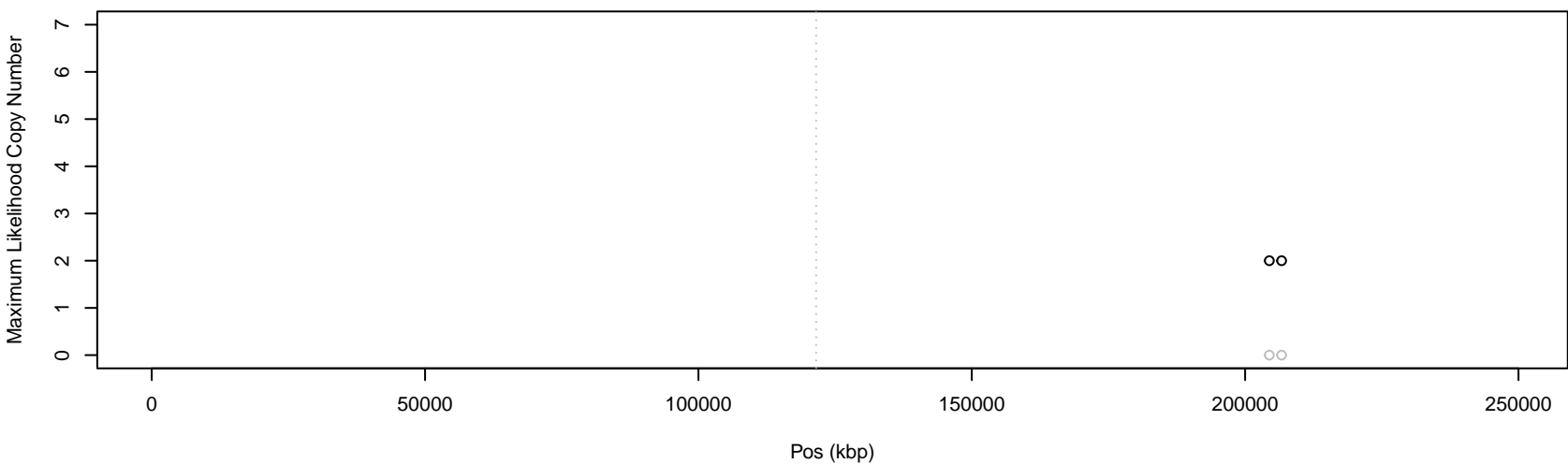
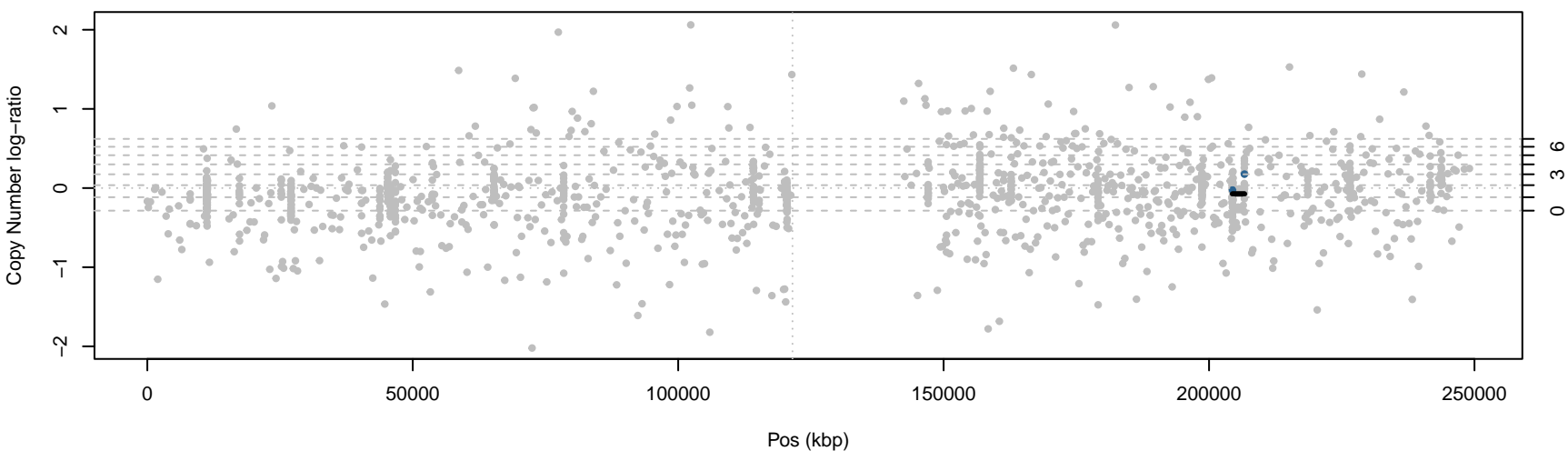
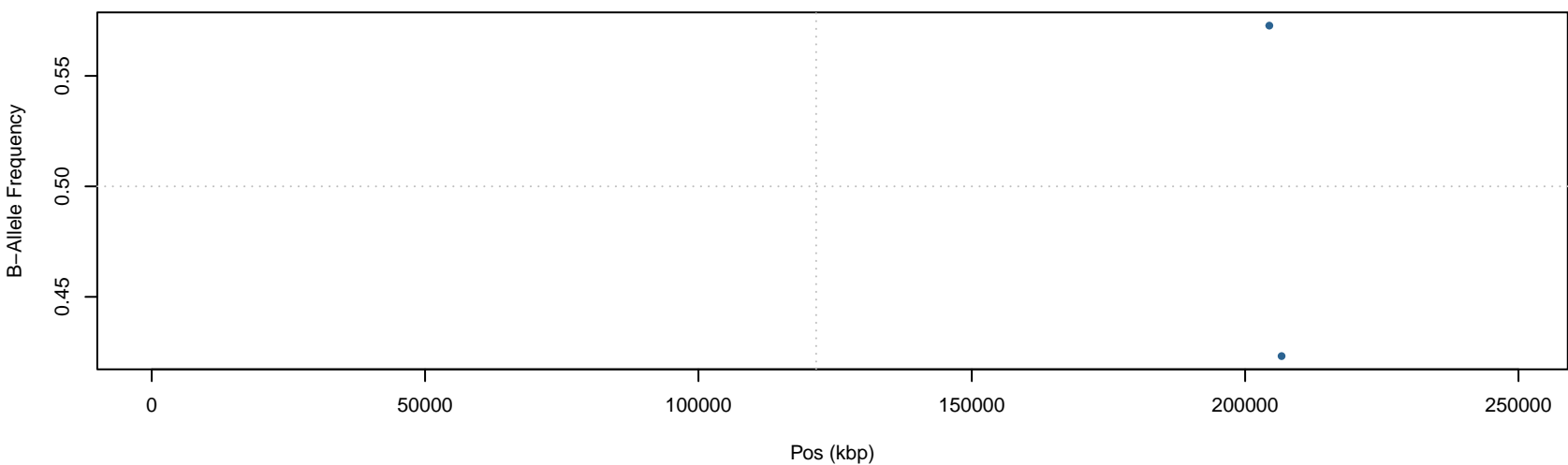
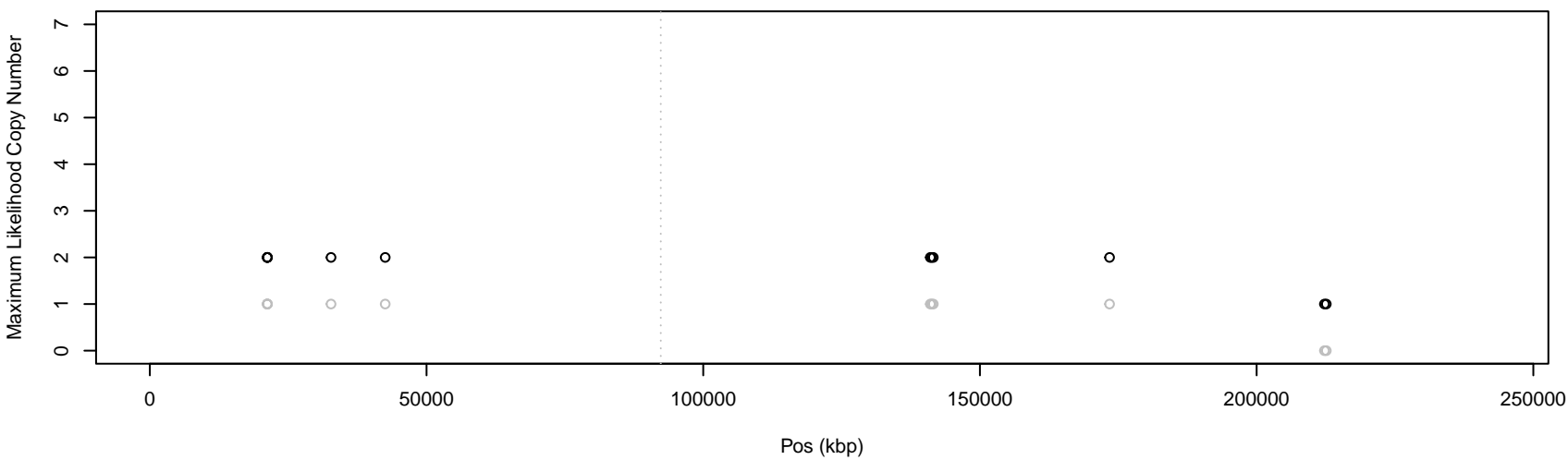
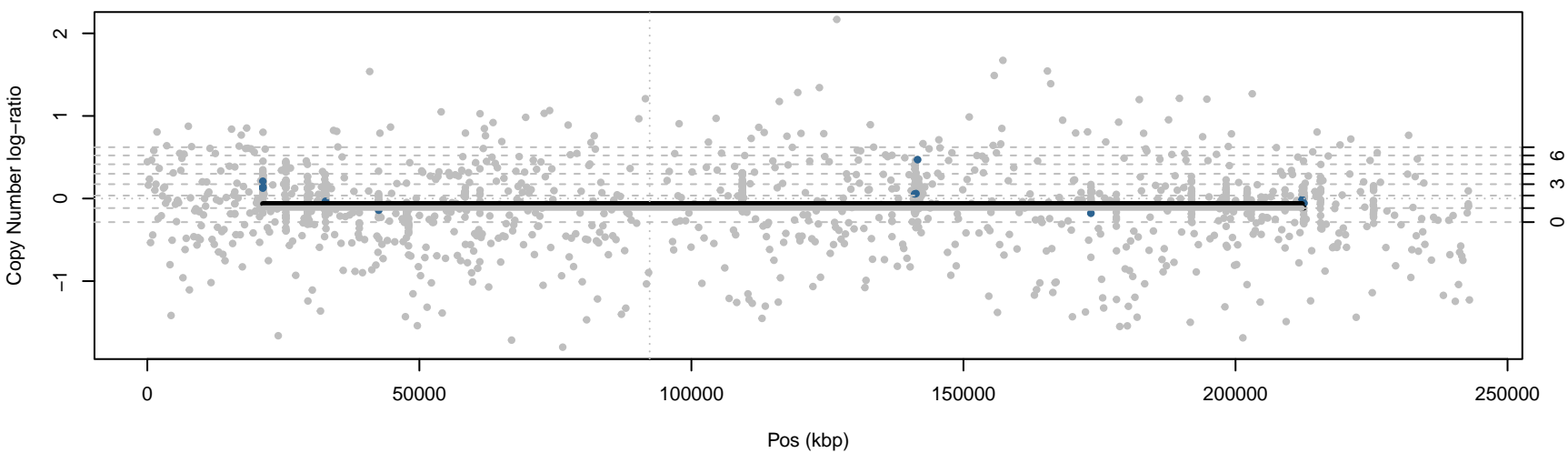
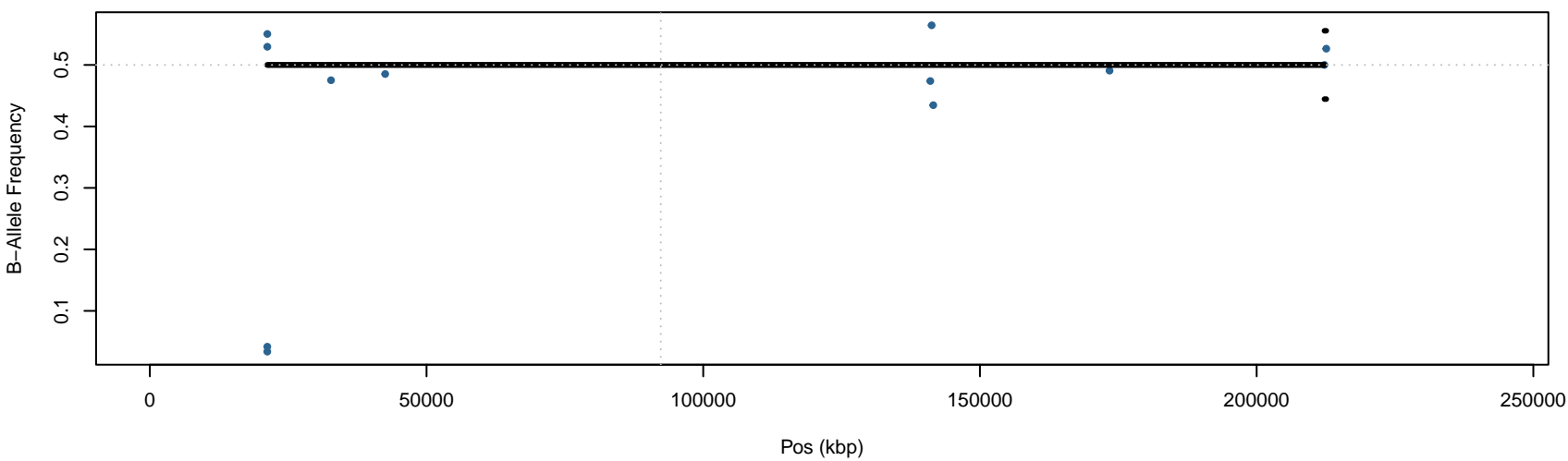


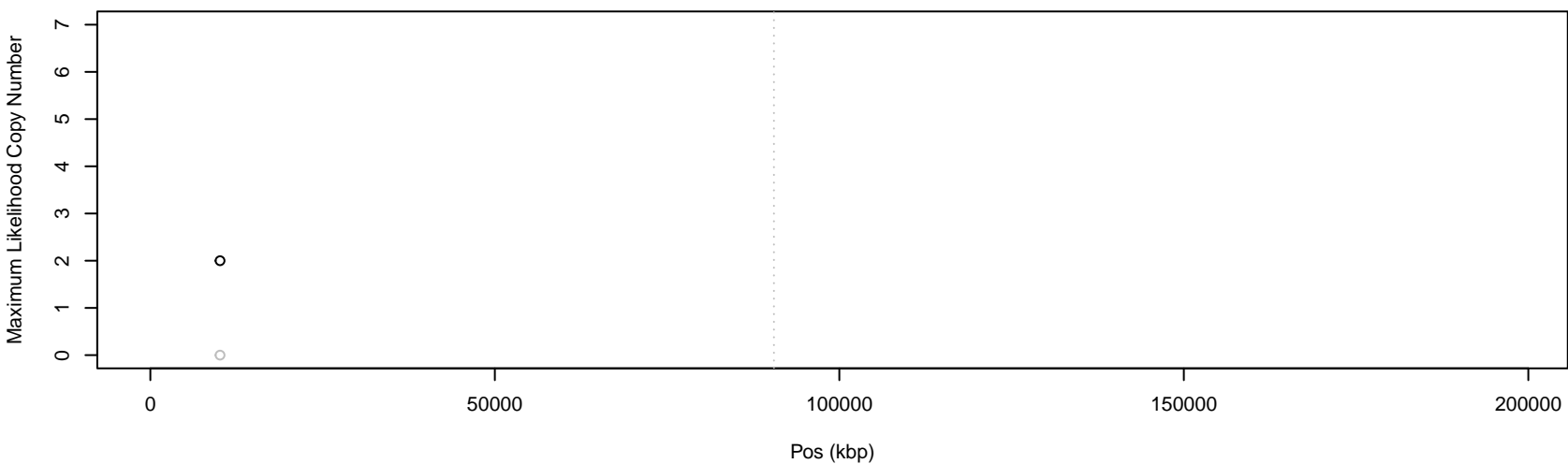
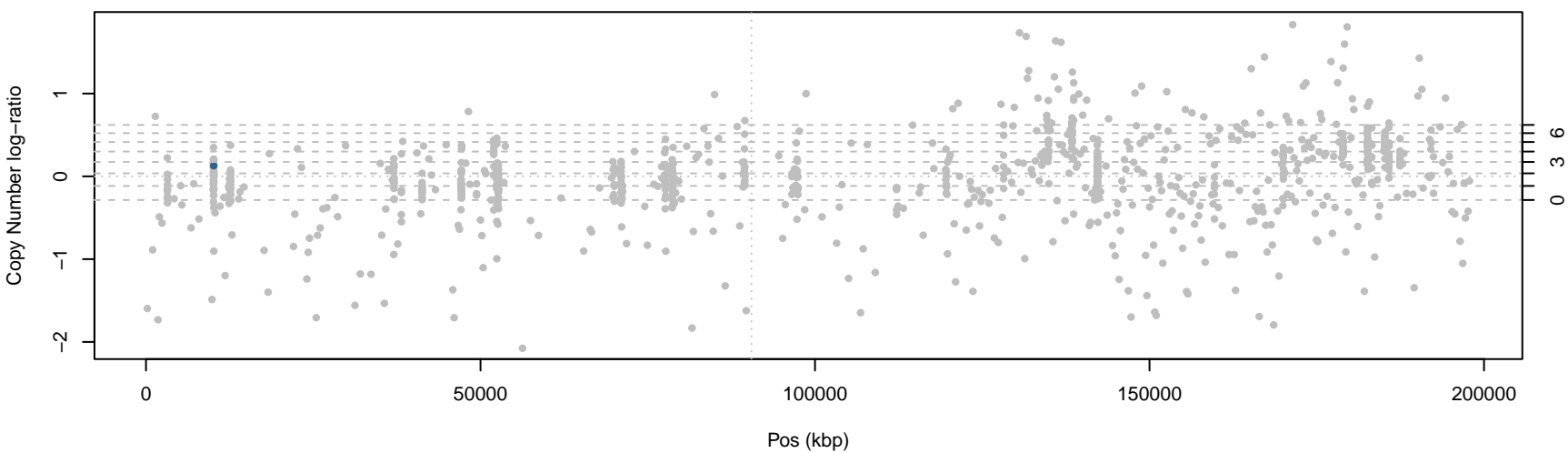
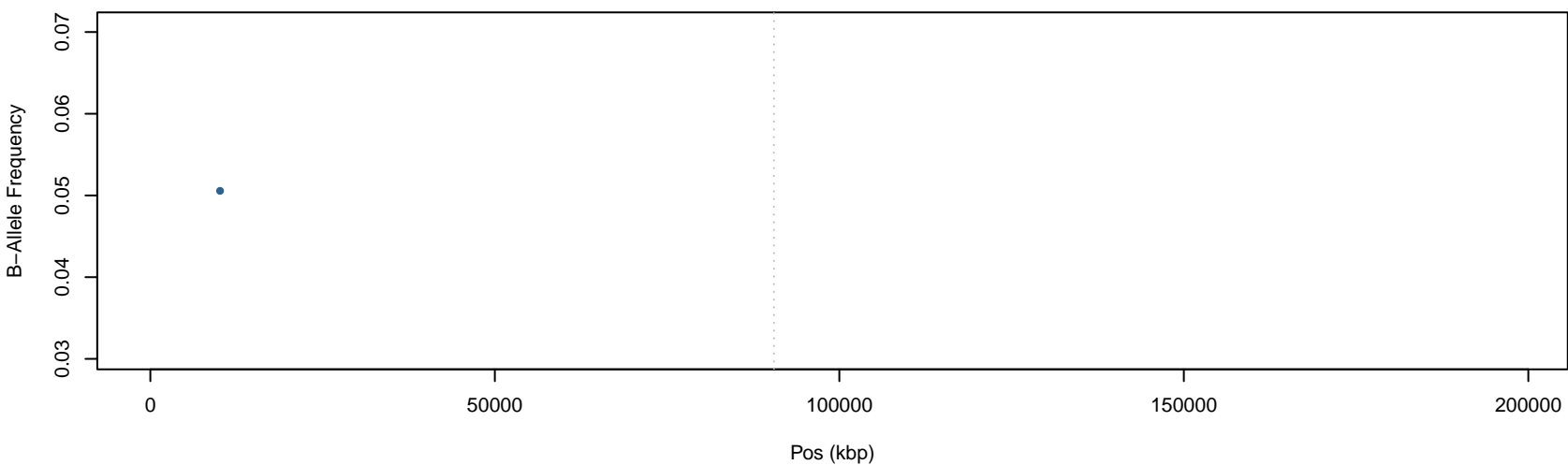
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 1



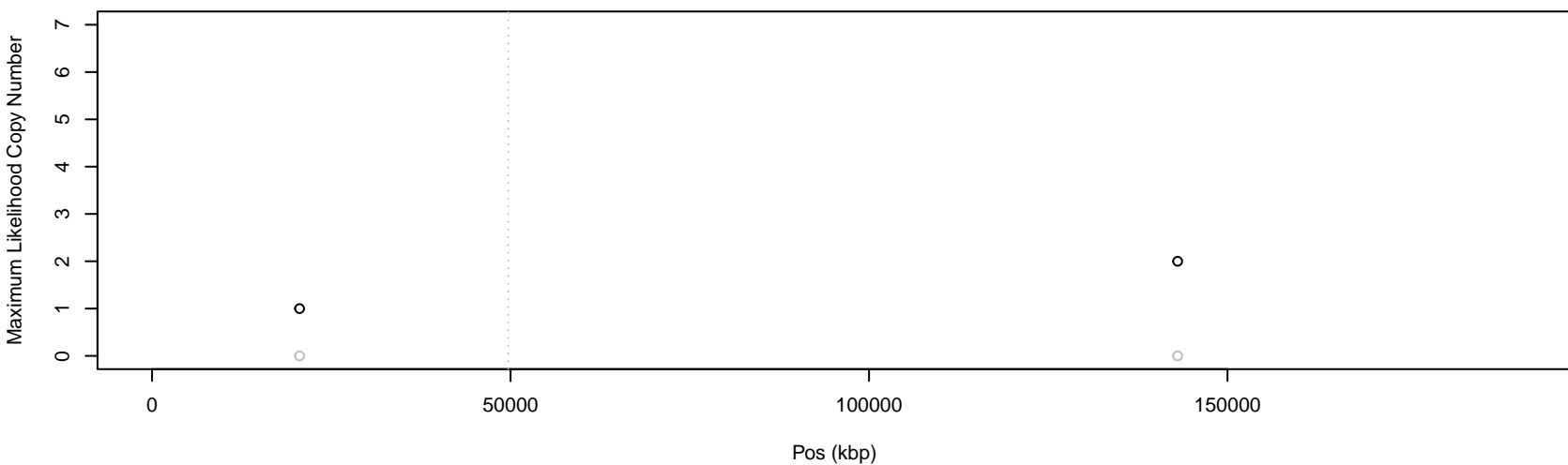
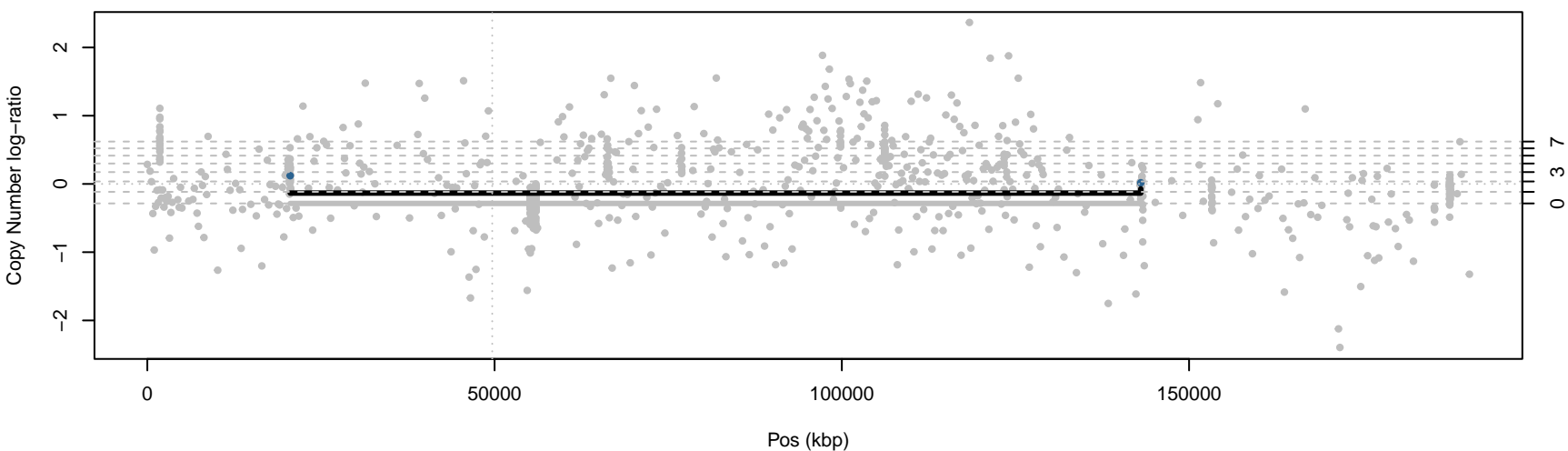
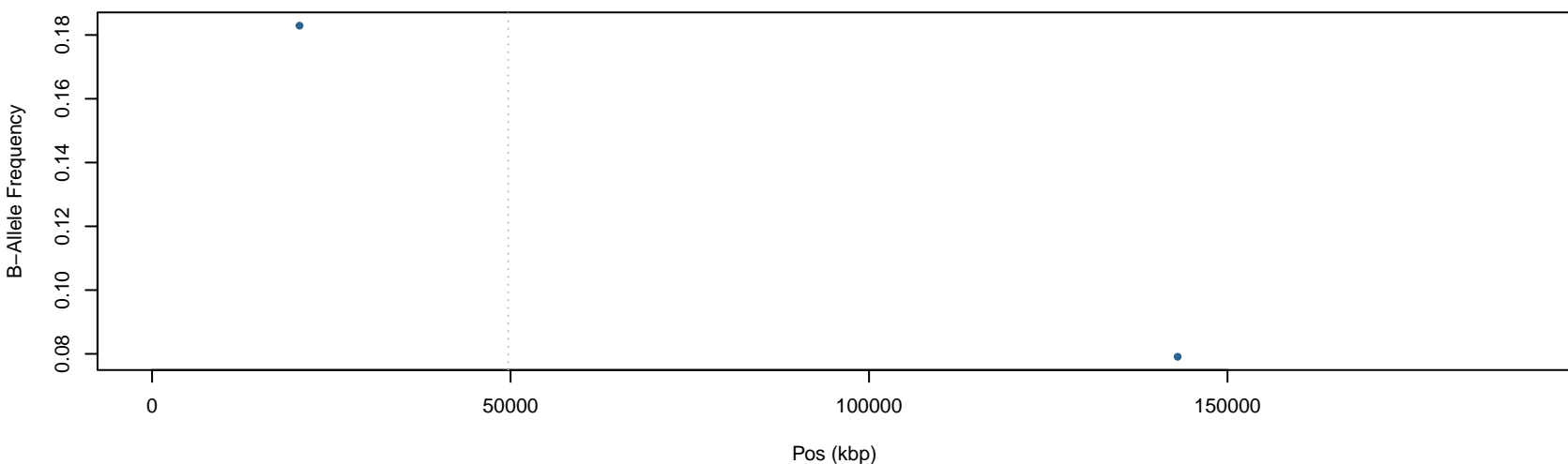
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 2



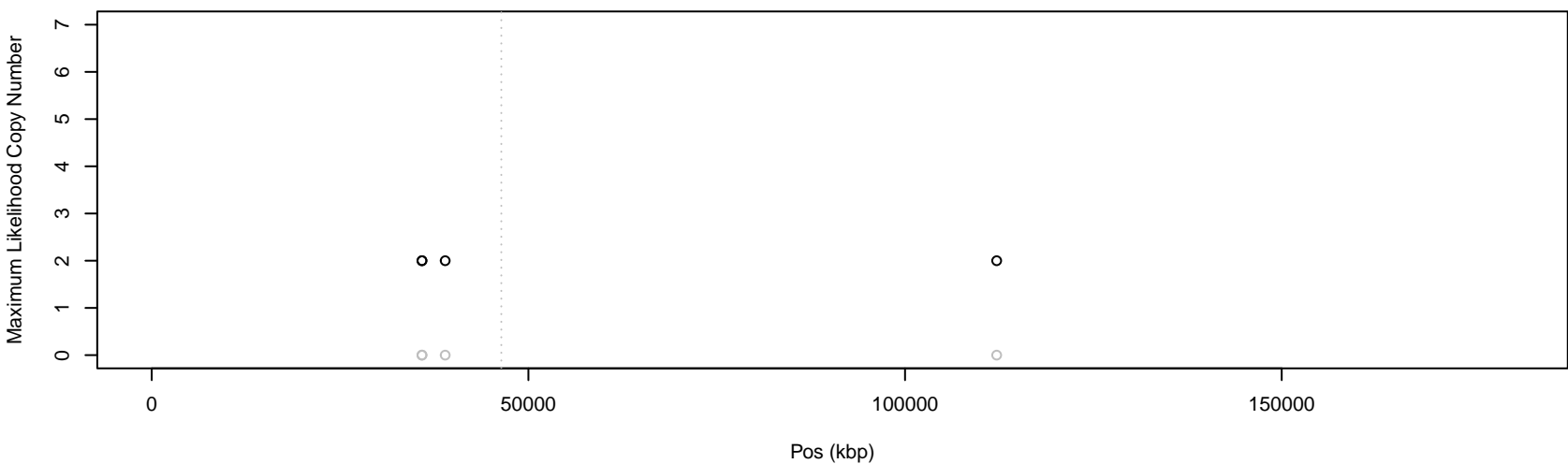
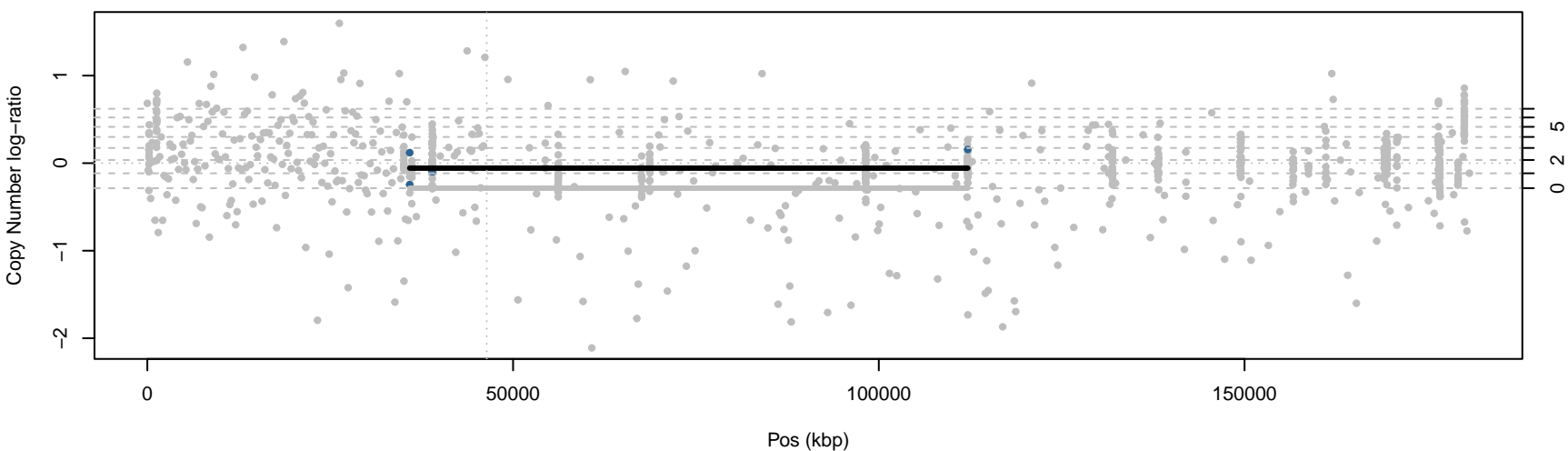
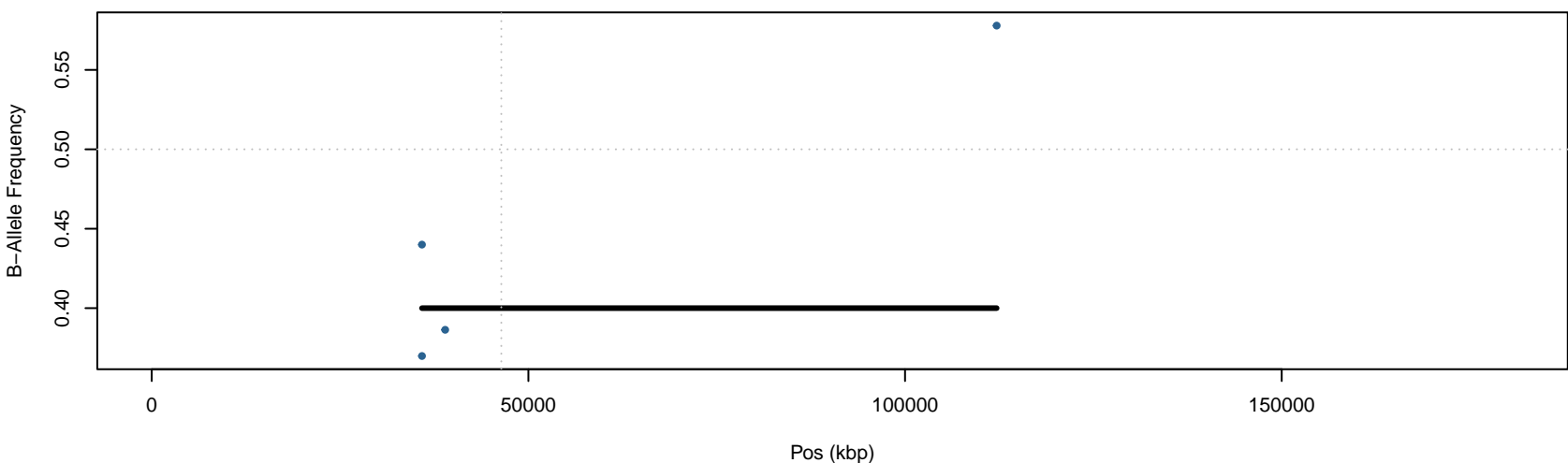
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 3



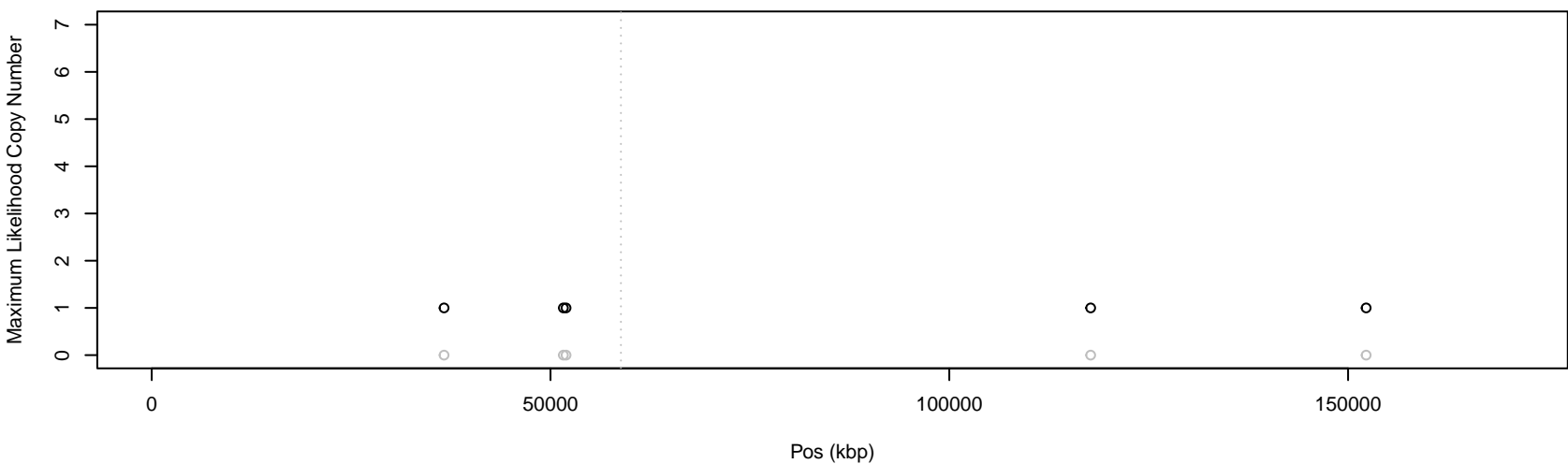
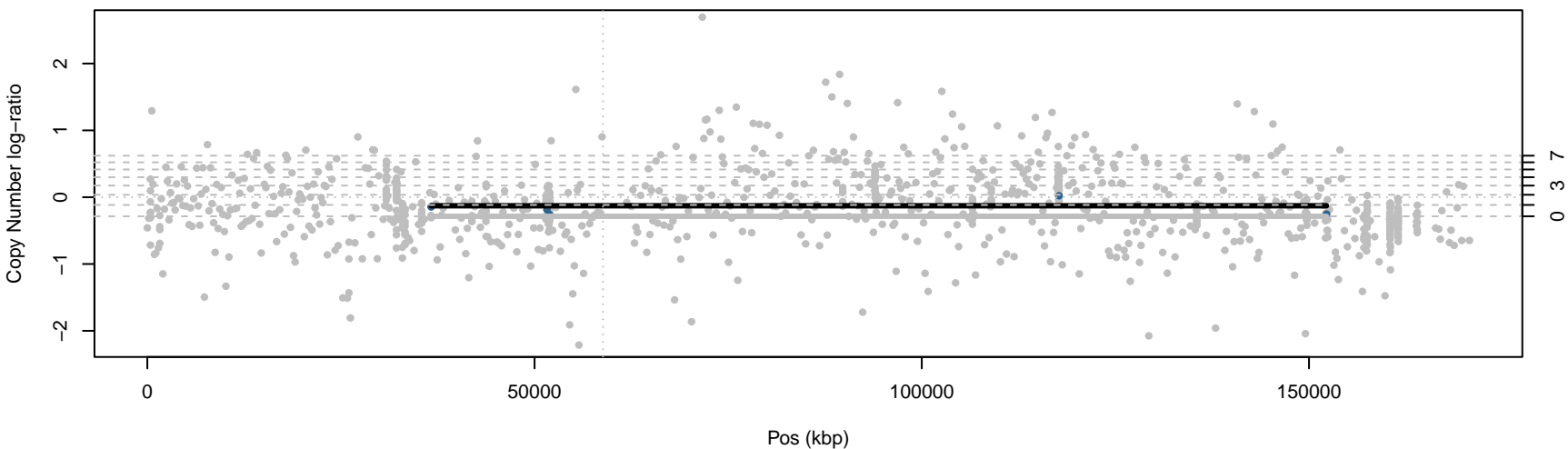
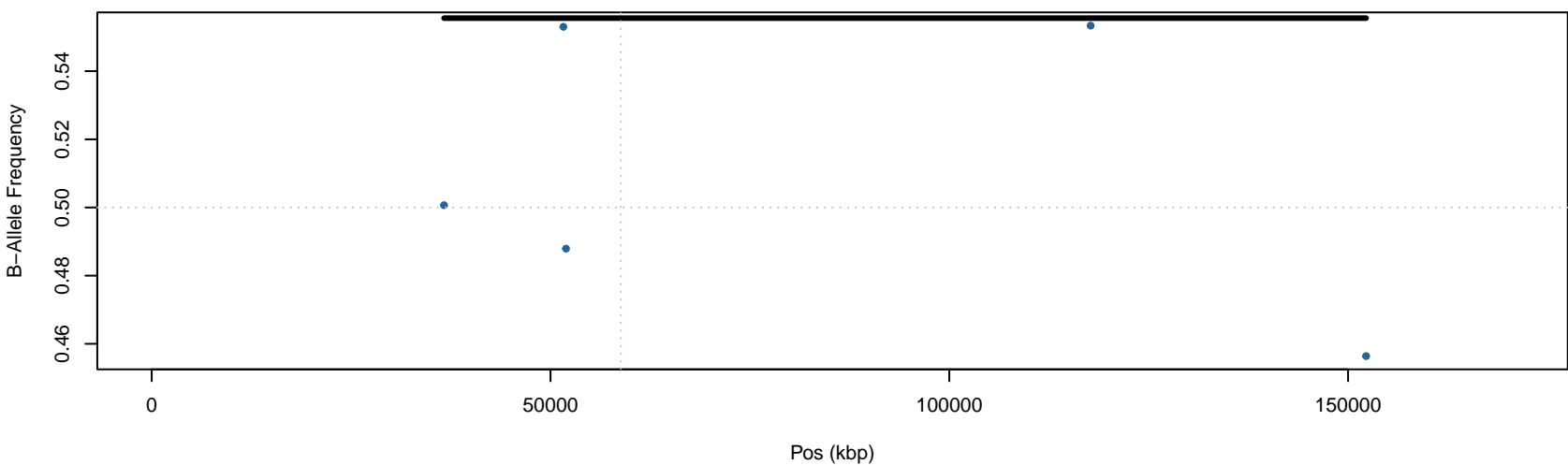
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 4



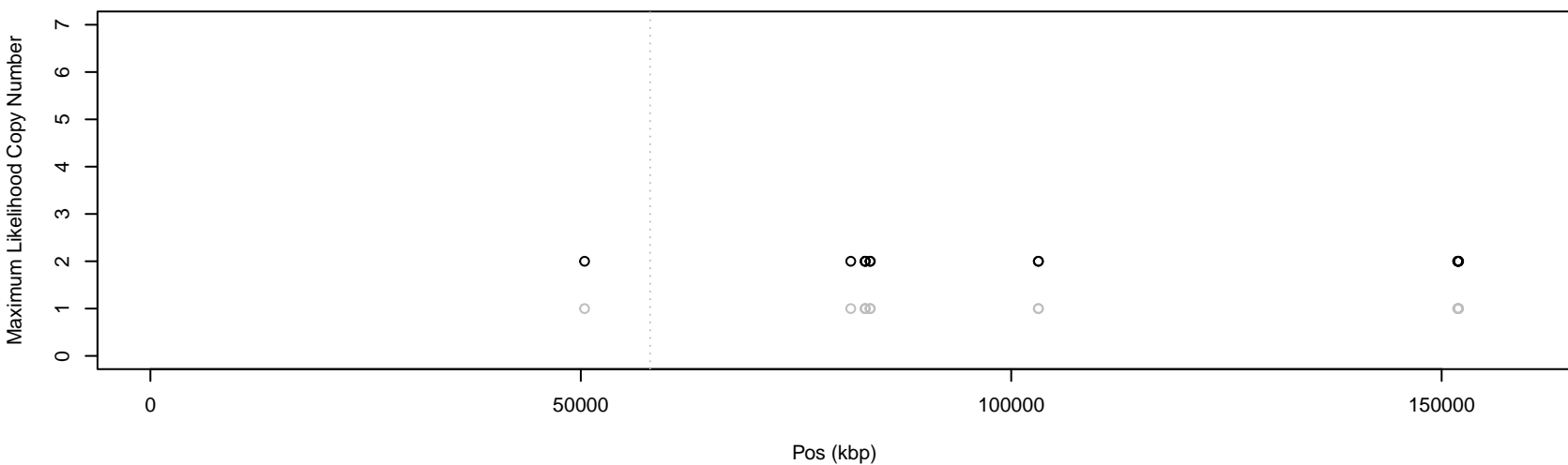
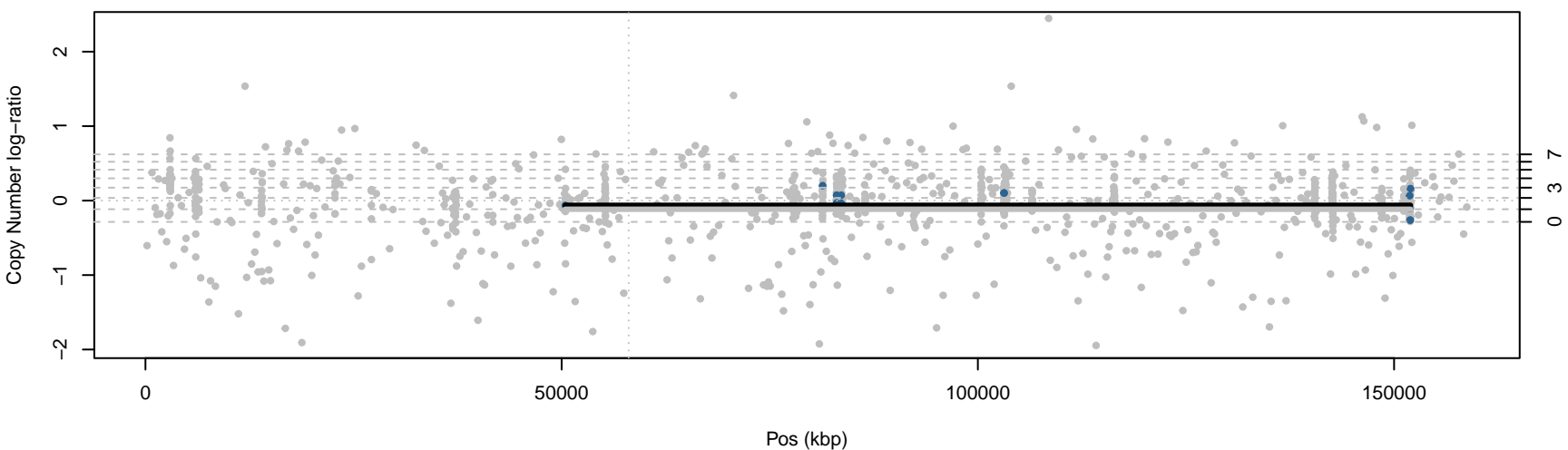
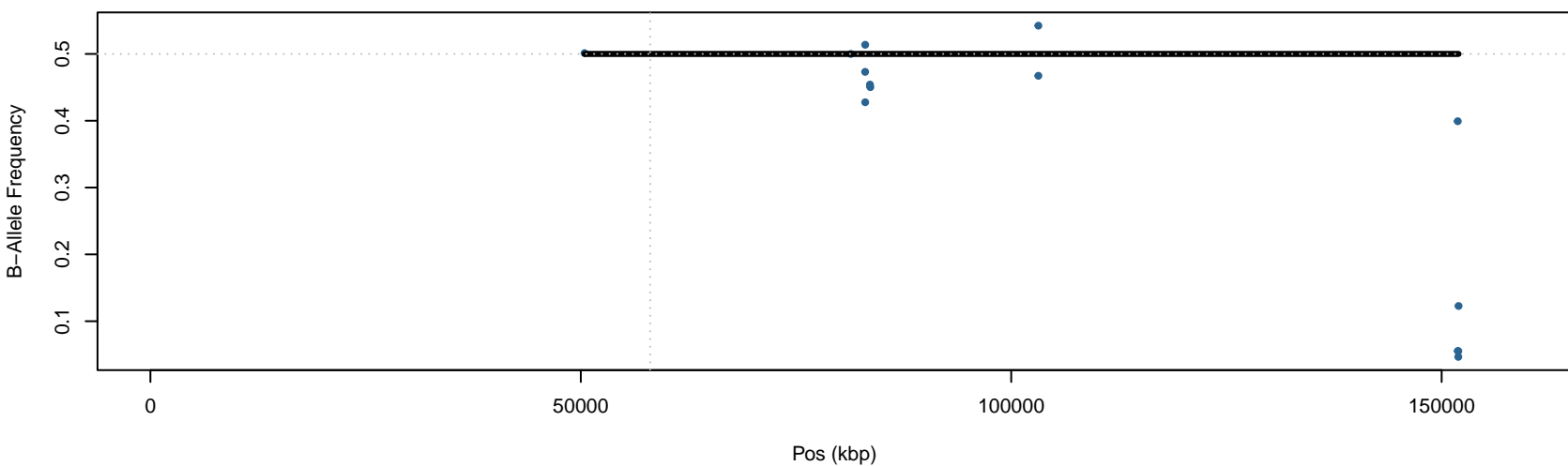
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 5



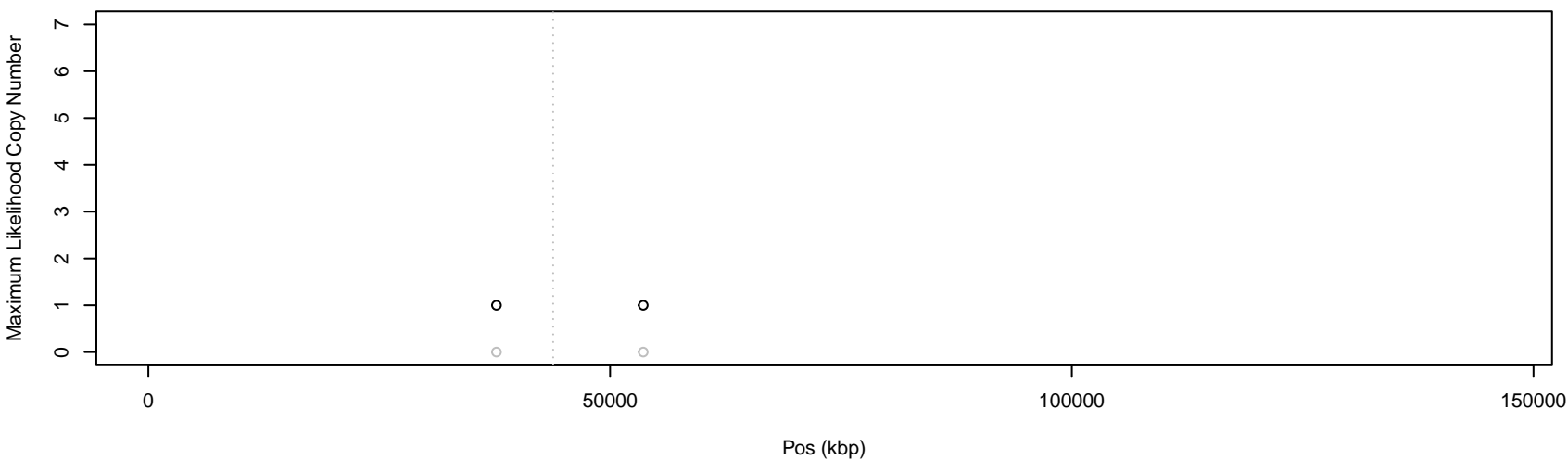
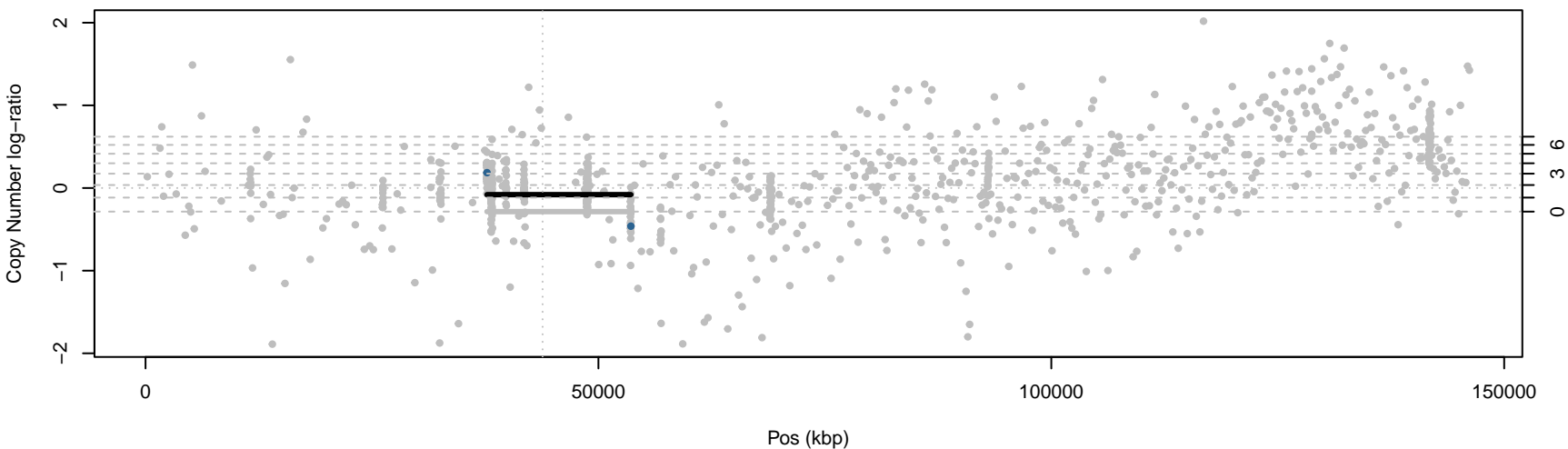
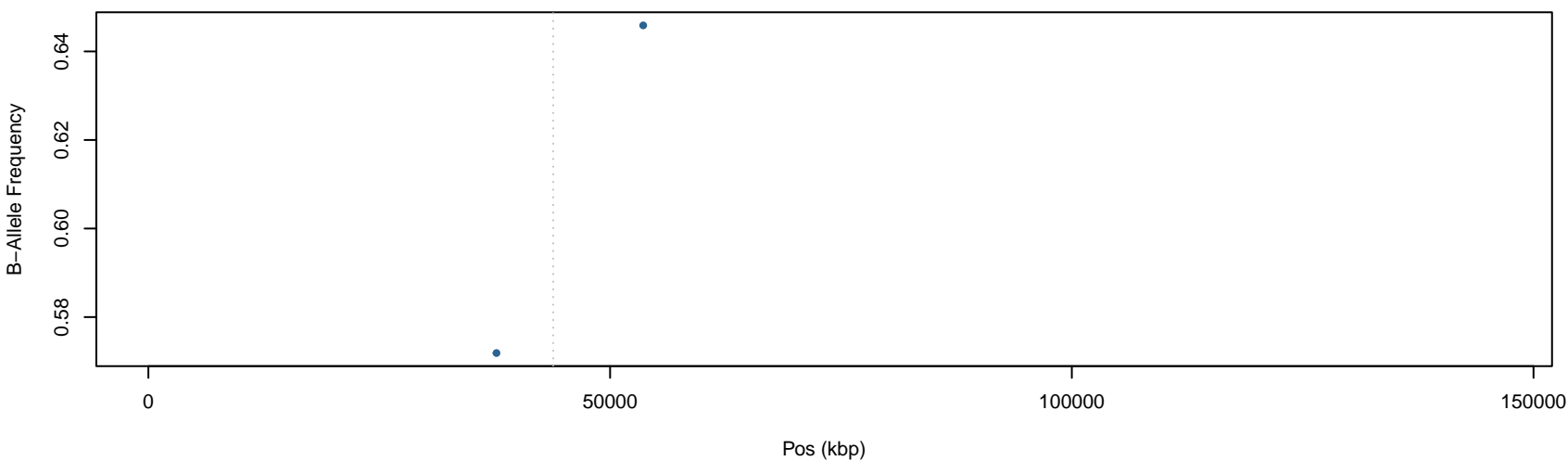
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 6



Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 7

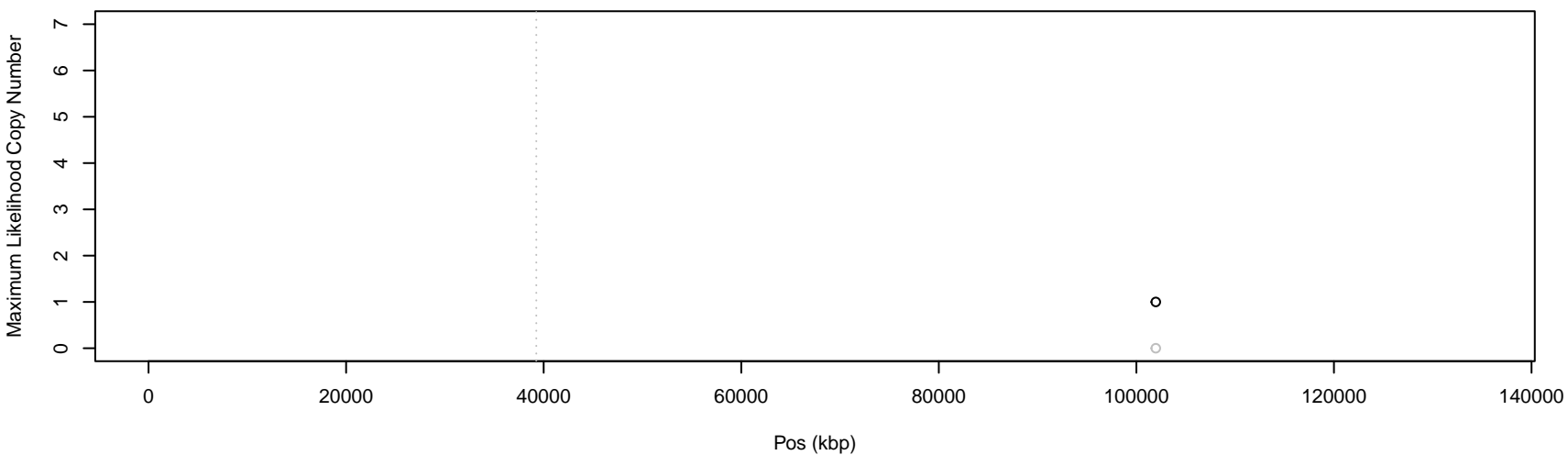
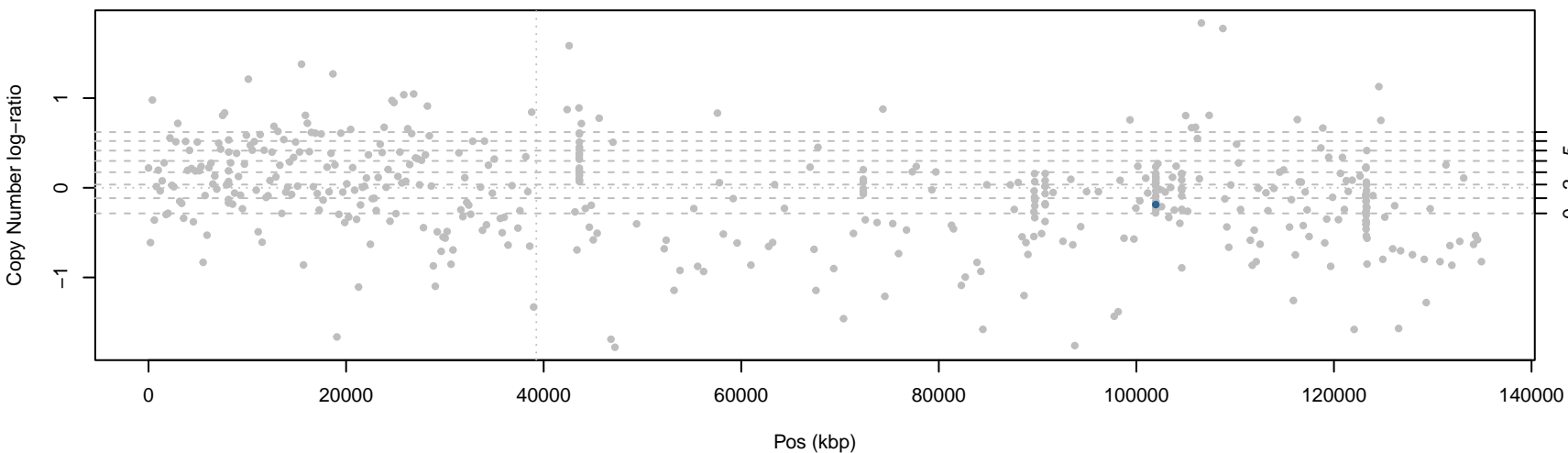
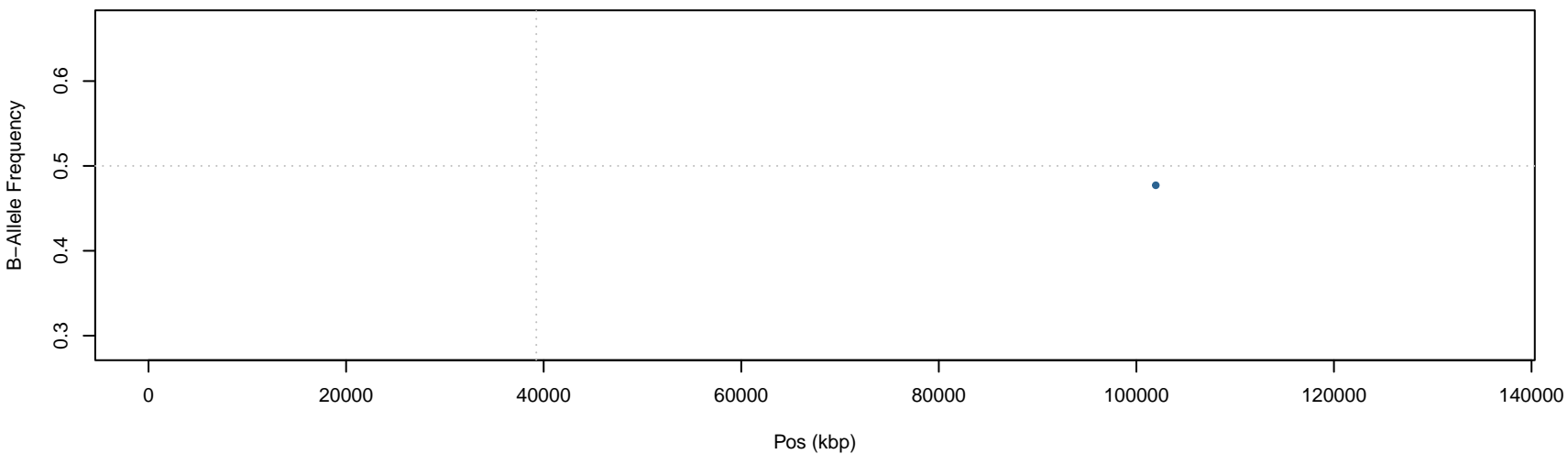


Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 8

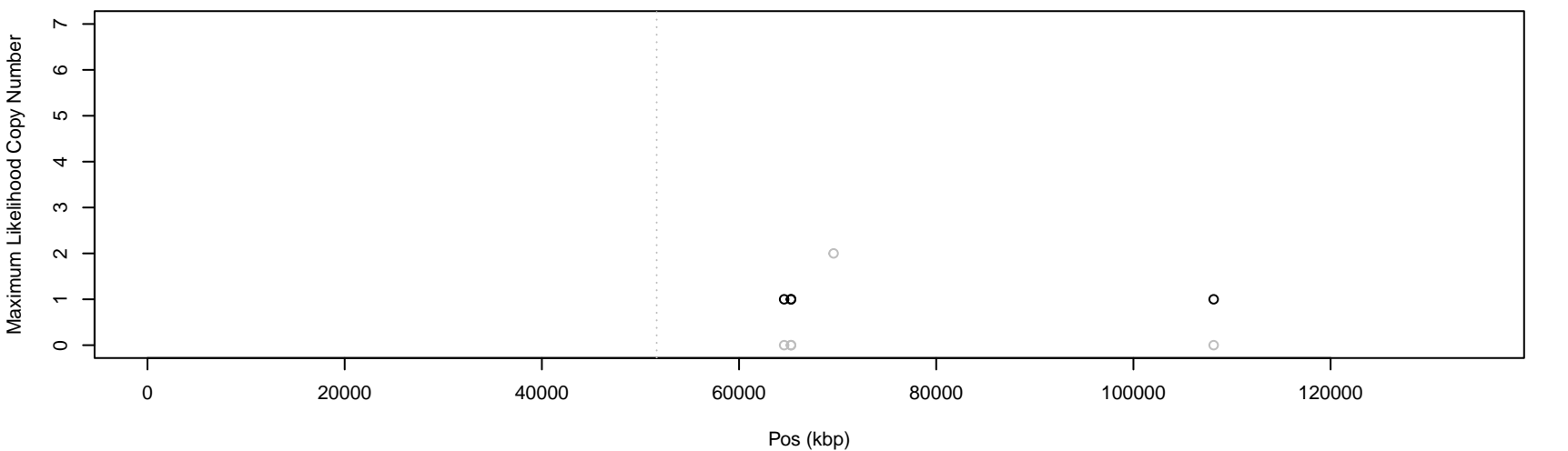
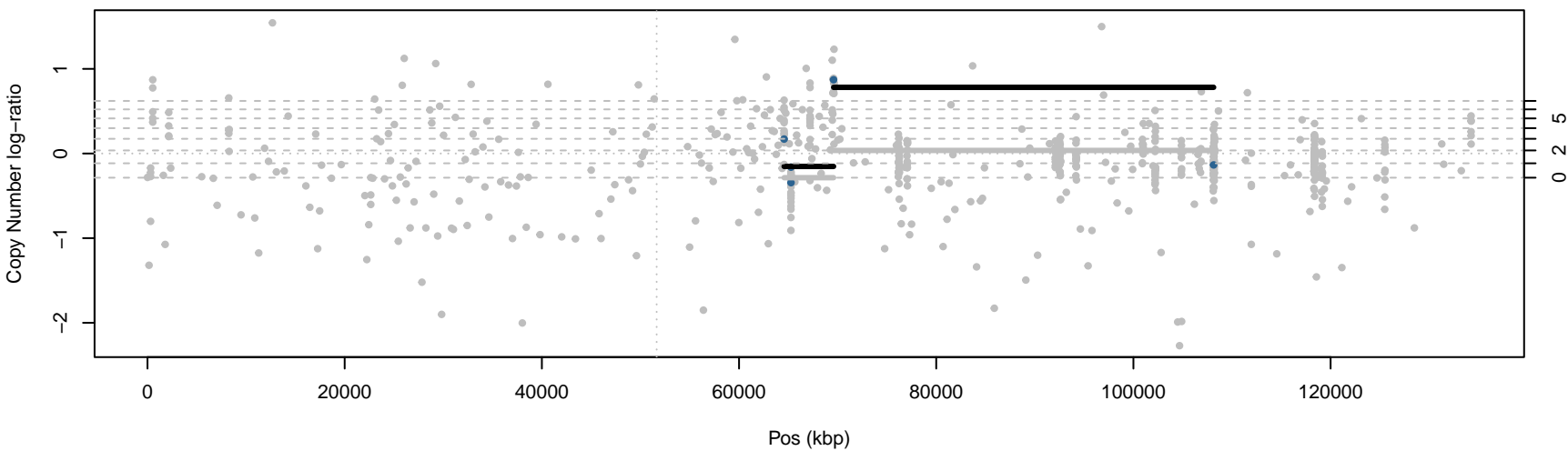
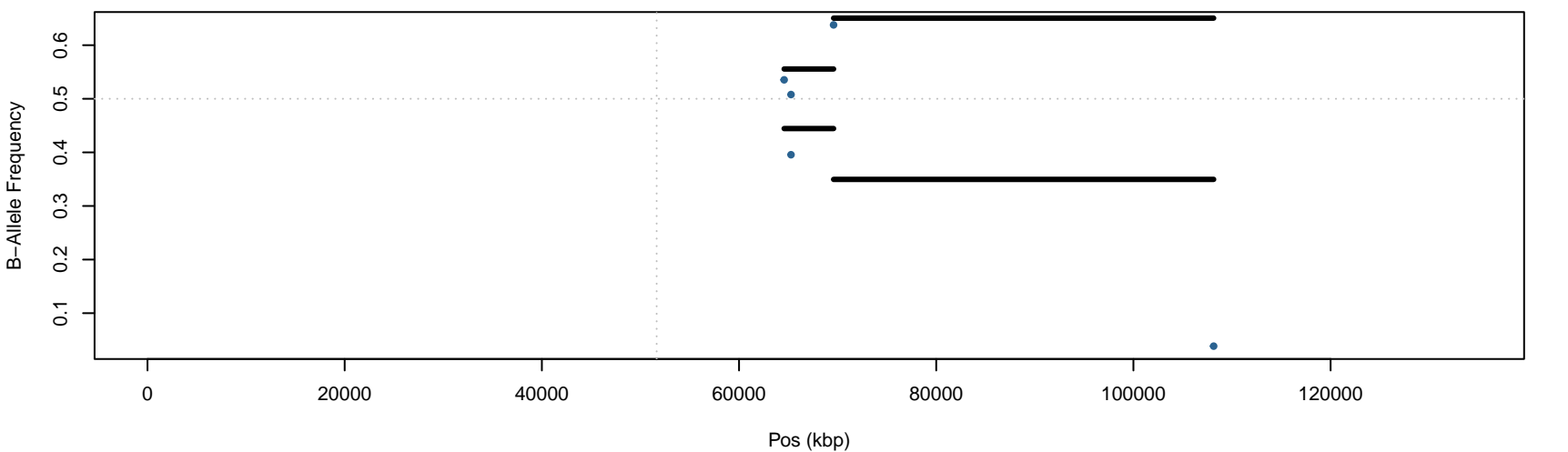




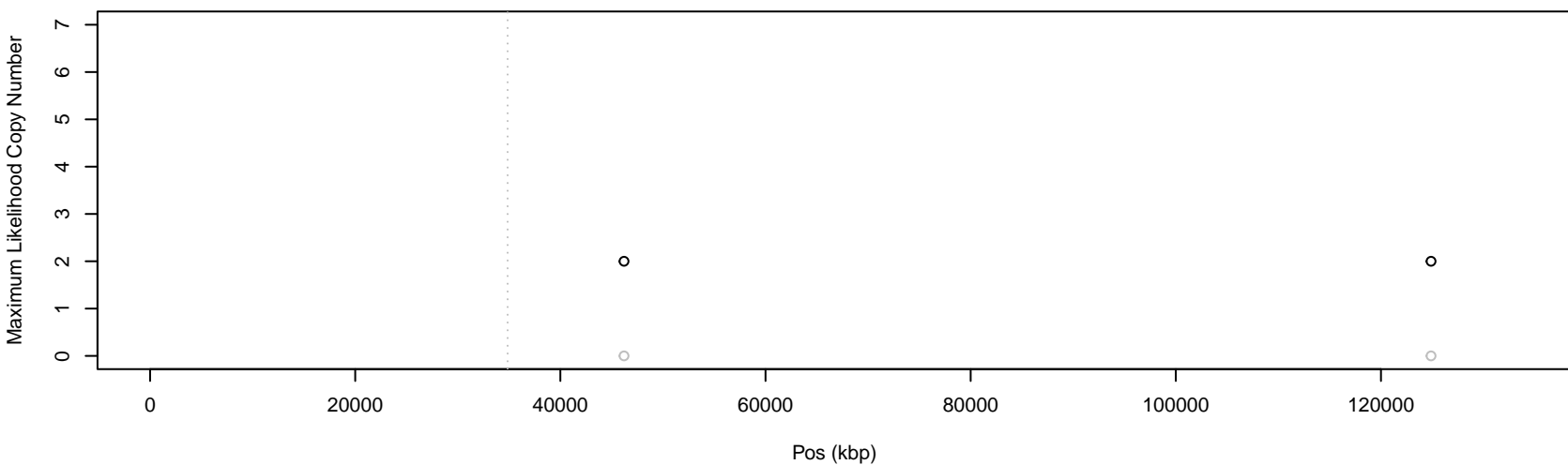
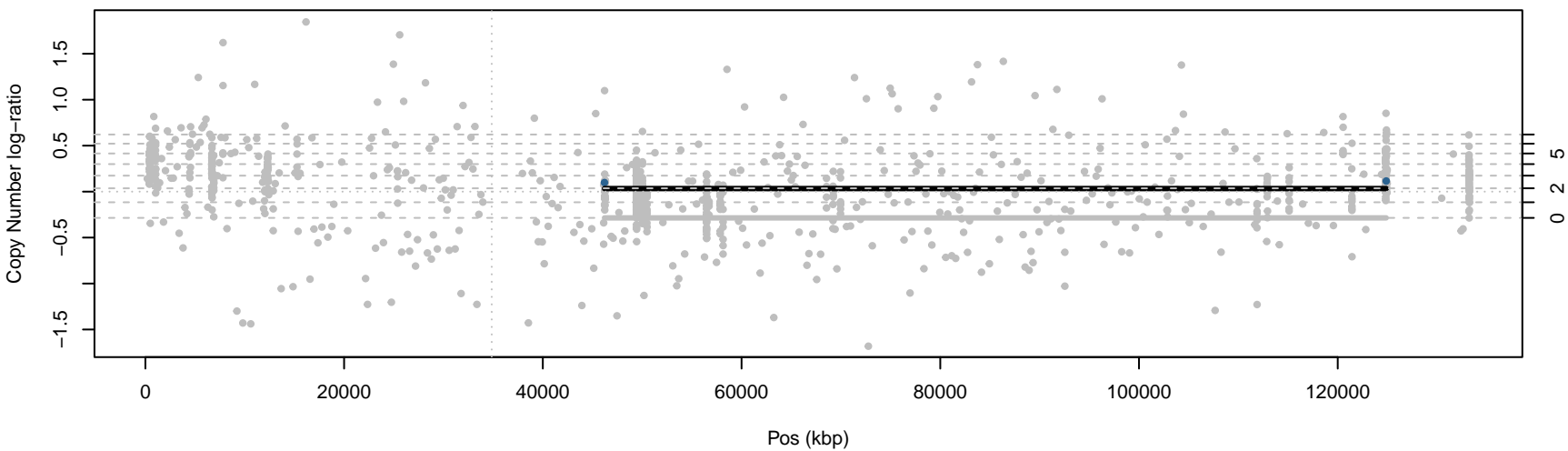
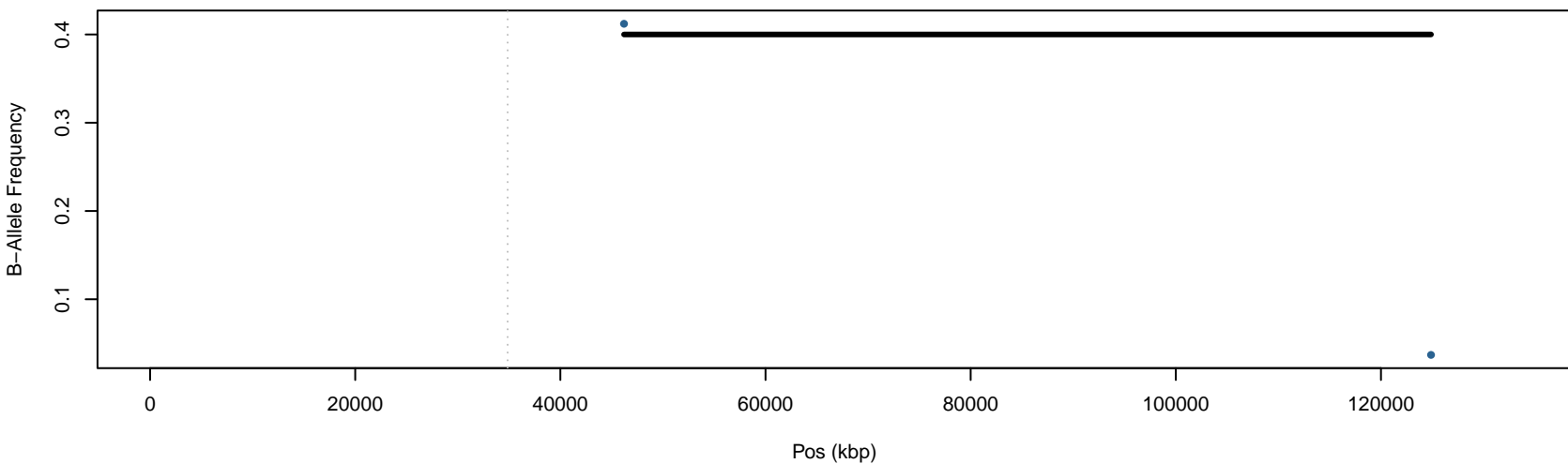
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 10



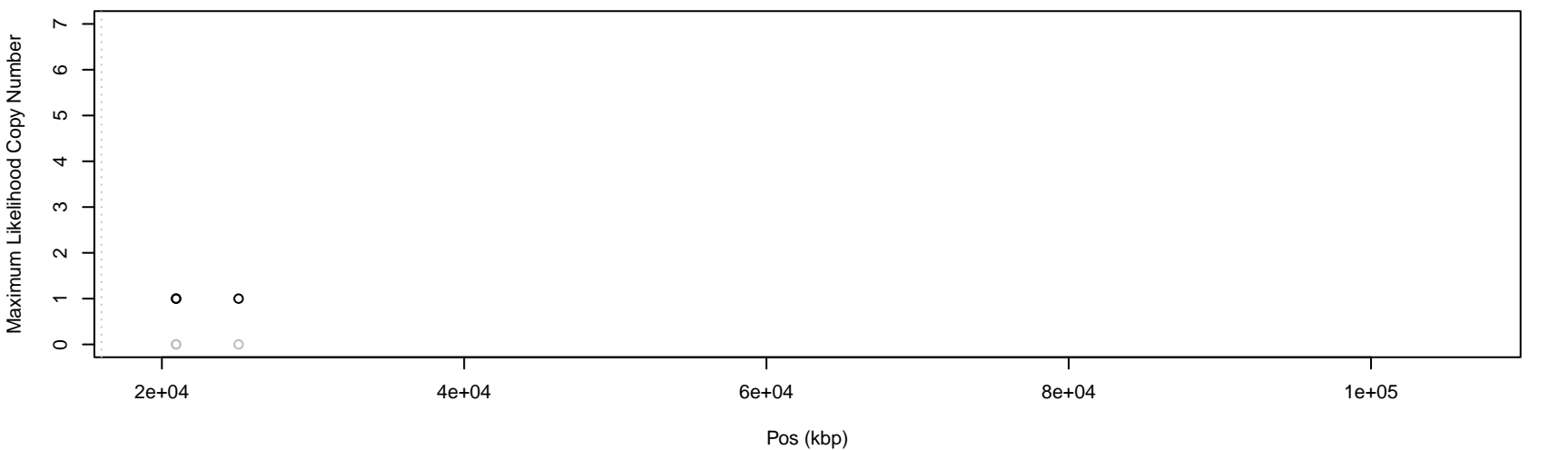
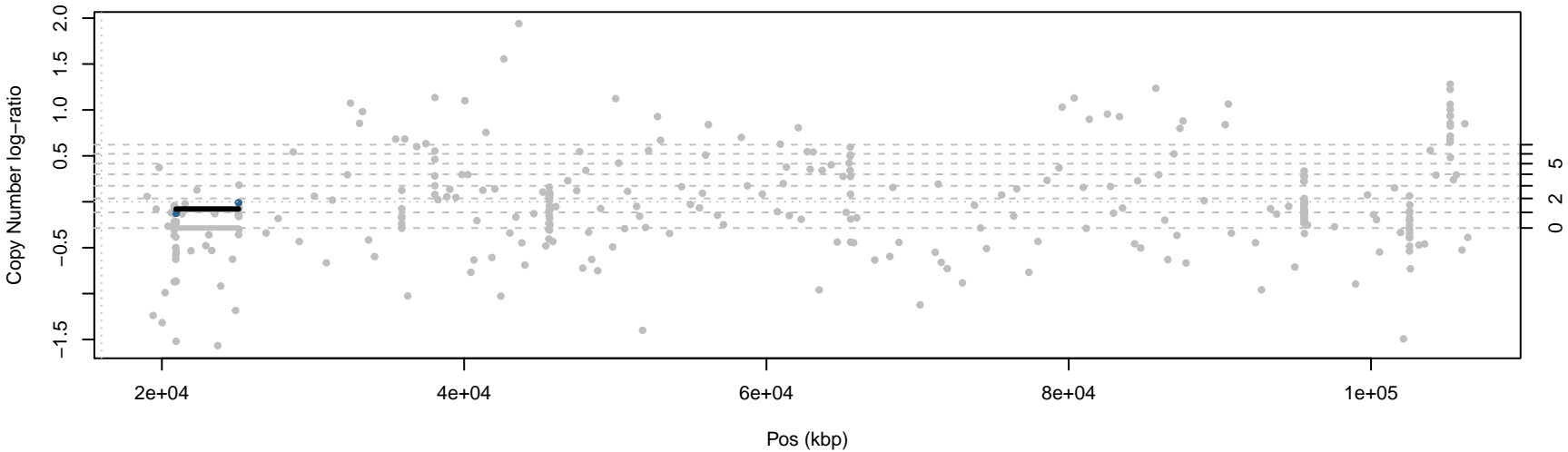
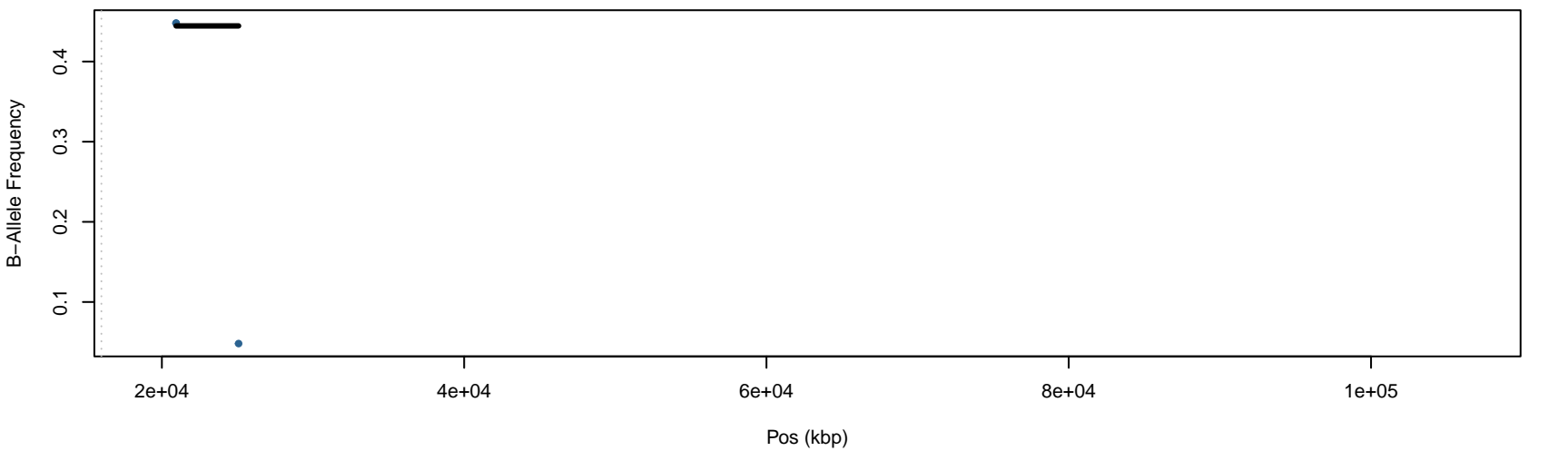
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 11



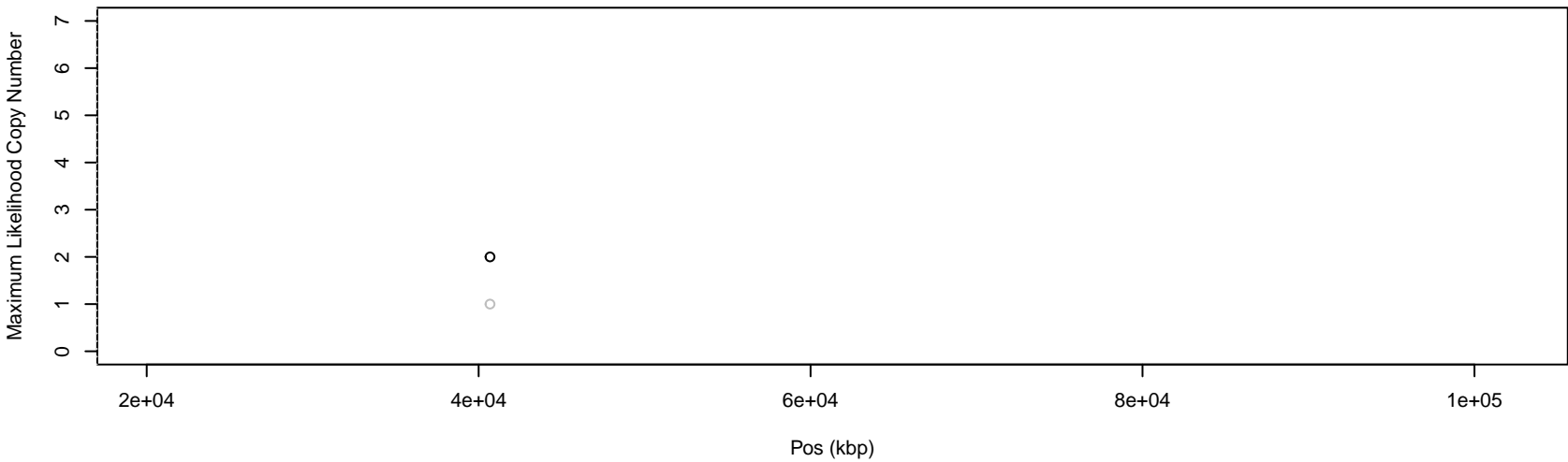
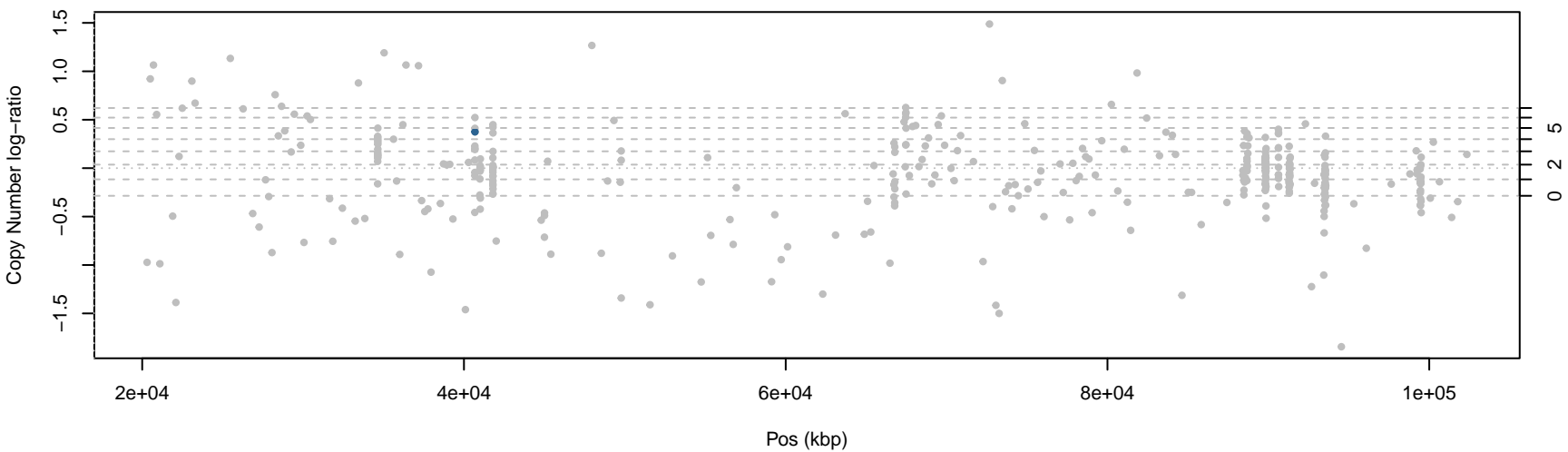
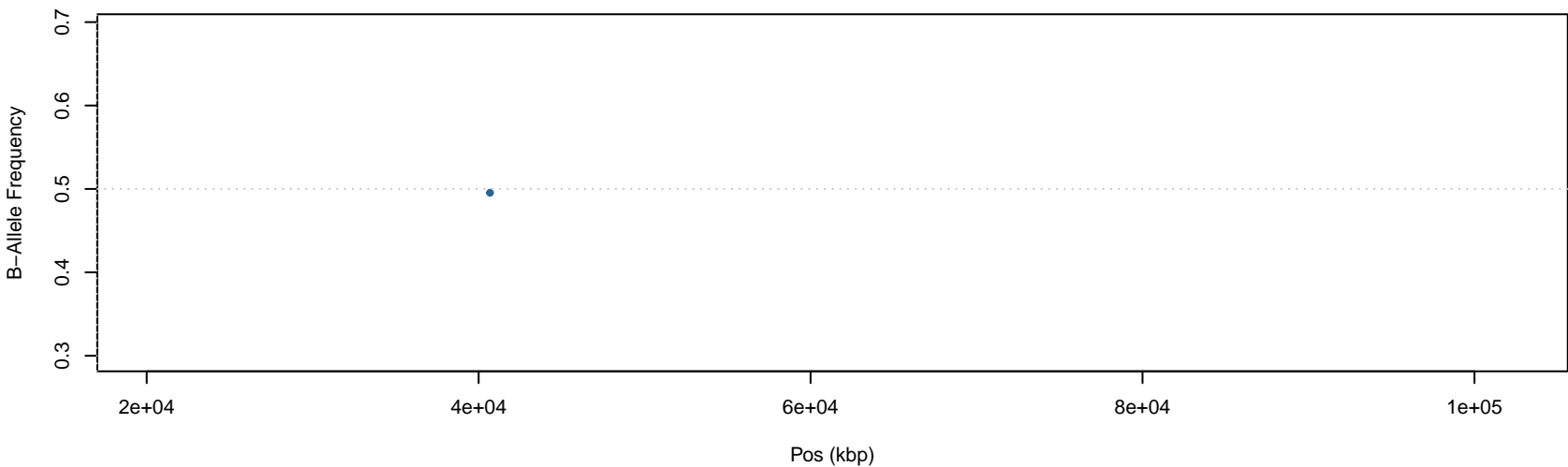
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 12



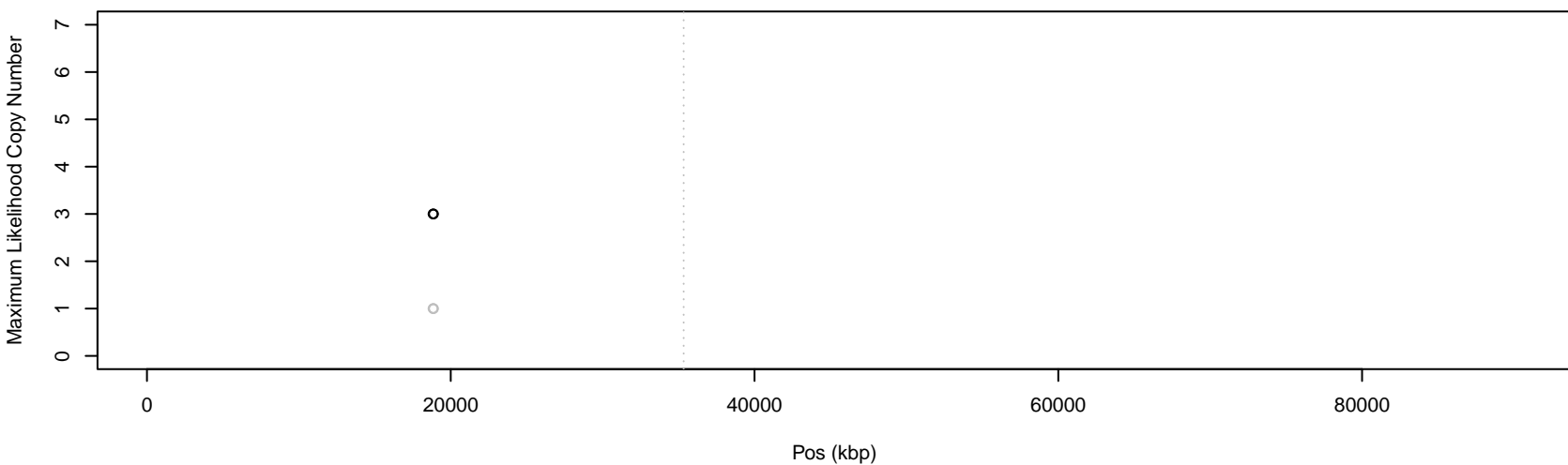
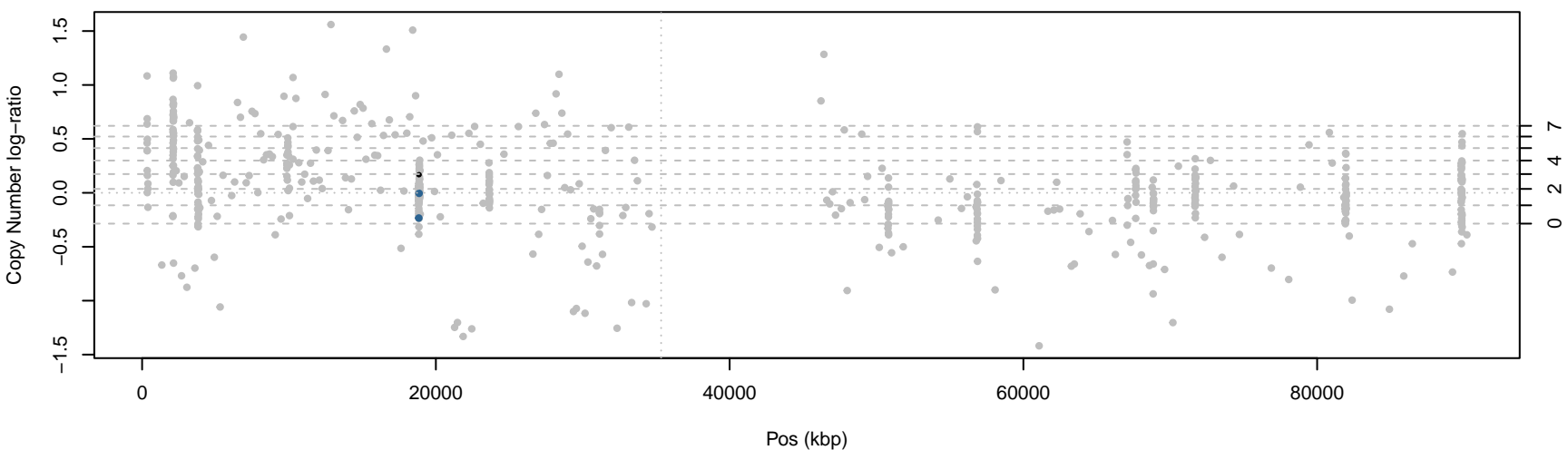
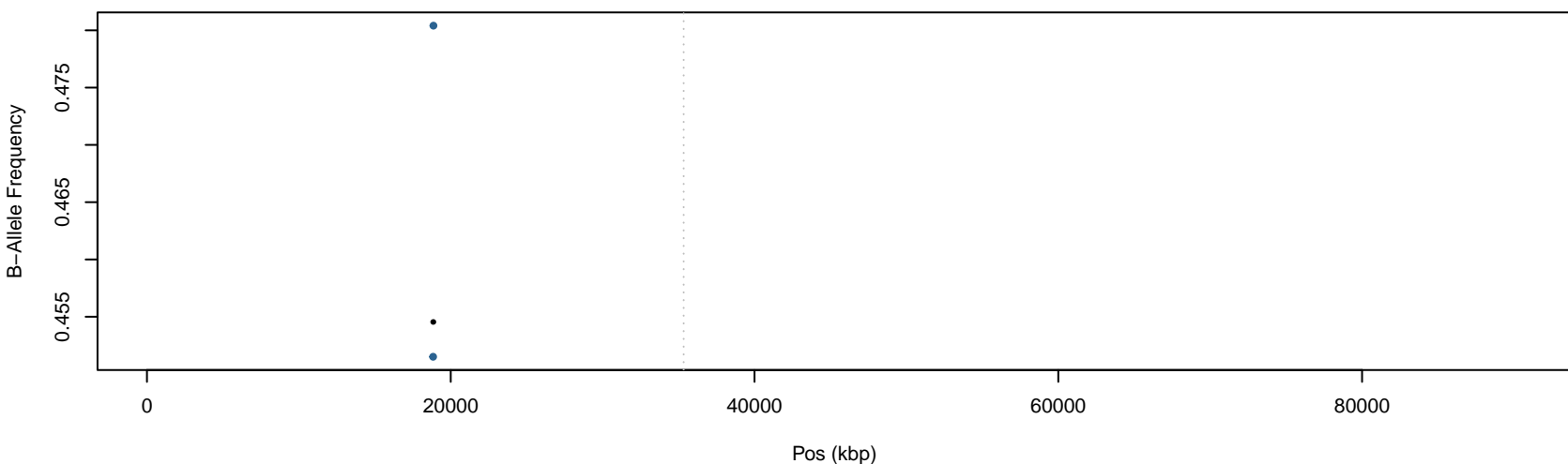
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 14



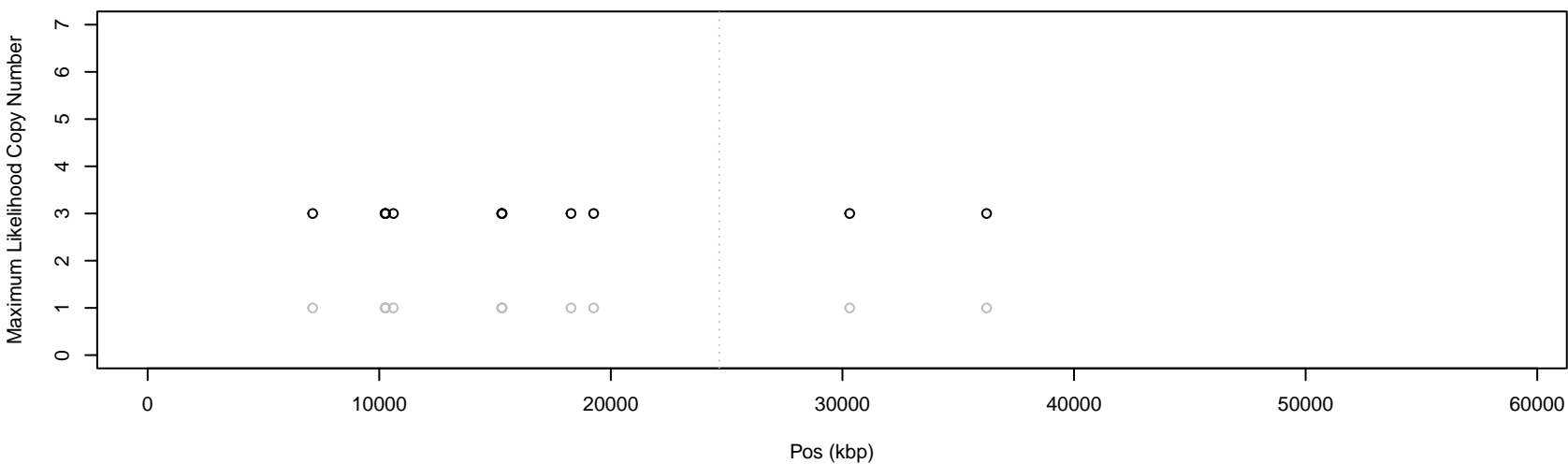
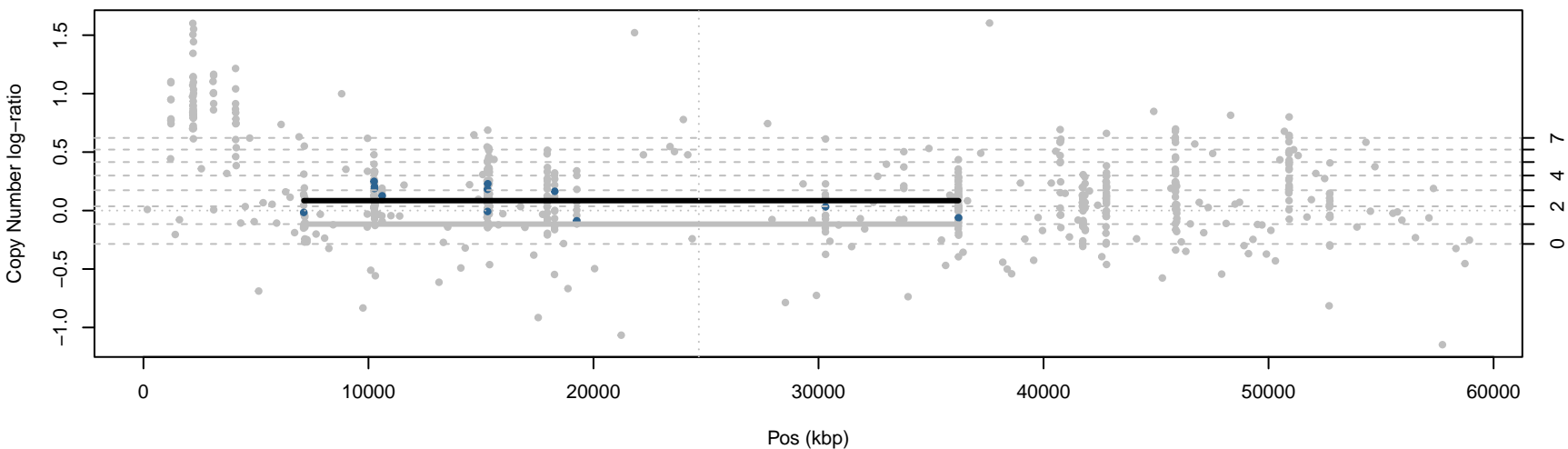
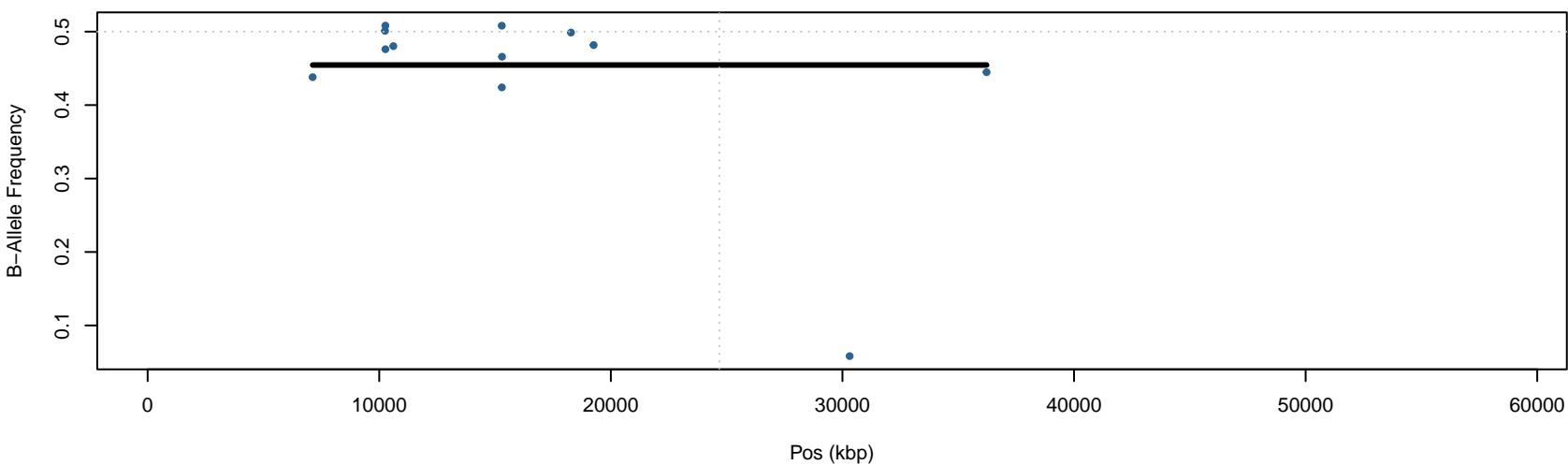
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 15



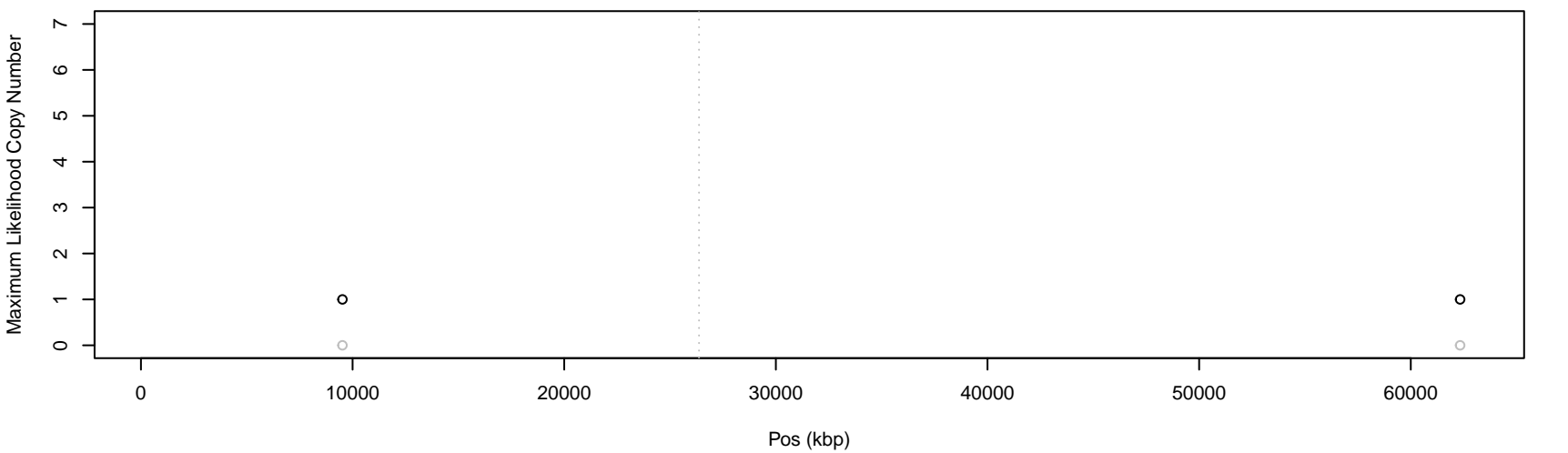
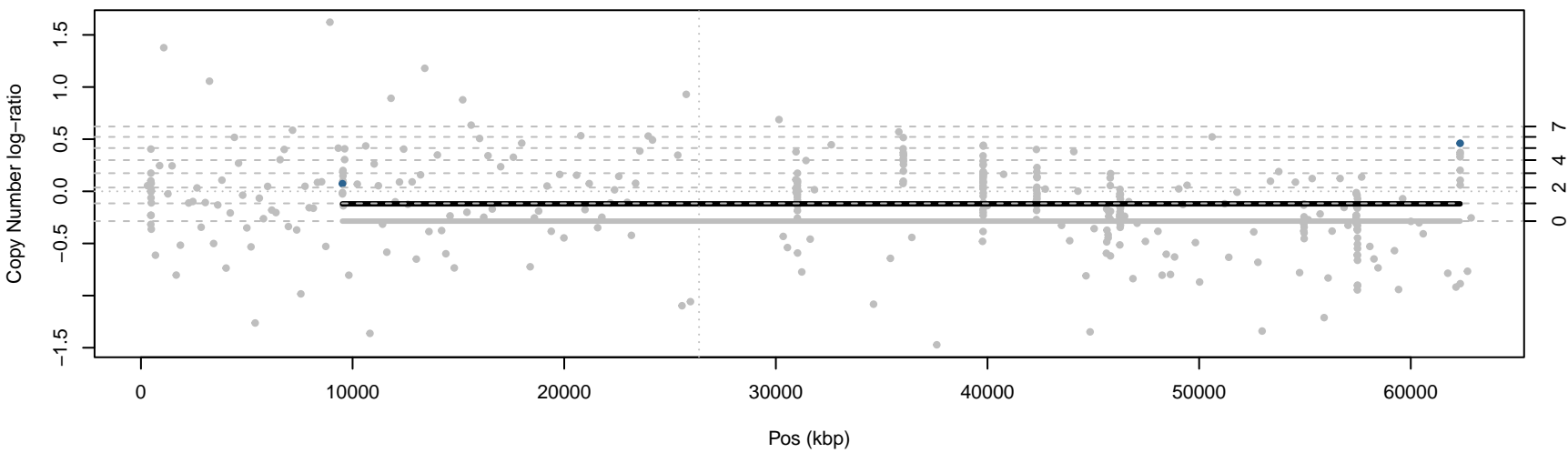
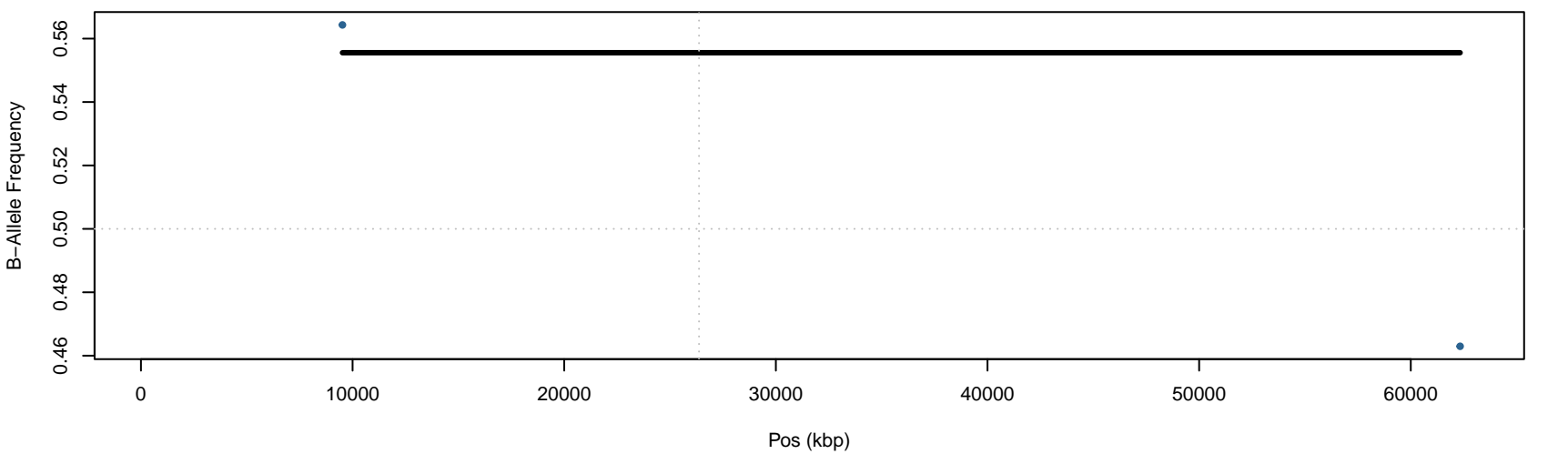
Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 16



Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 19

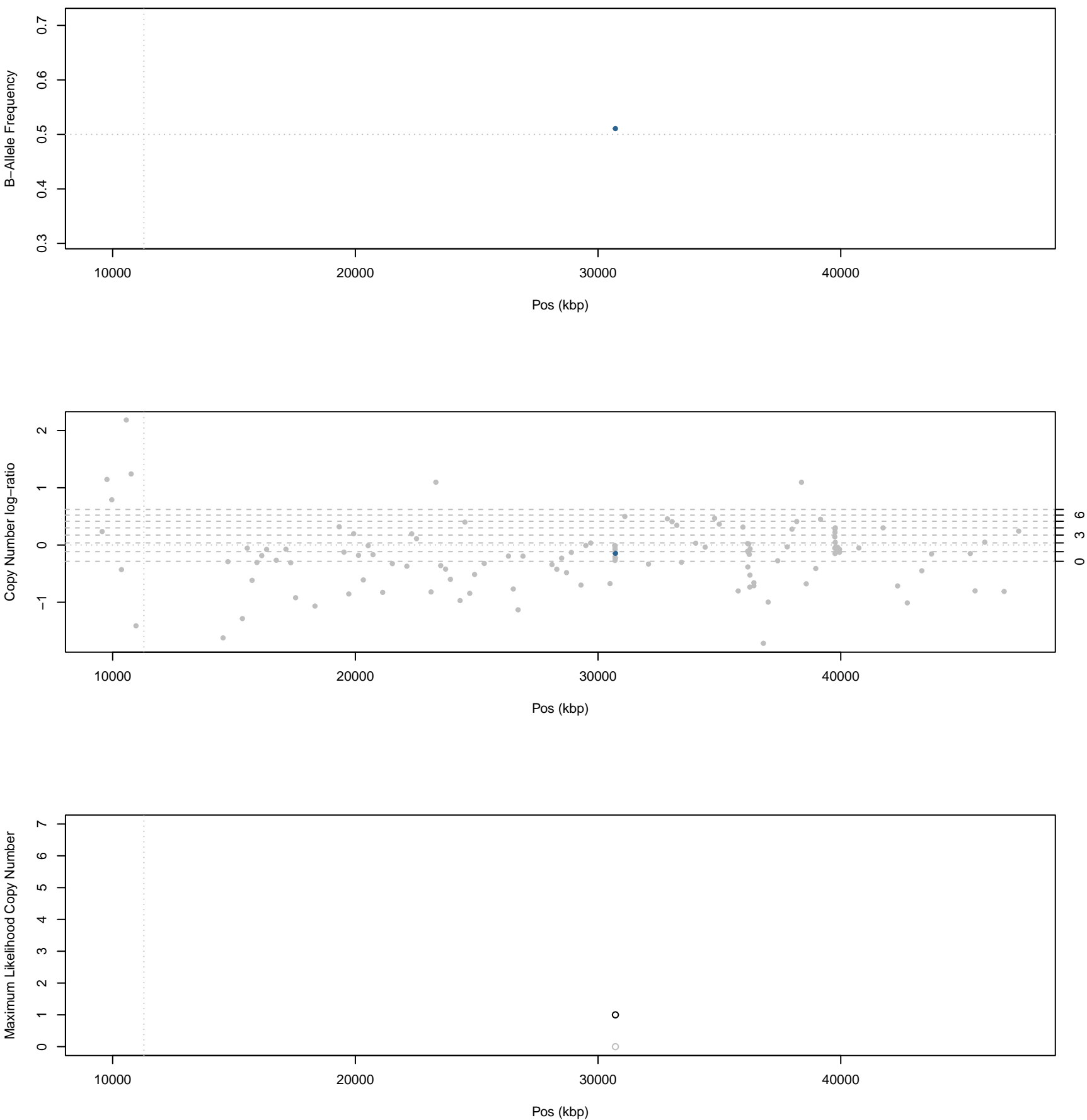


Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 20





Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 21



Purity: 0.2 Tumor ploidy: 1.752 SNV log-likelihood: 40.73 GoF: 94.1% Mean coverage: 520;712 Chromosome: 22

