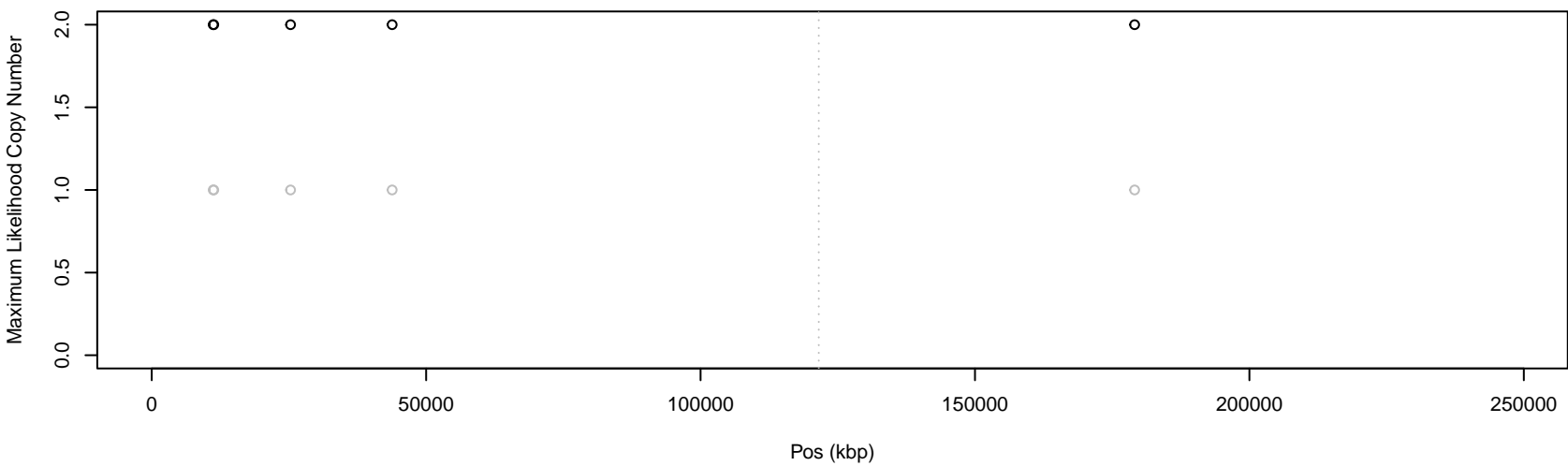
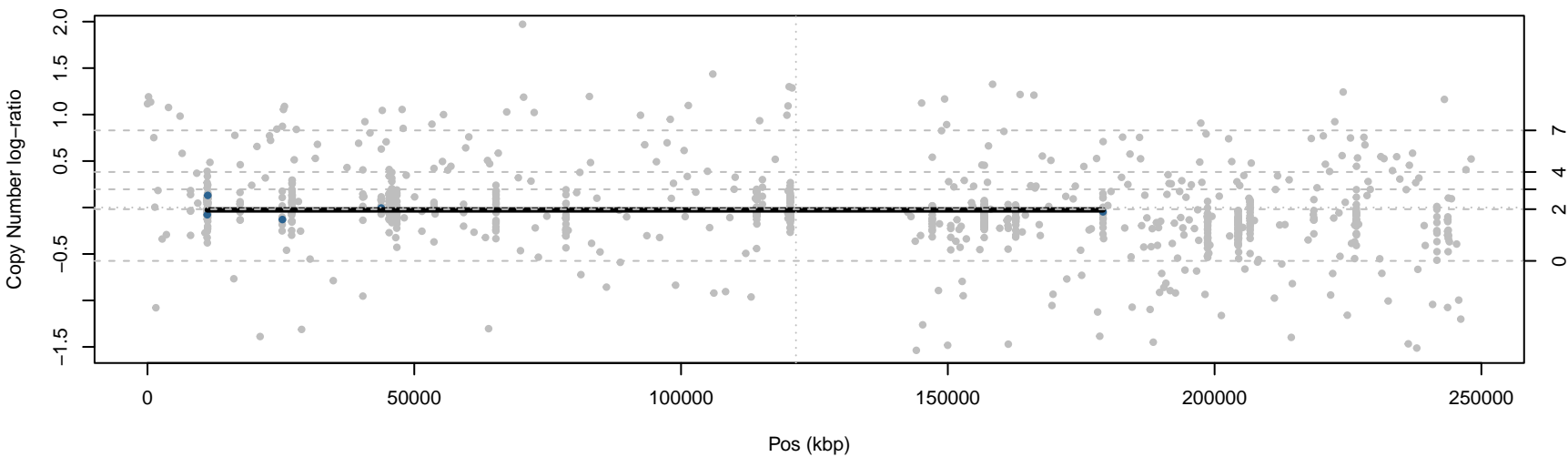
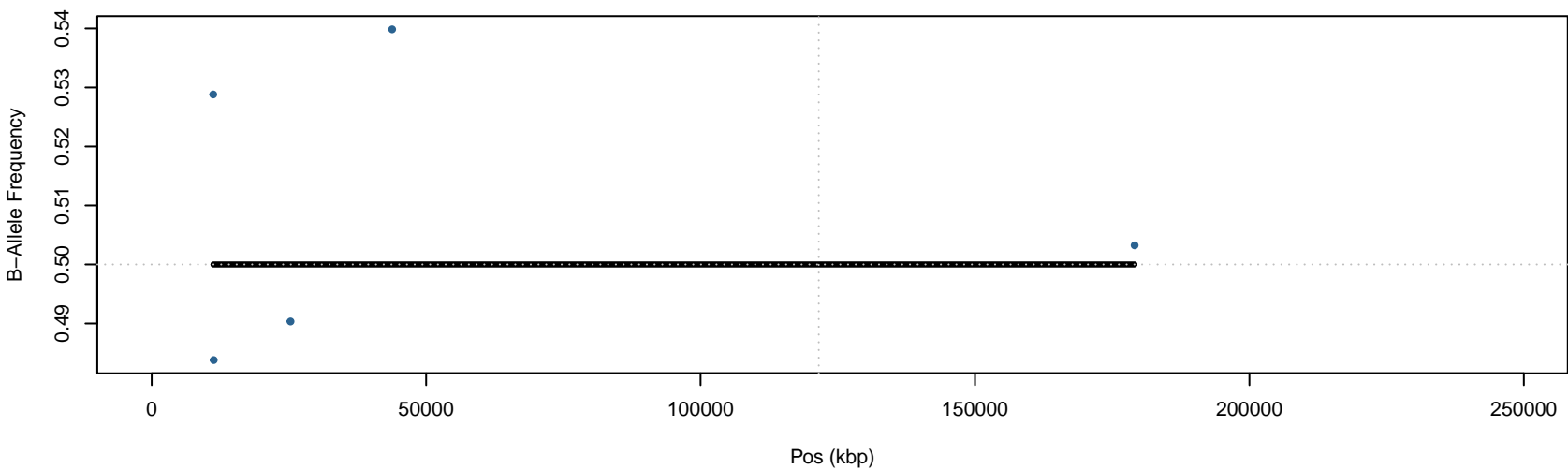
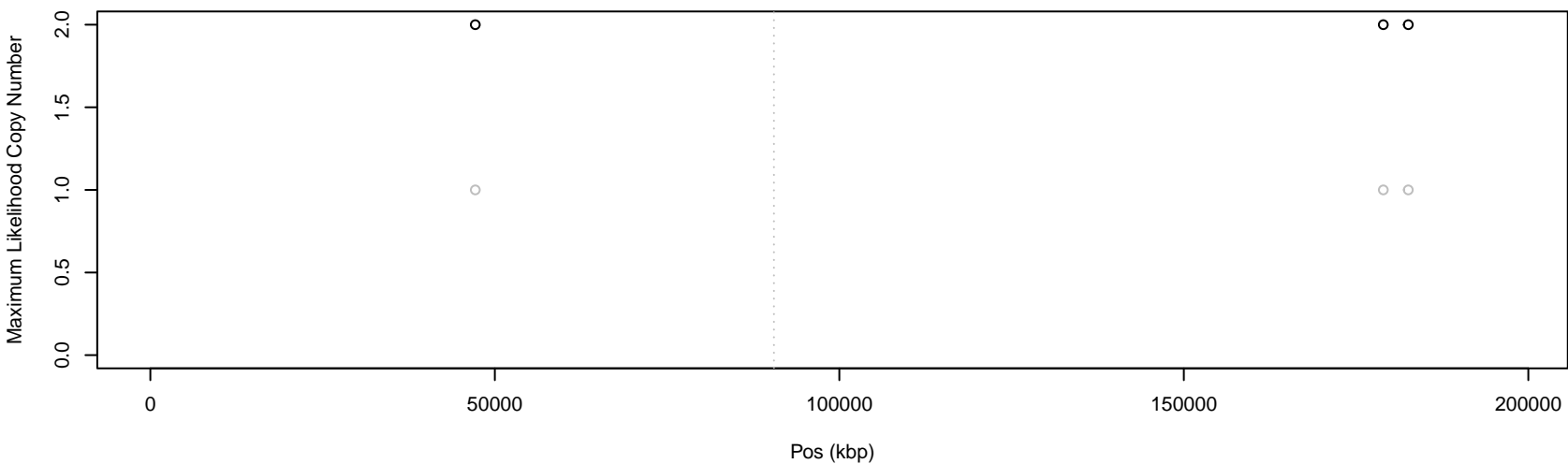
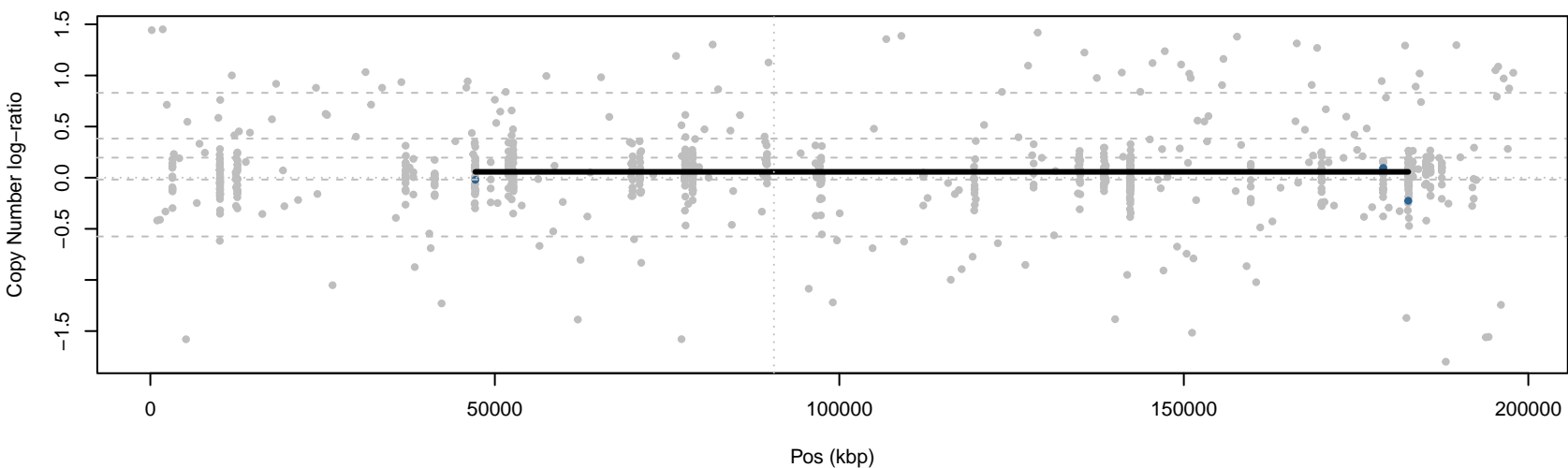
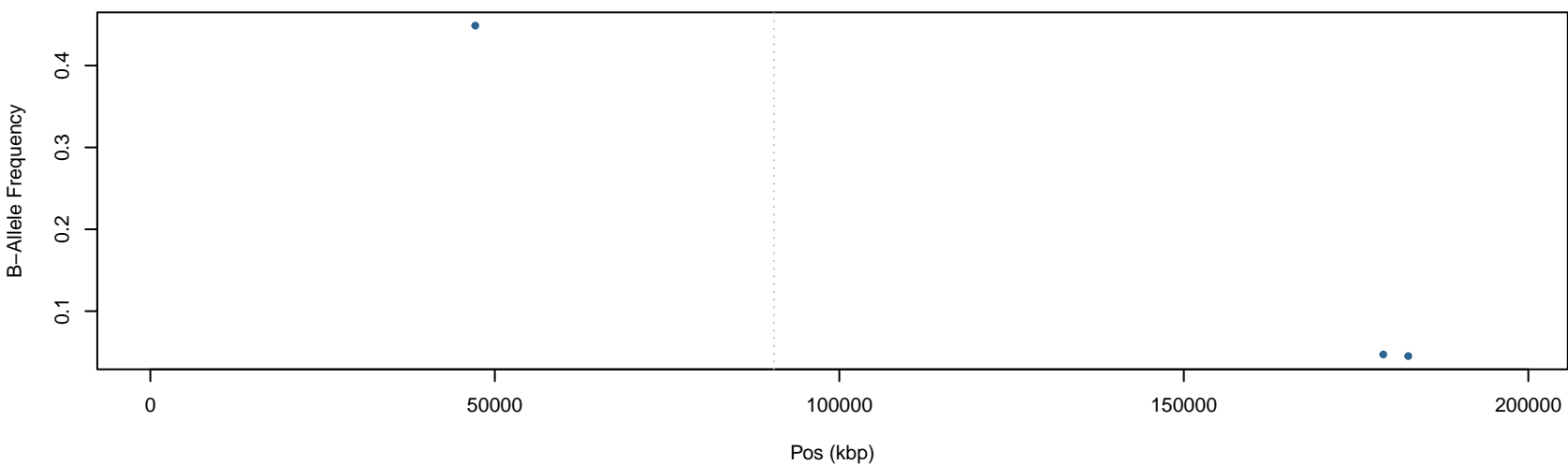


Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 1

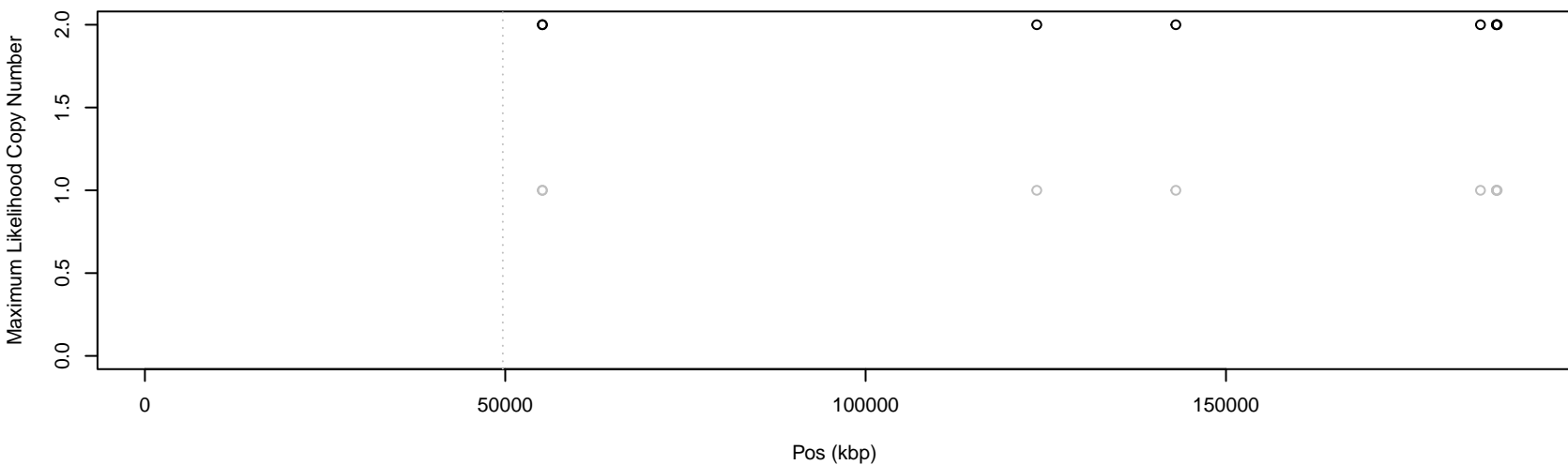
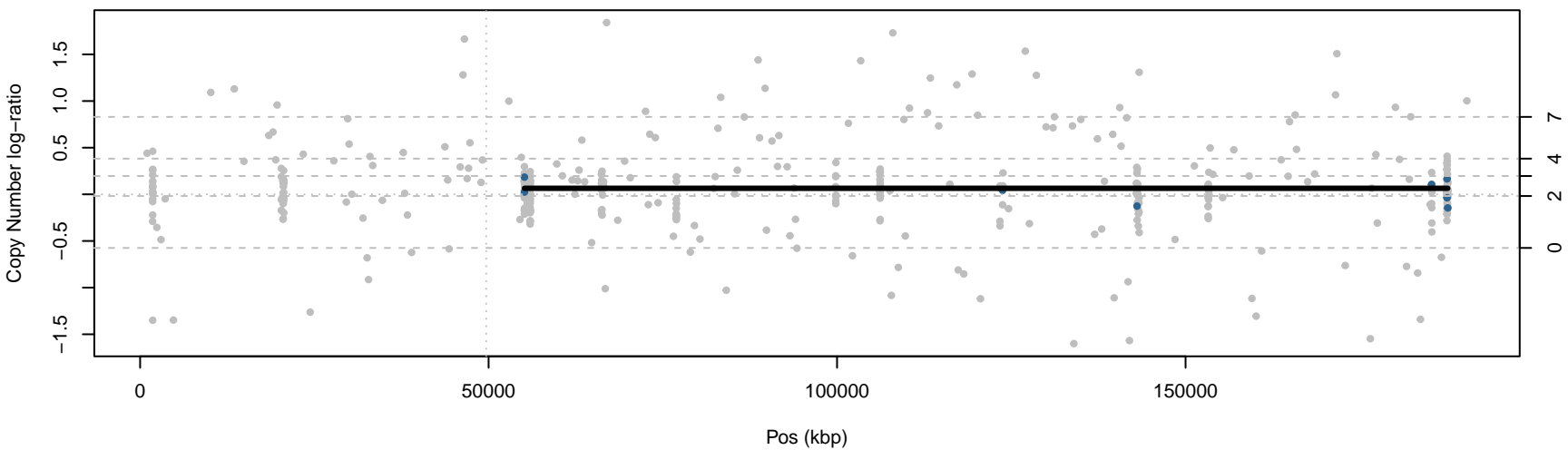
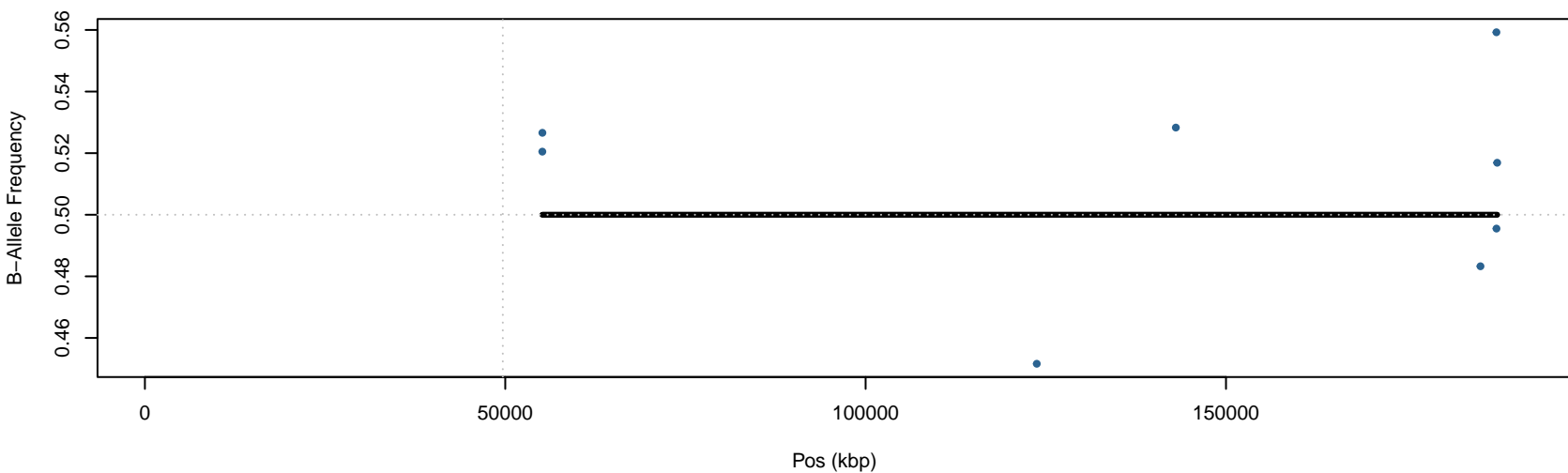




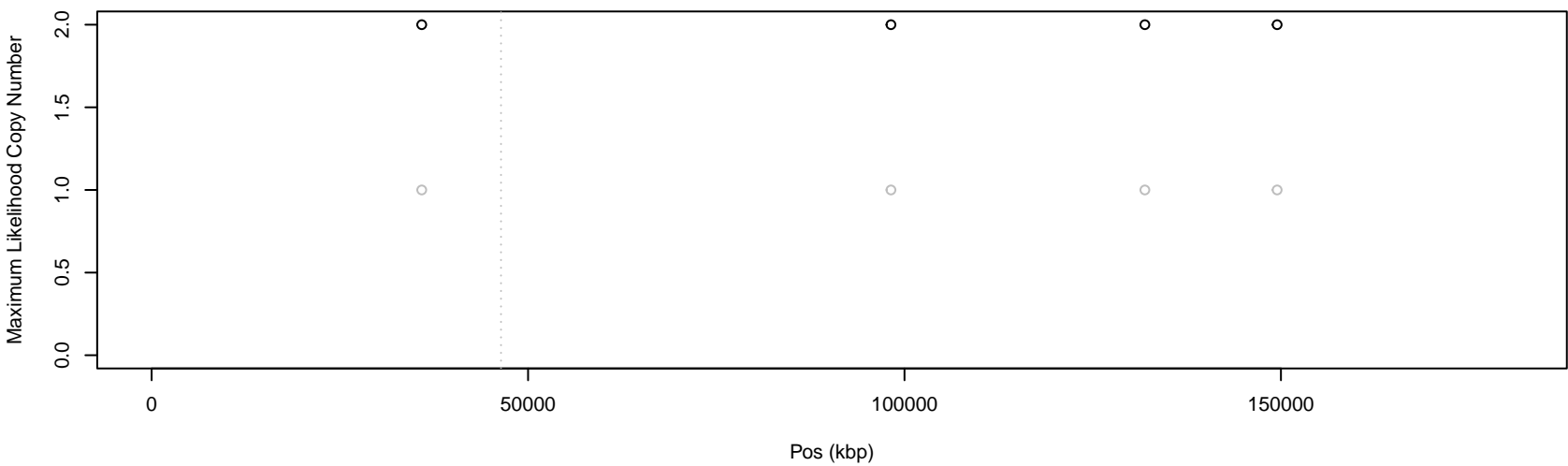
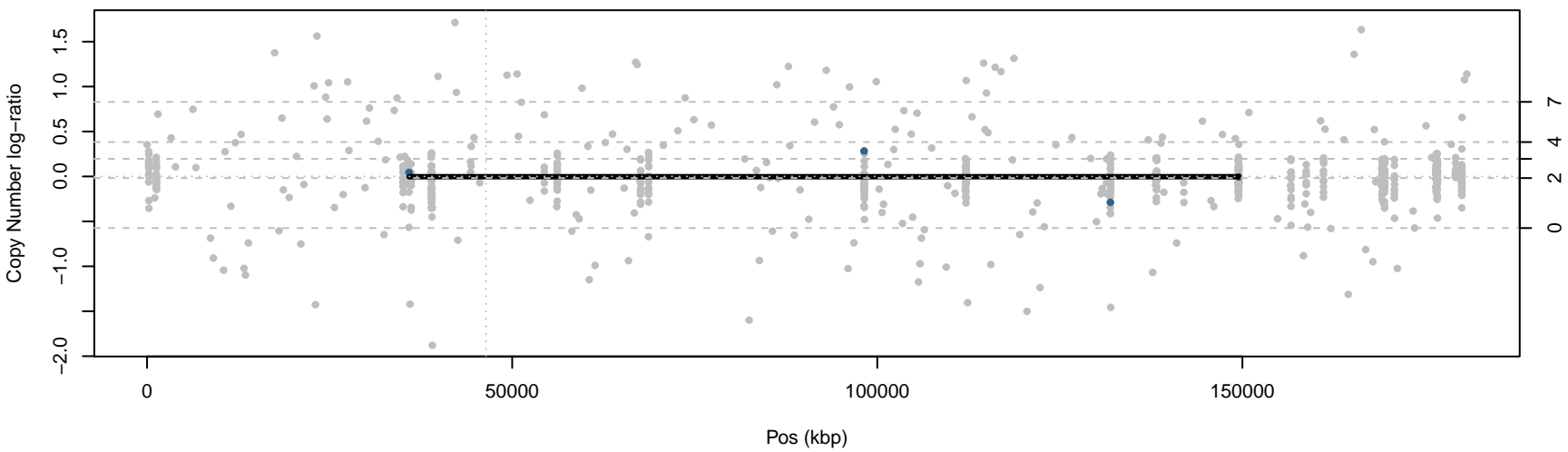
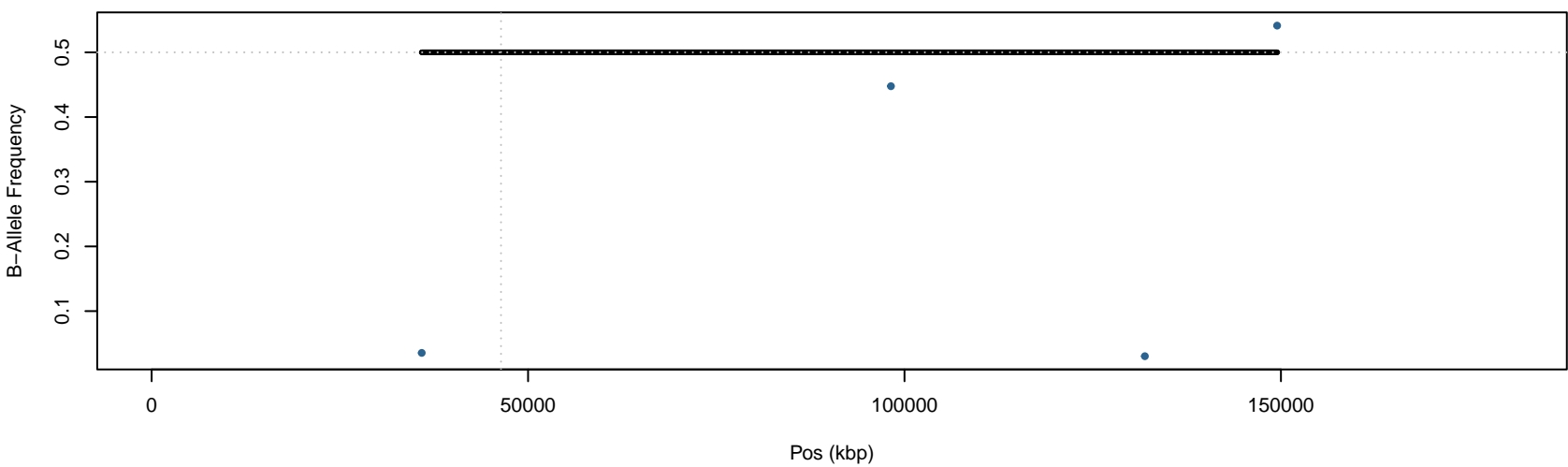
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 3



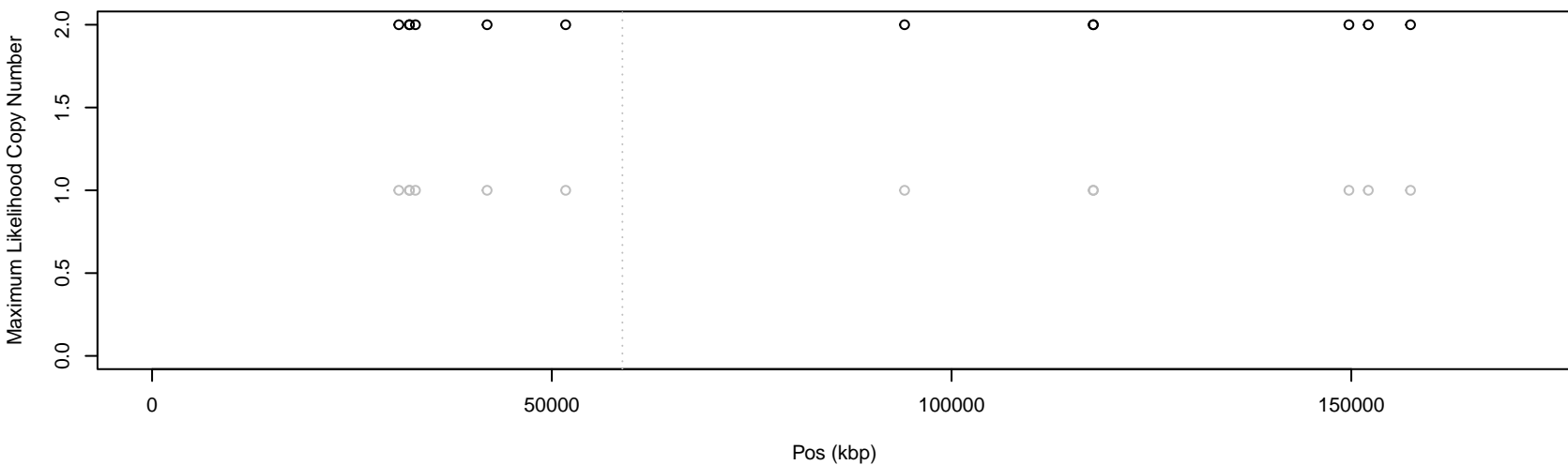
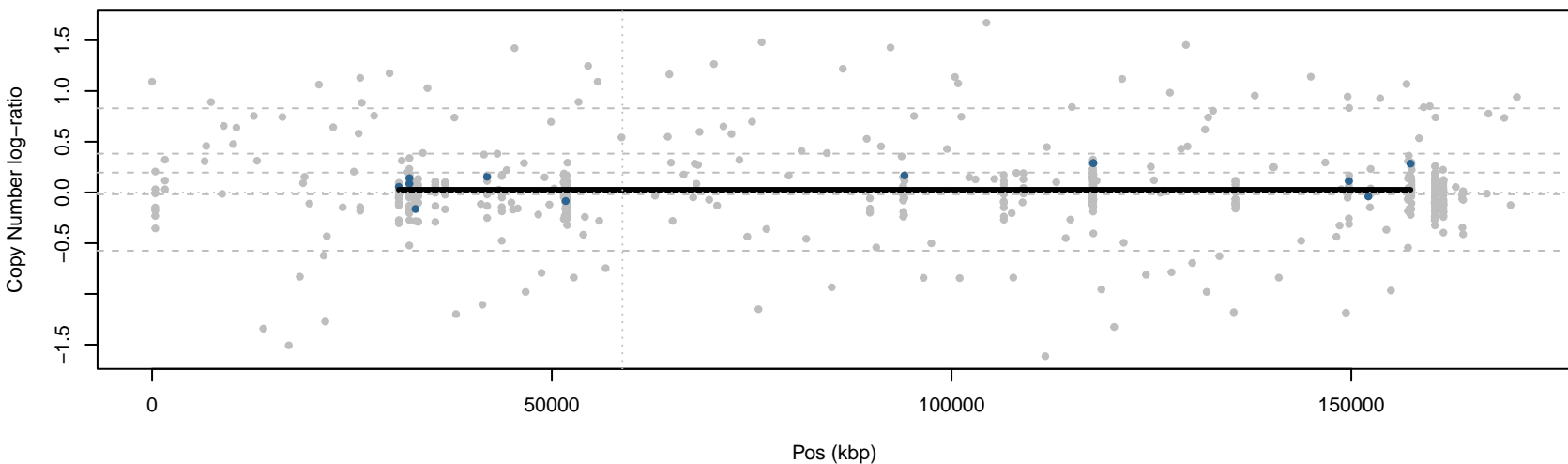
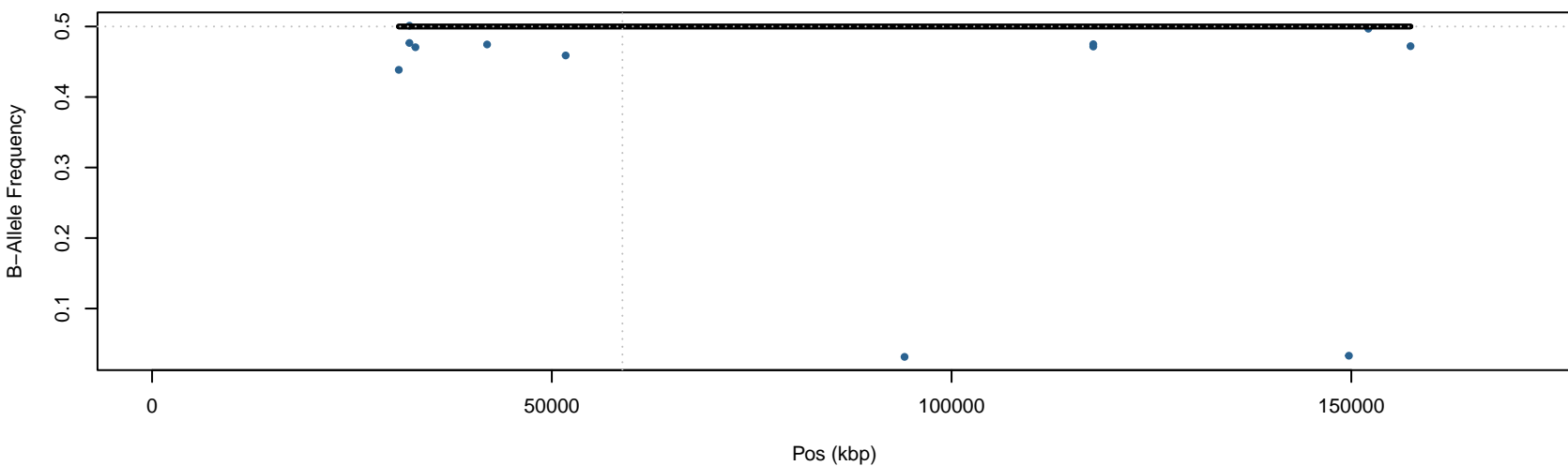
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 4



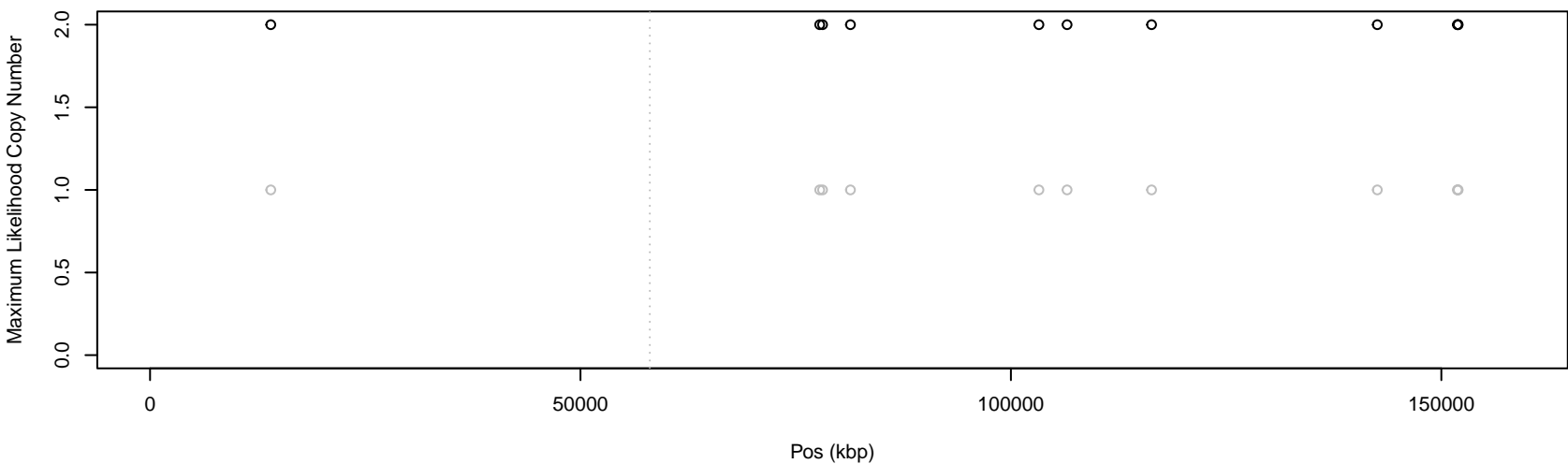
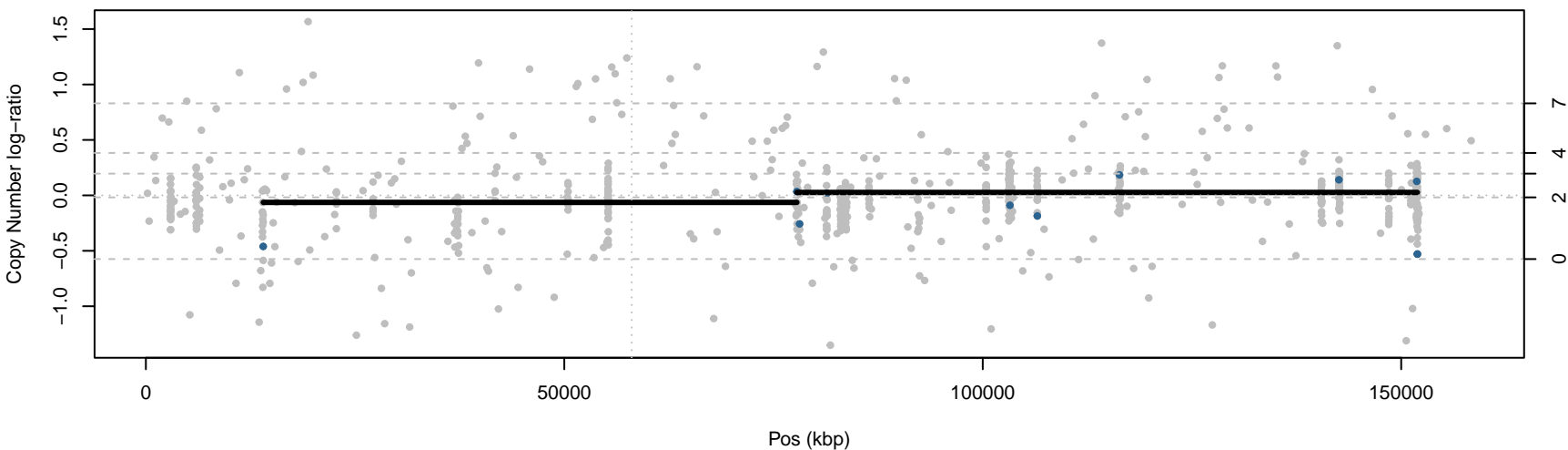
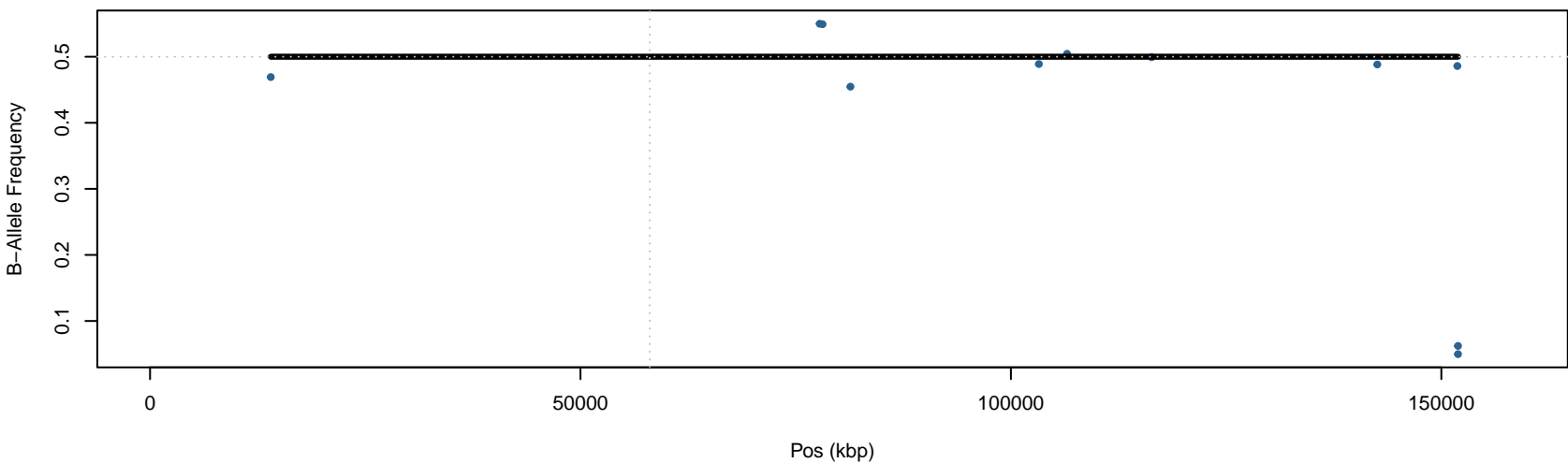
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 5



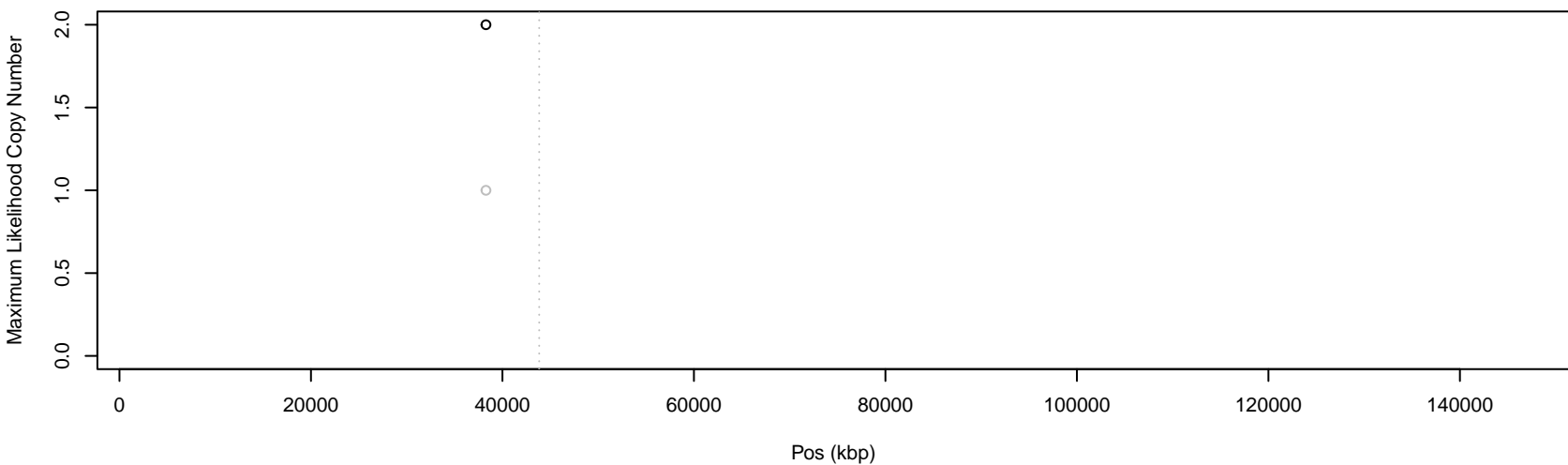
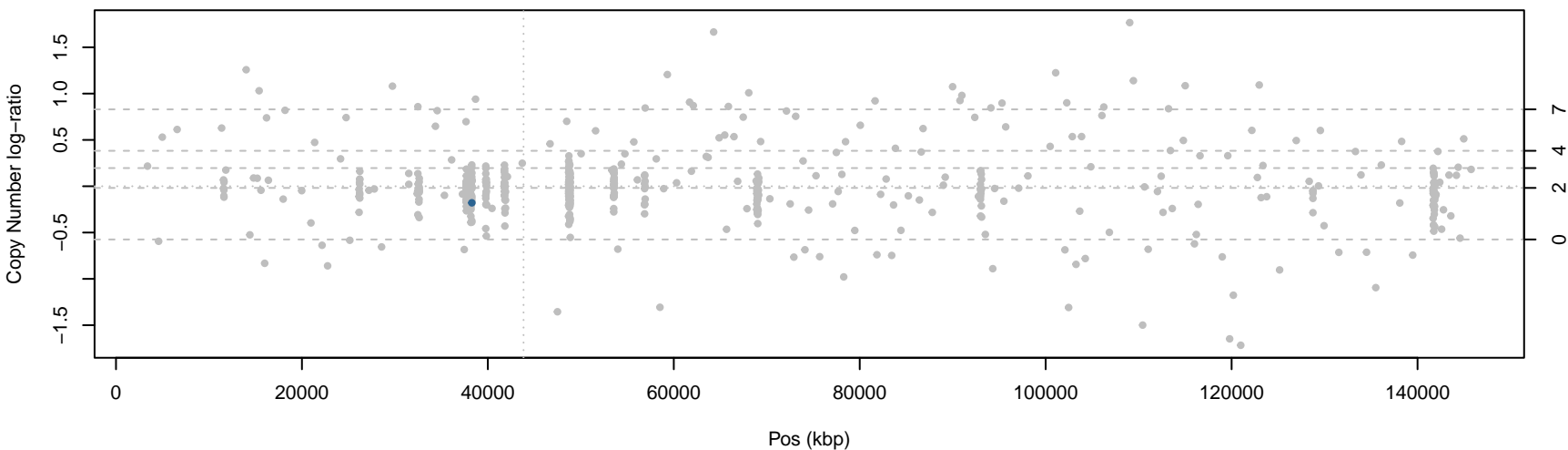
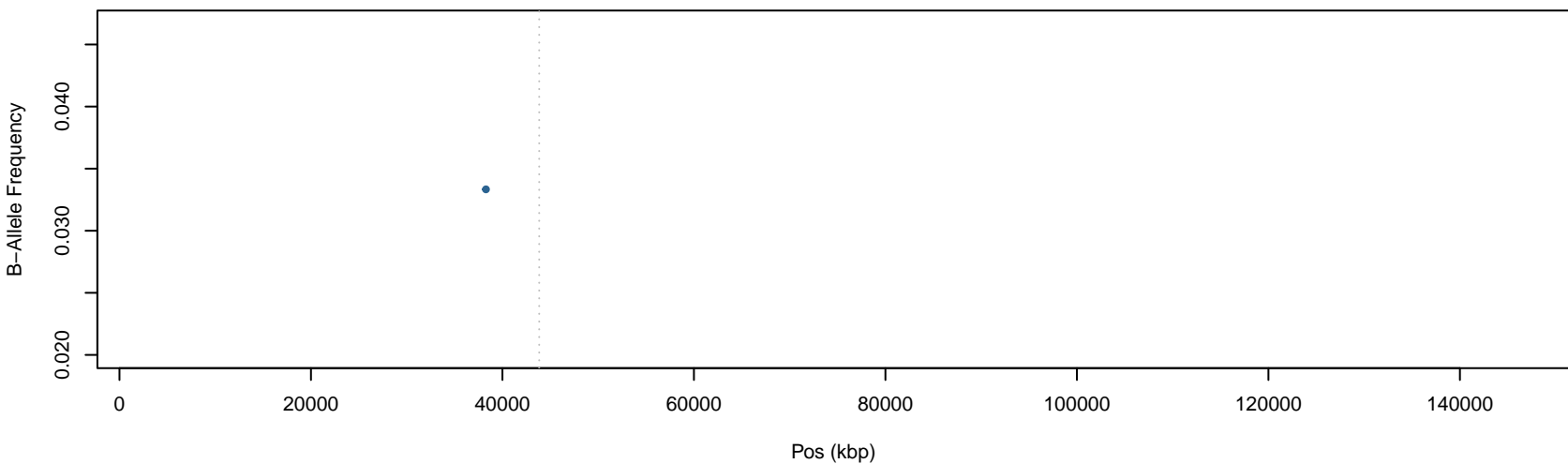
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 6



Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 7

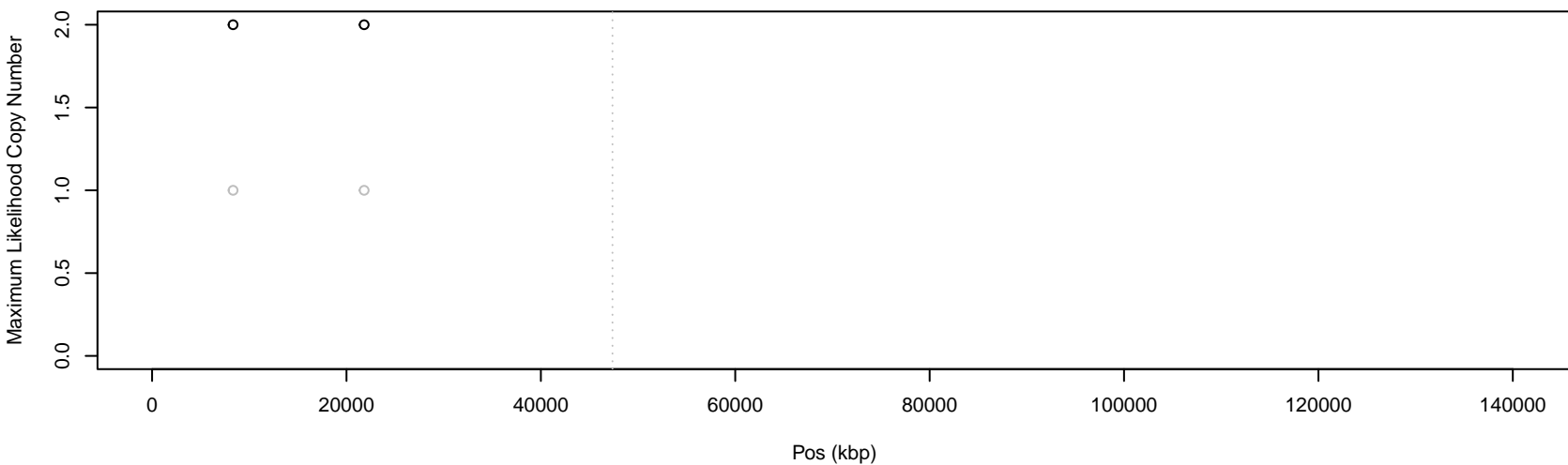
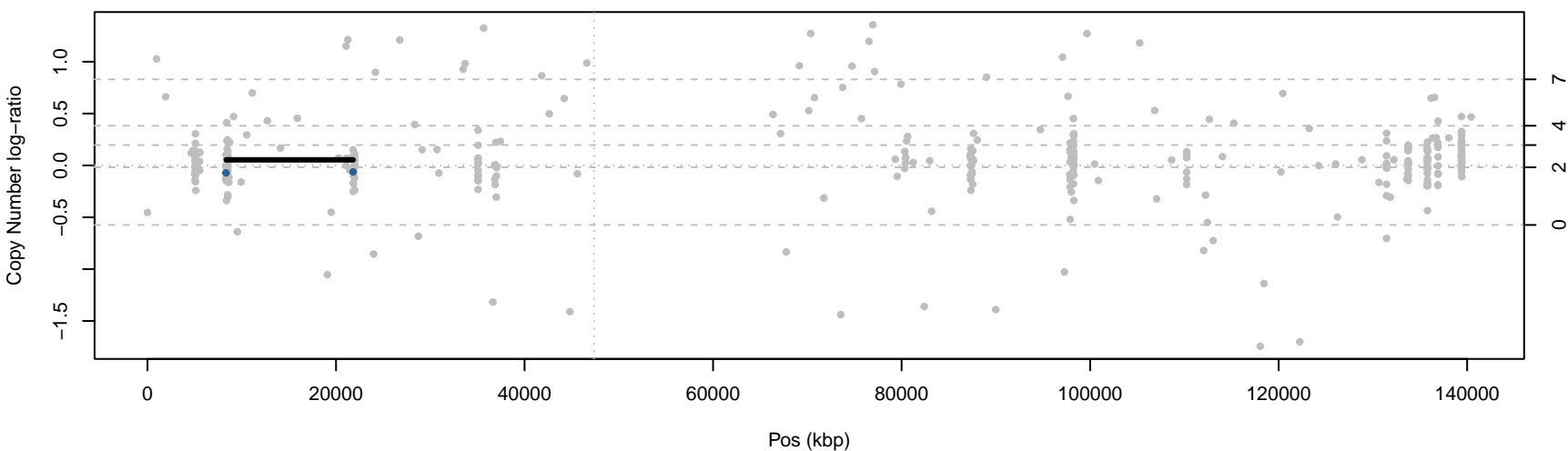
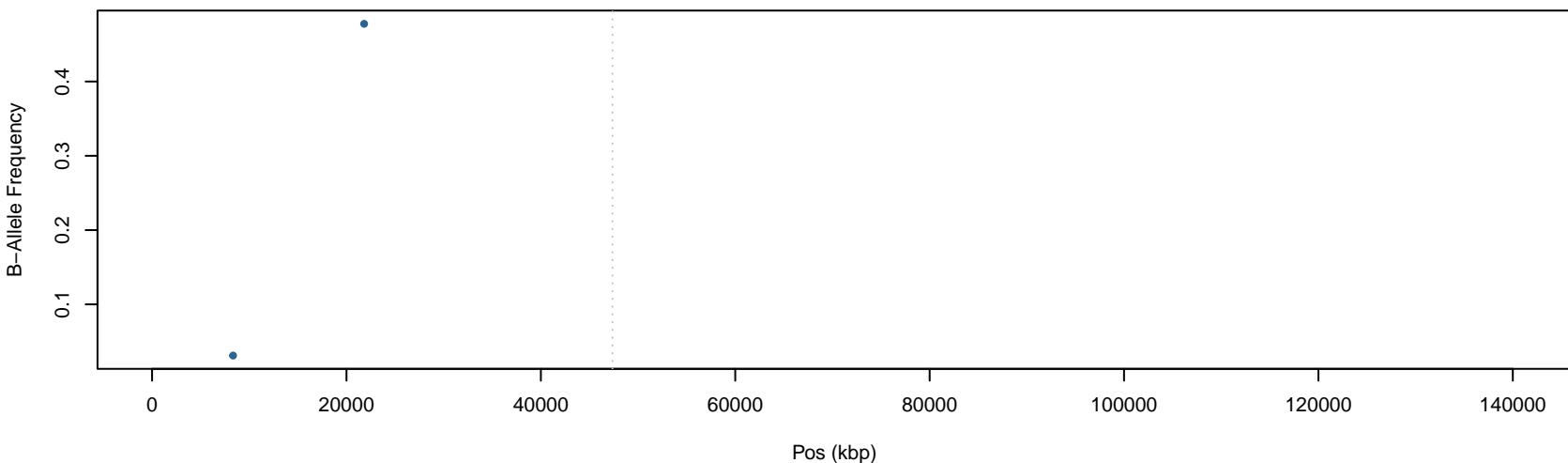


Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 8

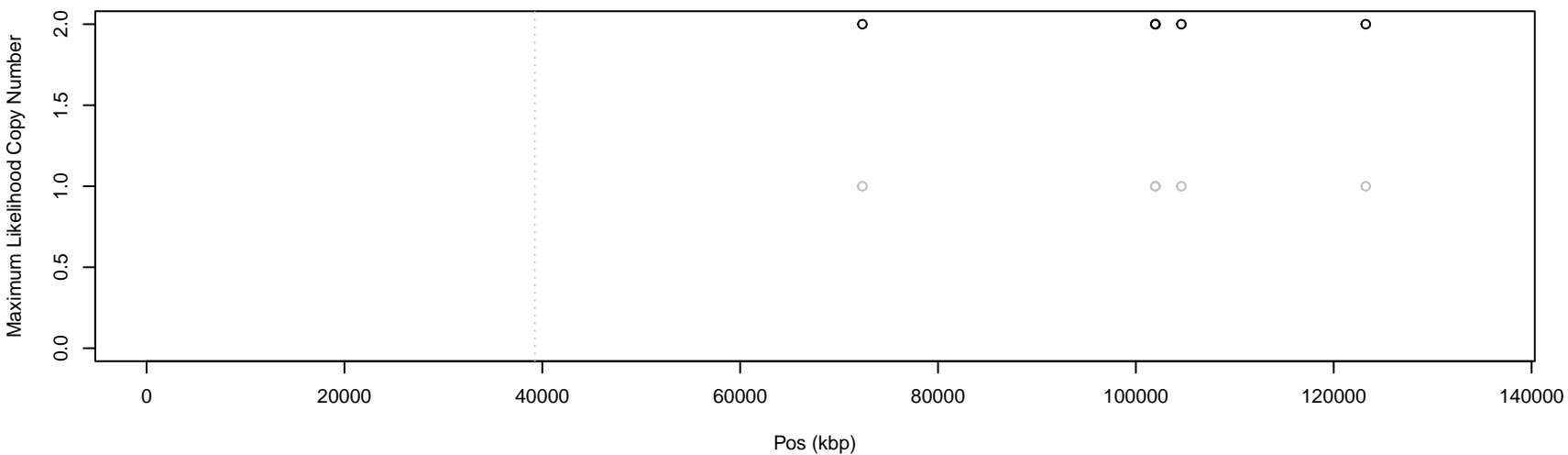
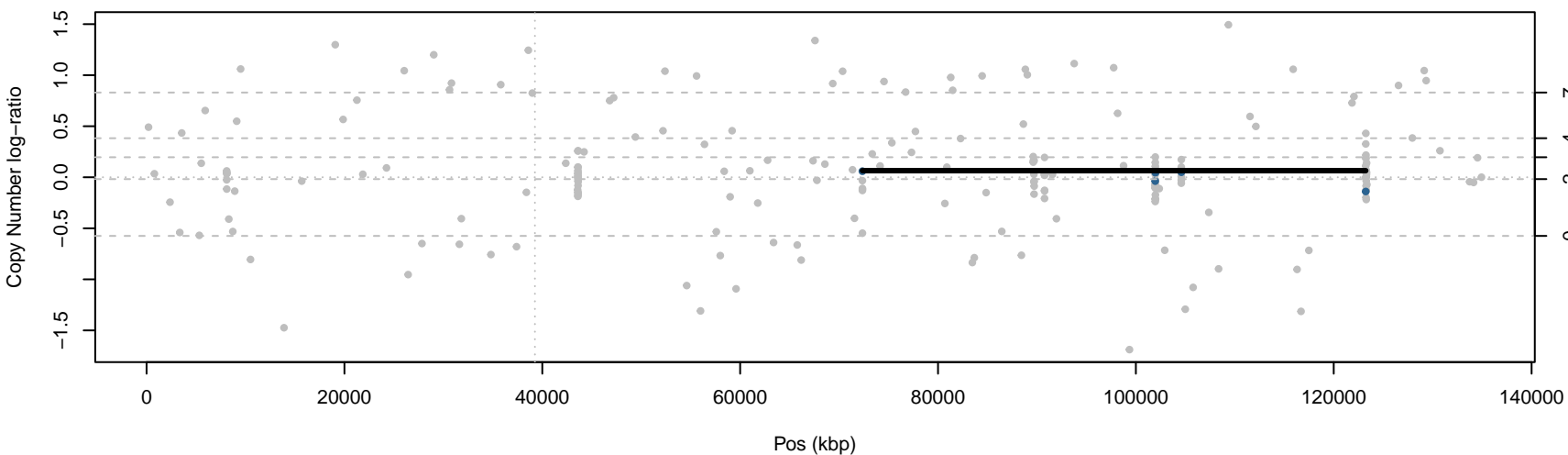
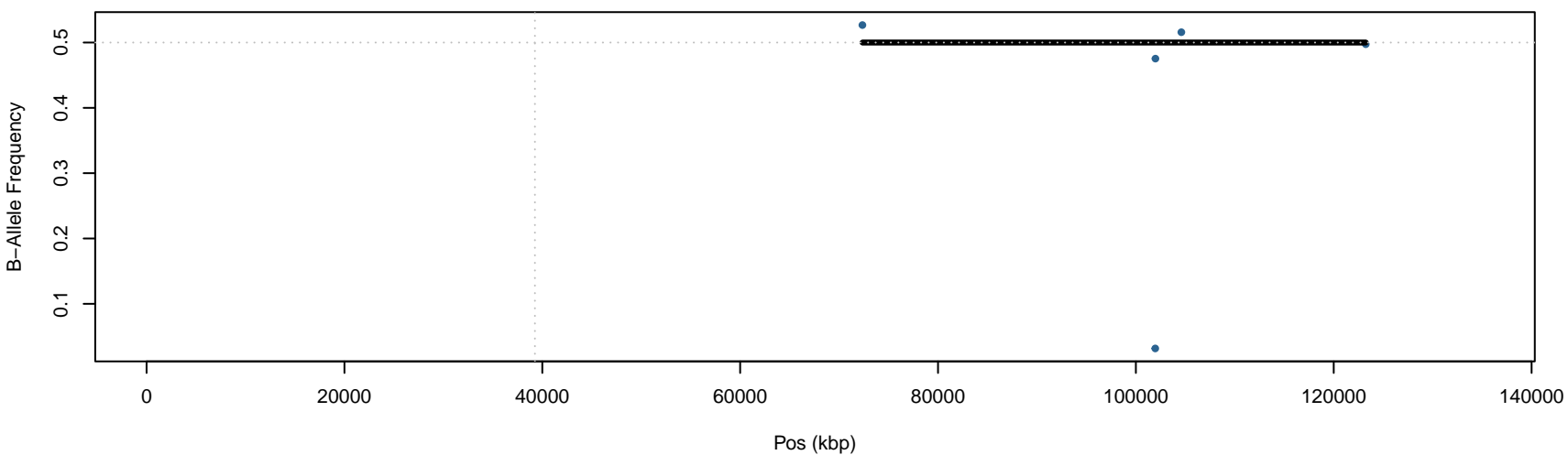




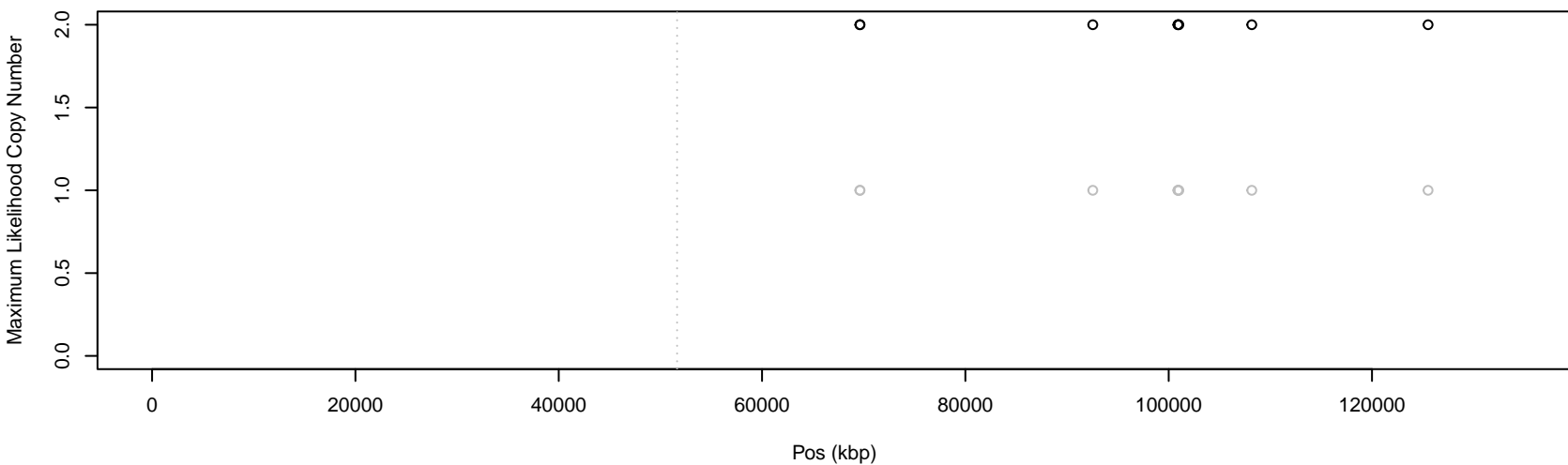
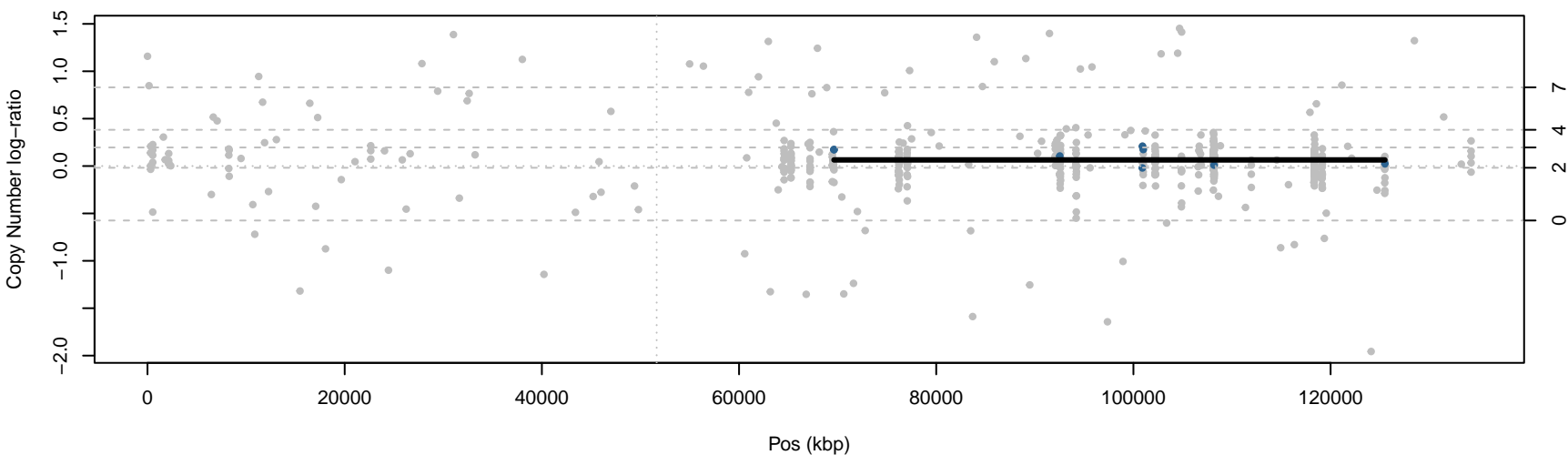
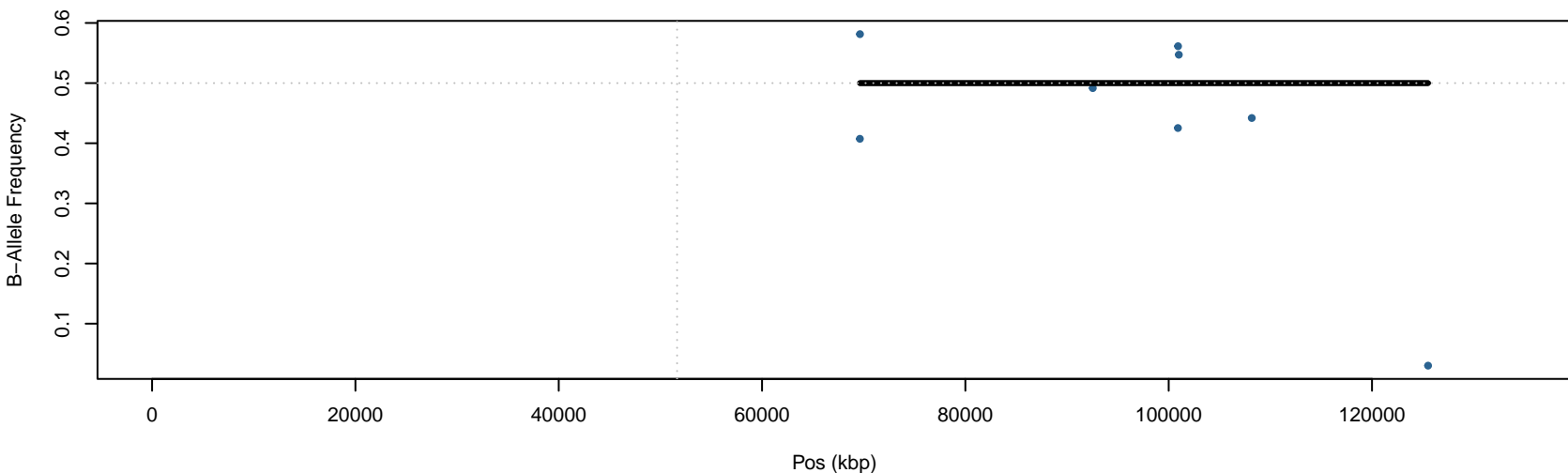
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 9



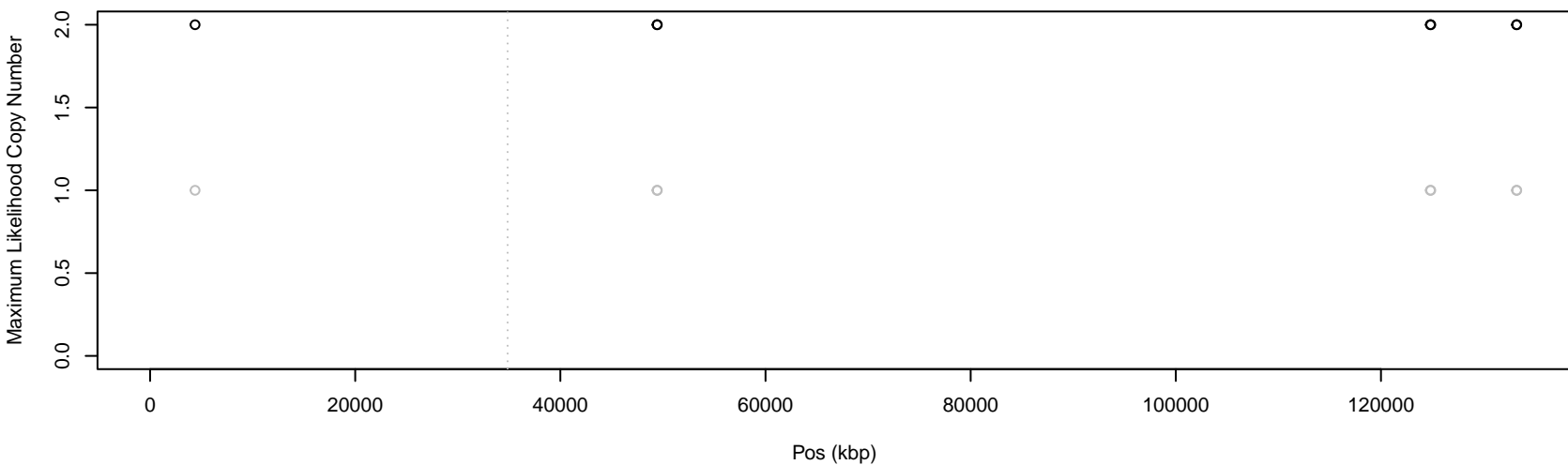
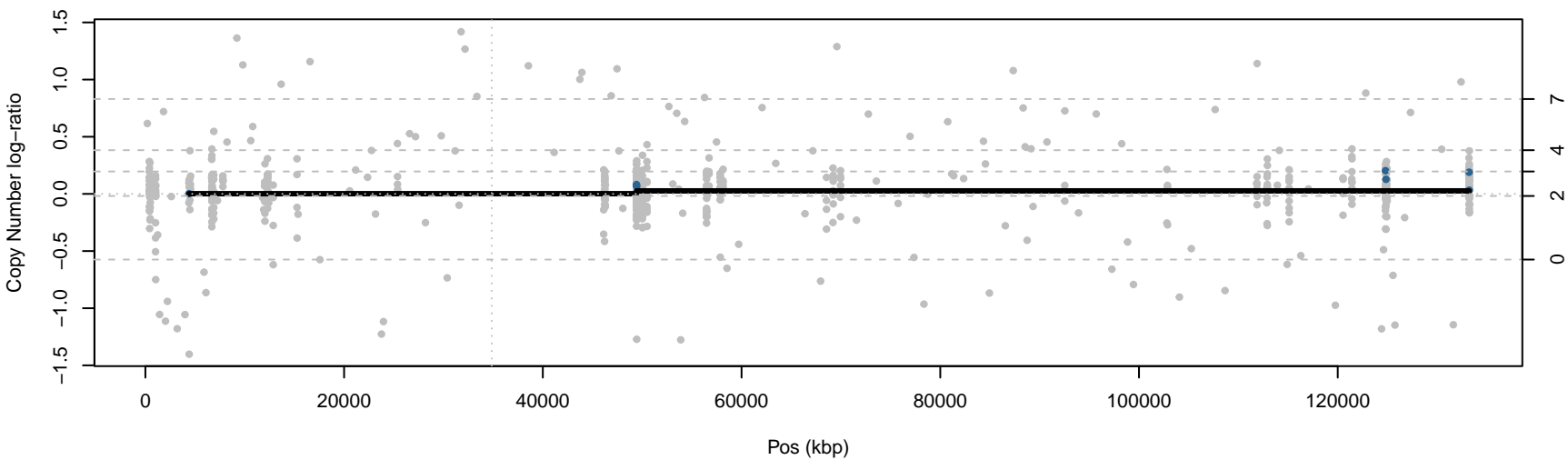
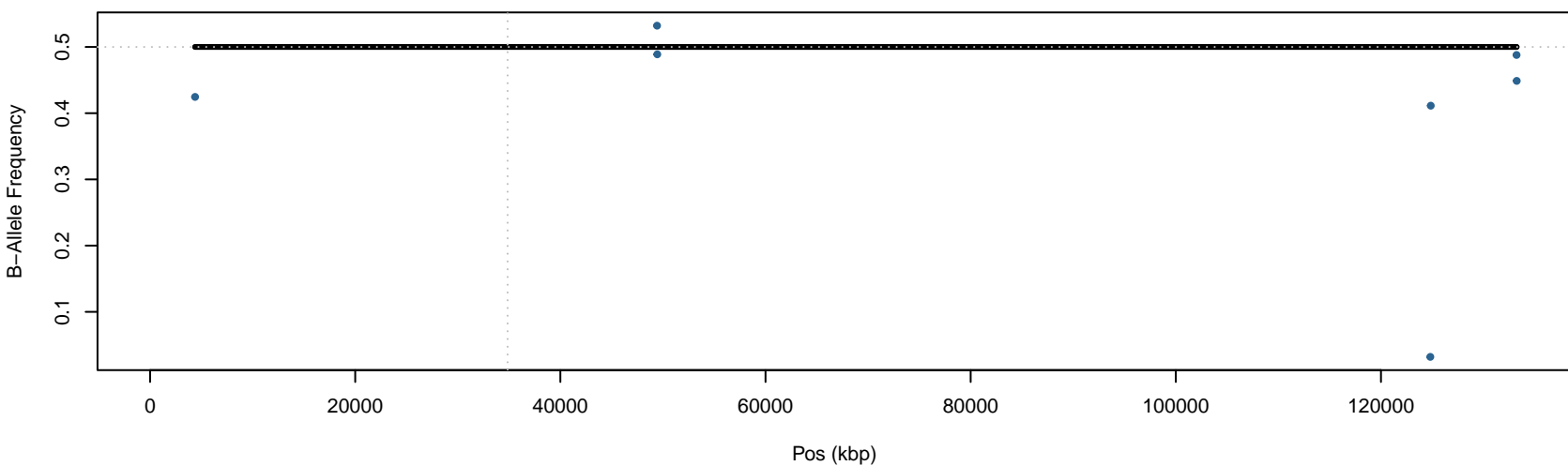
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 10



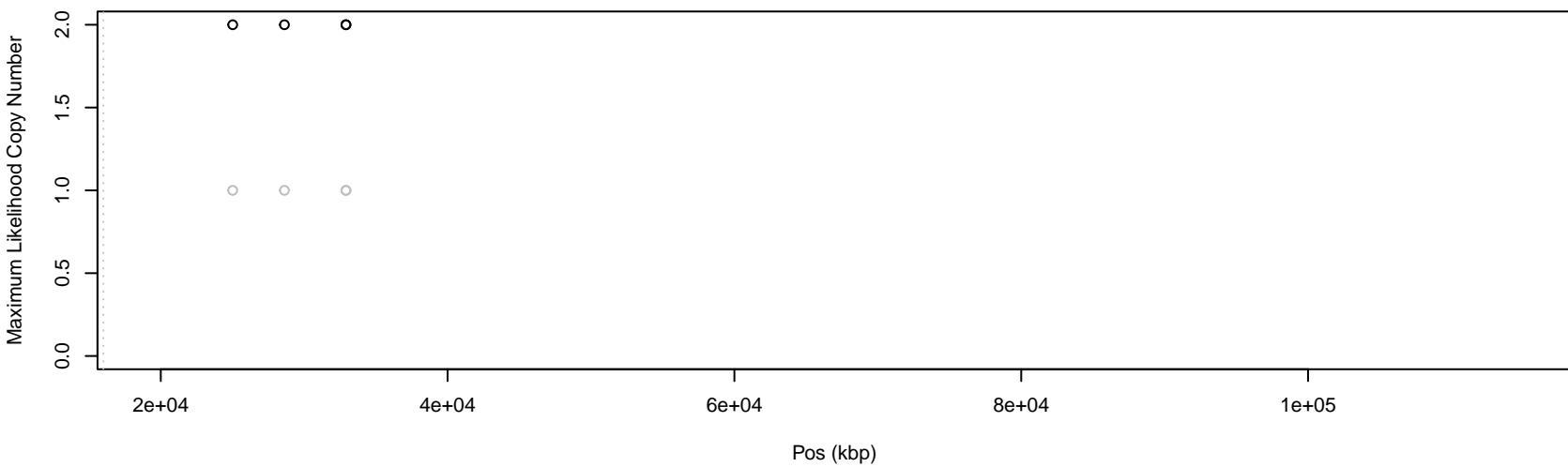
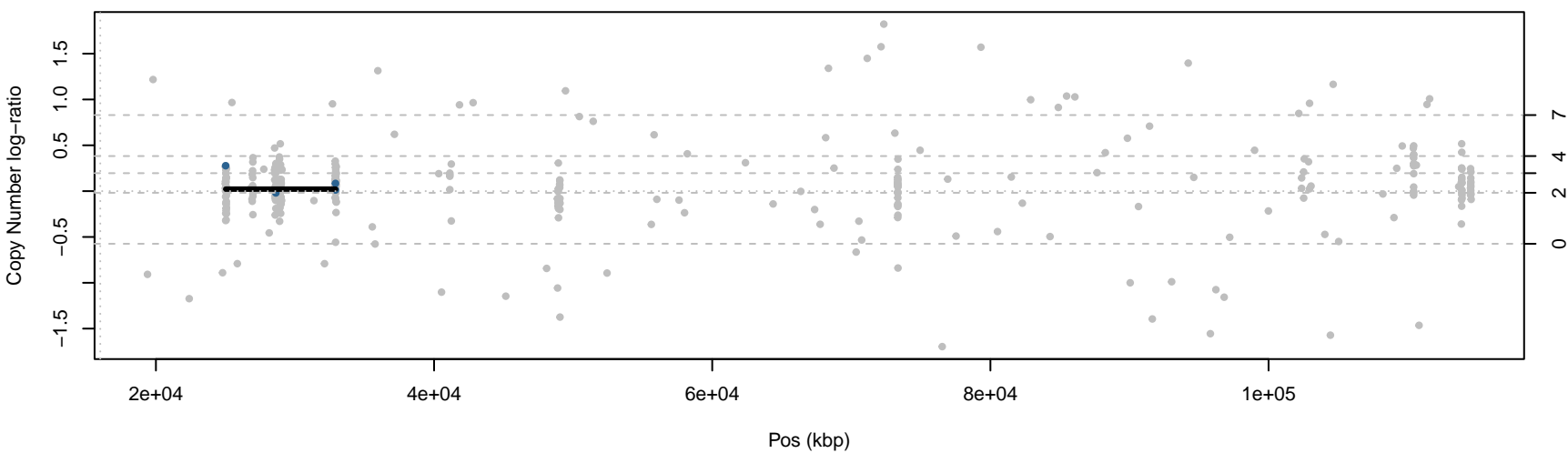
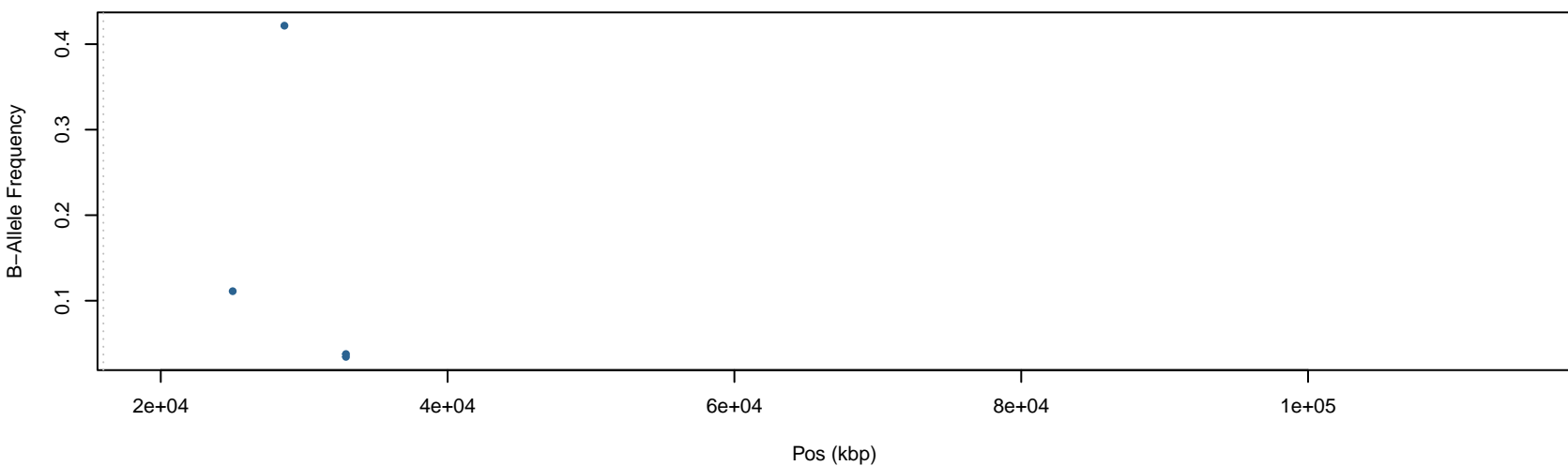
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 11



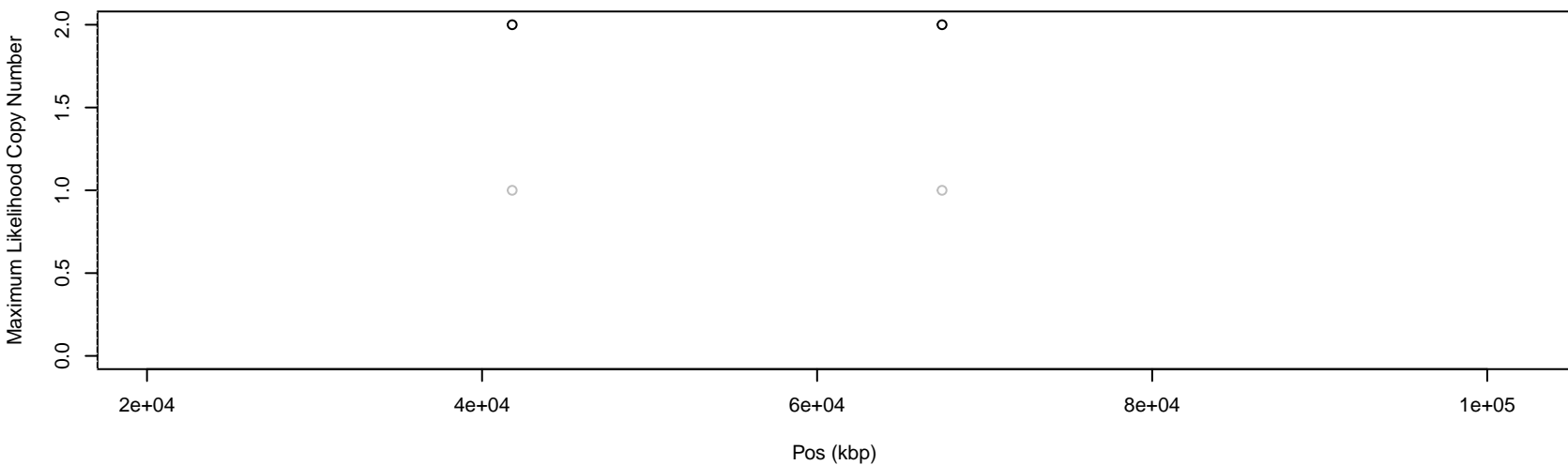
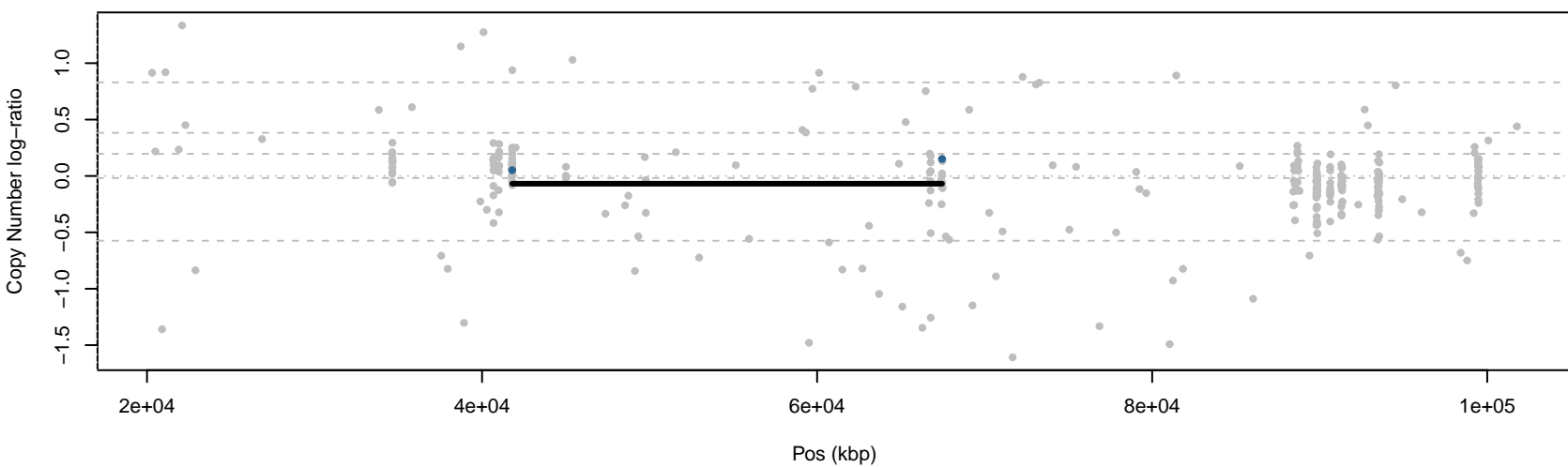
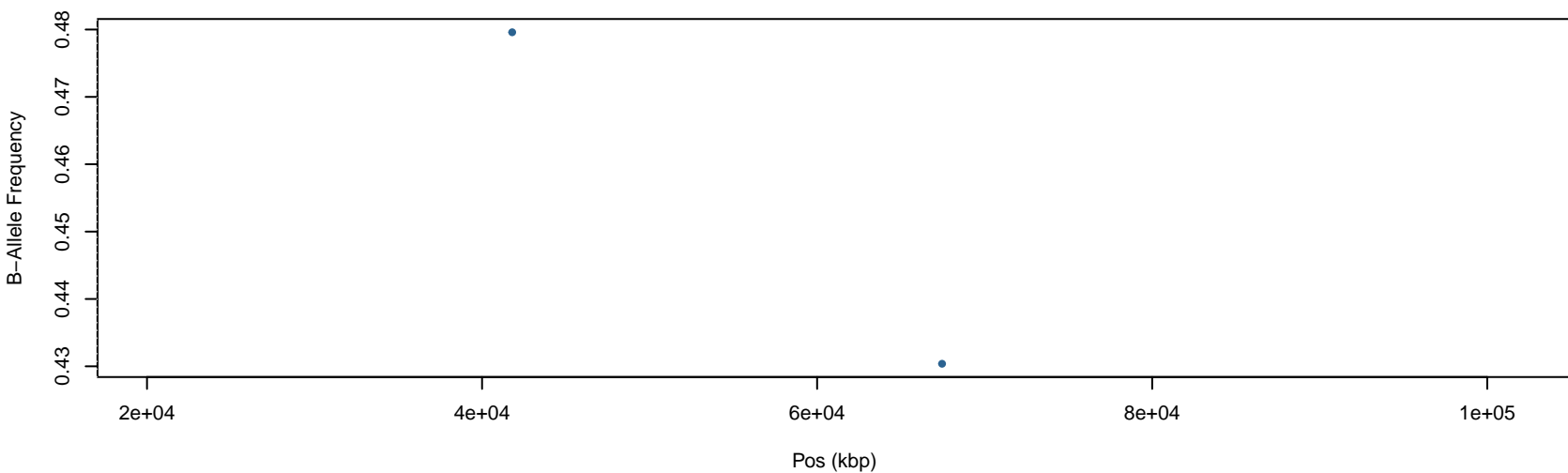
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 12



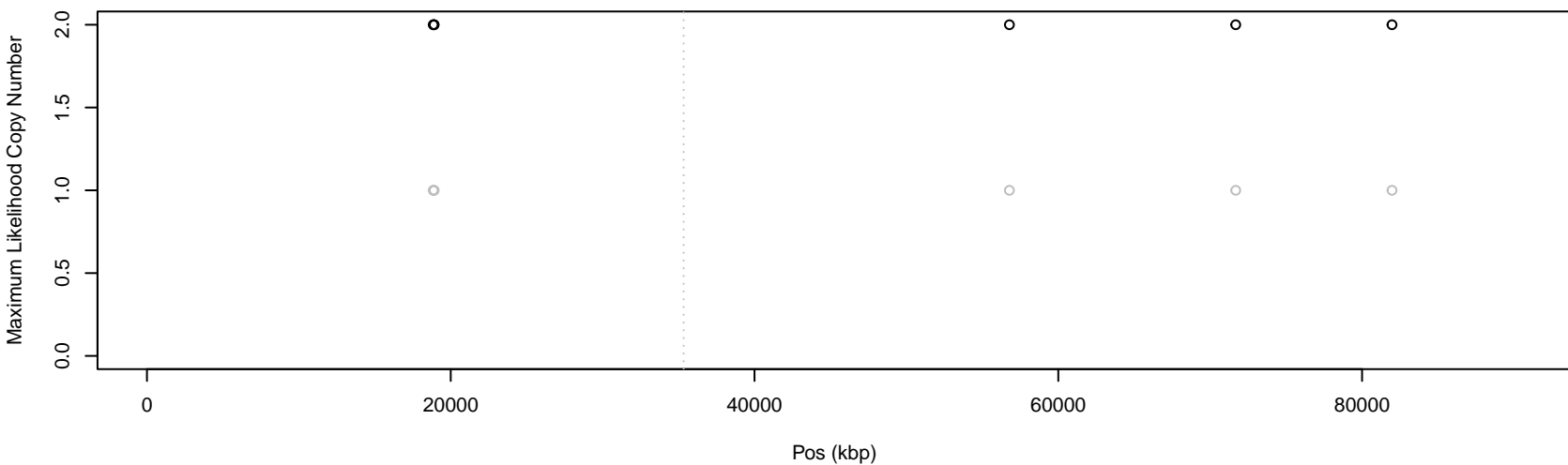
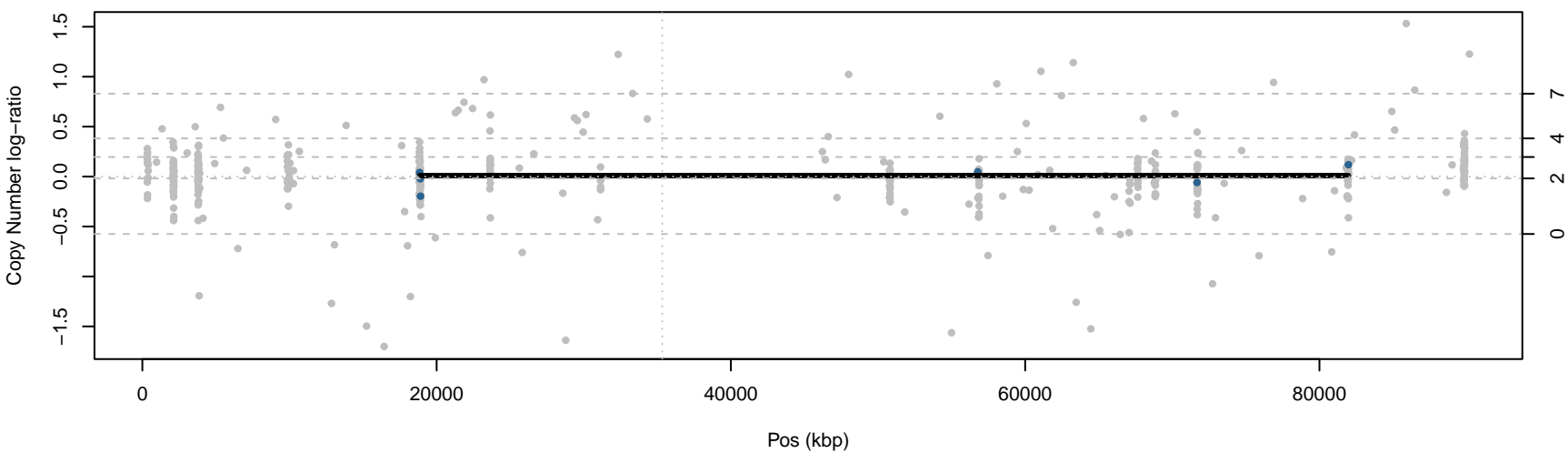
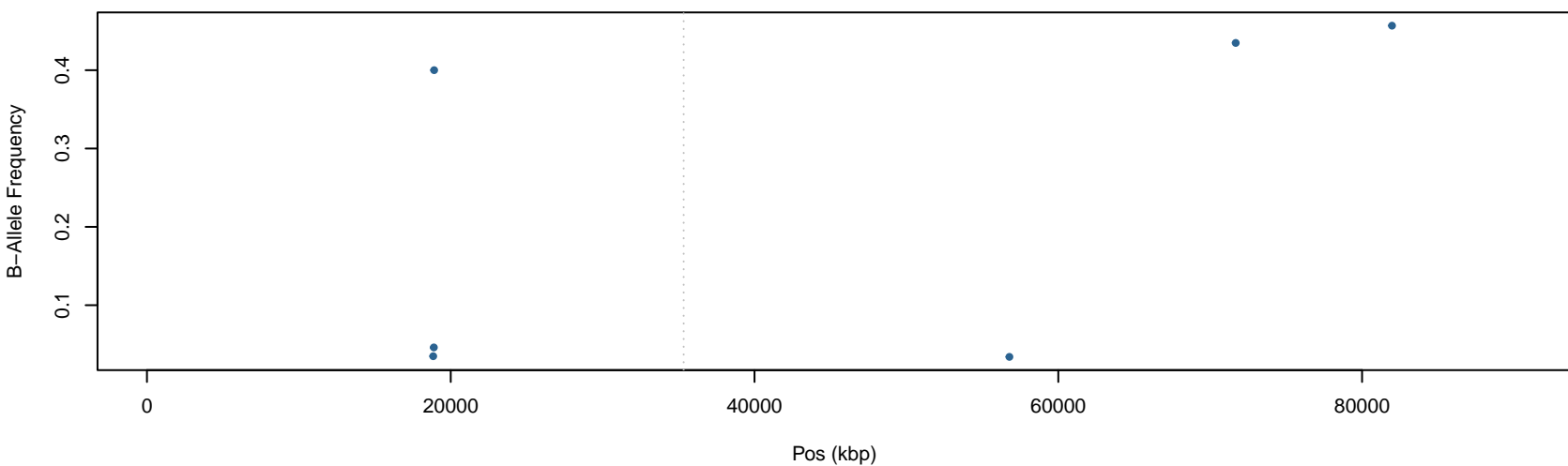
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 13



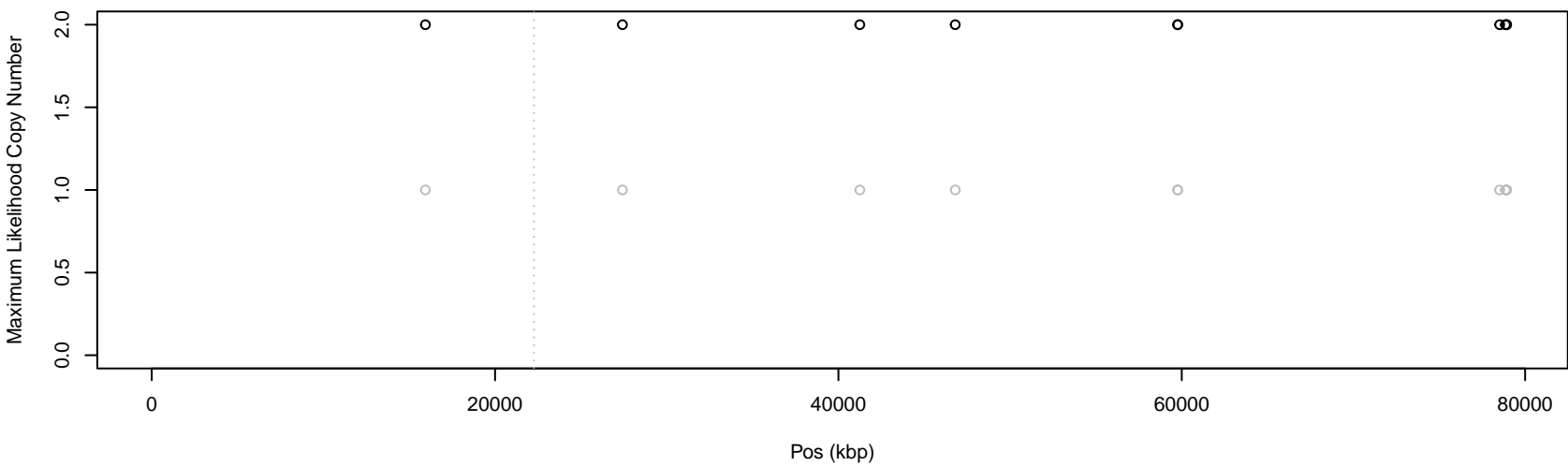
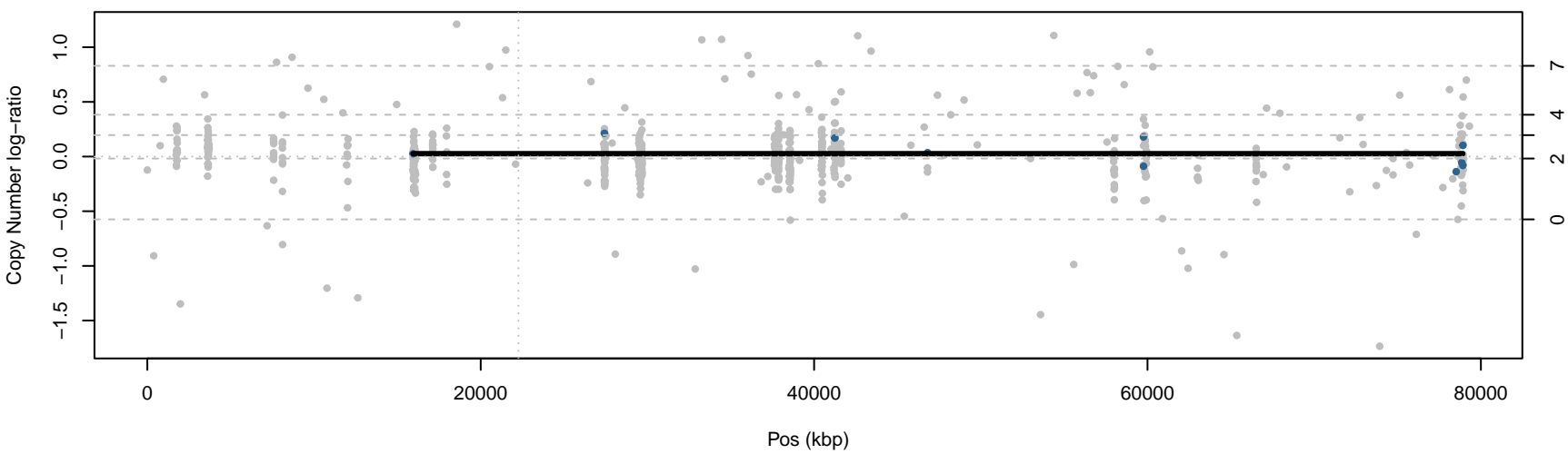
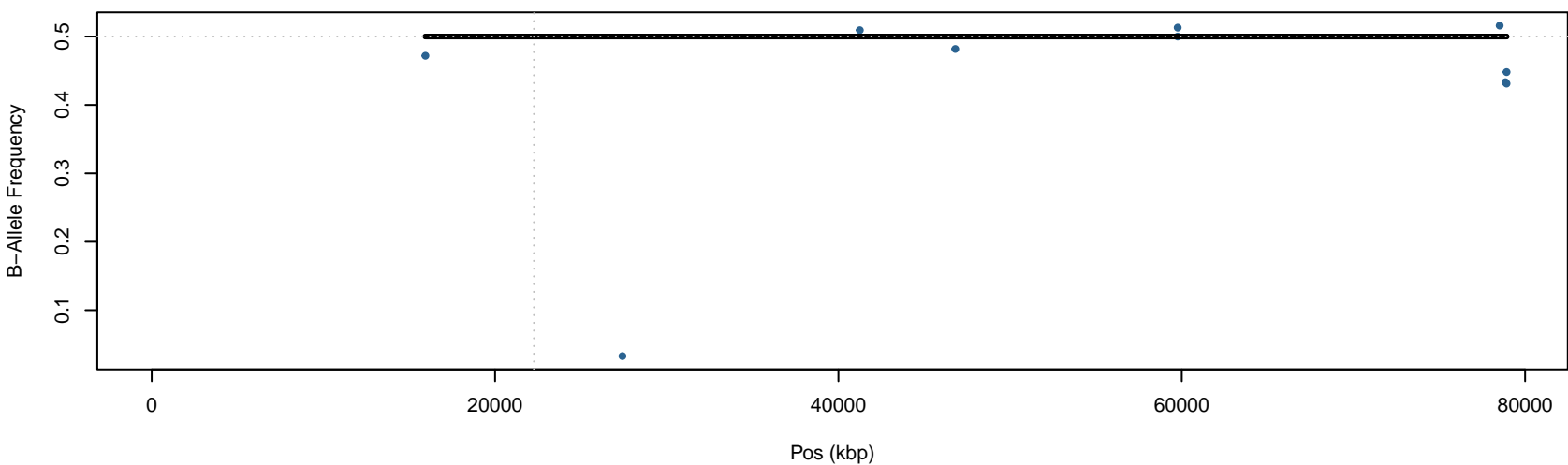
Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 15



Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 16

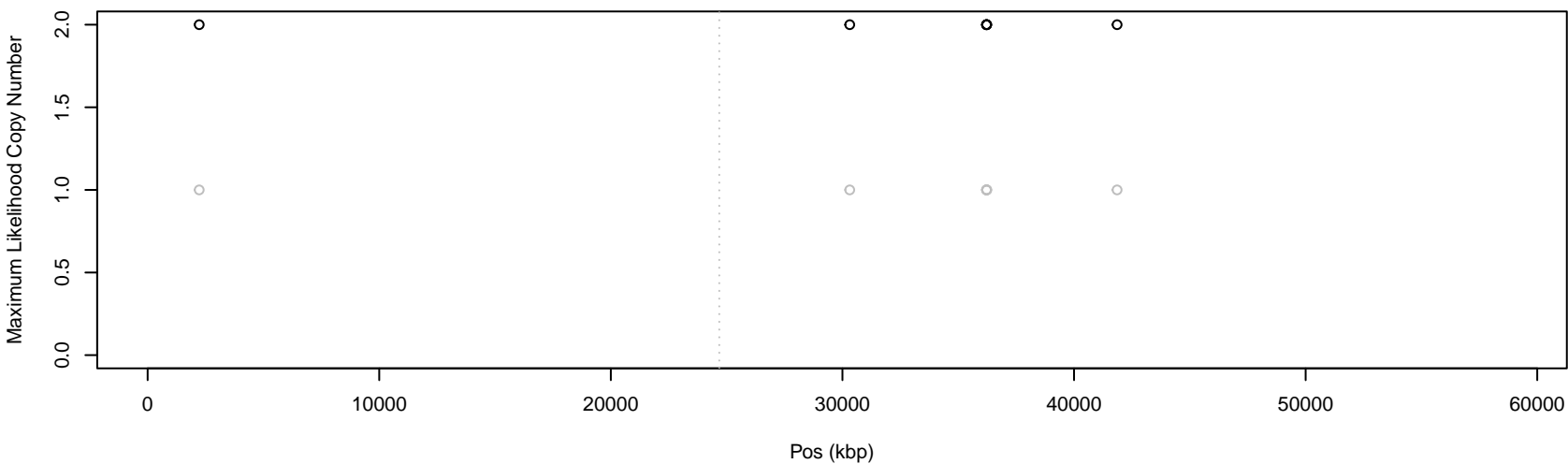
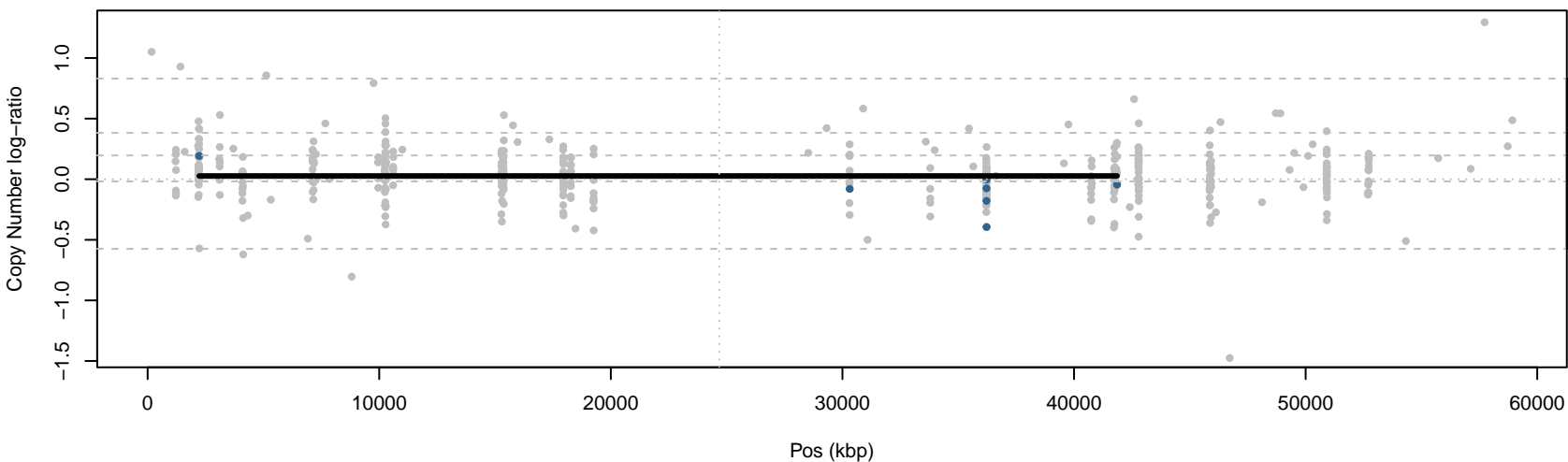
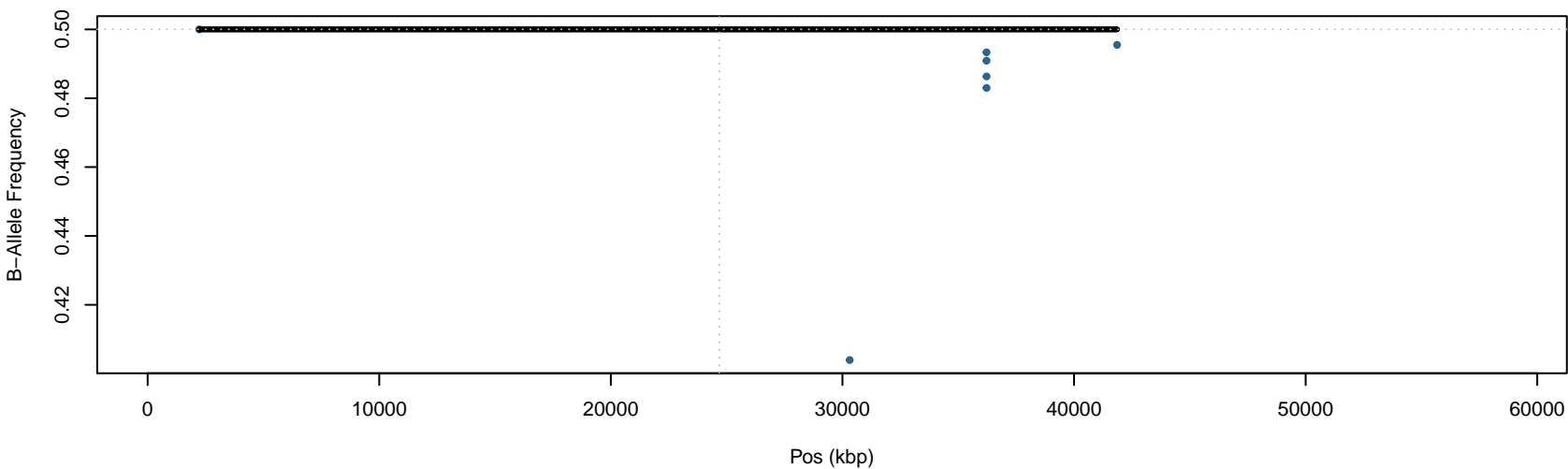


Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 17

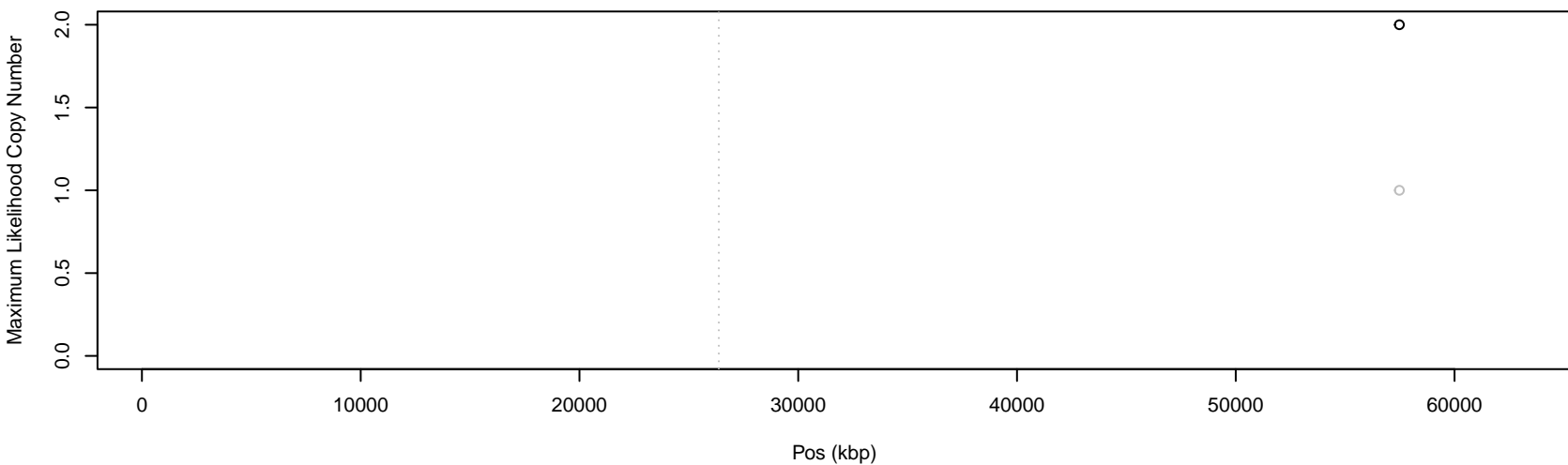
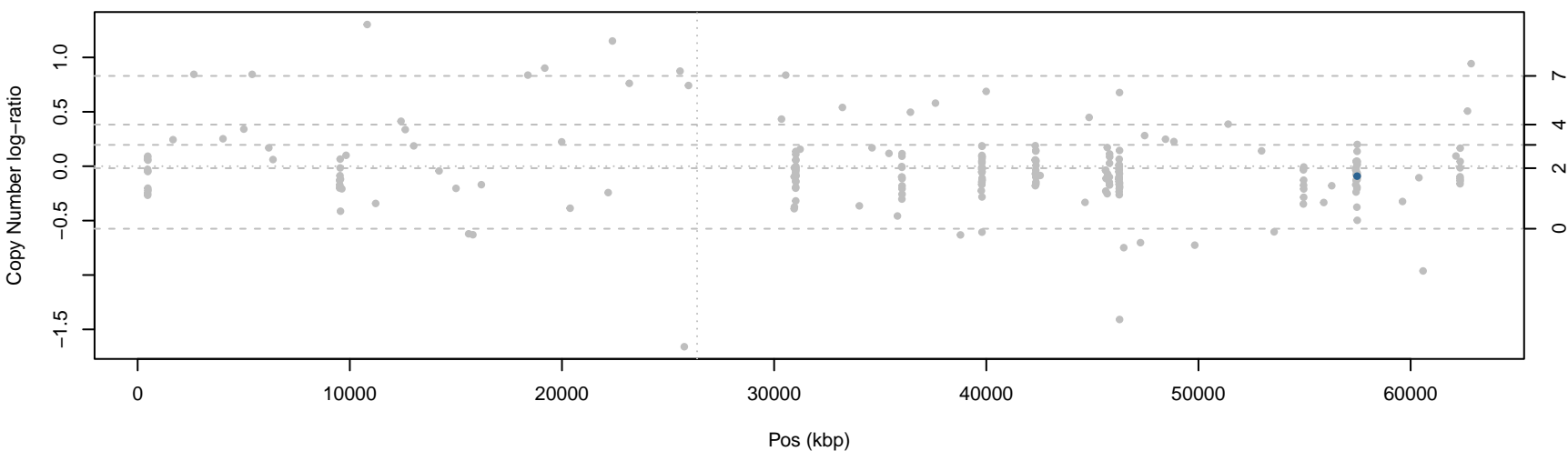
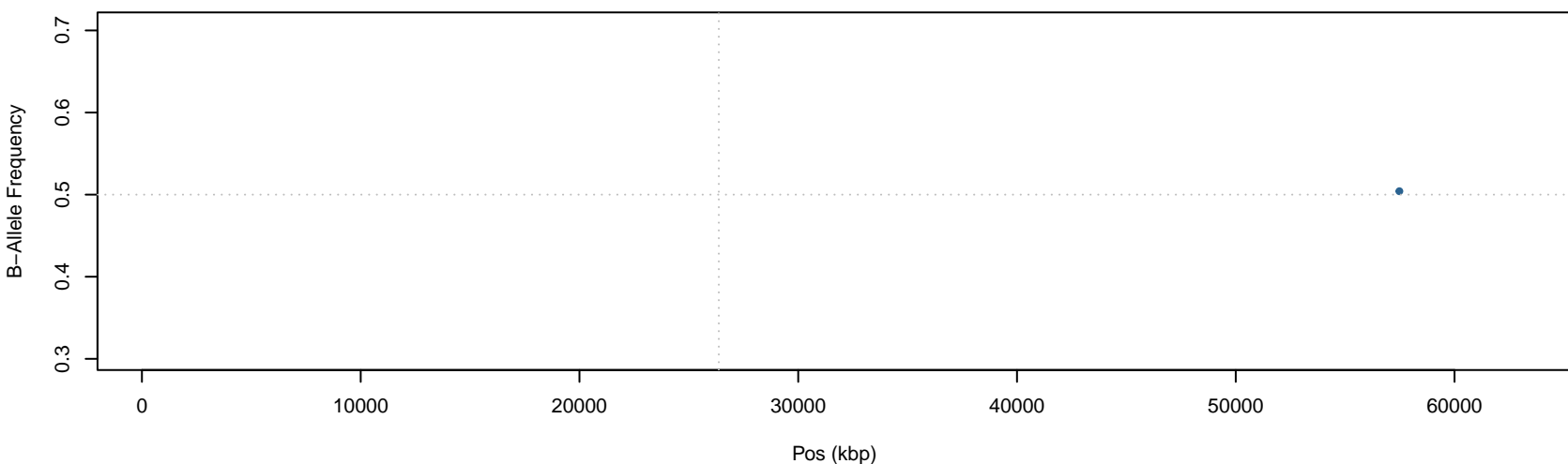




Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 19



Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 20



Purity: 0.32 Tumor ploidy: 2.079 SNV log-likelihood: 70.29 GoF: 88.4% Mean coverage: 399;593 Chromosome: 22

