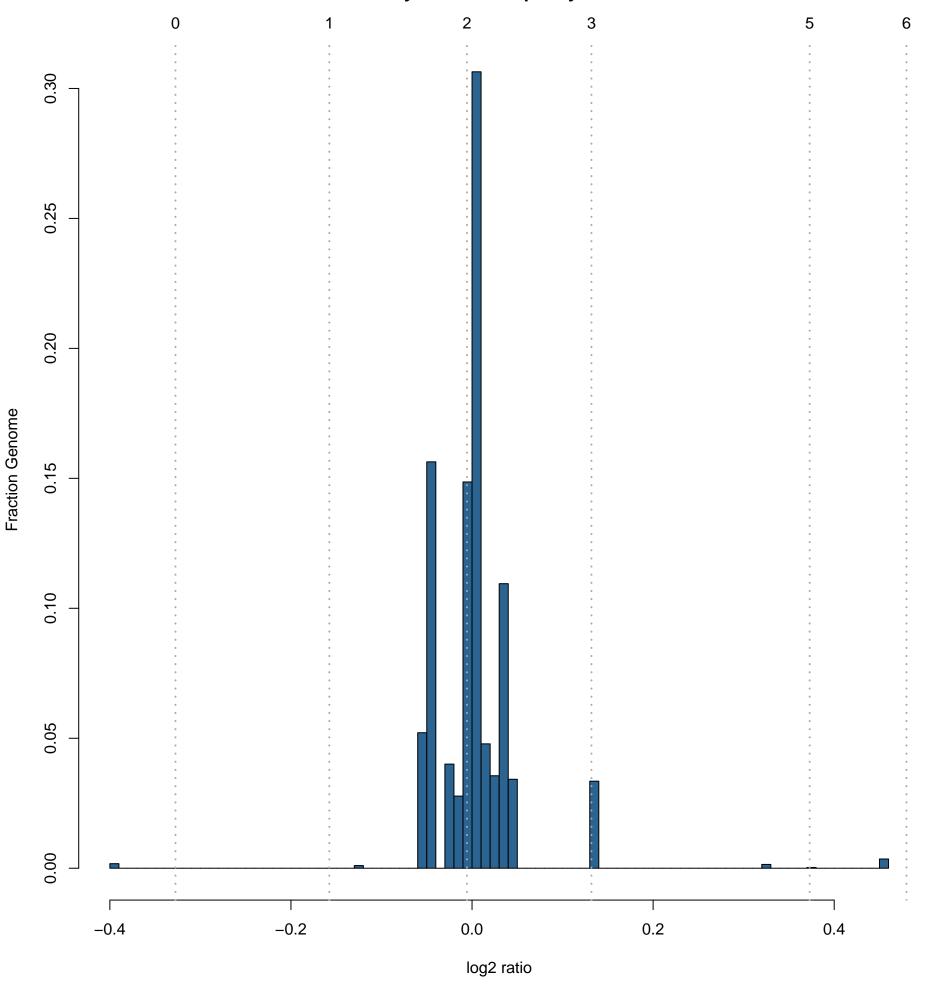
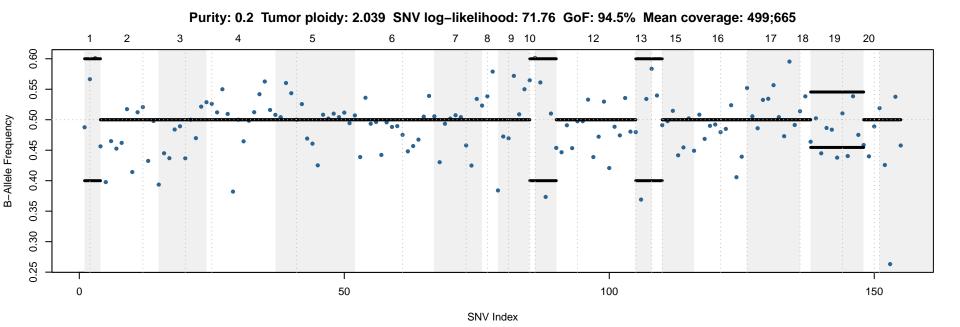
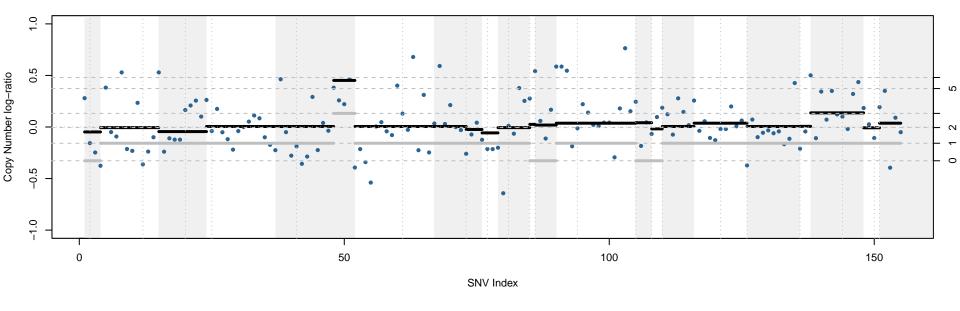
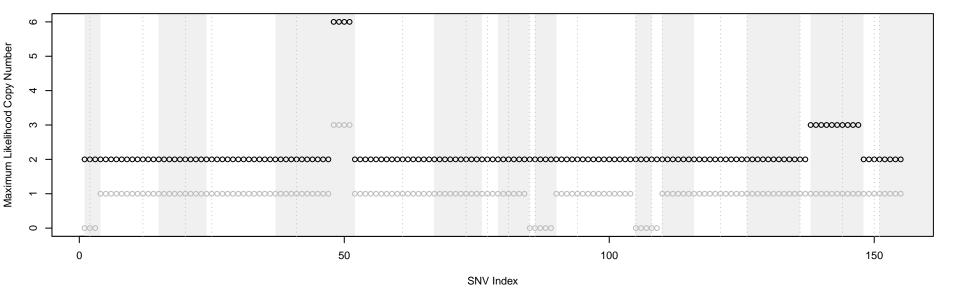
Purity: 0.2 Tumor ploidy: 2.039

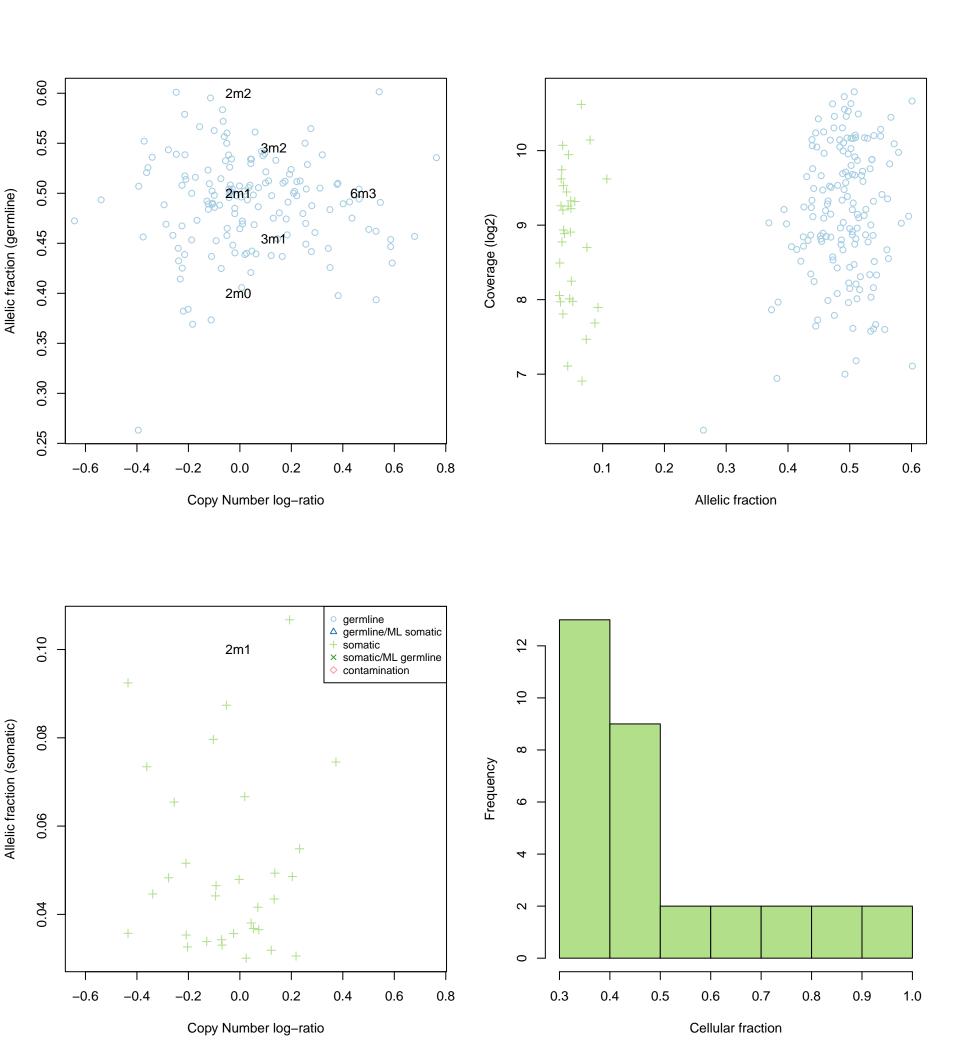




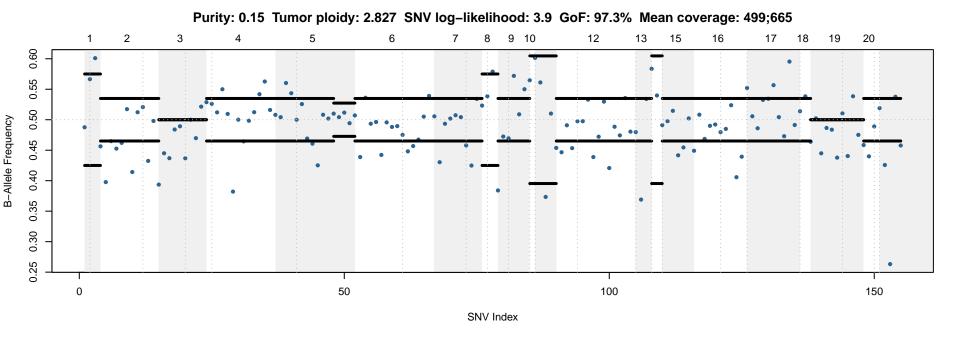
## SCNA-fit log-likelihood: -7368.17



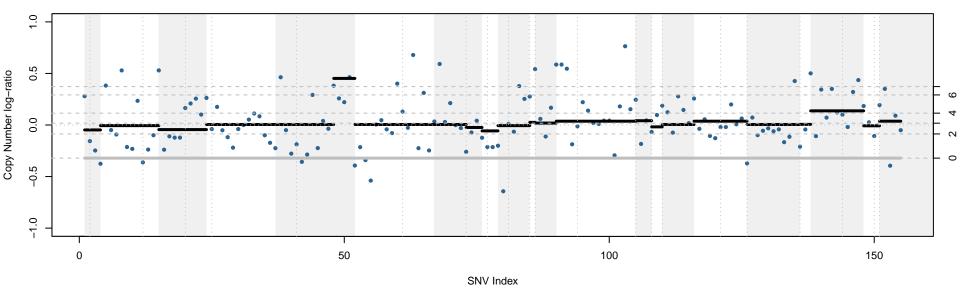


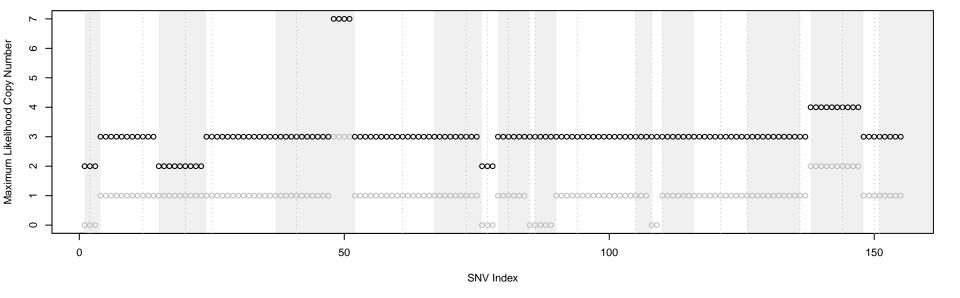


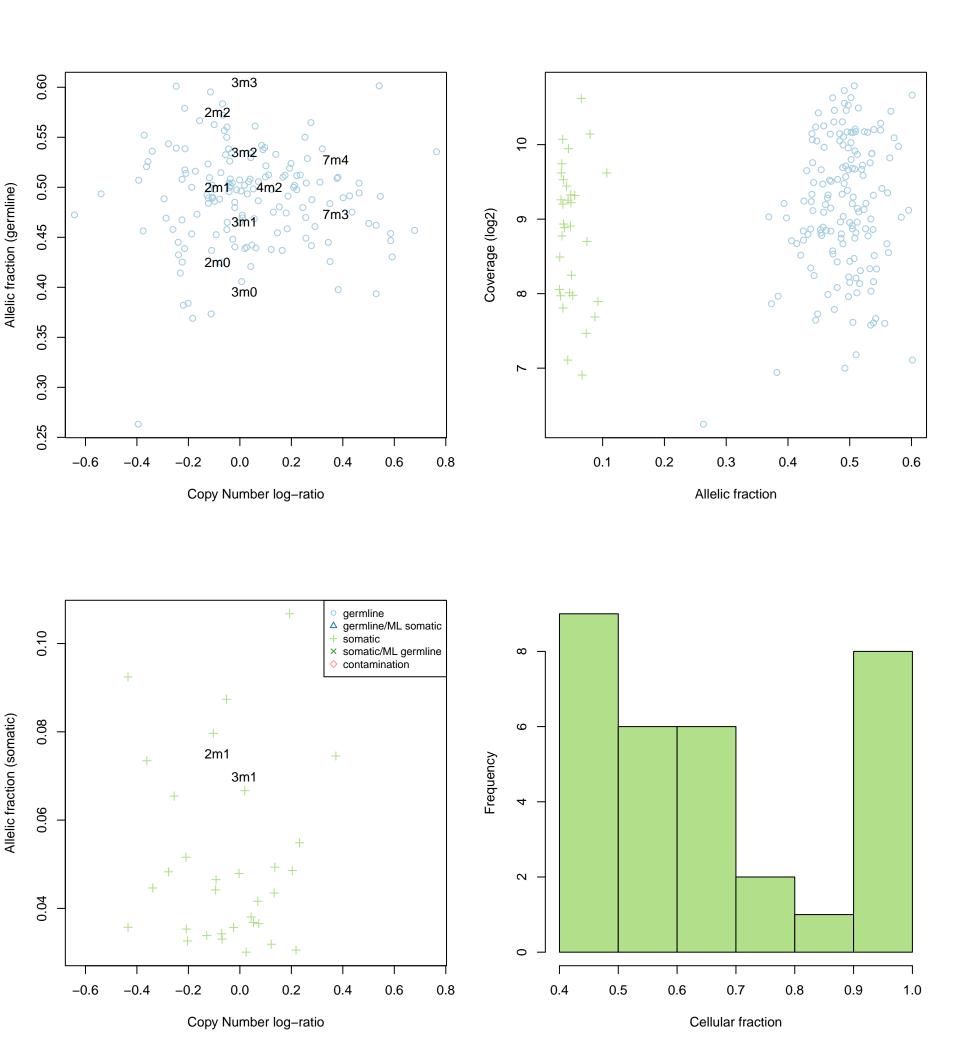
Purity: 0.15 Tumor ploidy: 2.827 0 2 6 7 3 0.30 0.25 0.20 Fraction Genome 0.15 0.10 0.05 0.00 -0.4 -0.2 0.0 0.2 0.4 log2 ratio



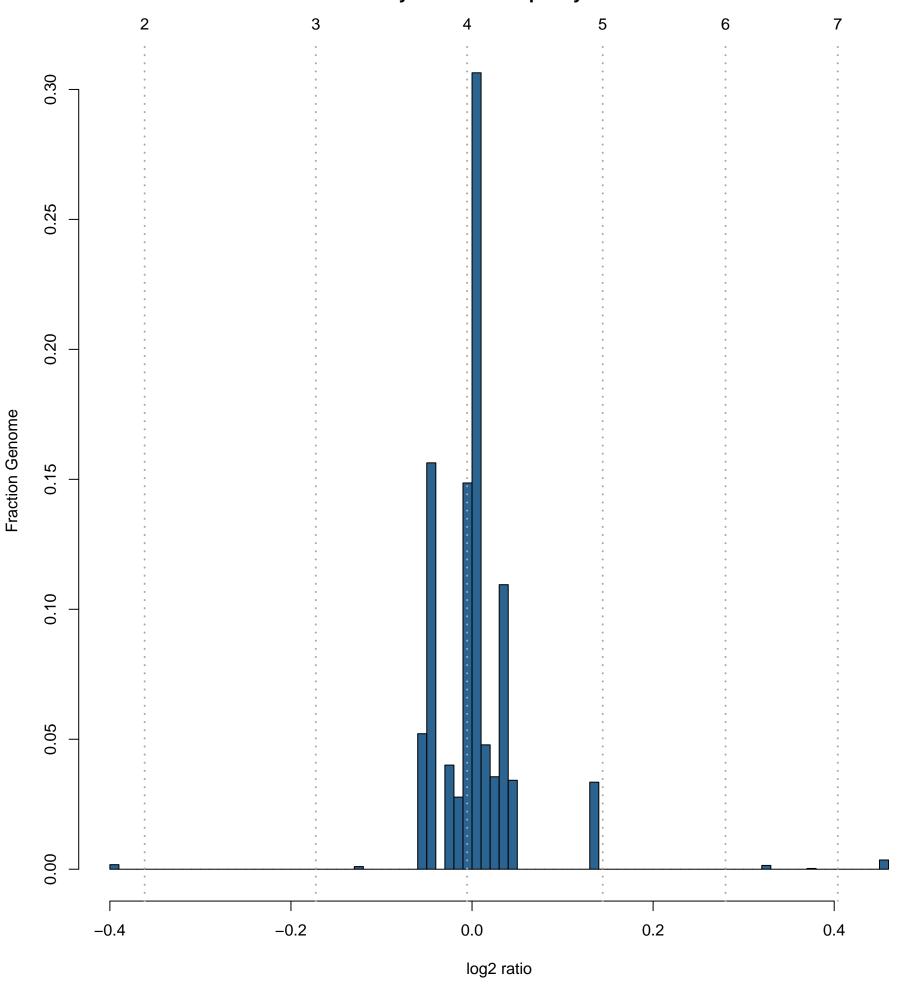
## SCNA-fit log-likelihood: -7353.8

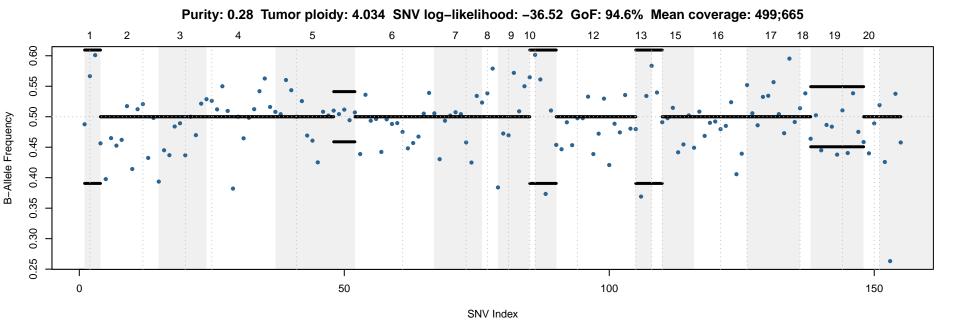




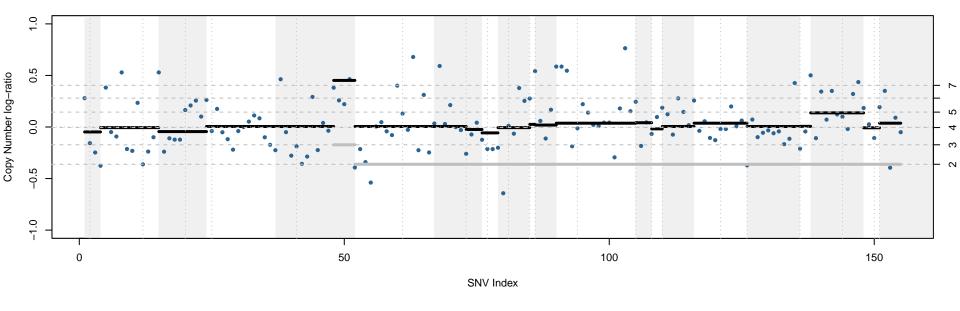


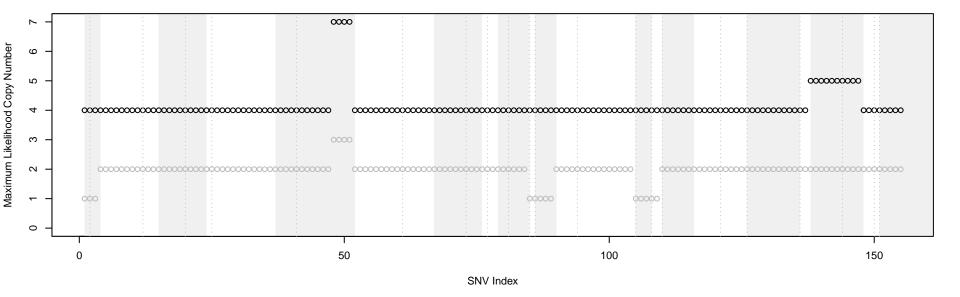
Purity: 0.28 Tumor ploidy: 4.034

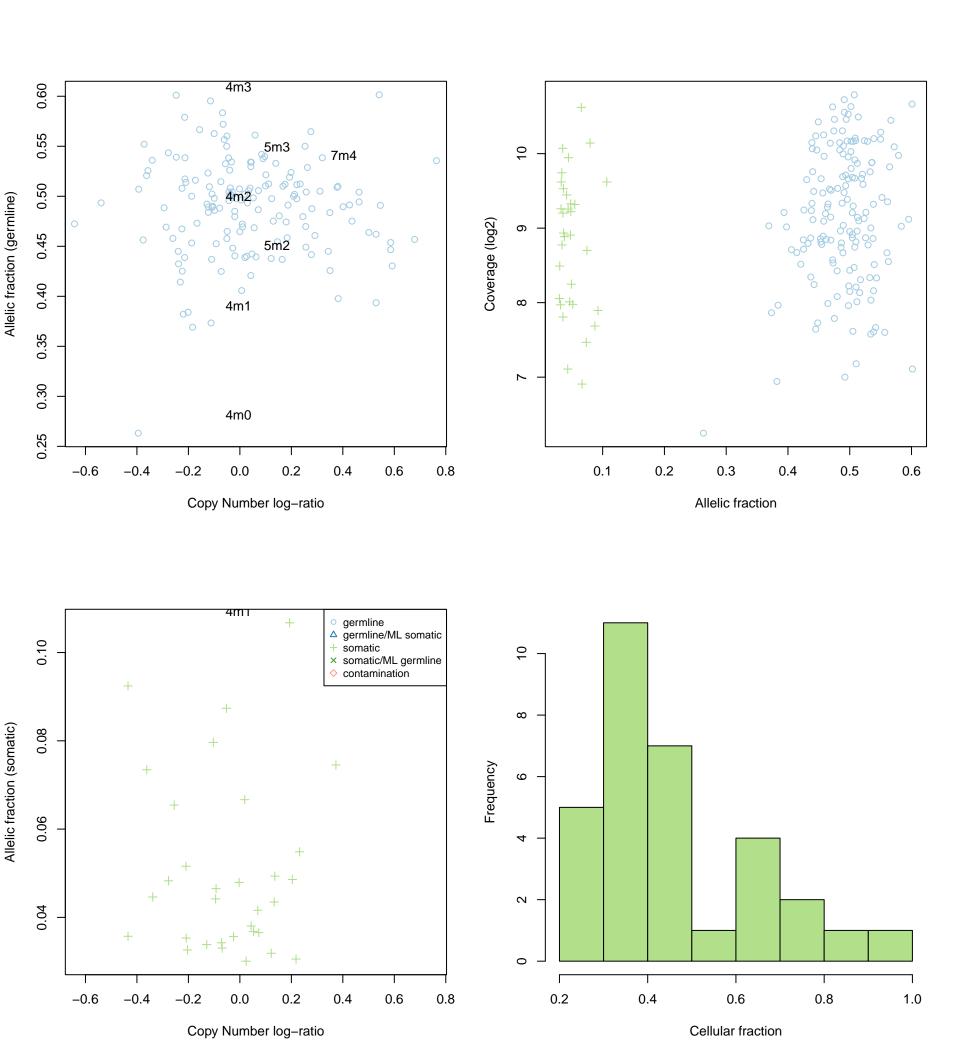




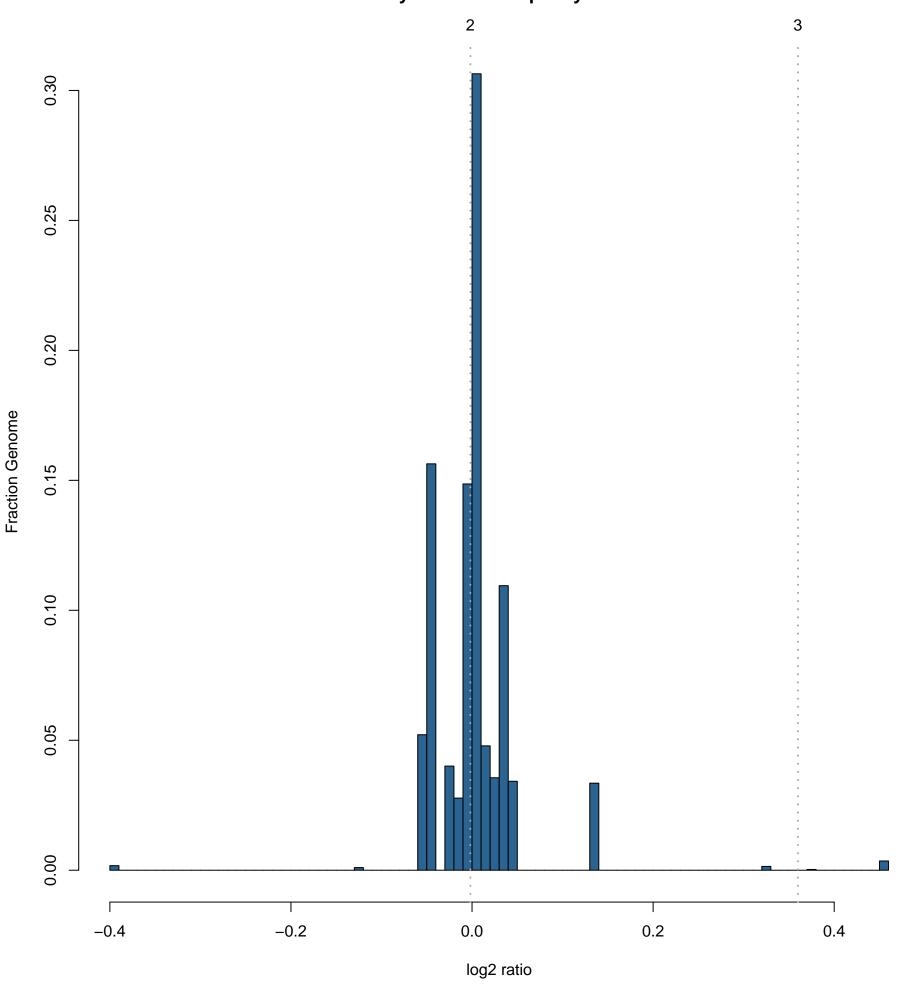
## SCNA-fit log-likelihood: -7368.51

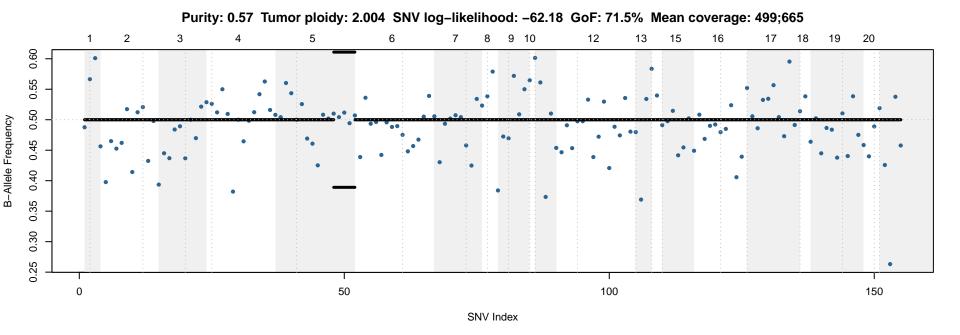




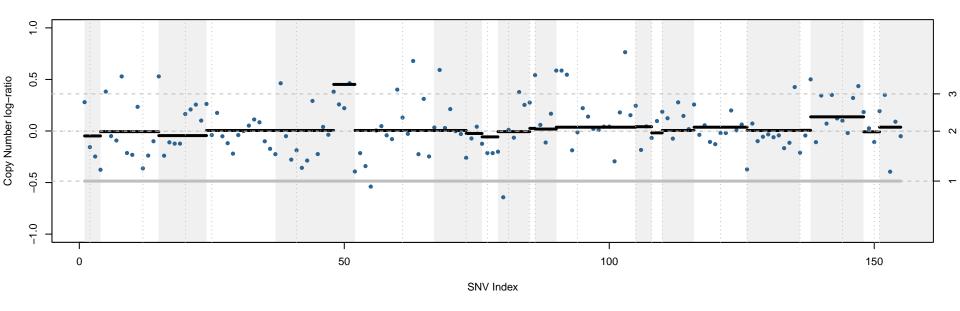


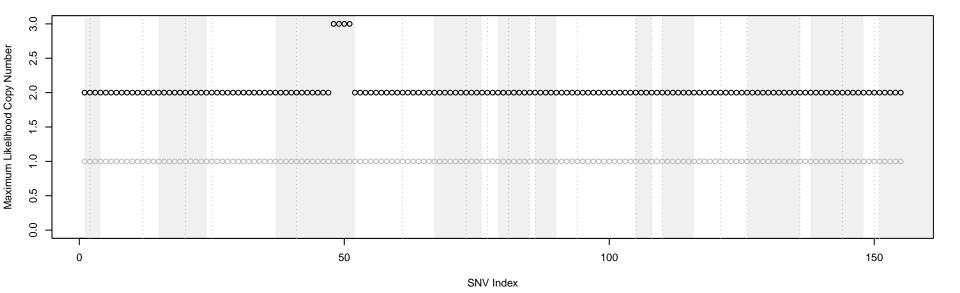
Purity: 0.57 Tumor ploidy: 2.004

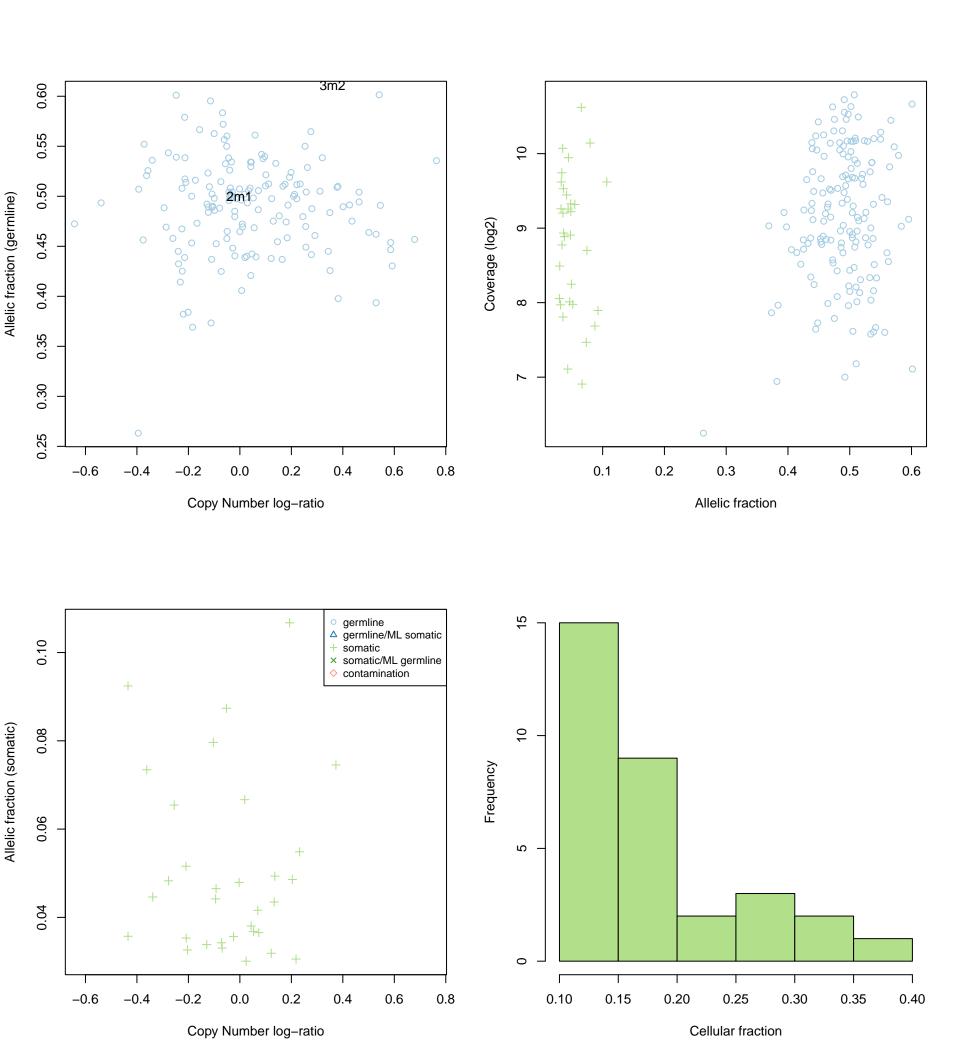




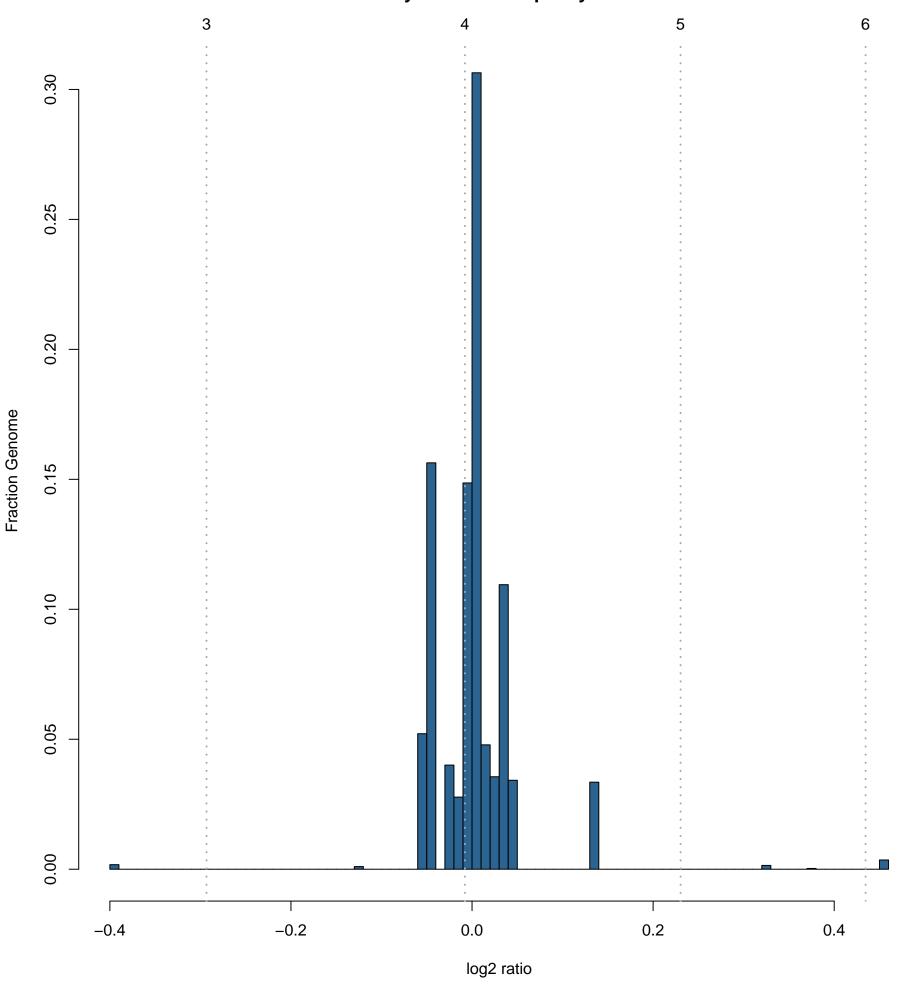
# SCNA-fit log-likelihood: -7425.86

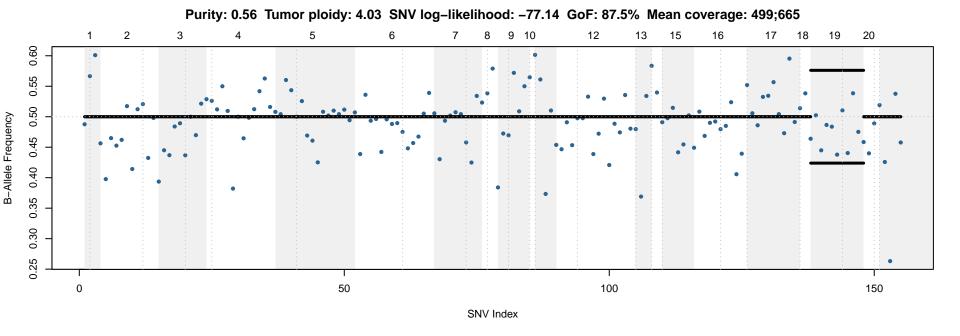




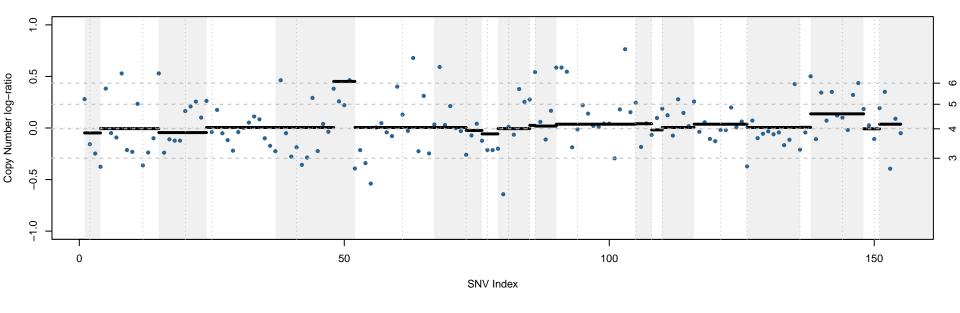


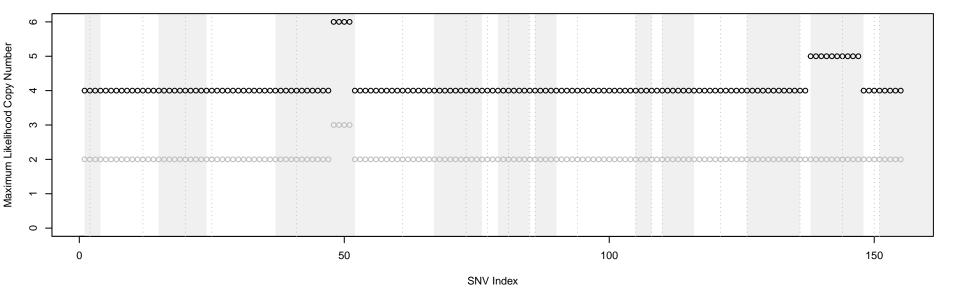
Purity: 0.56 Tumor ploidy: 4.03

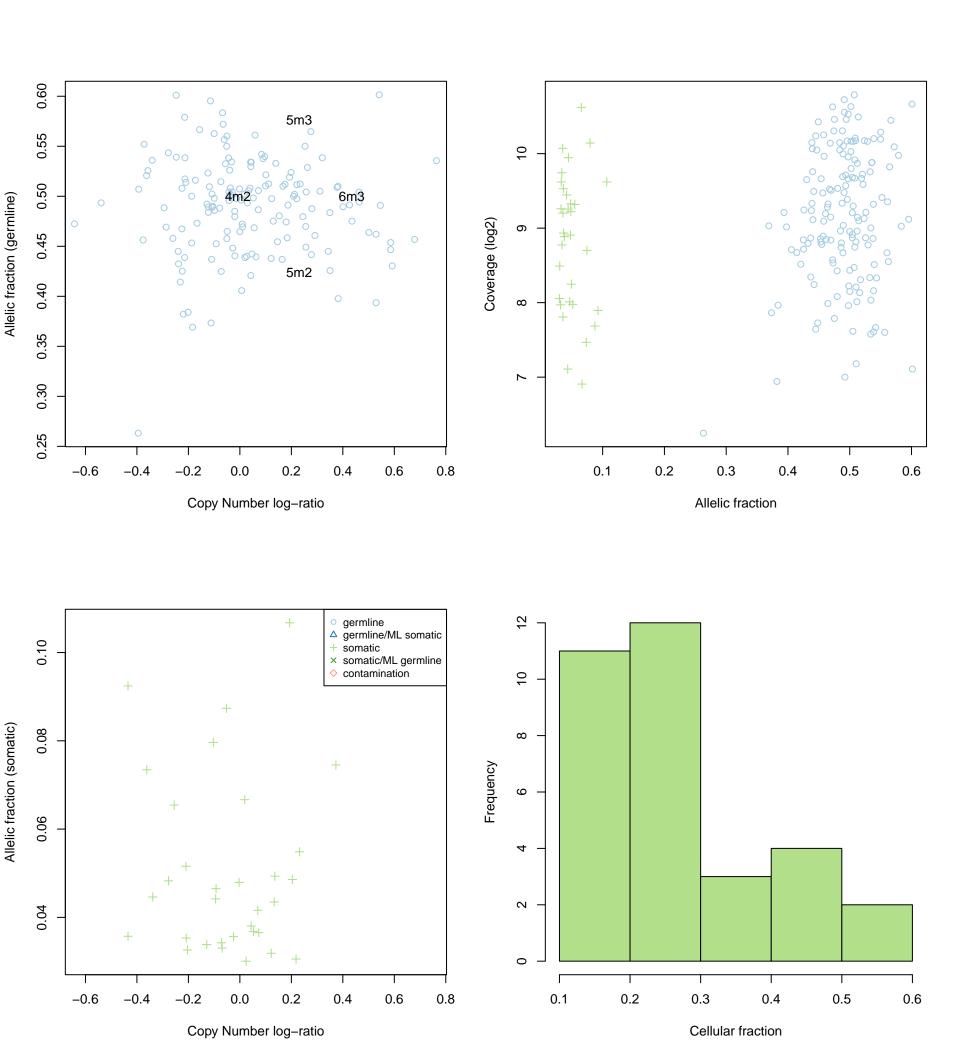




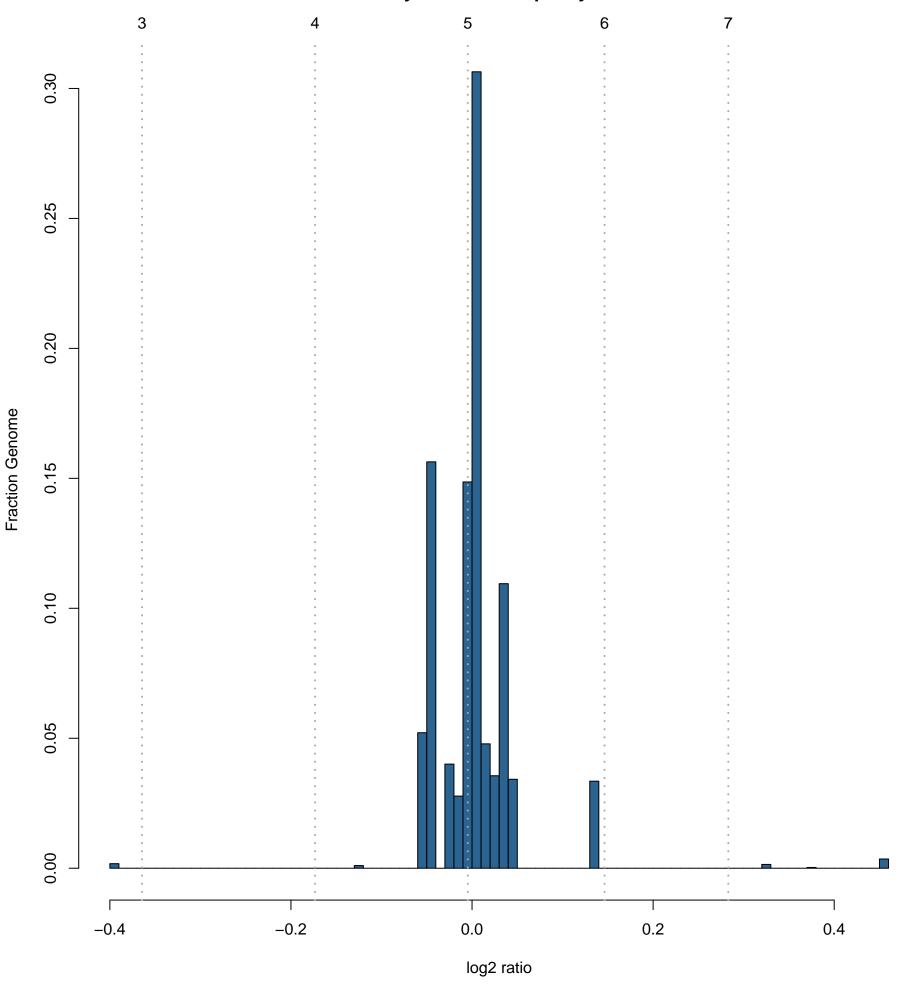
# SCNA-fit log-likelihood: -7397.1

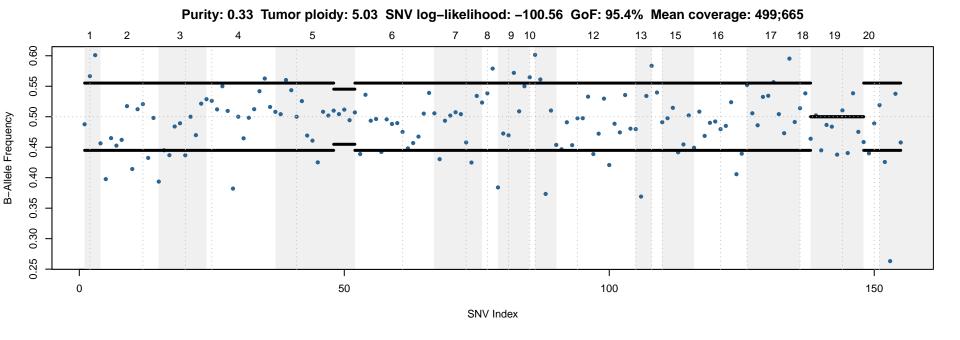




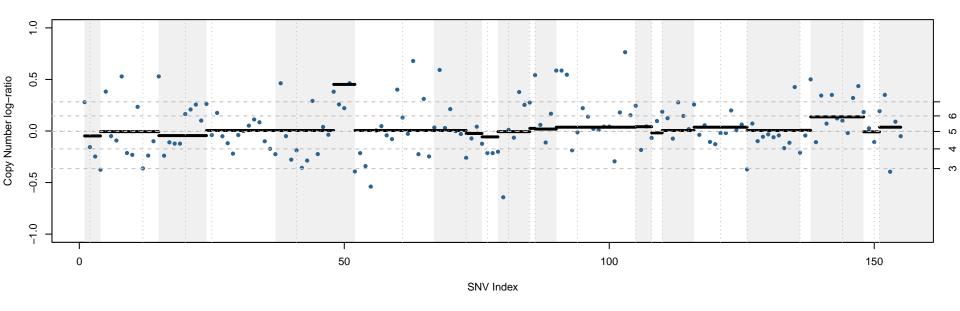


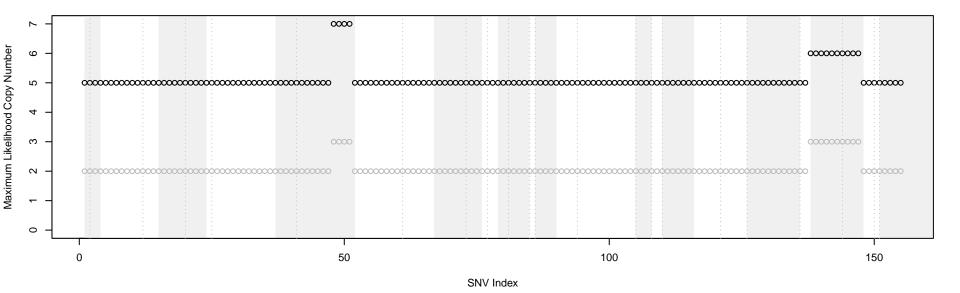
Purity: 0.33 Tumor ploidy: 5.03

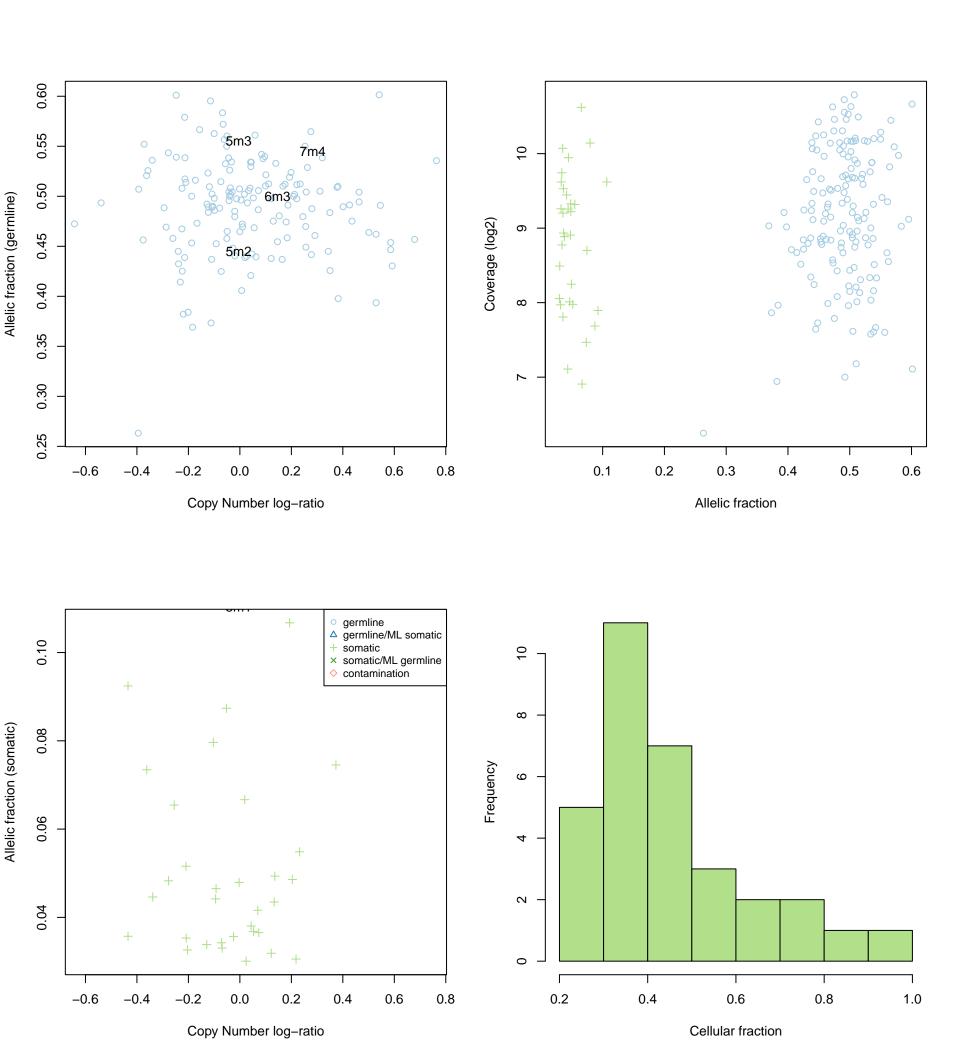




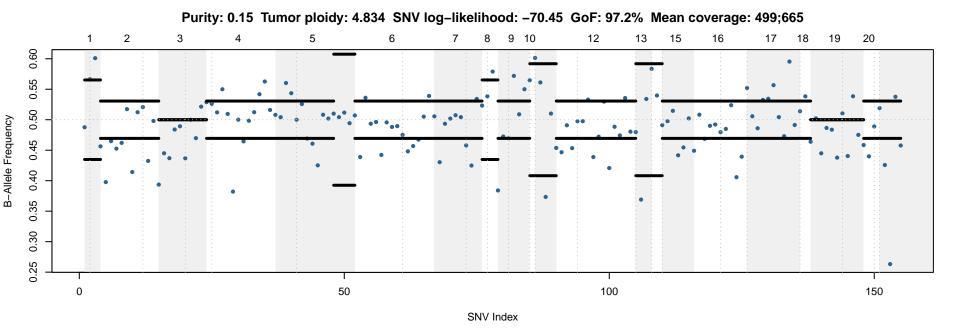
## SCNA-fit log-likelihood: -7376.87



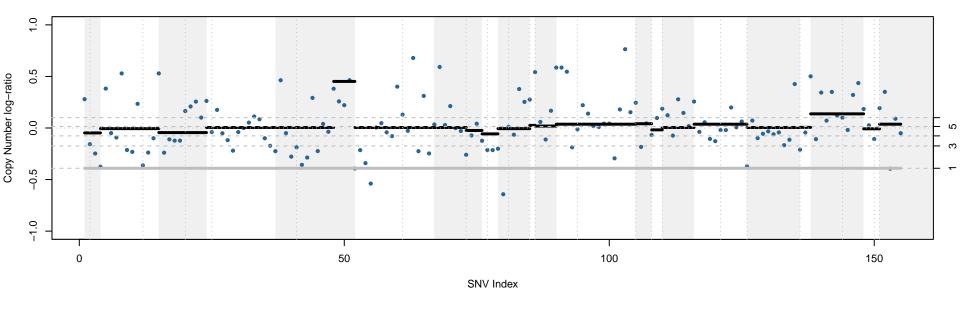


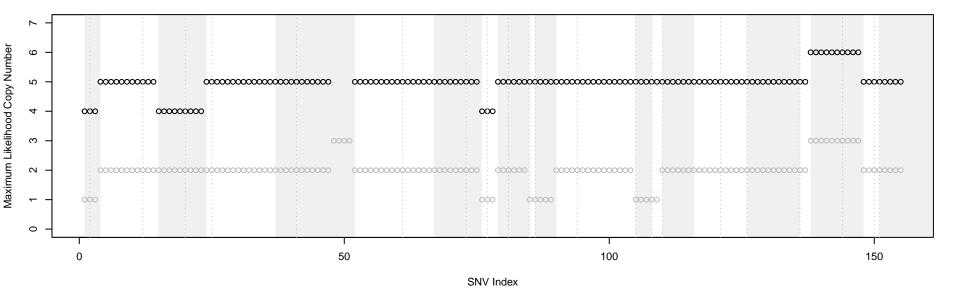


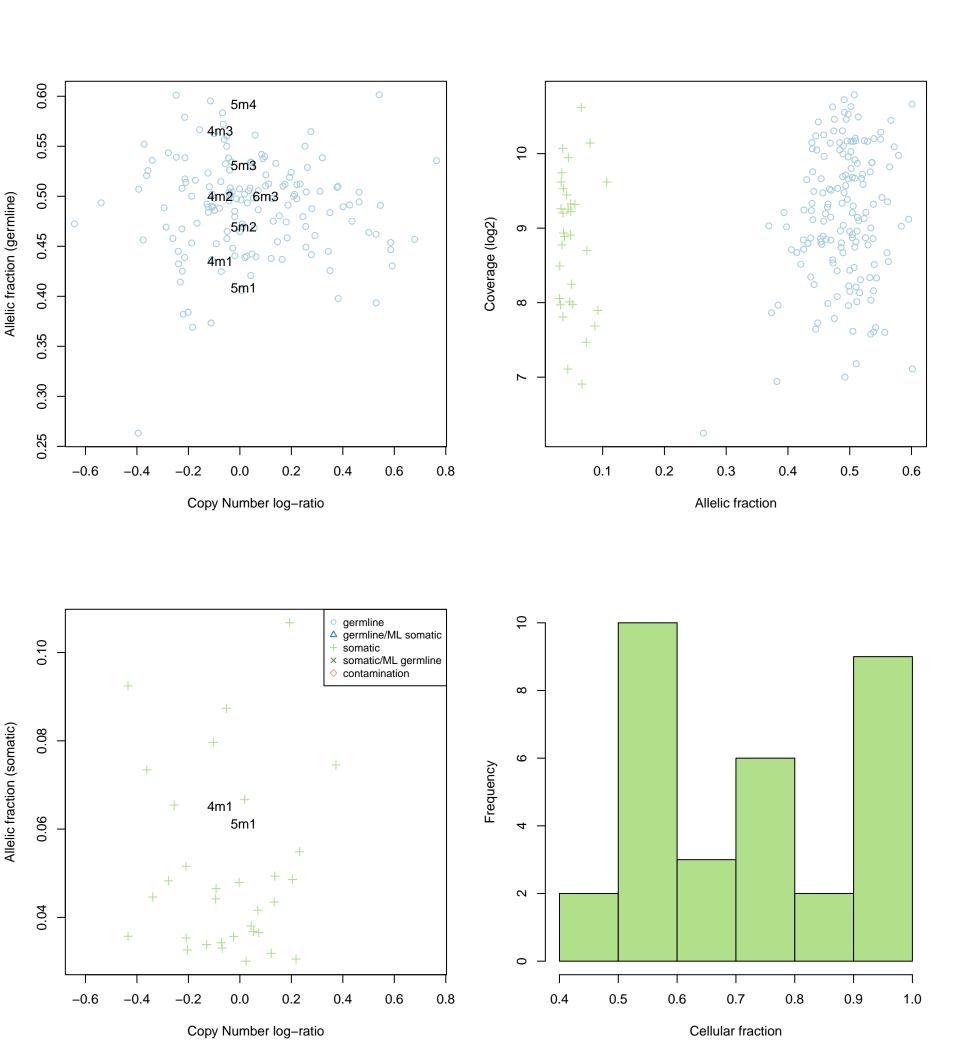
Purity: 0.15 Tumor ploidy: 4.834 5 3 1 0.30 0.25 0.20 Fraction Genome 0.15 0.10 0.05 0.00 -0.4 -0.2 0.0 0.2 0.4 log2 ratio



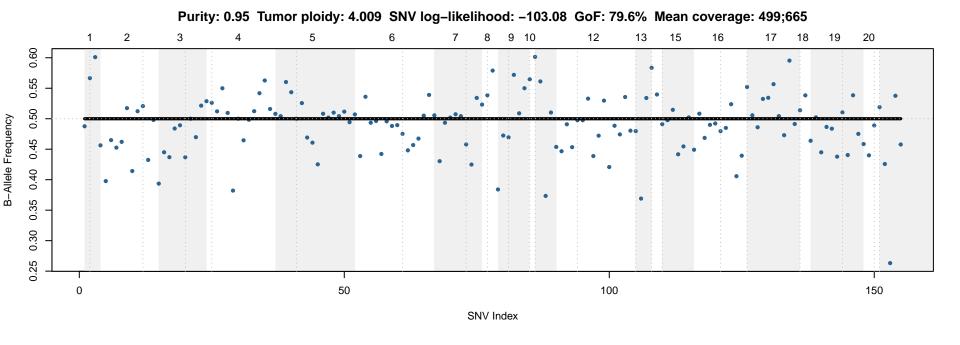
## SCNA-fit log-likelihood: -7464.66



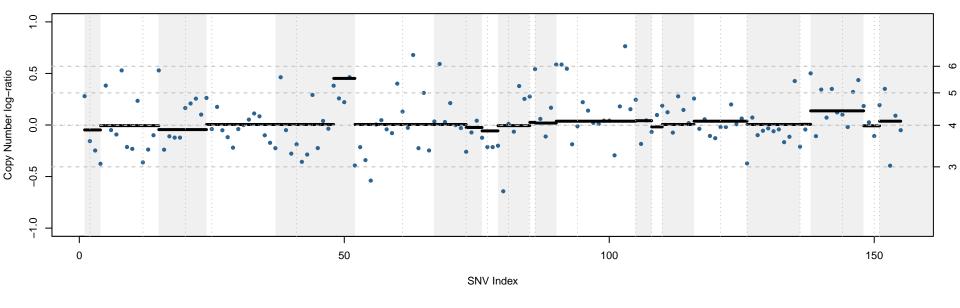


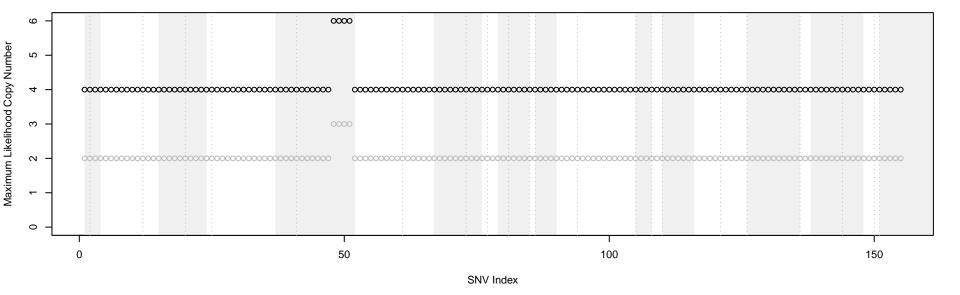


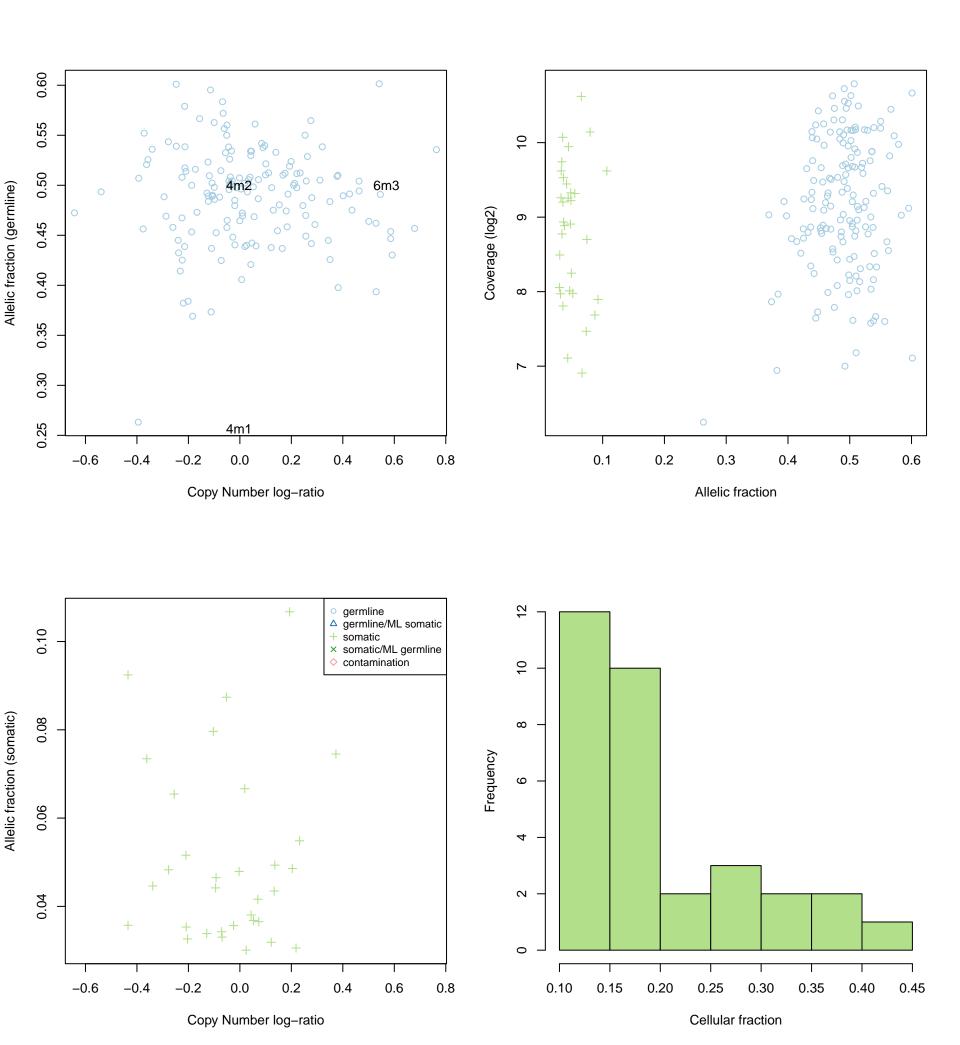
Purity: 0.95 Tumor ploidy: 4.009 3 5 0.30 0.25 0.20 Fraction Genome 0.15 0.10 0.05 0.00 -0.4 -0.2 0.0 0.2 0.4 log2 ratio



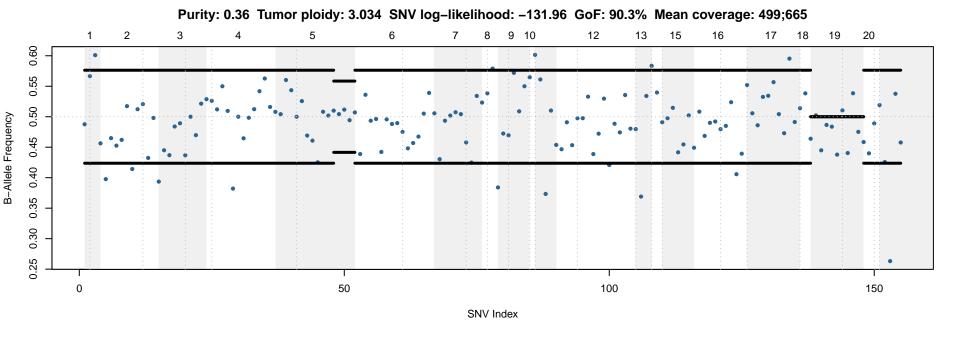
# SCNA-fit log-likelihood: -7426.15



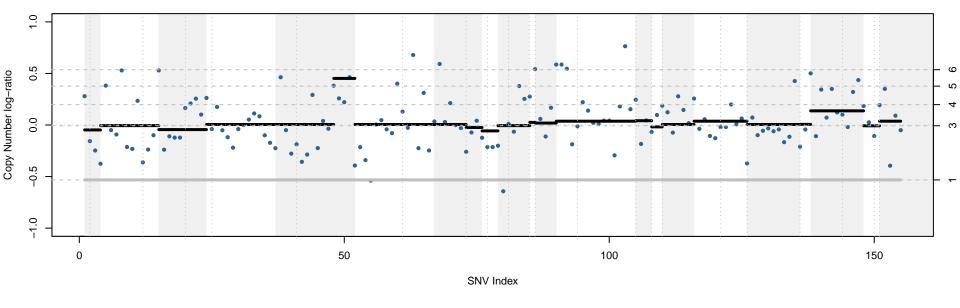


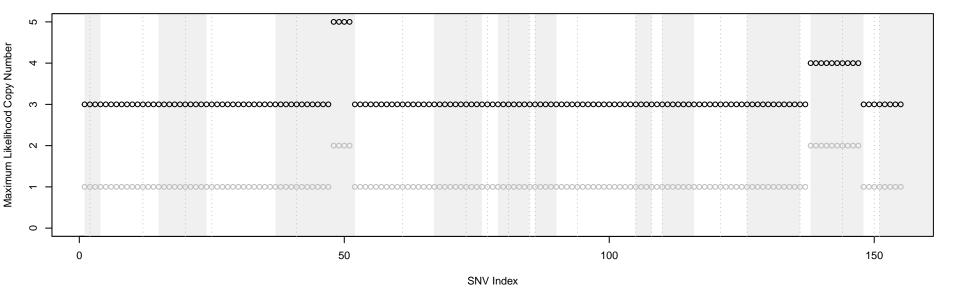


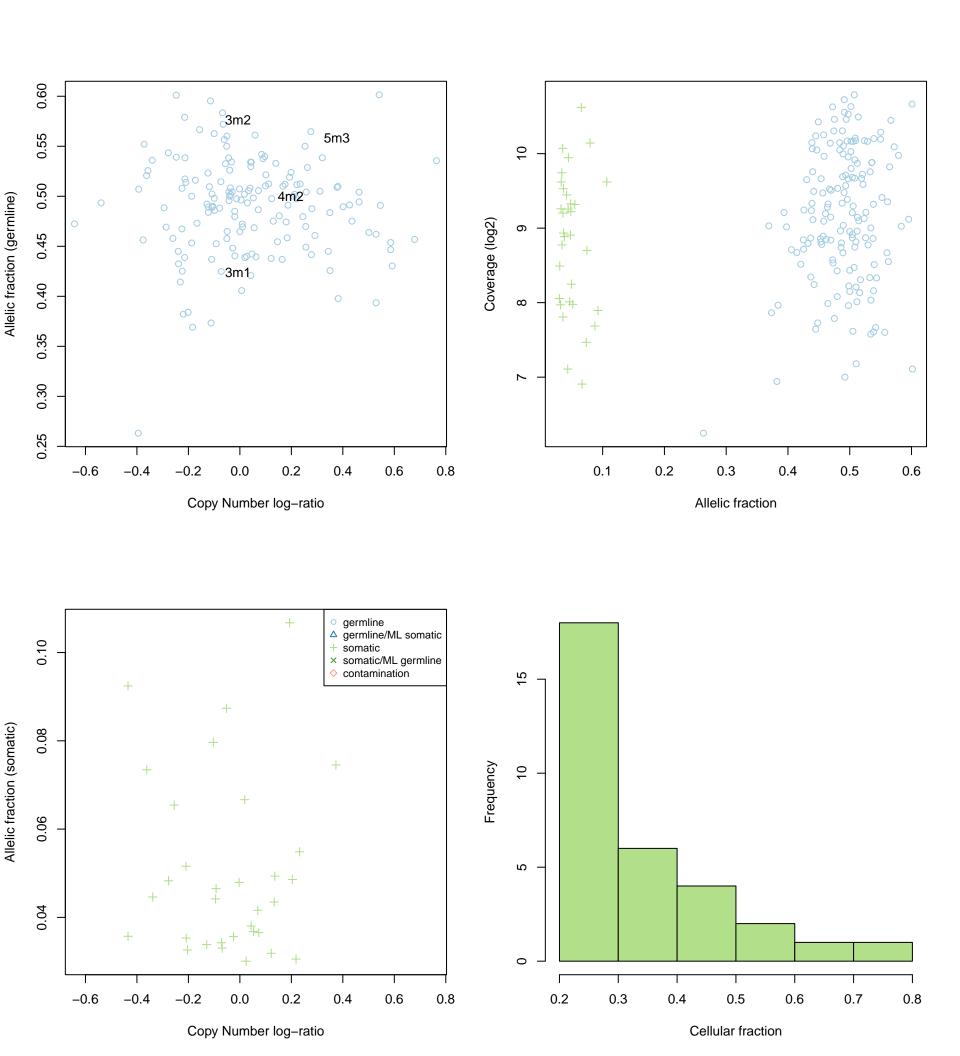
Purity: 0.36 Tumor ploidy: 3.034 5 3 0.30 0.25 0.20 Fraction Genome 0.15 0.05 0.00 -0.4 -0.2 0.0 0.2 0.4 log2 ratio



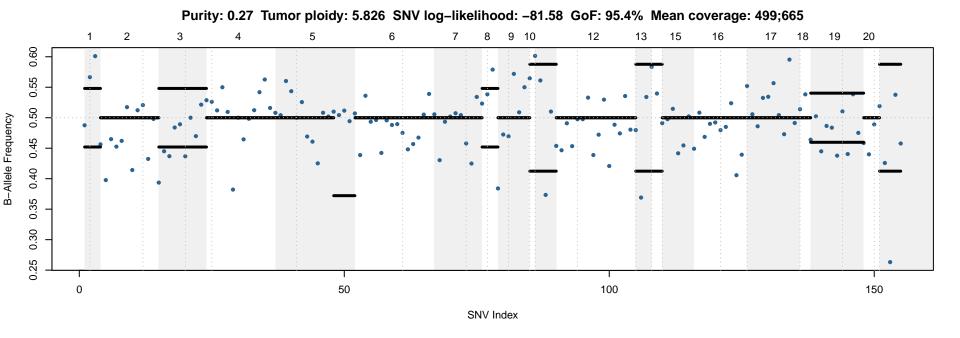
## SCNA-fit log-likelihood: -7383.72



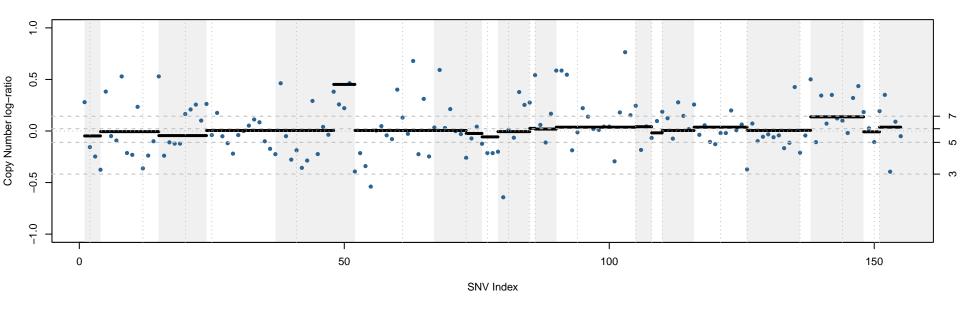


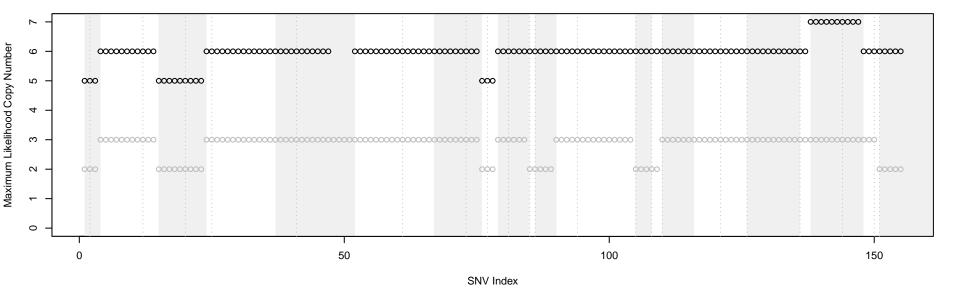


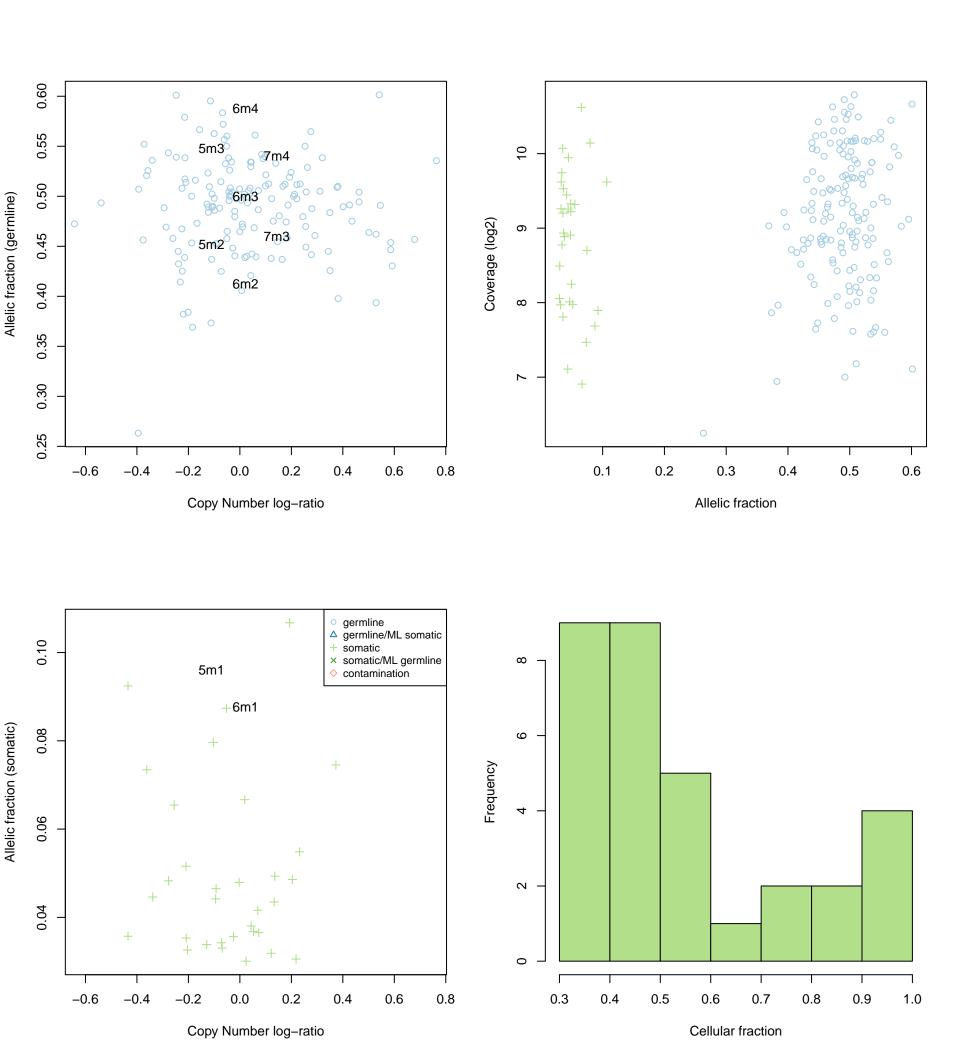
Purity: 0.27 Tumor ploidy: 5.826 3 0.30 0.25 0.20 Fraction Genome 0.15 0.10 0.05 0.00 -0.4 -0.2 0.0 0.2 0.4 log2 ratio



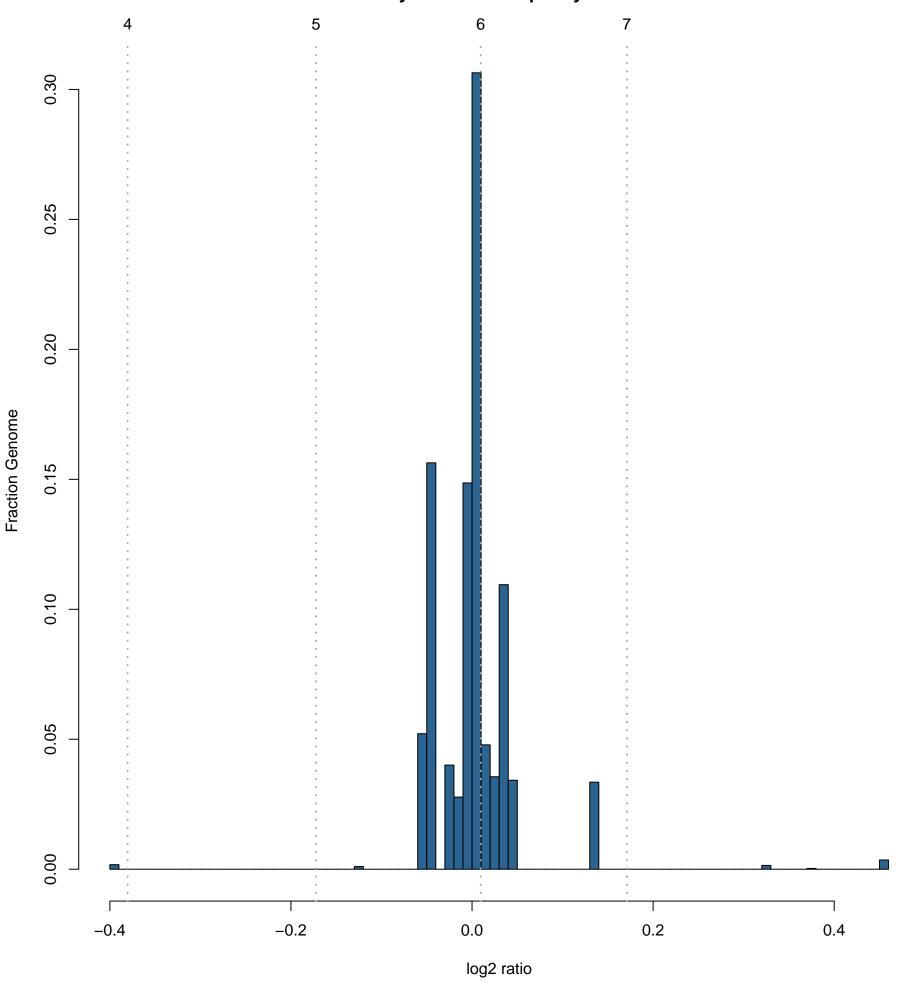
## SCNA-fit log-likelihood: -7522.27

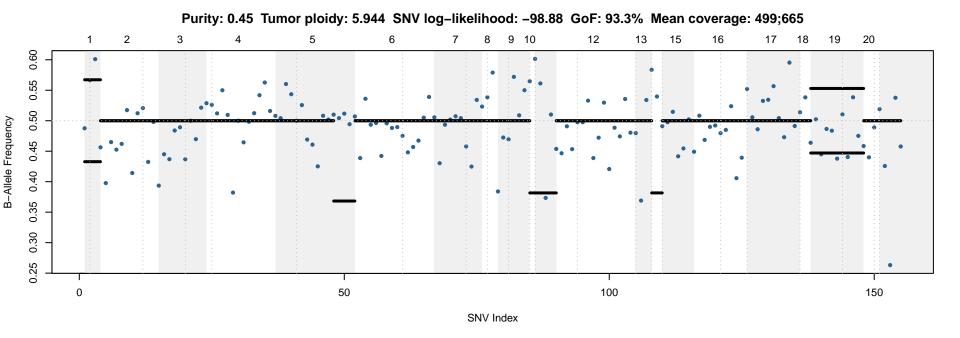




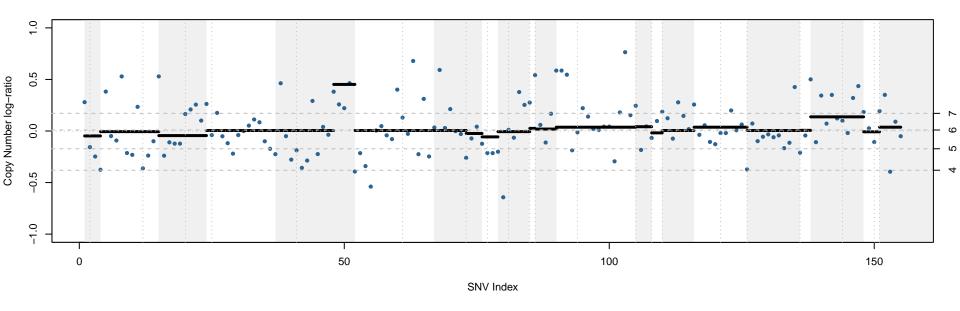


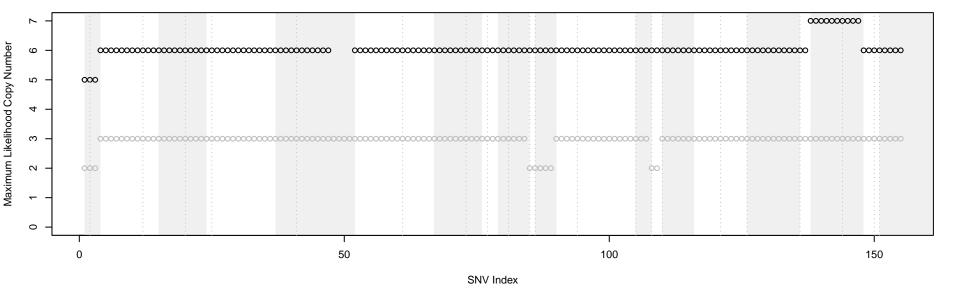
Purity: 0.45 Tumor ploidy: 5.944

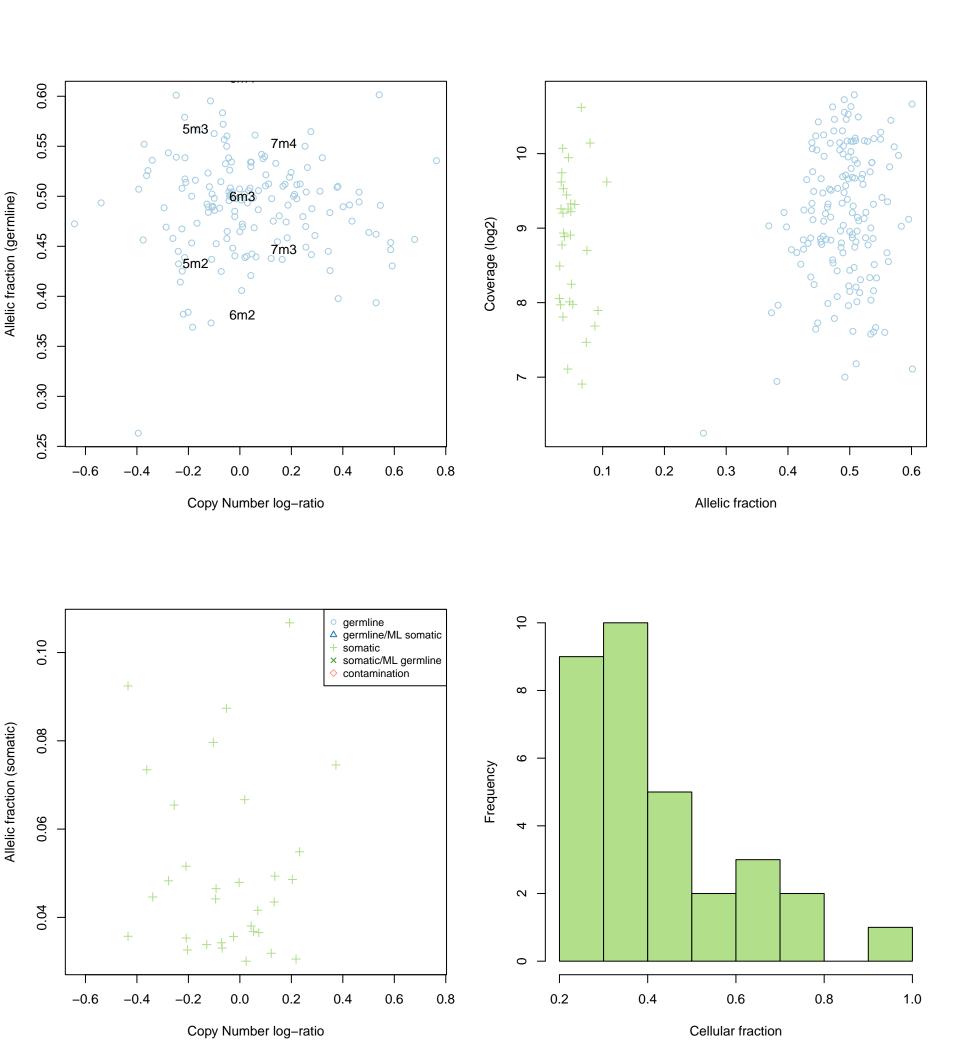




## SCNA-fit log-likelihood: -7551.44

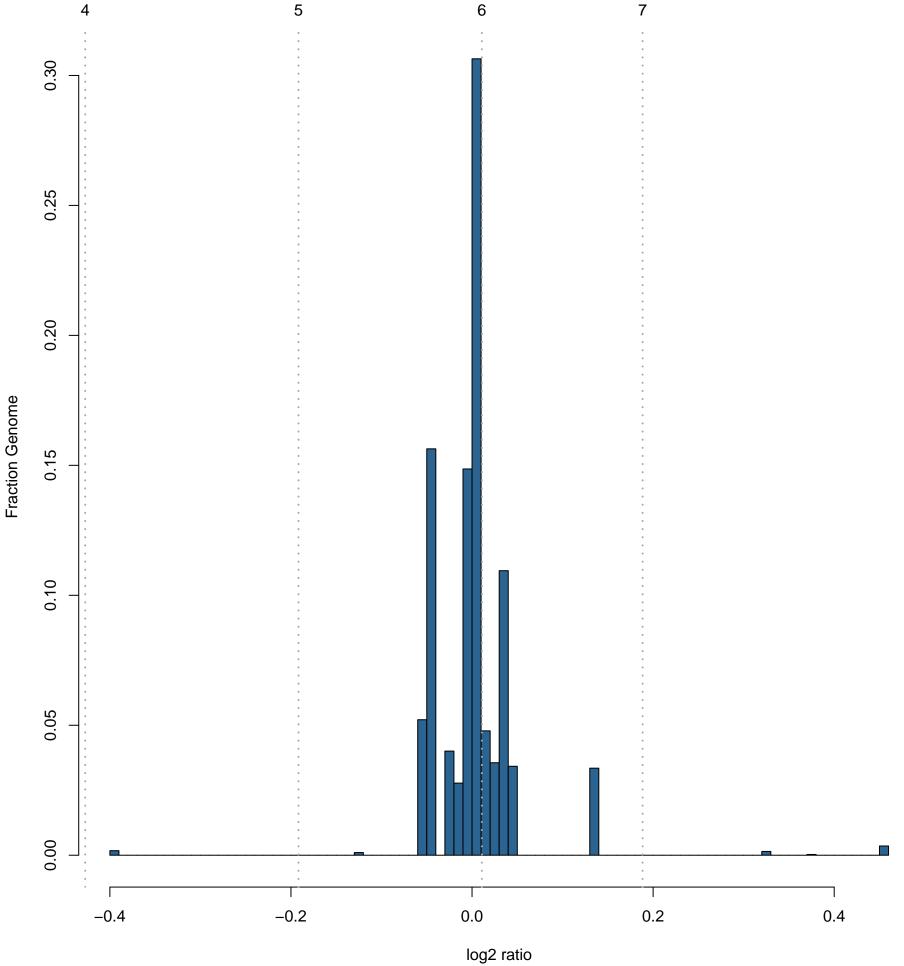


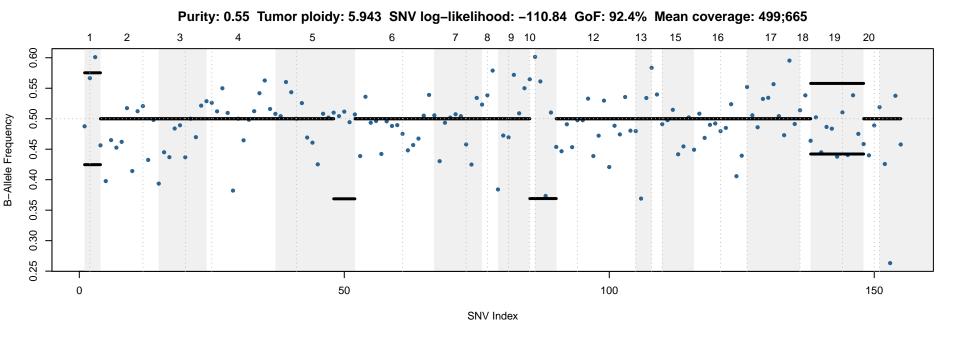




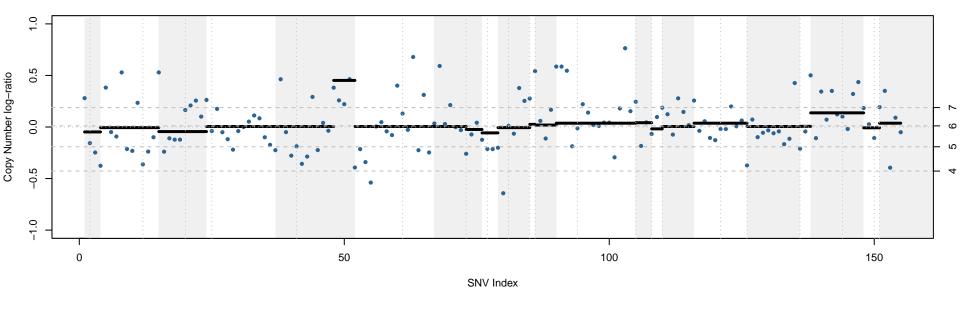
Purity: 0.55 Tumor ploidy: 5.943

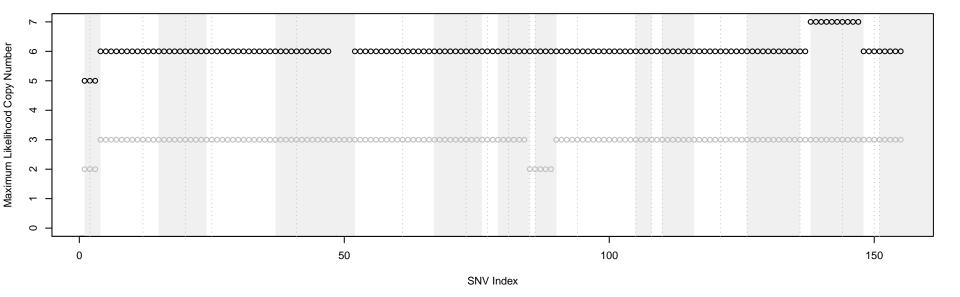
6 7

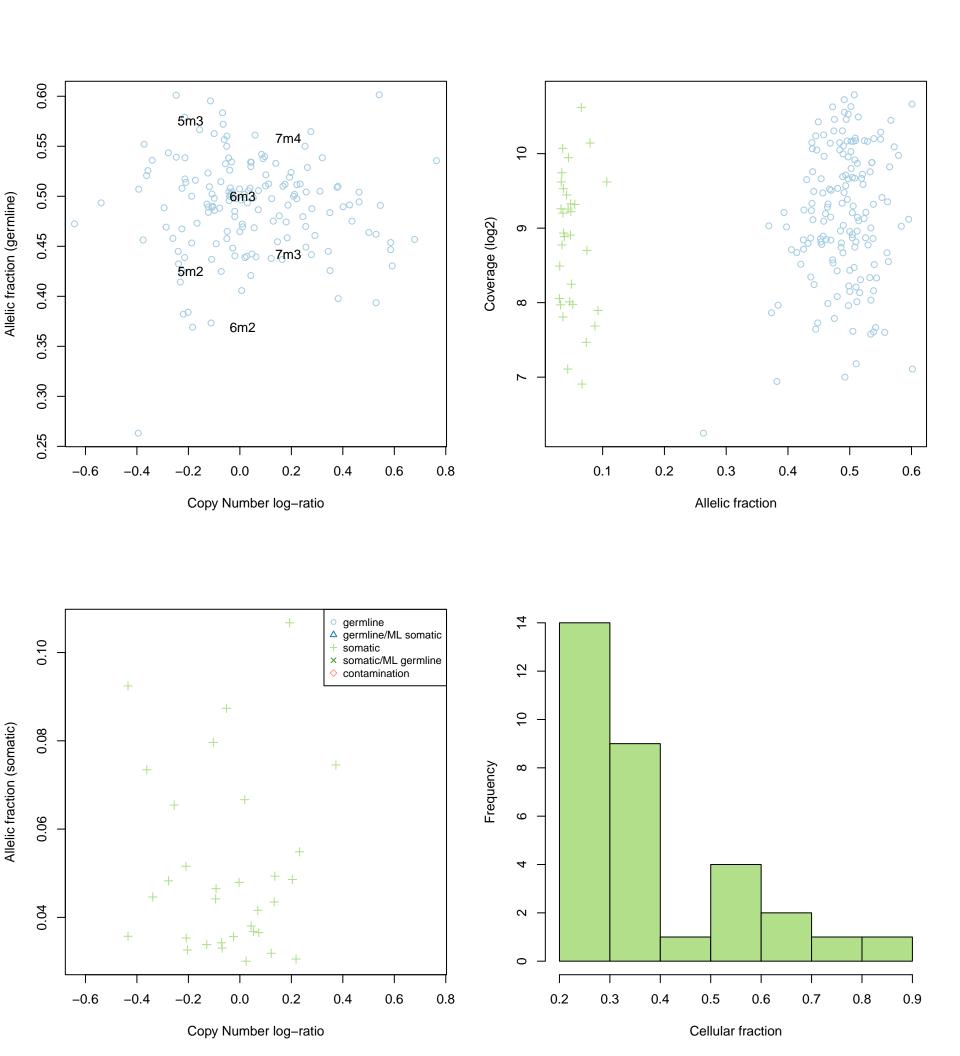




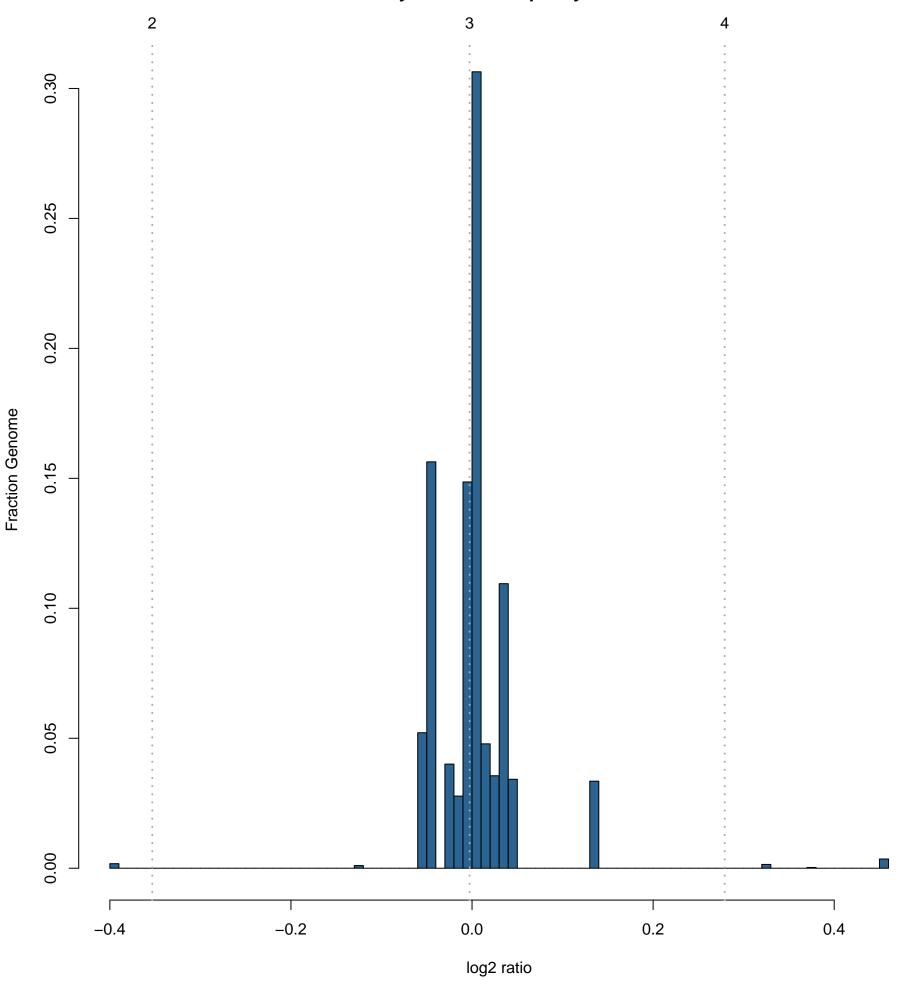
## SCNA-fit log-likelihood: -7592.55

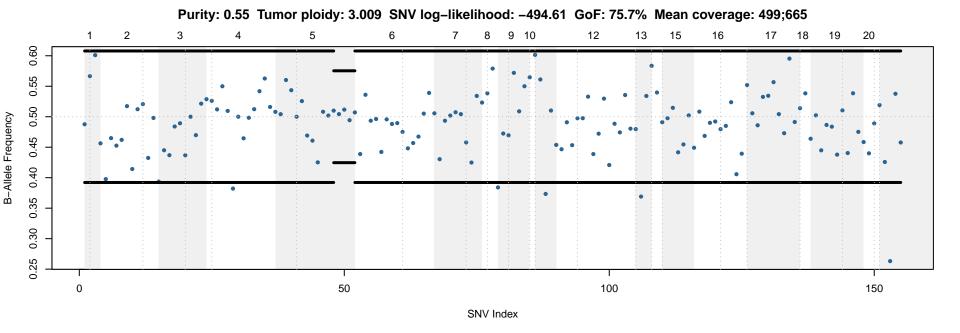




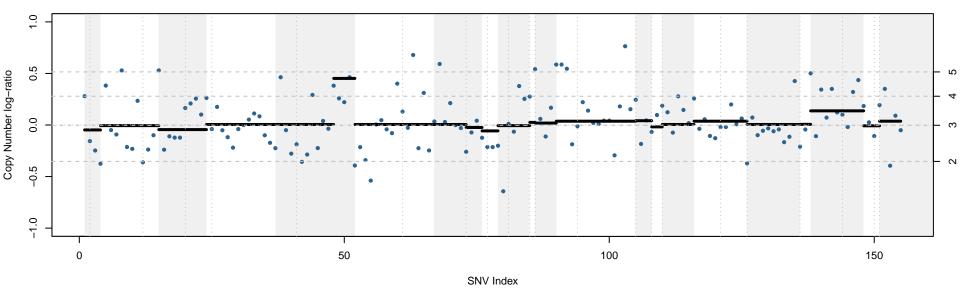


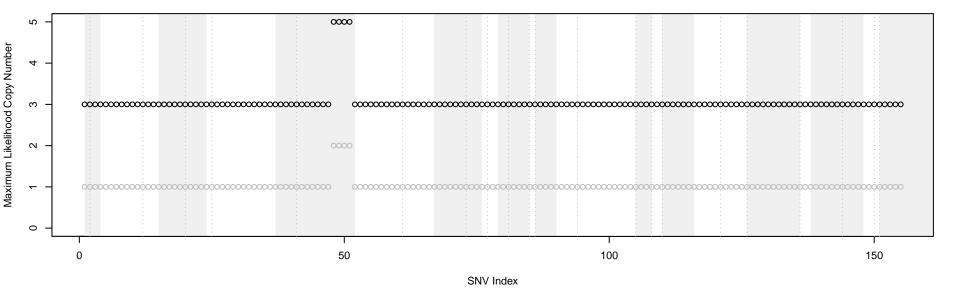
Purity: 0.55 Tumor ploidy: 3.009

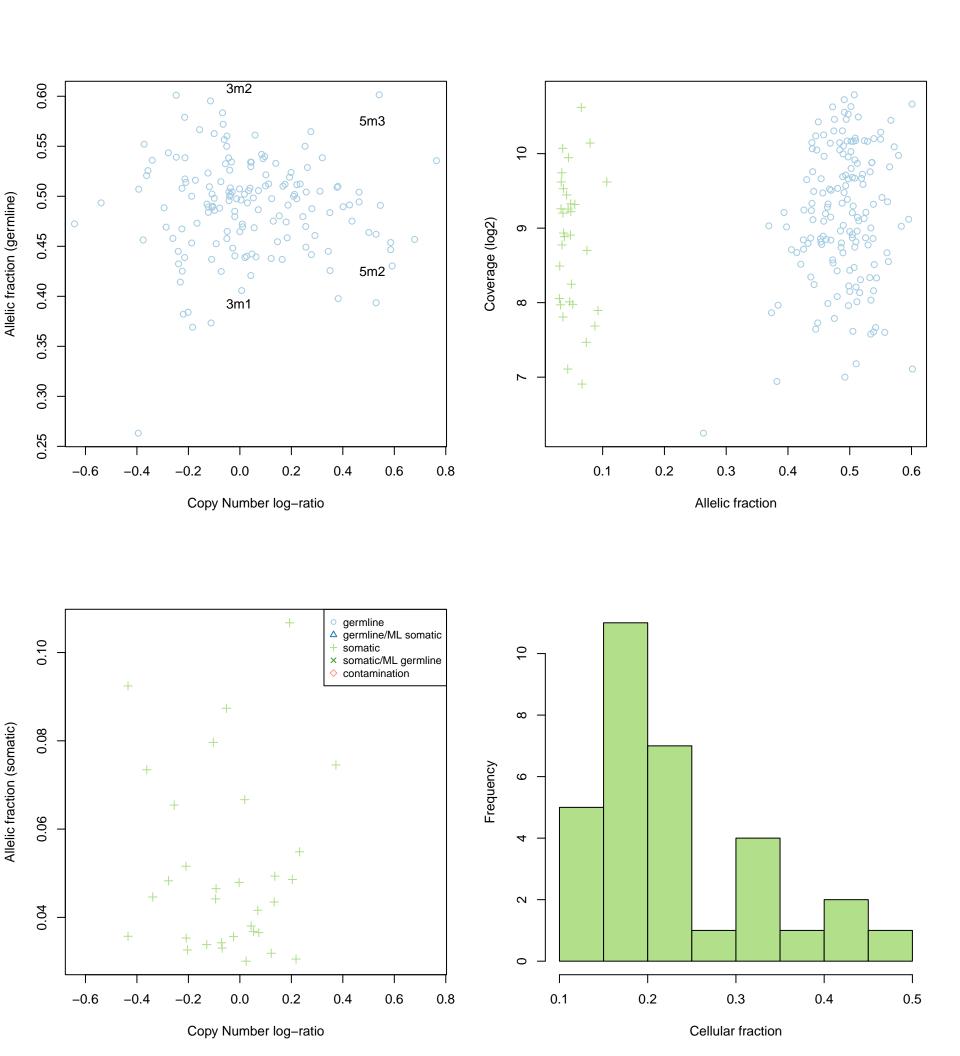




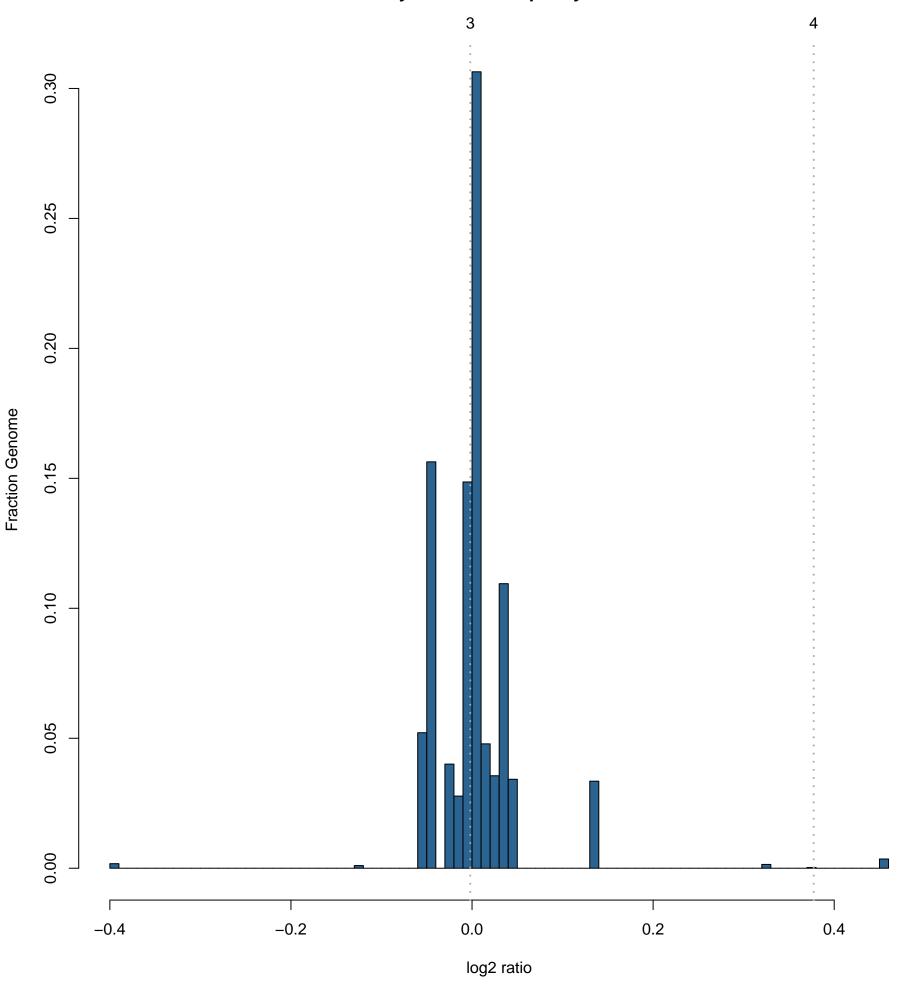
## SCNA-fit log-likelihood: -7424.13

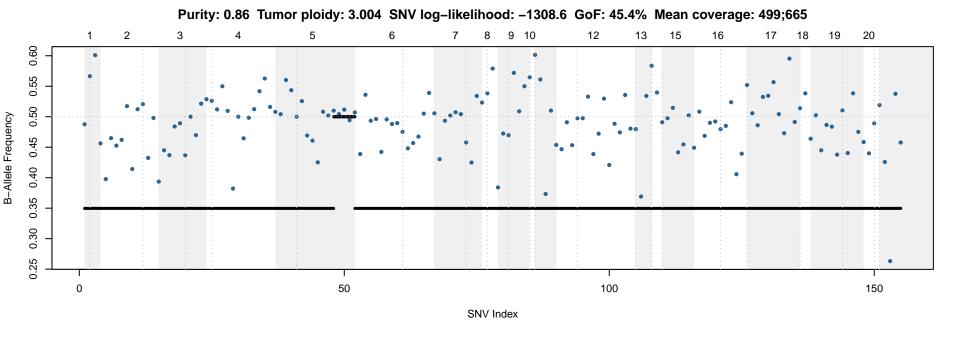






Purity: 0.86 Tumor ploidy: 3.004





## SCNA-fit log-likelihood: -7425.96

