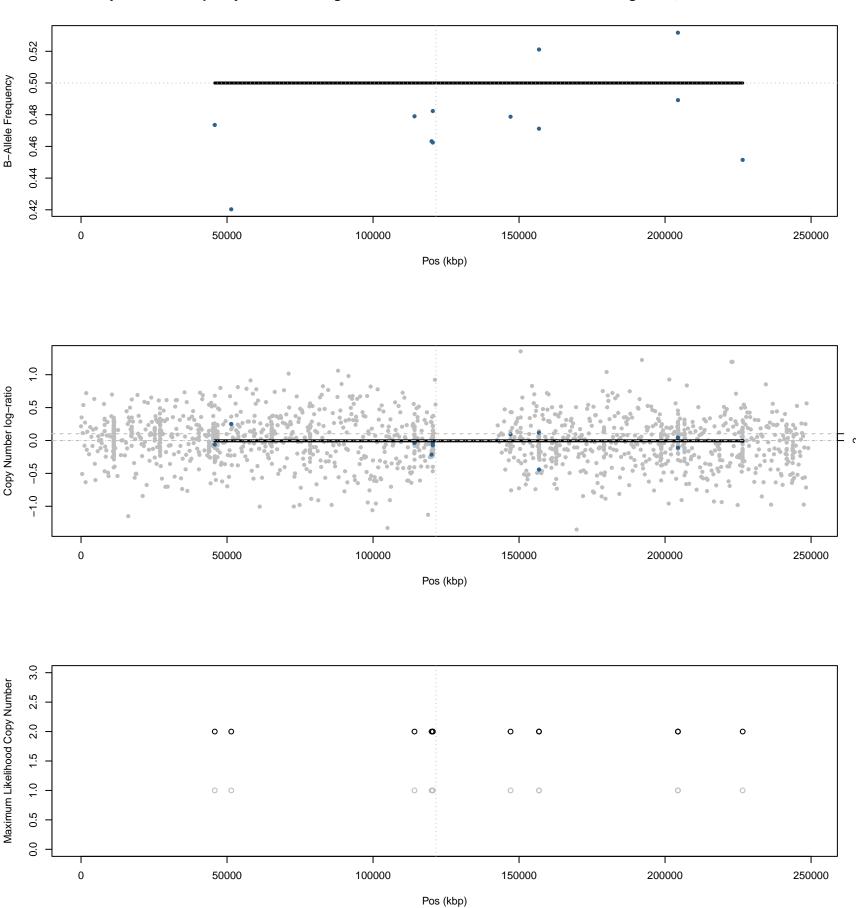
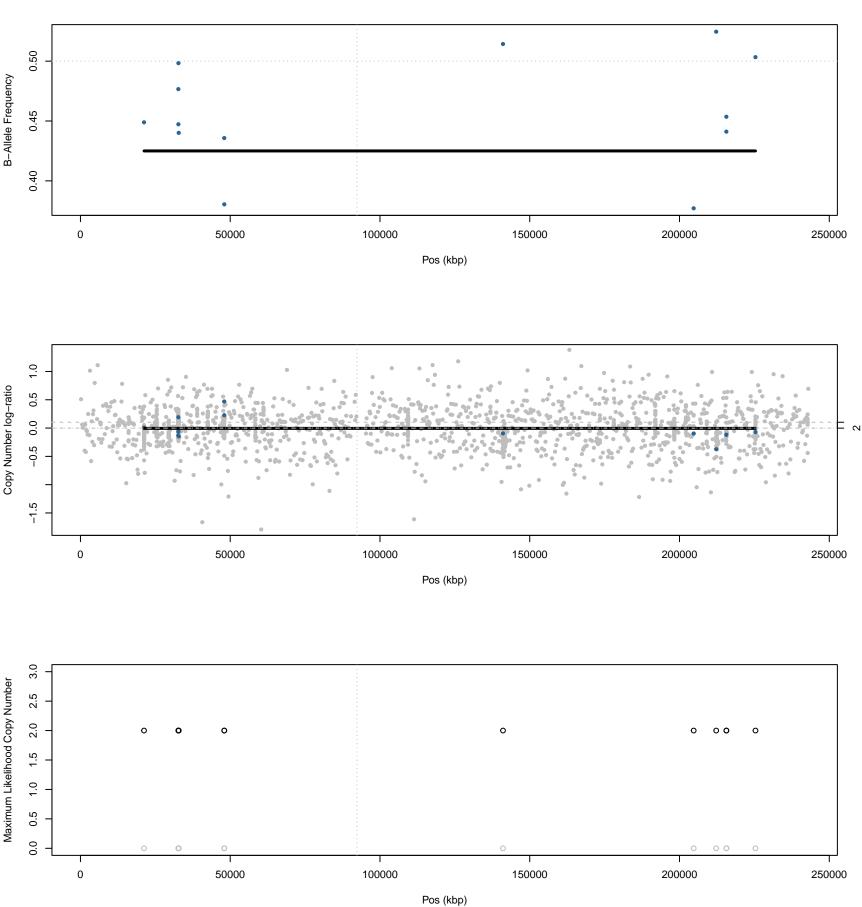
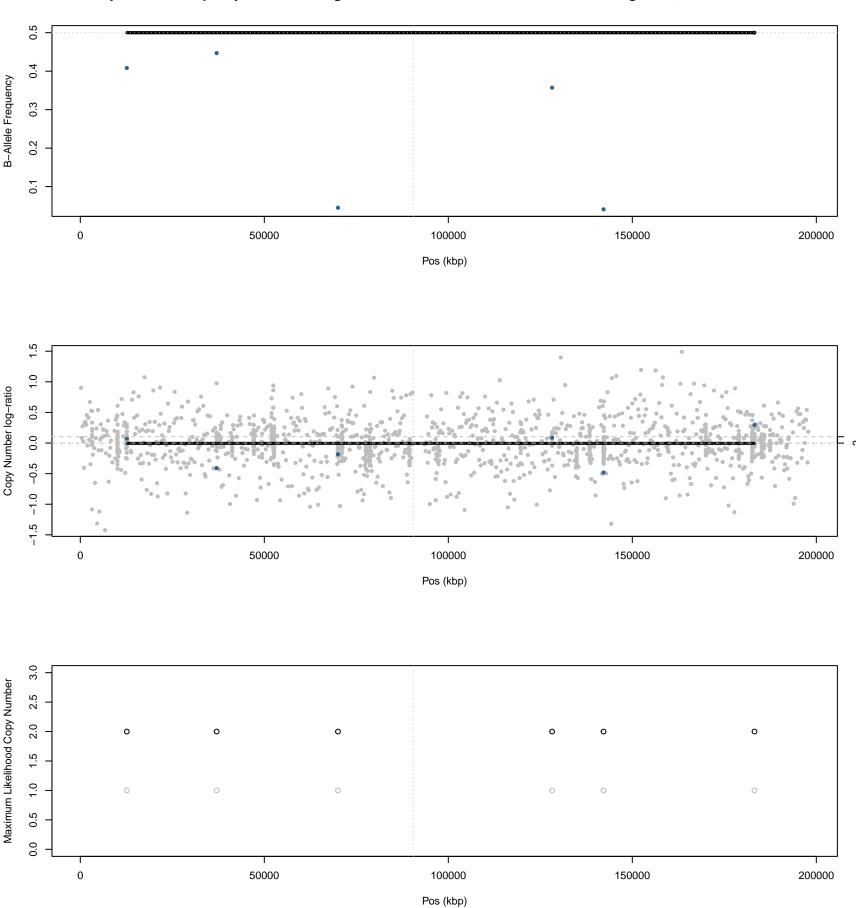
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 1



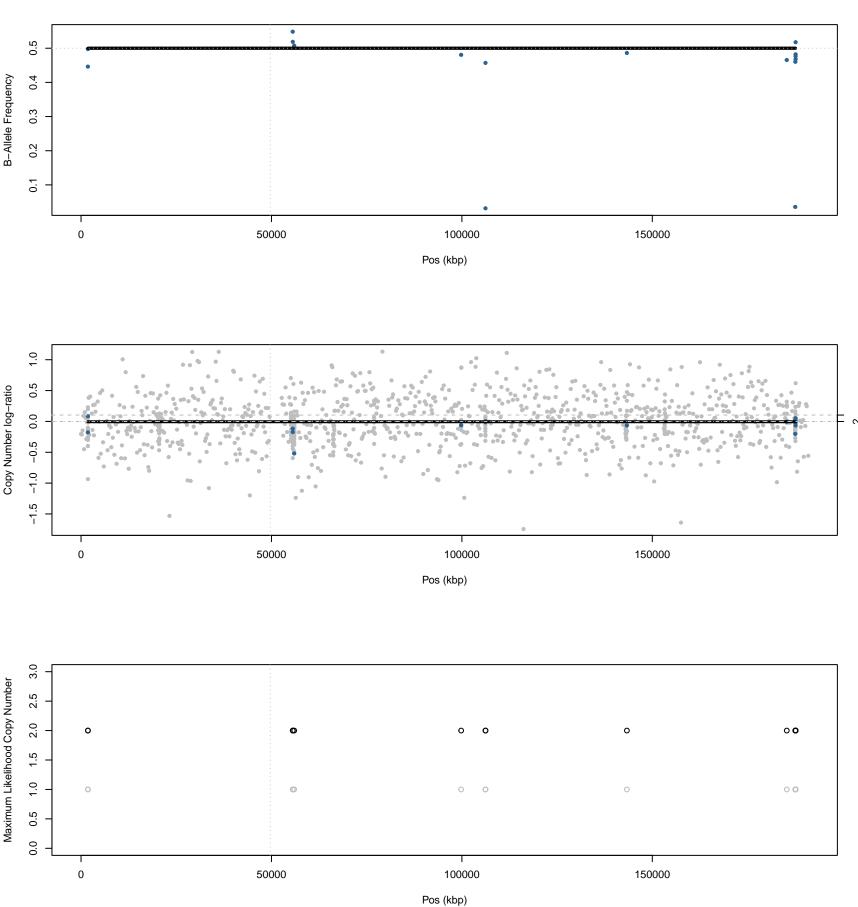
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 2



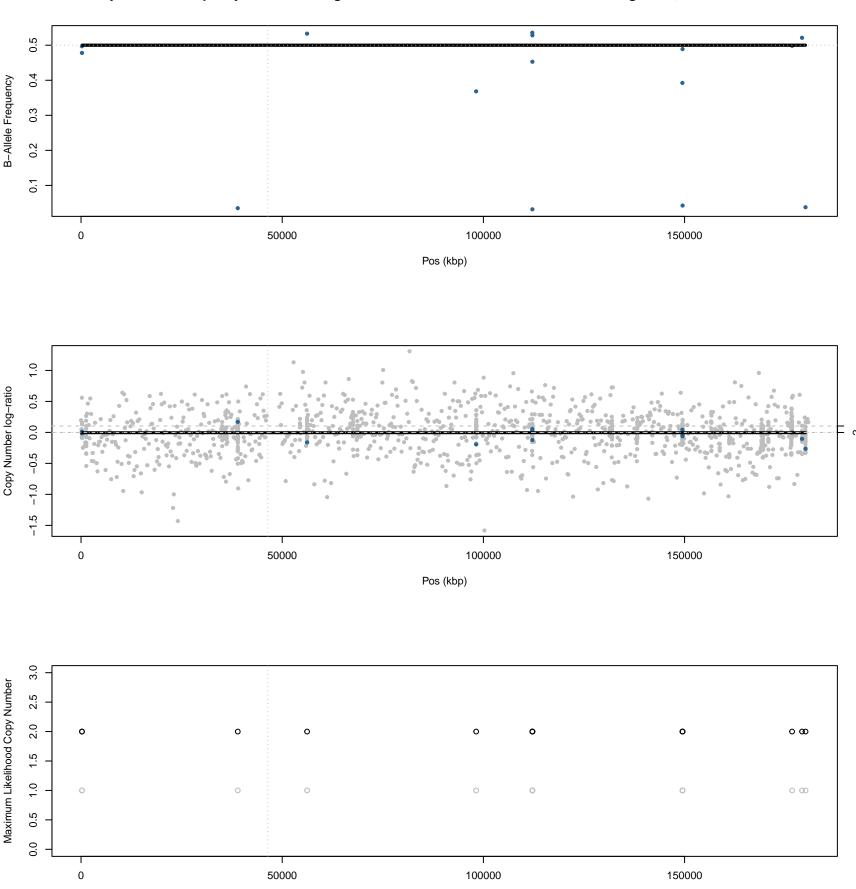
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 3



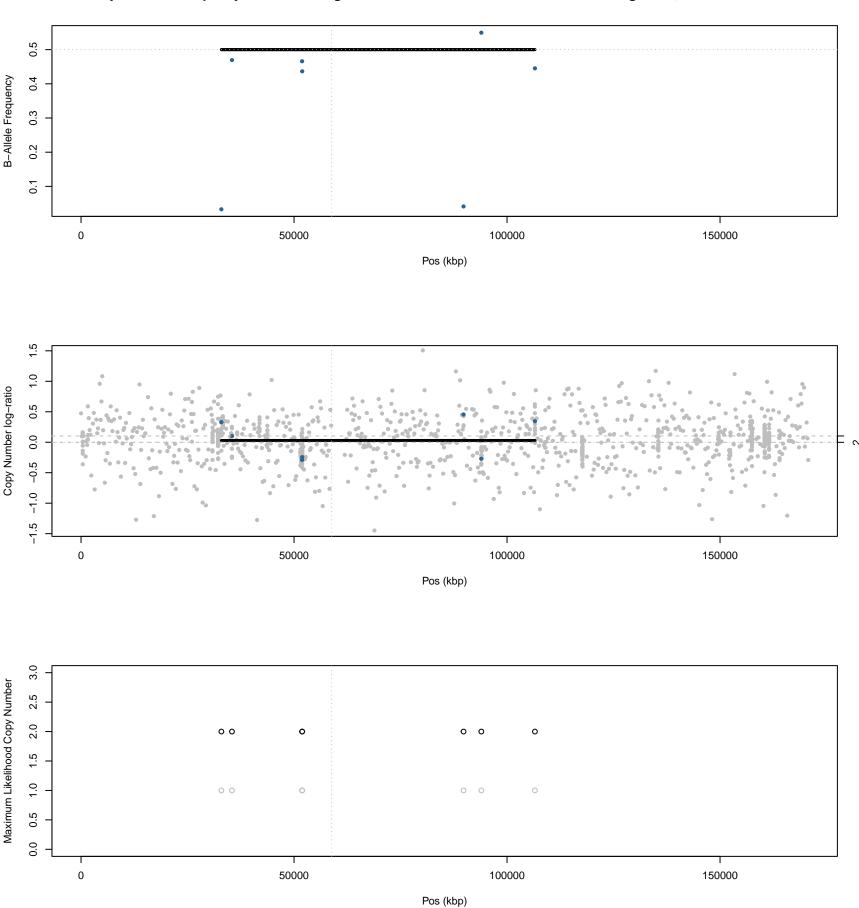
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 4



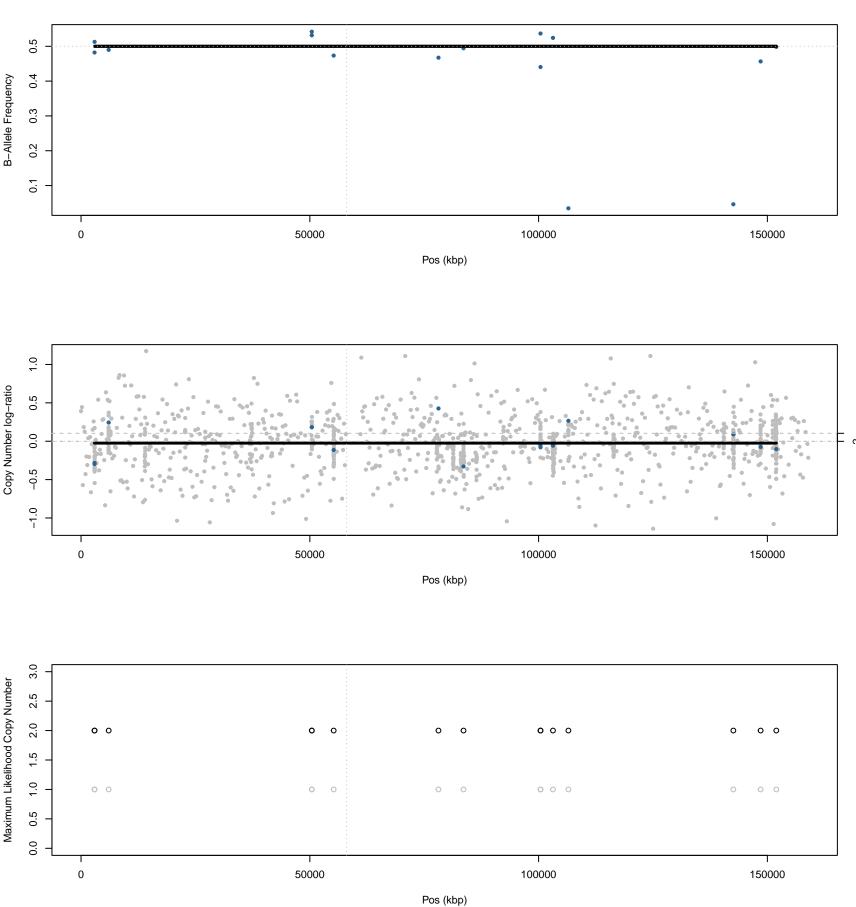
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 5



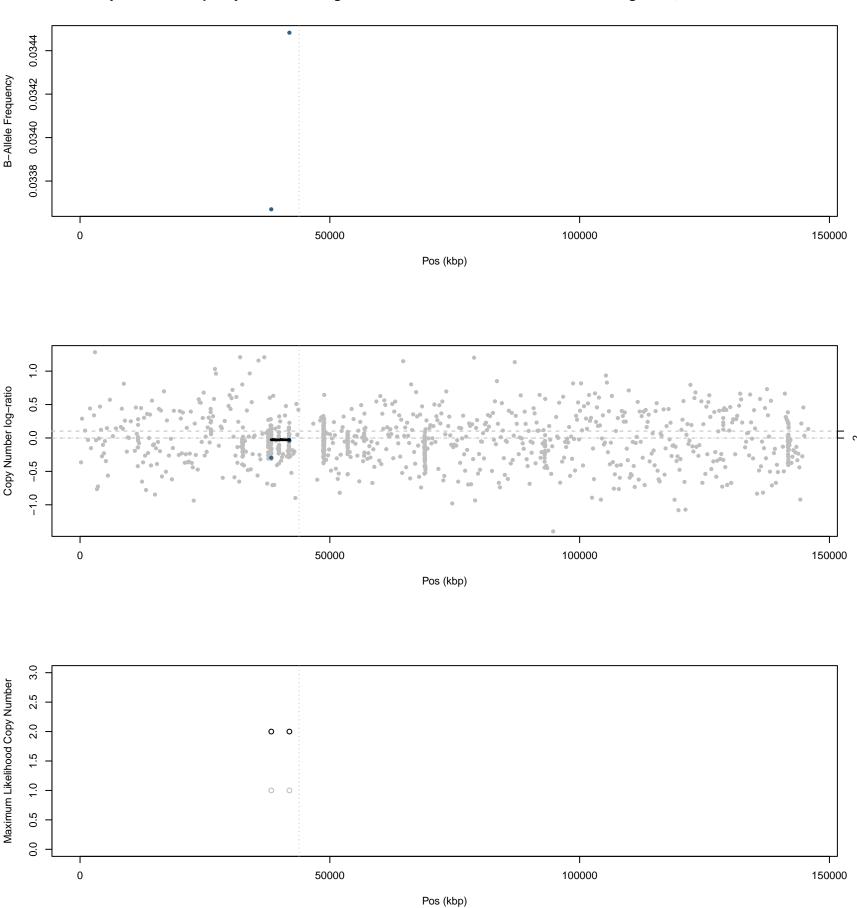
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 6



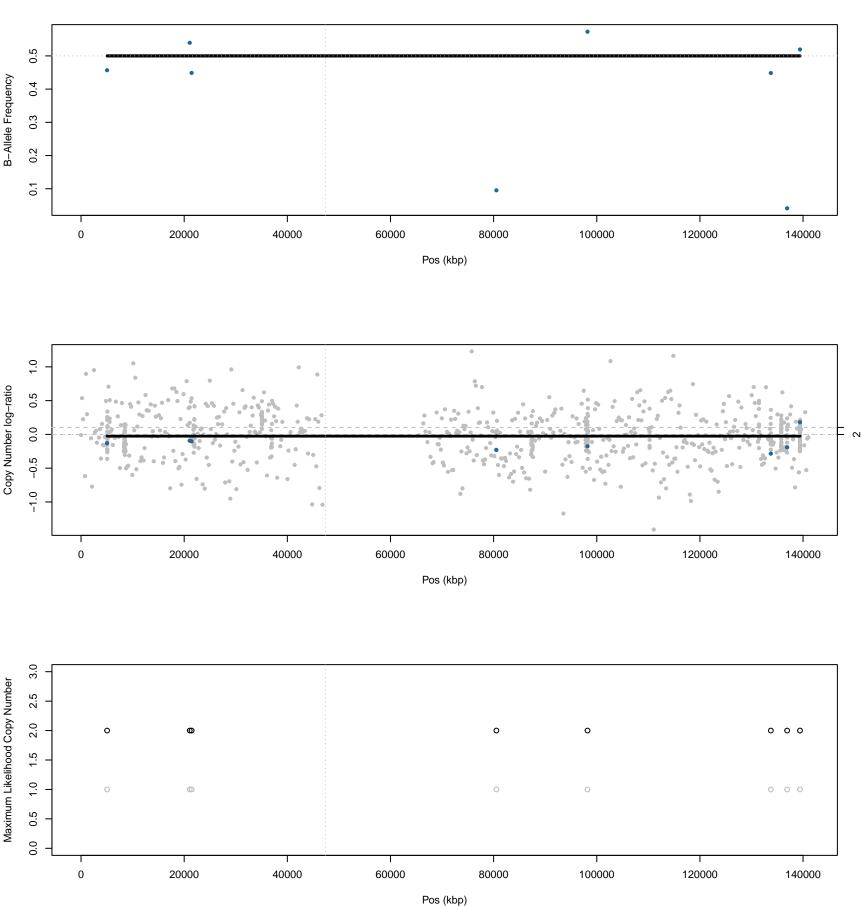
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 7



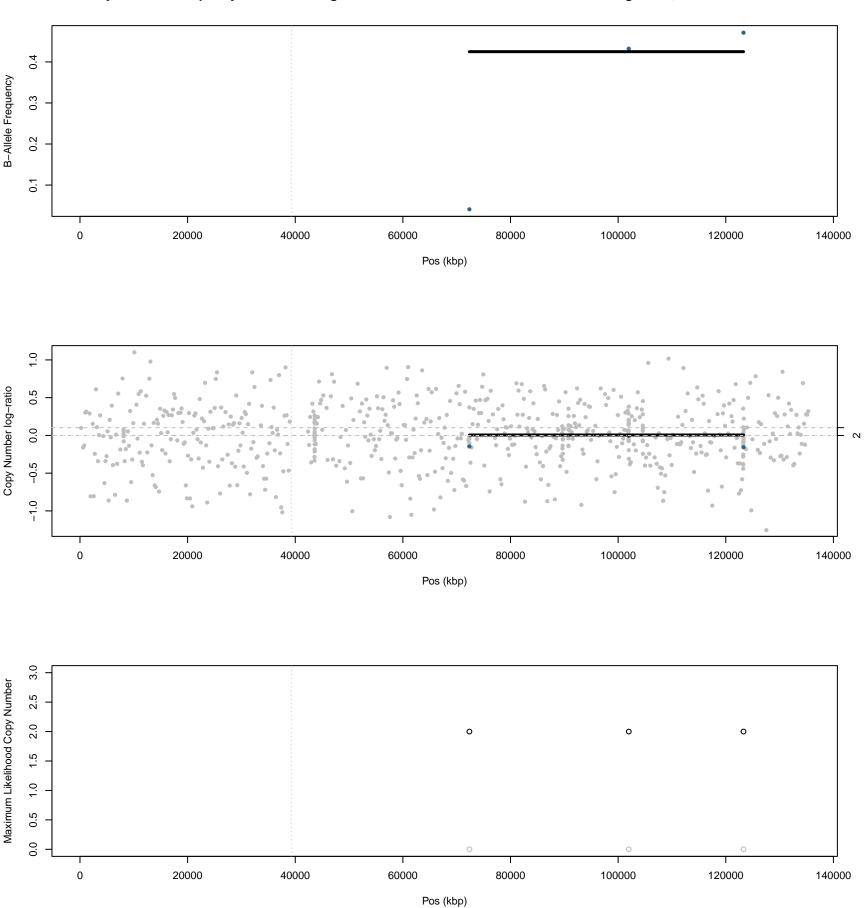
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 8



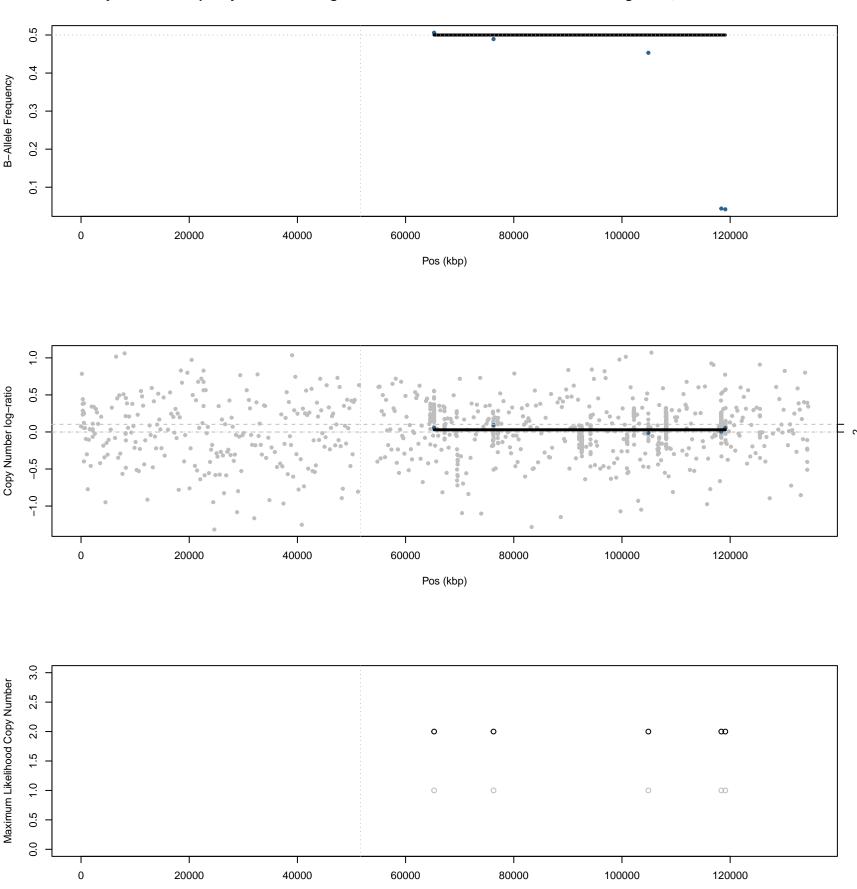
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 9



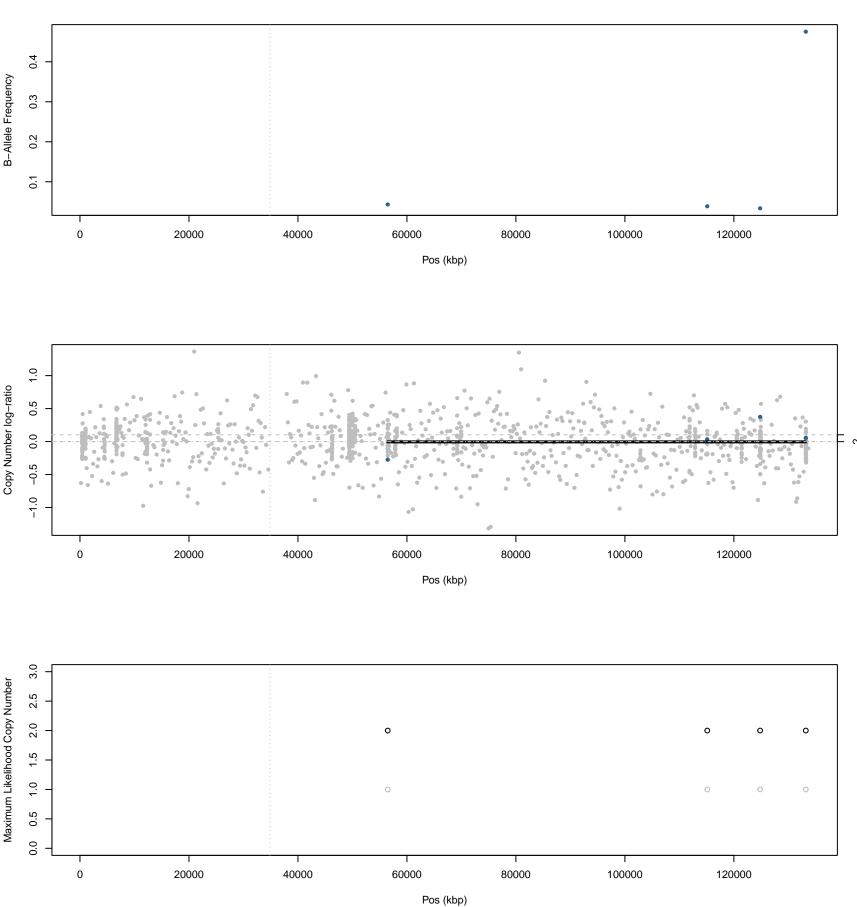
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 10



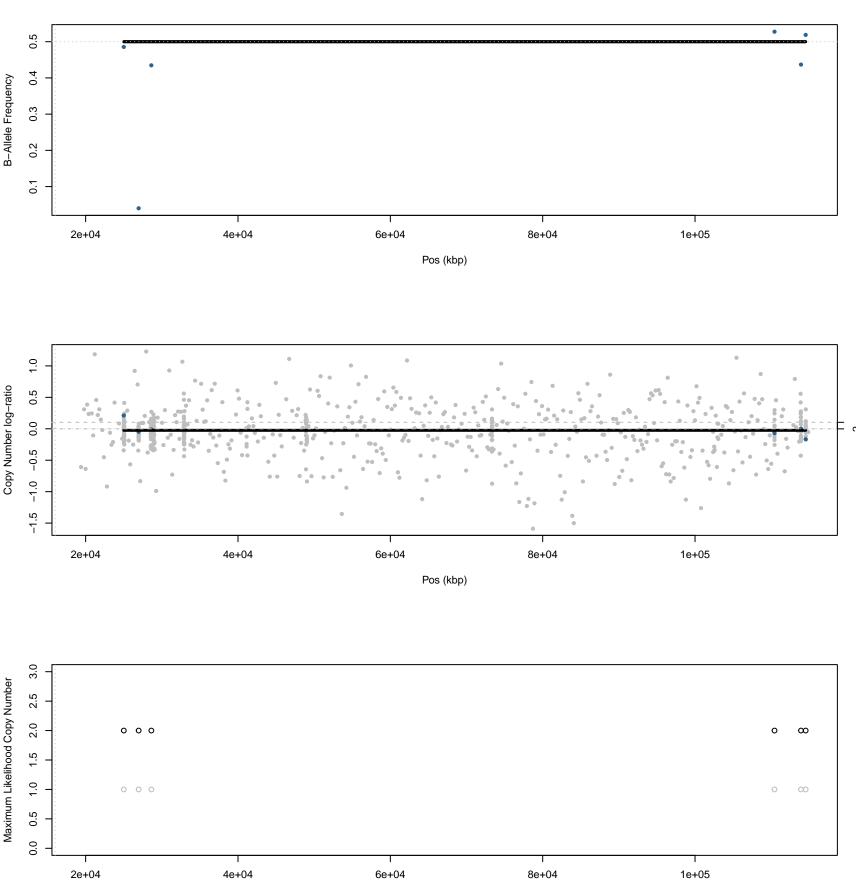
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 11



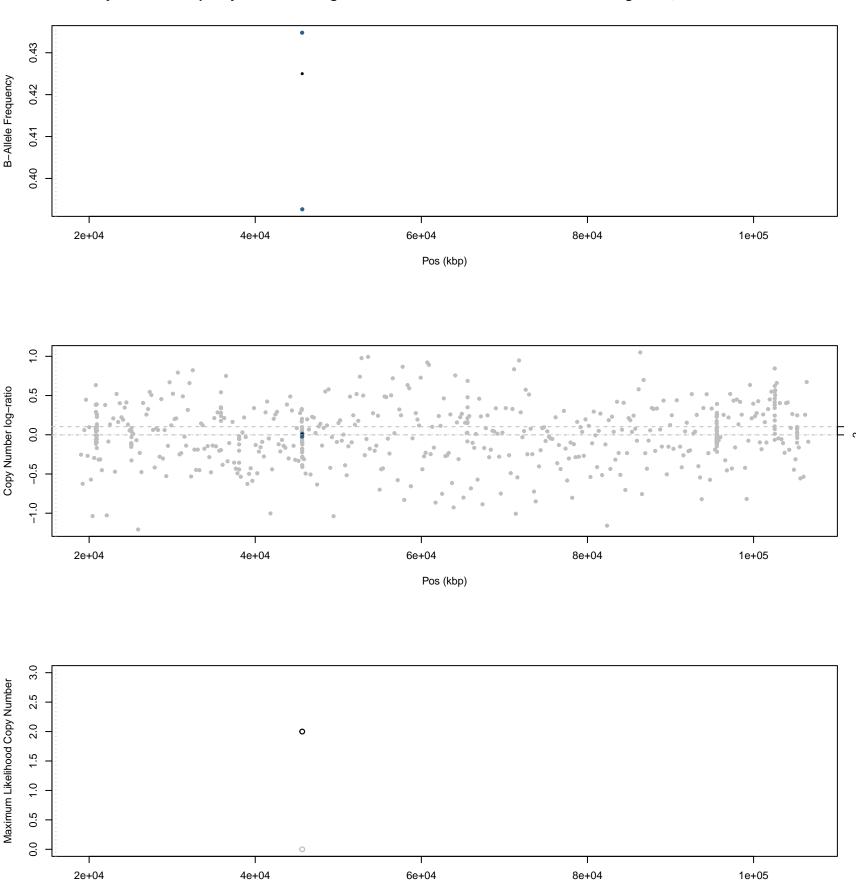
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 12



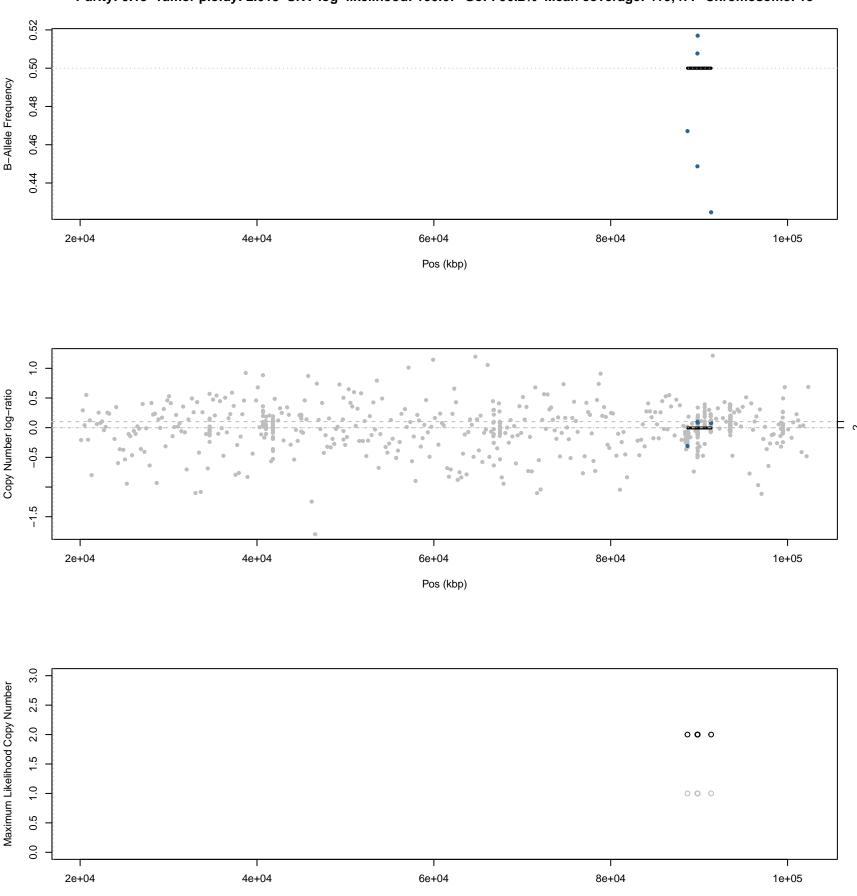
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 13



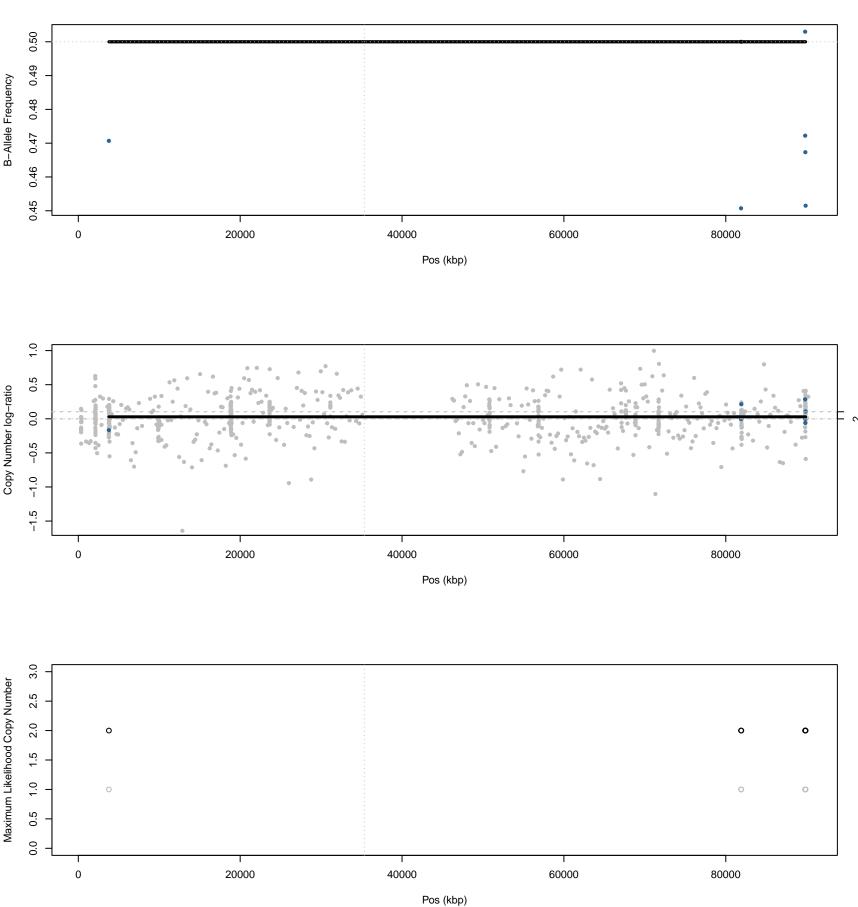
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 14



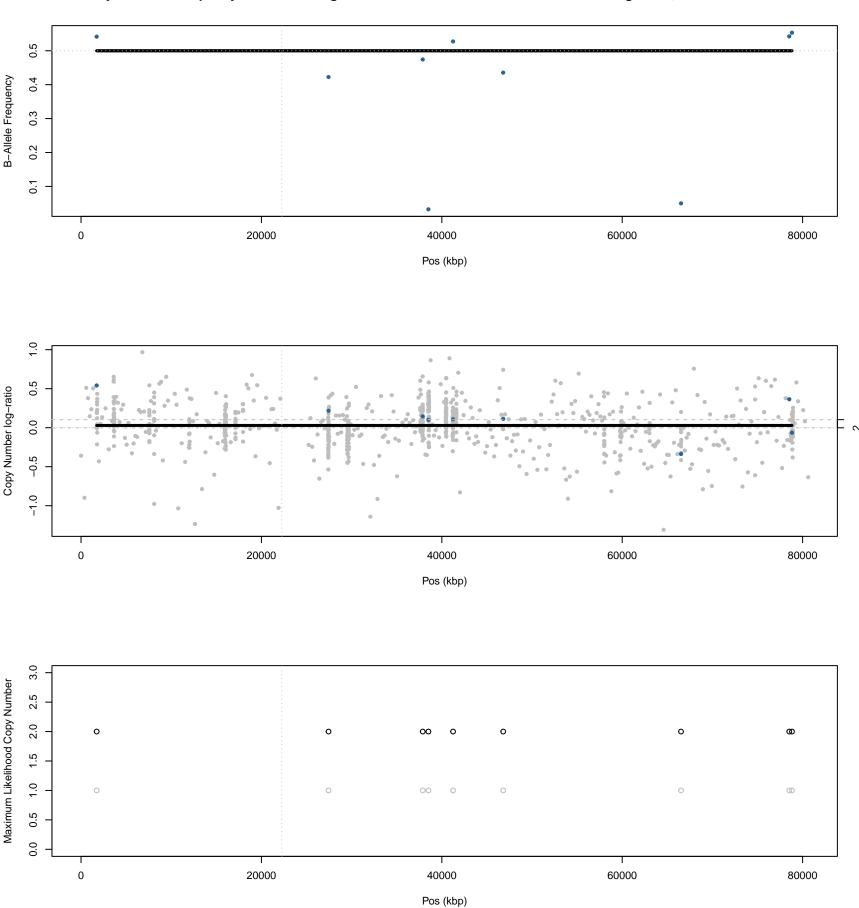
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 15



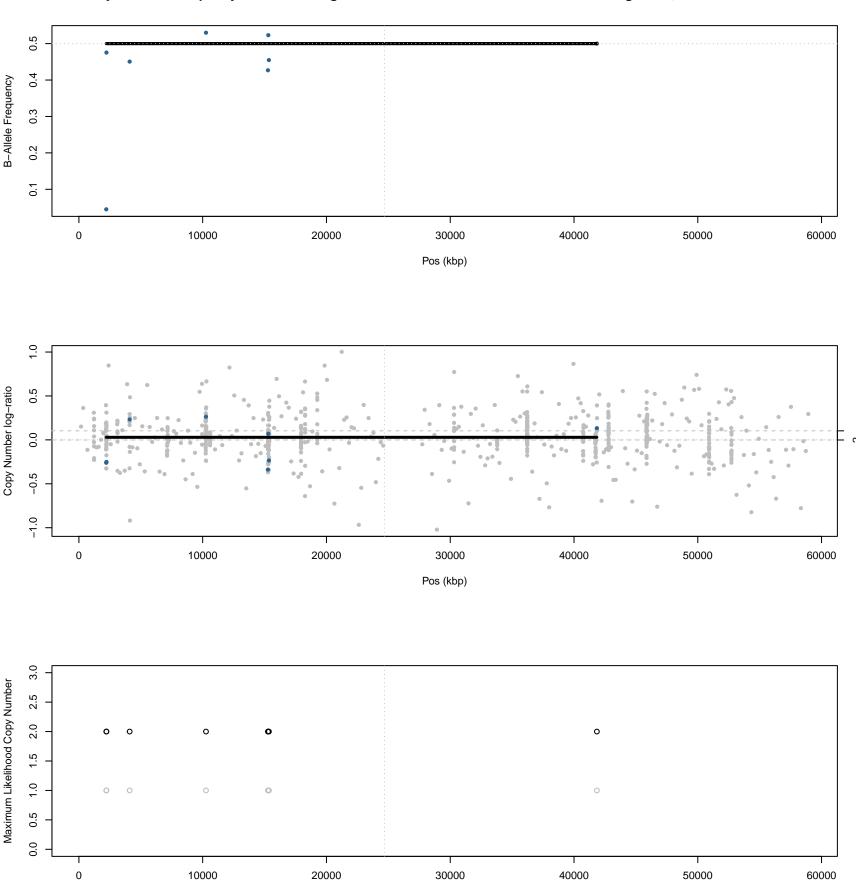
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 16



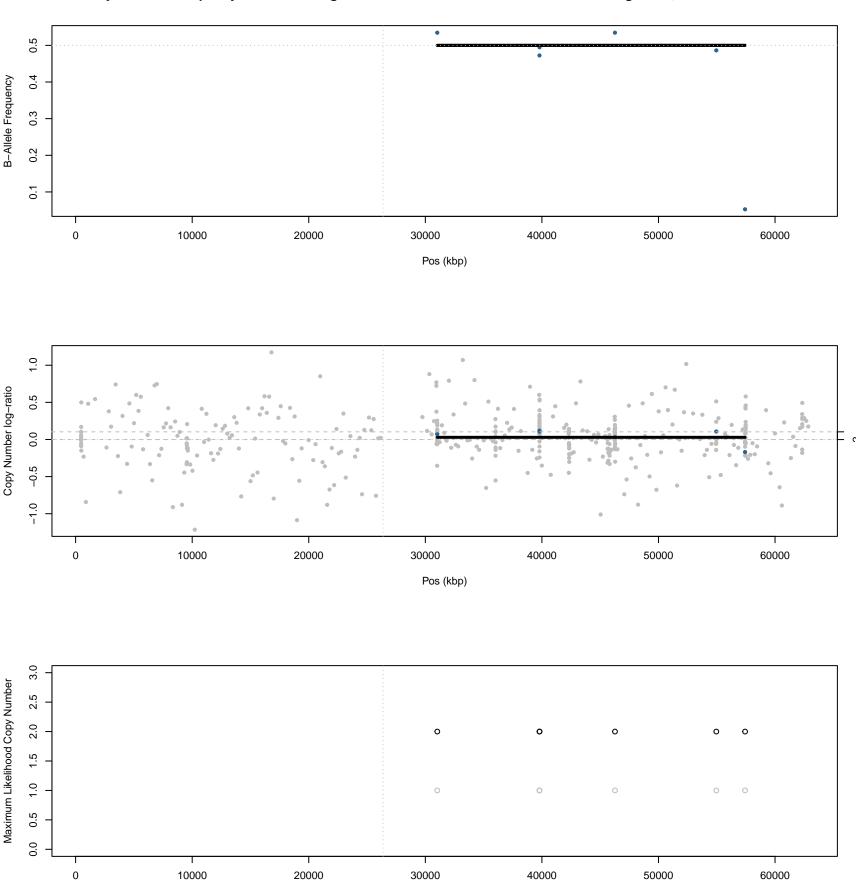
Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 17



Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 19



Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 20



Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477 Chromosome: 22

