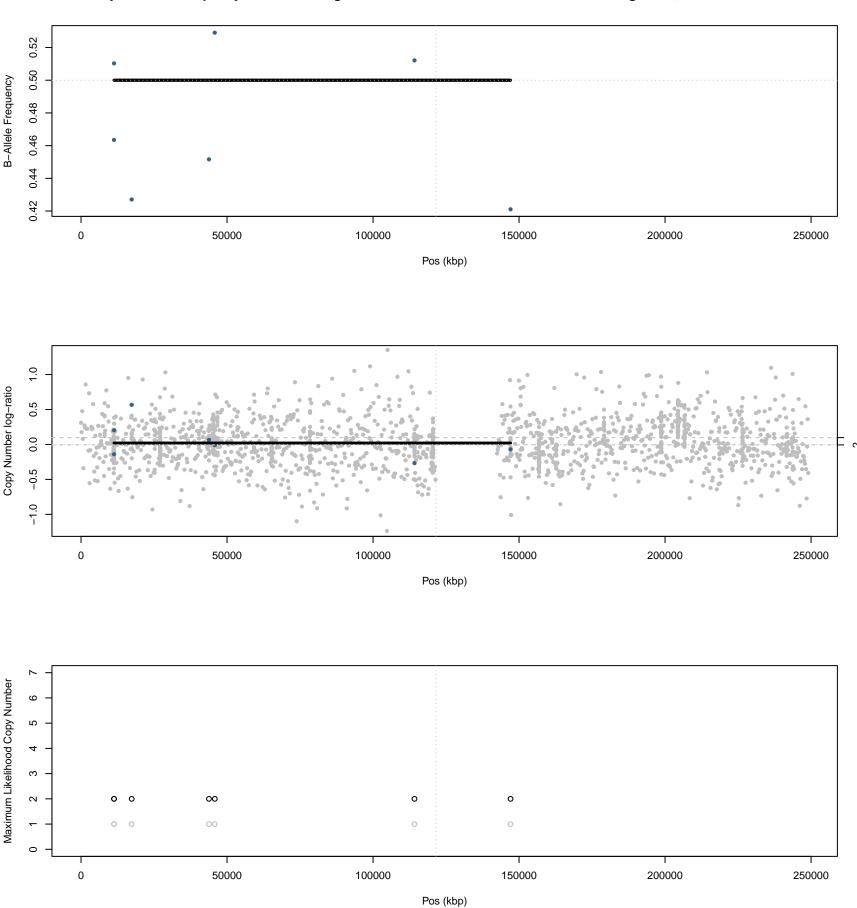
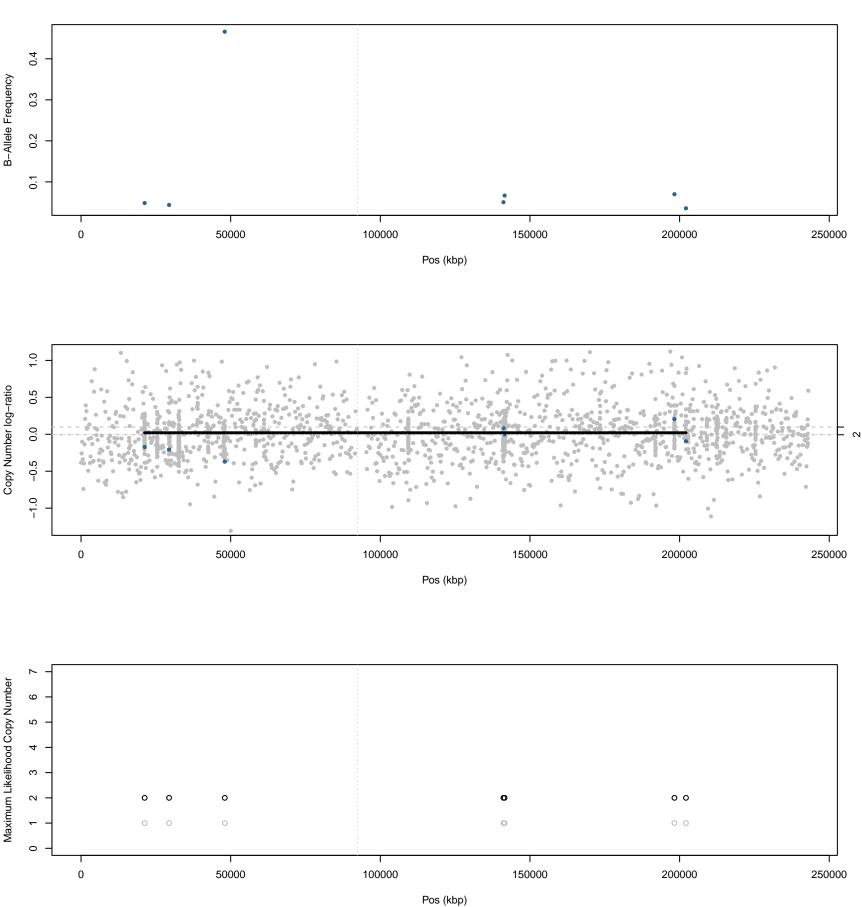
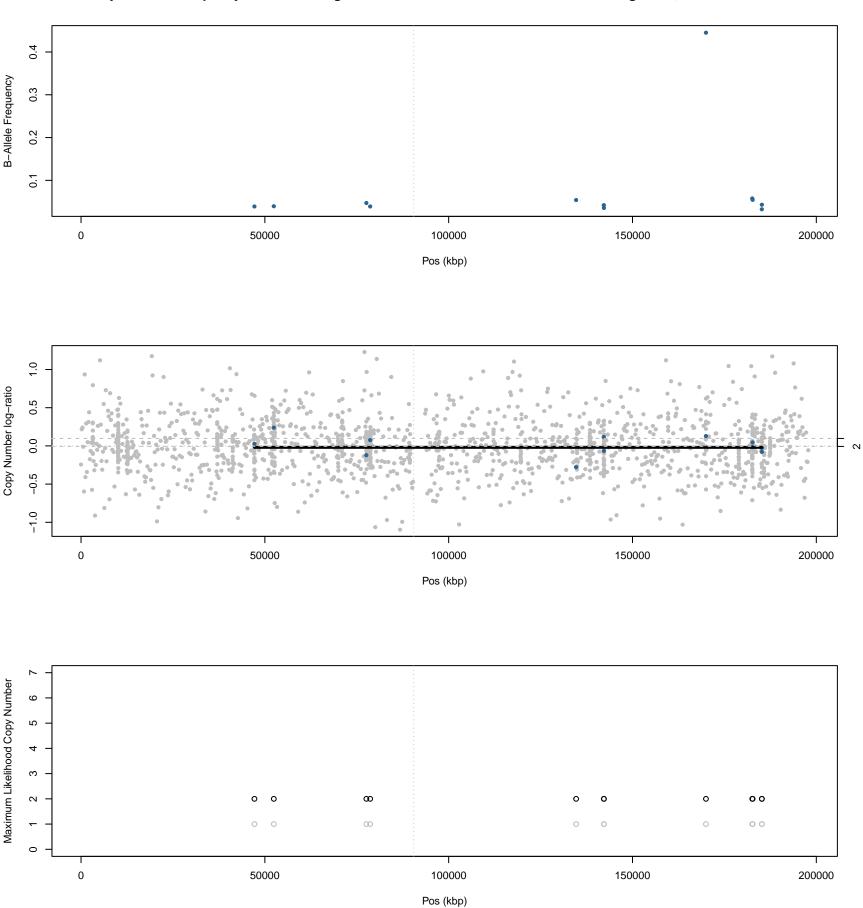
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 1



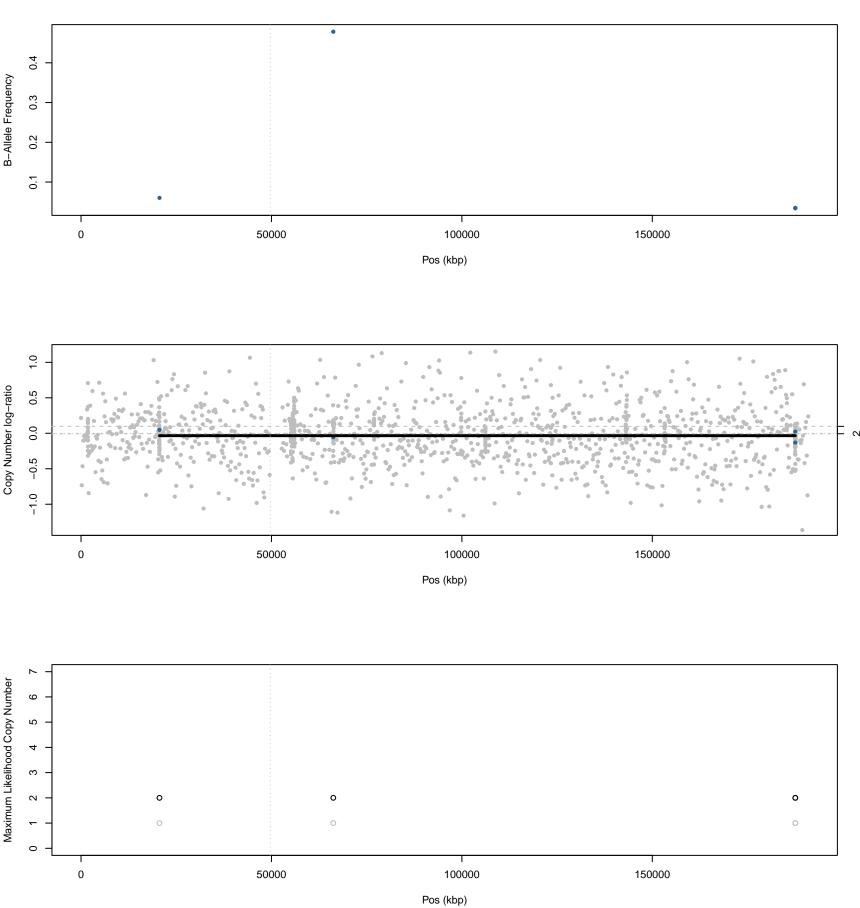
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 2



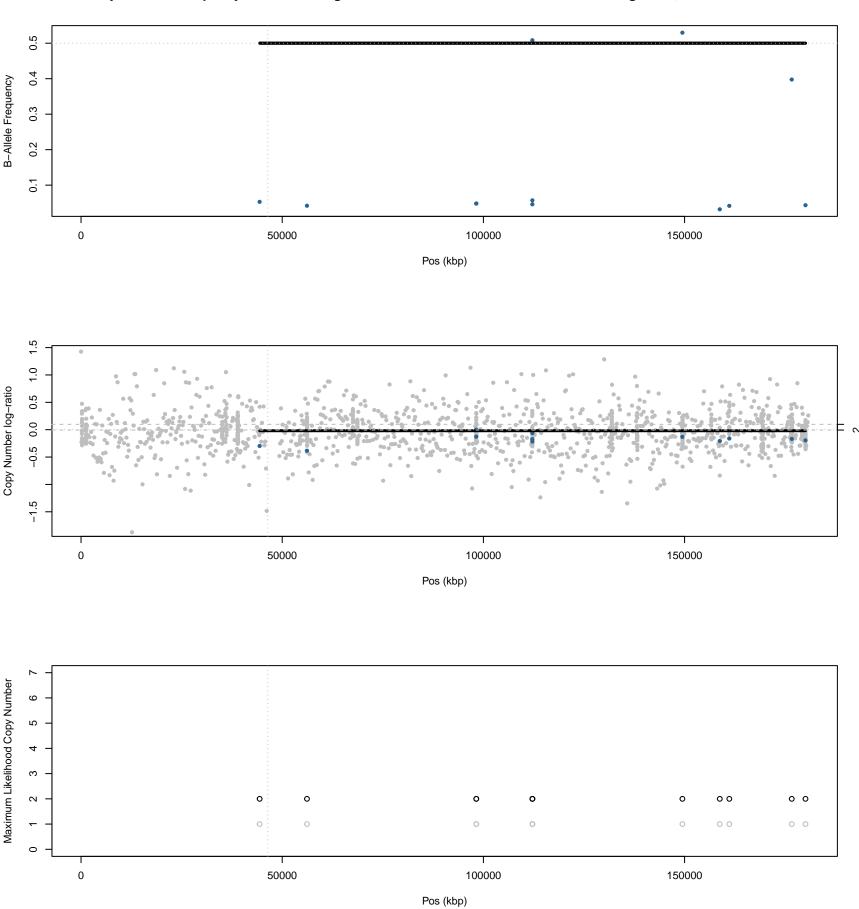
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 3



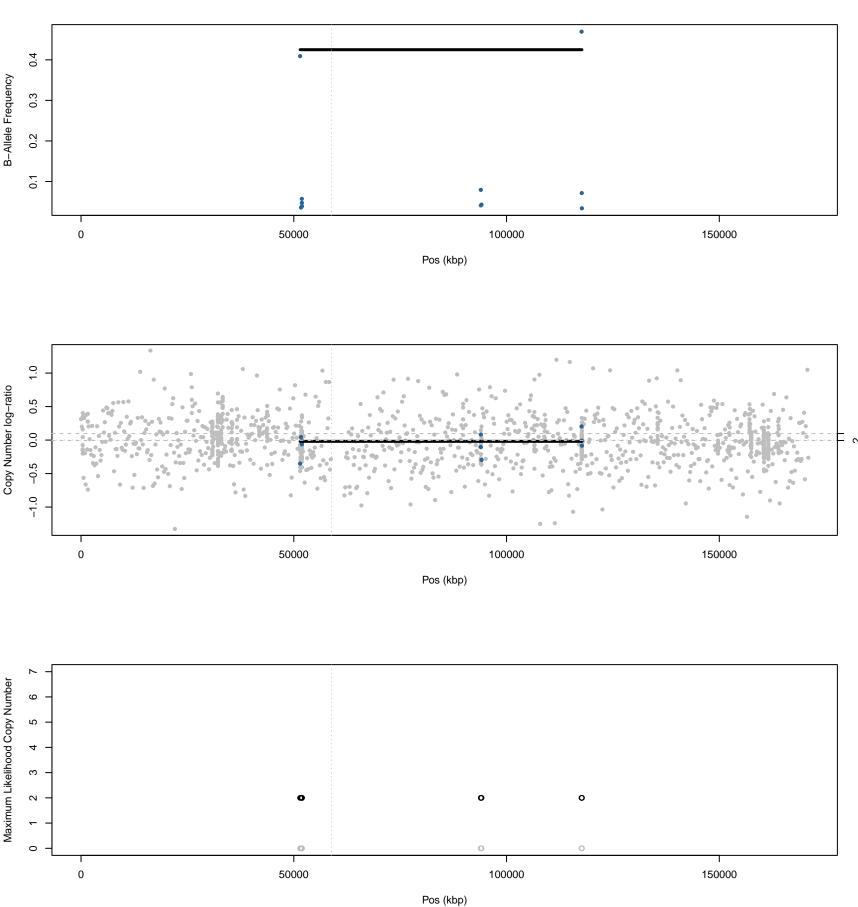
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 4



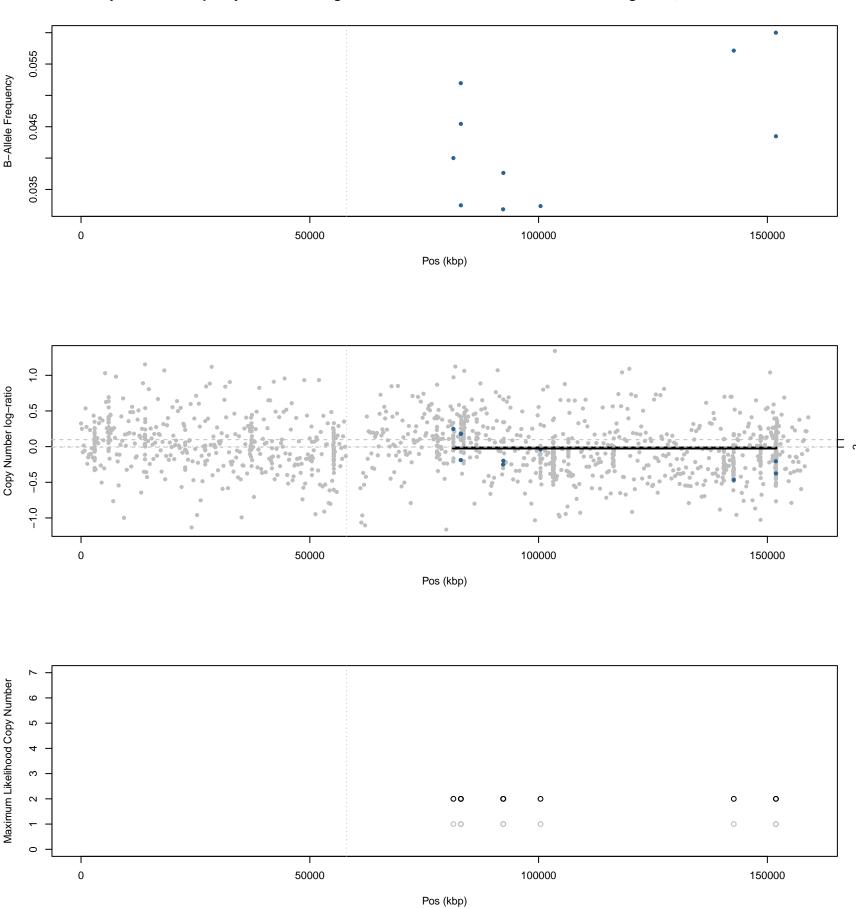
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 5



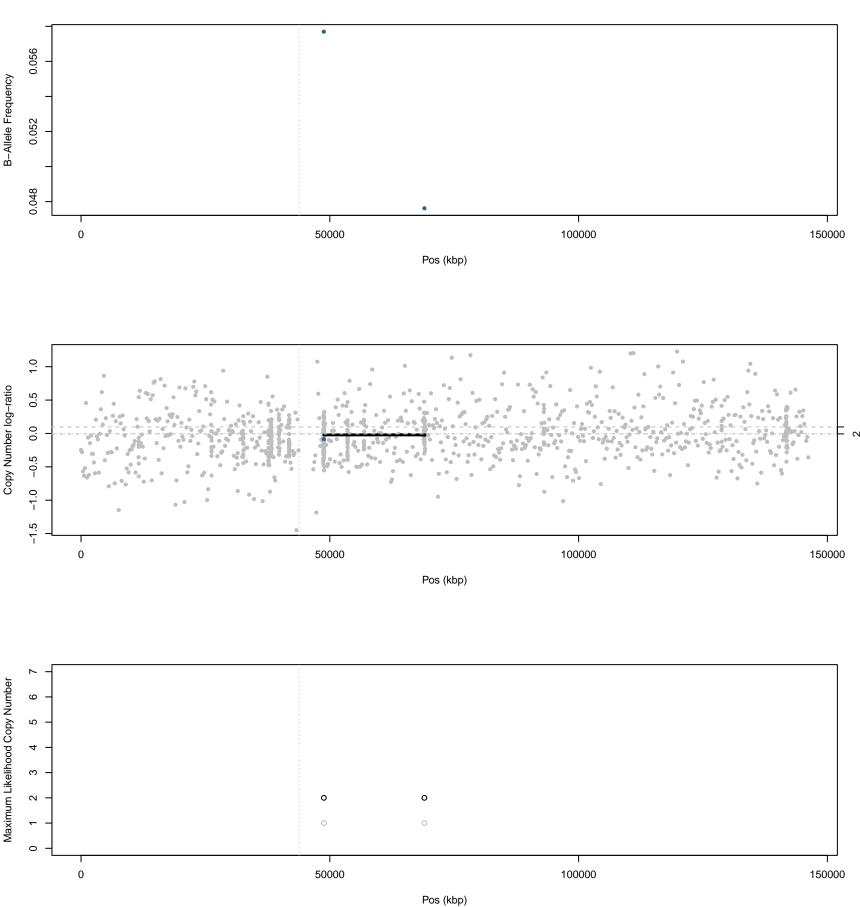
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 6



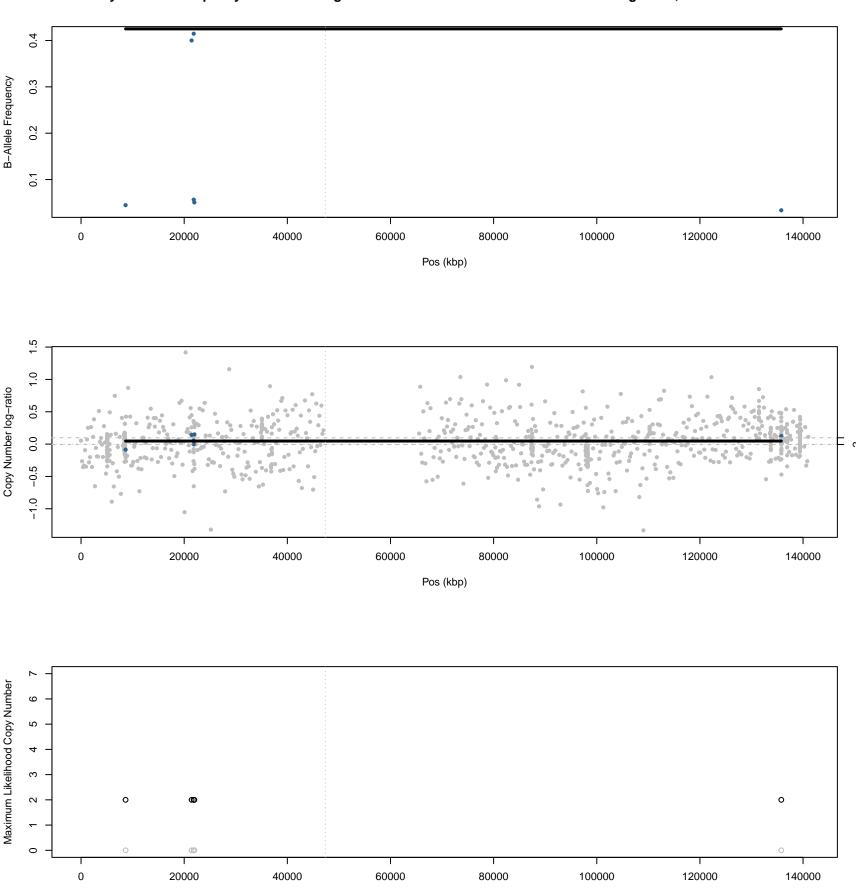
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 7



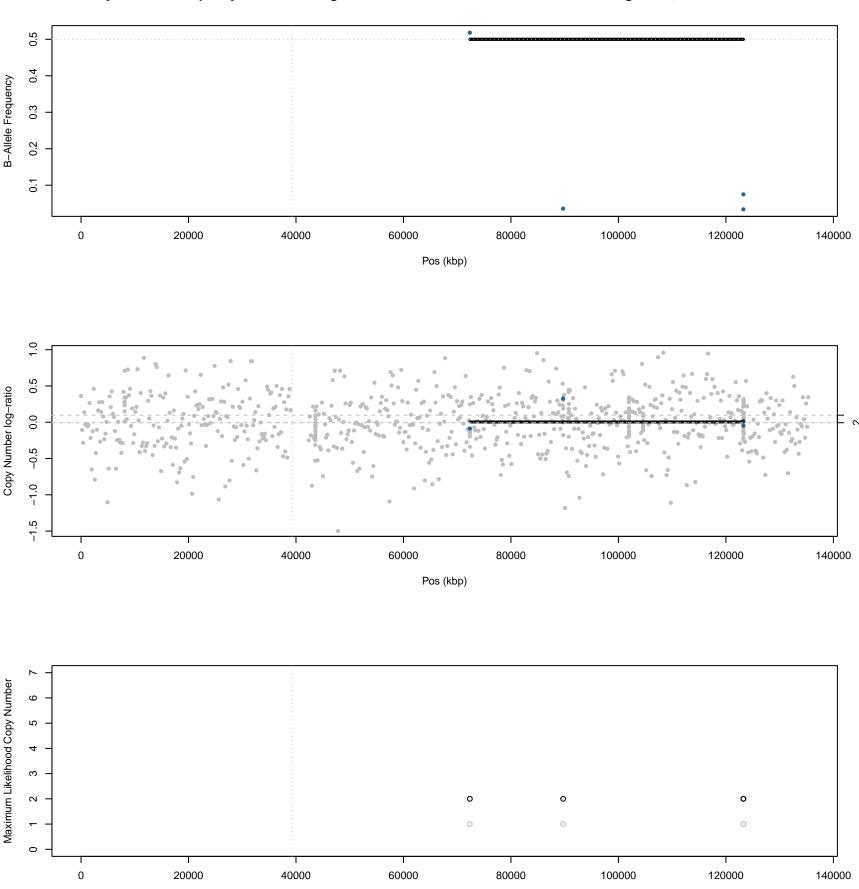
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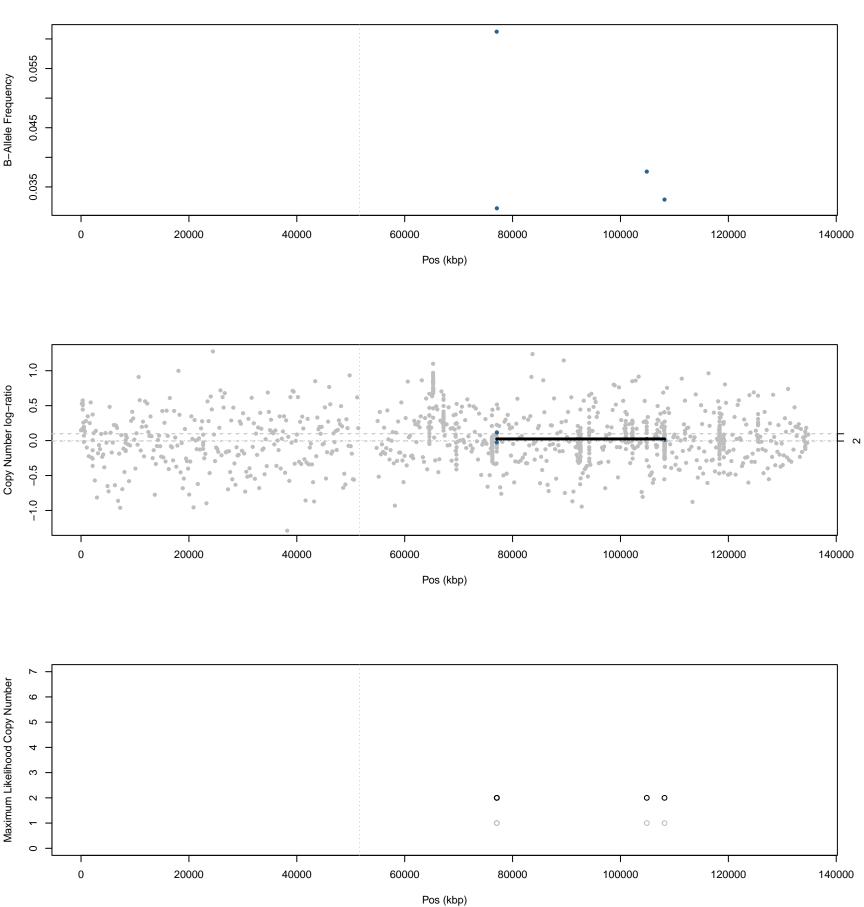
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 9



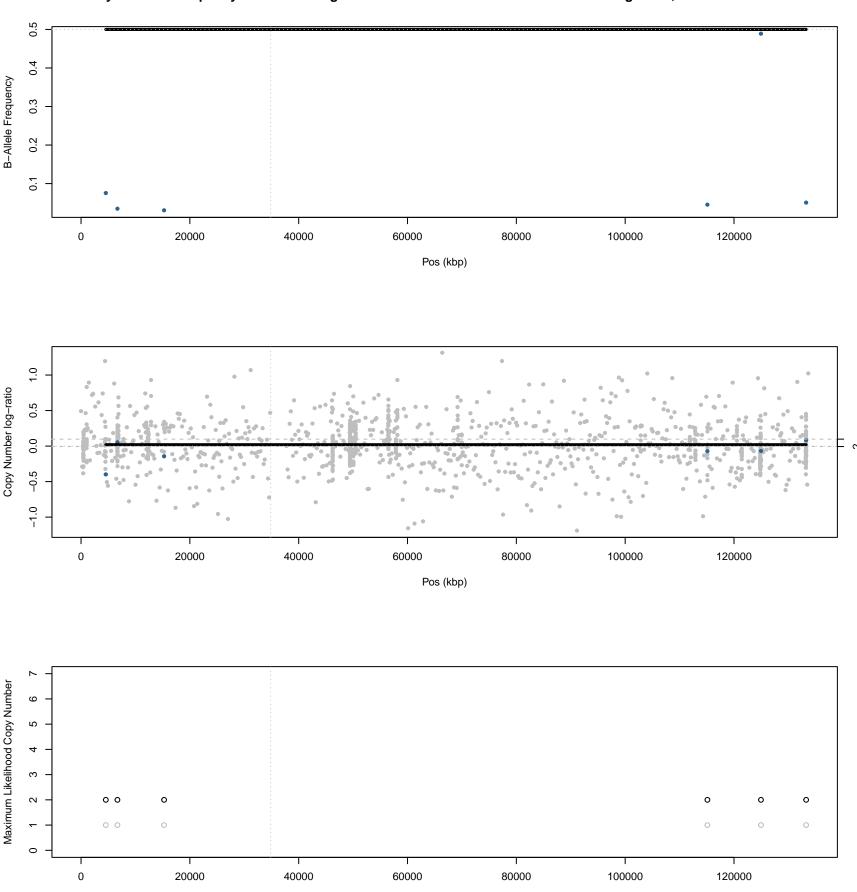
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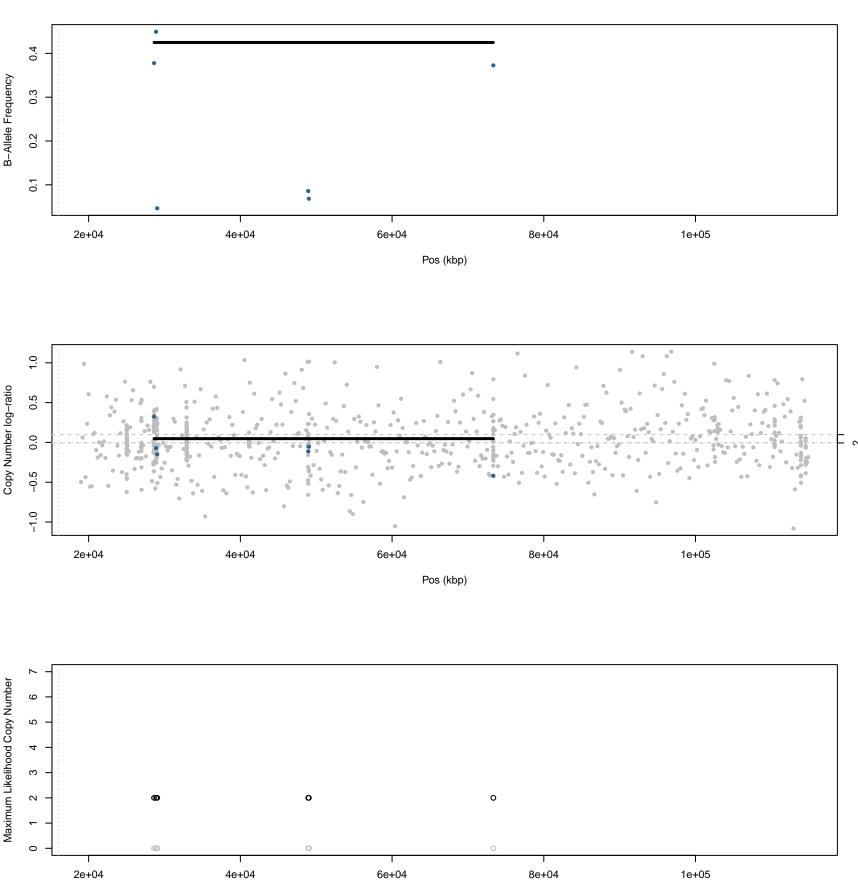
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 11



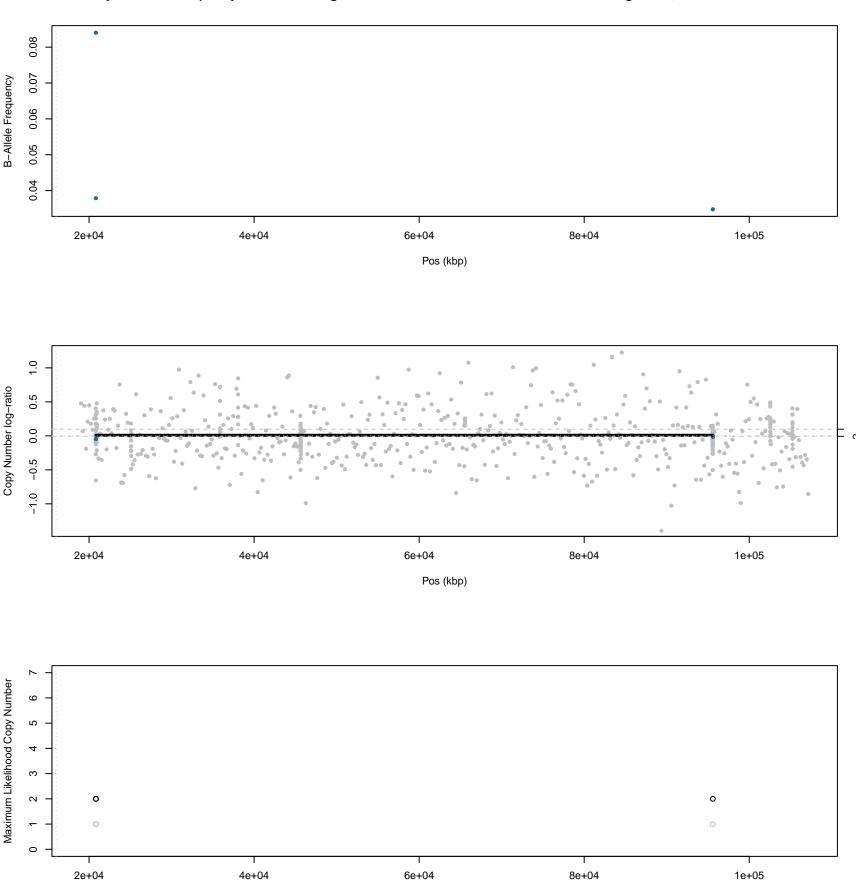
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 12



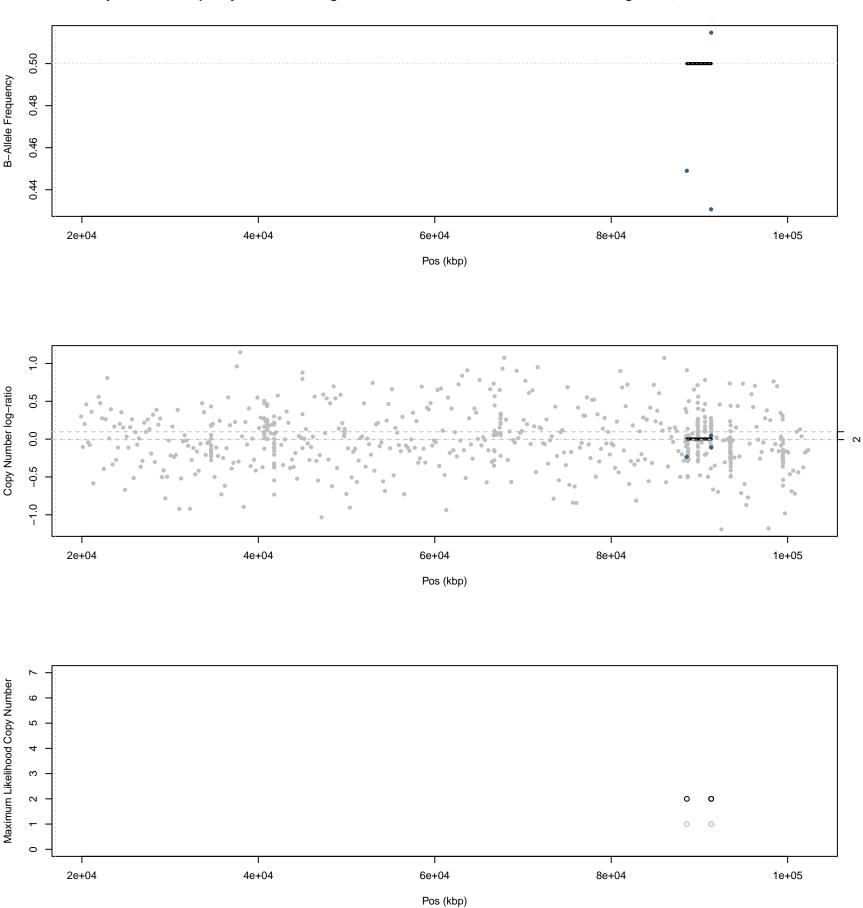
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 13



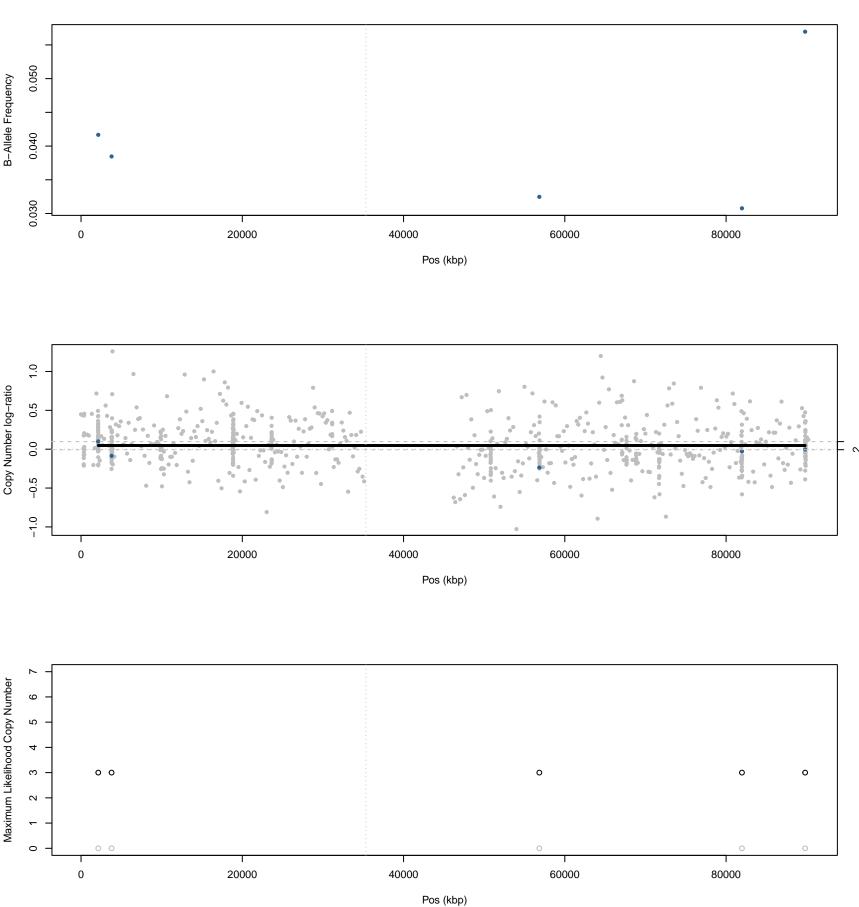
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 14



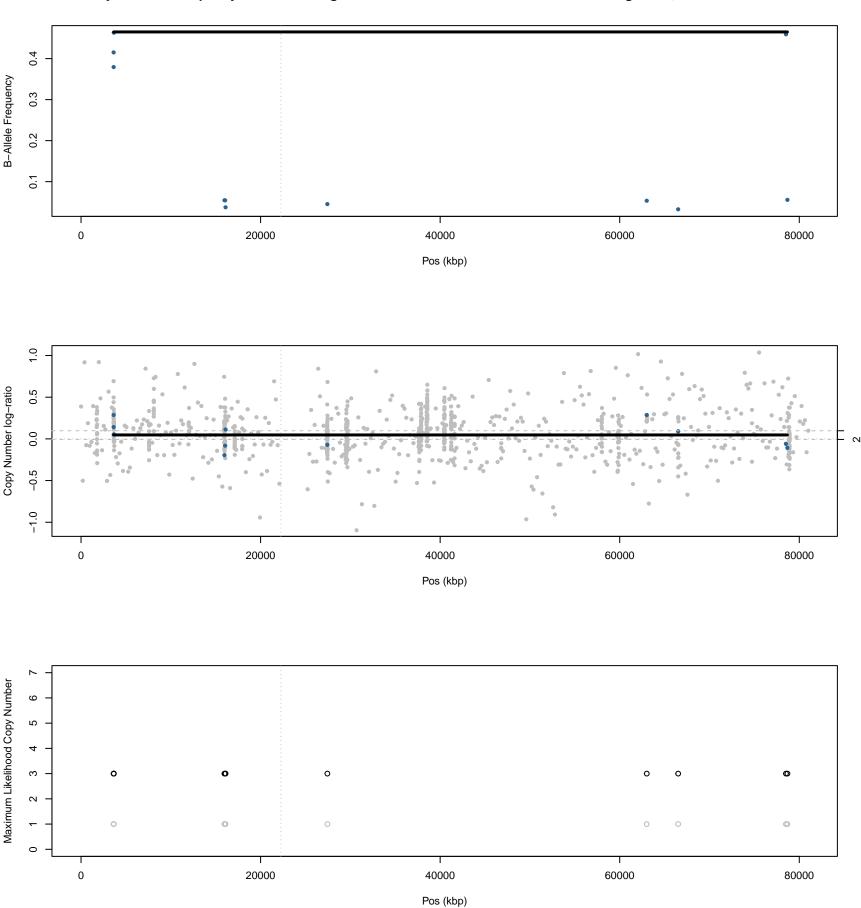
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 15



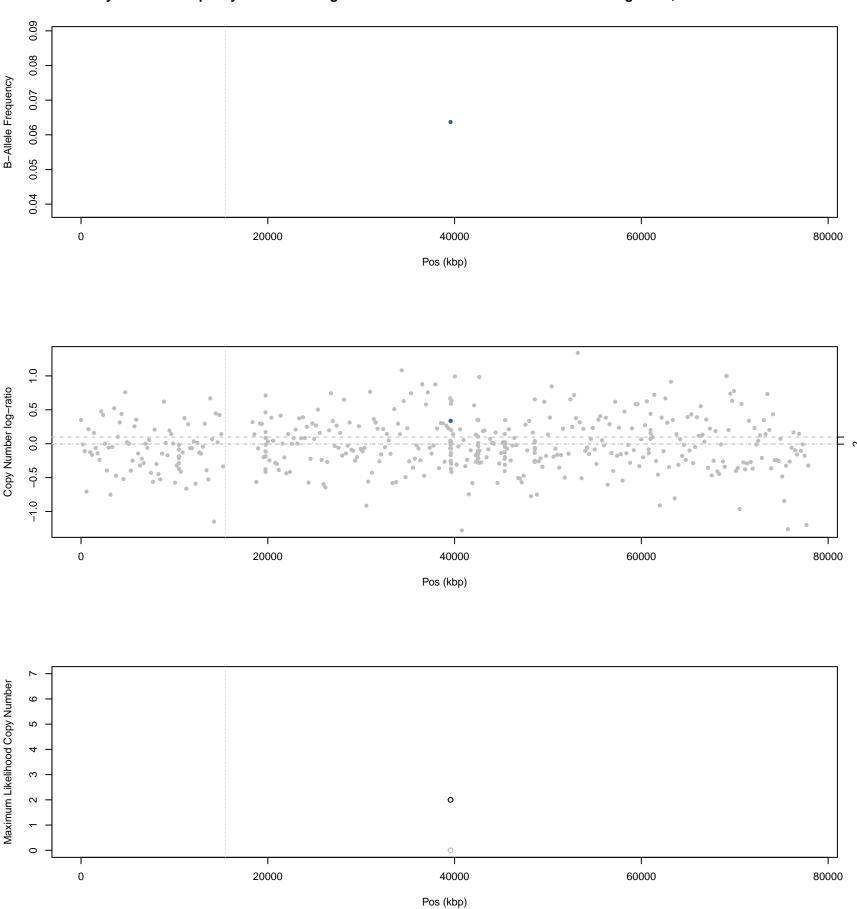
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 16



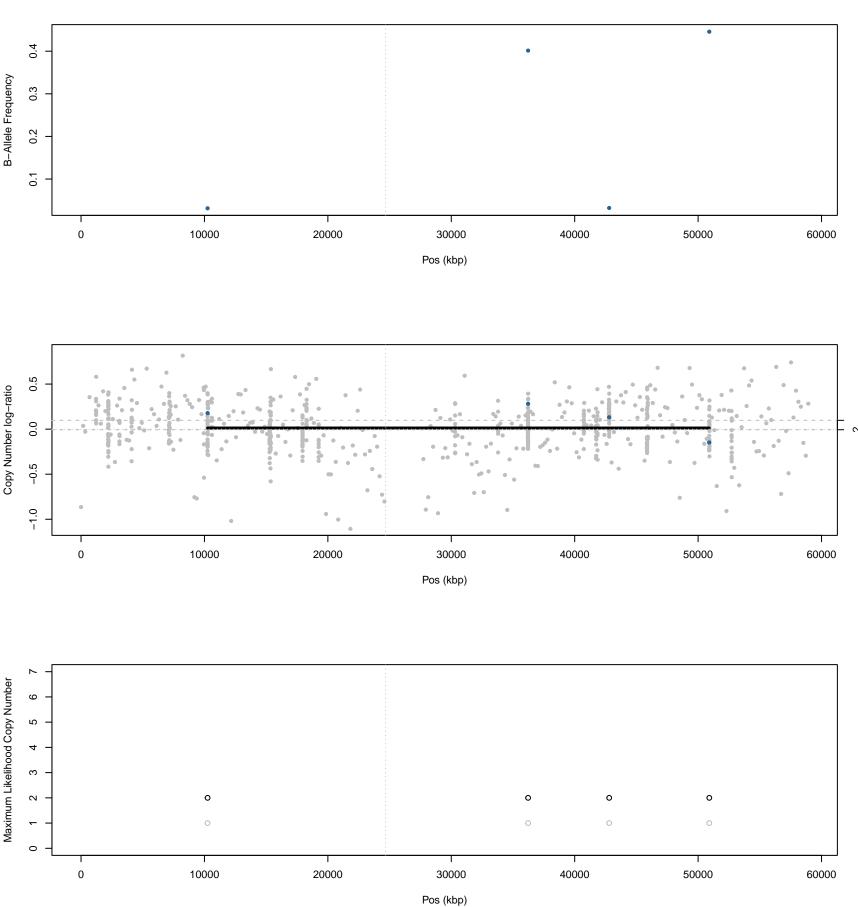
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 17



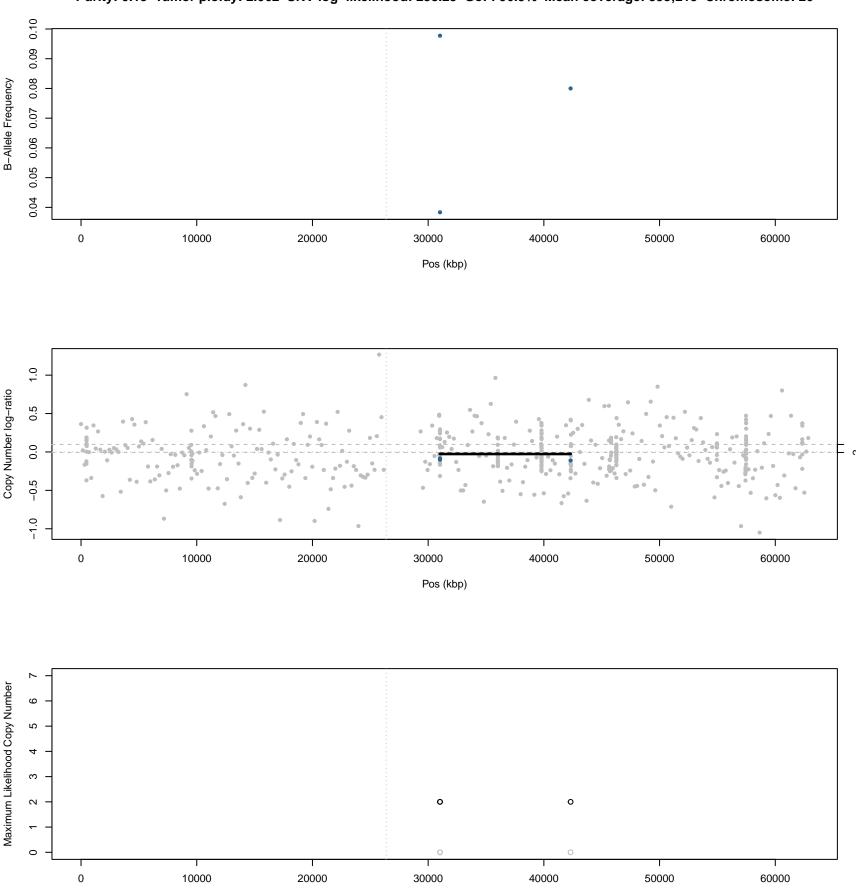
Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 18



Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 19



Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 20



Purity: 0.15 Tumor ploidy: 2.062 SNV log-likelihood: 258.23 GoF: 96.3% Mean coverage: 353;215 Chromosome: 22

