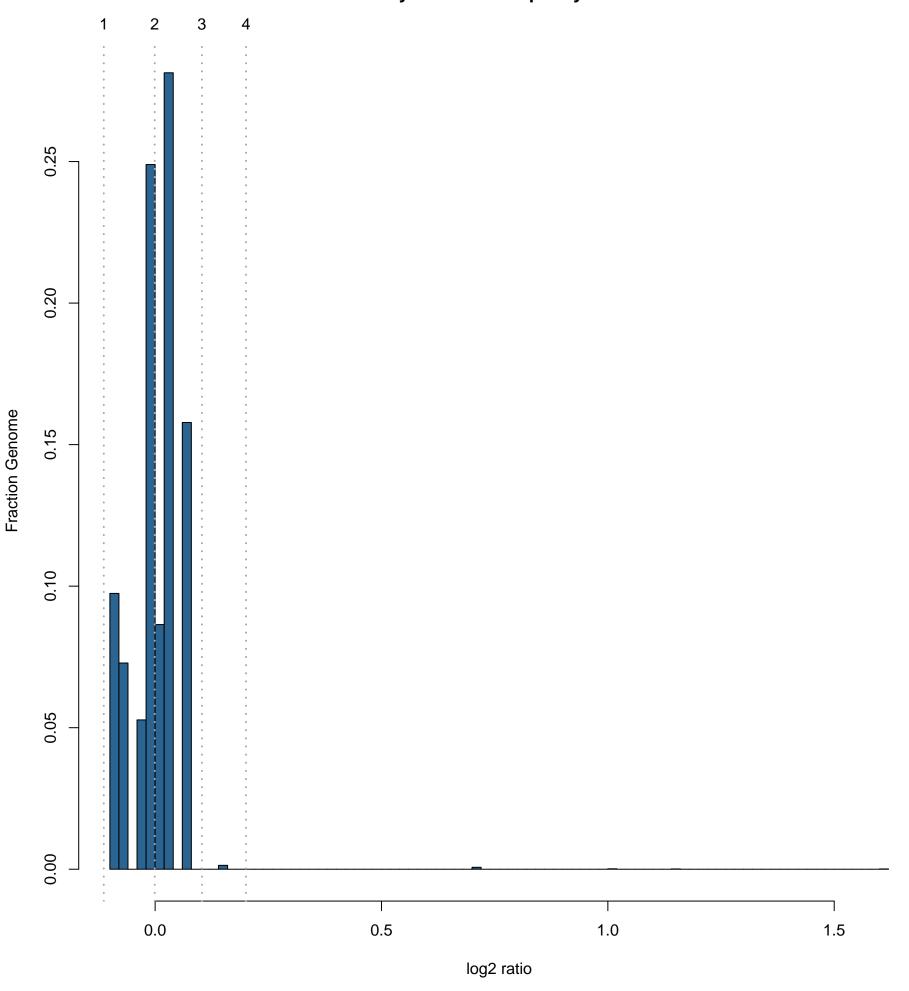
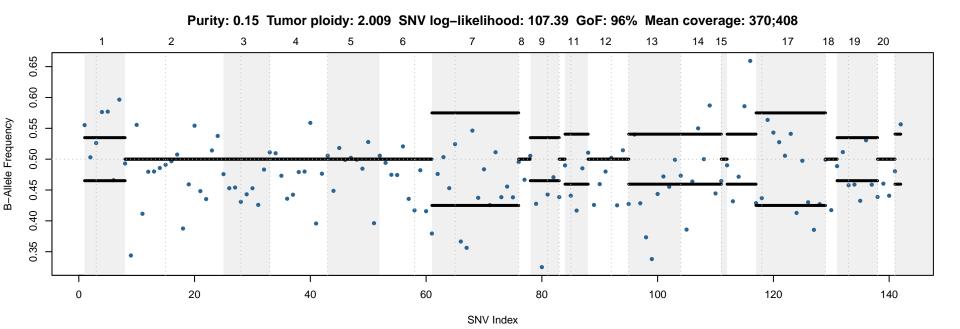
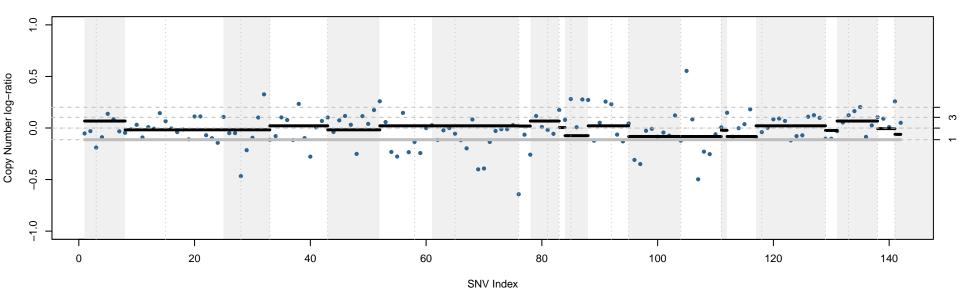
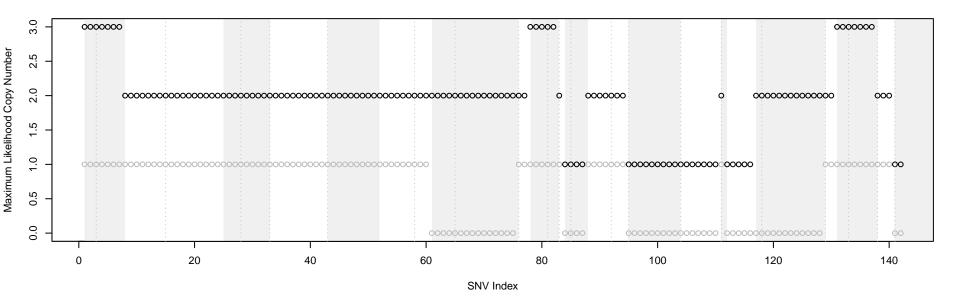
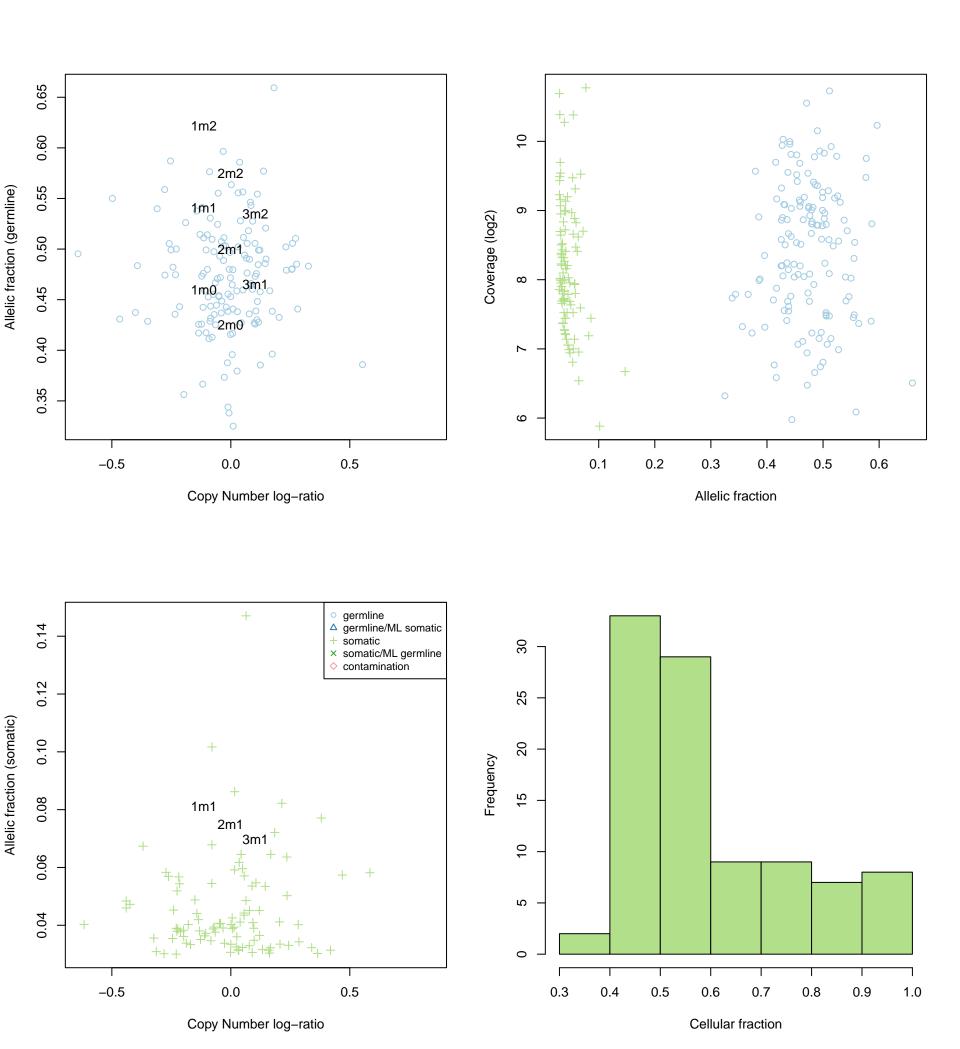
Purity: 0.15 Tumor ploidy: 2.009



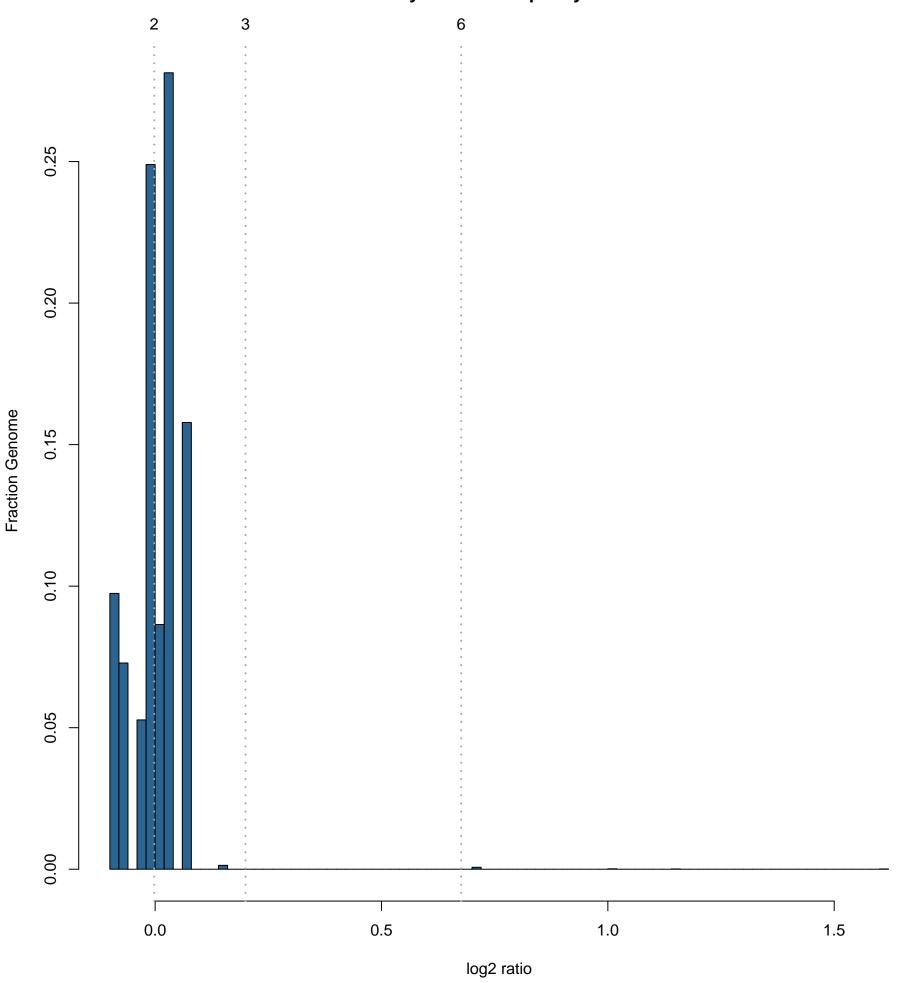


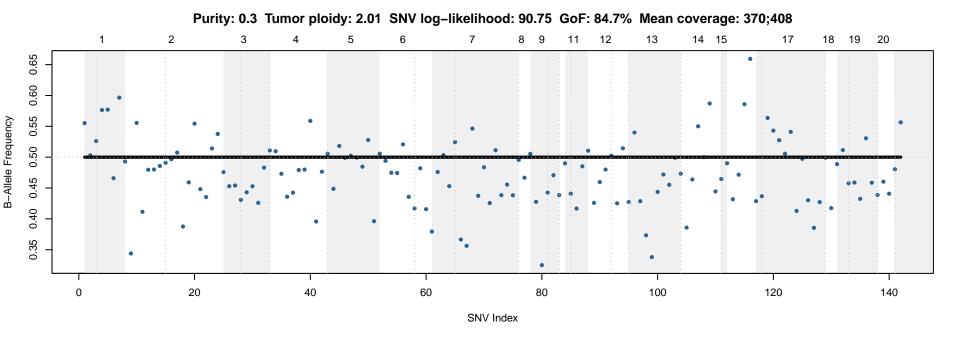


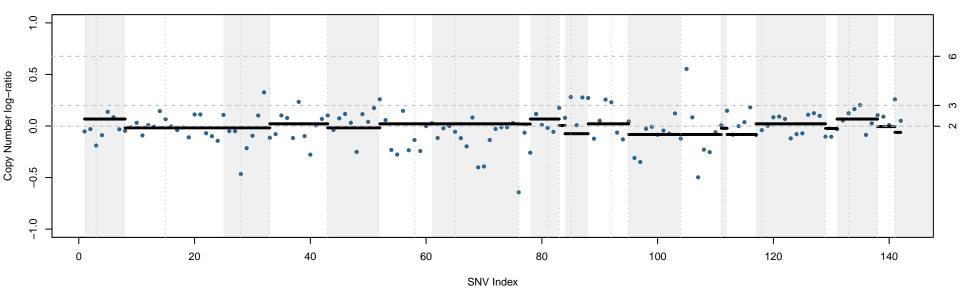


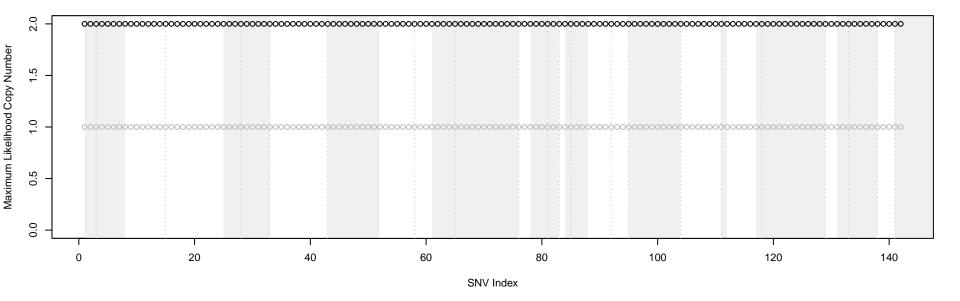


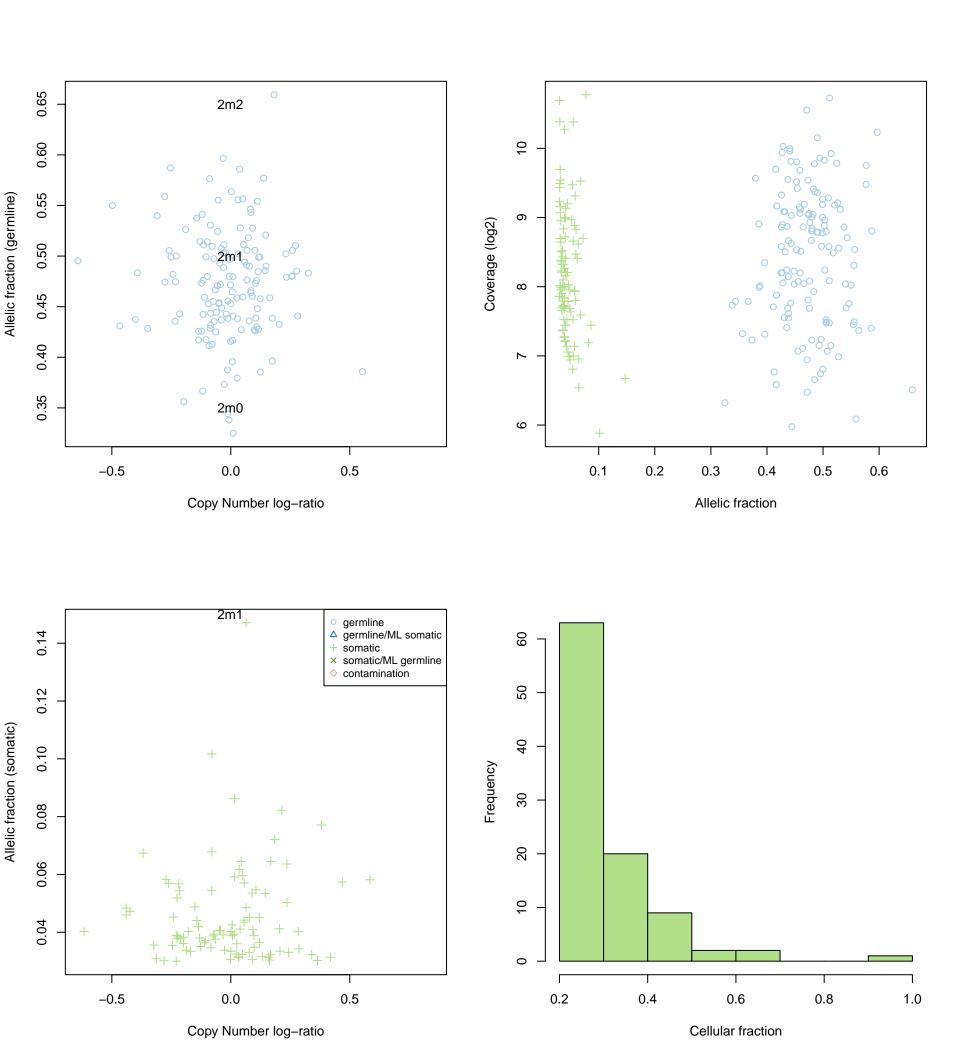
Purity: 0.3 Tumor ploidy: 2.01



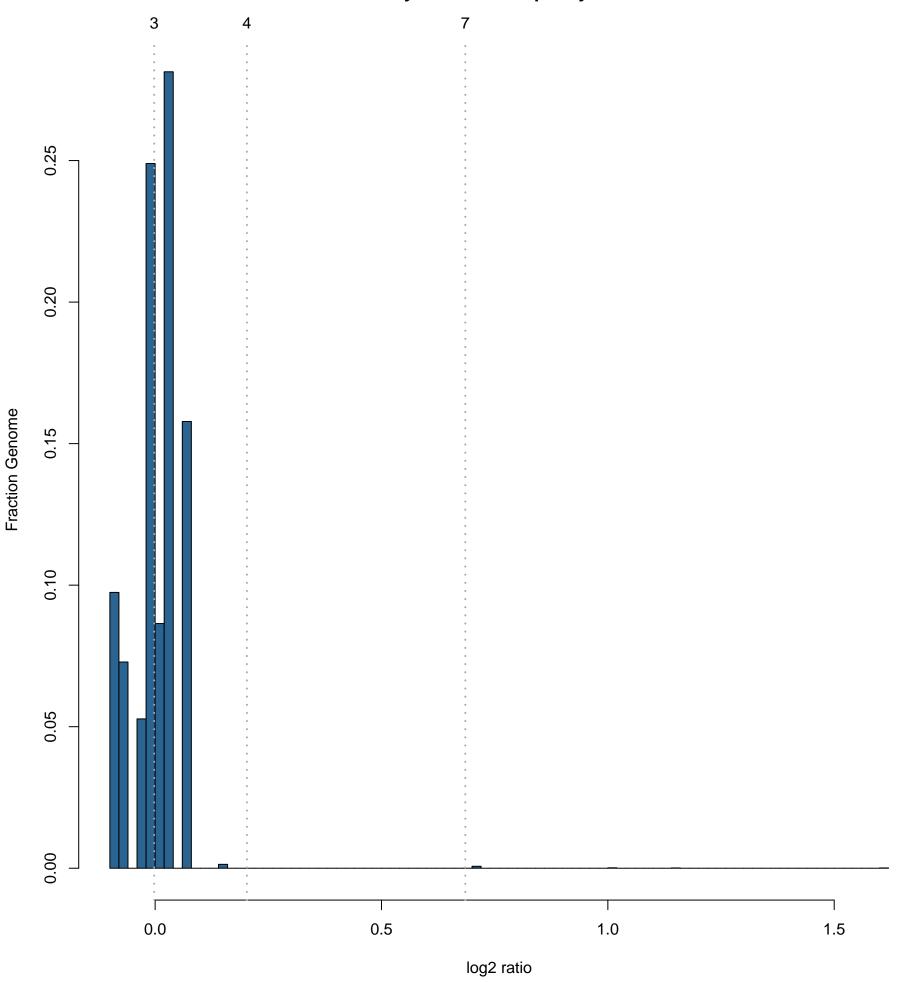


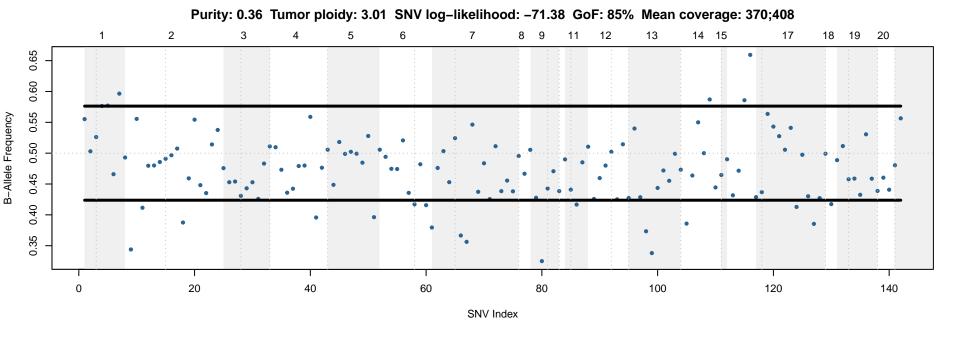


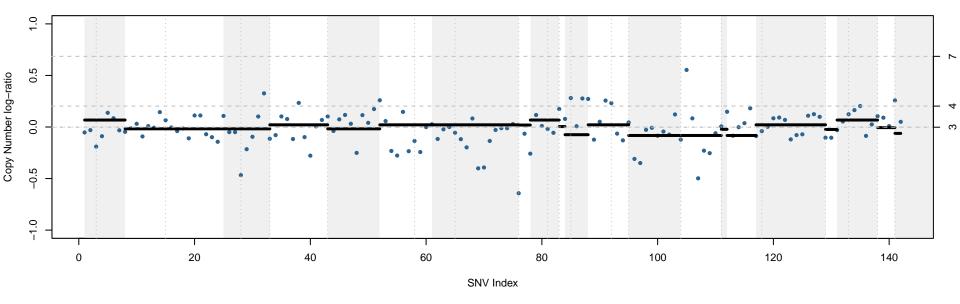


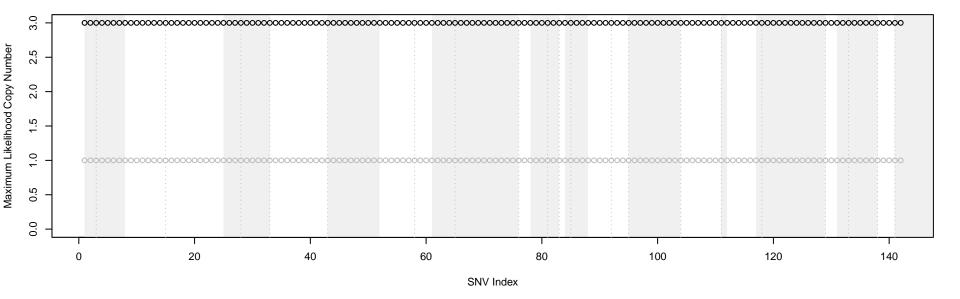


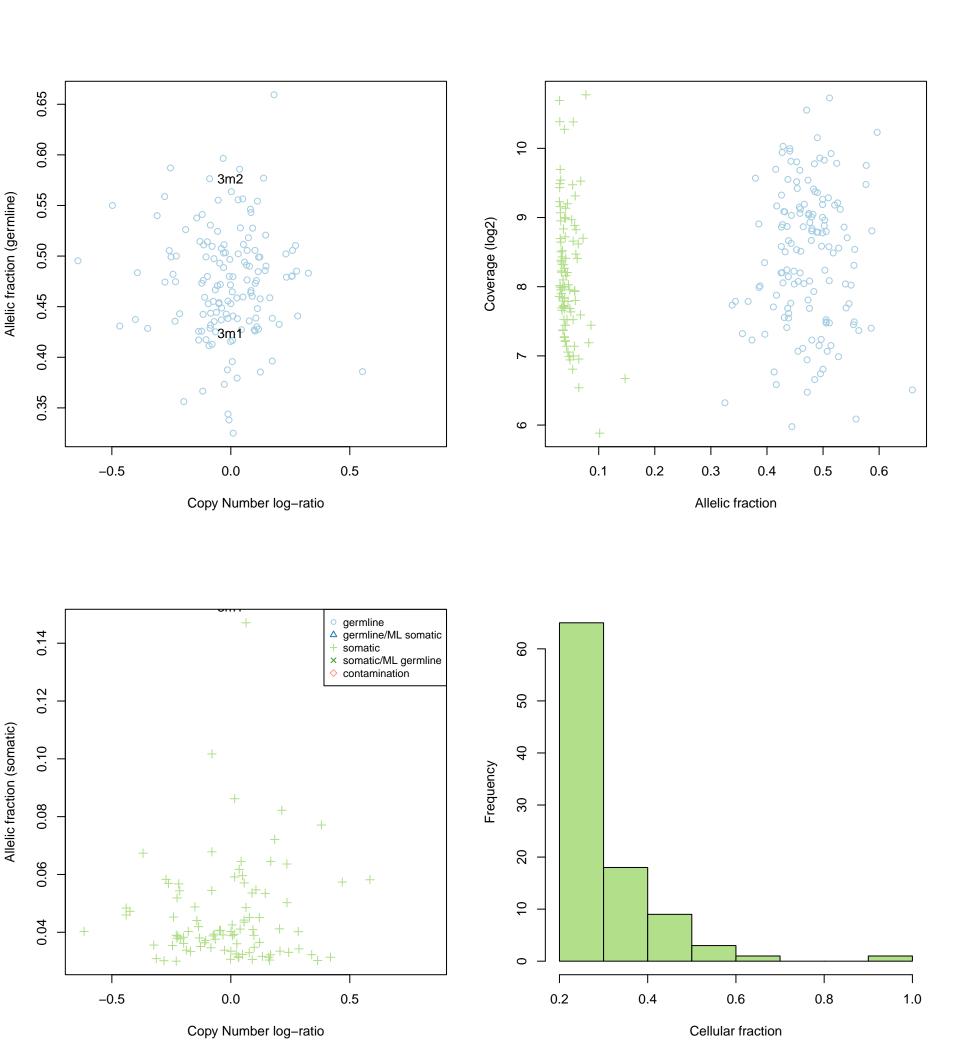
Purity: 0.36 Tumor ploidy: 3.01



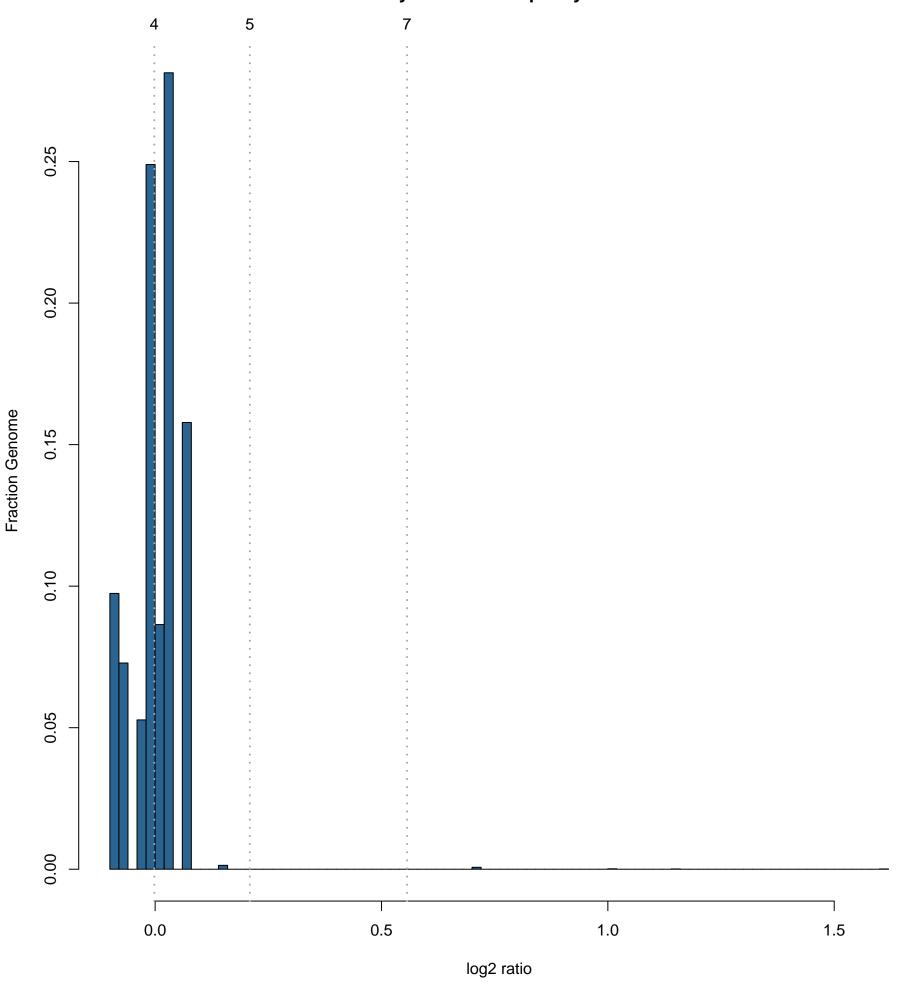


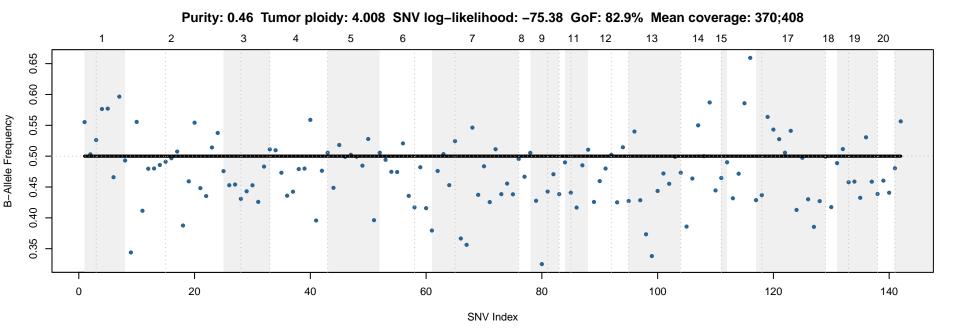


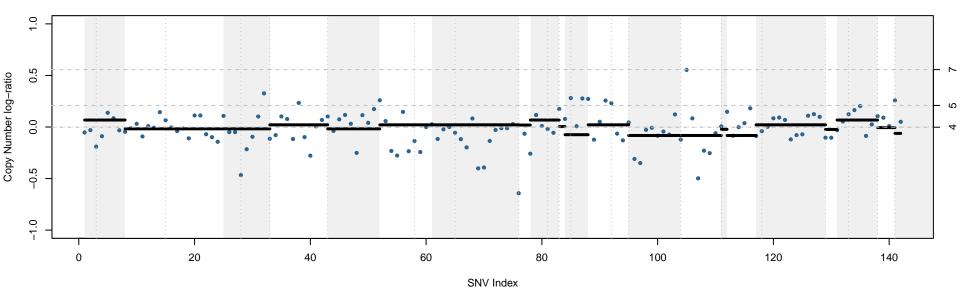


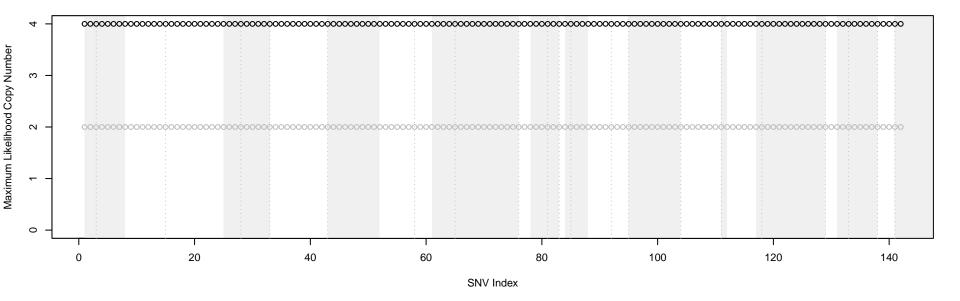


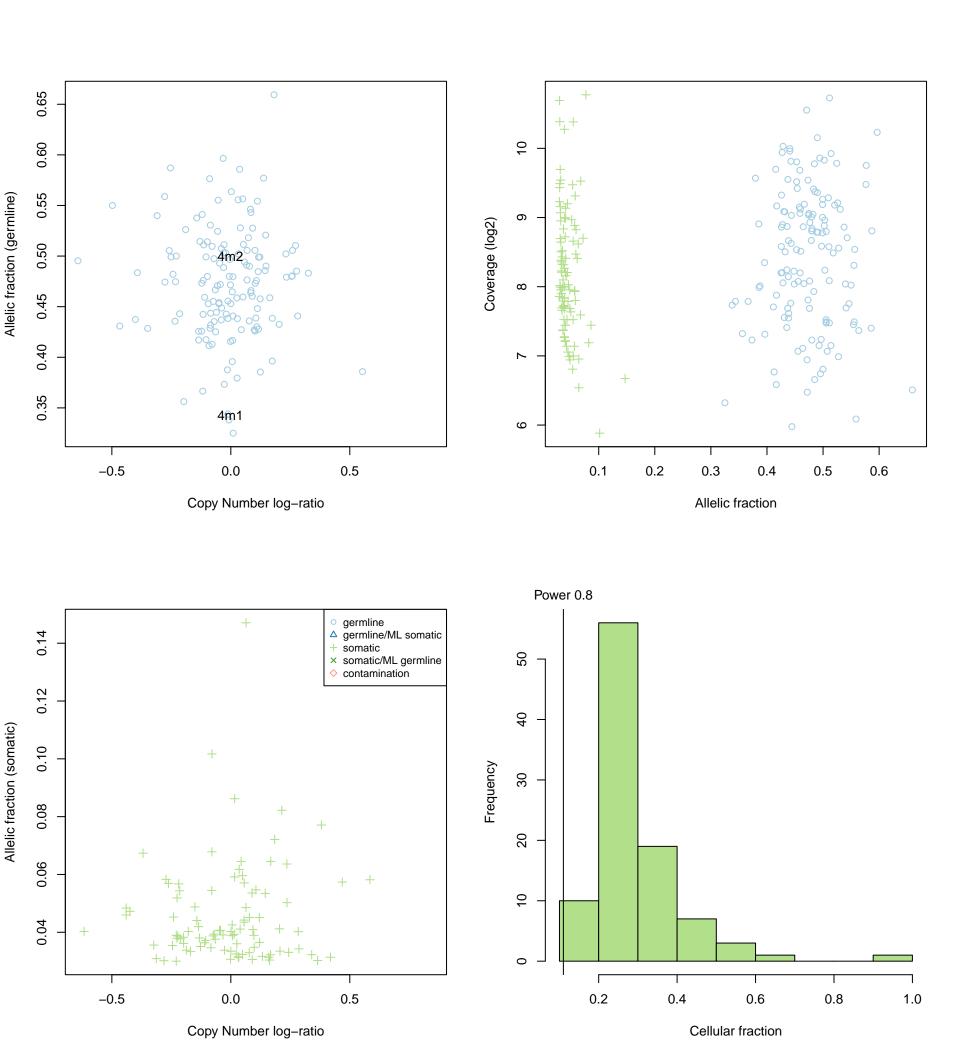
Purity: 0.46 Tumor ploidy: 4.008



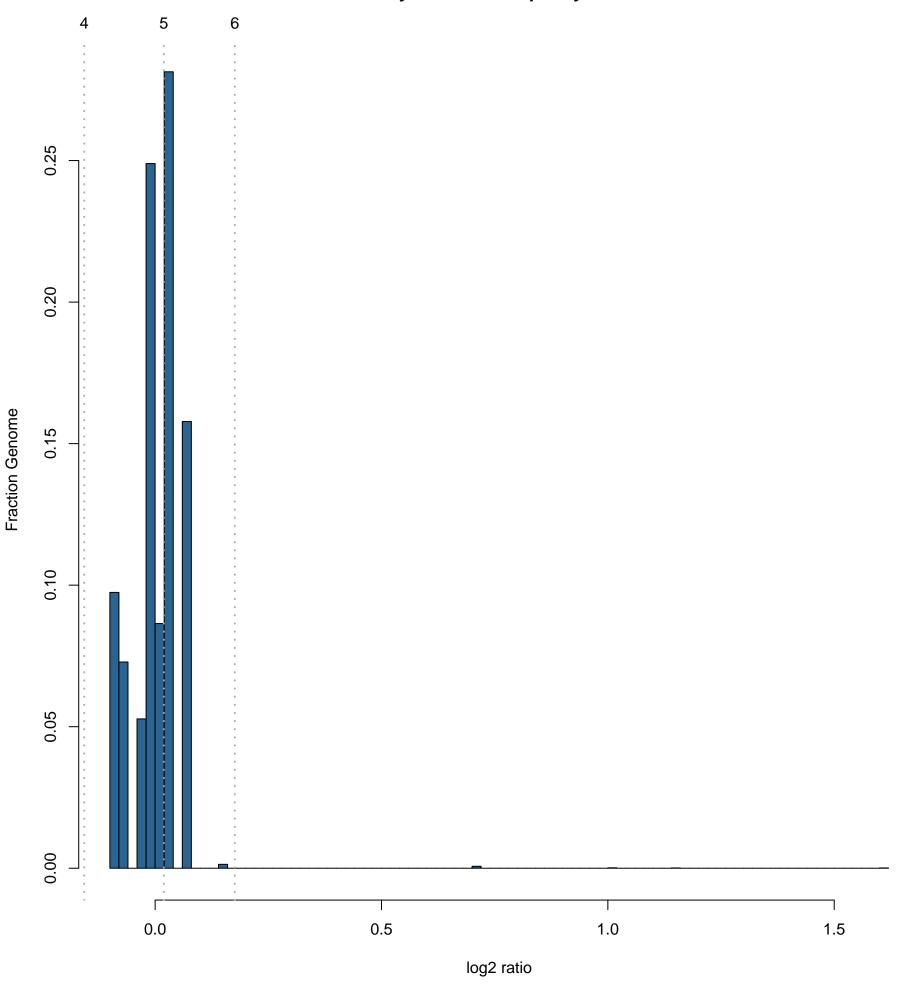


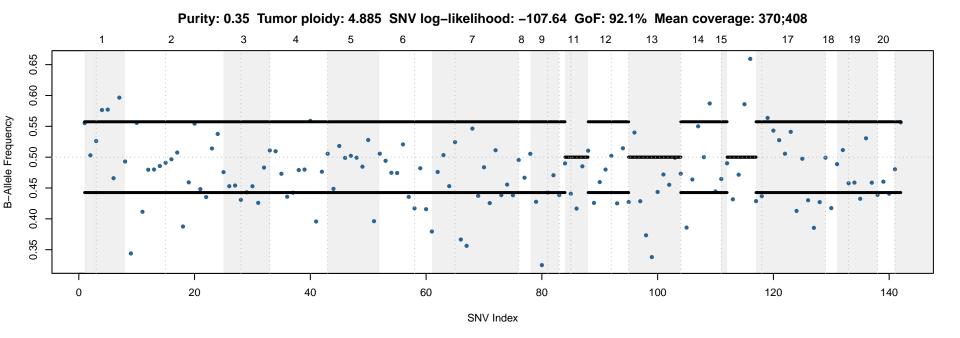


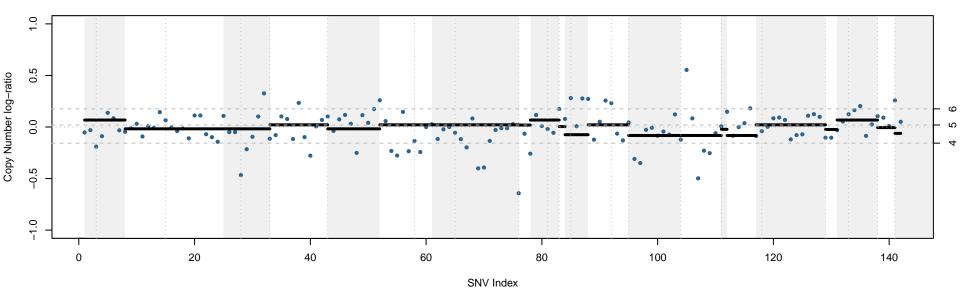


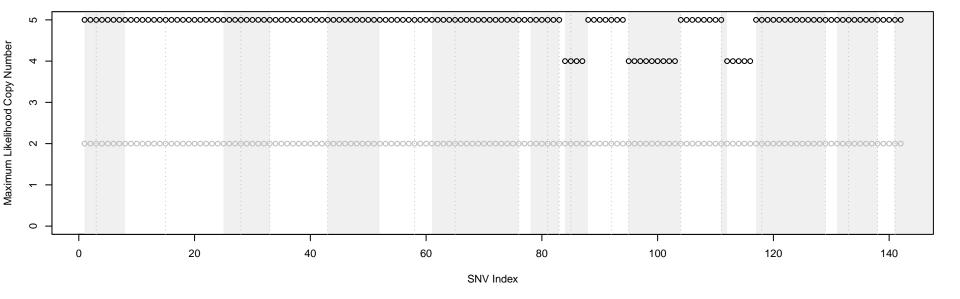


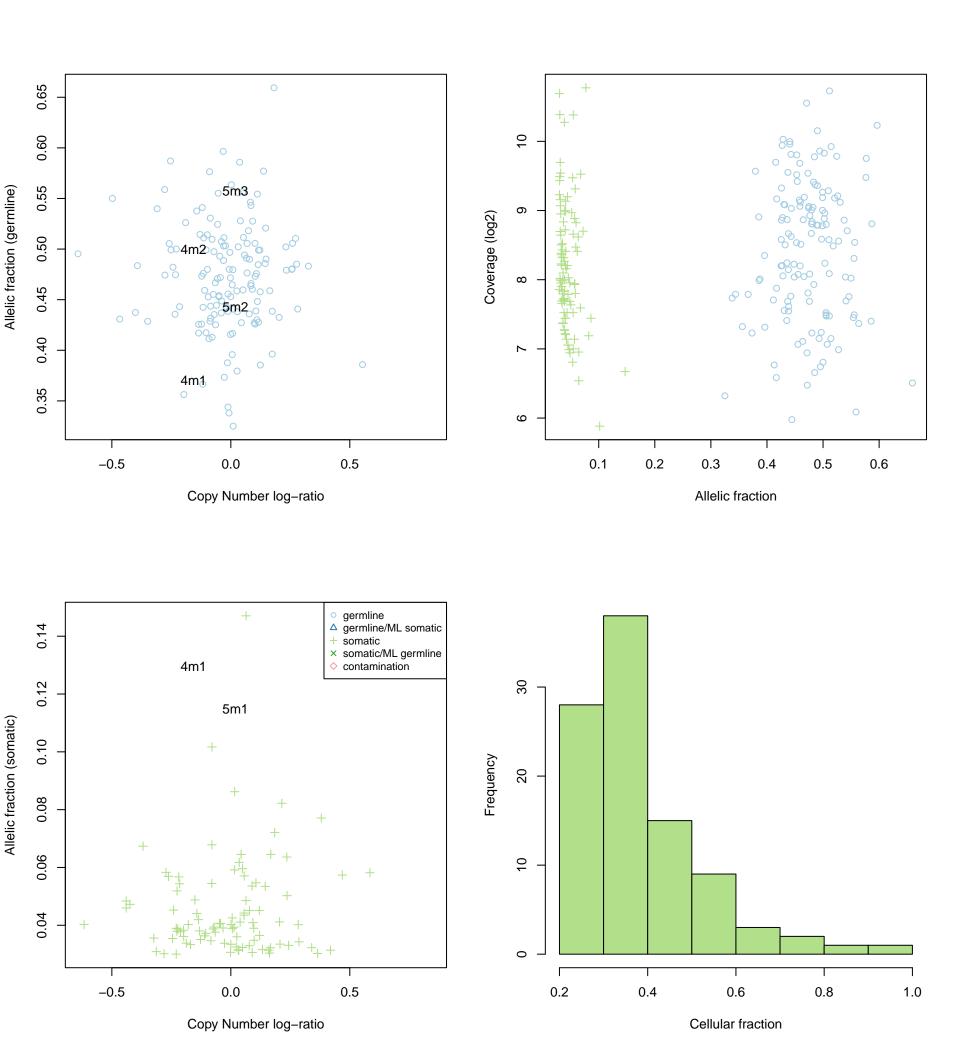
Purity: 0.35 Tumor ploidy: 4.885



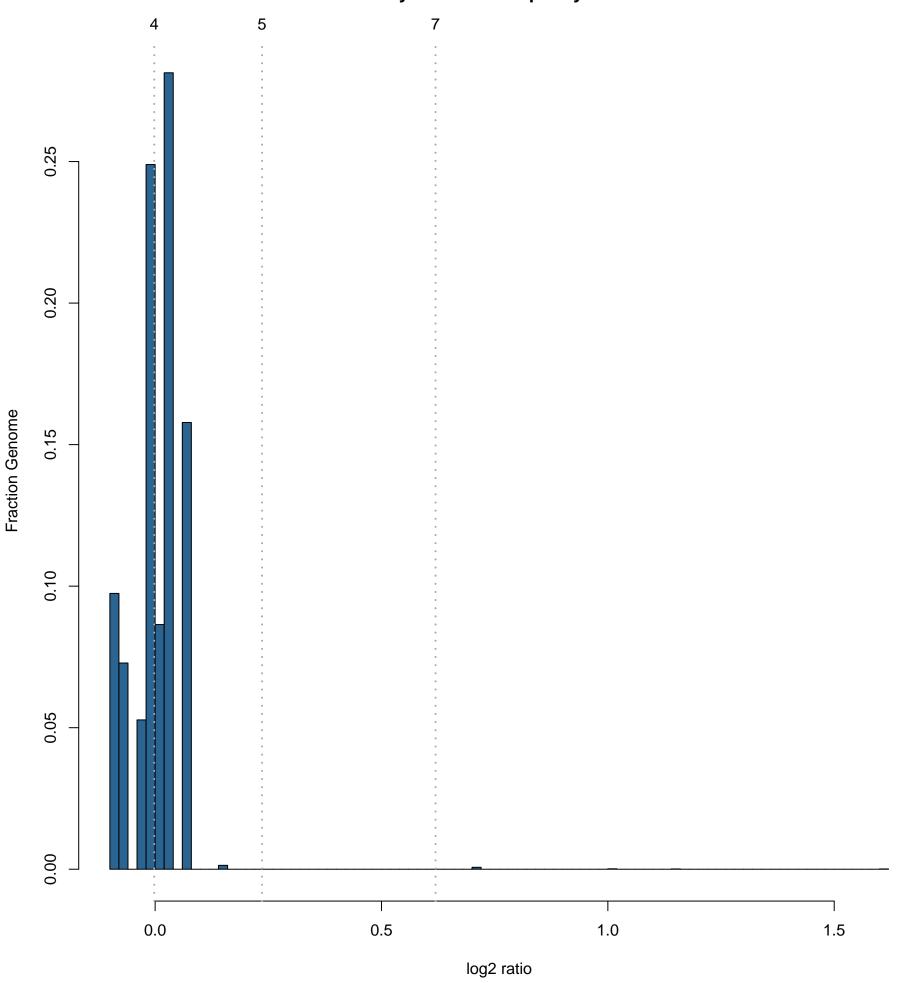


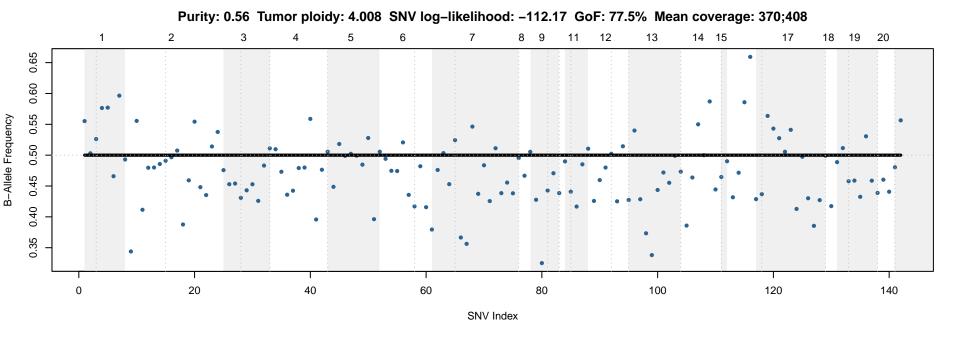


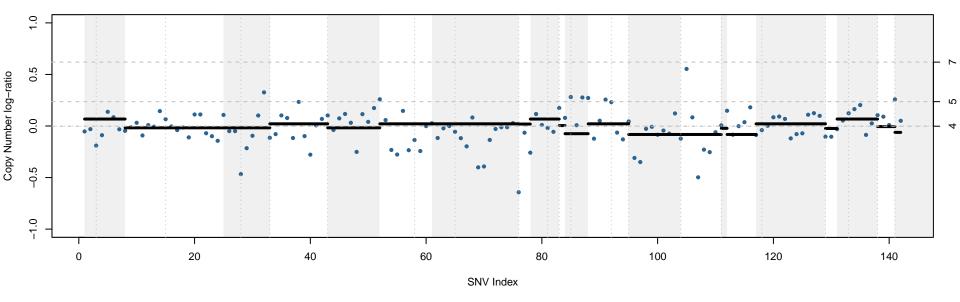


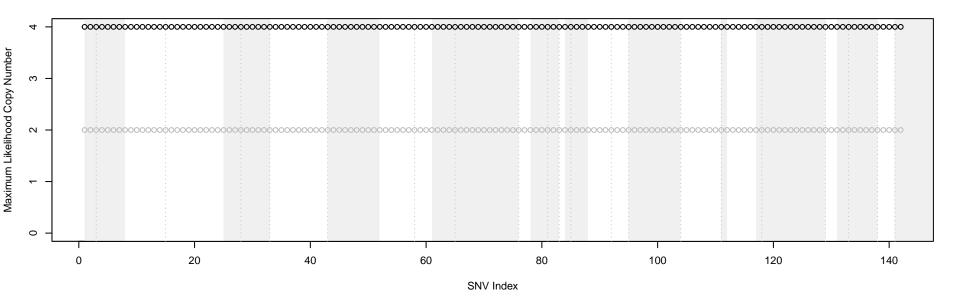


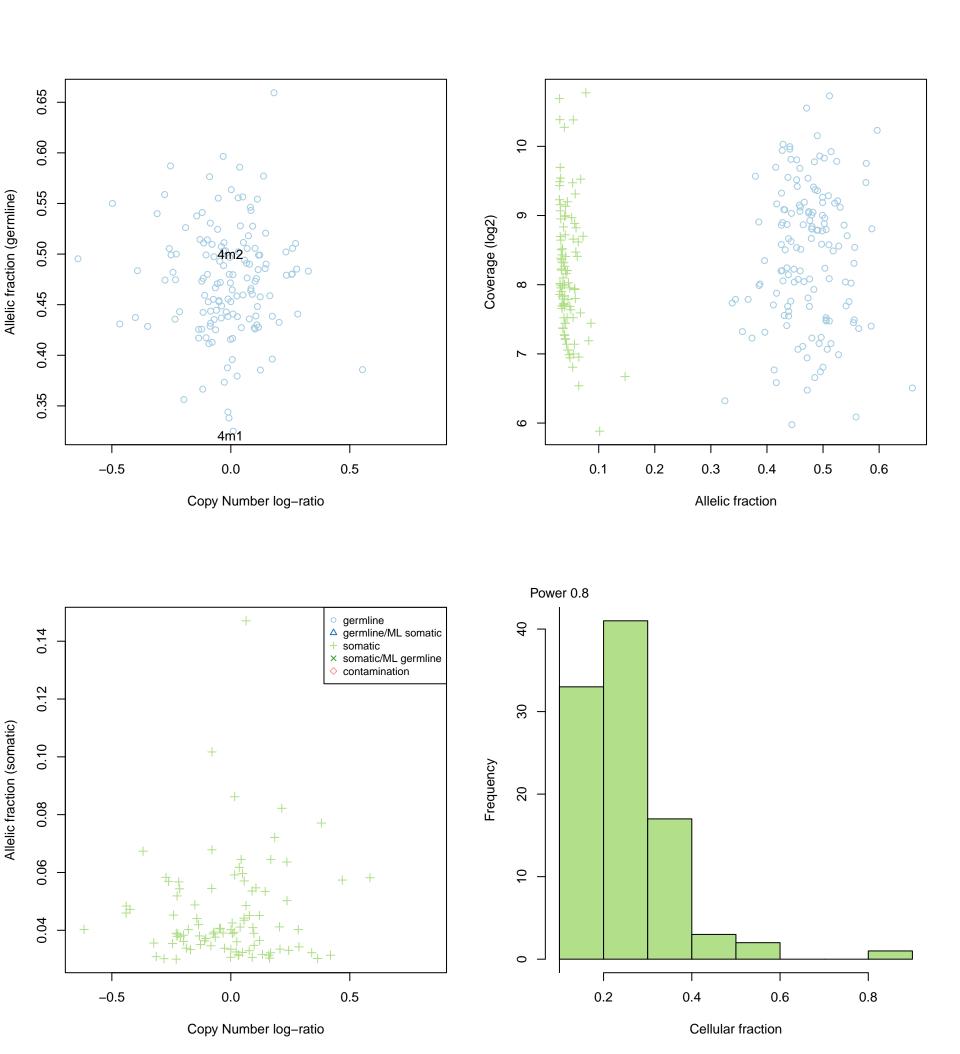
Purity: 0.56 Tumor ploidy: 4.008



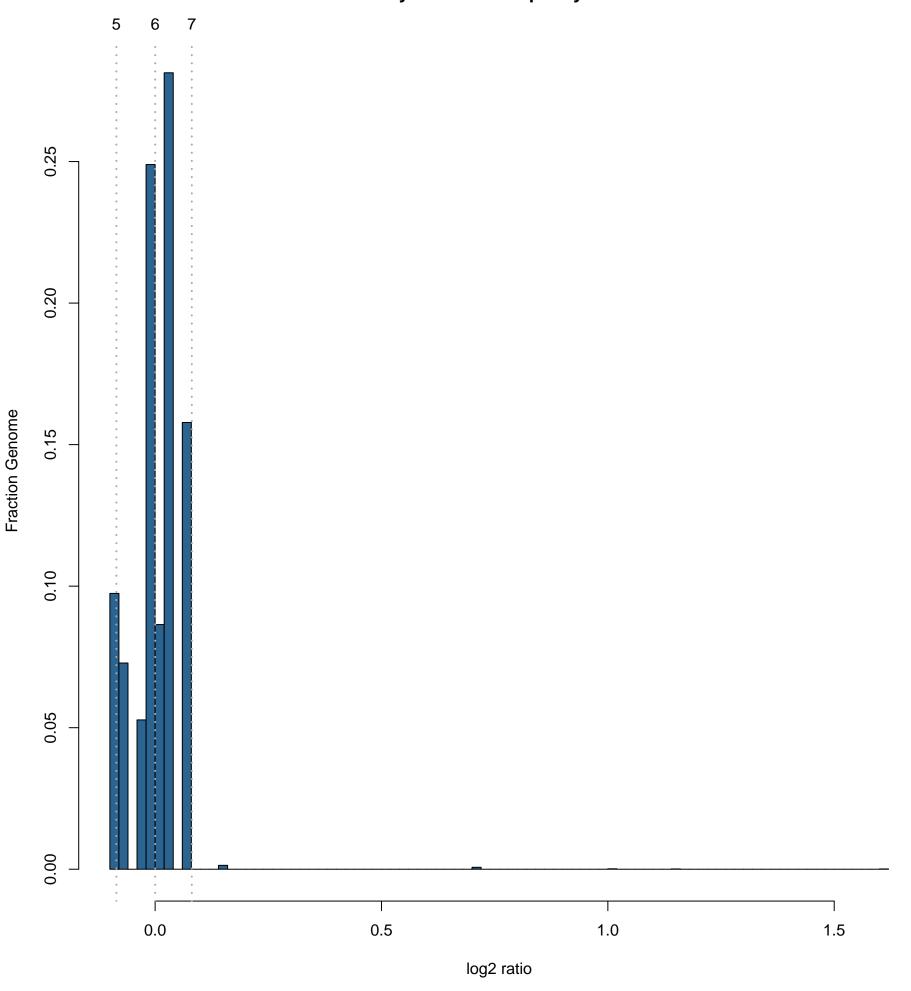


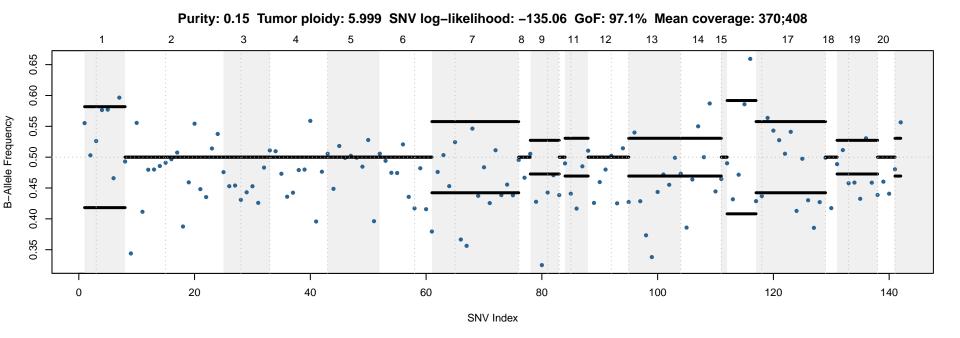


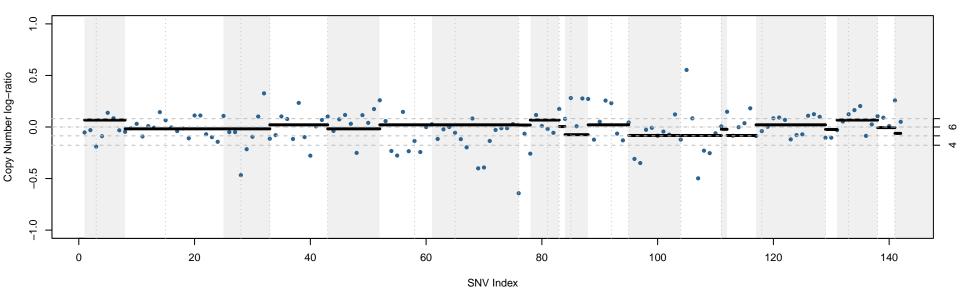


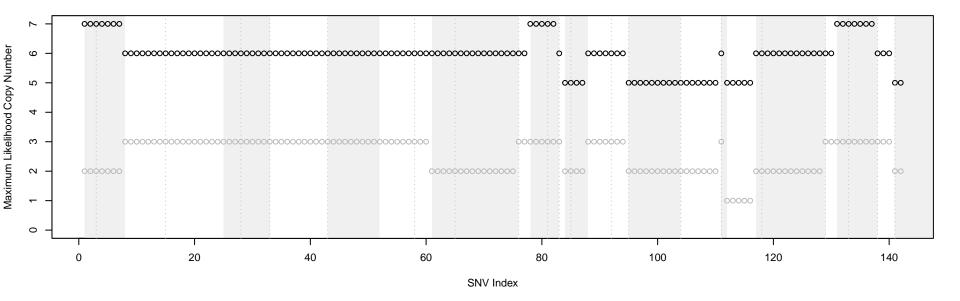


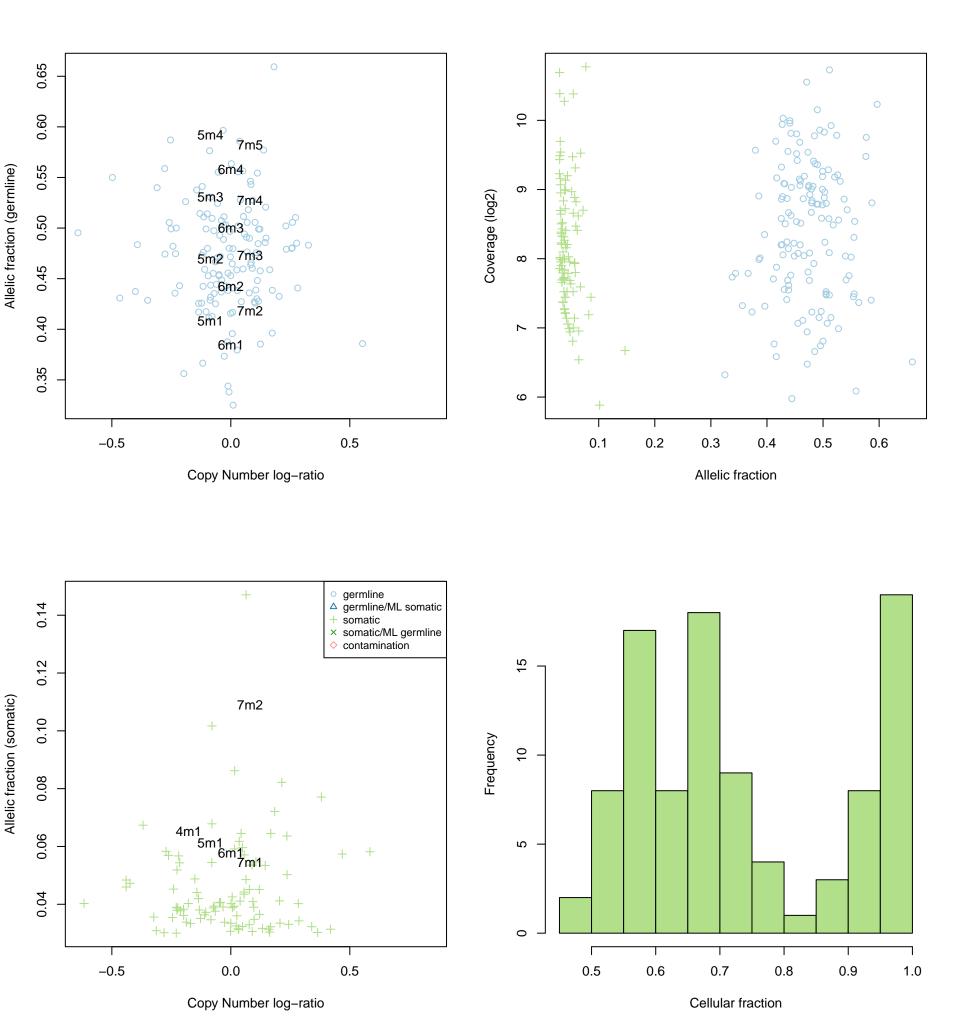
Purity: 0.15 Tumor ploidy: 5.999



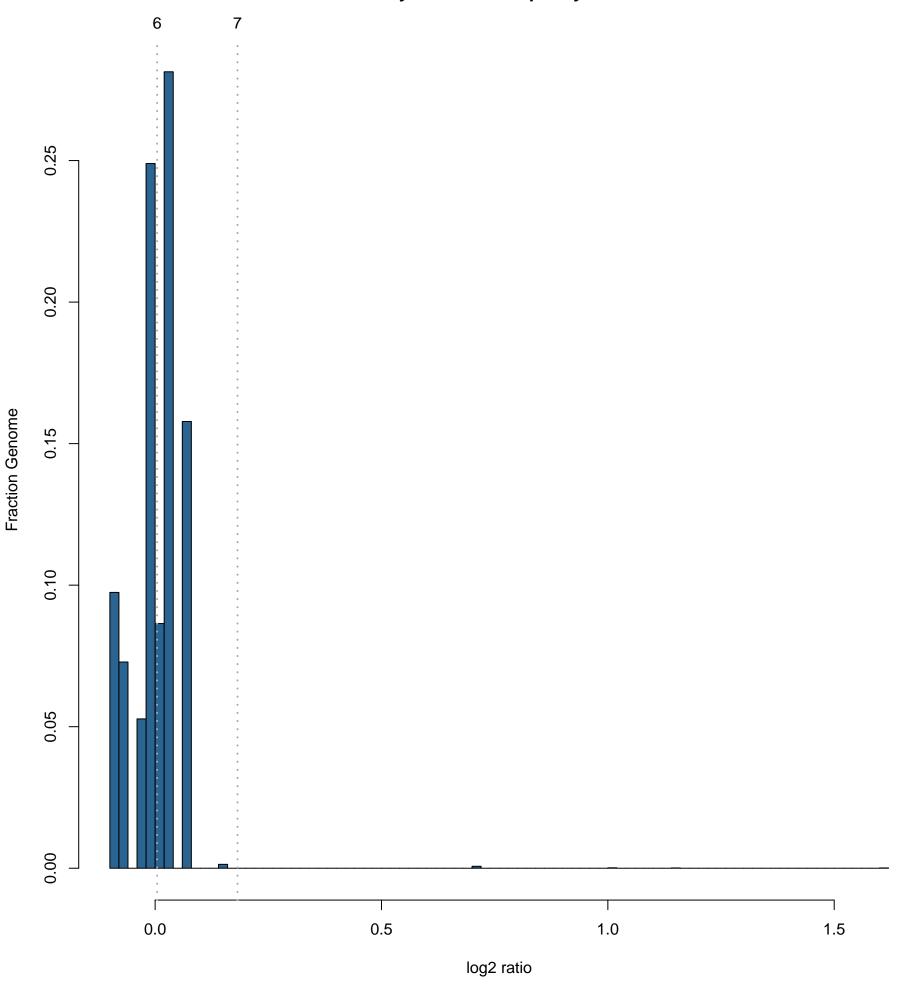


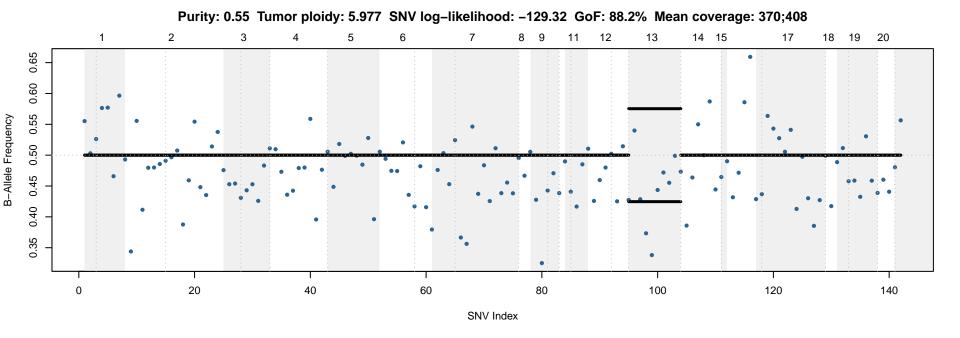


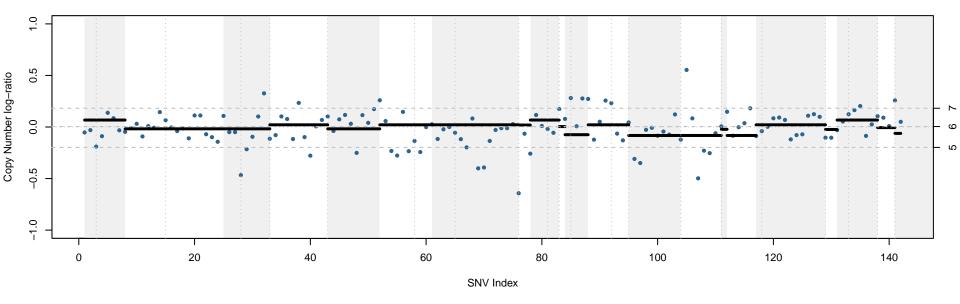


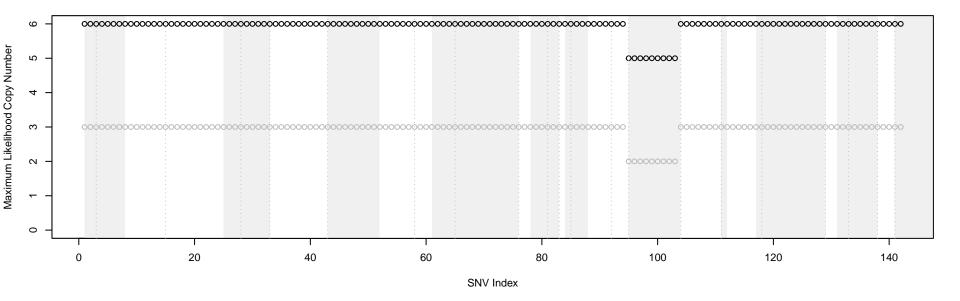


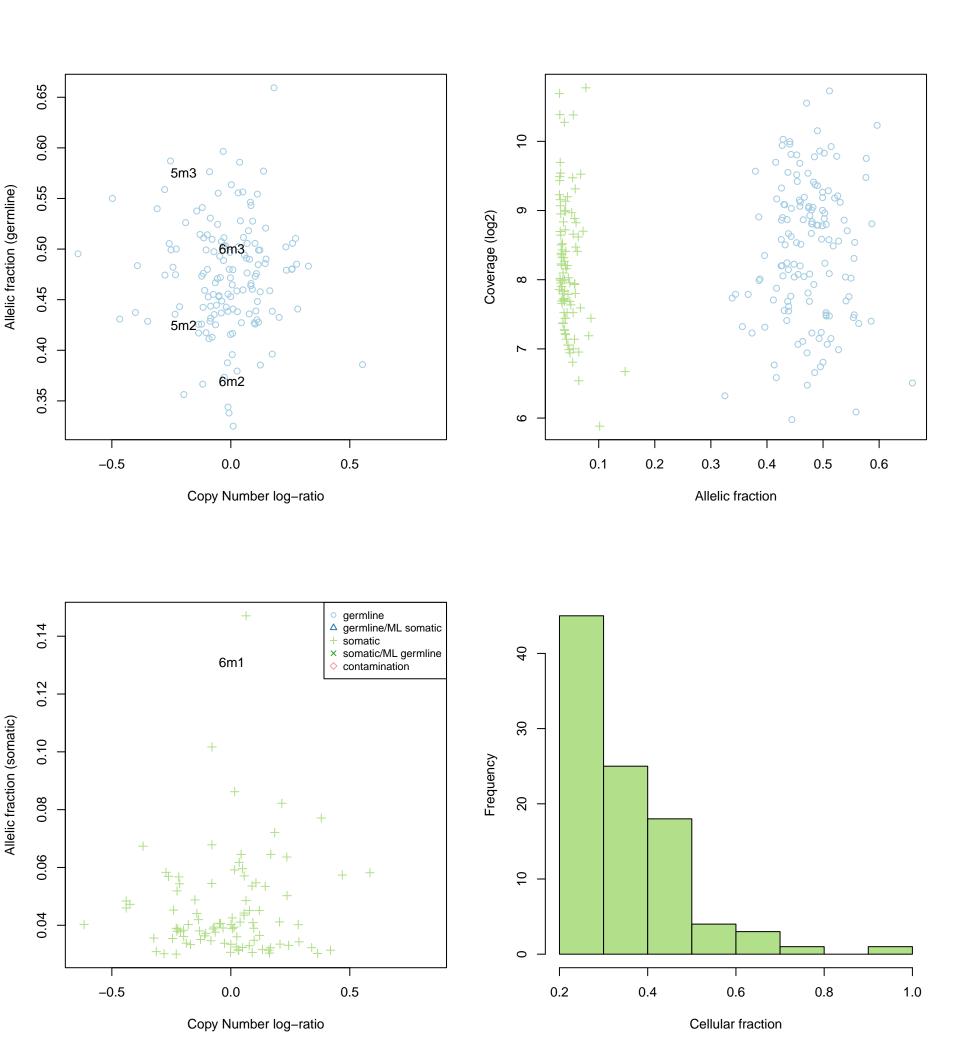
Purity: 0.55 Tumor ploidy: 5.977



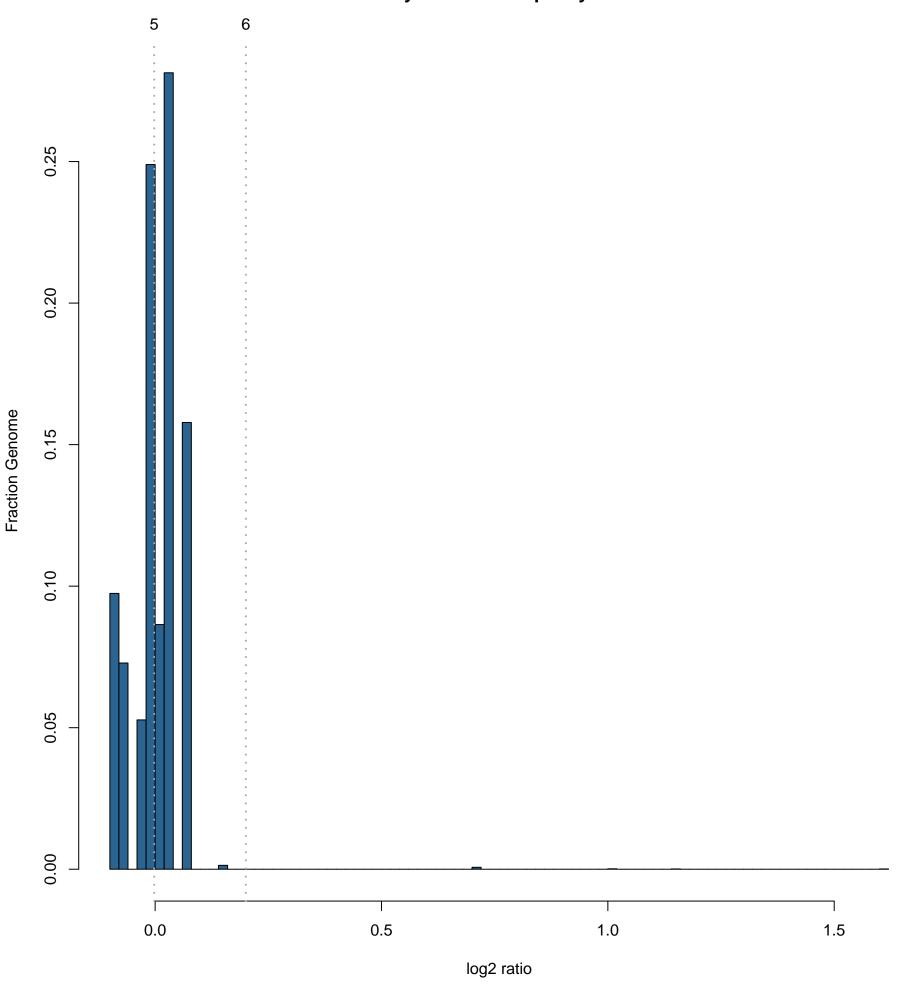


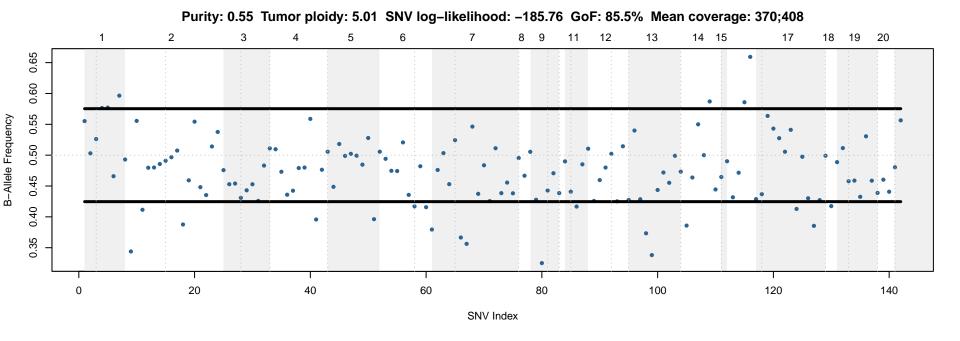


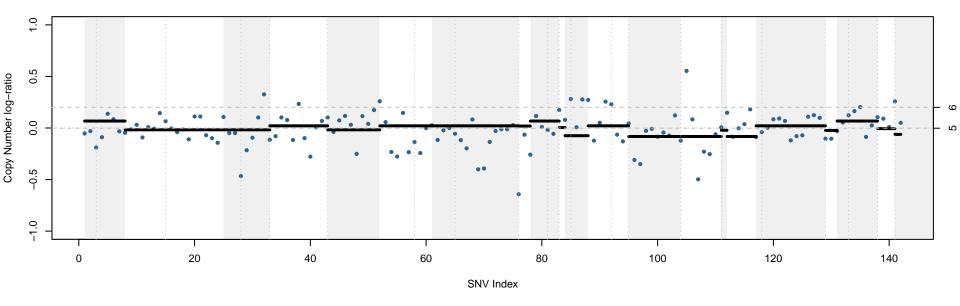


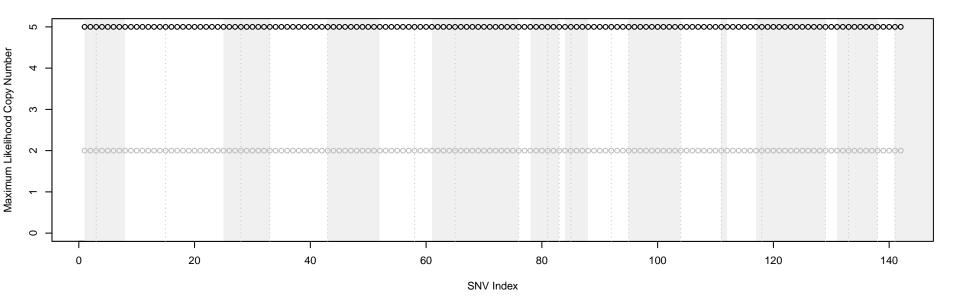


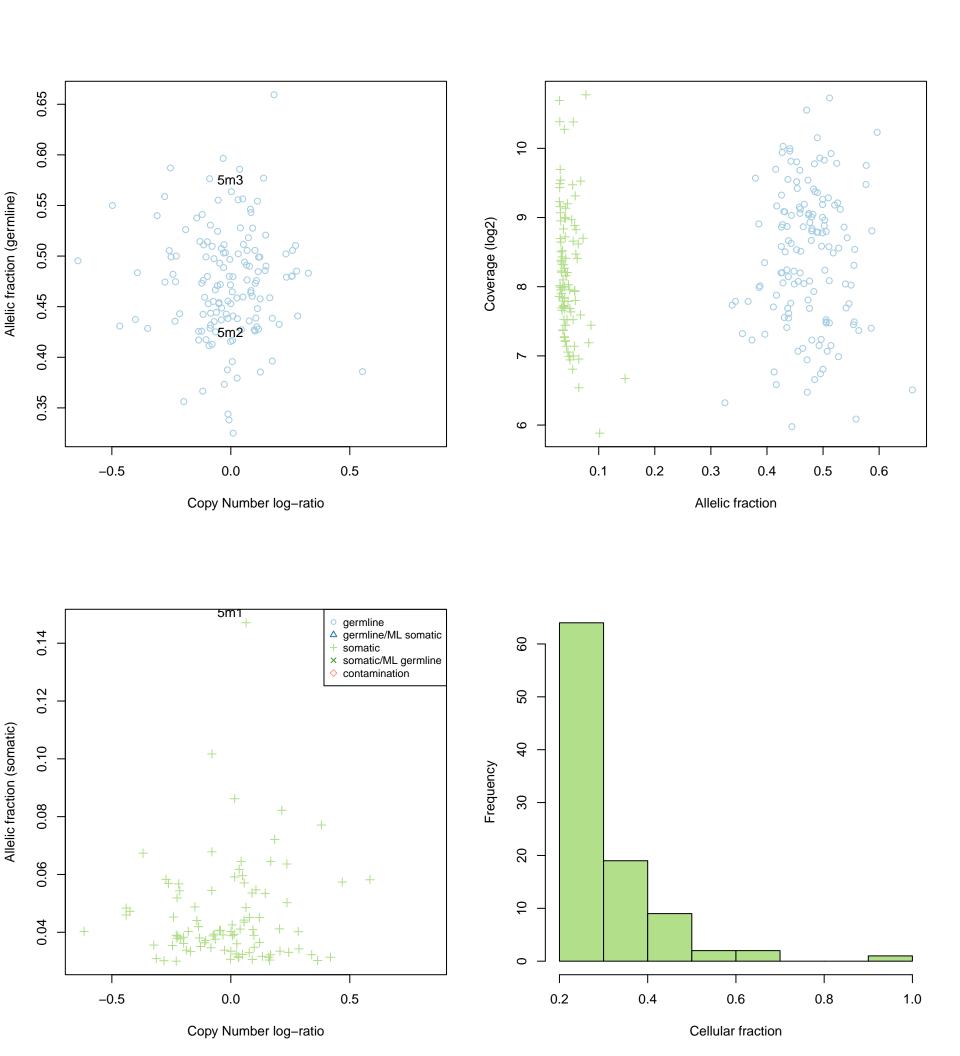
Purity: 0.55 Tumor ploidy: 5.01



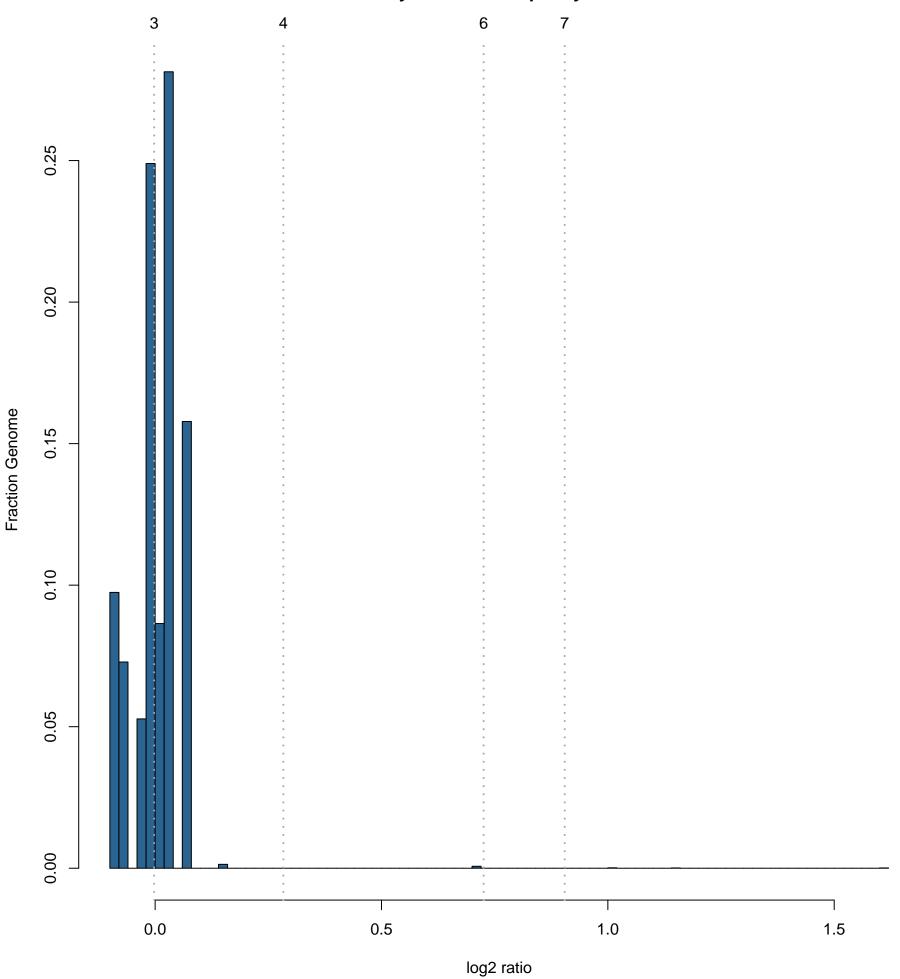


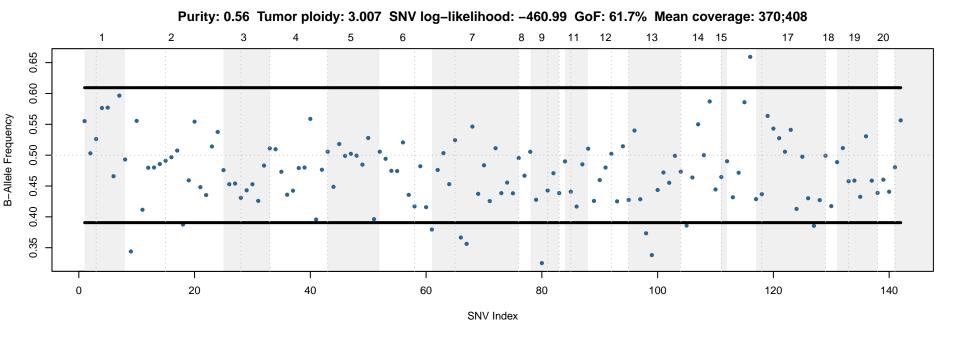


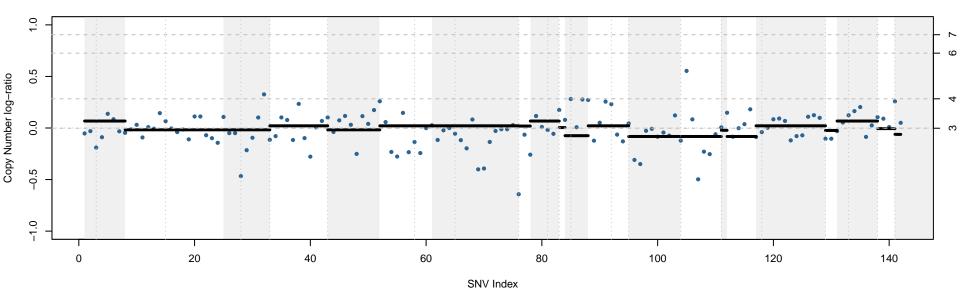


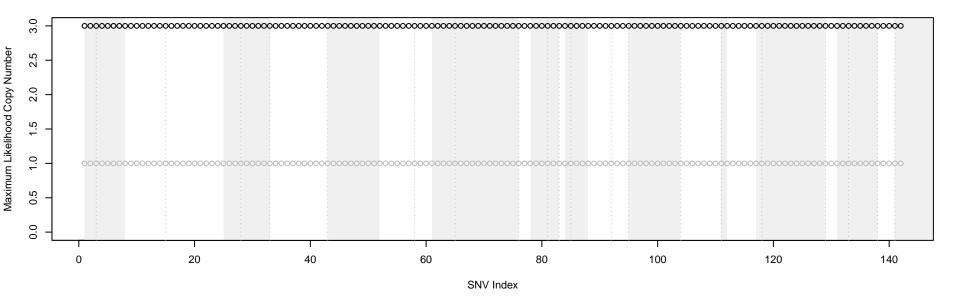


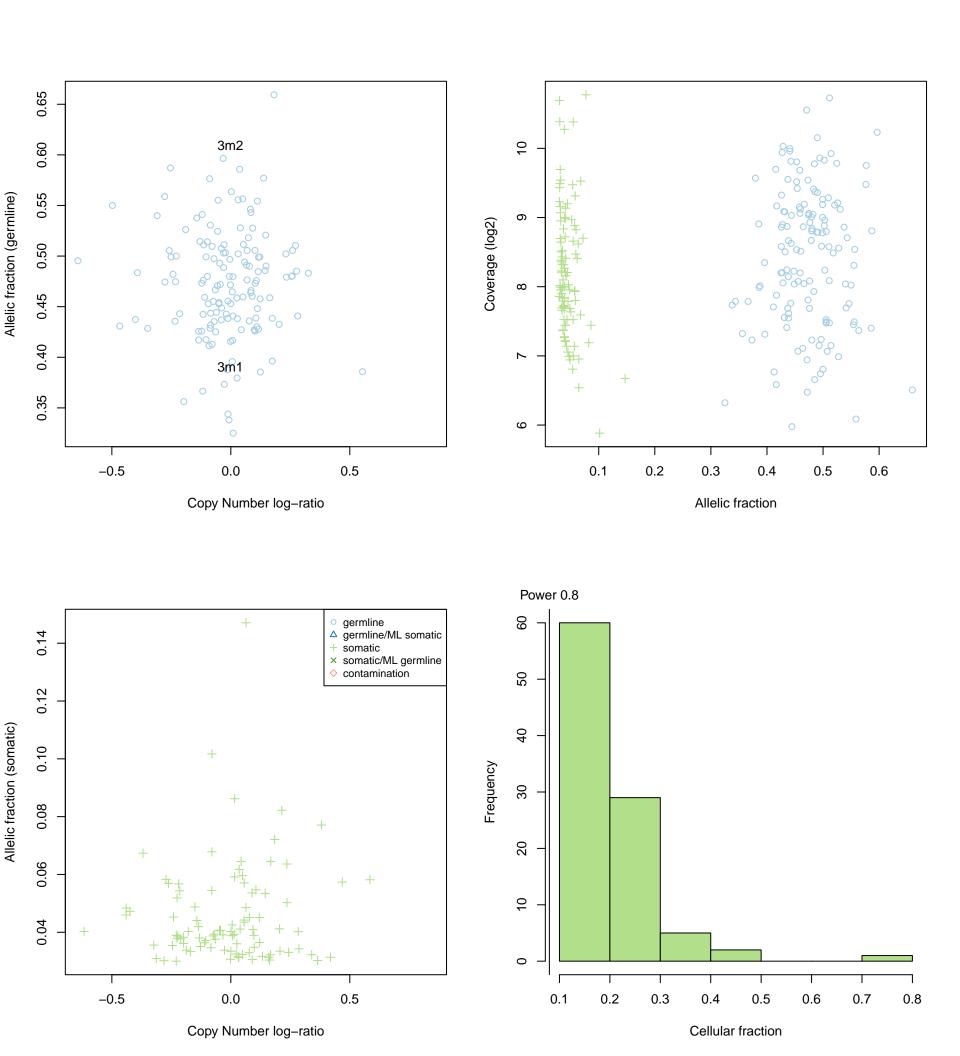
Purity: 0.56 Tumor ploidy: 3.007





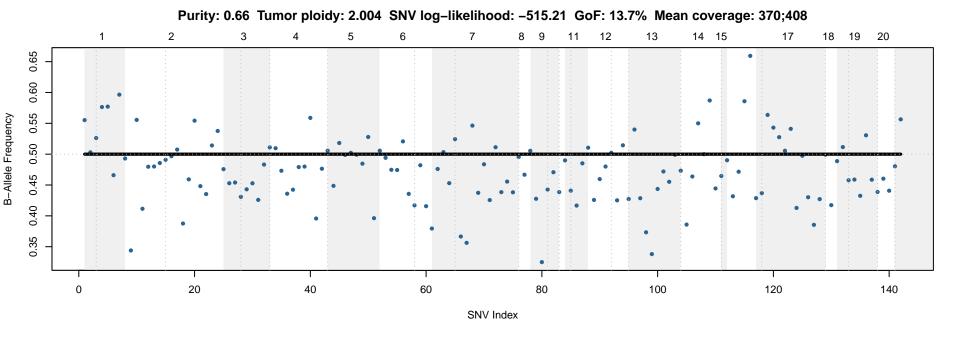


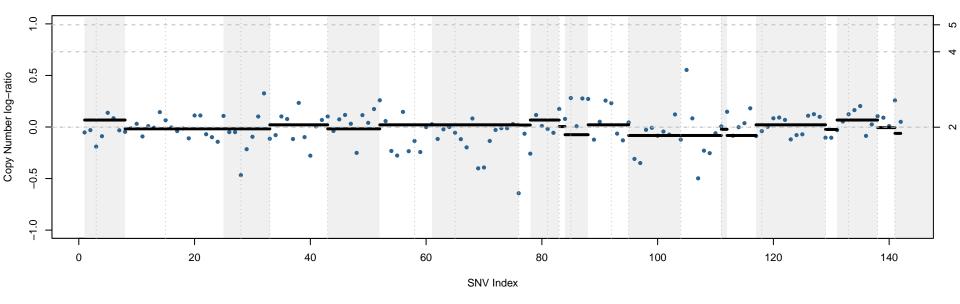


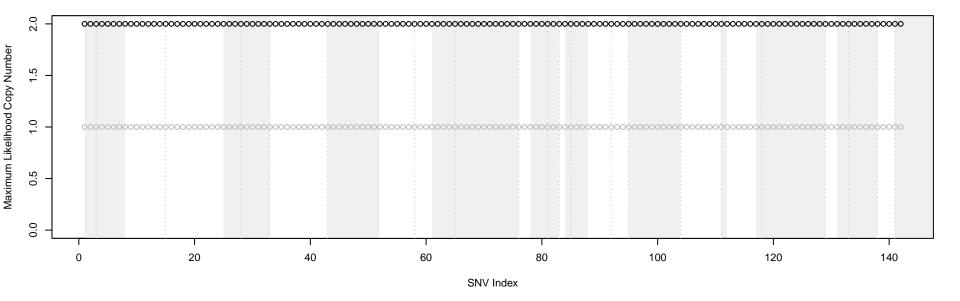


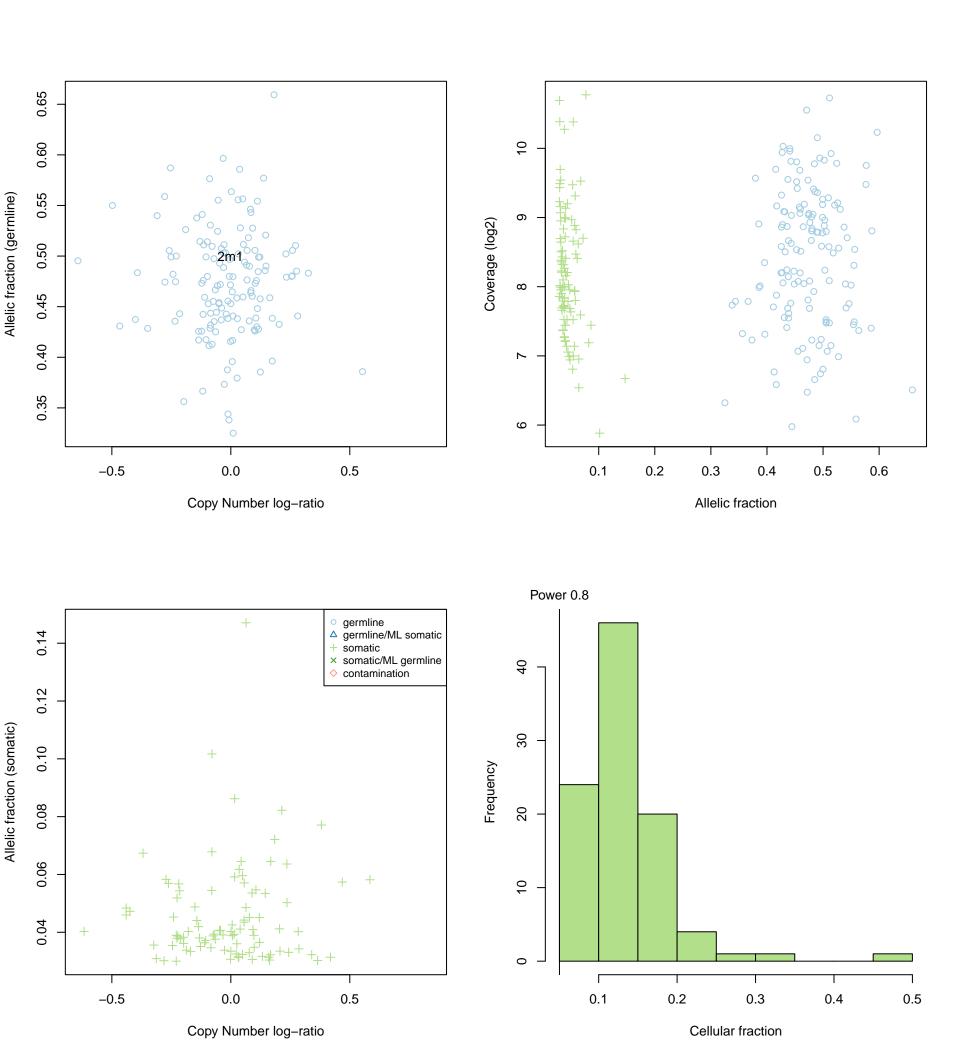
Purity: 0.66 Tumor ploidy: 2.004 2 7 6 0.25 0.20 Fraction Genome 0.15 0.10 0.05 0.00 0.0 0.5 1.0 1.5

log2 ratio

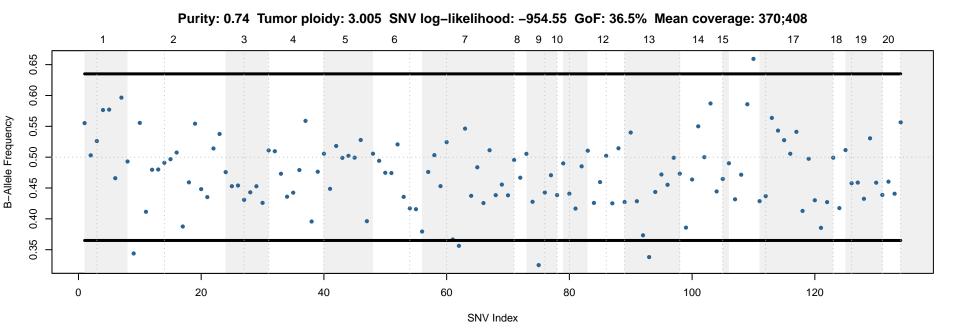


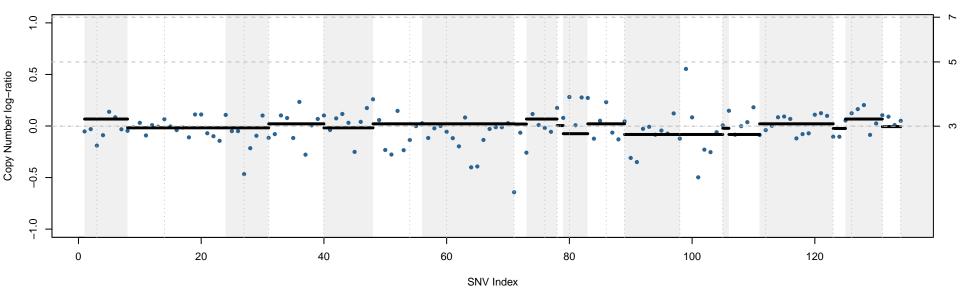


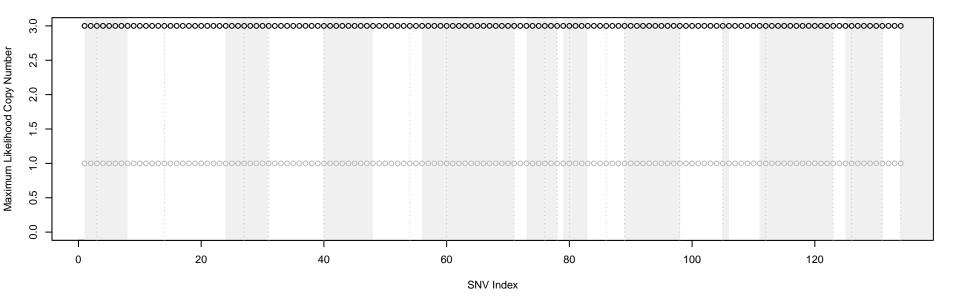


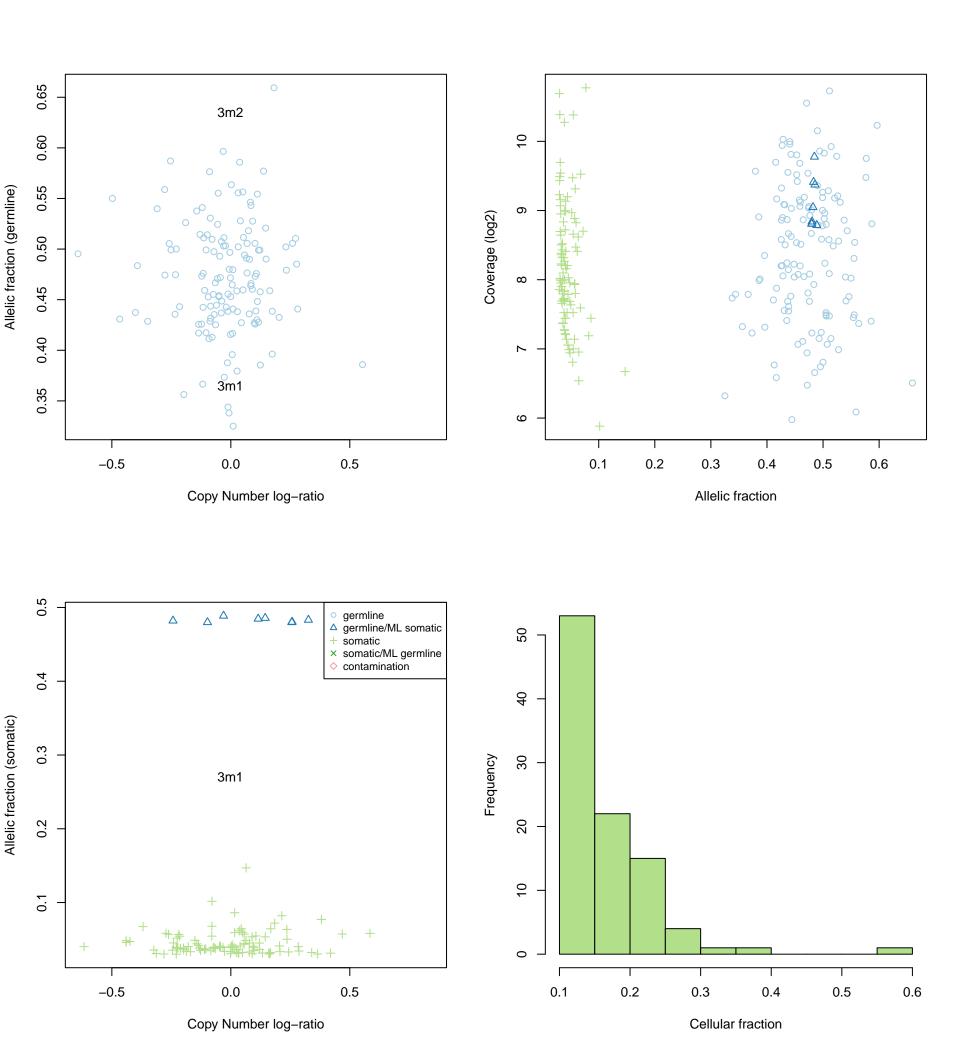


Purity: 0.74 Tumor ploidy: 3.005 3 0.25 0.20 Fraction Genome 0.15 0.10 0.00 0.0 0.5 1.0 1.5 log2 ratio



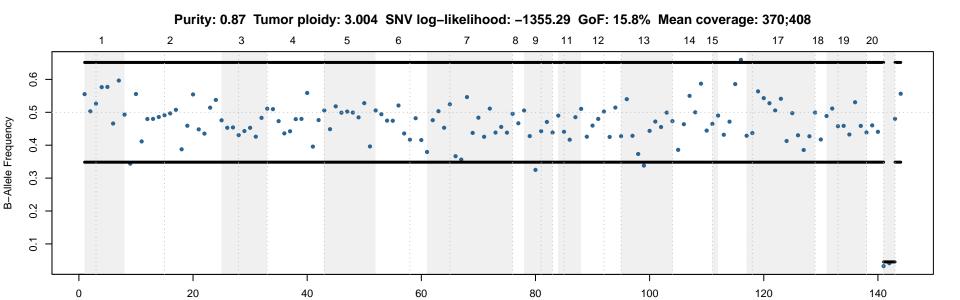






Purity: 0.87 Tumor ploidy: 3.004 3 7 0.25 0.20 Fraction Genome 0.15 0.10 0.00 0.0 0.5 1.0 1.5

log2 ratio



SCNA-fit log-likelihood: -14820.84

SNV Index

