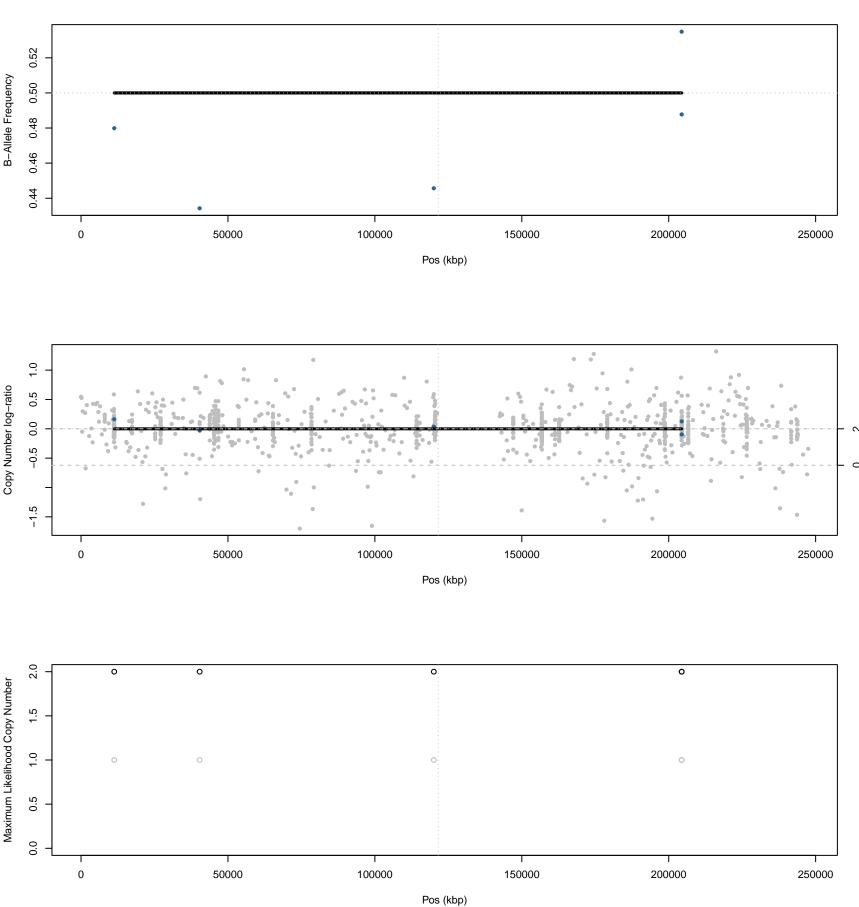
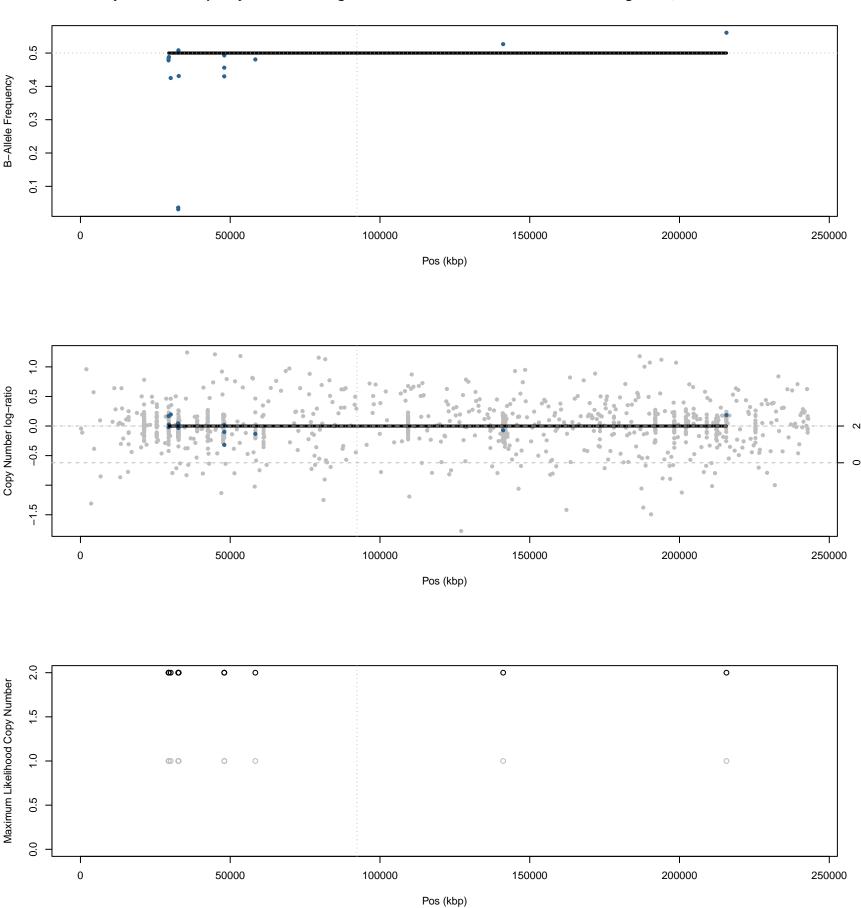
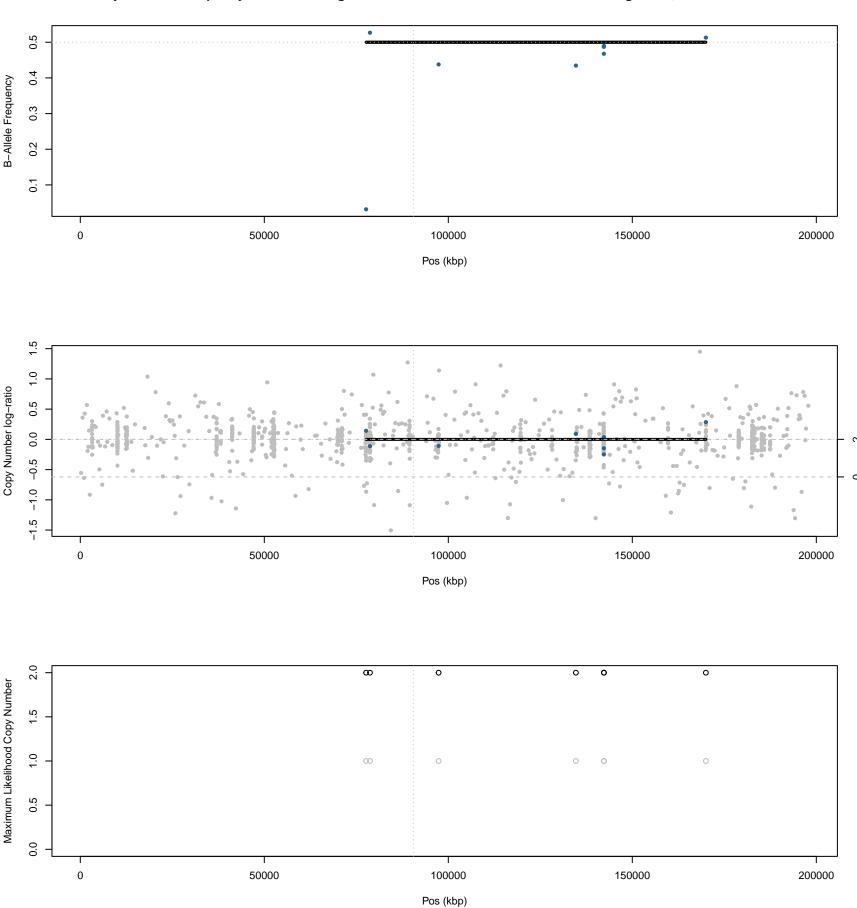
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 1



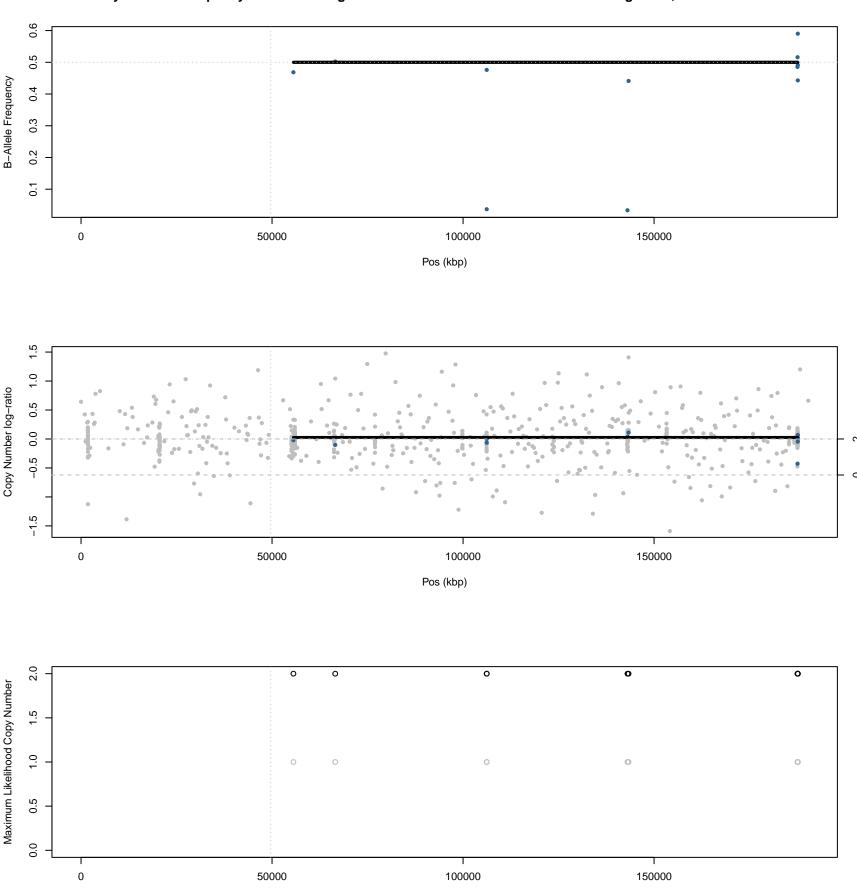
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 2



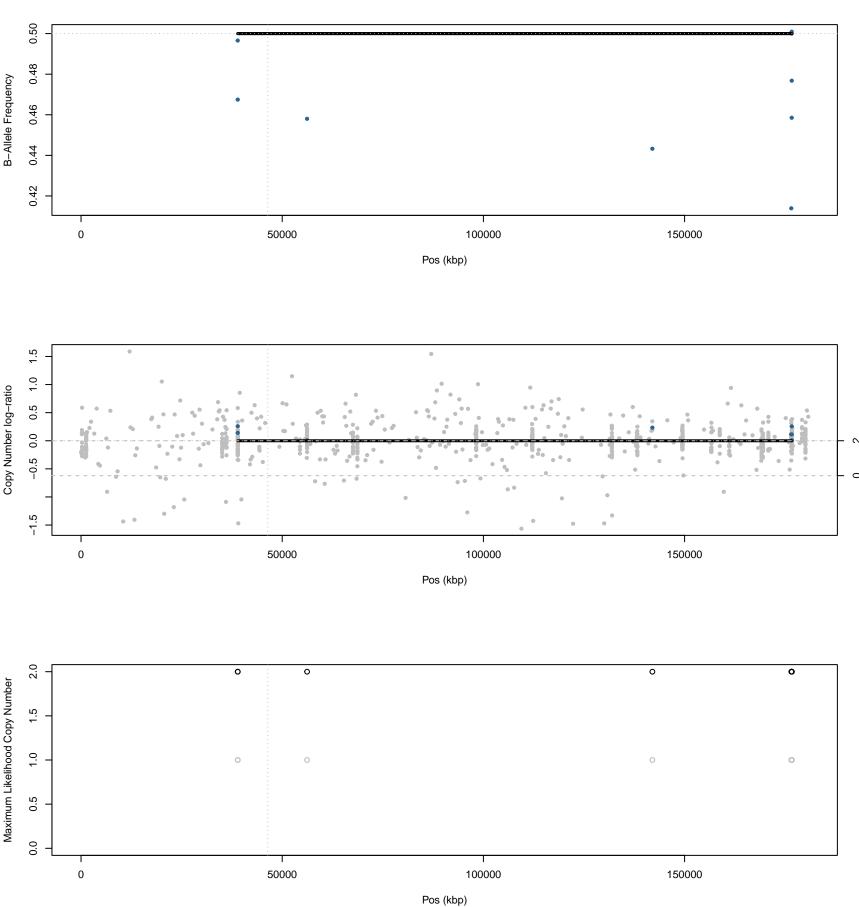
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 3



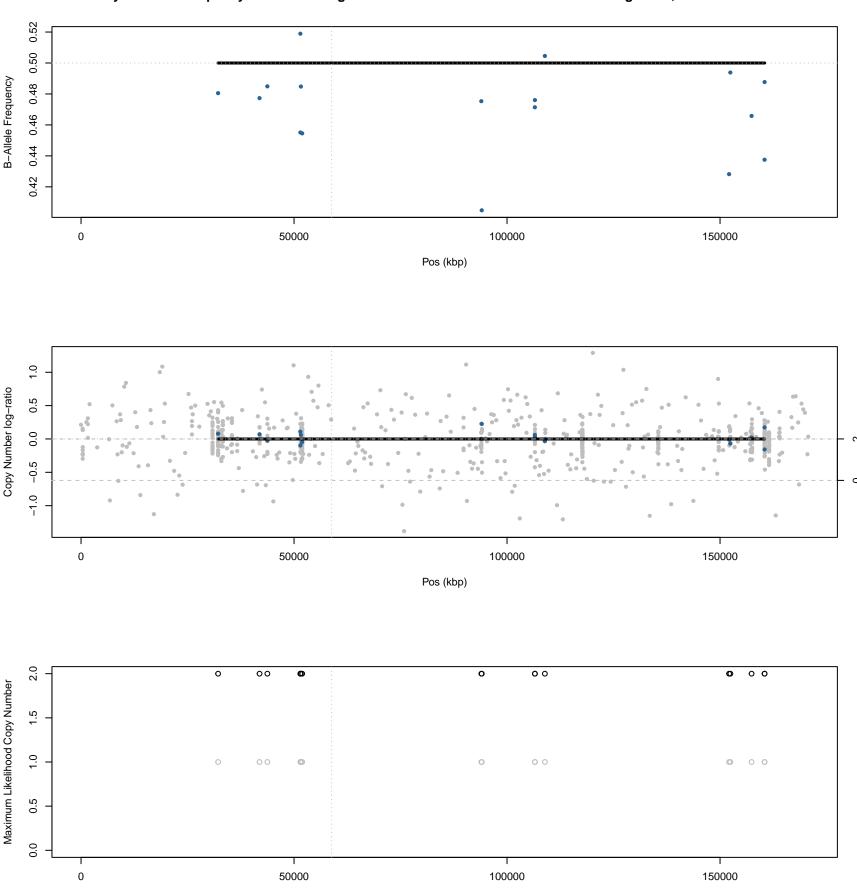
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 4



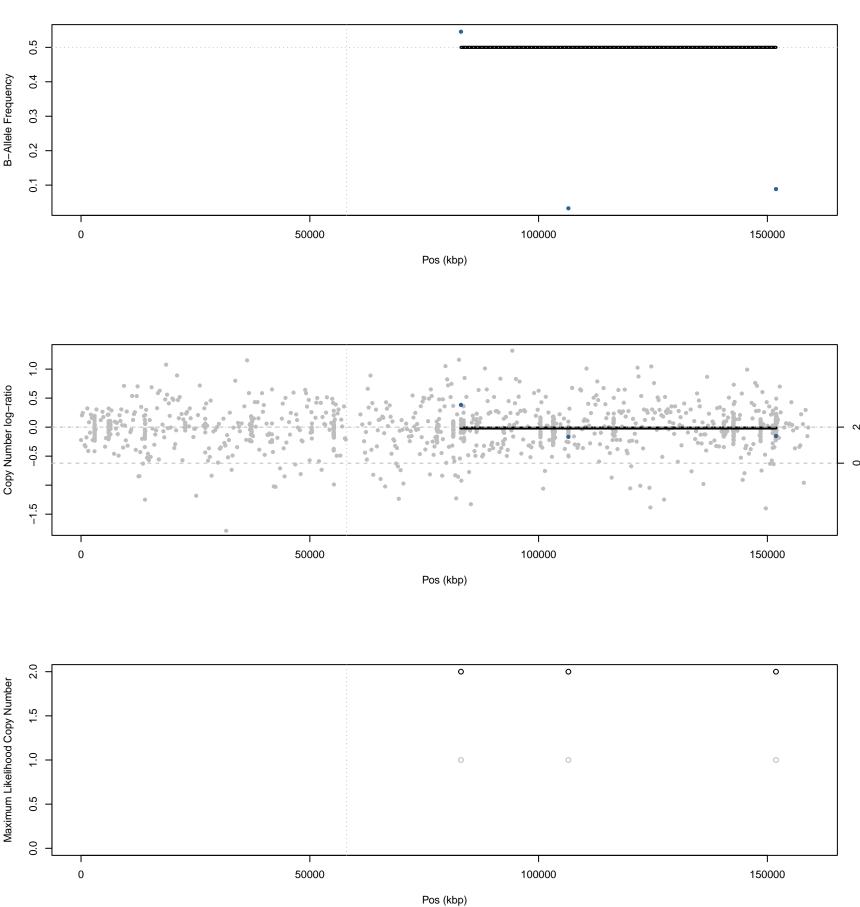
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 5



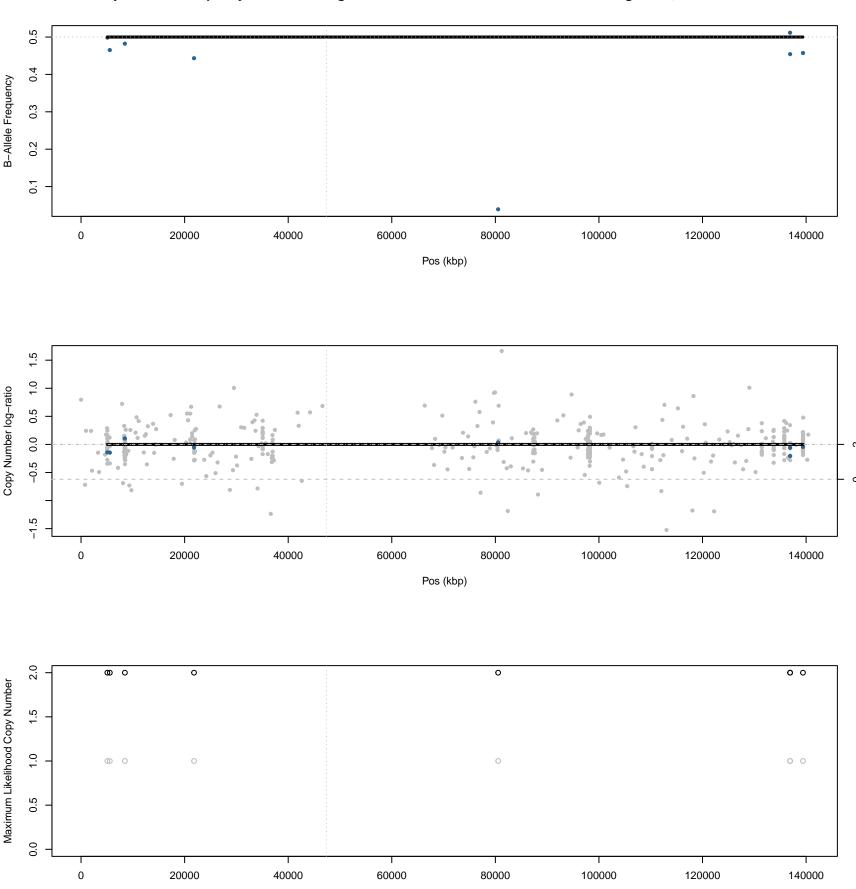
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 6



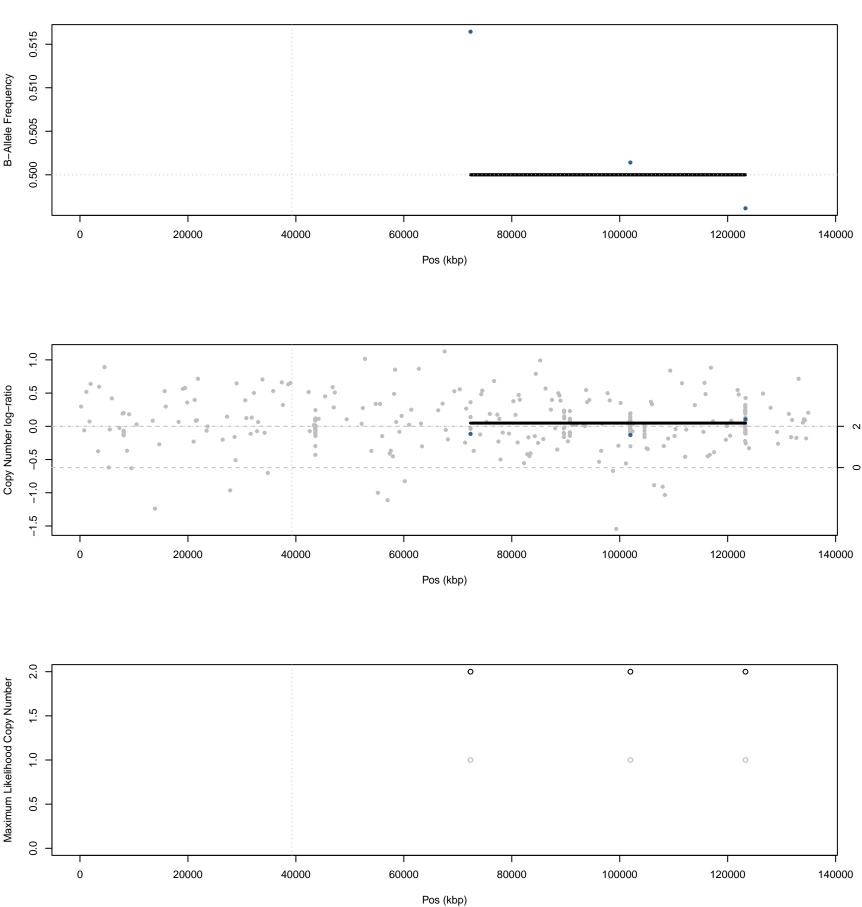
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 7



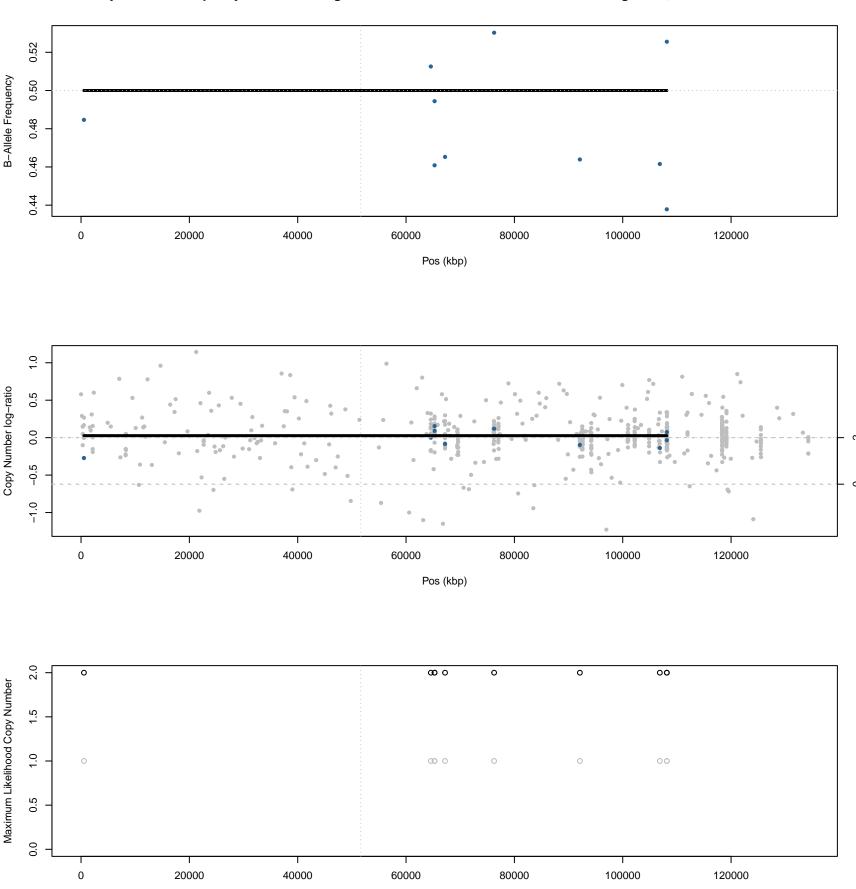
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 9



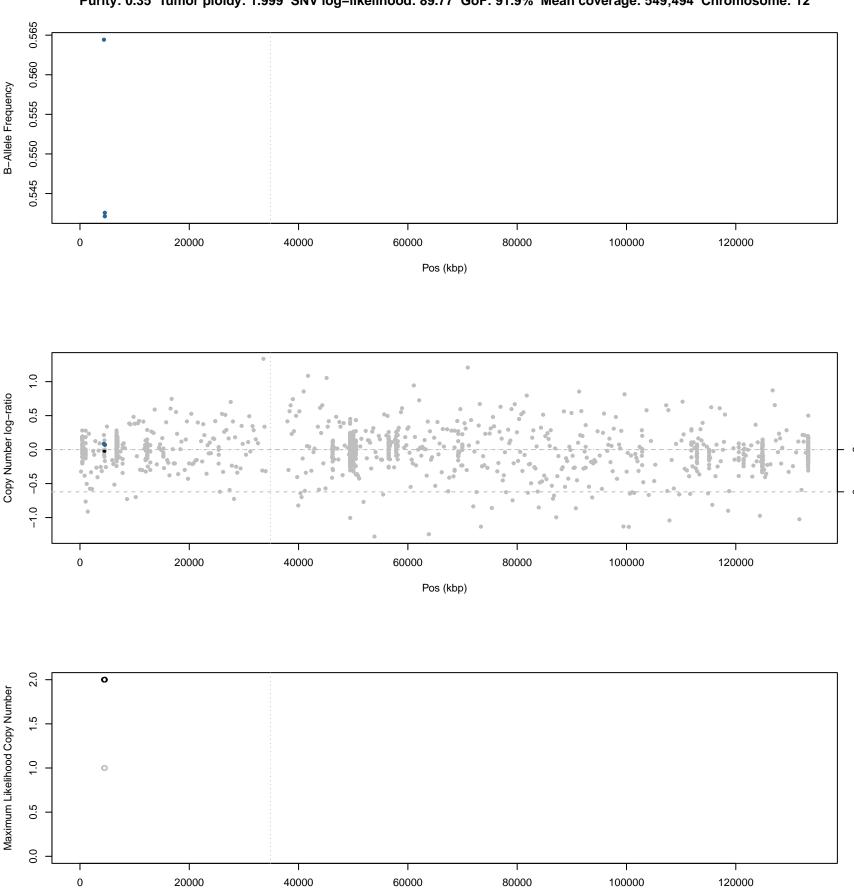
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 10



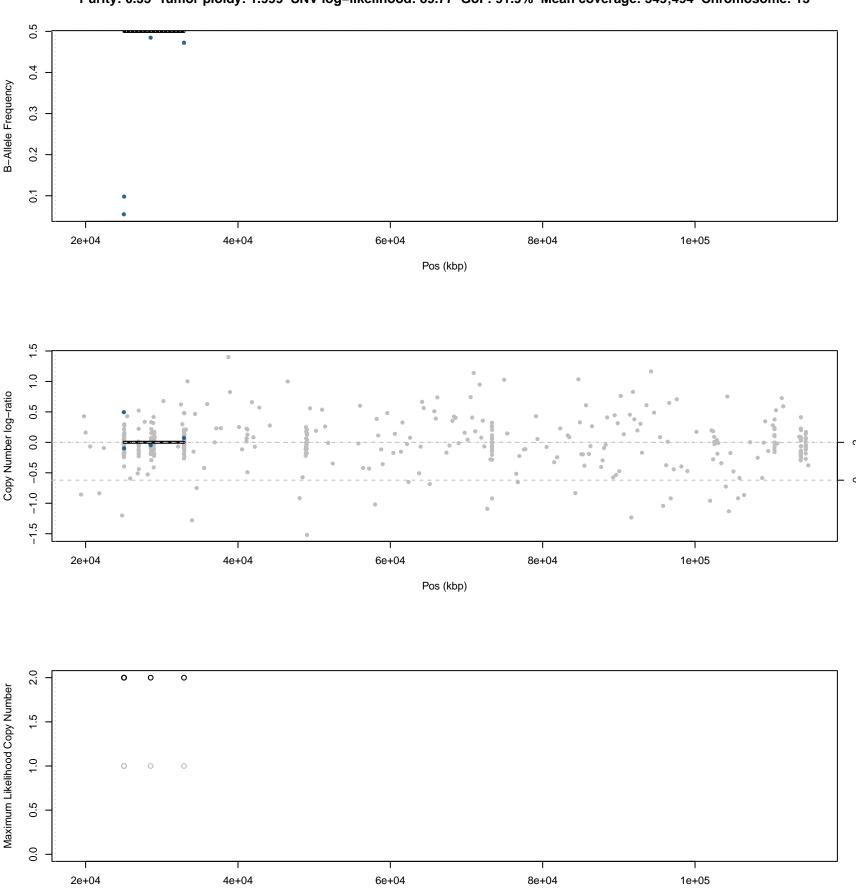
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 11



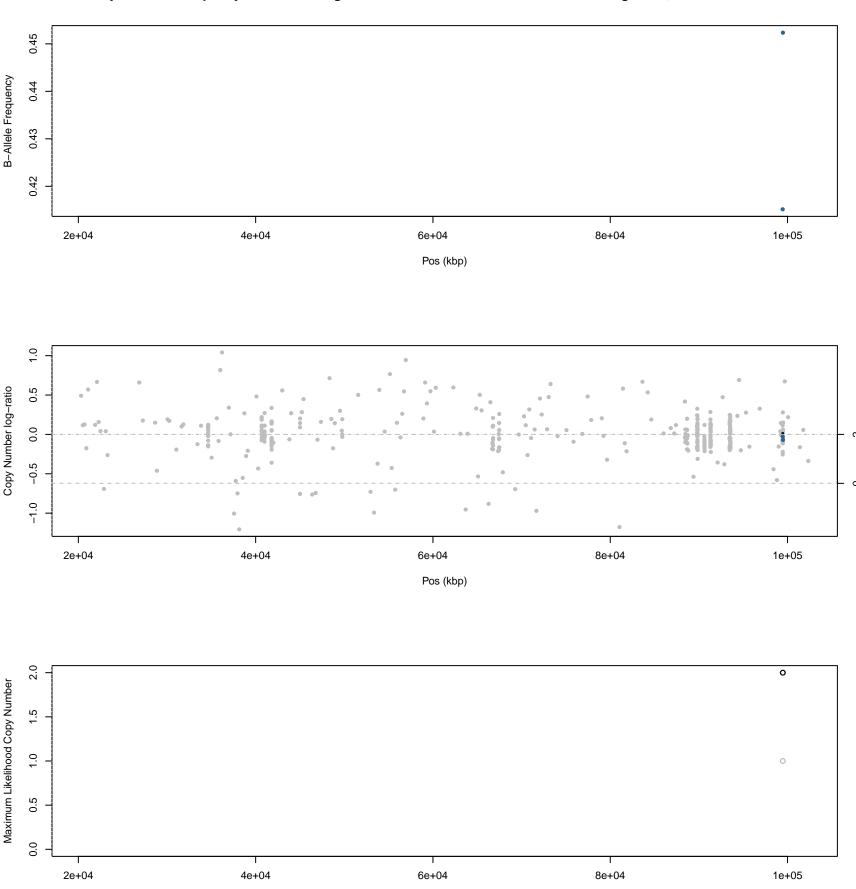
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 12



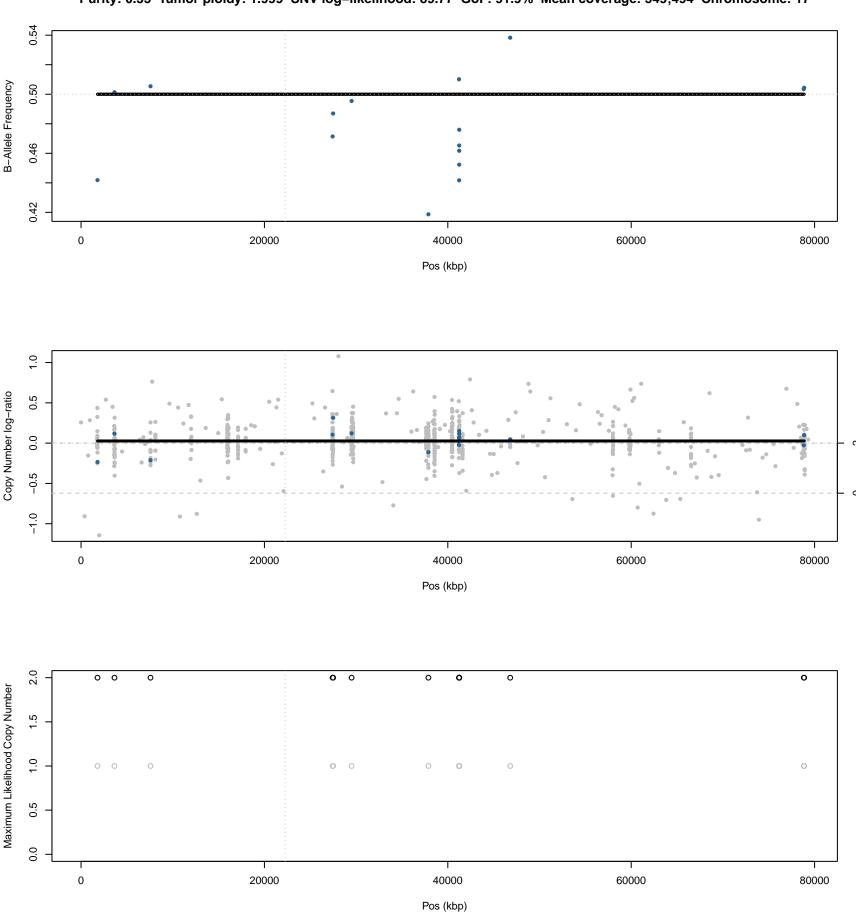
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 13



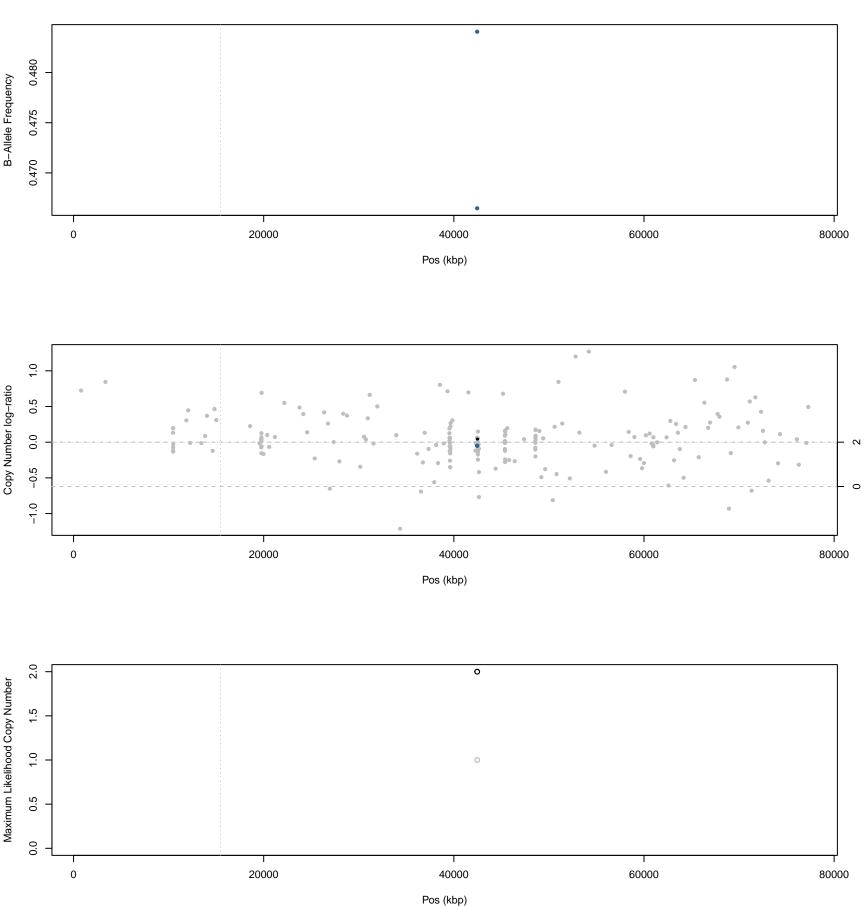
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 15



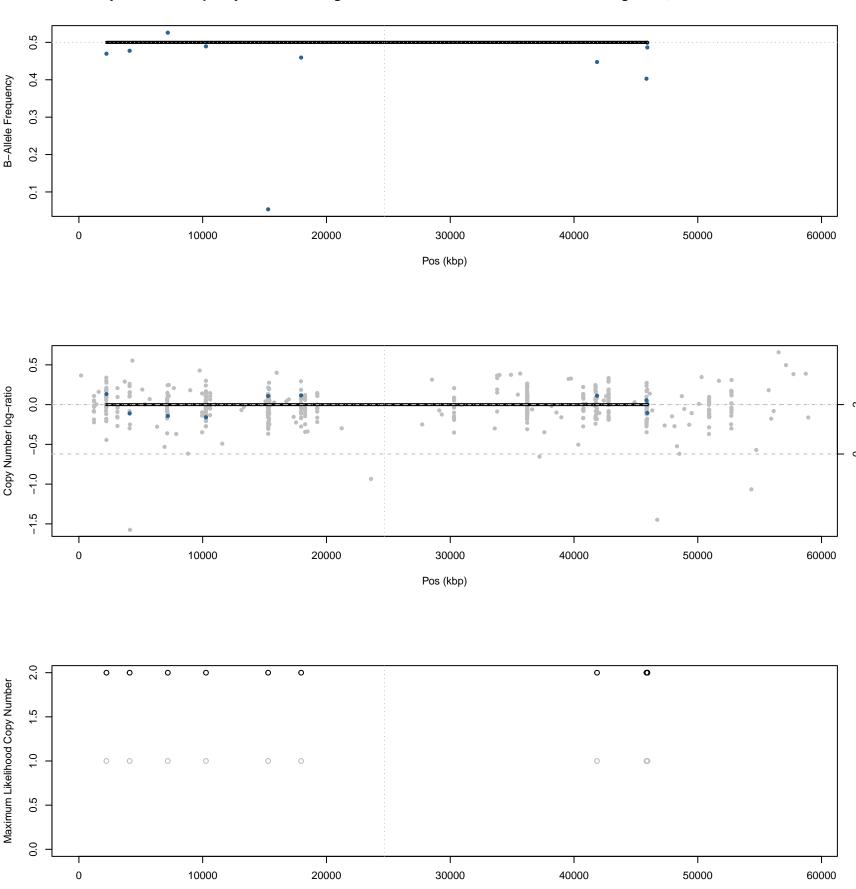
Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 17



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 18



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 19



Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549;494 Chromosome: 20

