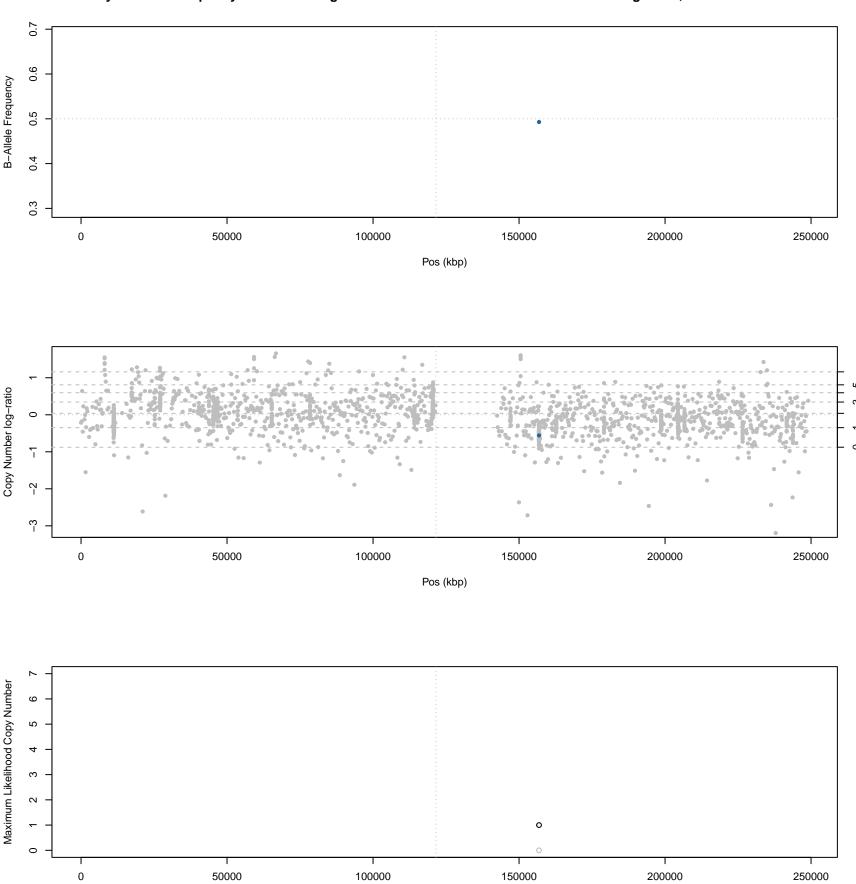
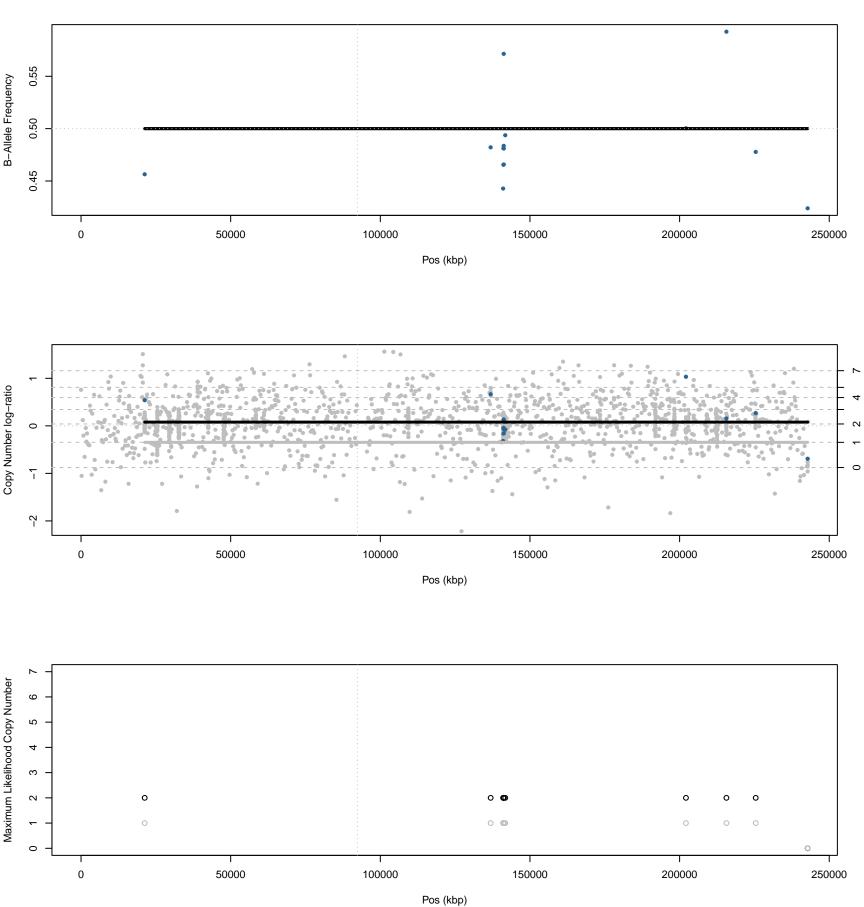
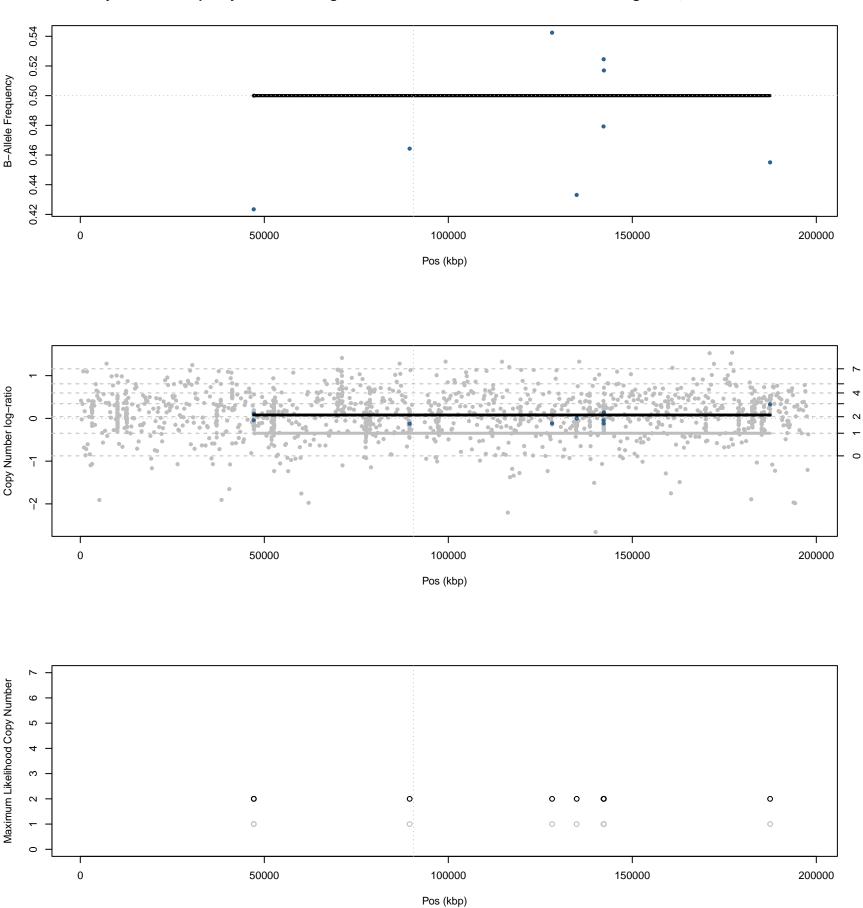
Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 1



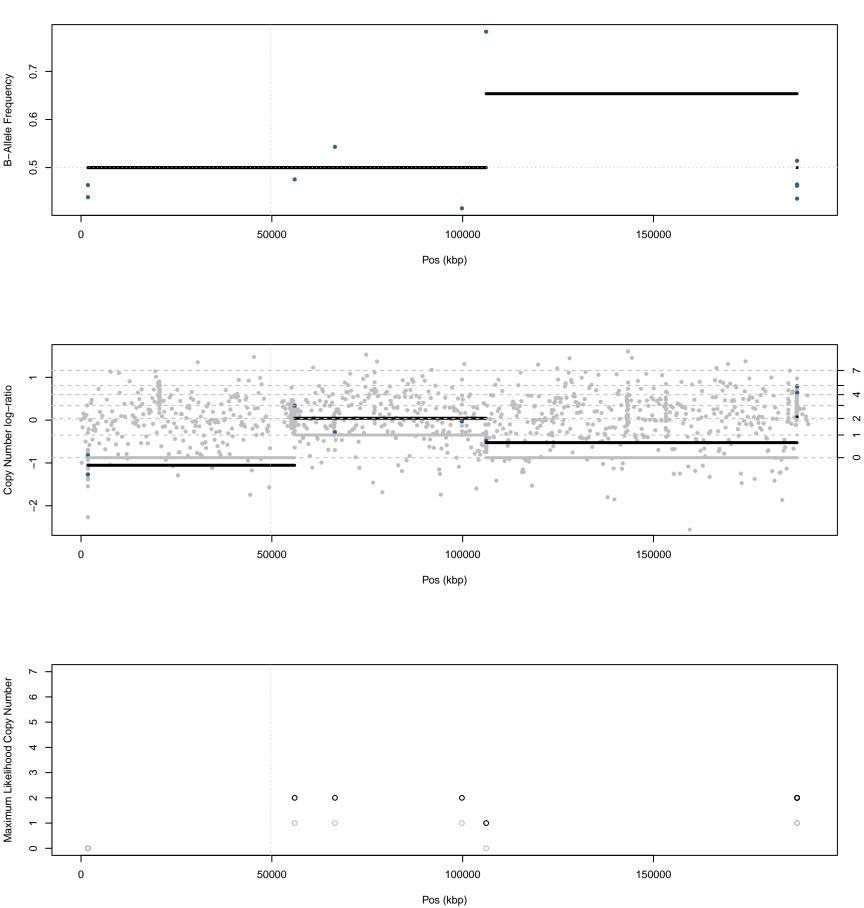
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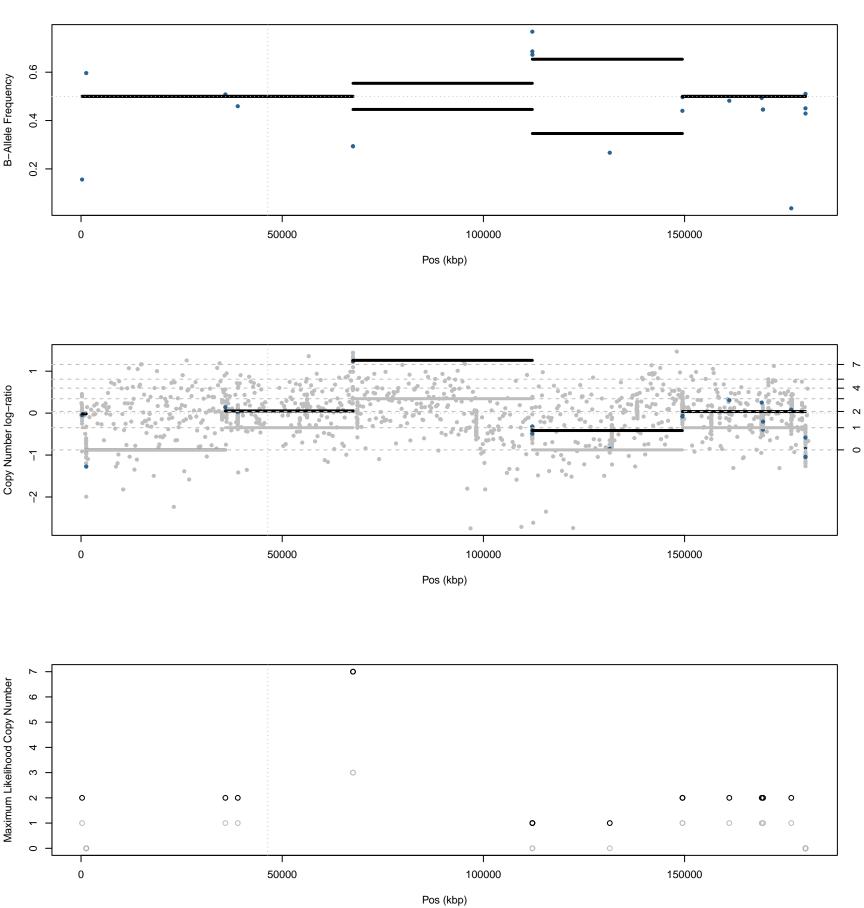
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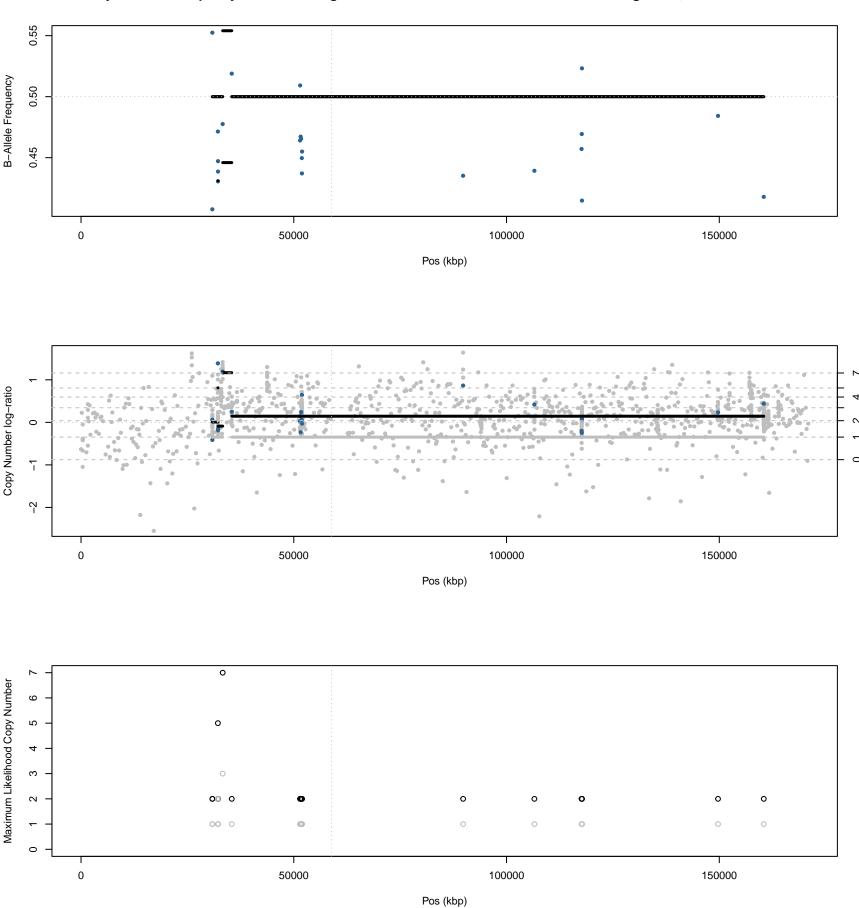
Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 4



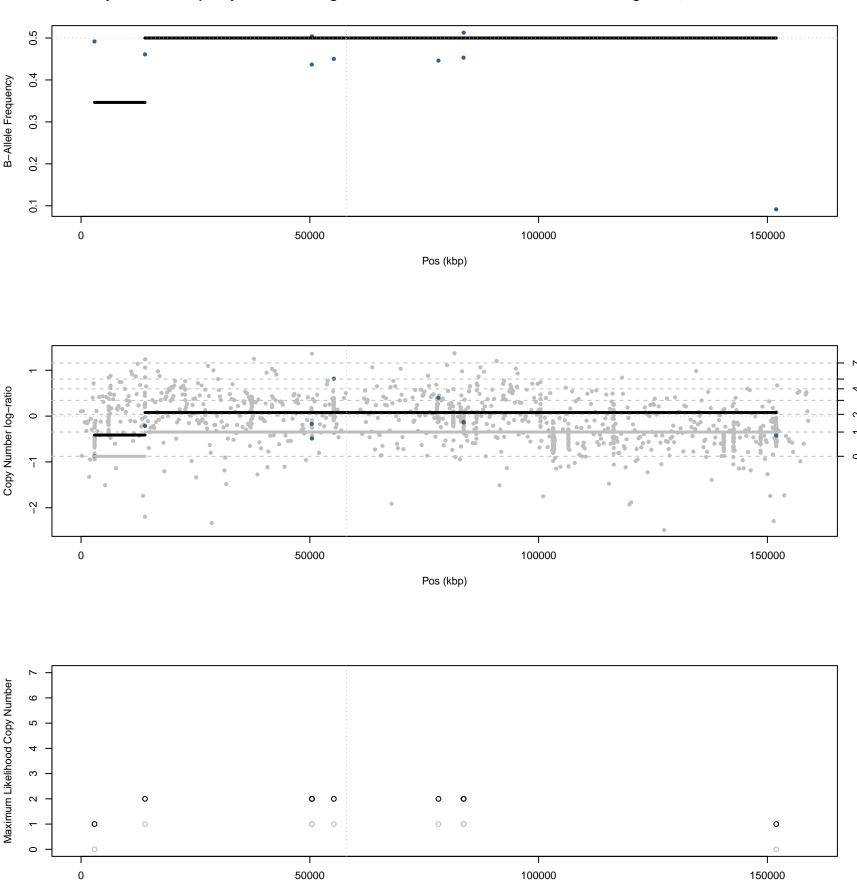
Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 5



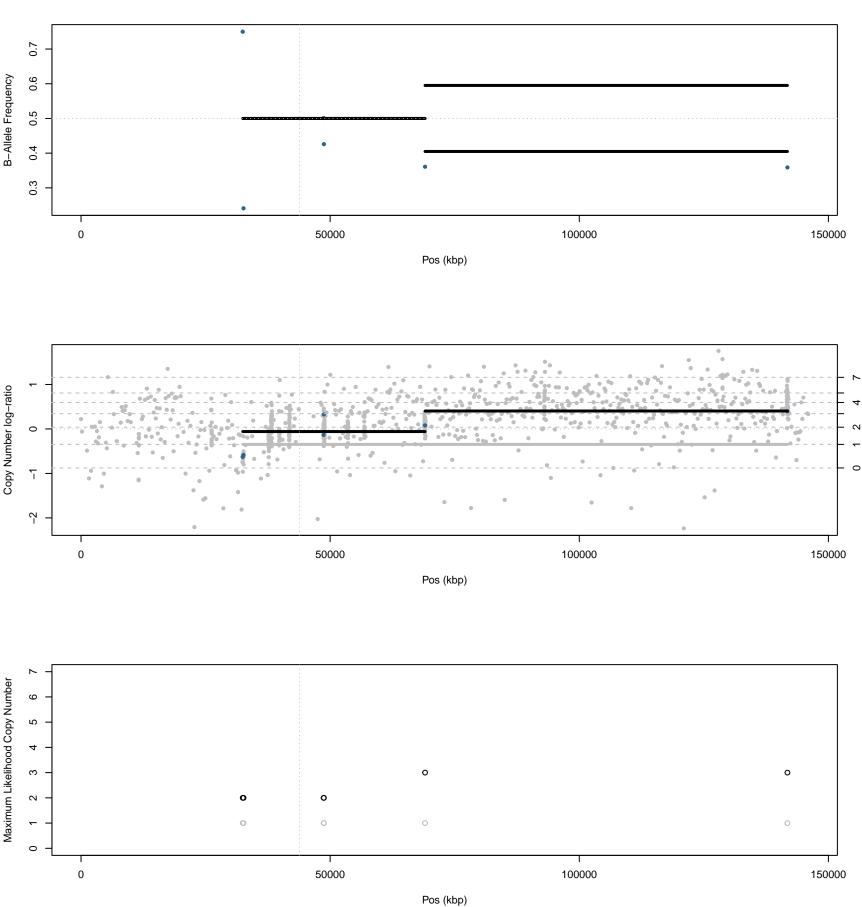
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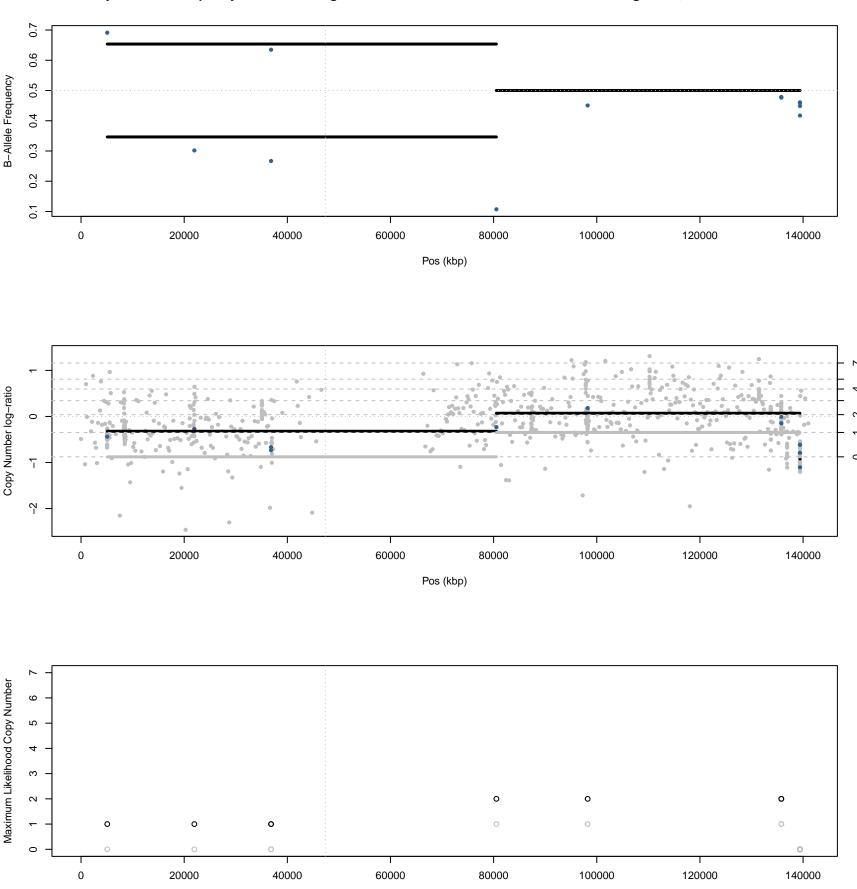
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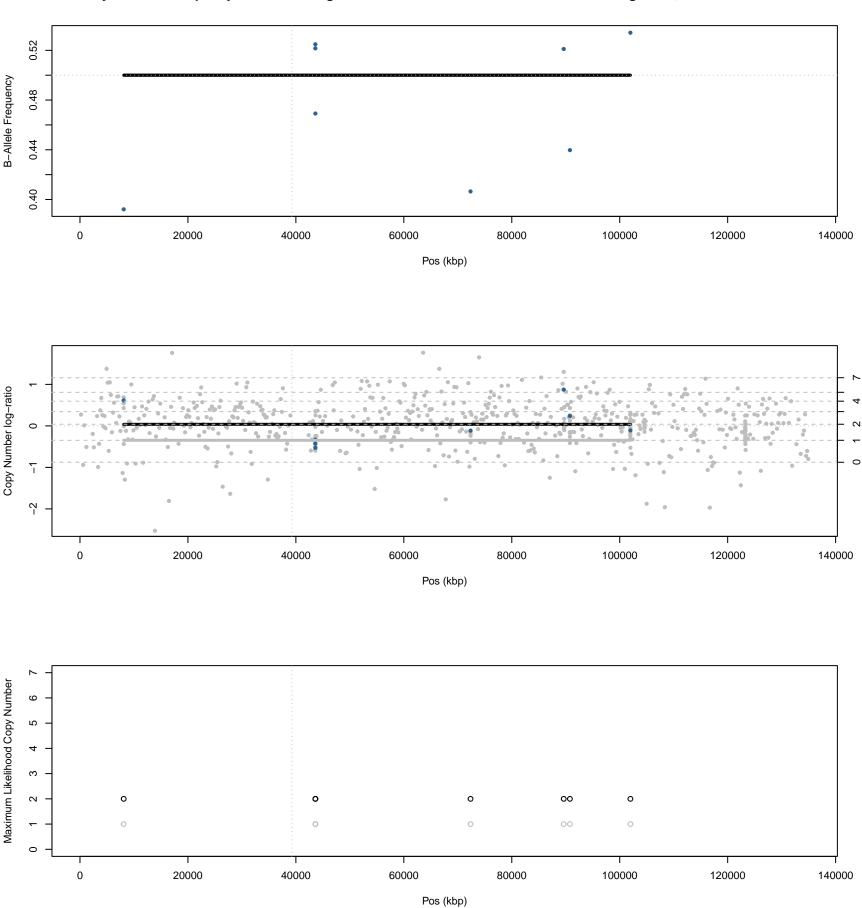
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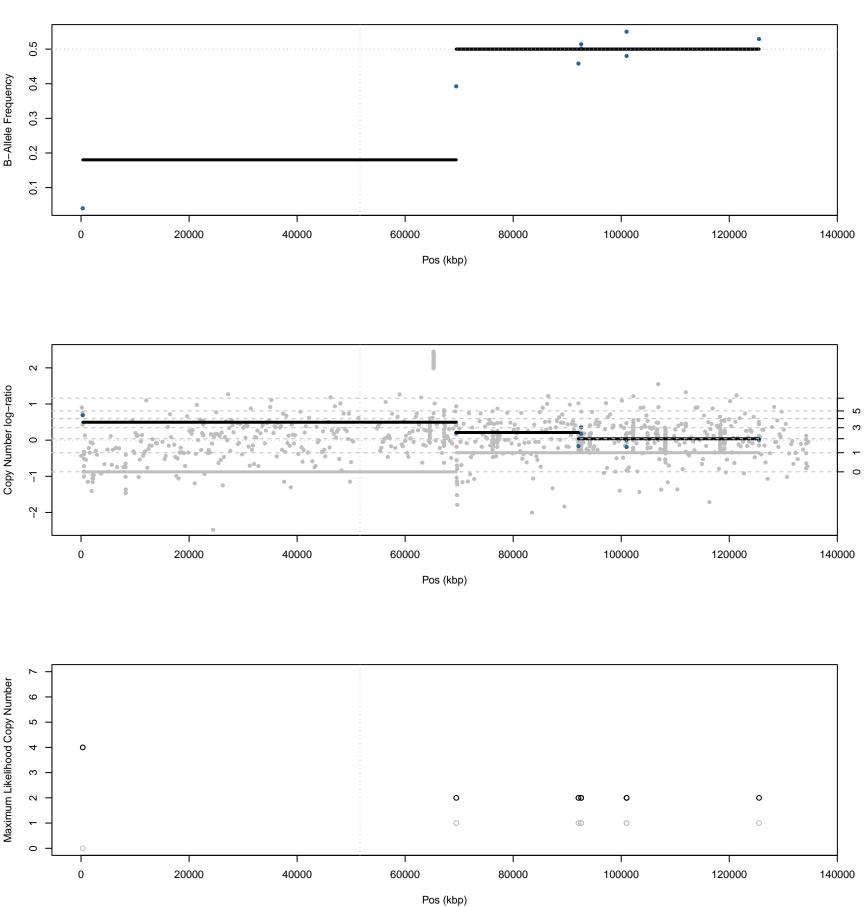
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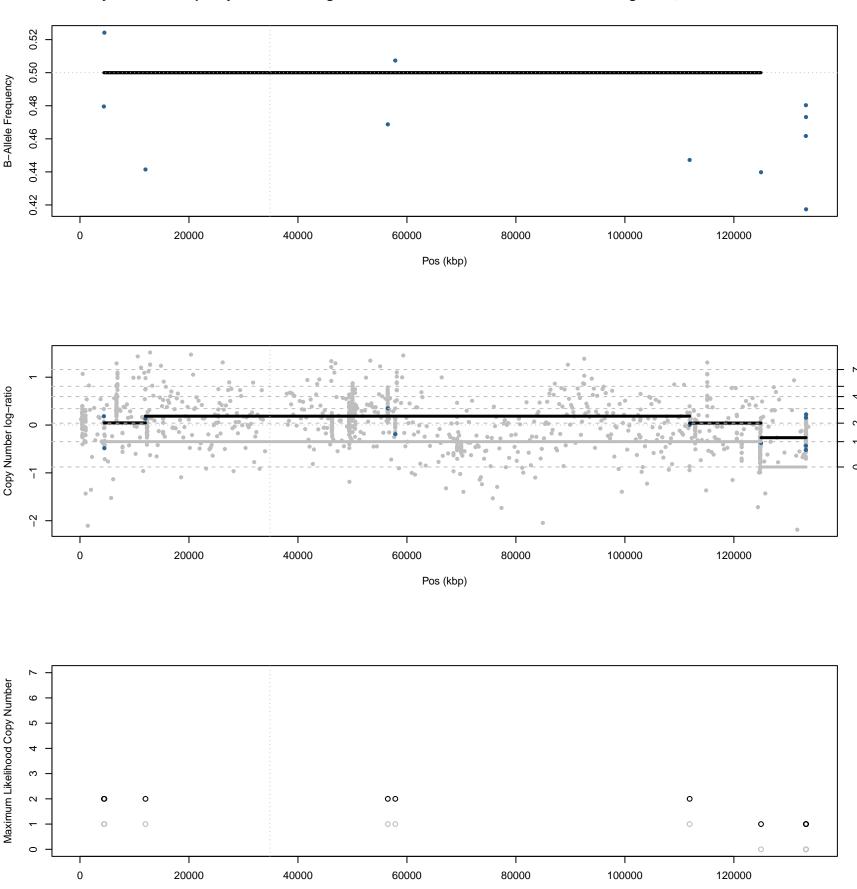
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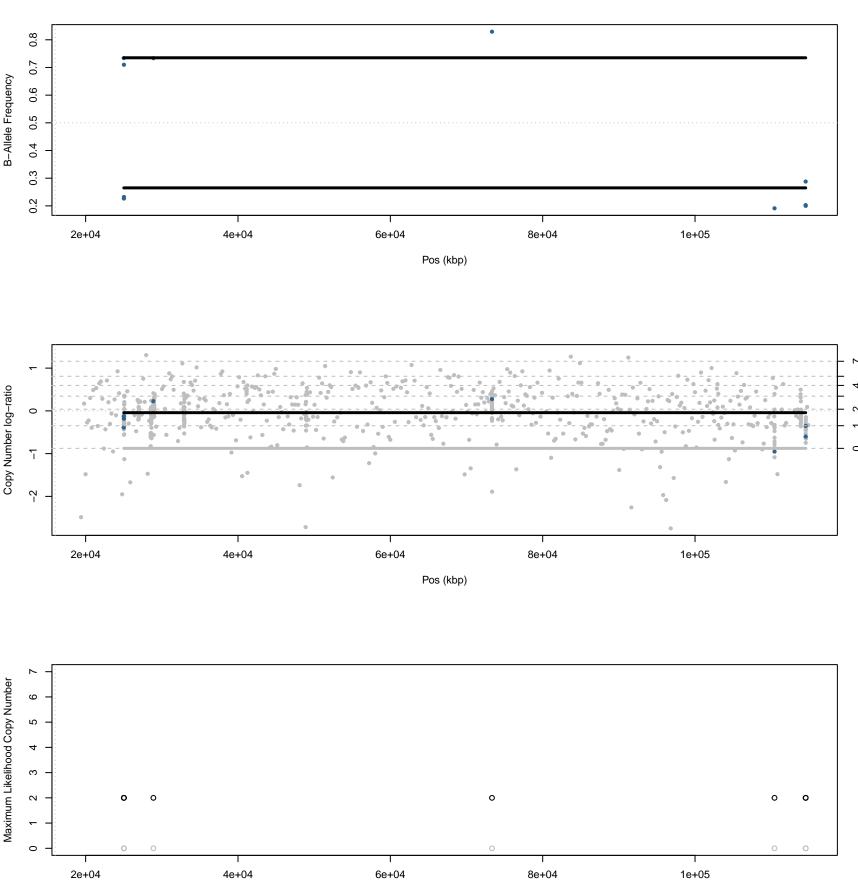
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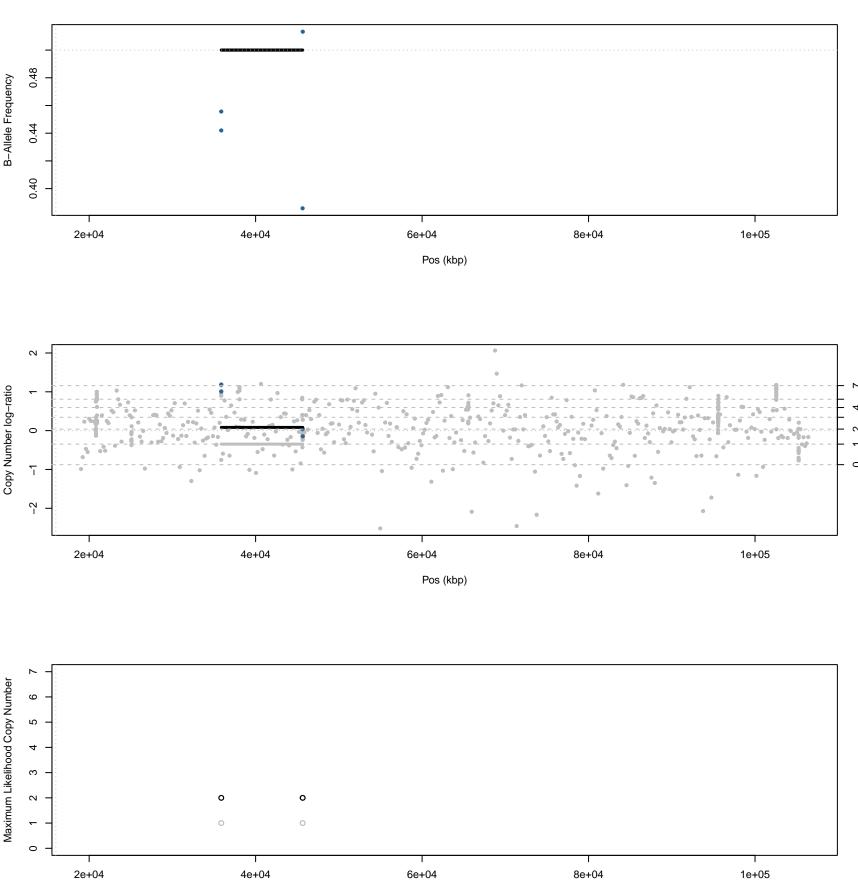
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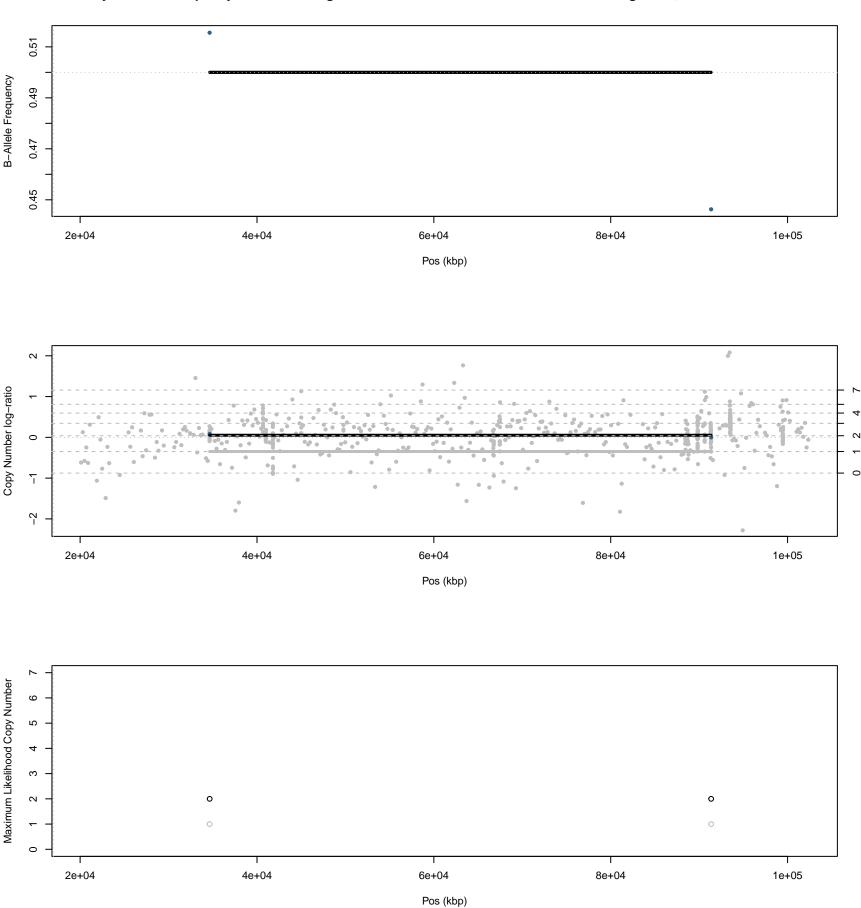
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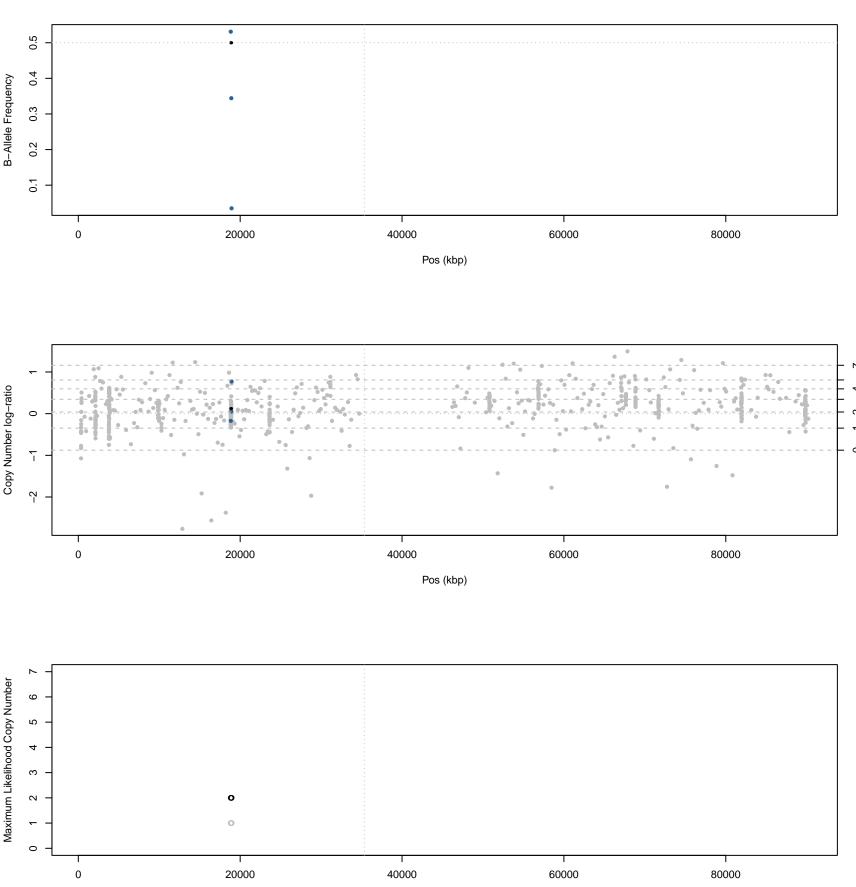
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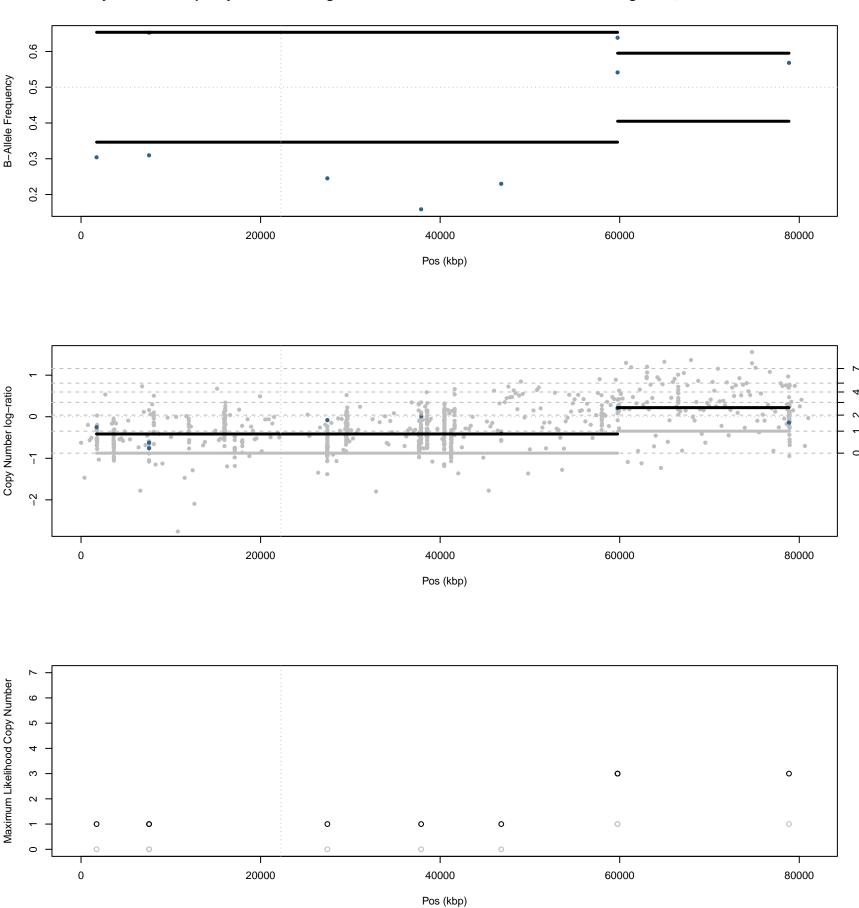
Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 15



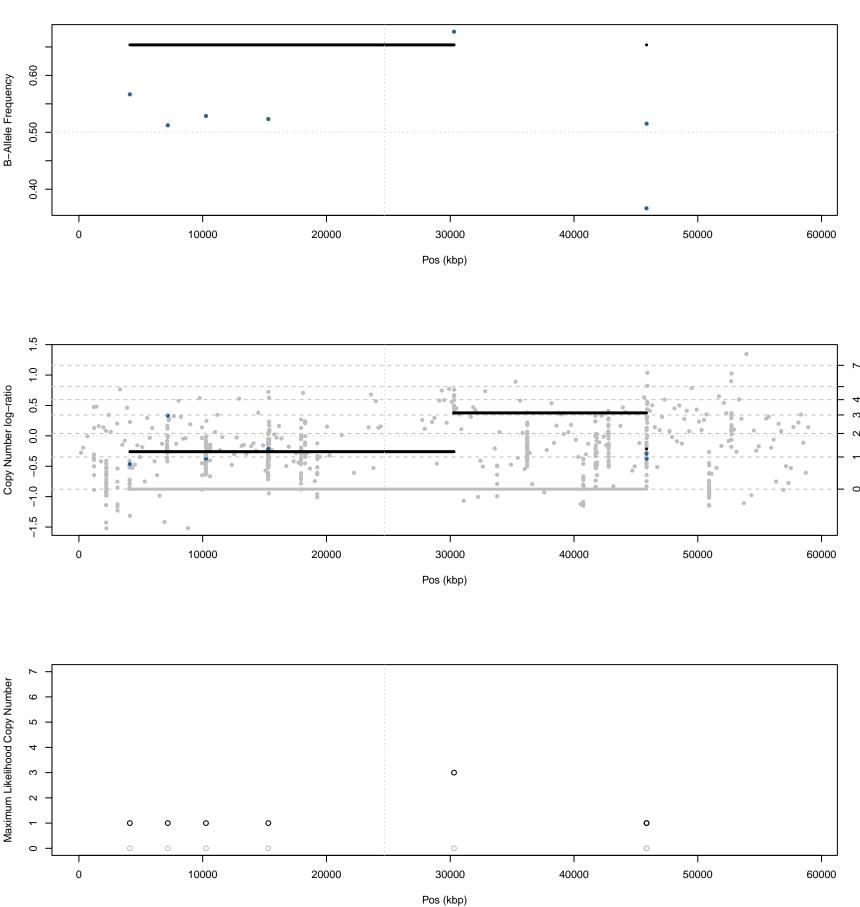
Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 16



Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 17



Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 19



Purity: 0.47 Tumor ploidy: 1.886 SNV log-likelihood: -124.56 GoF: 86.2% Mean coverage: 505;410 Chromosome: 20

