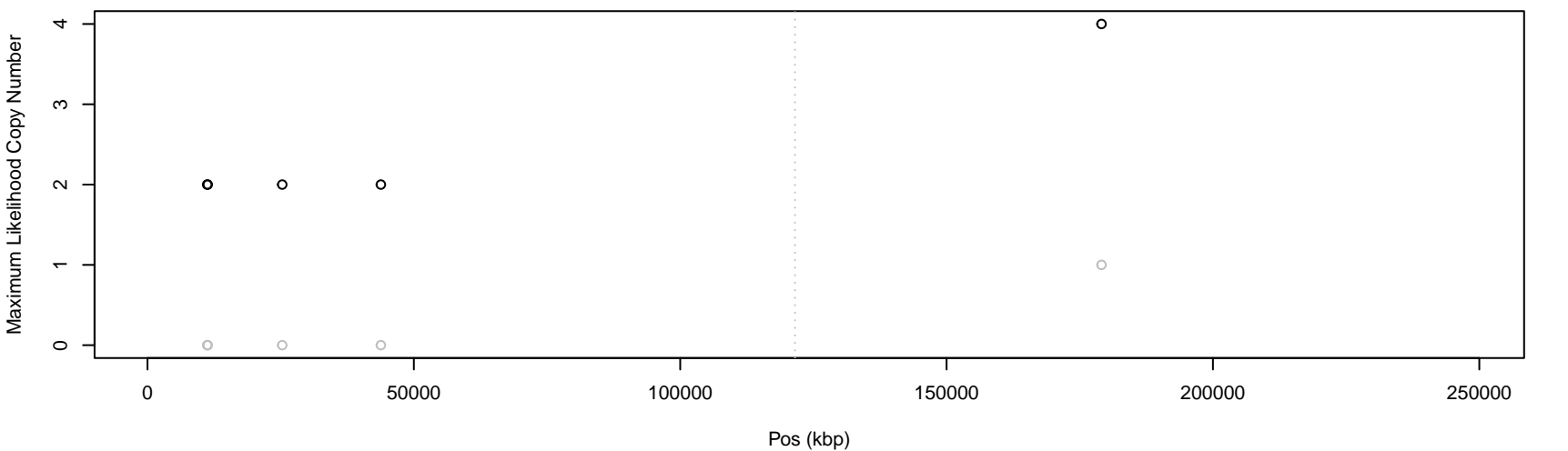
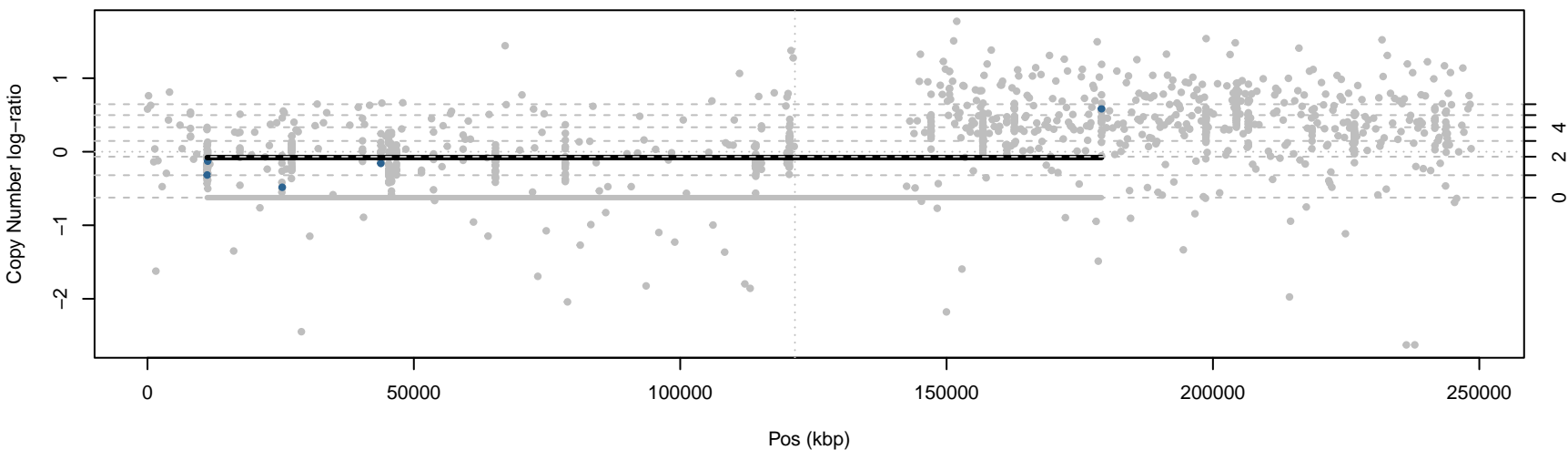
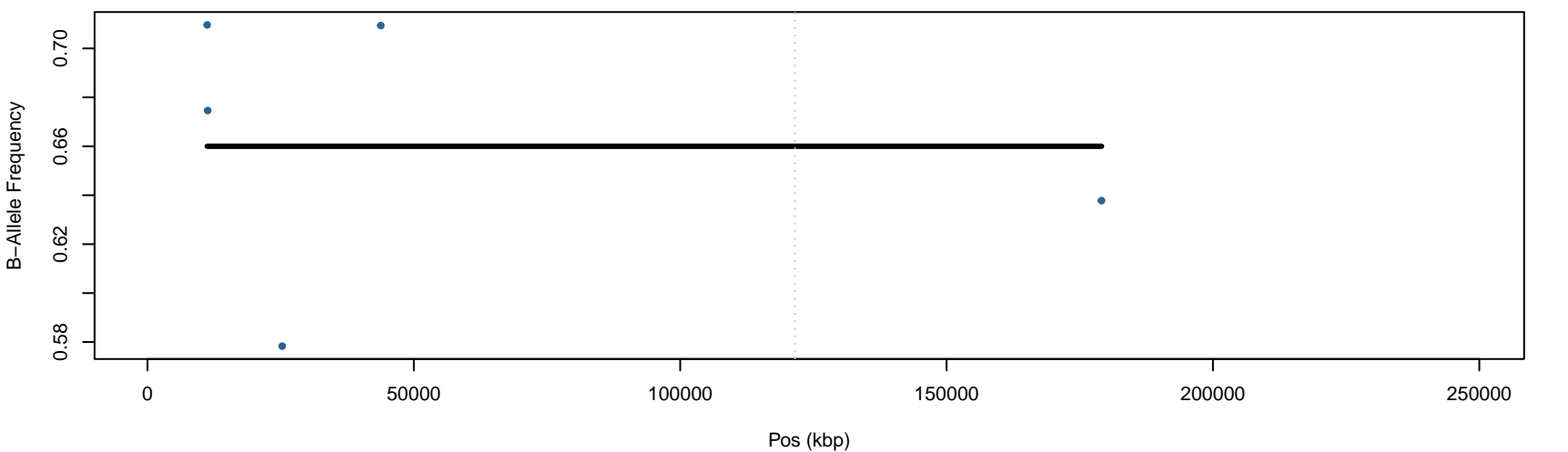
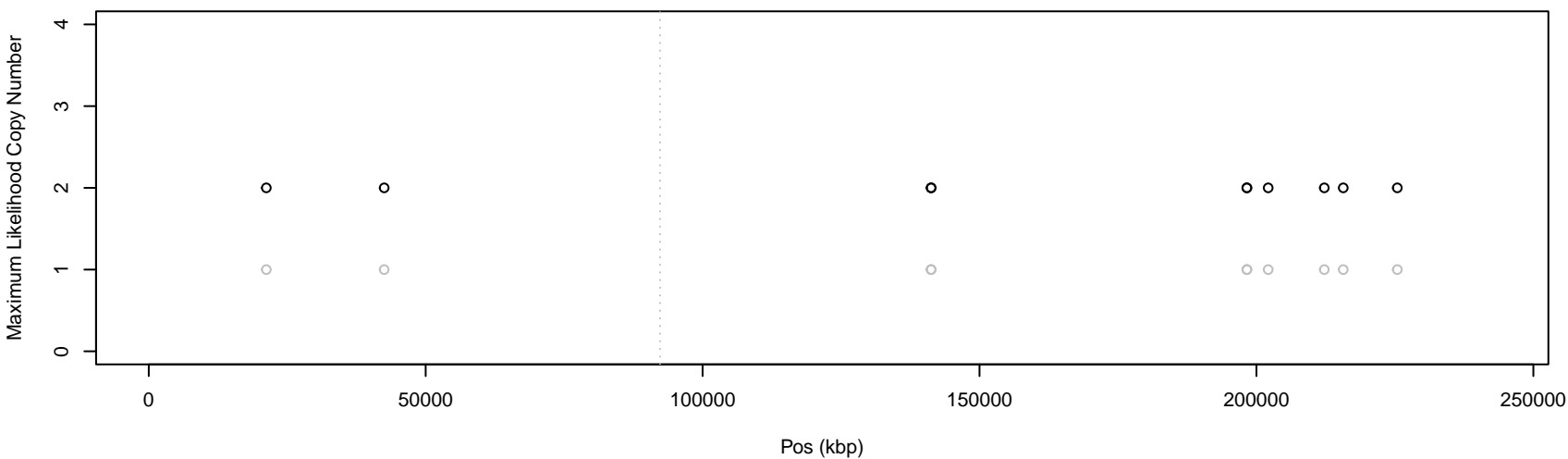
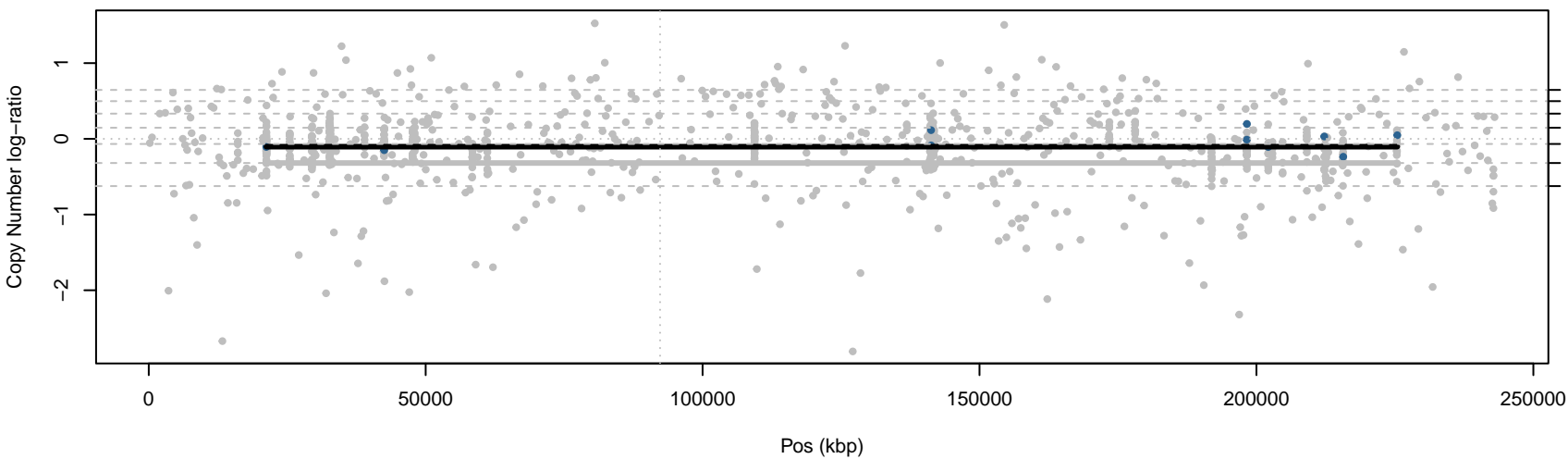
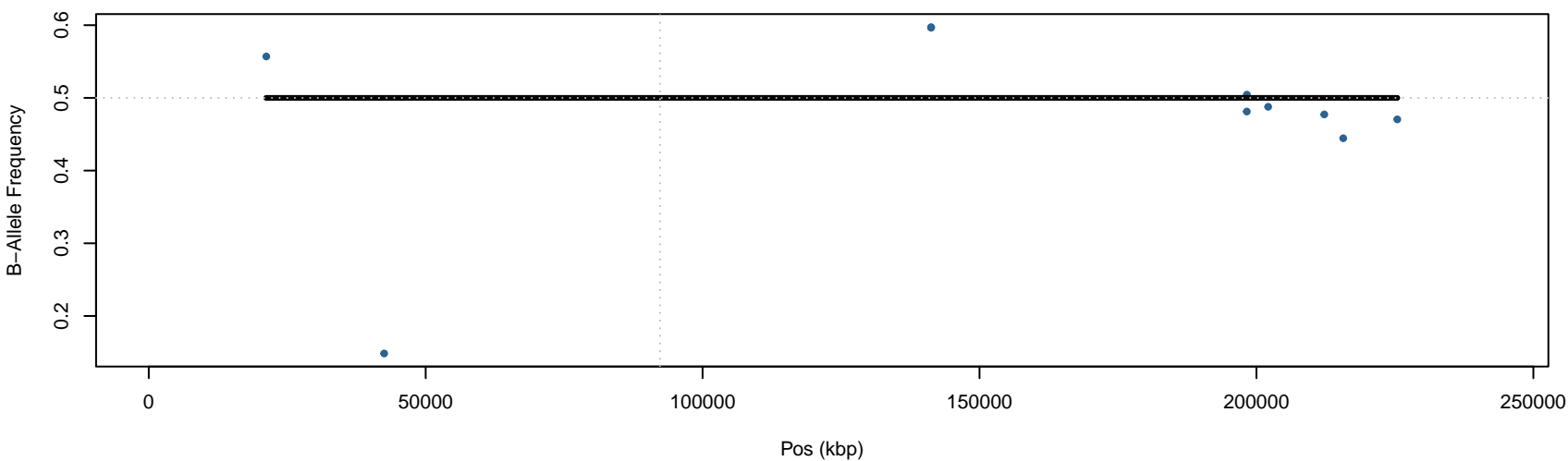


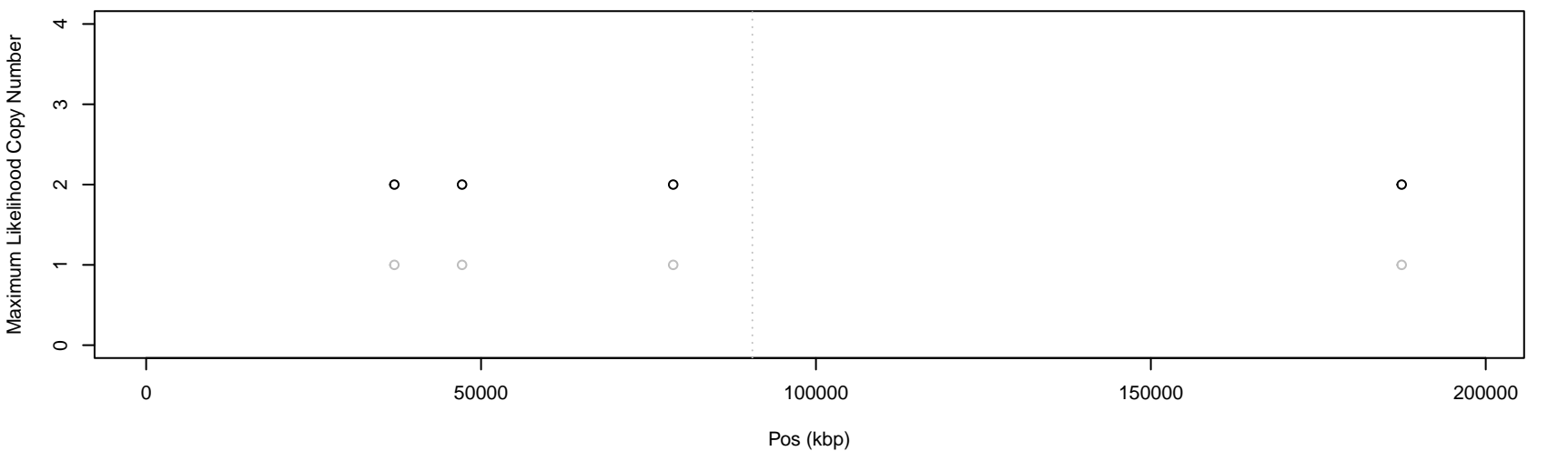
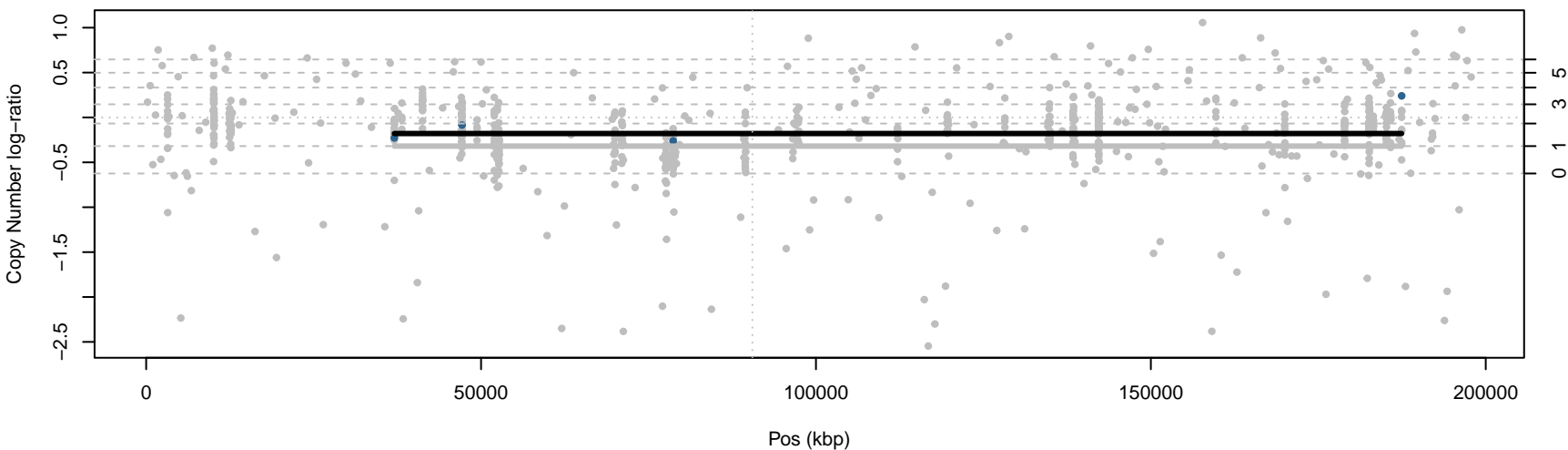
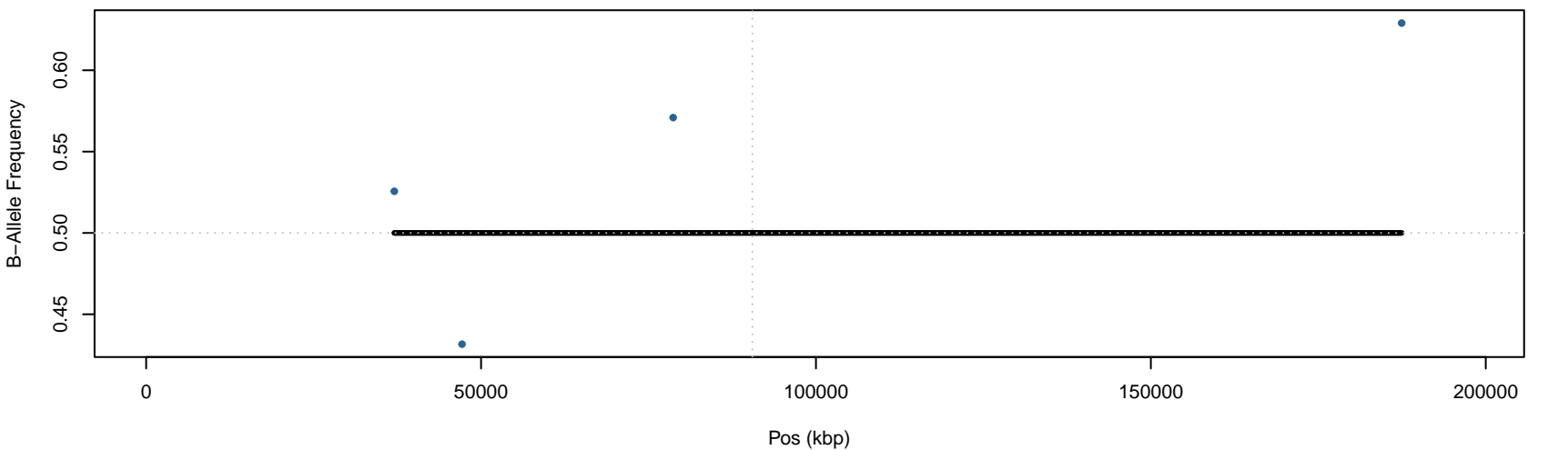
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 1



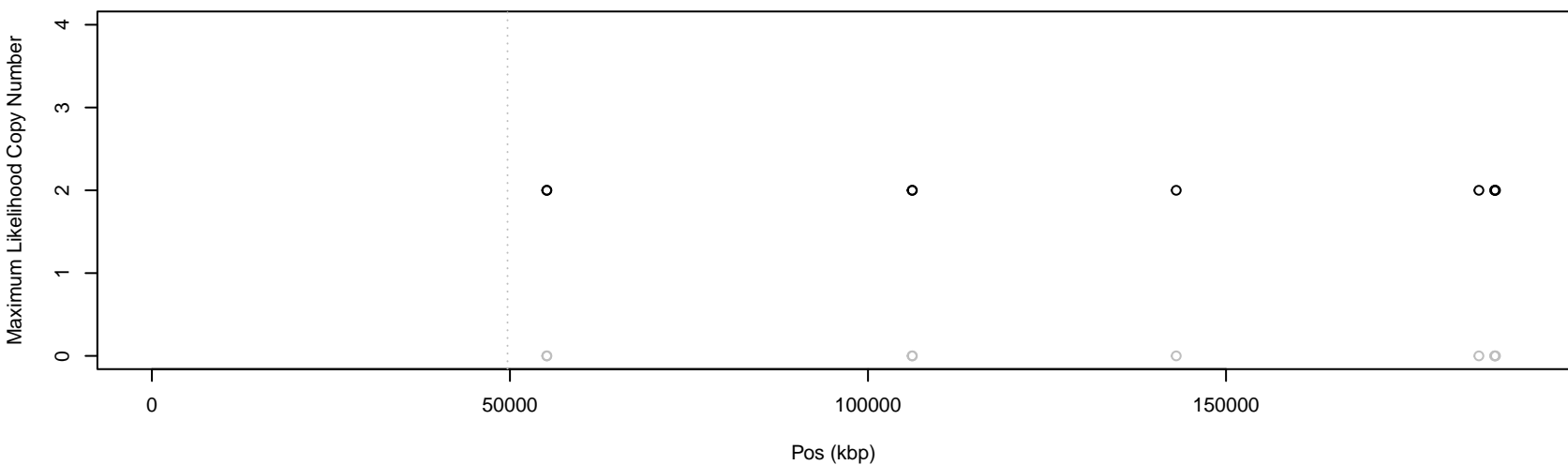
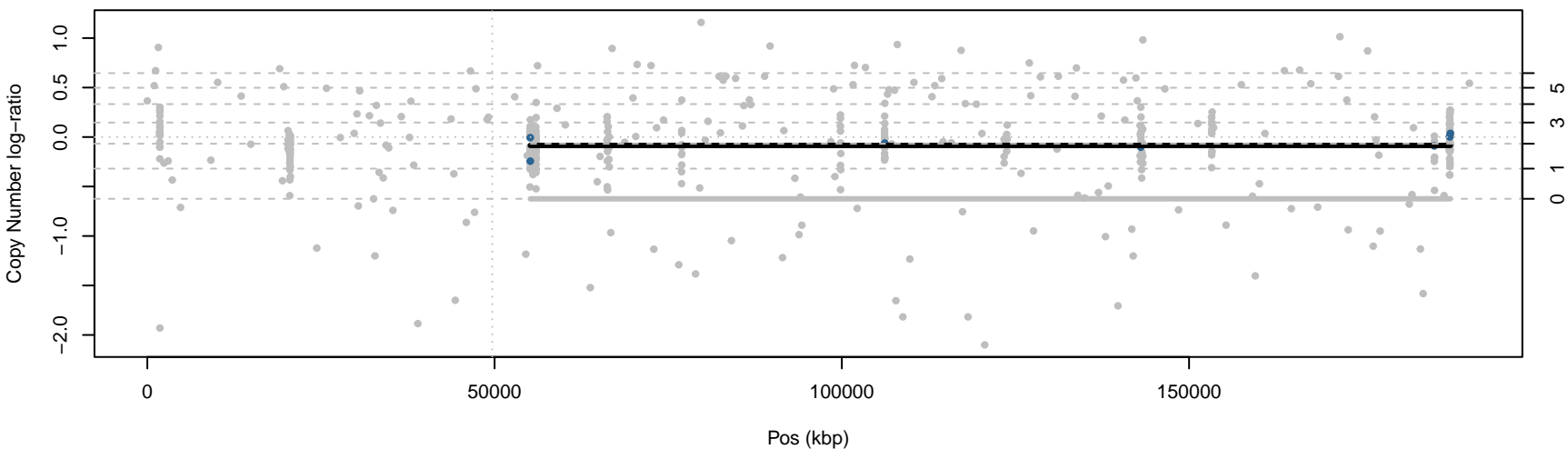
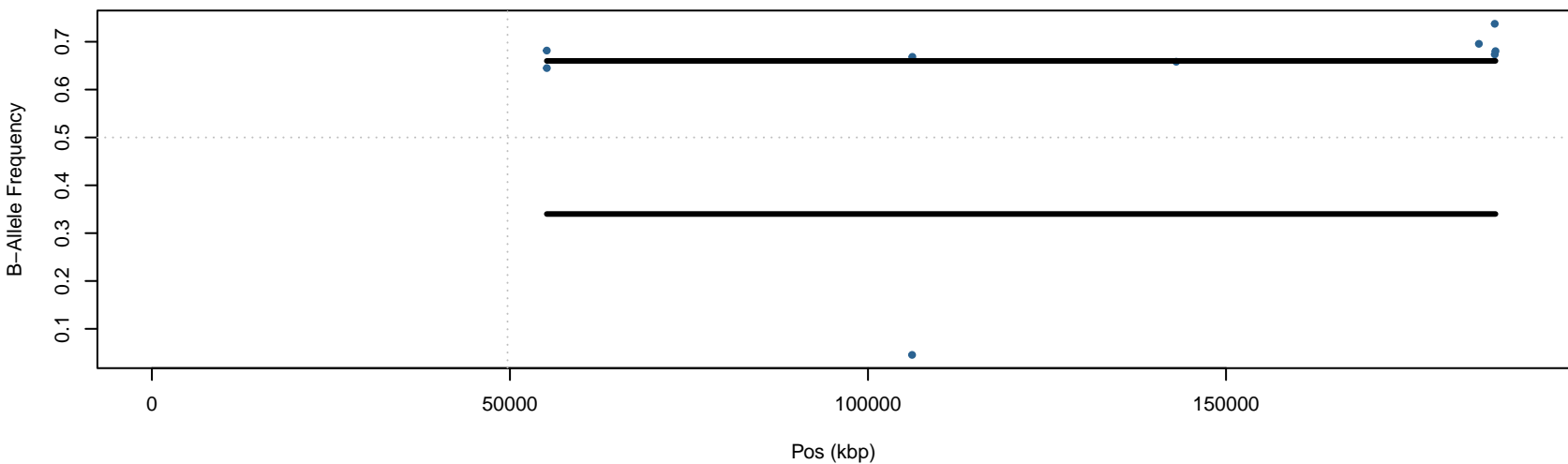
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 2



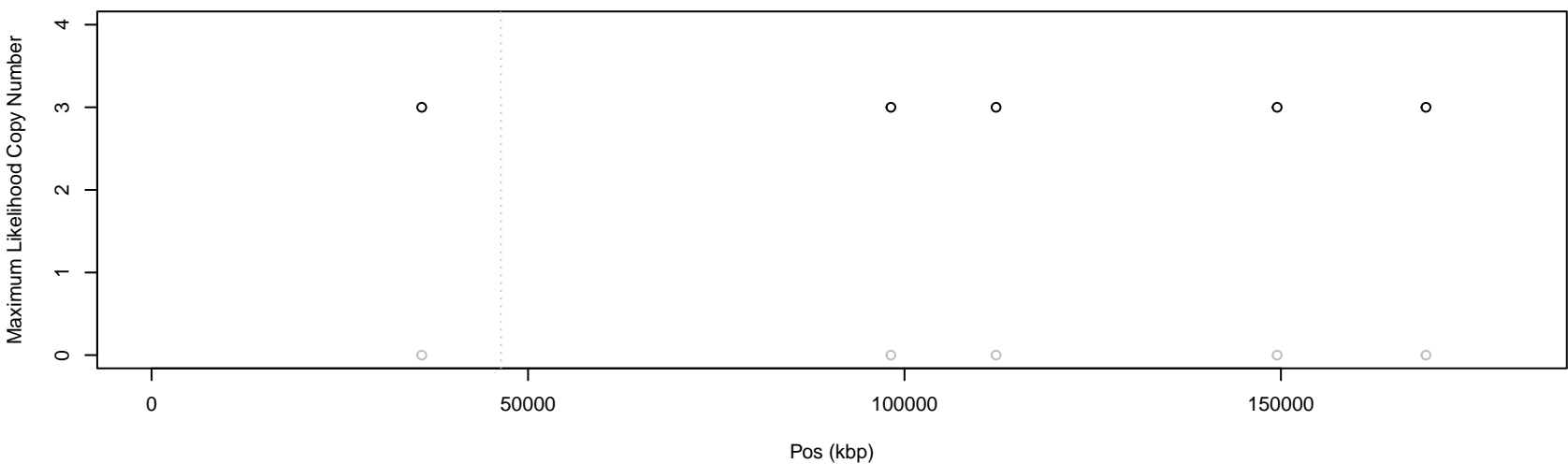
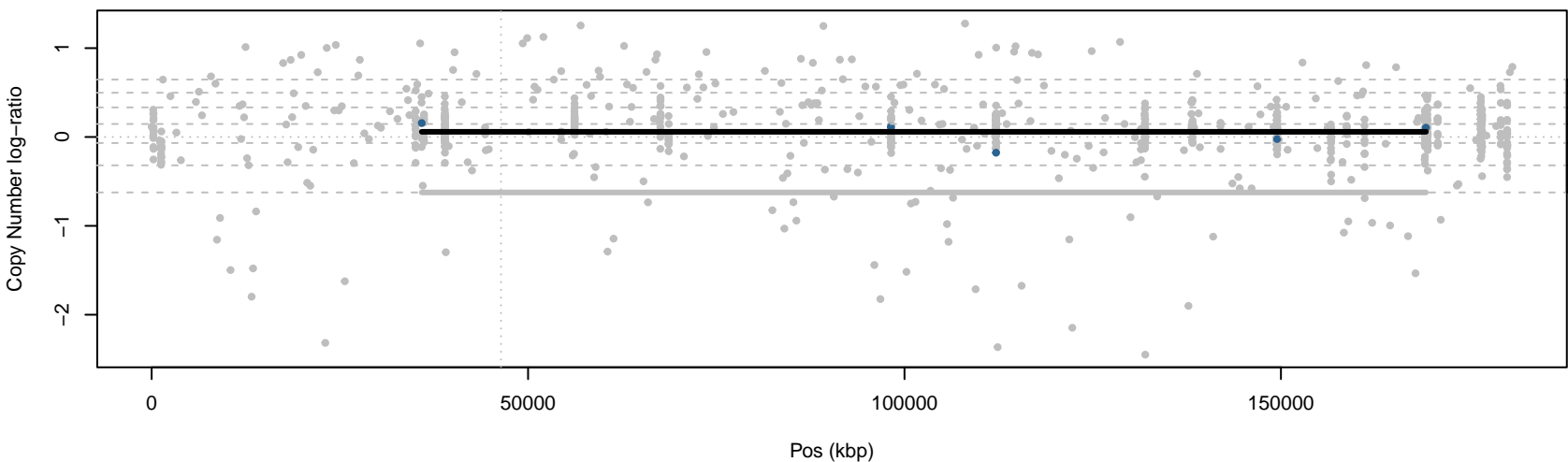
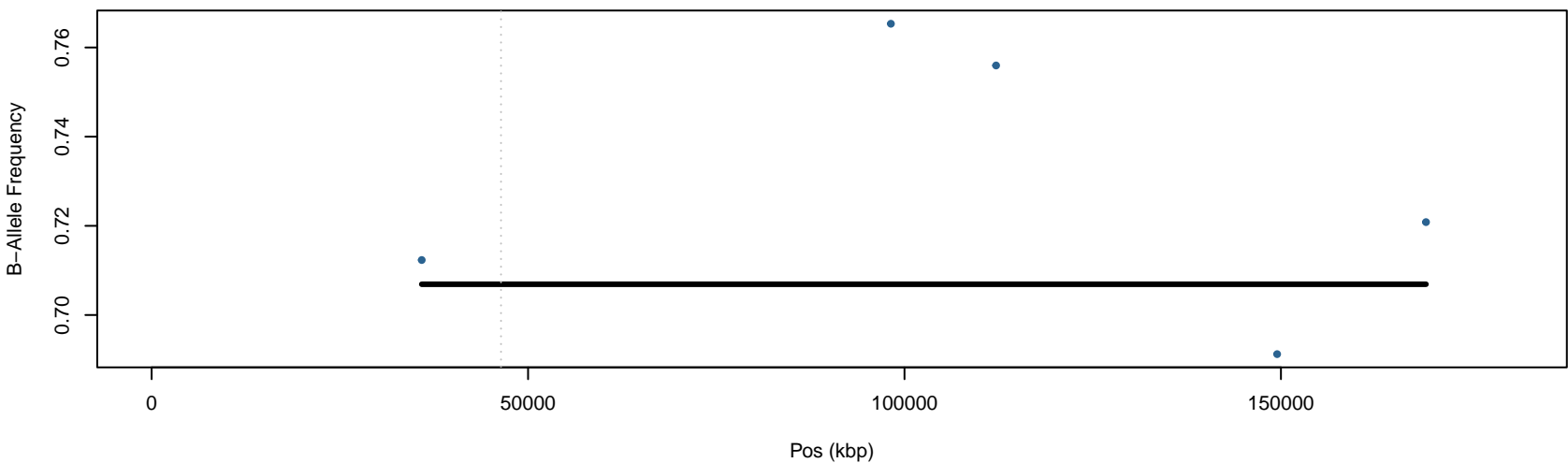
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 3



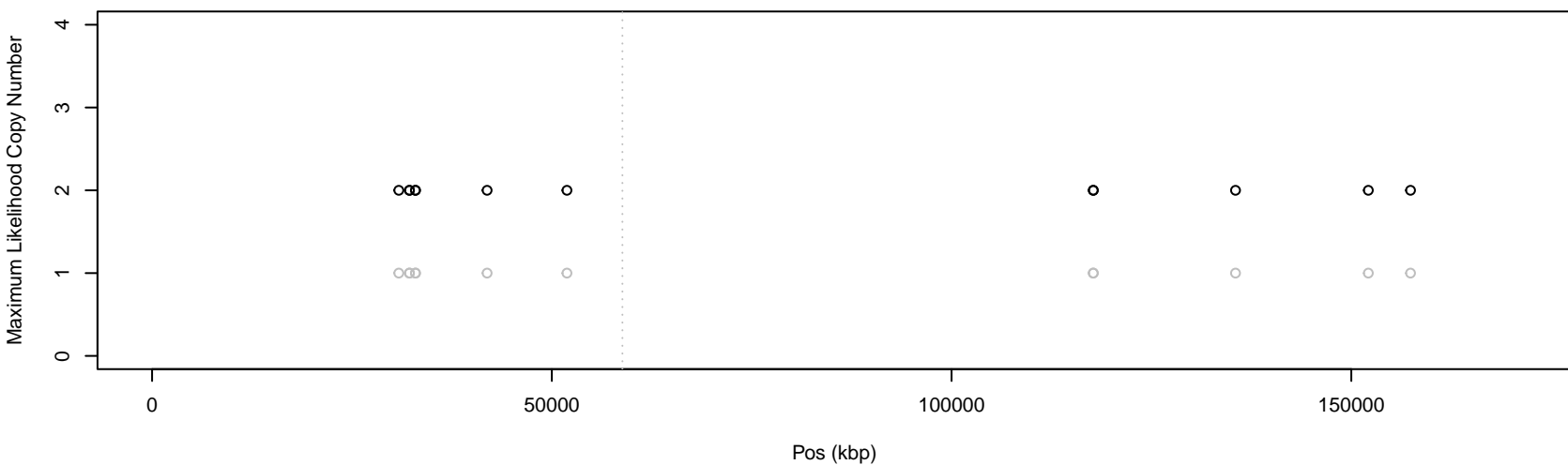
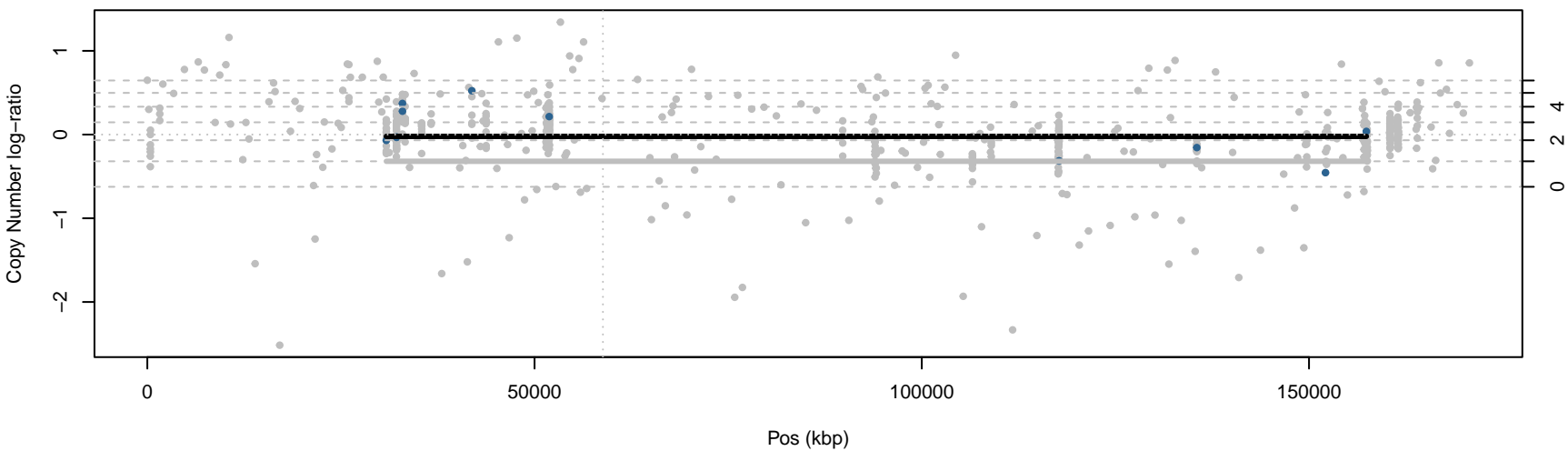
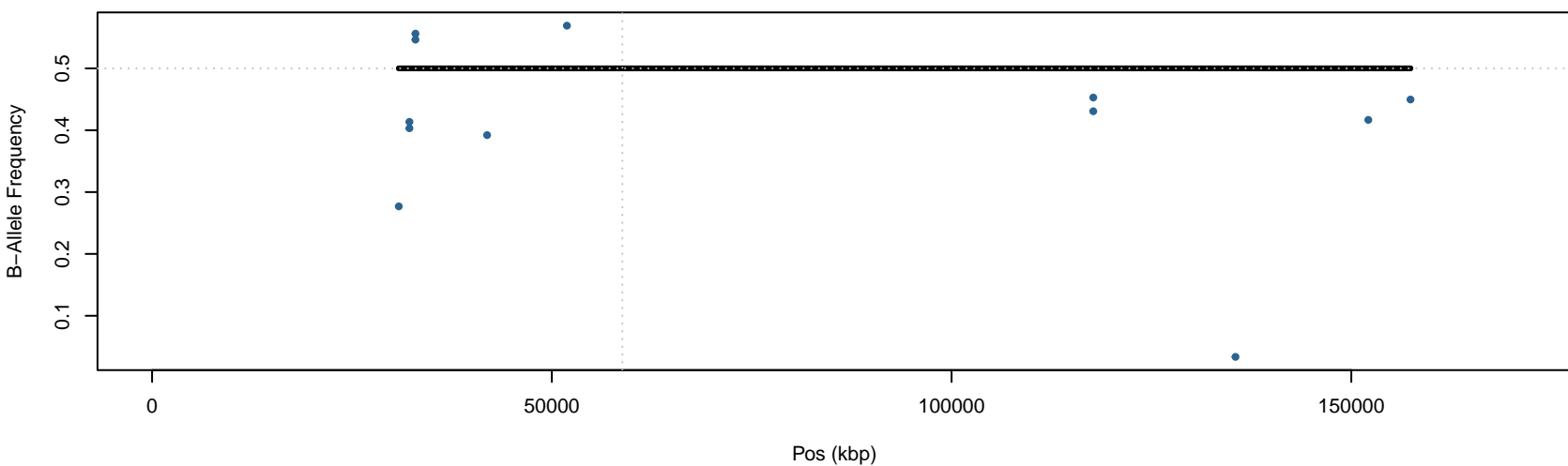
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 4



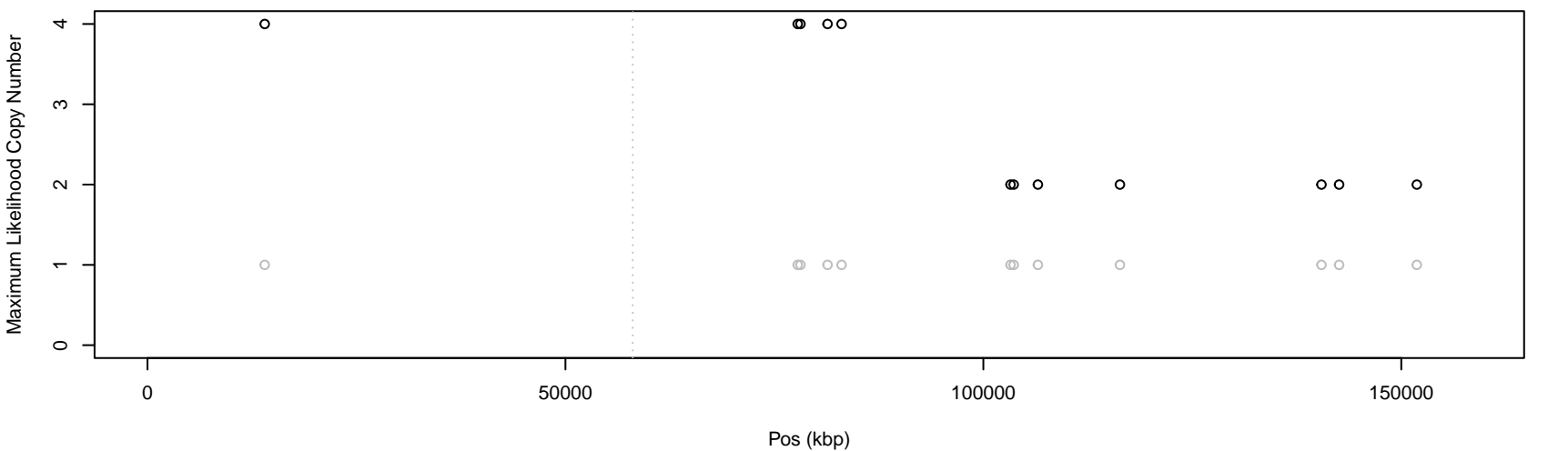
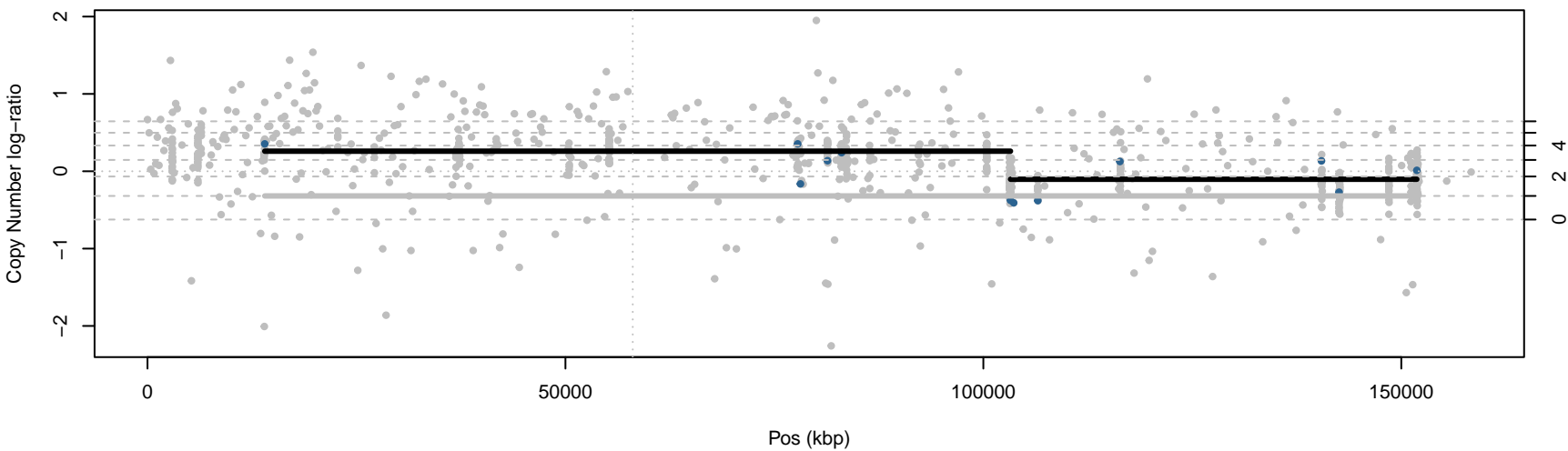
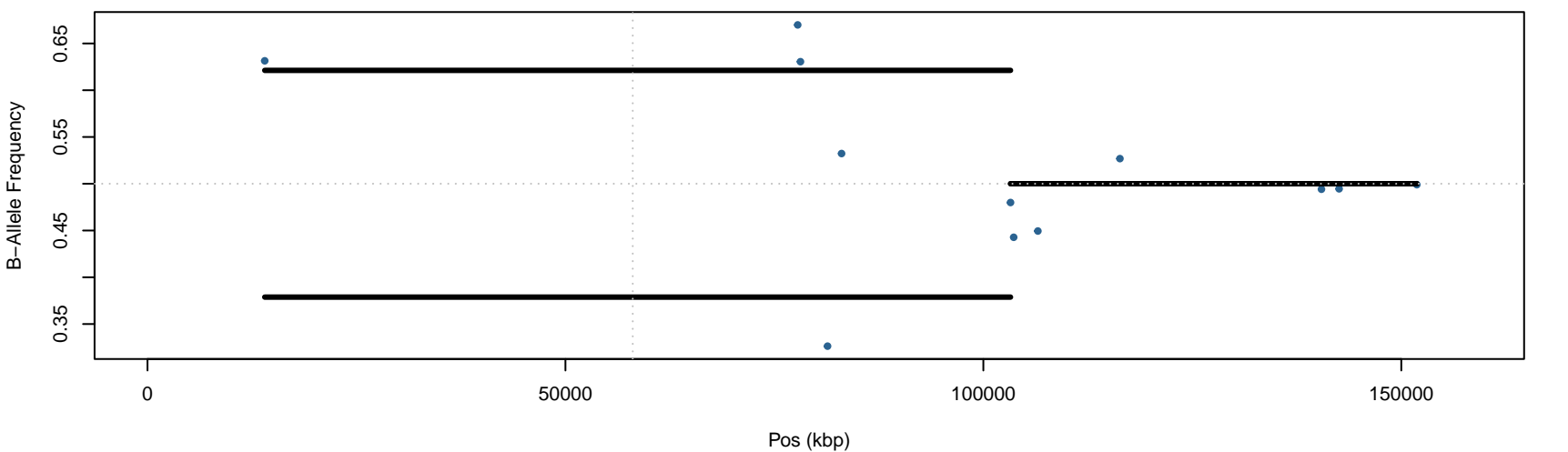
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 5



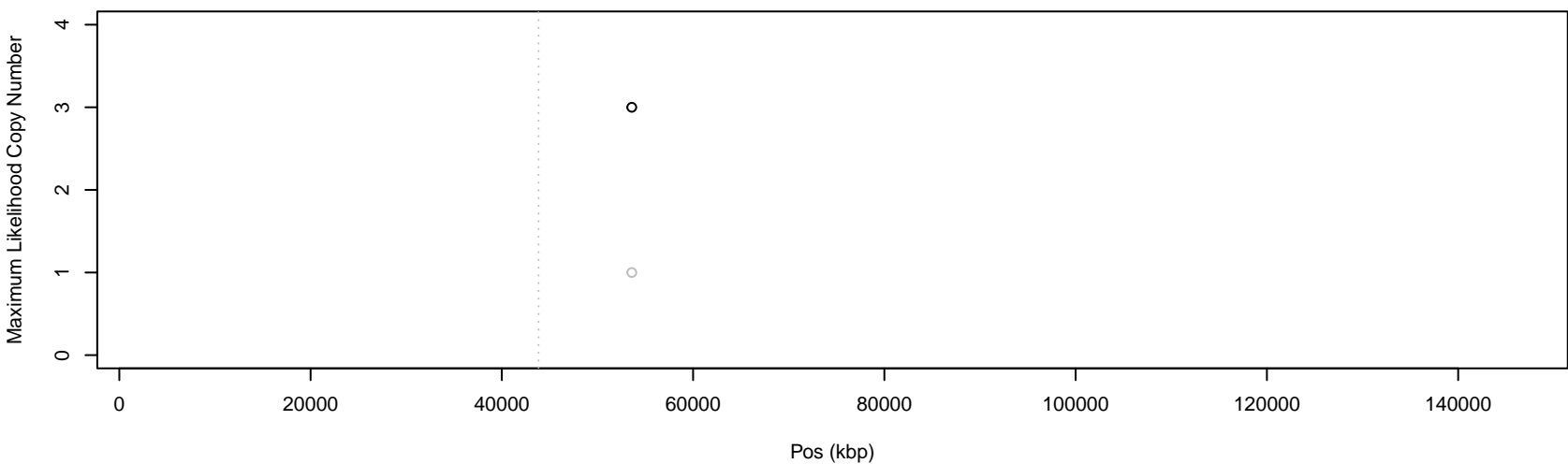
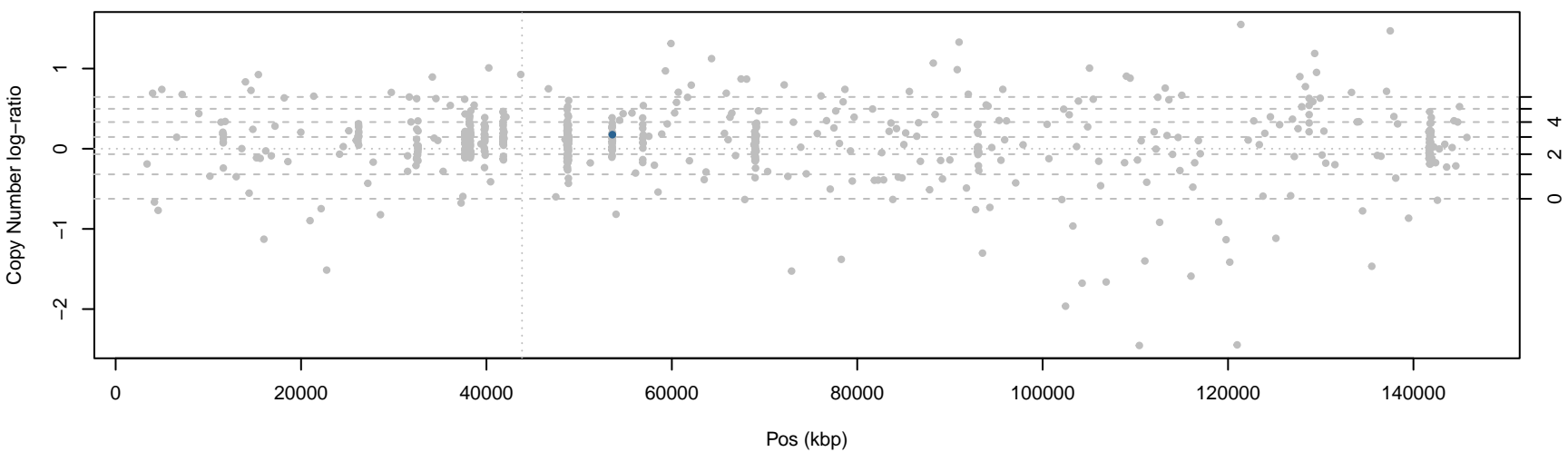
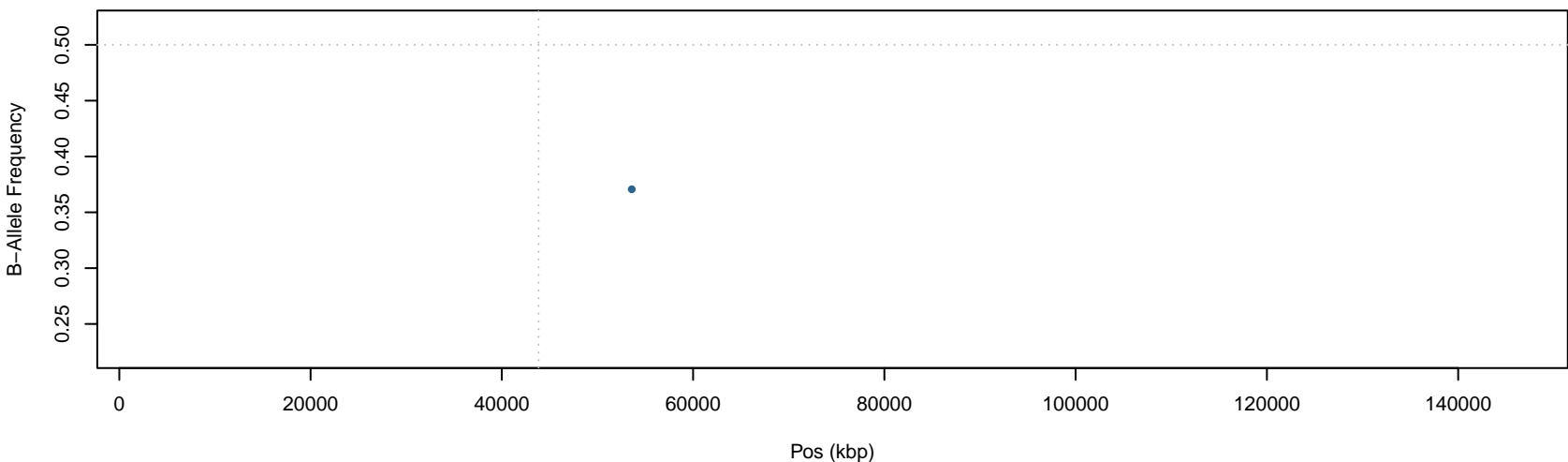
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 6



Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 7

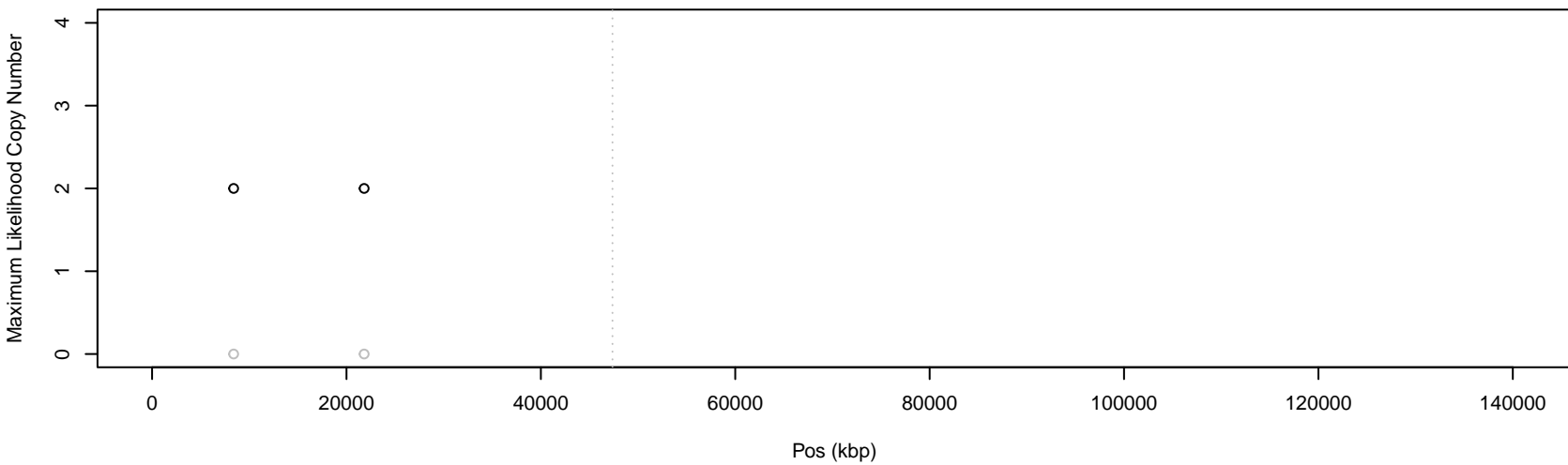
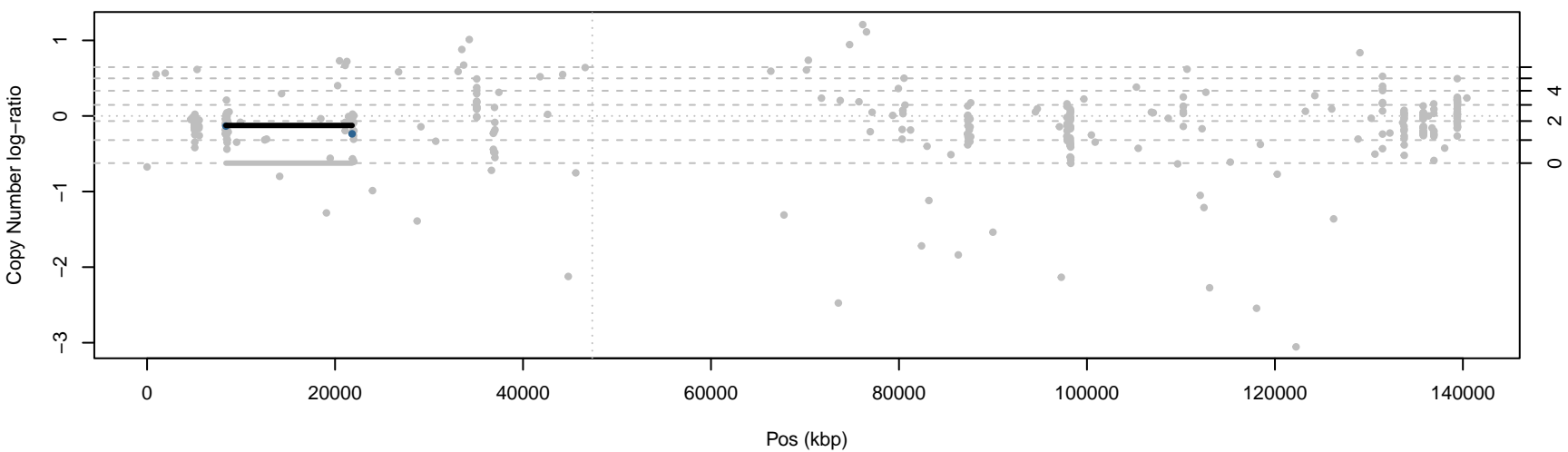
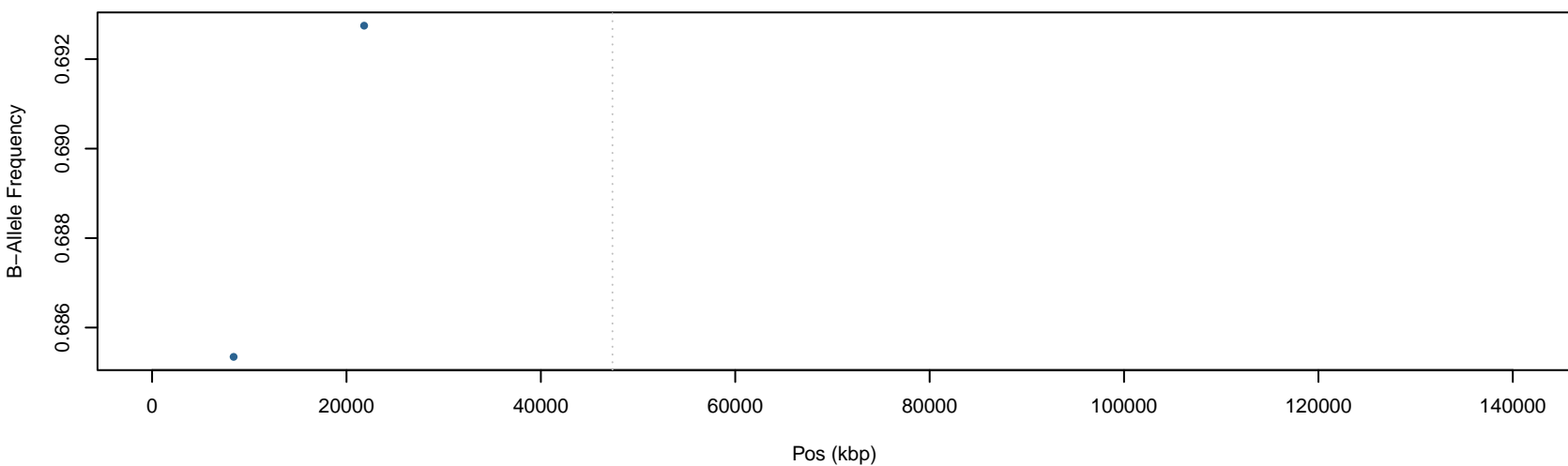


Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 8

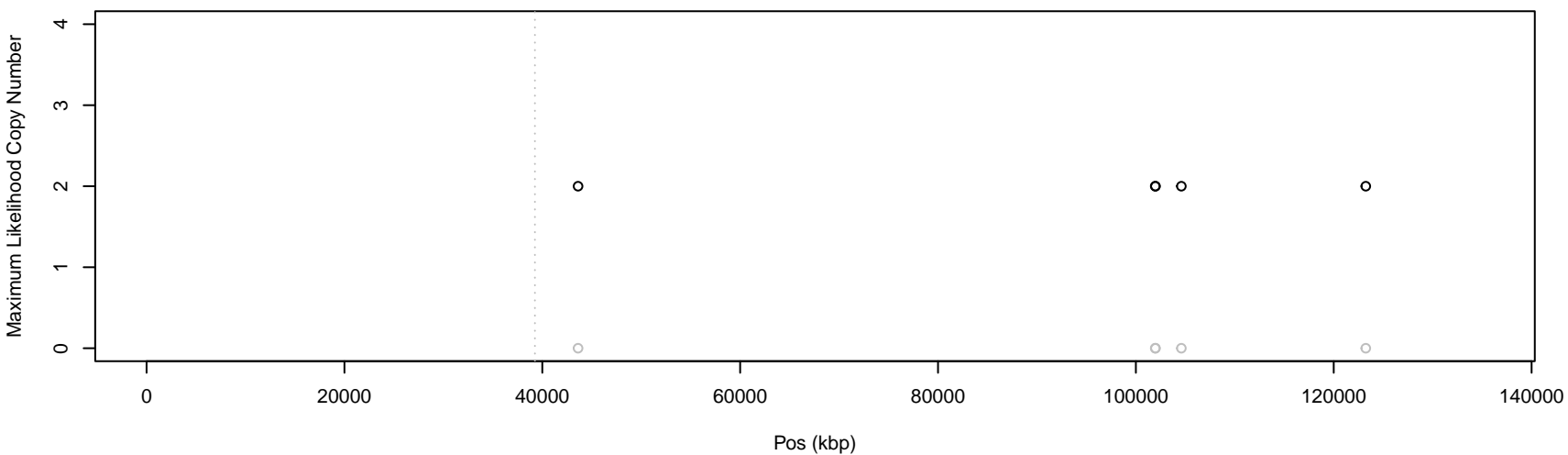
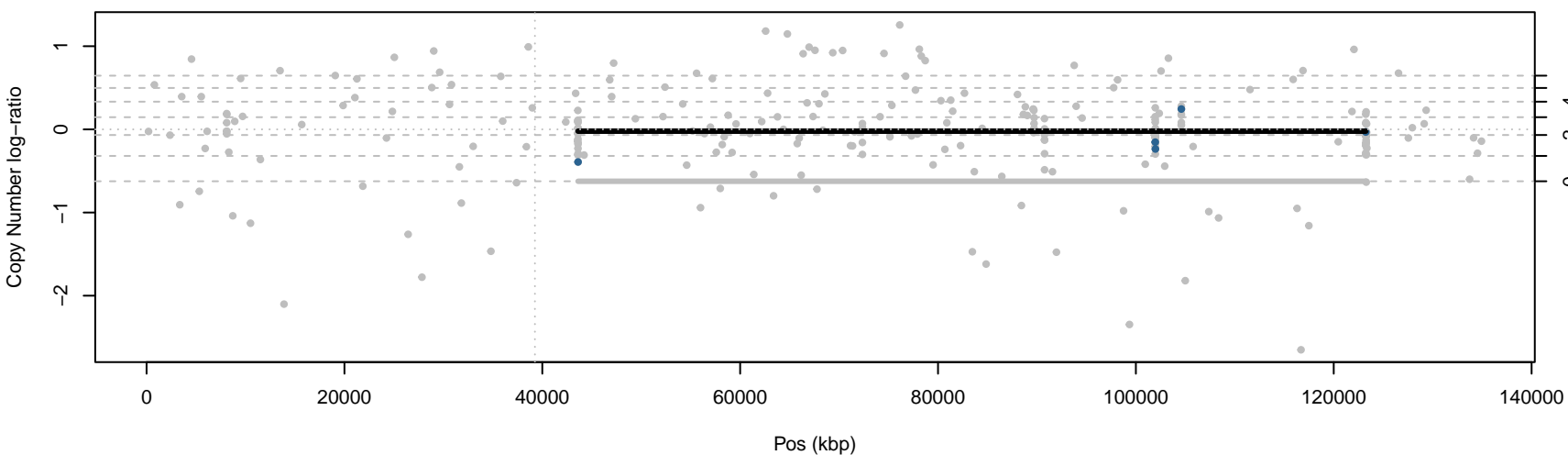
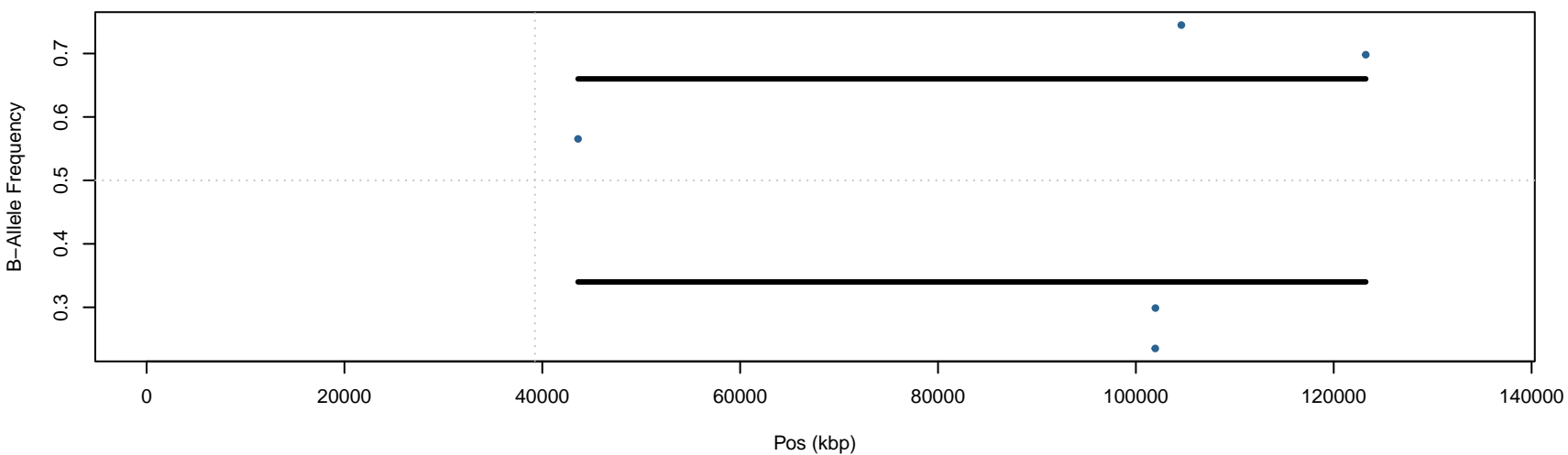




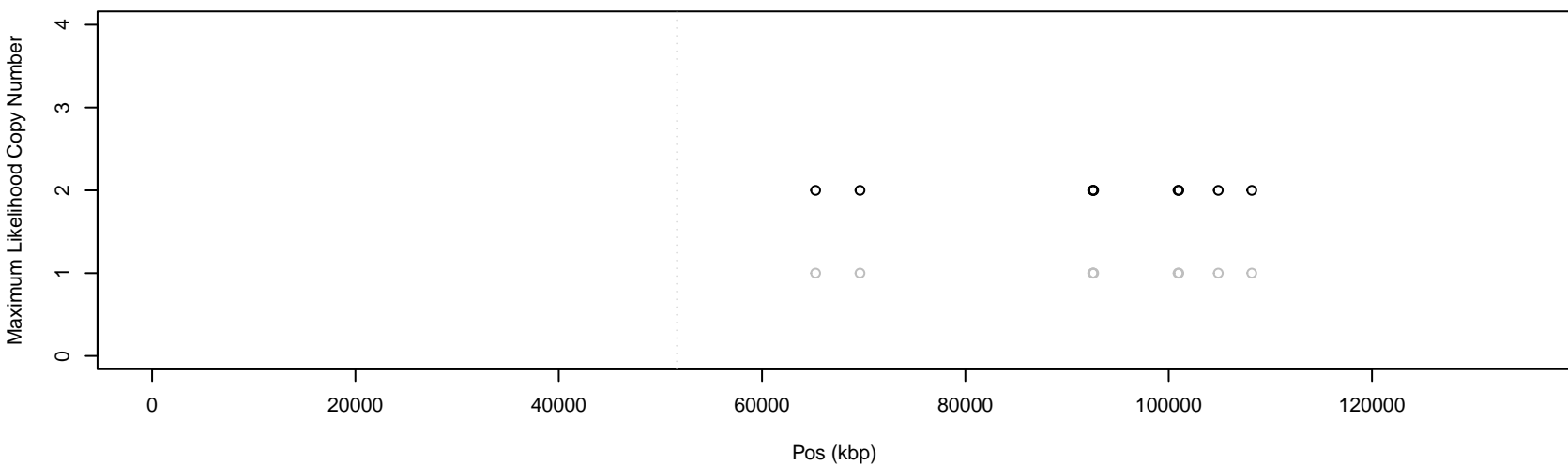
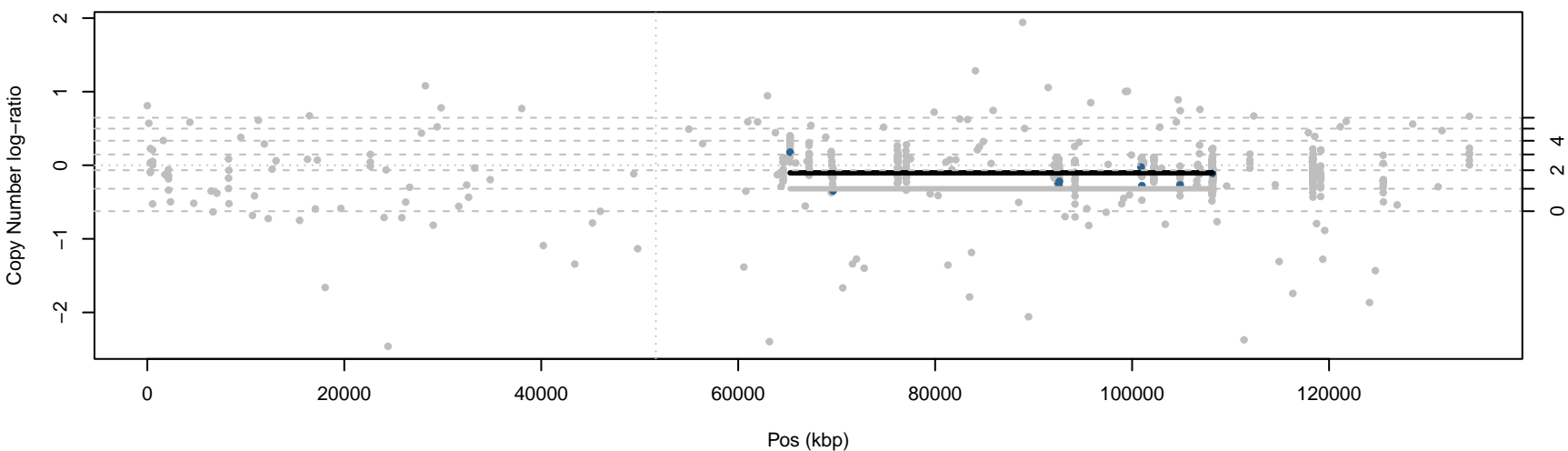
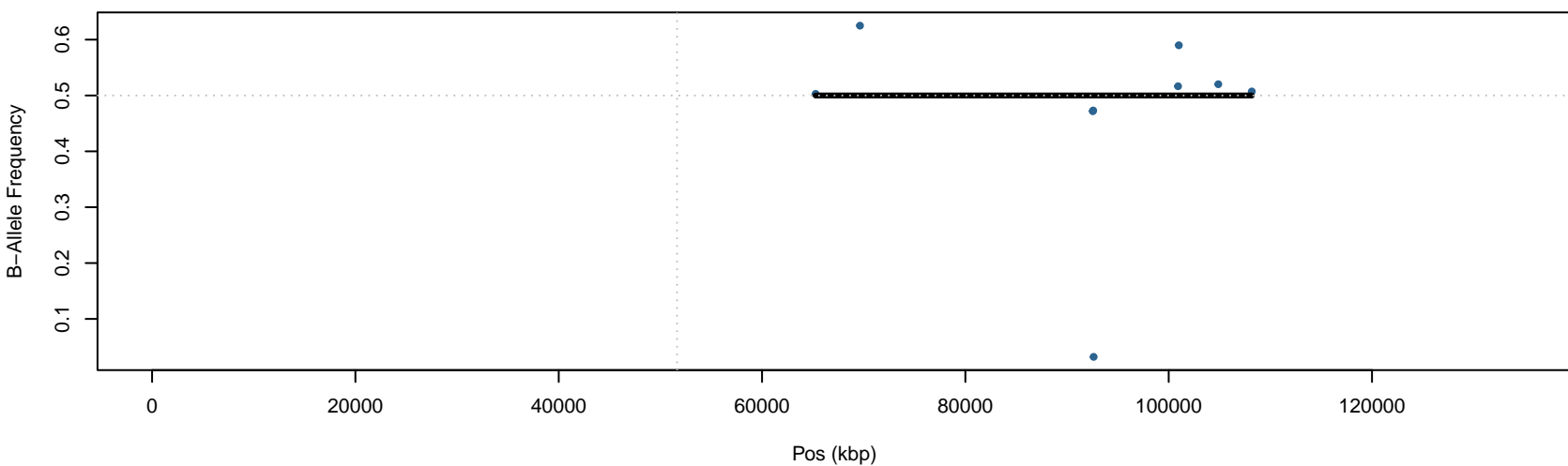
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 9



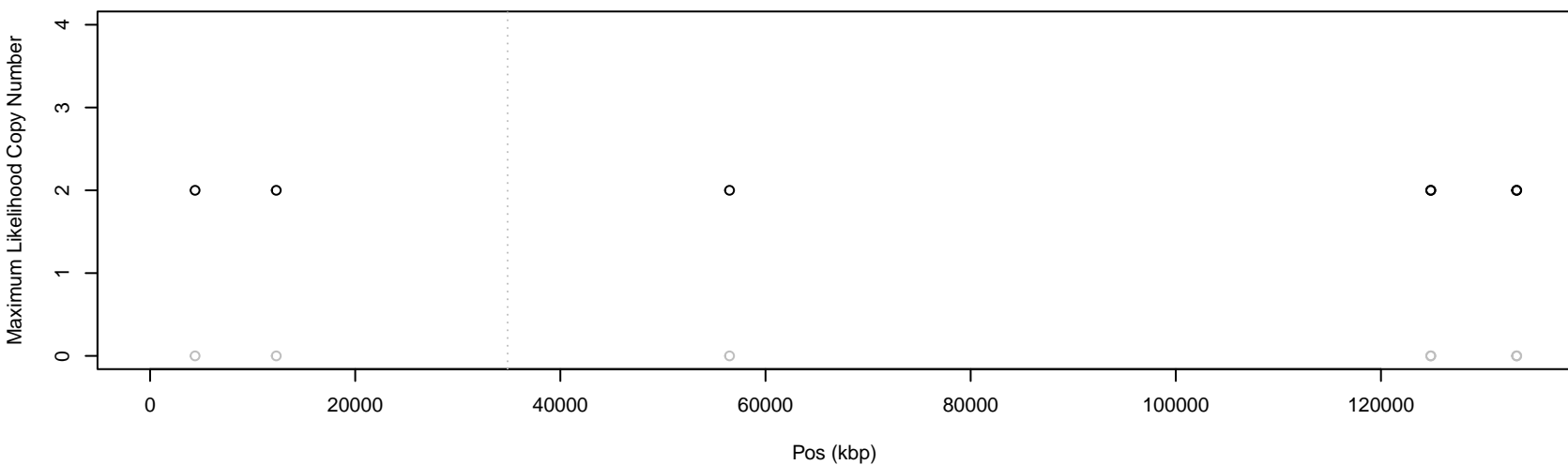
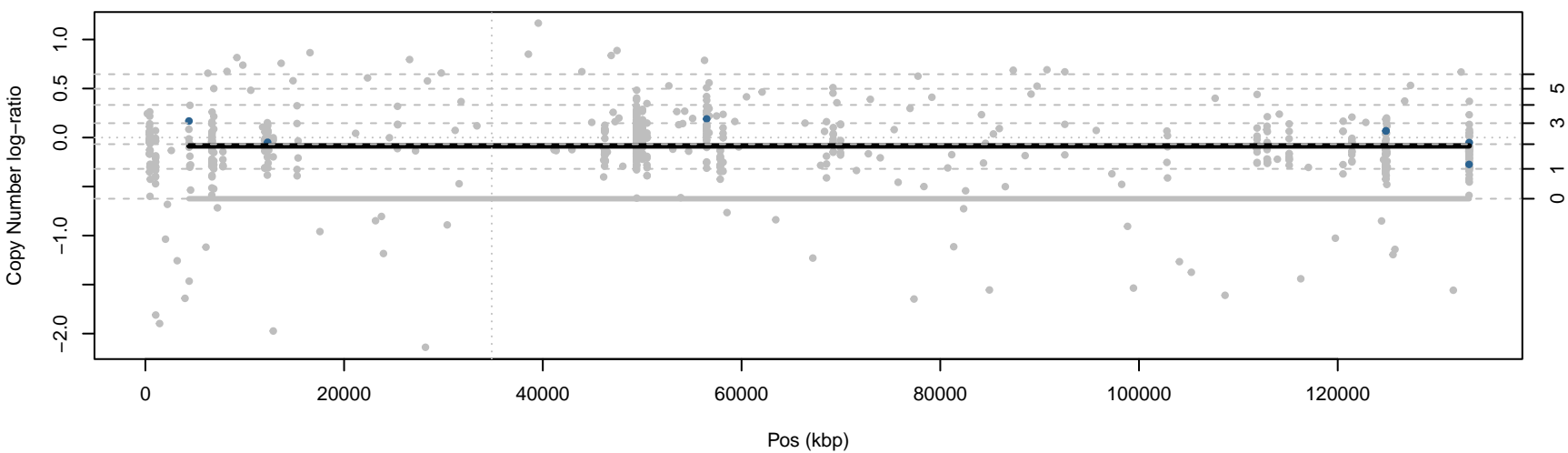
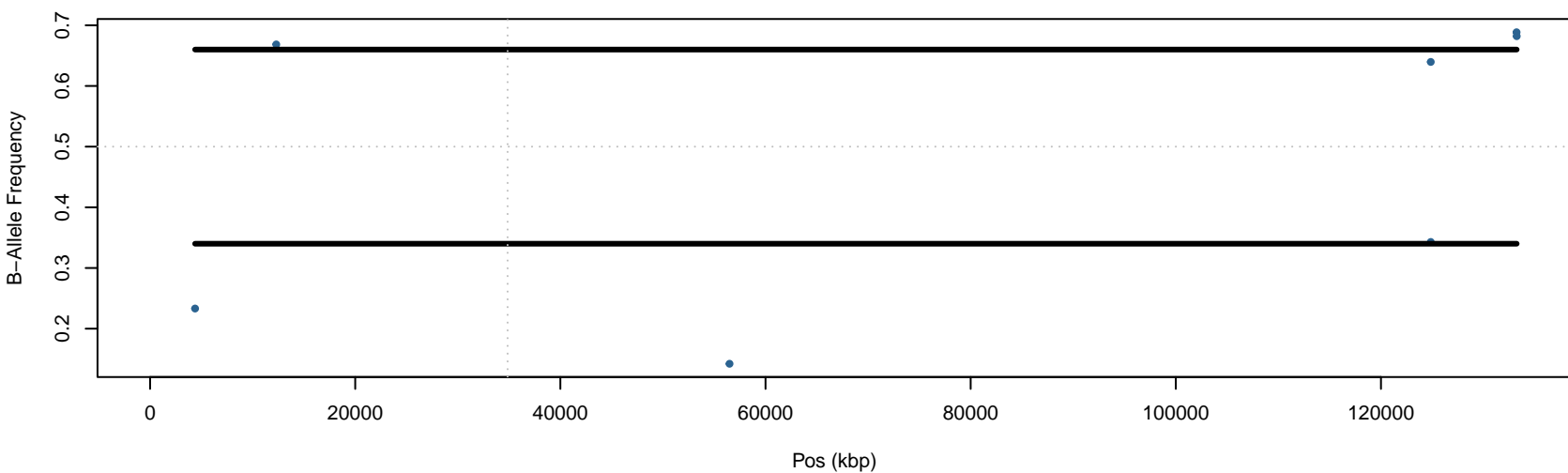
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 10



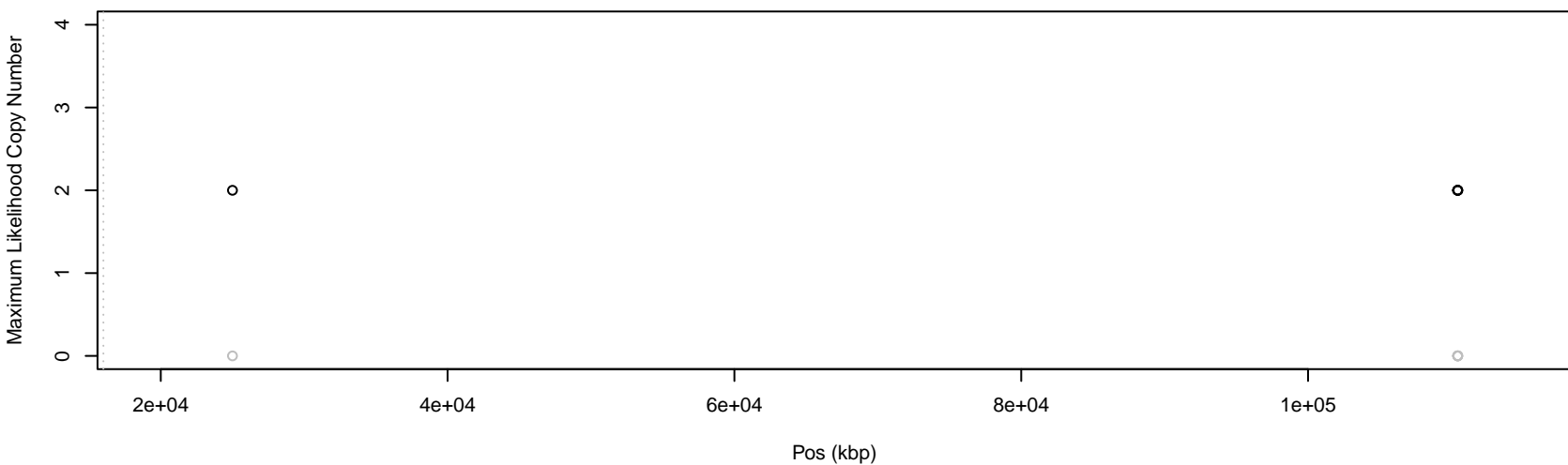
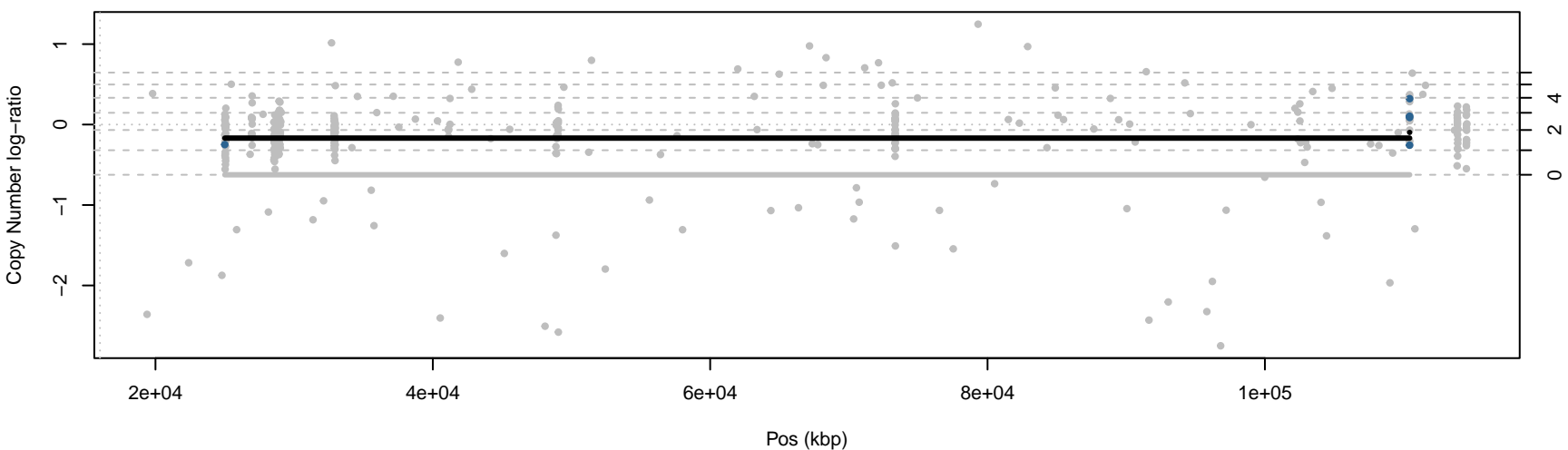
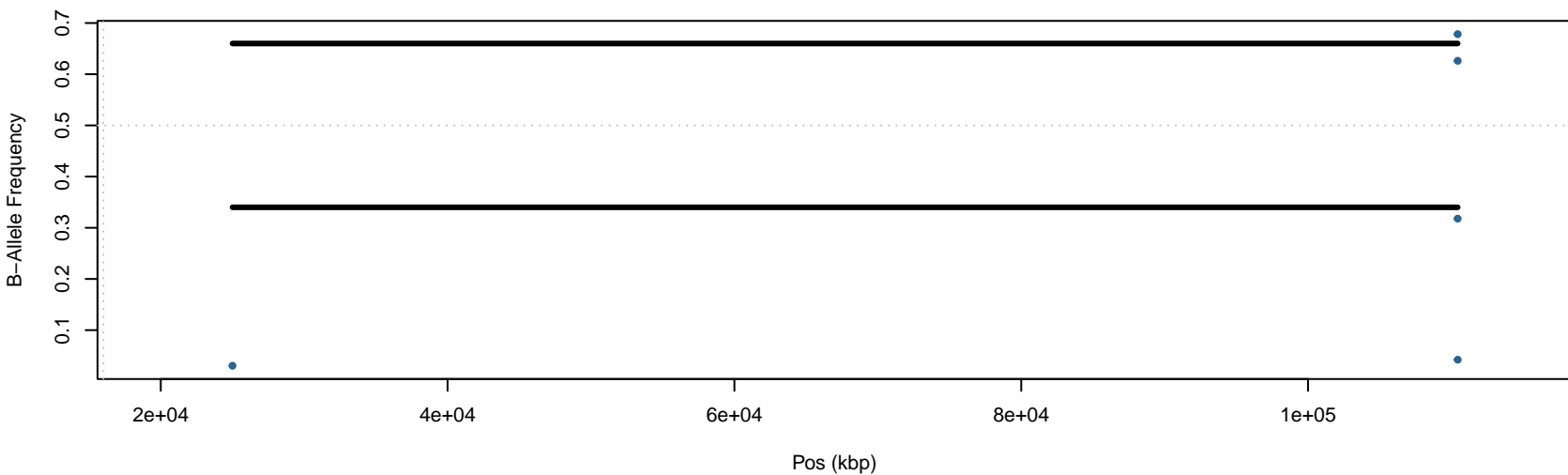
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 11



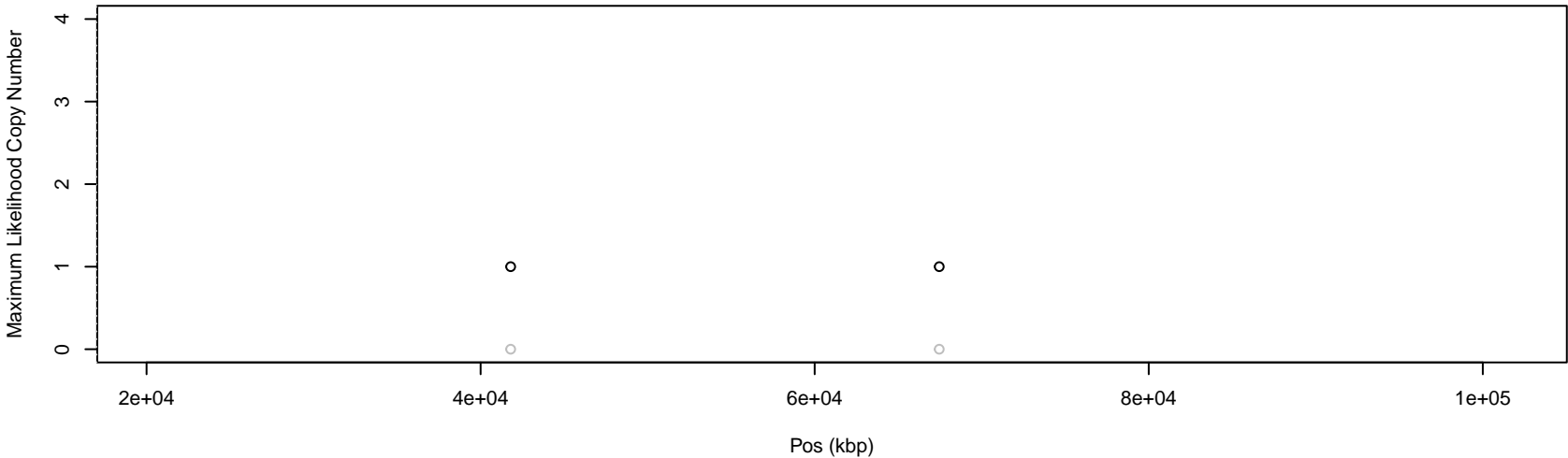
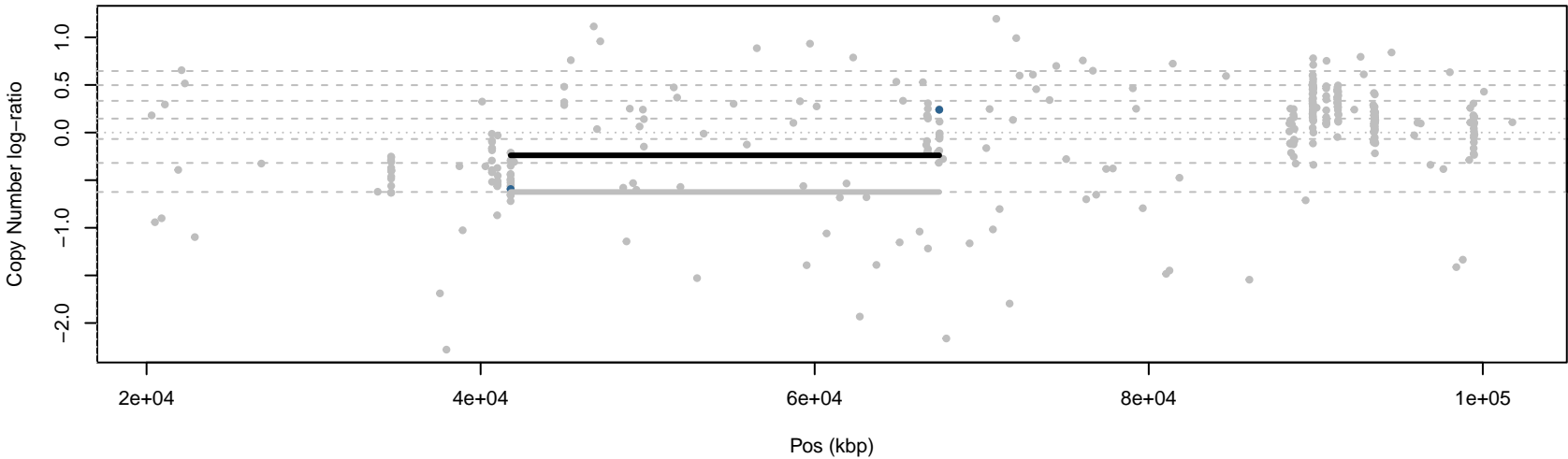
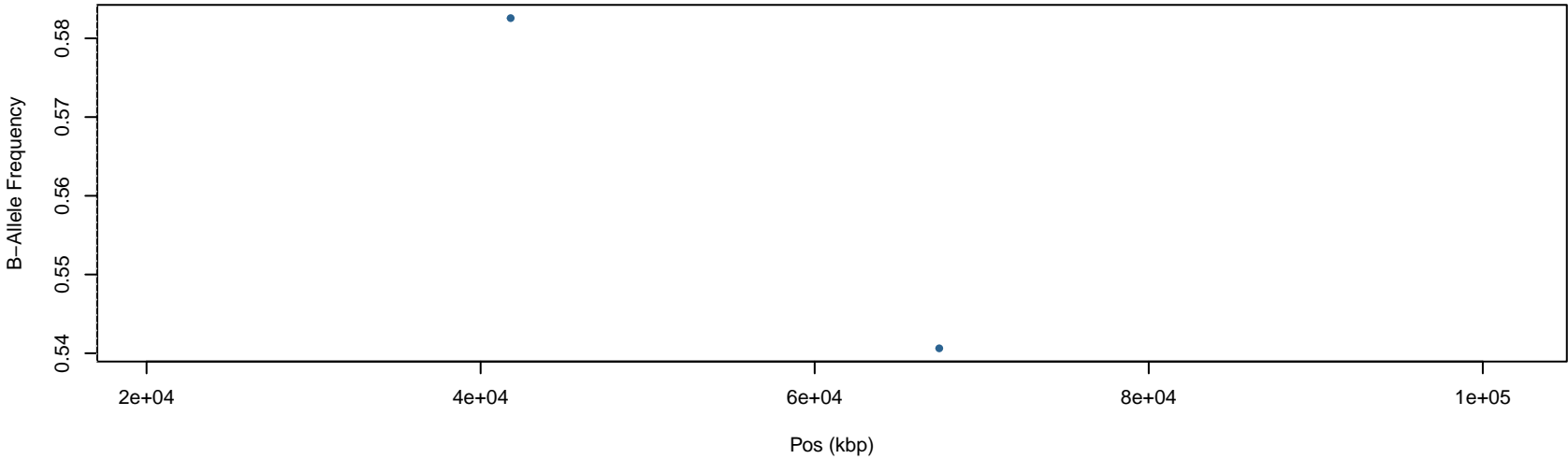
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 12



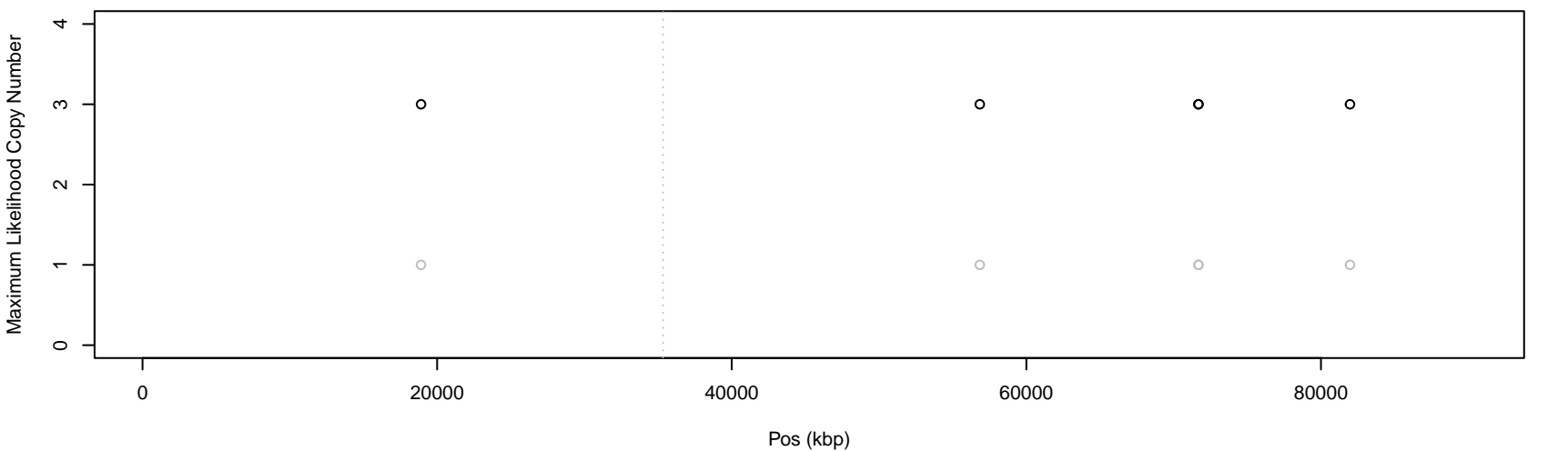
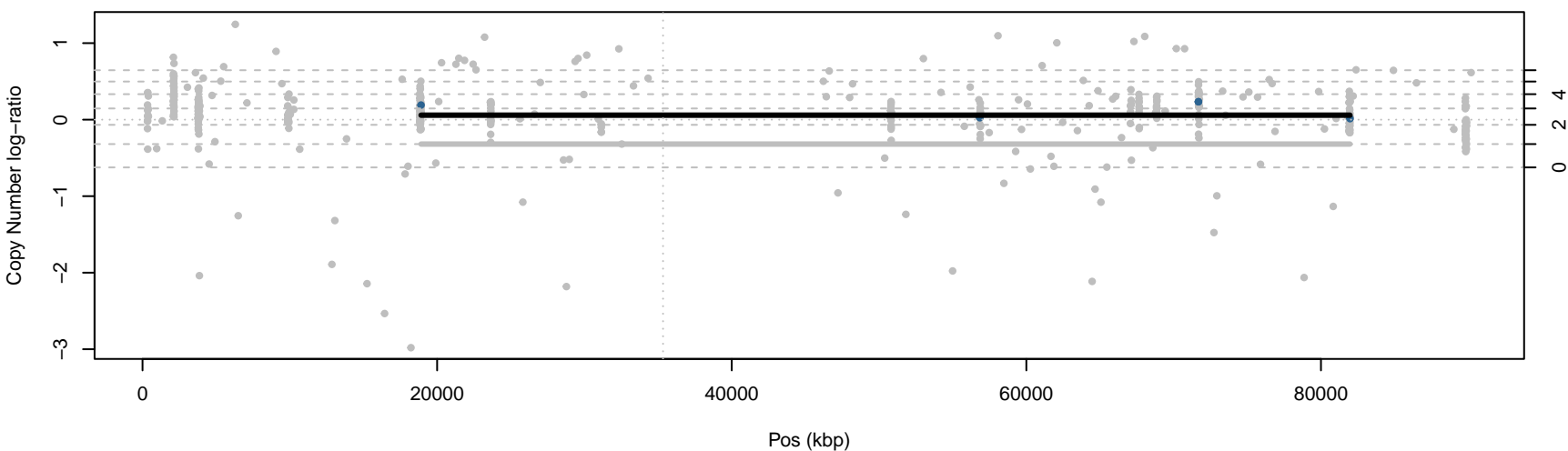
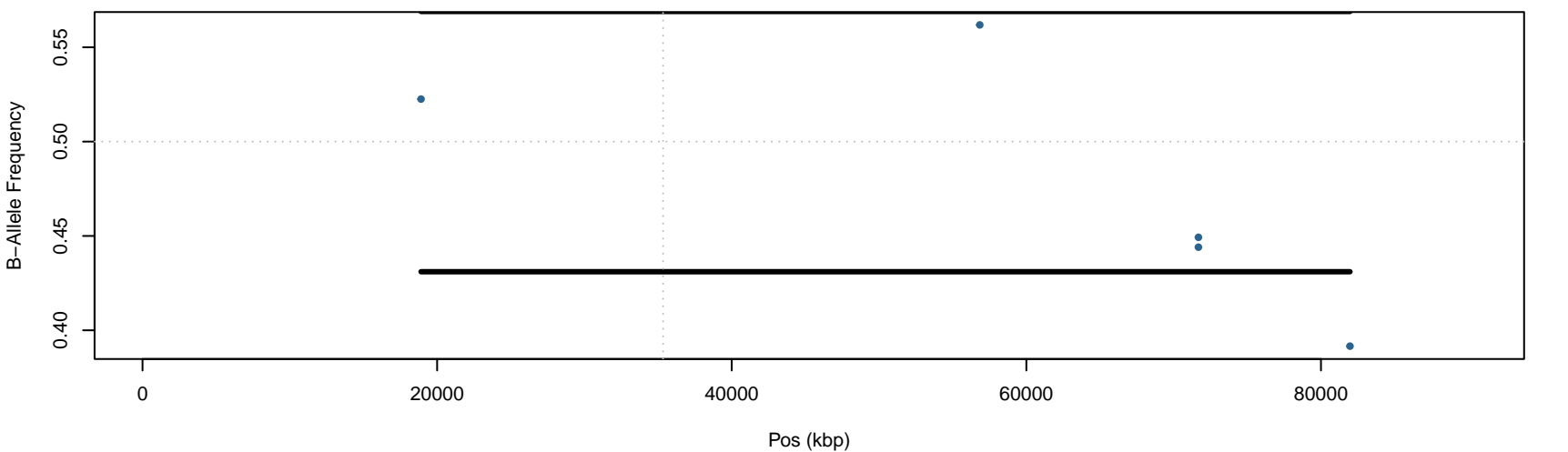
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 13



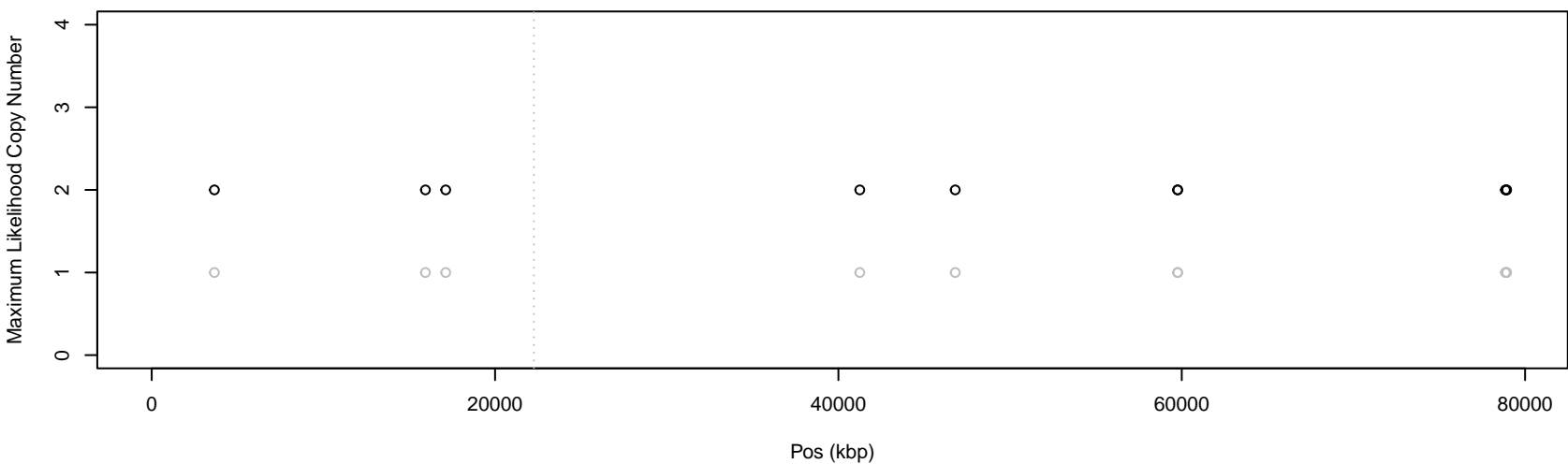
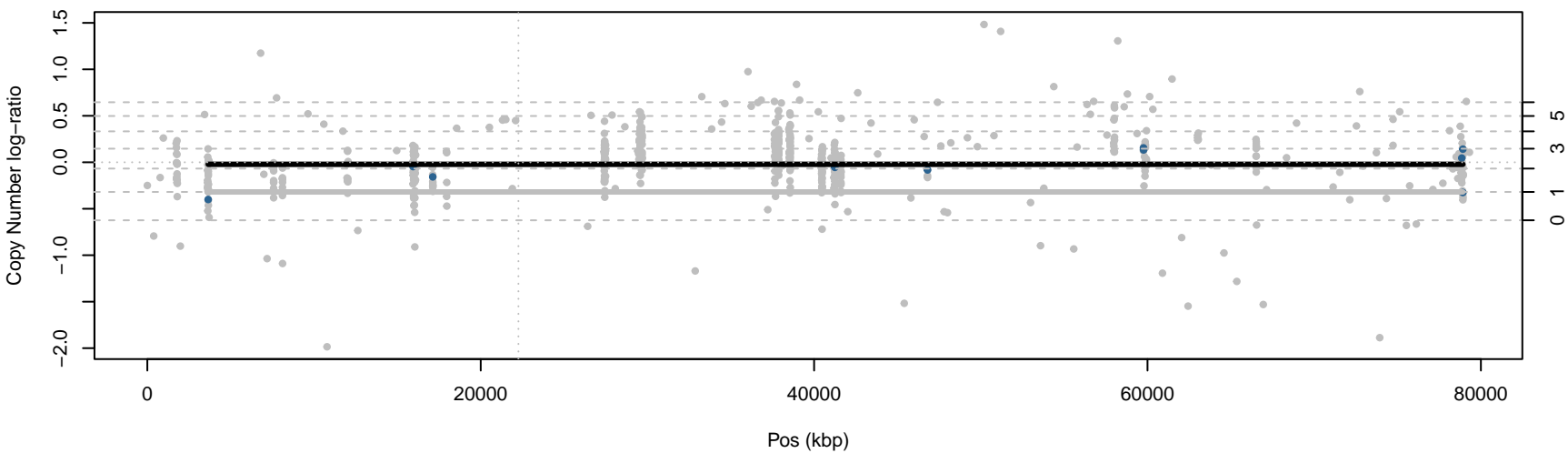
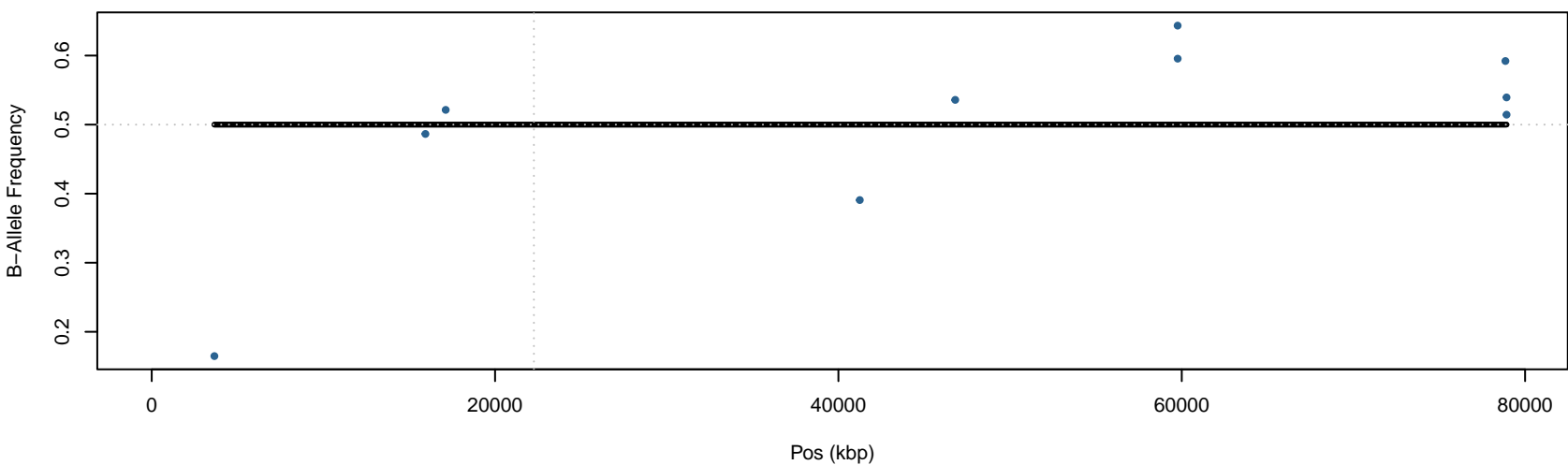
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 15



Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 16

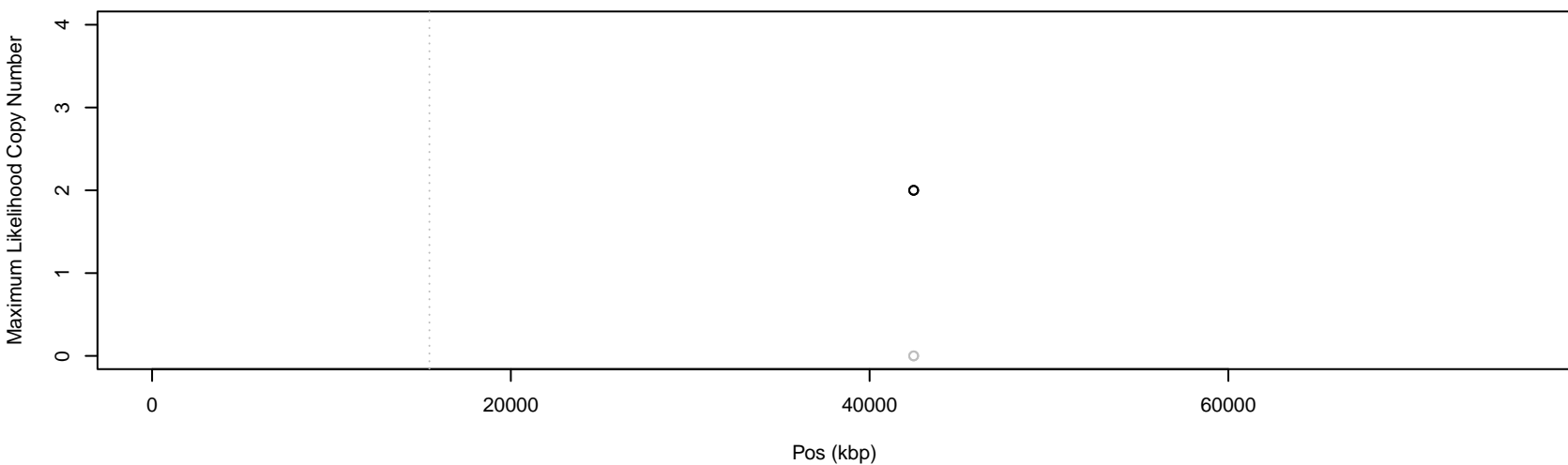
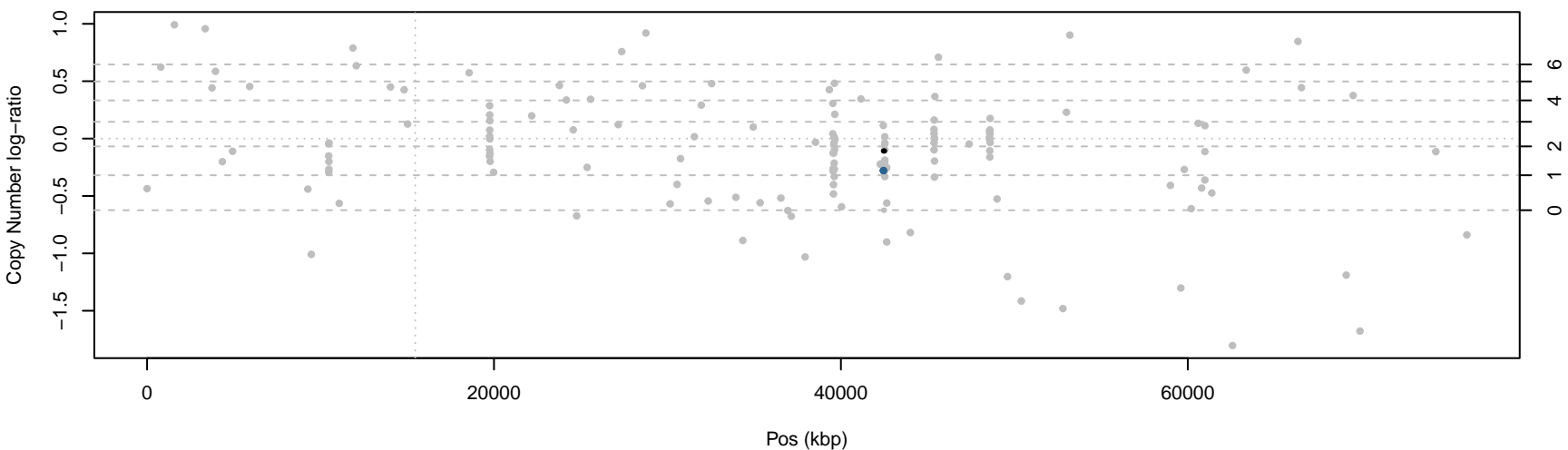
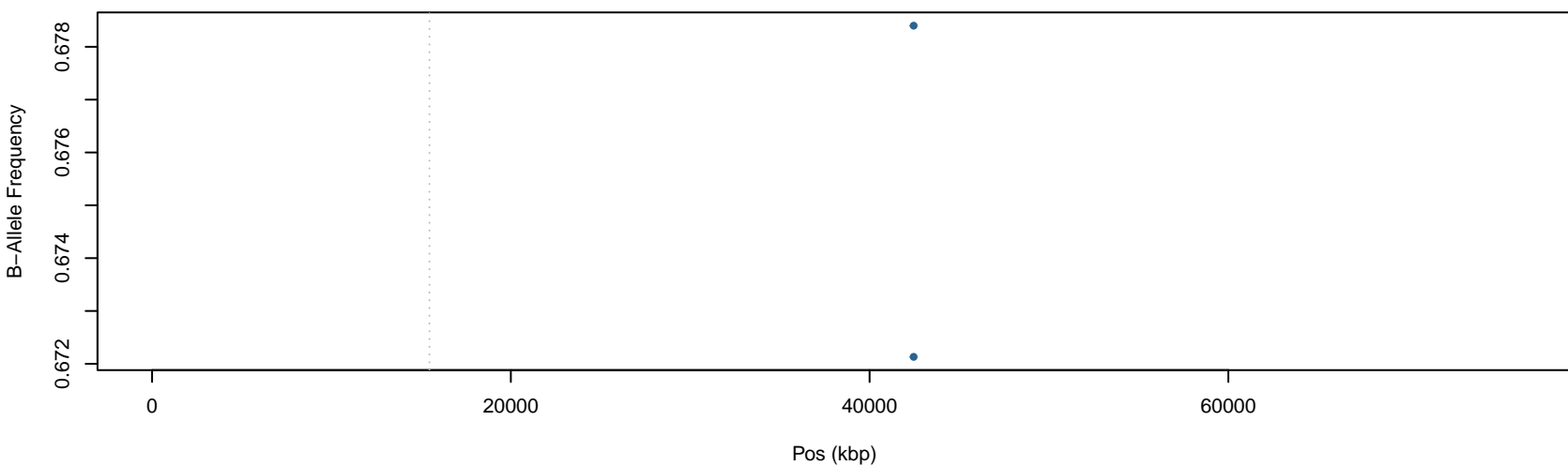


Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 17

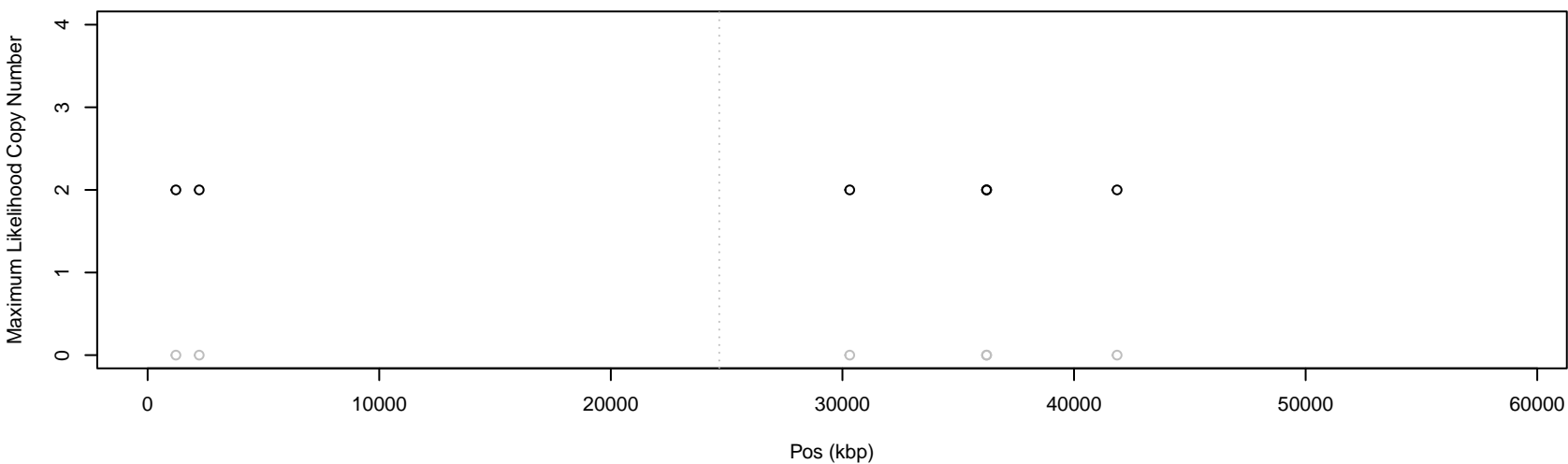
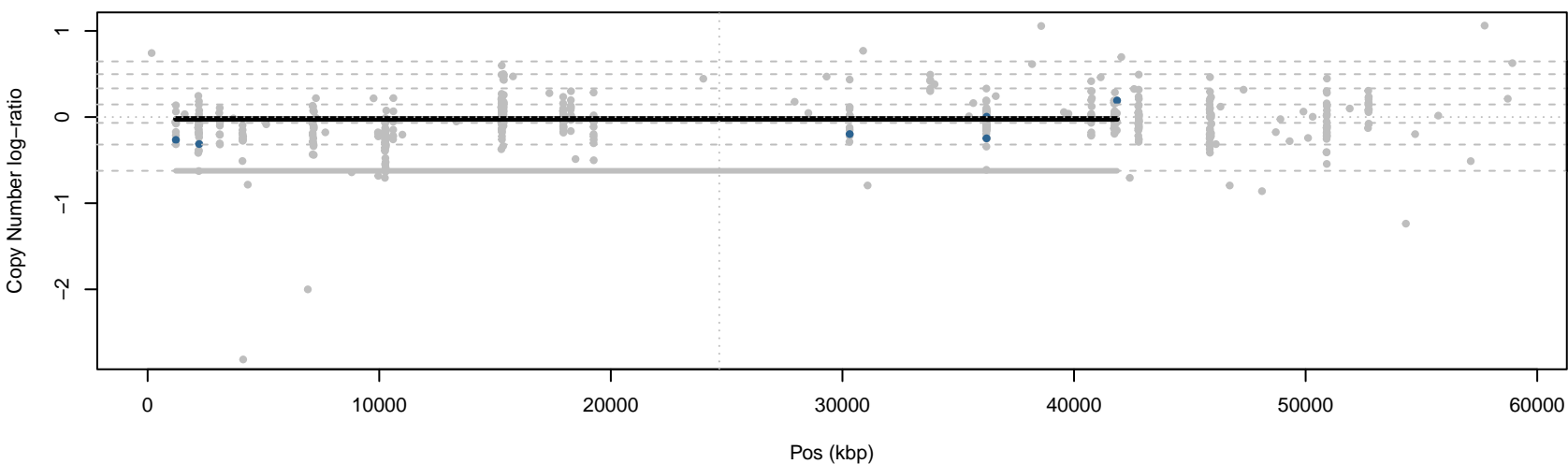
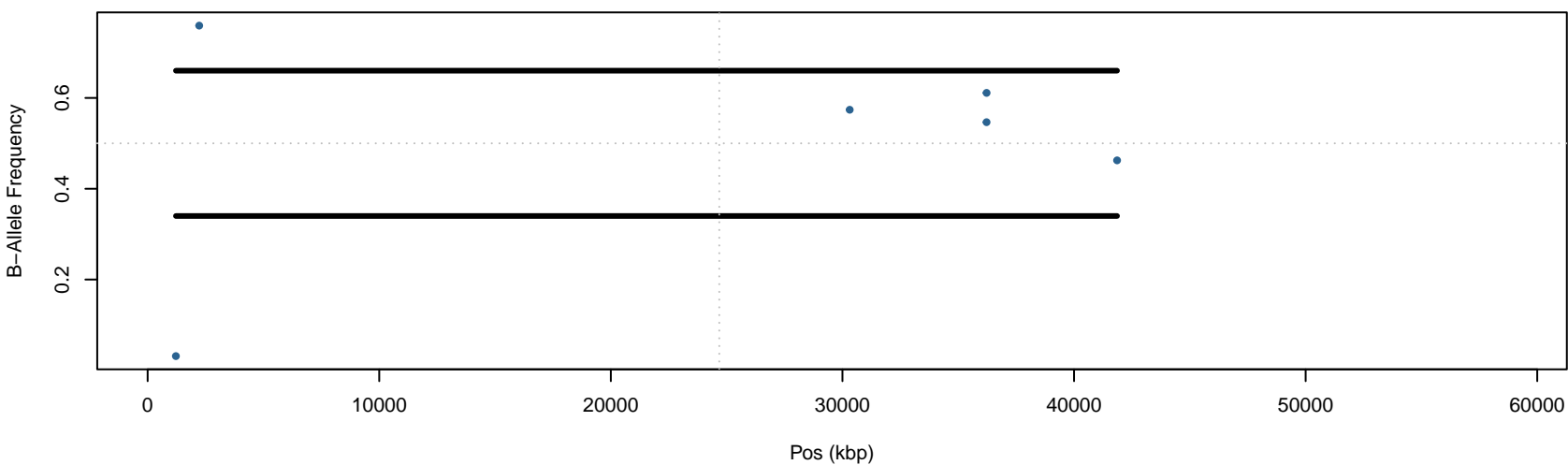




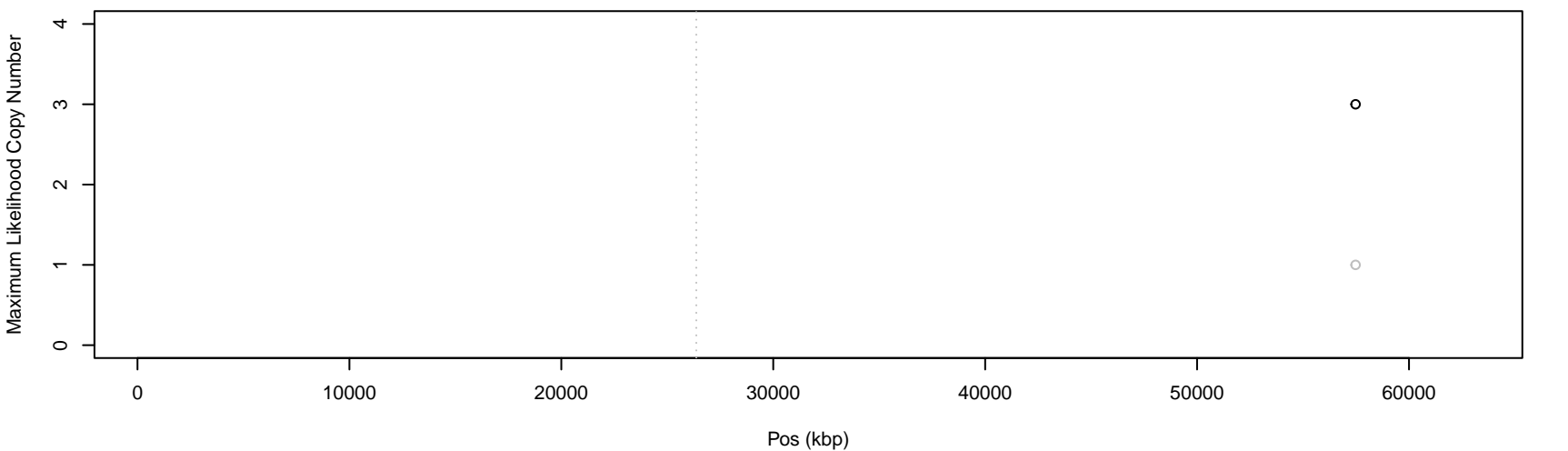
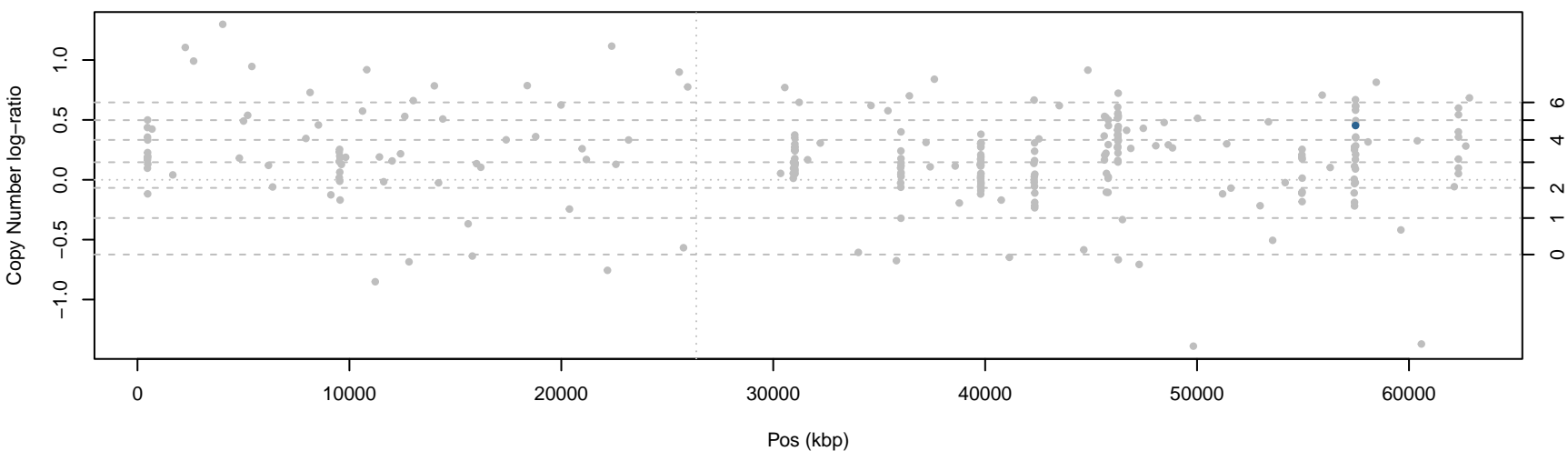
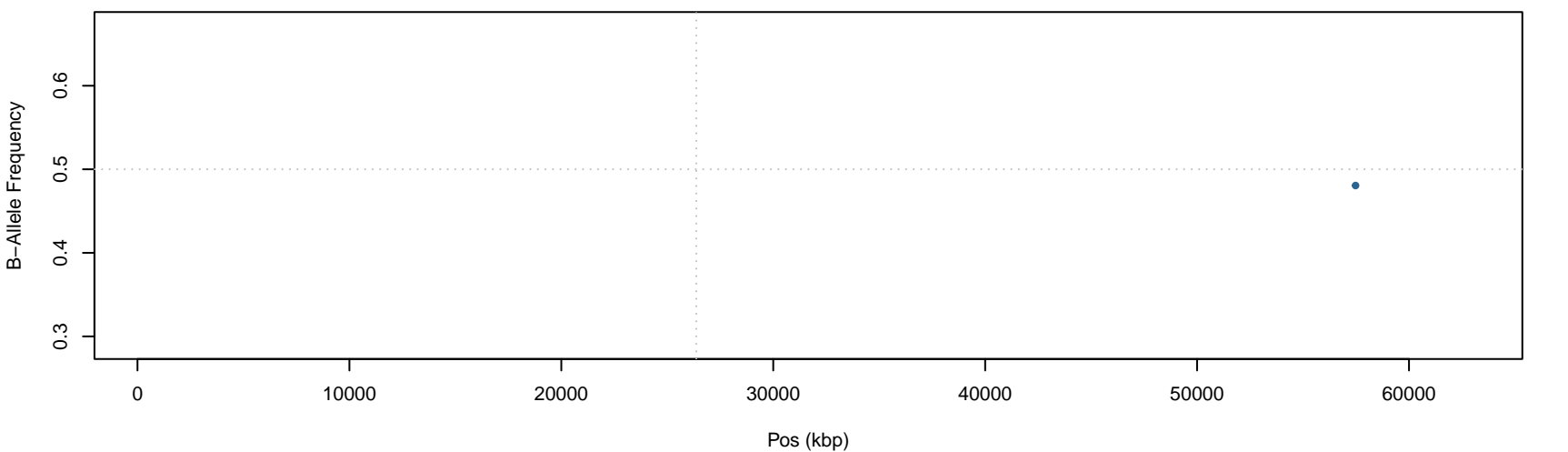
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 18



Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 19



Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 20



Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 22

