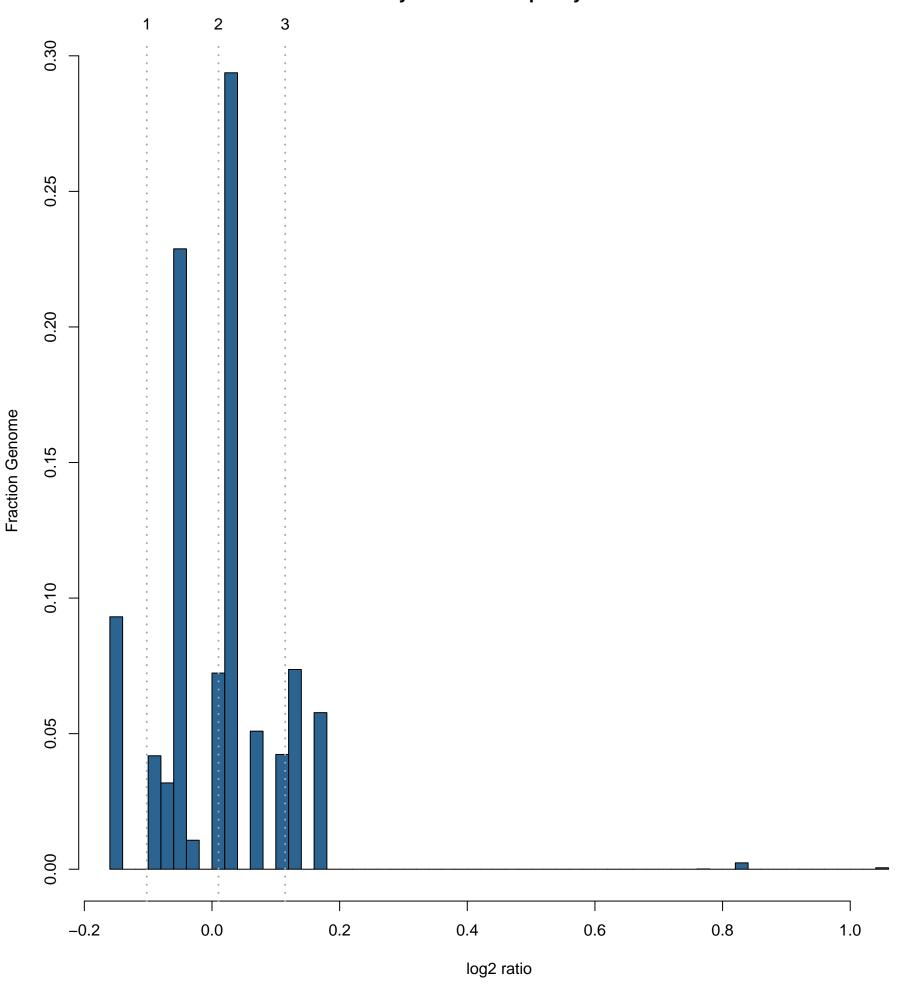
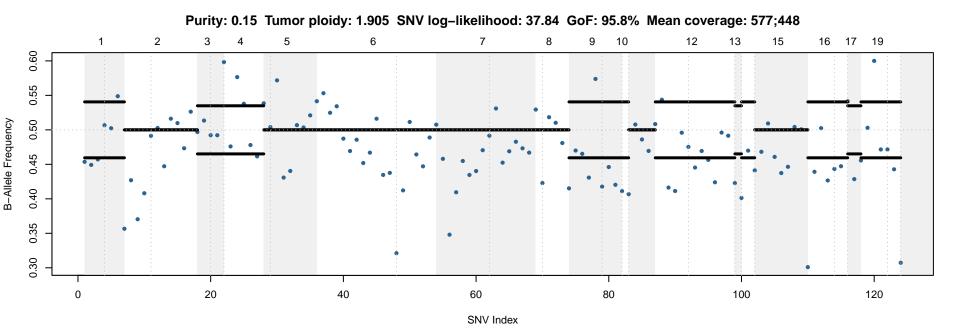
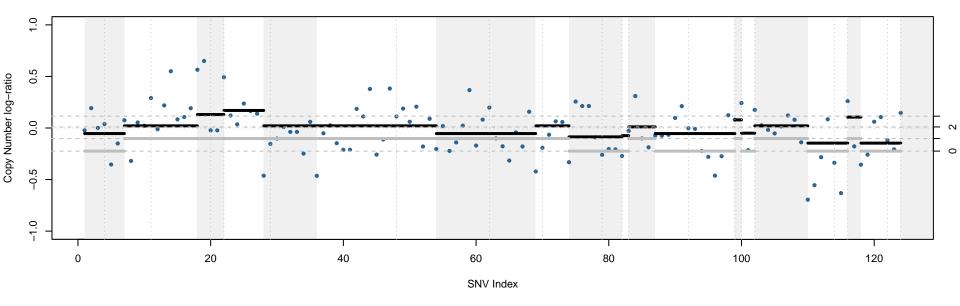
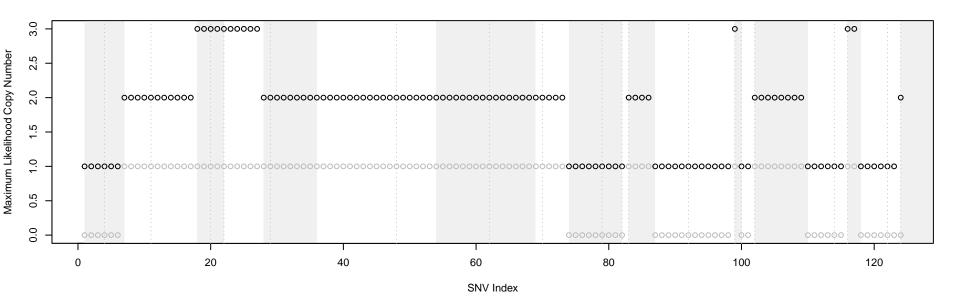
Purity: 0.15 Tumor ploidy: 1.905

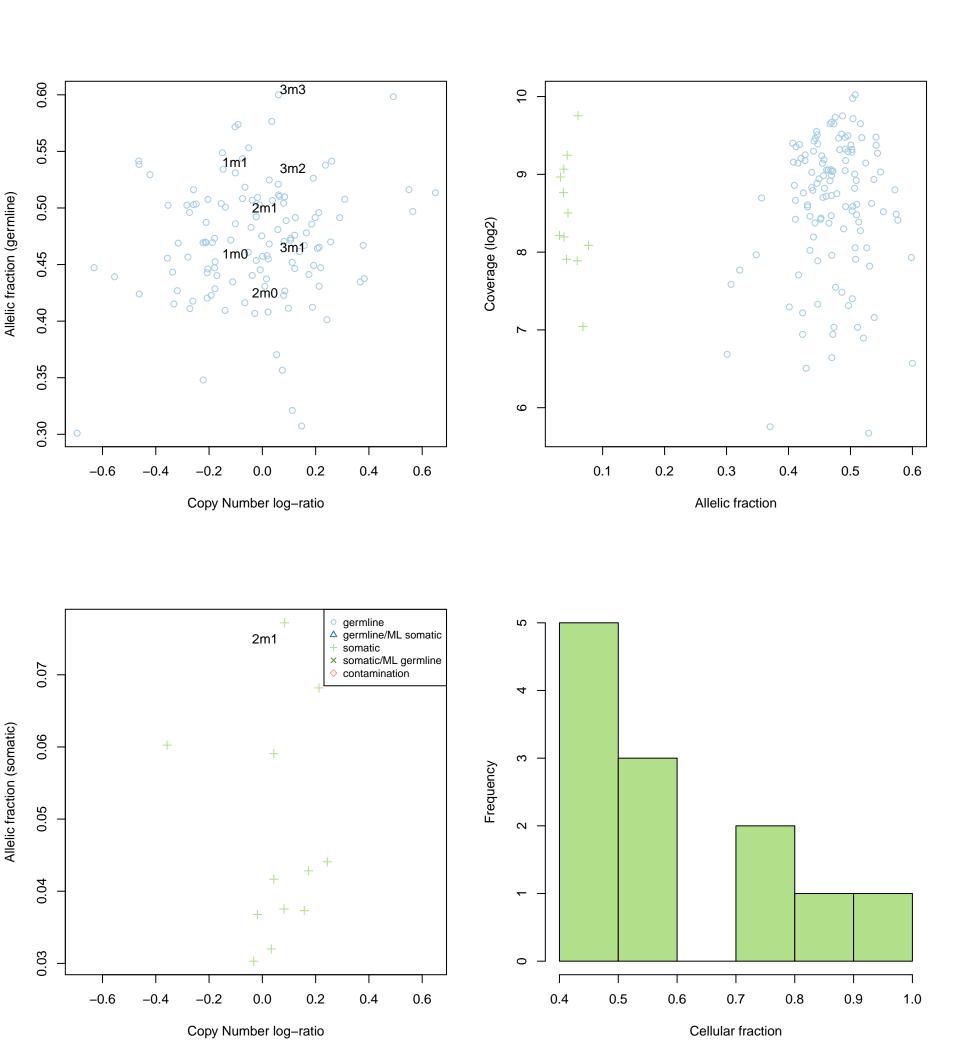




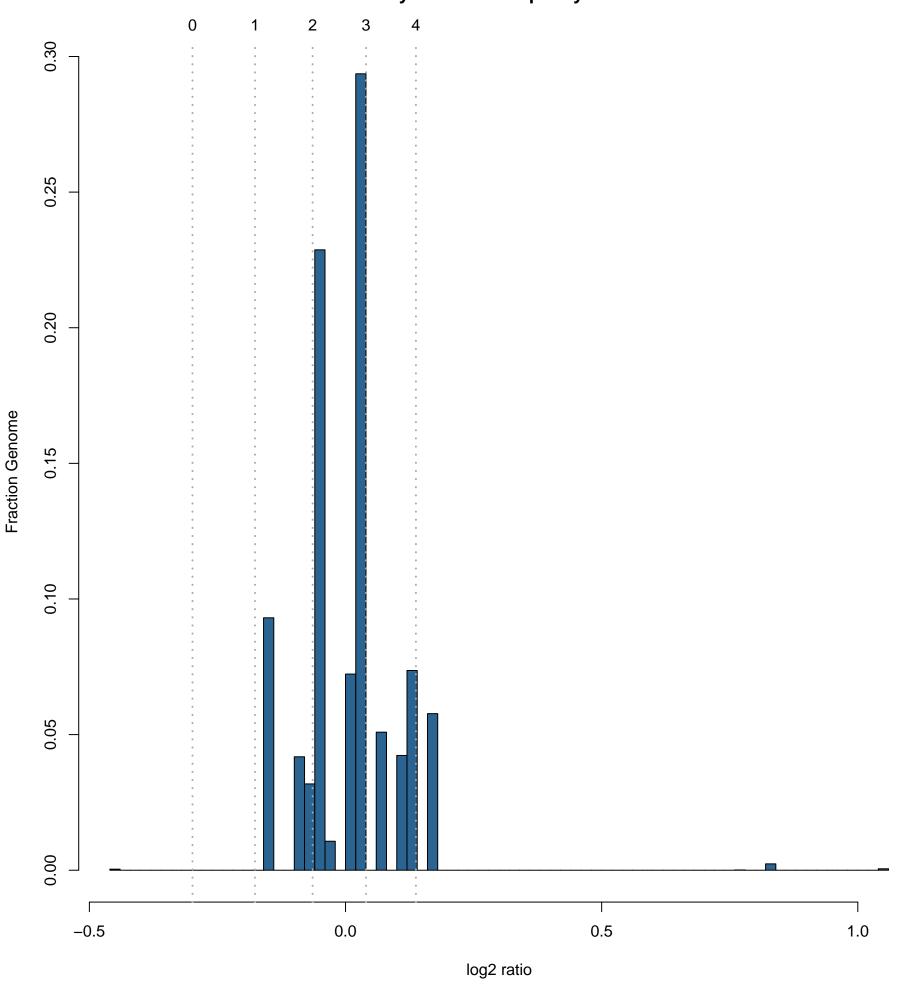
SCNA-fit log-likelihood: -14517.25

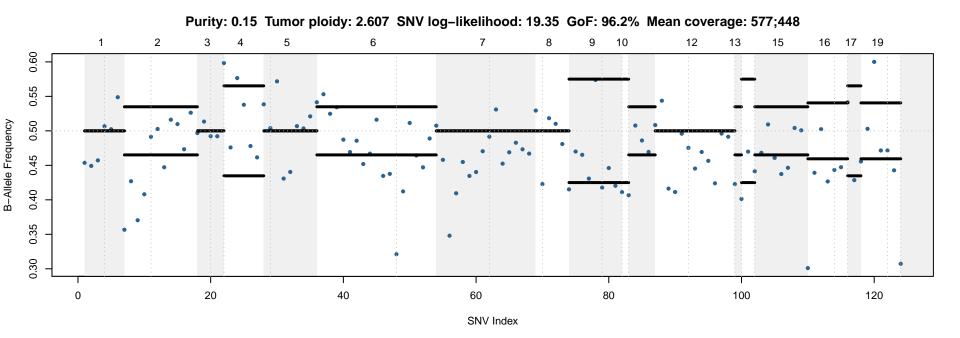




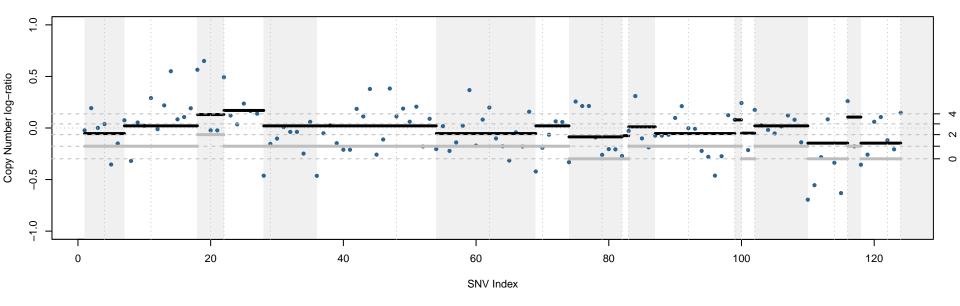


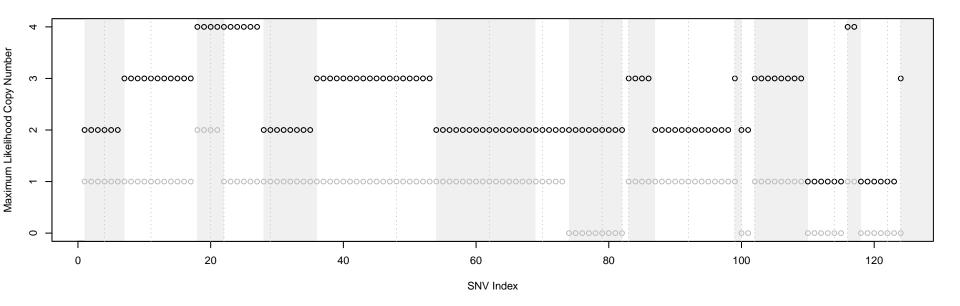
Purity: 0.15 Tumor ploidy: 2.607

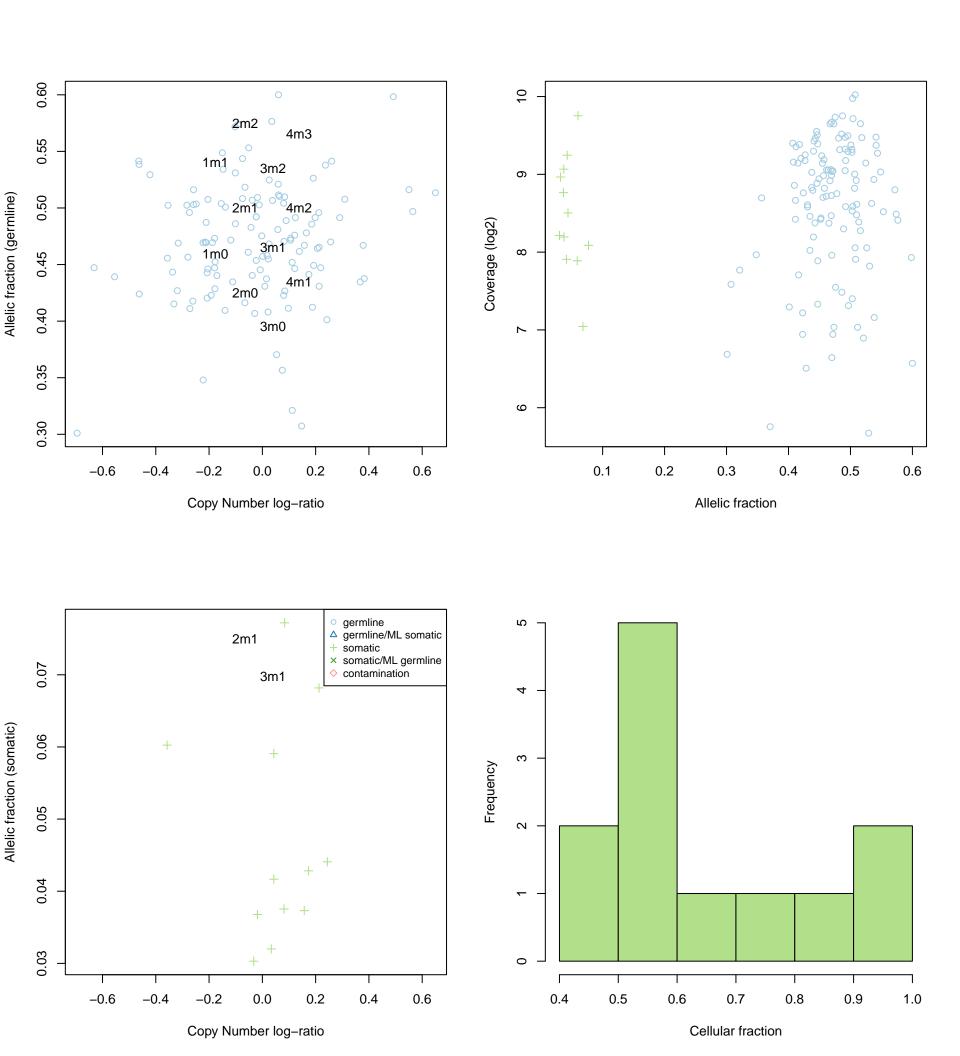




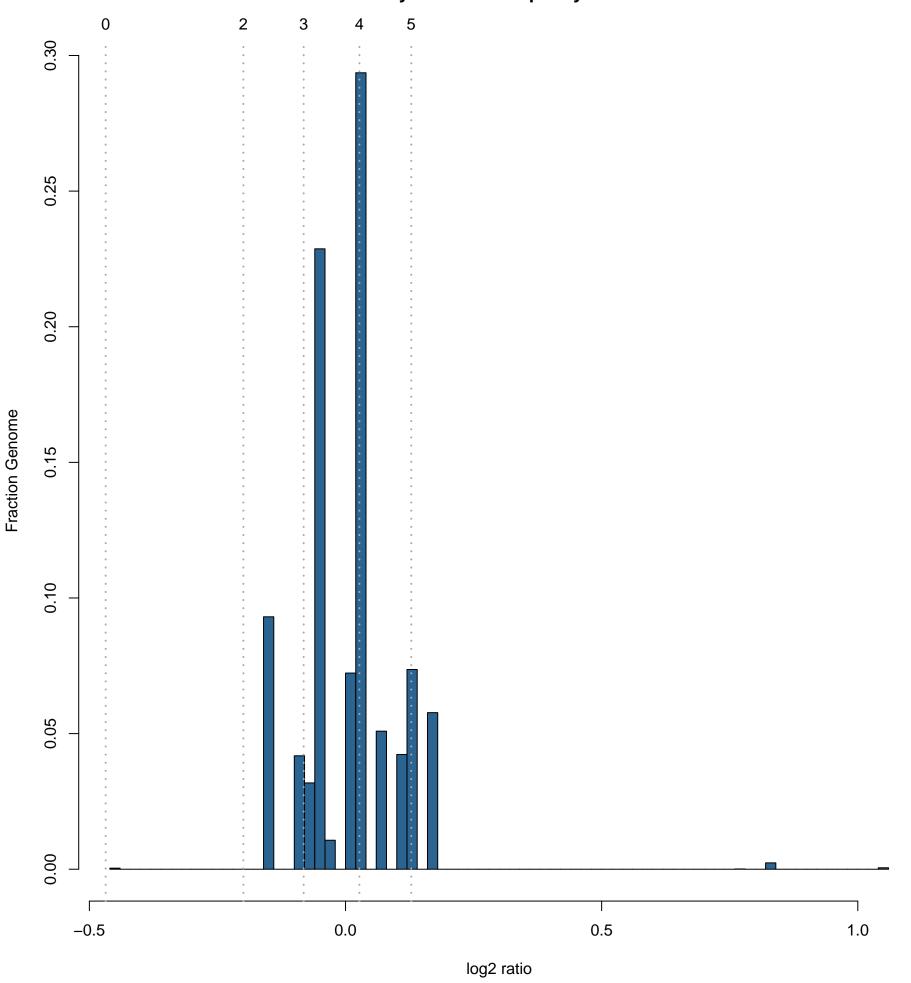
## SCNA-fit log-likelihood: -14501.14

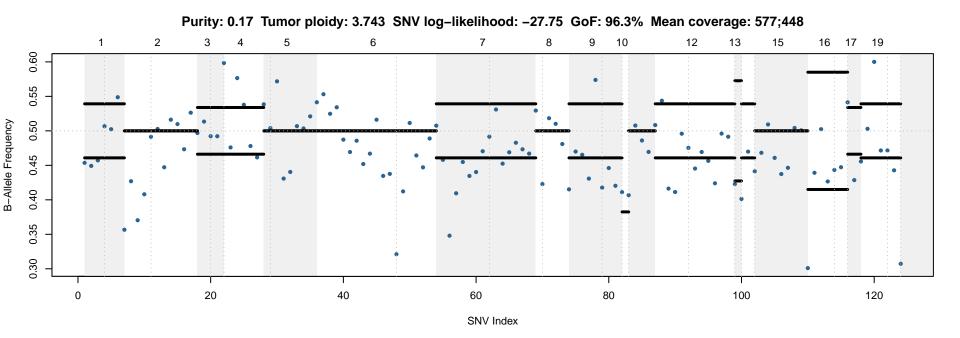




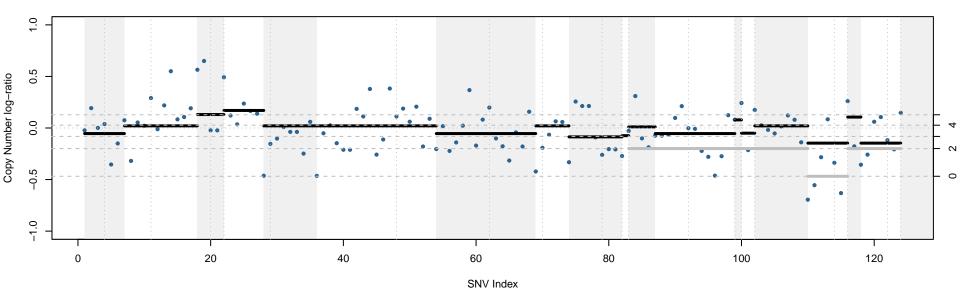


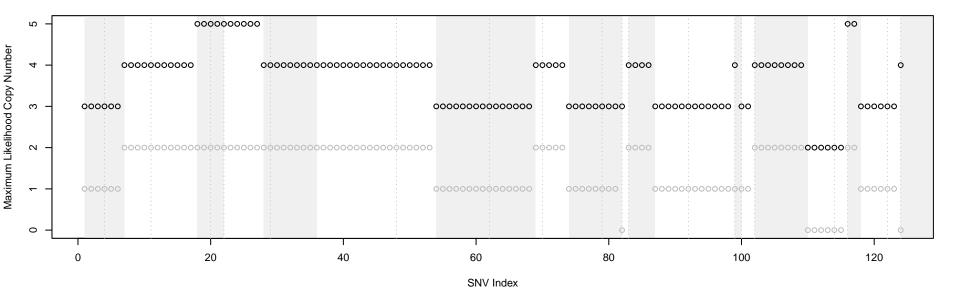
Purity: 0.17 Tumor ploidy: 3.743

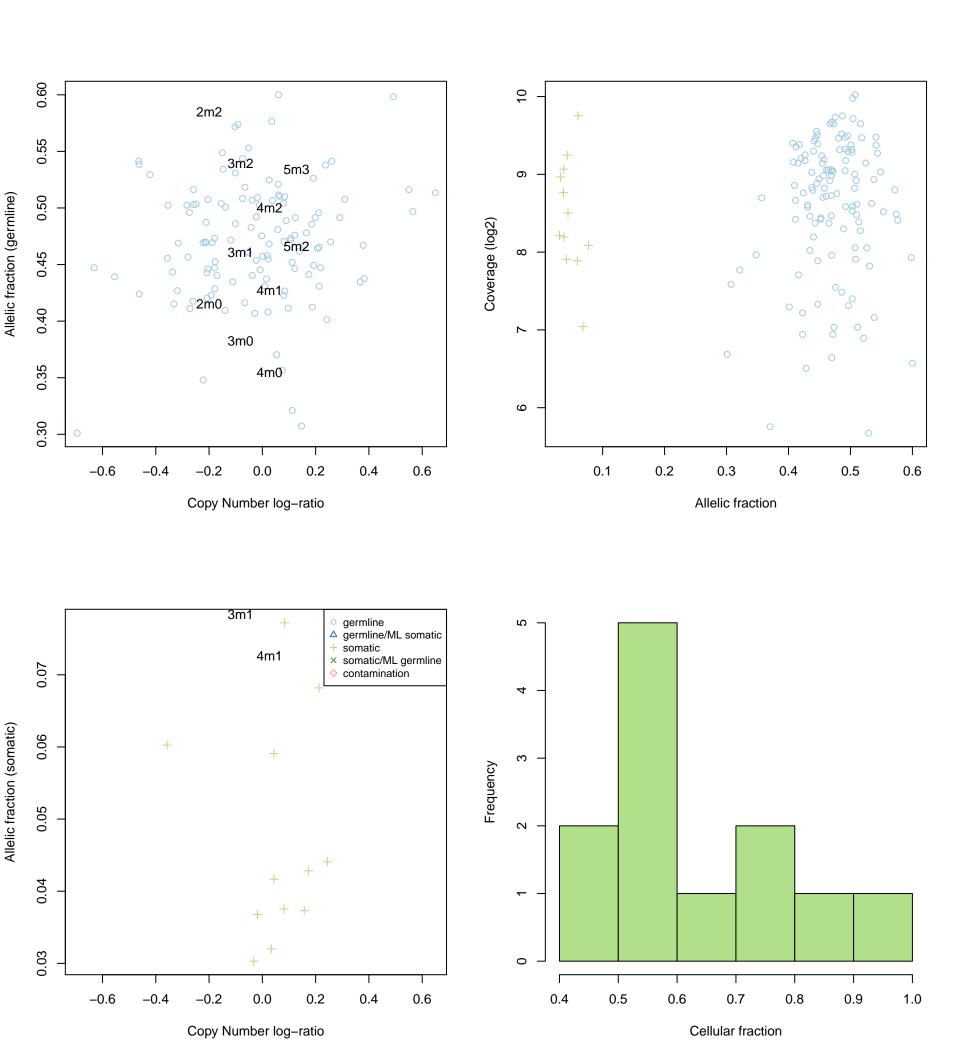




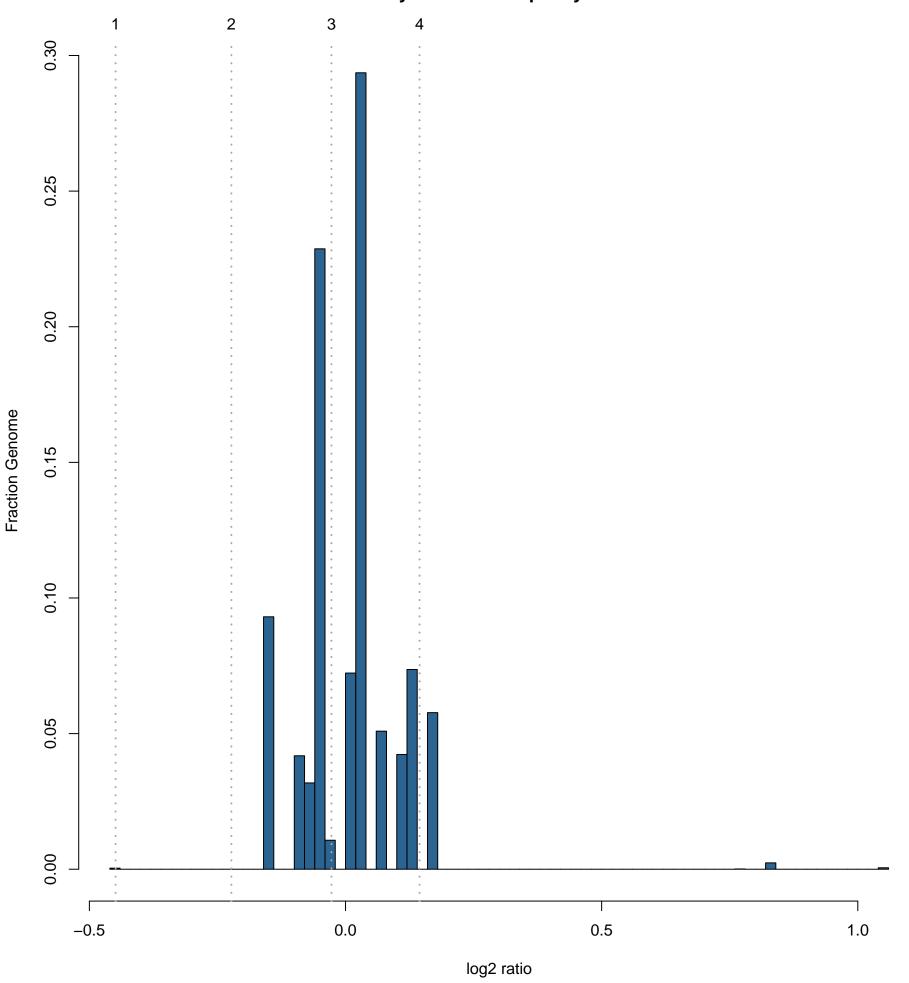
## SCNA-fit log-likelihood: -14502.4

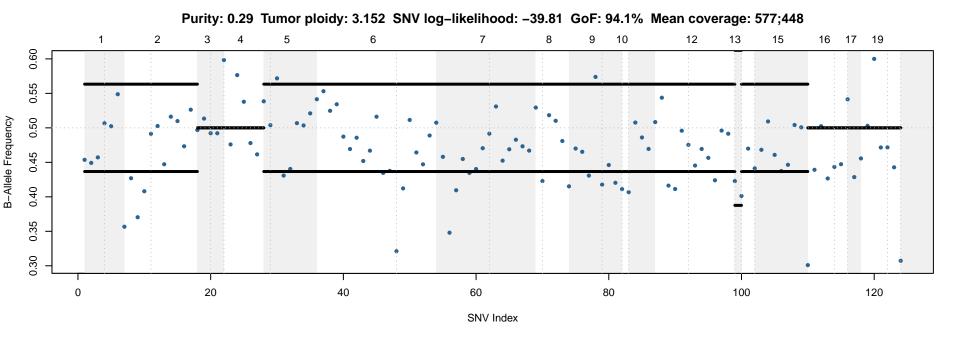




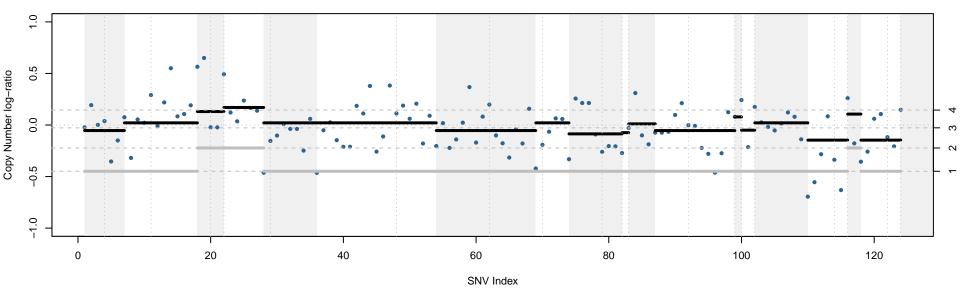


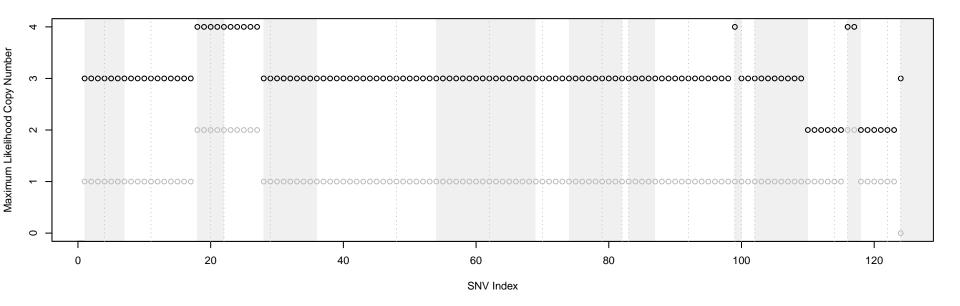
Purity: 0.29 Tumor ploidy: 3.152

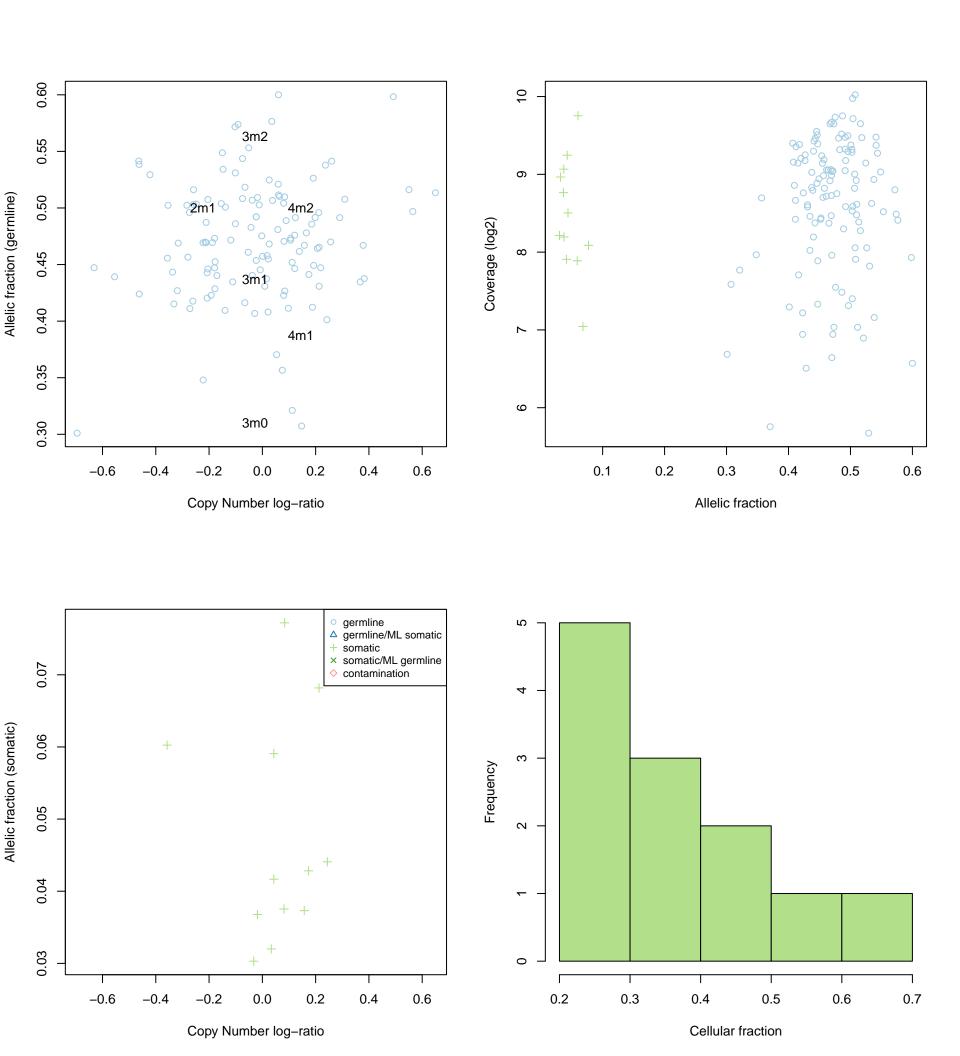




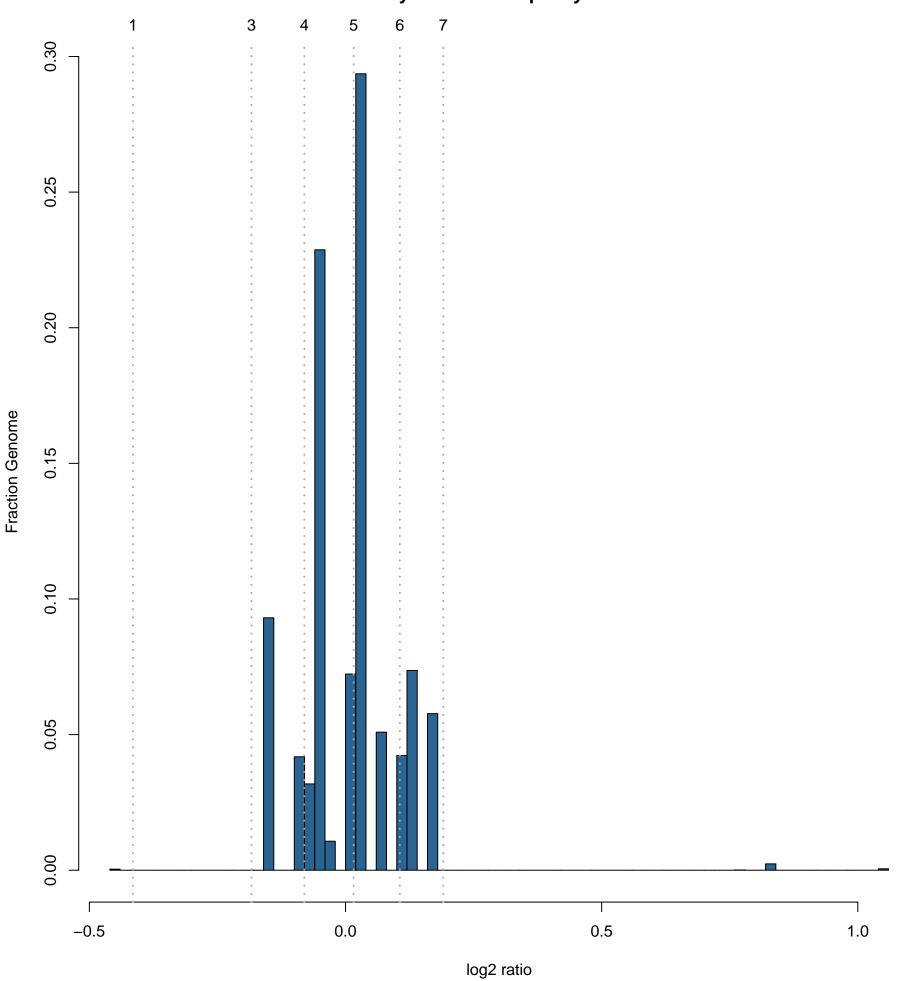
## SCNA-fit log-likelihood: -14562.08

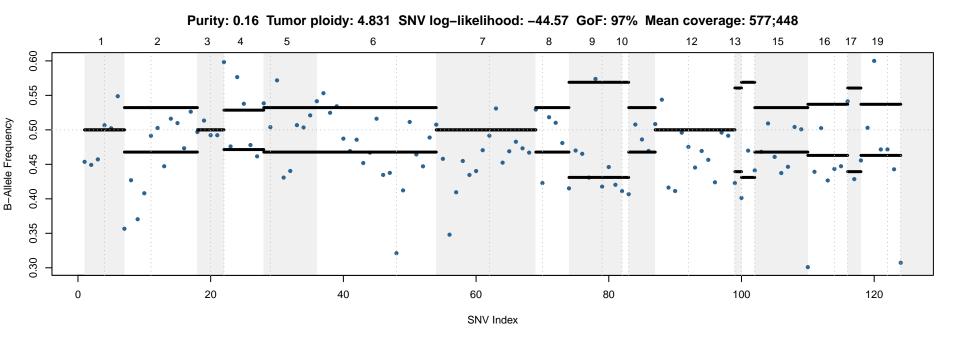




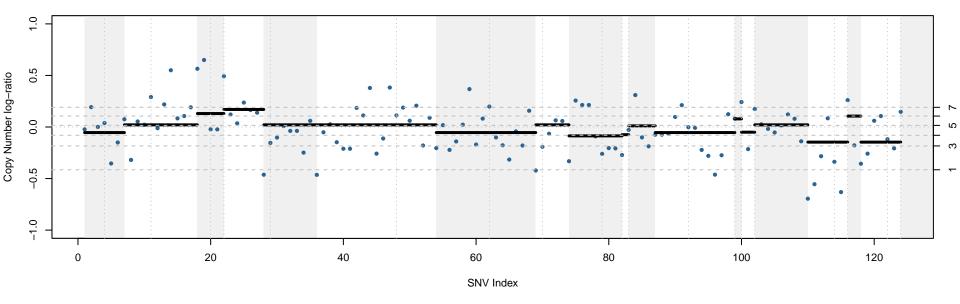


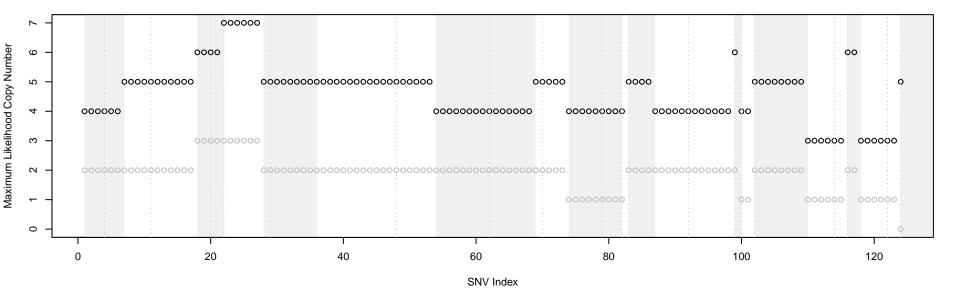
Purity: 0.16 Tumor ploidy: 4.831

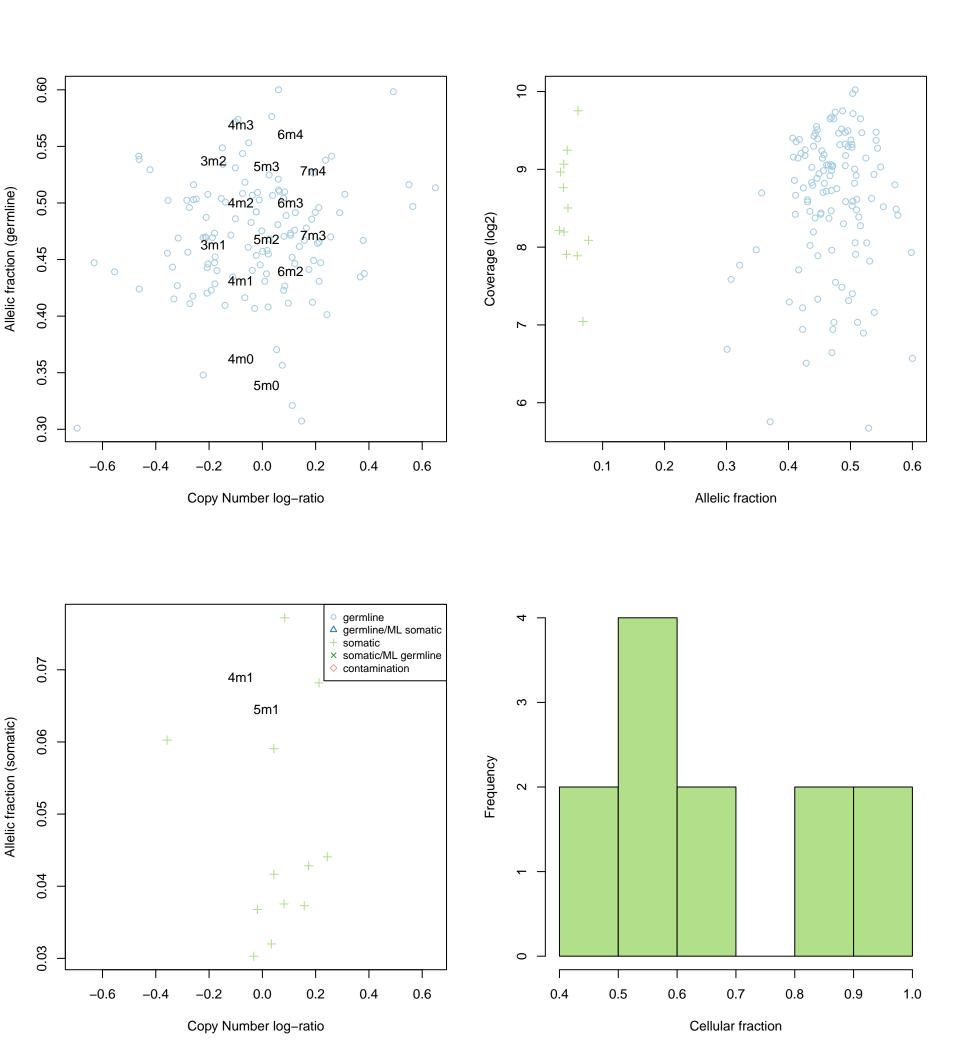




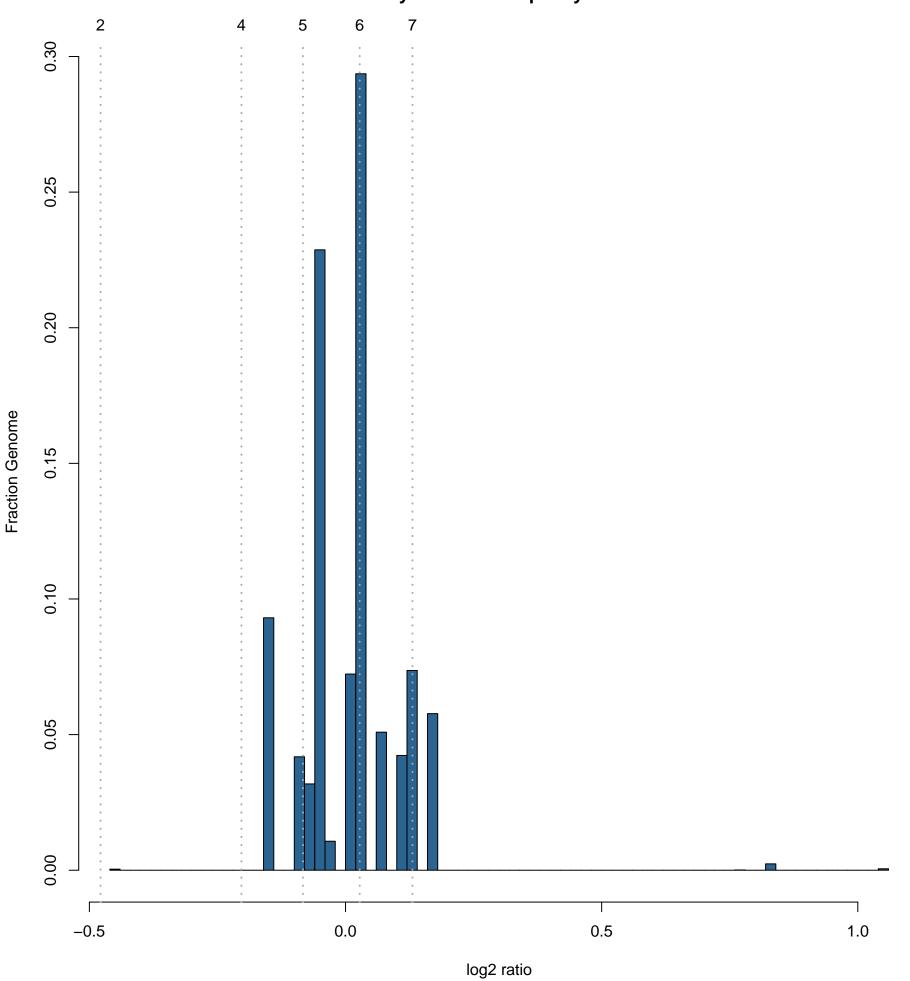
## SCNA-fit log-likelihood: -14495.11

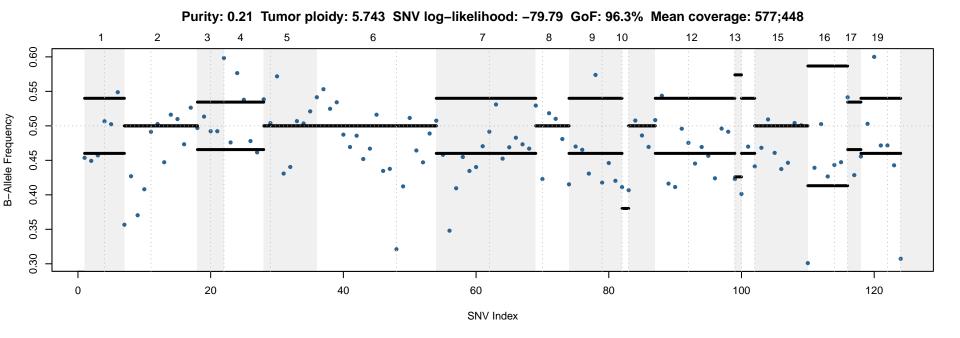




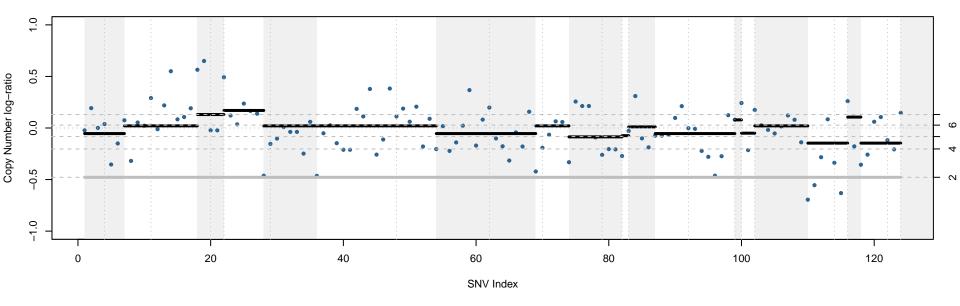


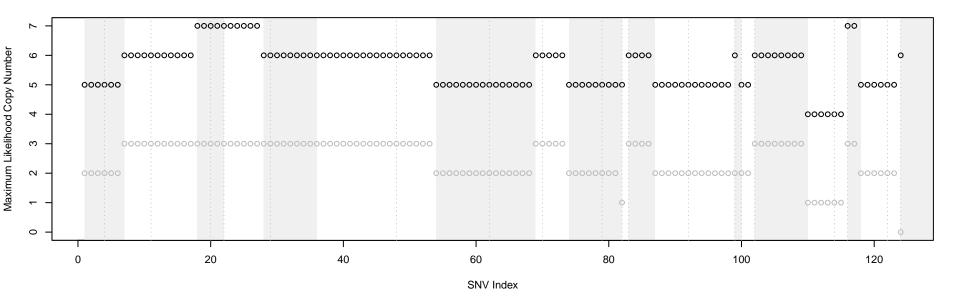
Purity: 0.21 Tumor ploidy: 5.743

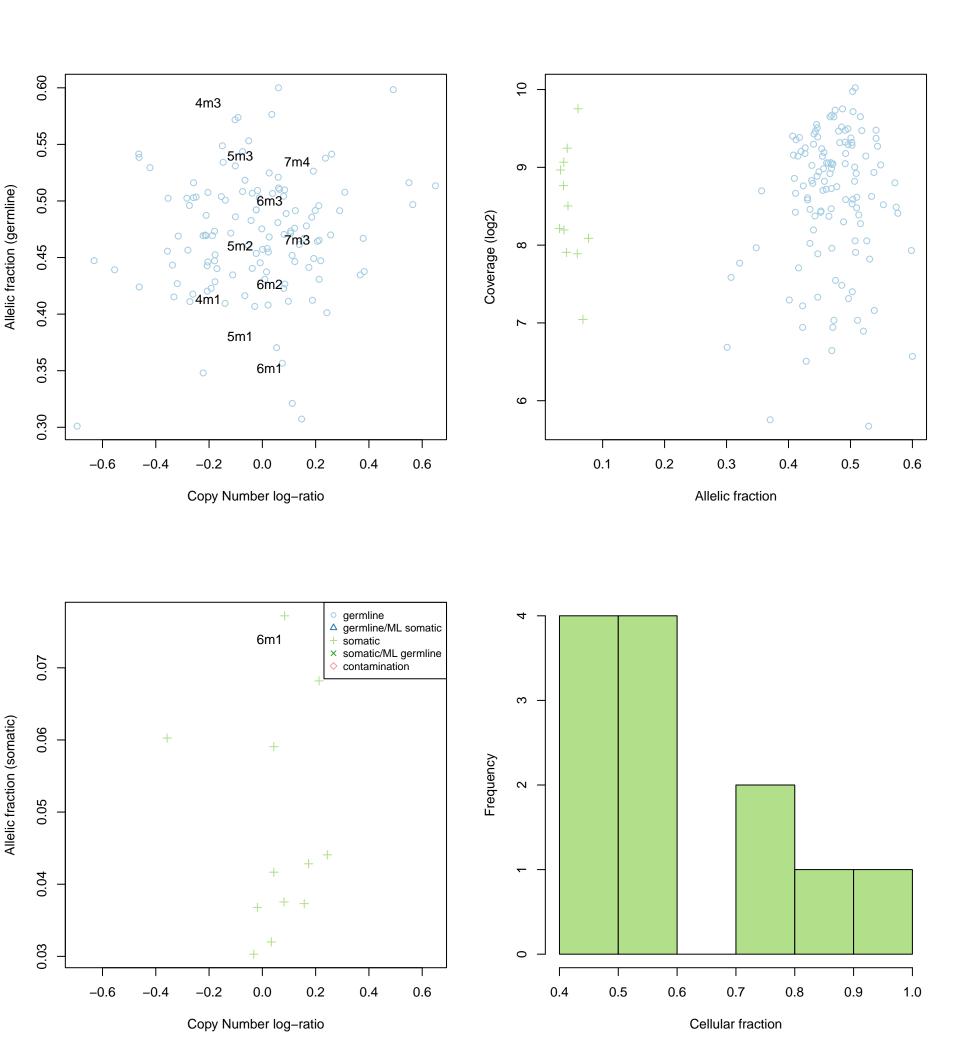




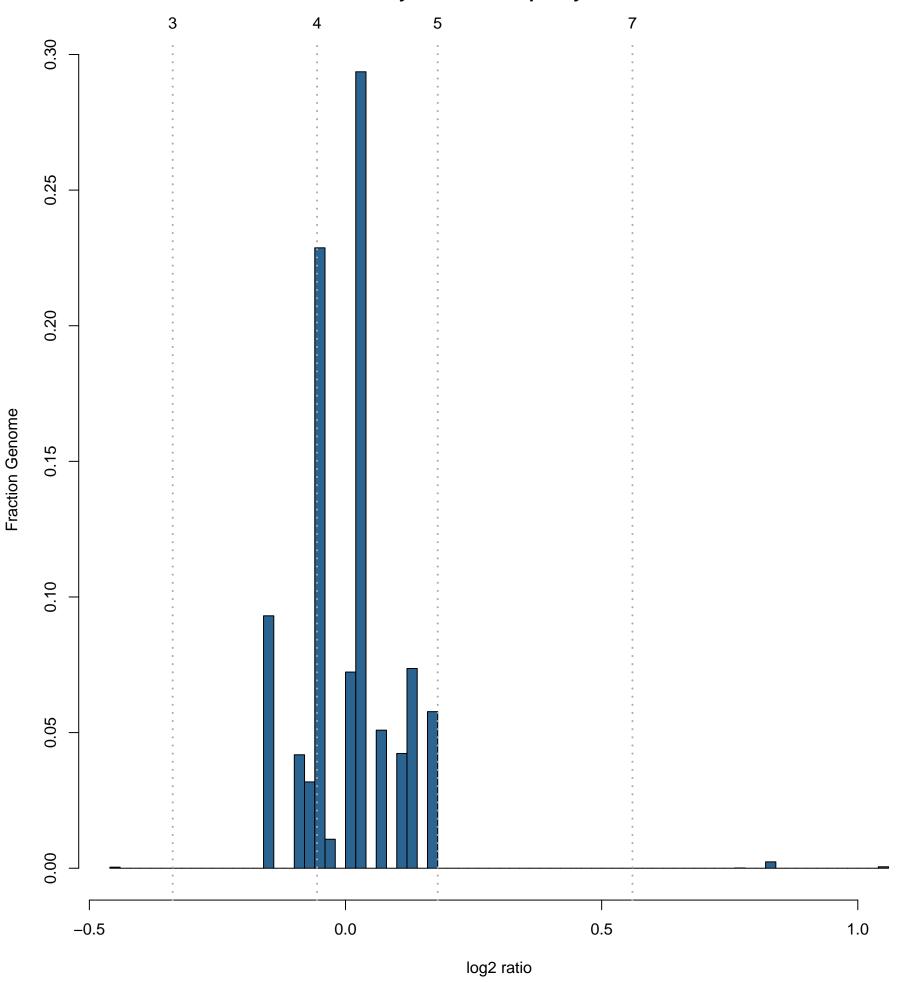
### SCNA-fit log-likelihood: -14503.49

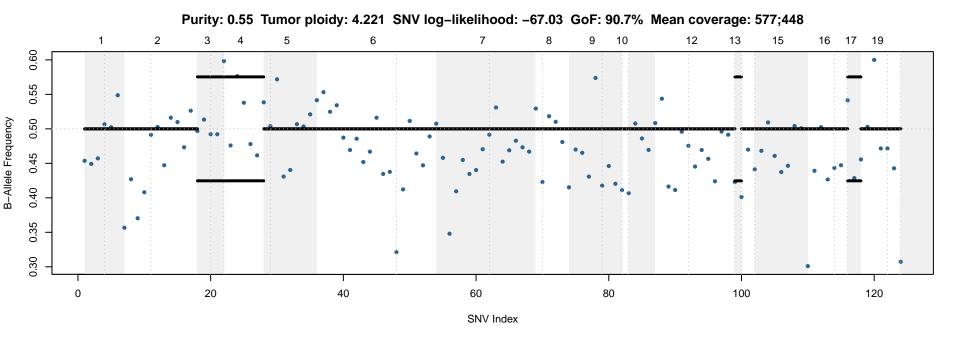




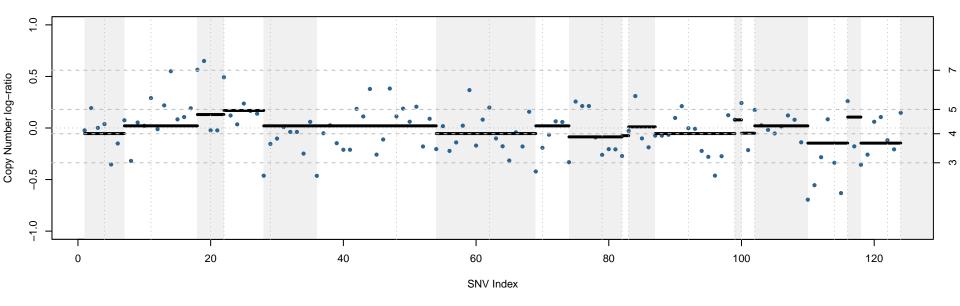


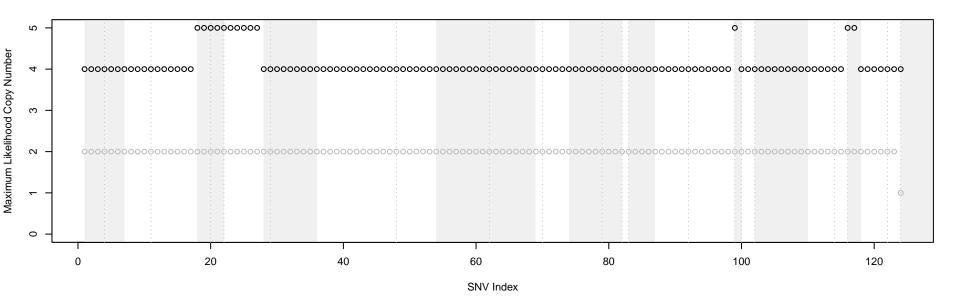
Purity: 0.55 Tumor ploidy: 4.221

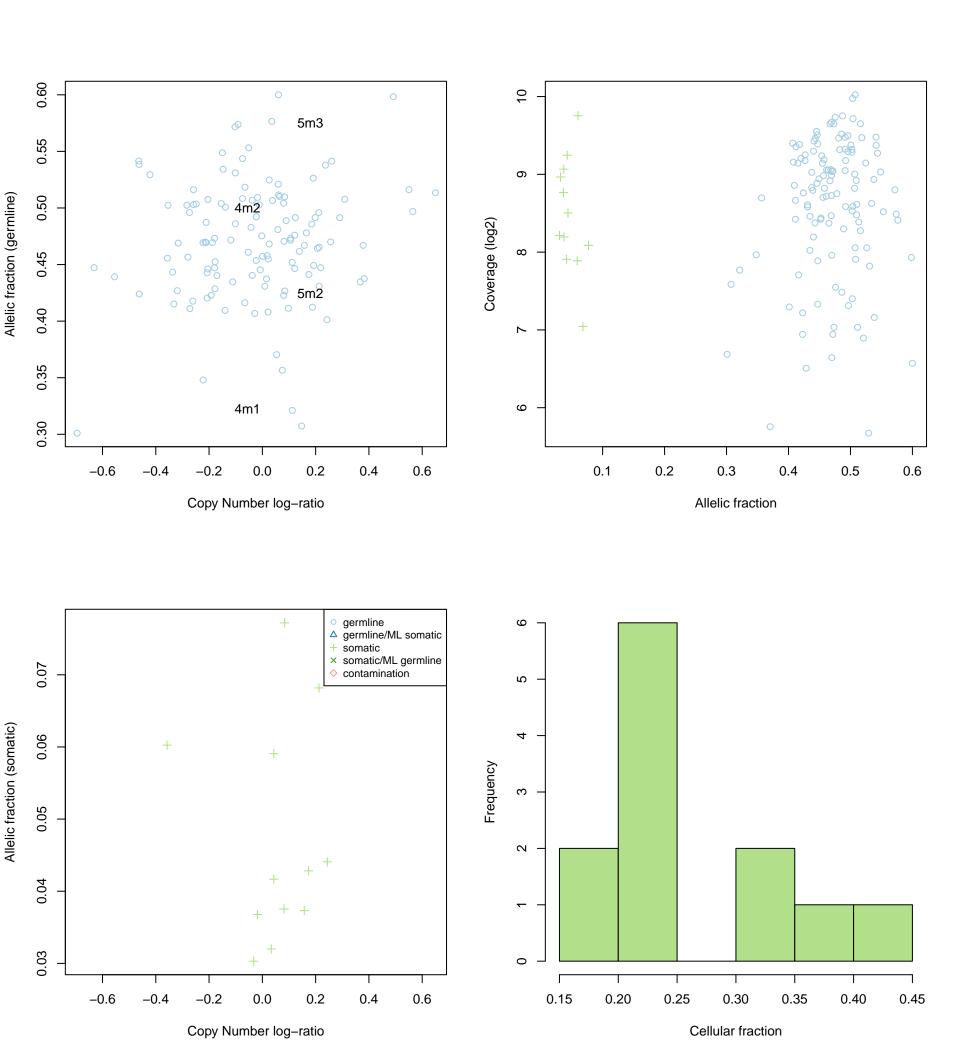




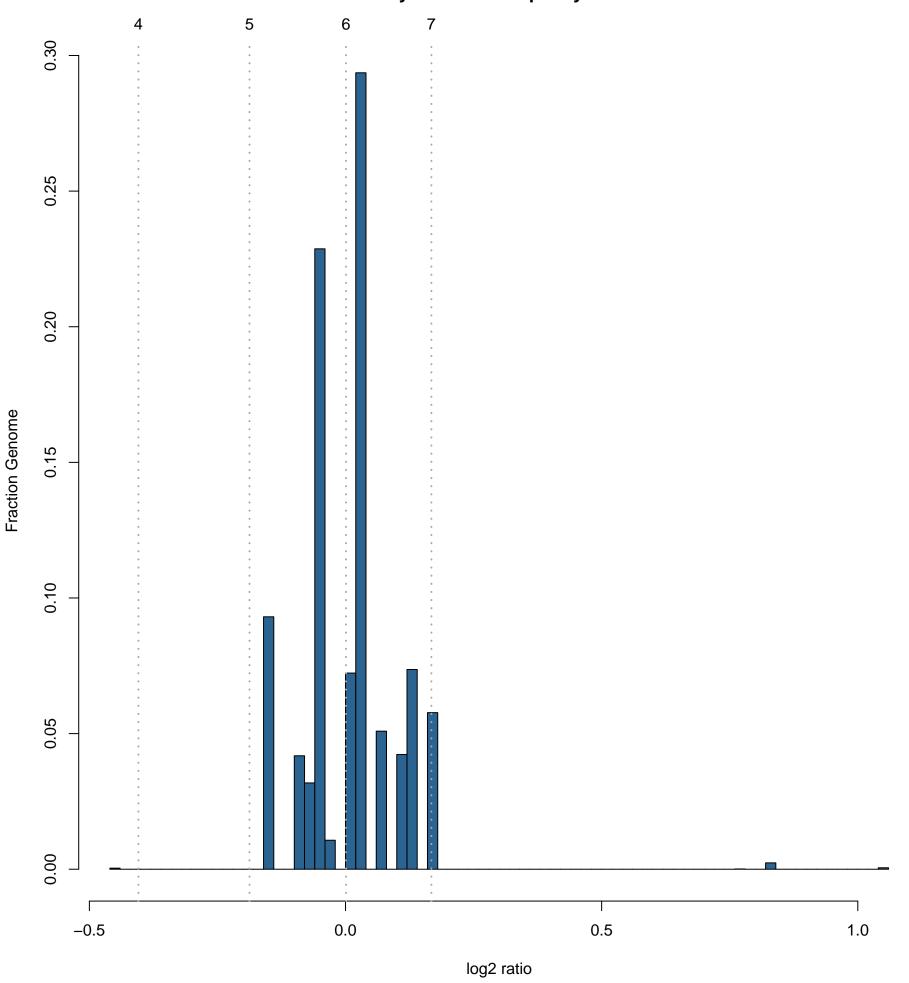
## SCNA-fit log-likelihood: -14640.63

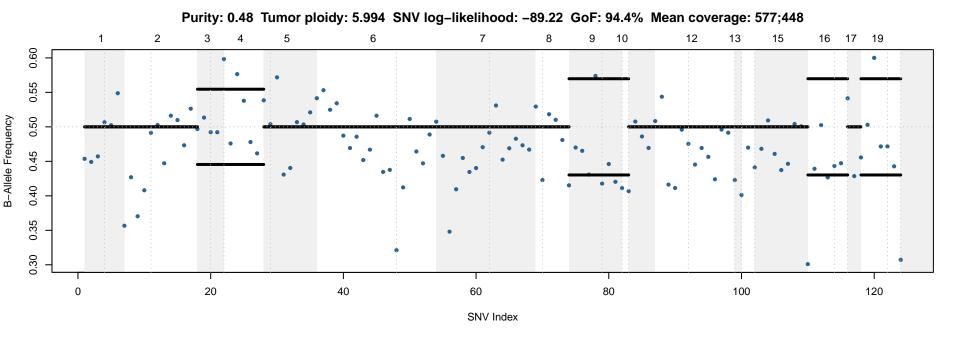




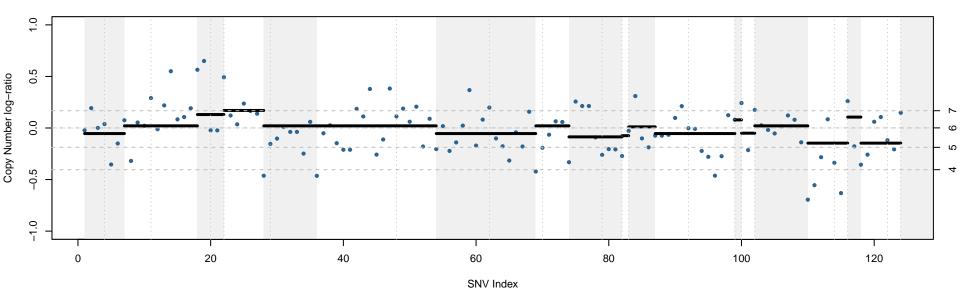


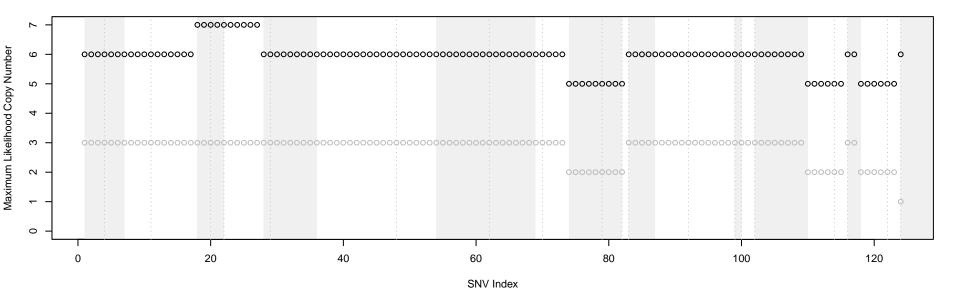
Purity: 0.48 Tumor ploidy: 5.994

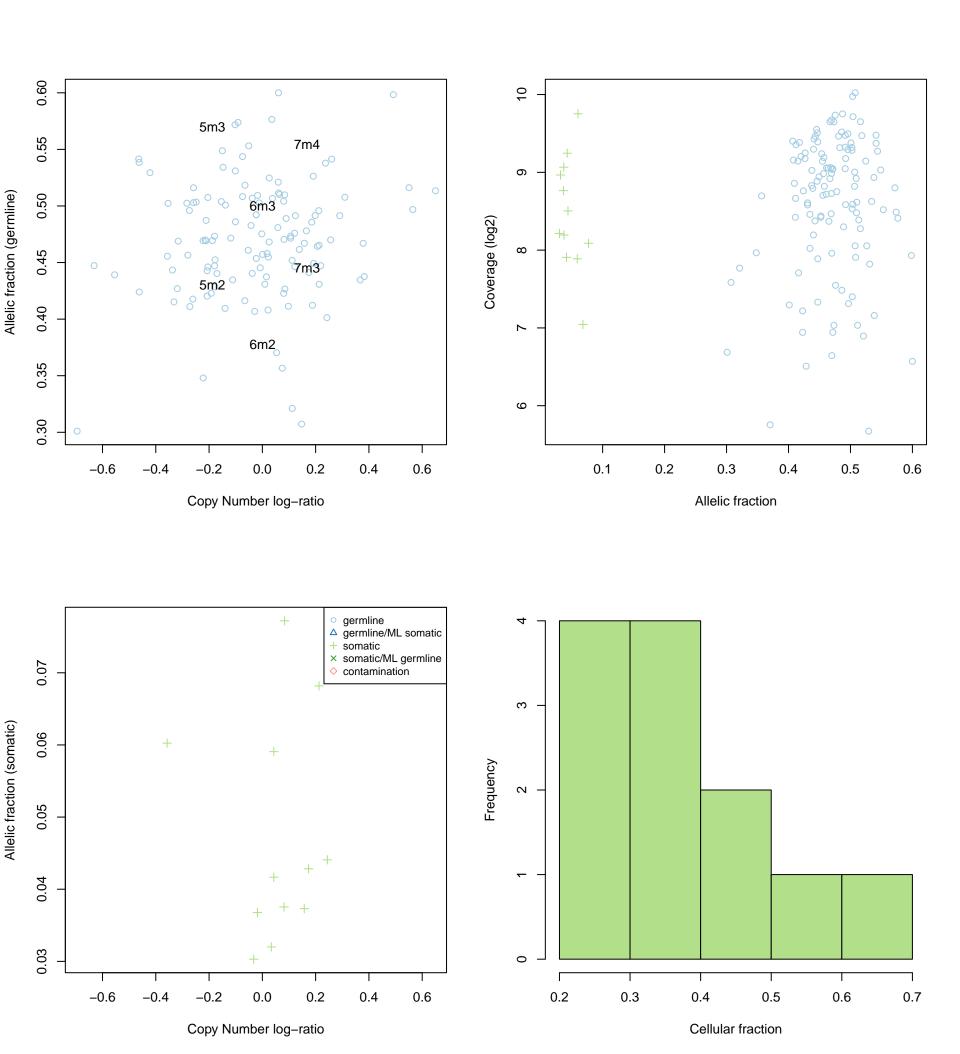




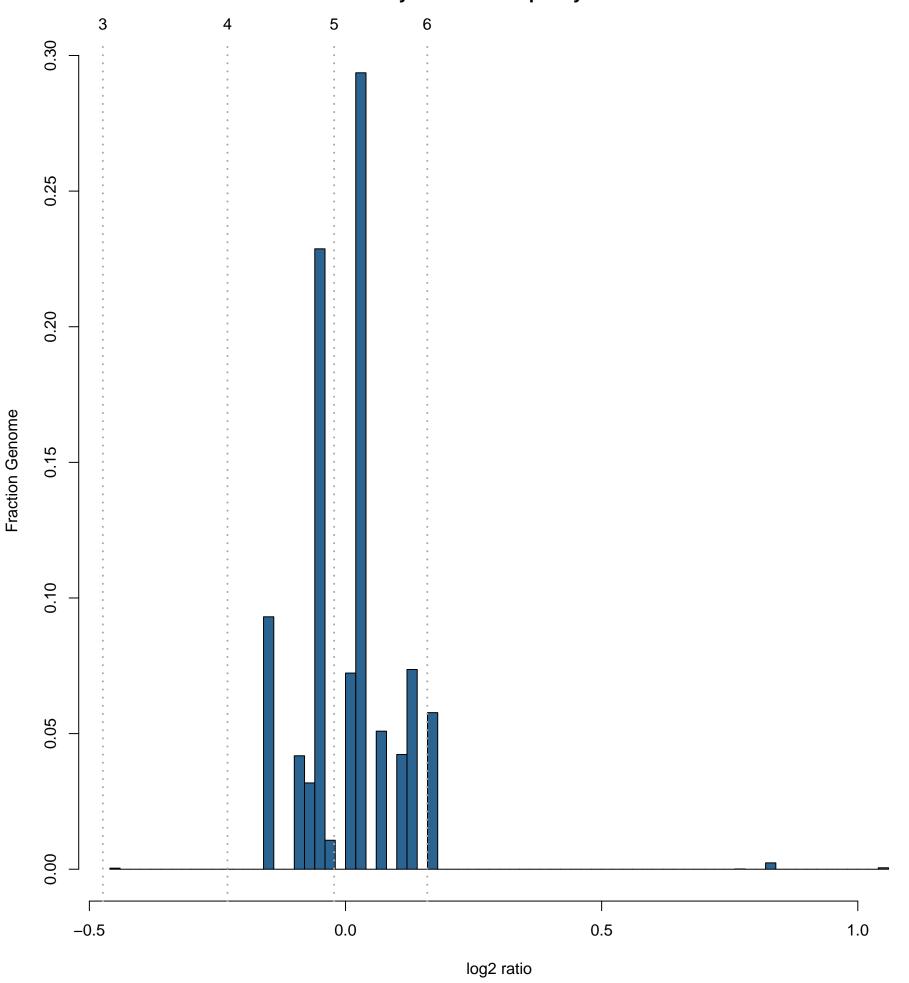
## SCNA-fit log-likelihood: -14583.85

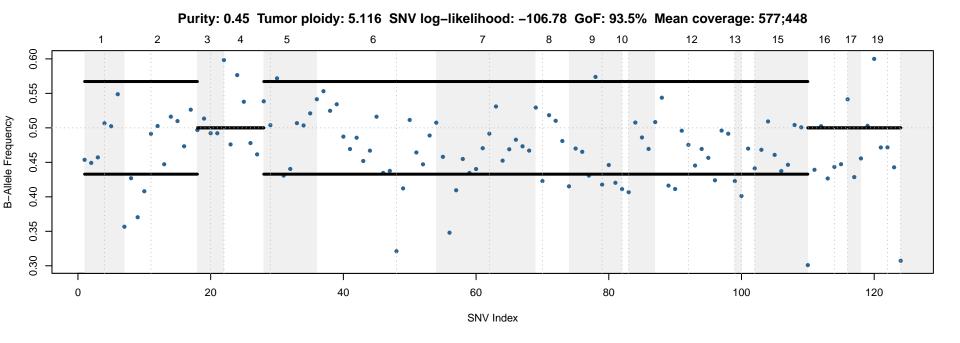




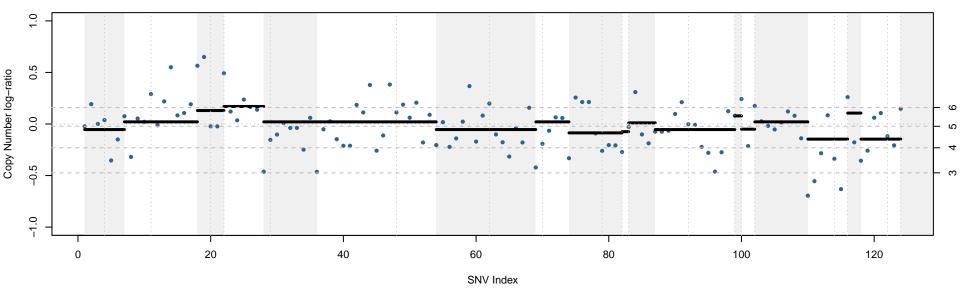


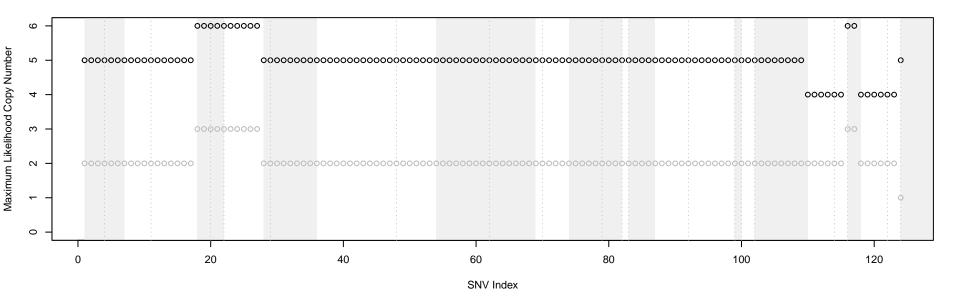
Purity: 0.45 Tumor ploidy: 5.116

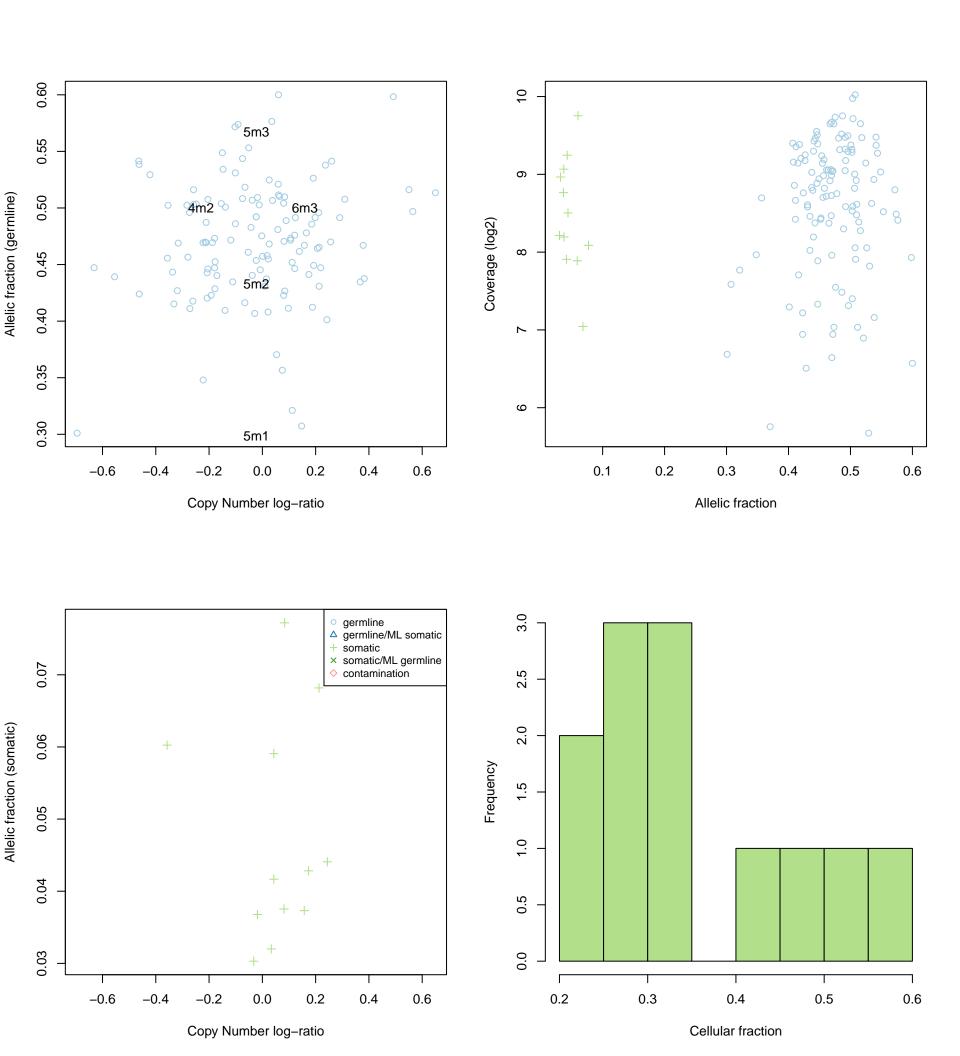




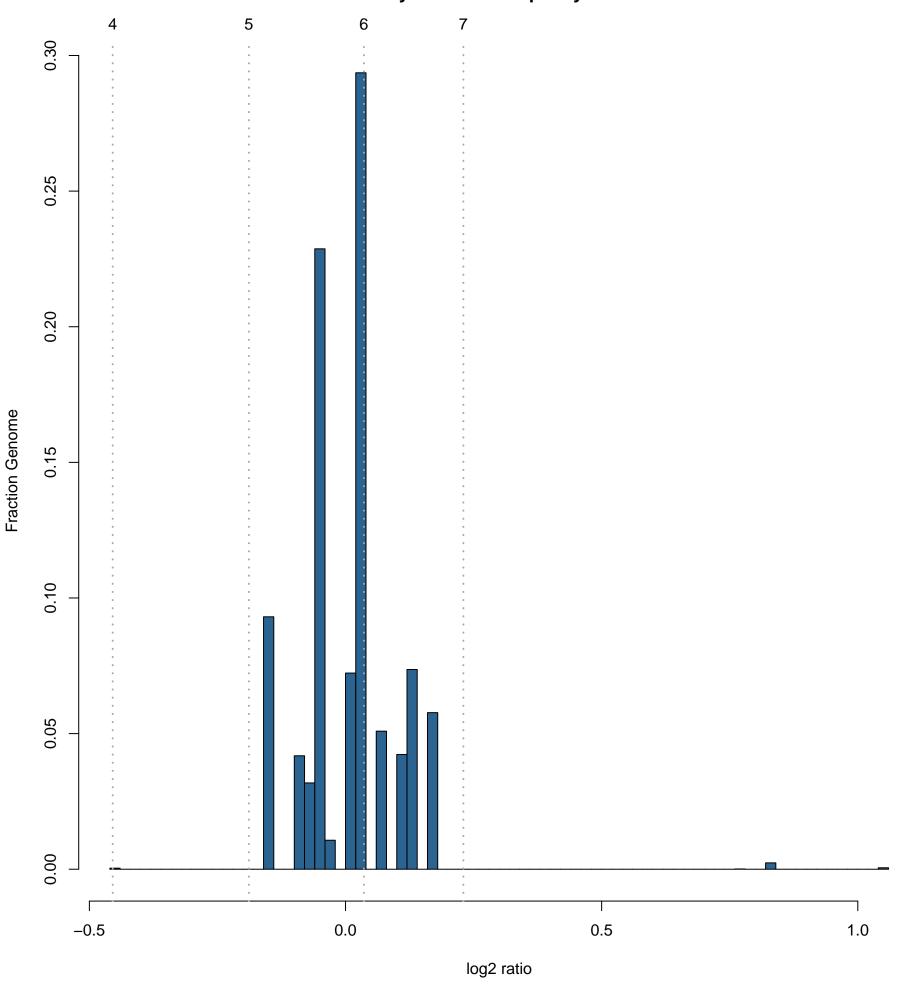
## SCNA-fit log-likelihood: -14574.4

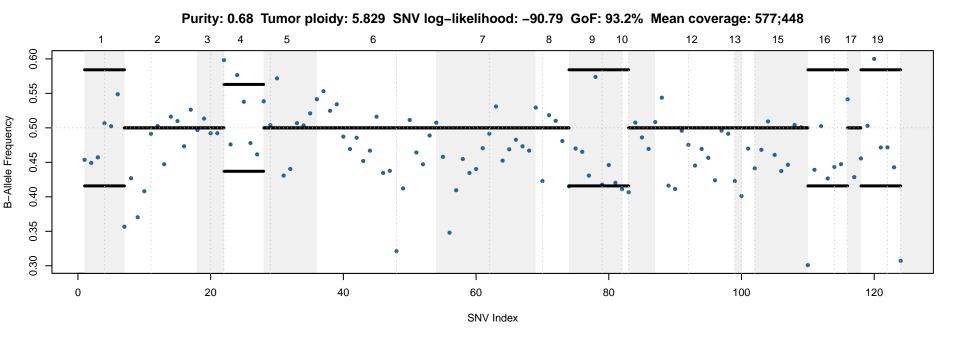




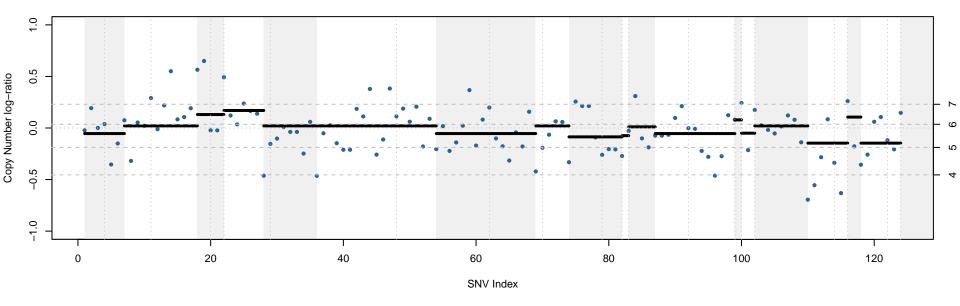


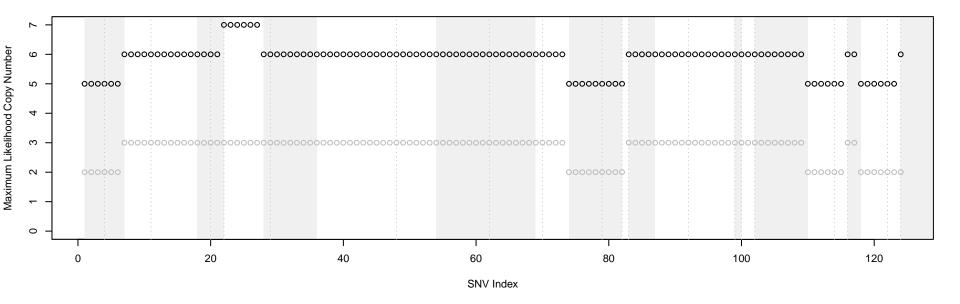
Purity: 0.68 Tumor ploidy: 5.829

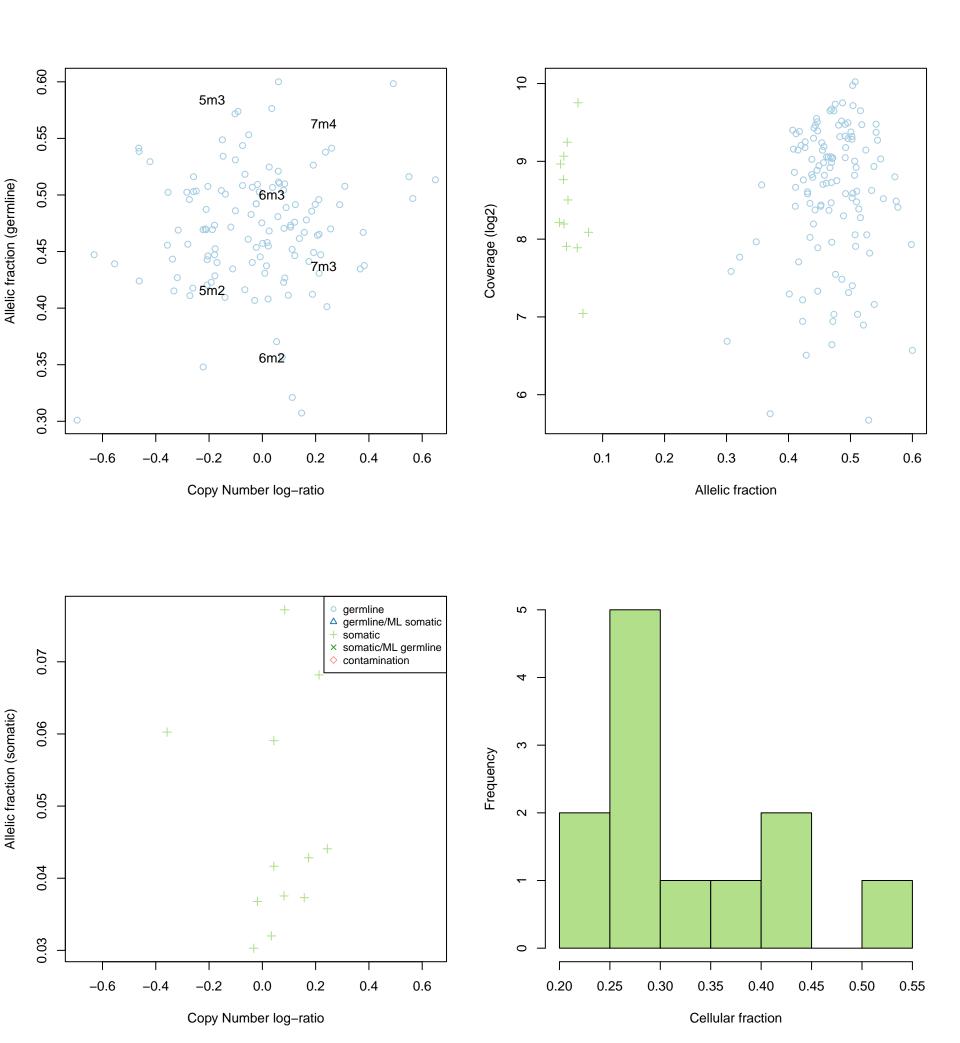




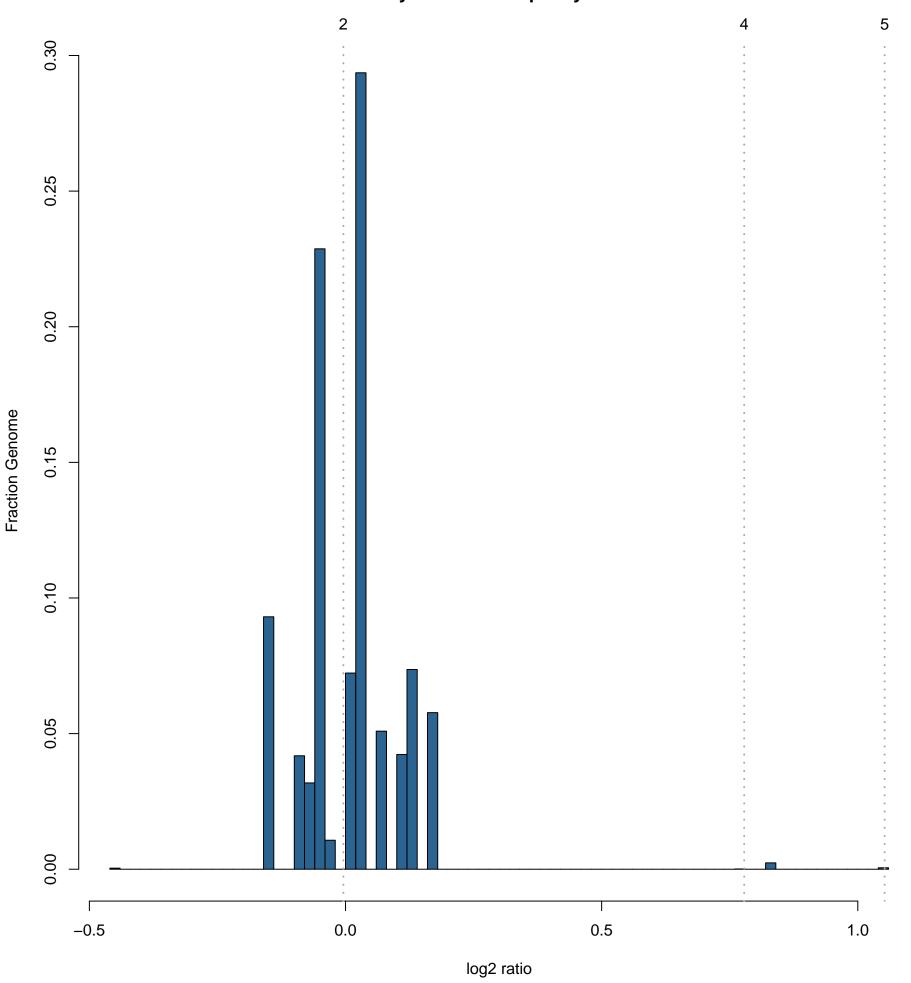
## SCNA-fit log-likelihood: -14657.24

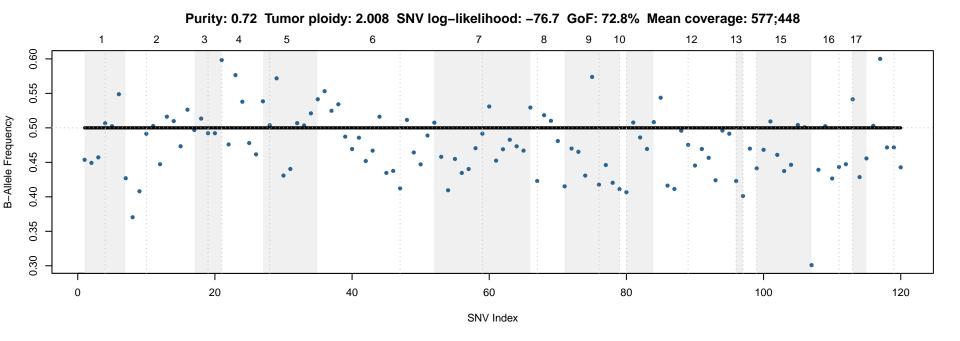




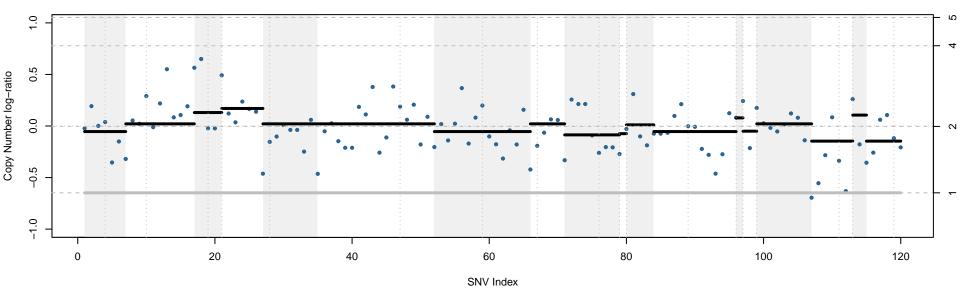


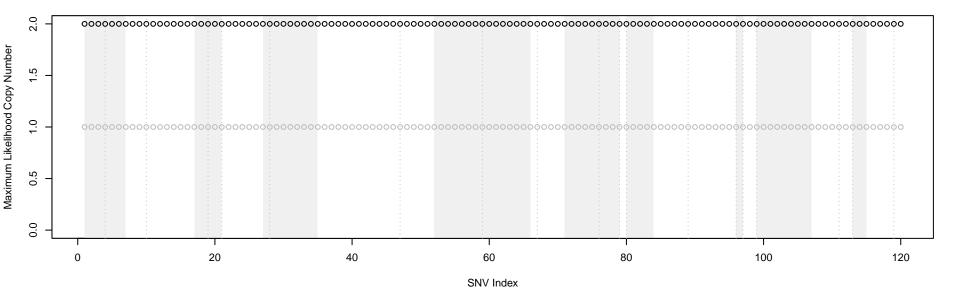
Purity: 0.72 Tumor ploidy: 2.008

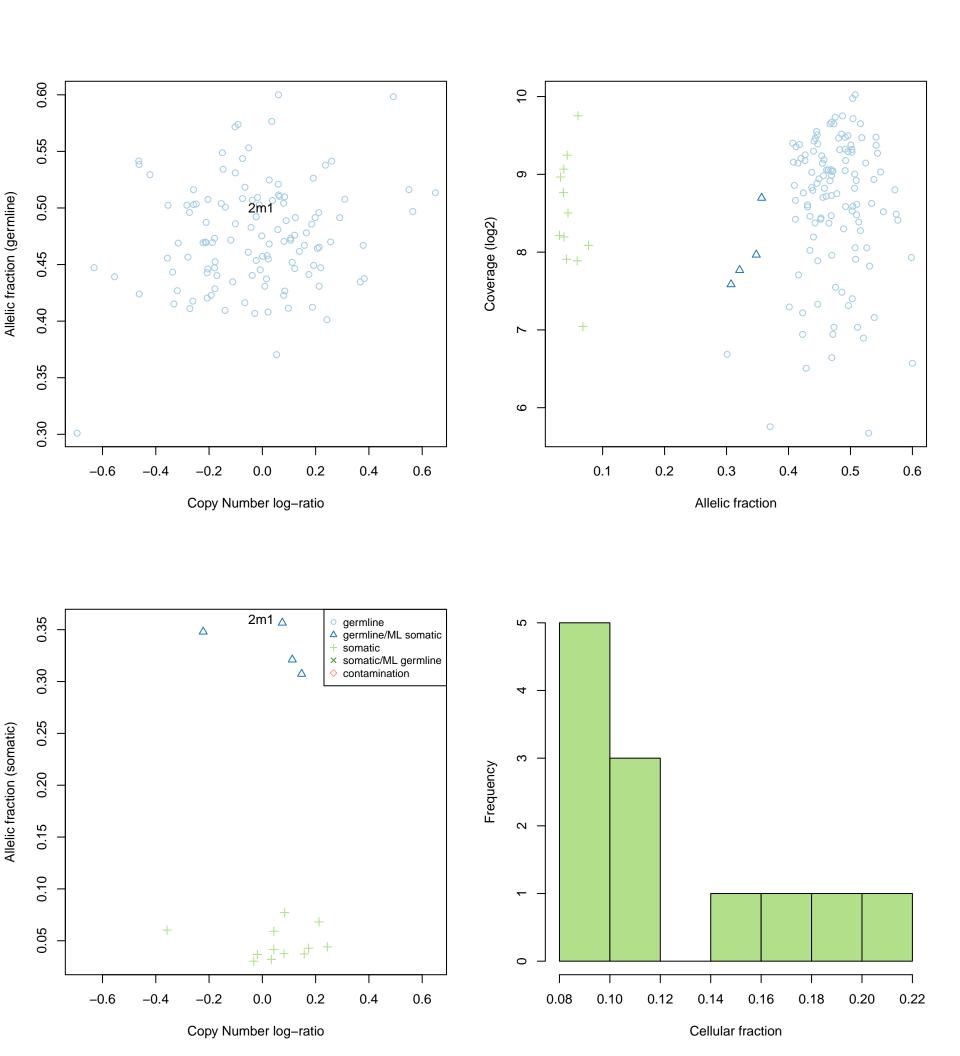




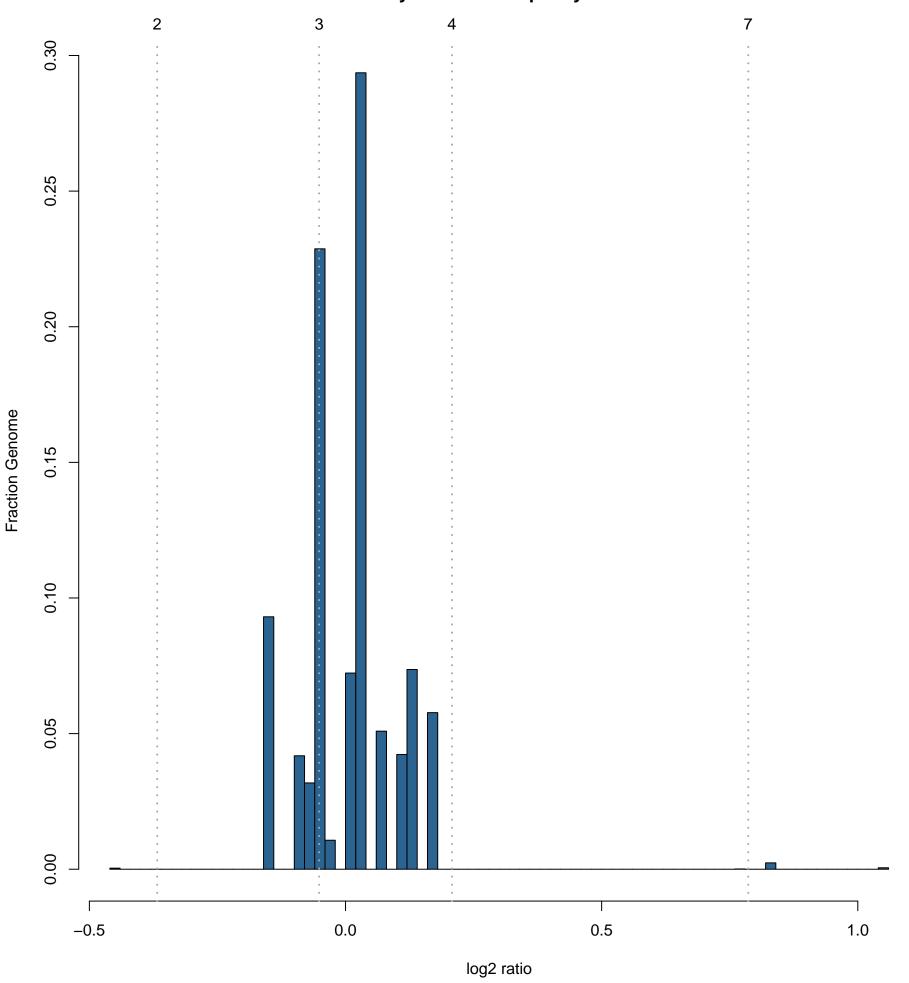
# SCNA-fit log-likelihood: -14736.02

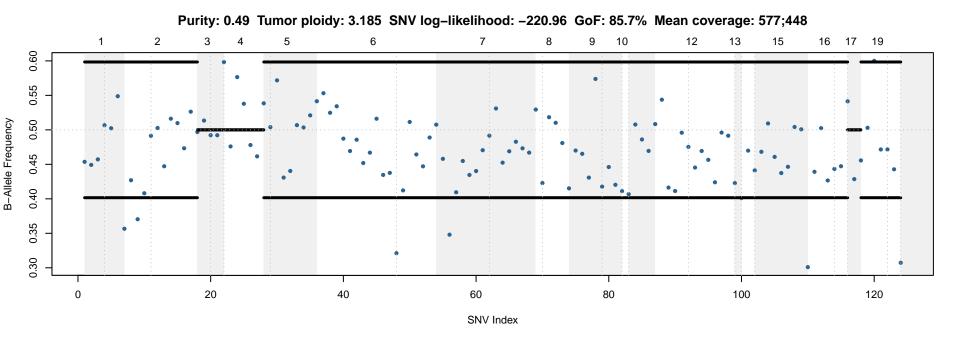




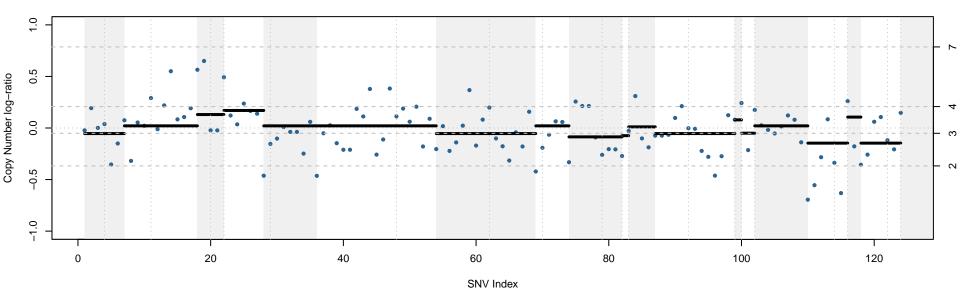


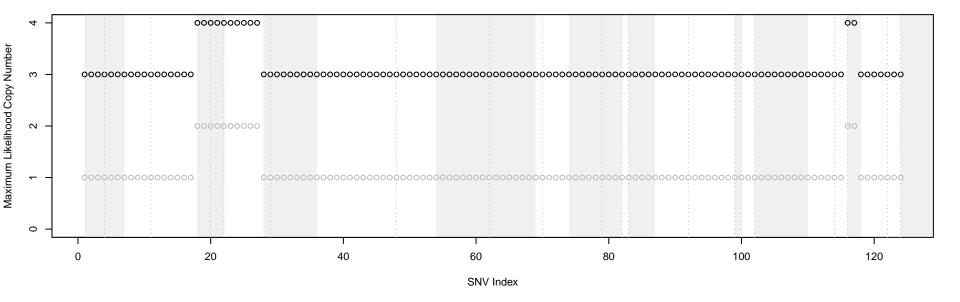
Purity: 0.49 Tumor ploidy: 3.185

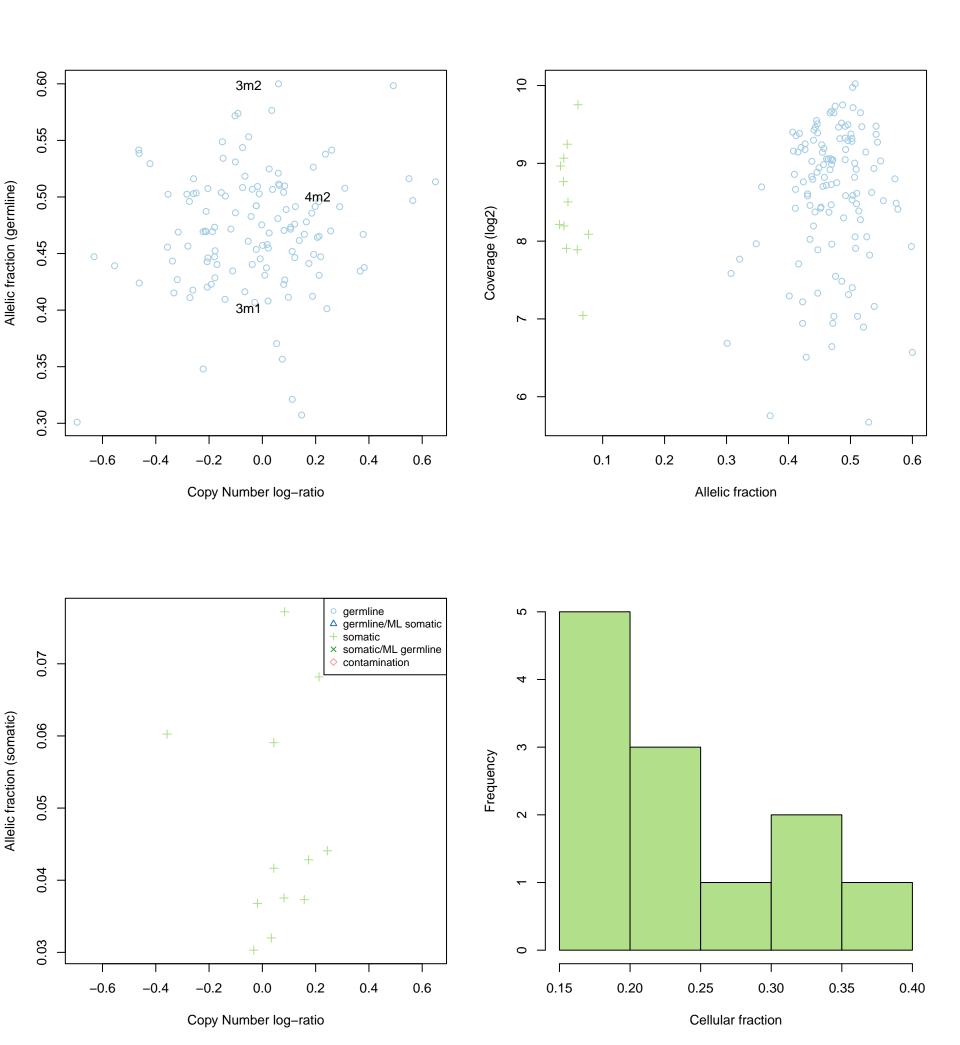




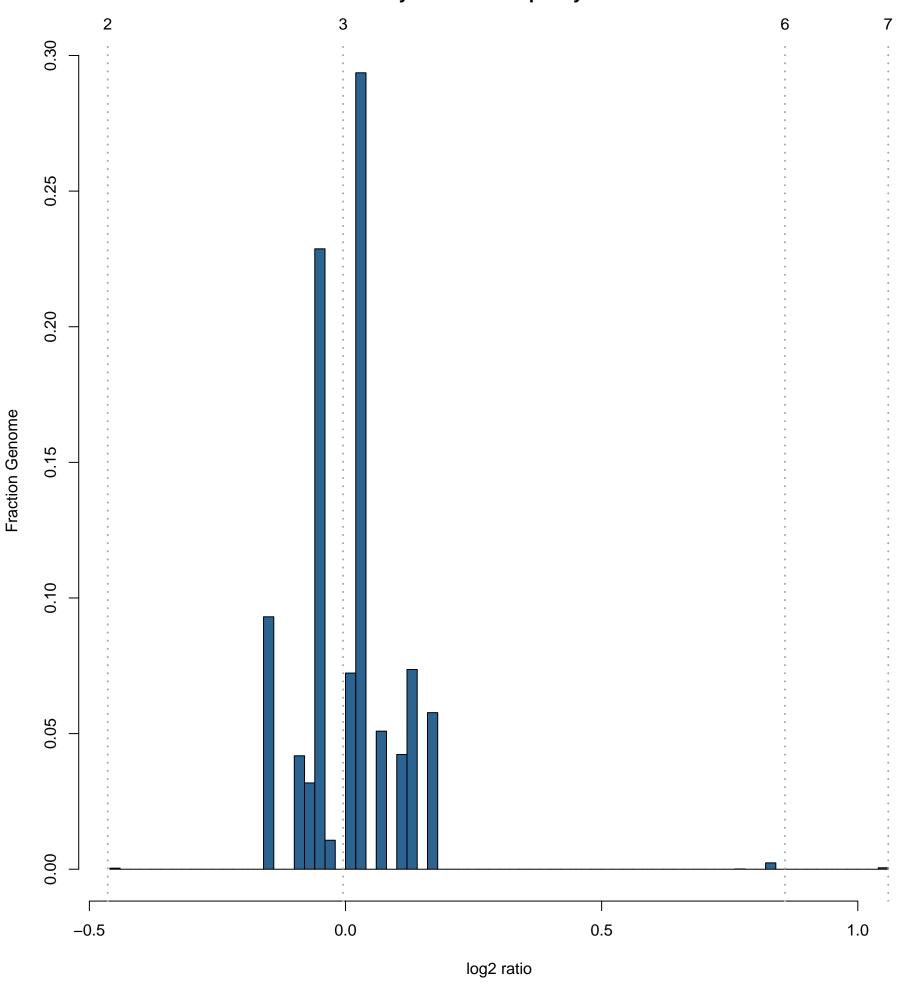
## SCNA-fit log-likelihood: -14634.1

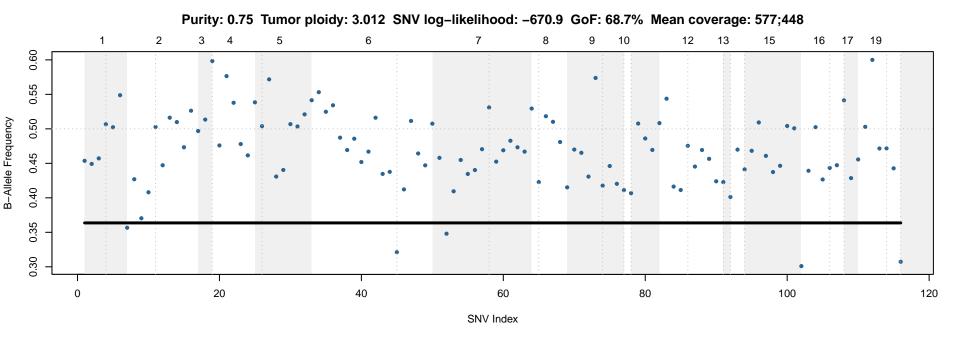






Purity: 0.75 Tumor ploidy: 3.012





# SCNA-fit log-likelihood: -14735.65

