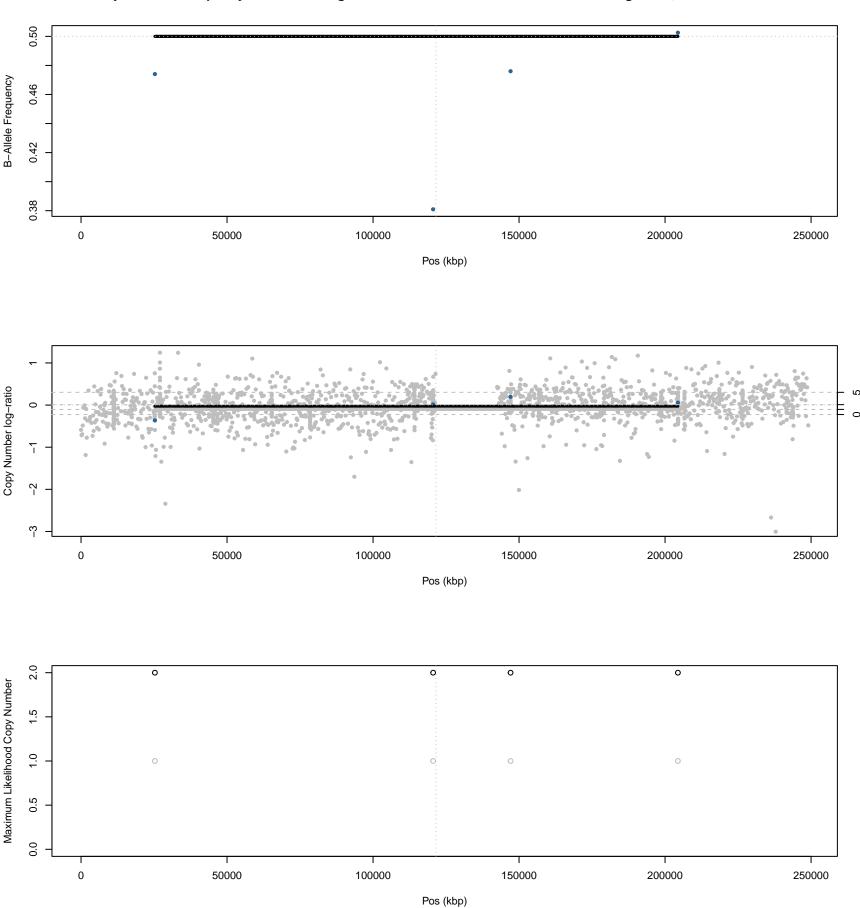
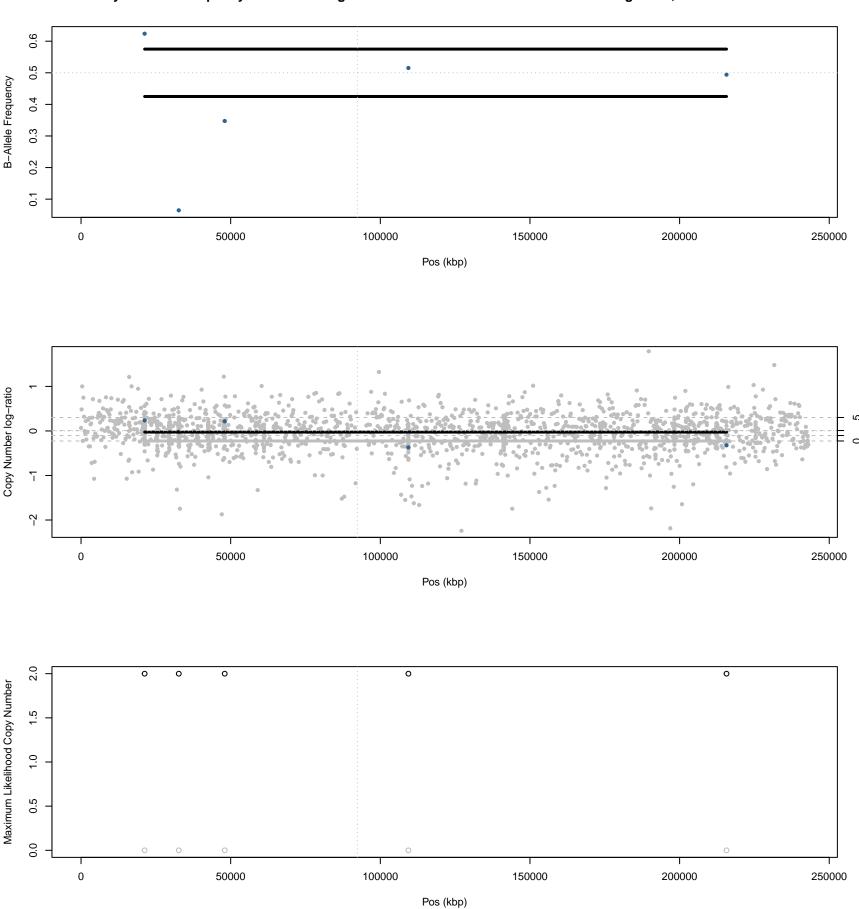
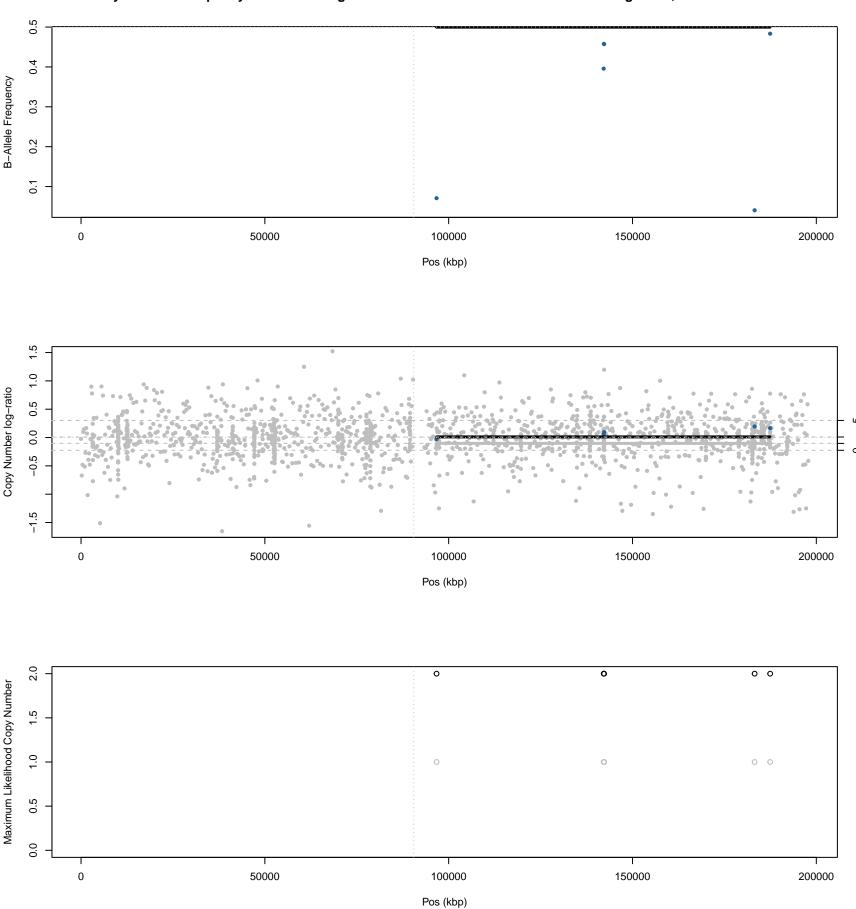
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 1



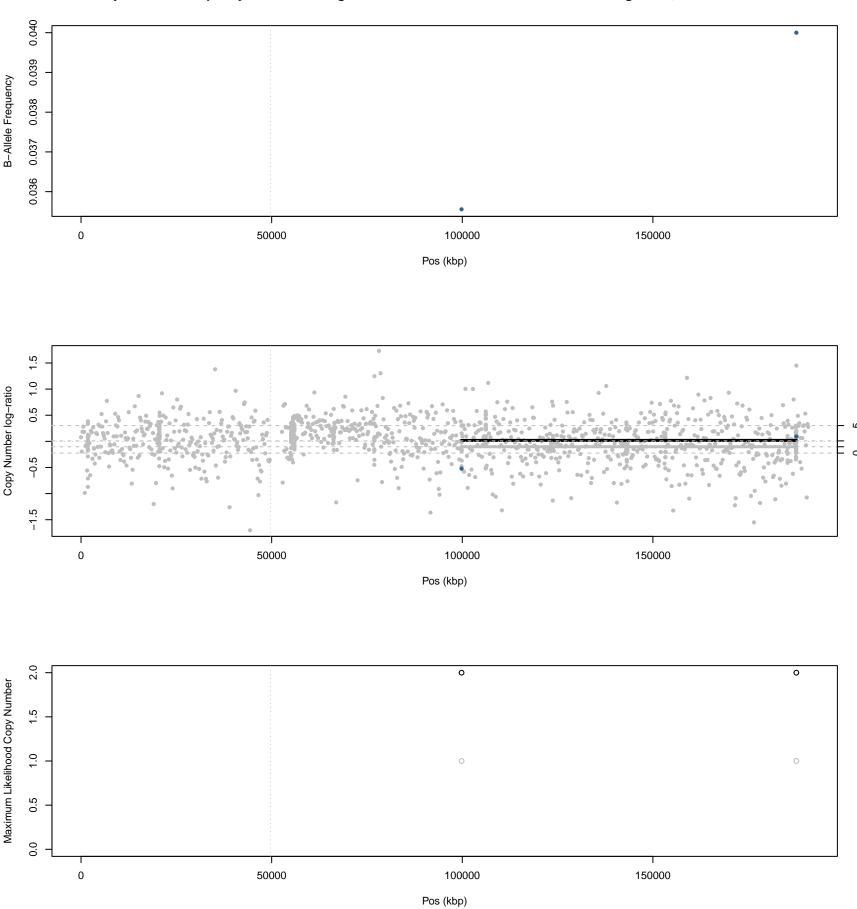
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 2



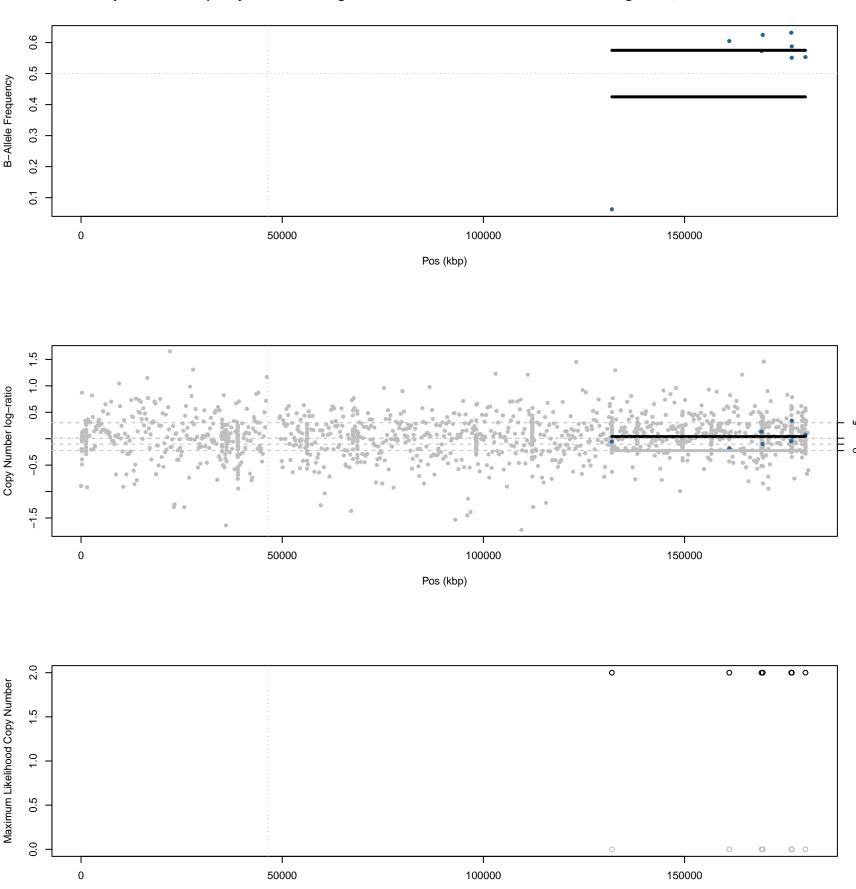
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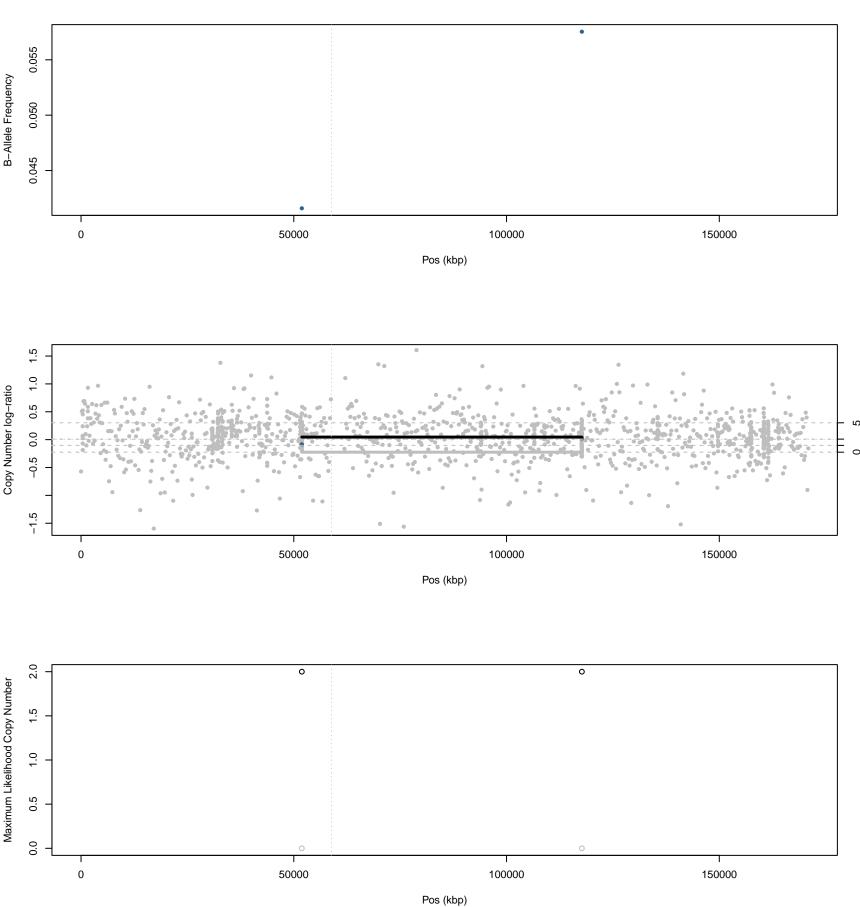
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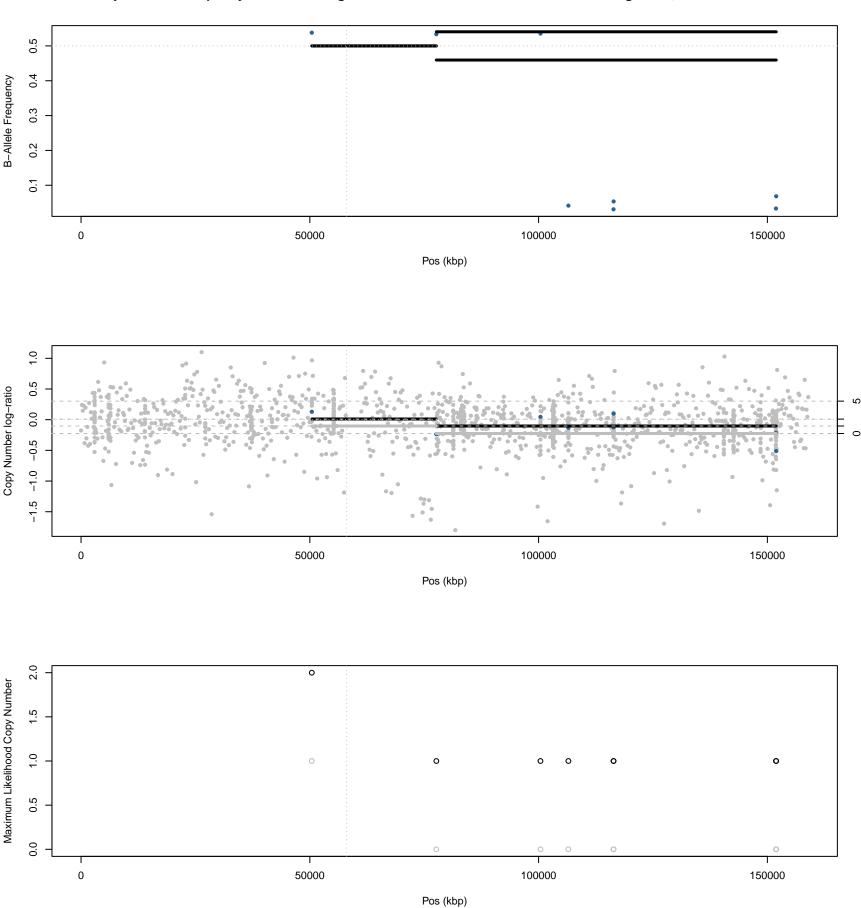
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 5



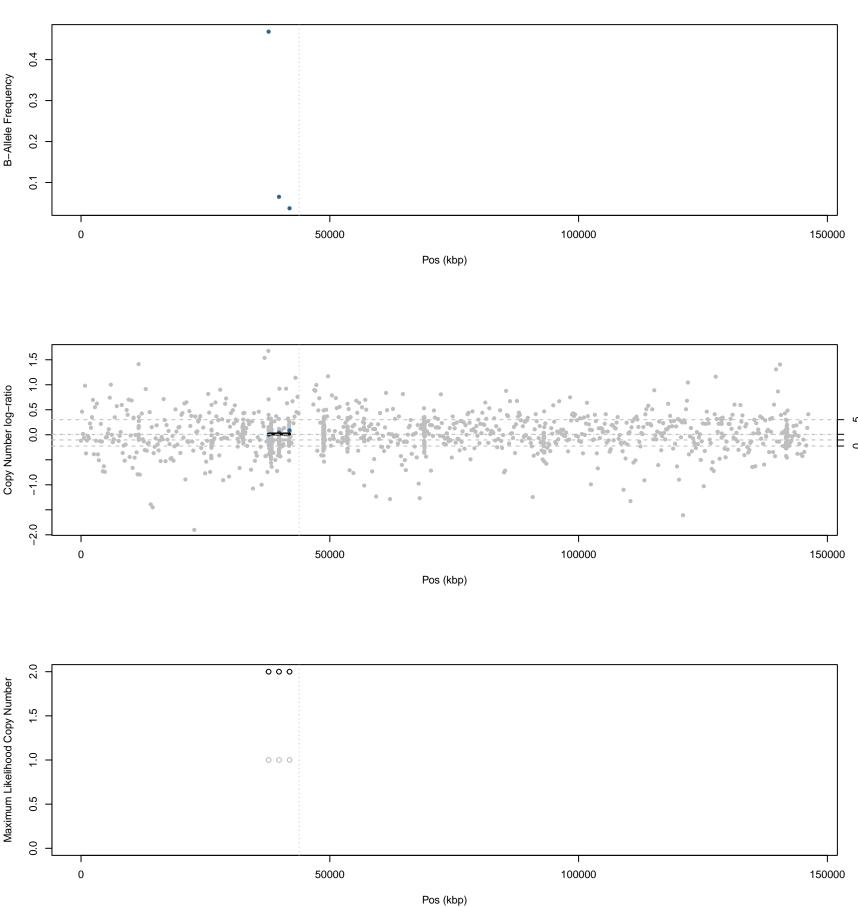
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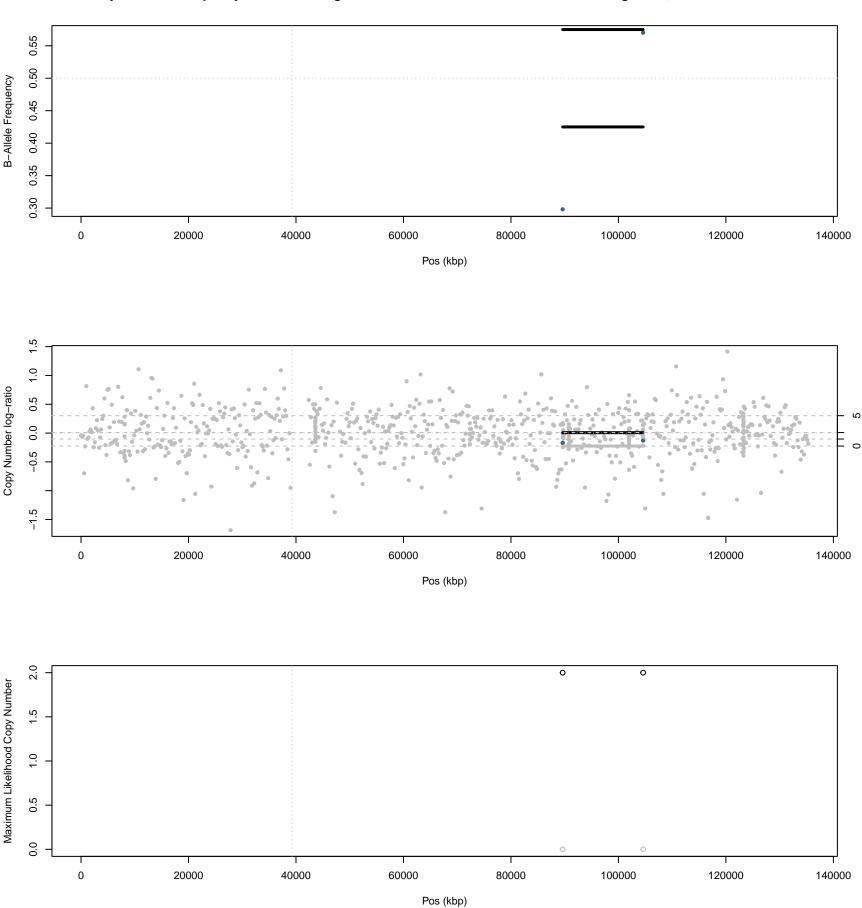
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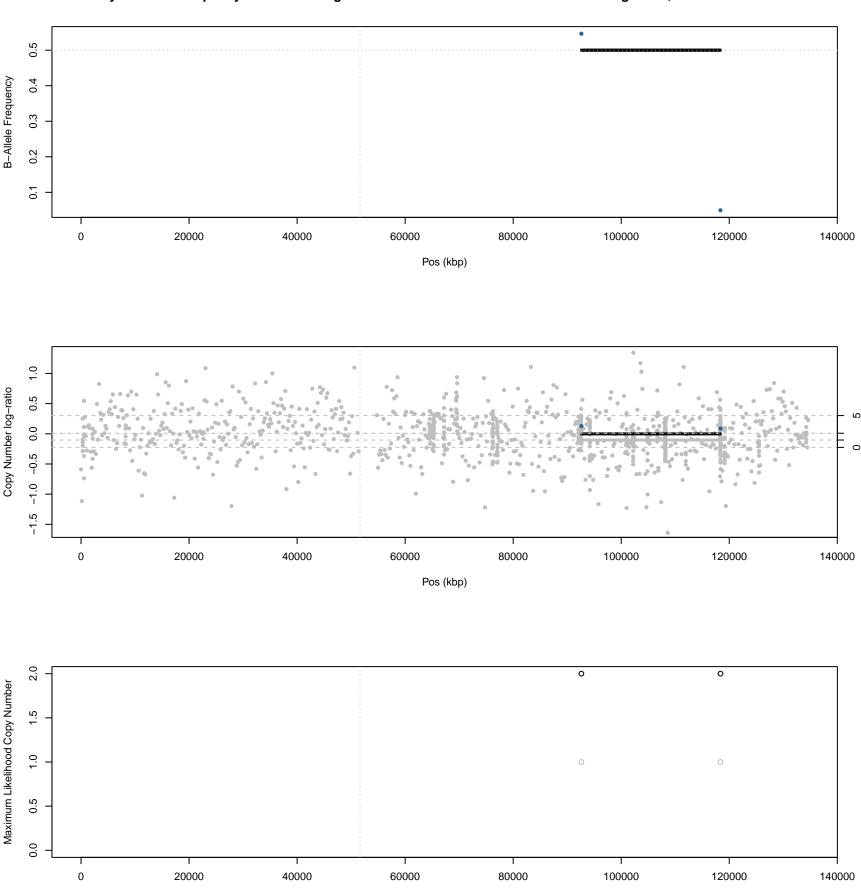
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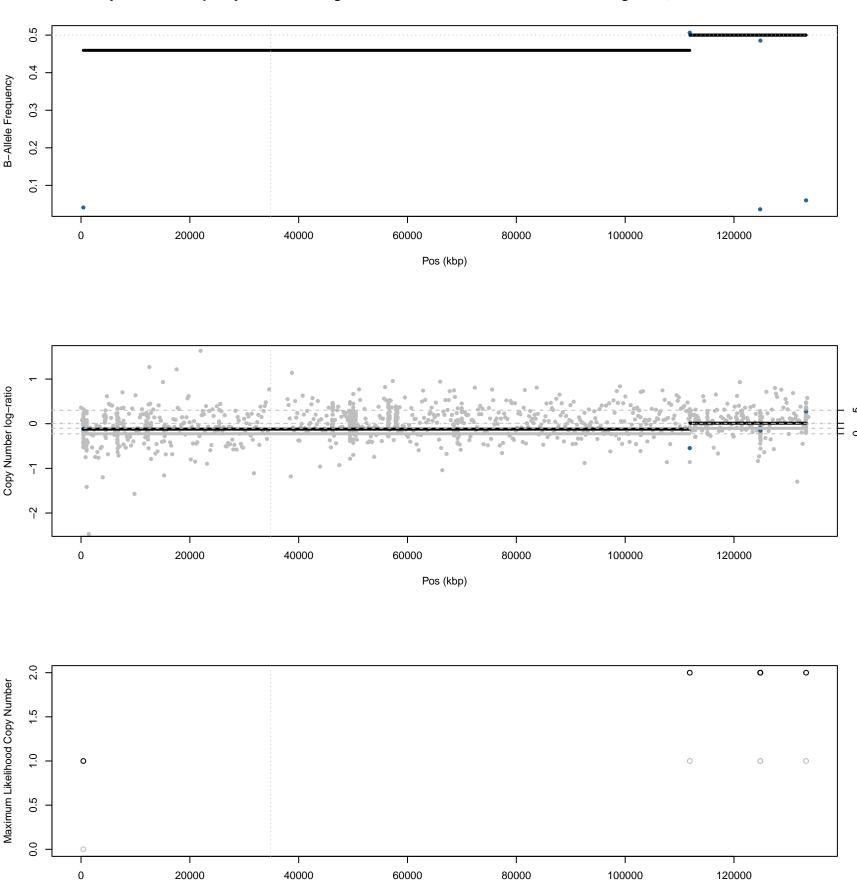
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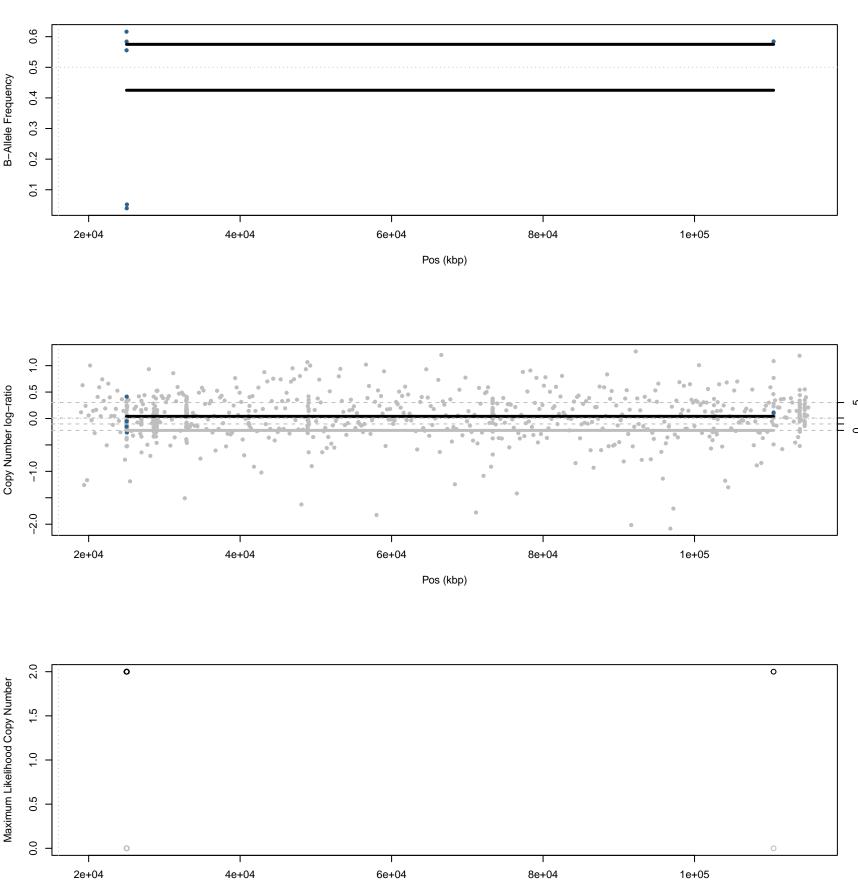
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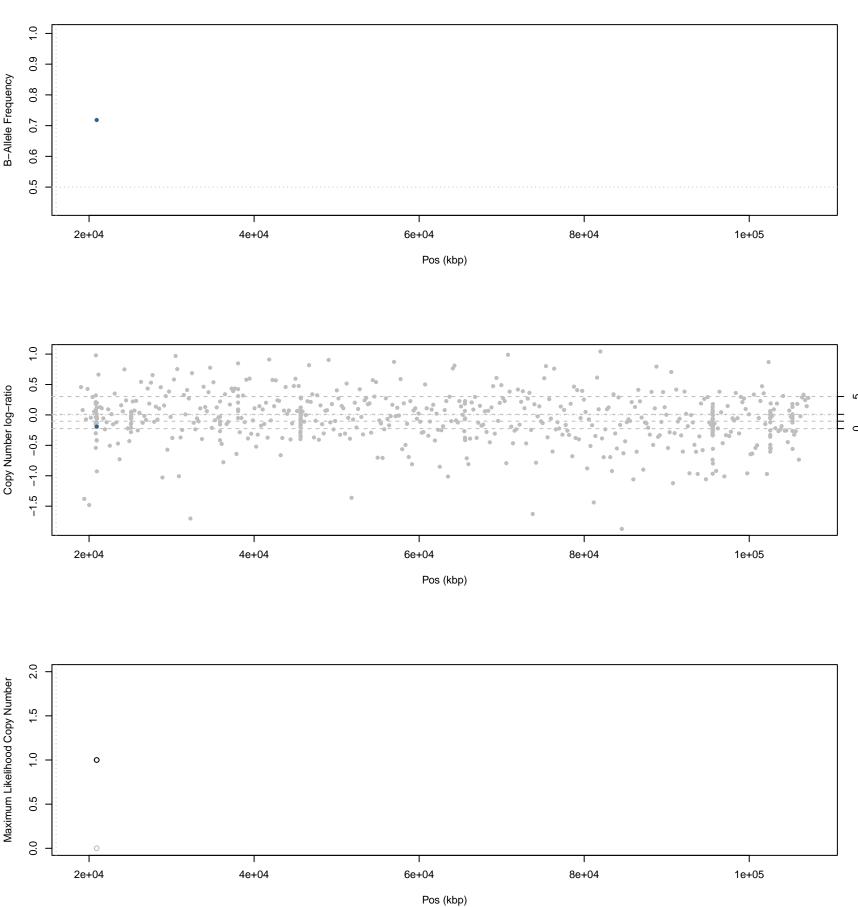
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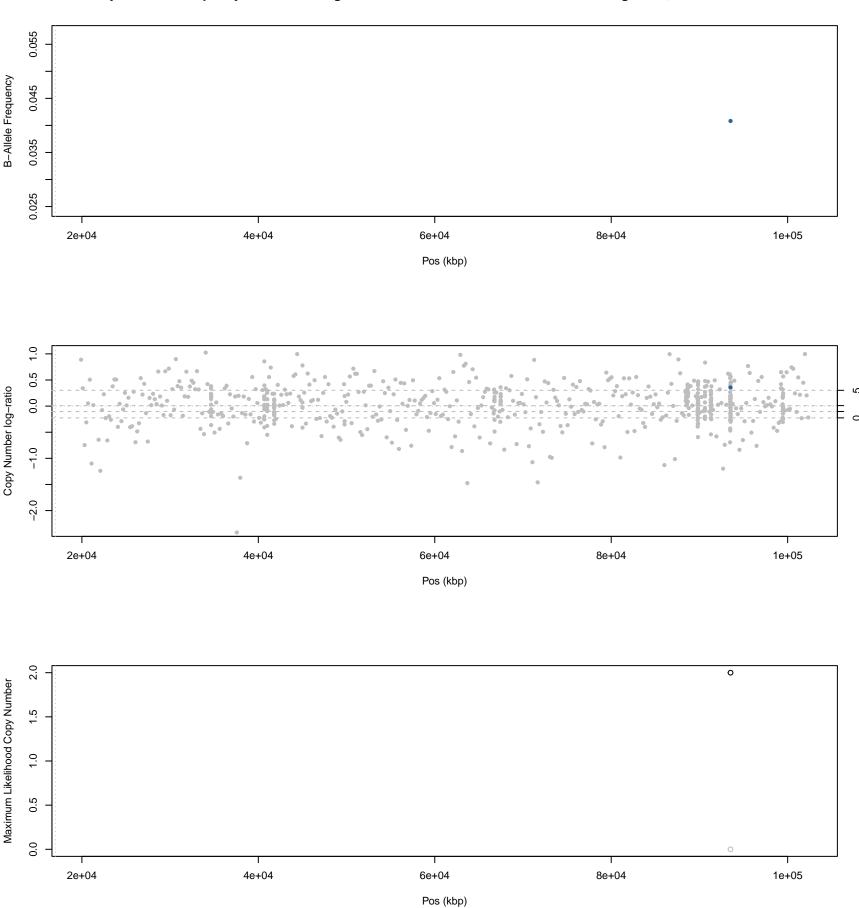
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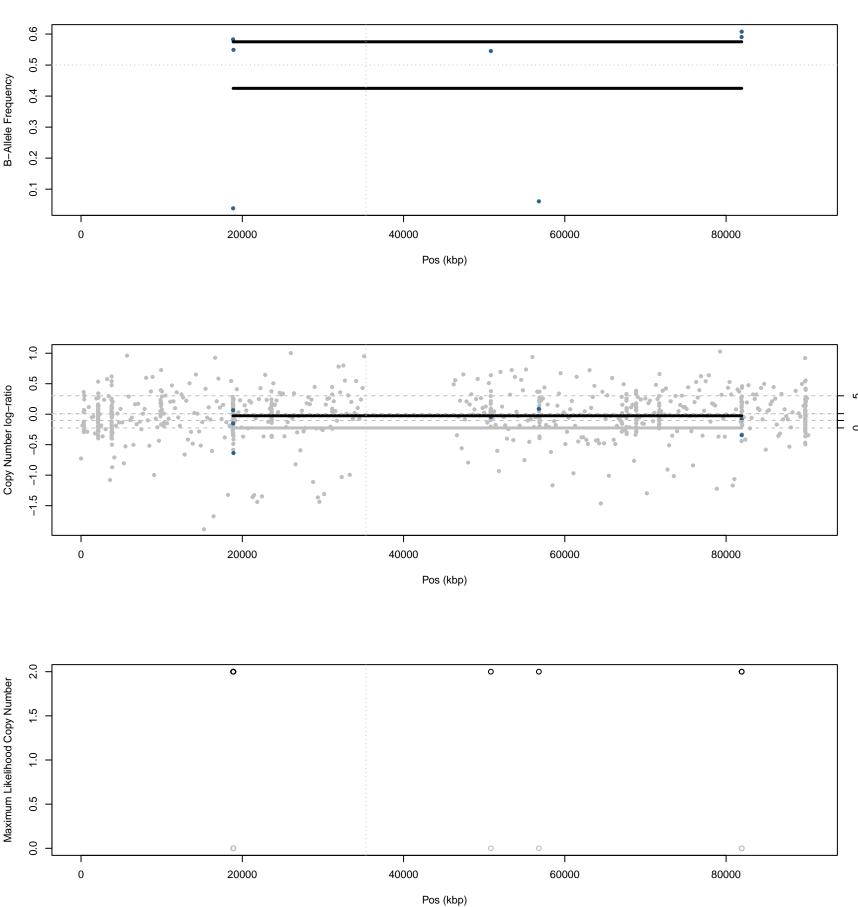
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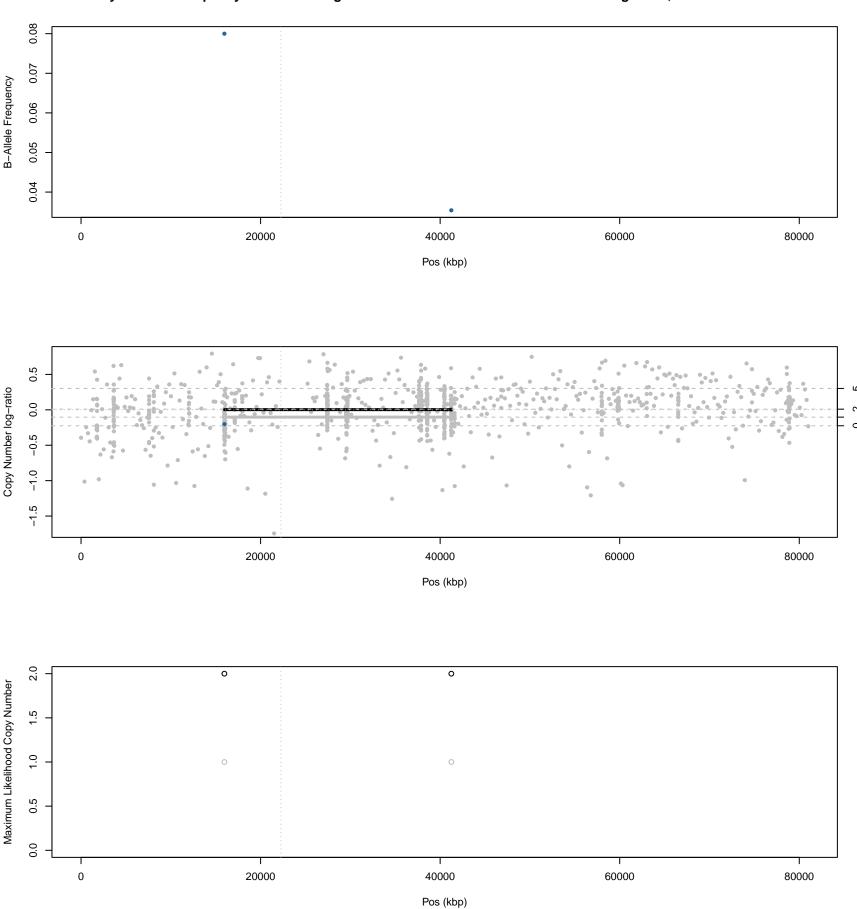
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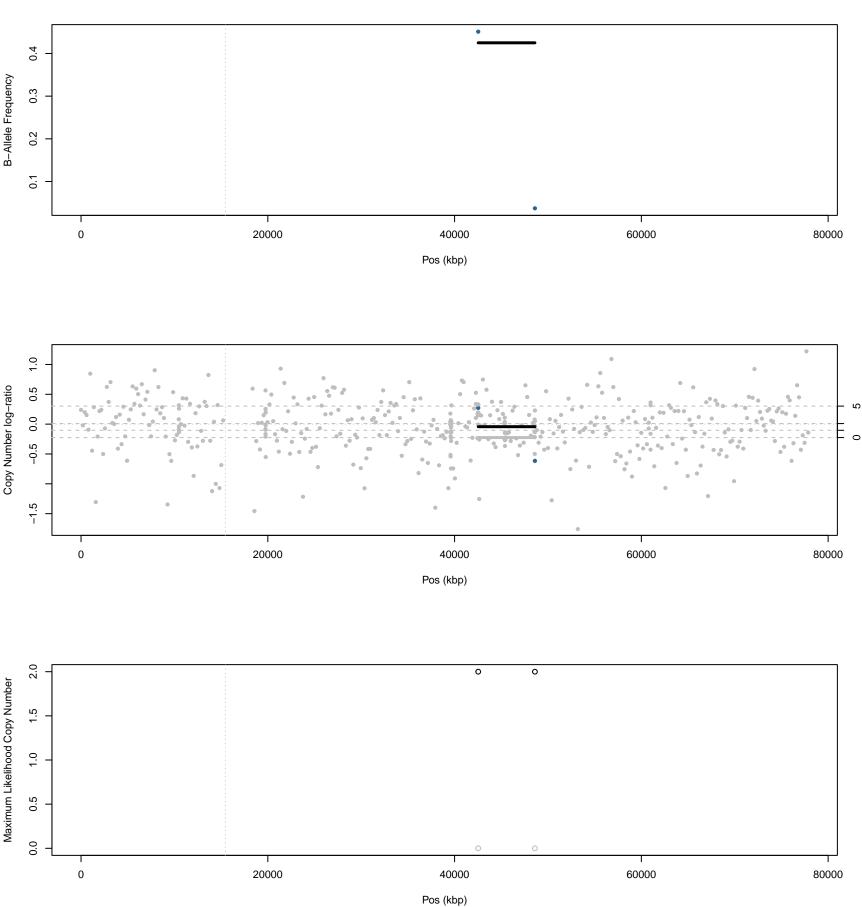
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 16



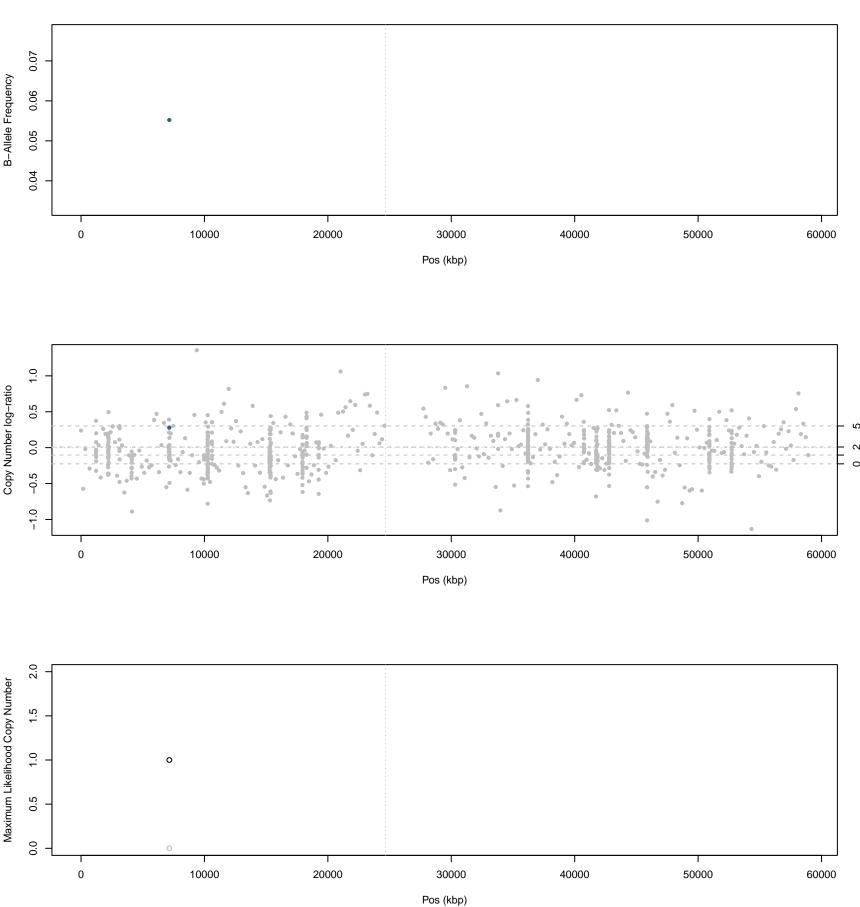
Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 17



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 18



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 19



Purity: 0.15 Tumor ploidy: 1.915 SNV log-likelihood: 12.83 GoF: 95.1% Mean coverage: 496;466 Chromosome: 22

