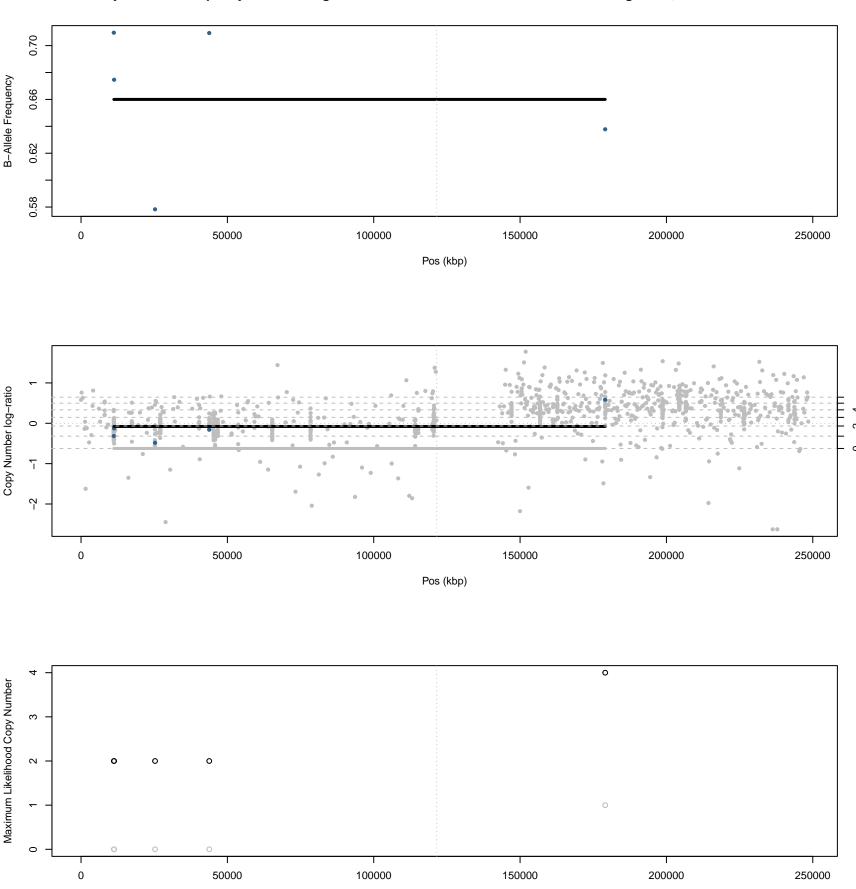
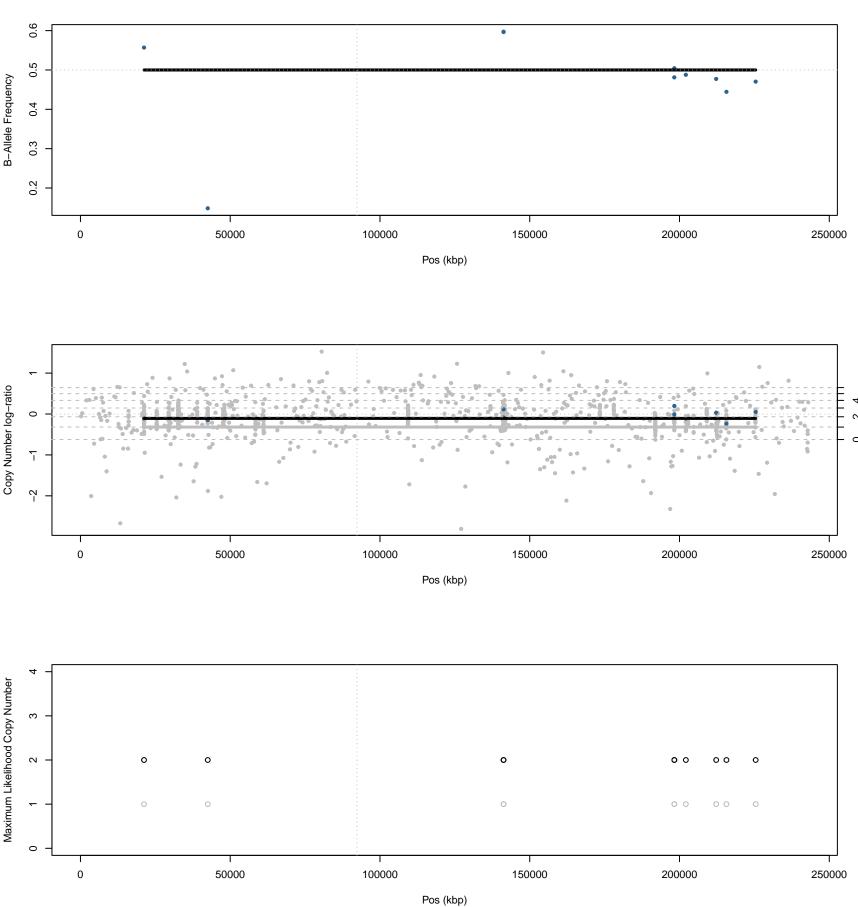
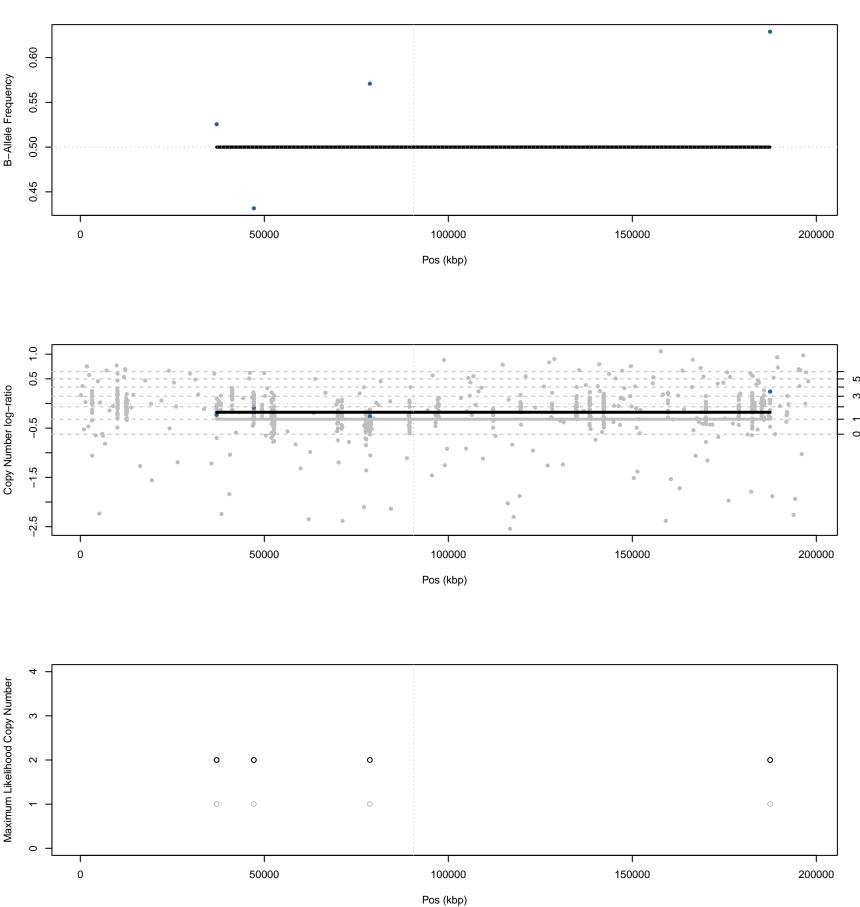
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 1



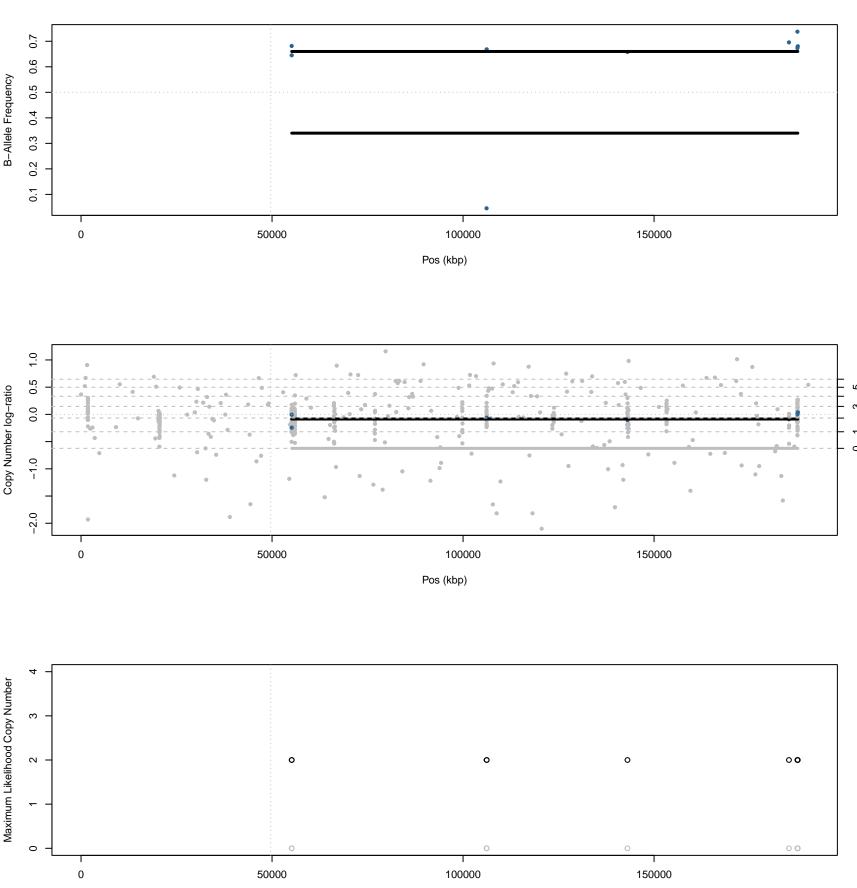
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 2



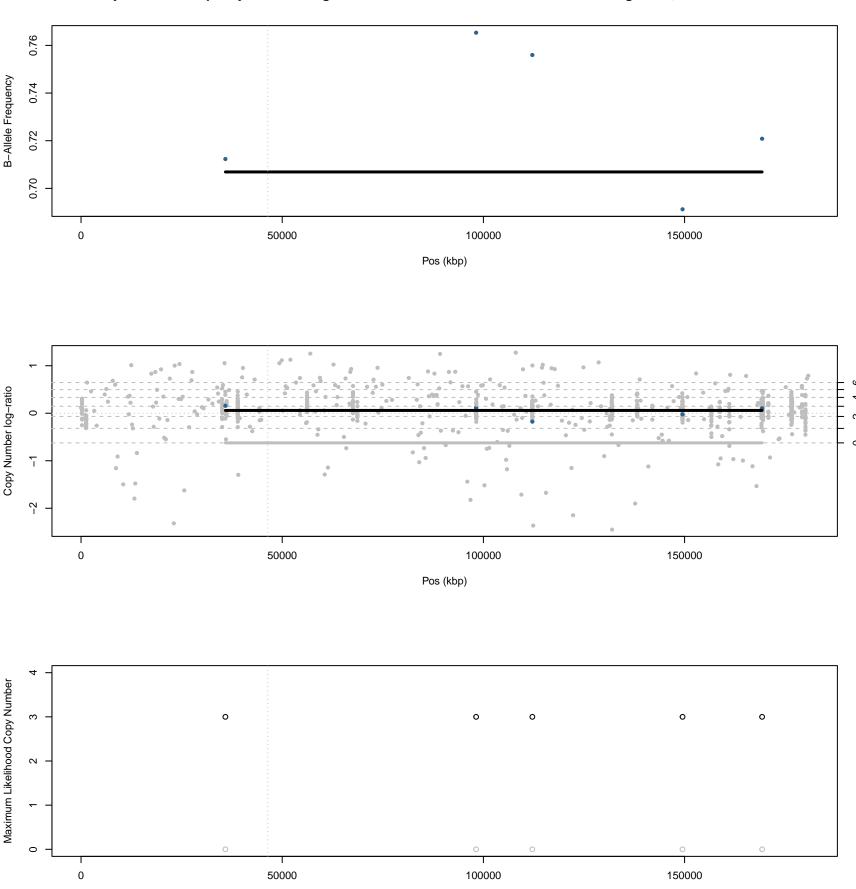
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 3



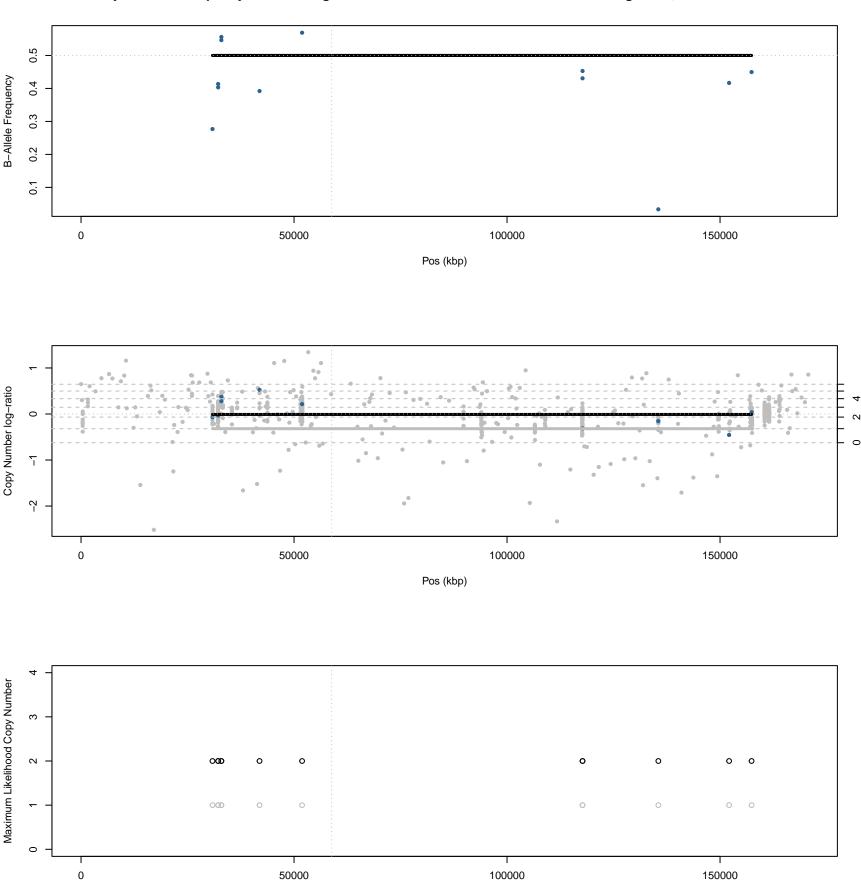
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 4



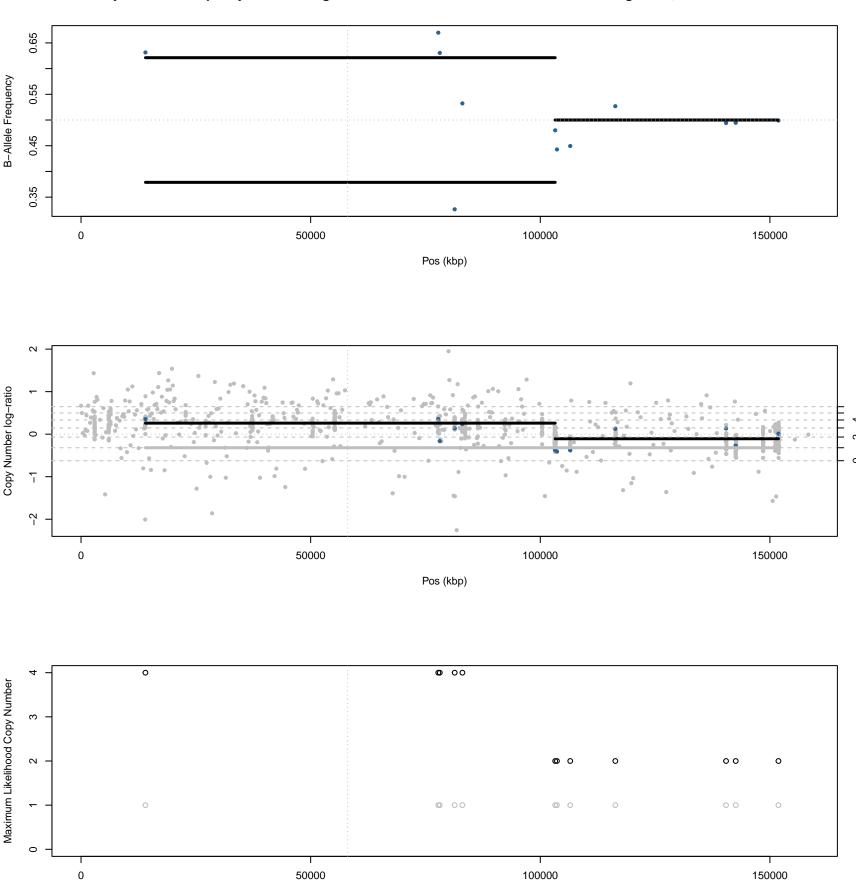
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 5



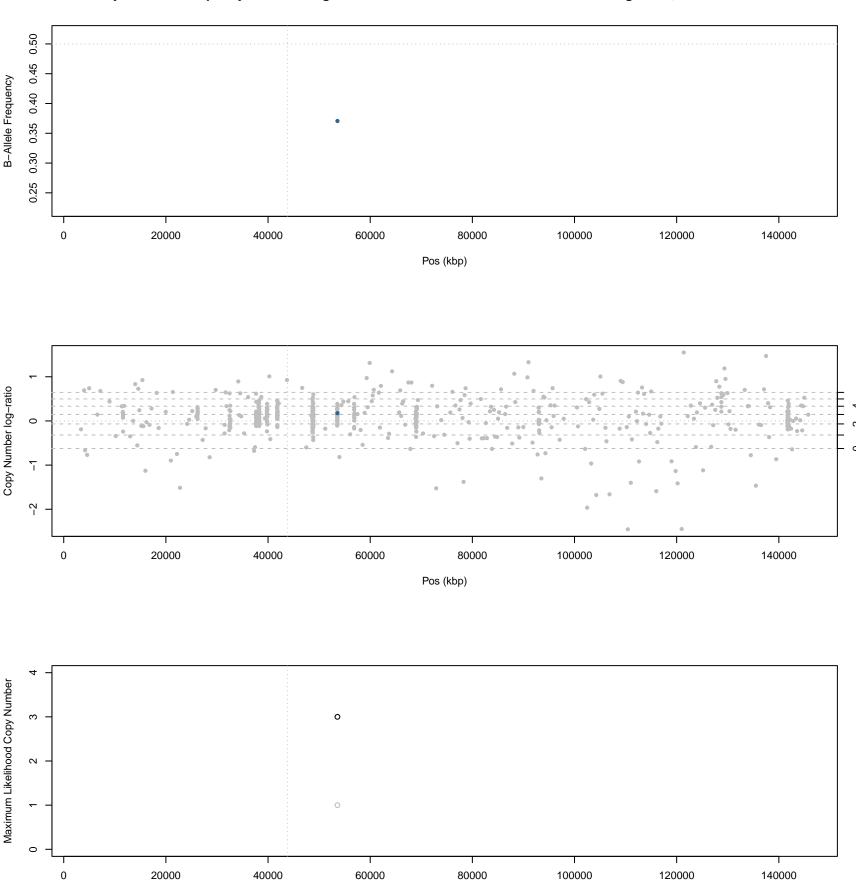
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 6



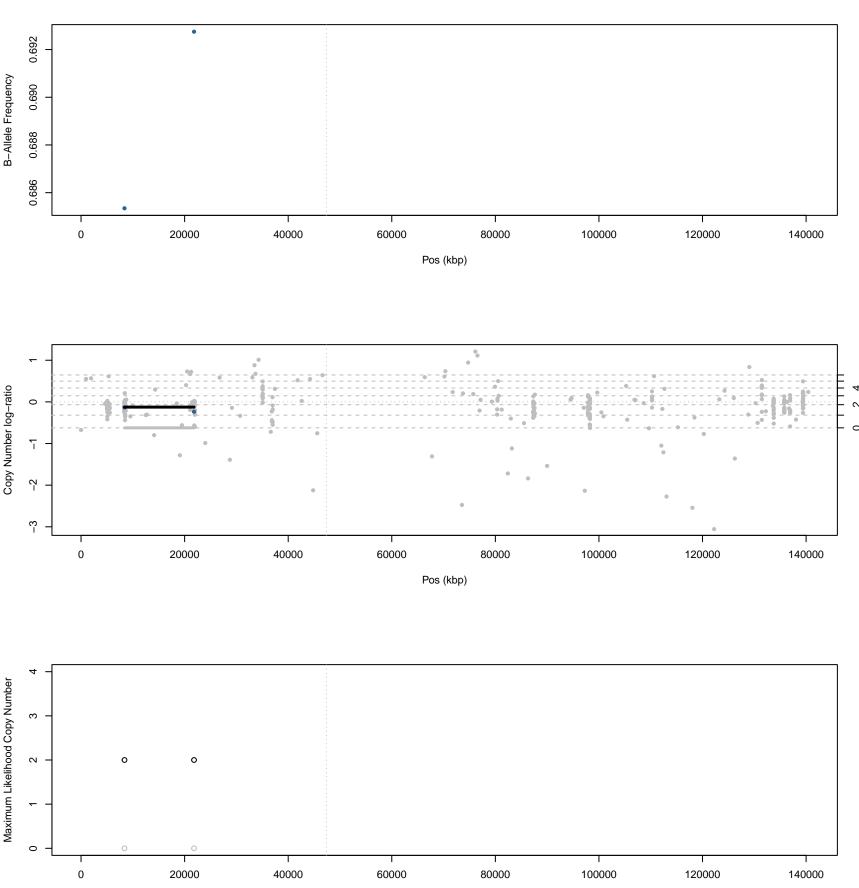
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 7



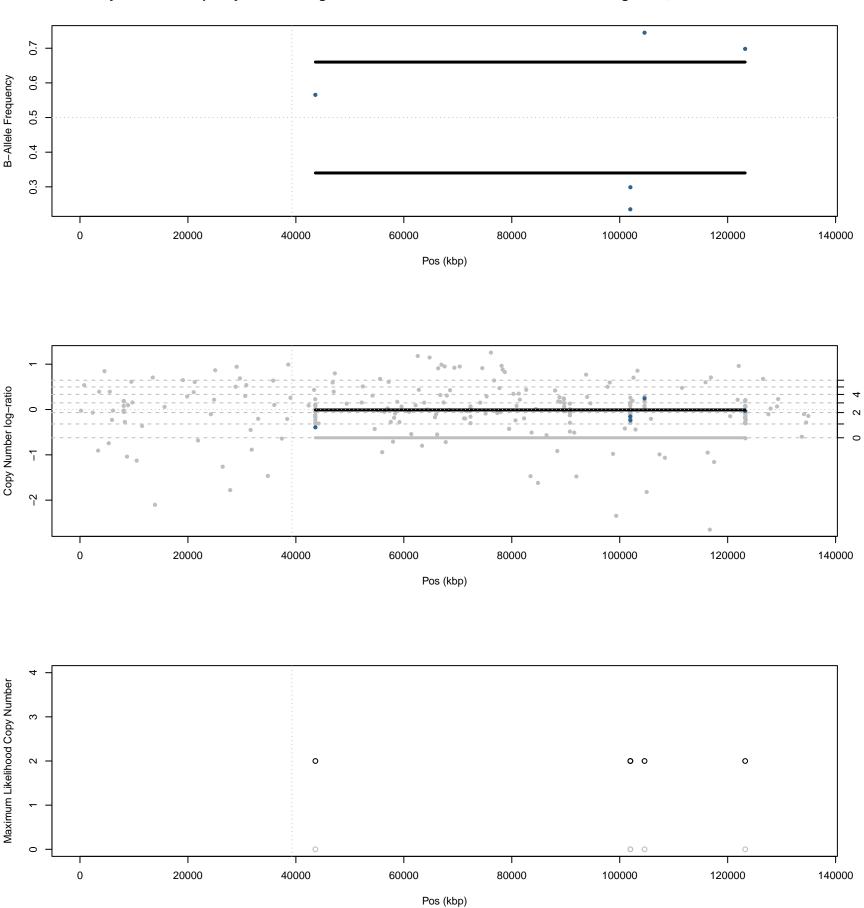
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 8

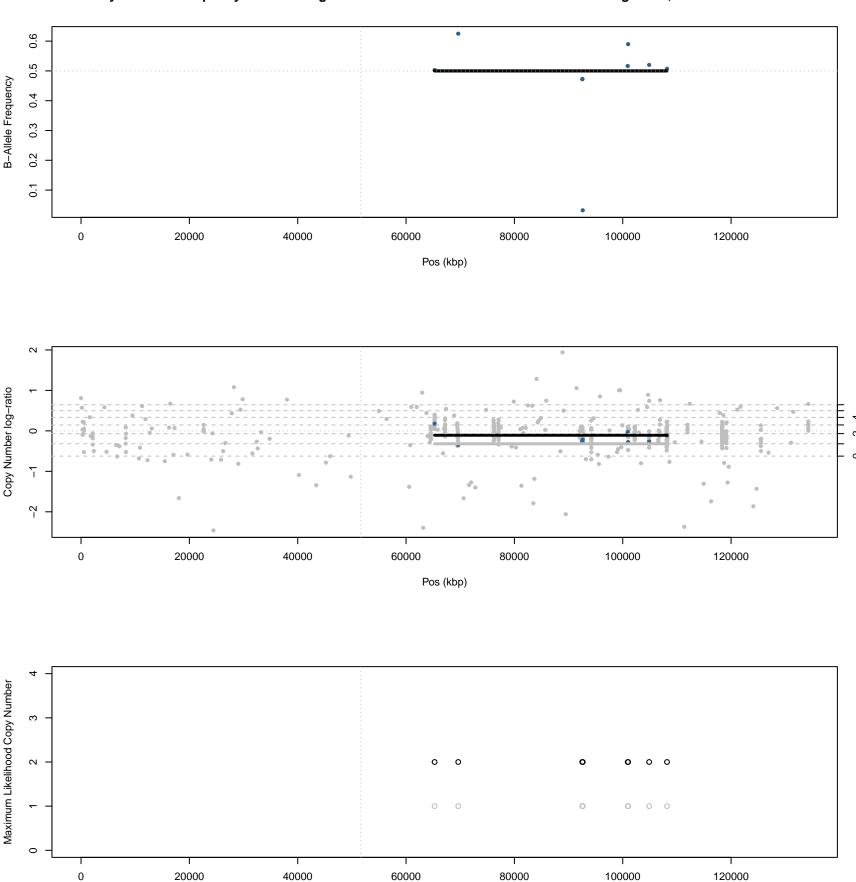


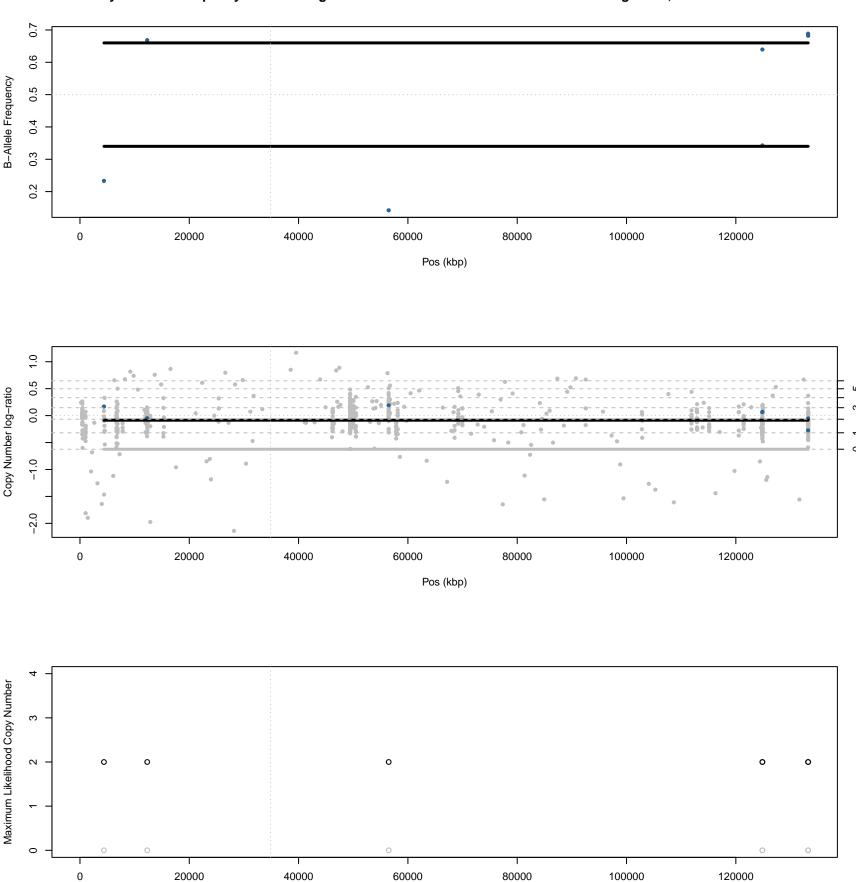
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 9



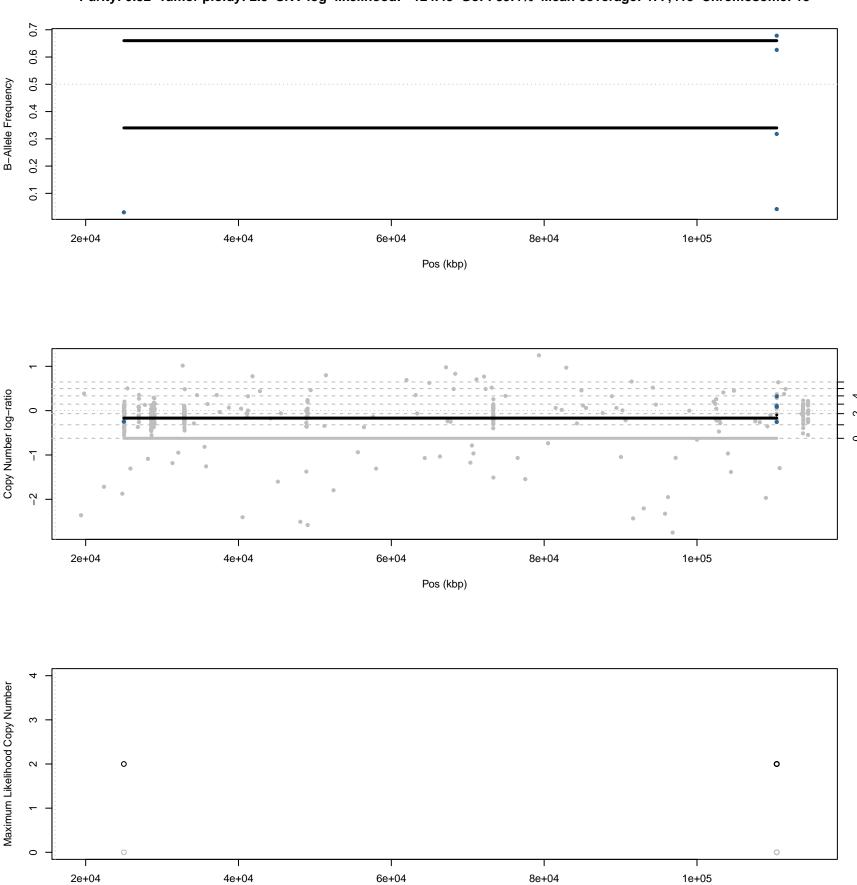
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 10



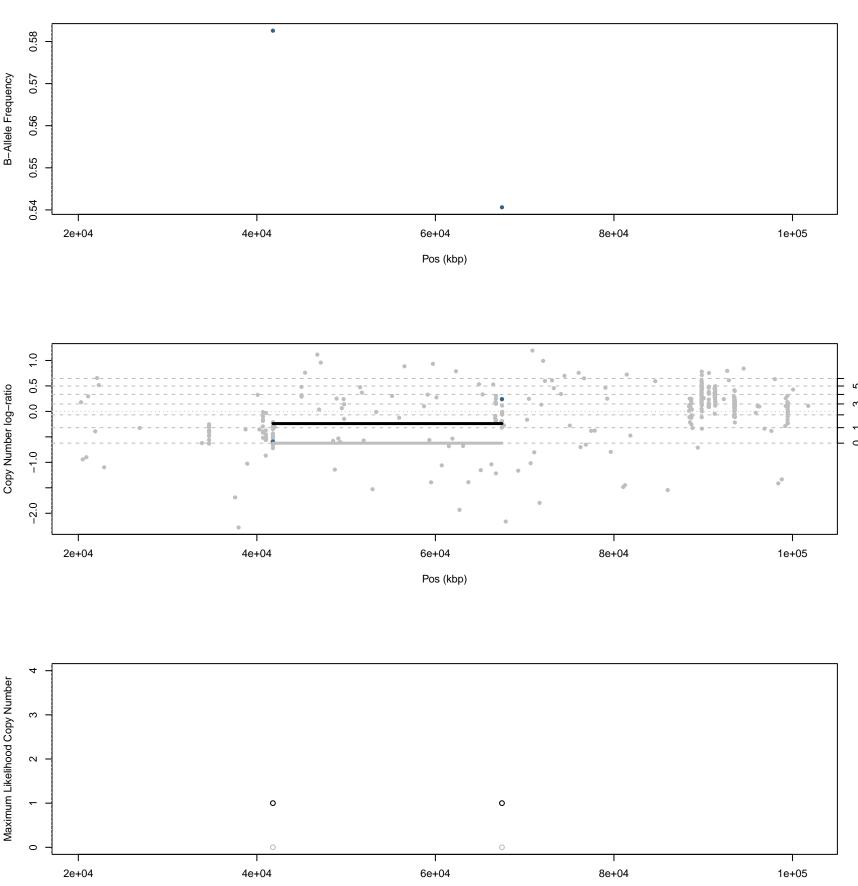




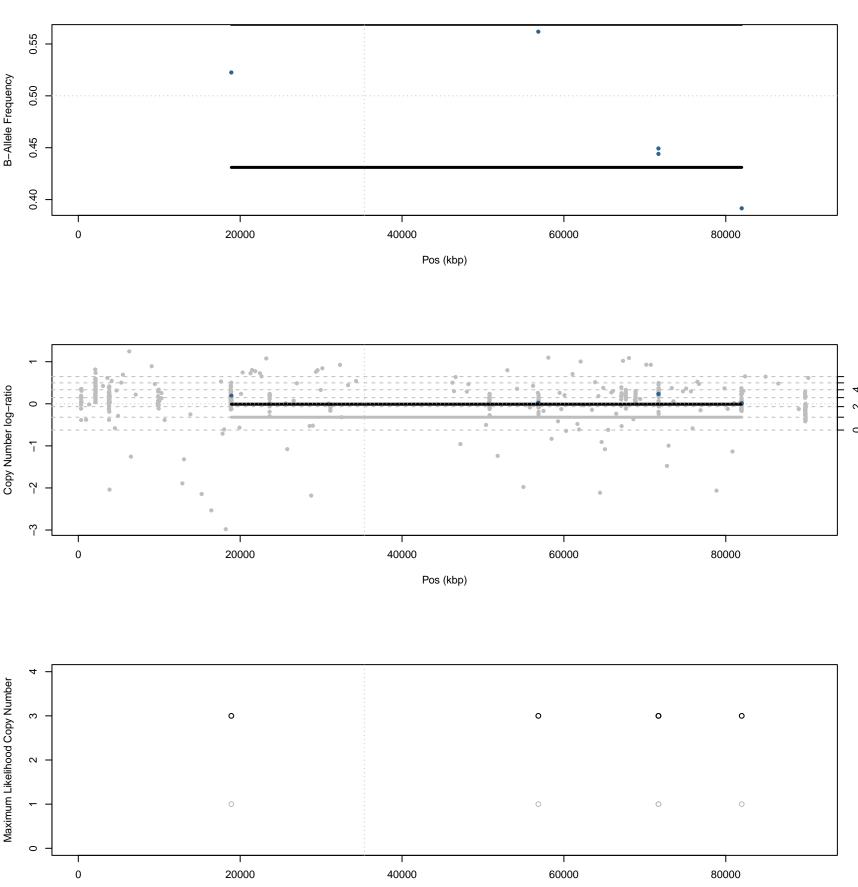
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 13

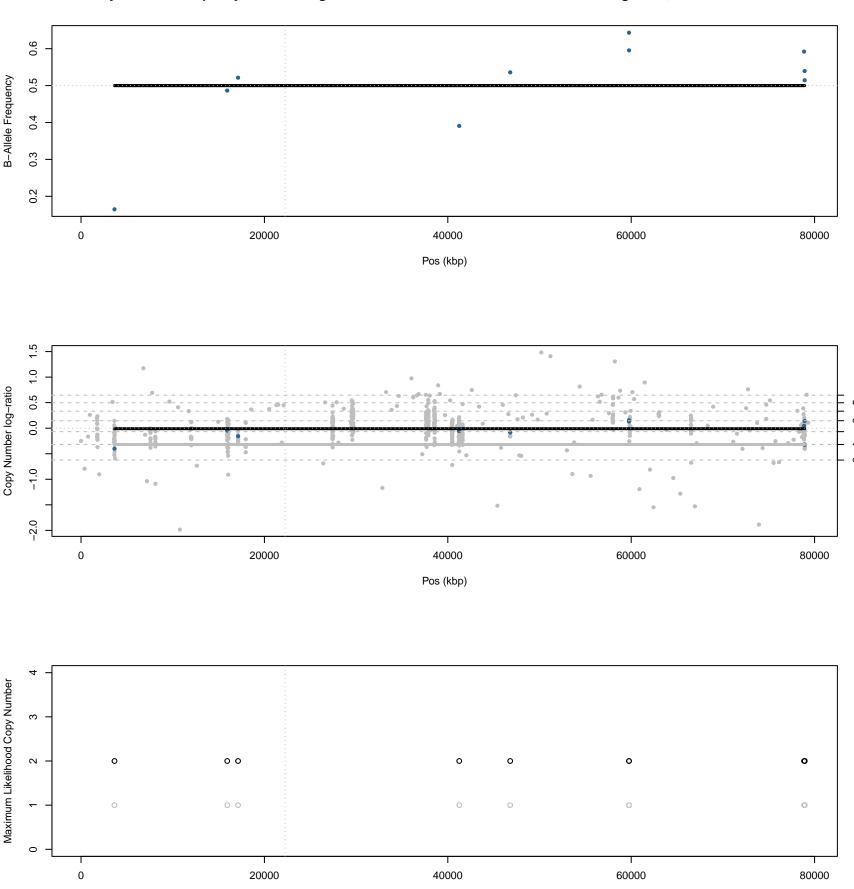


Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 15

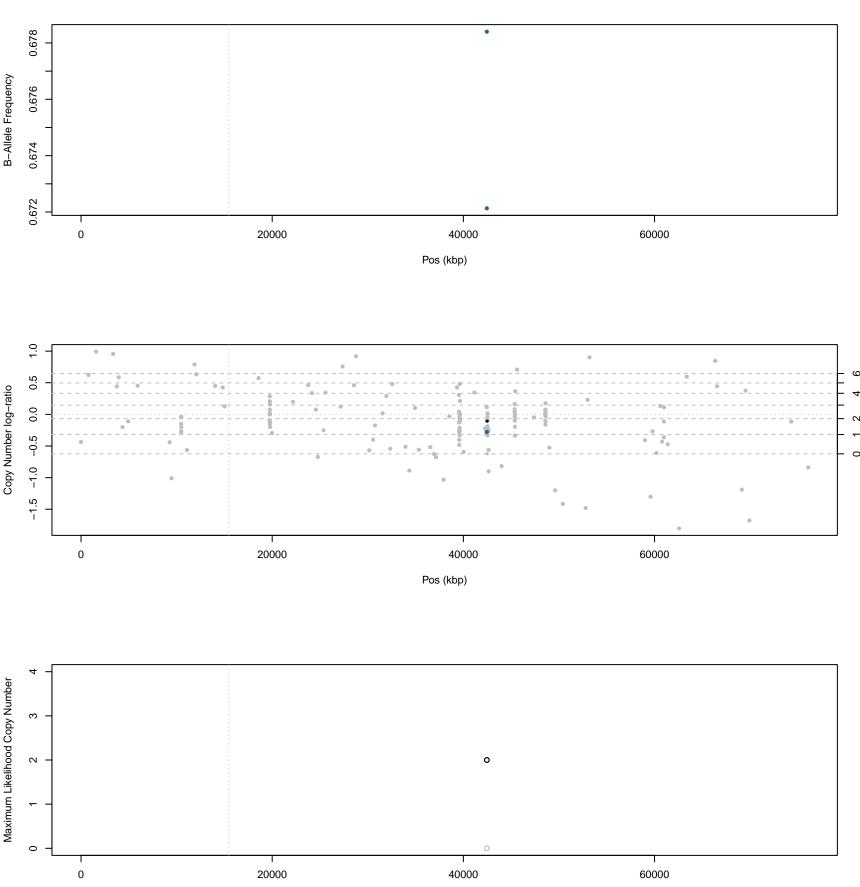


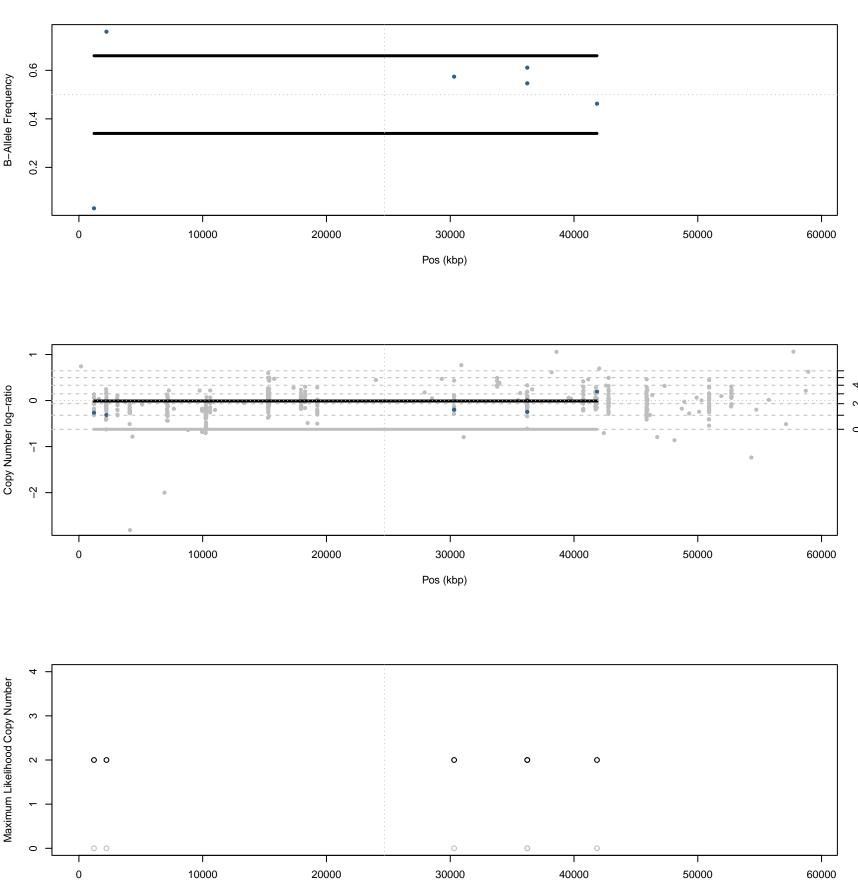
Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 16

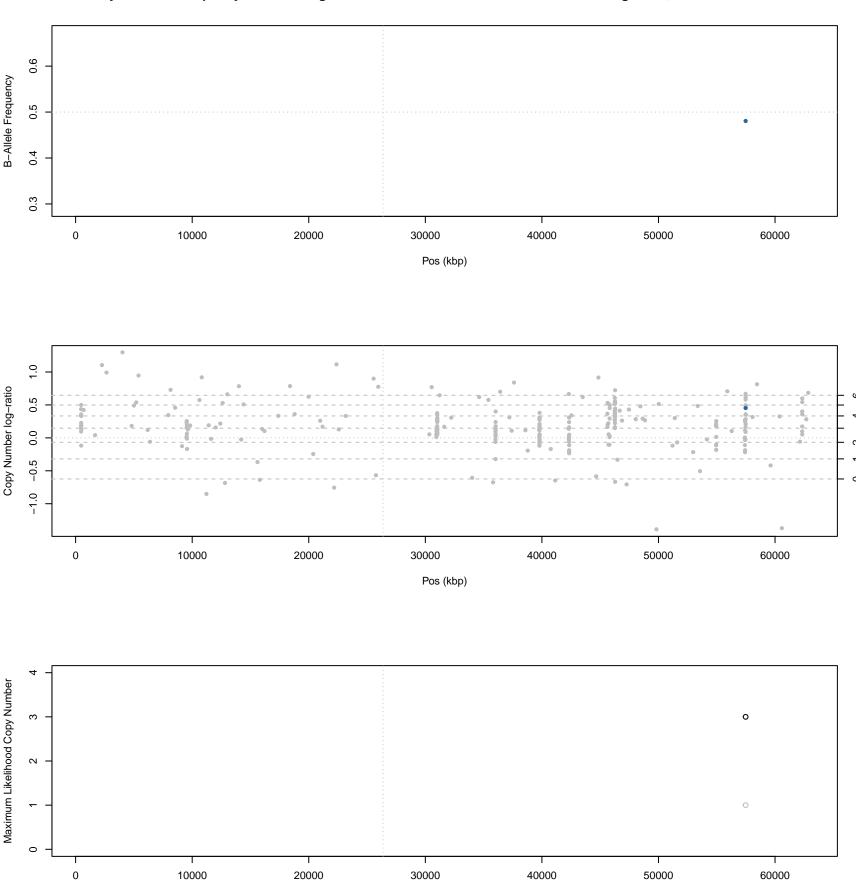


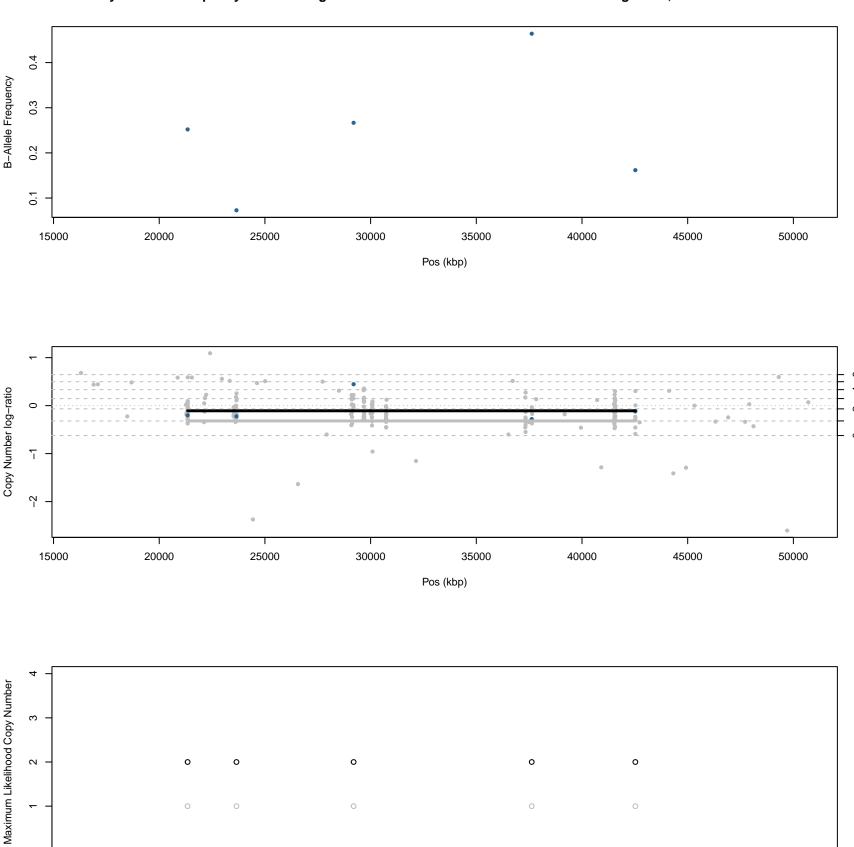


Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413 Chromosome: 18









Pos (kbp)