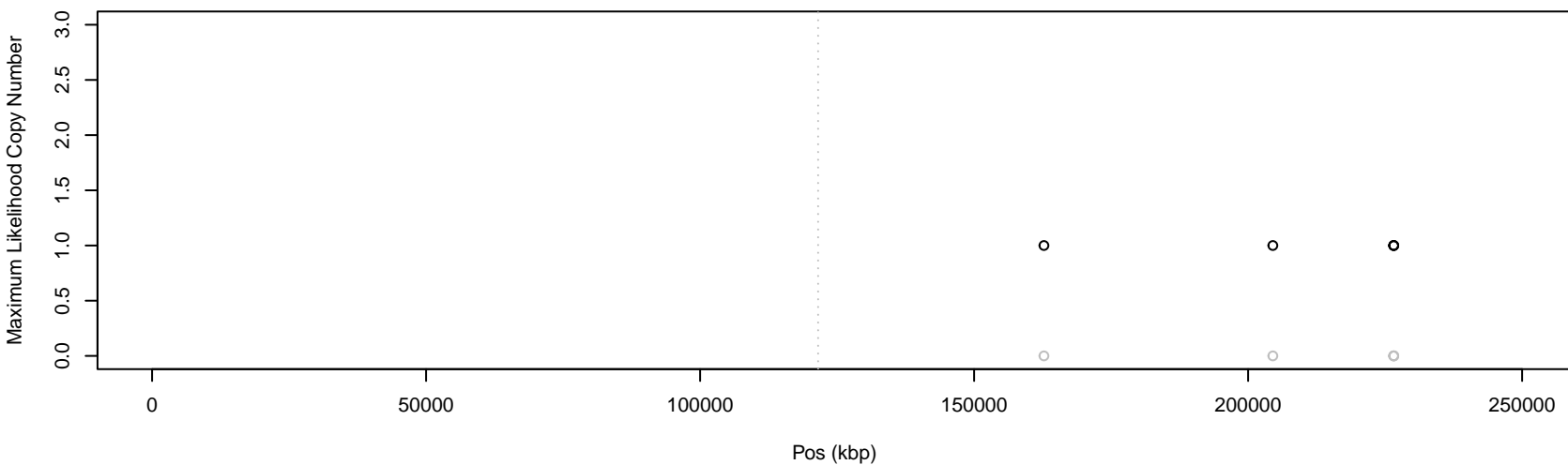
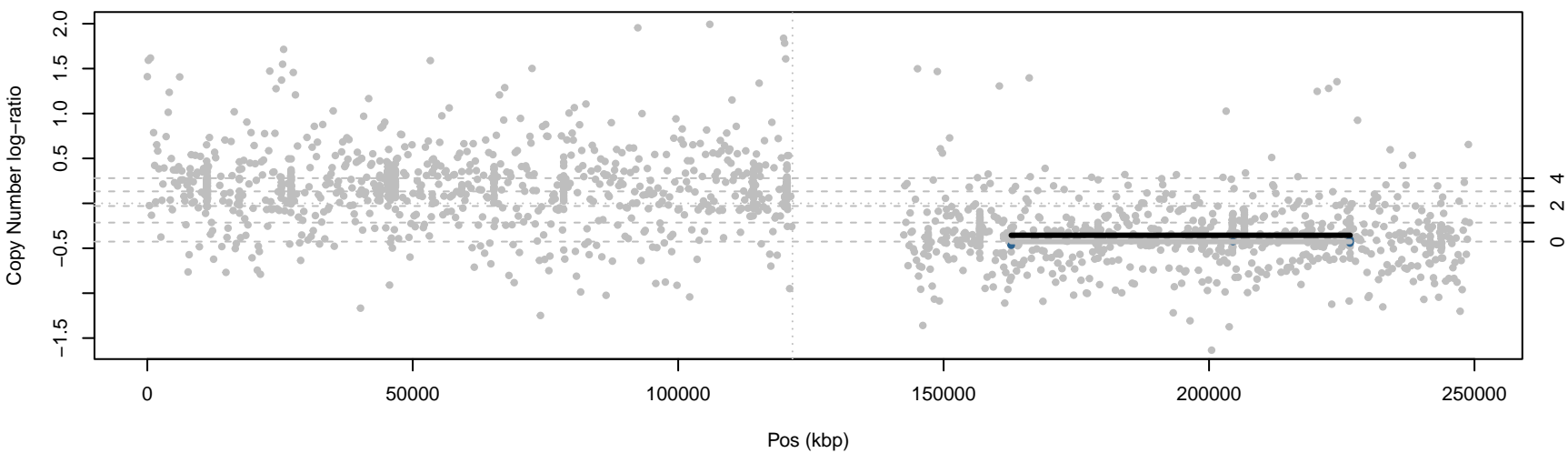
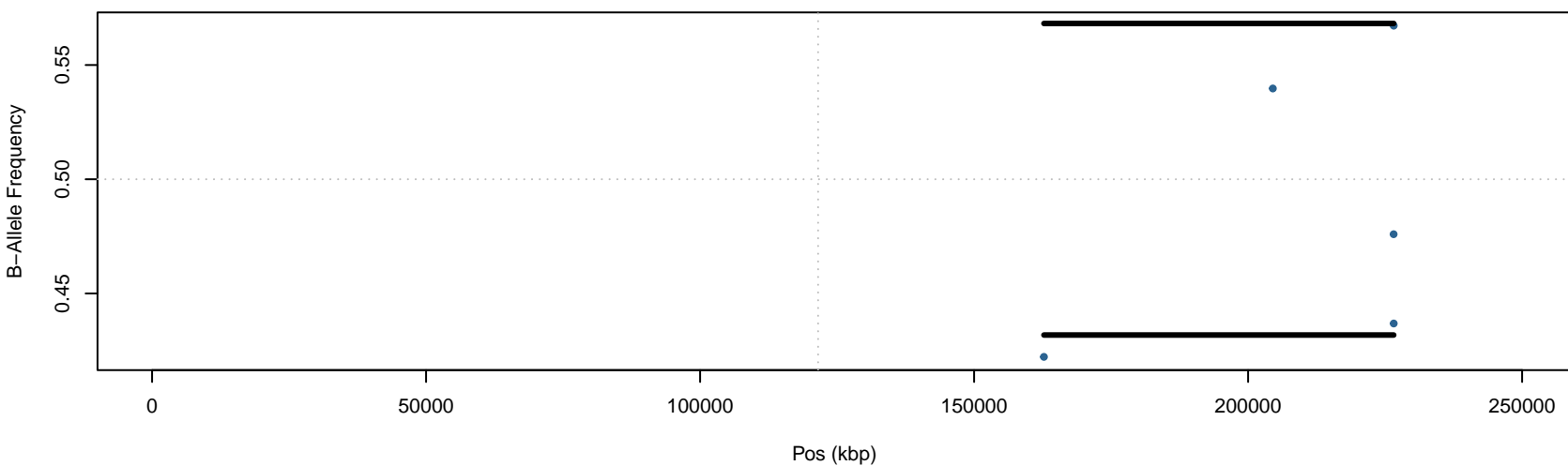
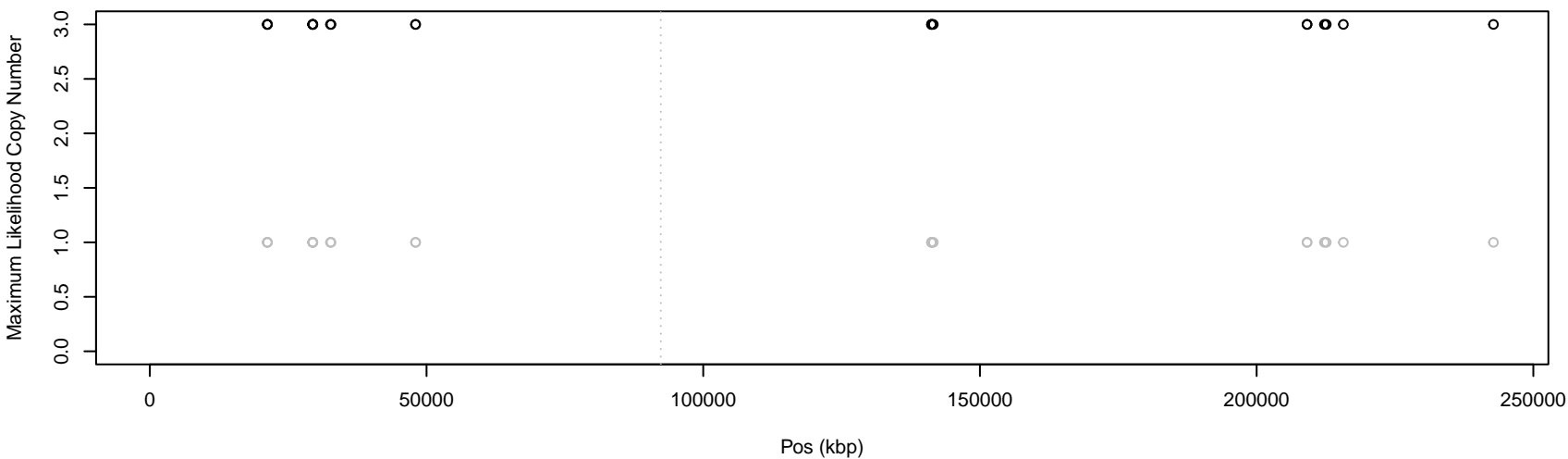
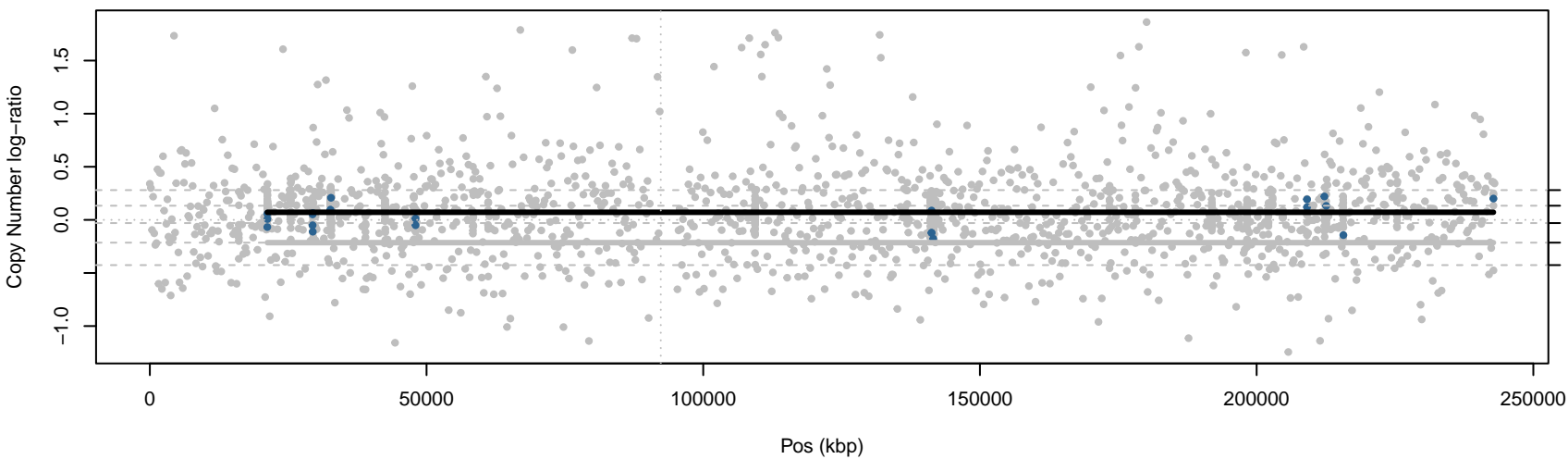
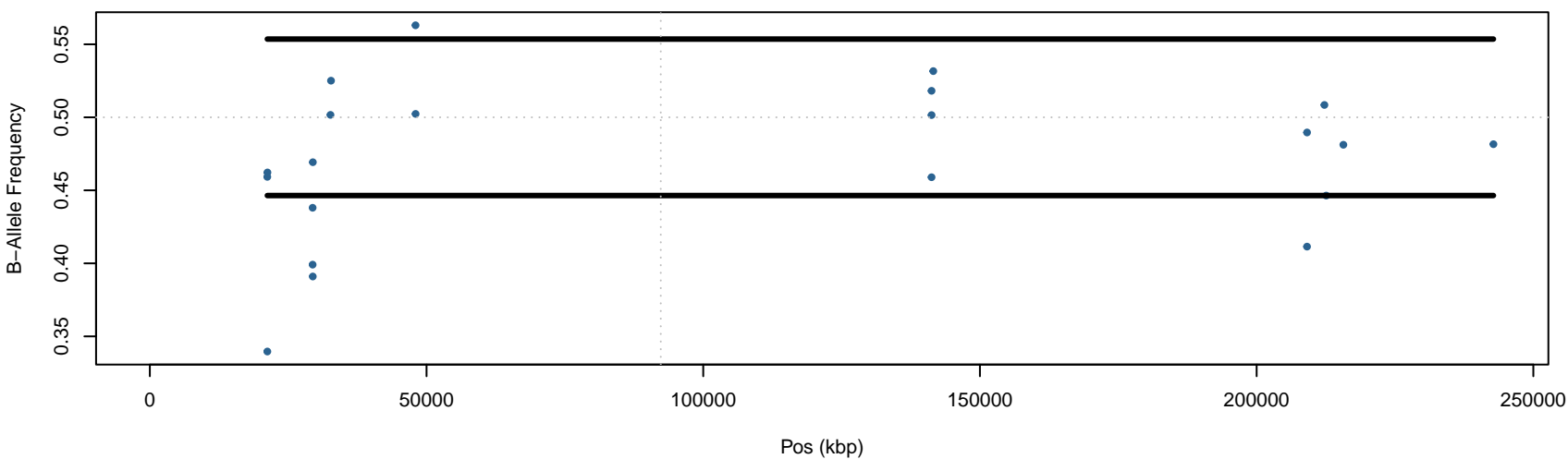


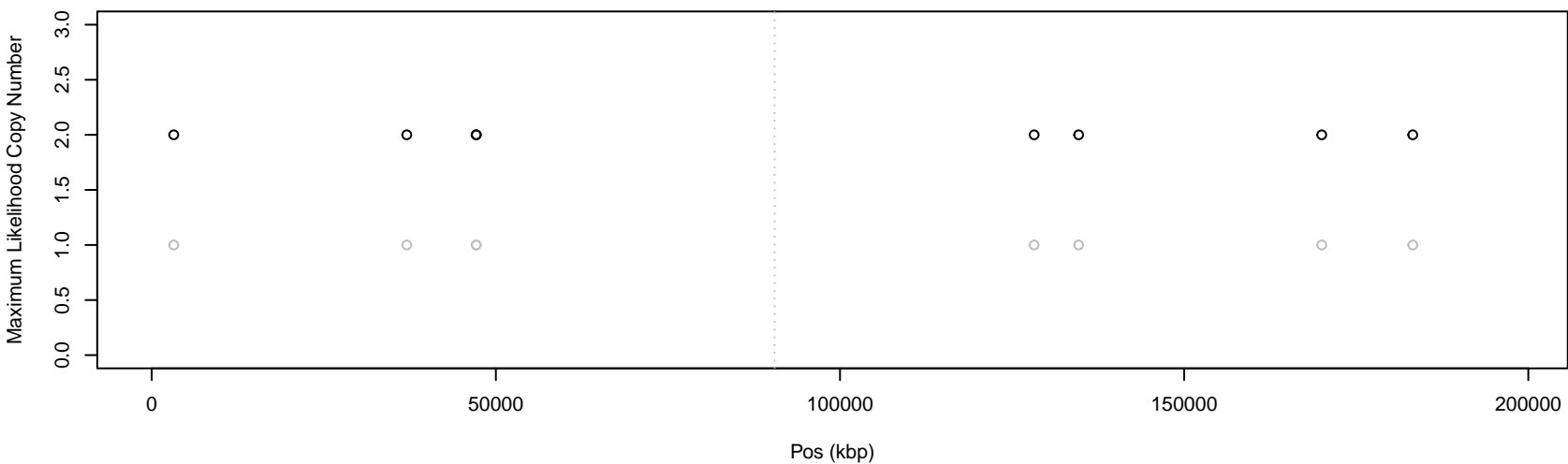
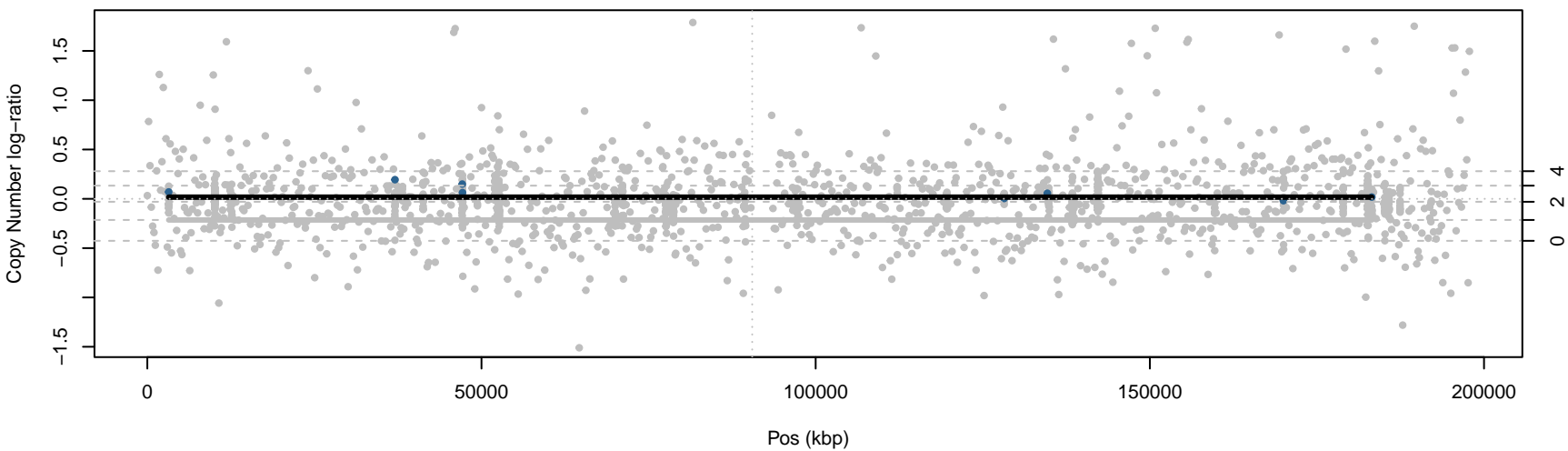
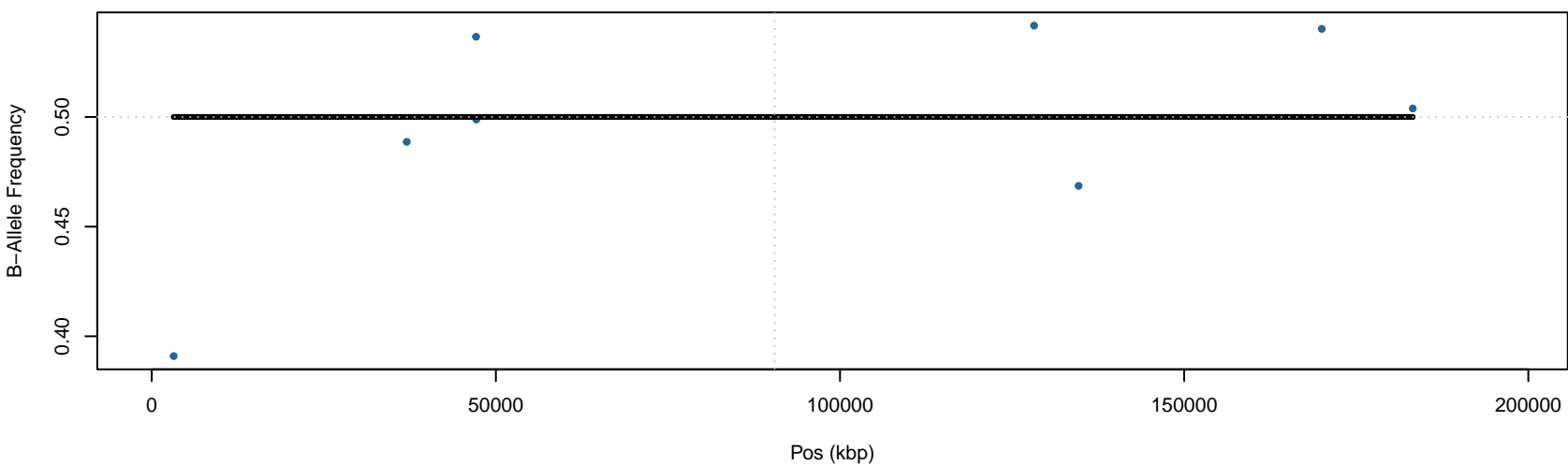
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 1



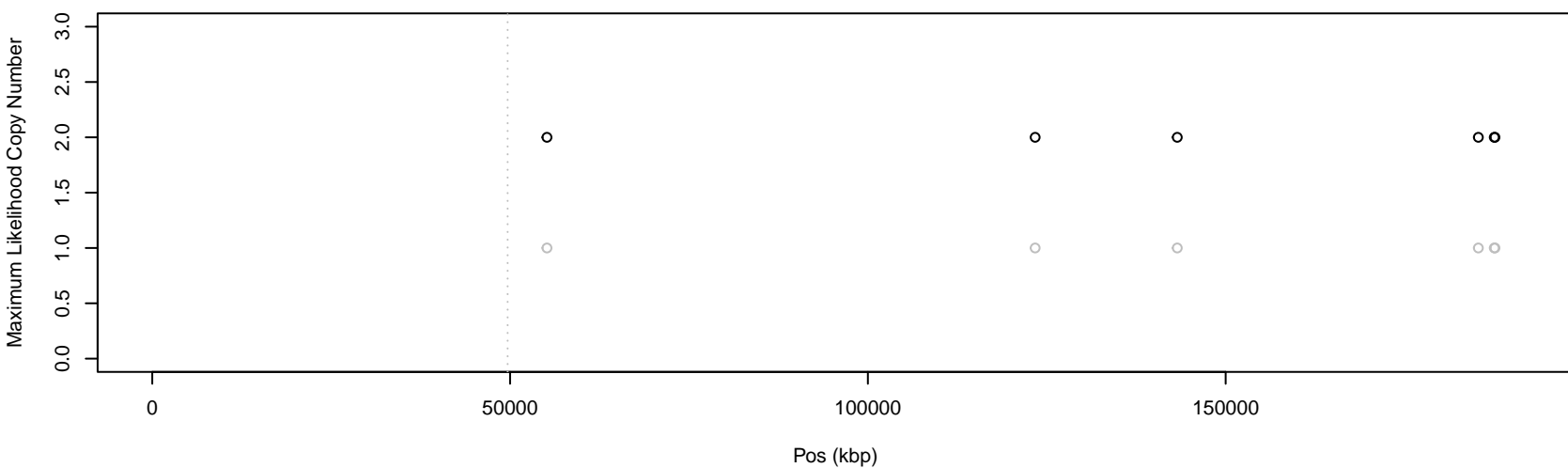
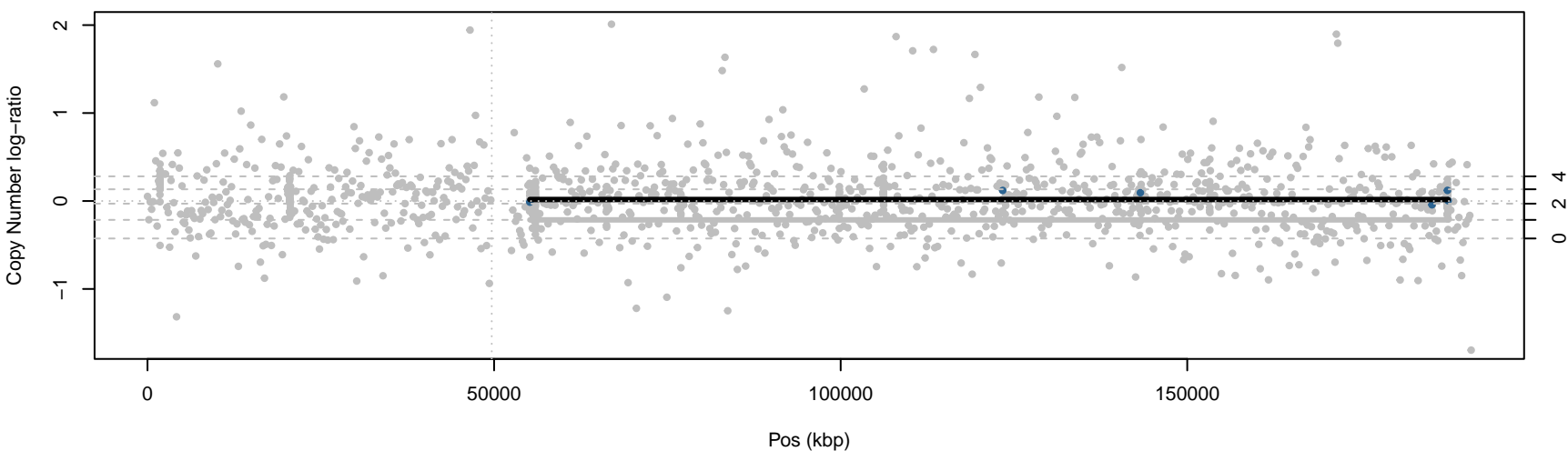
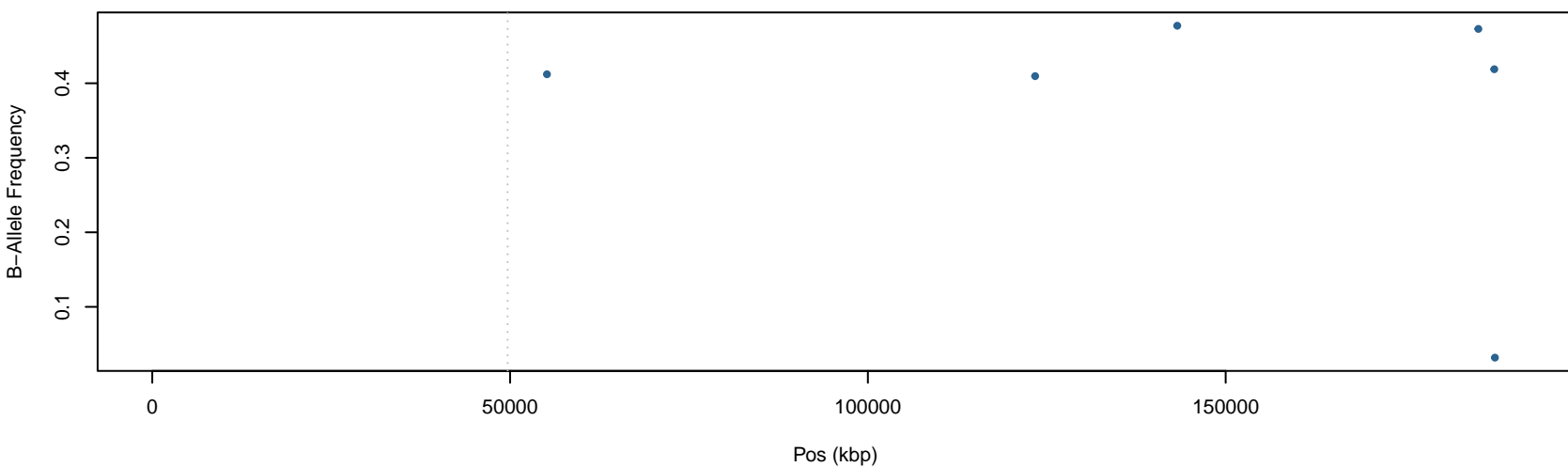
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 2



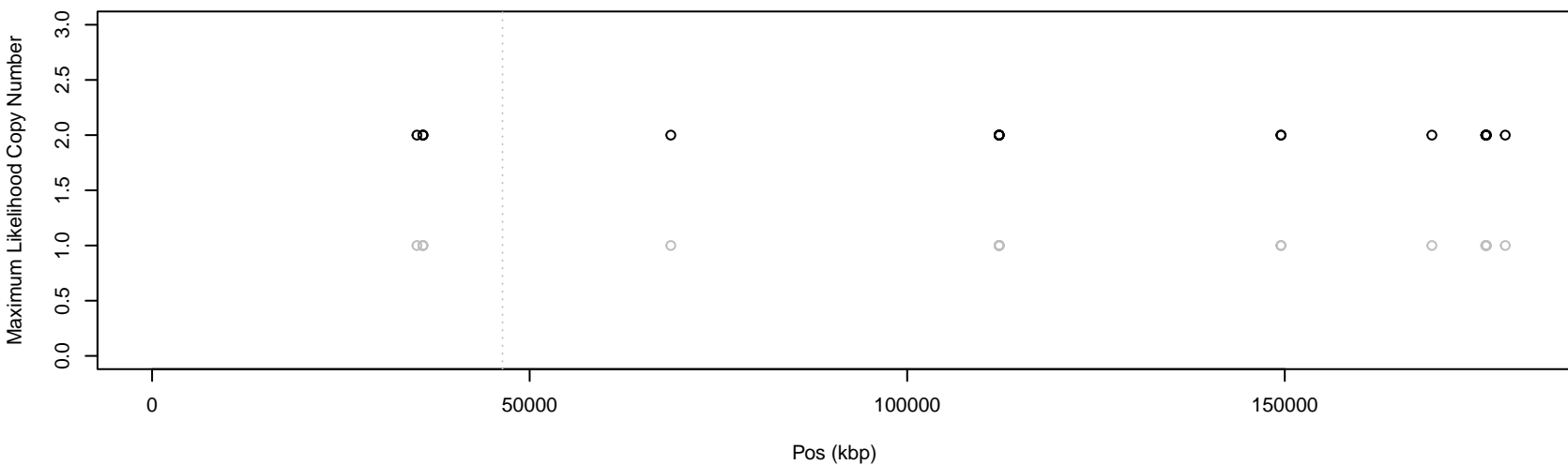
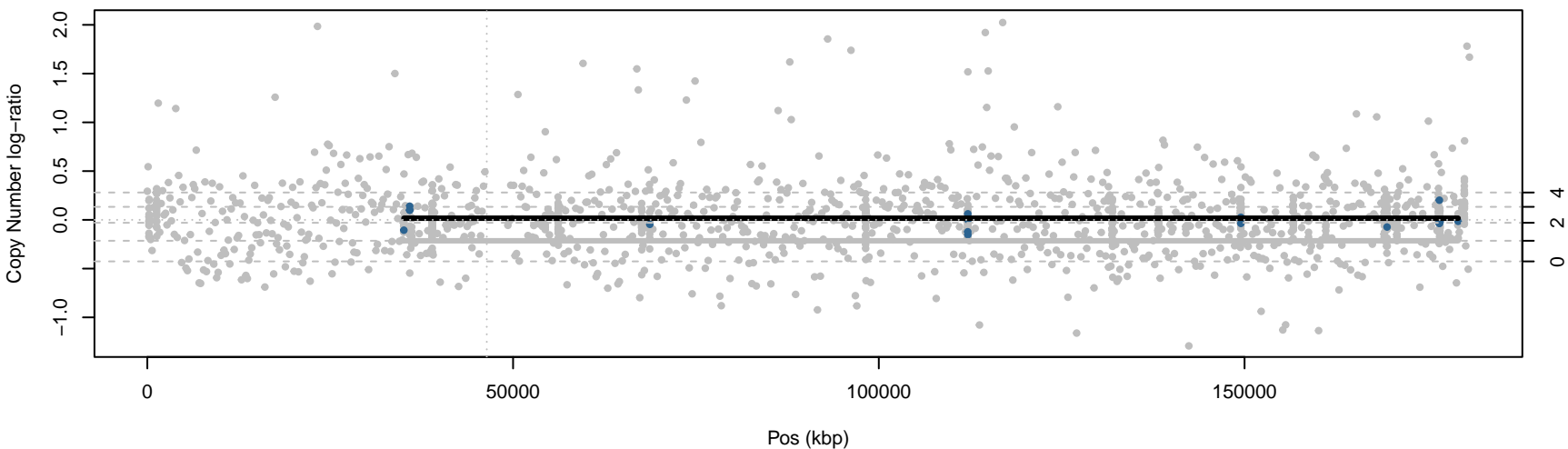
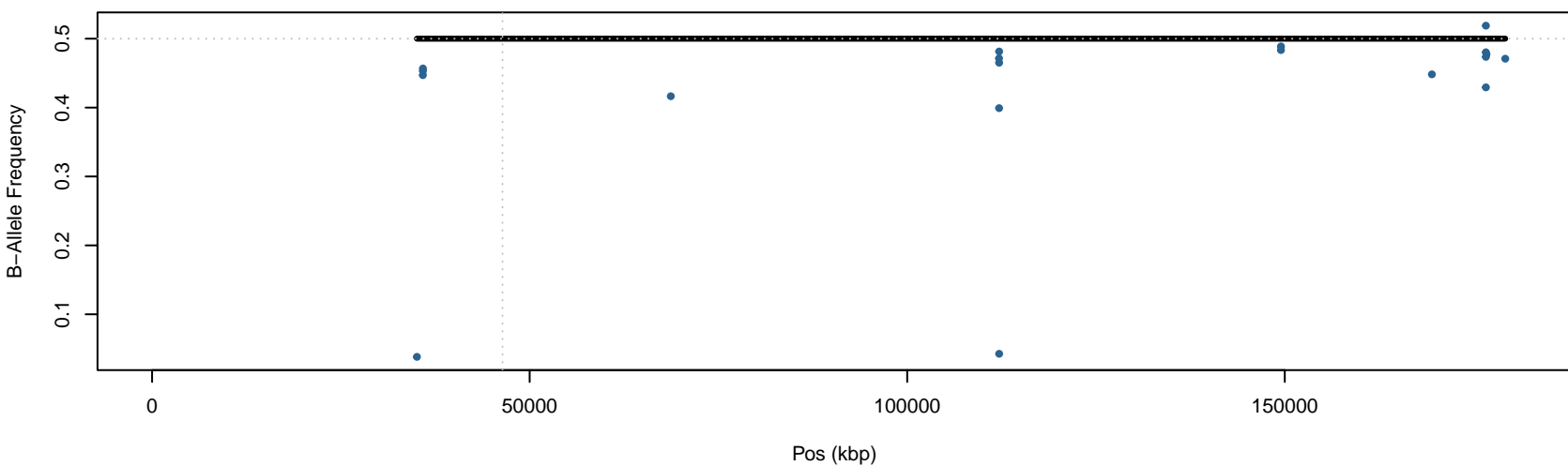
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 3



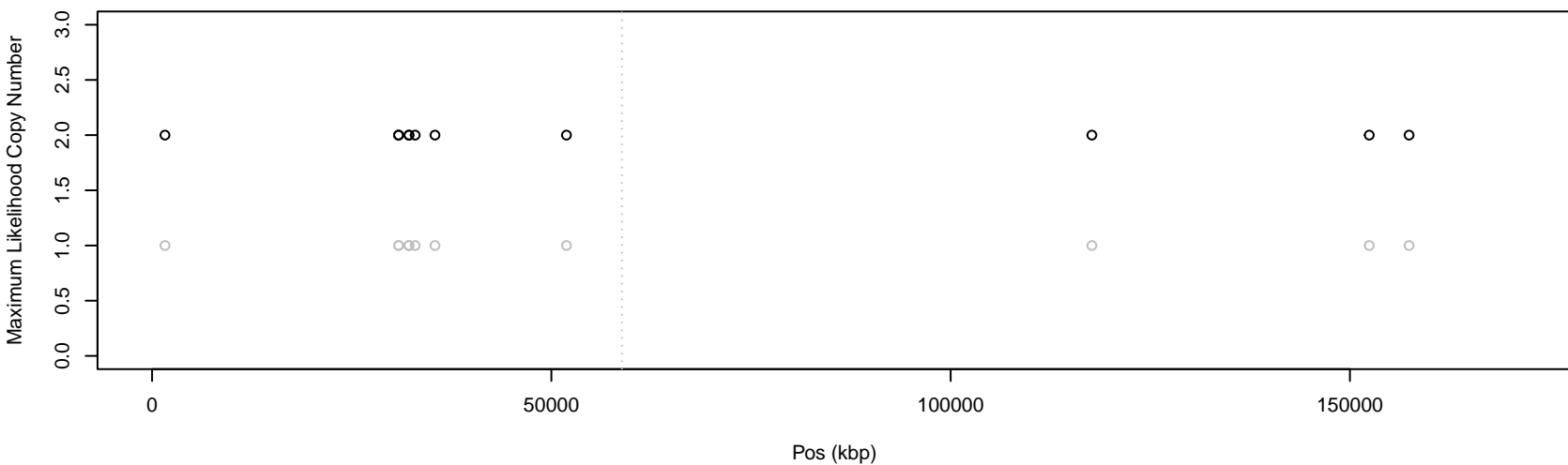
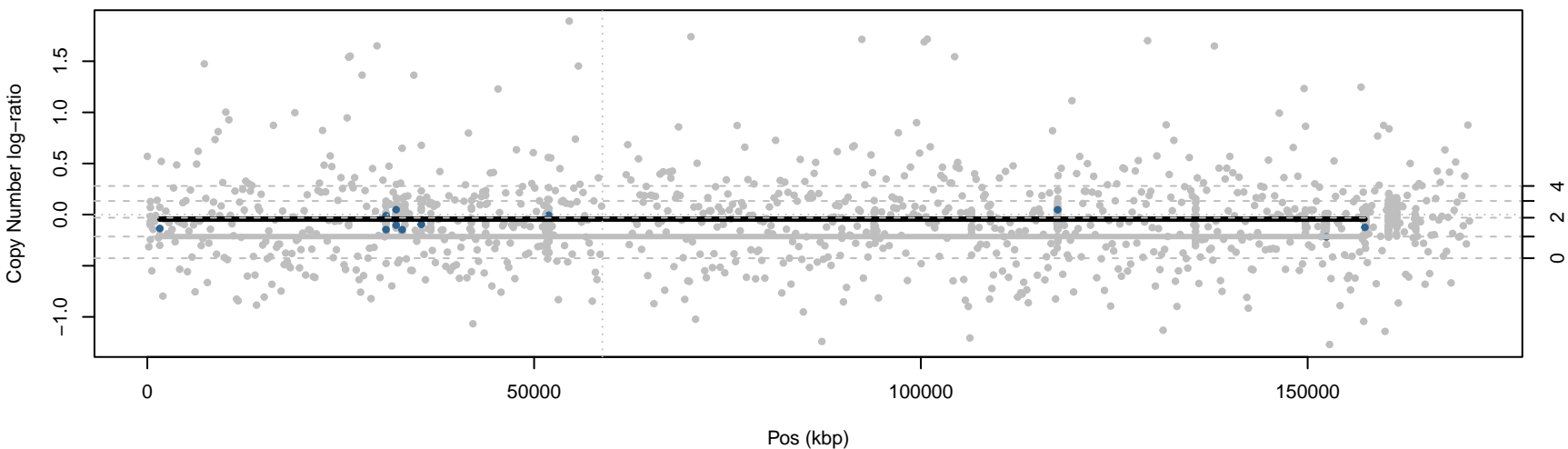
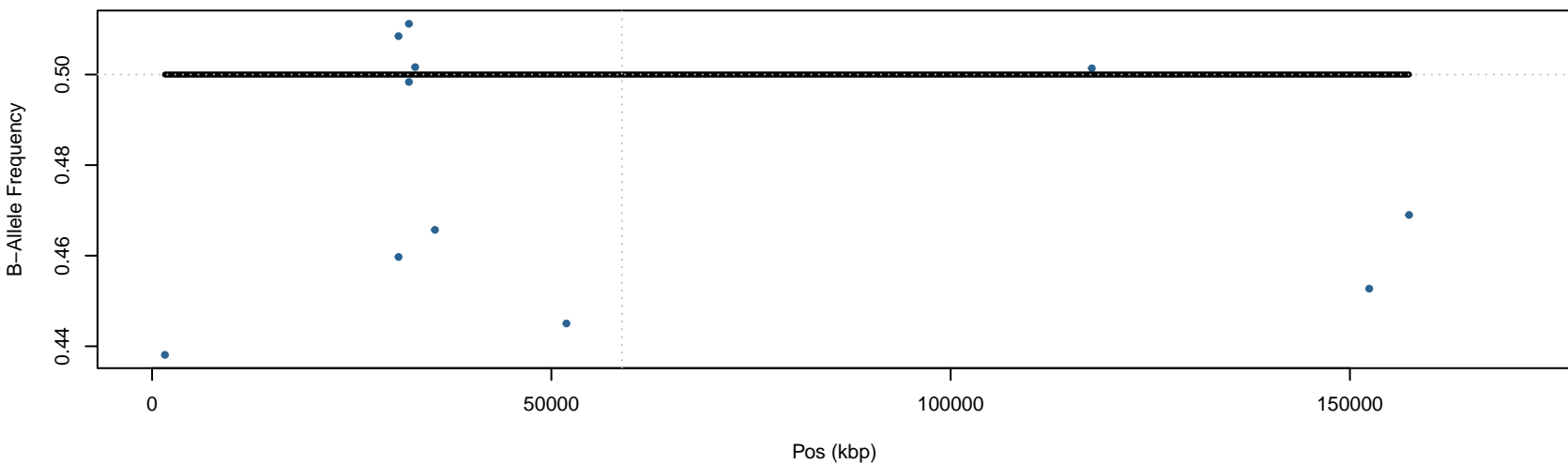
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565,782 Chromosome: 4



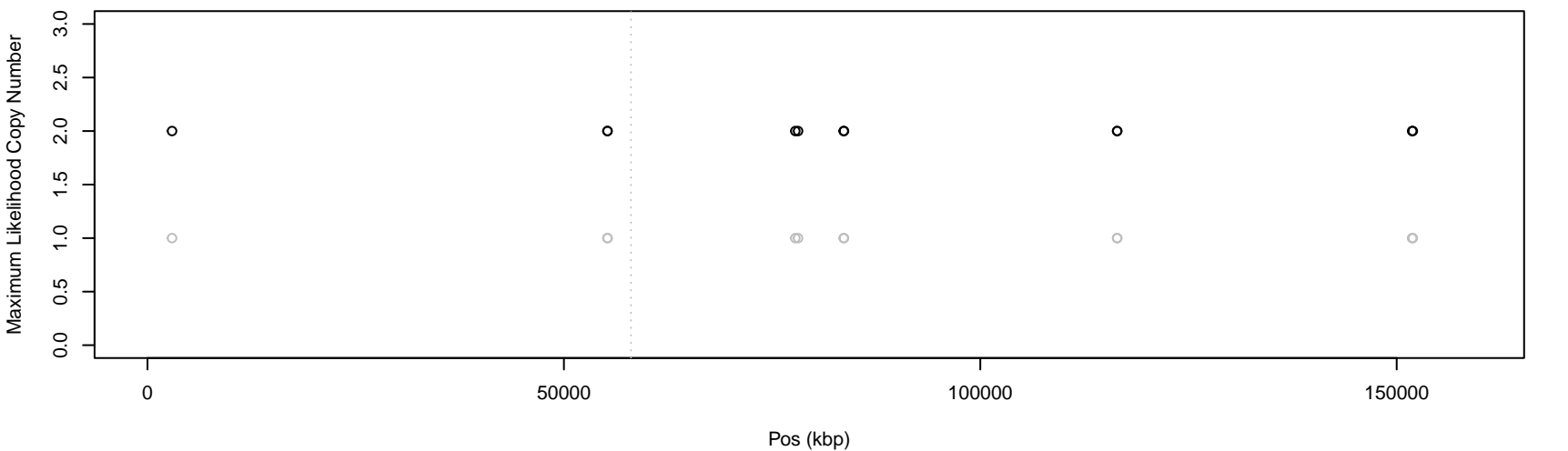
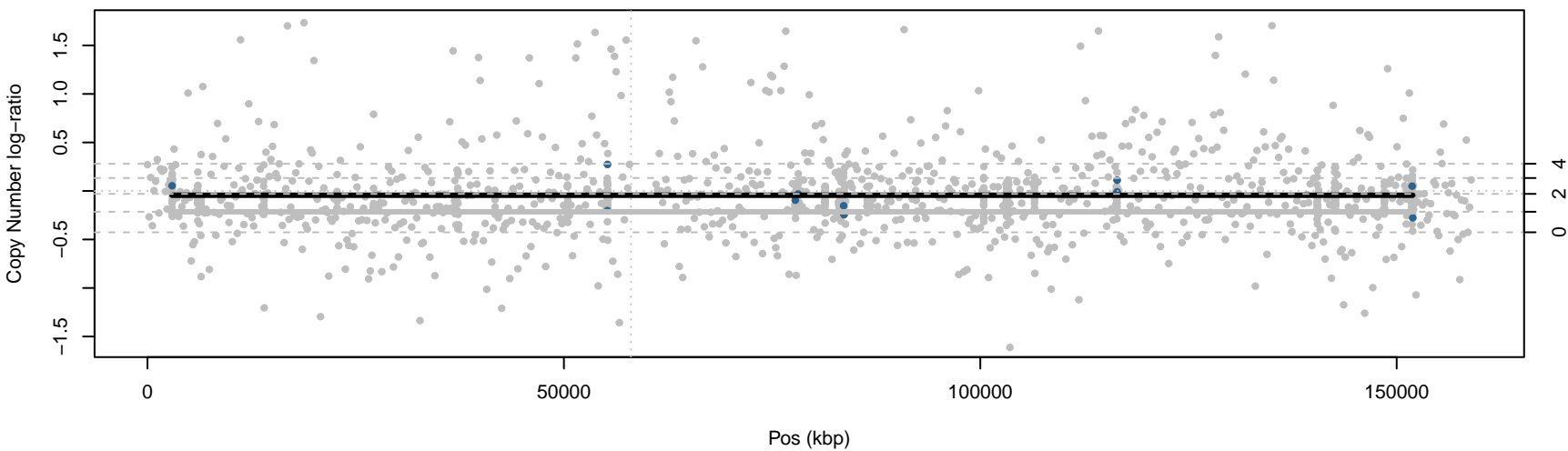
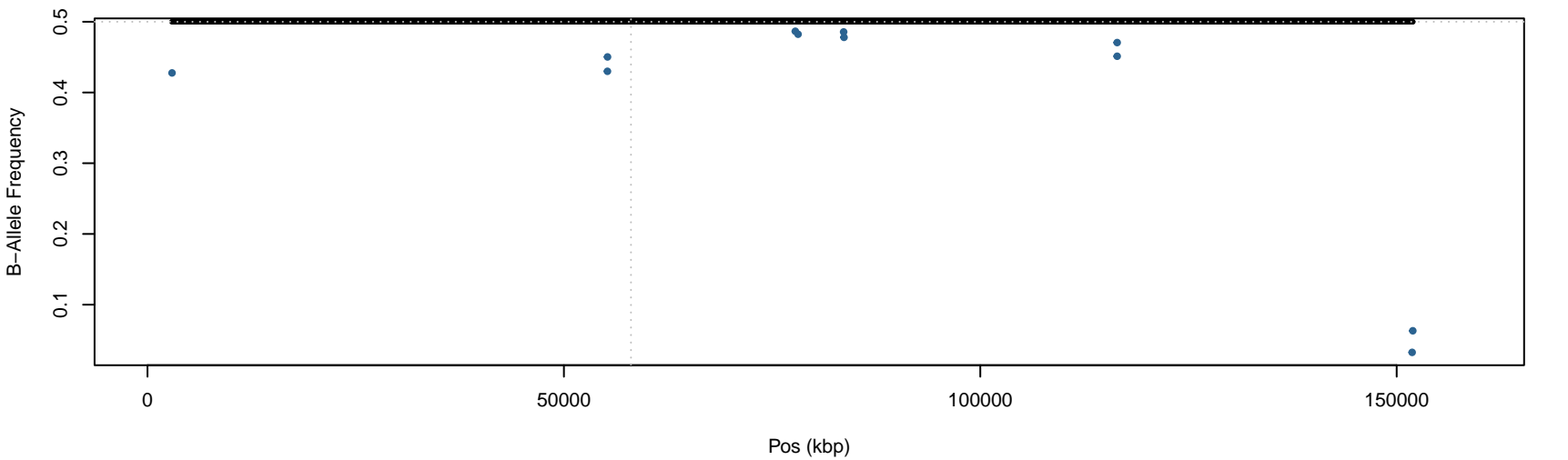
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 5



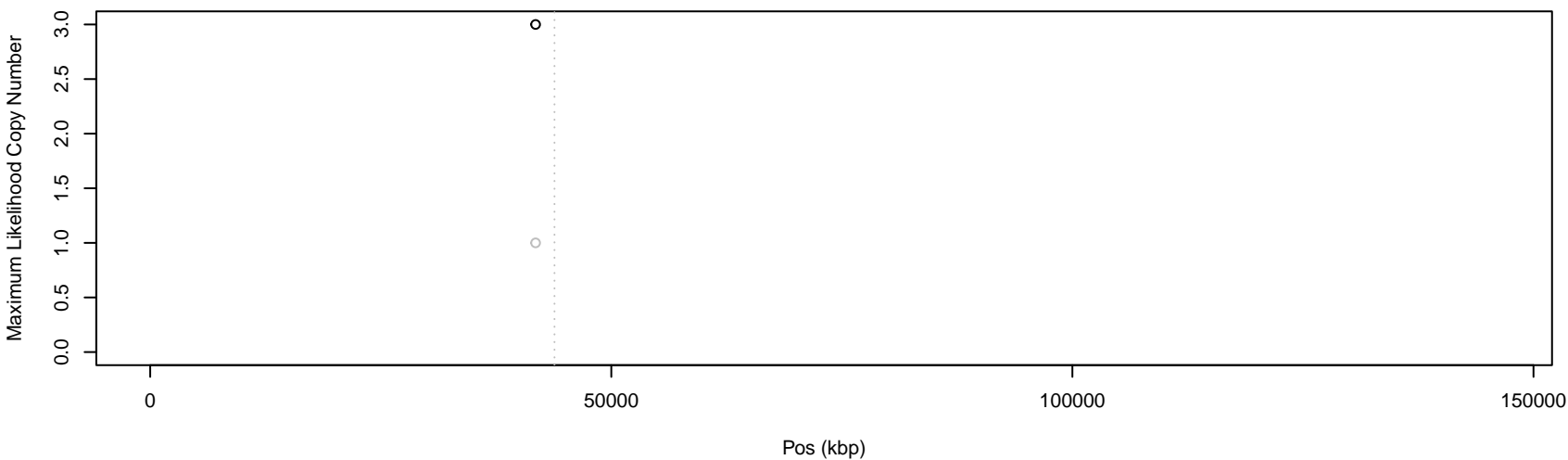
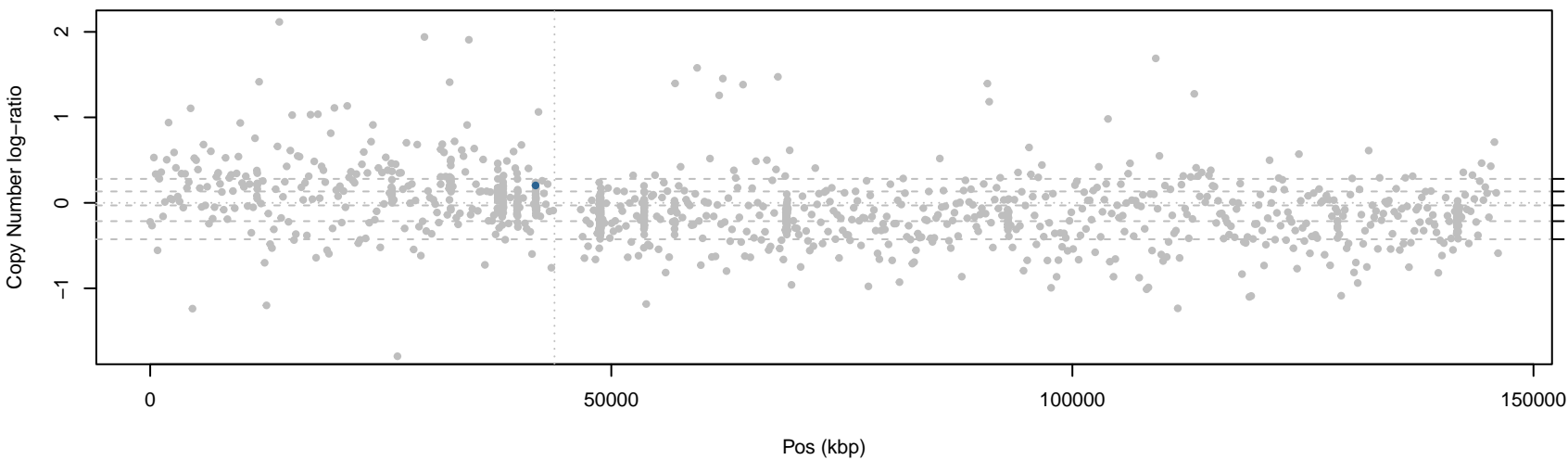
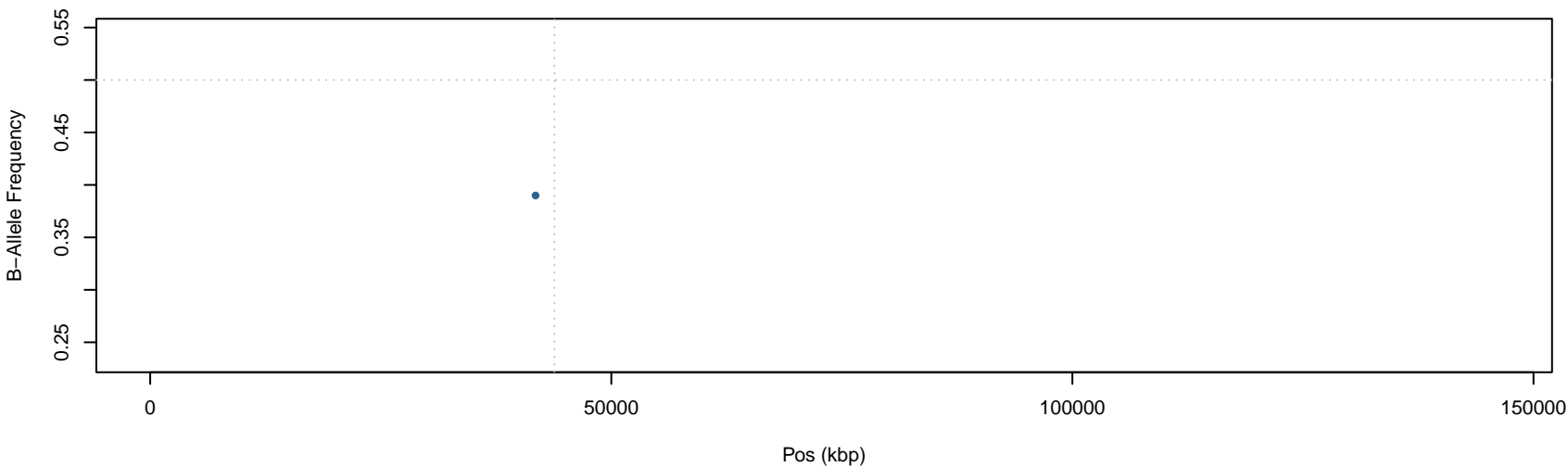
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 6



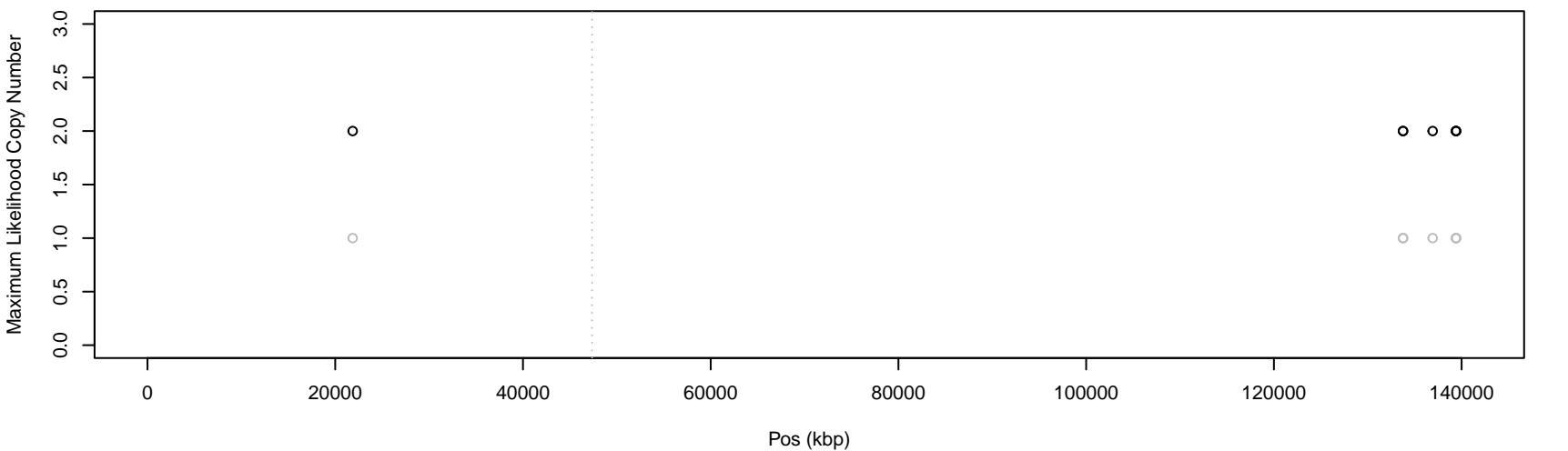
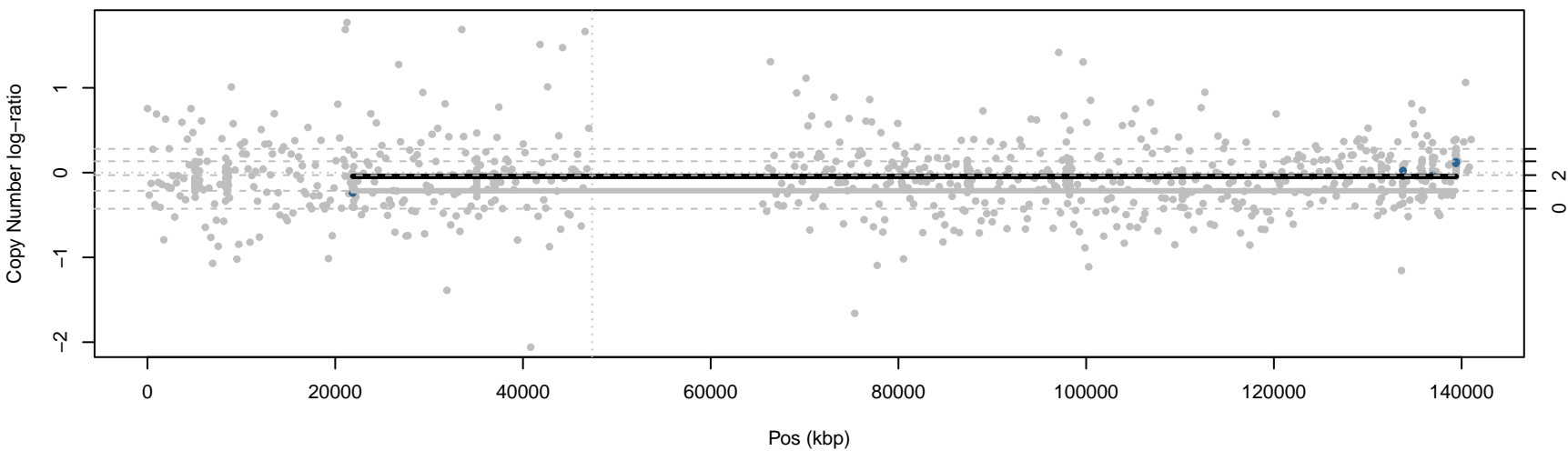
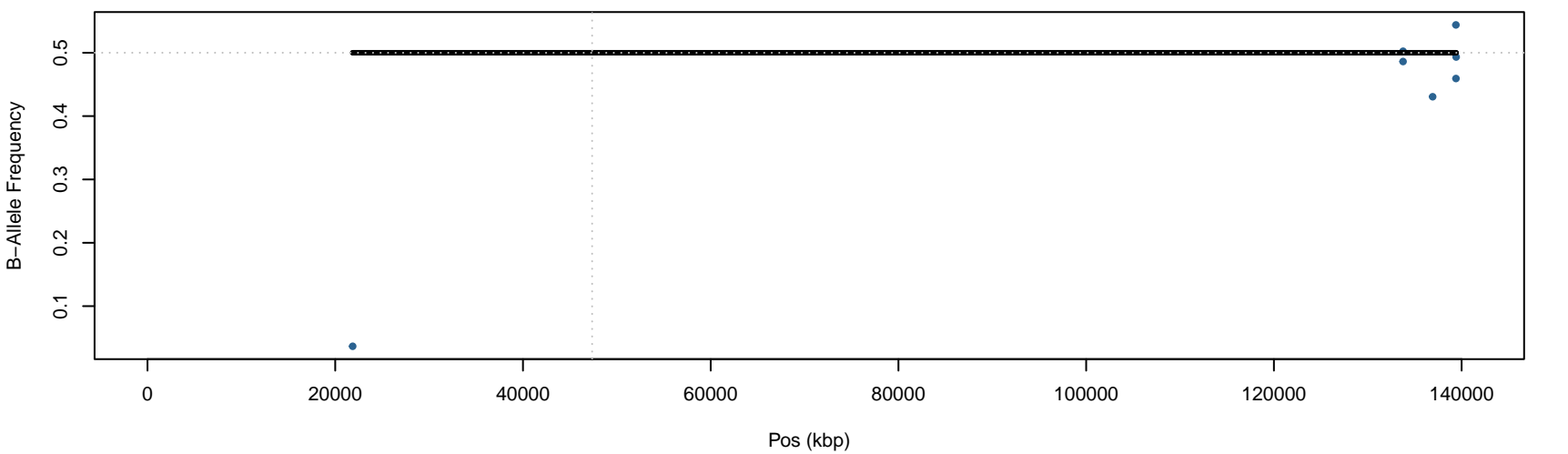
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 7



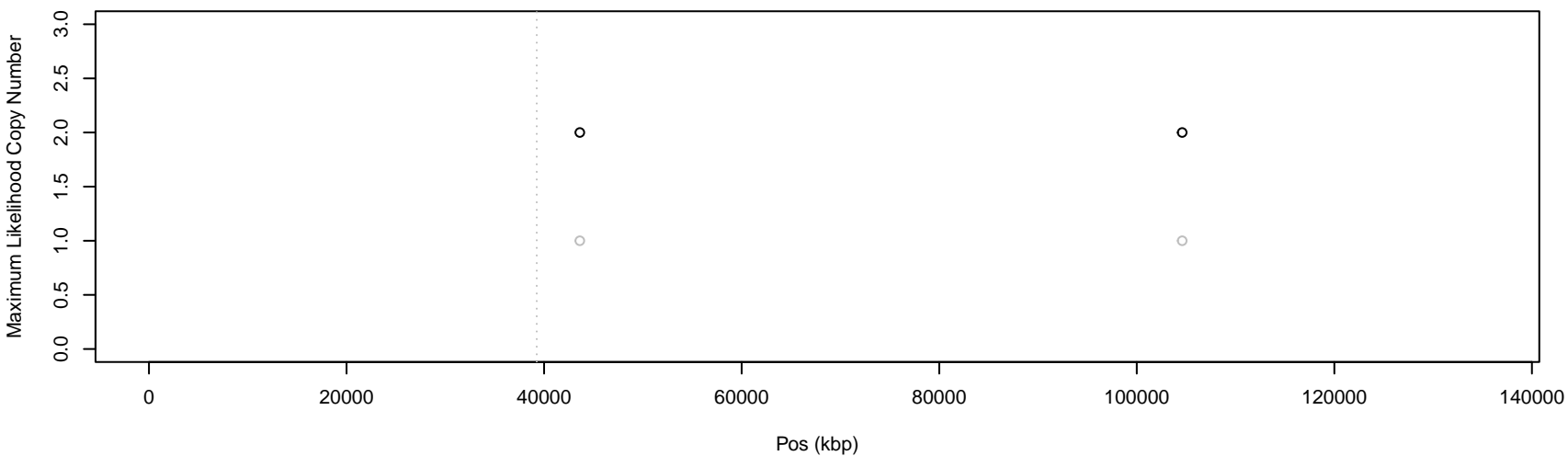
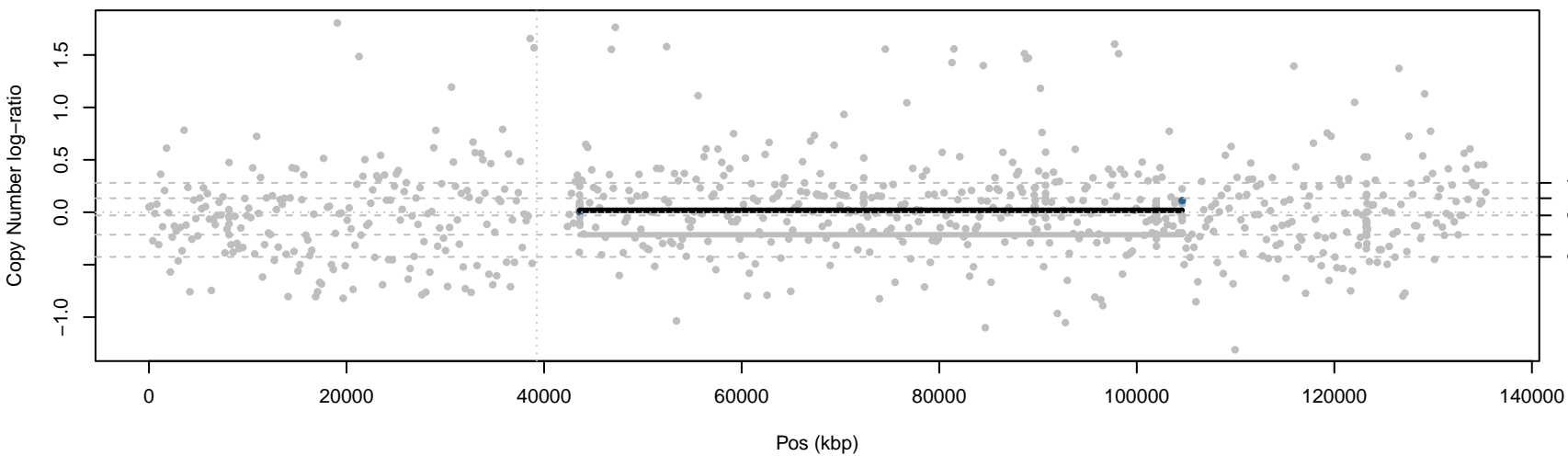
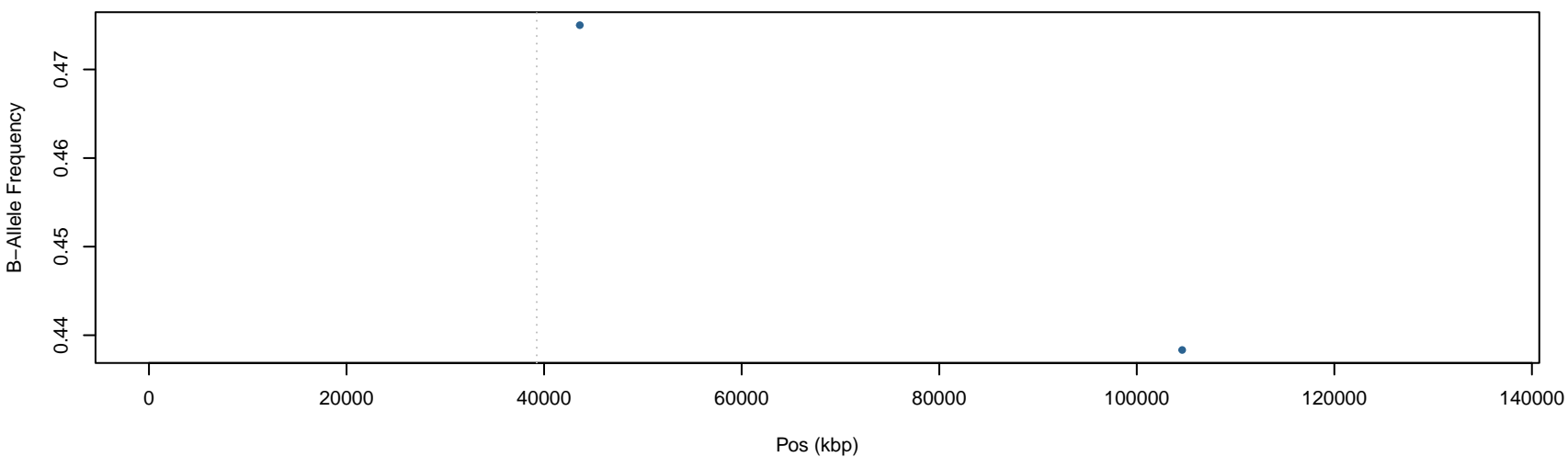
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 8



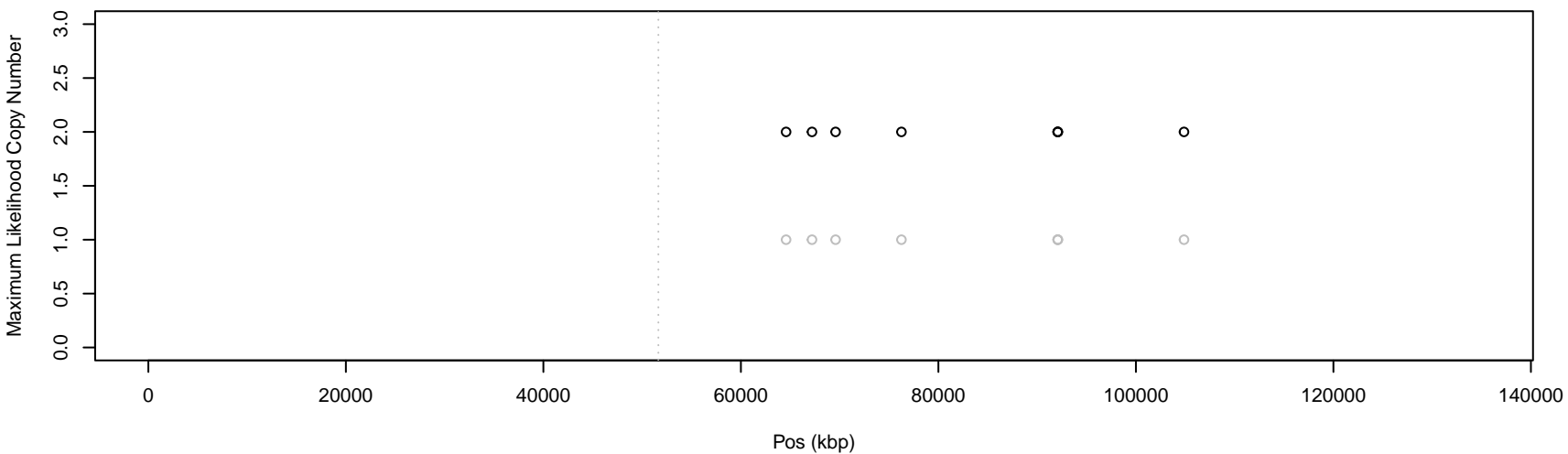
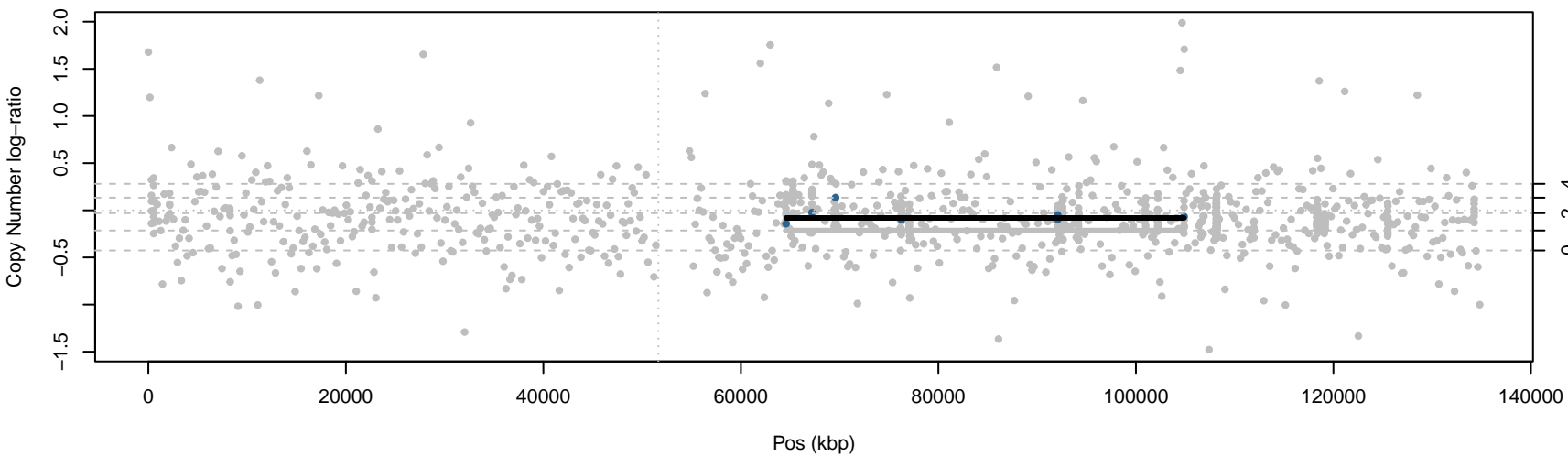
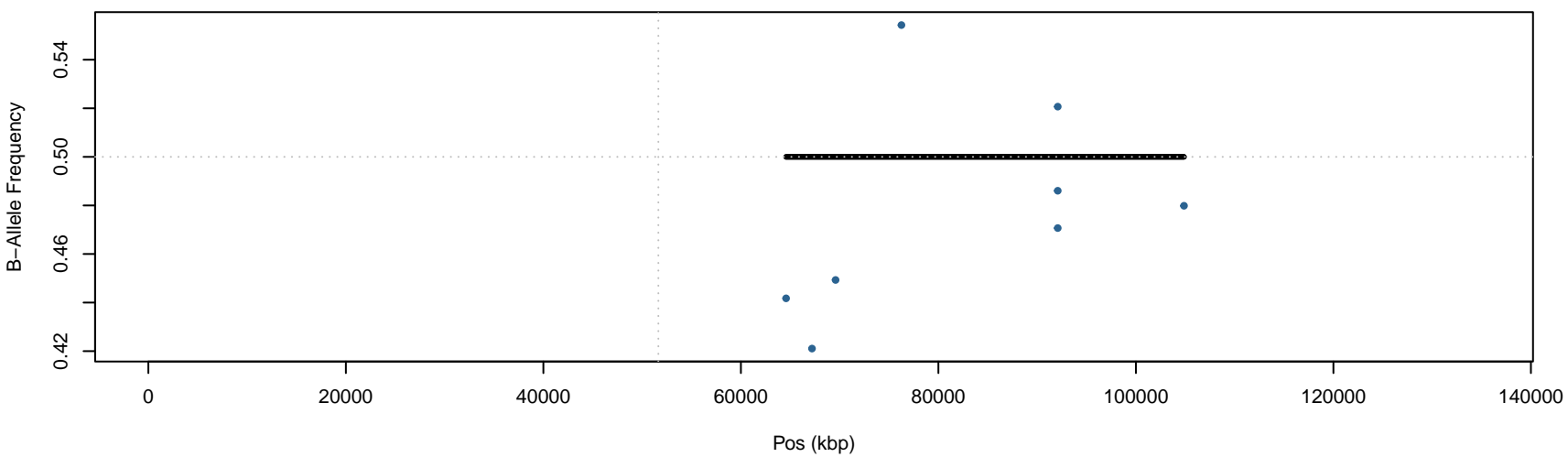
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 9



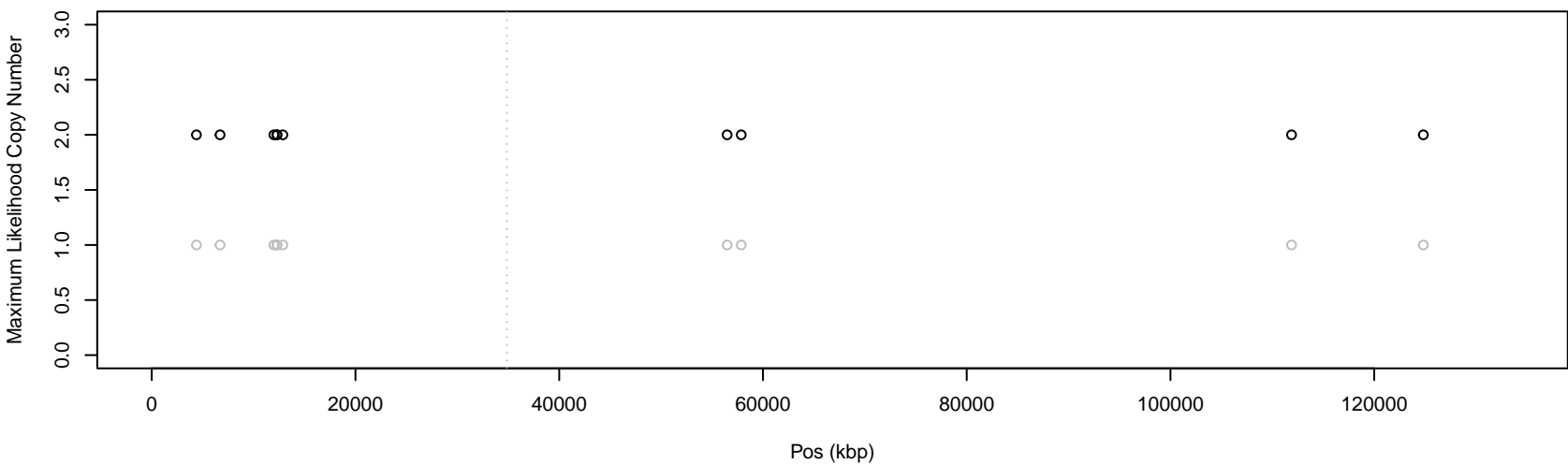
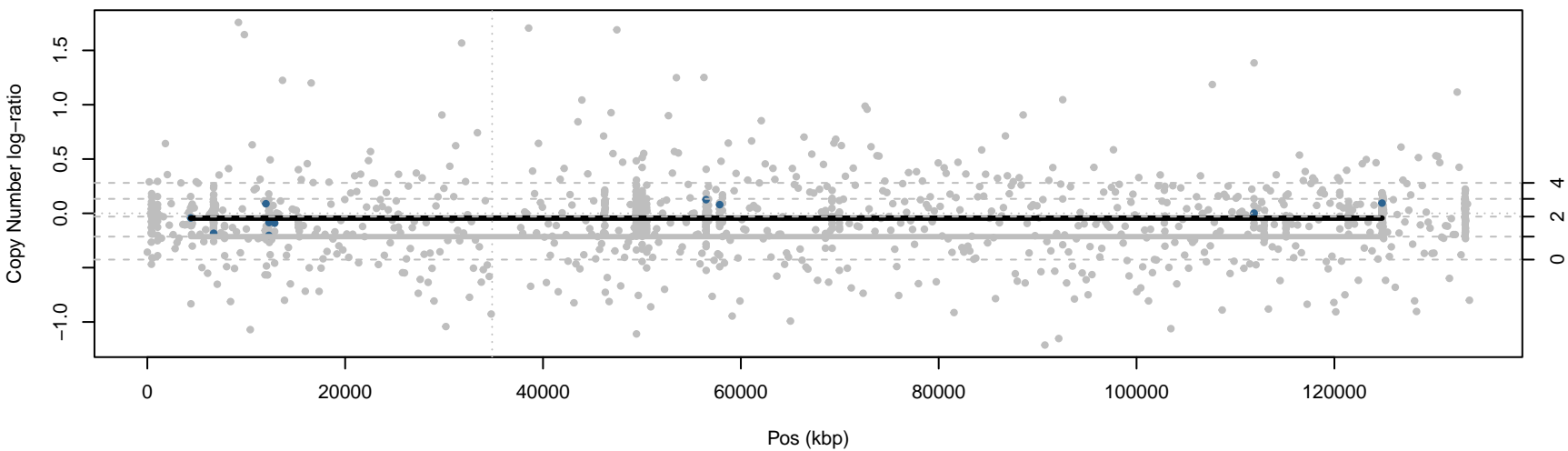
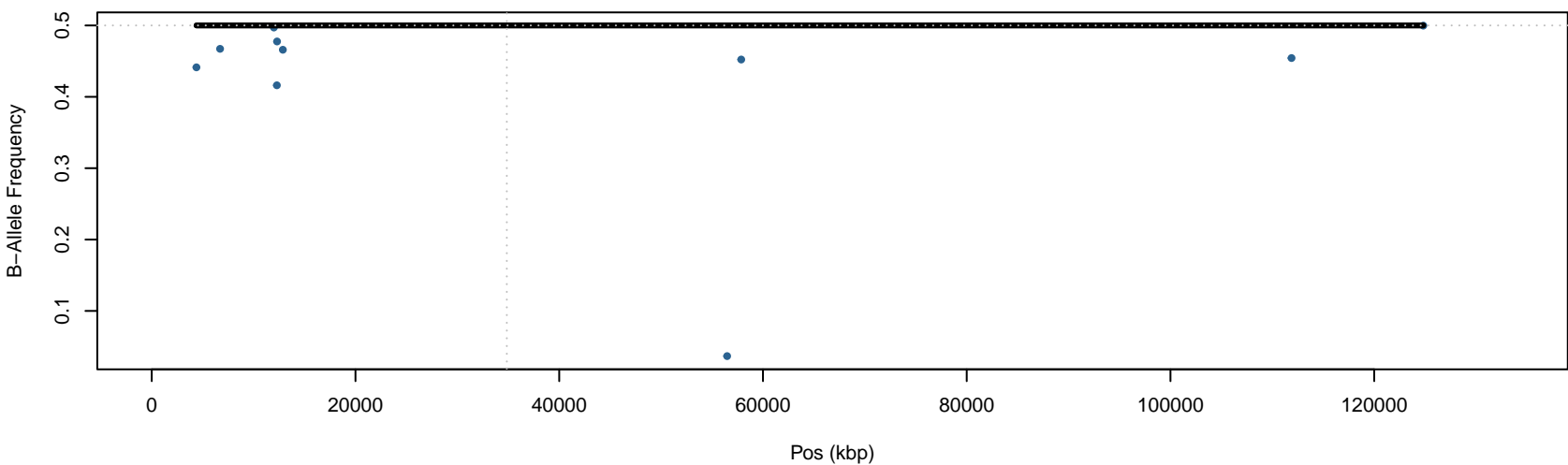
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 10



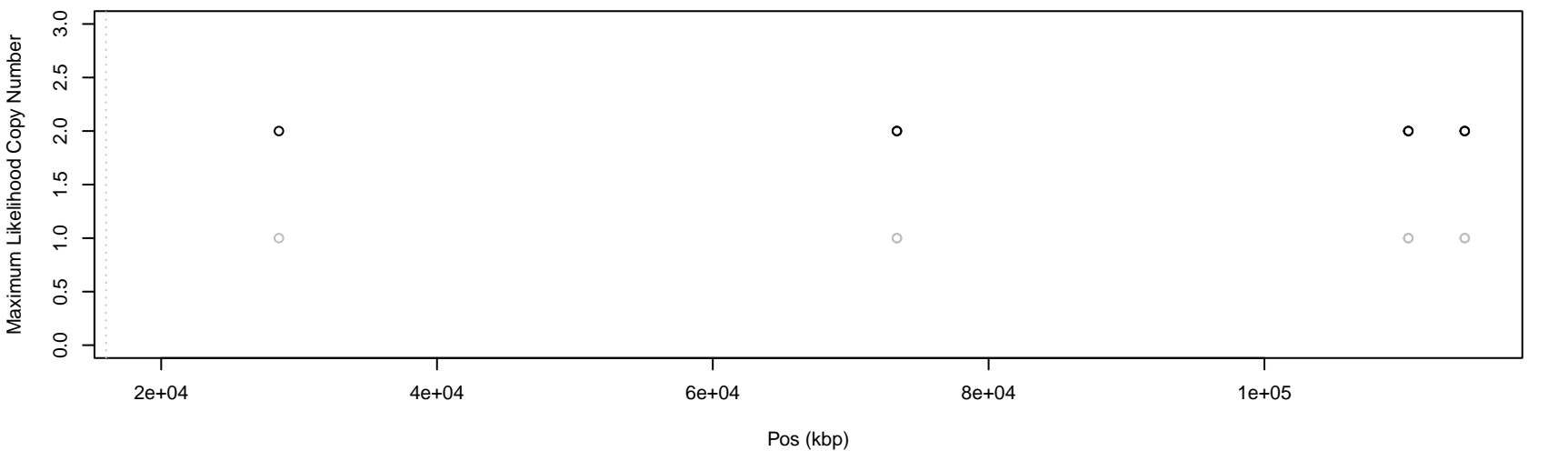
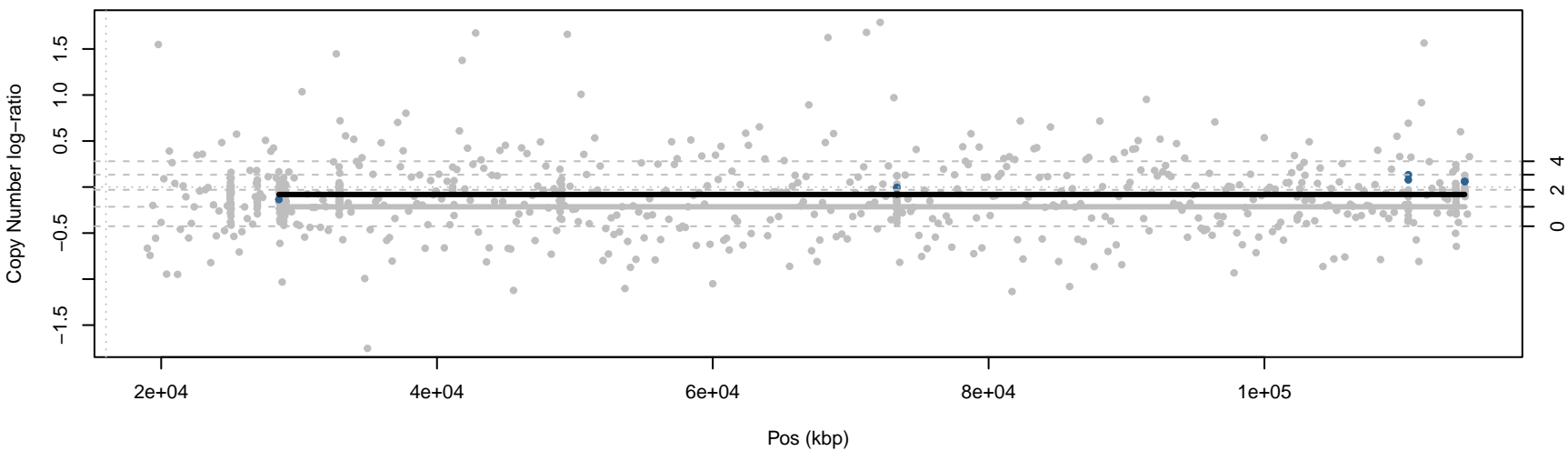
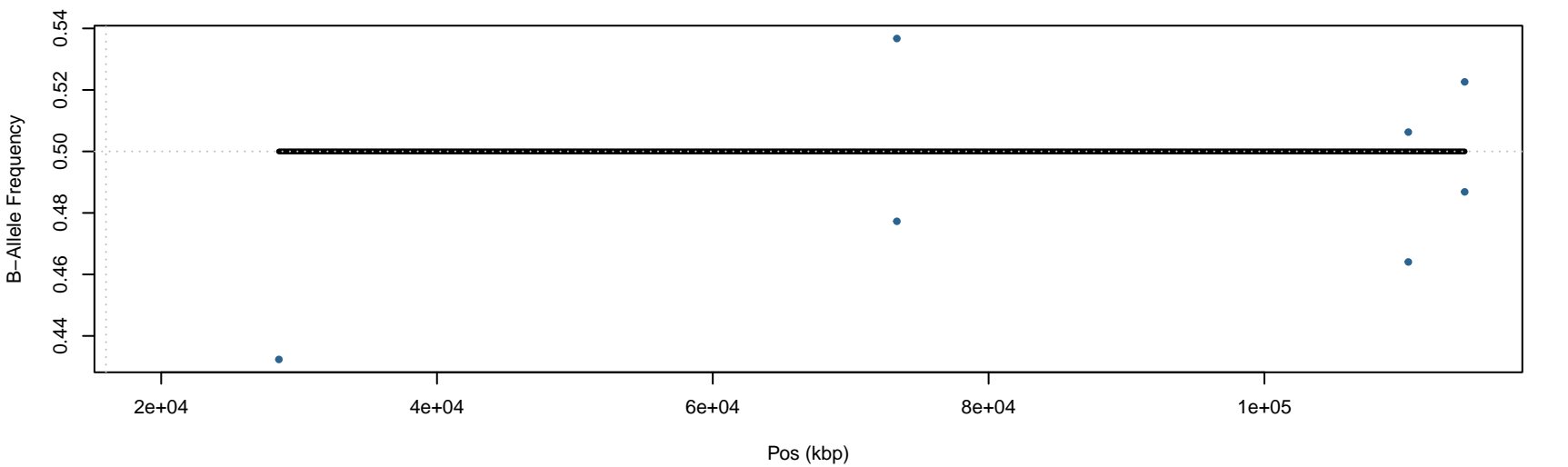
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 11



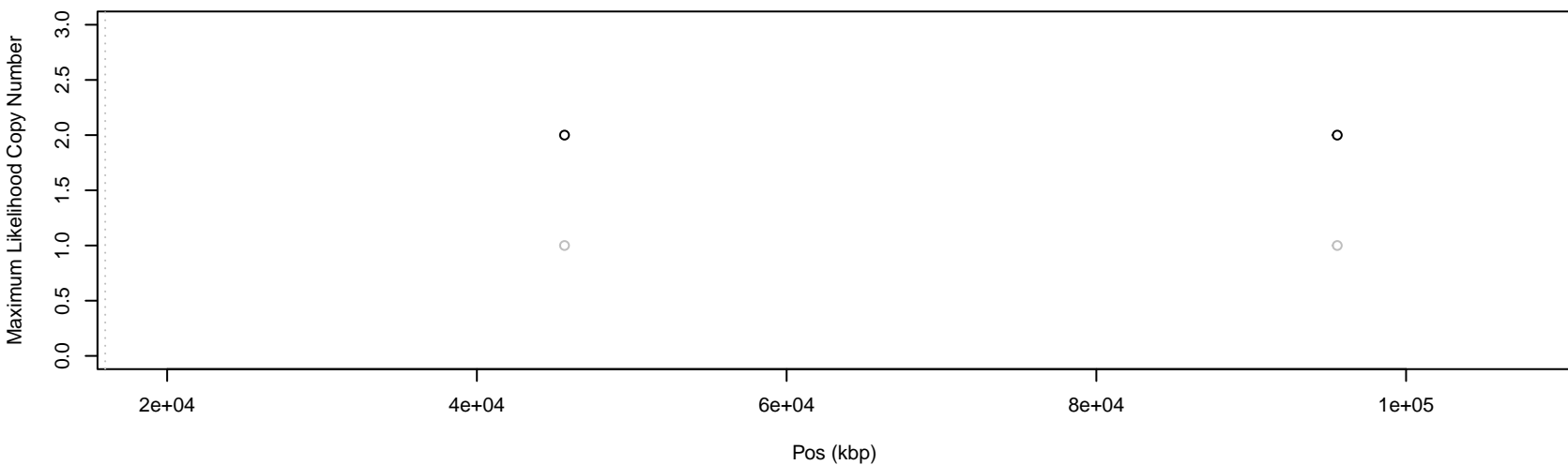
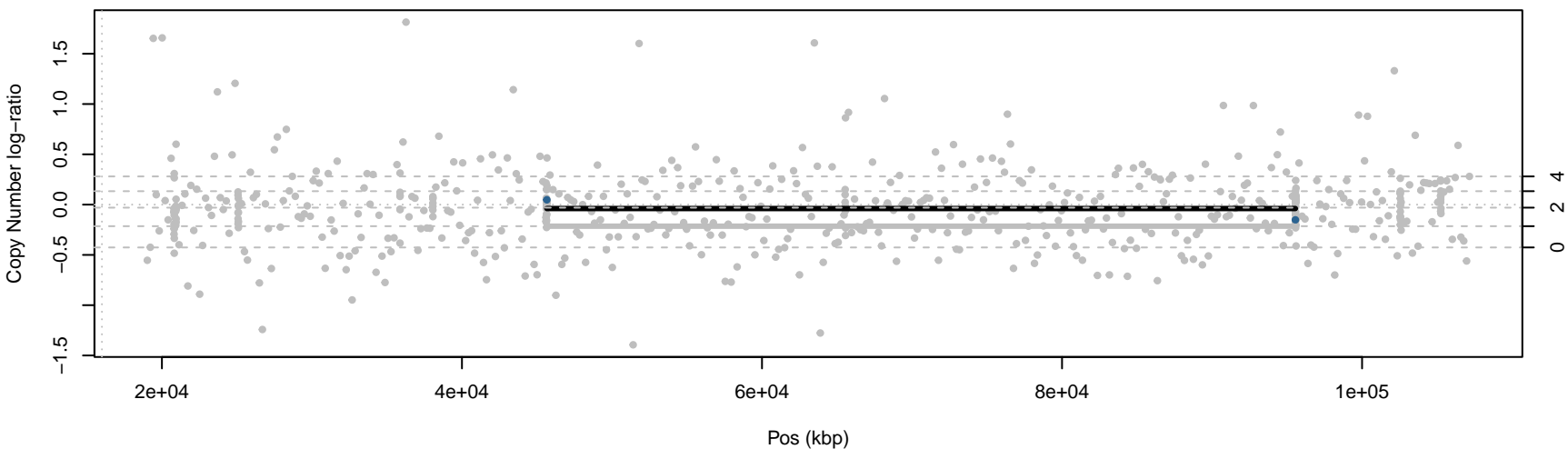
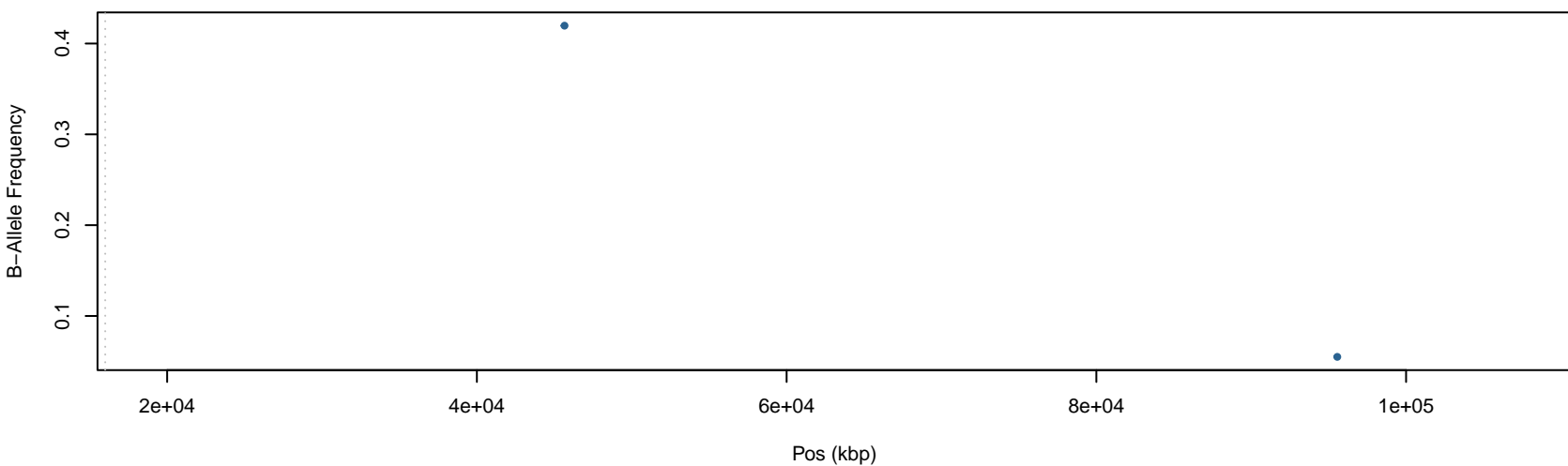
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 12



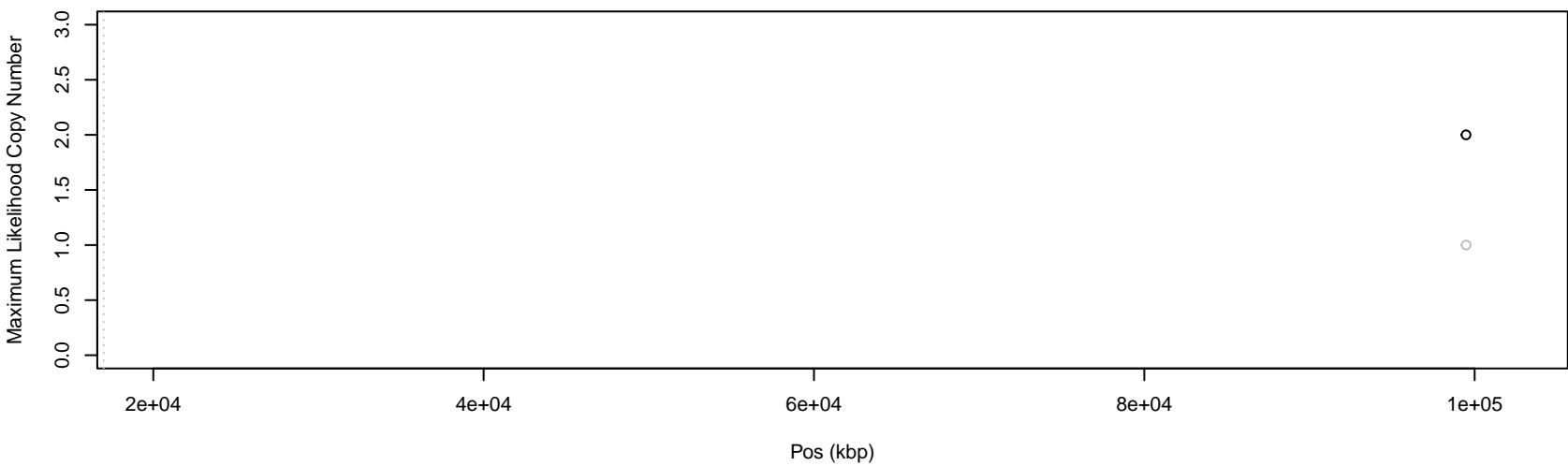
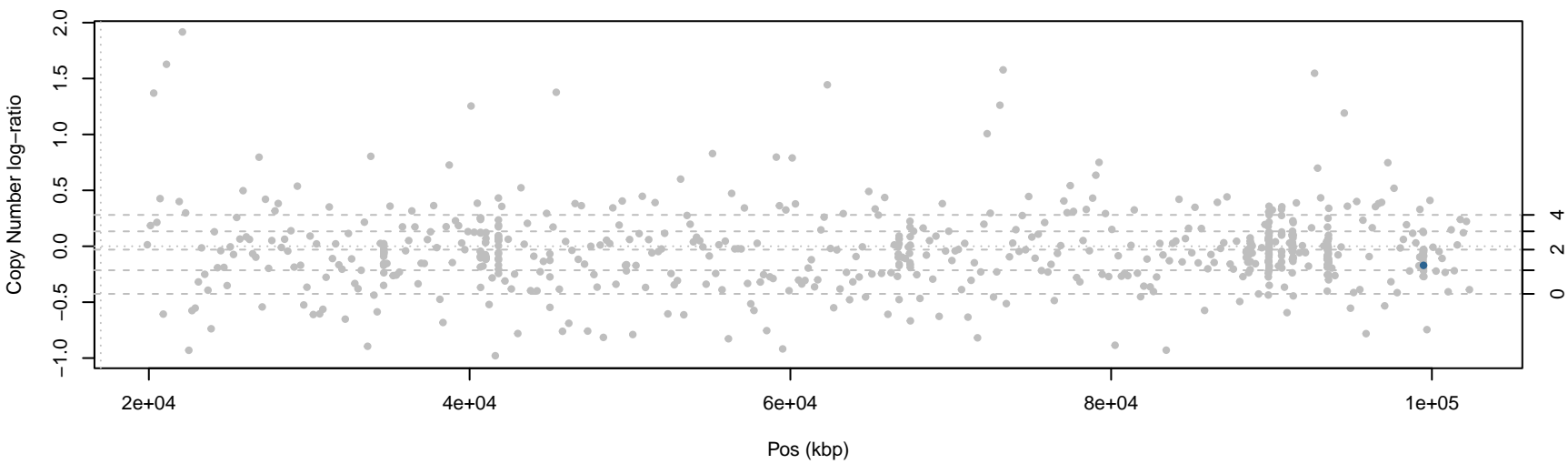
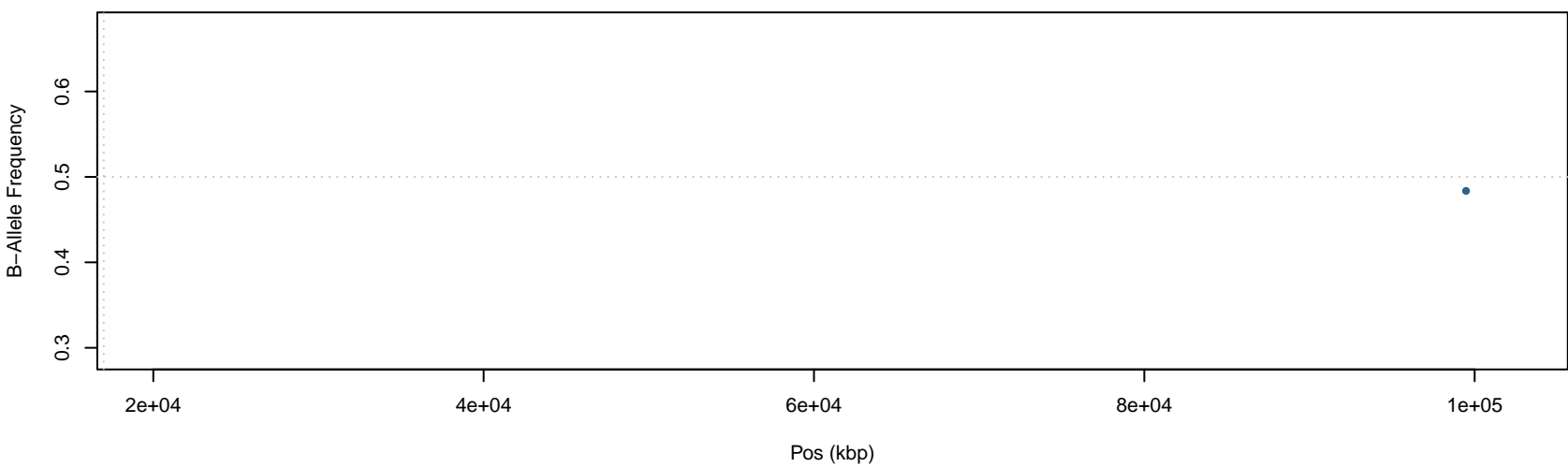
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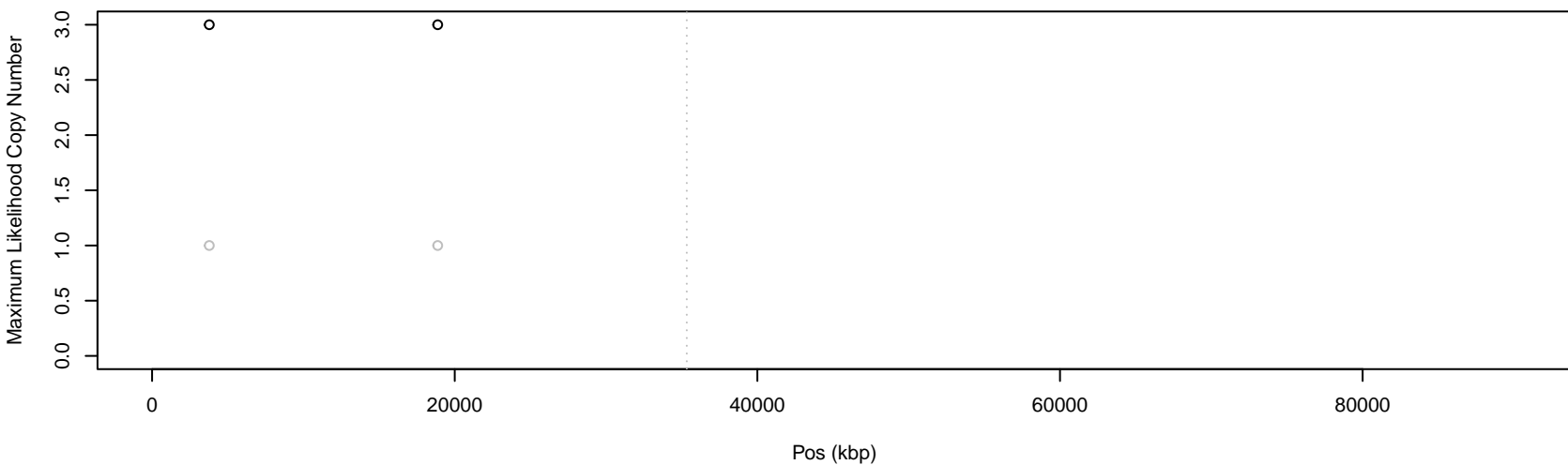
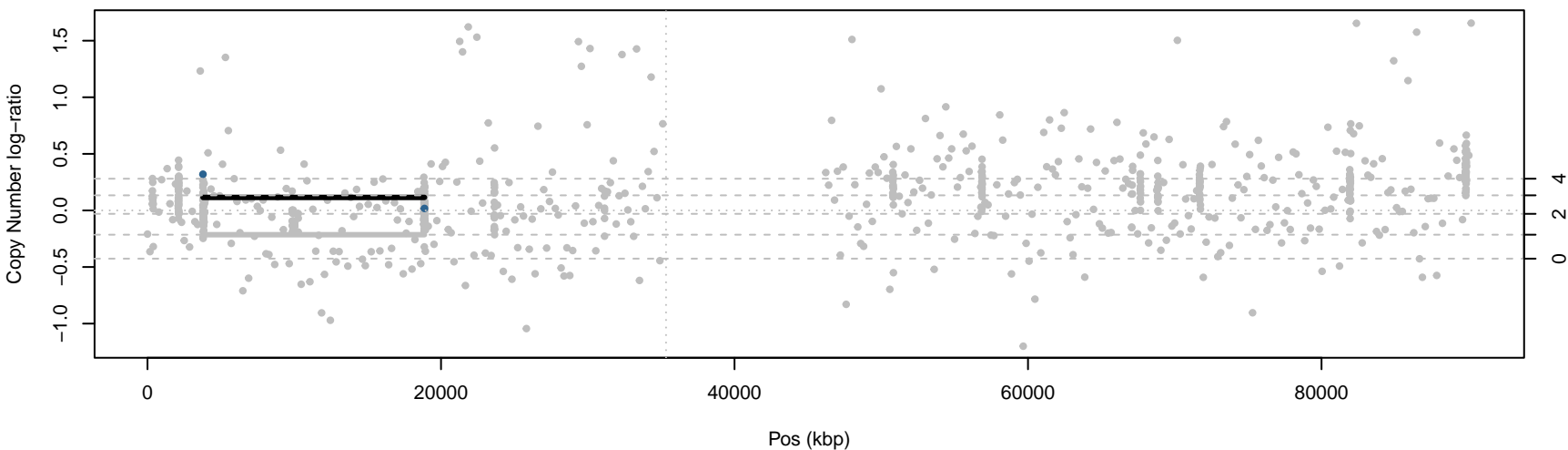
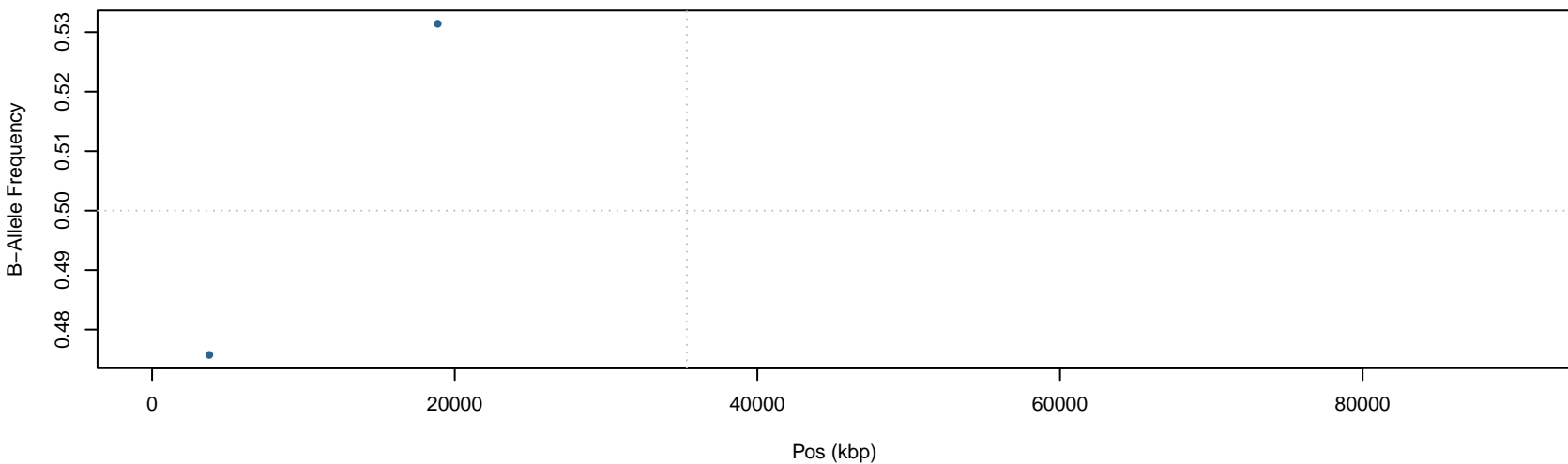
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 14



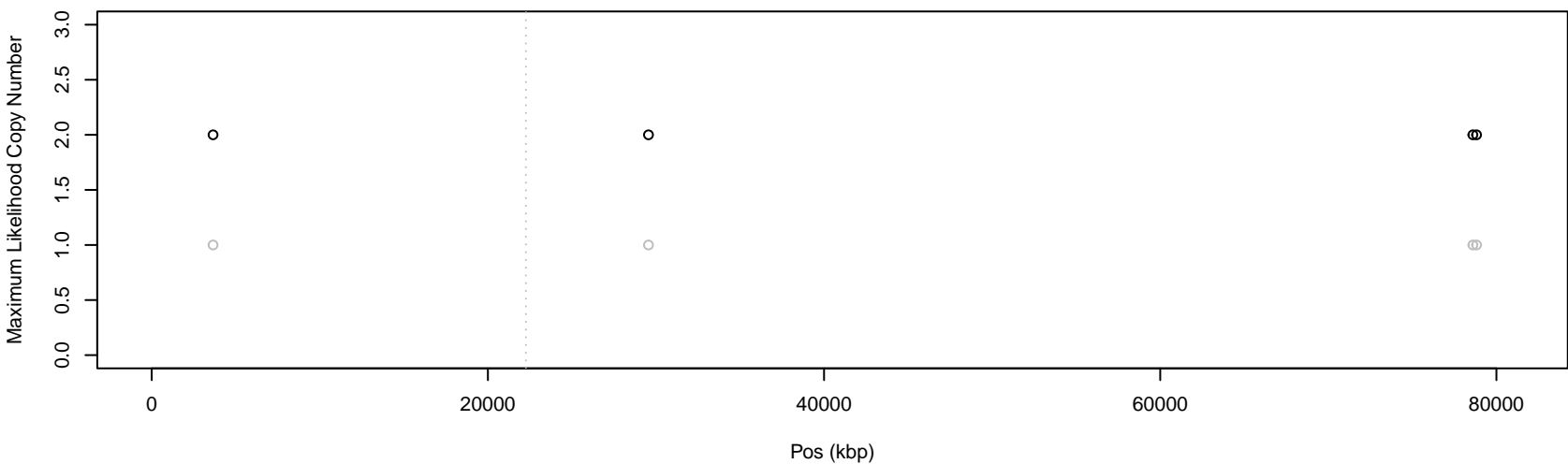
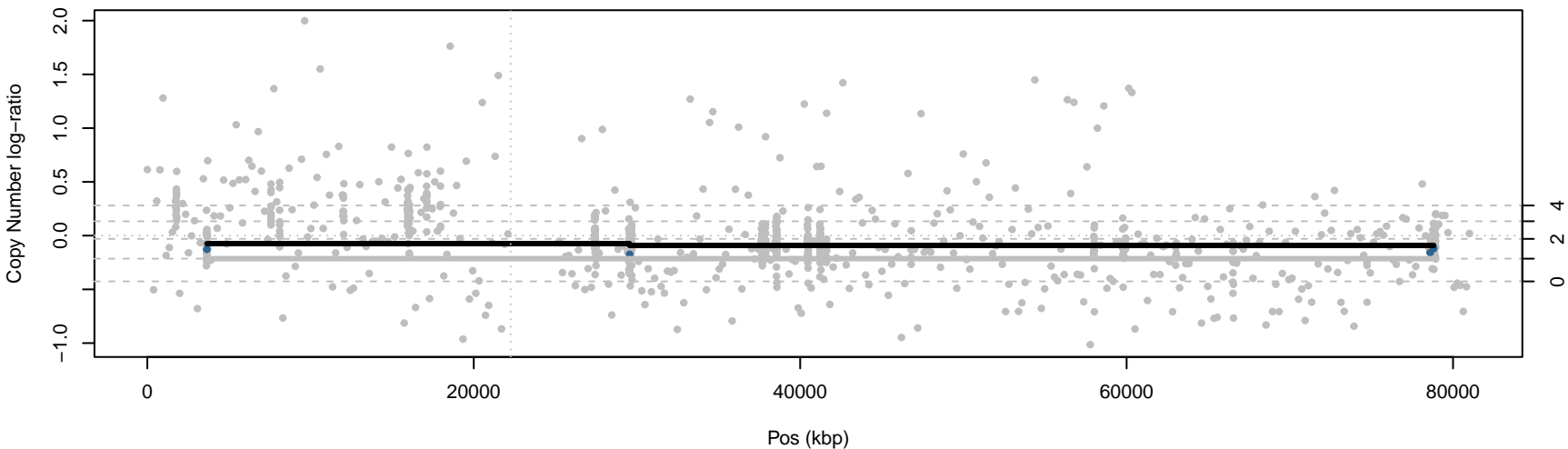
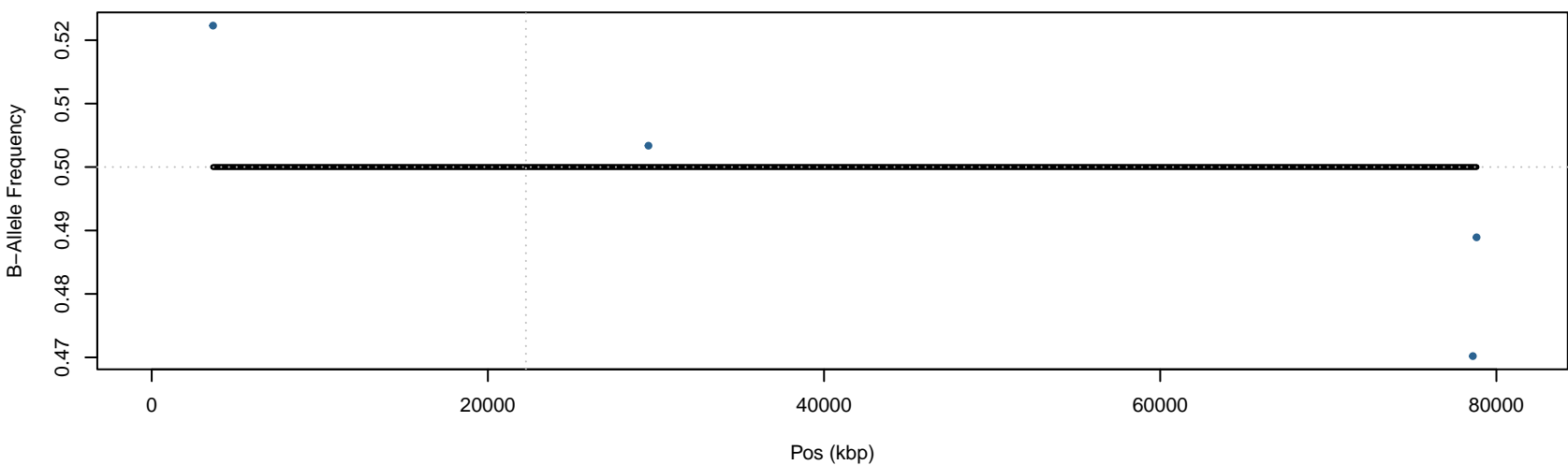
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 15



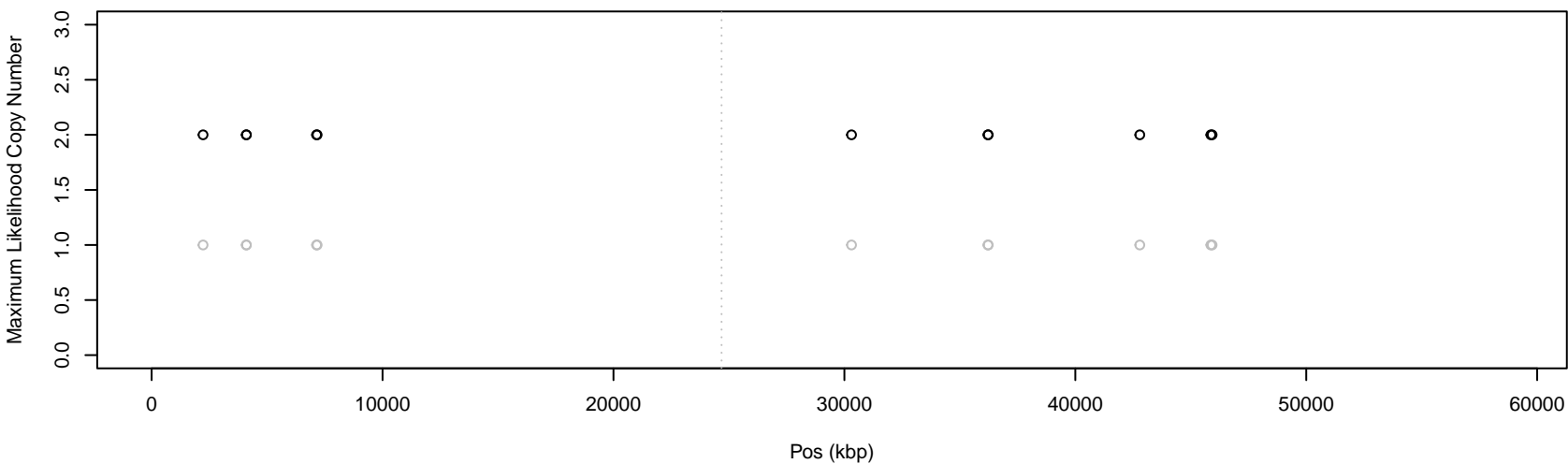
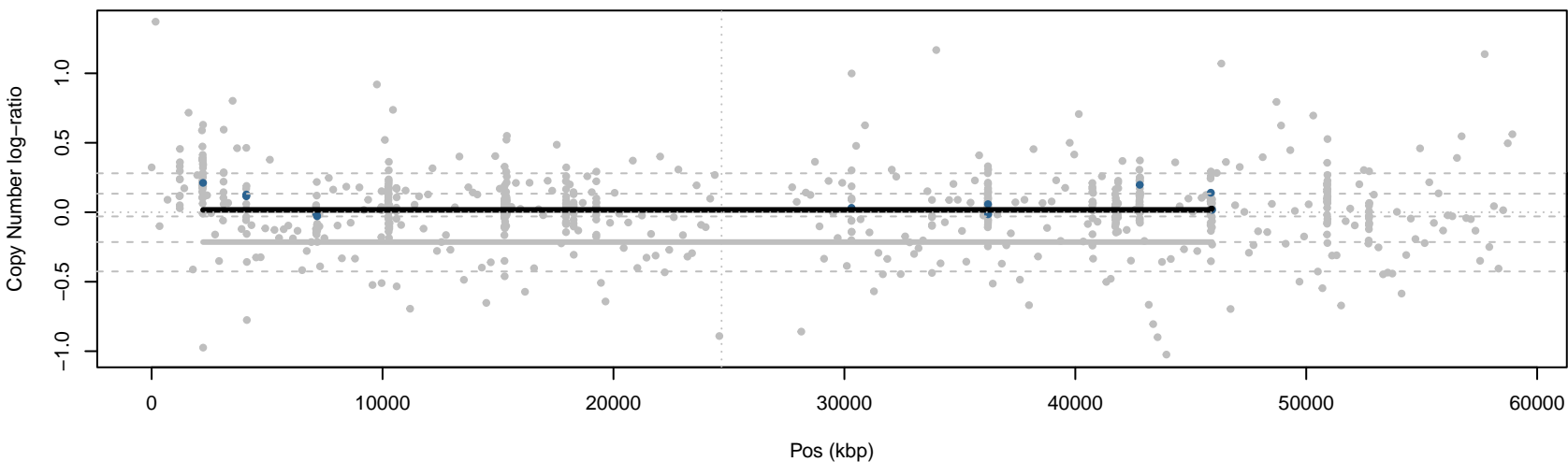
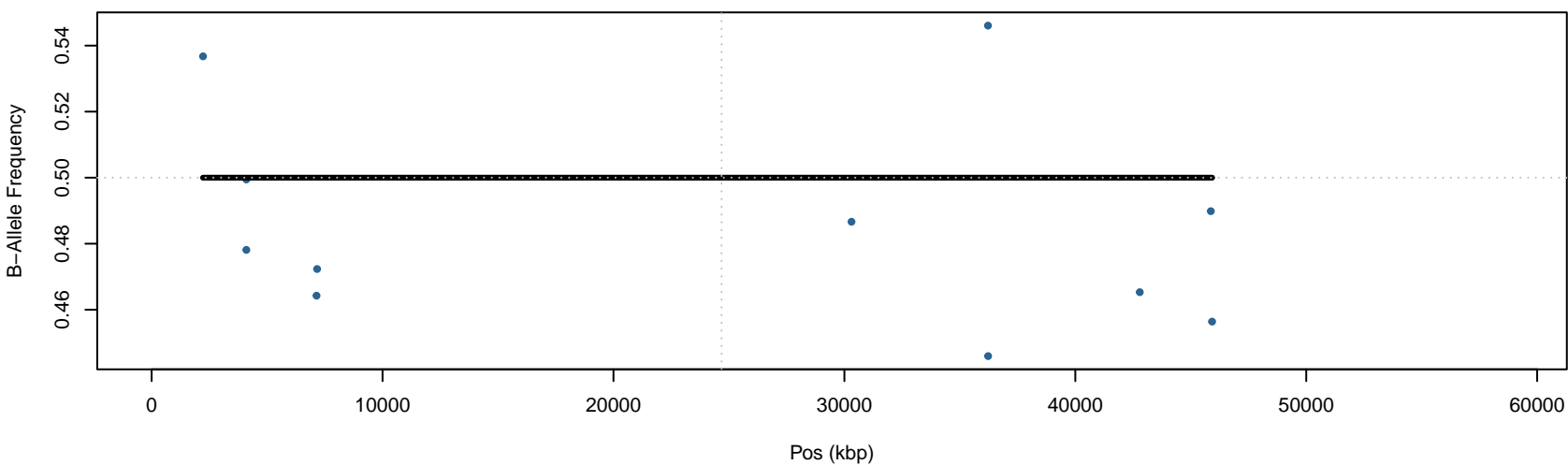
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 16



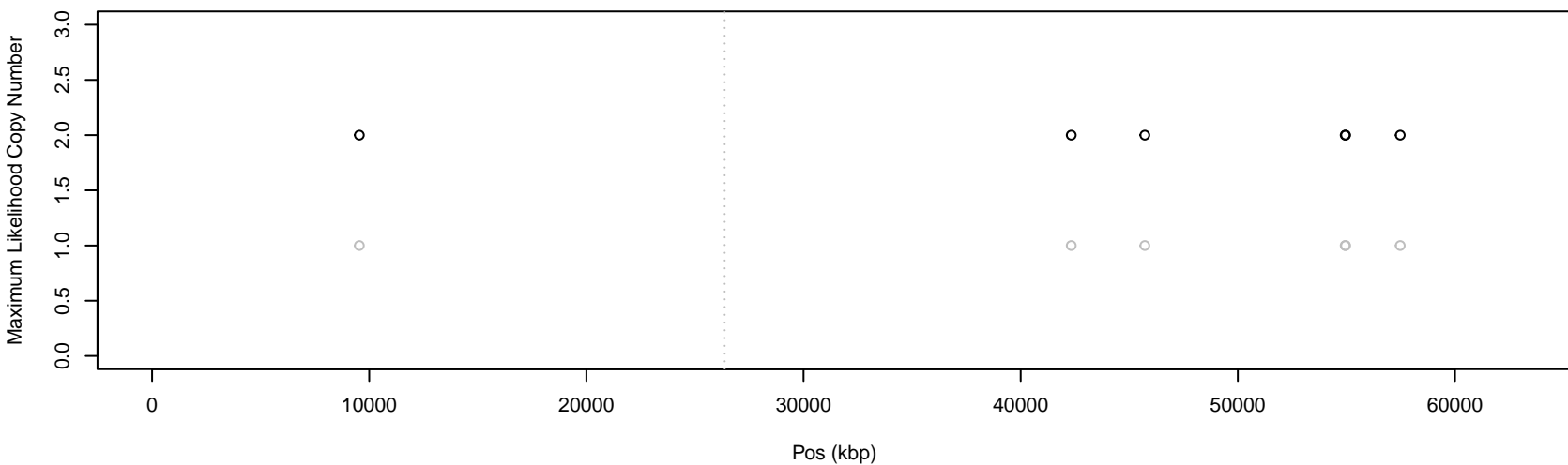
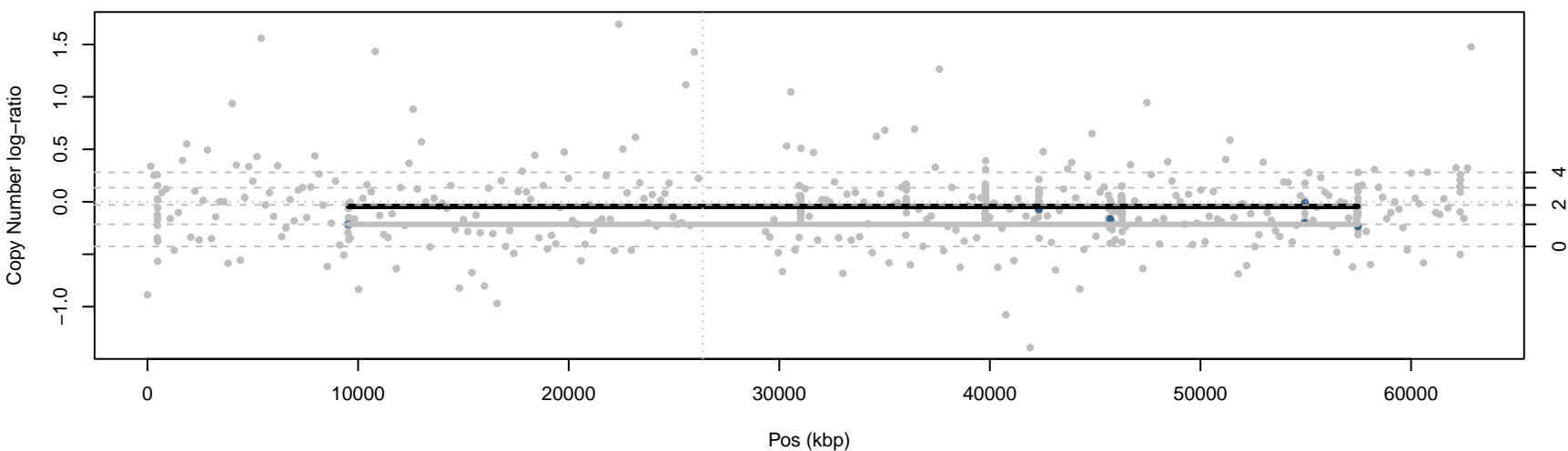
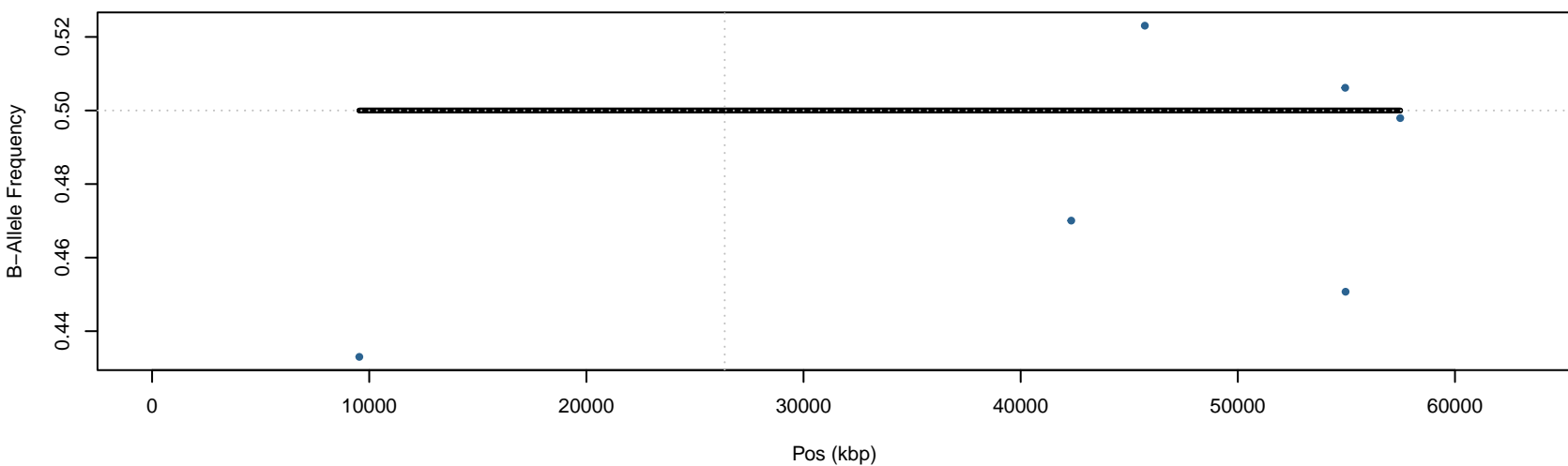
Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 17



Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 19



Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 20



Purity: 0.24 Tumor ploidy: 2.175 SNV log-likelihood: 80.83 GoF: 95.9% Mean coverage: 565;782 Chromosome: 22

