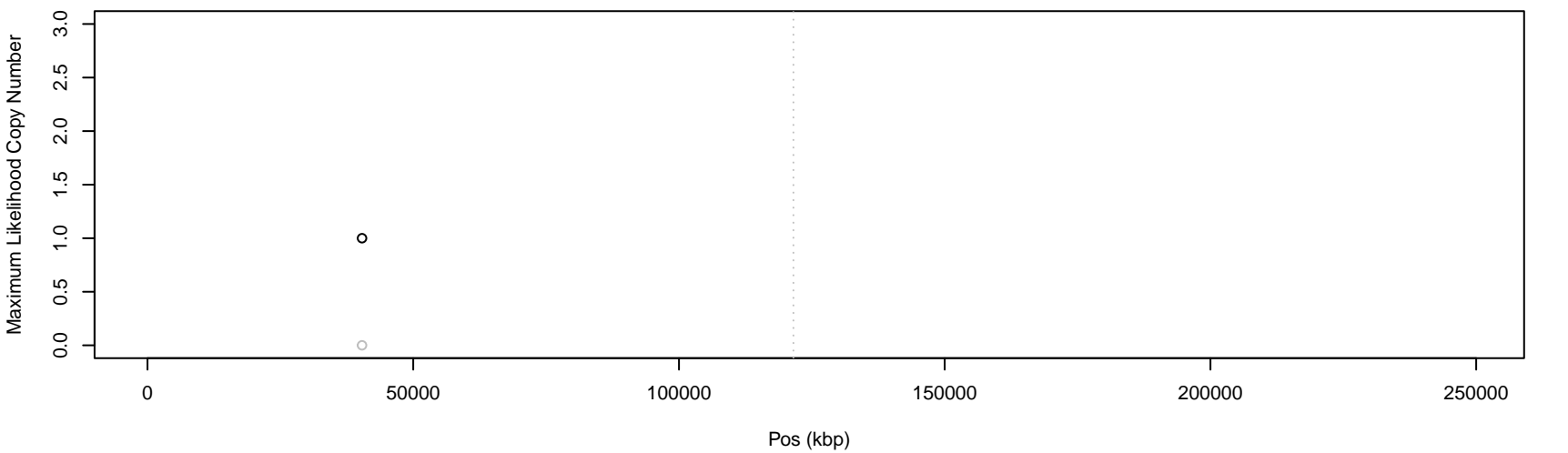
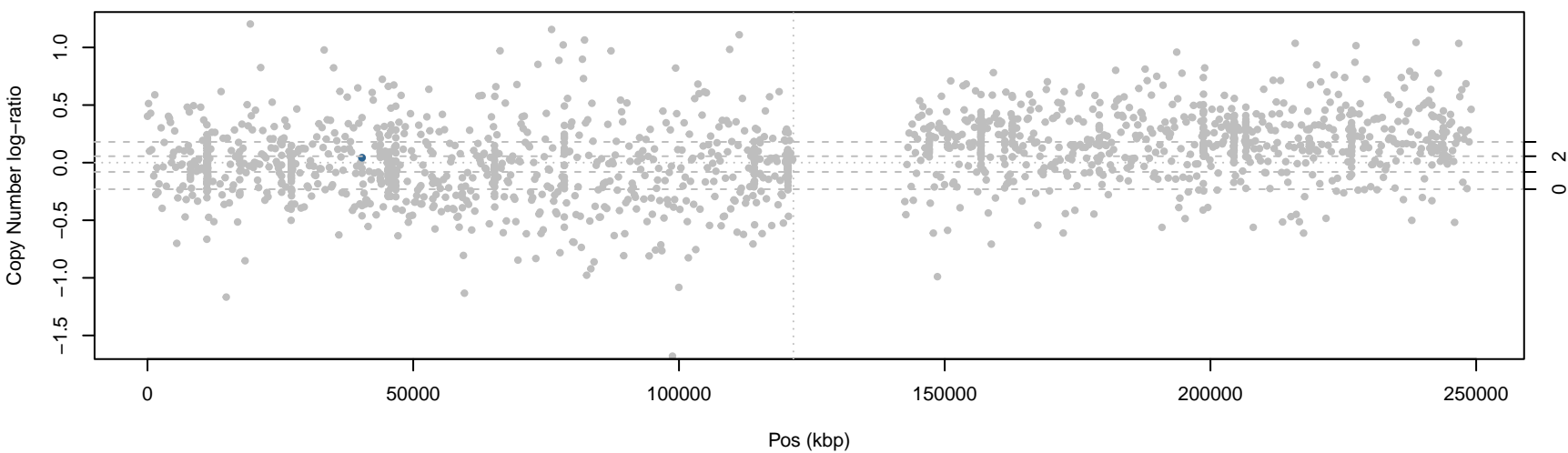
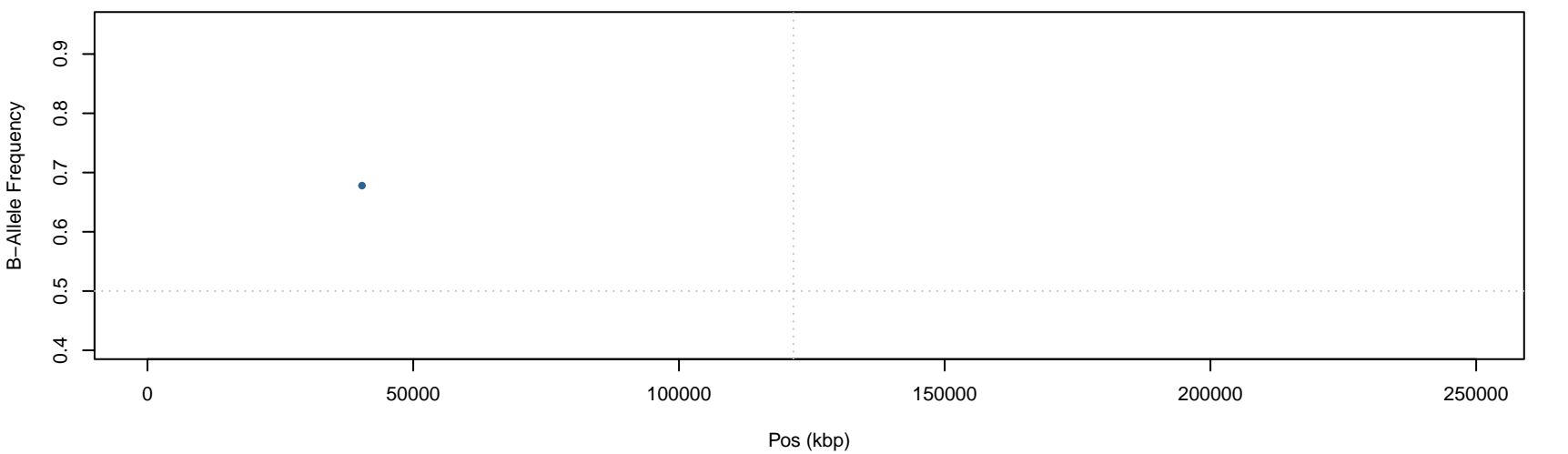
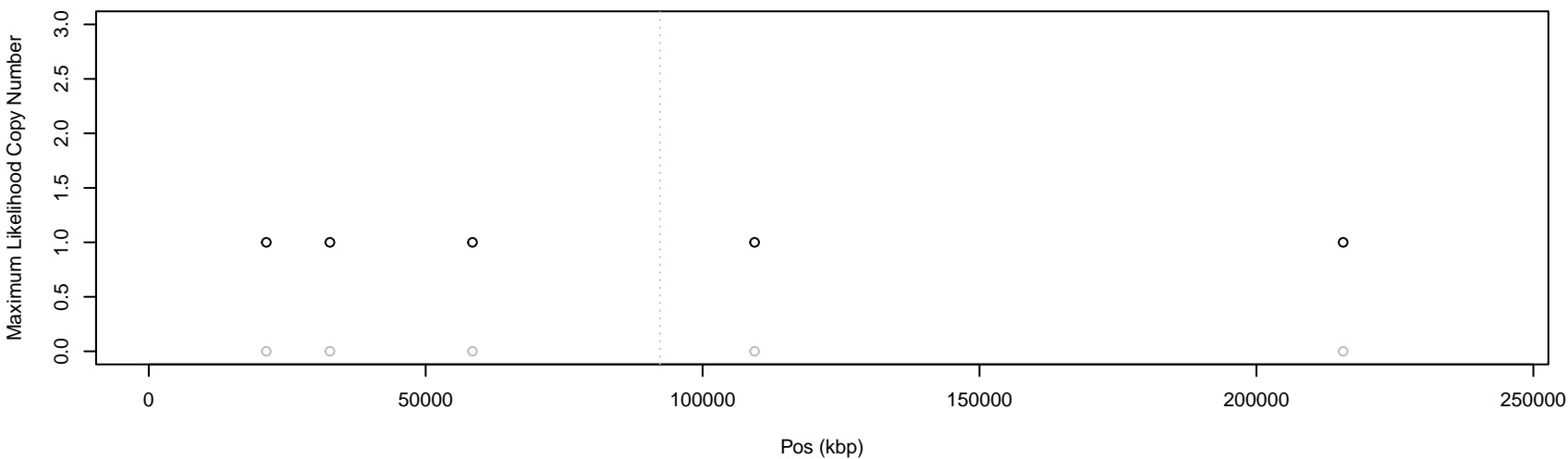
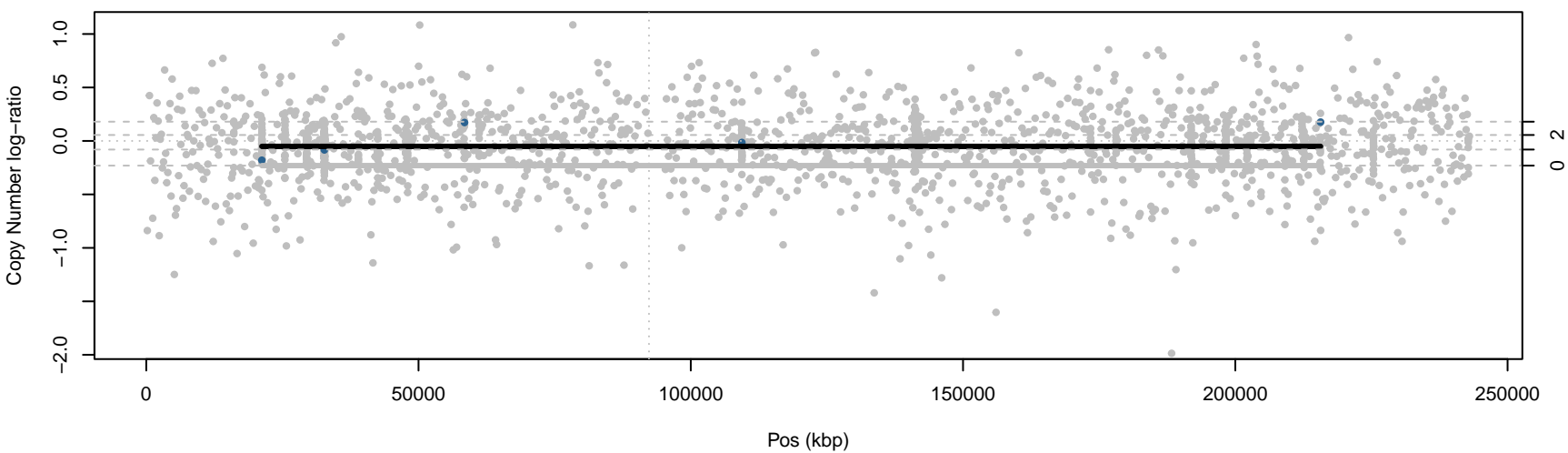
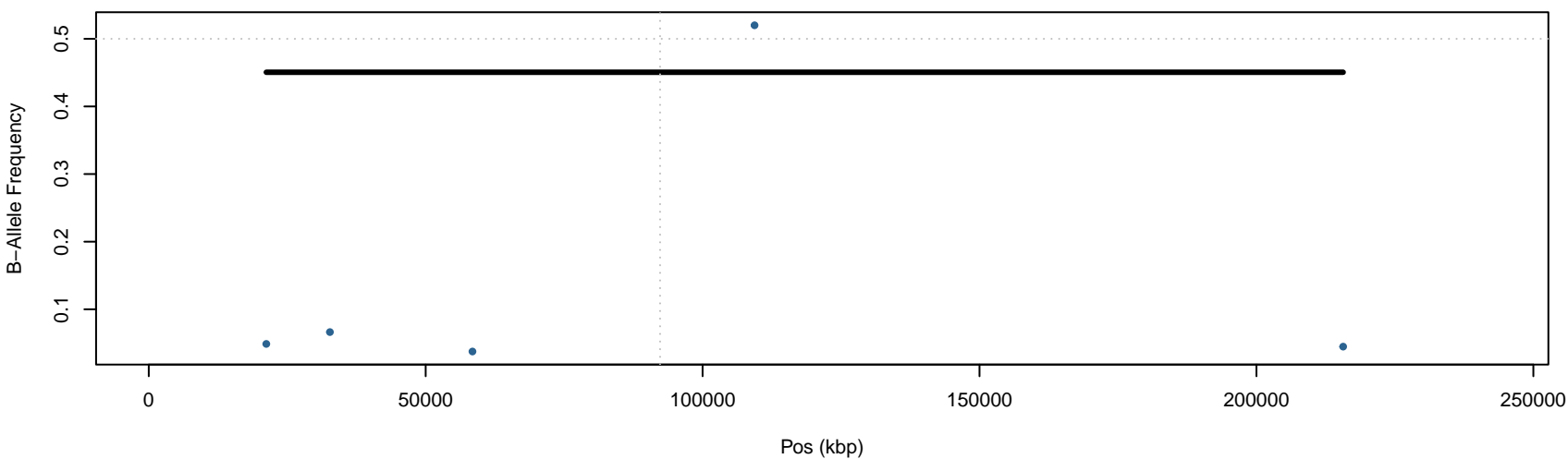


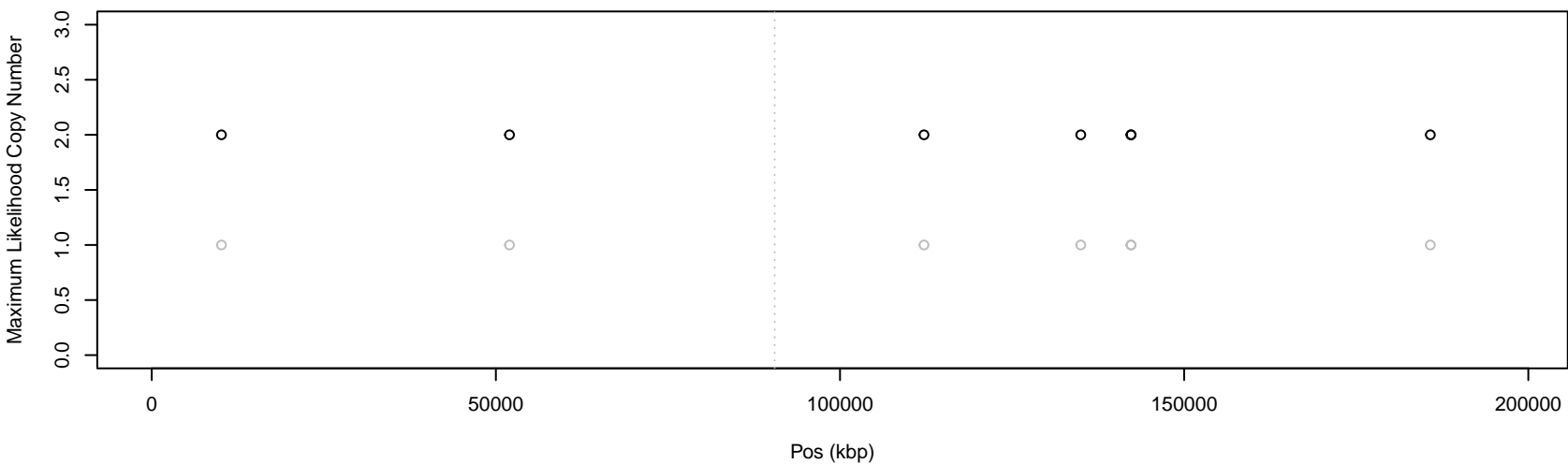
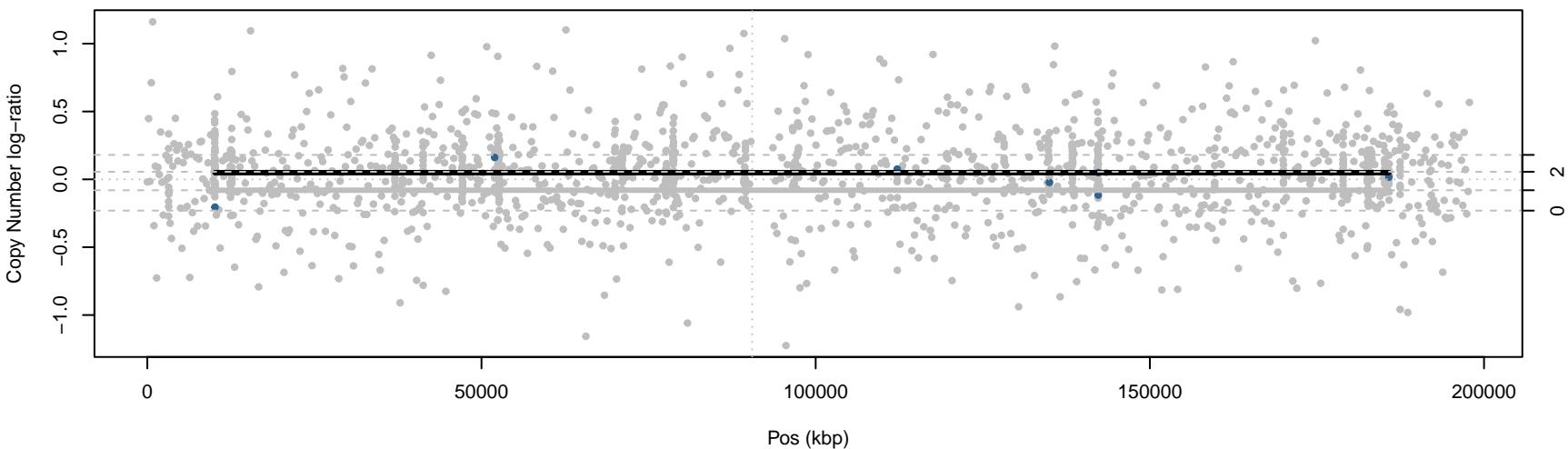
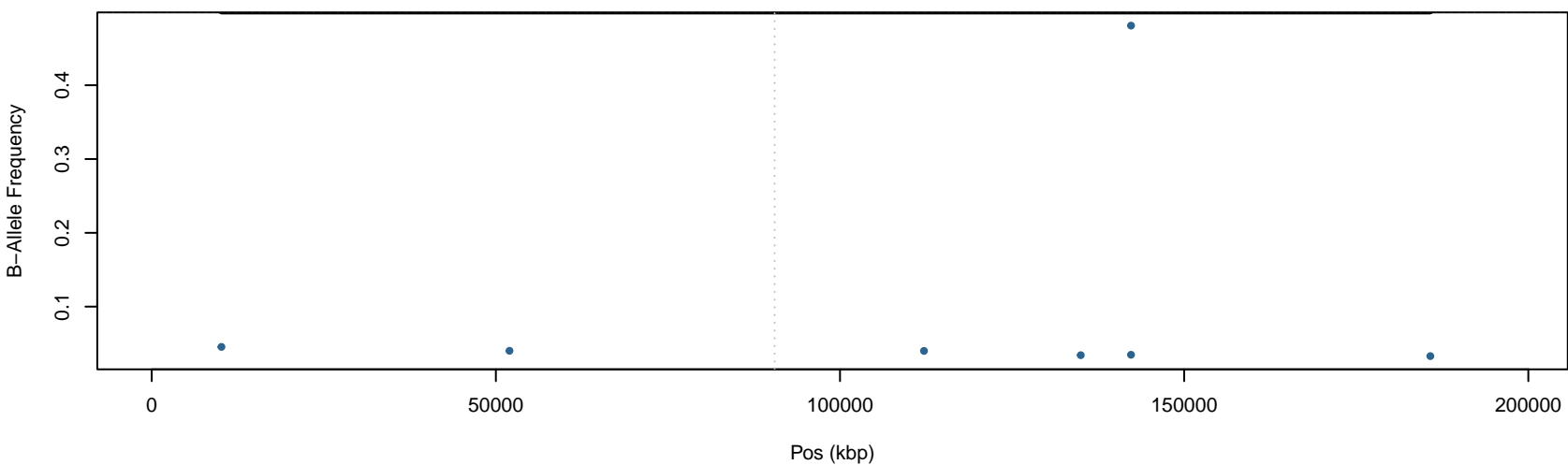
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 1



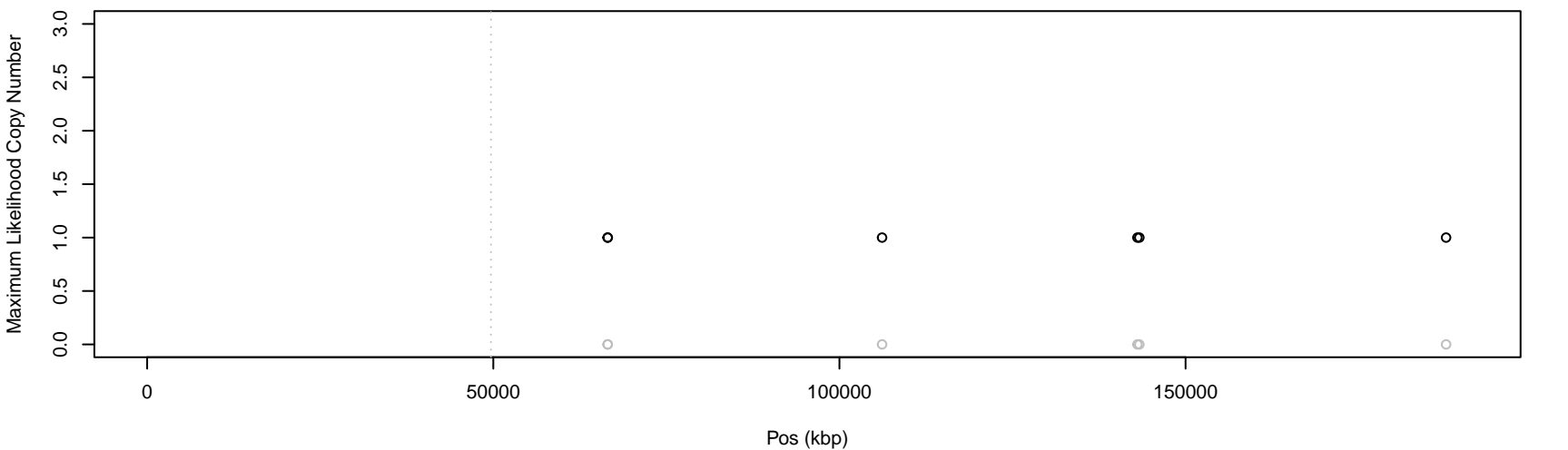
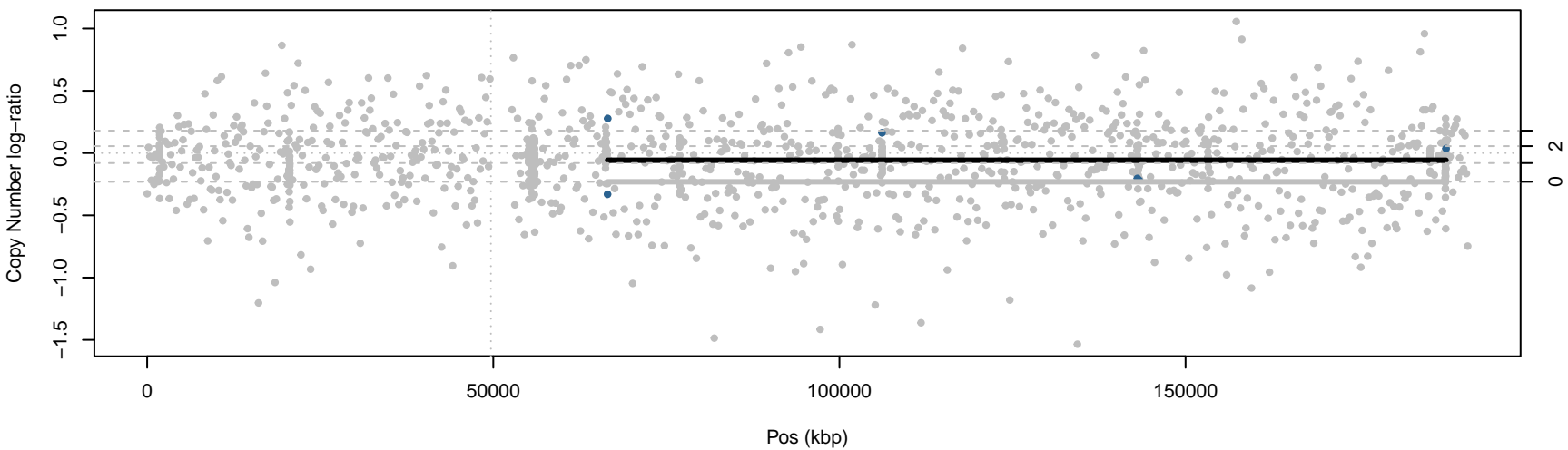
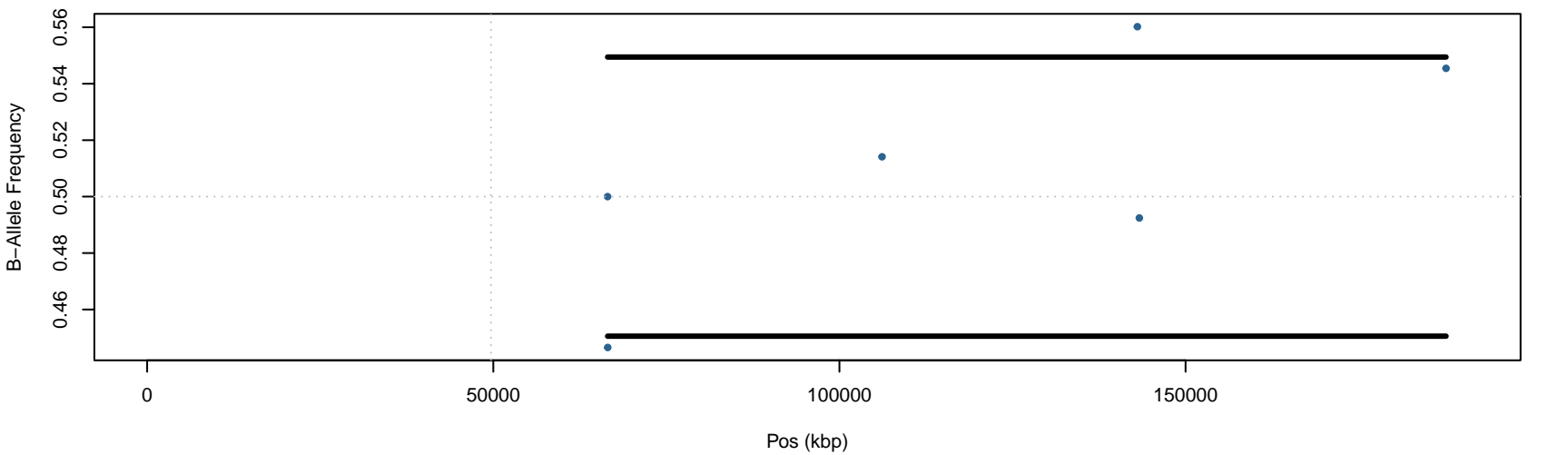
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 2



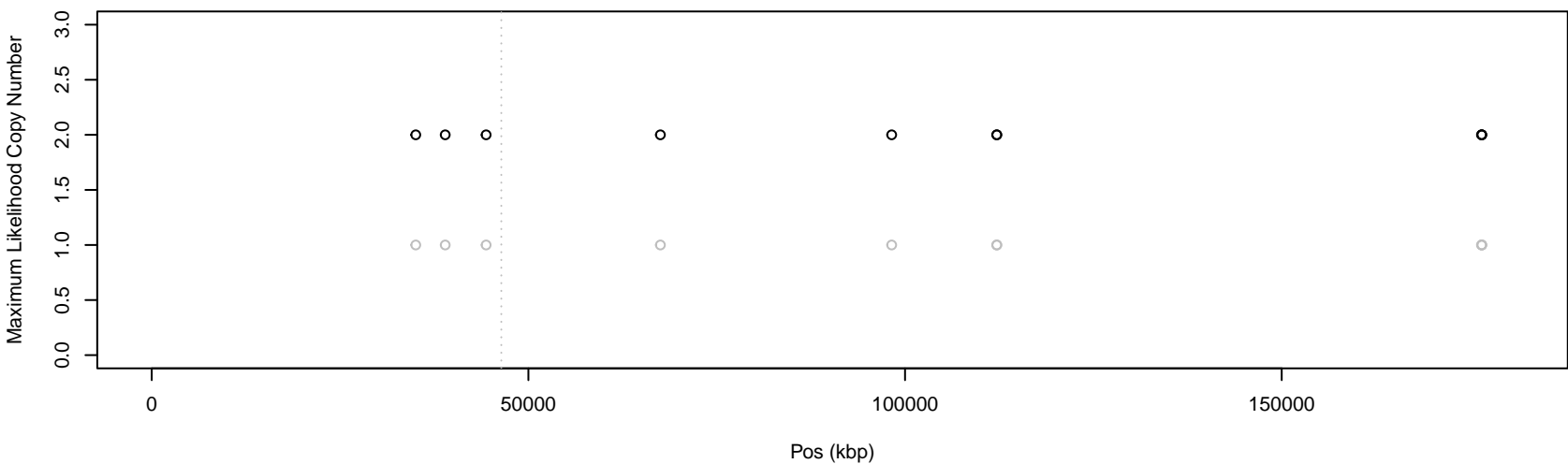
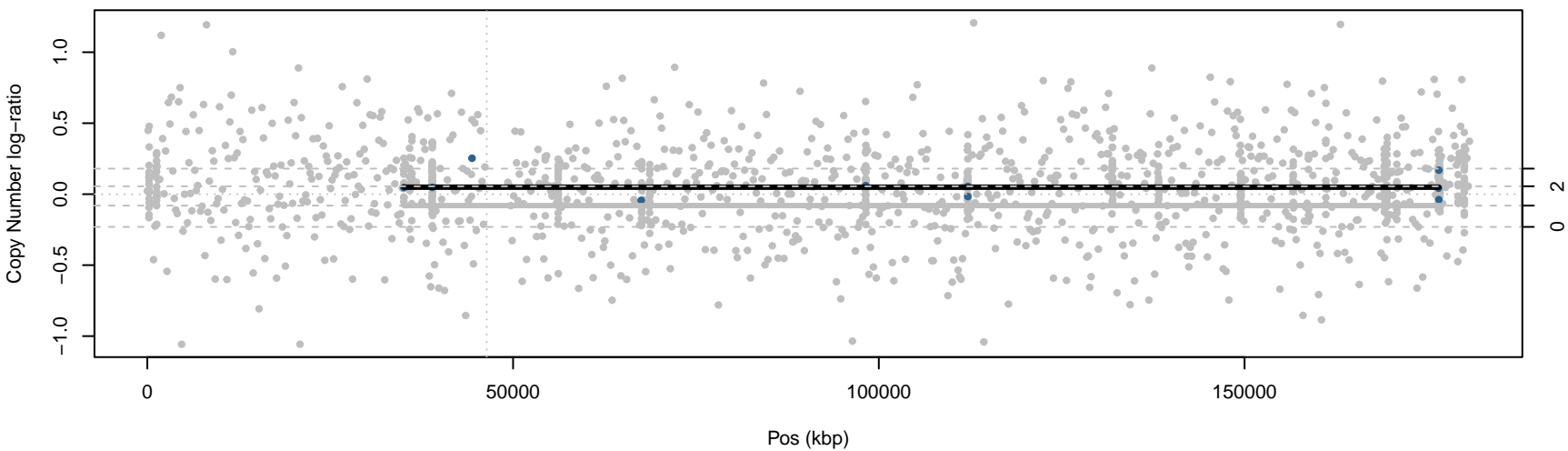
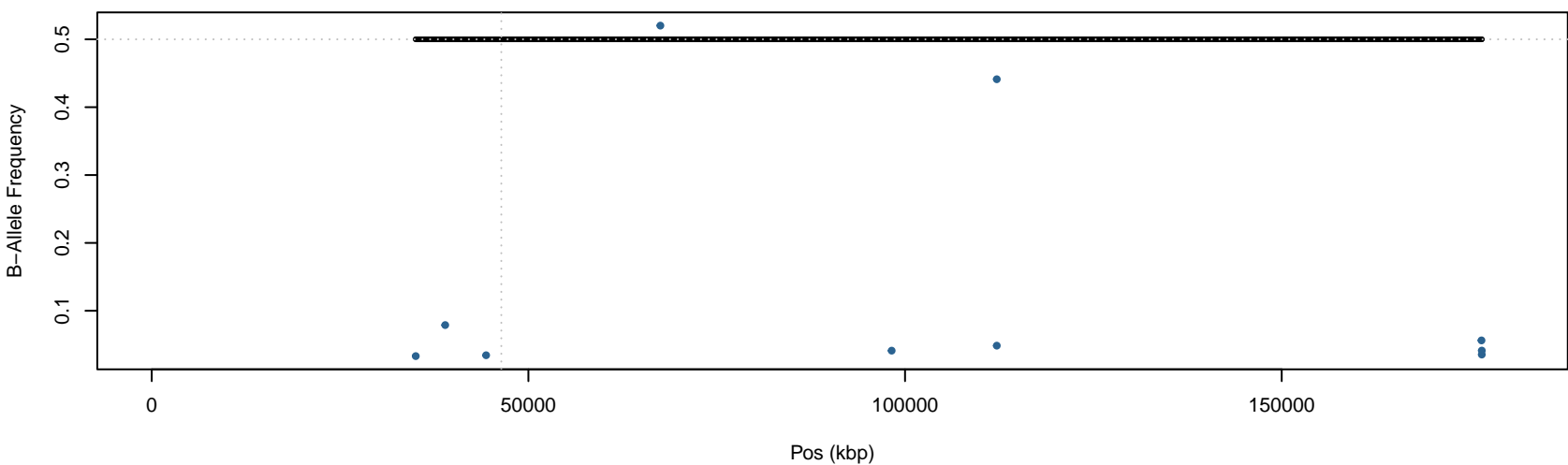
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 3



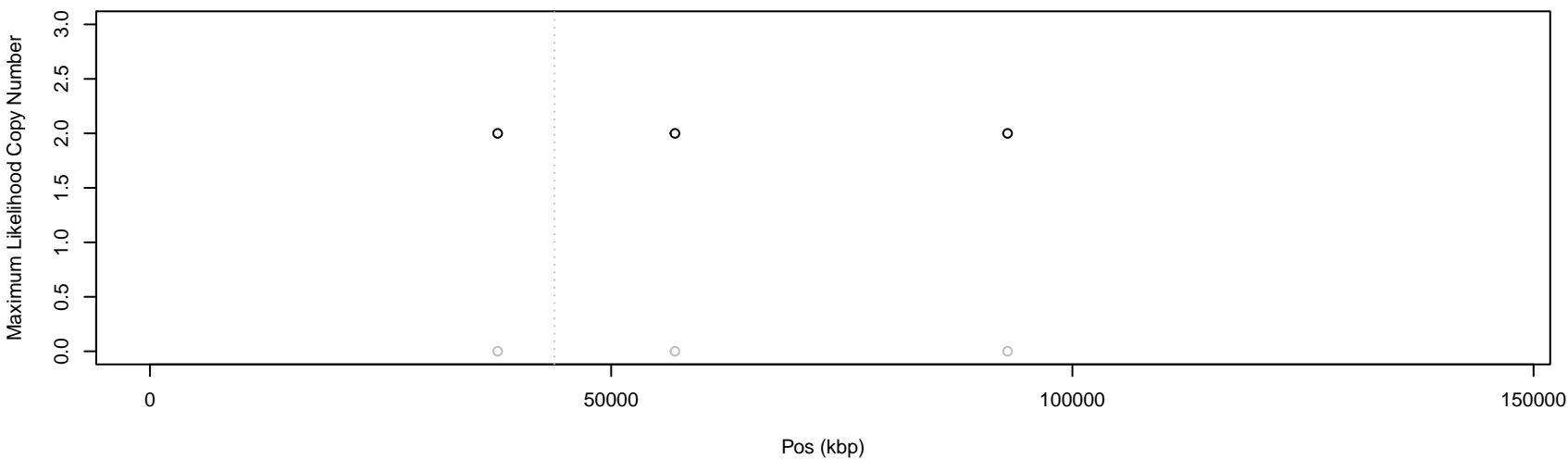
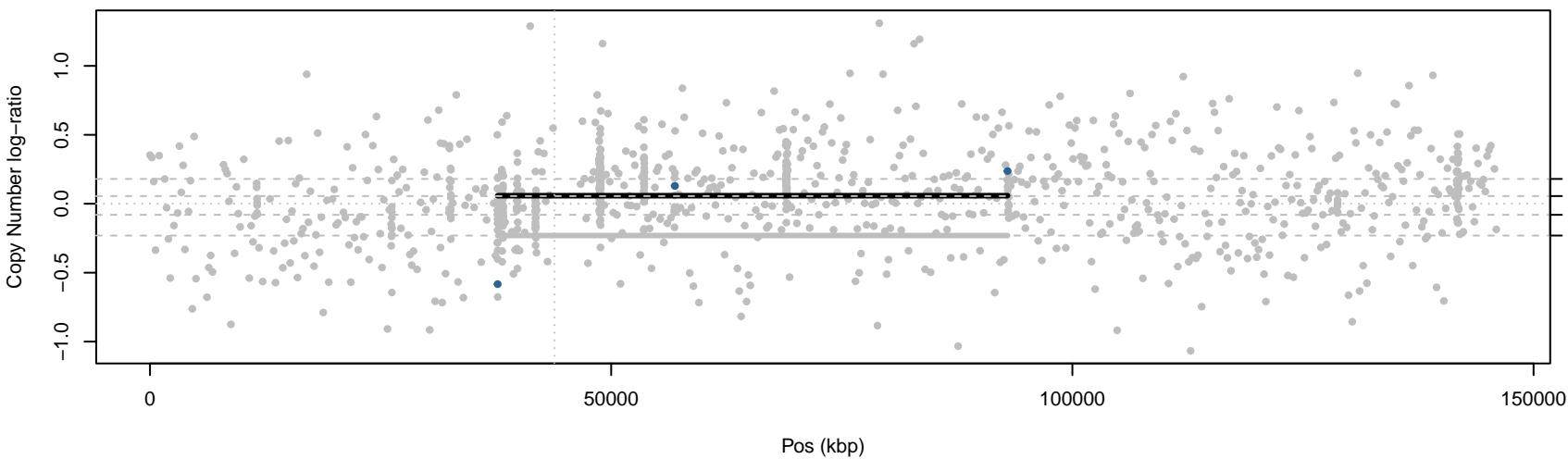
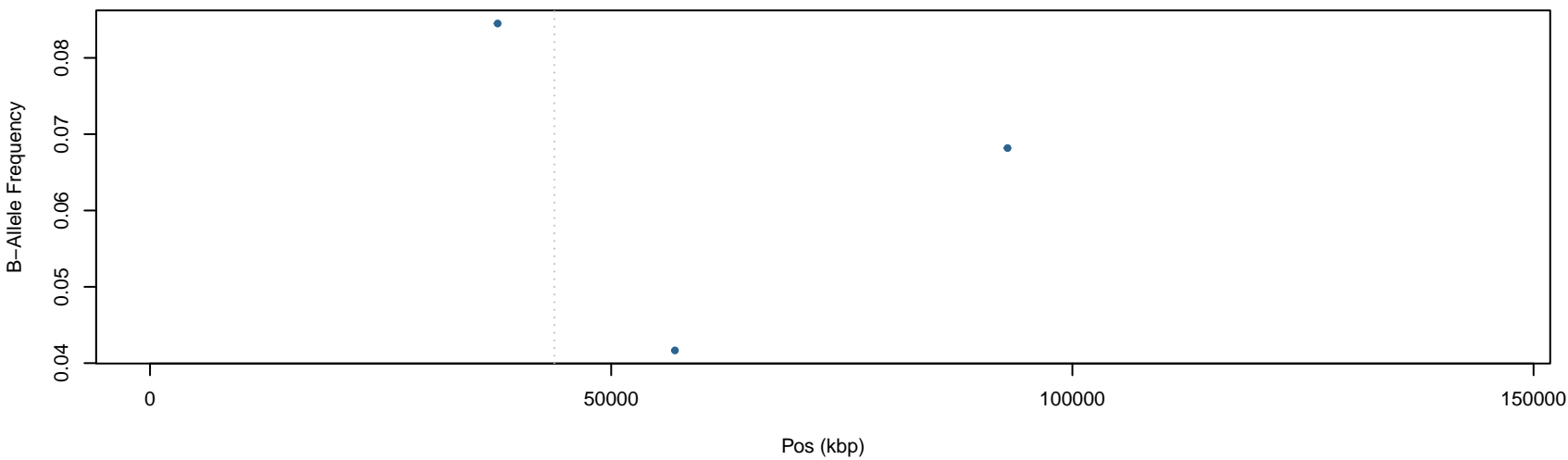
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 4



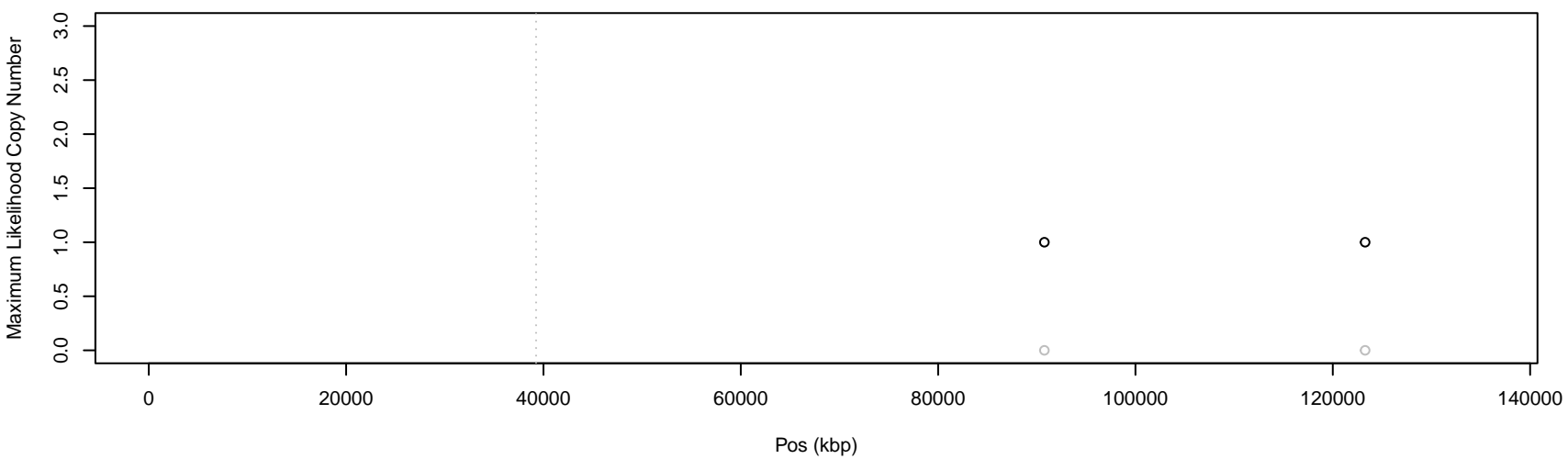
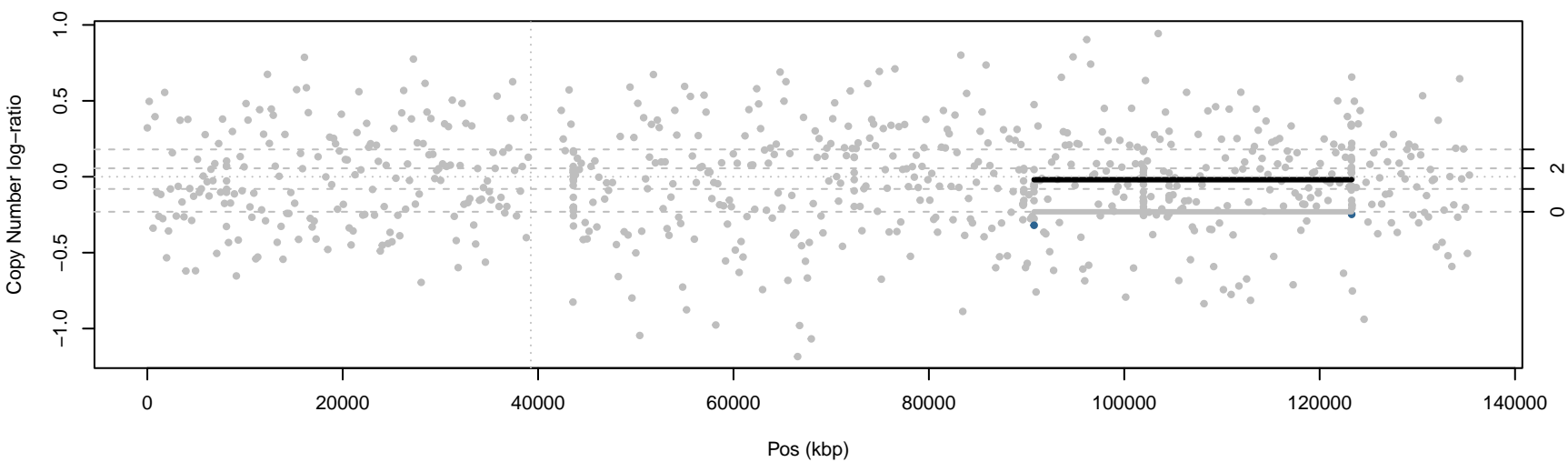
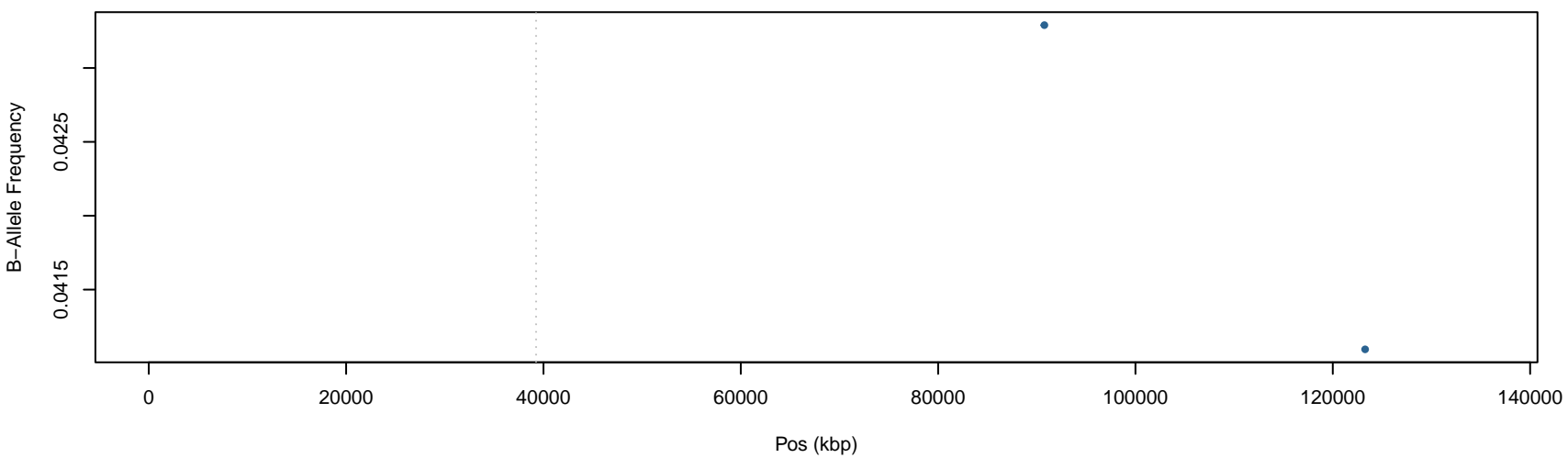
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 5



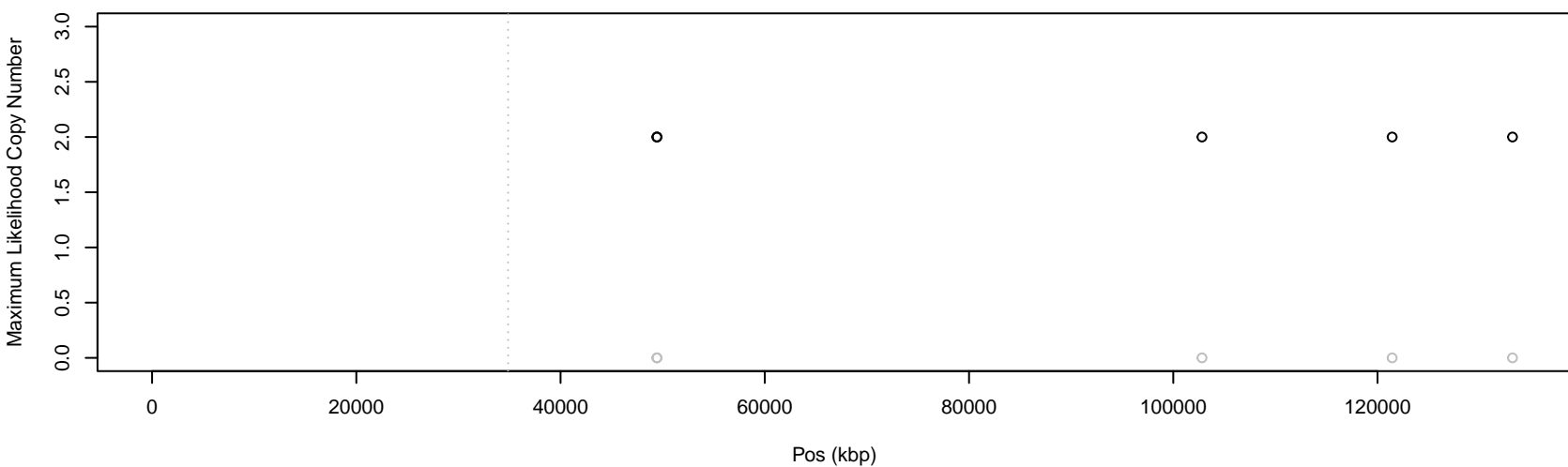
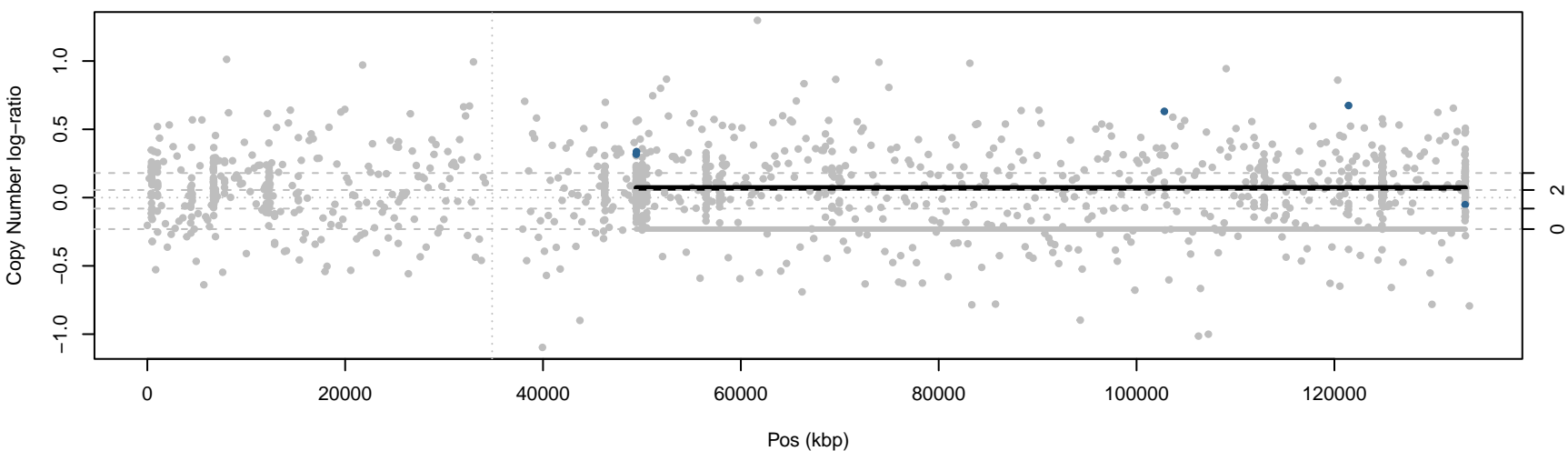
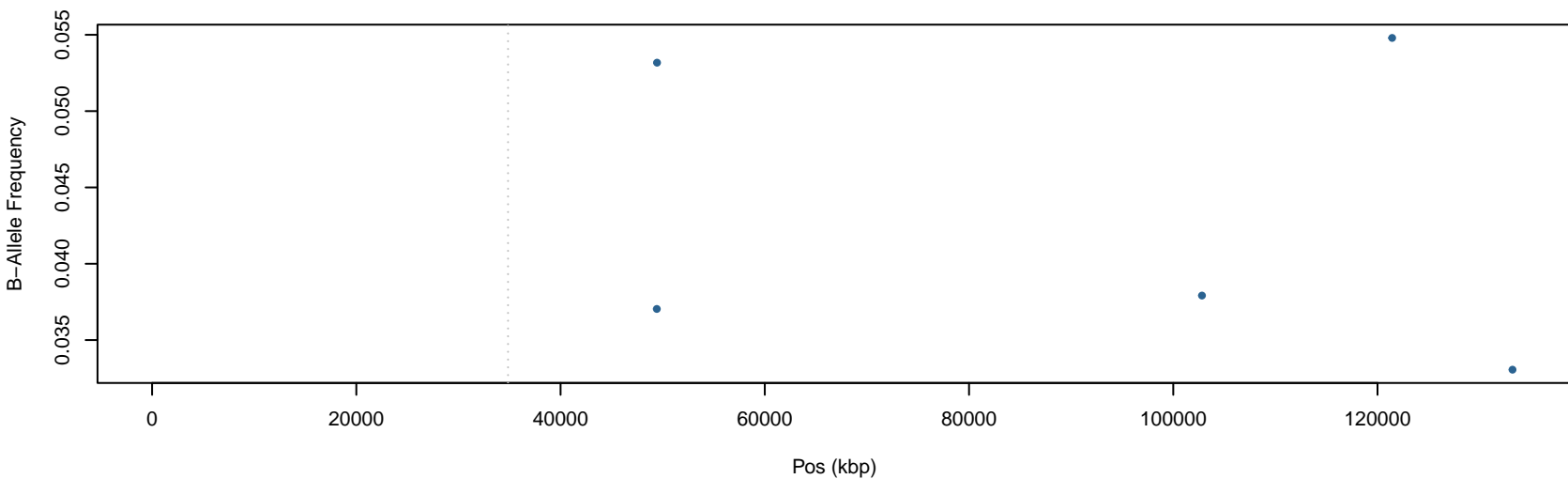
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 8



Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 10

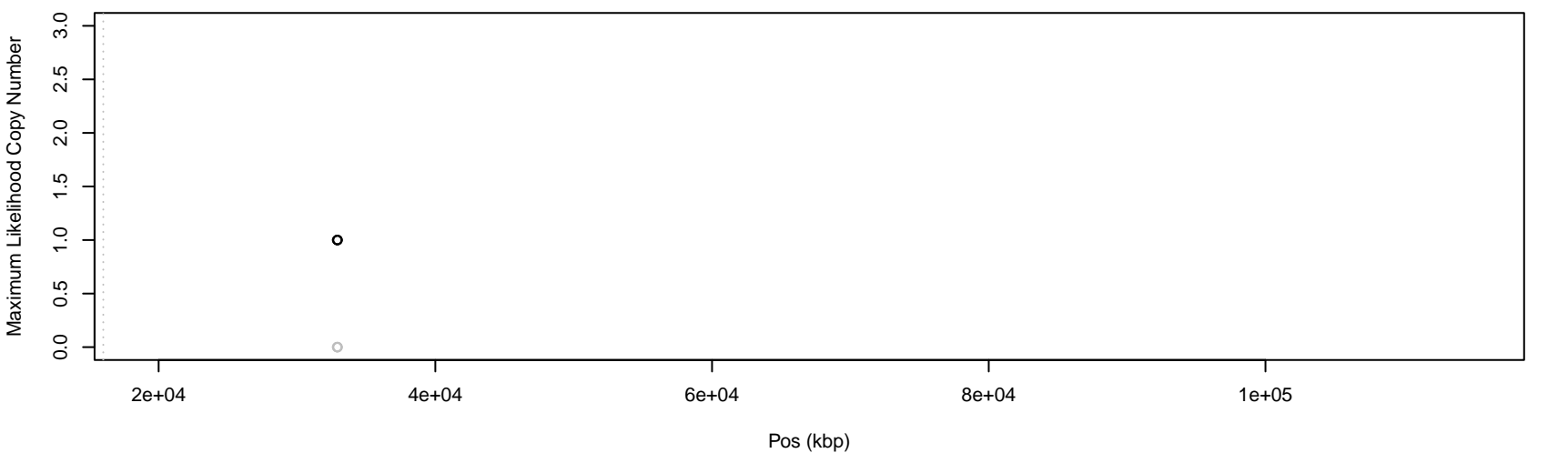
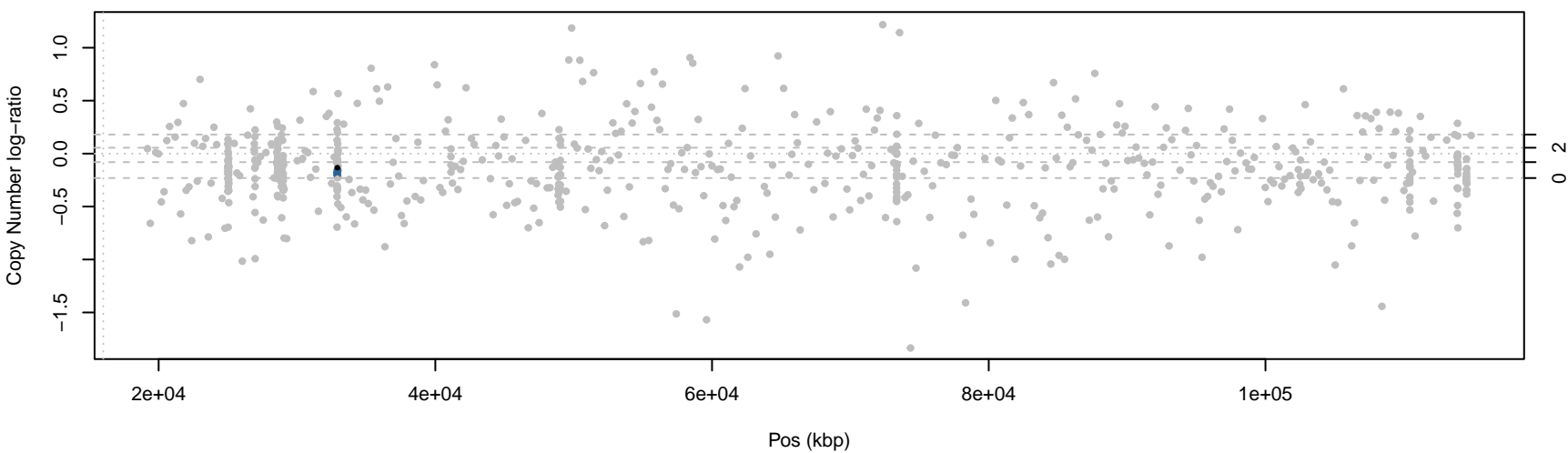
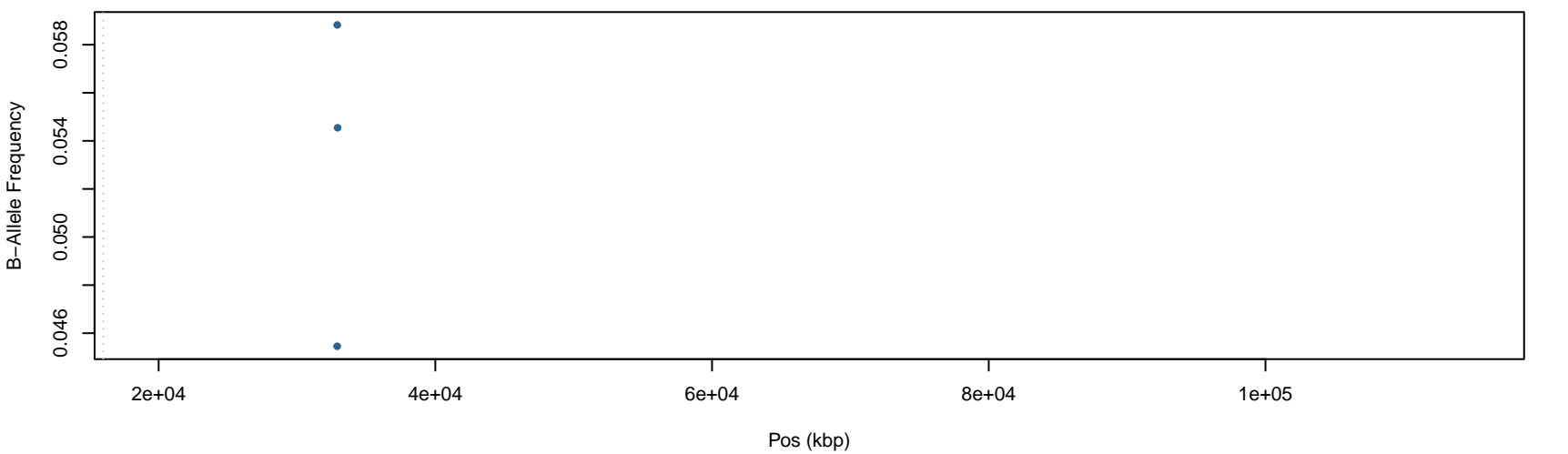


Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 12

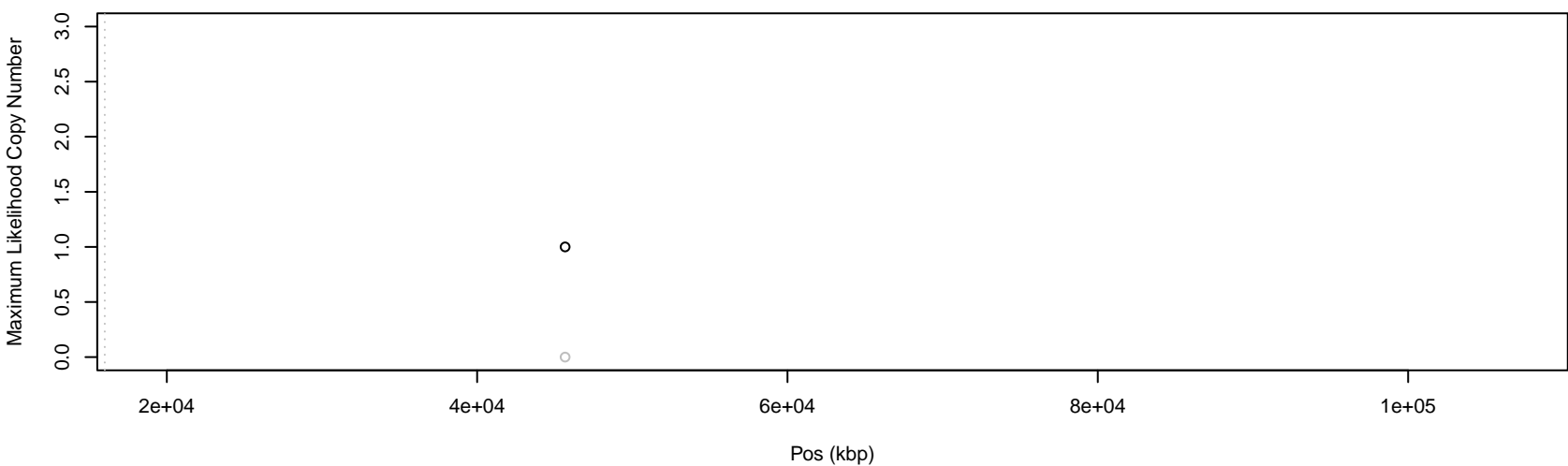
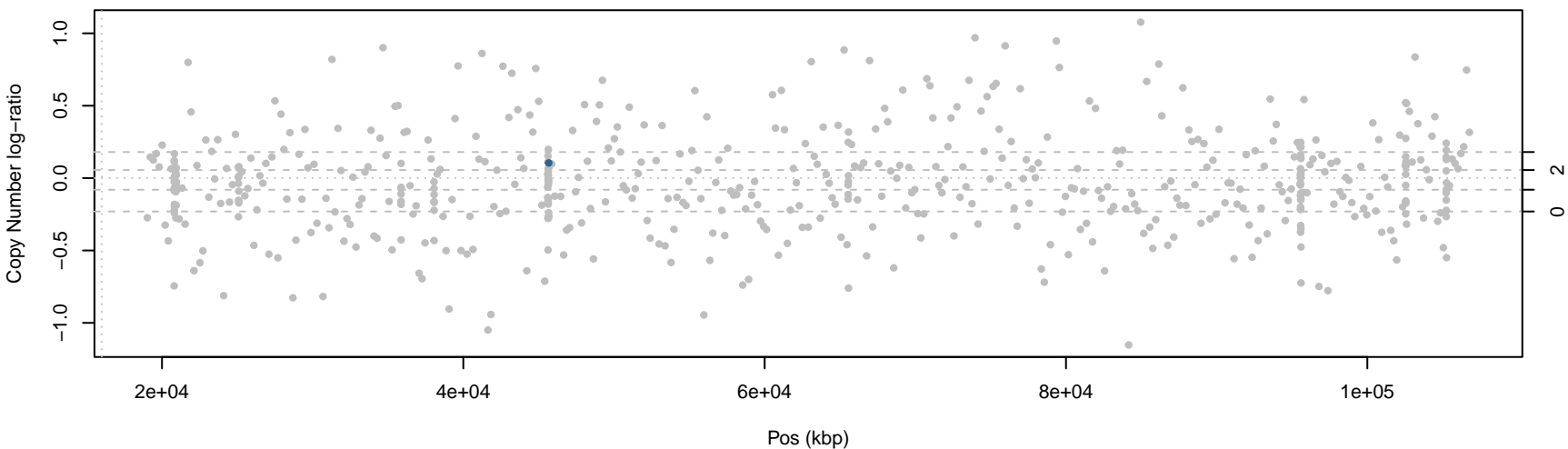
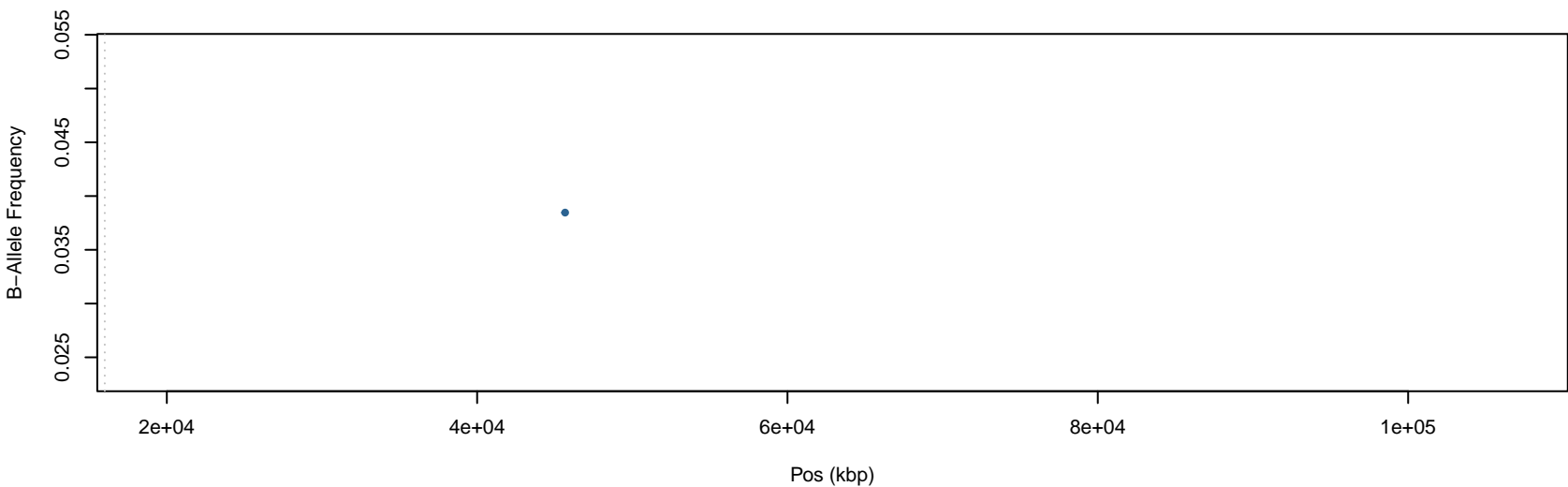




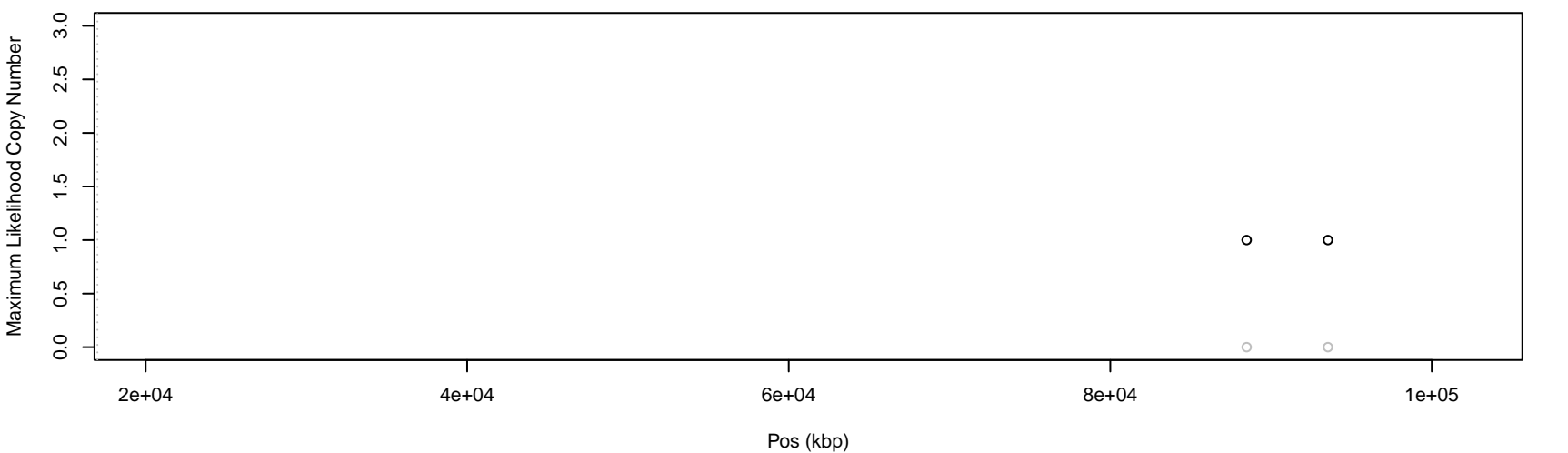
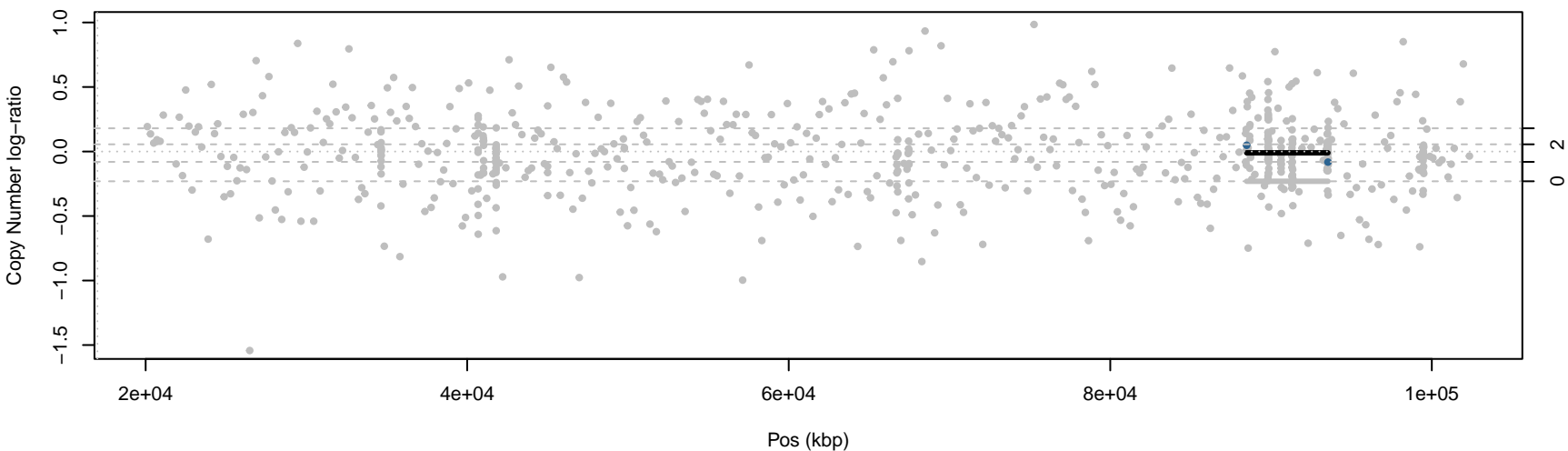
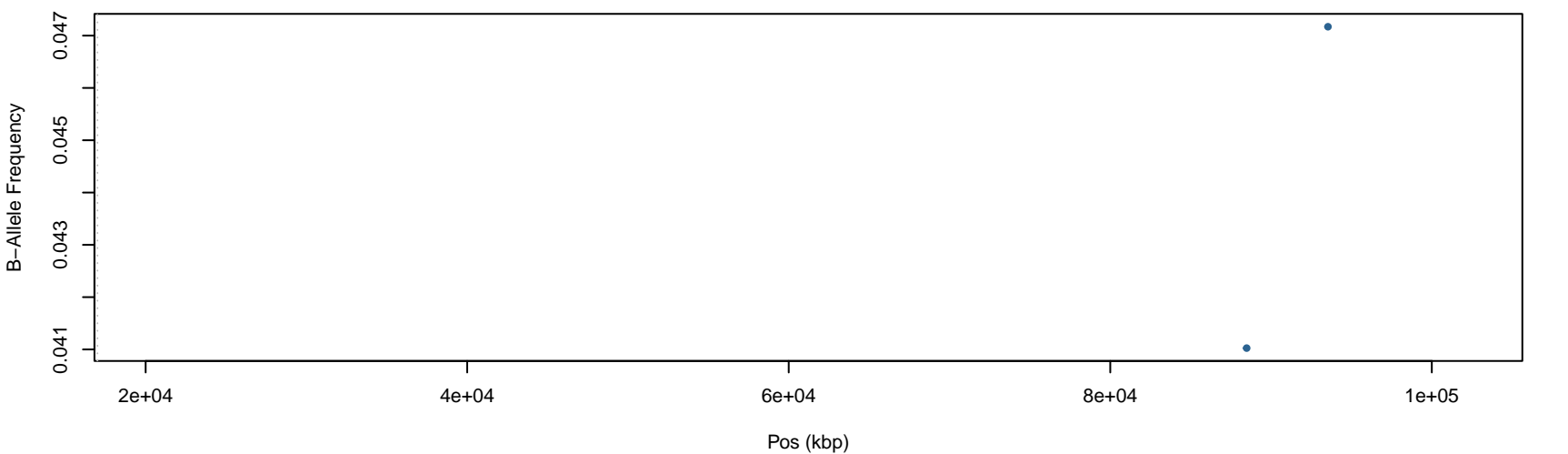
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 13



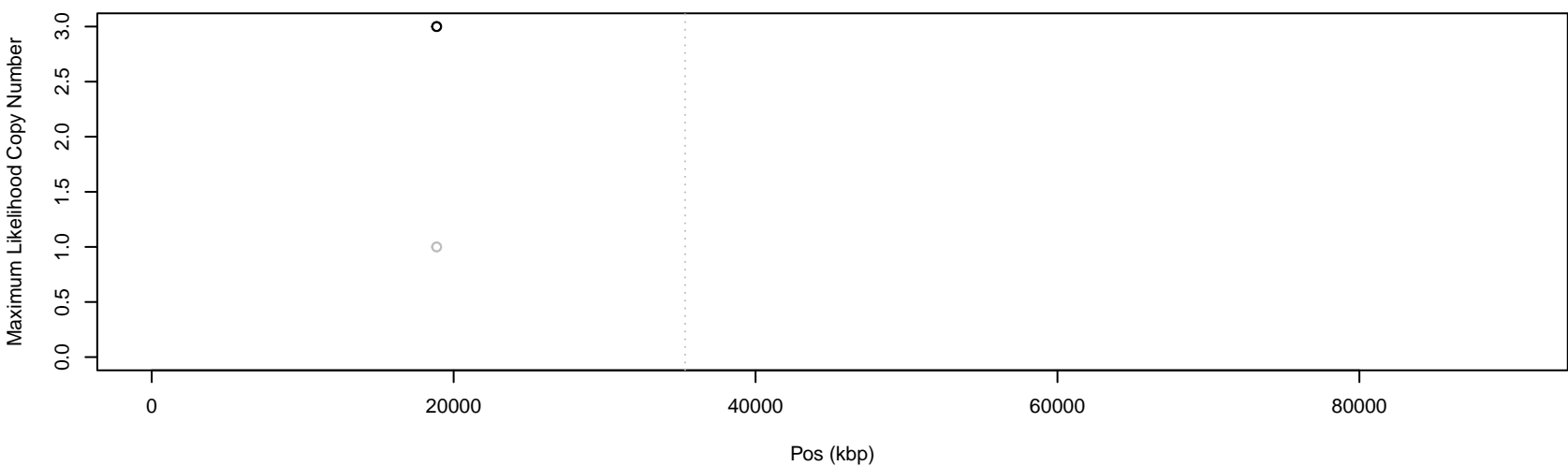
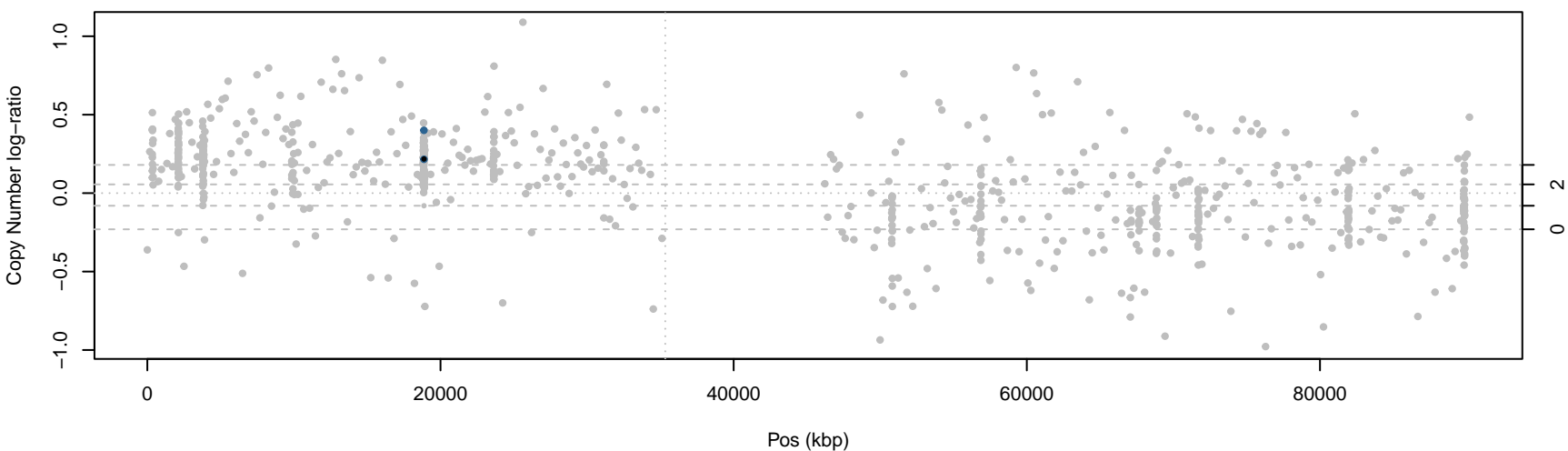
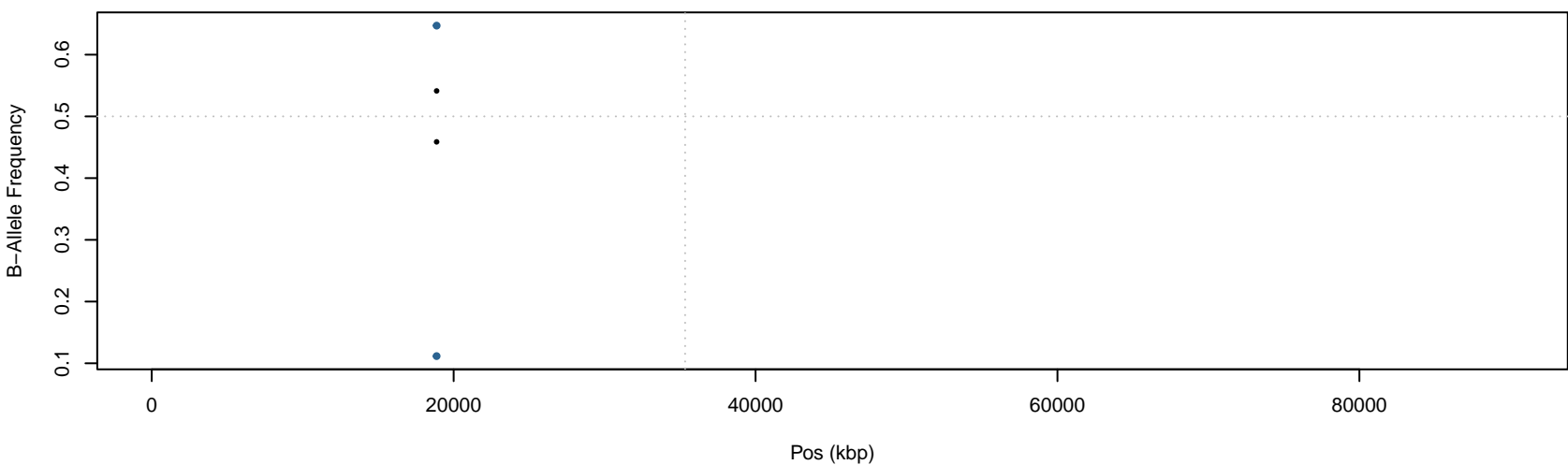
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 14



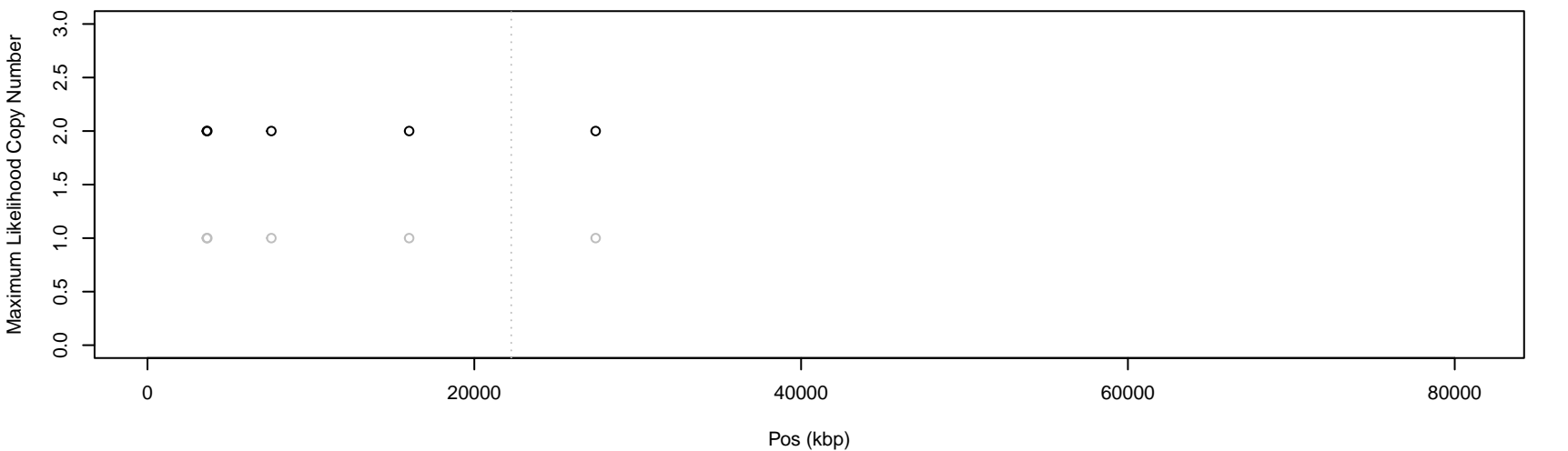
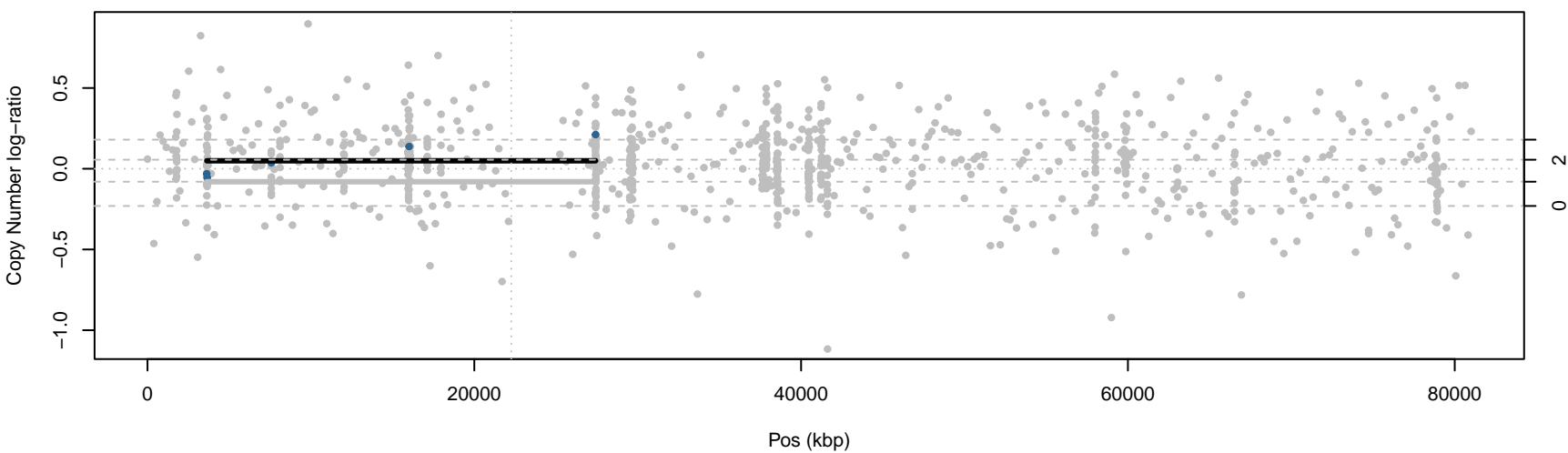
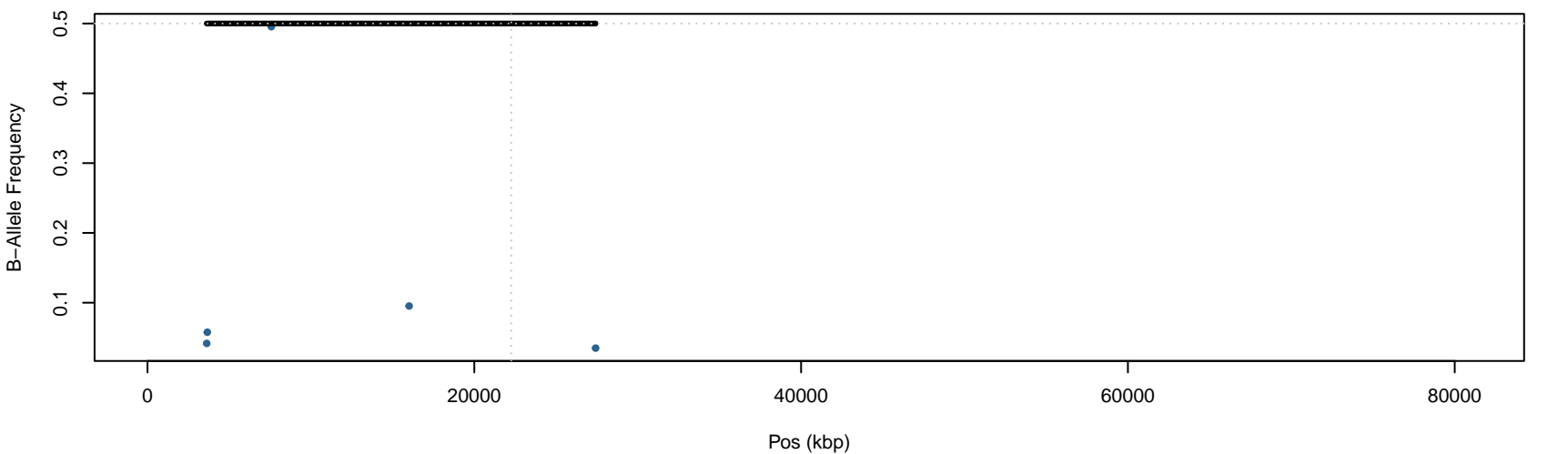
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 15



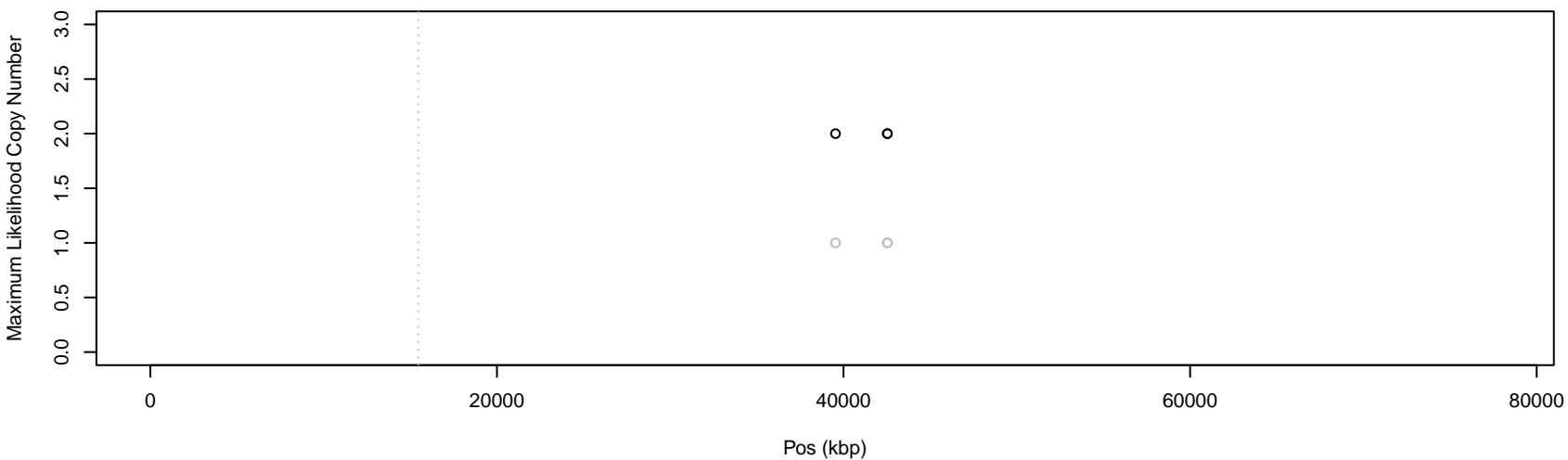
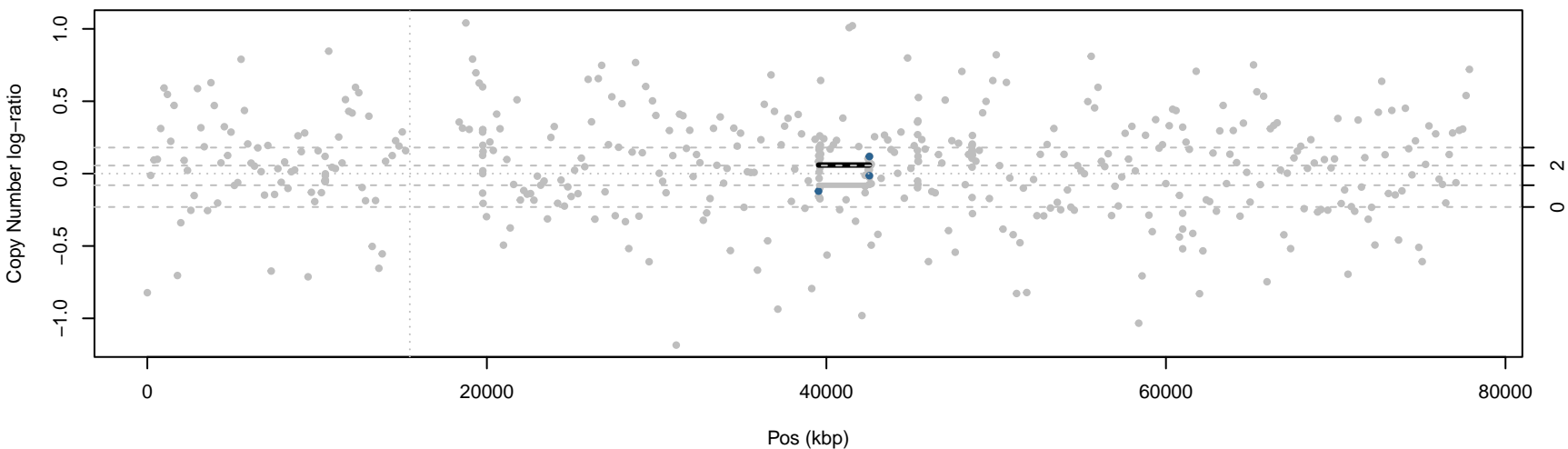
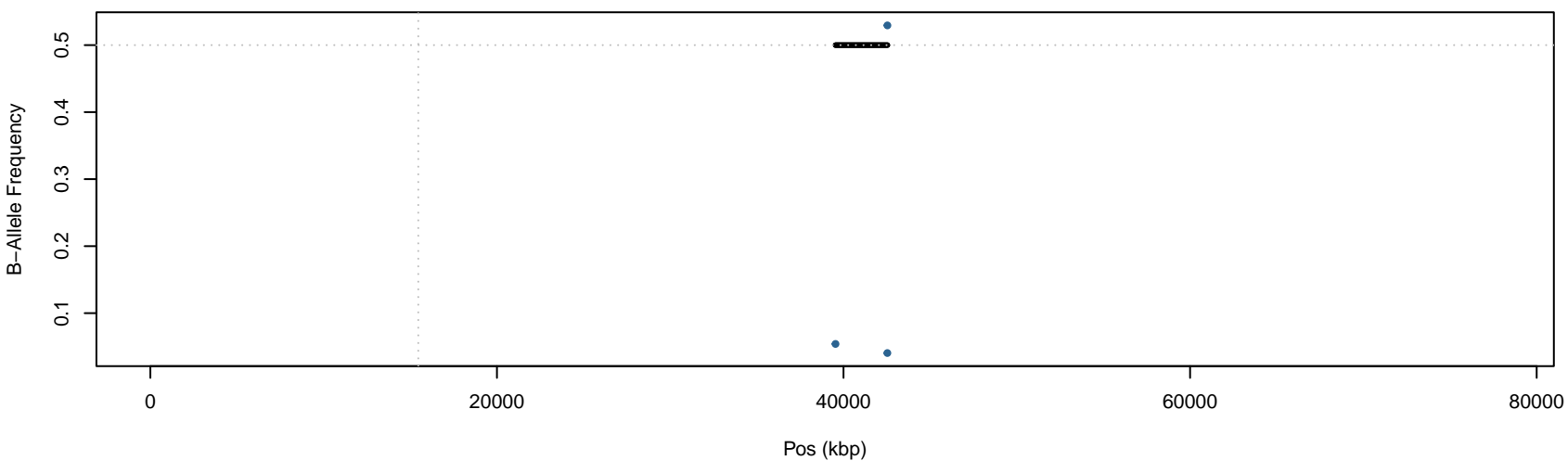
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 16



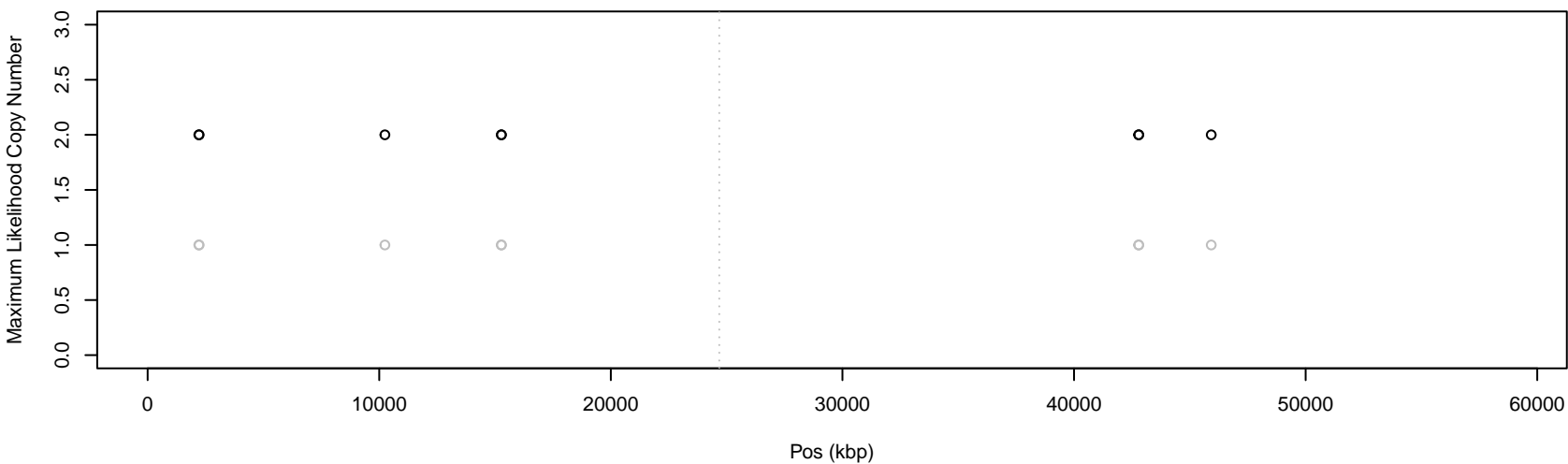
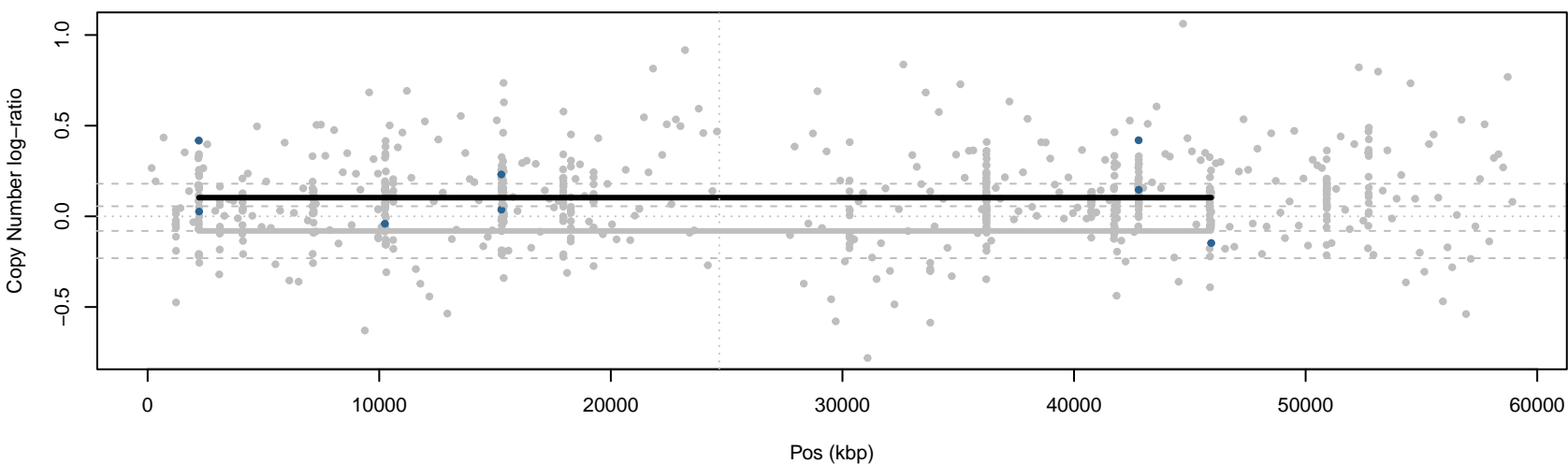
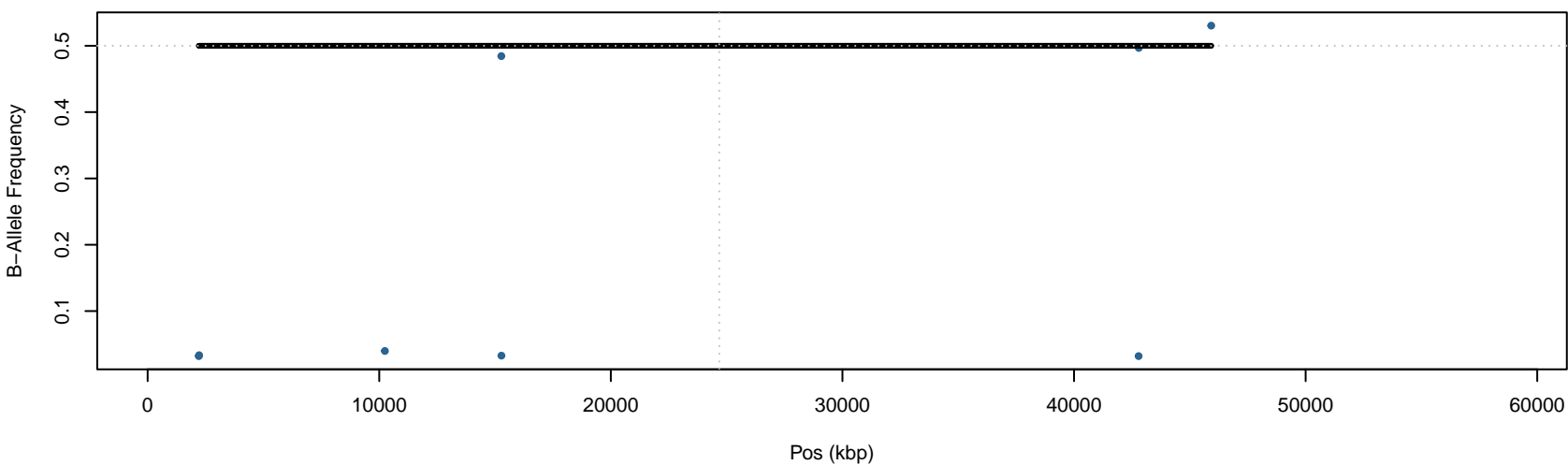
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 17



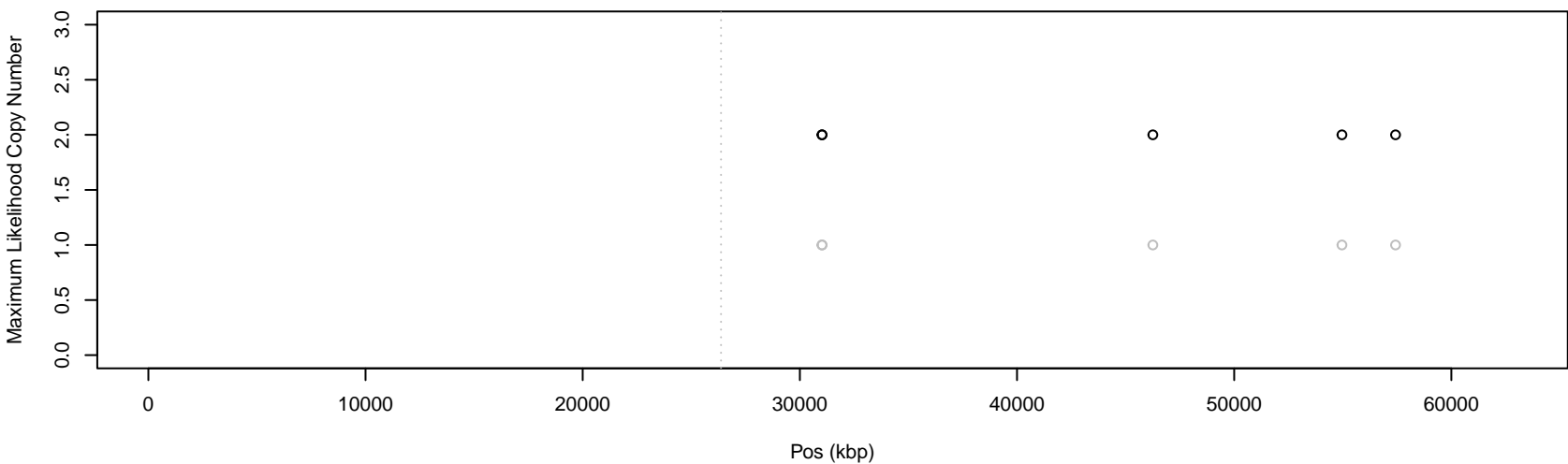
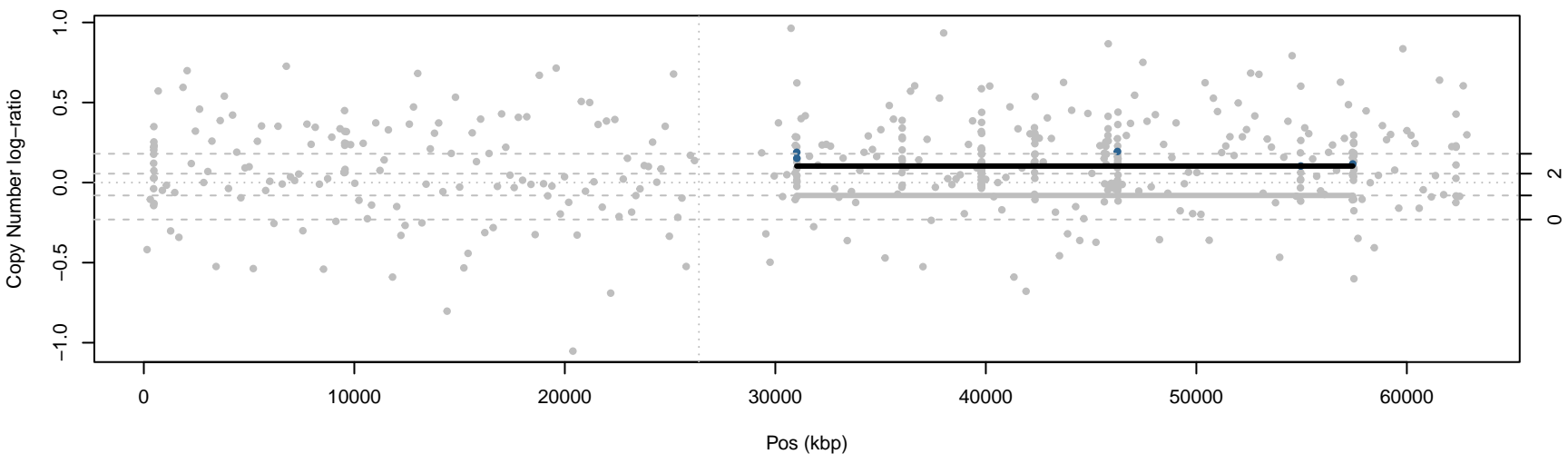
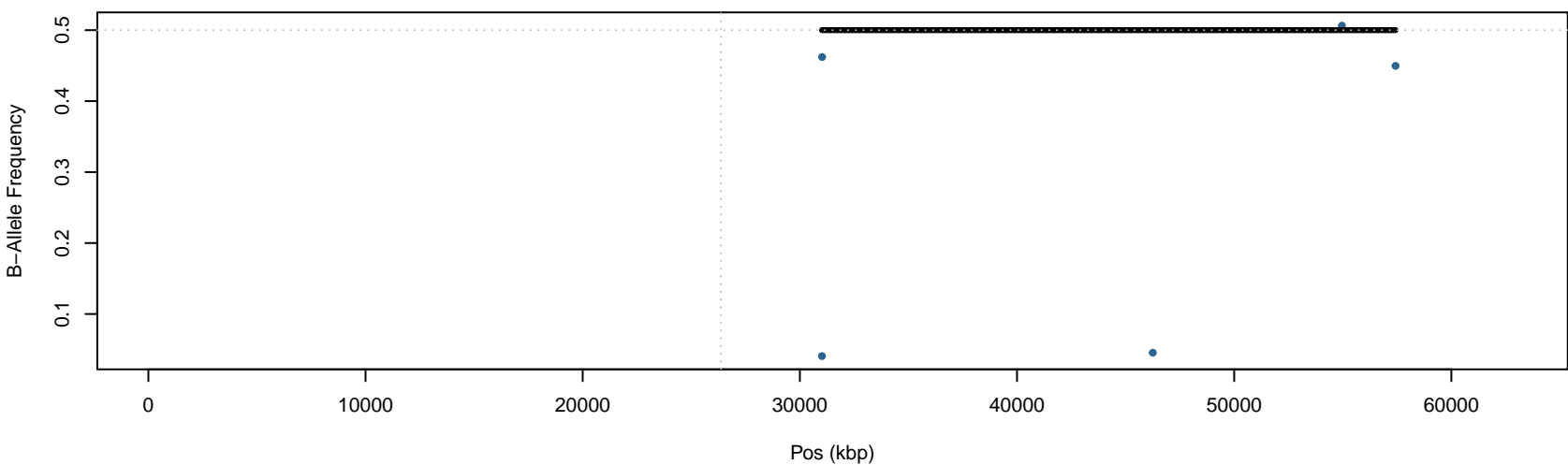
Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 18



Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 19

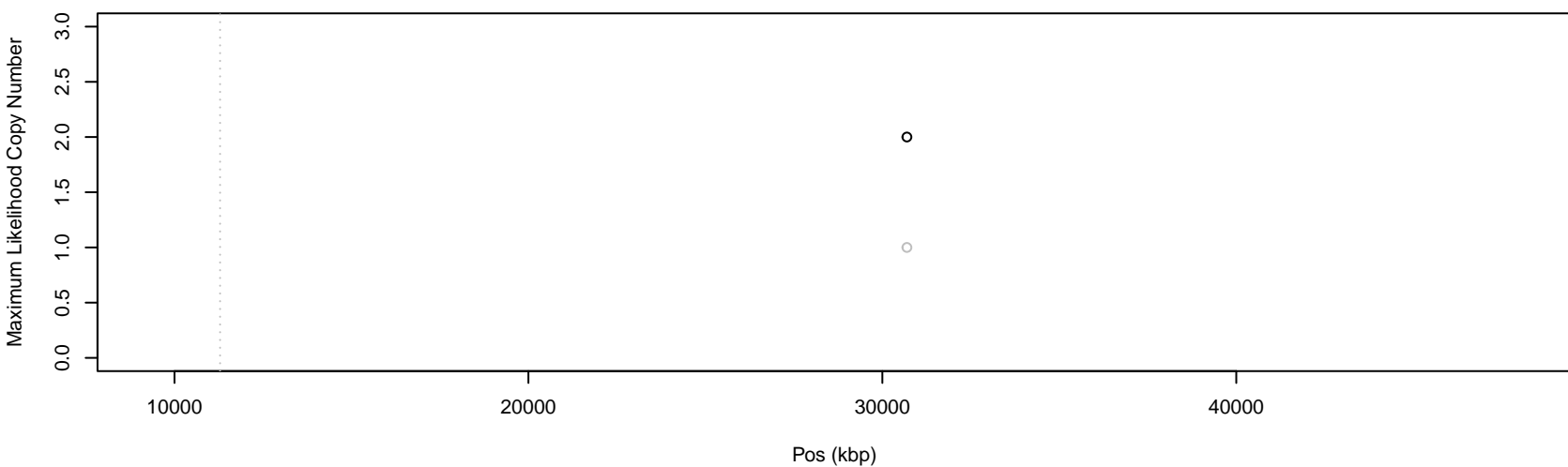
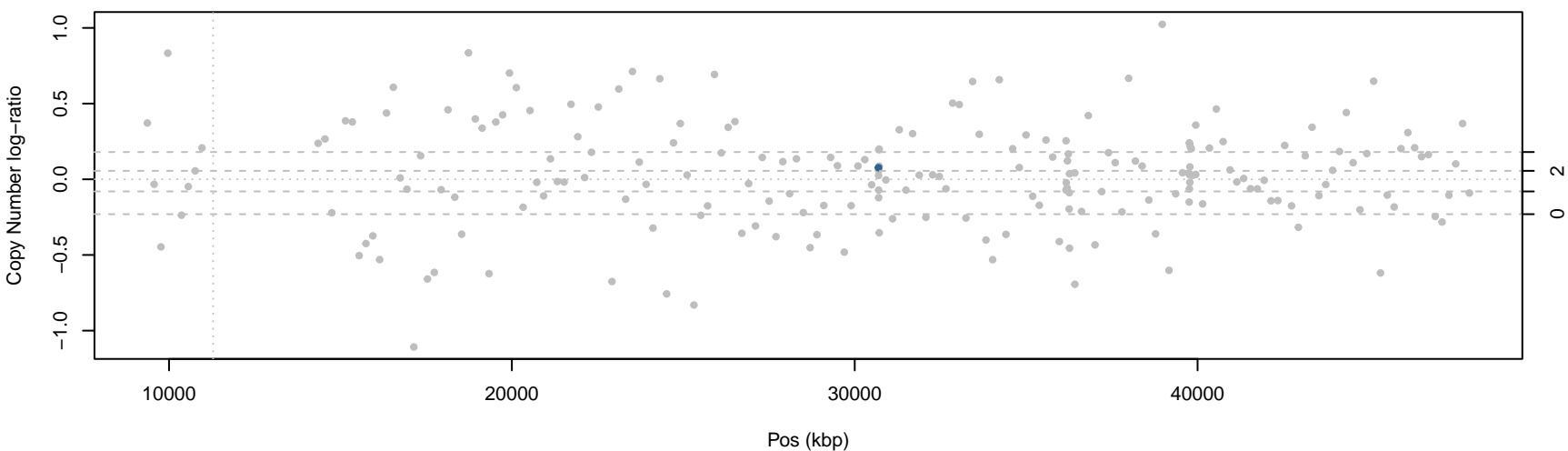
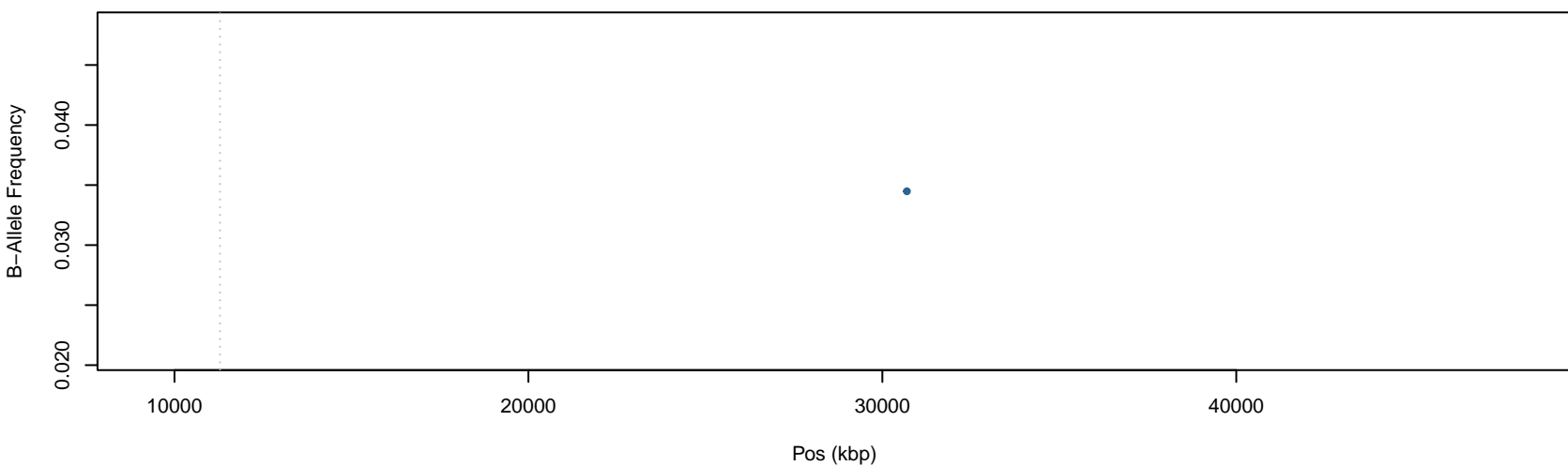


Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 20





Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 21



Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 402.31 GoF: 92.9% Mean coverage: 233;281 Chromosome: 22

