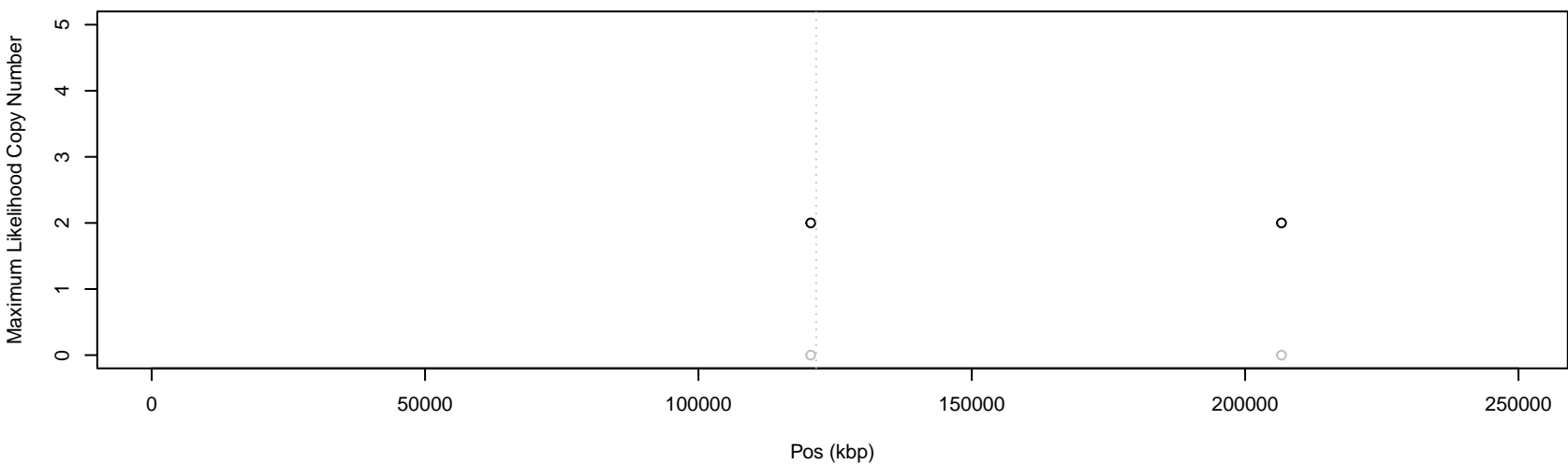
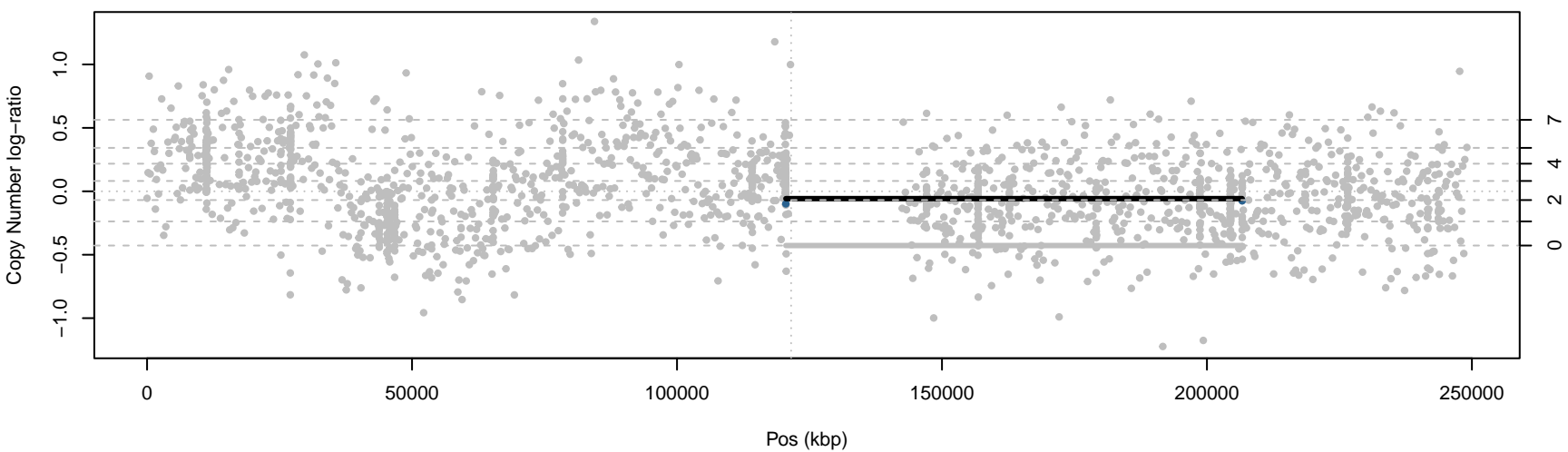
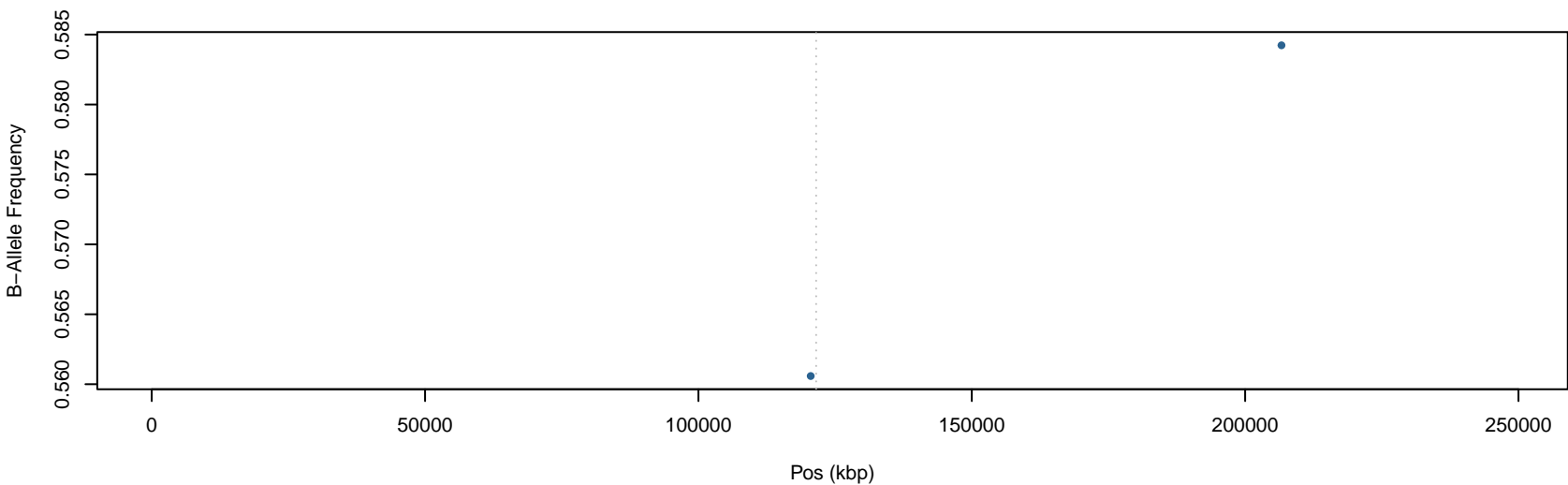
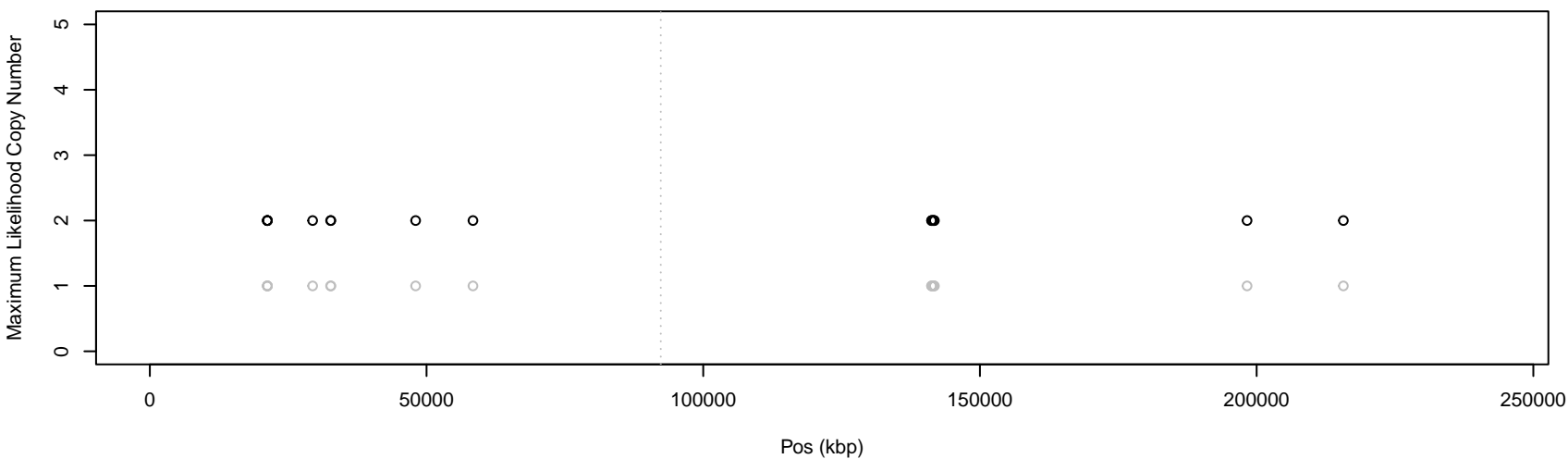
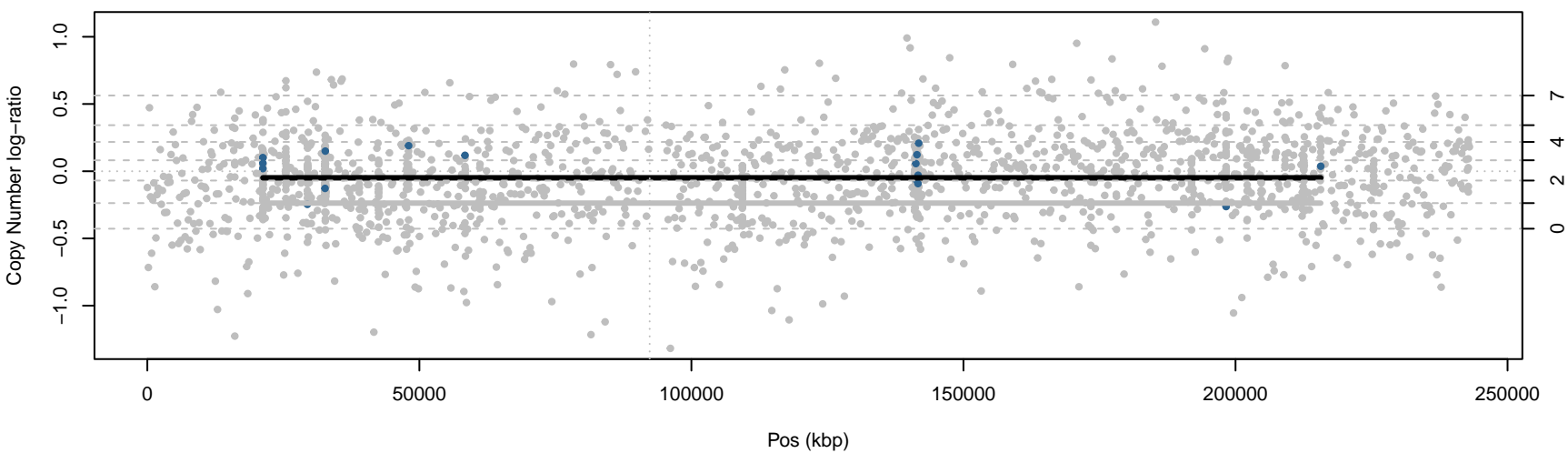
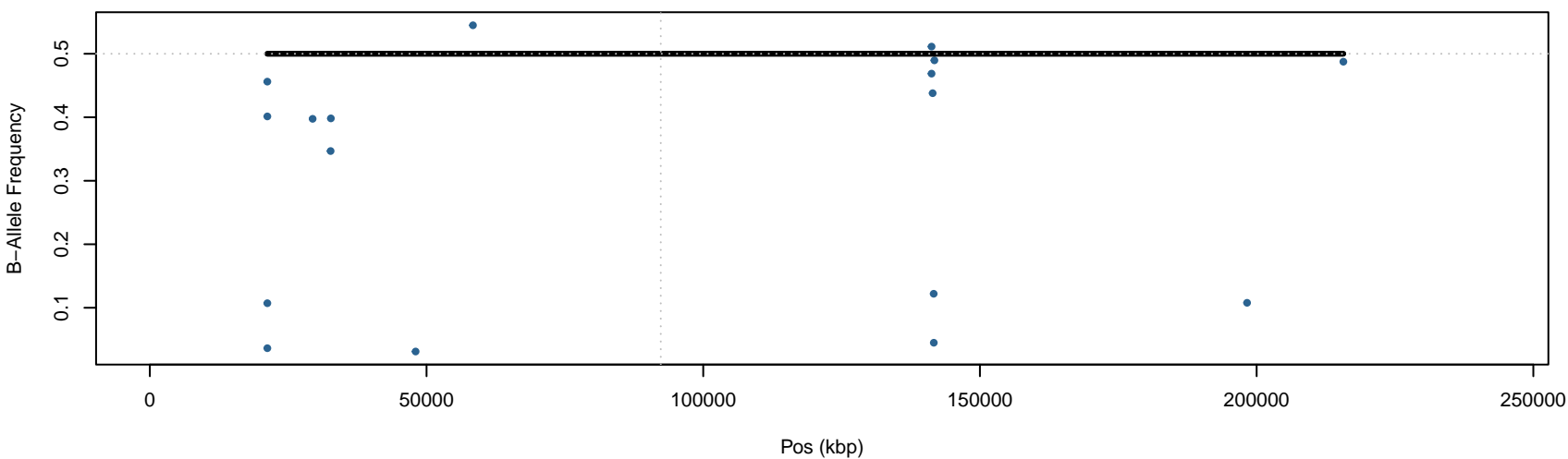


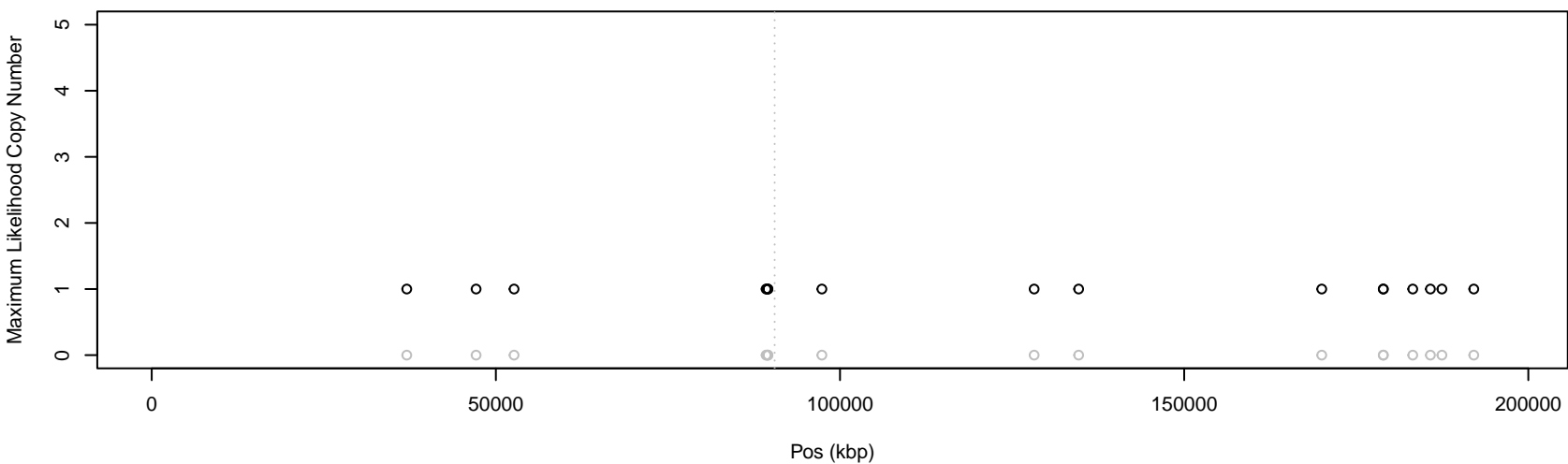
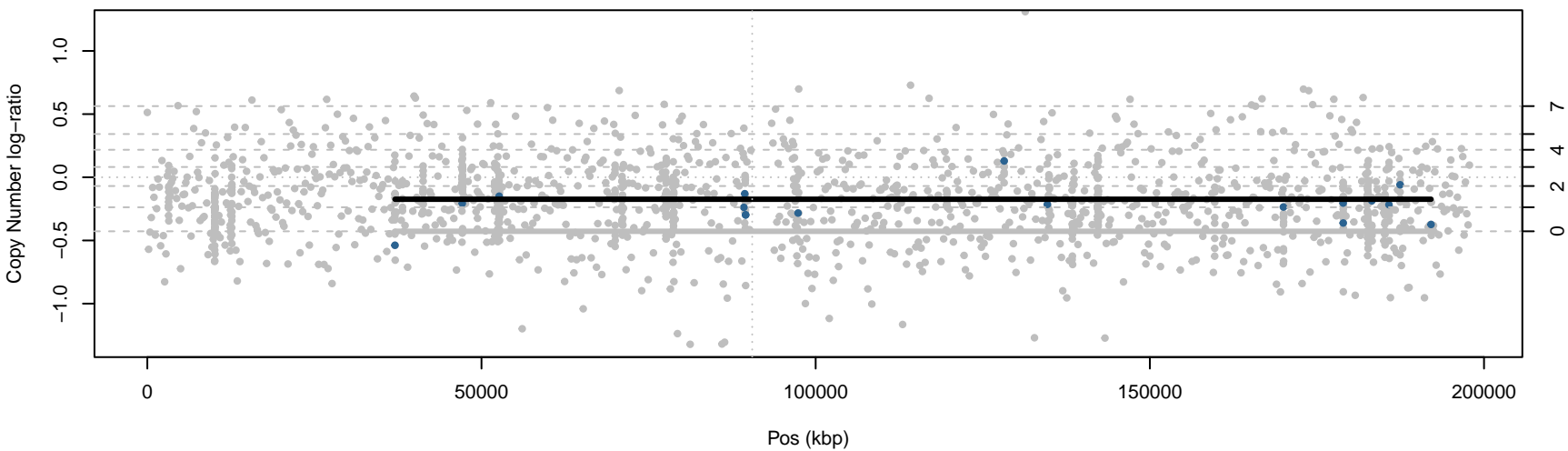
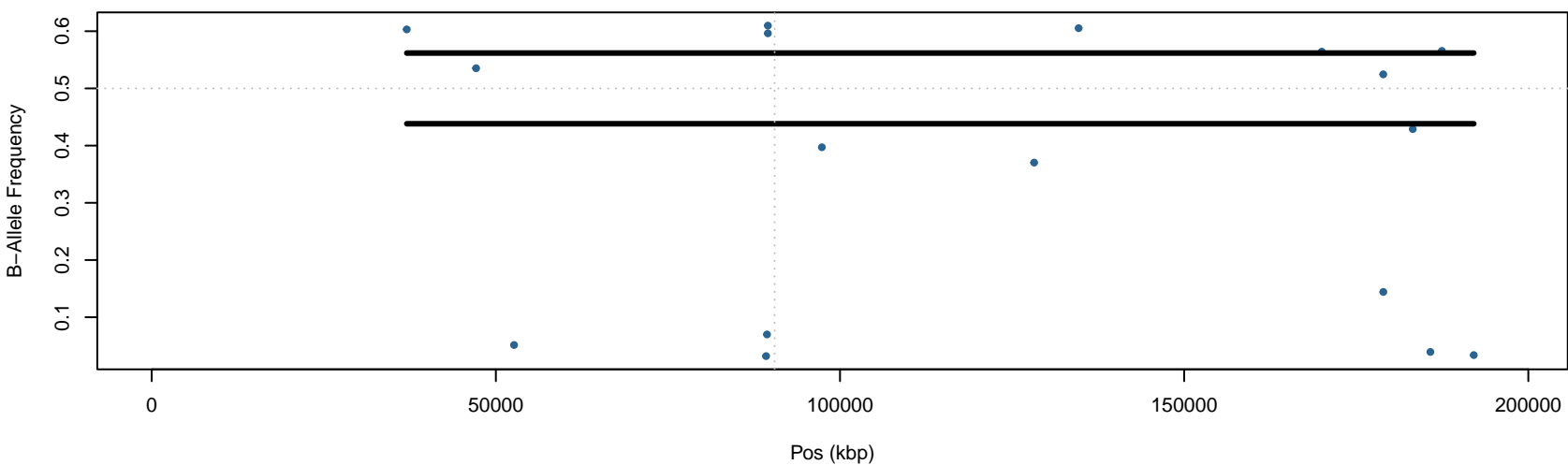
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 1



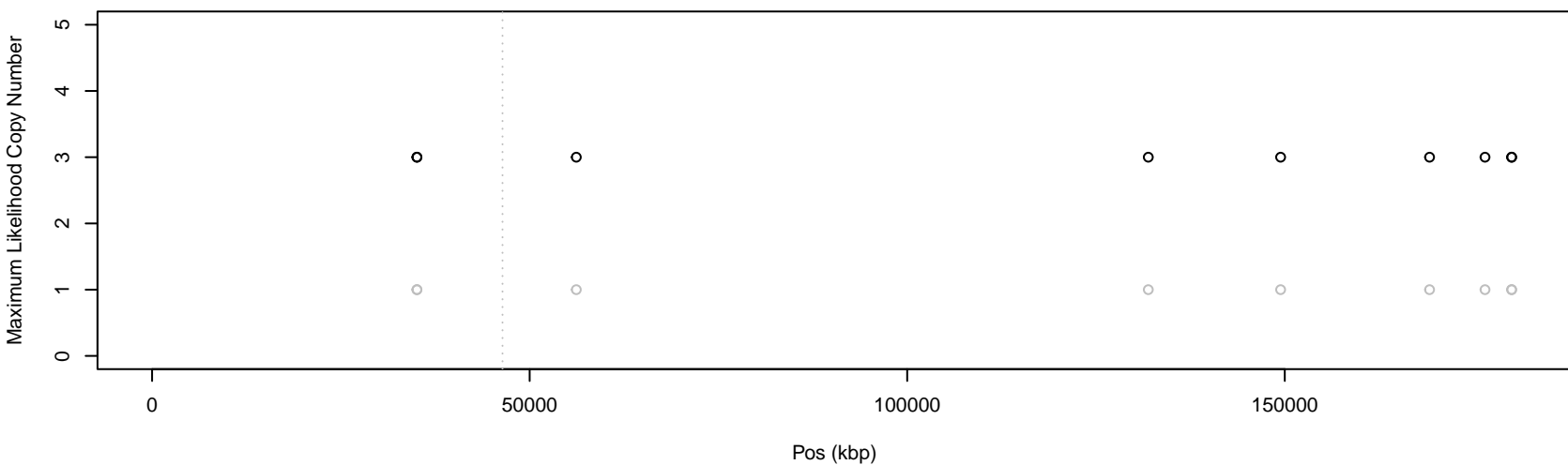
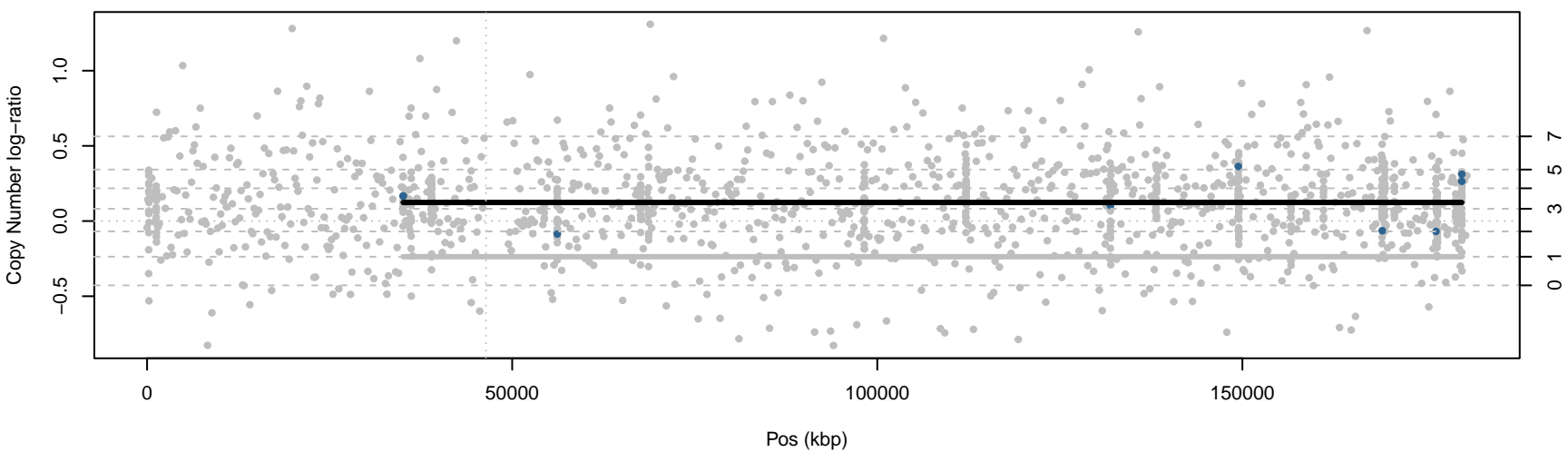
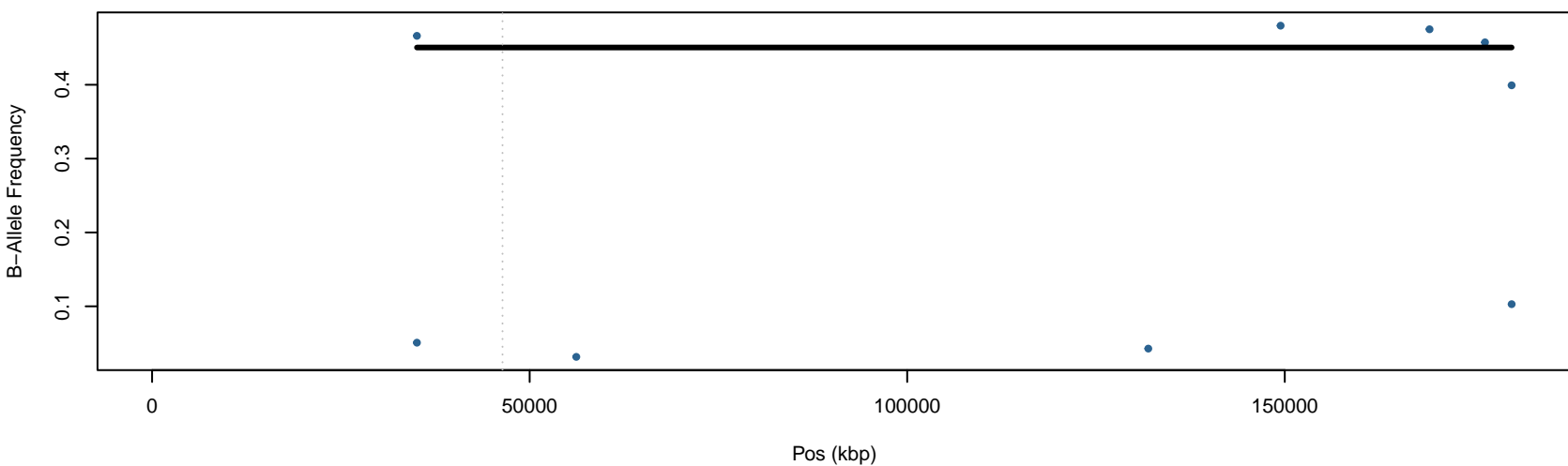
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 2



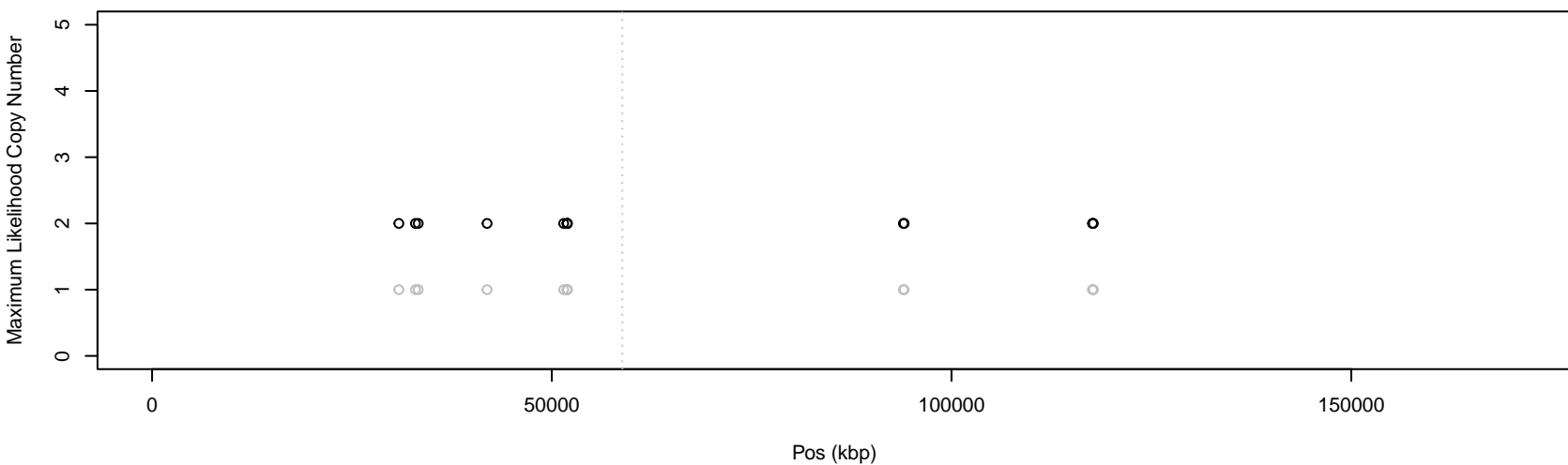
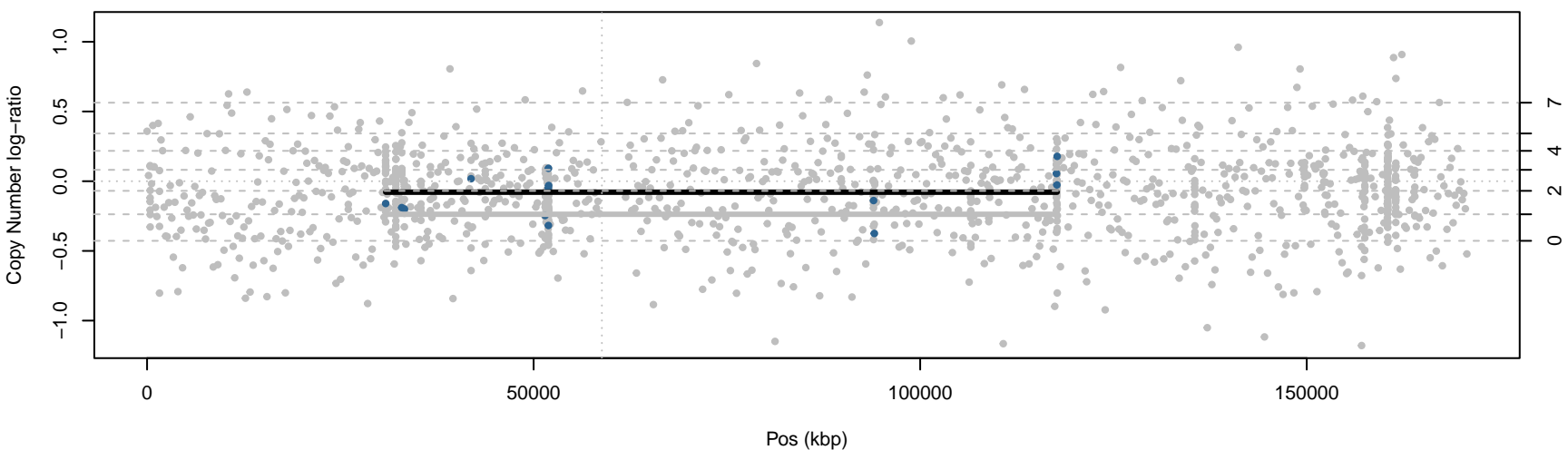
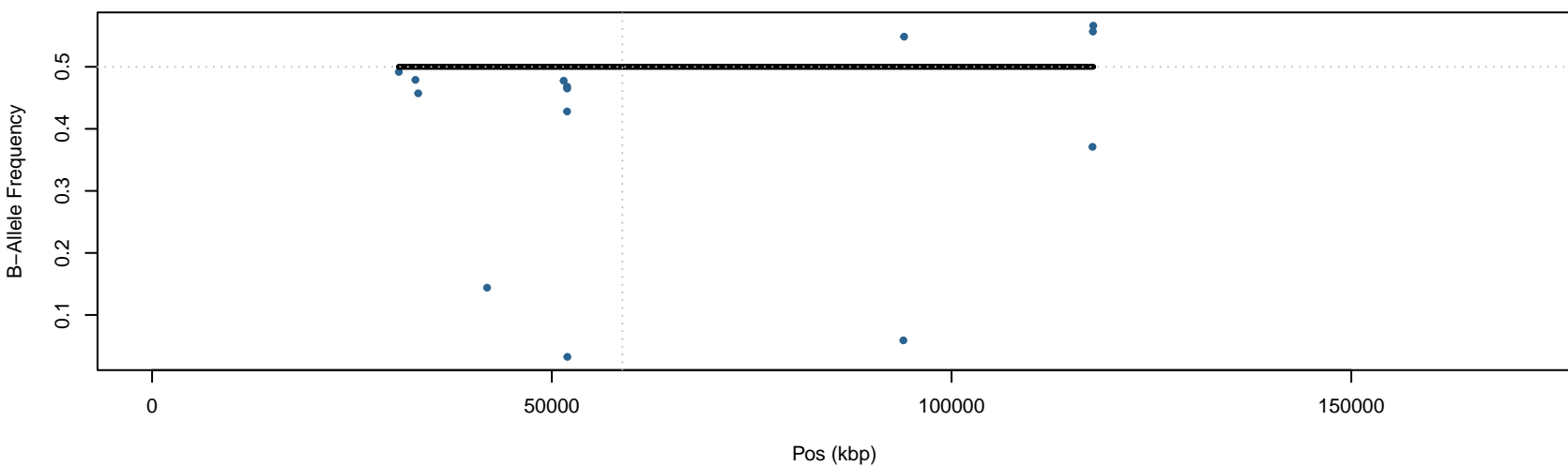
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 3



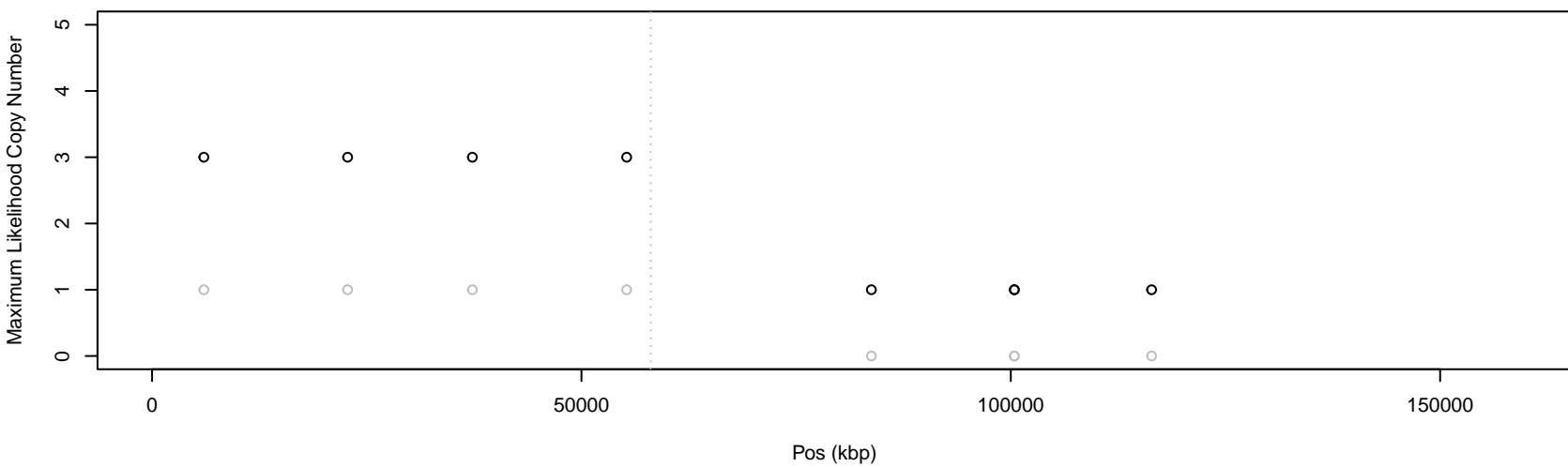
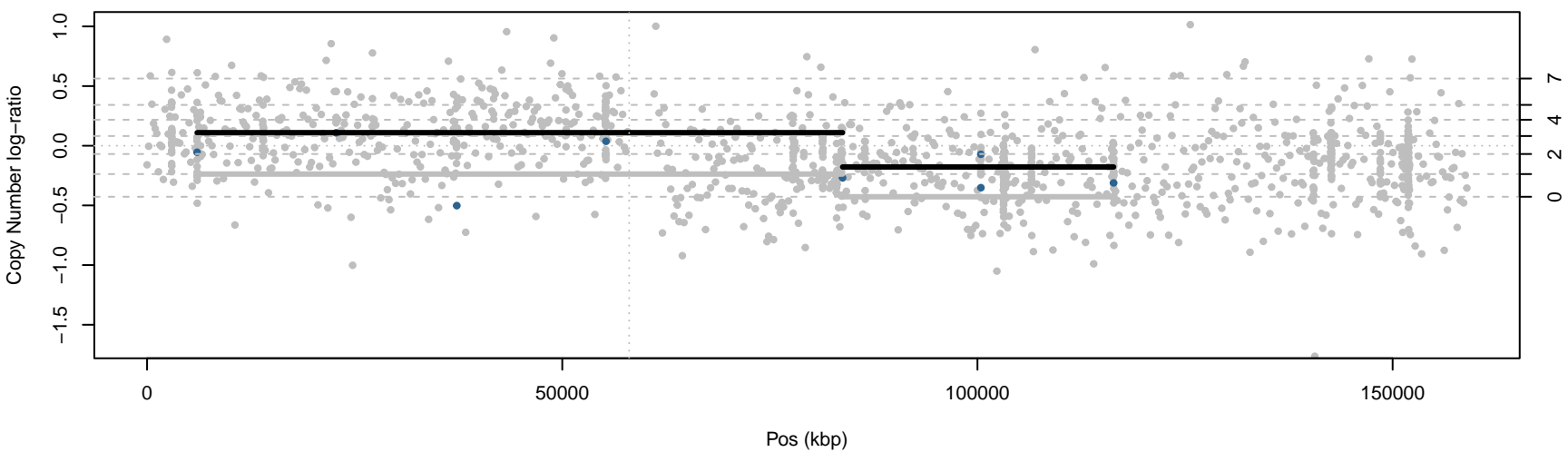
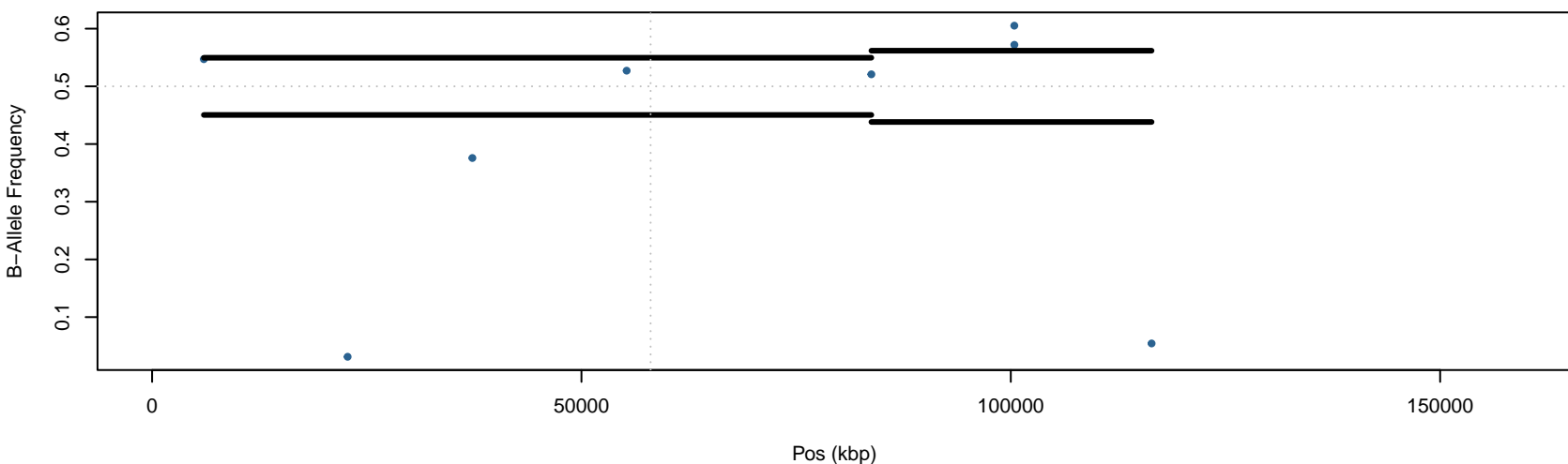
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 5



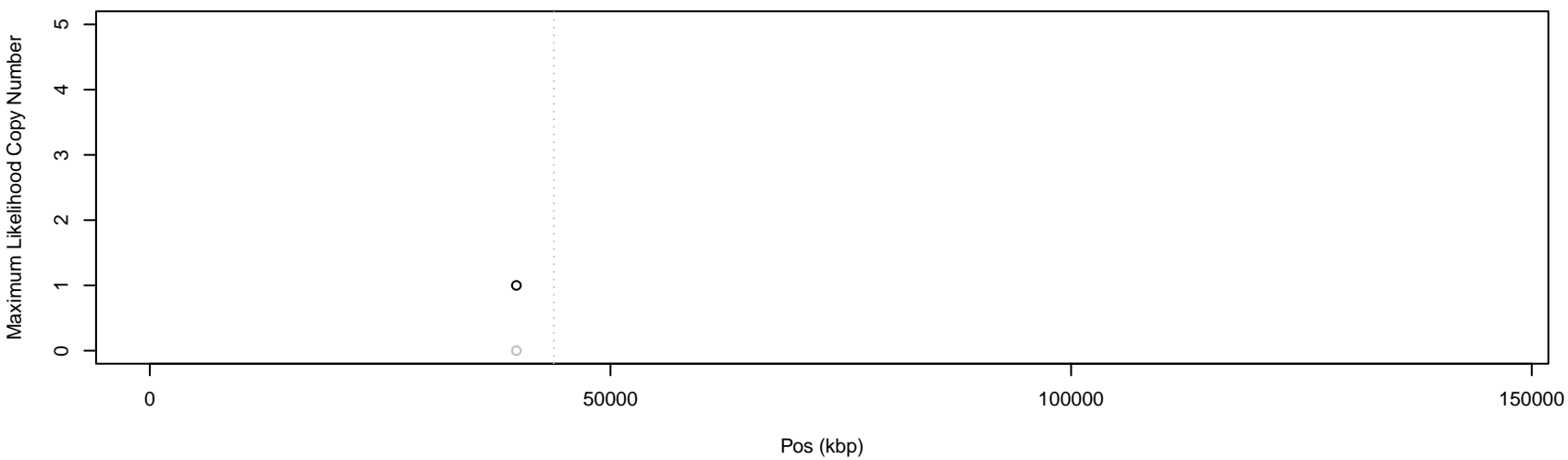
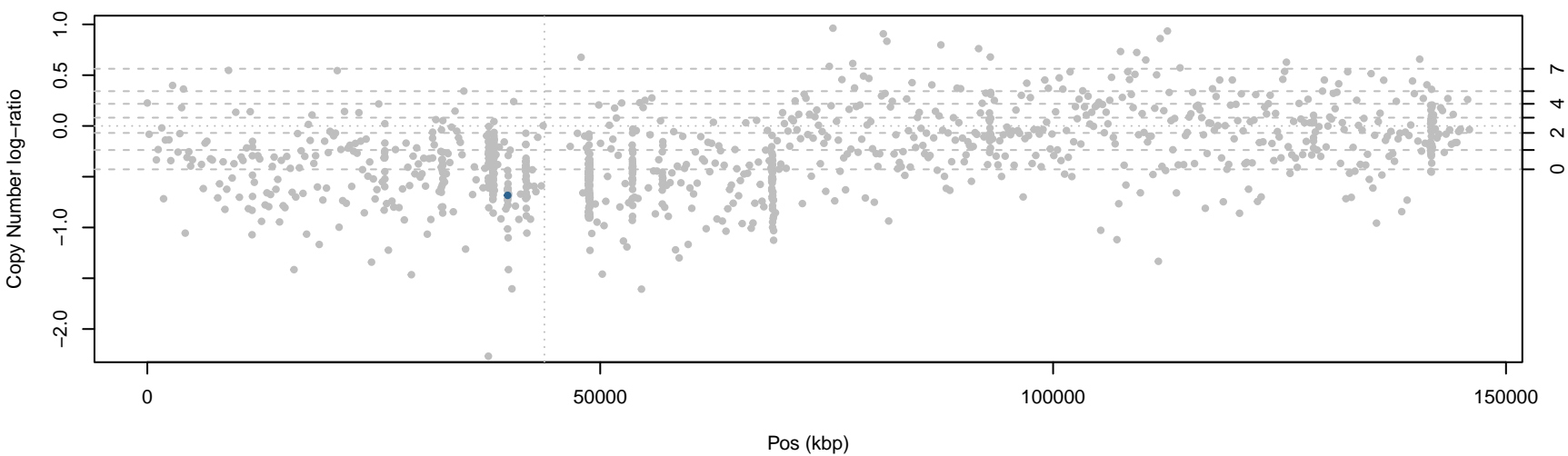
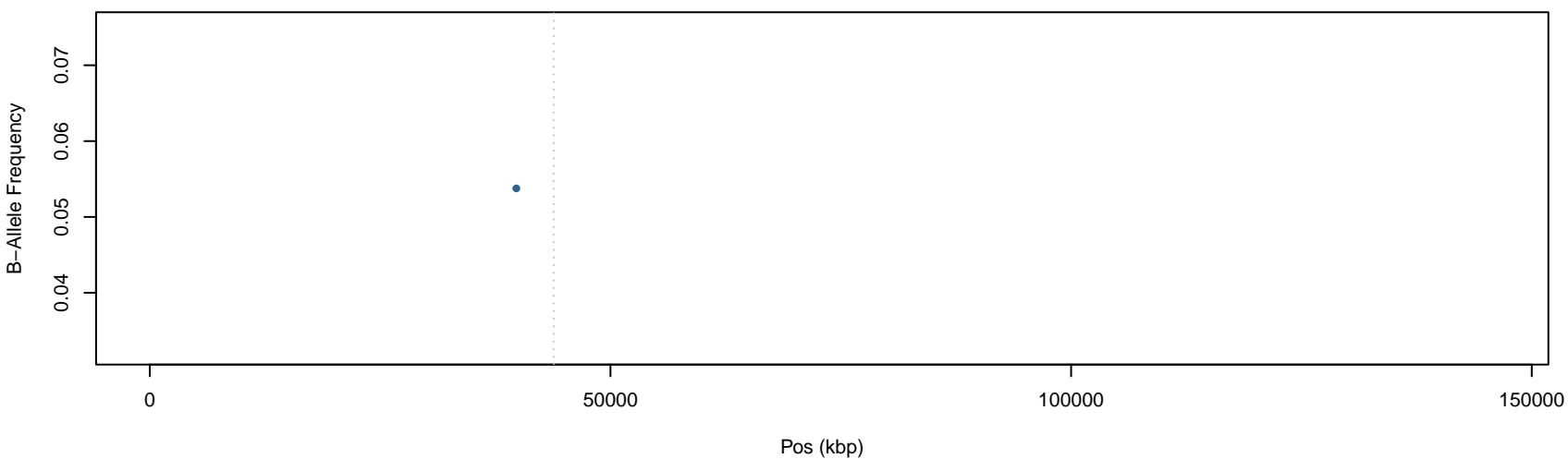
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 6



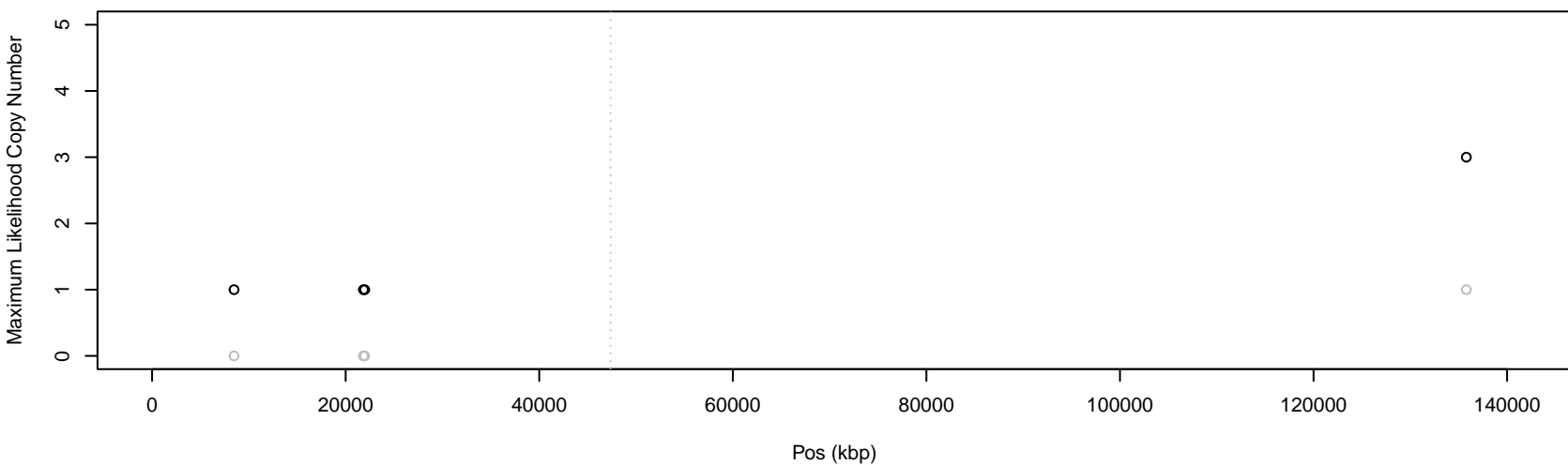
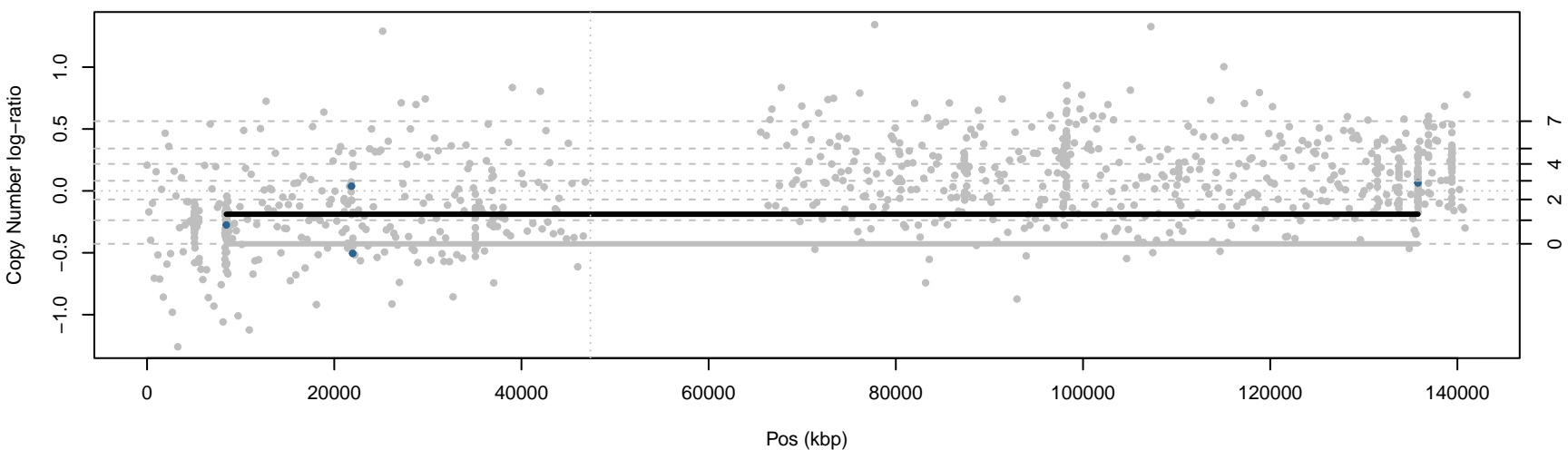
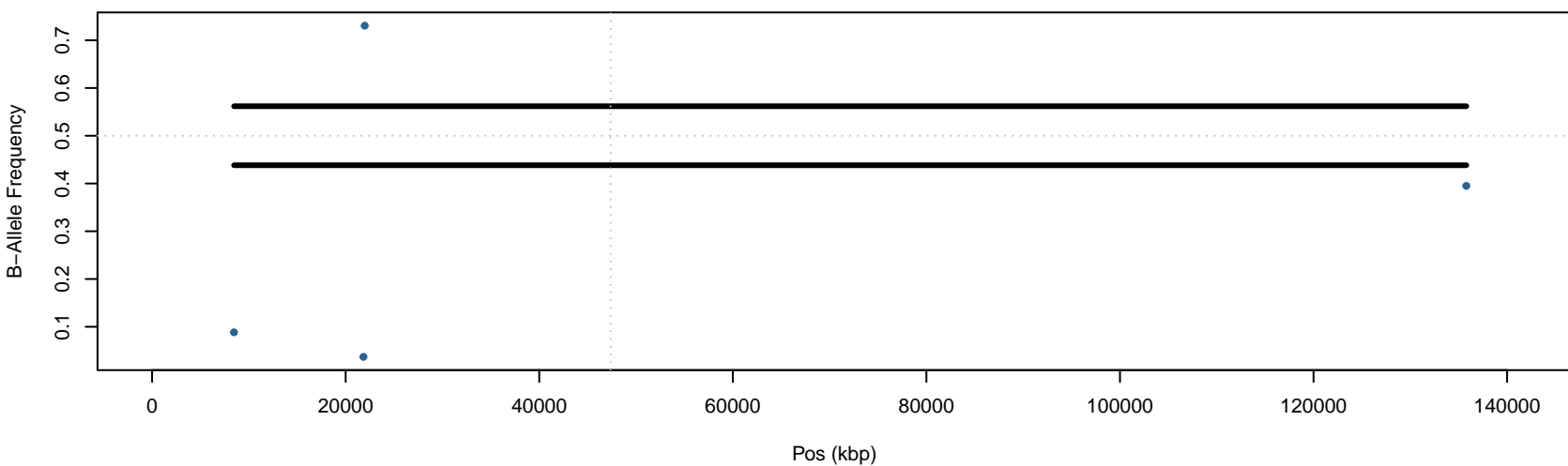
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 7



Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 8

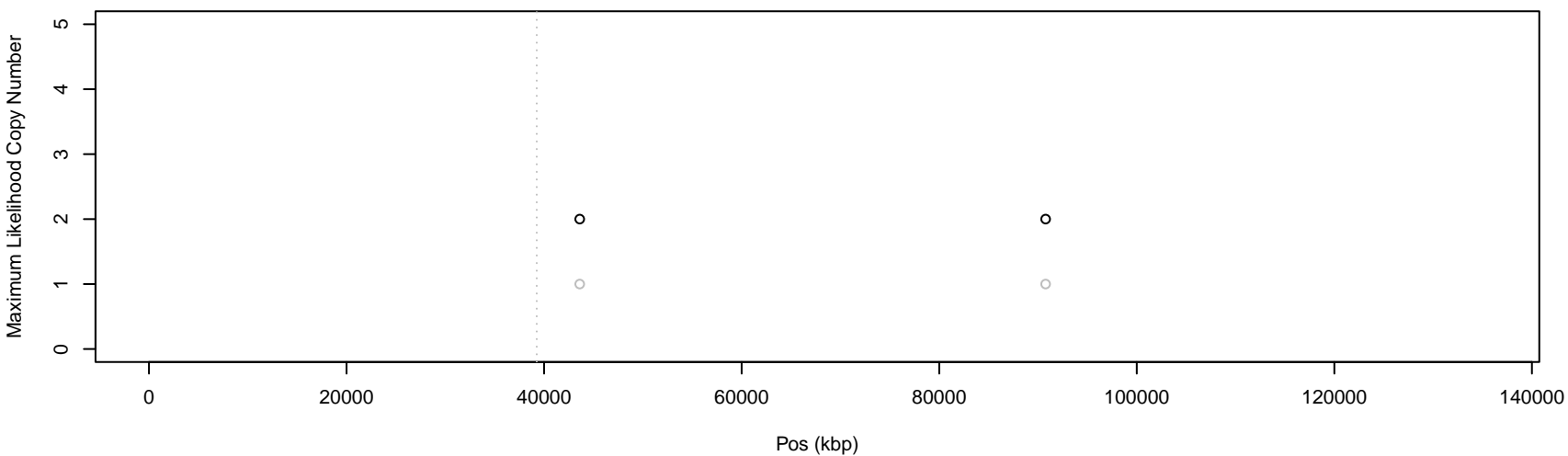
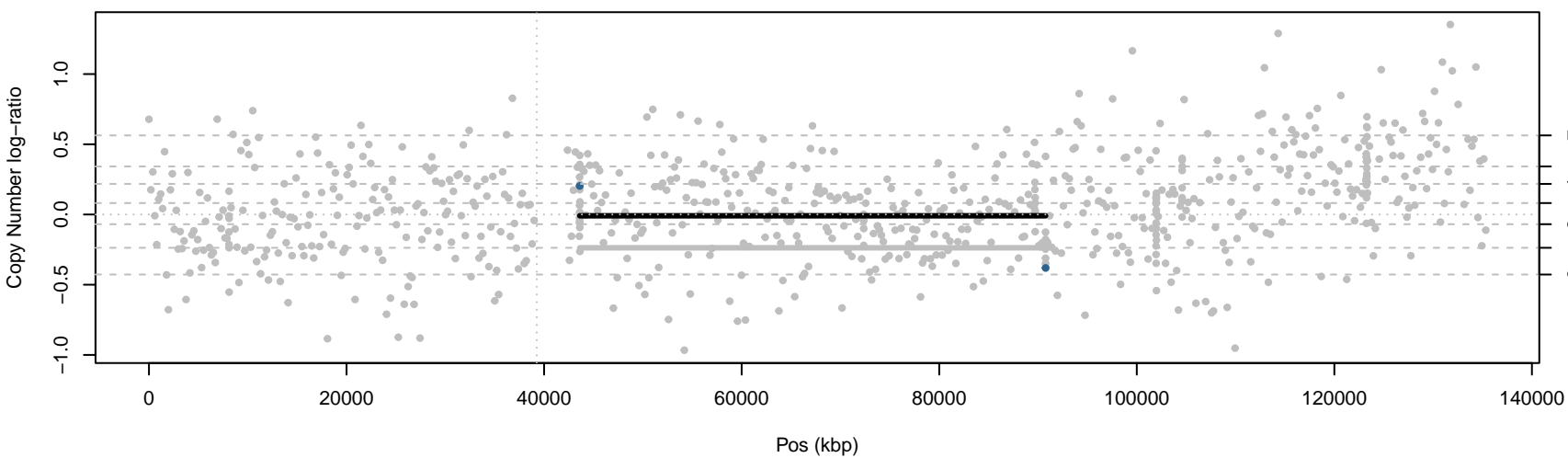
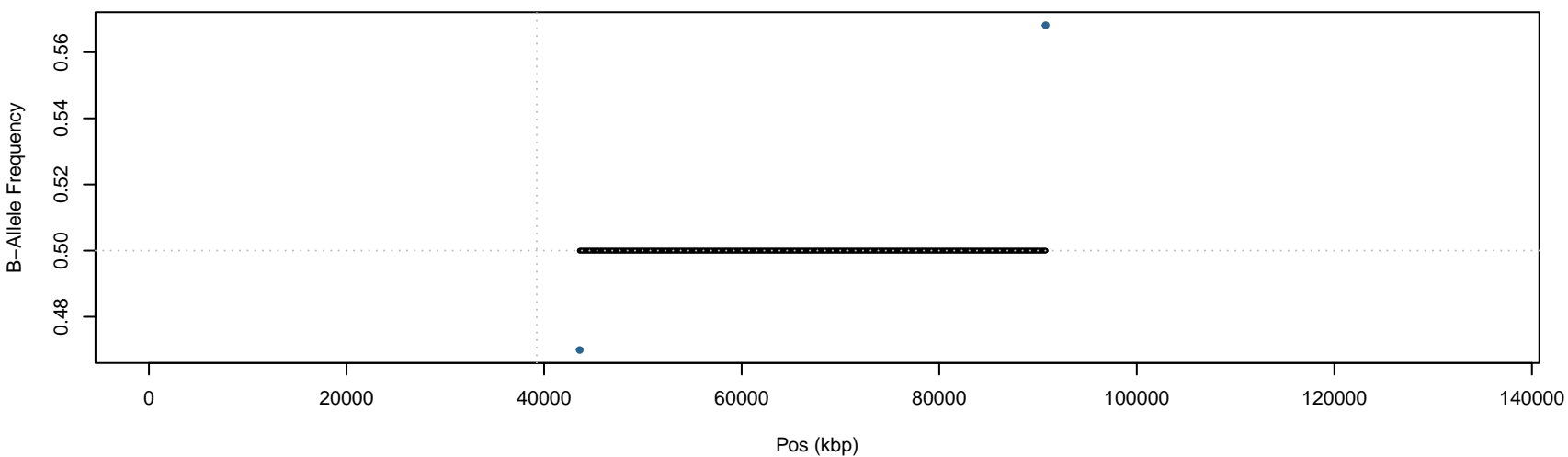


Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 9

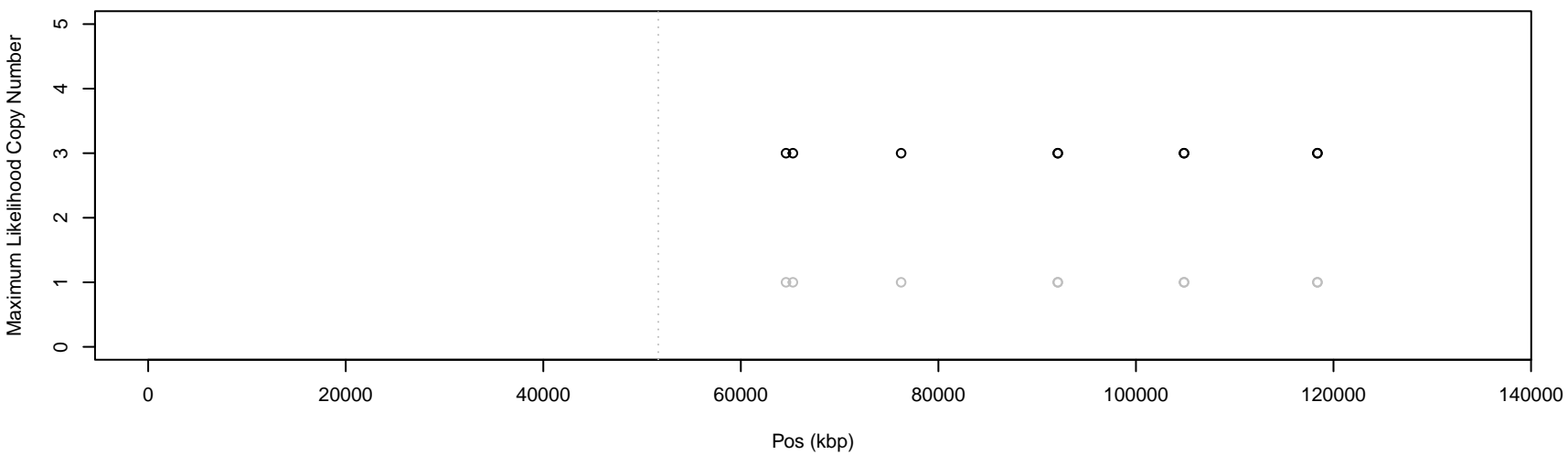
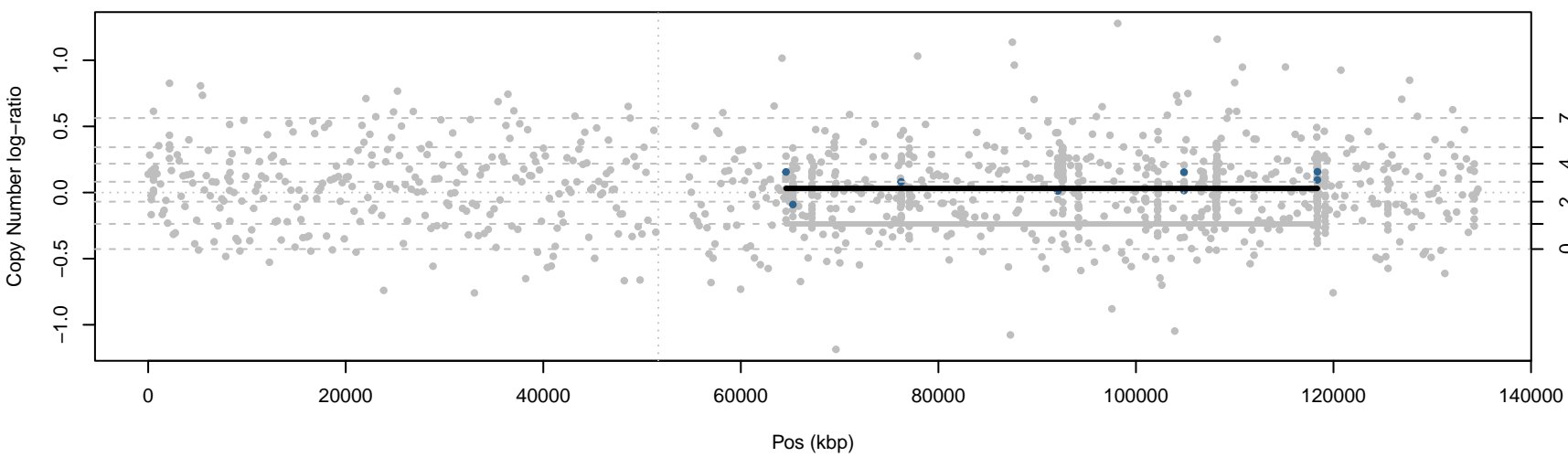
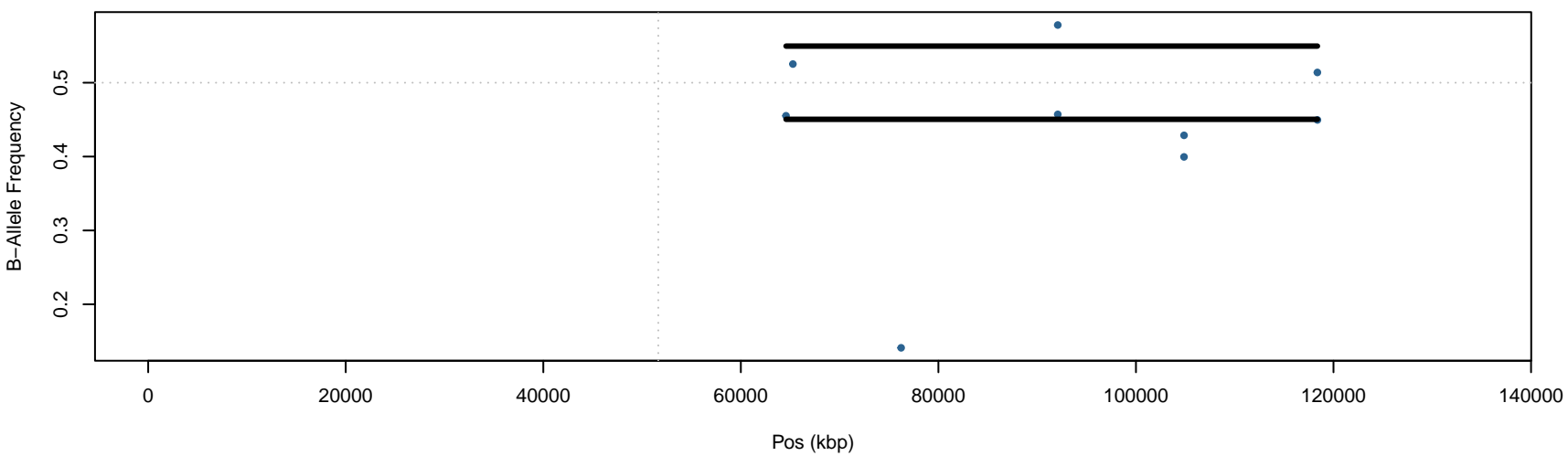




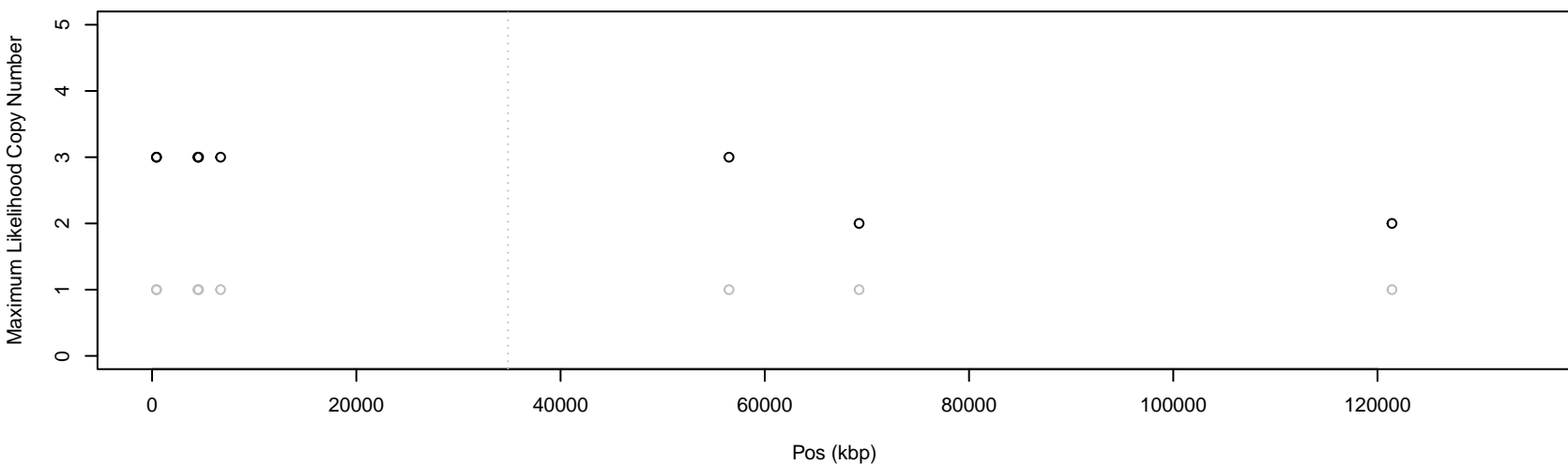
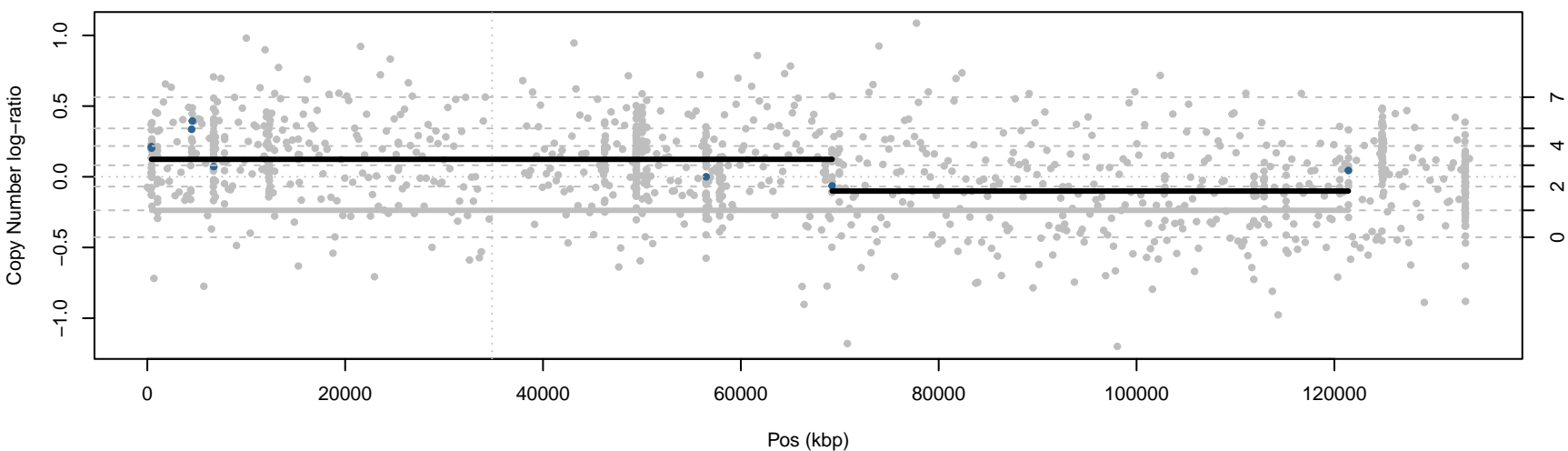
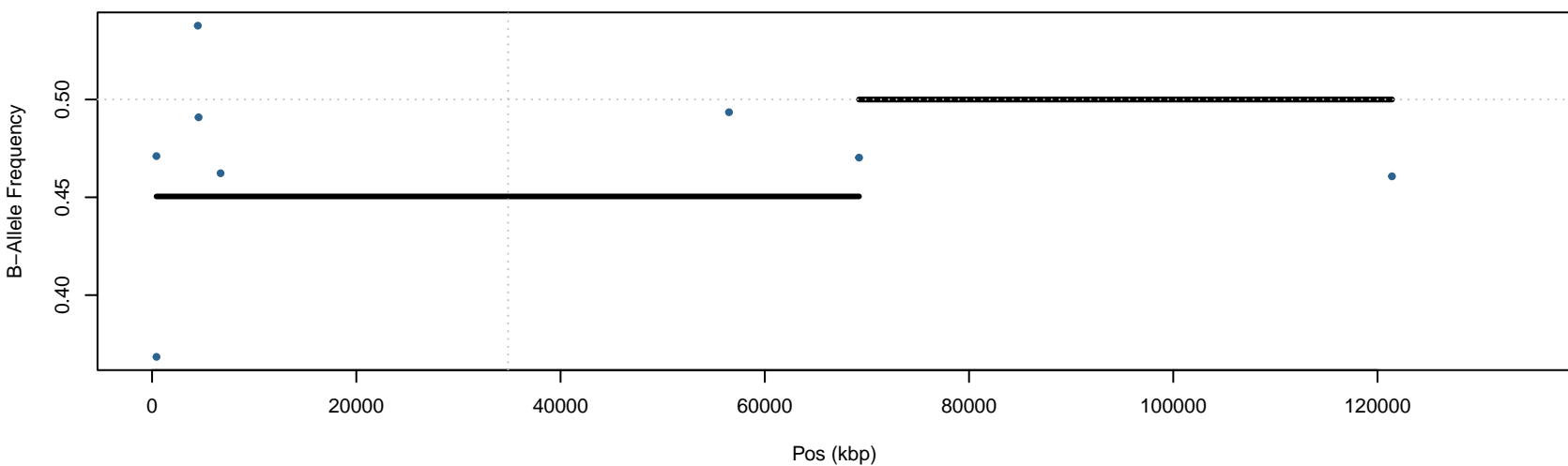
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 10



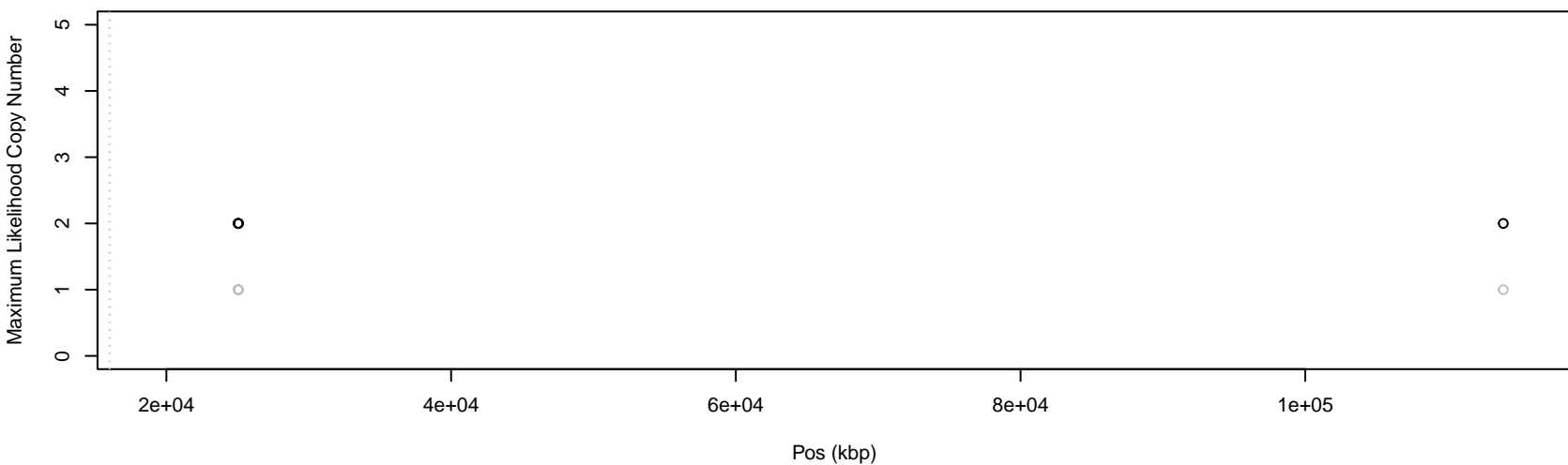
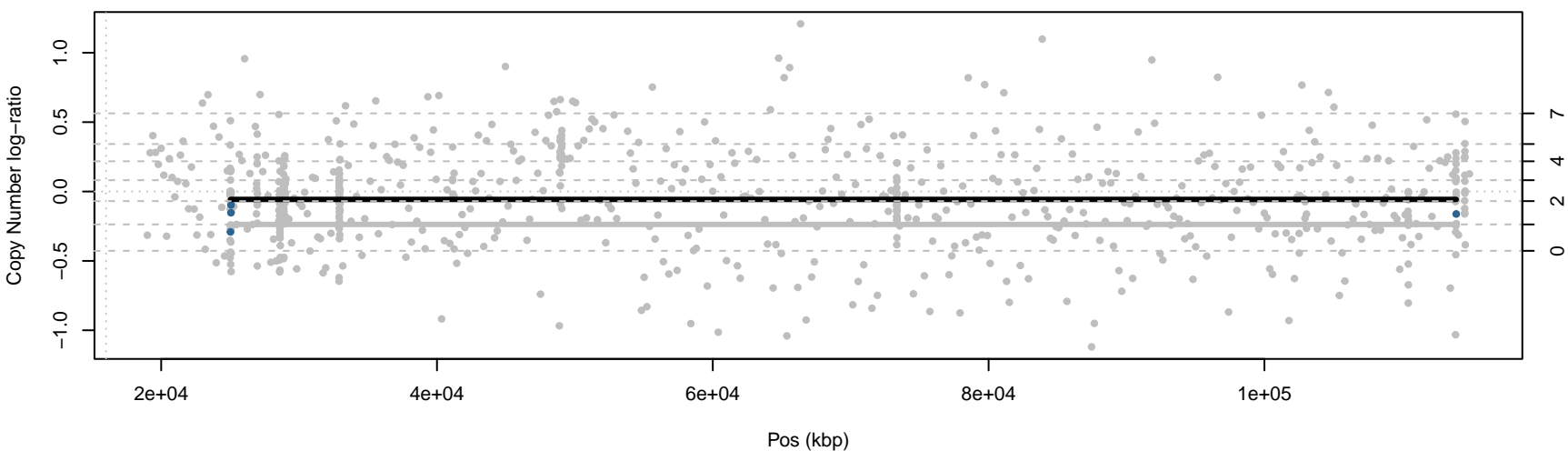
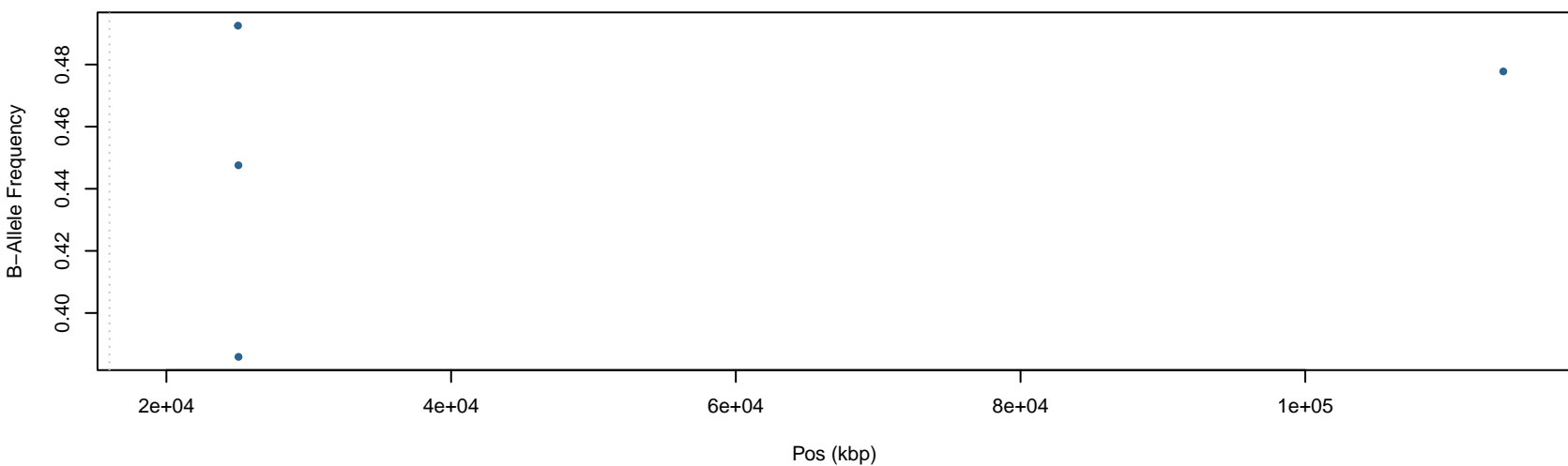
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 11



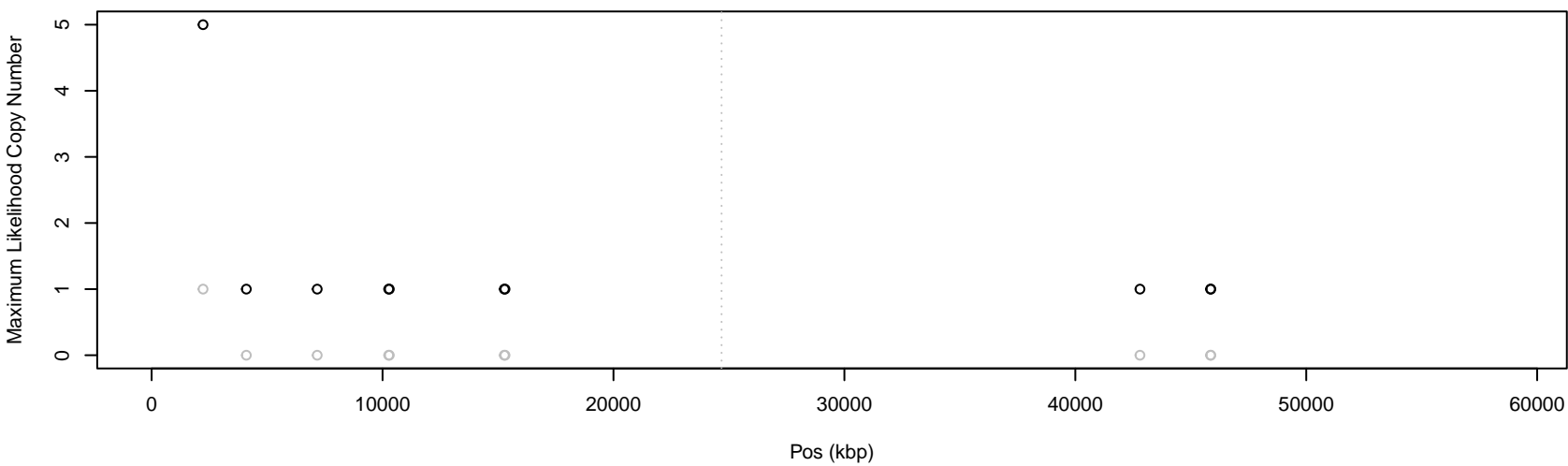
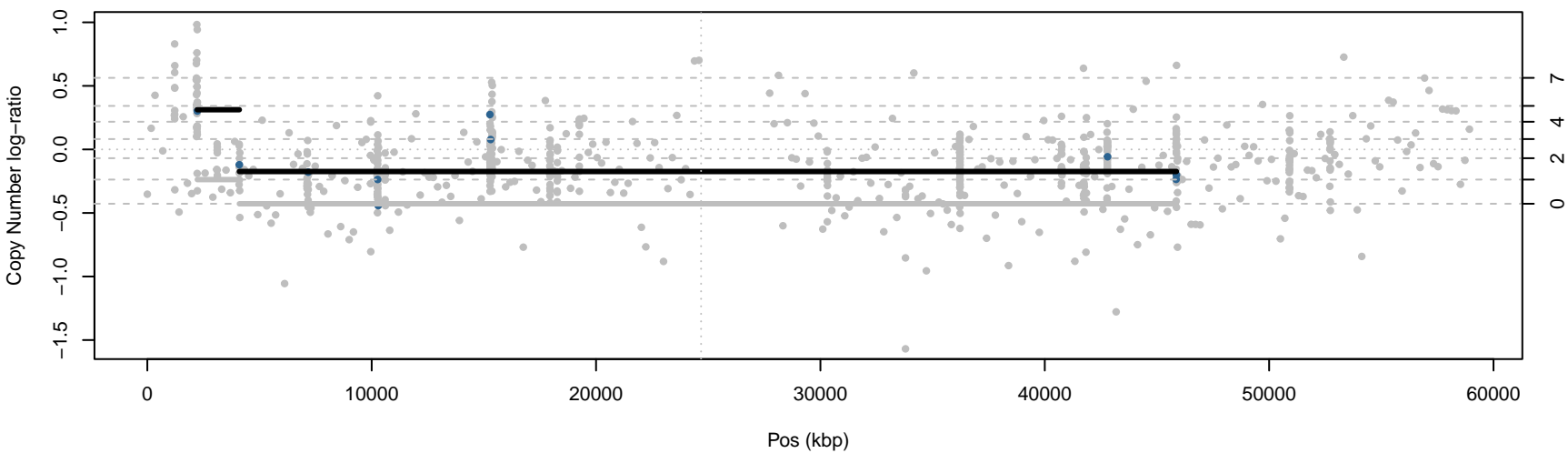
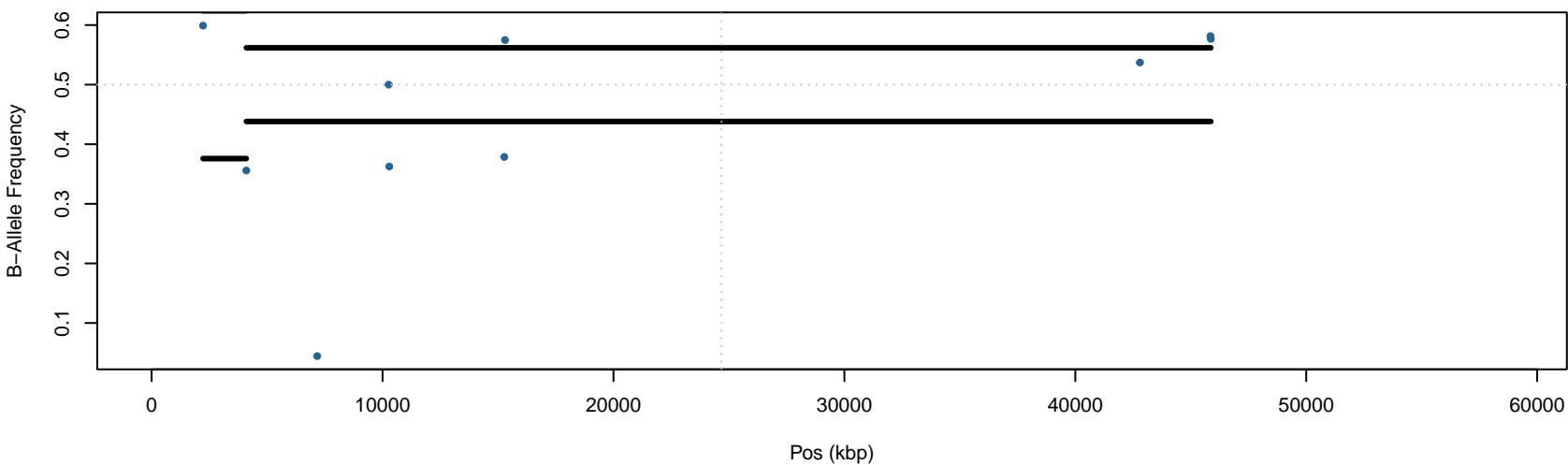
Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 12



Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 13



Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 19



Purity: 0.22 Tumor ploidy: 2.446 SNV log-likelihood: 2.98 GoF: 92.7% Mean coverage: 466;556 Chromosome: 20

