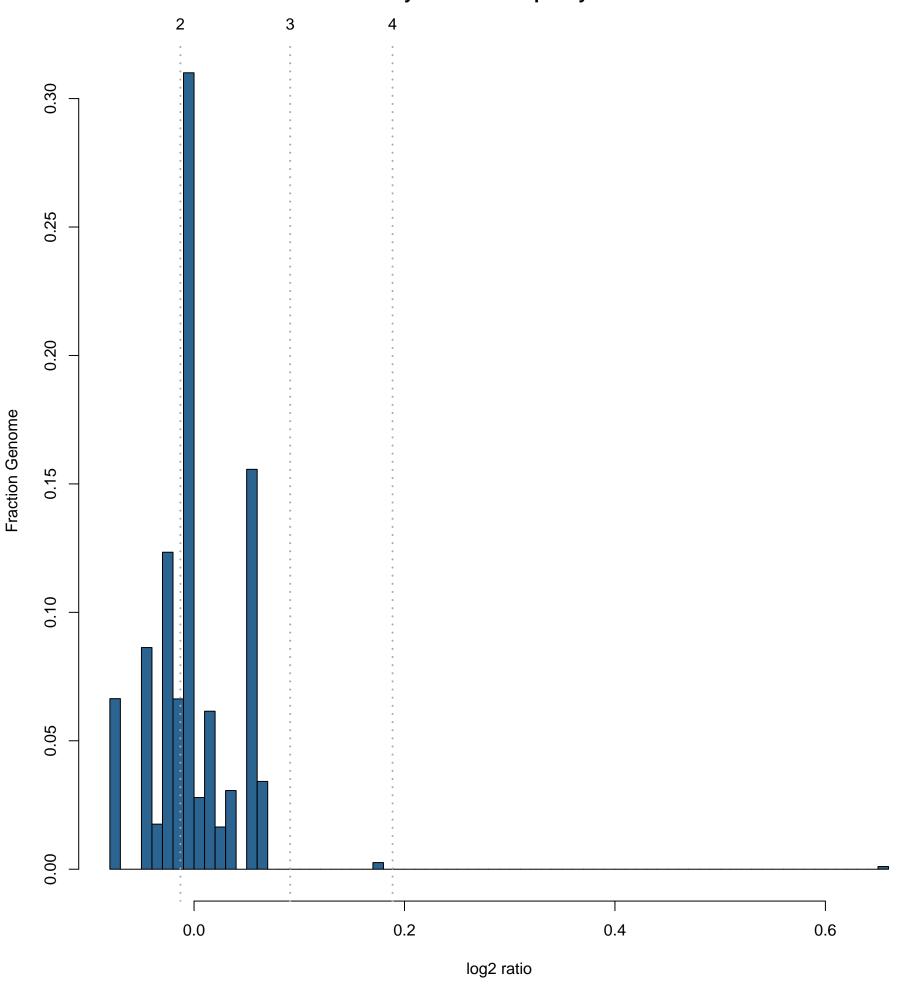
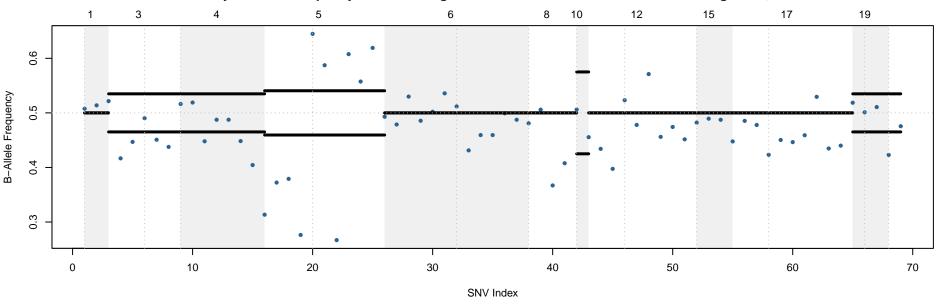
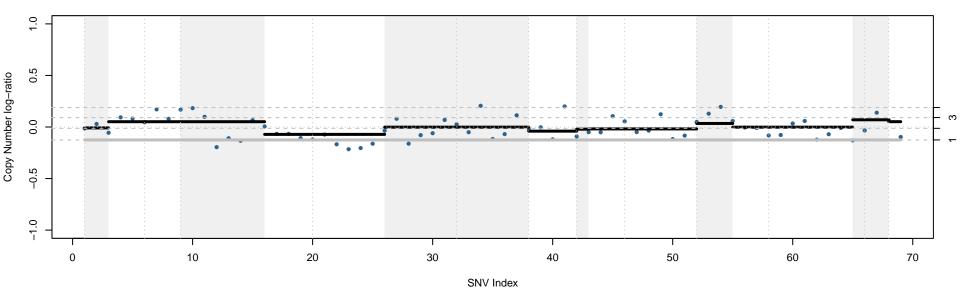
Purity: 0.15 Tumor ploidy: 2.12

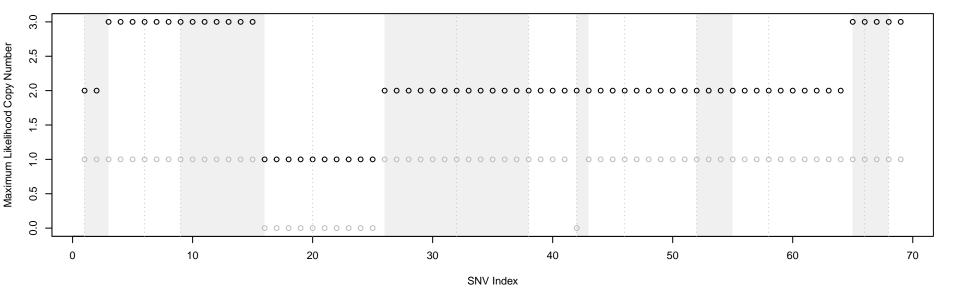


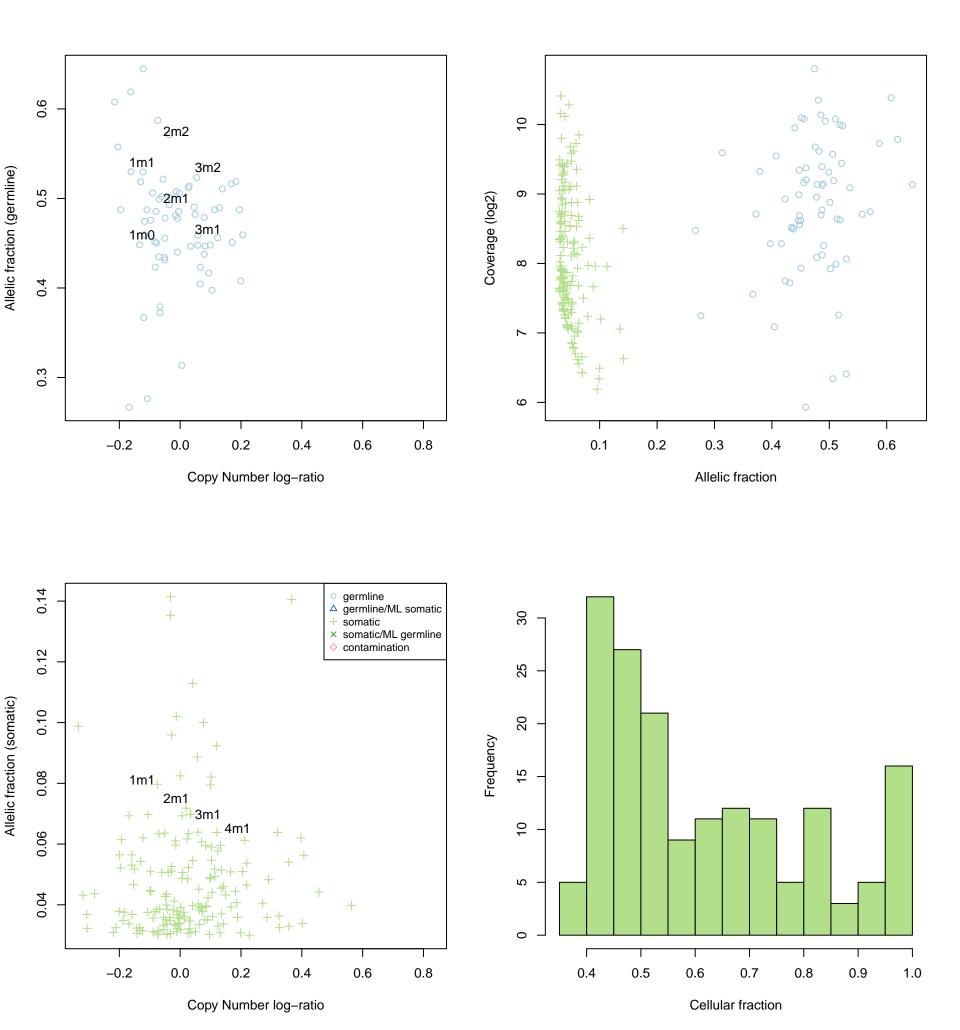
Purity: 0.15 Tumor ploidy: 2.12 SNV log-likelihood: 83.3 GoF: 95.9% Mean coverage: 373;394



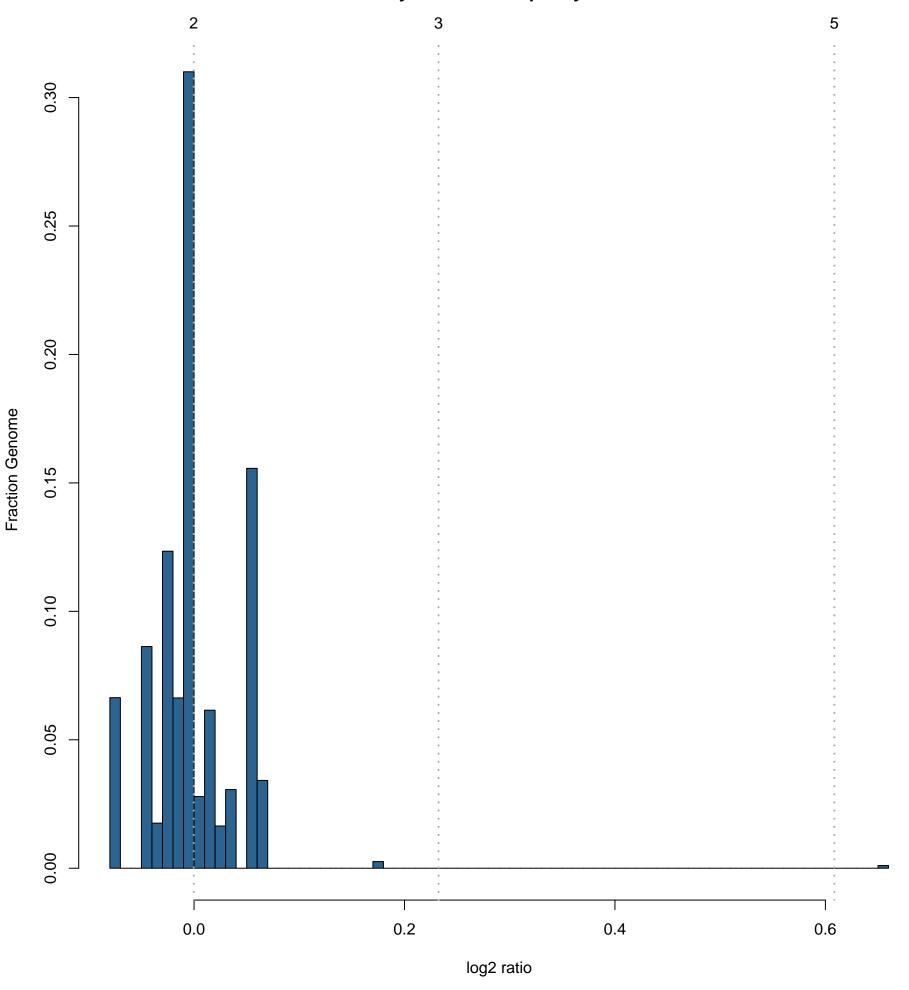
SCNA-fit log-likelihood: -1084.02

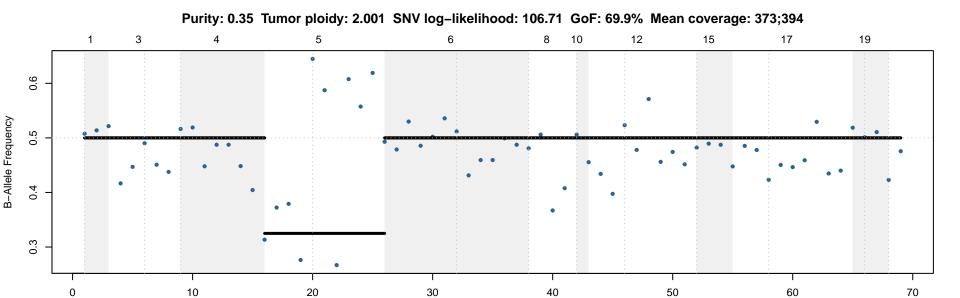






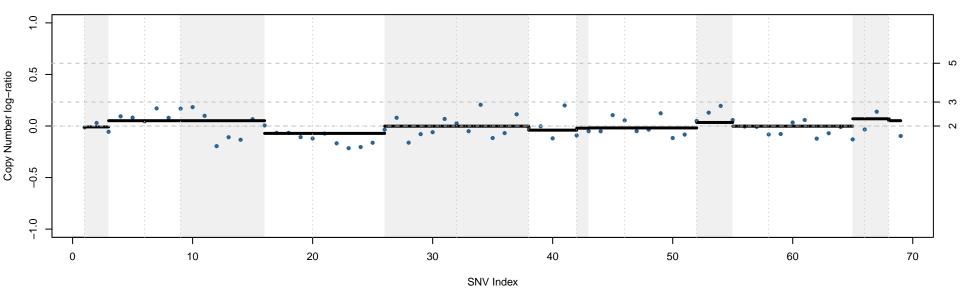
Purity: 0.35 Tumor ploidy: 2.001

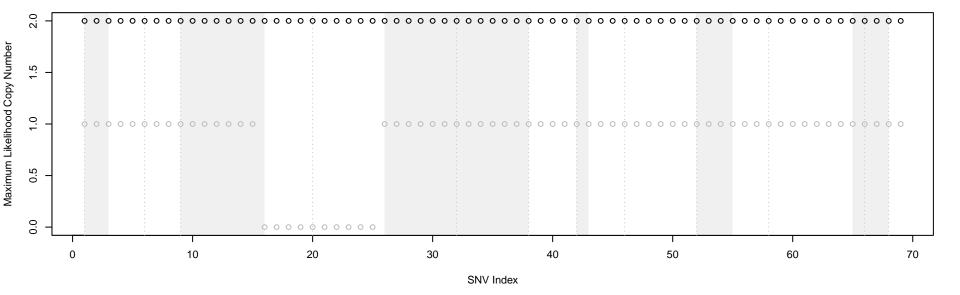


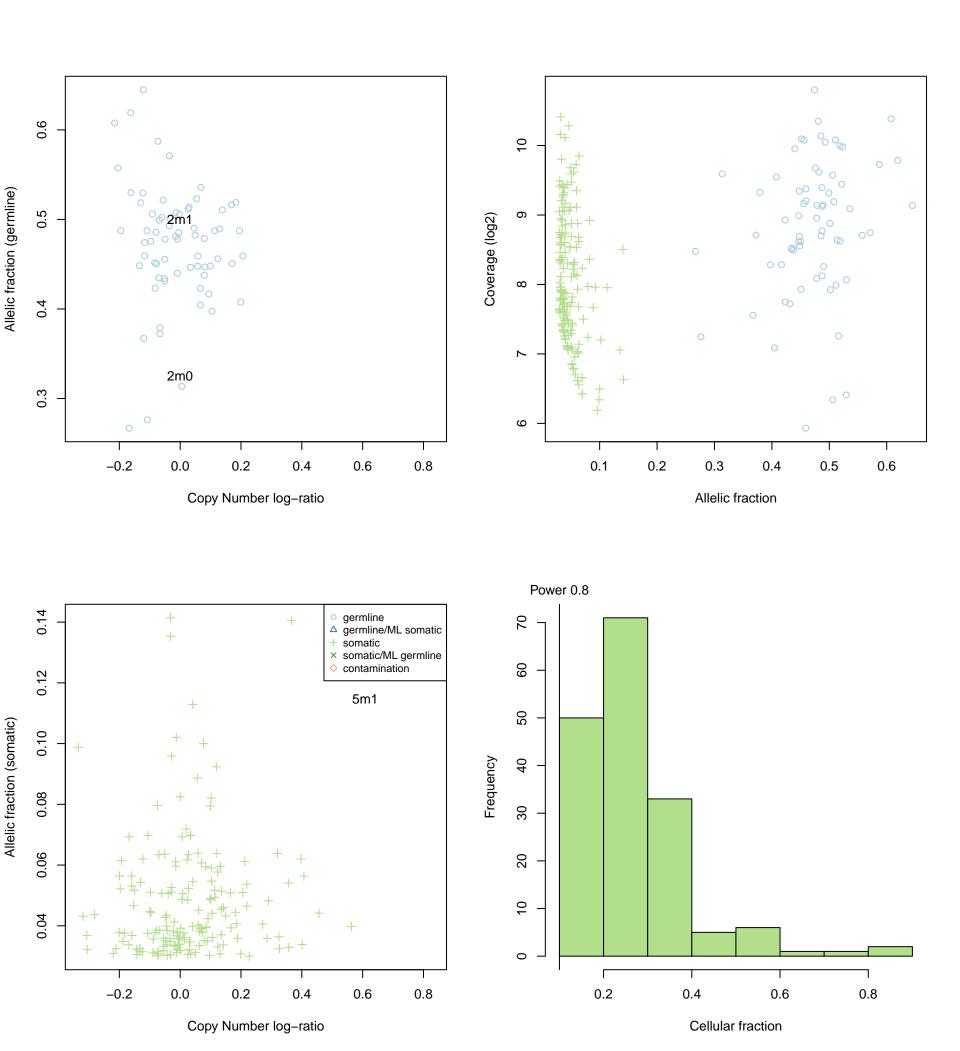


SCNA-fit log-likelihood: -1133.35

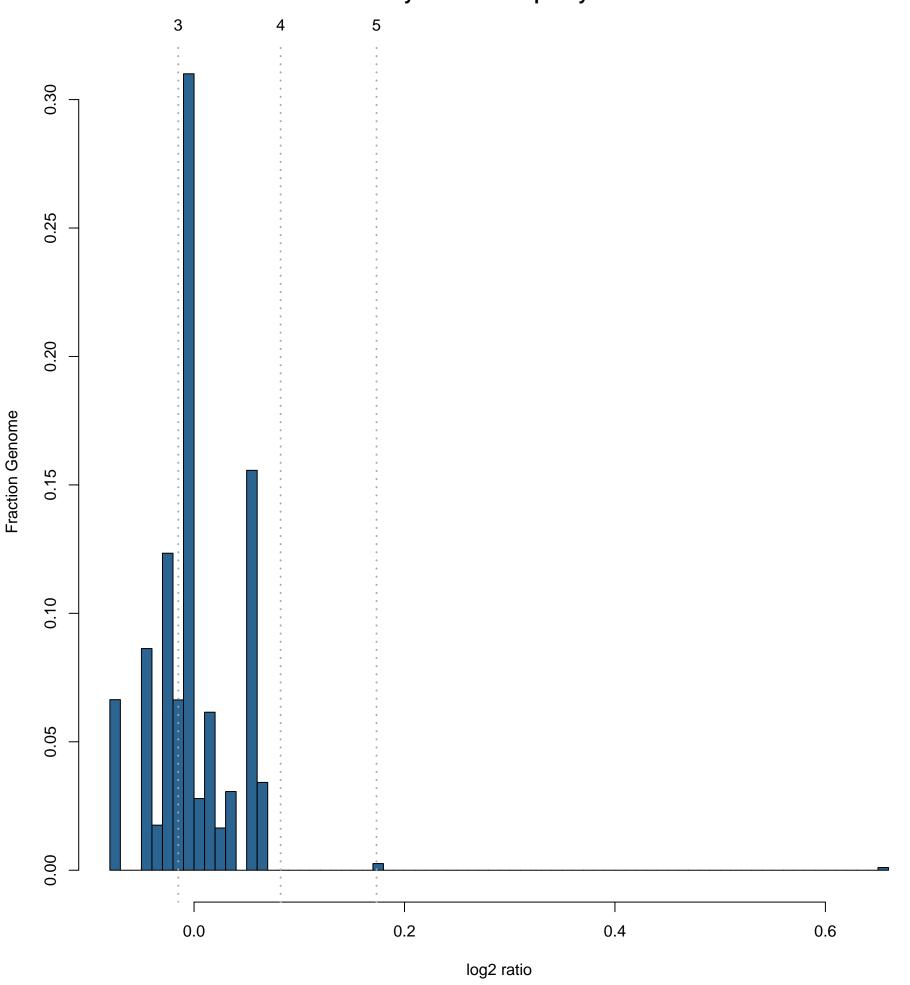
SNV Index

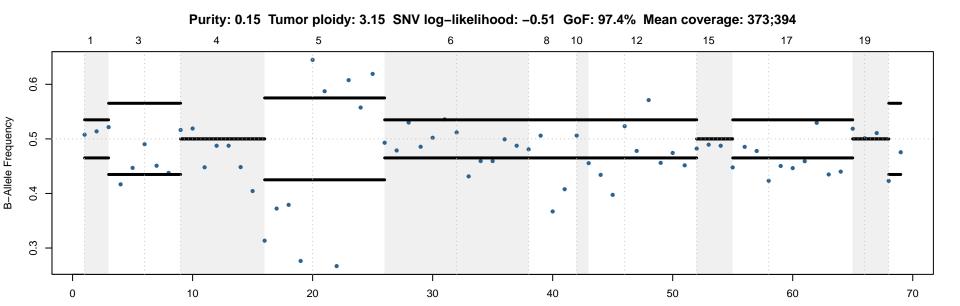






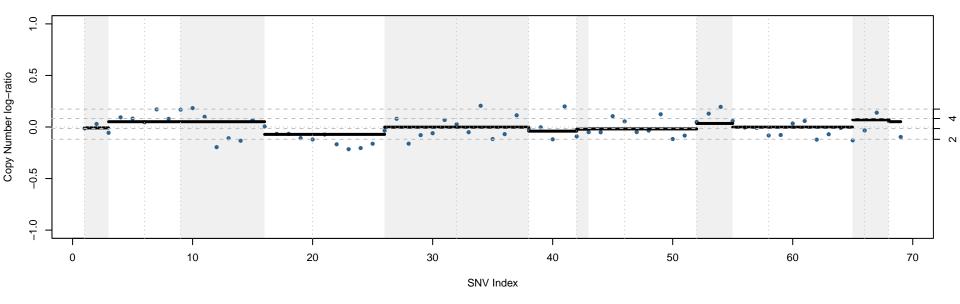
Purity: 0.15 Tumor ploidy: 3.15

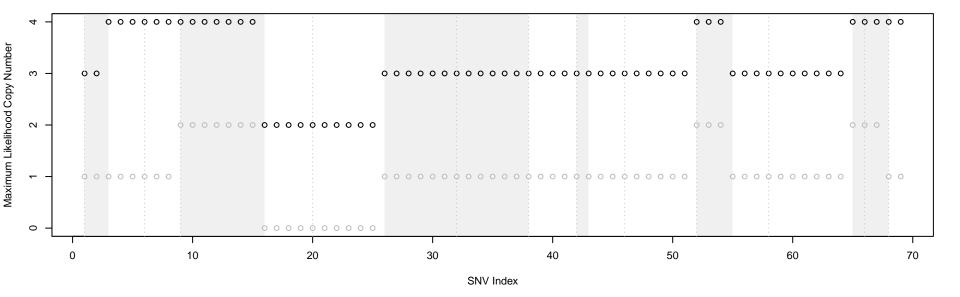


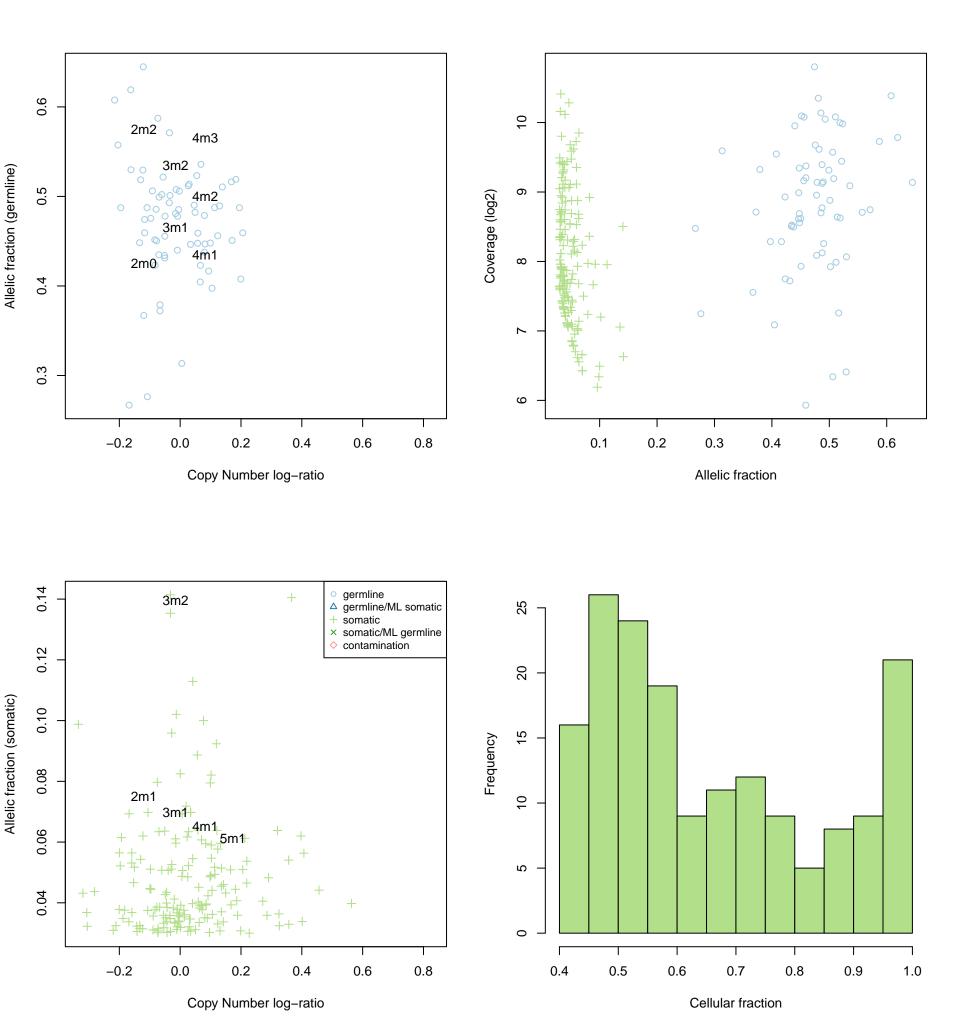


SCNA-fit log-likelihood: -1059.98

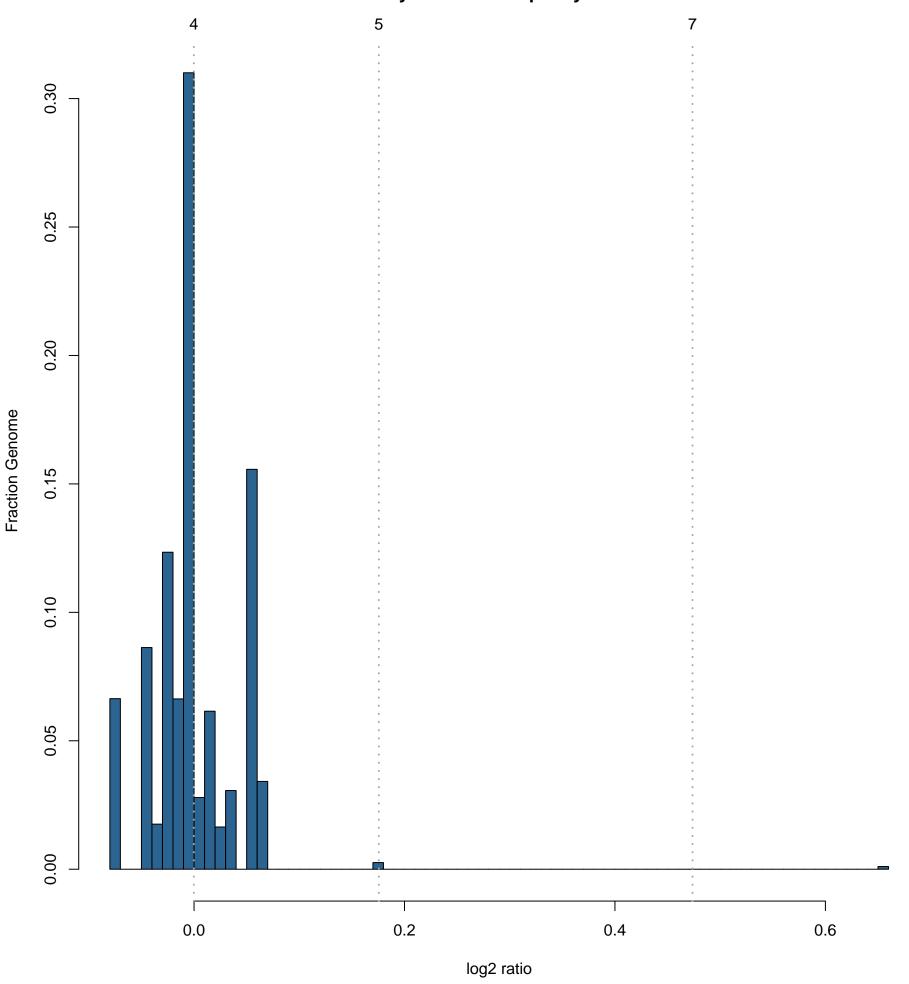
SNV Index



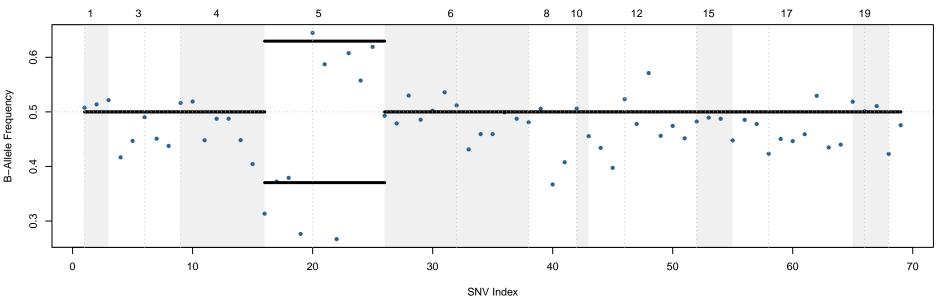




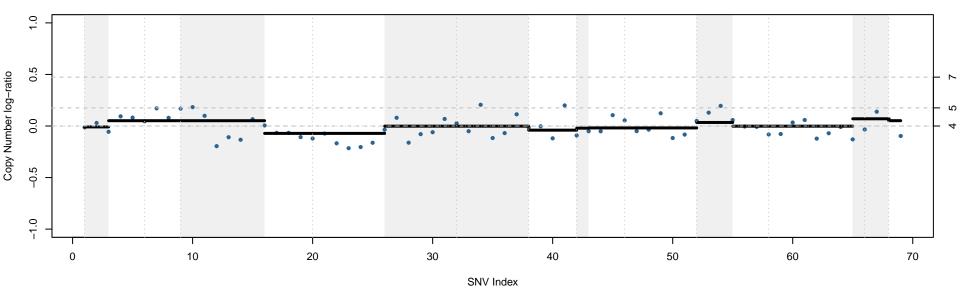
Purity: 0.35 Tumor ploidy: 4.001

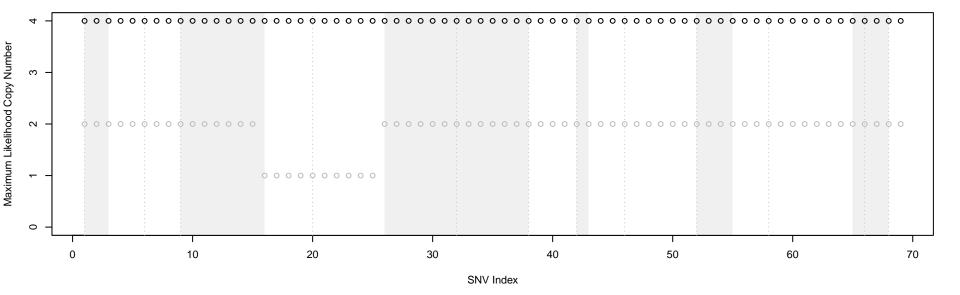


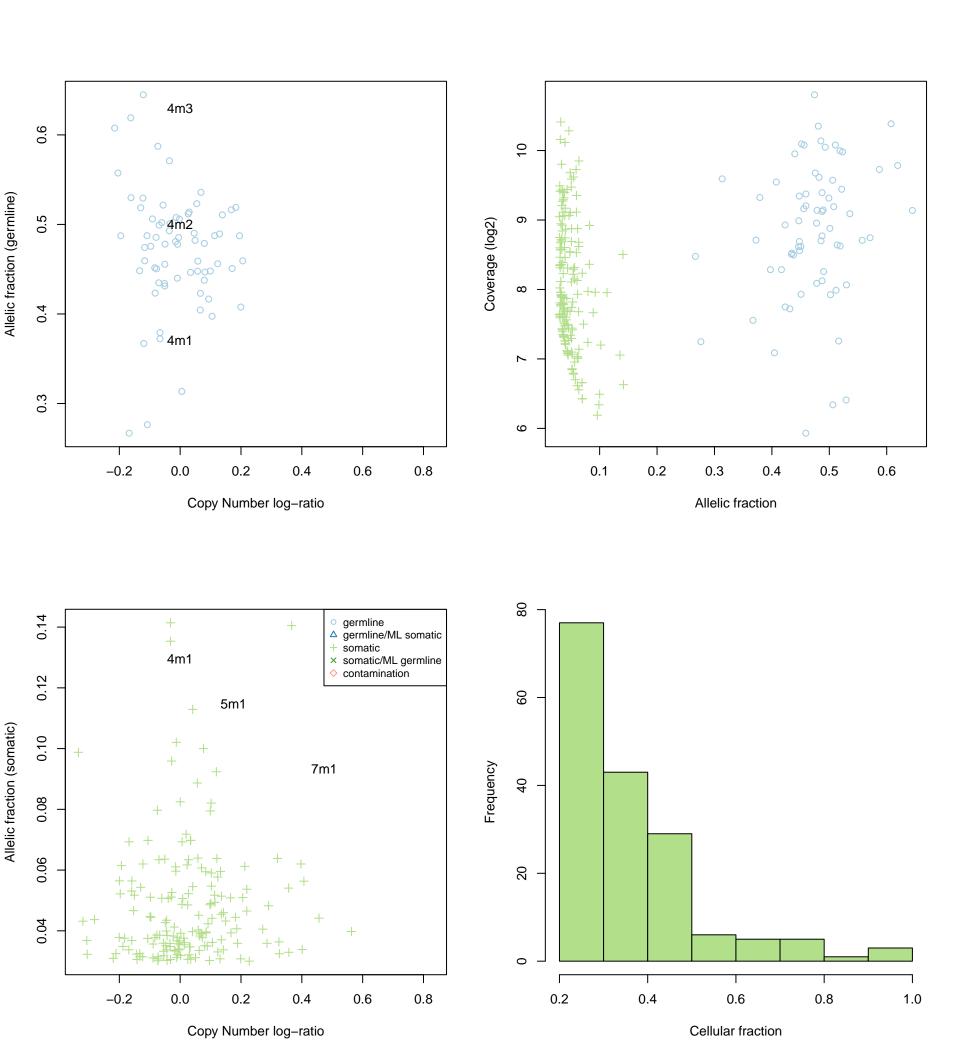




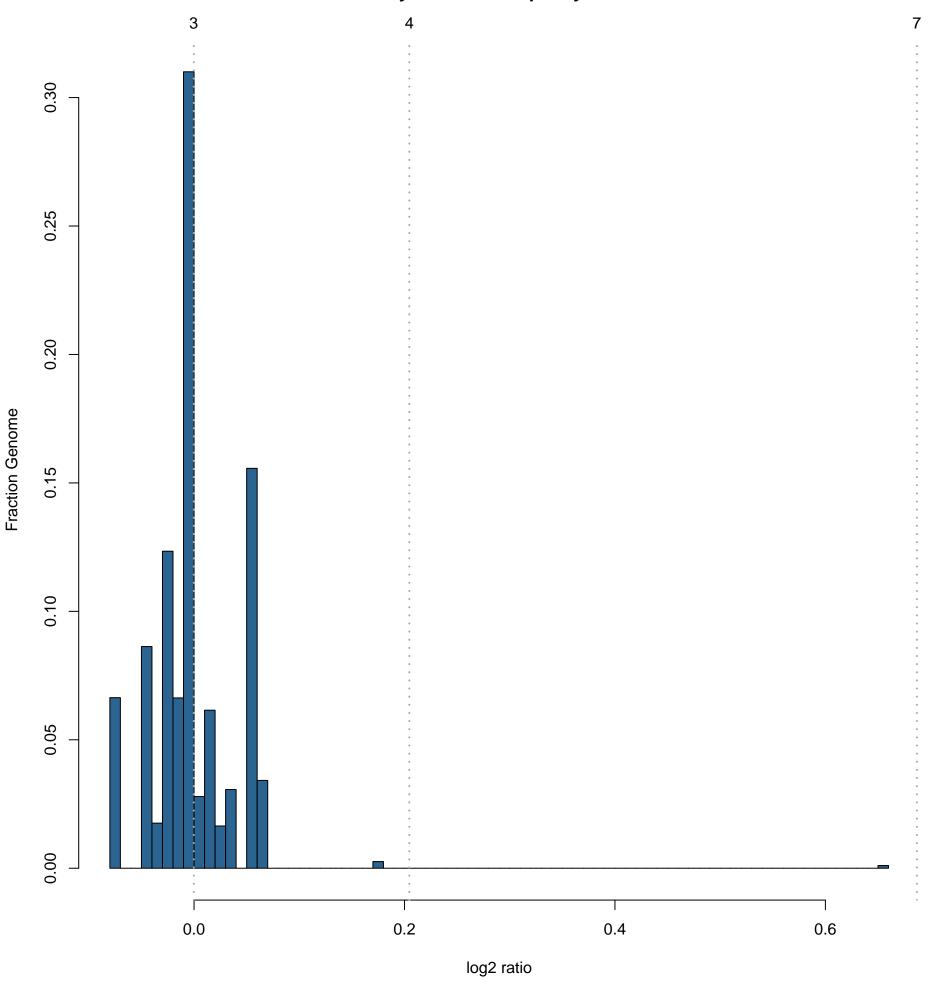
SCNA-fit log-likelihood: -1137.58

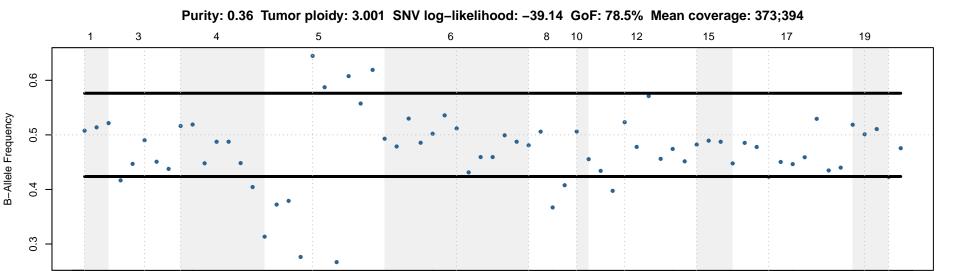






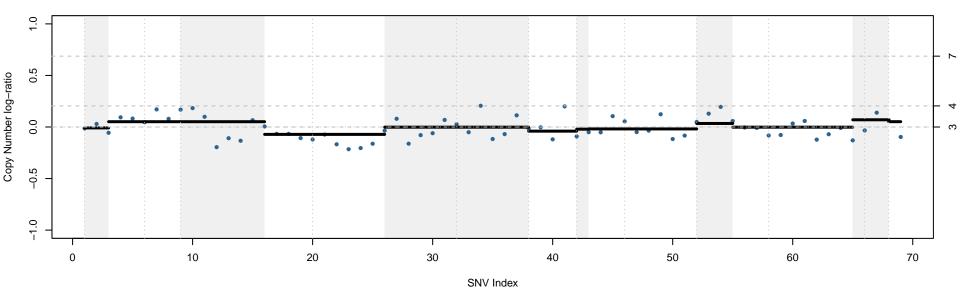
Purity: 0.36 Tumor ploidy: 3.001

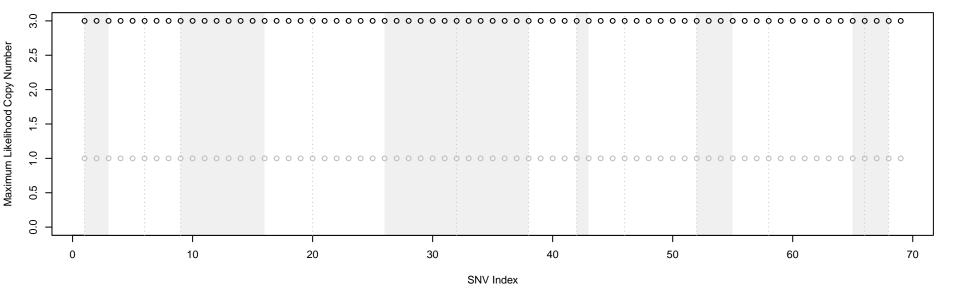


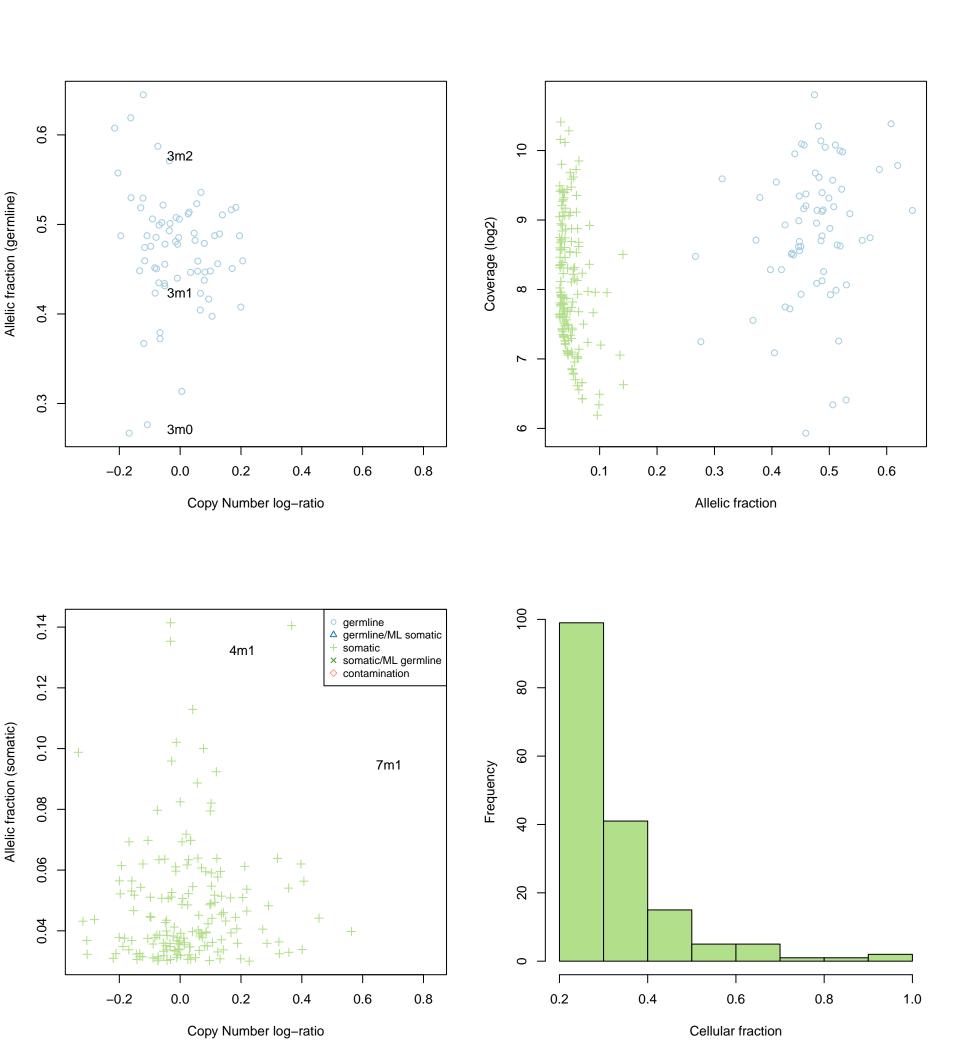


SCNA-fit log-likelihood: -1132.15

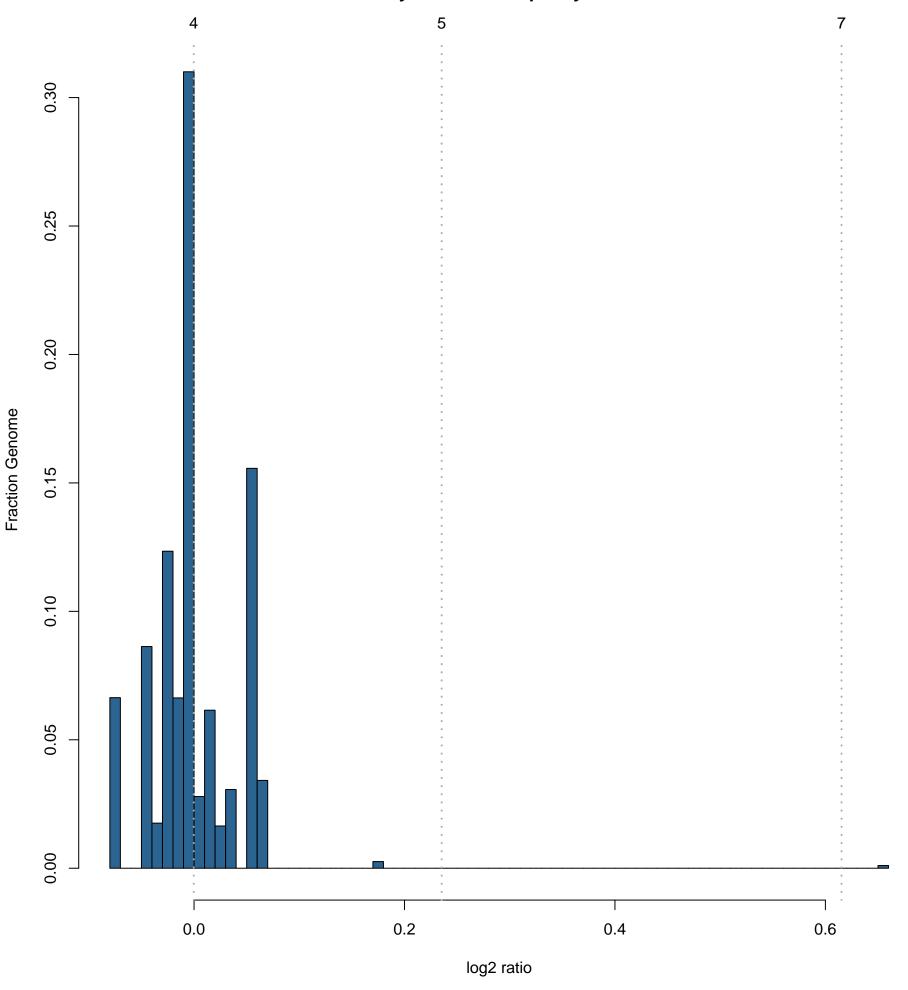
SNV Index

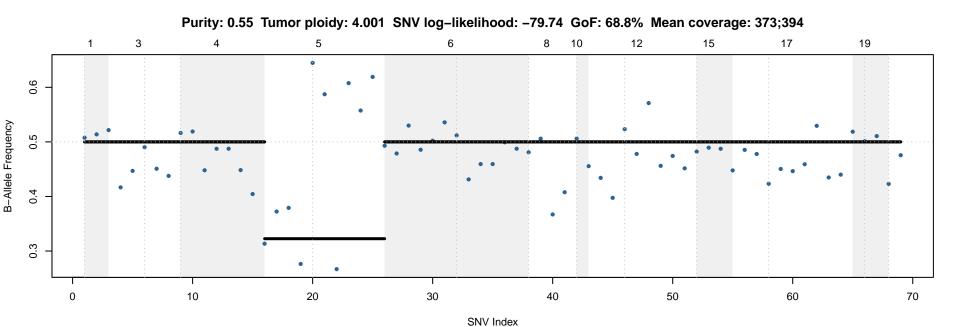




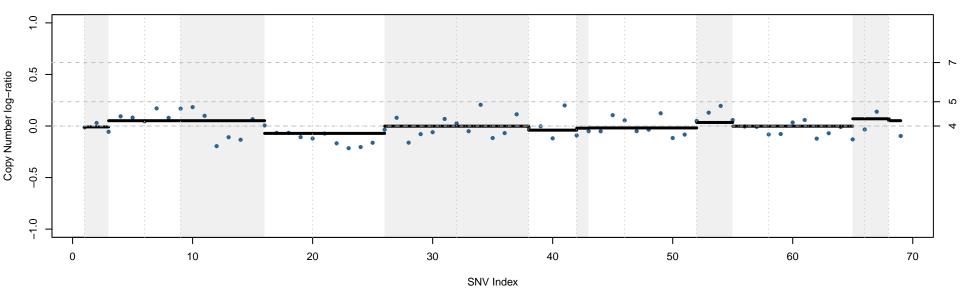


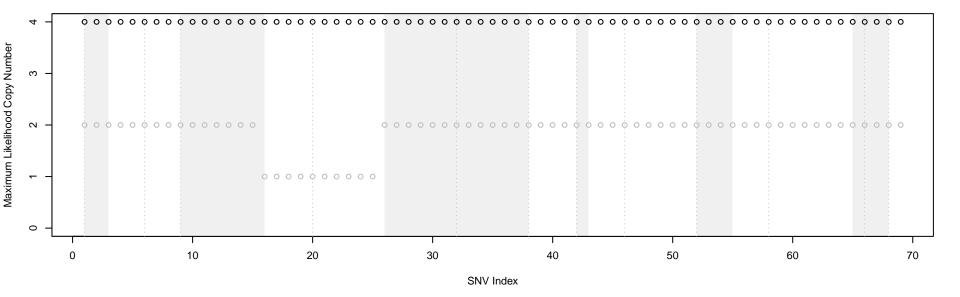
Purity: 0.55 Tumor ploidy: 4.001

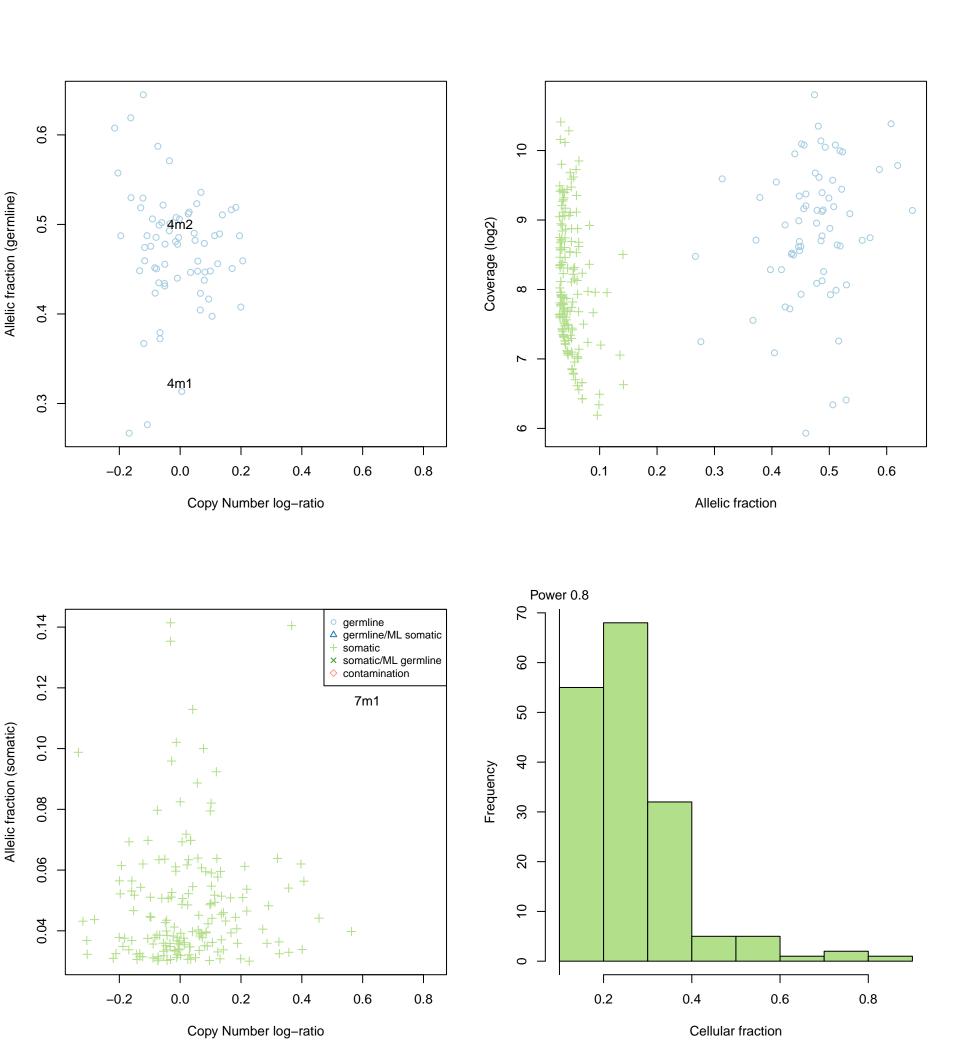




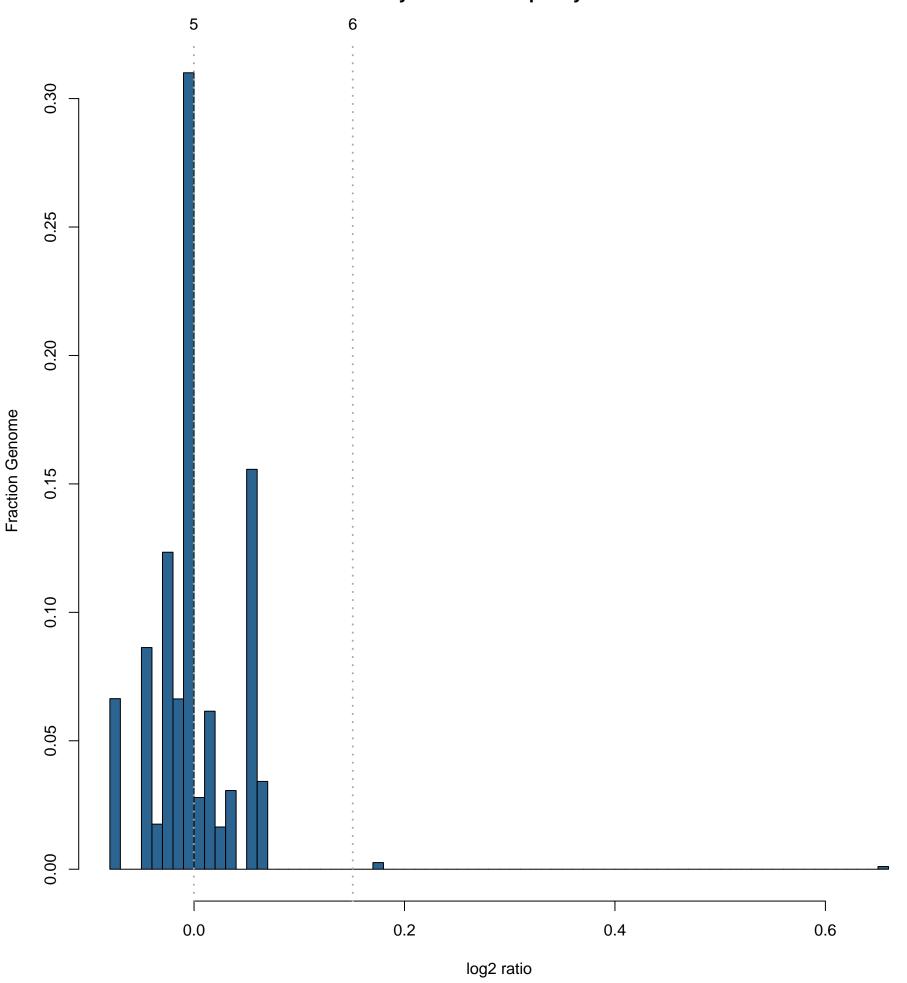
SCNA-fit log-likelihood: -1133.62



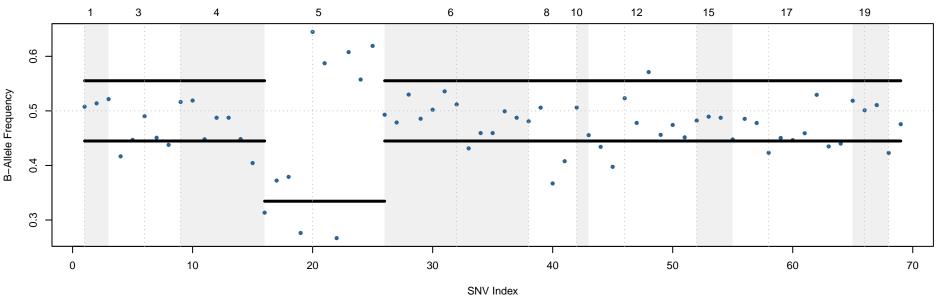




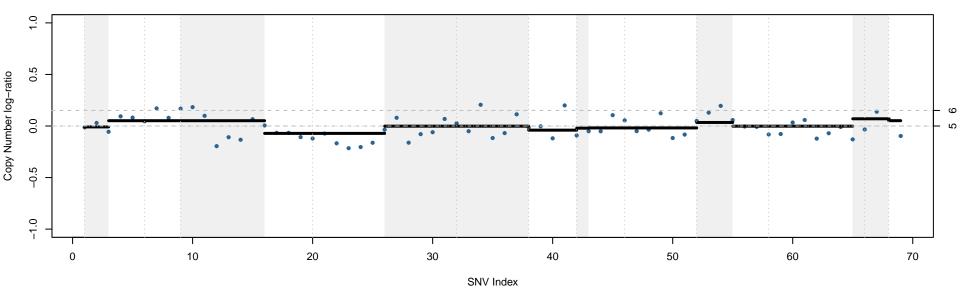
Purity: 0.33 Tumor ploidy: 5.001

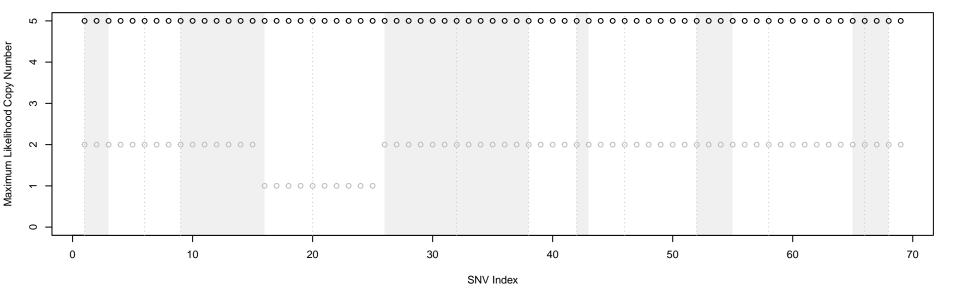


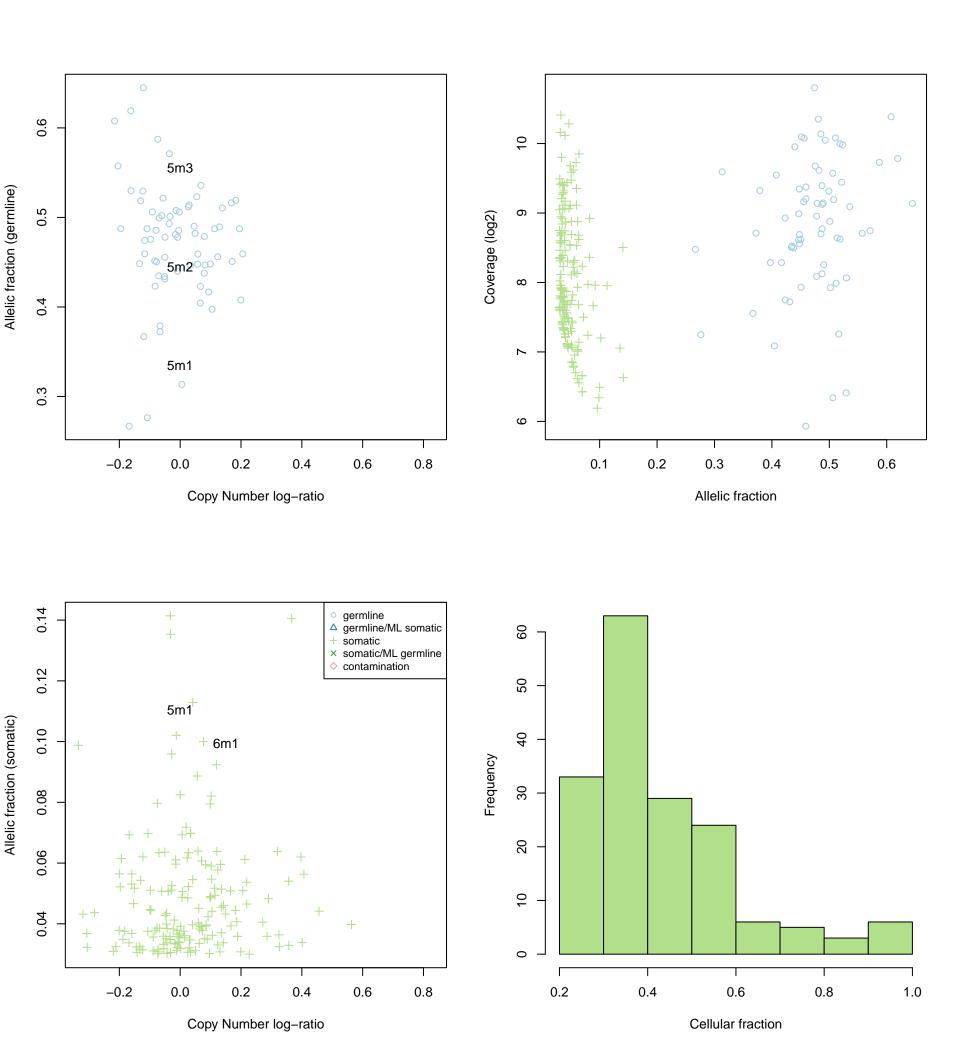




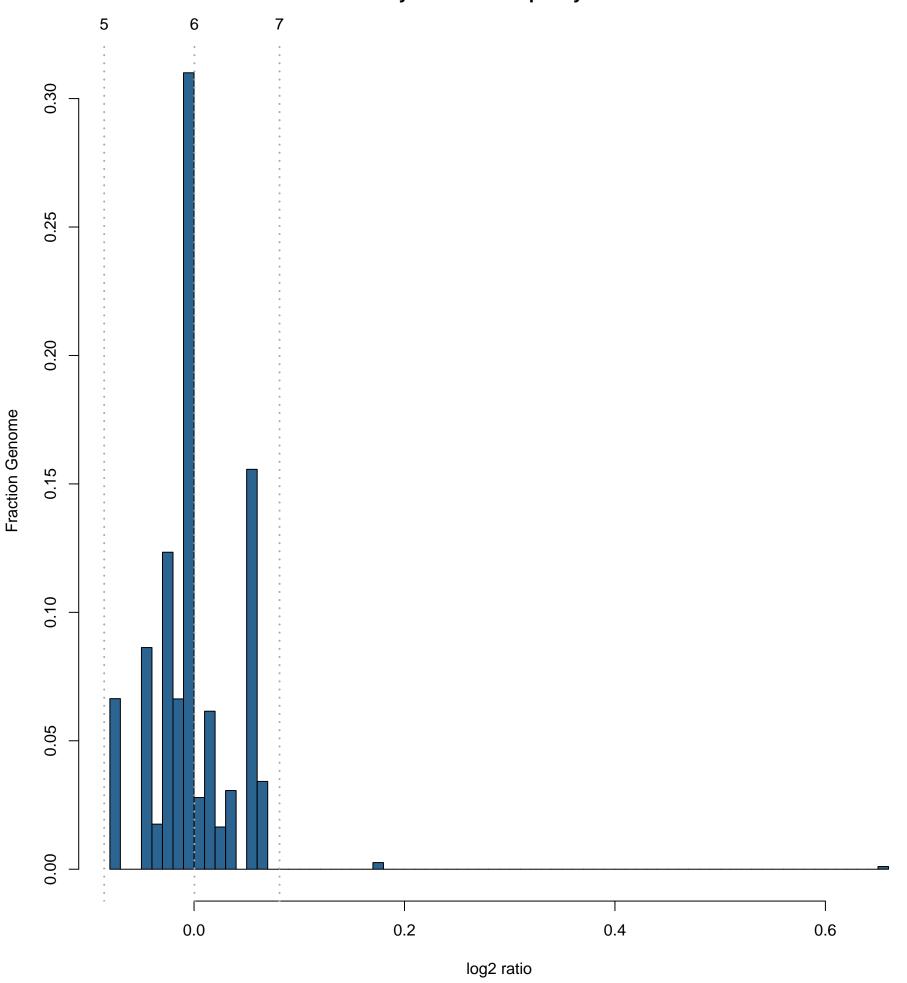
SCNA-fit log-likelihood: -1170.45



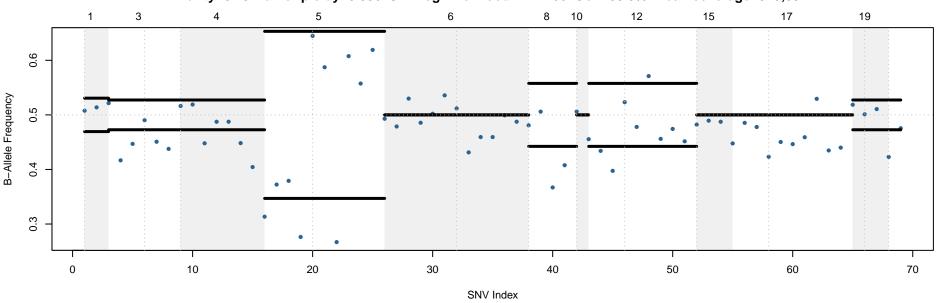




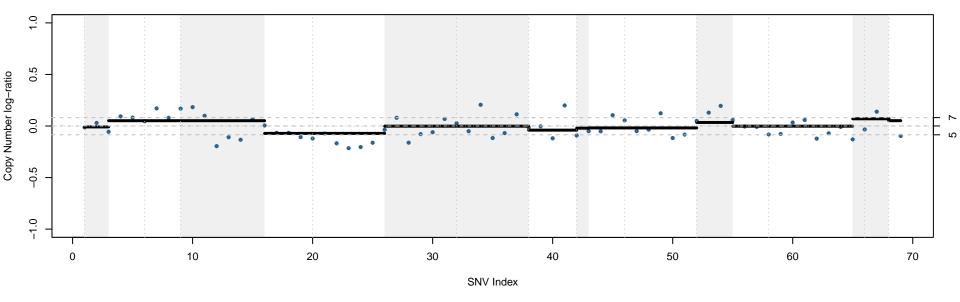
Purity: 0.15 Tumor ploidy: 5.996

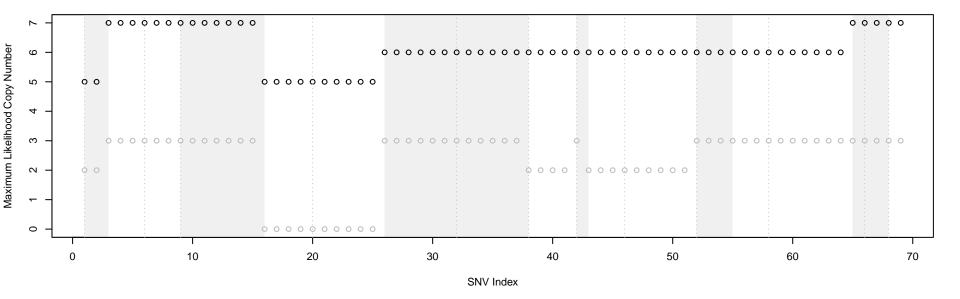


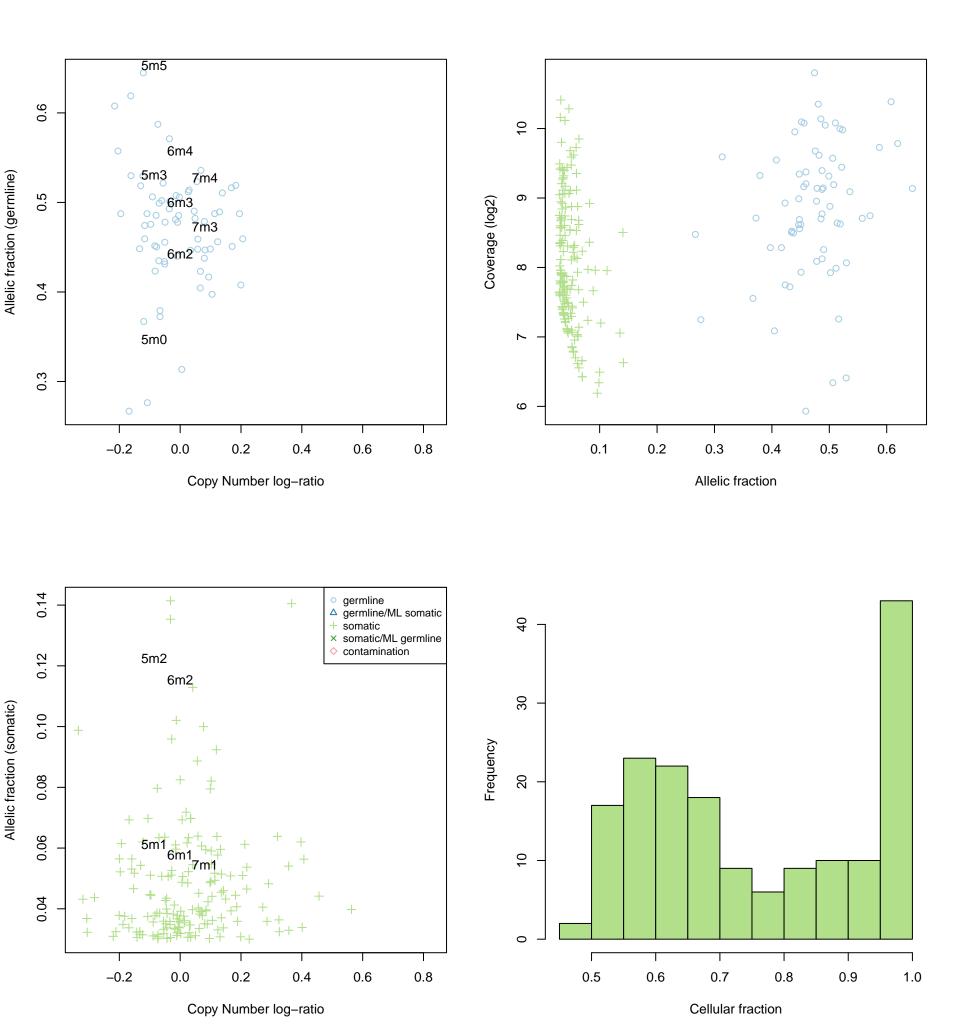
Purity: 0.15 Tumor ploidy: 5.996 SNV log-likelihood: -114.66 GoF: 98.5% Mean coverage: 373;394



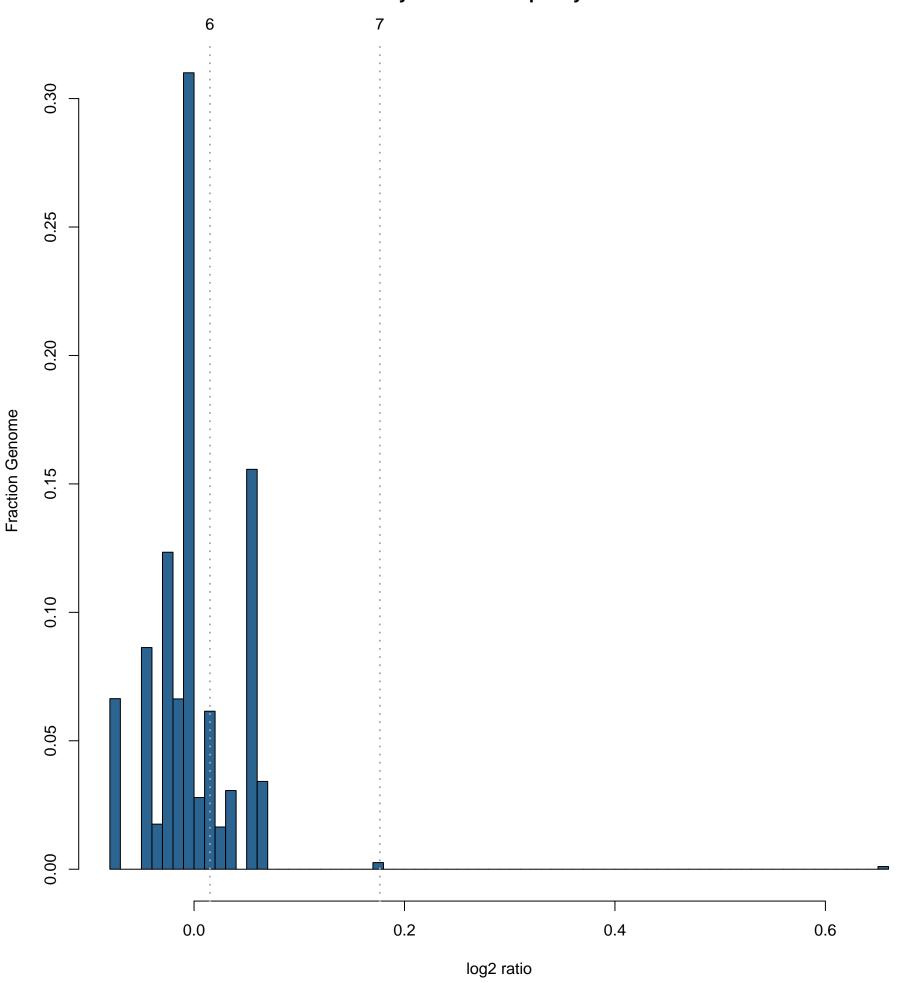
SCNA-fit log-likelihood: -1121.79

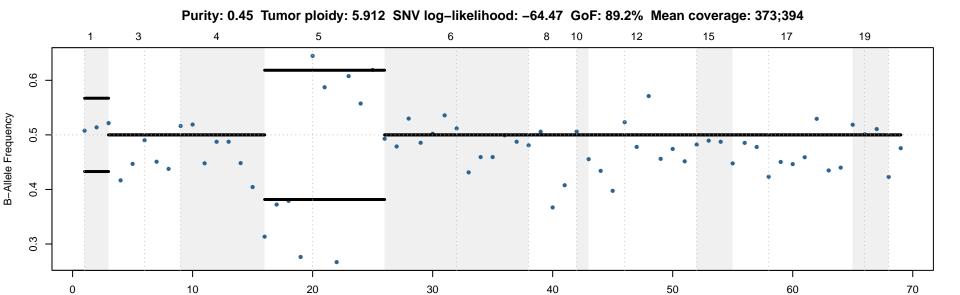






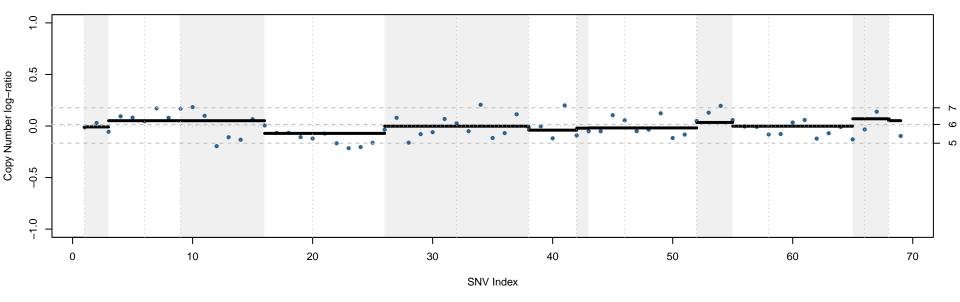
Purity: 0.45 Tumor ploidy: 5.912

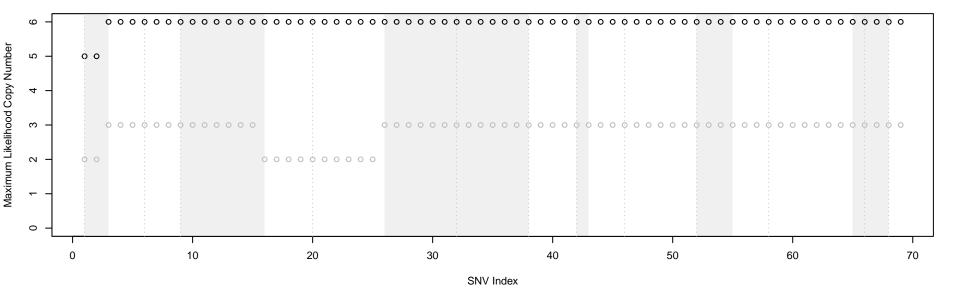


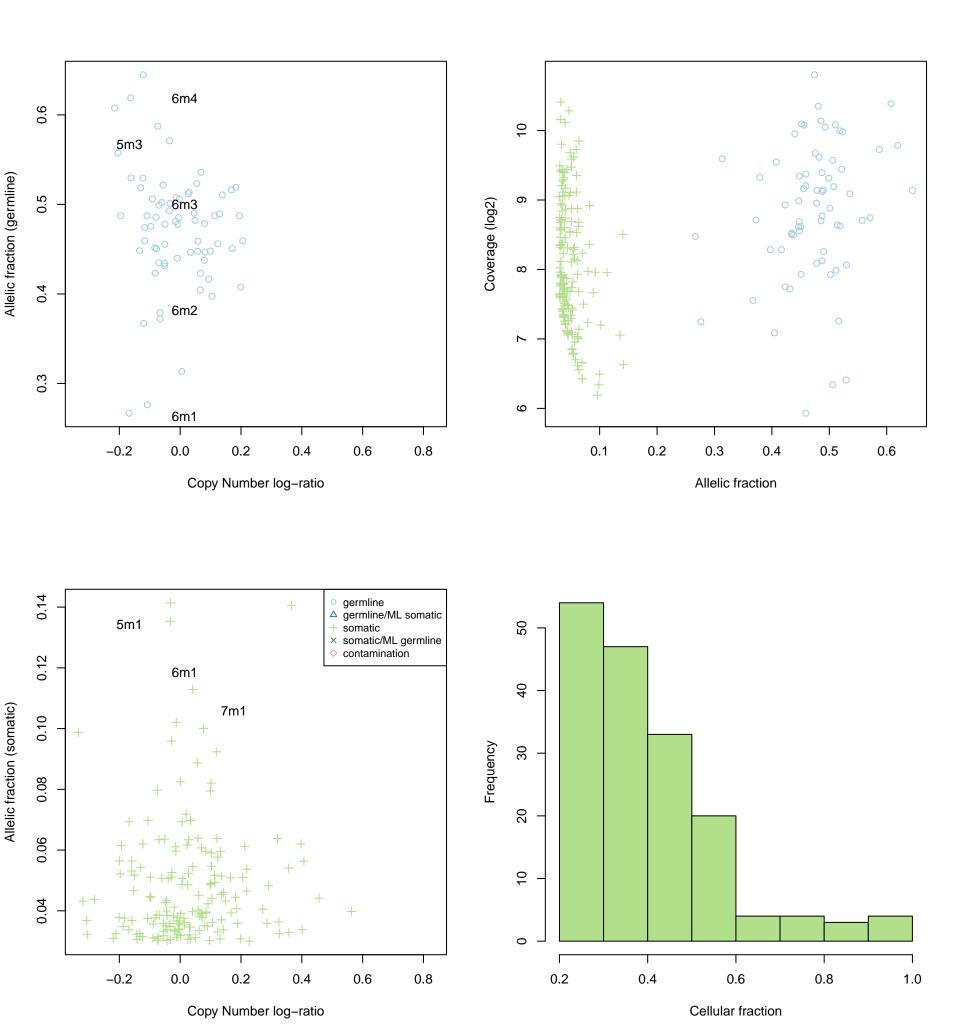


SCNA-fit log-likelihood: -1531.37

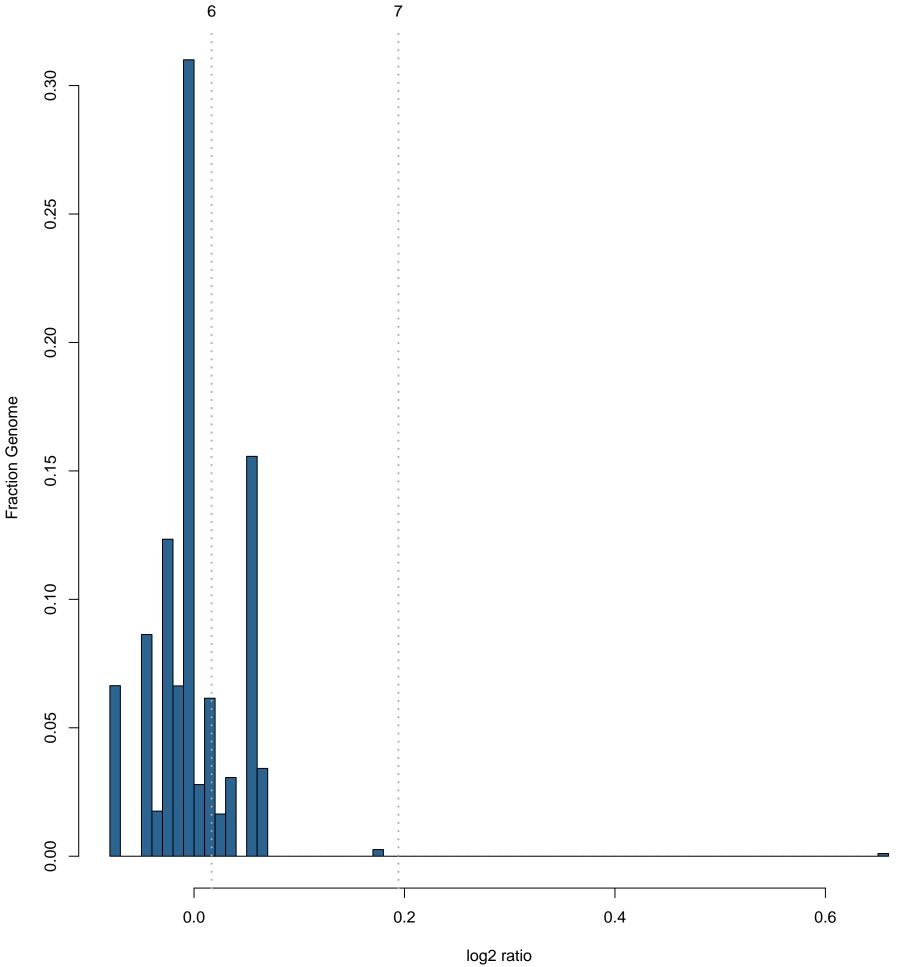
SNV Index

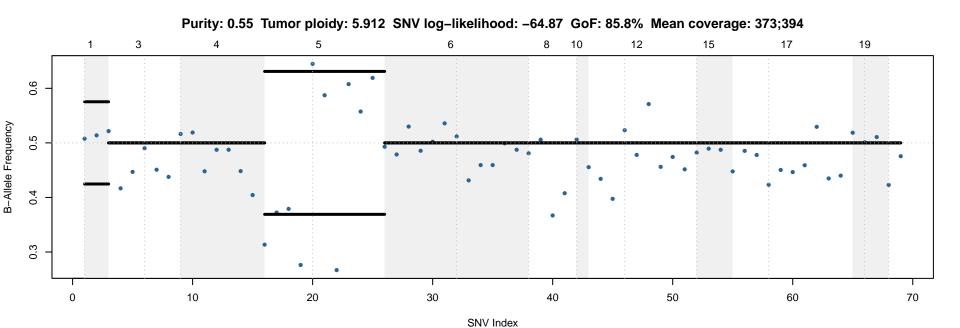




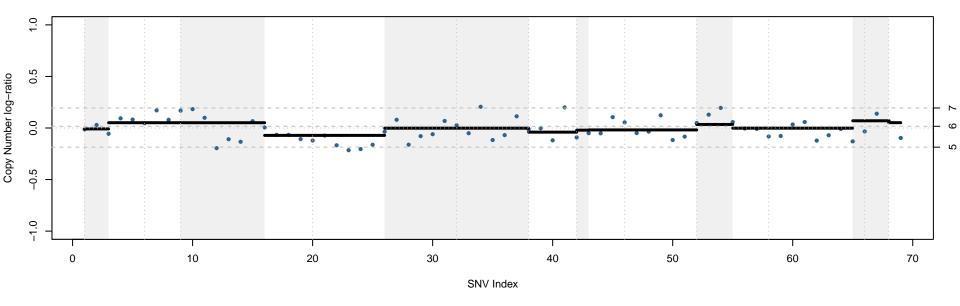


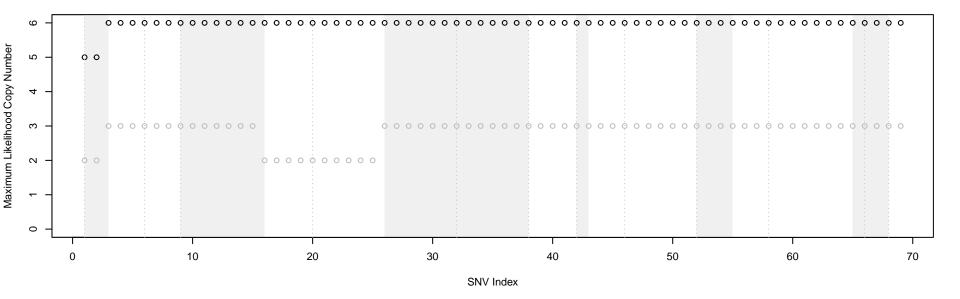
Purity: 0.55 Tumor ploidy: 5.912

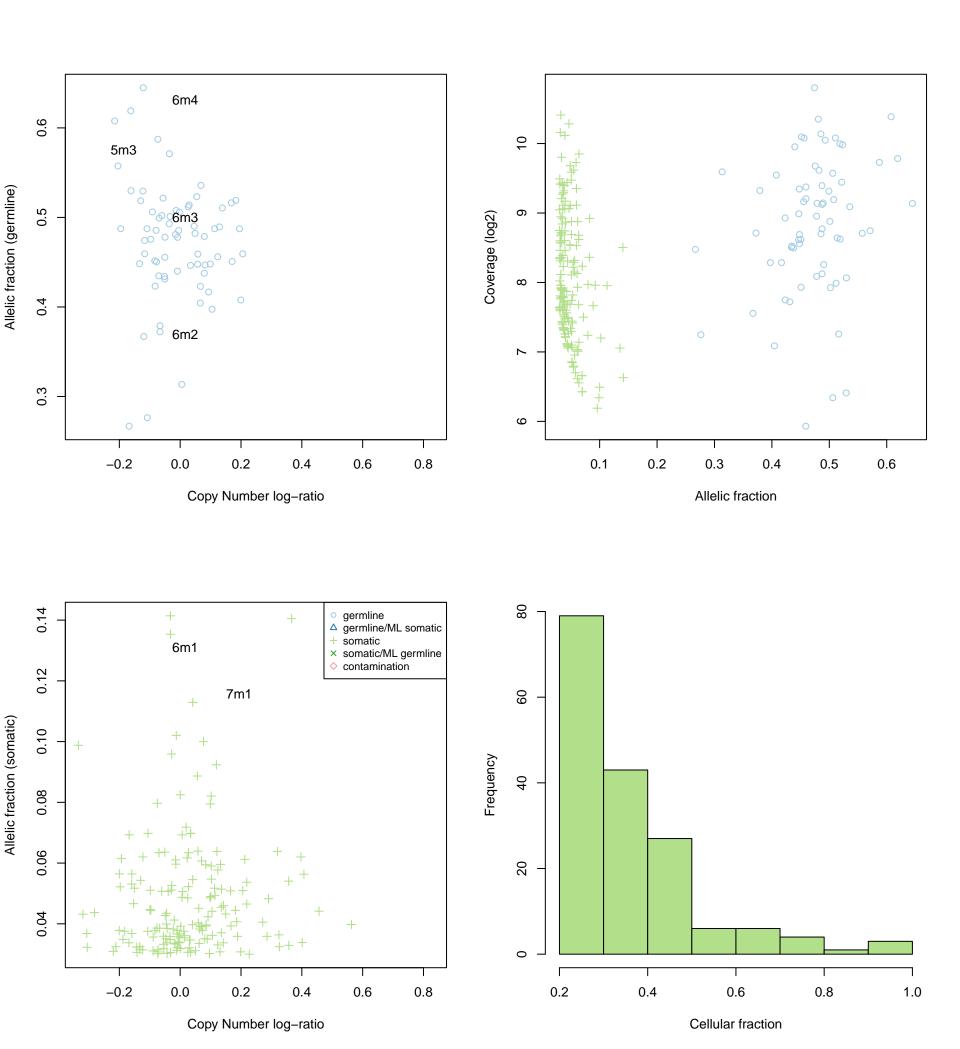




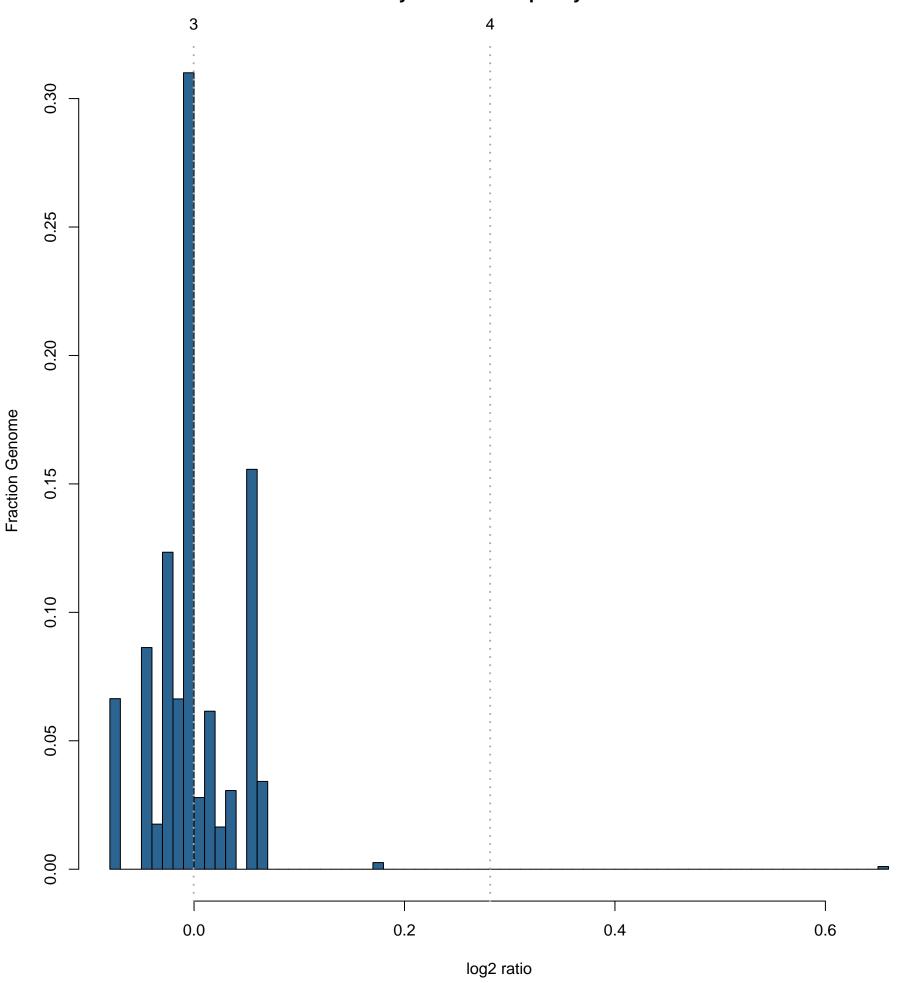
SCNA-fit log-likelihood: -1615.68



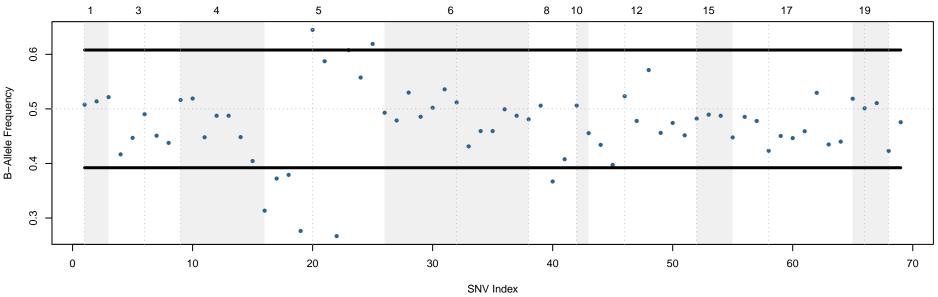




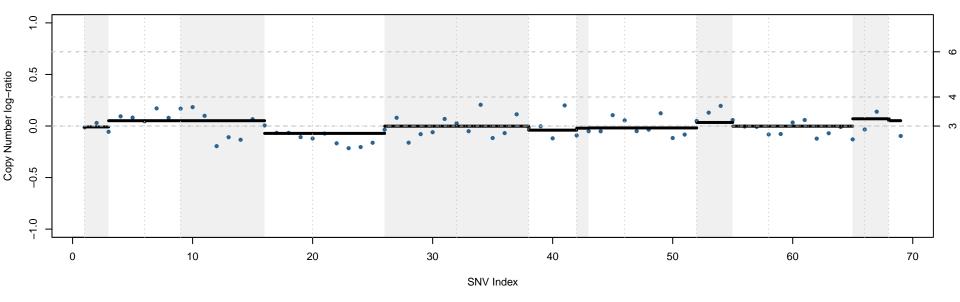
Purity: 0.55 Tumor ploidy: 3.001

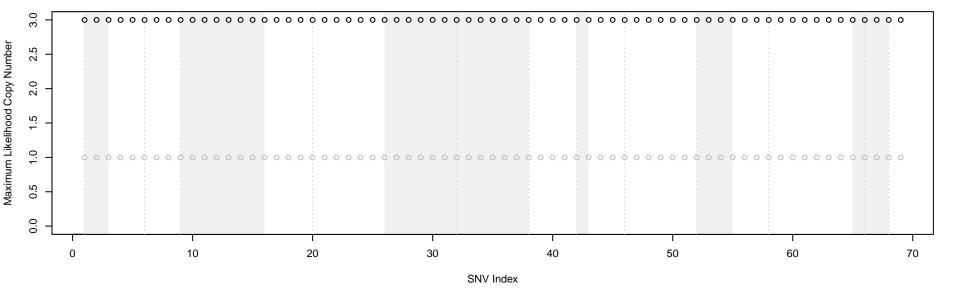


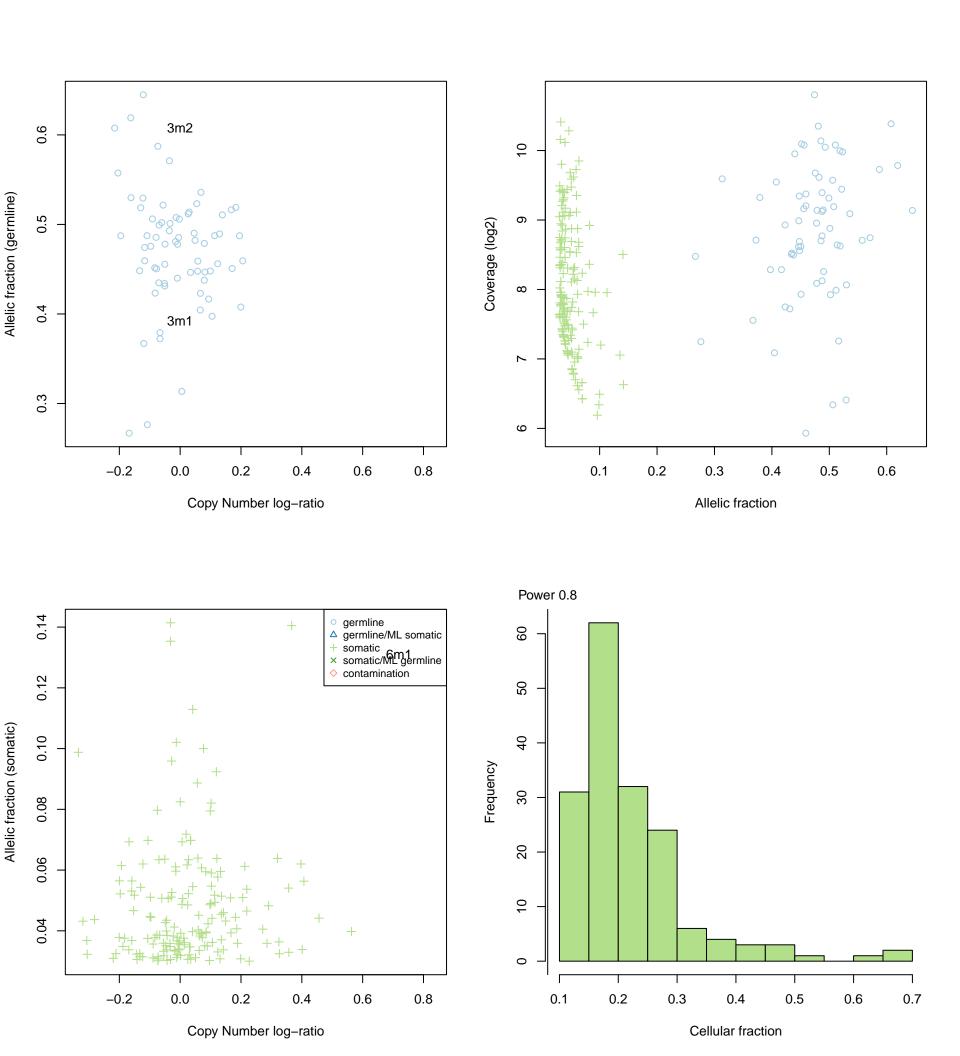




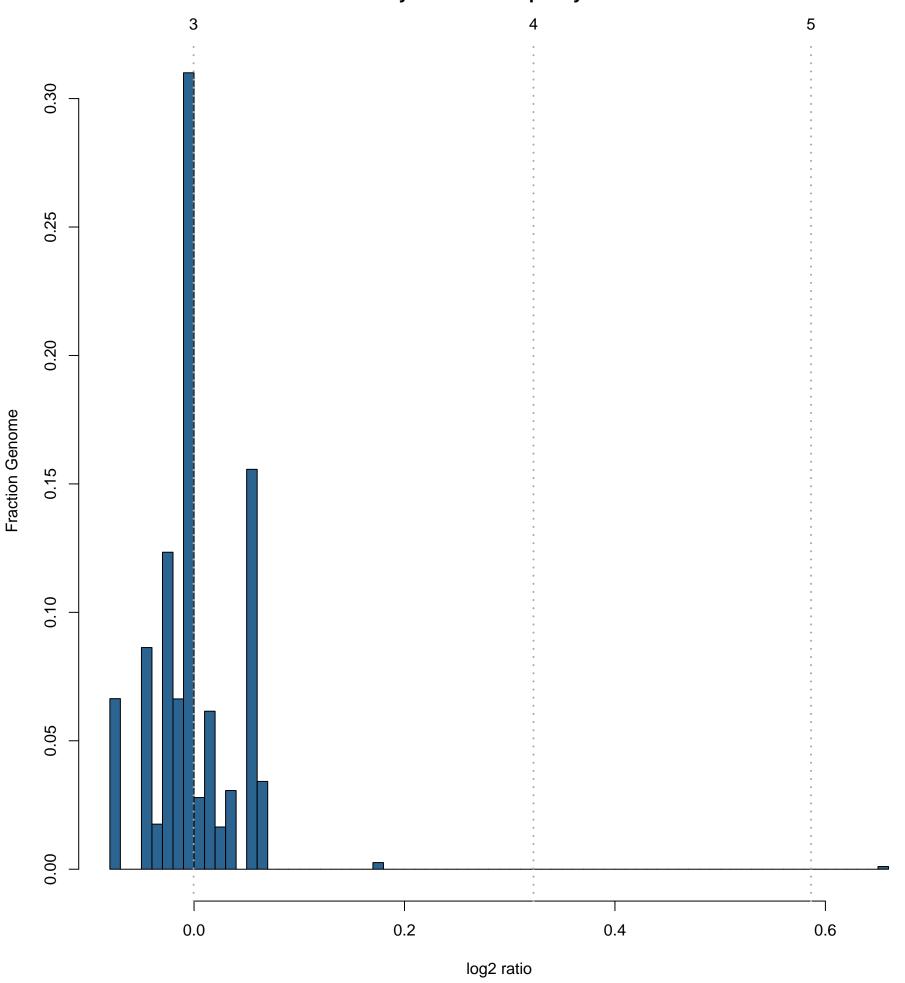
SCNA-fit log-likelihood: -1137.07



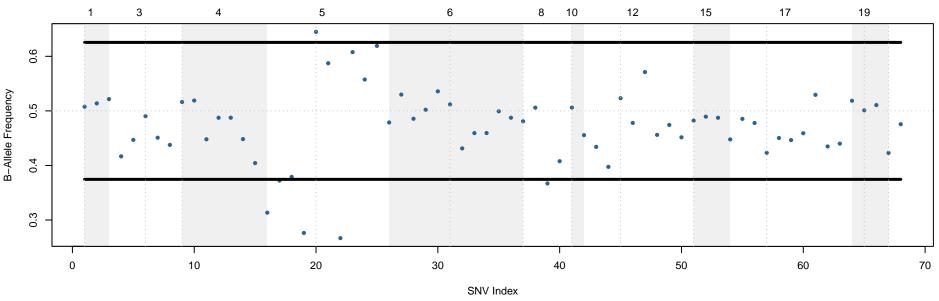




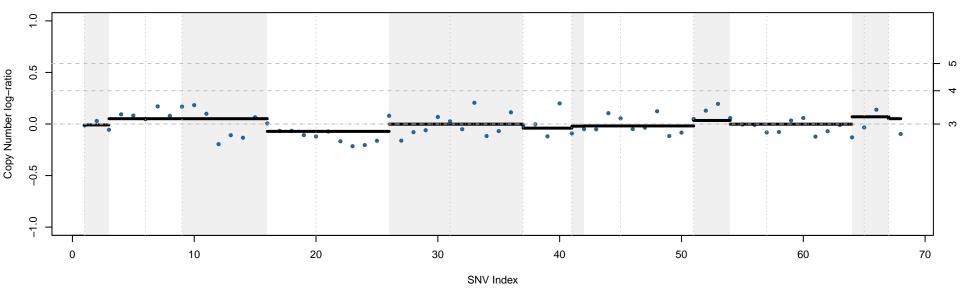
Purity: 0.67 Tumor ploidy: 3.001

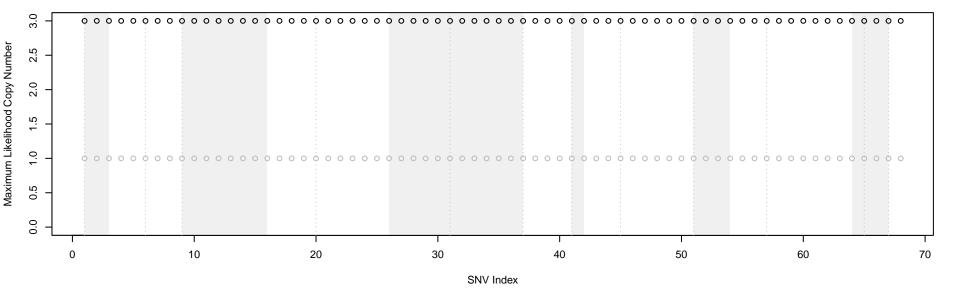


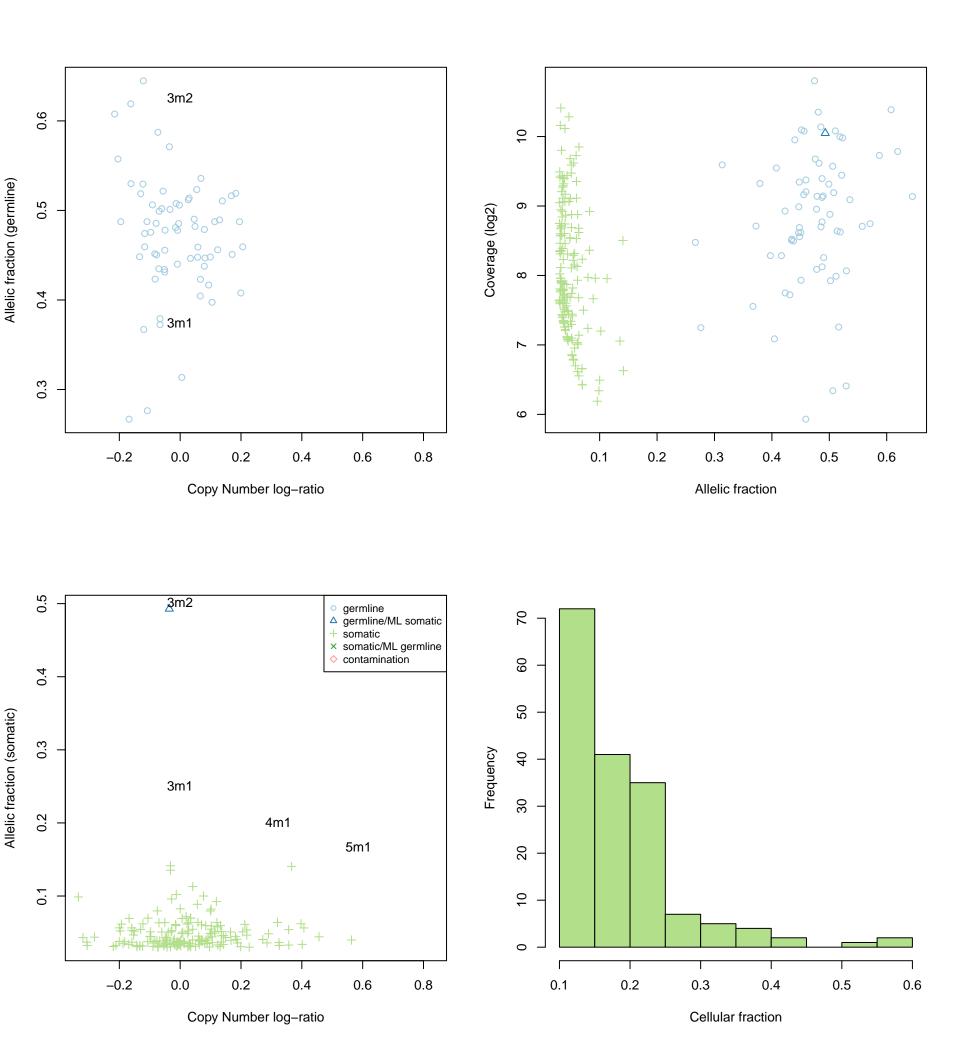




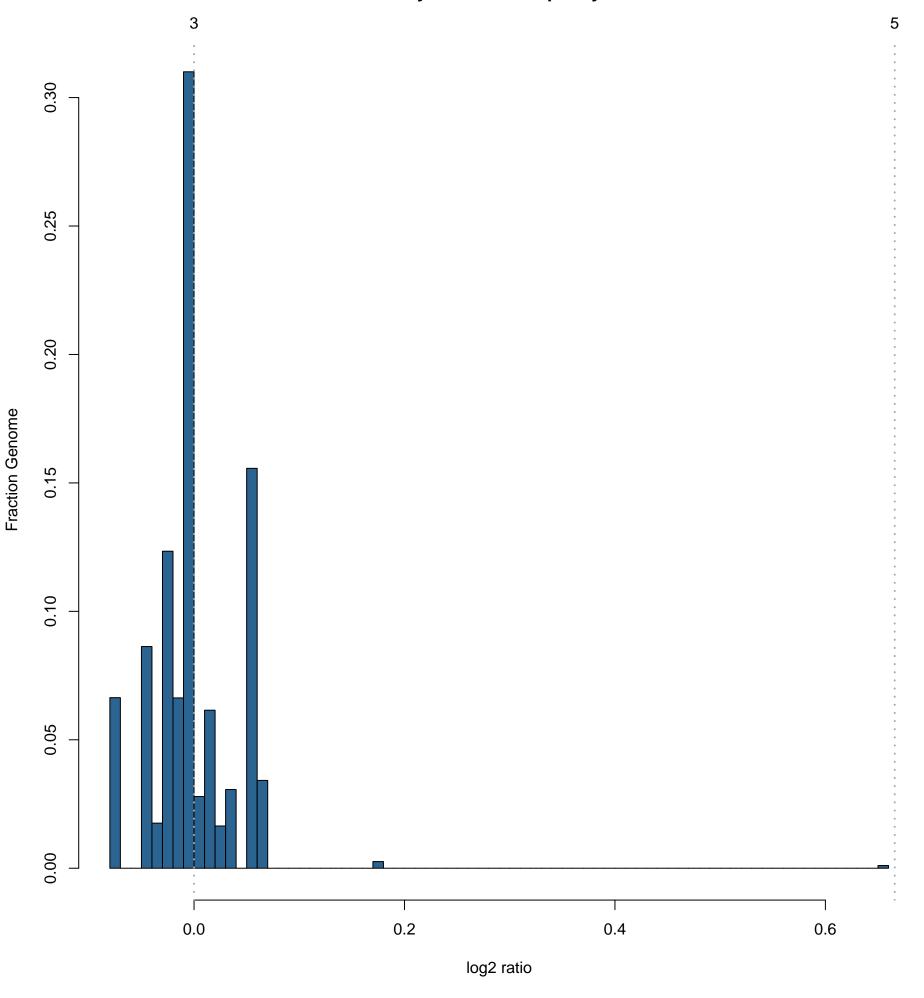
SCNA-fit log-likelihood: -1141.77

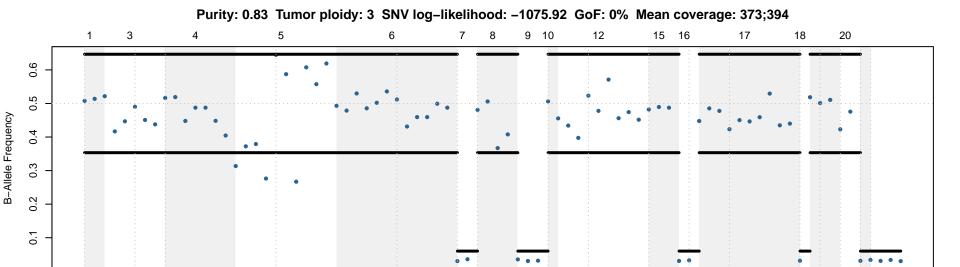






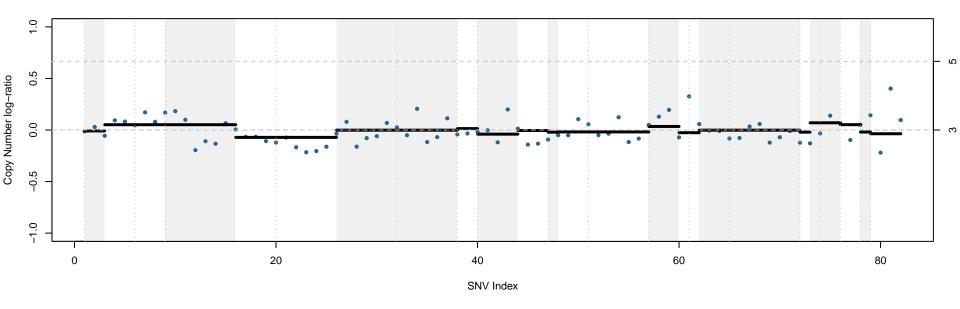
Purity: 0.83 Tumor ploidy: 3

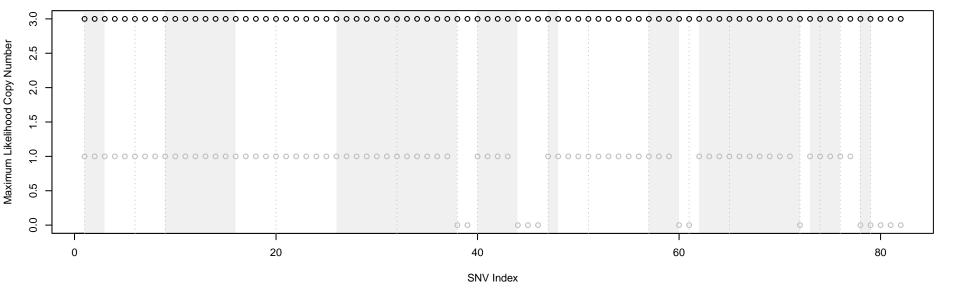


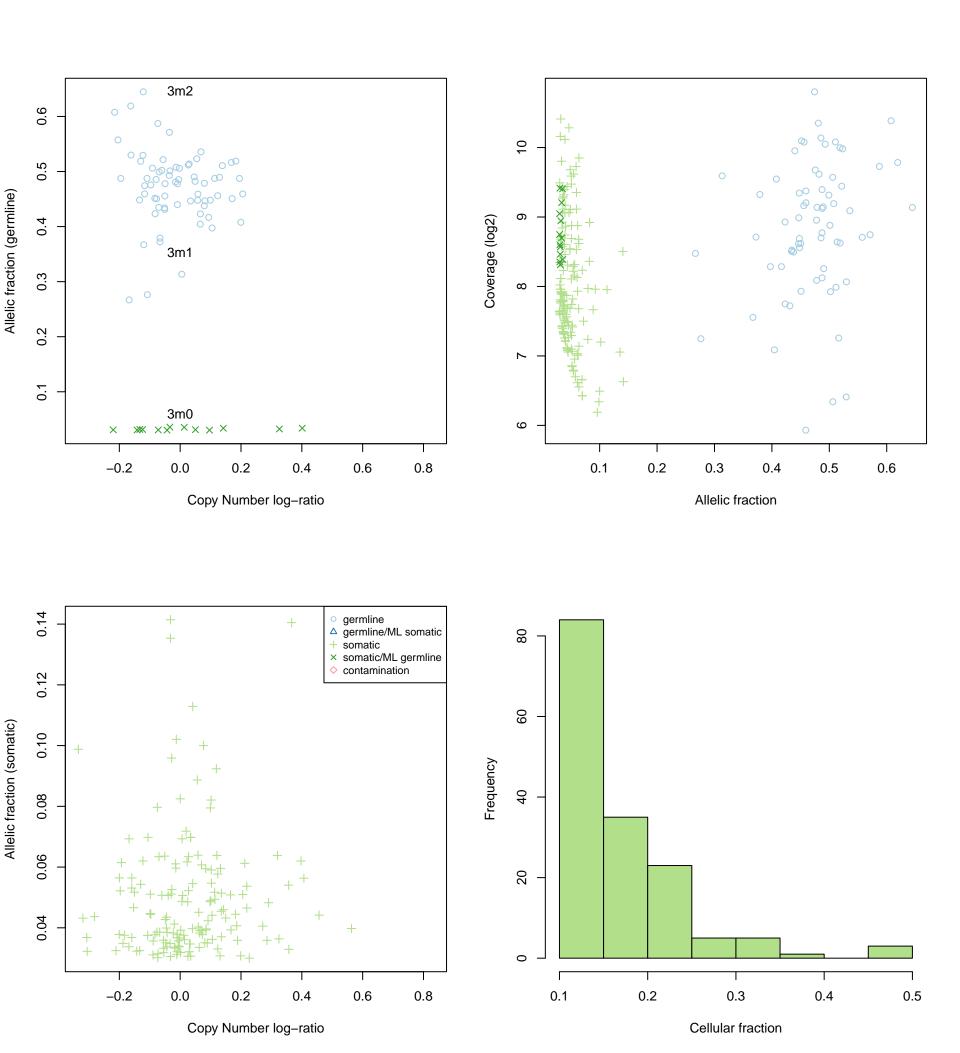


SCNA-fit log-likelihood: -1145.79

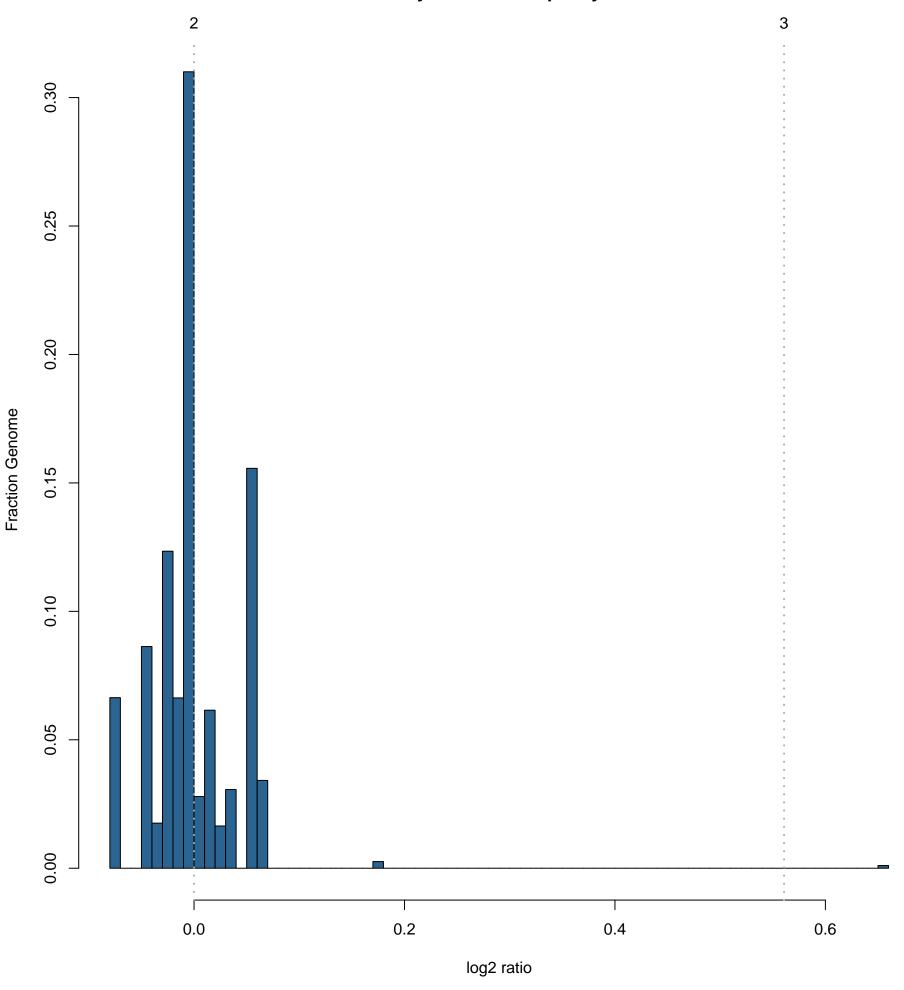
SNV Index

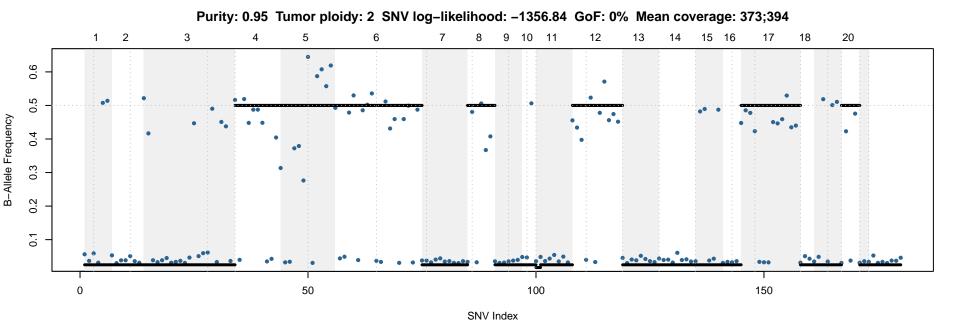






Purity: 0.95 Tumor ploidy: 2





SCNA-fit log-likelihood: -1147.57

