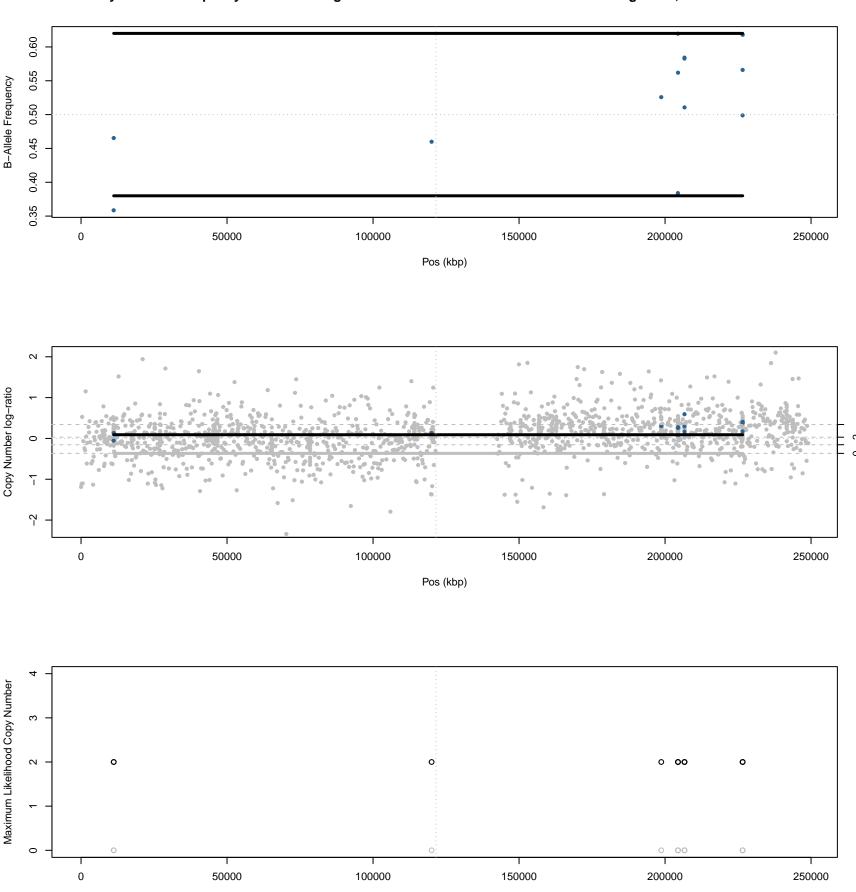
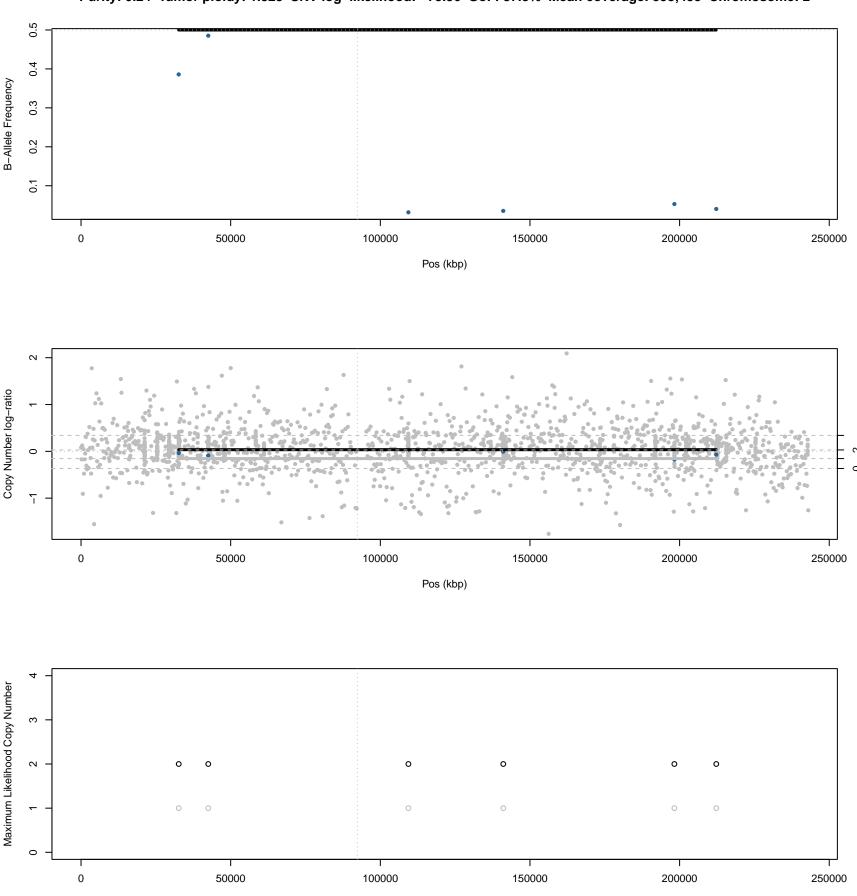
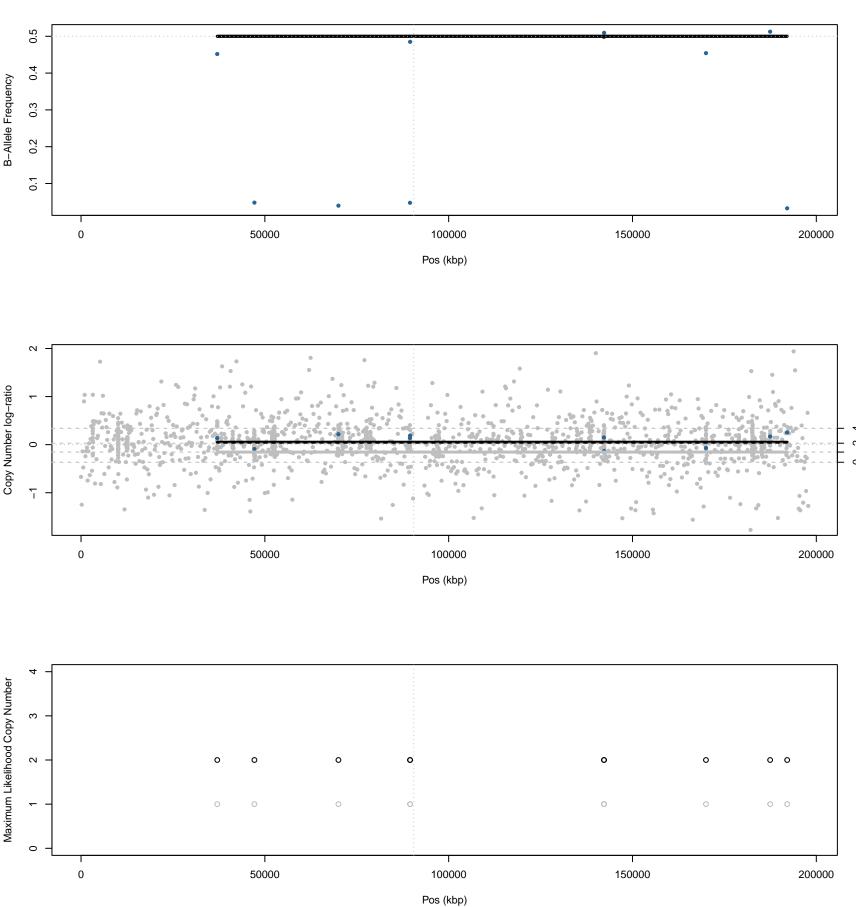
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 1



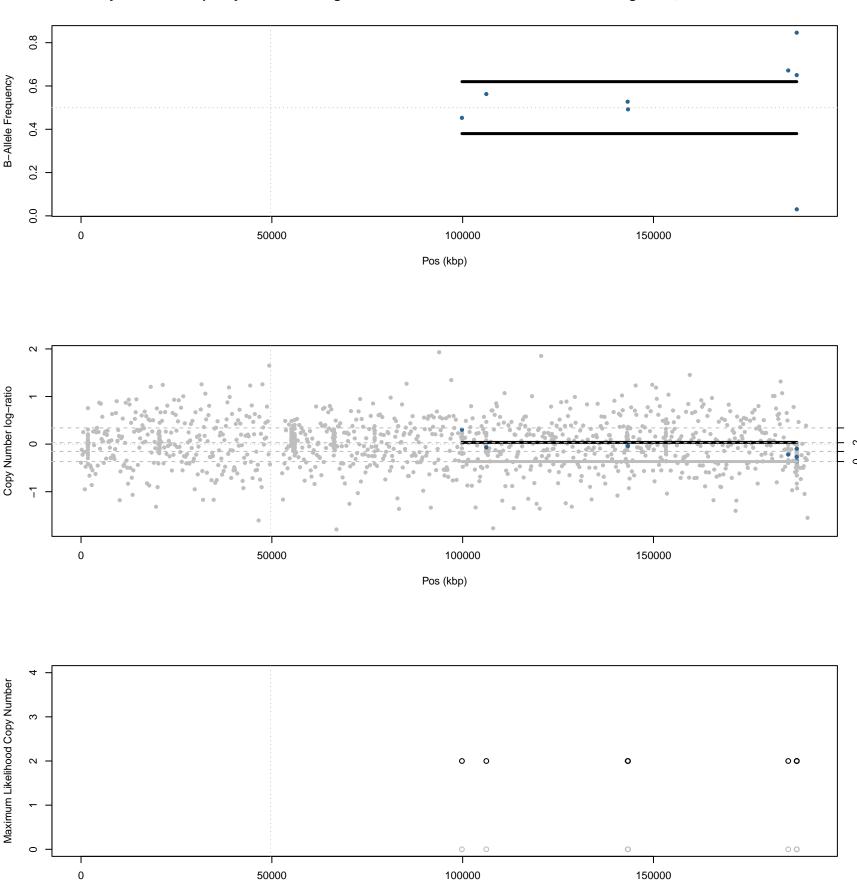
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 2



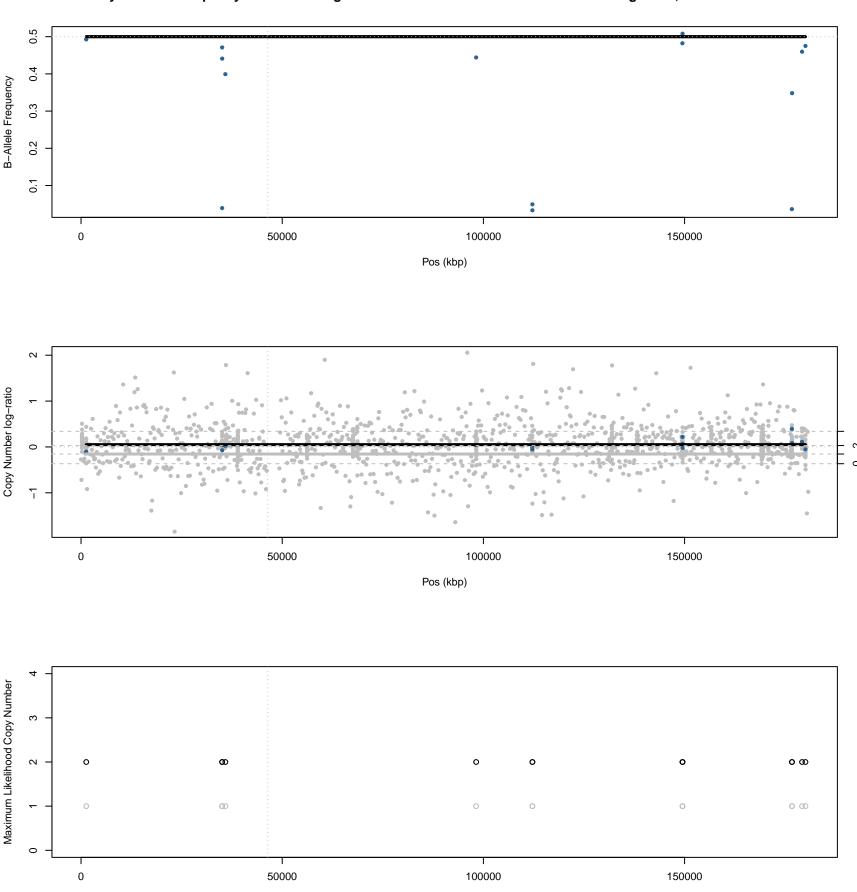
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 3



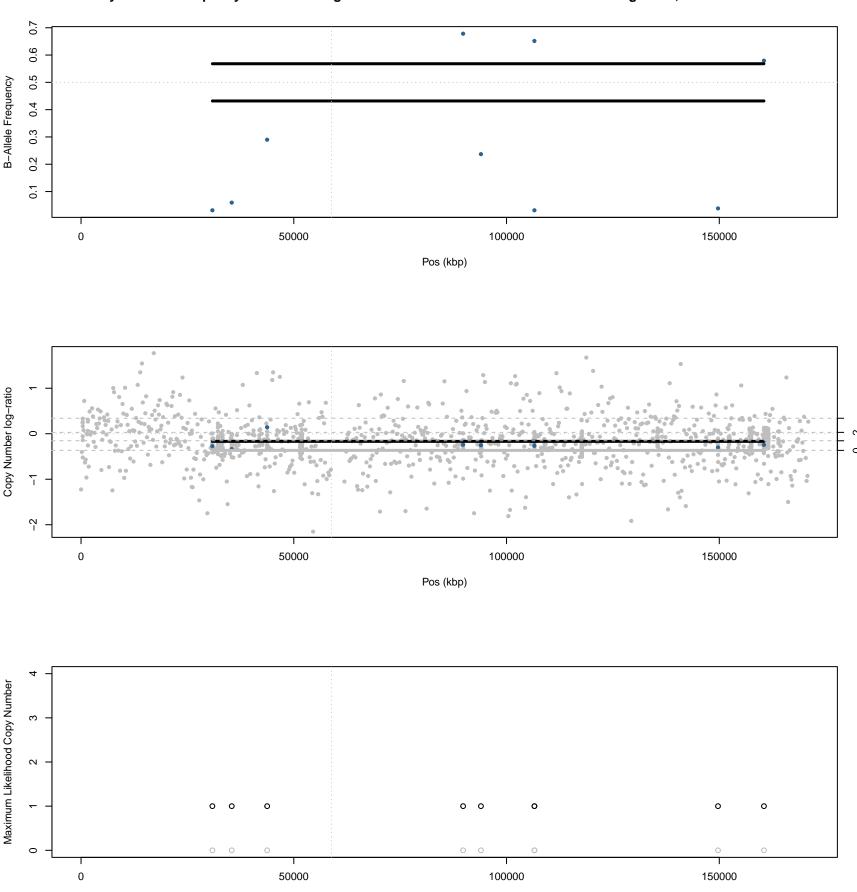
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 4



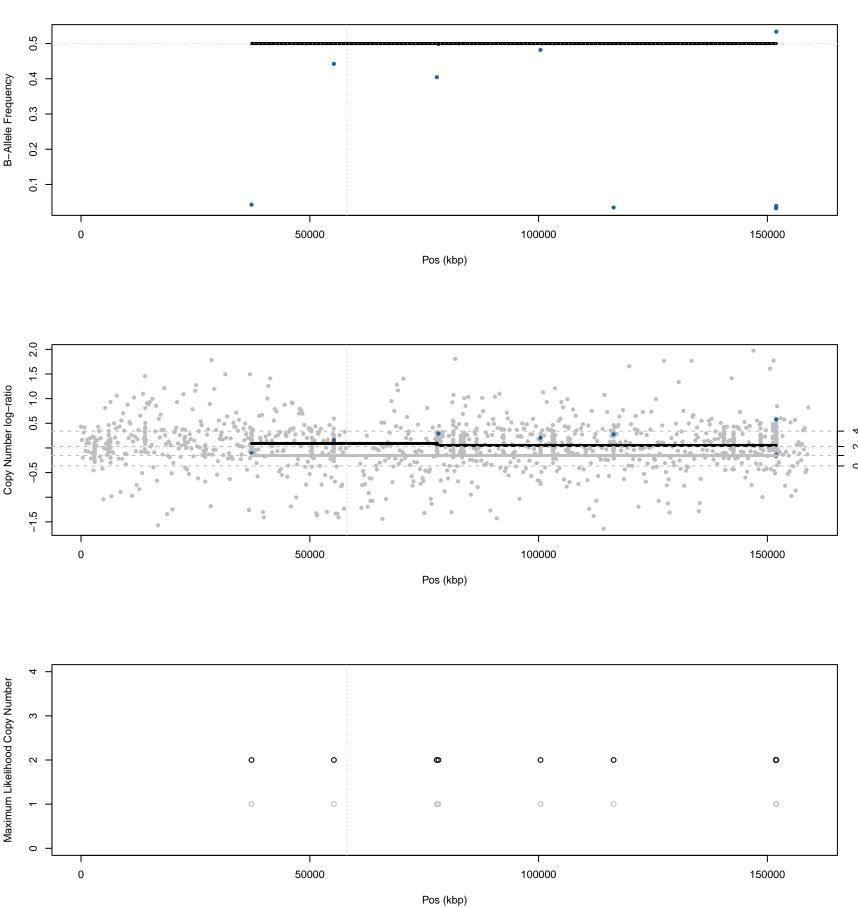
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 5



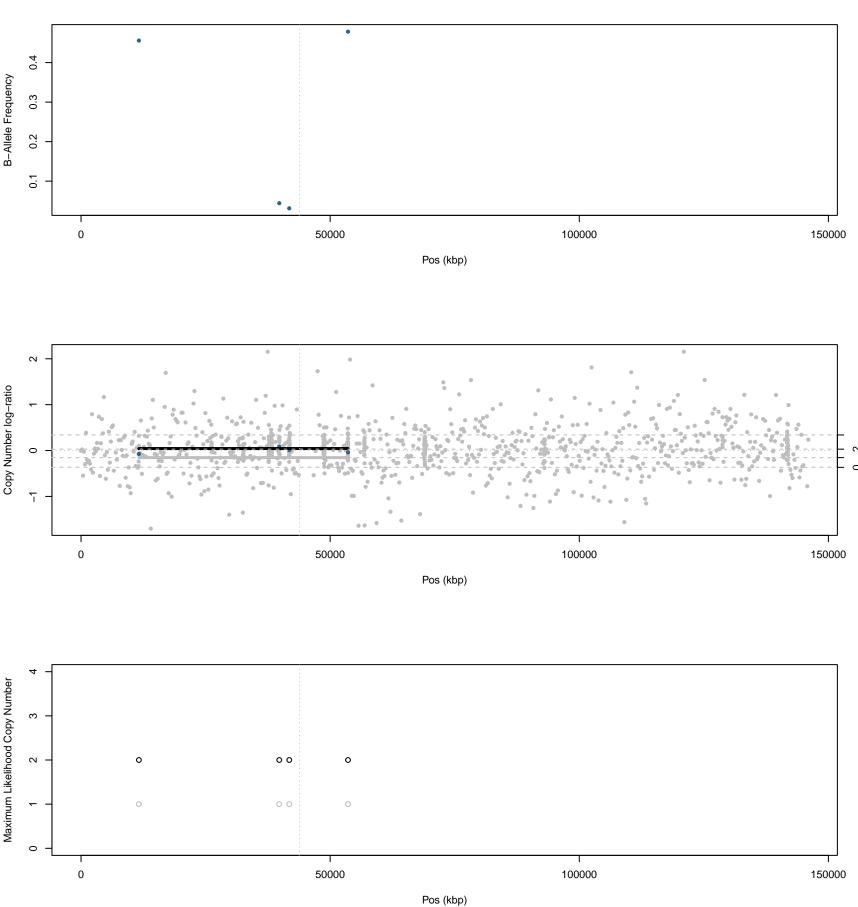
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 6



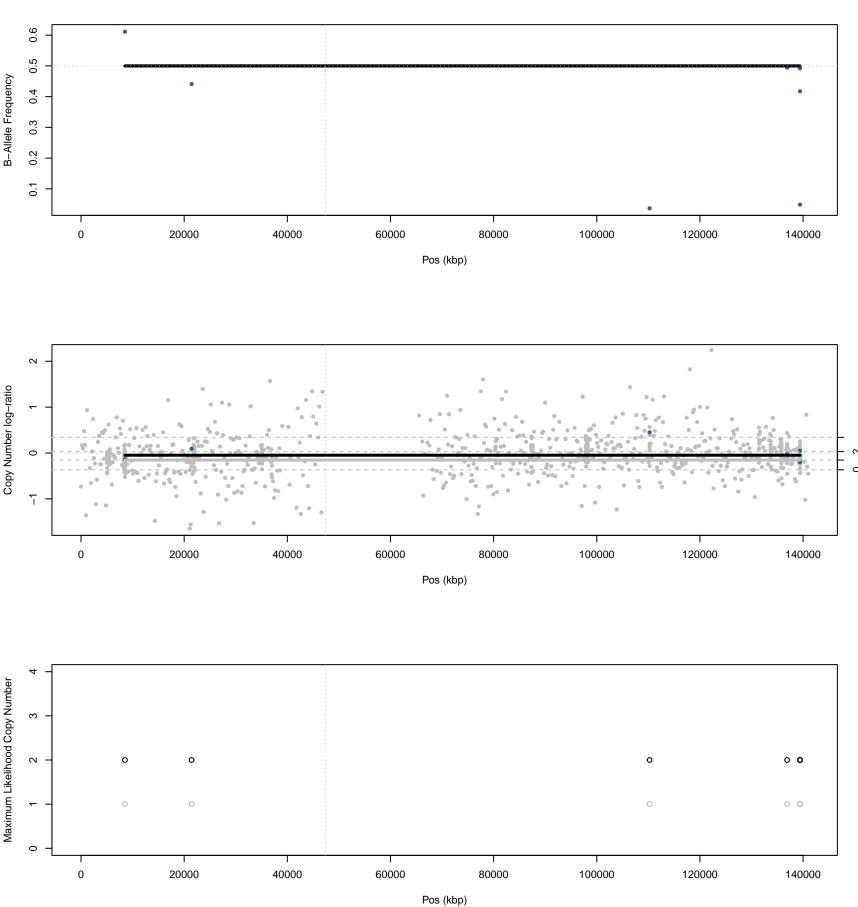
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 7



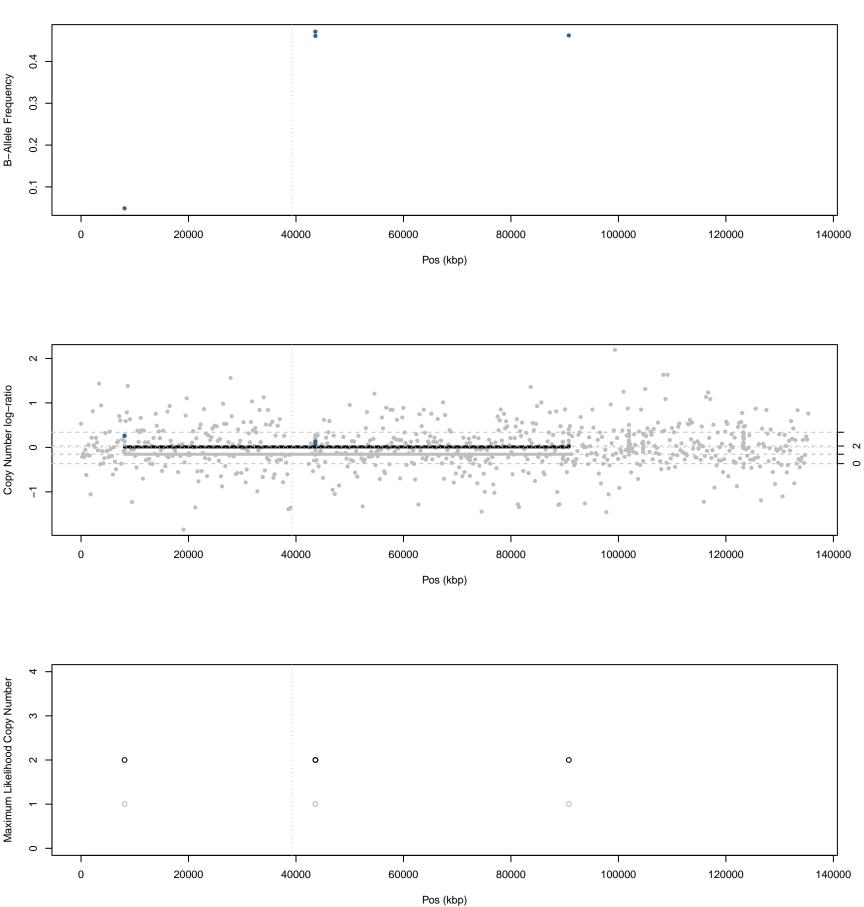
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 8



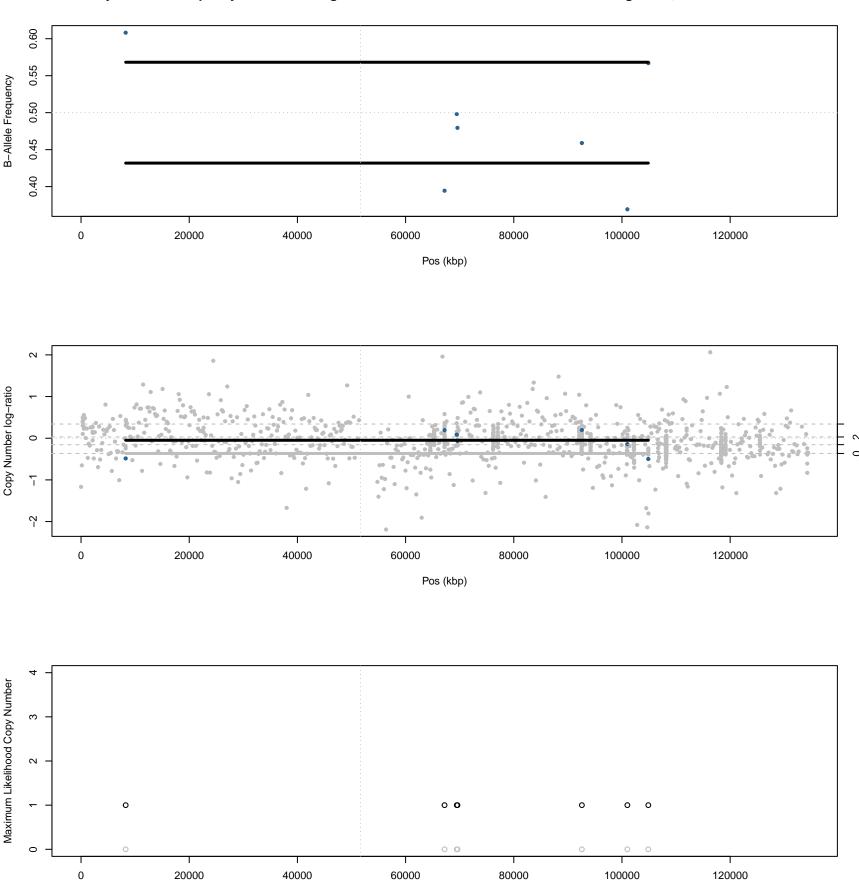
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 9



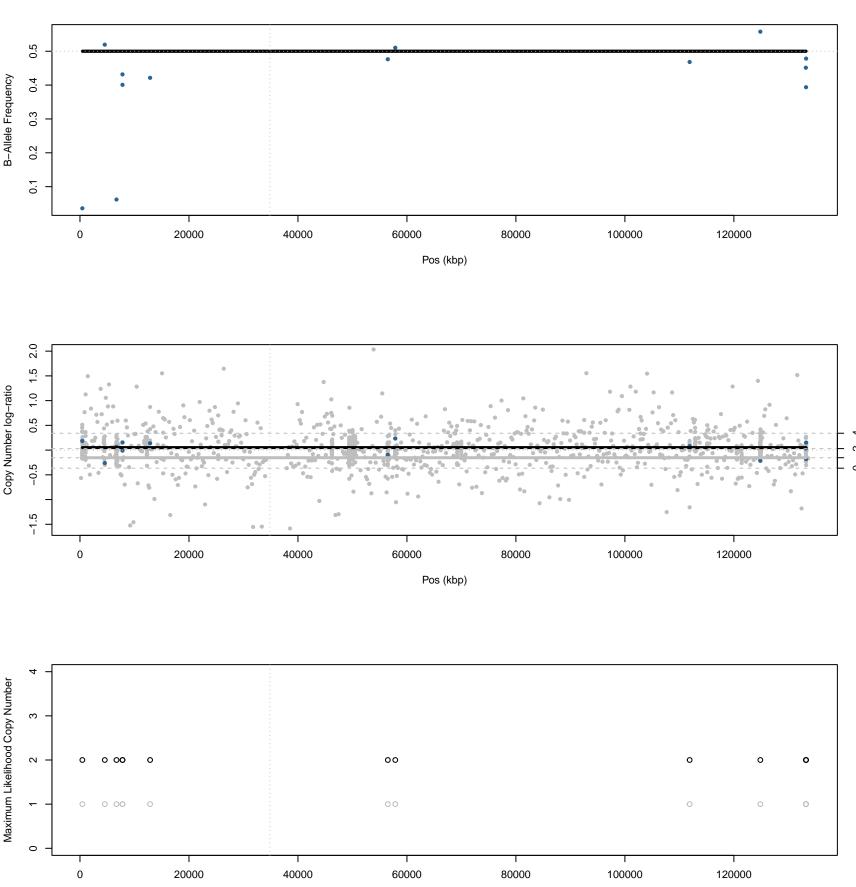
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 10



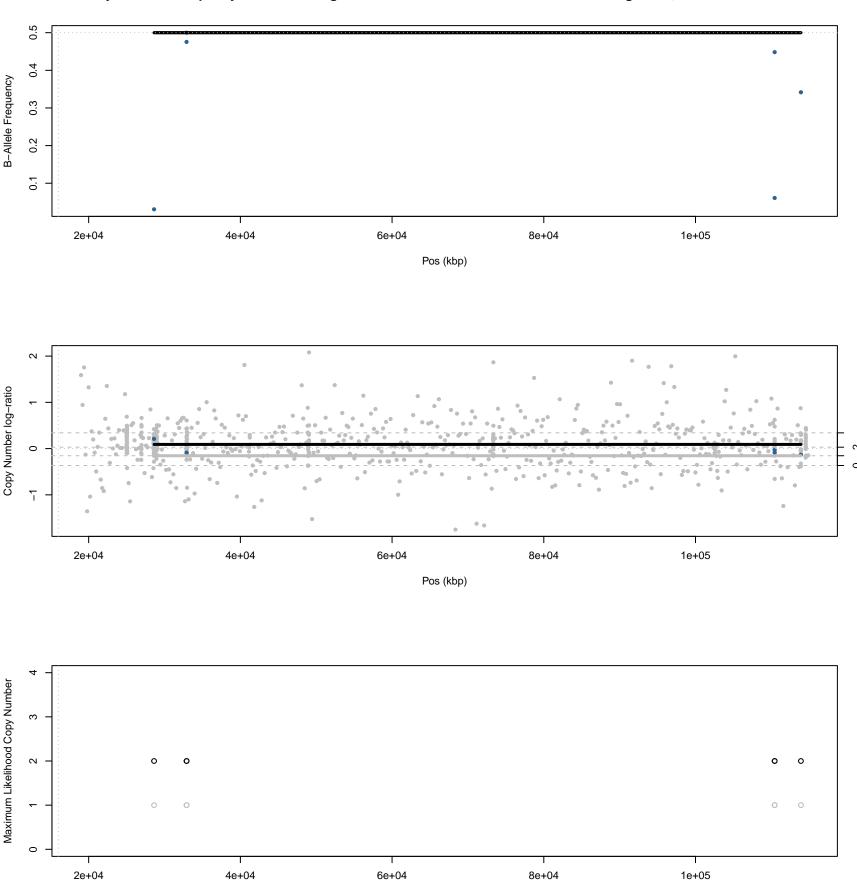
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 11



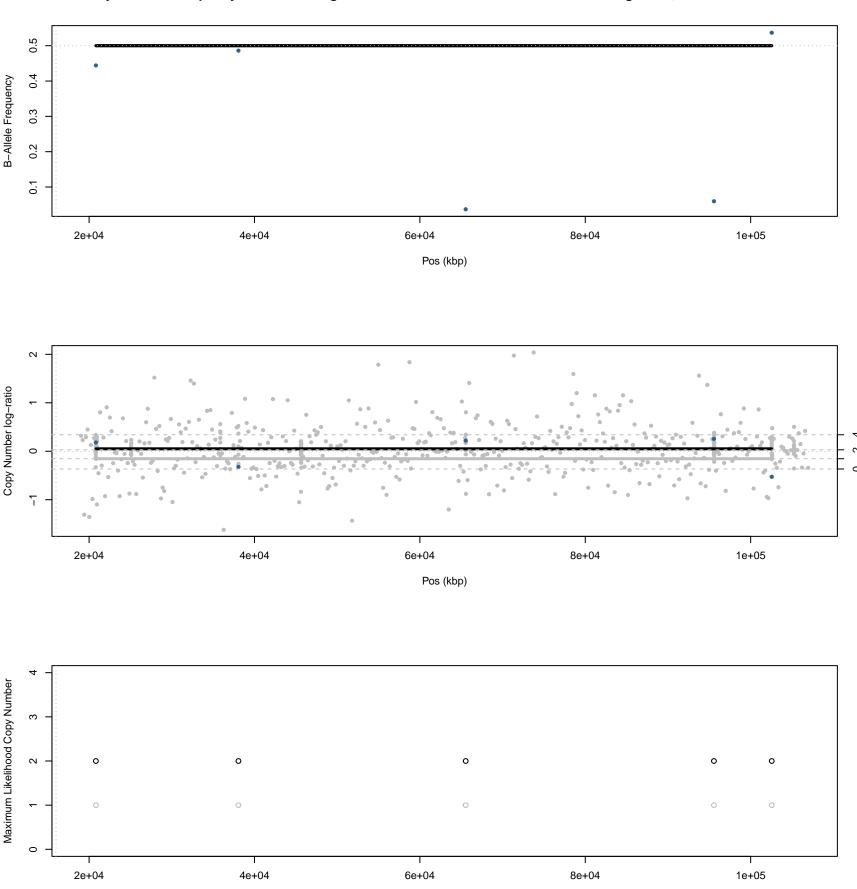
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 12



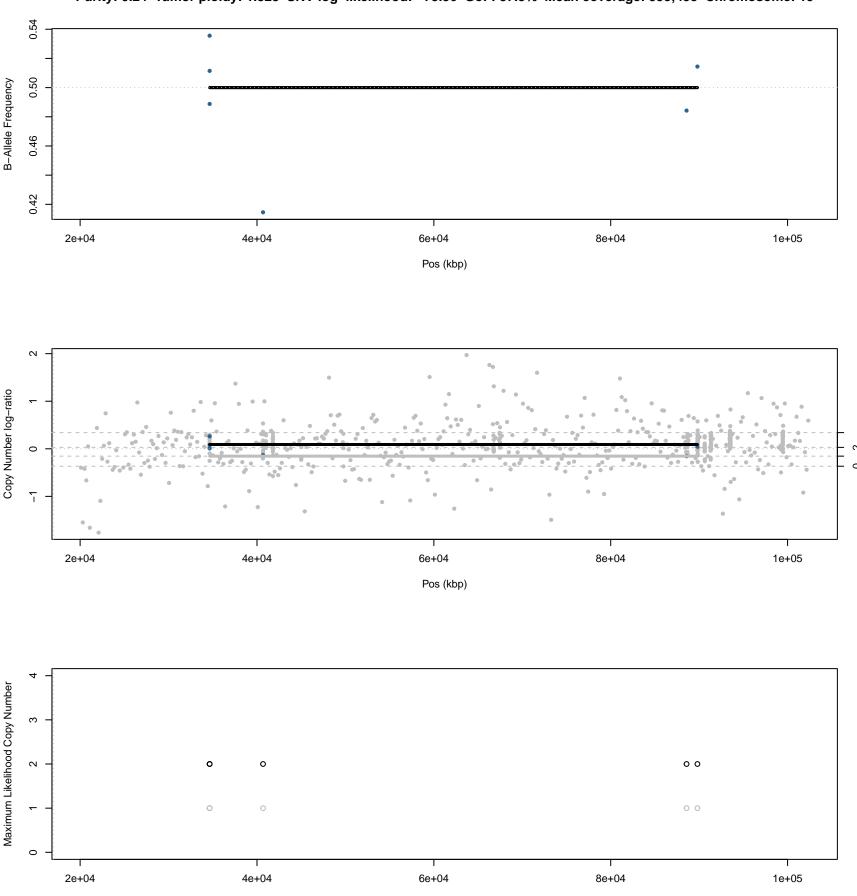
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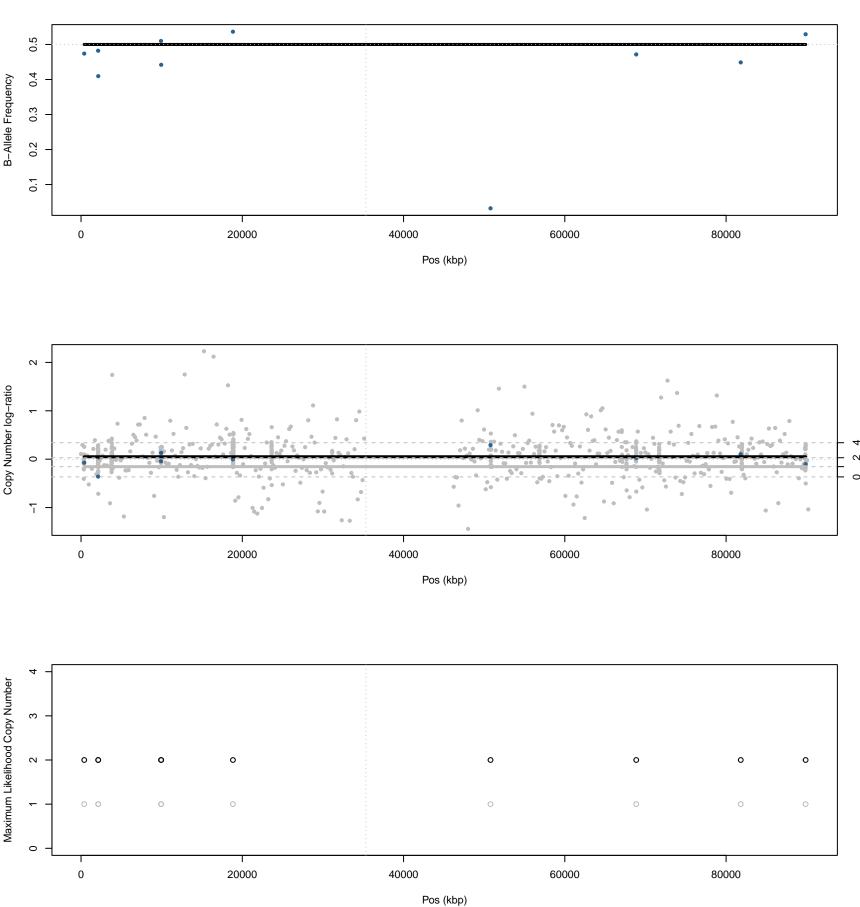
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 14



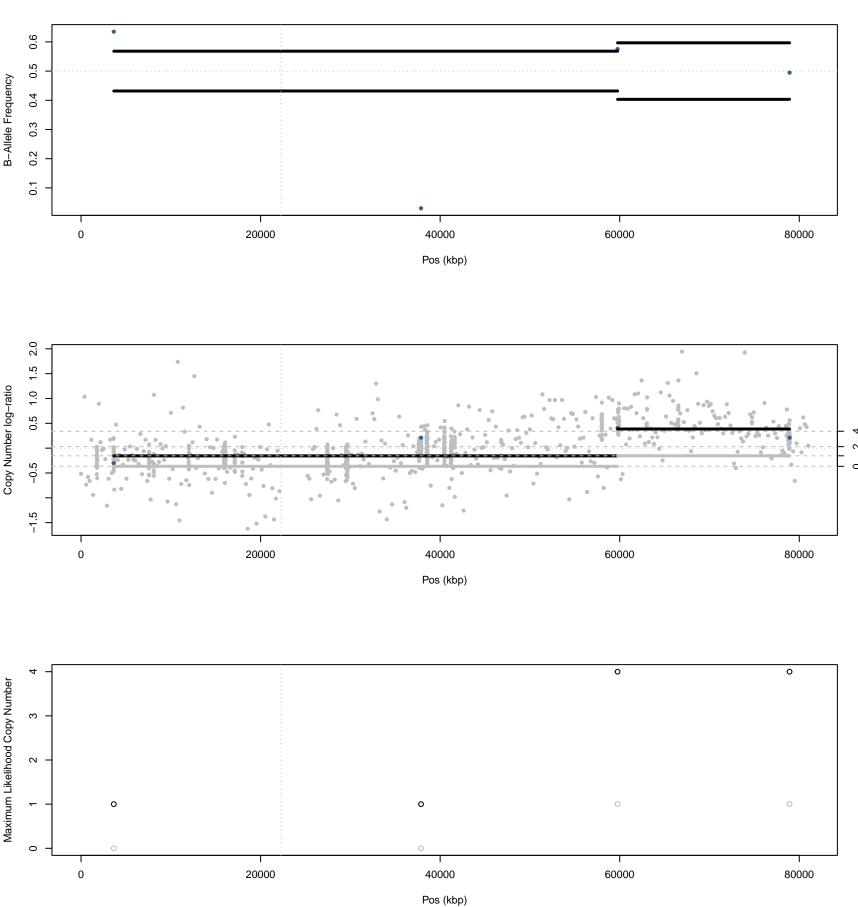
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 15



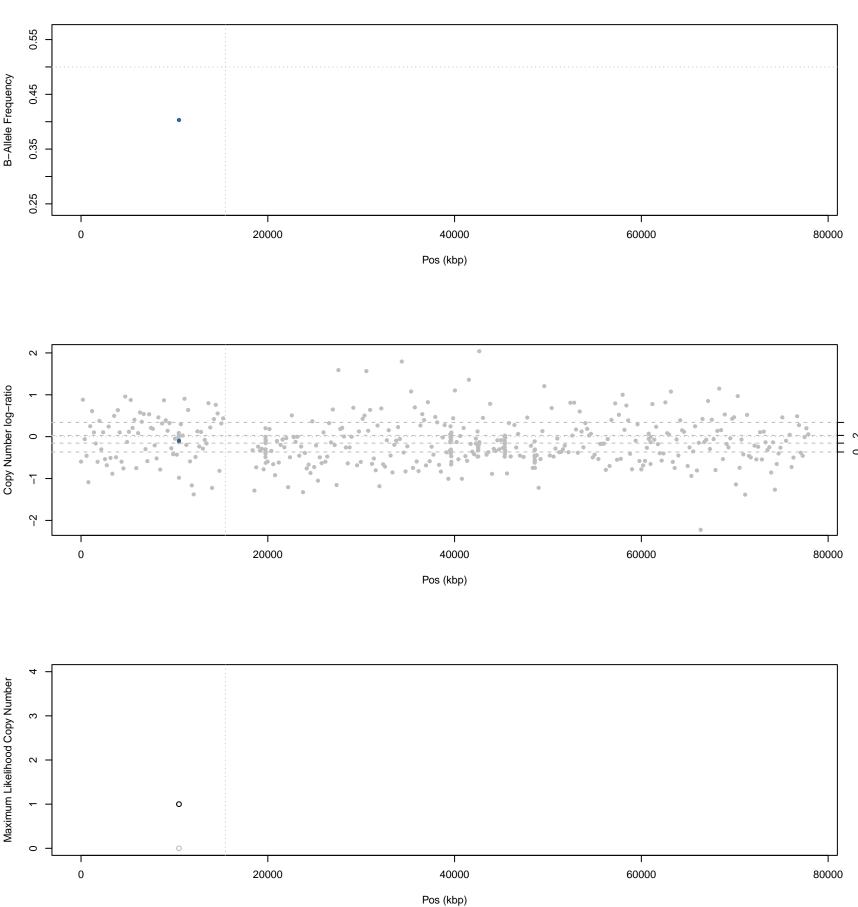
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 16



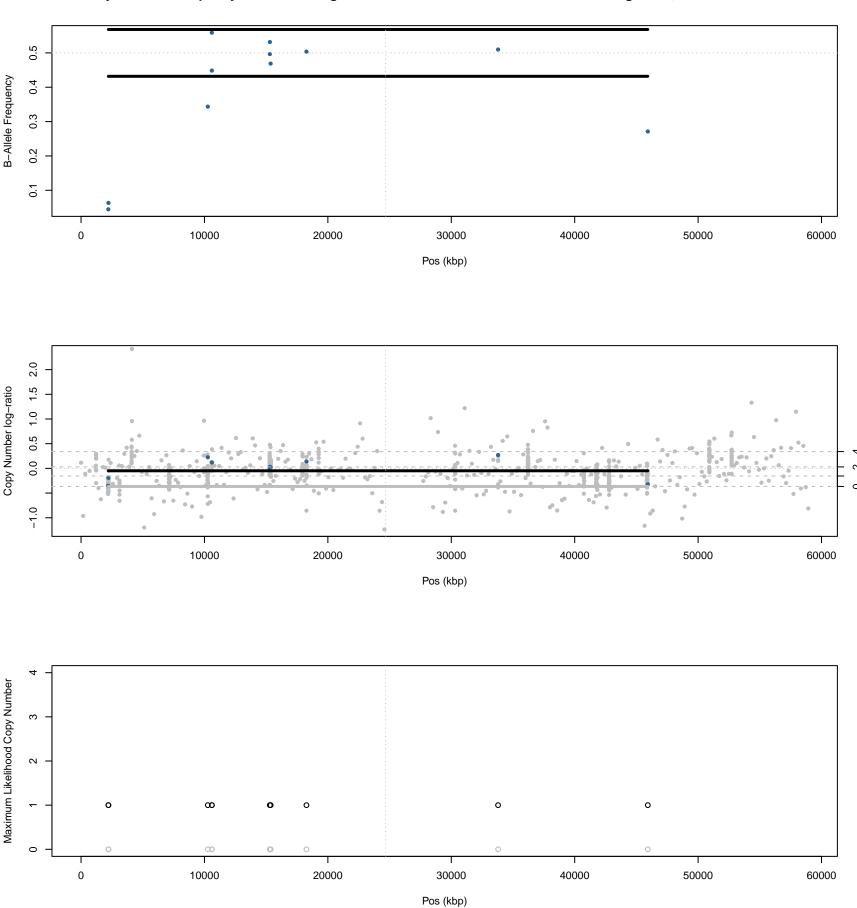
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 17



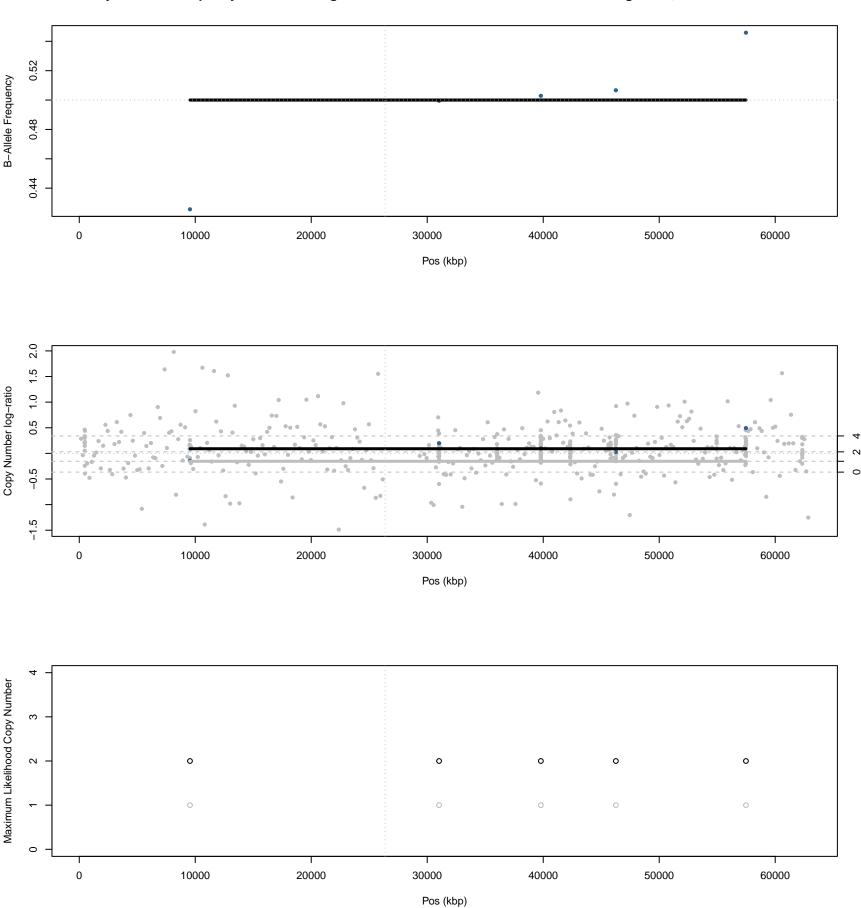
Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 18



Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 19



Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 20



Purity: 0.24 Tumor ploidy: 1.828 SNV log-likelihood: -73.36 GoF: 87.8% Mean coverage: 558;453 Chromosome: 22

