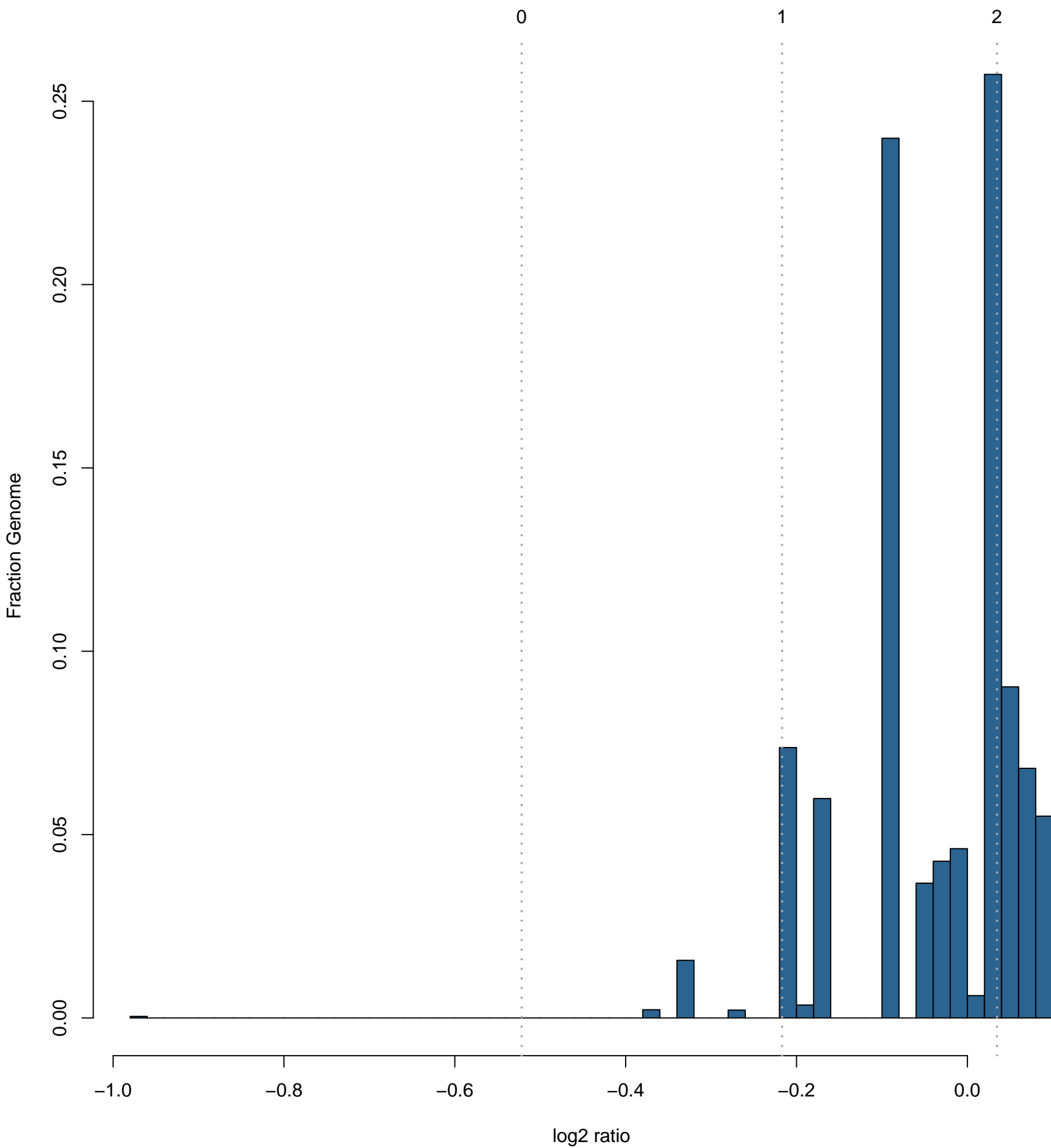
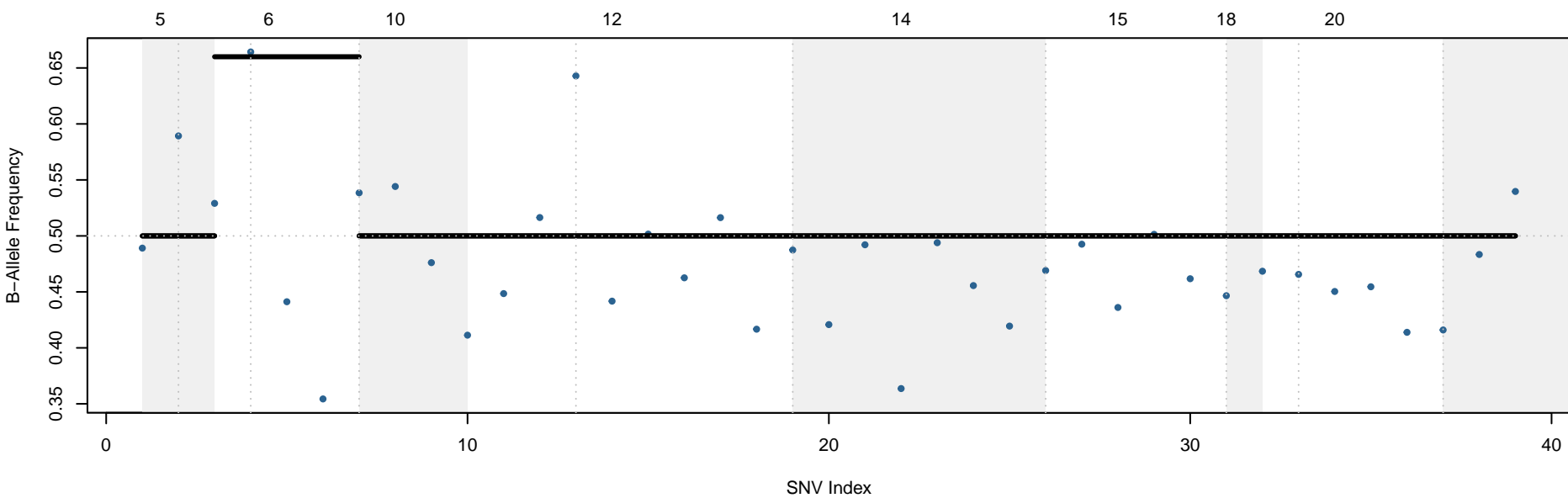


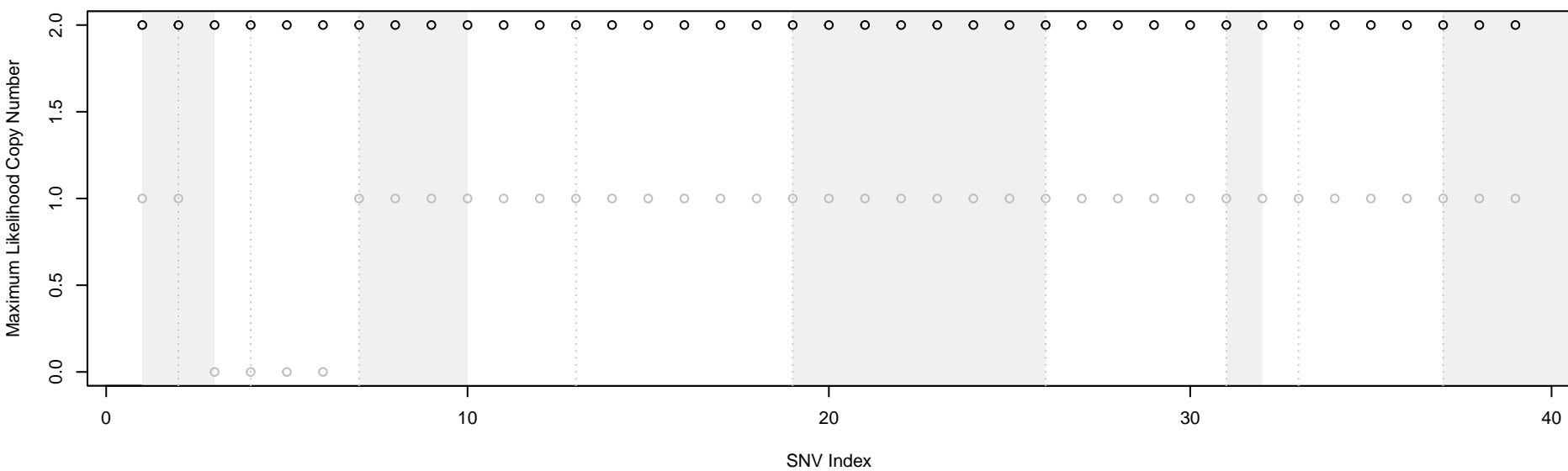
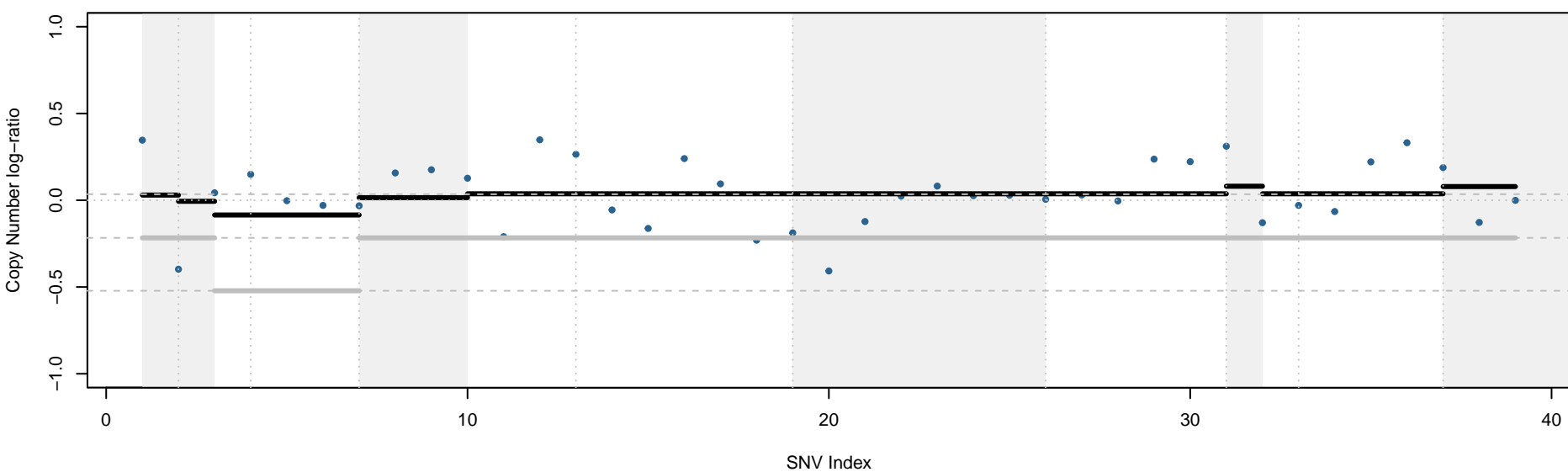
Purity: 0.32 Tumor ploidy: 1.852

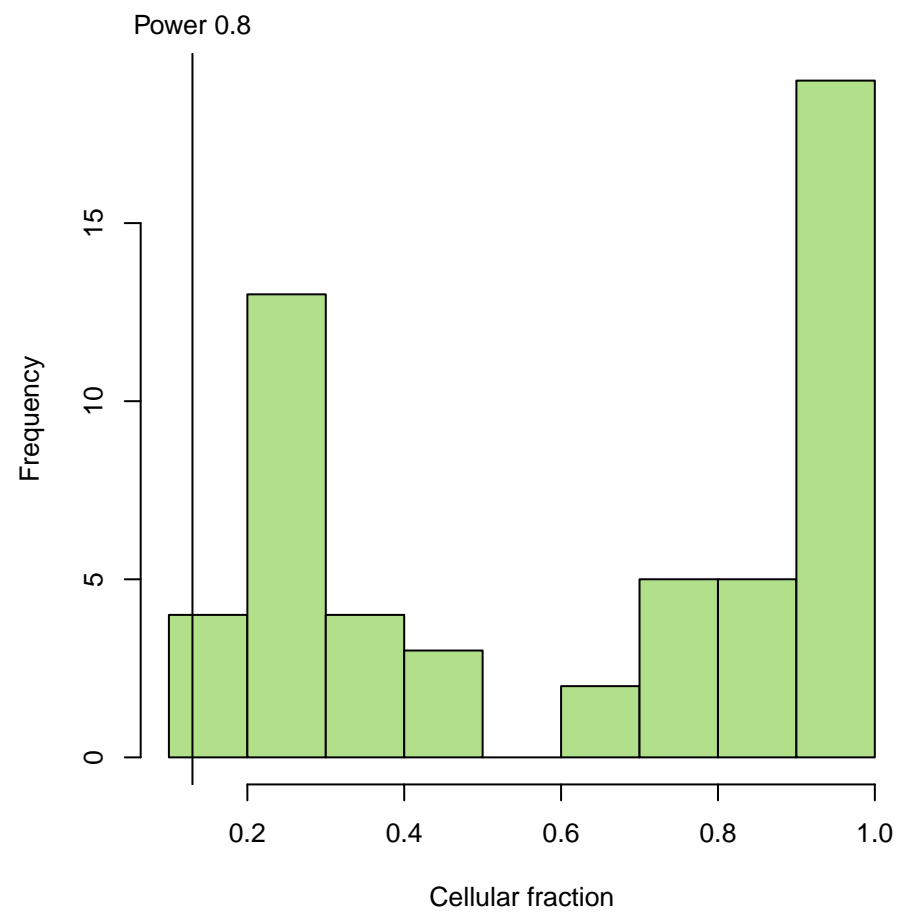
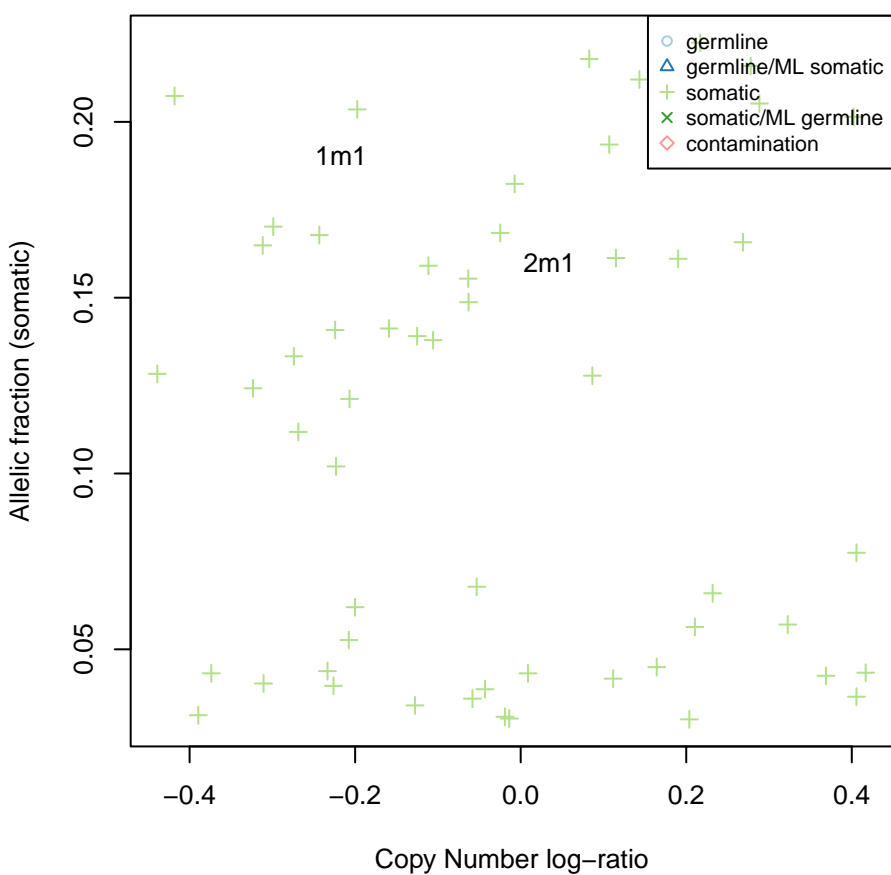
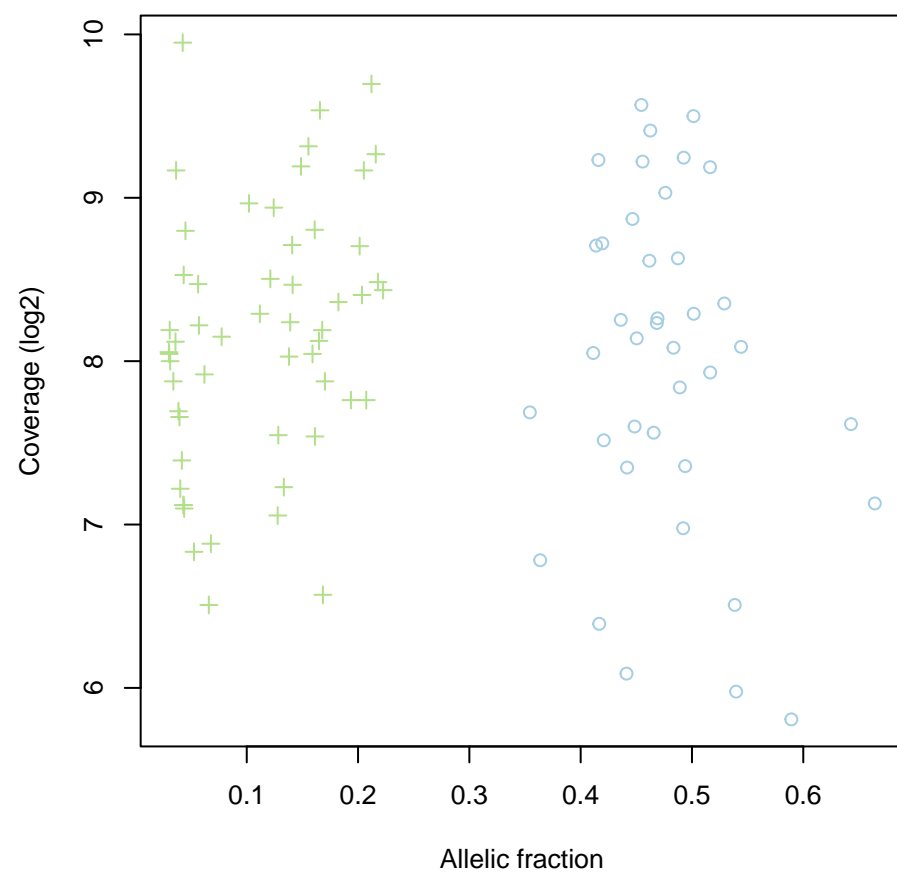
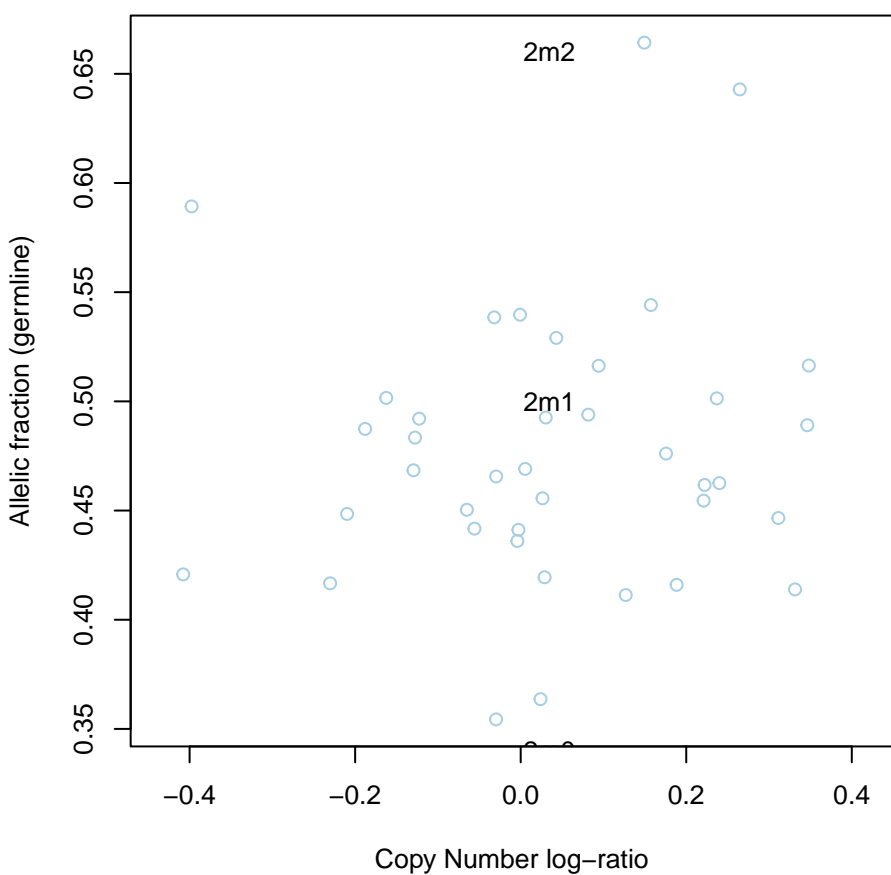


Purity: 0.32 Tumor ploidy: 1.852 SNV log-likelihood: 6.76 GoF: 87.3% Mean coverage: 315;324

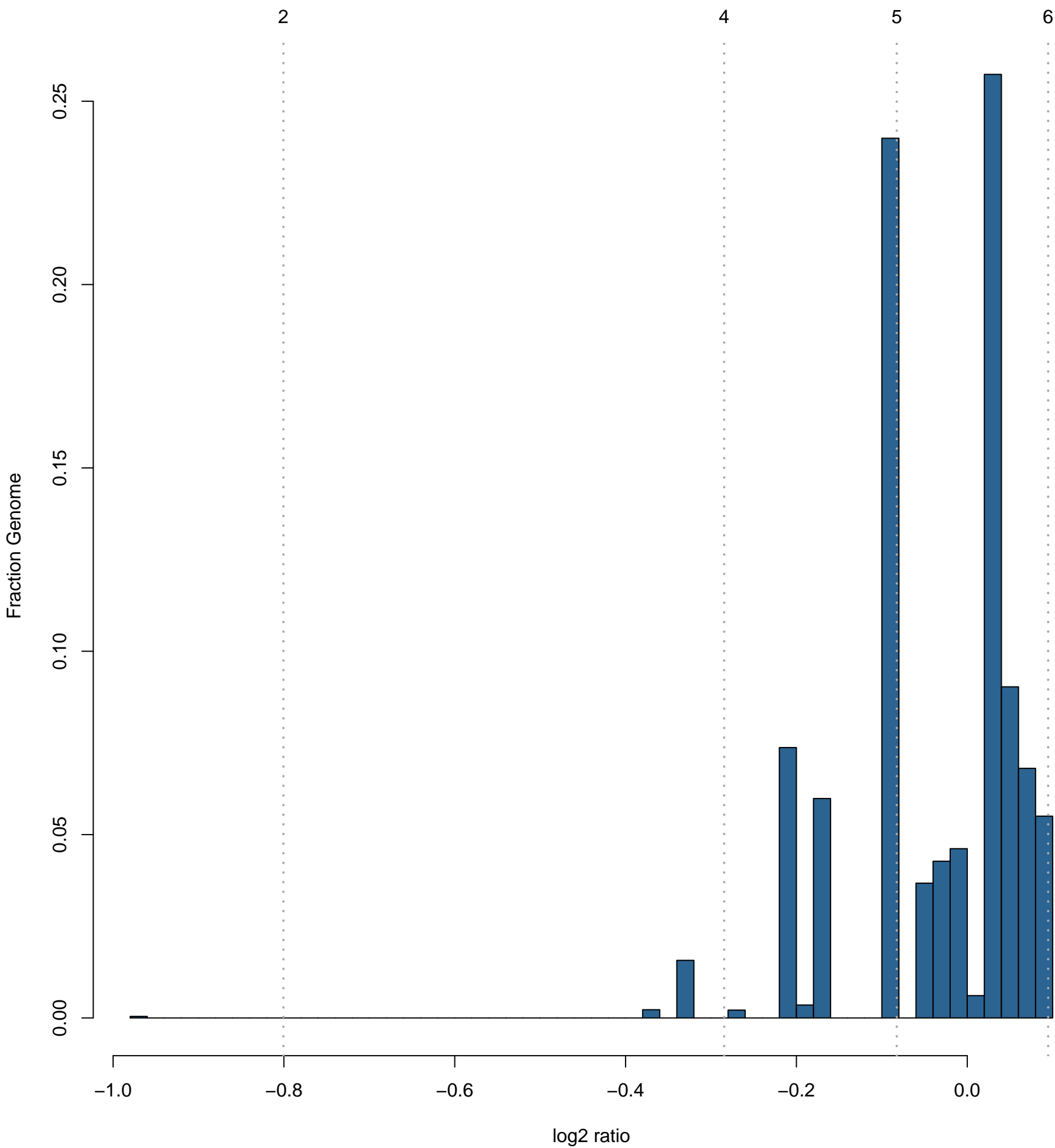


SCNA-fit log-likelihood: -624.43

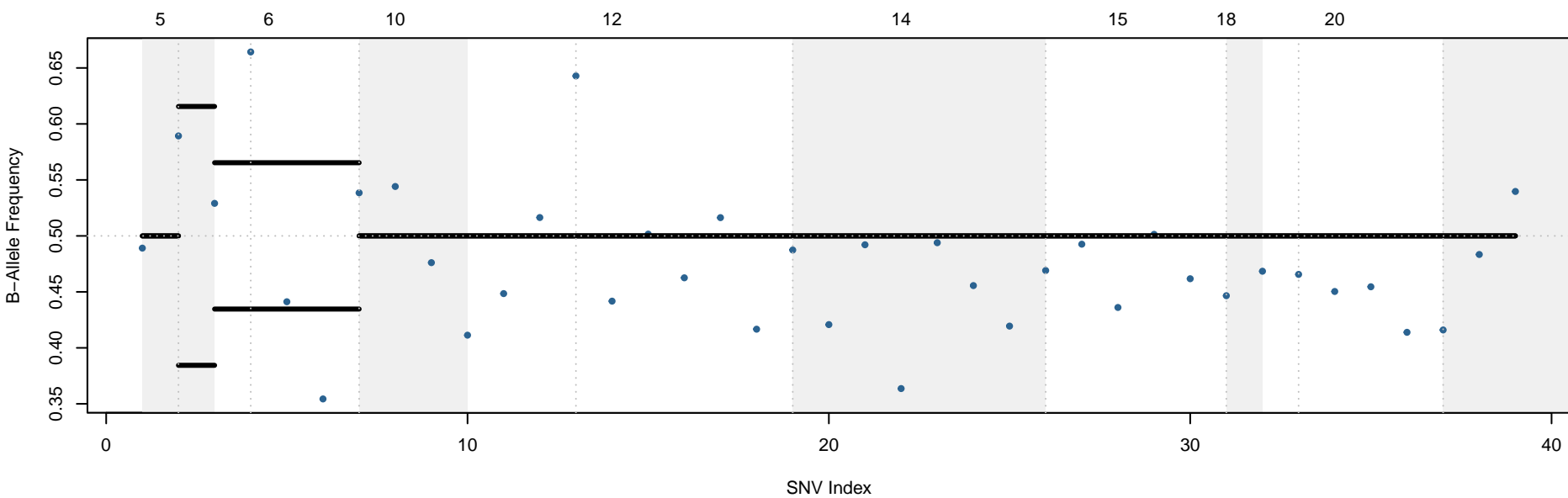




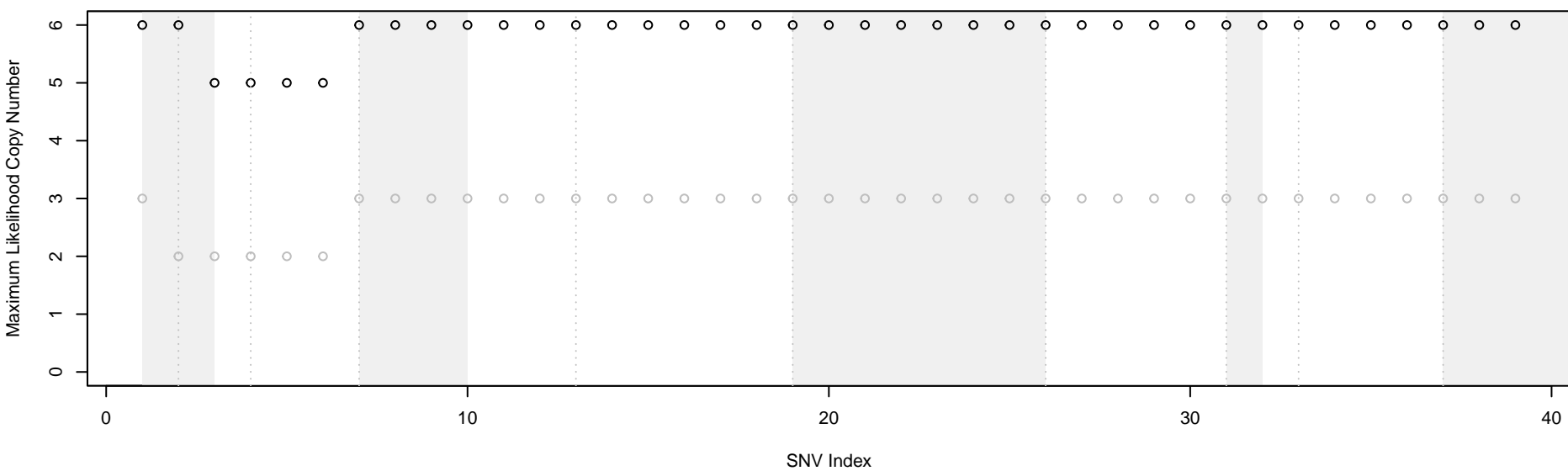
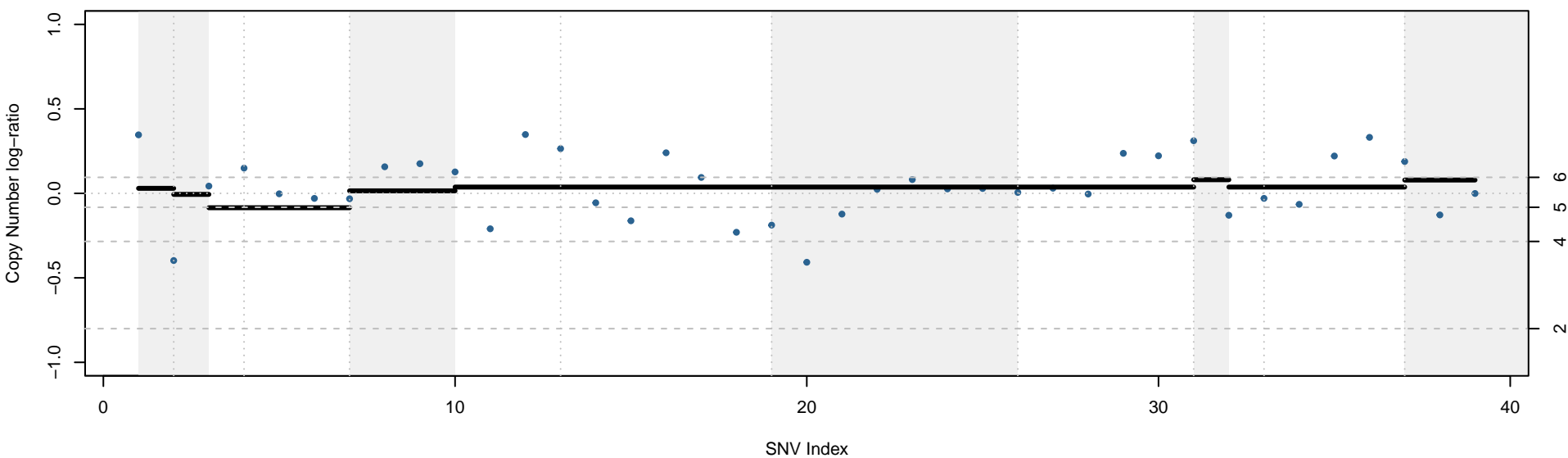
Purity: 0.43 Tumor ploidy: 5.45

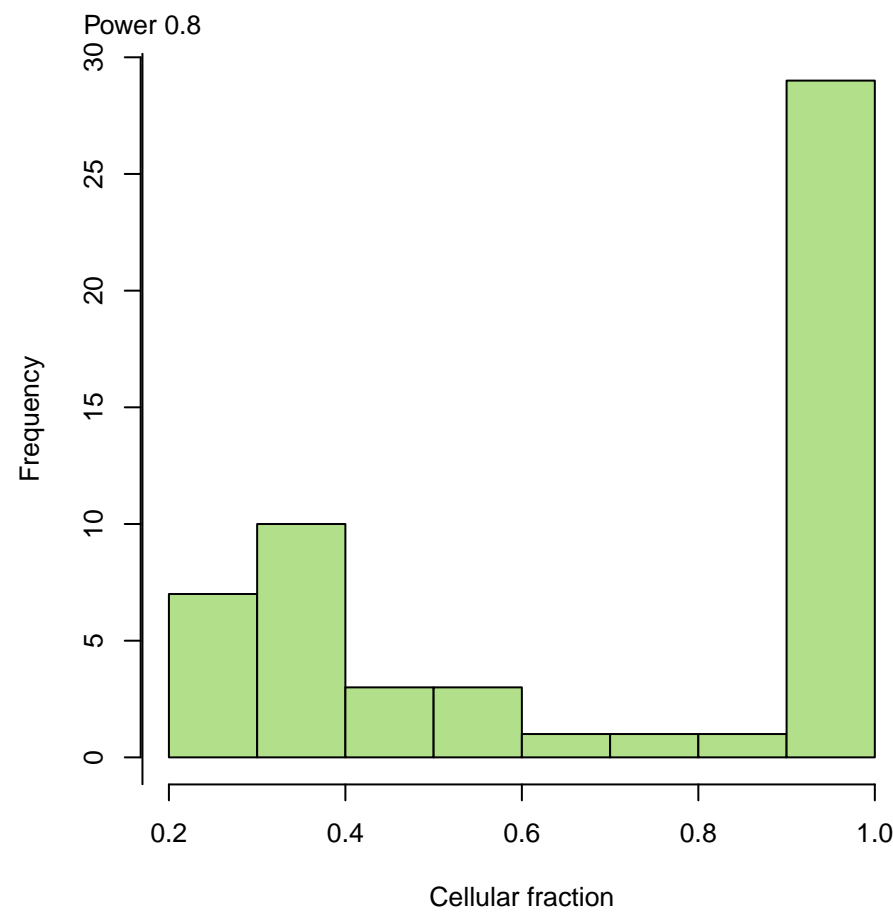
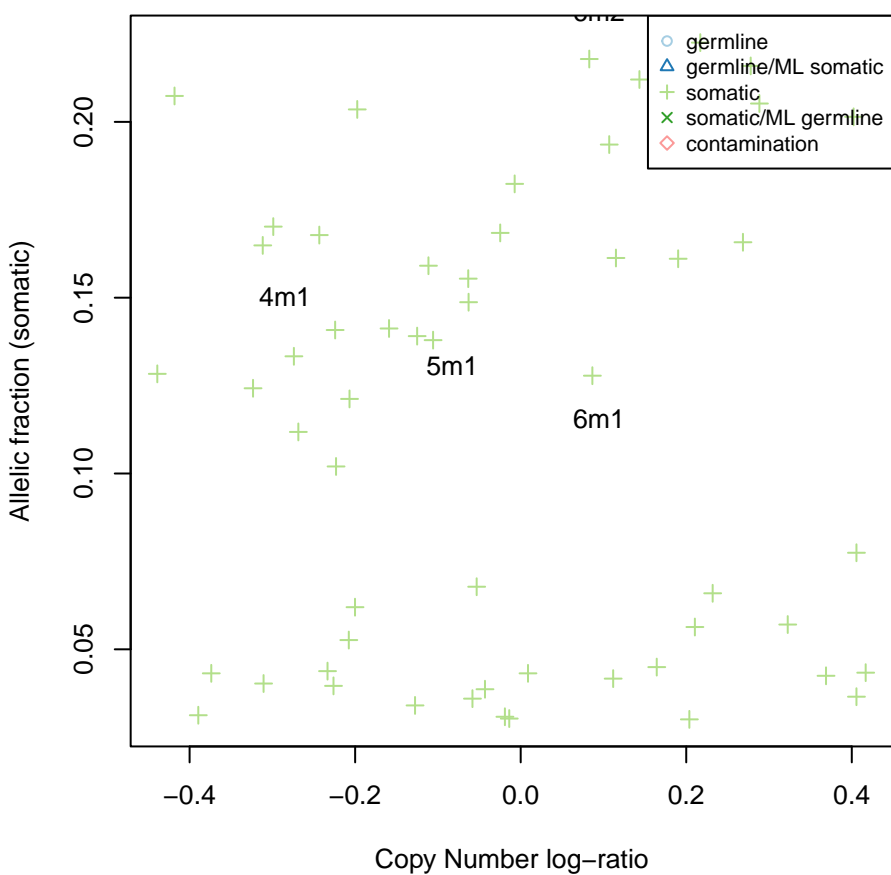
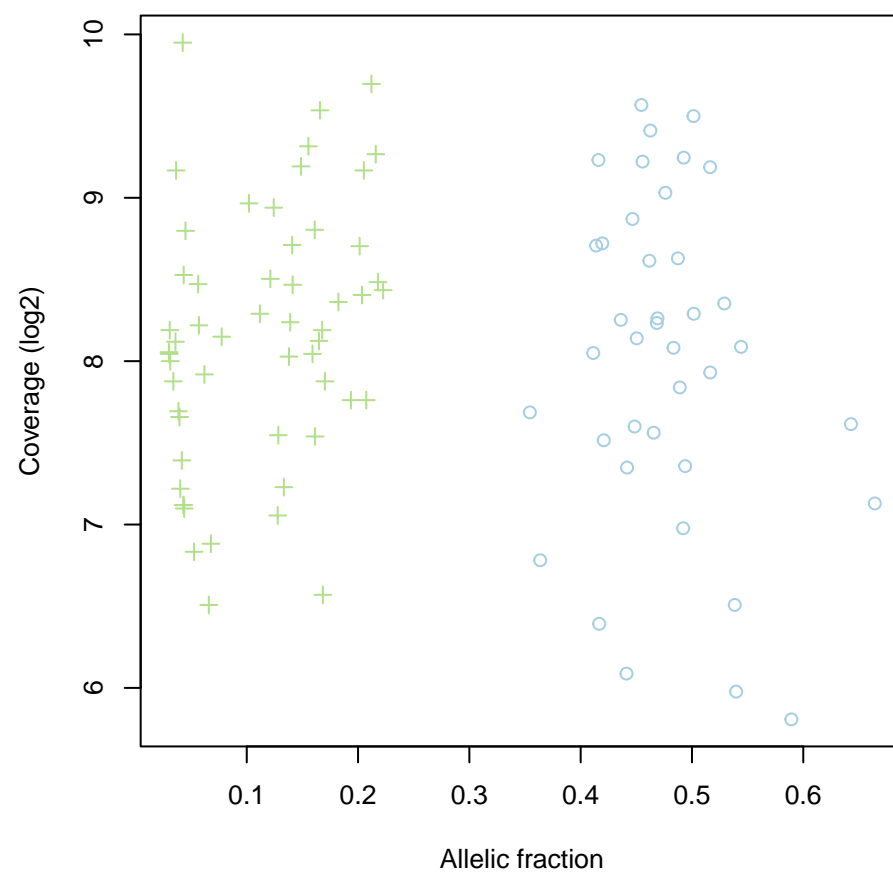
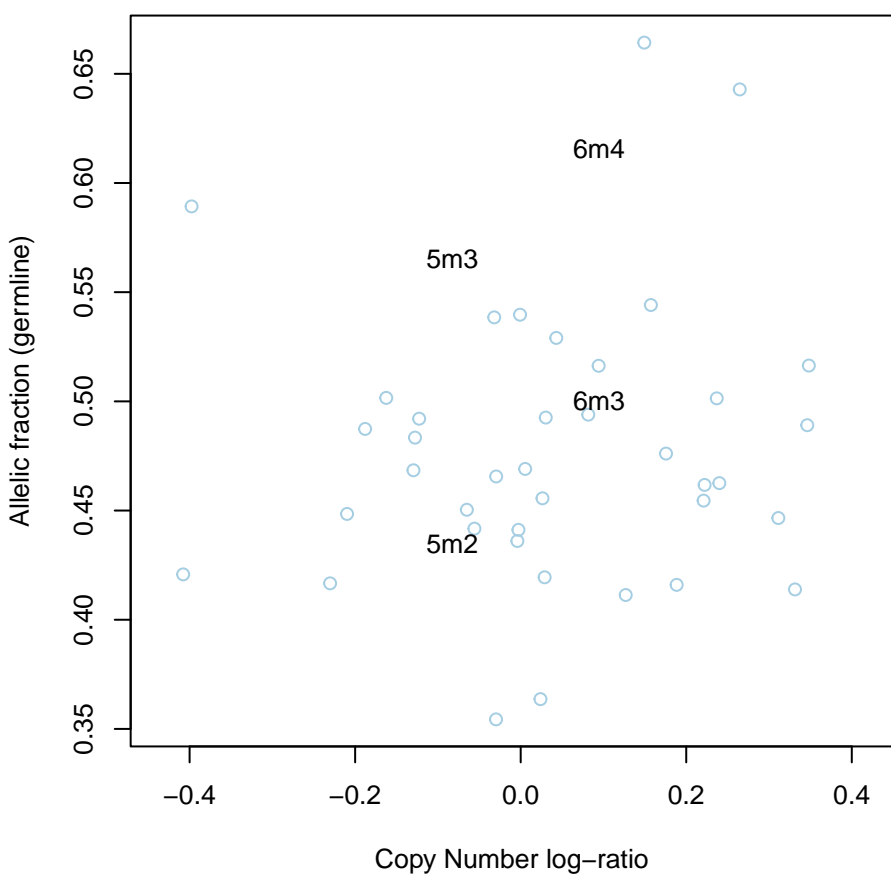


Purity: 0.43 Tumor ploidy: 5.45 SNV log-likelihood: -74.08 GoF: 92.8% Mean coverage: 315;324

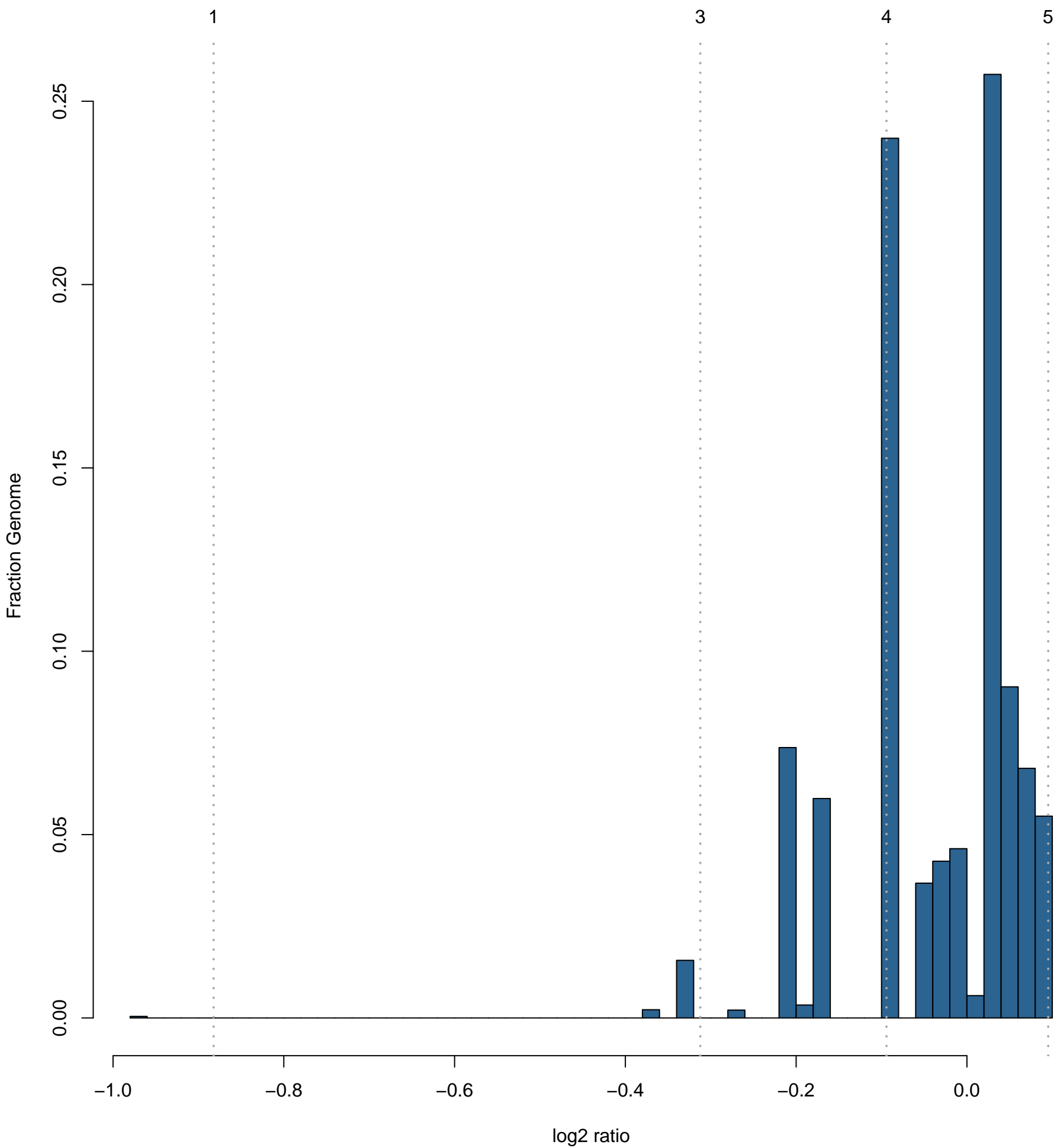


SCNA-fit log-likelihood: -448.81

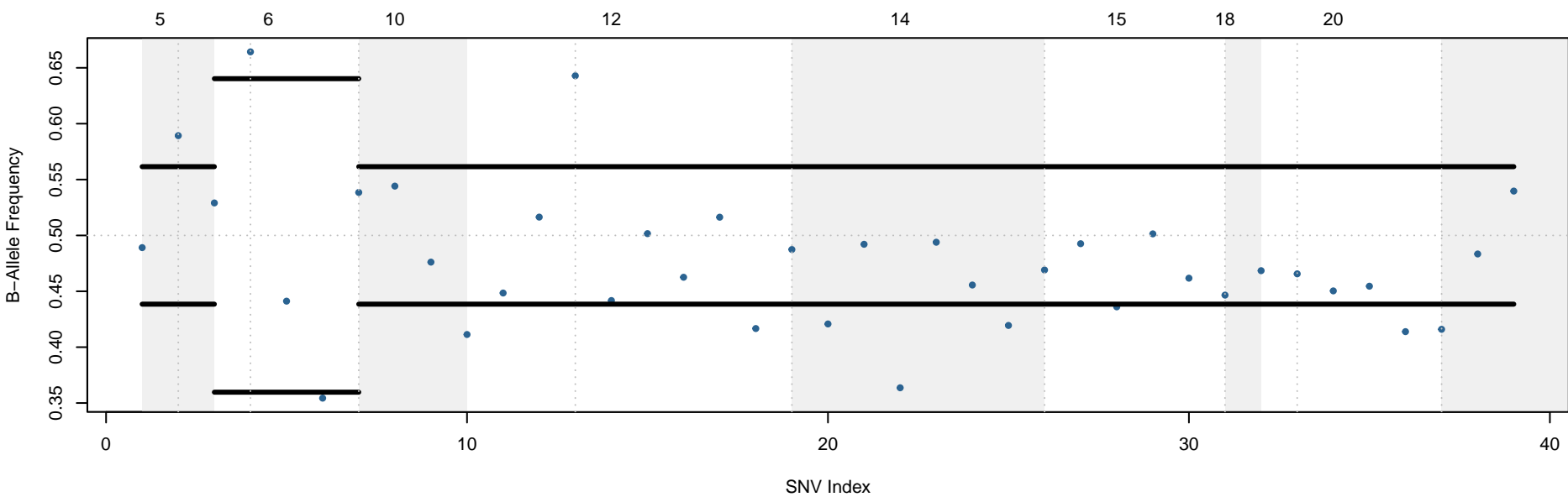




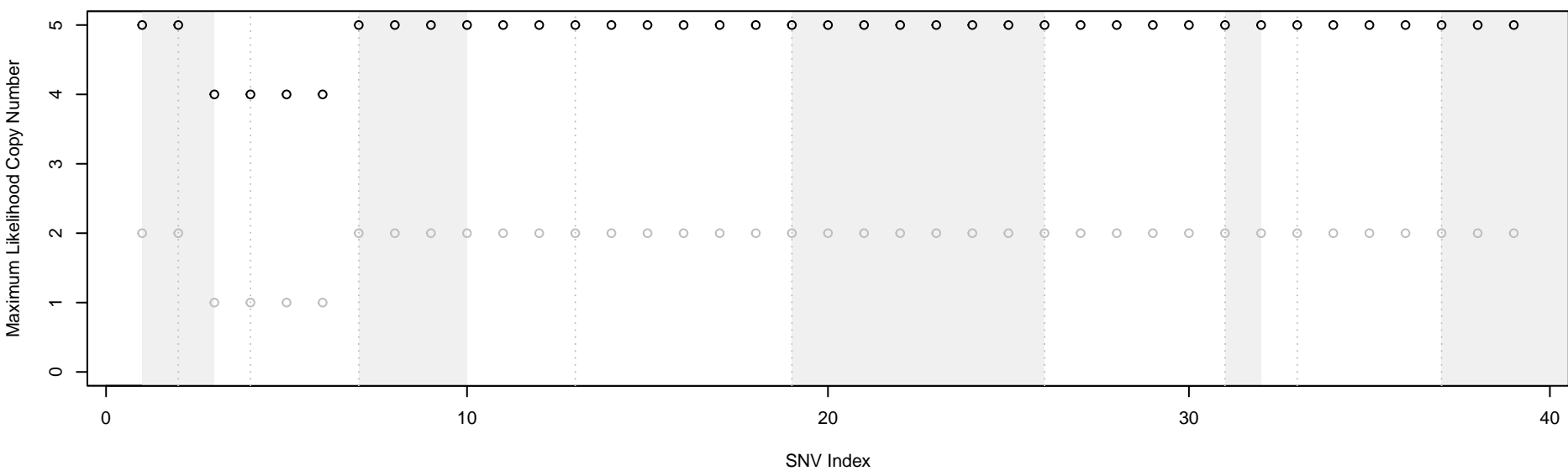
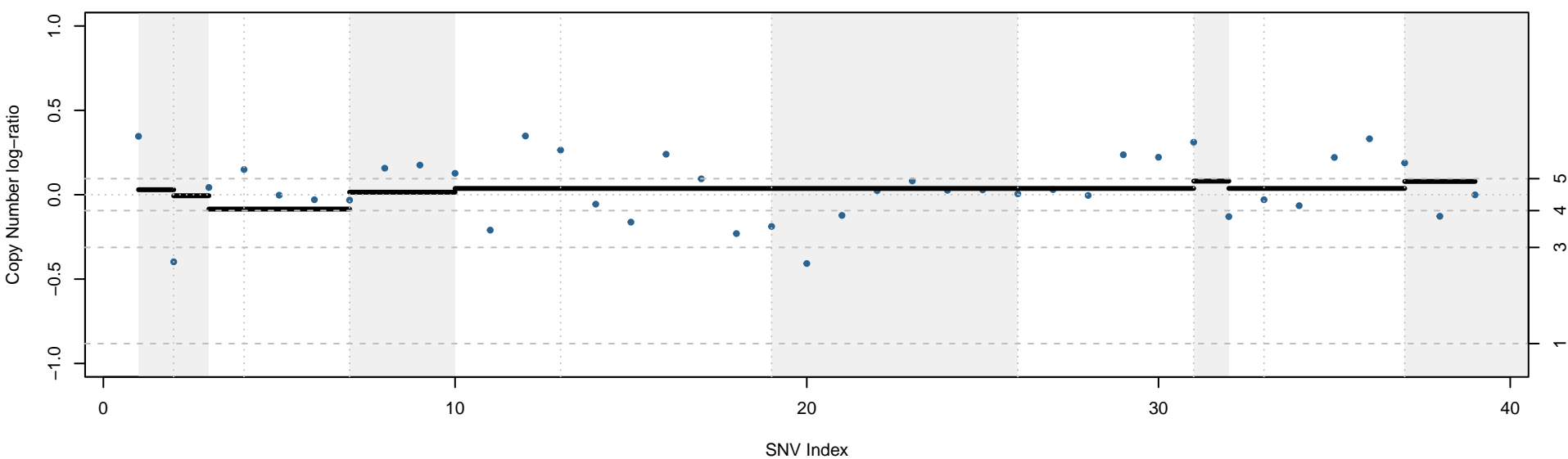
Purity: 0.39 Tumor ploidy: 4.481



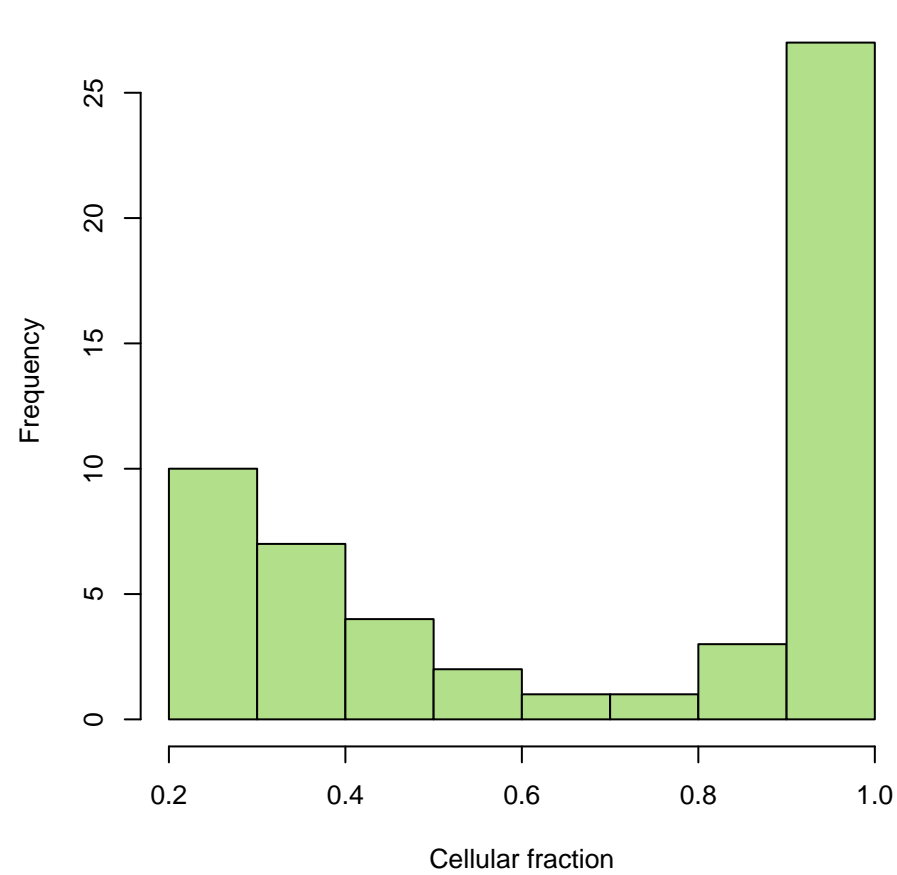
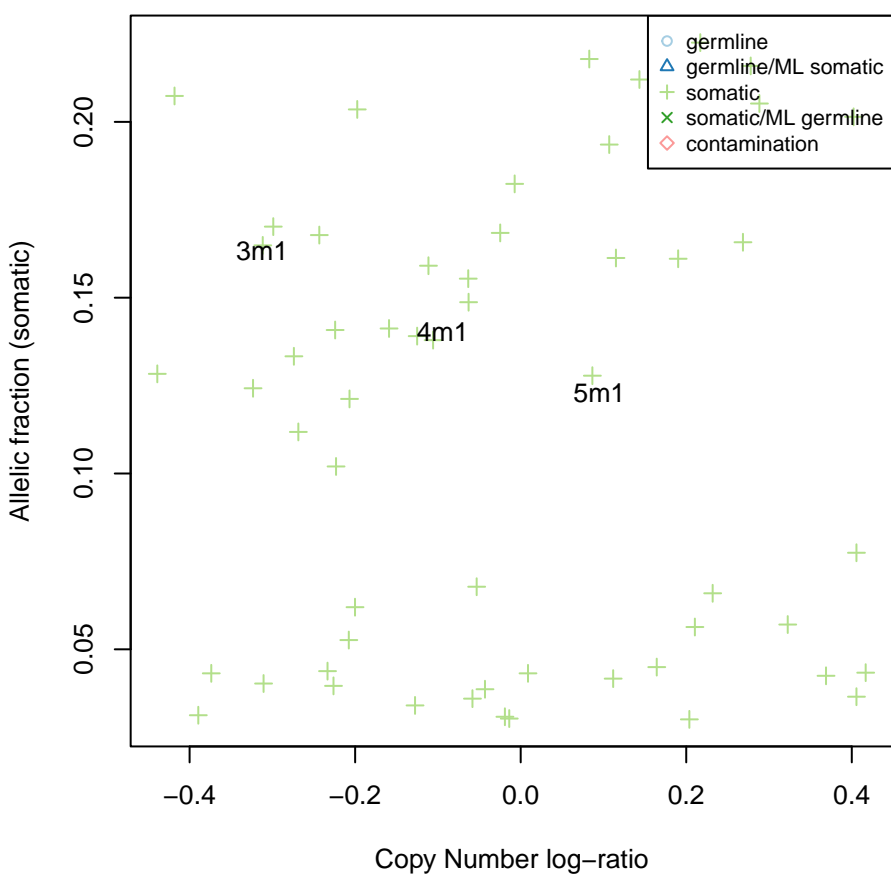
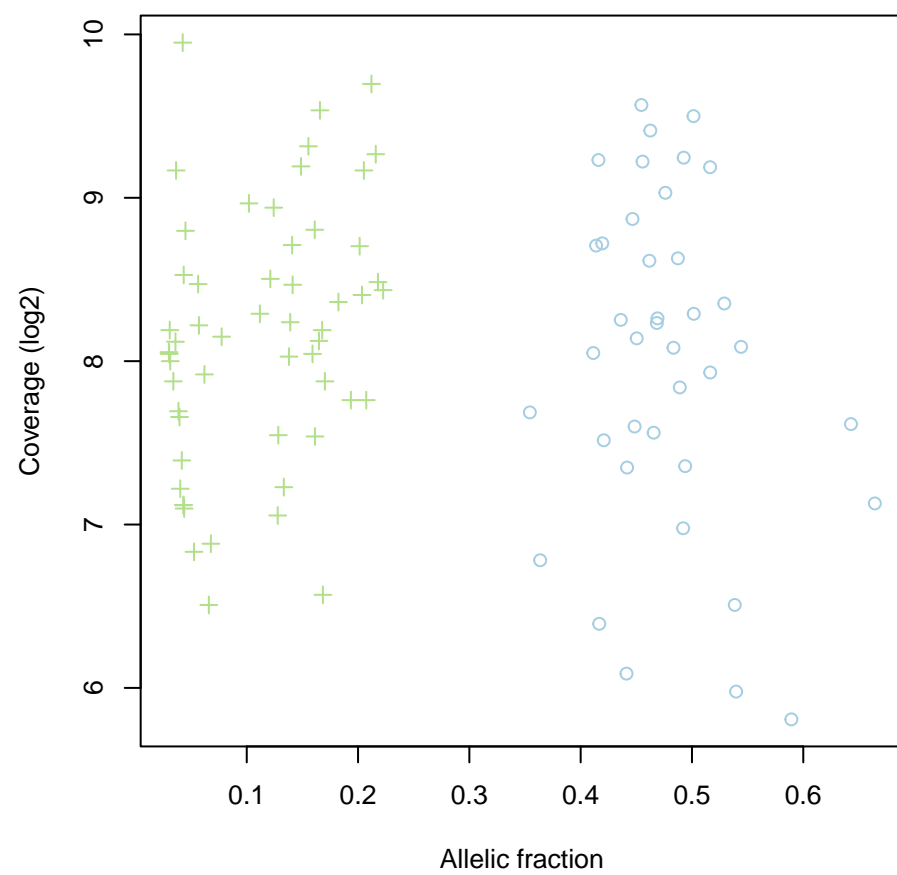
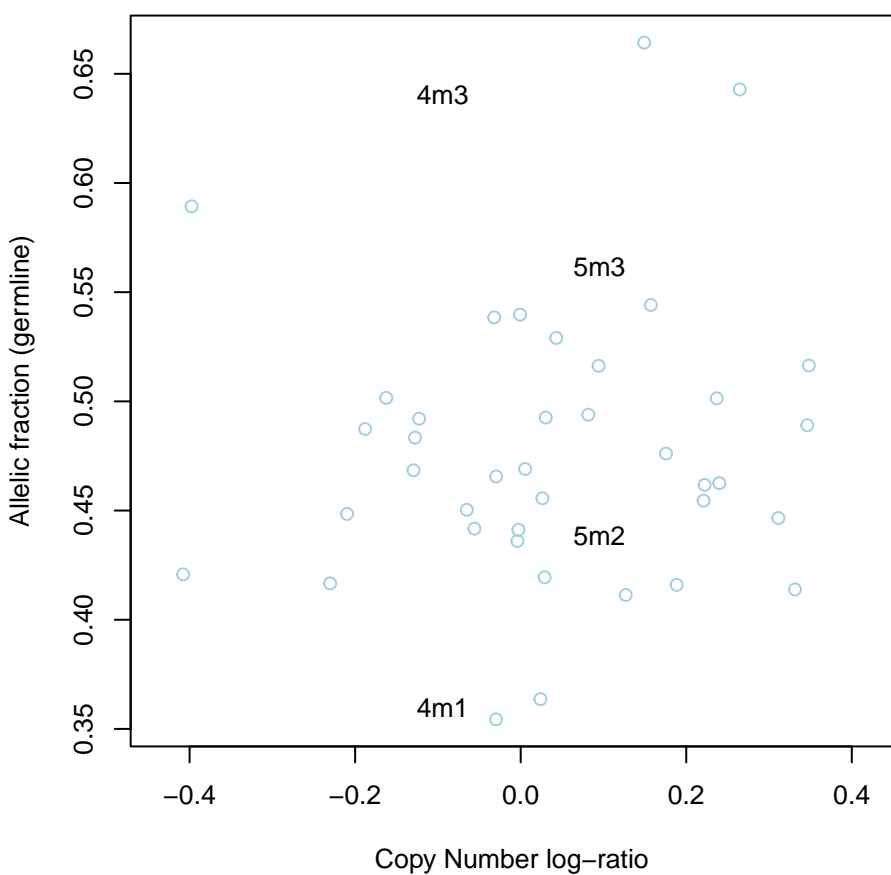
Purity: 0.39 Tumor ploidy: 4.481 SNV log-likelihood: -72.24 GoF: 92.4% Mean coverage: 315;324



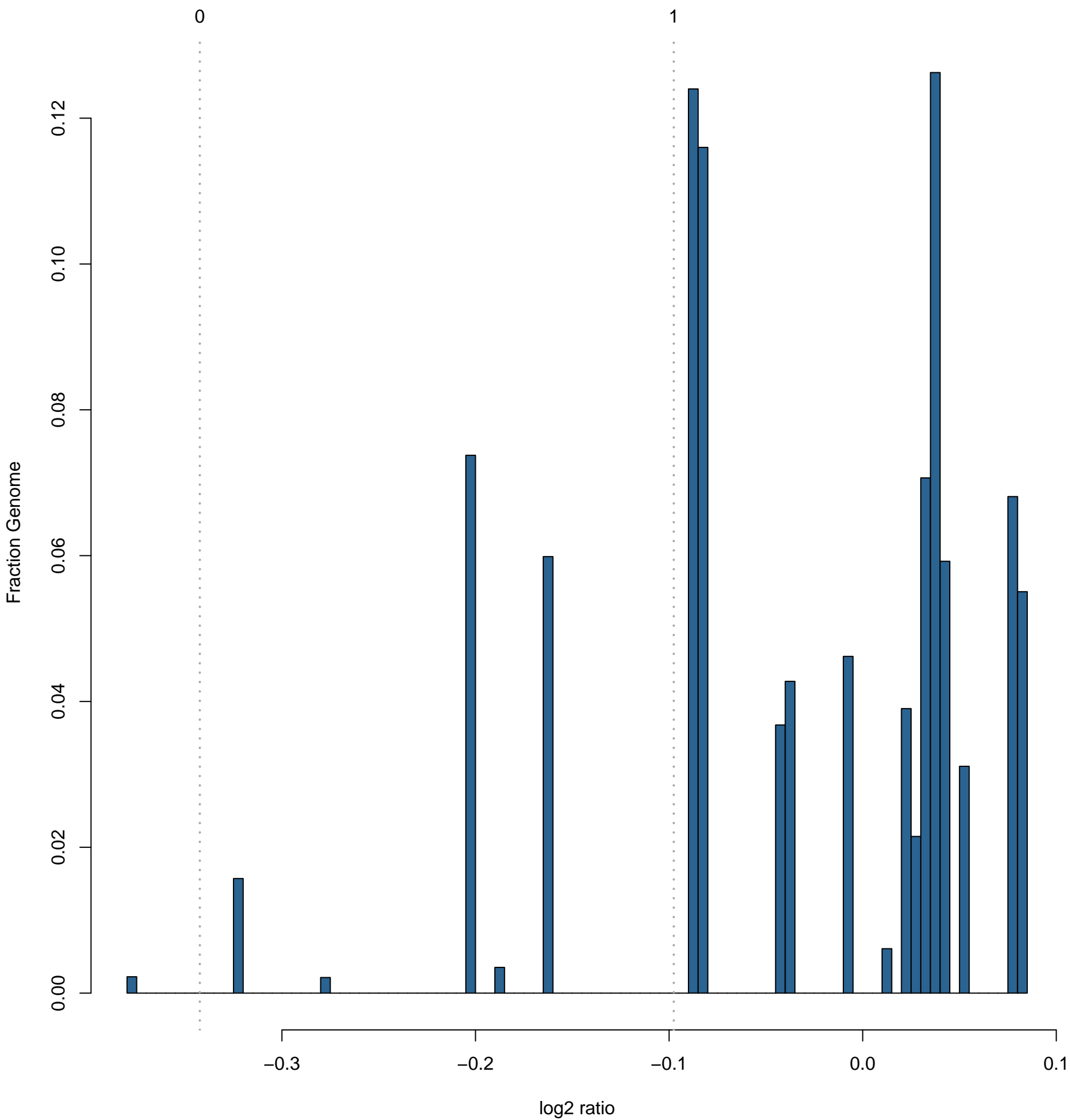
SCNA-fit log-likelihood: -458.18



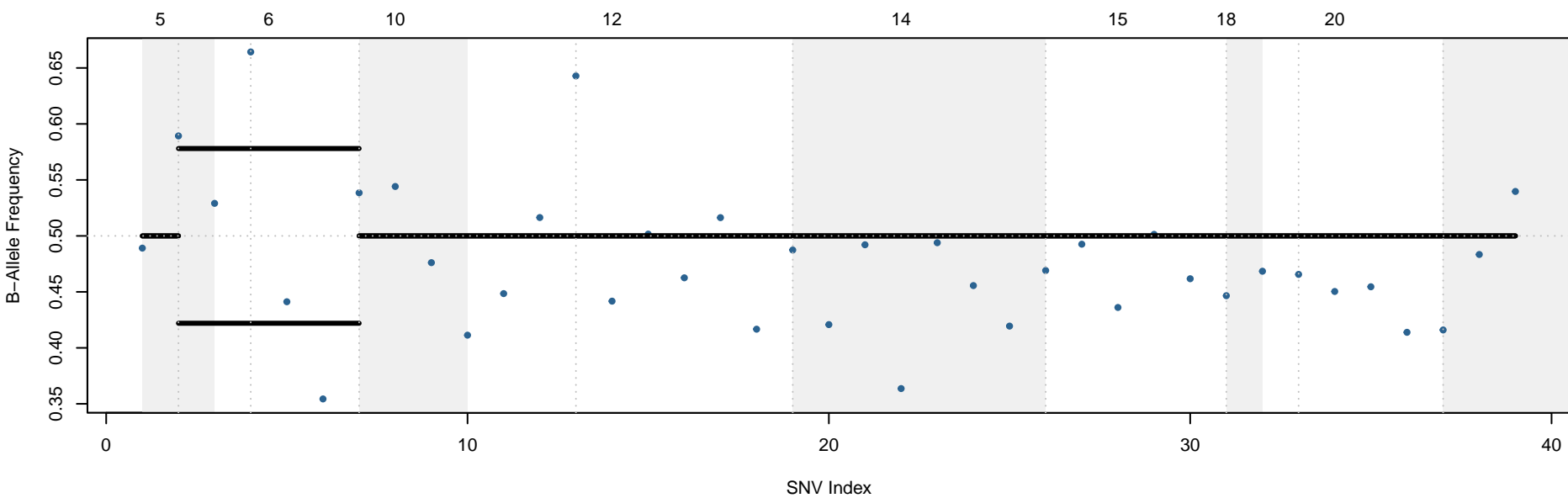




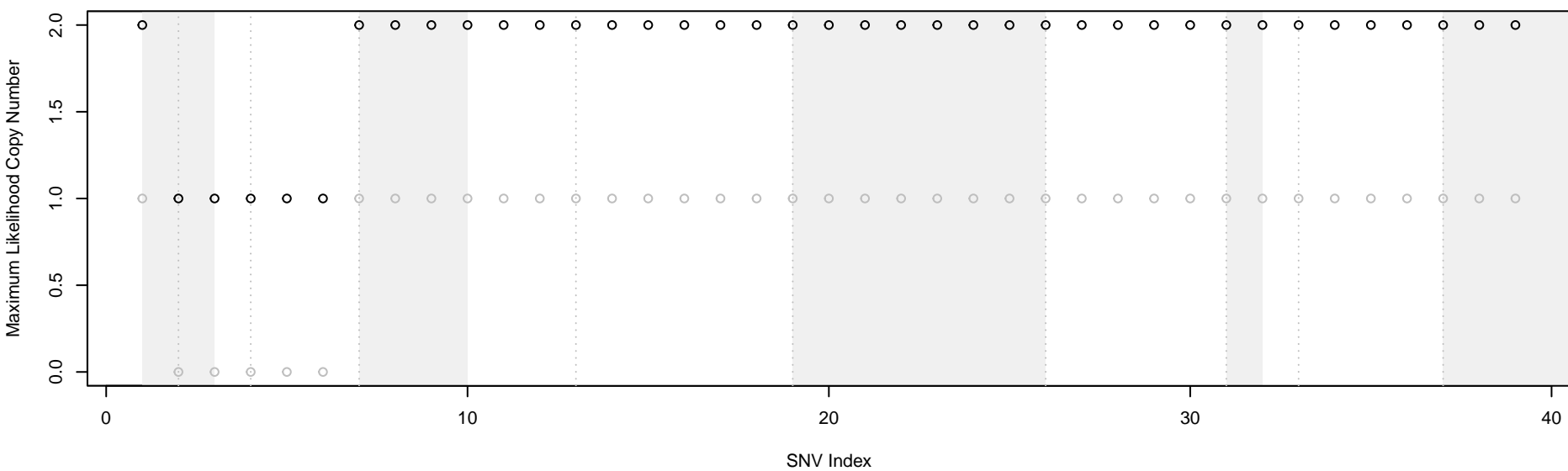
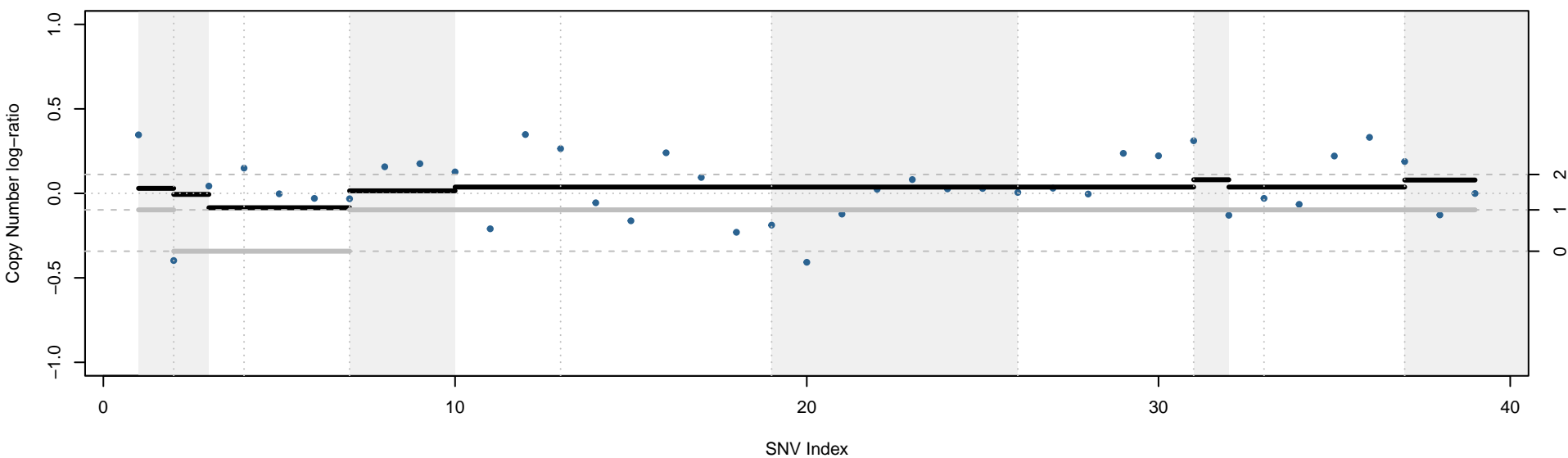
Purity: 0.27 Tumor ploidy: 1.448

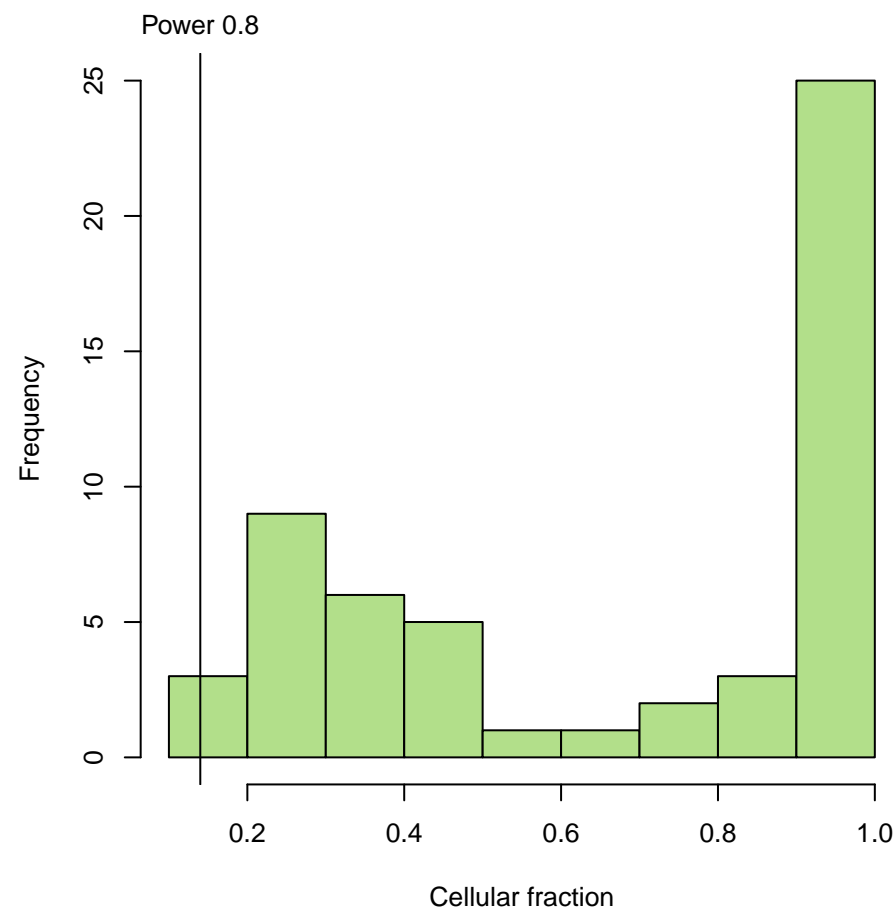
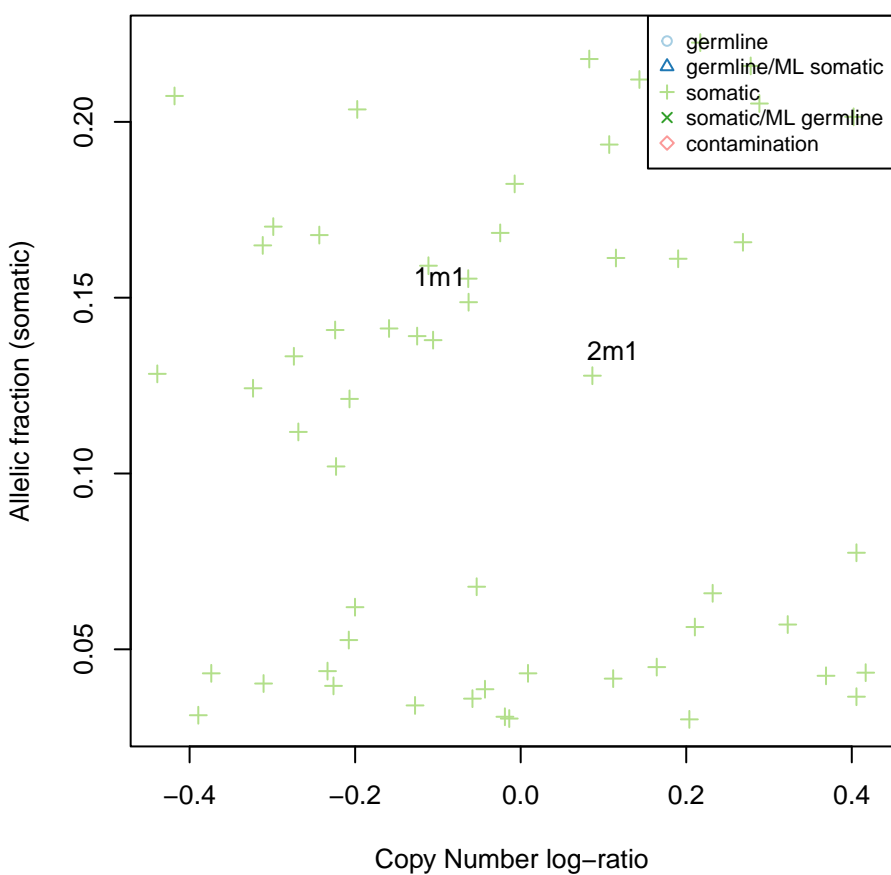
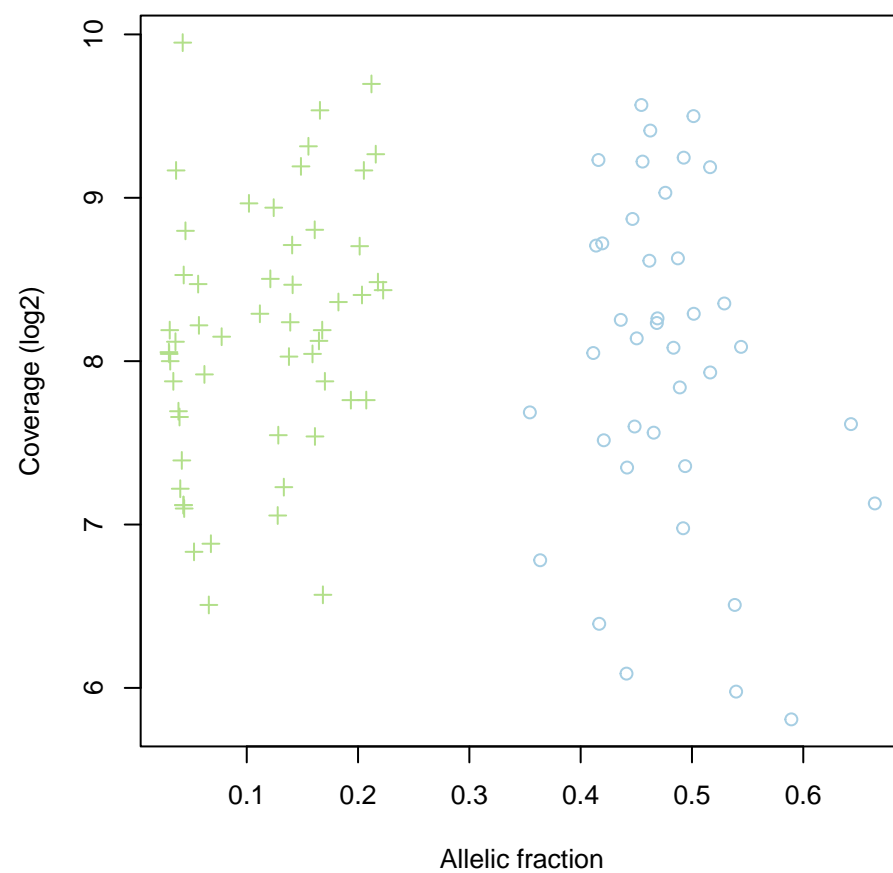
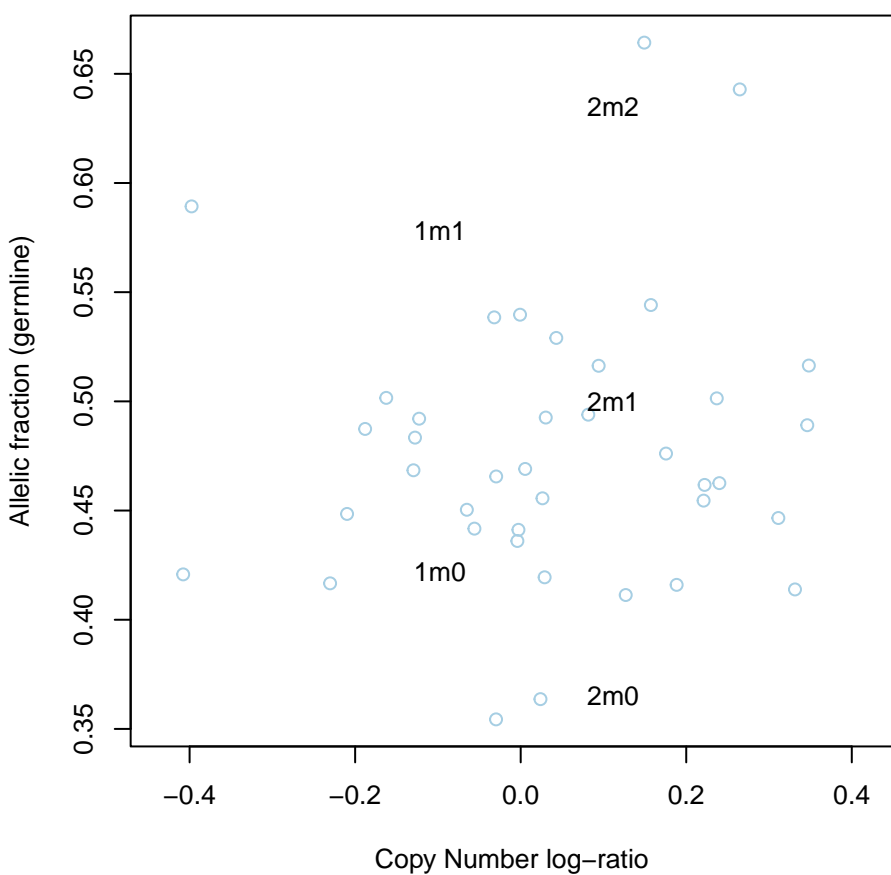


Purity: 0.27 Tumor ploidy: 1.448 SNV log-likelihood: -27.94 GoF: 90.7% Mean coverage: 315;324

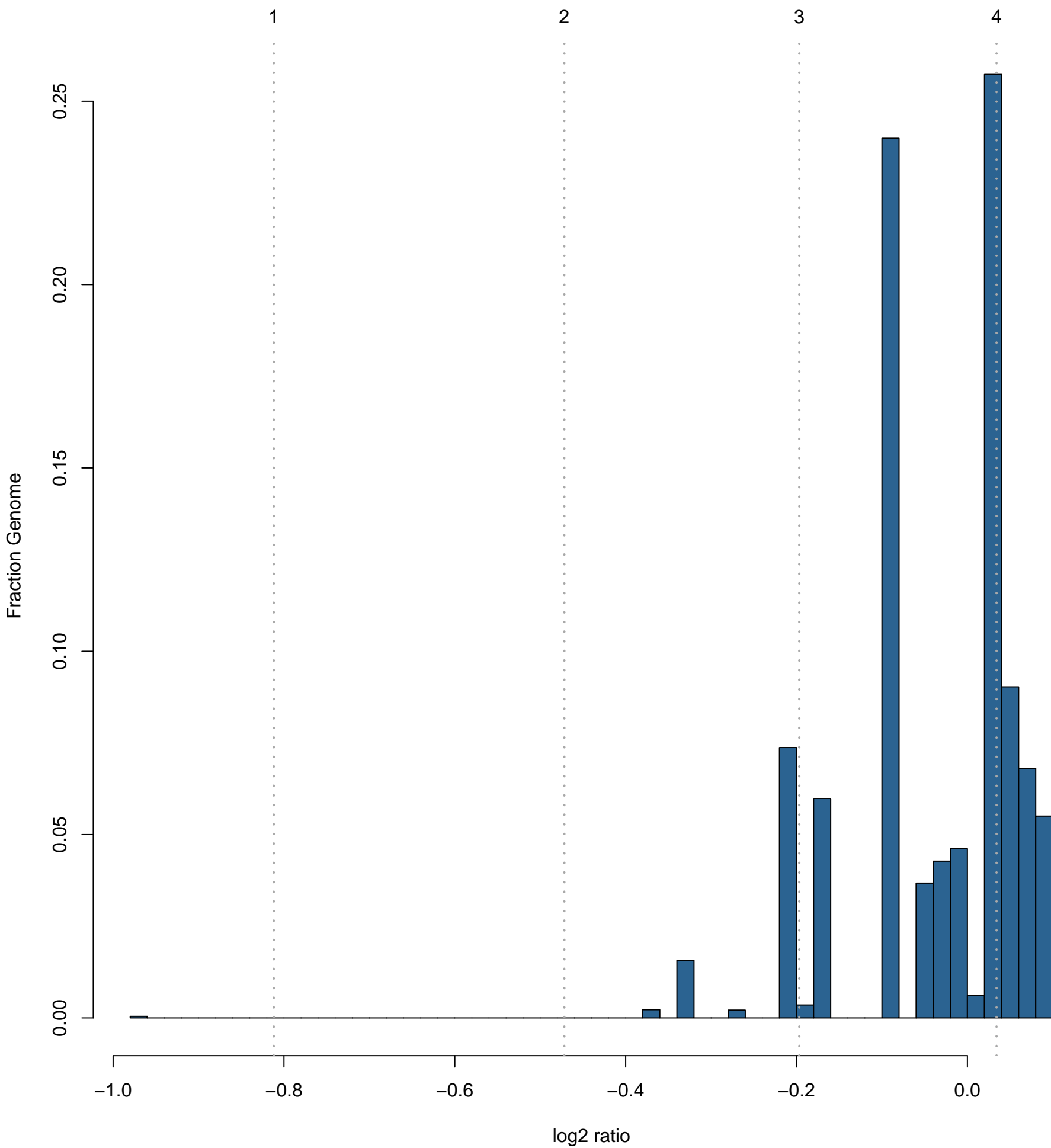


SCNA-fit log-likelihood: -642.33

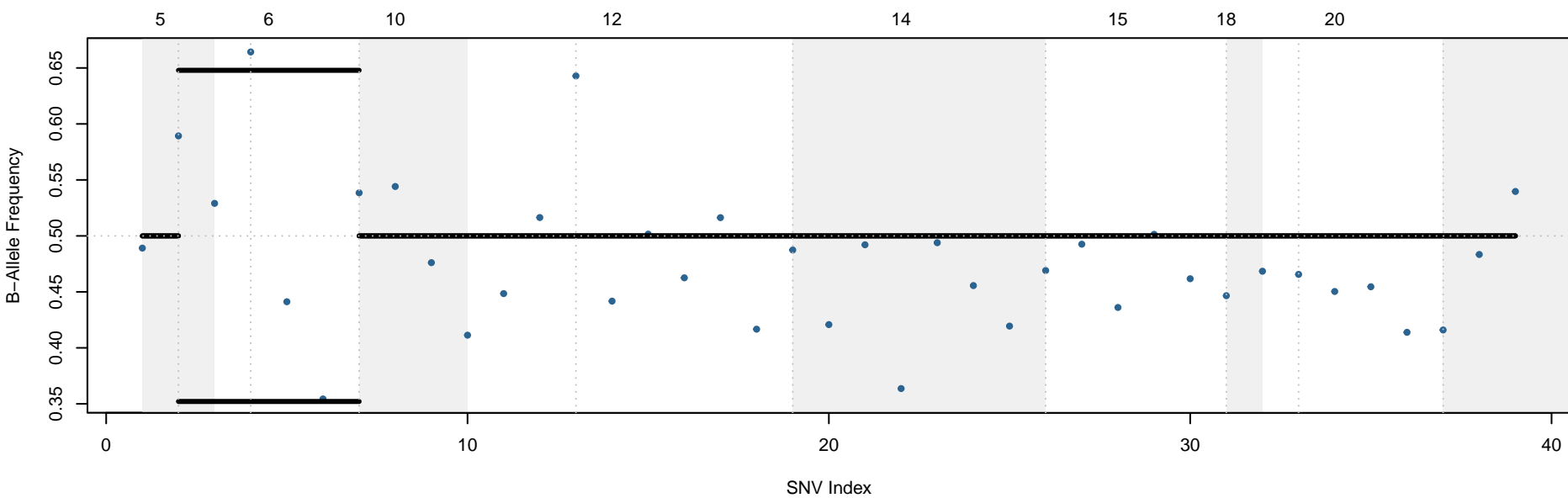




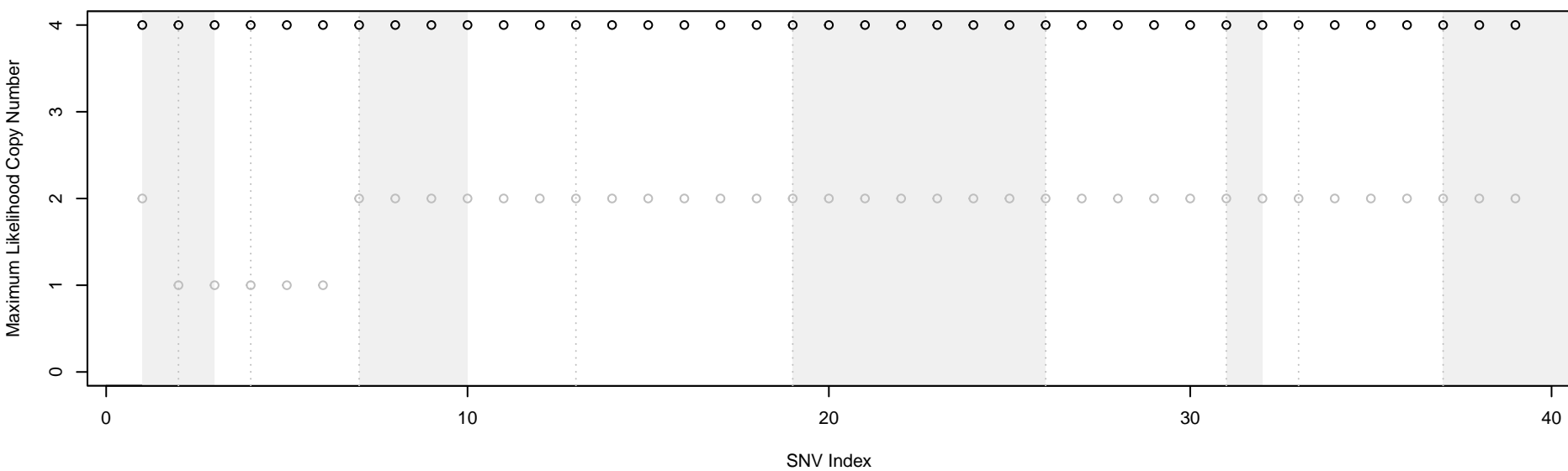
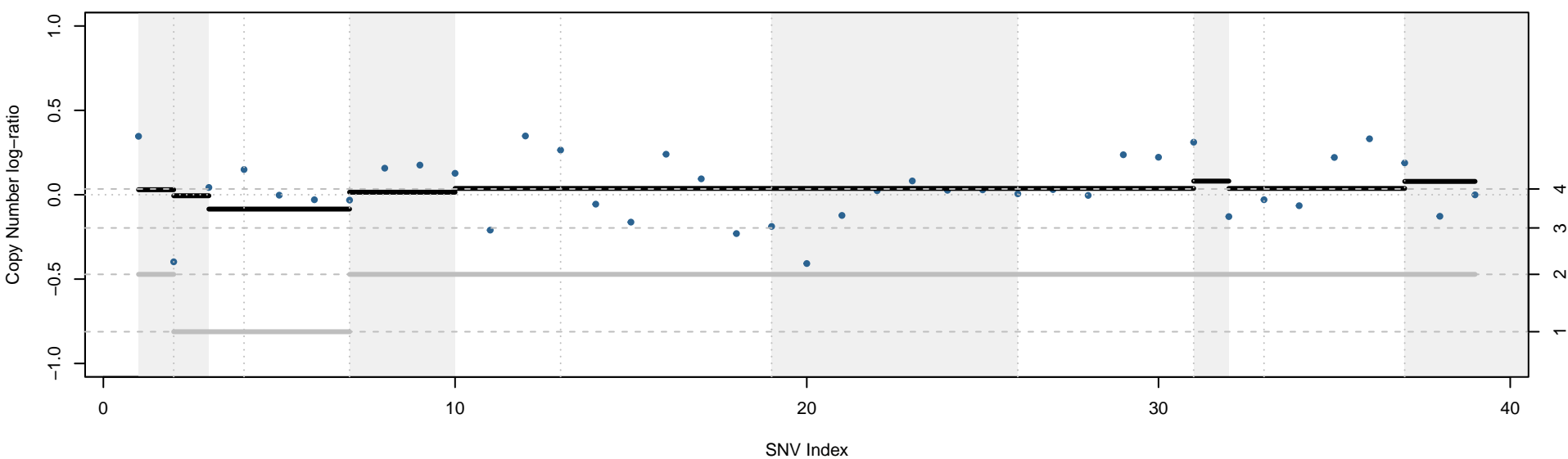
Purity: 0.42 Tumor ploidy: 3.842

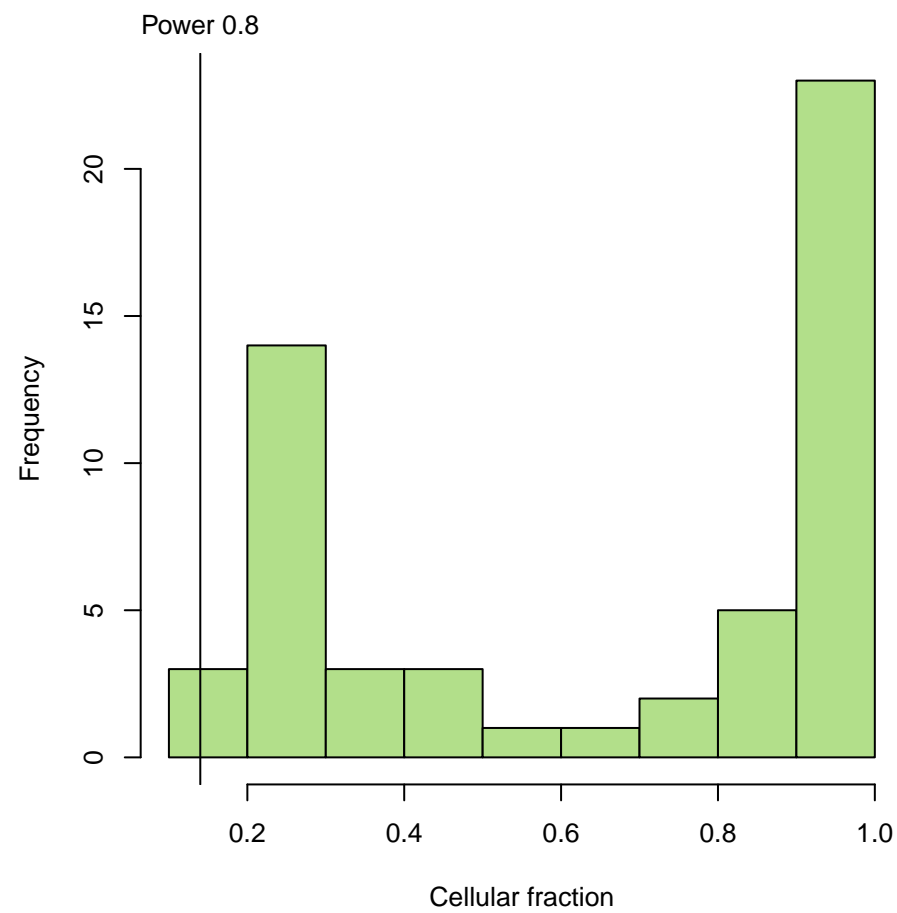
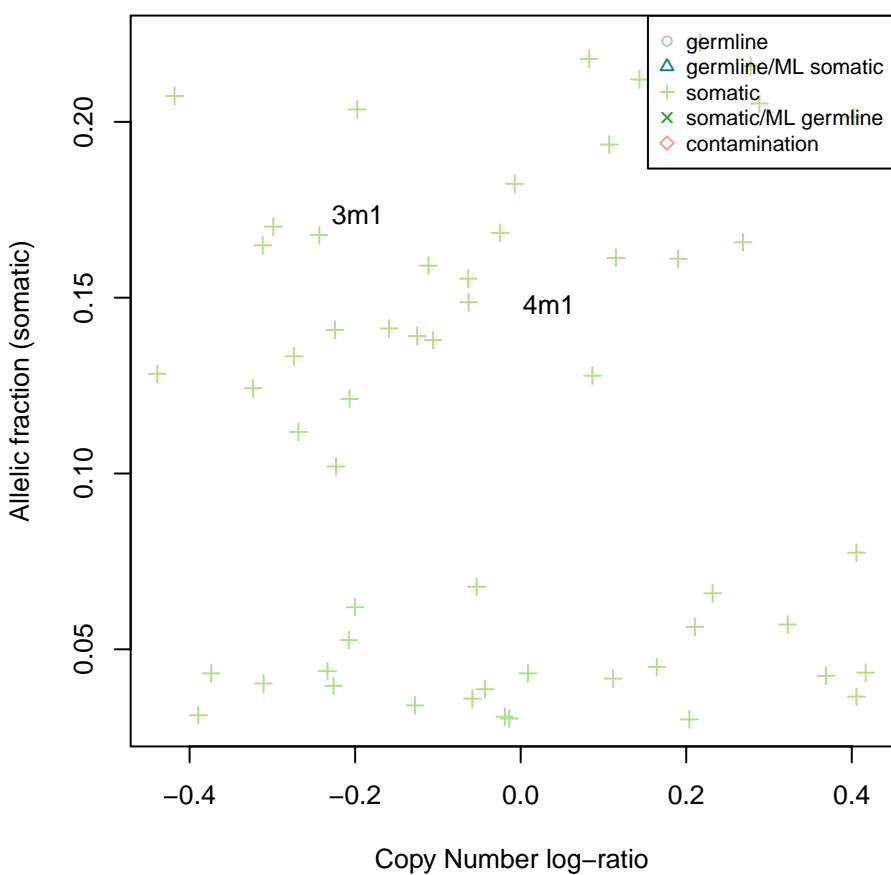
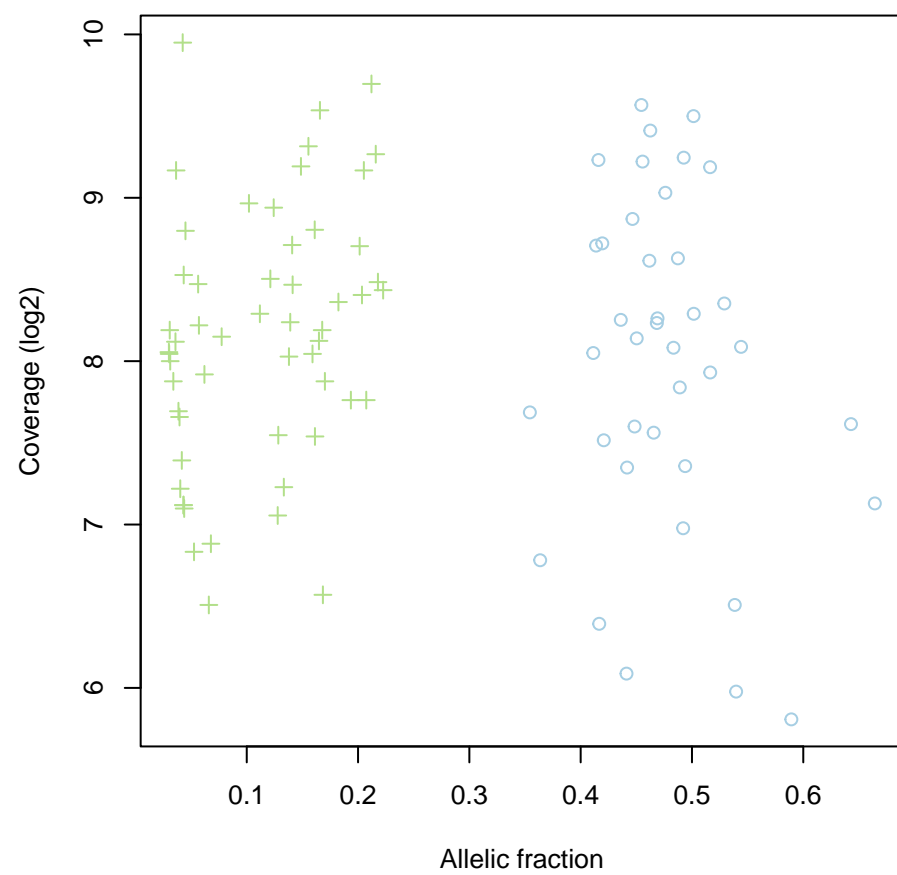
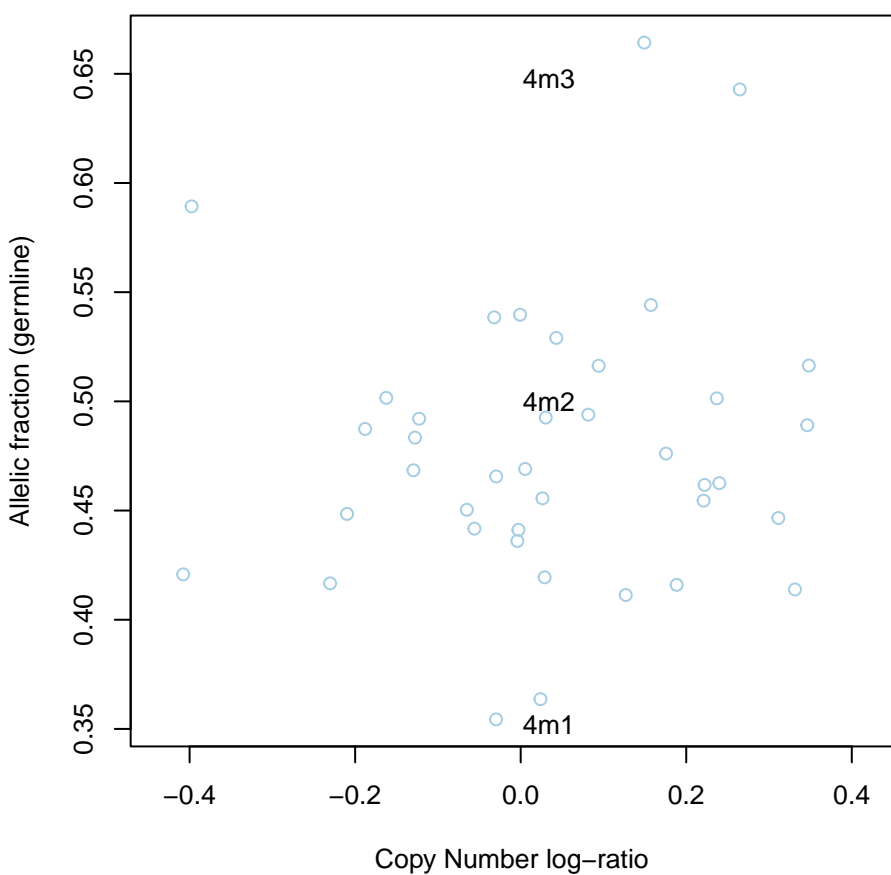


Purity: 0.42 Tumor ploidy: 3.842 SNV log-likelihood: -62.98 GoF: 88.8% Mean coverage: 315;324

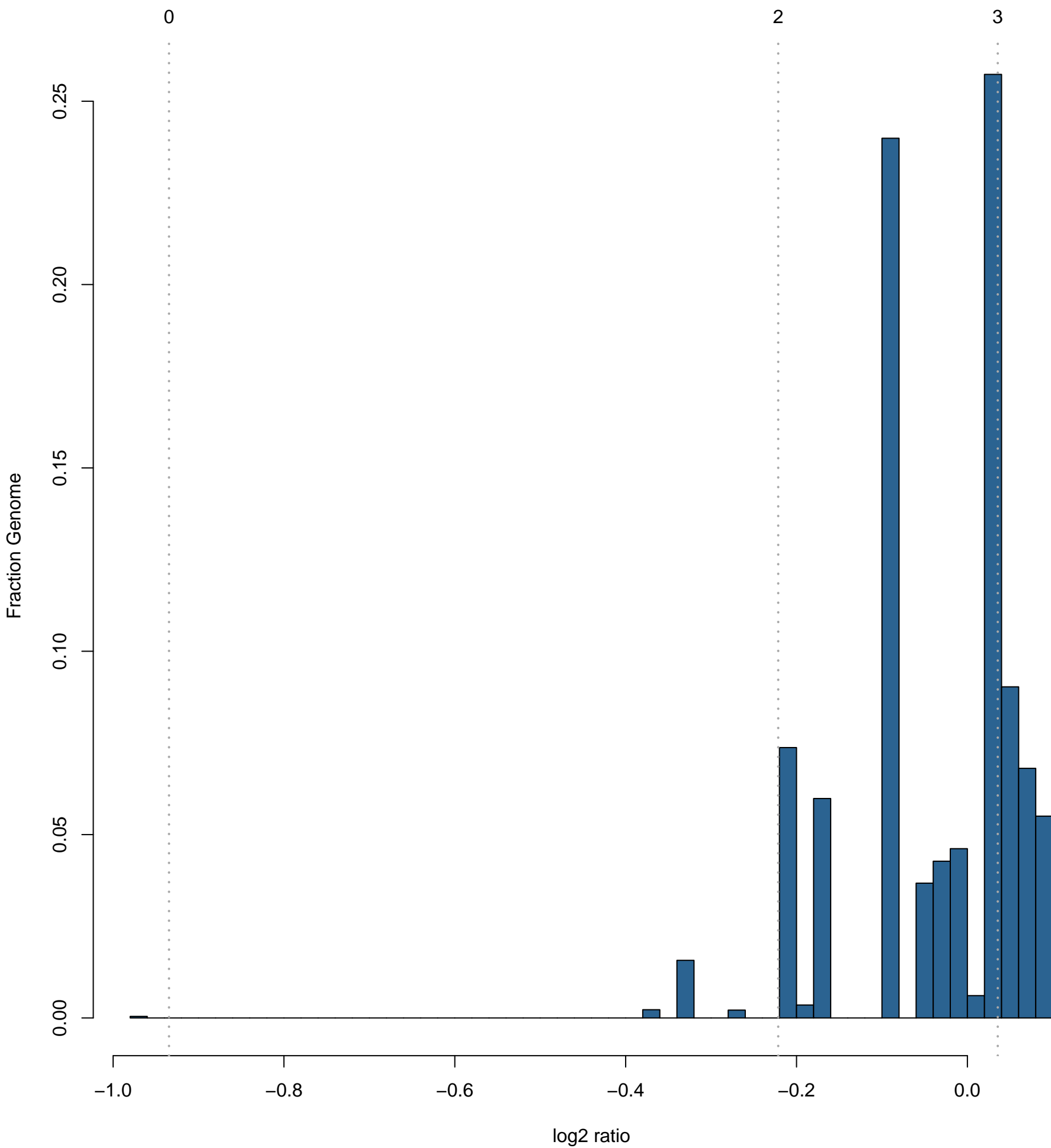


SCNA-fit log-likelihood: -601.19



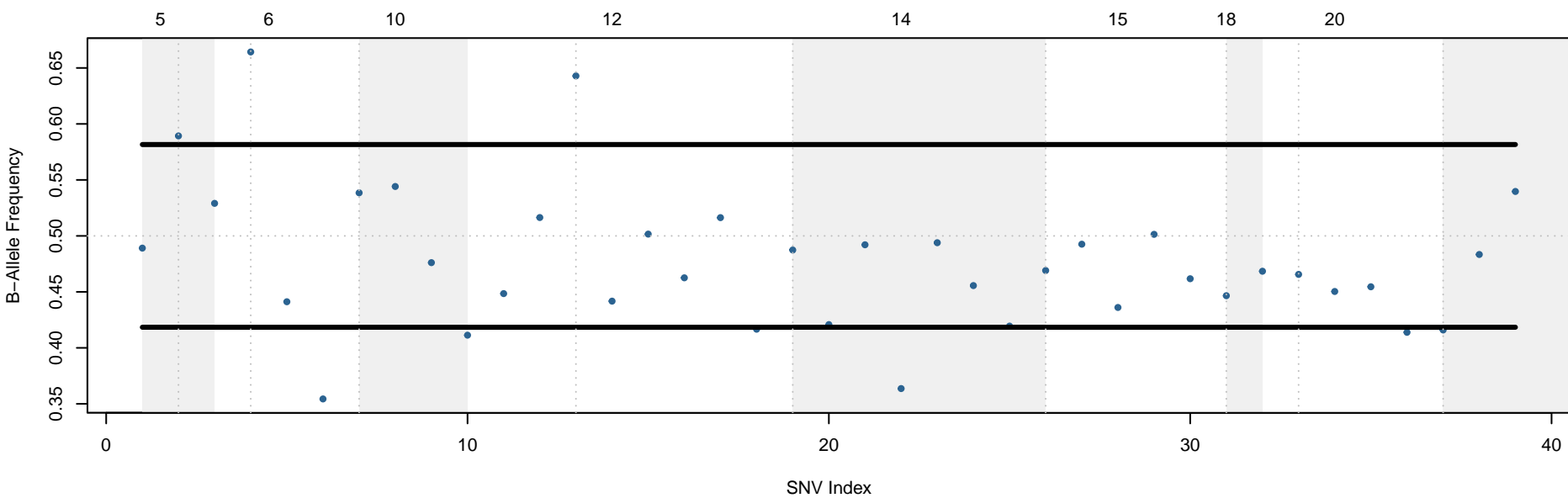


Purity: 0.39 Tumor ploidy: 2.851

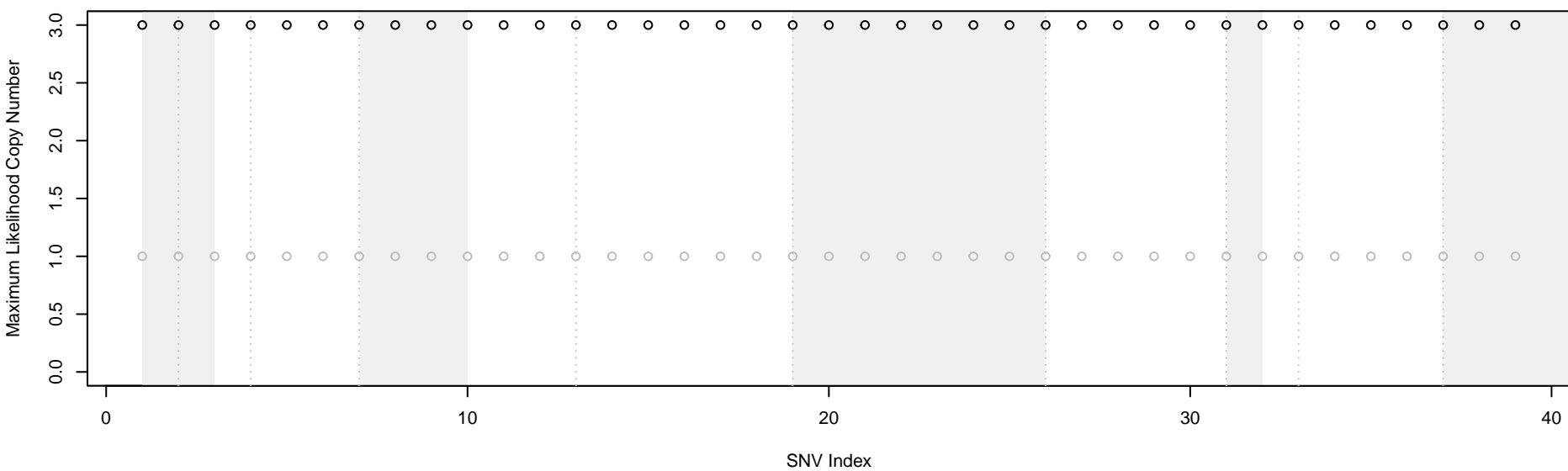
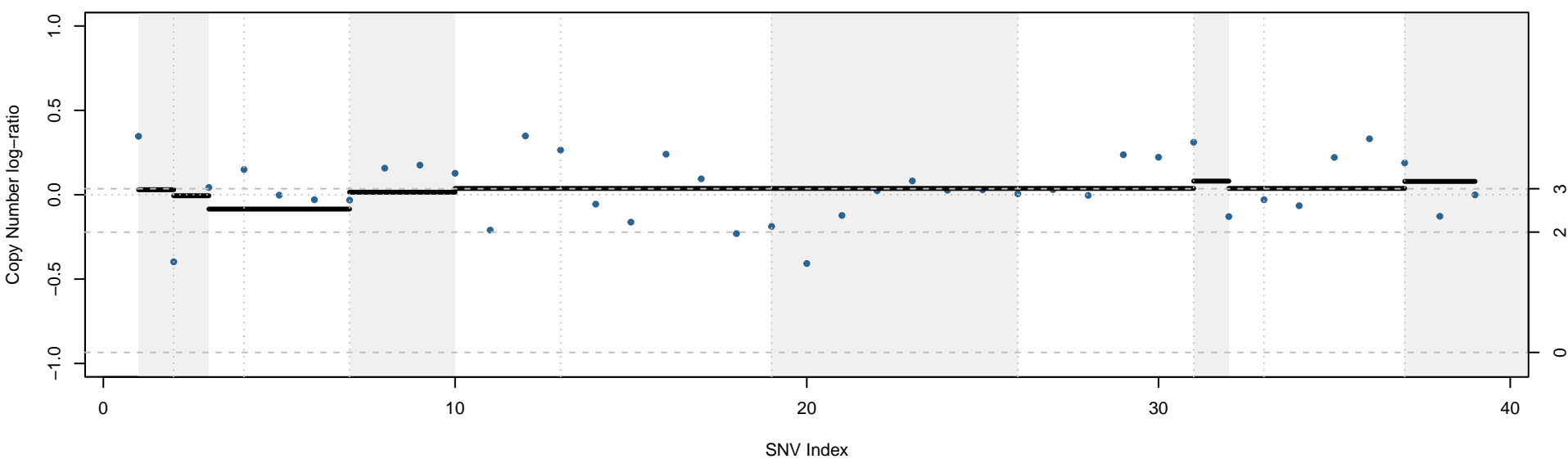


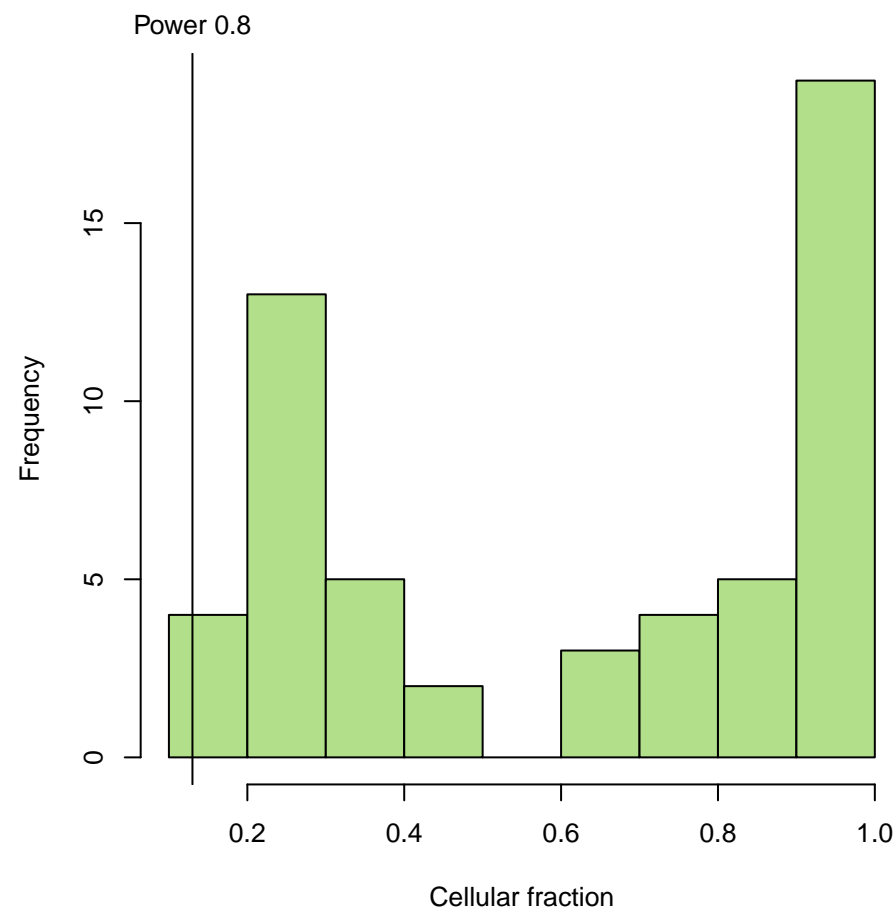
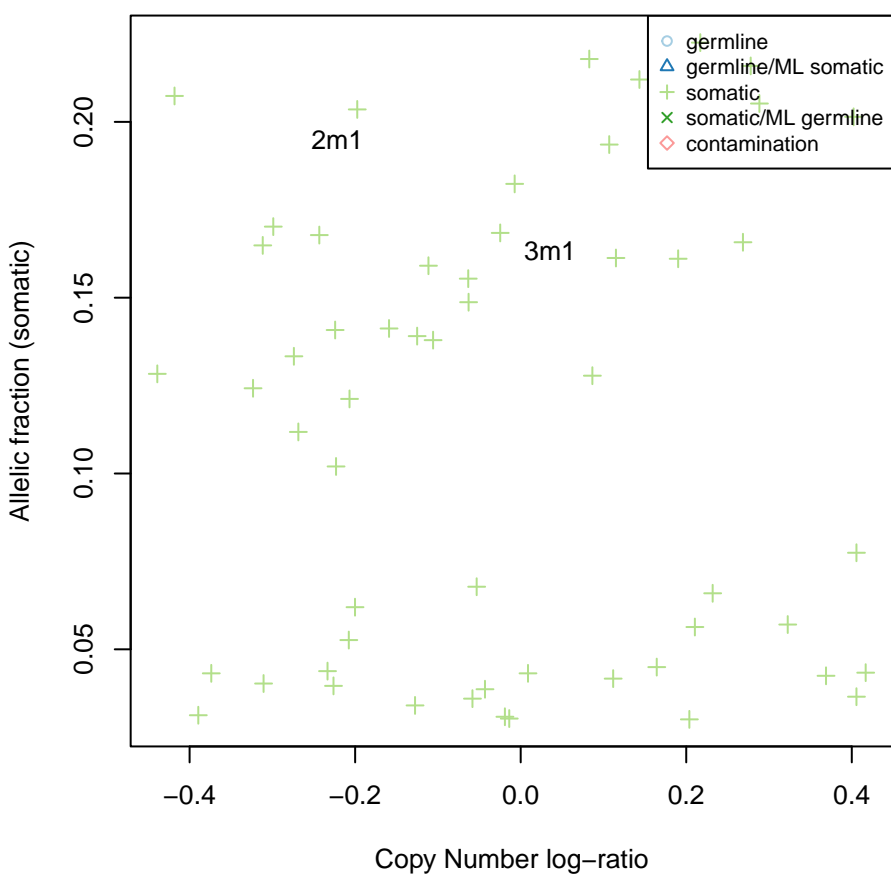
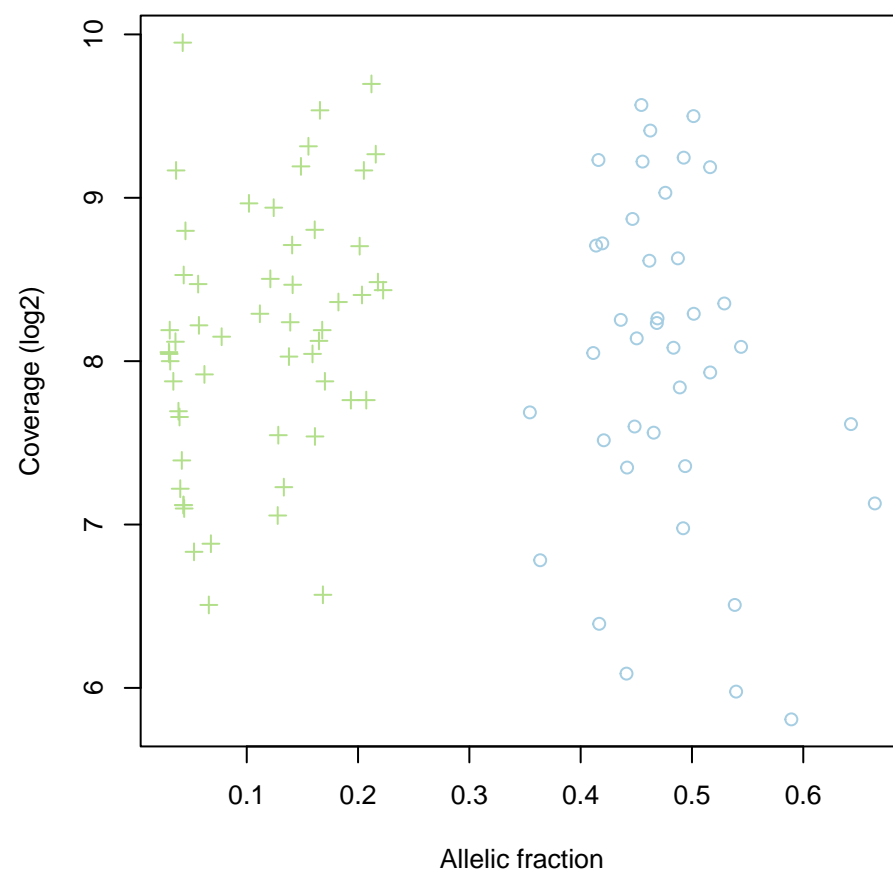
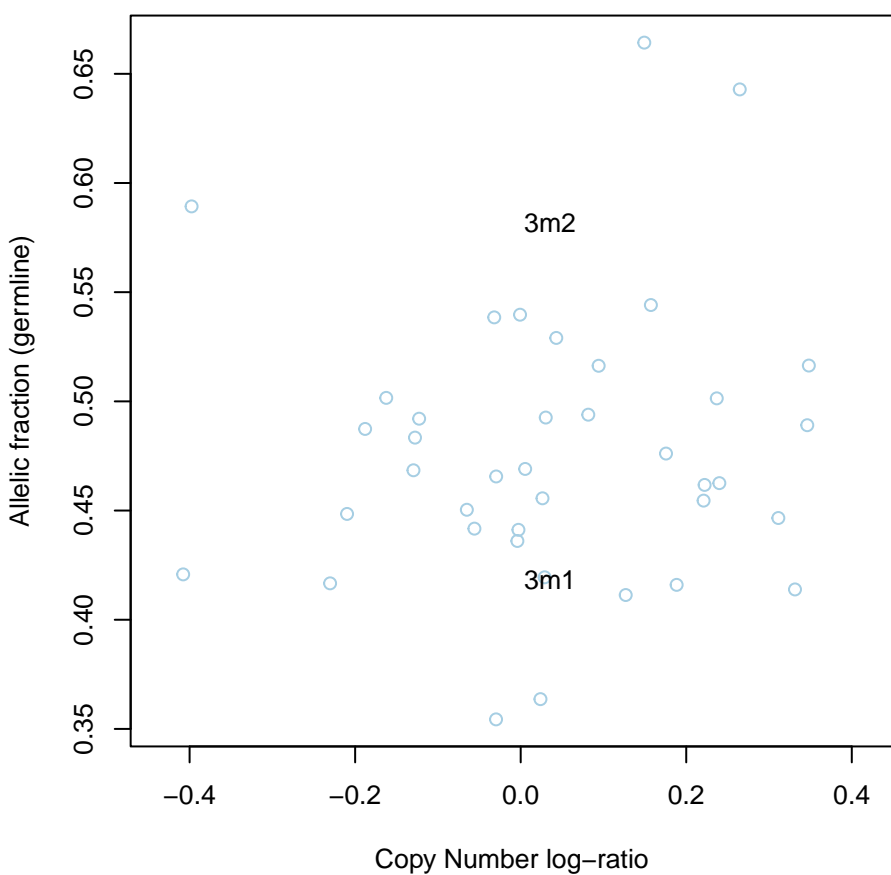


Purity: 0.39 Tumor ploidy: 2.851 SNV log-likelihood: -66.02 GoF: 86.6% Mean coverage: 315;324

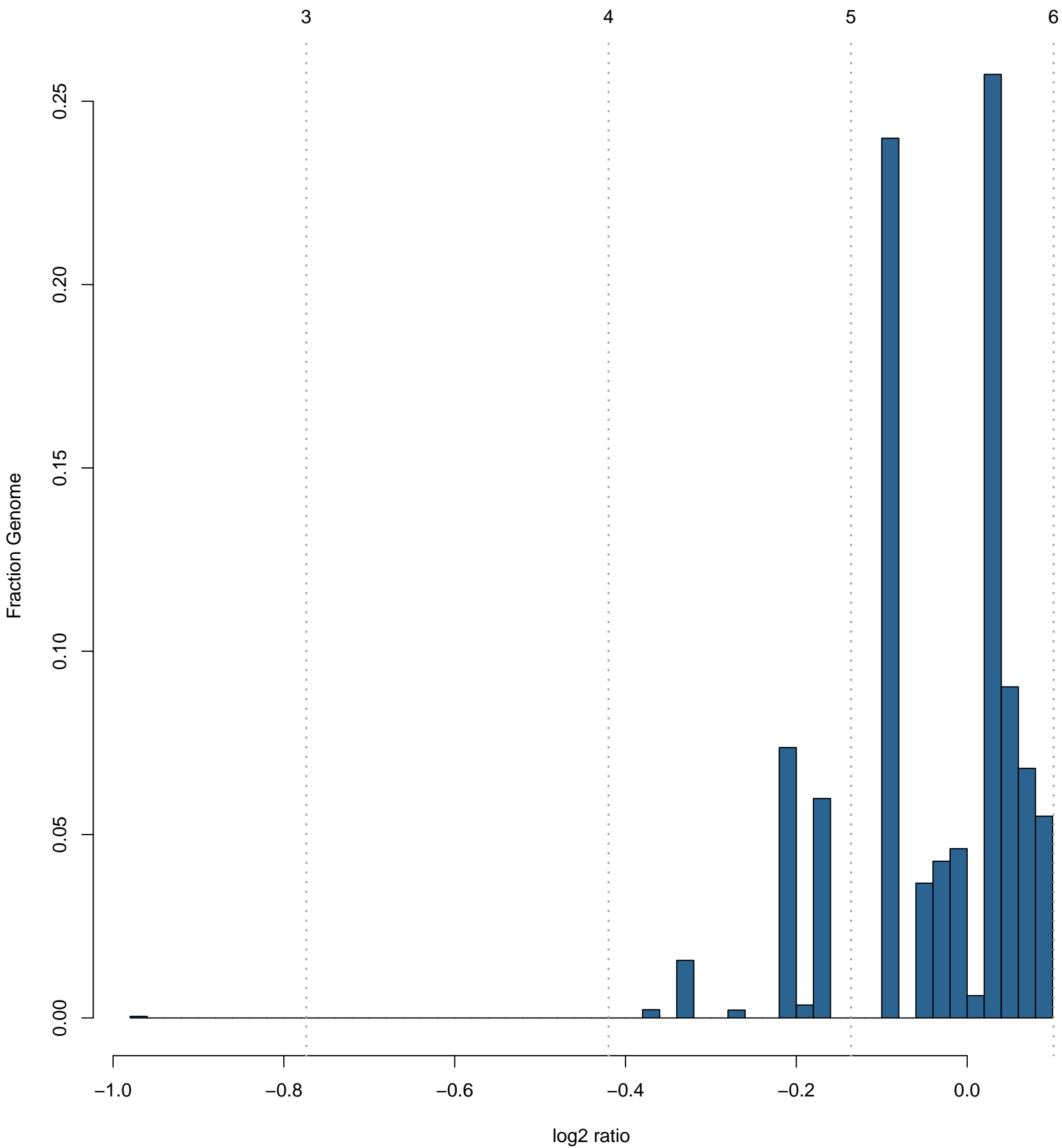


SCNA-fit log-likelihood: -619.87

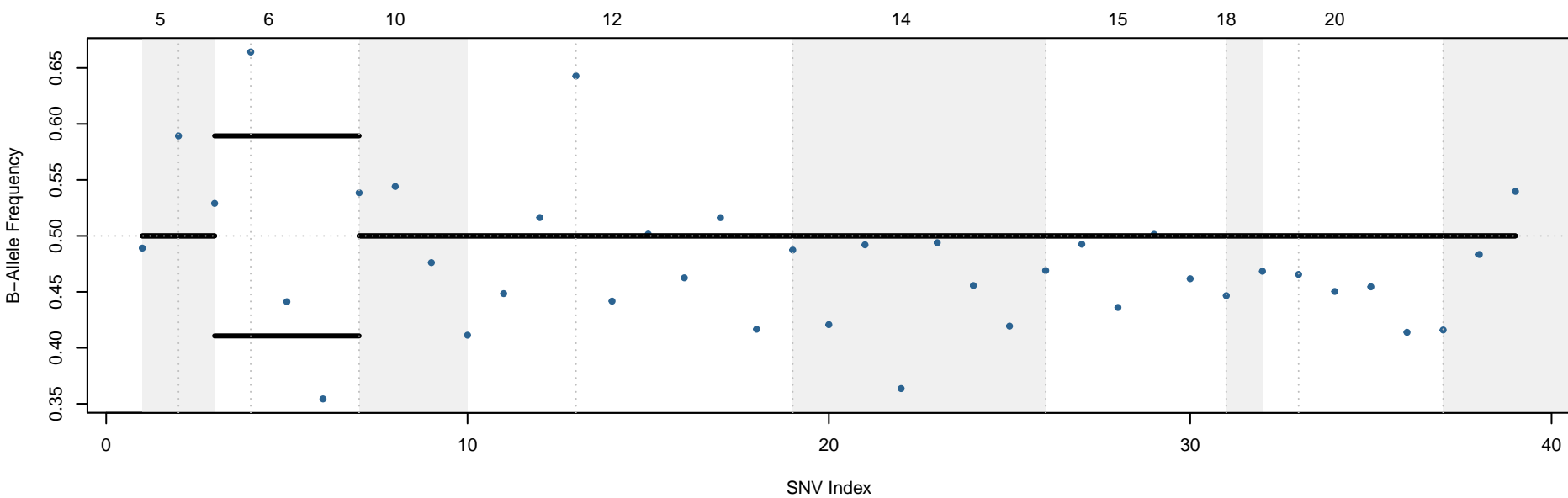




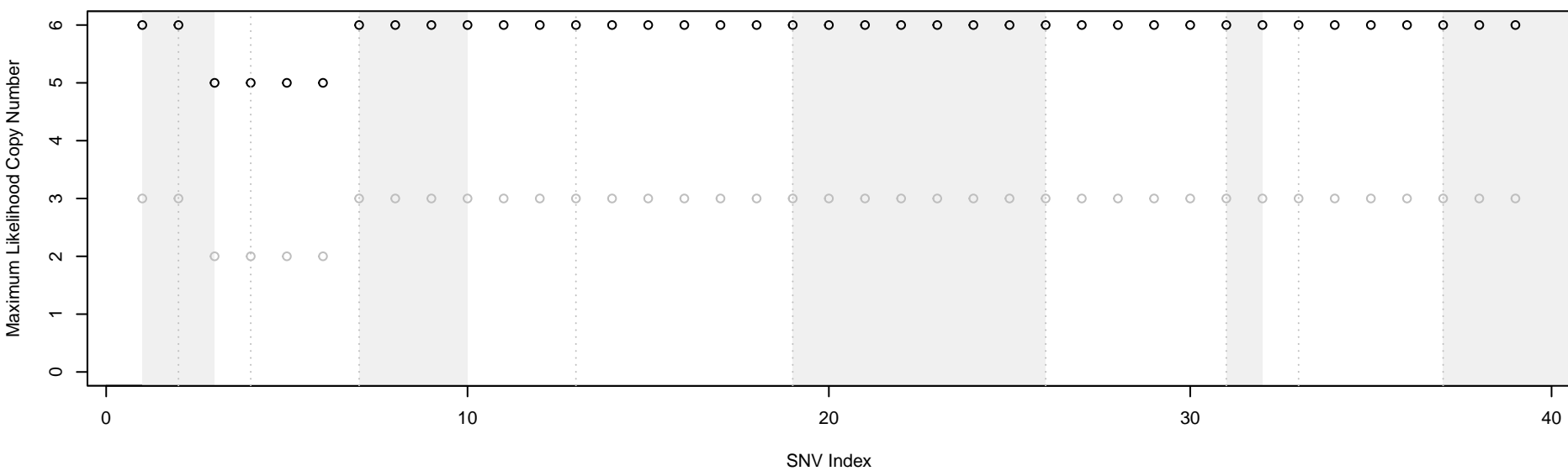
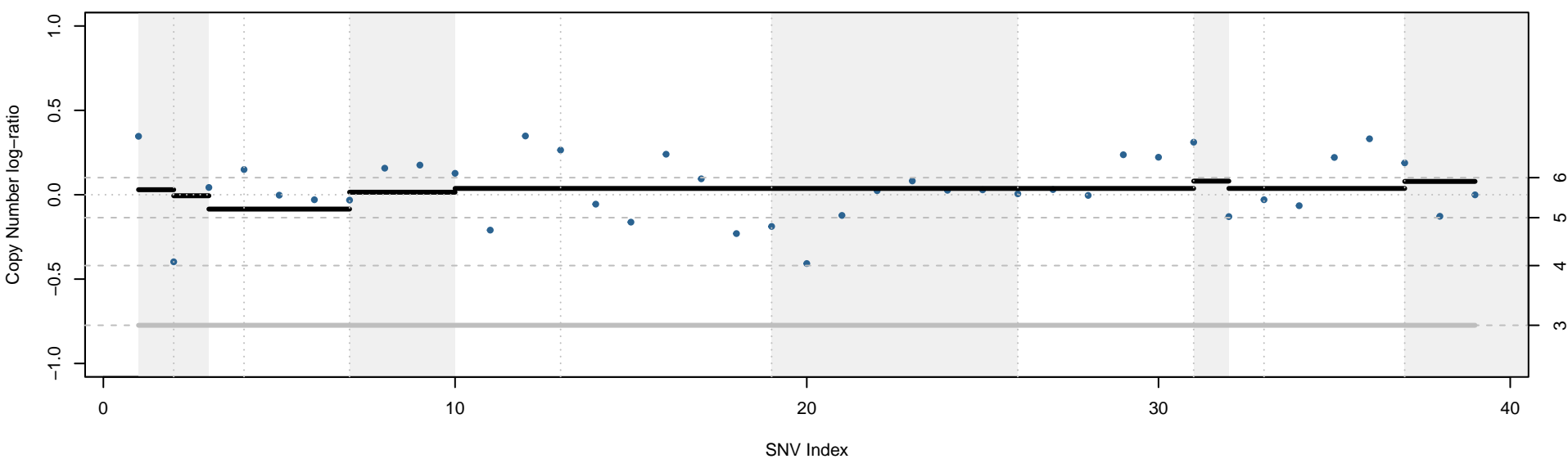
Purity: 0.77 Tumor ploidy: 5.553

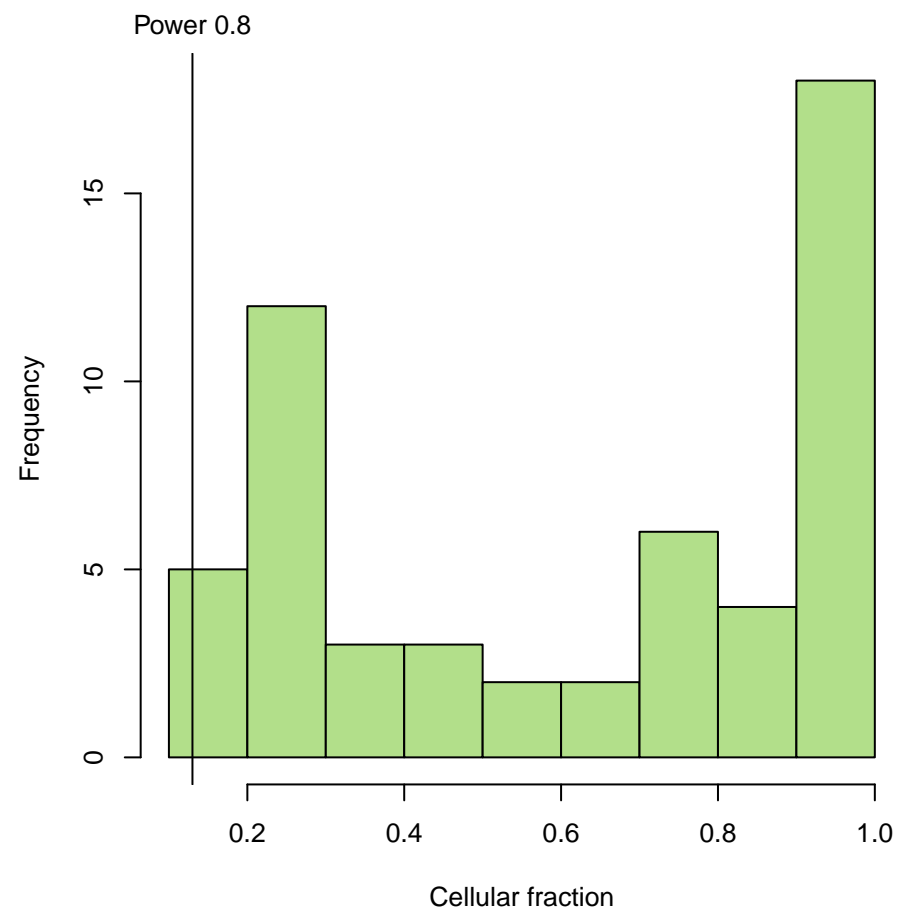
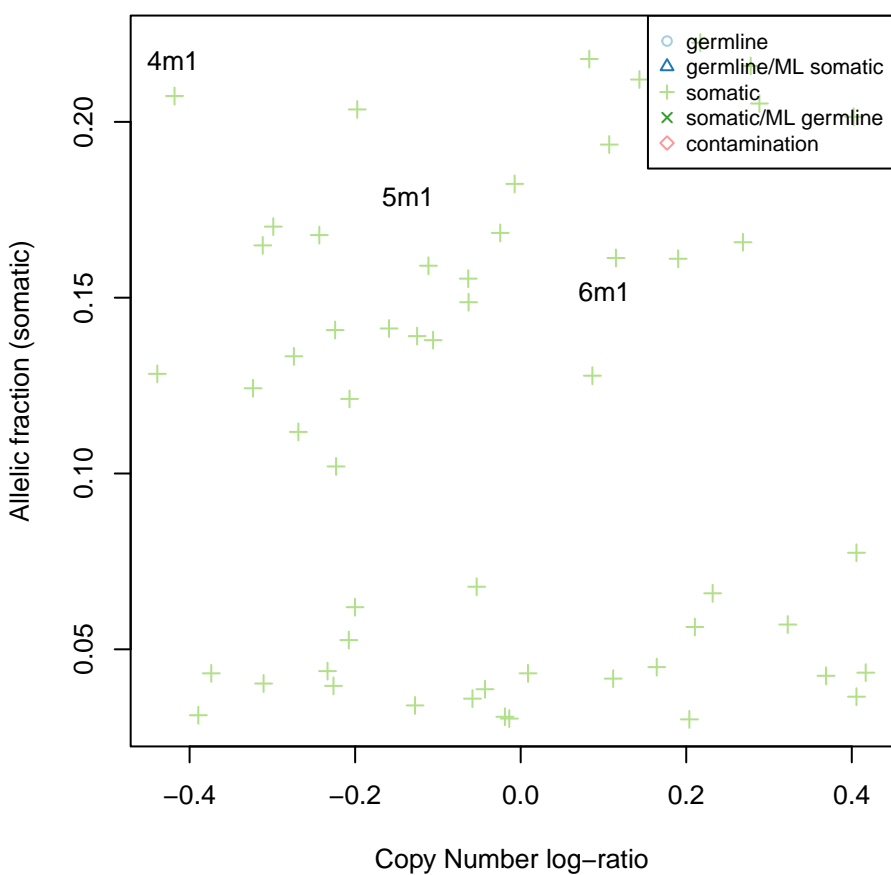
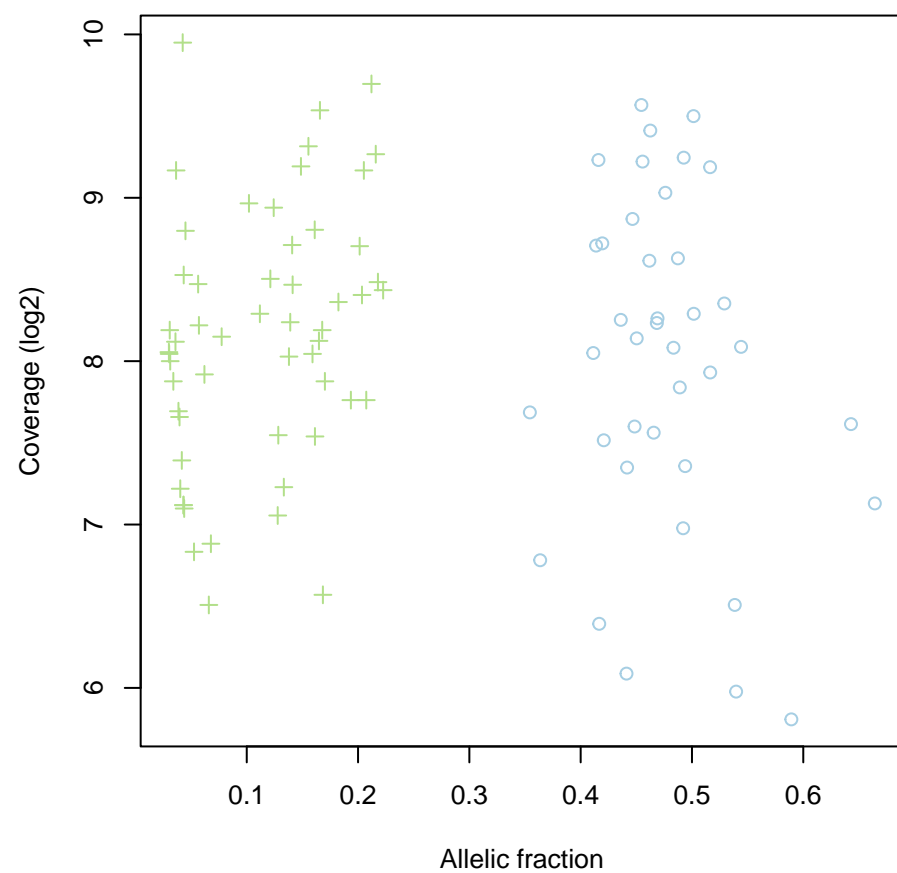
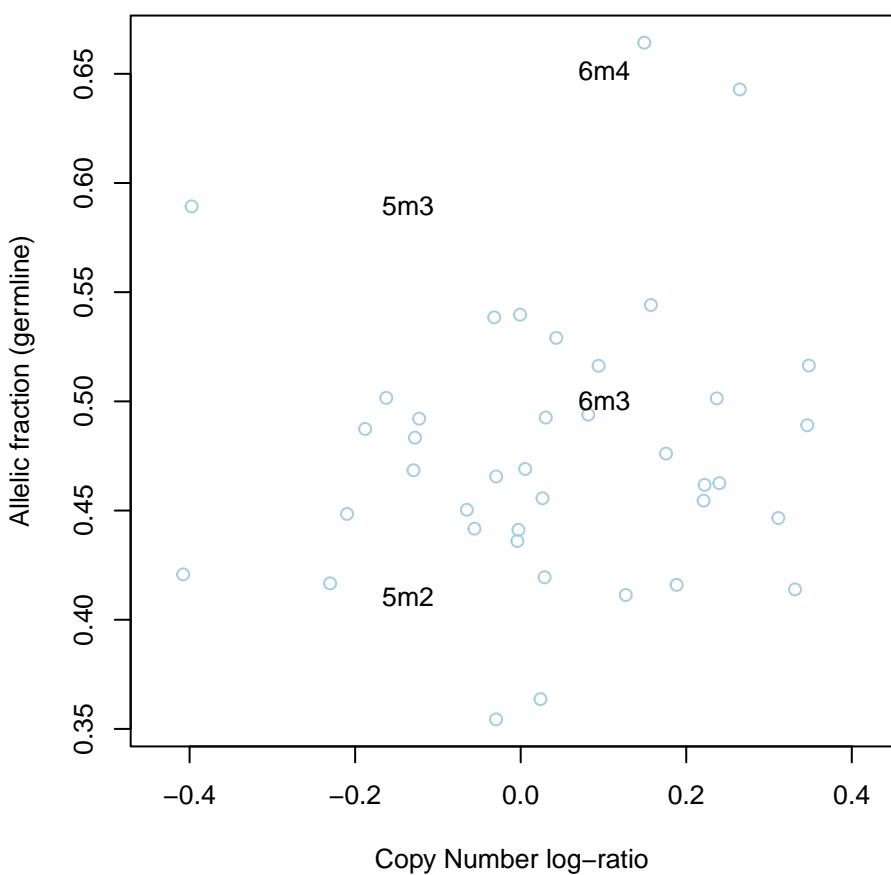


Purity: 0.77 Tumor ploidy: 5.553 SNV log-likelihood: -94.03 GoF: 87.6% Mean coverage: 315;324

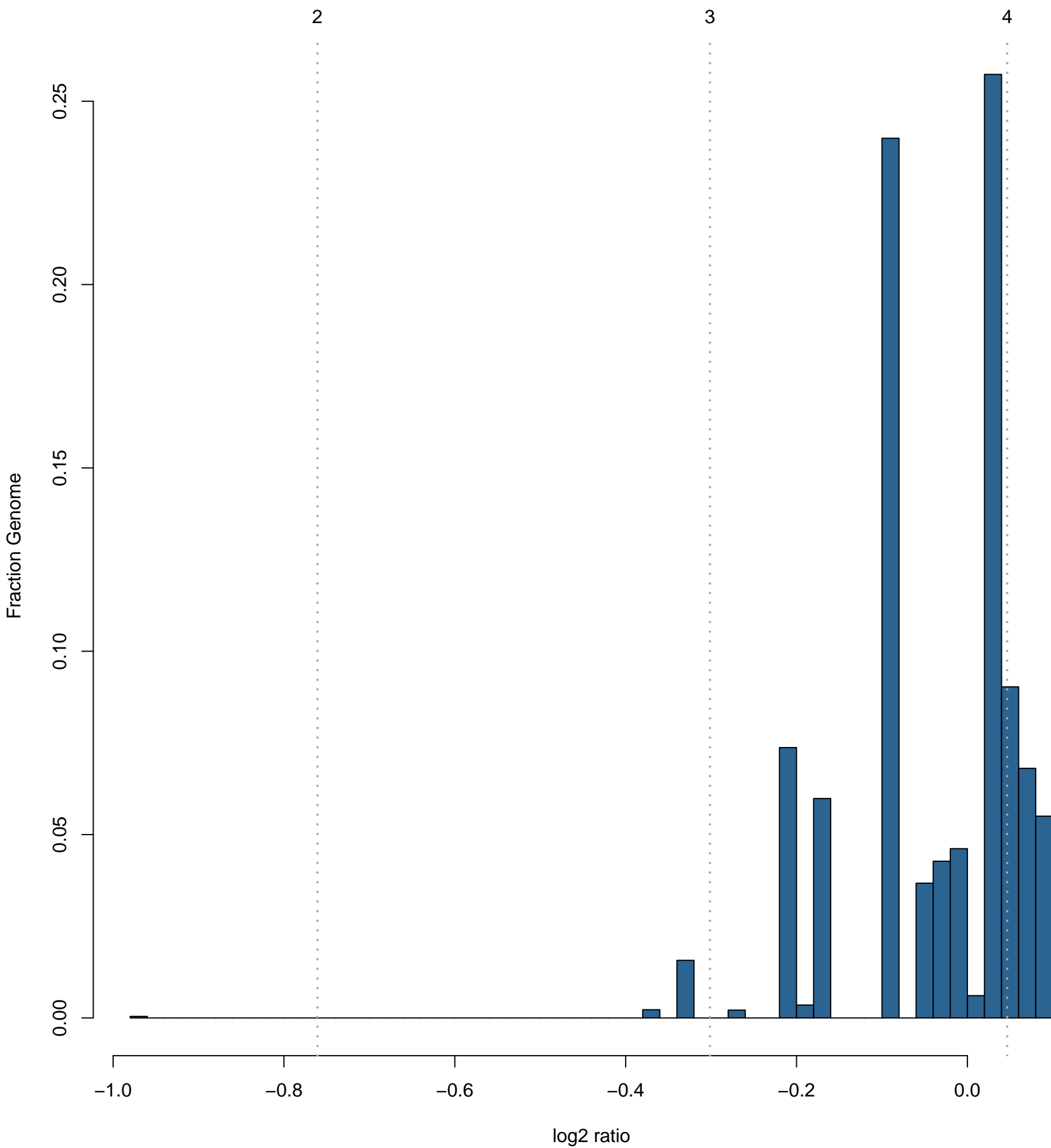


SCNA-fit log-likelihood: -576.51

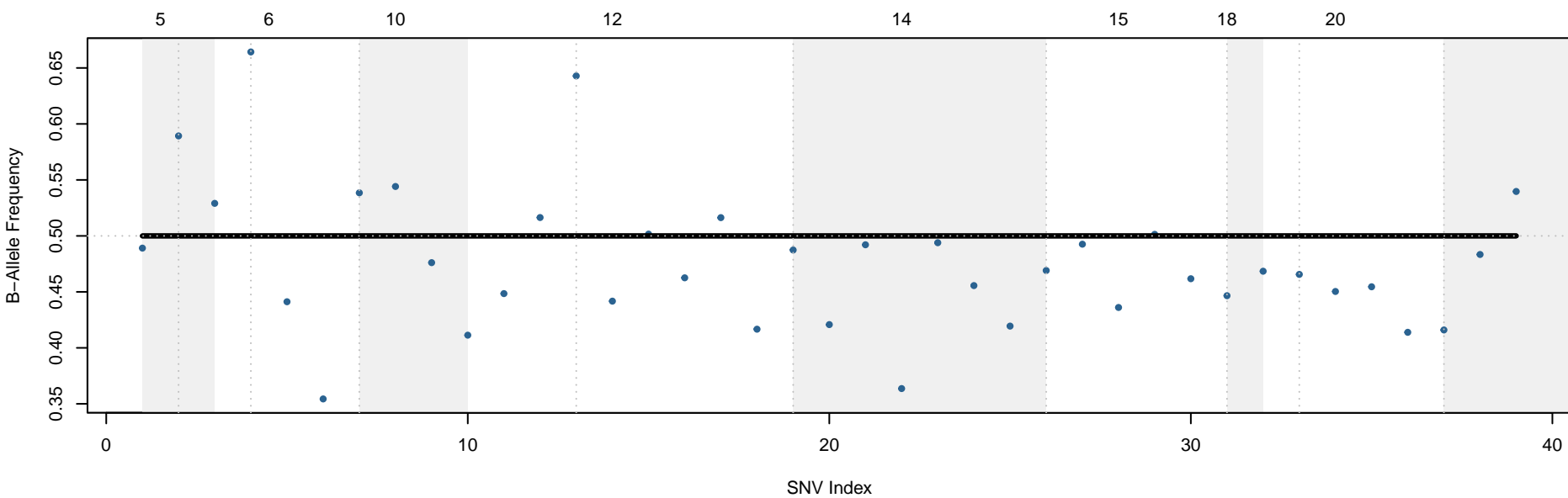




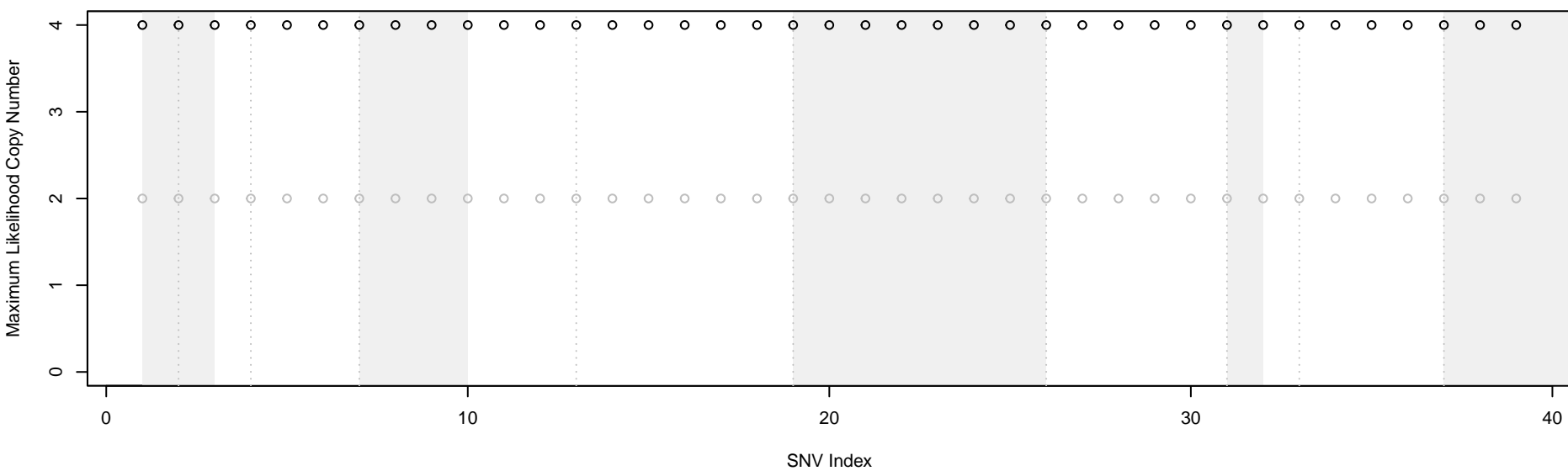
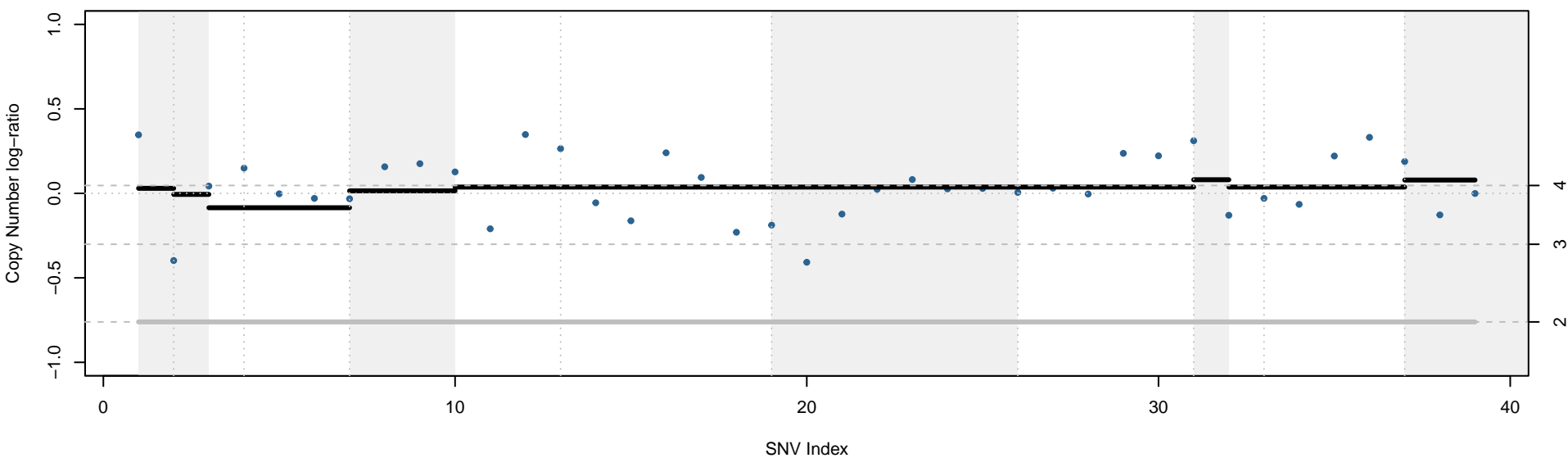
Purity: 0.75 Tumor ploidy: 3.852

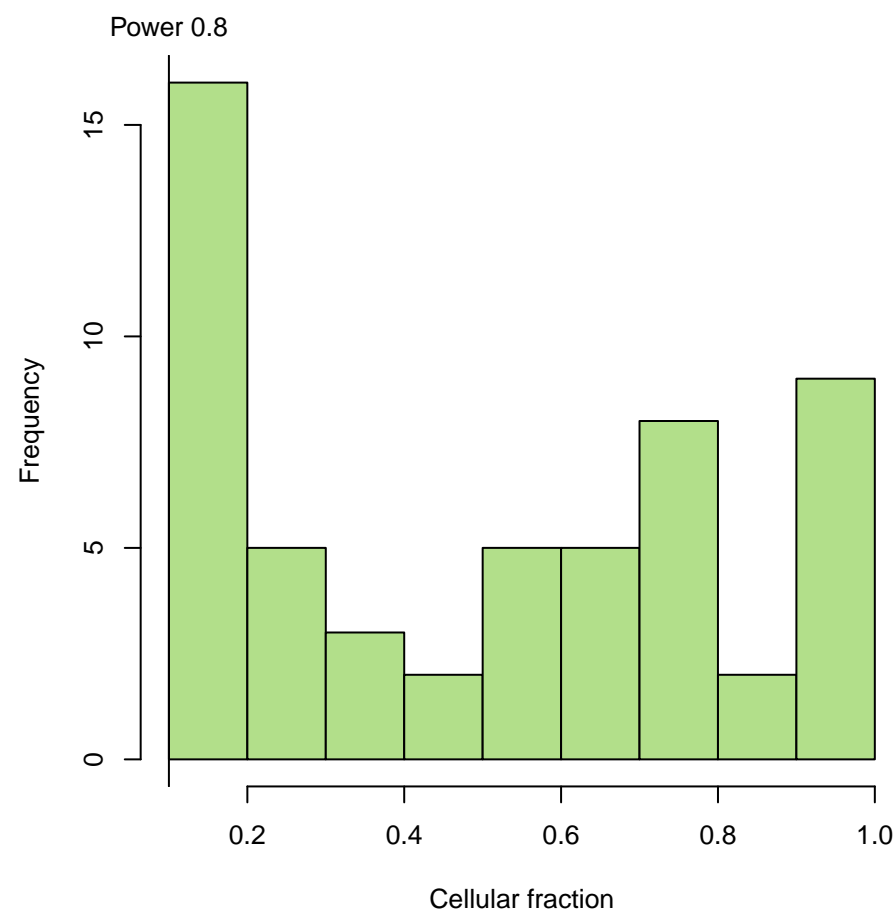
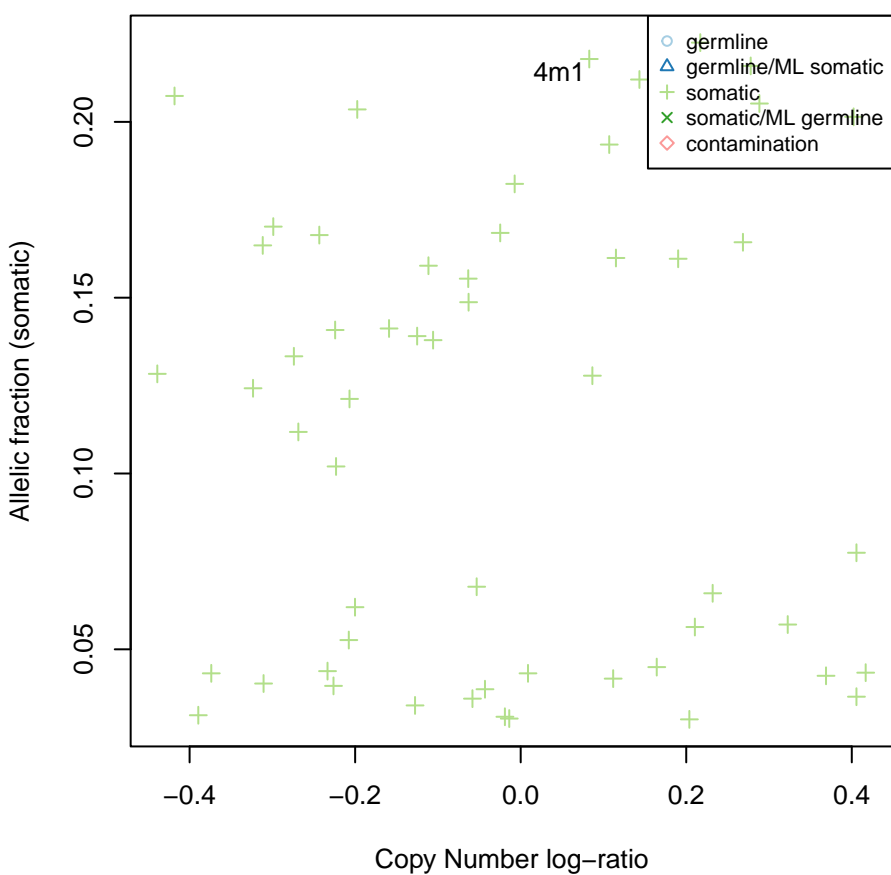
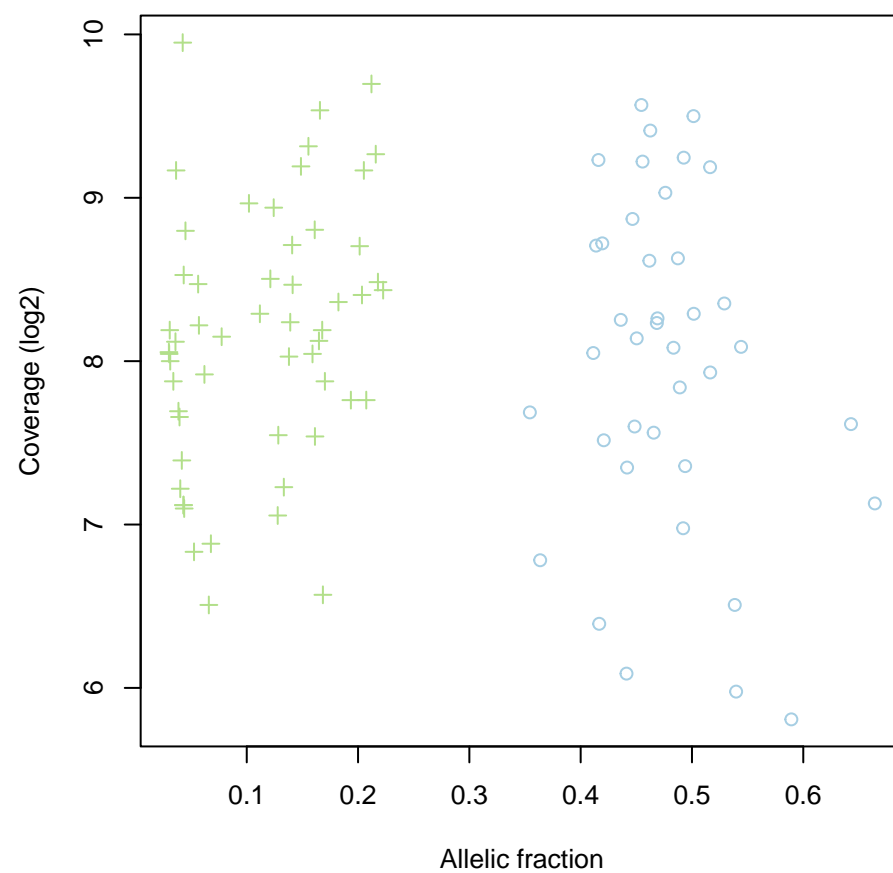
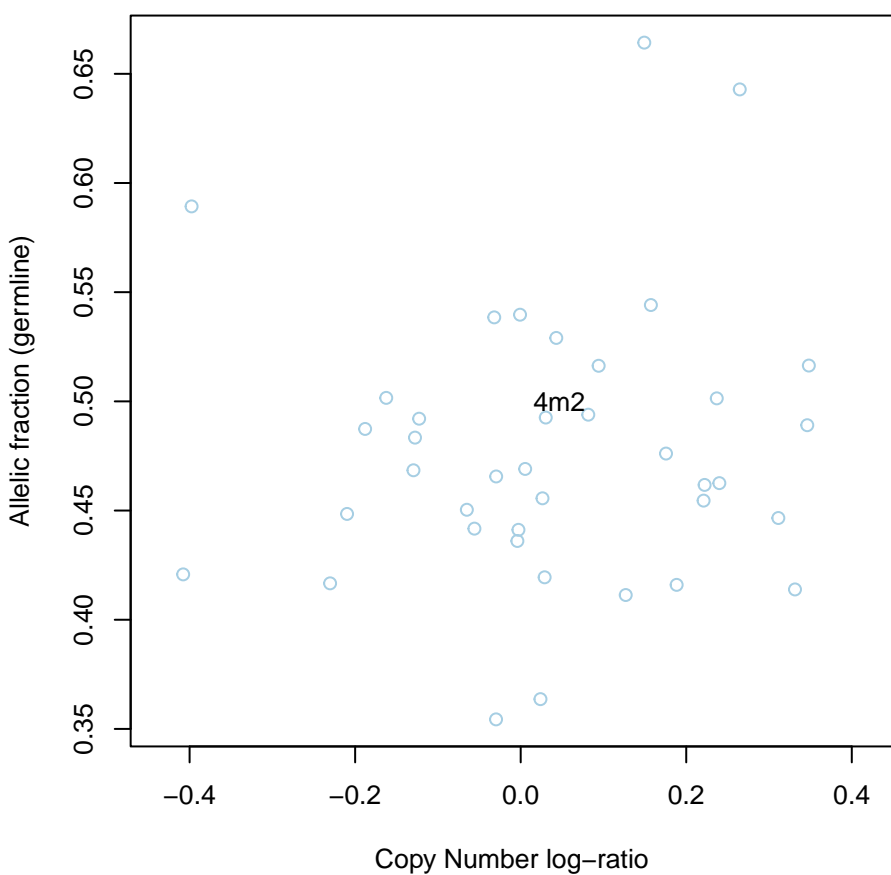


Purity: 0.75 Tumor ploidy: 3.852 SNV log-likelihood: -122.28 GoF: 73.2% Mean coverage: 315,324



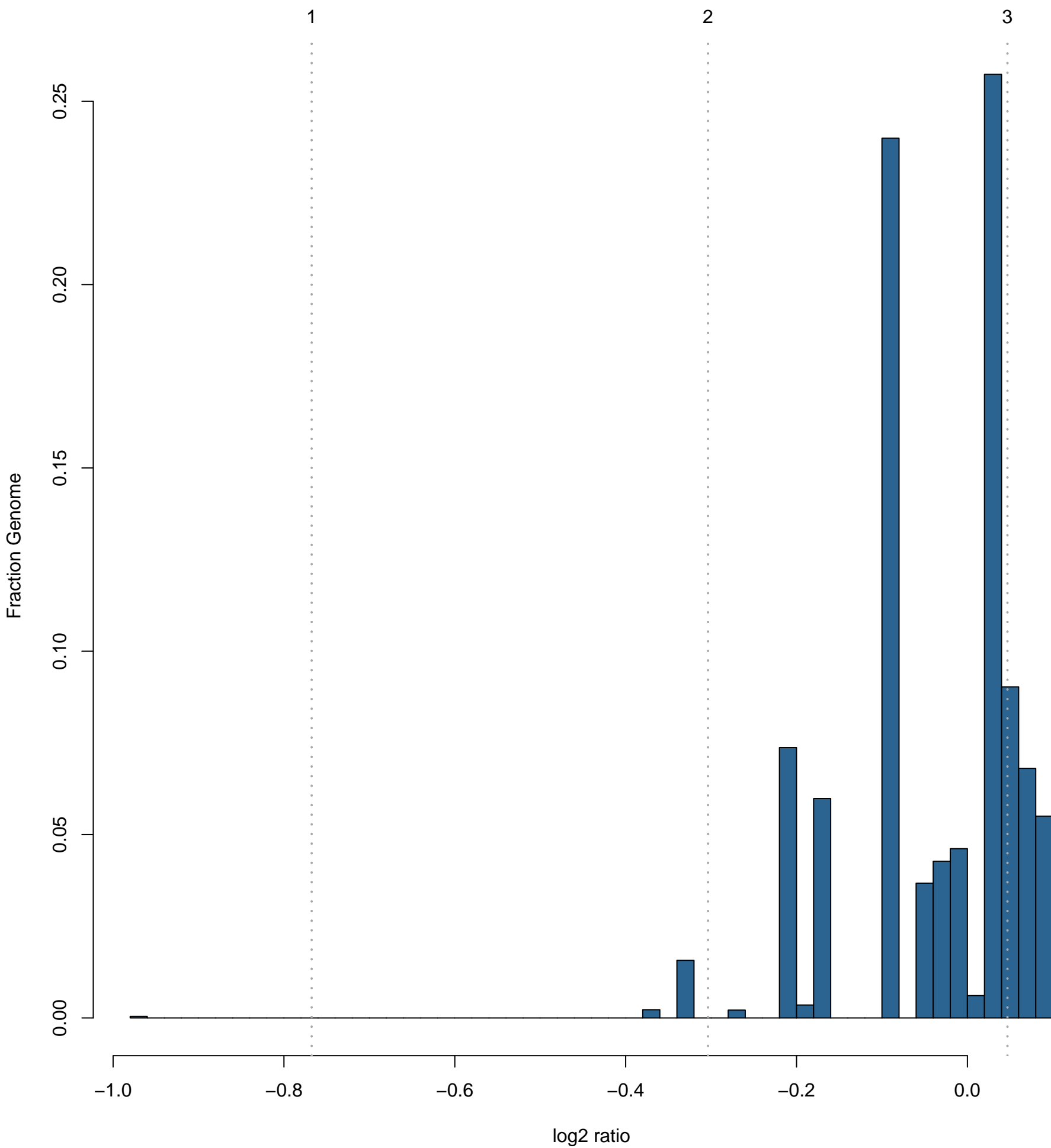
SCNA-fit log-likelihood: -767.45



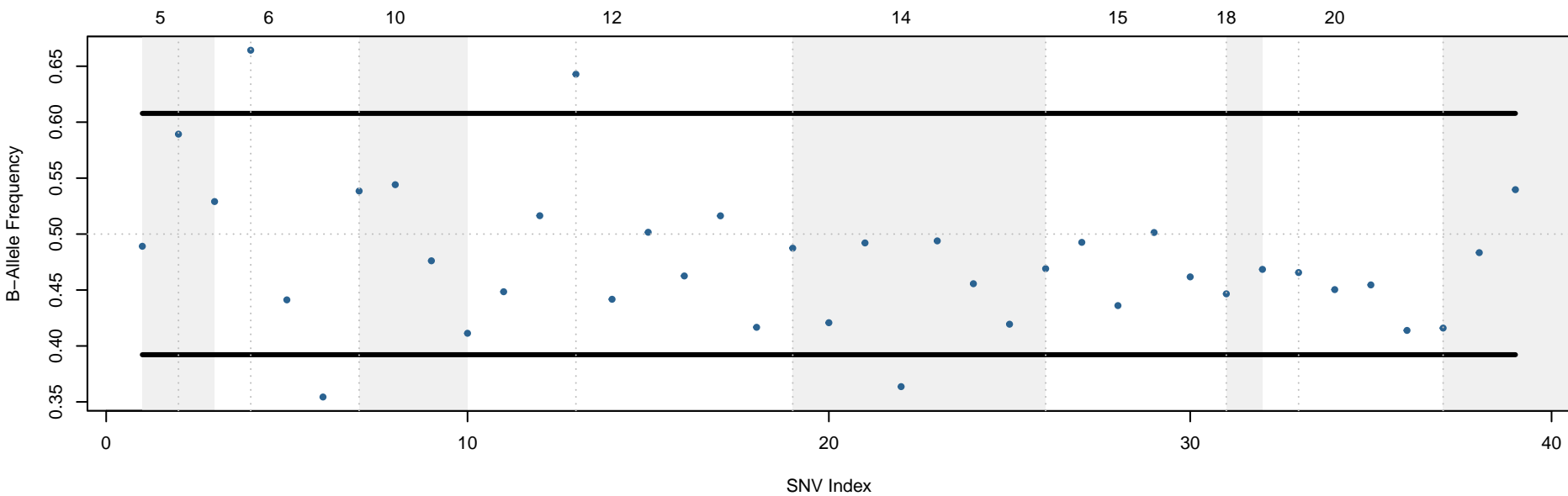




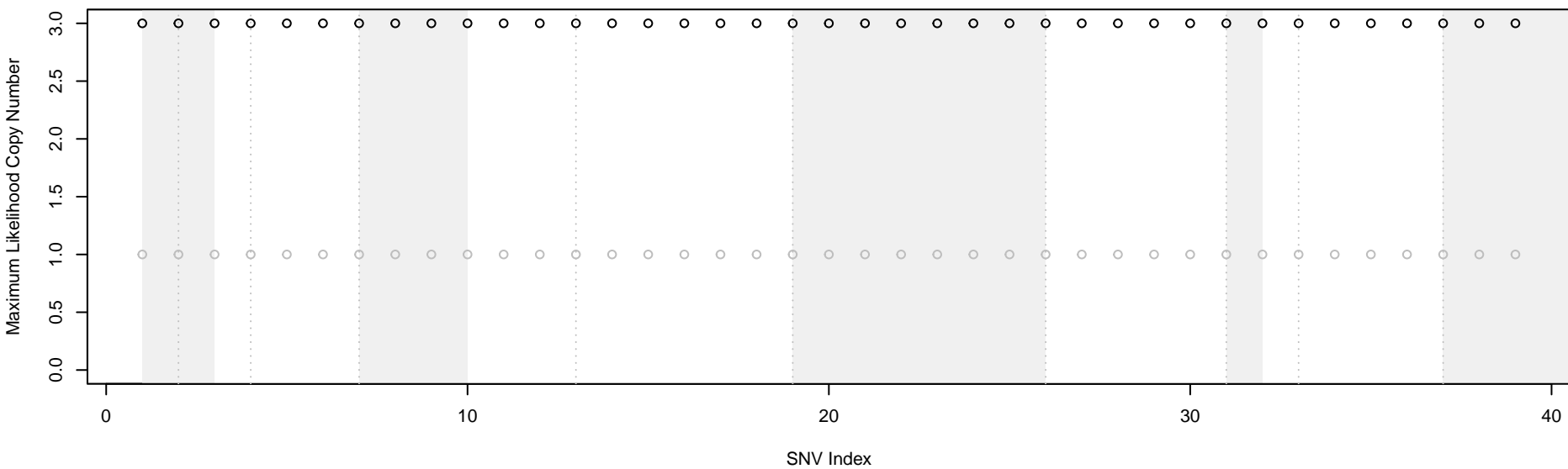
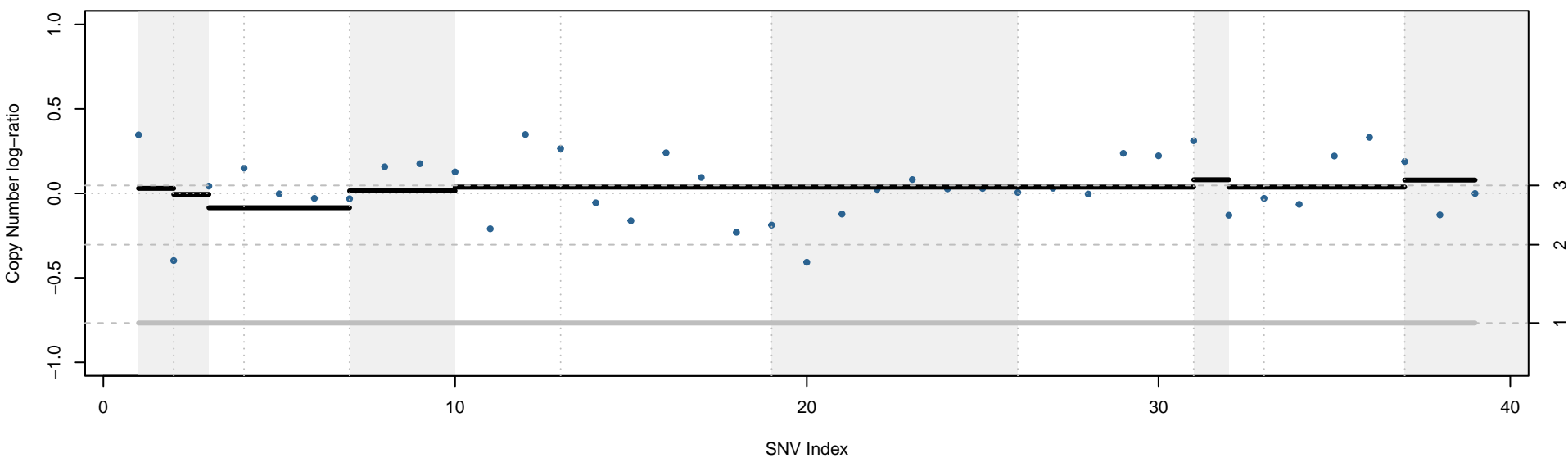
Purity: 0.55 Tumor ploidy: 2.852

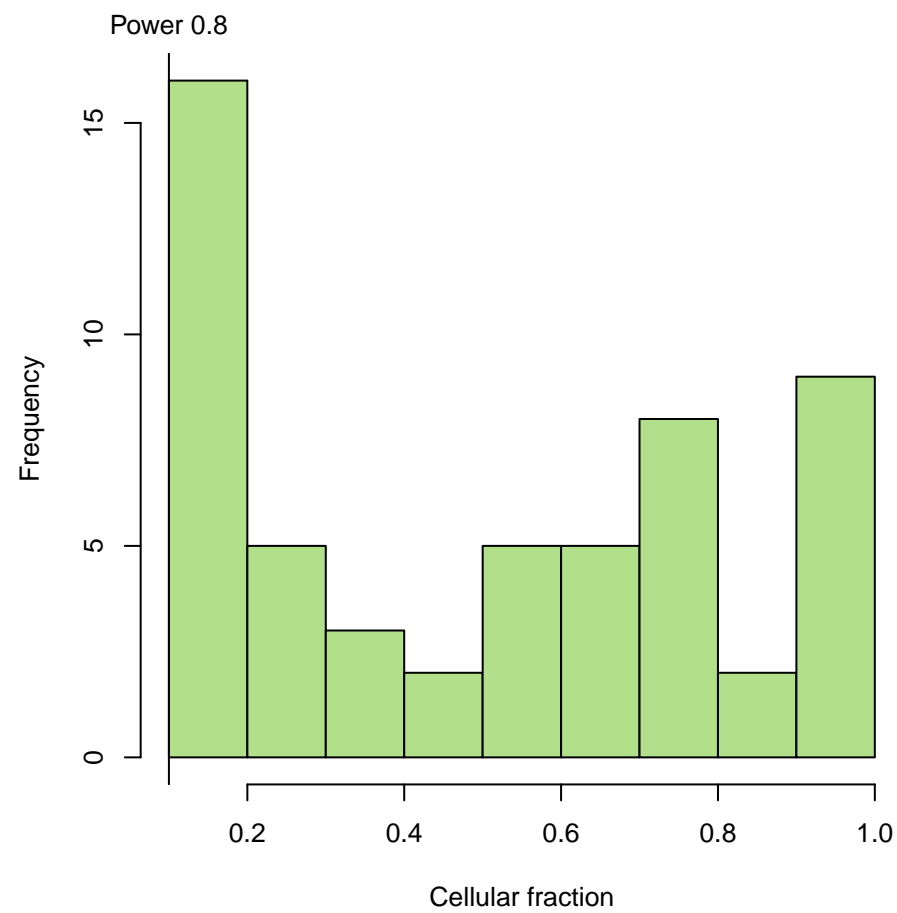
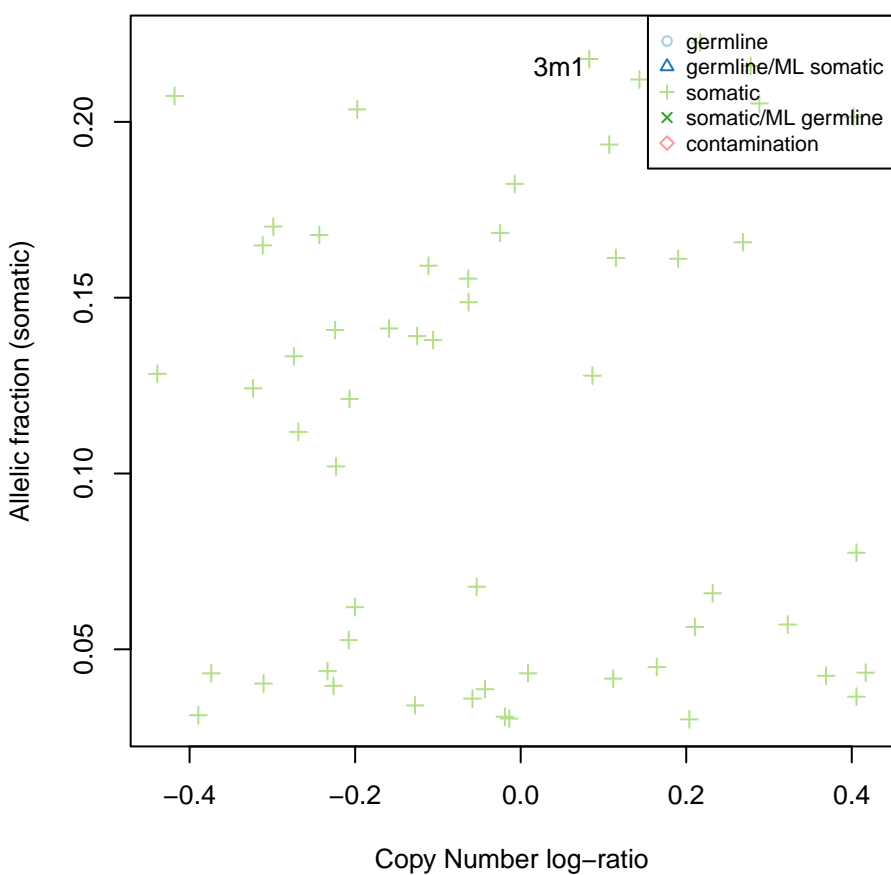
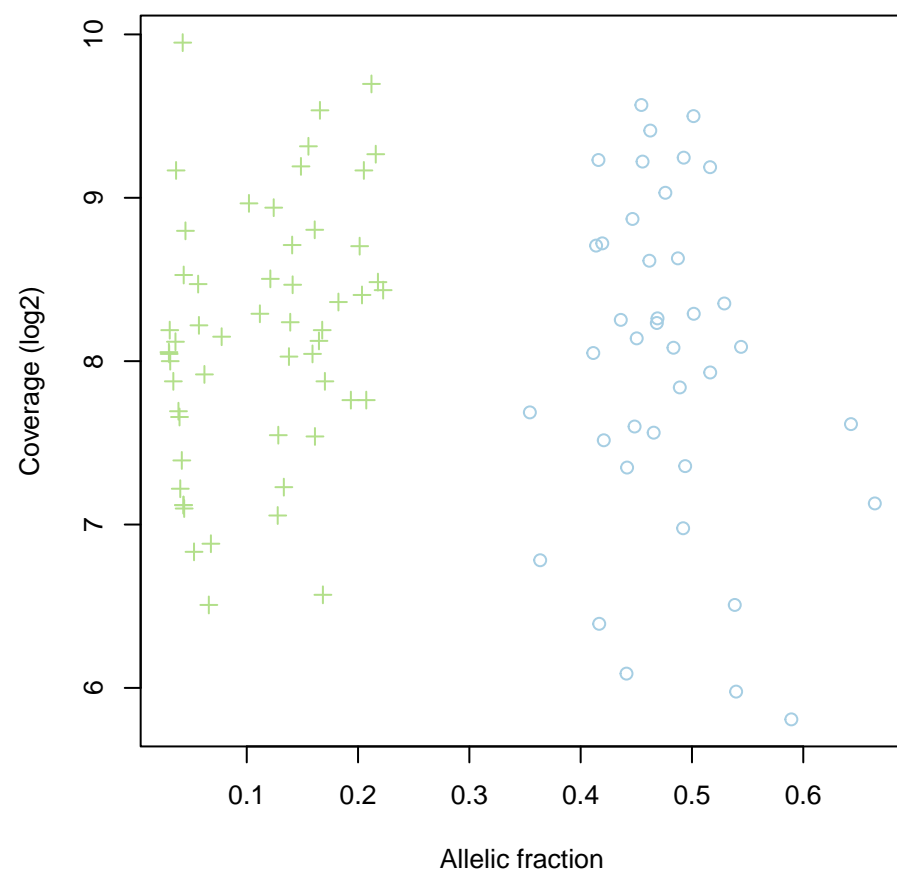
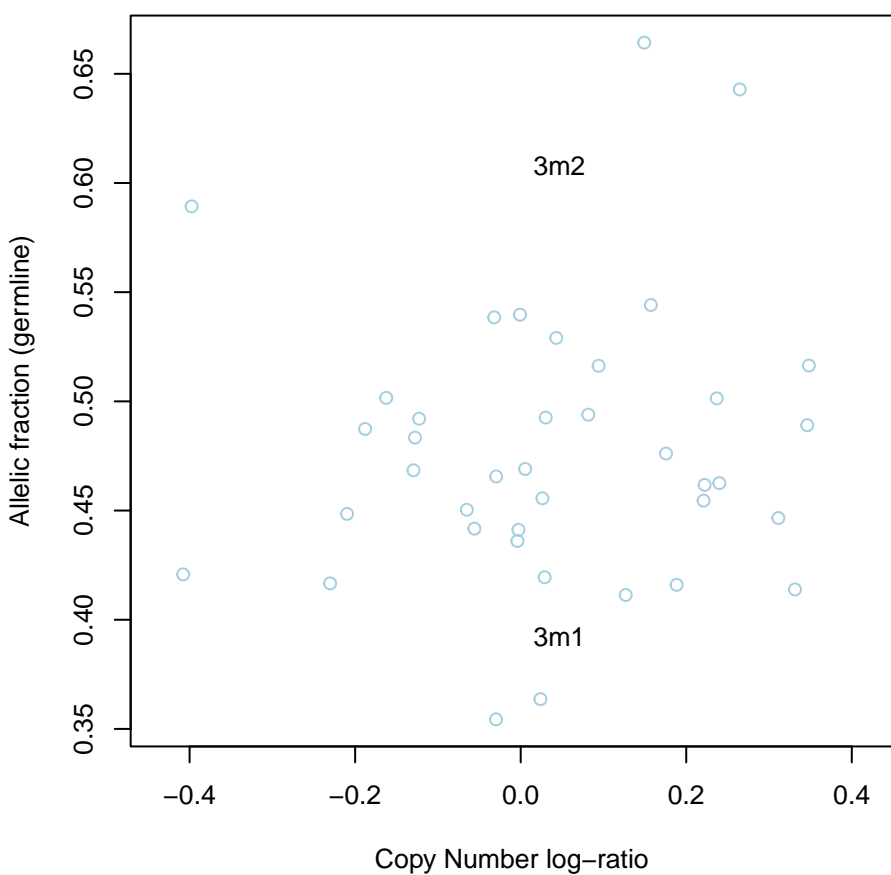


Purity: 0.55 Tumor ploidy: 2.852 SNV log-likelihood: -168.92 GoF: 71% Mean coverage: 315;324

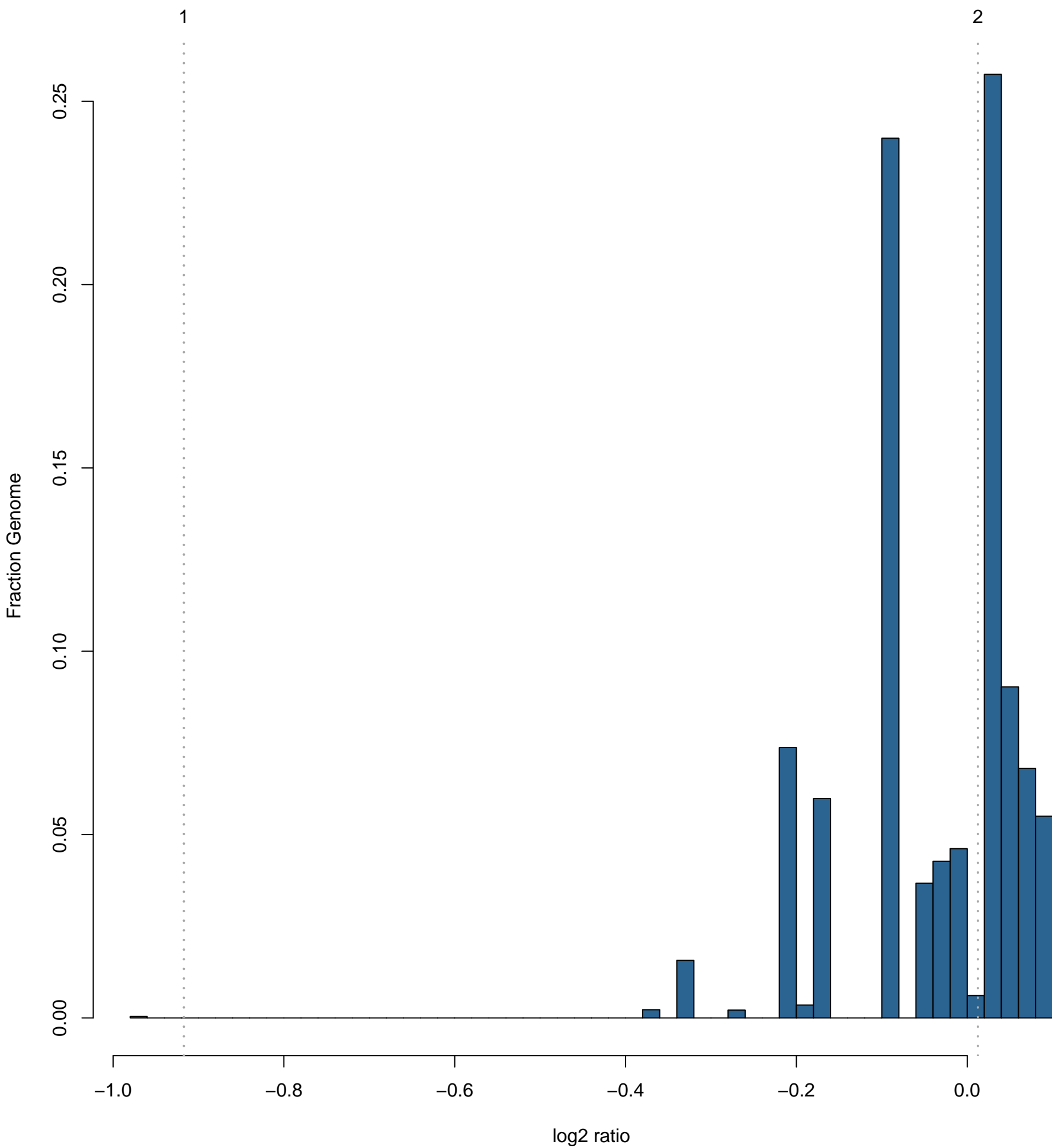


SCNA-fit log-likelihood: -780.31

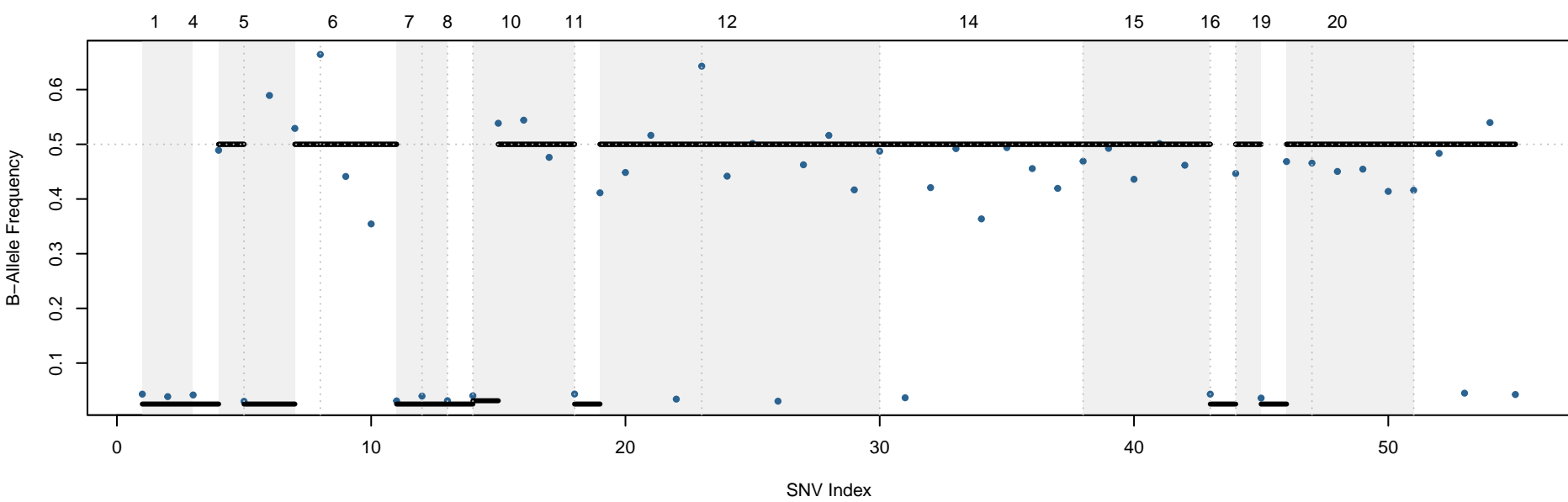




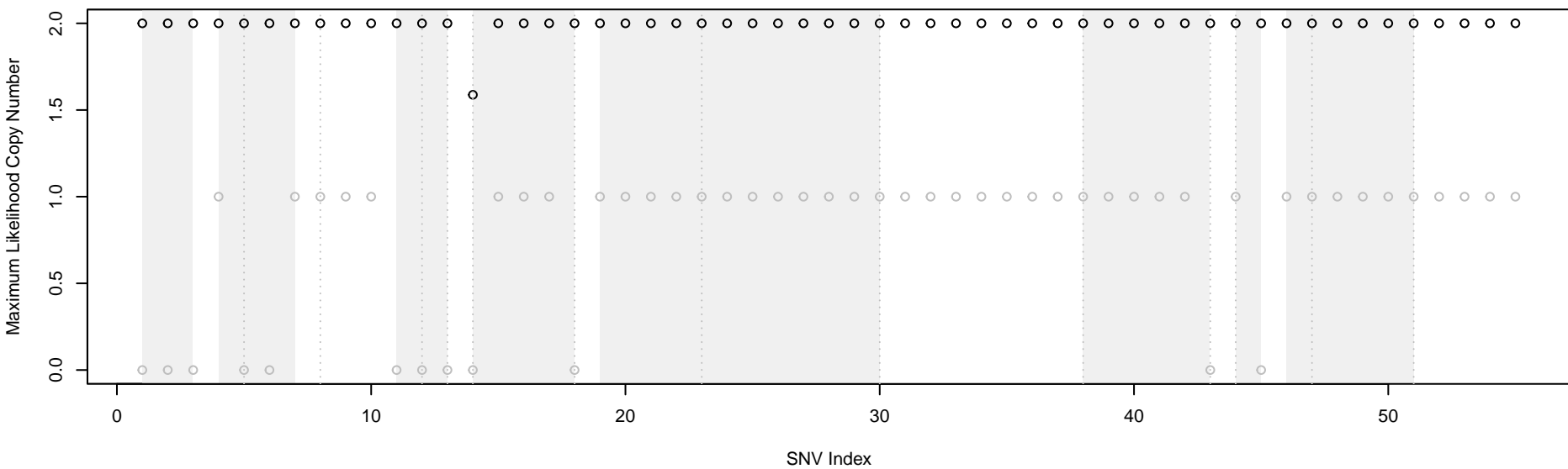
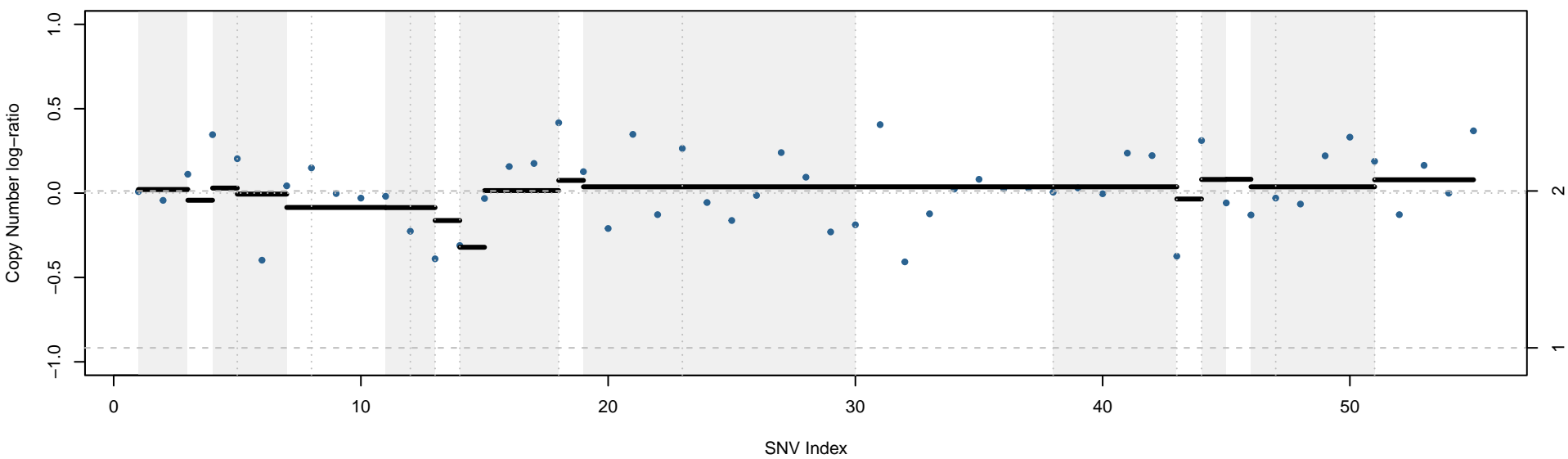
Purity: 0.95 Tumor ploidy: 1.982

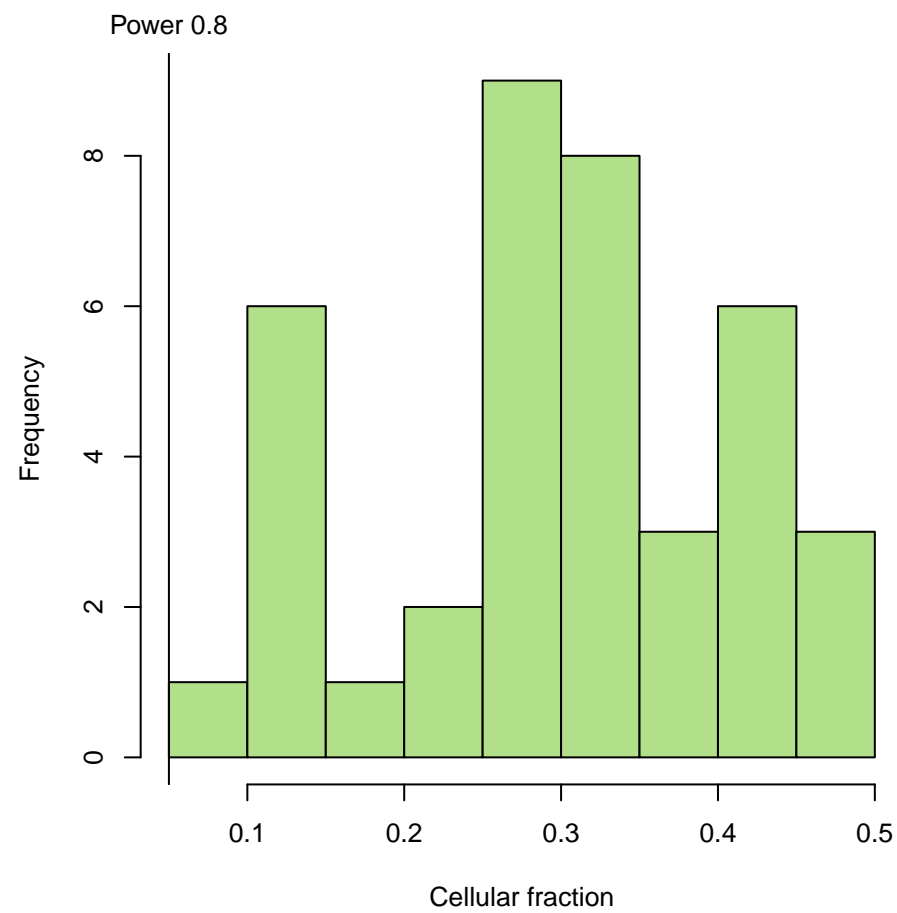
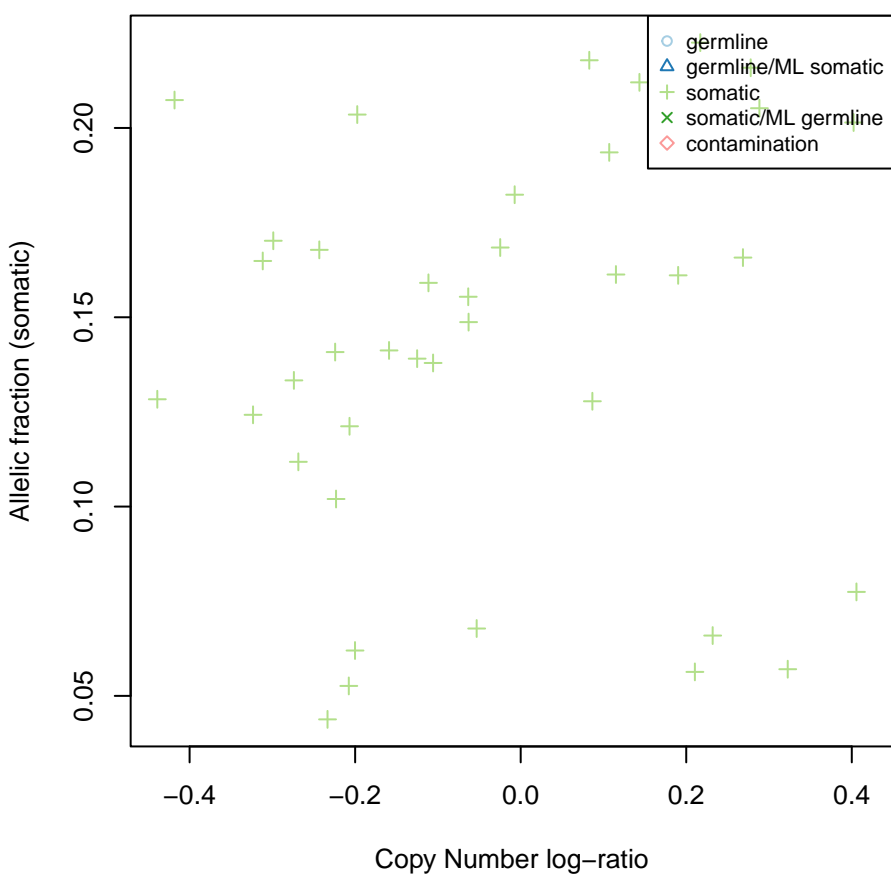
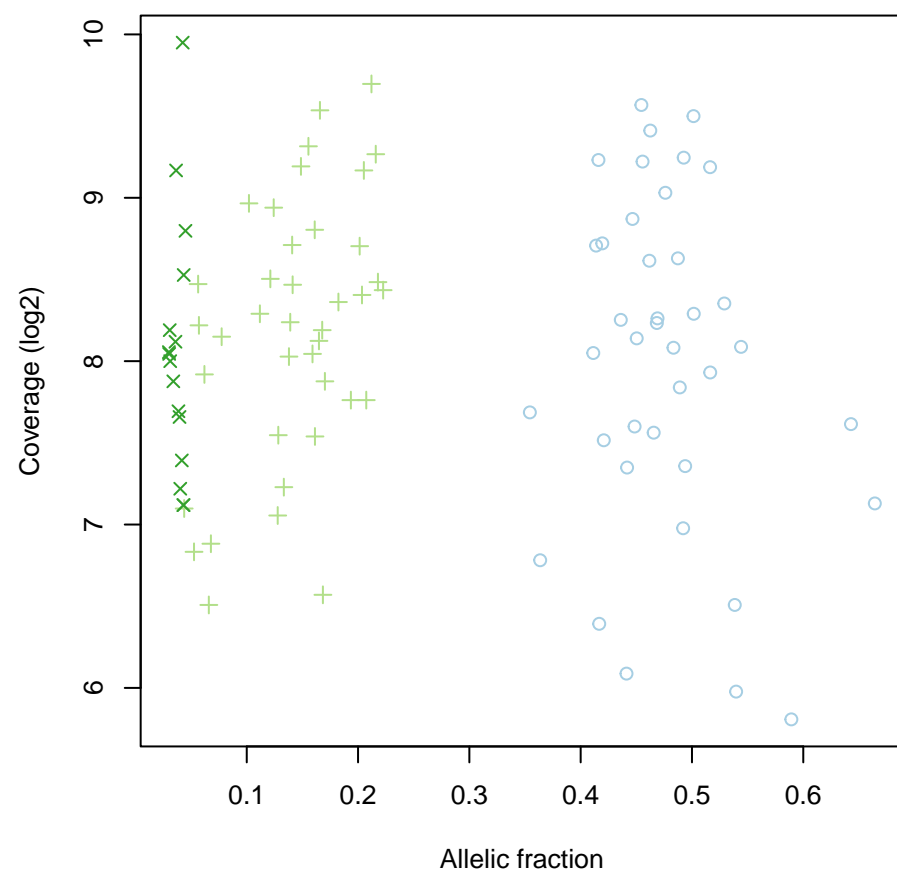
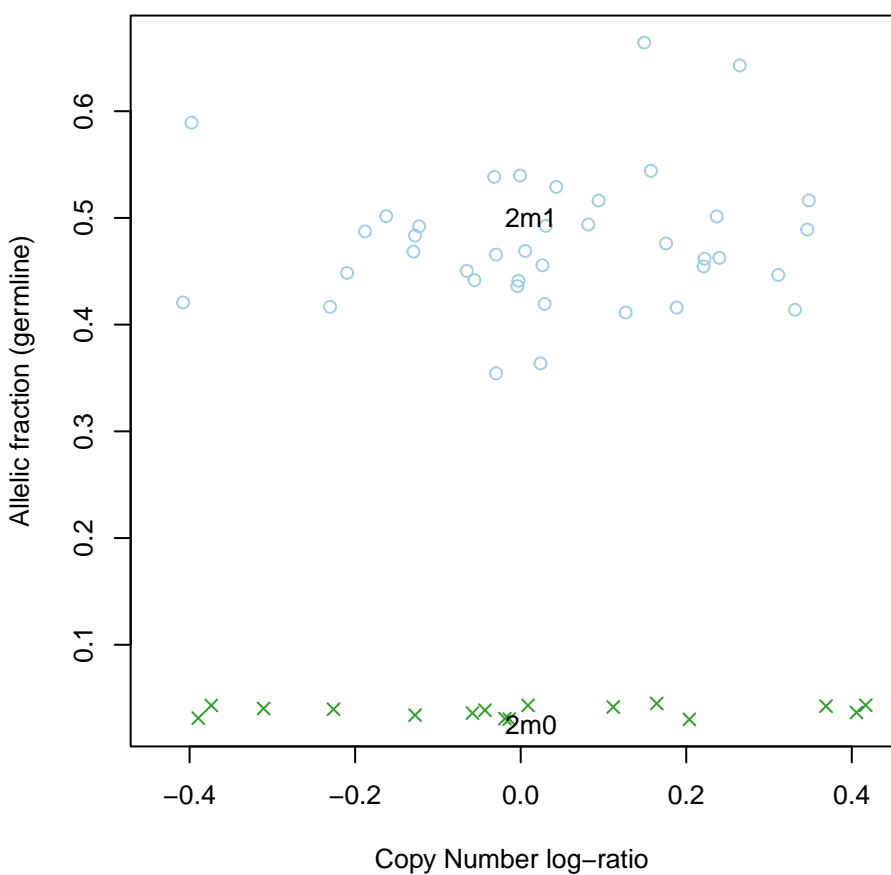


Purity: 0.95 Tumor ploidy: 1.982 SNV log-likelihood: -196.03 GoF: 0% Mean coverage: 315;324

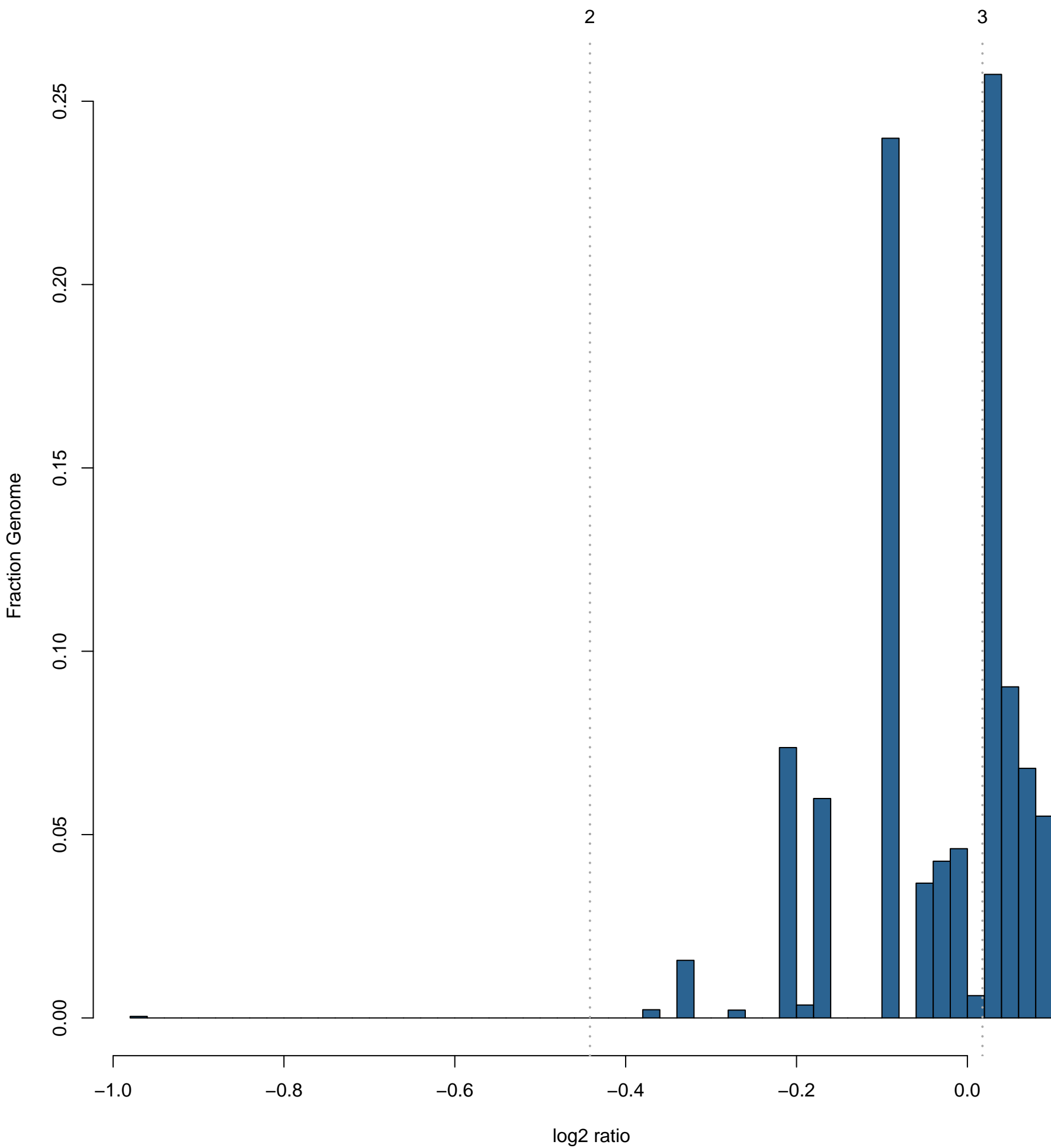


SCNA-fit log-likelihood: -953.92

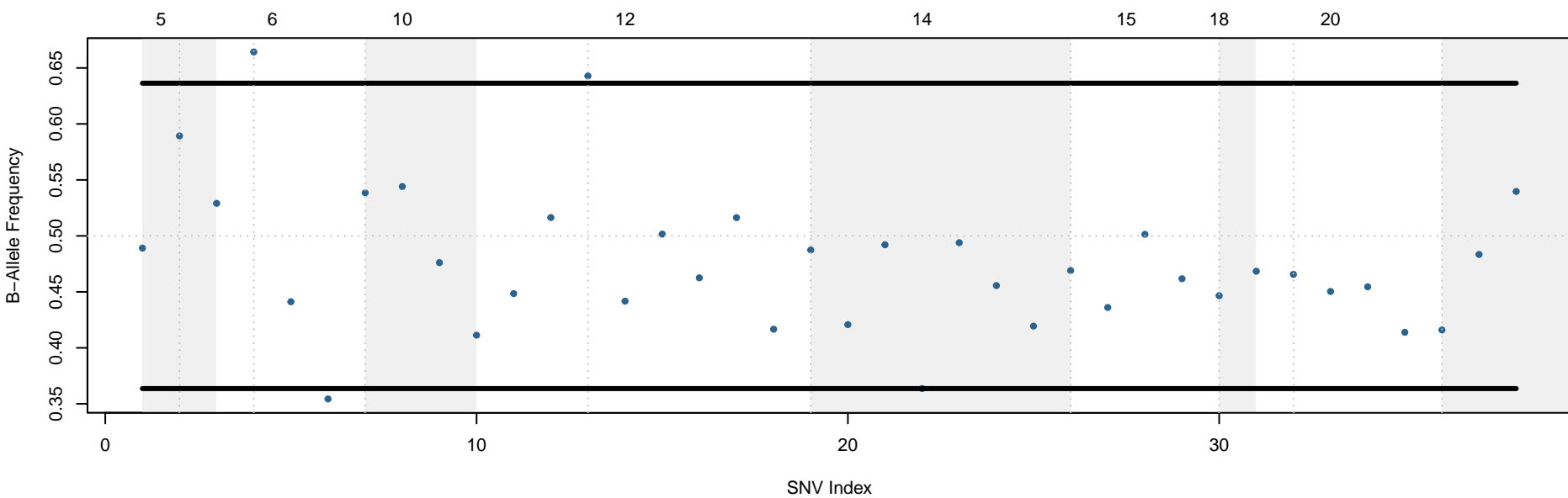




Purity: 0.75 Tumor ploidy: 2.955



Purity: 0.75 Tumor ploidy: 2.955 SNV log-likelihood: -357.18 GoF: 46.3% Mean coverage: 315,324



SCNA-fit log-likelihood: -883.61

