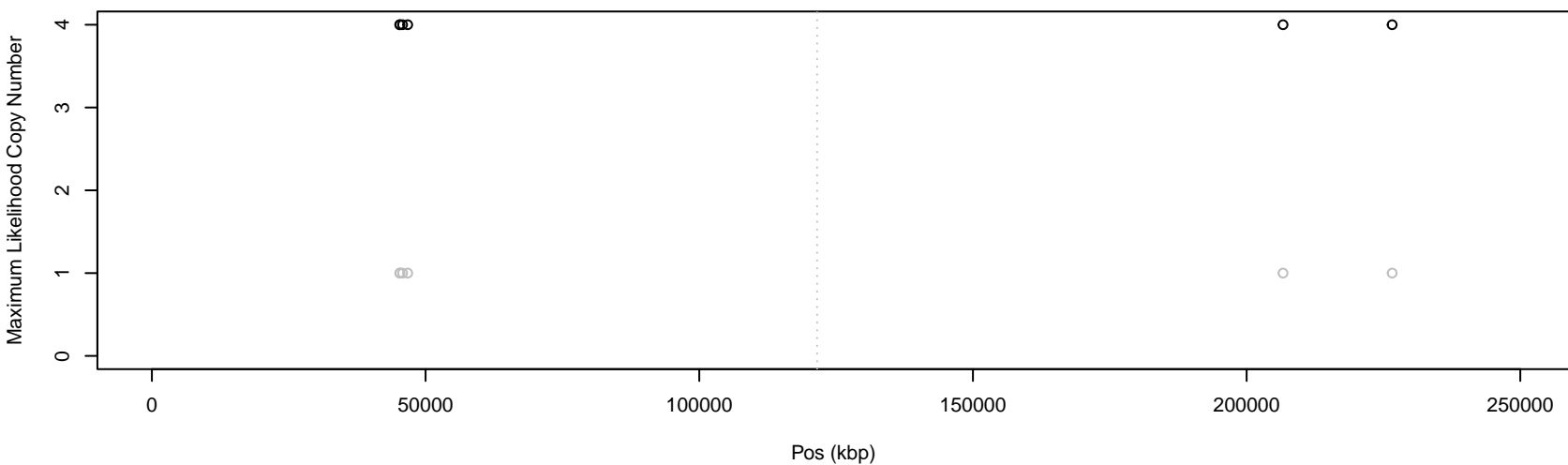
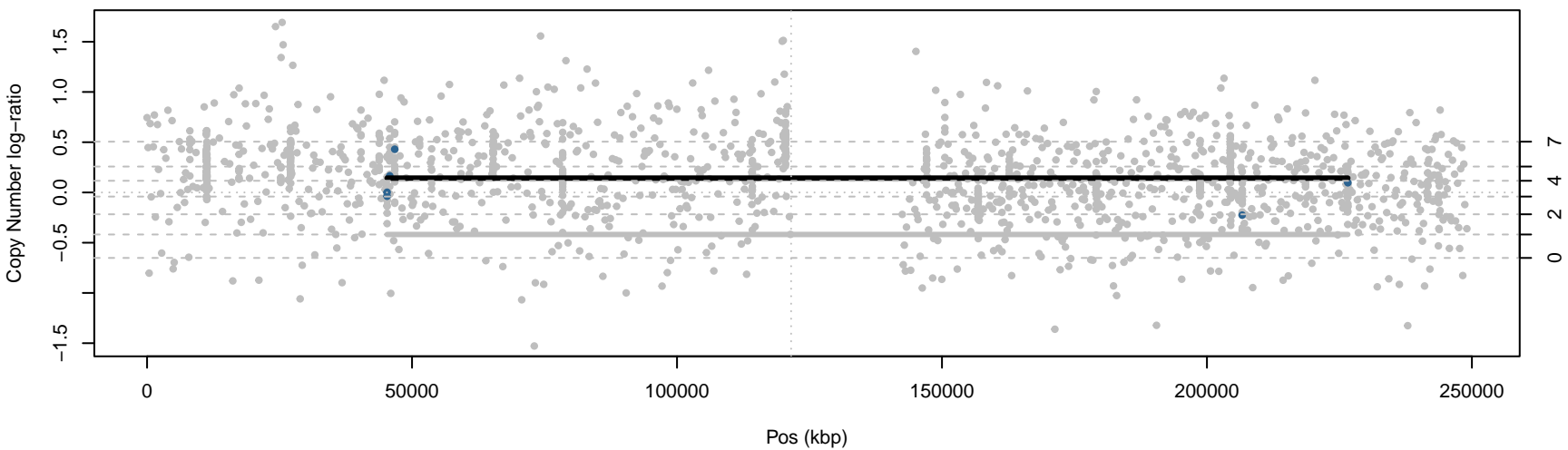
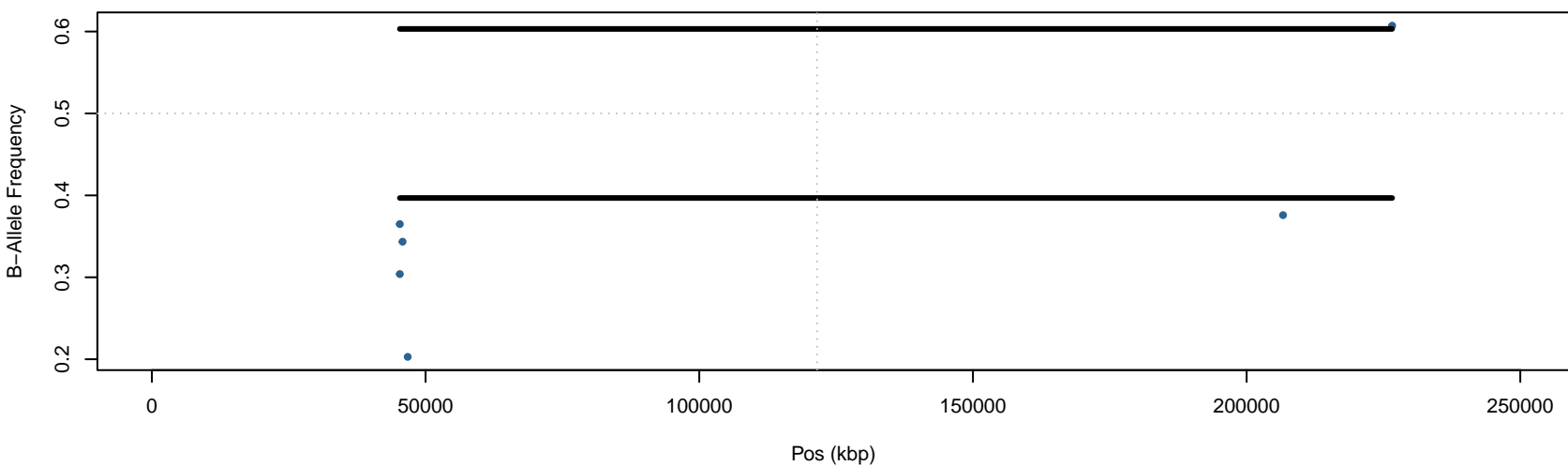
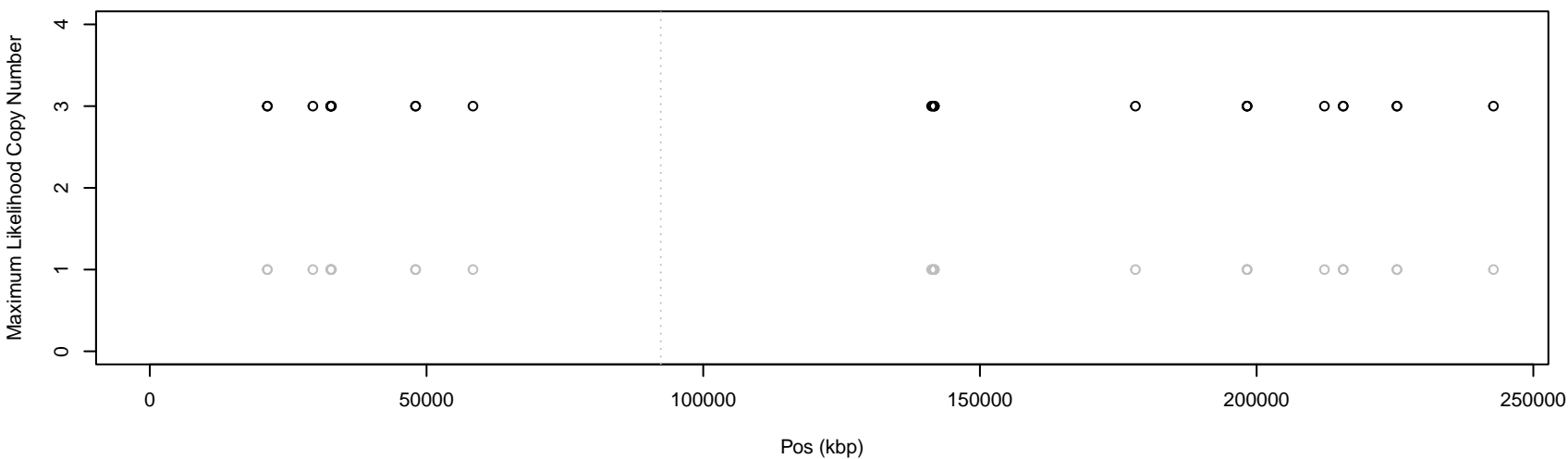
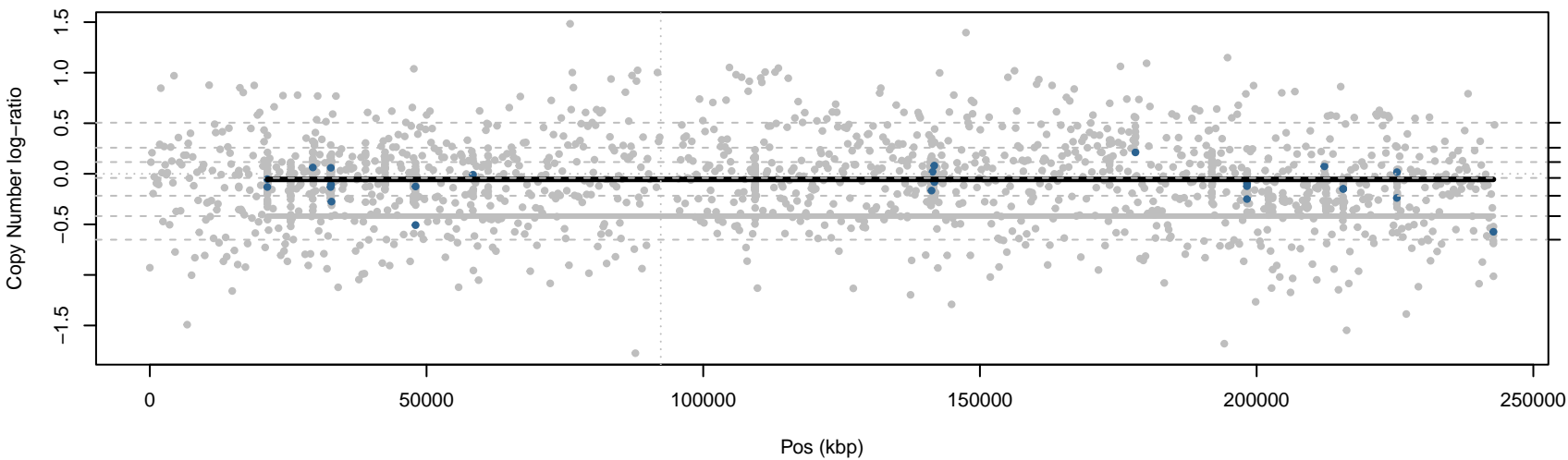
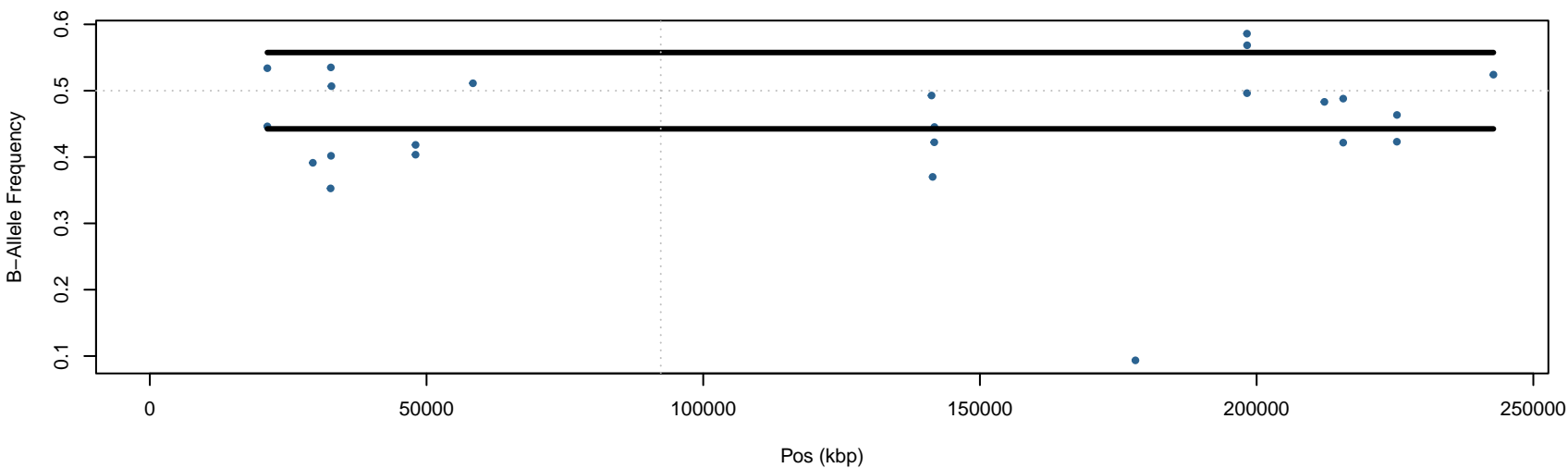


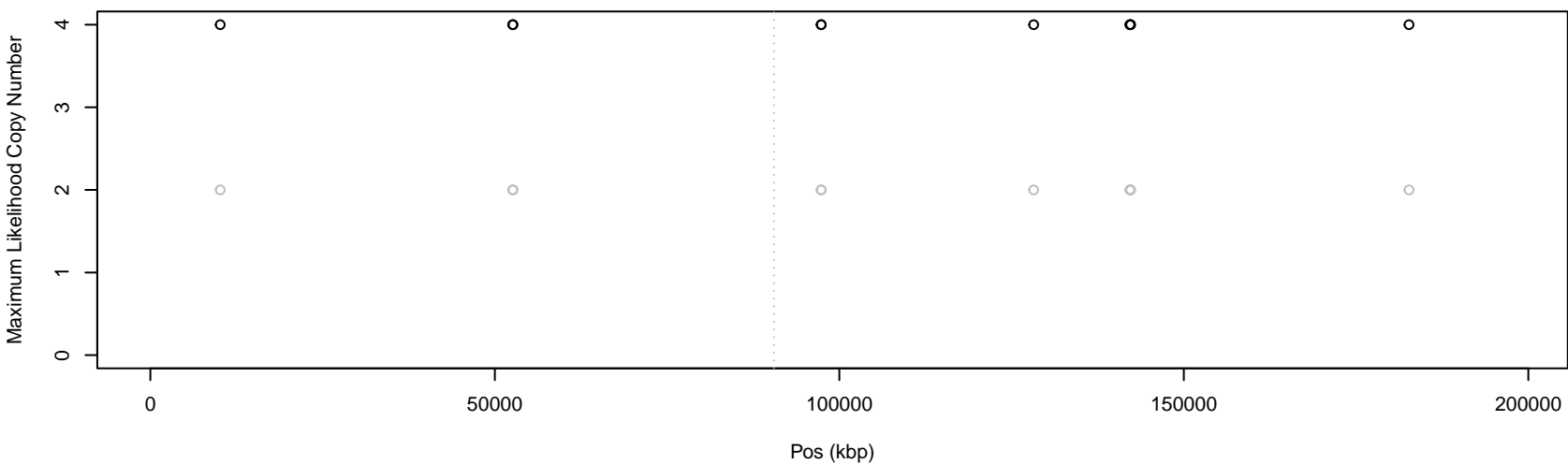
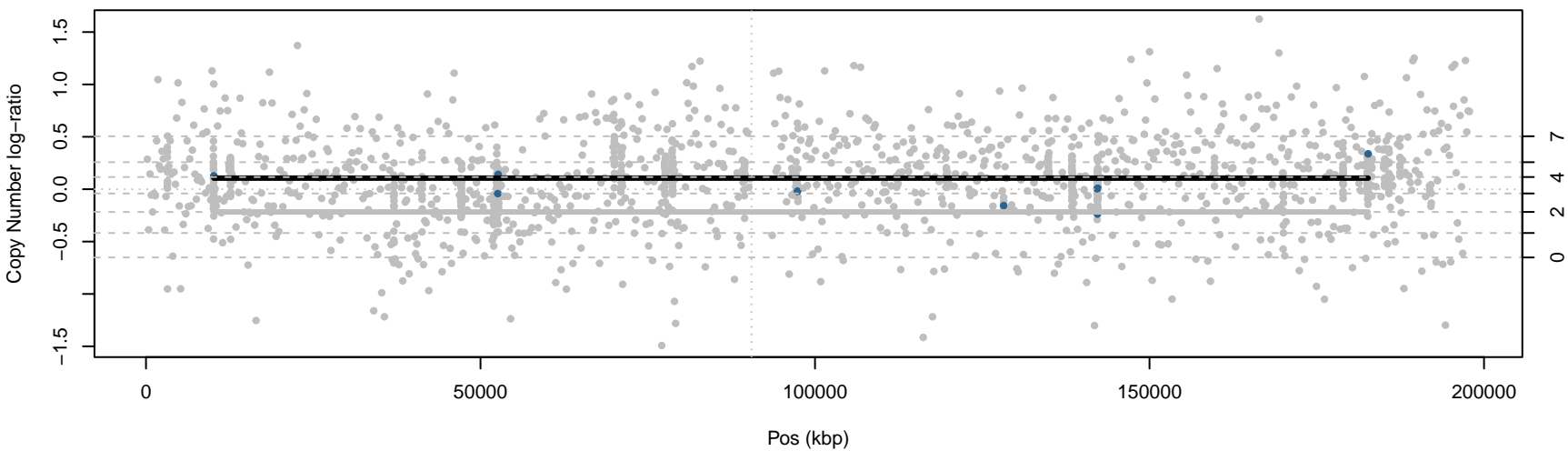
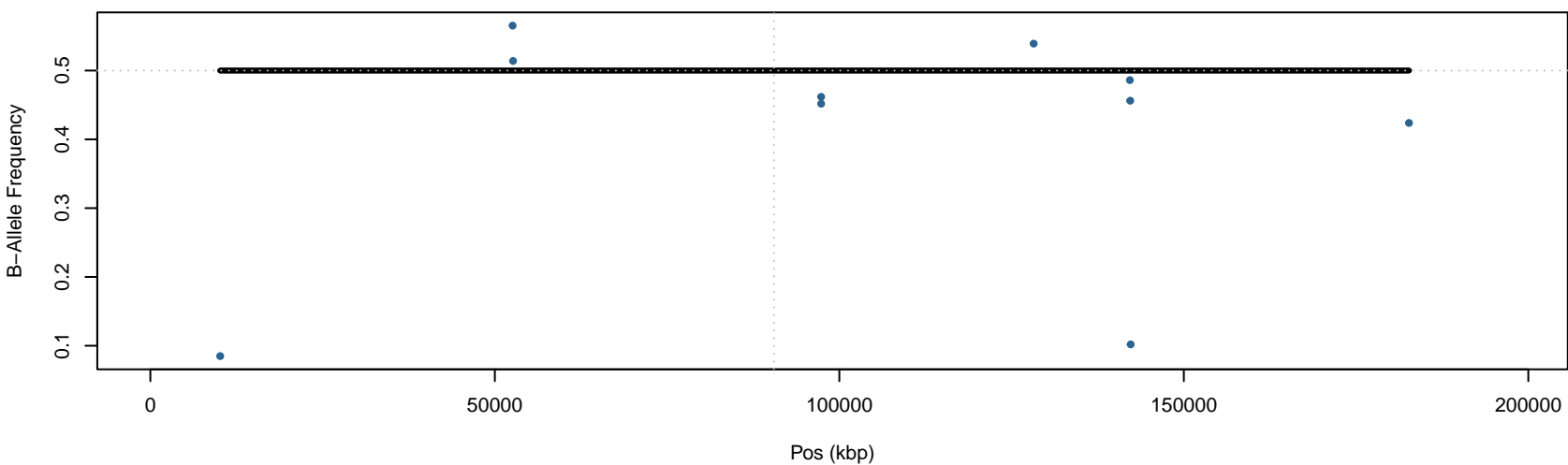
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 1



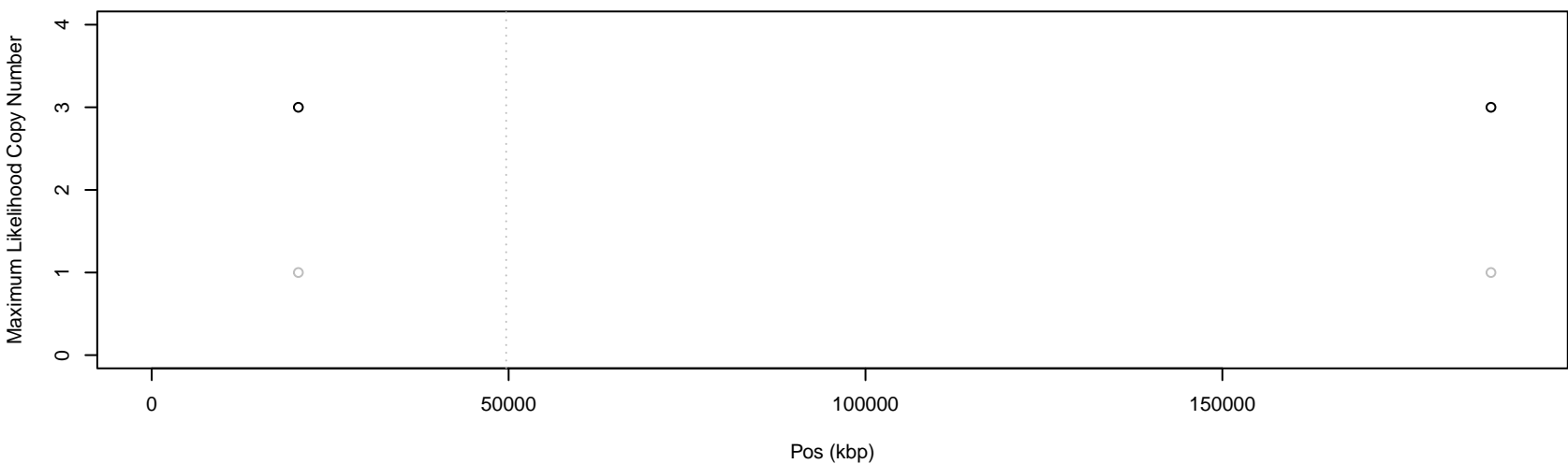
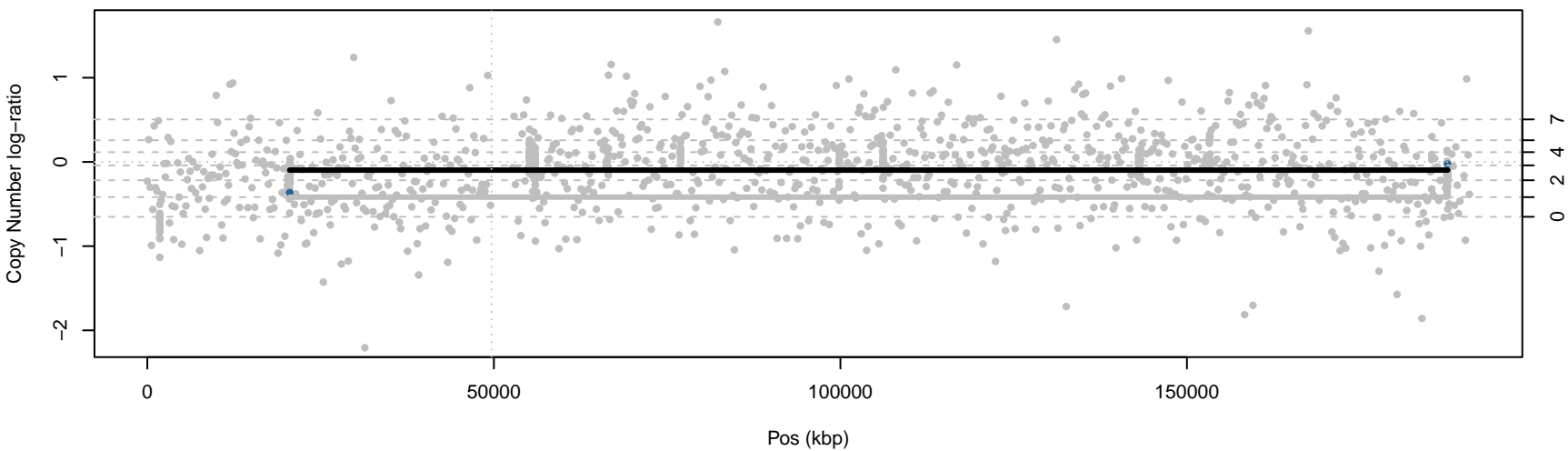
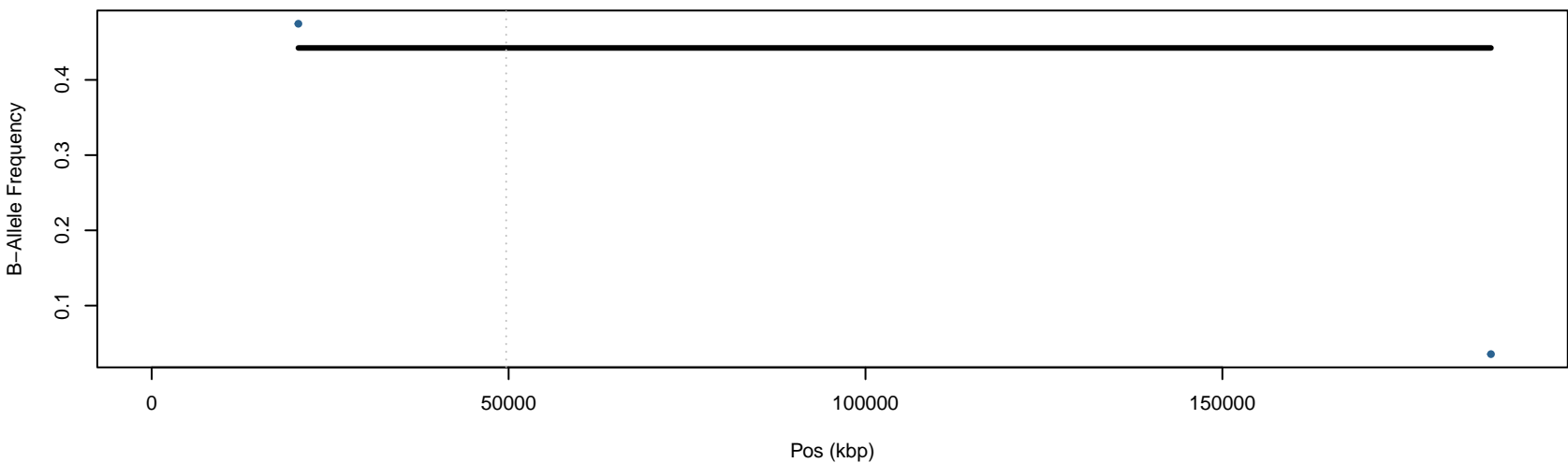
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 2



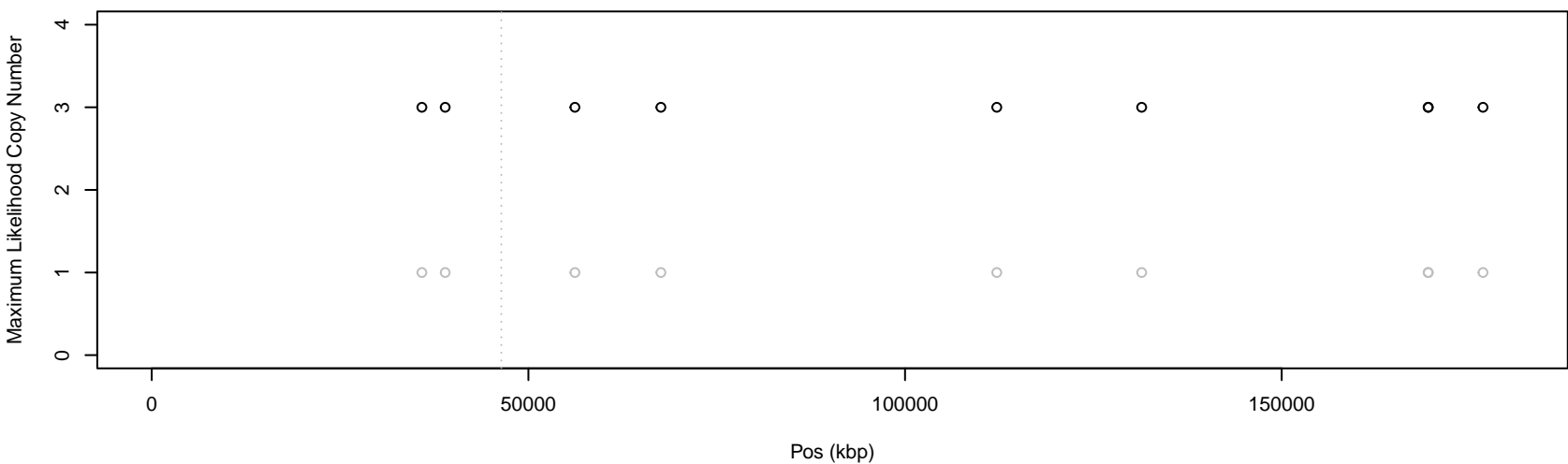
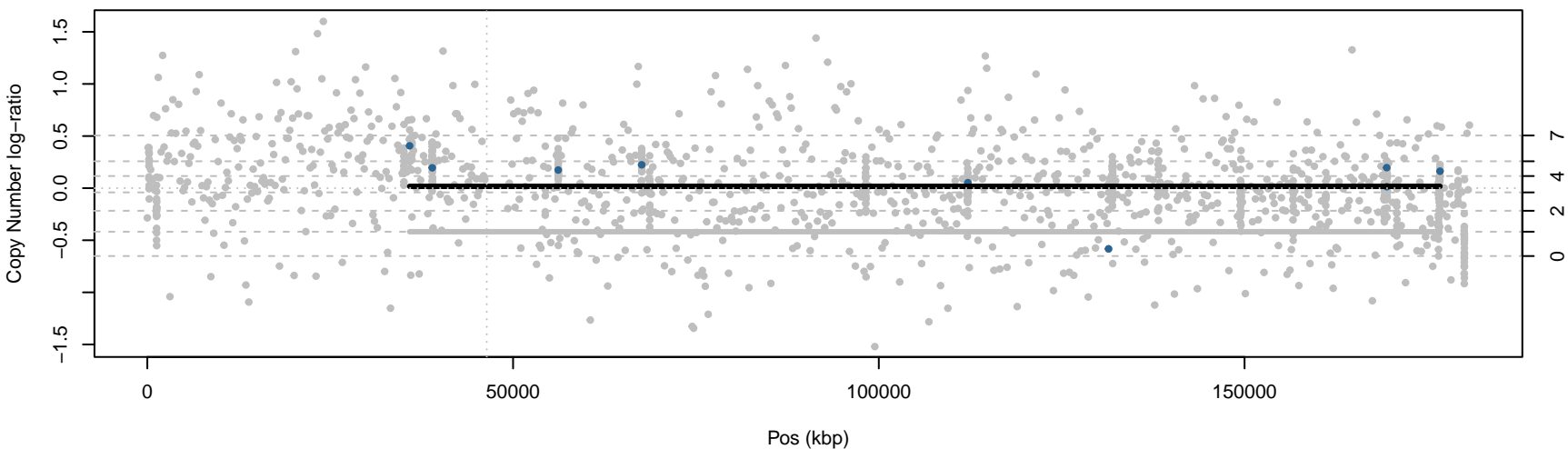
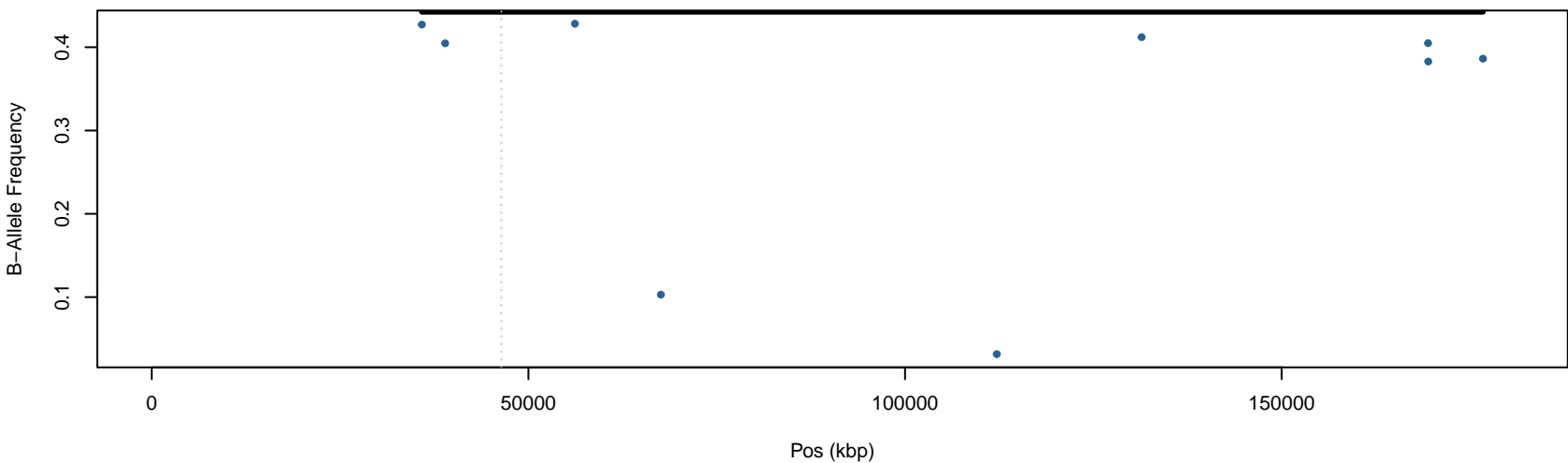
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 3



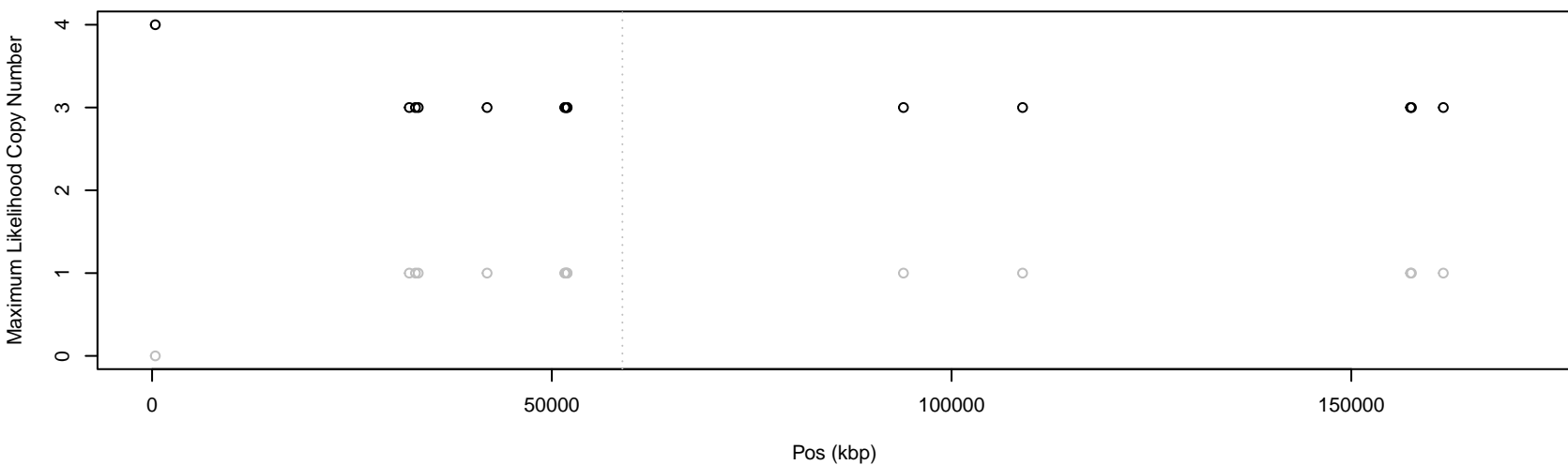
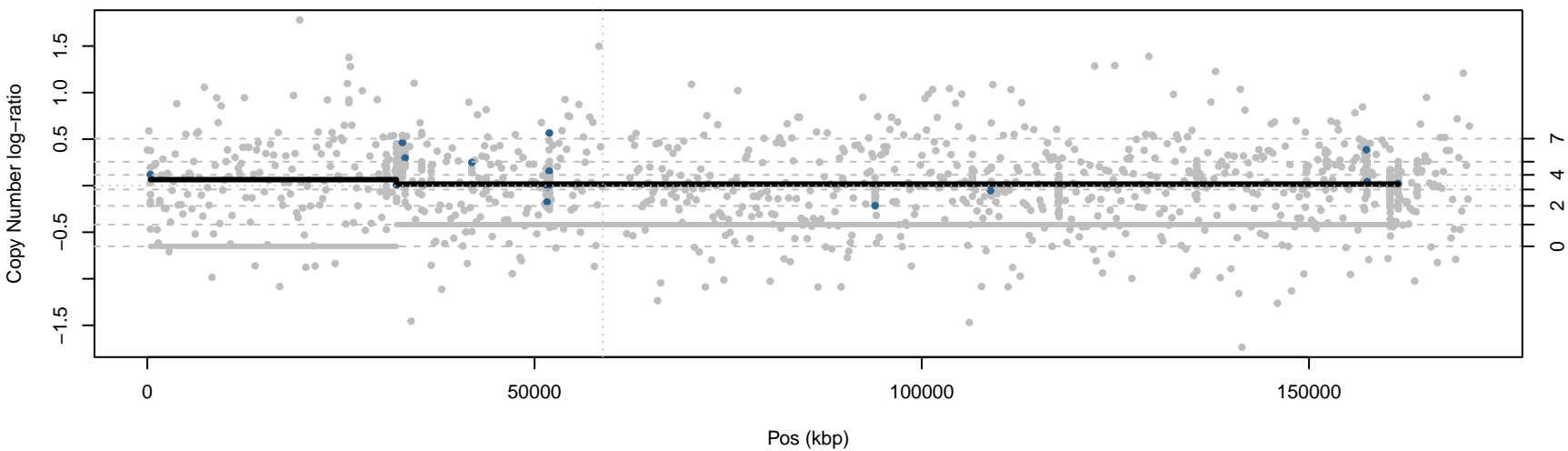
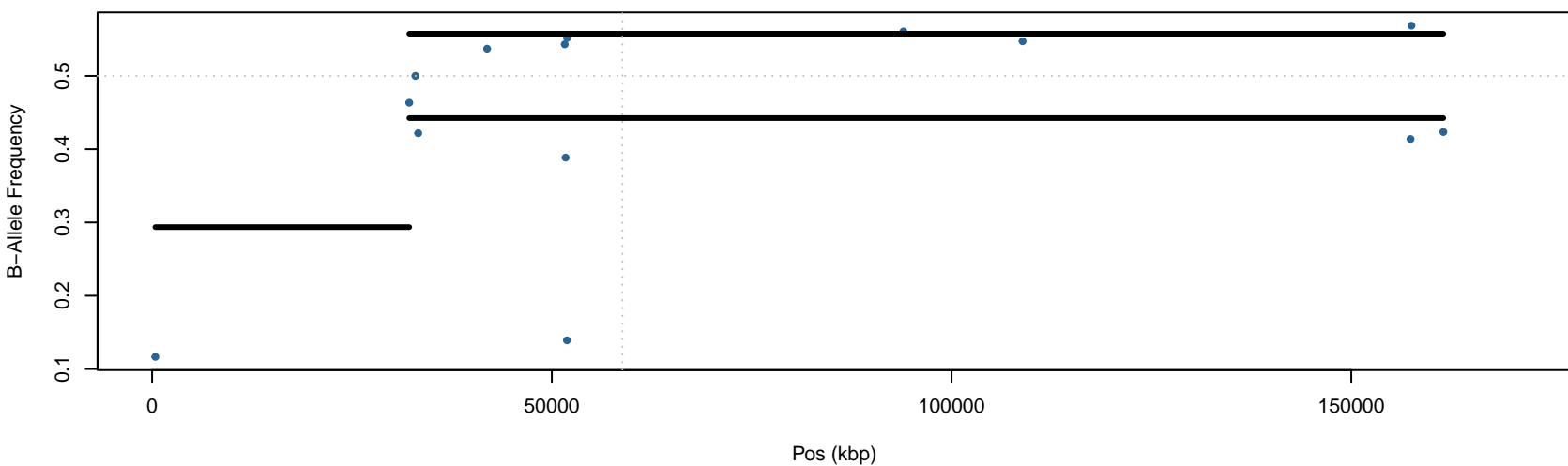
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 4



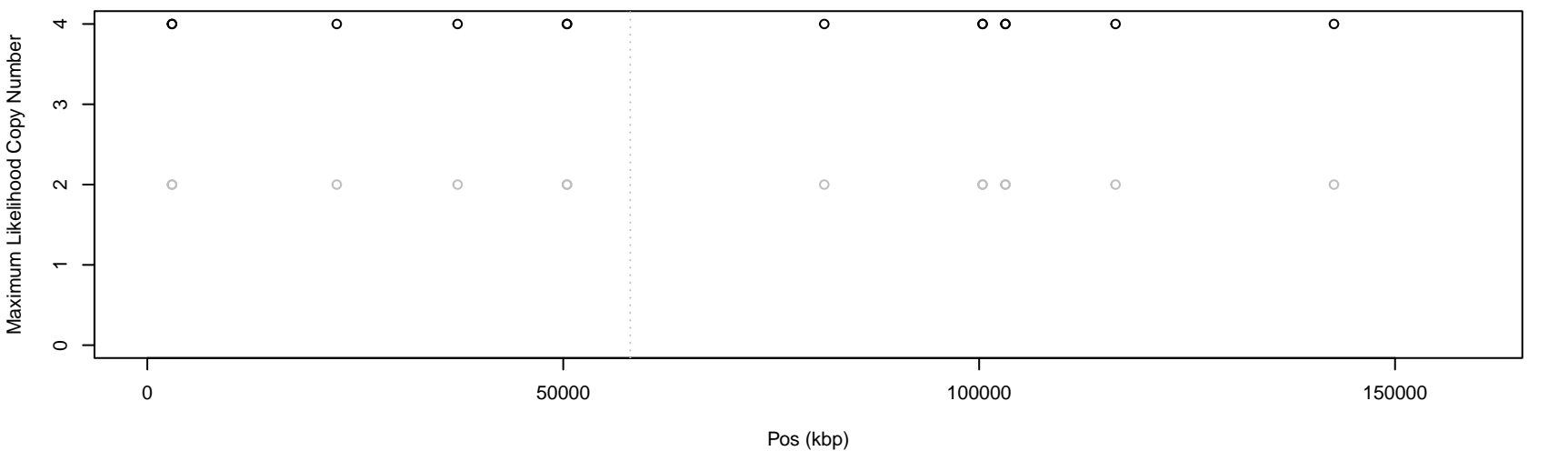
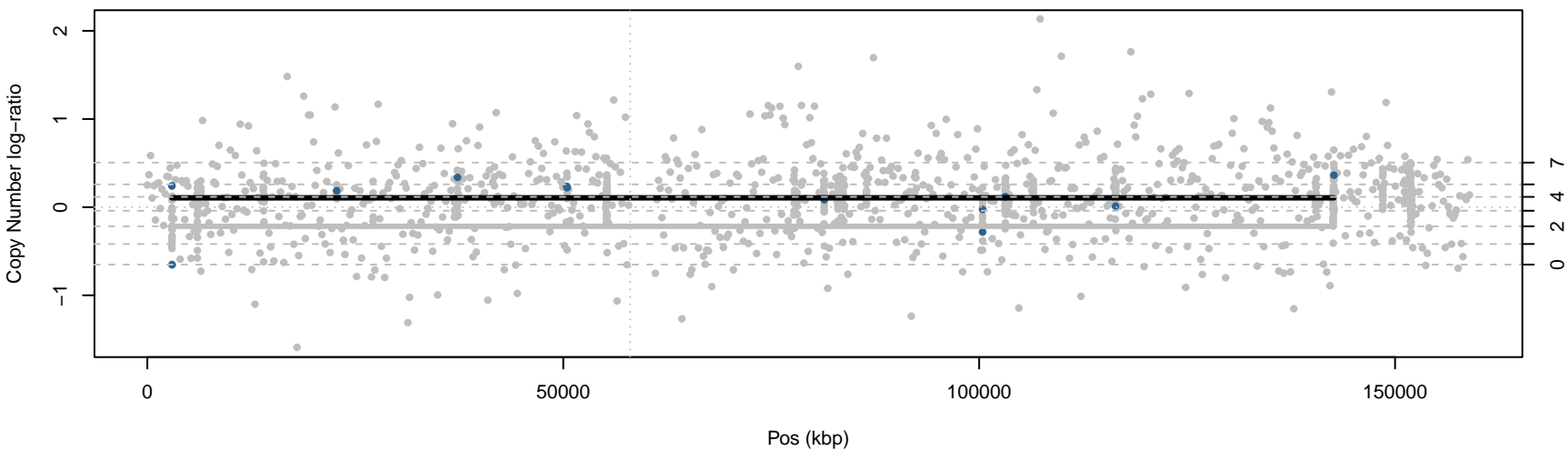
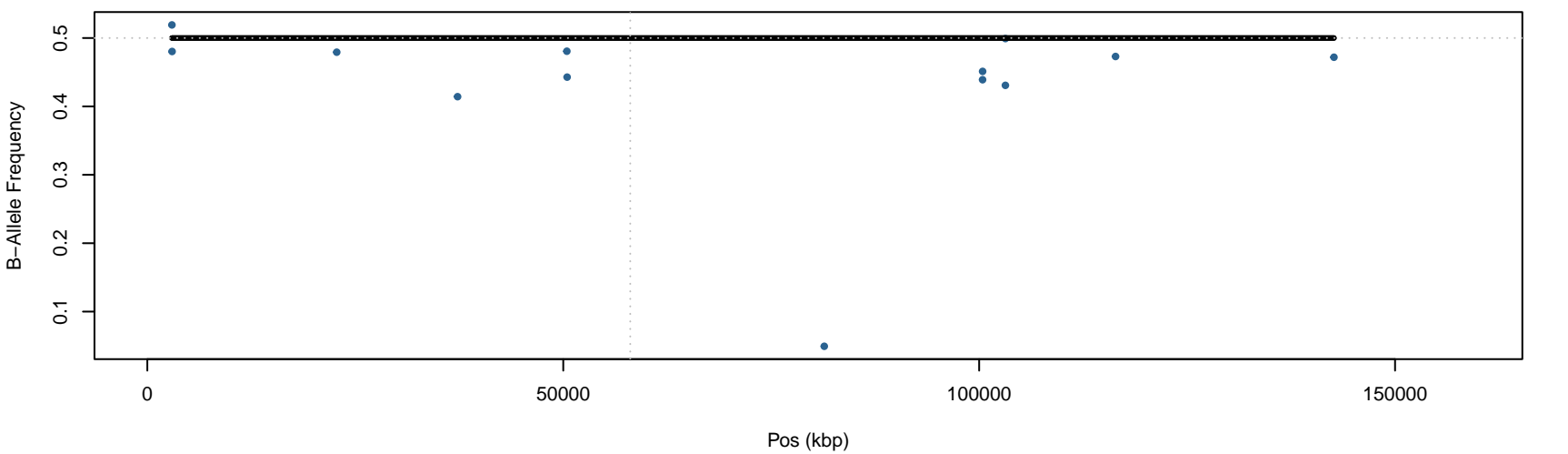
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 5



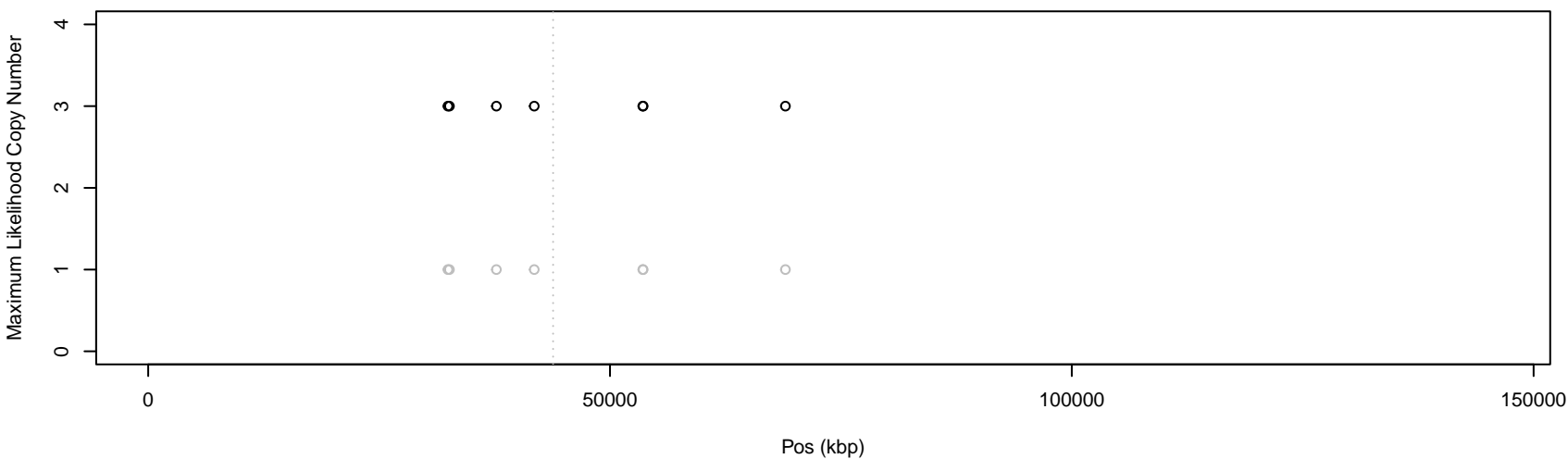
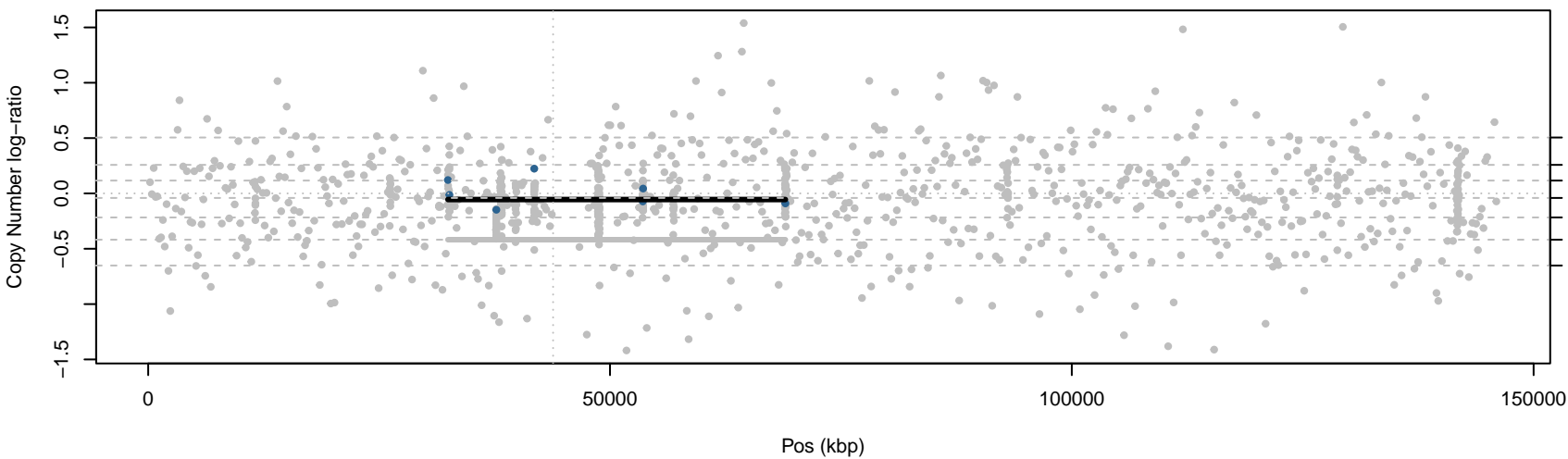
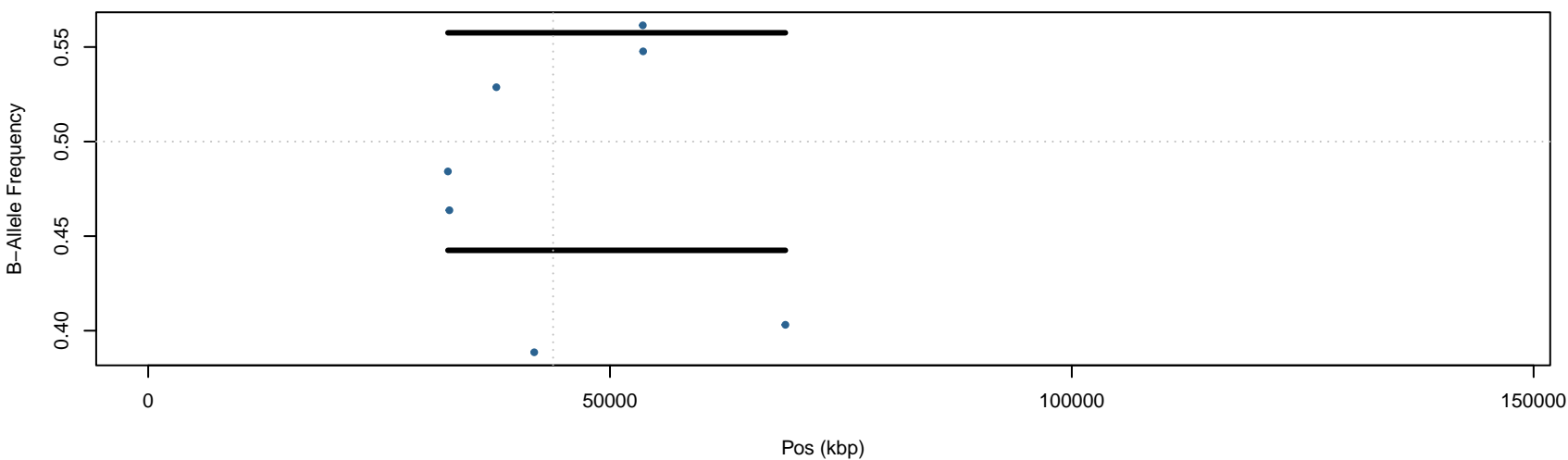
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 6



Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 7

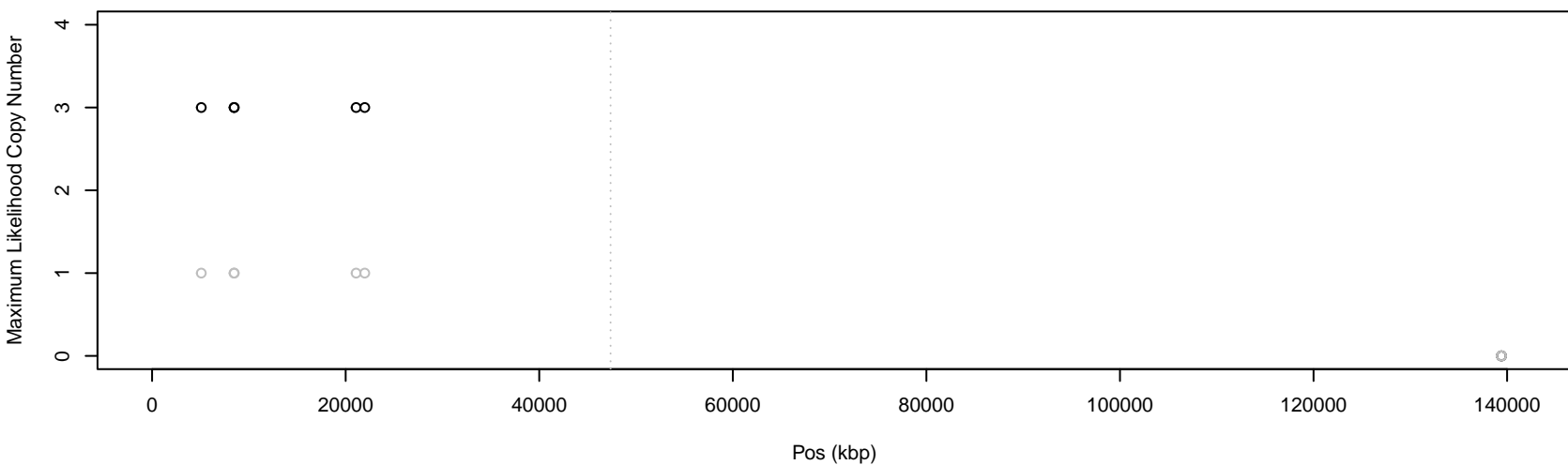
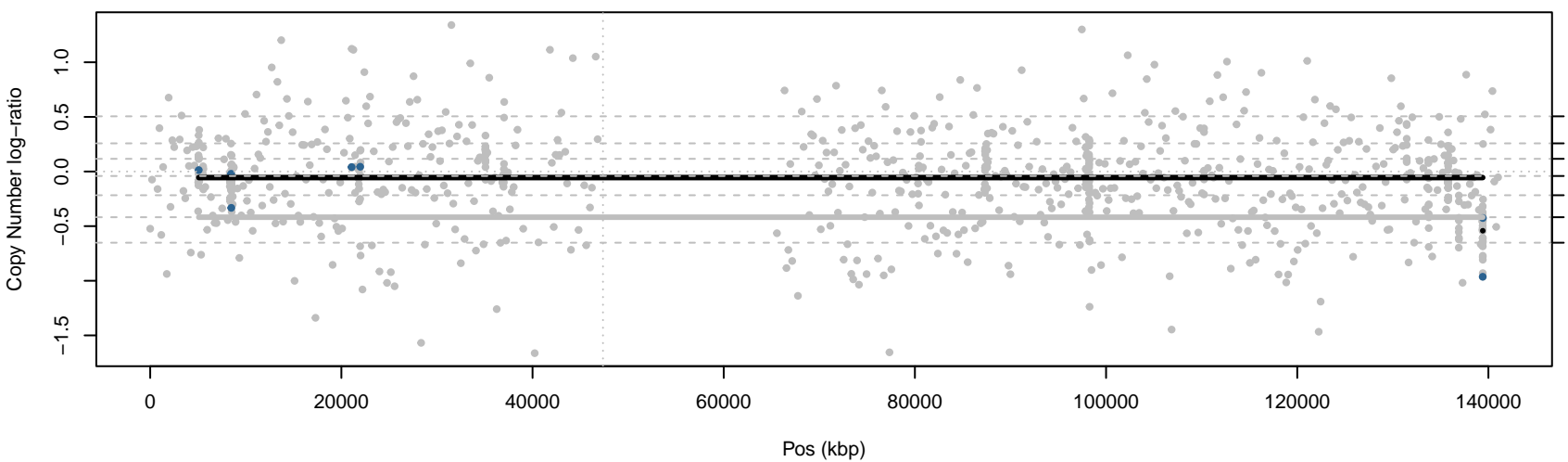
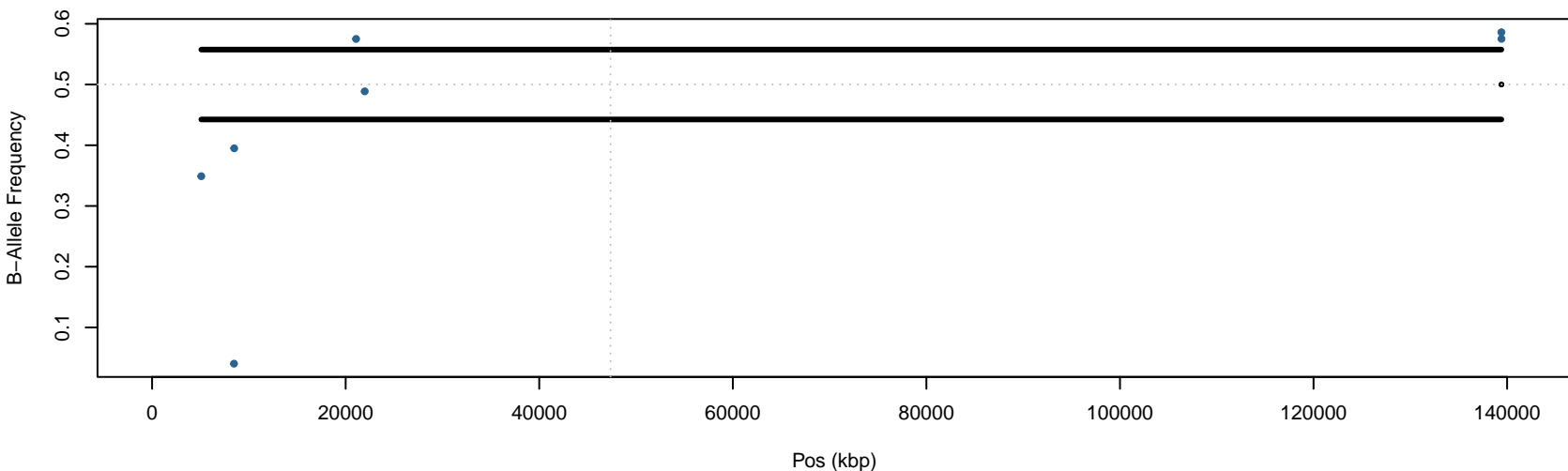


Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 8

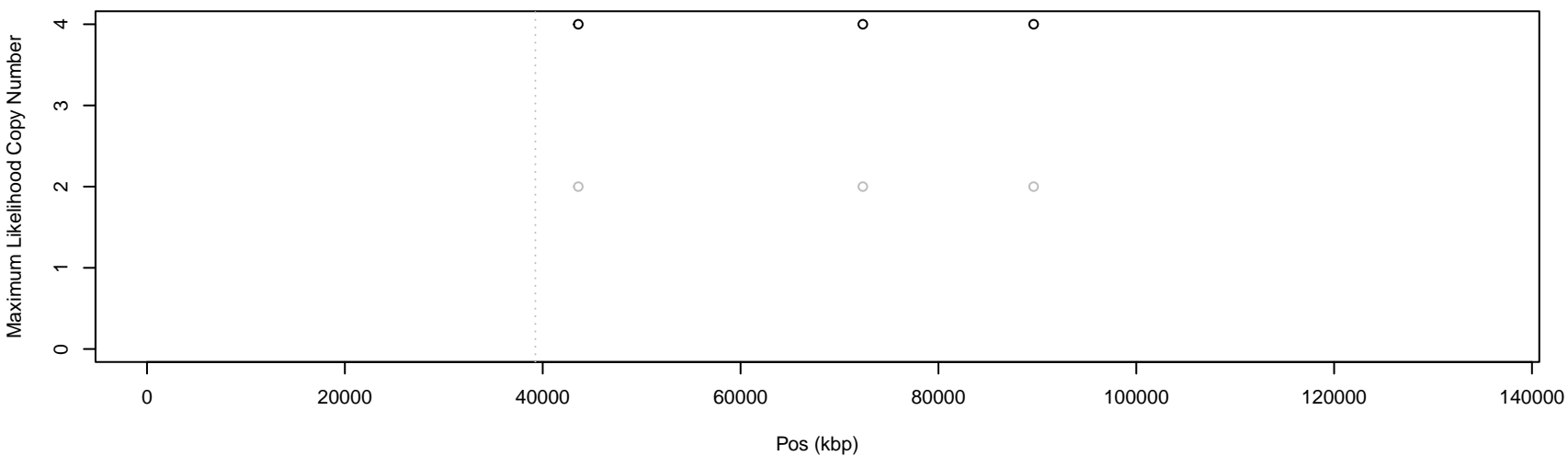
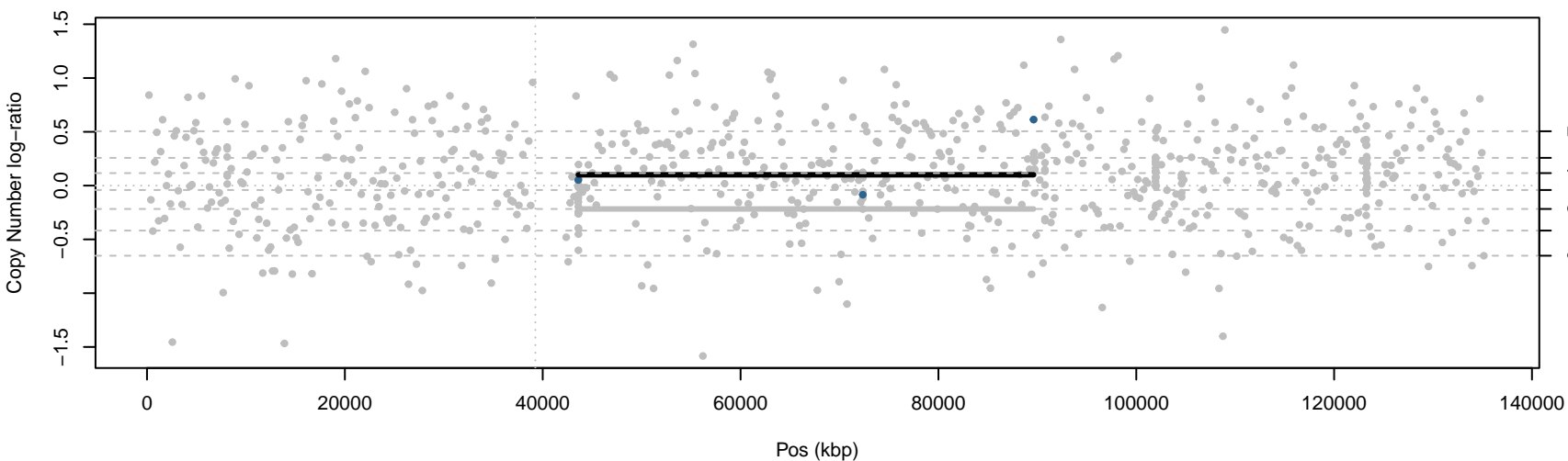
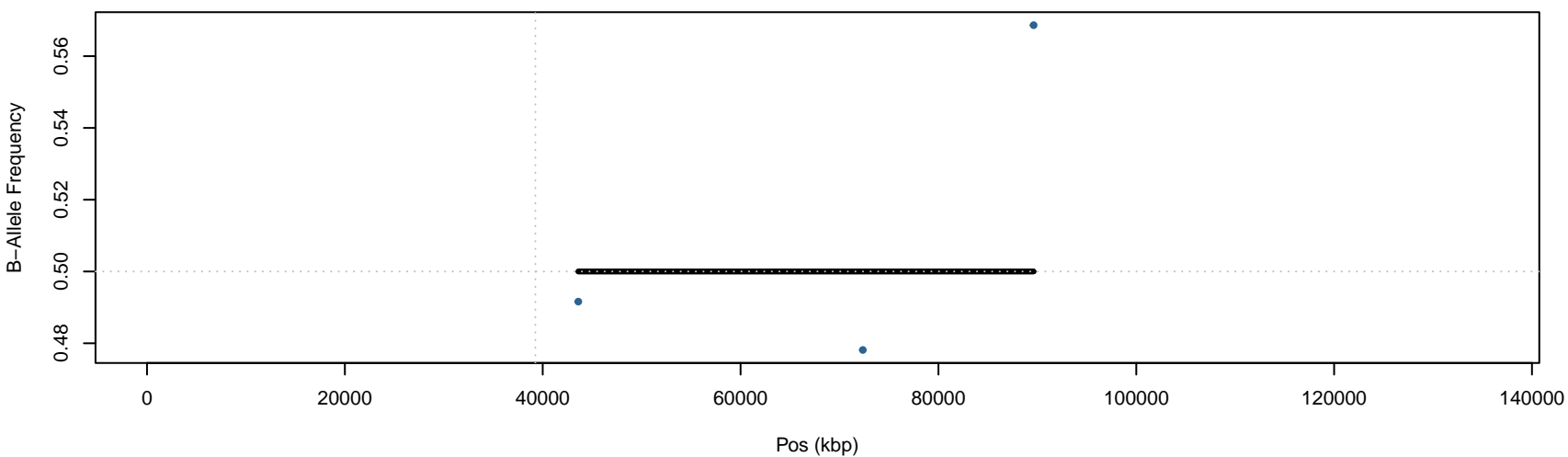




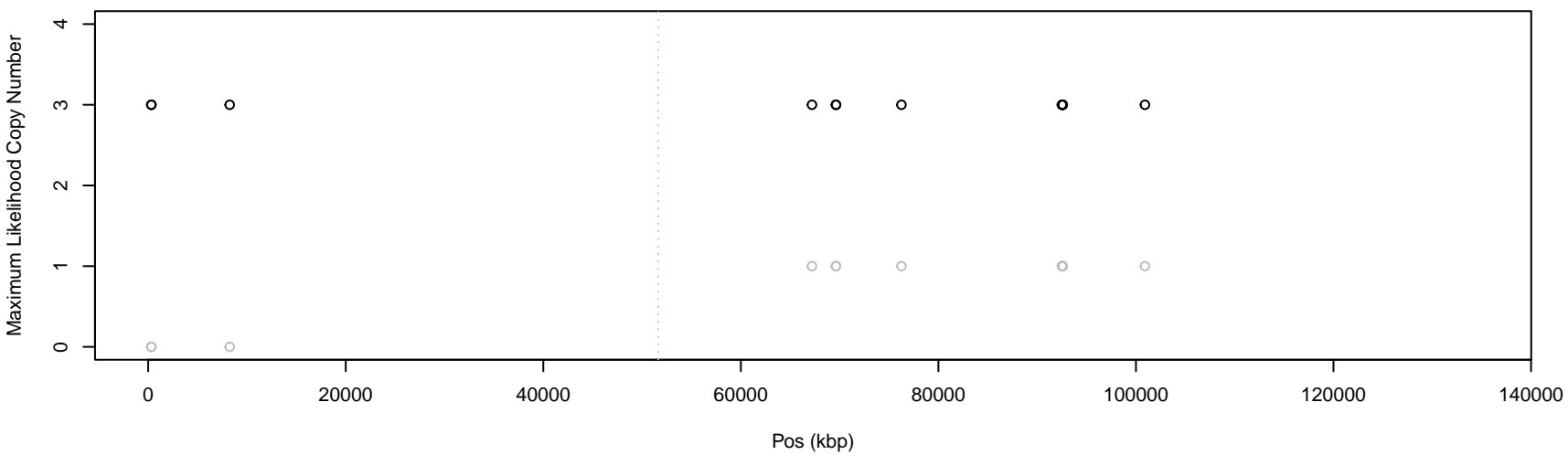
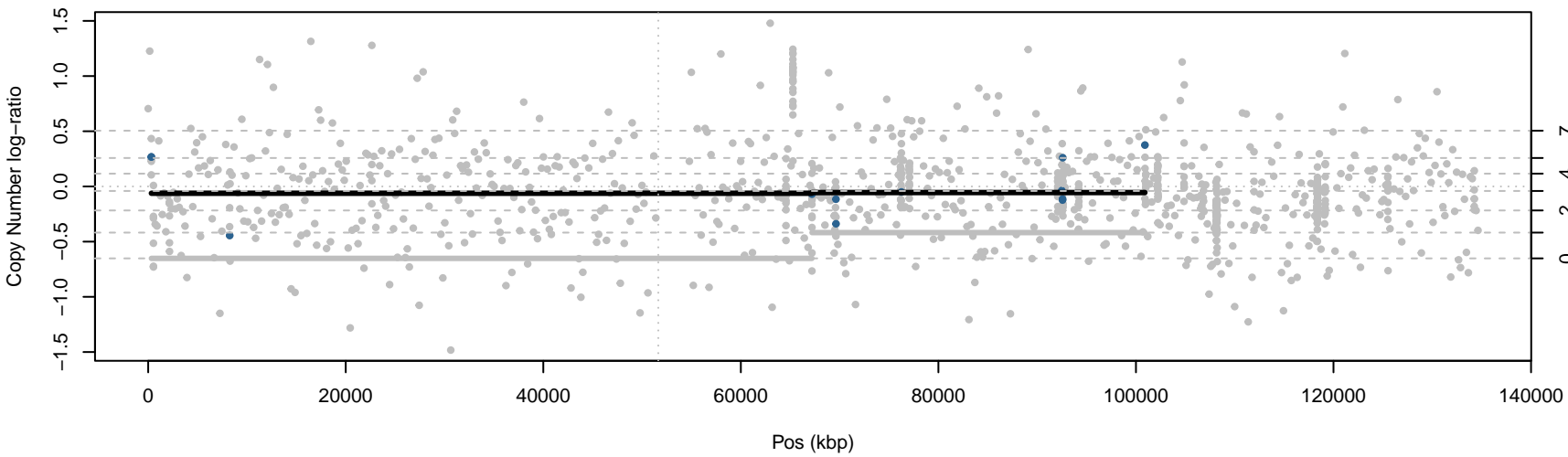
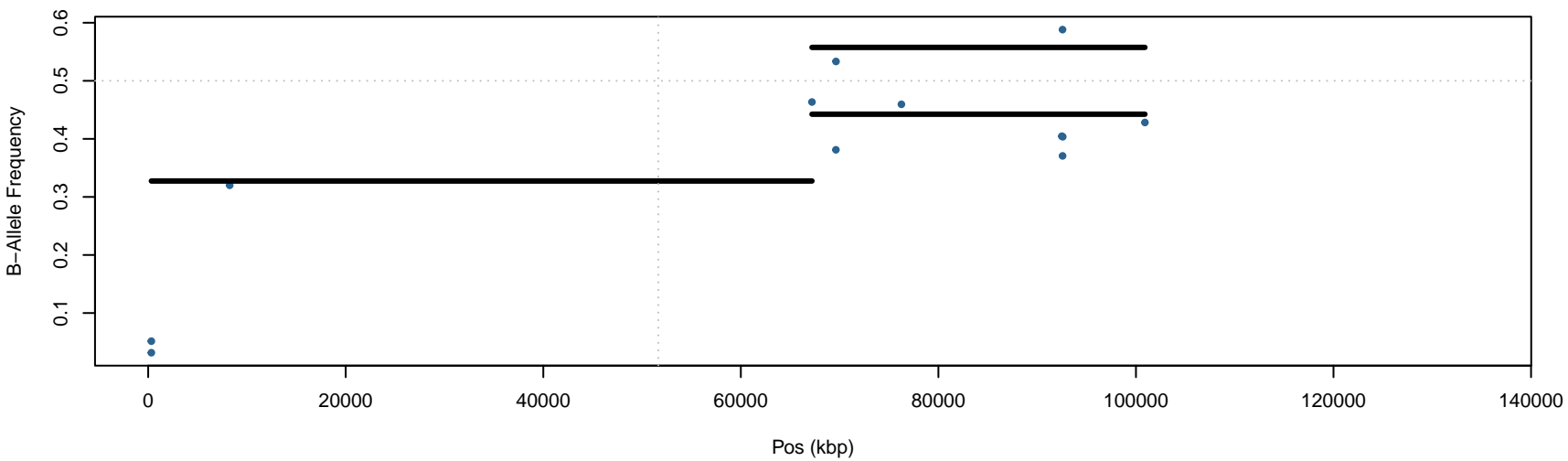
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 9



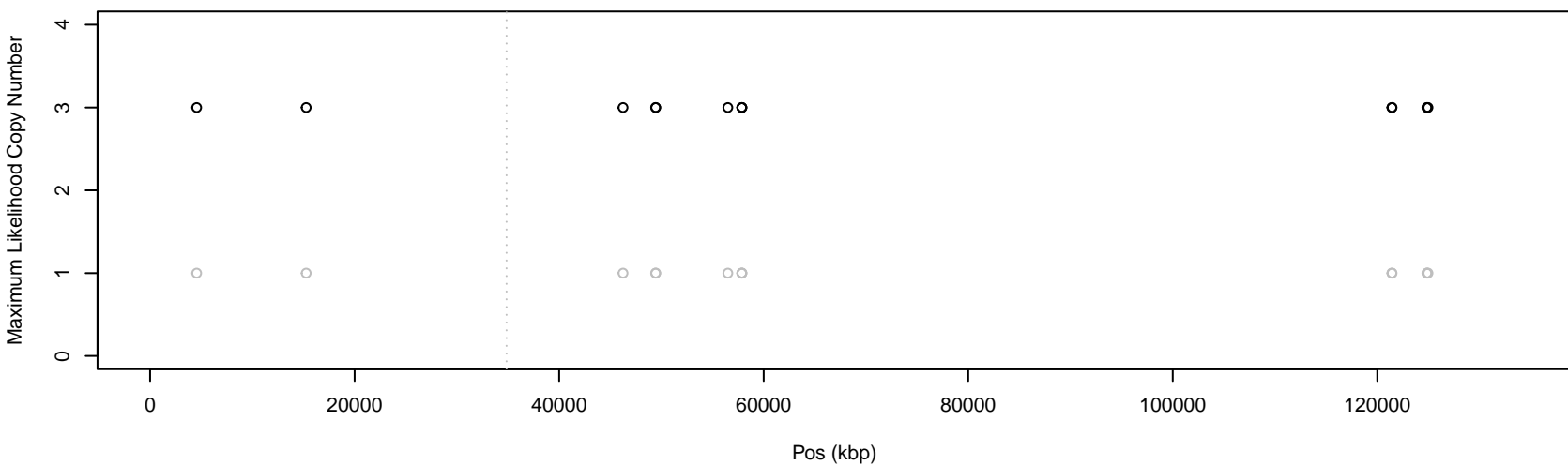
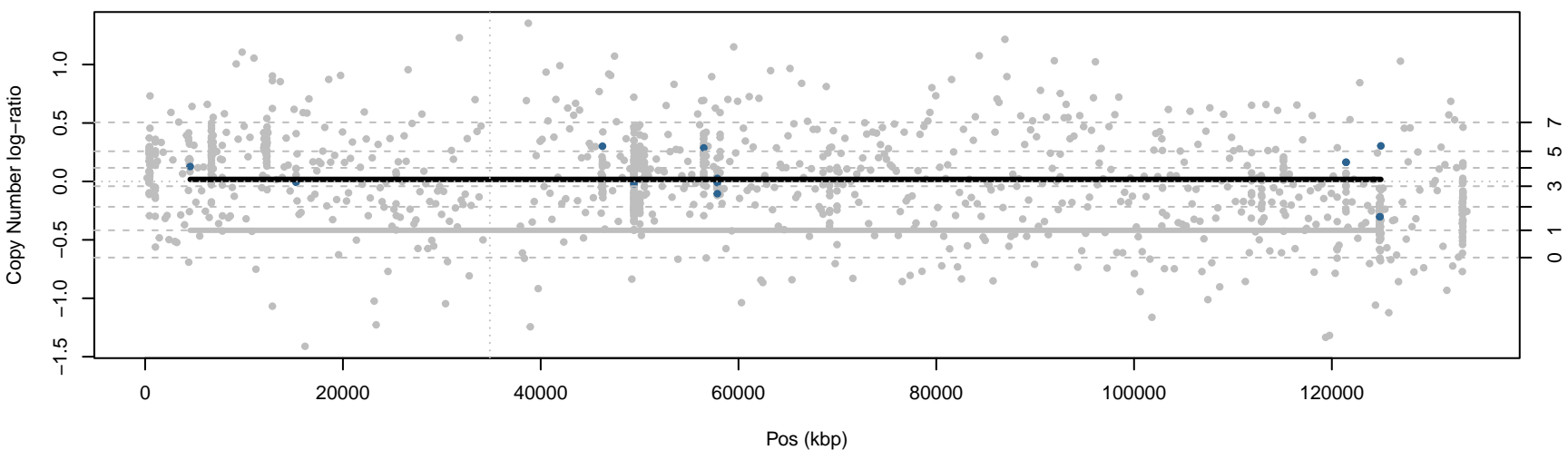
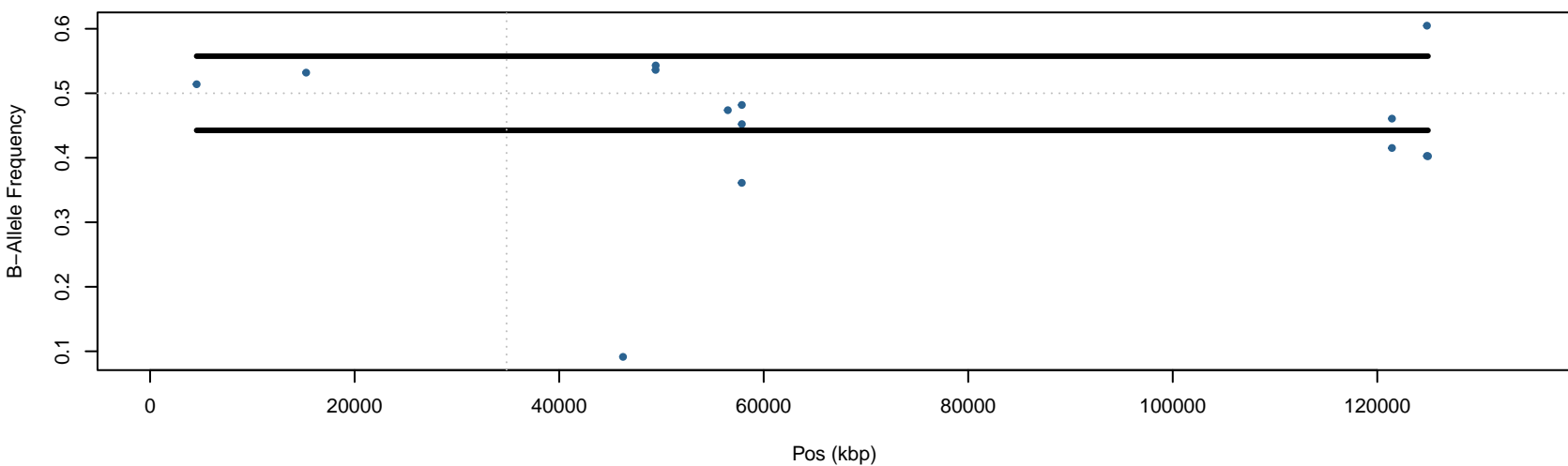
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 10



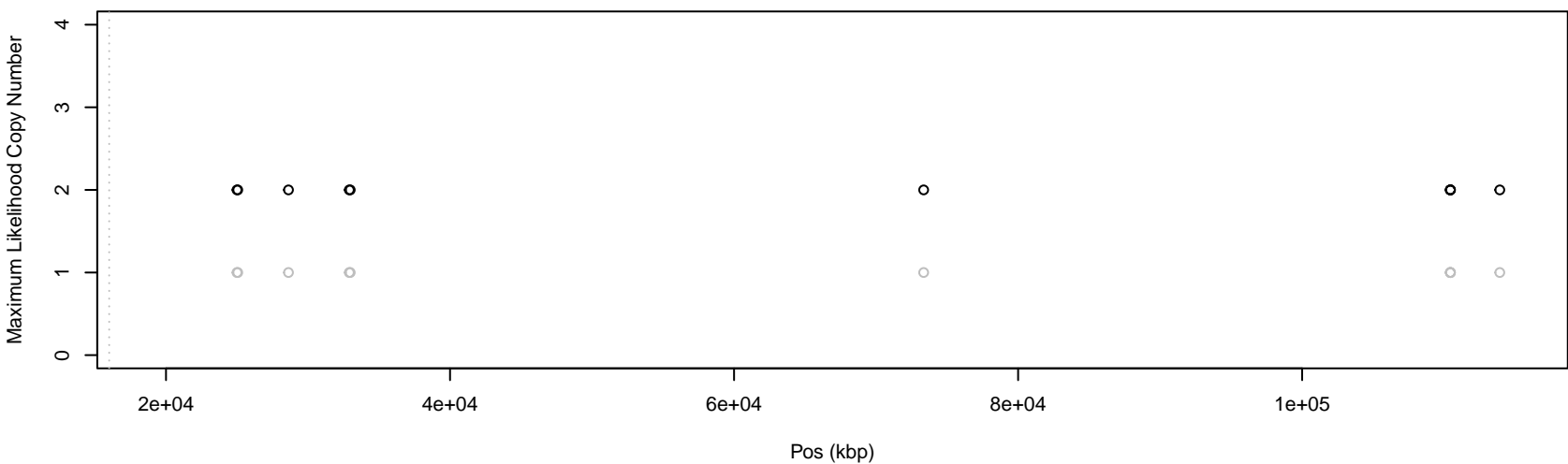
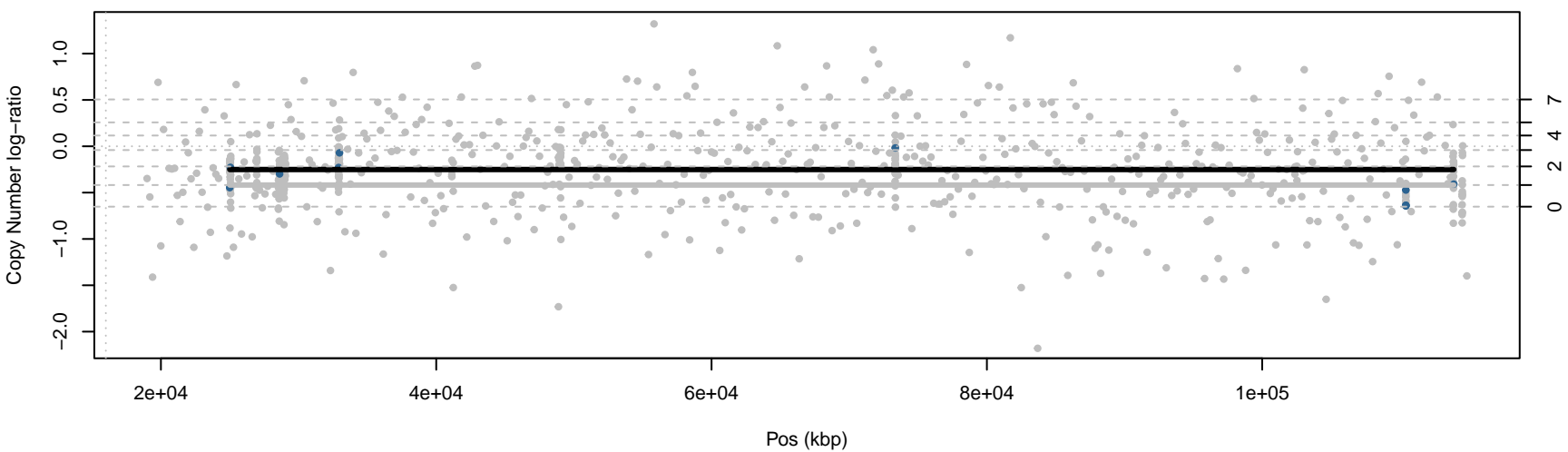
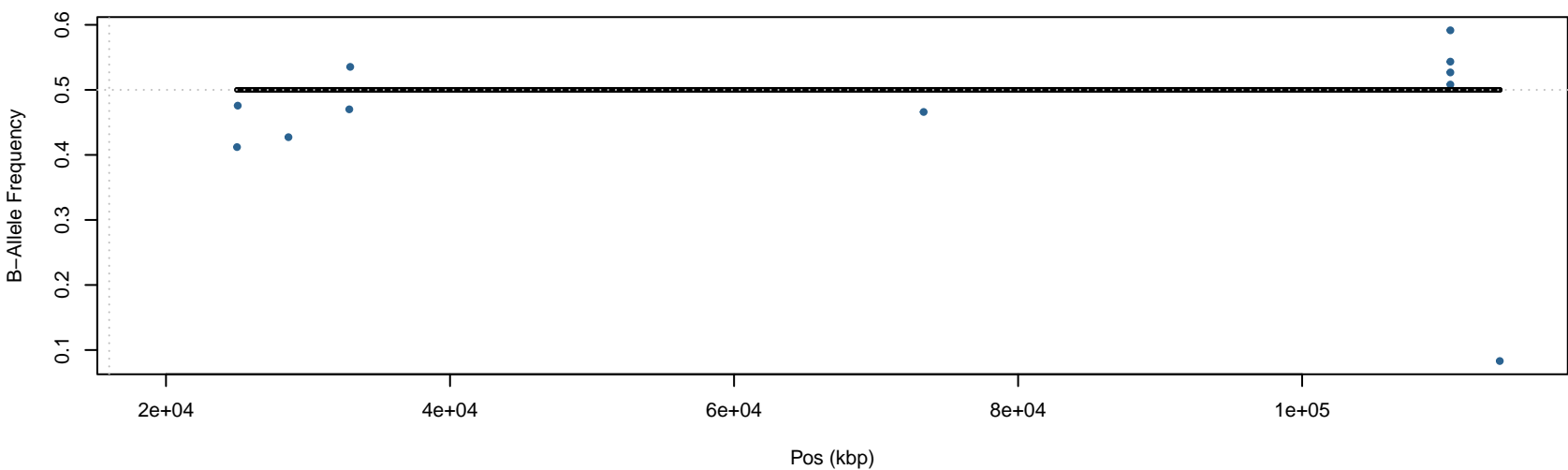
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 11



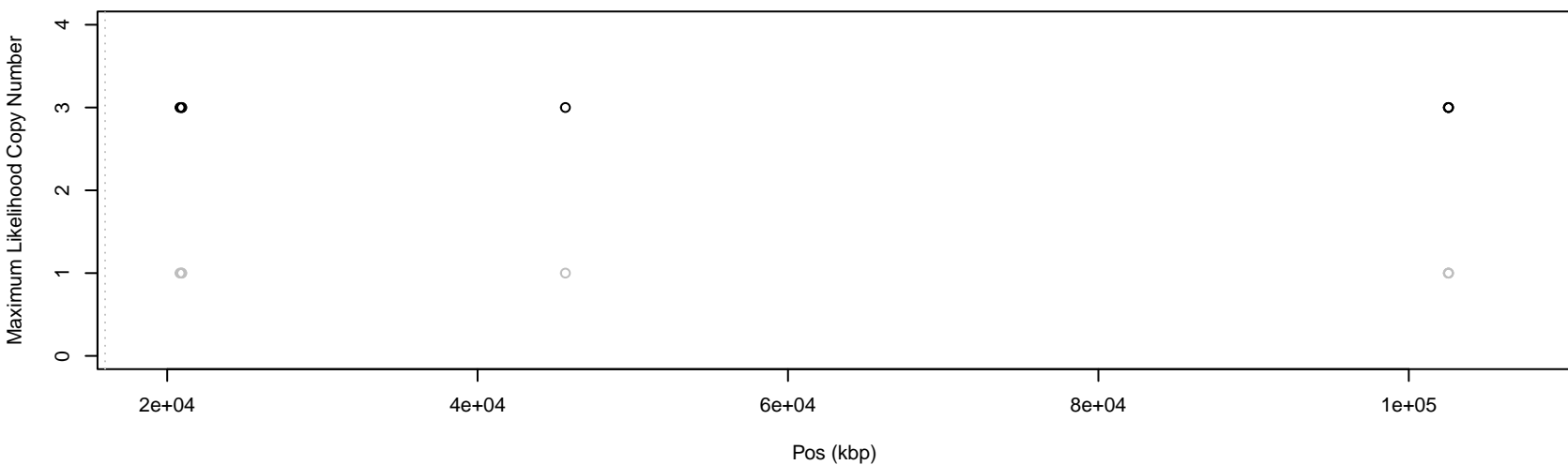
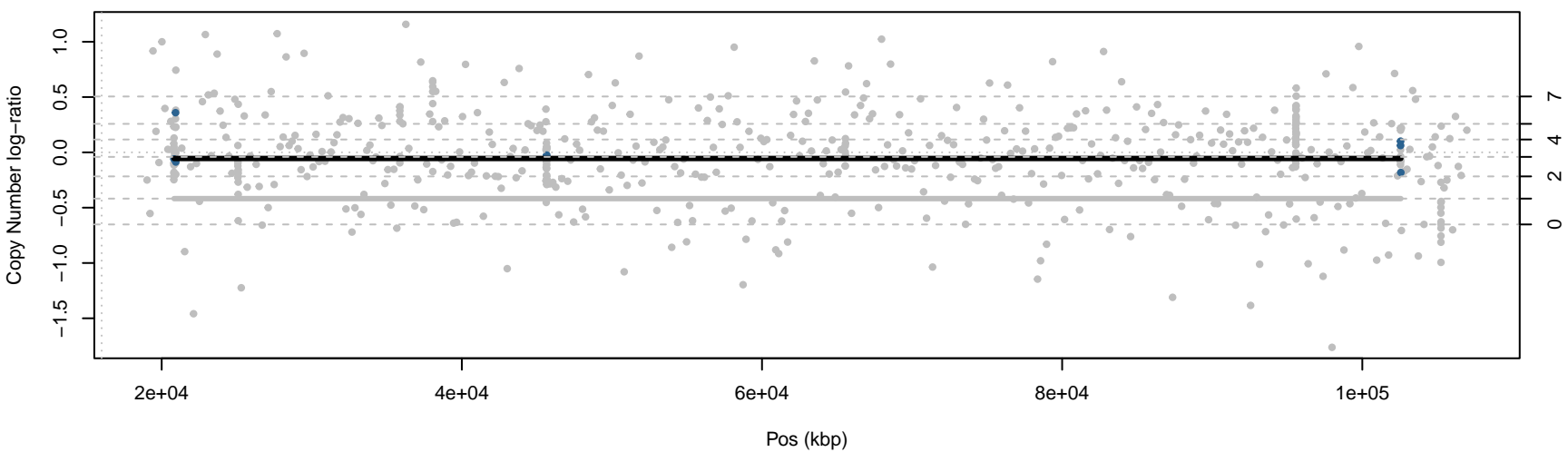
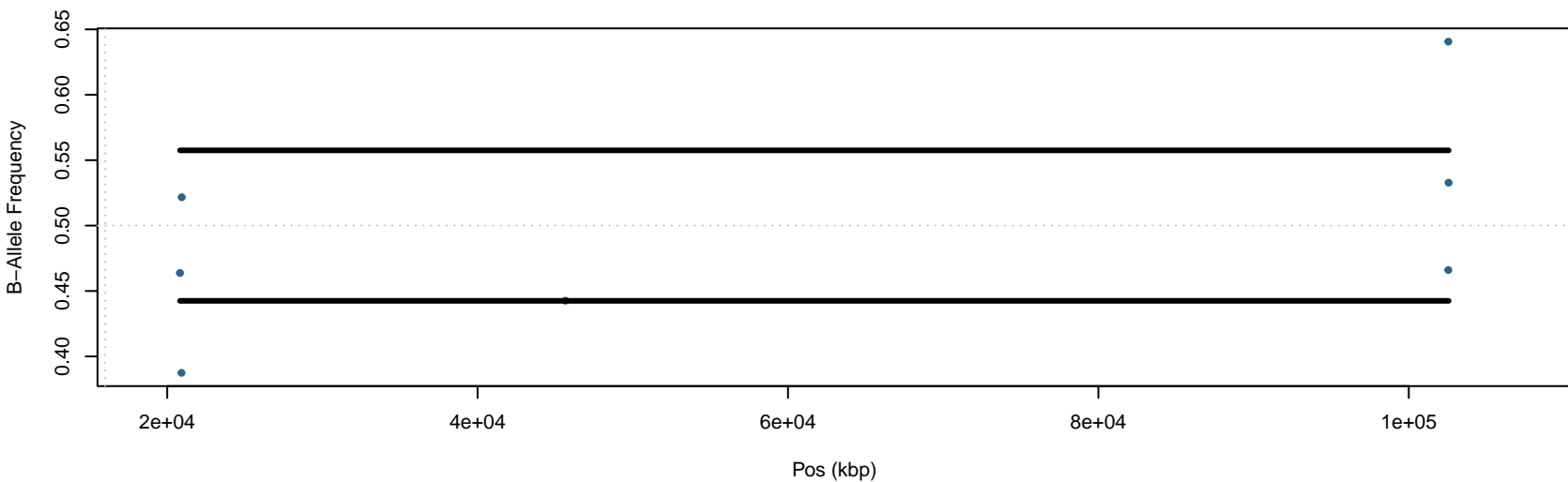
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 12



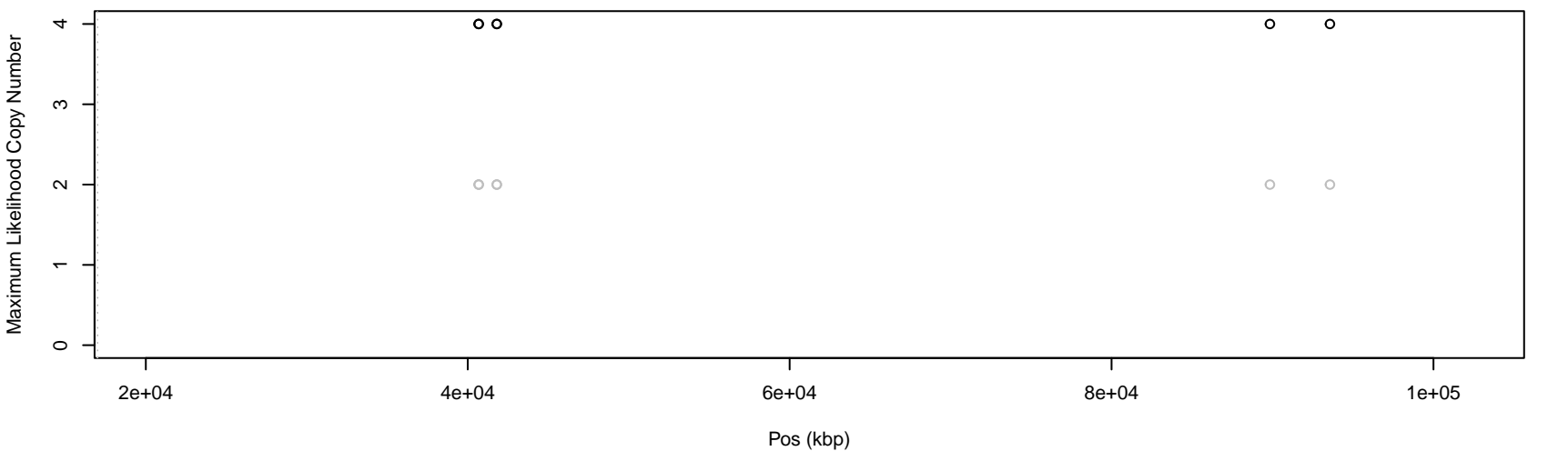
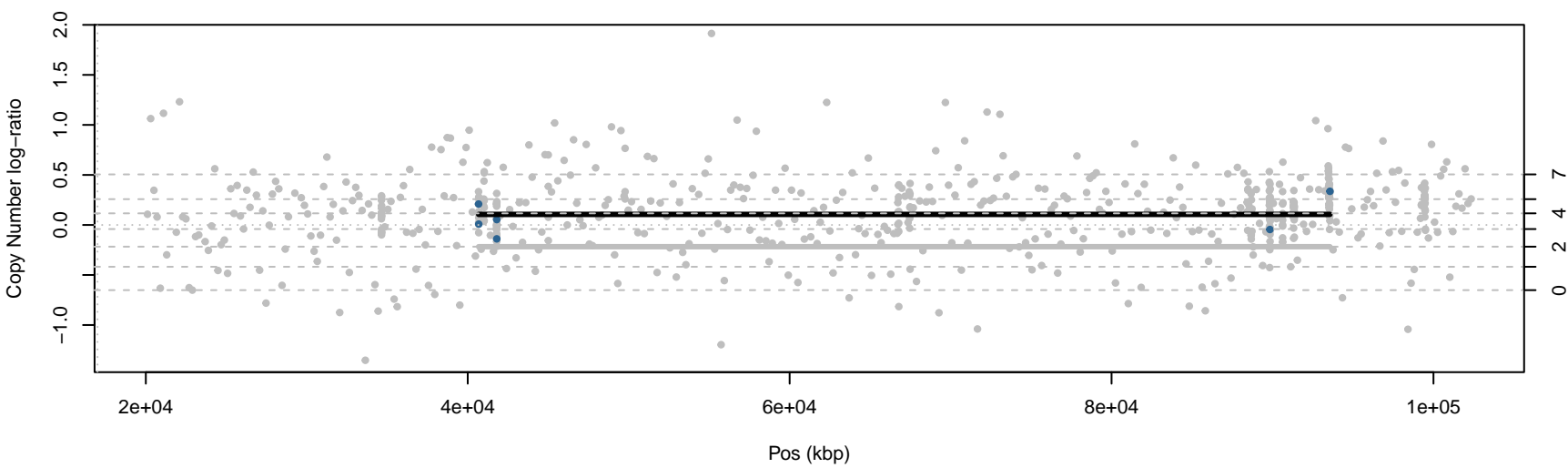
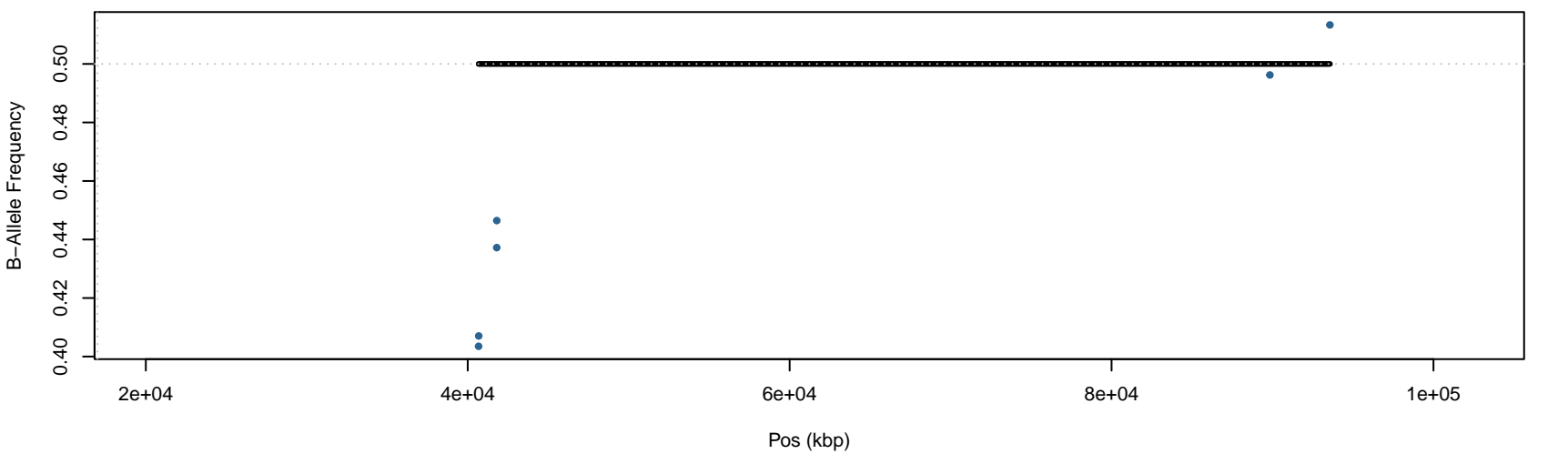
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 13



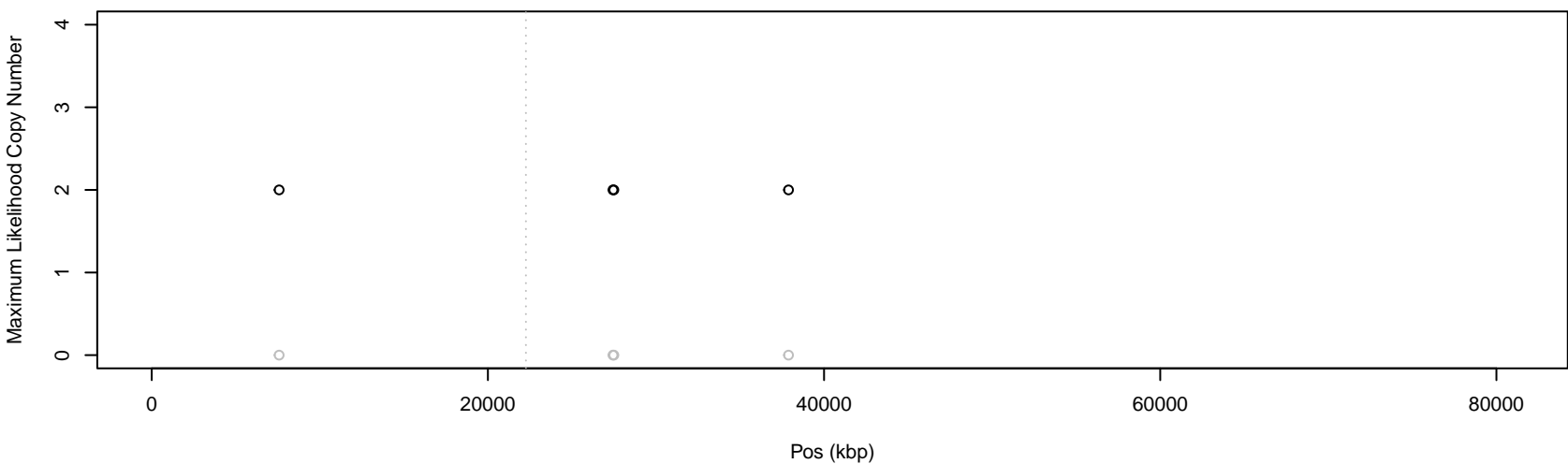
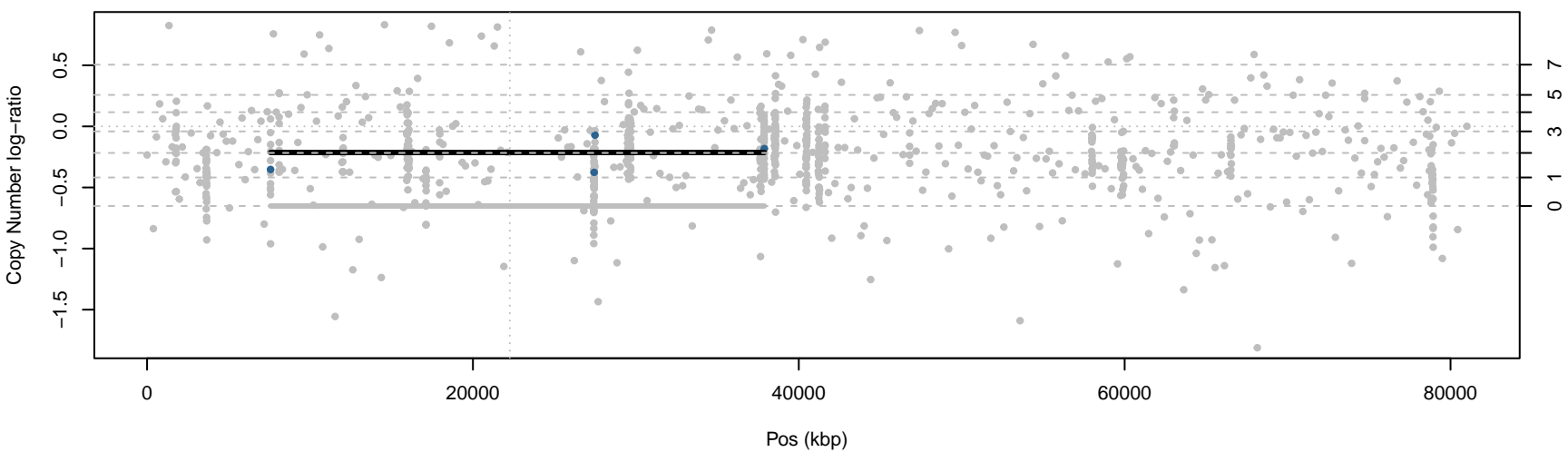
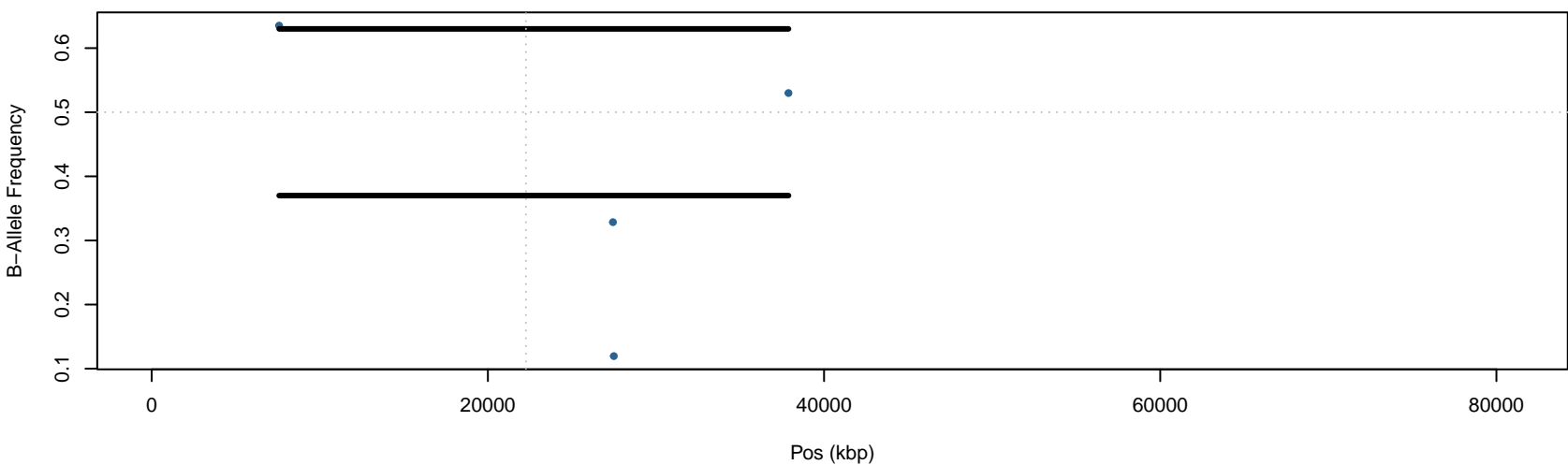
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 14



Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 15

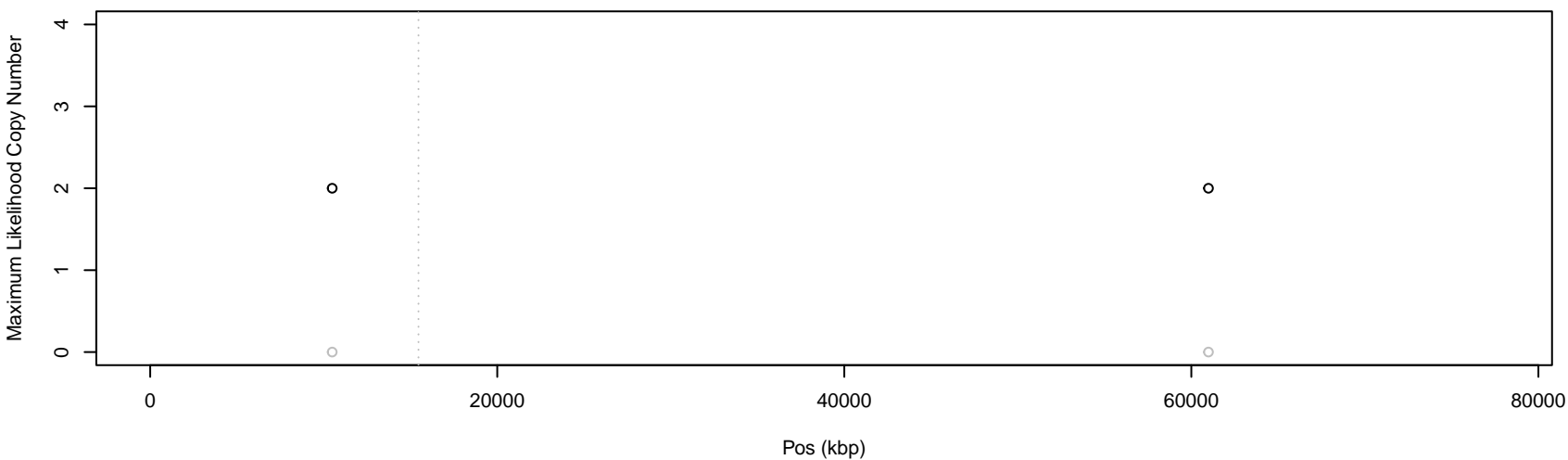
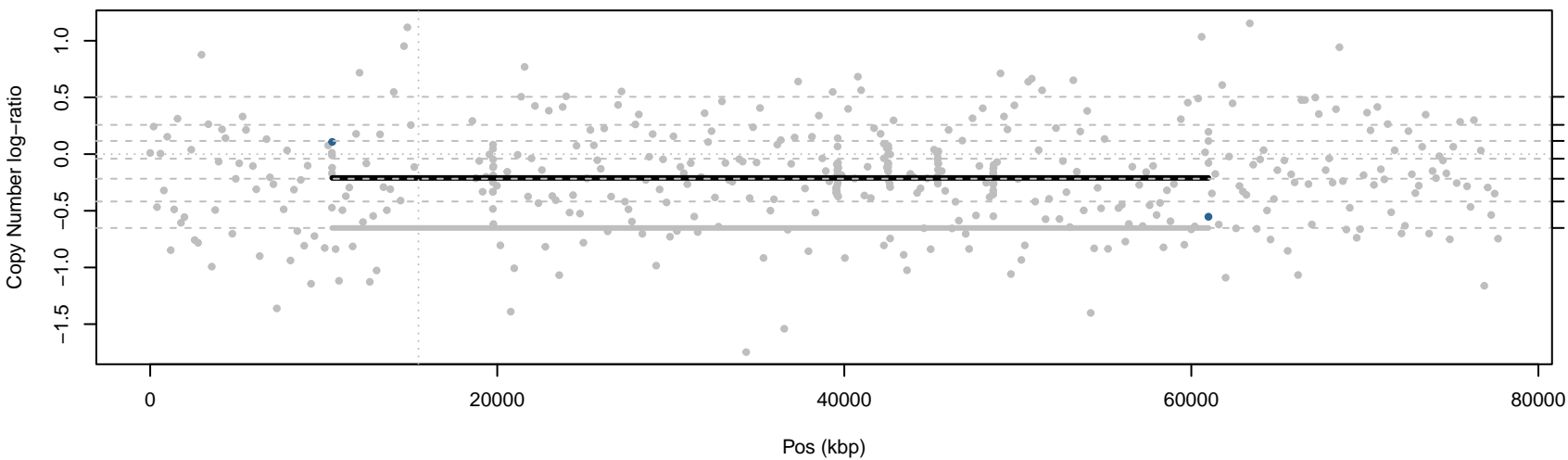
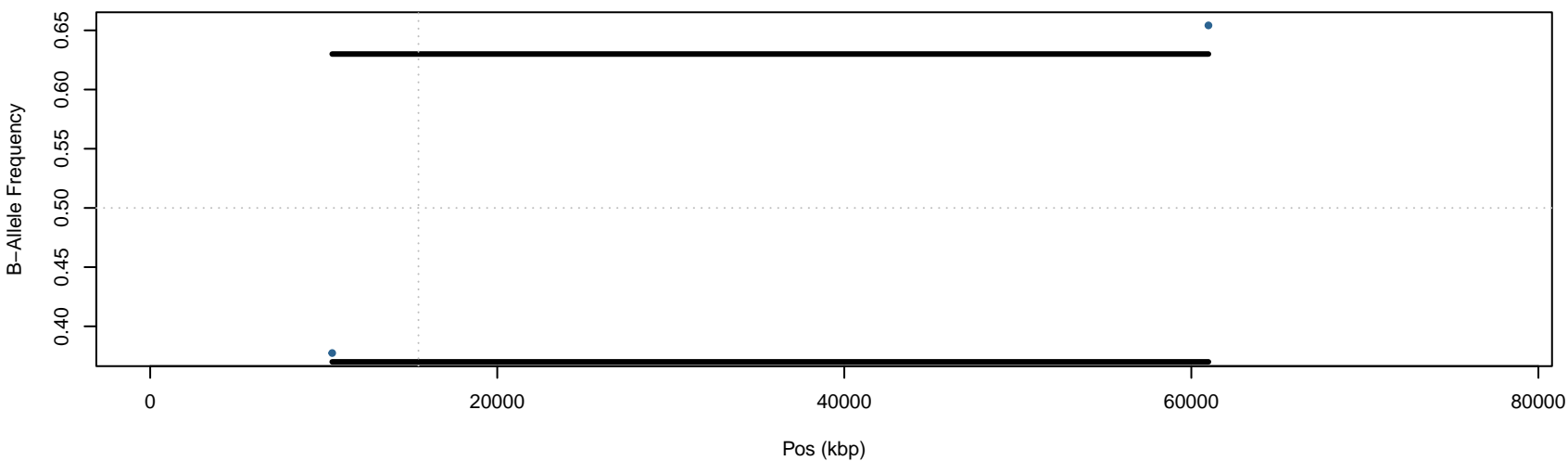


Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 17

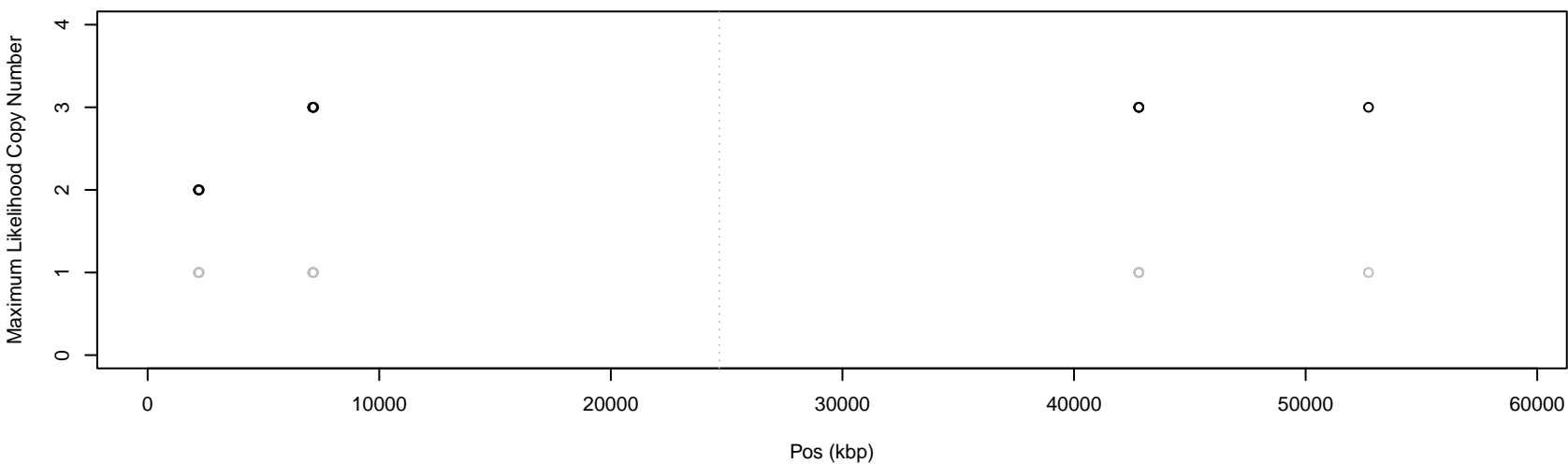
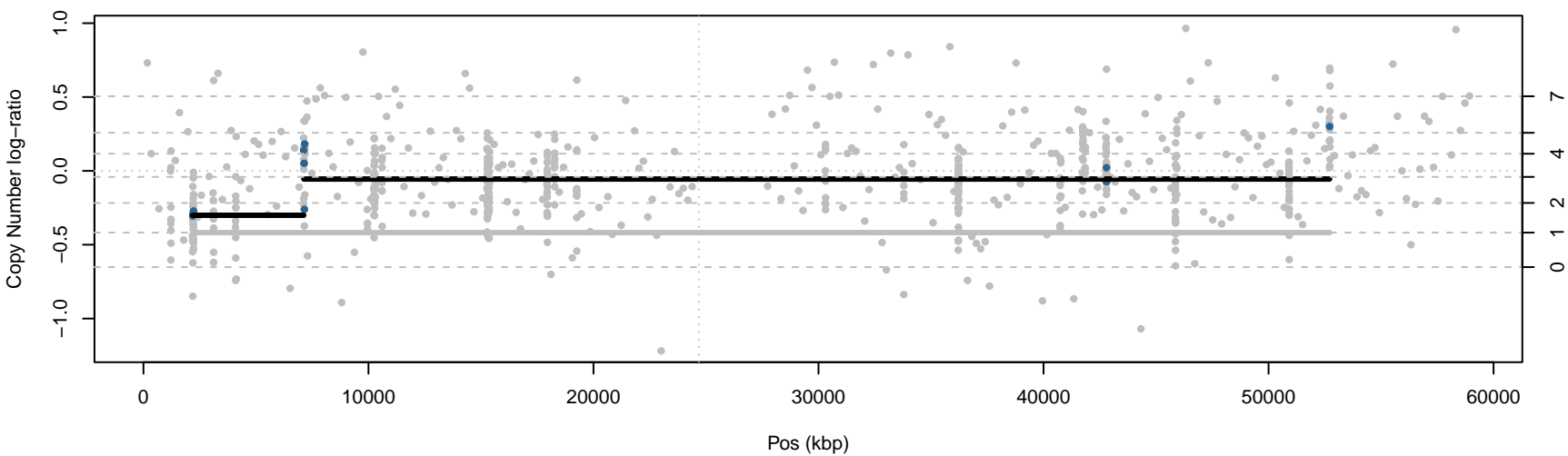
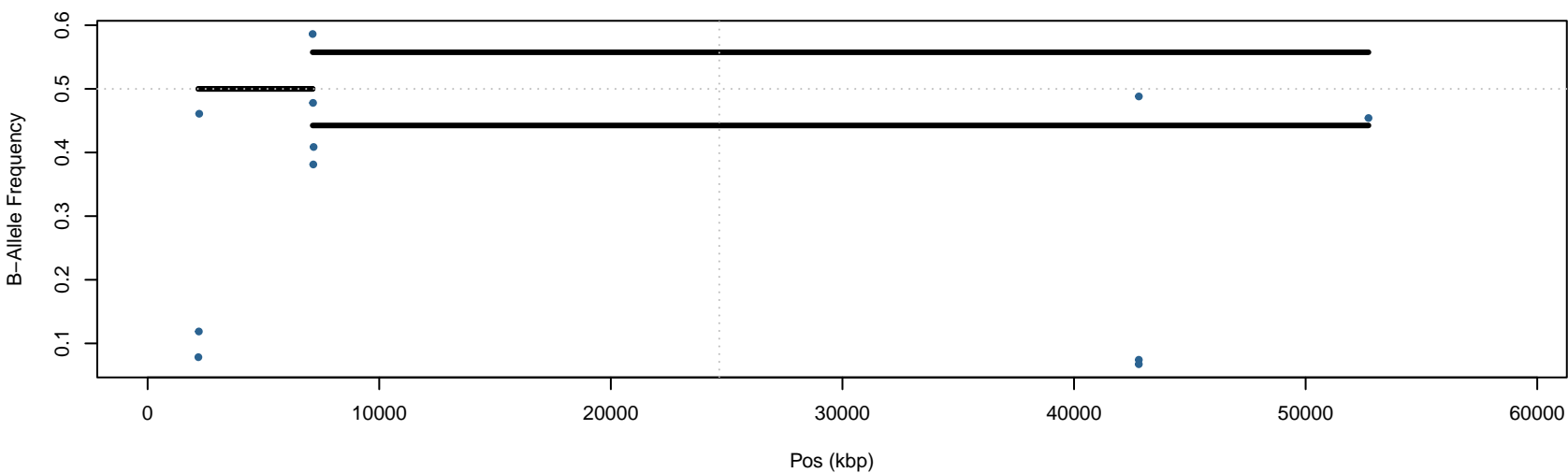




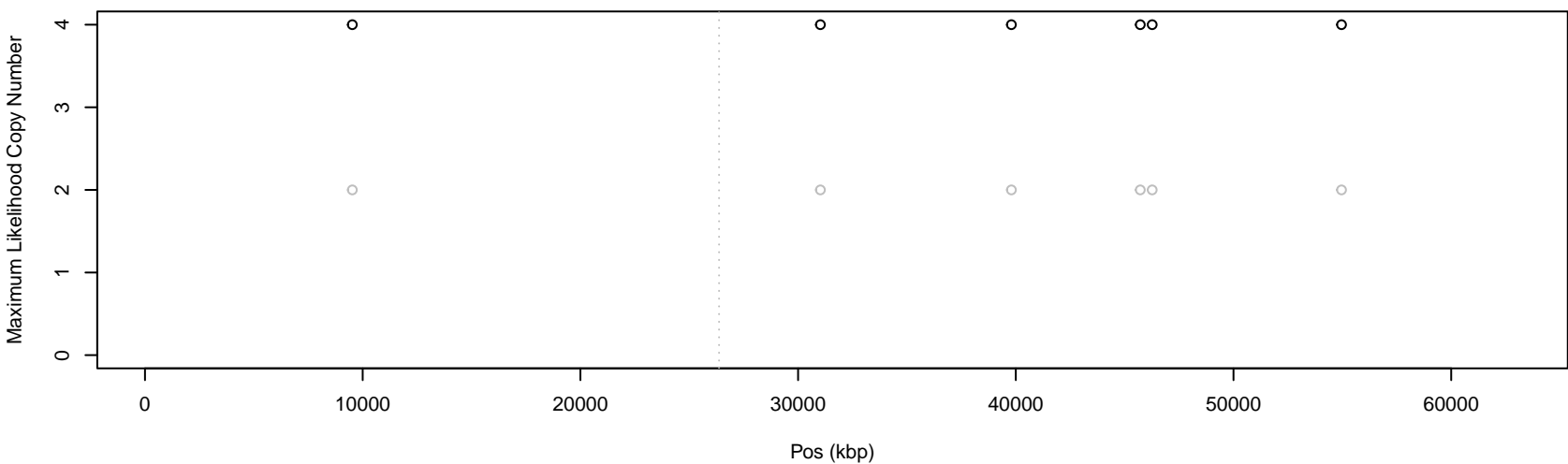
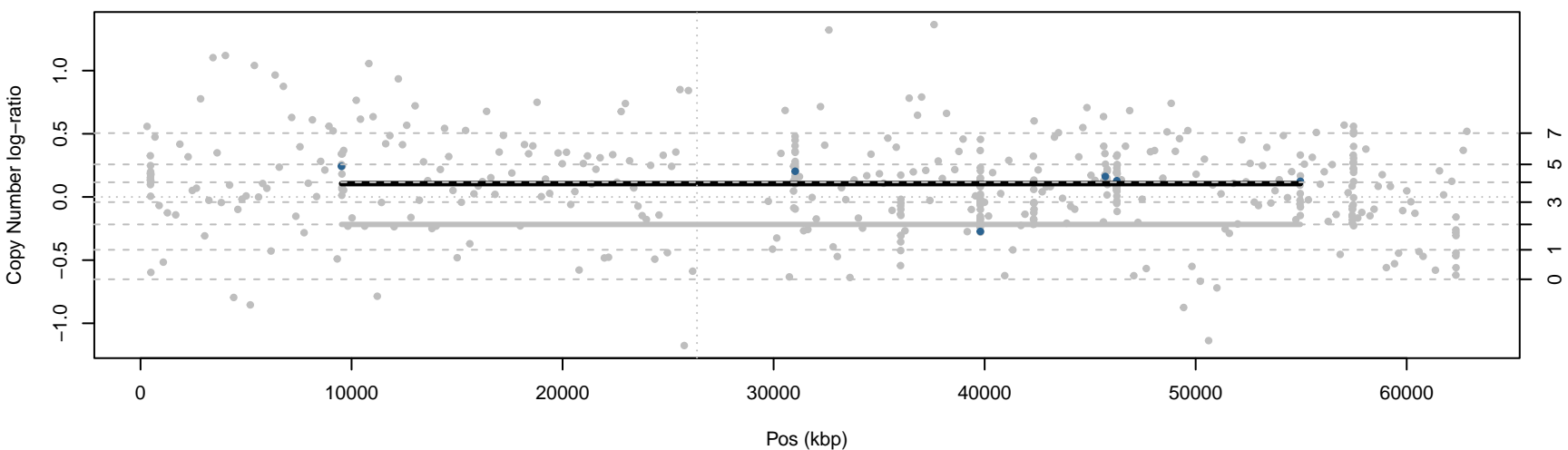
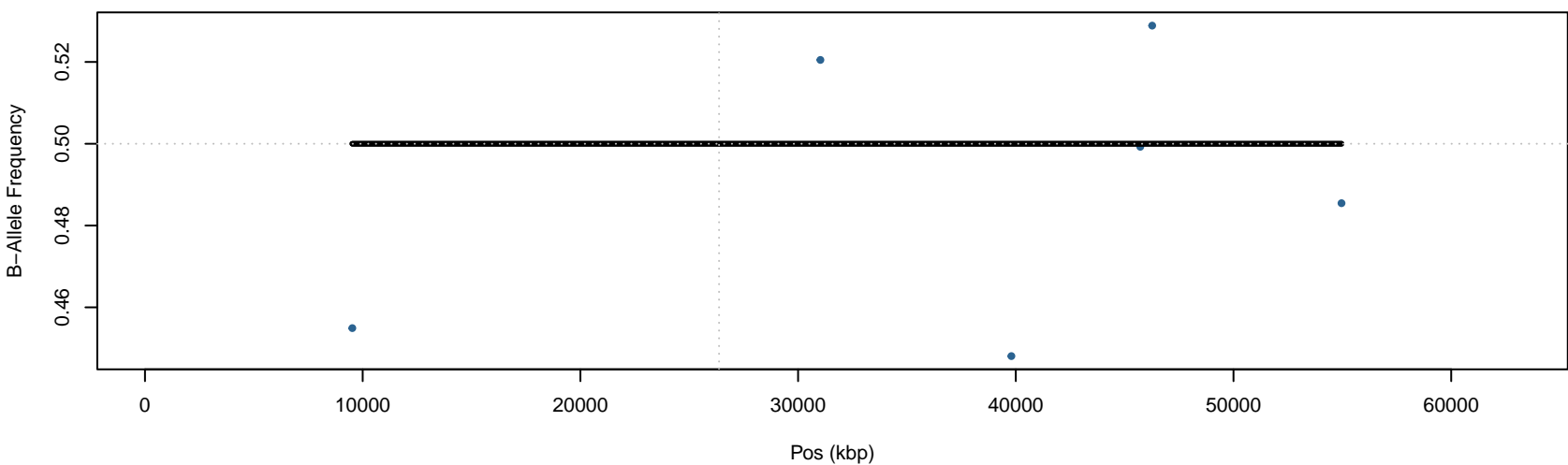
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 18



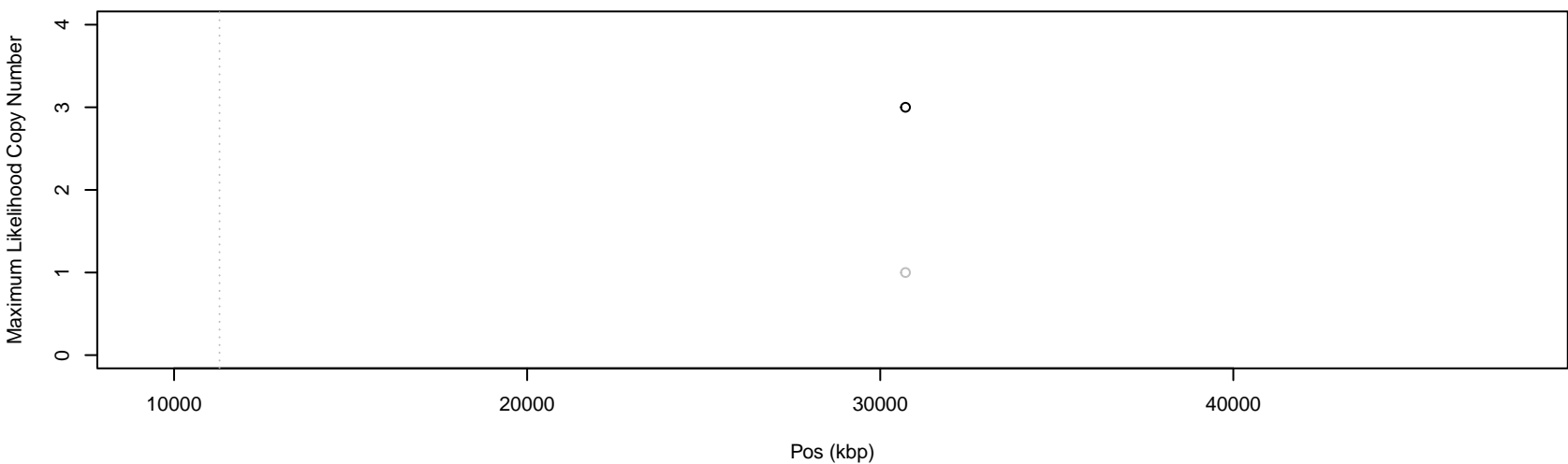
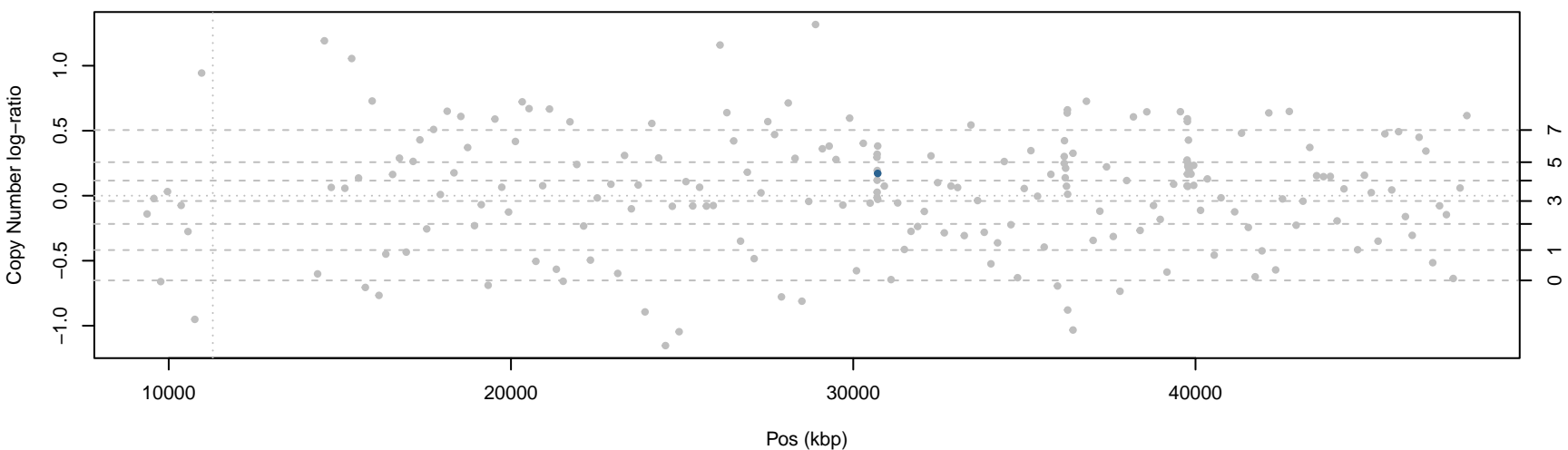
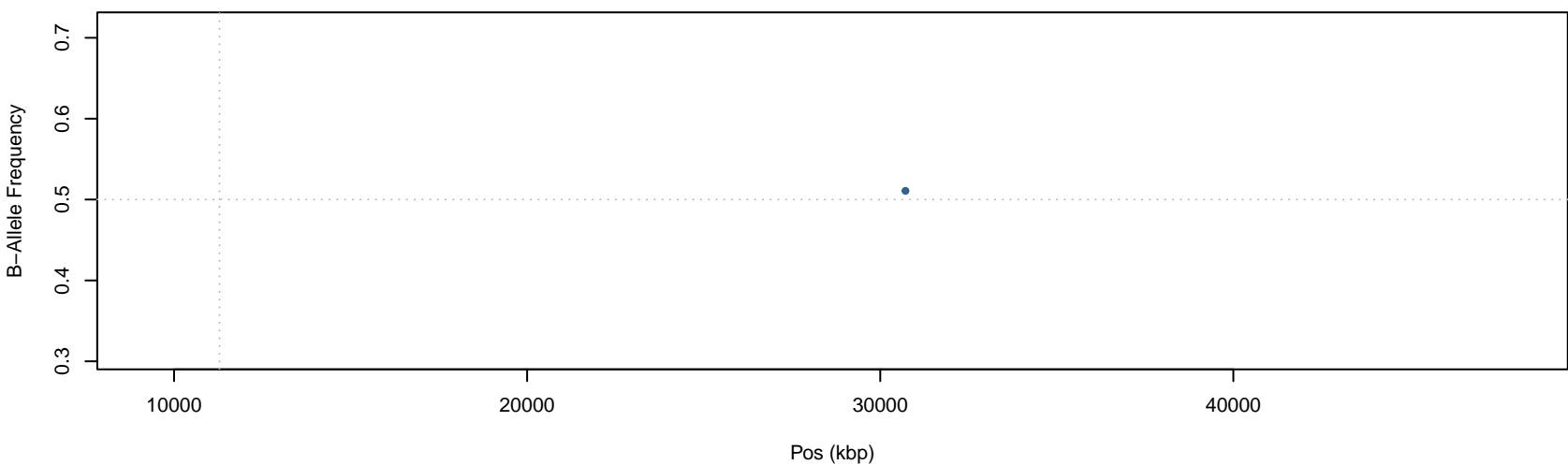
Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 19



Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 20



Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 21



Purity: 0.26 Tumor ploidy: 3.25 SNV log-likelihood: -52.12 GoF: 95.4% Mean coverage: 486;543 Chromosome: 22

