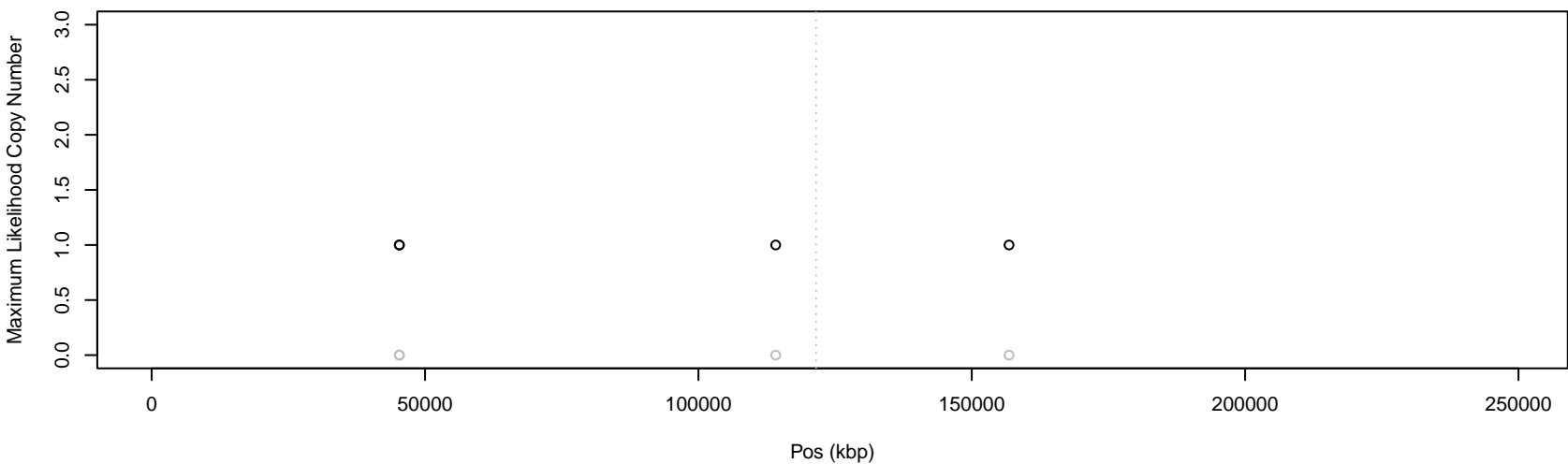
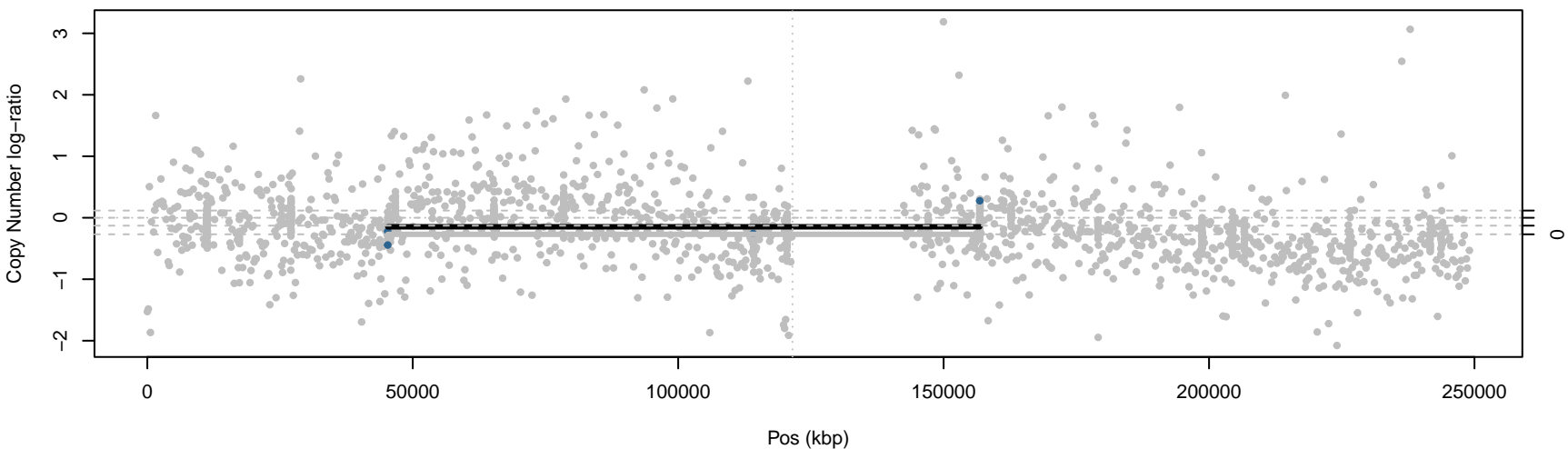
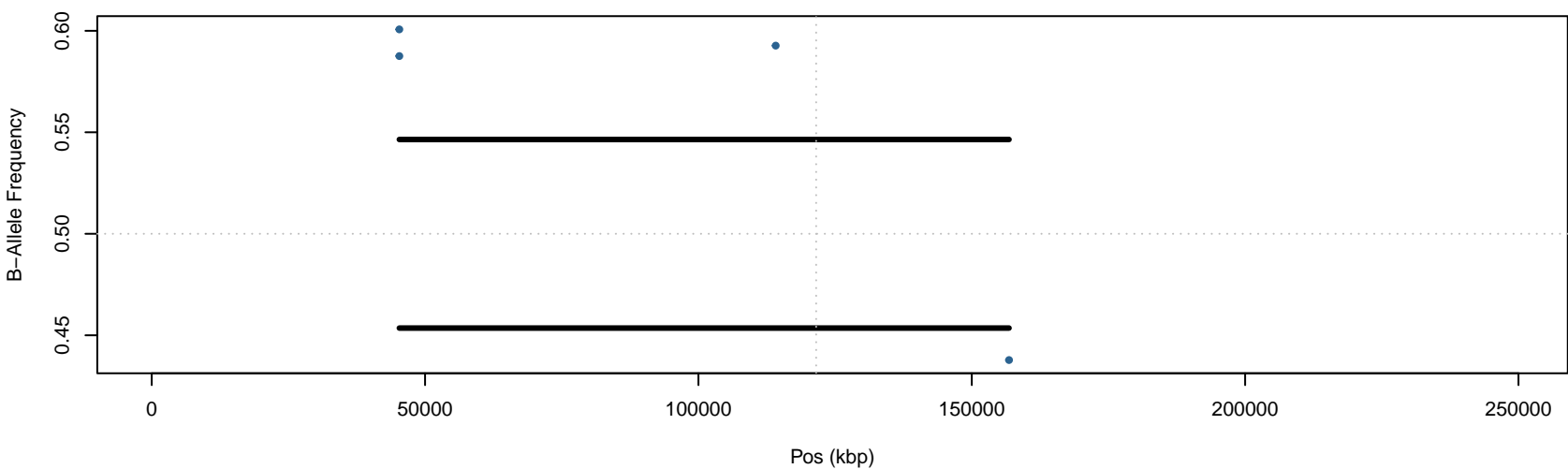
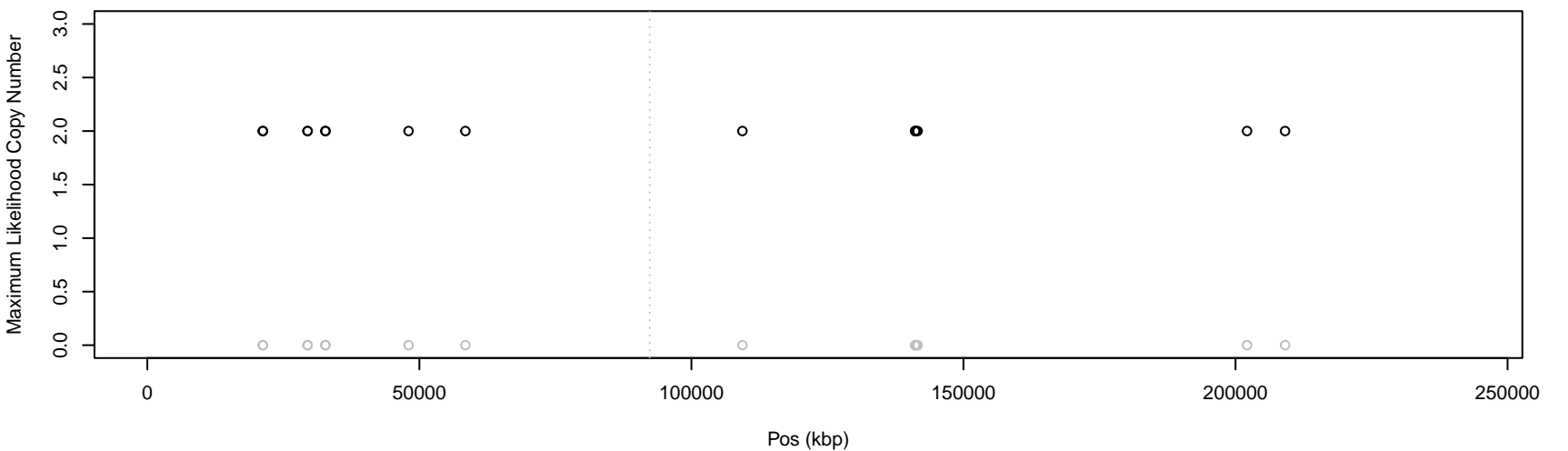
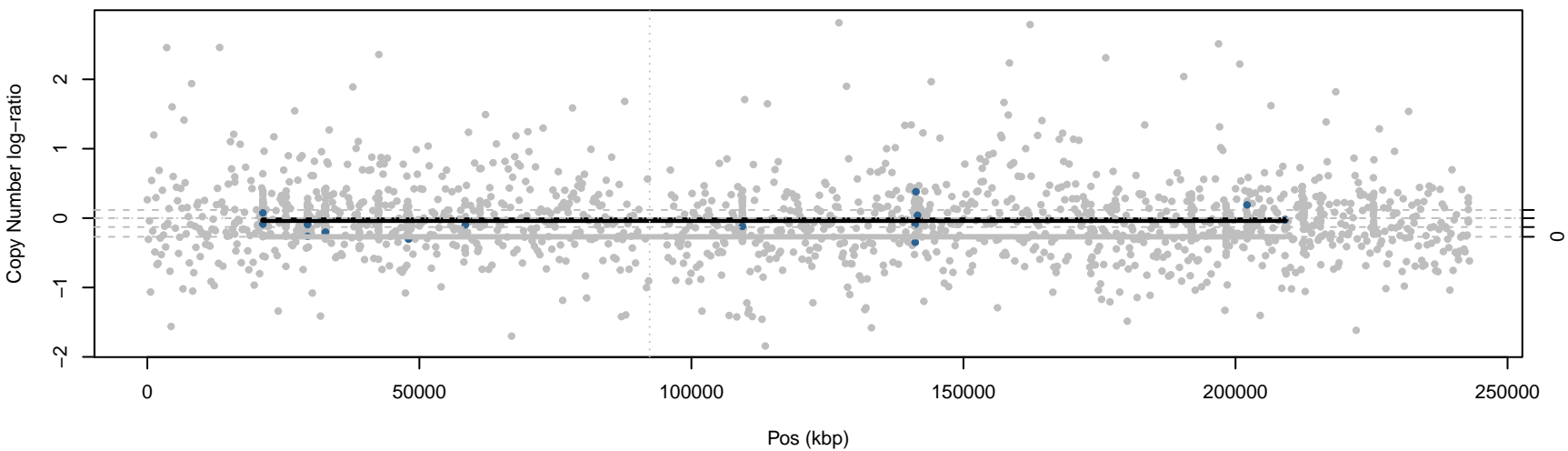
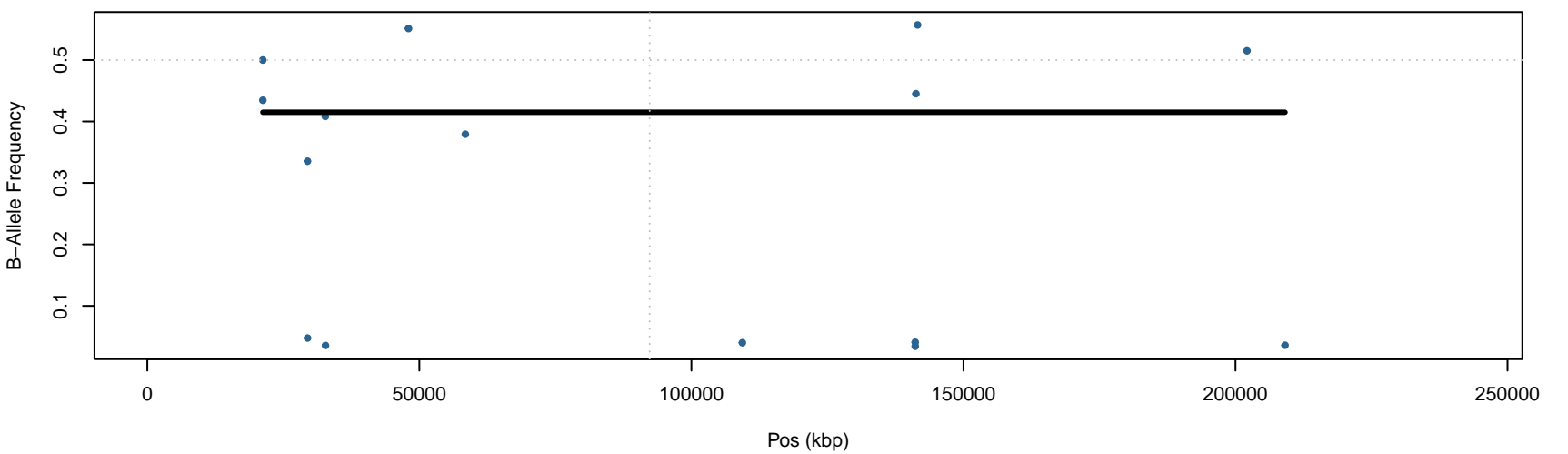


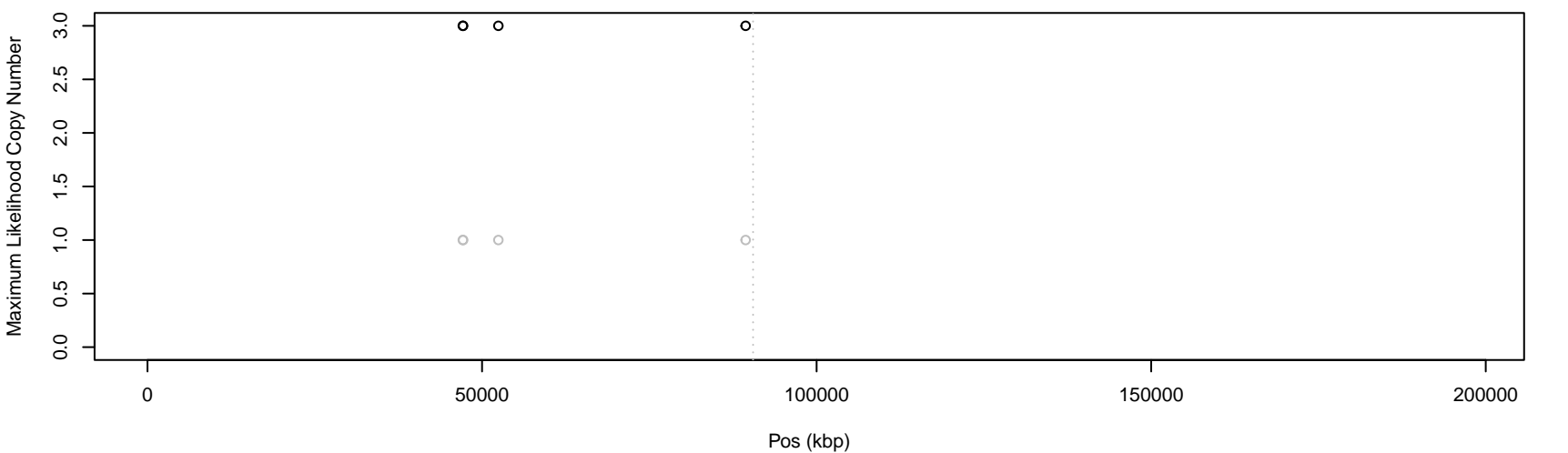
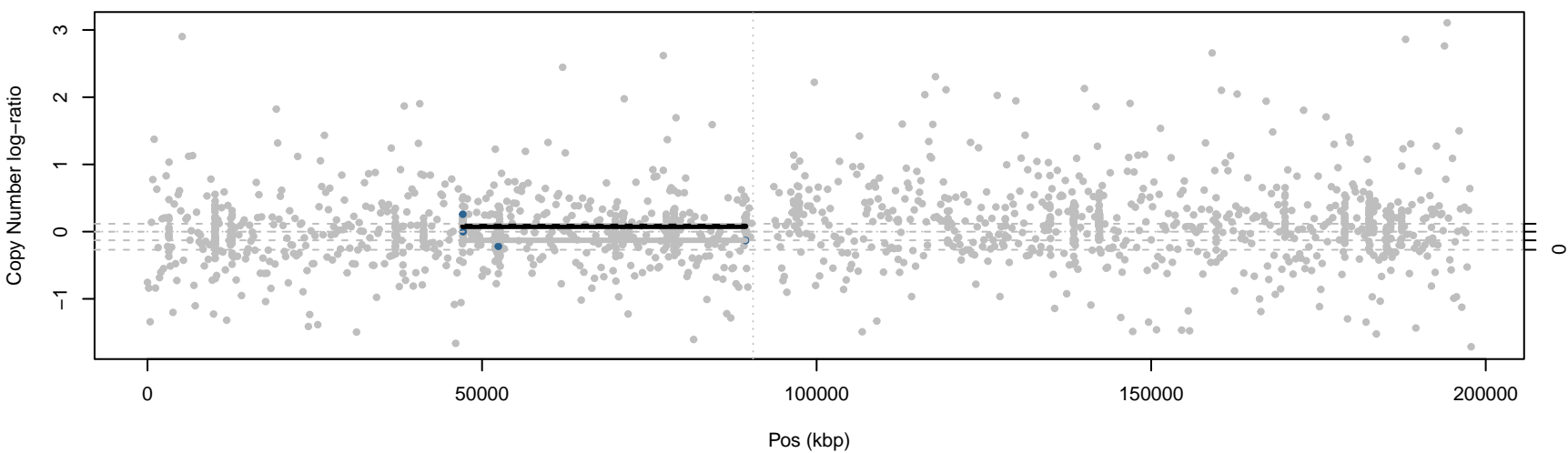
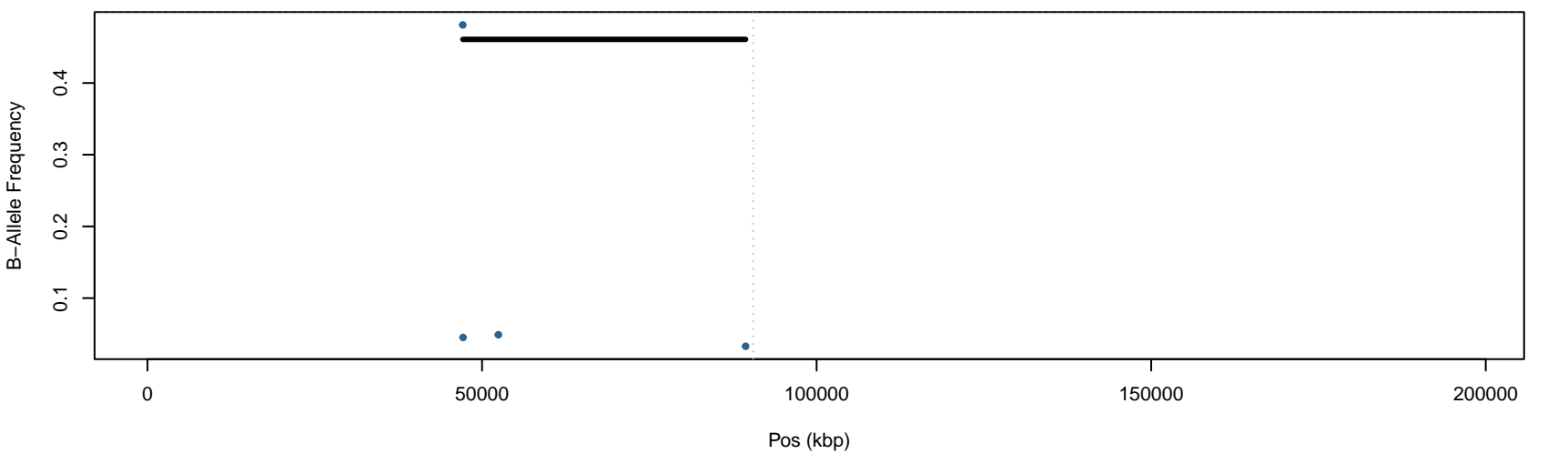
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 1



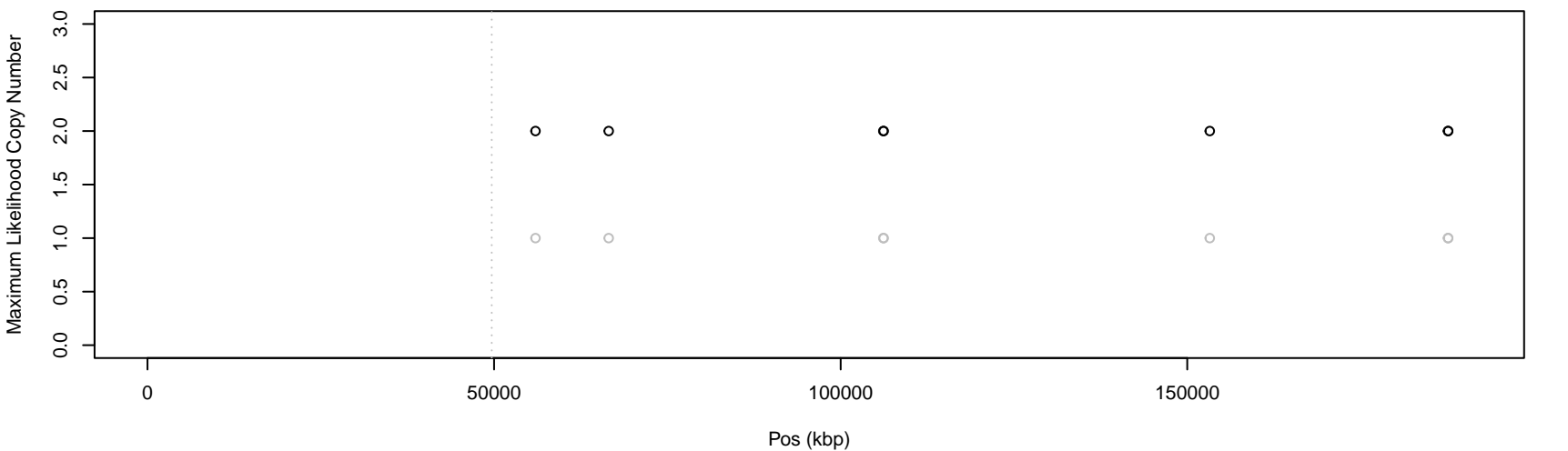
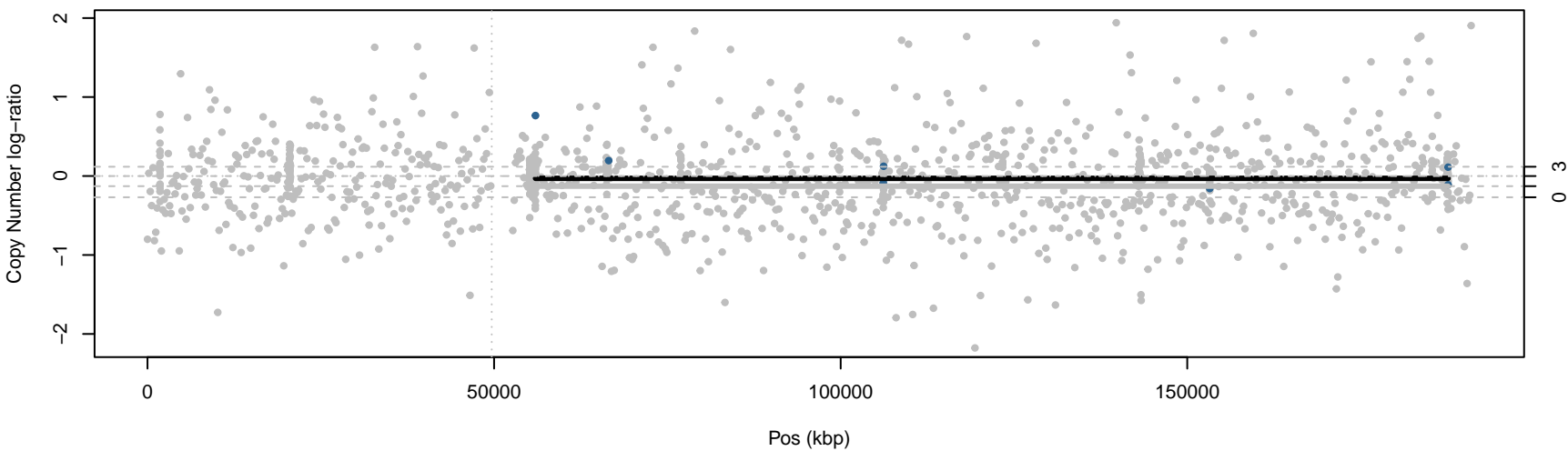
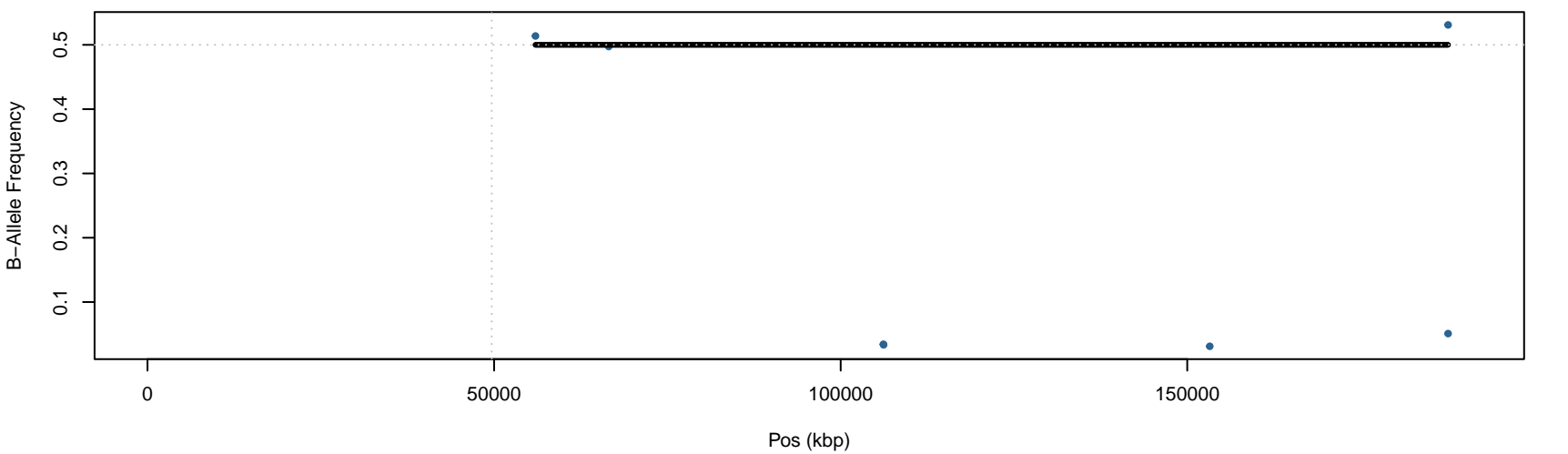
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 2



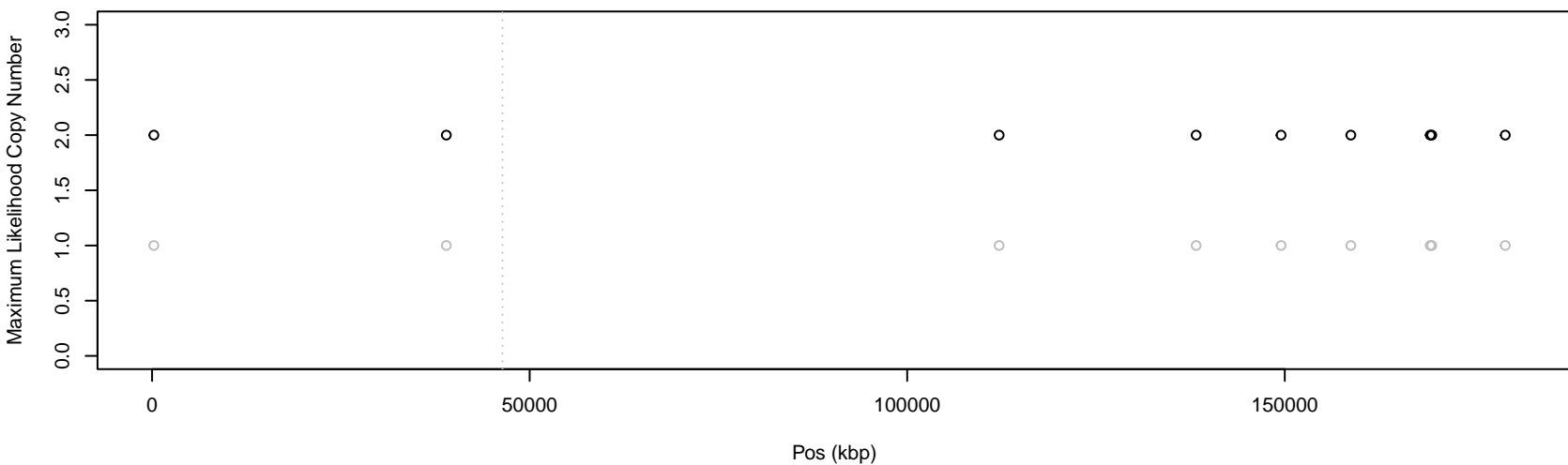
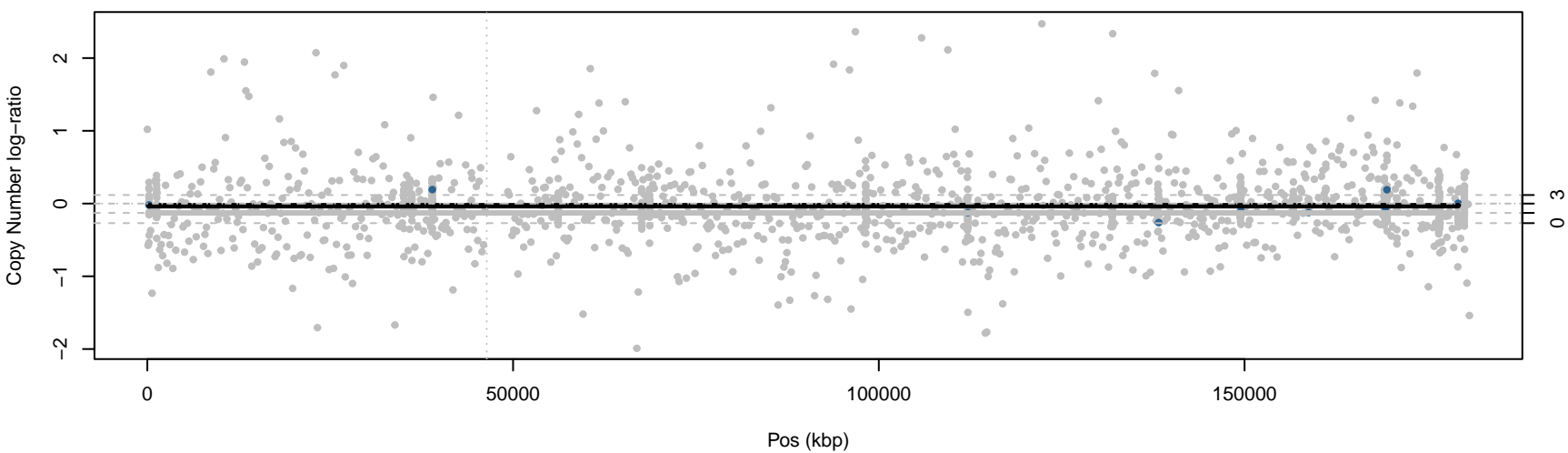
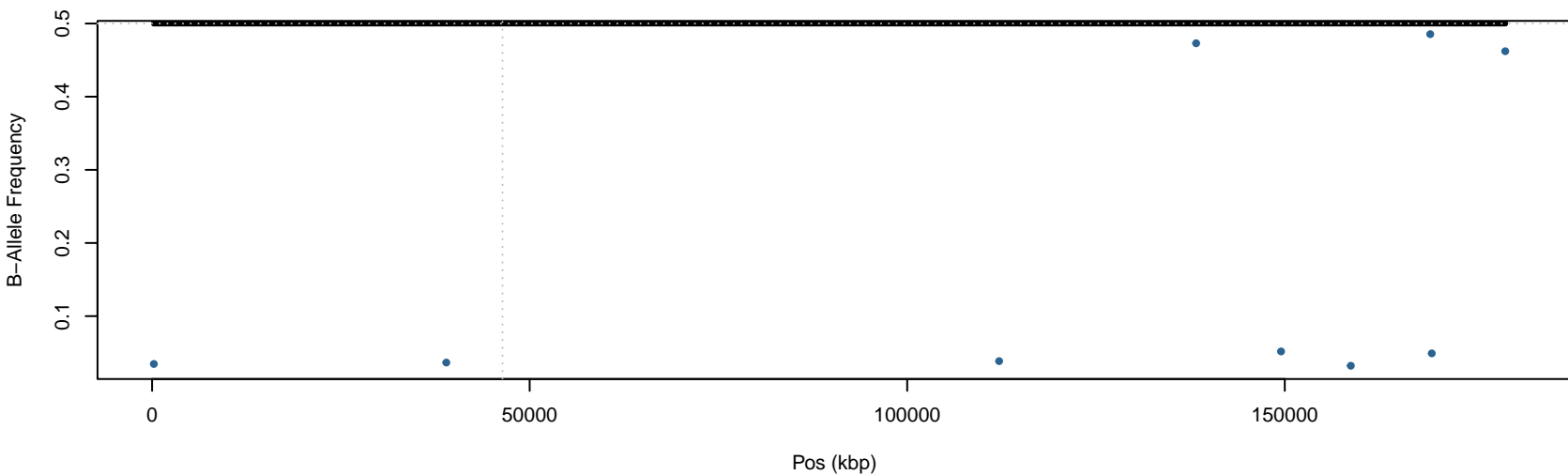
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 3



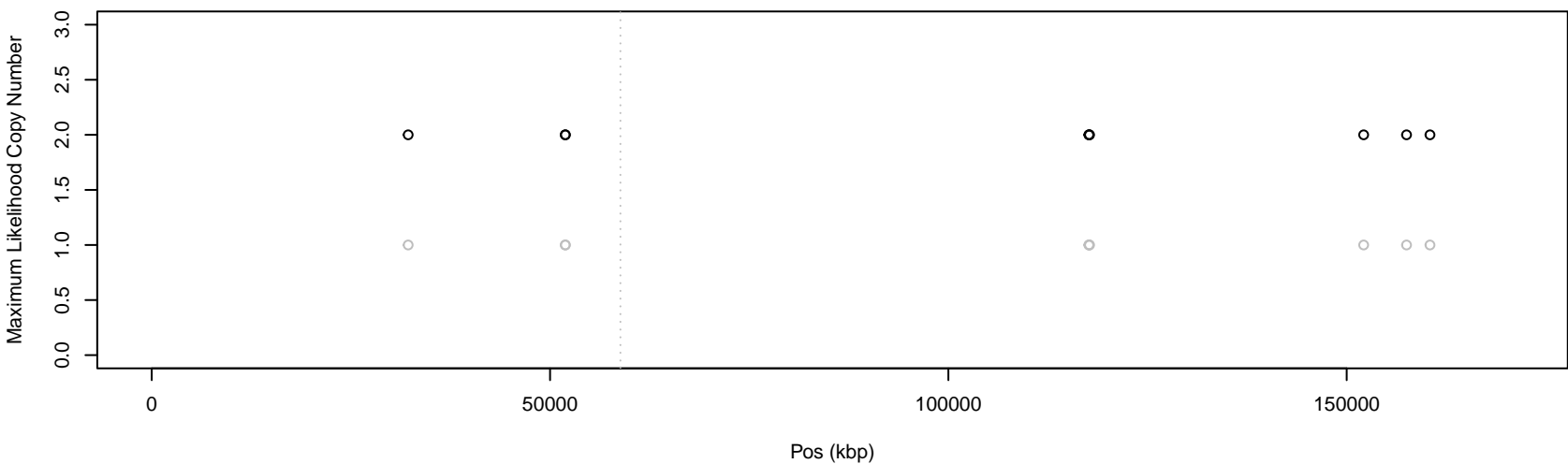
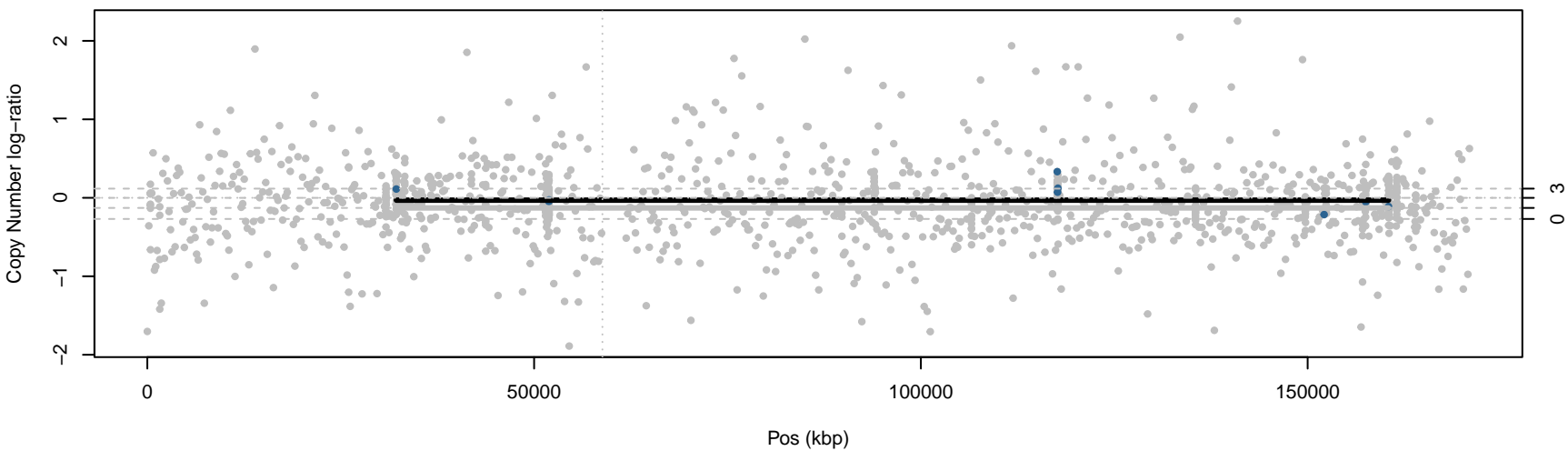
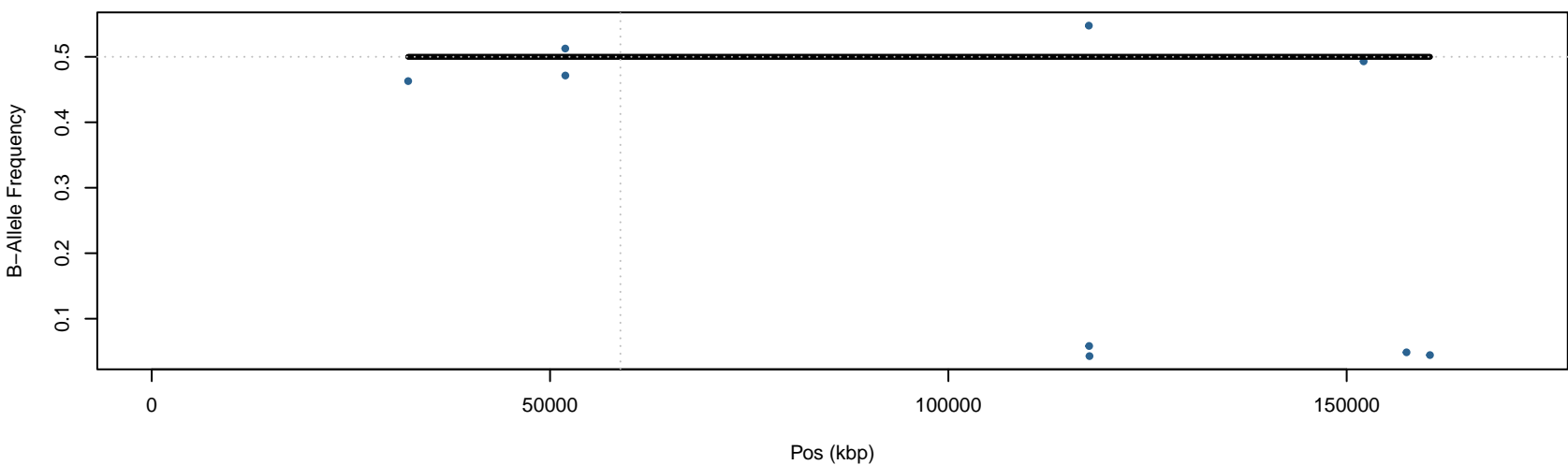
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 4



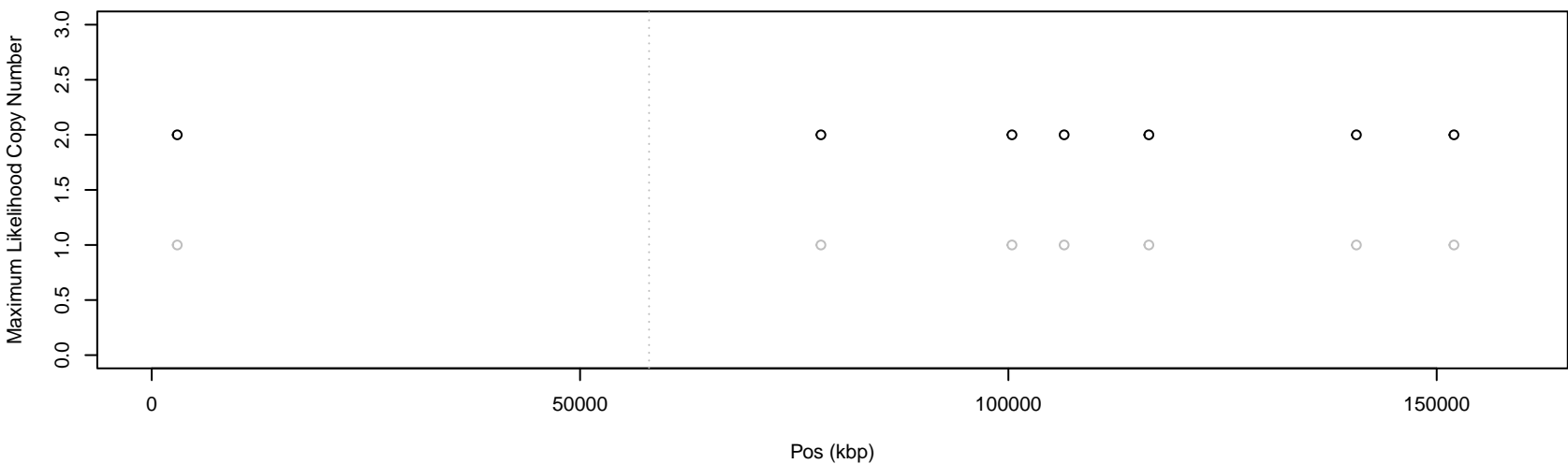
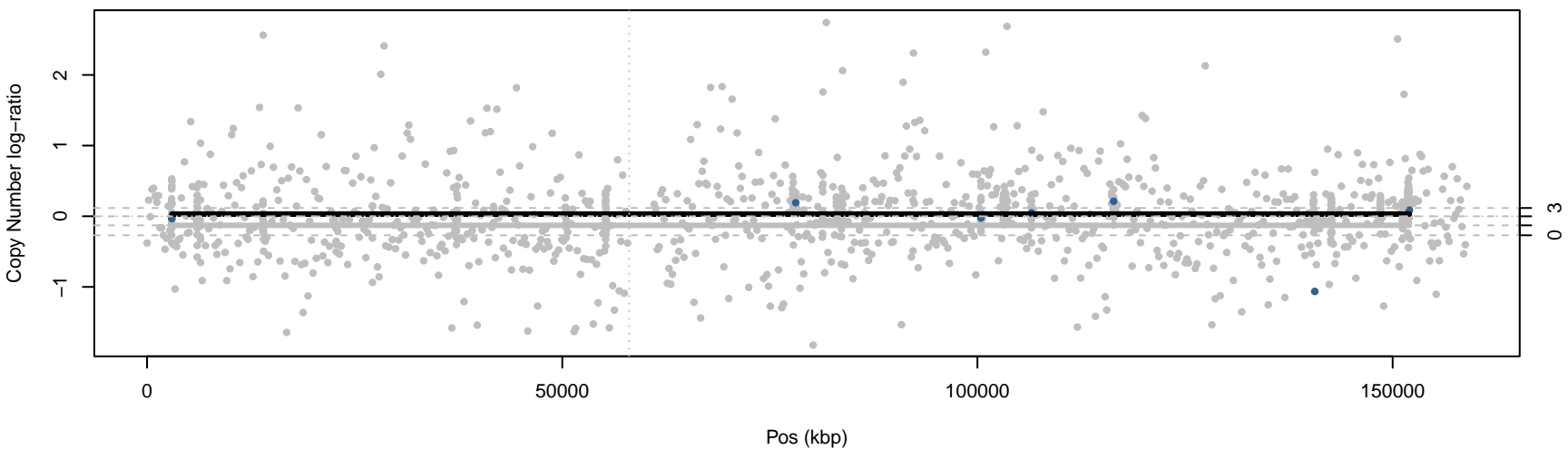
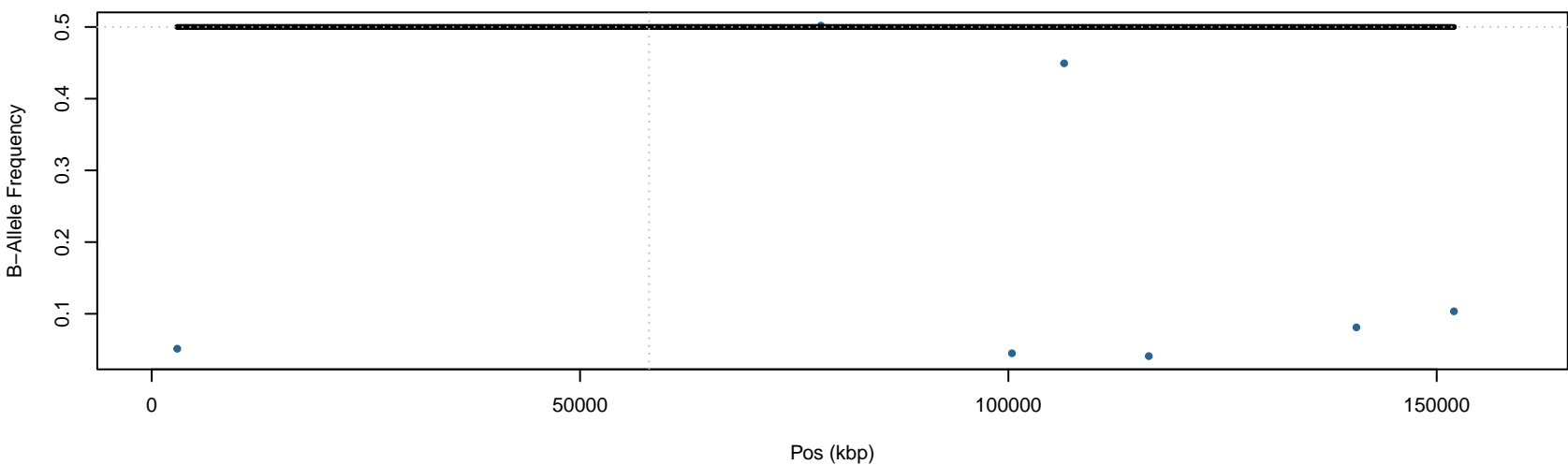
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 5



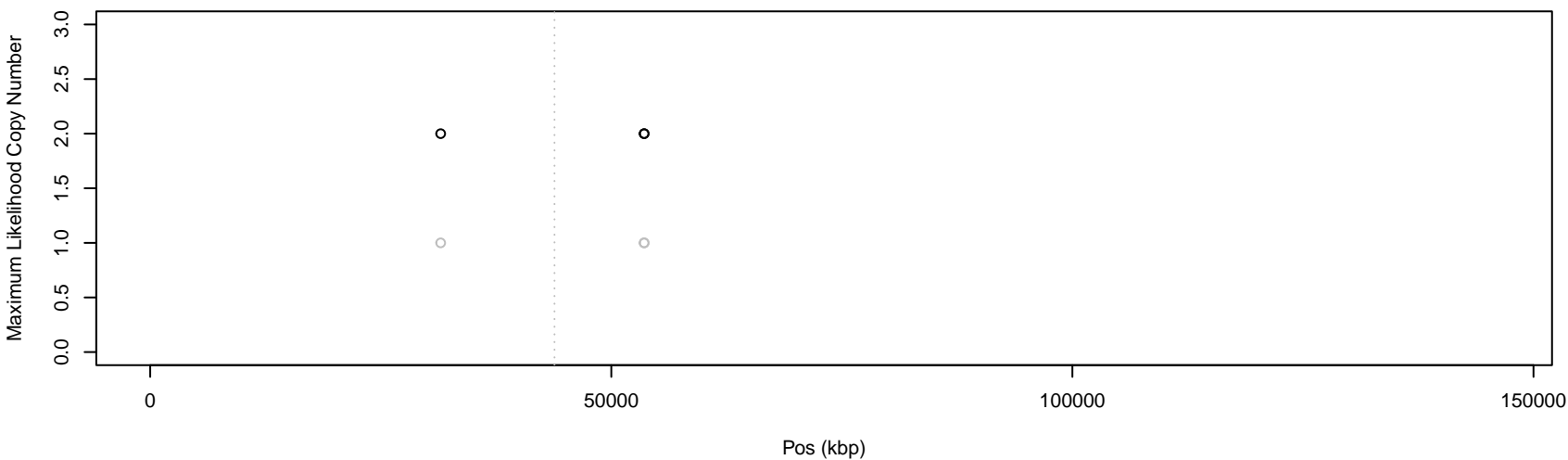
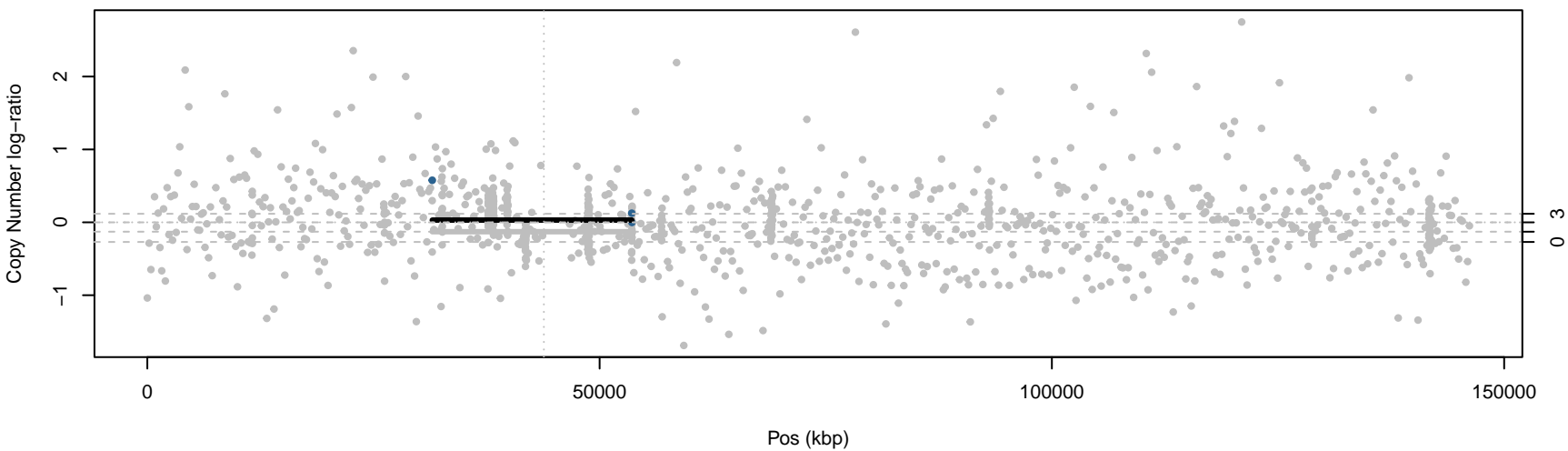
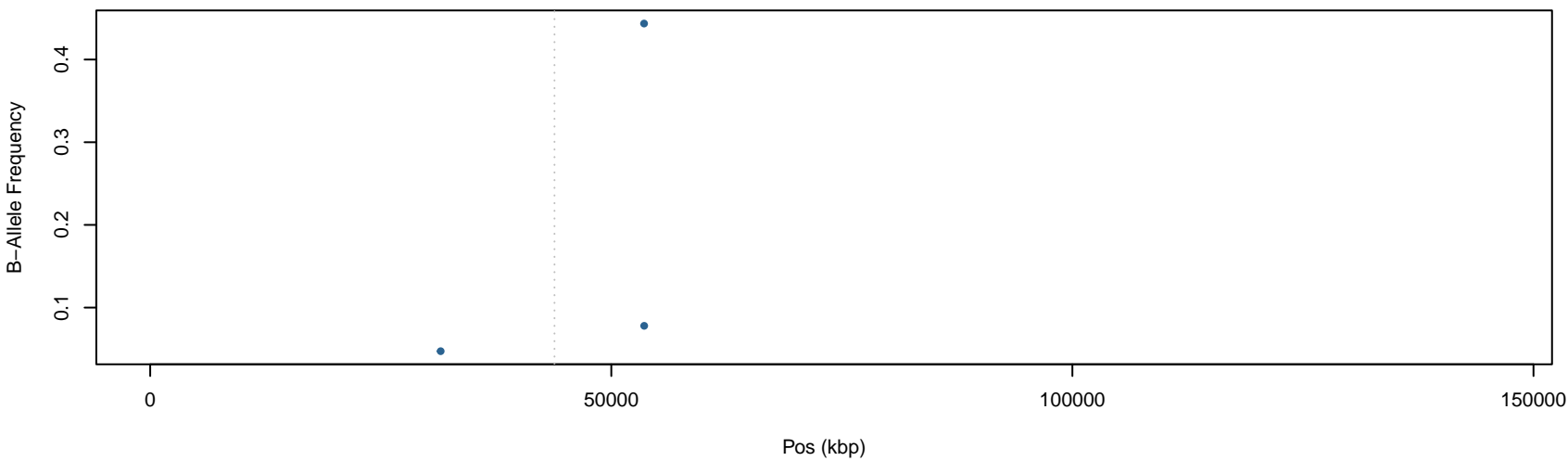
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 6



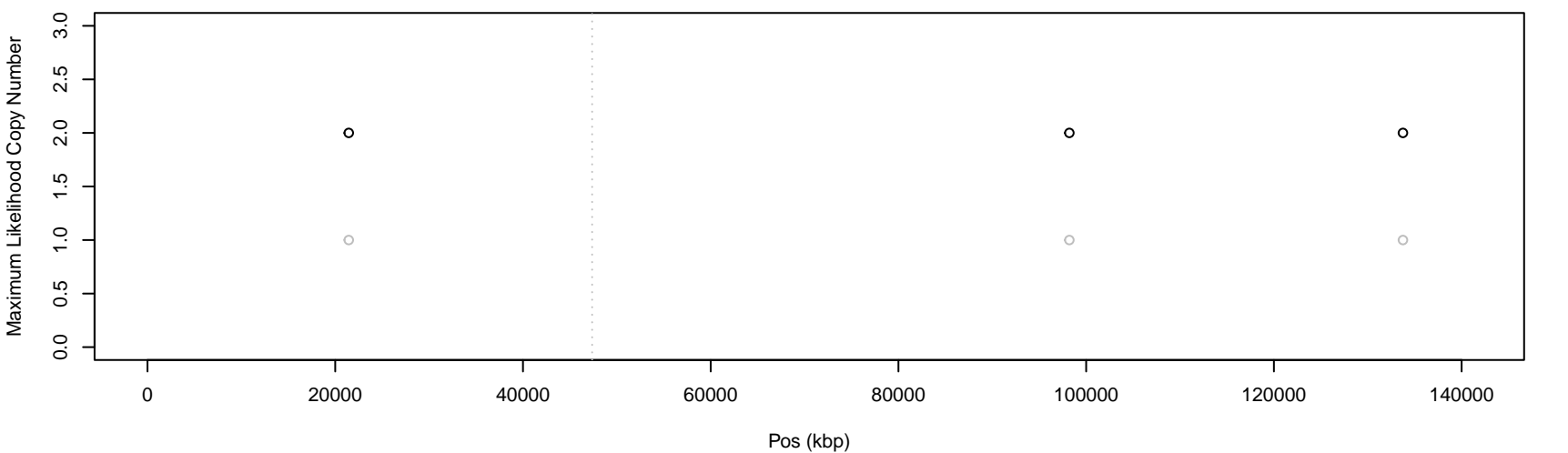
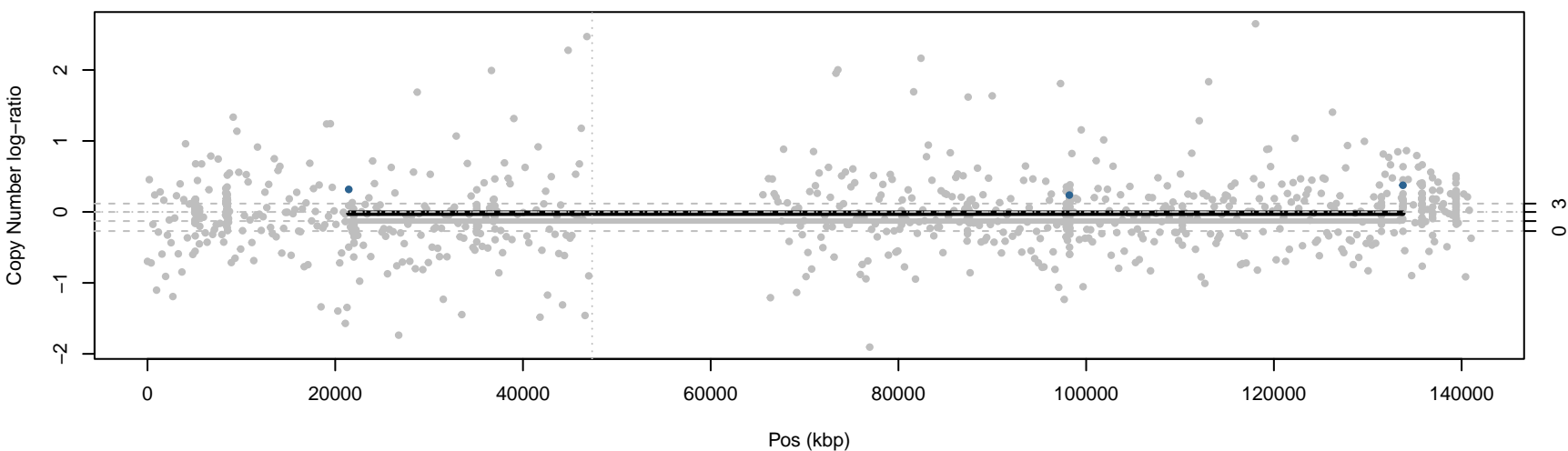
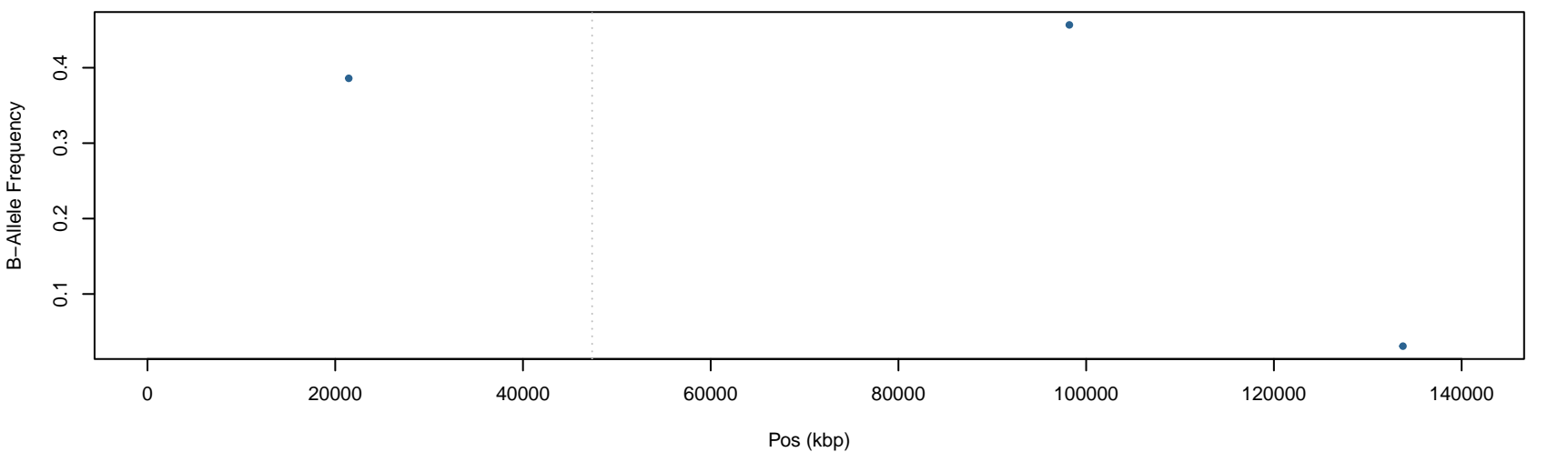
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 7



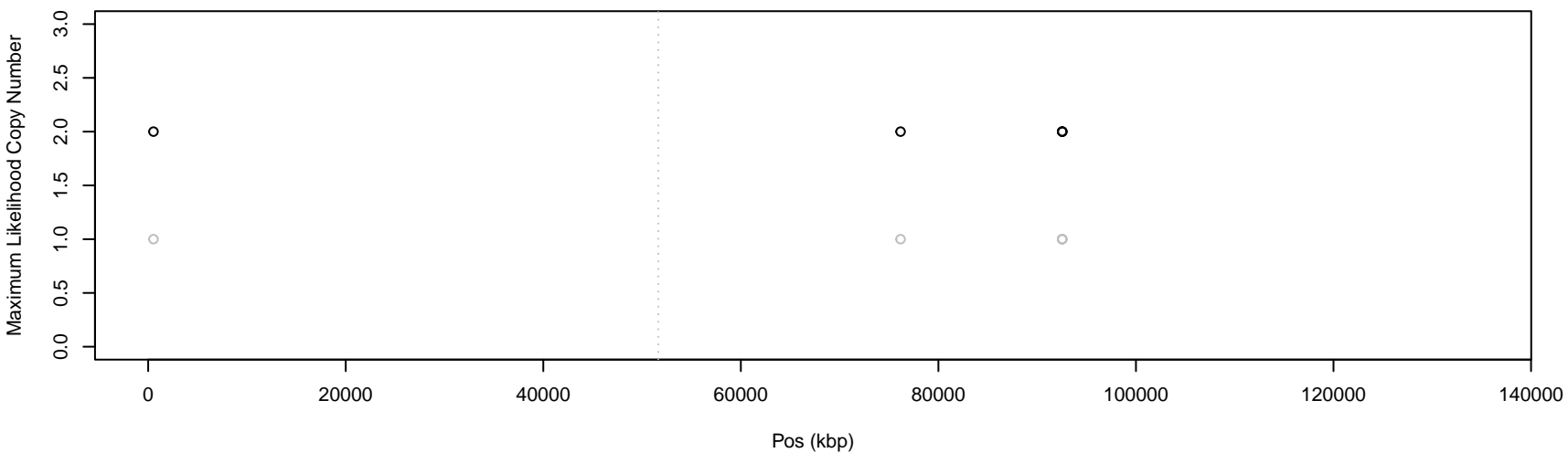
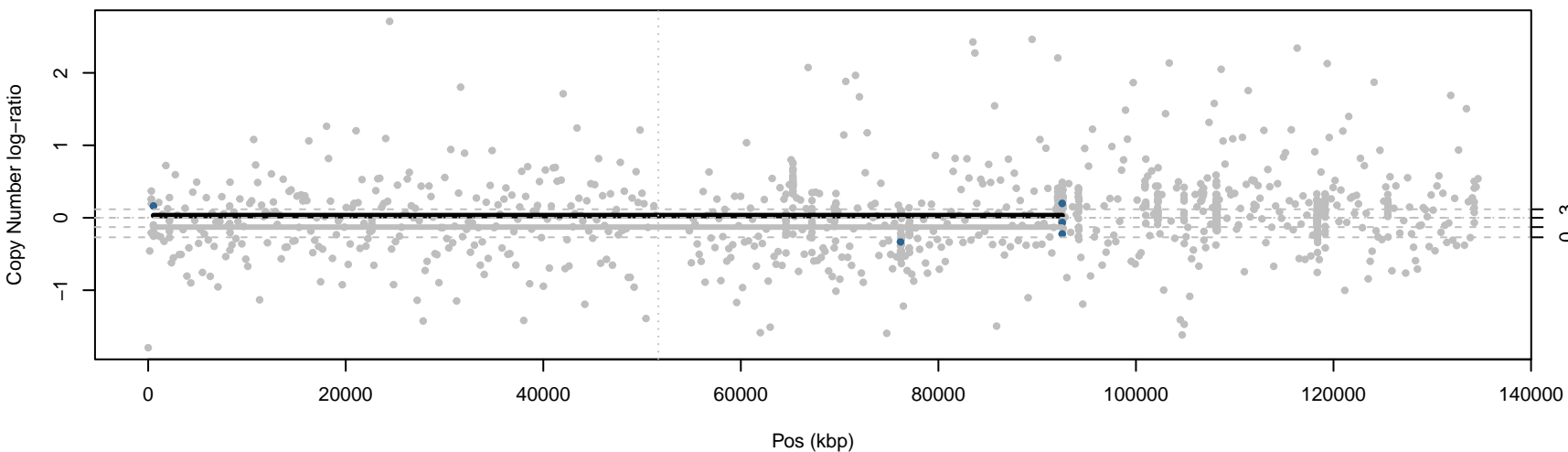
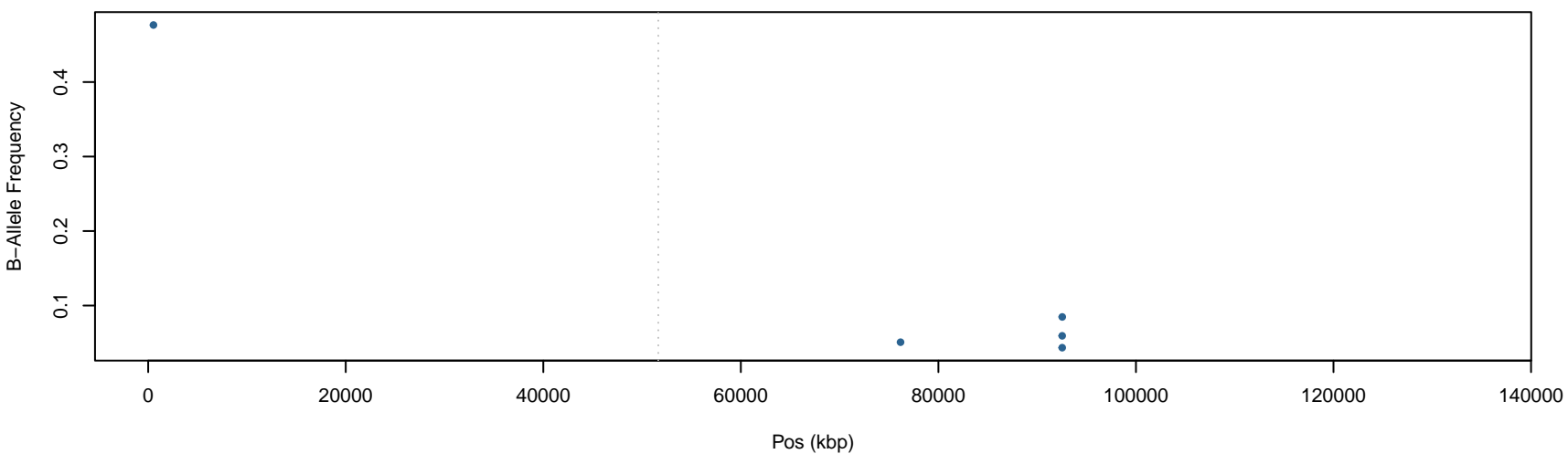
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 8



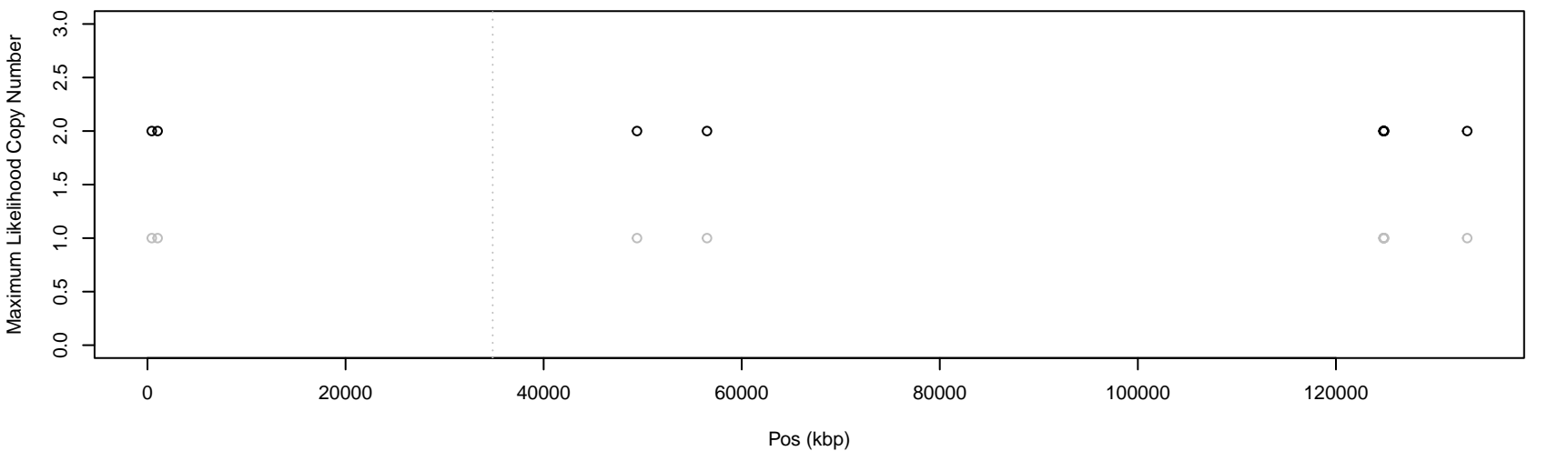
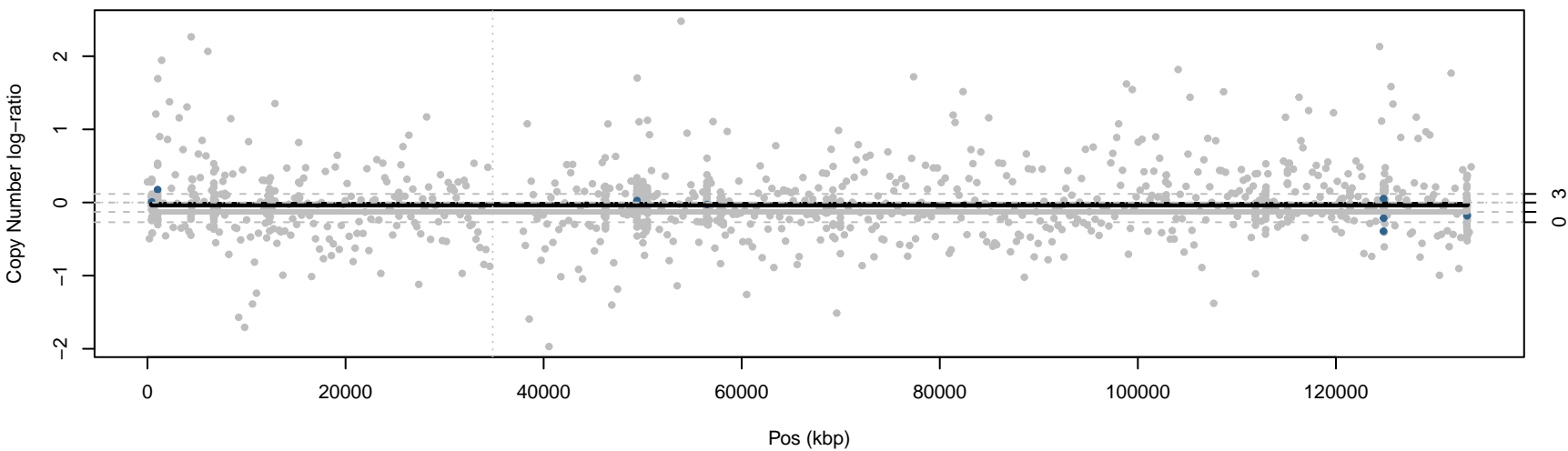
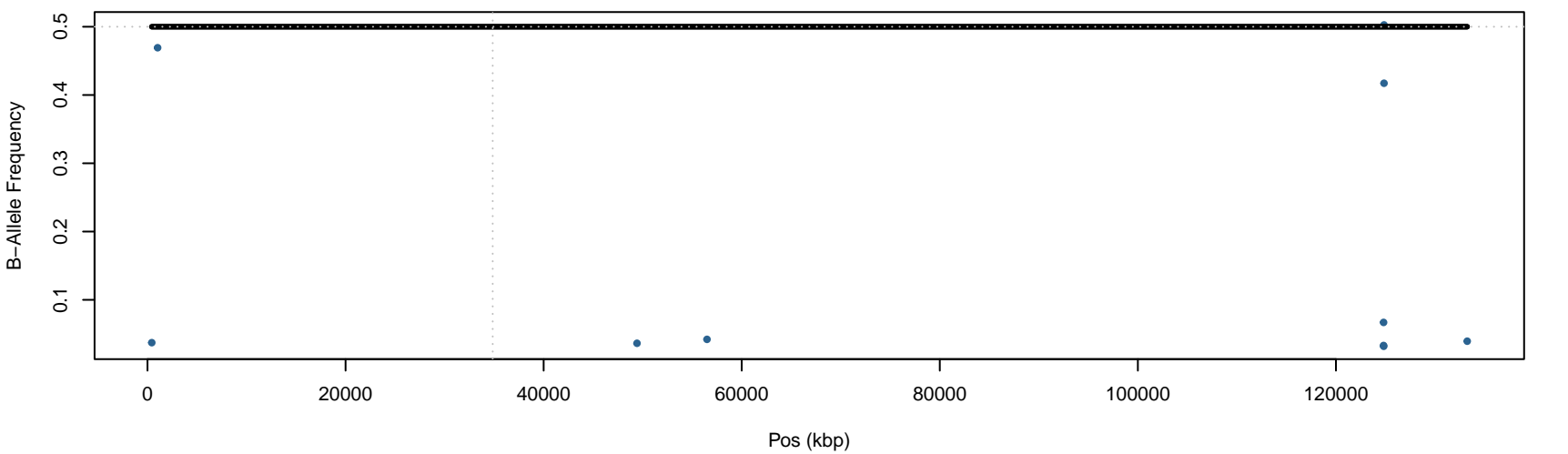
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 9



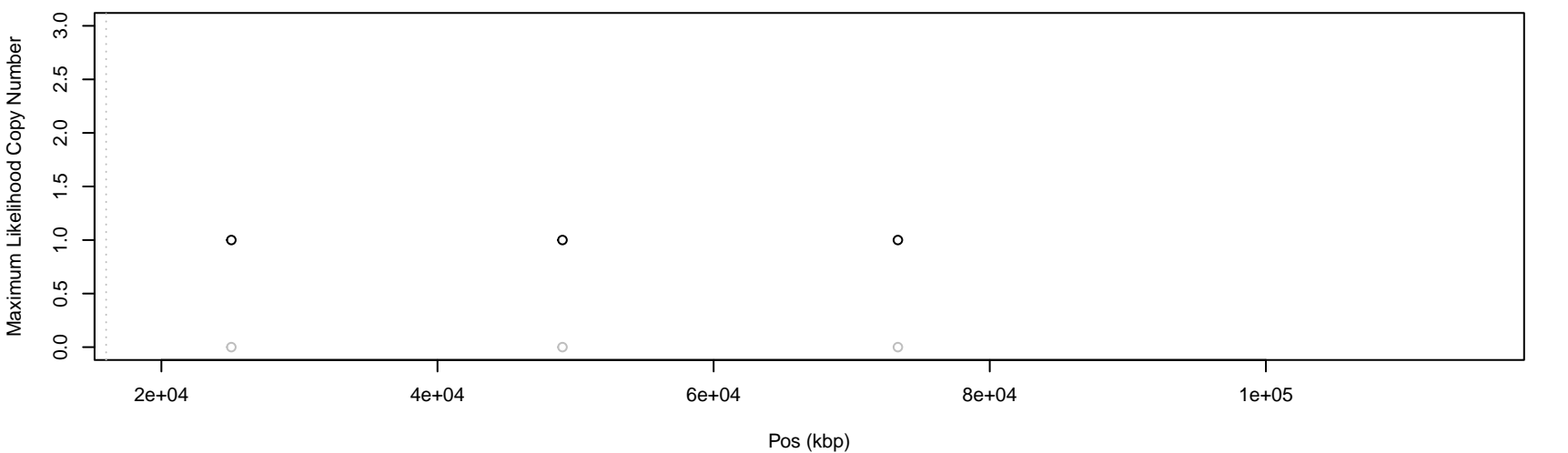
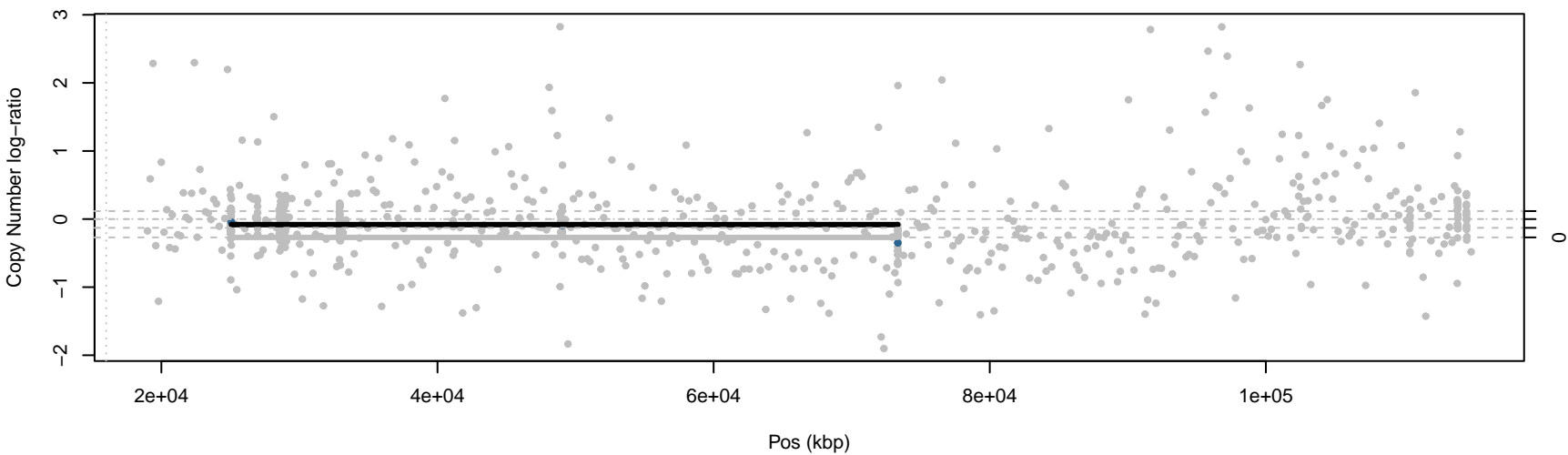
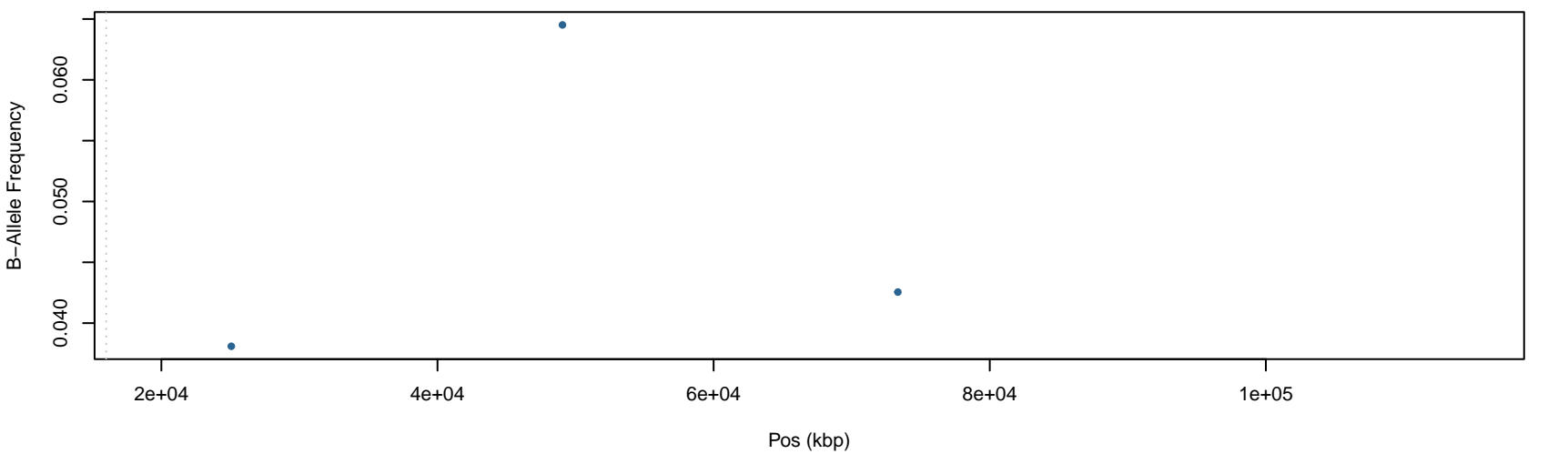
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 11



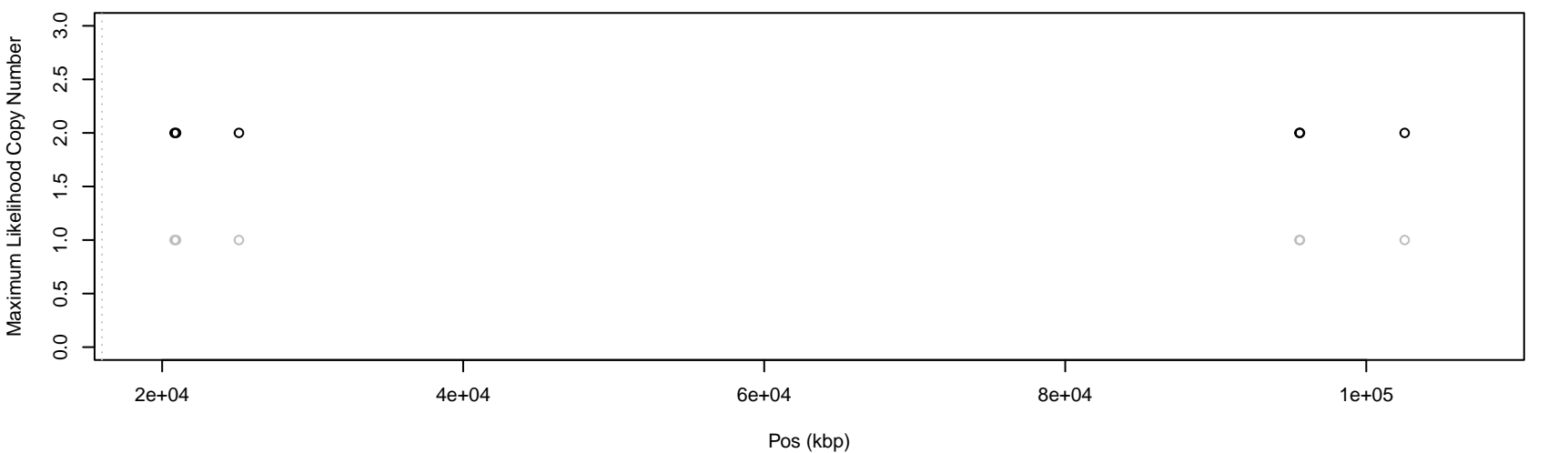
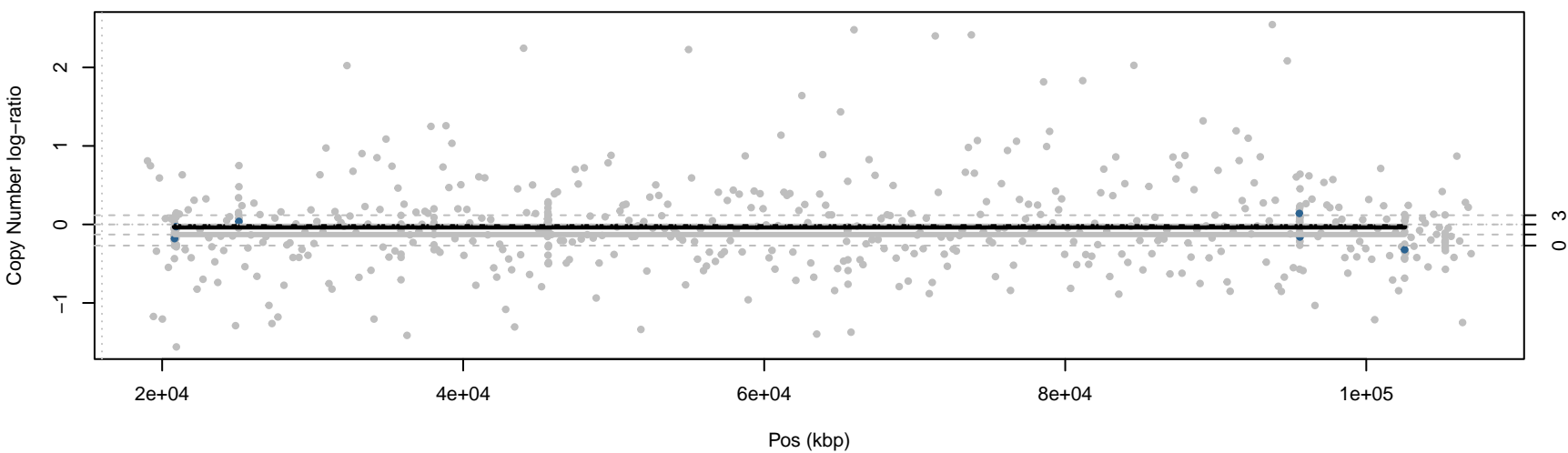
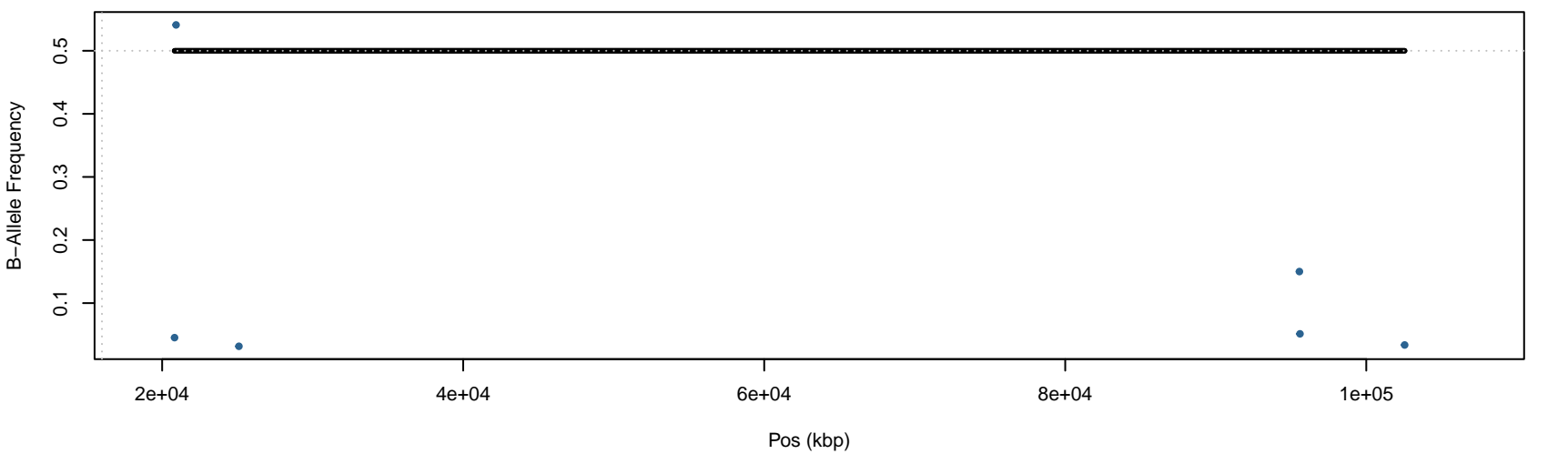
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 12



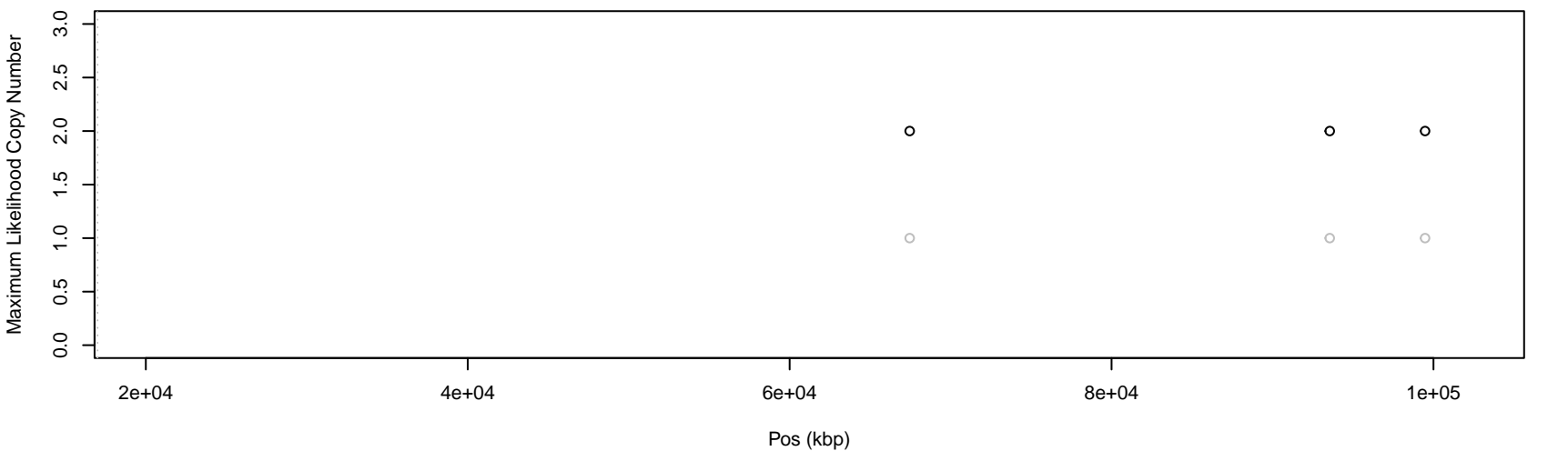
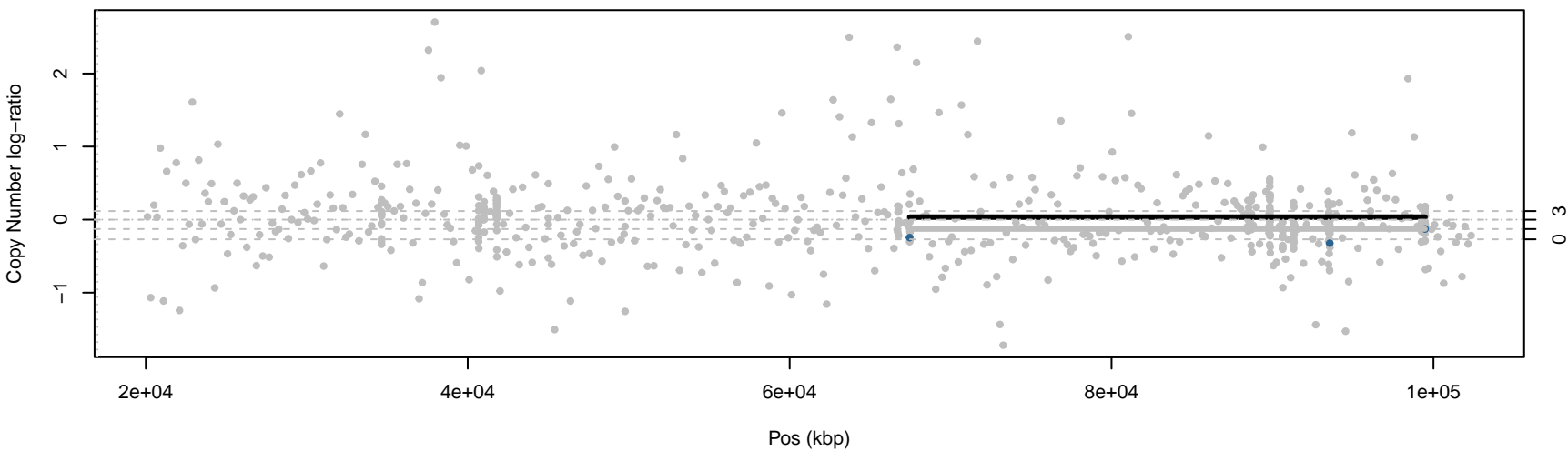
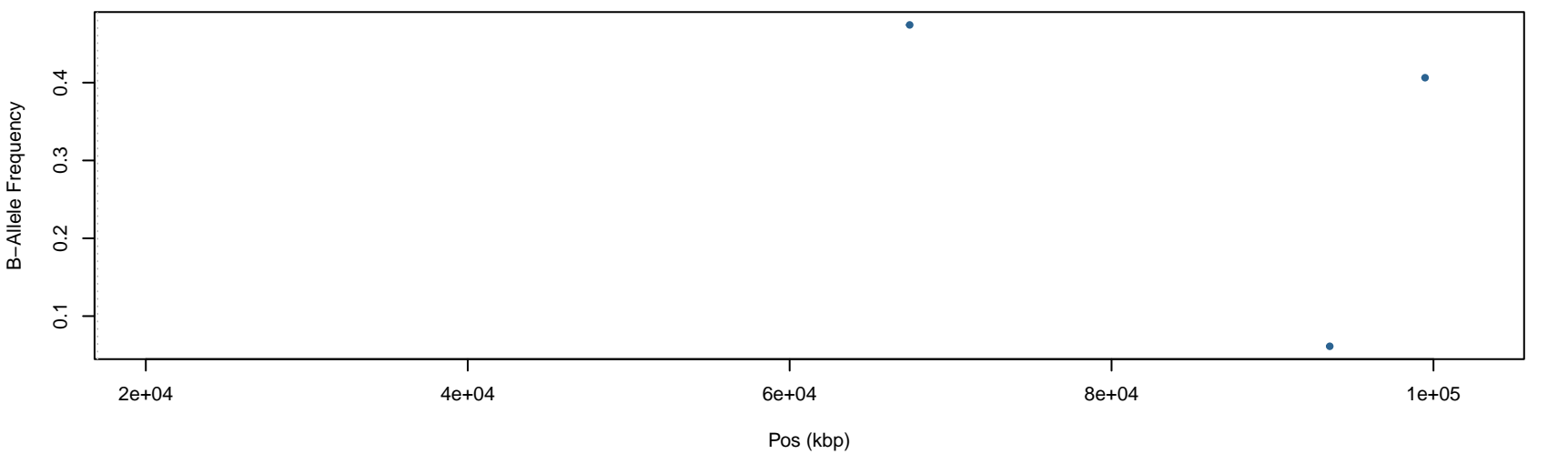
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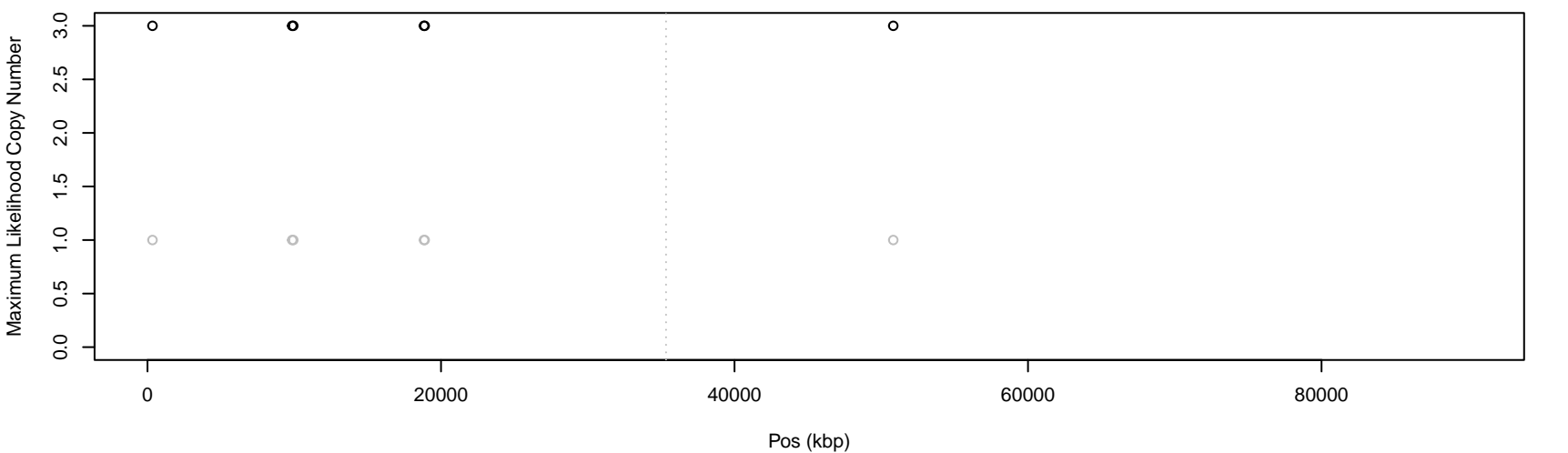
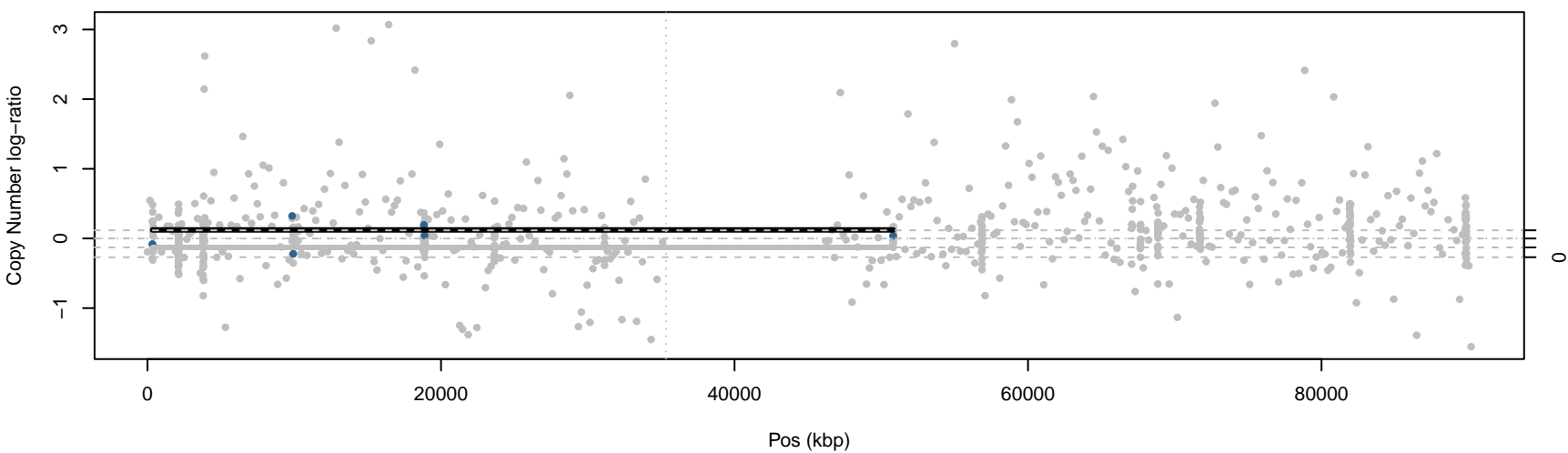
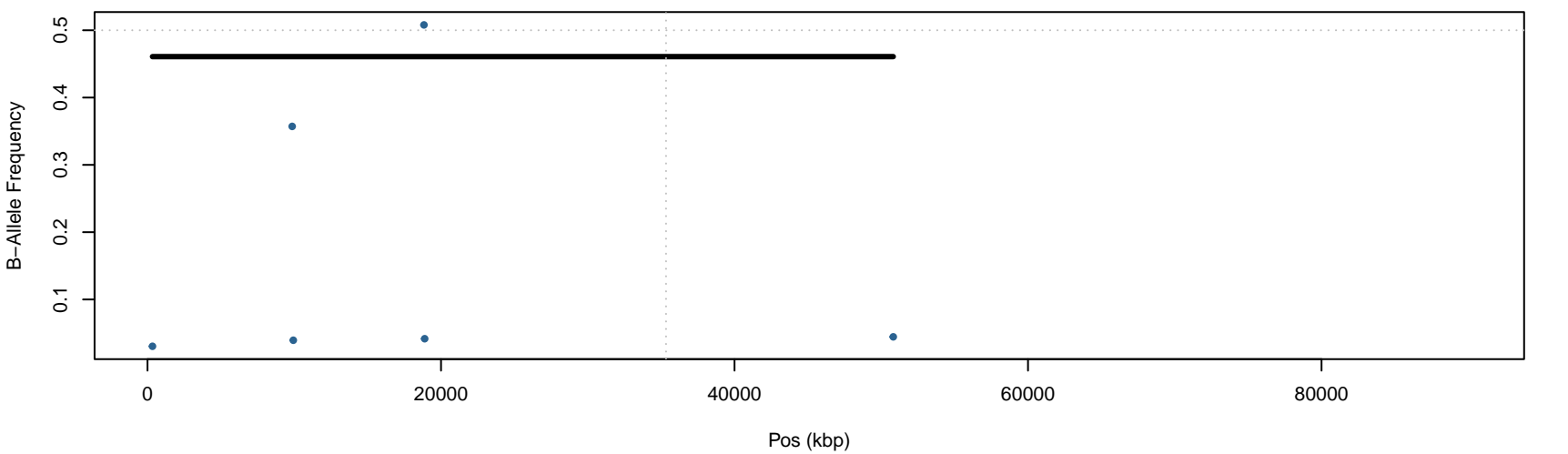
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 14



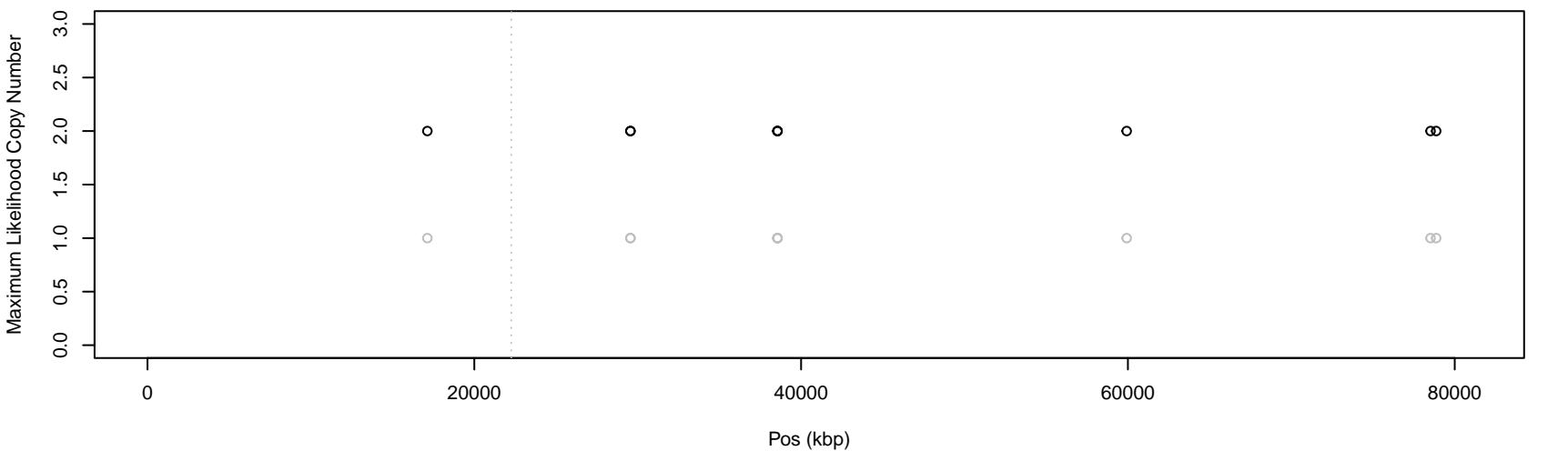
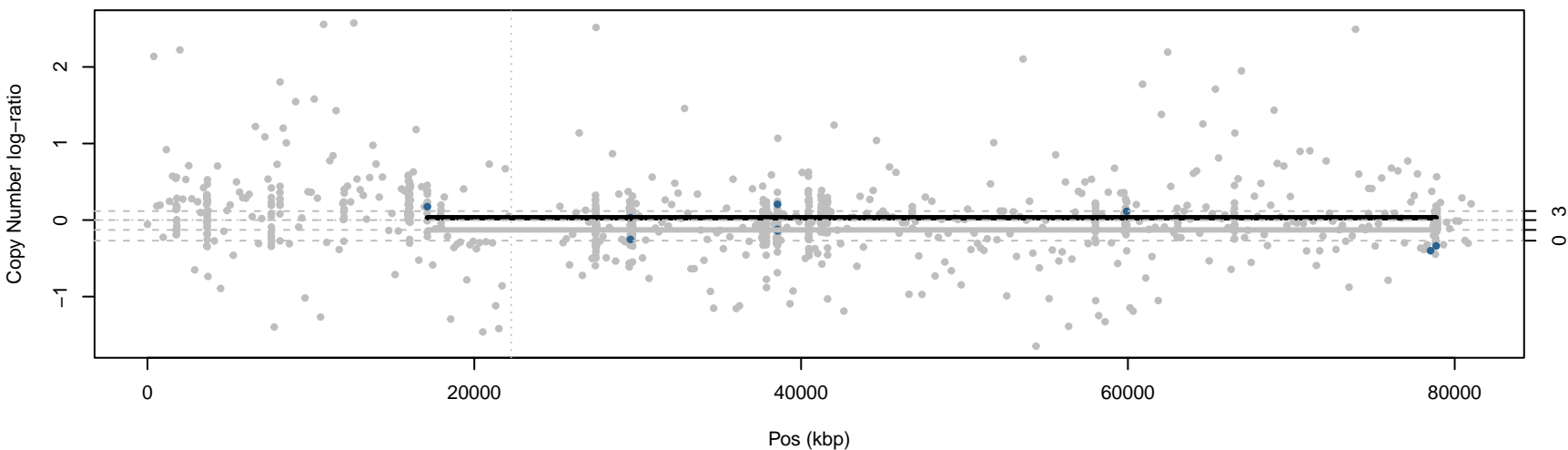
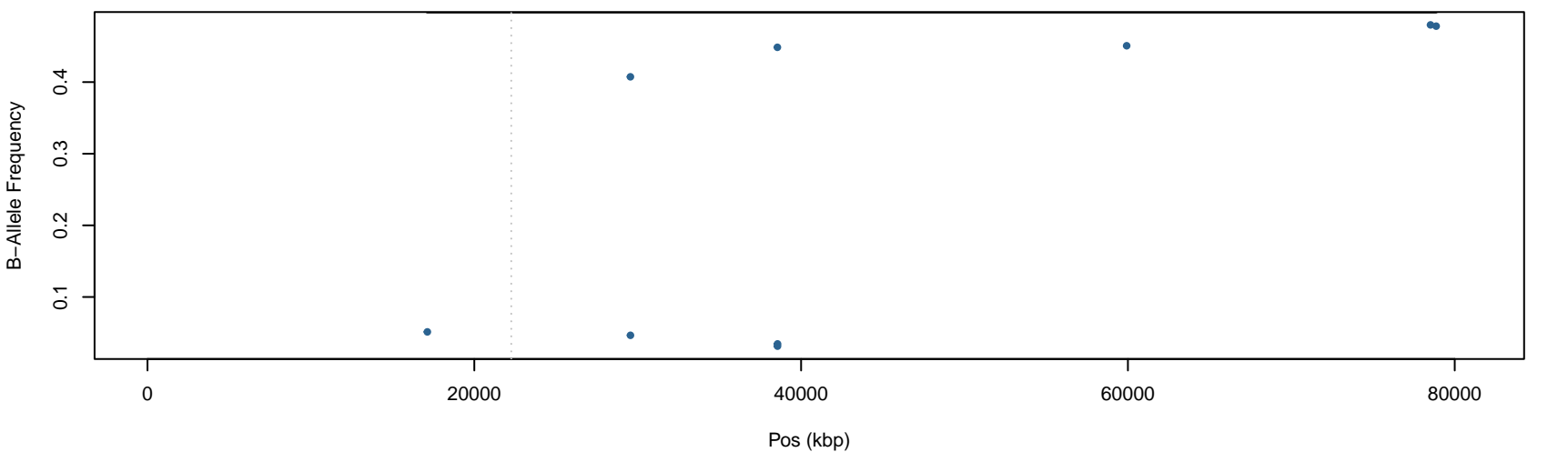
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 15



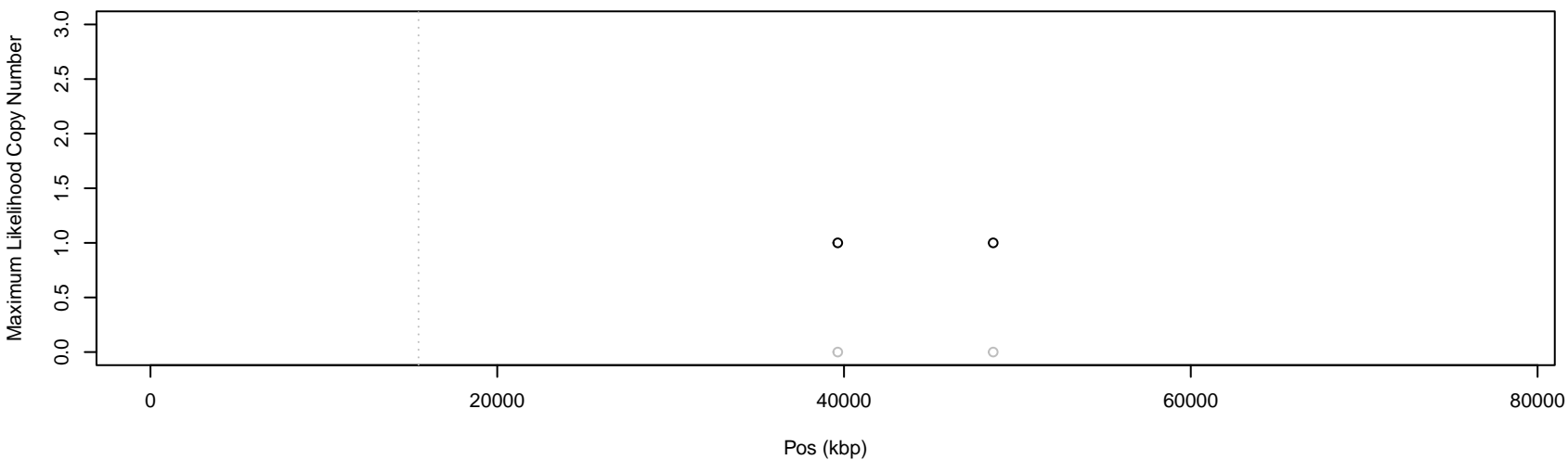
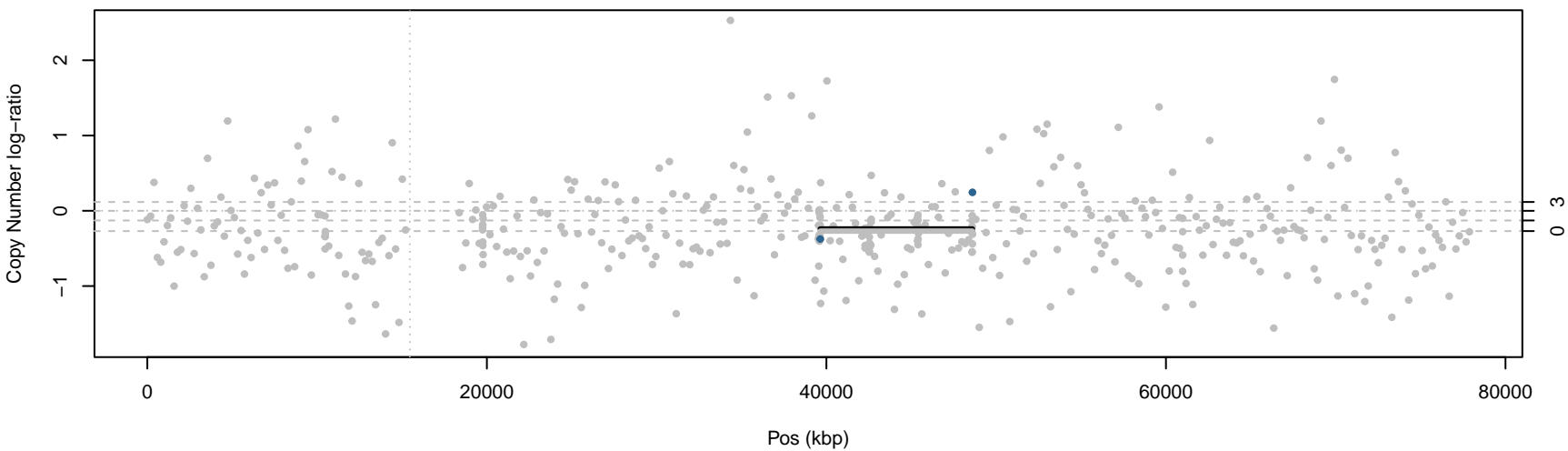
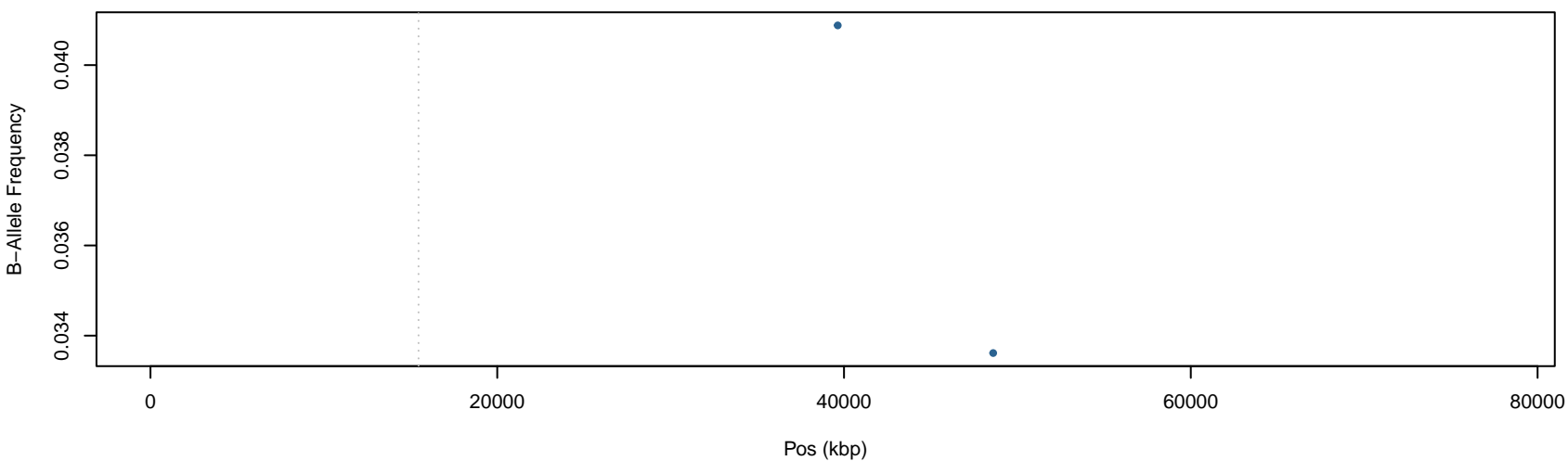
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 16



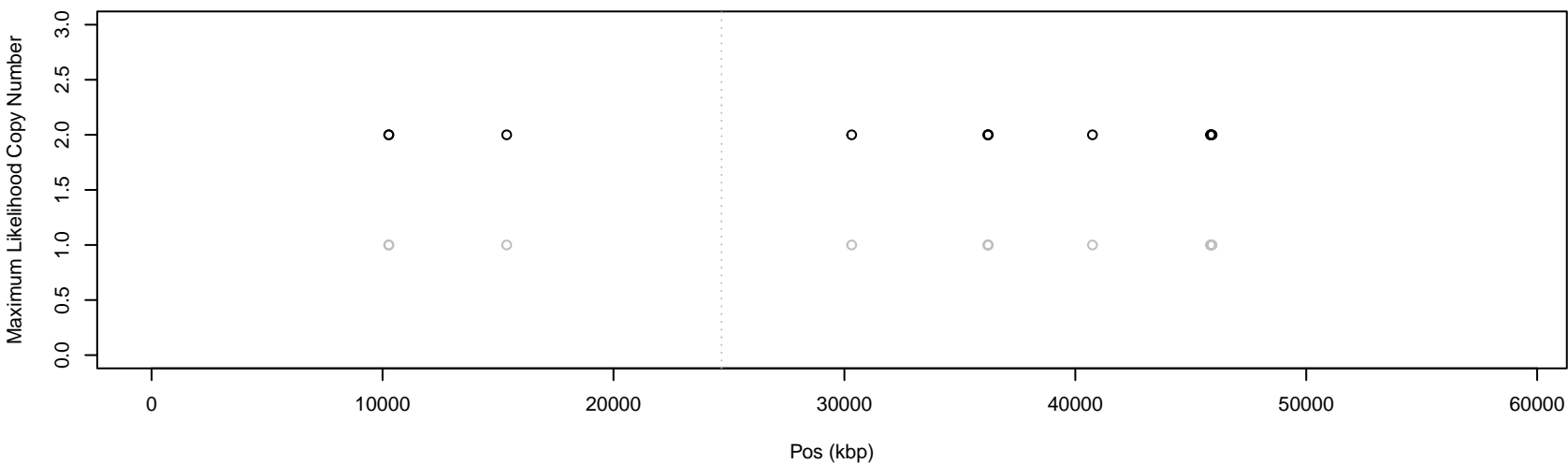
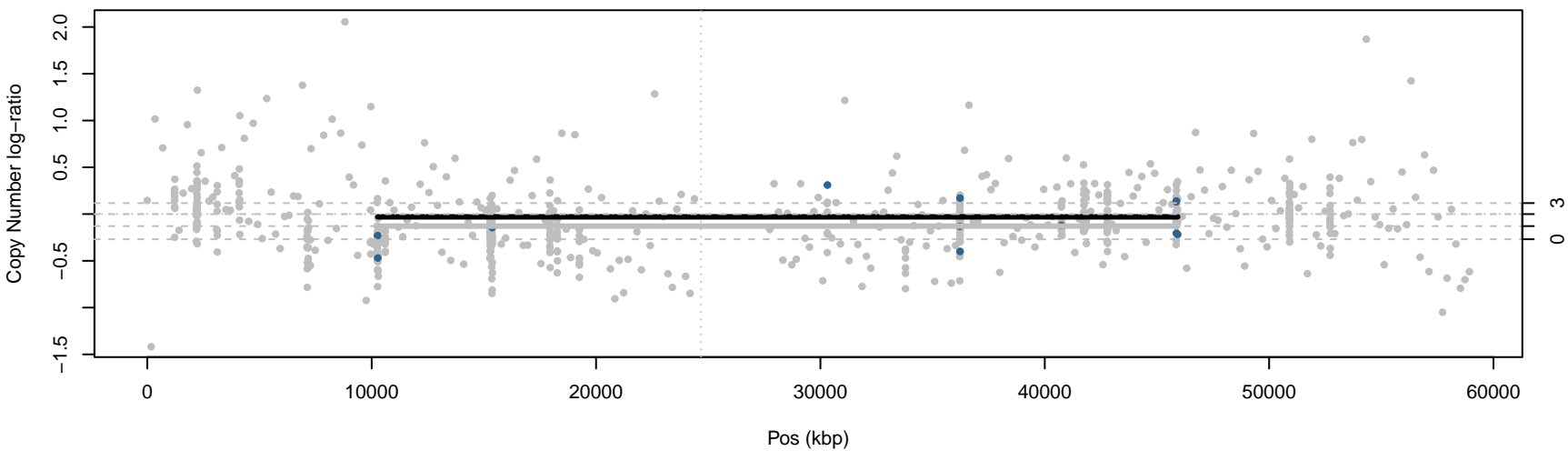
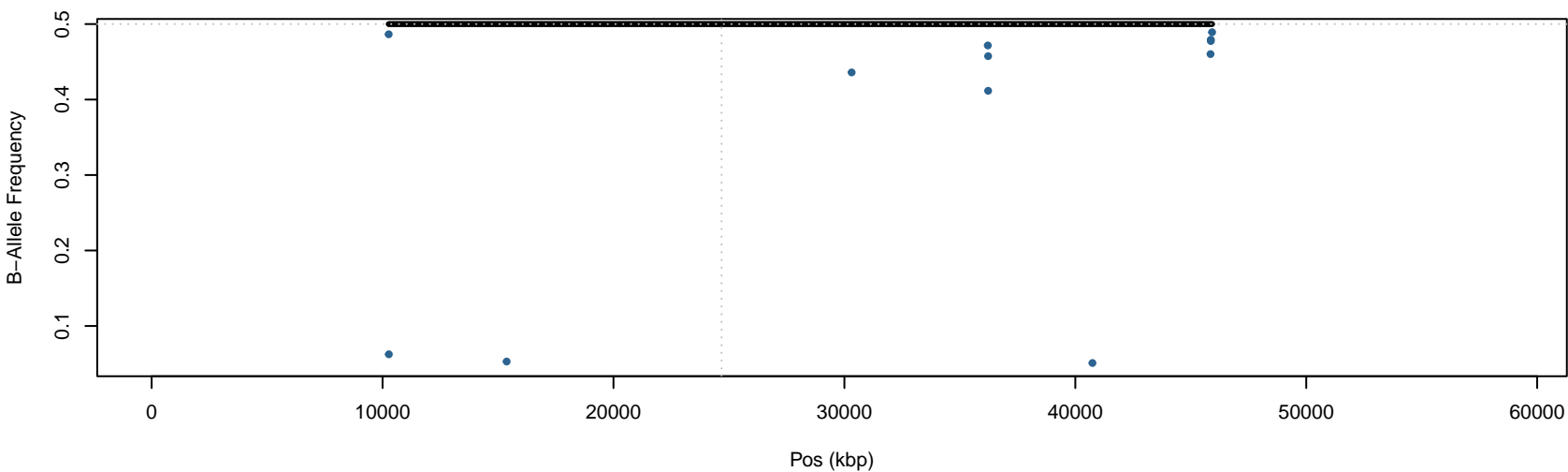
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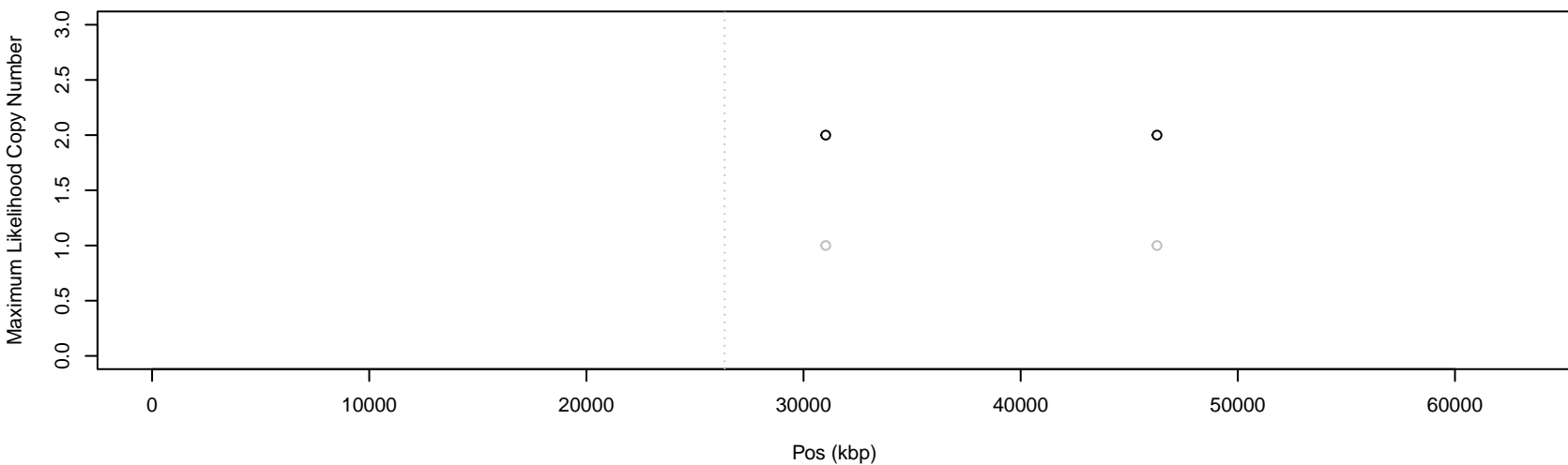
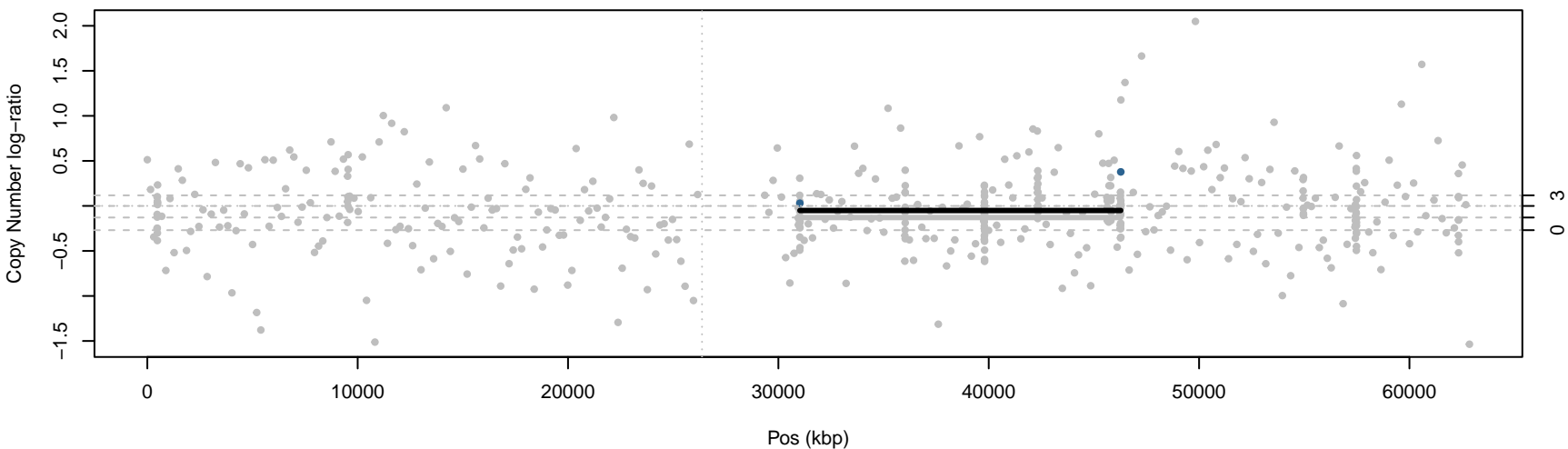
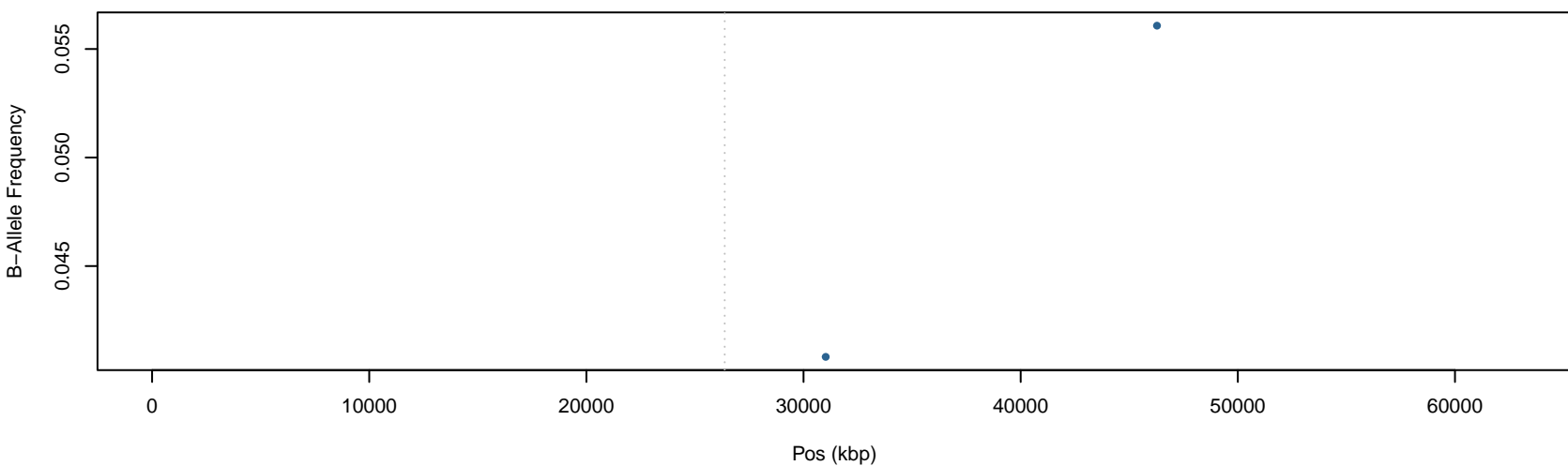
Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 18



Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 19



Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 20



Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281 Chromosome: 21

