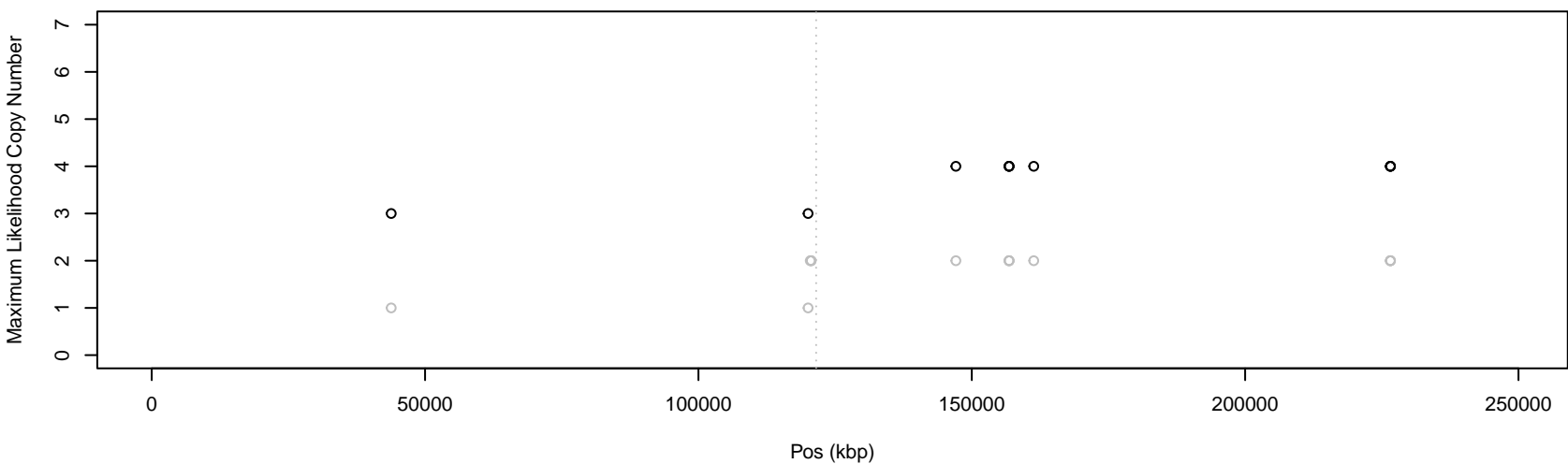
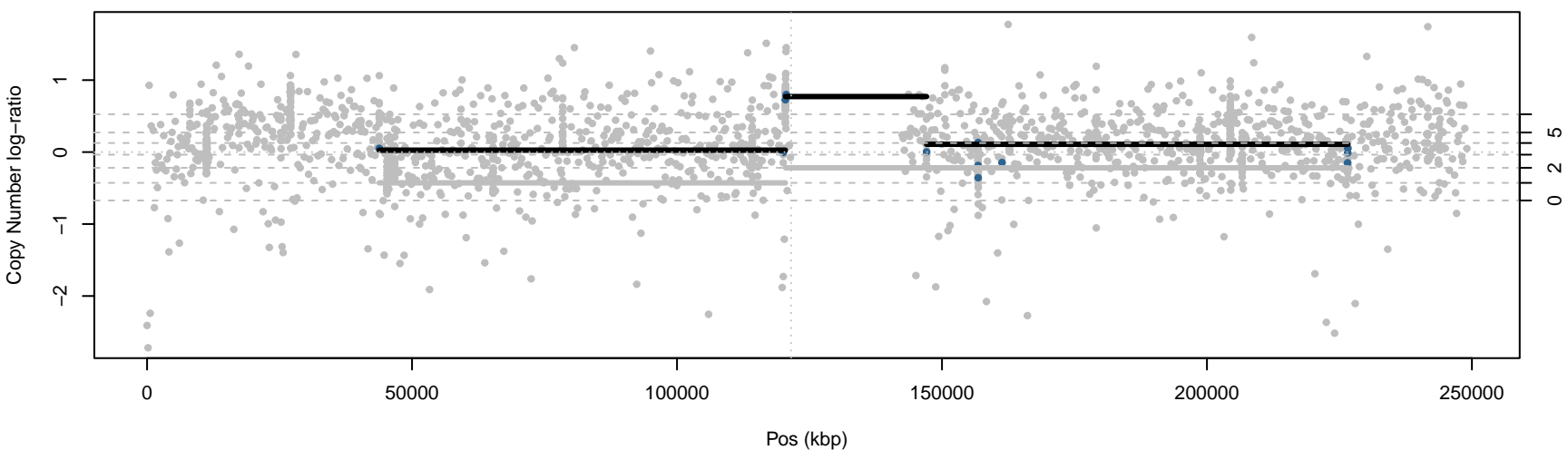
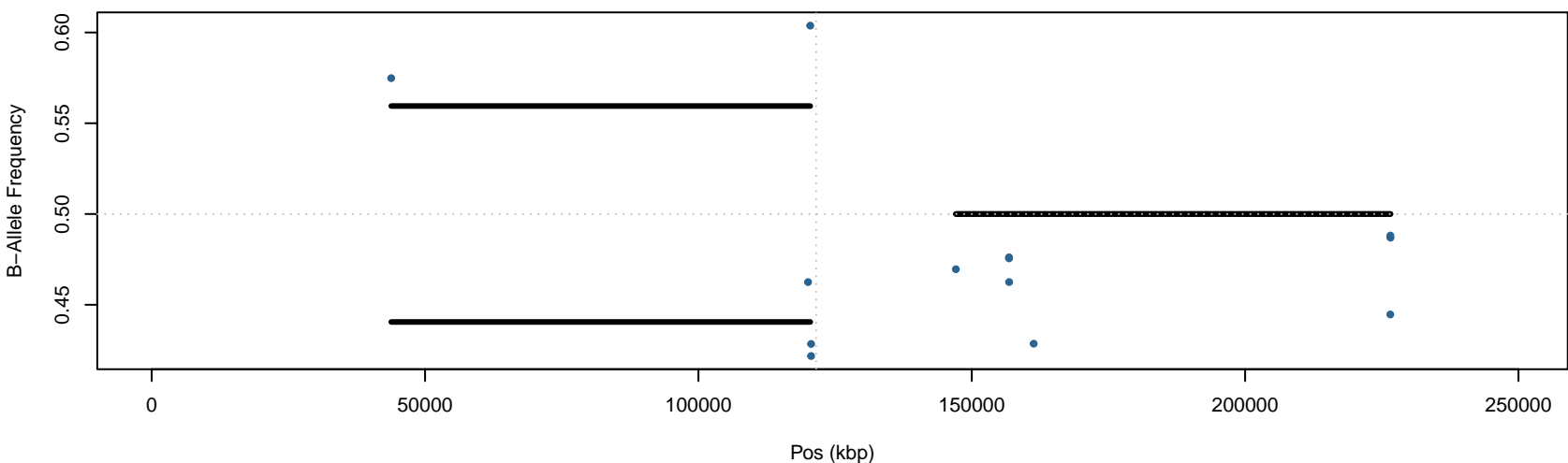
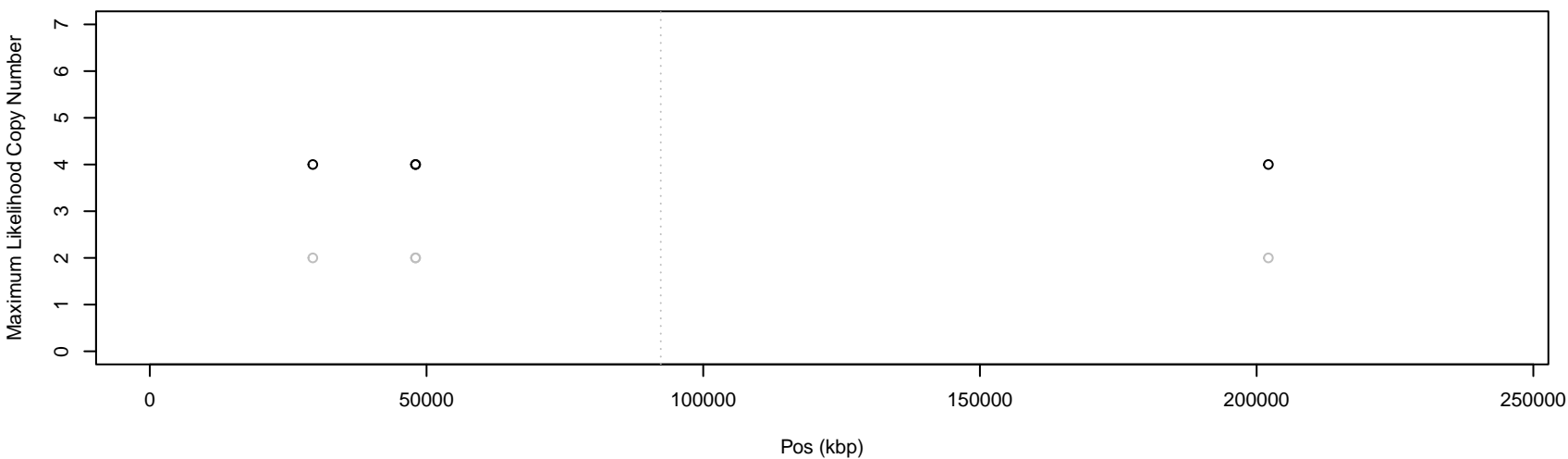
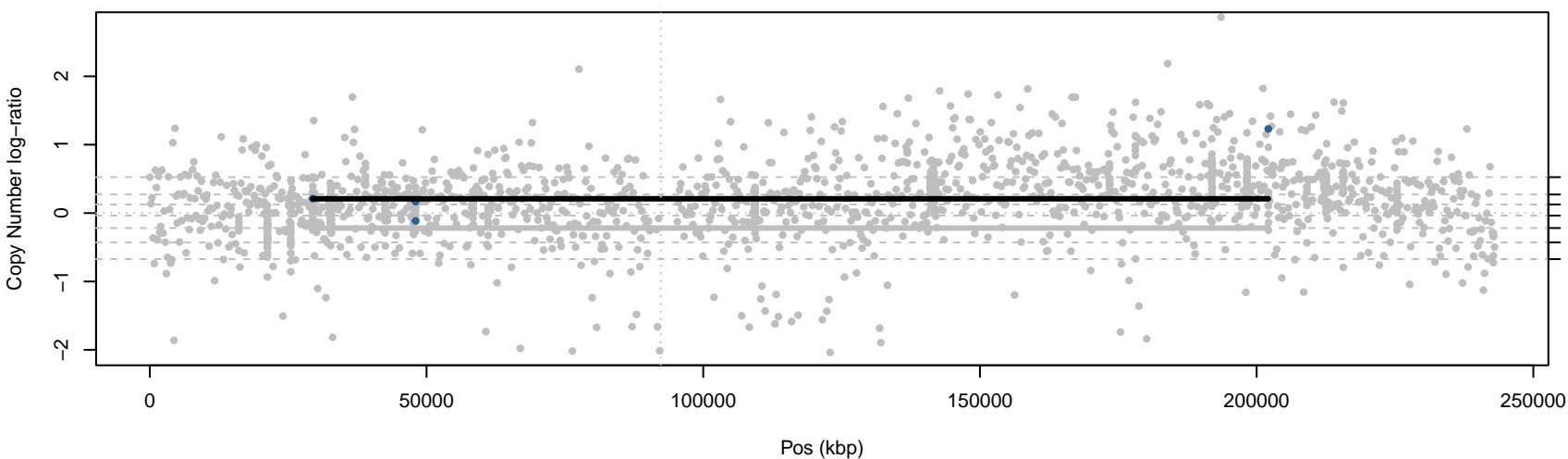
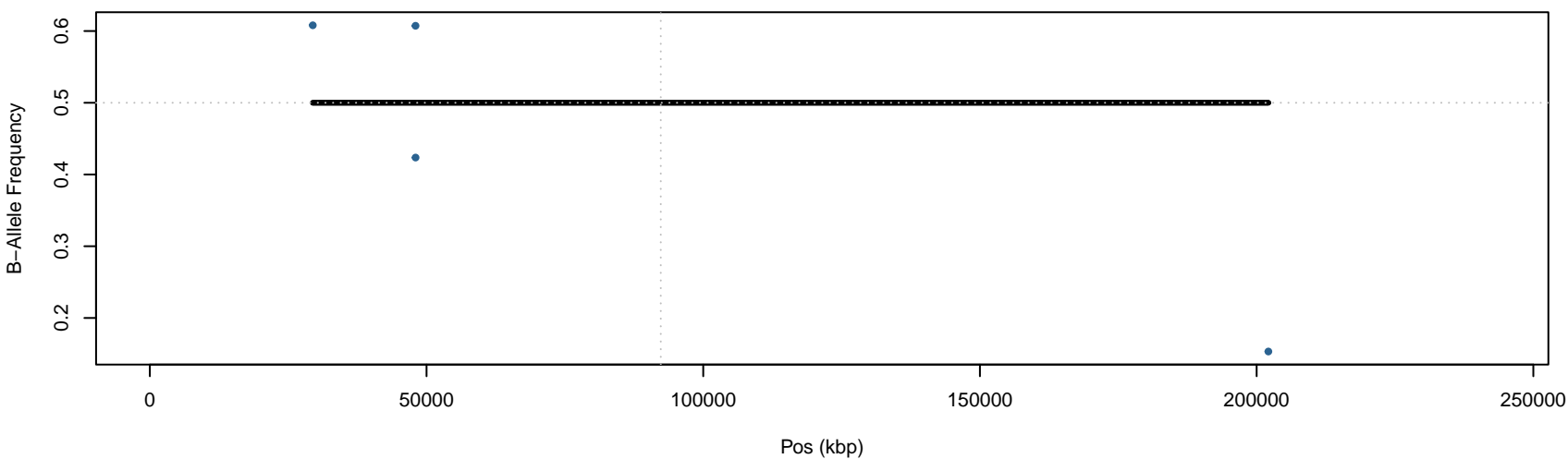


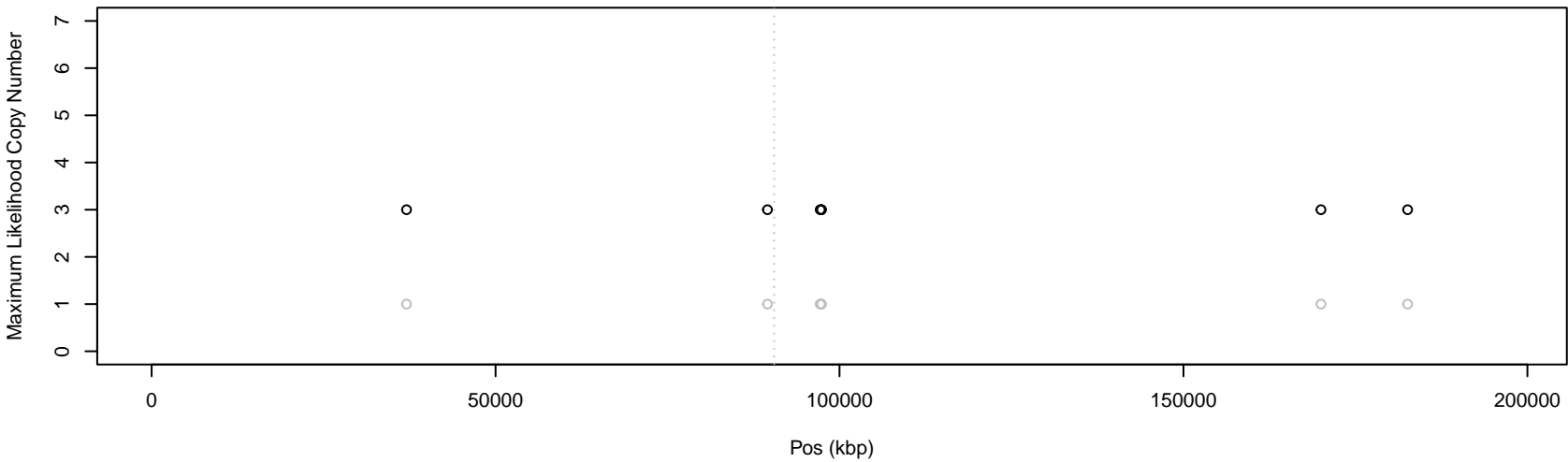
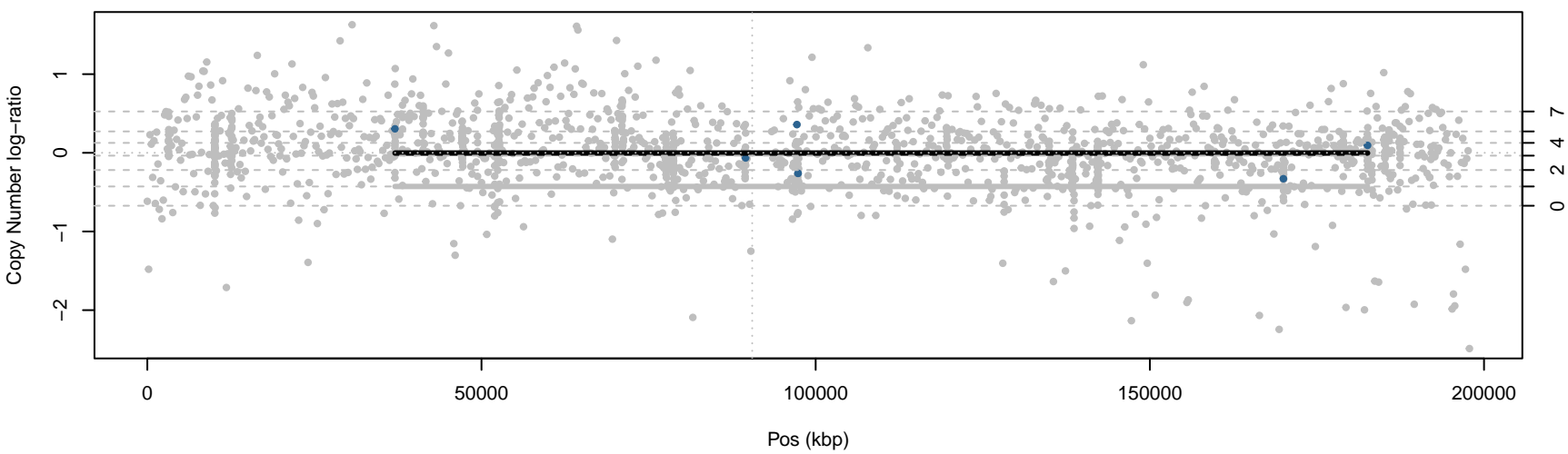
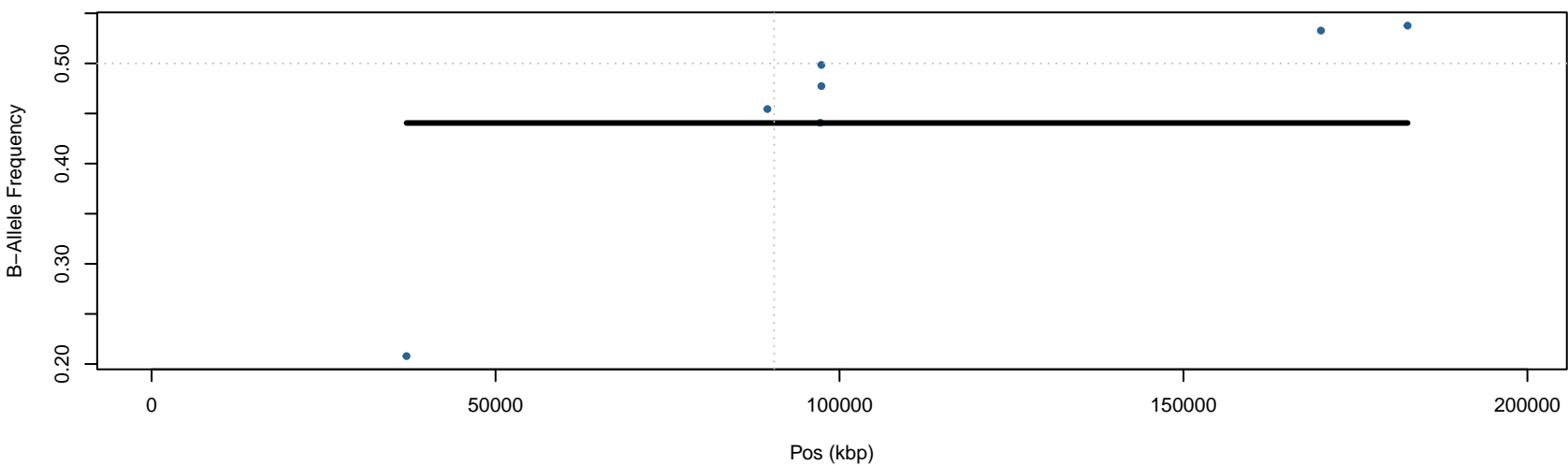
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 1



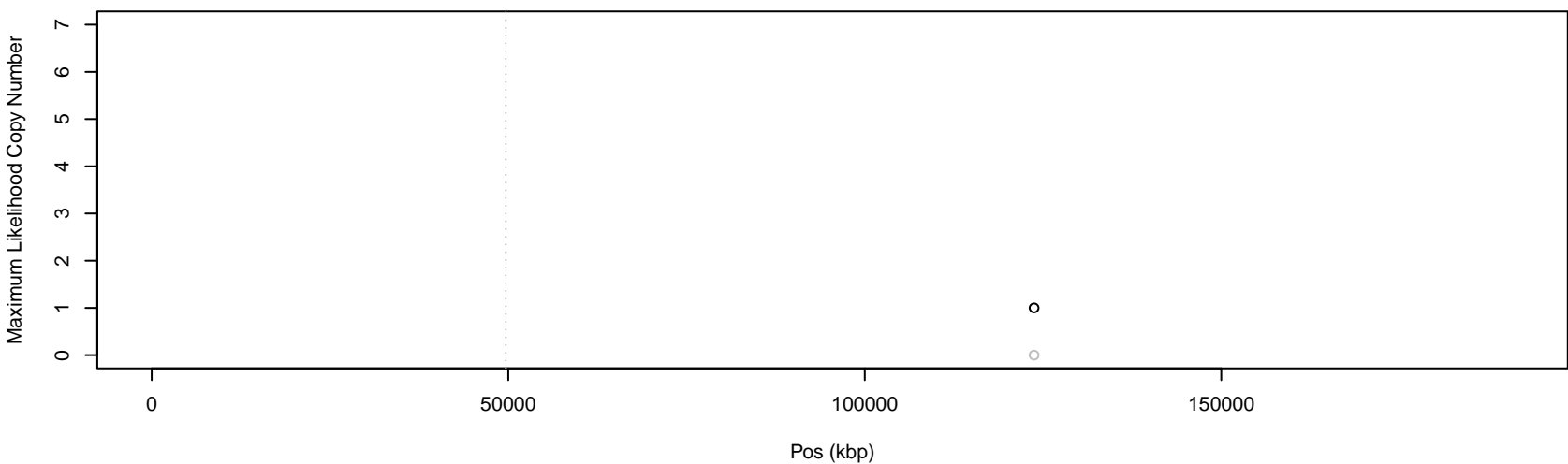
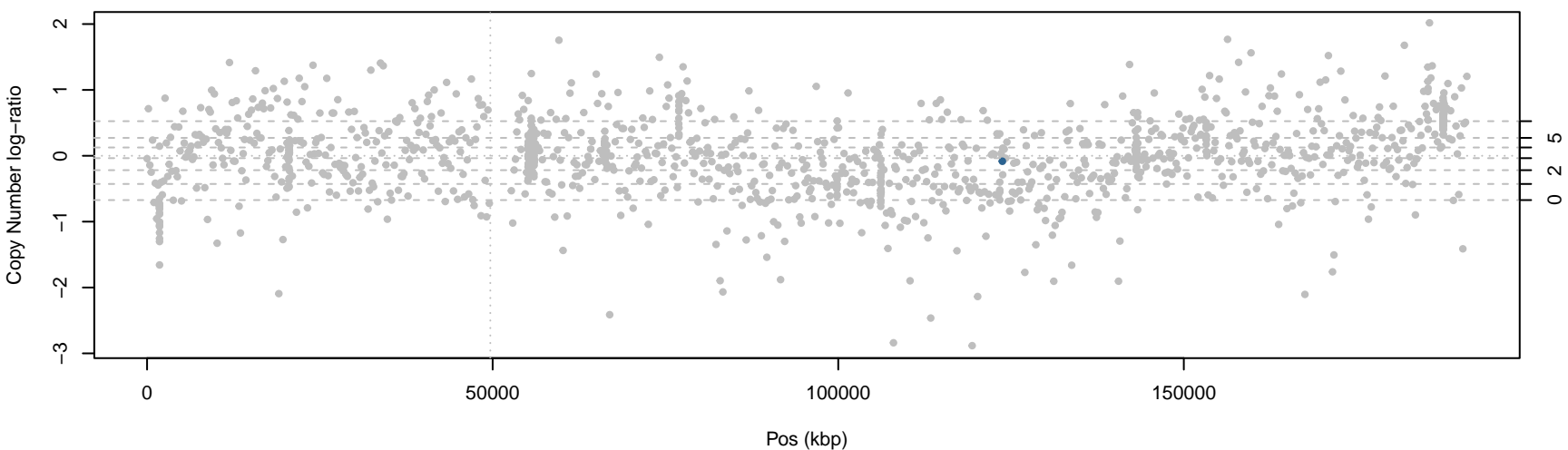
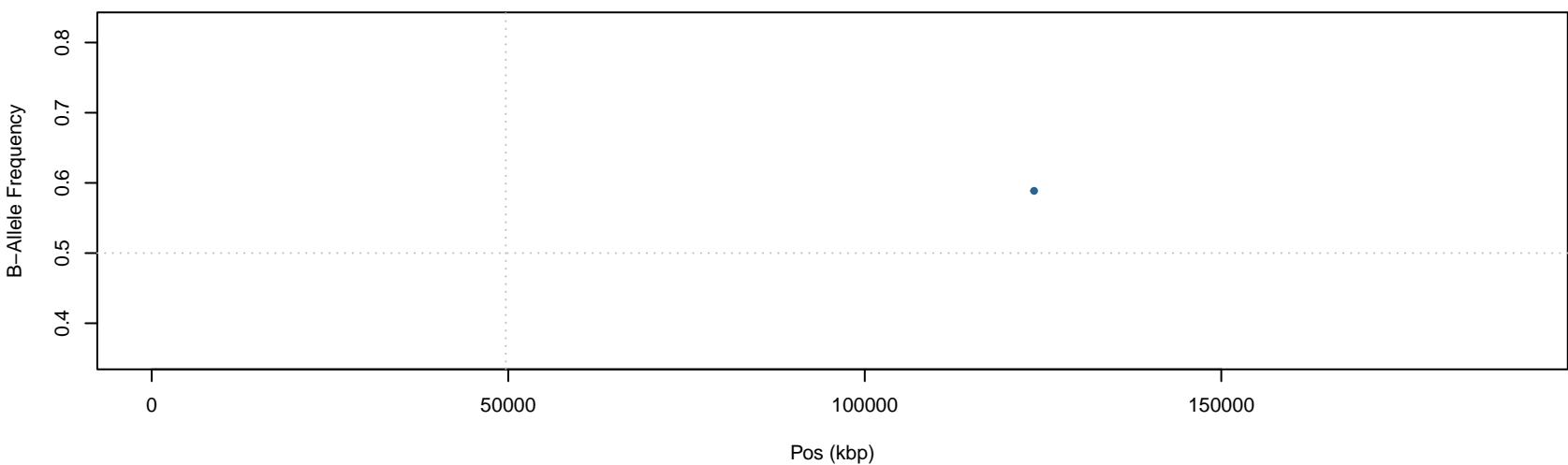
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 2



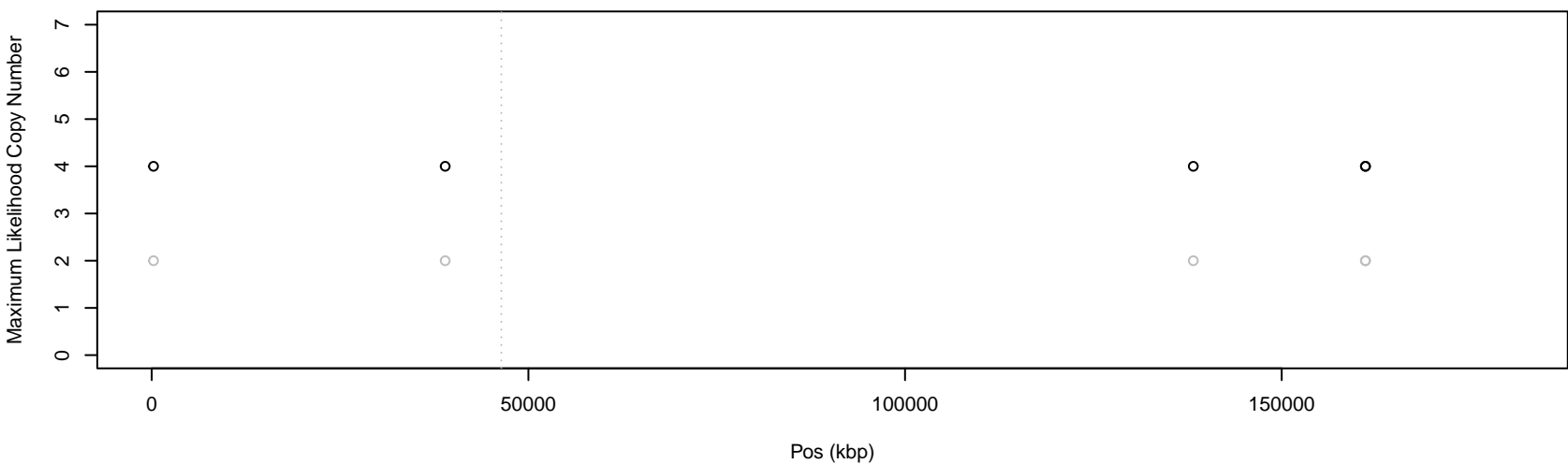
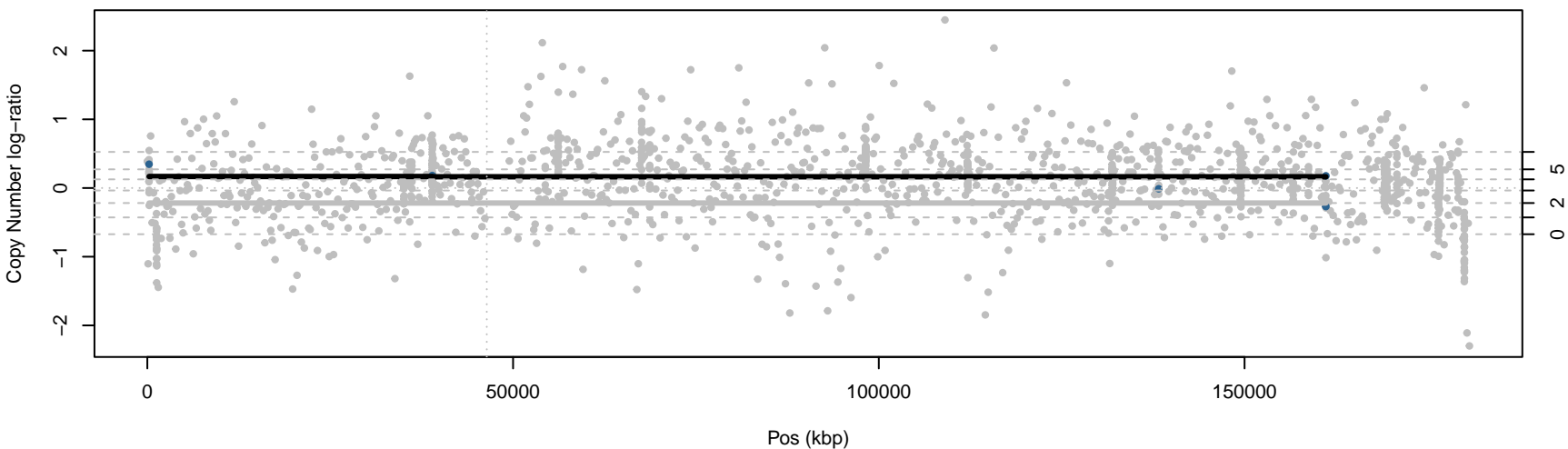
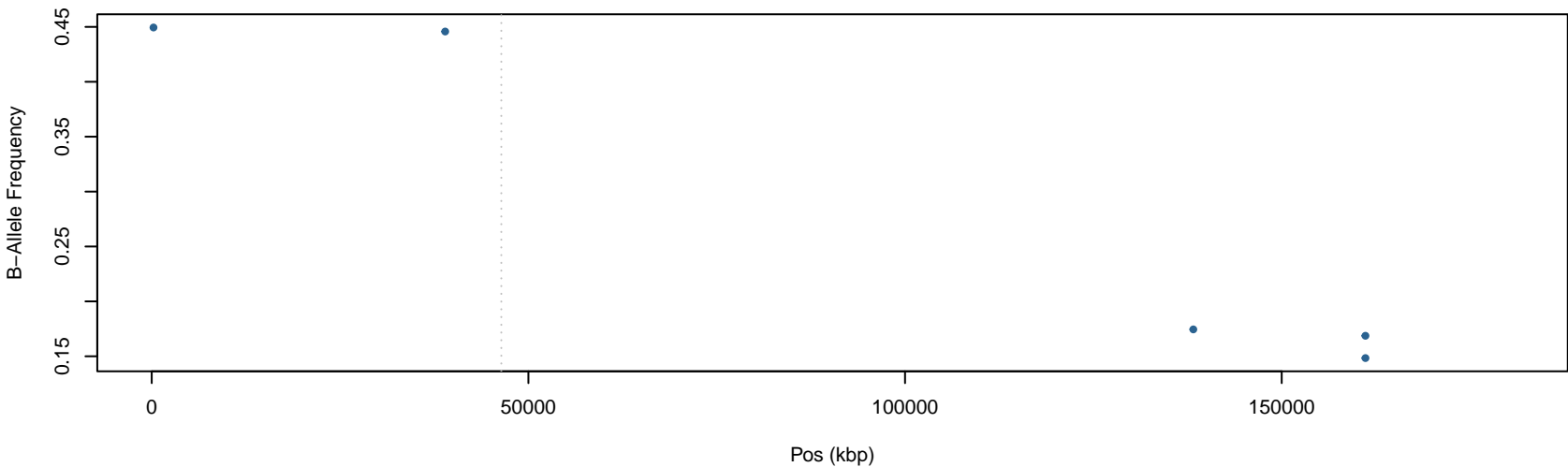
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 3



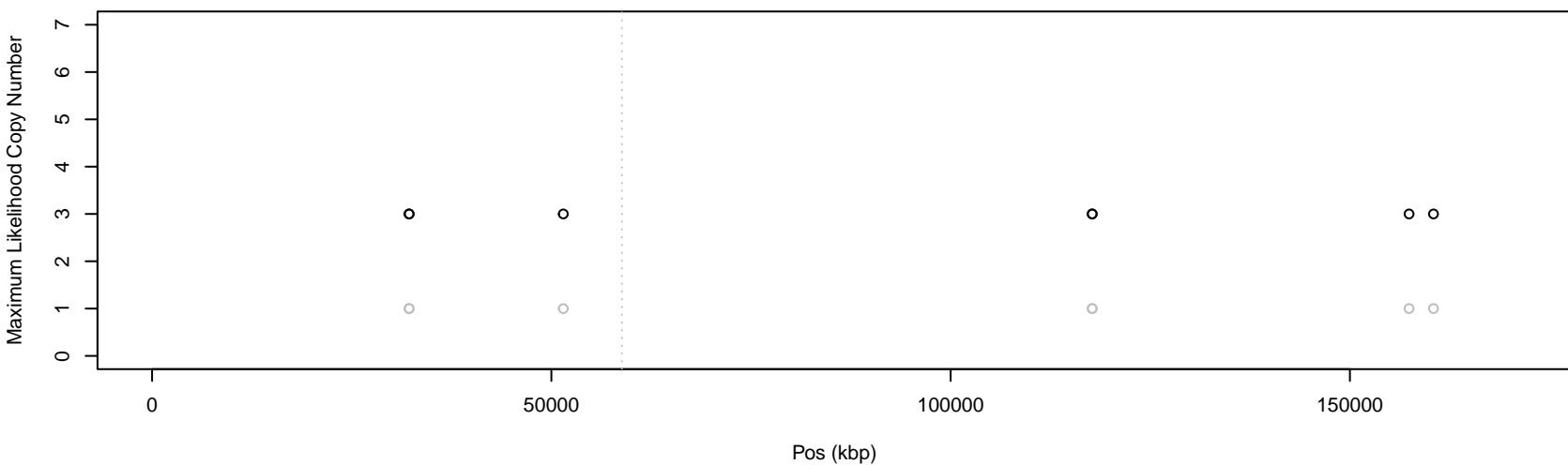
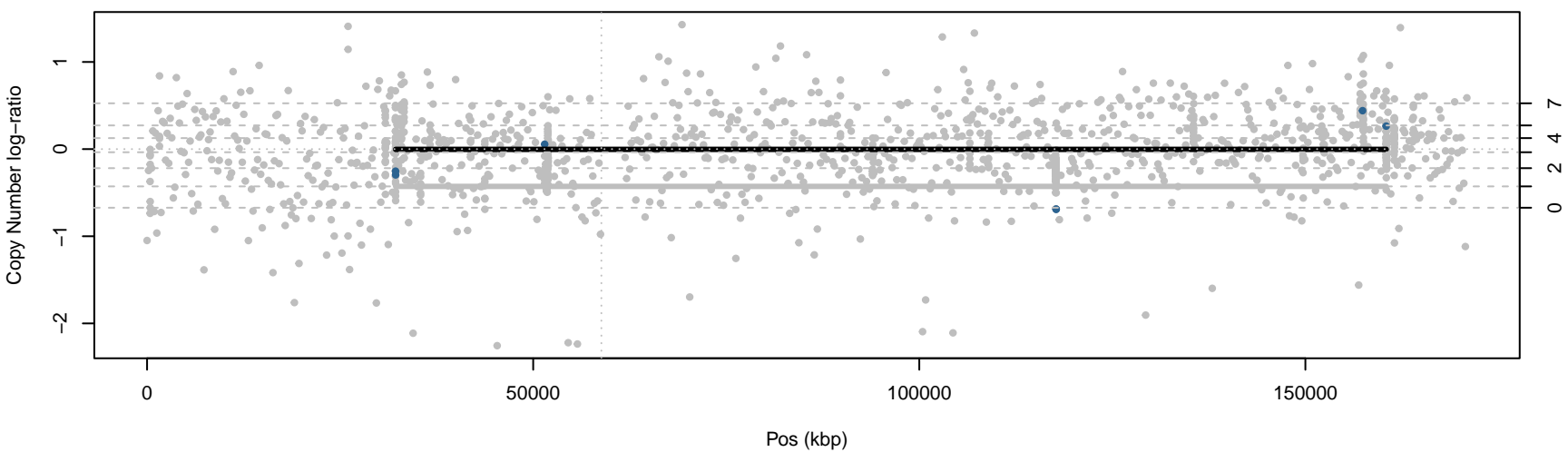
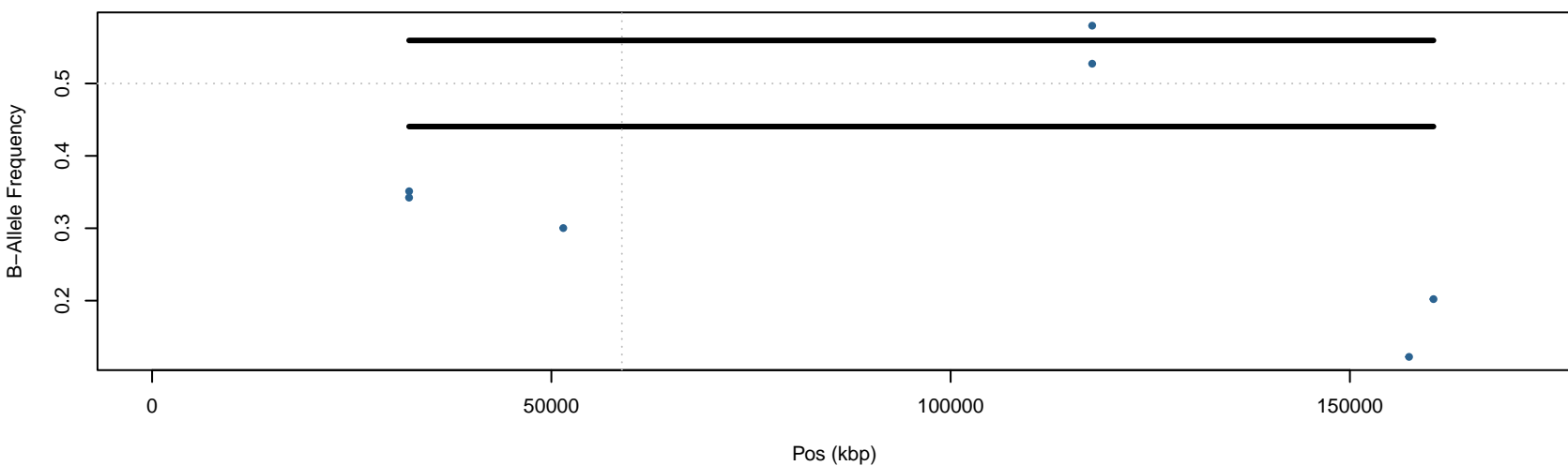
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 4



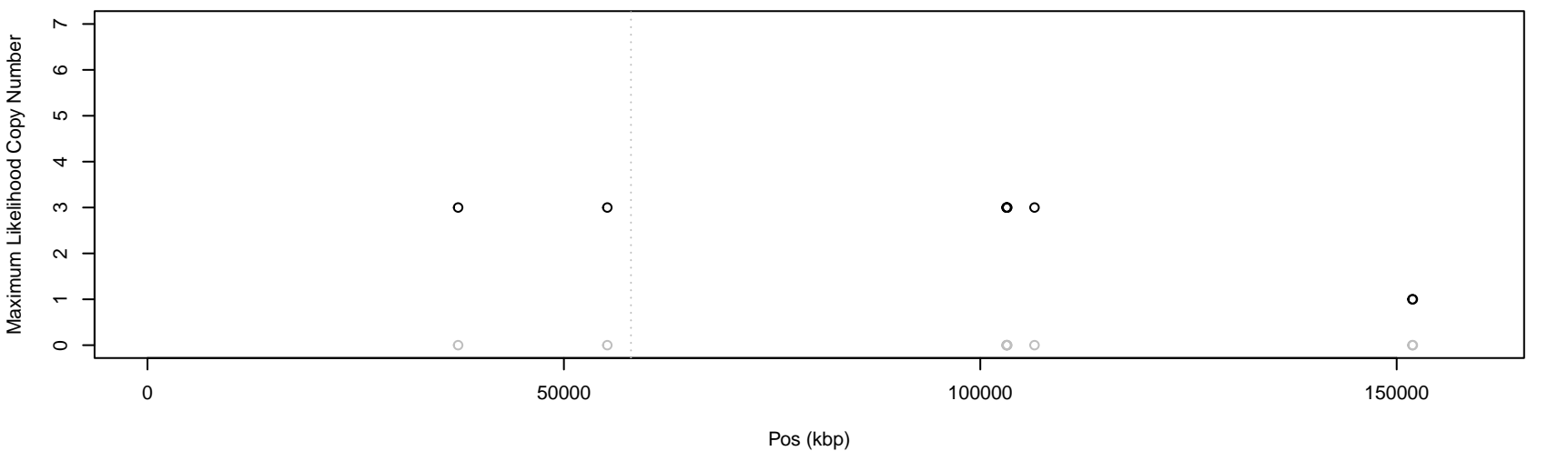
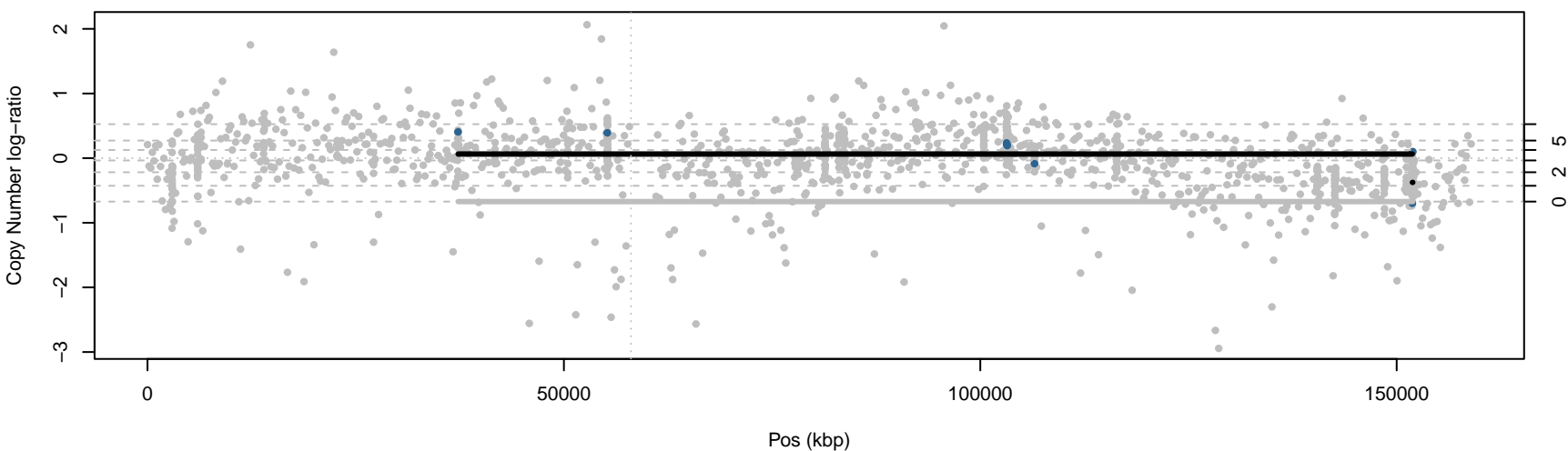
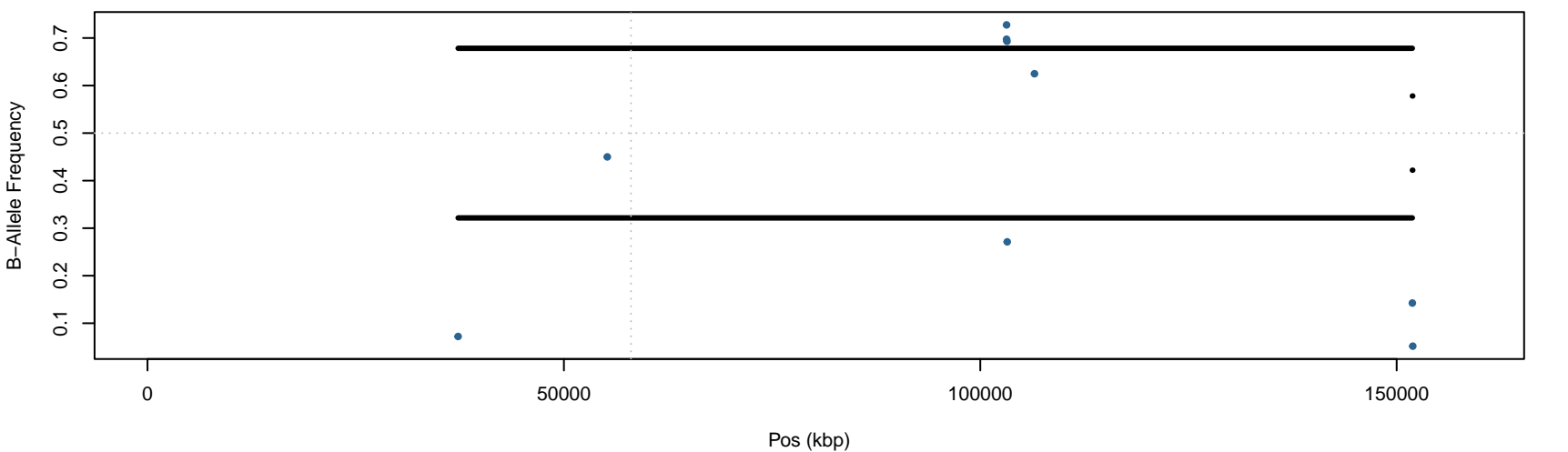
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 5



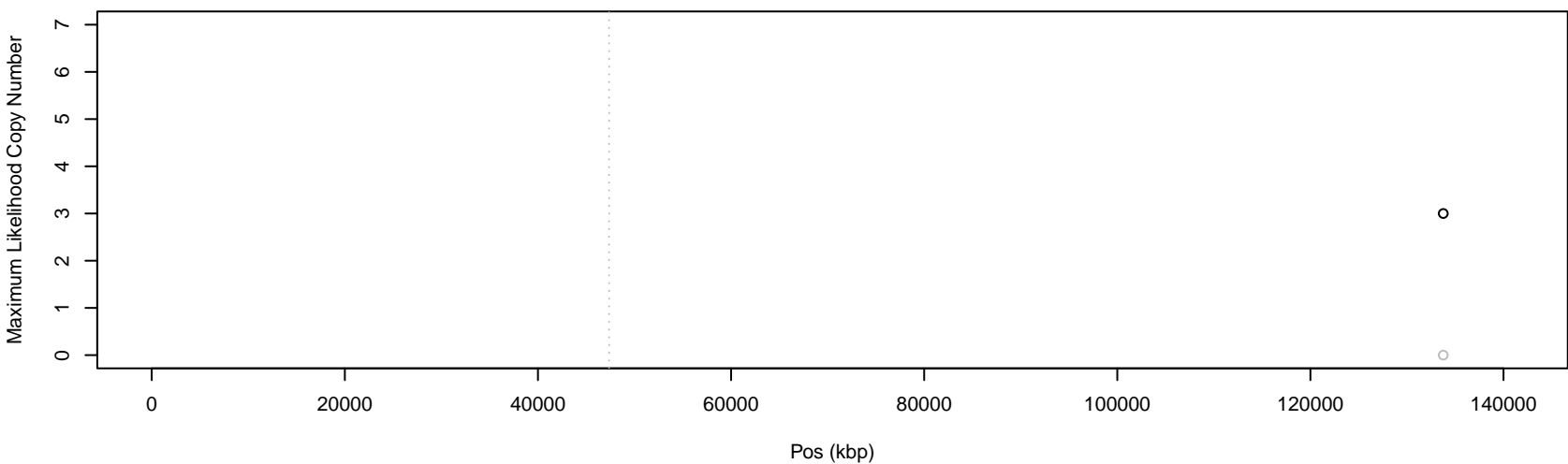
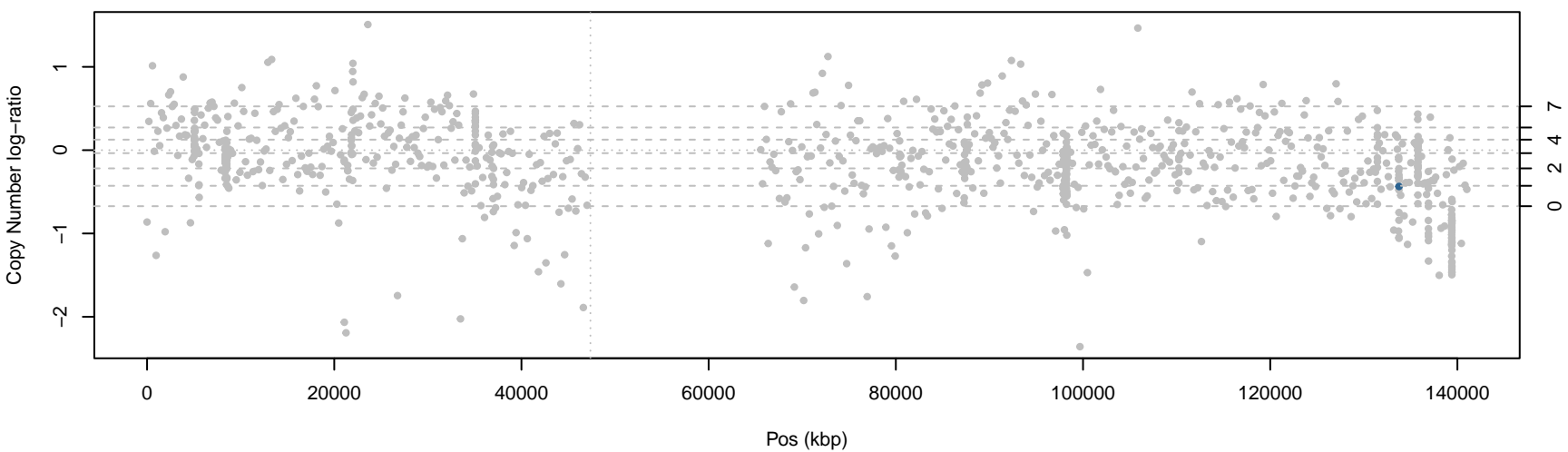
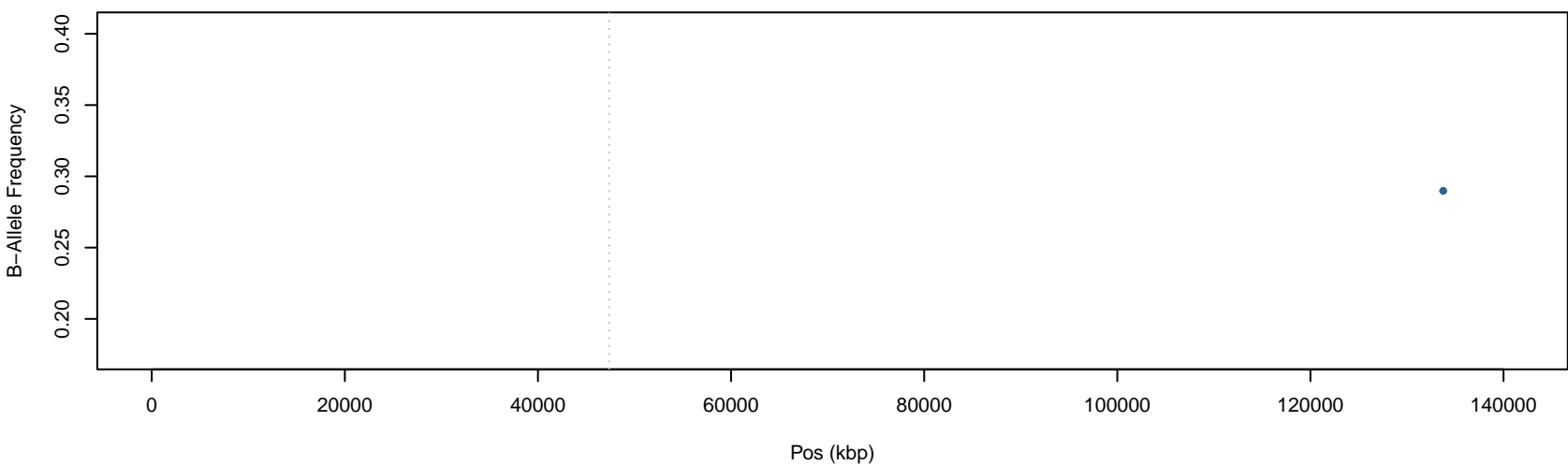
Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 6



Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 7

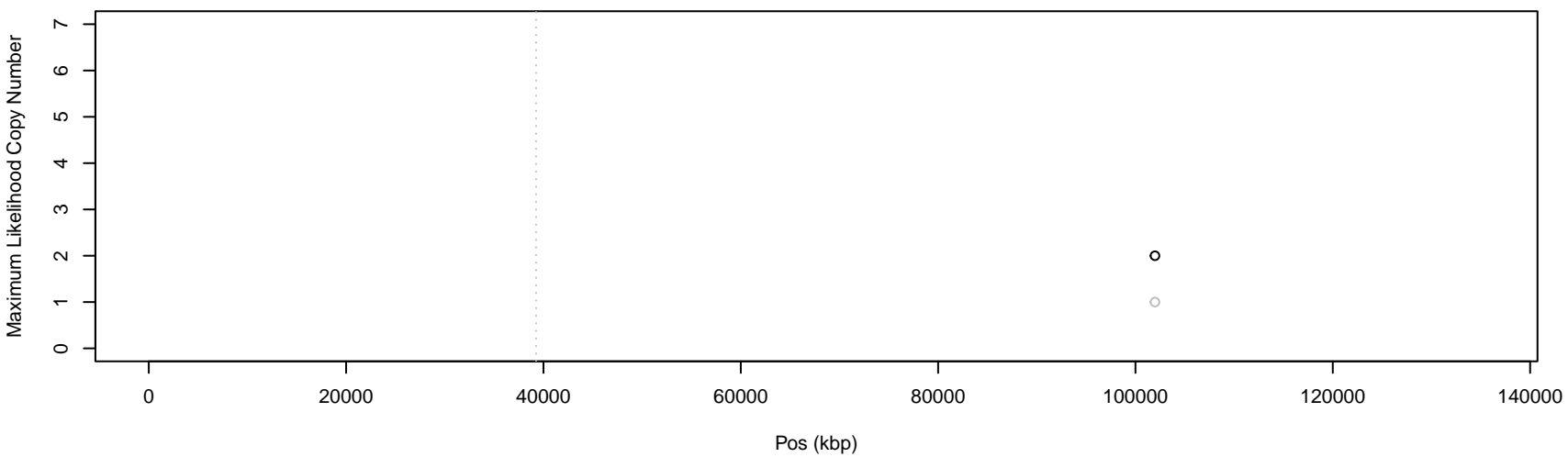
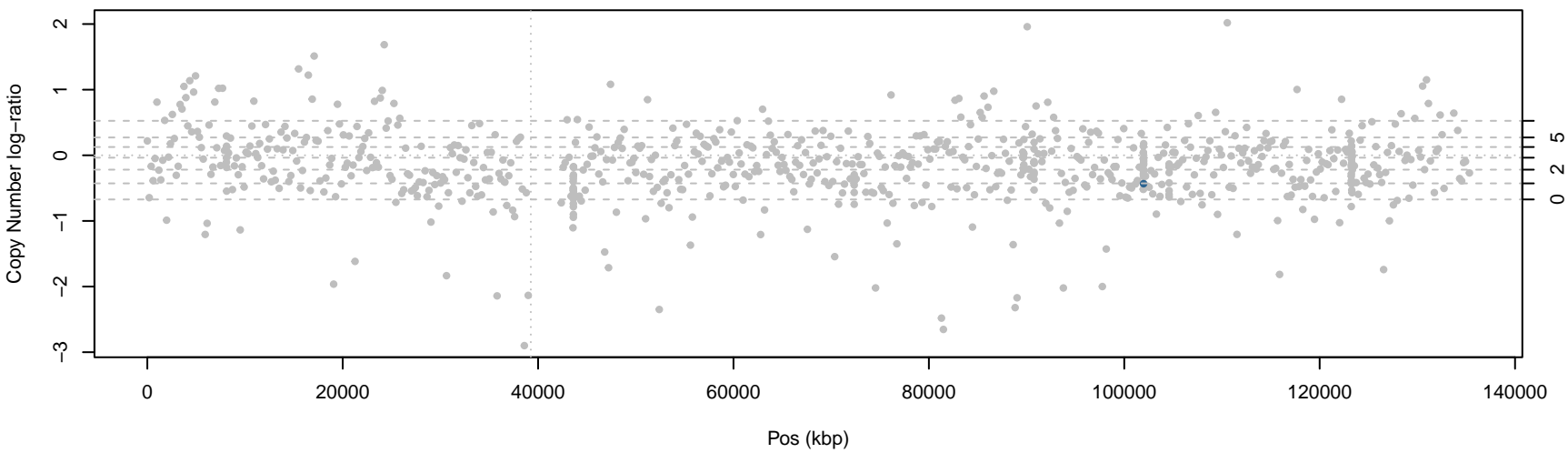
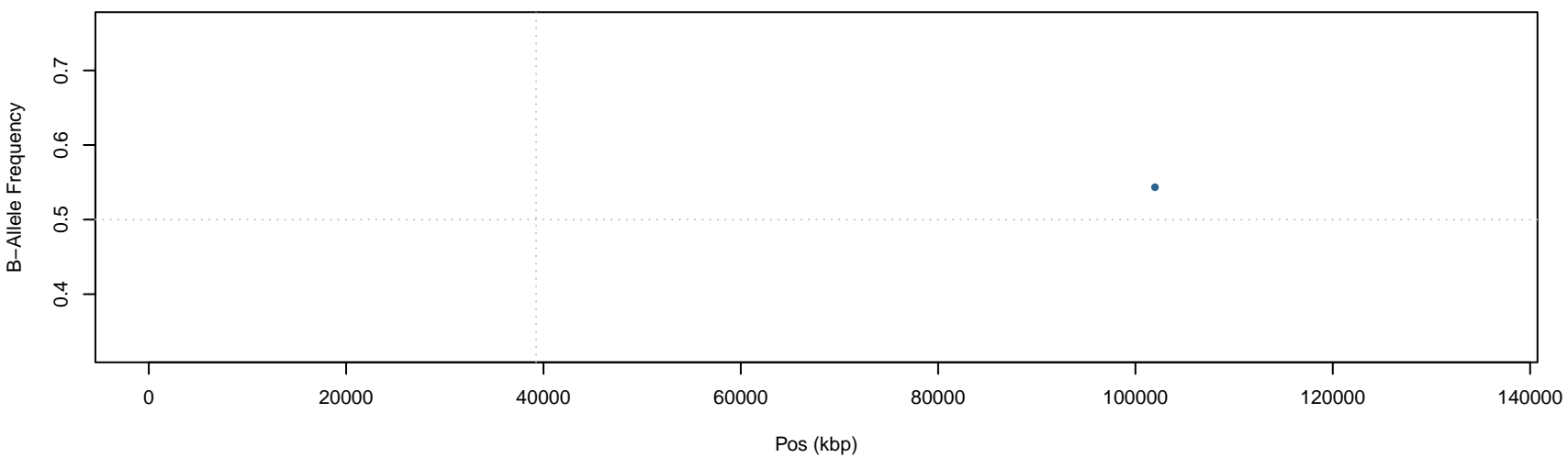


Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 9

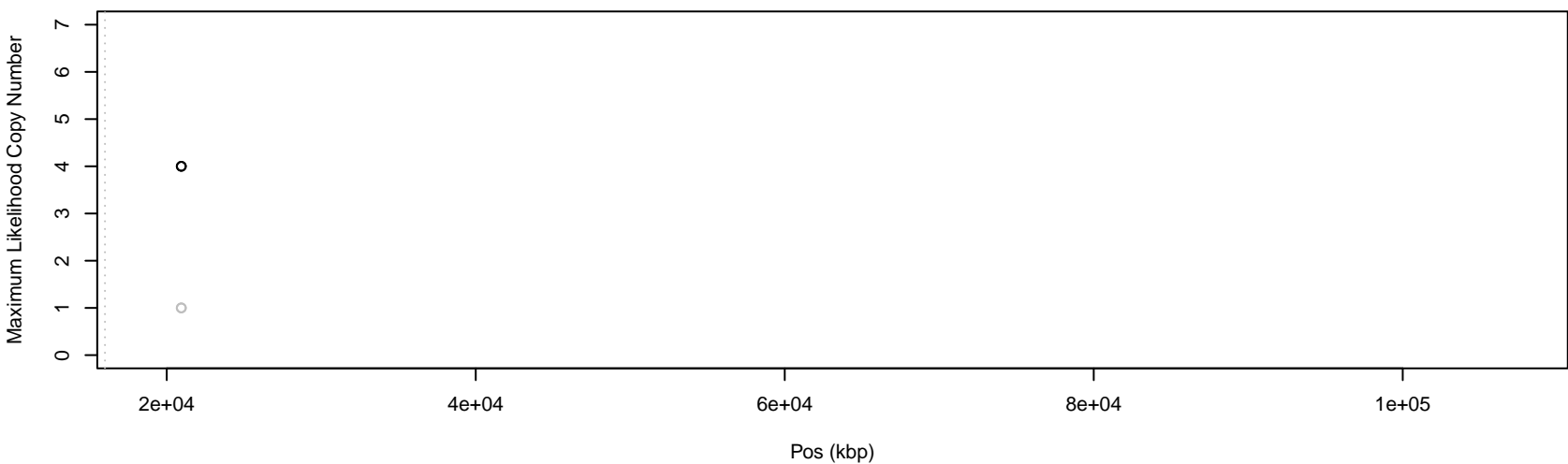
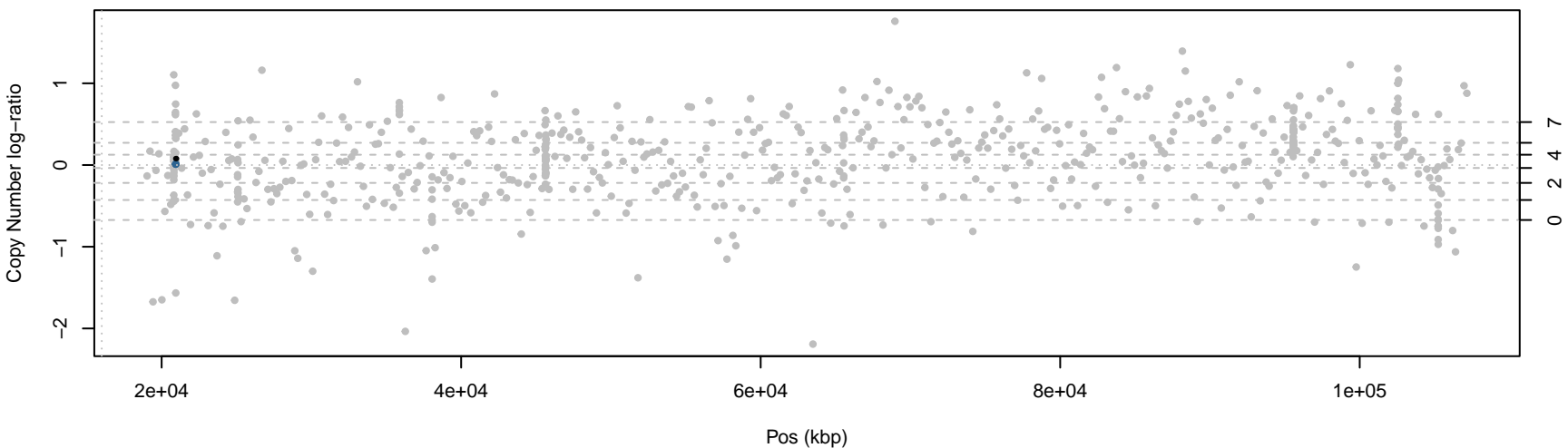
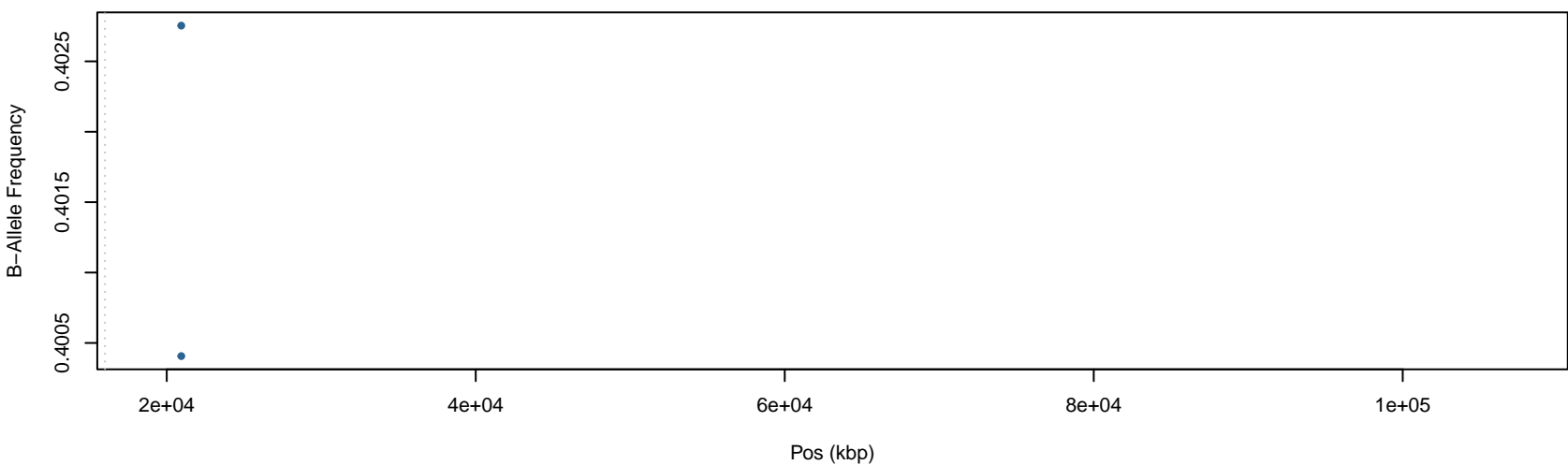




Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 10



Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 14



Purity: 0.27 Tumor ploidy: 3.213 SNV log-likelihood: -63.89 GoF: 93.5% Mean coverage: 586;597 Chromosome: 19

