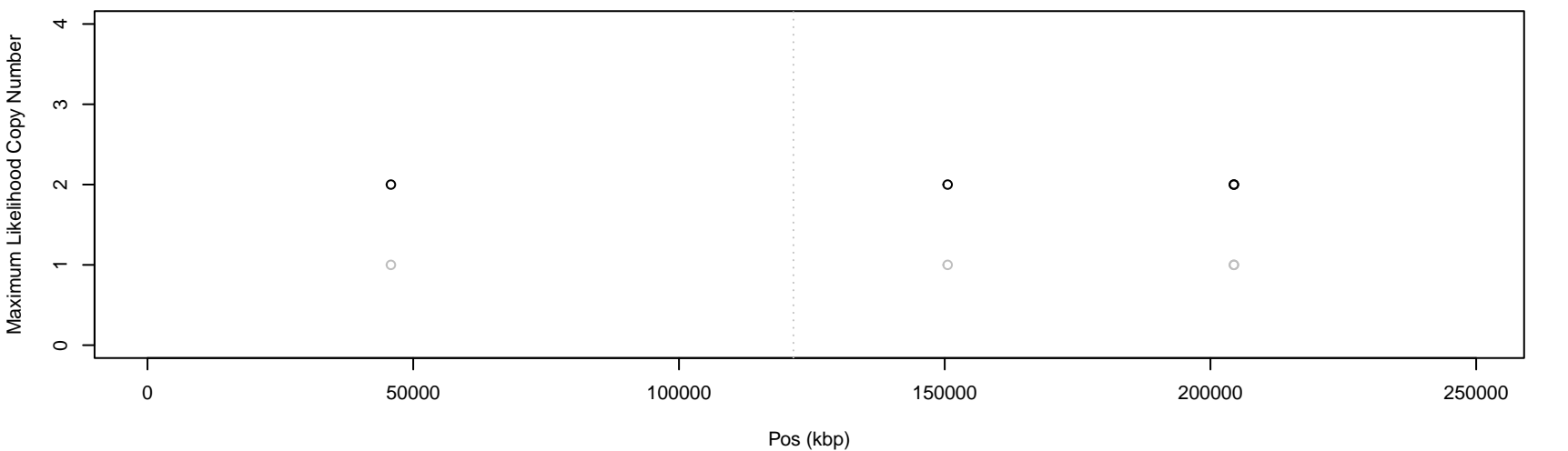
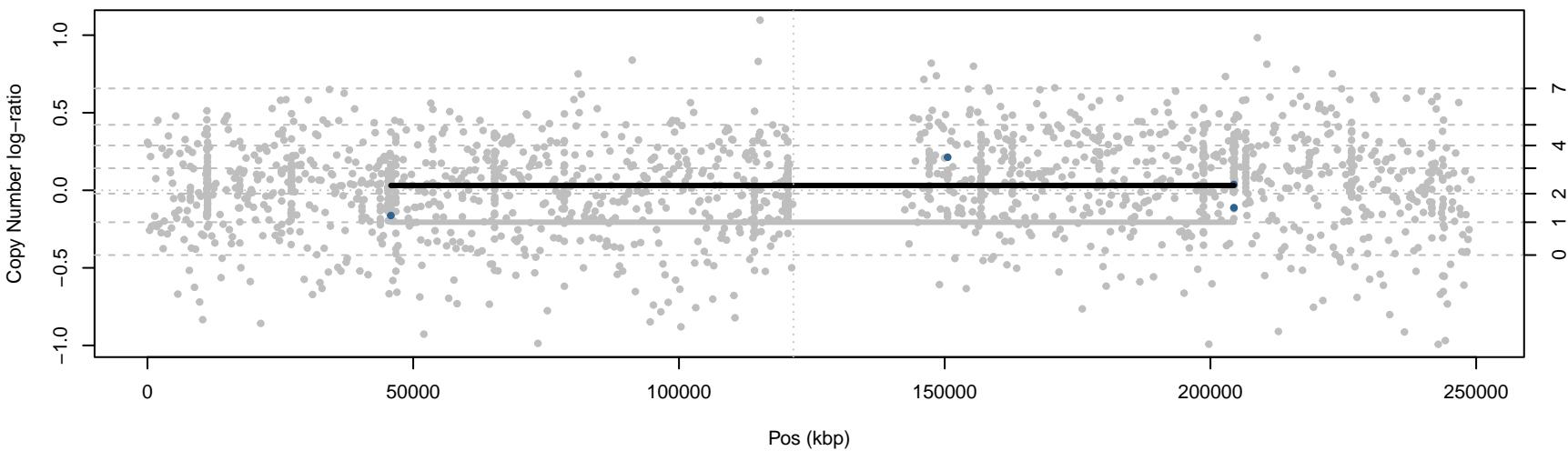
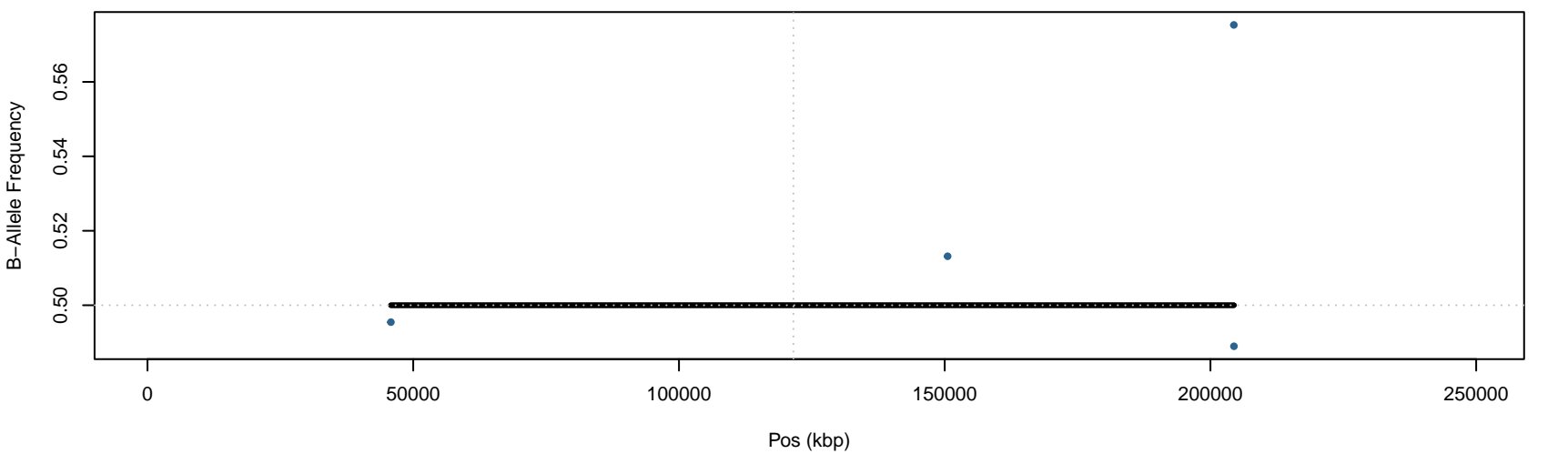
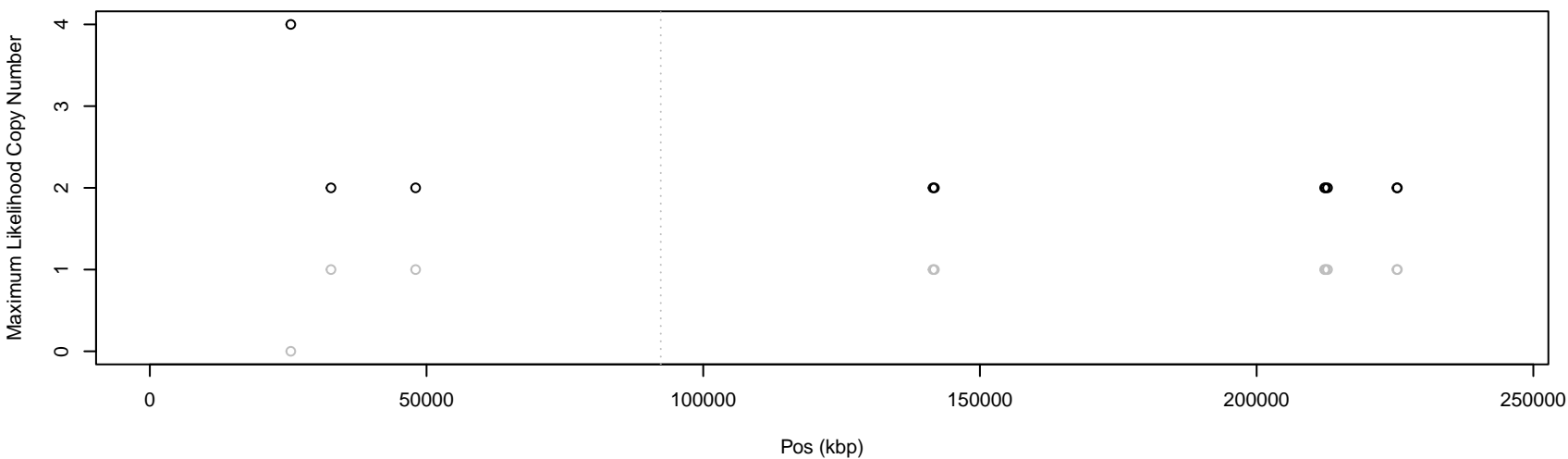
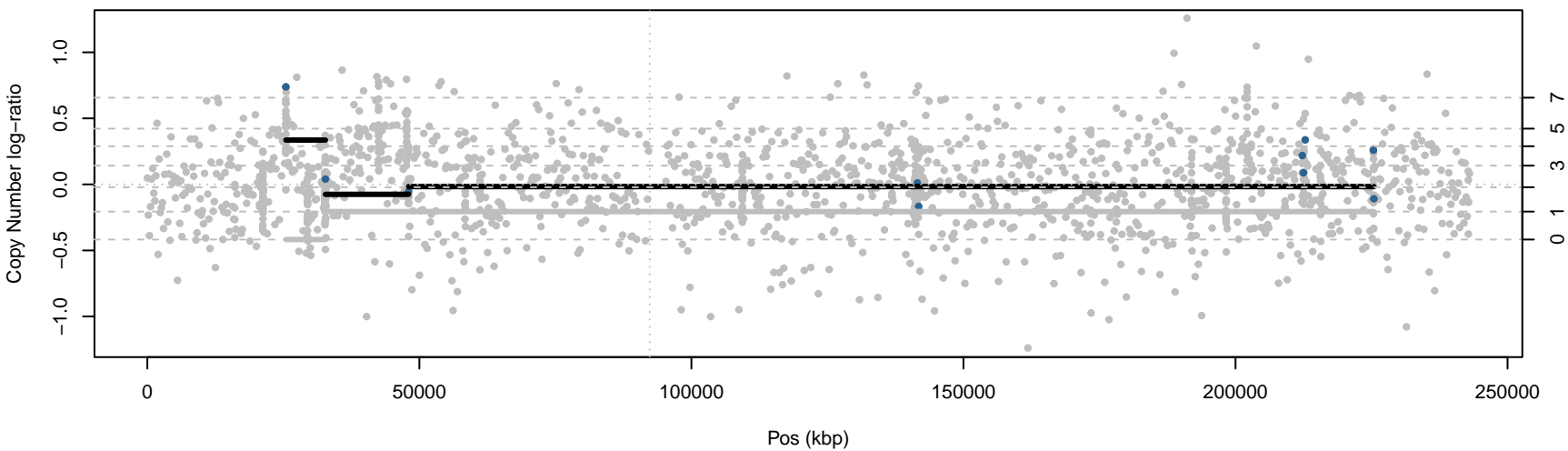
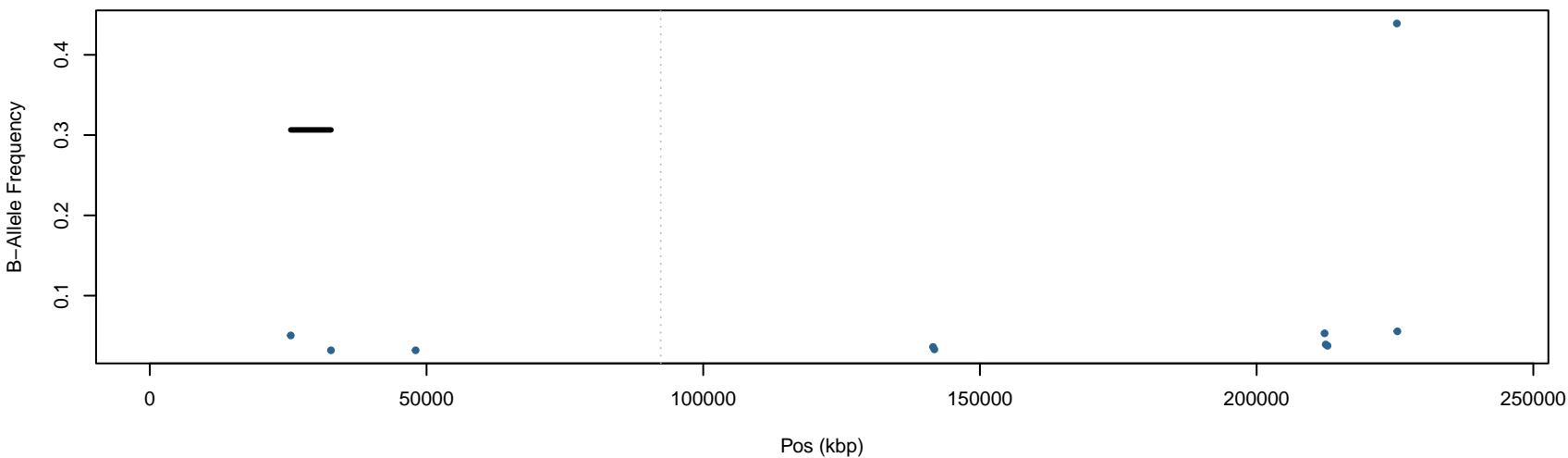


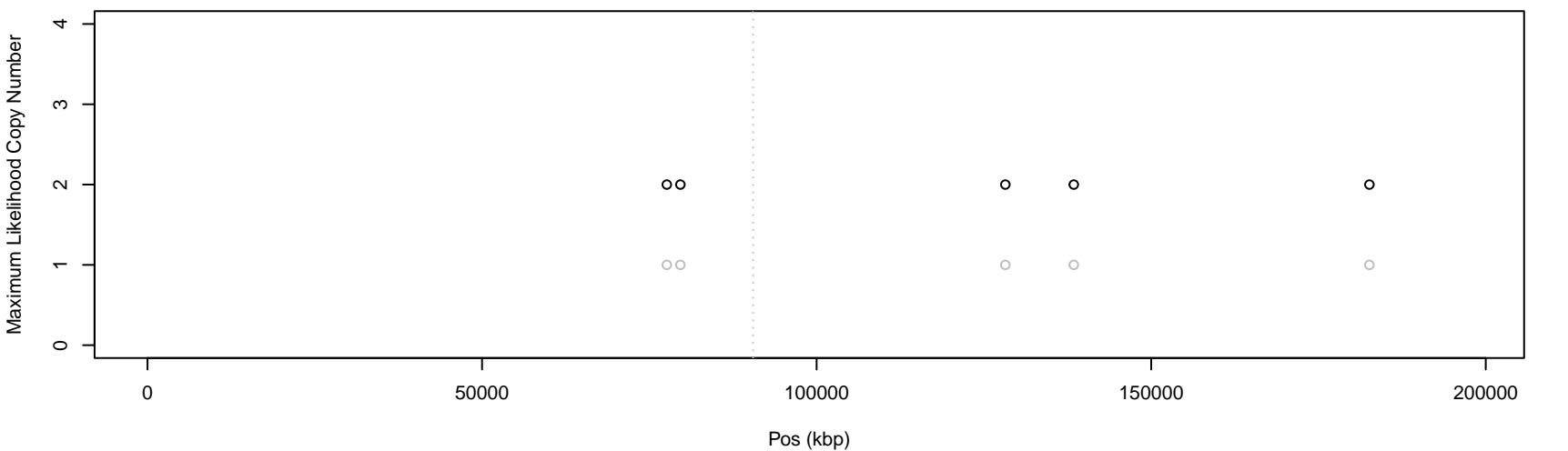
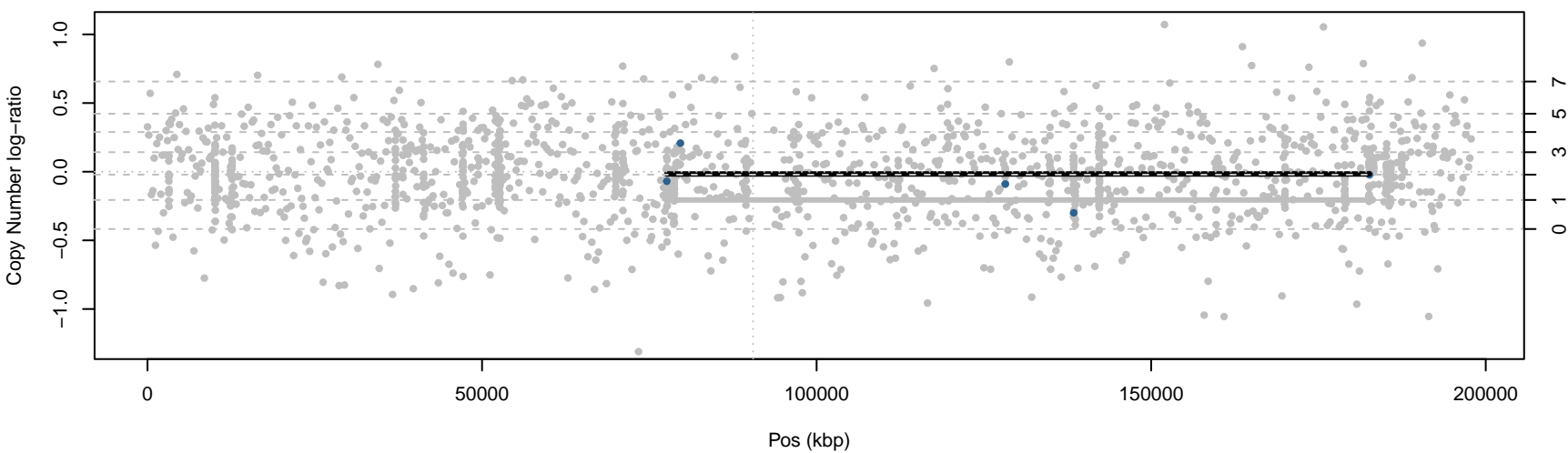
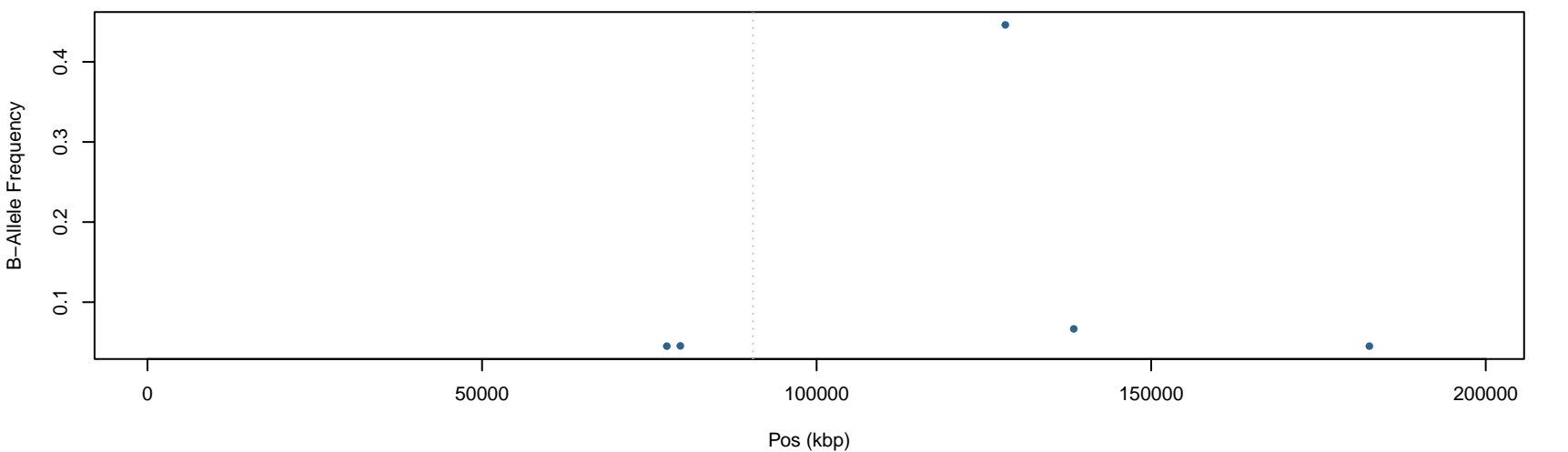
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 1



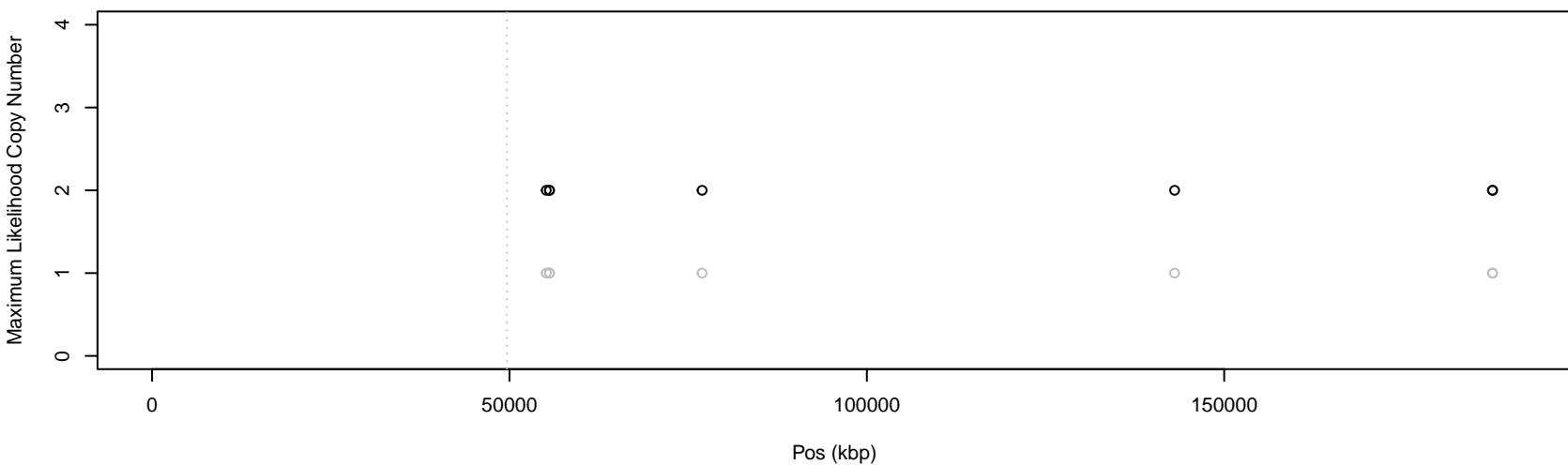
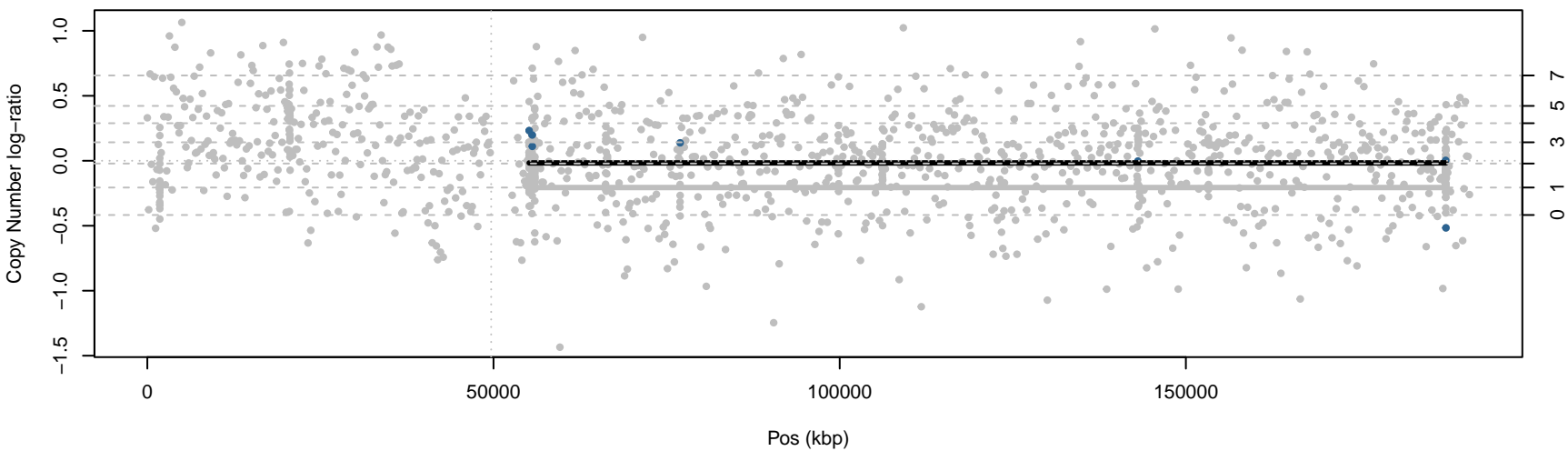
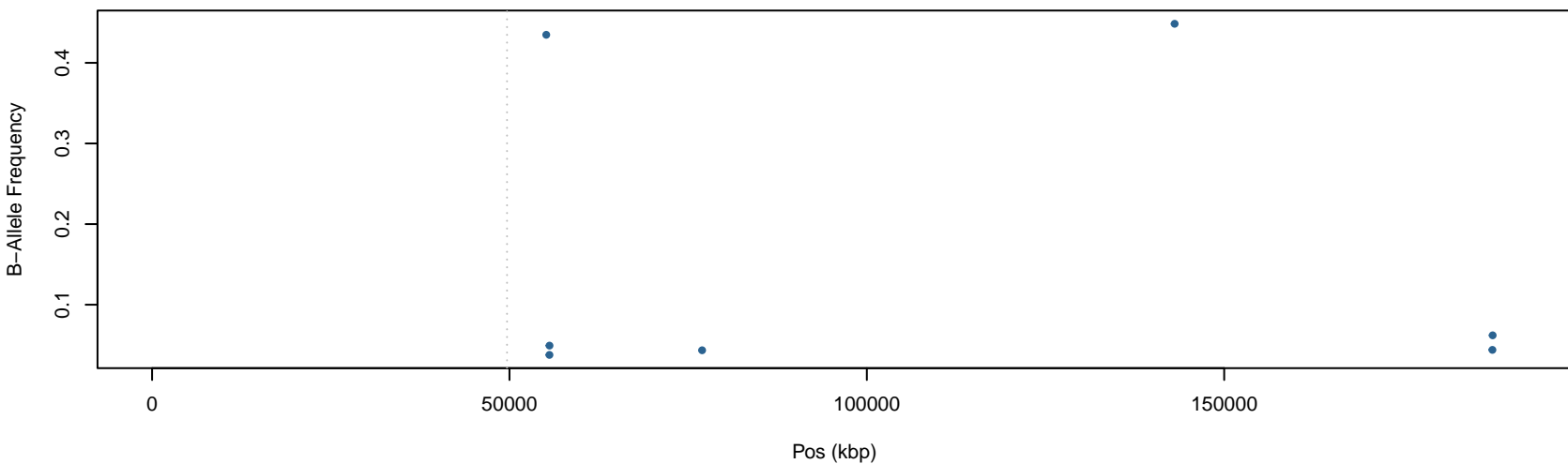
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 2



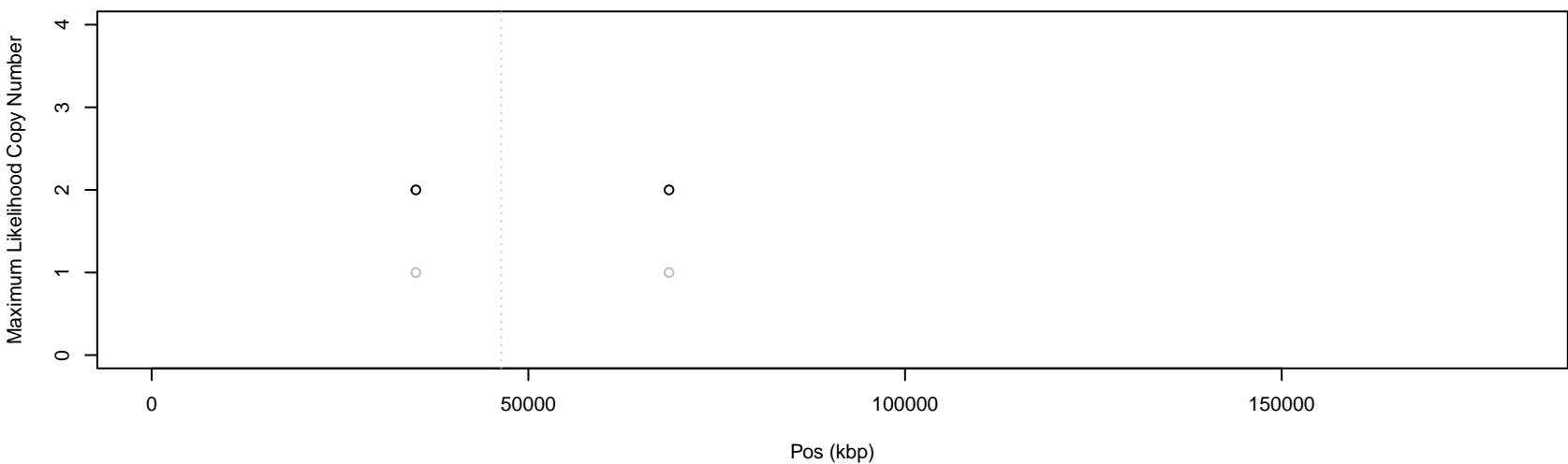
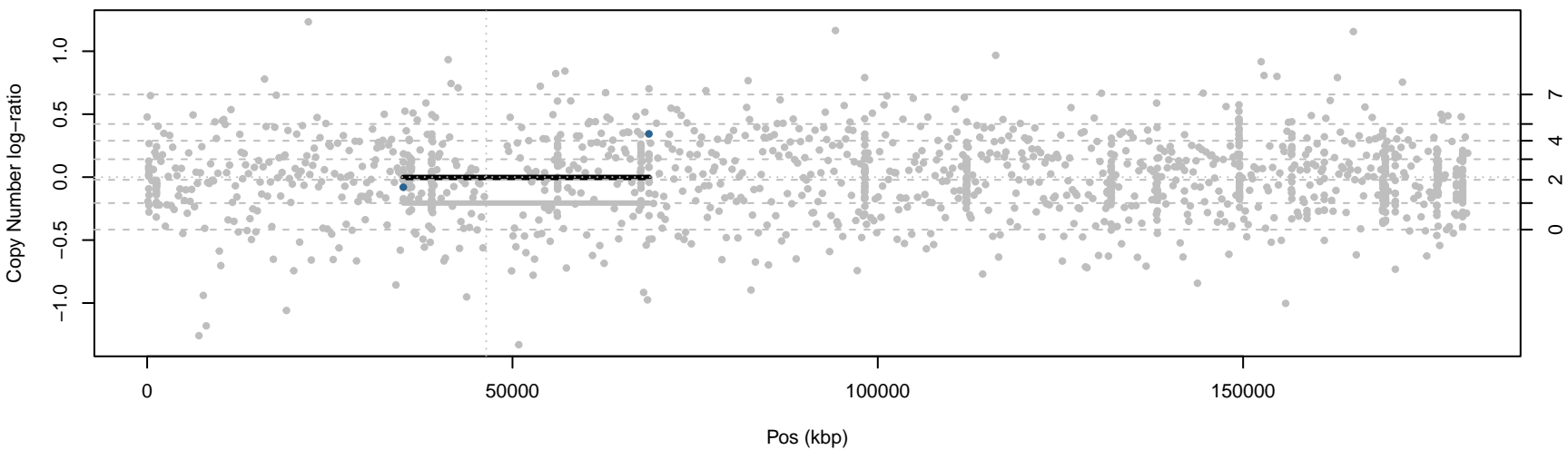
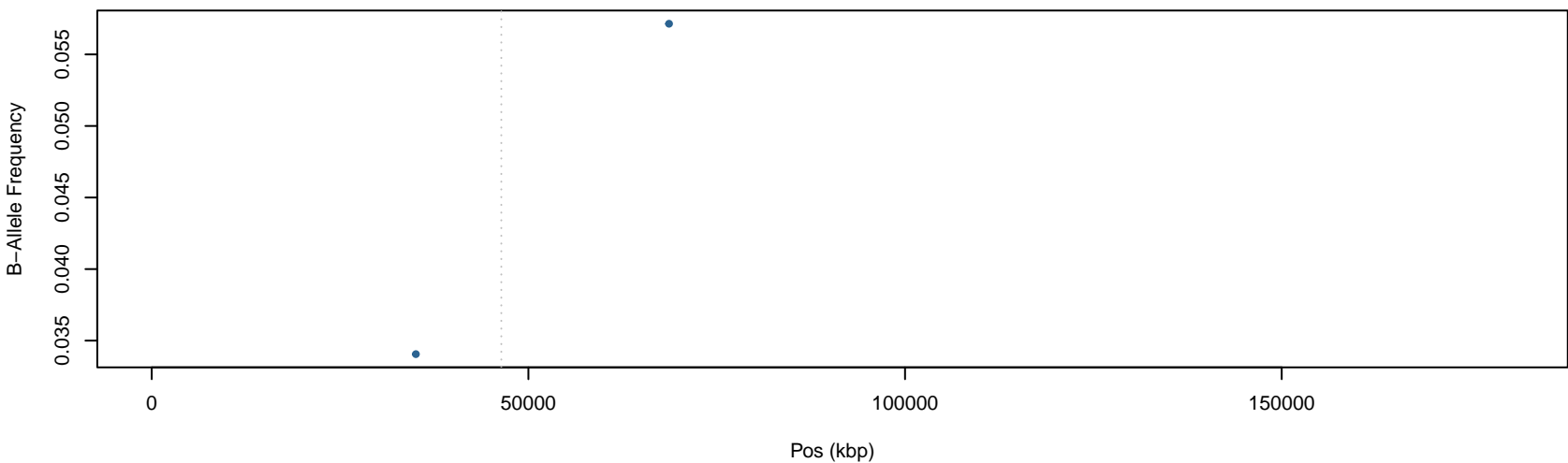
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 3



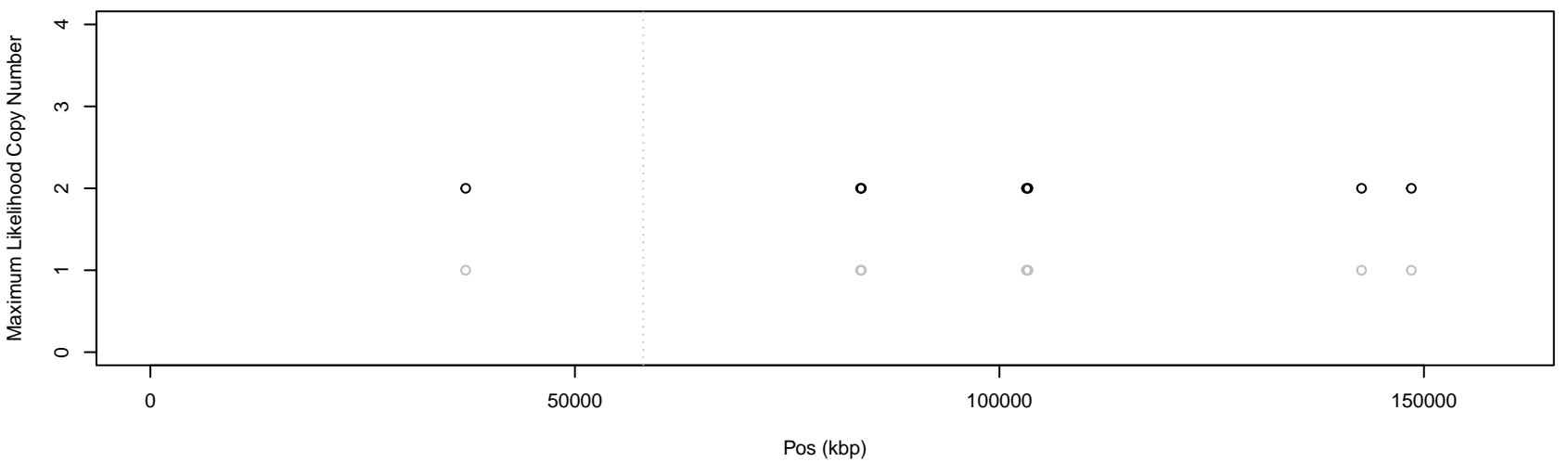
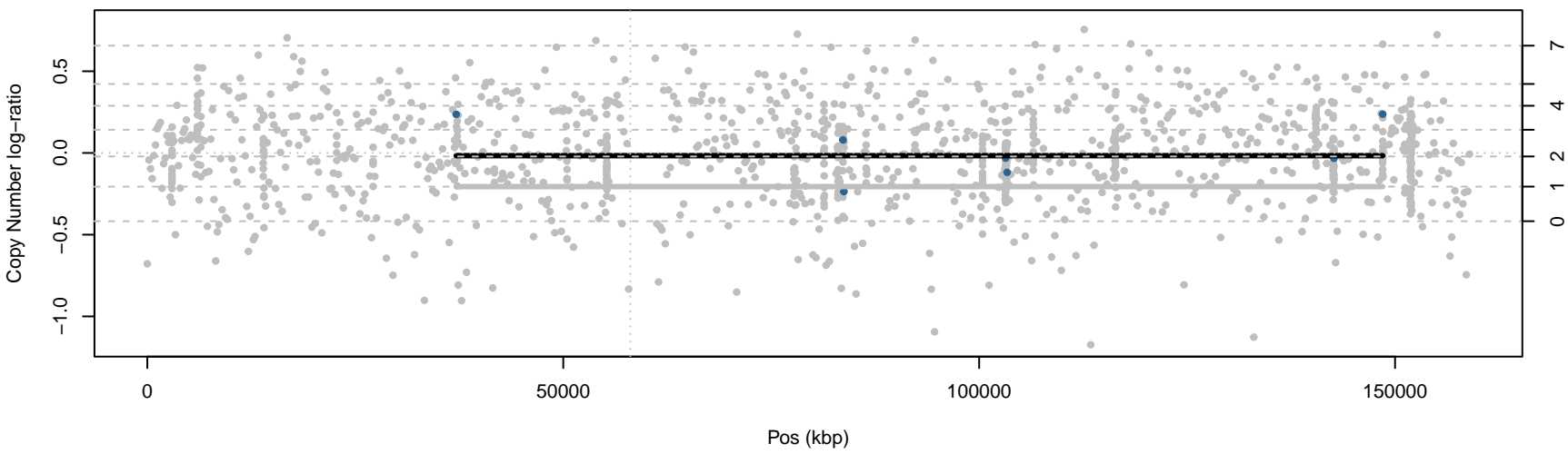
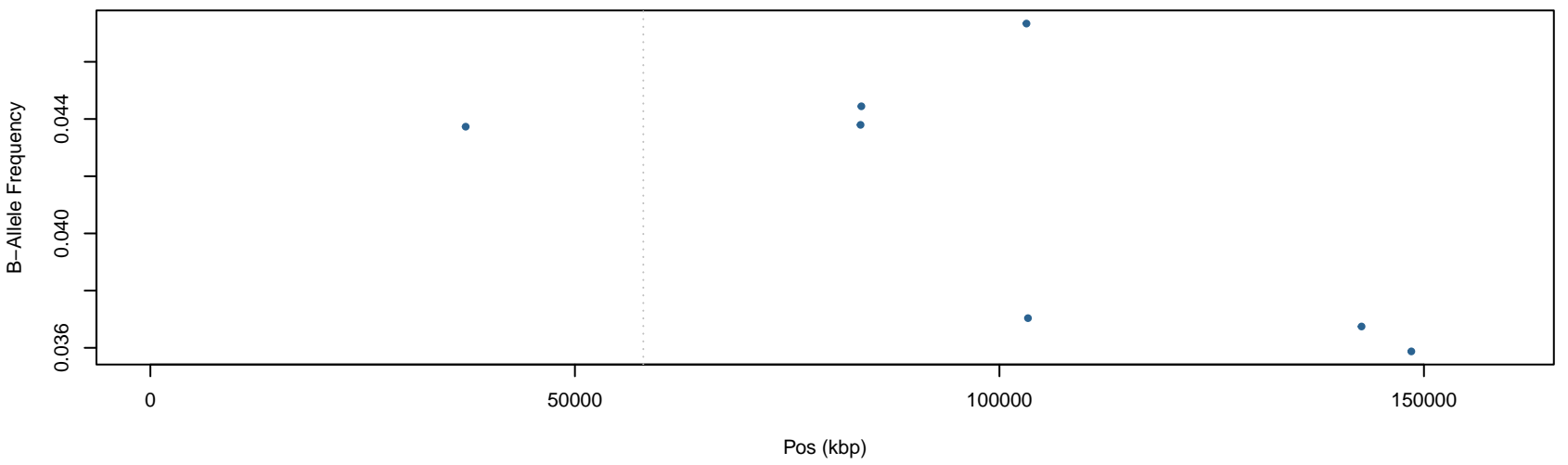
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 4



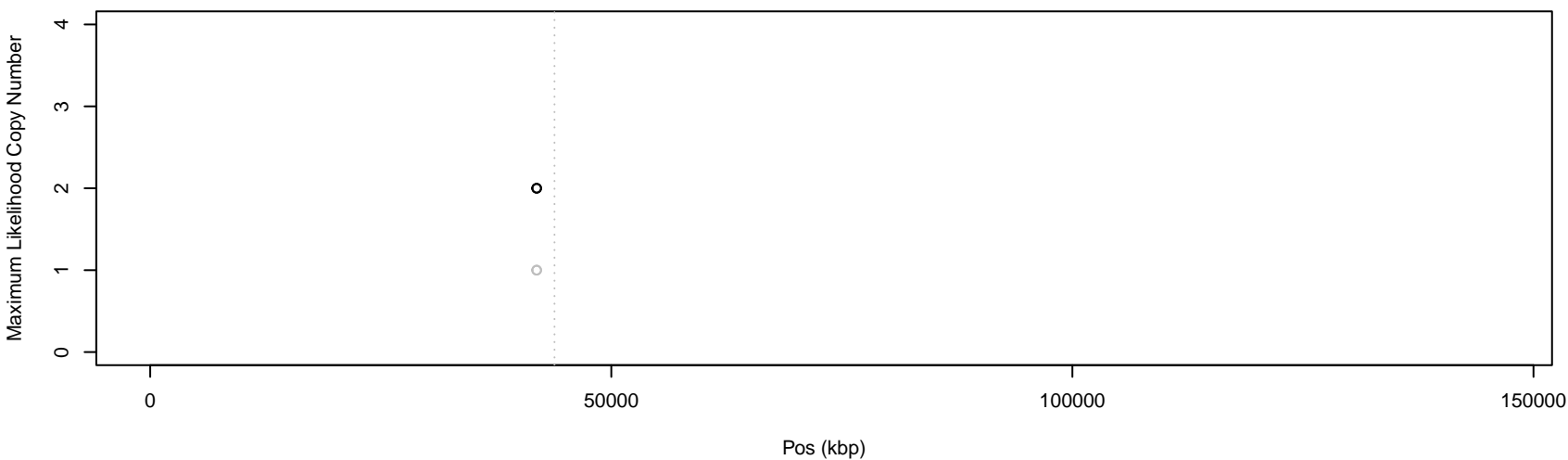
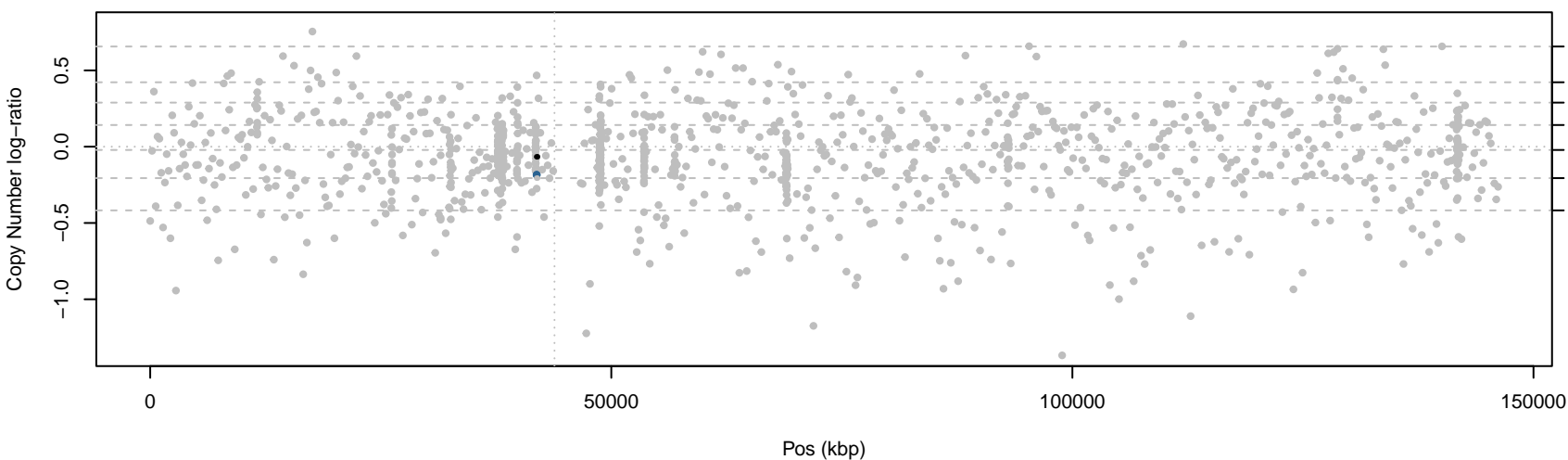
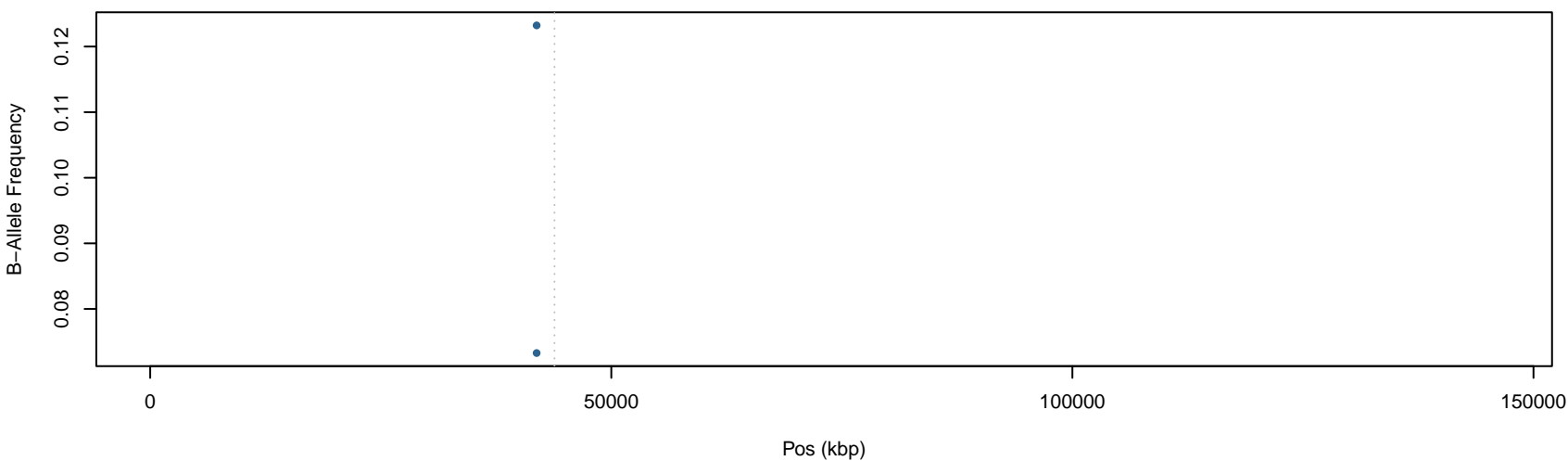
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 5



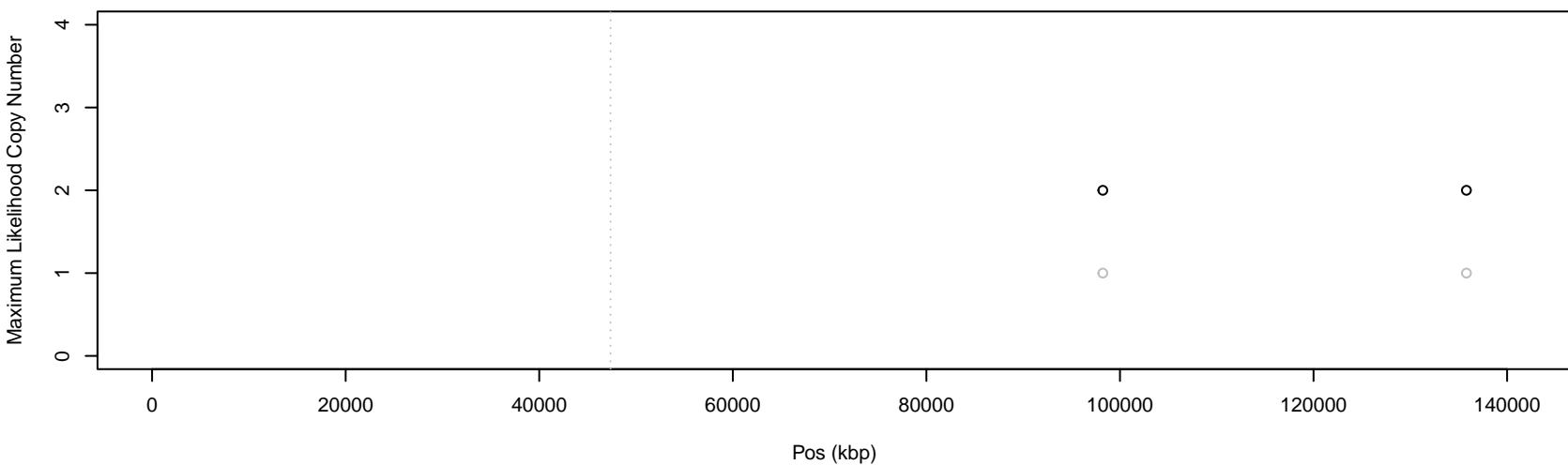
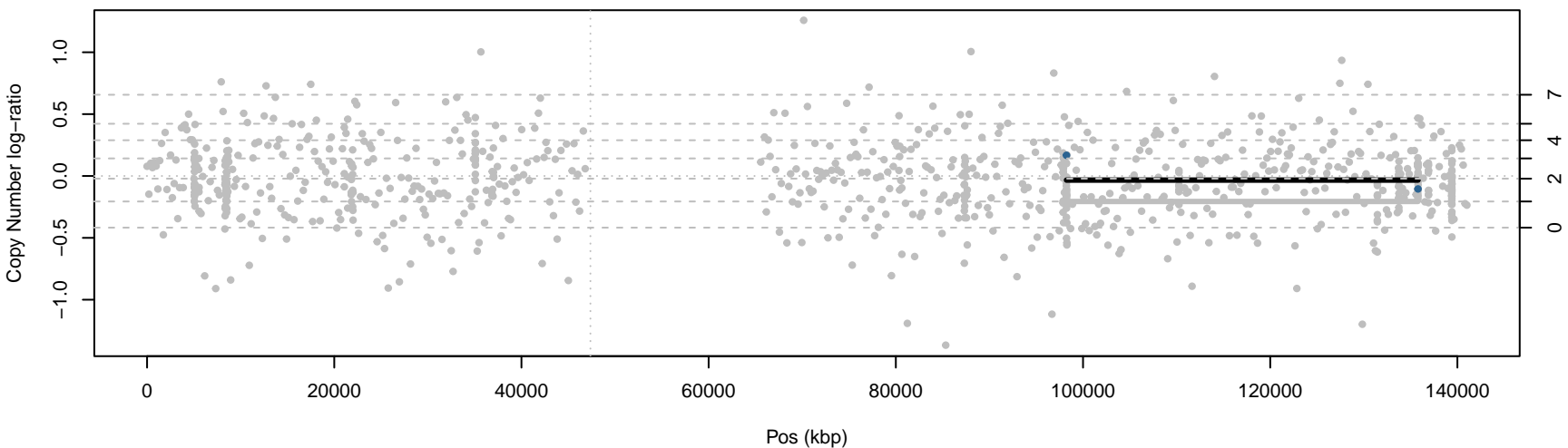
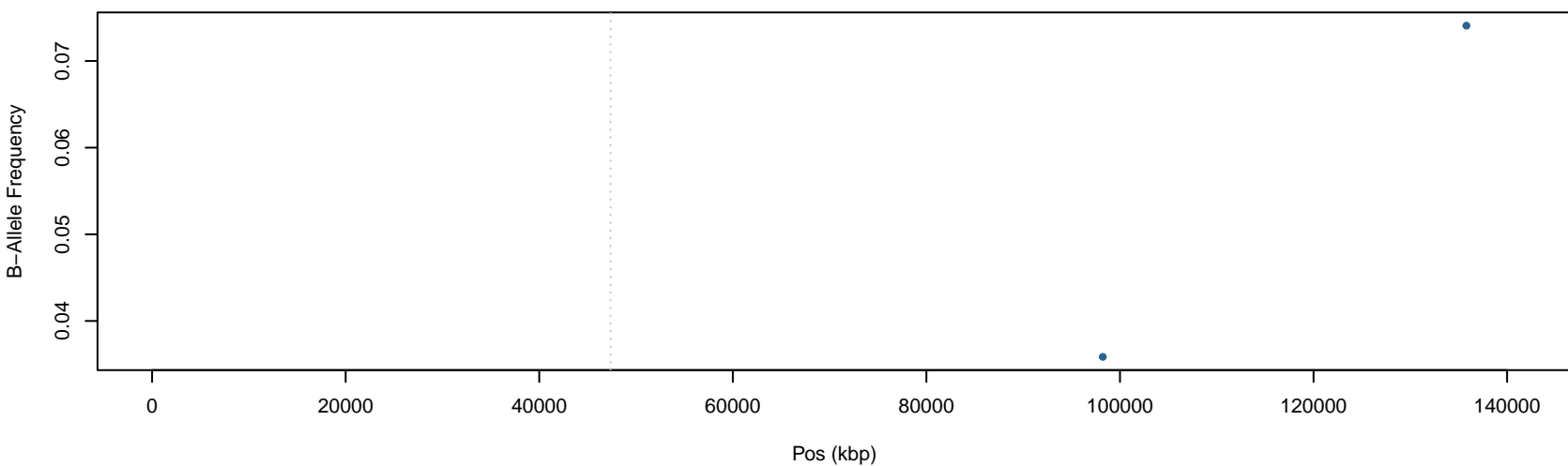
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 7



Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 8

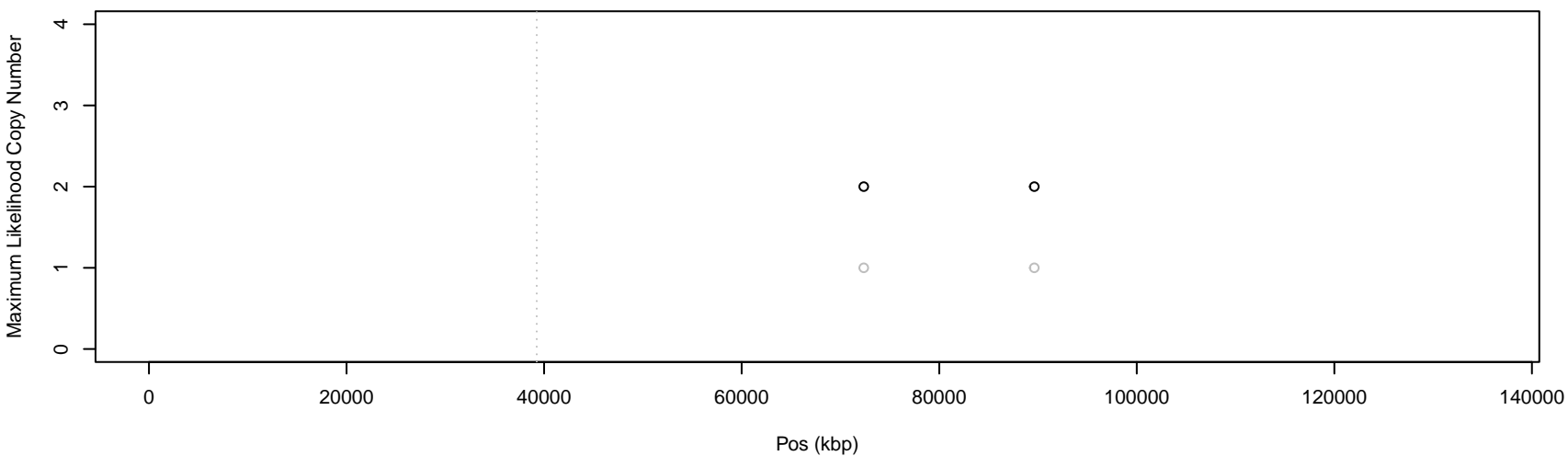
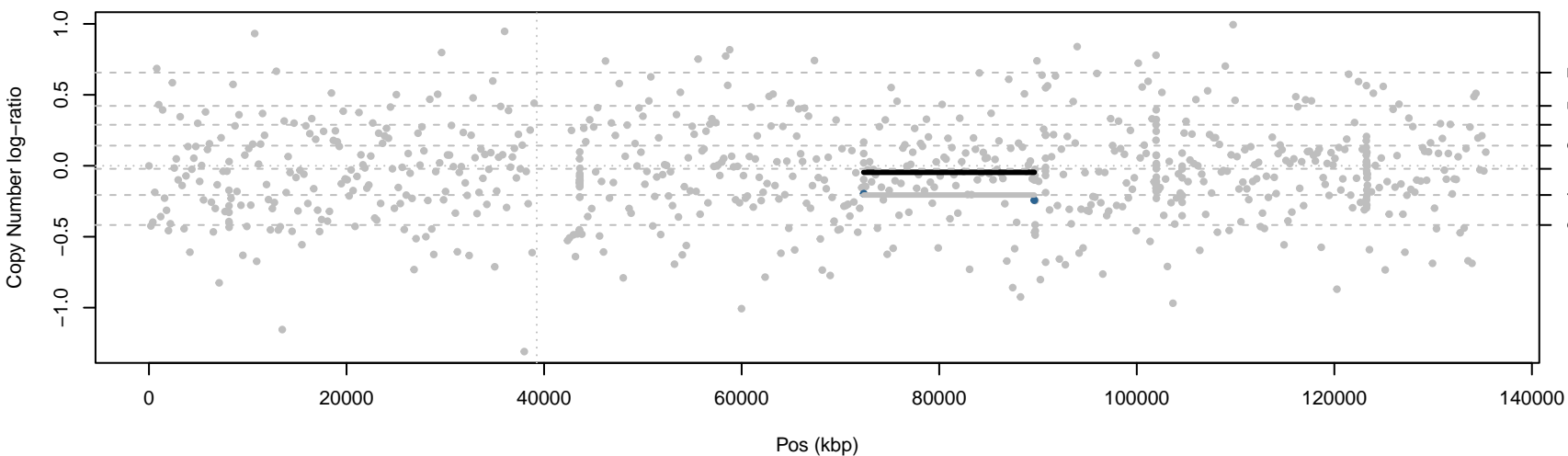
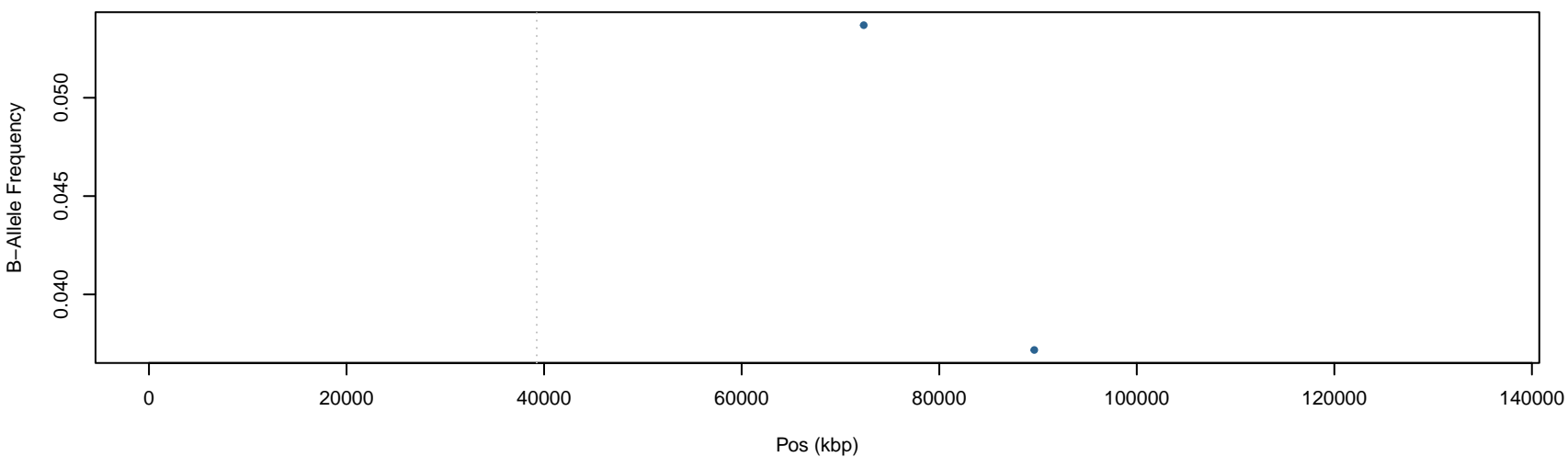


Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 9

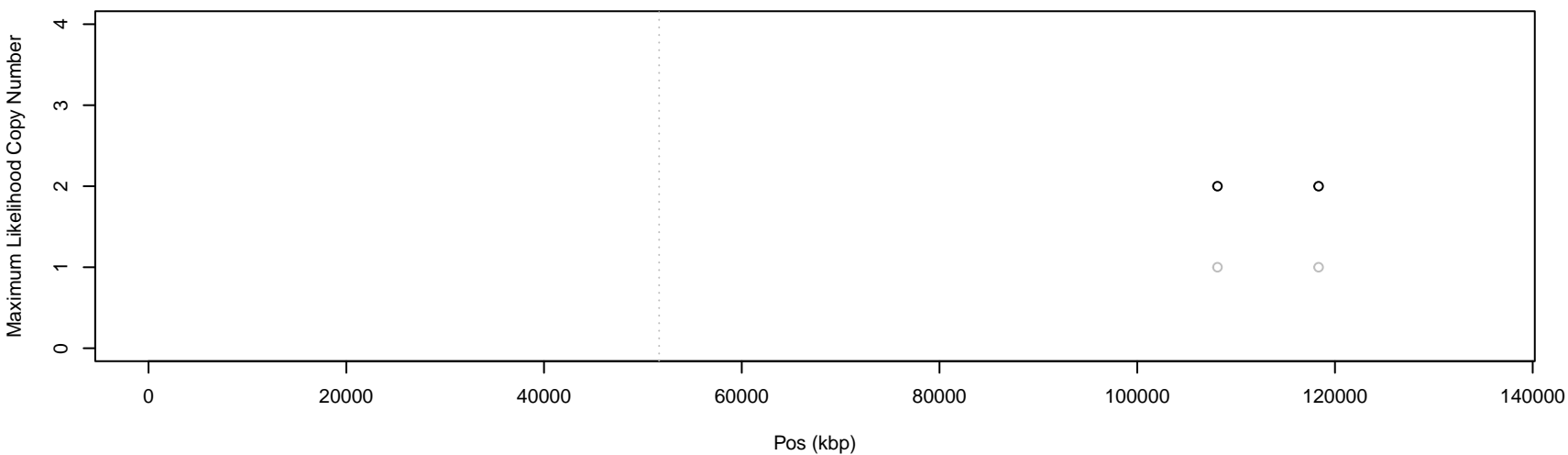
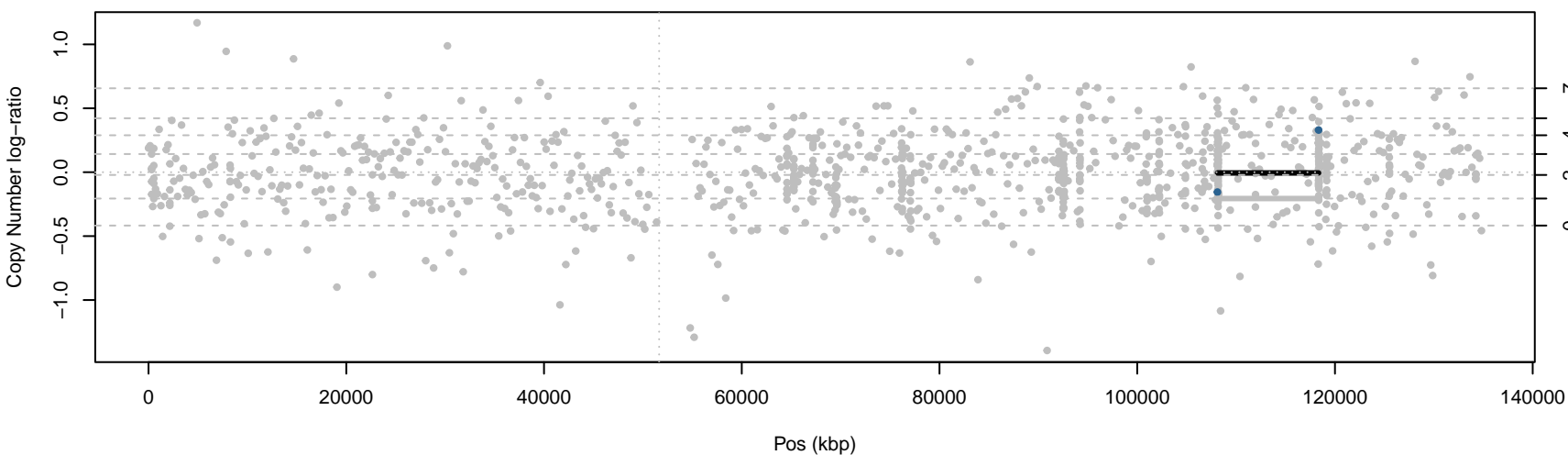
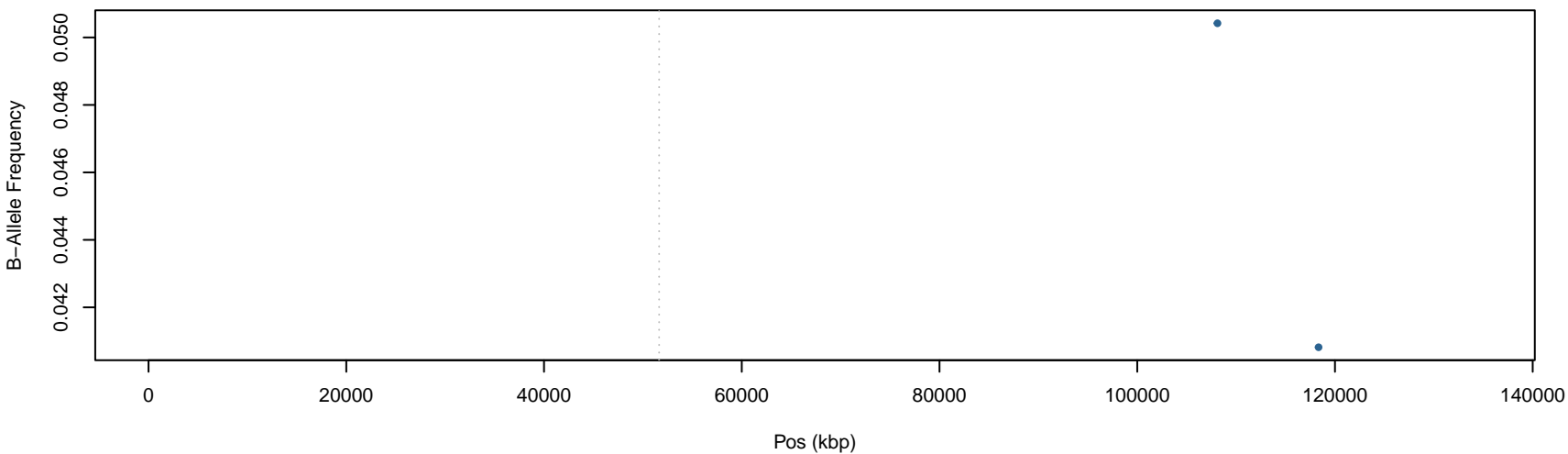




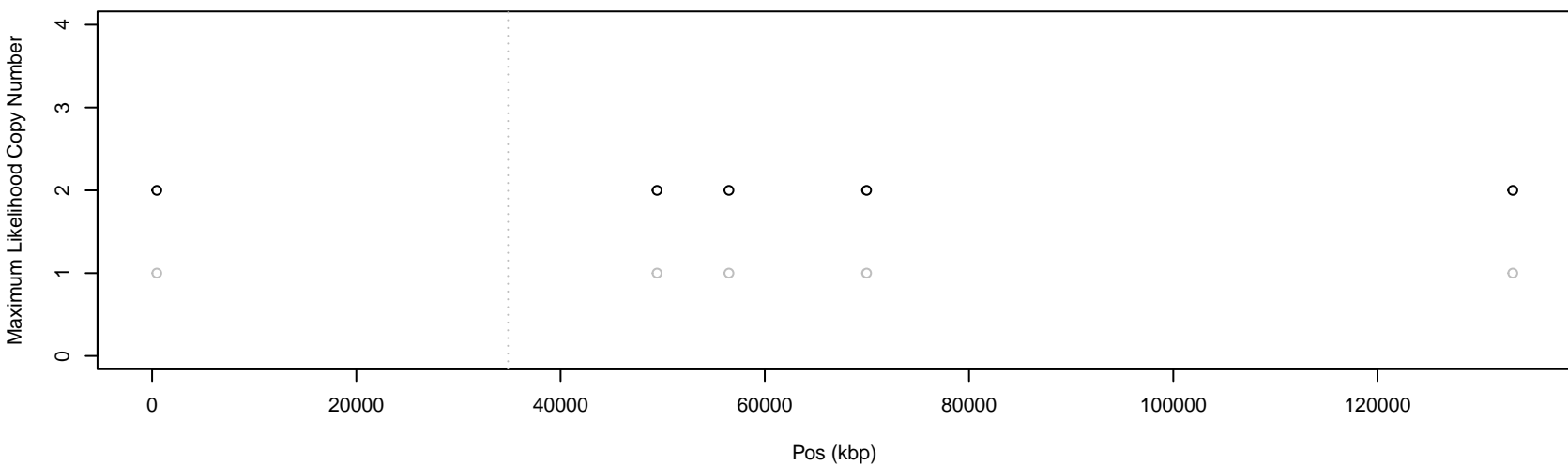
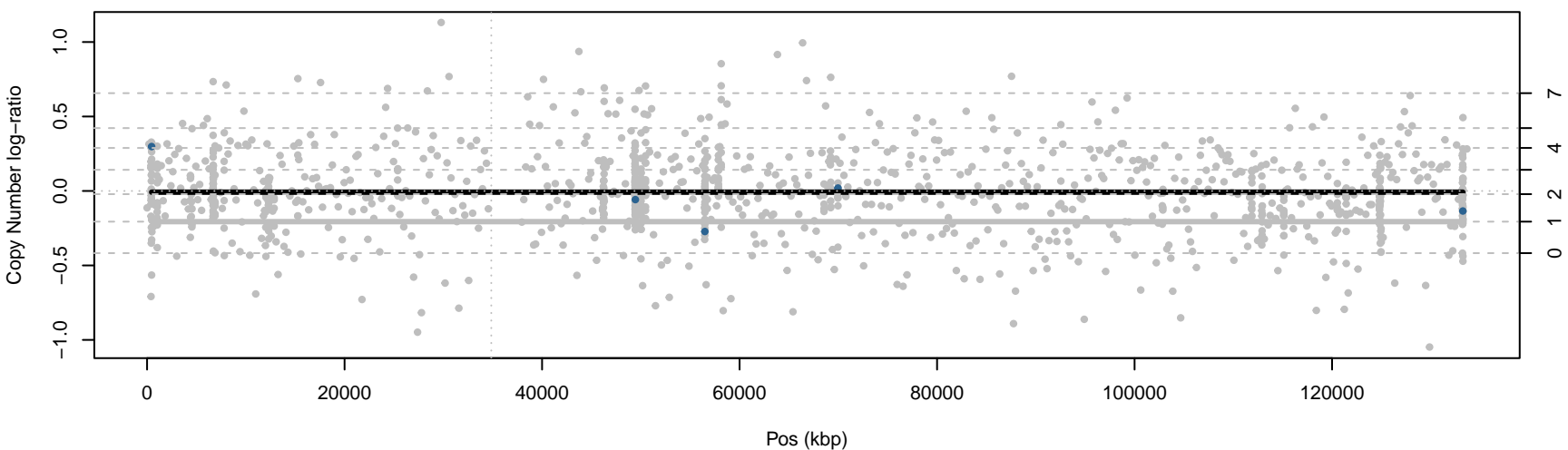
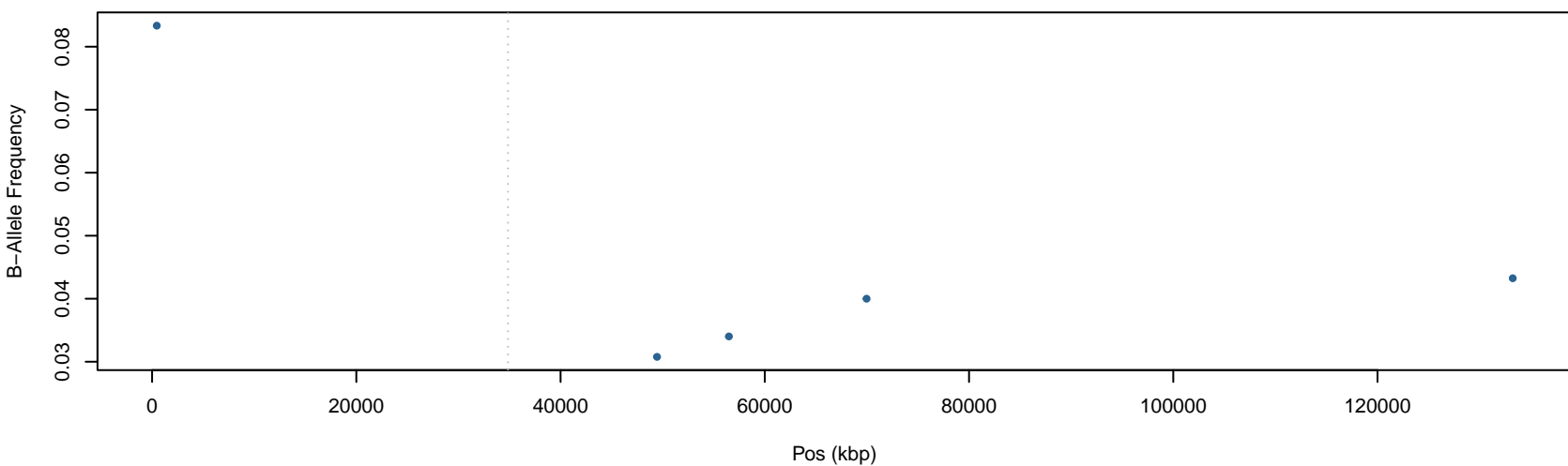
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 10



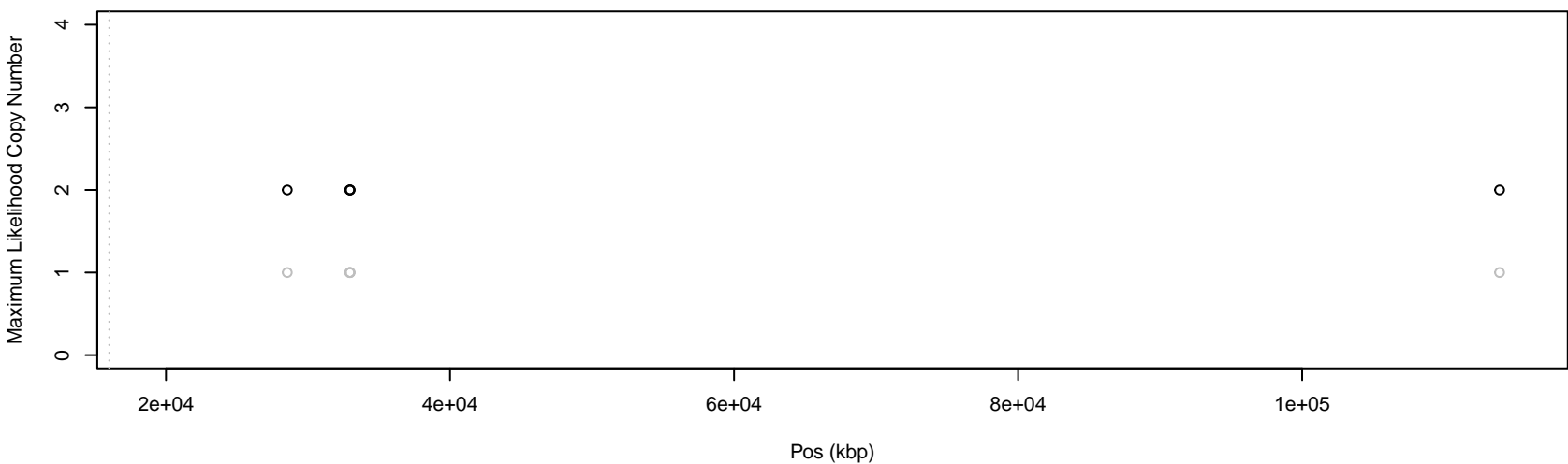
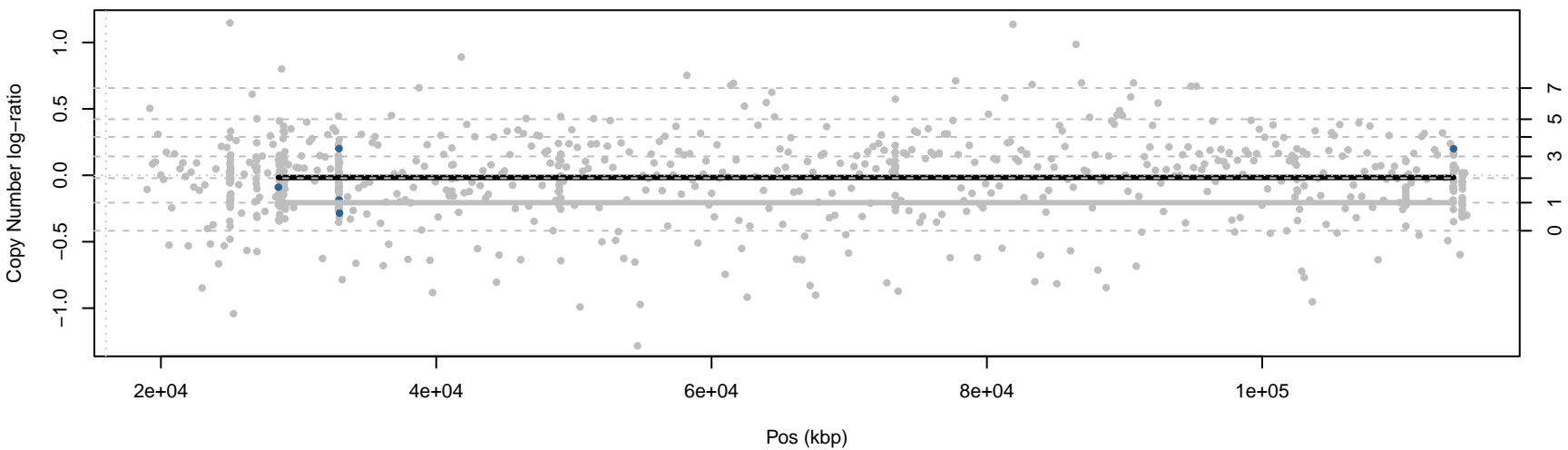
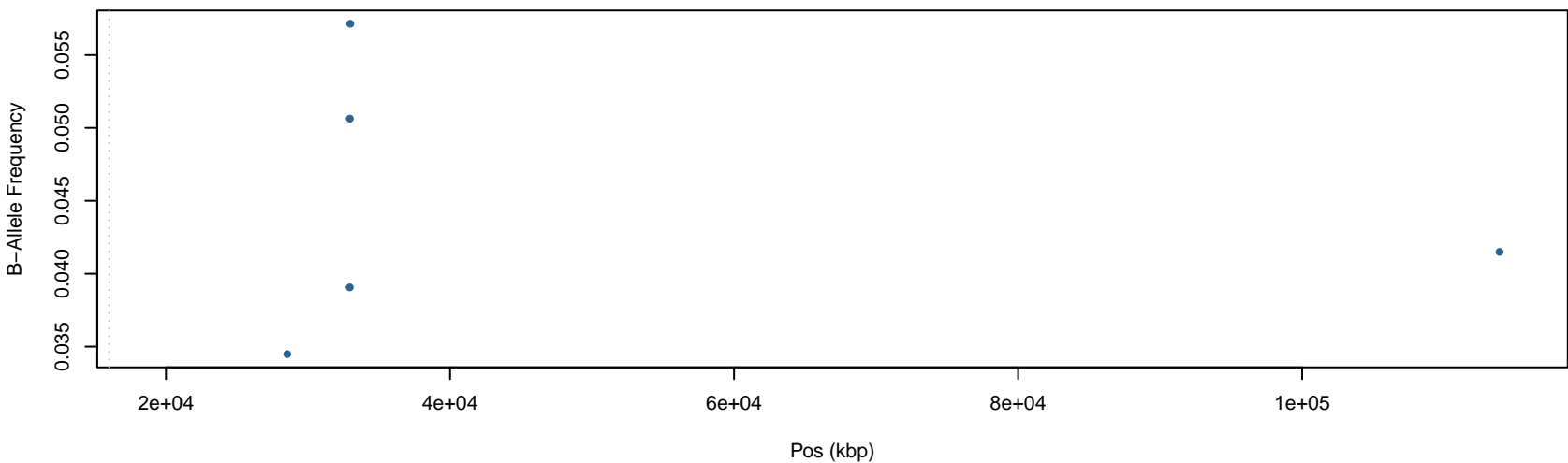
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 11



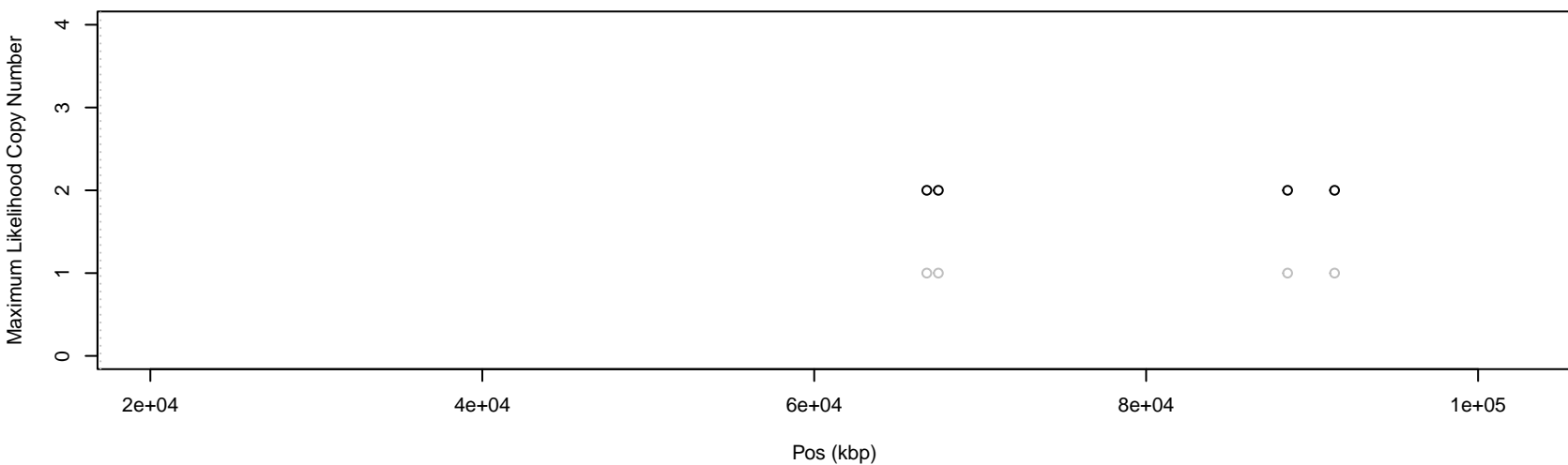
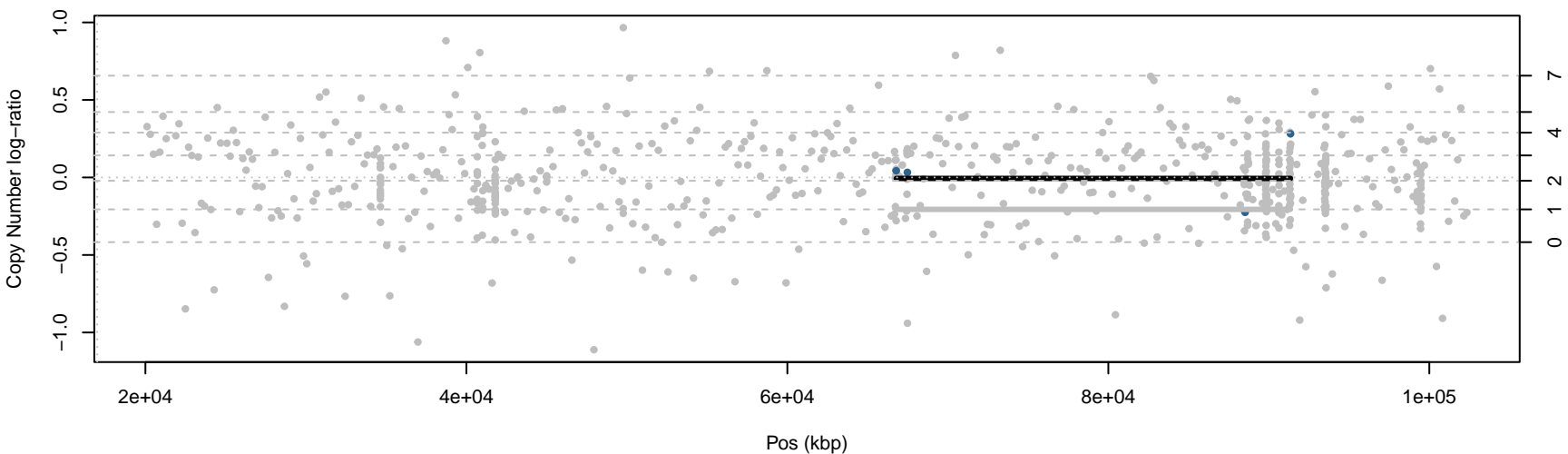
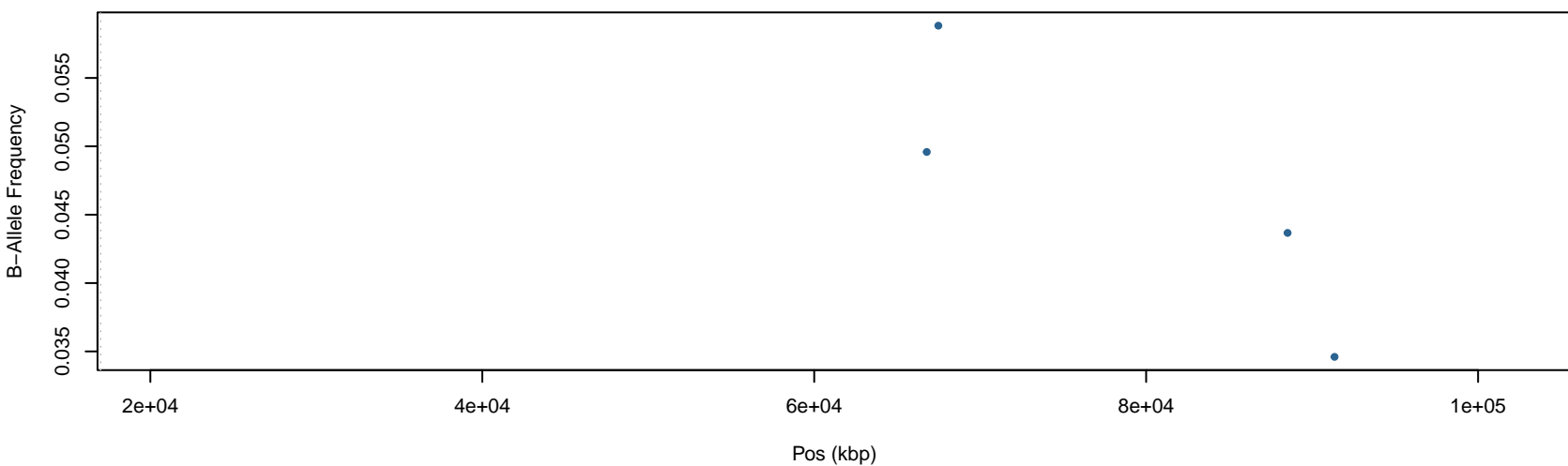
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 12



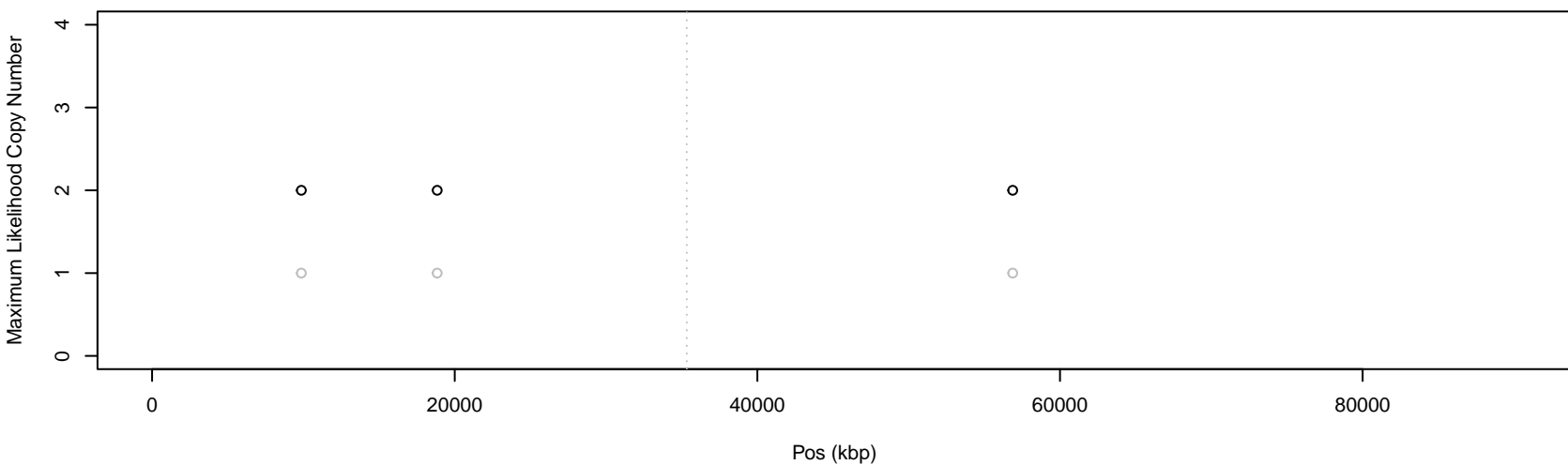
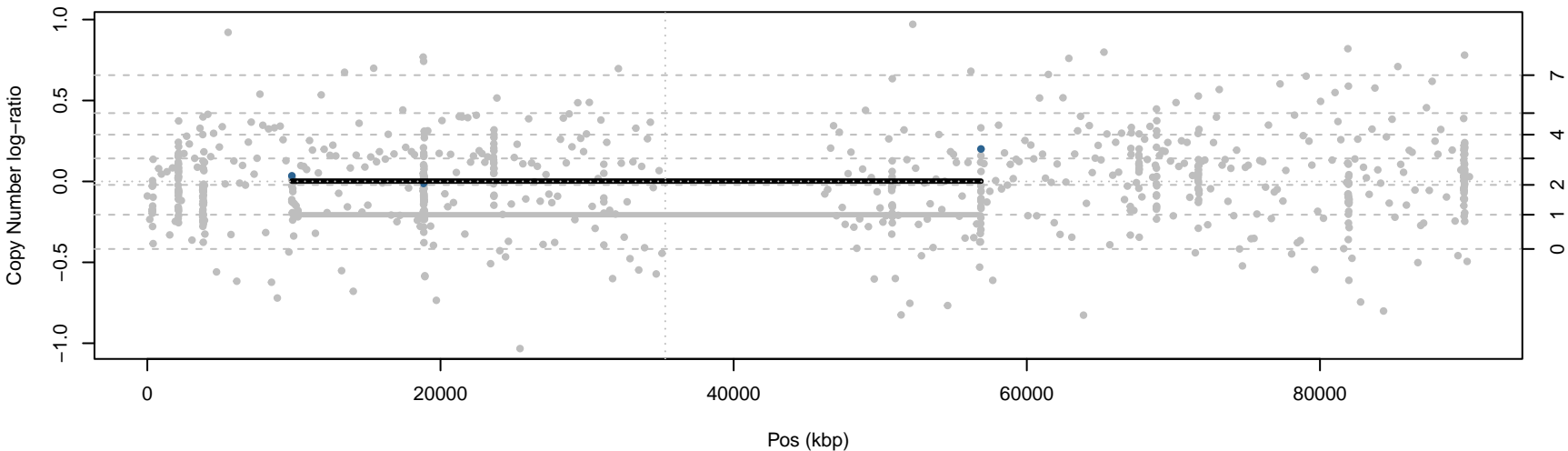
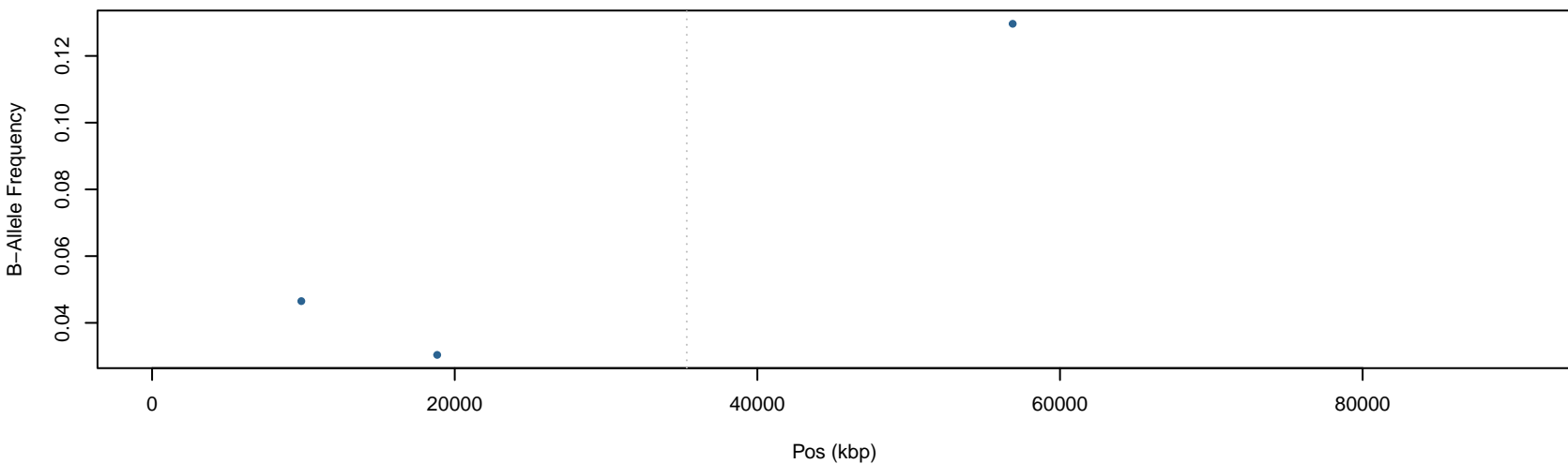
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 13



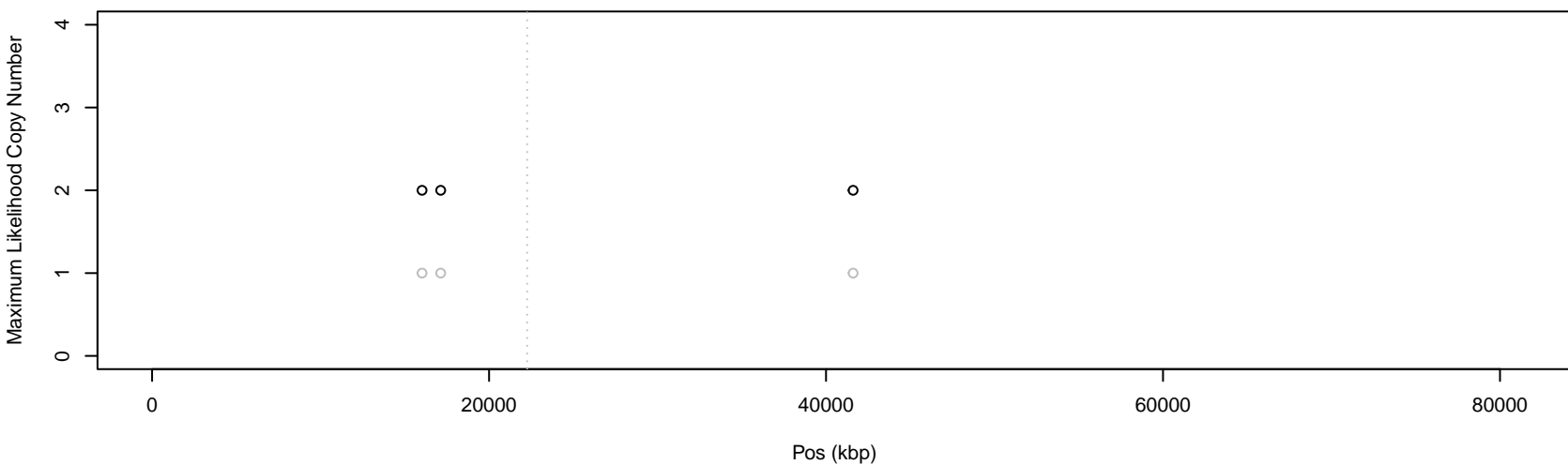
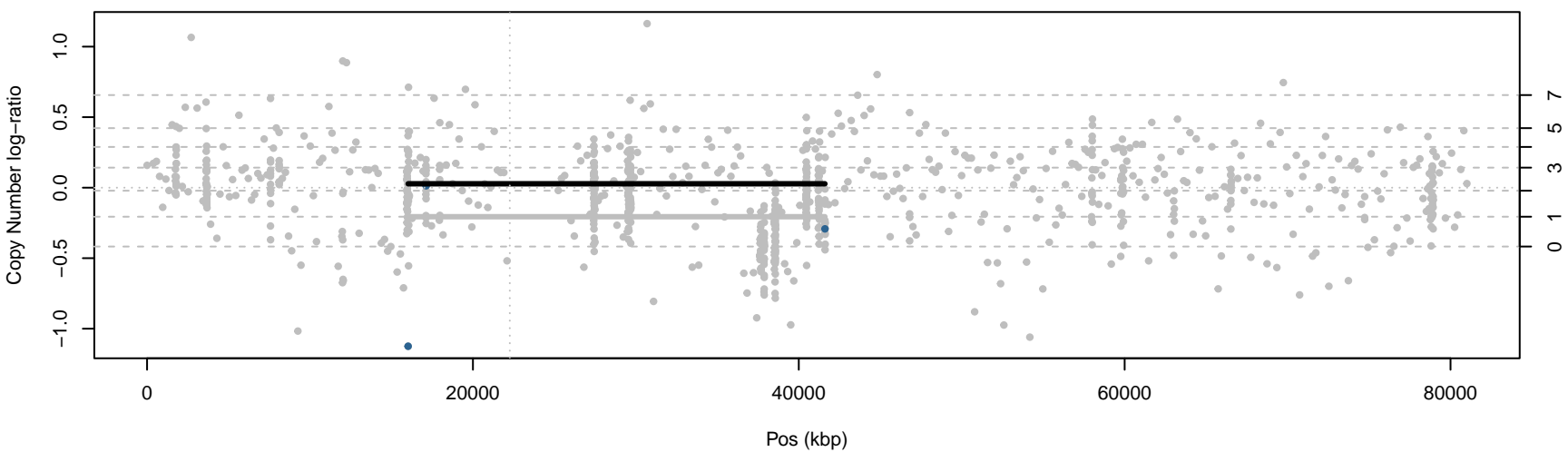
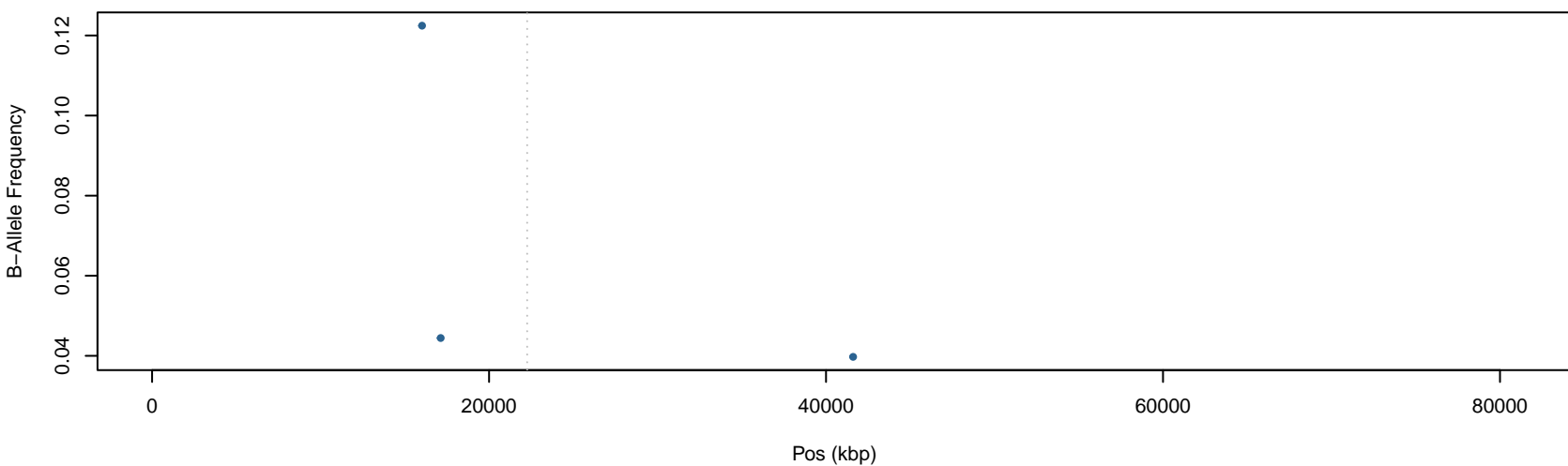
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 15



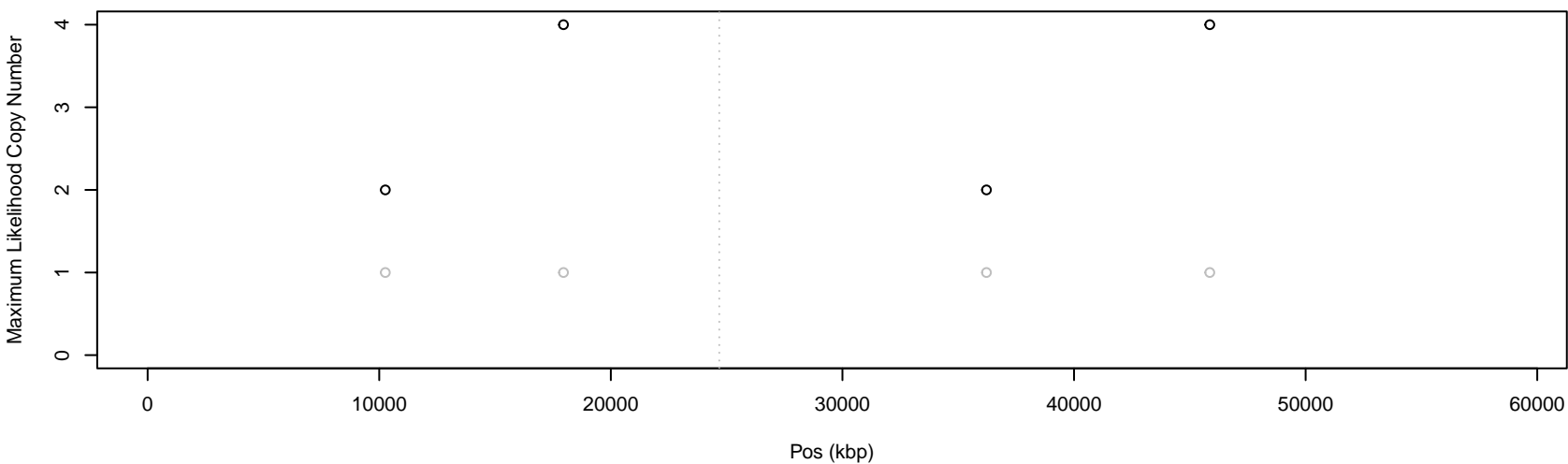
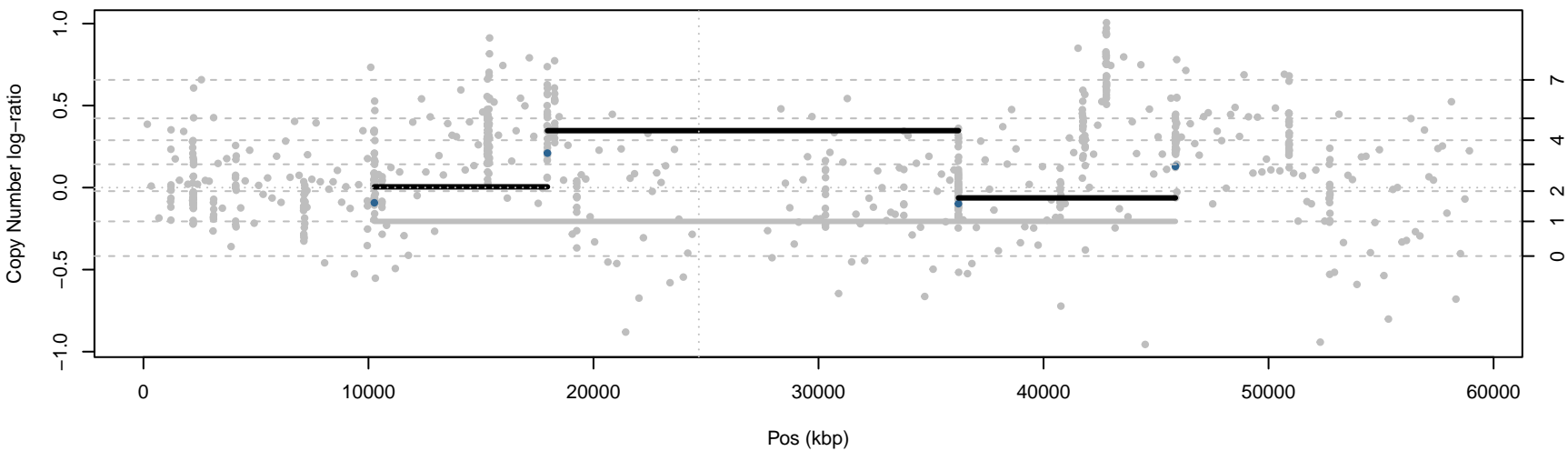
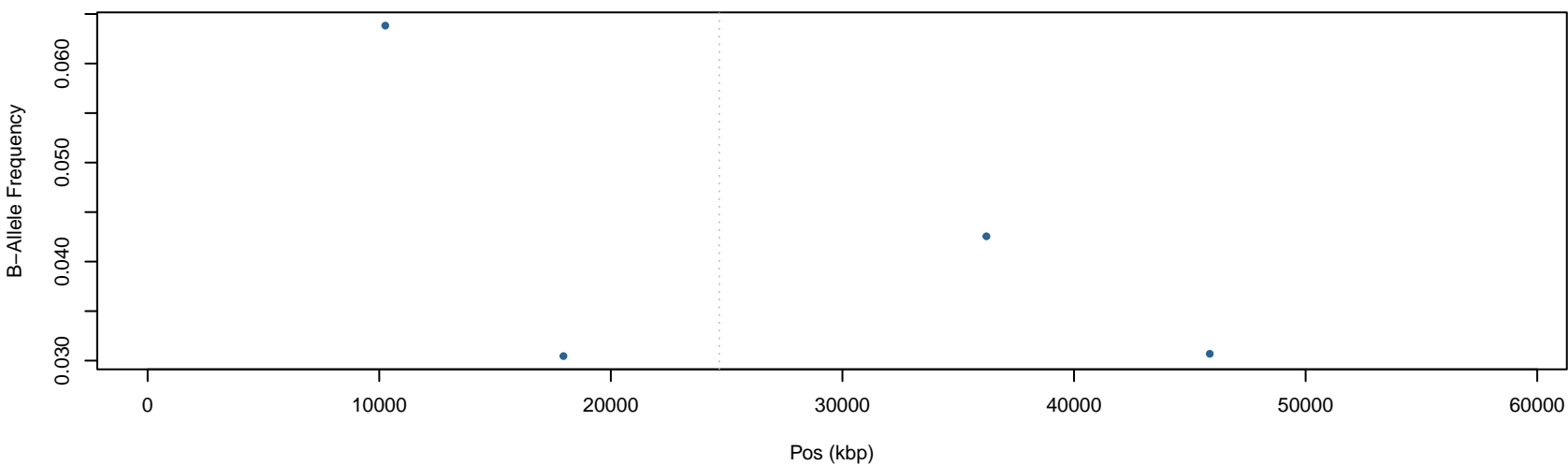
Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 16



Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 17

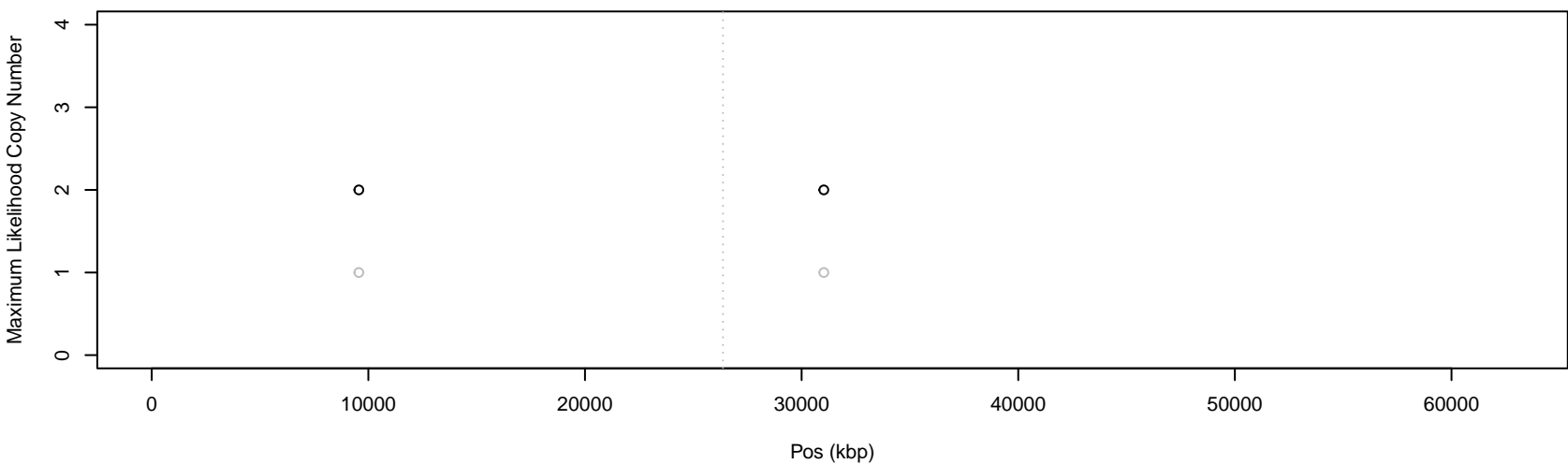
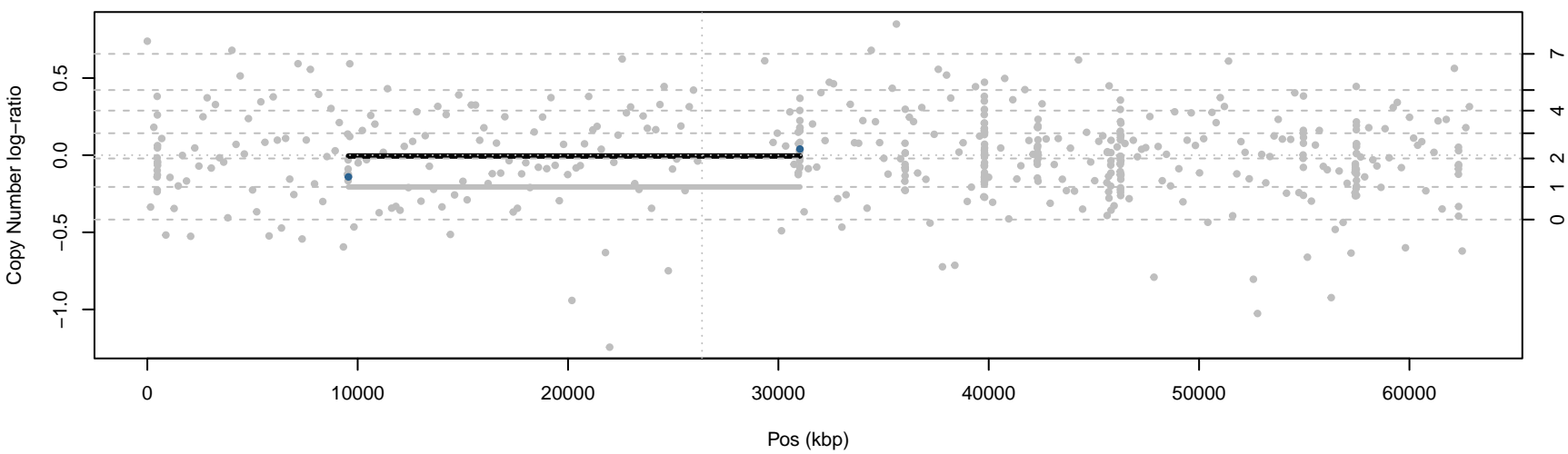
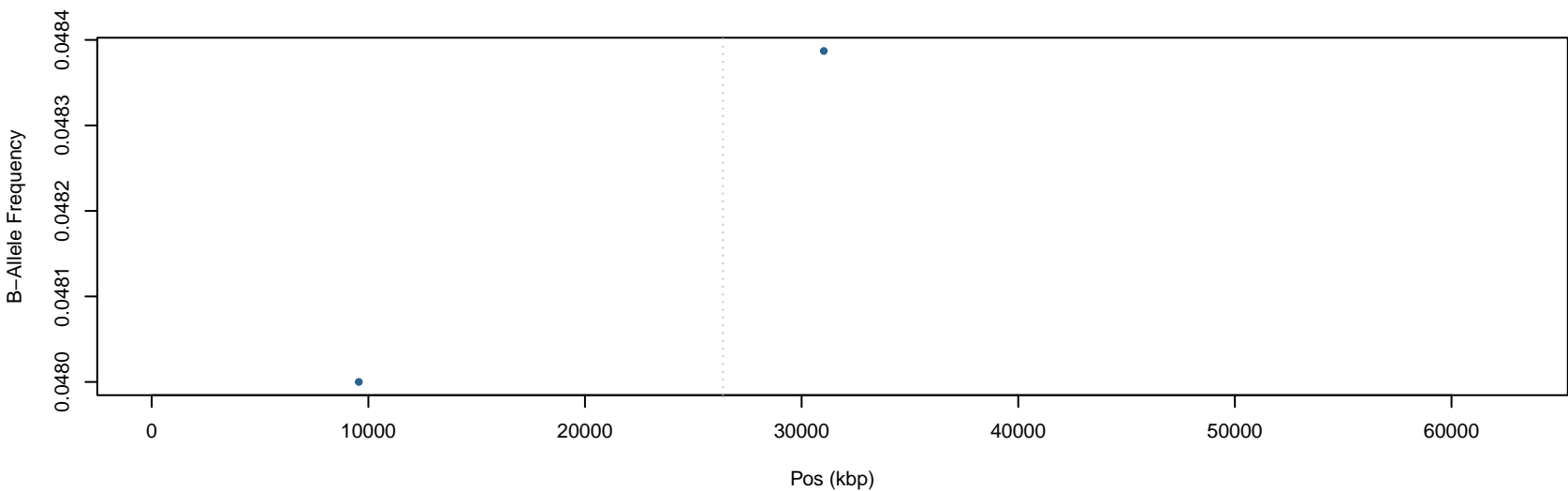


Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 19





Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 20



Purity: 0.24 Tumor ploidy: 2.125 SNV log-likelihood: 921.8 GoF: 87.3% Mean coverage: 149;174 Chromosome: 22

