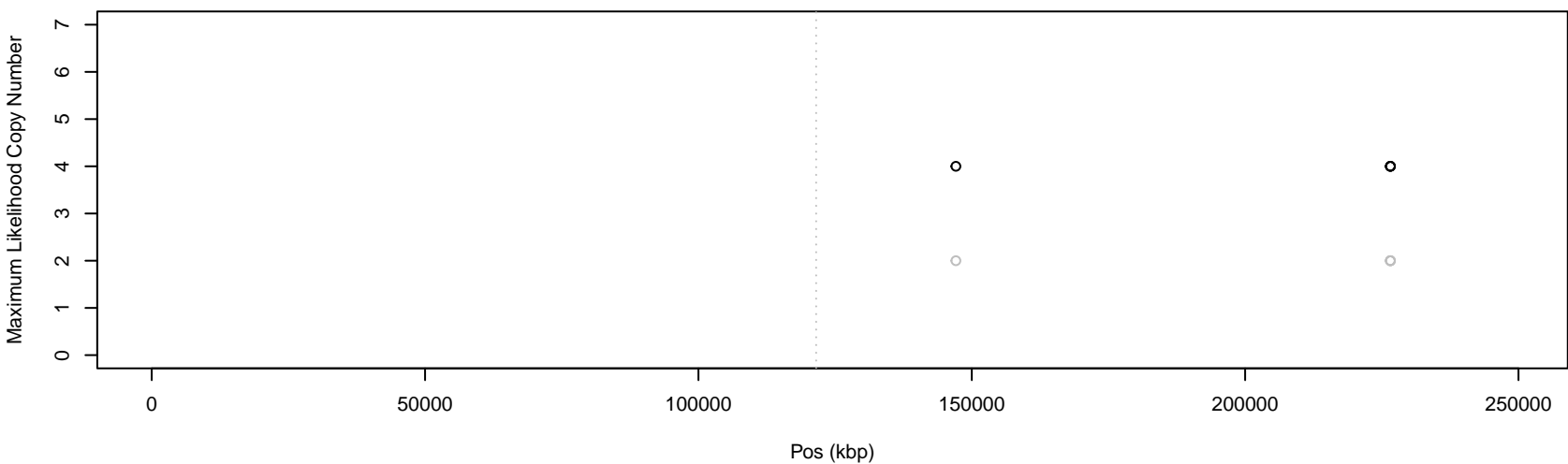
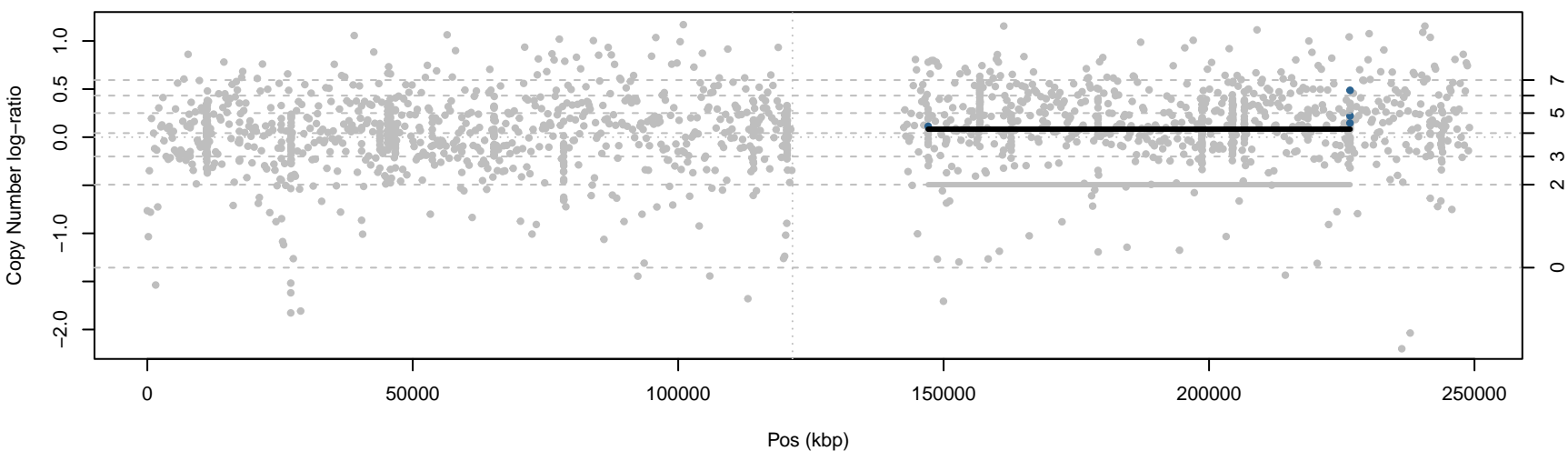
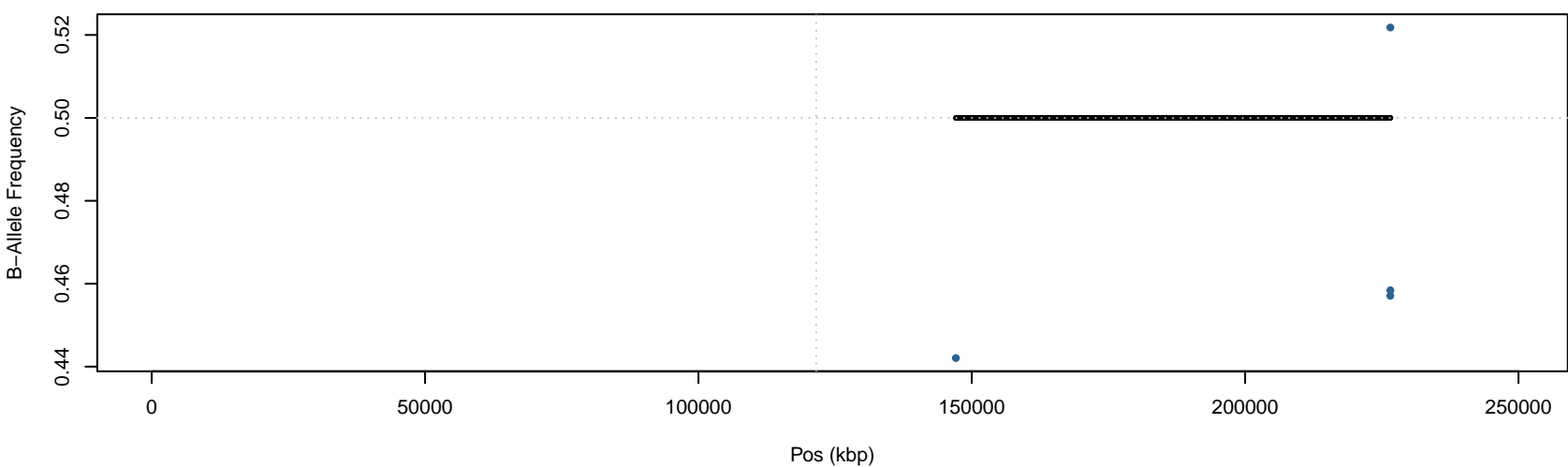
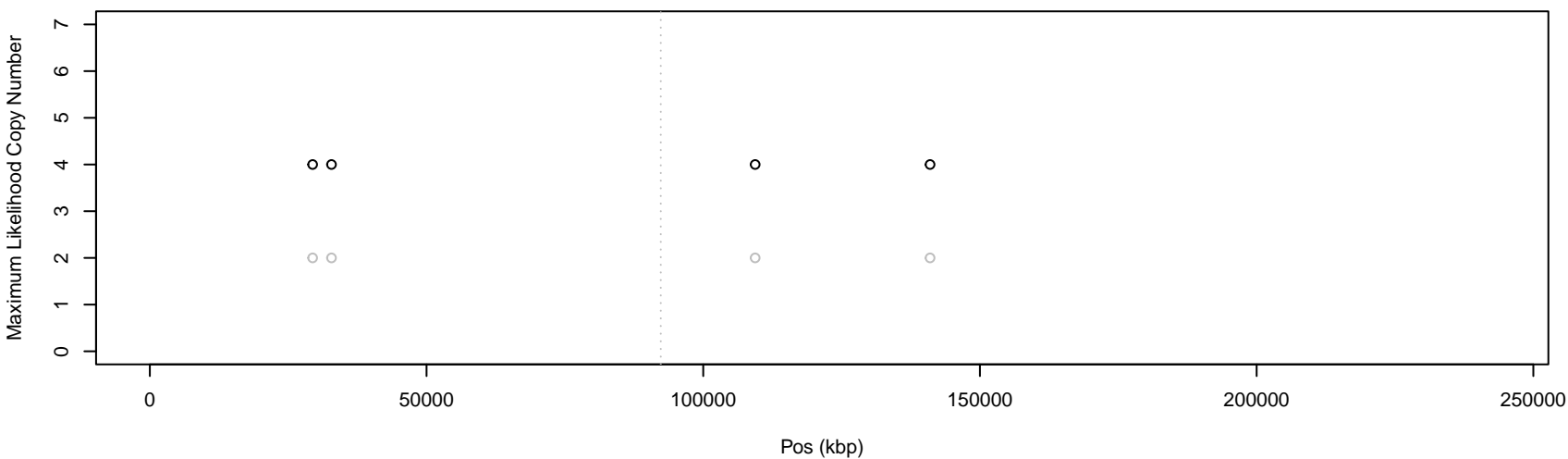
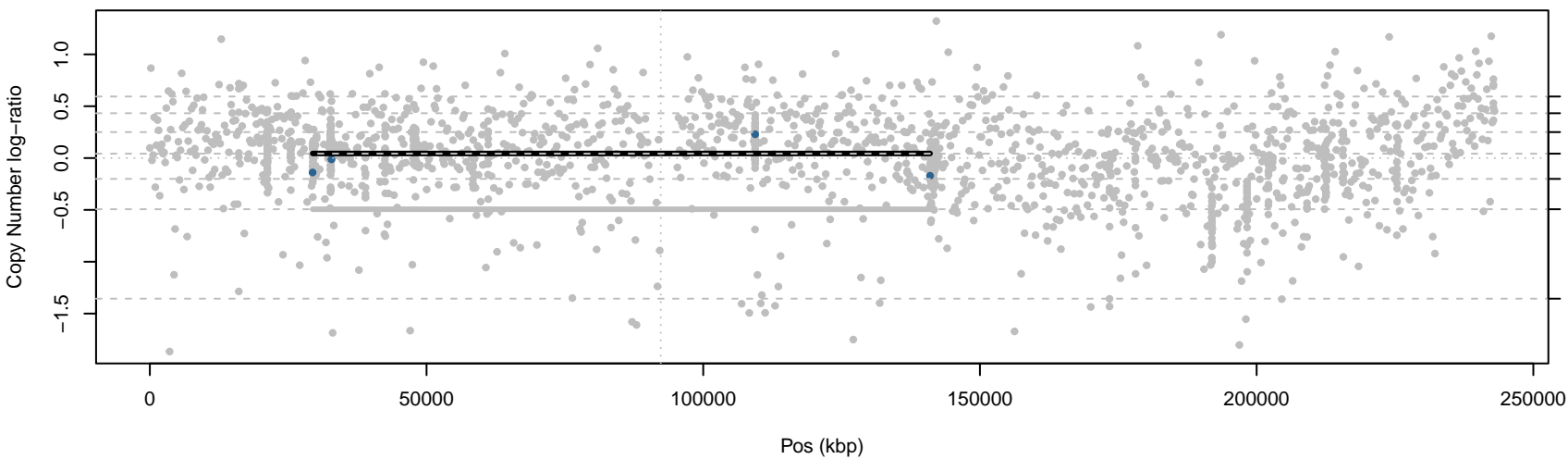
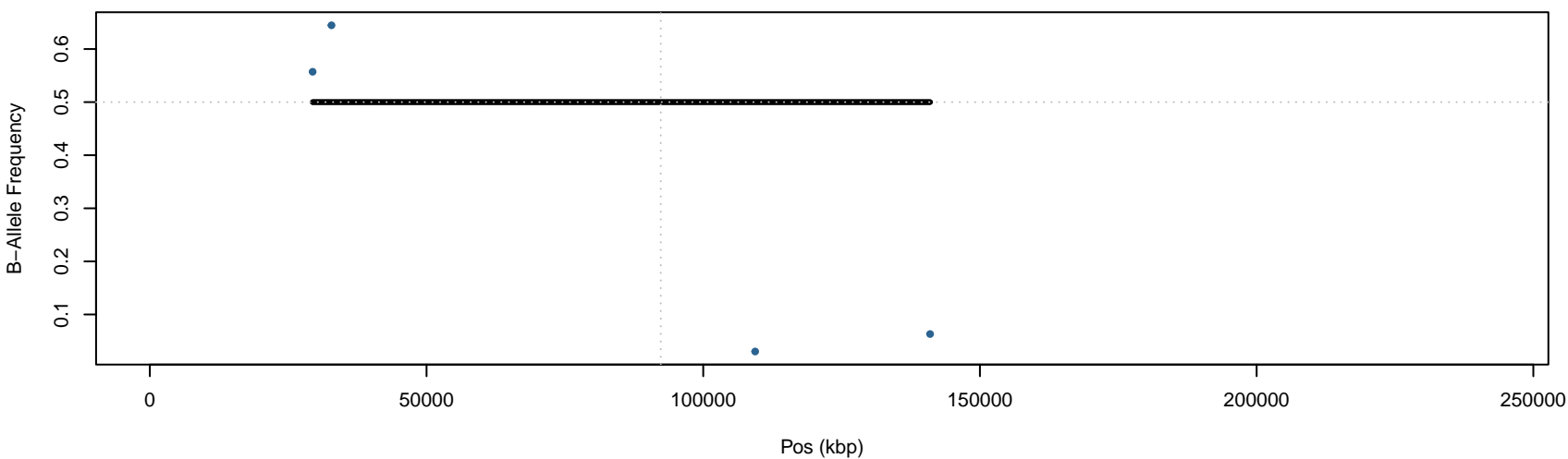


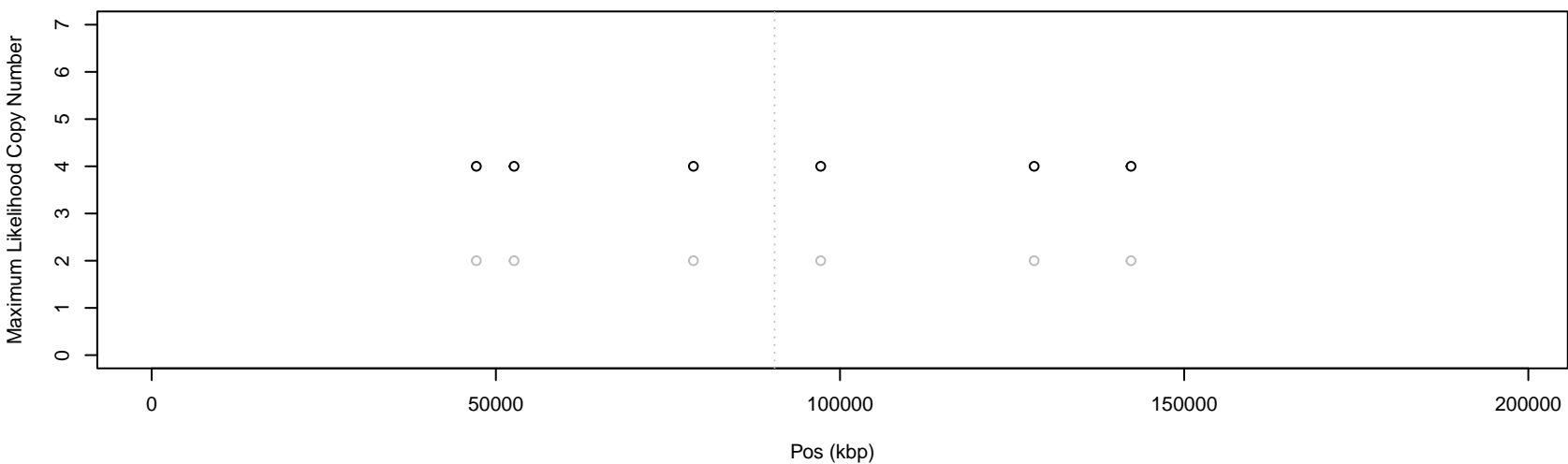
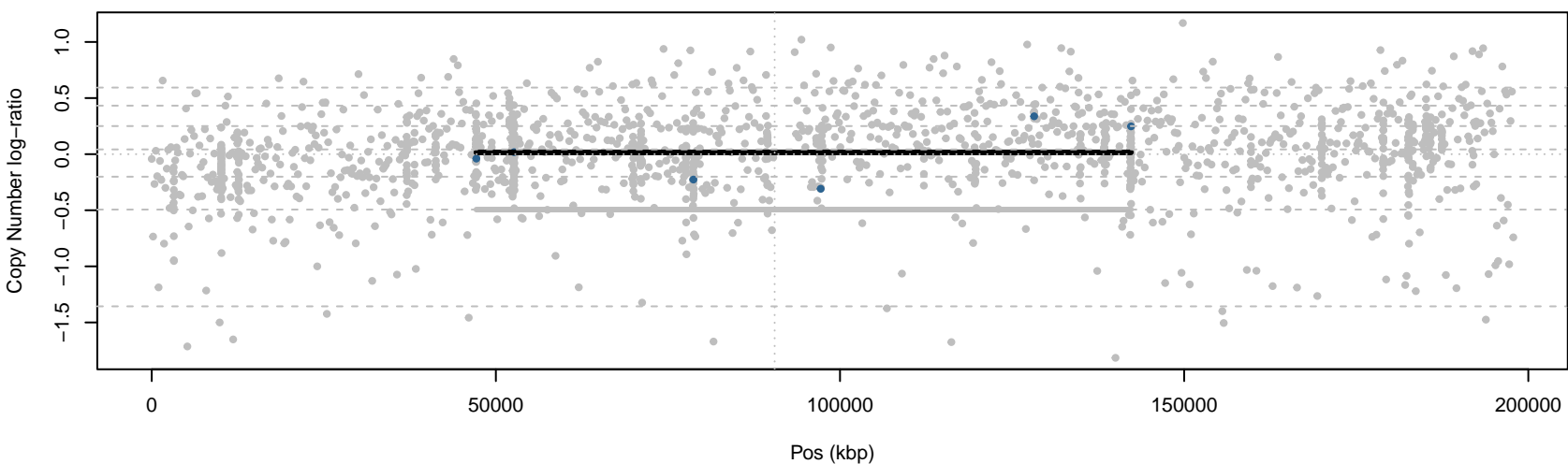
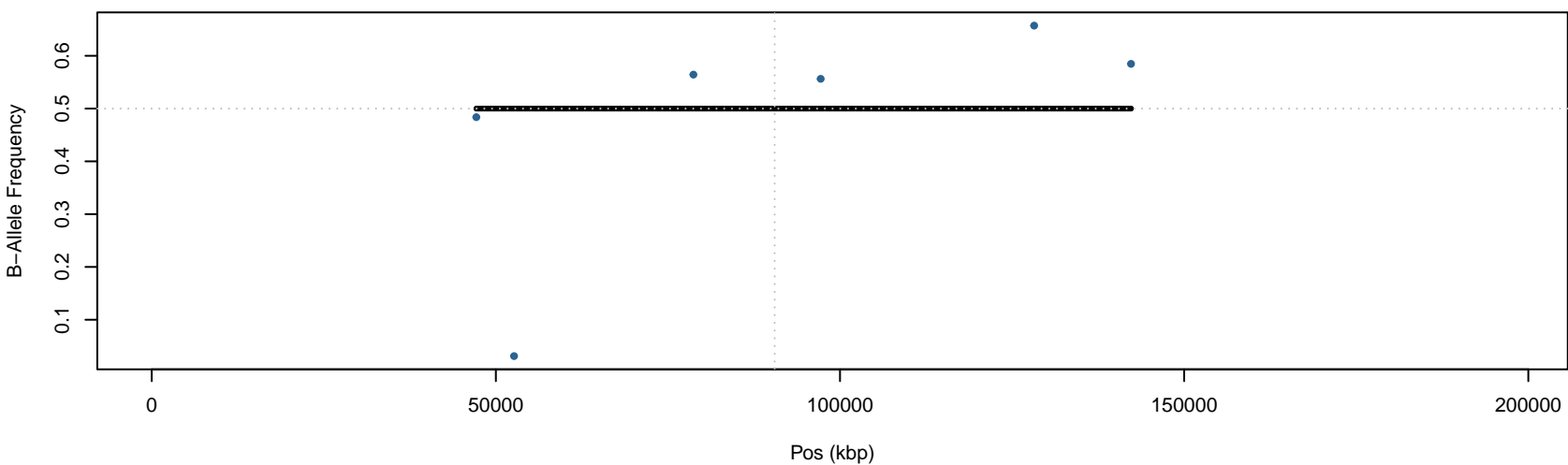
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 1



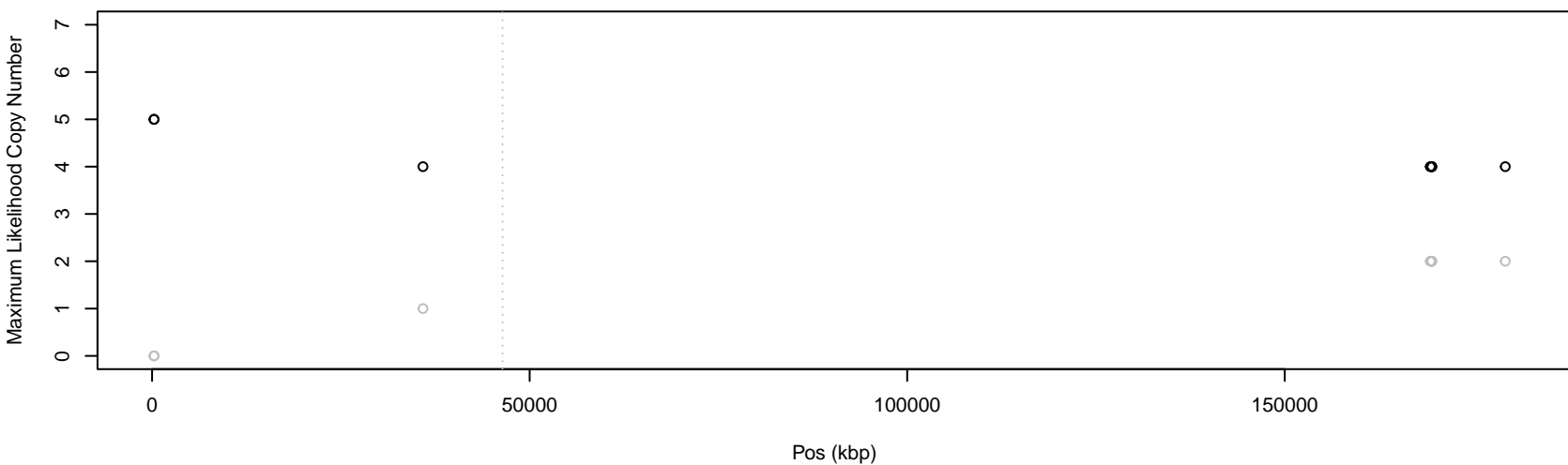
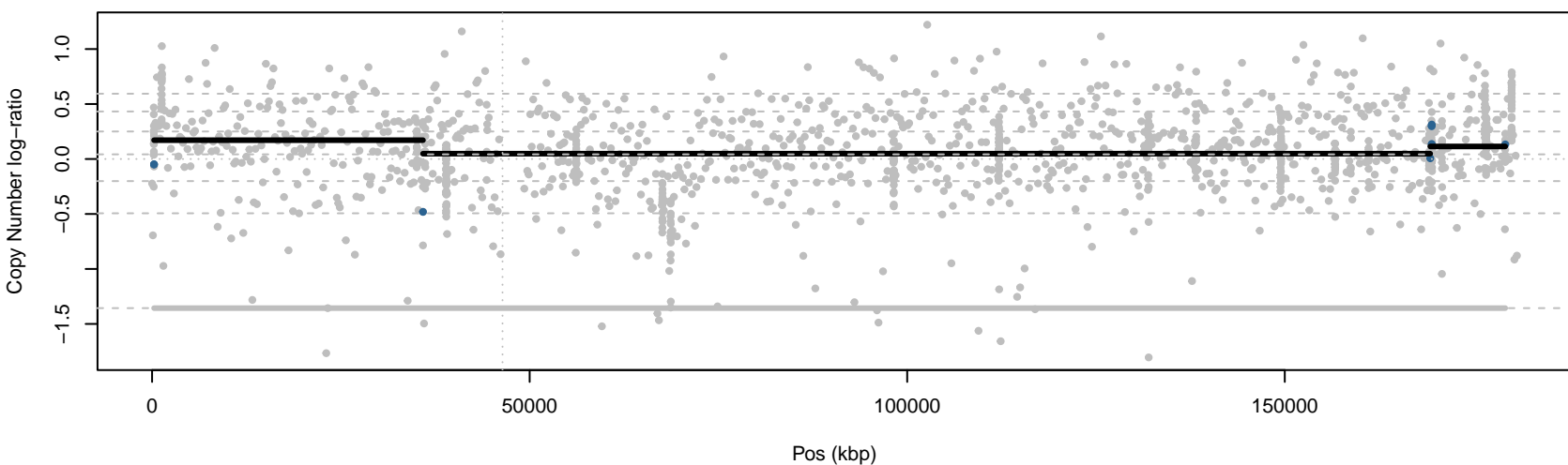
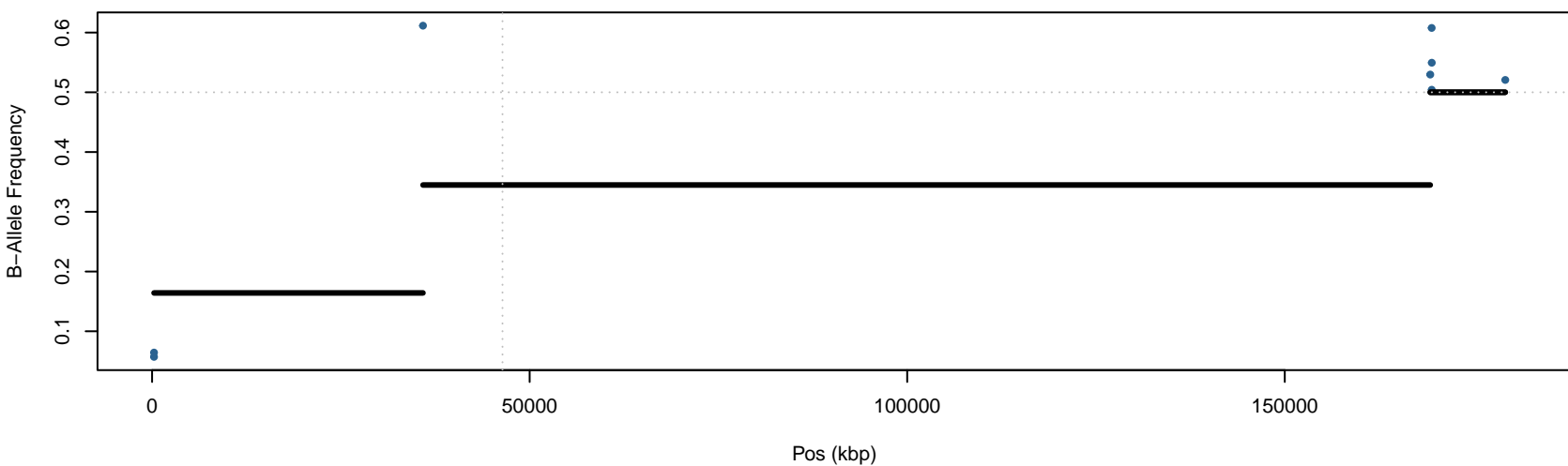
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 2



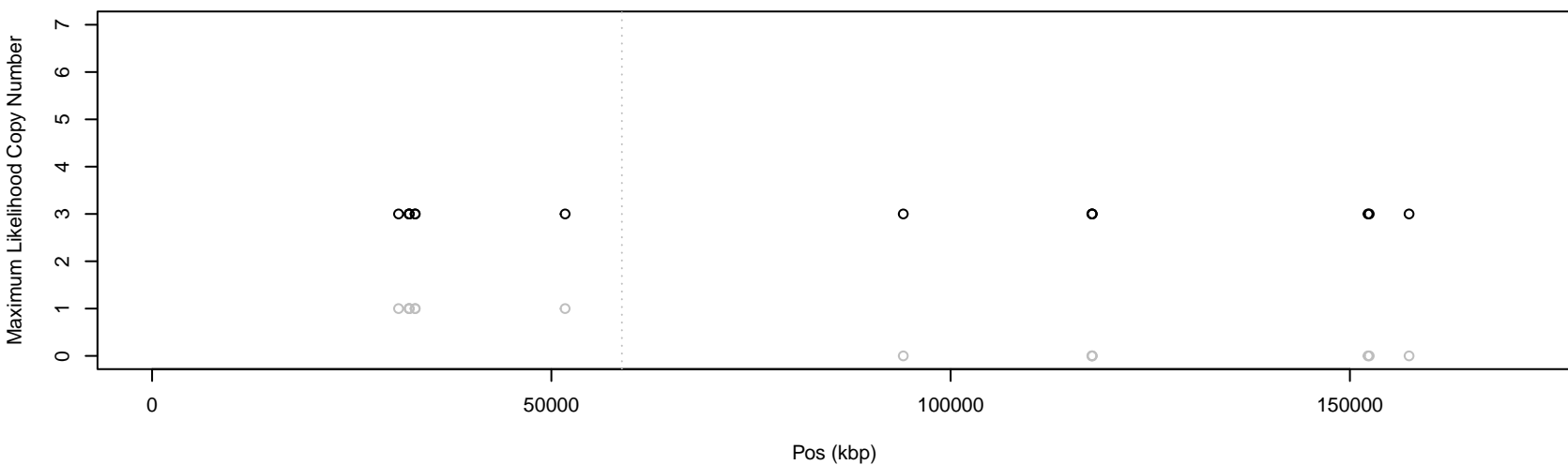
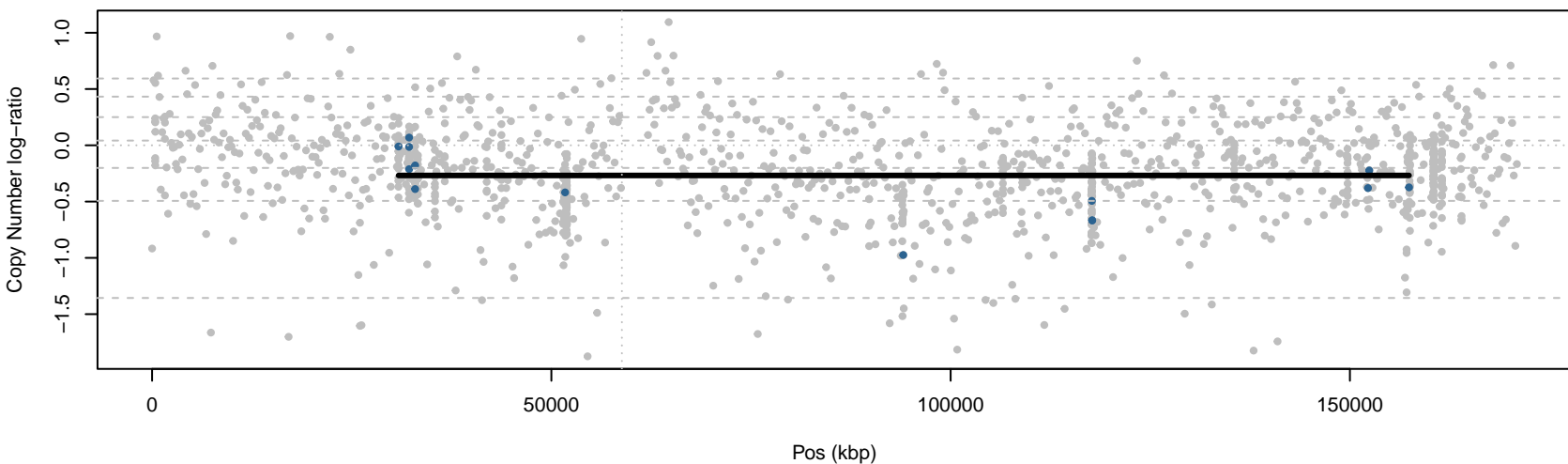
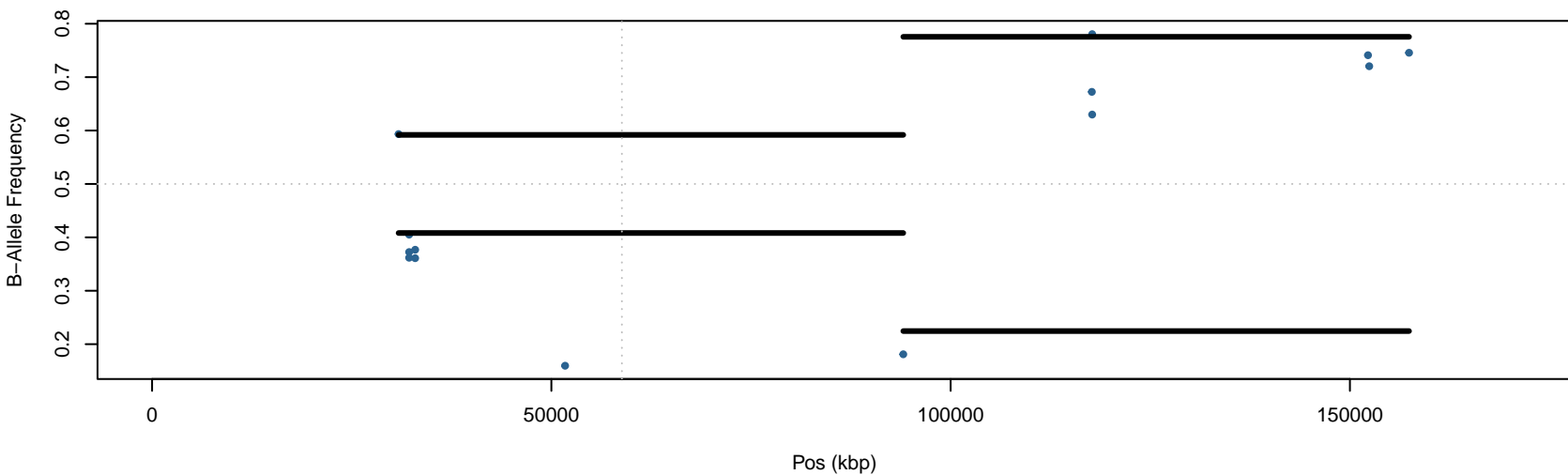
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 3



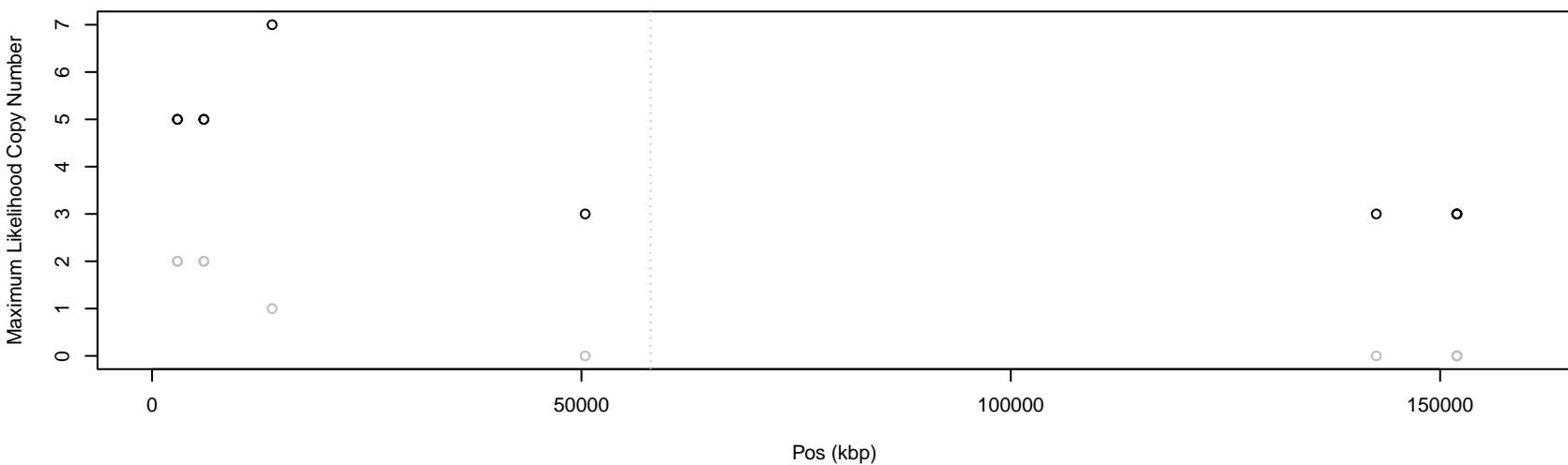
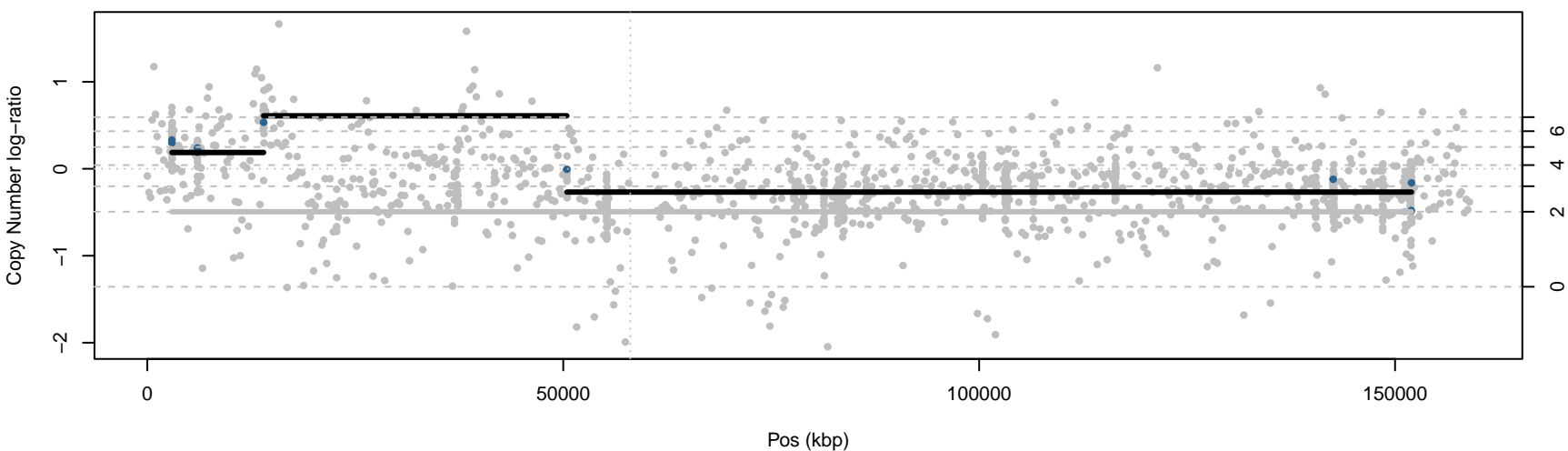
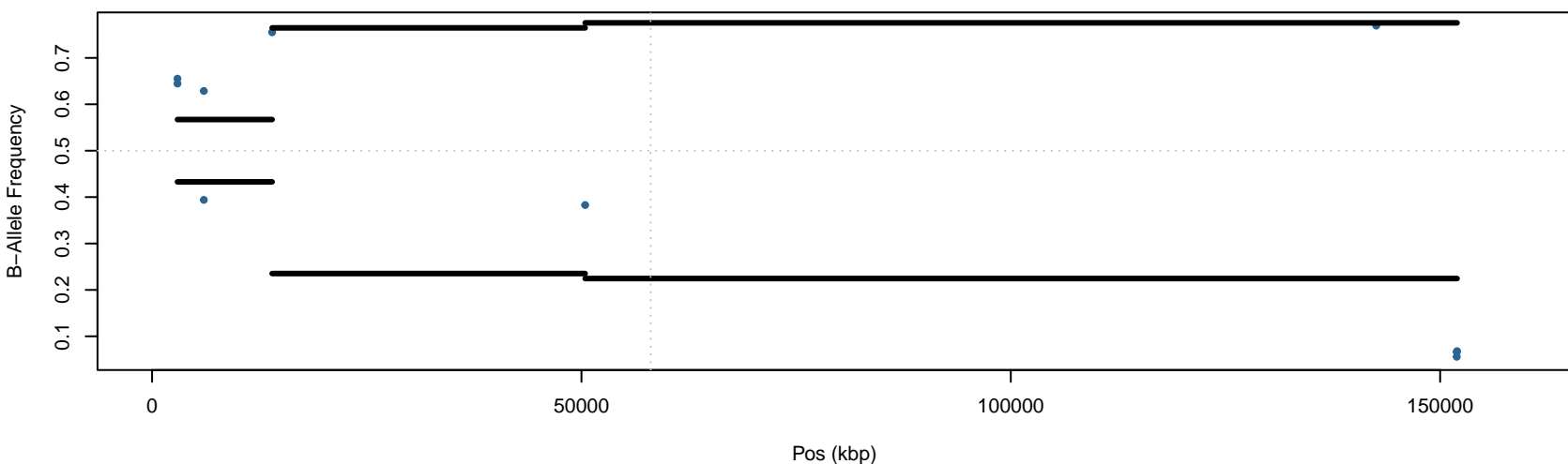
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 5



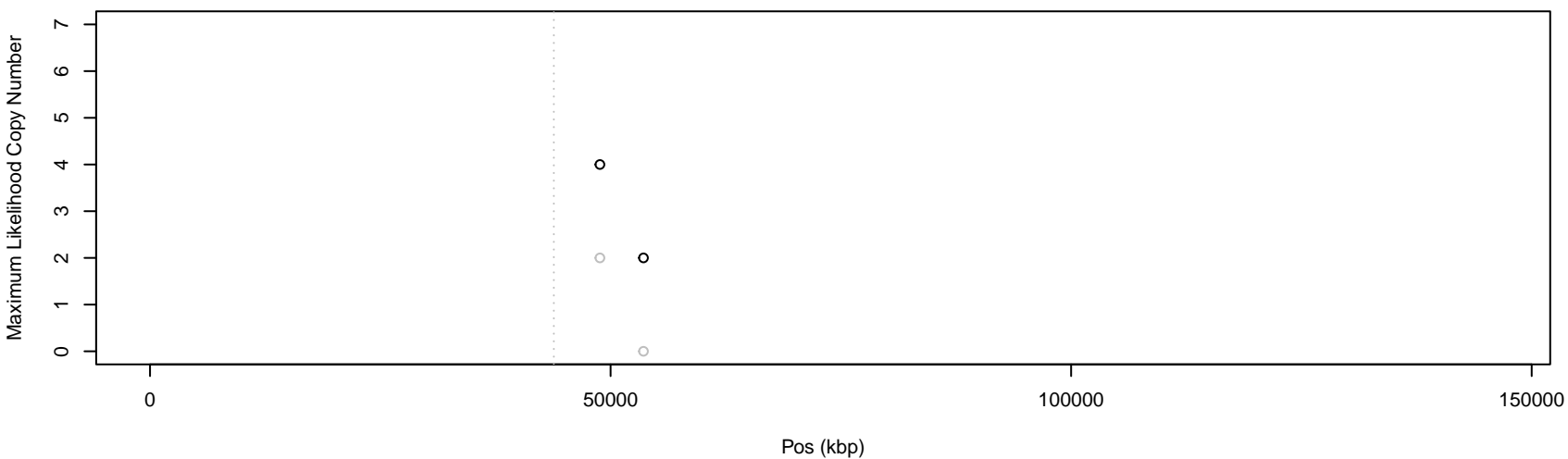
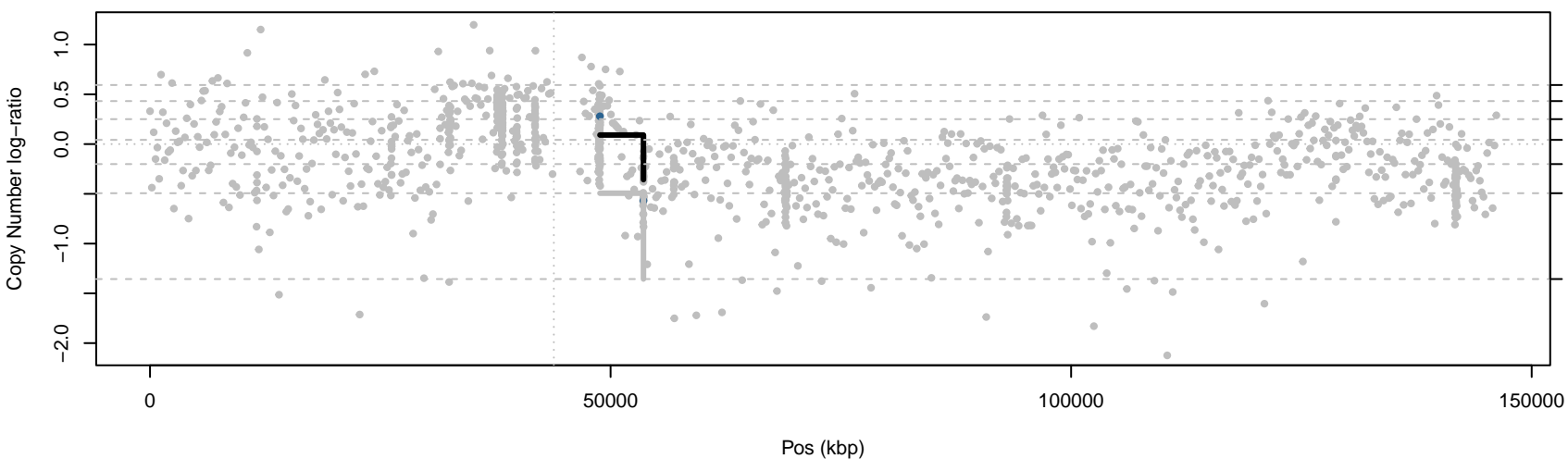
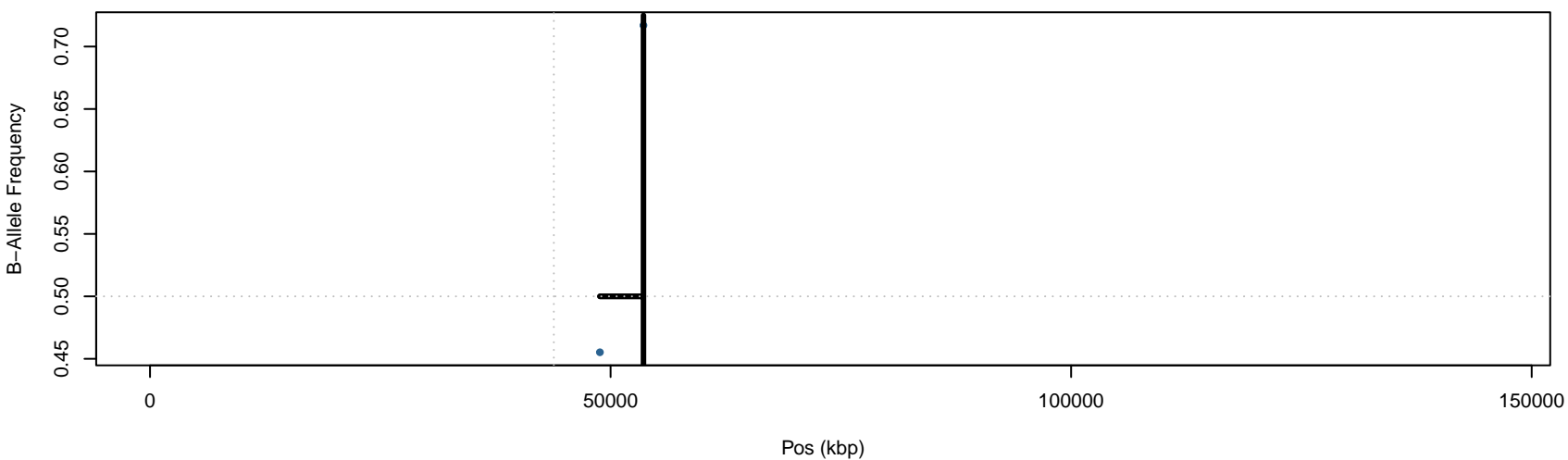
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 6



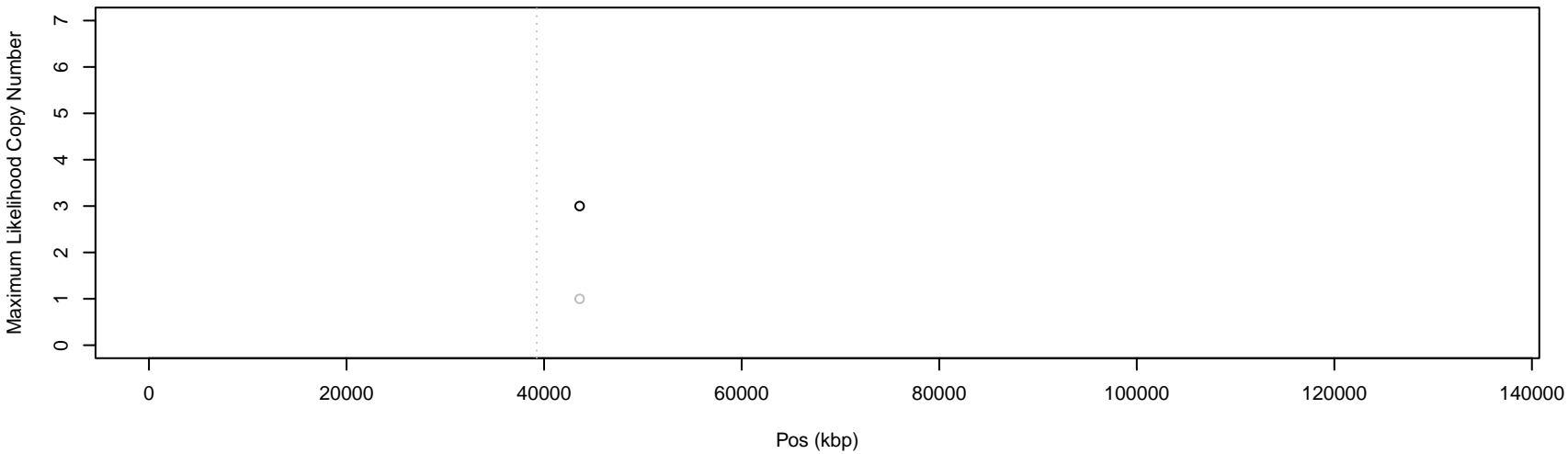
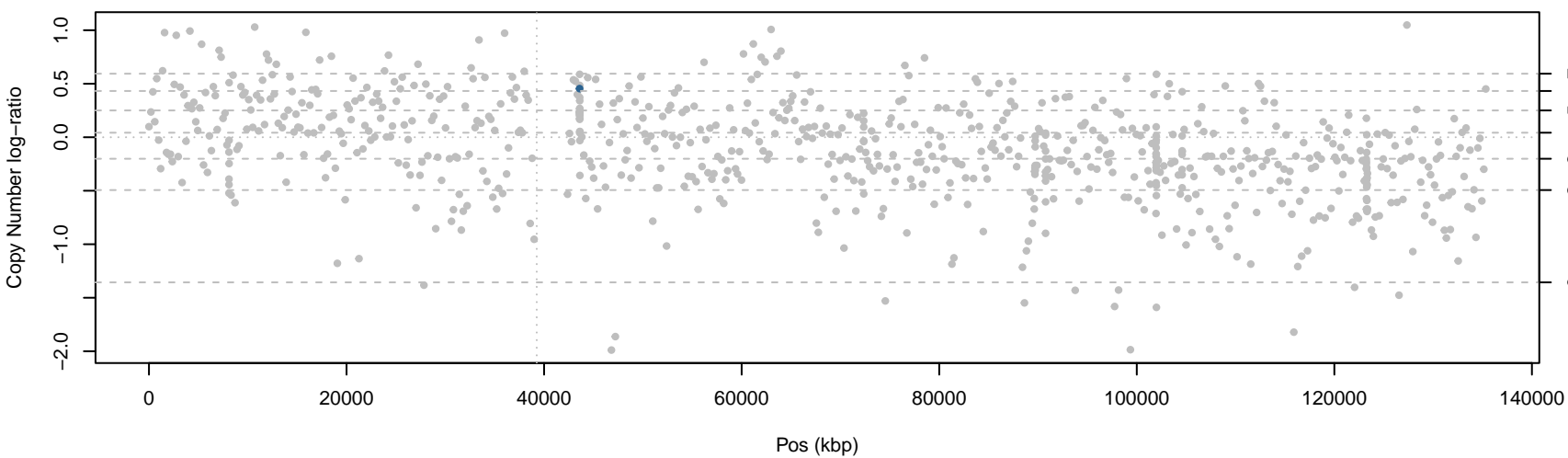
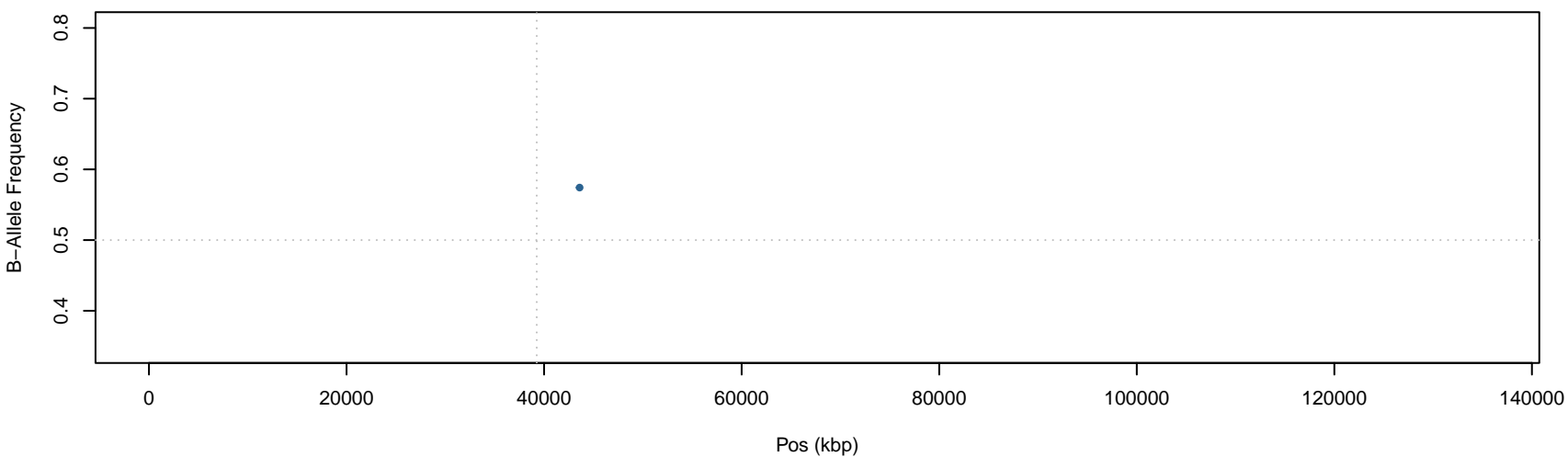
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 7



Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 8

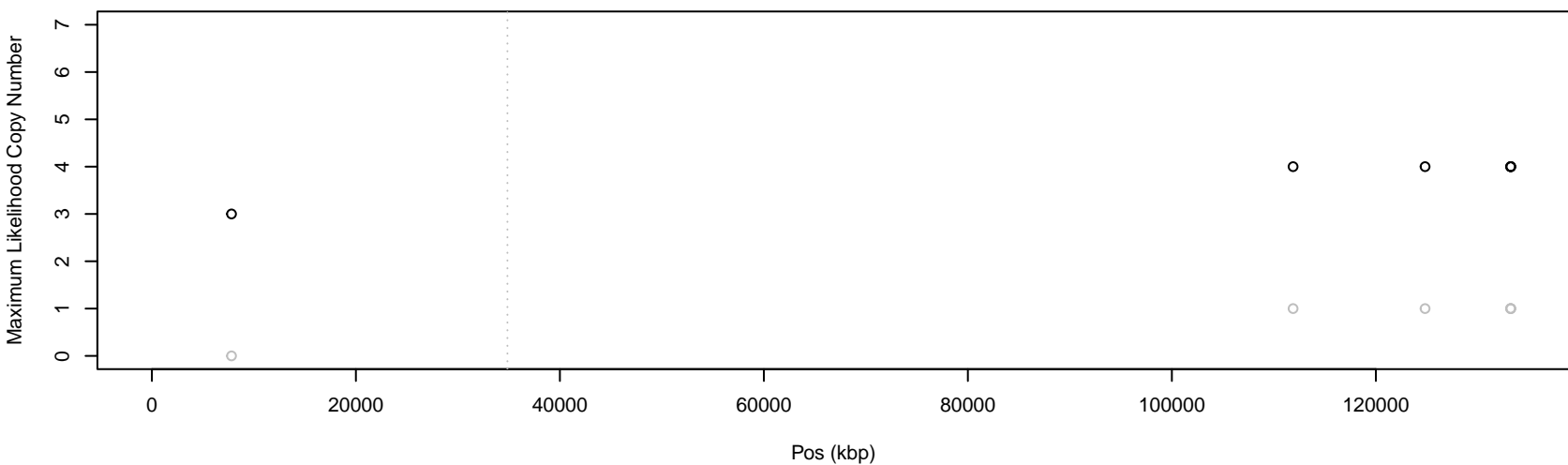
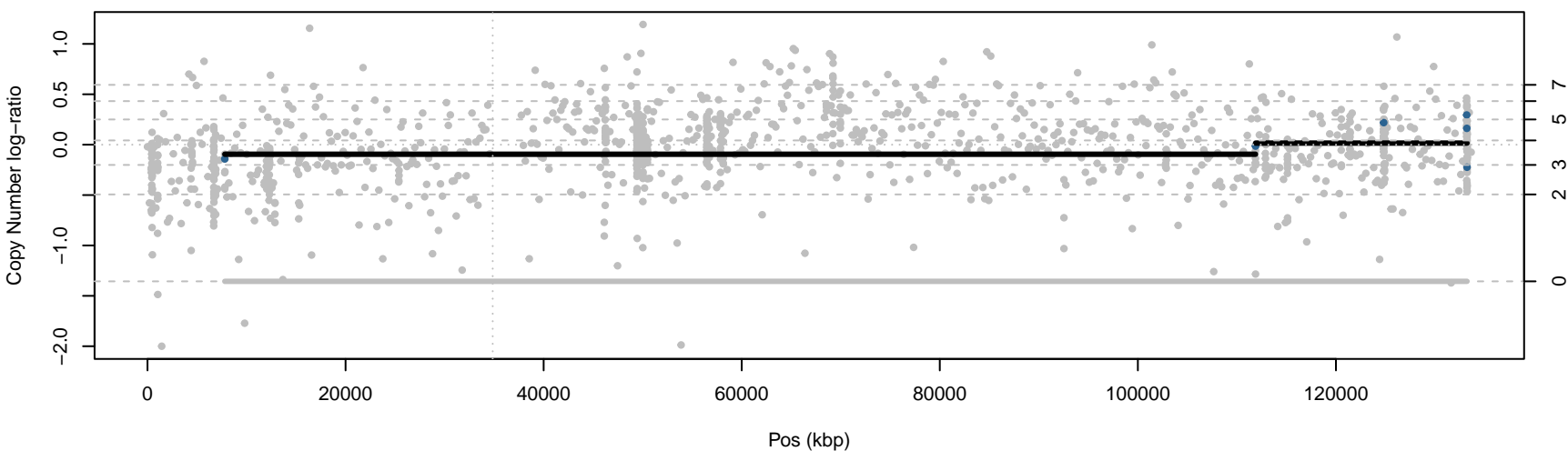
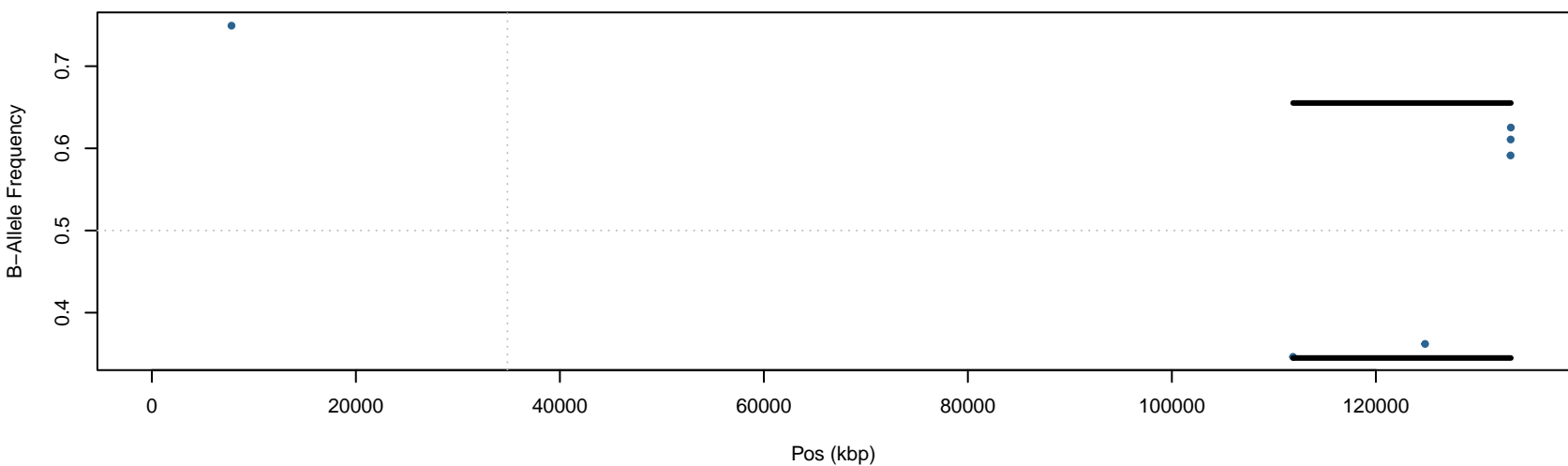


Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 10

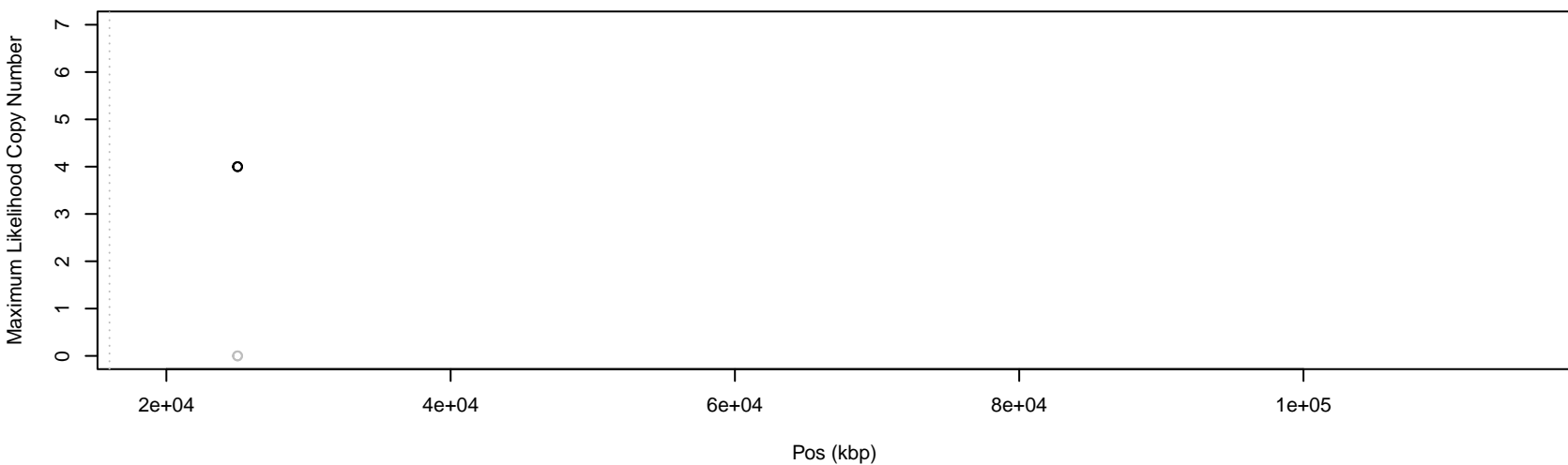
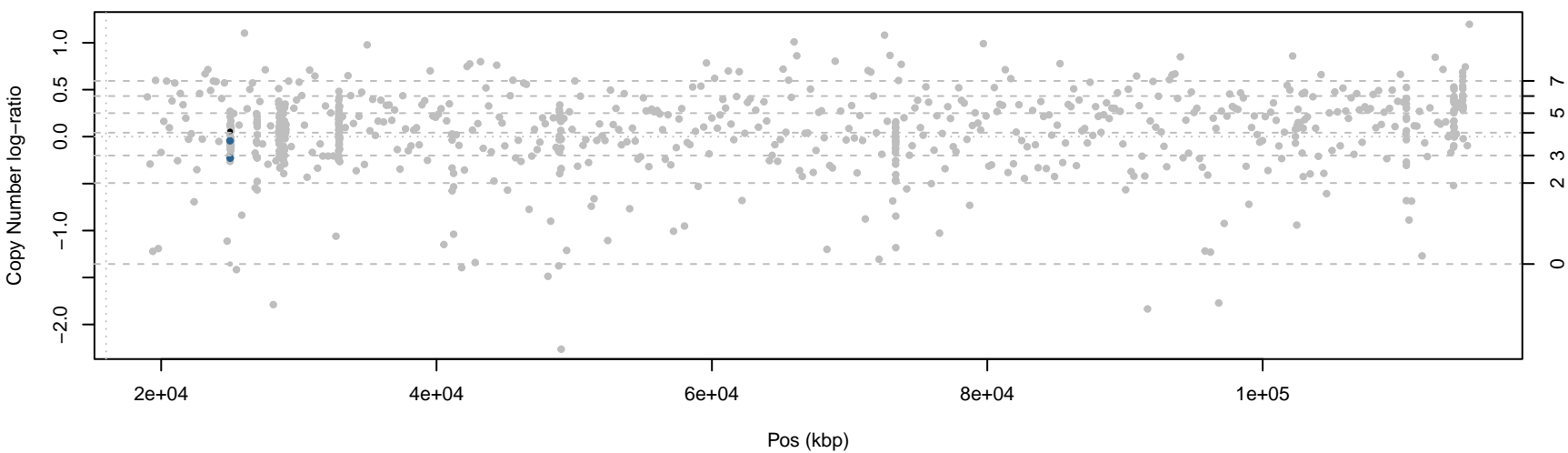
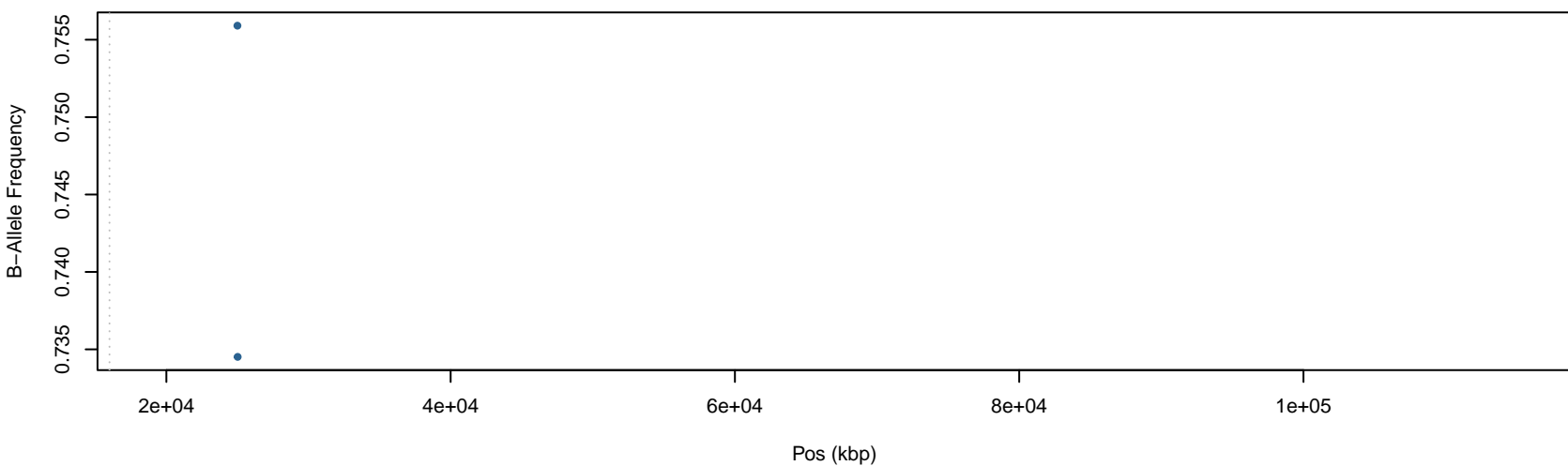




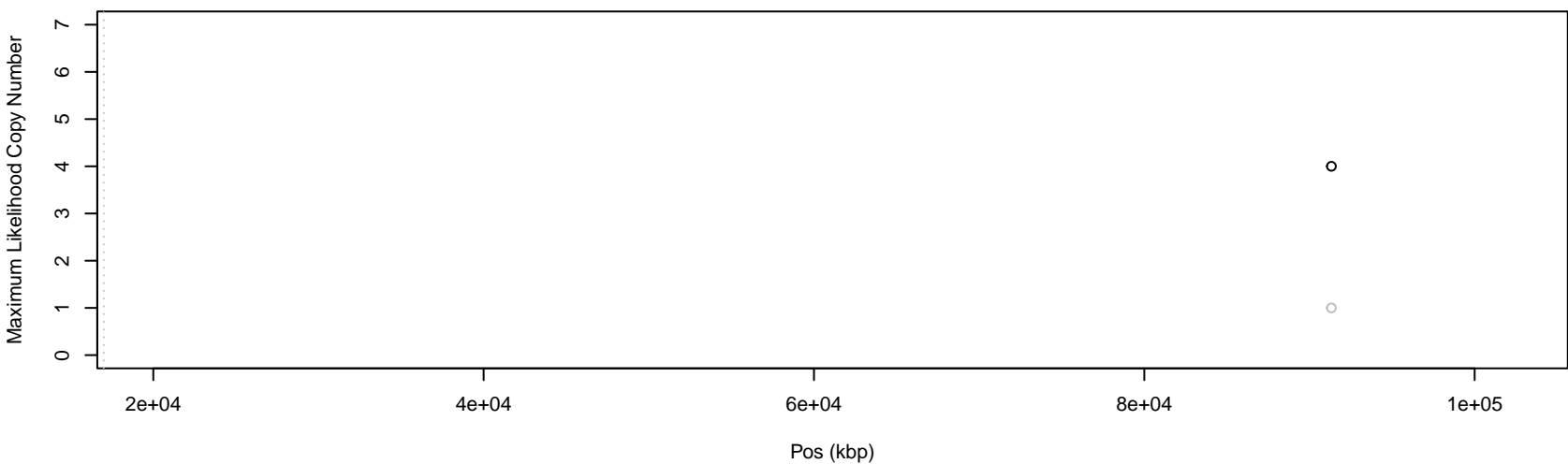
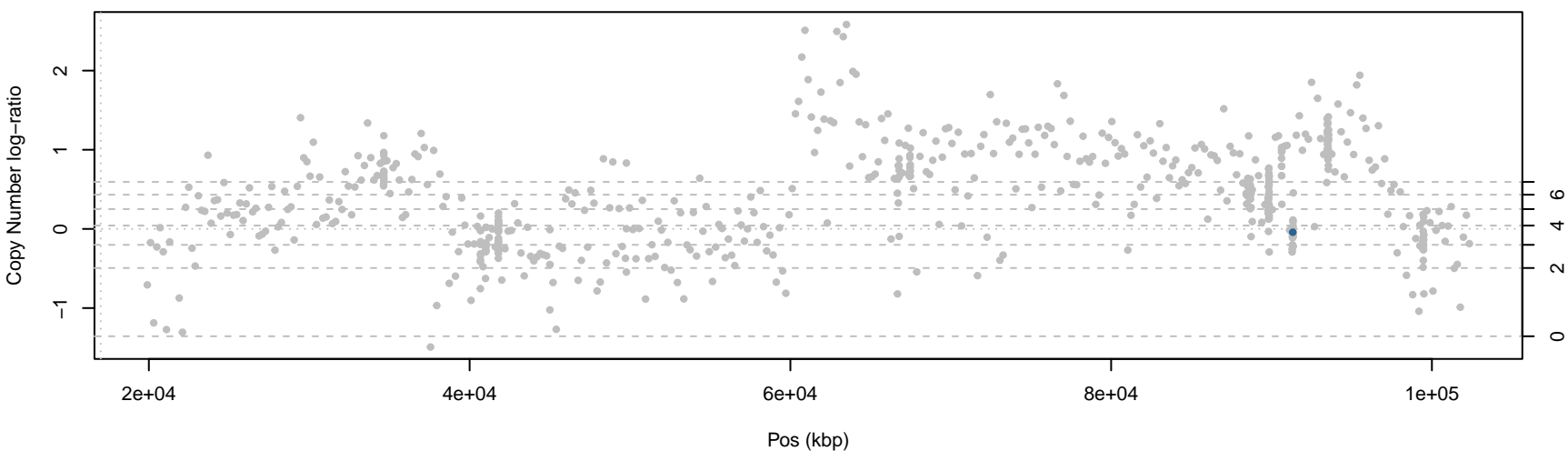
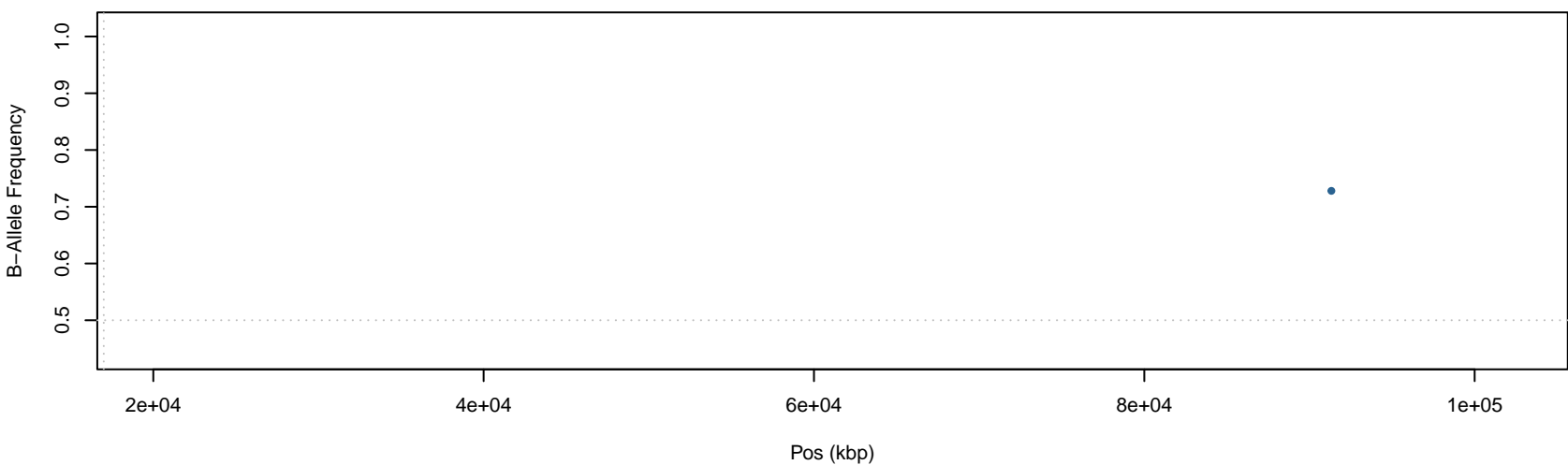
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 12



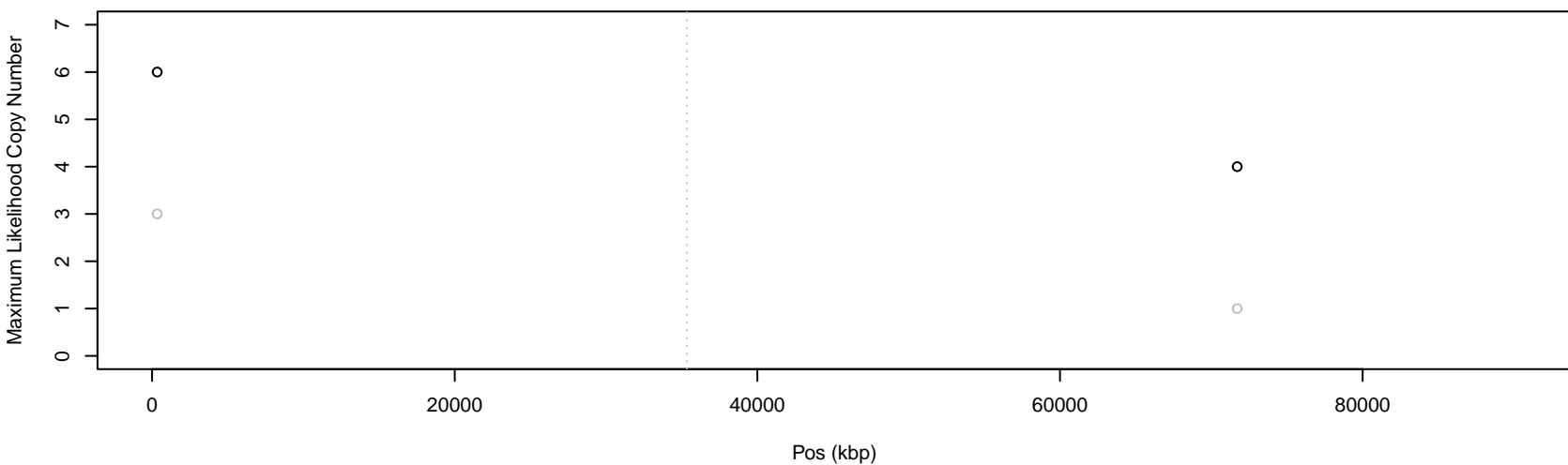
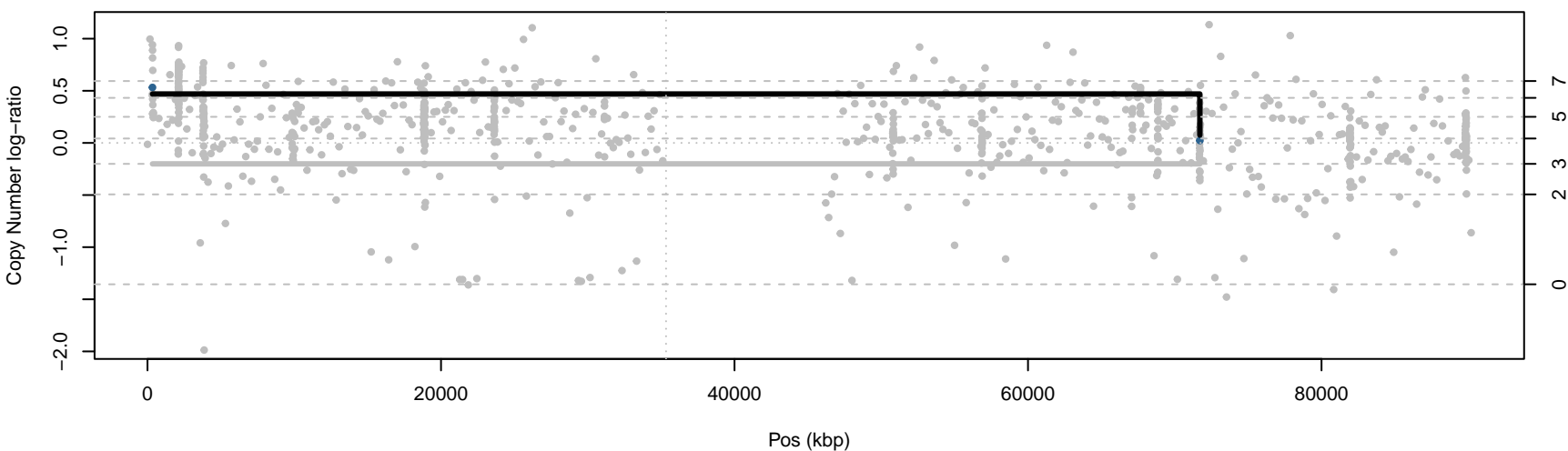
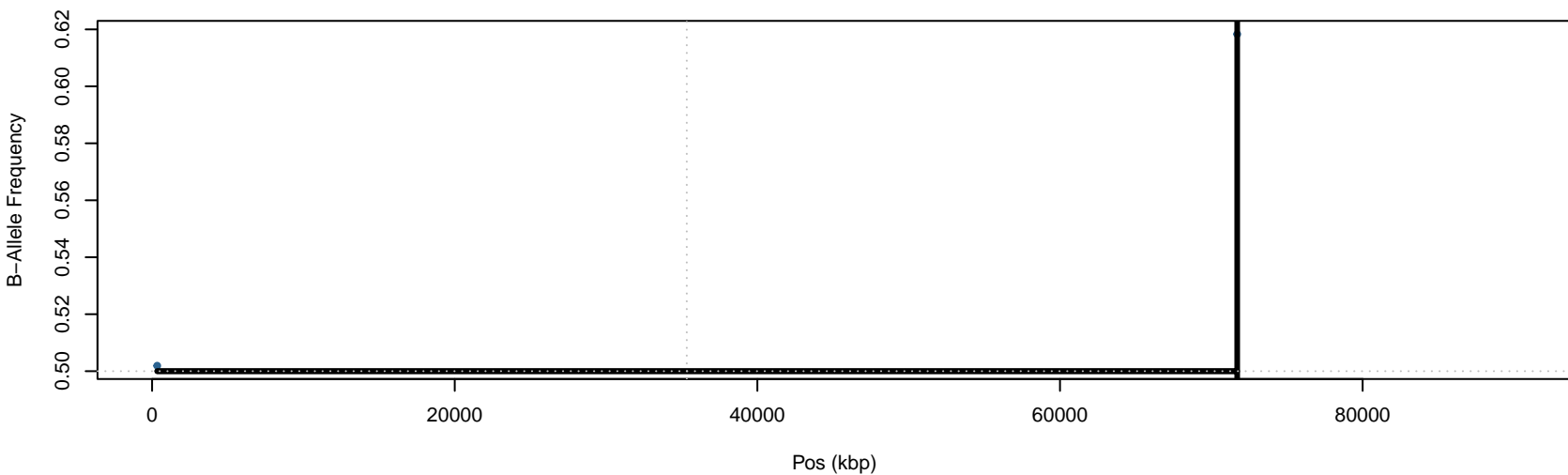
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 13



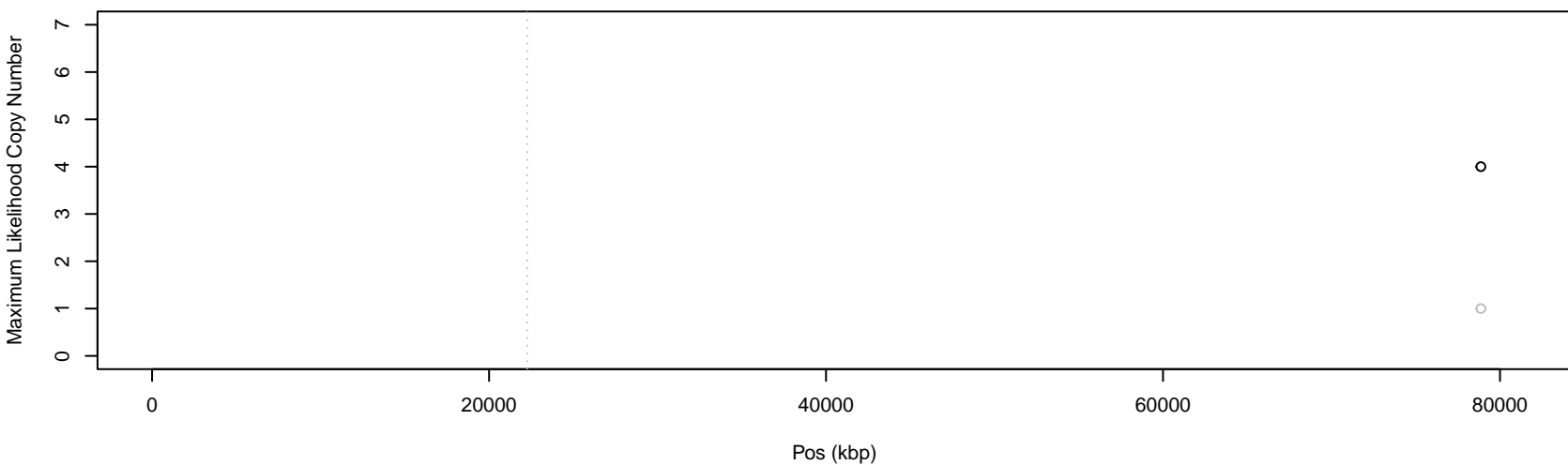
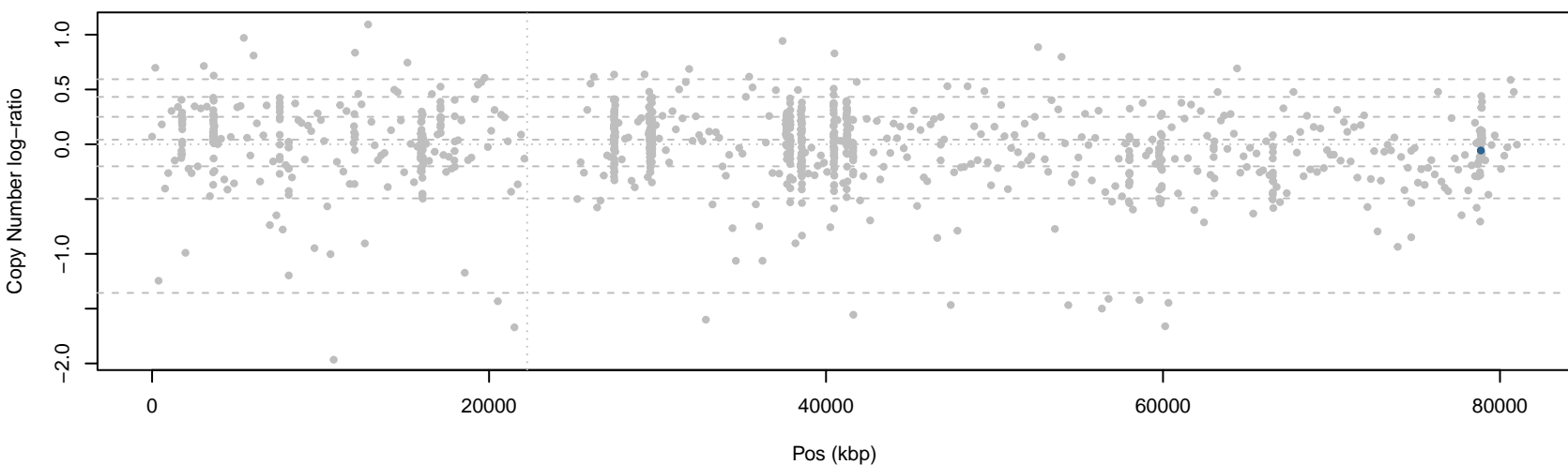
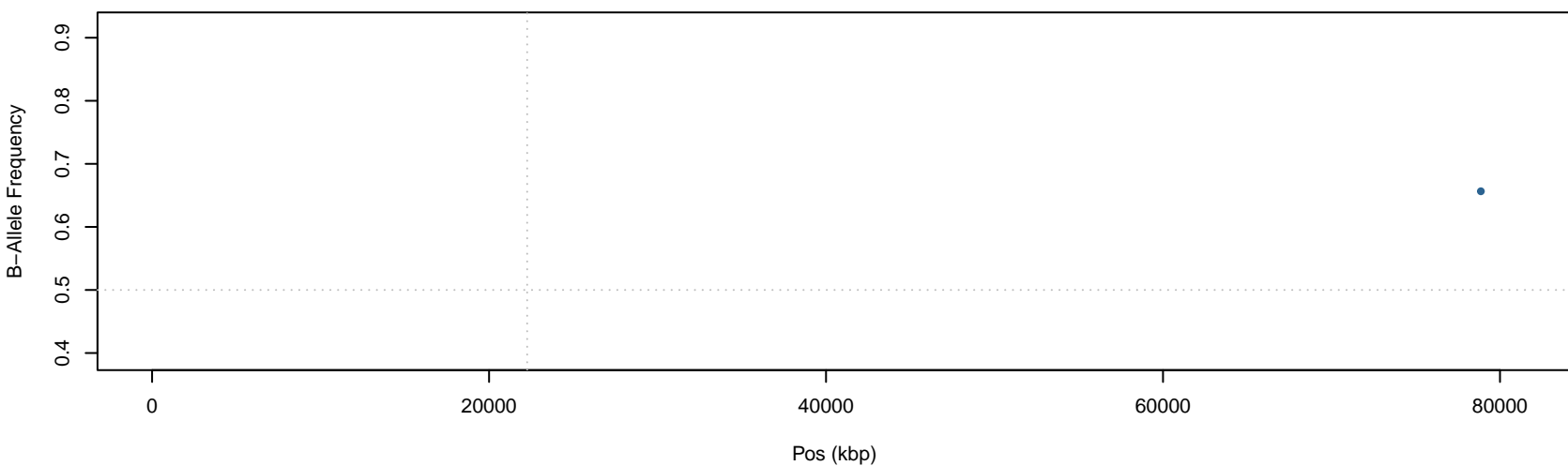
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 15



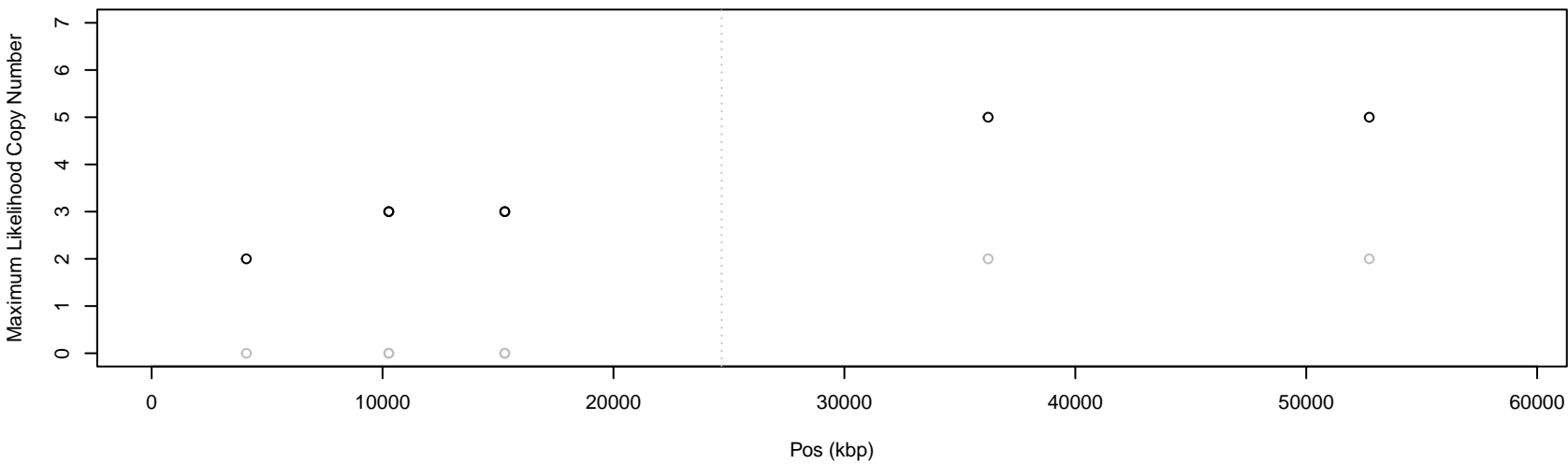
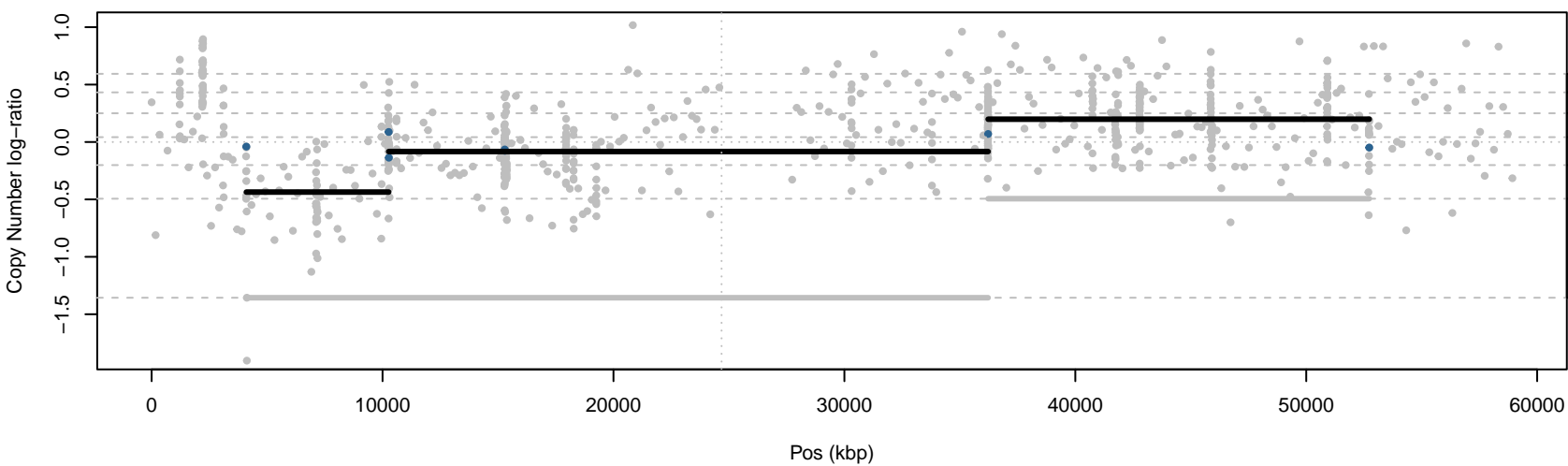
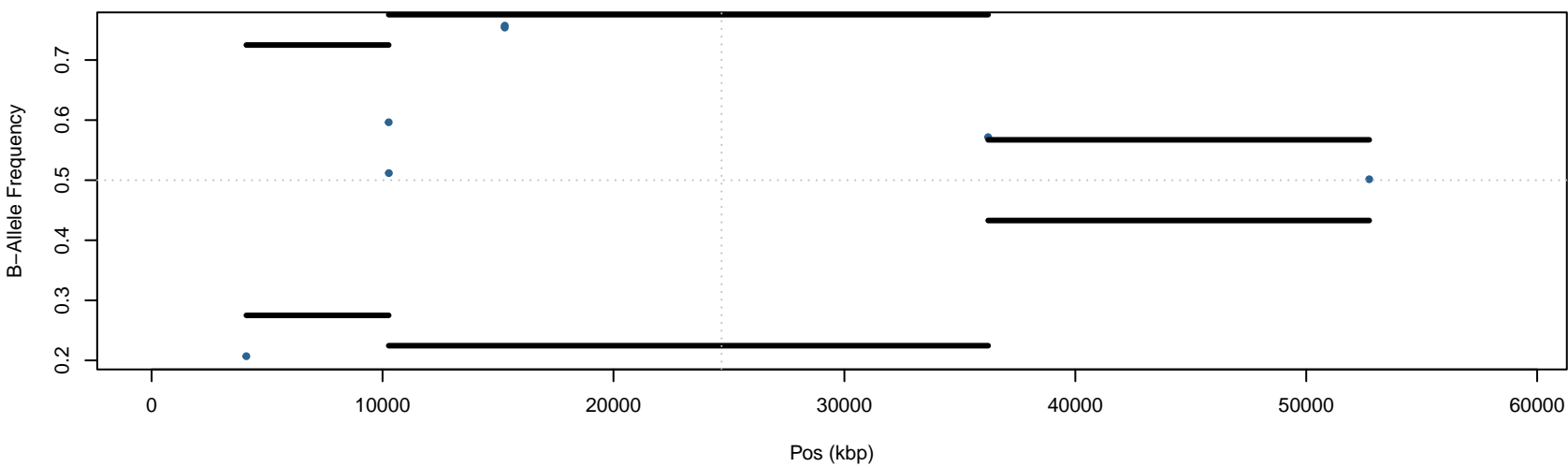
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 16



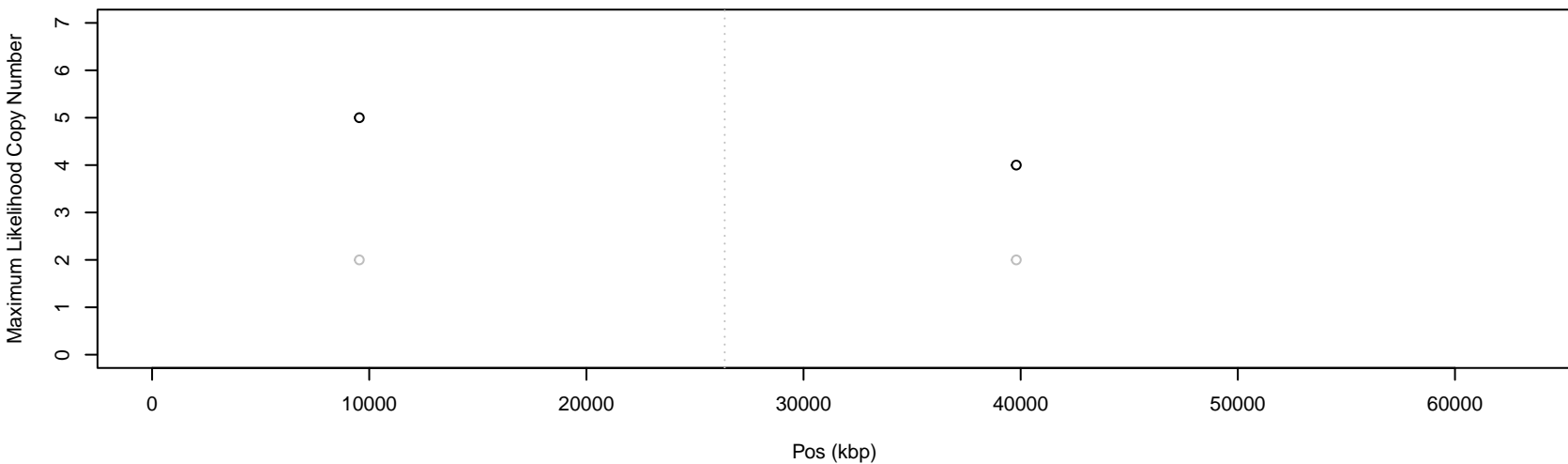
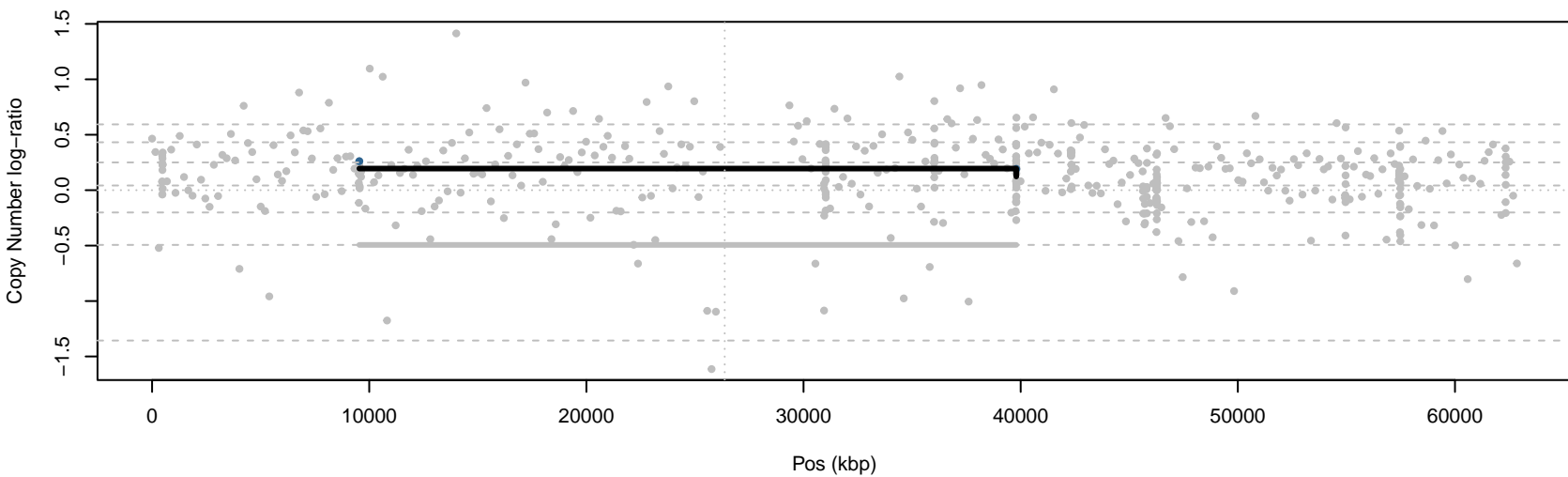
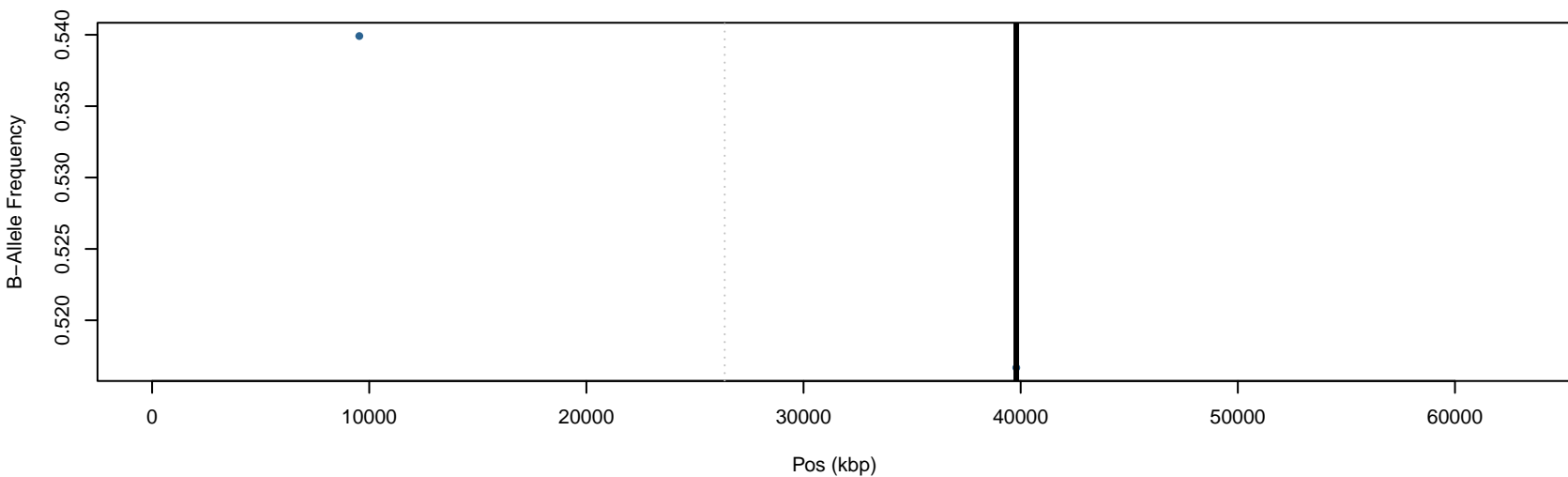
Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 17



Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 19



Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 20



Purity: 0.45 Tumor ploidy: 3.814 SNV log-likelihood: -99.22 GoF: 89.3% Mean coverage: 561;493 Chromosome: 22

