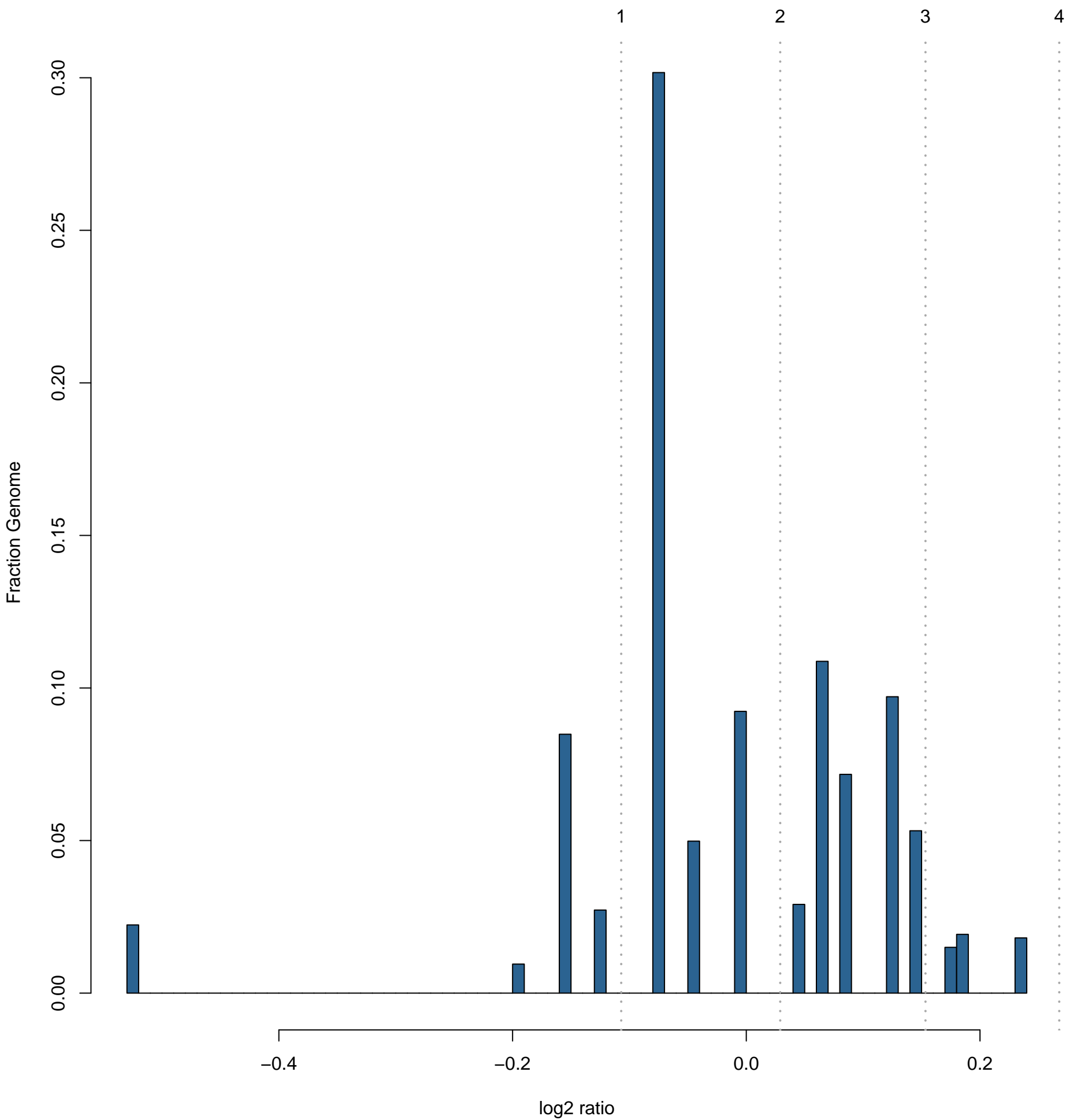
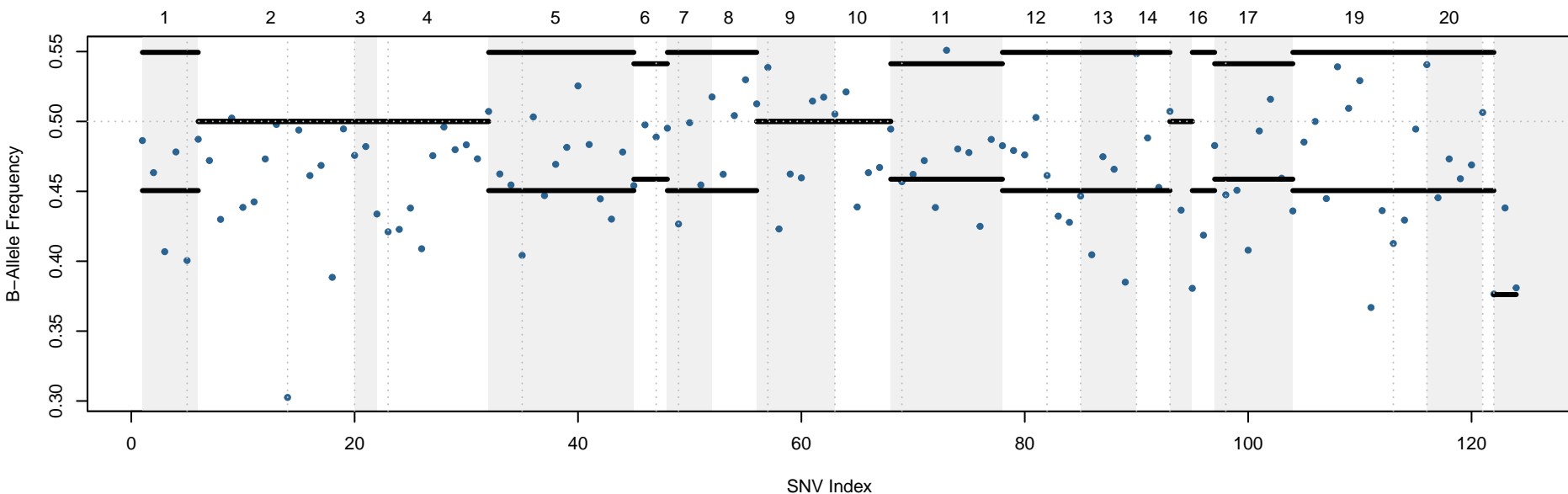


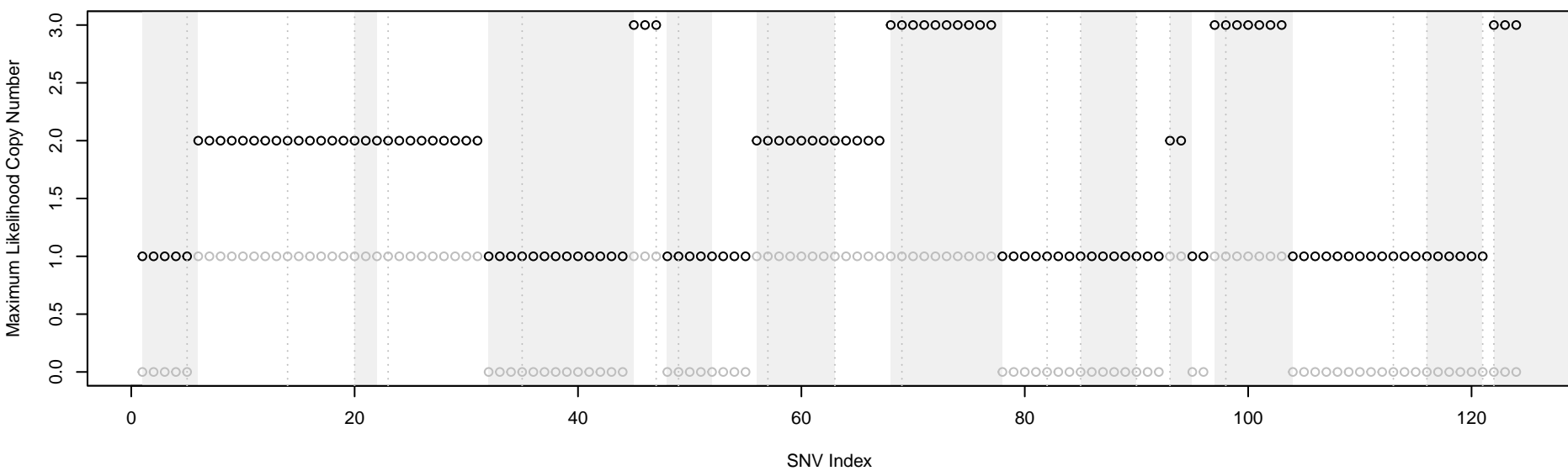
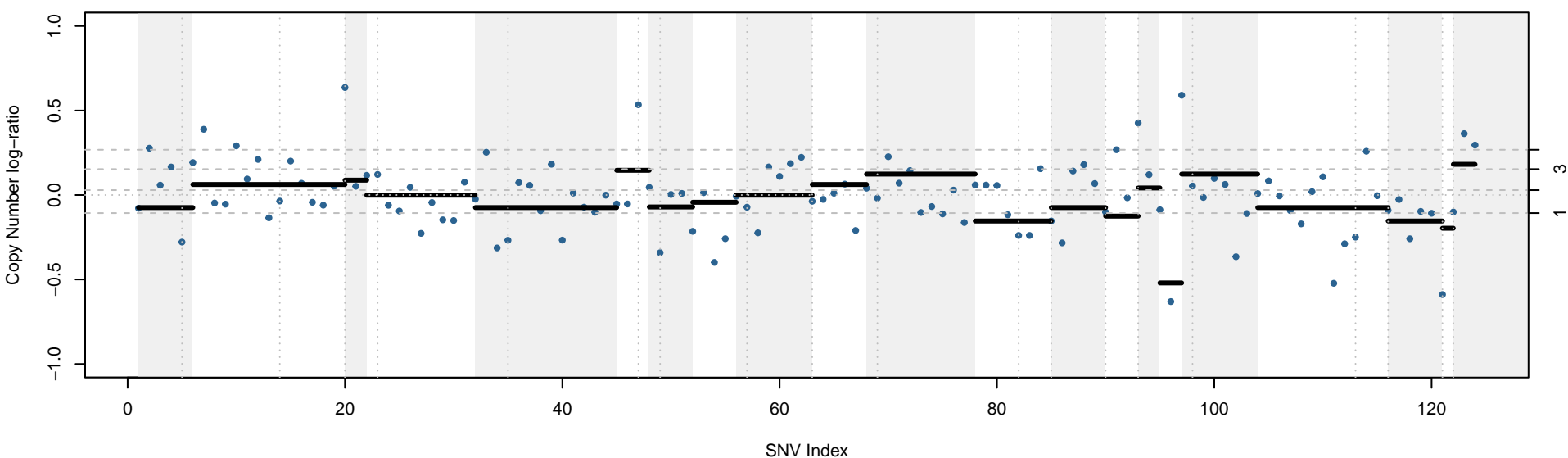
Purity: 0.18 Tumor ploidy: 1.779

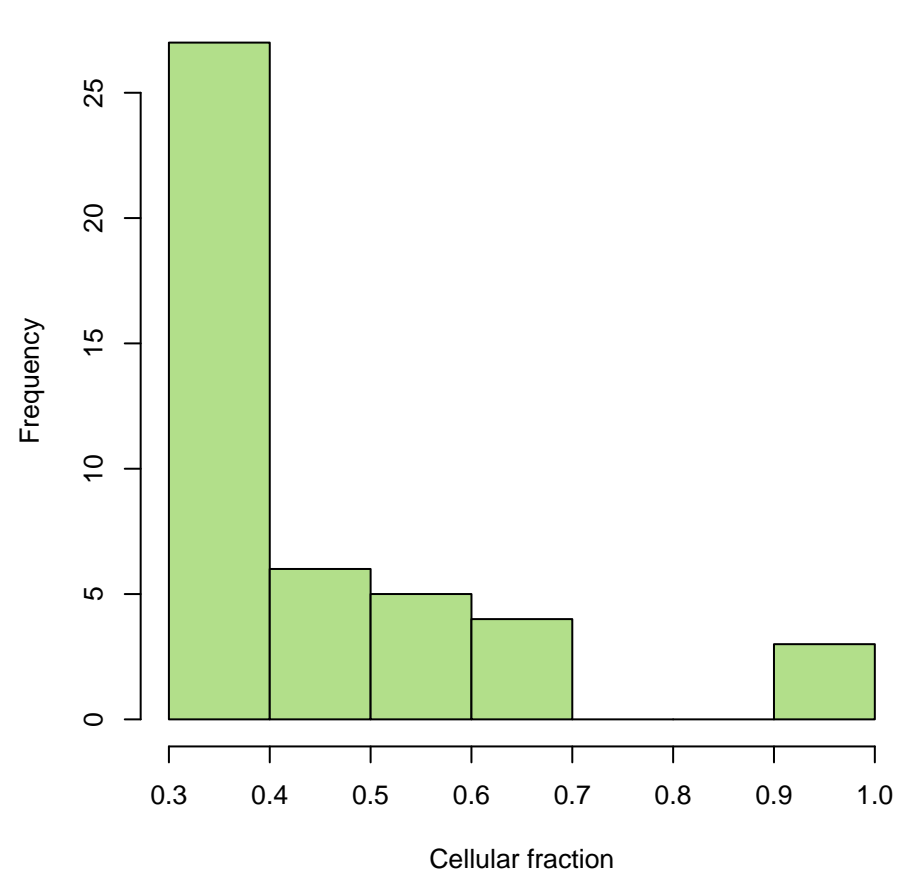
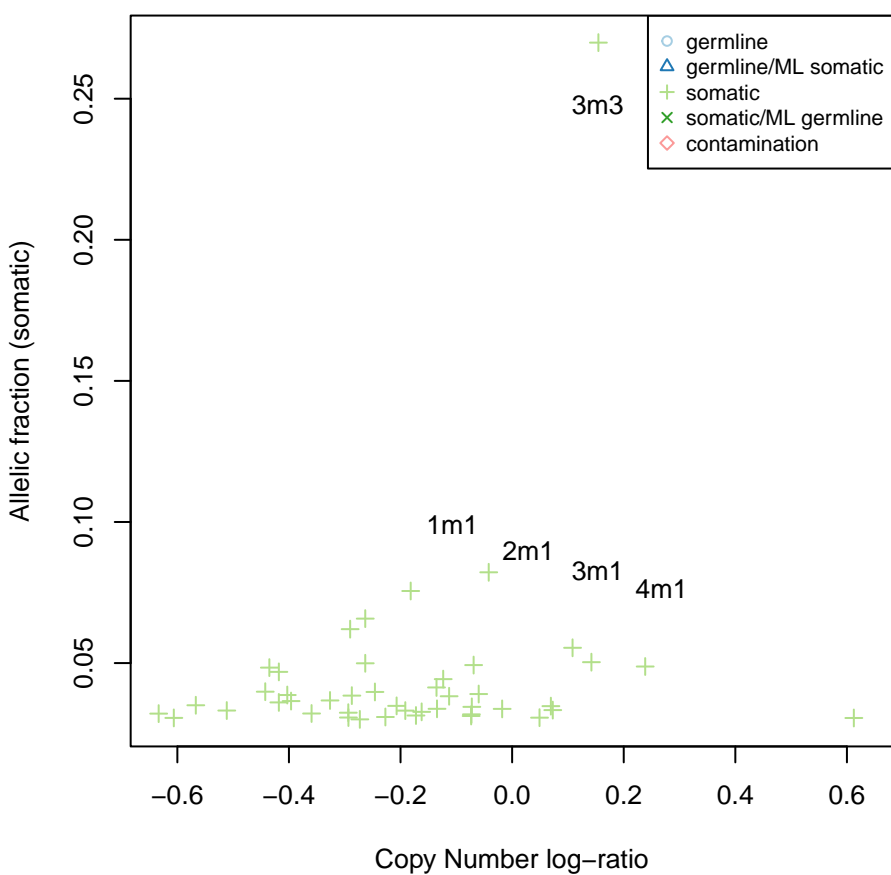
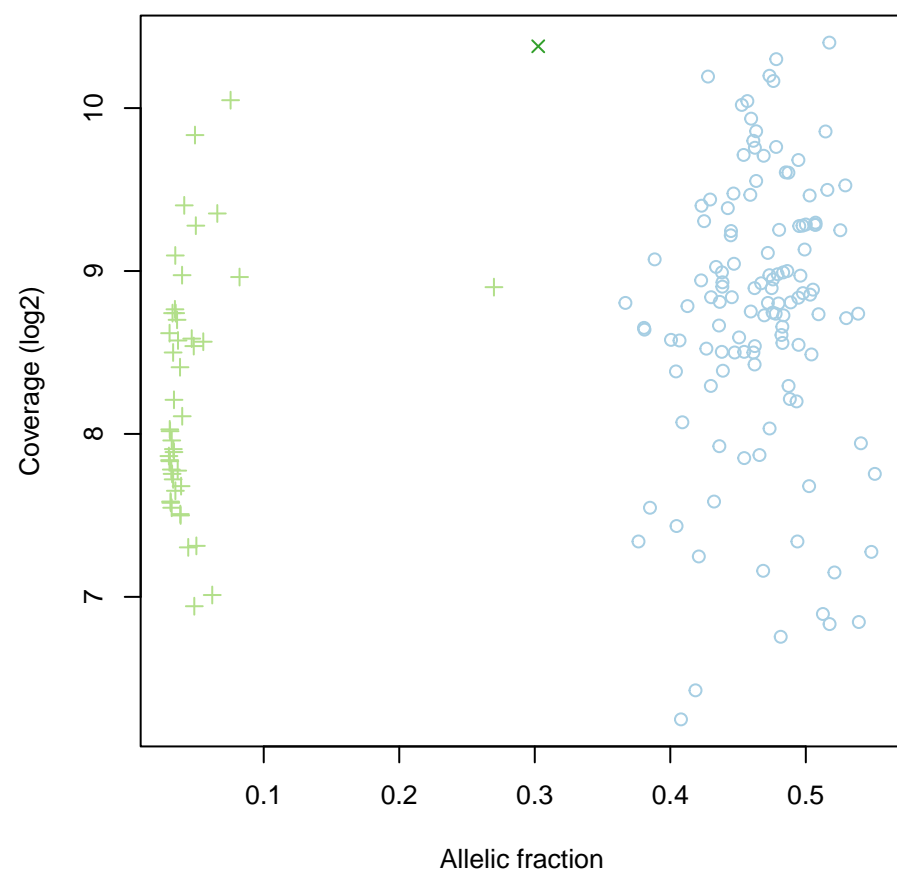
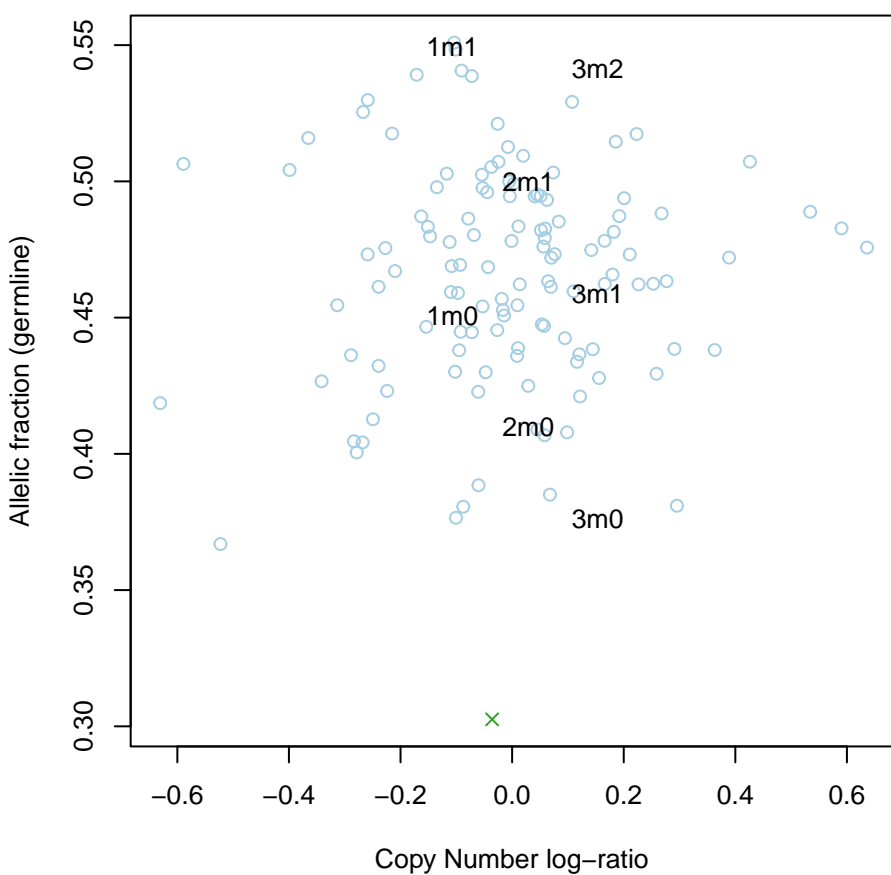


Purity: 0.18 Tumor ploidy: 1.779 SNV log-likelihood: 122.92 GoF: 95.9% Mean coverage: 464;227

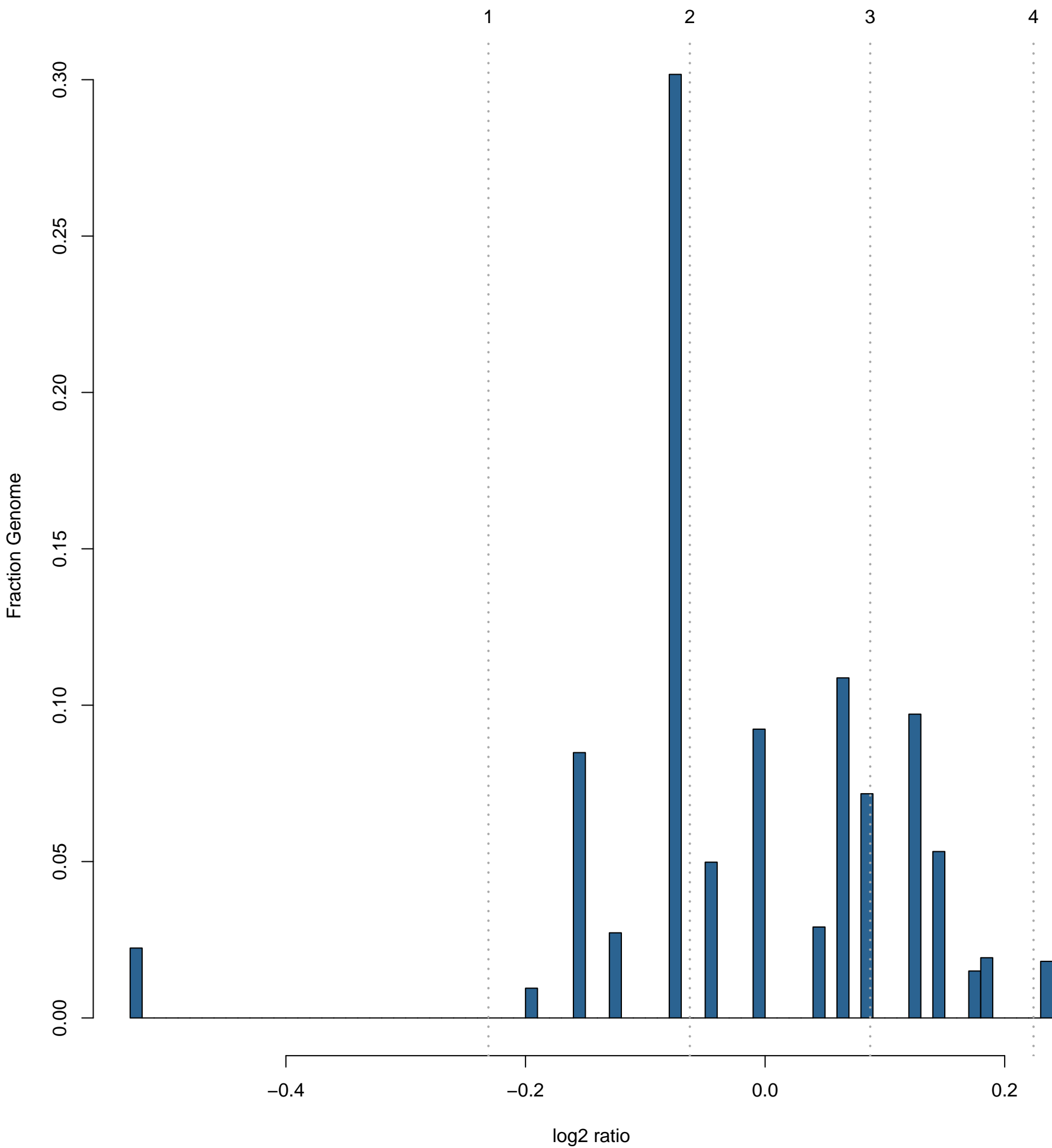


SCNA-fit log-likelihood: -12552.84

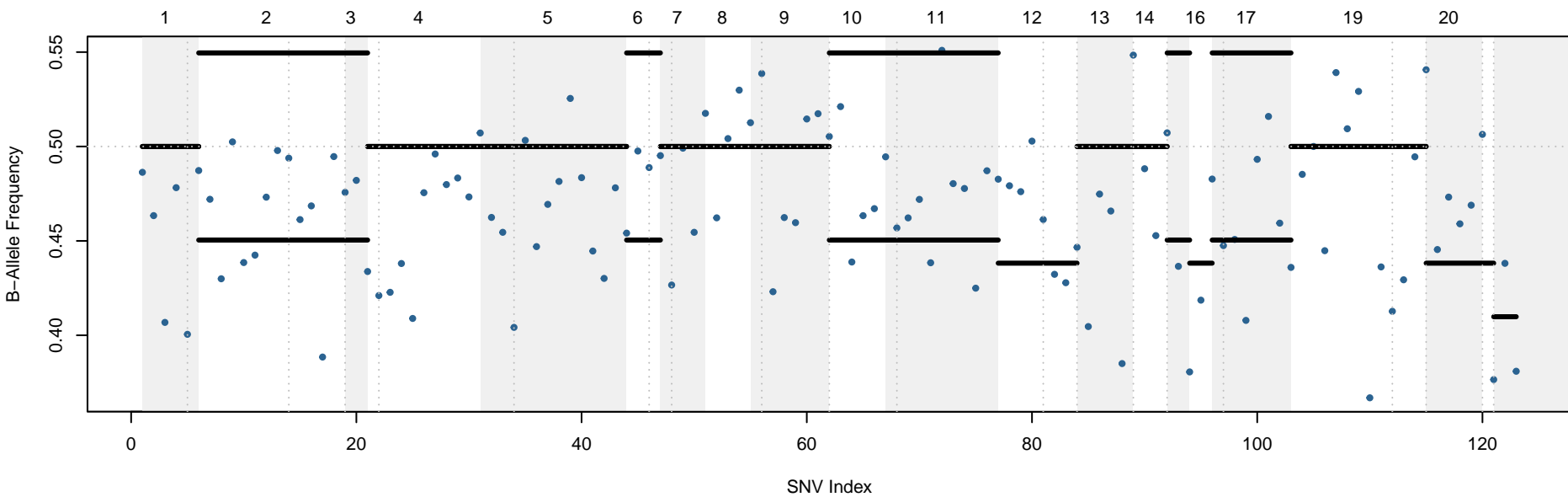




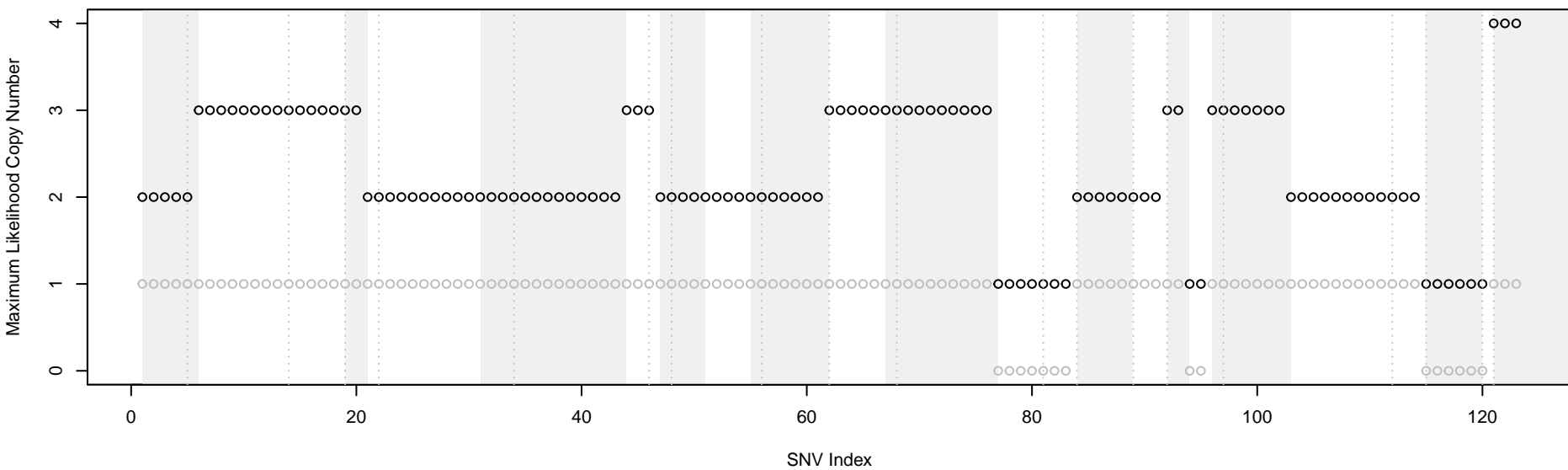
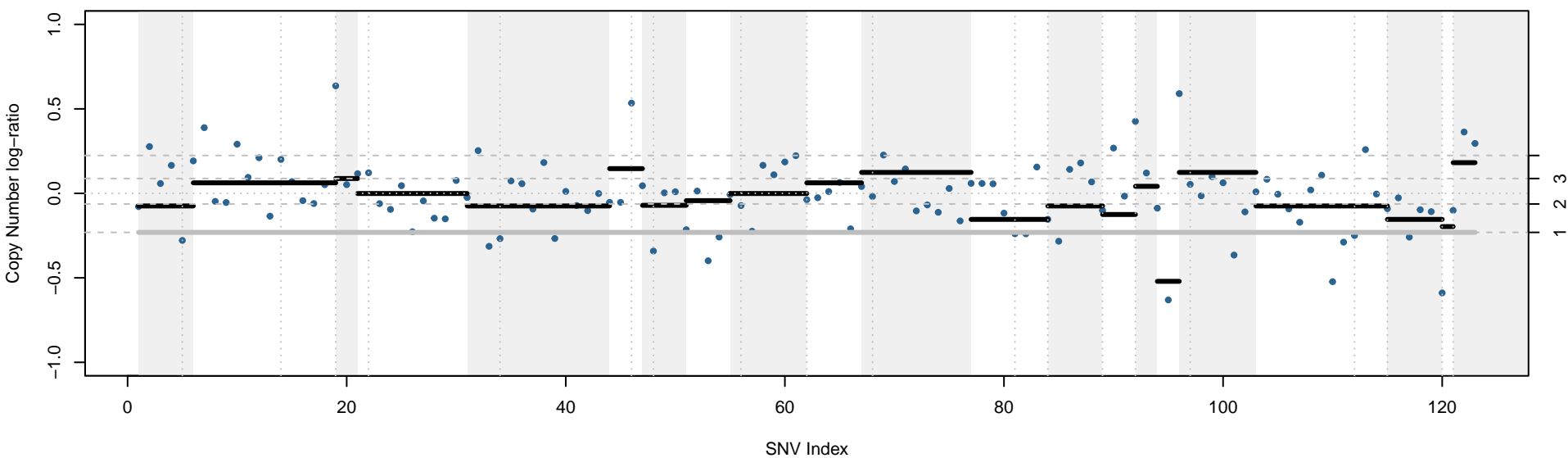
Purity: 0.22 Tumor ploidy: 2.405

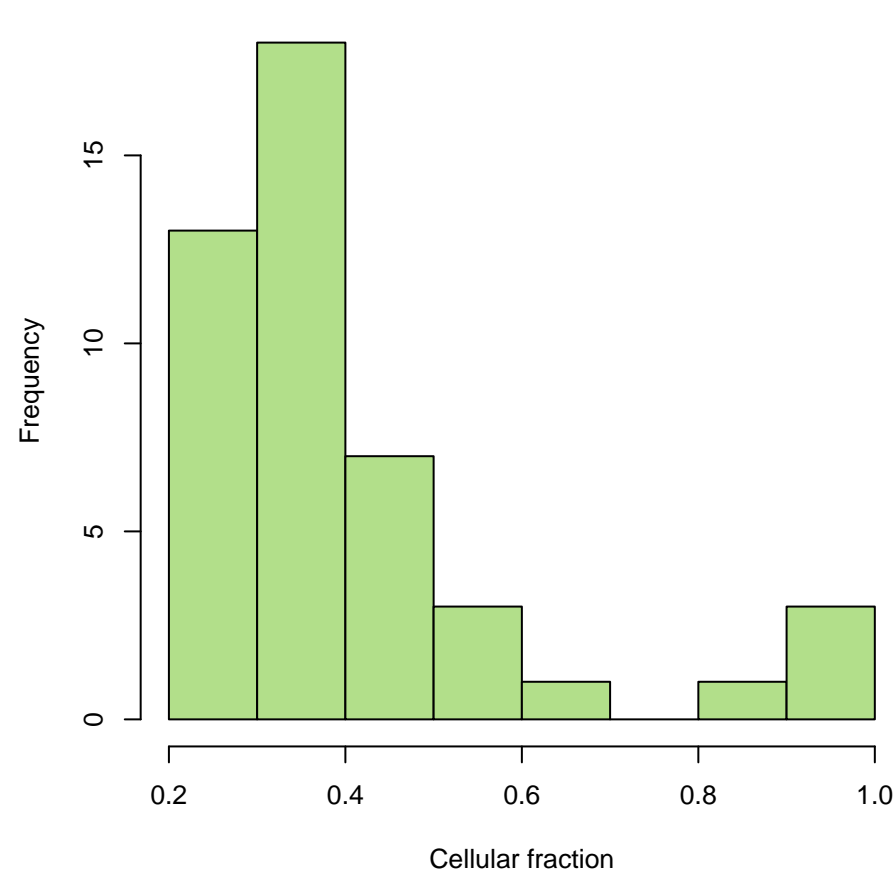
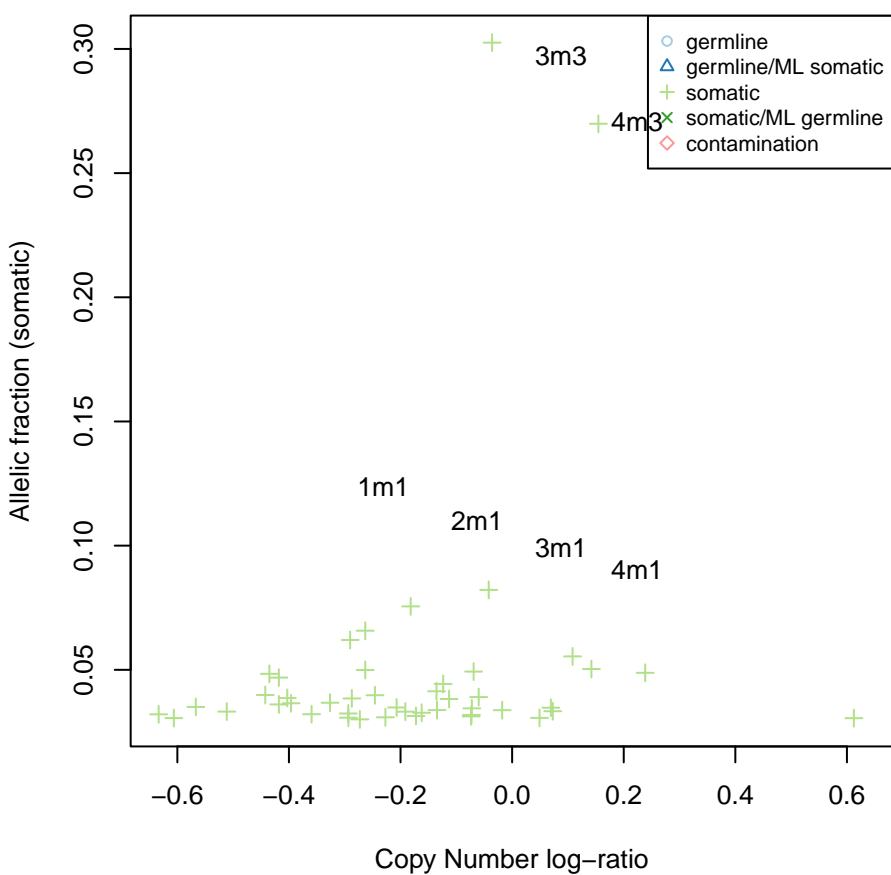
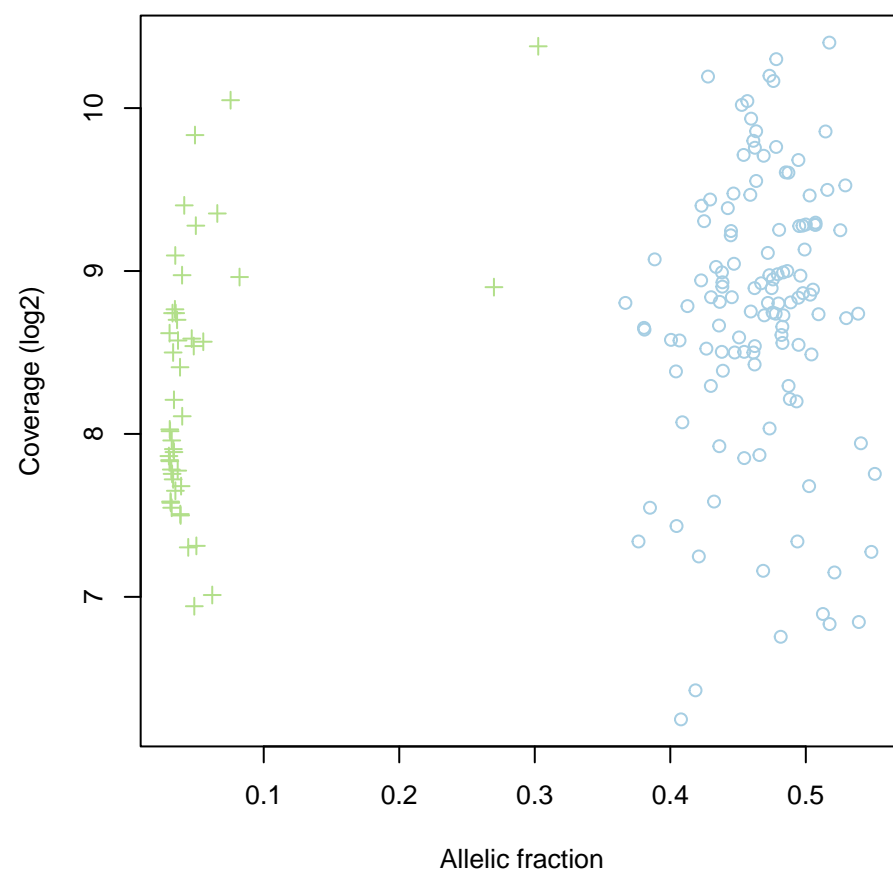
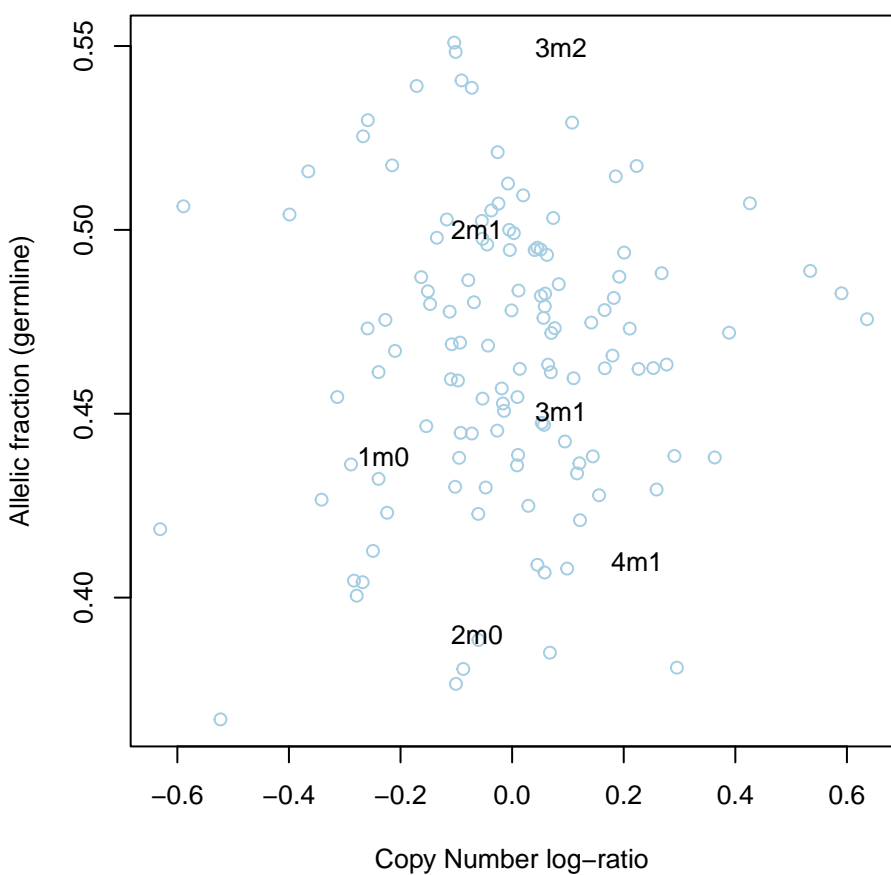


Purity: 0.22 Tumor ploidy: 2.405 SNV log-likelihood: 66.41 GoF: 94.1% Mean coverage: 464;227

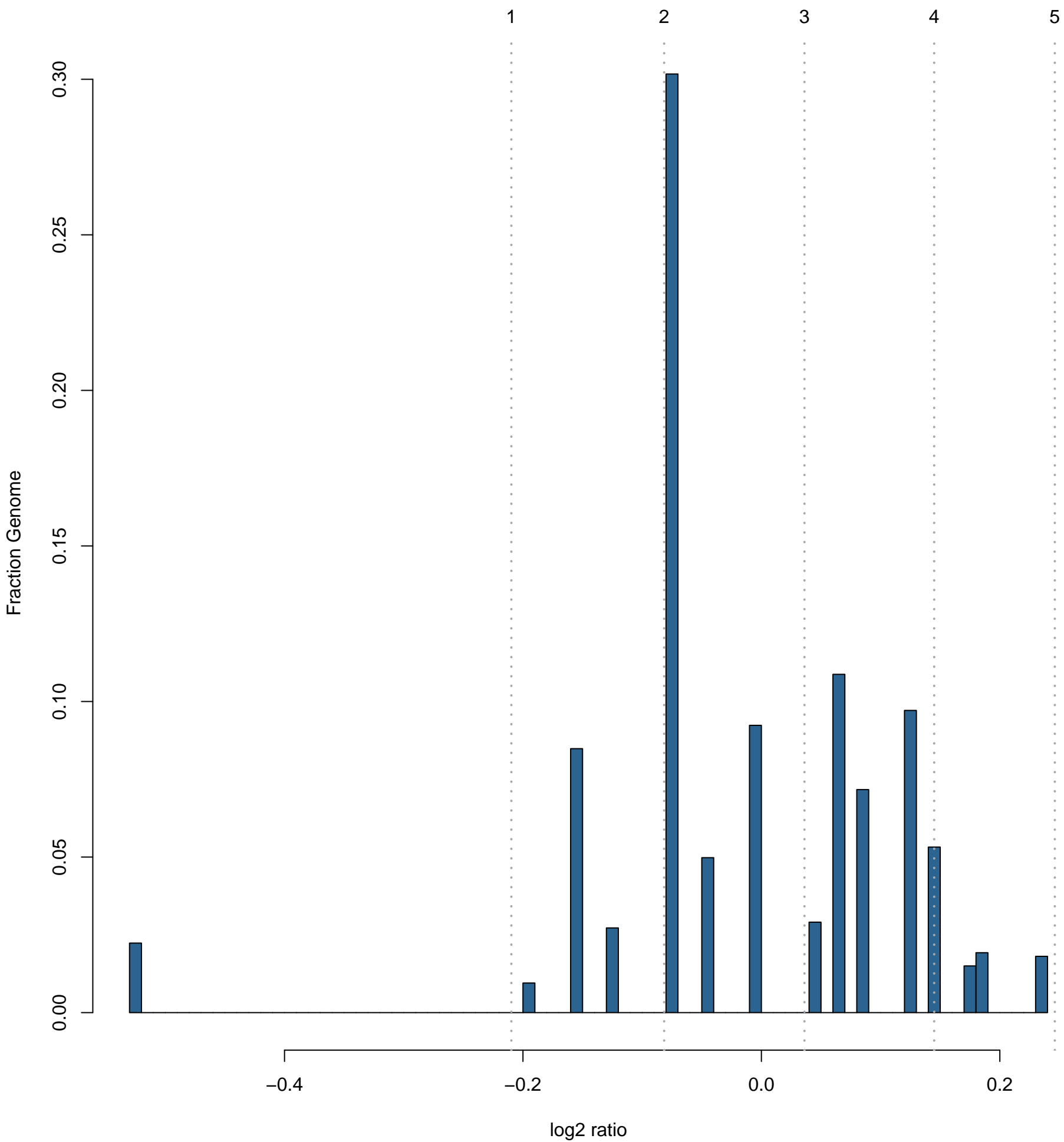


SCNA-fit log-likelihood: -12524.75

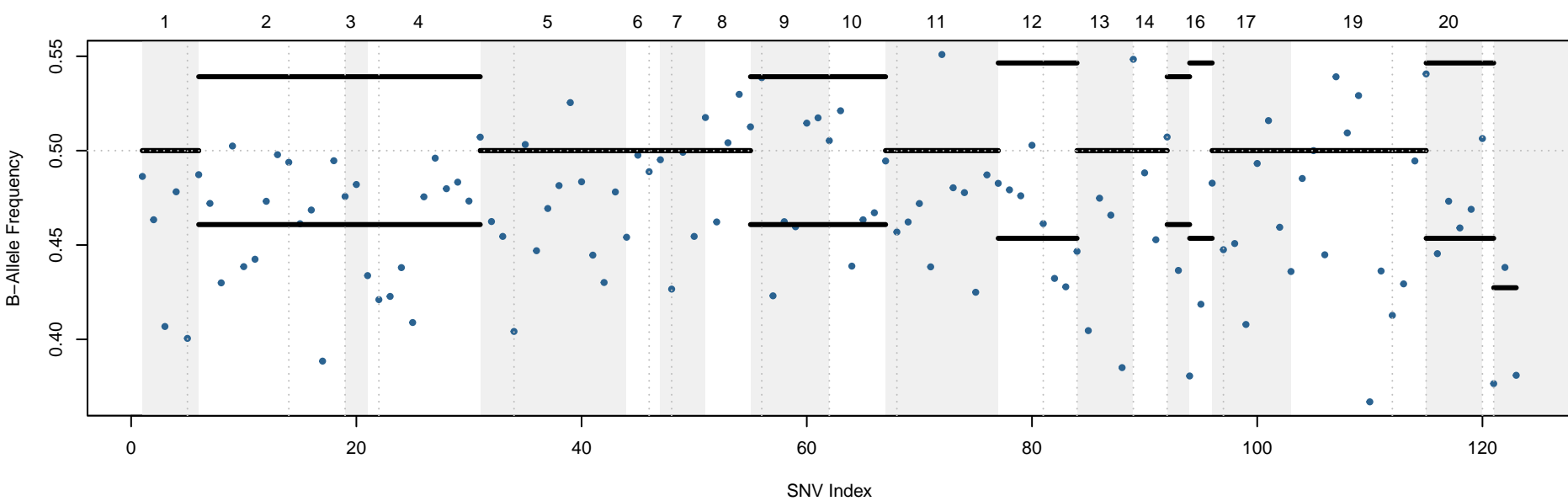




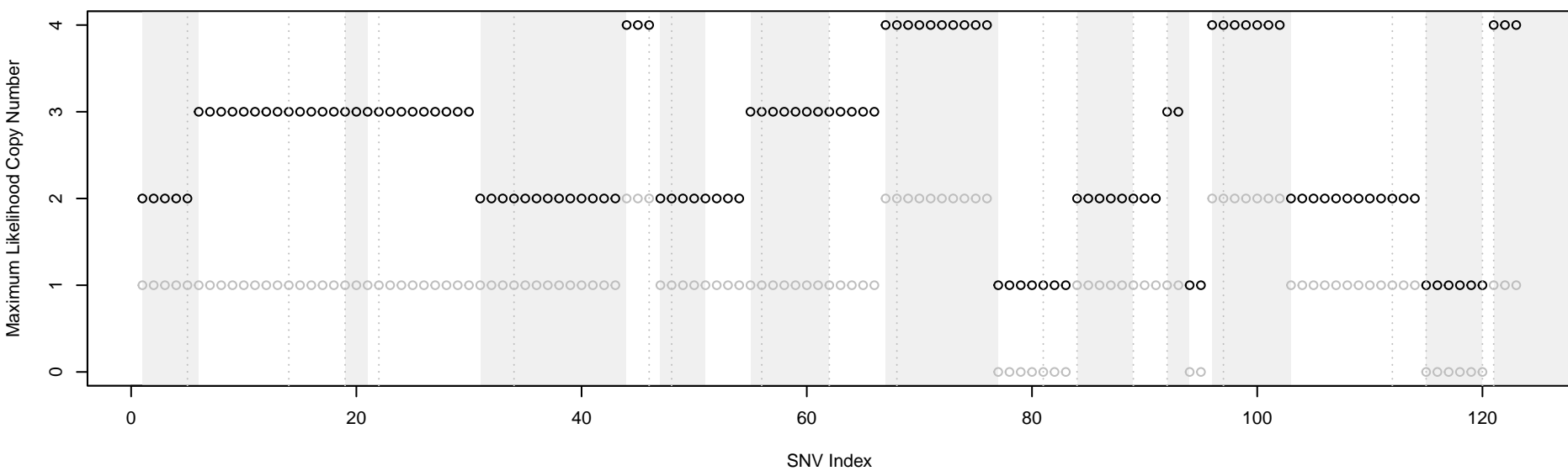
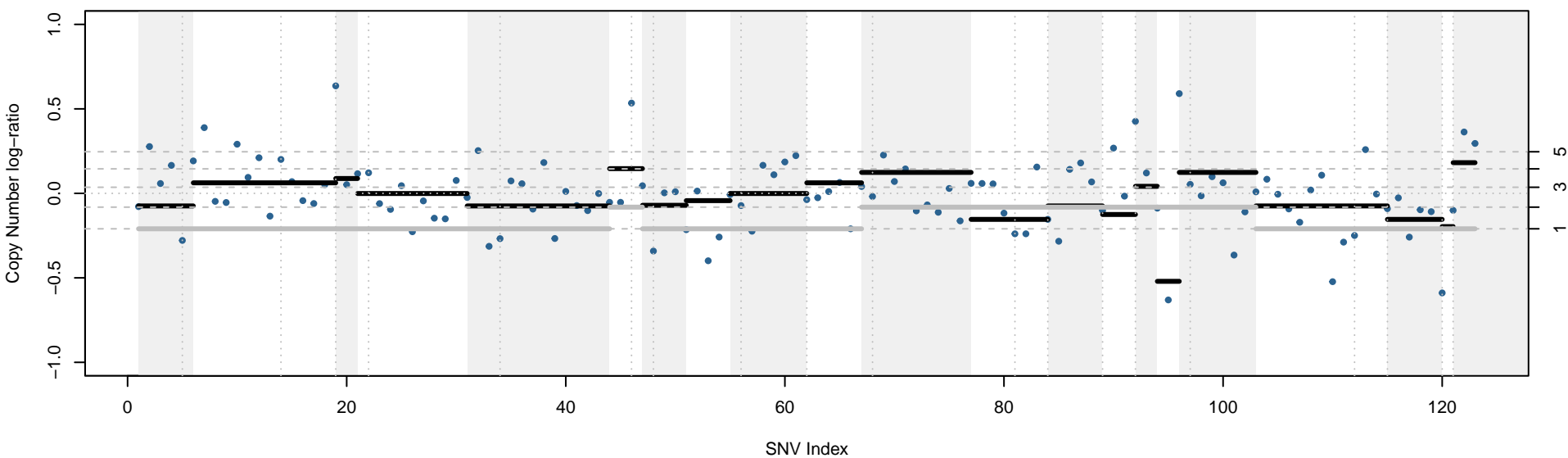
Purity: 0.17 Tumor ploidy: 2.684



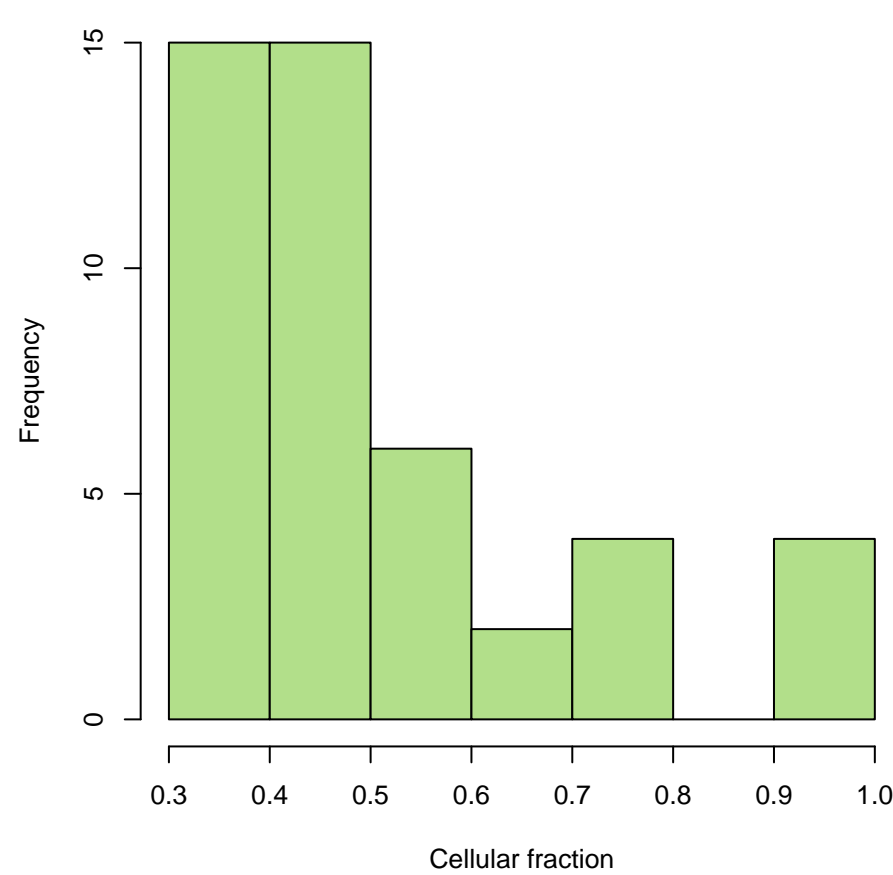
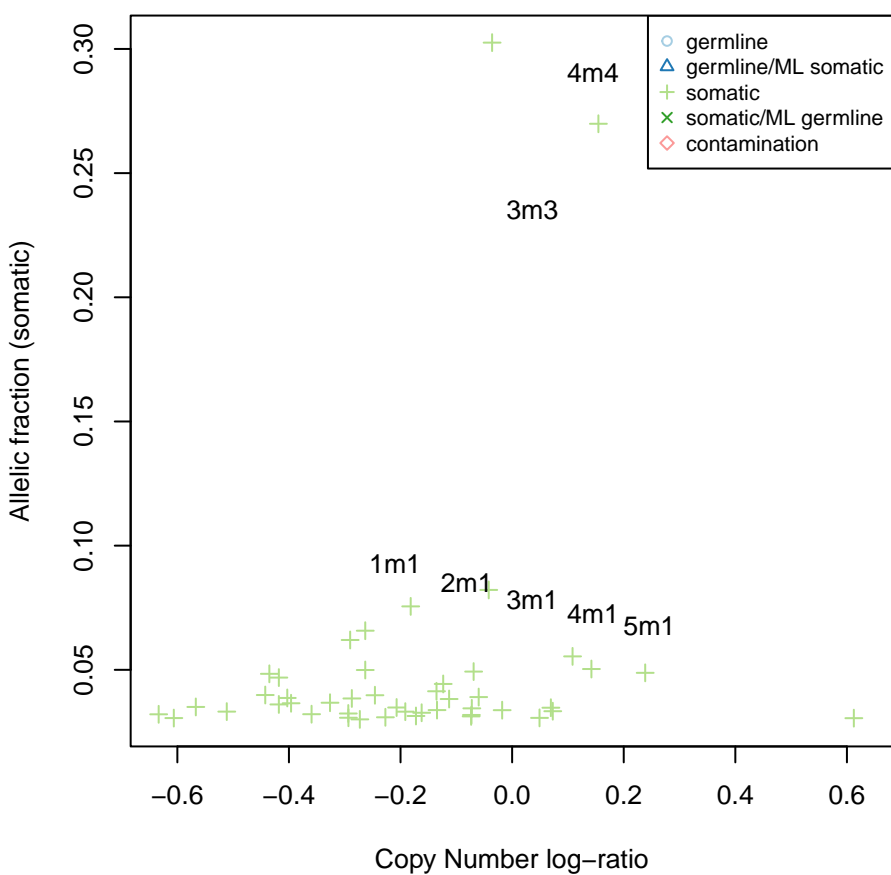
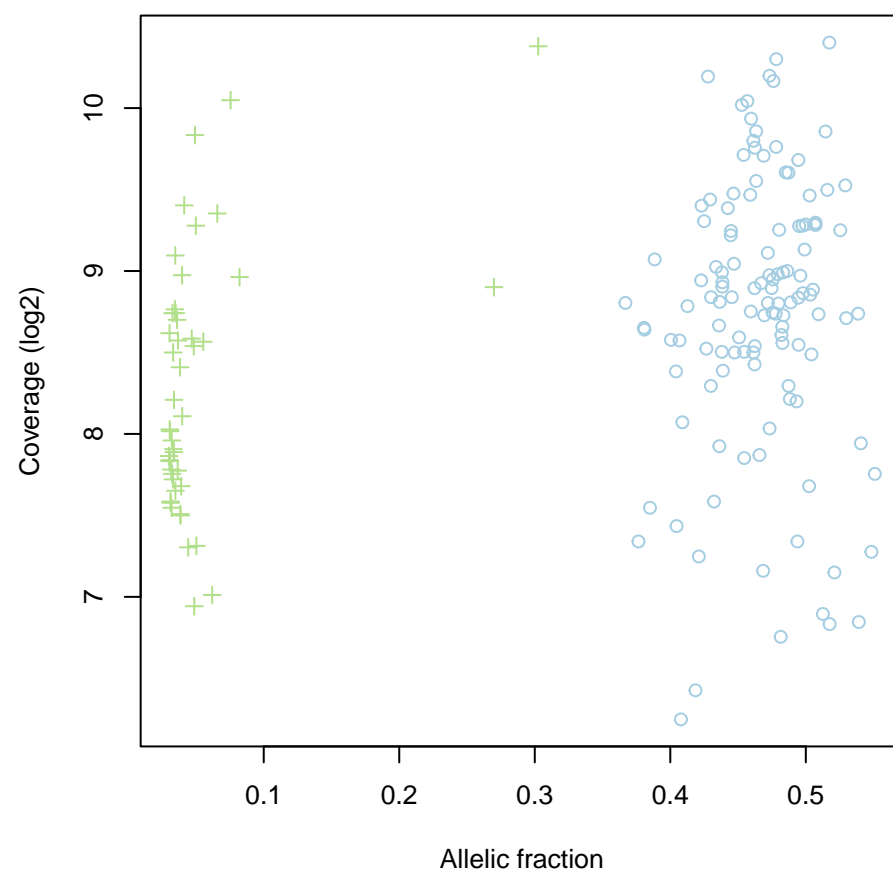
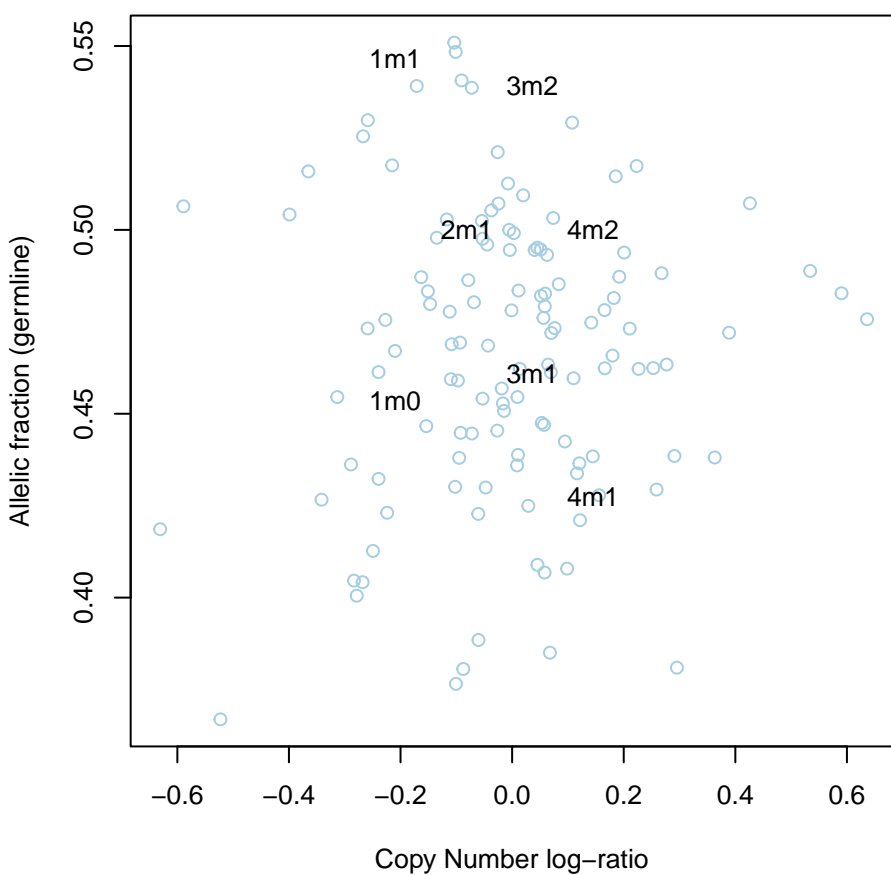
Purity: 0.17 Tumor ploidy: 2.684 SNV log-likelihood: 71.42 GoF: 96.1% Mean coverage: 464;227



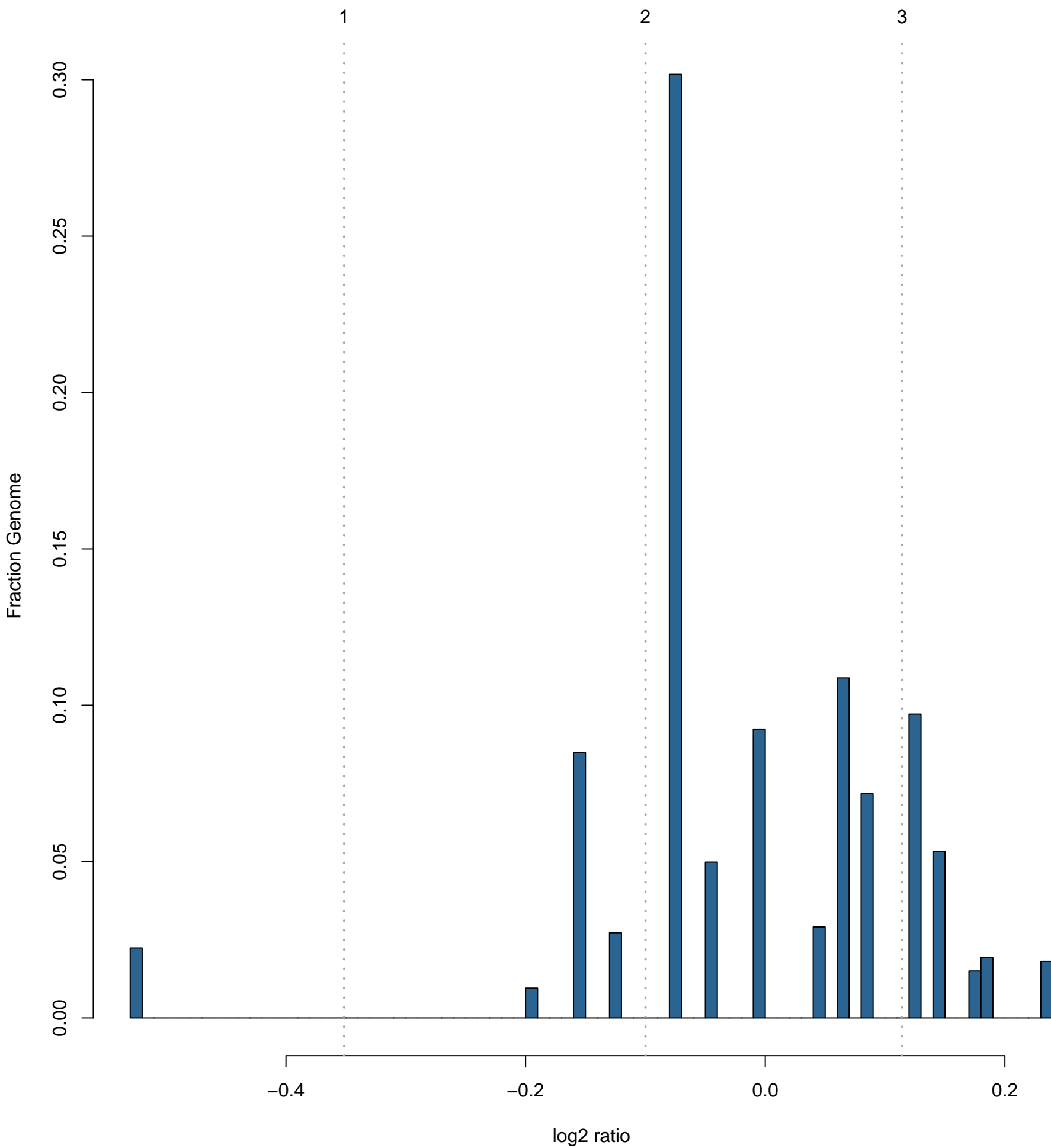
SCNA-fit log-likelihood: -12518.69



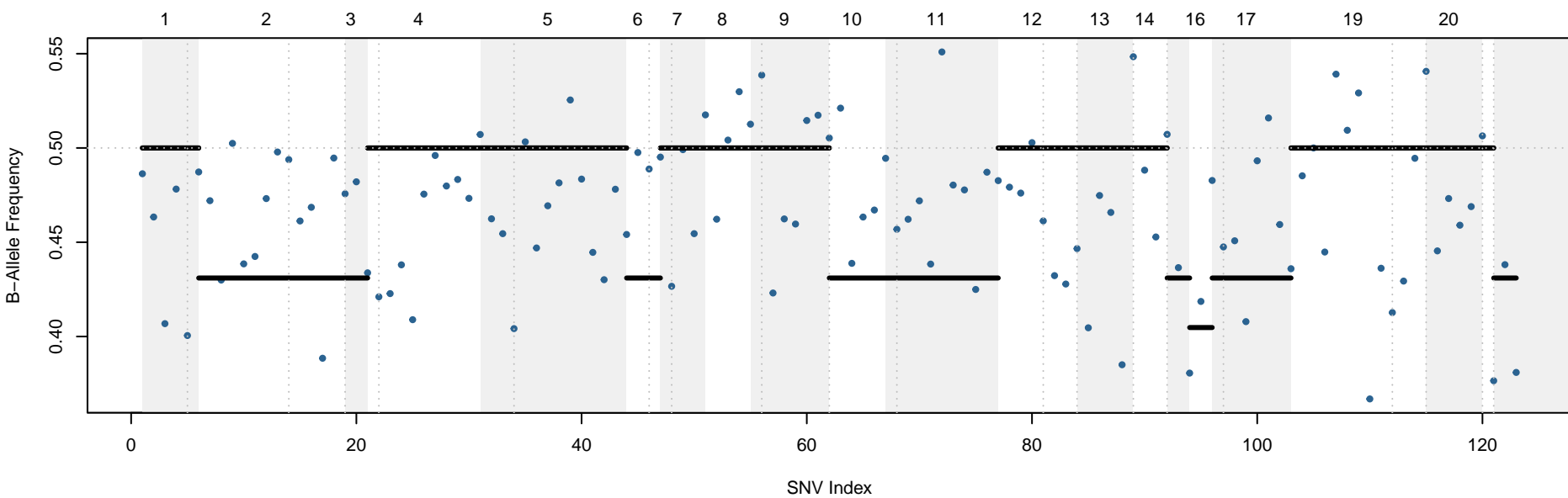




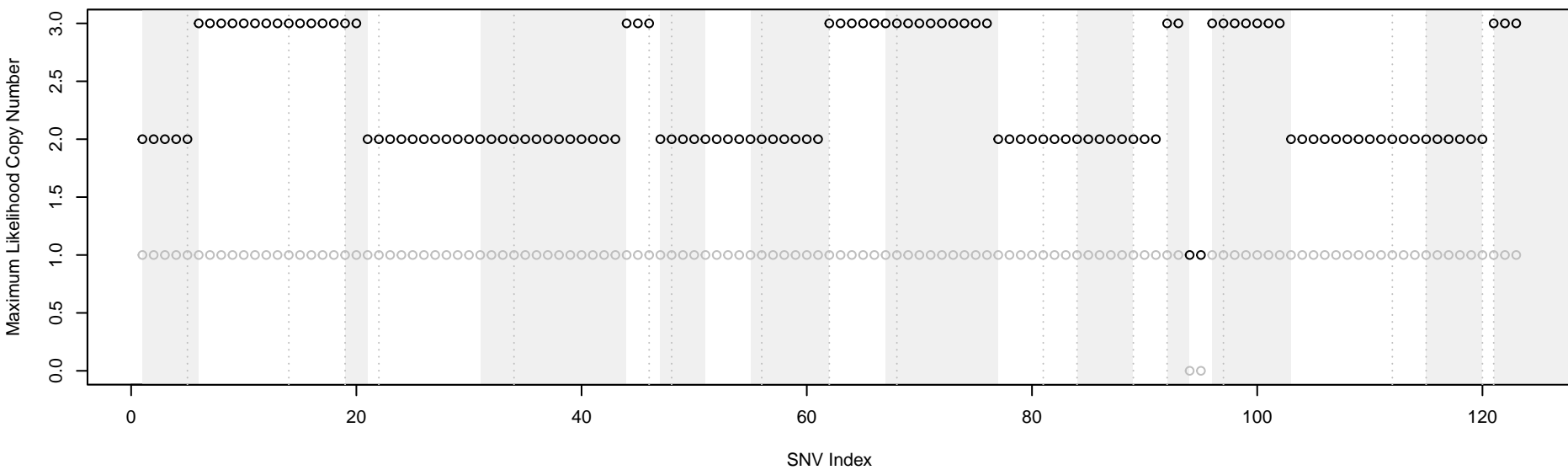
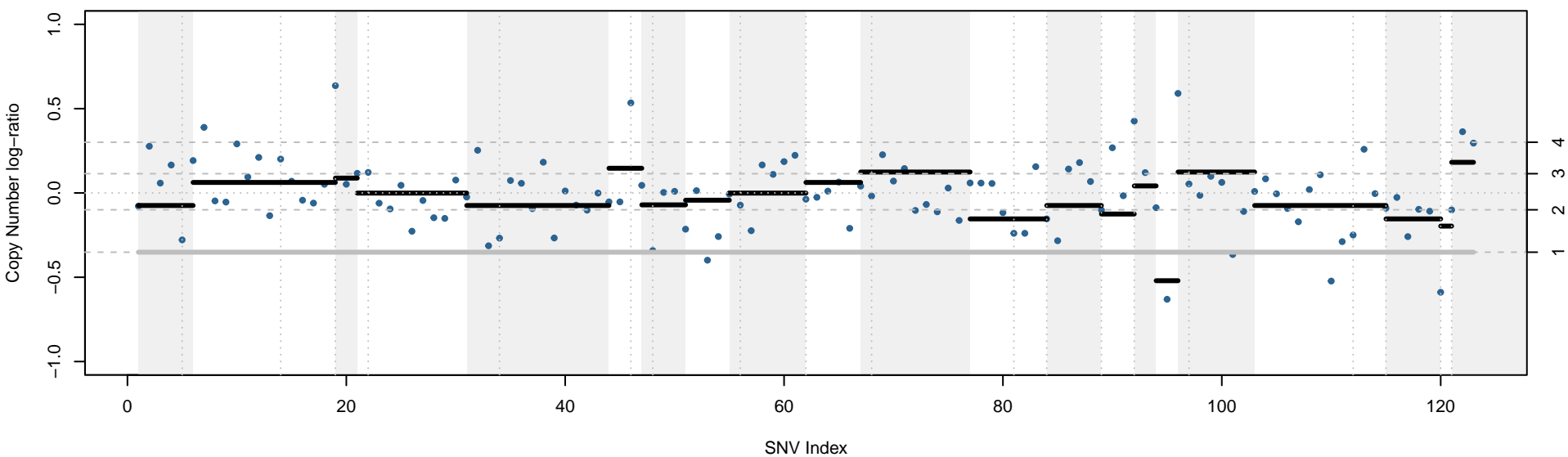
Purity: 0.32 Tumor ploidy: 2.448

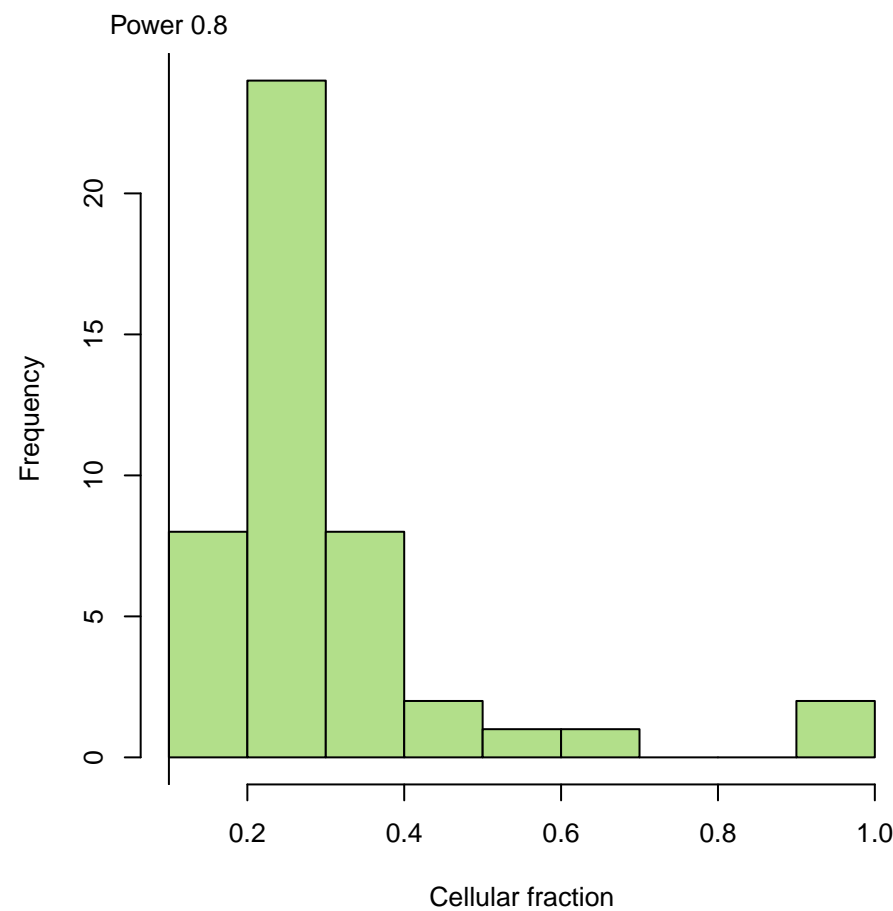
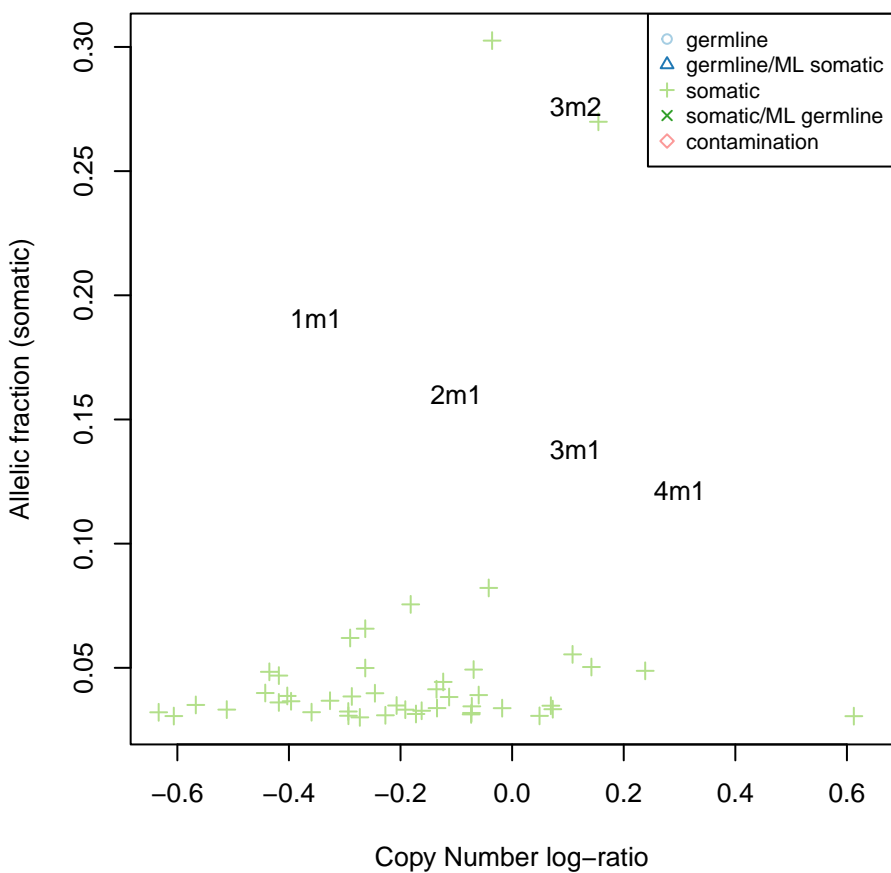
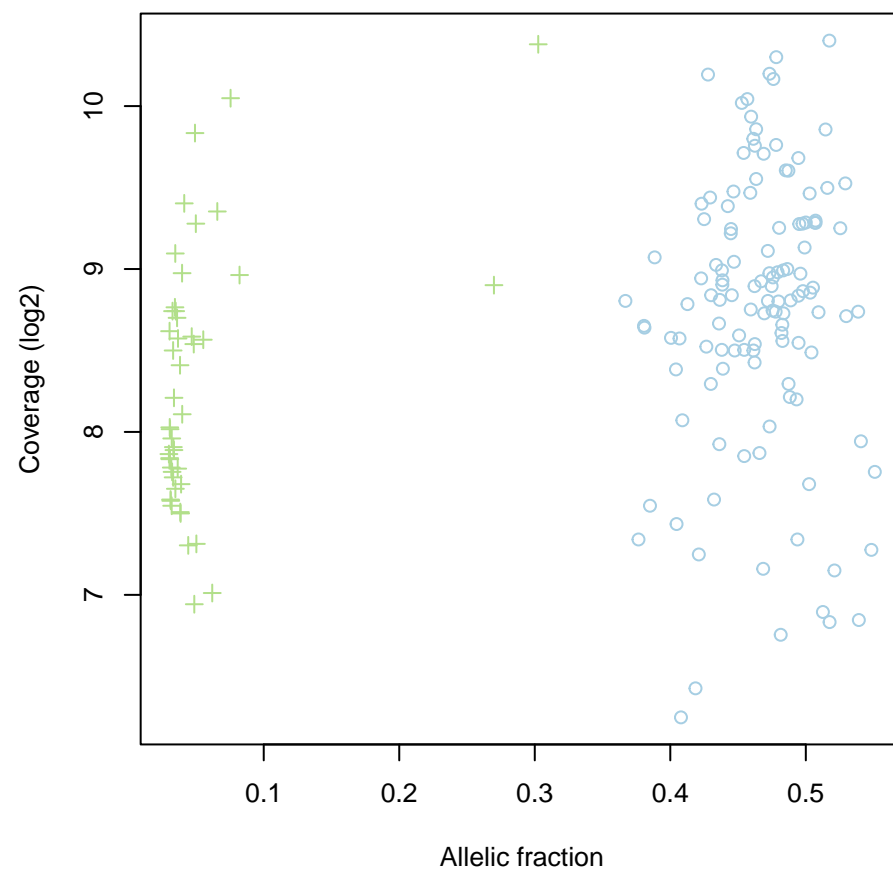
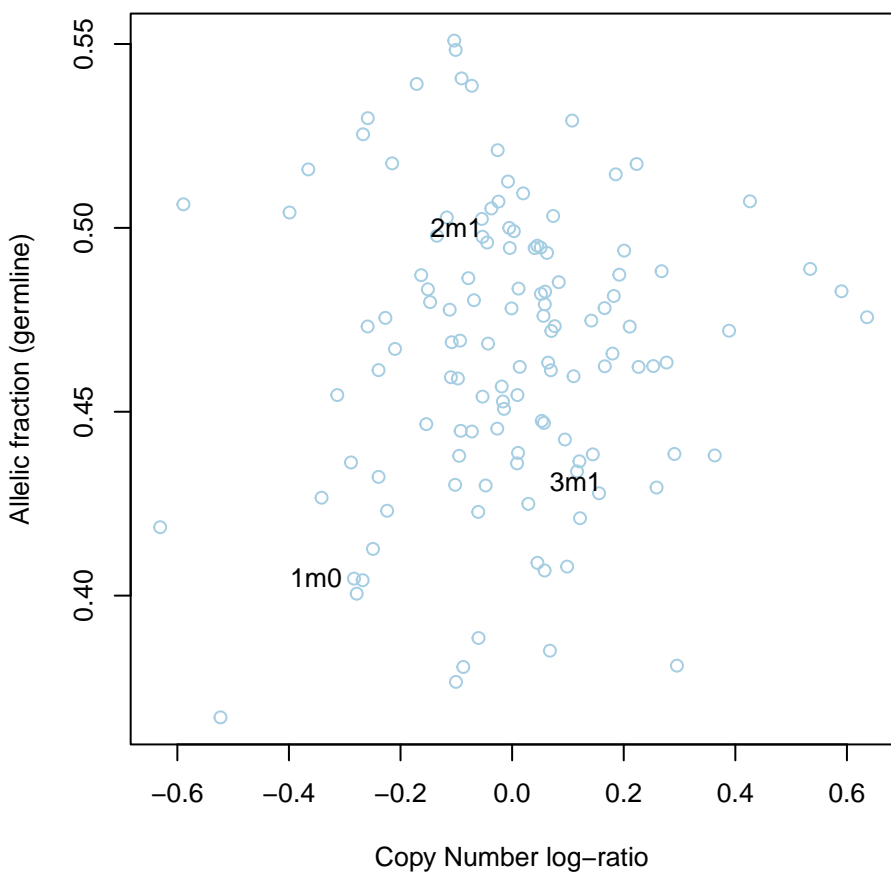


Purity: 0.32 Tumor ploidy: 2.448 SNV log-likelihood: 22.04 GoF: 88.3% Mean coverage: 464;227

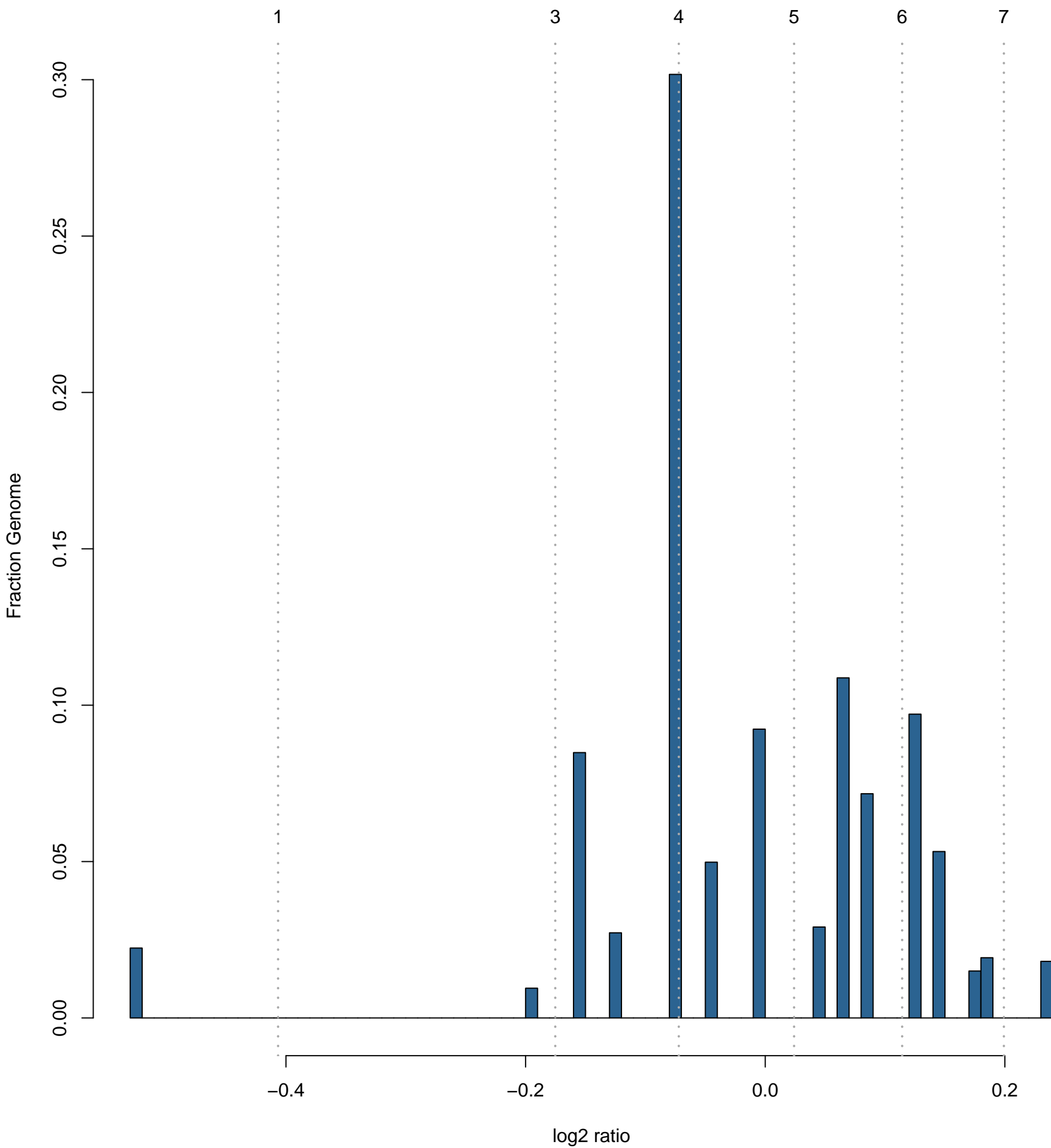


SCNA-fit log-likelihood: -12519.94

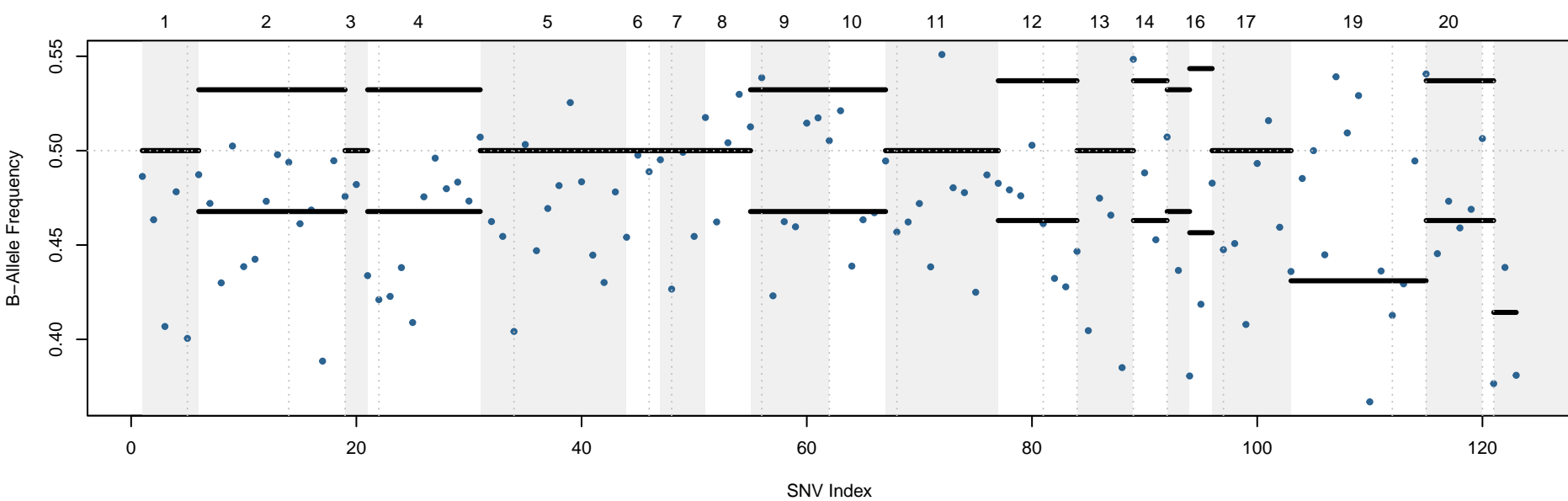




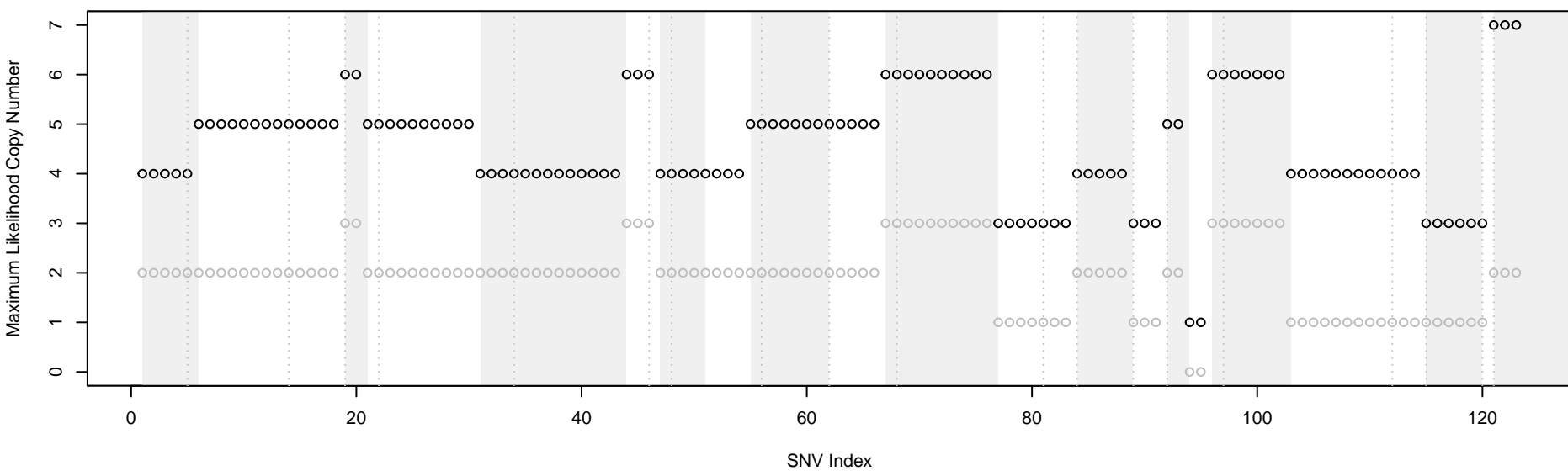
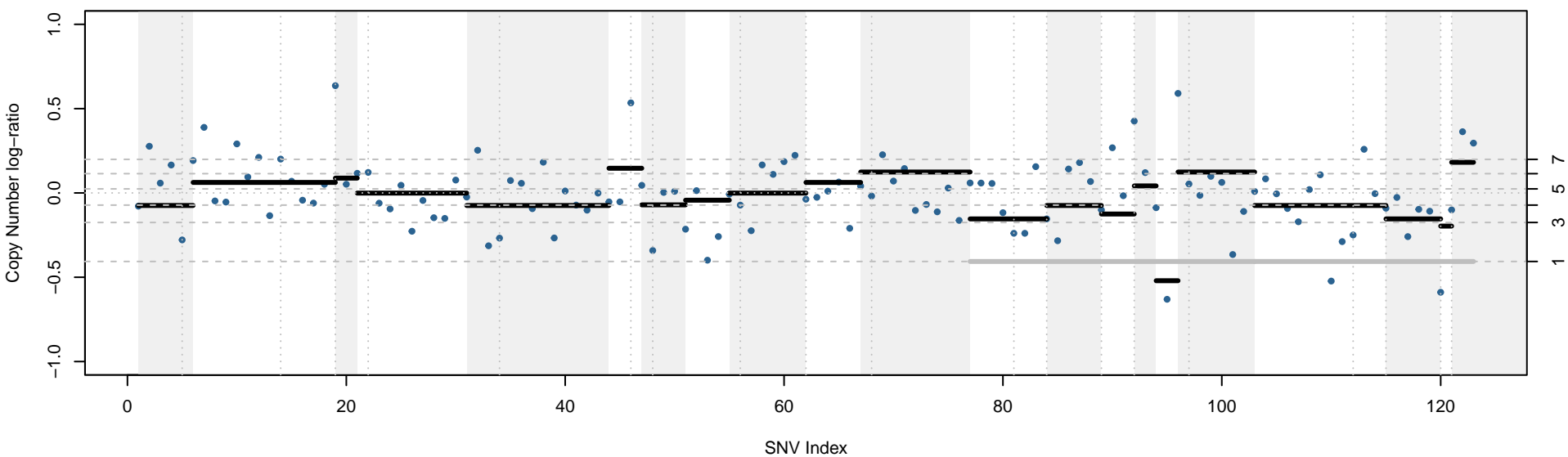
Purity: 0.16 Tumor ploidy: 4.743

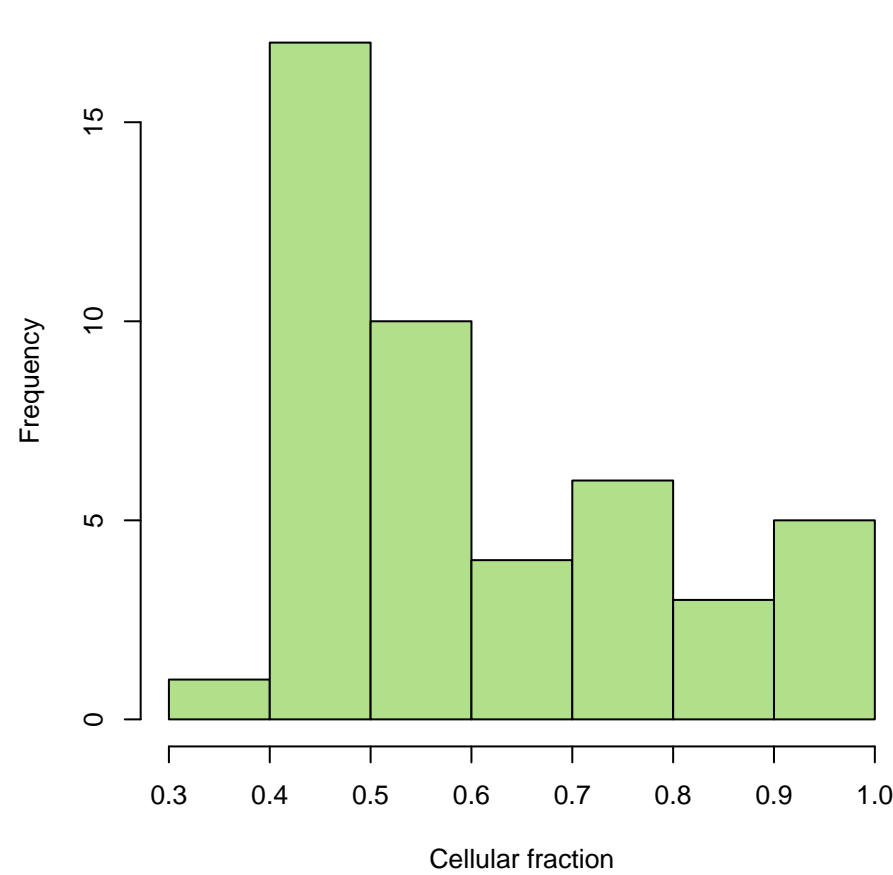
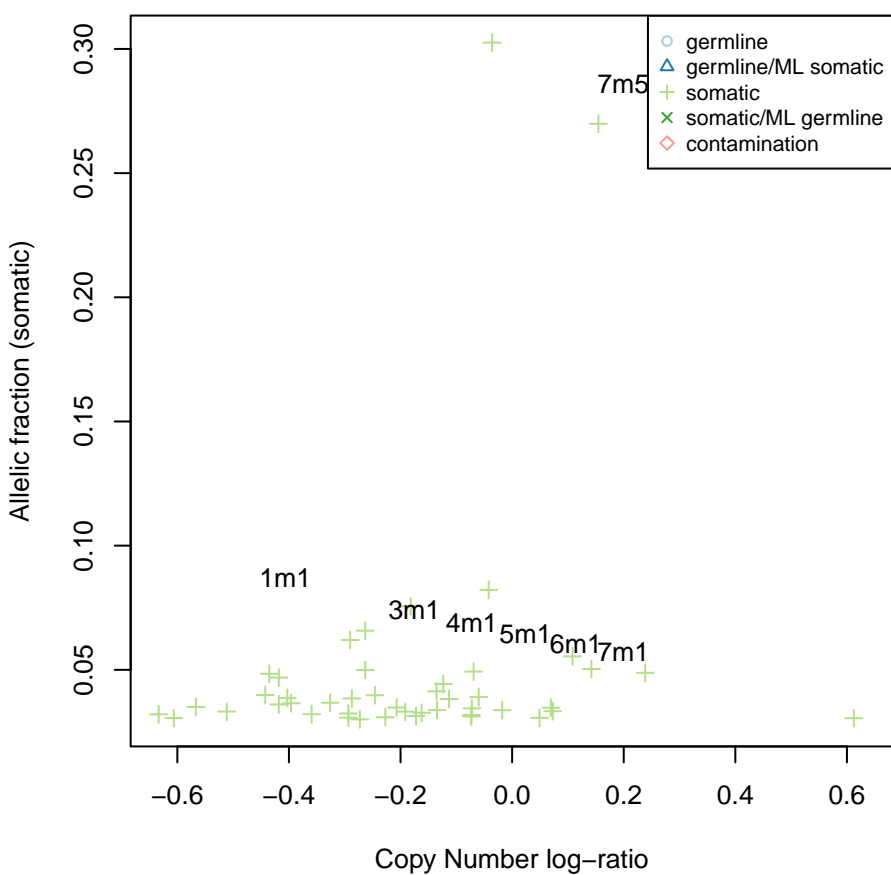
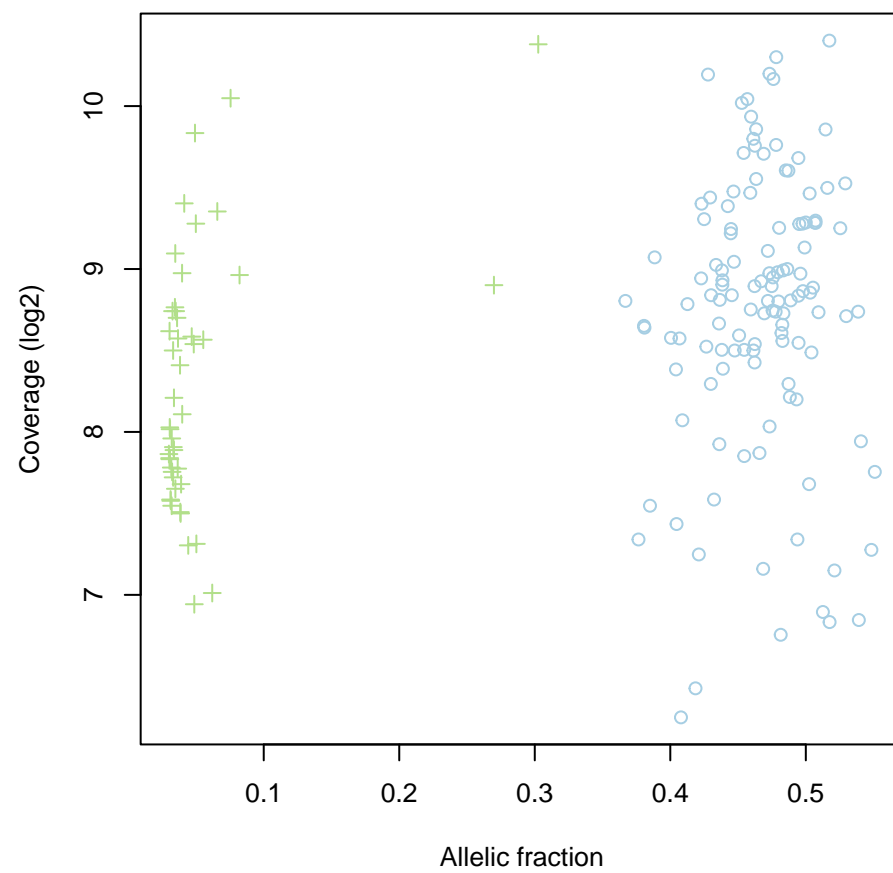
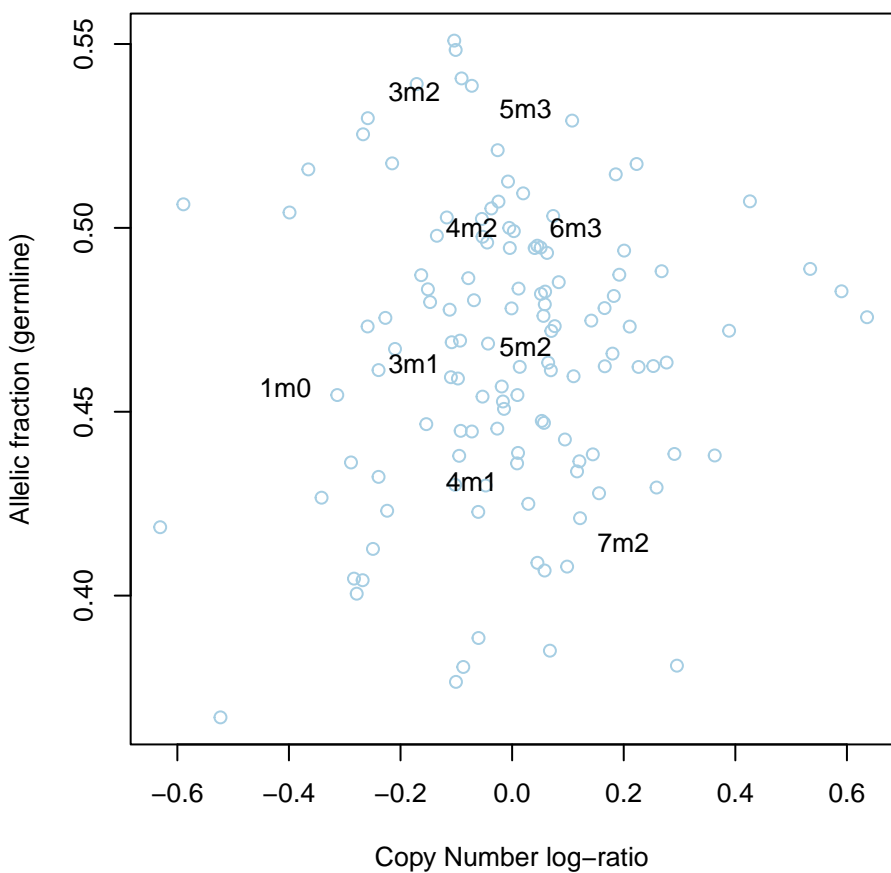


Purity: 0.16 Tumor ploidy: 4.743 SNV log-likelihood: -13.45 GoF: 97.3% Mean coverage: 464;227

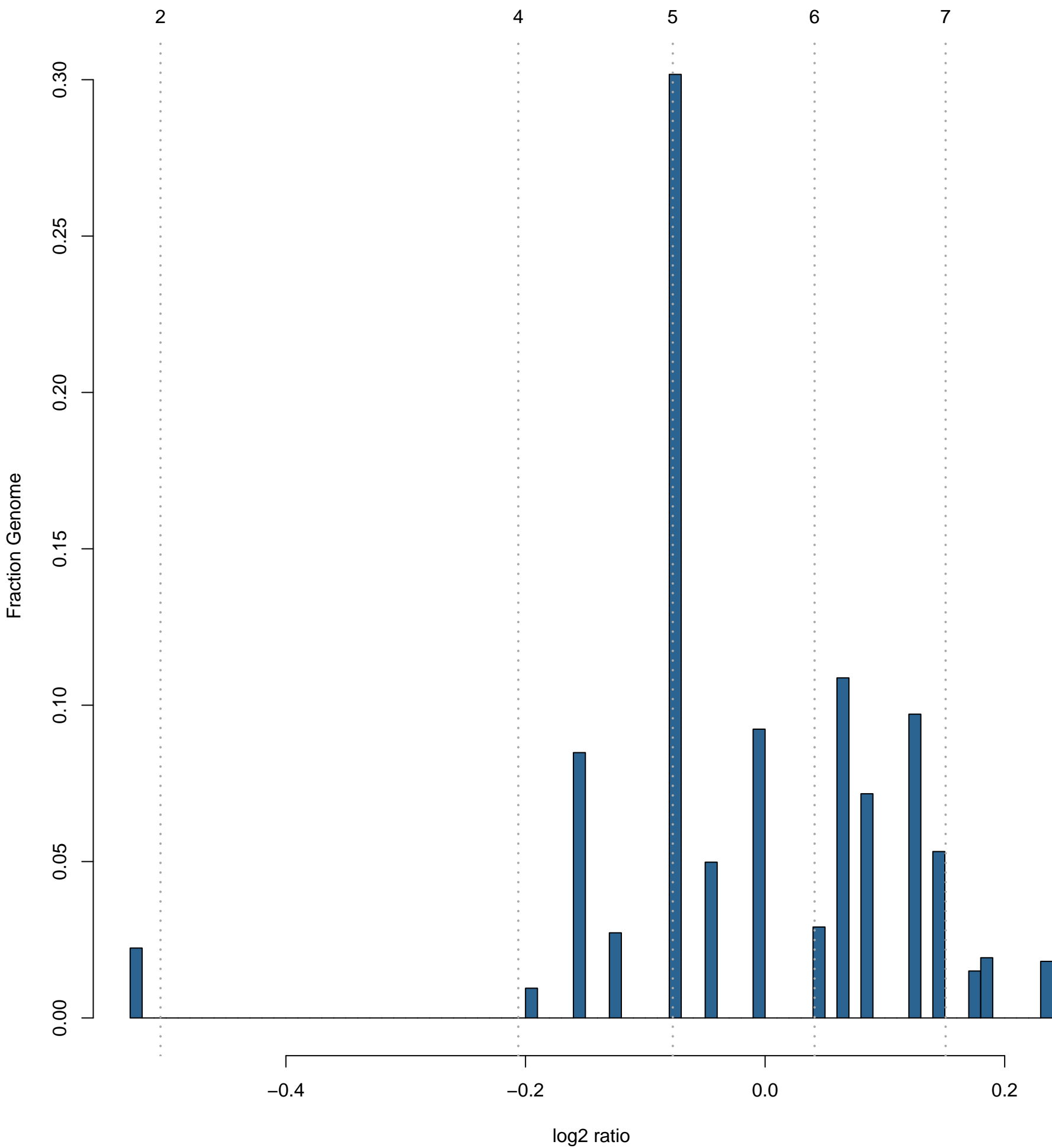


SCNA-fit log-likelihood: -12483.93



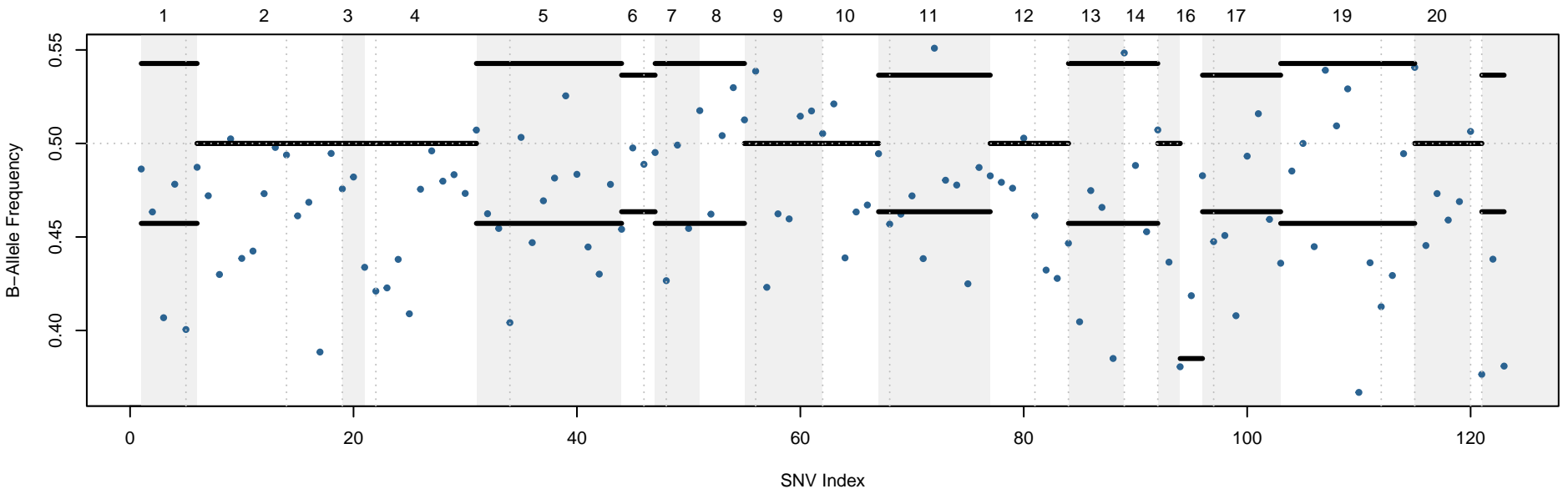


Purity: 0.23 Tumor ploidy: 5.642

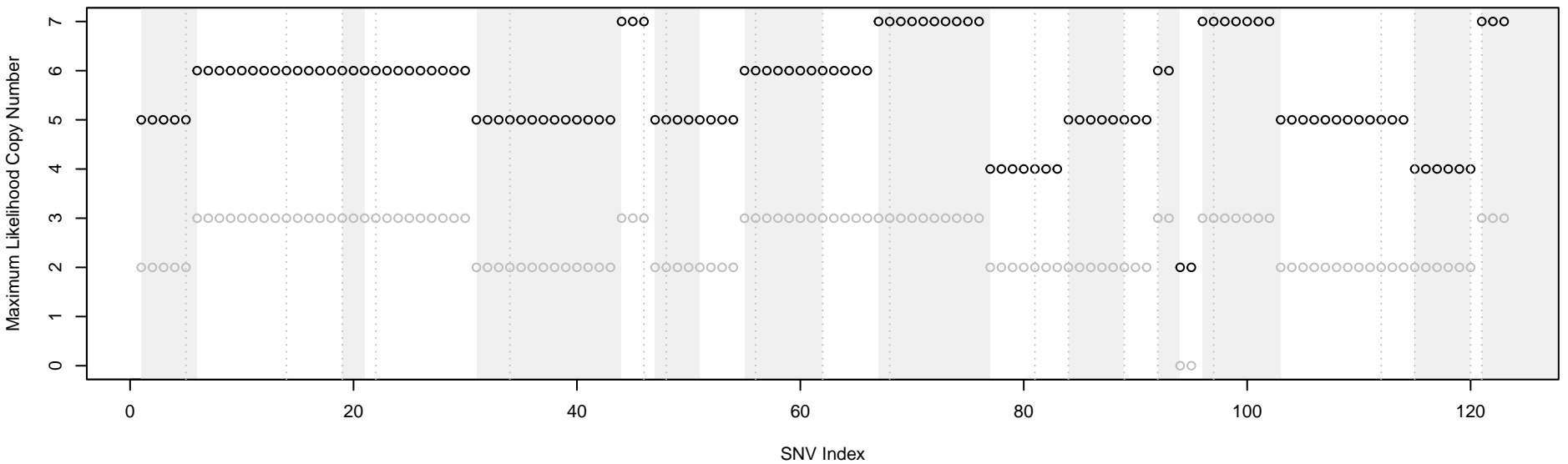
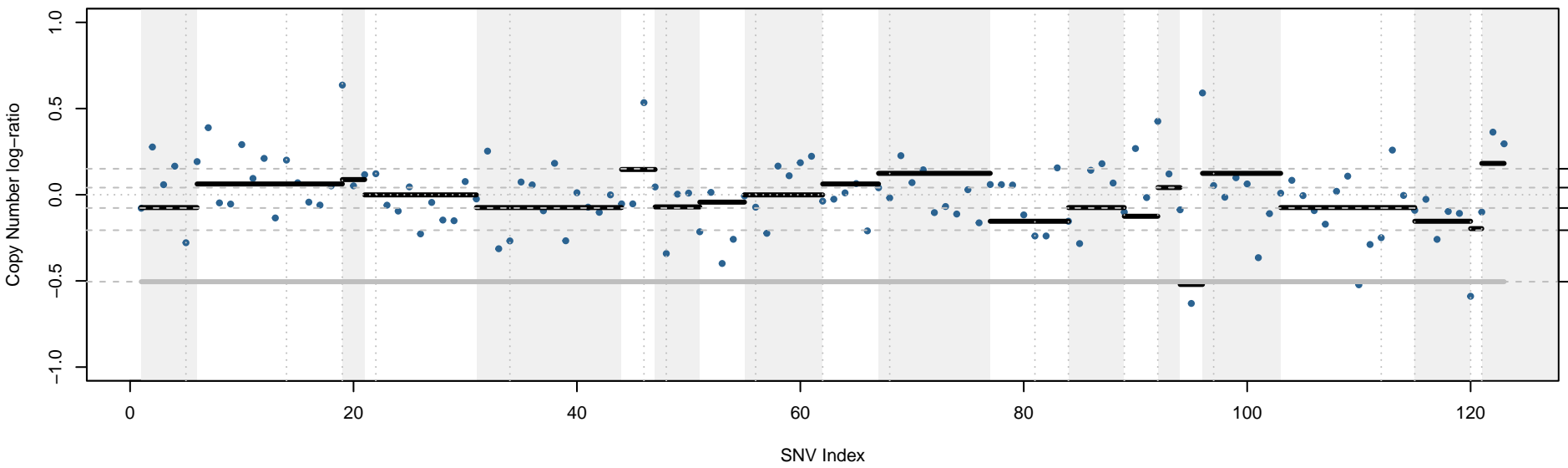


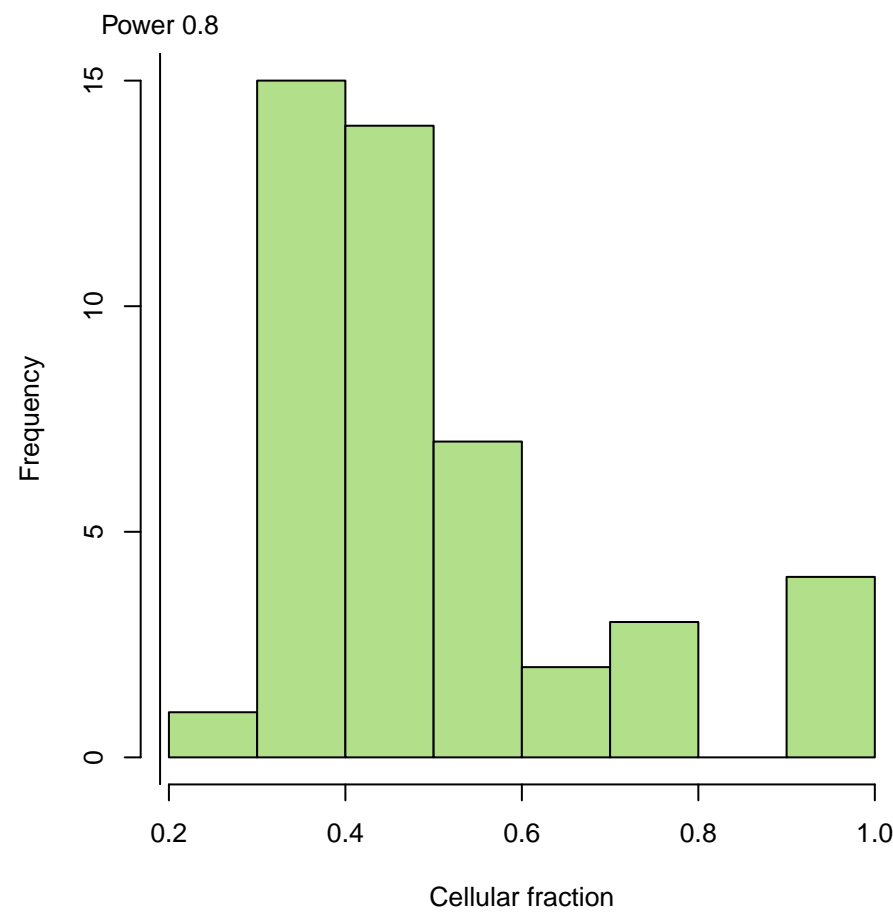
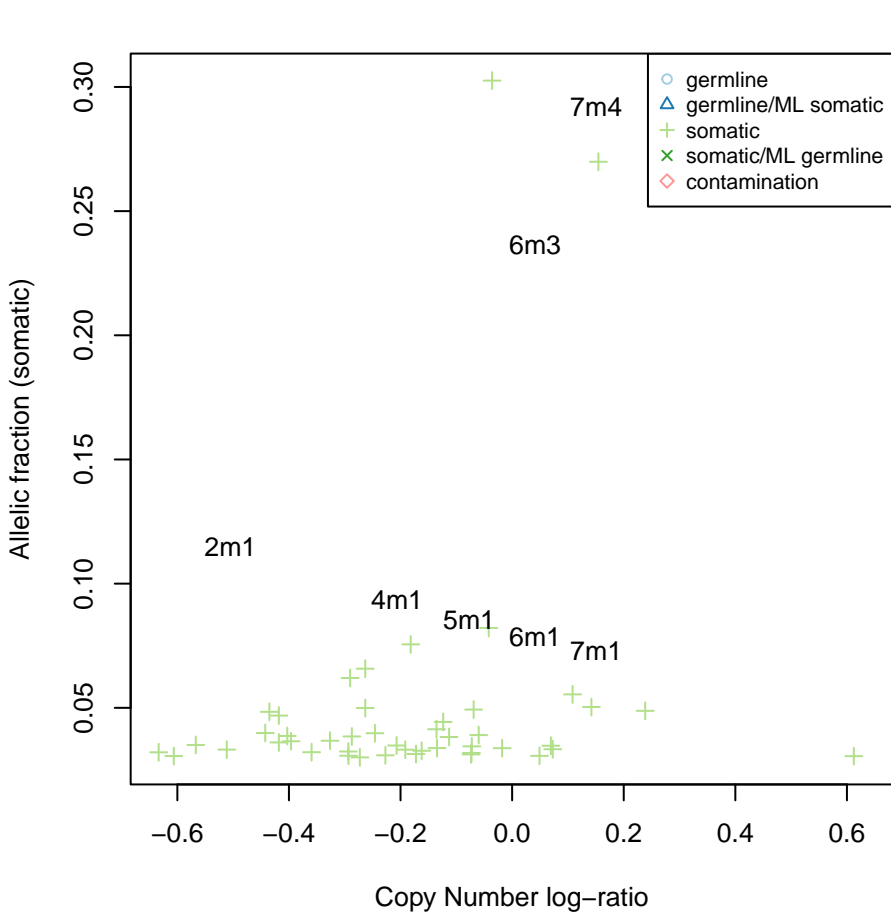
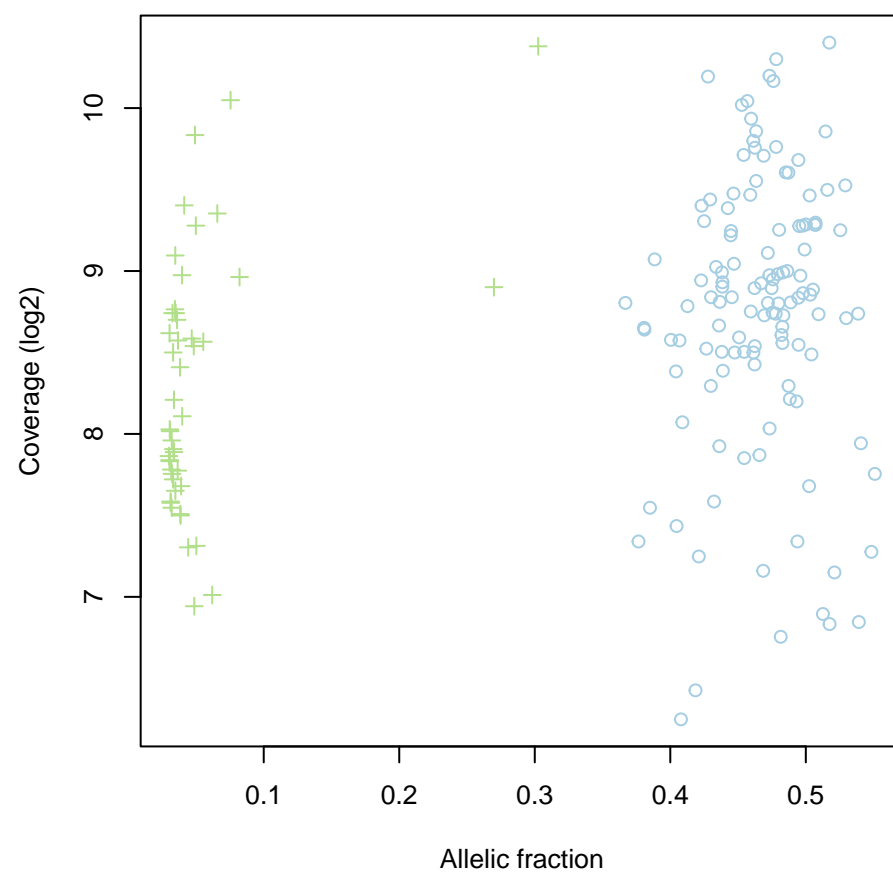
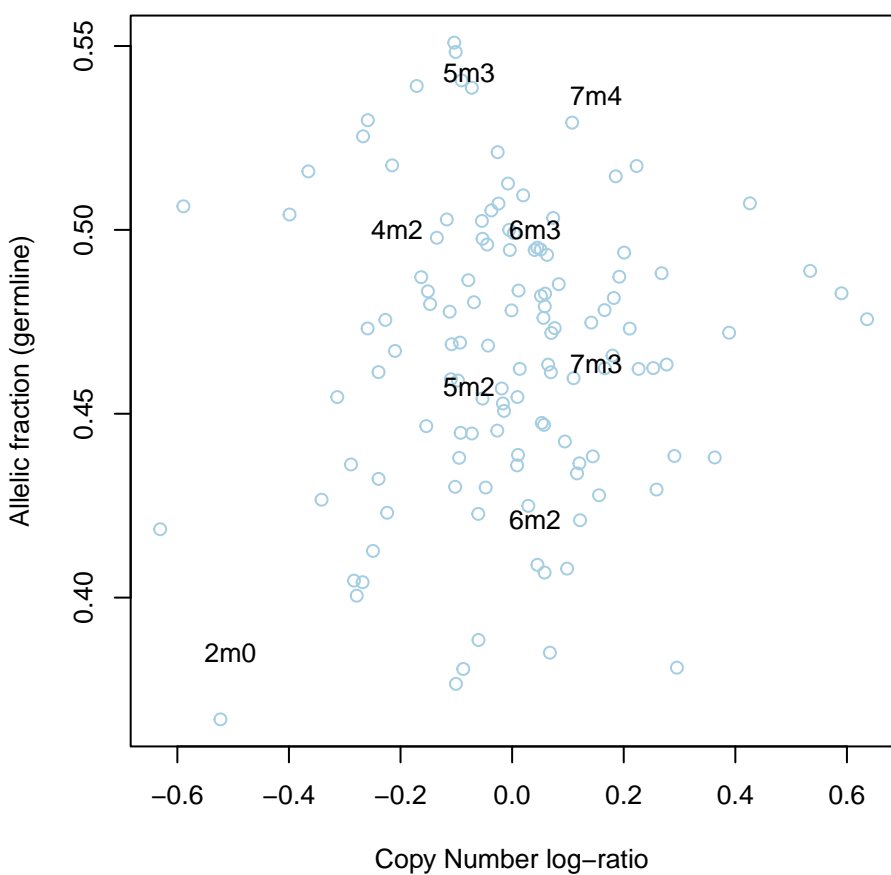


Purity: 0.23 Tumor ploidy: 5.642 SNV log-likelihood: -38.11 GoF: 96.7% Mean coverage: 464;227

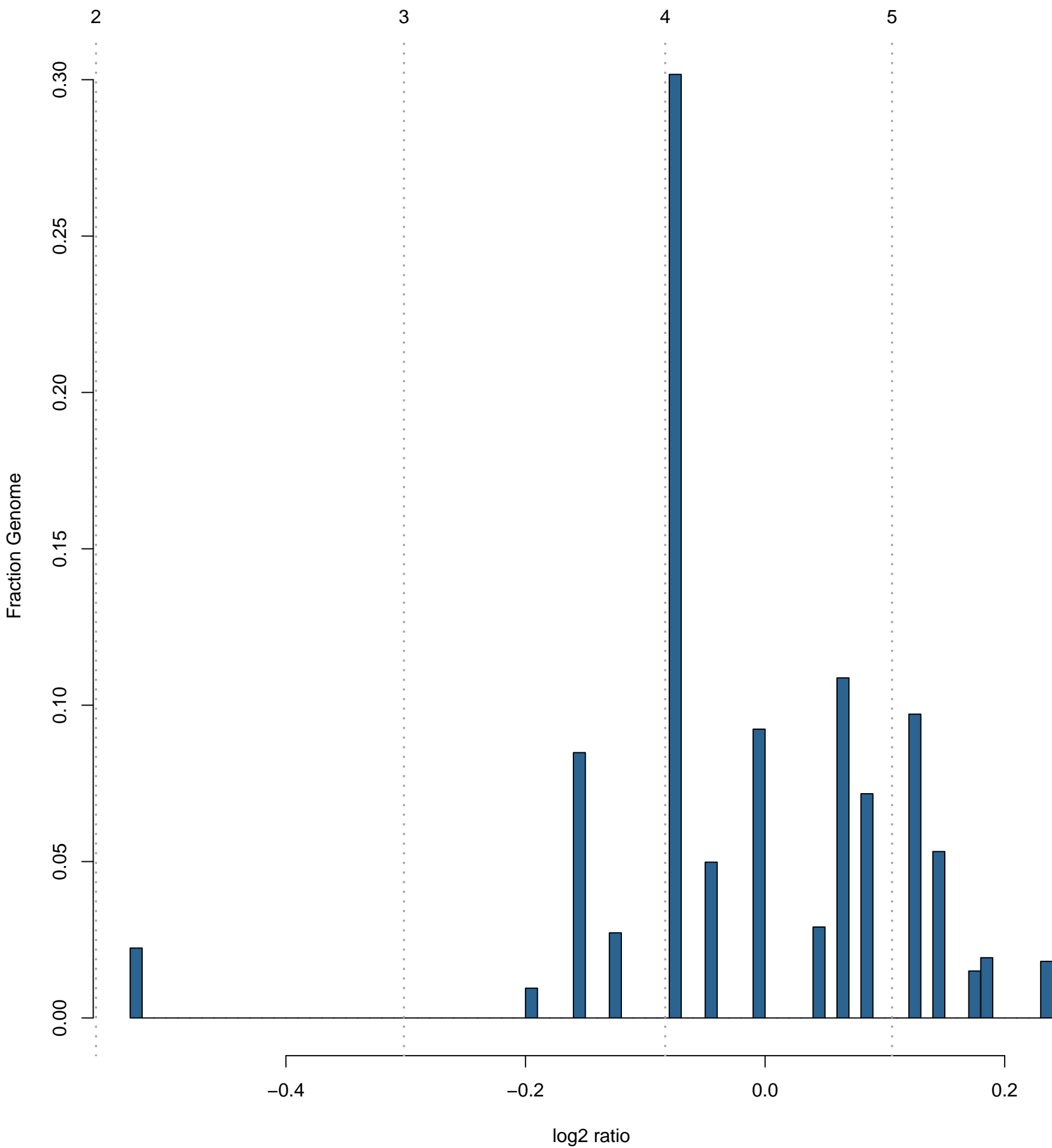


SCNA-fit log-likelihood: -12485.86

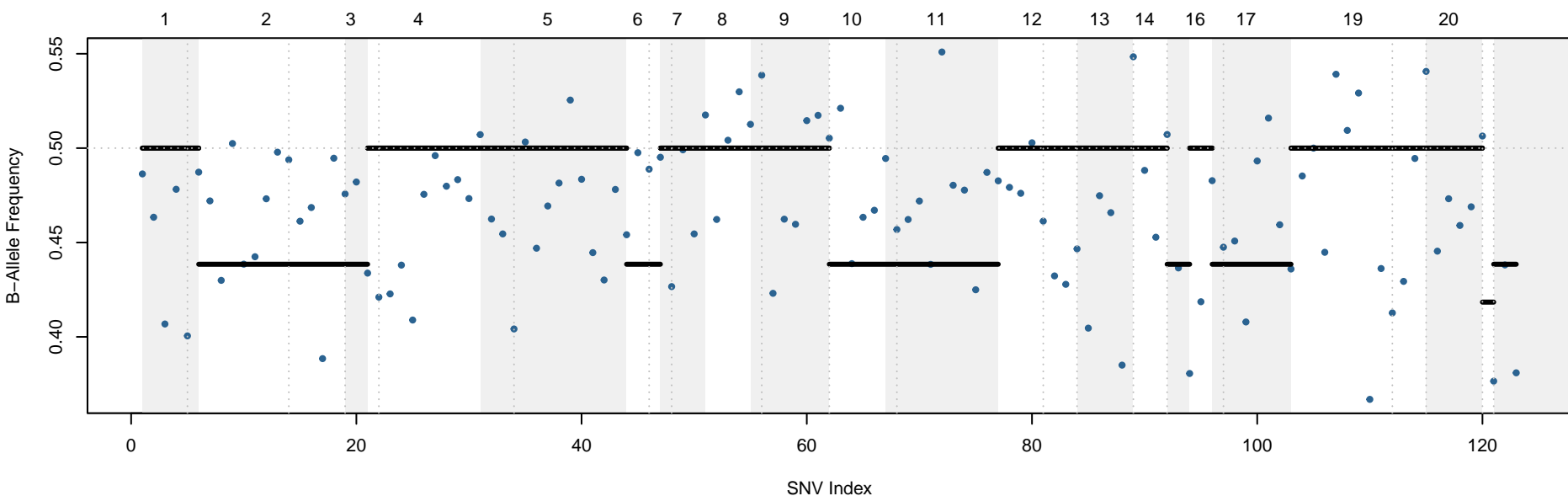




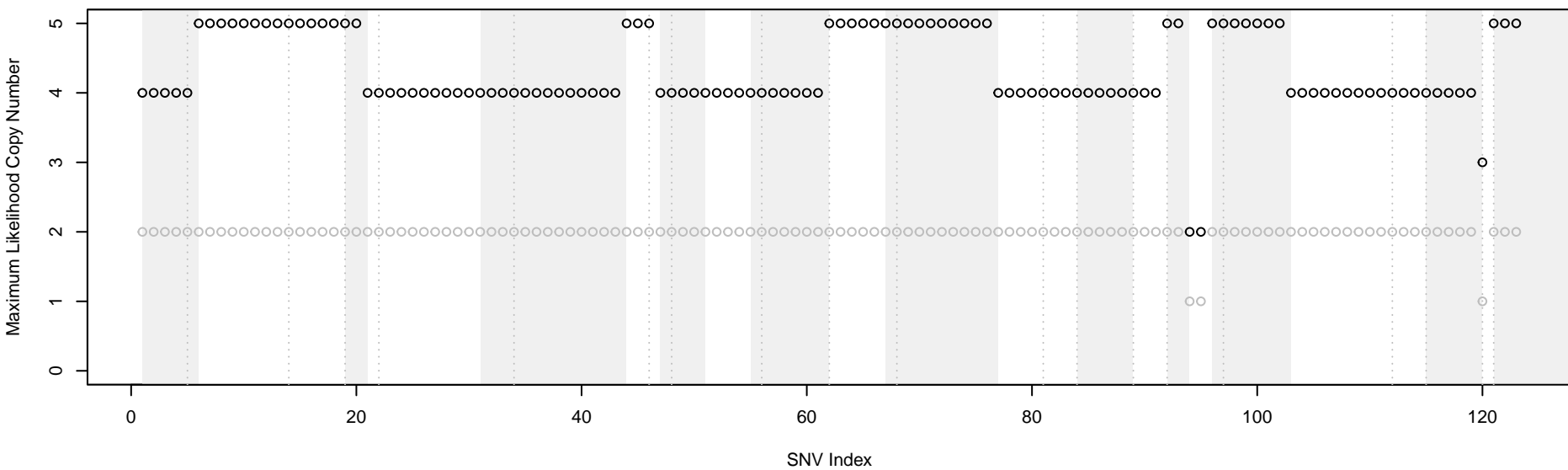
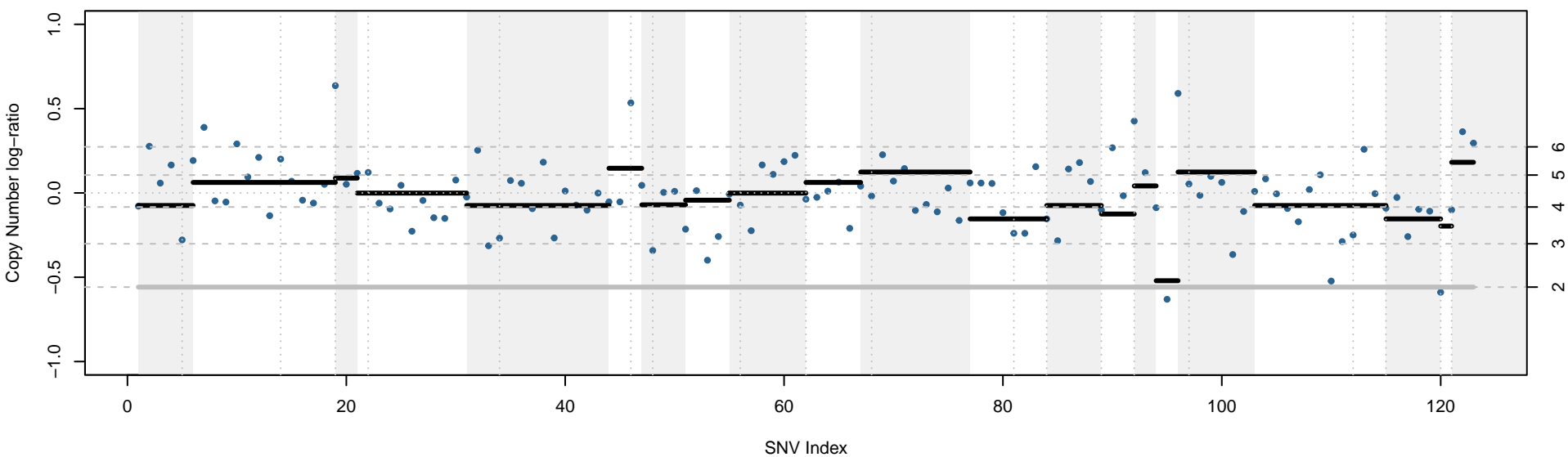
Purity: 0.39 Tumor ploidy: 4.424

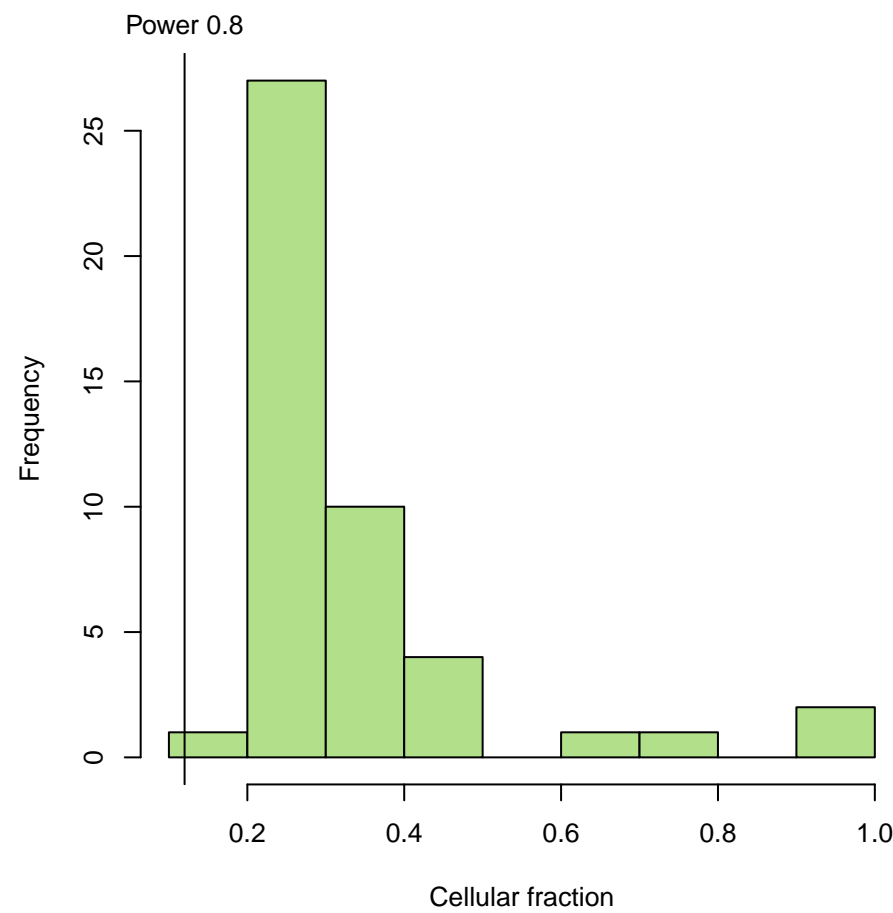
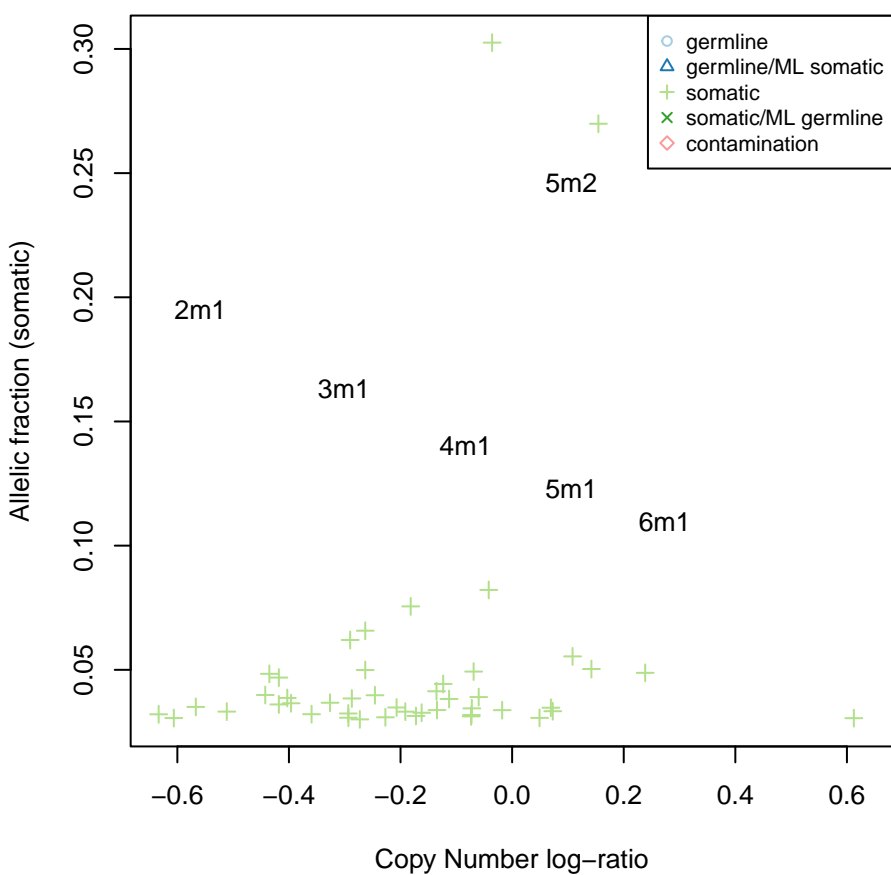
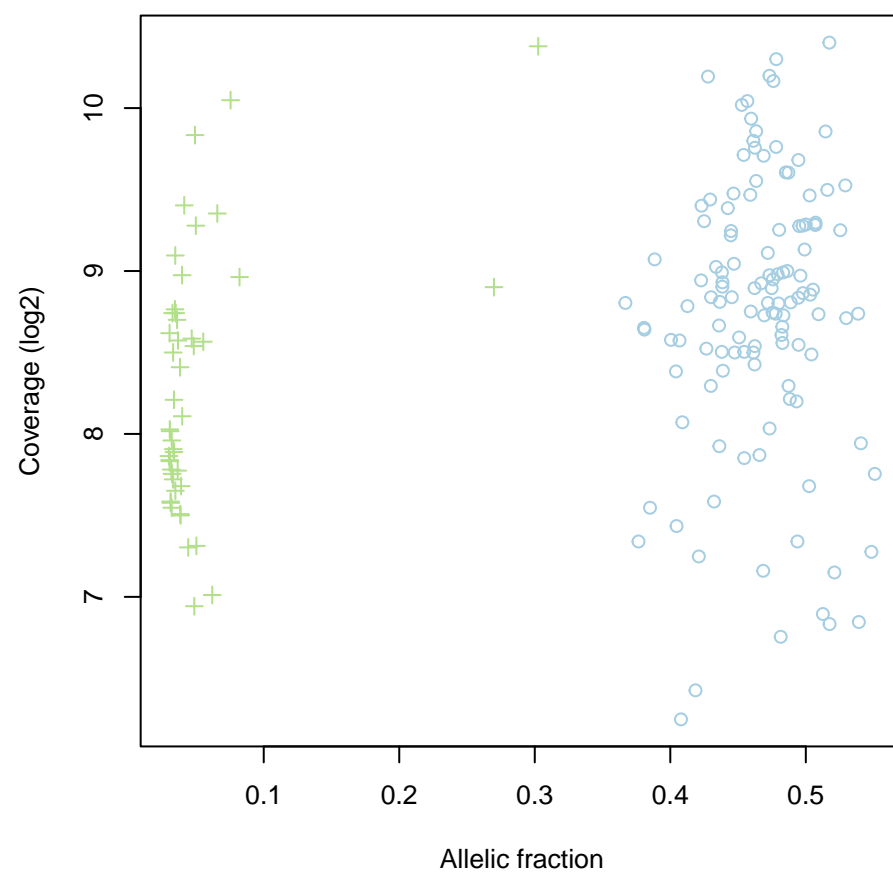
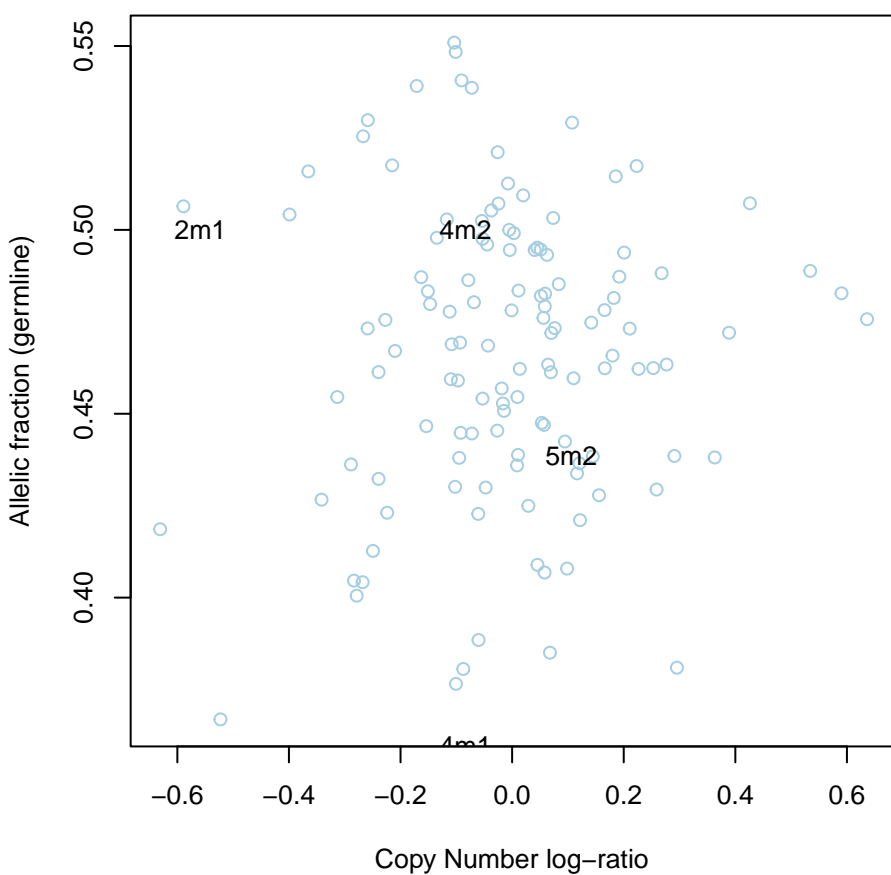


Purity: 0.39 Tumor ploidy: 4.424 SNV log-likelihood: -49.08 GoF: 90.5% Mean coverage: 464;227

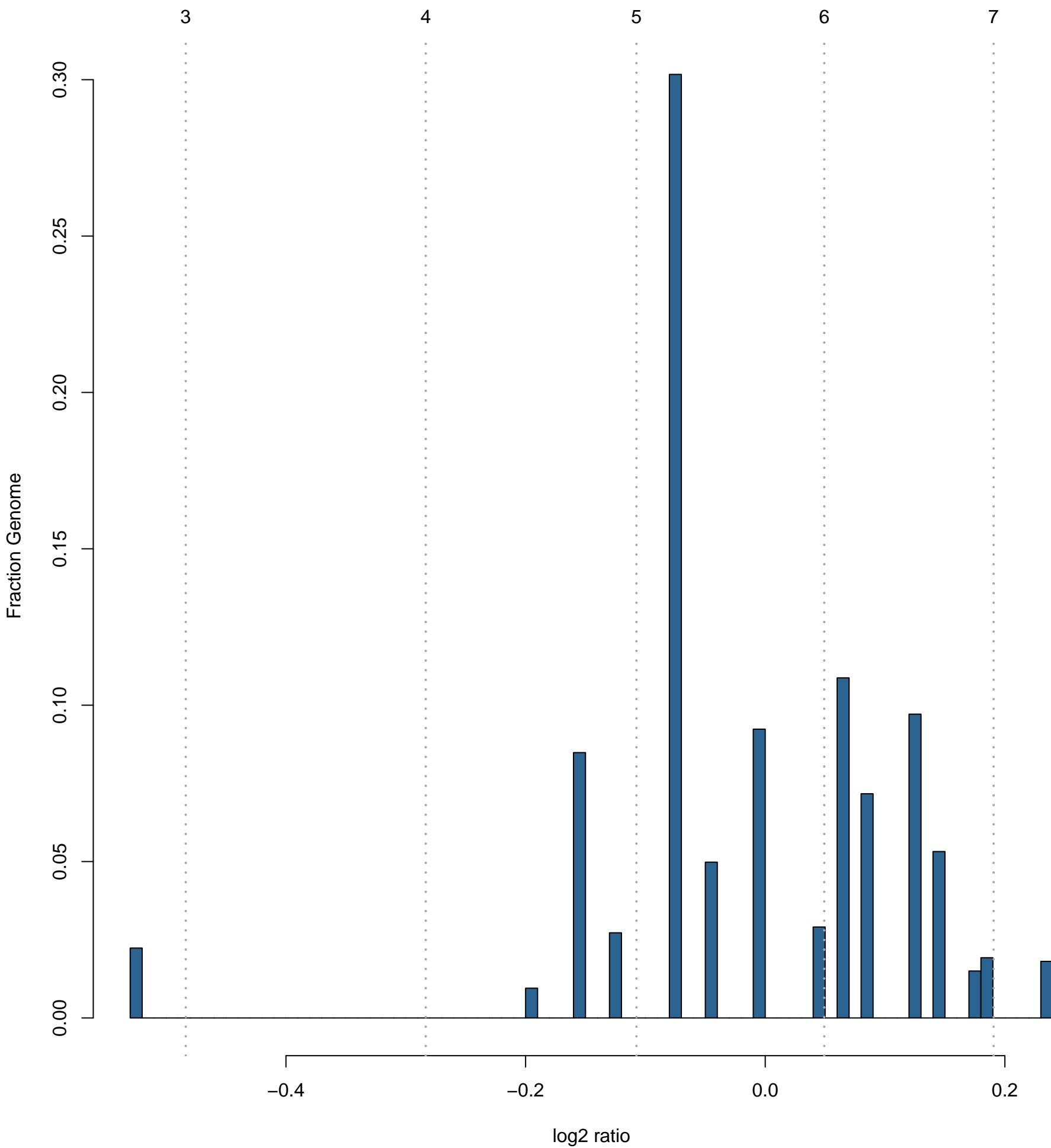


SCNA-fit log-likelihood: -12504.09

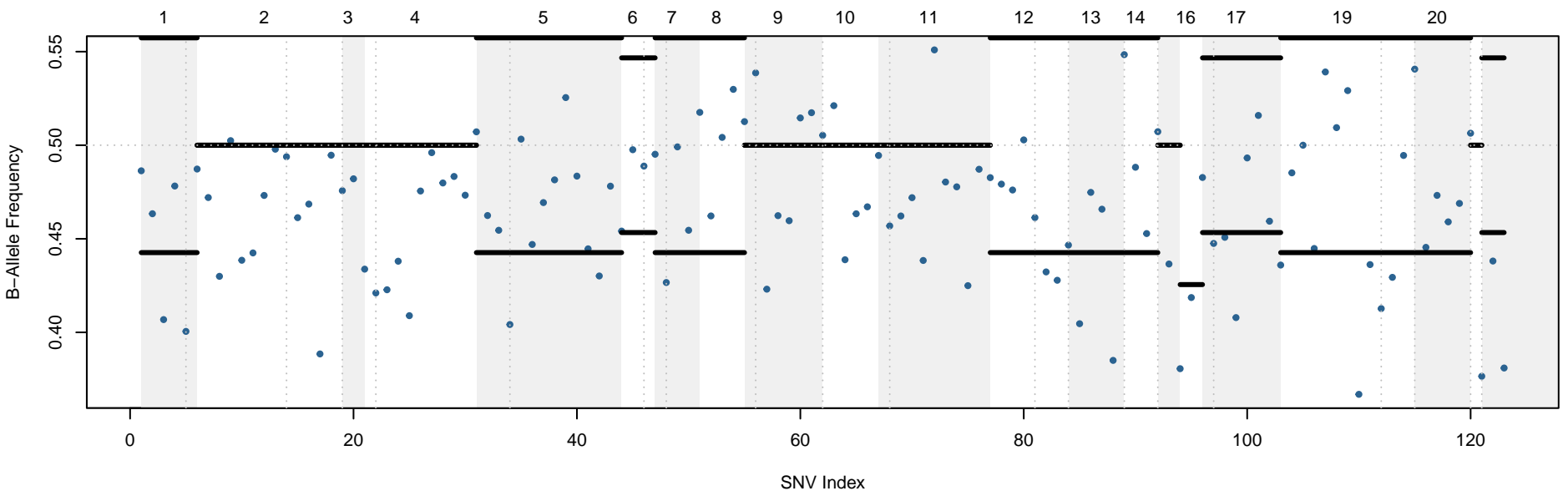




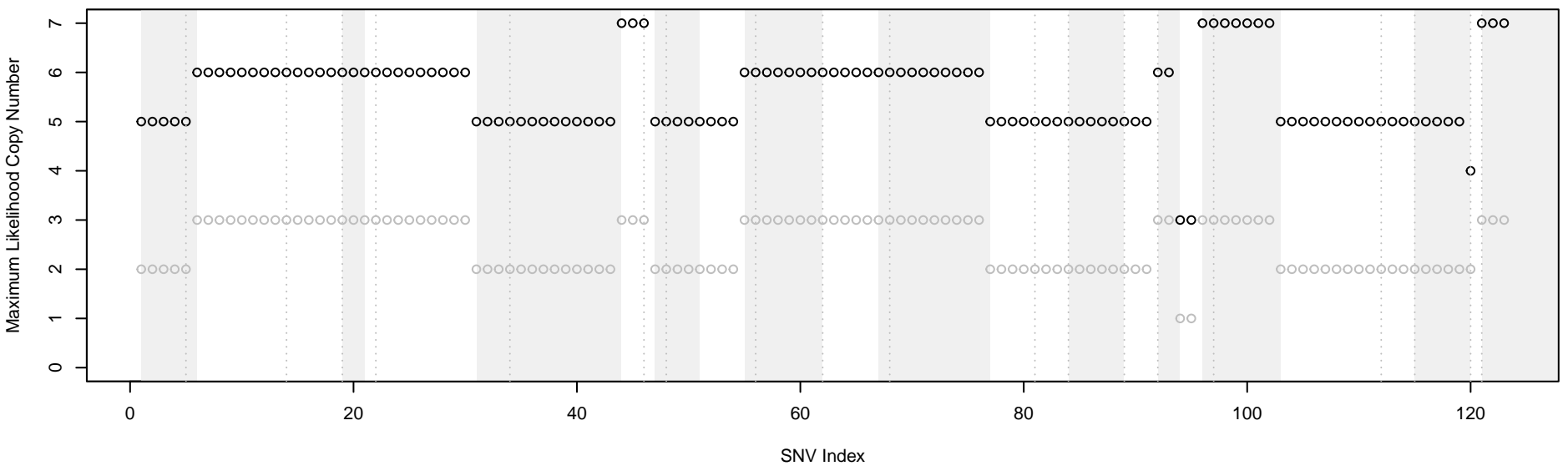
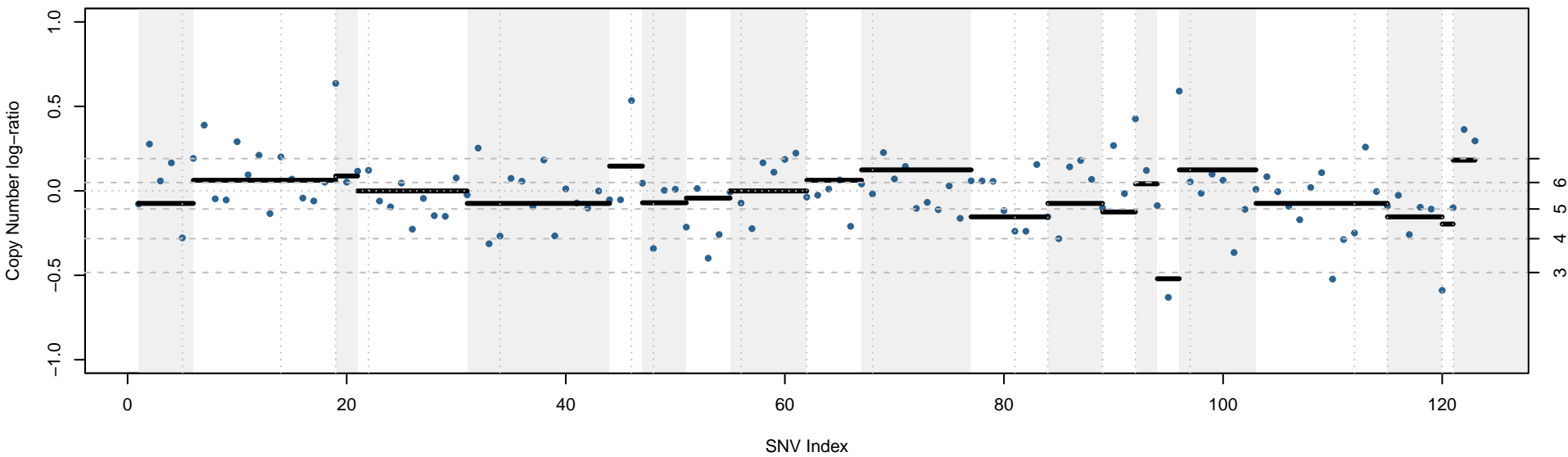
Purity: 0.35 Tumor ploidy: 5.674

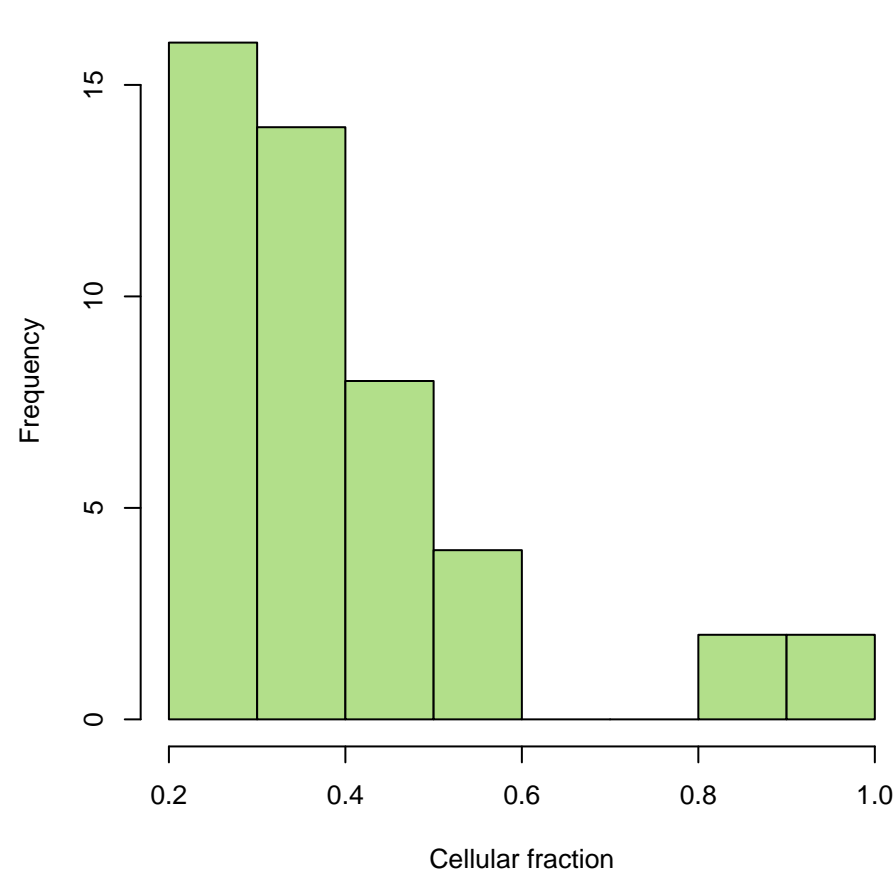
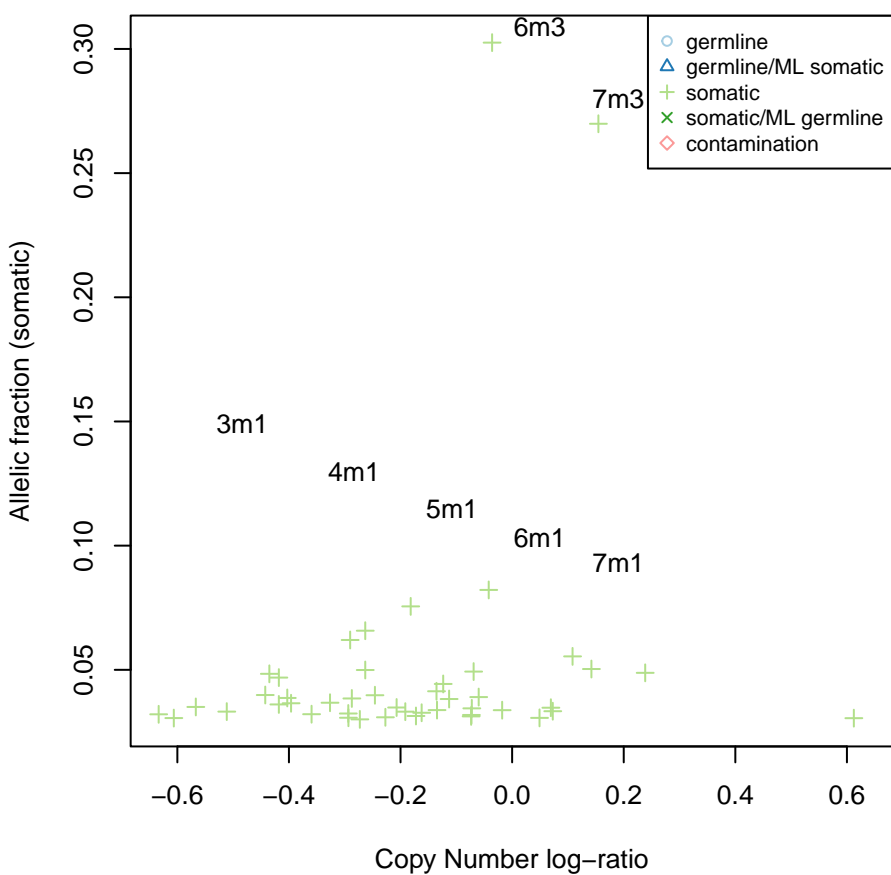
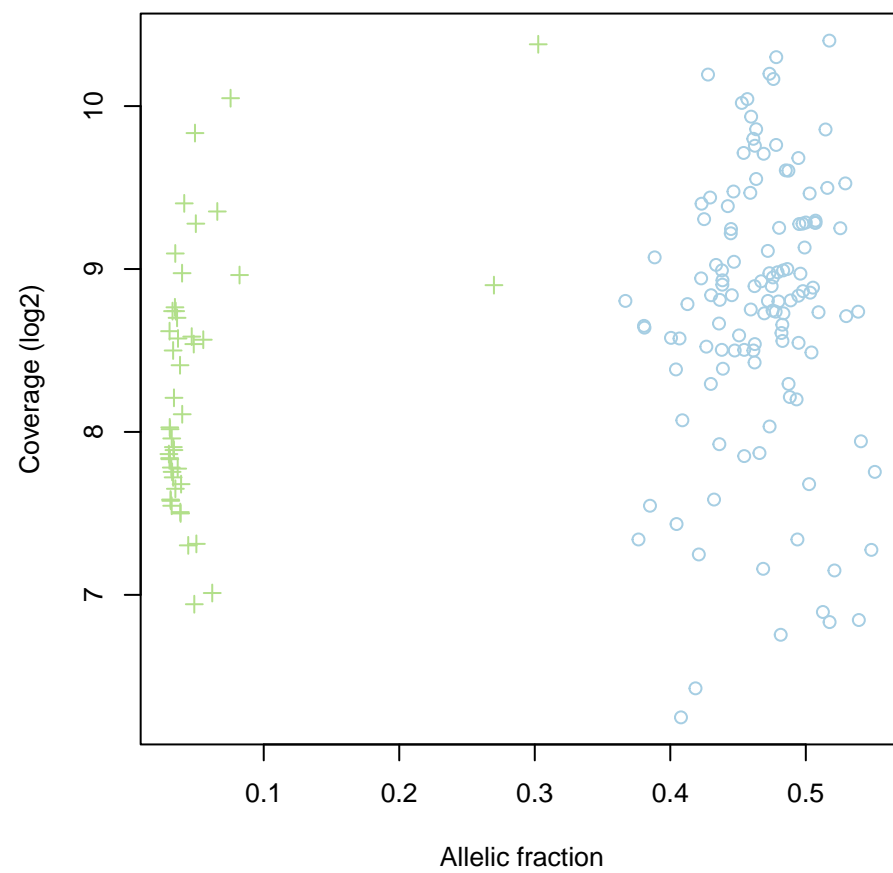
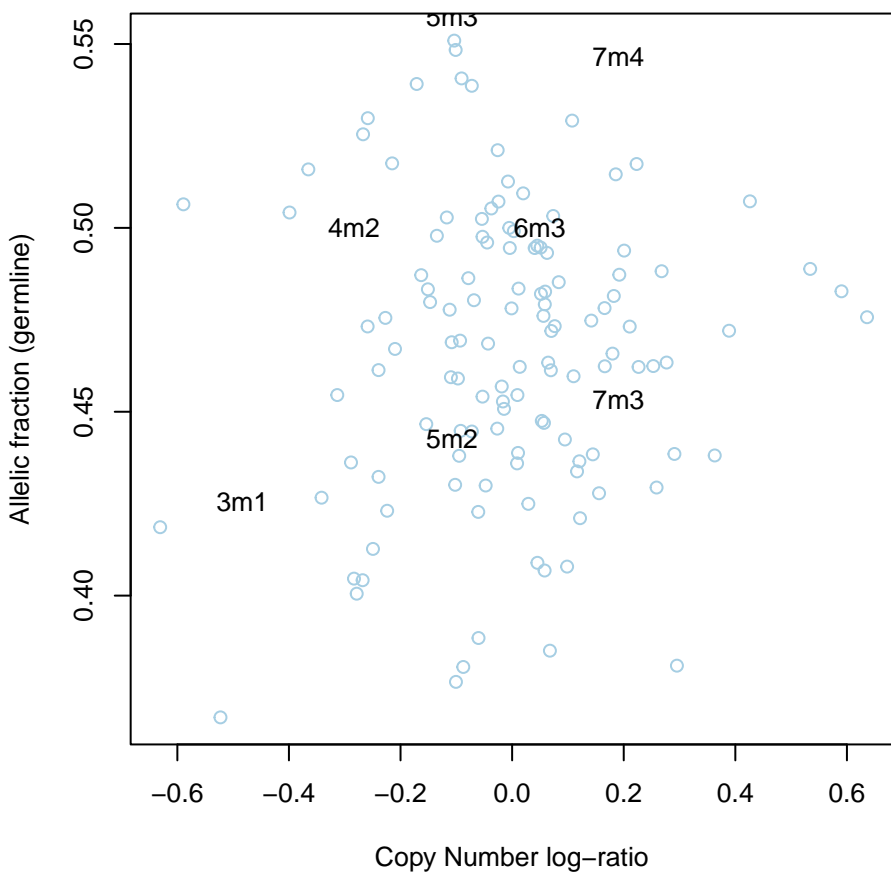


Purity: 0.35 Tumor ploidy: 5.674 SNV log-likelihood: -54.18 GoF: 94.3% Mean coverage: 464;227



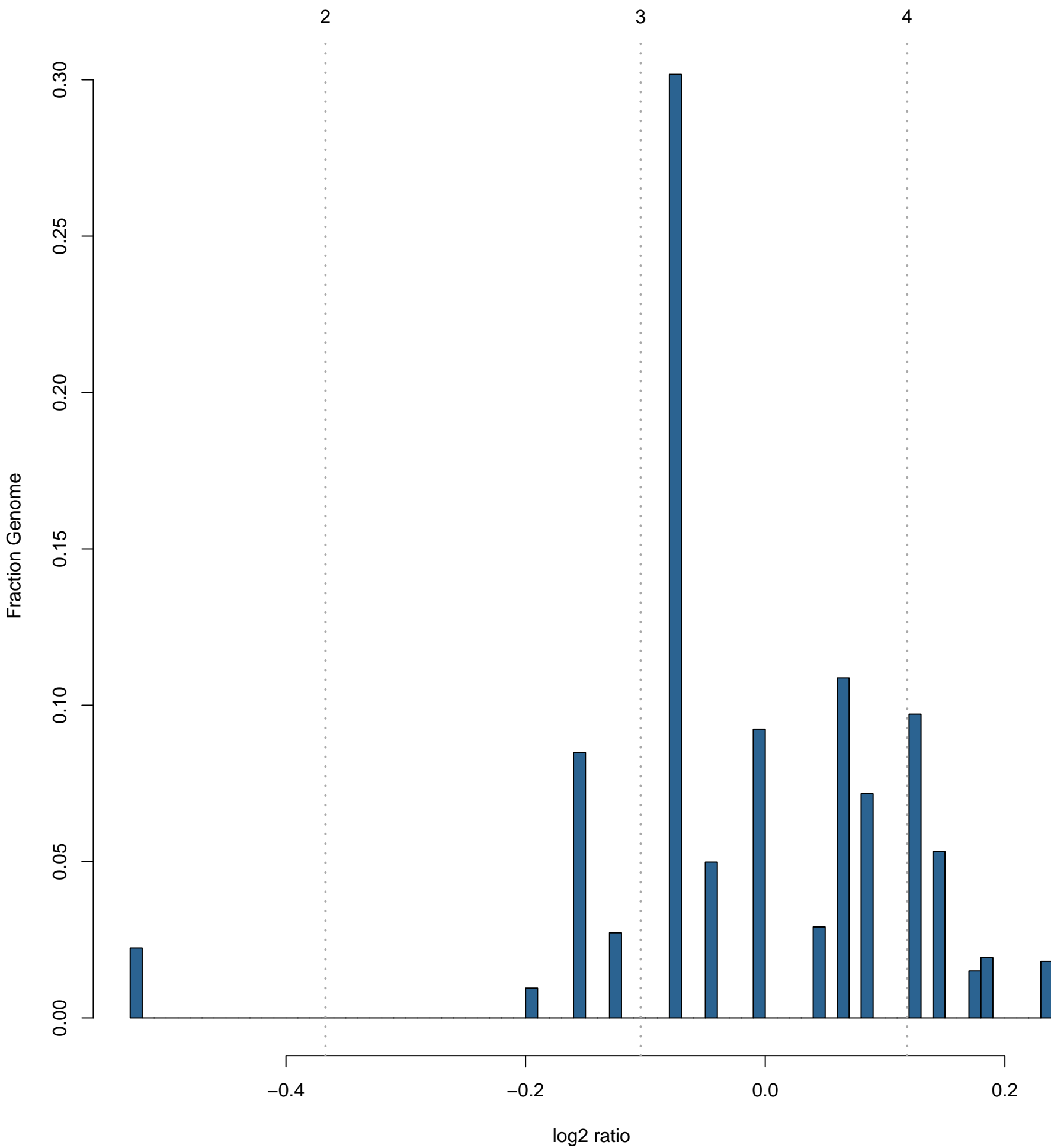
SCNA-fit log-likelihood: -12497.63



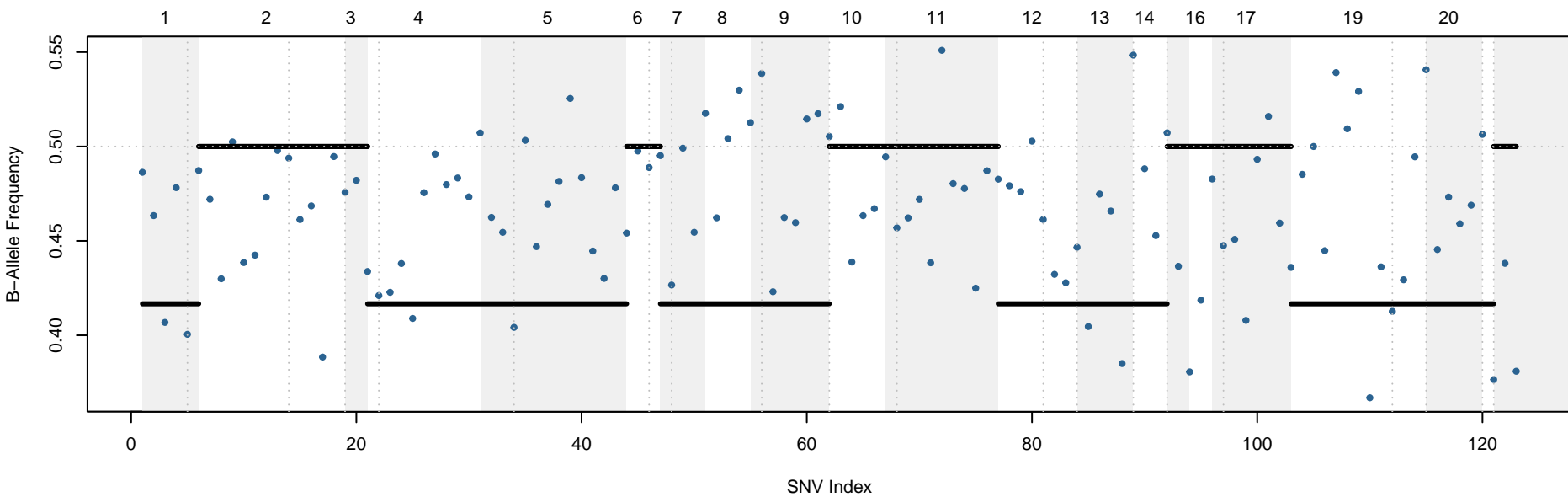




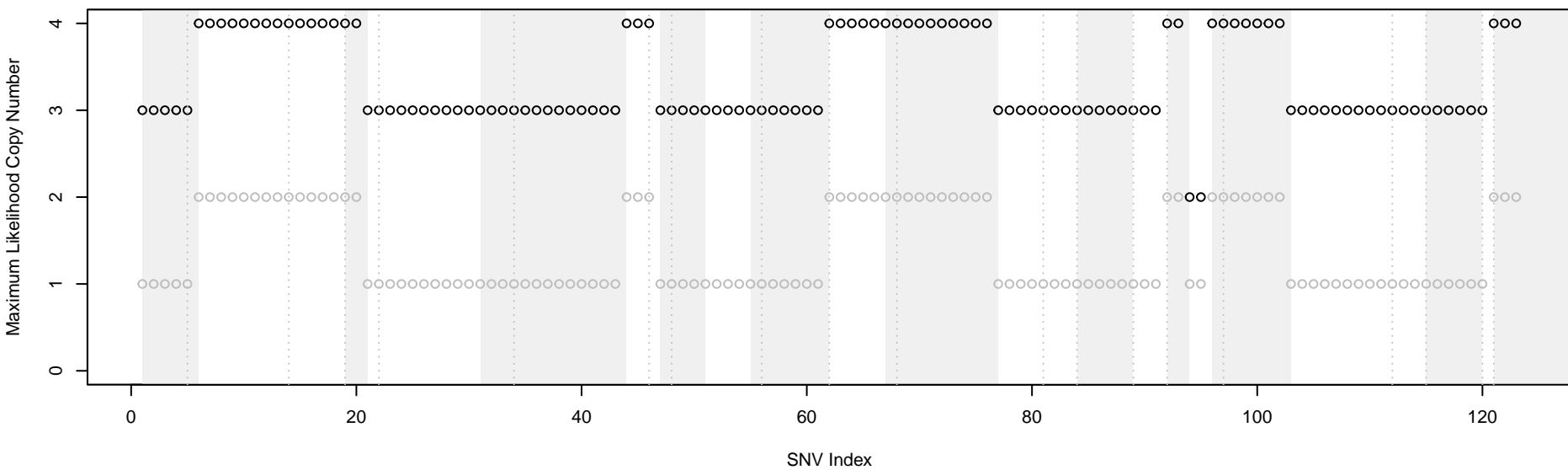
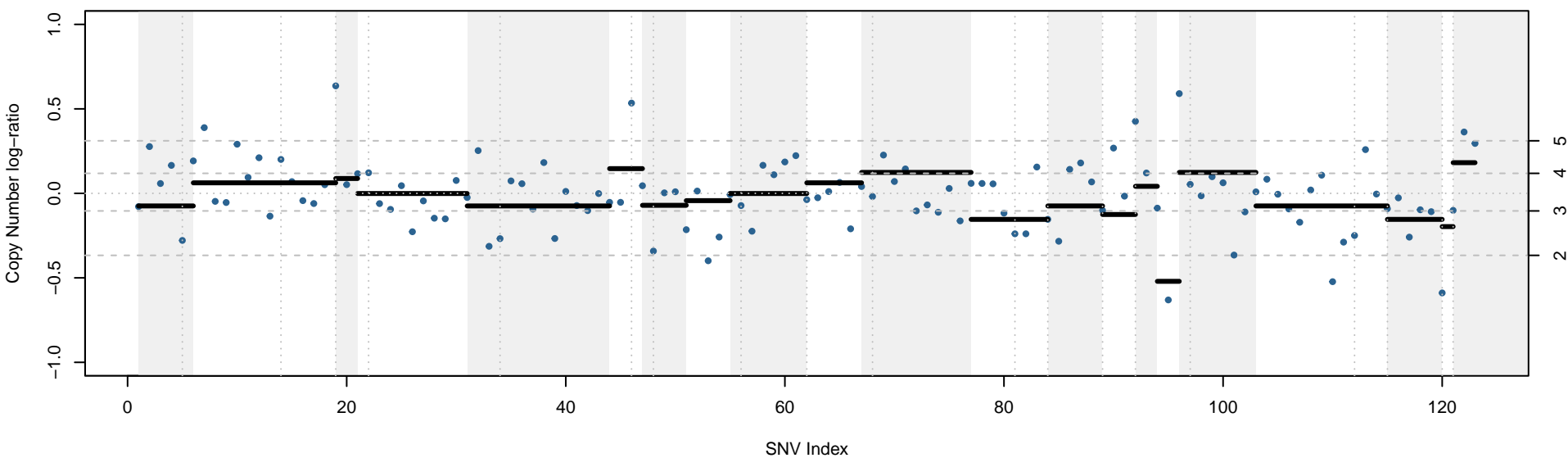
Purity: 0.4 Tumor ploidy: 3.448

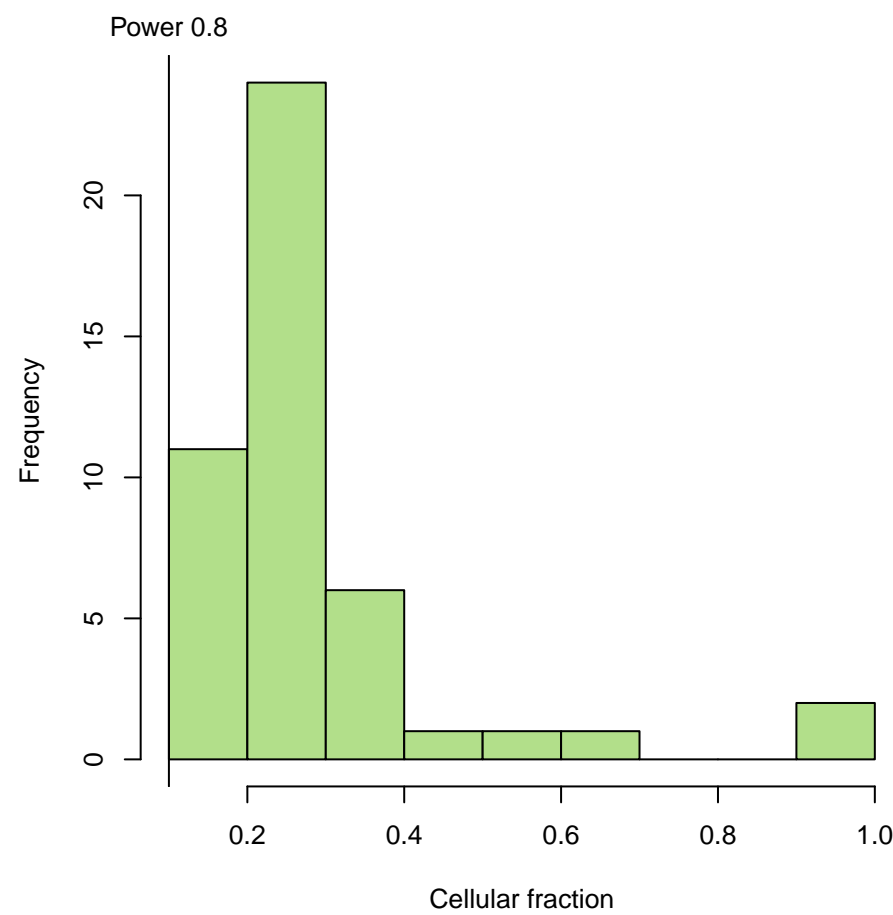
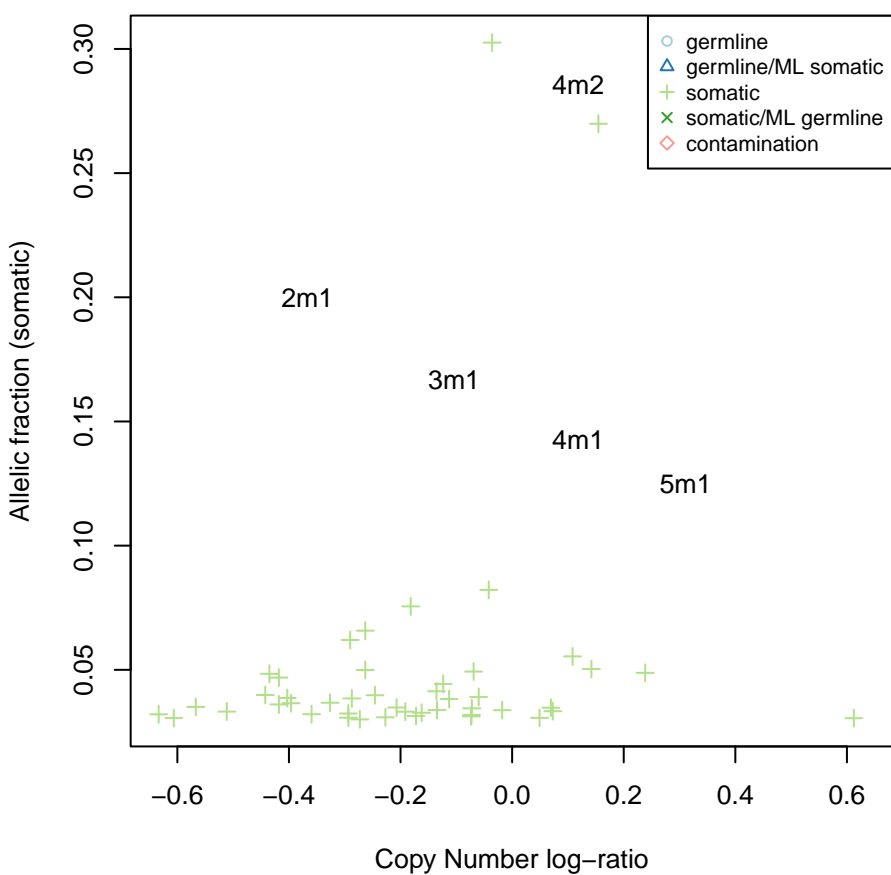
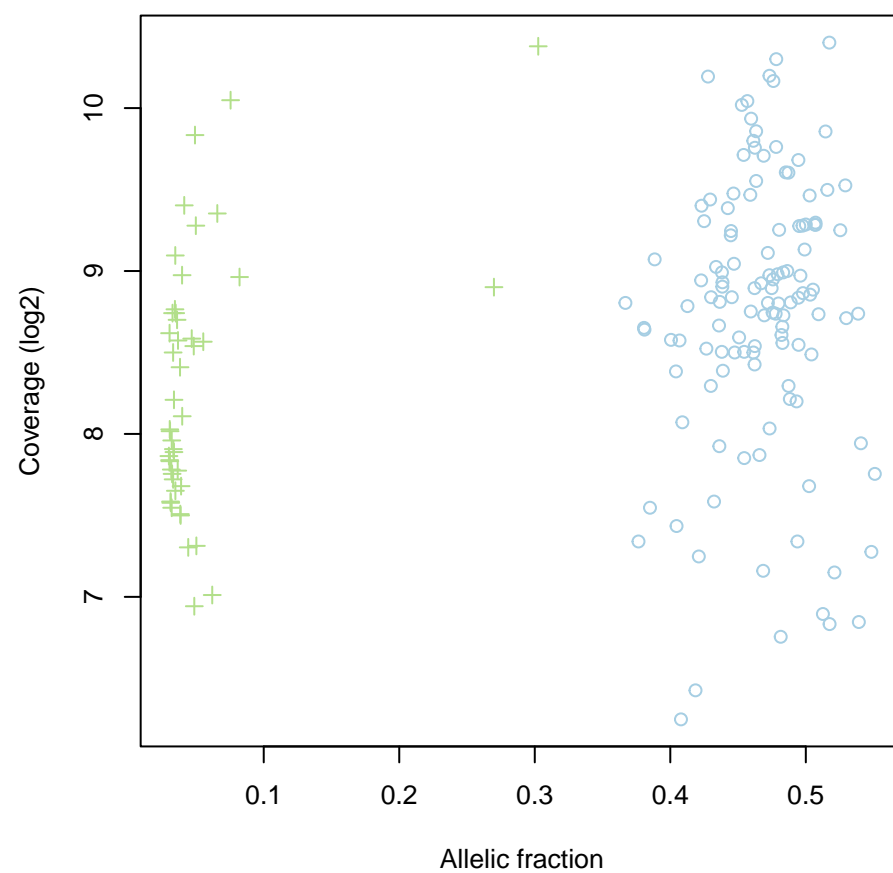
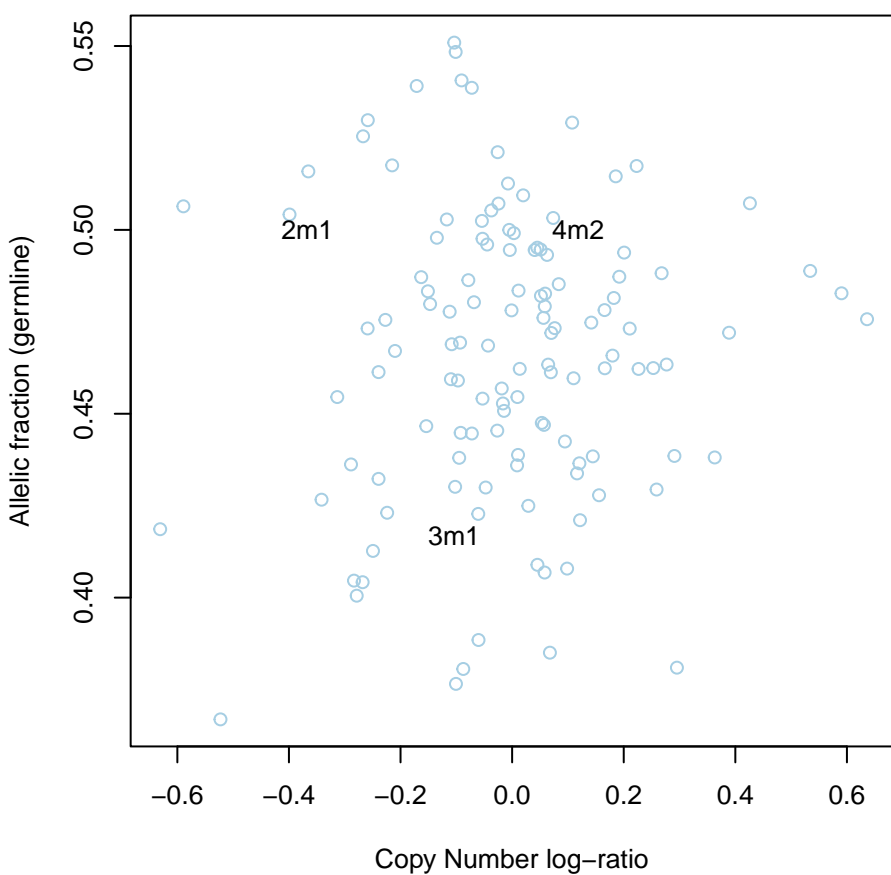


Purity: 0.4 Tumor ploidy: 3.448 SNV log-likelihood: -90.34 GoF: 86% Mean coverage: 464;227

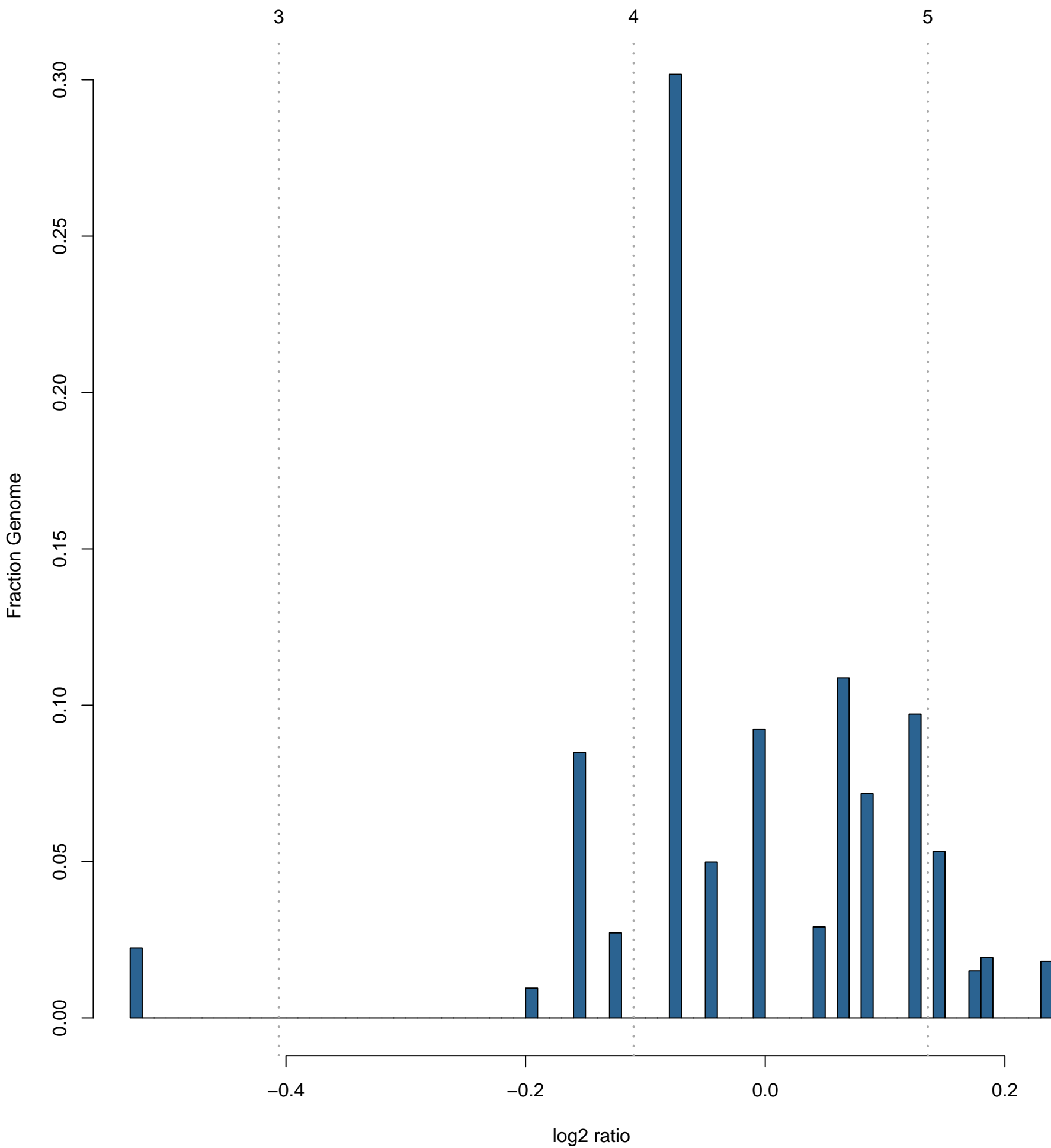


SCNA-fit log-likelihood: -12521.14

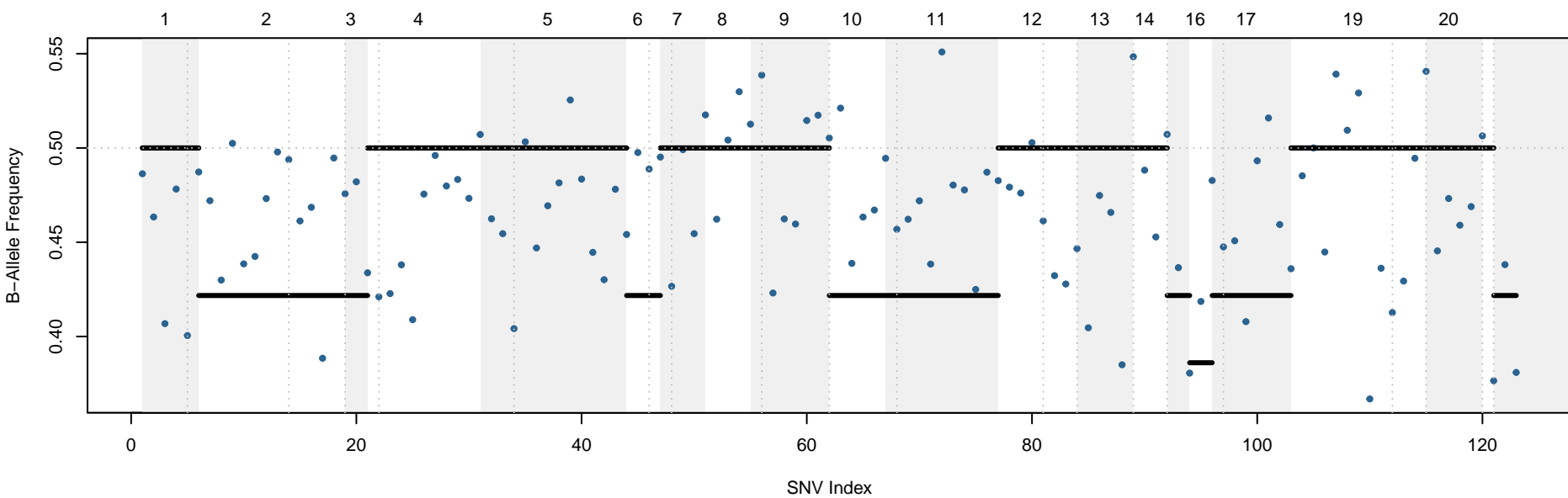




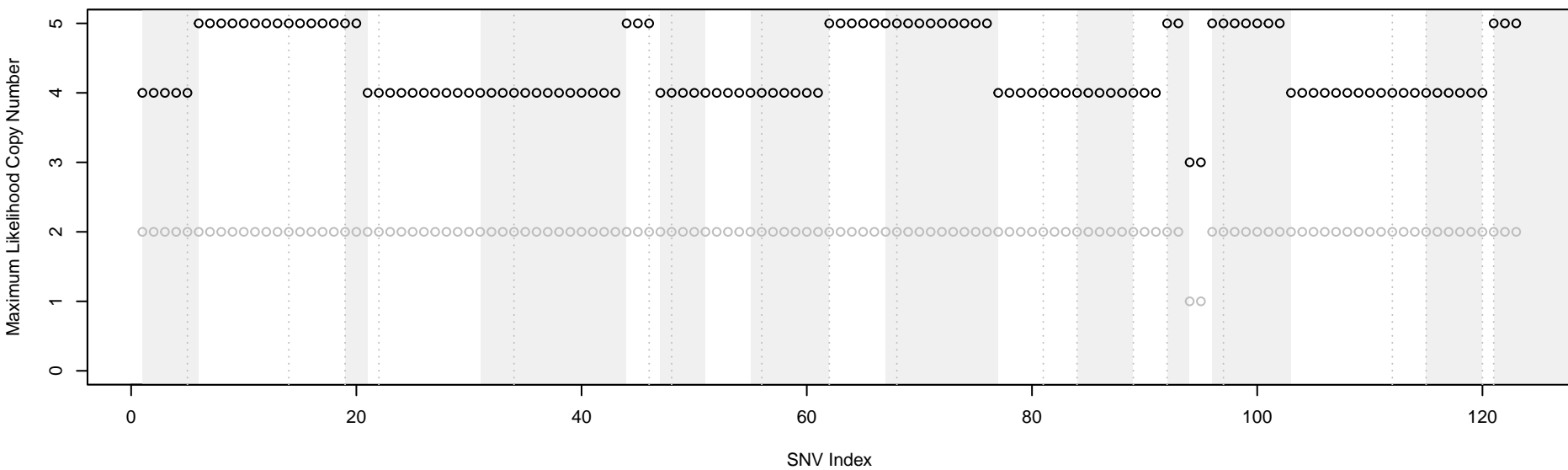
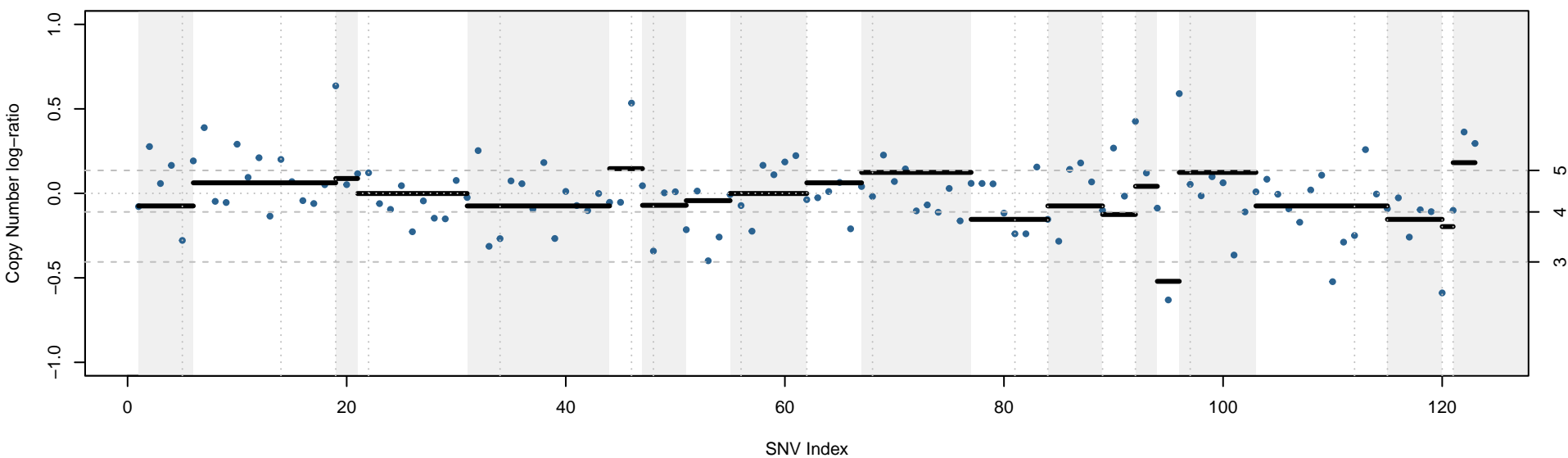
Purity: 0.59 Tumor ploidy: 4.426

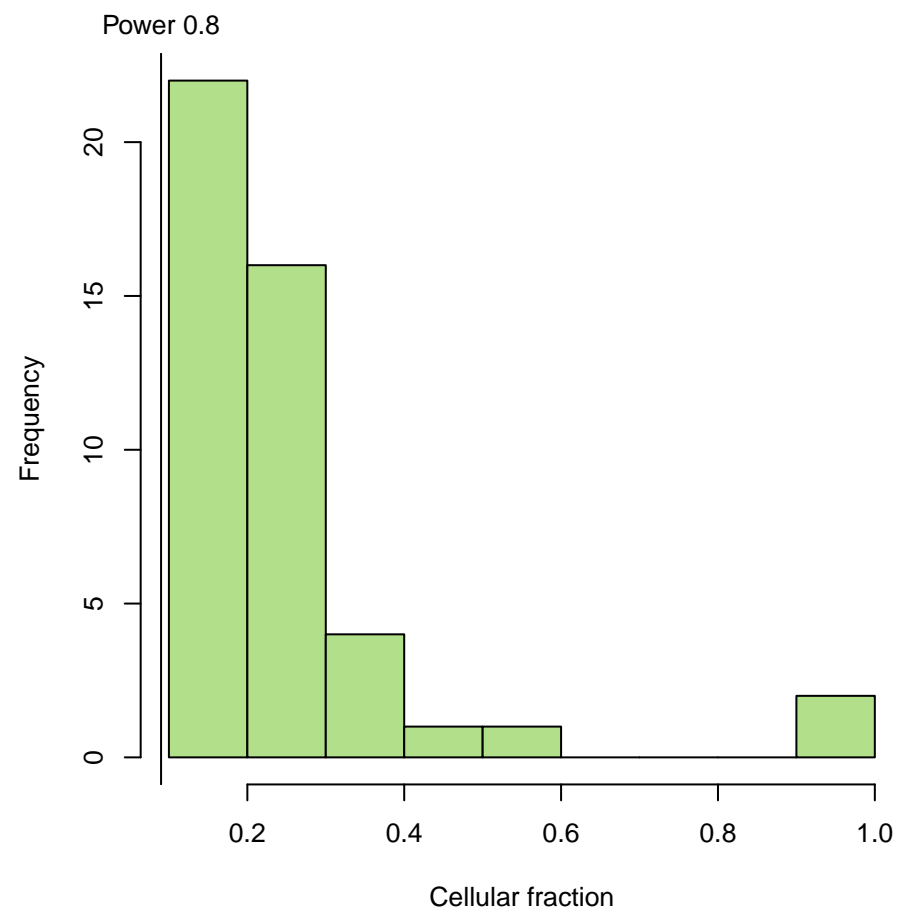
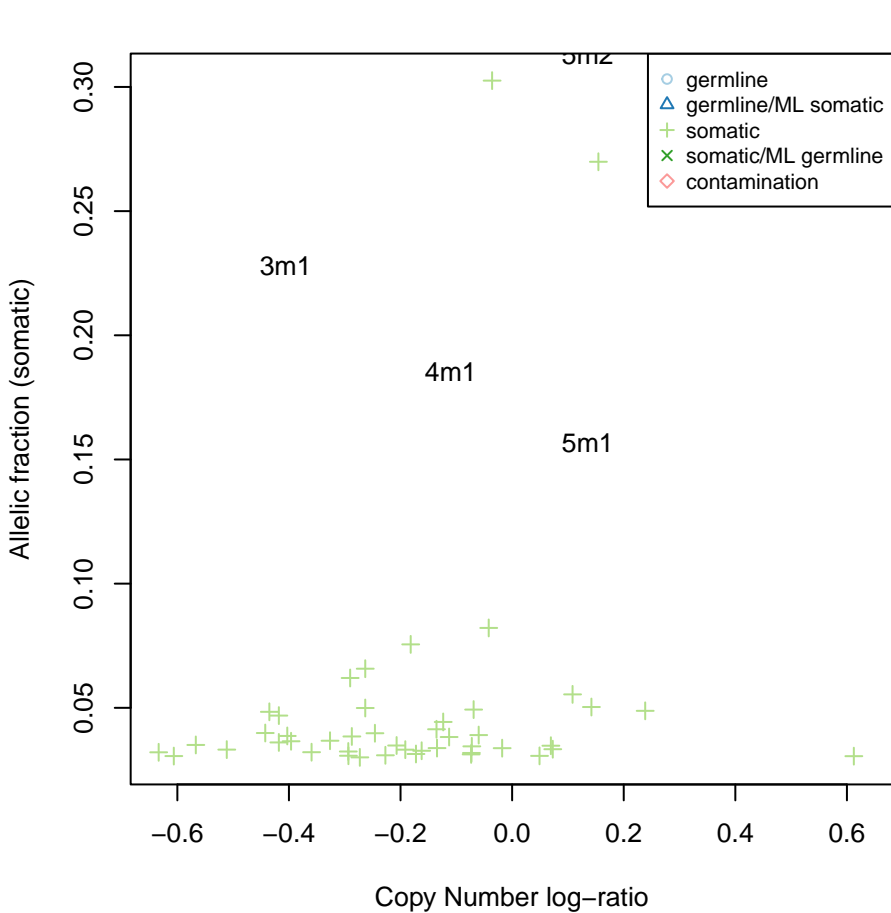
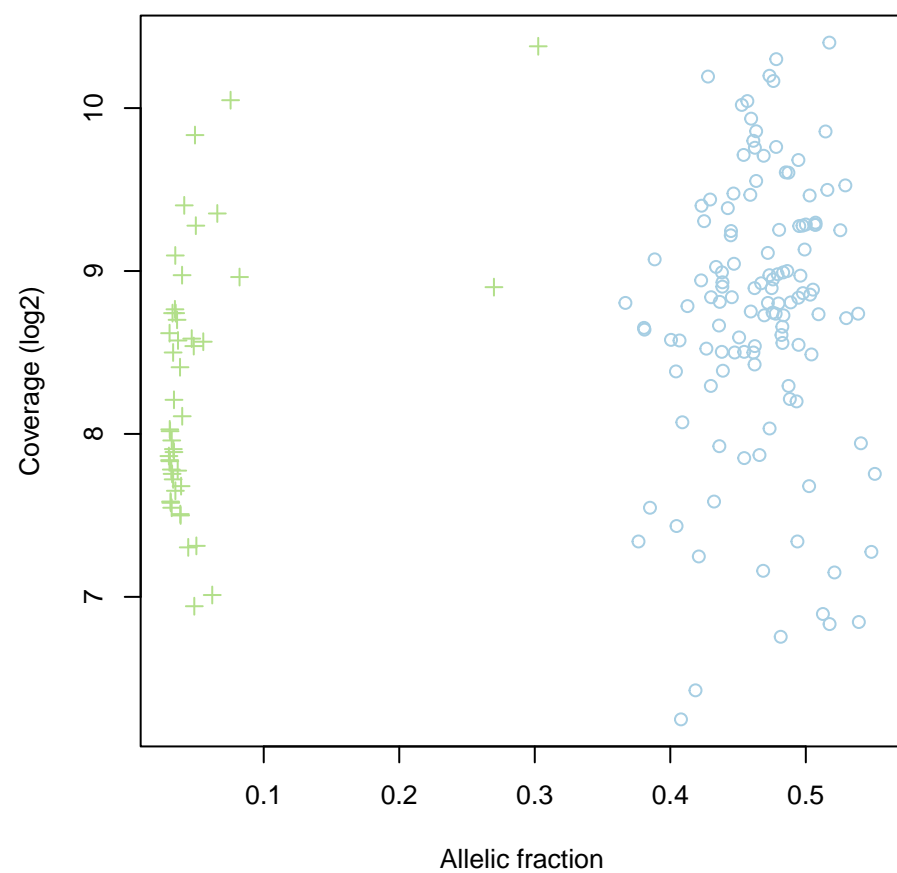
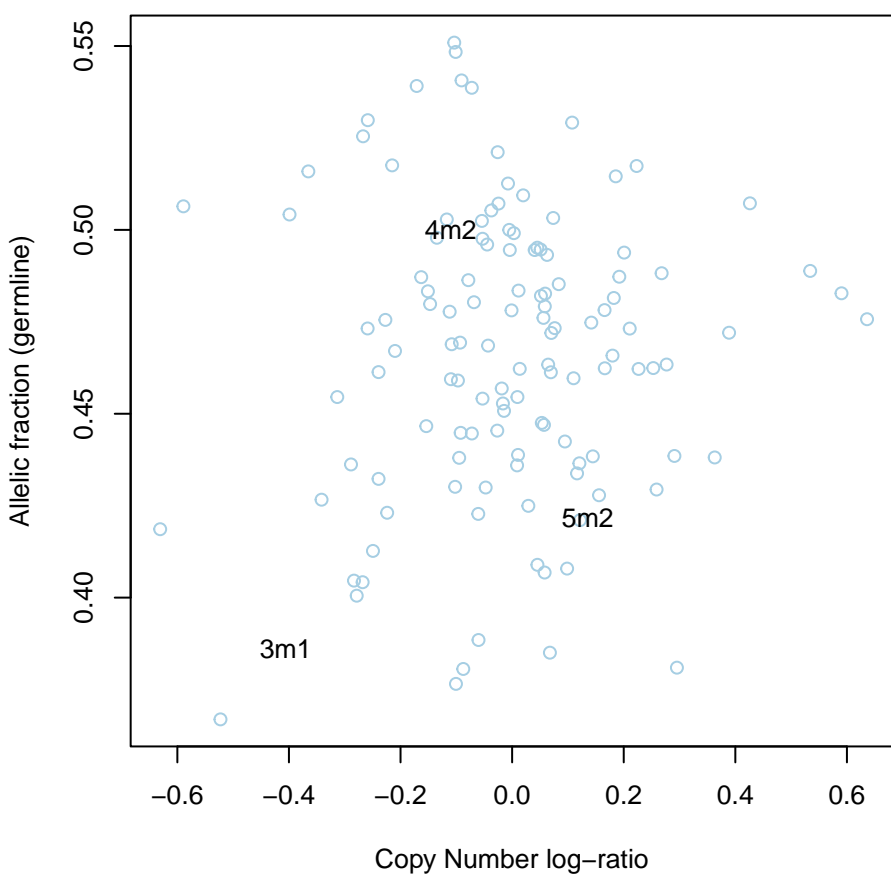


Purity: 0.59 Tumor ploidy: 4.426 SNV log-likelihood: -115.01 GoF: 83.6% Mean coverage: 464,227

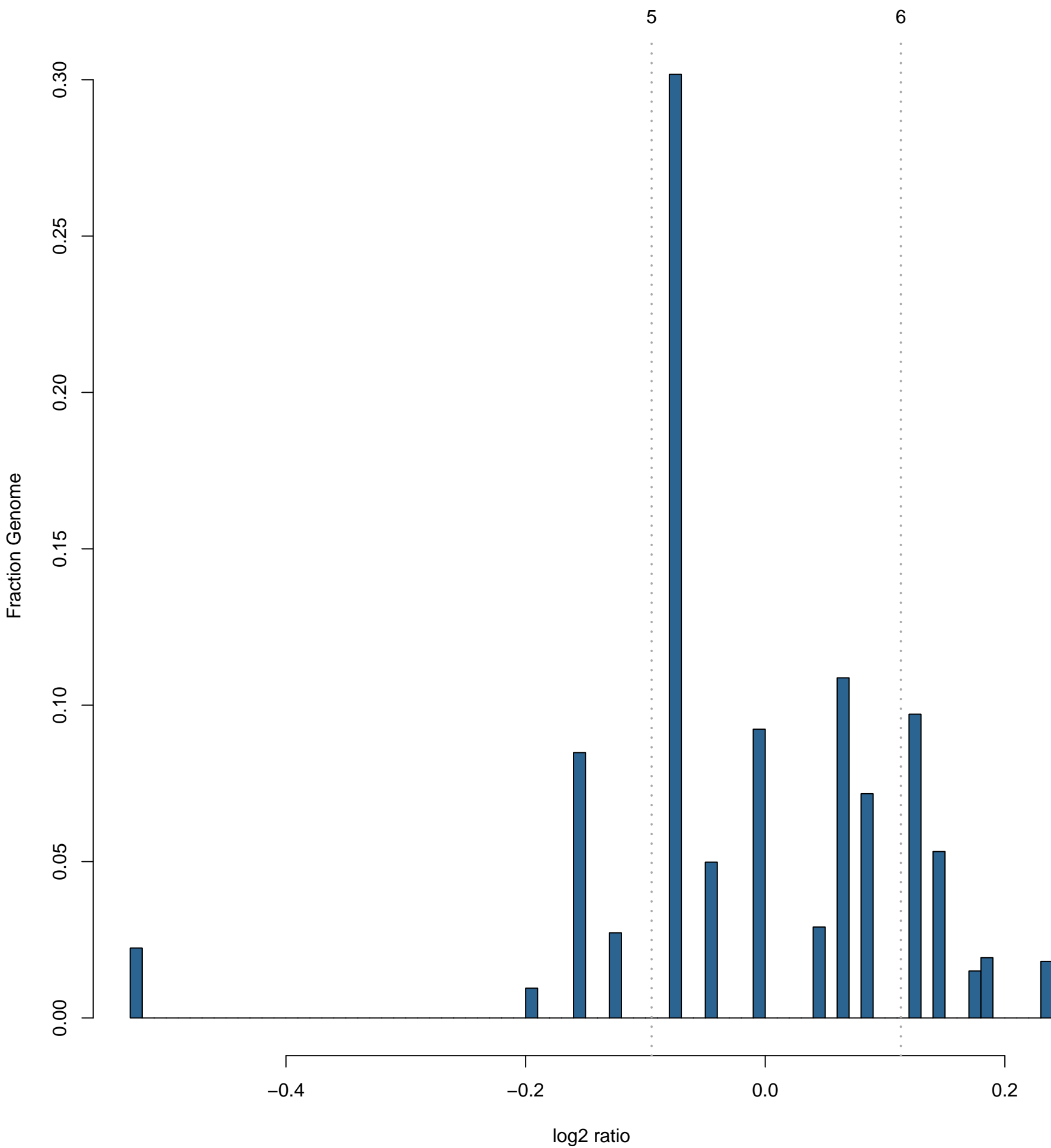


SCNA-fit log-likelihood: -12527.1

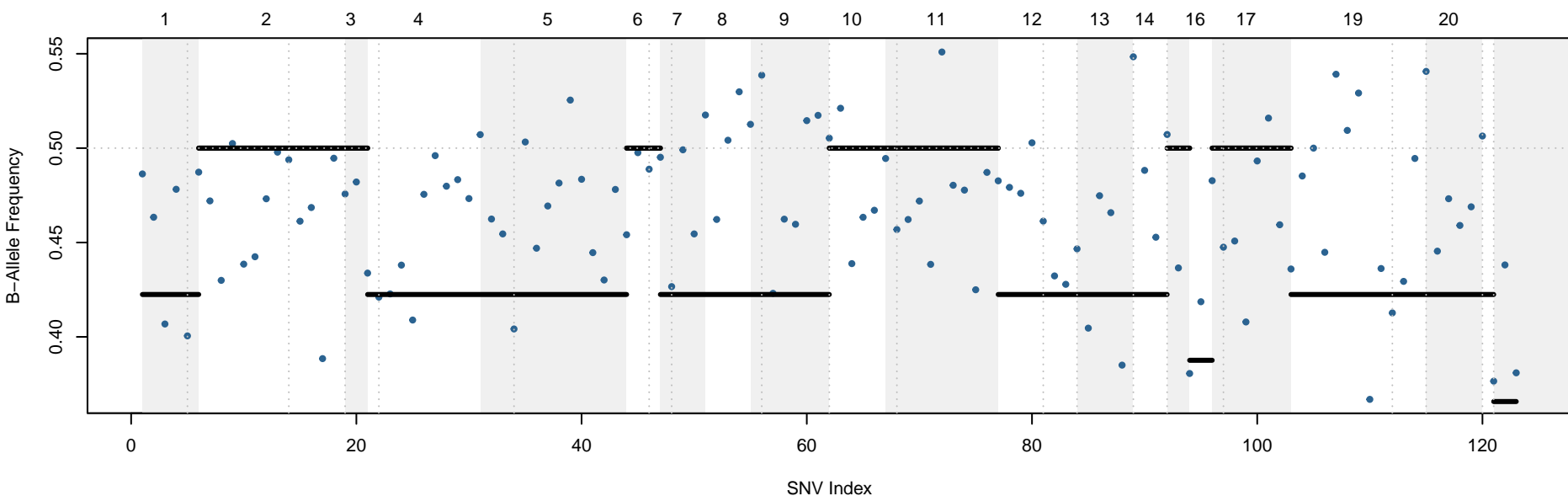




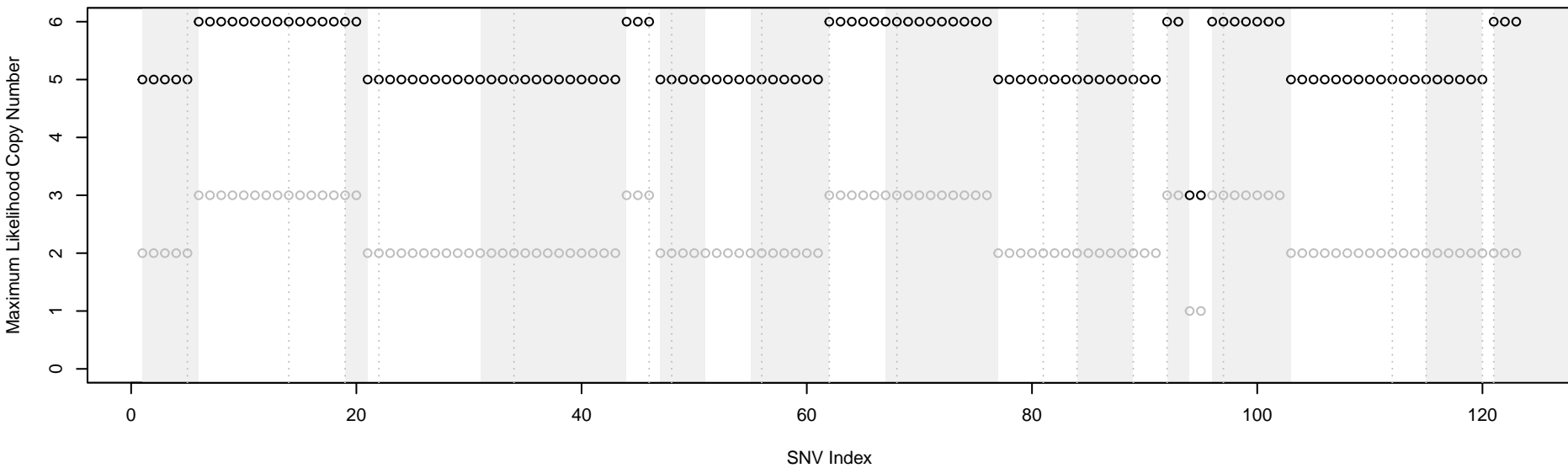
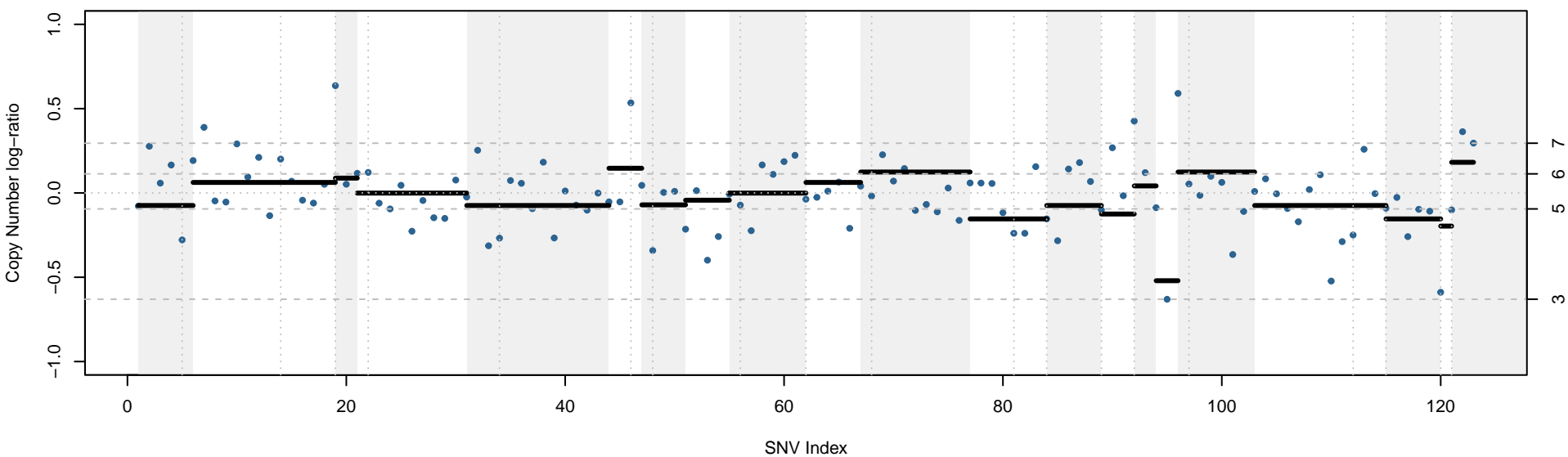
Purity: 0.58 Tumor ploidy: 5.438



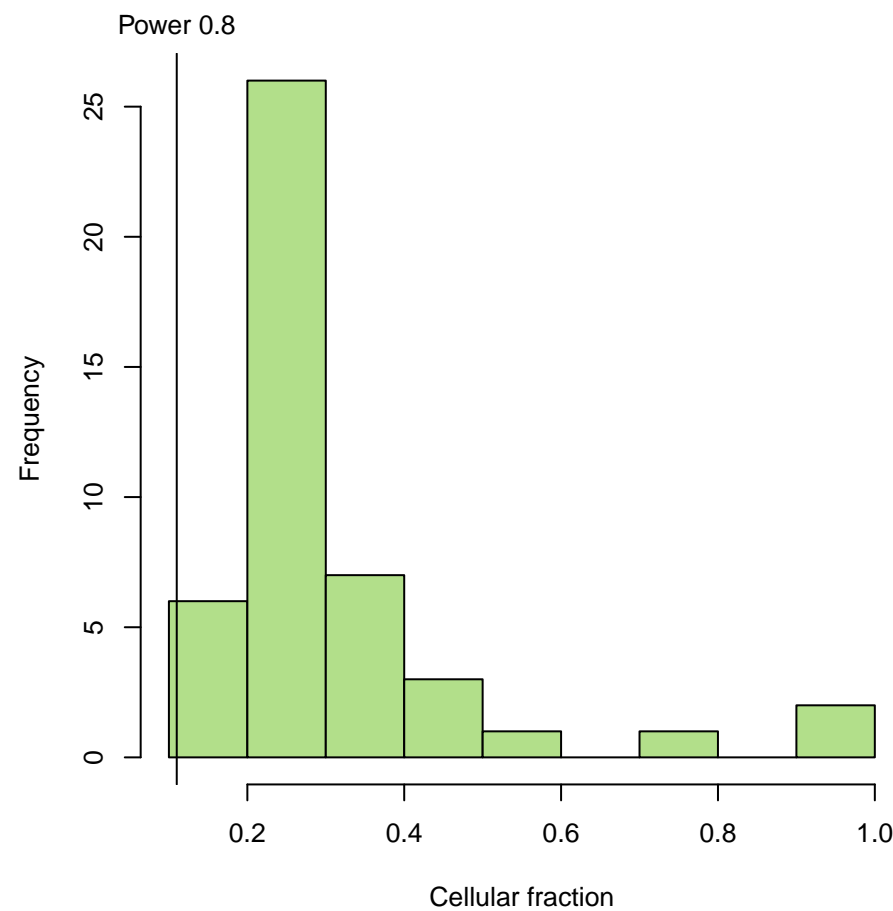
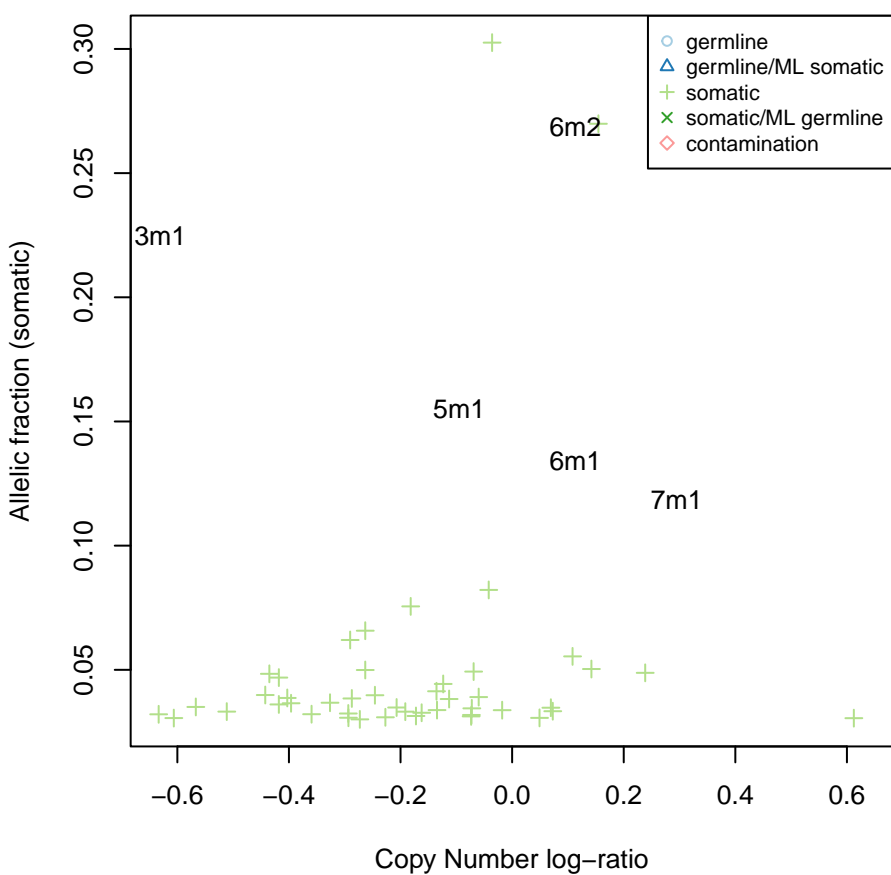
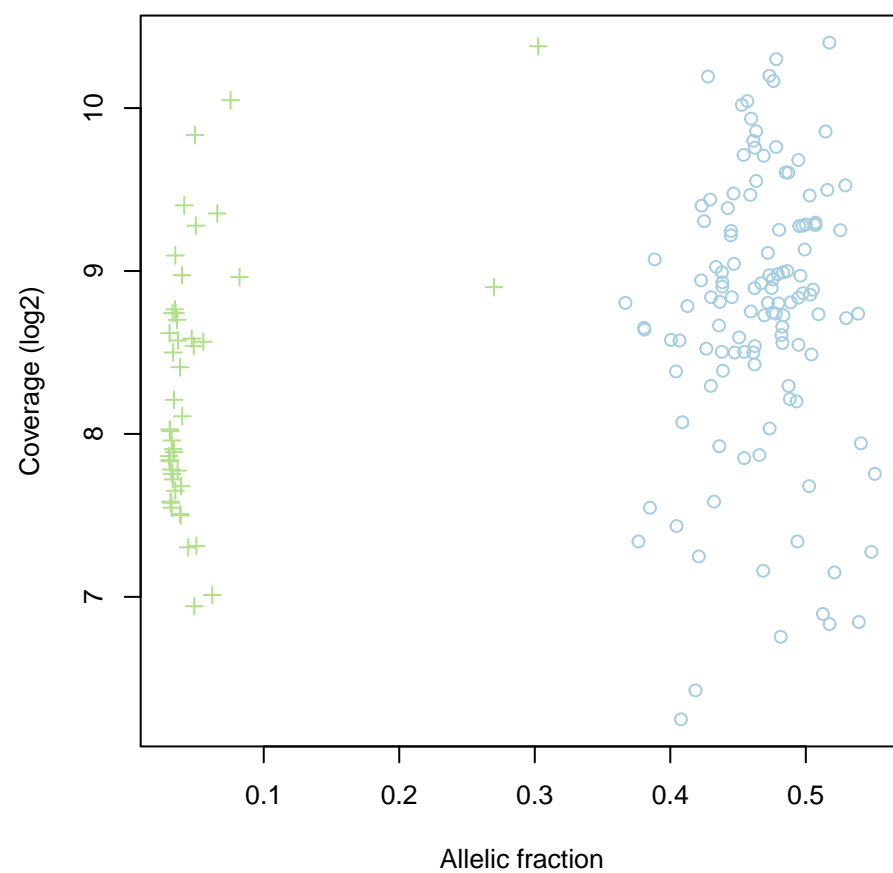
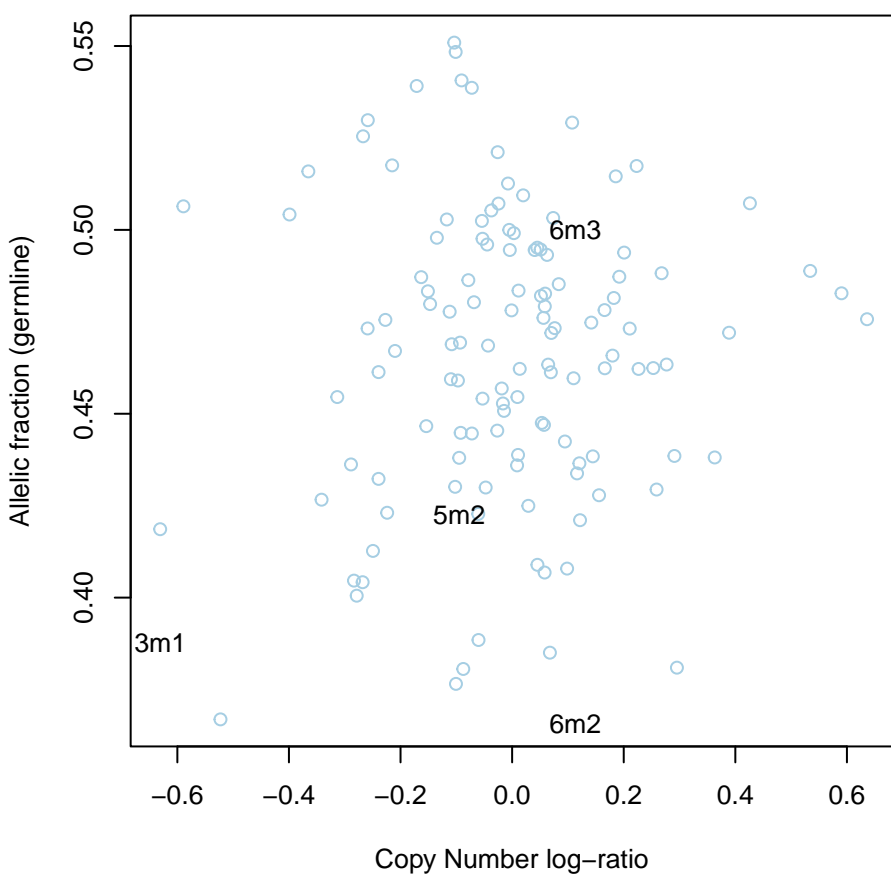
Purity: 0.58 Tumor ploidy: 5.438 SNV log-likelihood: -124.13 GoF: 88.3% Mean coverage: 464,227



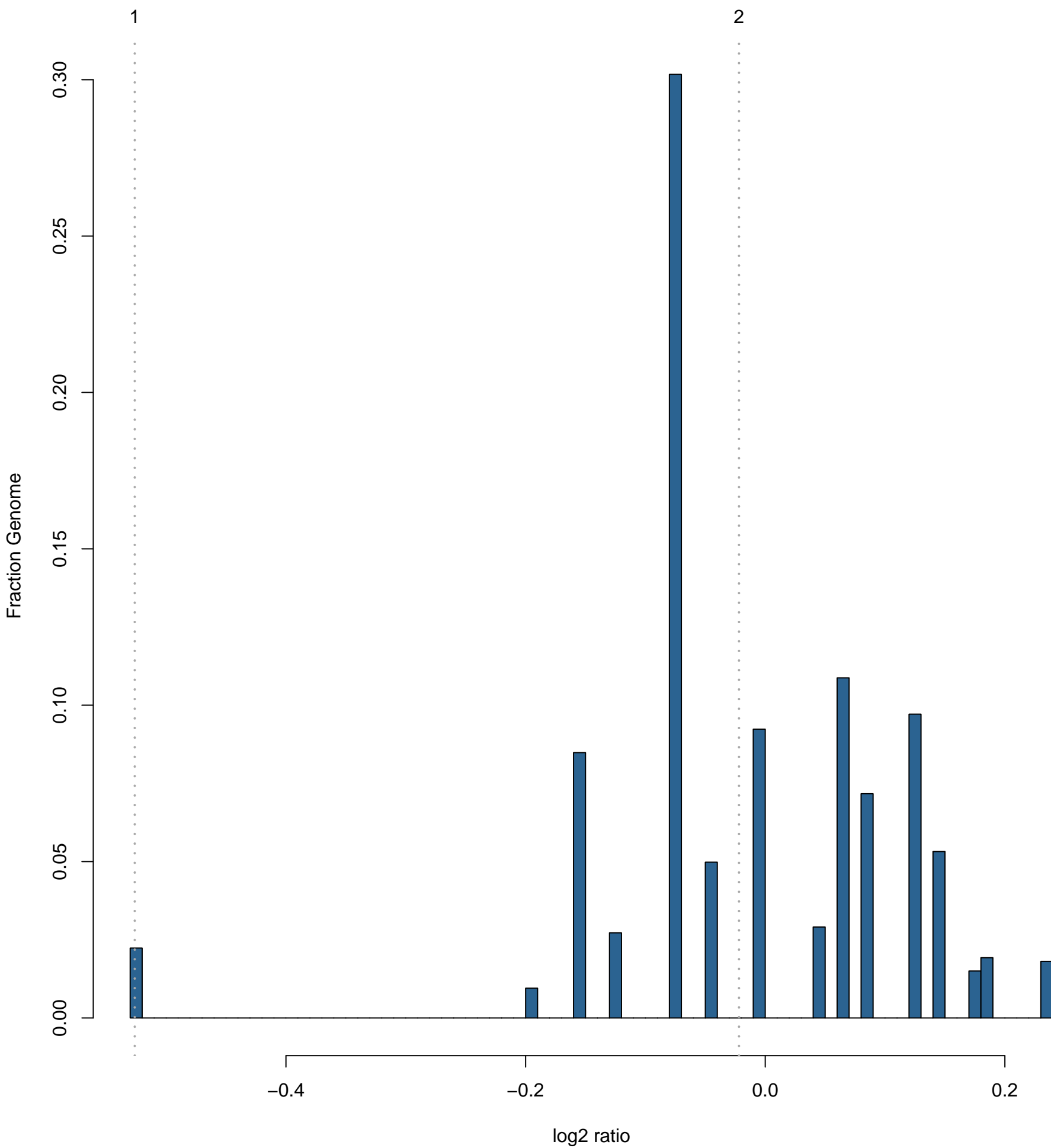
SCNA-fit log-likelihood: -12511.55



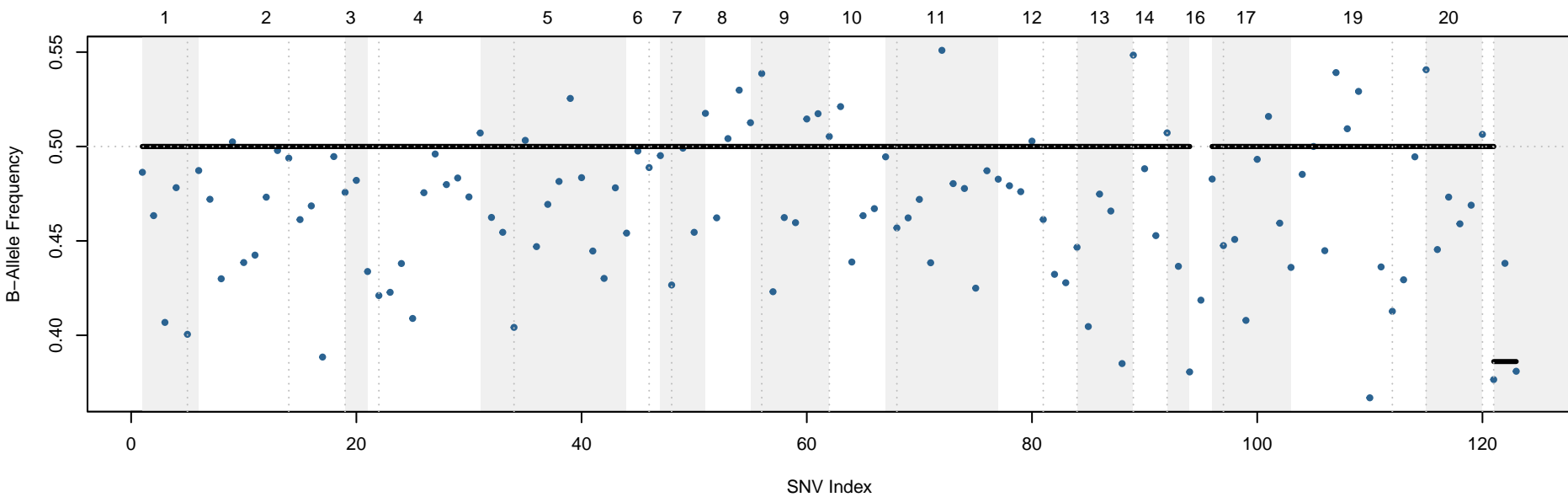




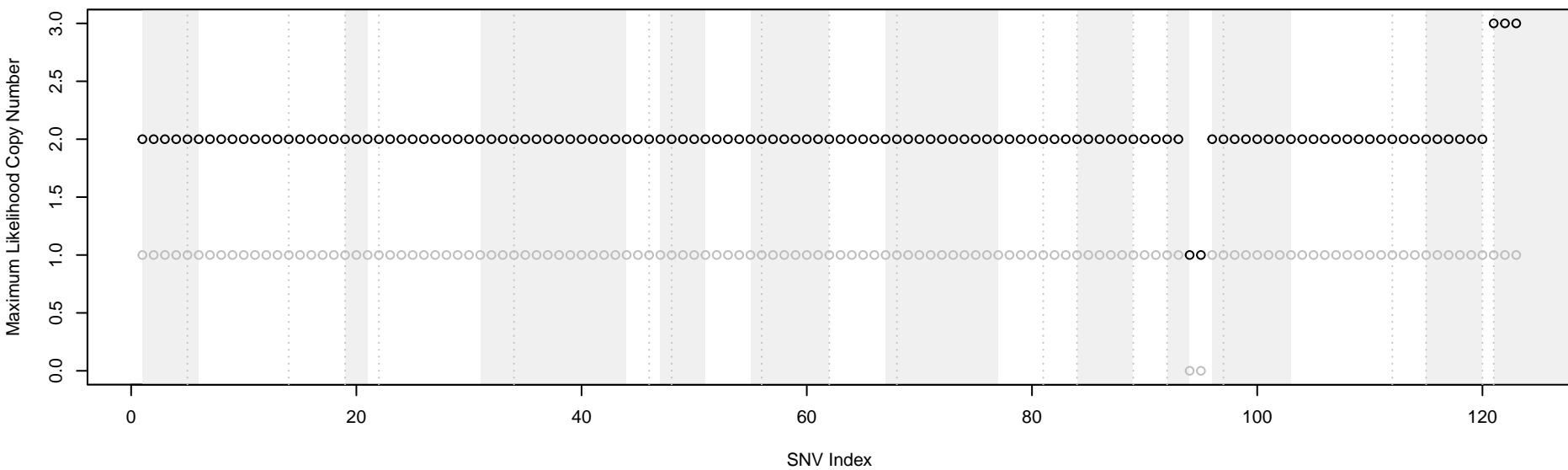
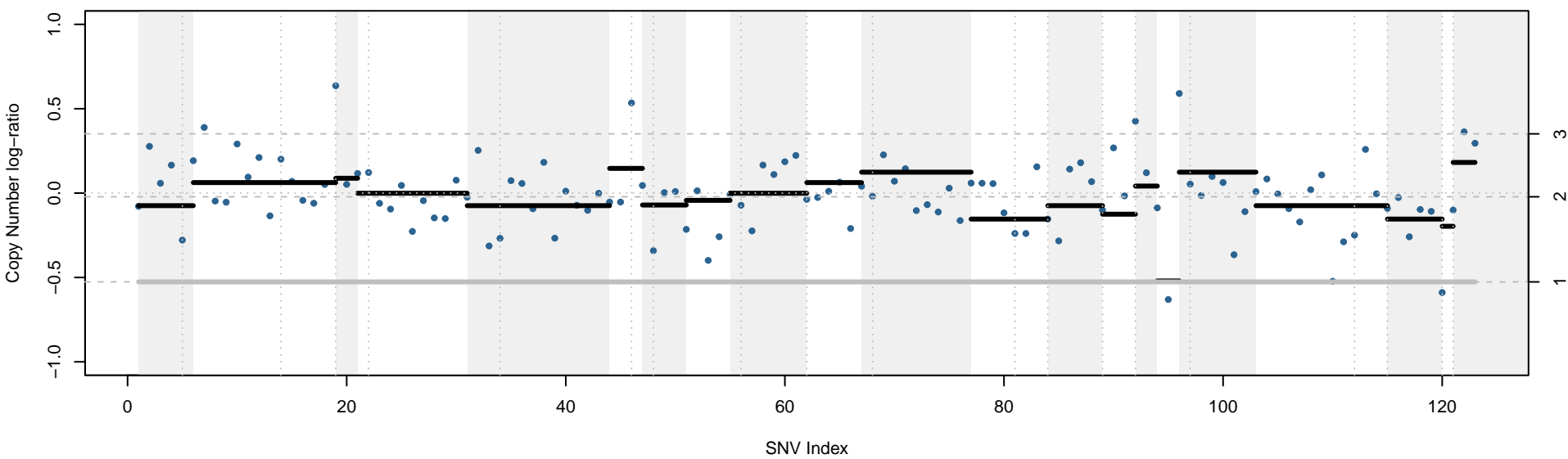
Purity: 0.59 Tumor ploidy: 2.052

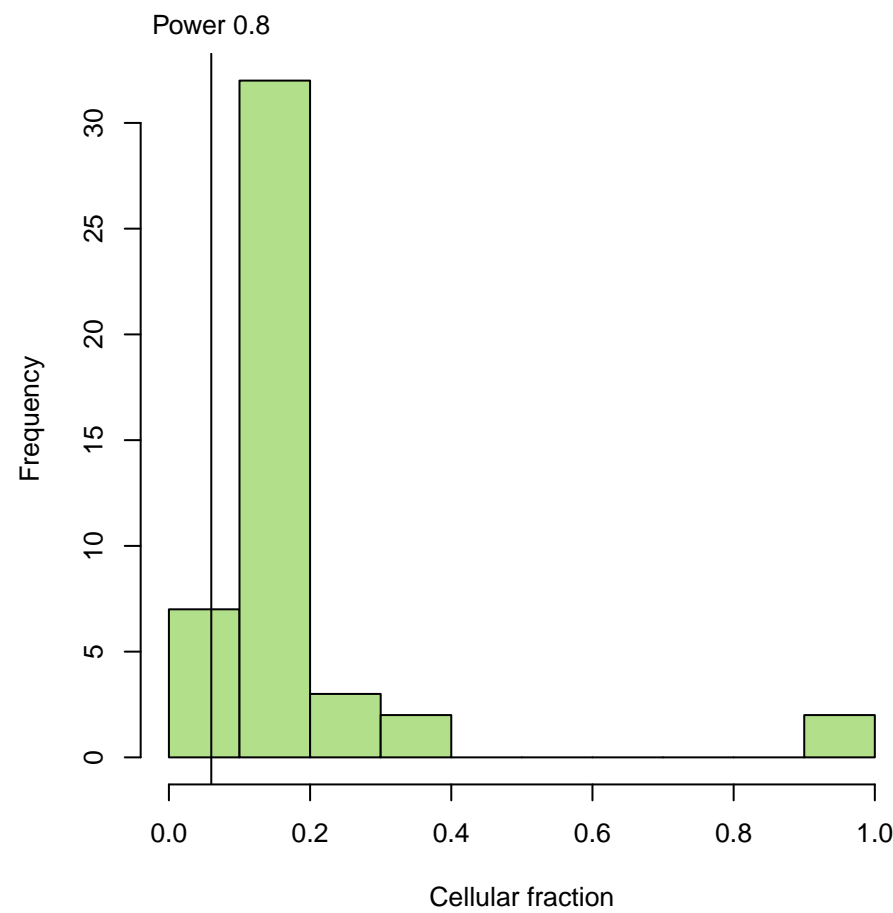
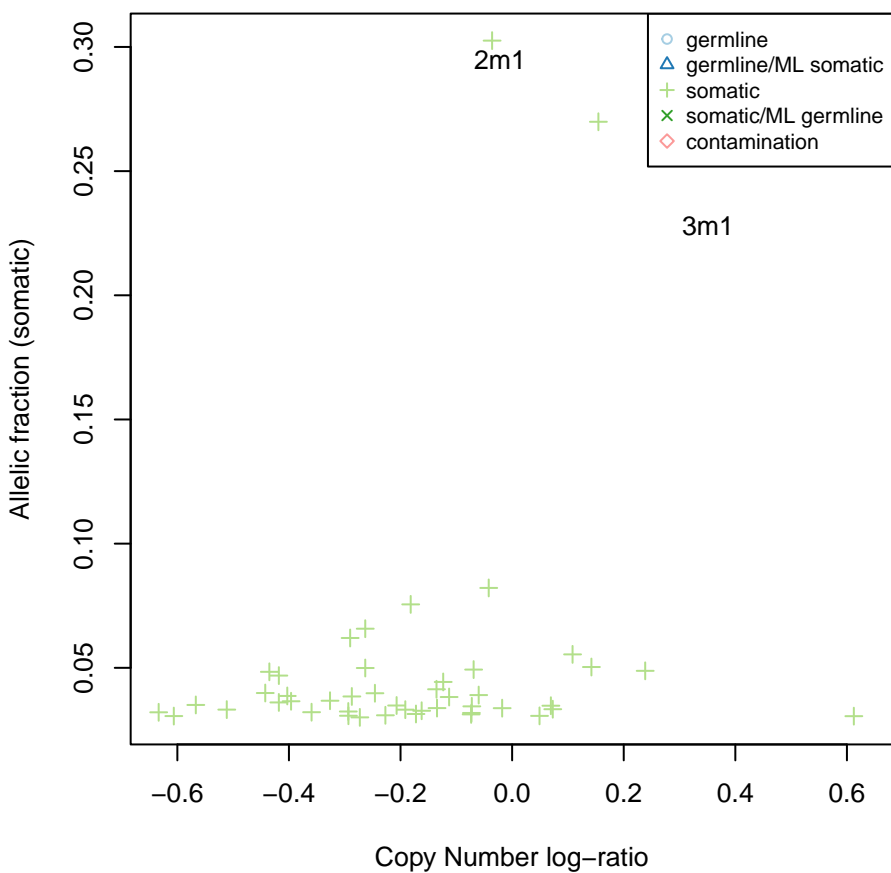
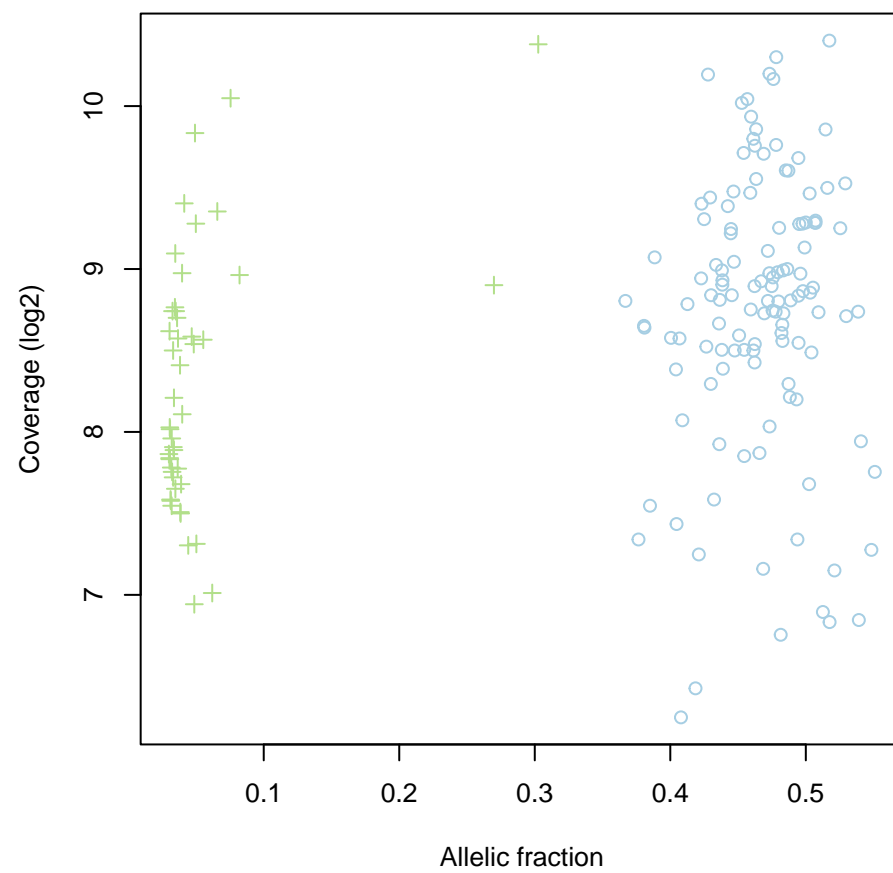
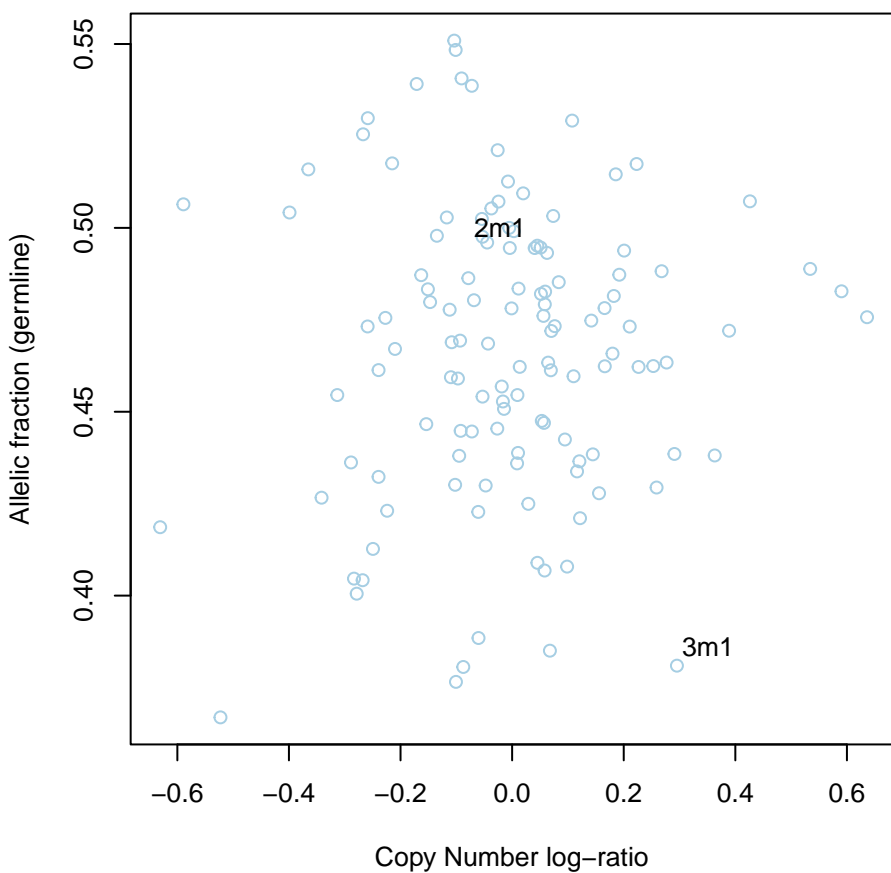


Purity: 0.59 Tumor ploidy: 2.052 SNV log-likelihood: -130.44 GoF: 55% Mean coverage: 464;227

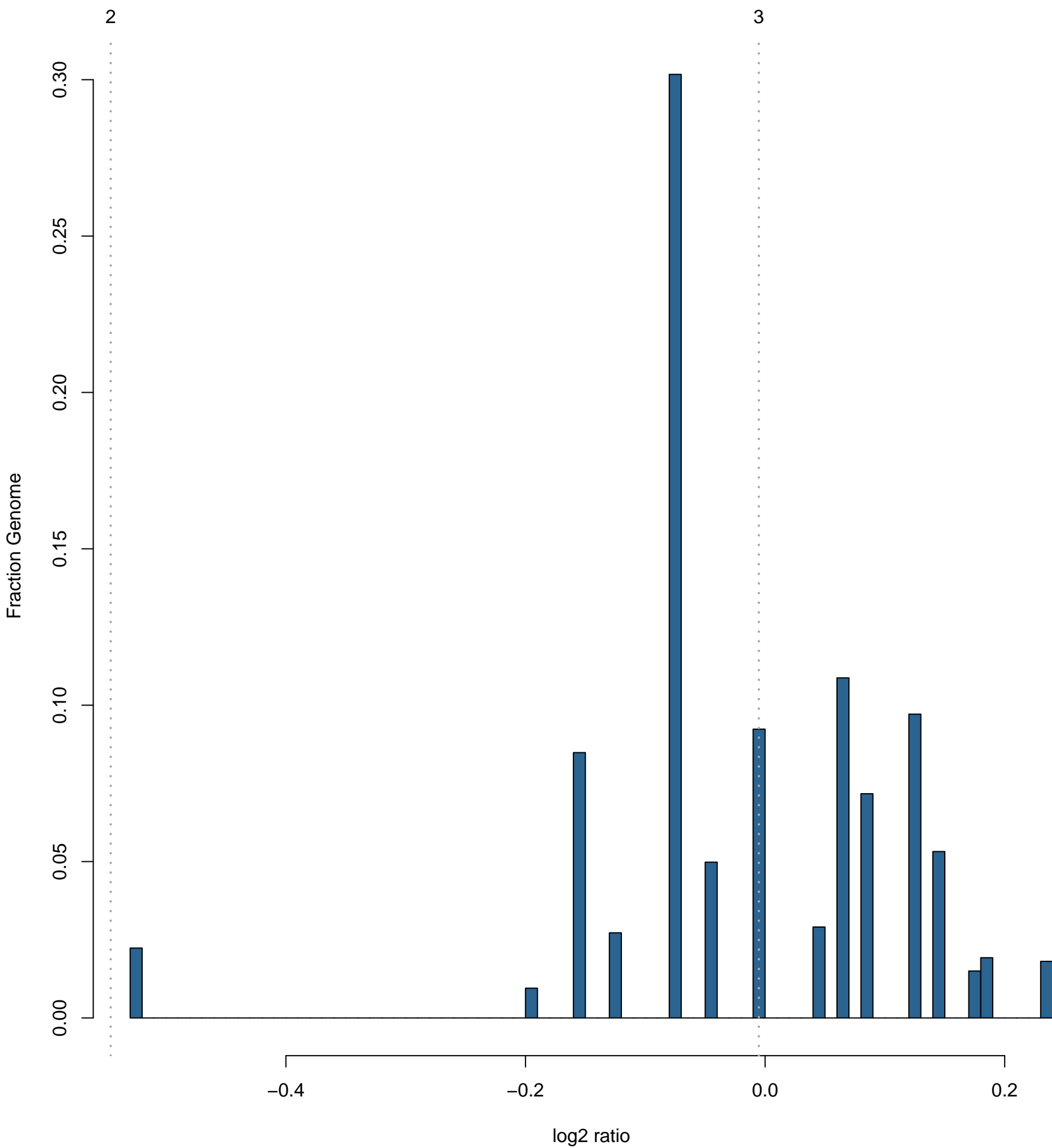


SCNA-fit log-likelihood: -12620.27

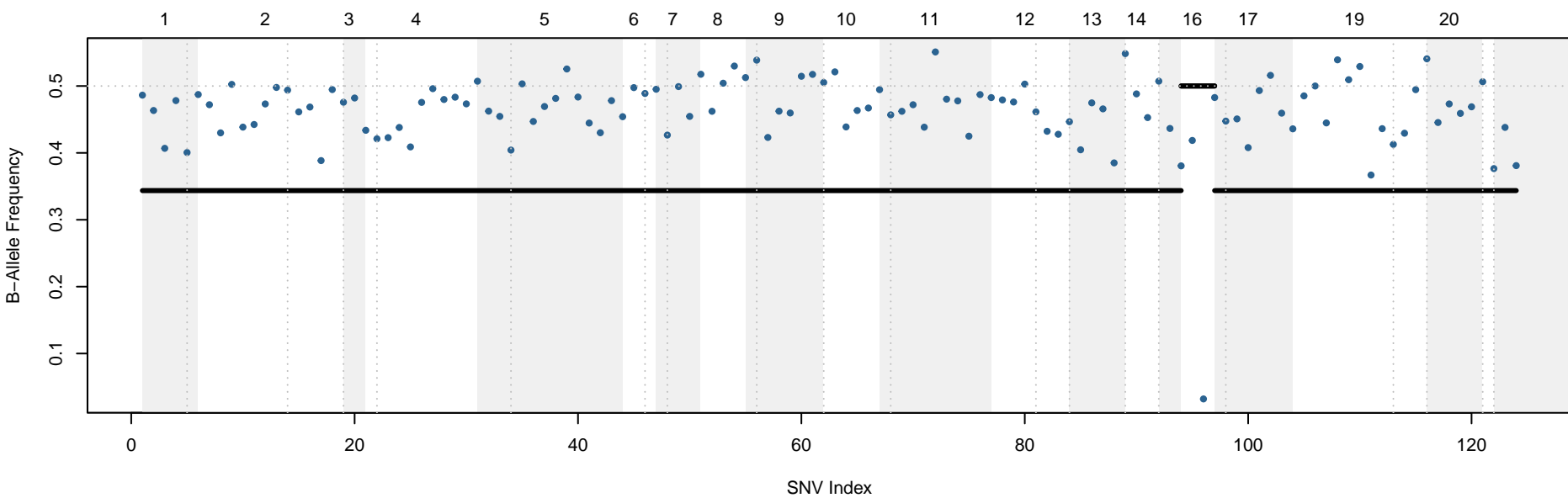




Purity: 0.91 Tumor ploidy: 3.012



Purity: 0.91 Tumor ploidy: 3.012 SNV log-likelihood: -1320.42 GoF: 25.3% Mean coverage: 464;227



SCNA-fit log-likelihood: -12624.35

