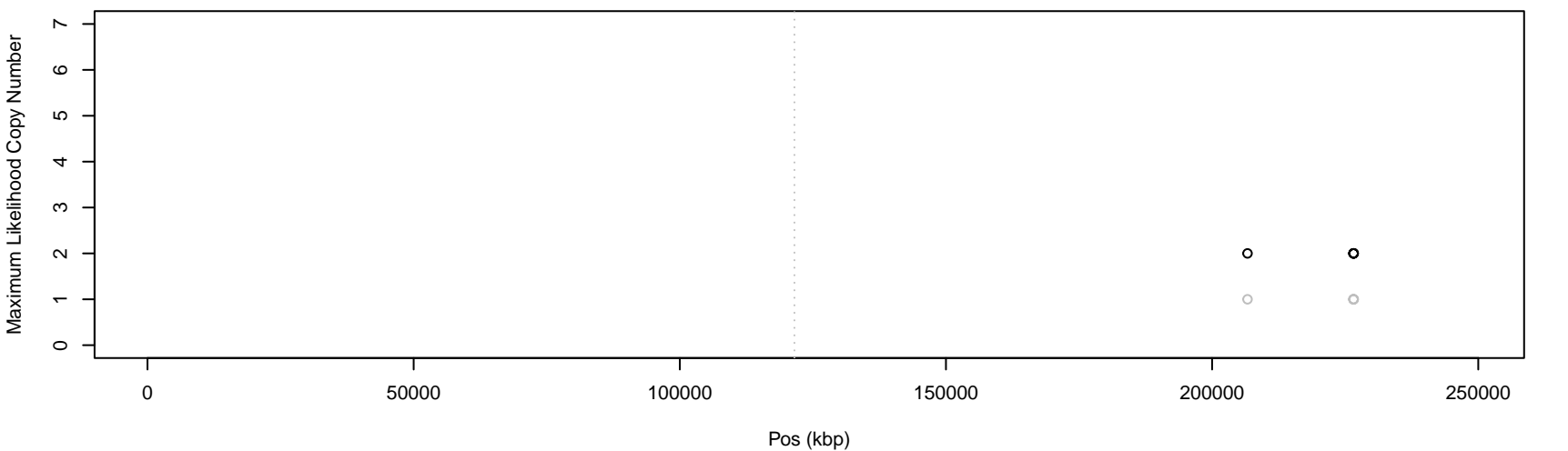
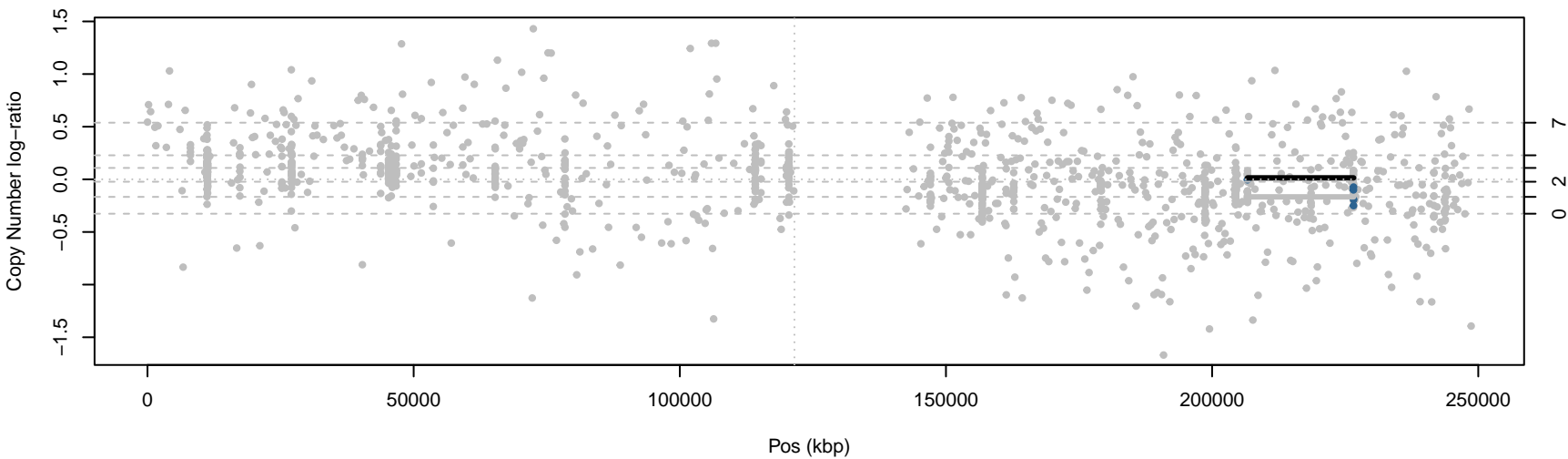
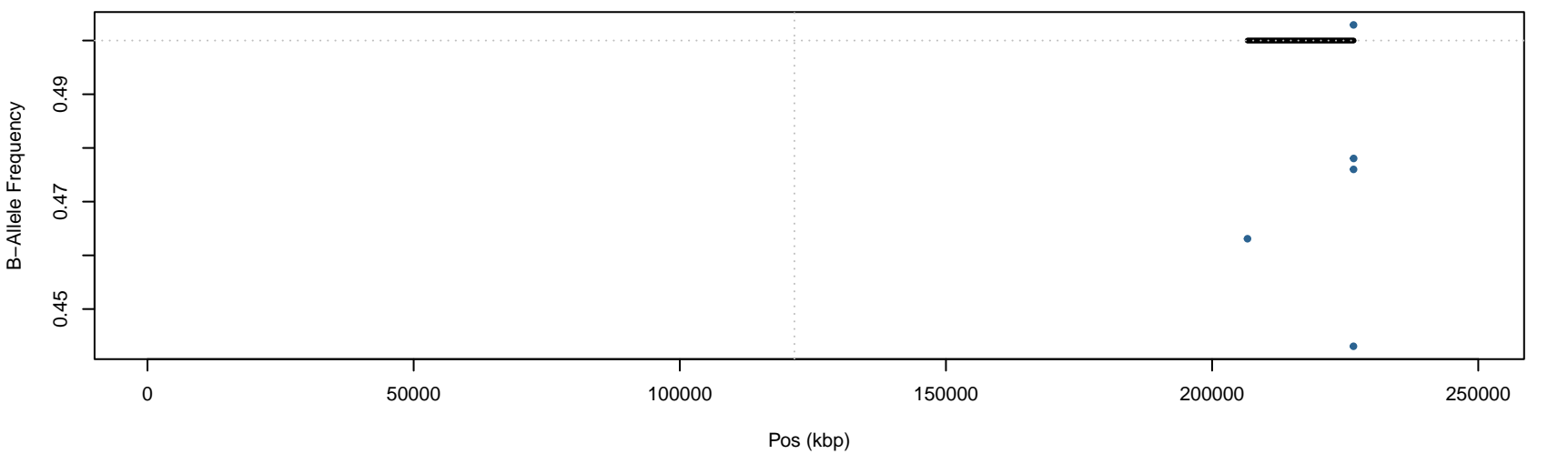
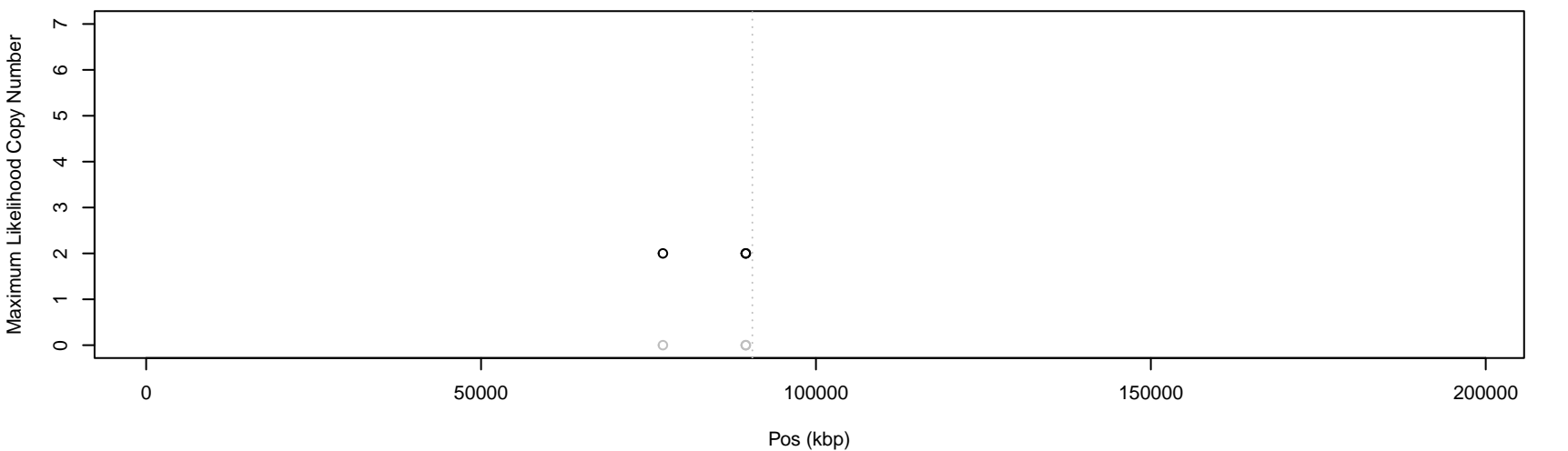
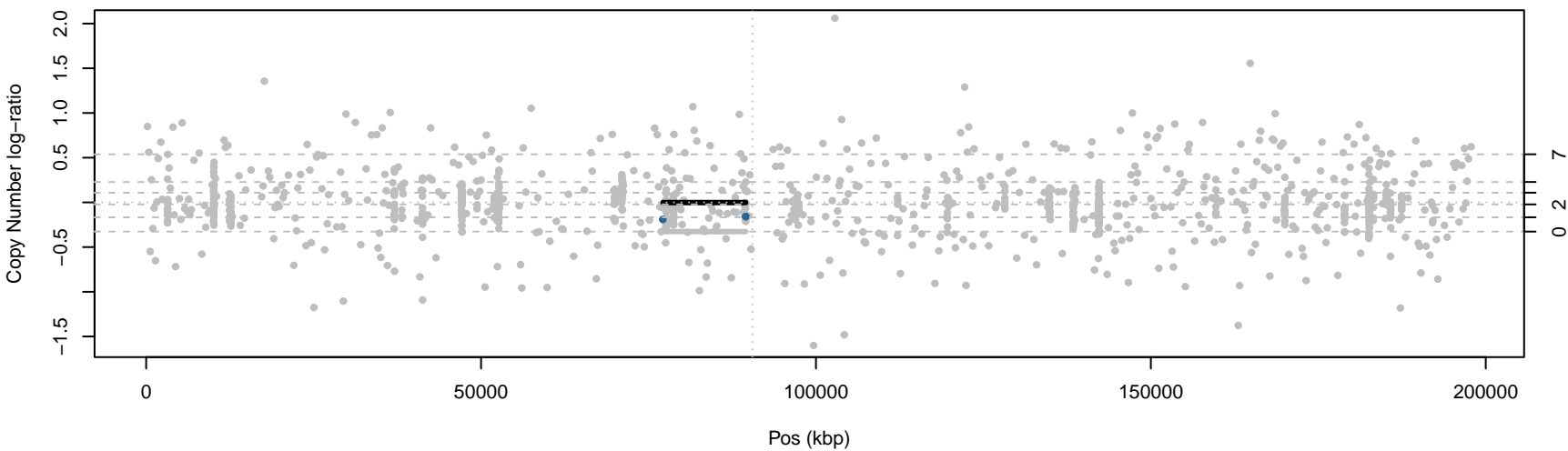
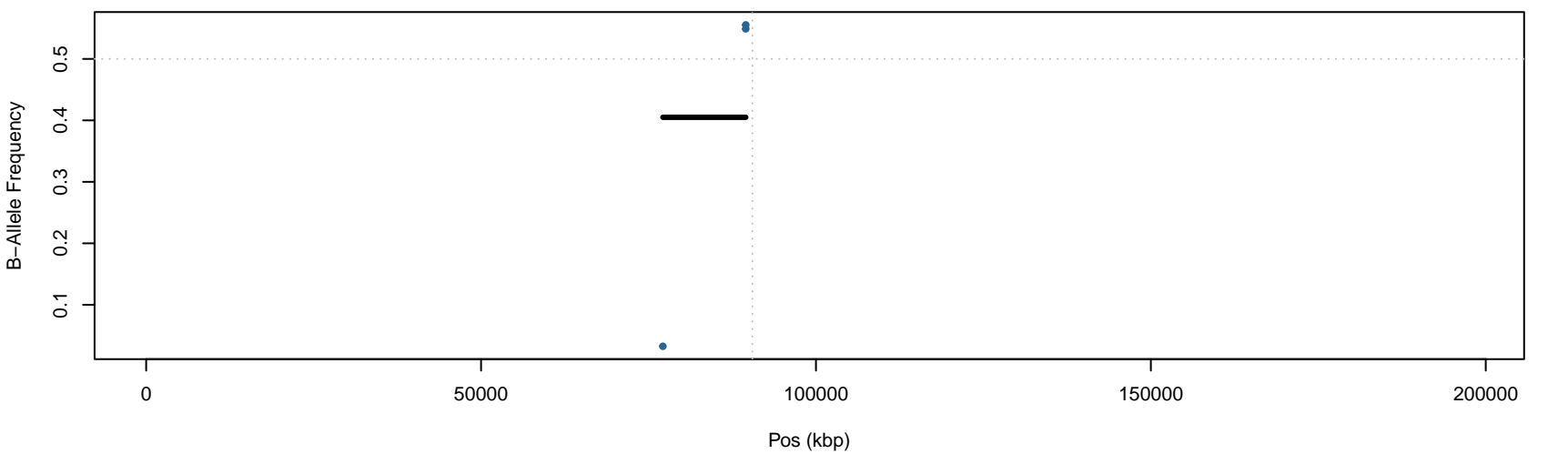


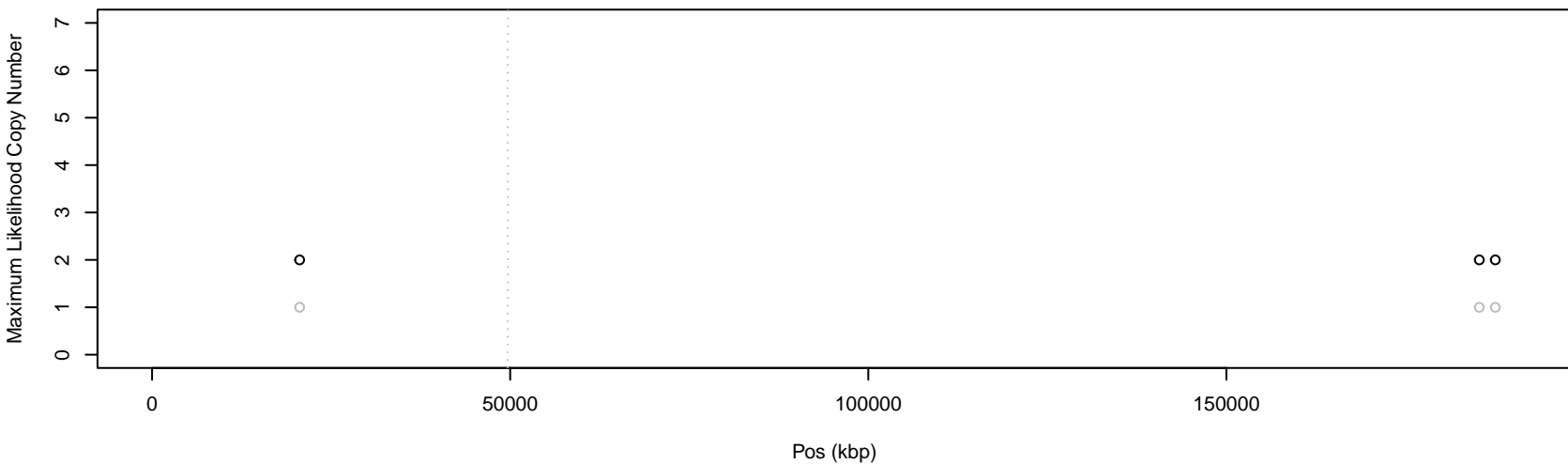
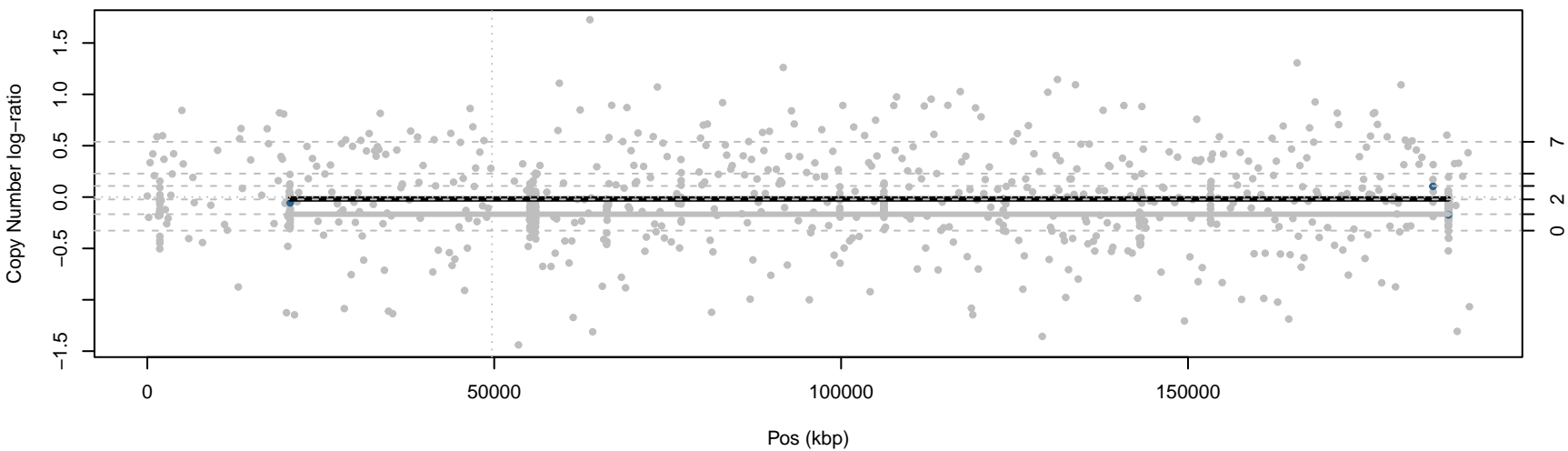
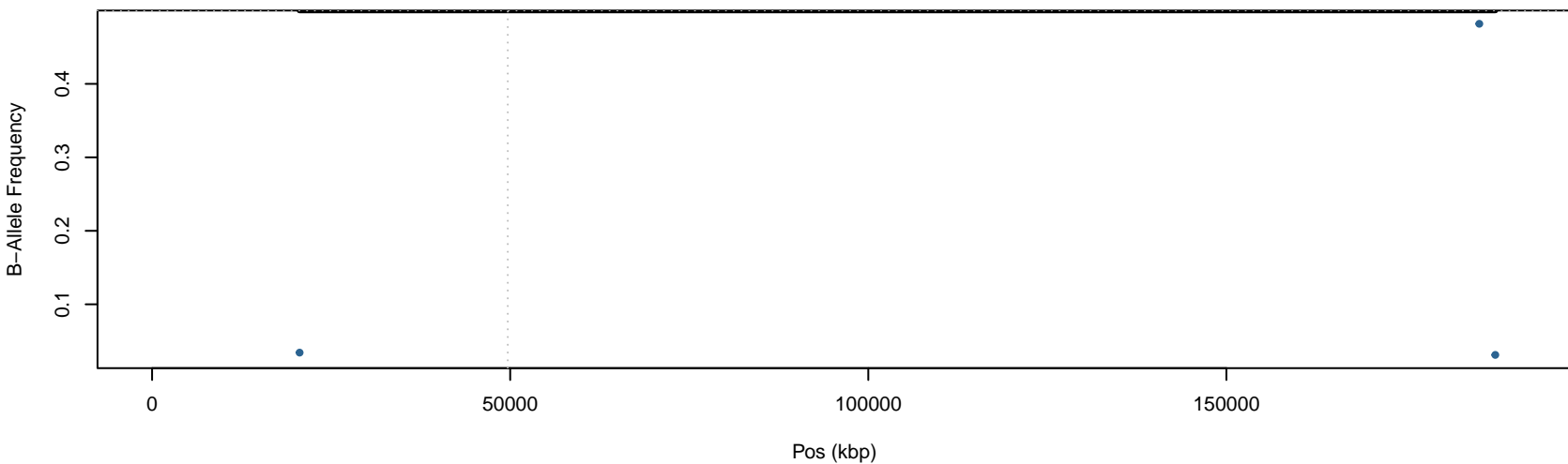
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 1



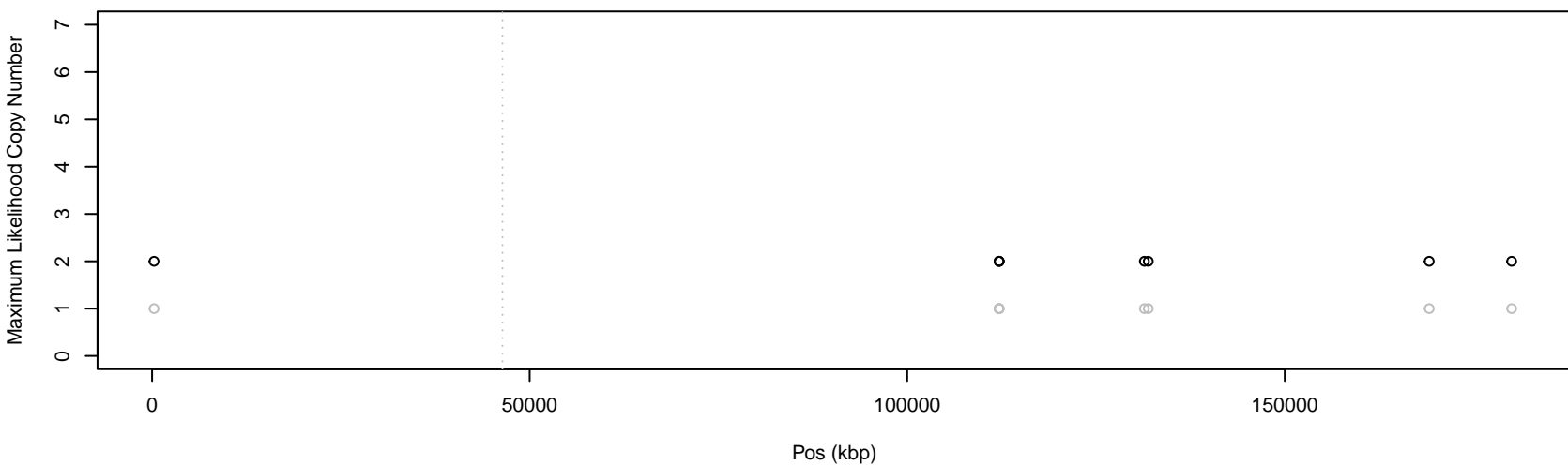
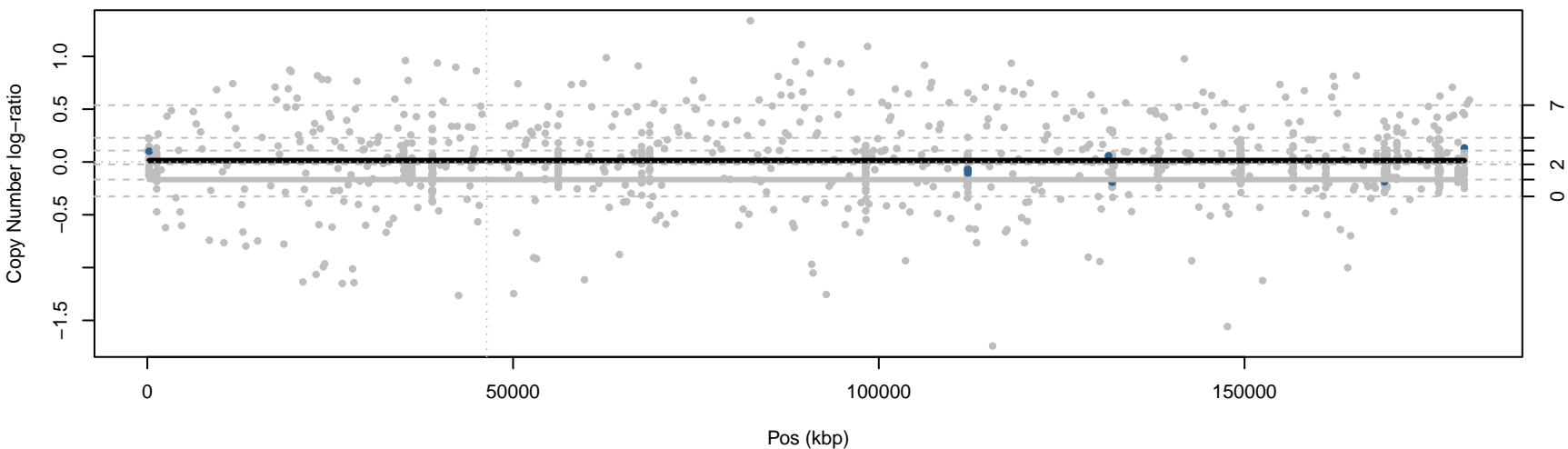
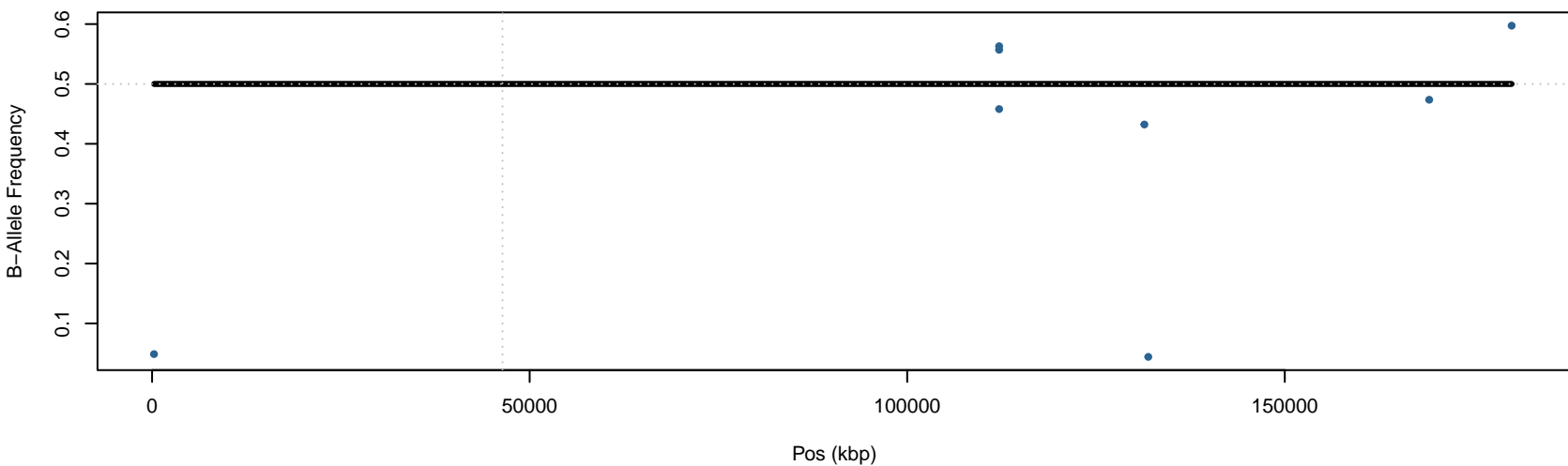
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 3



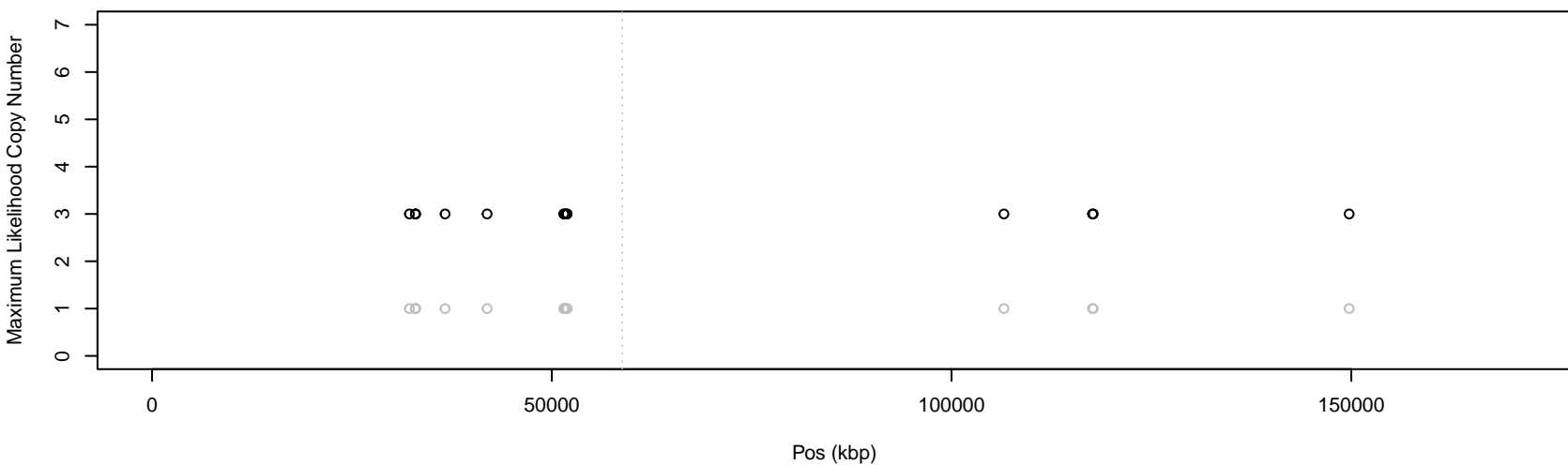
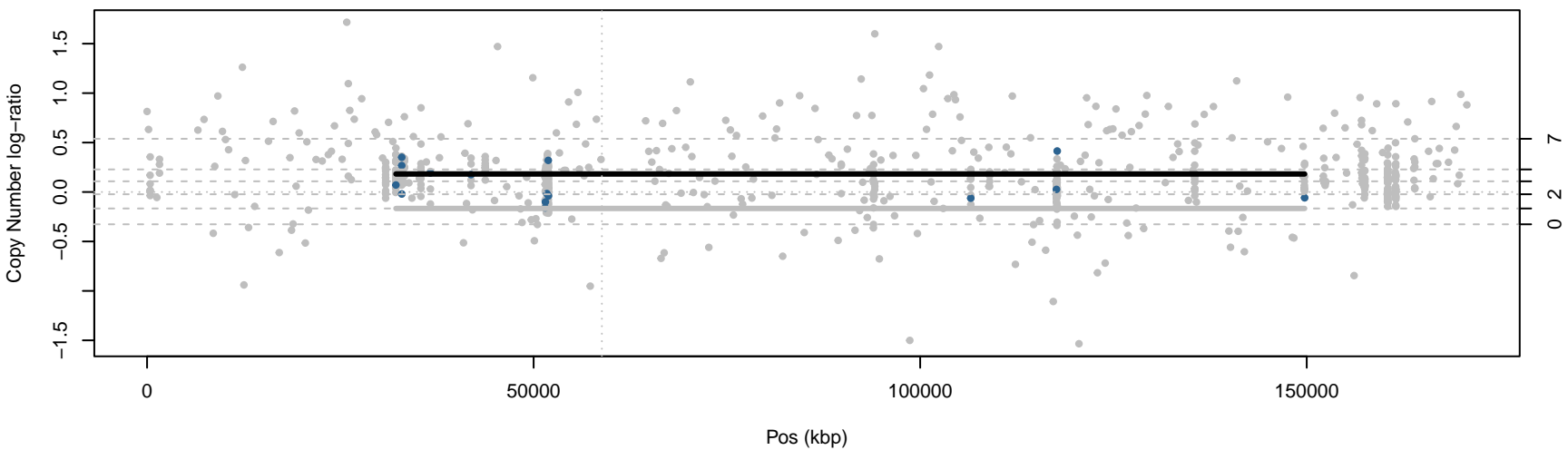
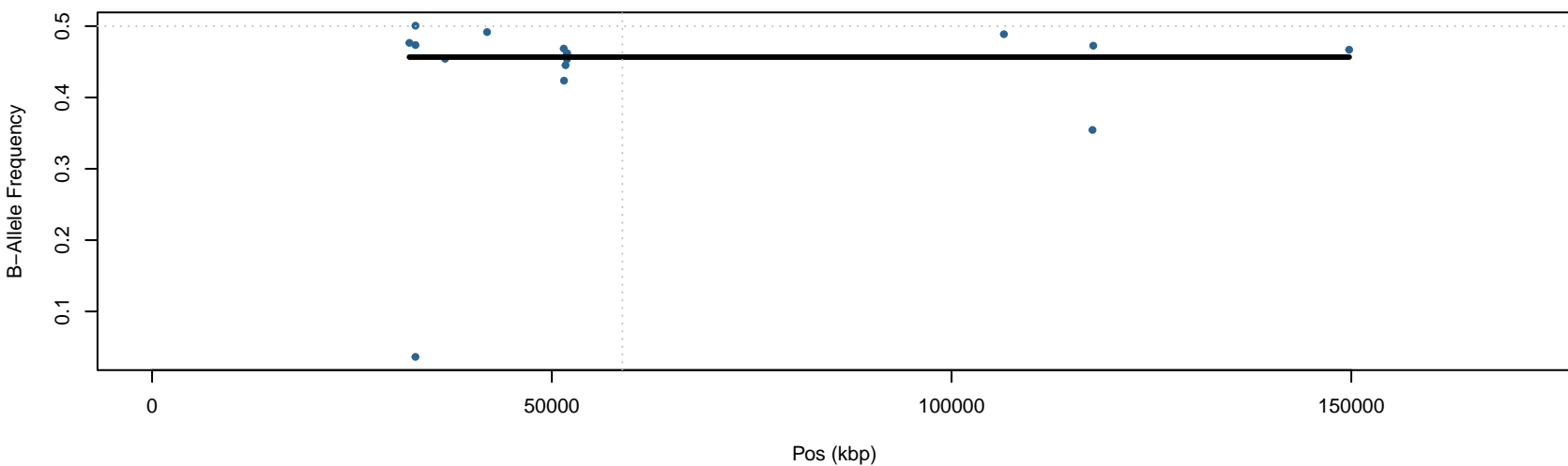
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 4



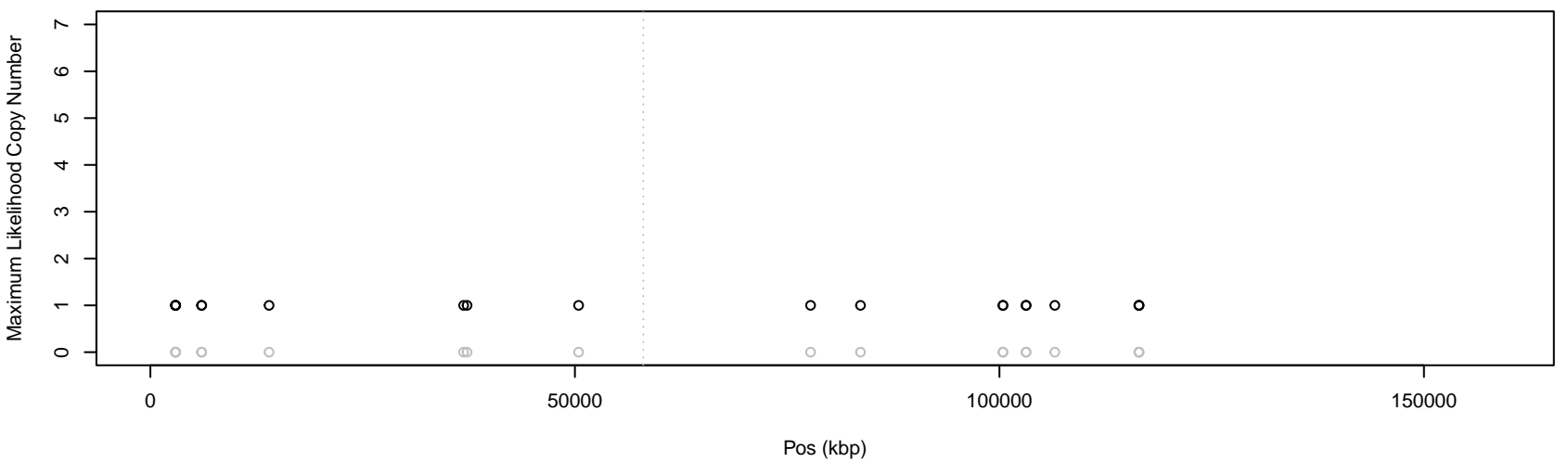
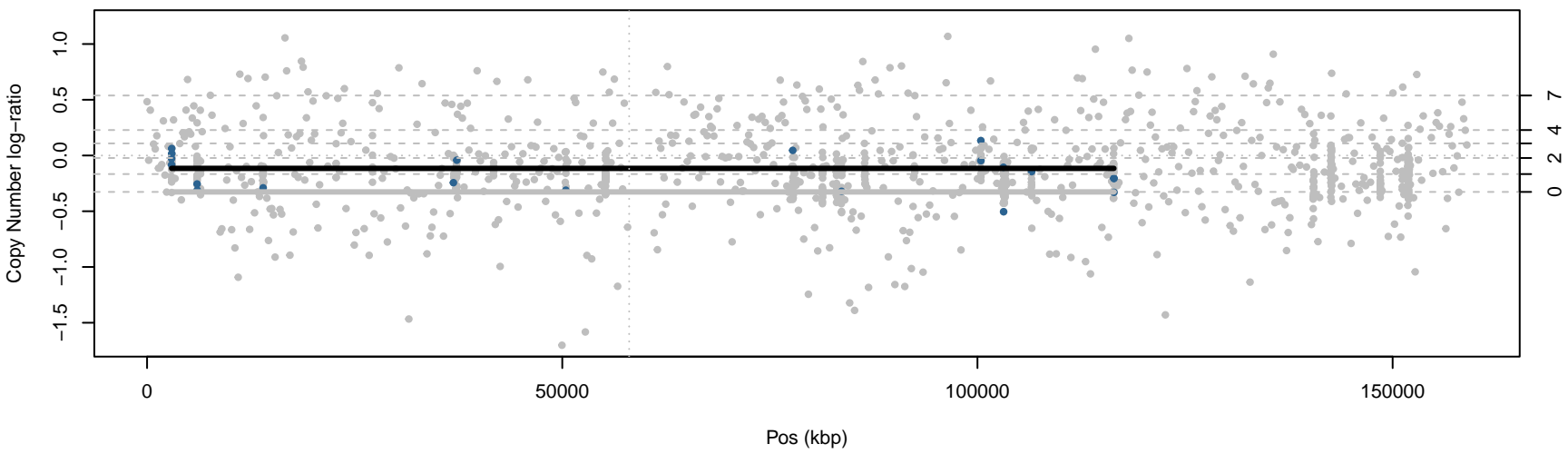
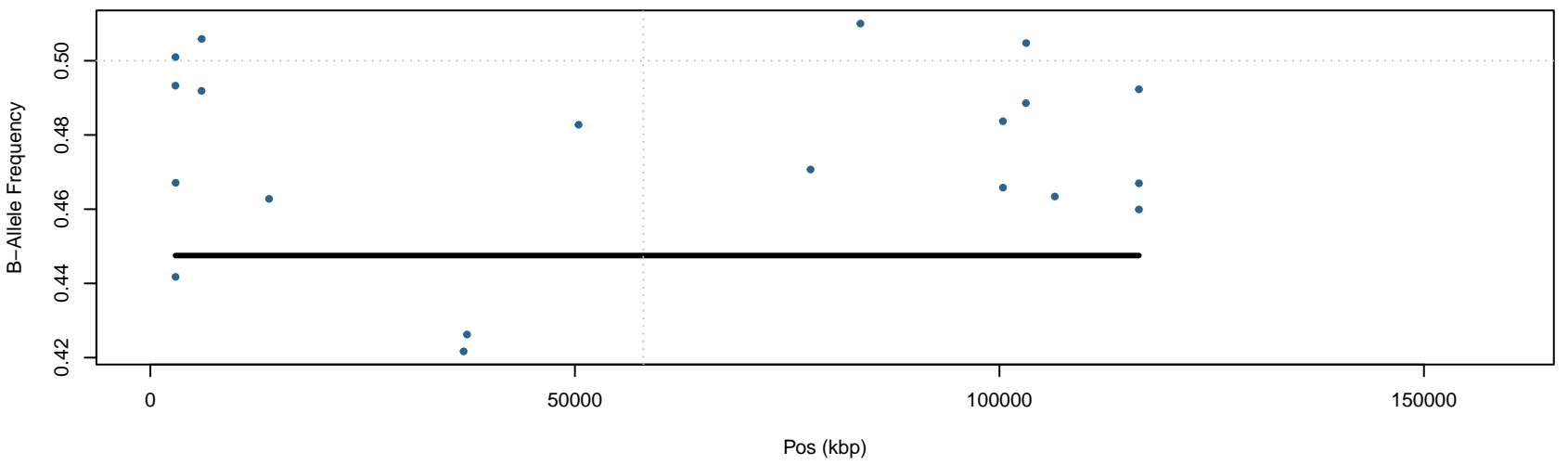
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 5



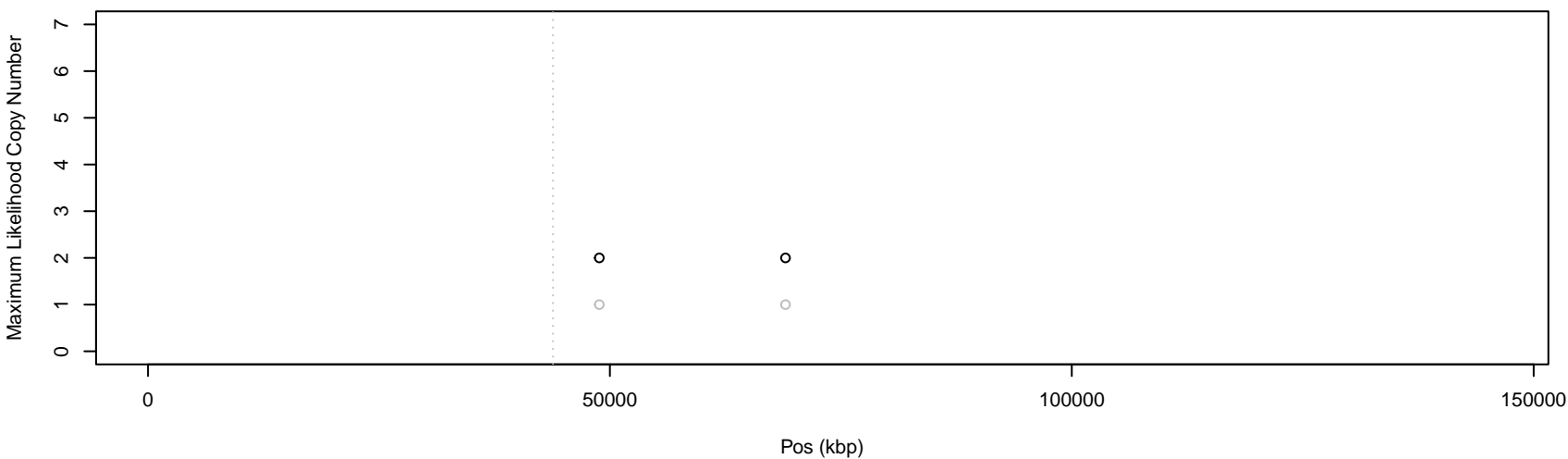
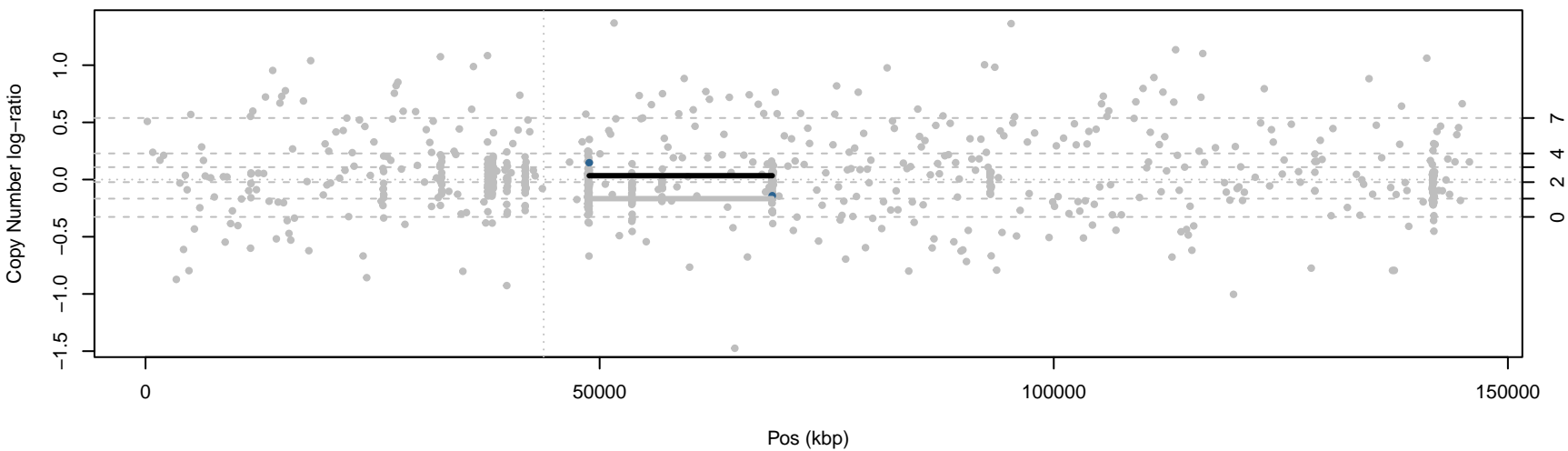
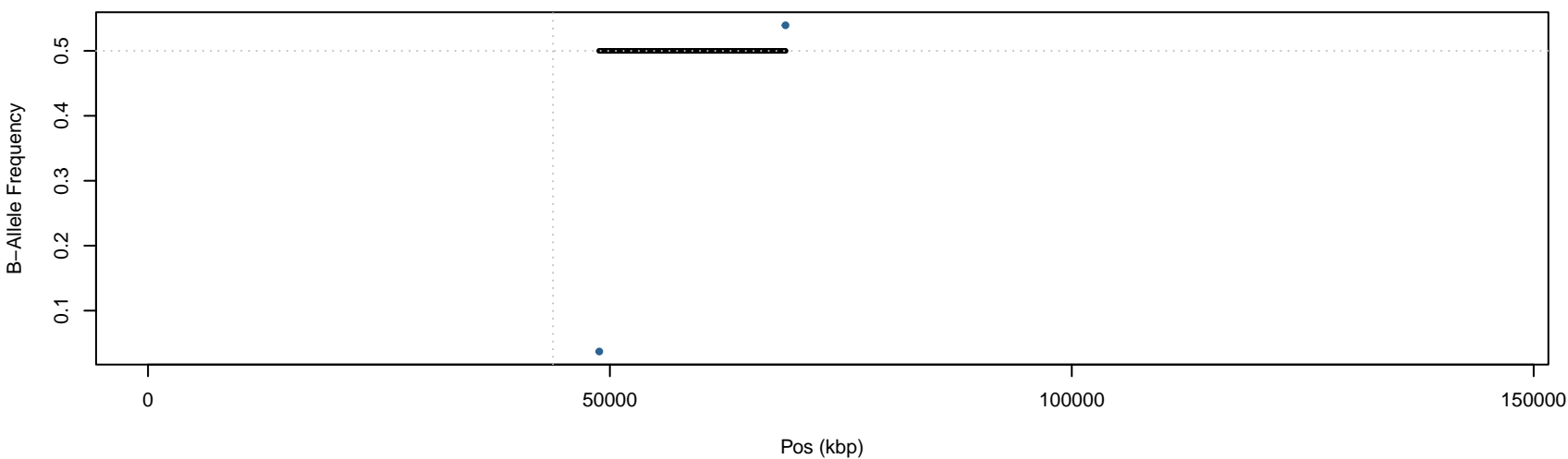
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 6



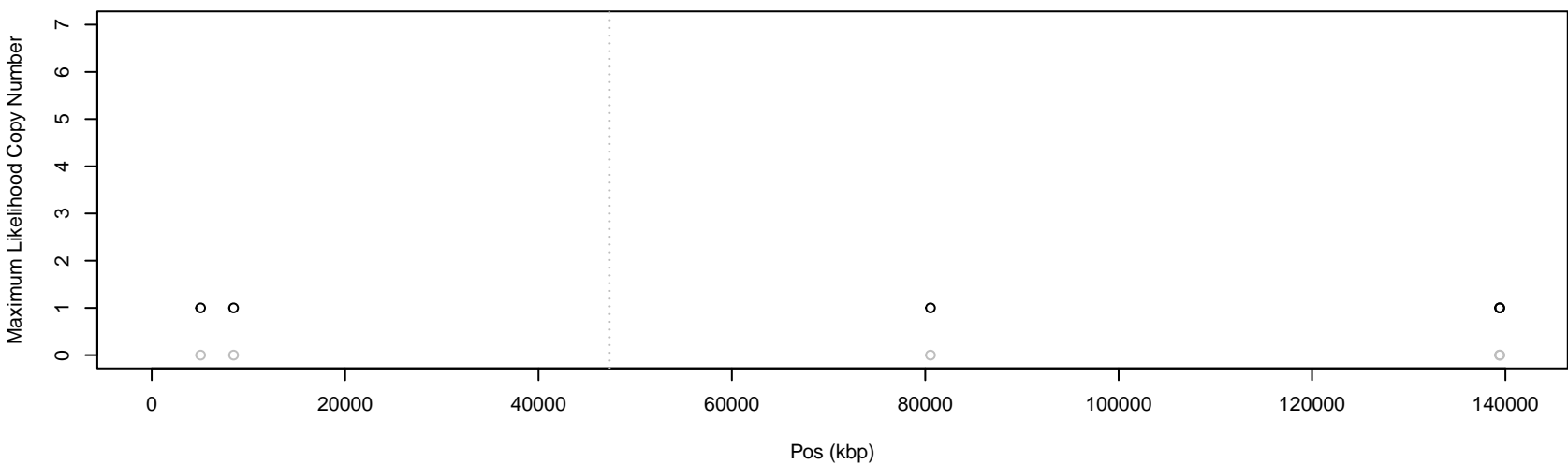
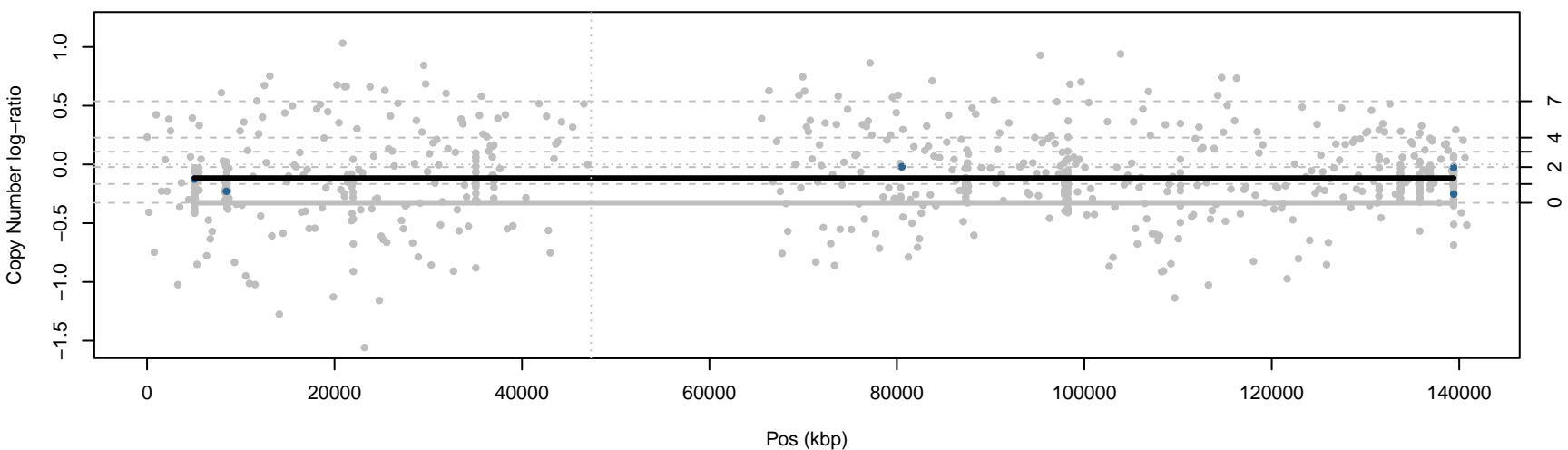
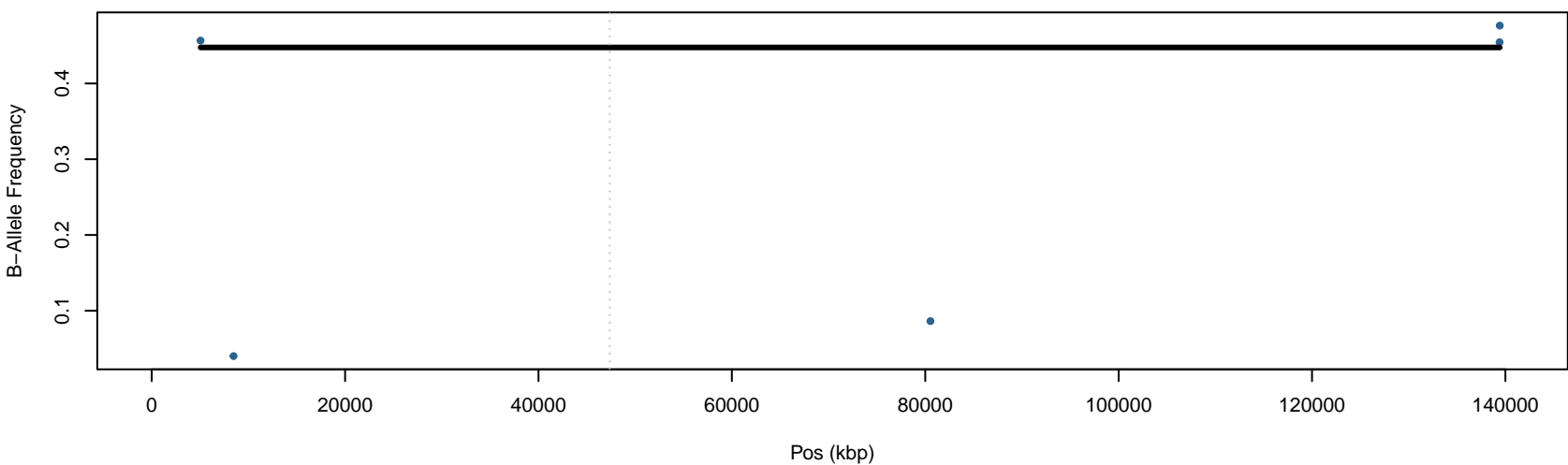
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 7



Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 8

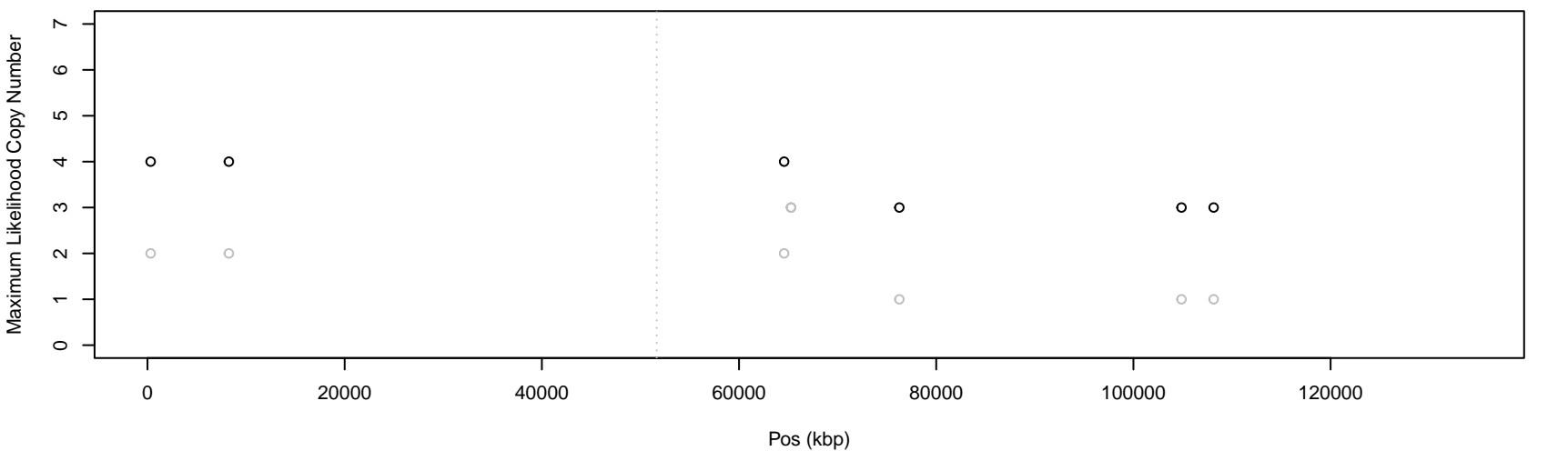
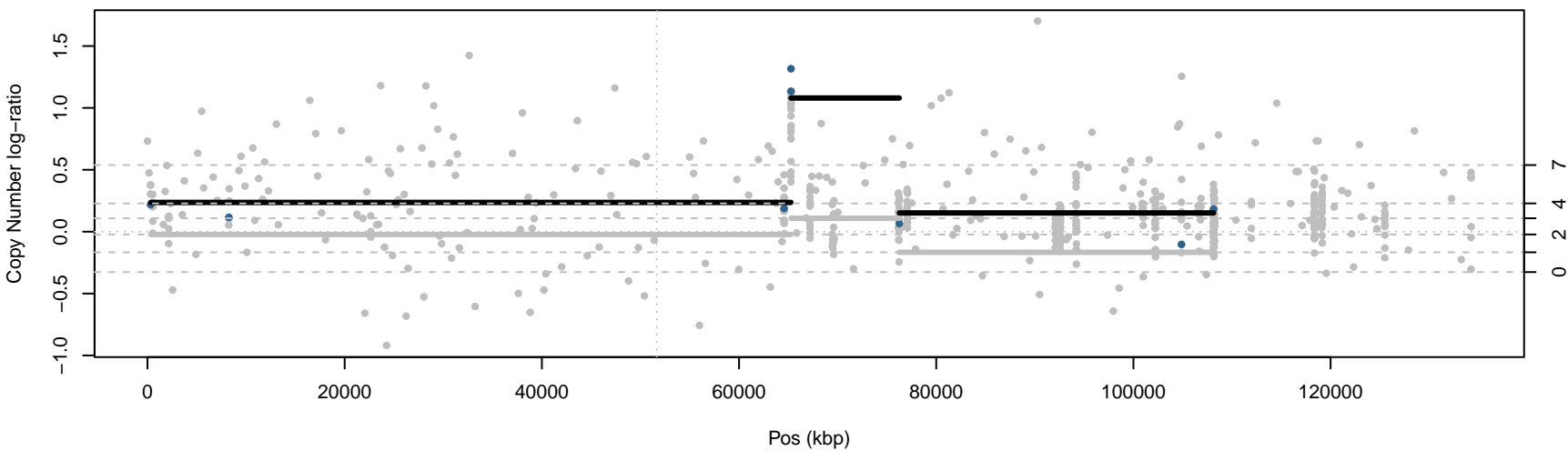
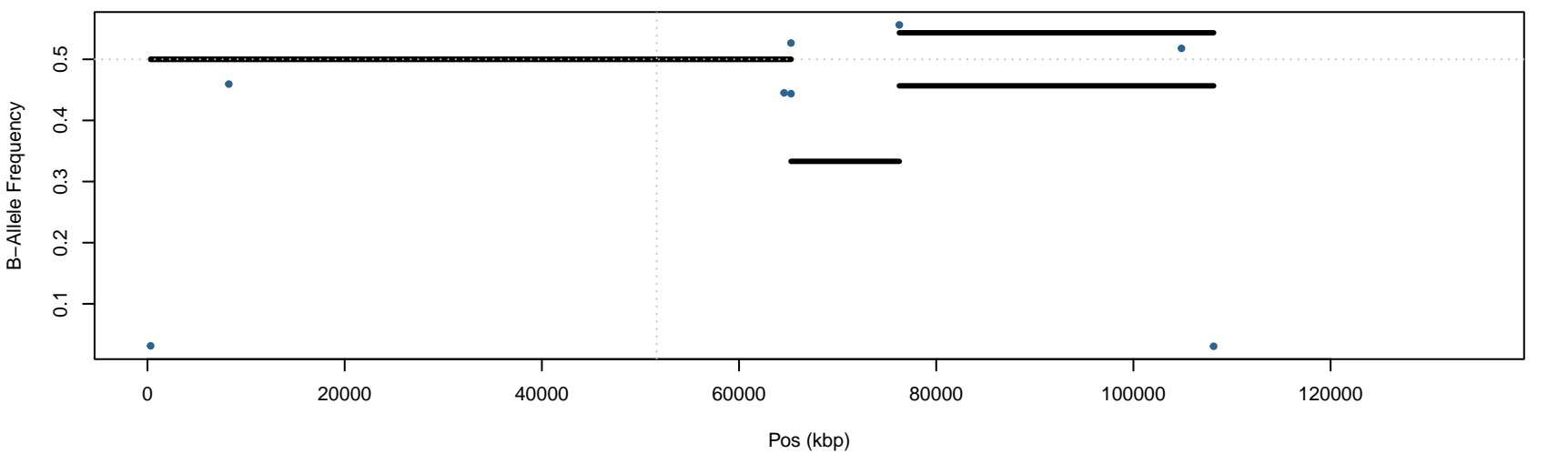


Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 9

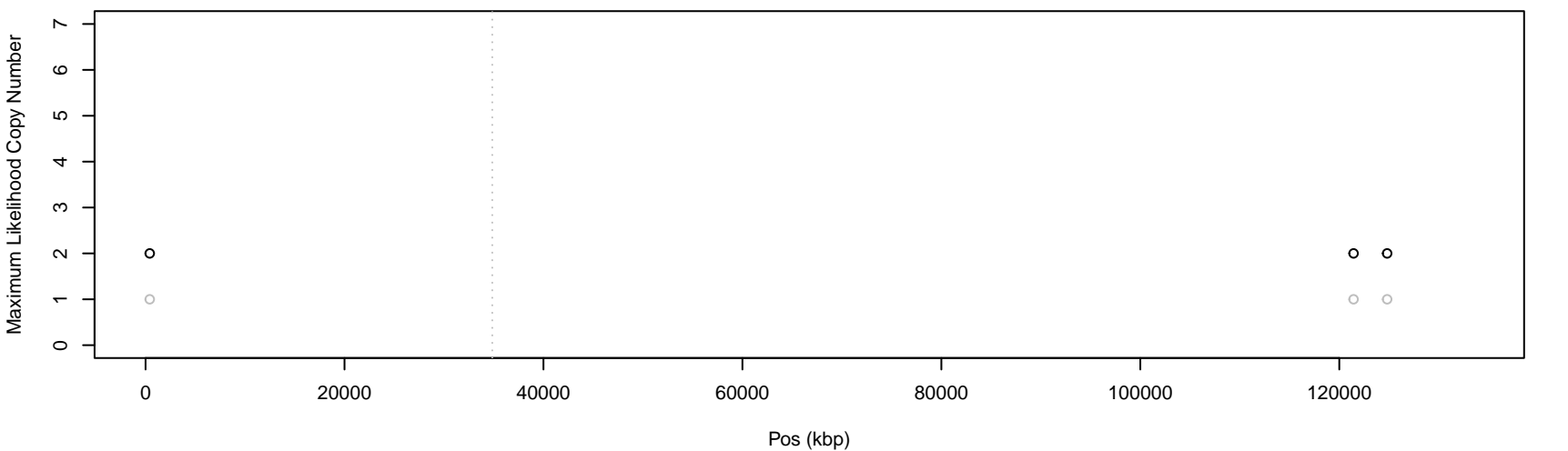
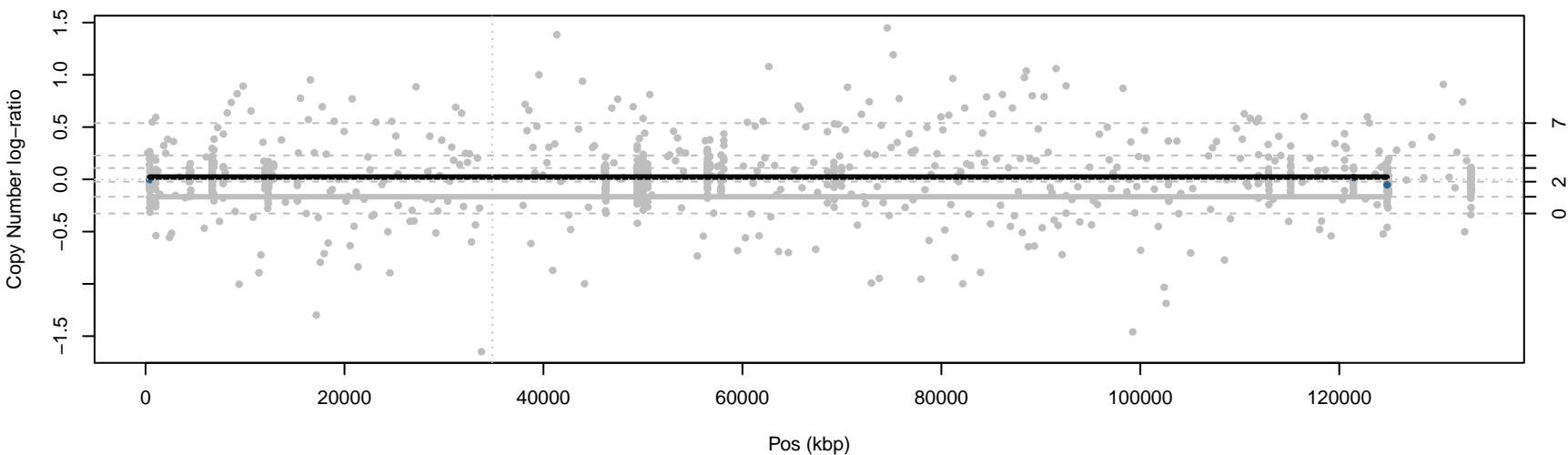
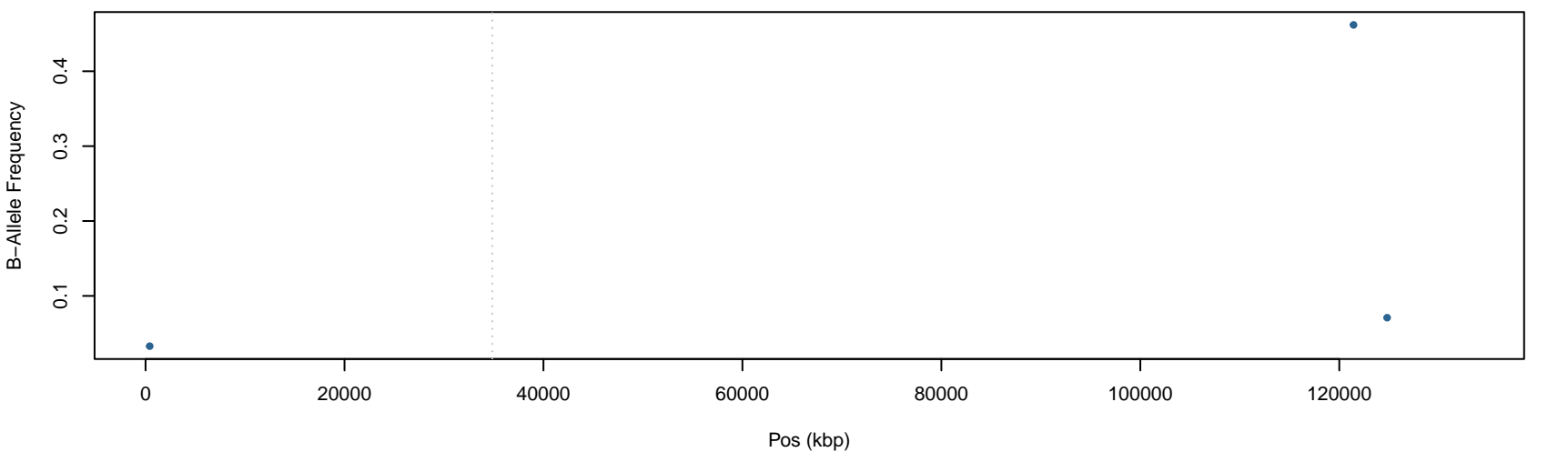




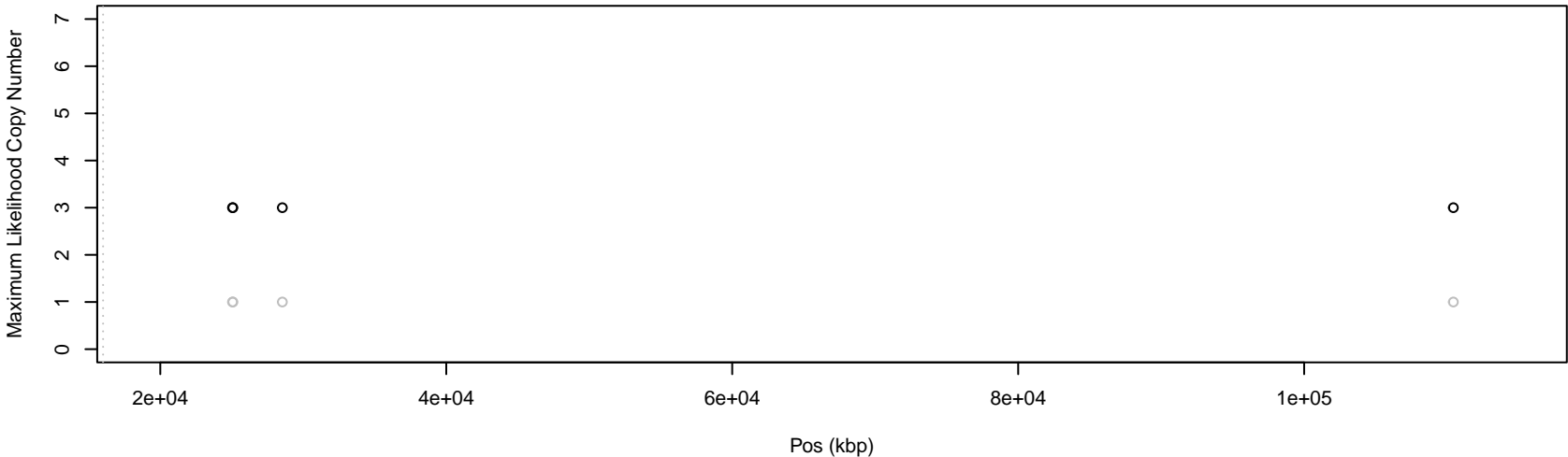
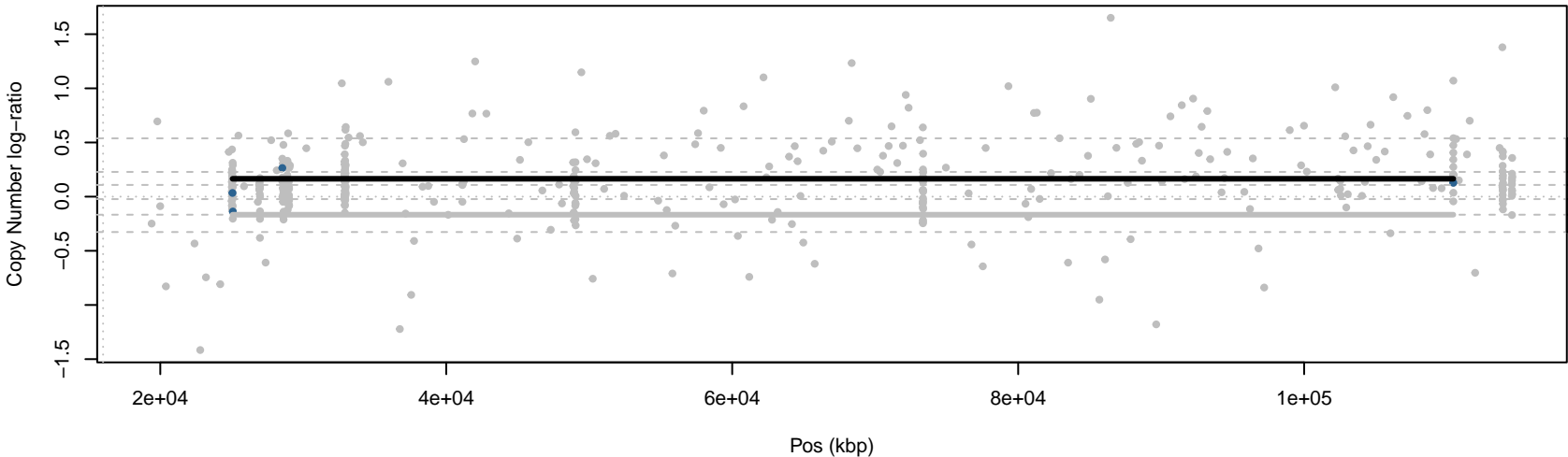
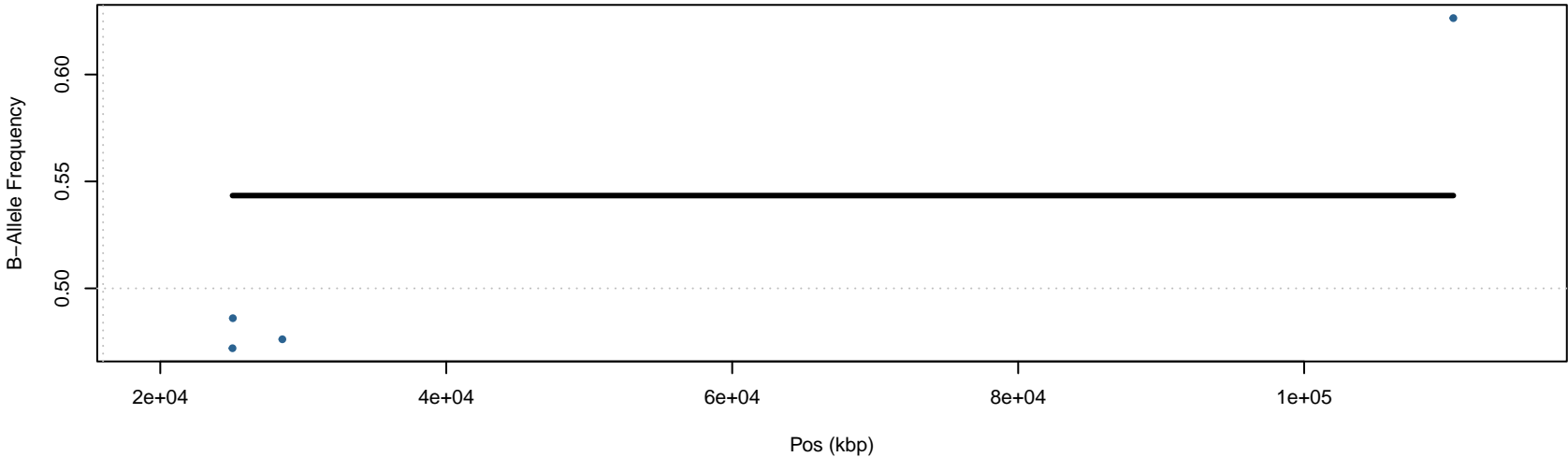
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 11



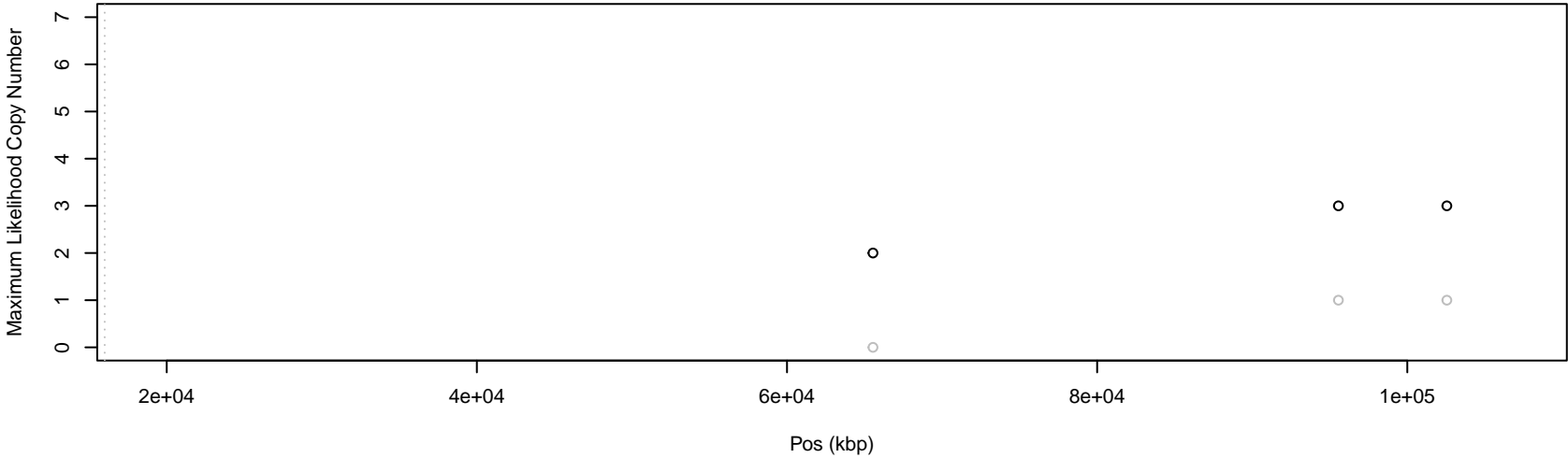
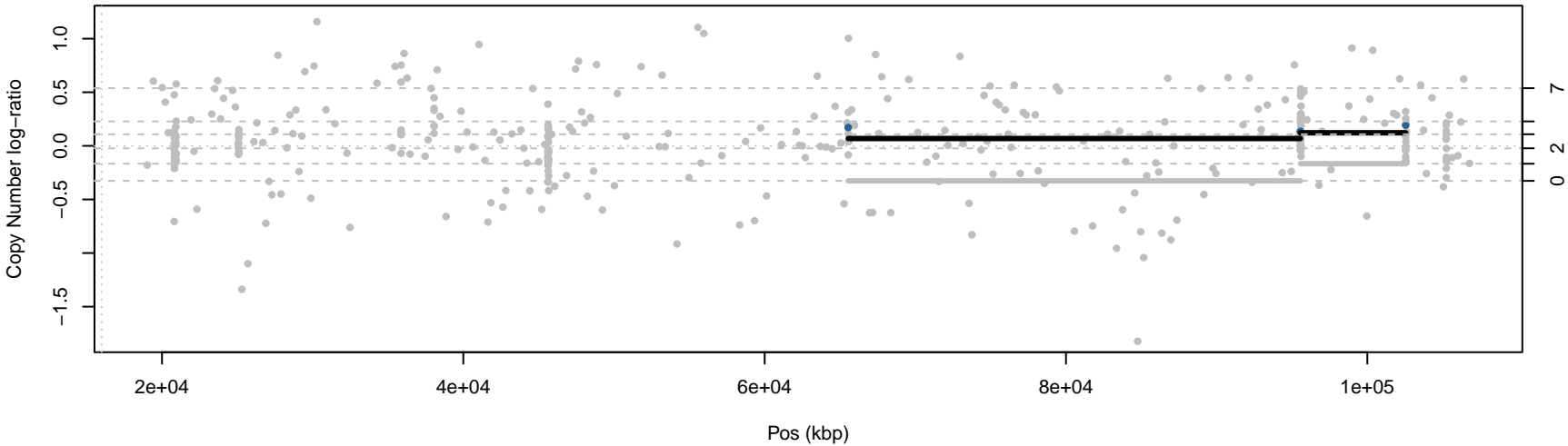
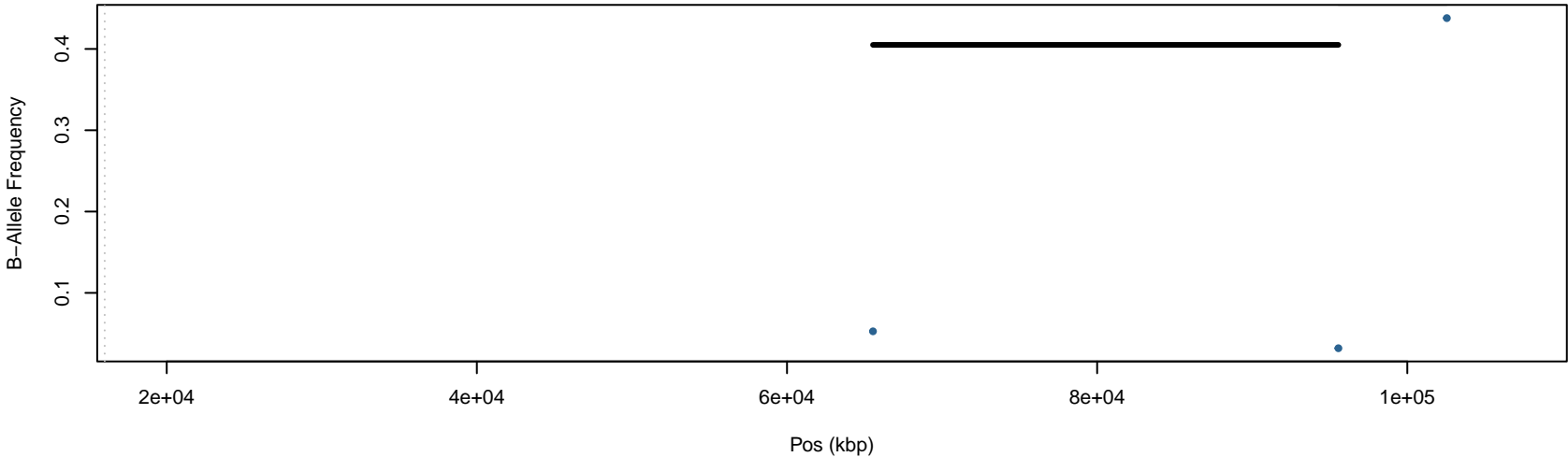
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 12



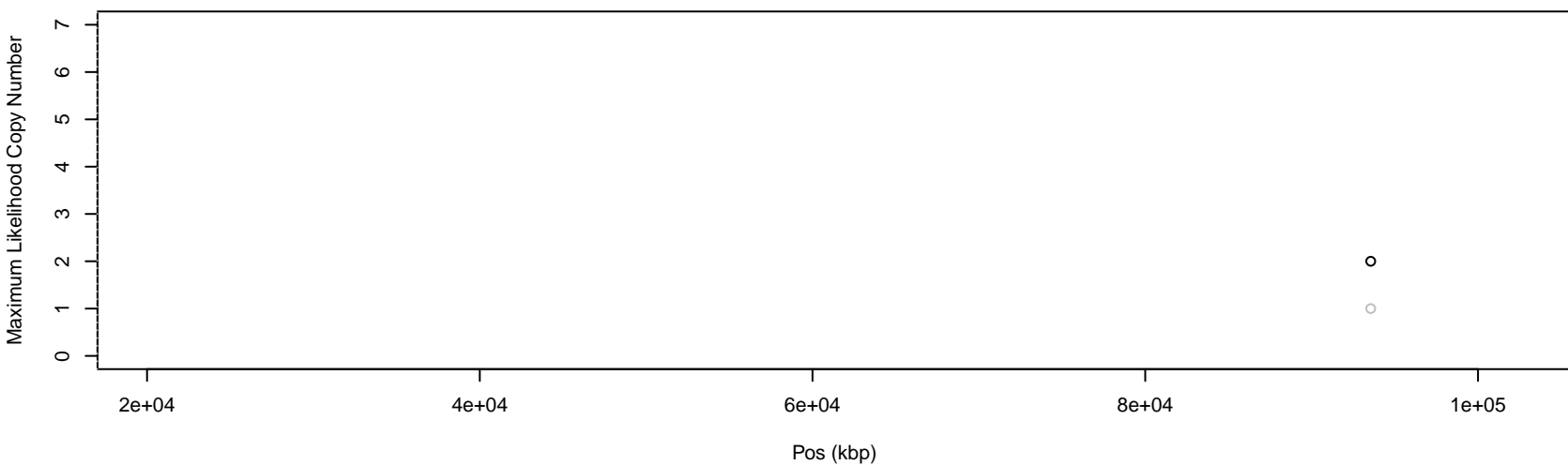
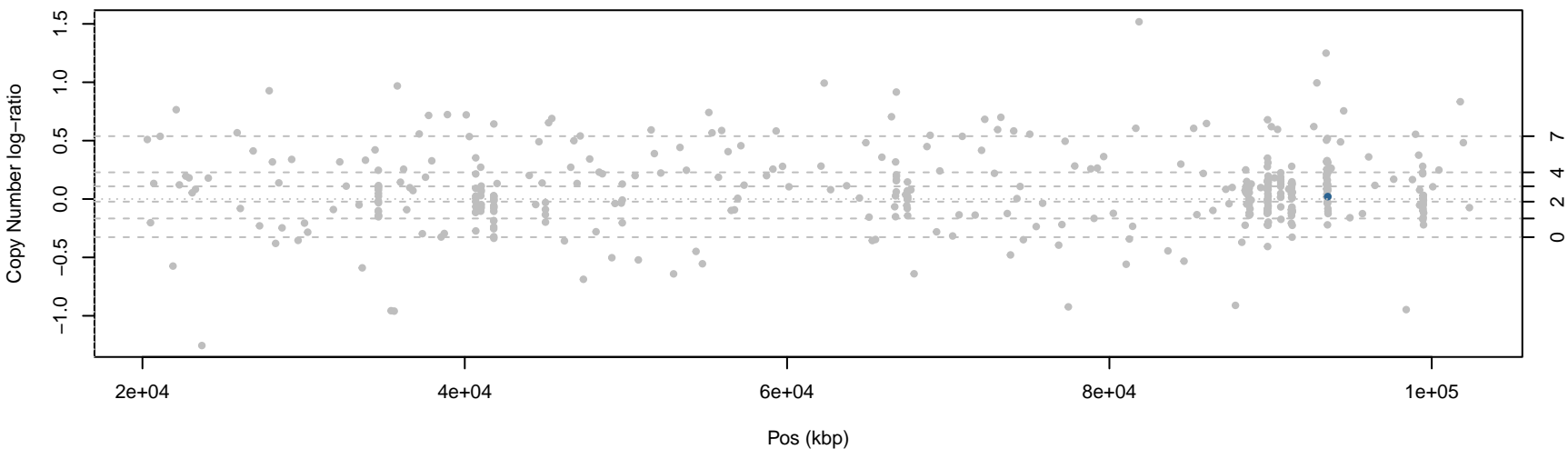
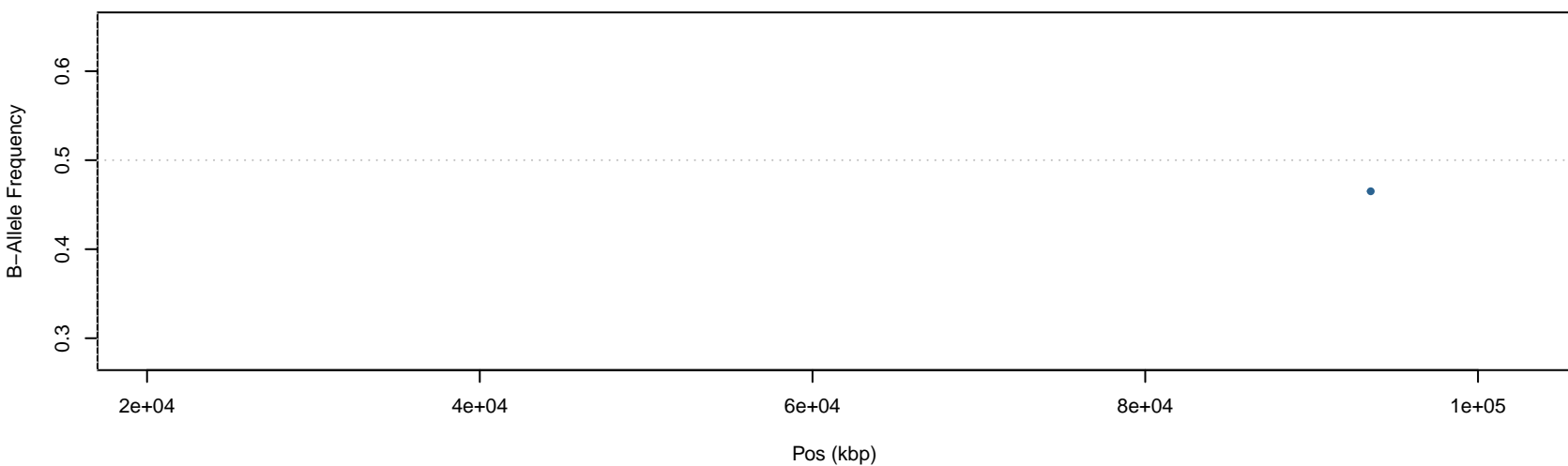
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 13



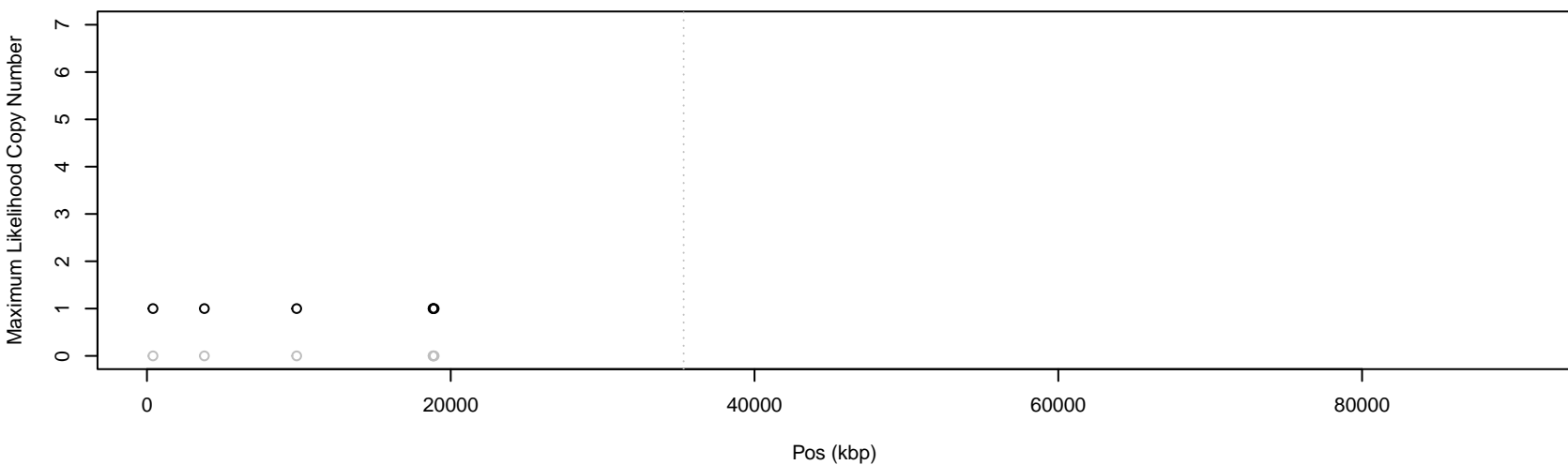
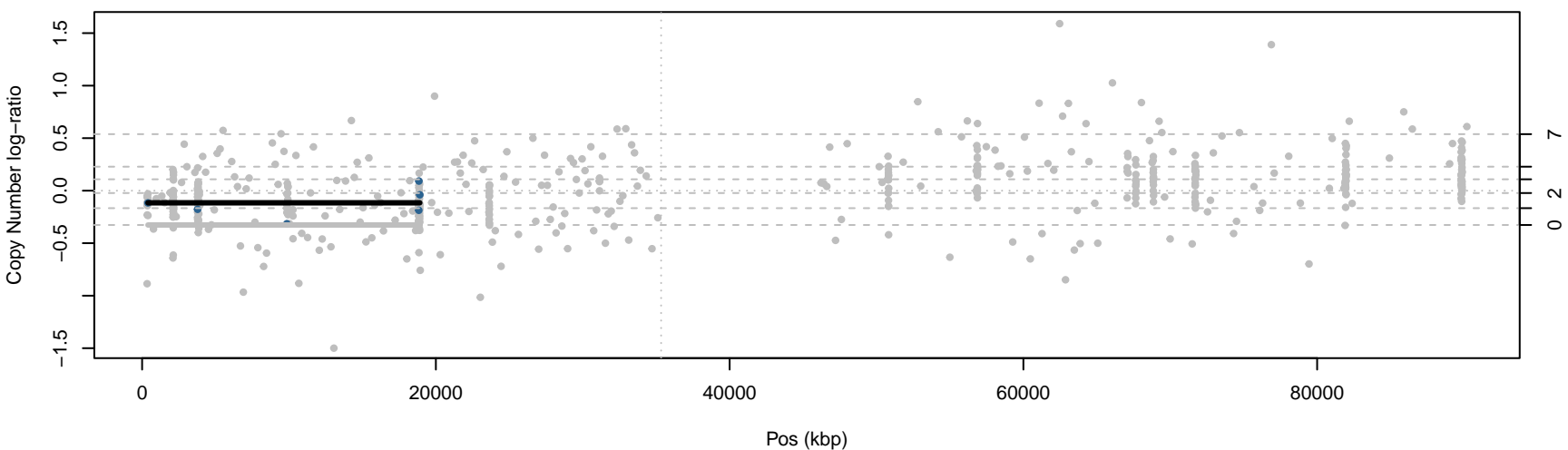
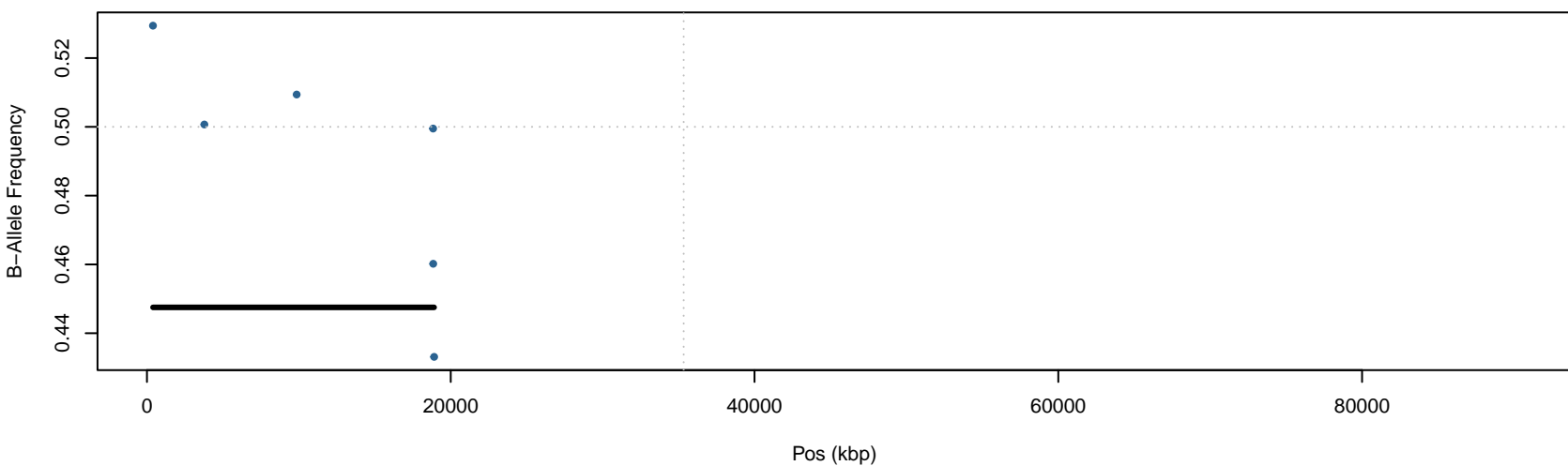
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 14



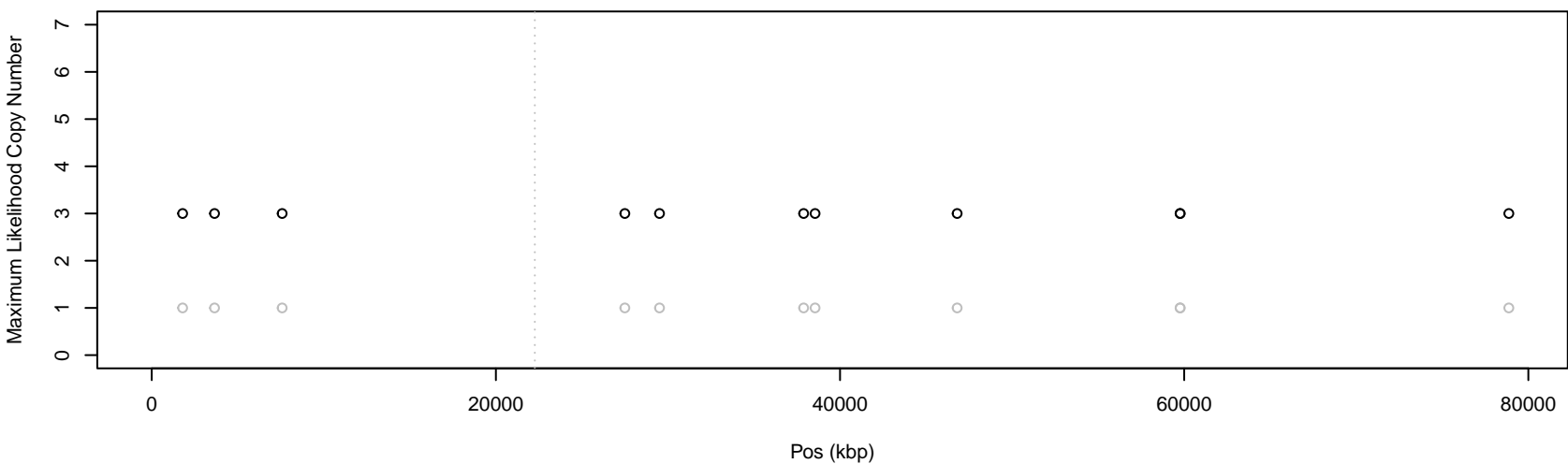
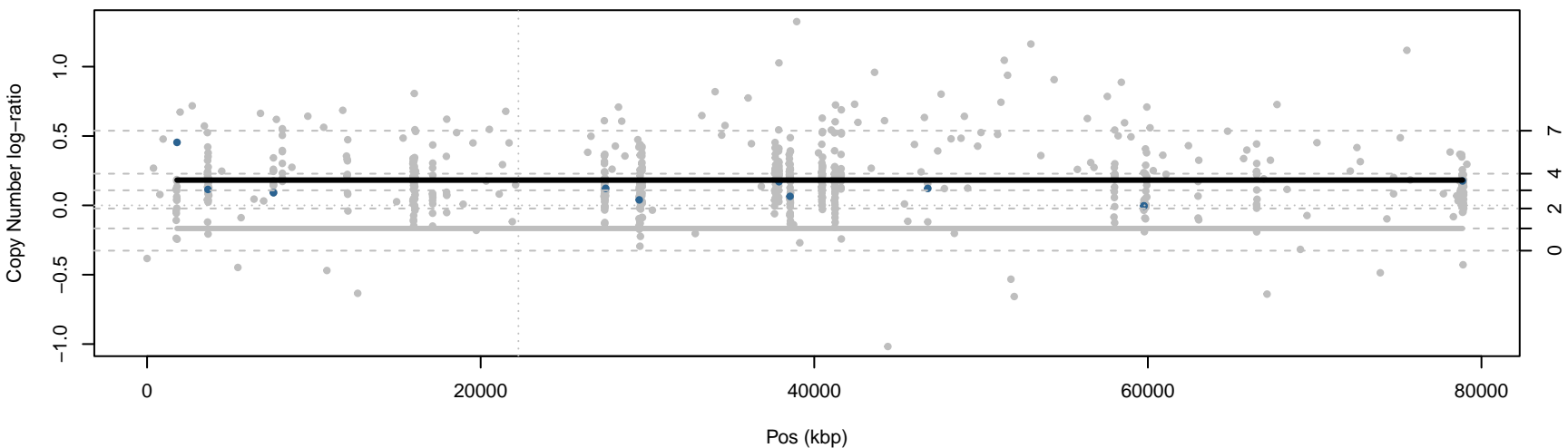
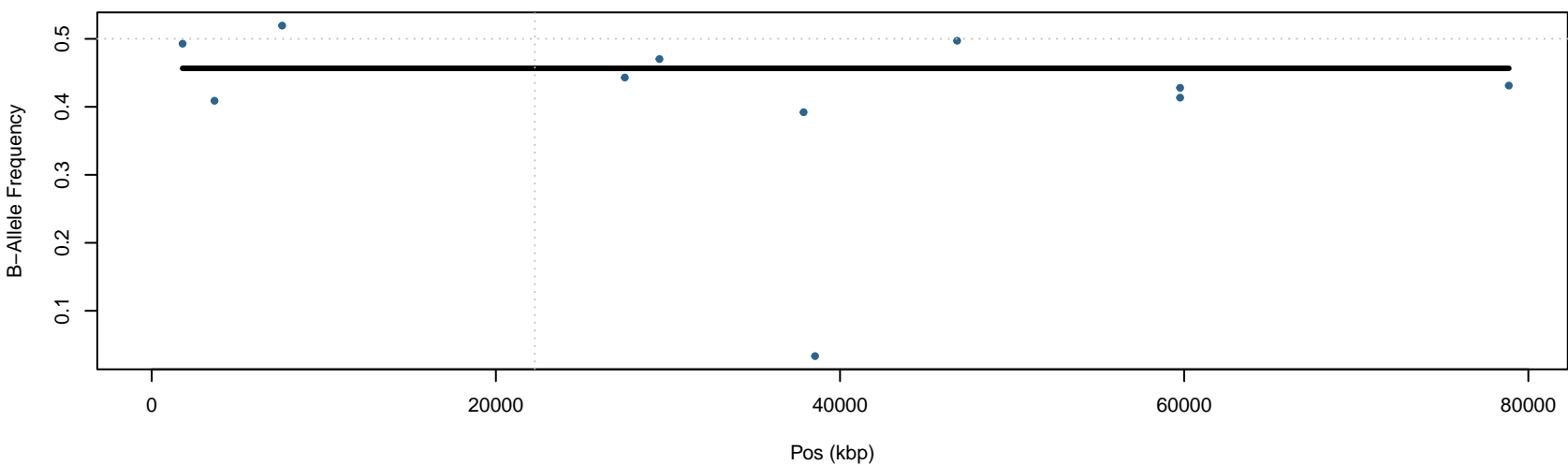
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 15



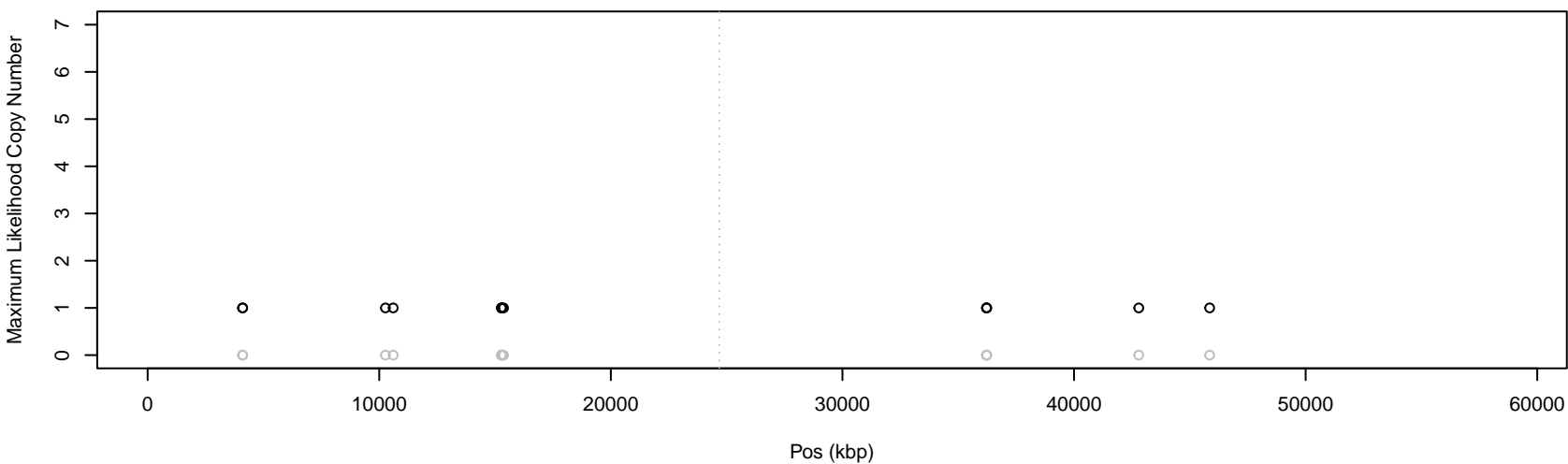
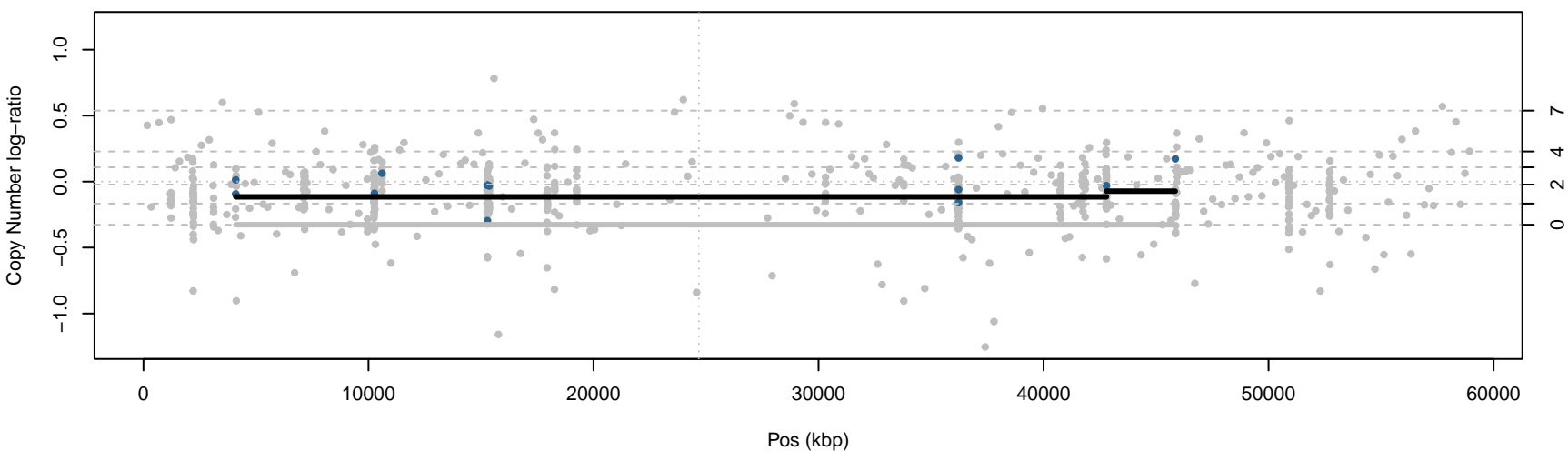
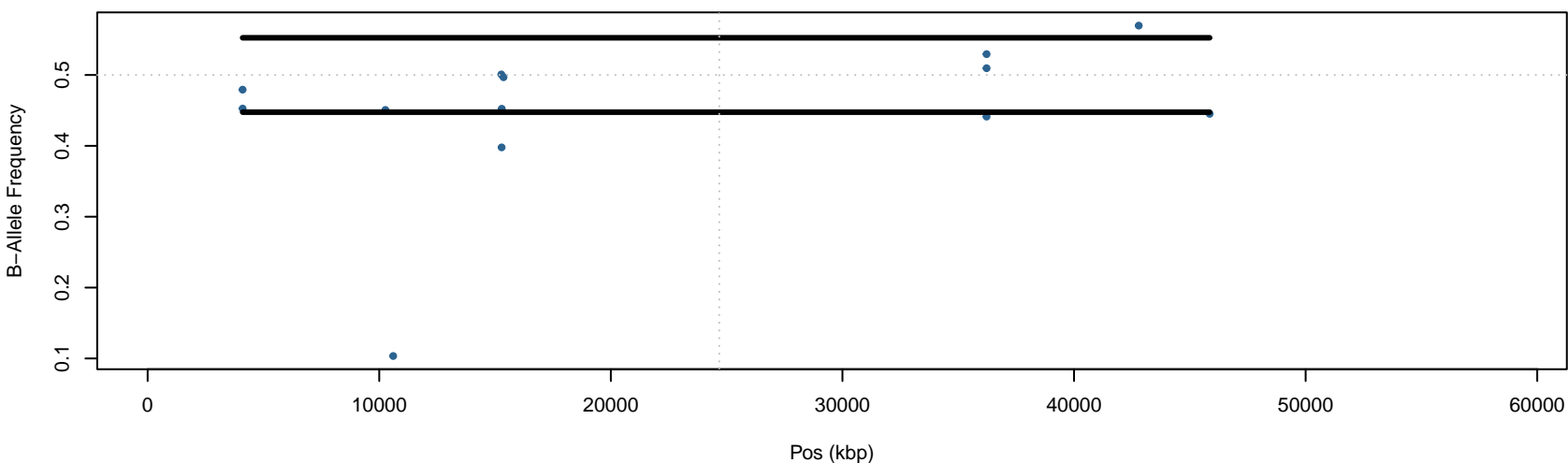
Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 16



Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 17

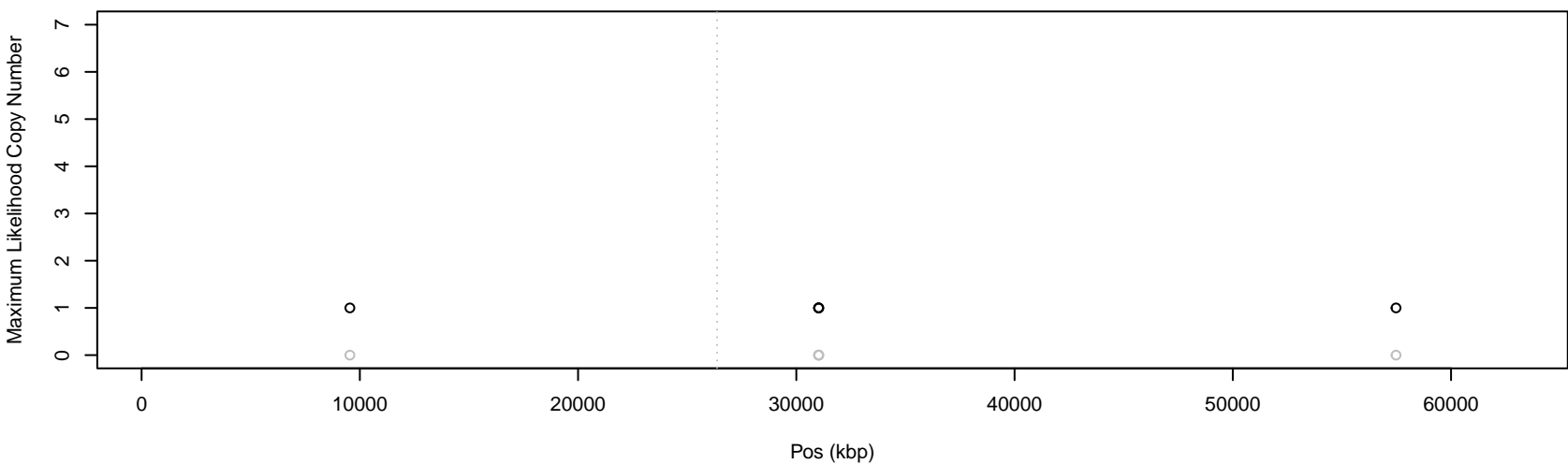
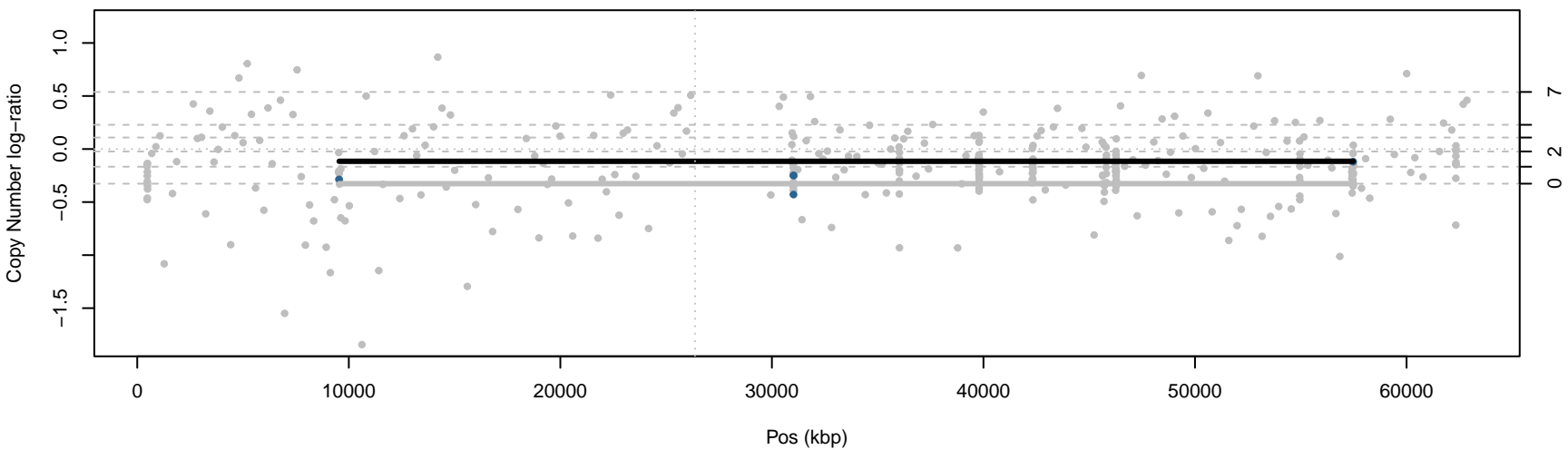
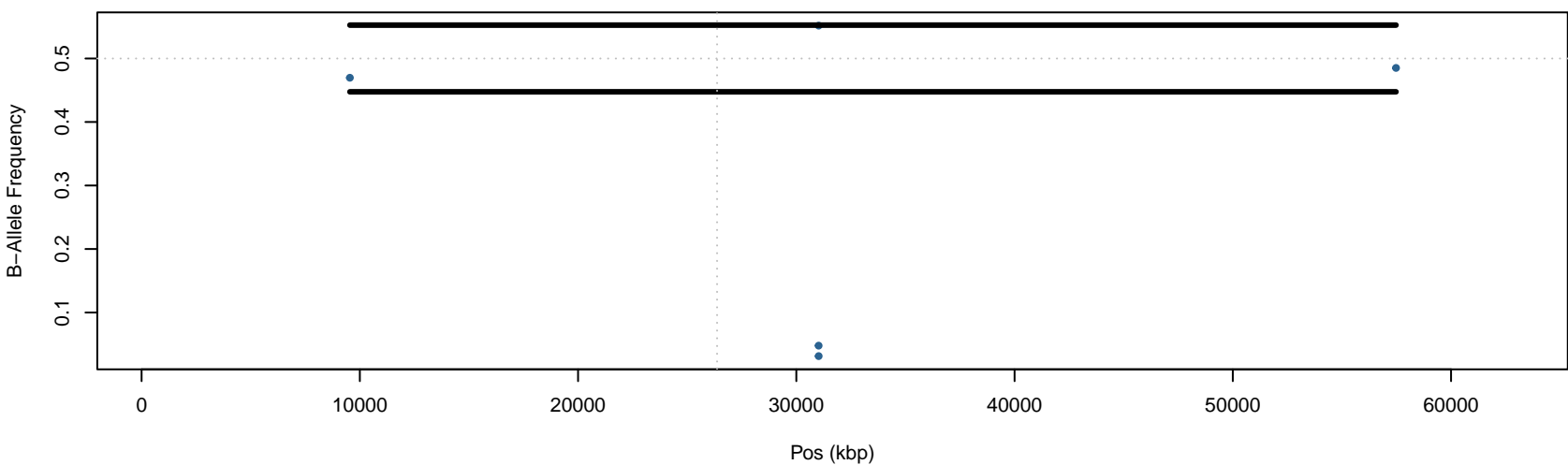


Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 19

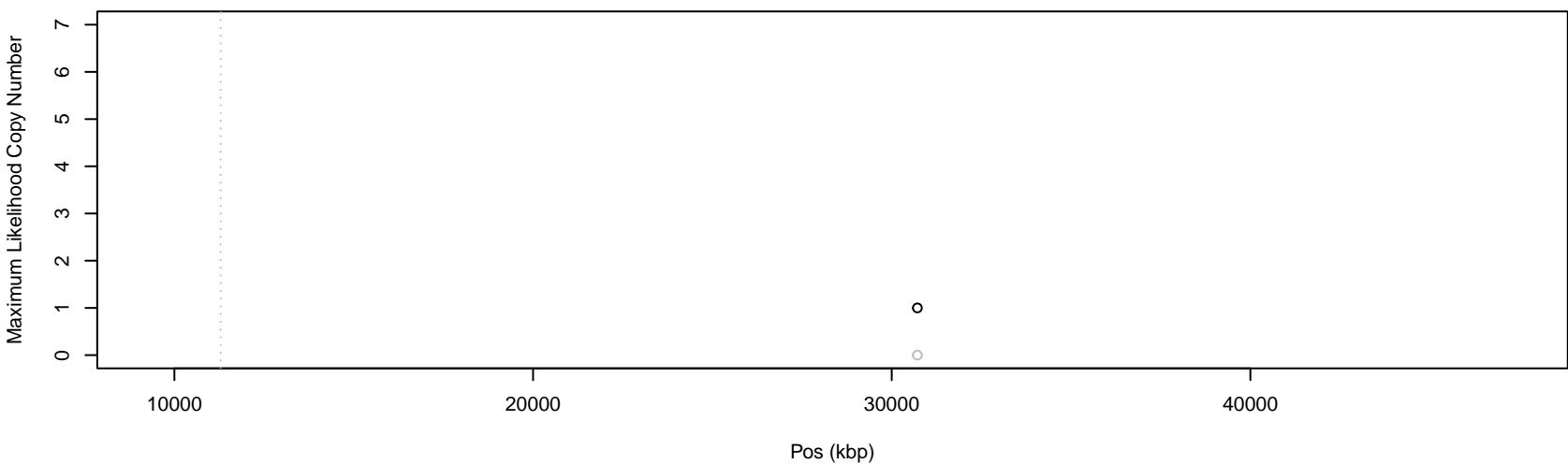
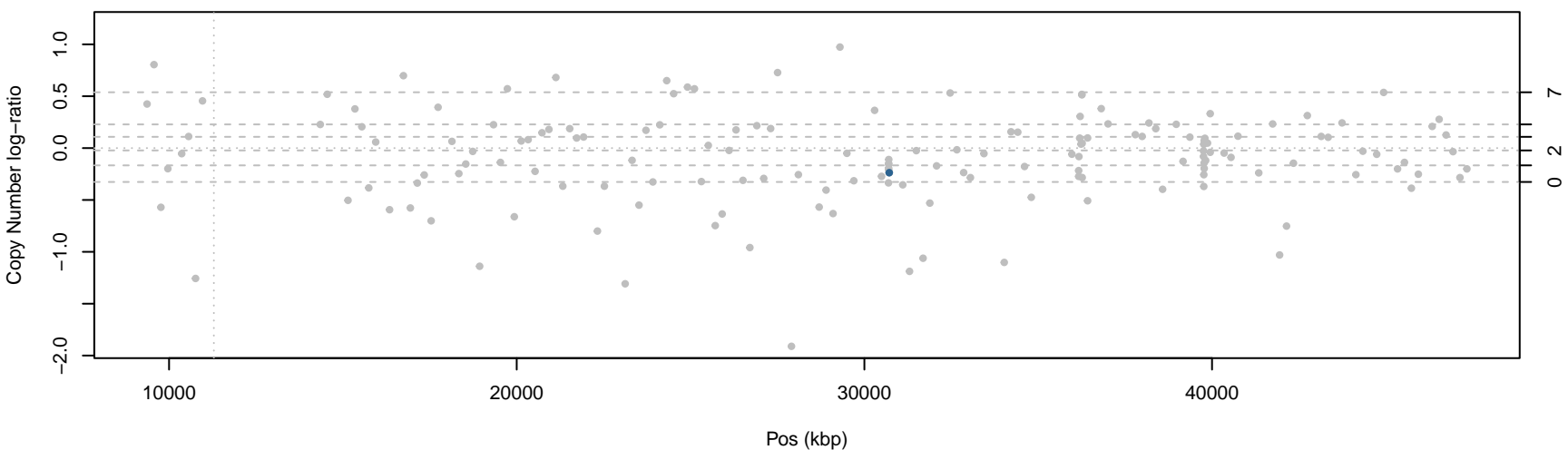
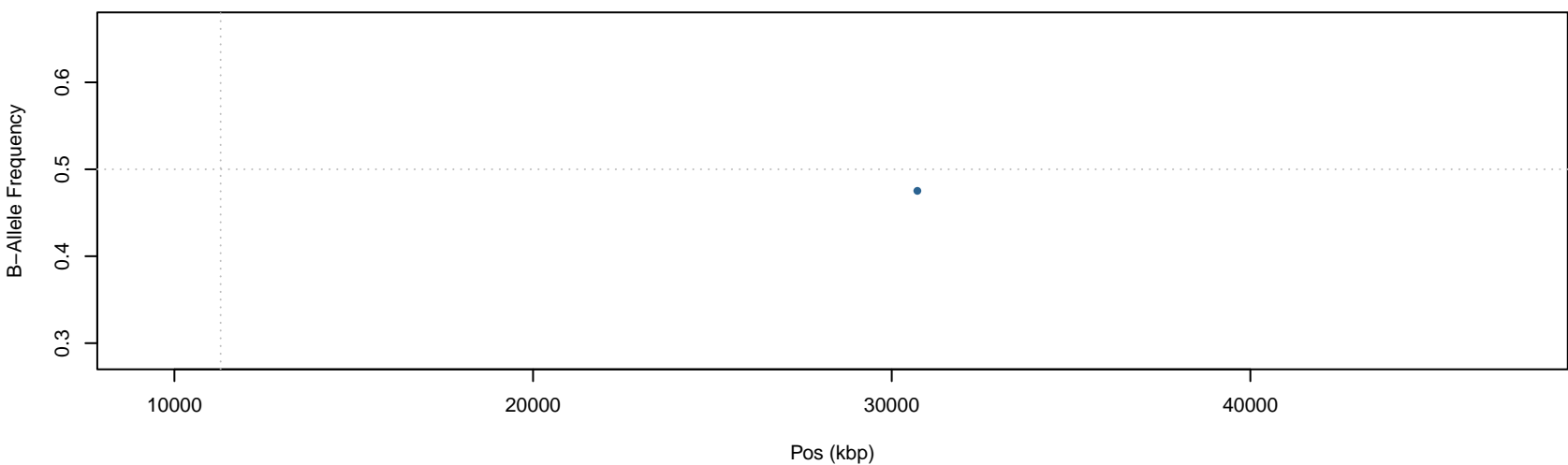




Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 20



Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 21



Purity: 0.19 Tumor ploidy: 2.166 SNV log-likelihood: 89.03 GoF: 95.4% Mean coverage: 597;603 Chromosome: 22

