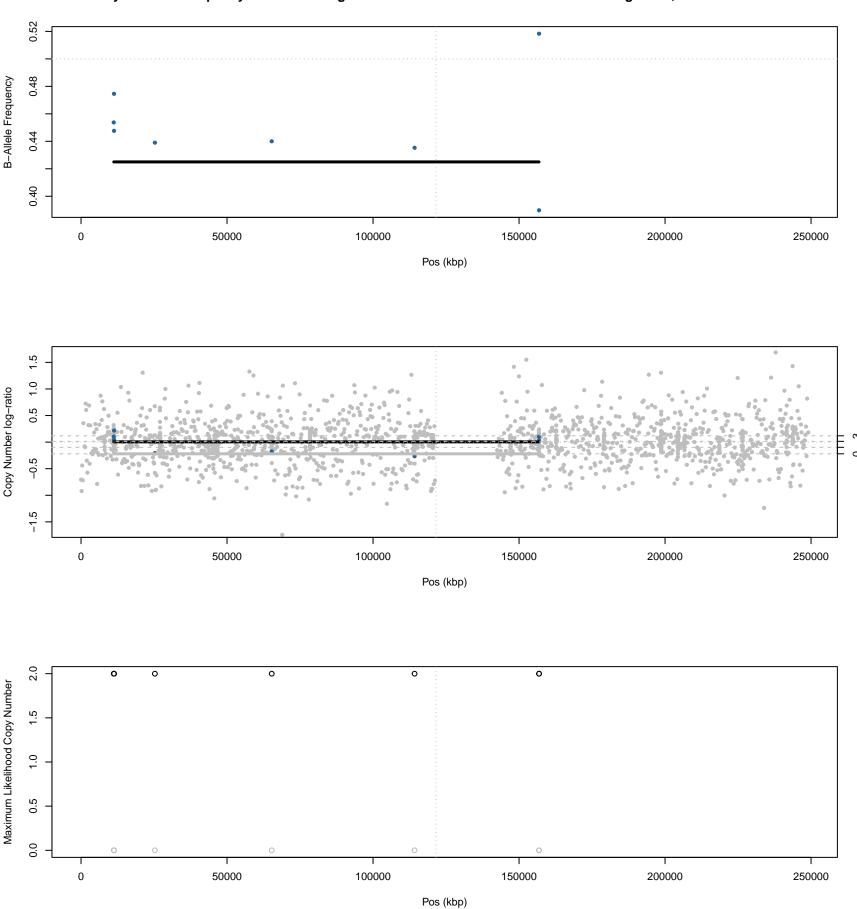
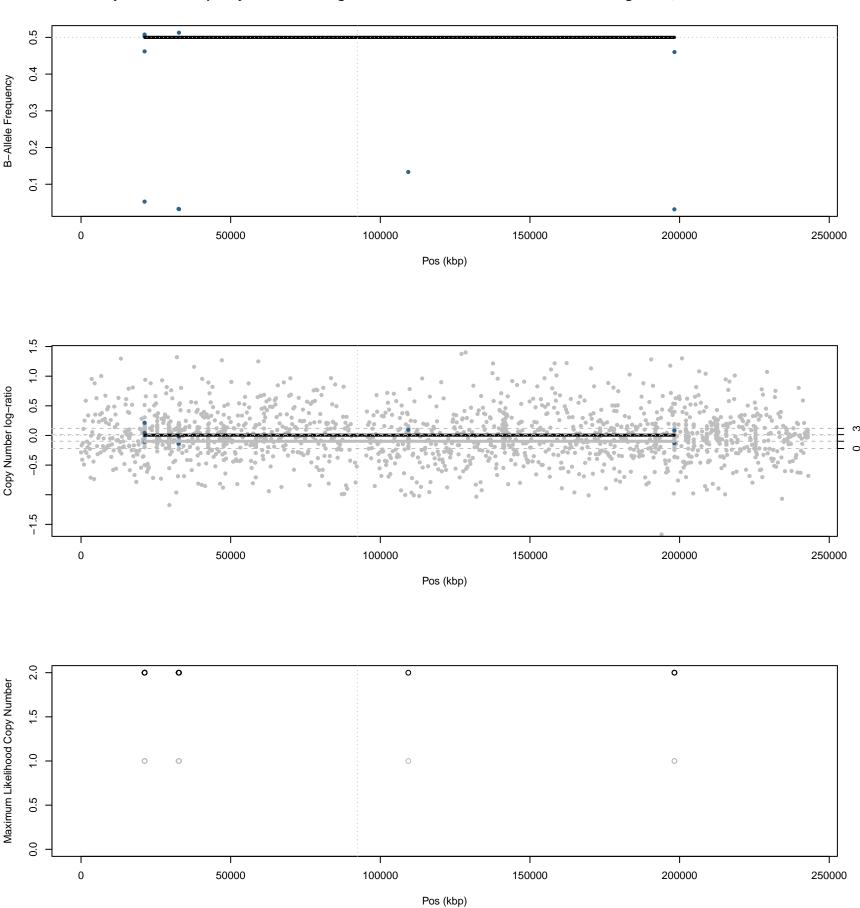
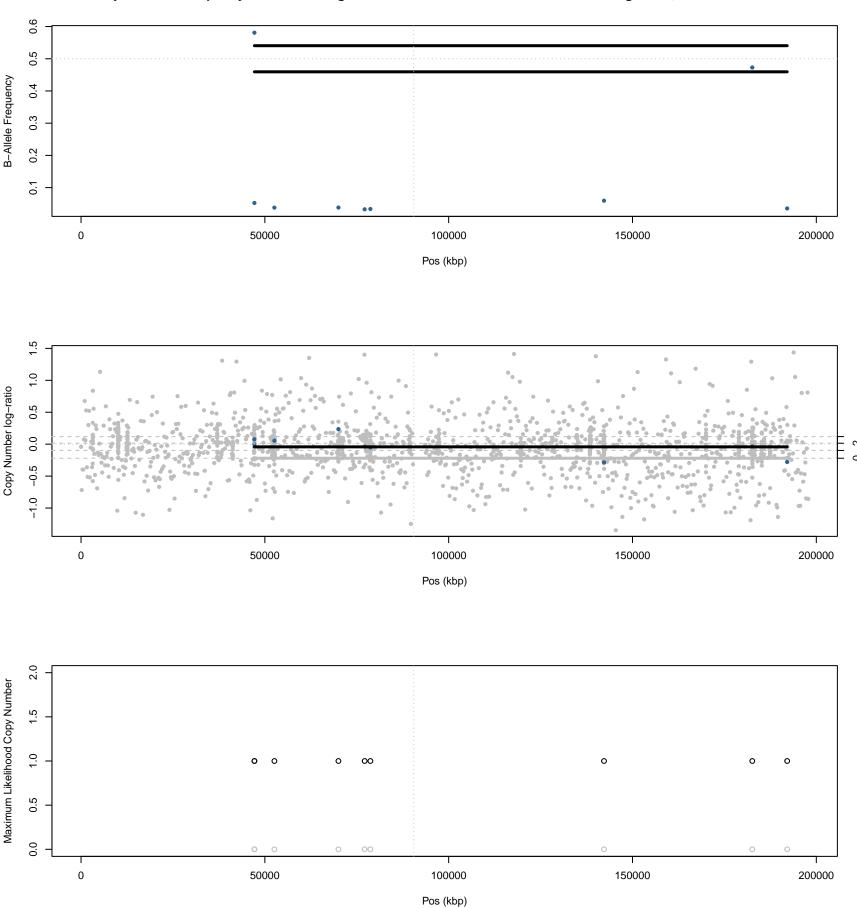
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 1



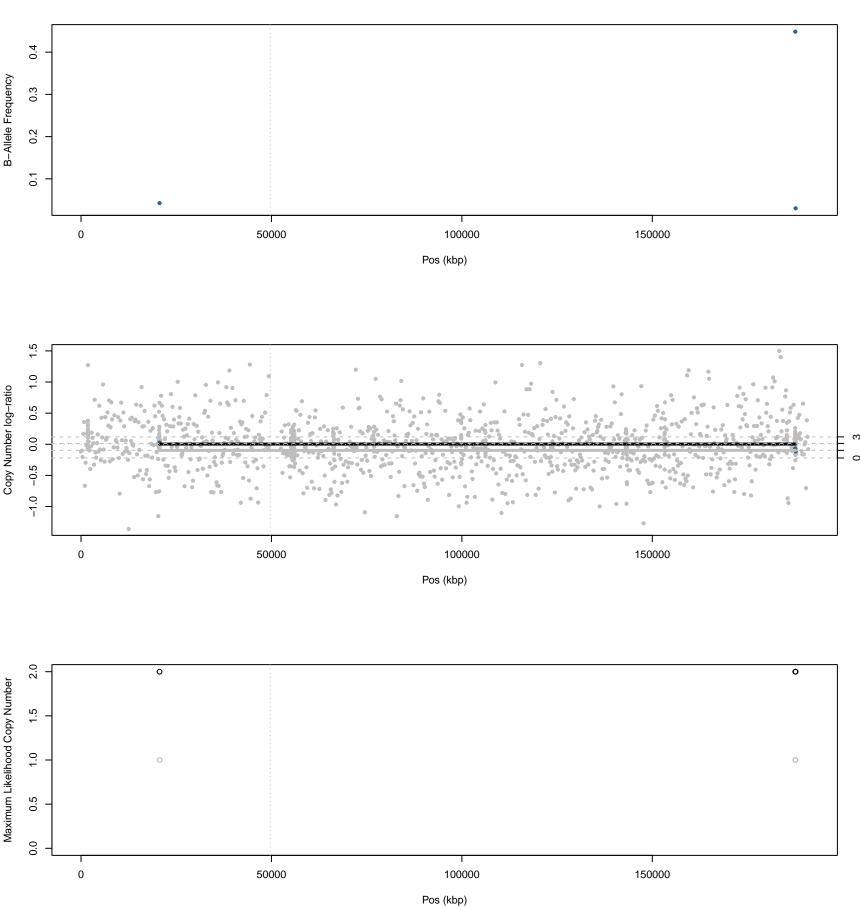
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 2



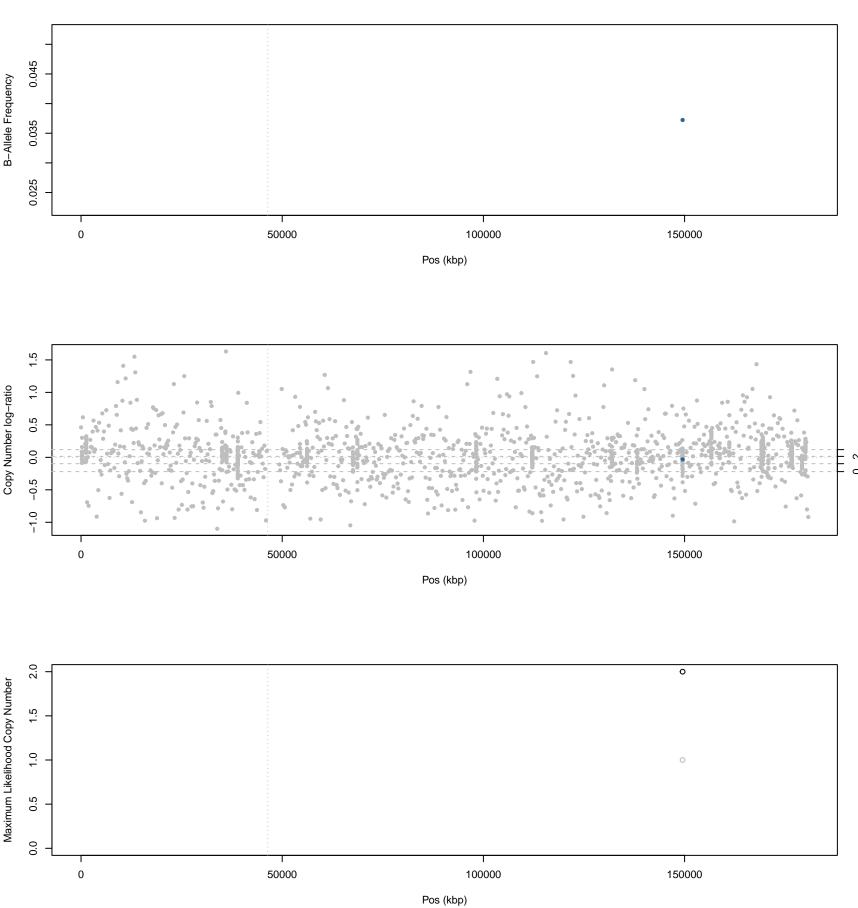
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 3



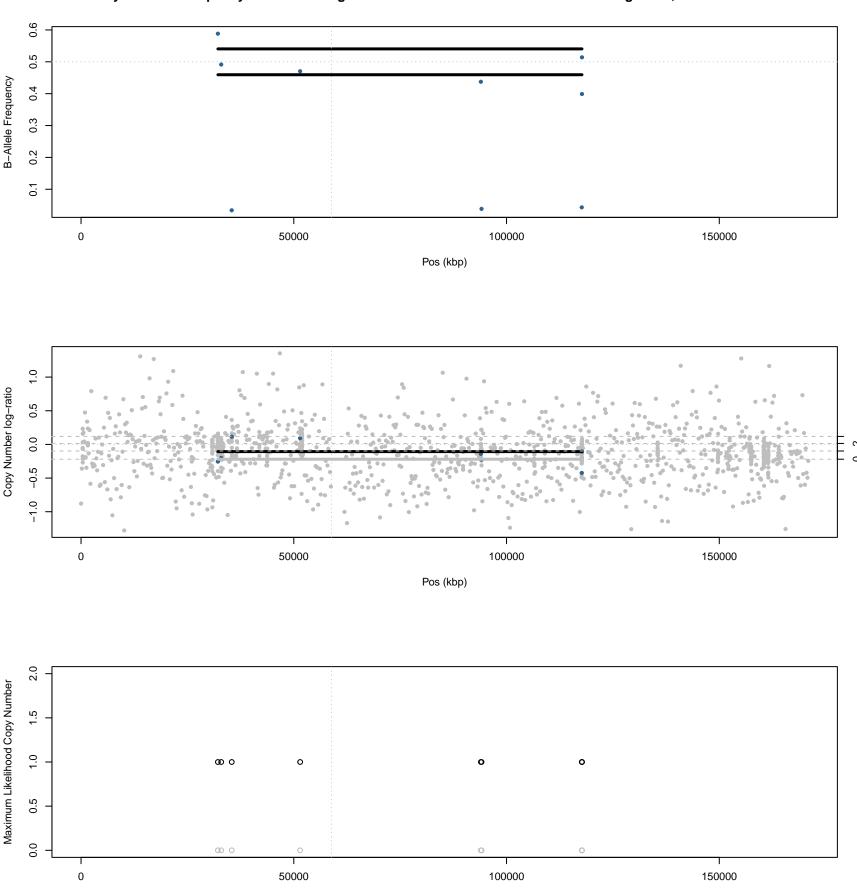
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 4



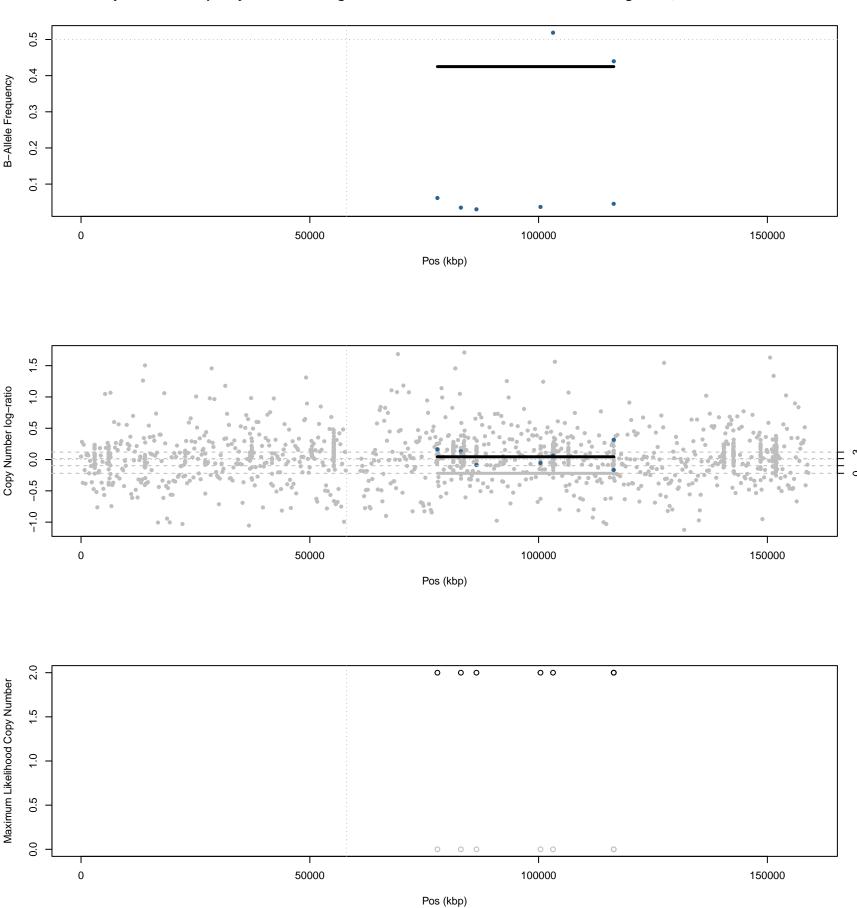
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 5



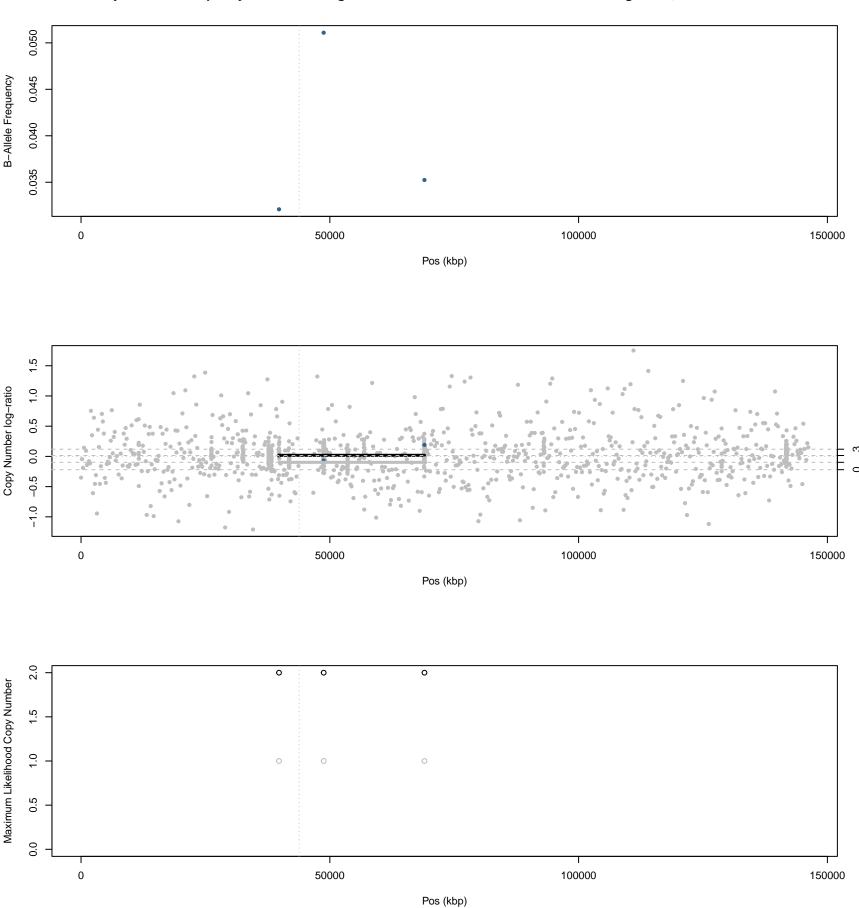
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 6



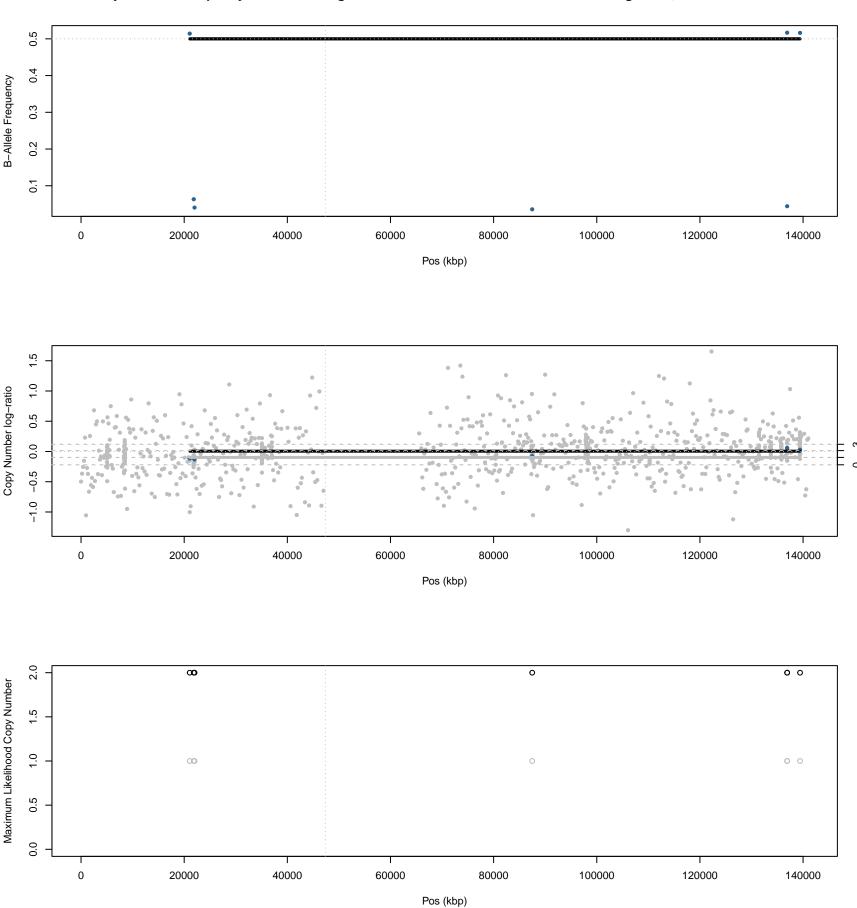
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 7



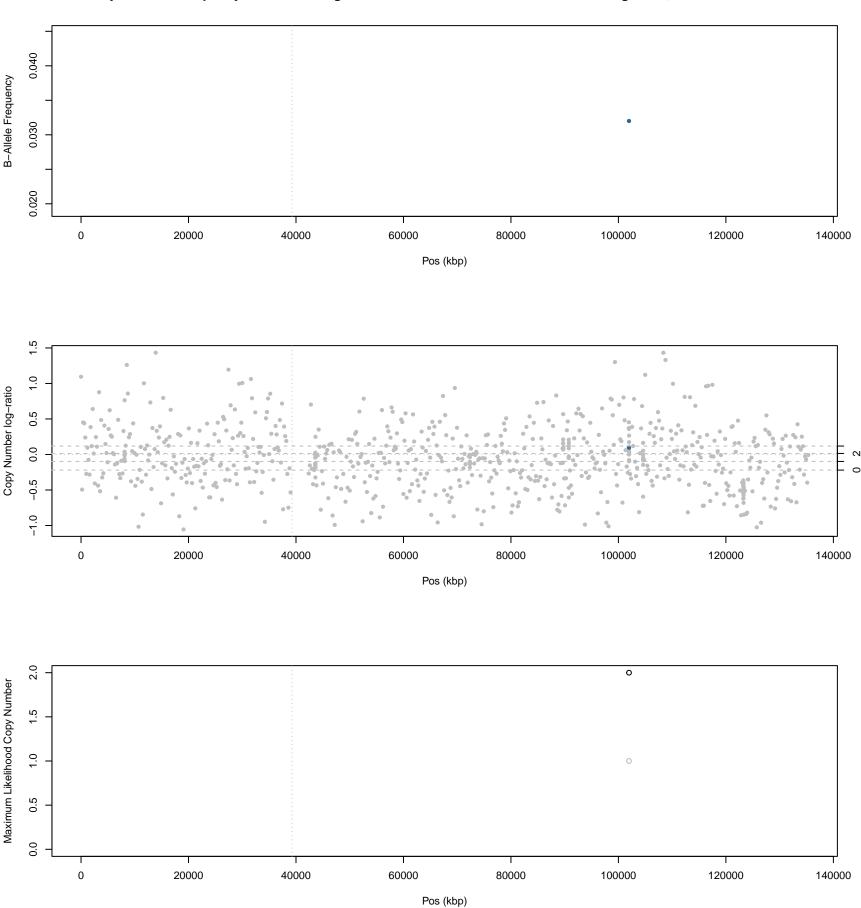
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 8



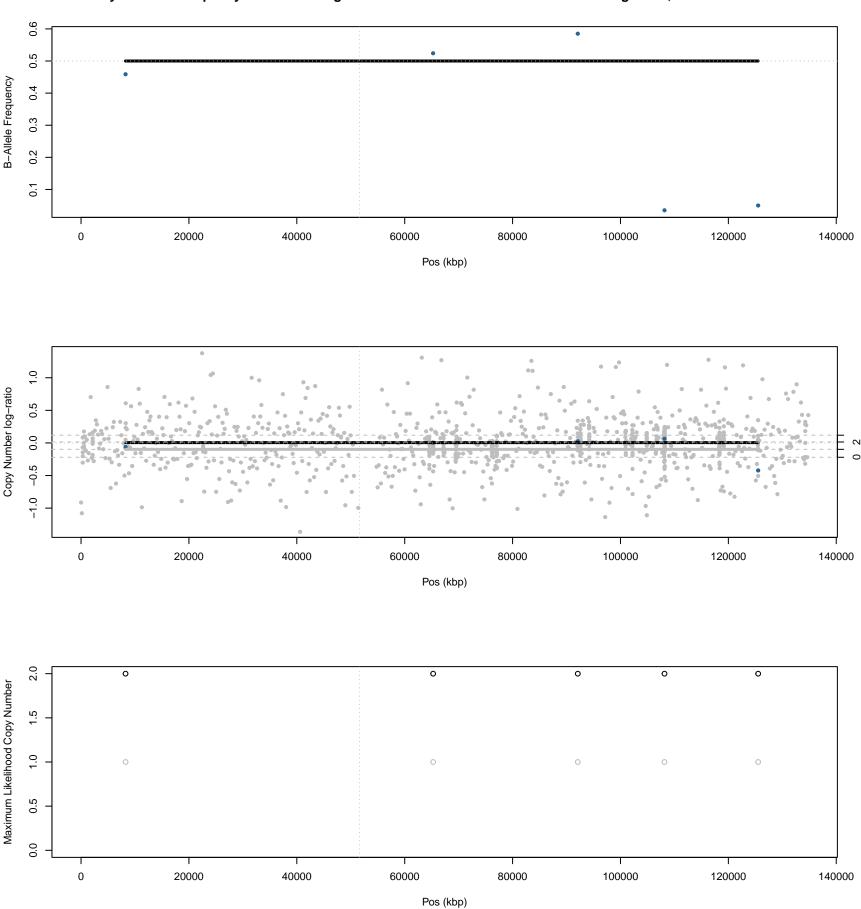
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 9



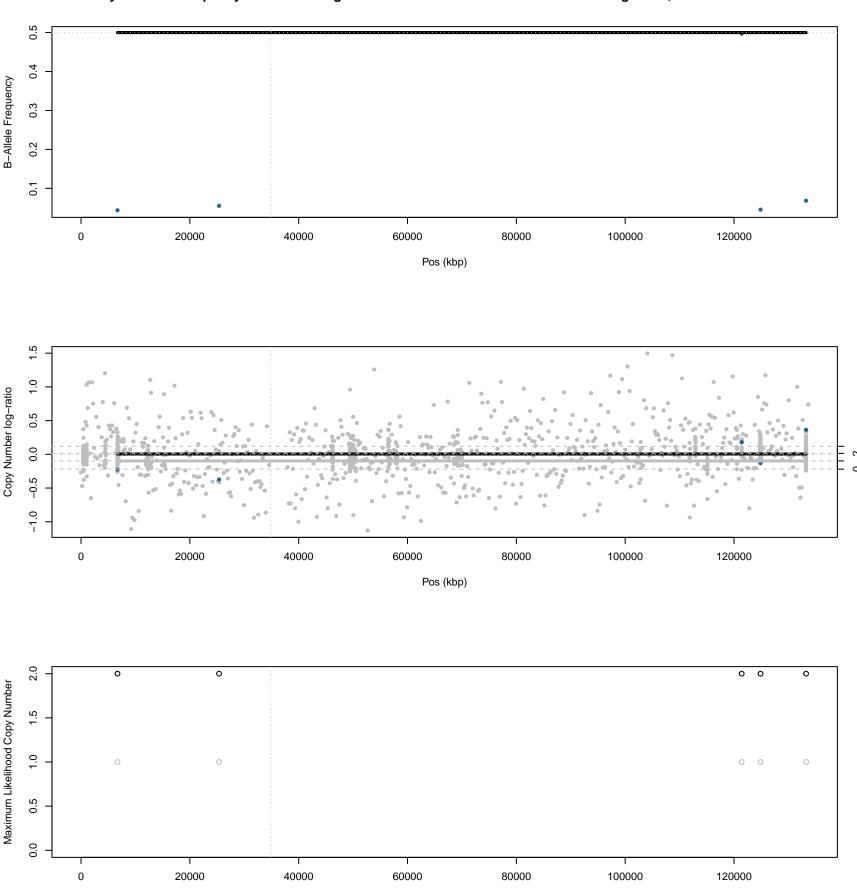
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 10



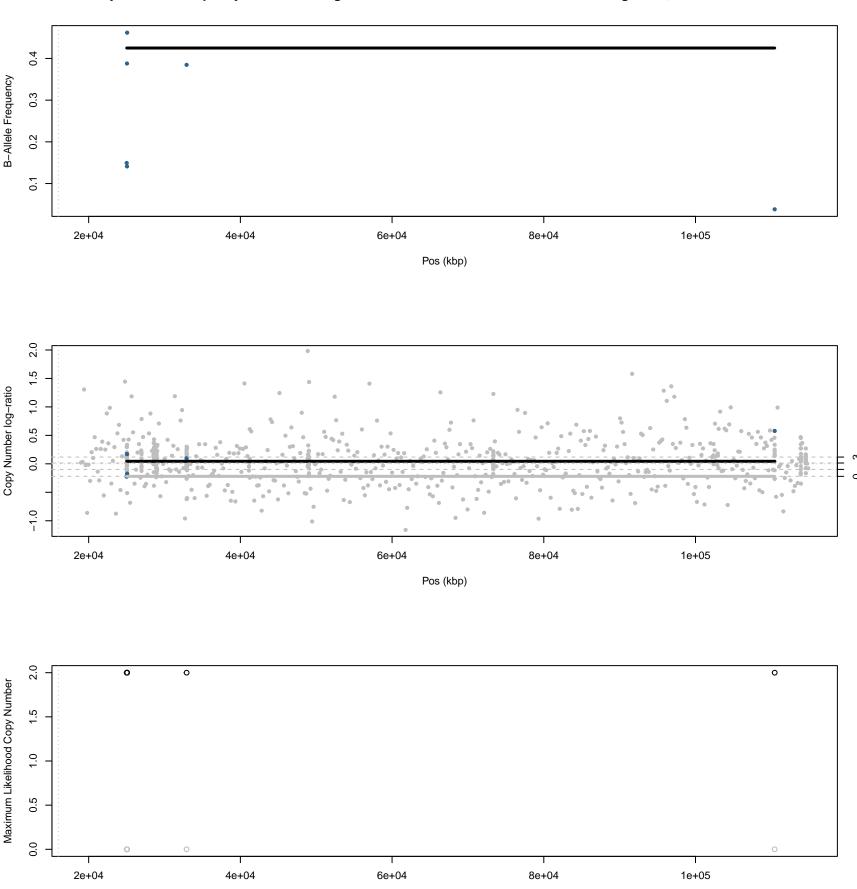
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 11



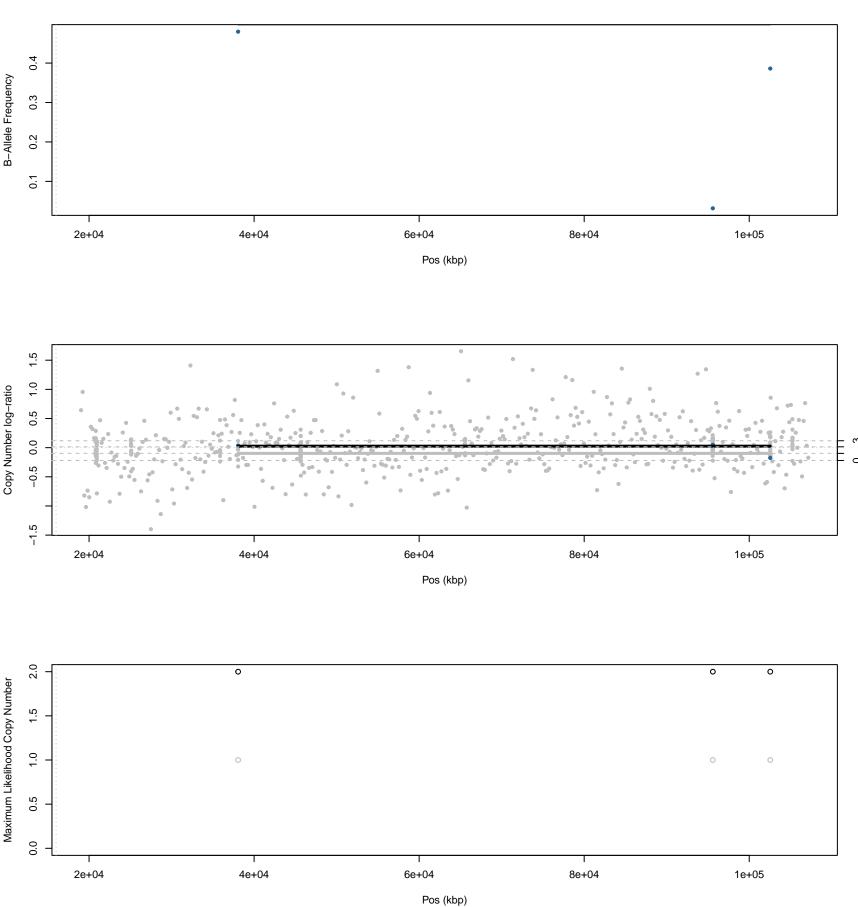
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 12



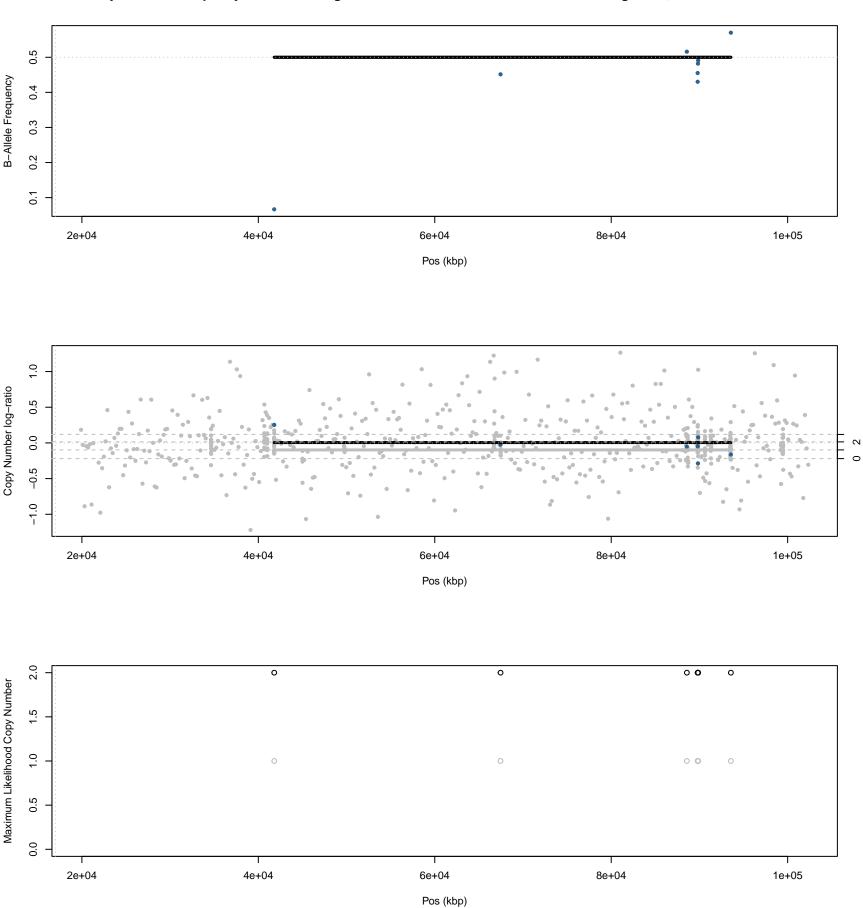
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 13



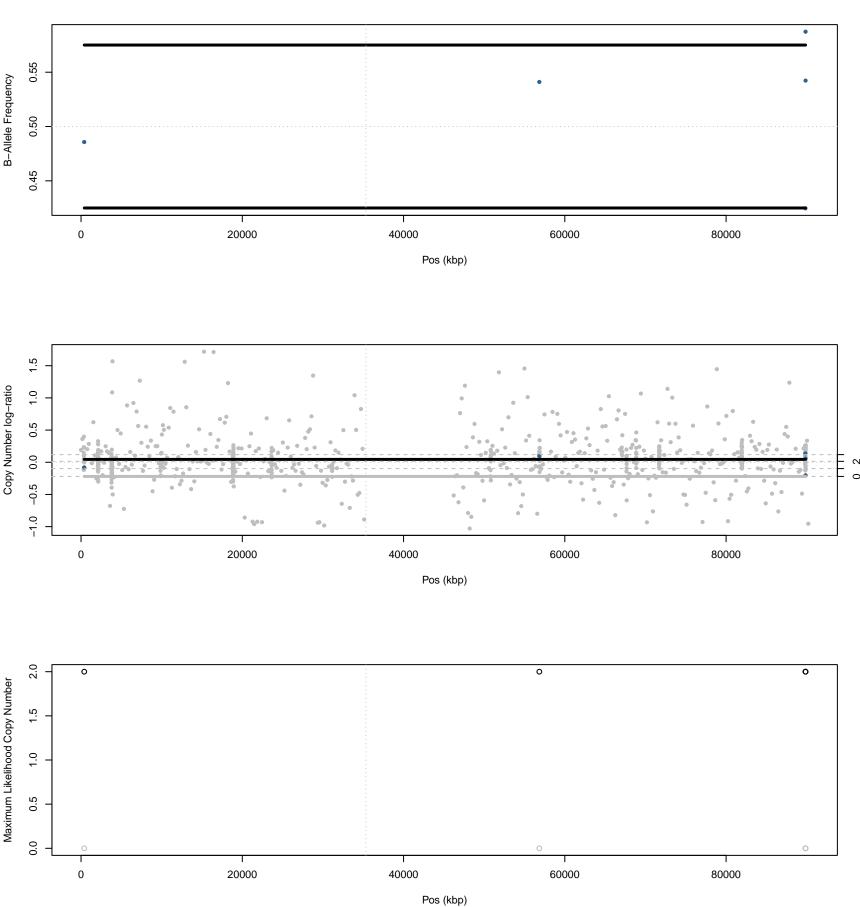
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 14



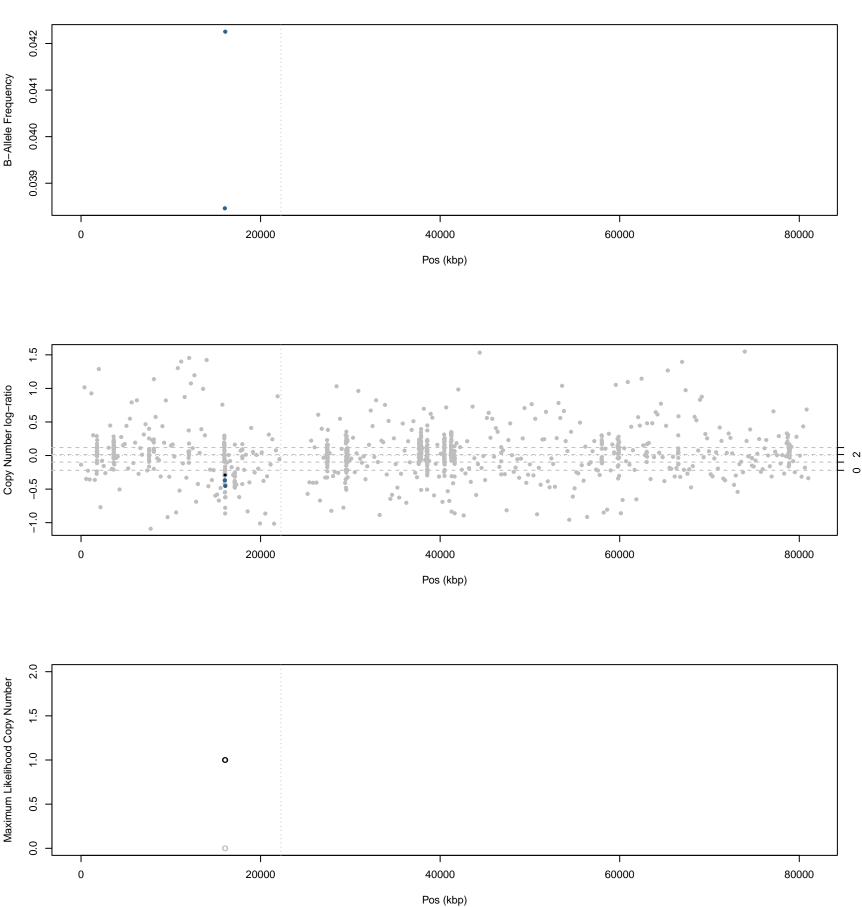
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 15



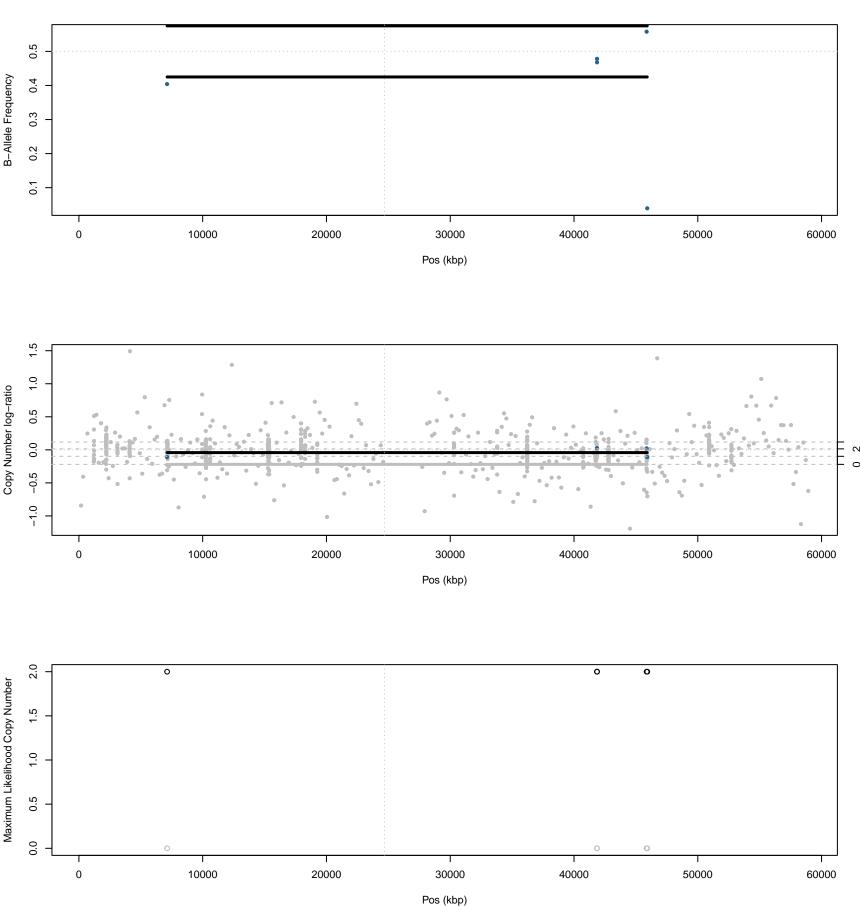
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 16



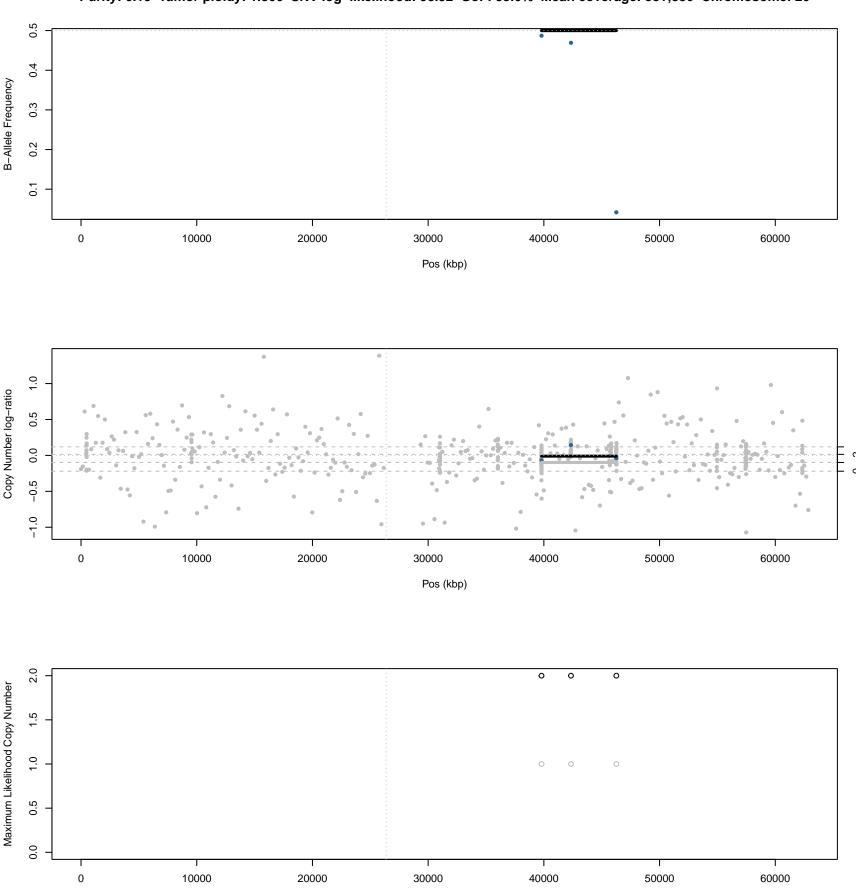
Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 17



Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 19



Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 20



Purity: 0.15 Tumor ploidy: 1.866 SNV log-likelihood: 98.32 GoF: 95.6% Mean coverage: 381;350 Chromosome: 22

