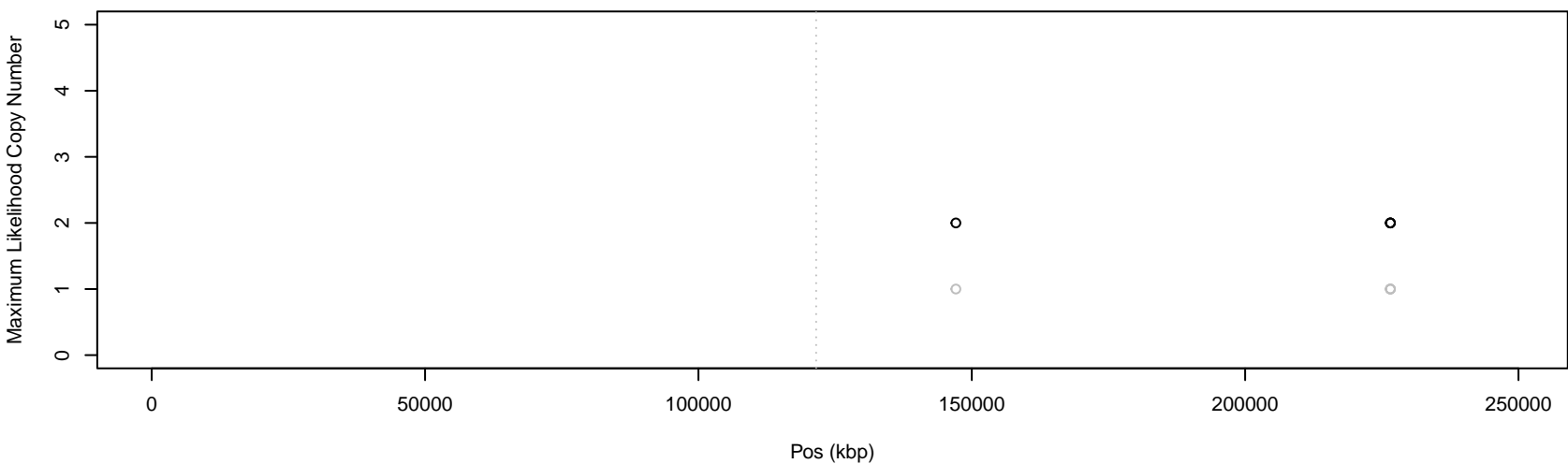
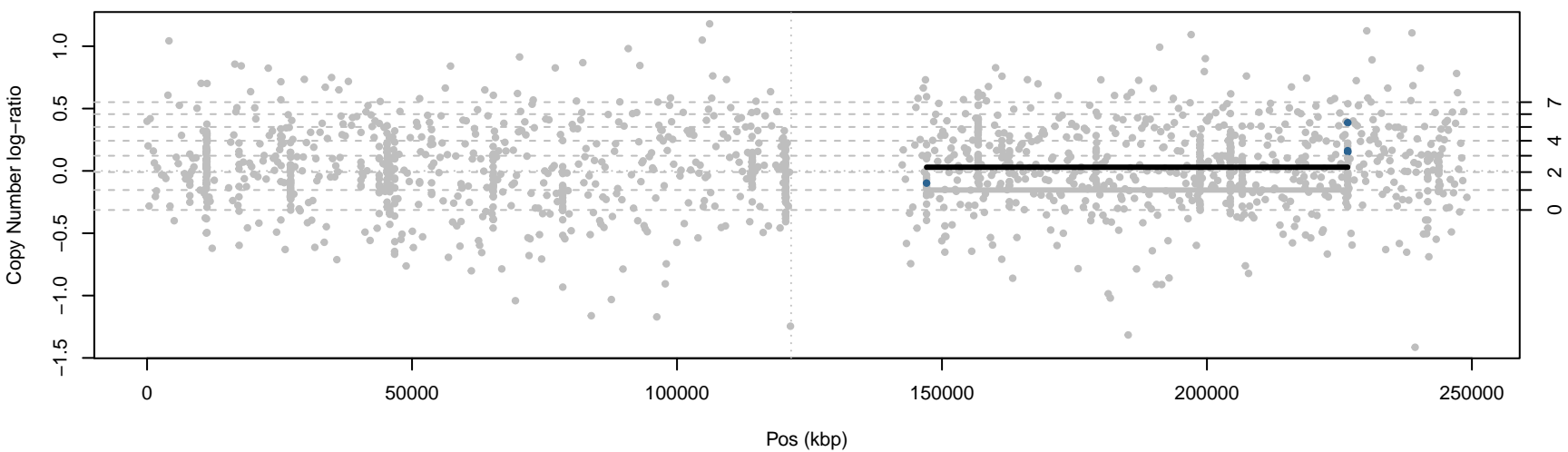
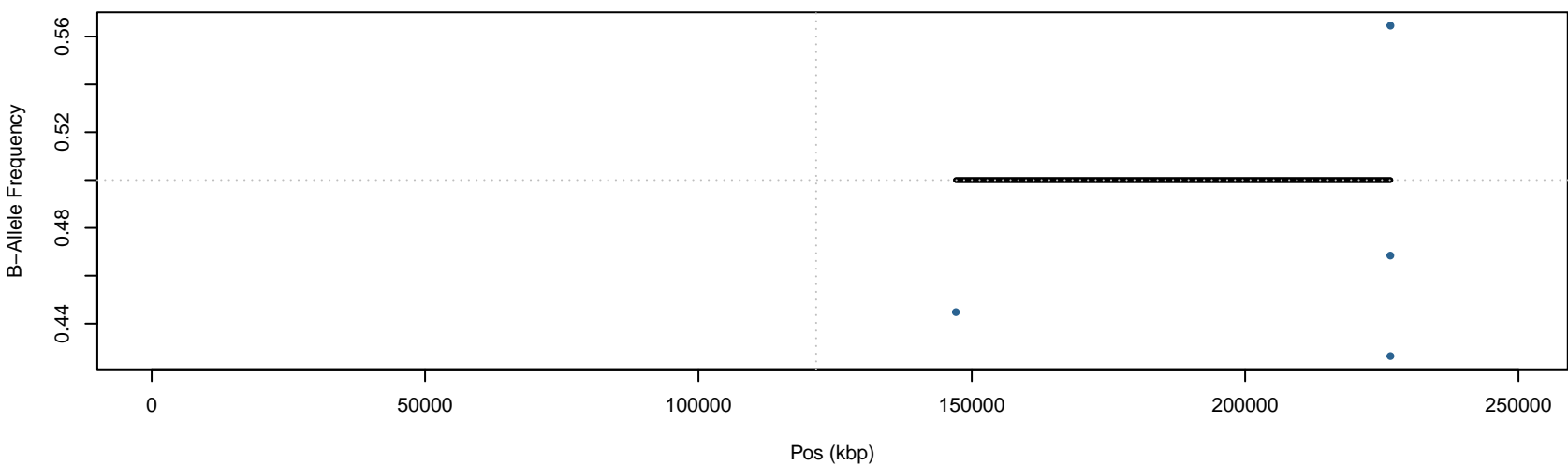
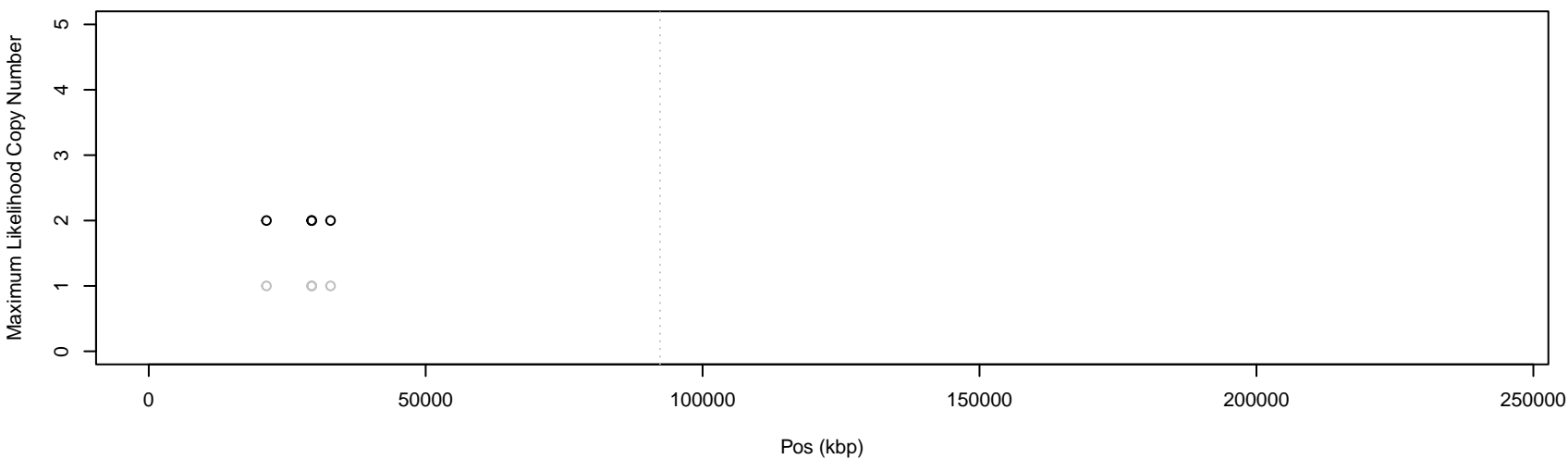
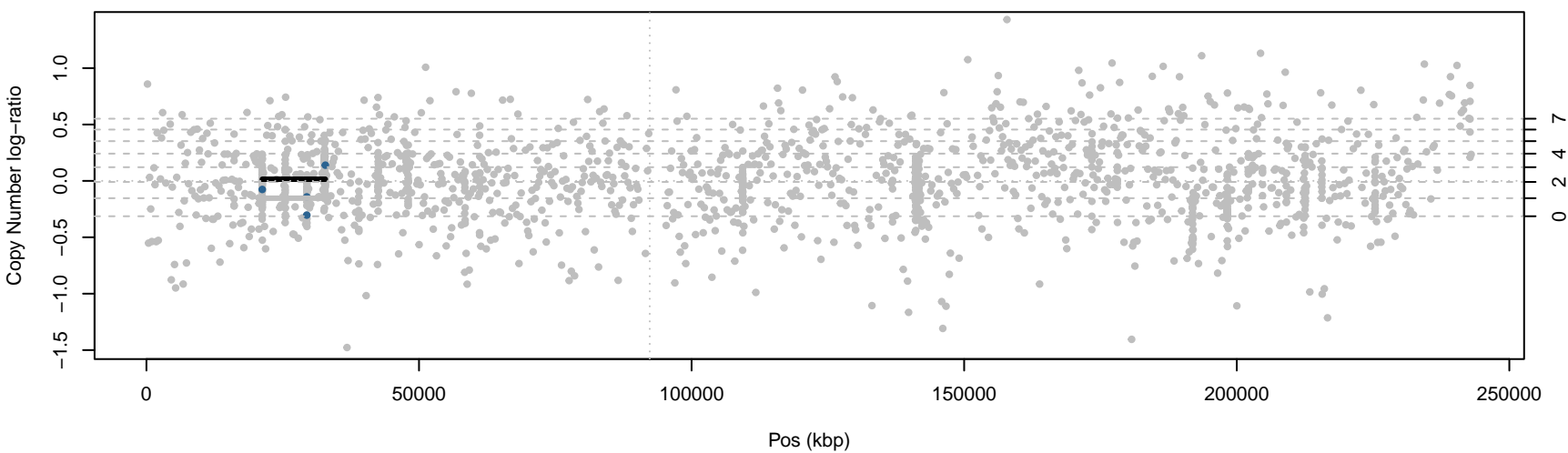
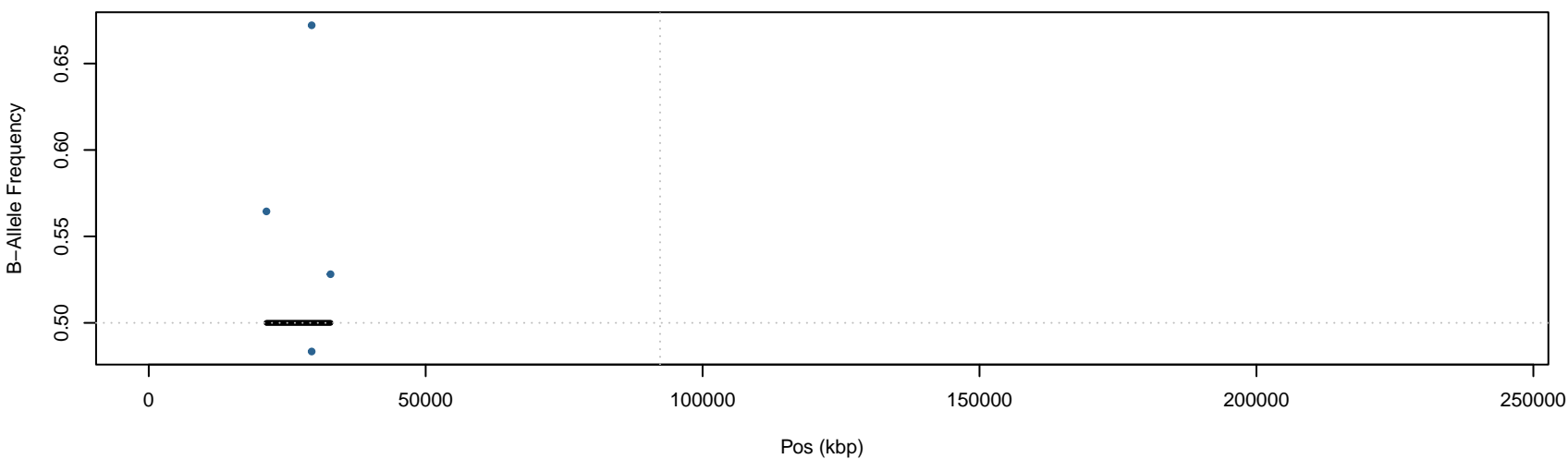


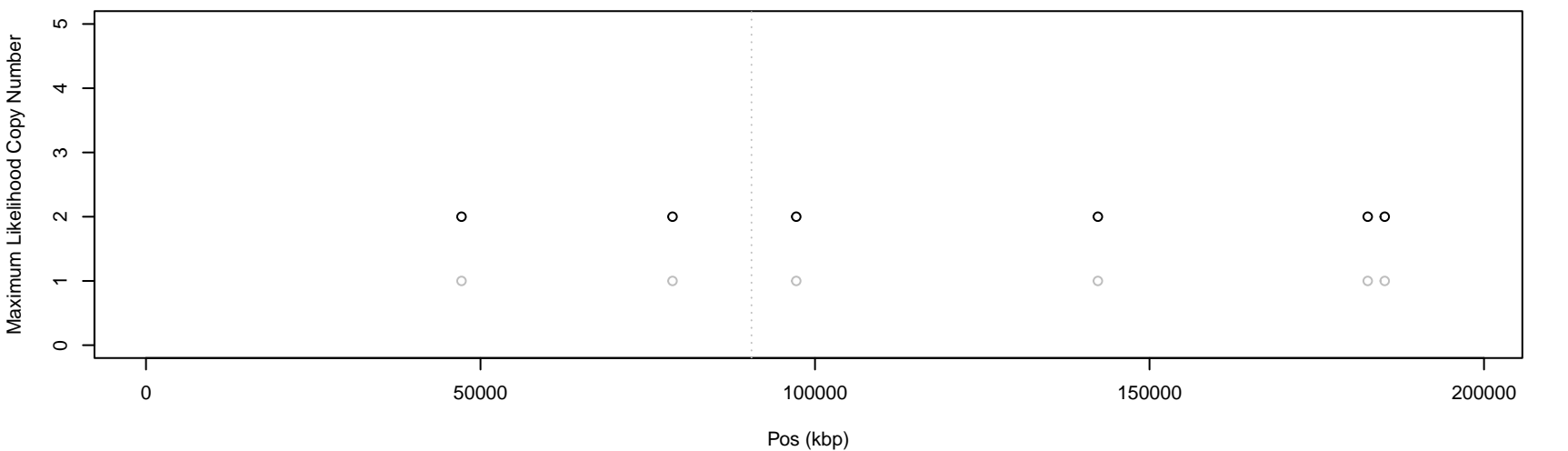
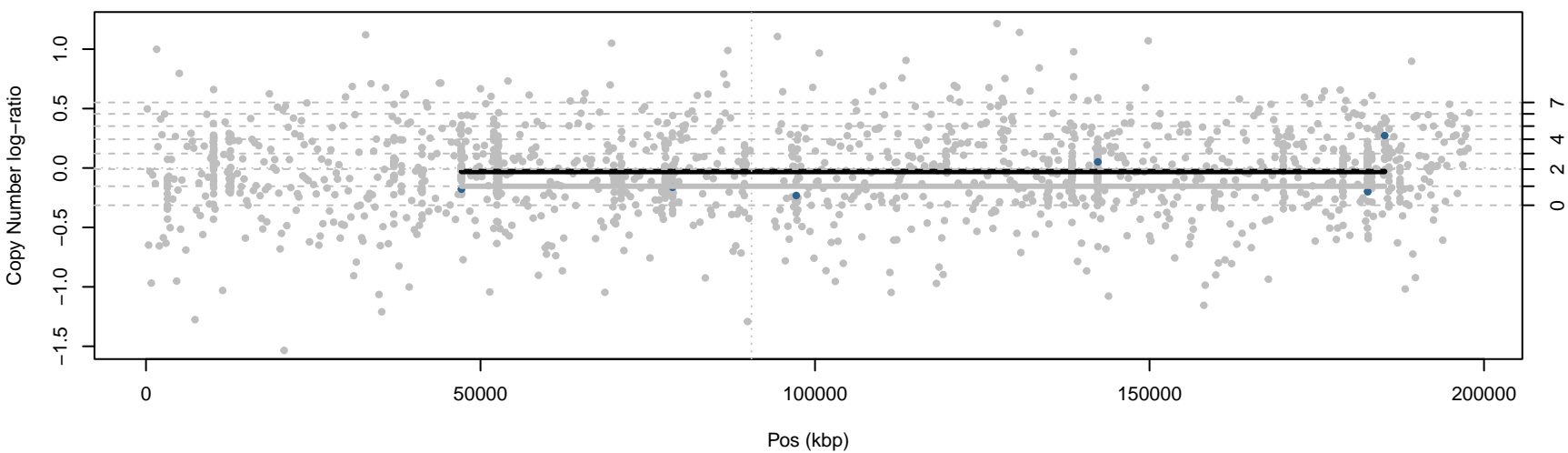
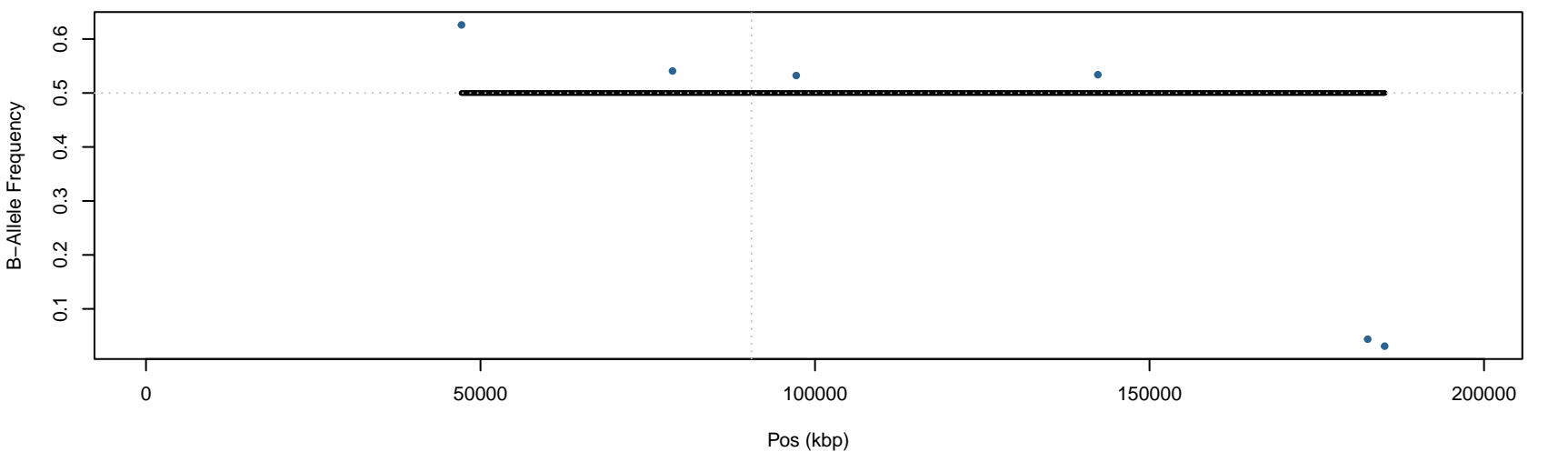
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 1



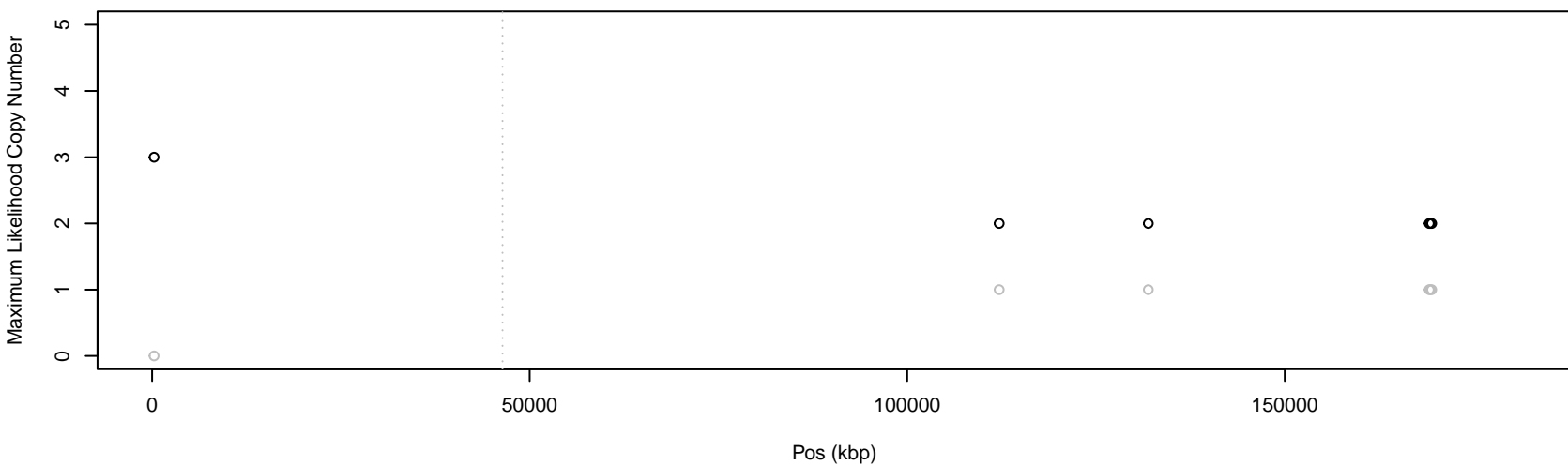
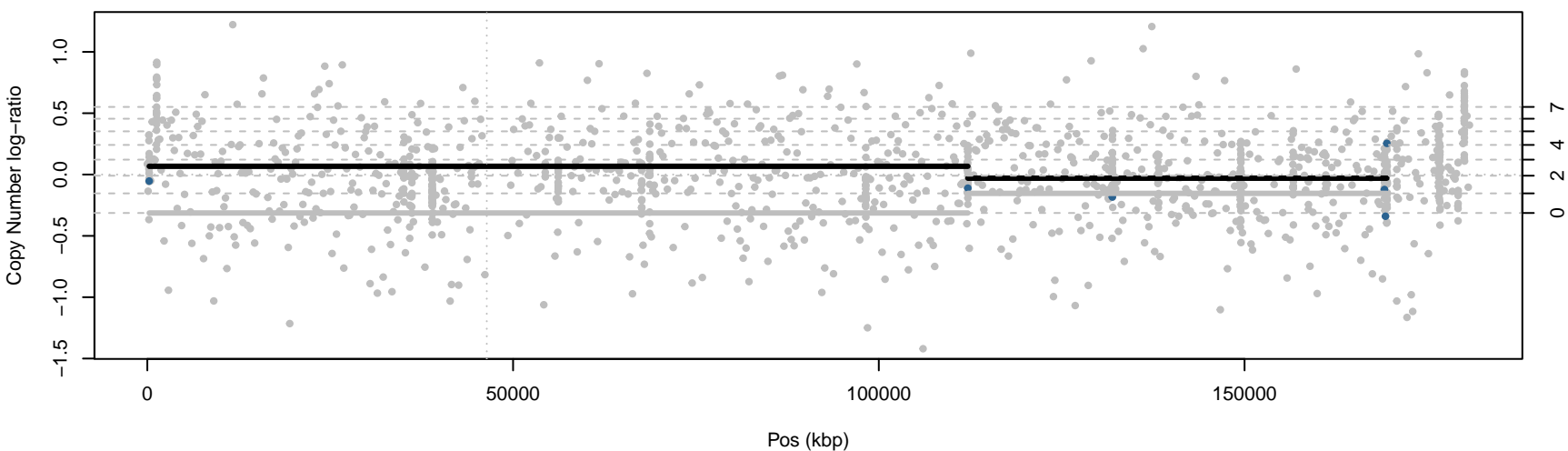
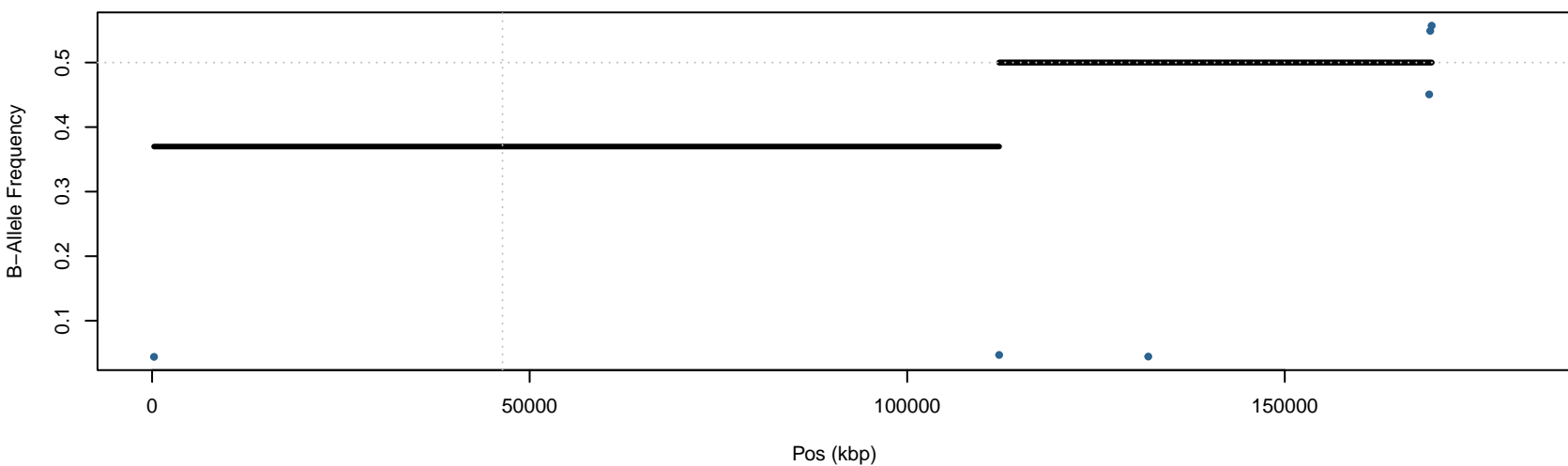
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 2



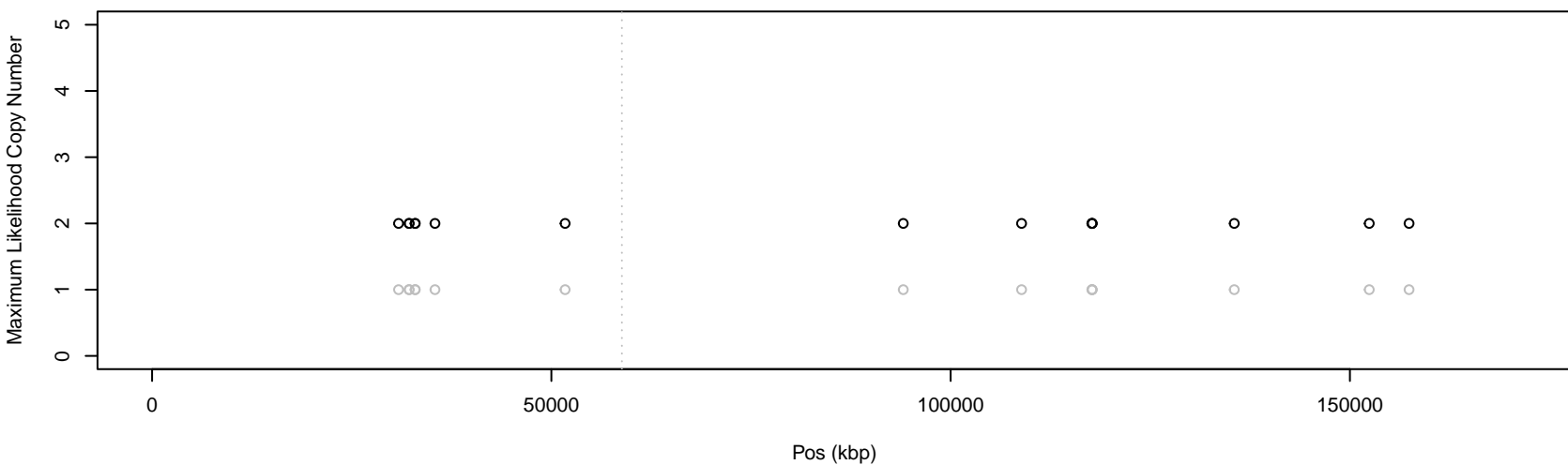
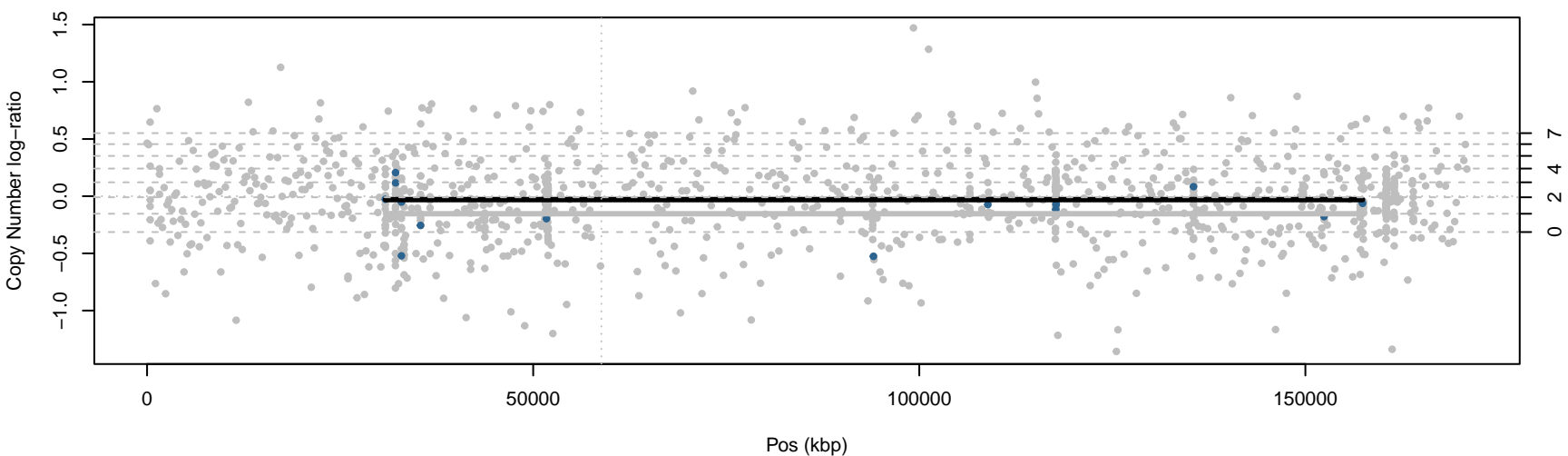
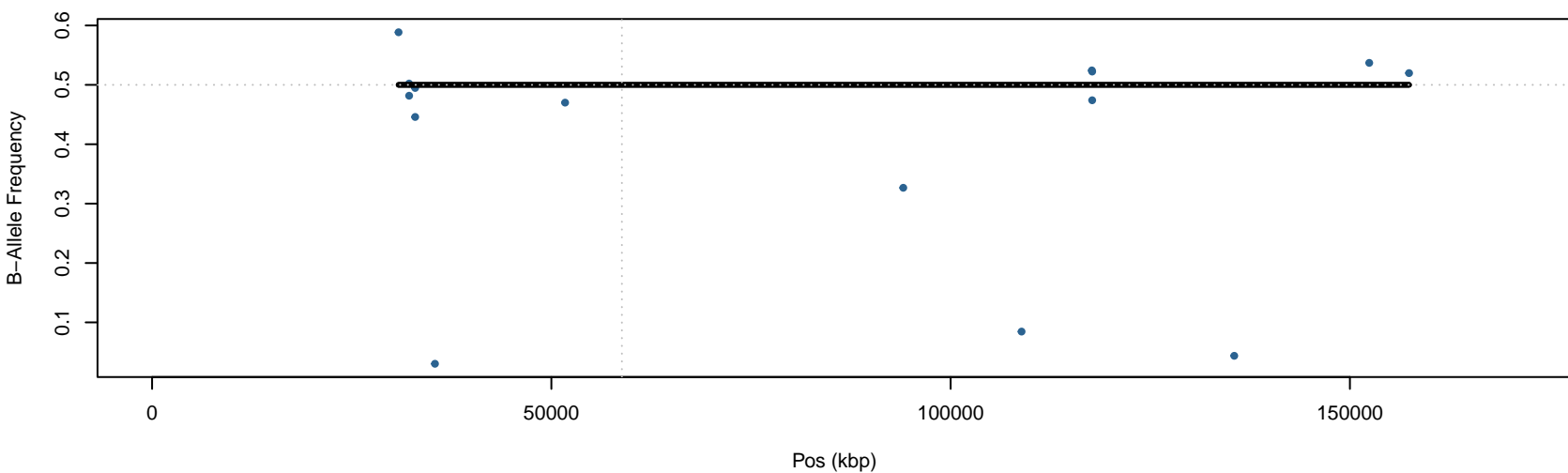
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 3



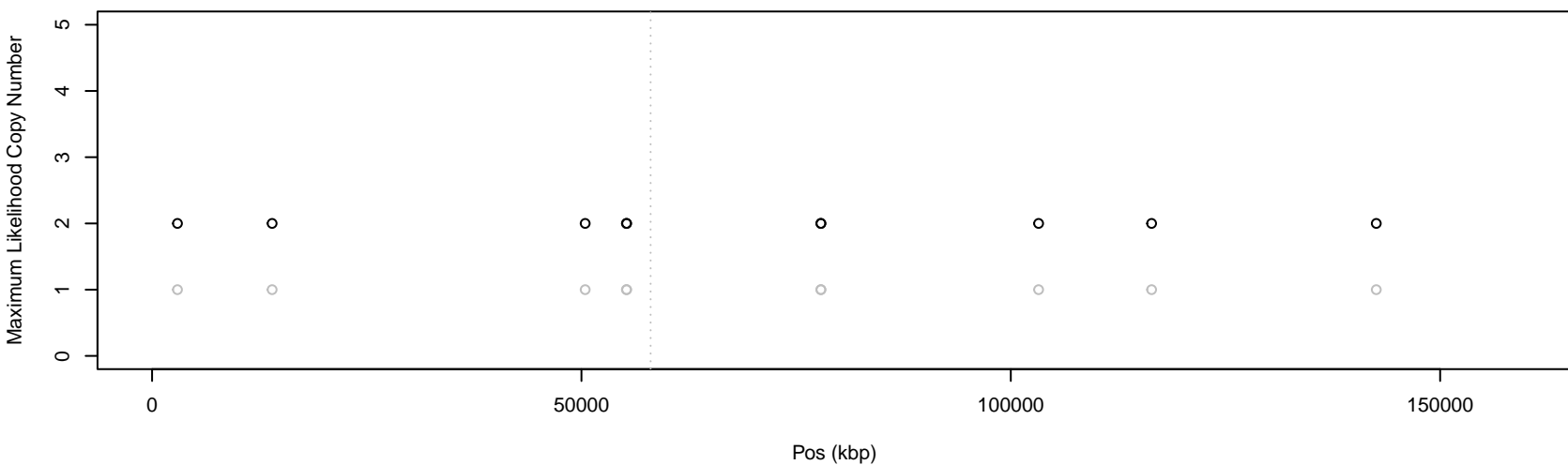
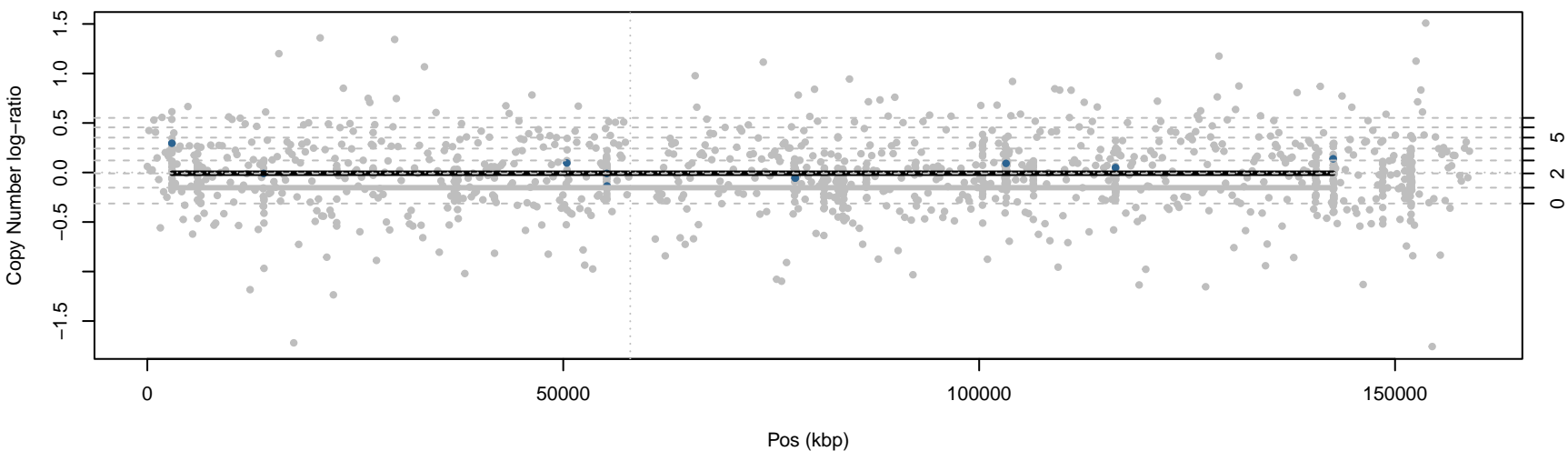
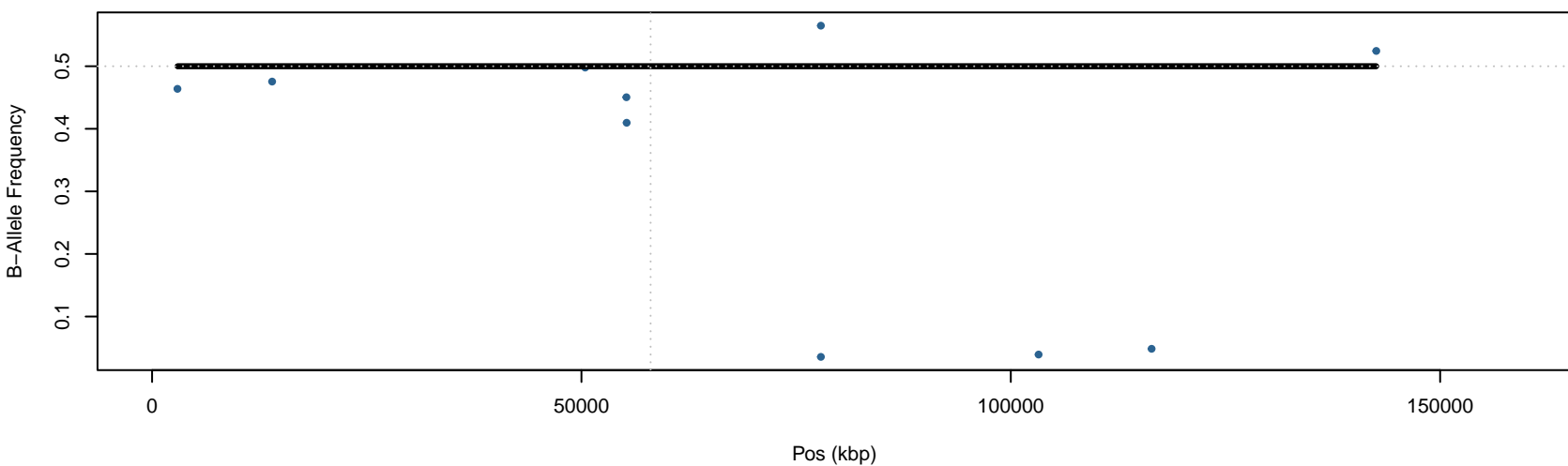
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 5



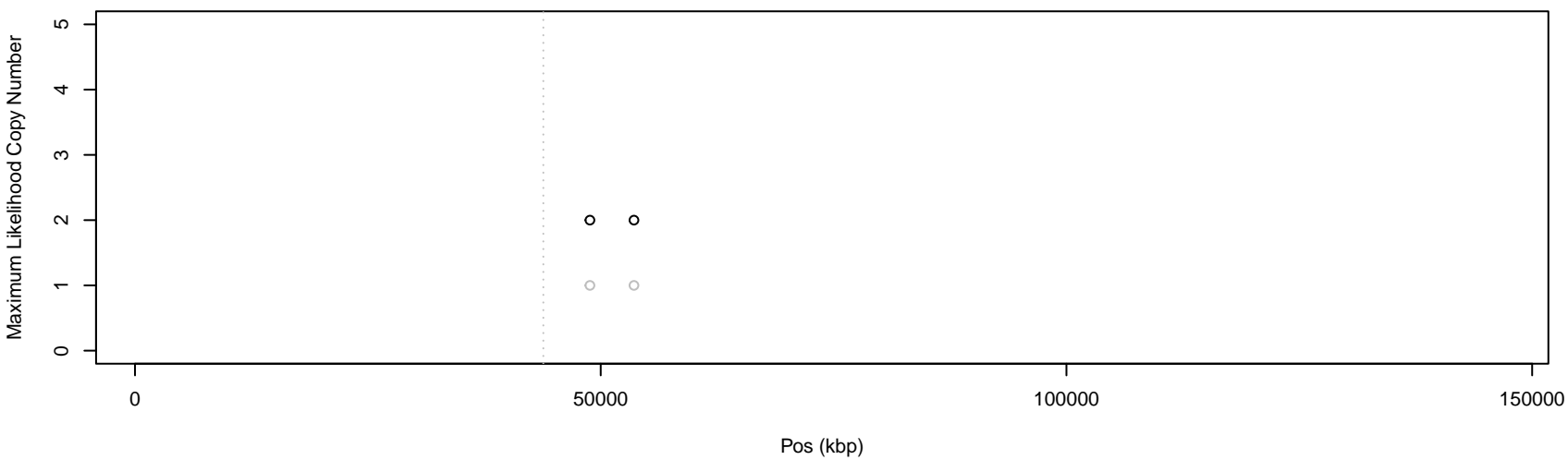
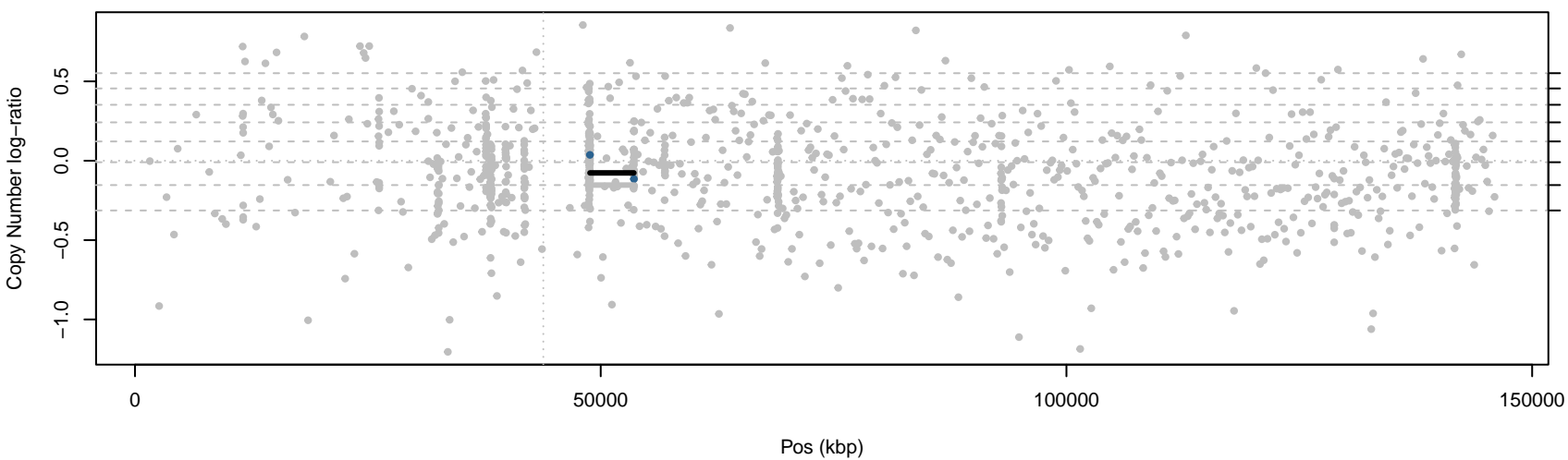
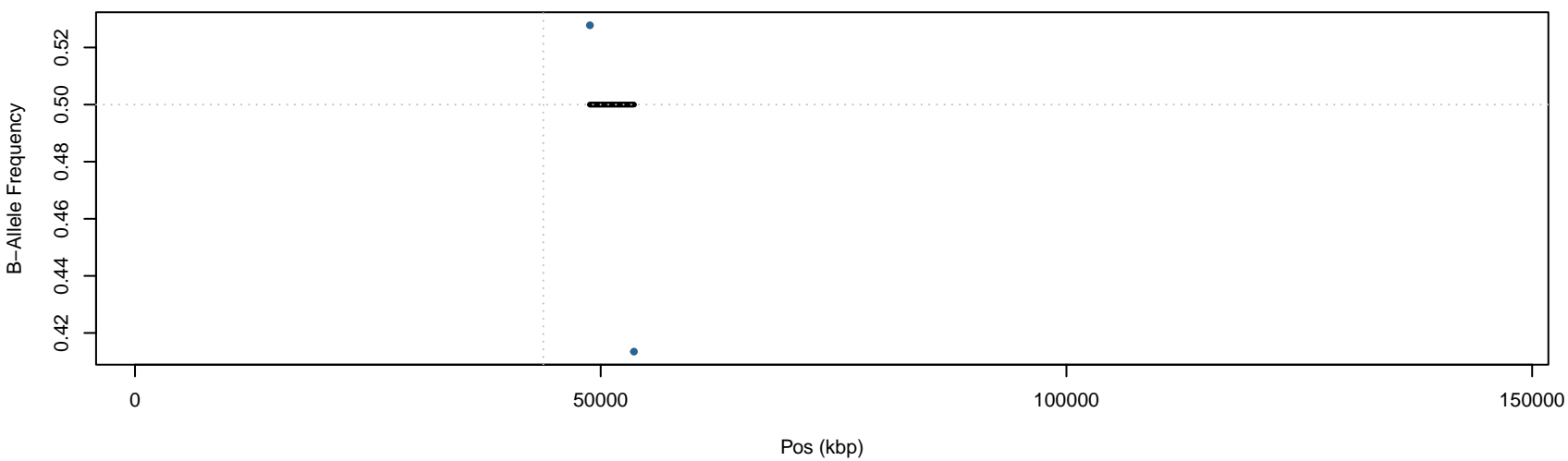
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 6



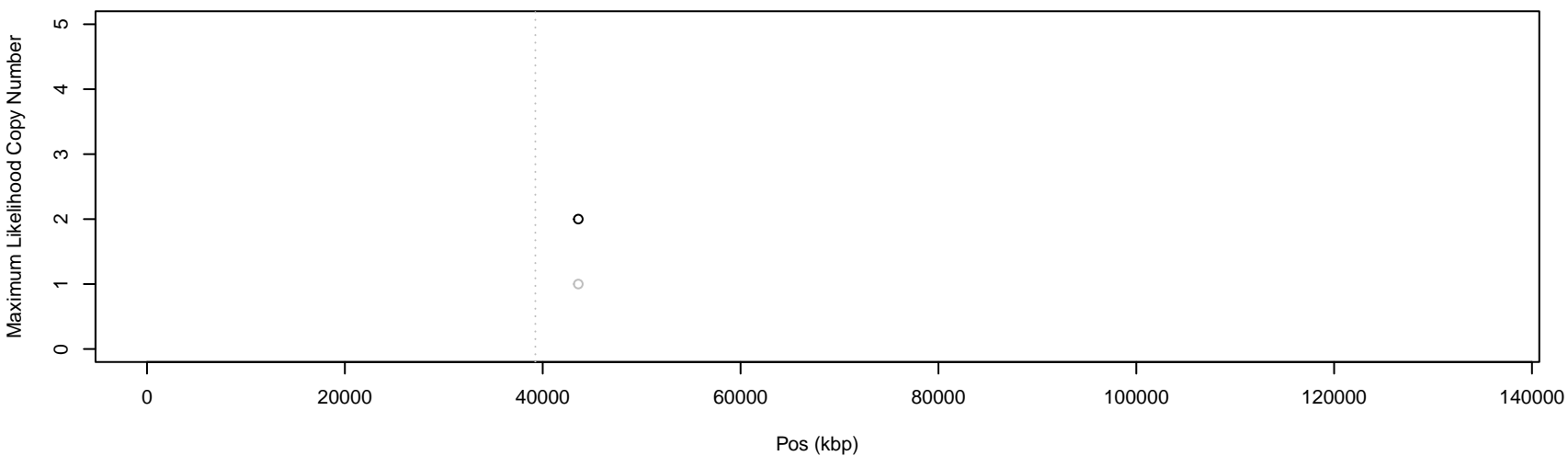
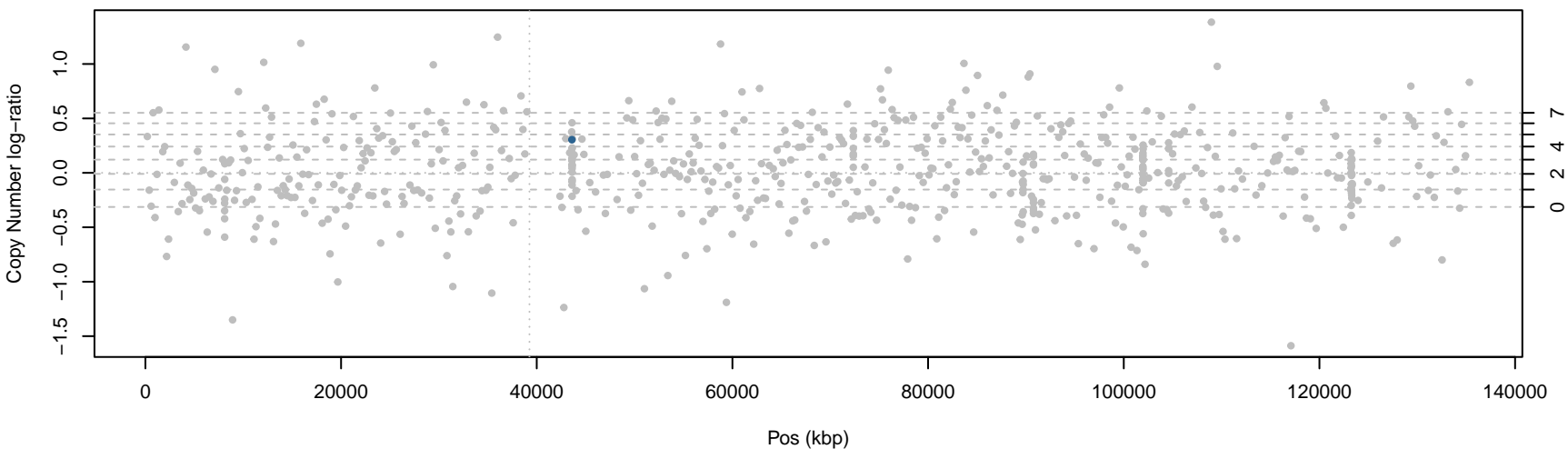
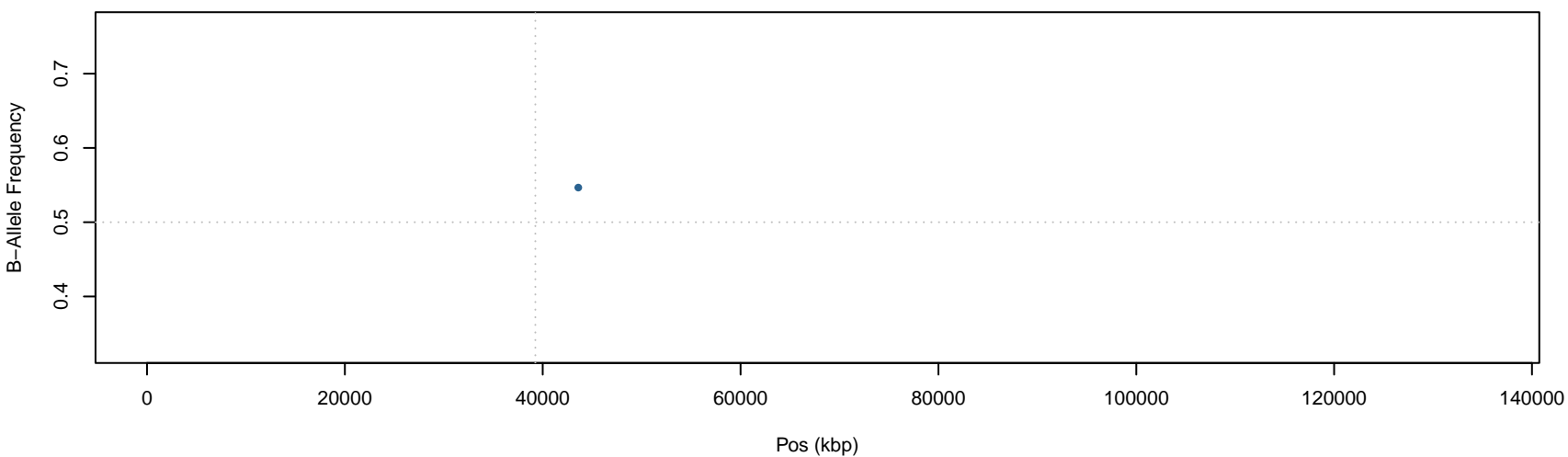
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 7



Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 8

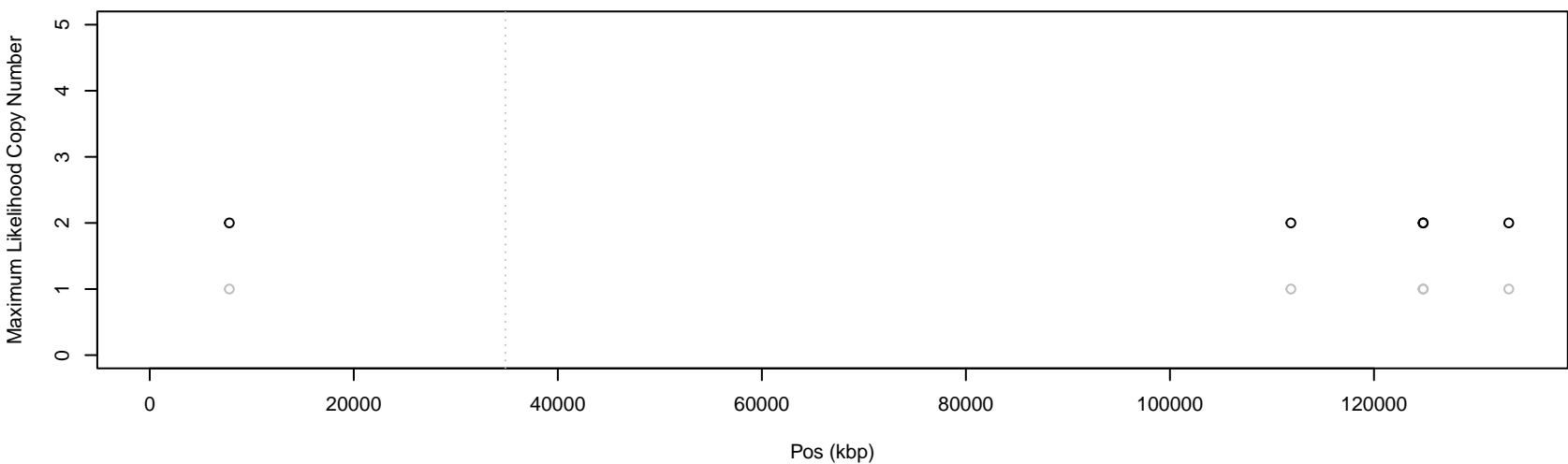
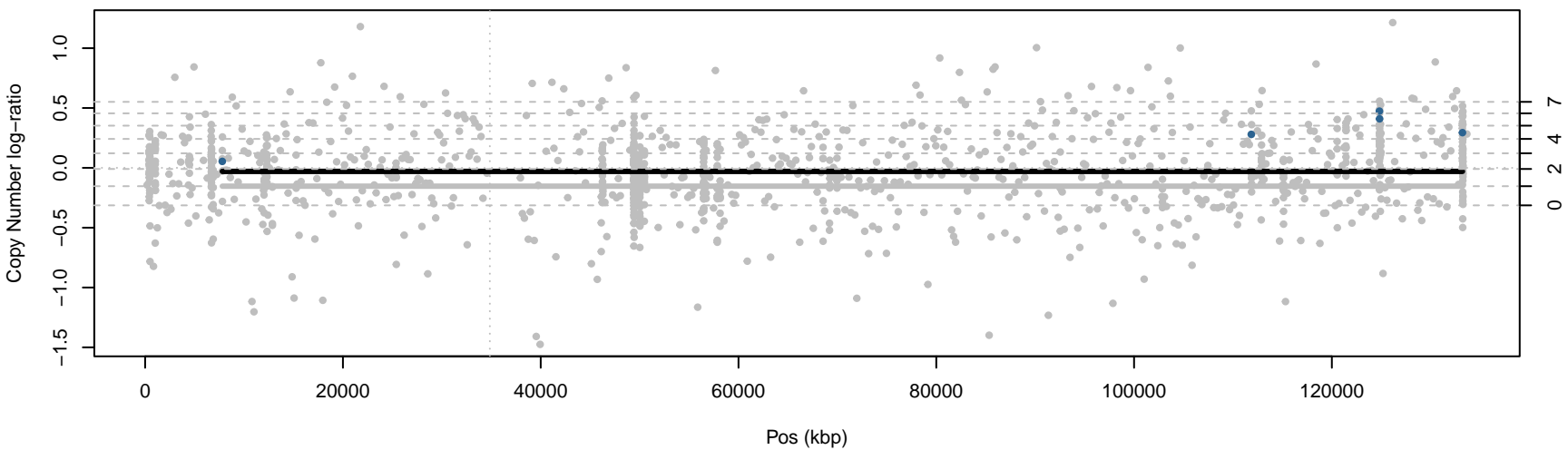
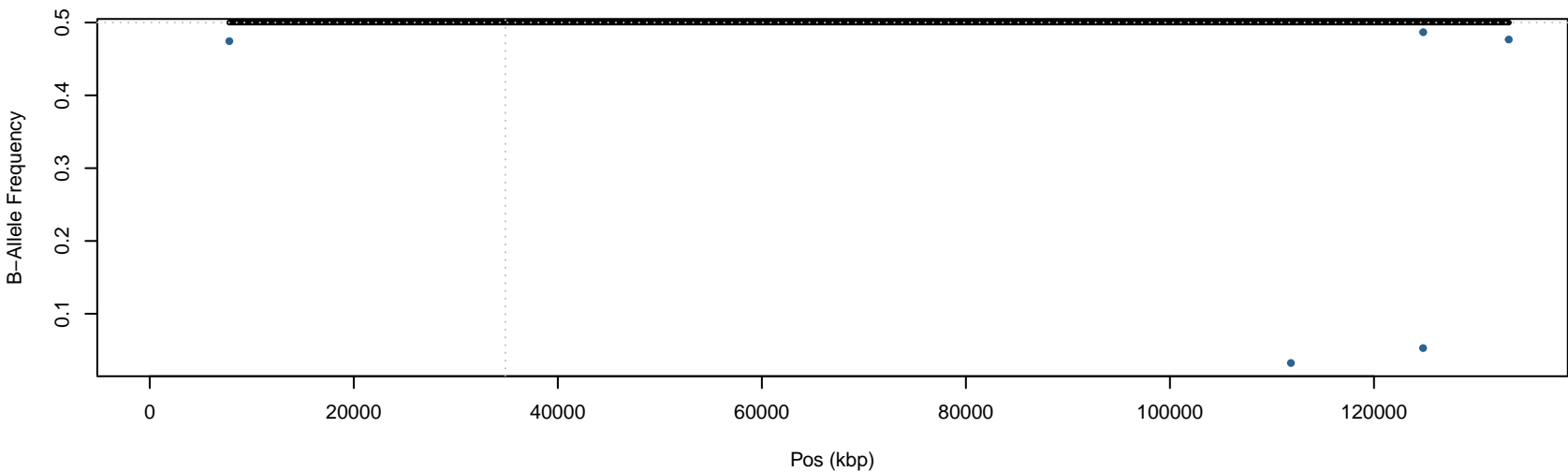


Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 10

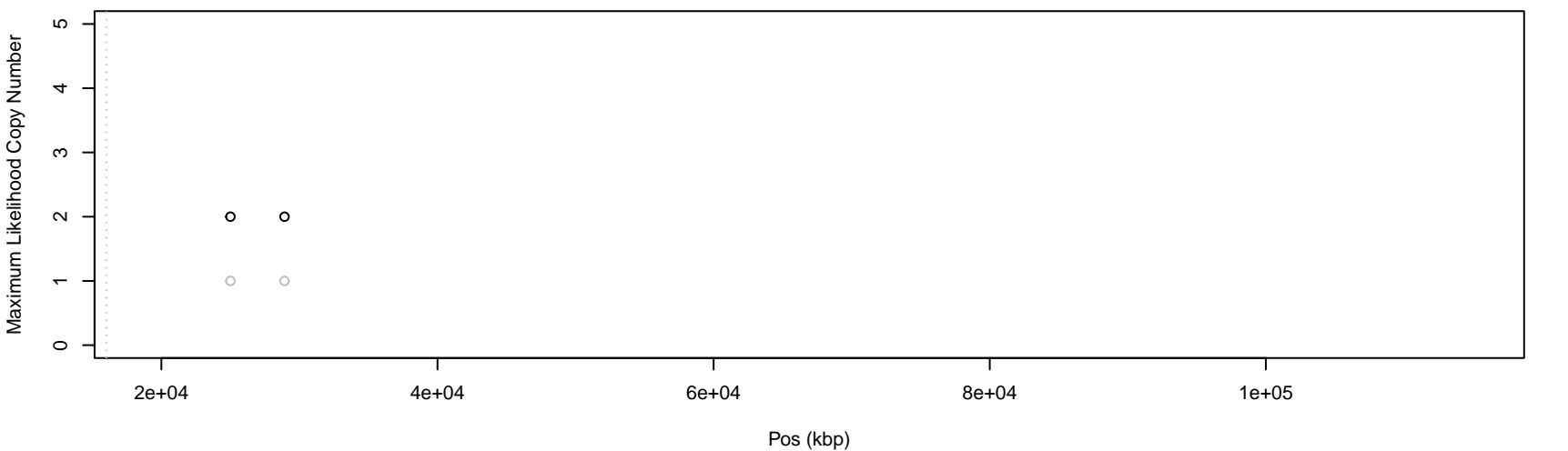
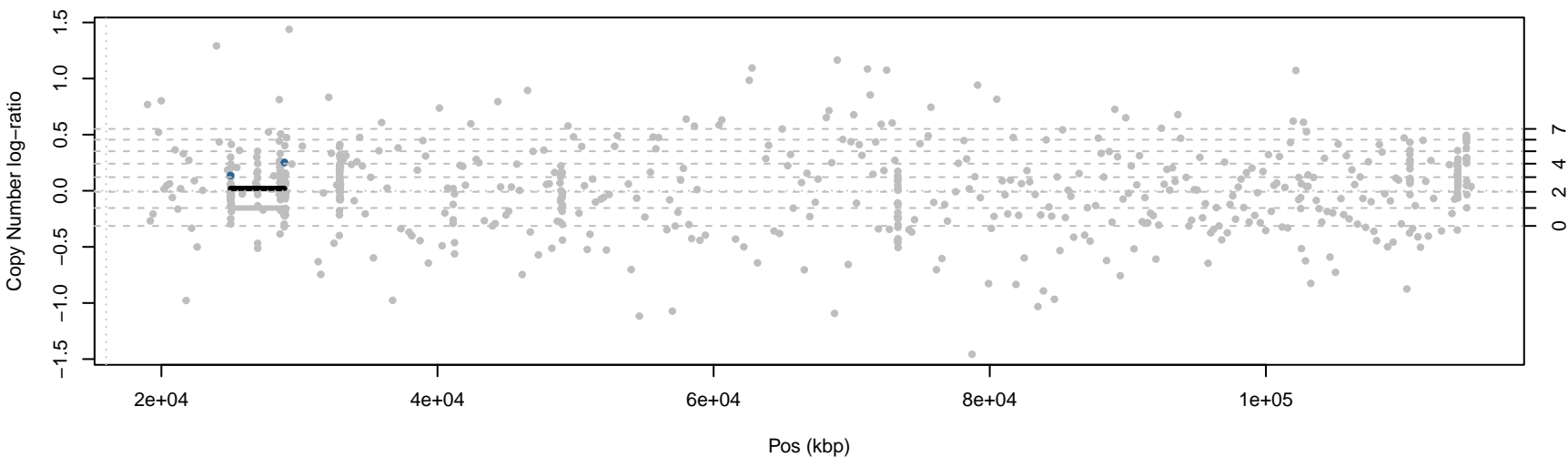
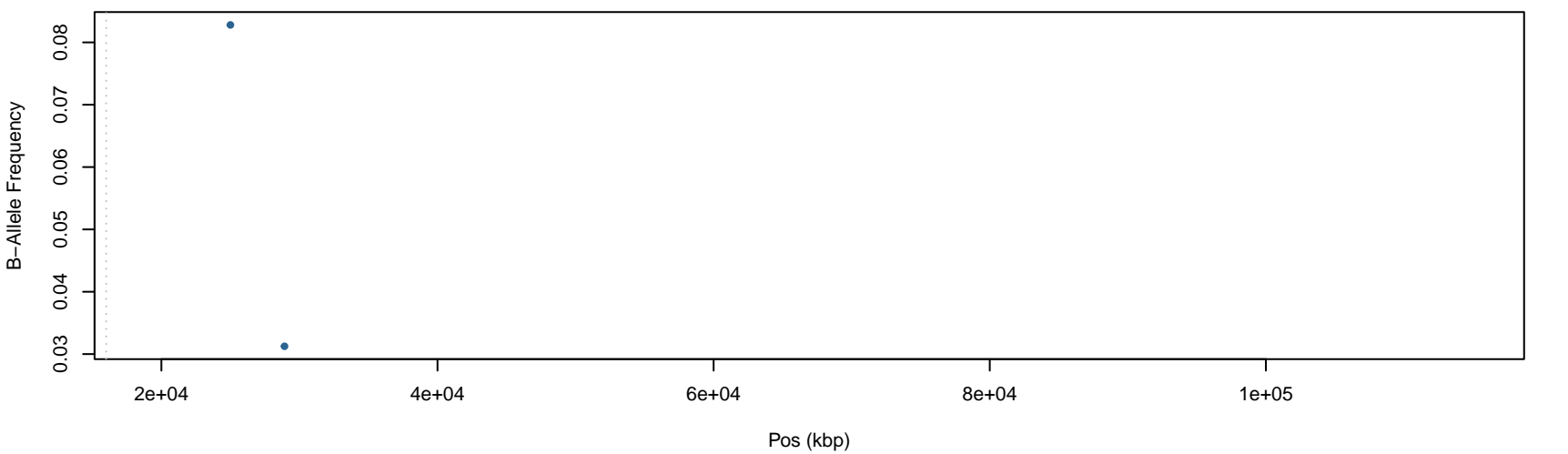




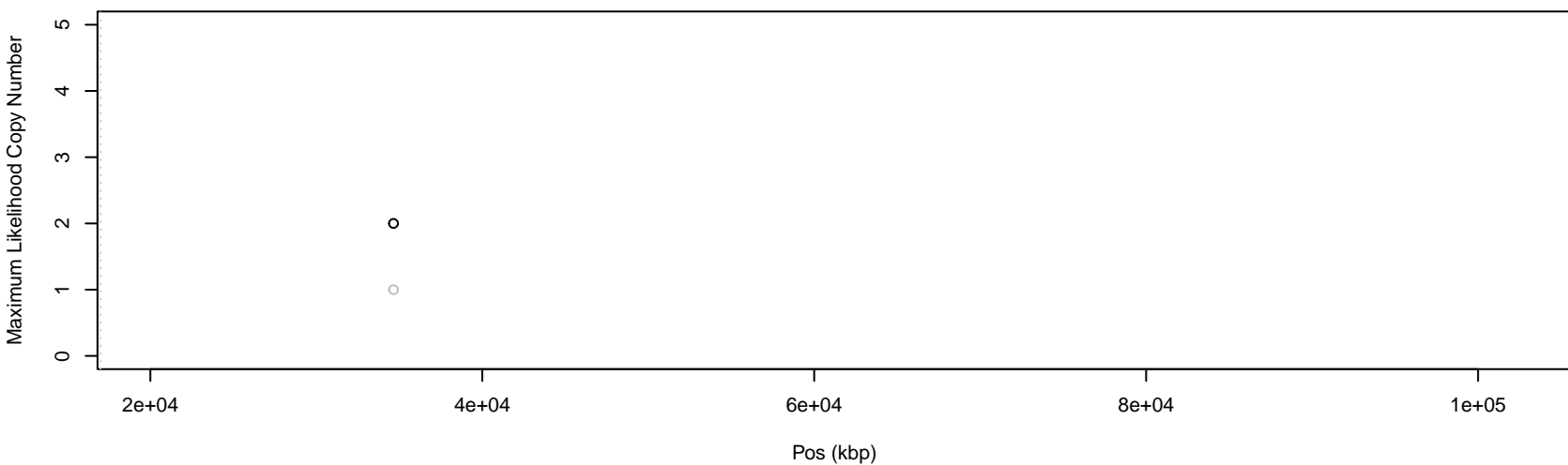
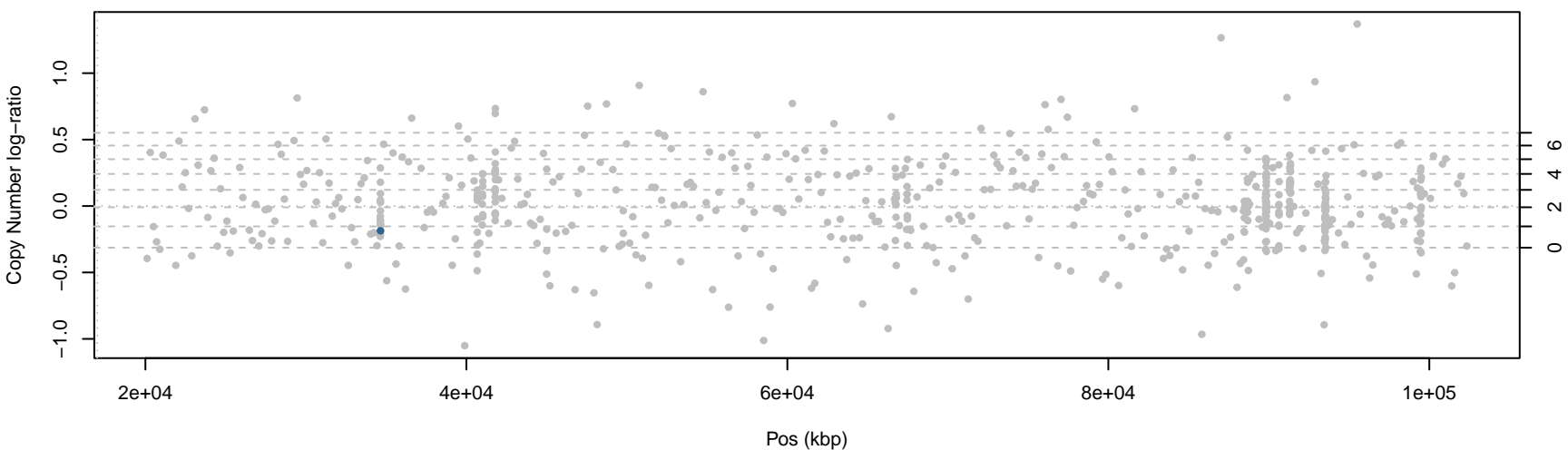
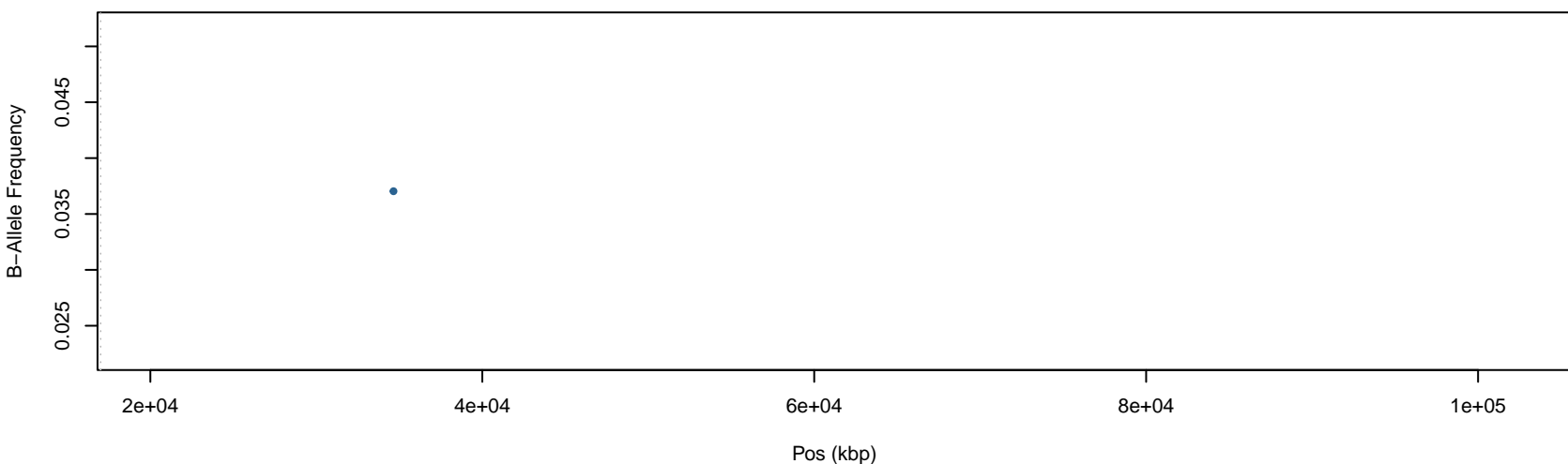
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 12



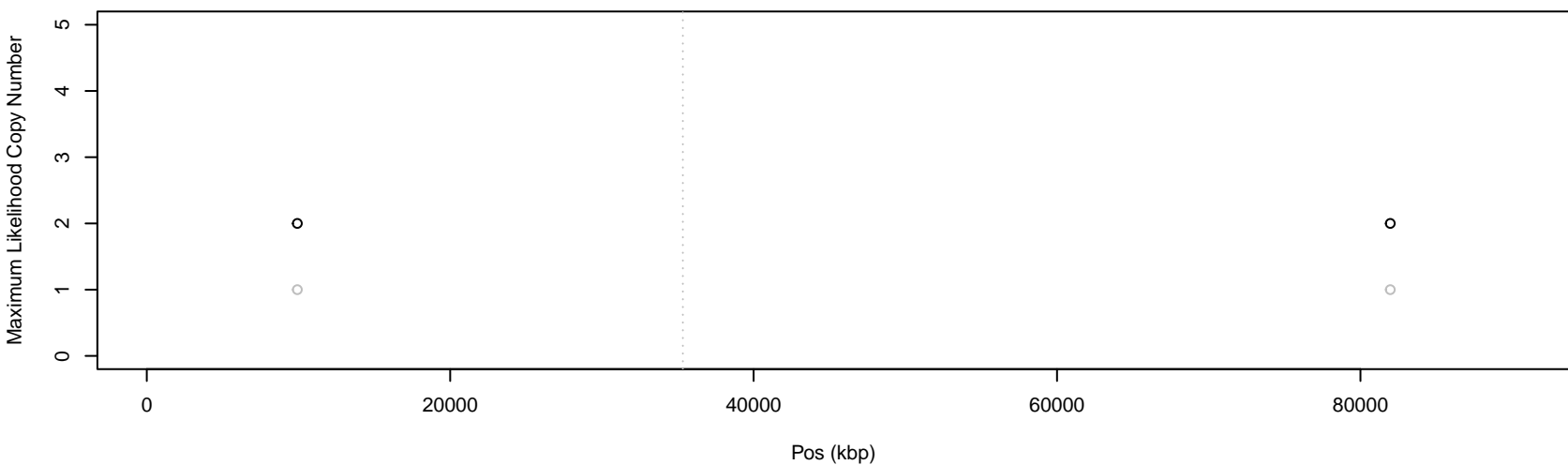
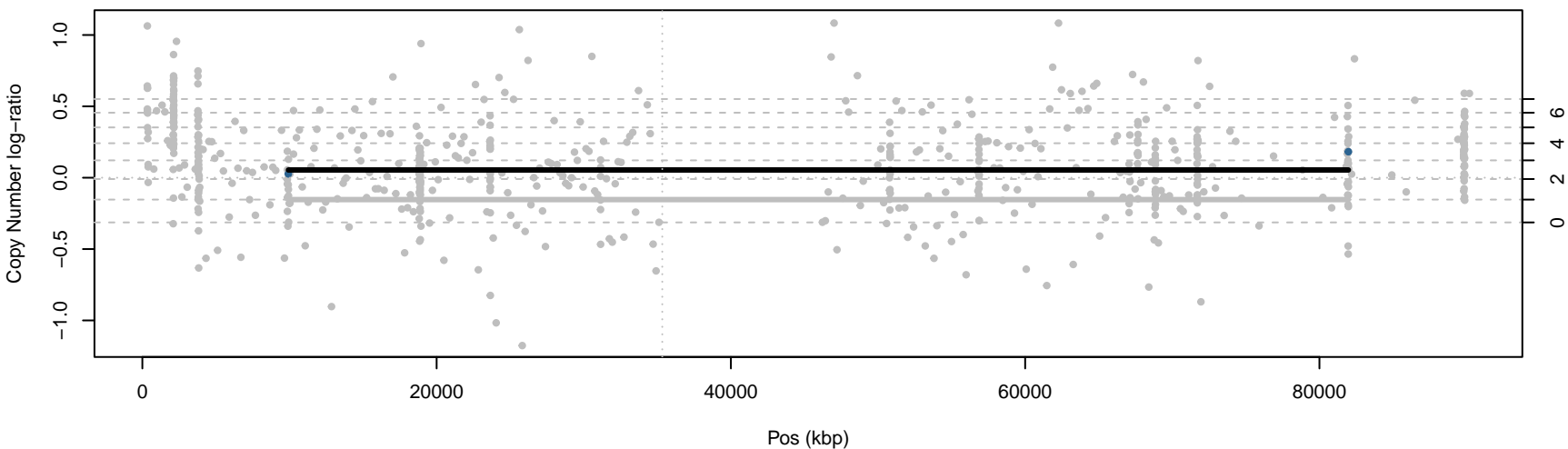
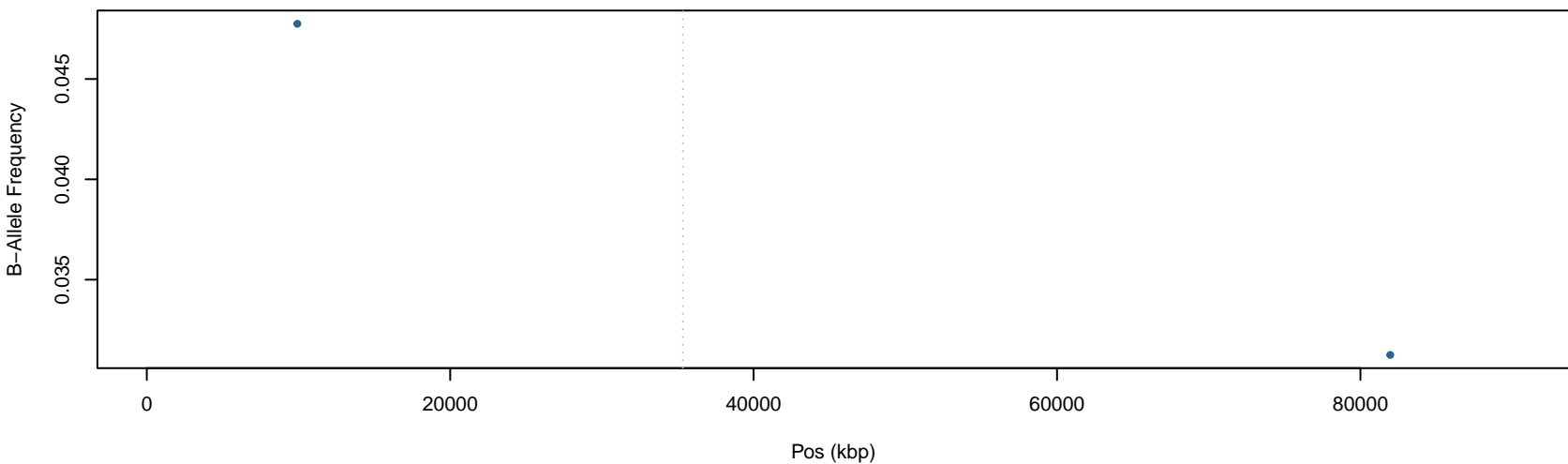
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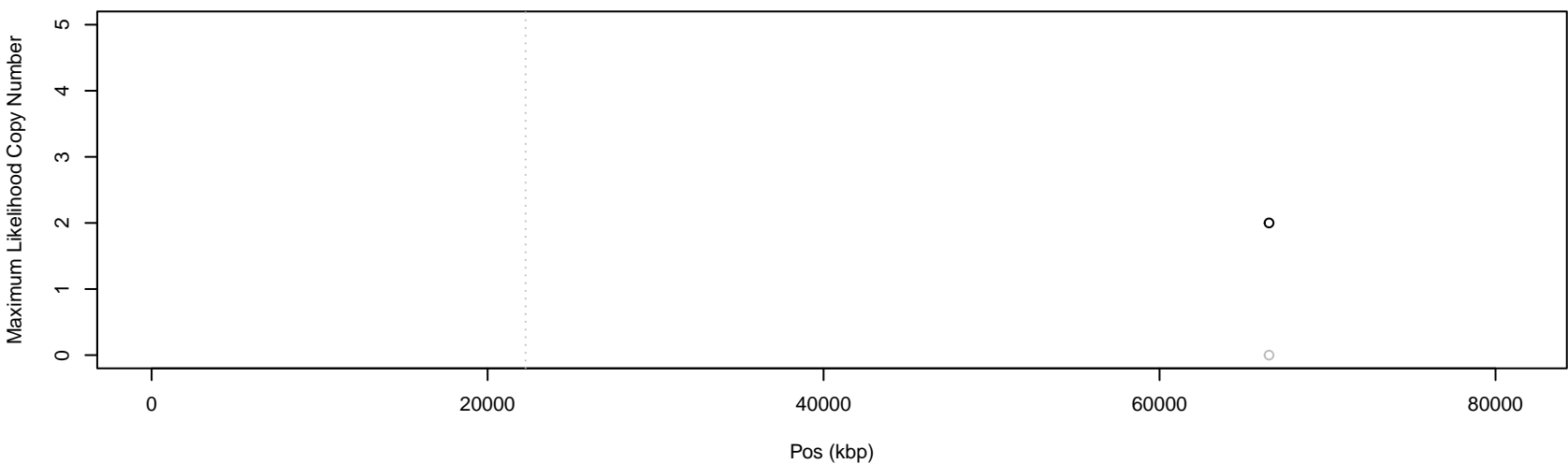
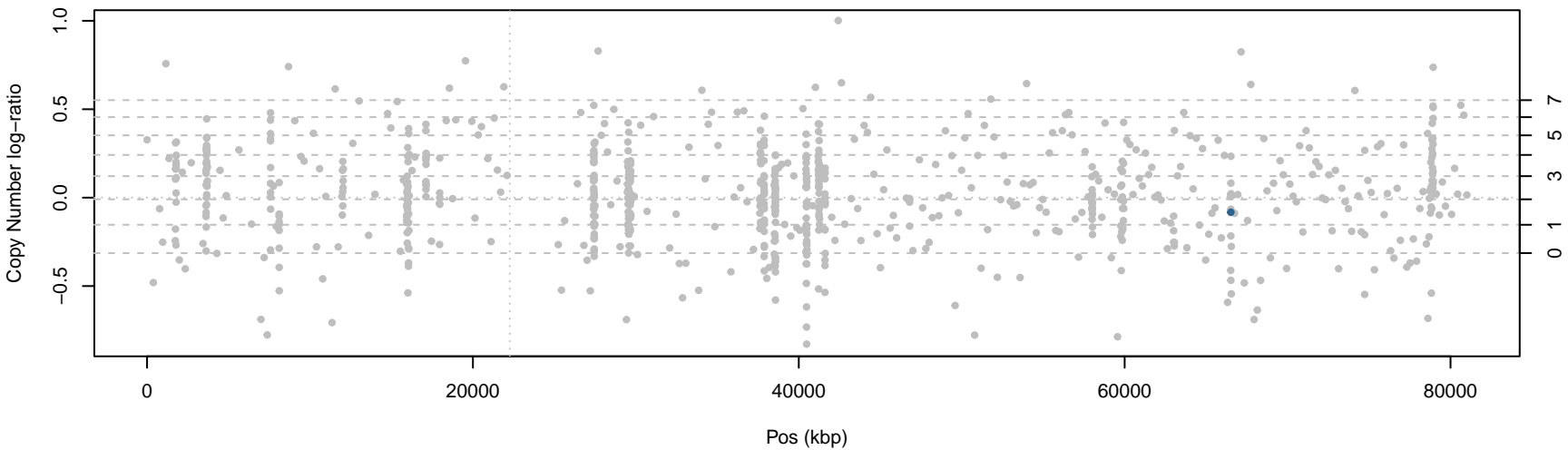
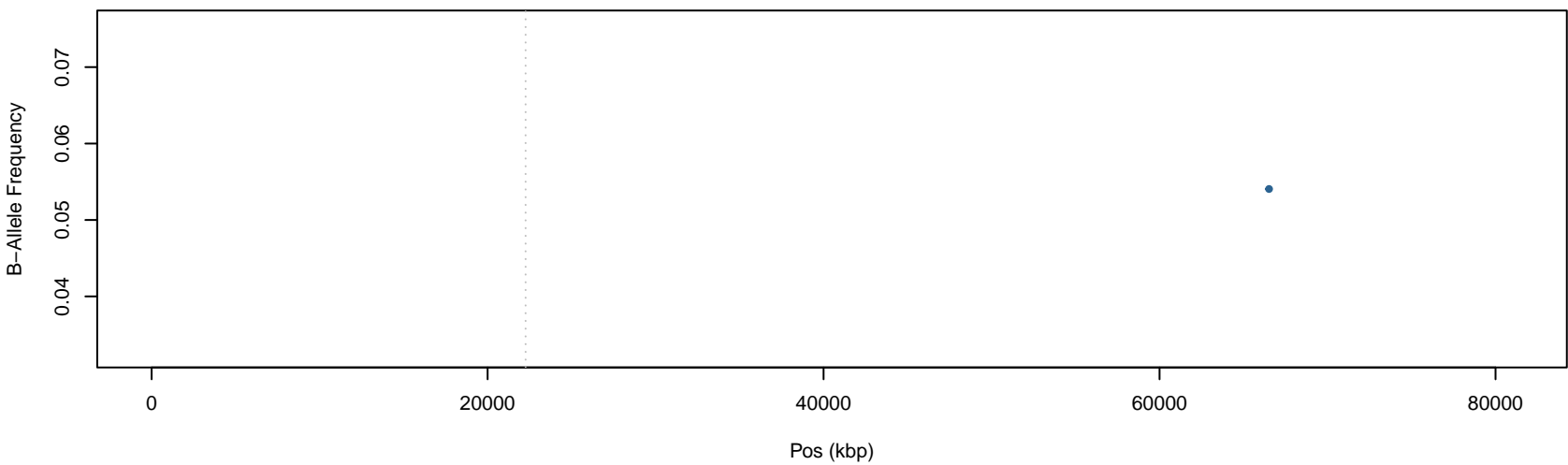
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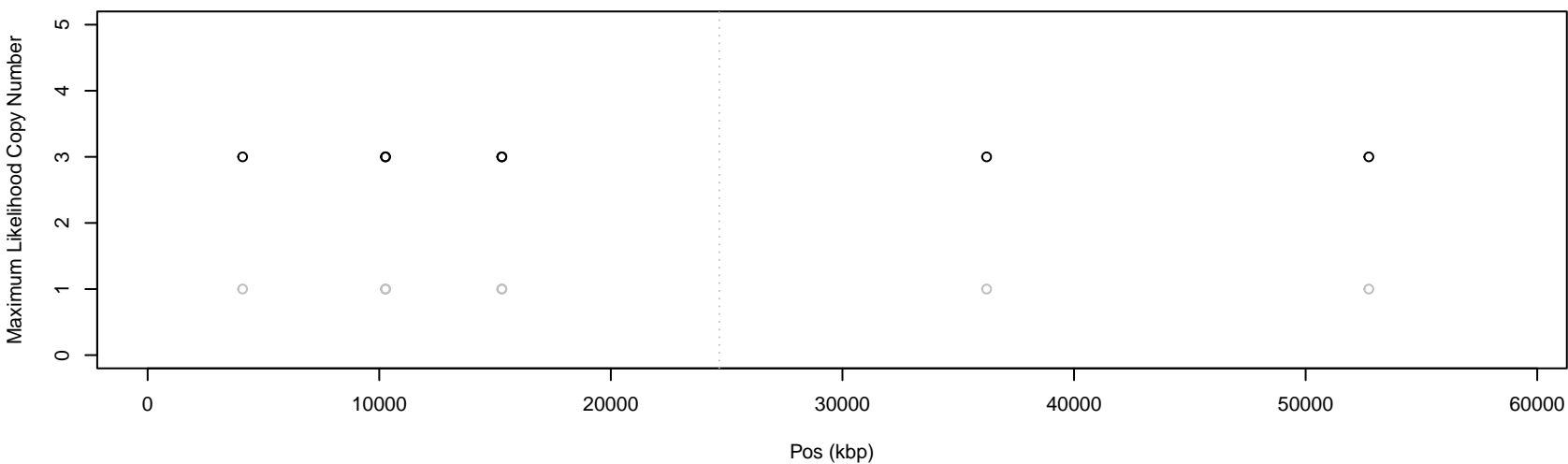
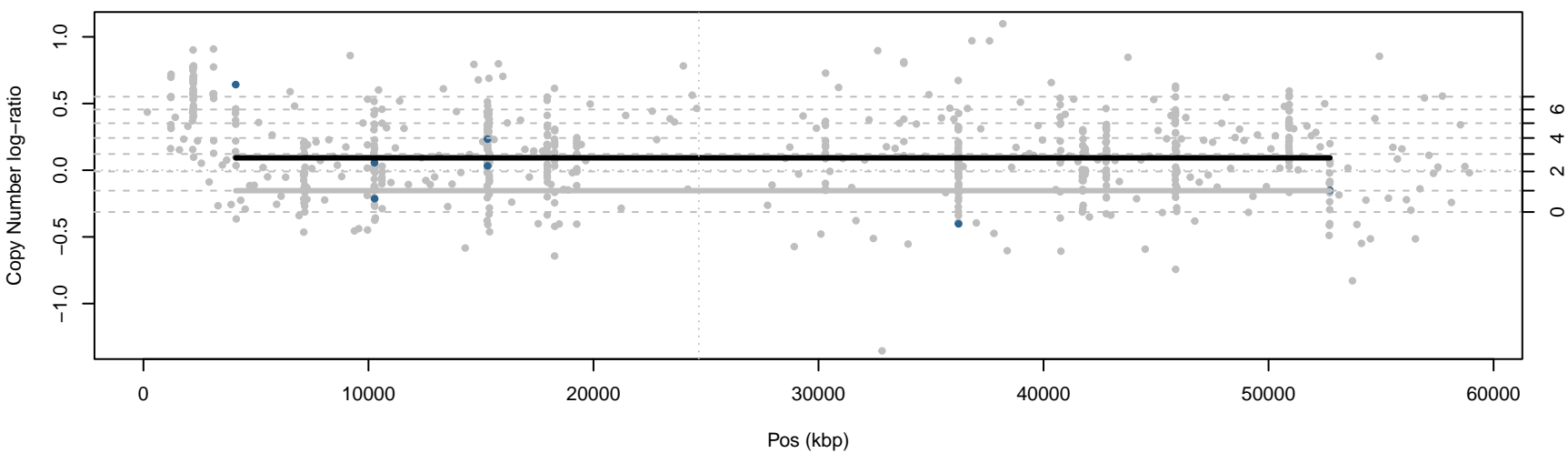
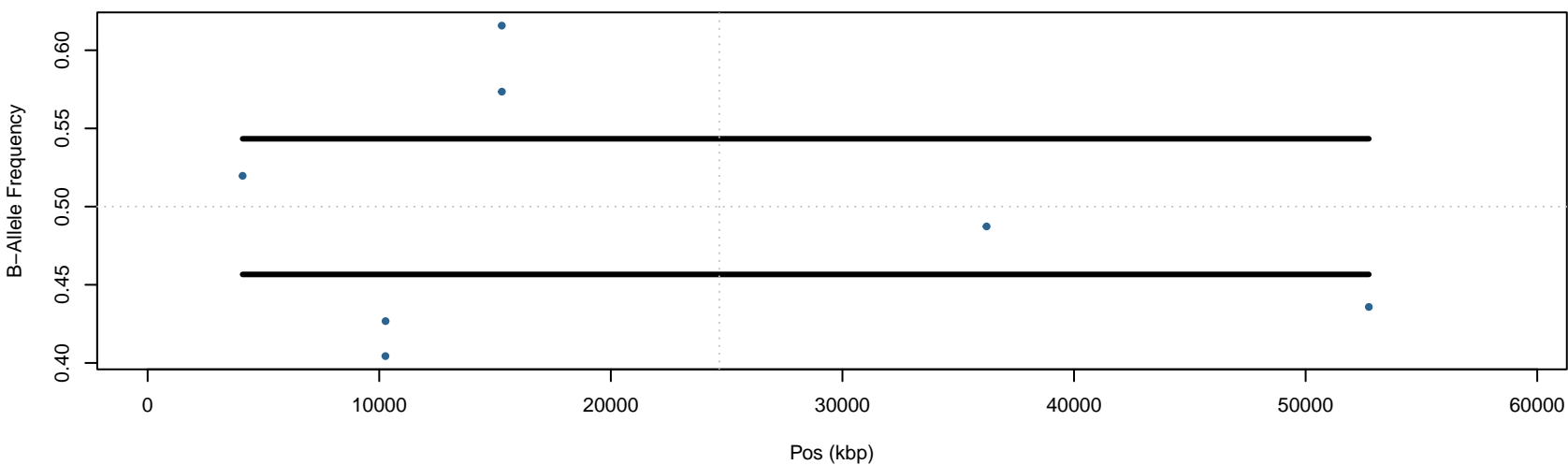
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 16



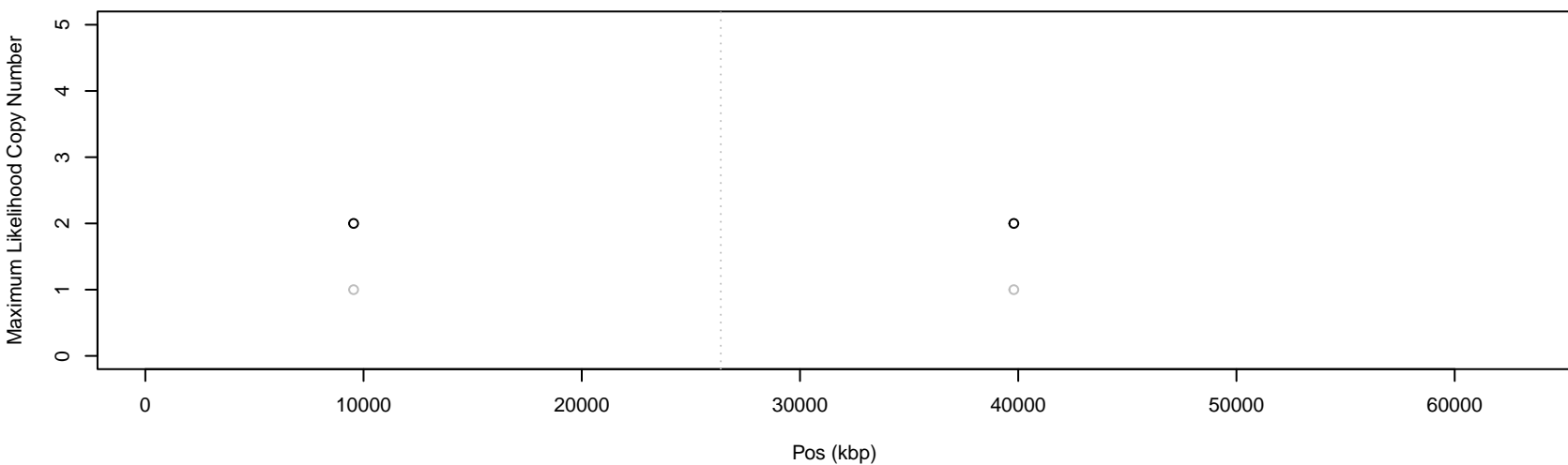
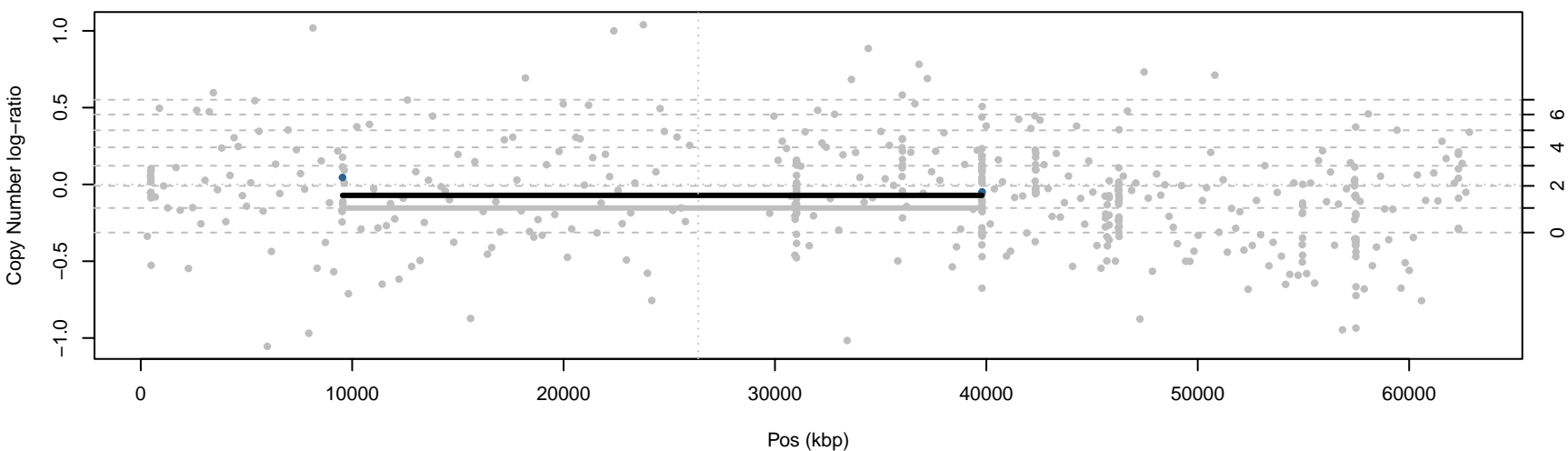
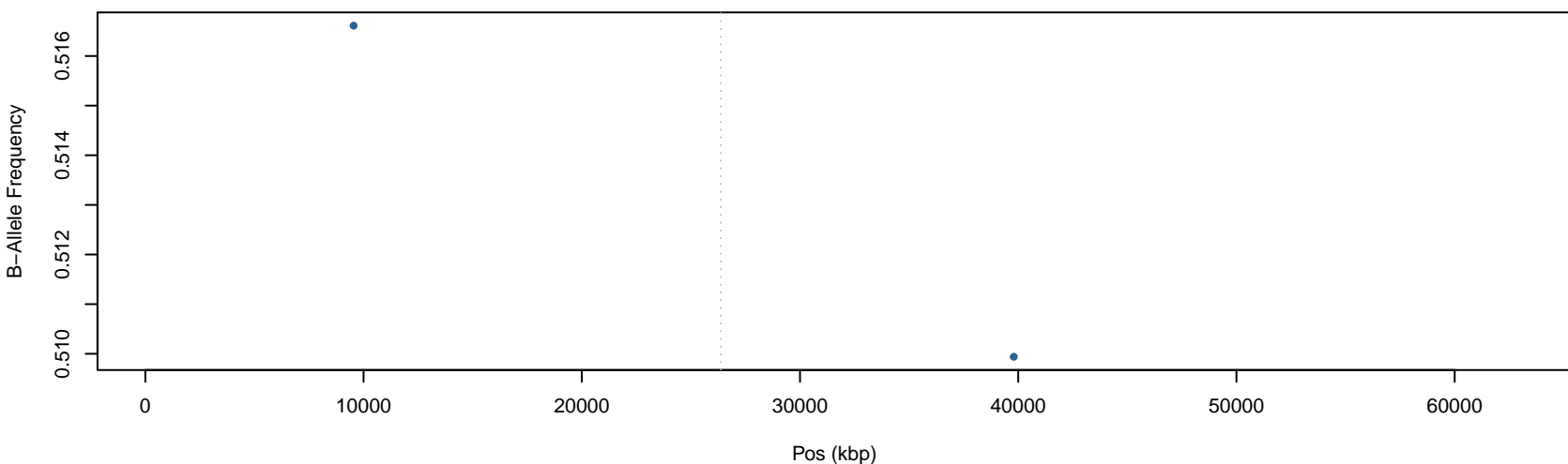
Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 17



Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 19



Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 20



Purity: 0.19 Tumor ploidy: 2.069 SNV log-likelihood: 60.5 GoF: 93.3% Mean coverage: 473;439 Chromosome: 22

