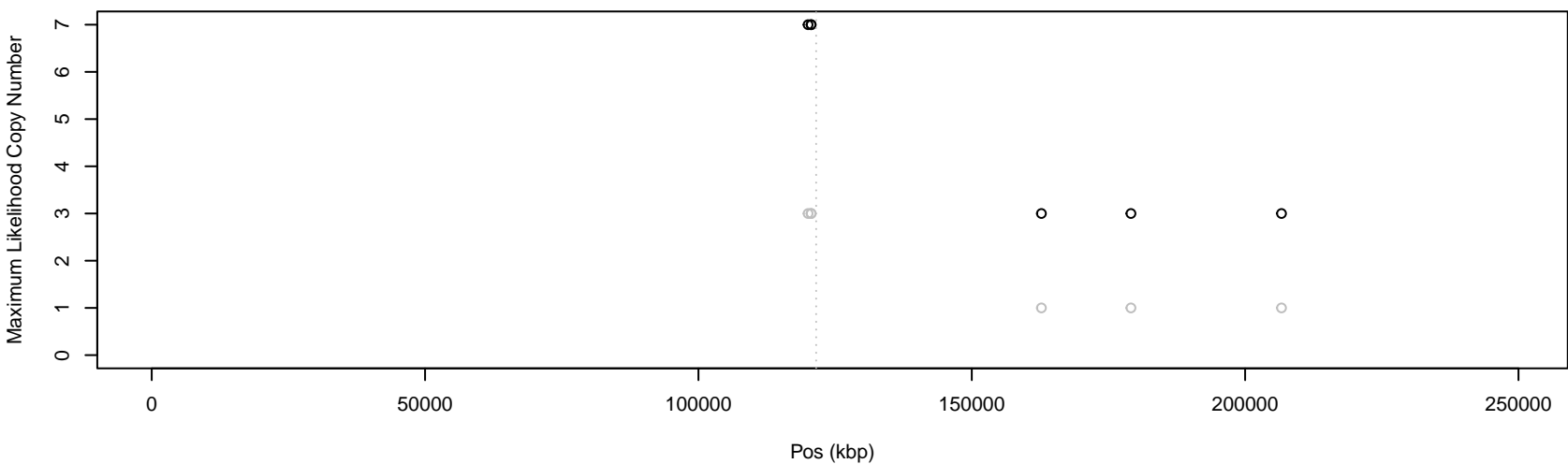
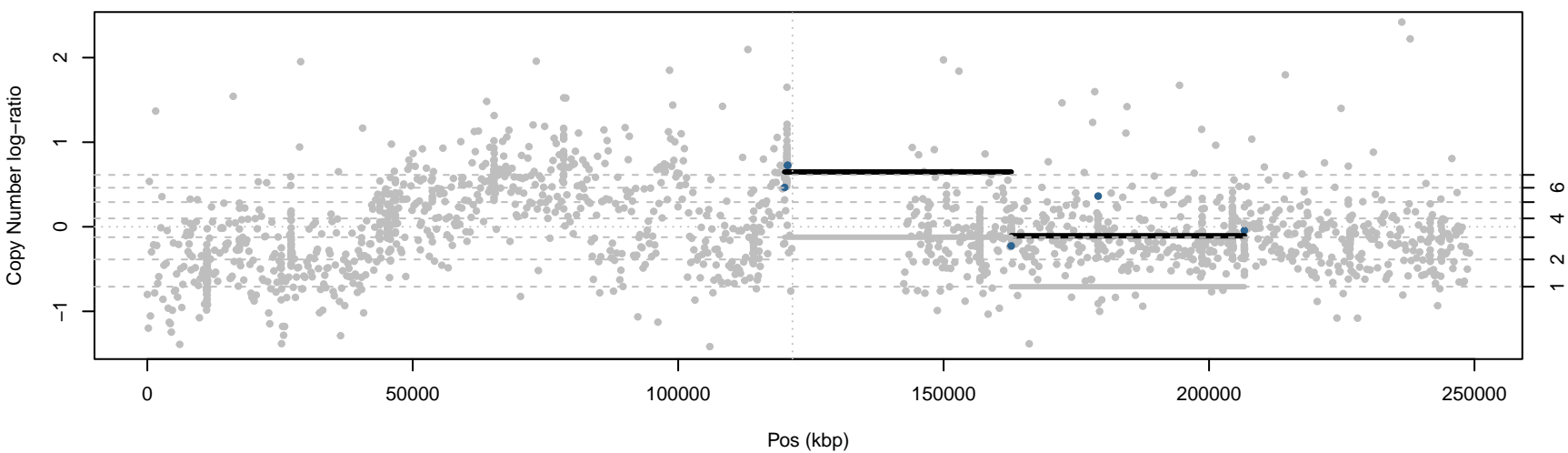
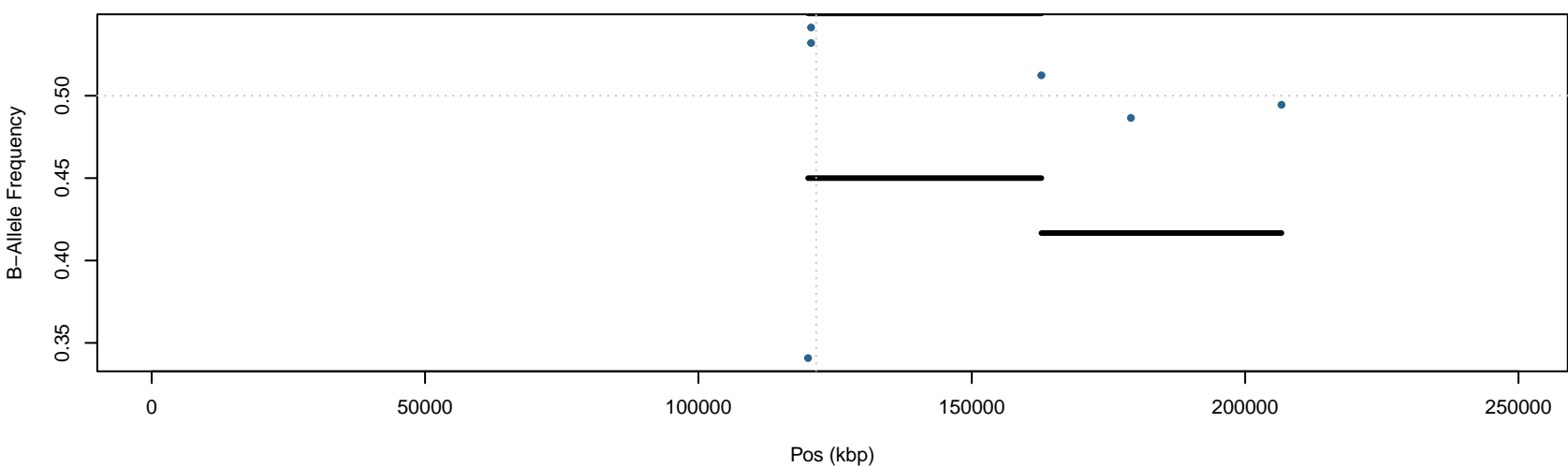
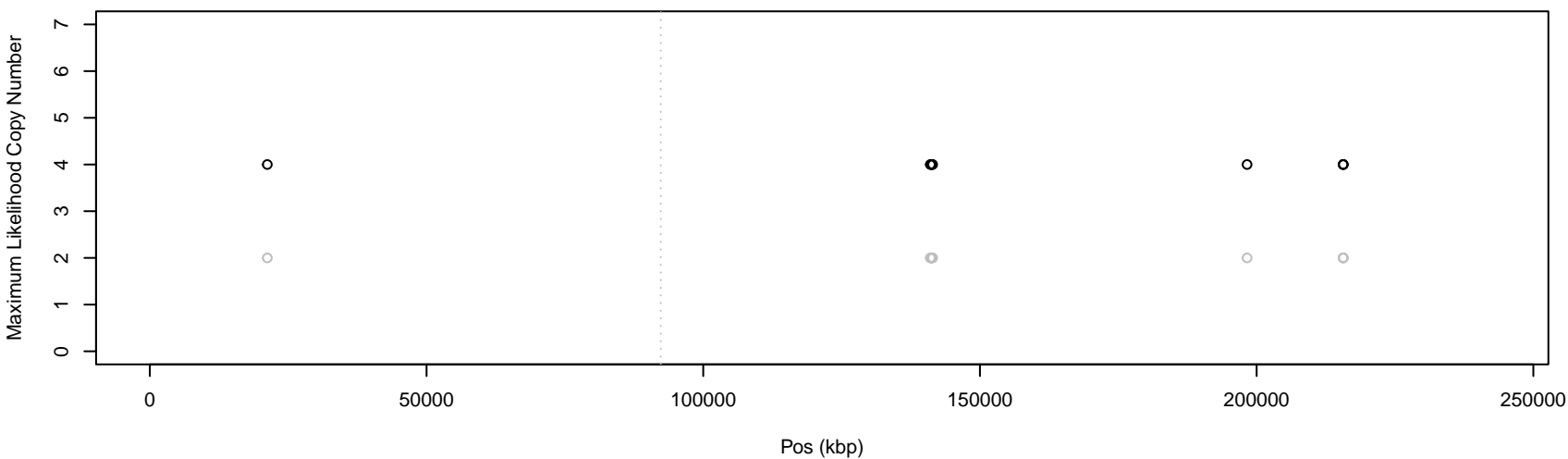
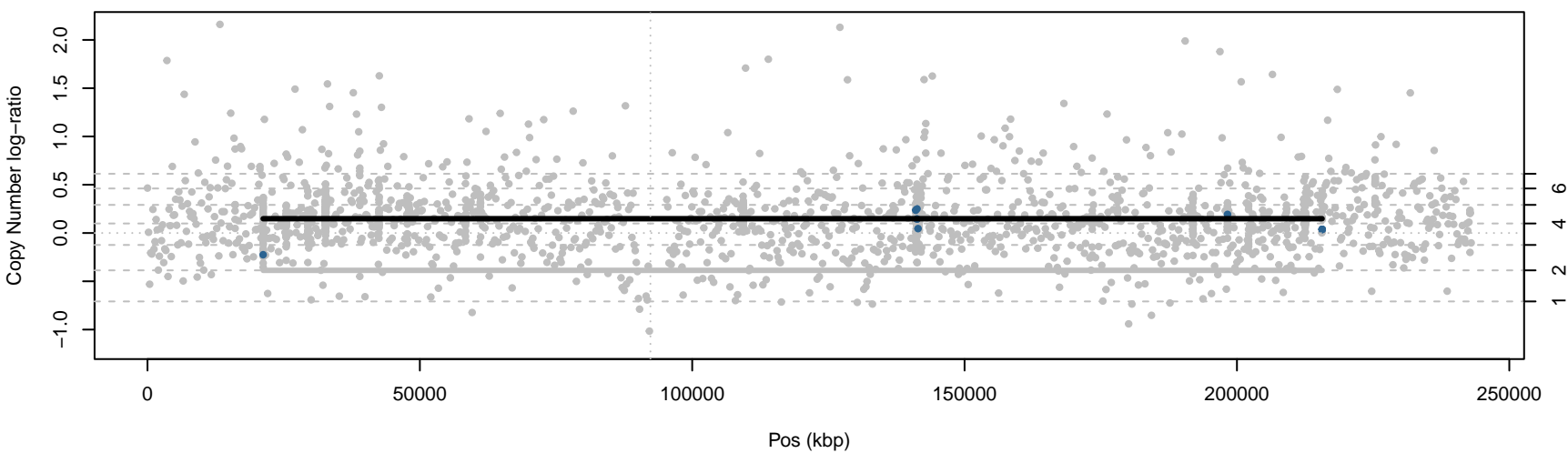
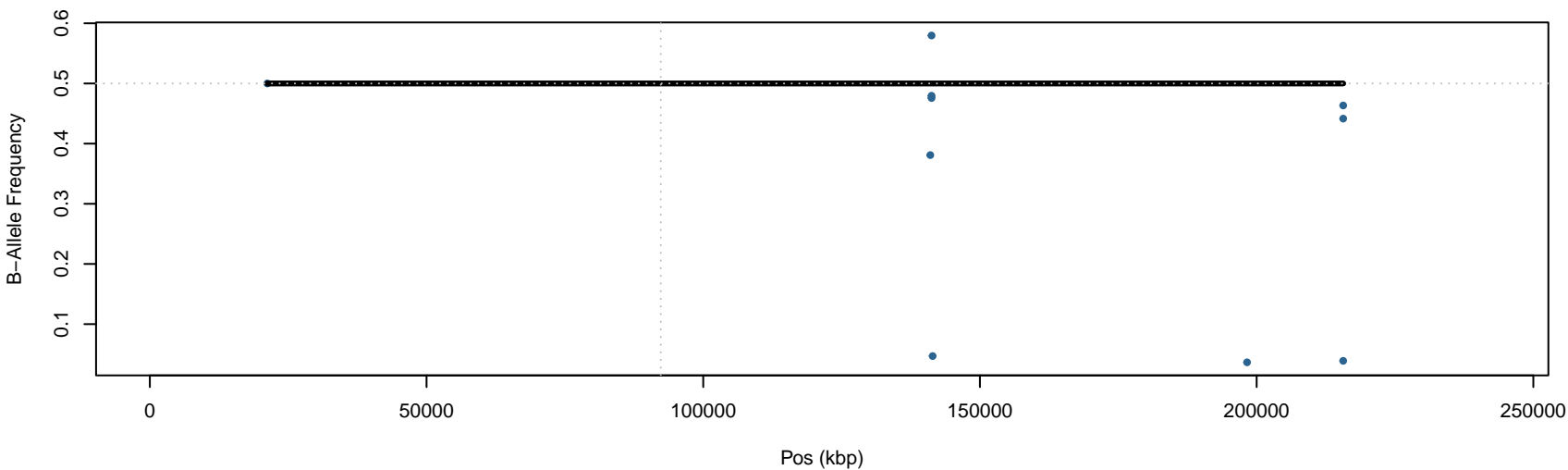


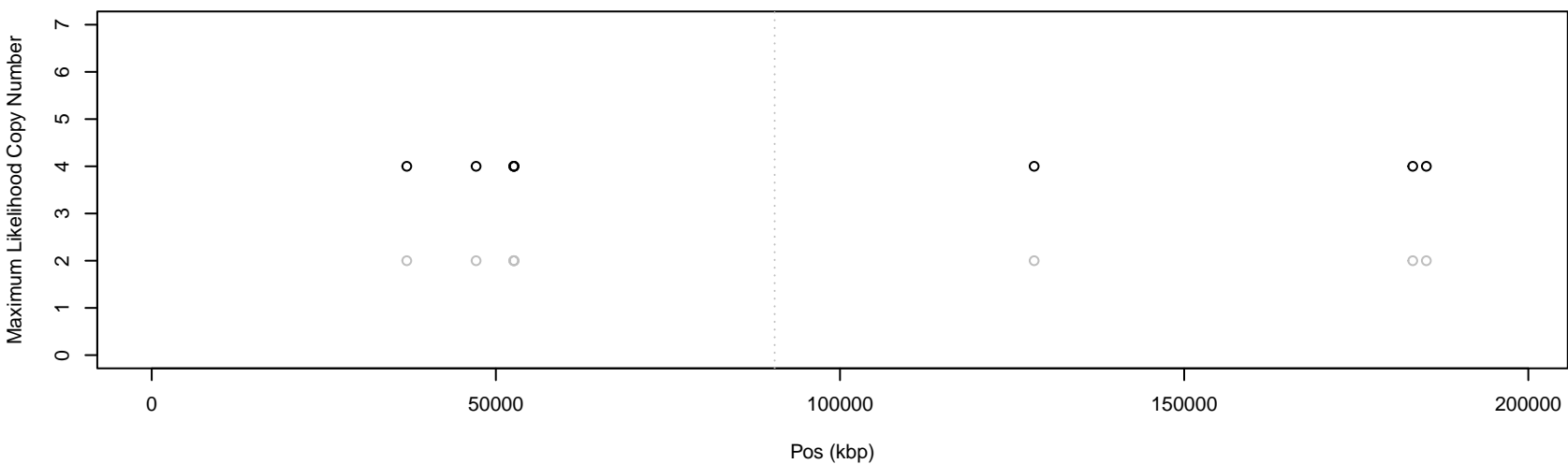
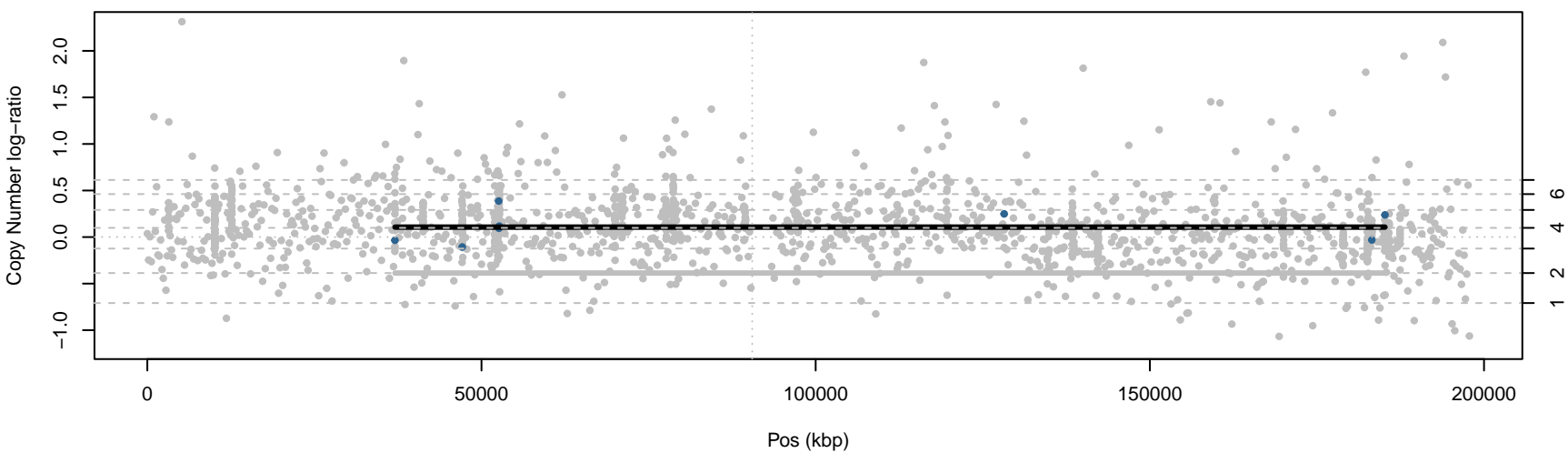
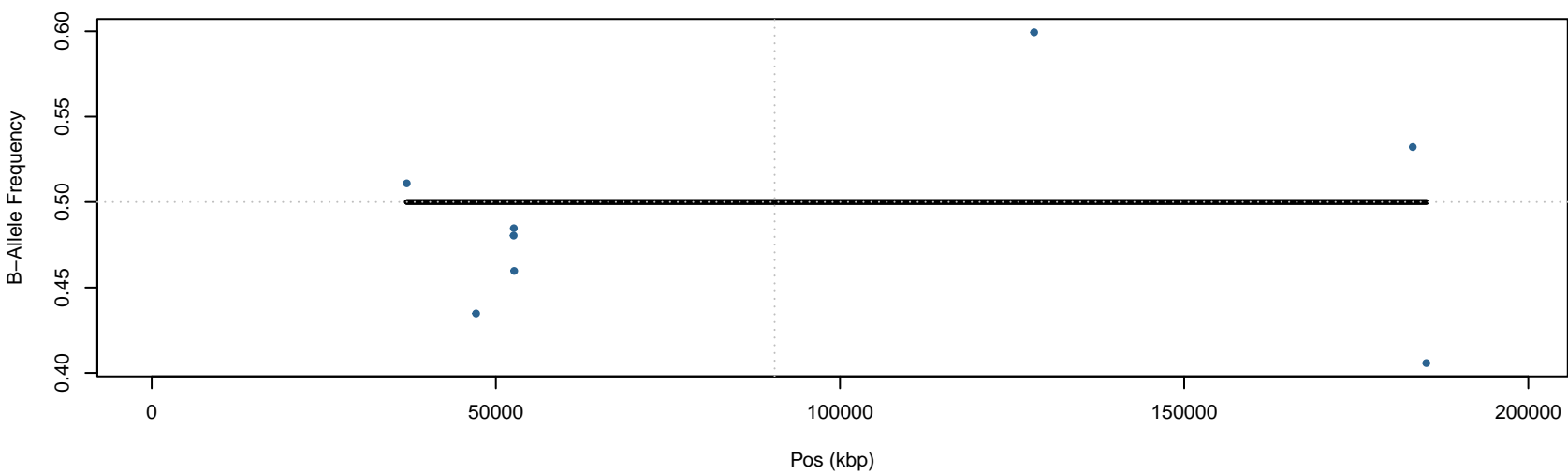
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 1



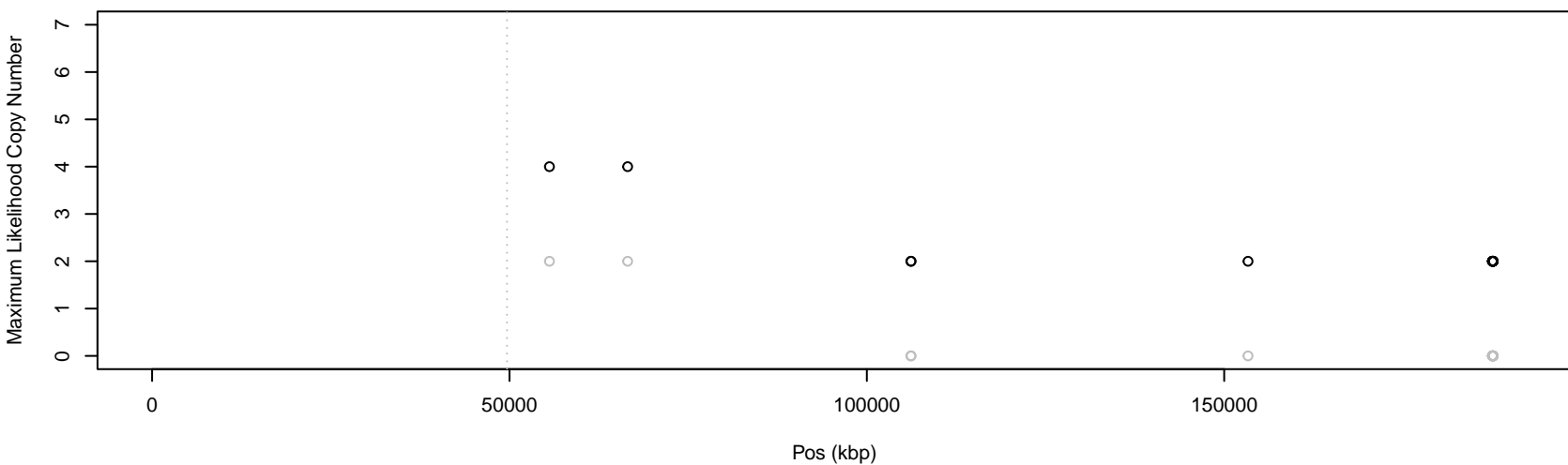
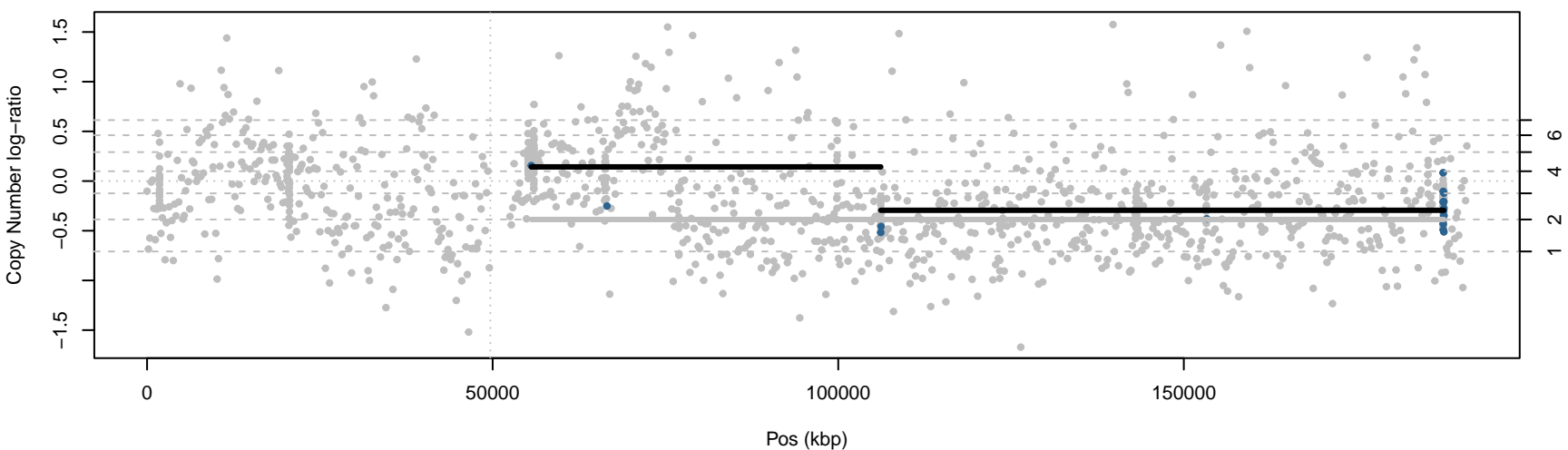
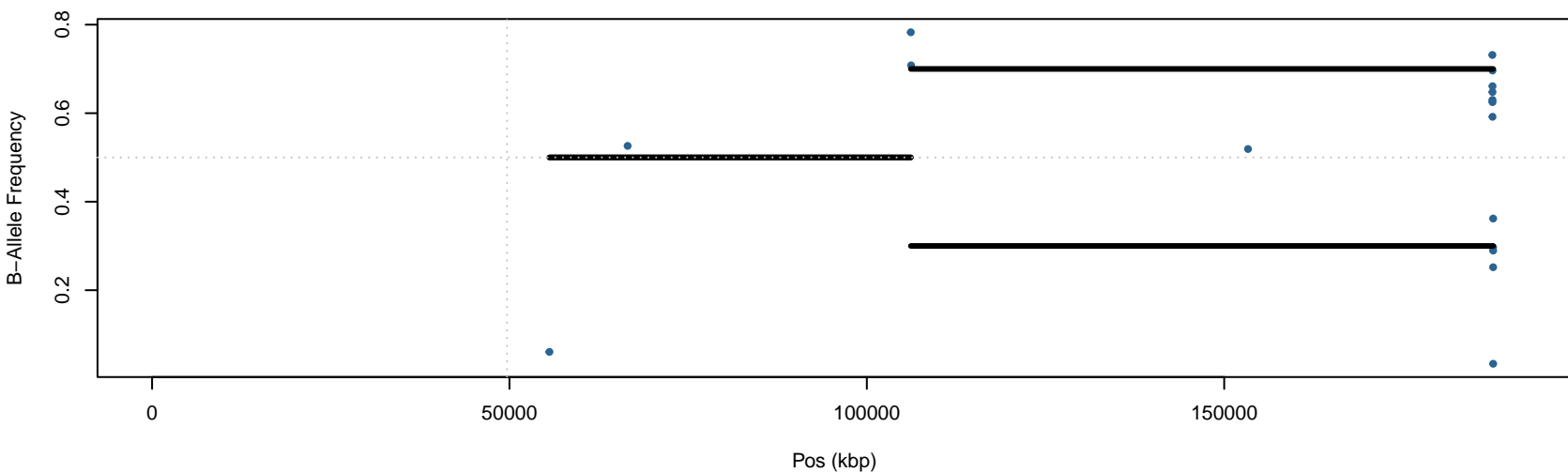
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 2



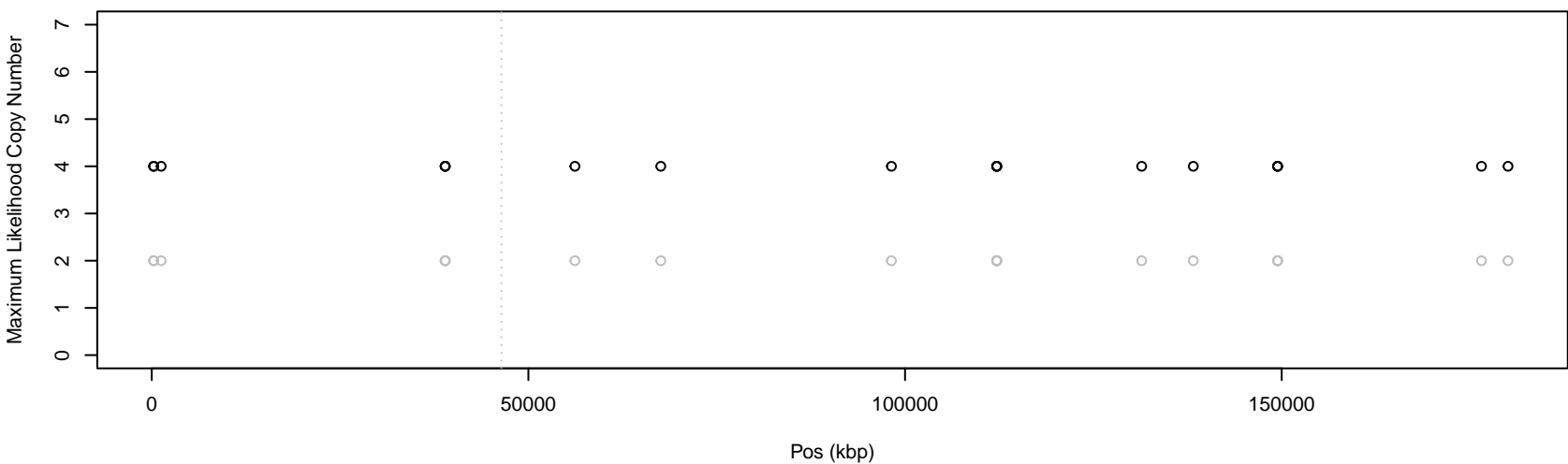
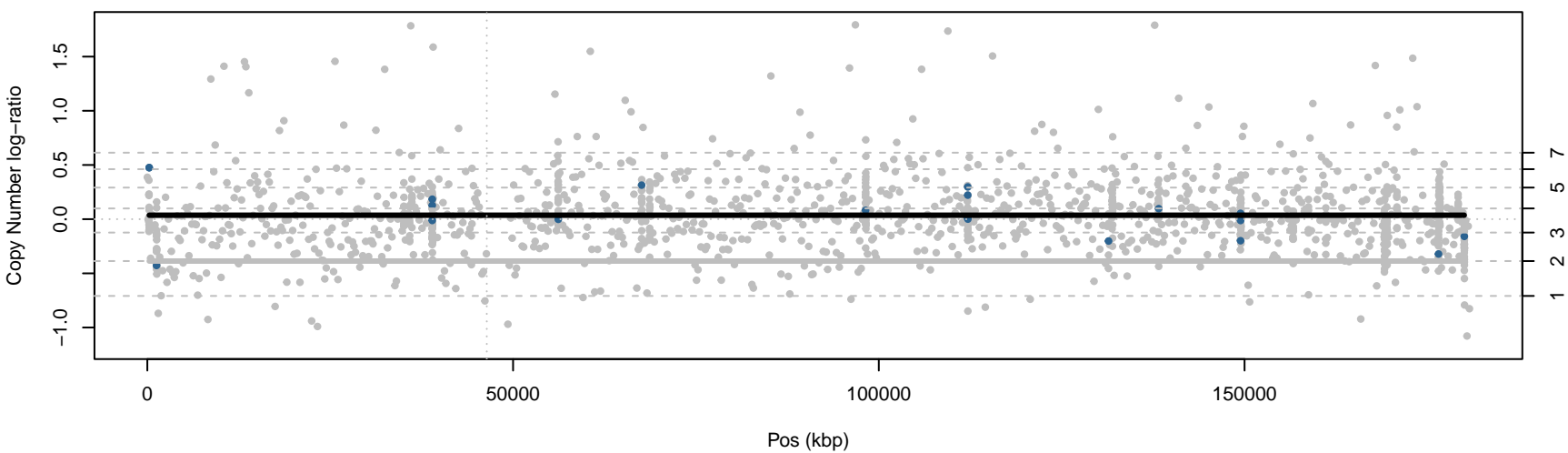
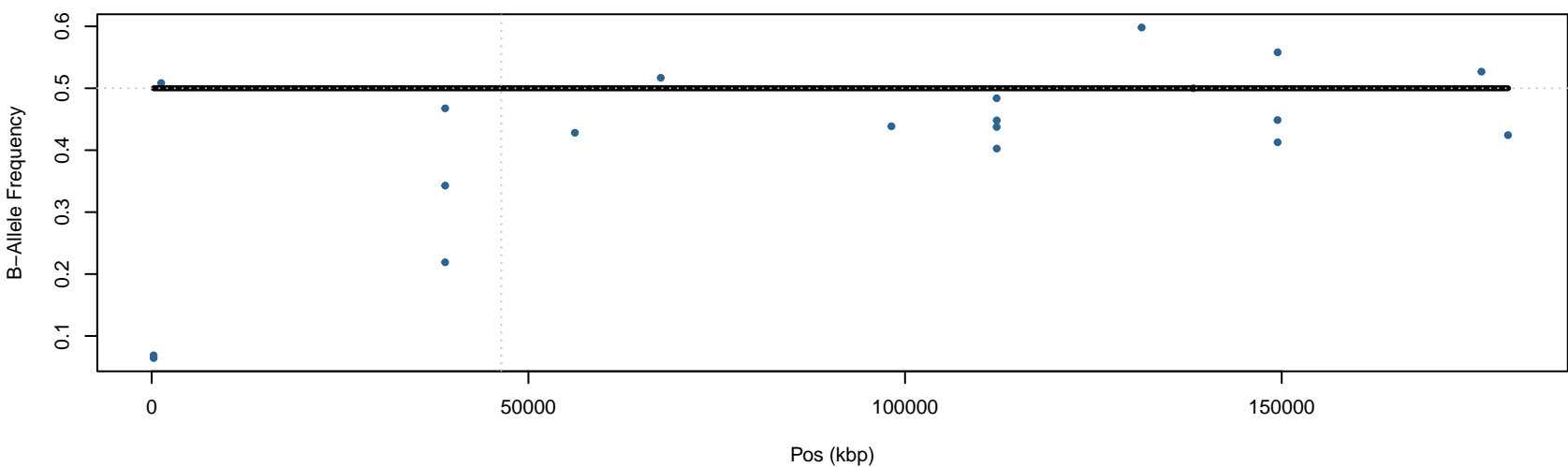
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 3



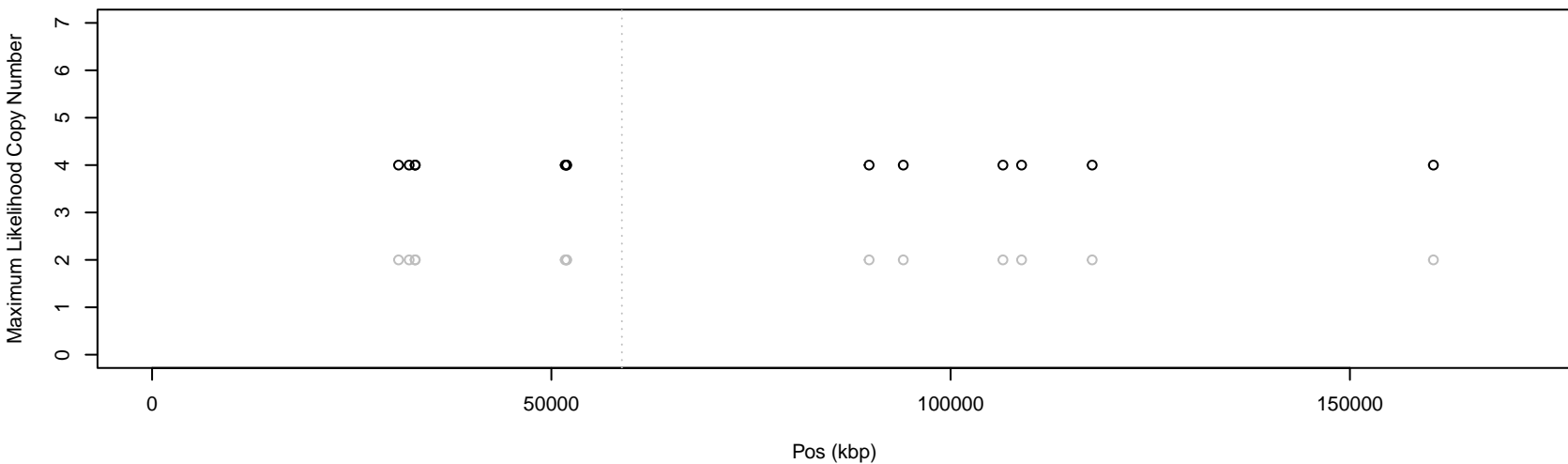
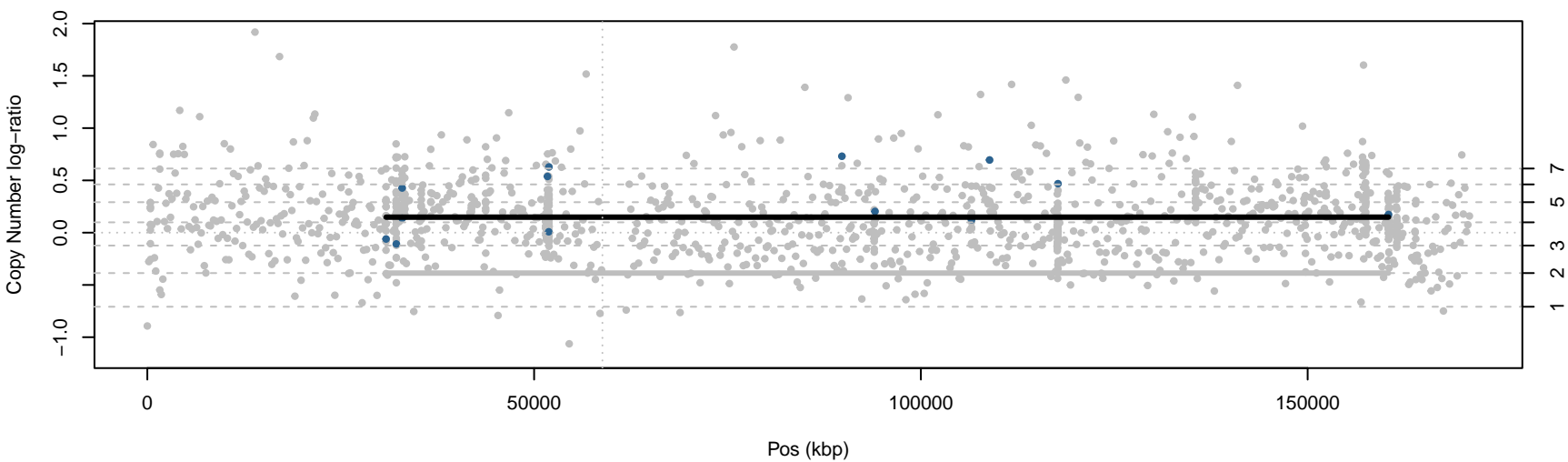
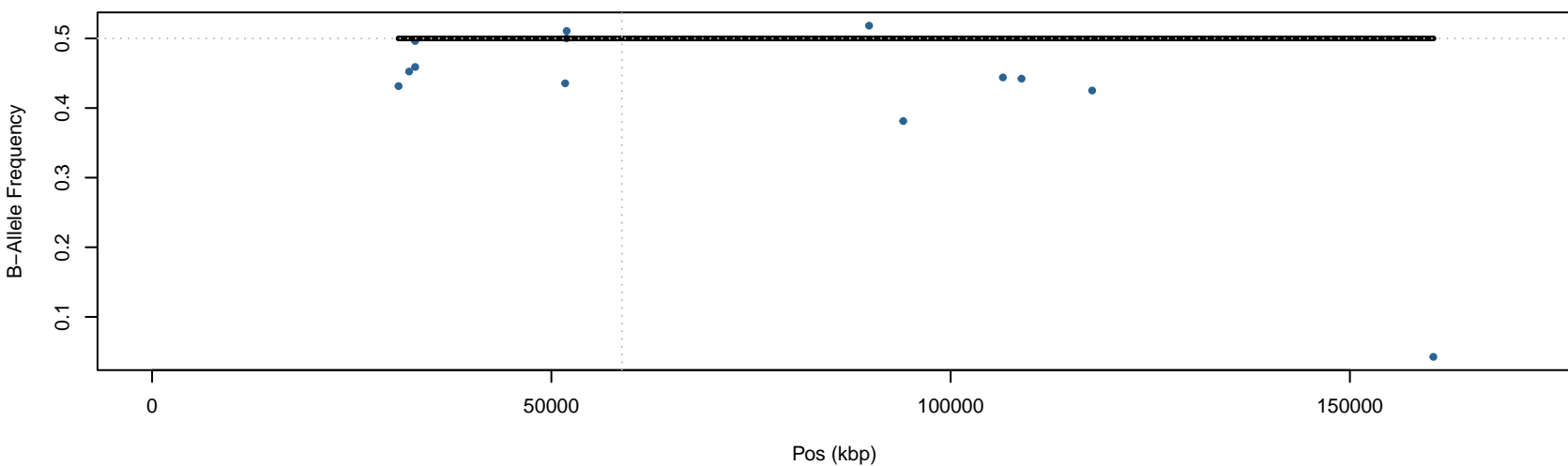
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 4



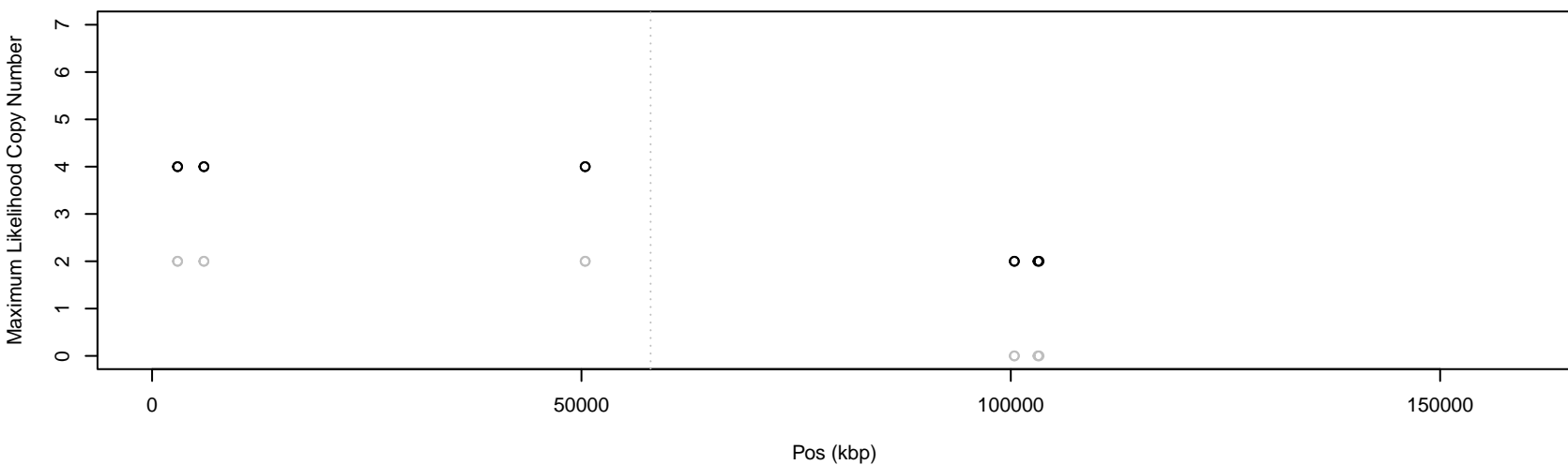
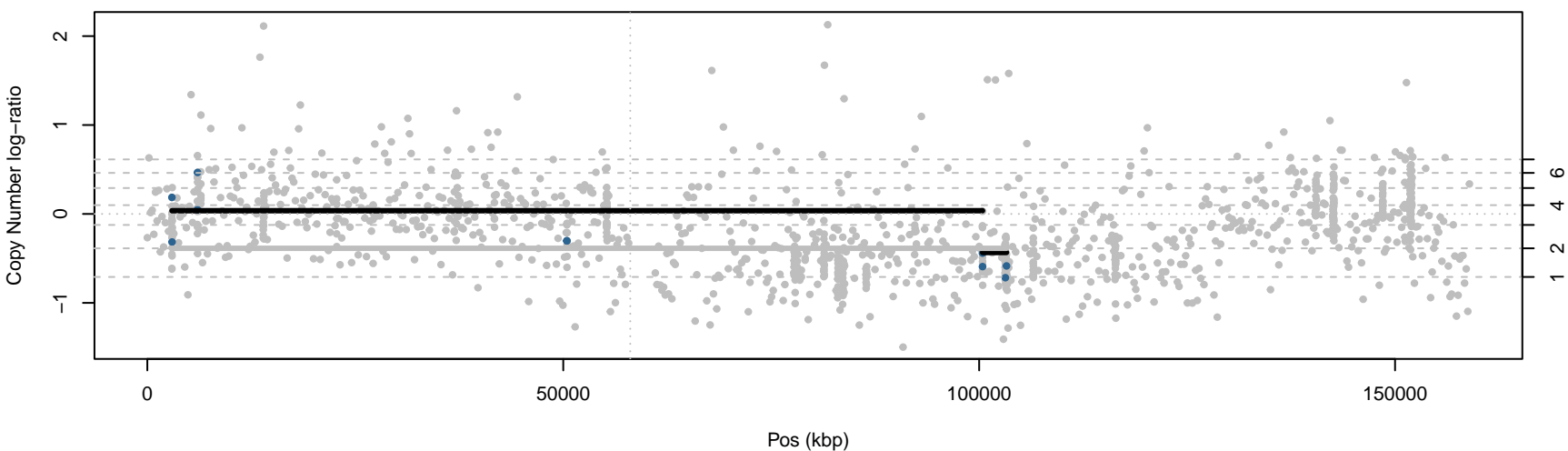
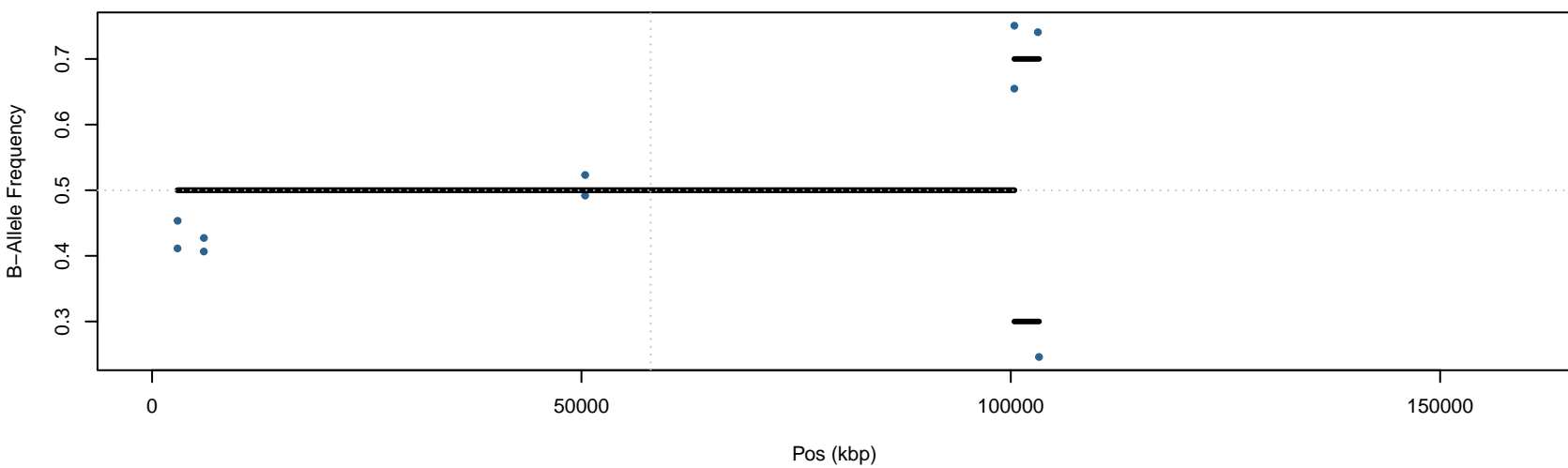
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 5



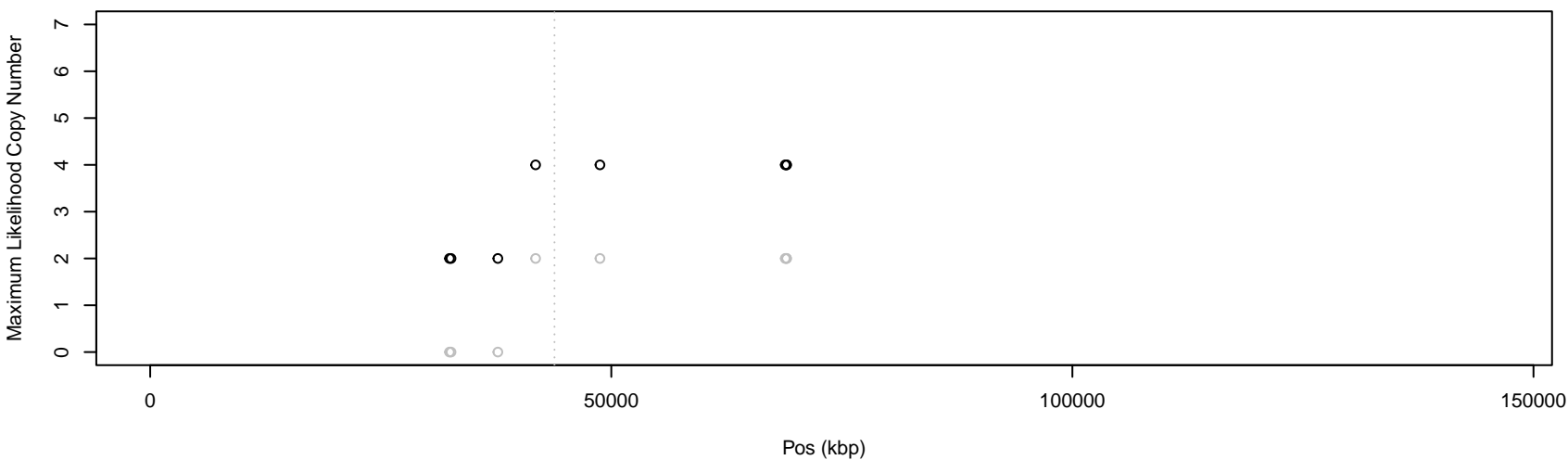
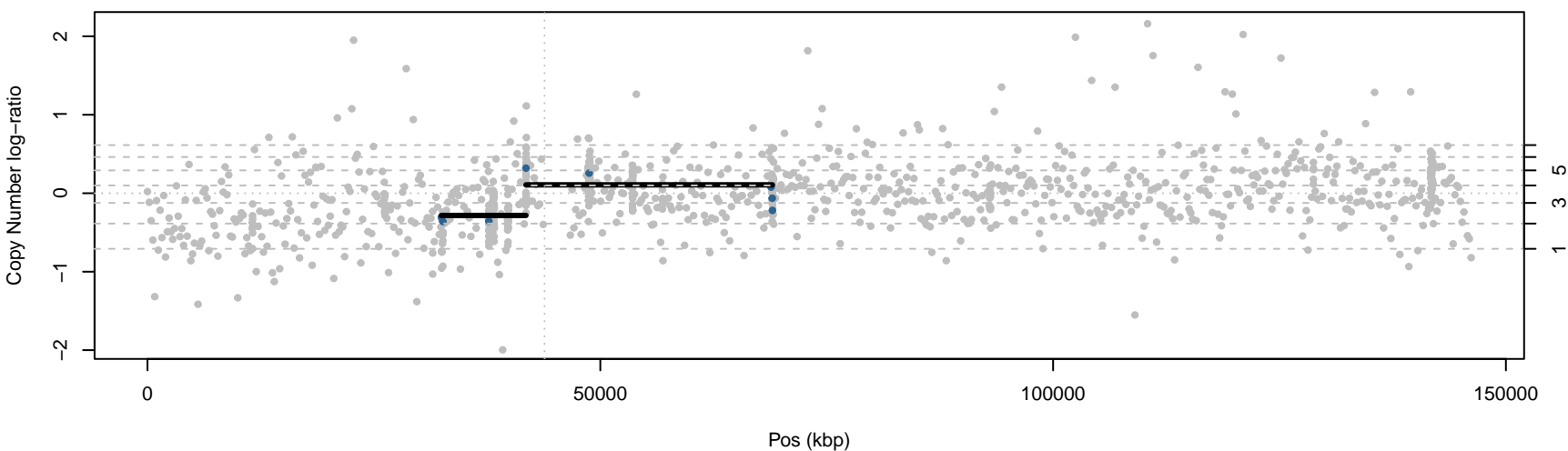
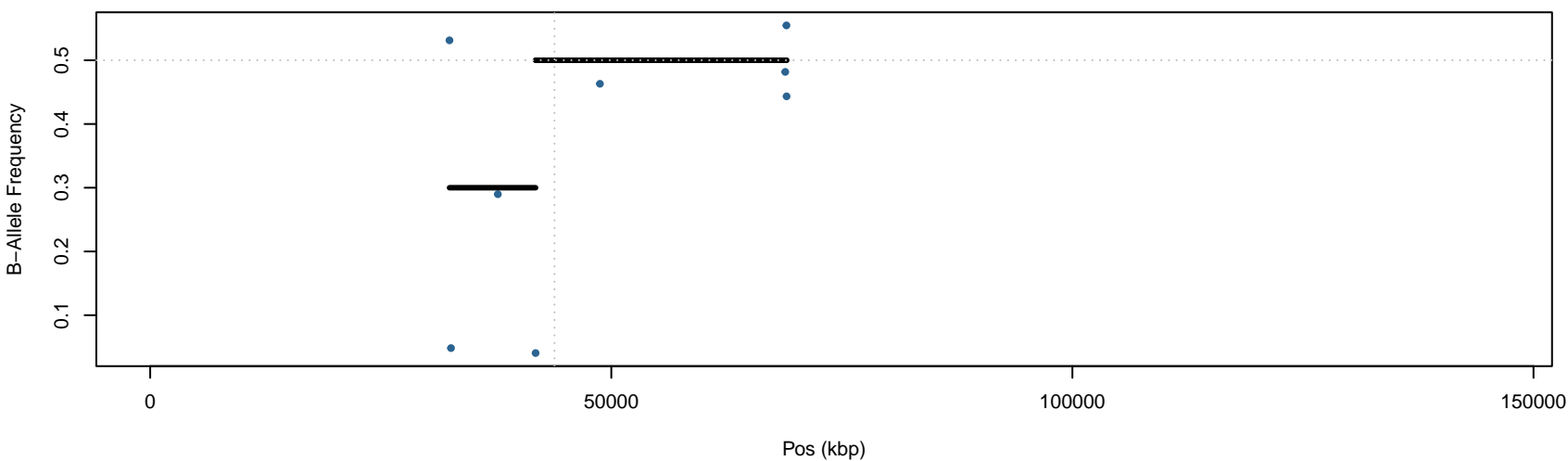
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 6



Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 7

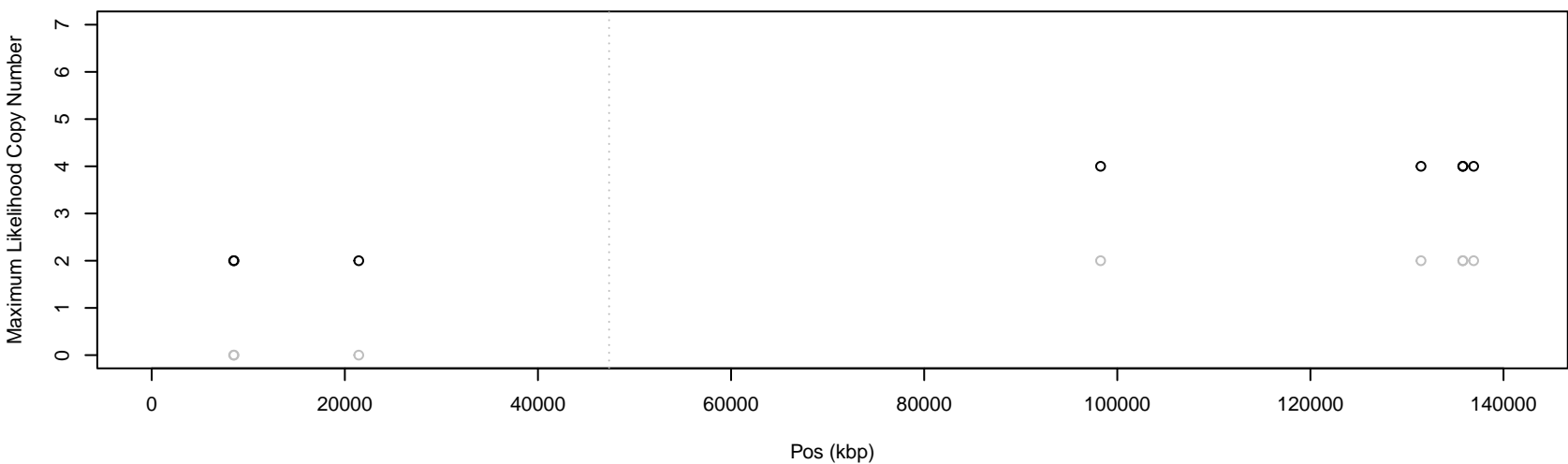
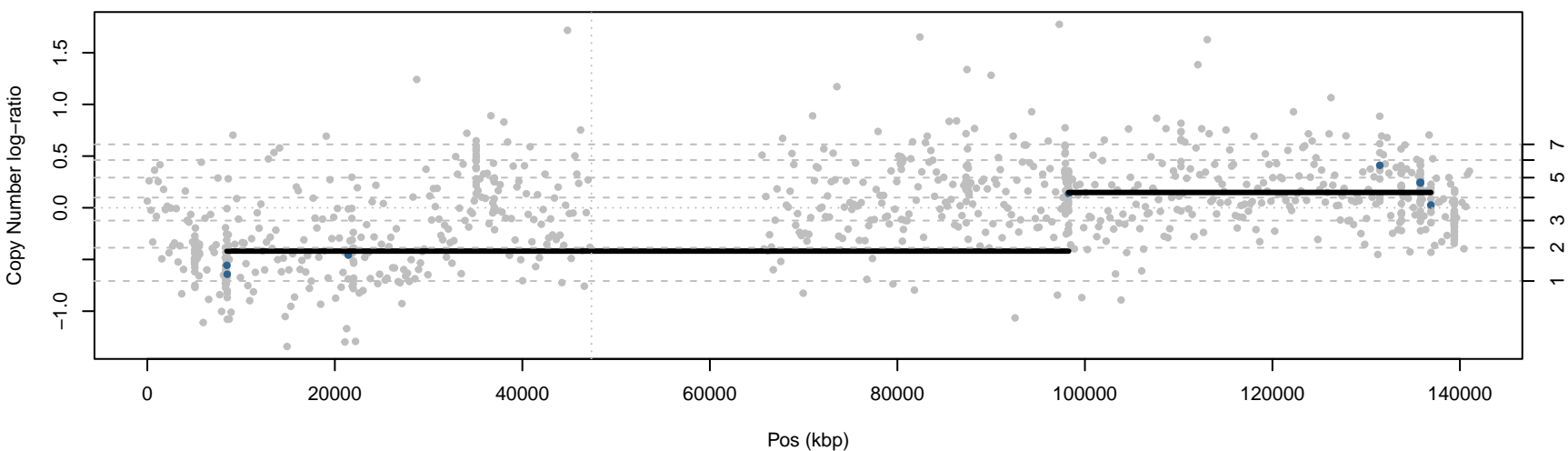
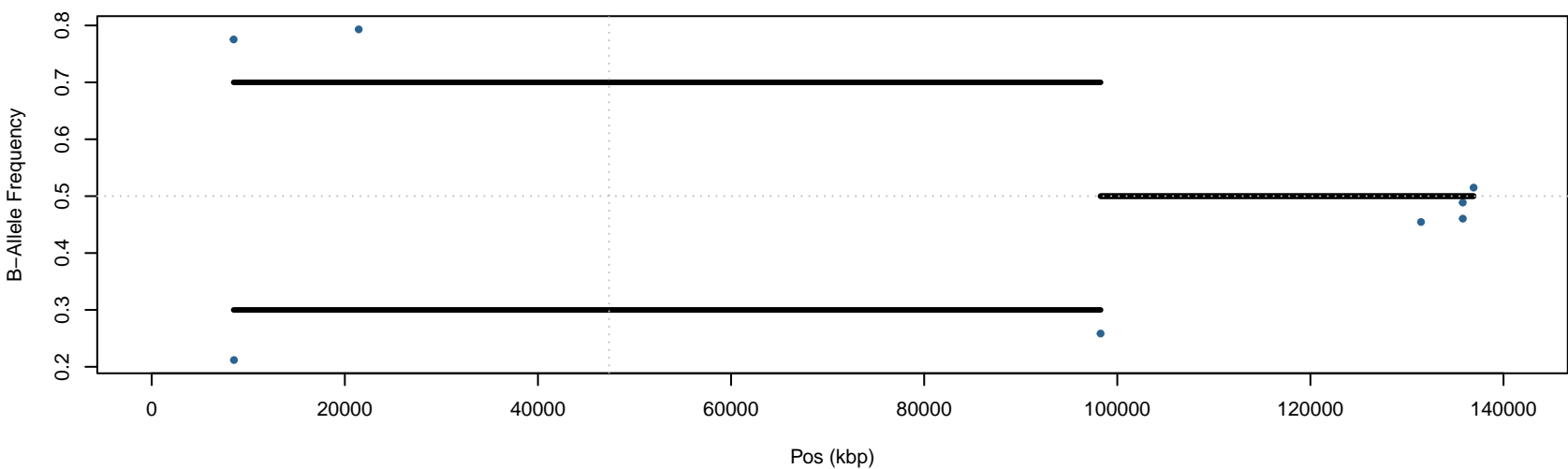


Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 8

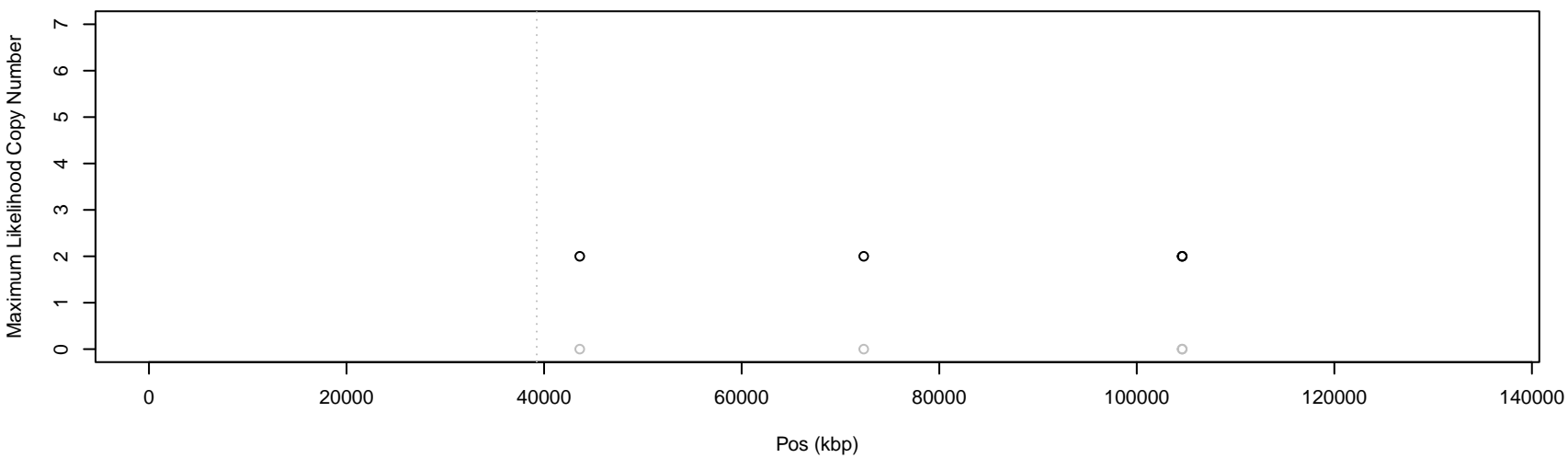
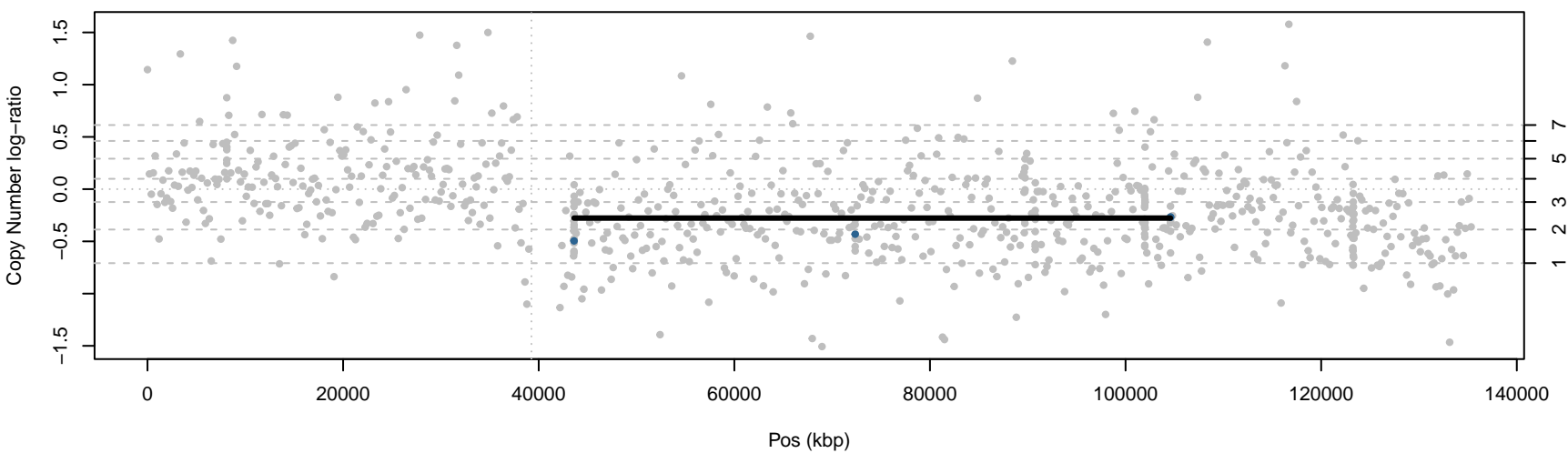
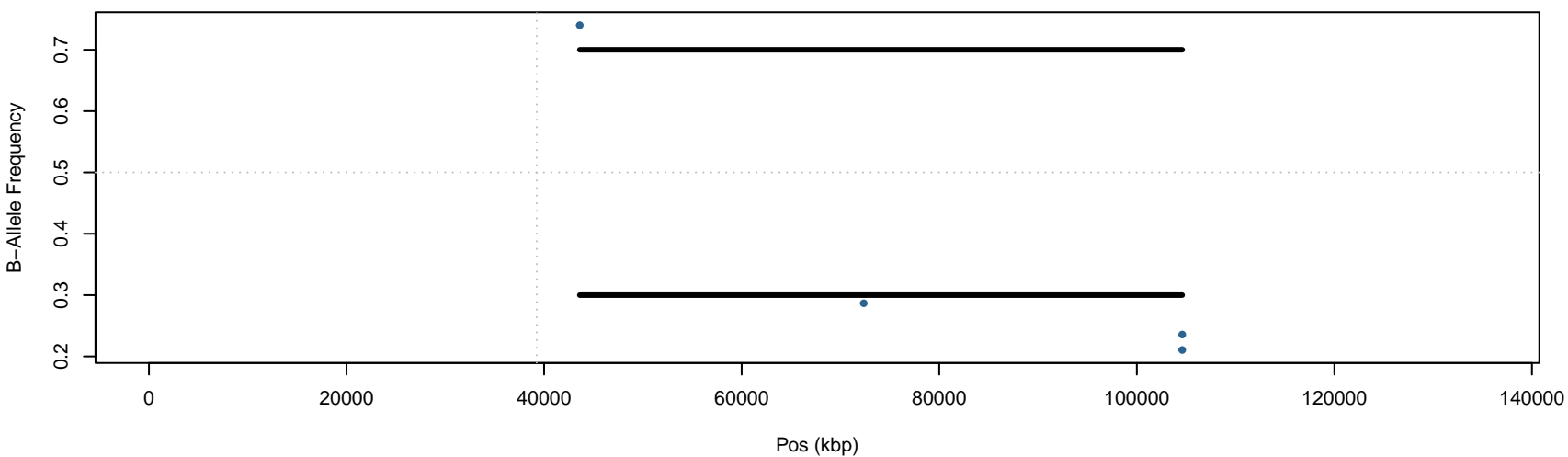




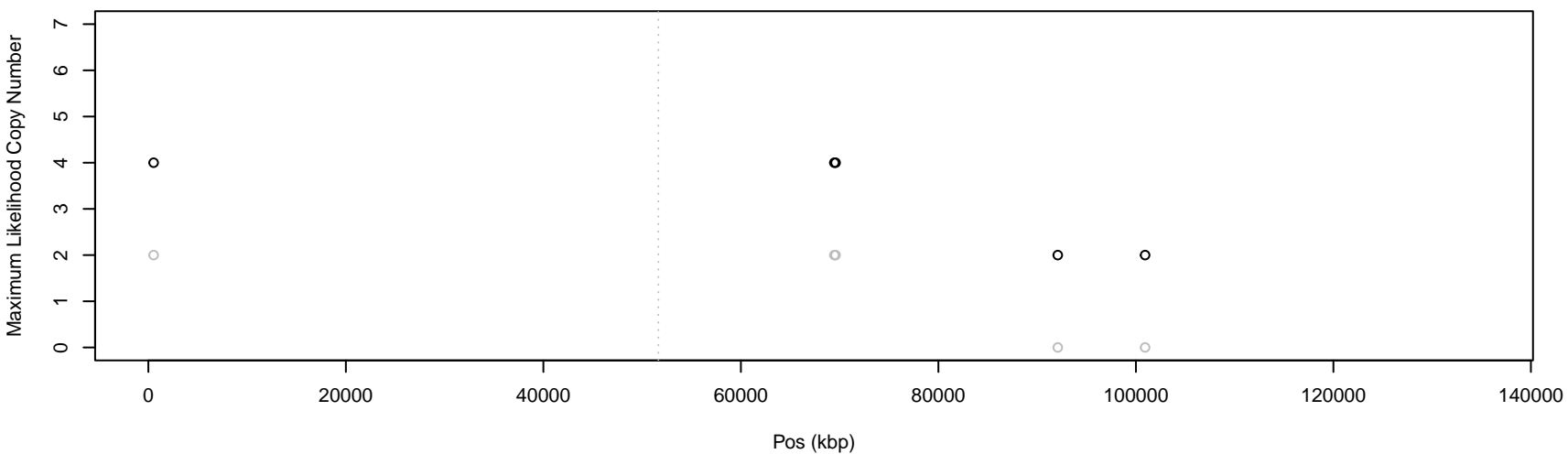
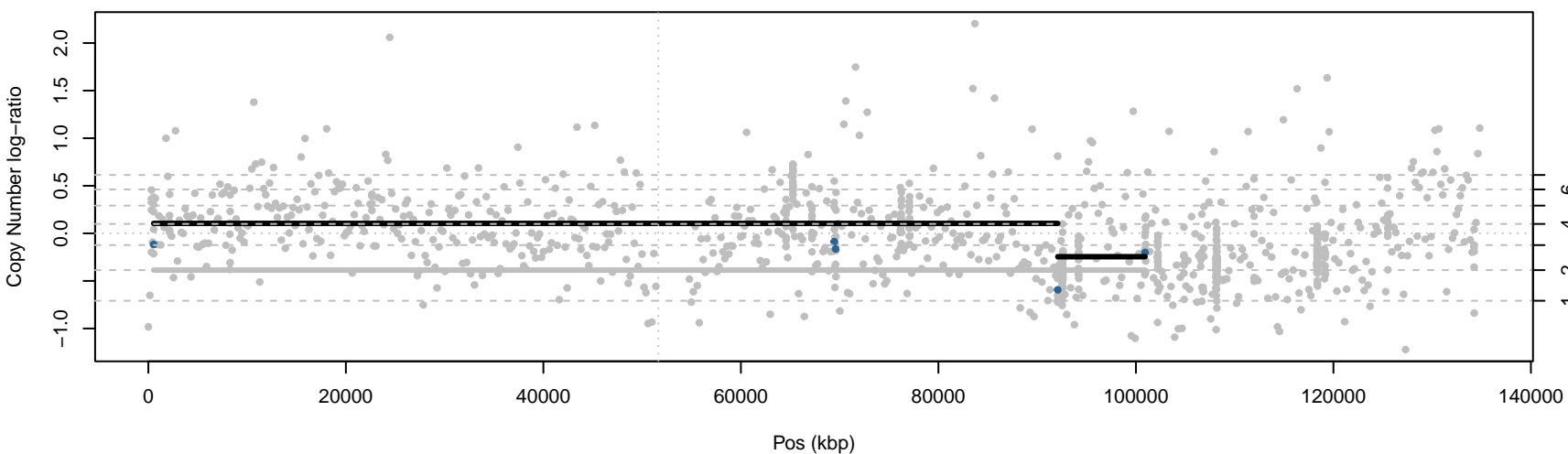
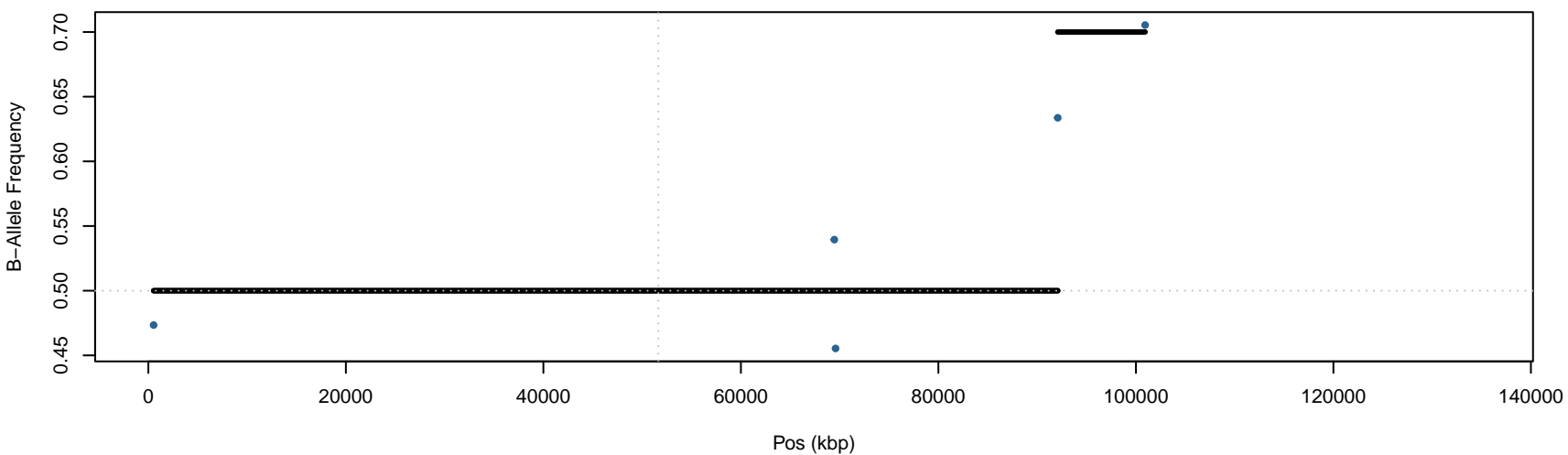
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 9



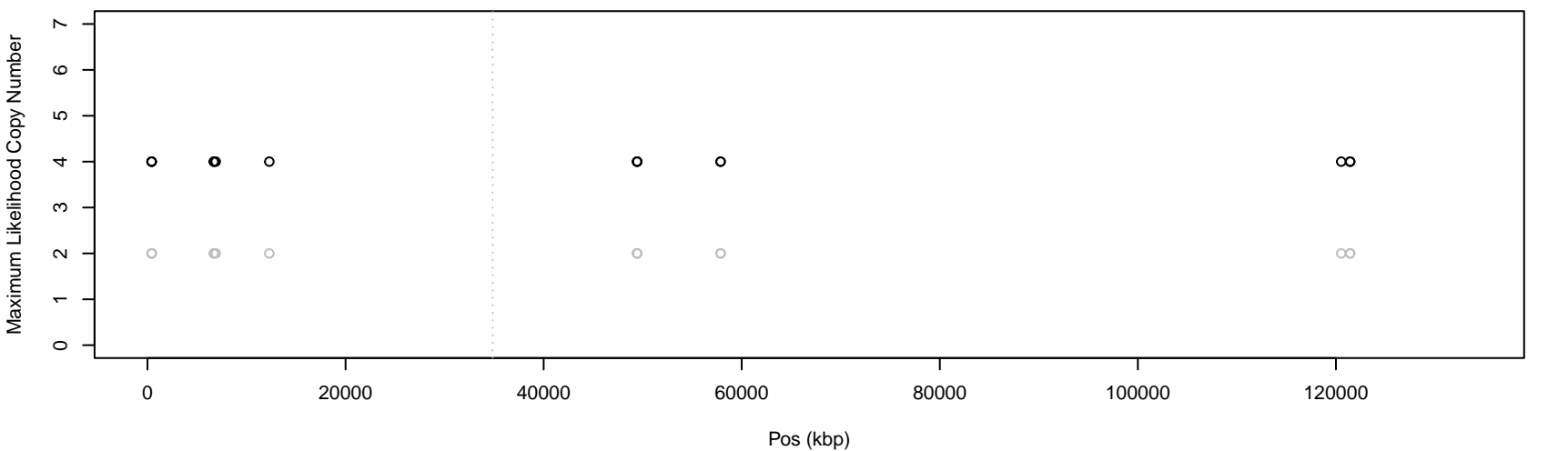
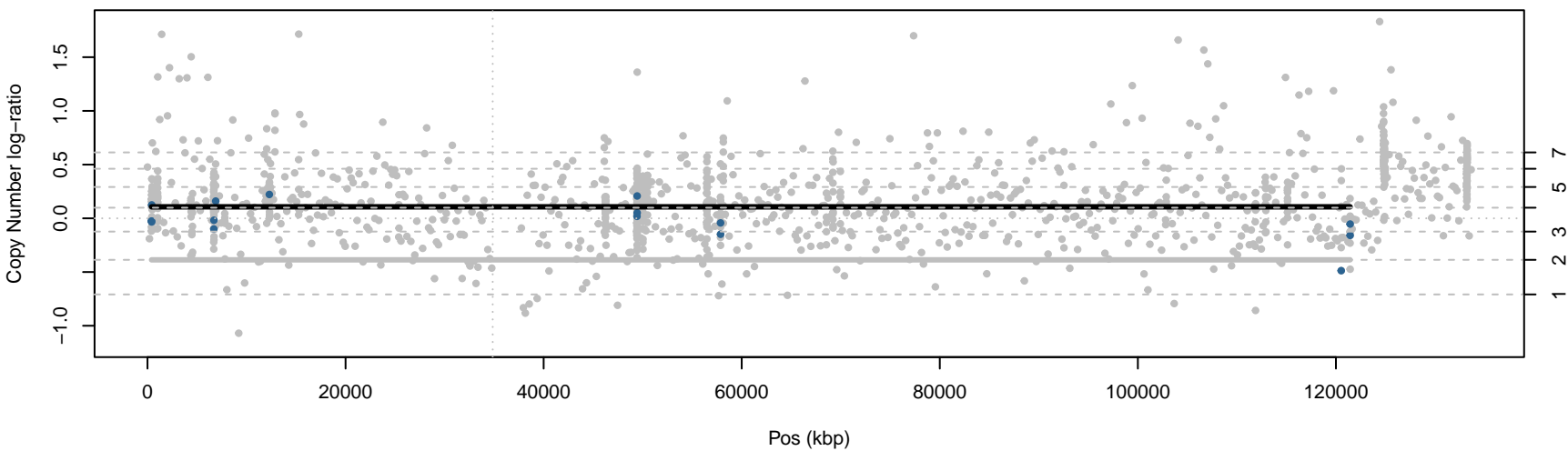
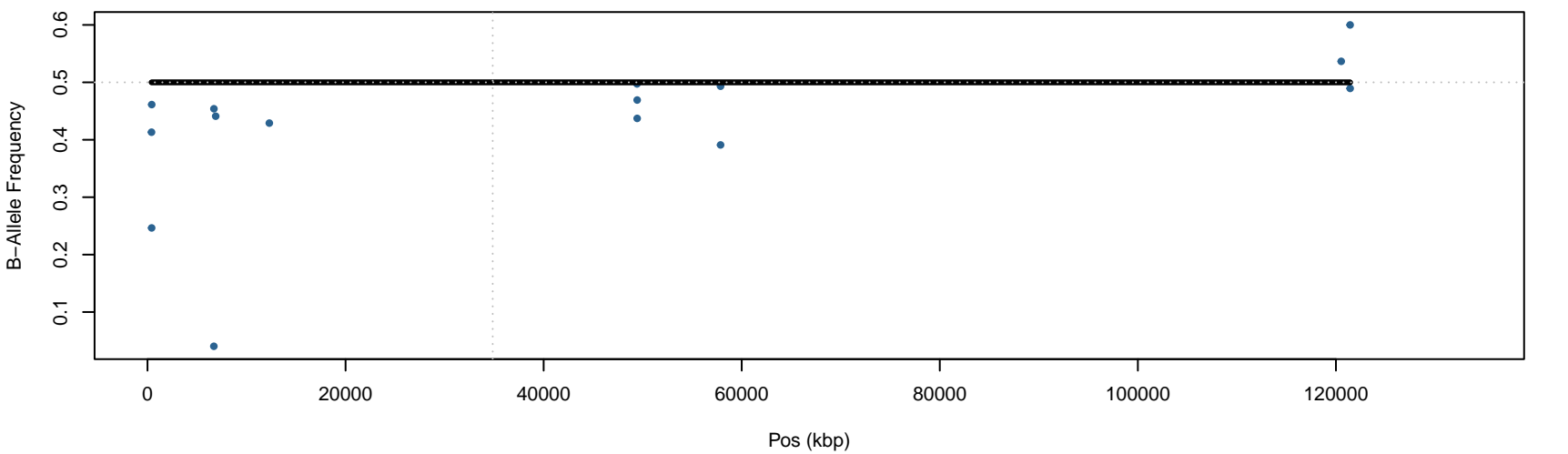
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 10



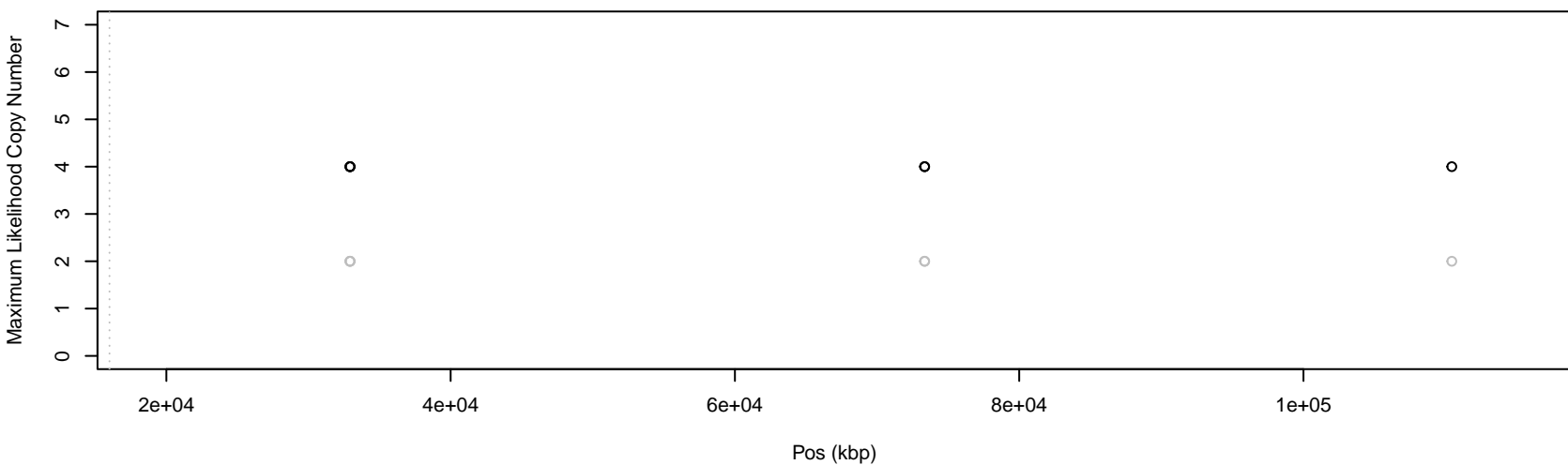
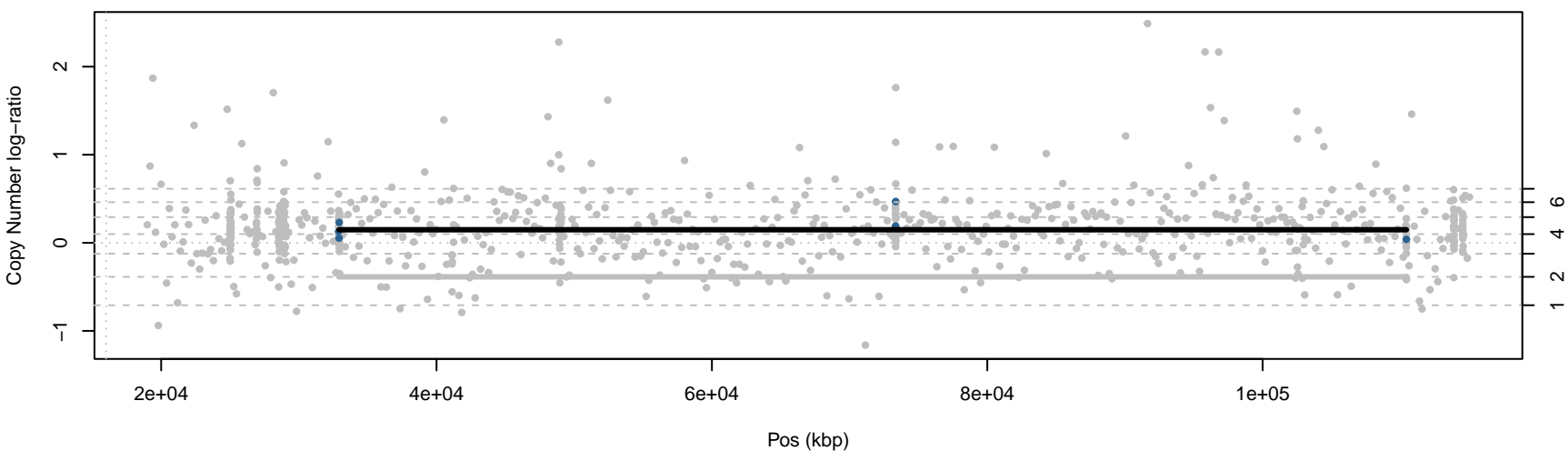
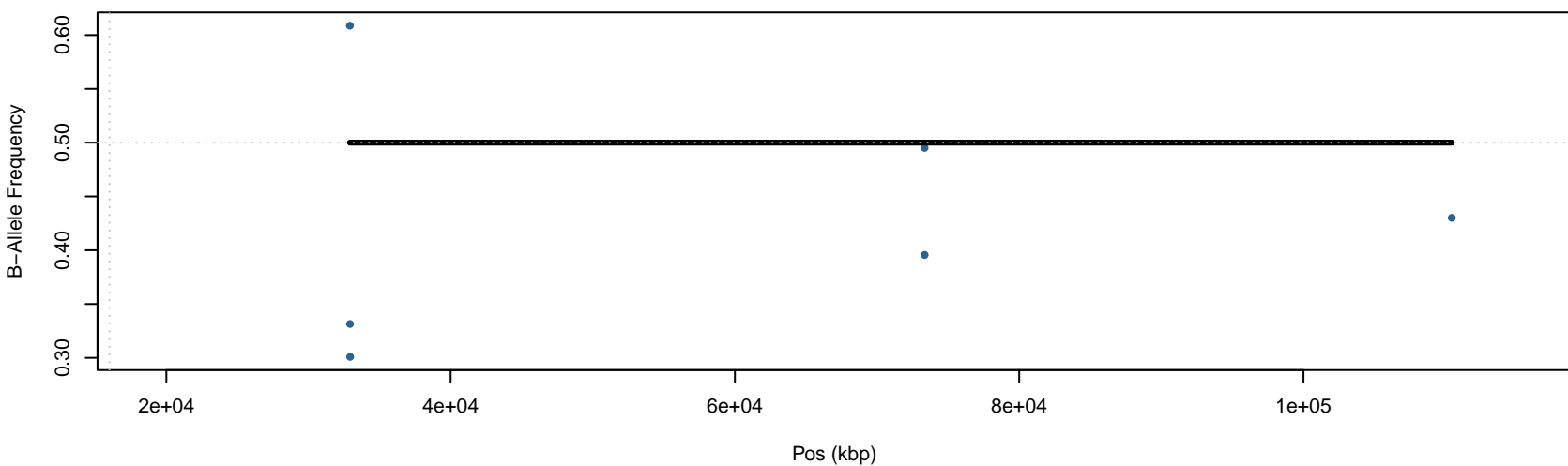
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 11



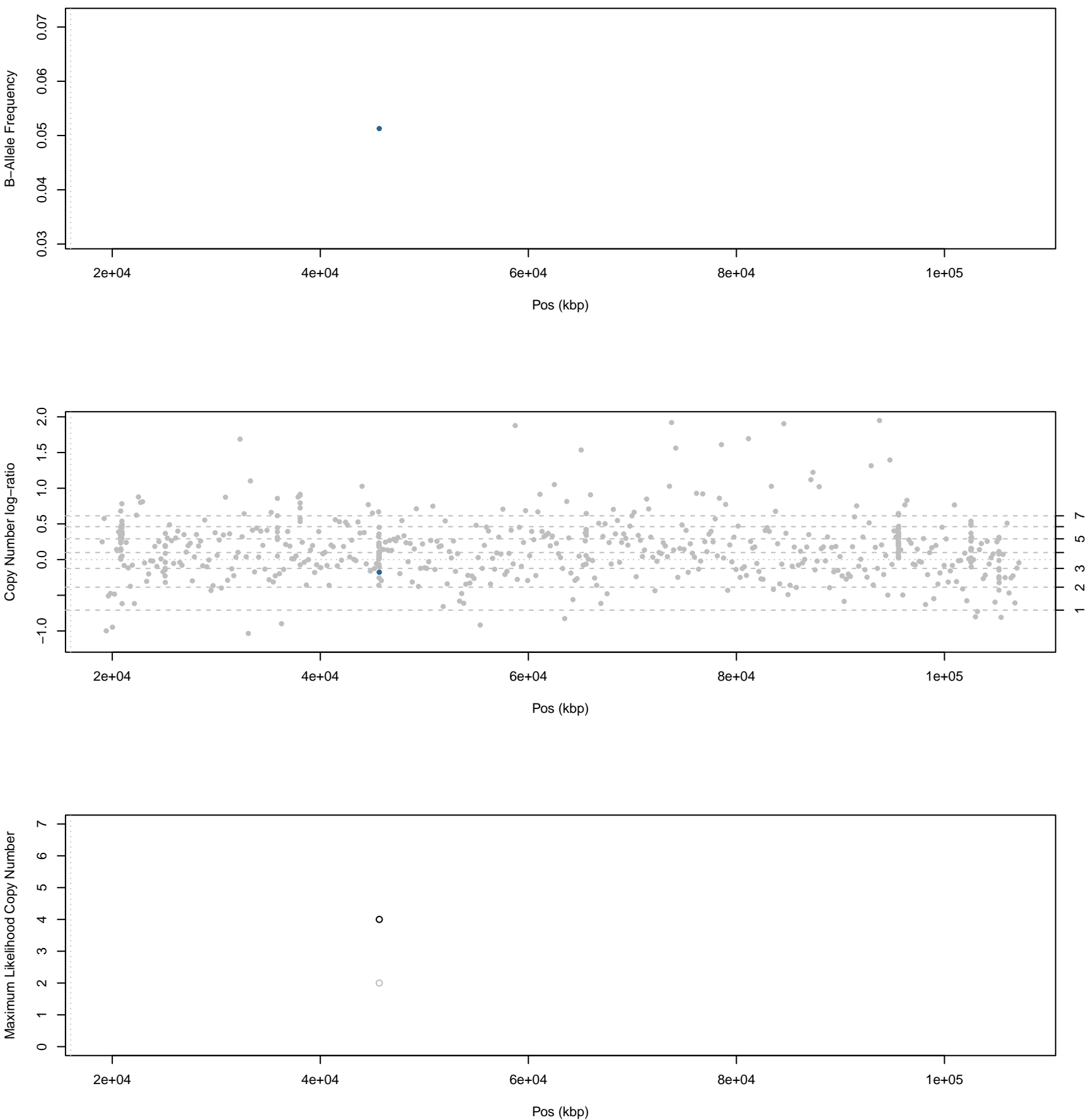
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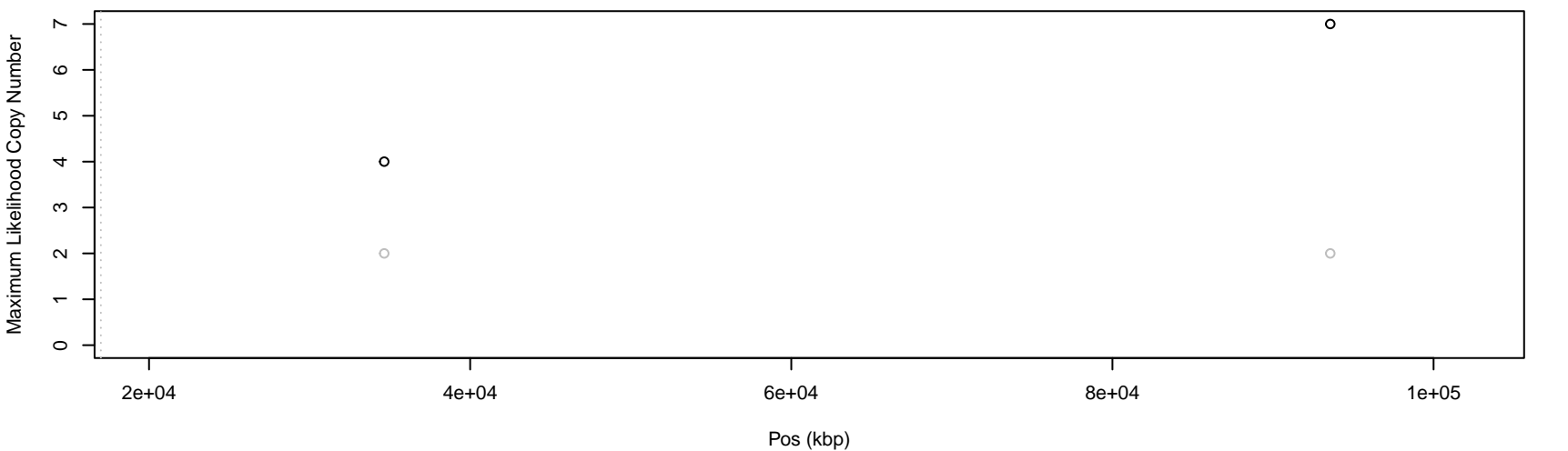
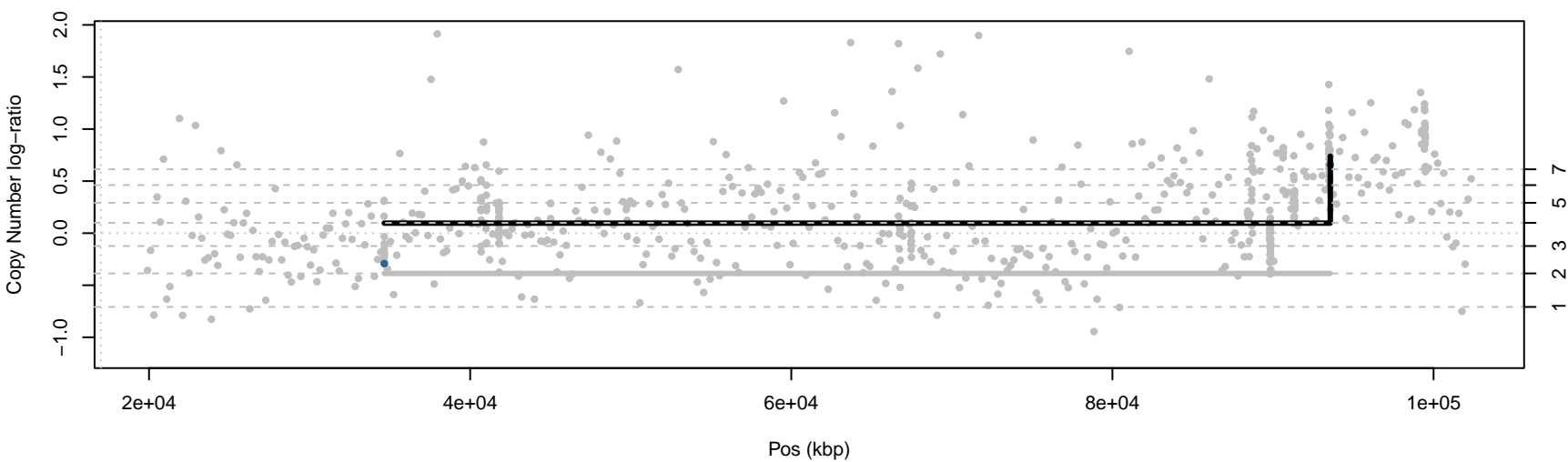
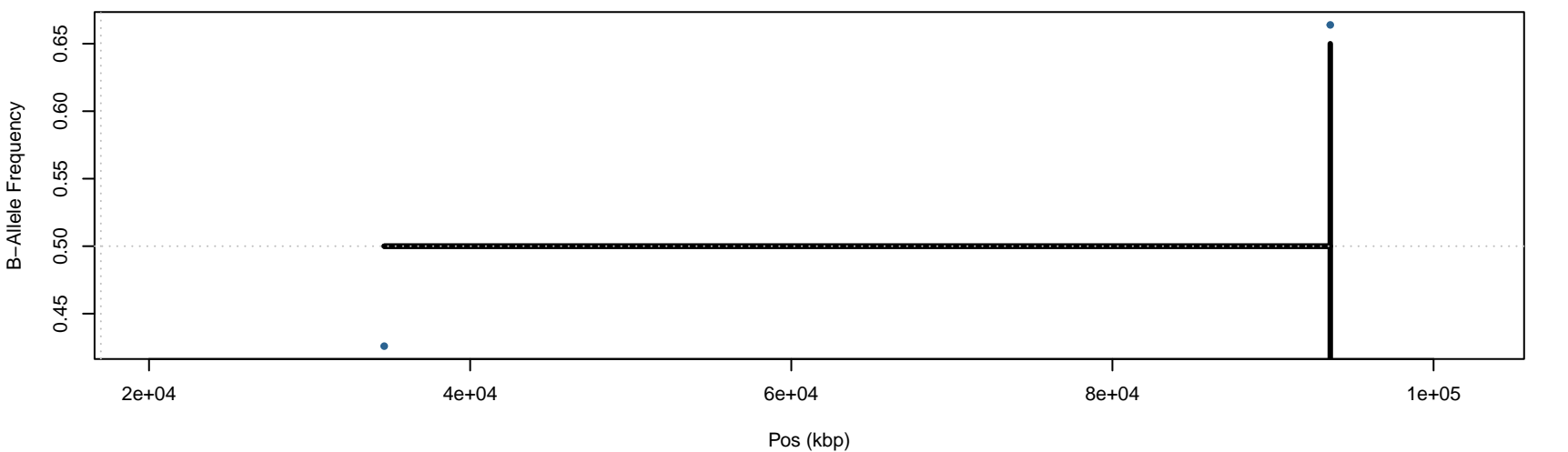
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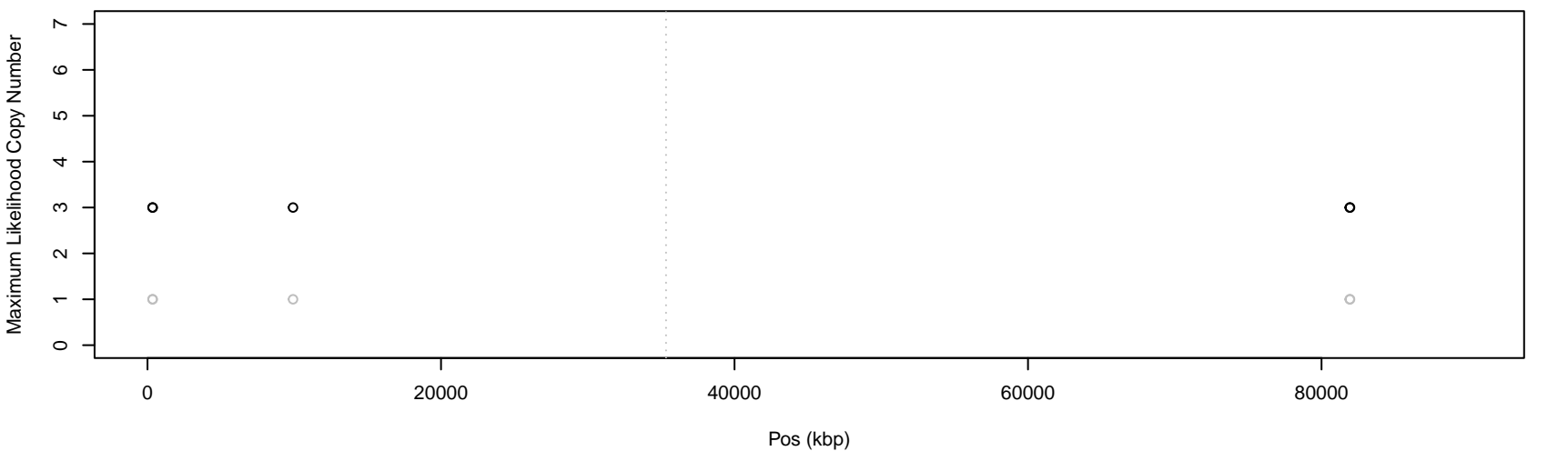
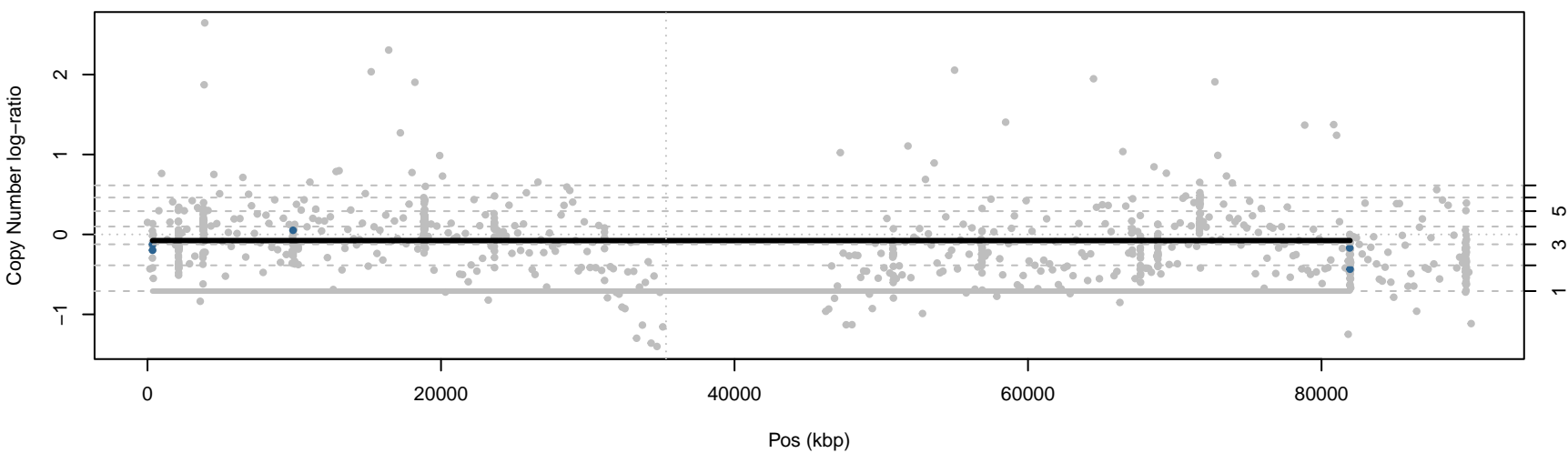
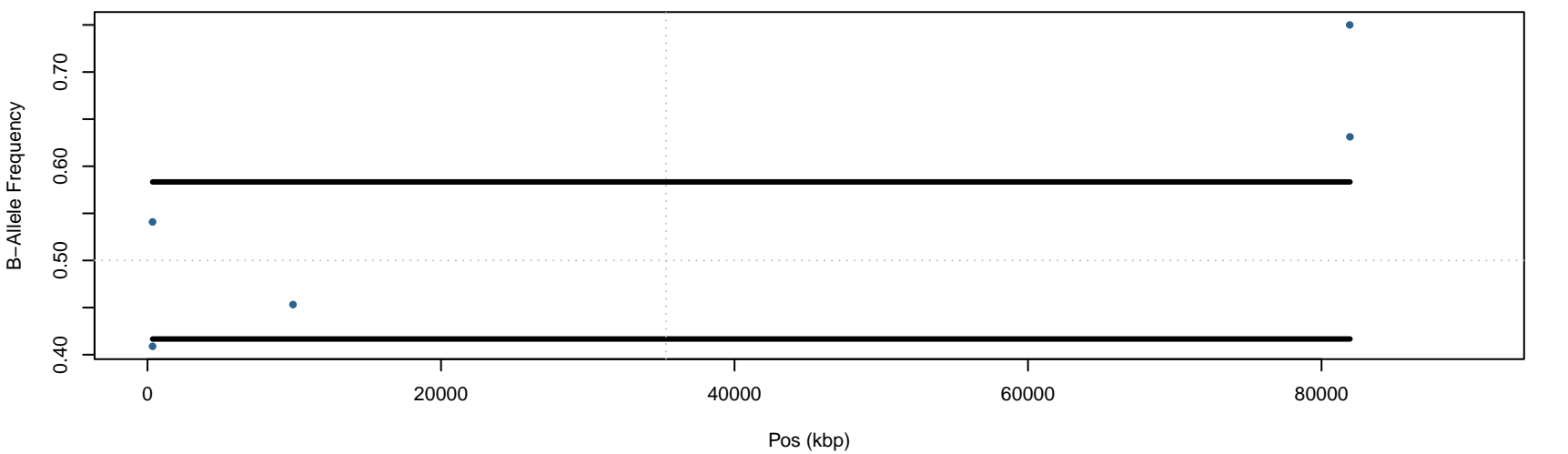
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 14



Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 15

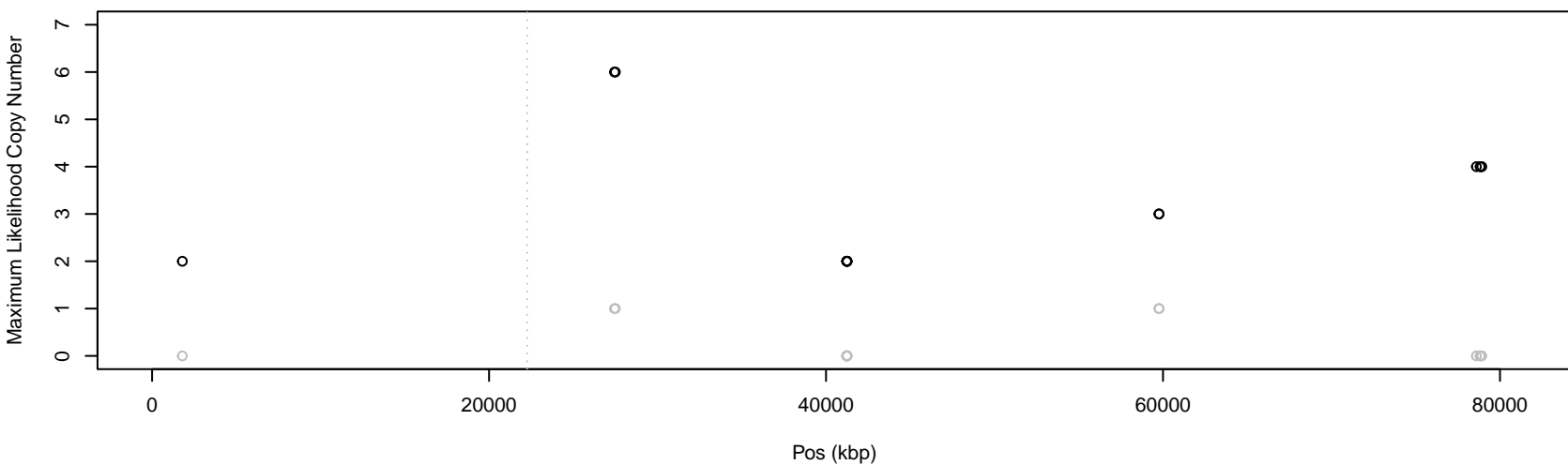
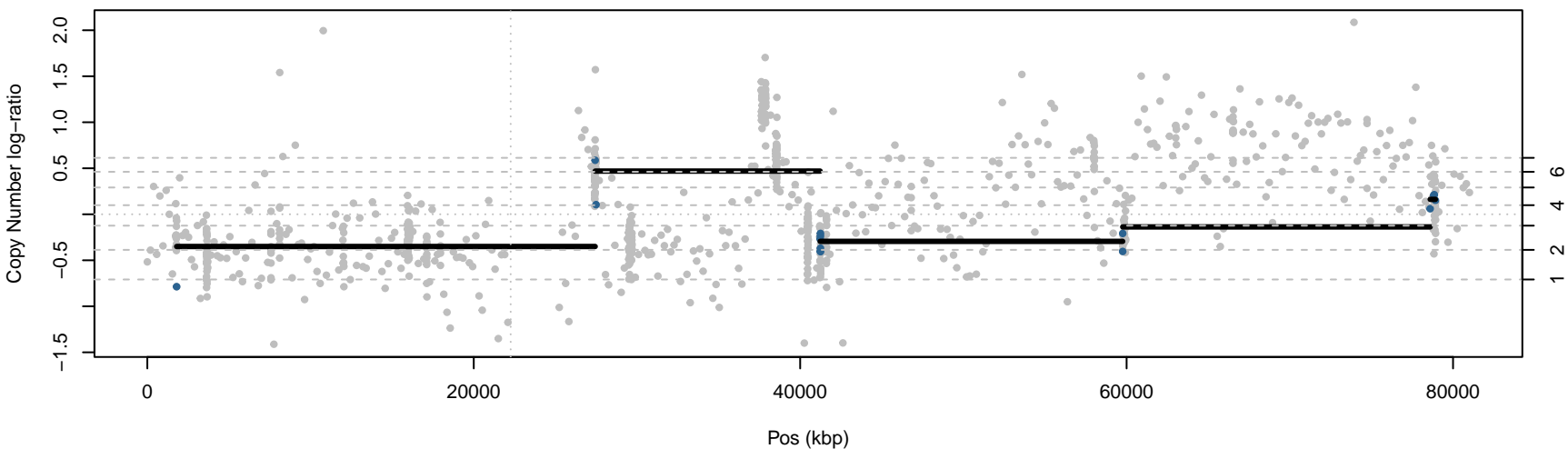
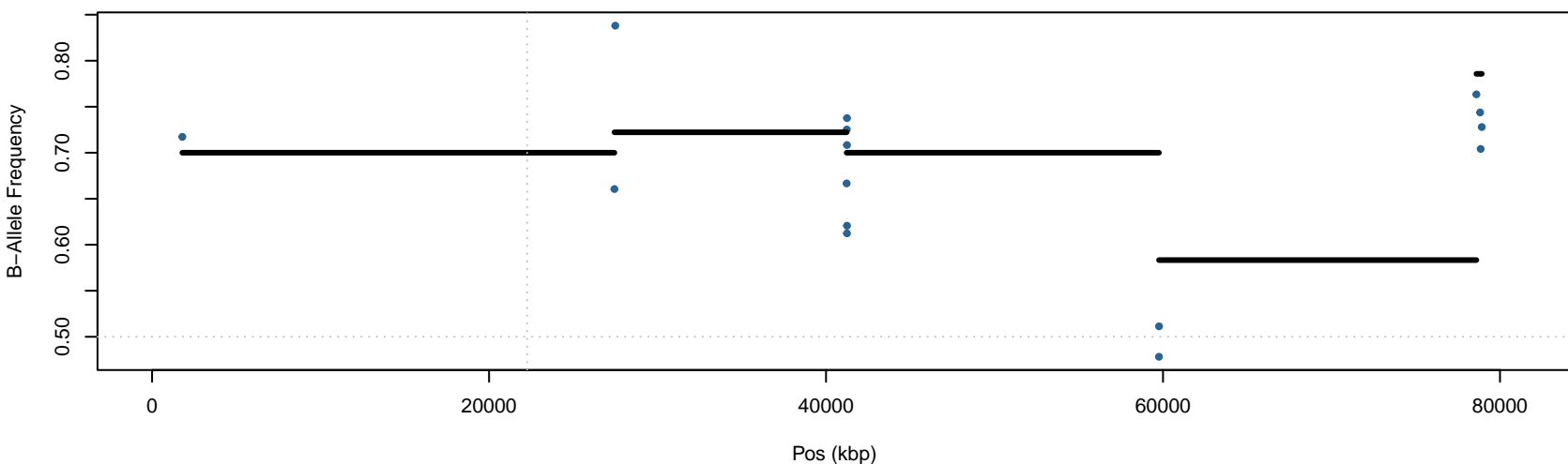


Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 16

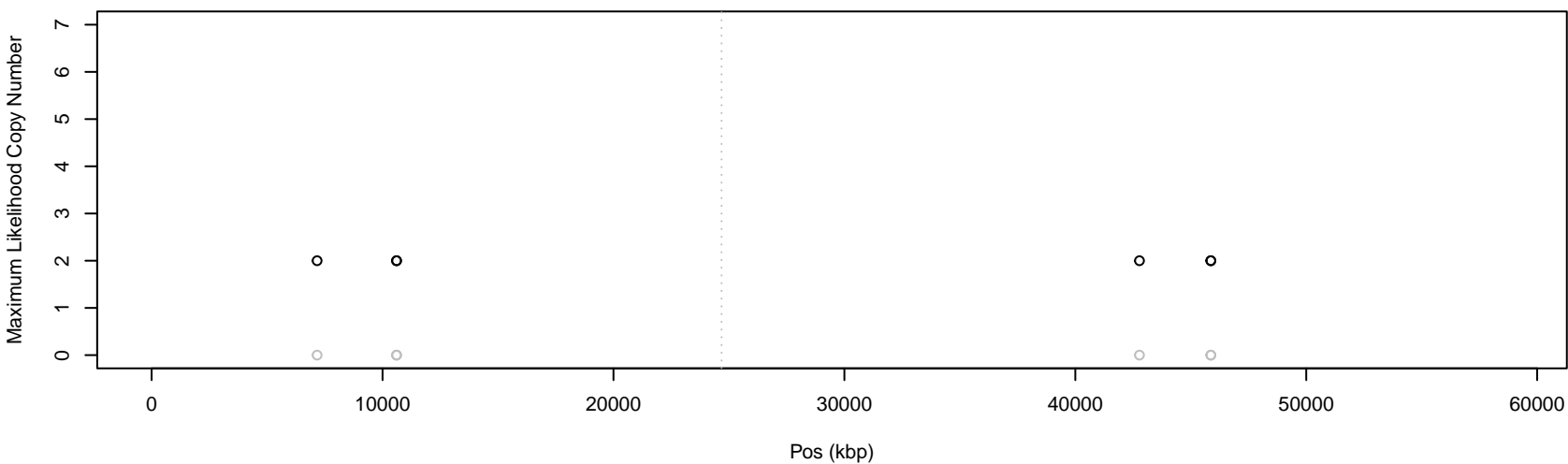
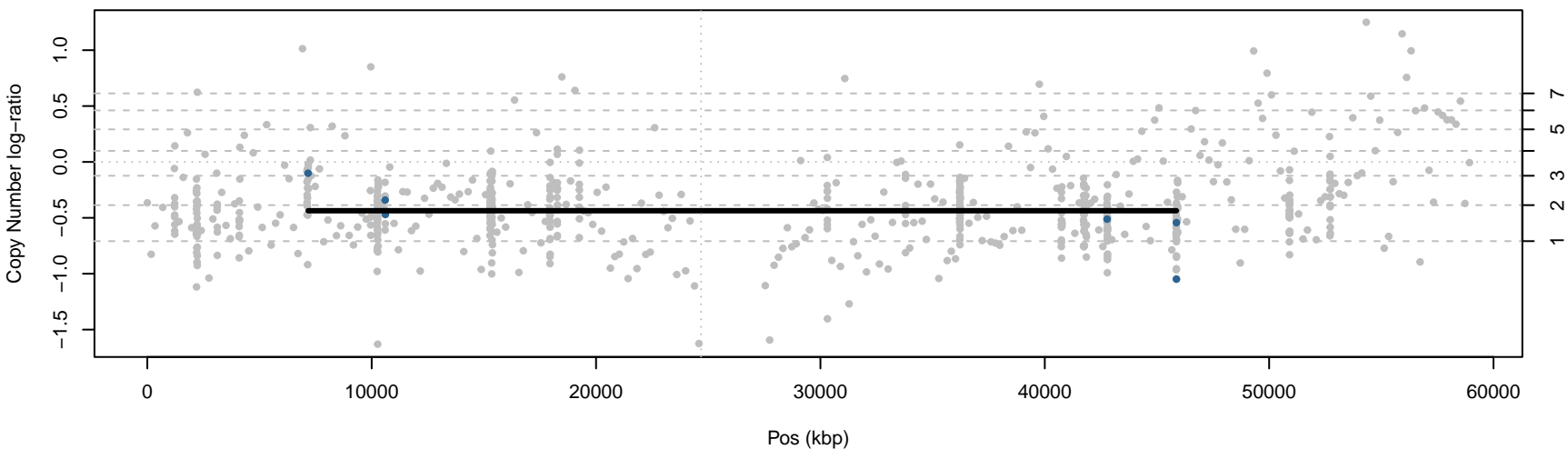
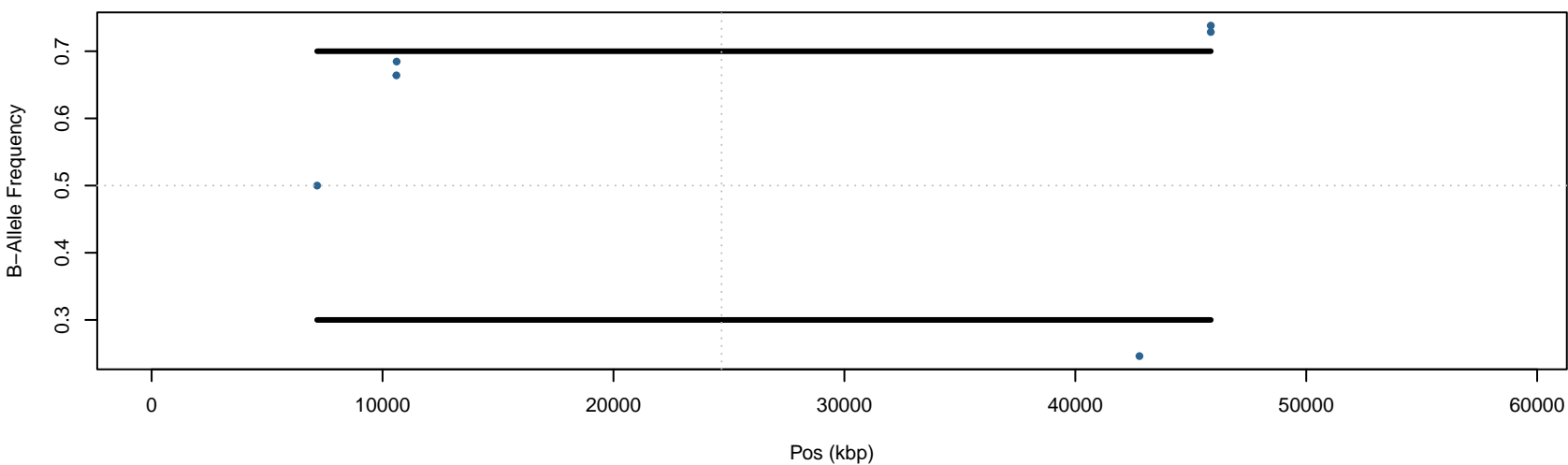




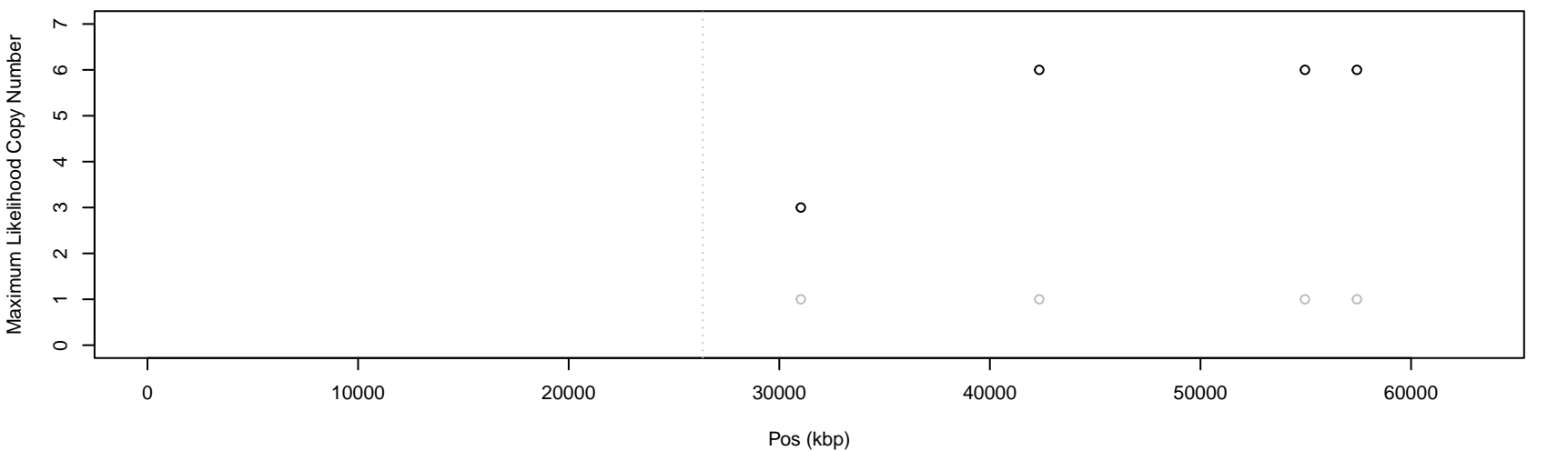
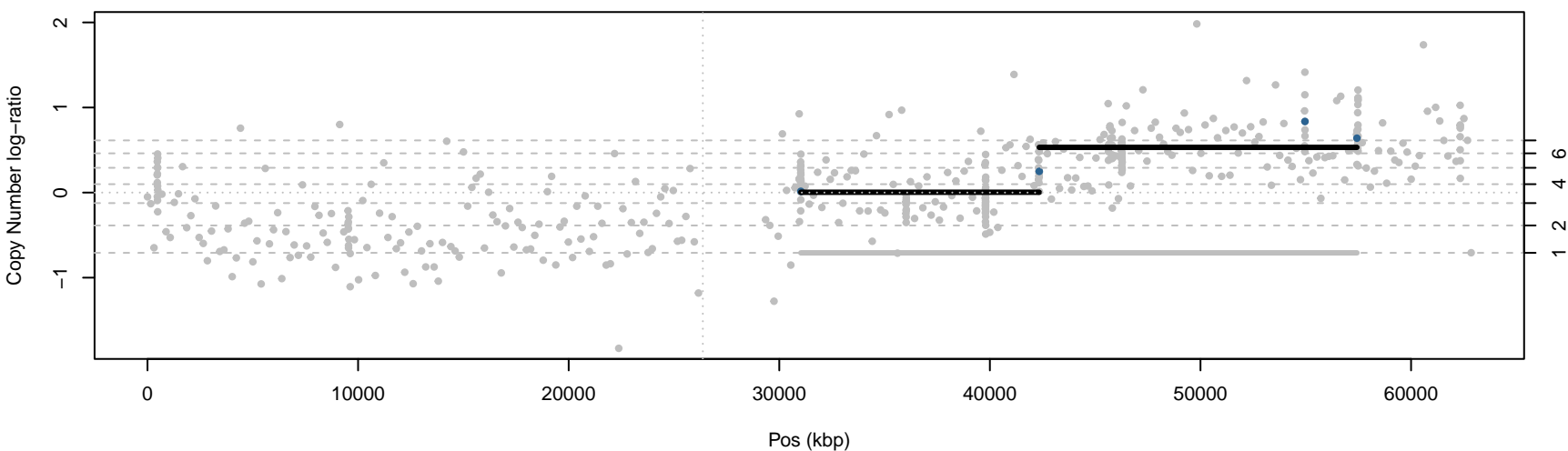
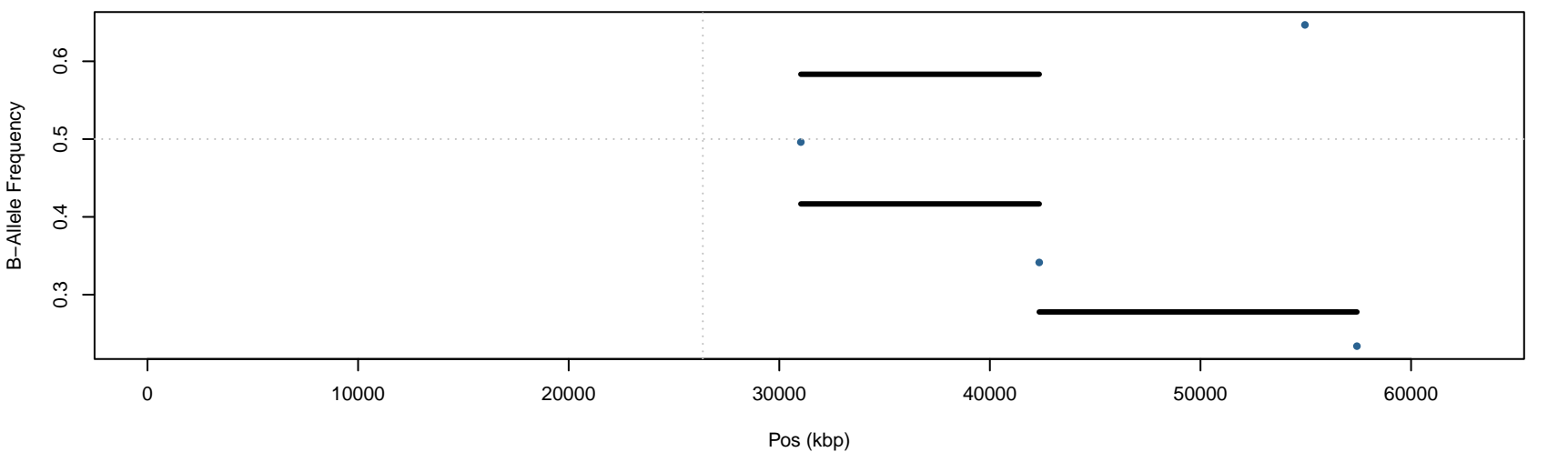
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 17



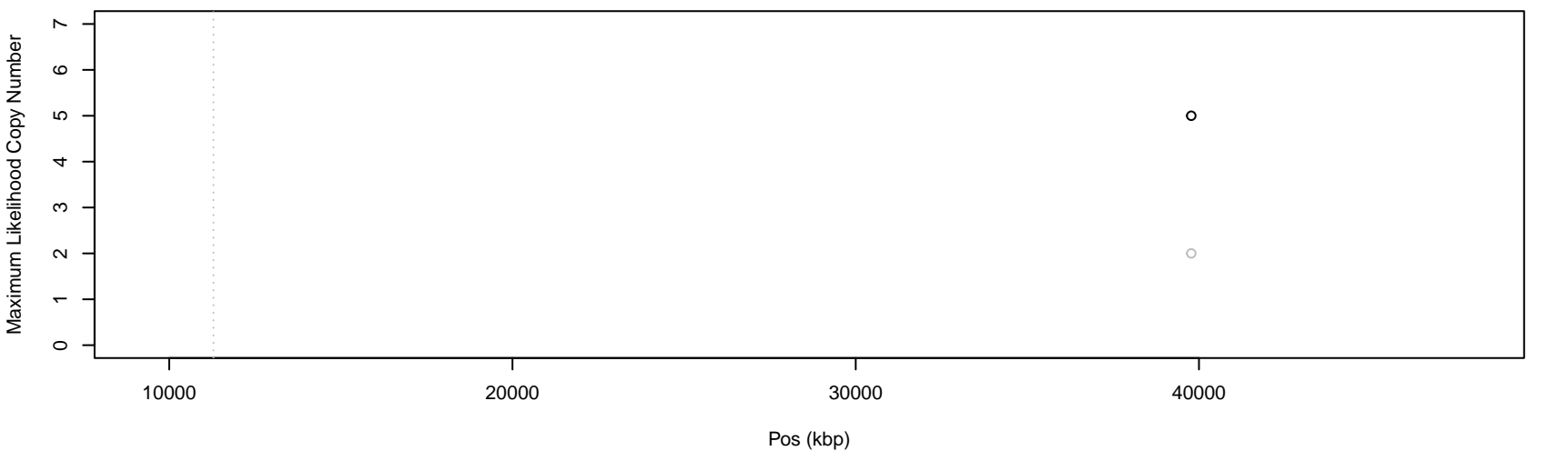
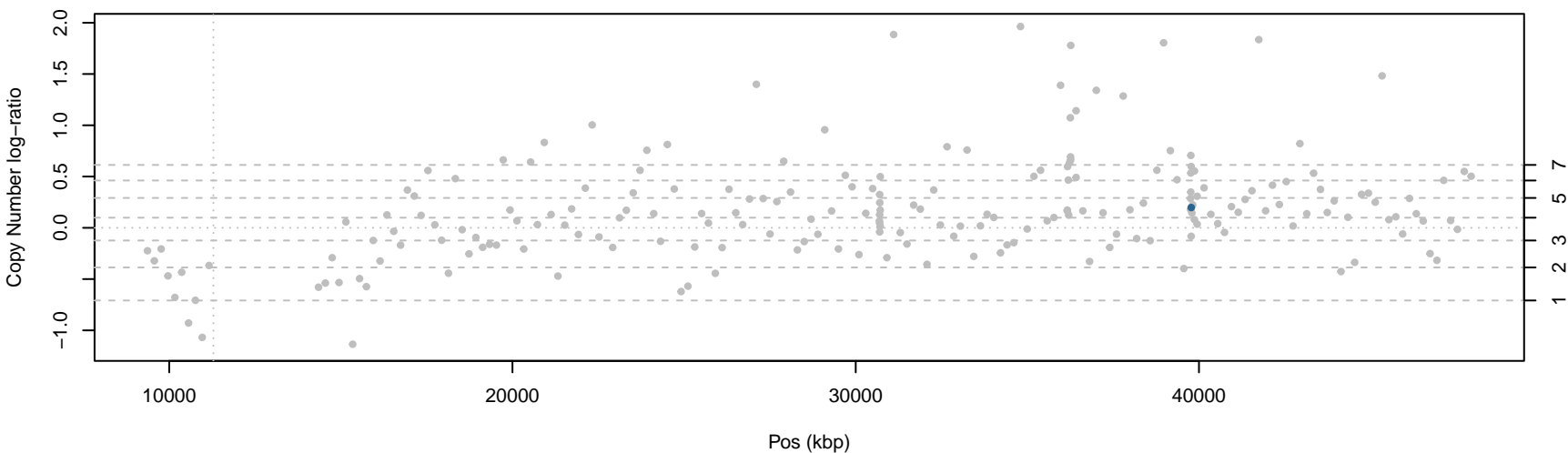
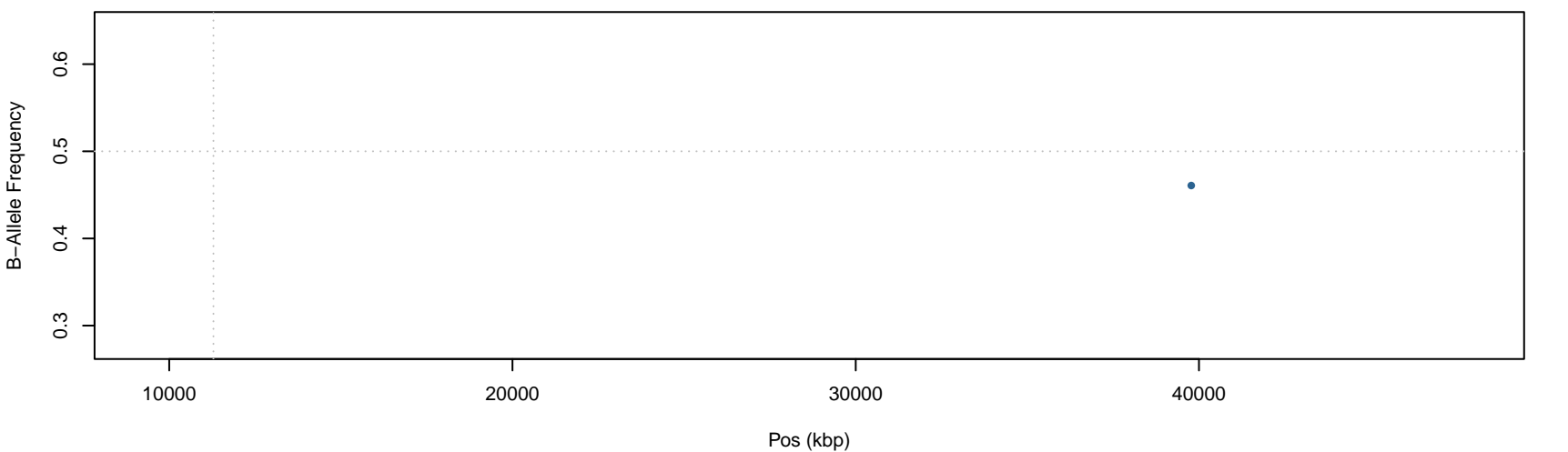
Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 19



Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 20



Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 21



Purity: 0.4 Tumor ploidy: 3.537 SNV log-likelihood: -156.08 GoF: 88.3% Mean coverage: 276;465 Chromosome: 22

