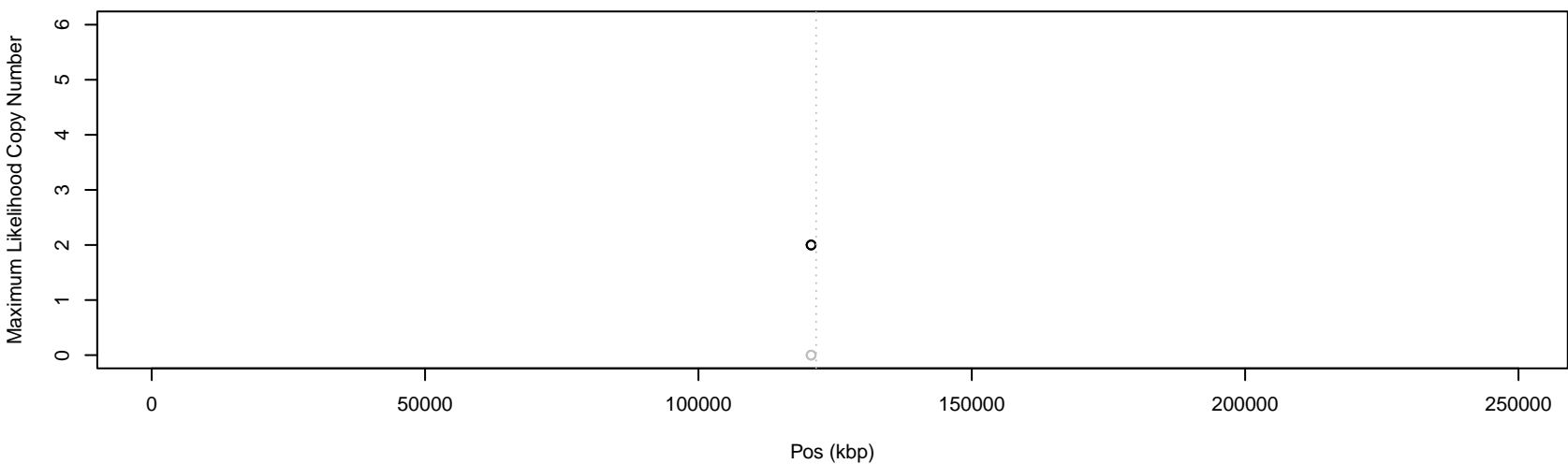
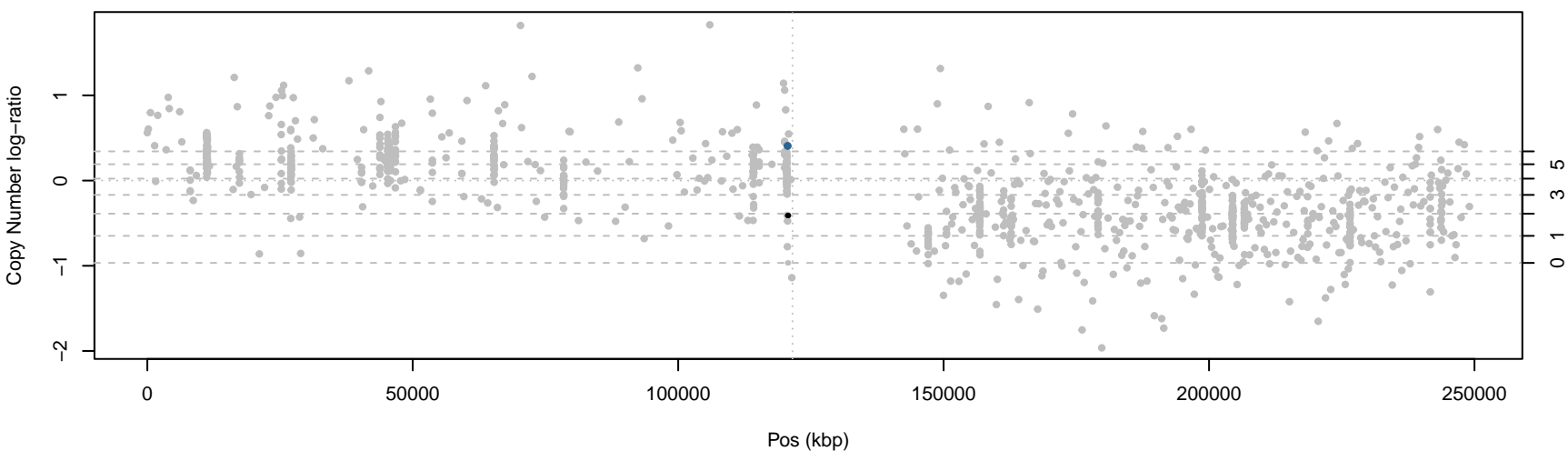
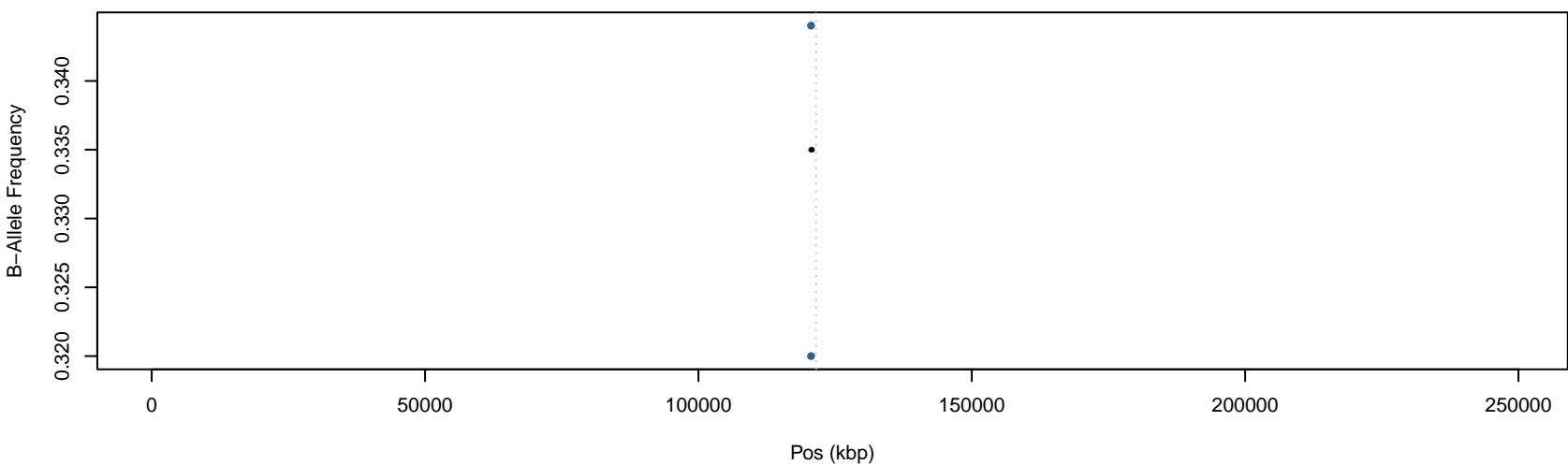
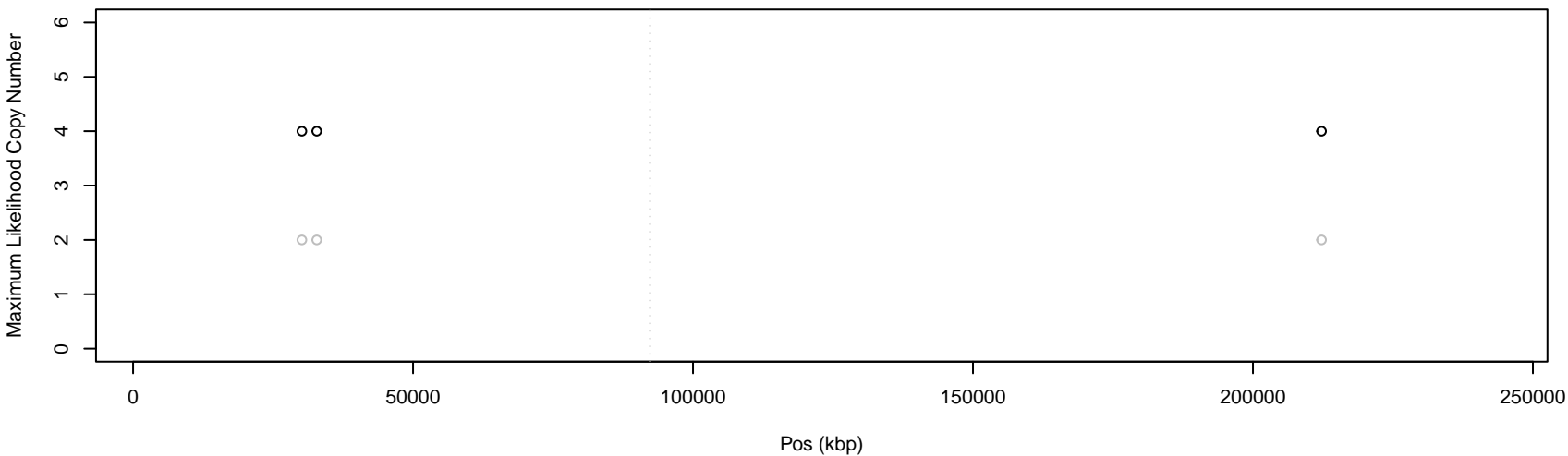
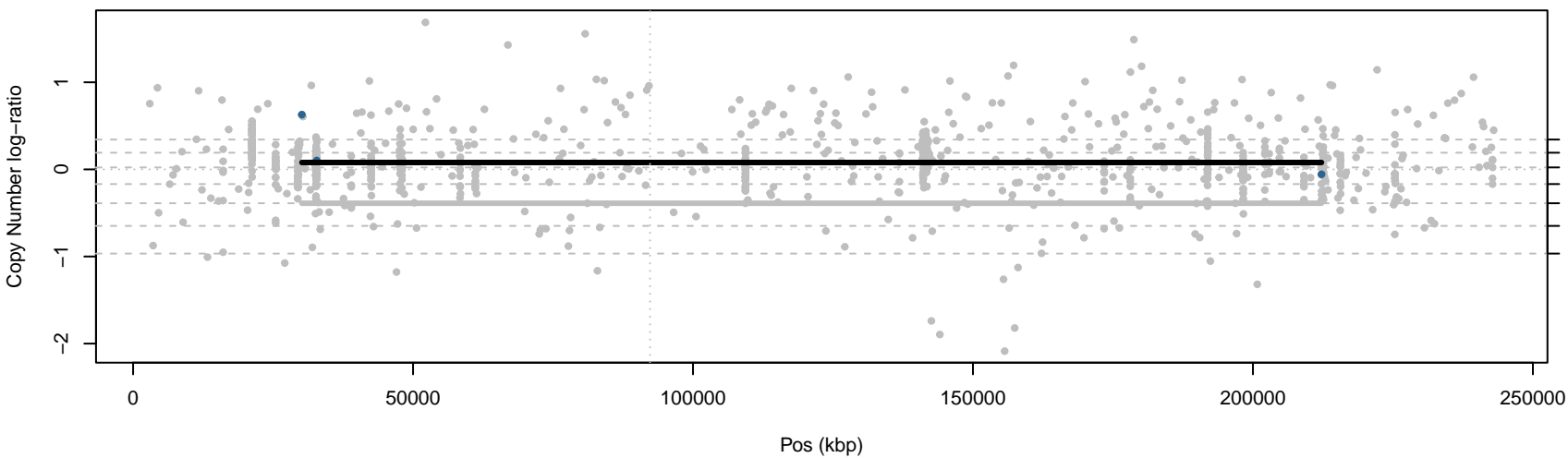
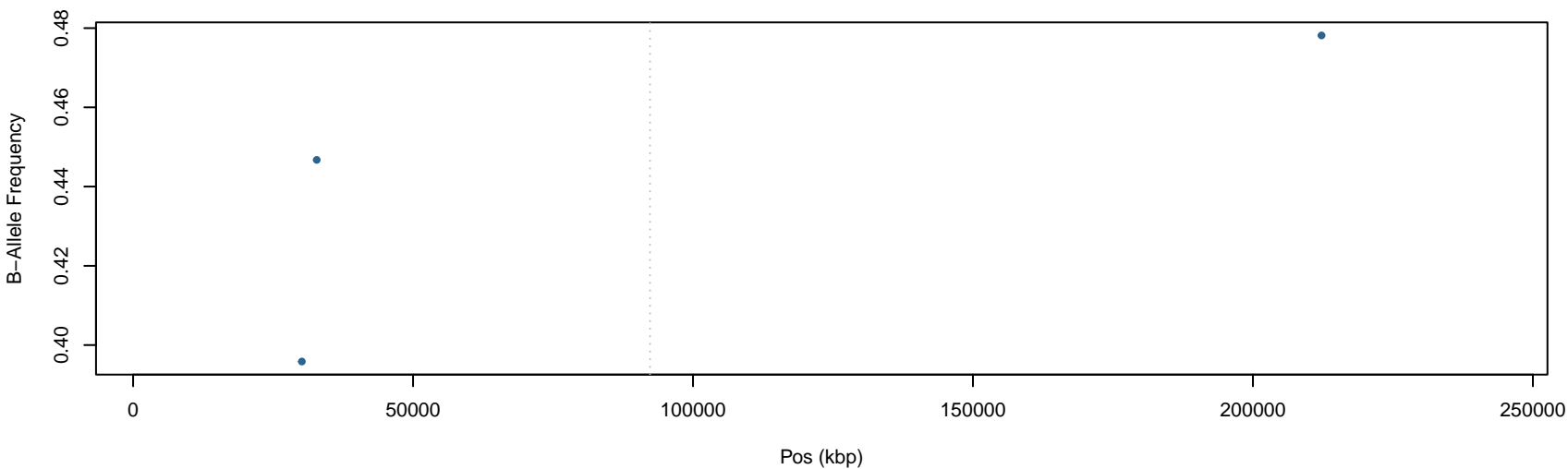


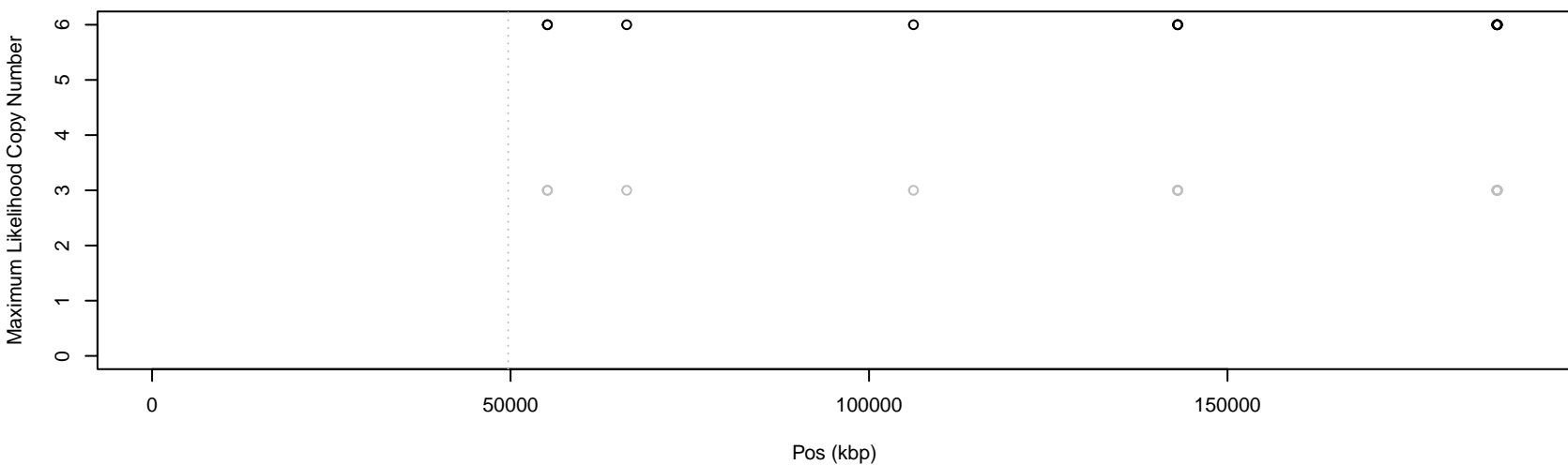
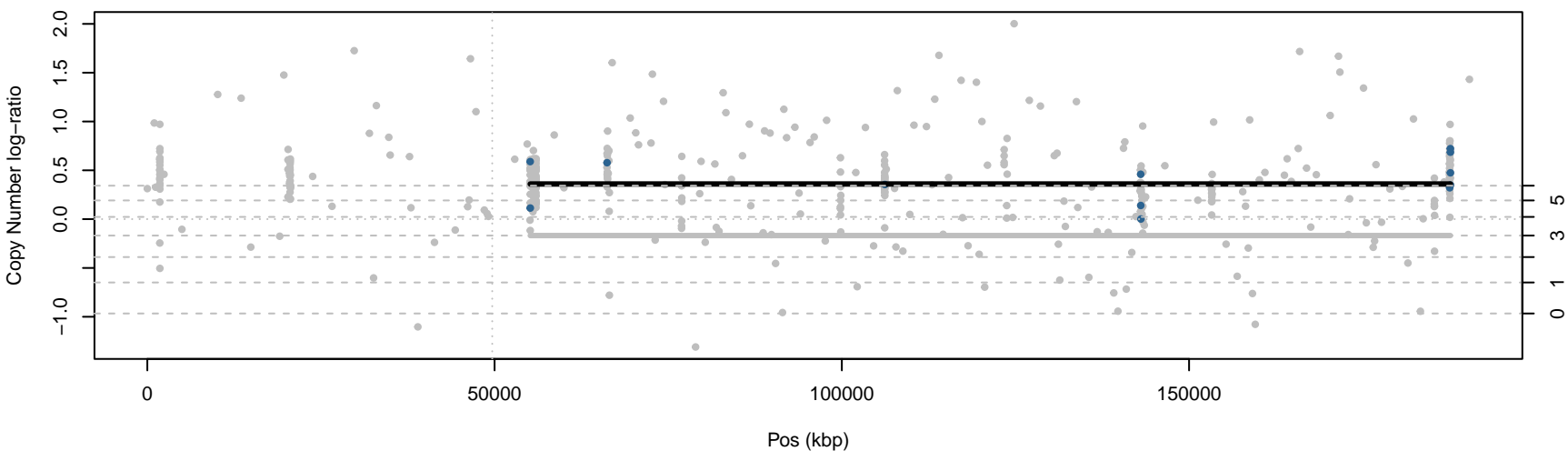
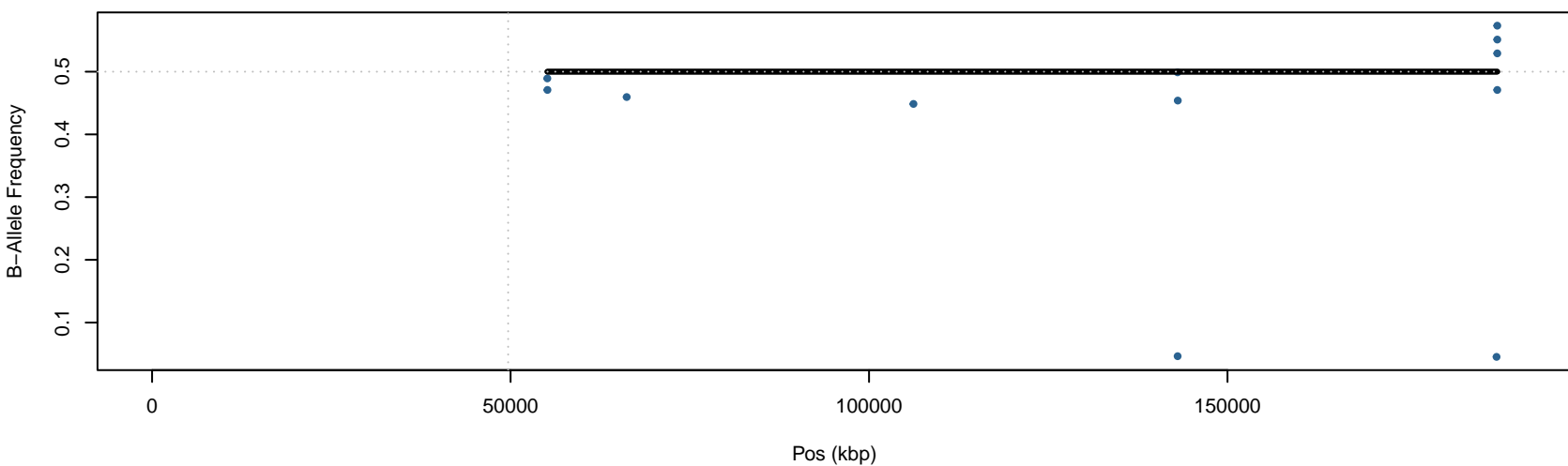
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 1



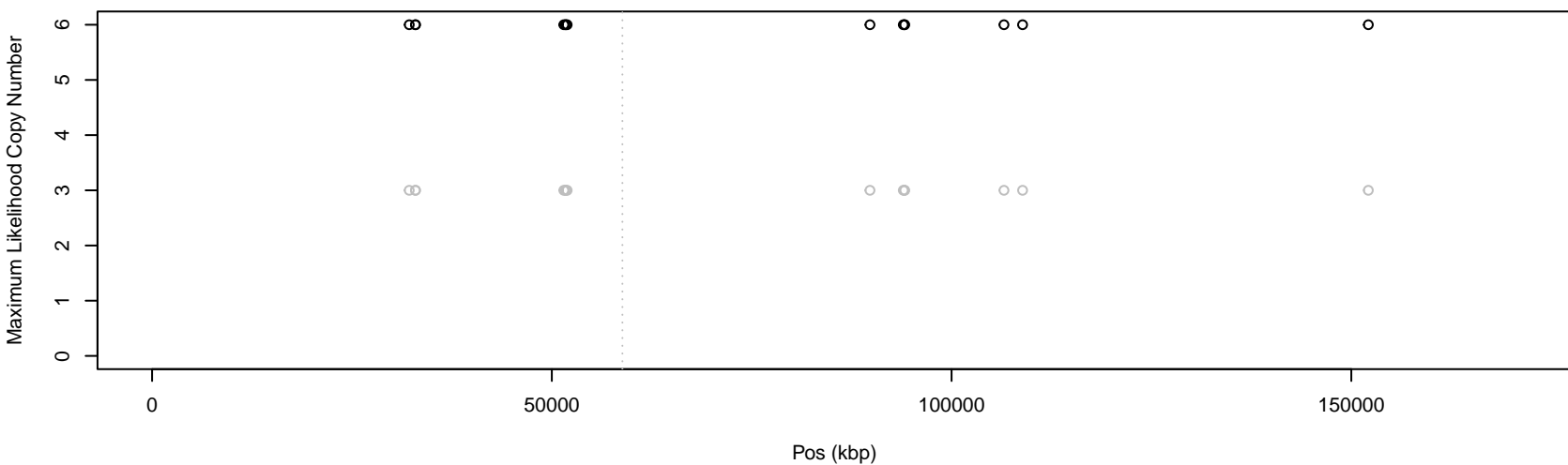
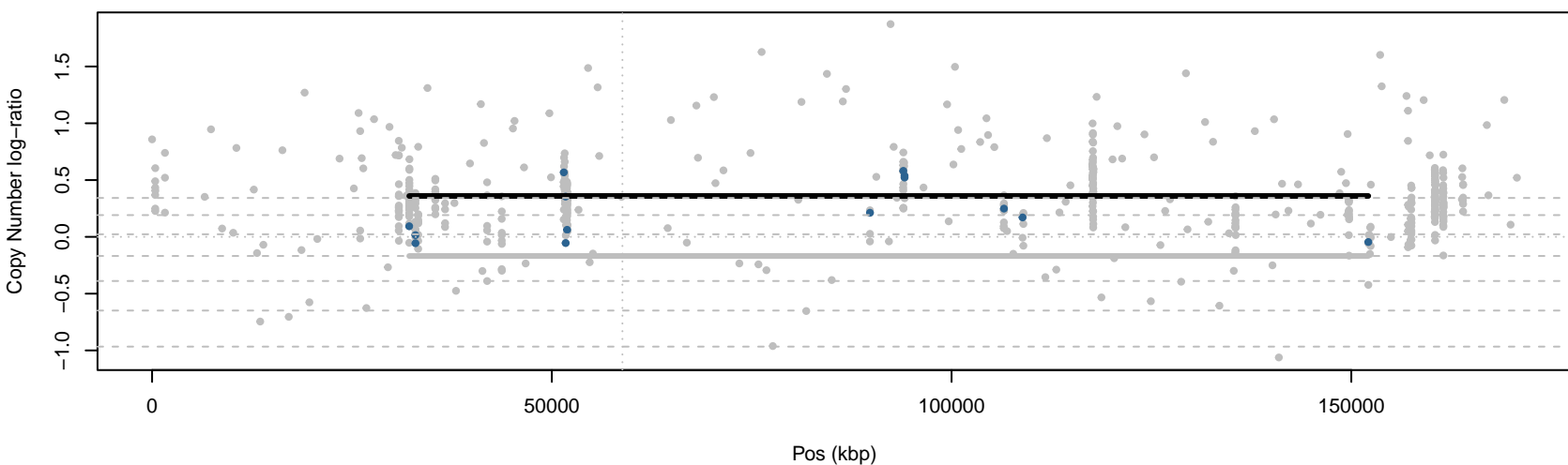
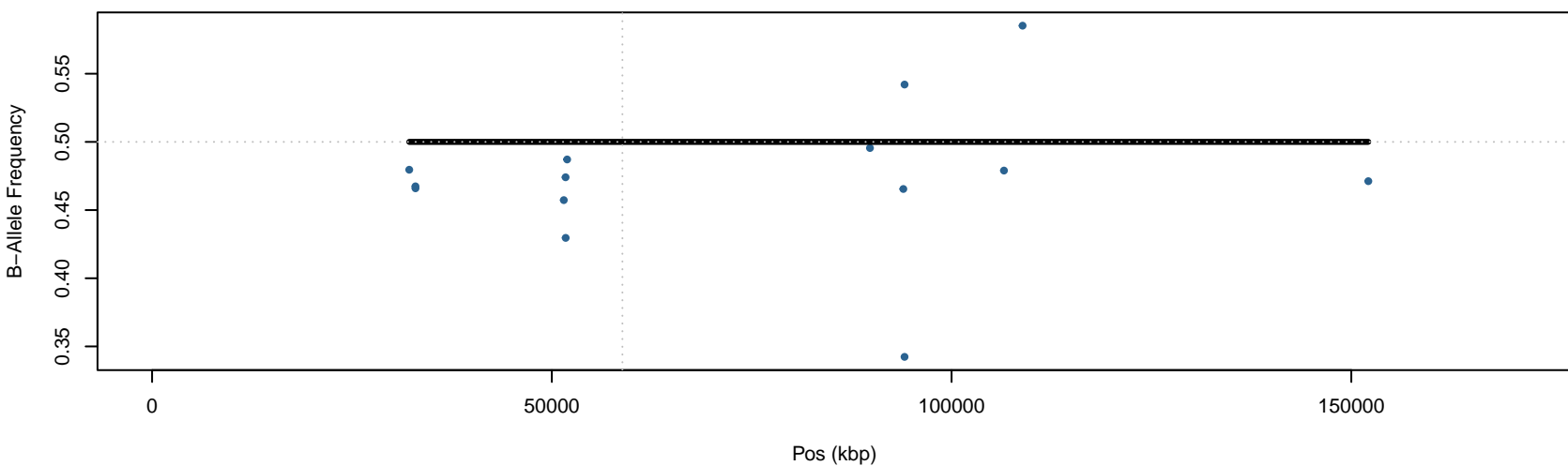
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 2



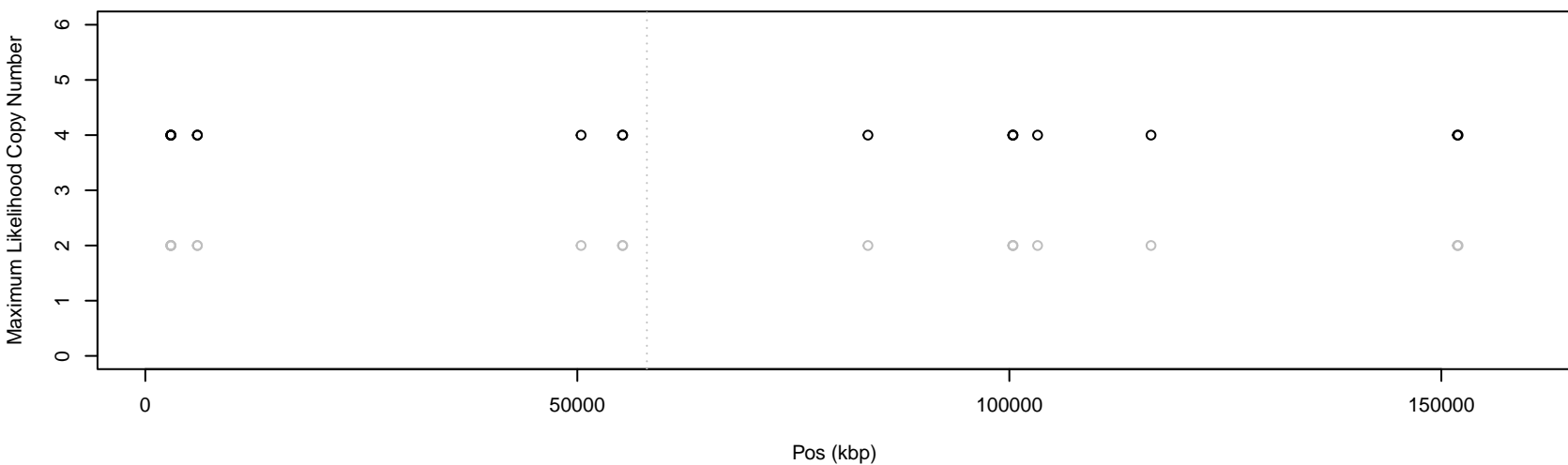
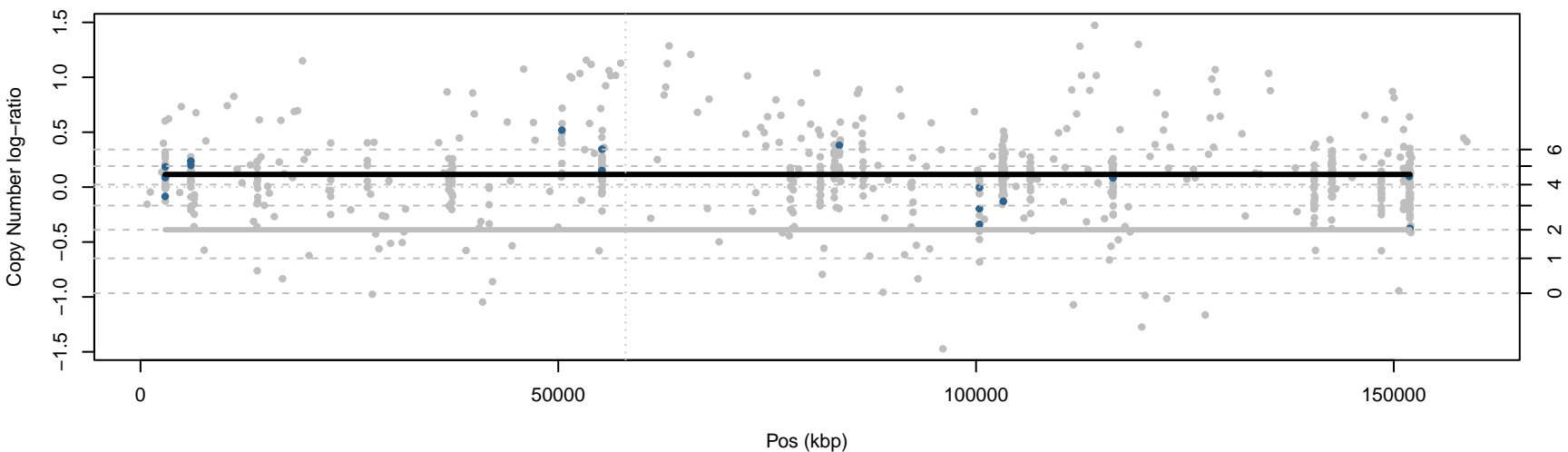
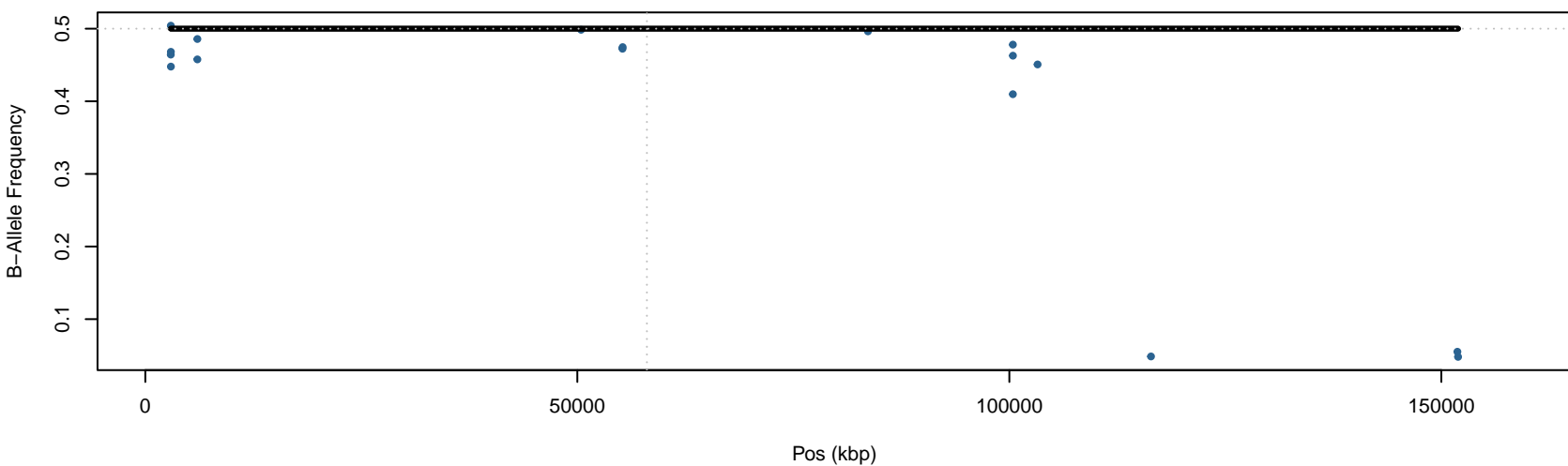
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 4



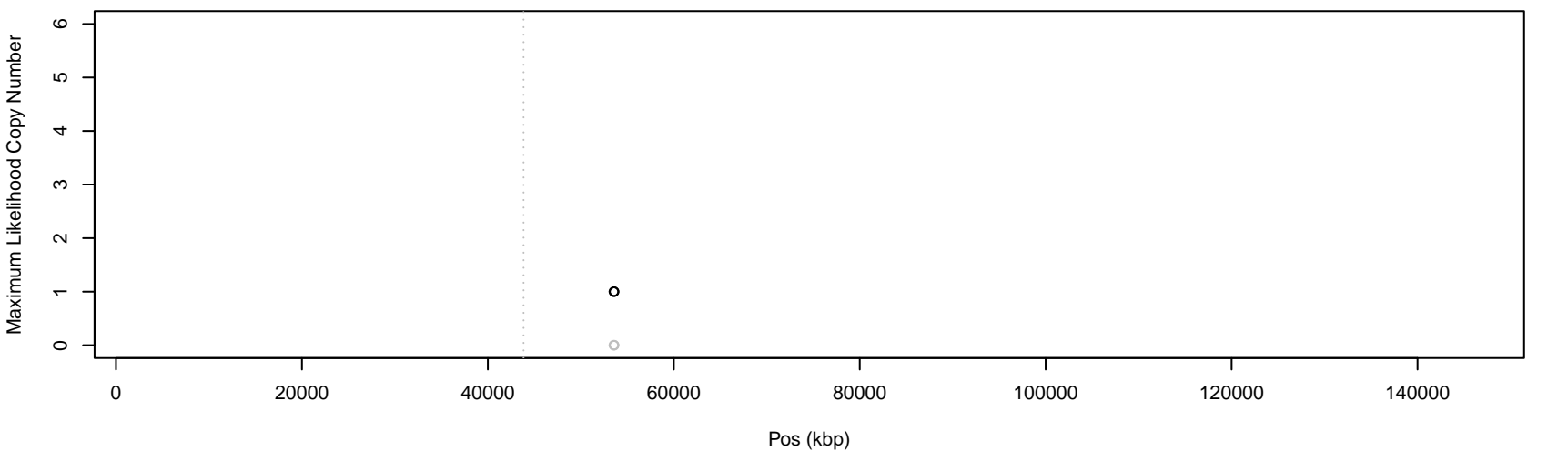
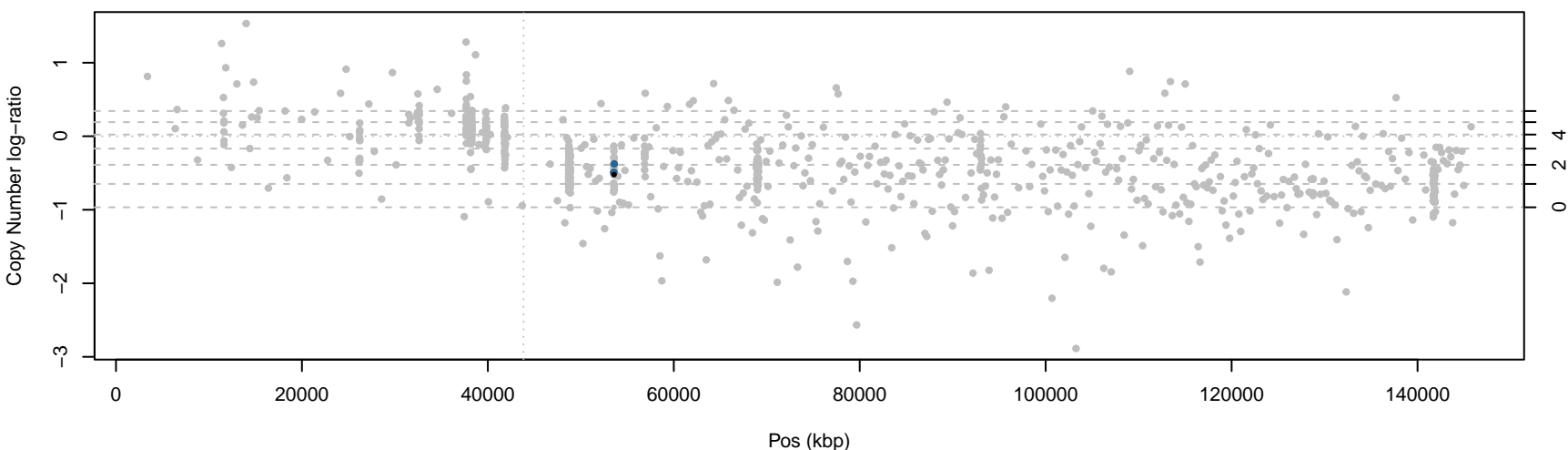
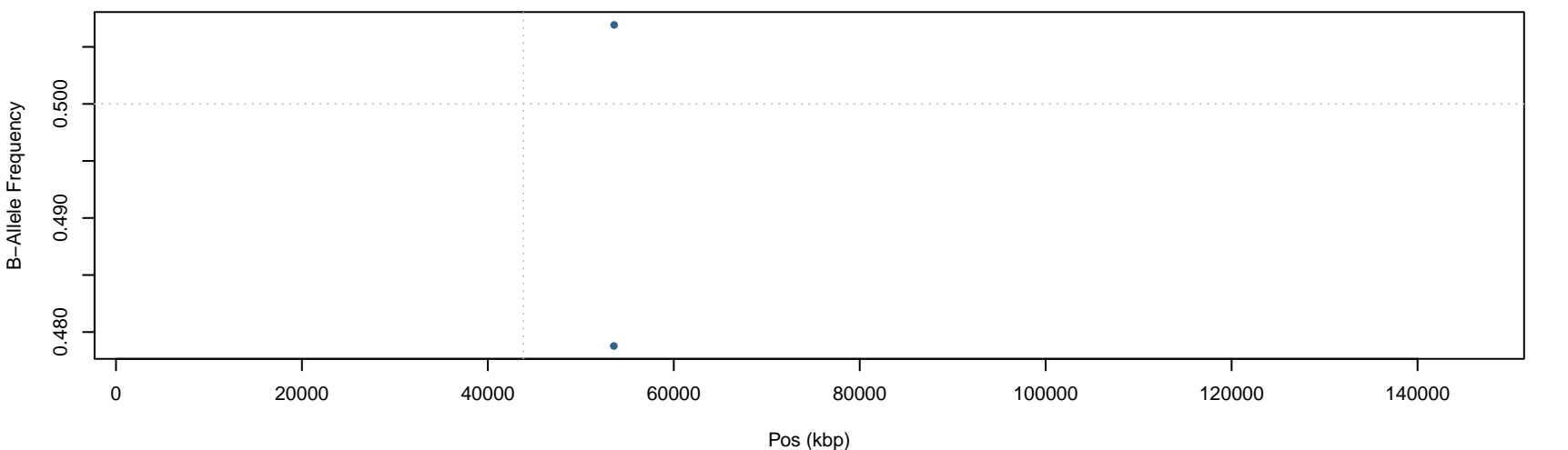
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 6



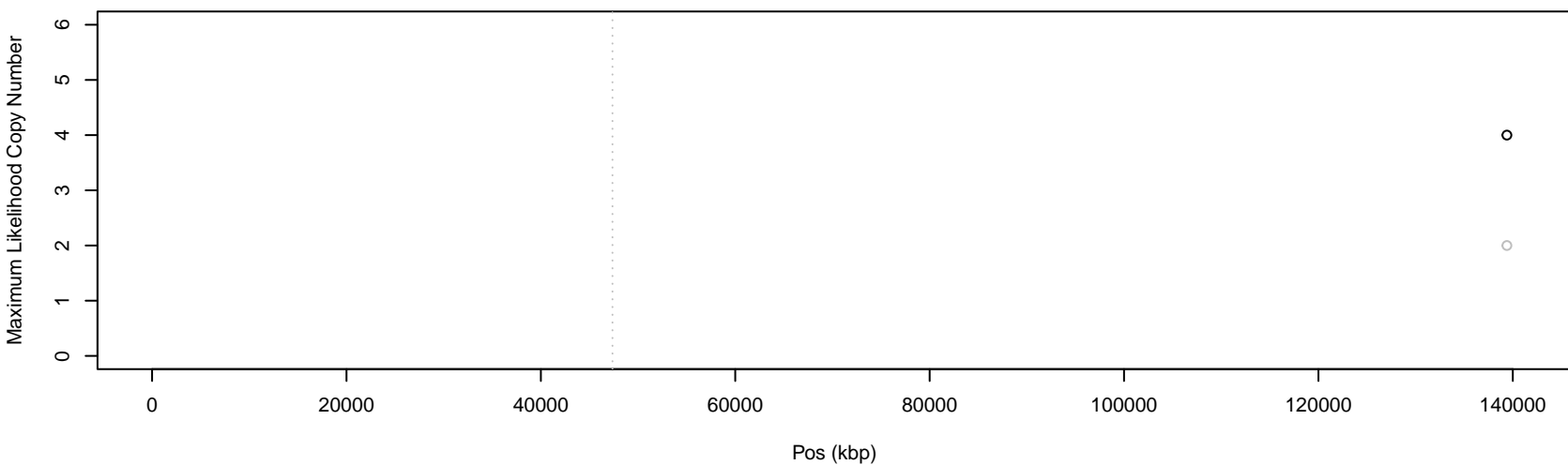
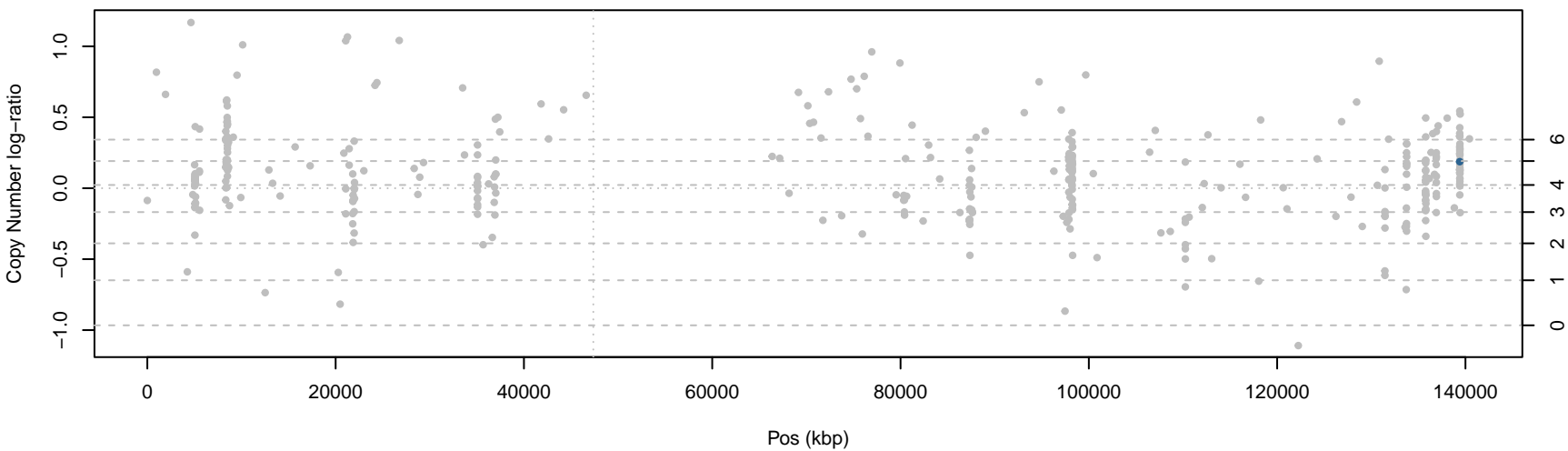
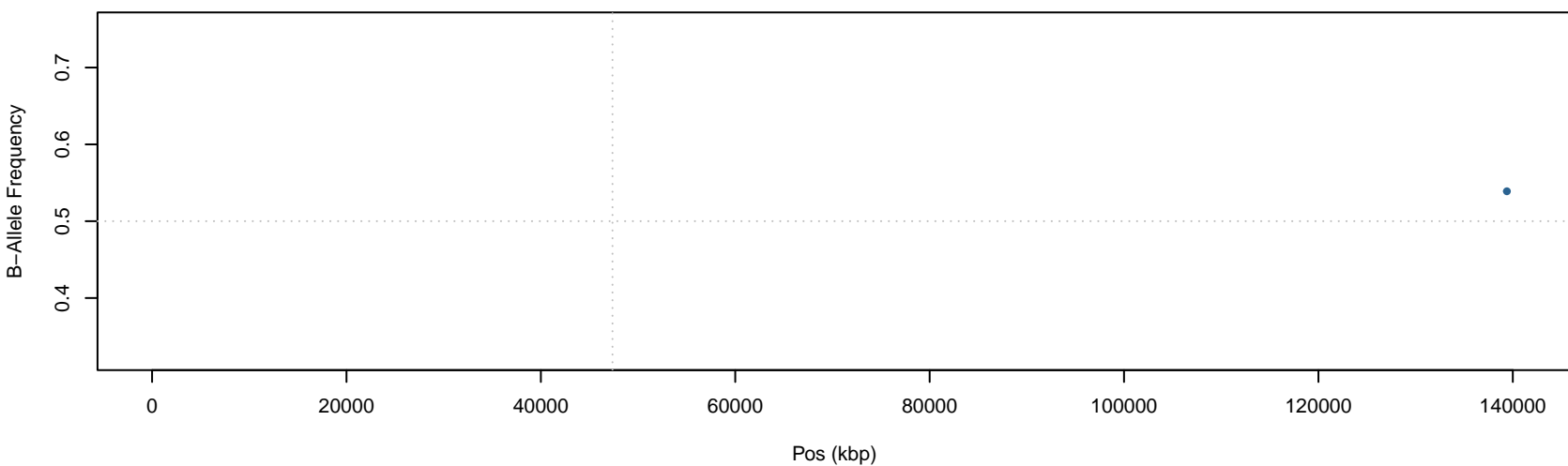
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 7



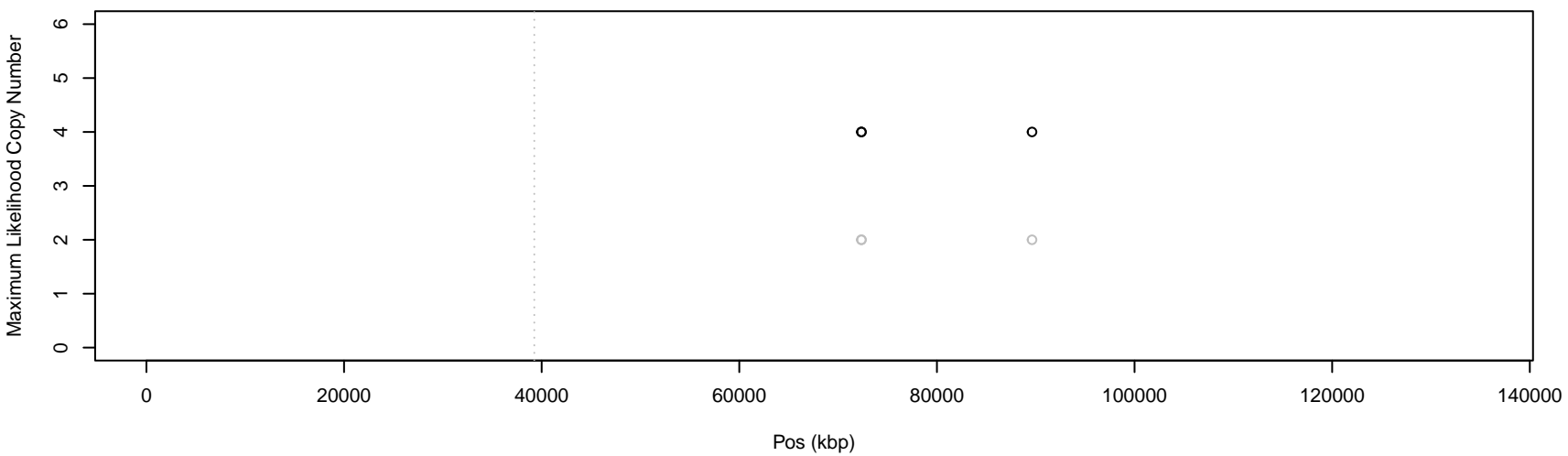
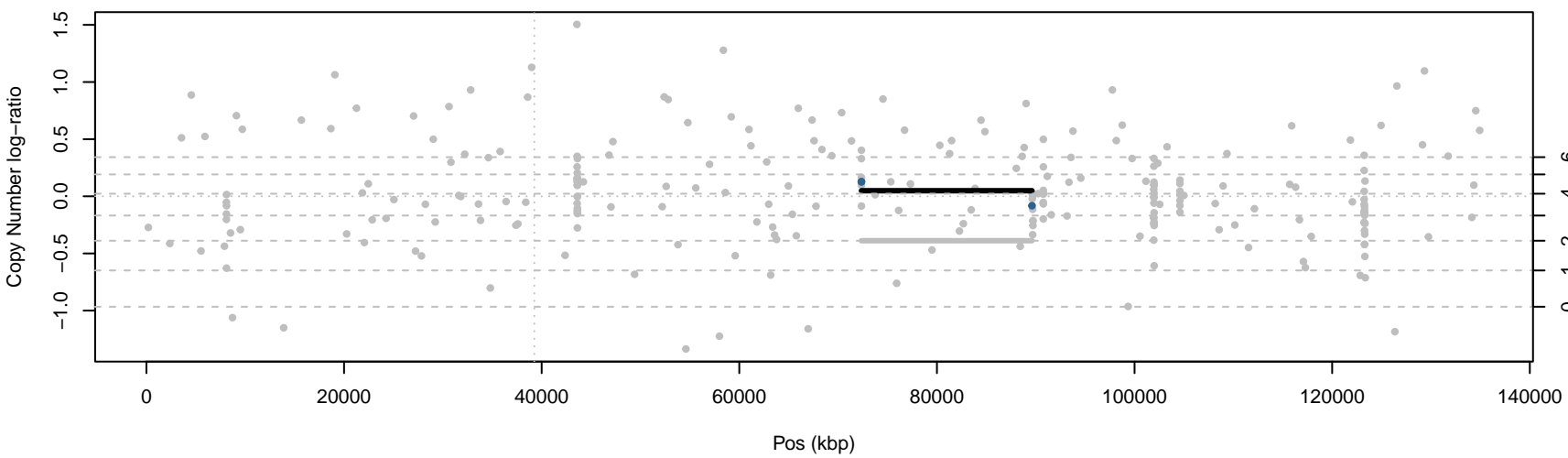
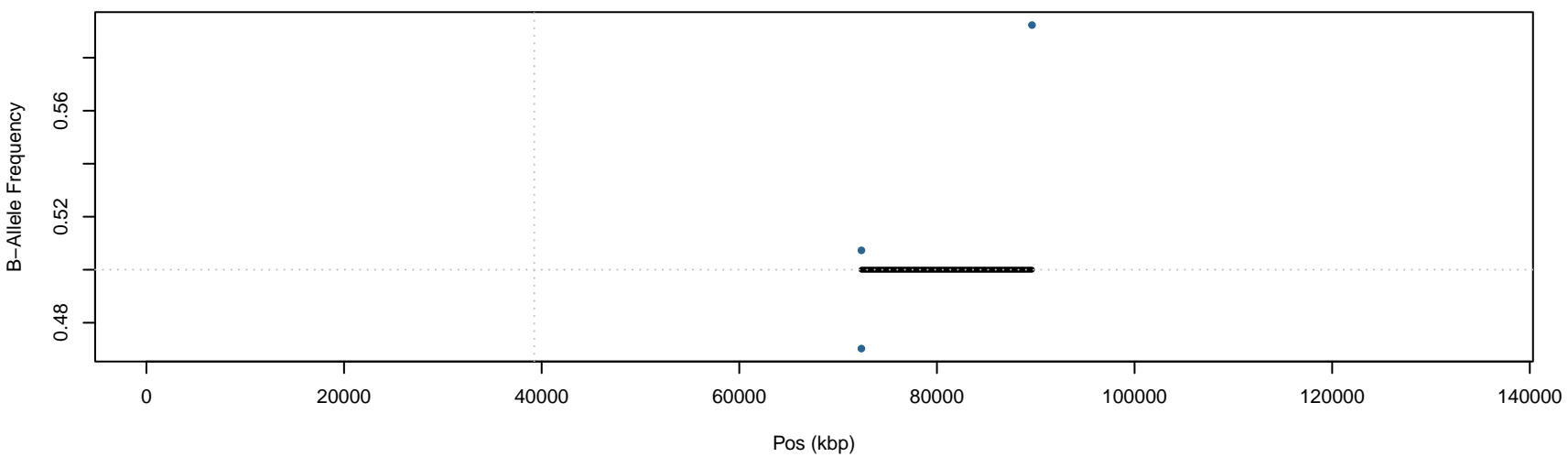
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 8



Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 9

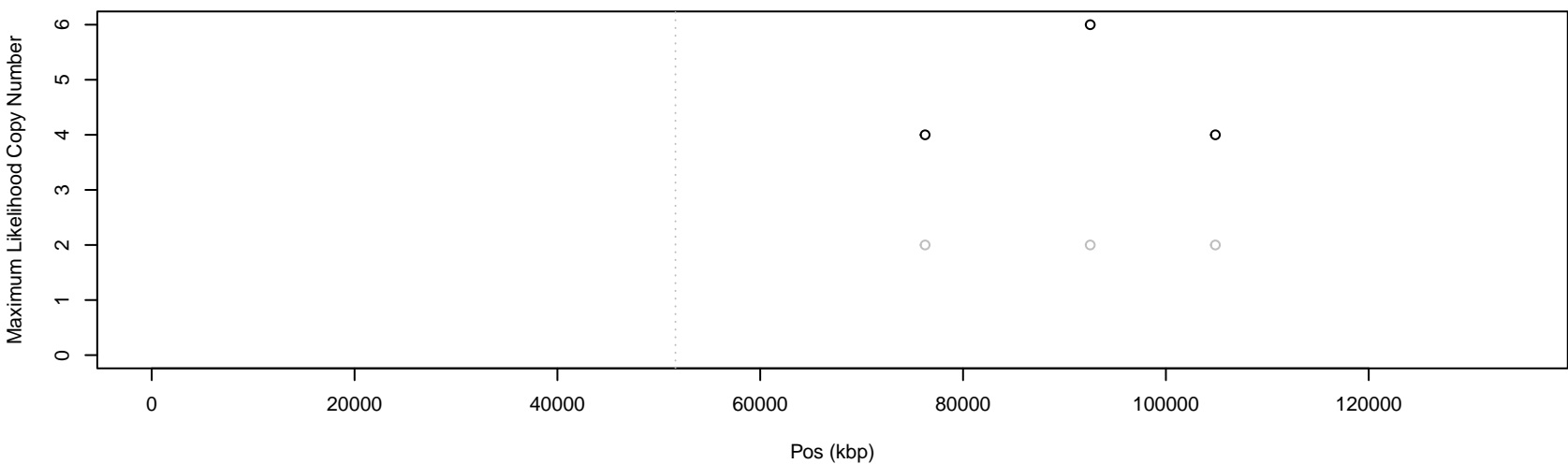
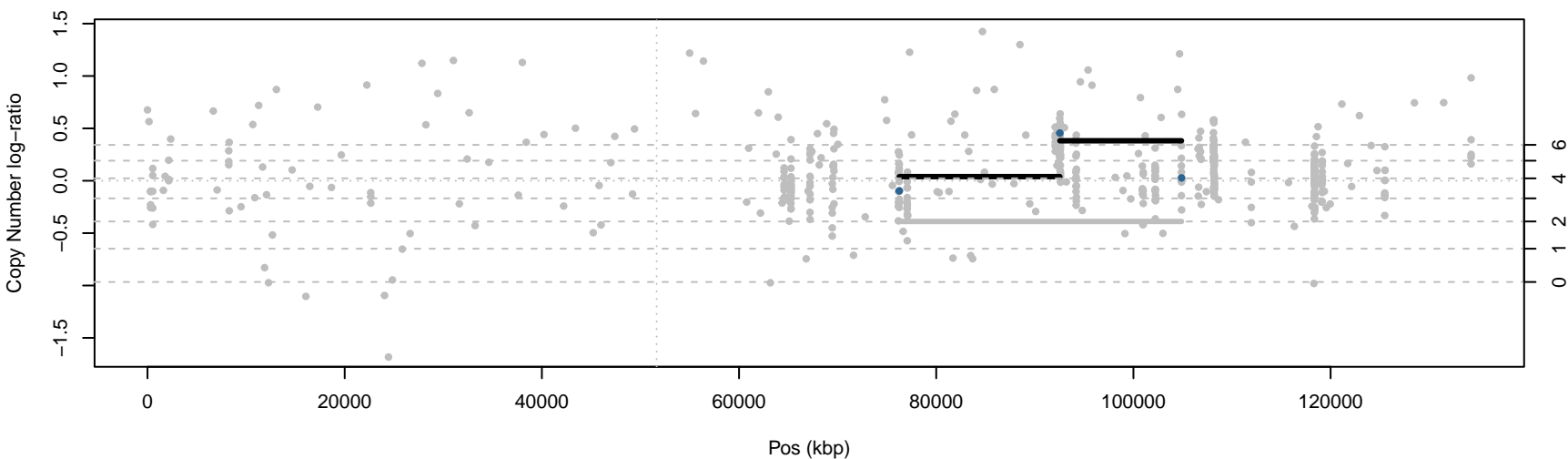
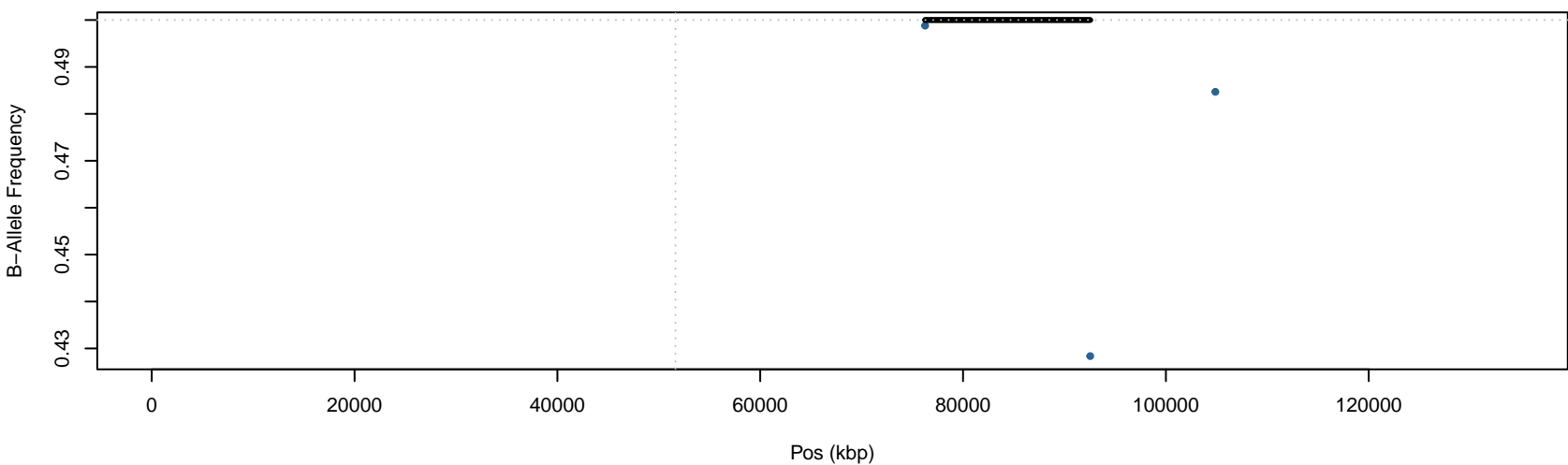


Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 10

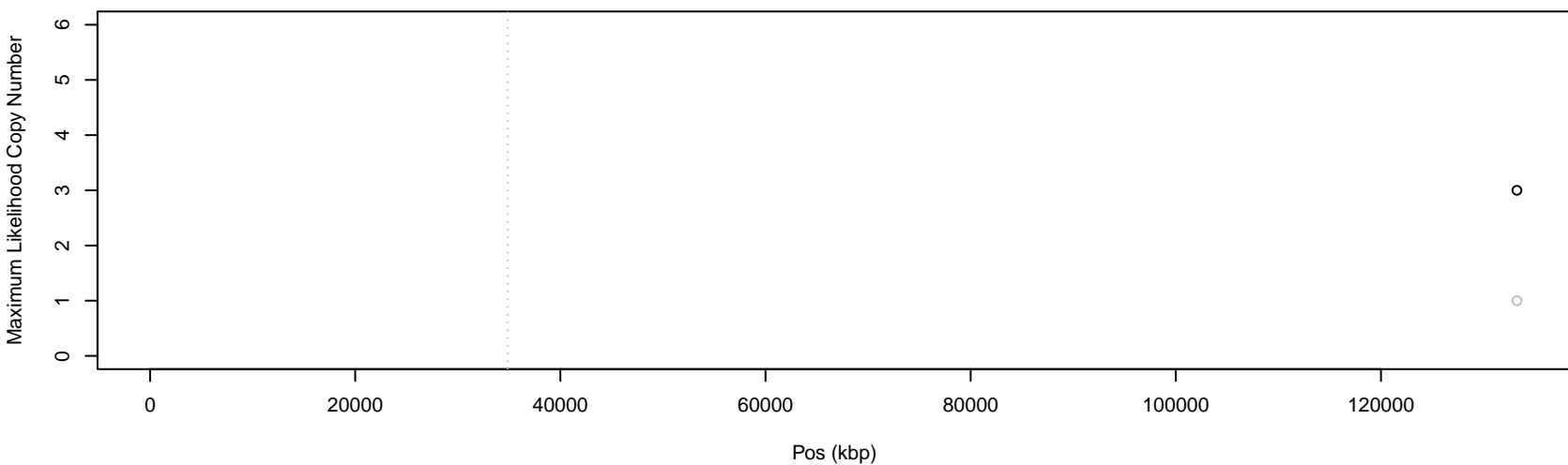
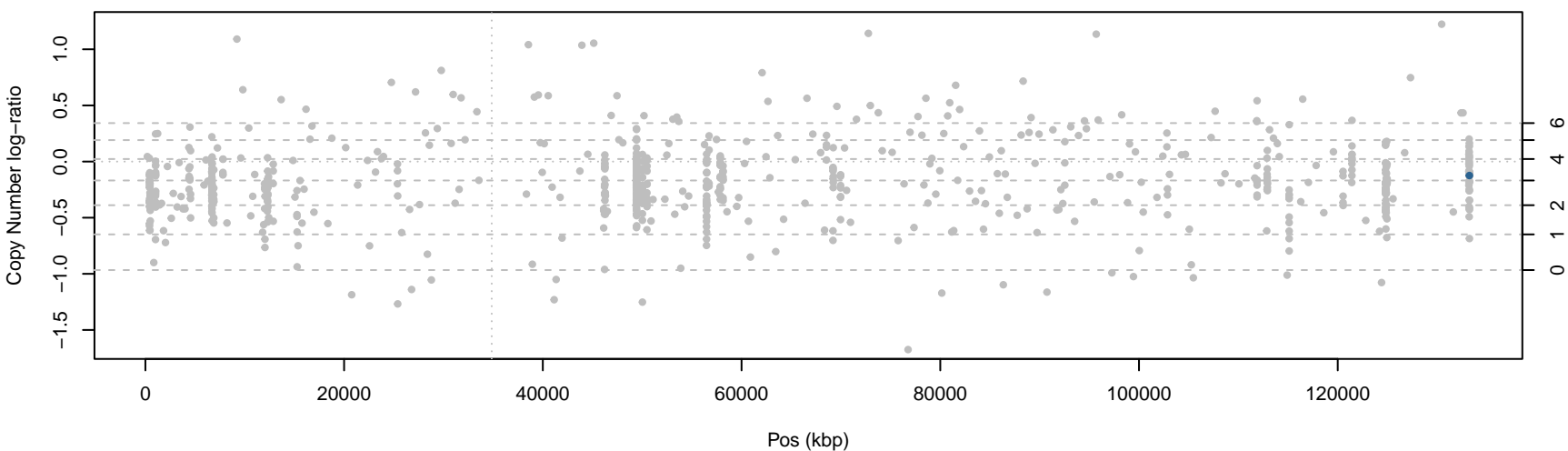
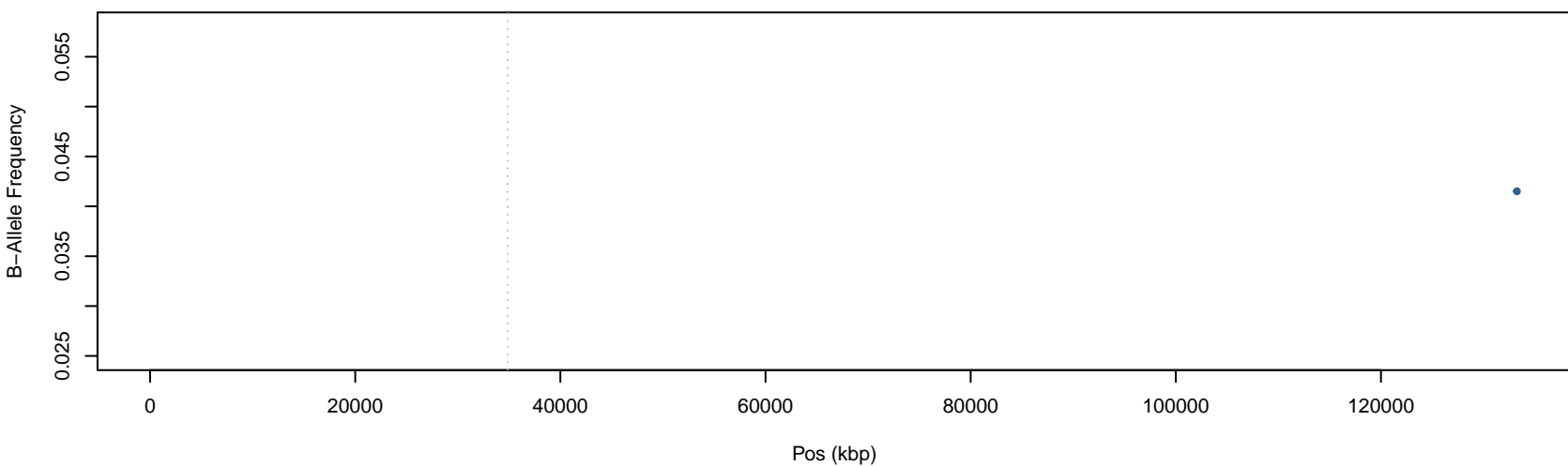




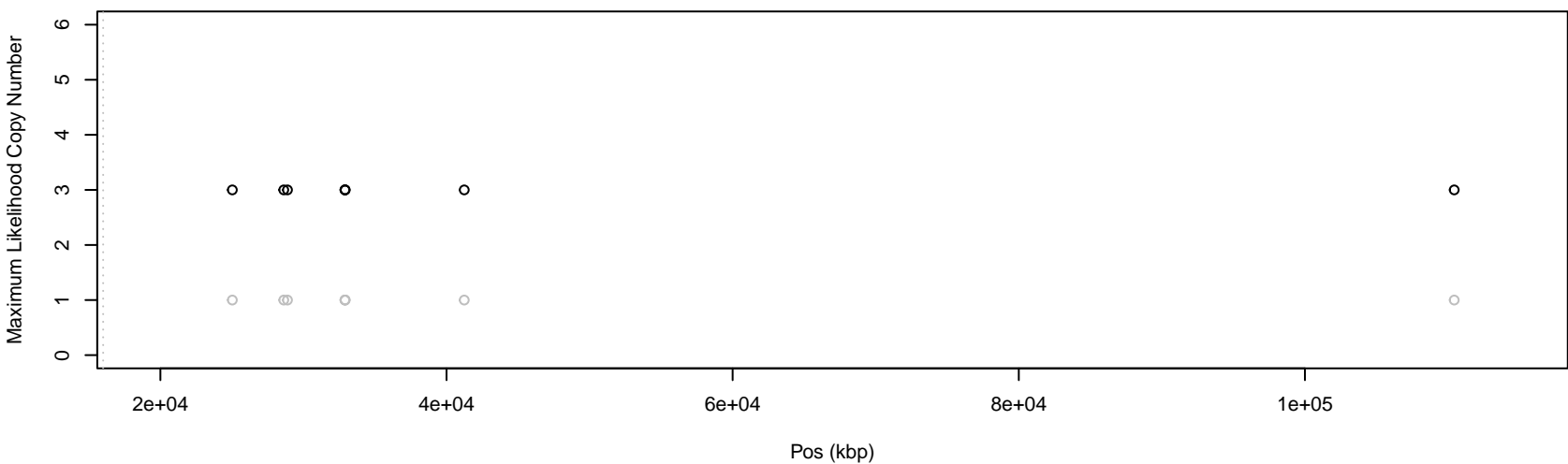
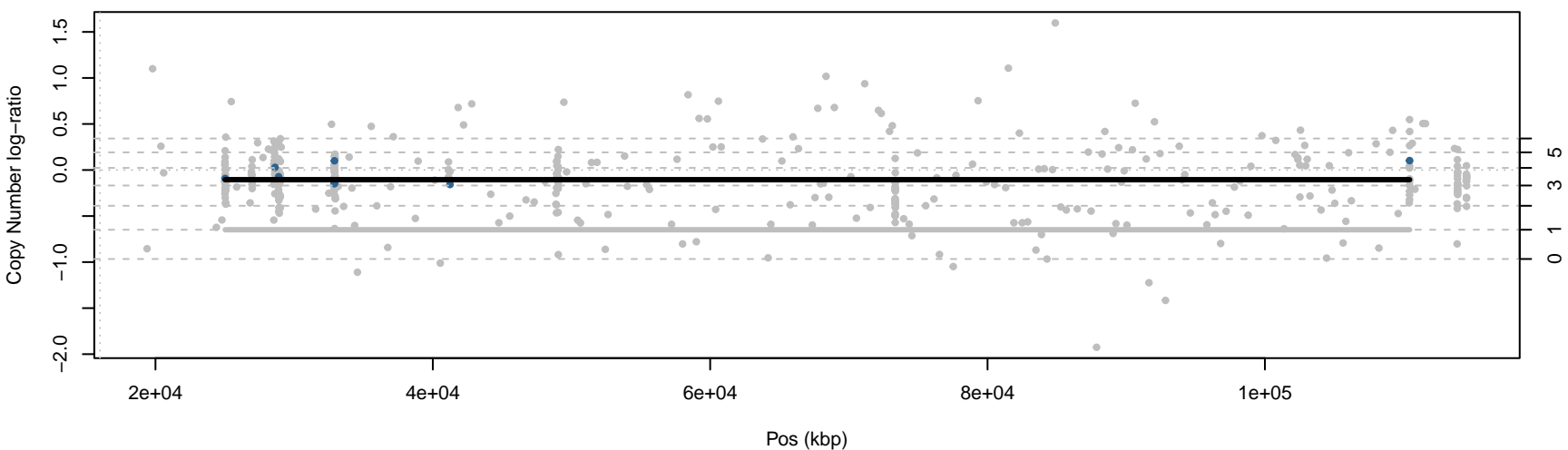
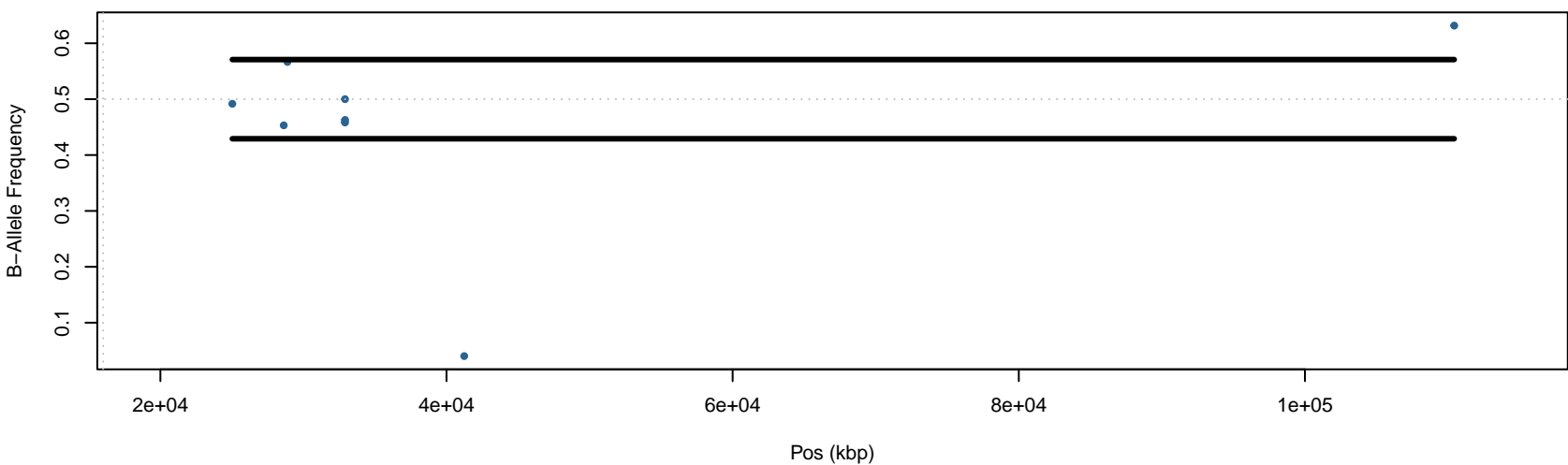
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 11



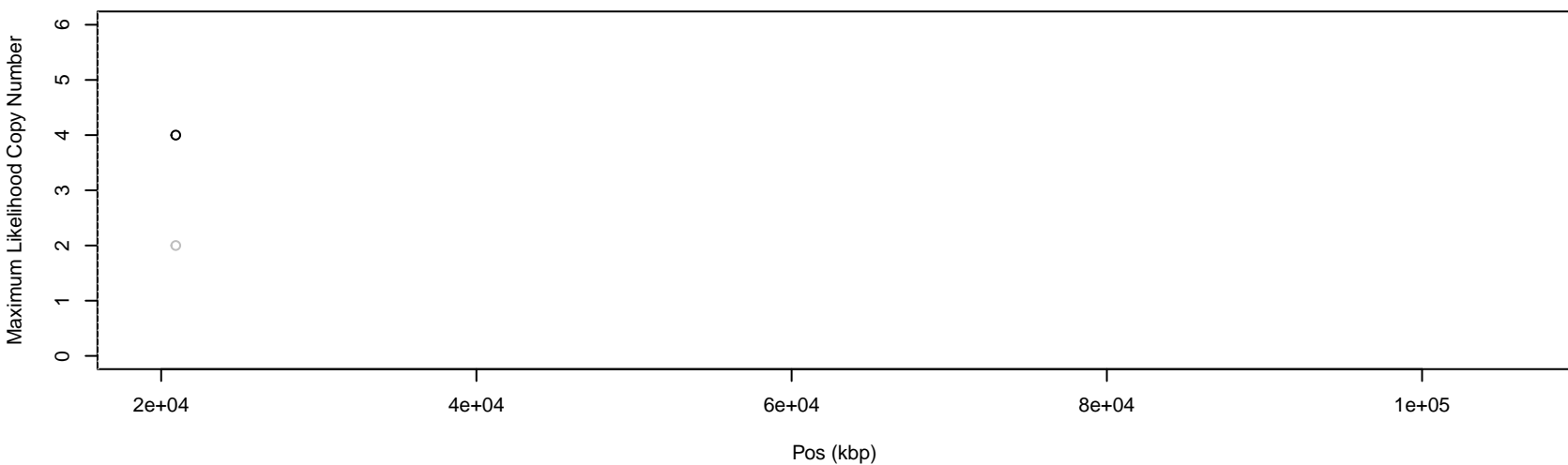
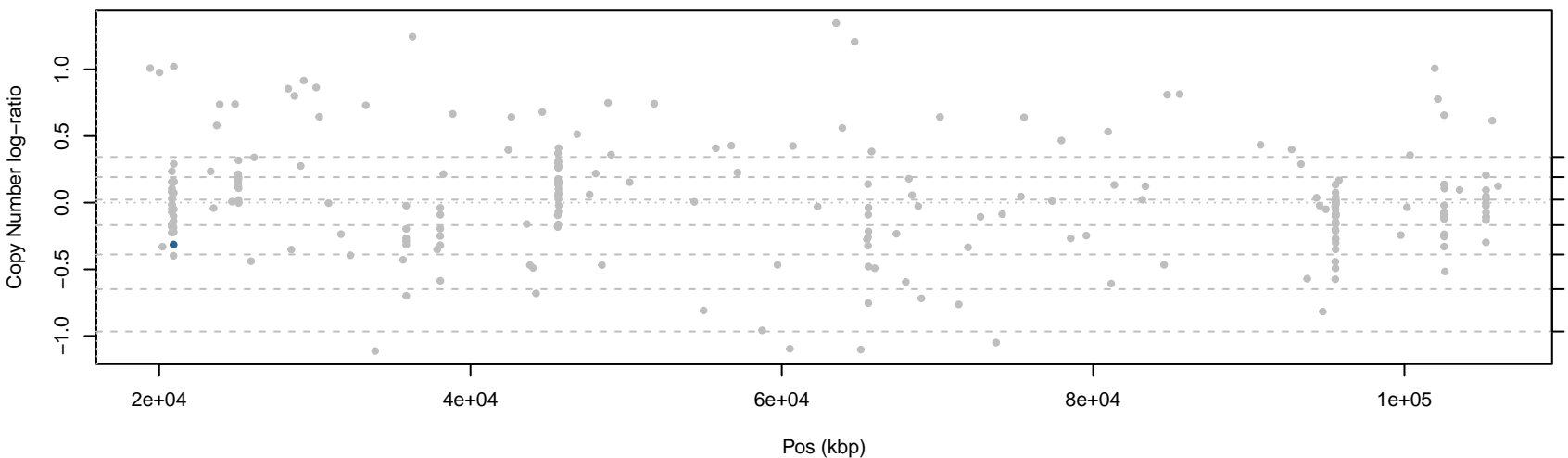
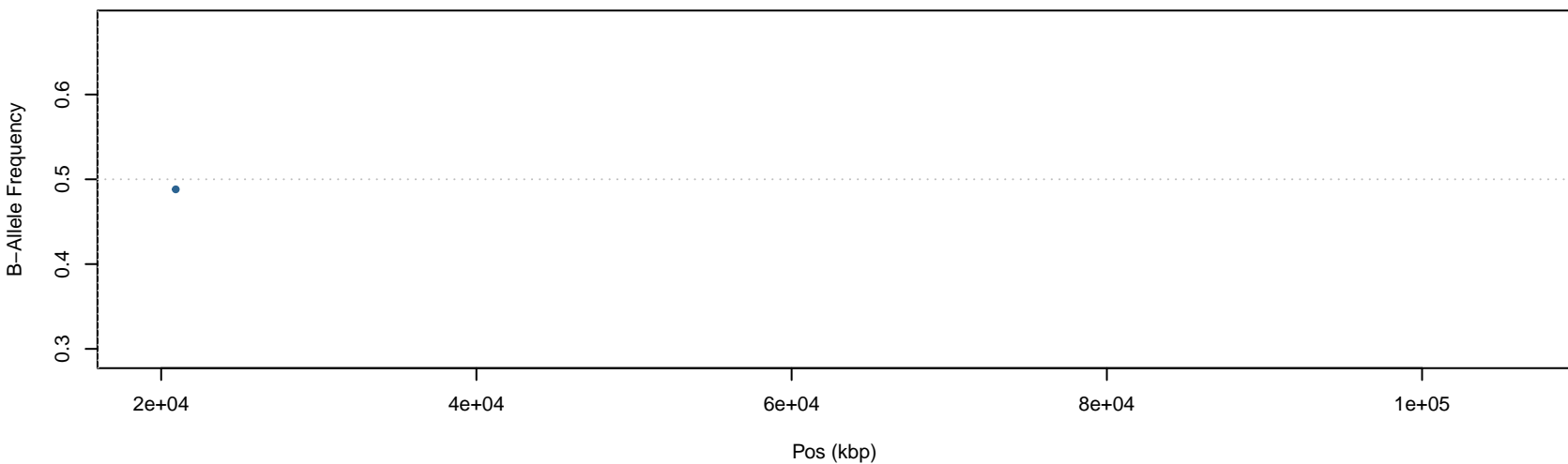
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 12



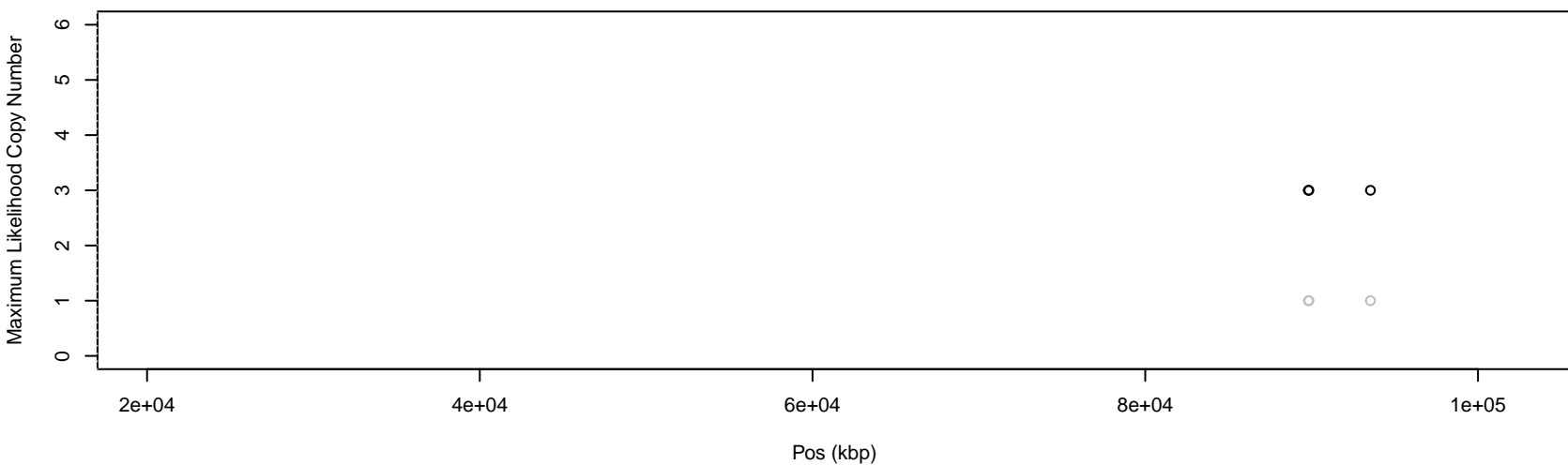
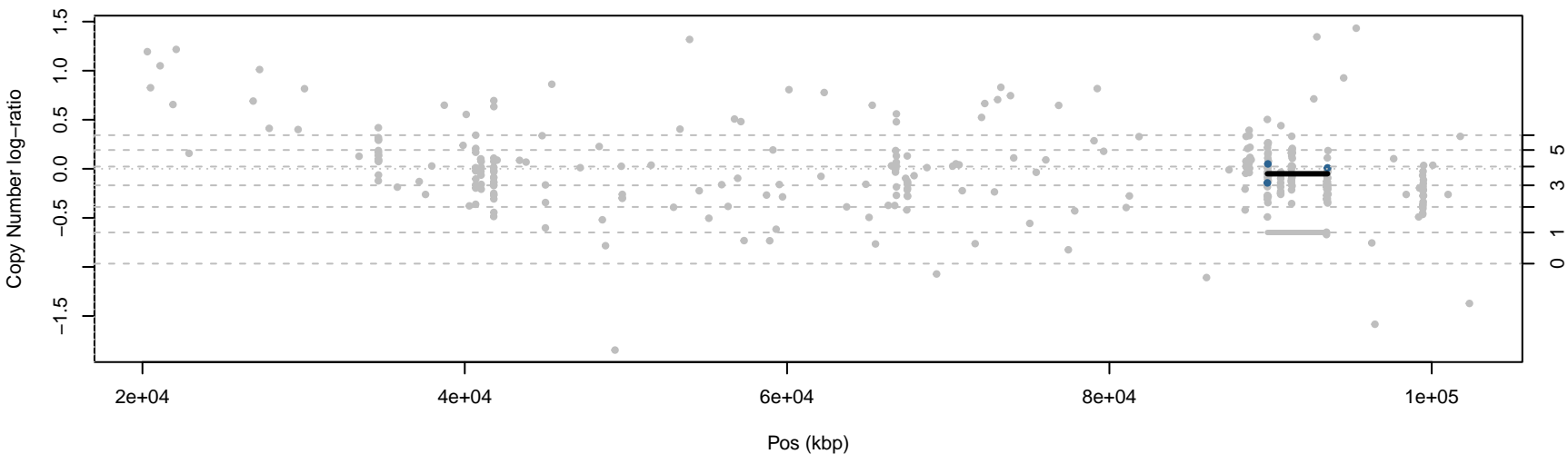
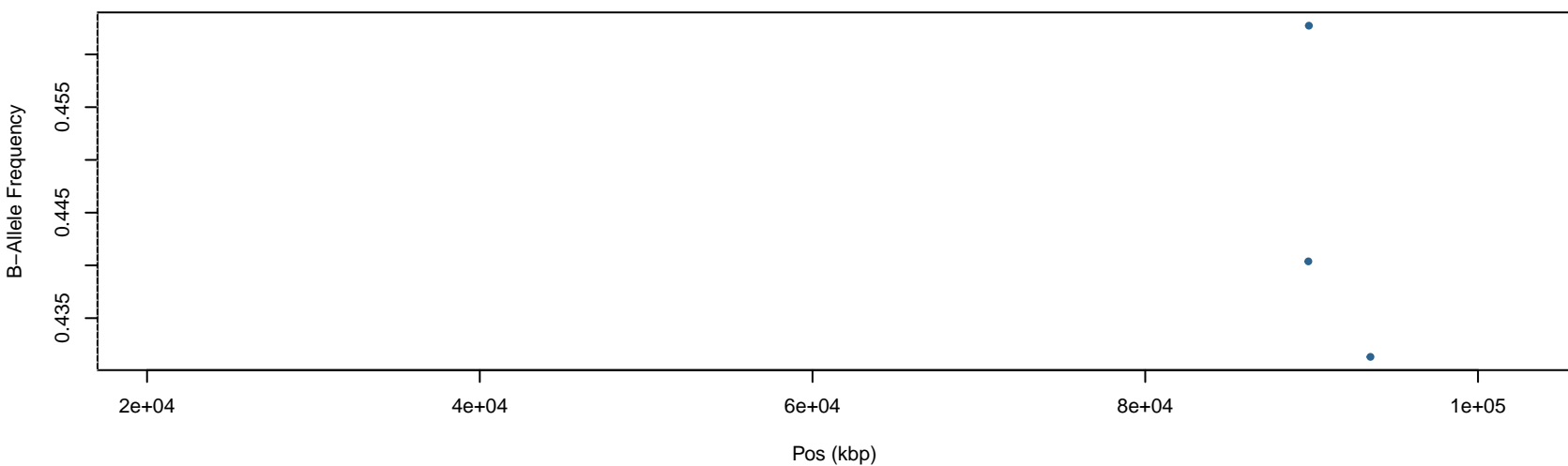
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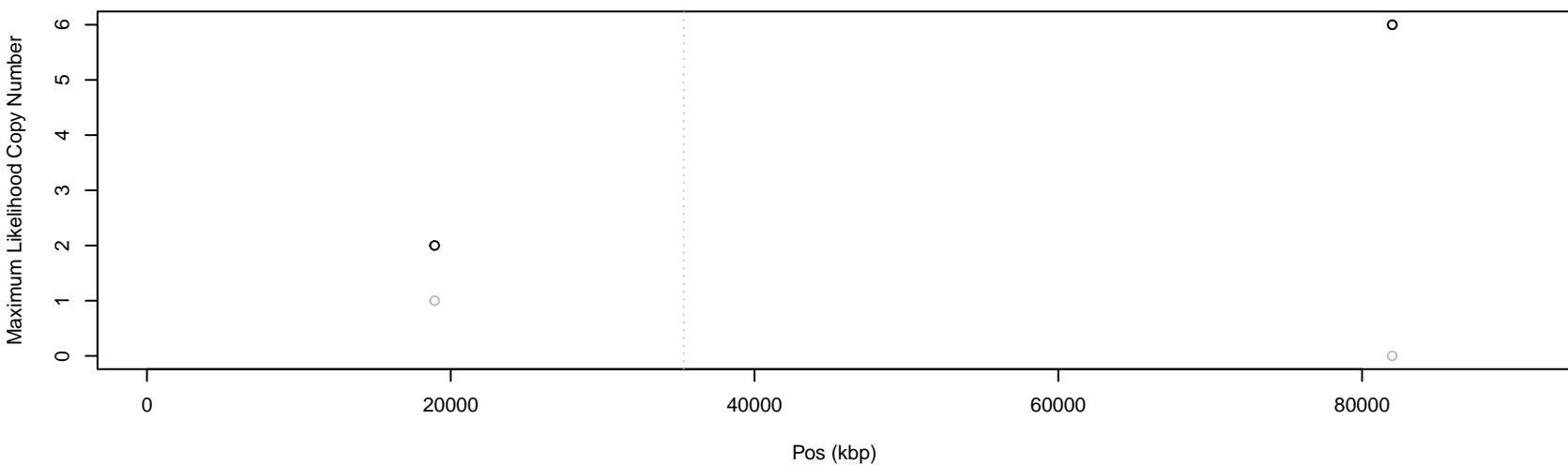
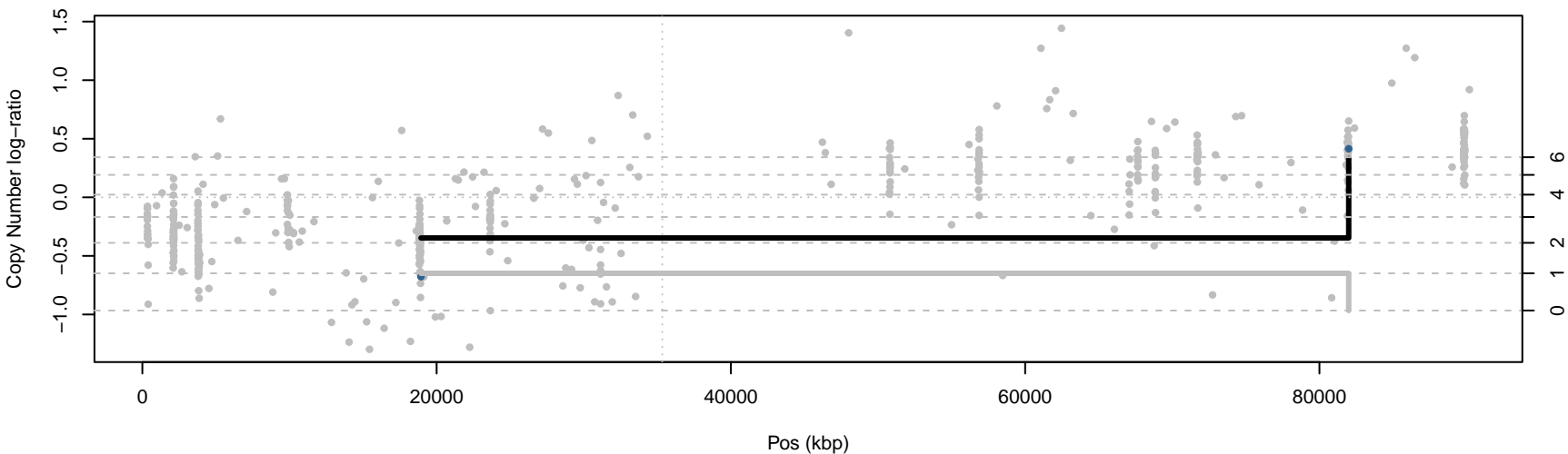
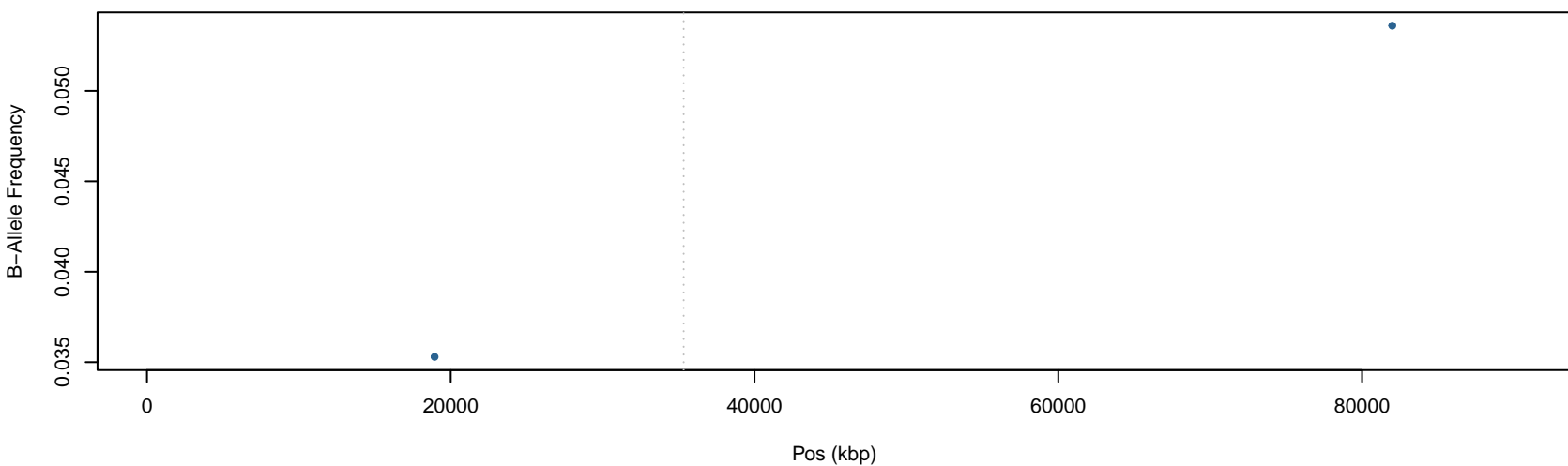
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 14



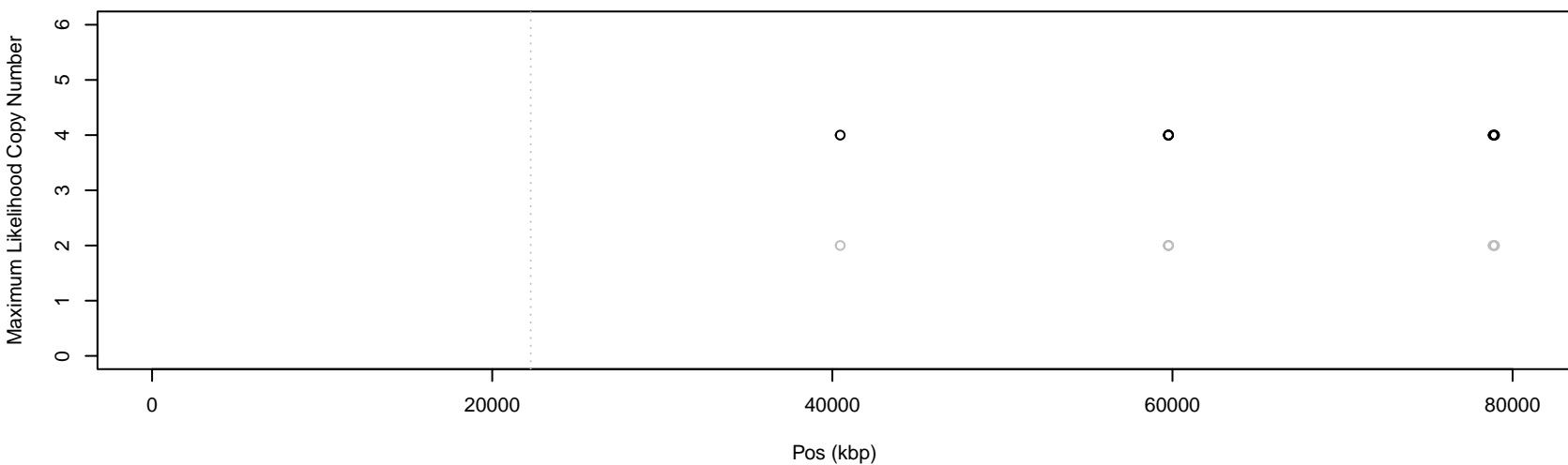
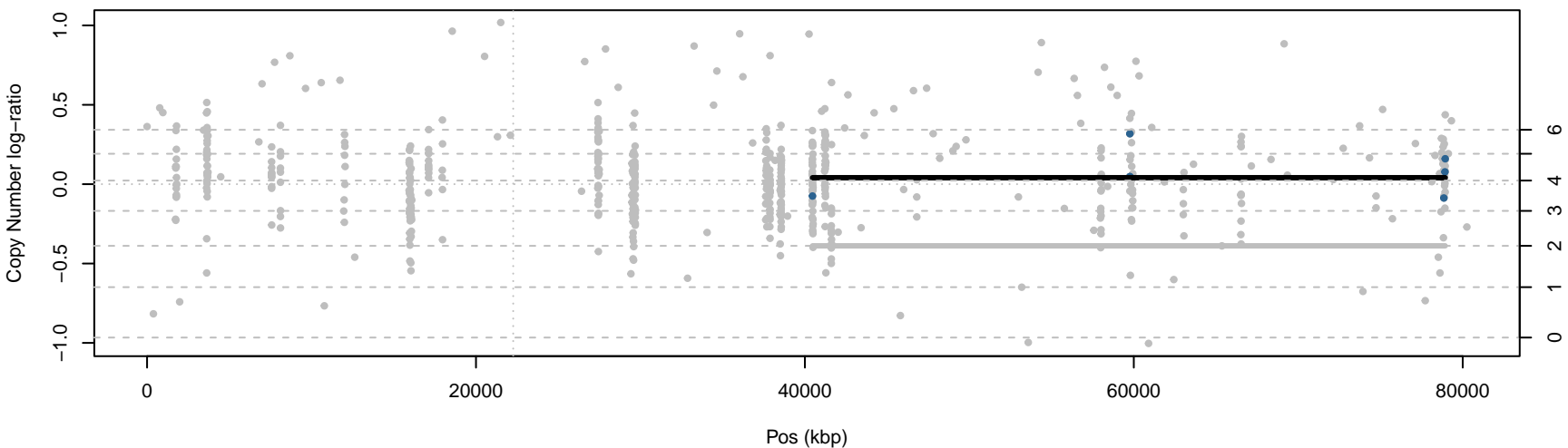
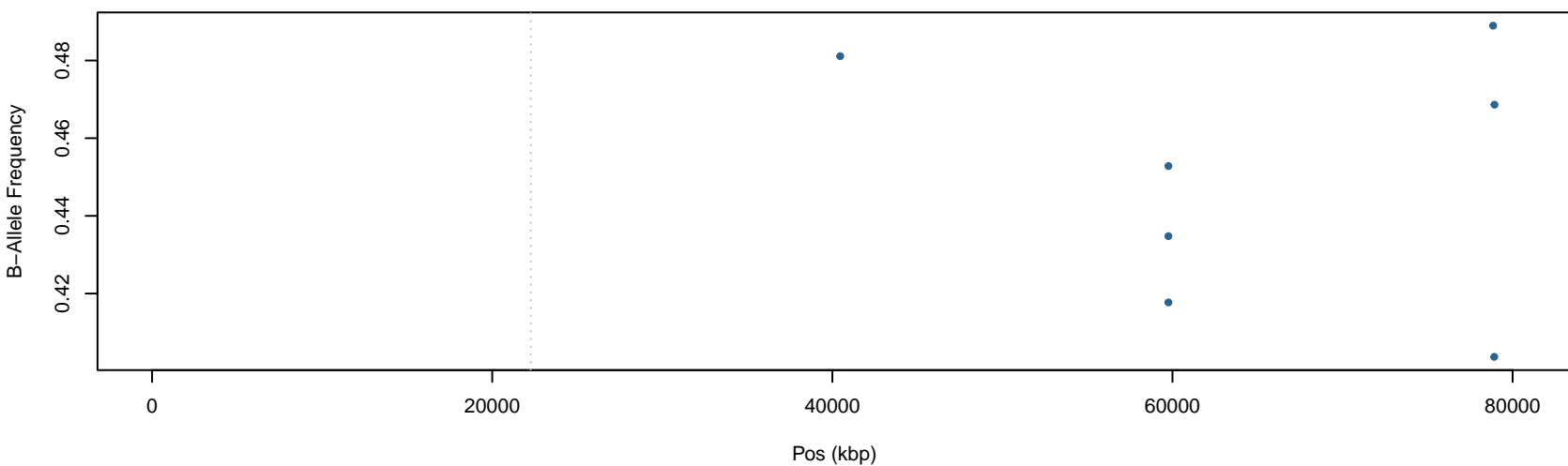
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 15



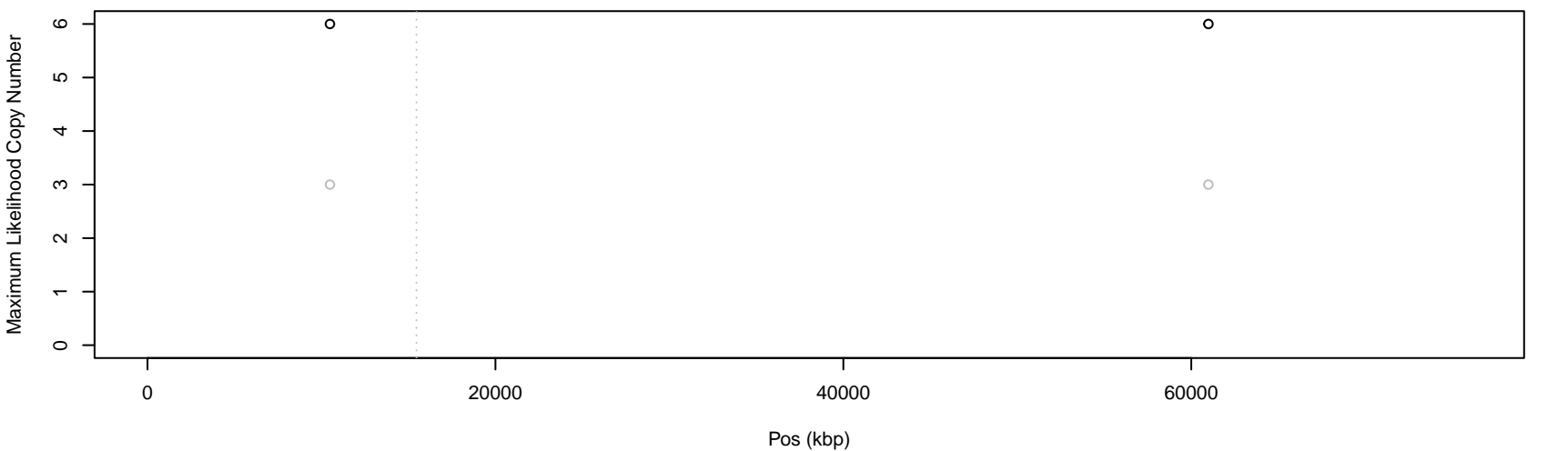
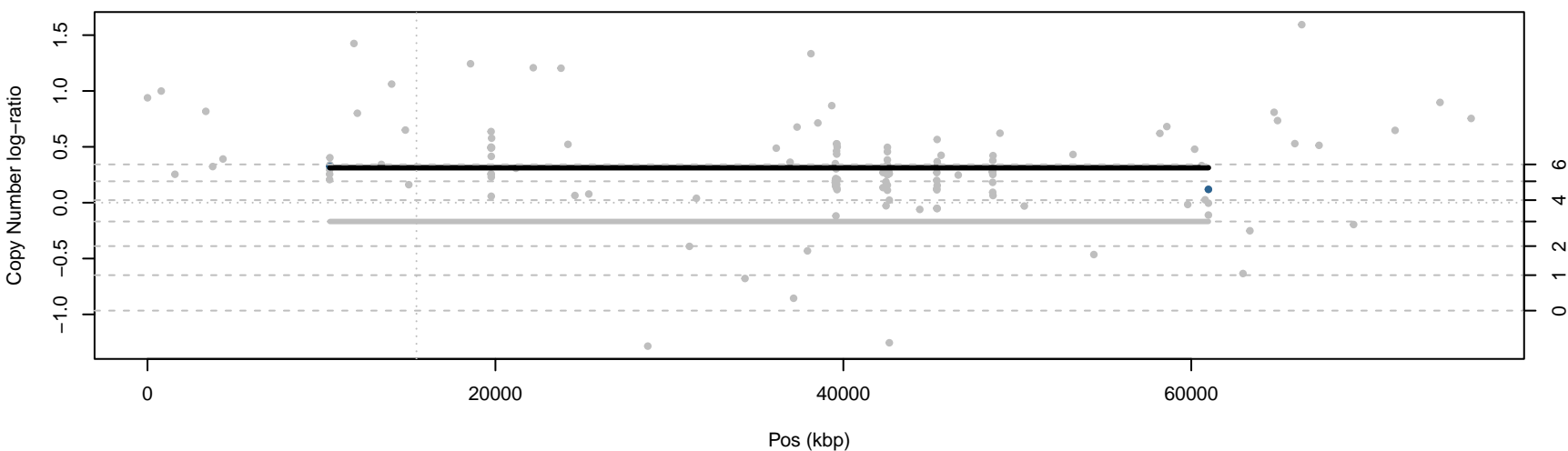
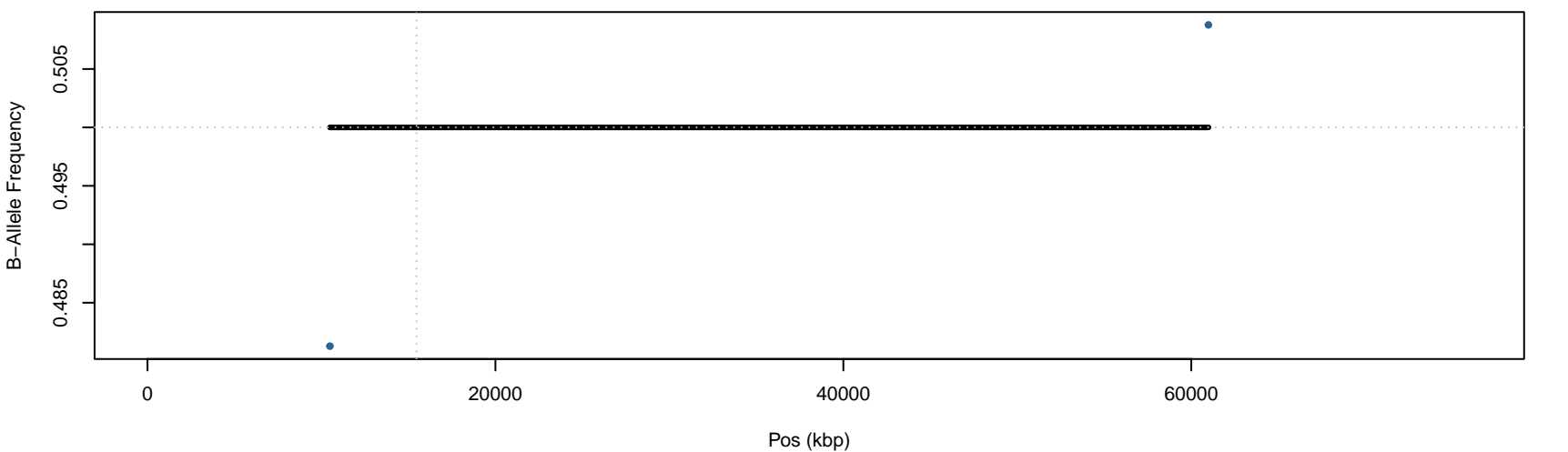
Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 16



Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 17

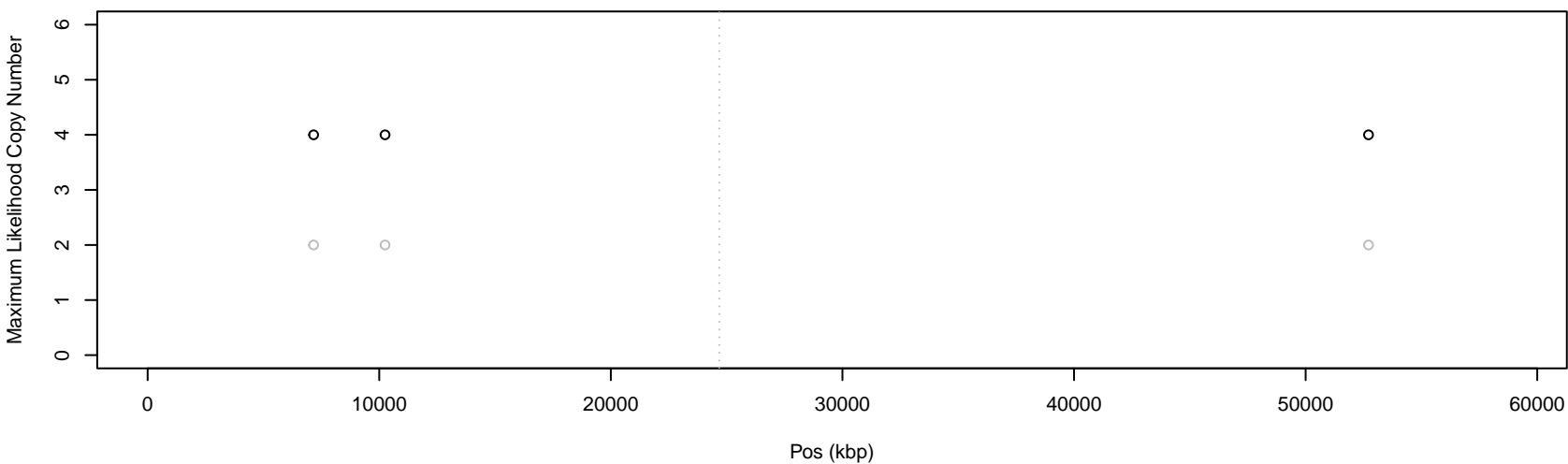
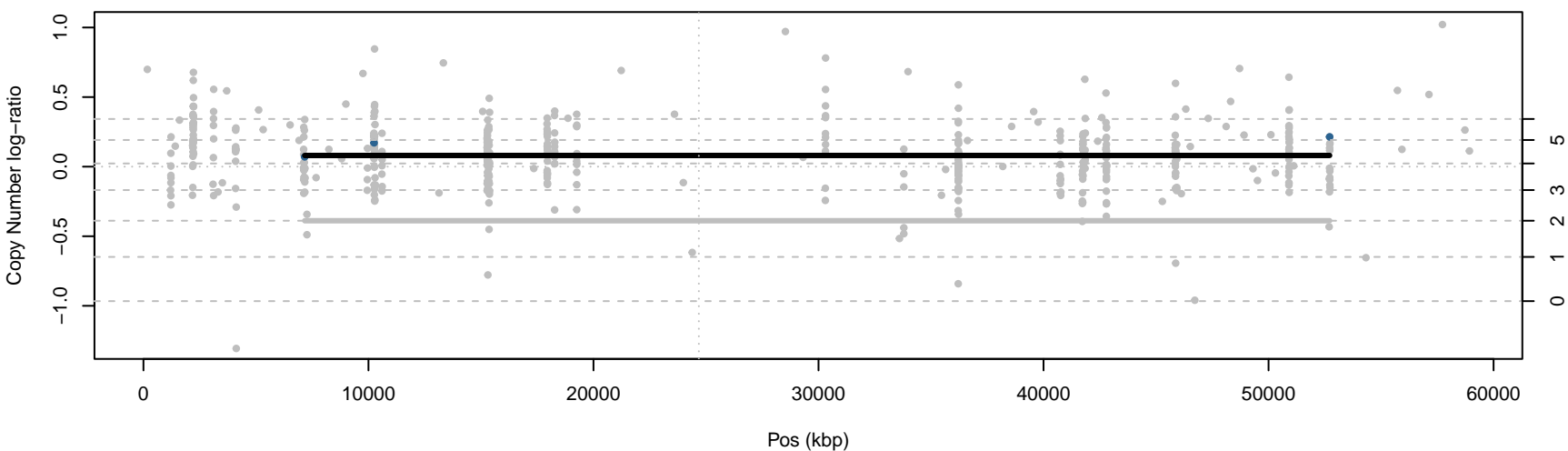
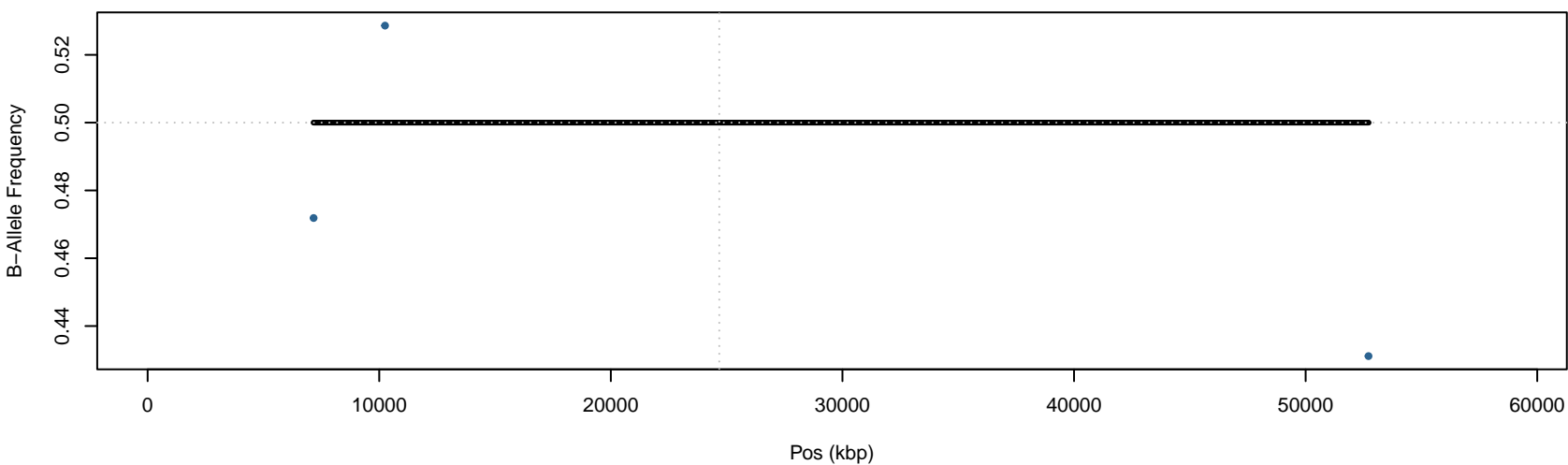


Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 18

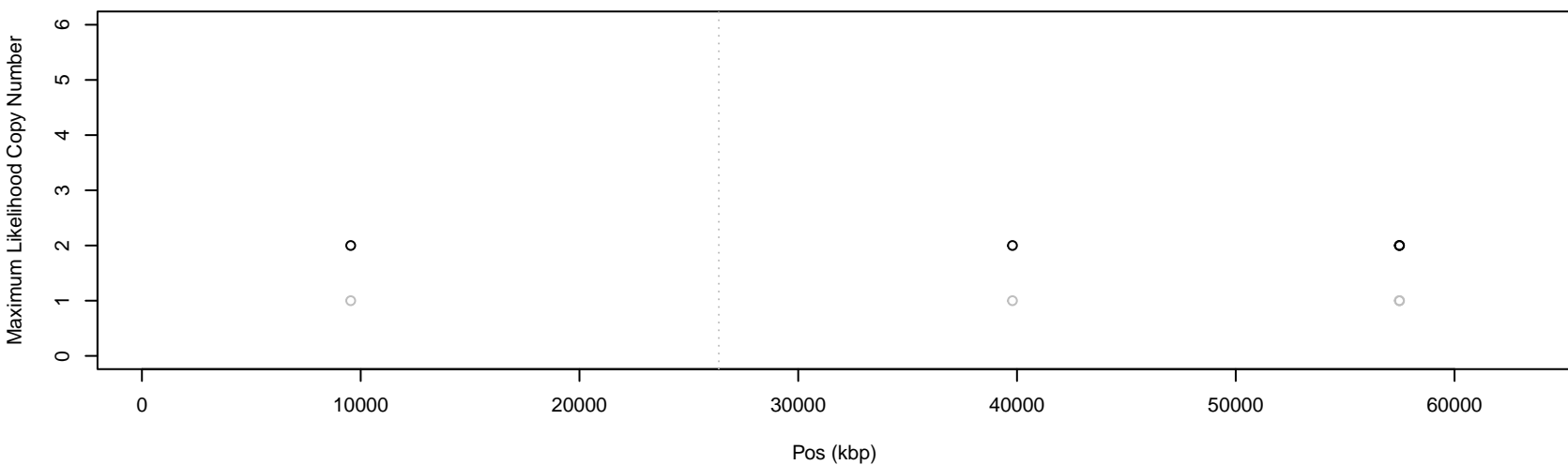
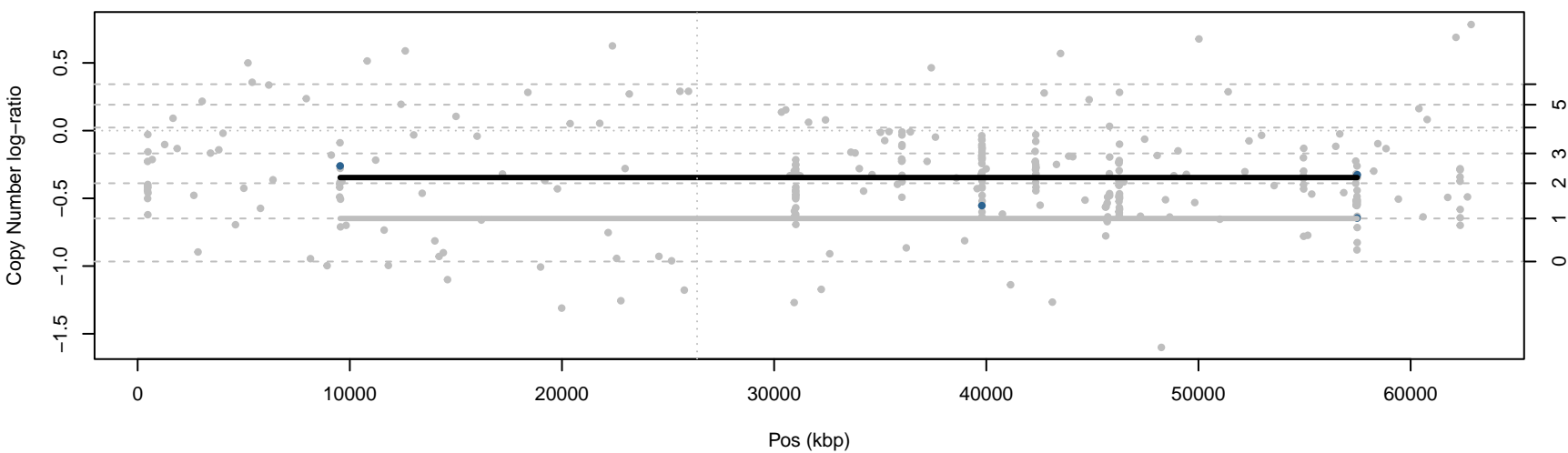
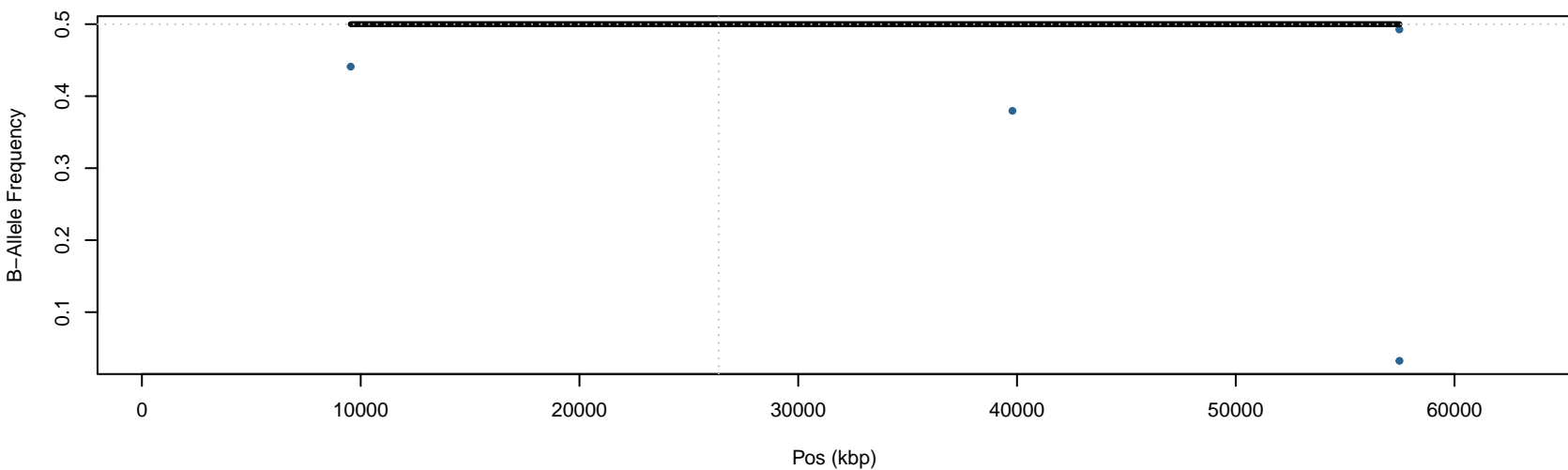




Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 19



Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 20



Purity: 0.33 Tumor ploidy: 3.874 SNV log-likelihood: -29.33 GoF: 91.1% Mean coverage: 454;525 Chromosome: 22

