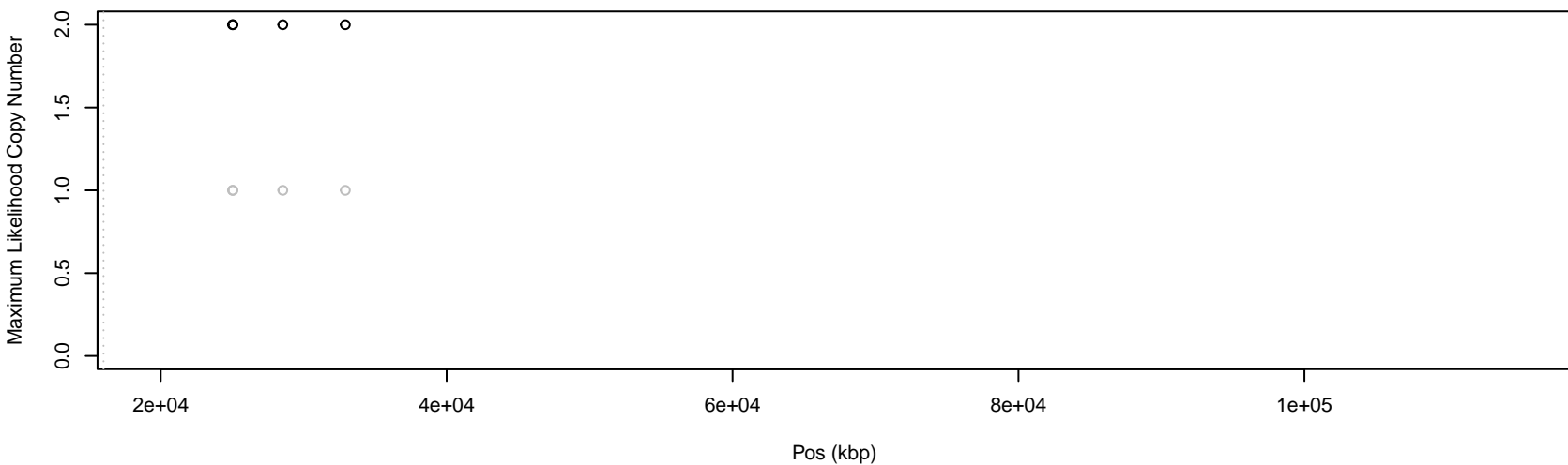
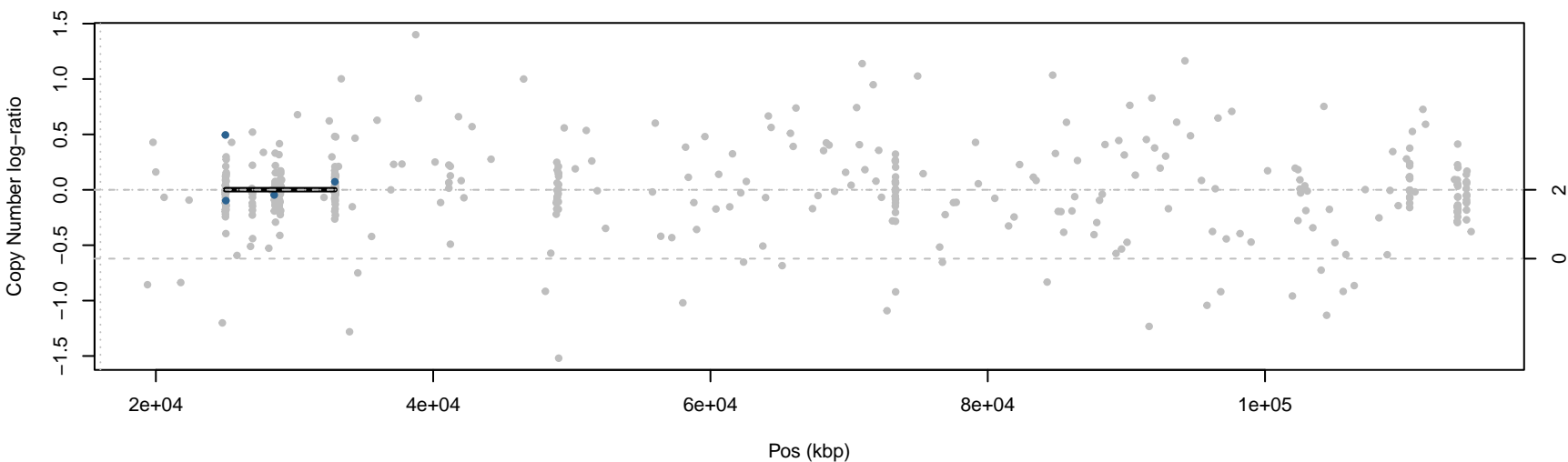
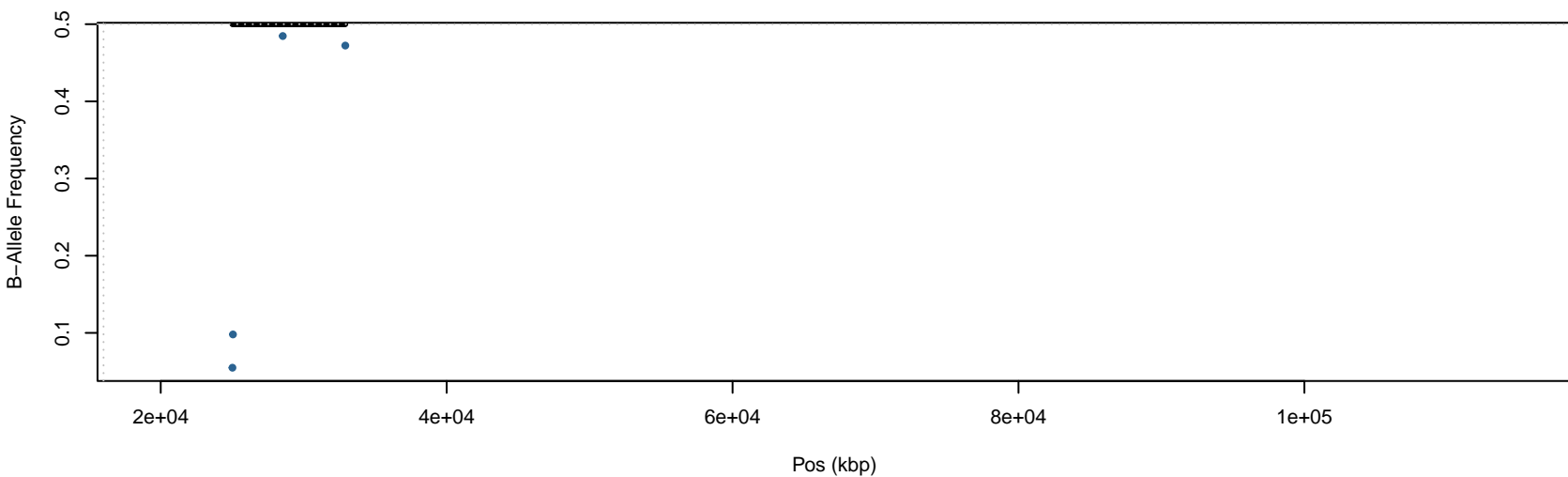
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 11



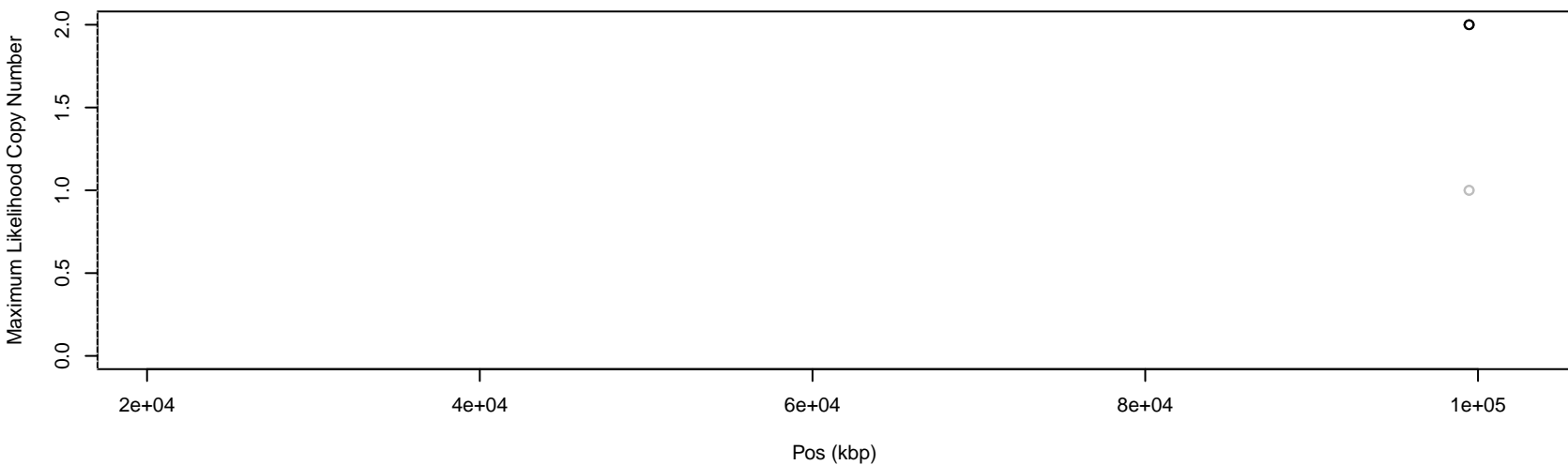
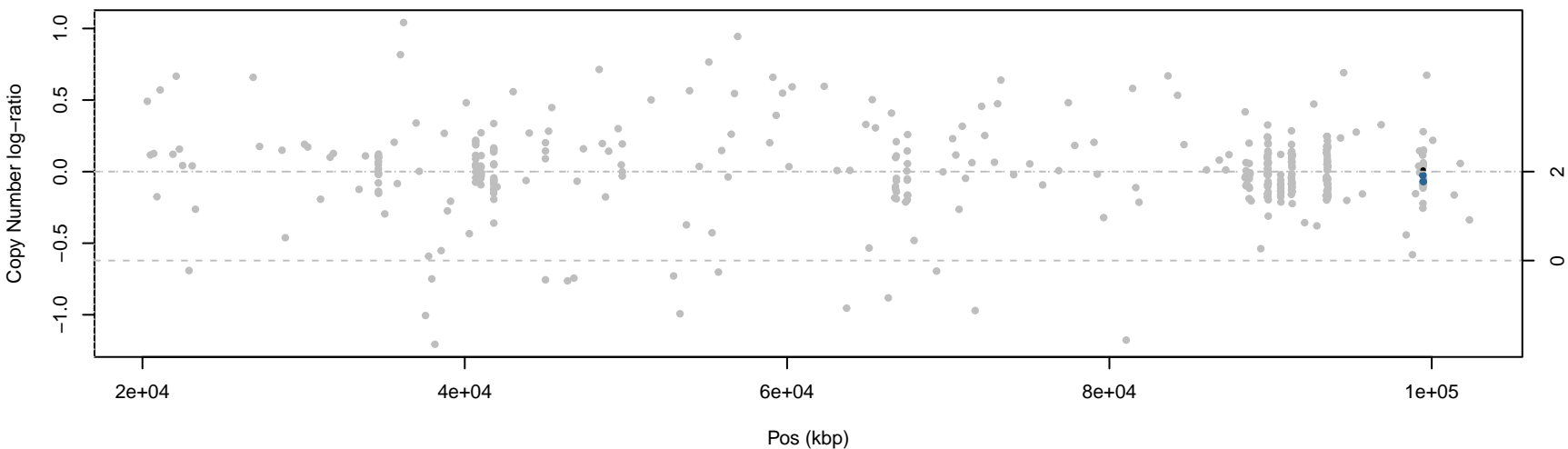
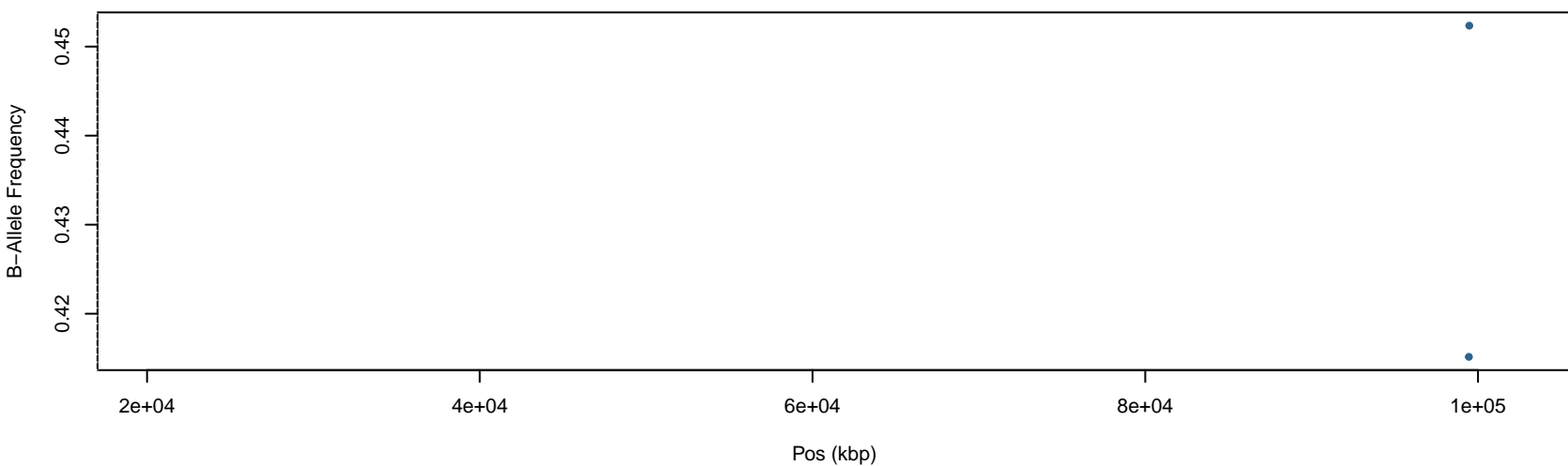
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 12



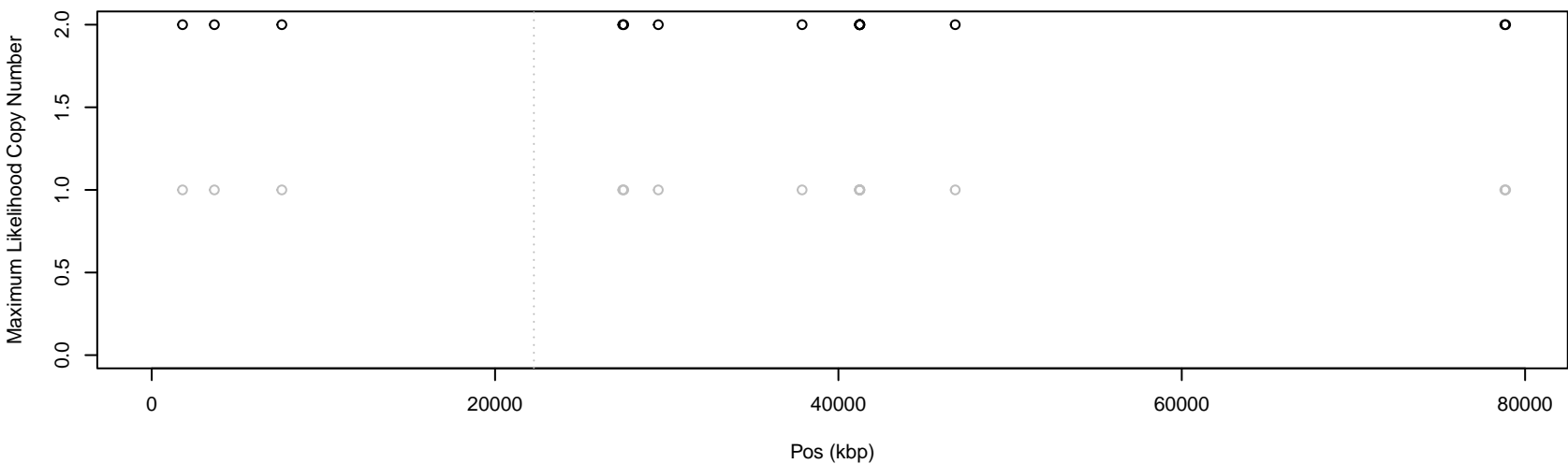
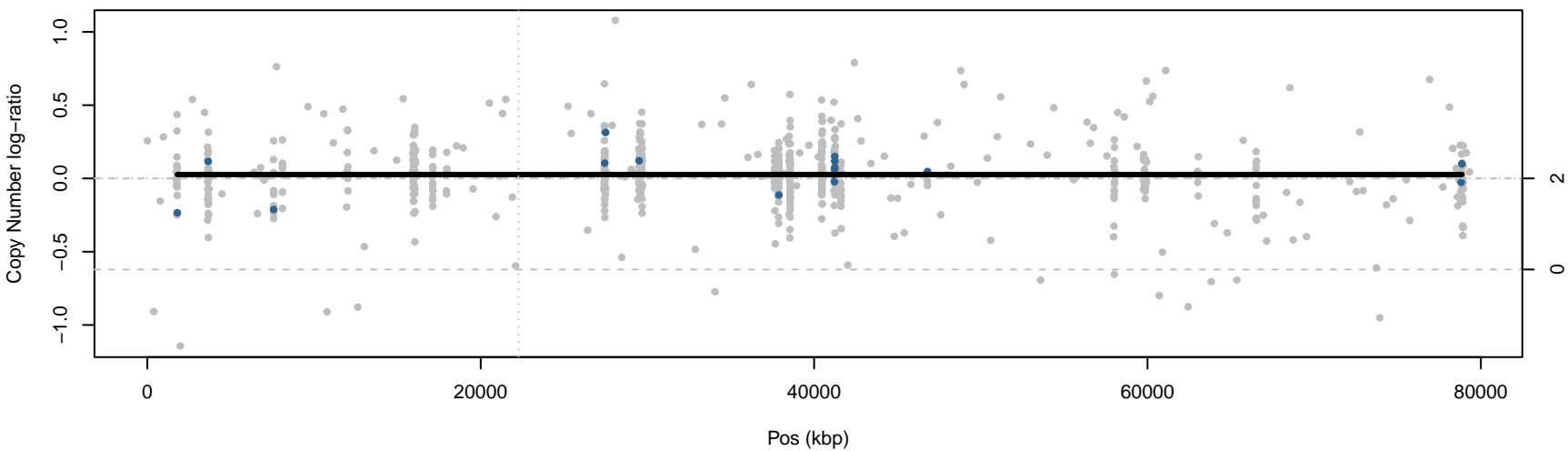
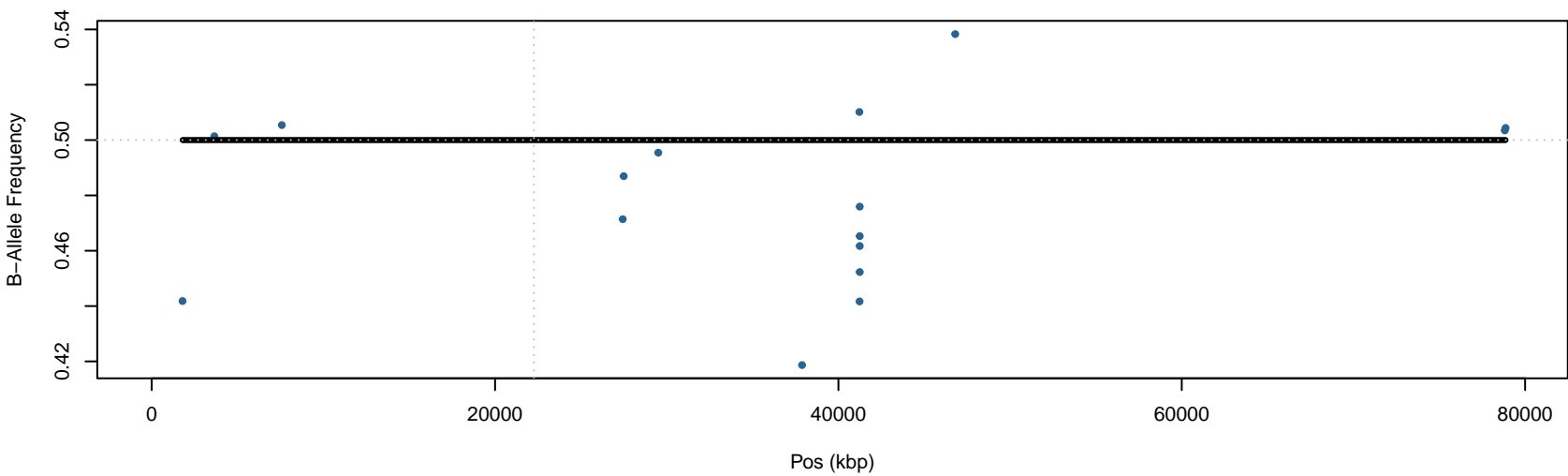
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 13



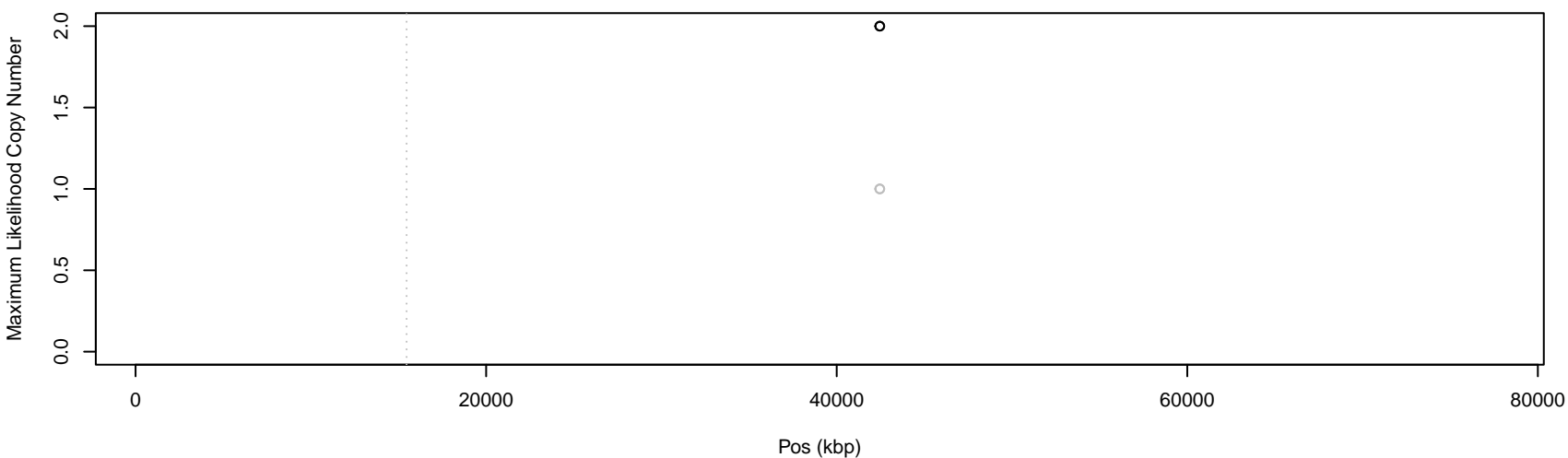
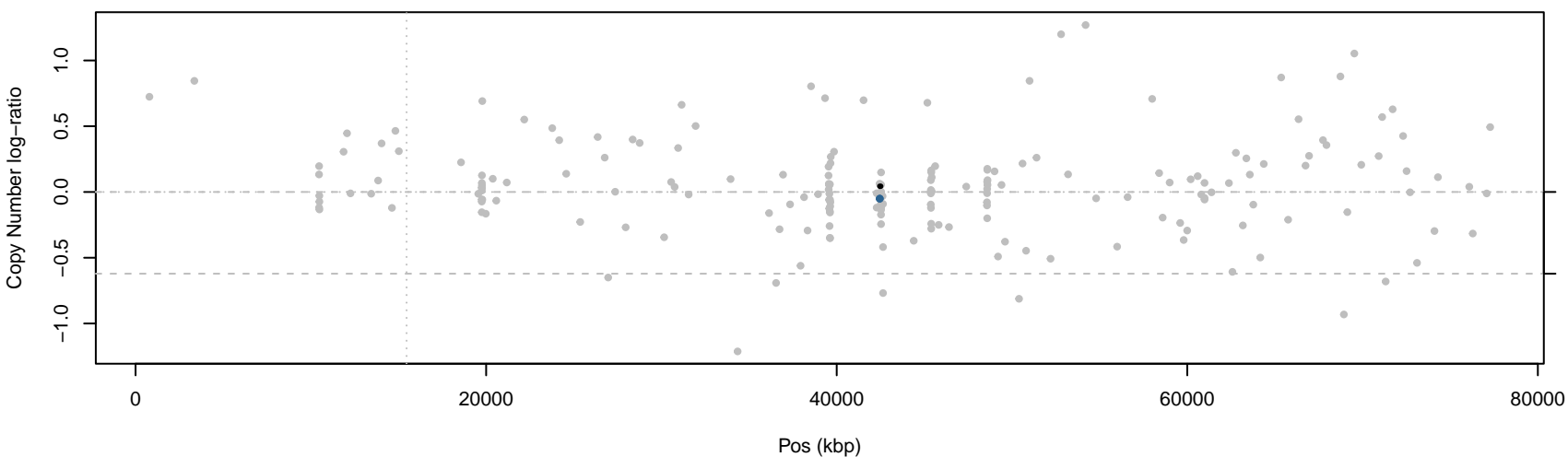
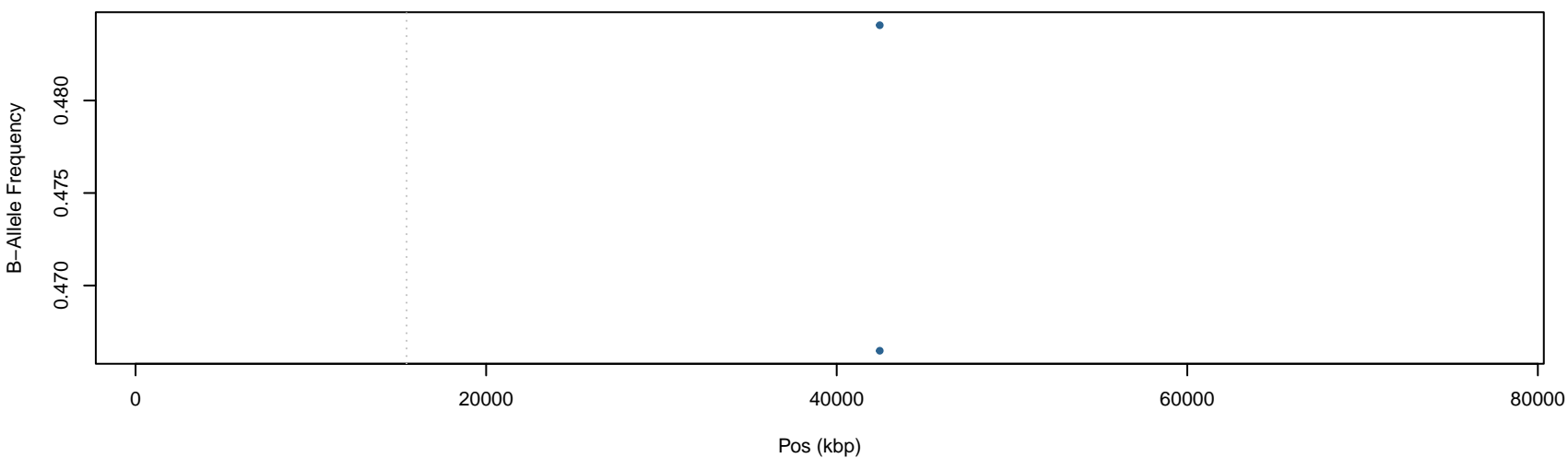
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 15



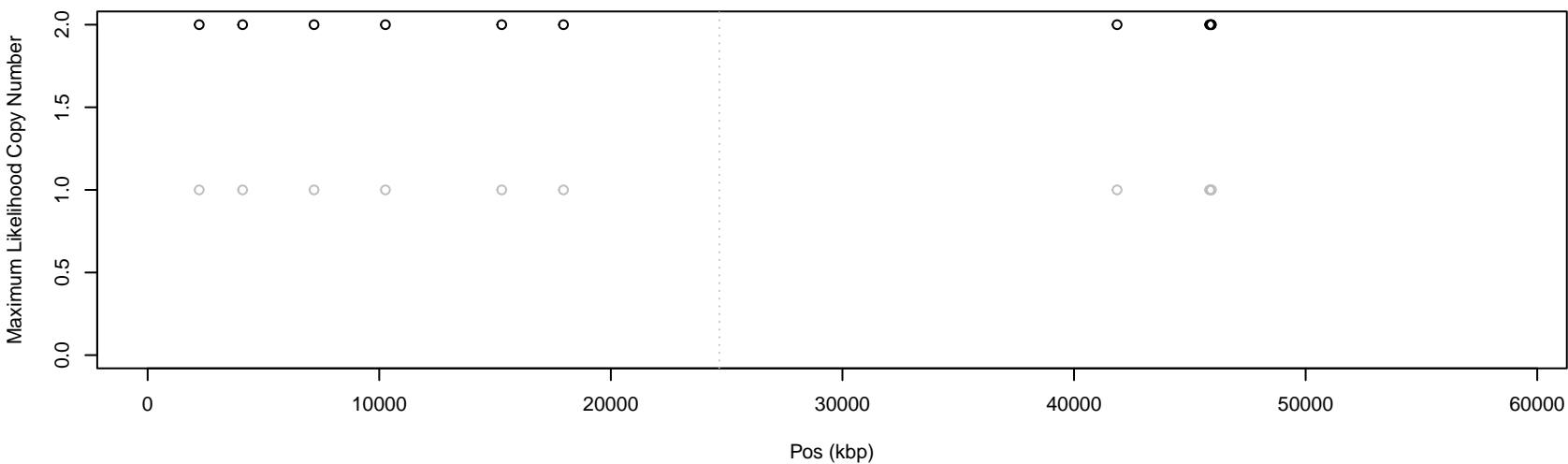
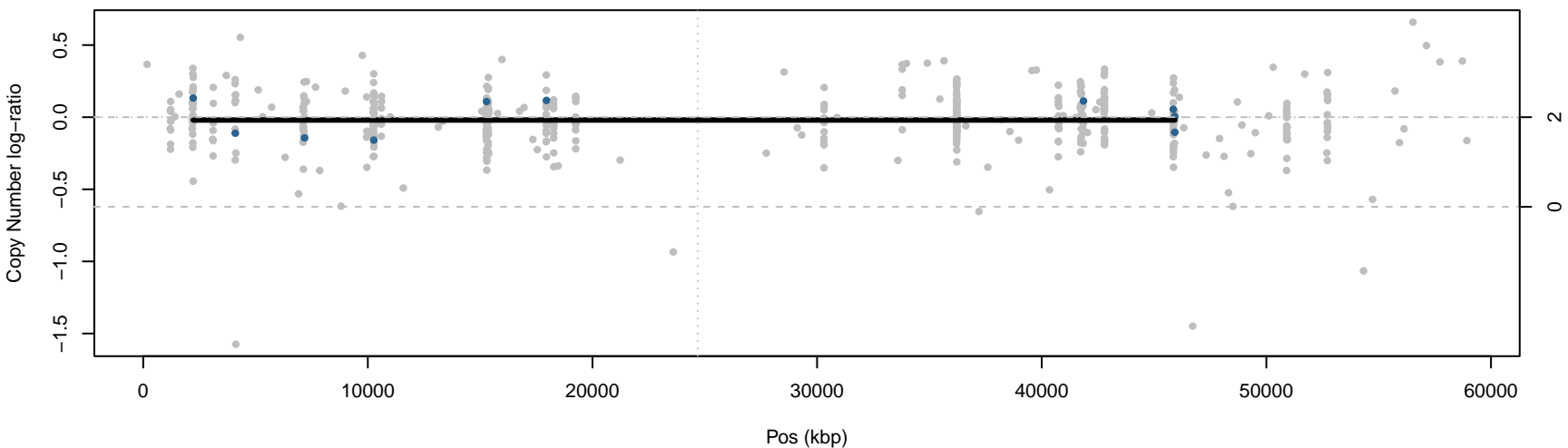
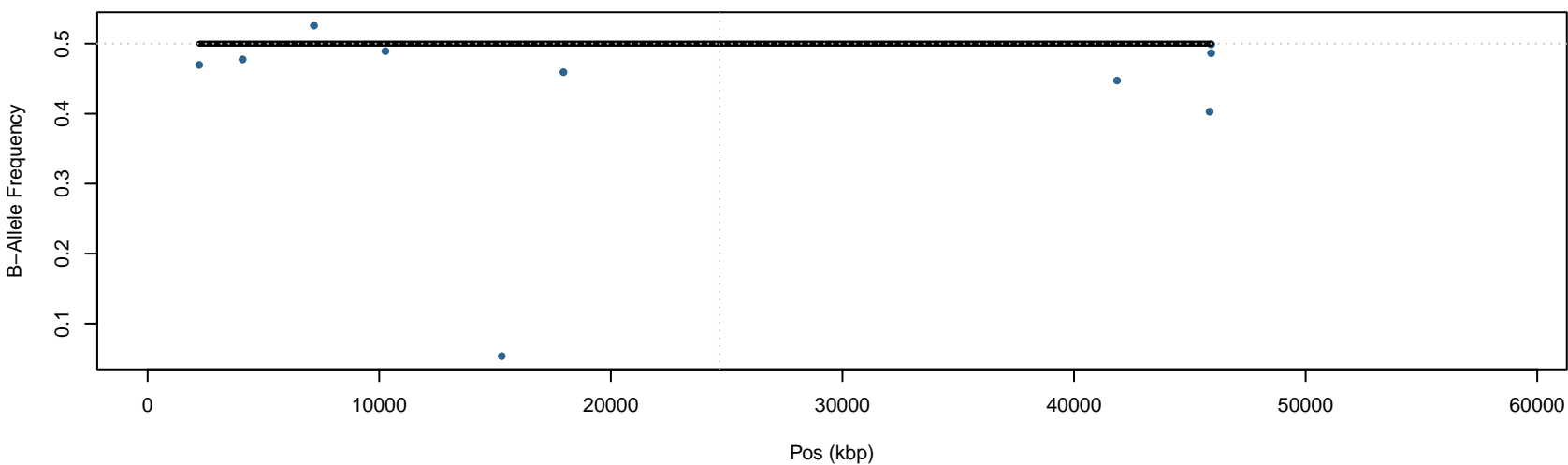
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 17



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 18

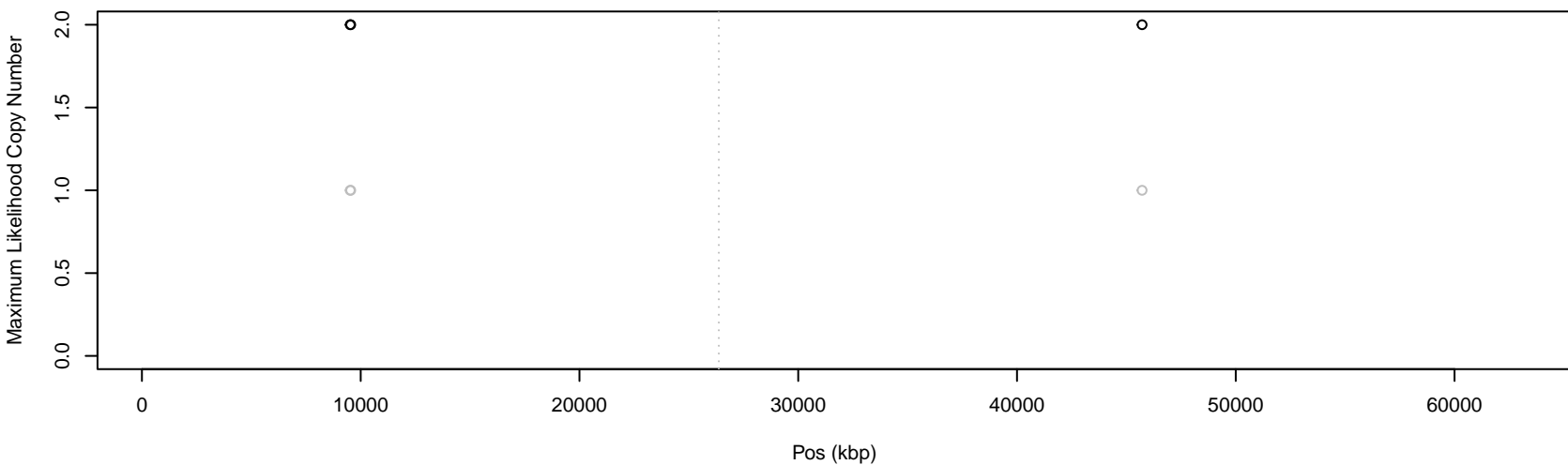
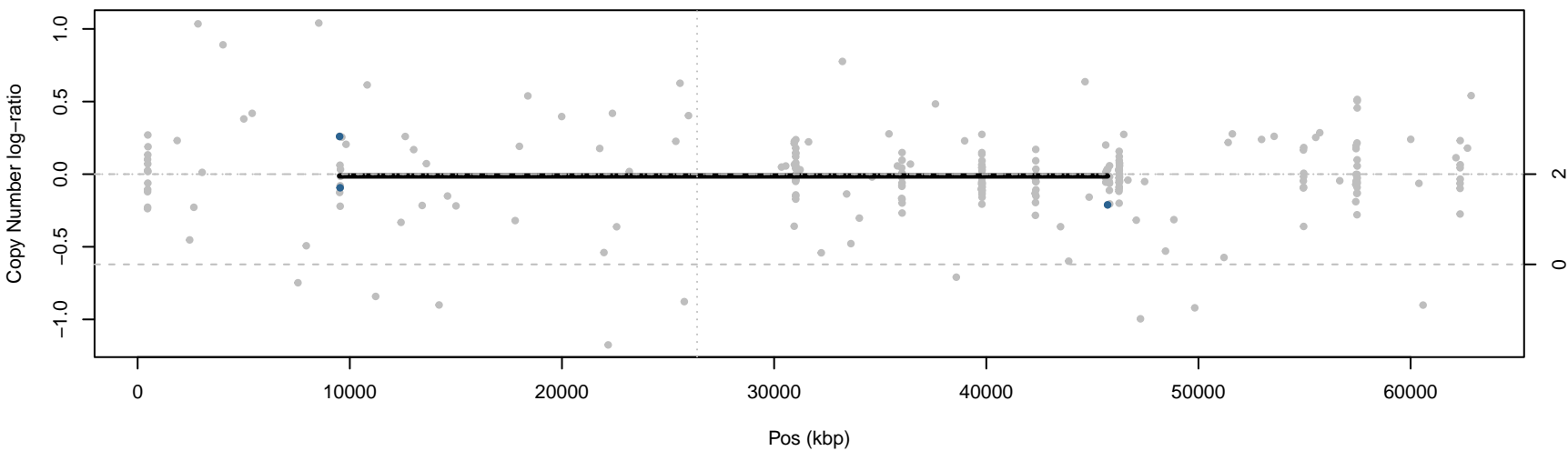
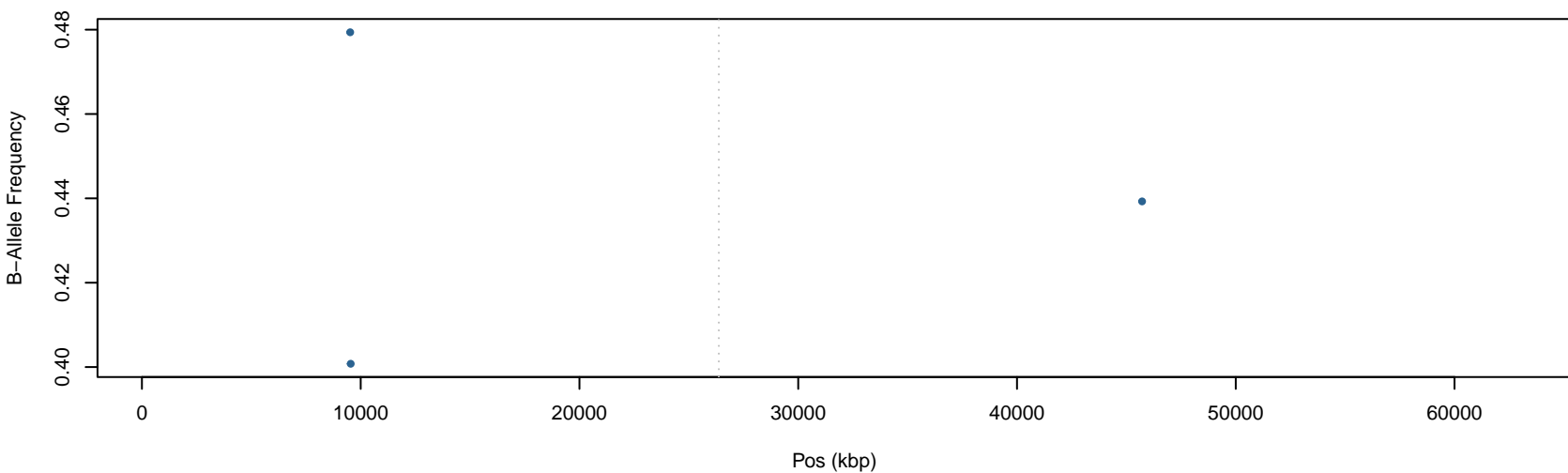


Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 19





Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 20



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 22

