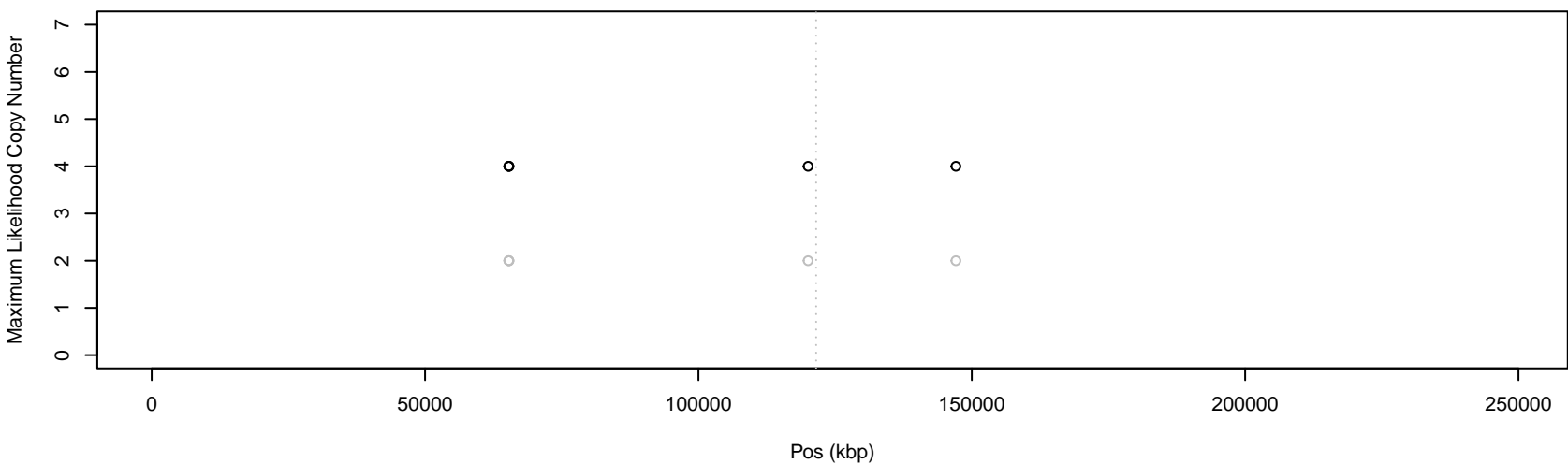
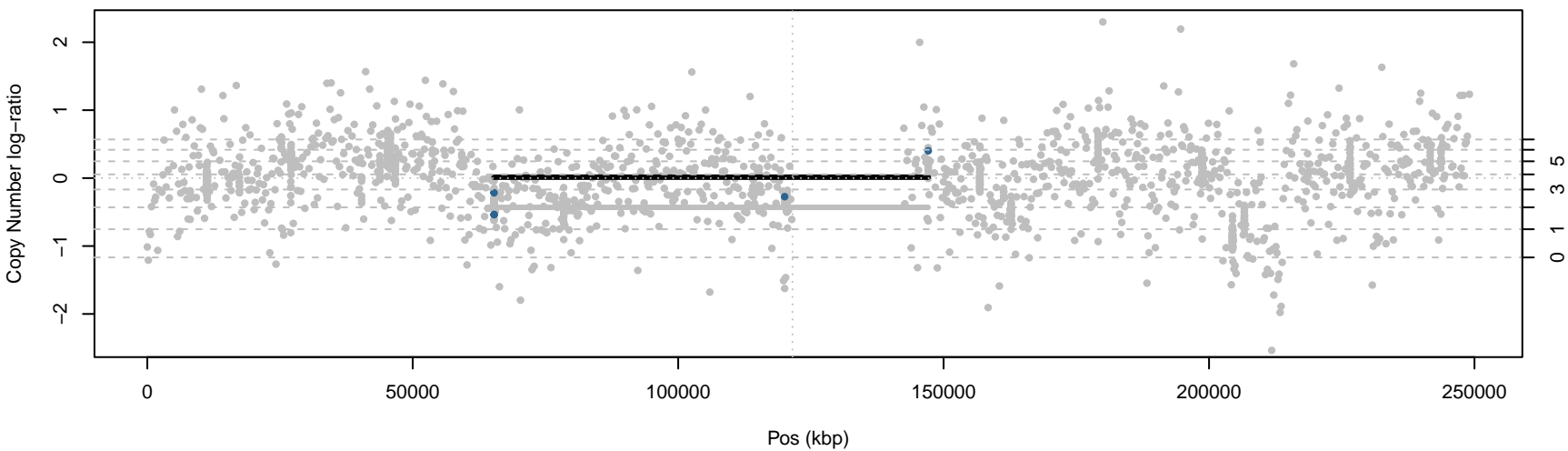
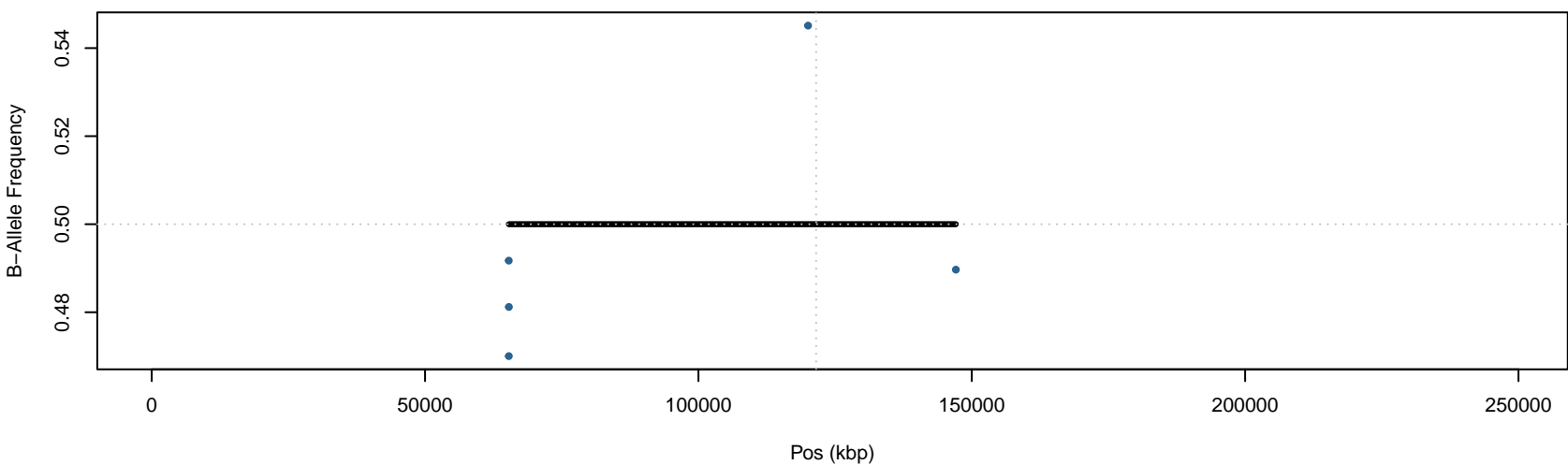
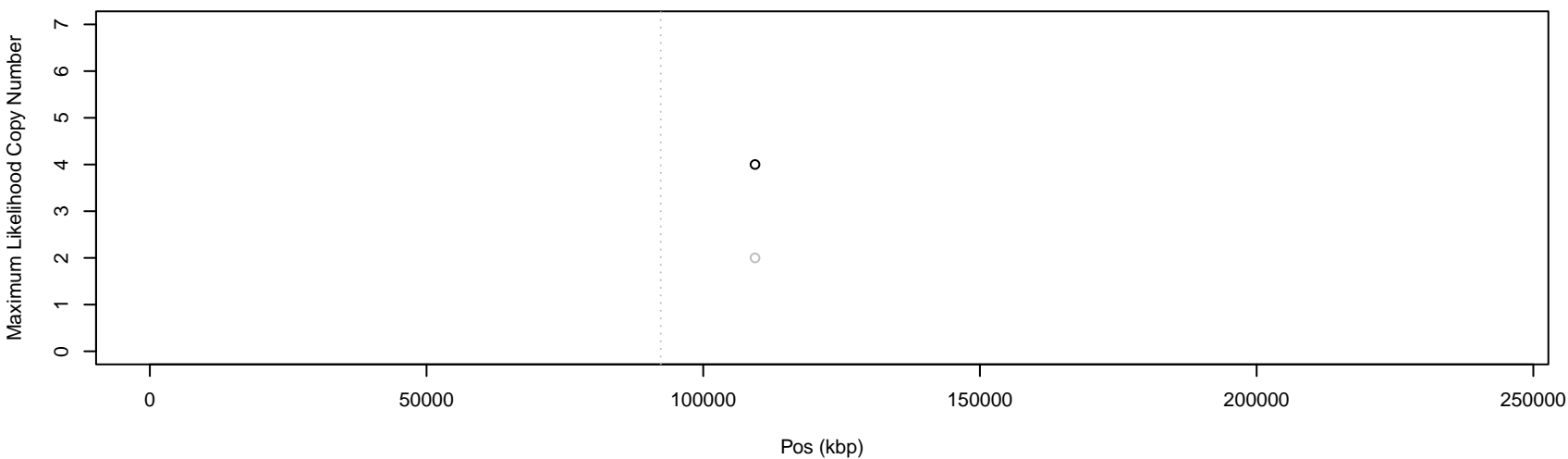
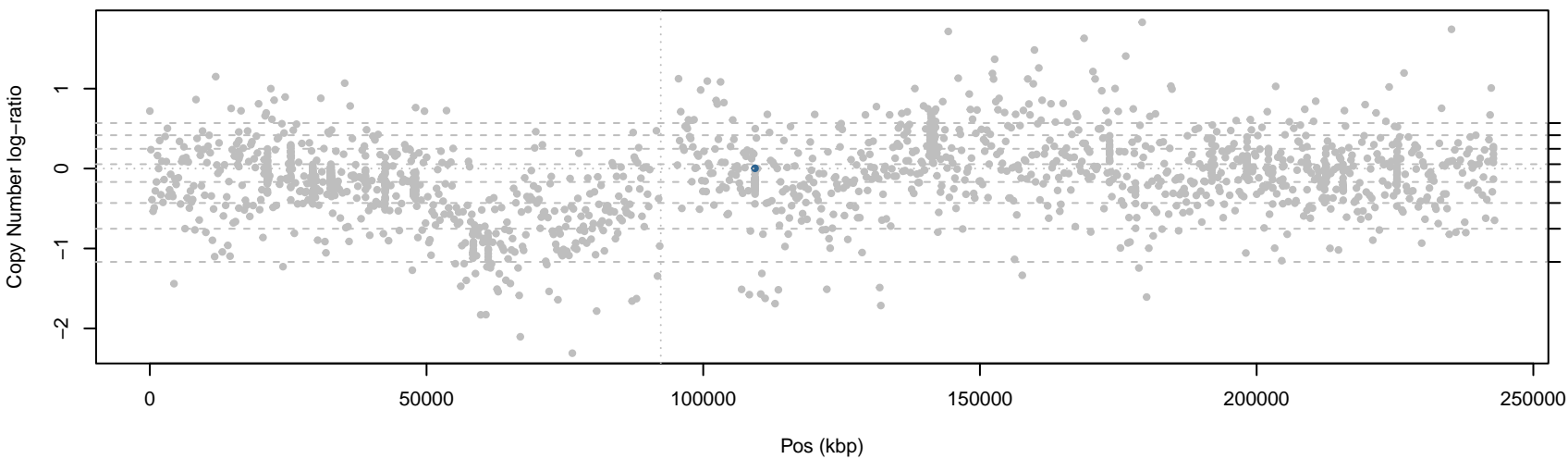
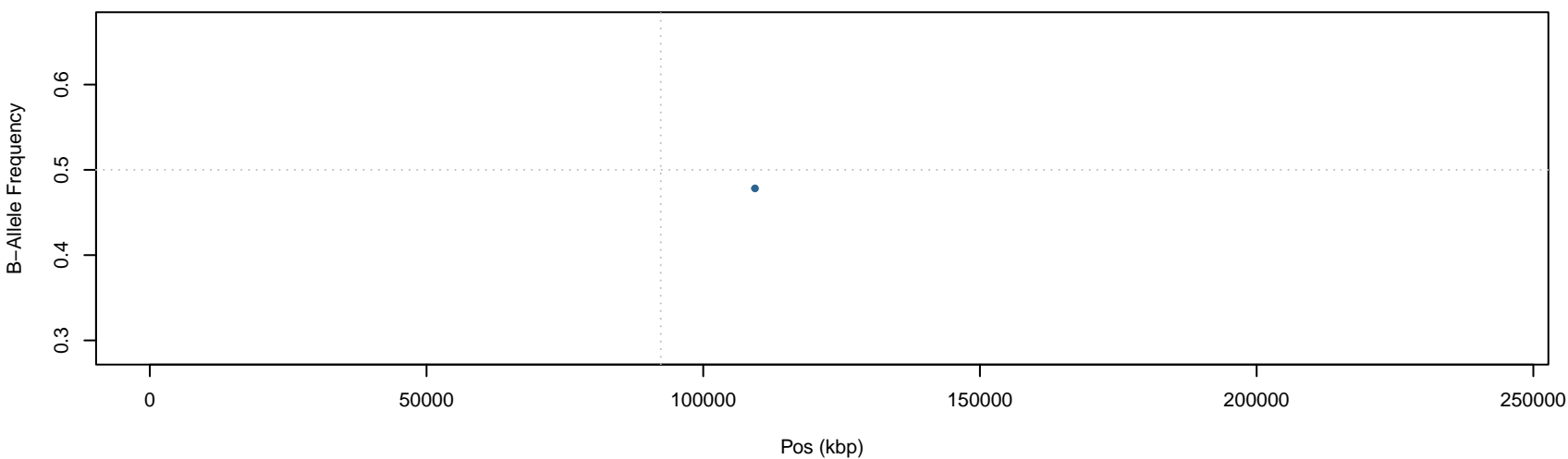


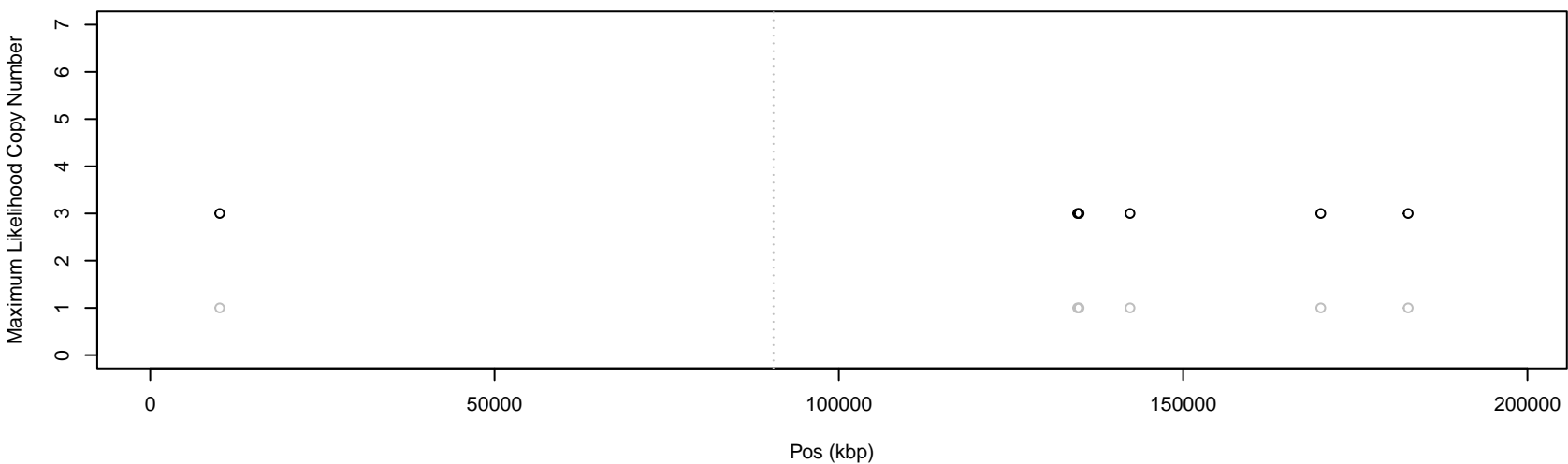
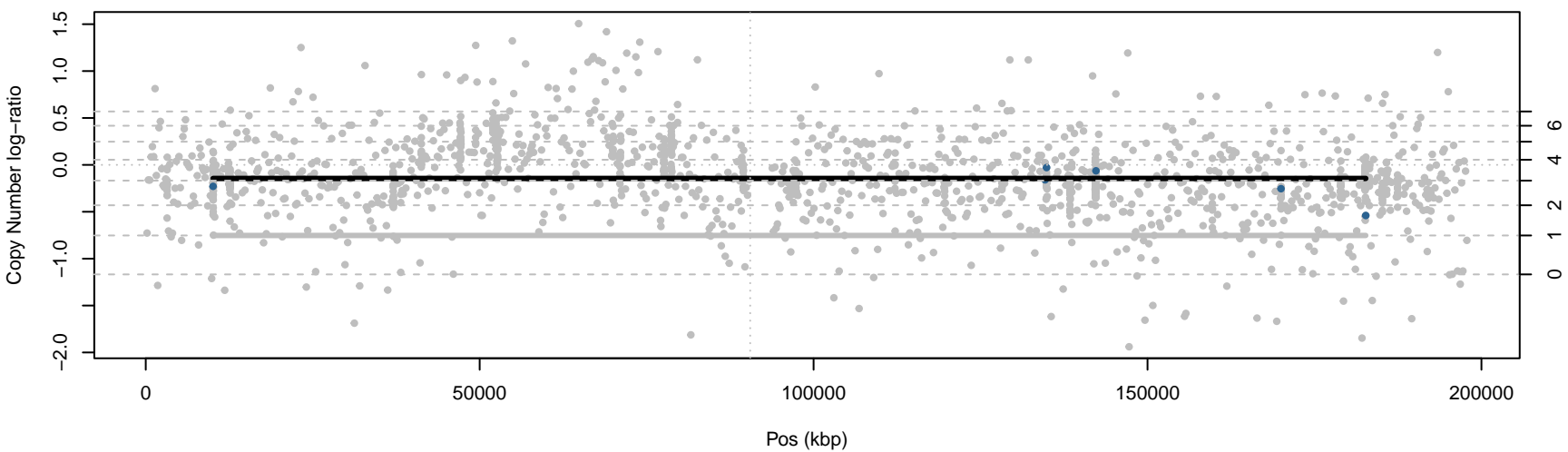
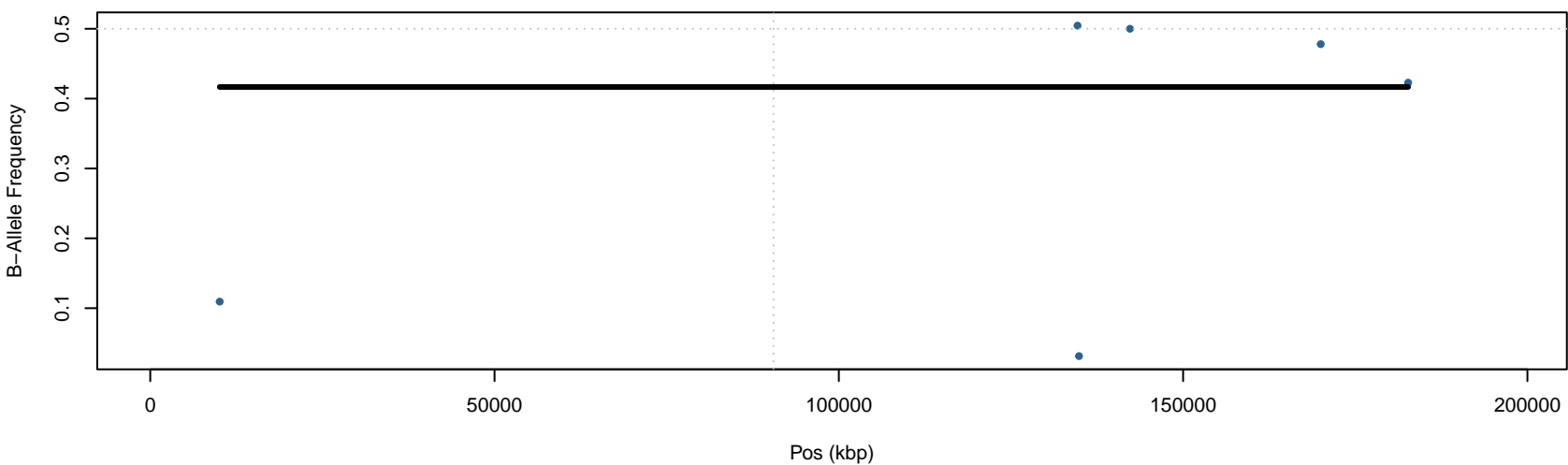
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 1



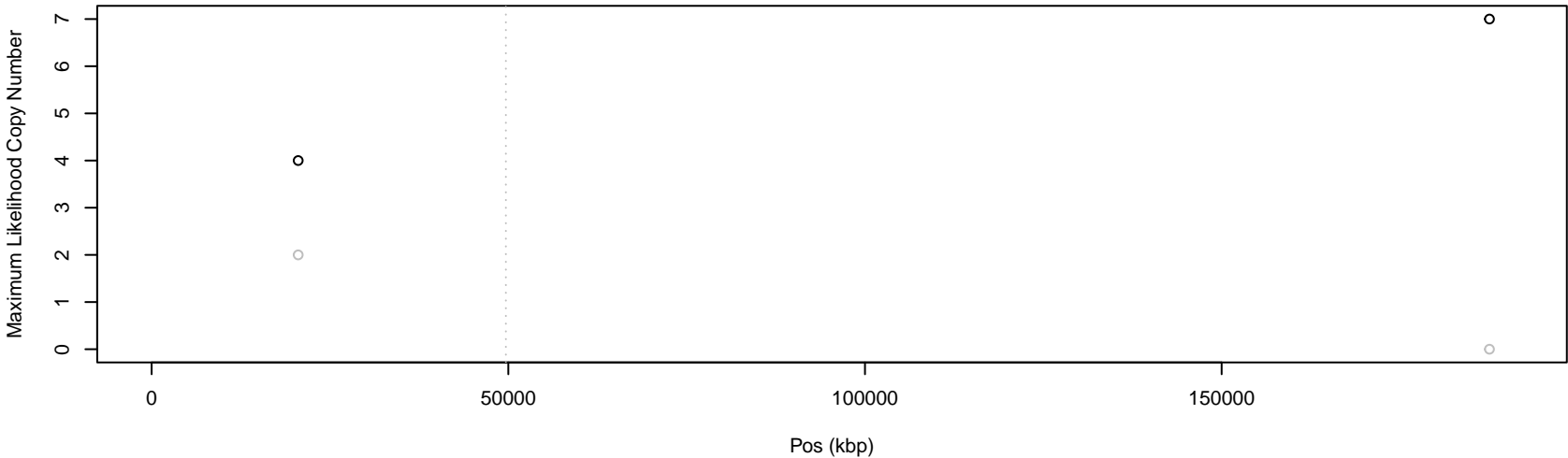
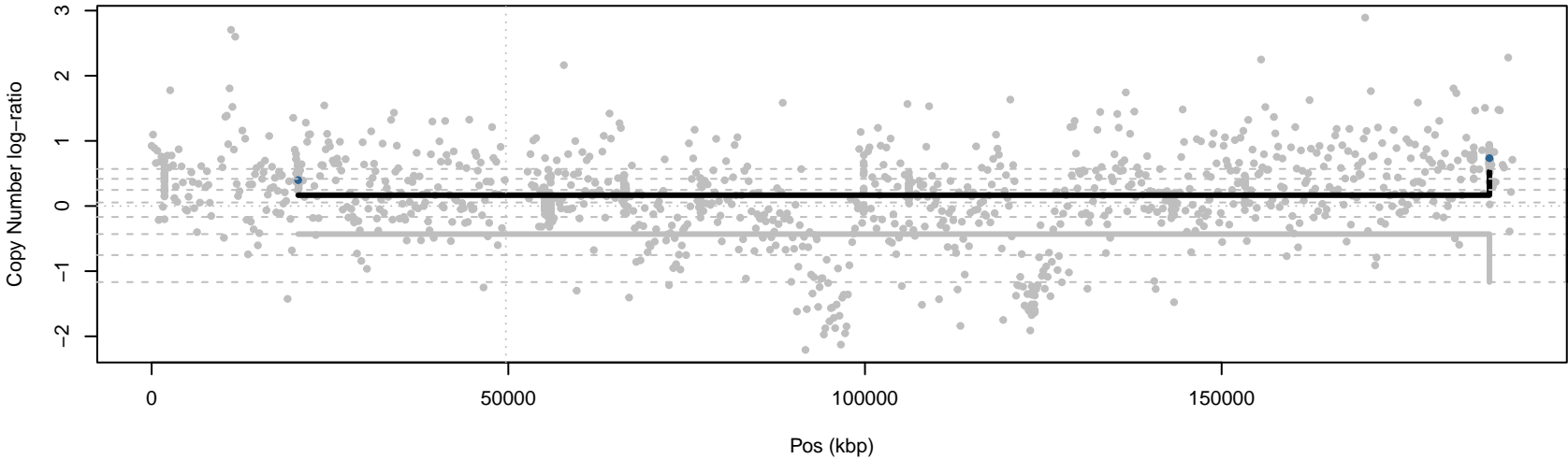
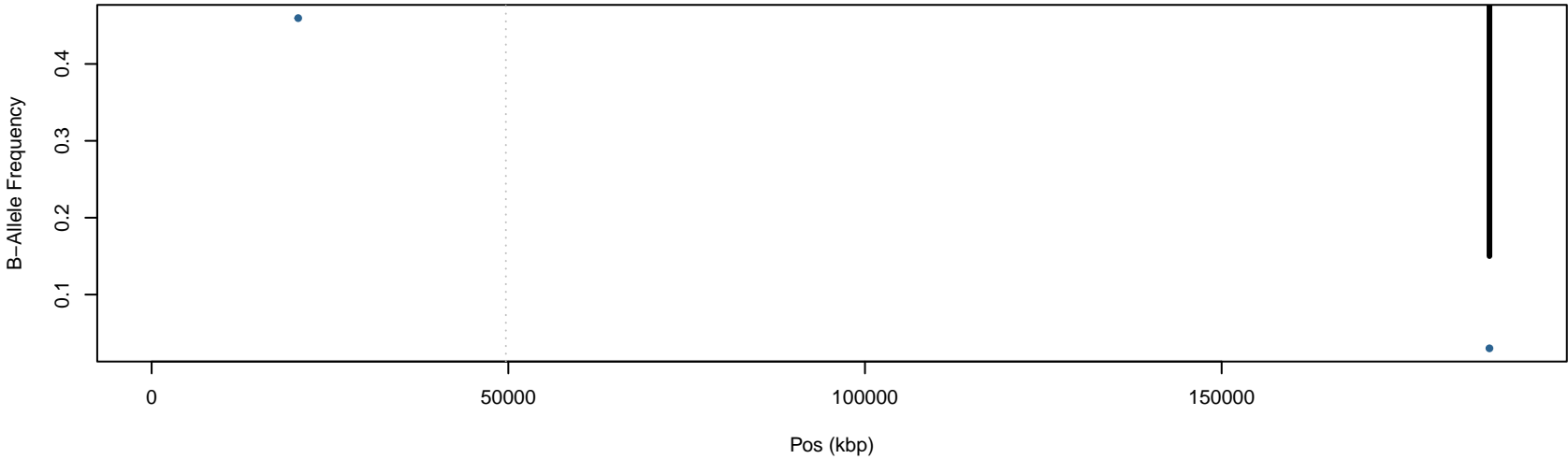
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 2



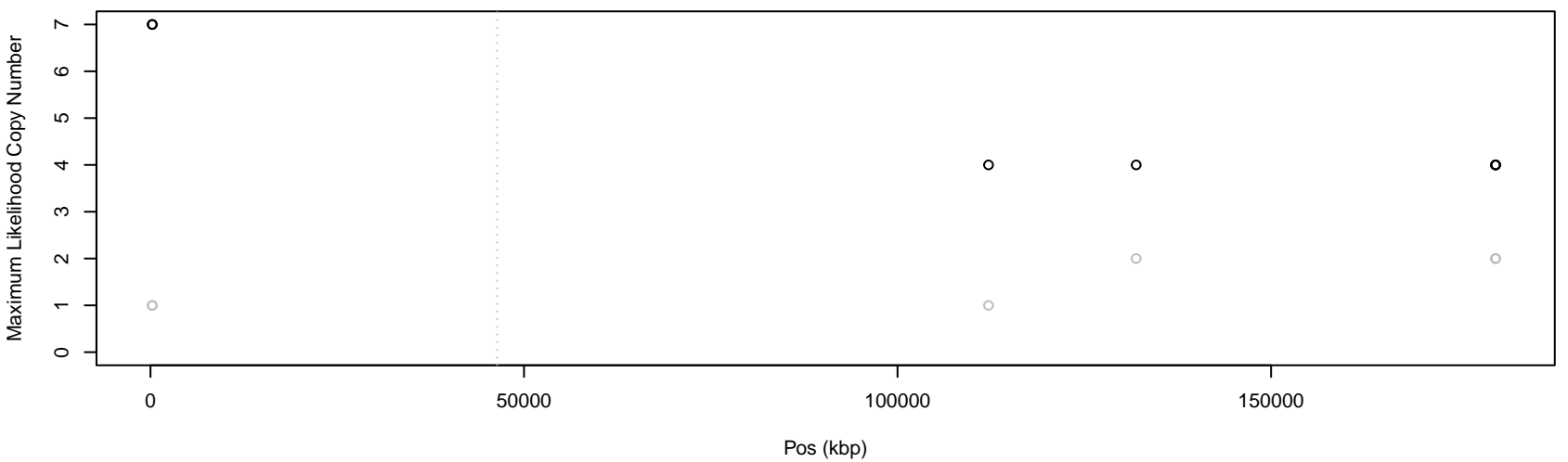
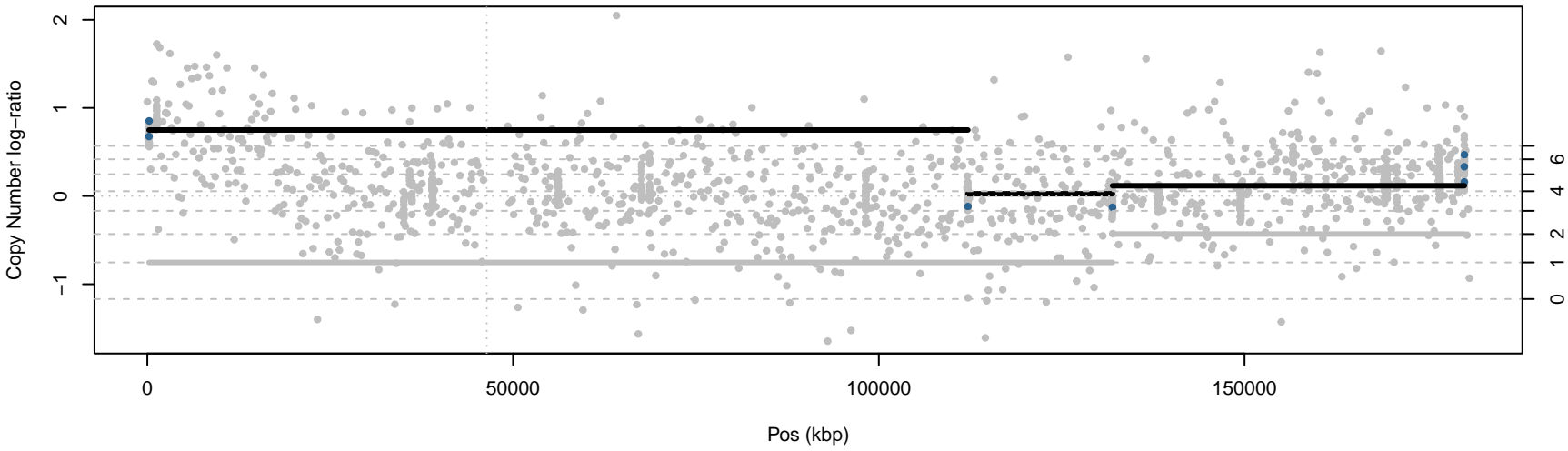
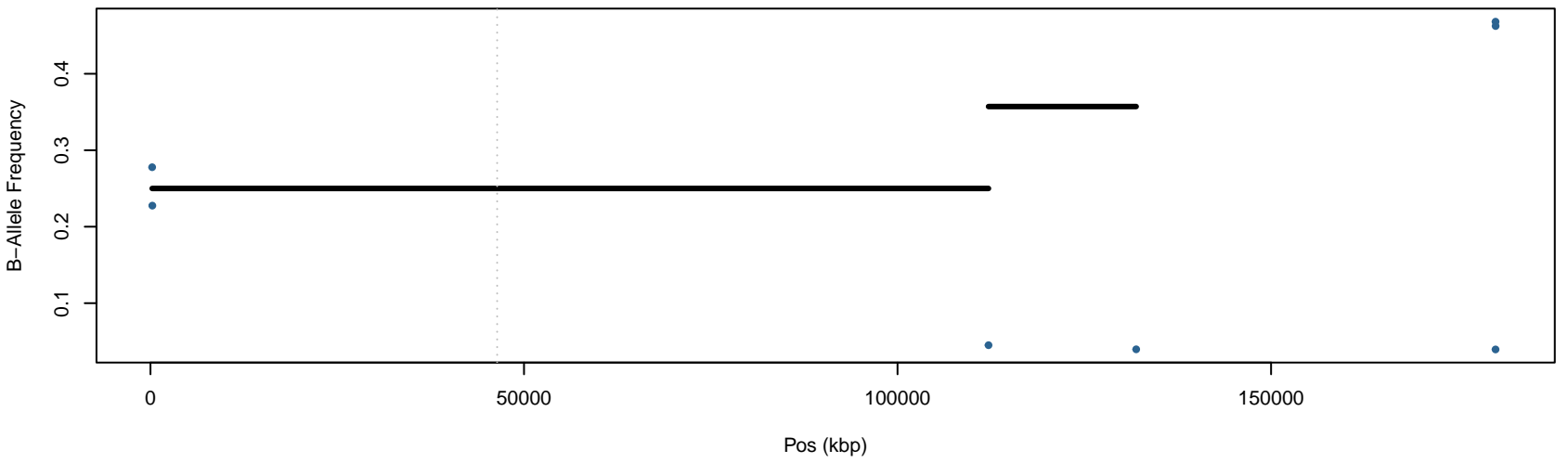
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 3



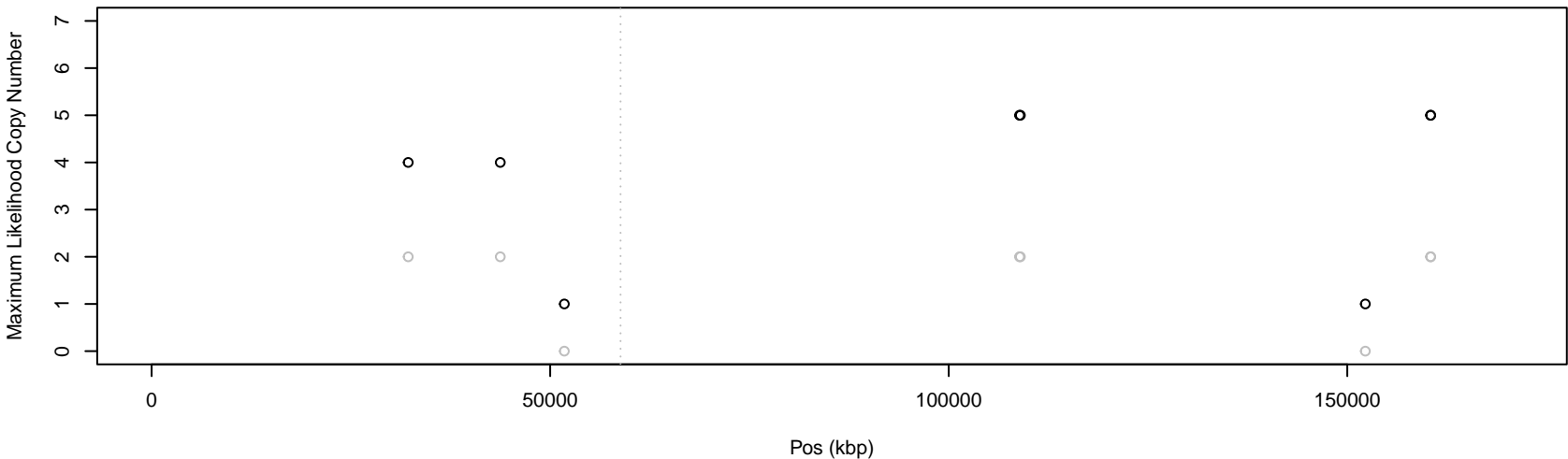
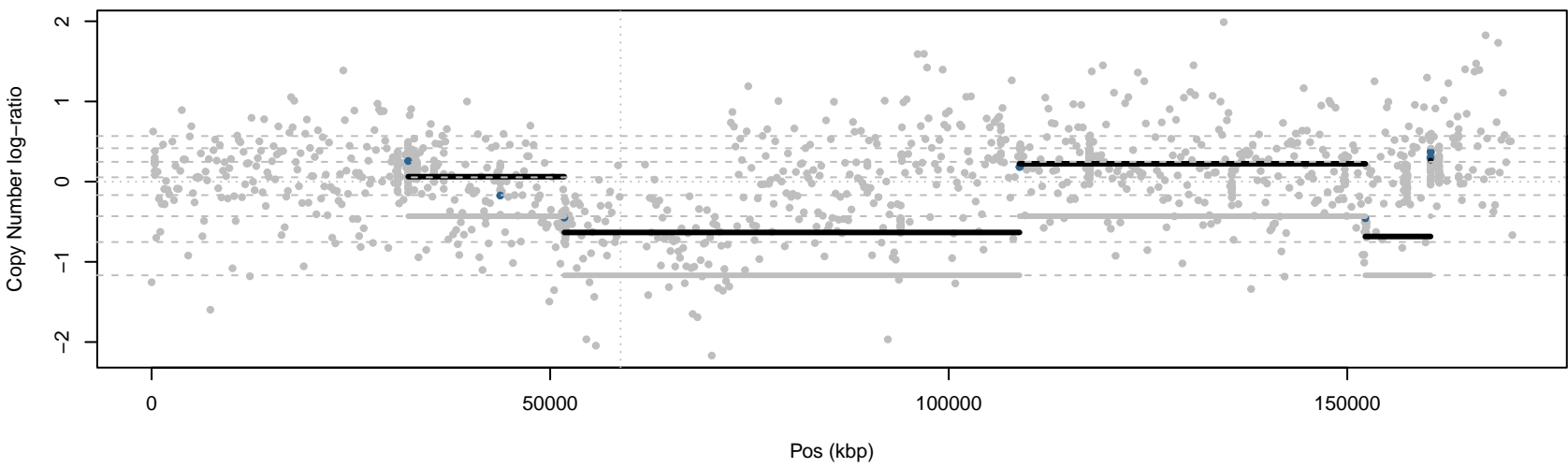
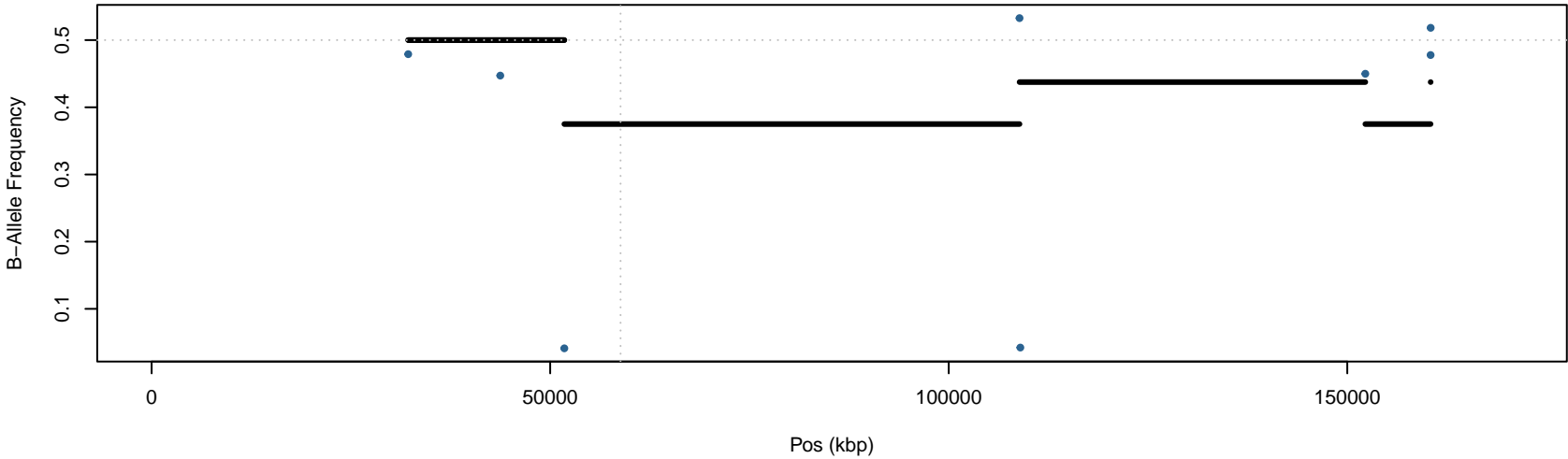
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 4



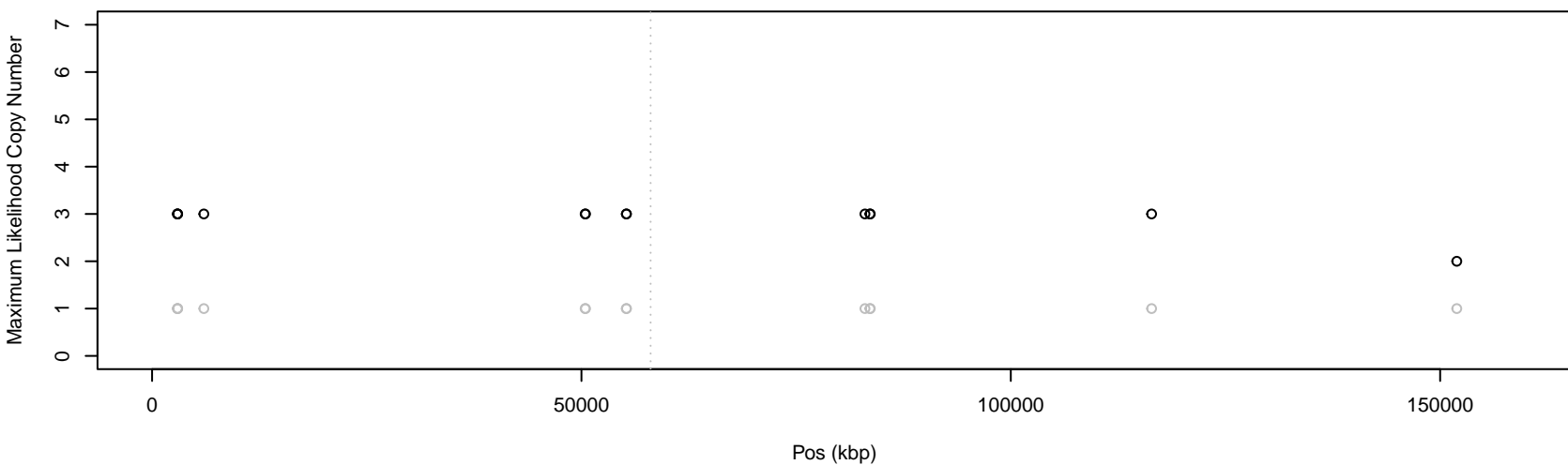
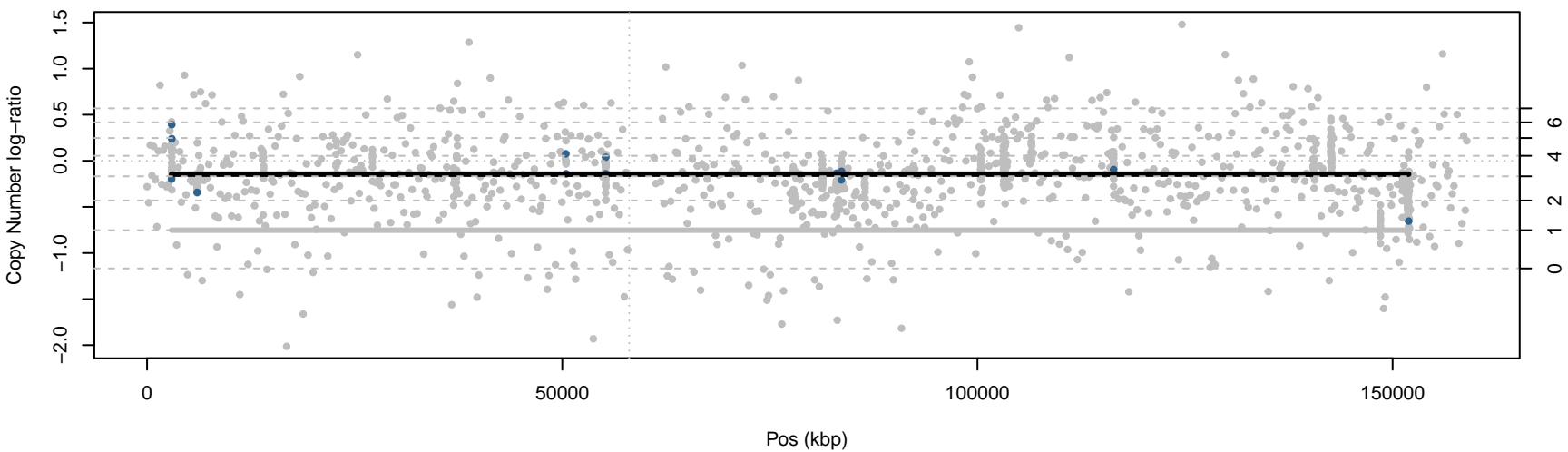
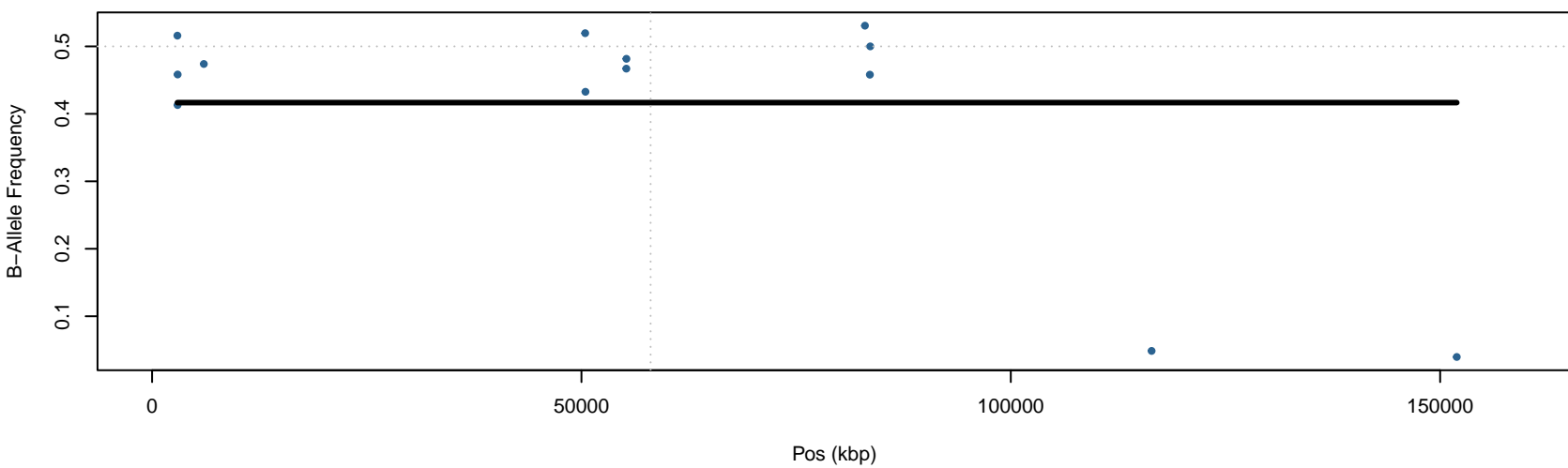
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 5

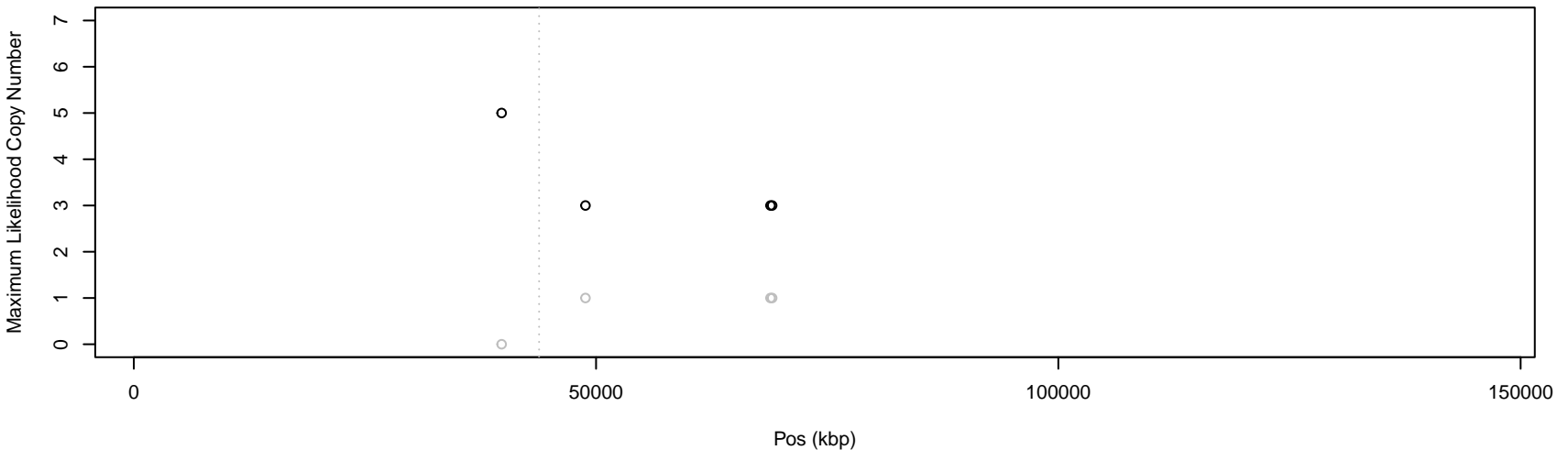
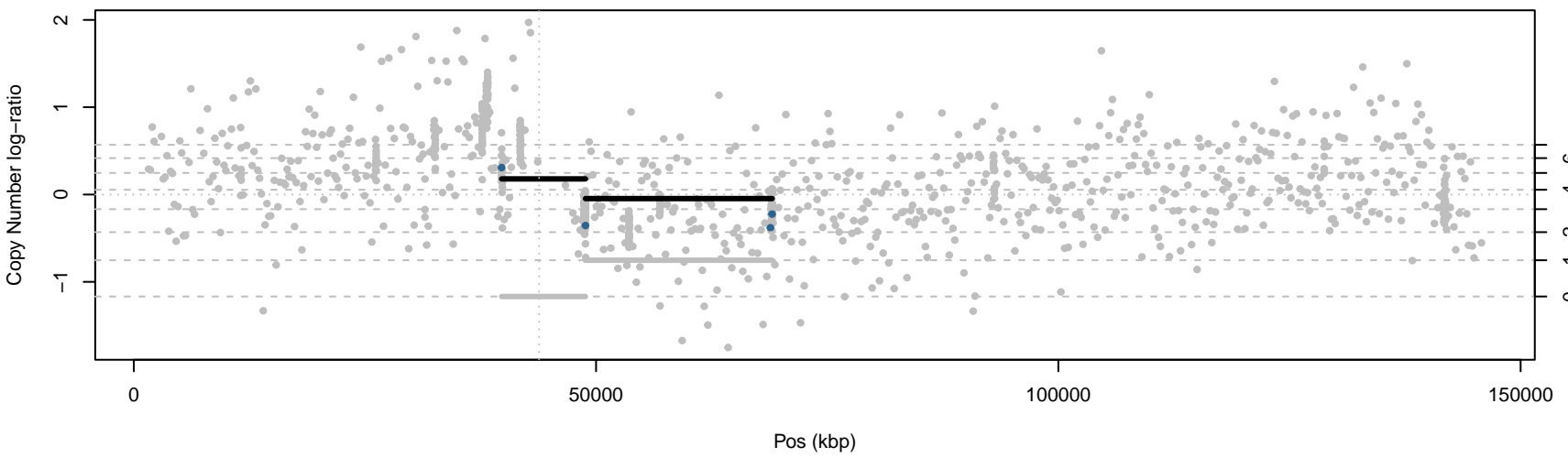
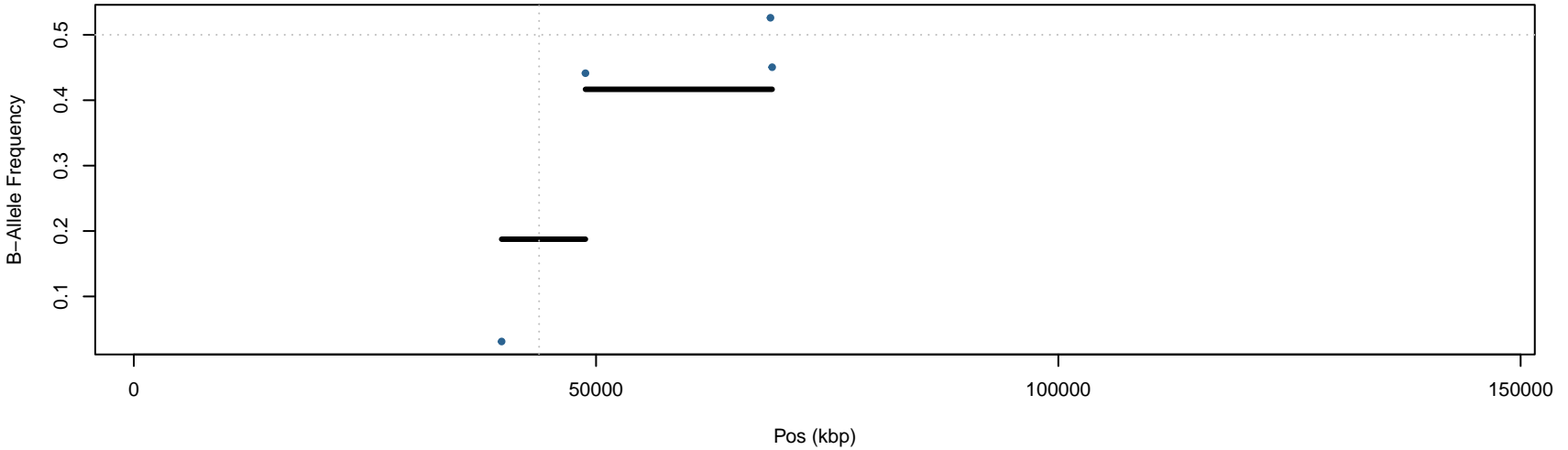


Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 6



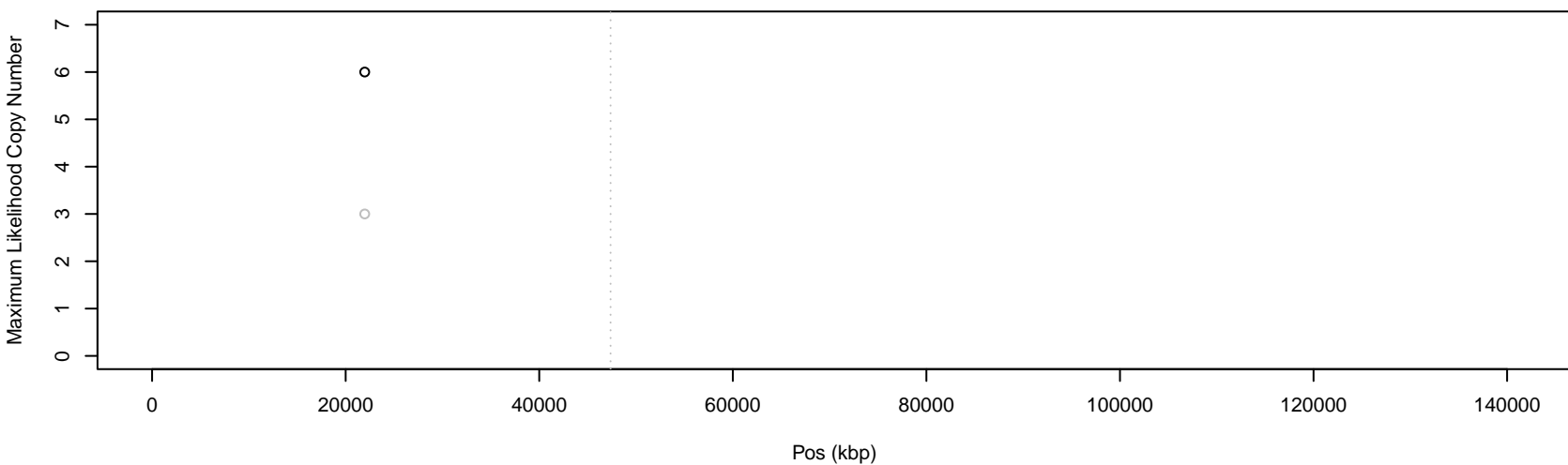
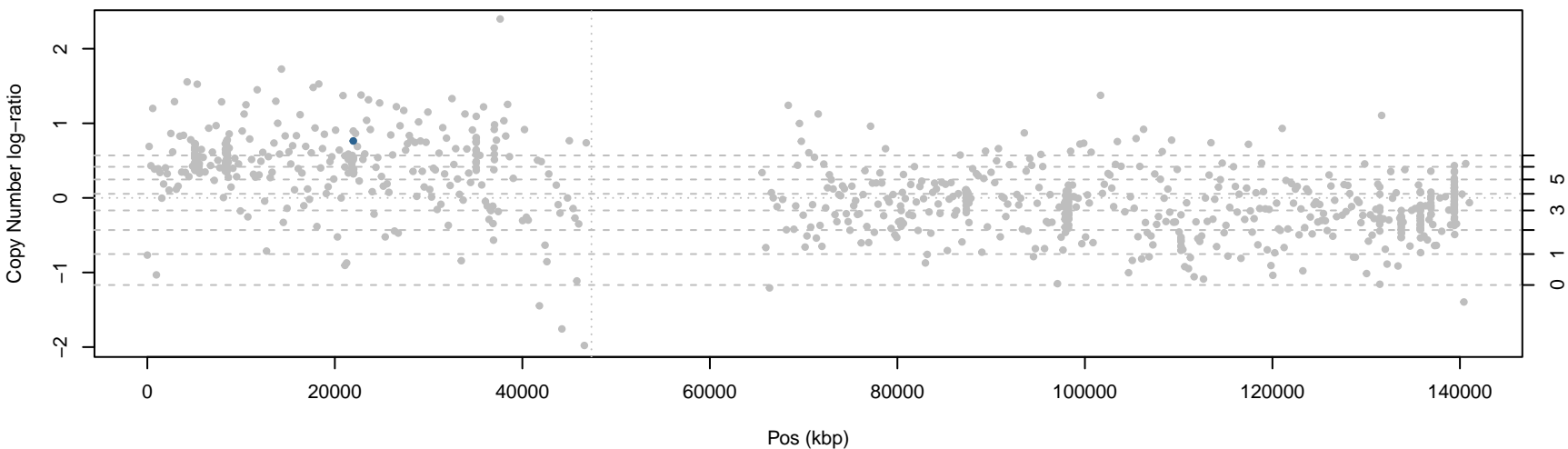
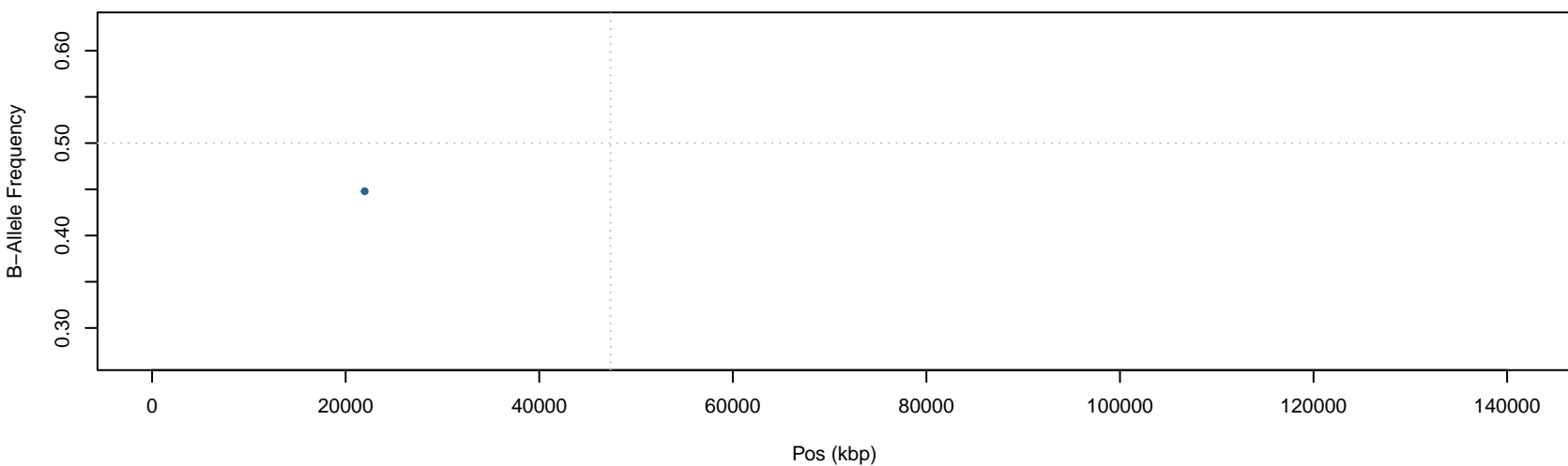
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 7



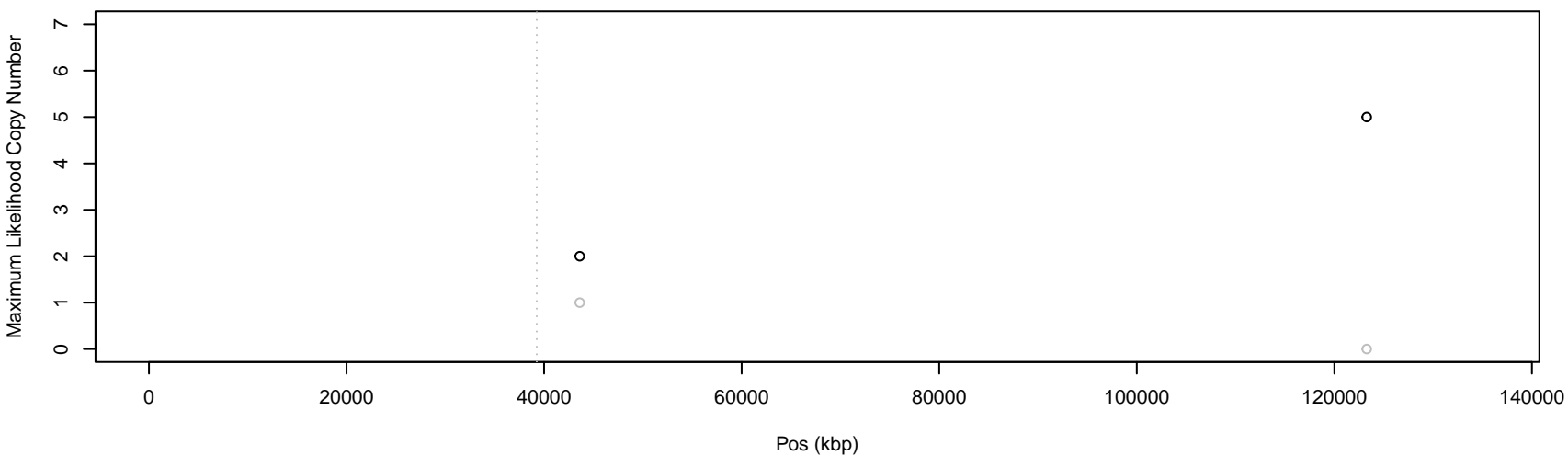
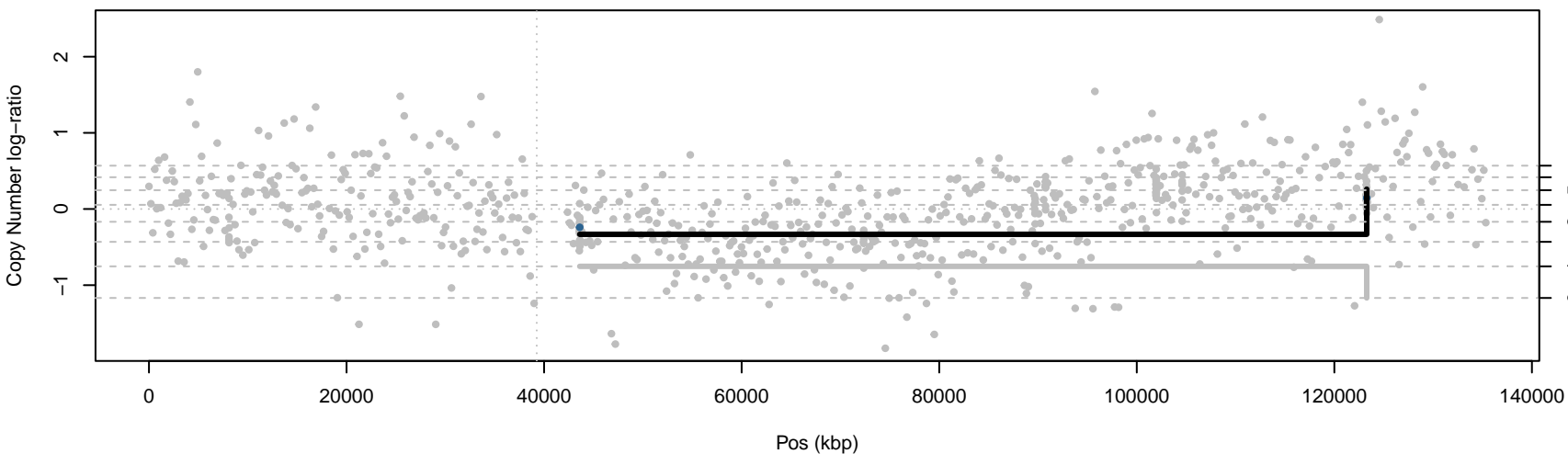
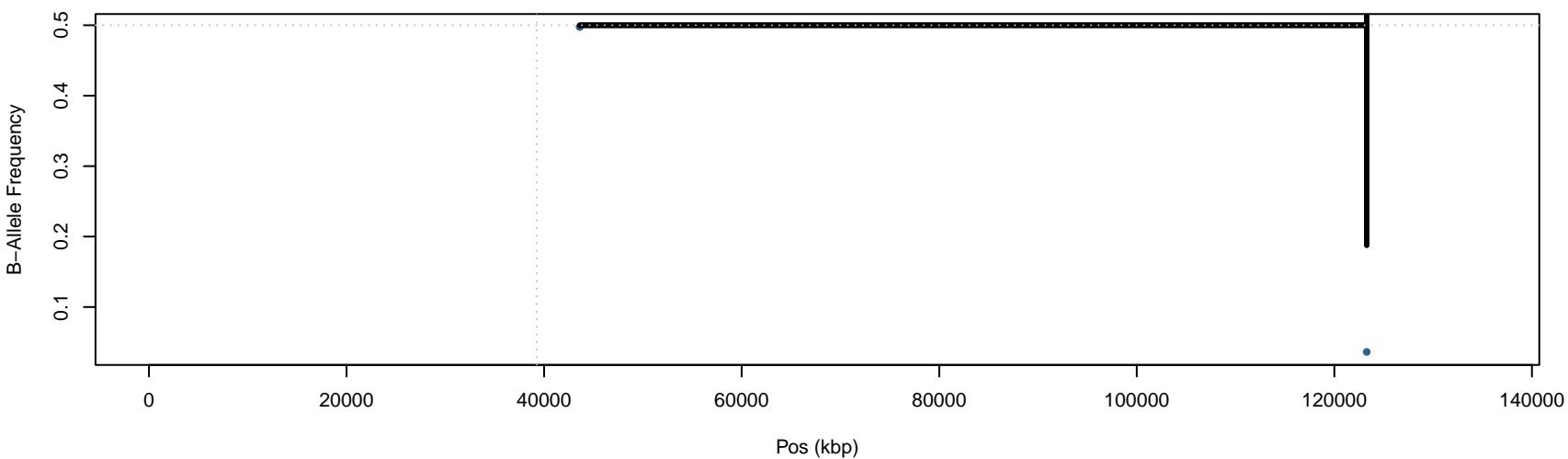




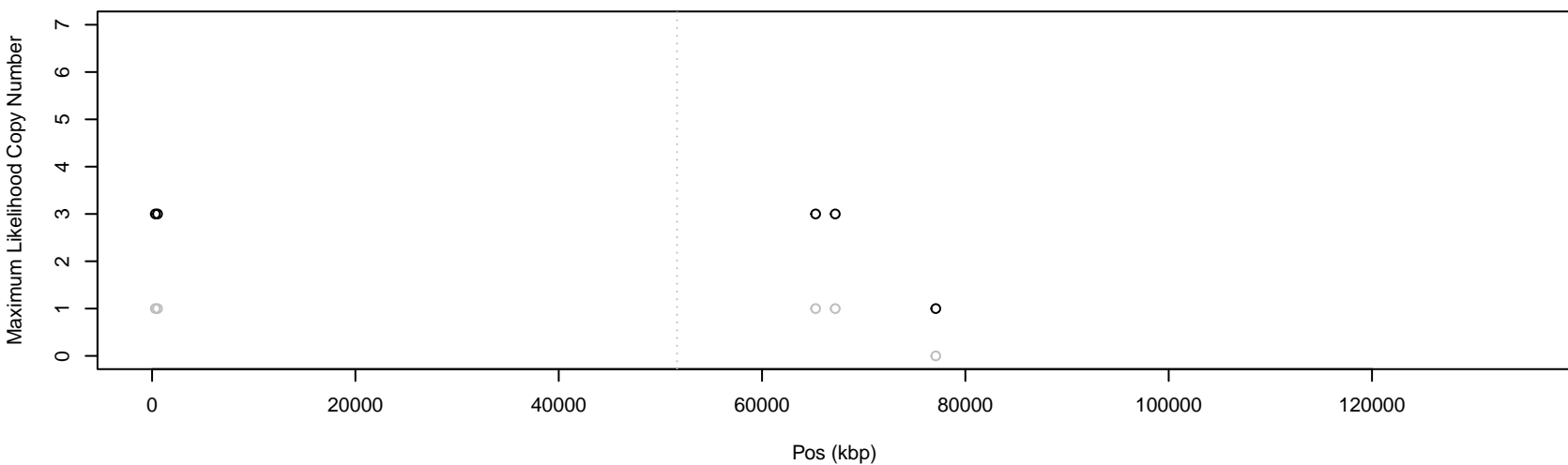
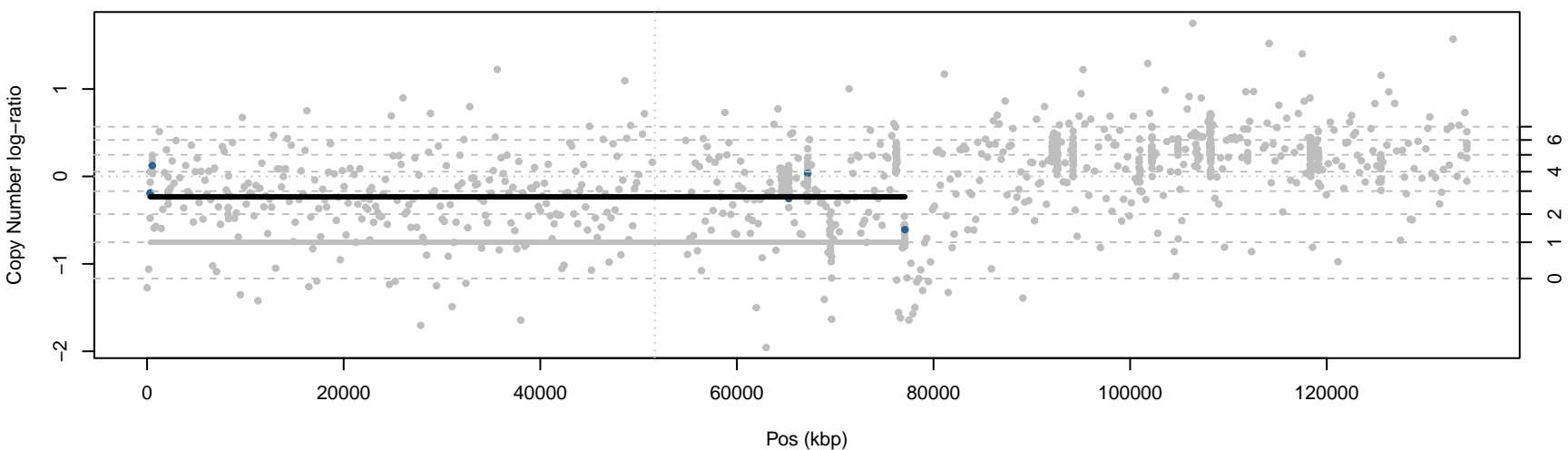
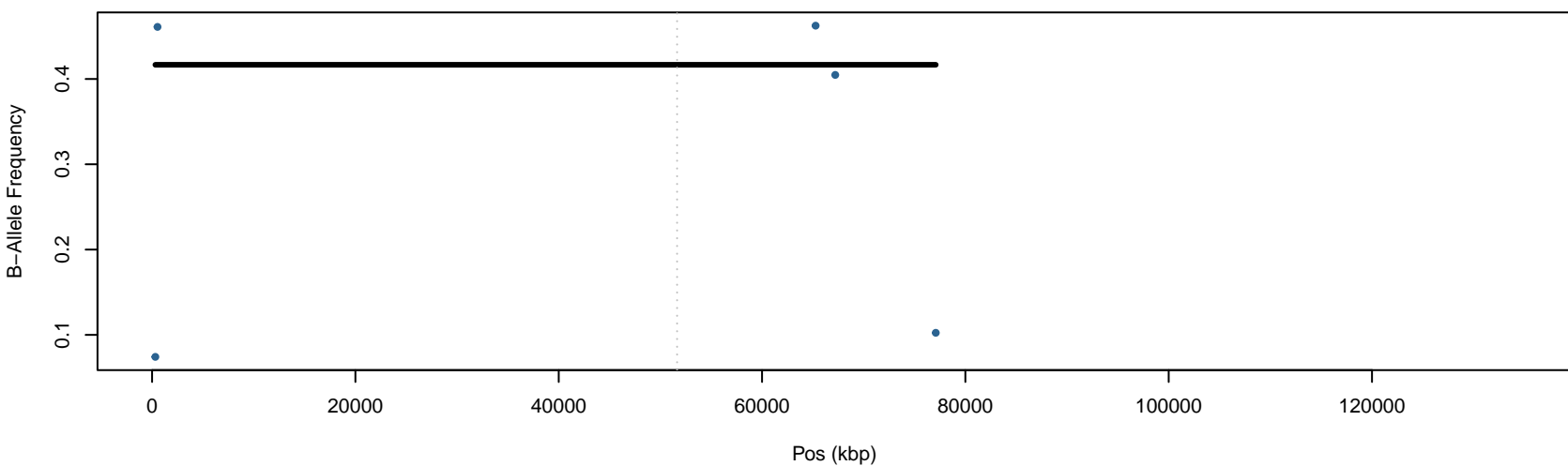
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505,489 Chromosome: 9



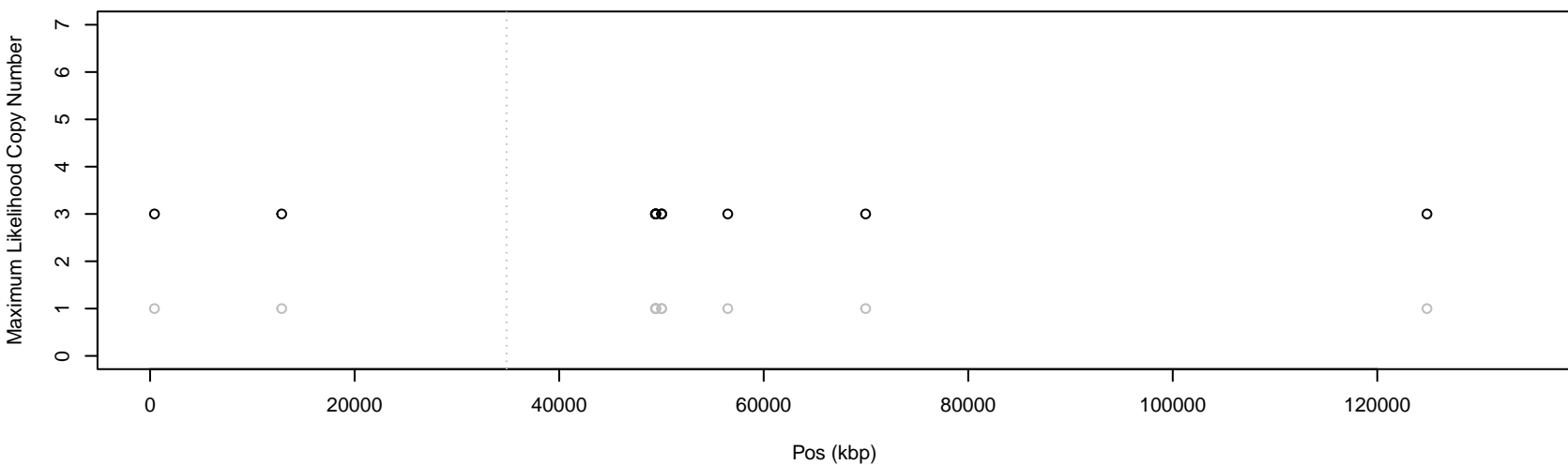
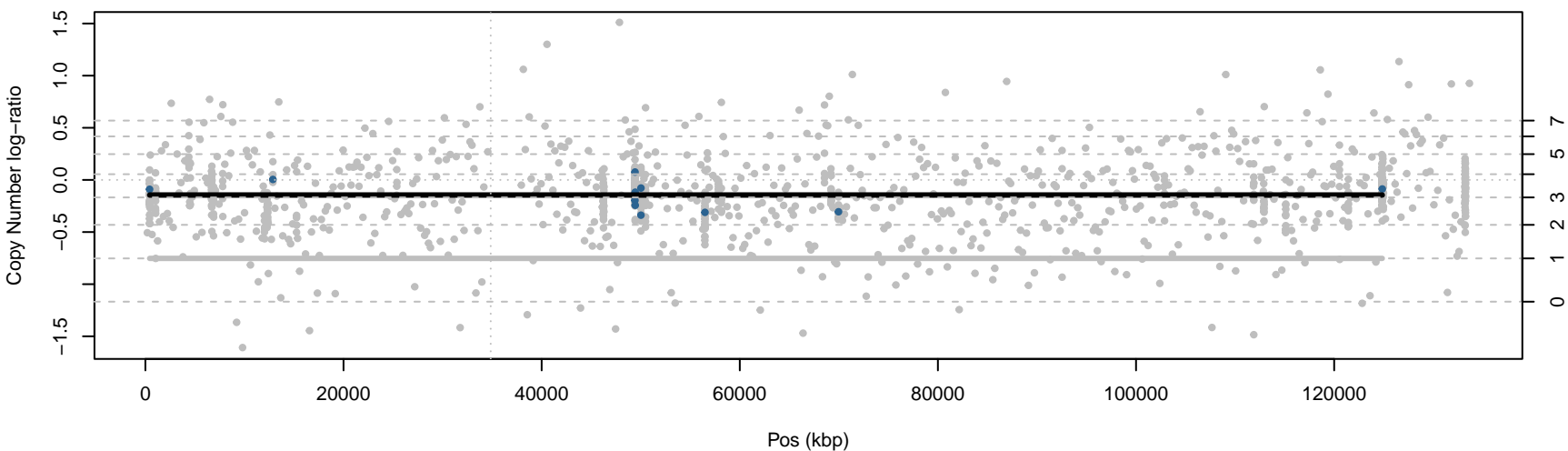
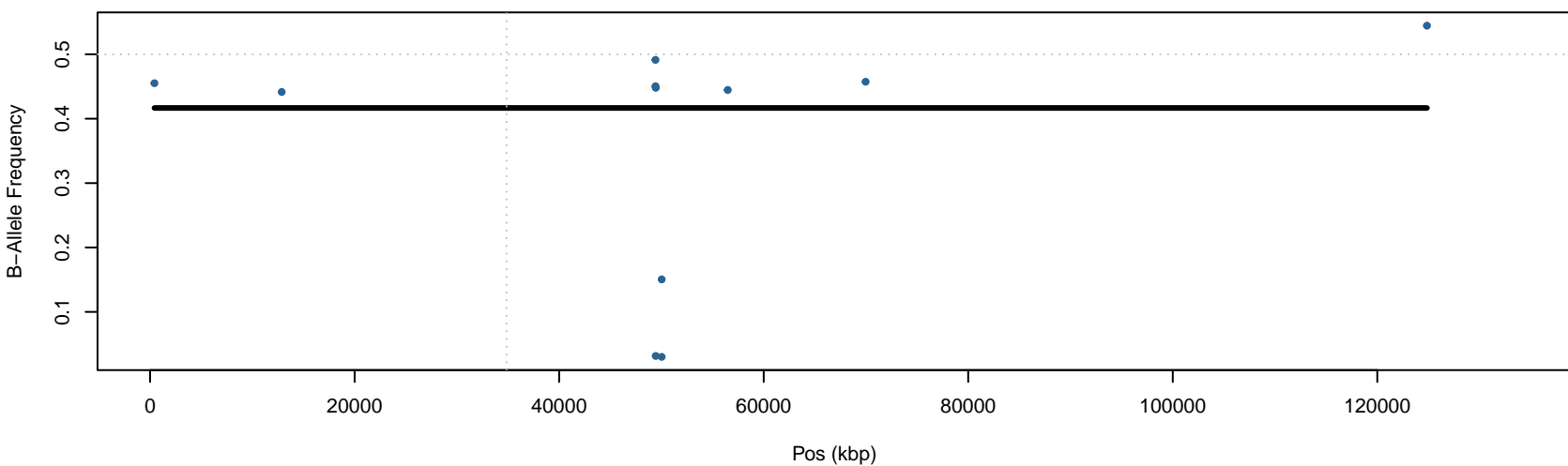
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 10



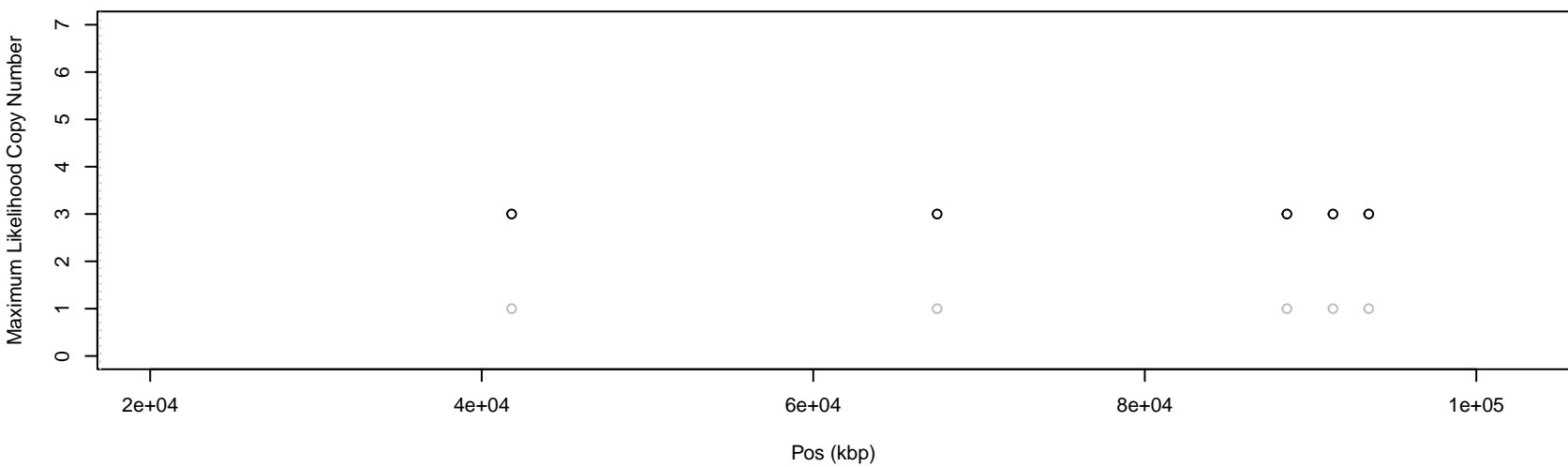
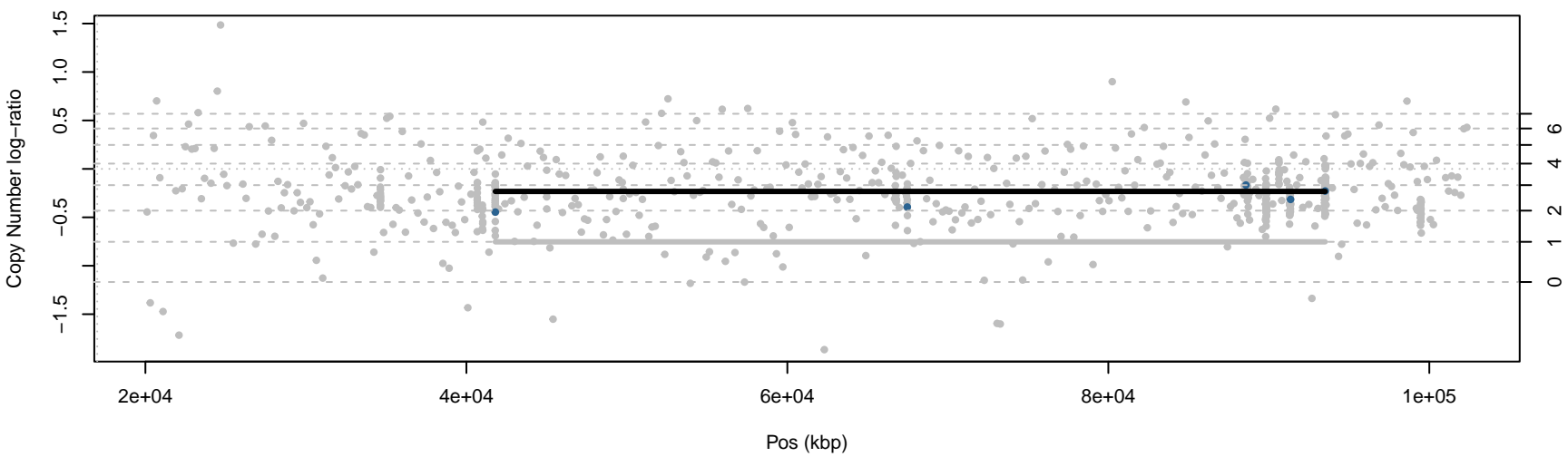
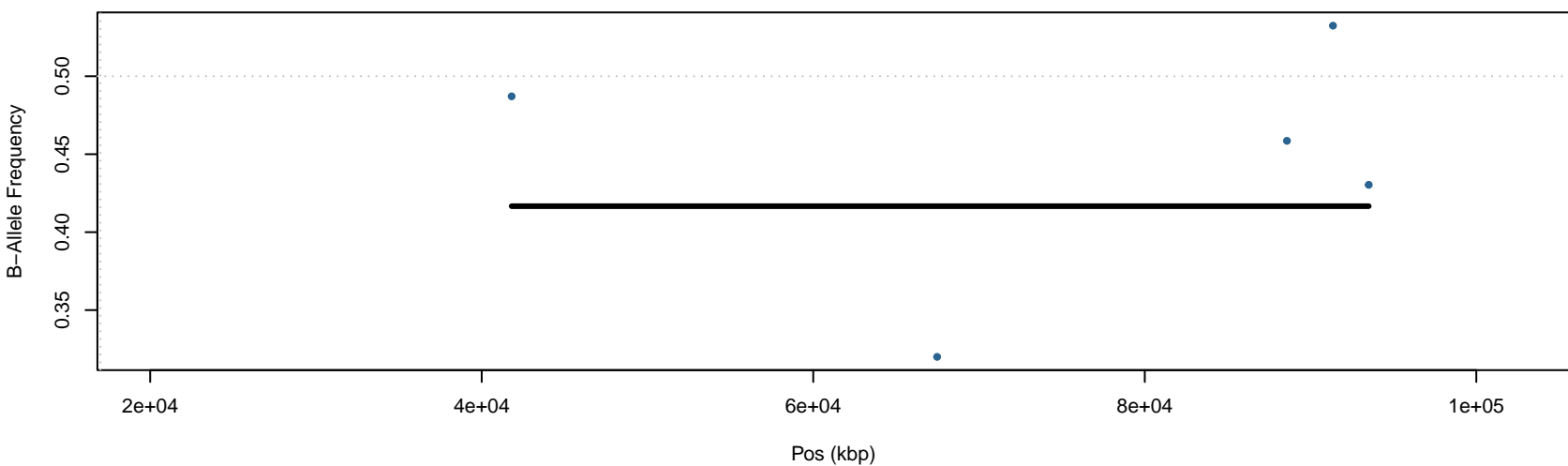
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 11



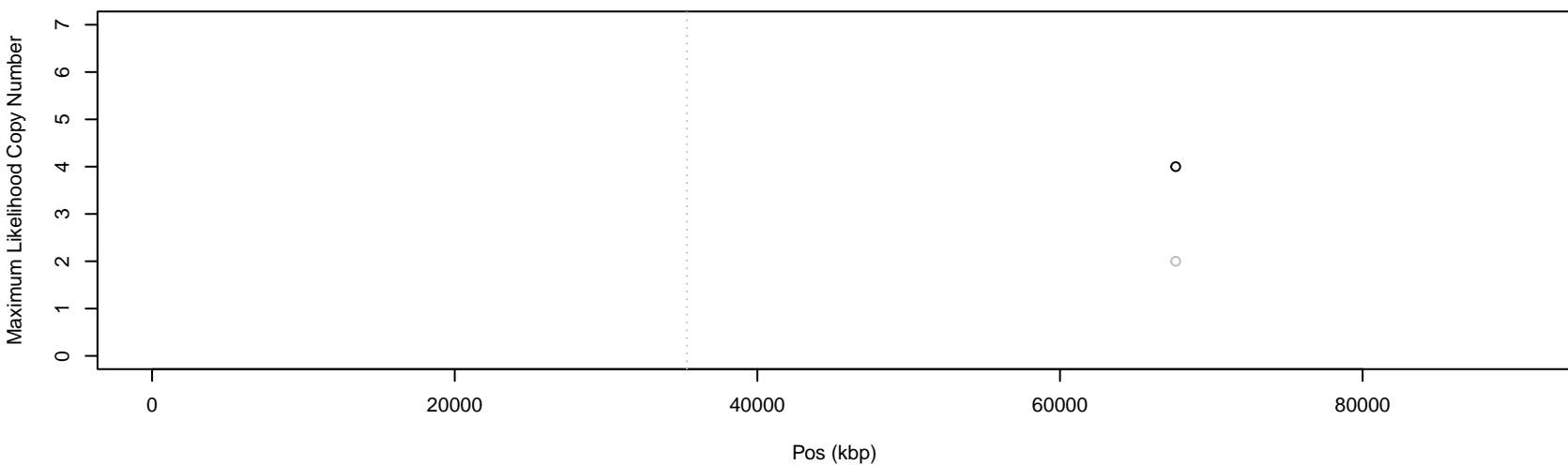
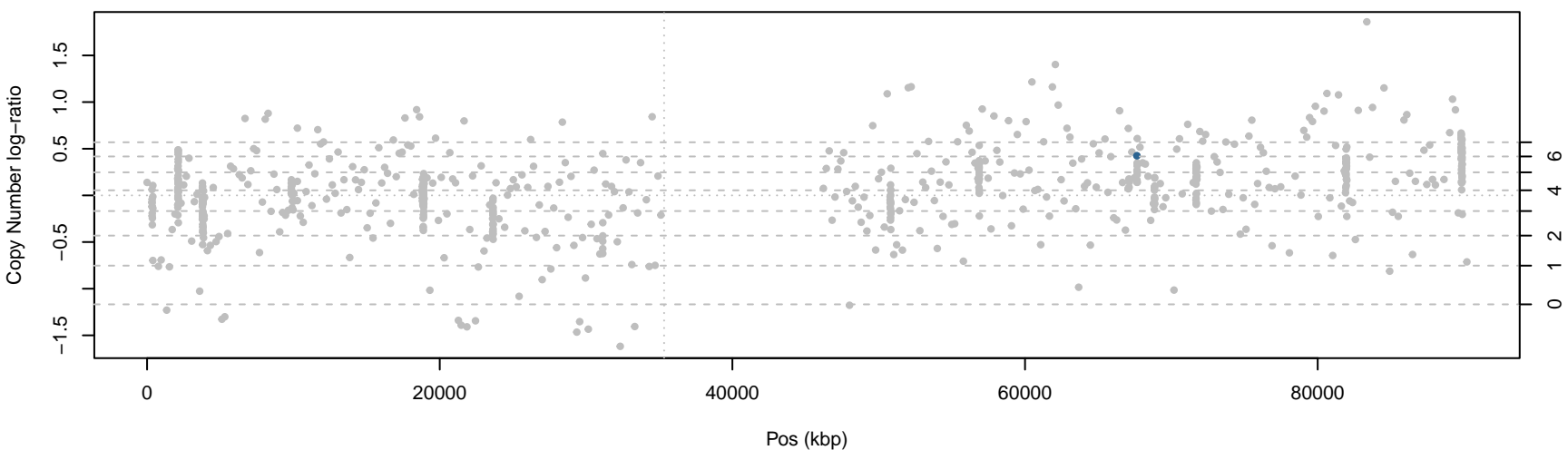
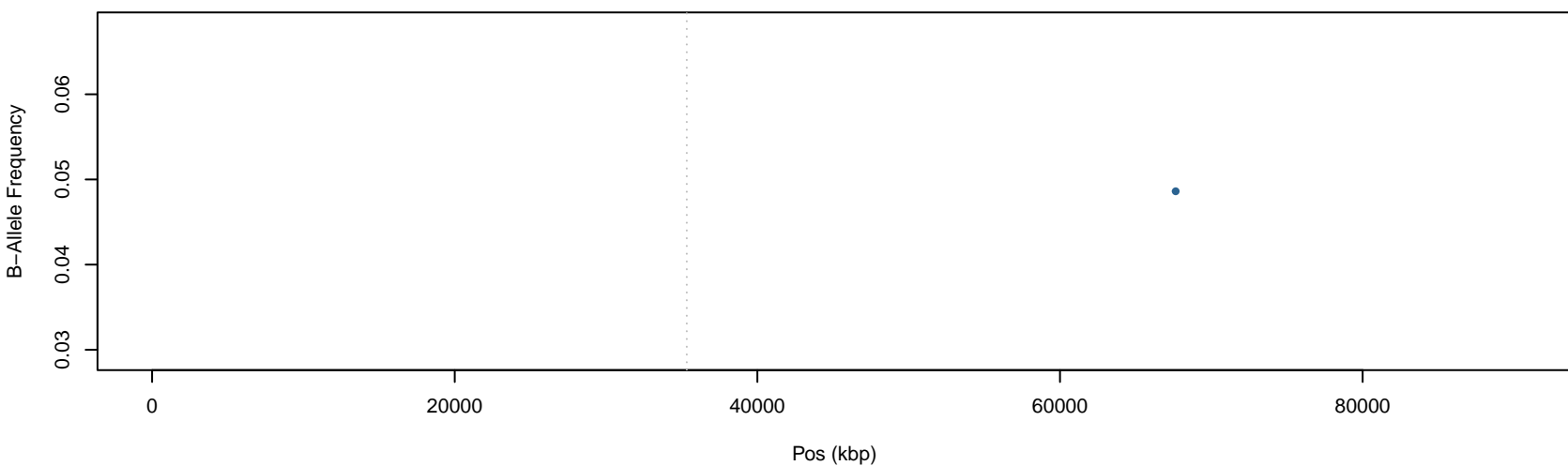
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 12



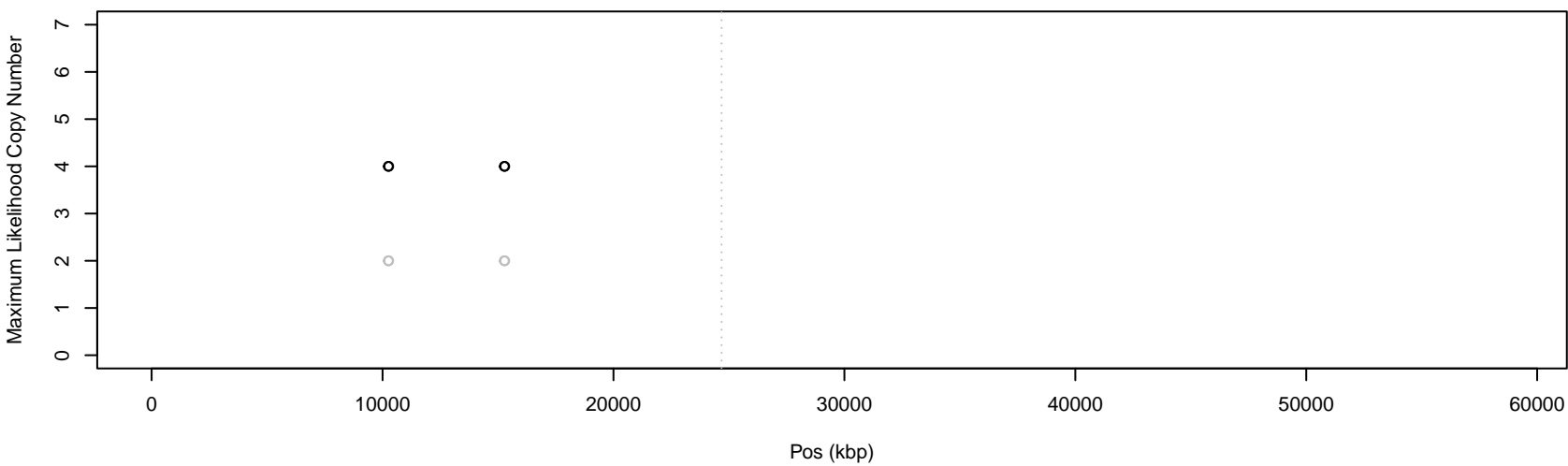
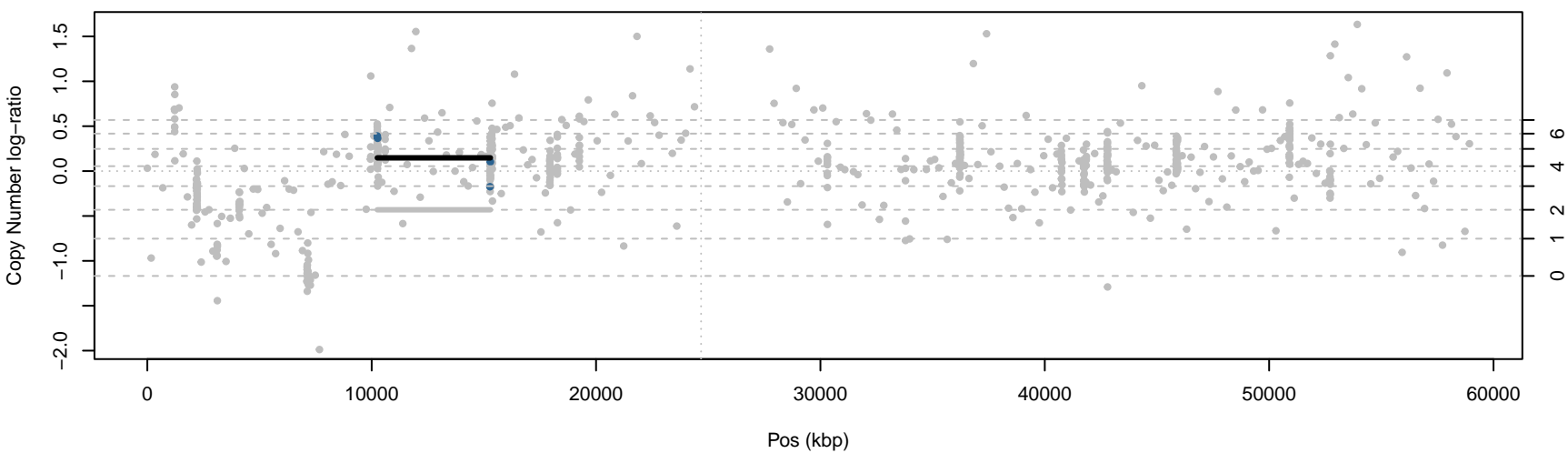
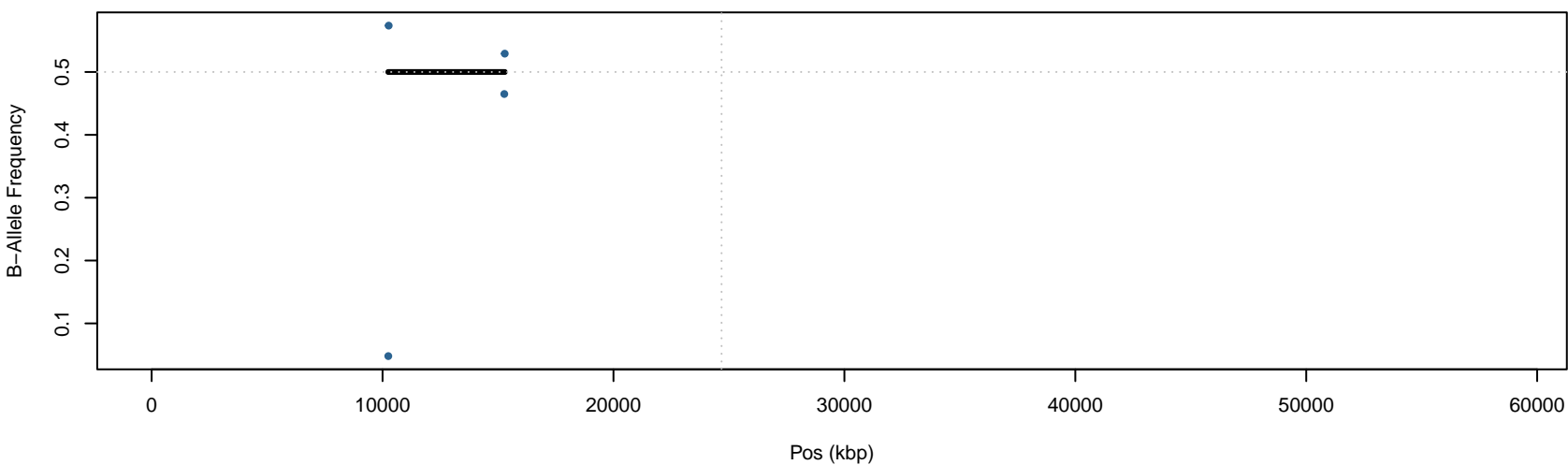
Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 15



Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 16



Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 19



Purity: 0.4 Tumor ploidy: 3.74 SNV log-likelihood: -57.73 GoF: 87% Mean coverage: 505;489 Chromosome: 21

