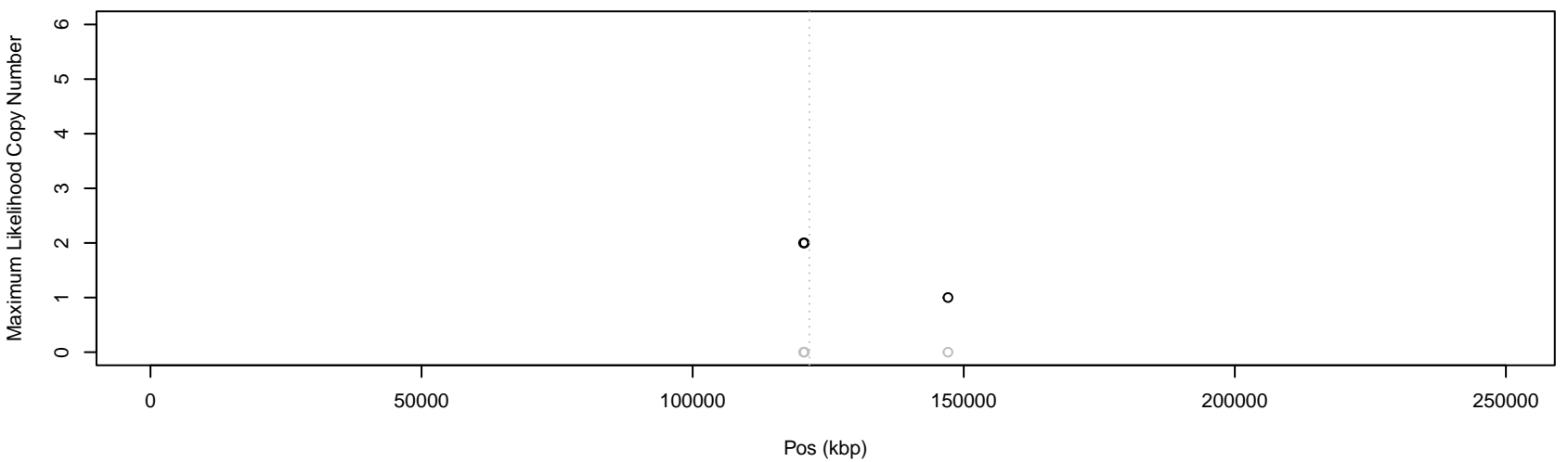
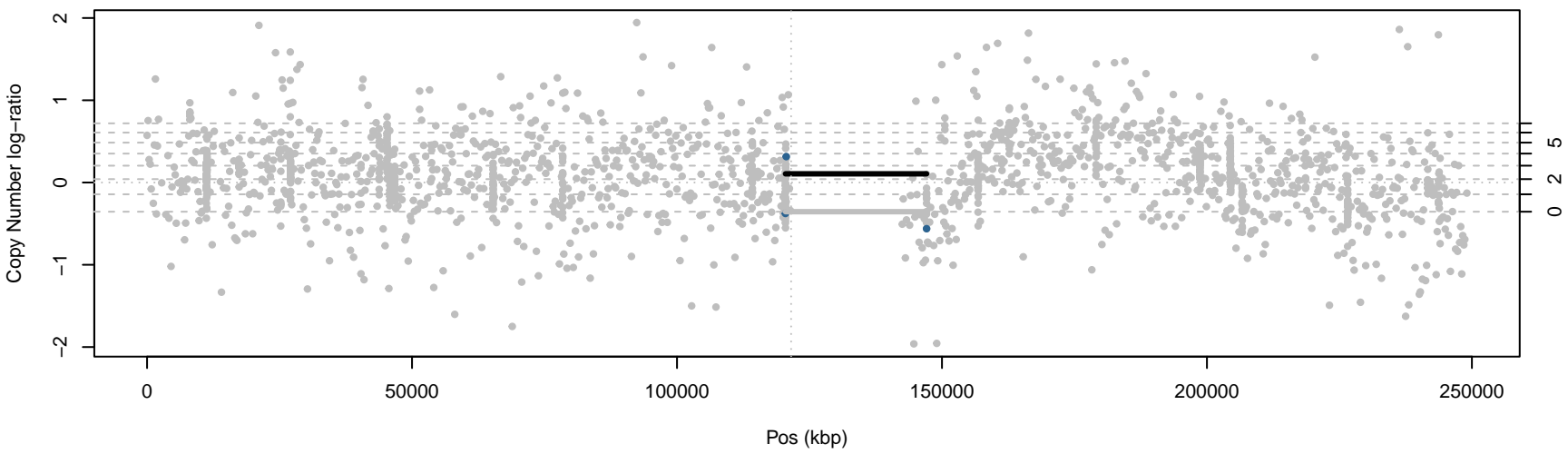
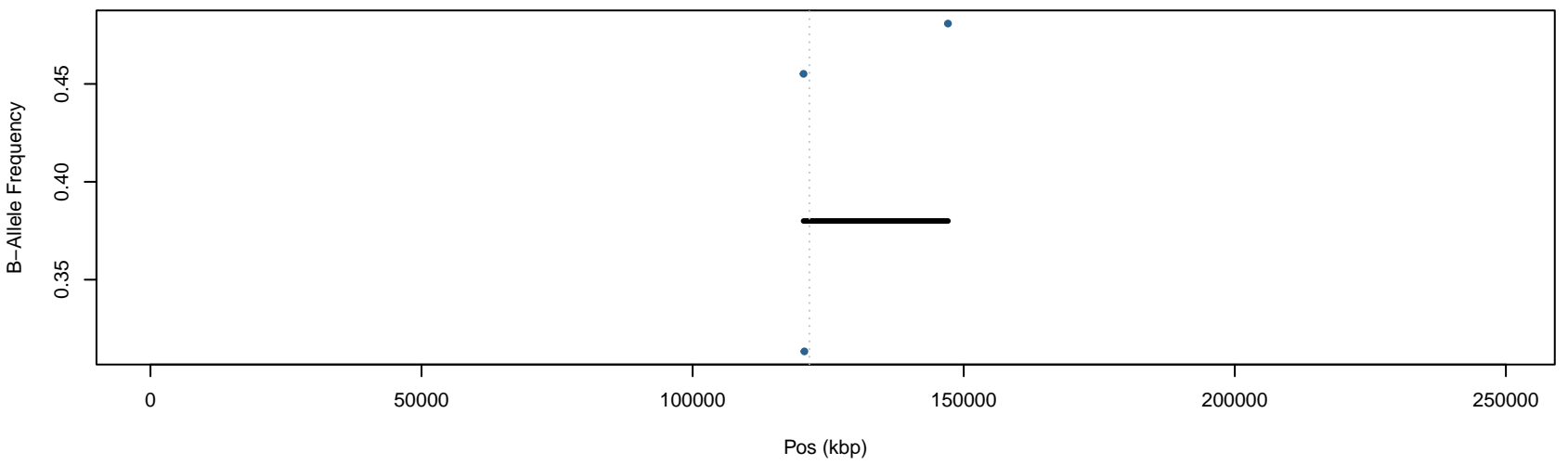
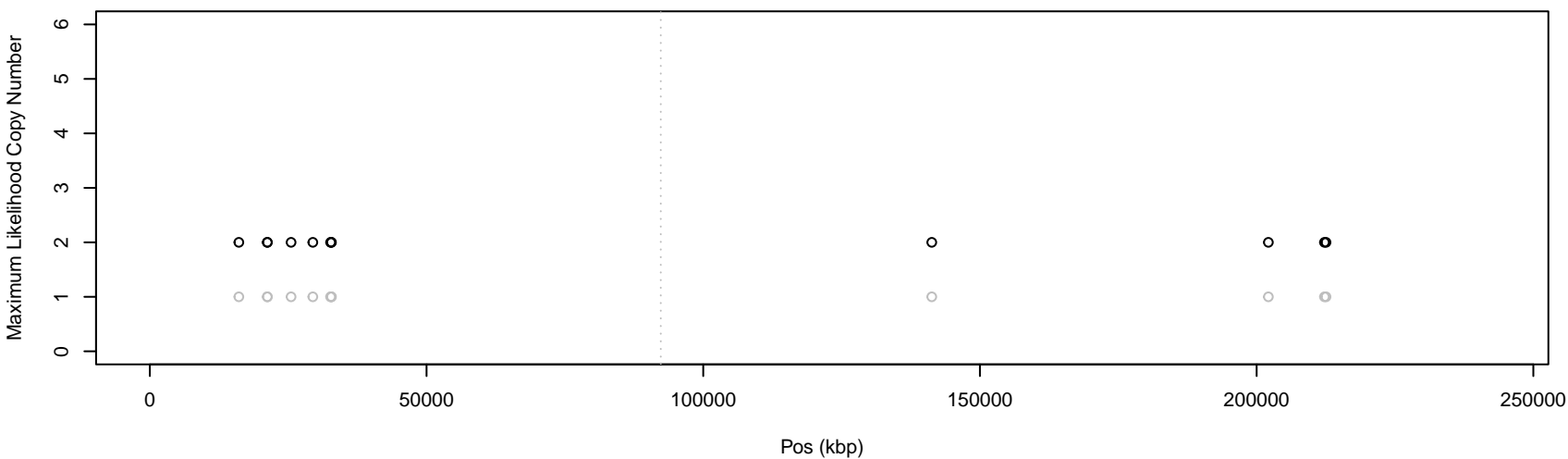
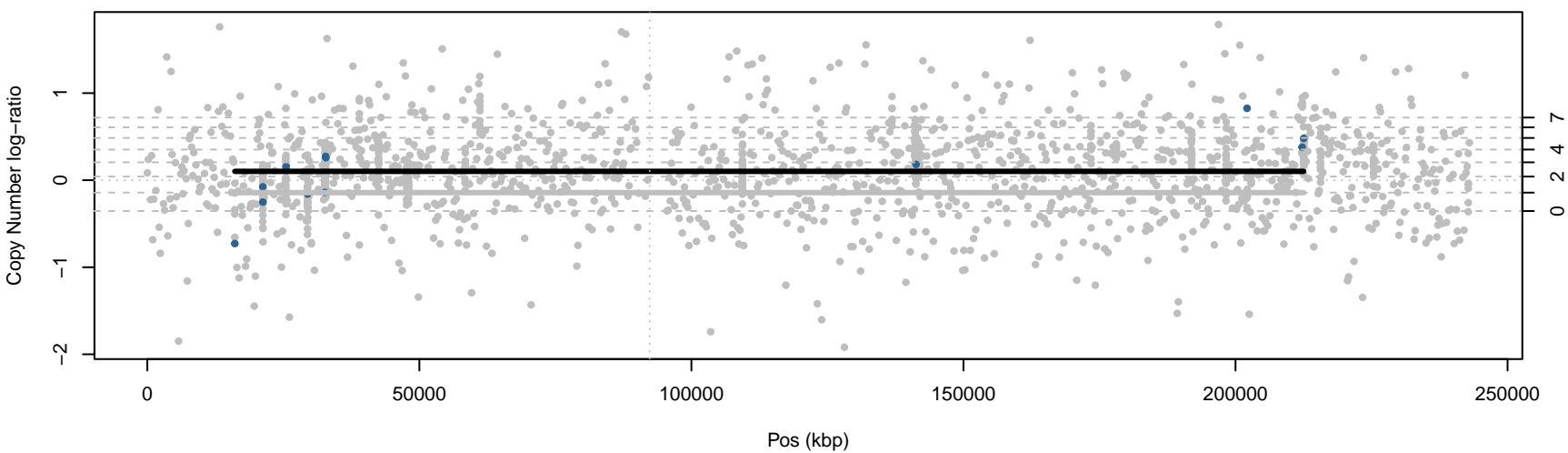
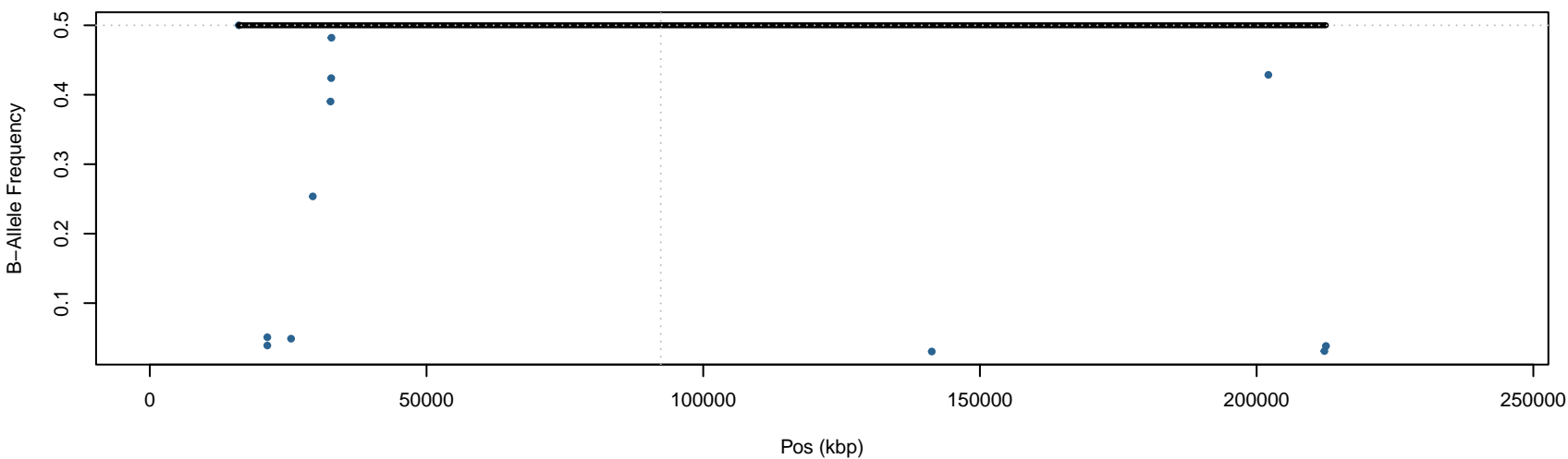


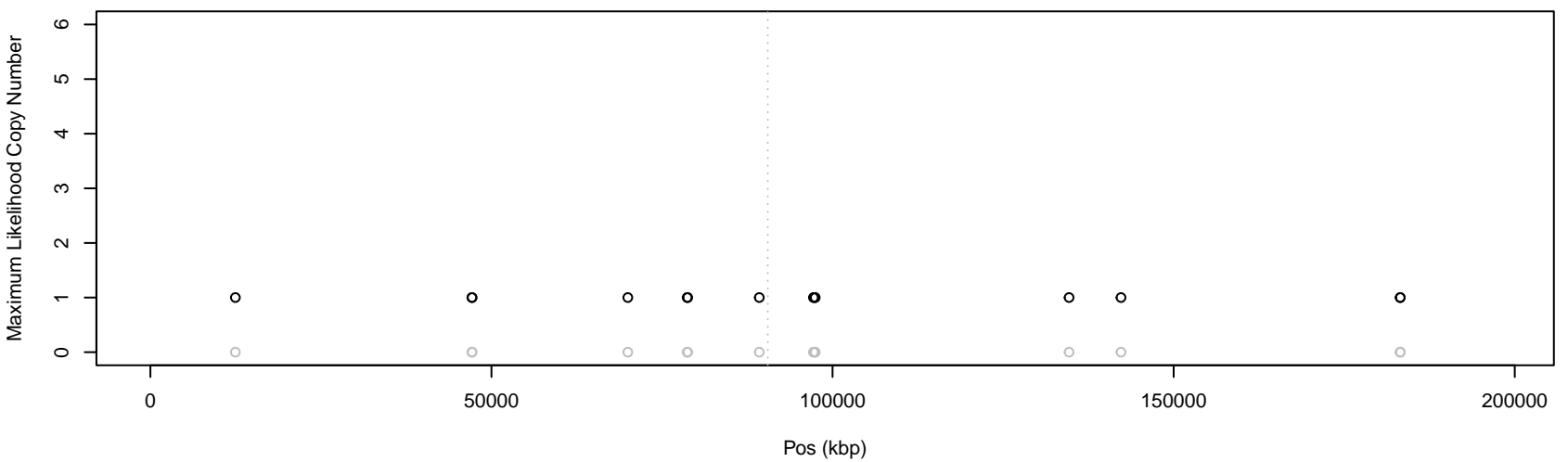
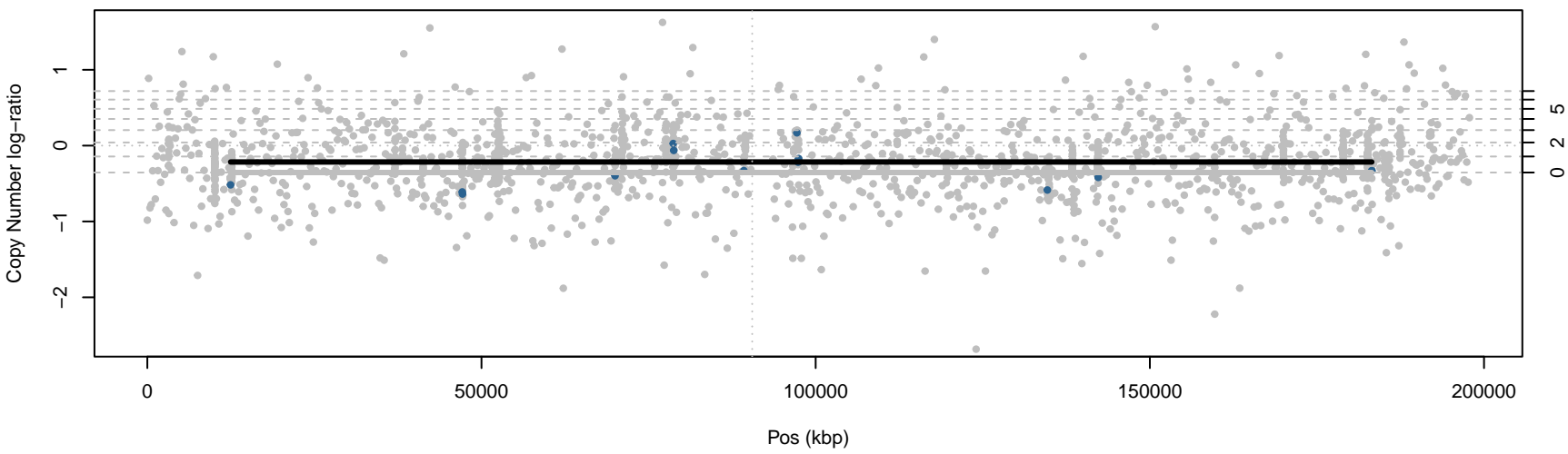
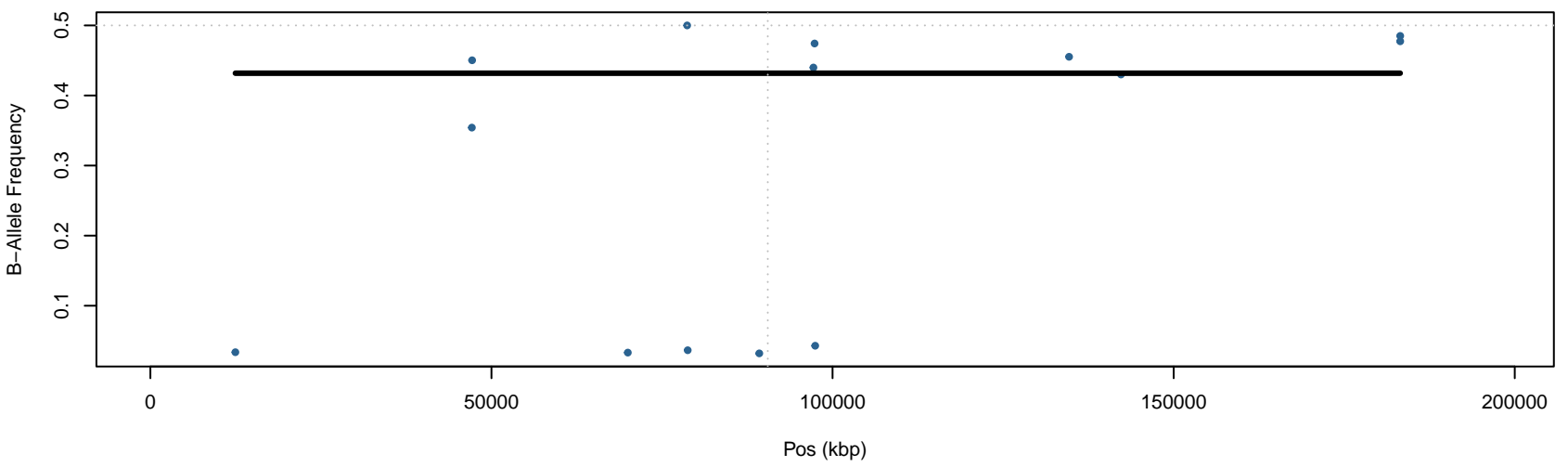
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 1



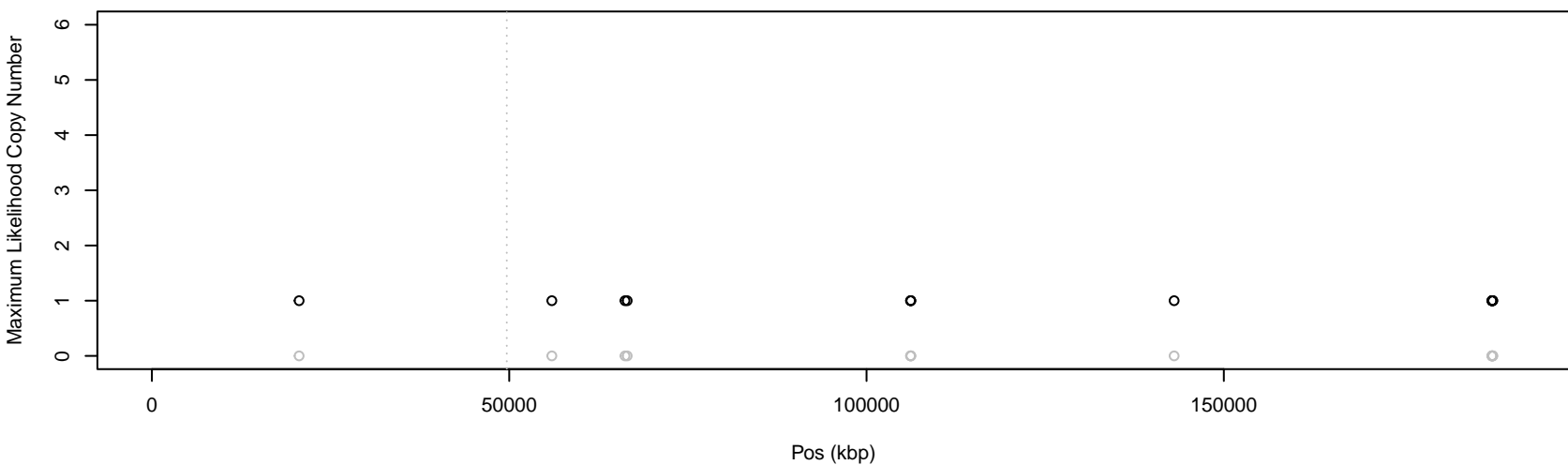
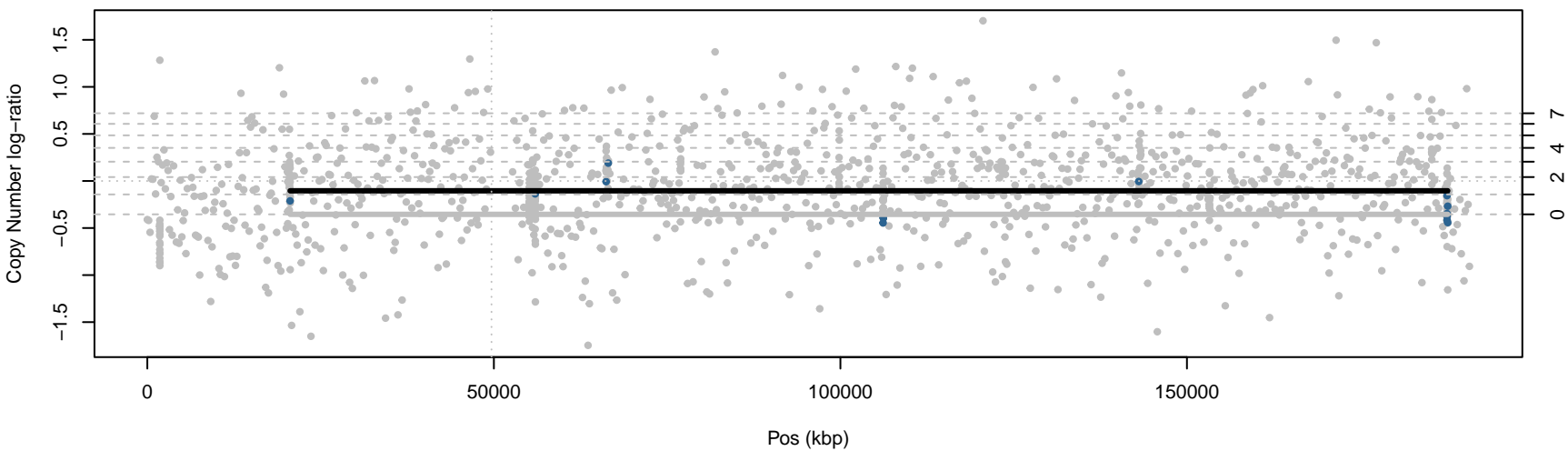
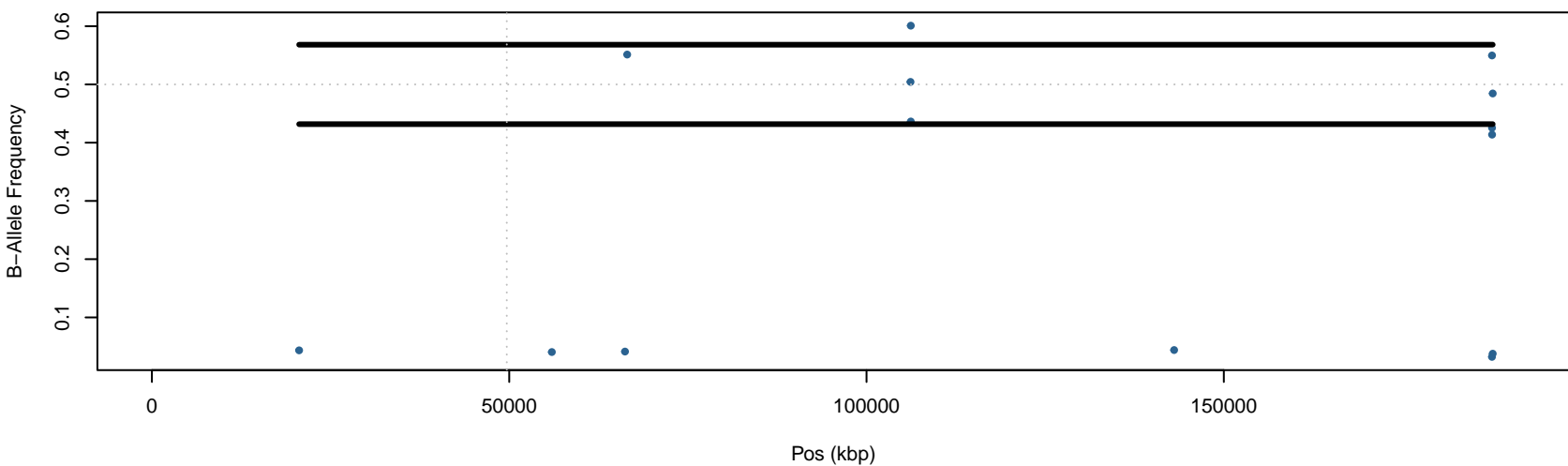
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 2



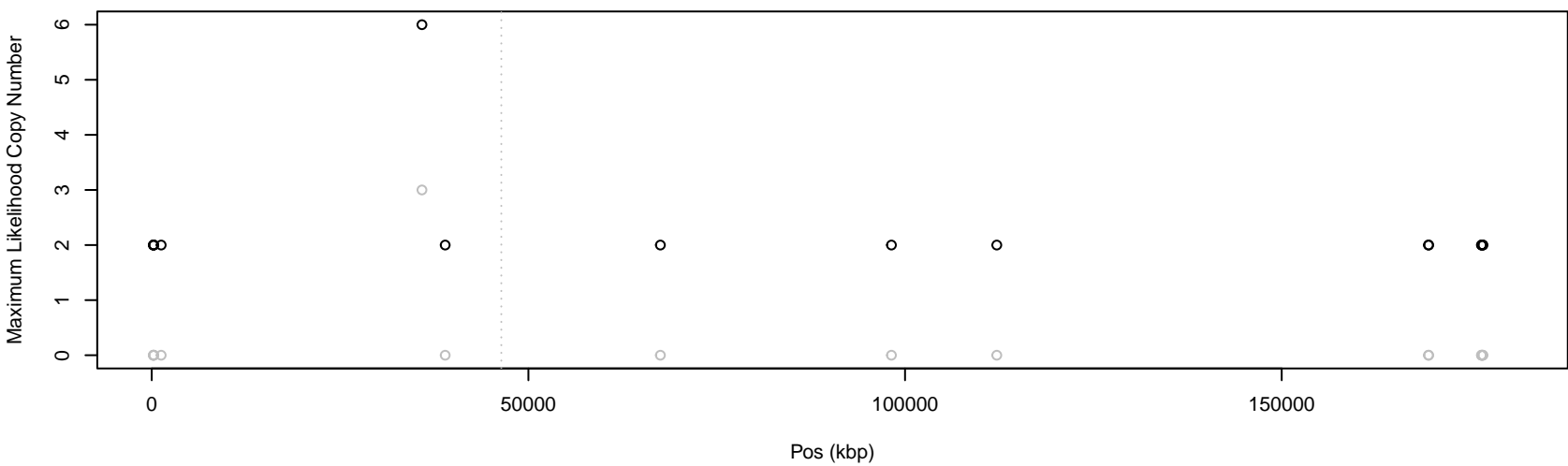
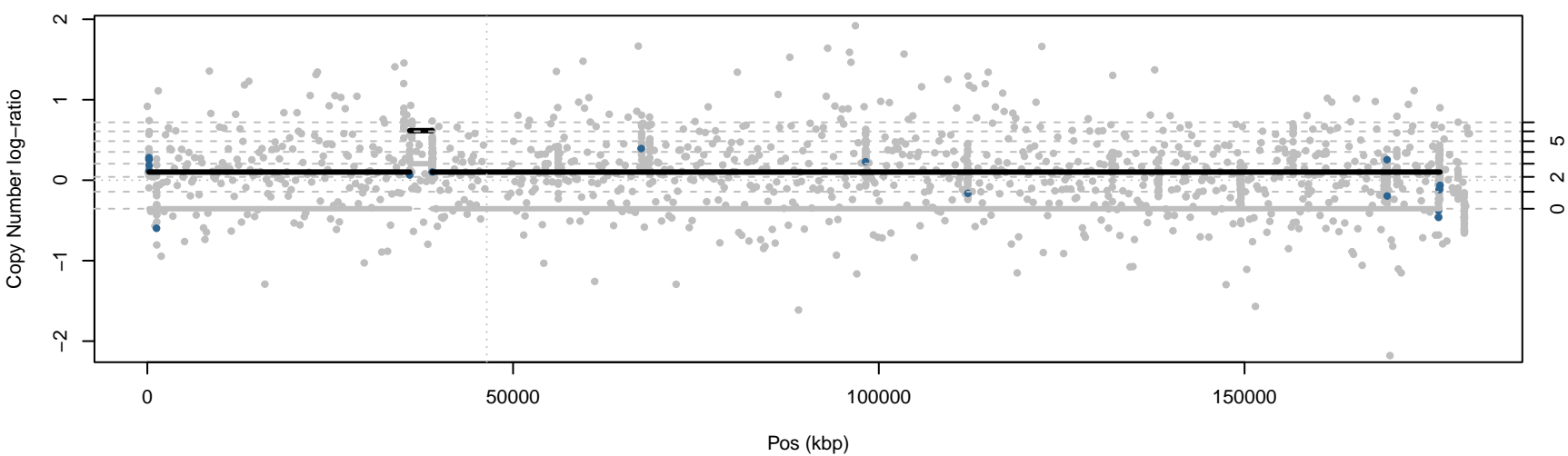
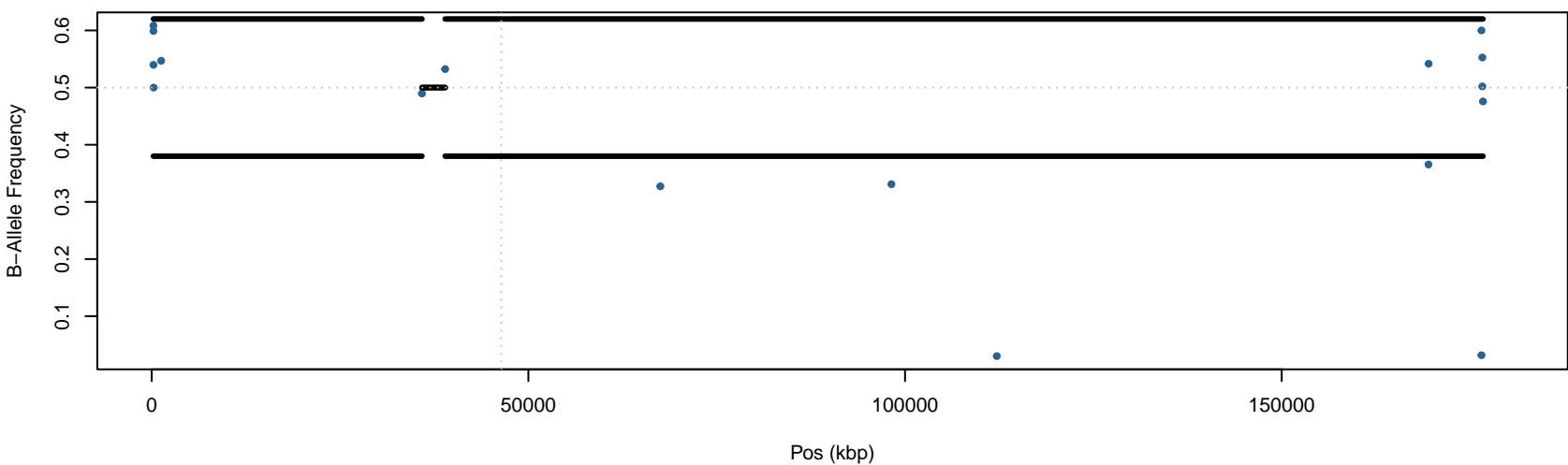
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 3



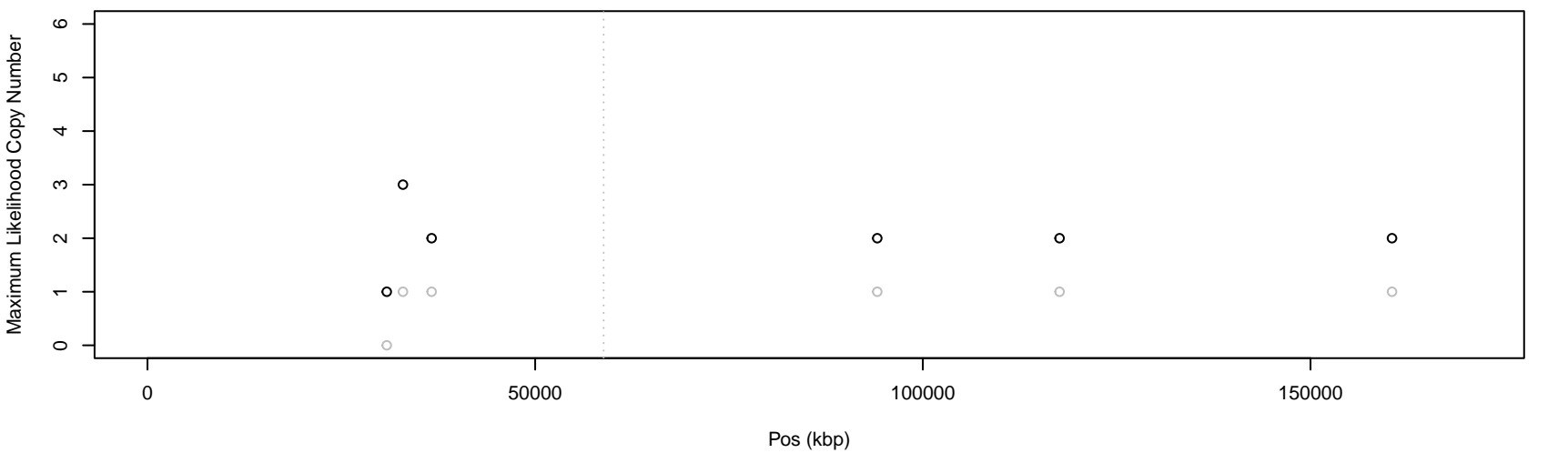
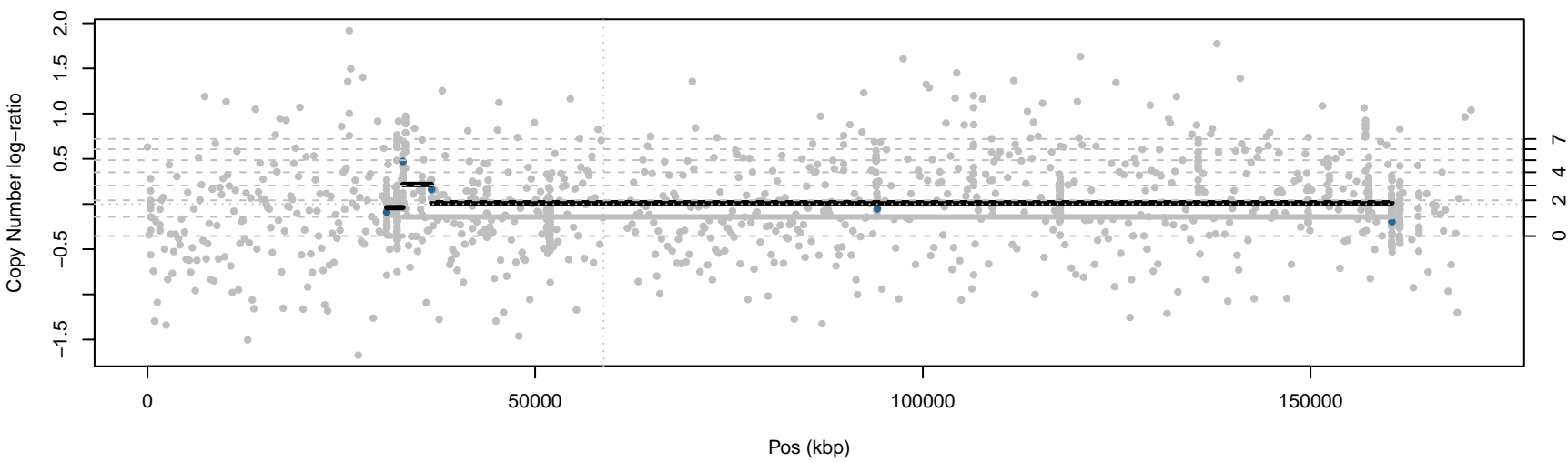
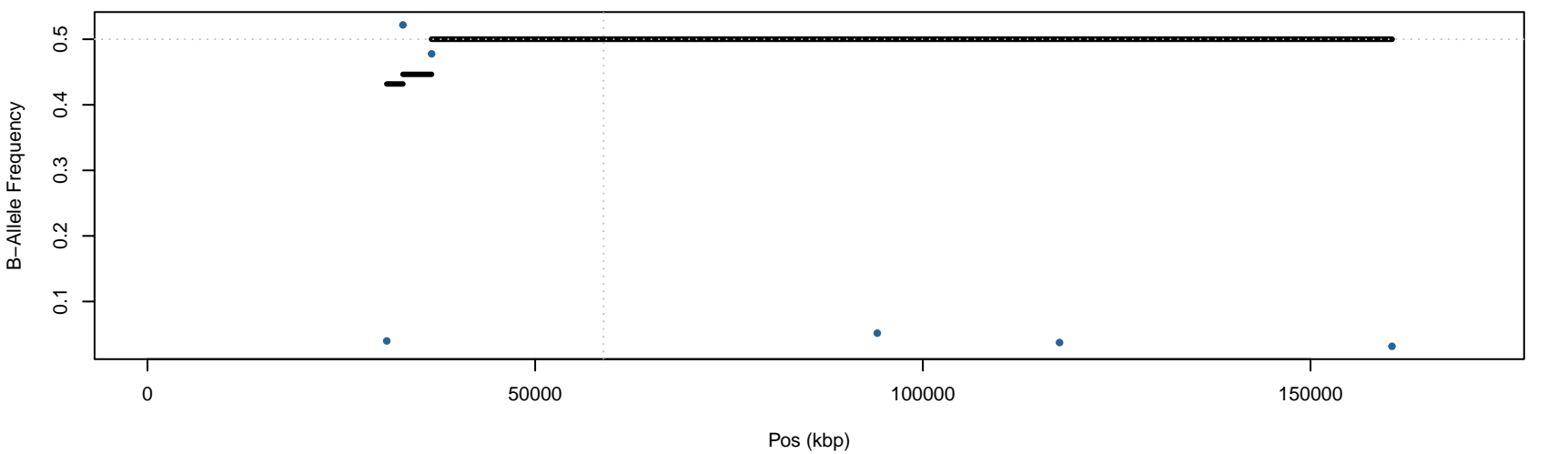
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 4



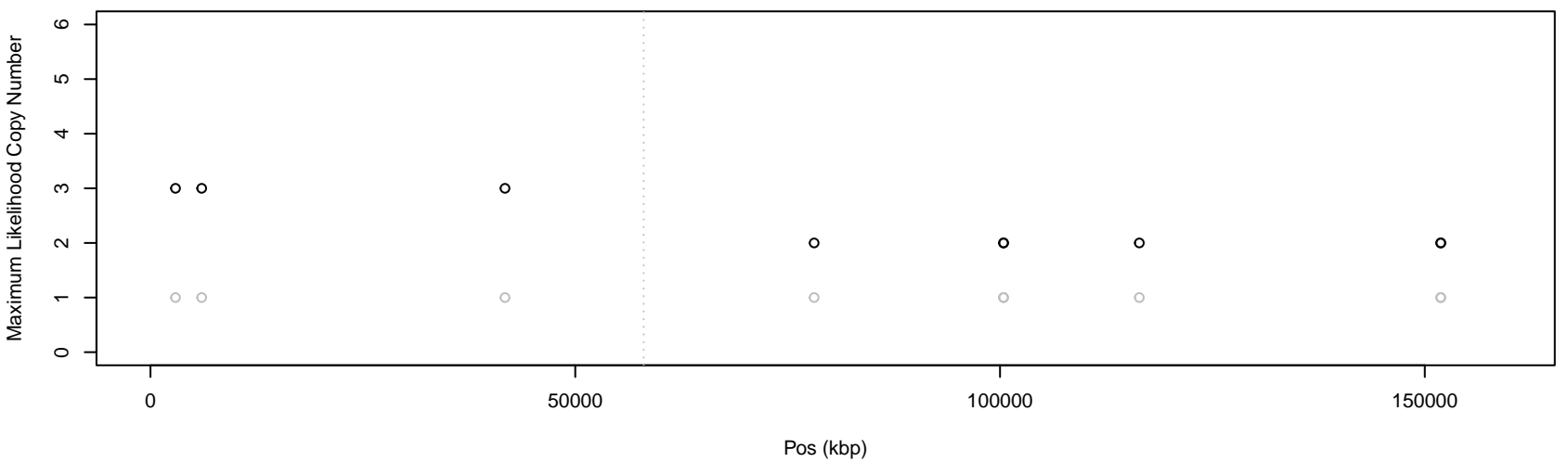
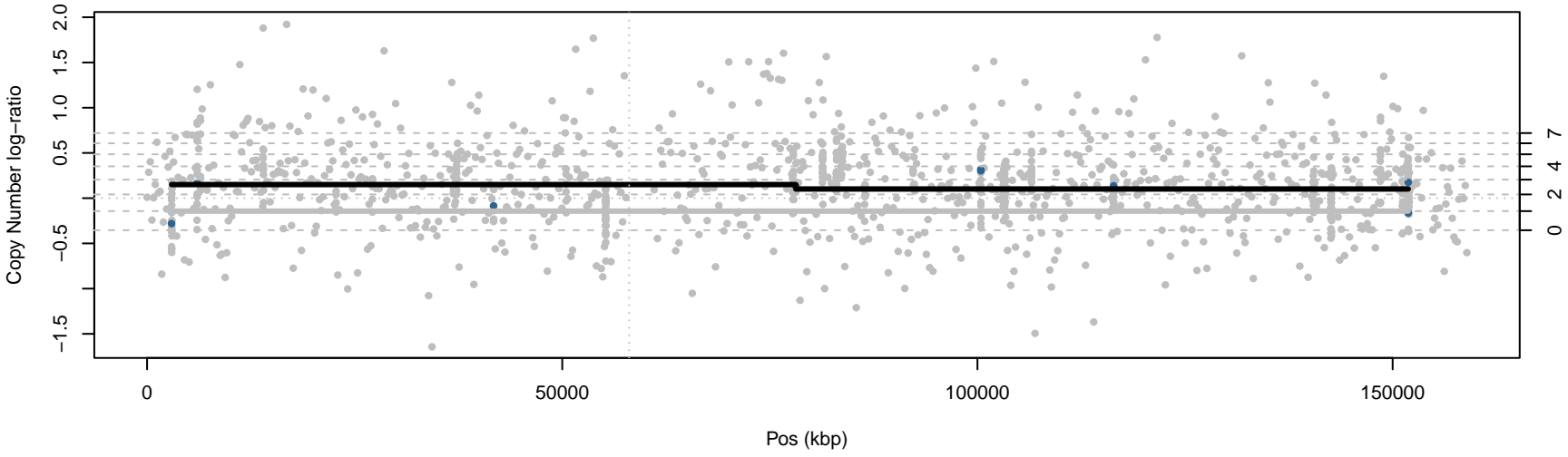
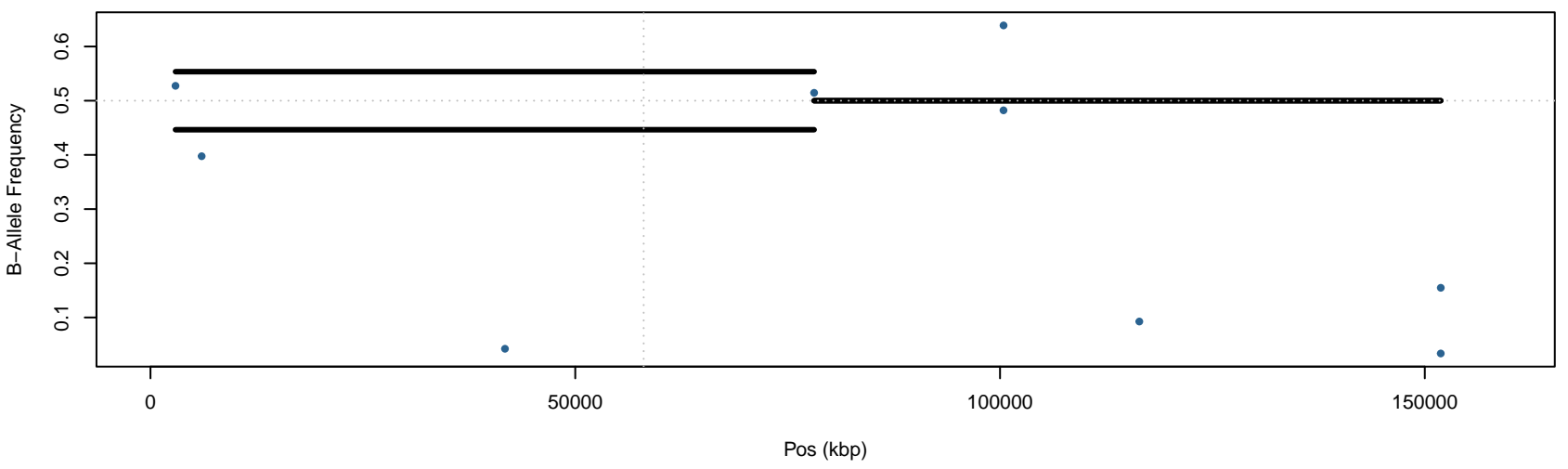
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 5



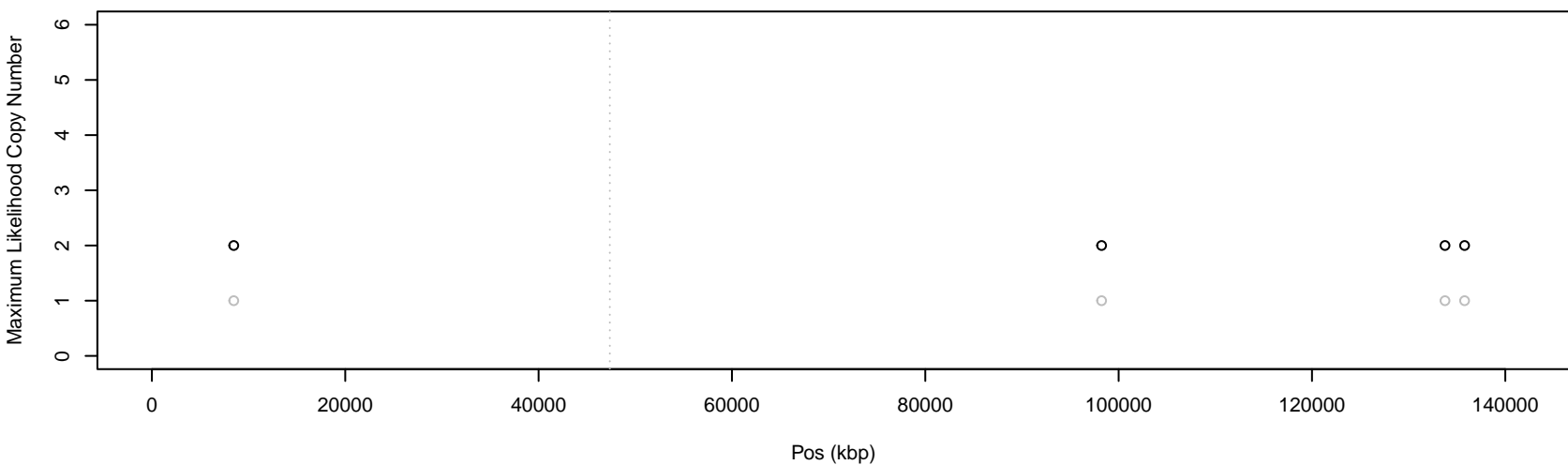
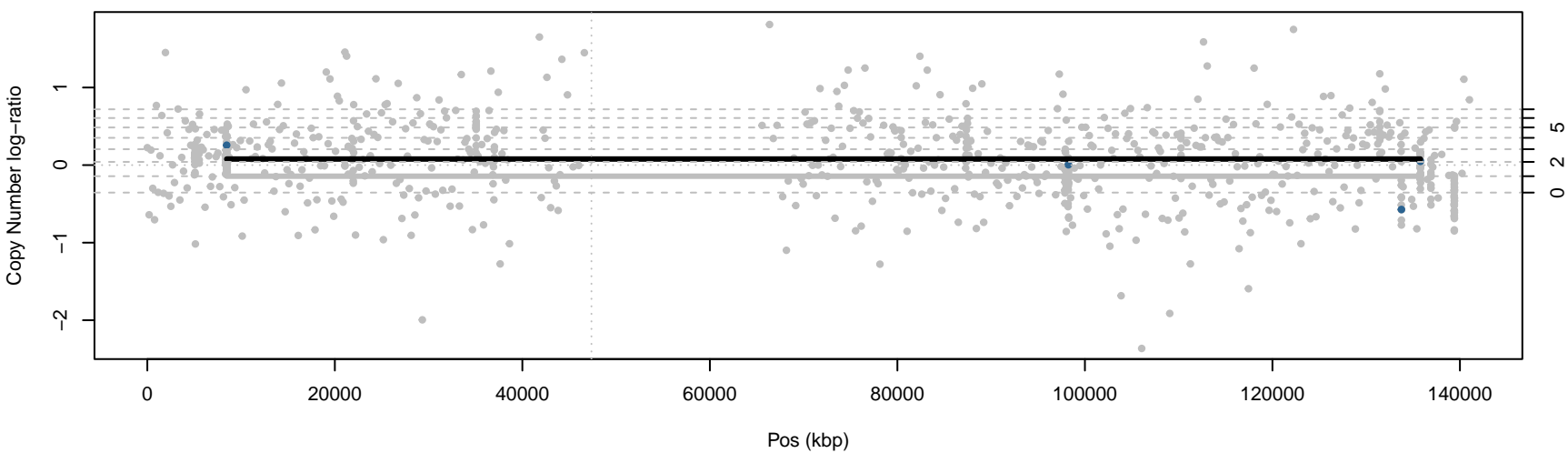
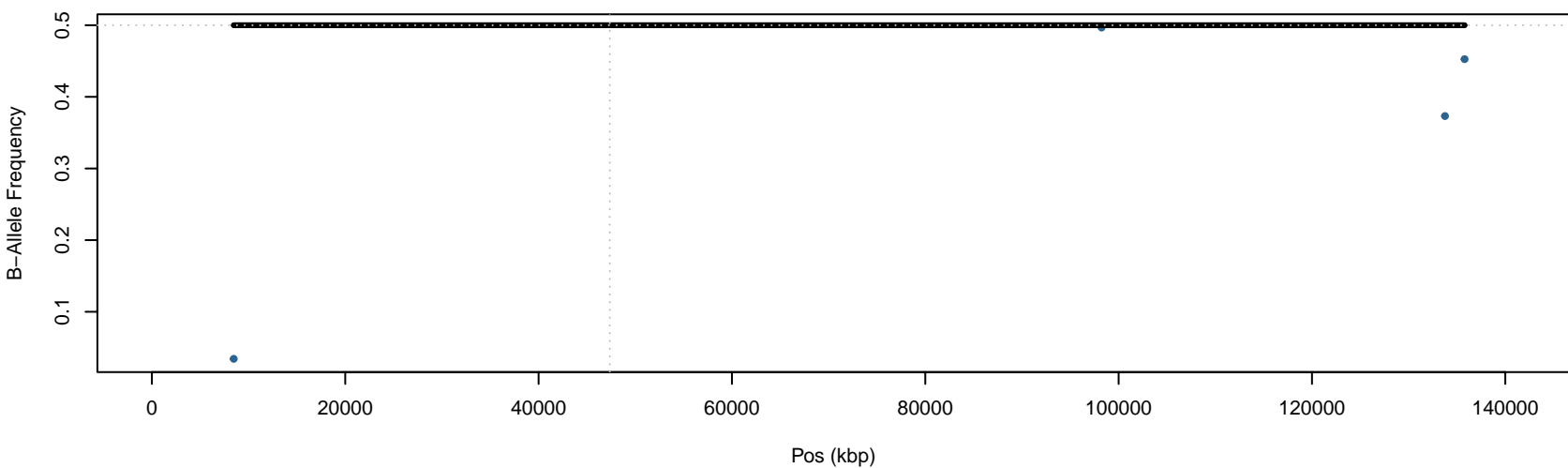
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 6



Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 7

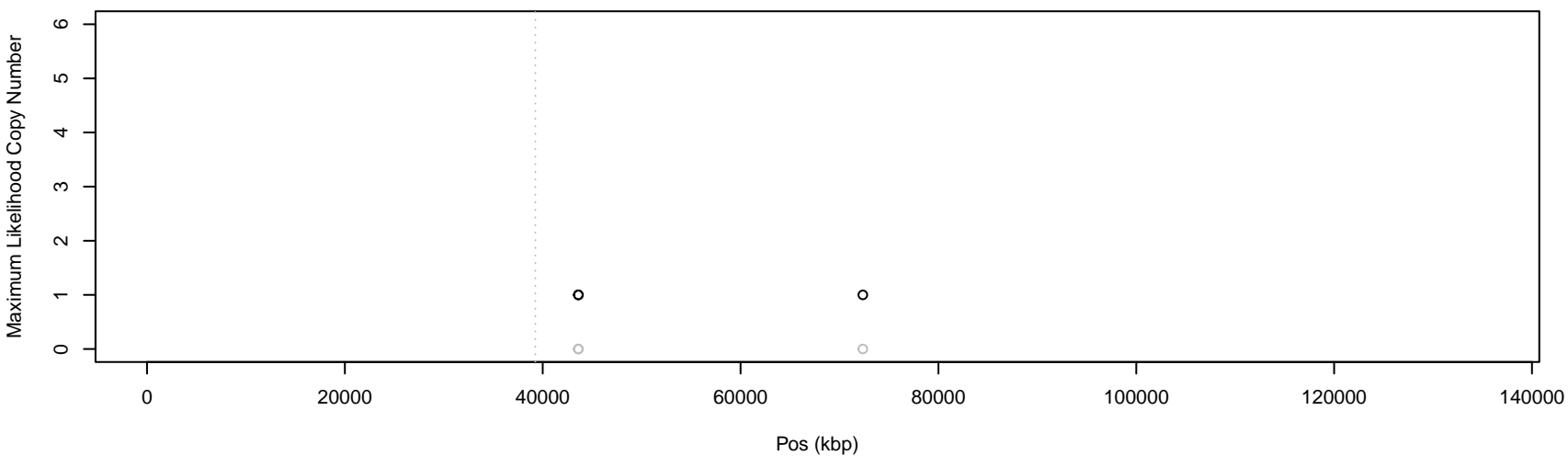
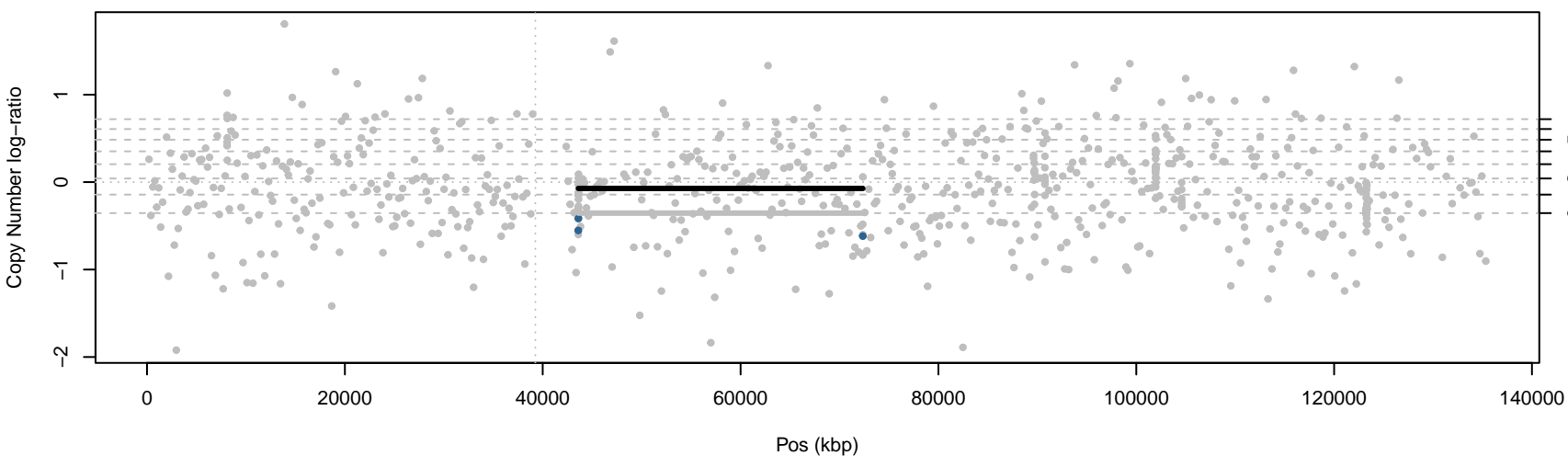
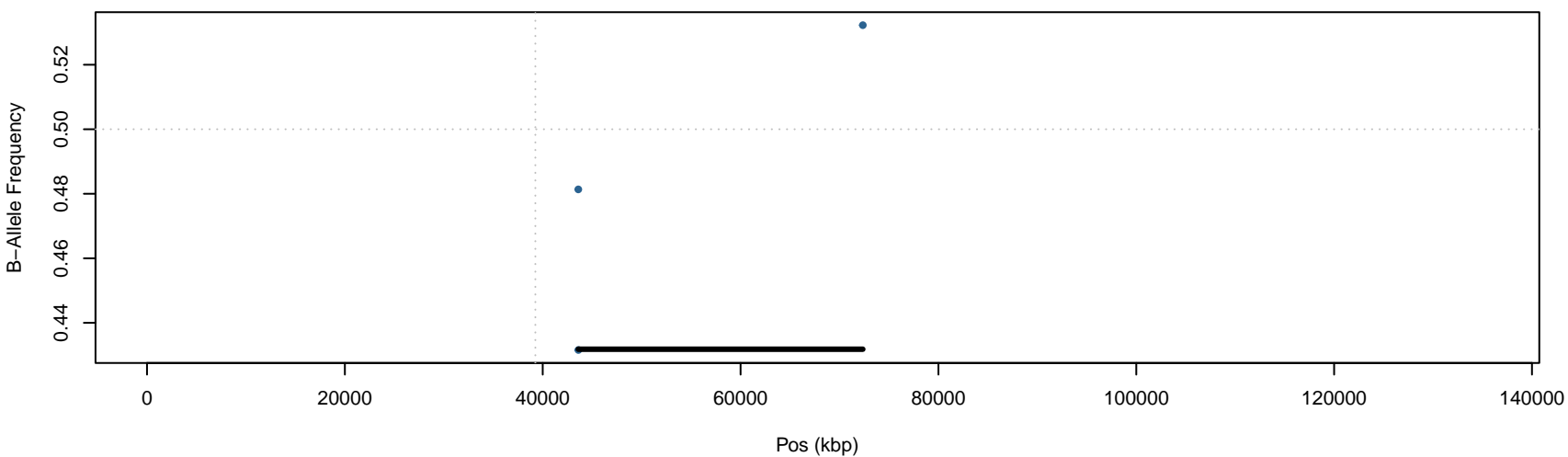


Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 9

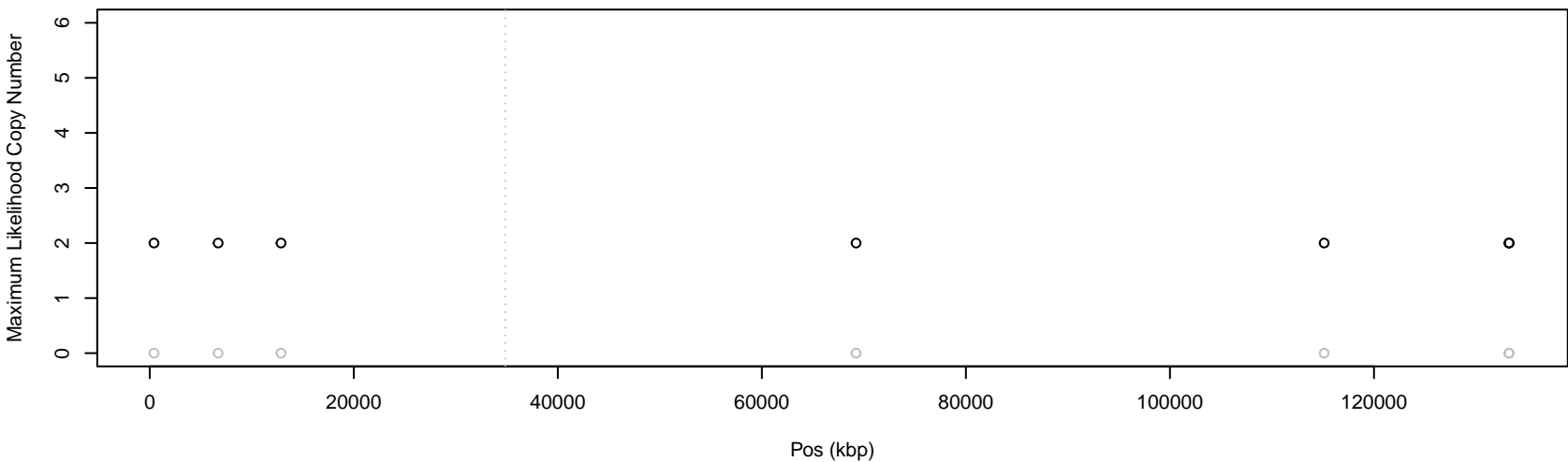
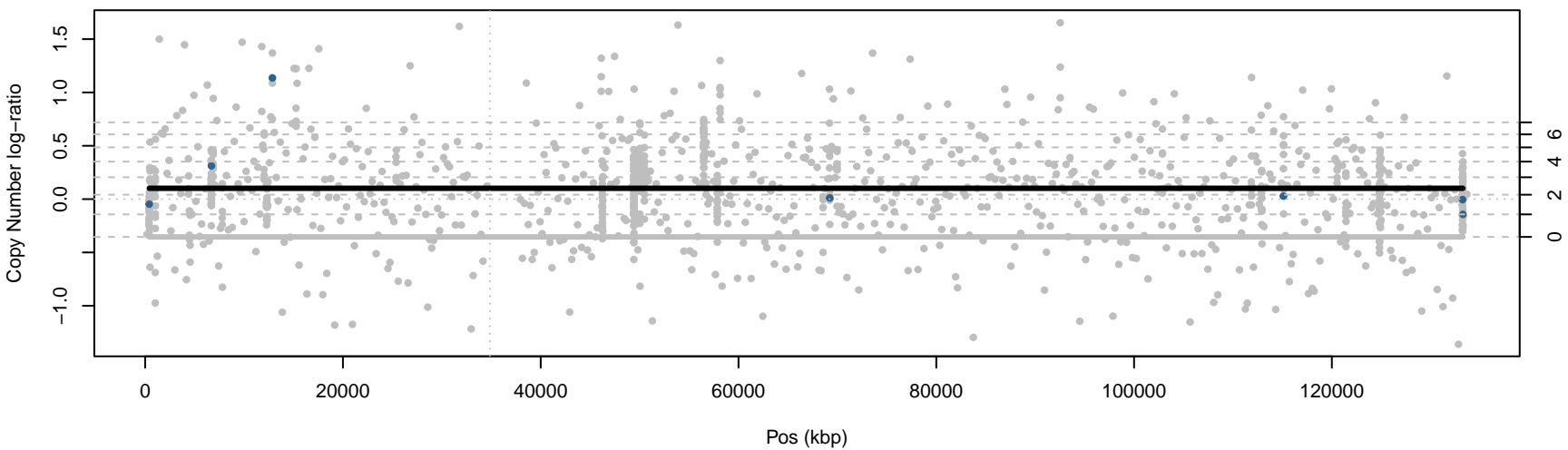
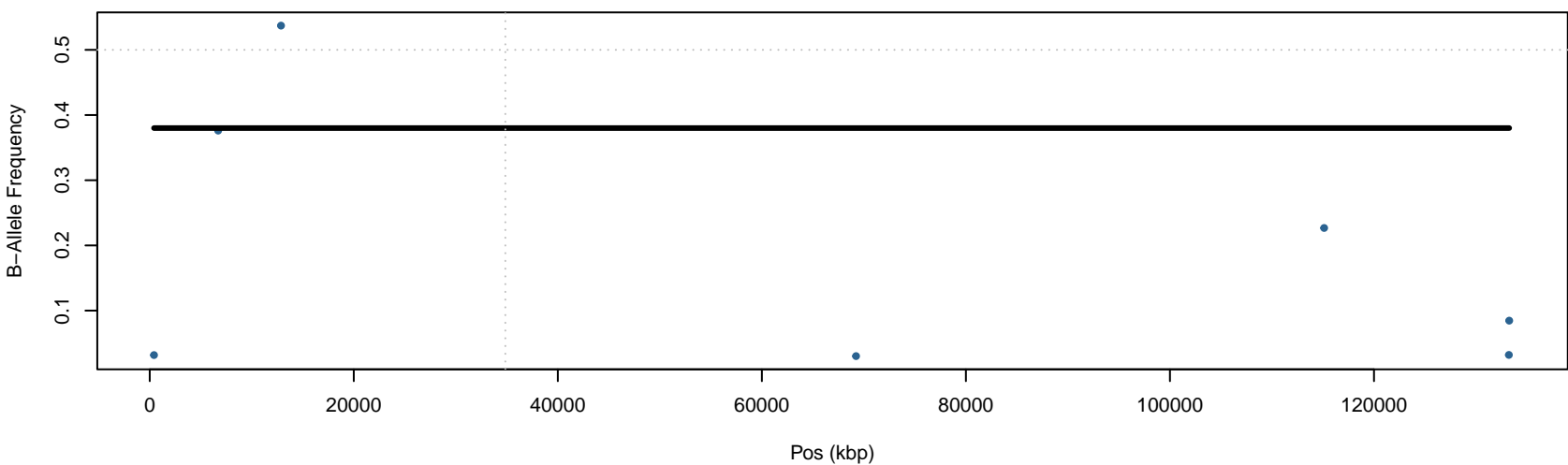




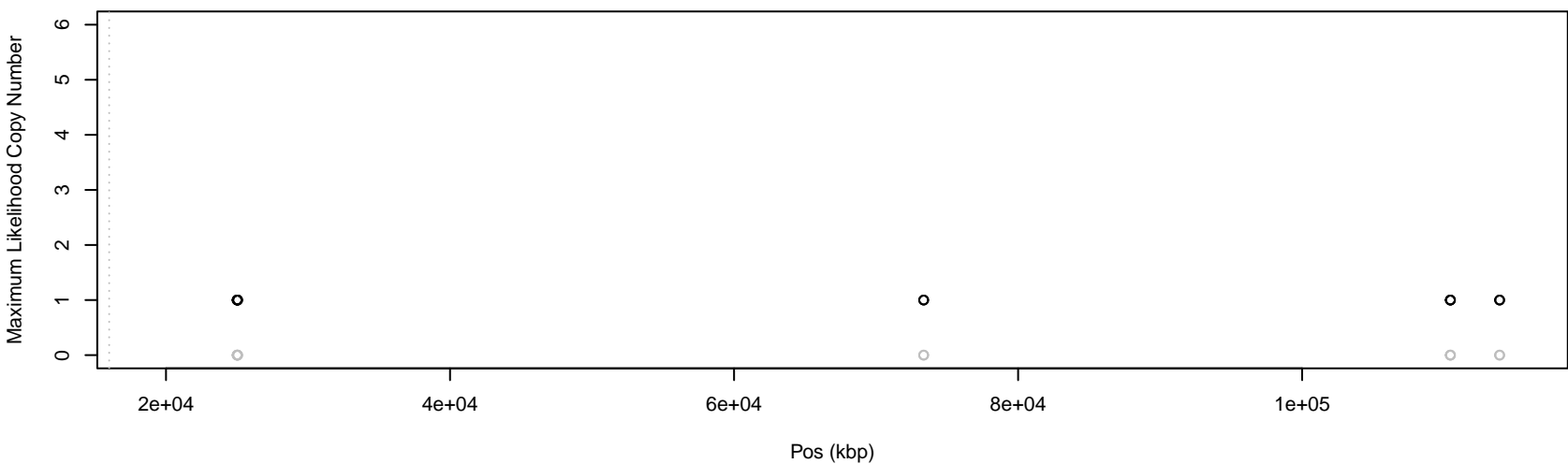
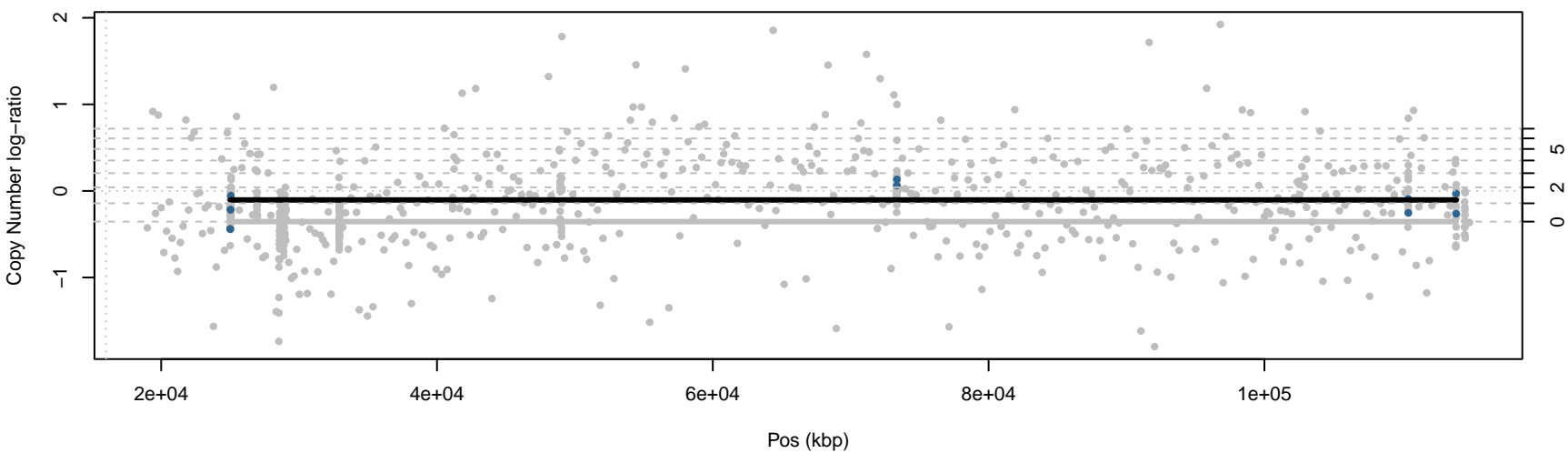
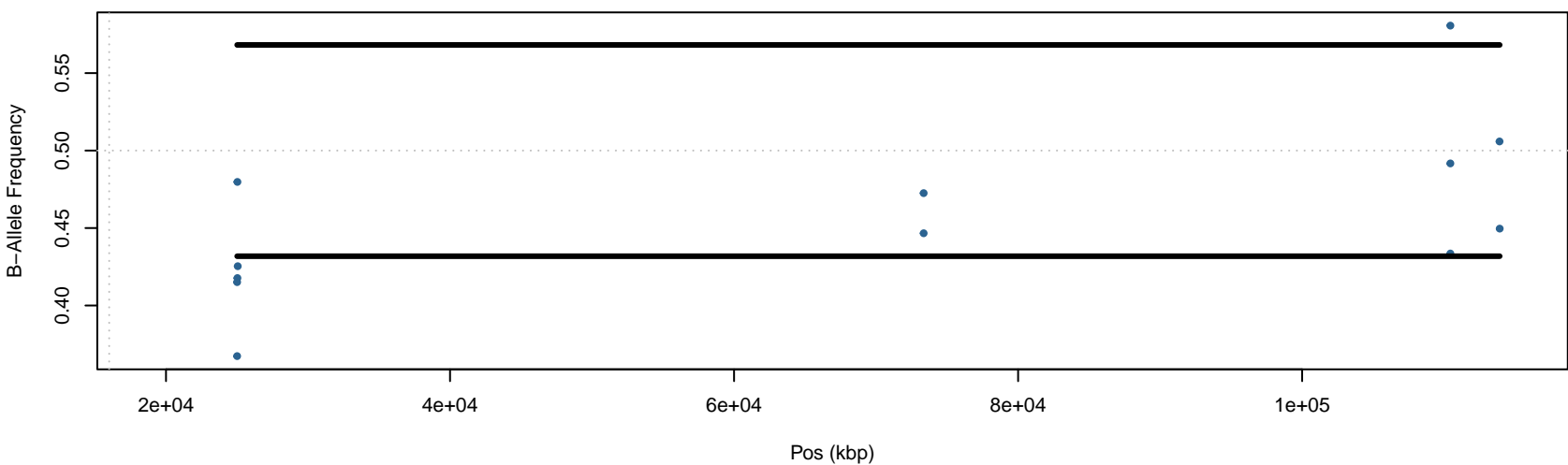
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 10



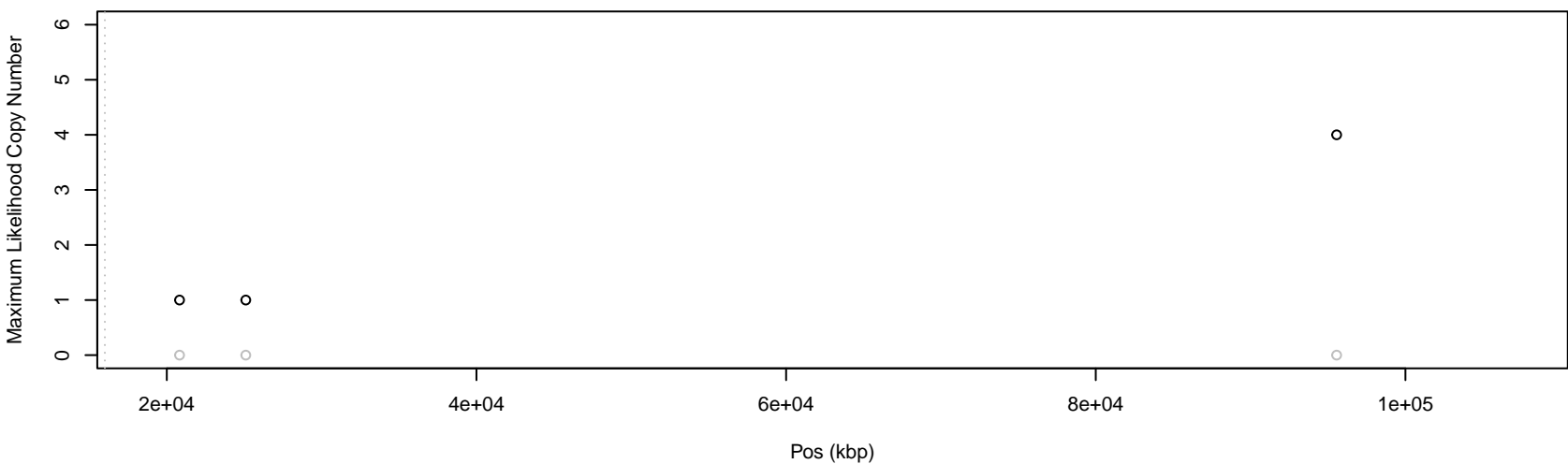
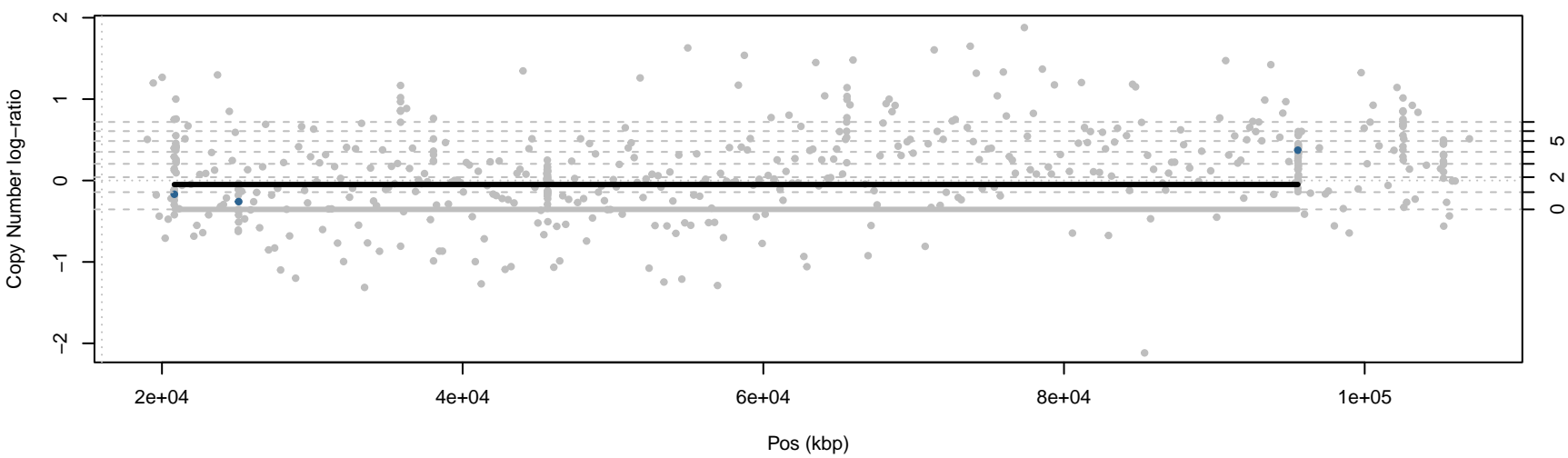
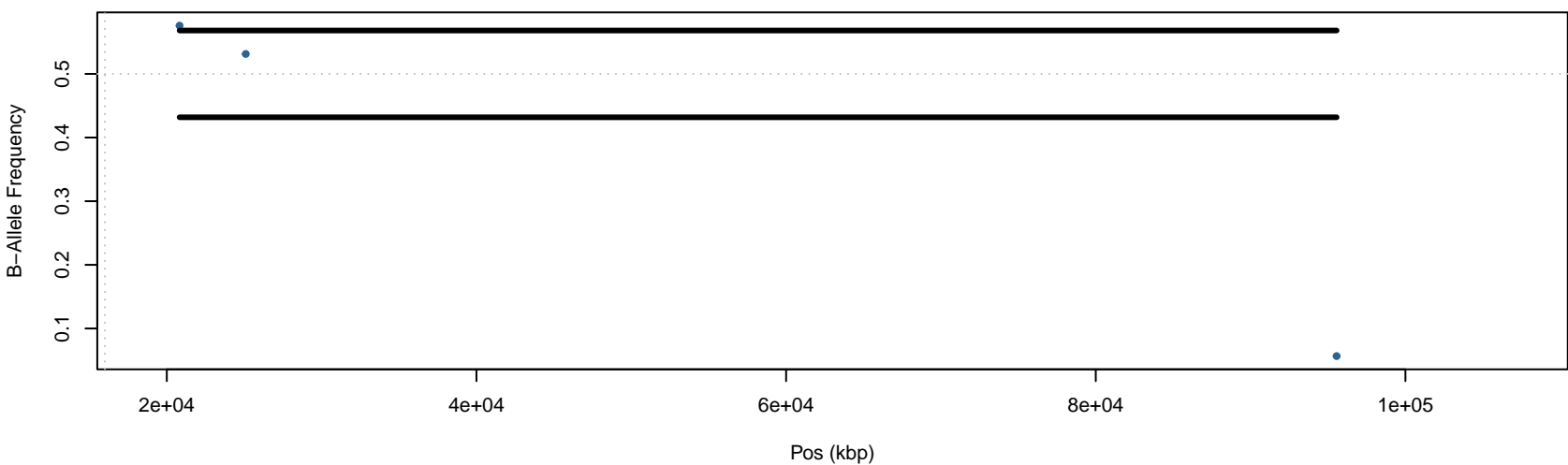
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 12



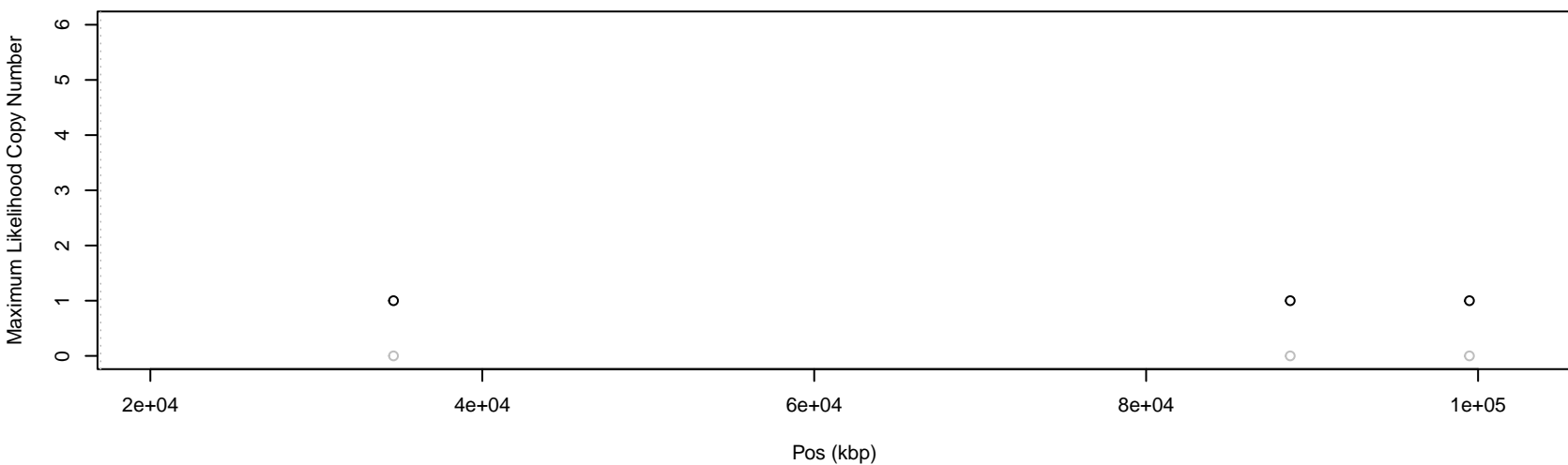
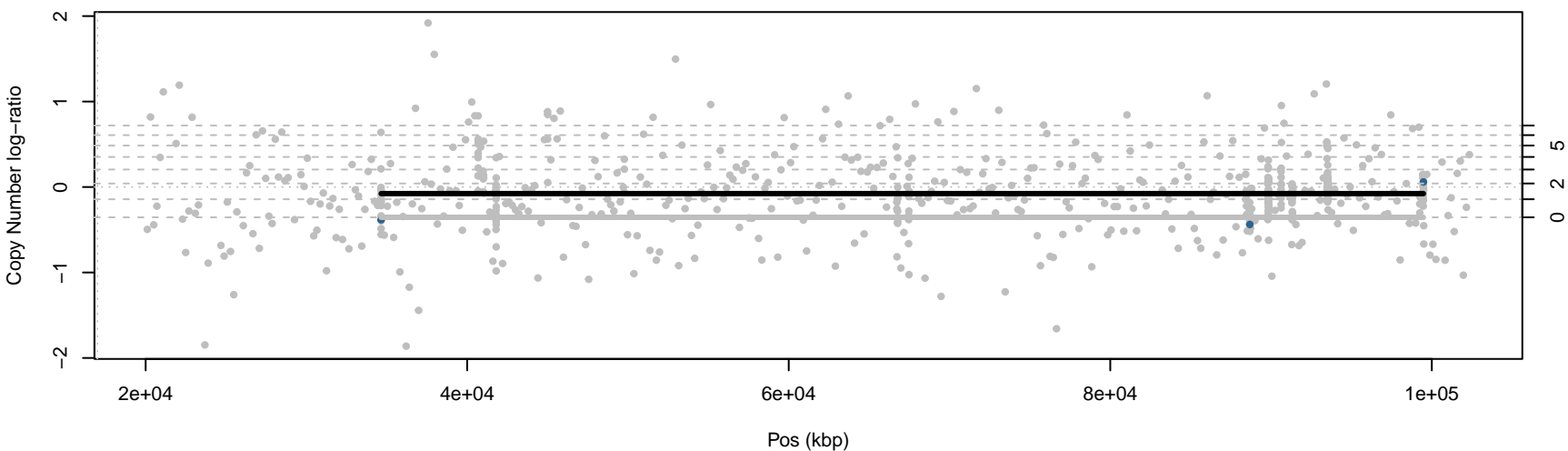
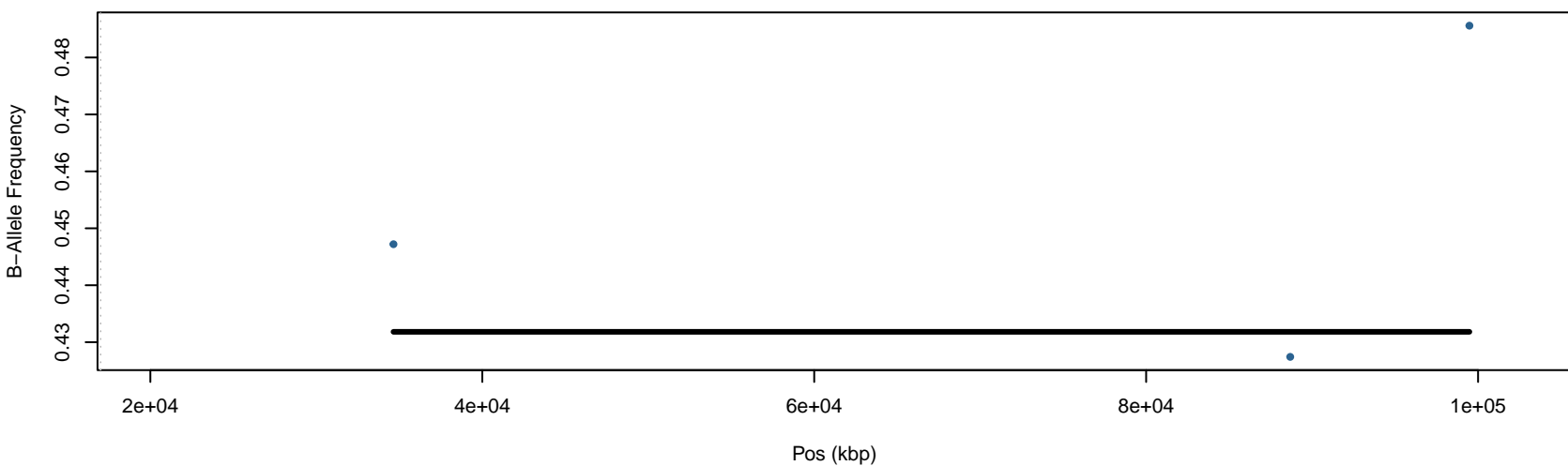
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 13



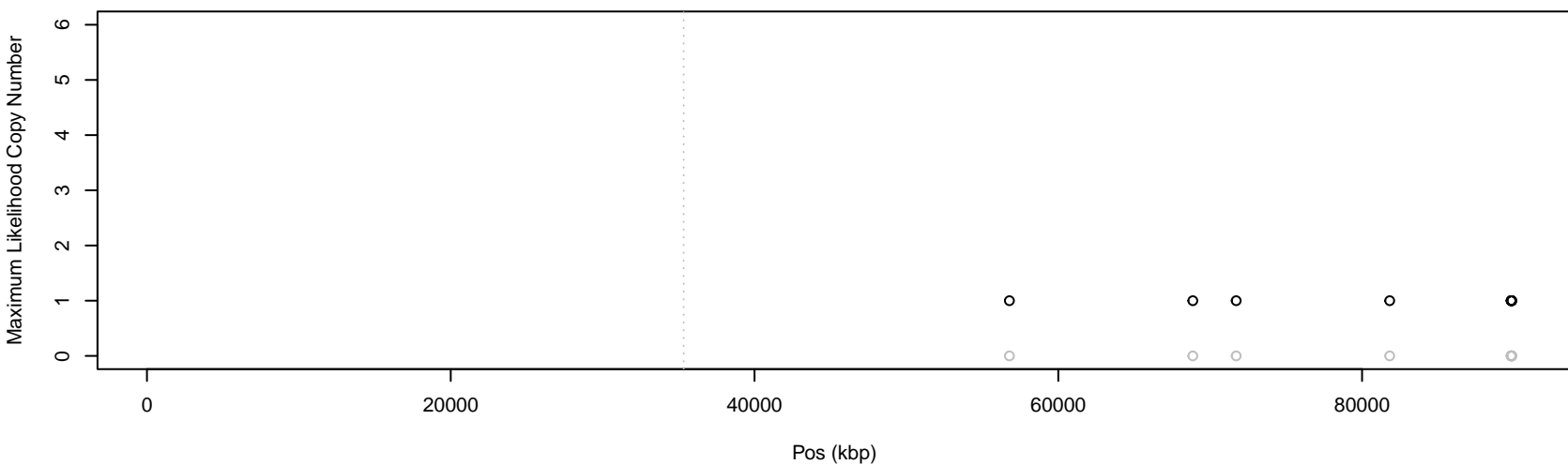
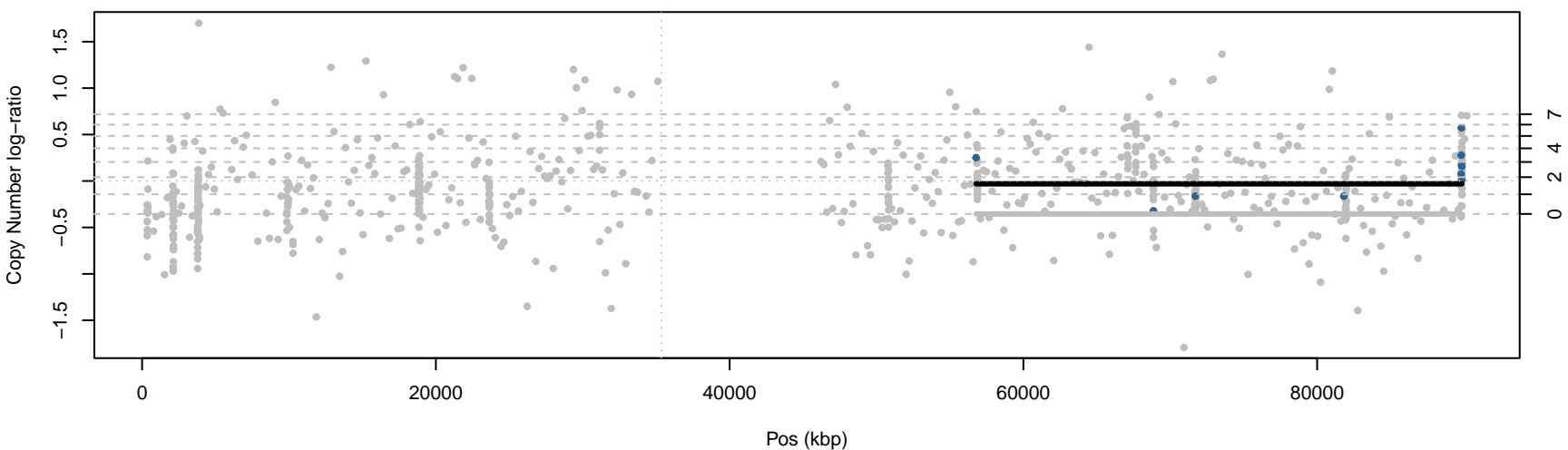
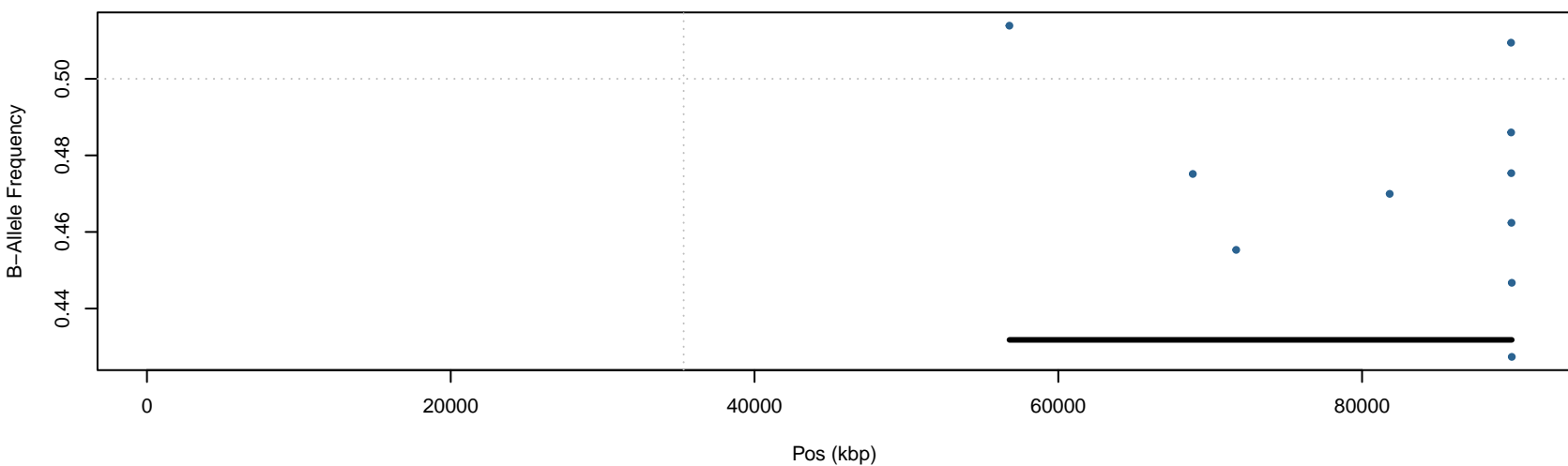
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 14



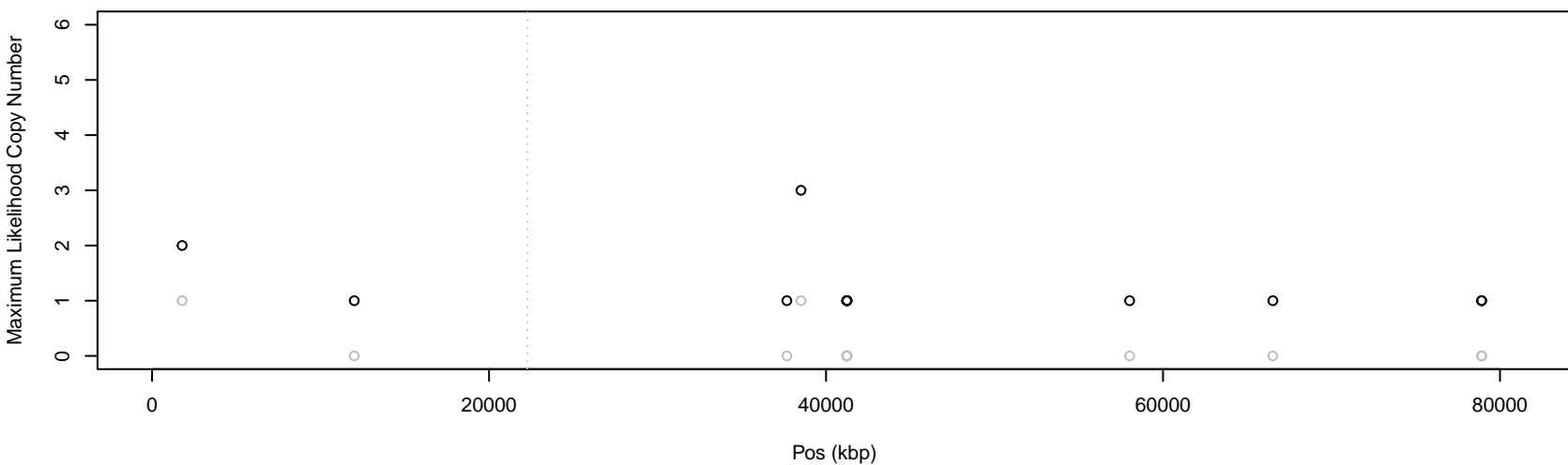
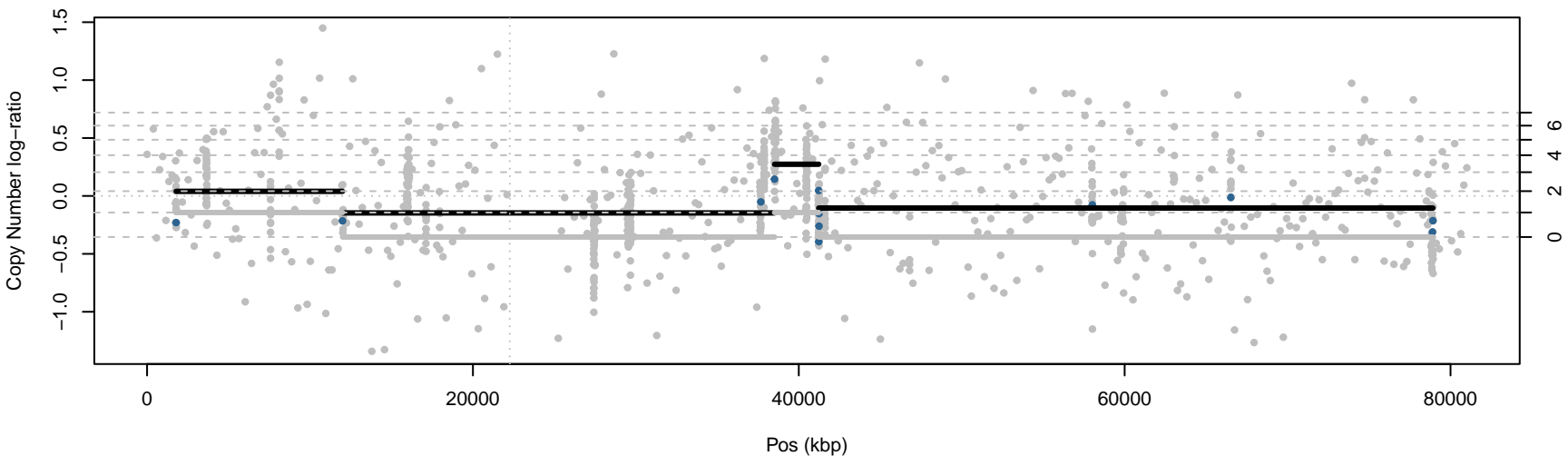
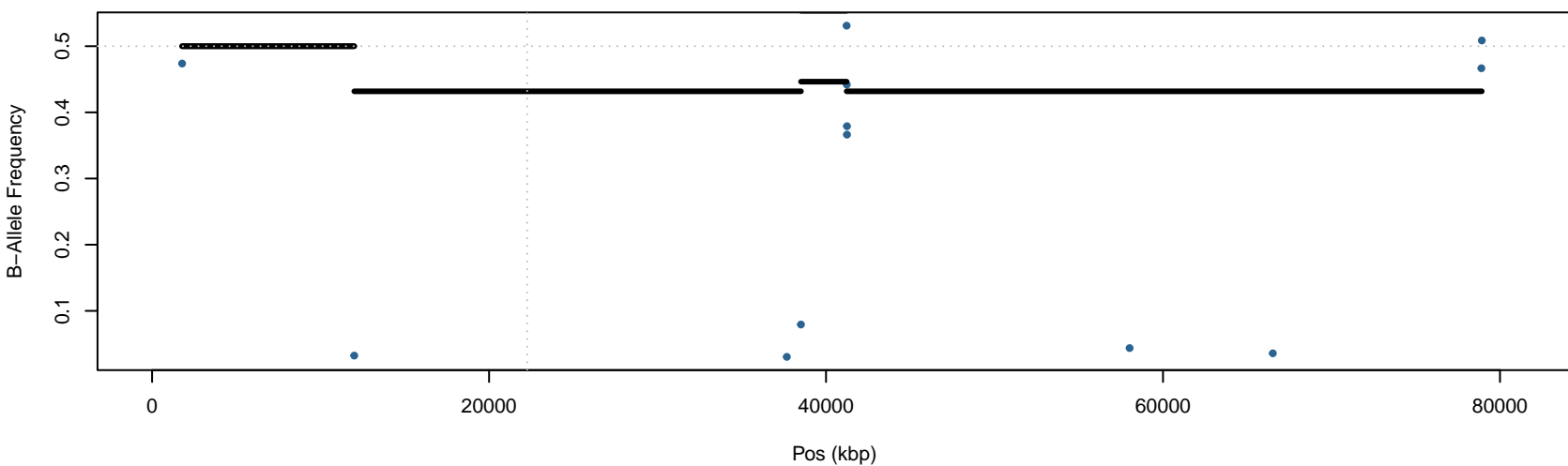
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 15



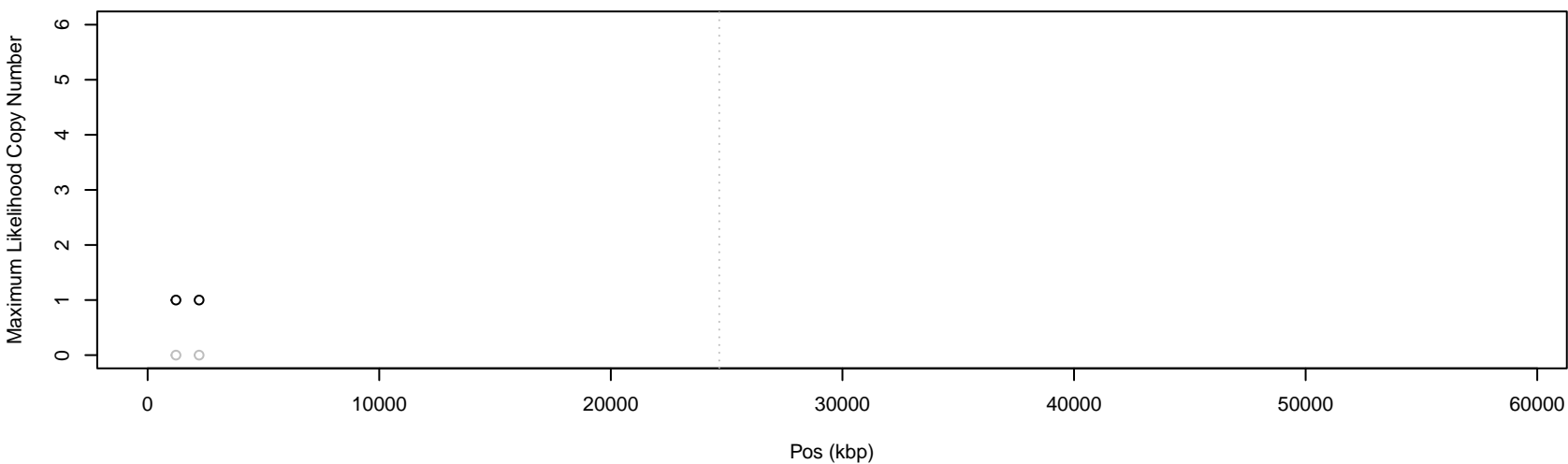
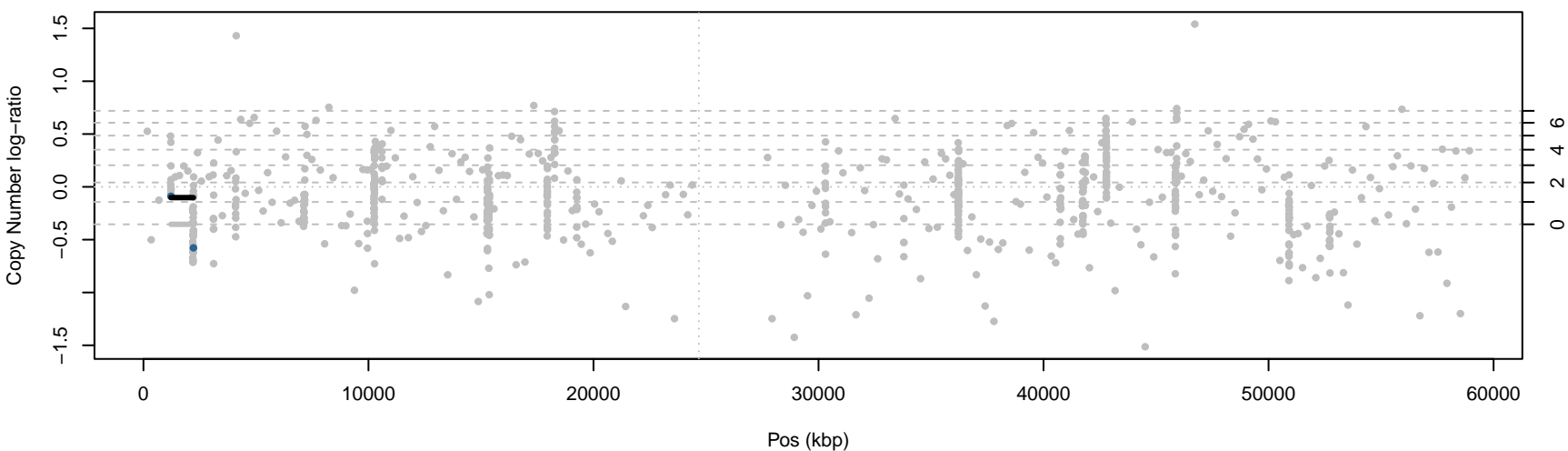
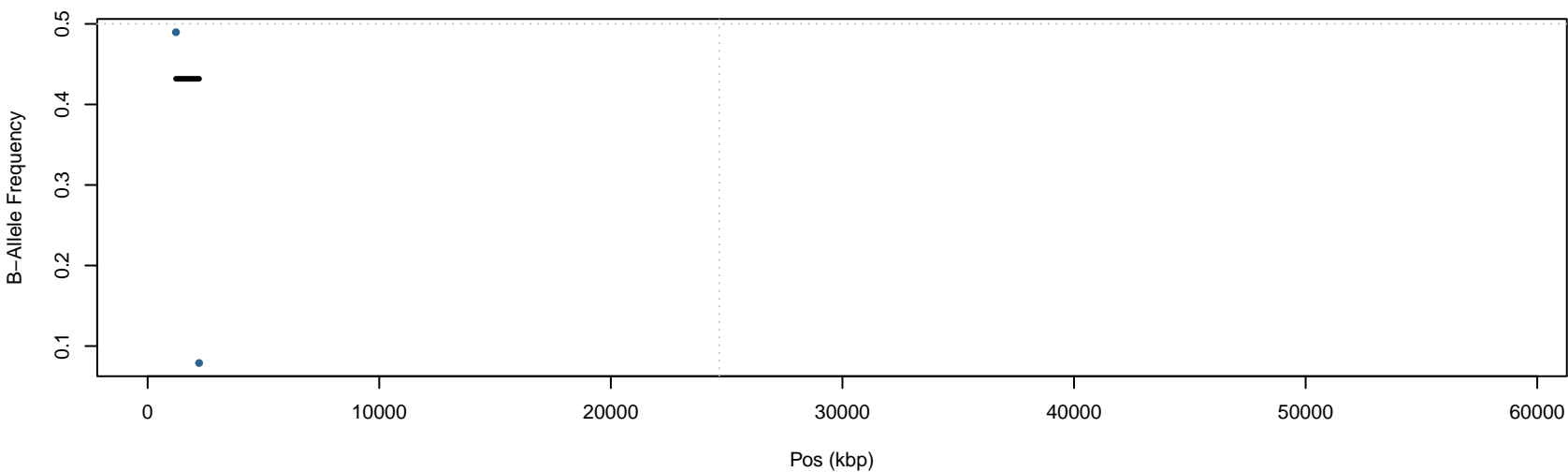
Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 16



Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 17

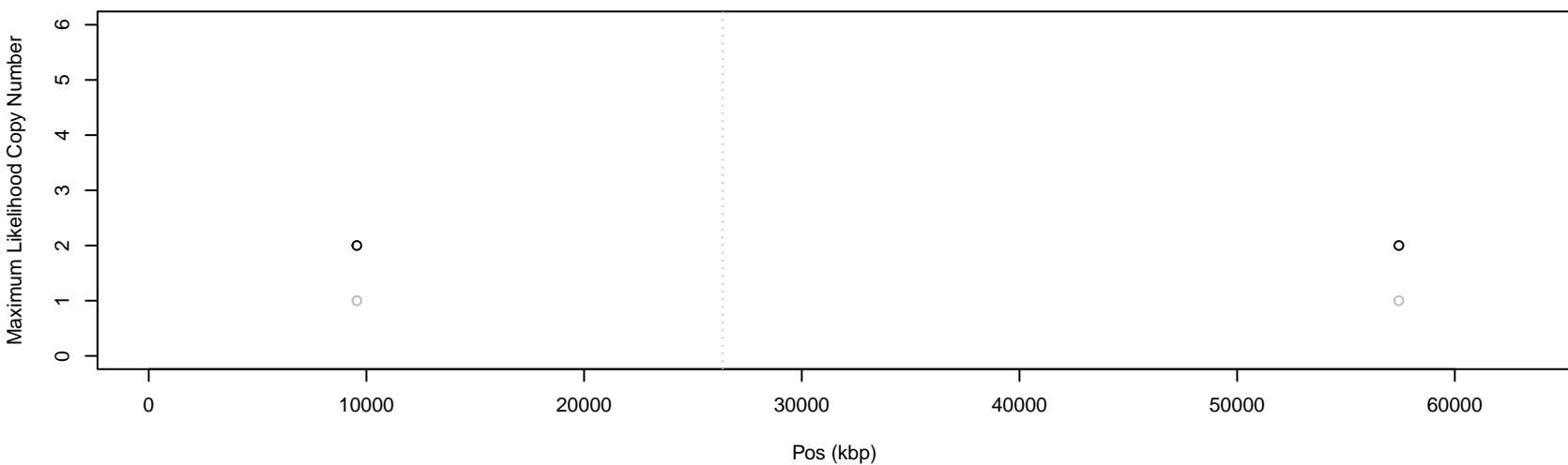
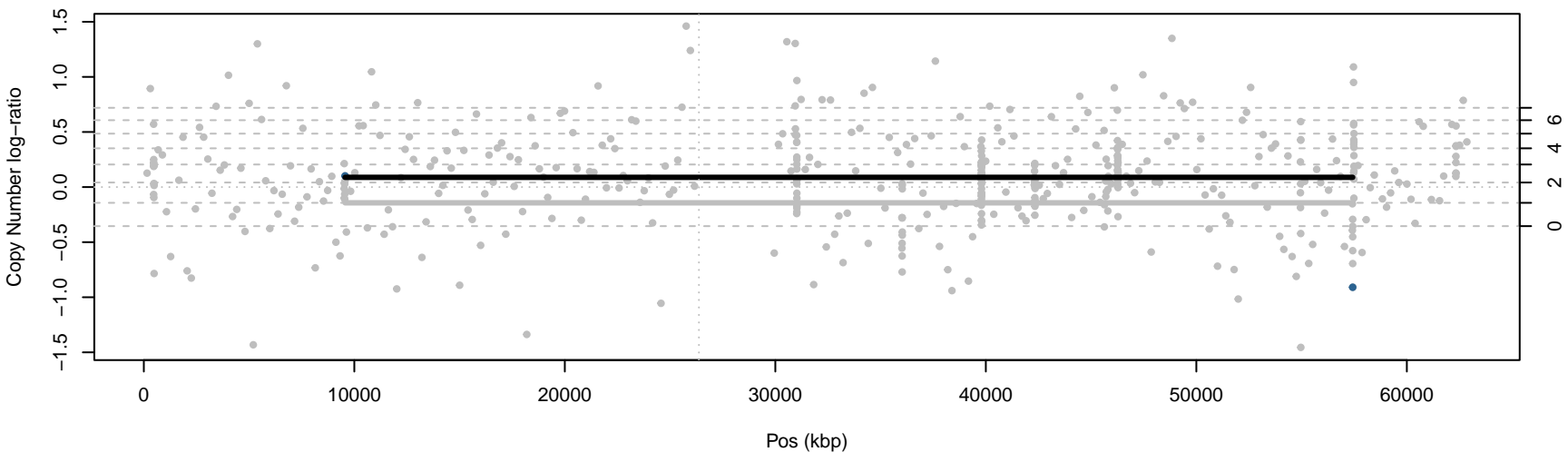
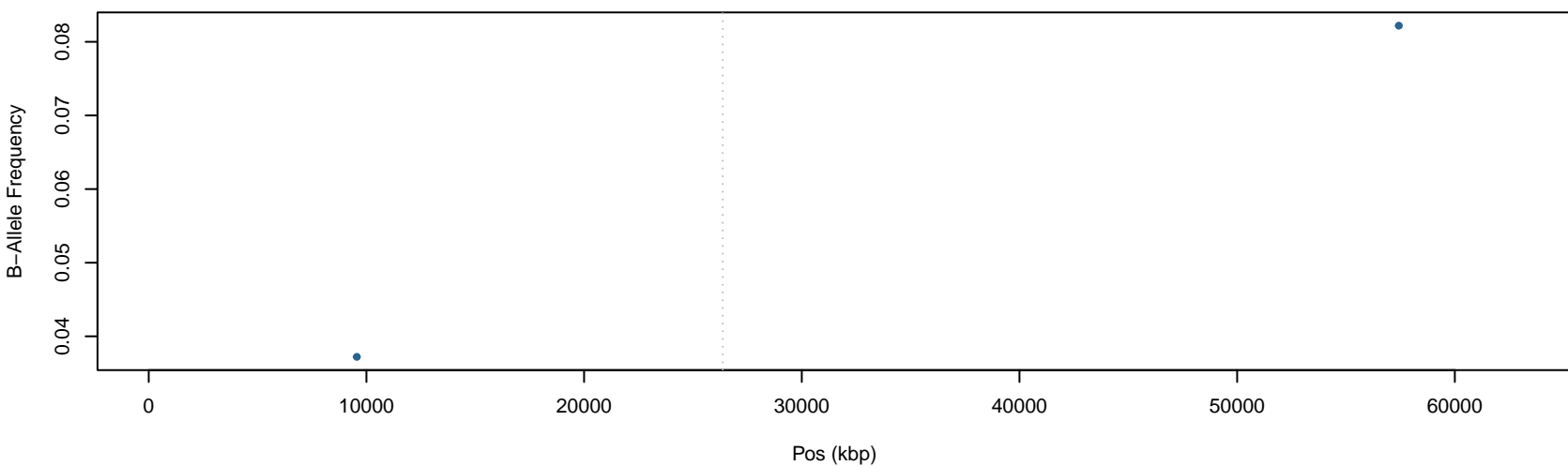


Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 19

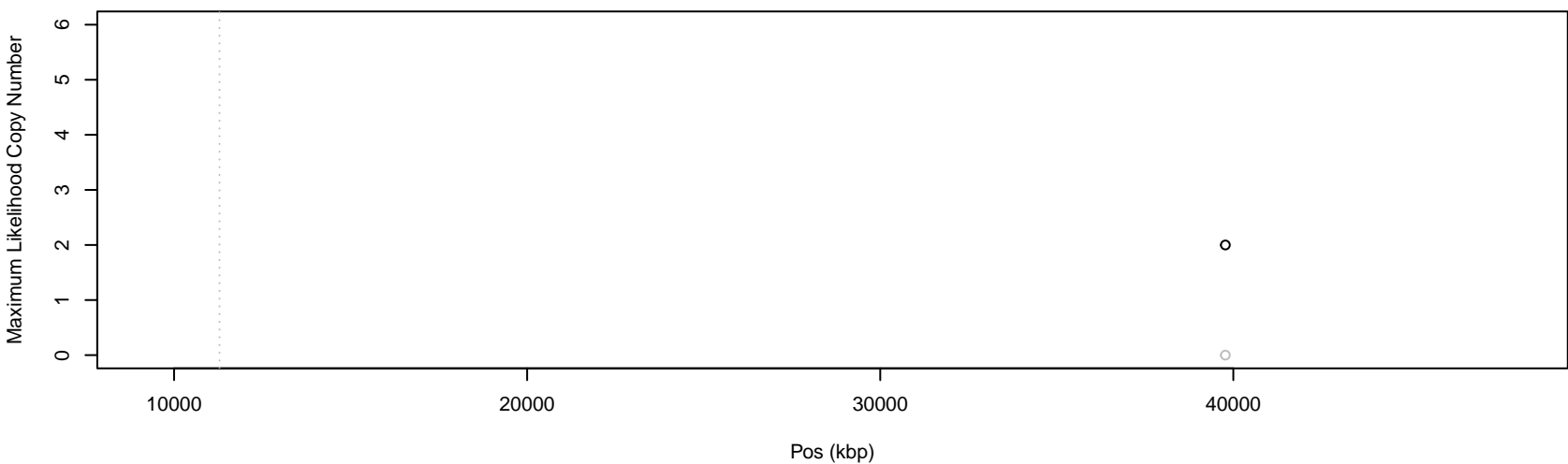
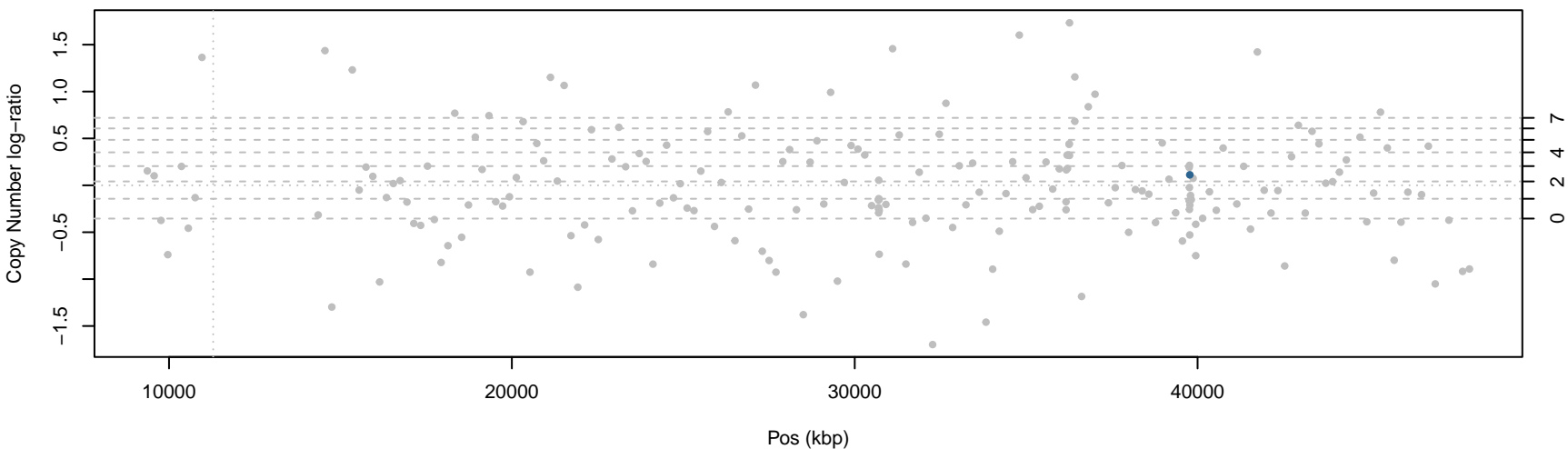
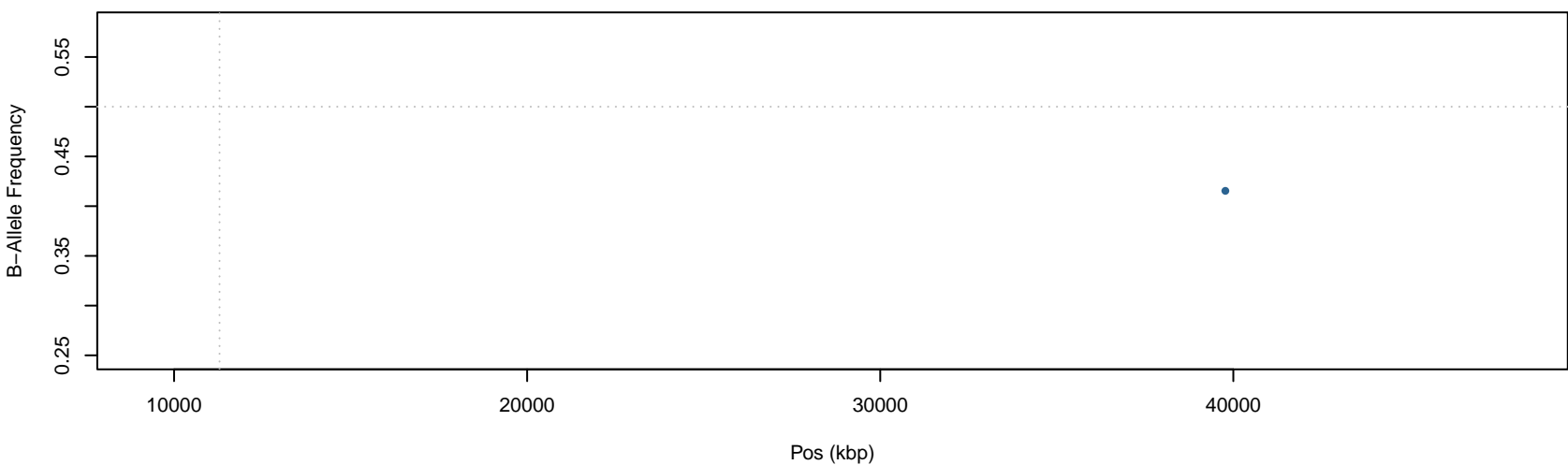




Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 20



Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401;324 Chromosome: 21



Purity: 0.24 Tumor ploidy: 1.765 SNV log-likelihood: 34.68 GoF: 90% Mean coverage: 401,324 Chromosome: 22

