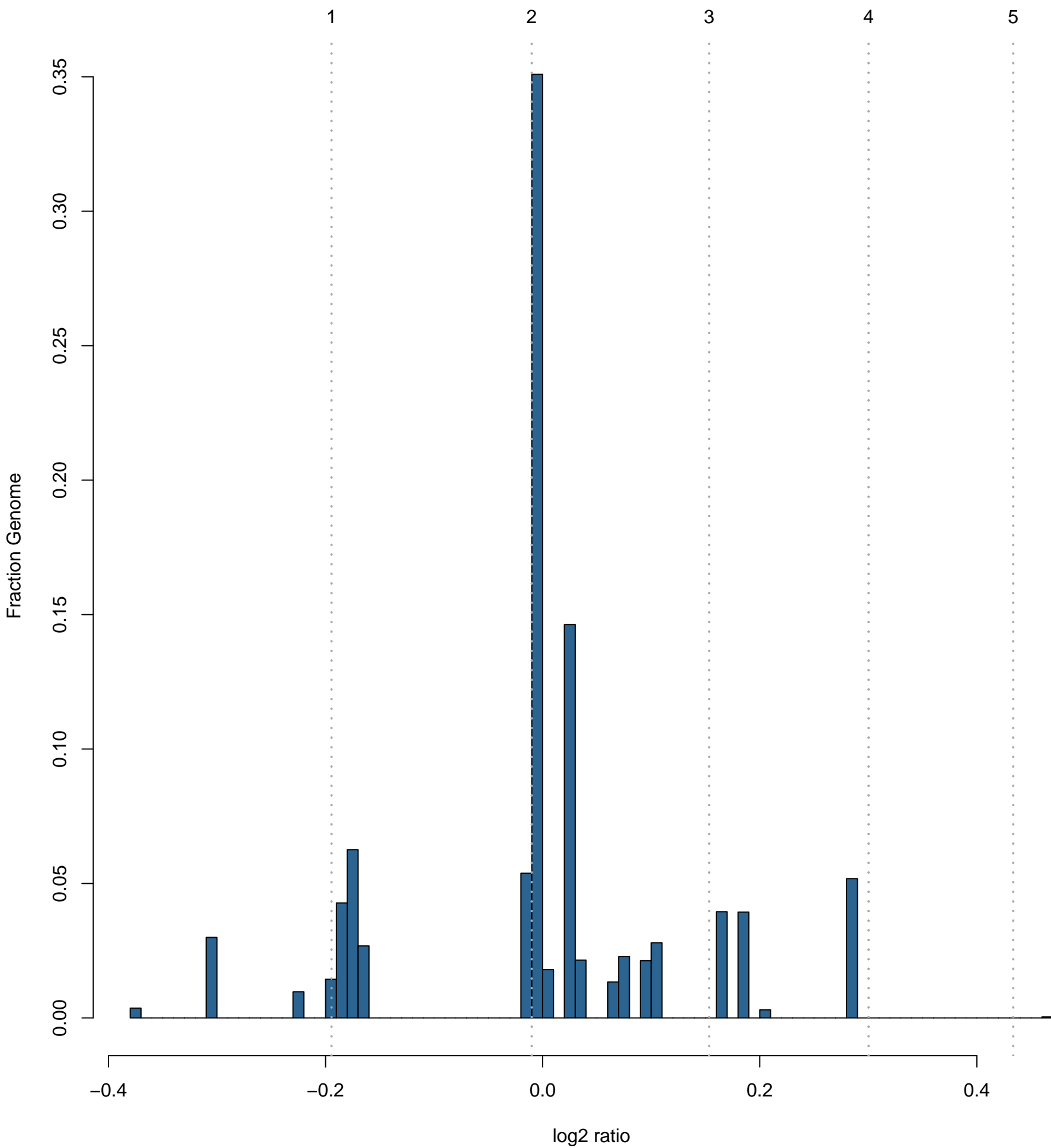
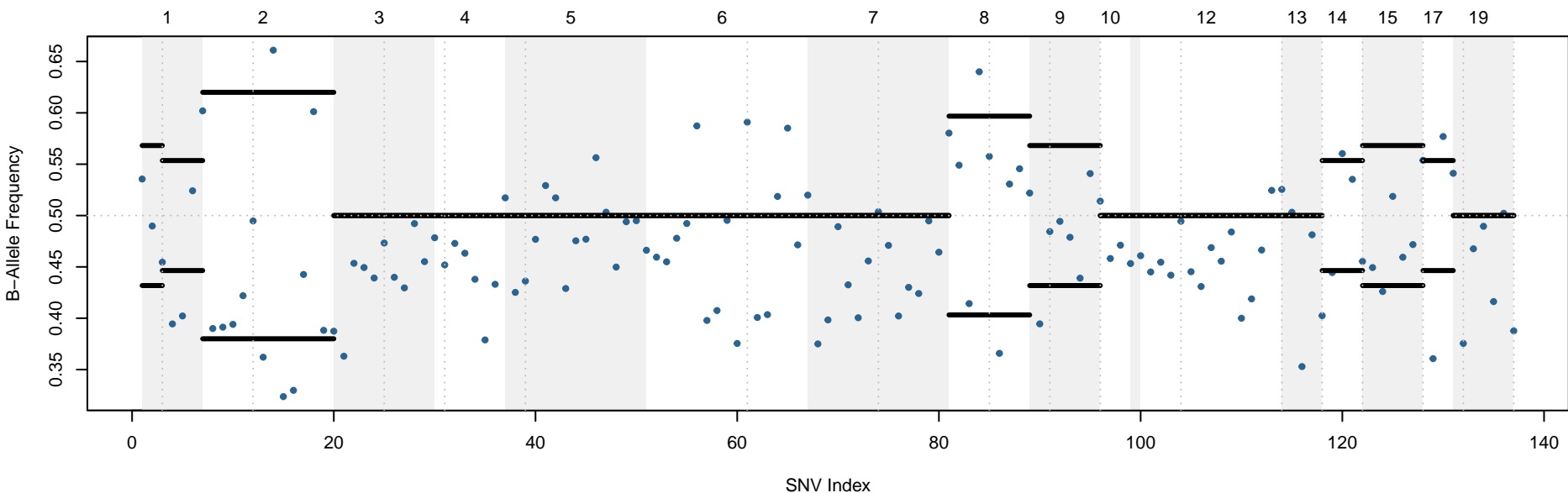


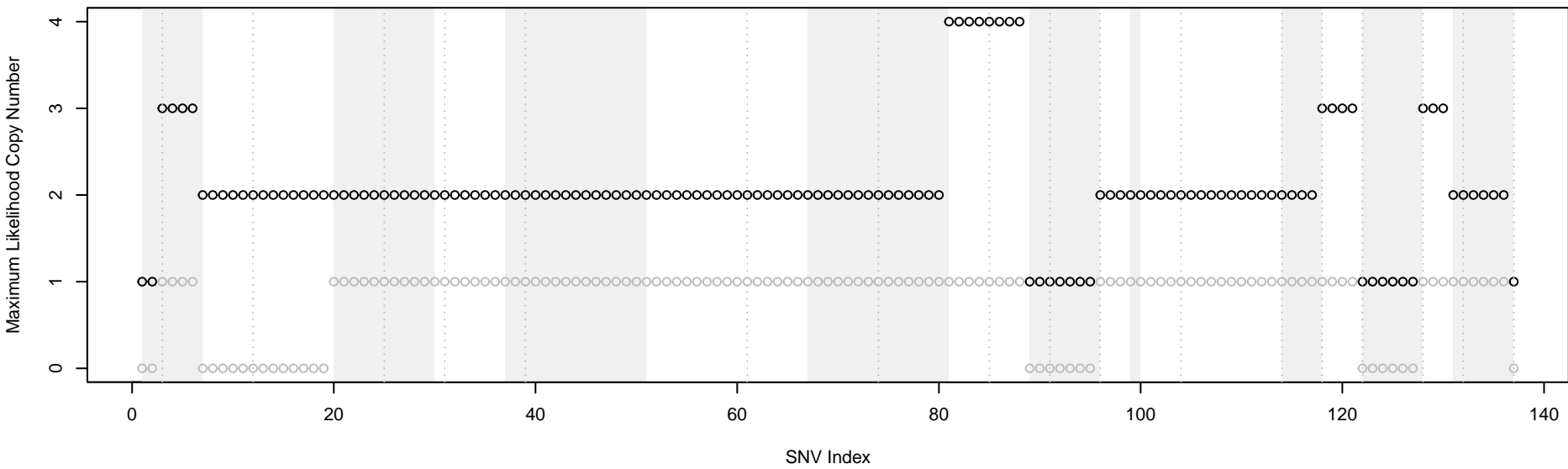
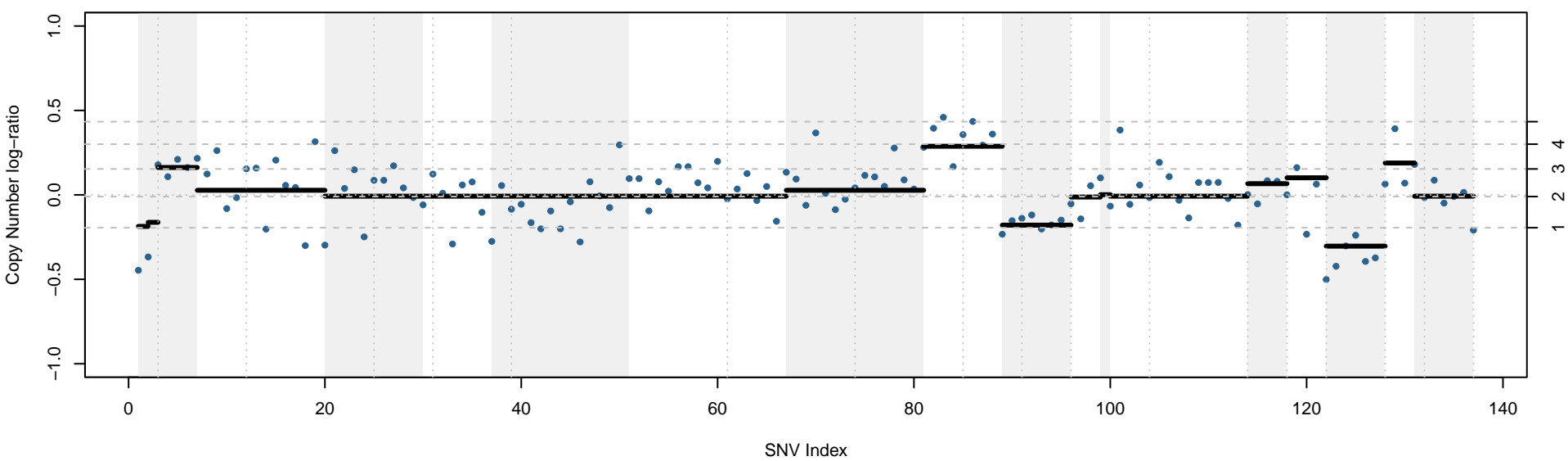
Purity: 0.24 Tumor ploidy: 2.058

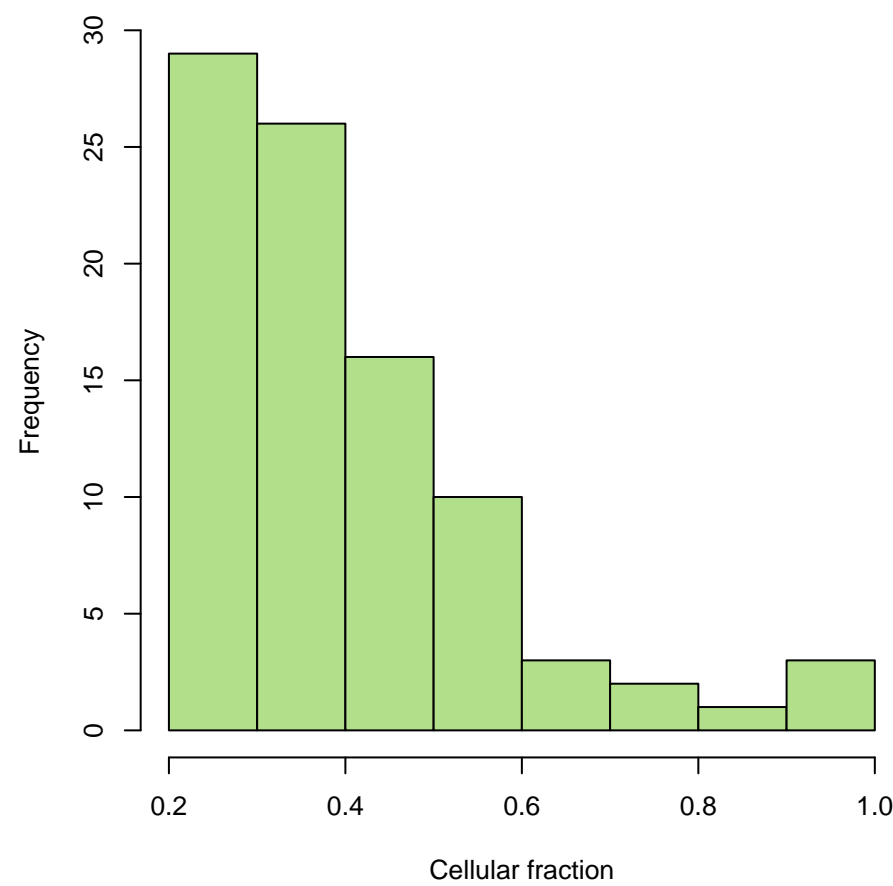
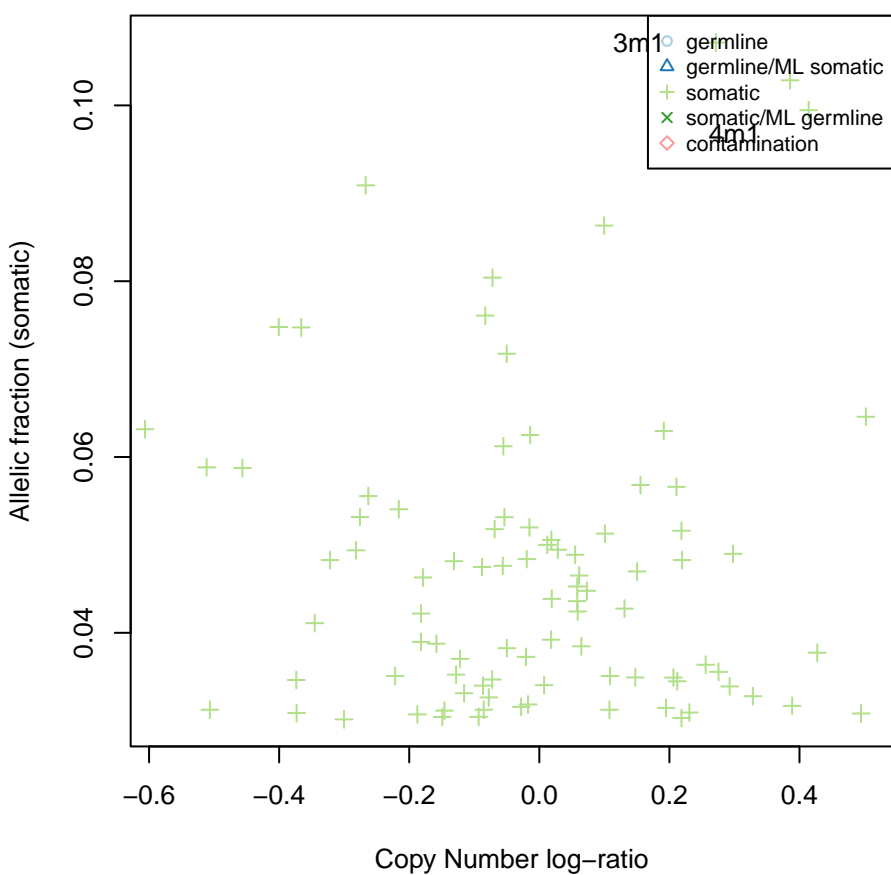
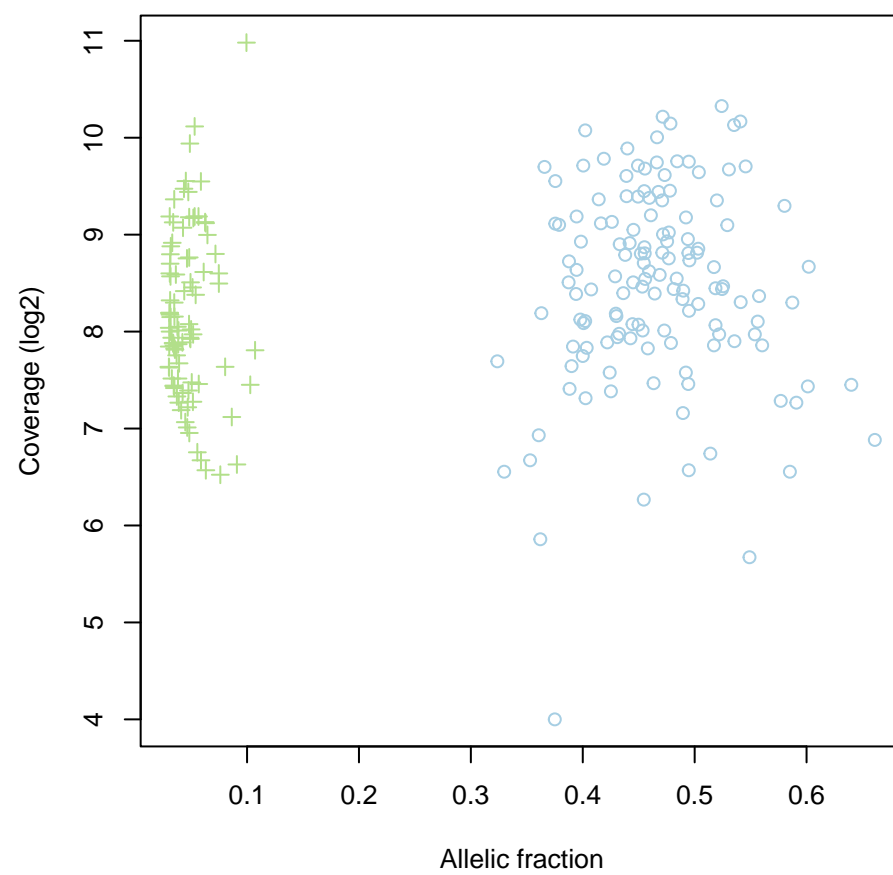
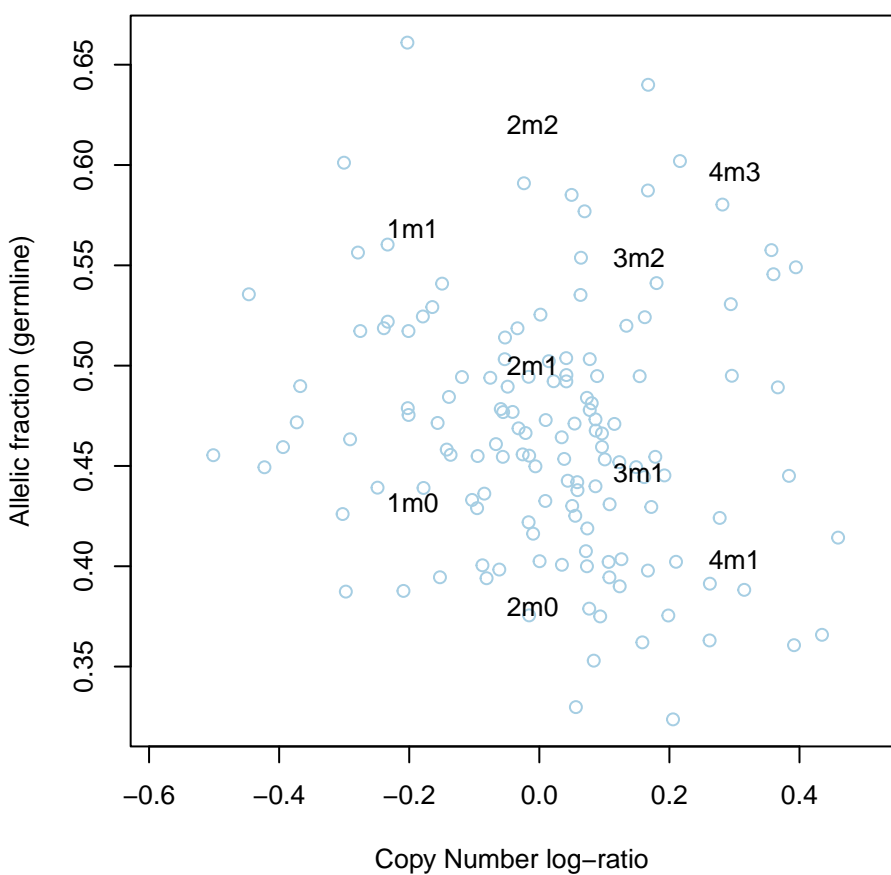


Purity: 0.24 Tumor ploidy: 2.058 SNV log-likelihood: 67.22 GoF: 91% Mean coverage: 385;403

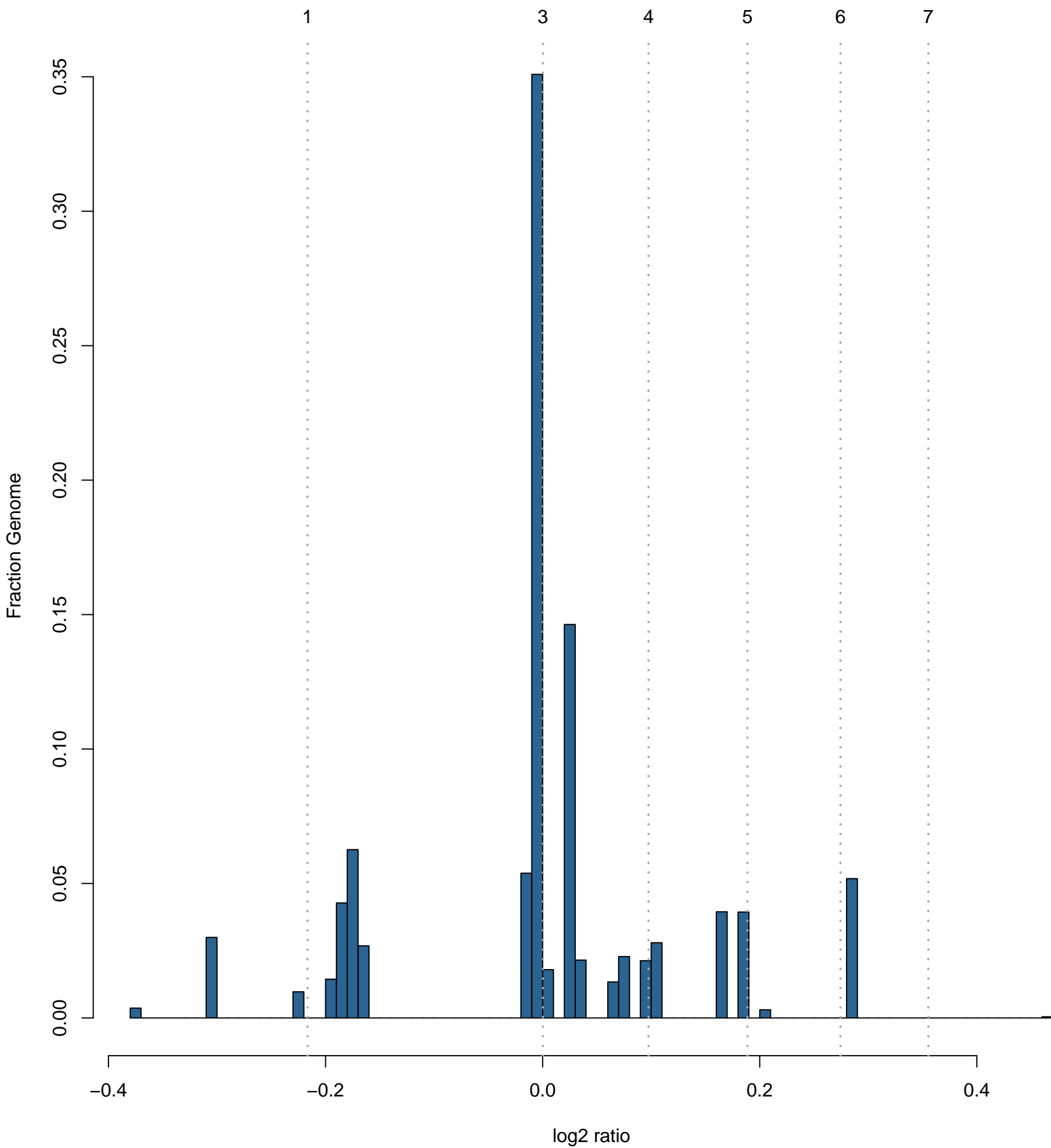


SCNA-fit log-likelihood: -2869.65

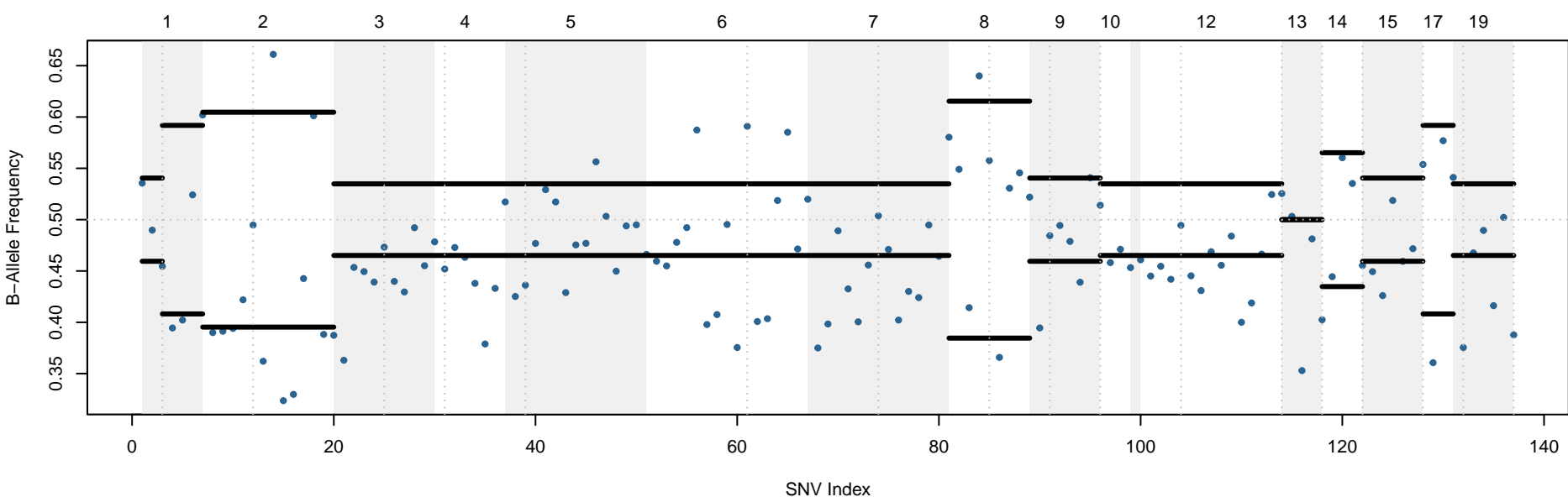




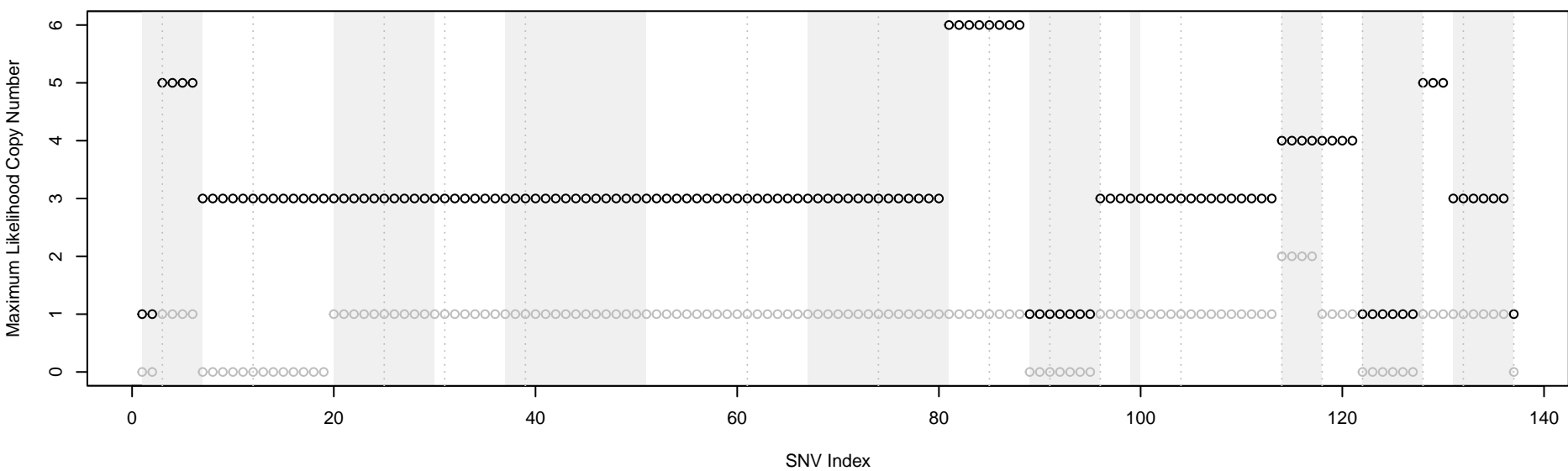
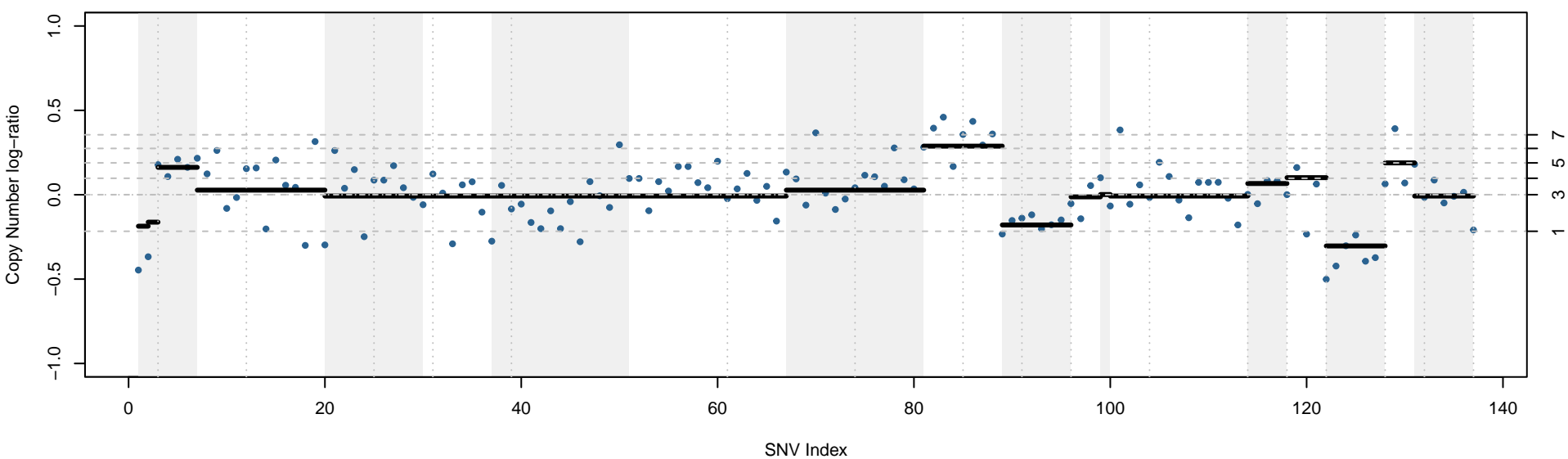
Purity: 0.15 Tumor ploidy: 2.998

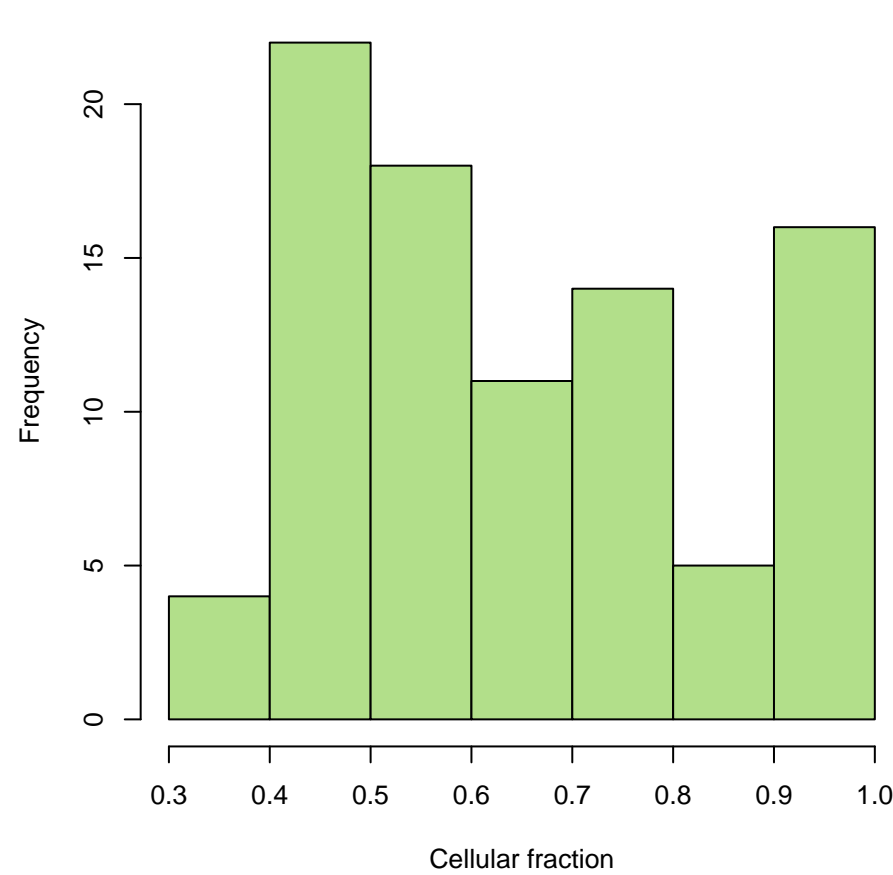
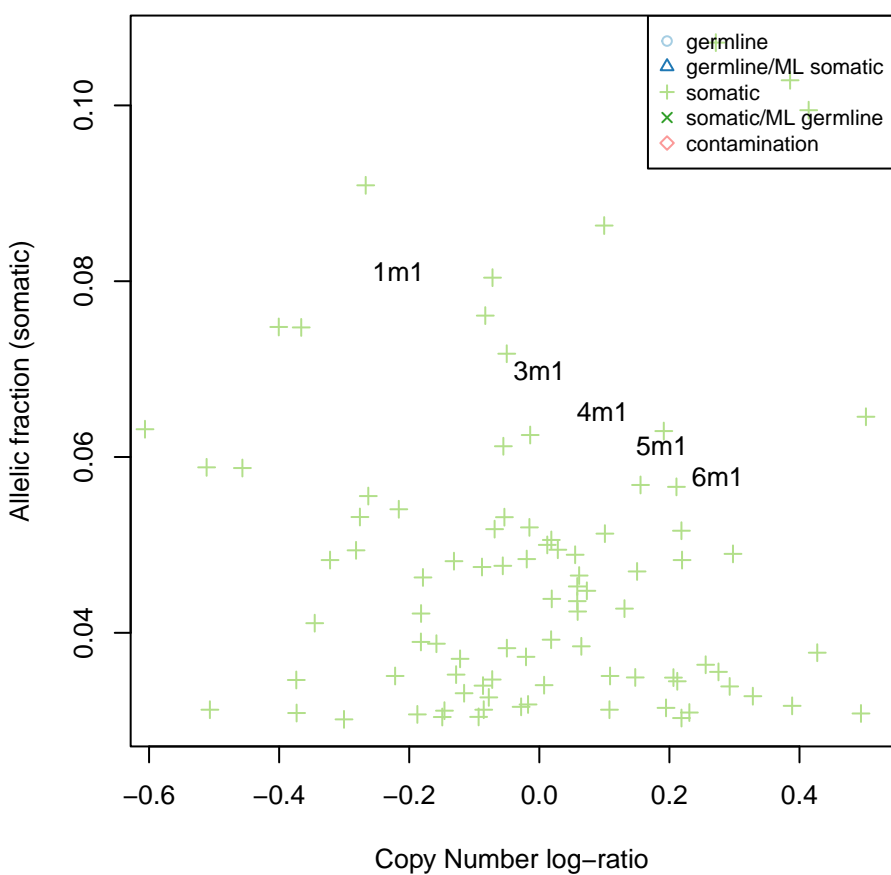
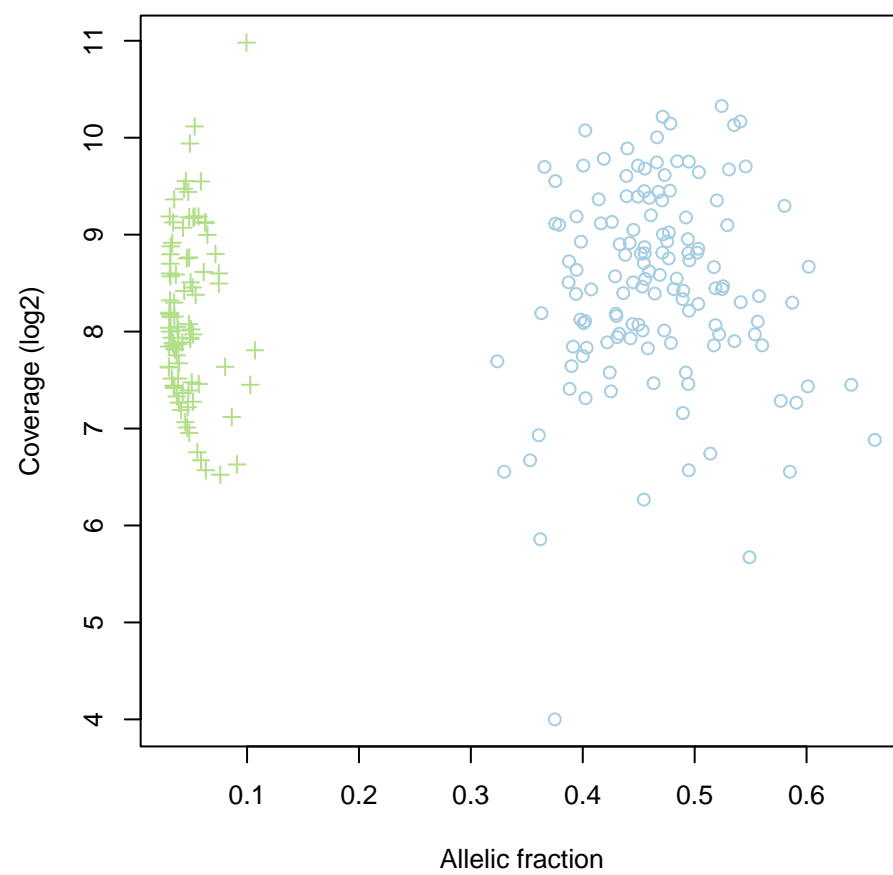
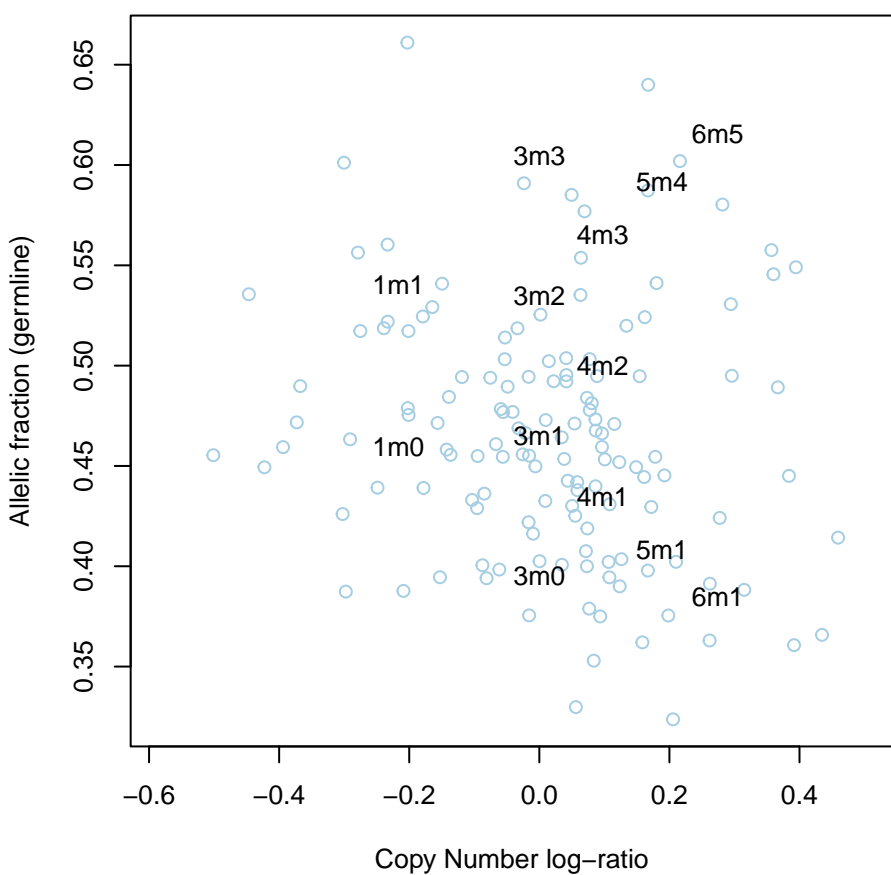


Purity: 0.15 Tumor ploidy: 2.998 SNV log-likelihood: -3.56 GoF: 97.1% Mean coverage: 385,403

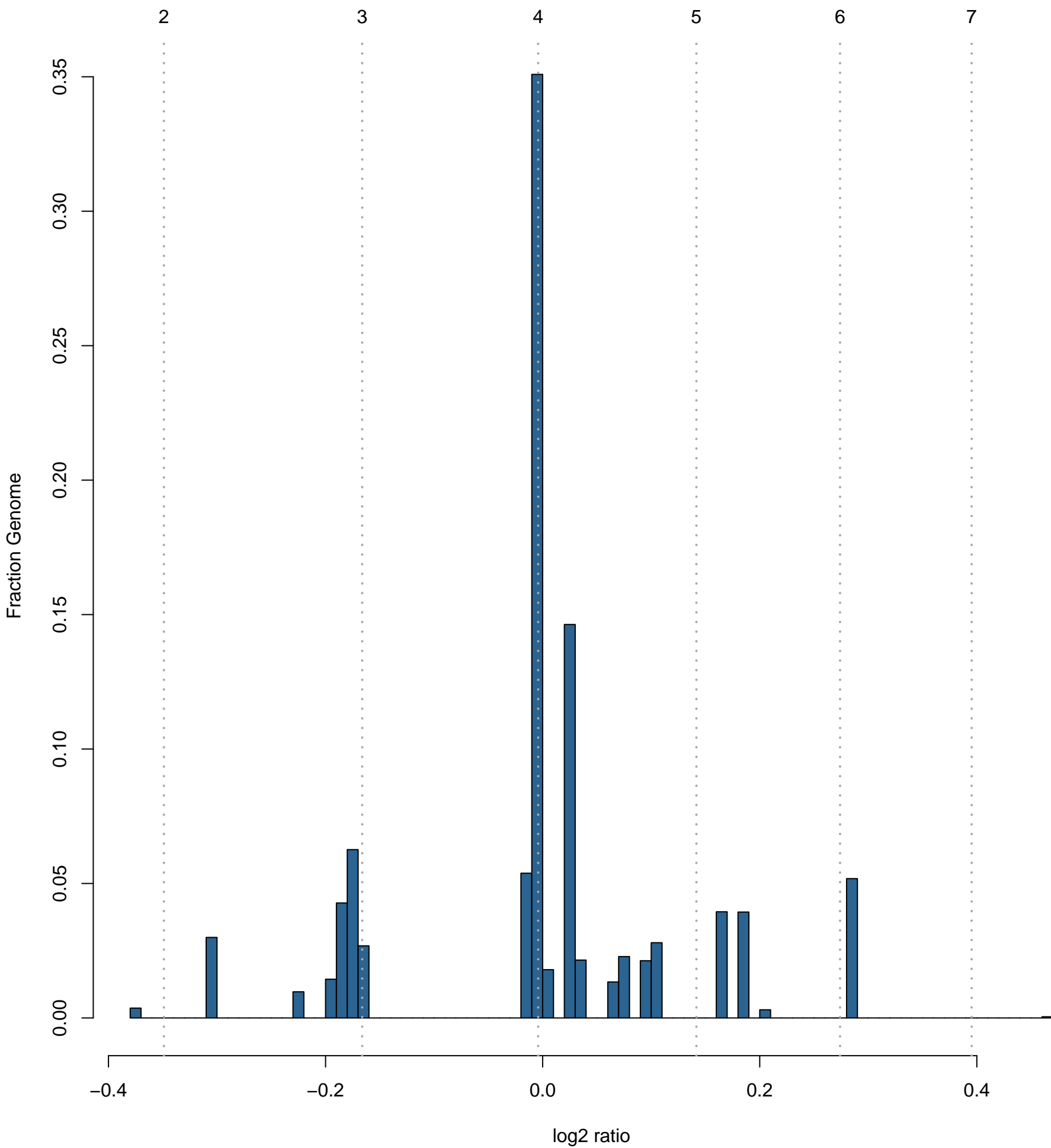


SCNA-fit log-likelihood: -2810.21

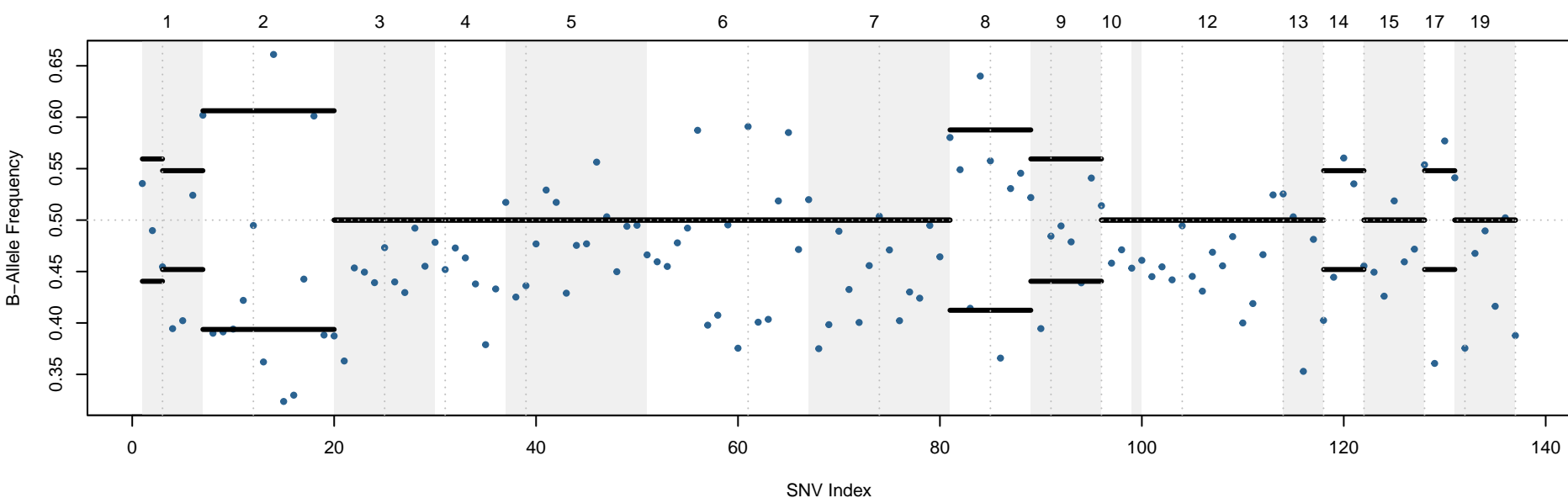




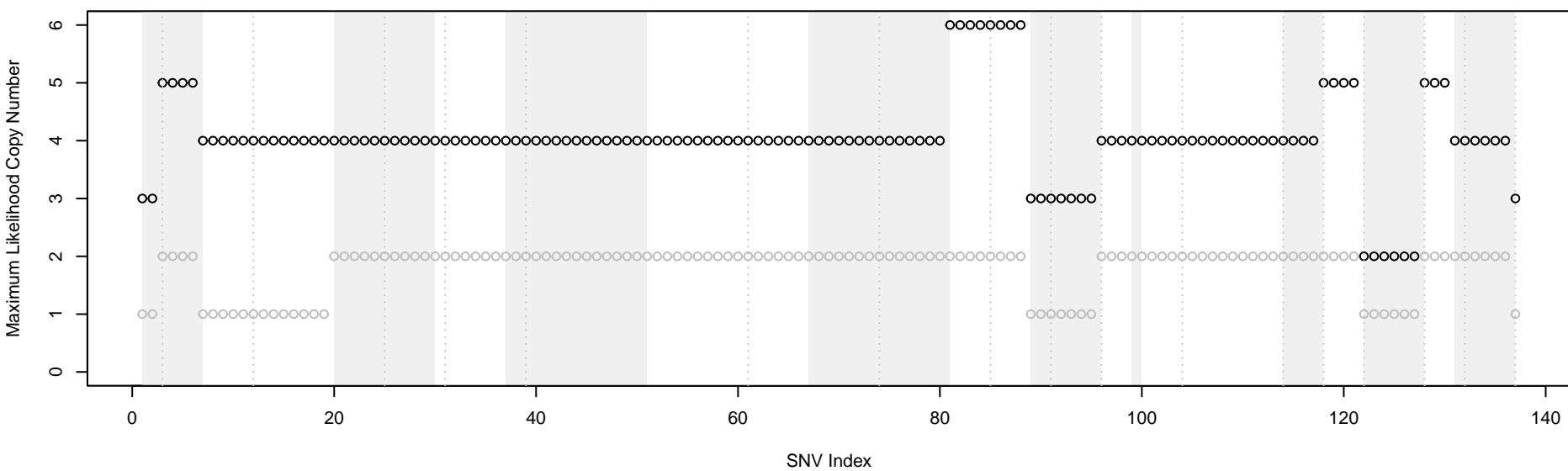
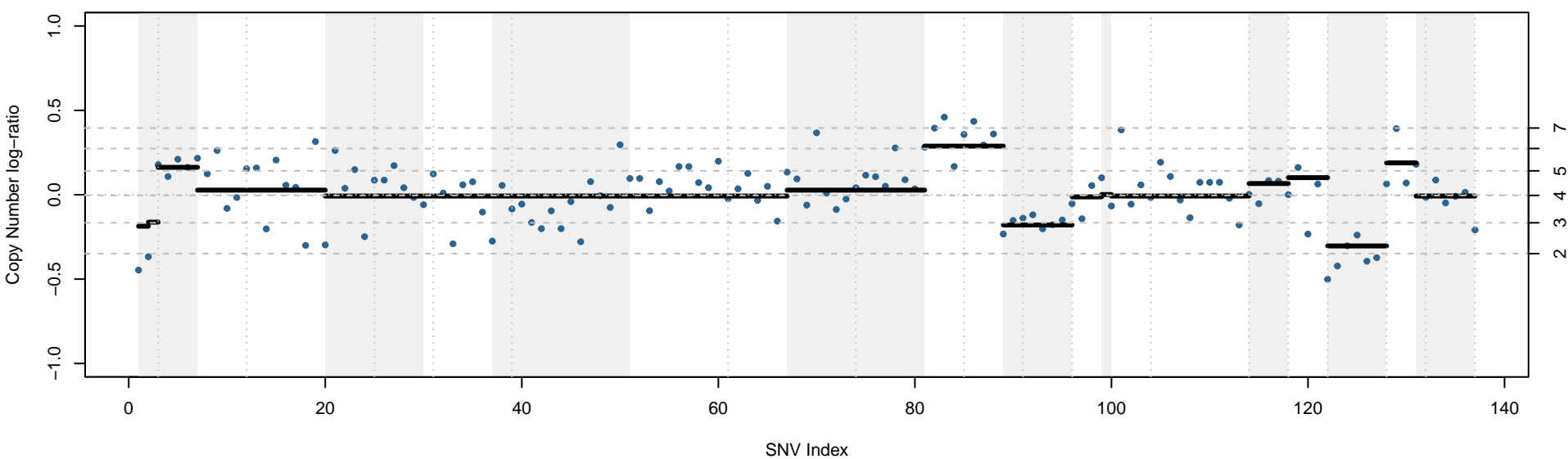
Purity: 0.27 Tumor ploidy: 4.027



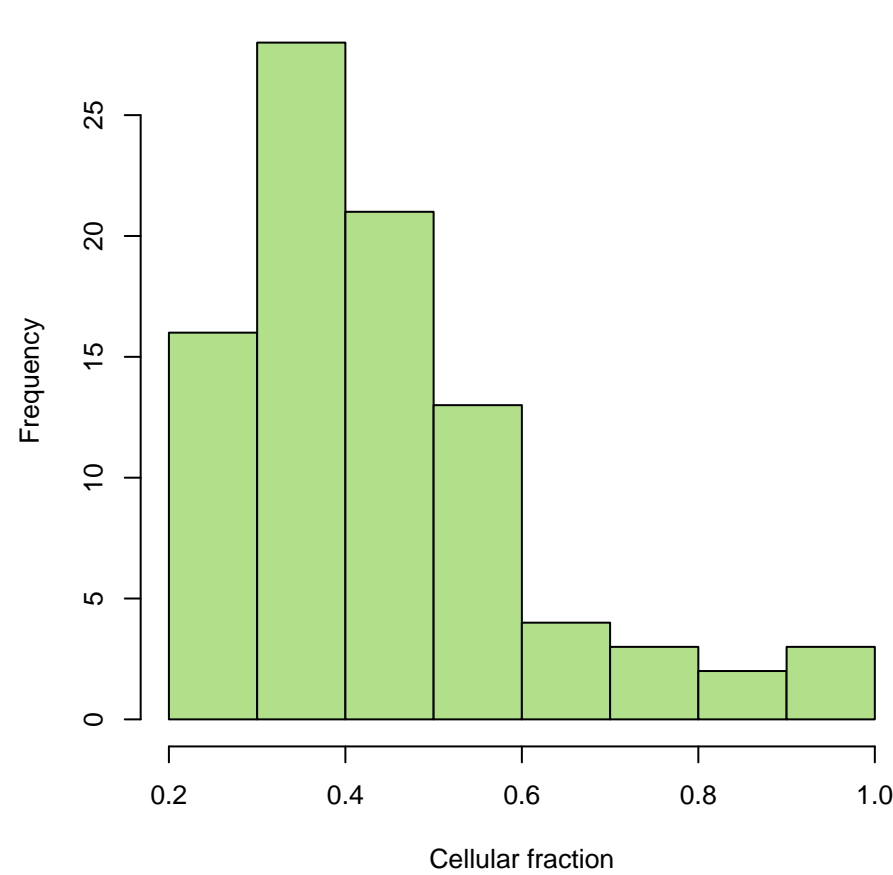
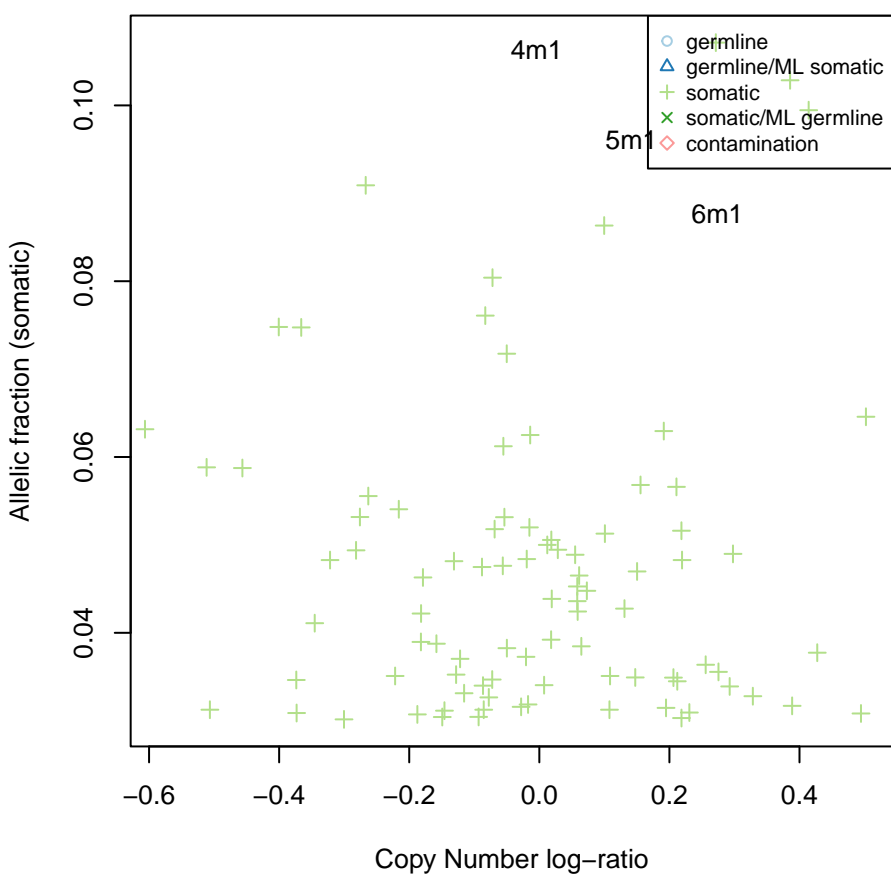
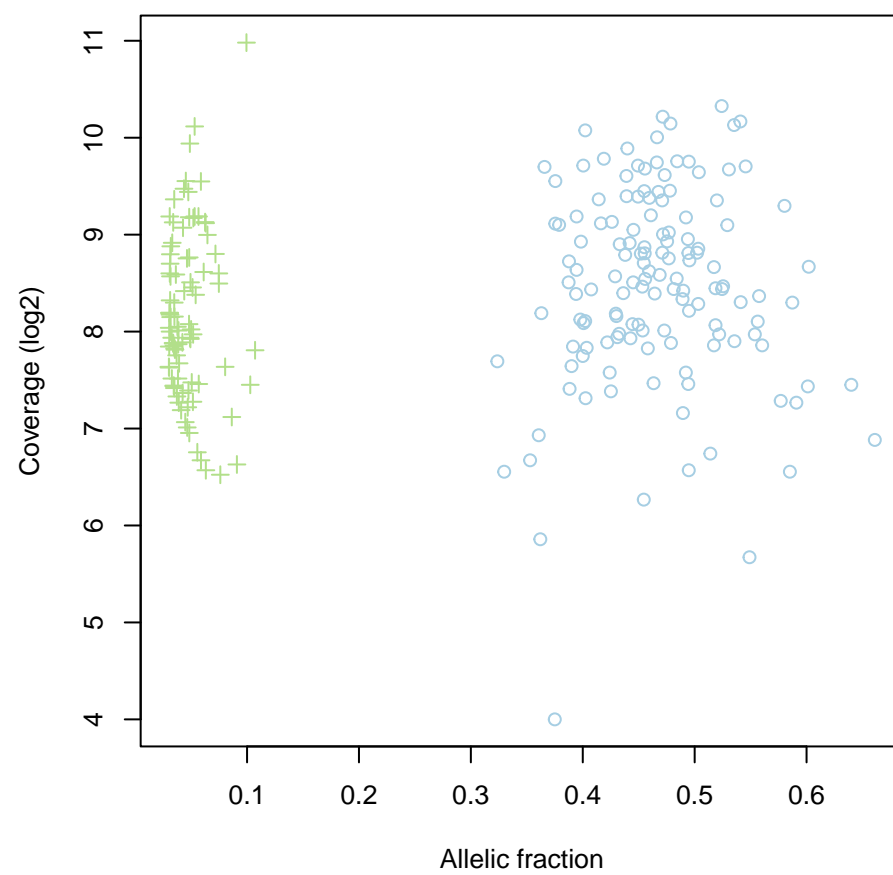
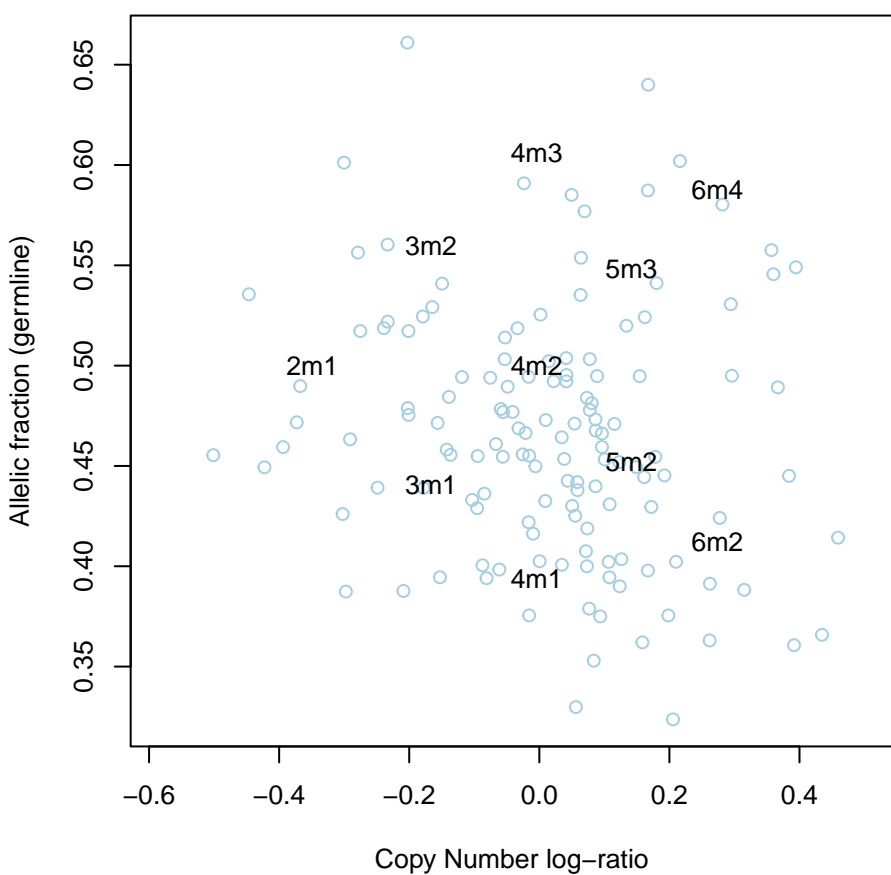
Purity: 0.27 Tumor ploidy: 4.027 SNV log-likelihood: -63.16 GoF: 92.3% Mean coverage: 385;403



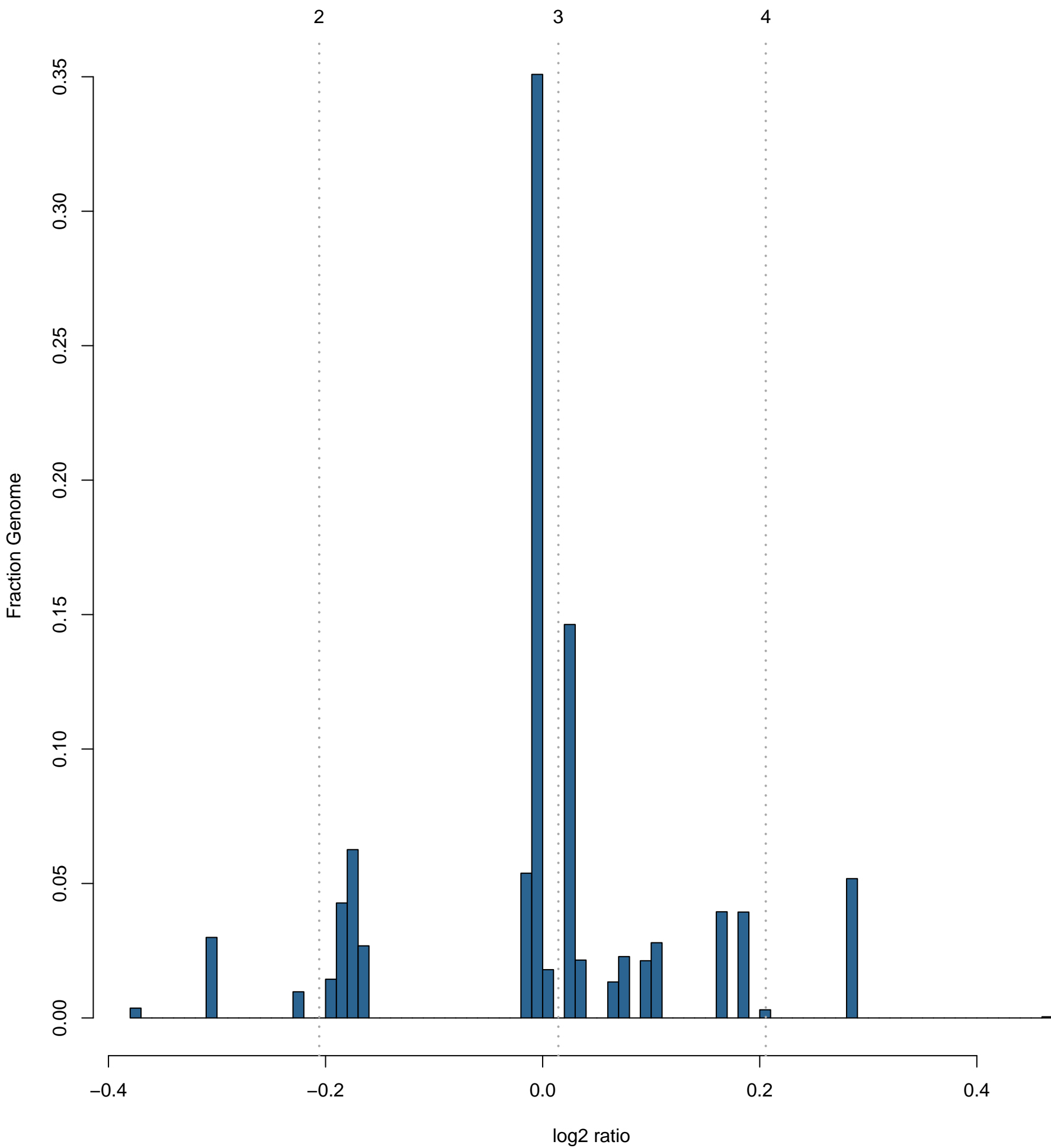
SCNA-fit log-likelihood: -2798.54



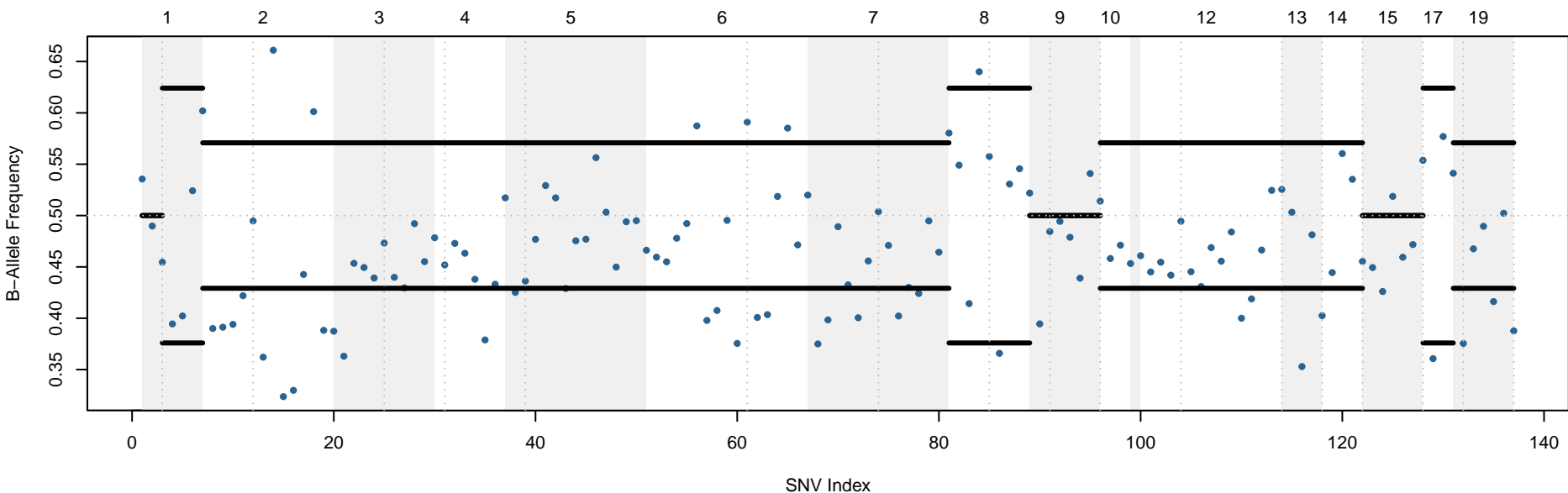




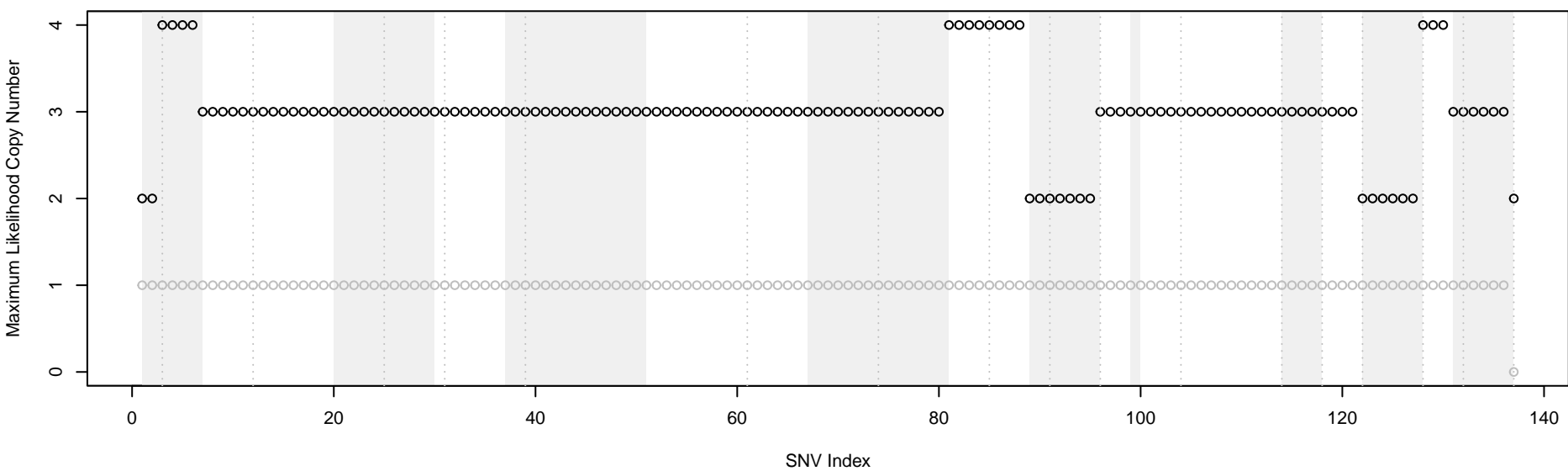
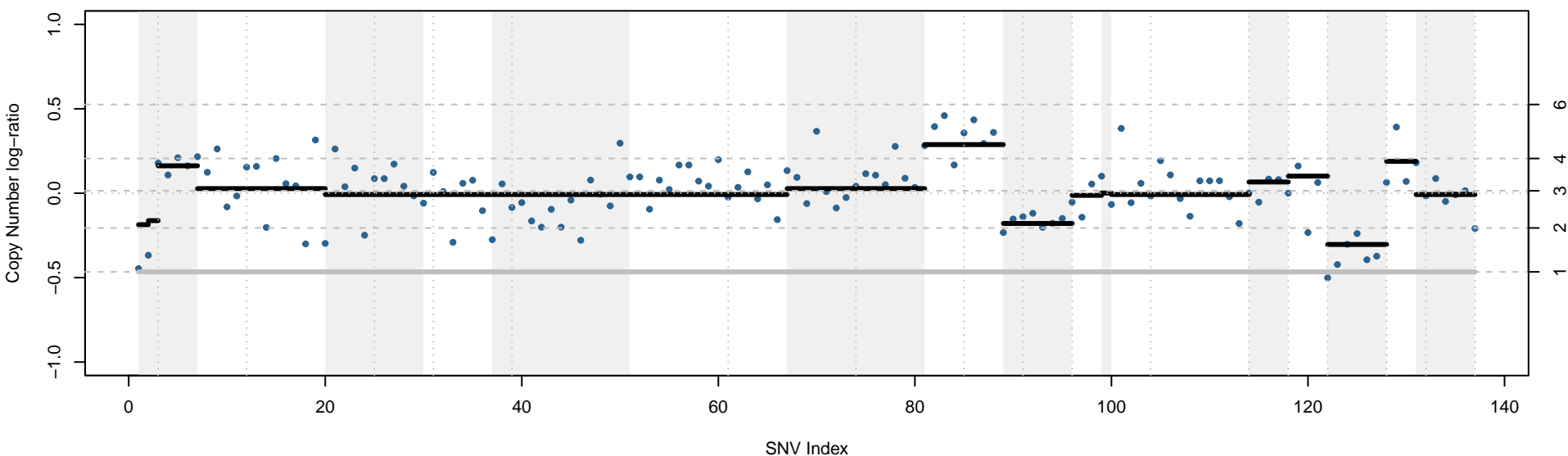
Purity: 0.33 Tumor ploidy: 2.93

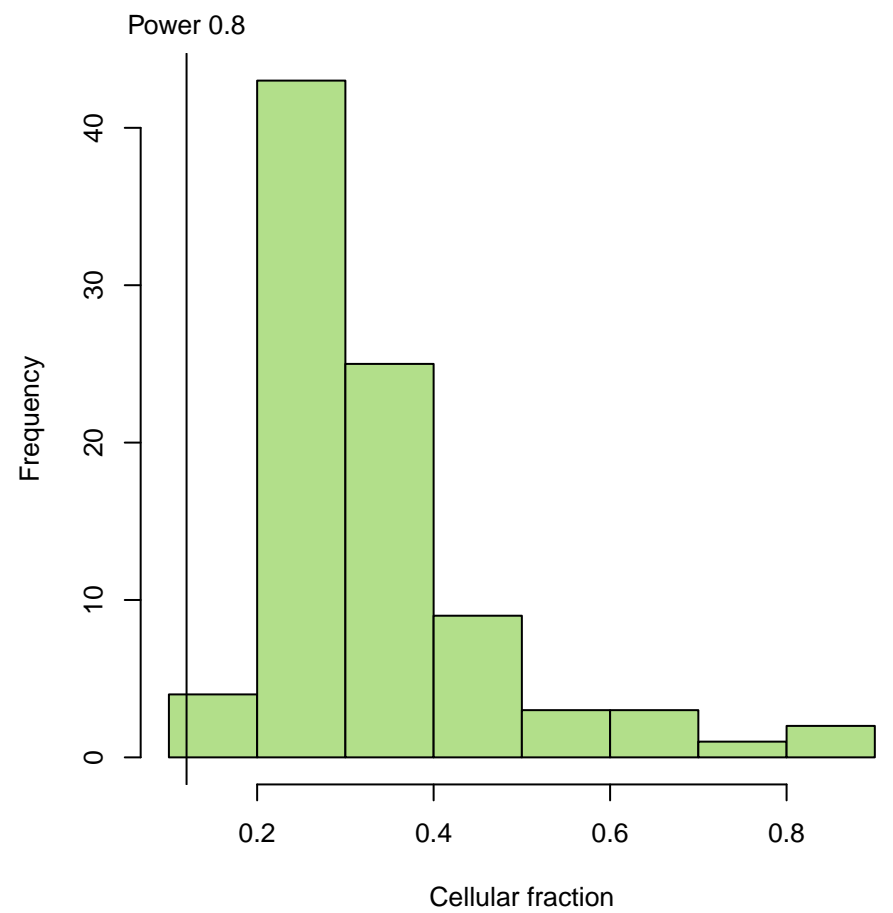
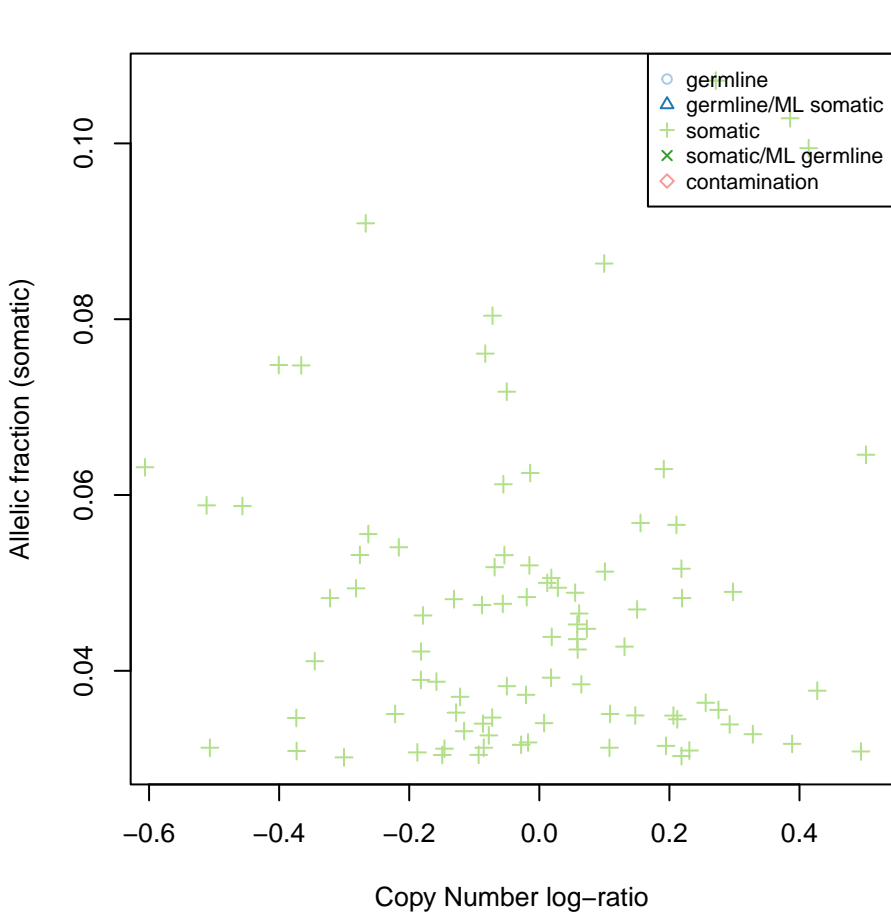
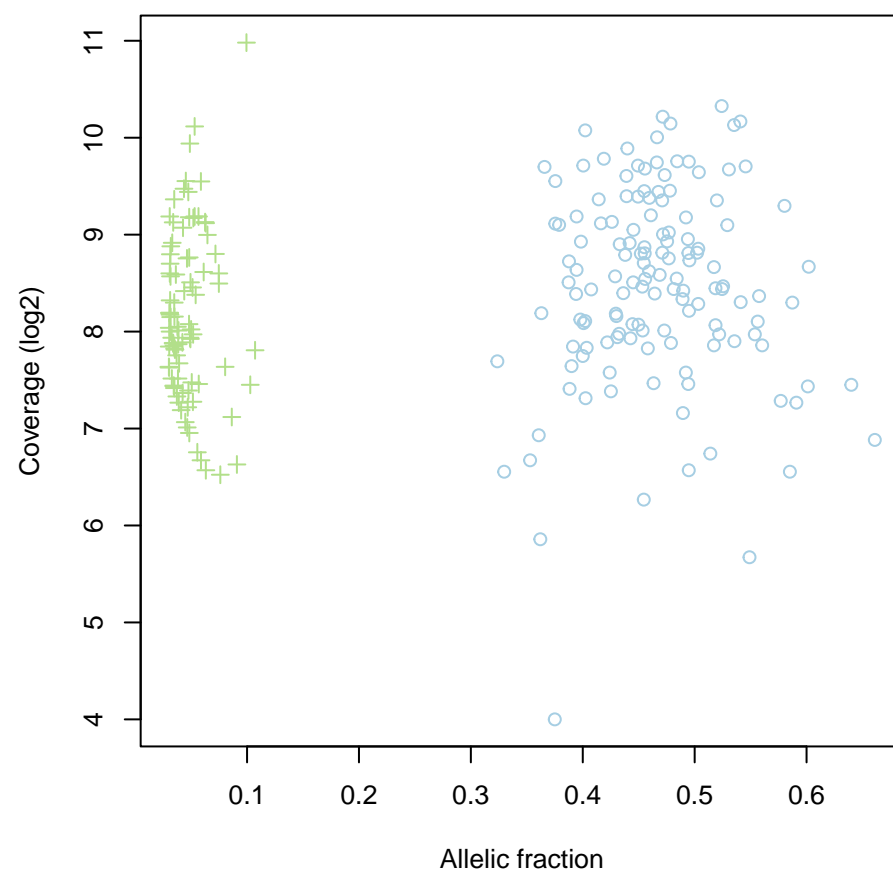
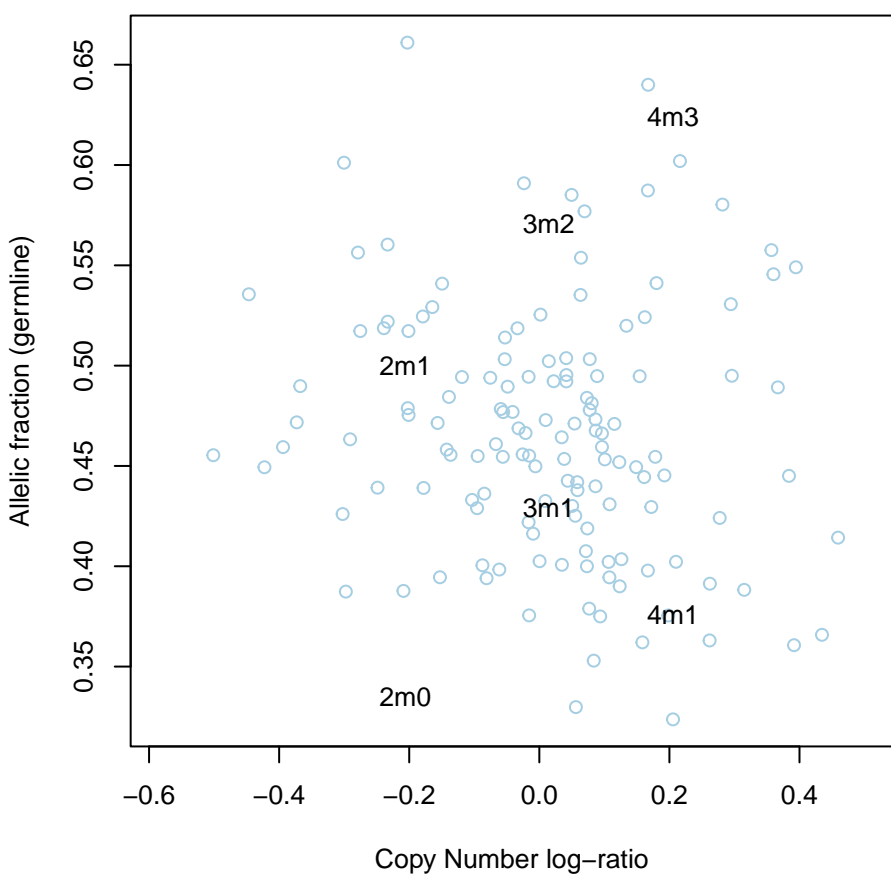


Purity: 0.33 Tumor ploidy: 2.93 SNV log-likelihood: -37.39 GoF: 87.1% Mean coverage: 385,403

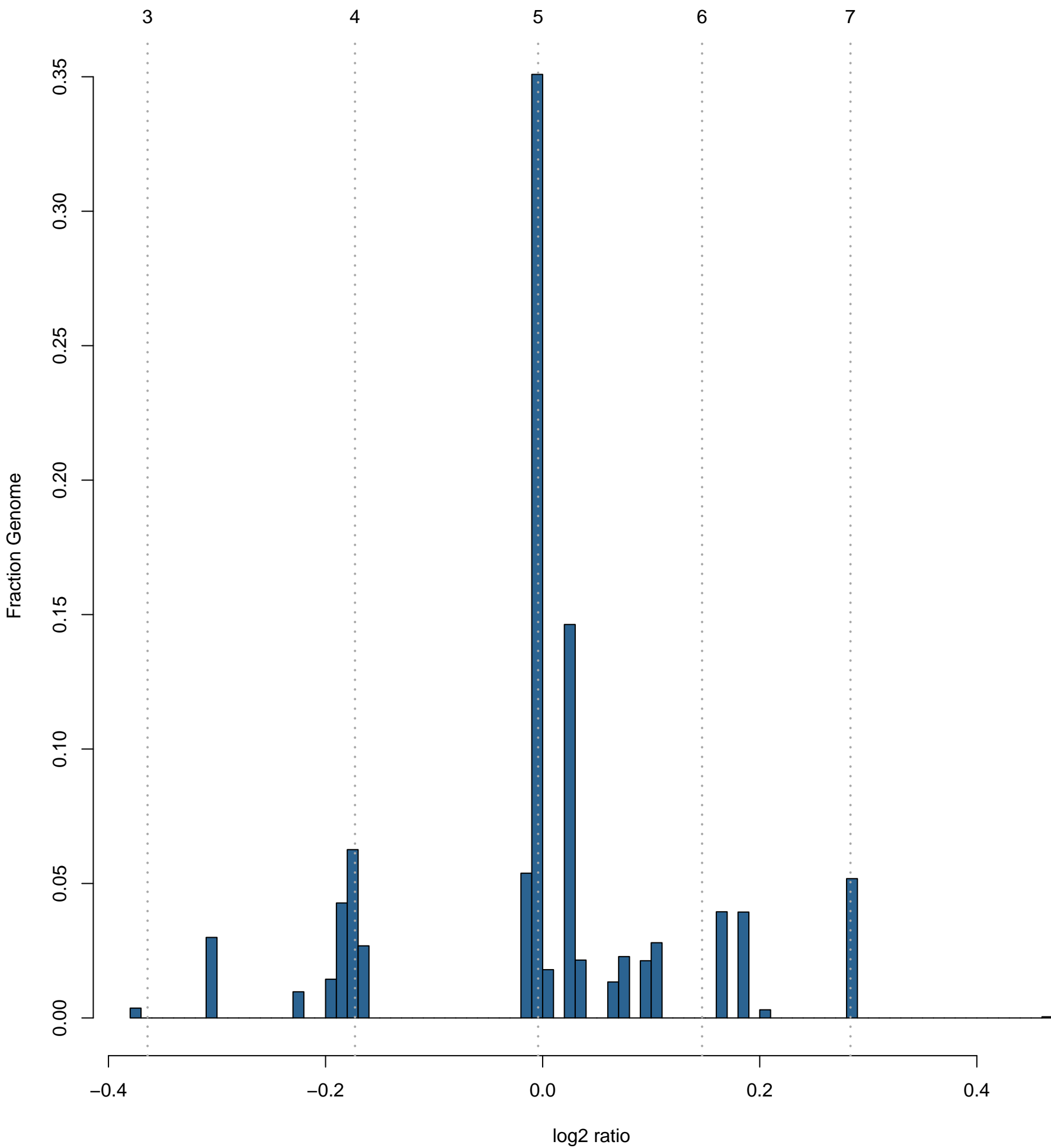


SCNA-fit log-likelihood: -2926.89

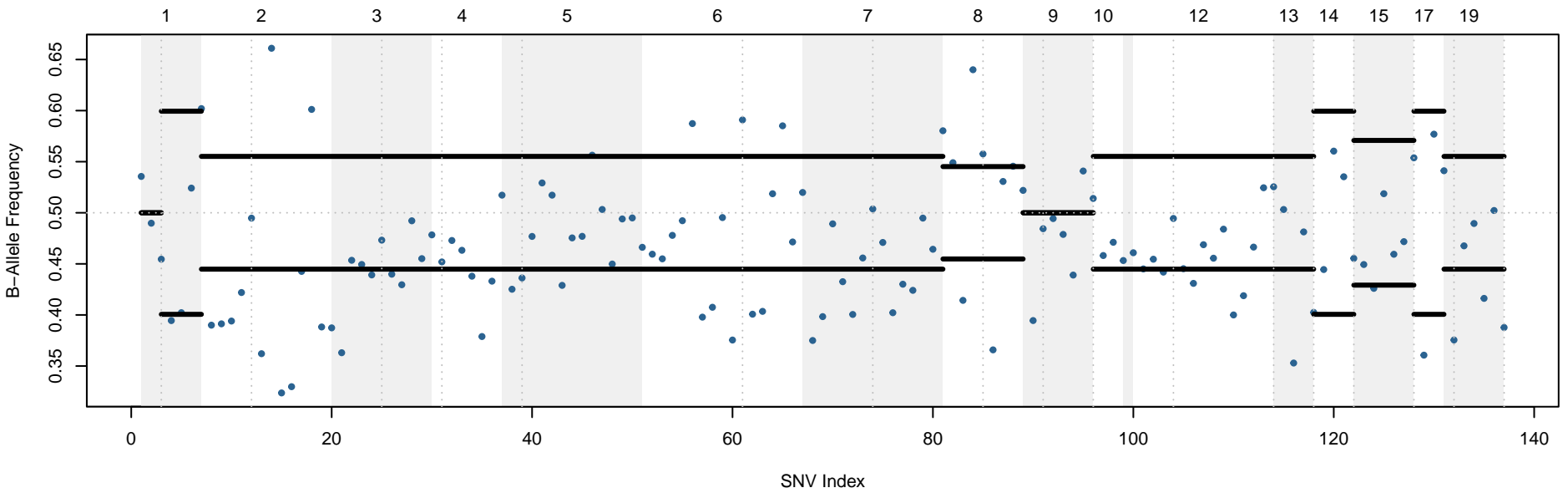




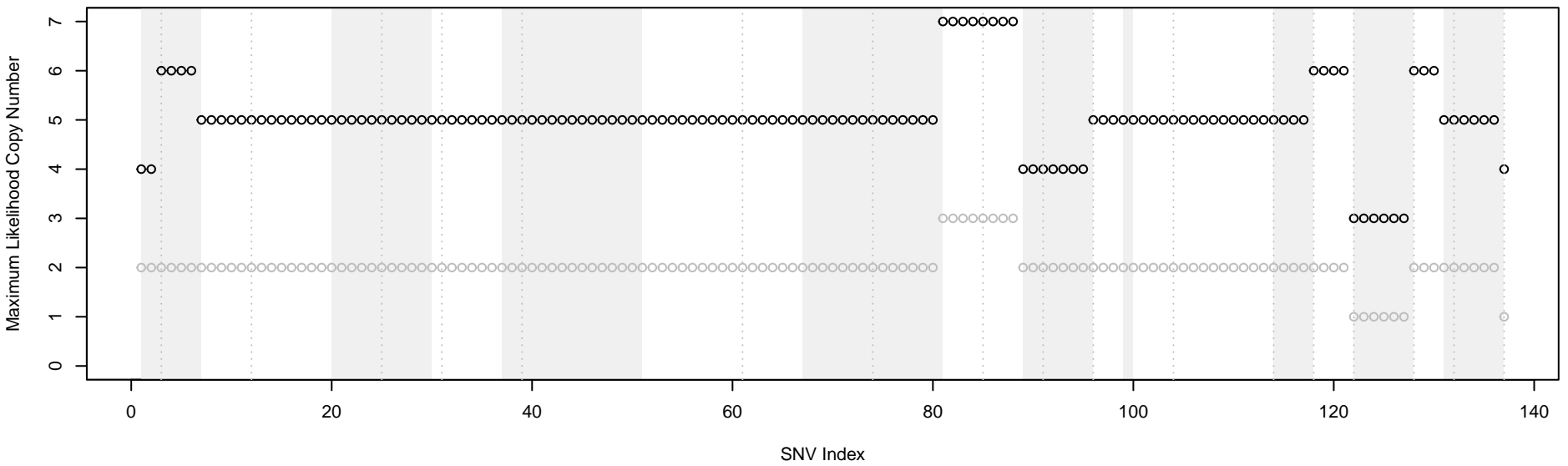
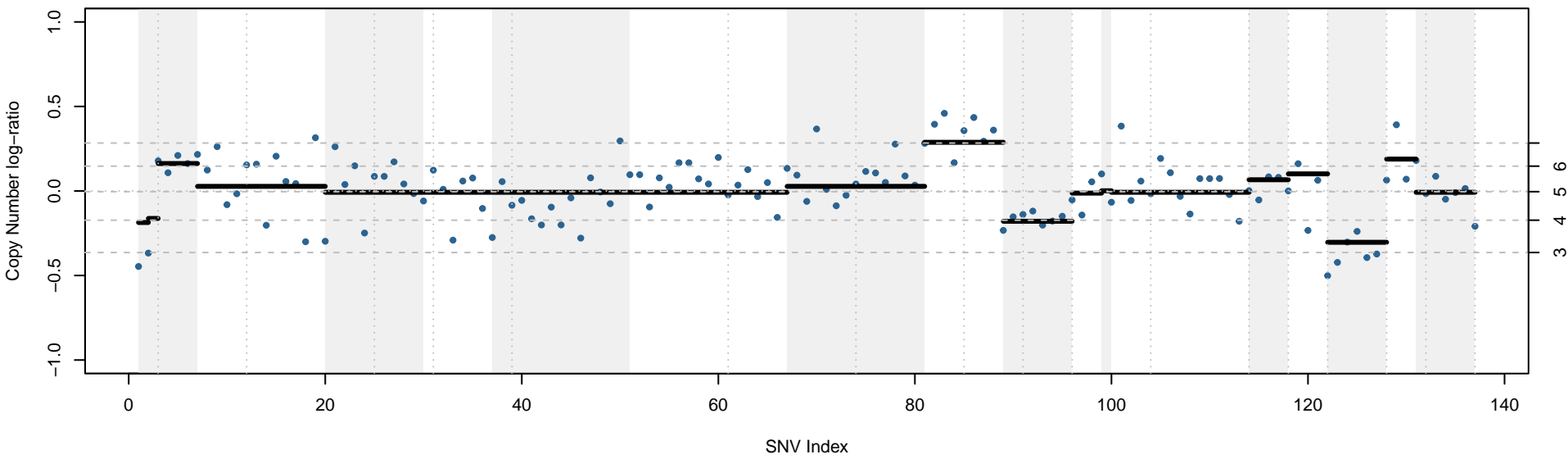
Purity: 0.33 Tumor ploidy: 5.026

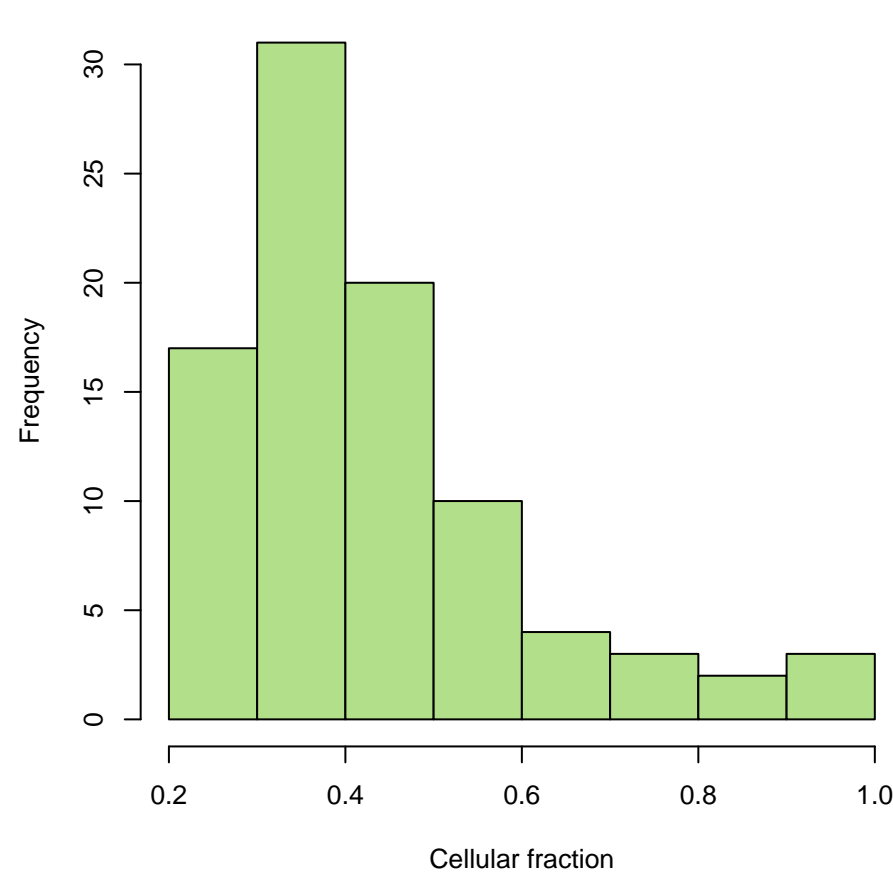
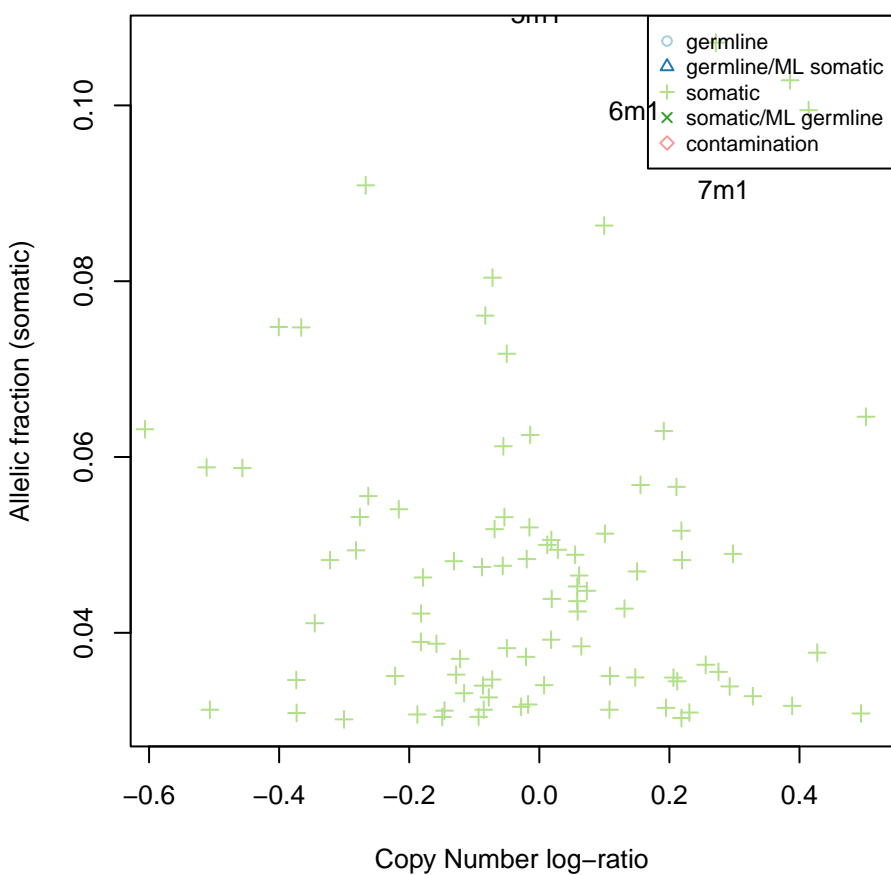
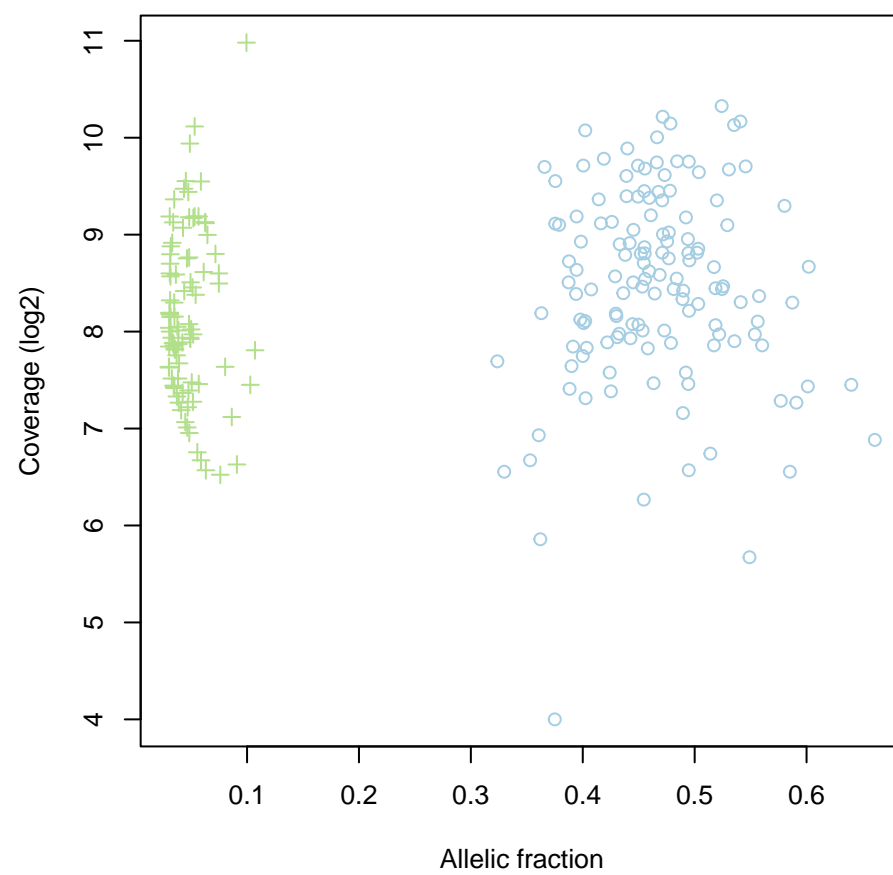
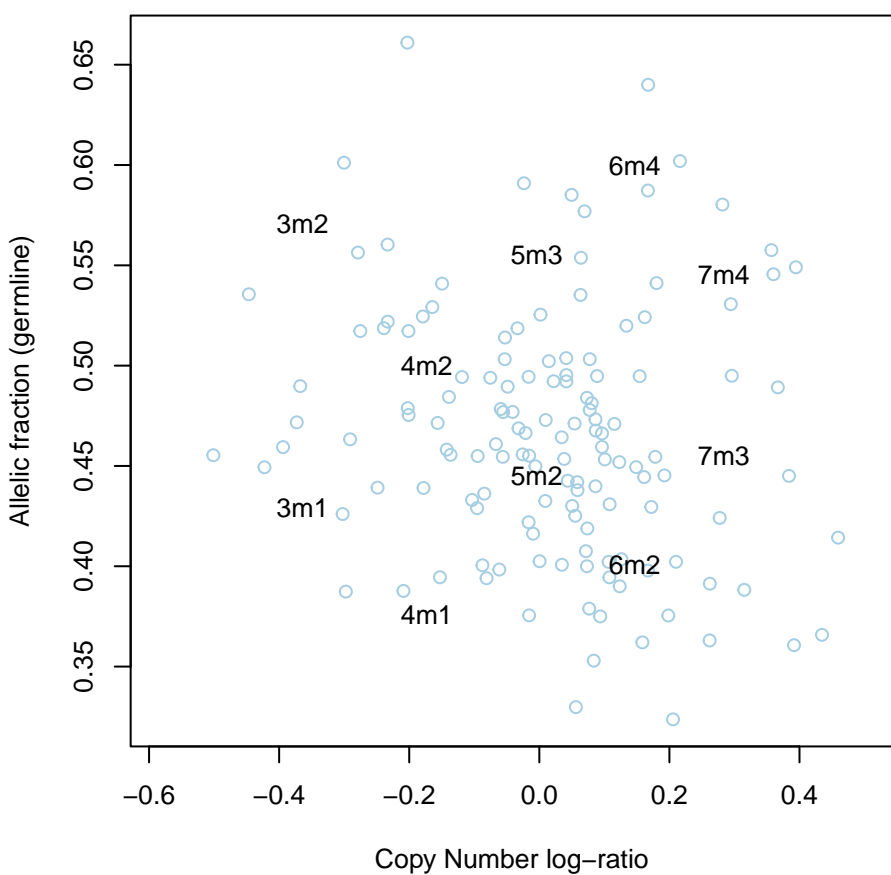


Purity: 0.33 Tumor ploidy: 5.026 SNV log-likelihood: -119.72 GoF: 92.9% Mean coverage: 385,403

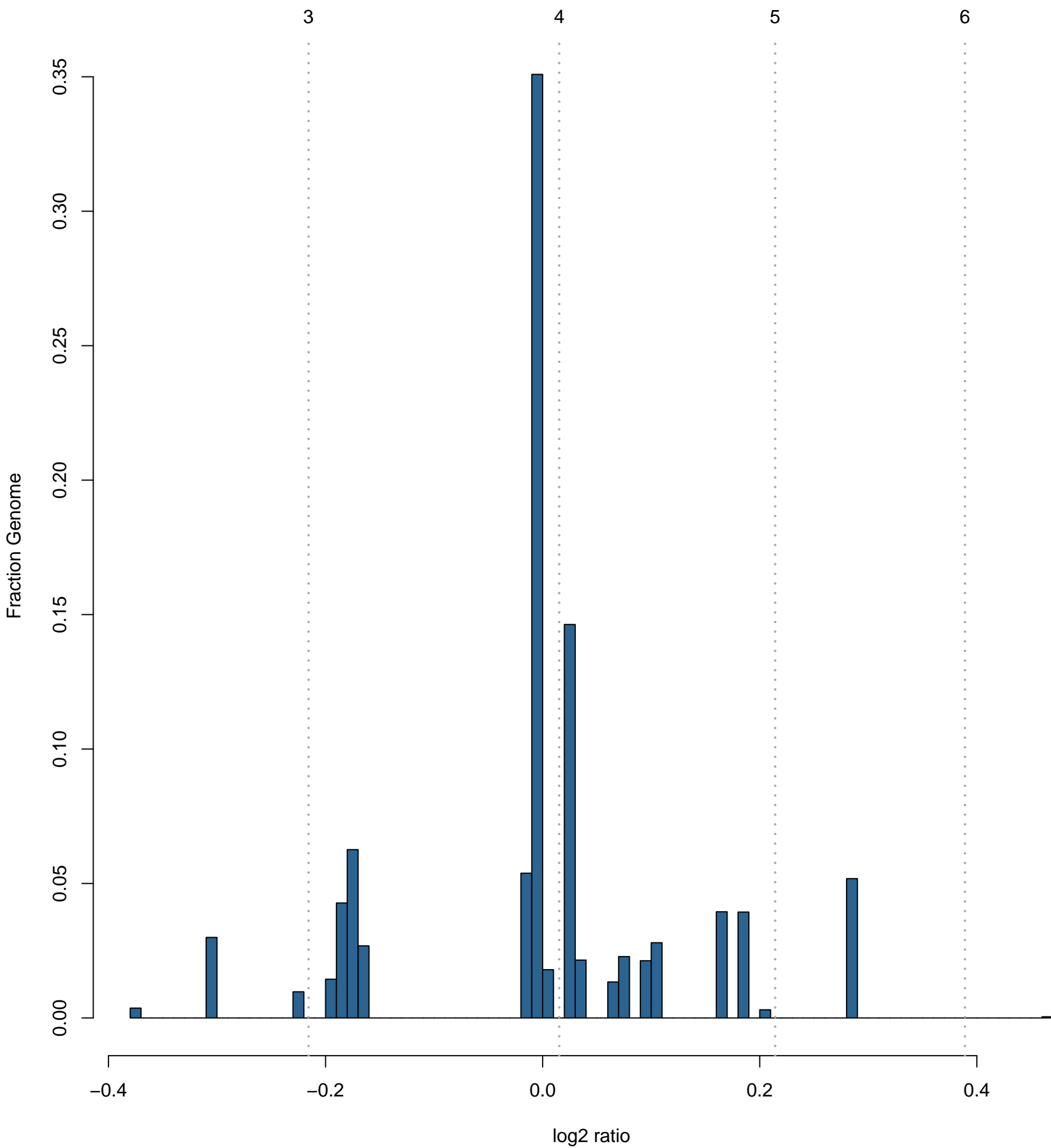


SCNA-fit log-likelihood: -2803.32



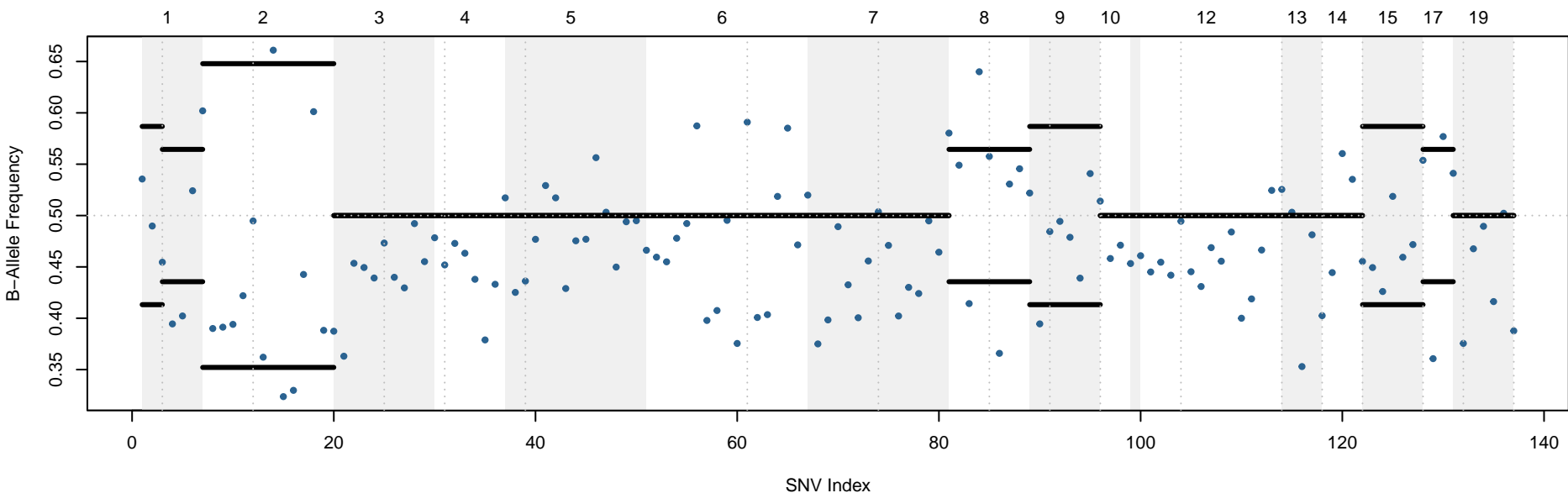


Purity: 0.42 Tumor ploidy: 3.929

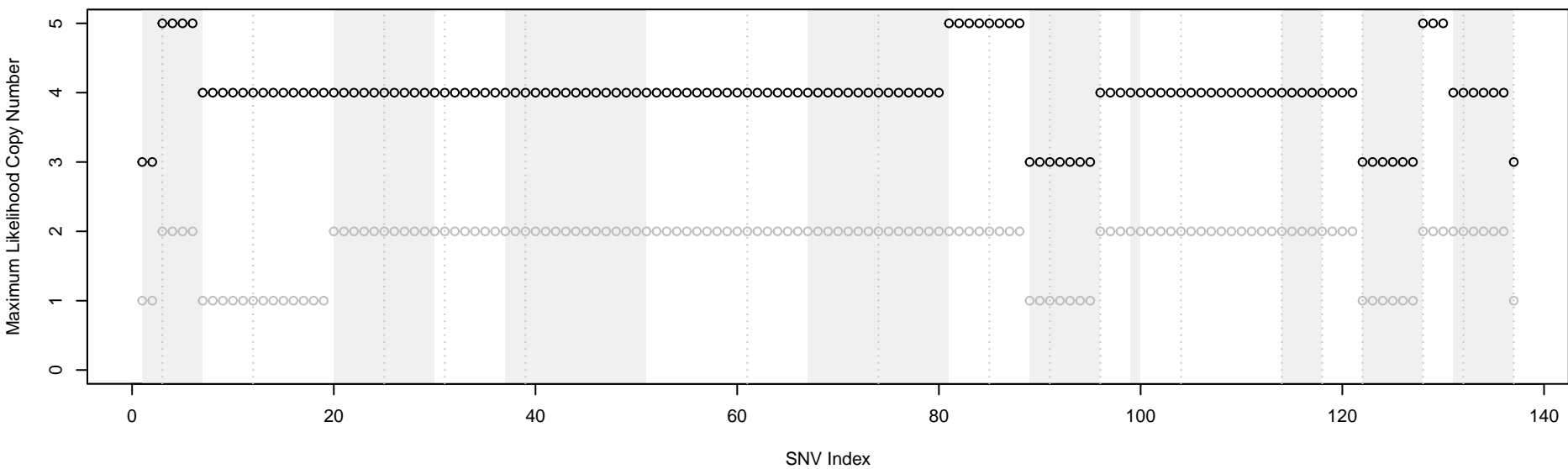
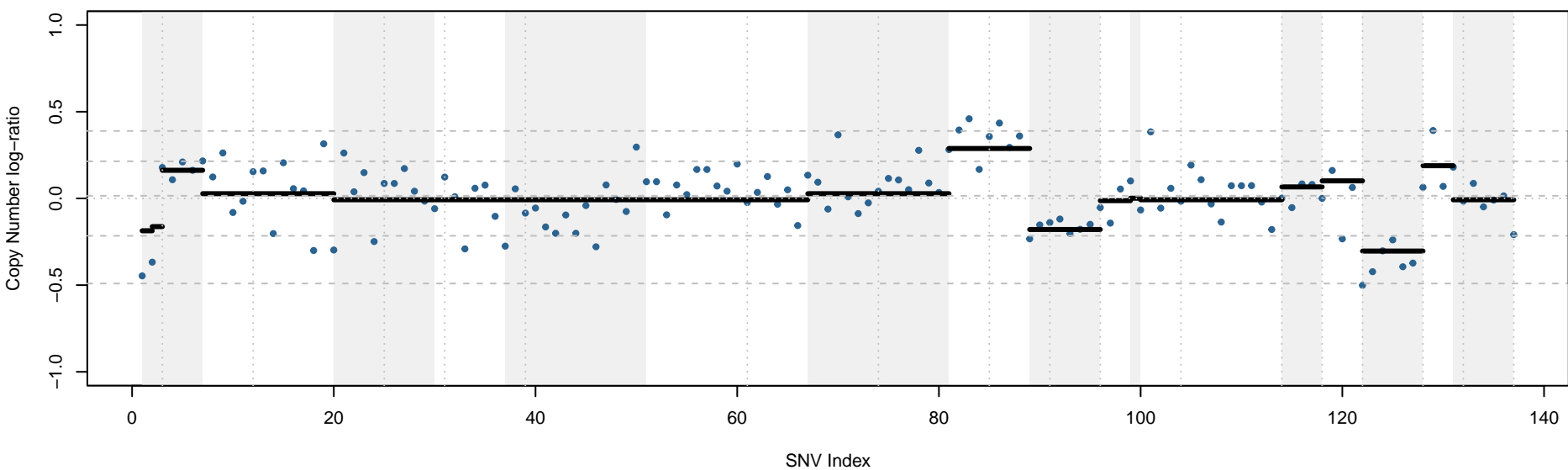


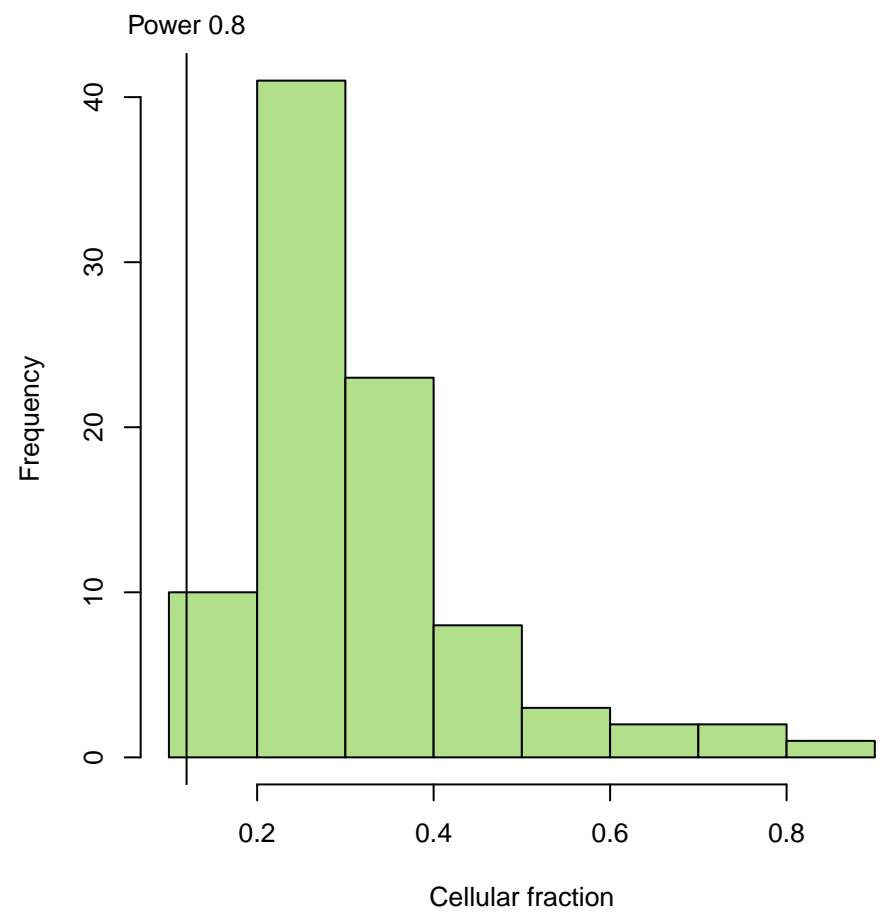
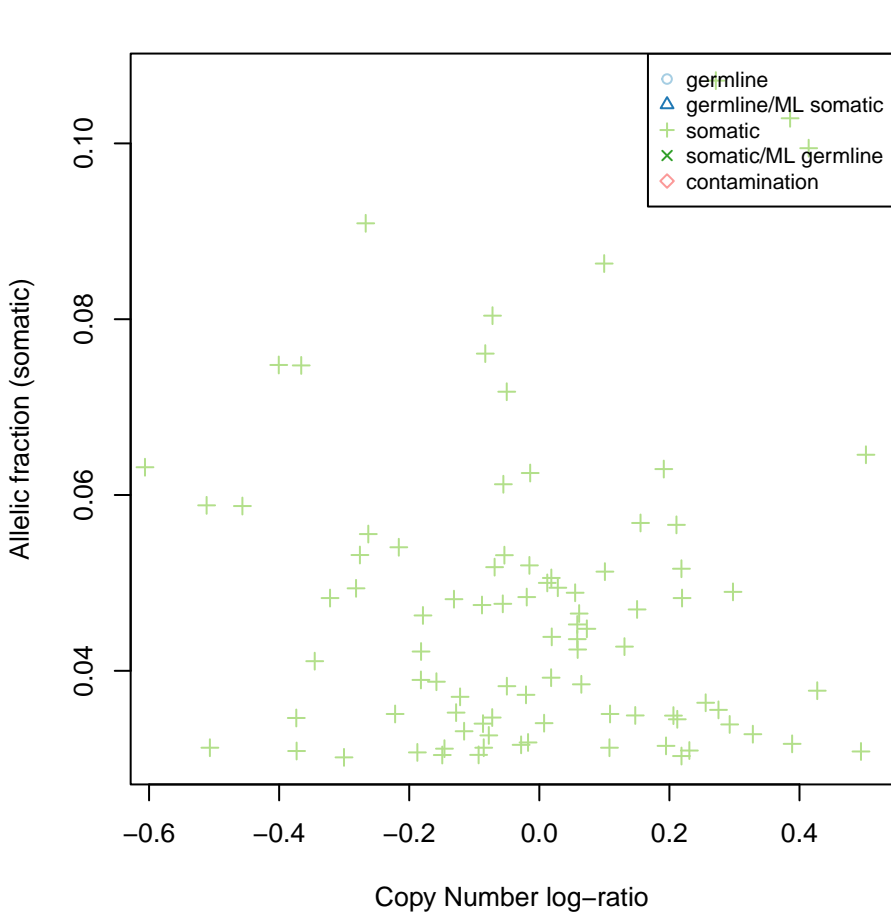
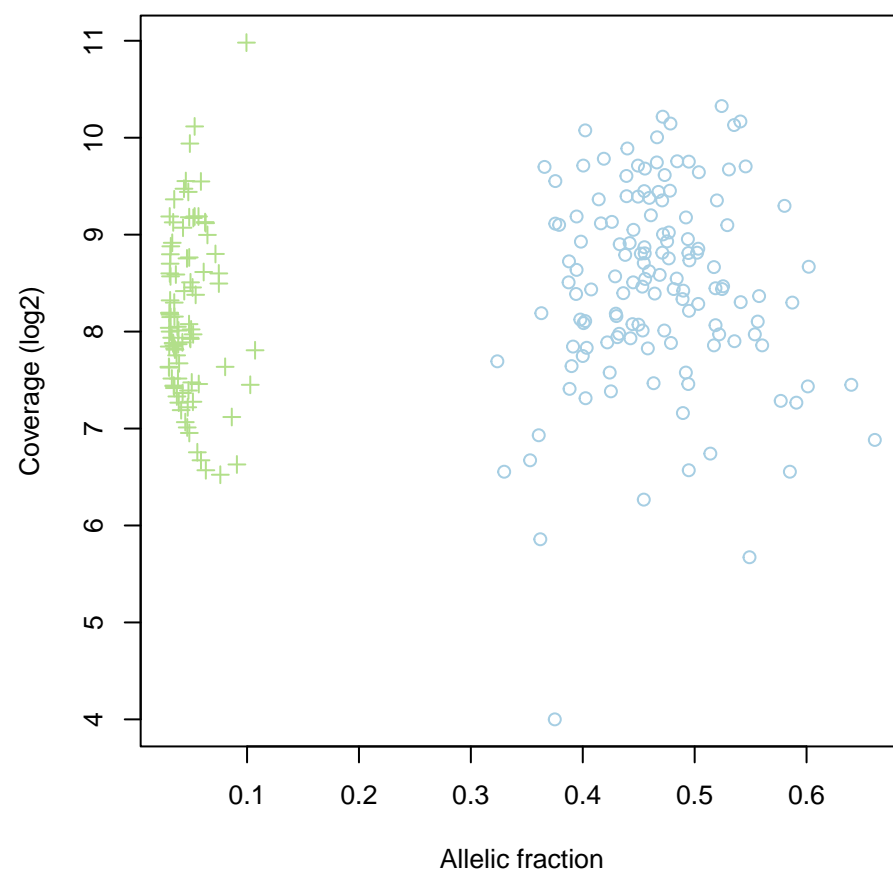
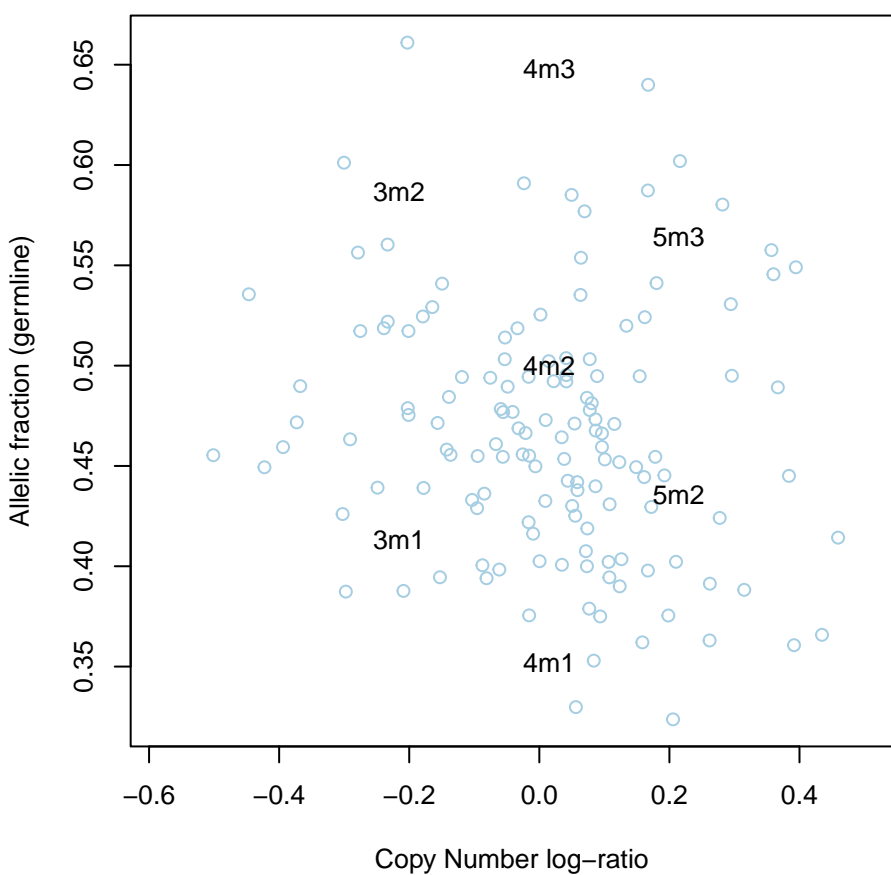


Purity: 0.42 Tumor ploidy: 3.929 SNV log-likelihood: -106.75 GoF: 84.4% Mean coverage: 385,403

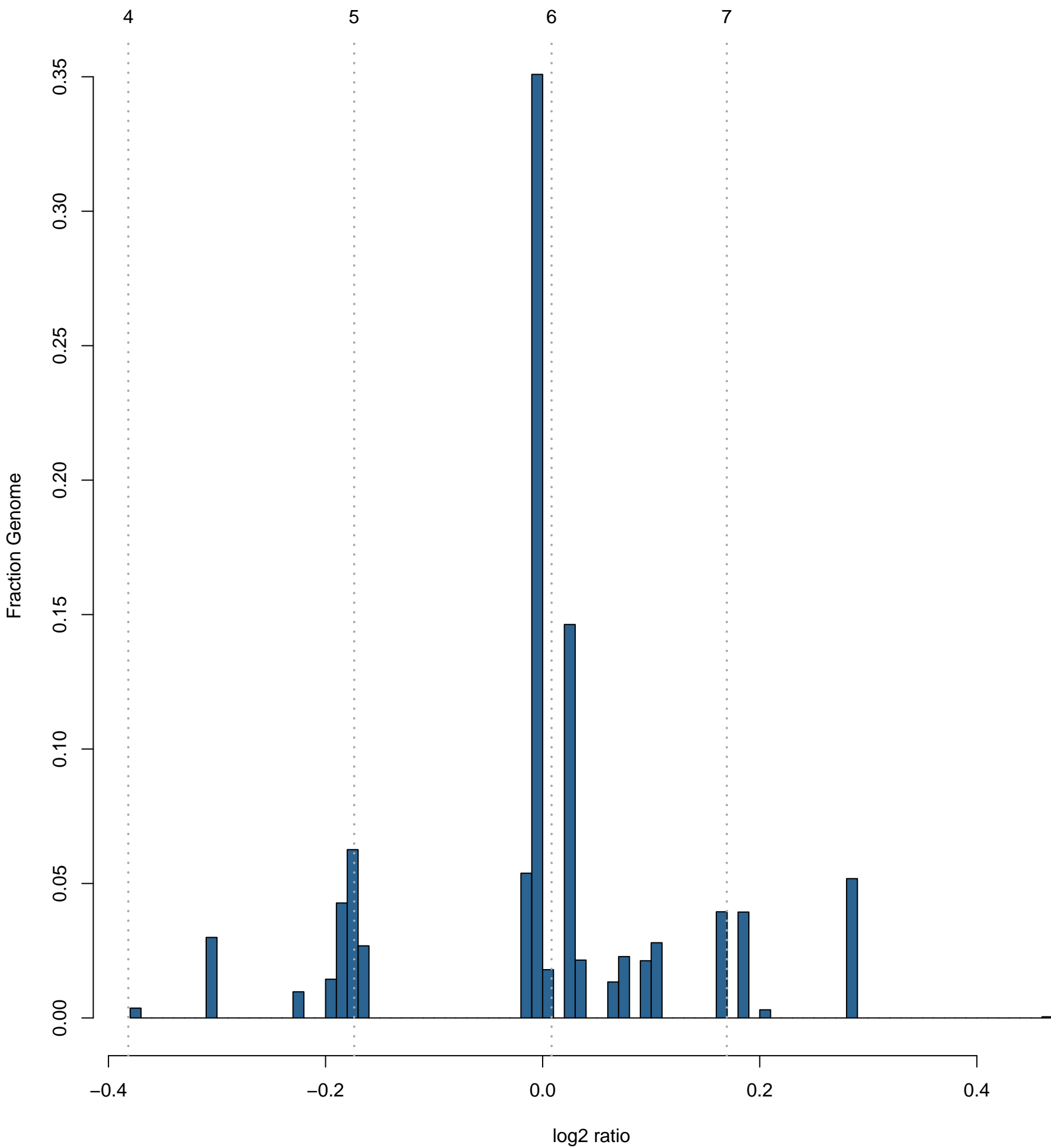


SCNA-fit log-likelihood: -2931.2

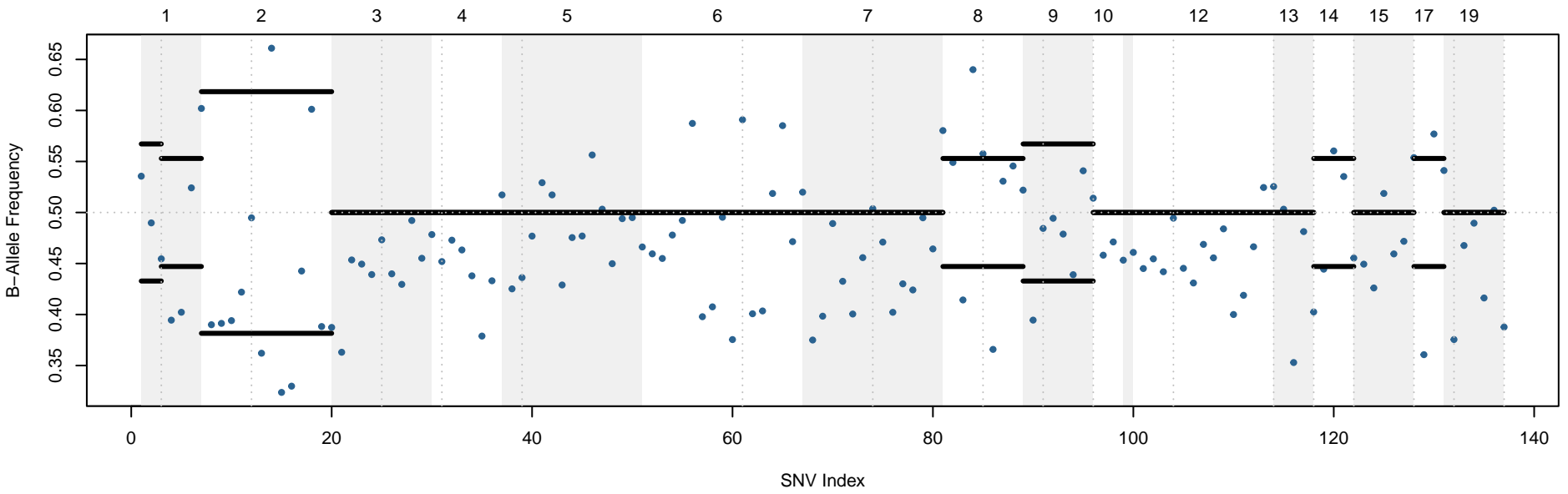




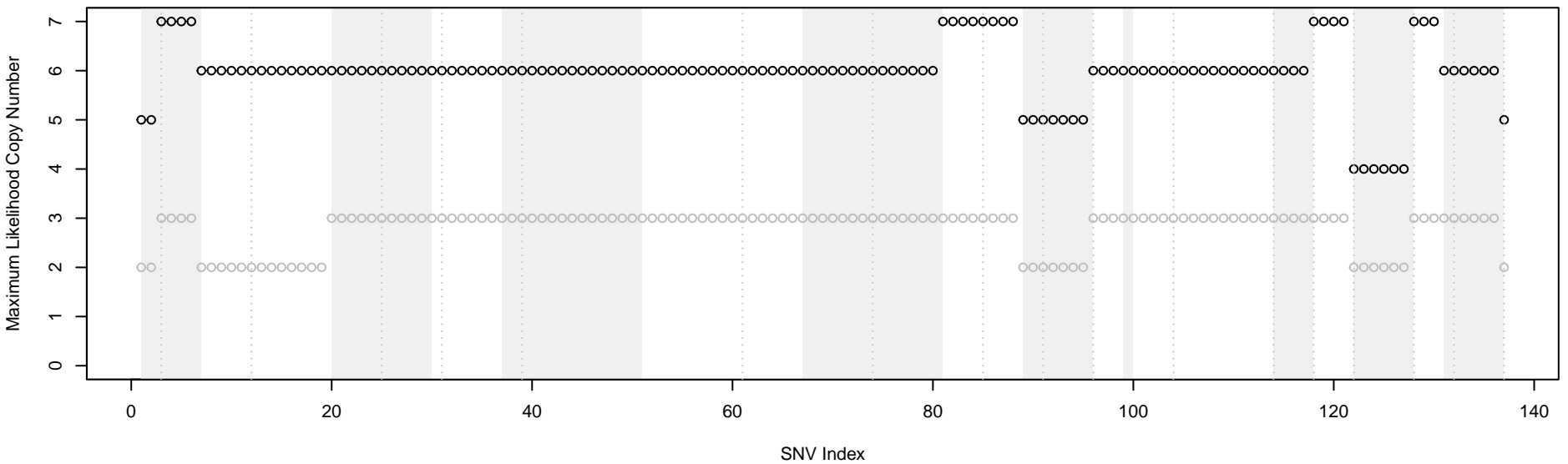
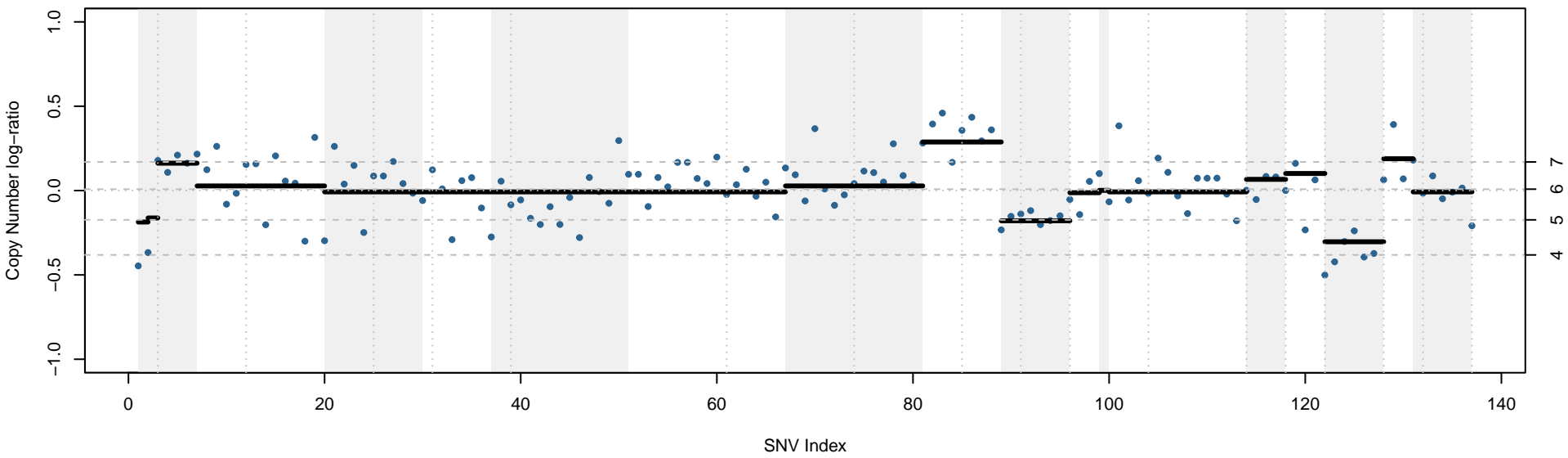
Purity: 0.45 Tumor ploidy: 5.952

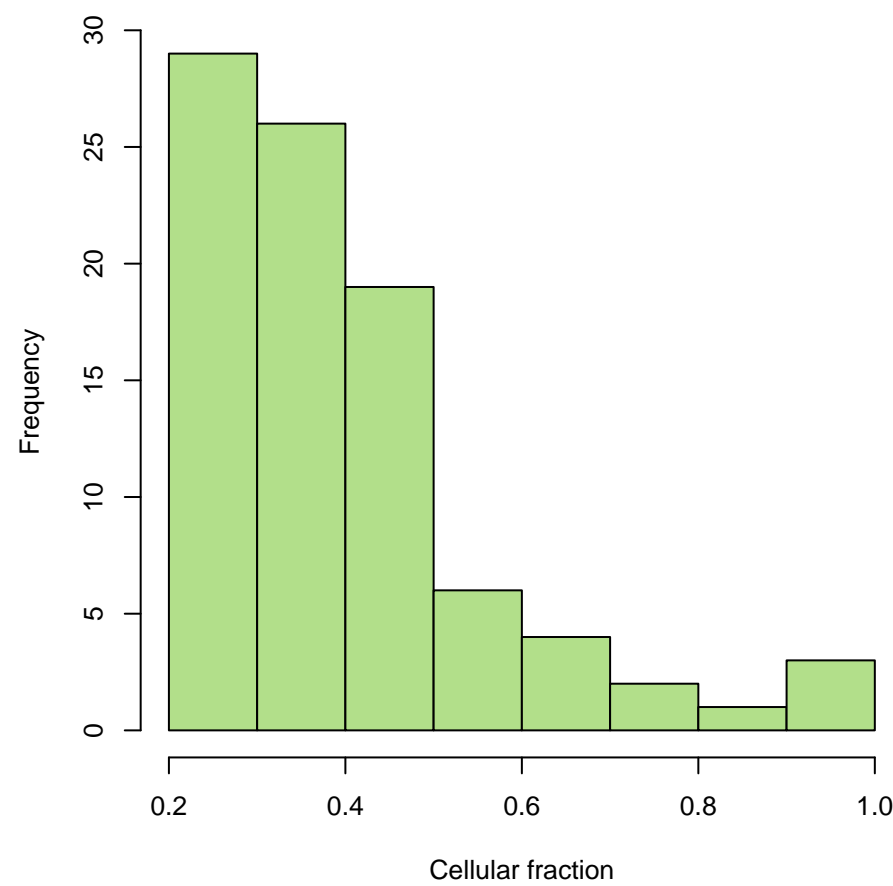
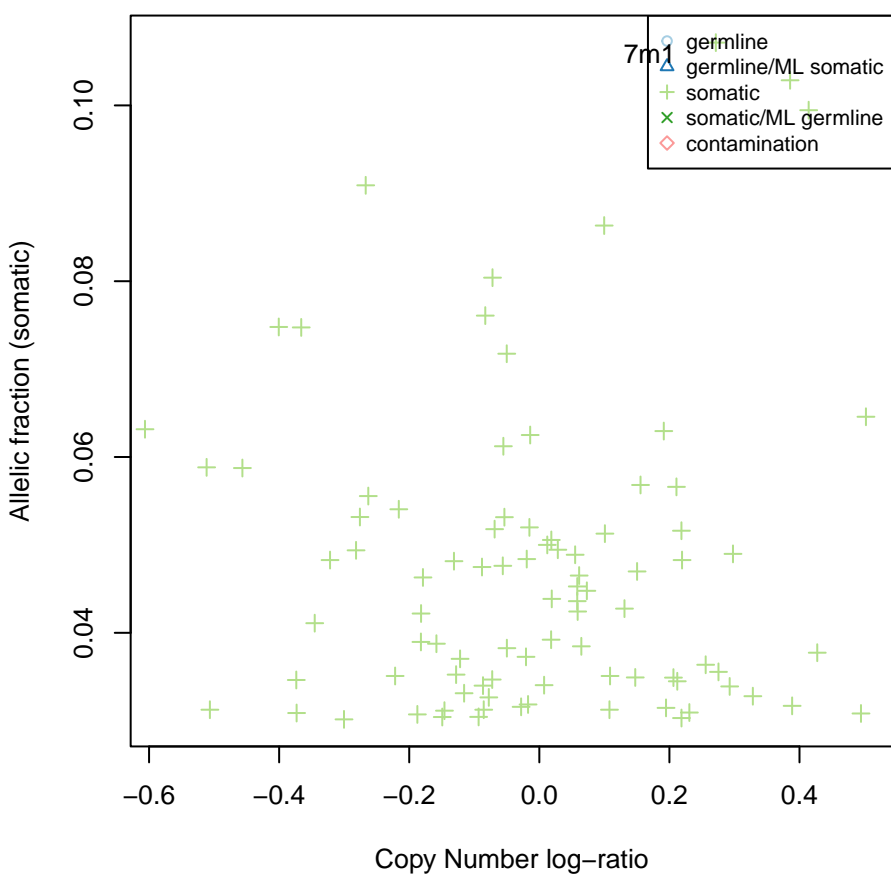
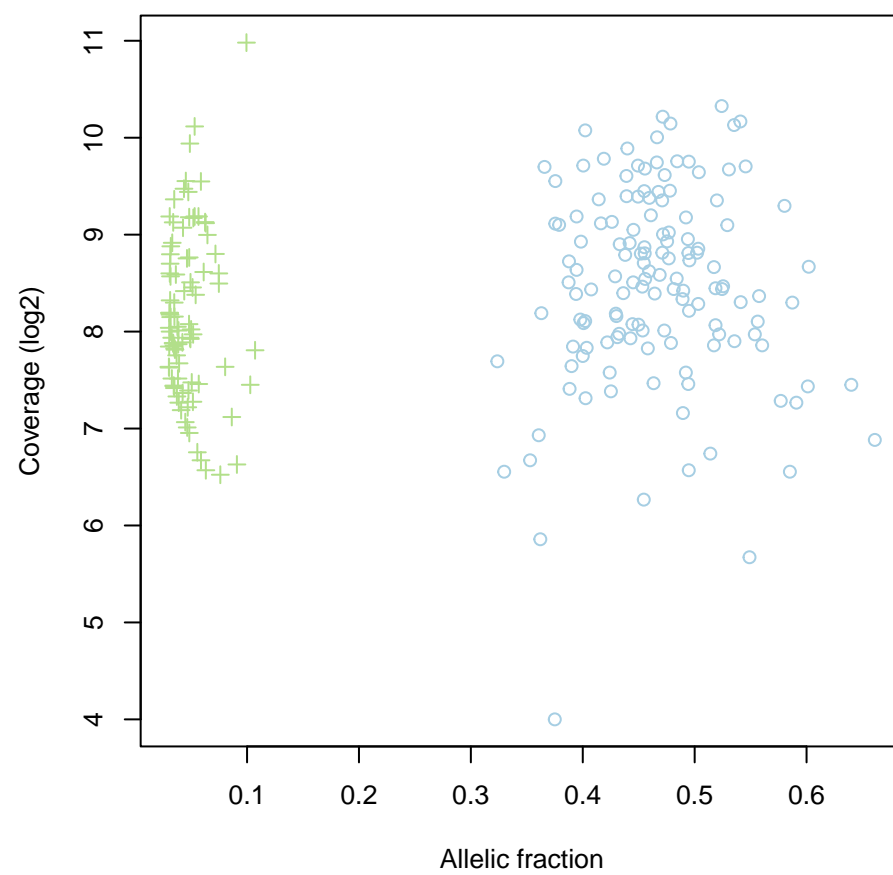
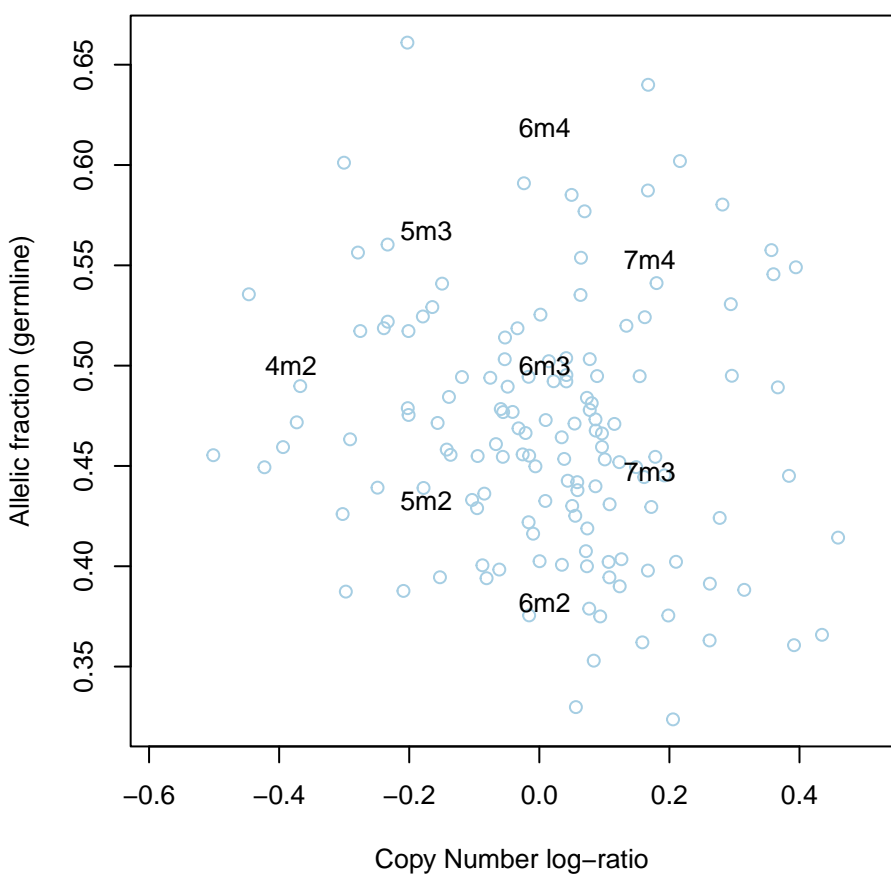


Purity: 0.45 Tumor ploidy: 5.952 SNV log-likelihood: -145.22 GoF: 90.4% Mean coverage: 385,403

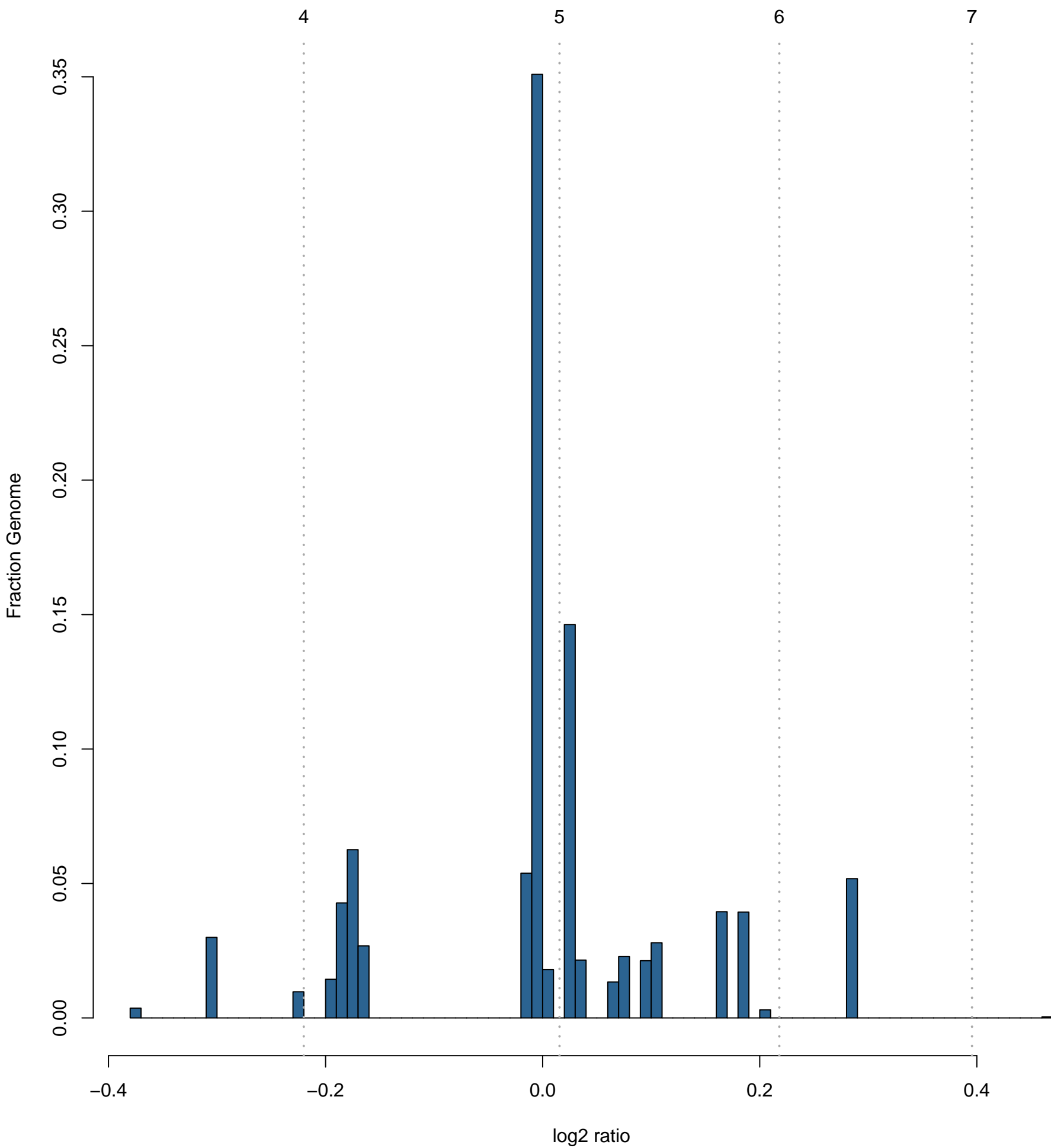


SCNA-fit log-likelihood: -2930.39

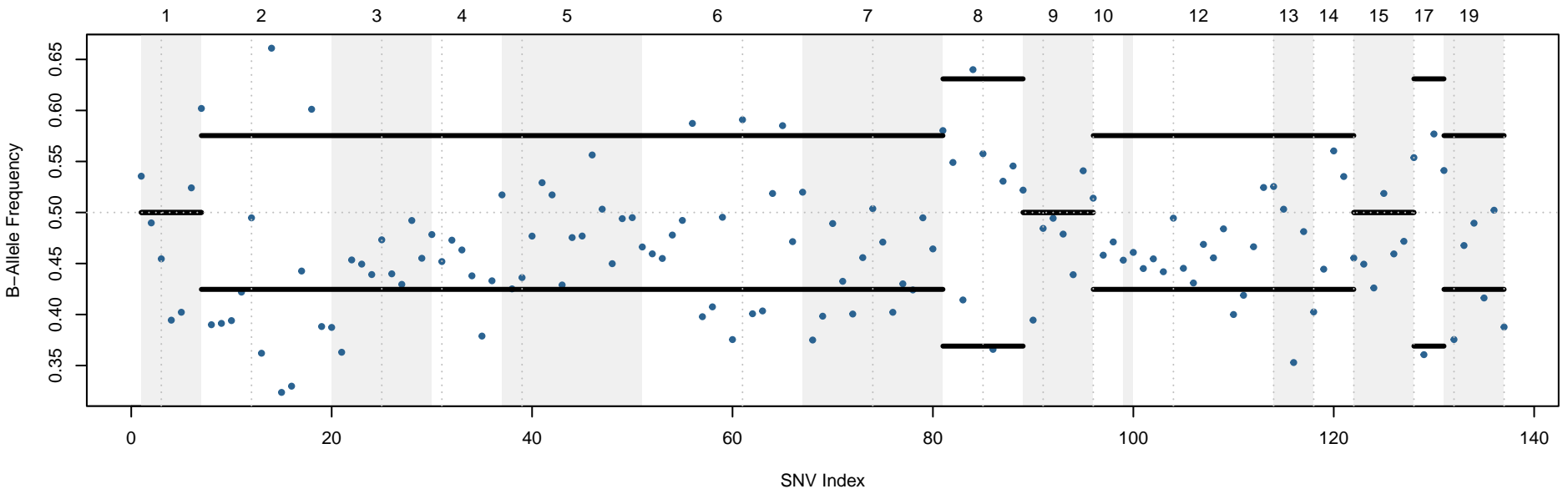




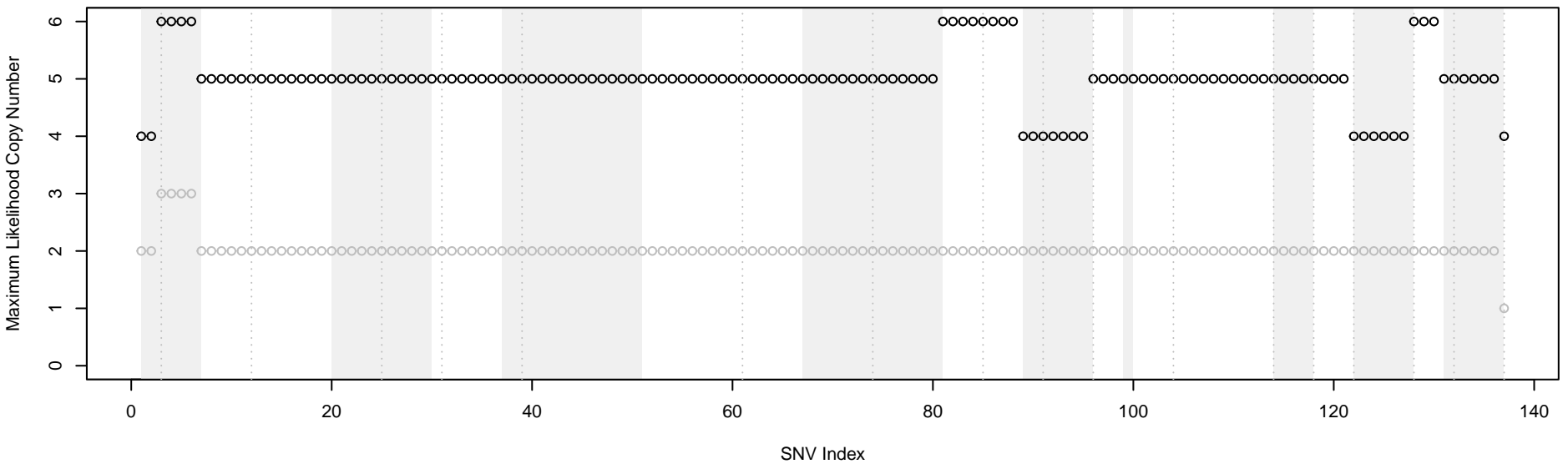
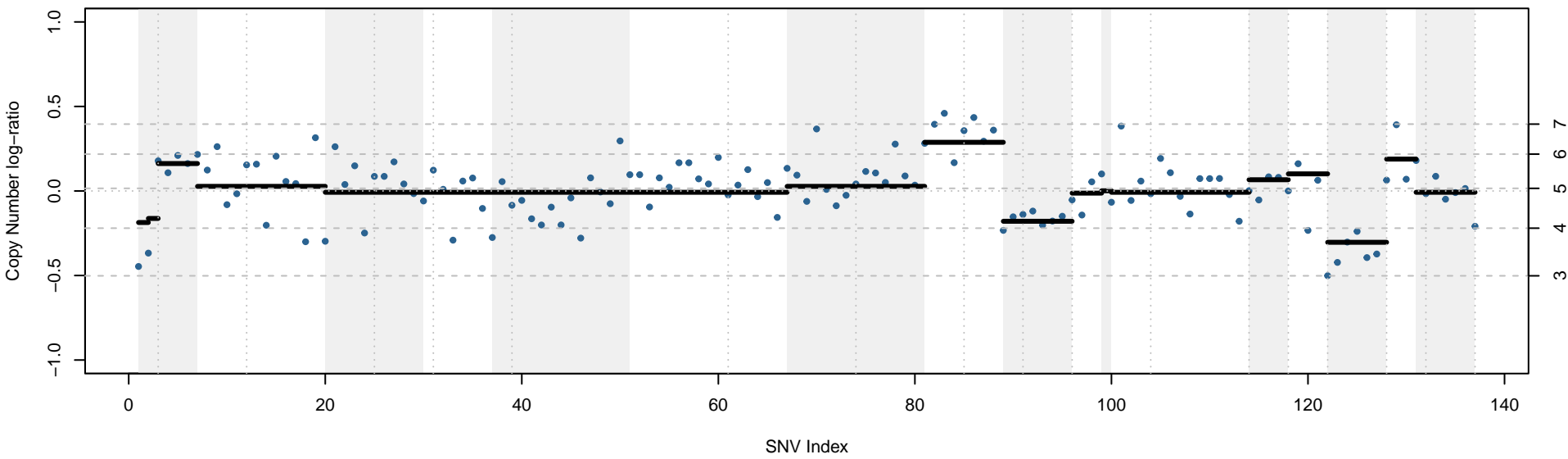
Purity: 0.55 Tumor ploidy: 4.929

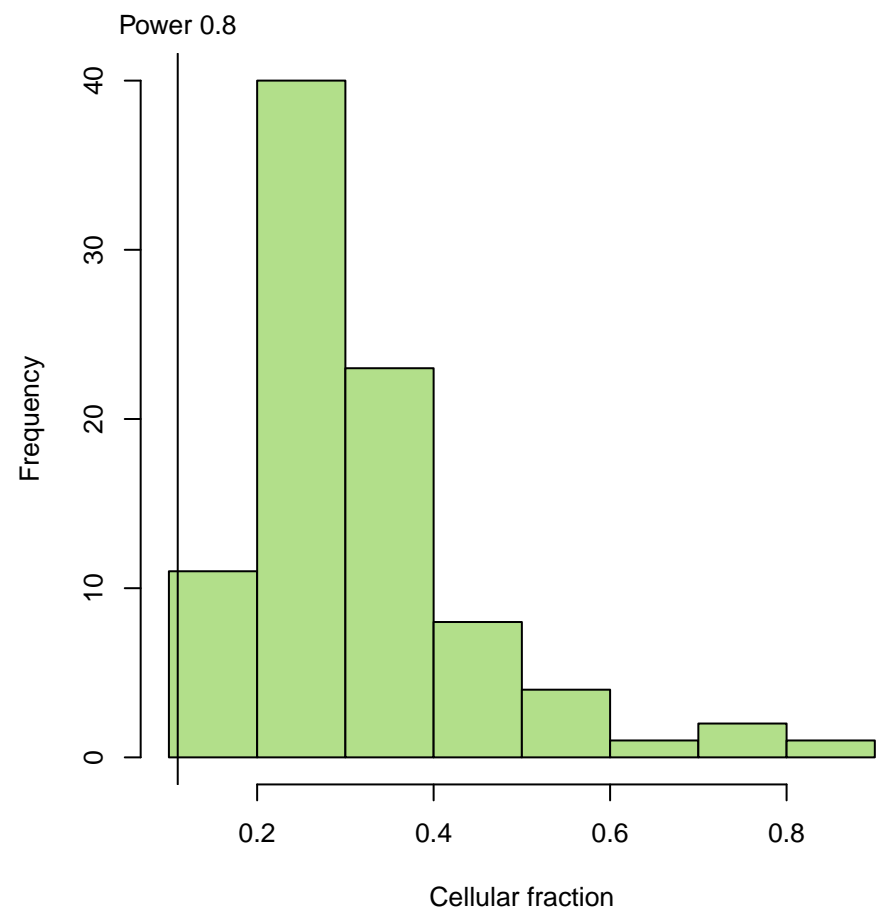
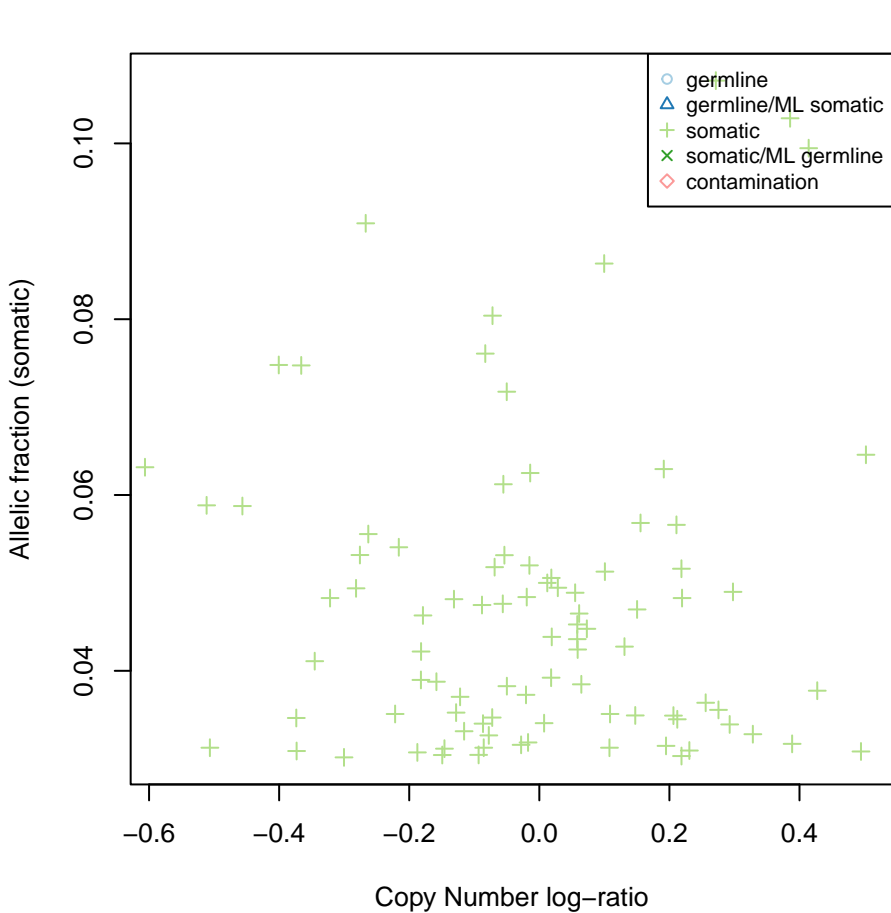
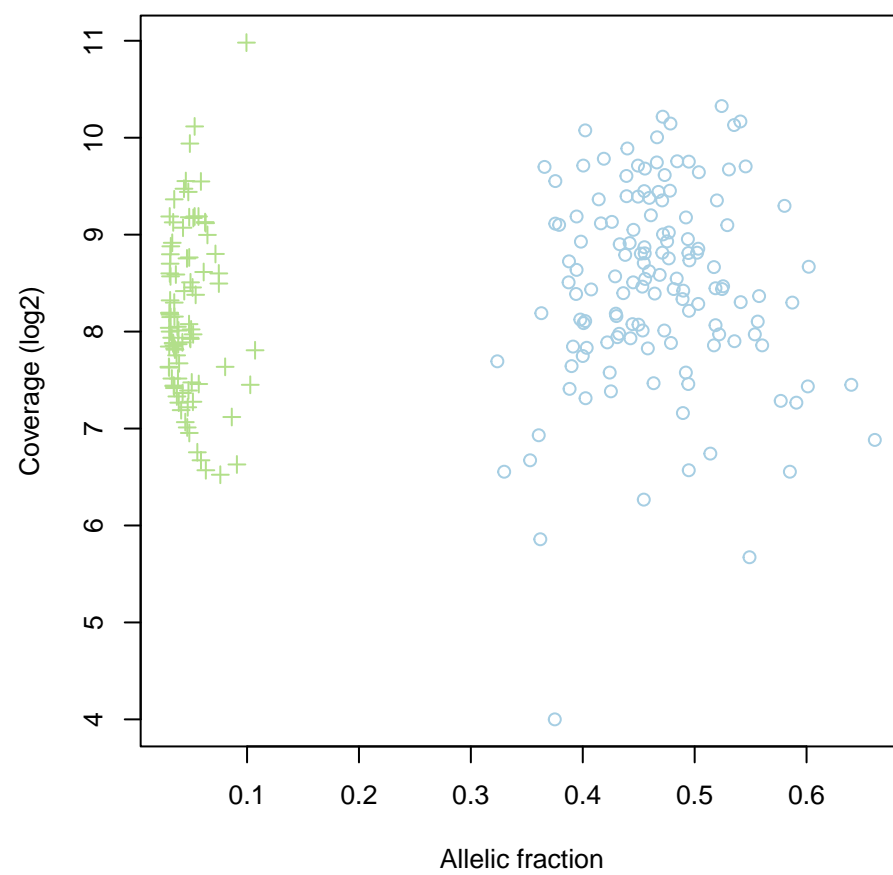
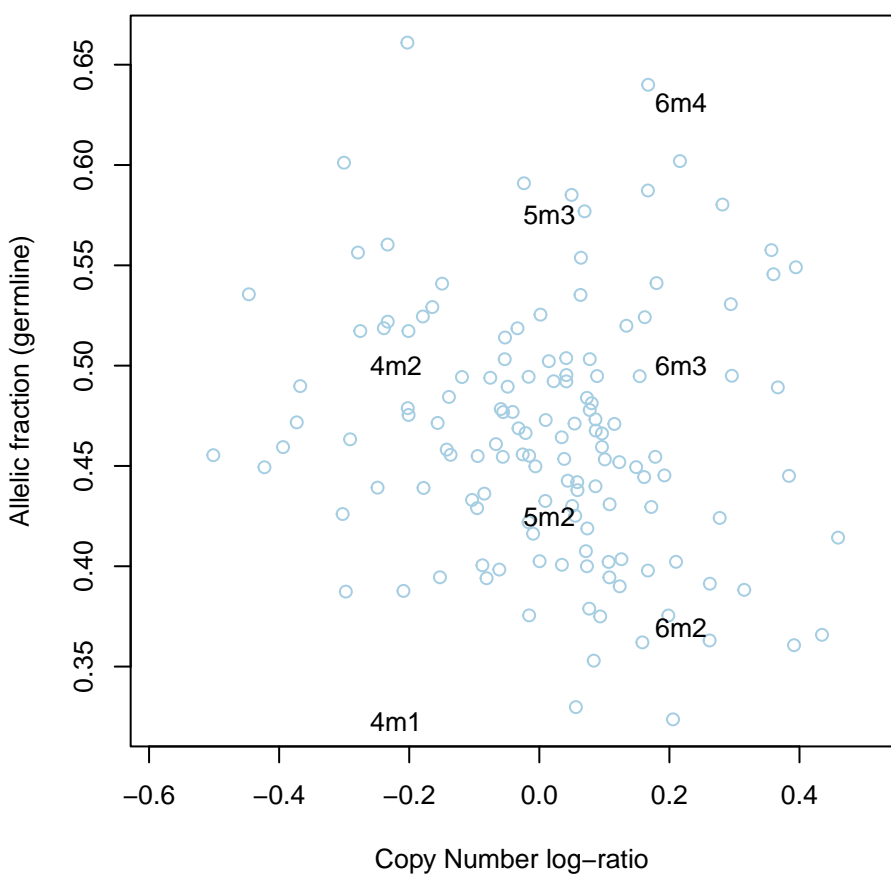


Purity: 0.55 Tumor ploidy: 4.929 SNV log-likelihood: -177.38 GoF: 84.8% Mean coverage: 385,403



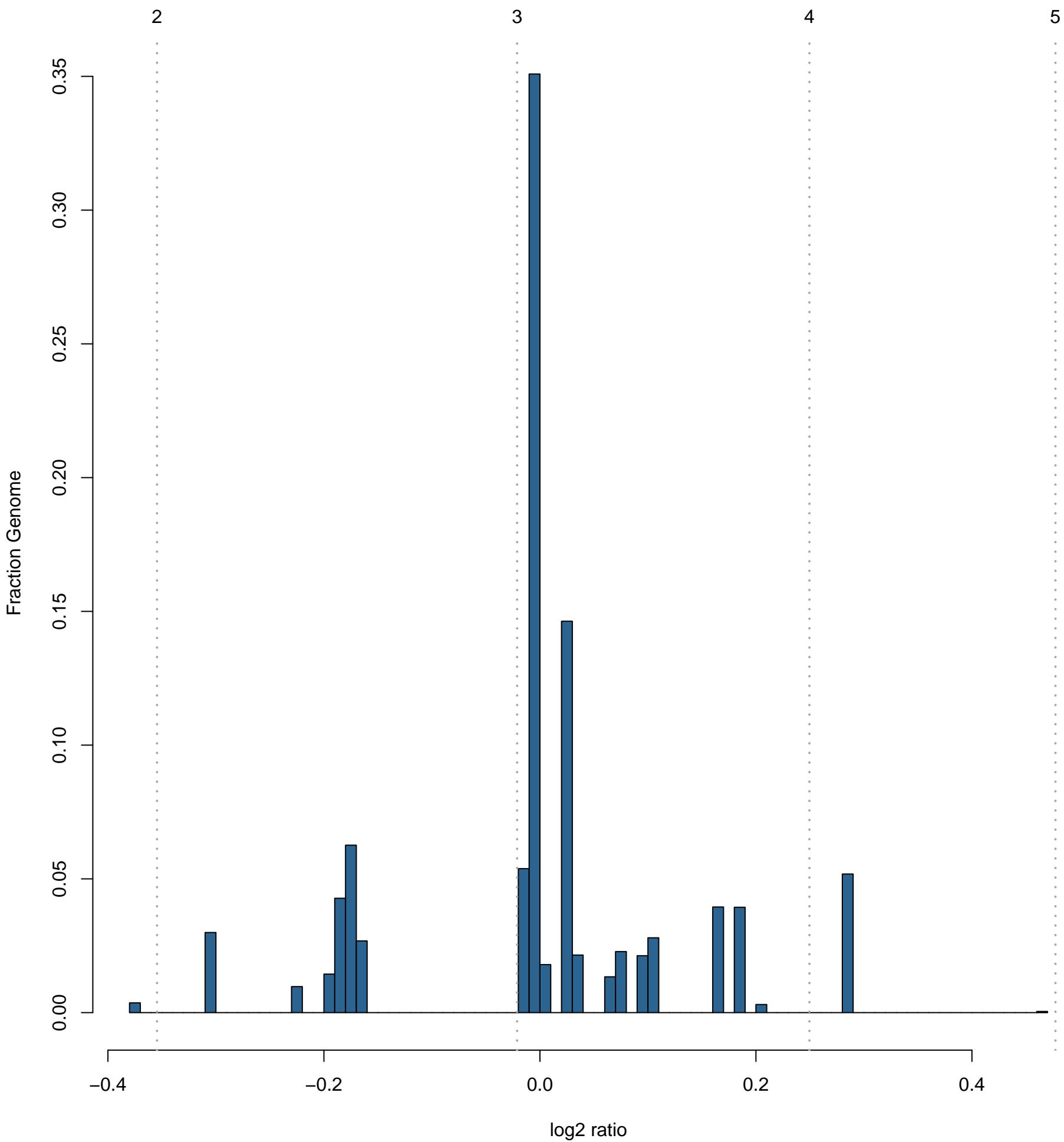
SCNA-fit log-likelihood: -2935.55



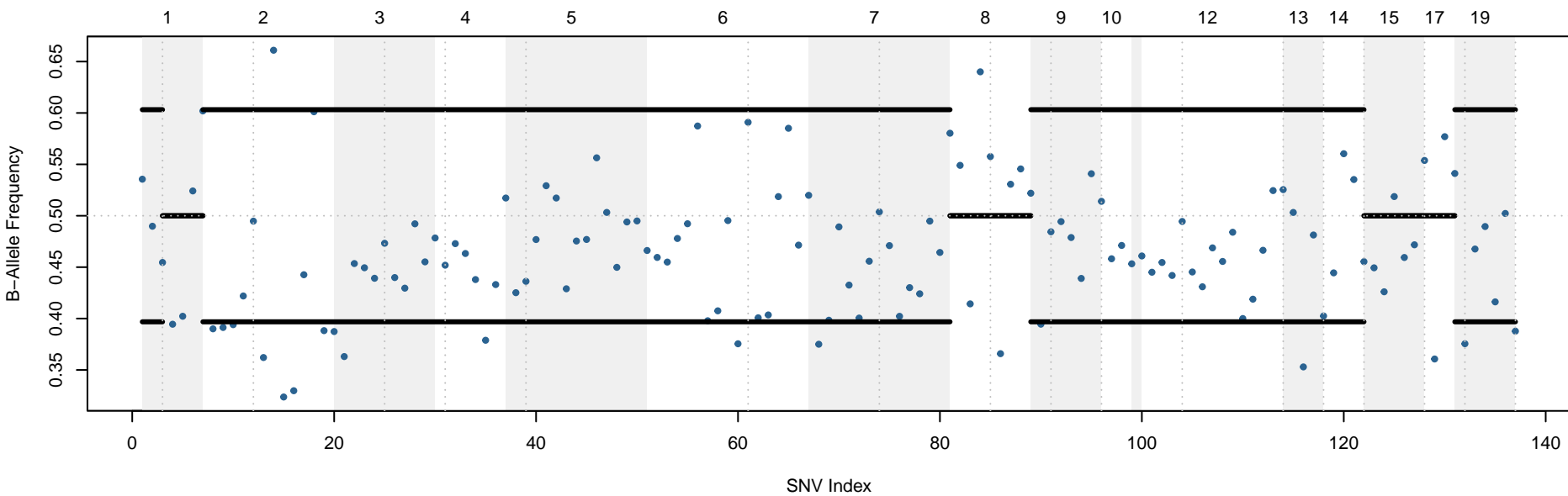




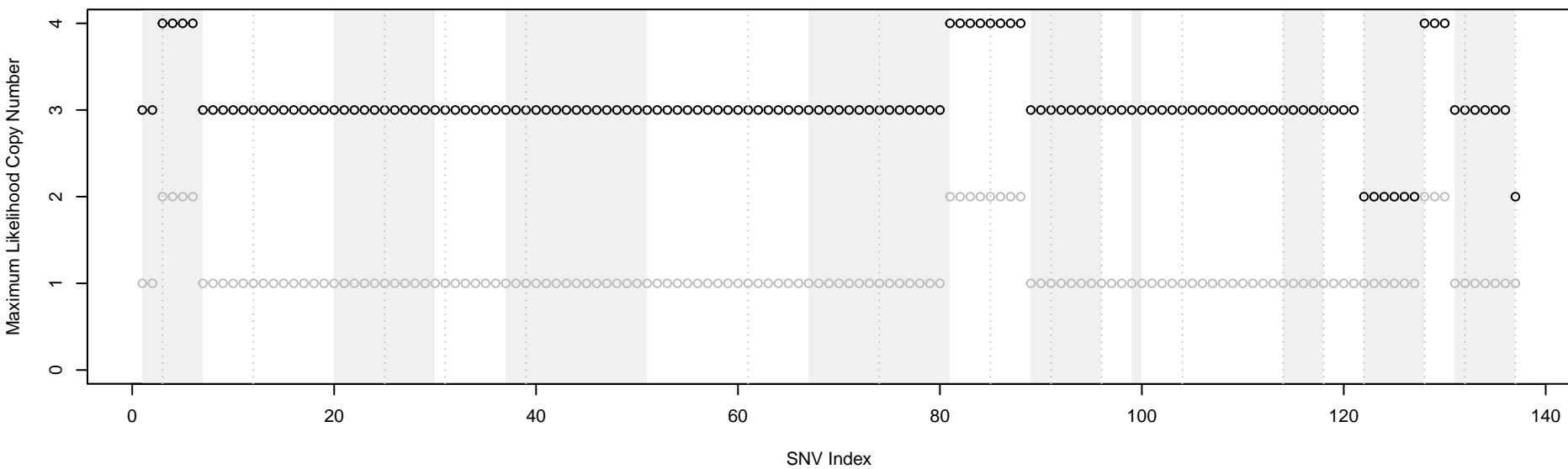
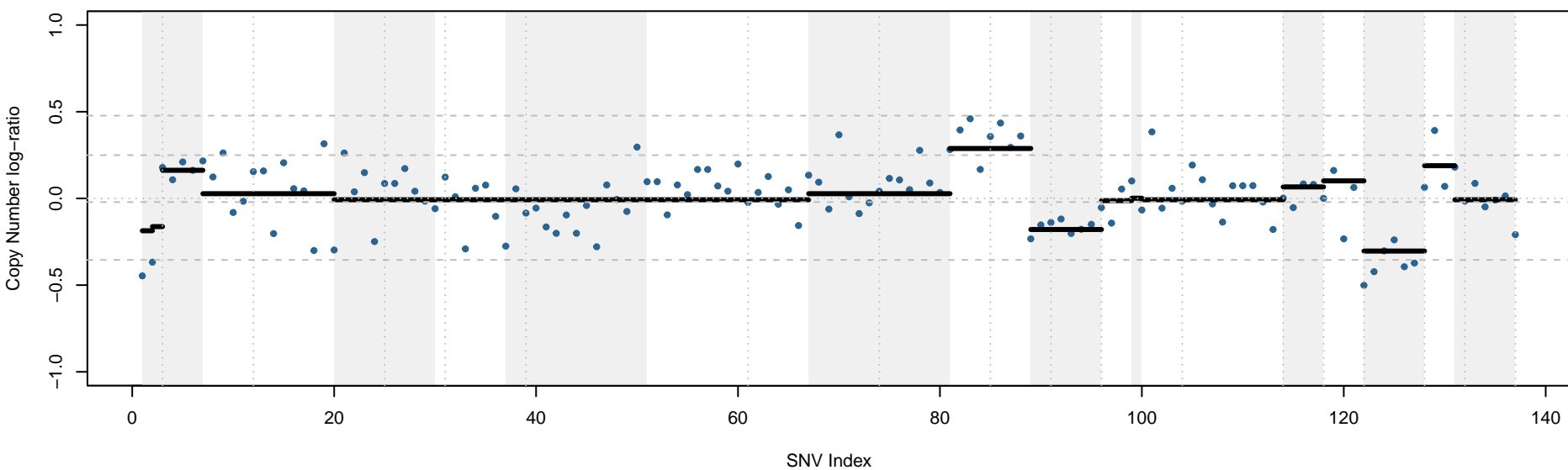
Purity: 0.52 Tumor ploidy: 3.071

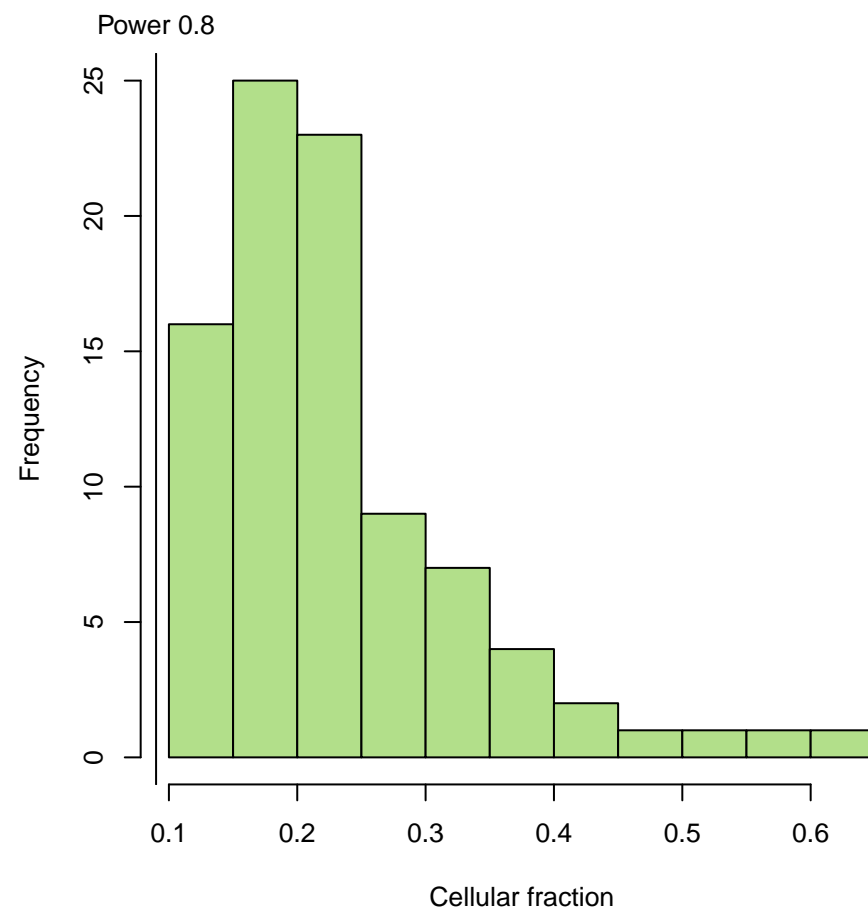
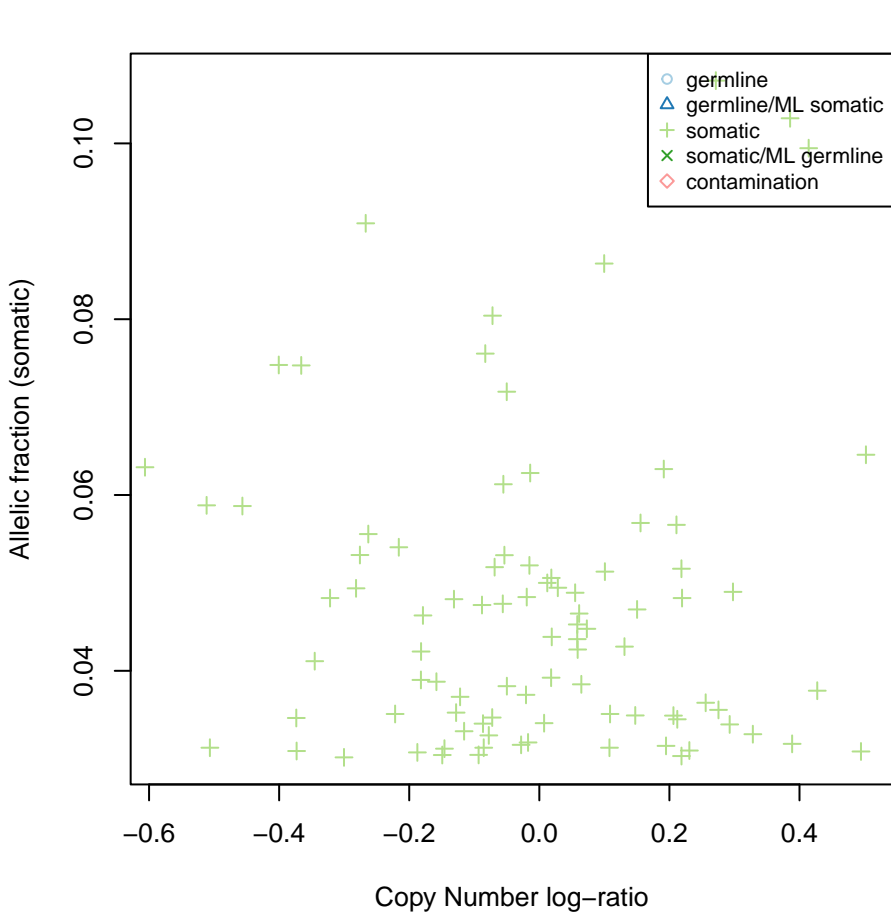
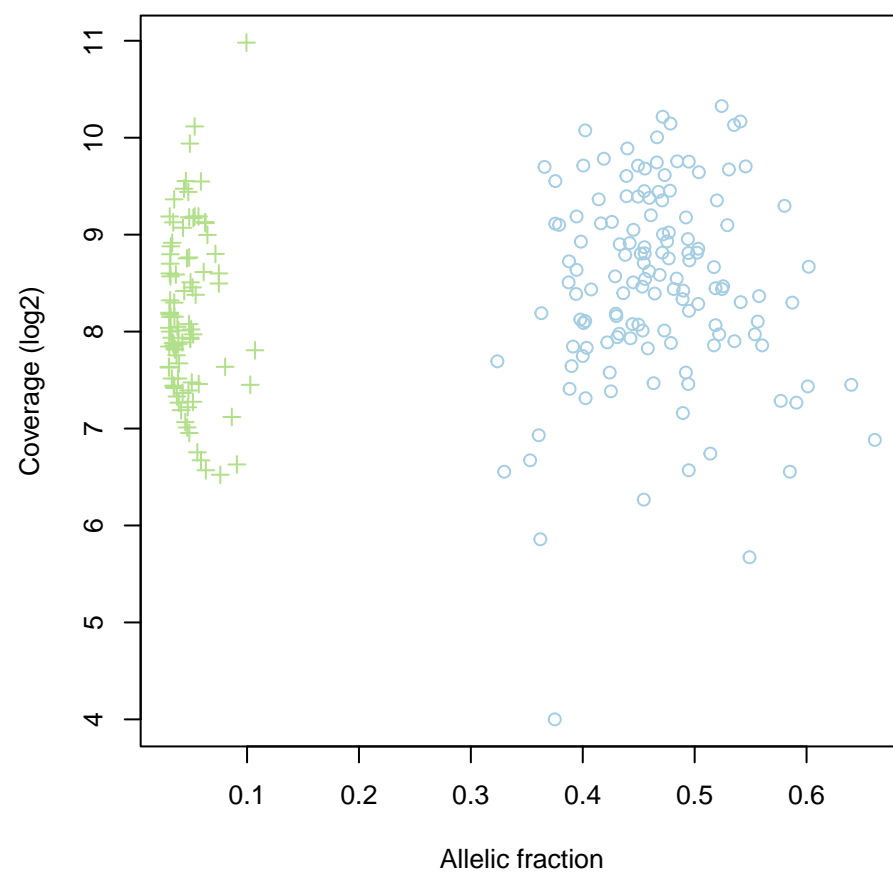
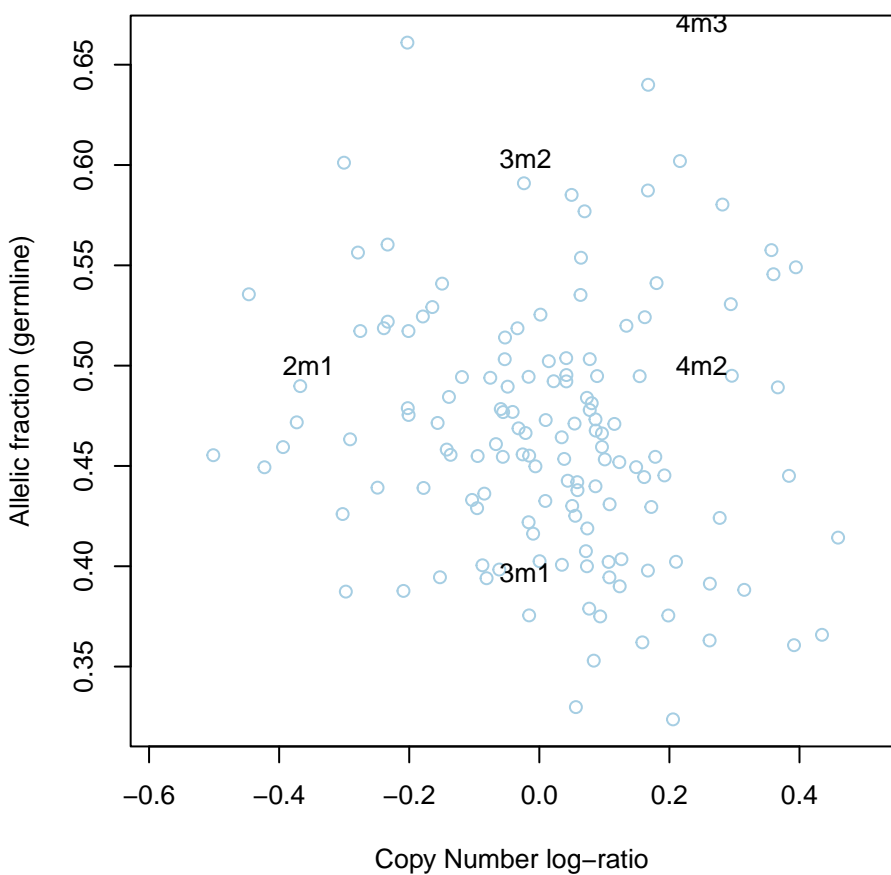


Purity: 0.52 Tumor ploidy: 3.071 SNV log-likelihood: -346.16 GoF: 68% Mean coverage: 385;403

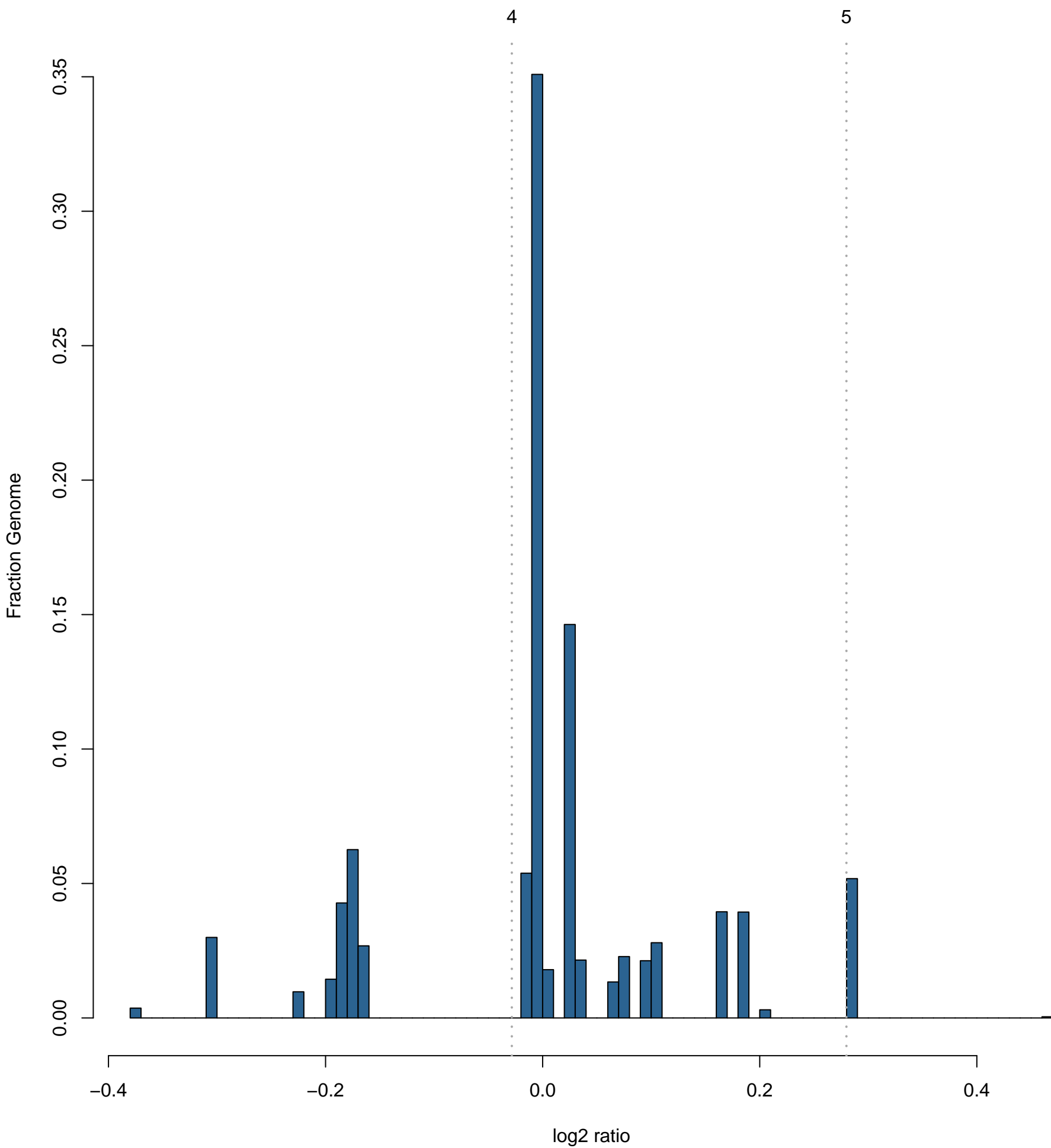


SCNA-fit log-likelihood: -3471.79

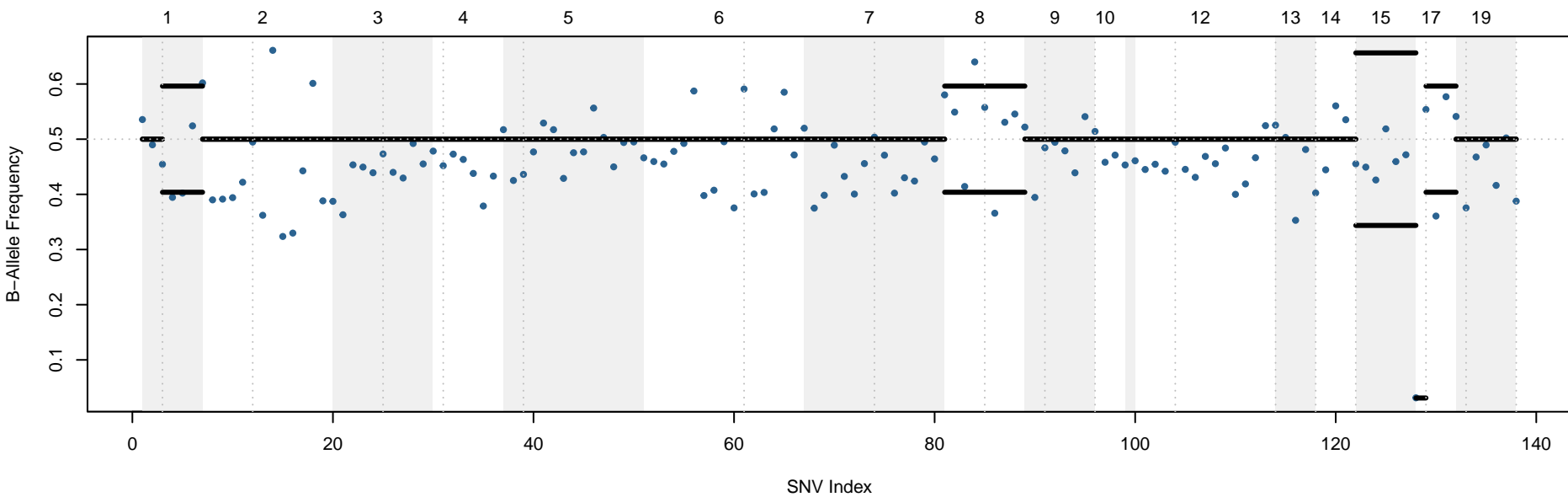




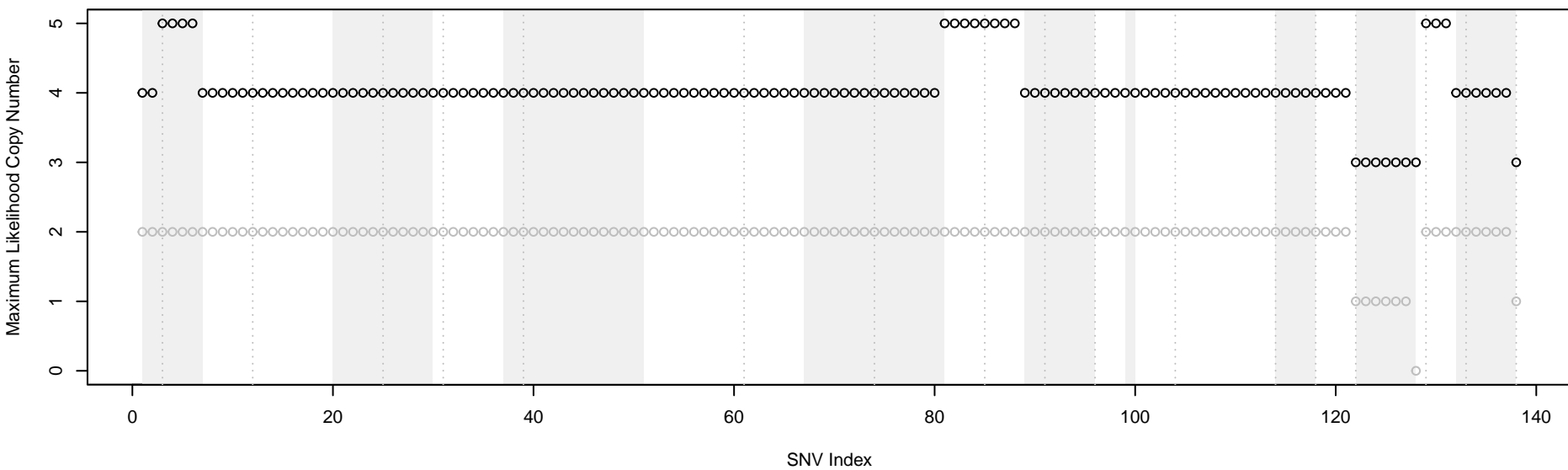
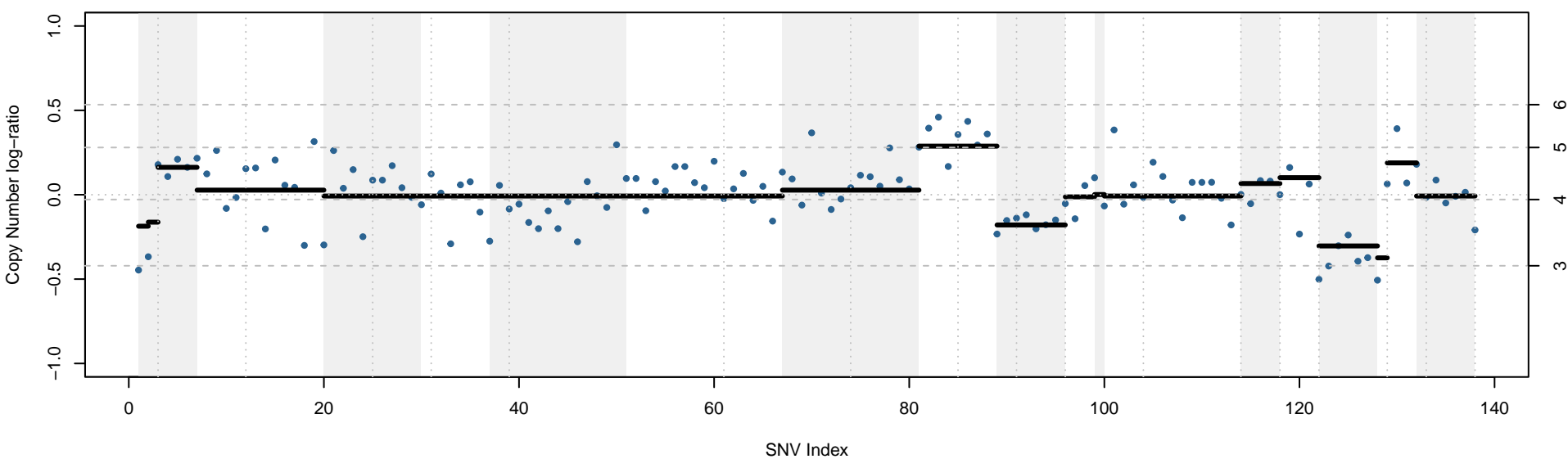
Purity: 0.91 Tumor ploidy: 4.083

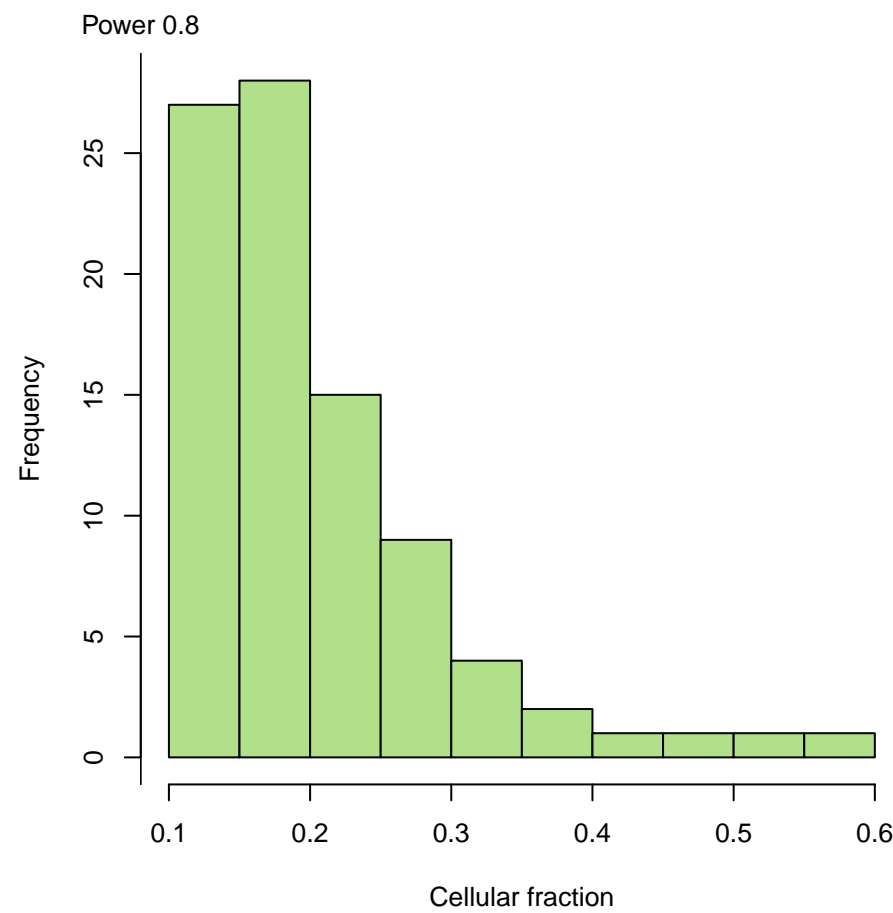
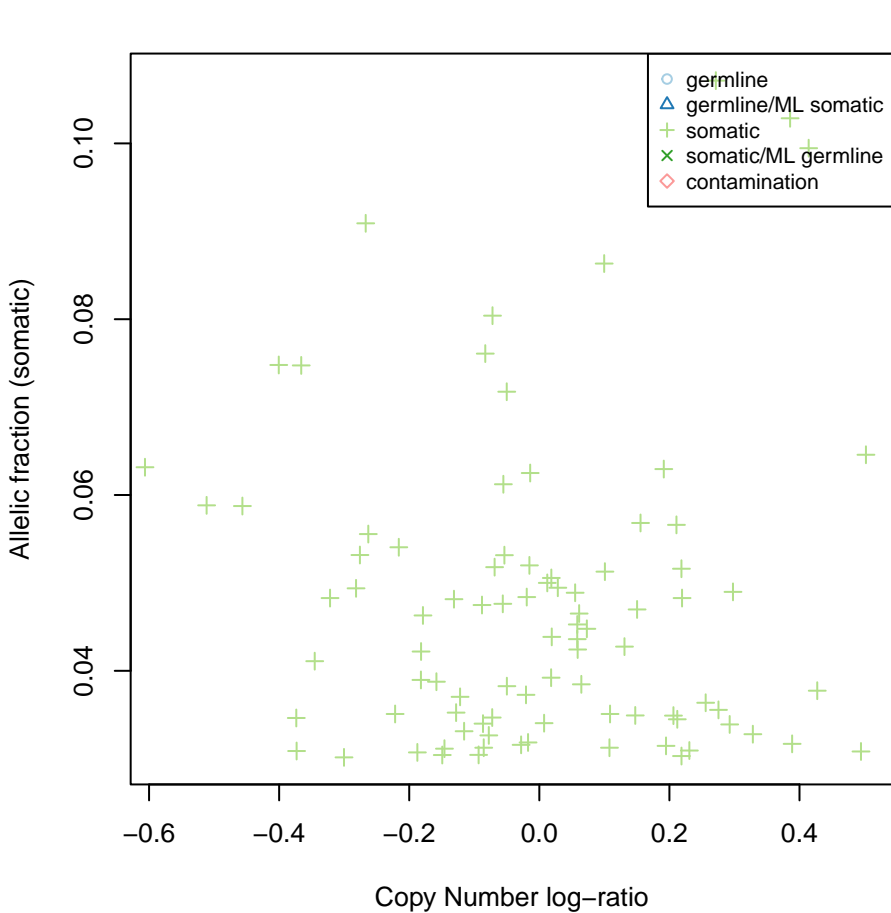
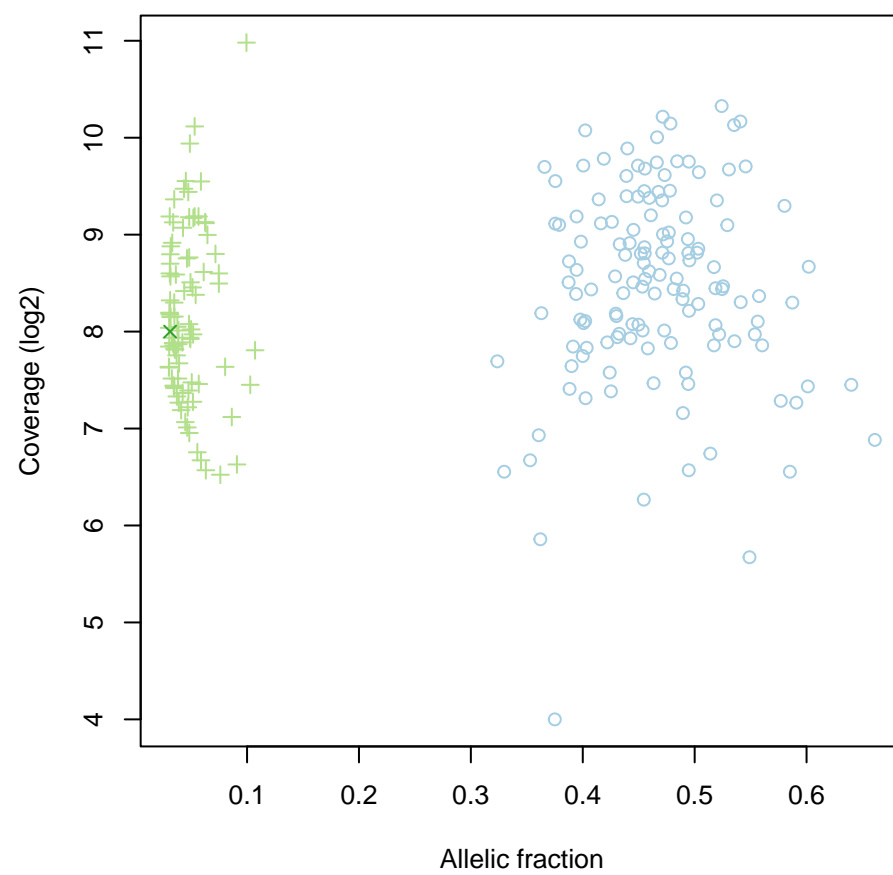
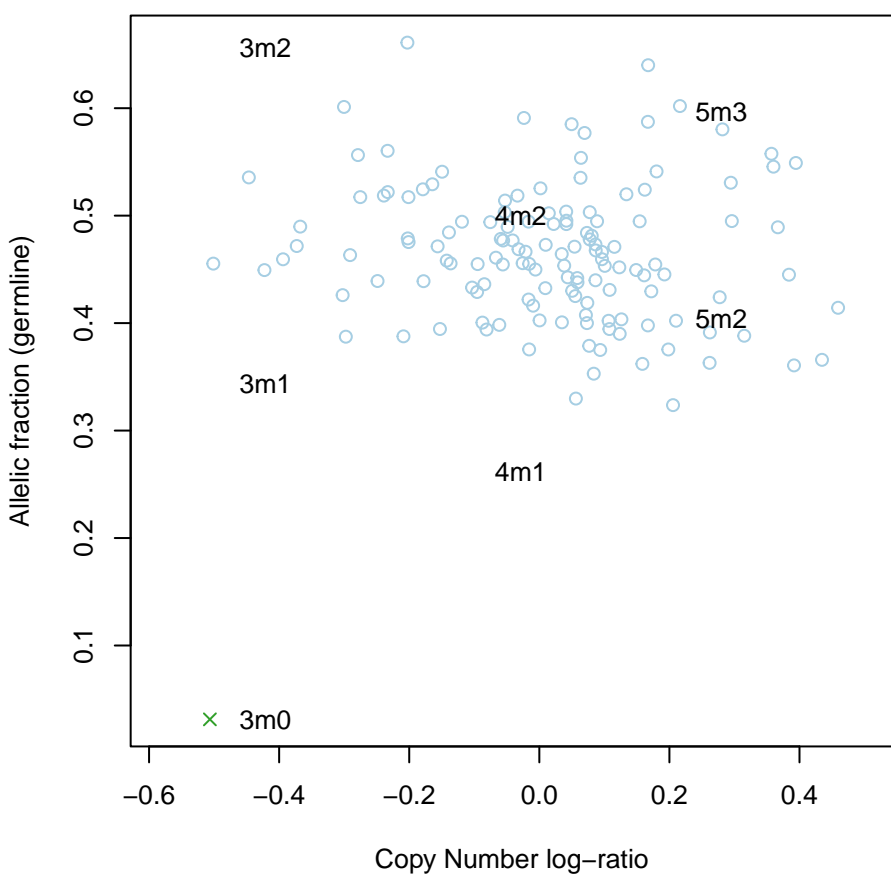


Purity: 0.91 Tumor ploidy: 4.083 SNV log-likelihood: -366.77 GoF: 58% Mean coverage: 385;403

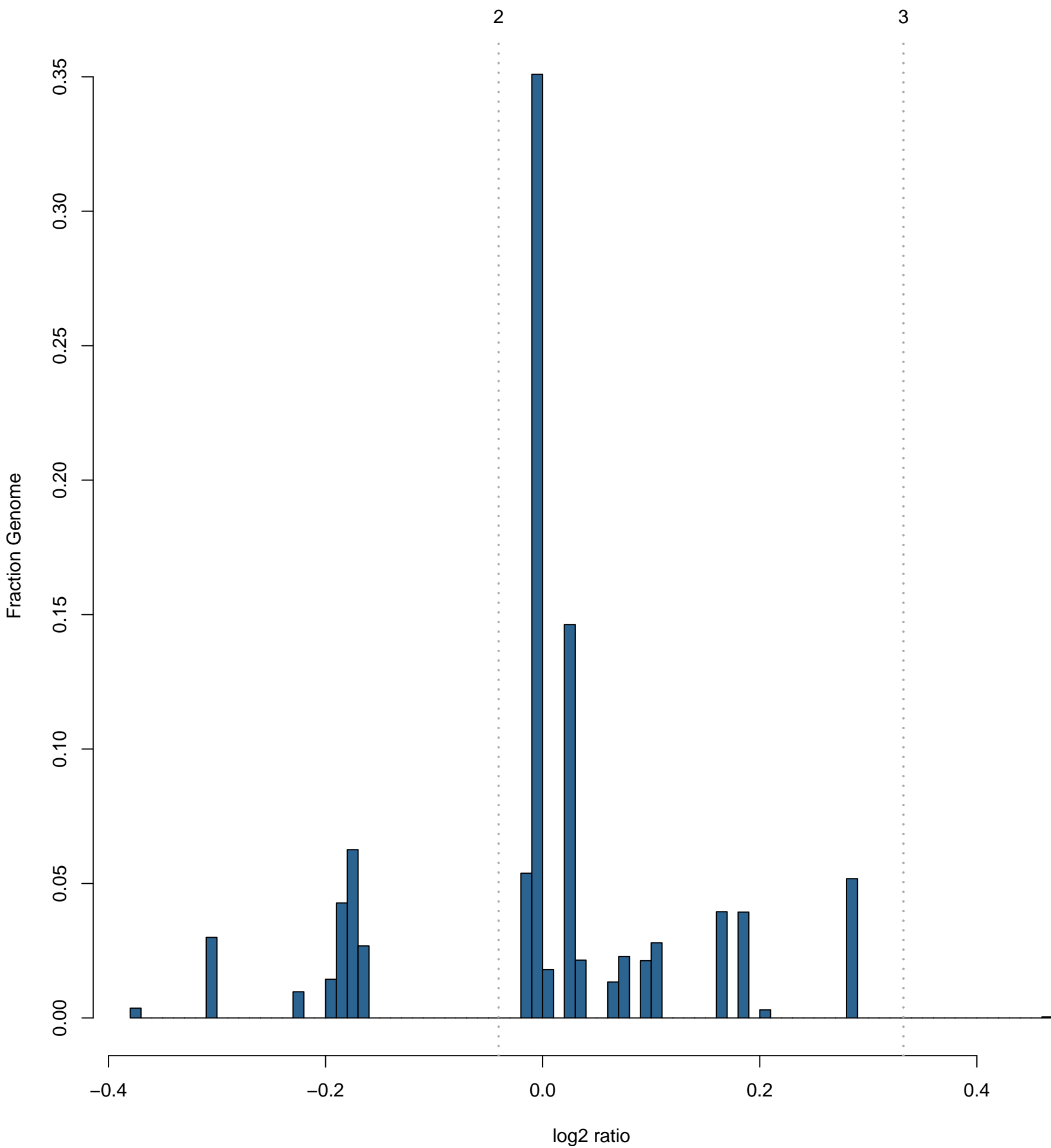


SCNA-fit log-likelihood: -3602.89

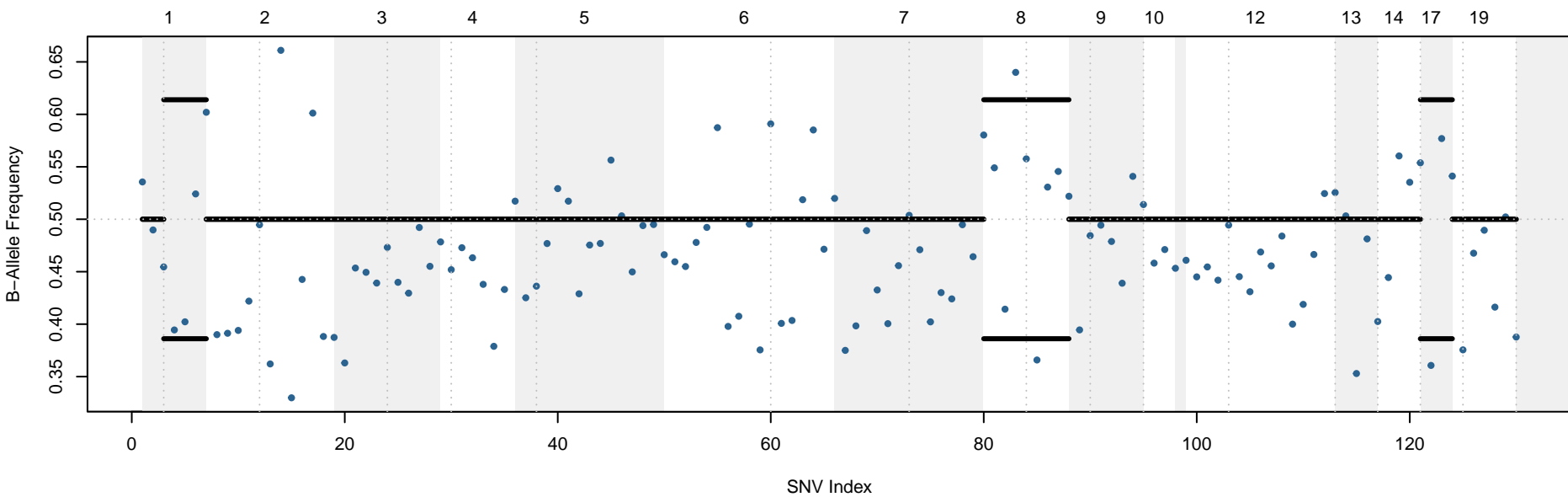




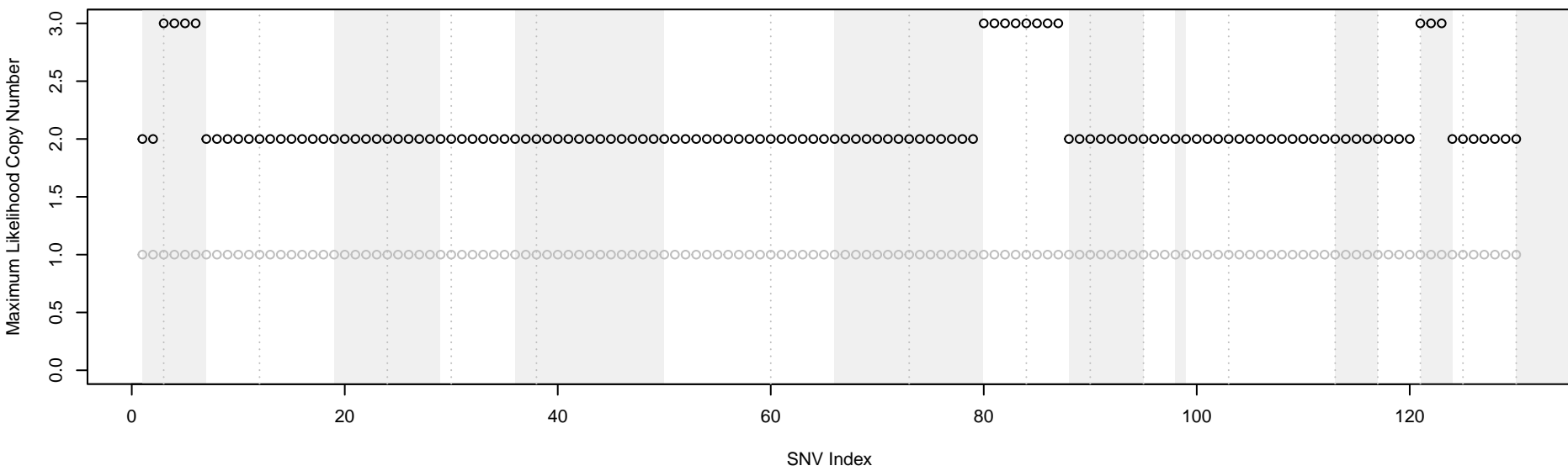
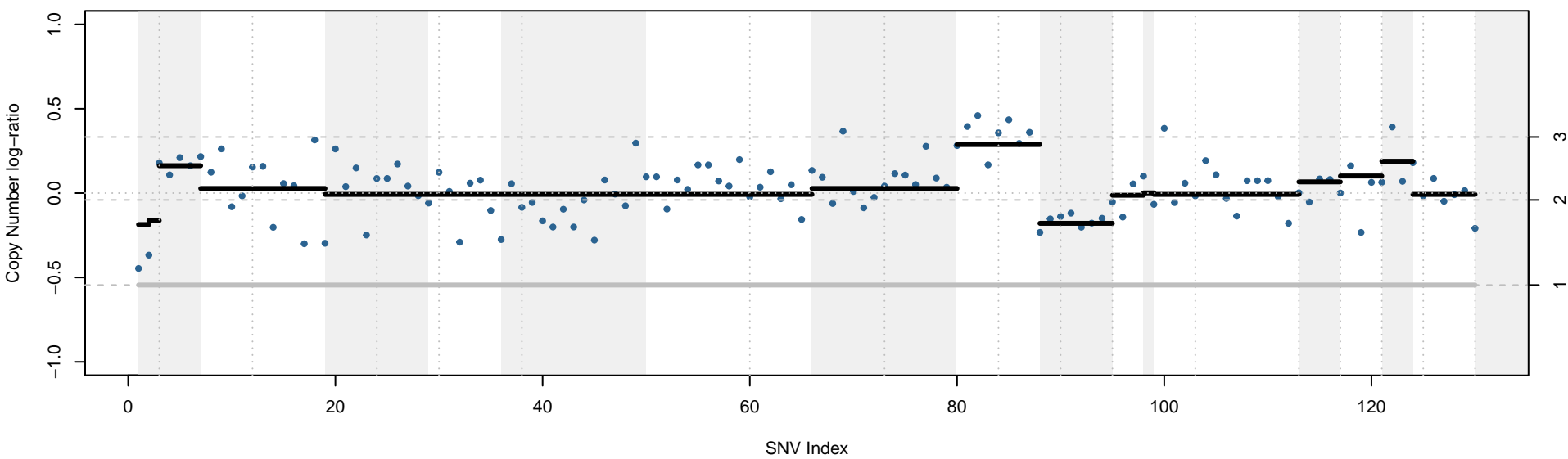
Purity: 0.59 Tumor ploidy: 2.097



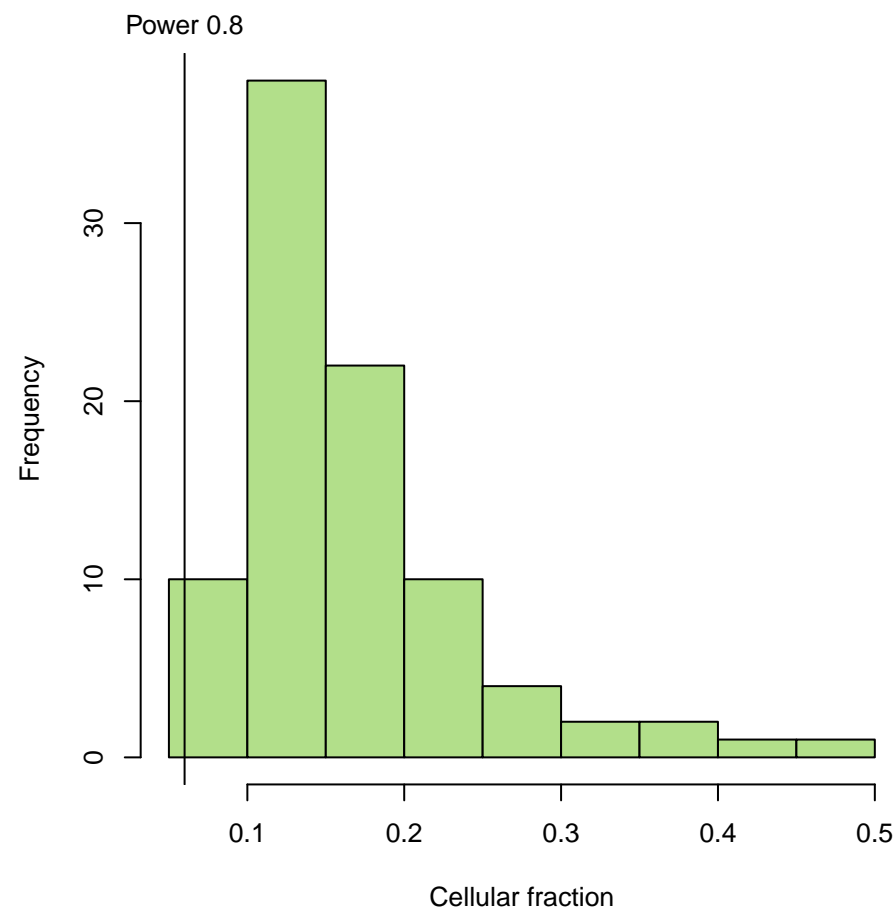
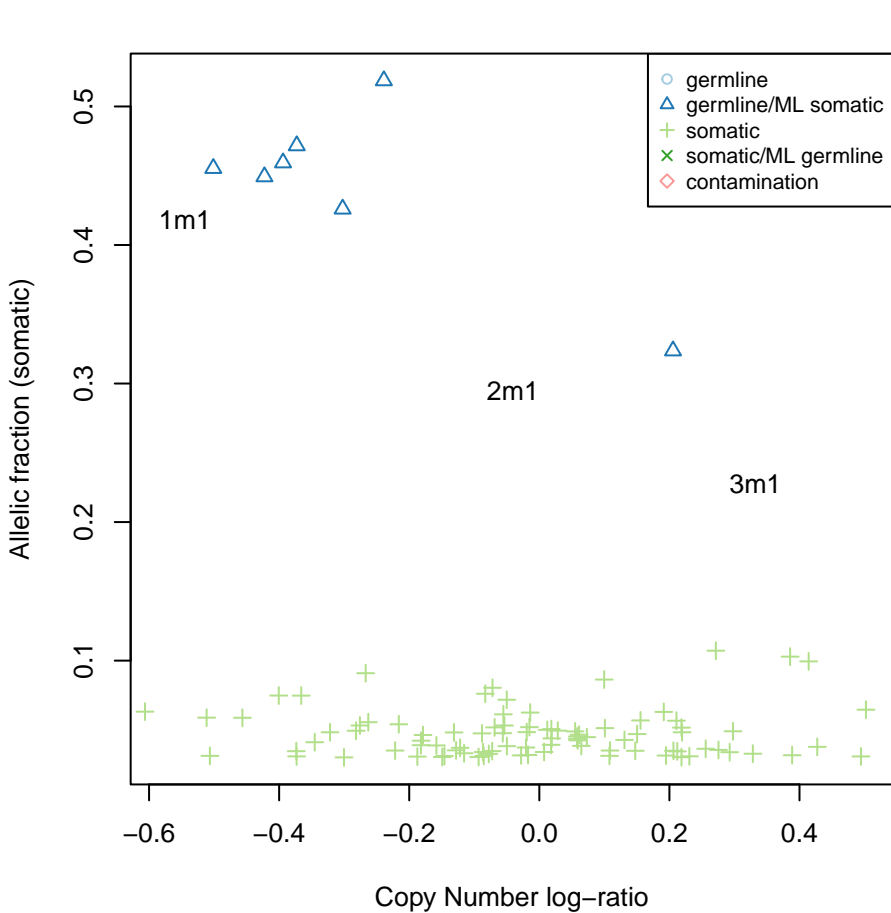
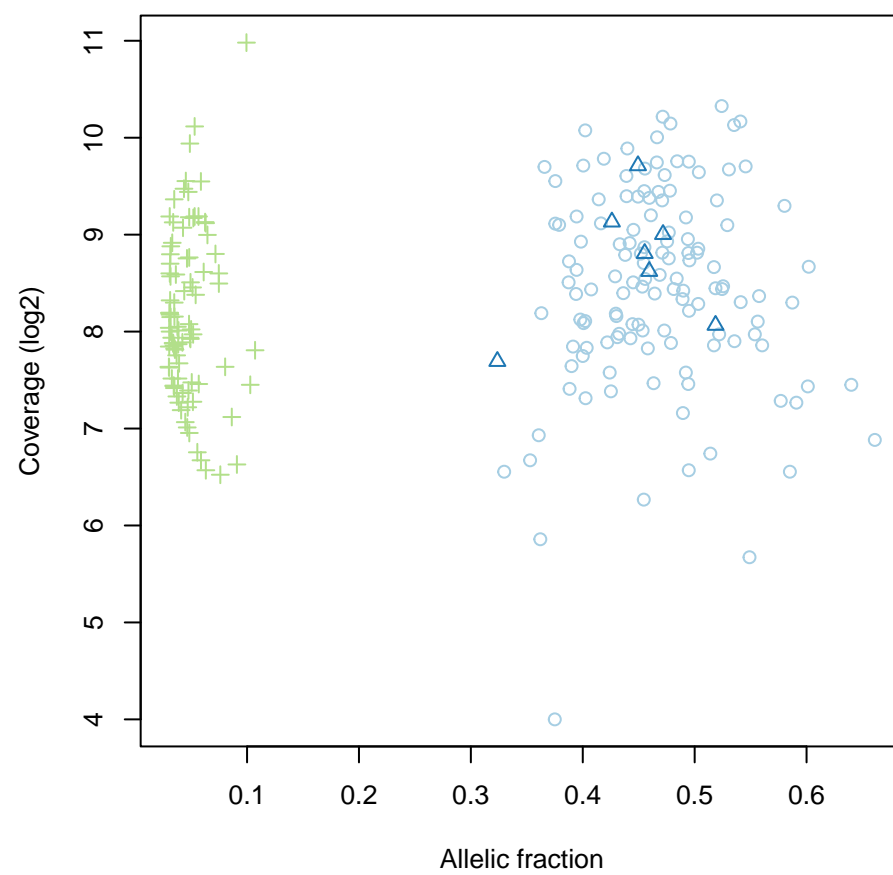
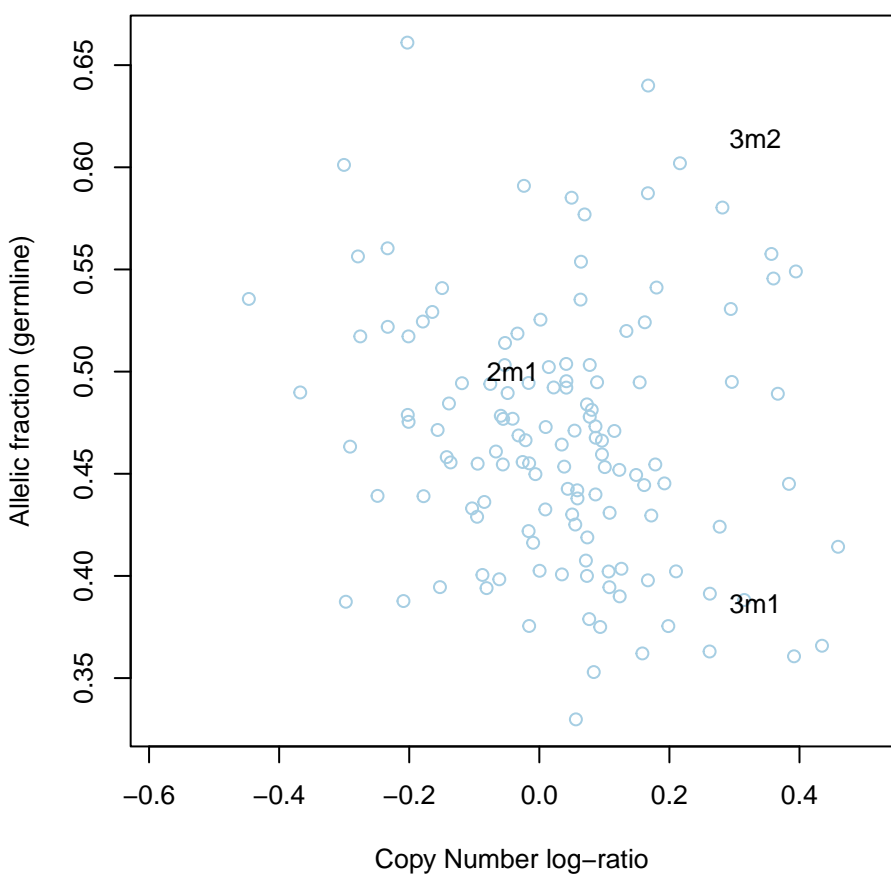
Purity: 0.59 Tumor ploidy: 2.097 SNV log-likelihood: -455.89 GoF: 31.7% Mean coverage: 385,403



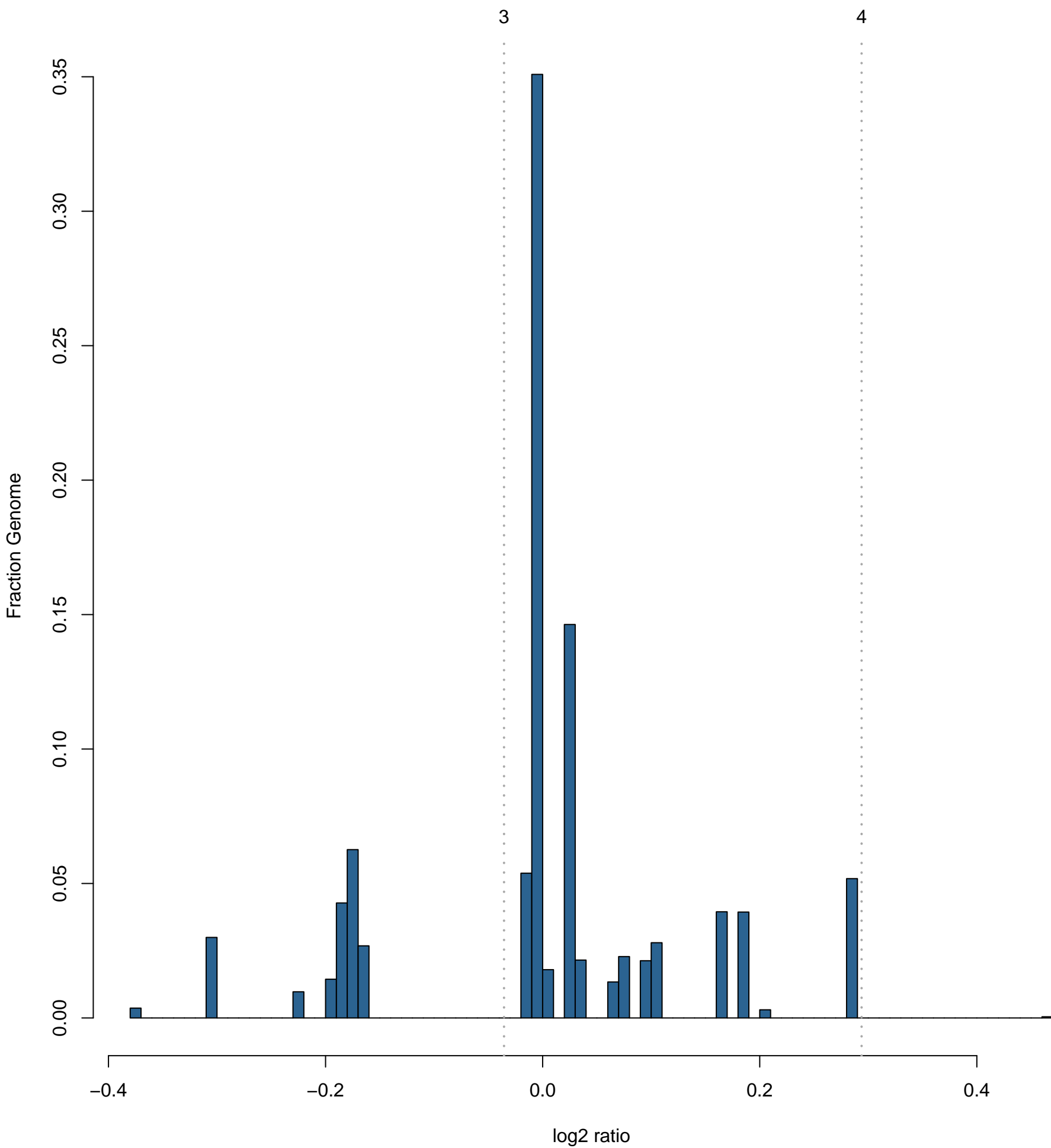
SCNA-fit log-likelihood: -3960.96



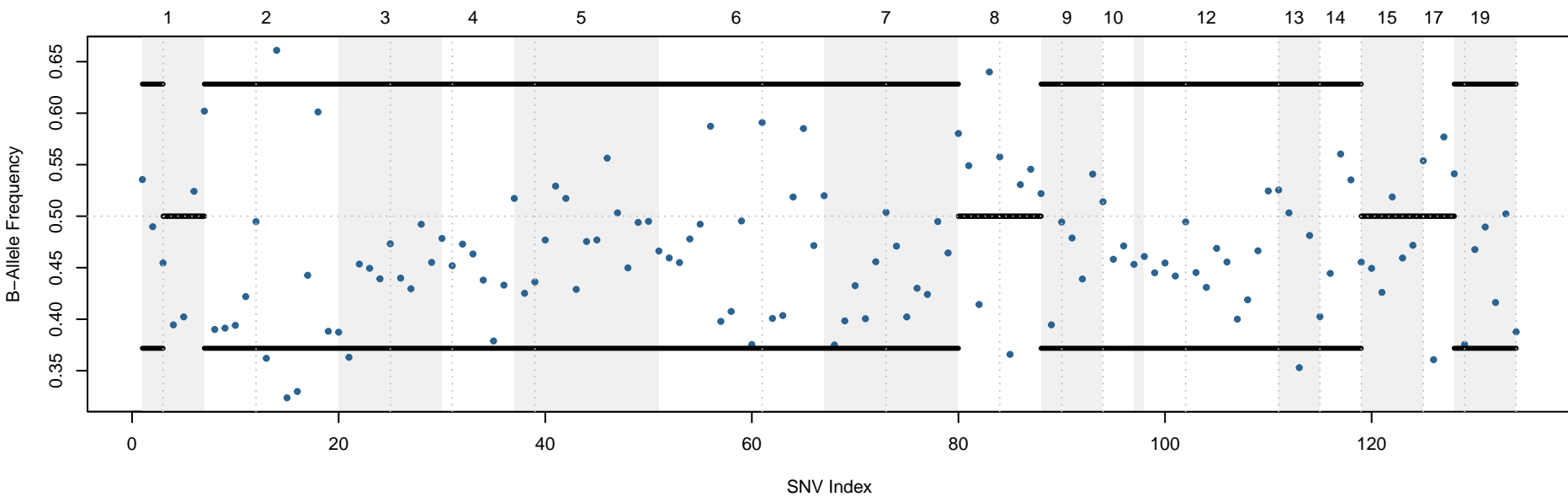




Purity: 0.69 Tumor ploidy: 3.097



Purity: 0.69 Tumor ploidy: 3.097 SNV log-likelihood: -709.59 GoF: 45.7% Mean coverage: 385,403



SCNA-fit log-likelihood: -3697.92

