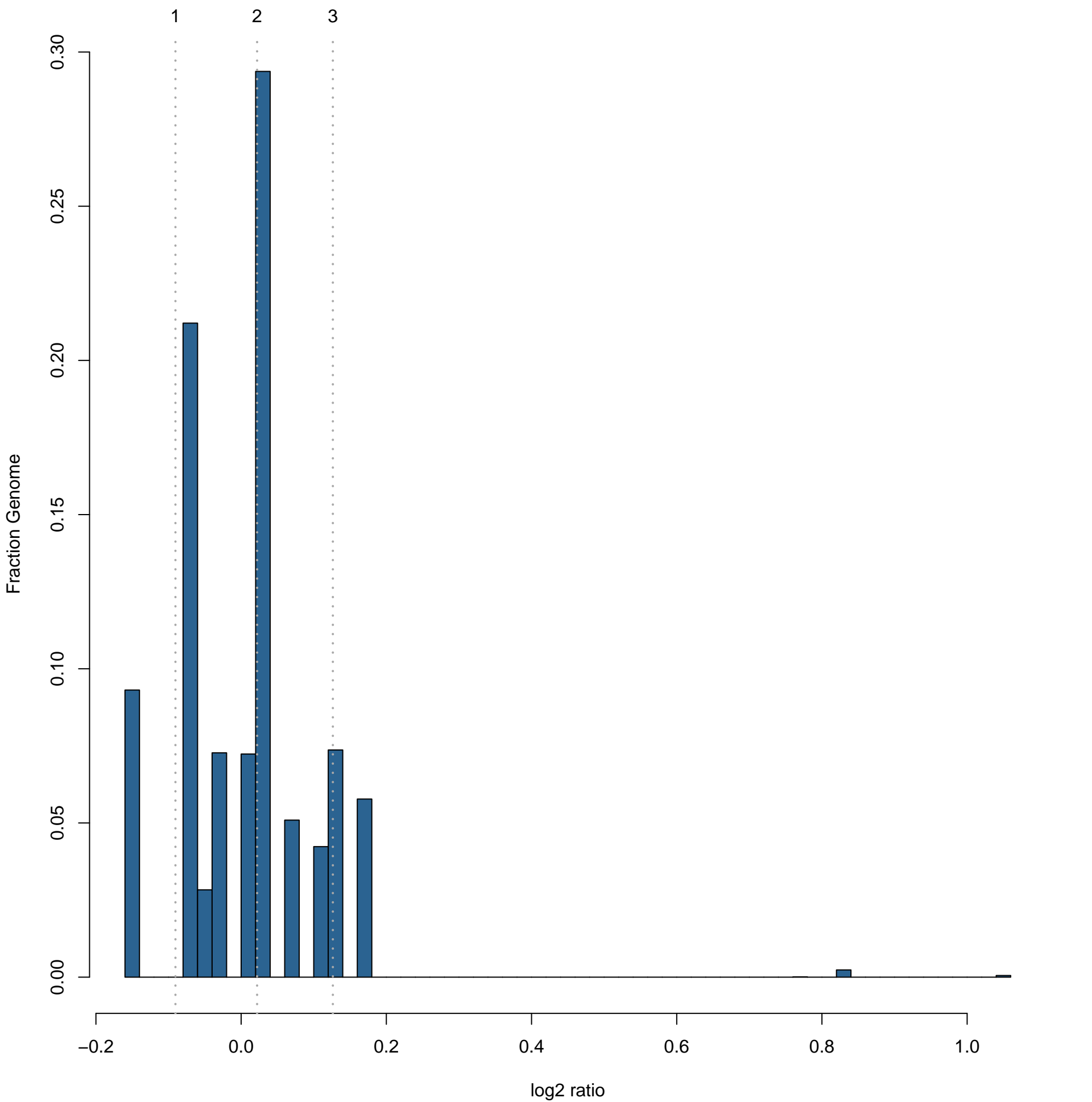
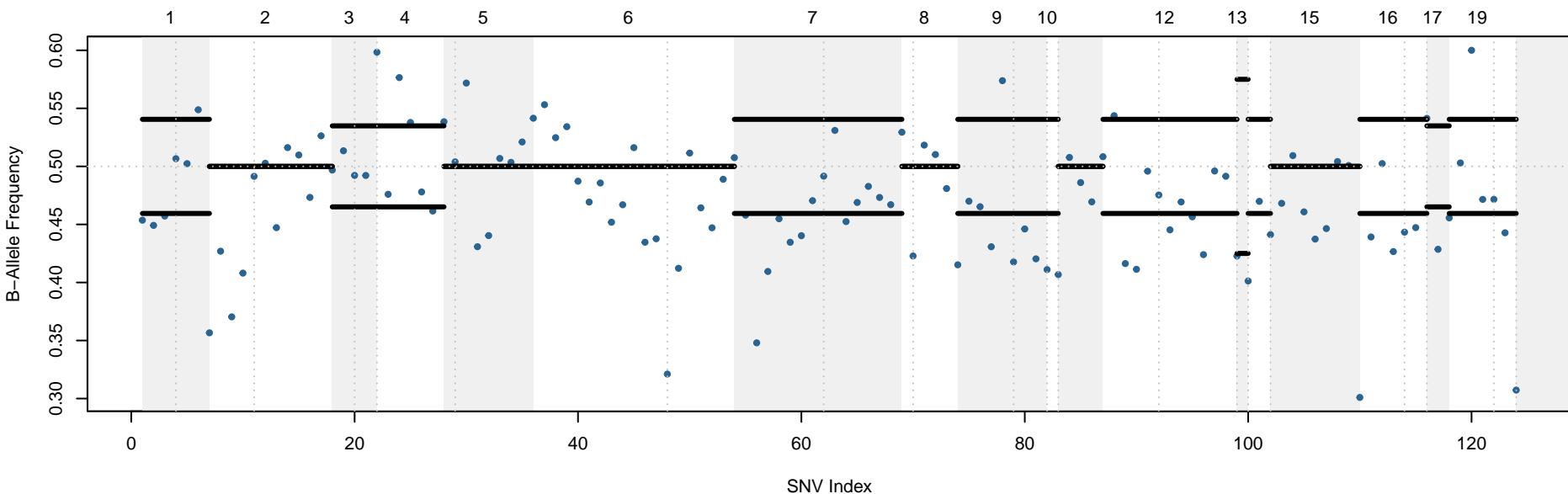


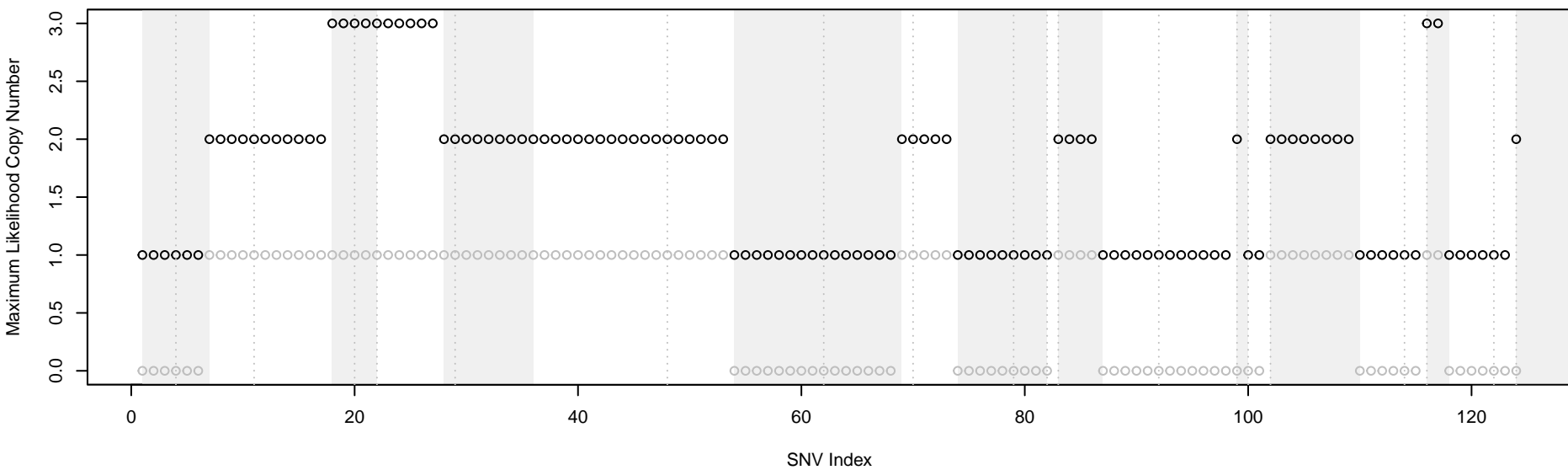
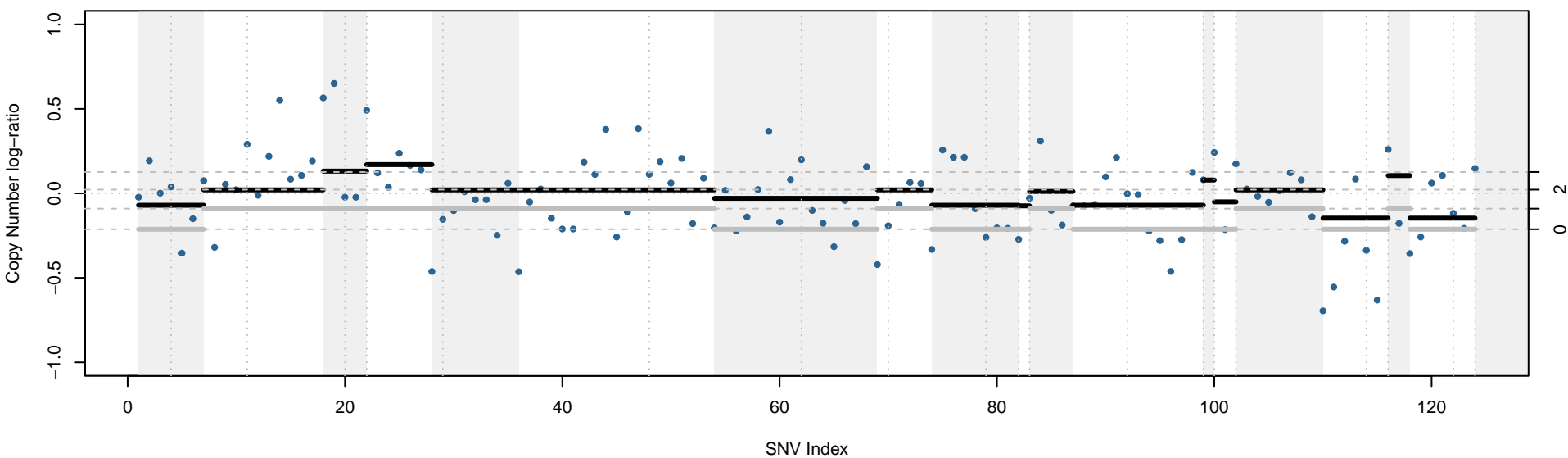
Purity: 0.15 Tumor ploidy: 1.798

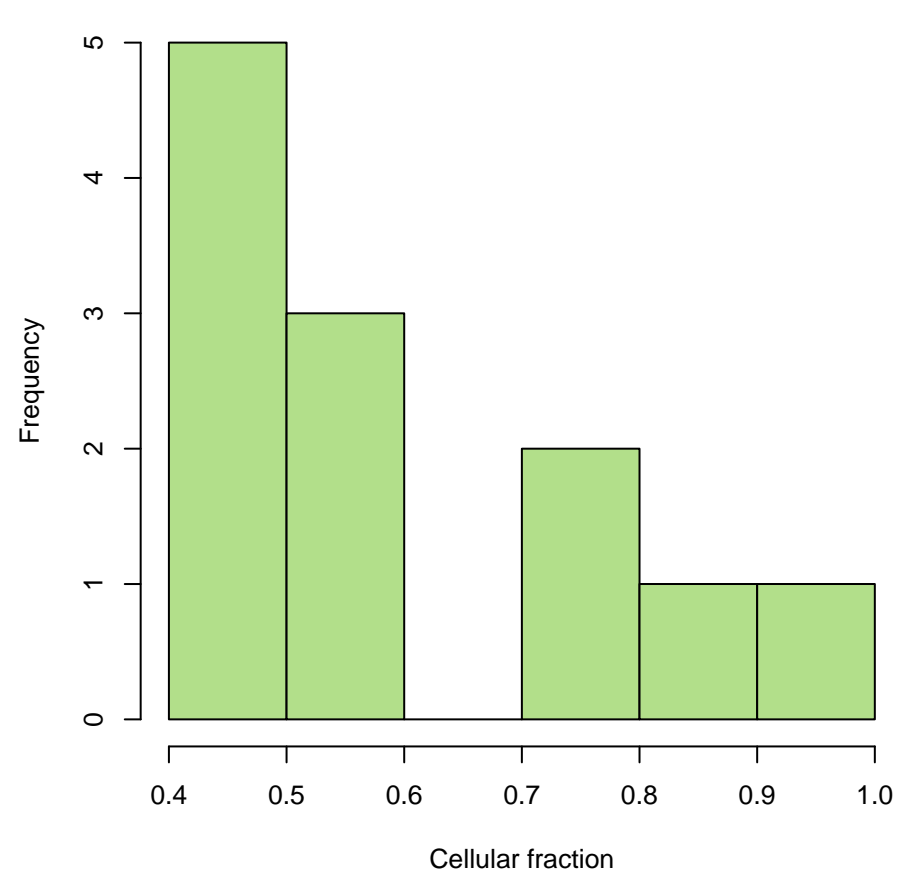
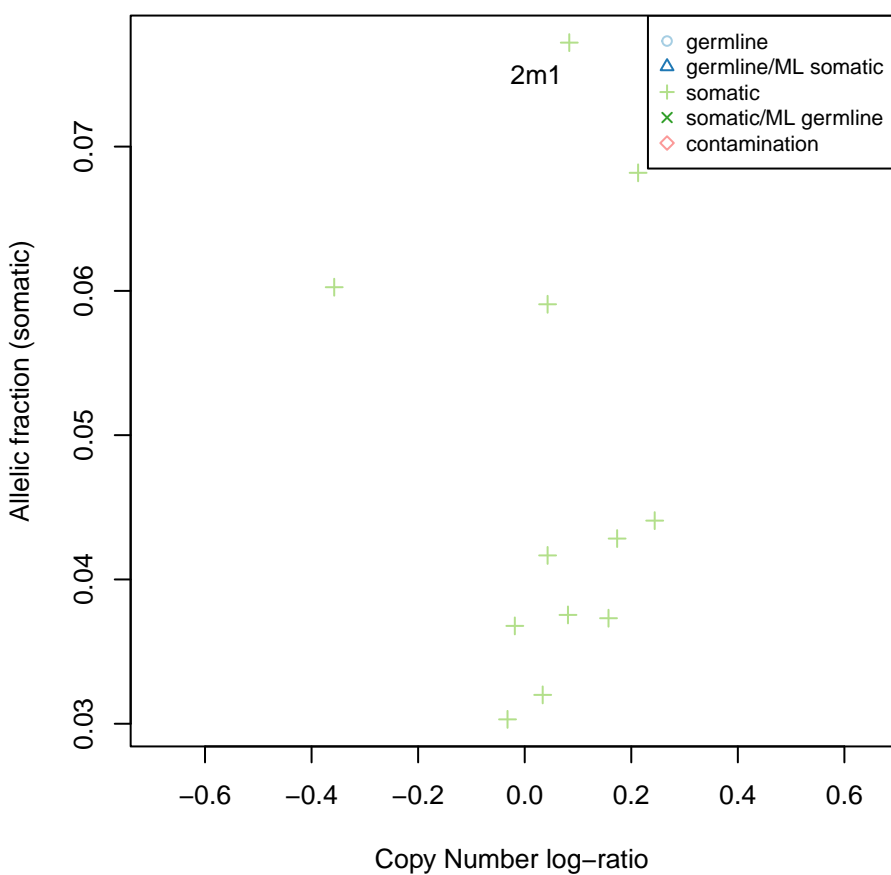
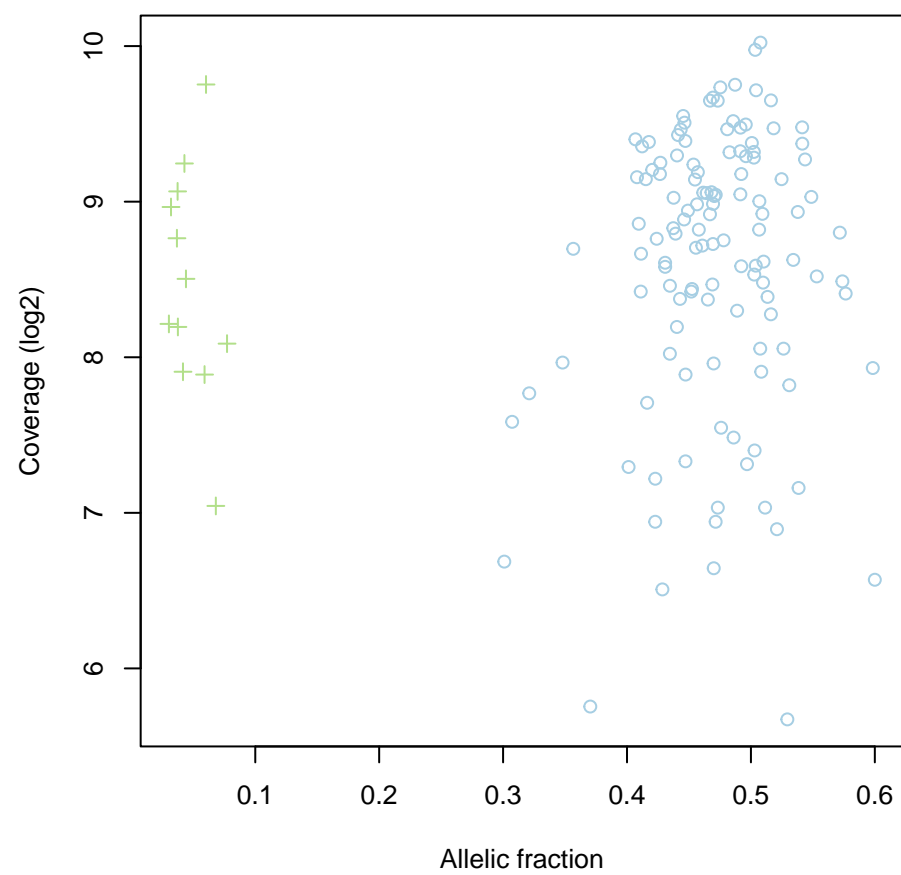
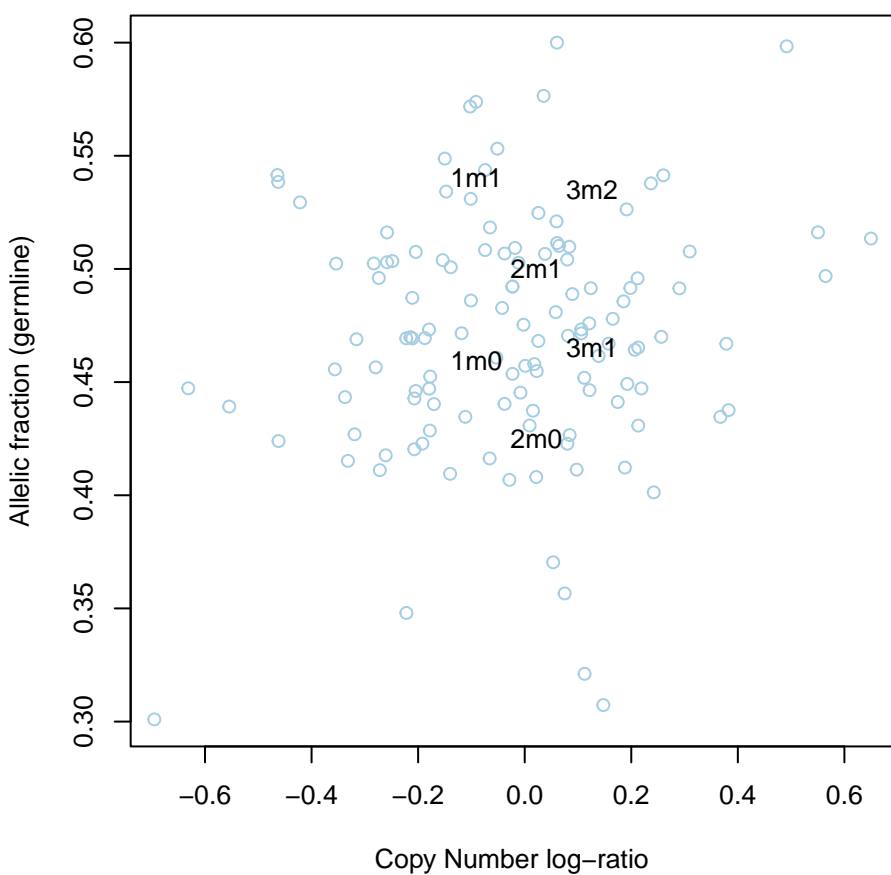


Purity: 0.15 Tumor ploidy: 1.798 SNV log-likelihood: 45.95 GoF: 95.8% Mean coverage: 577;448

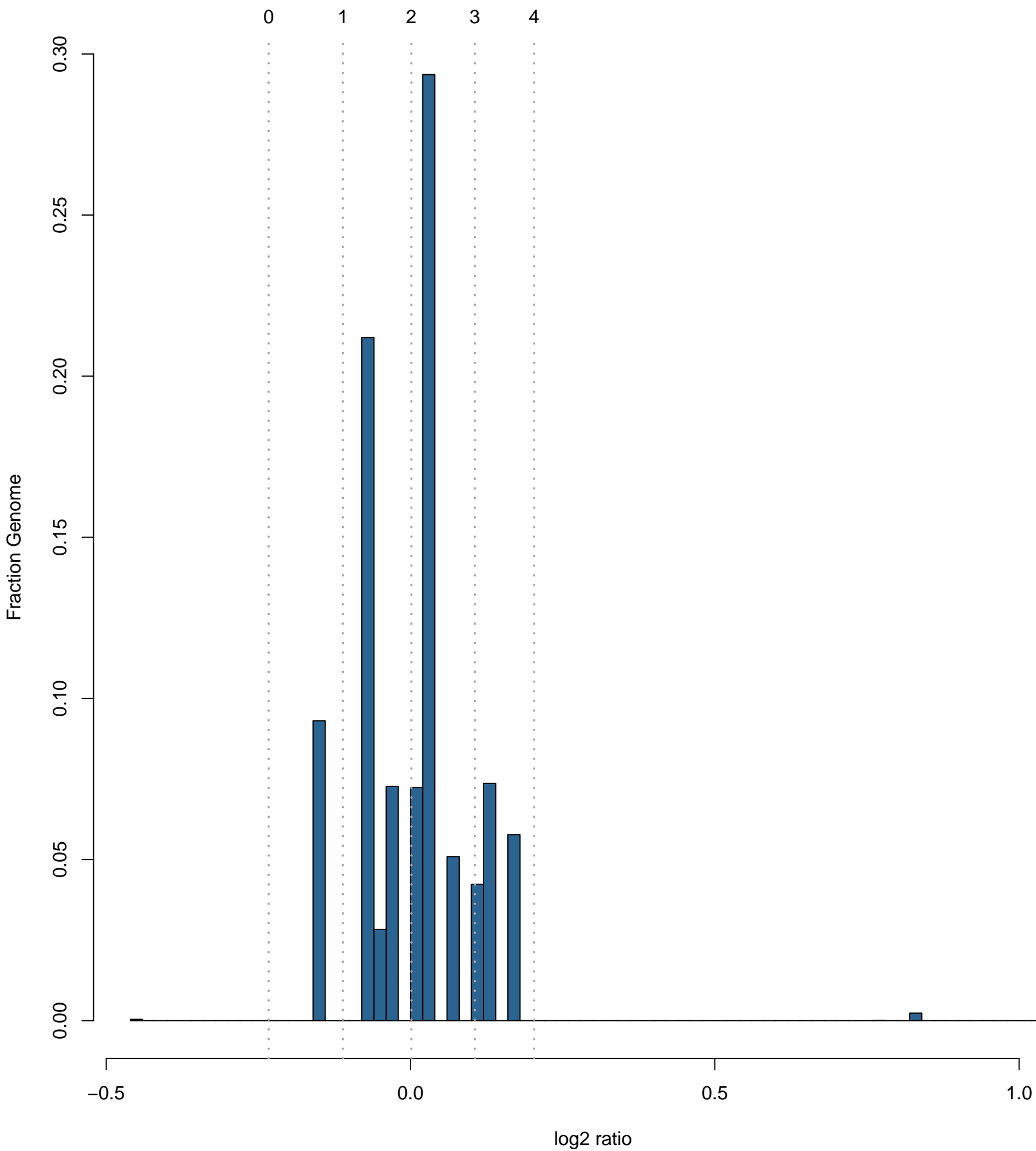


SCNA-fit log-likelihood: -14518.55

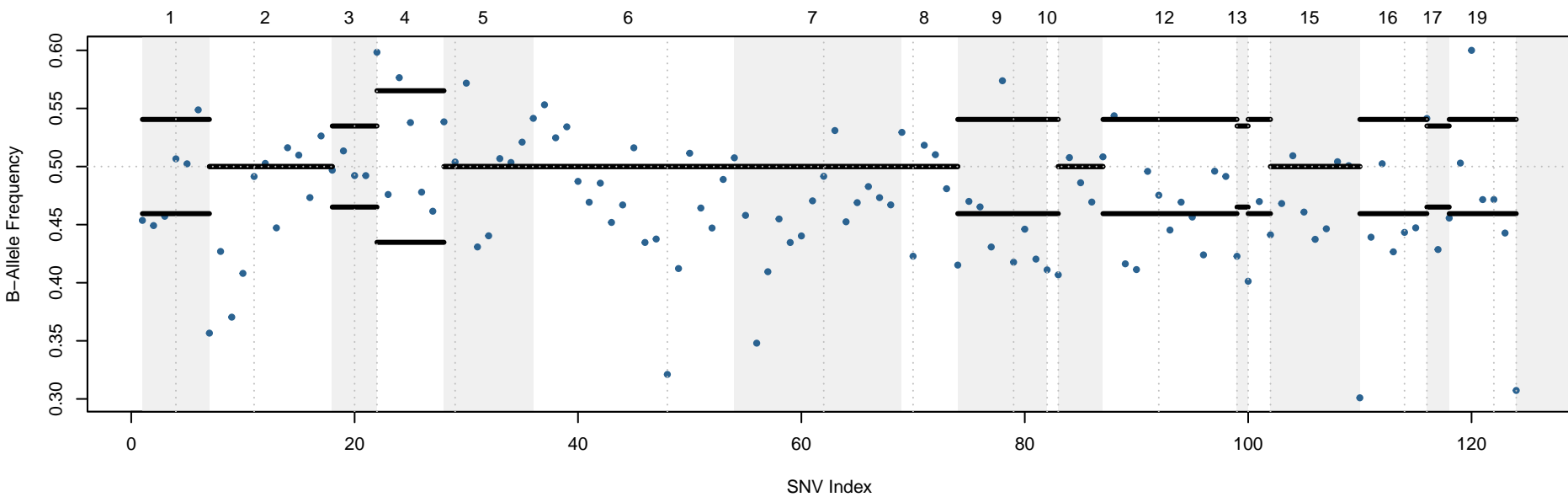




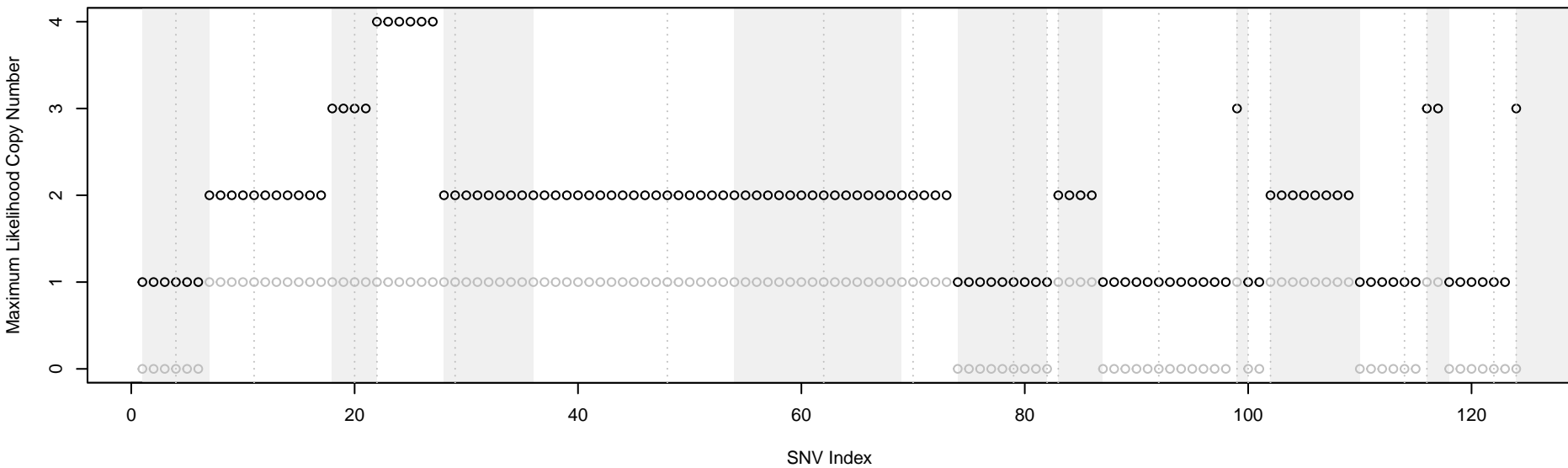
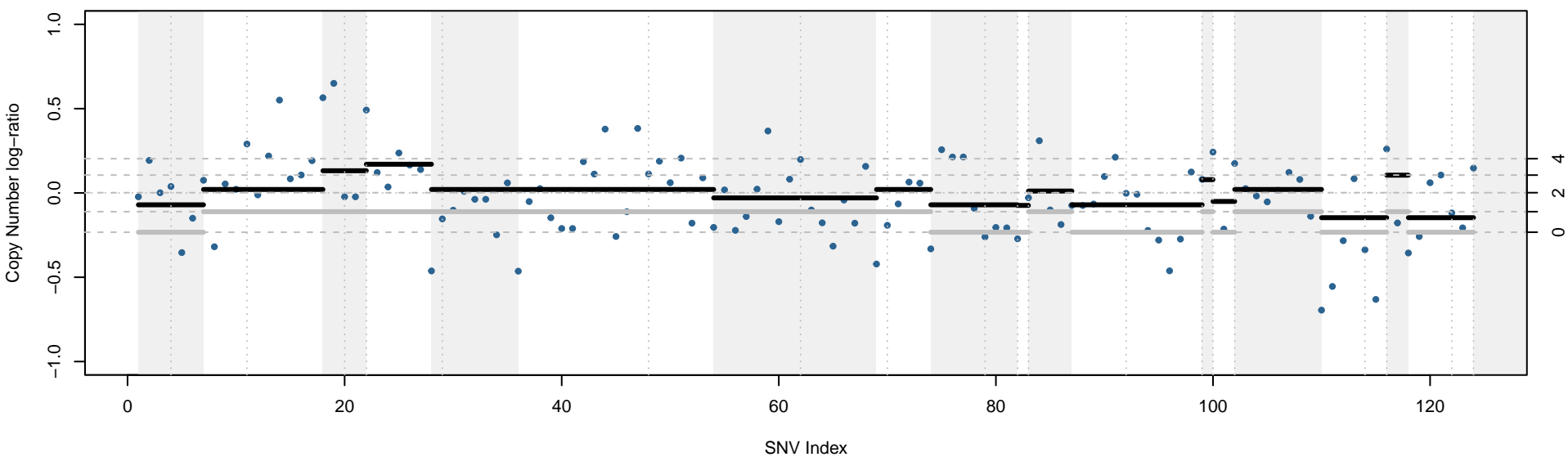
Purity: 0.15 Tumor ploidy: 1.987

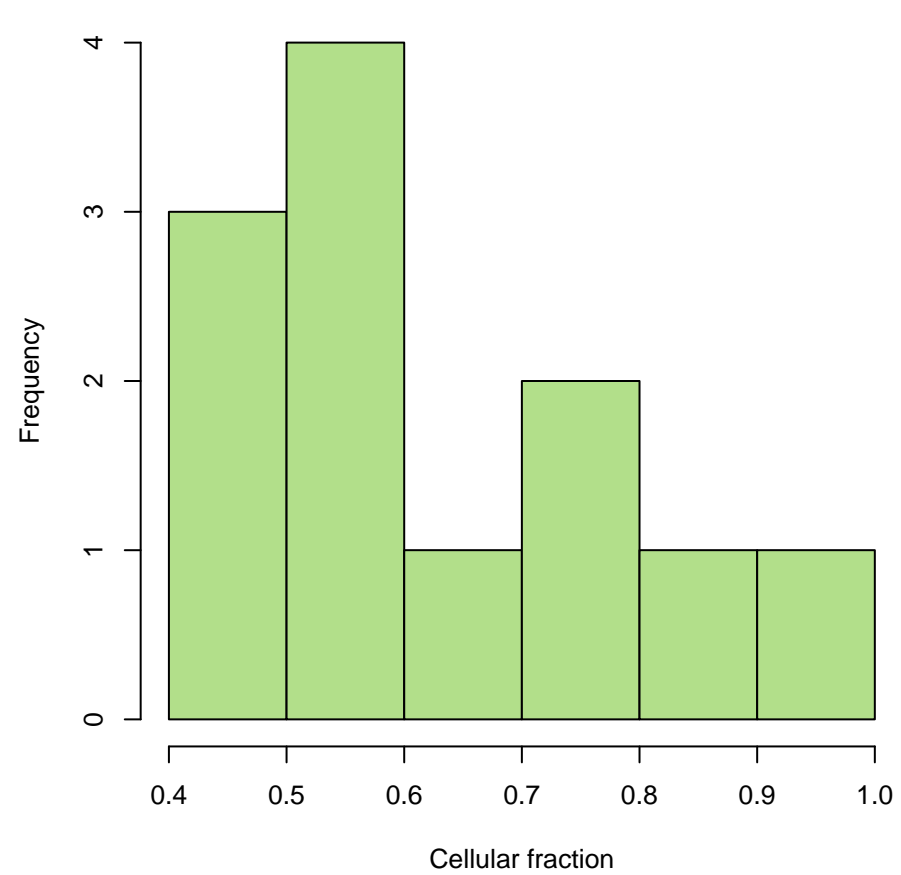
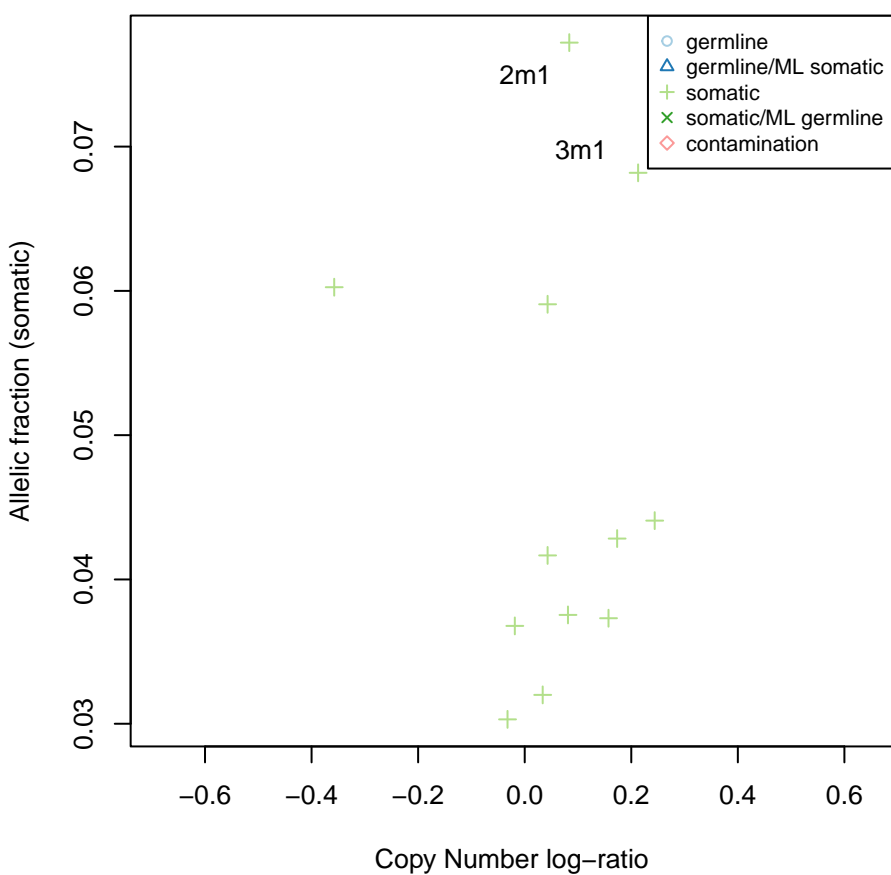
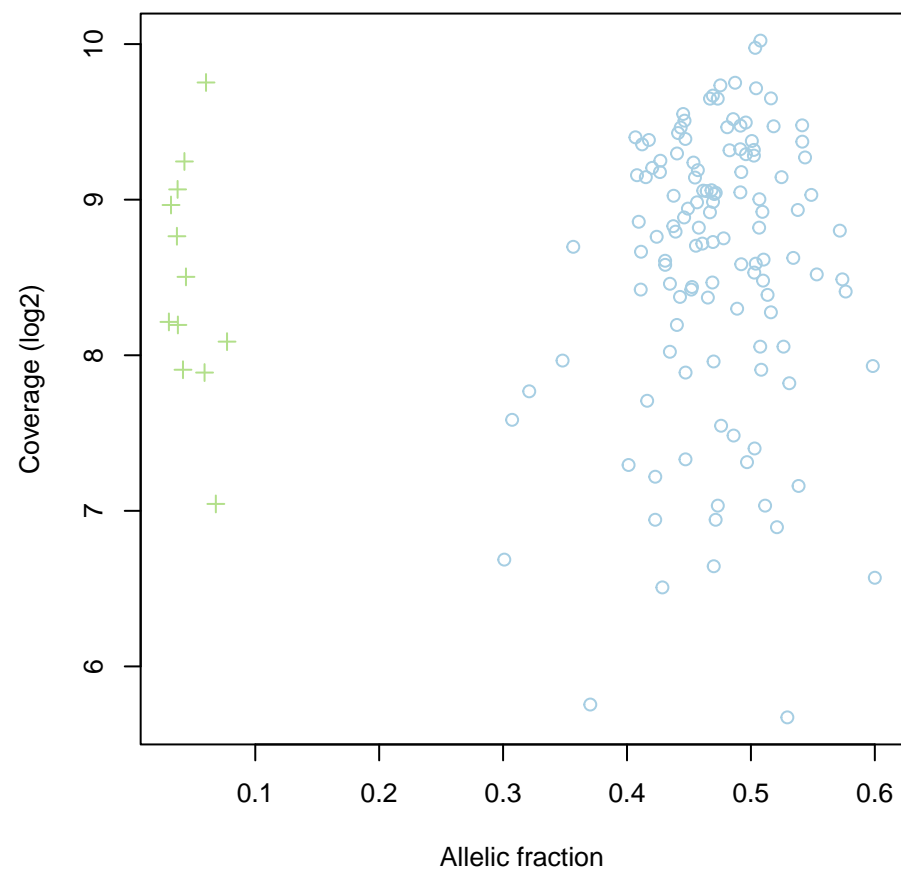
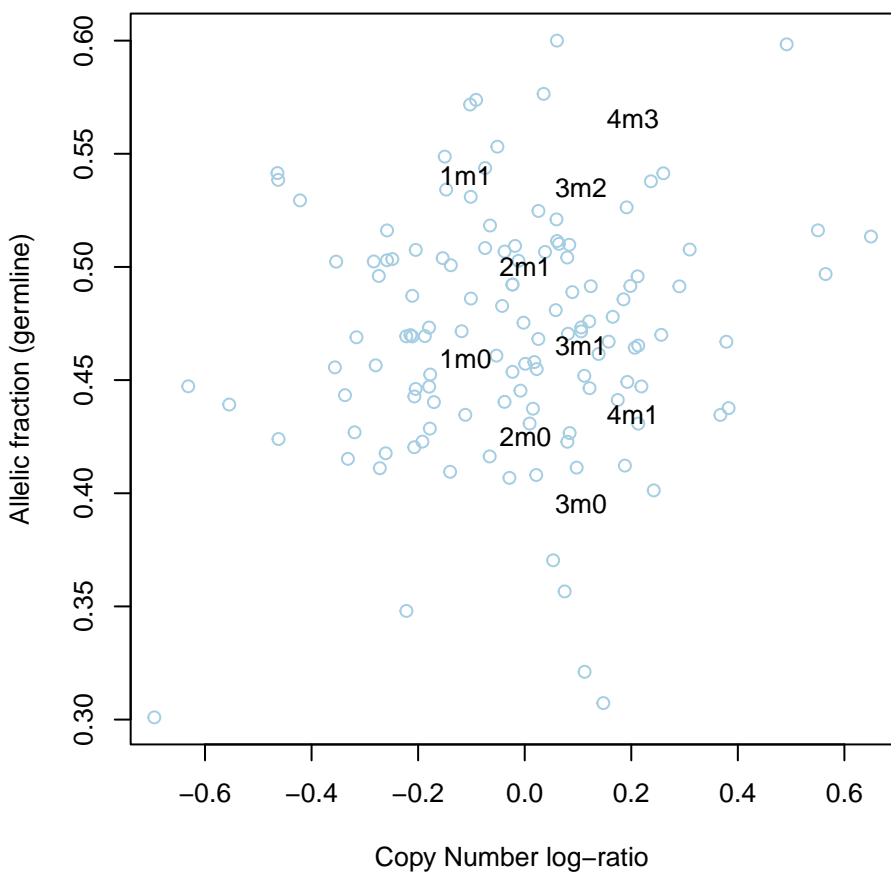


Purity: 0.15 Tumor ploidy: 1.987 SNV log-likelihood: 35.86 GoF: 95.8% Mean coverage: 577;448

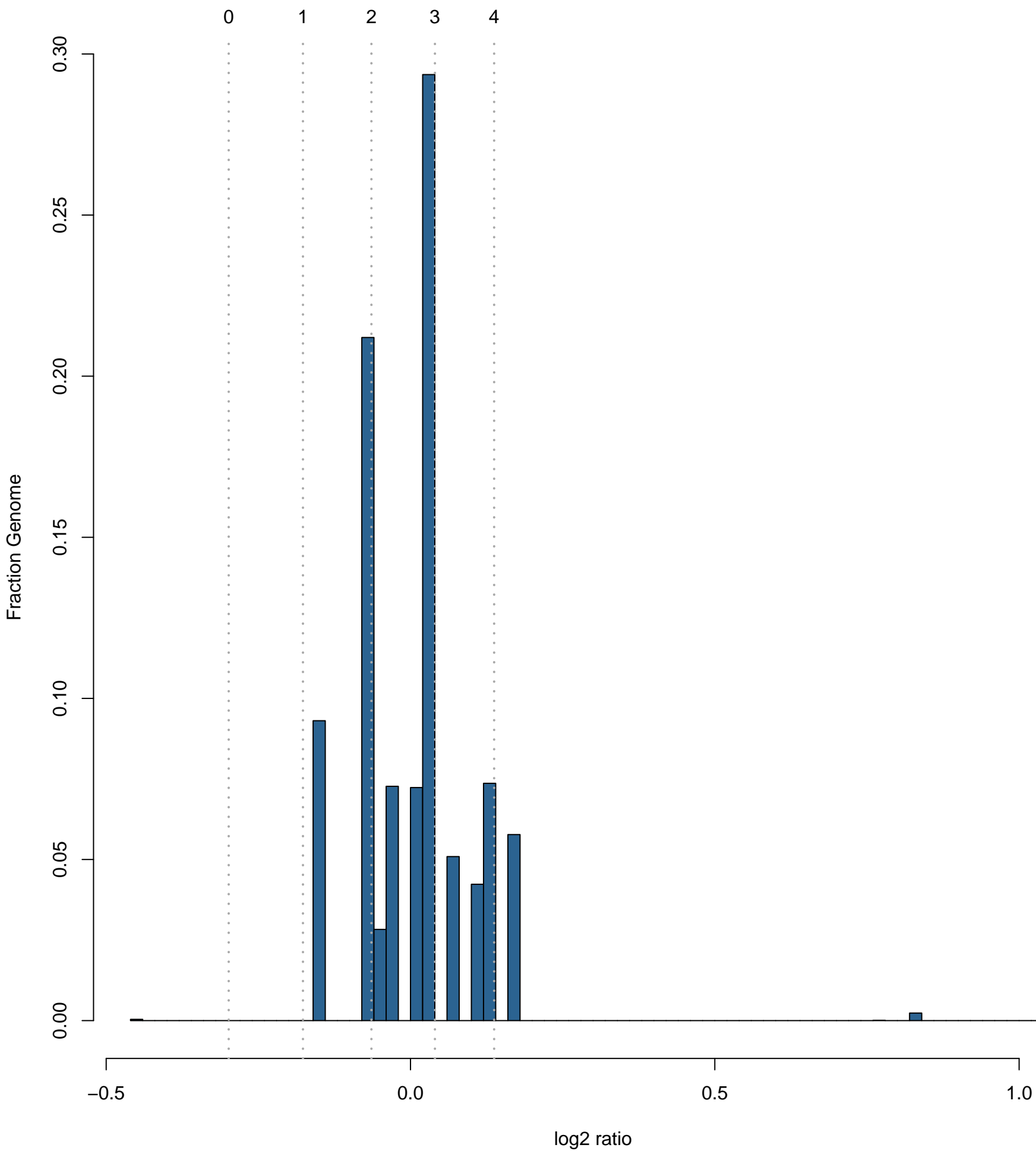


SCNA-fit log-likelihood: -14515.05

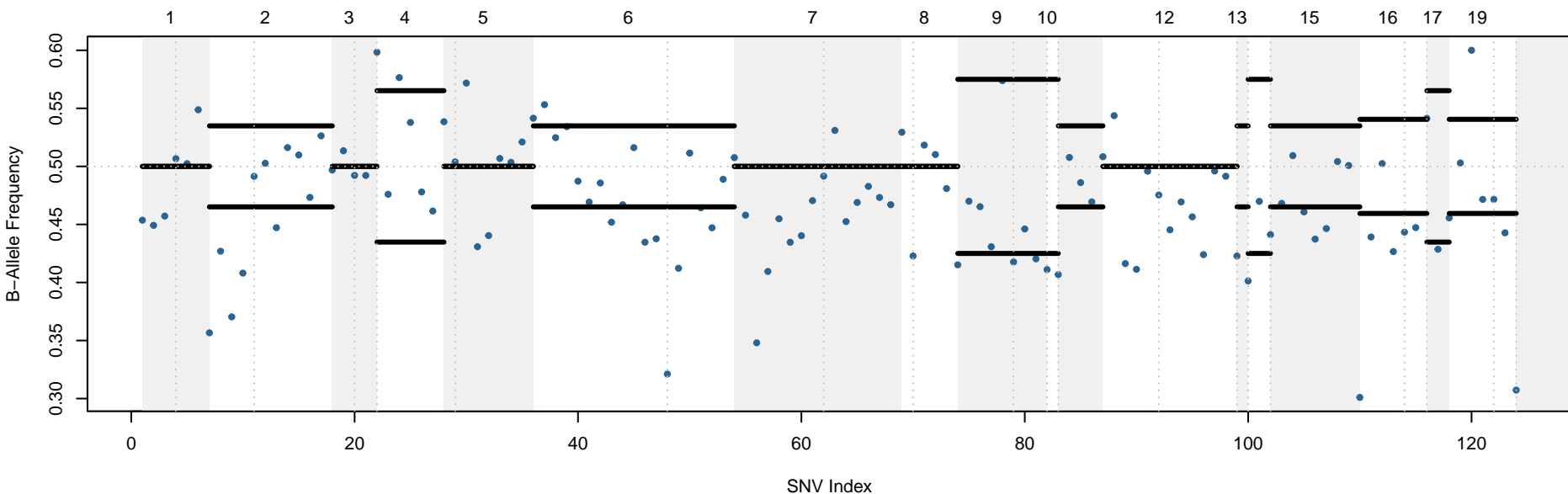




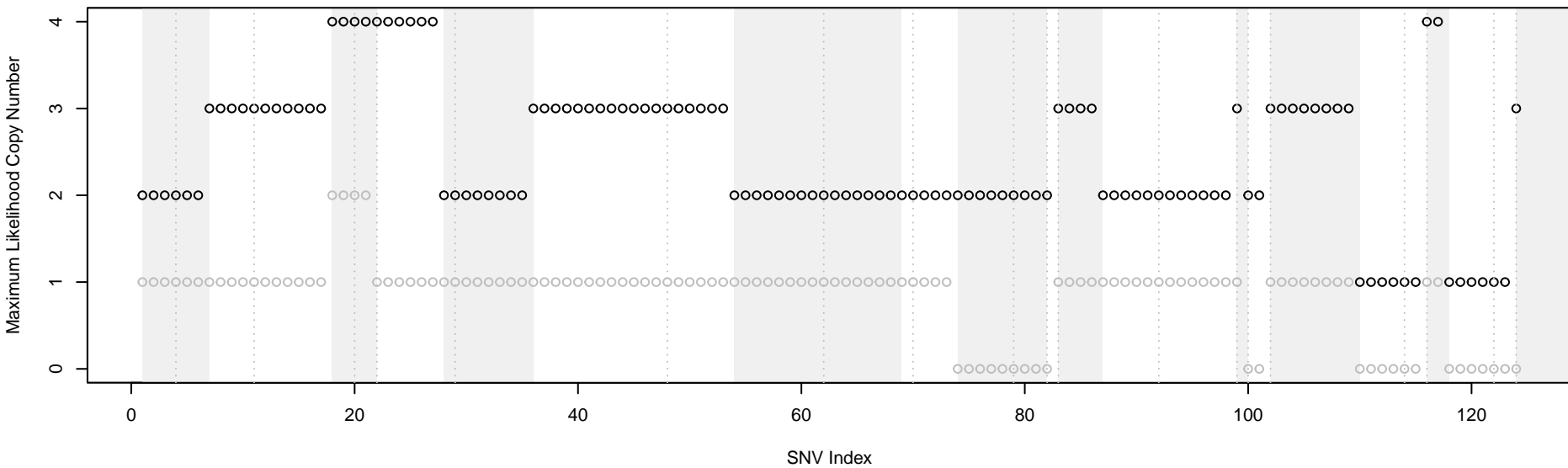
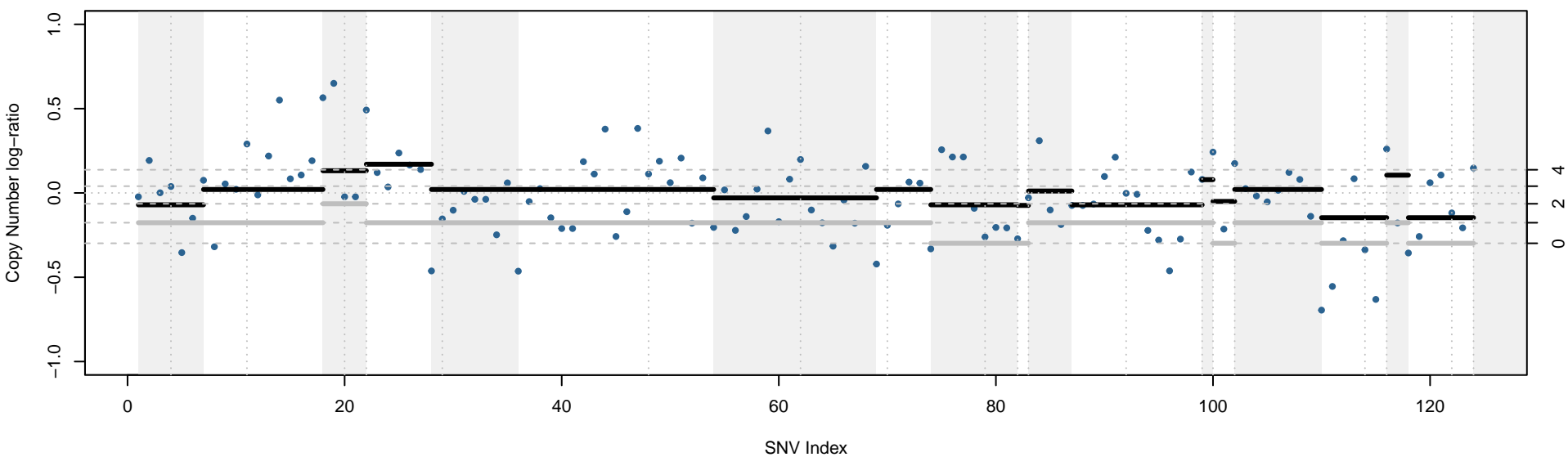
Purity: 0.15 Tumor ploidy: 2.607



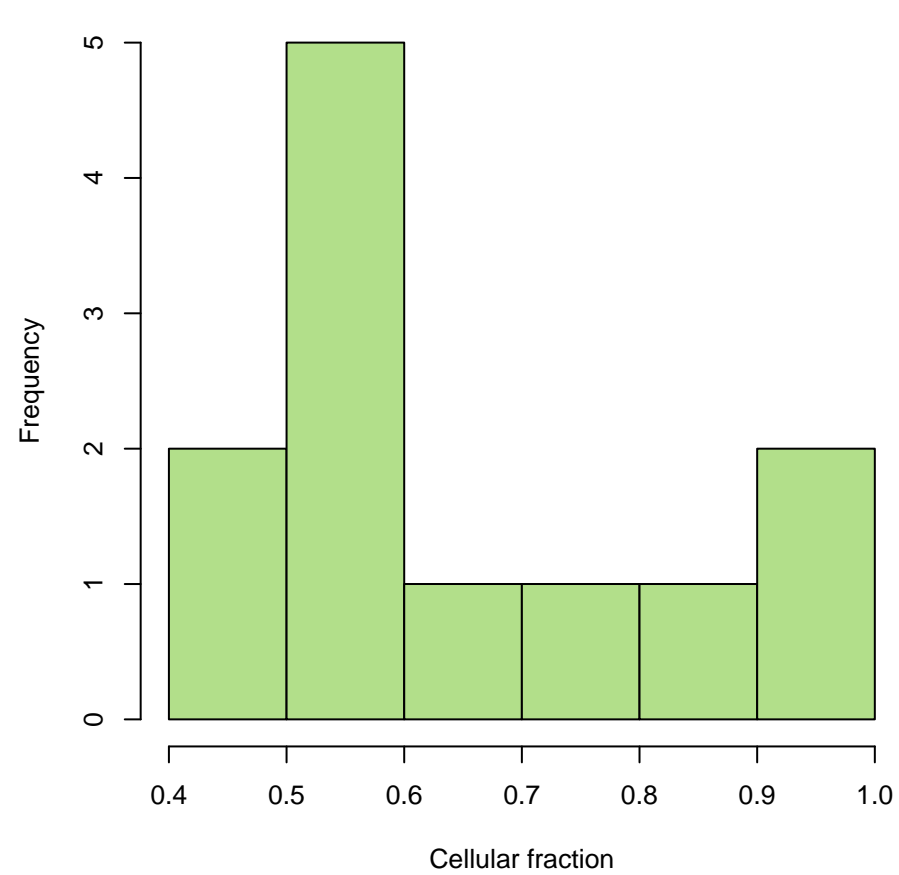
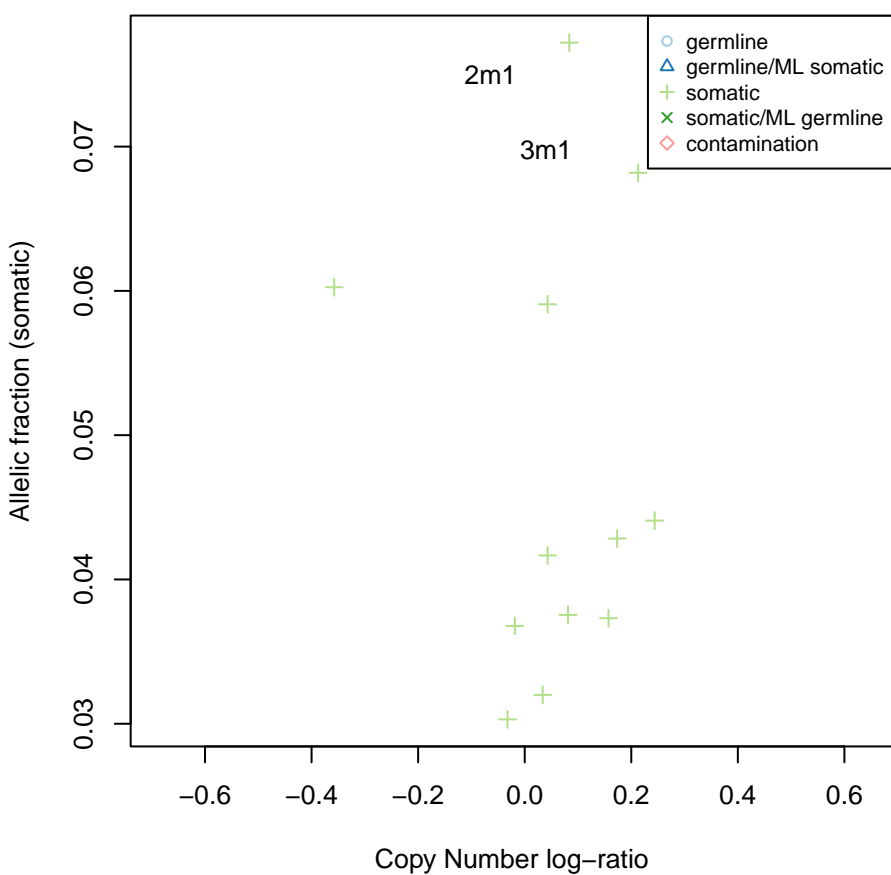
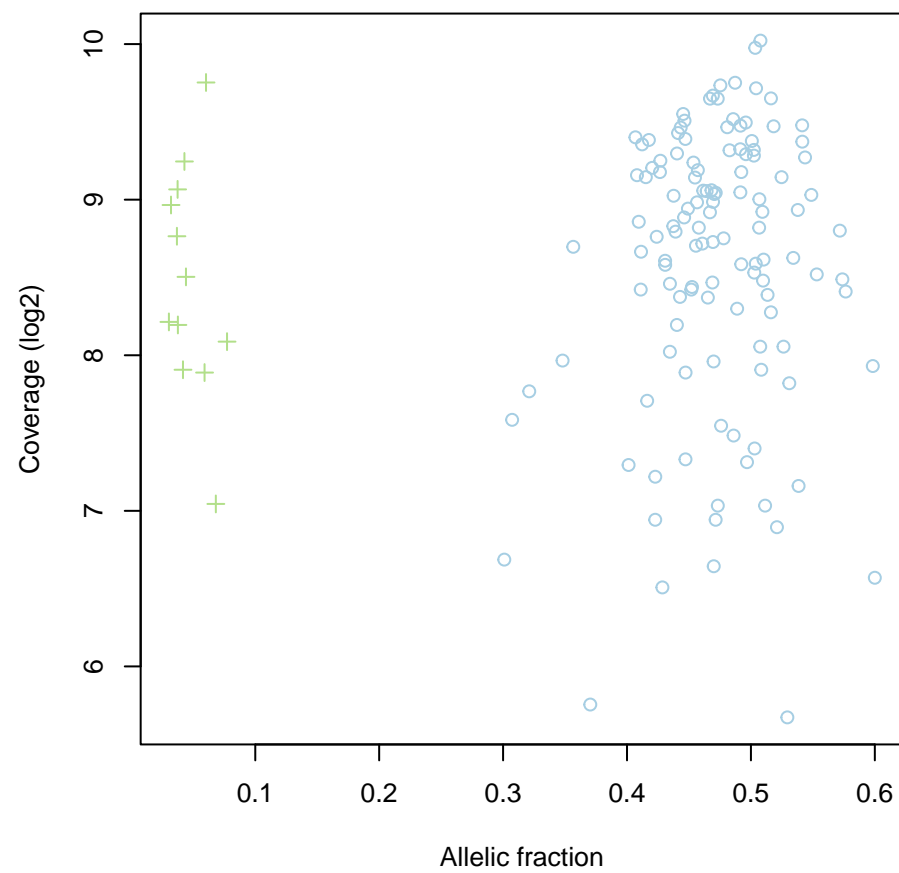
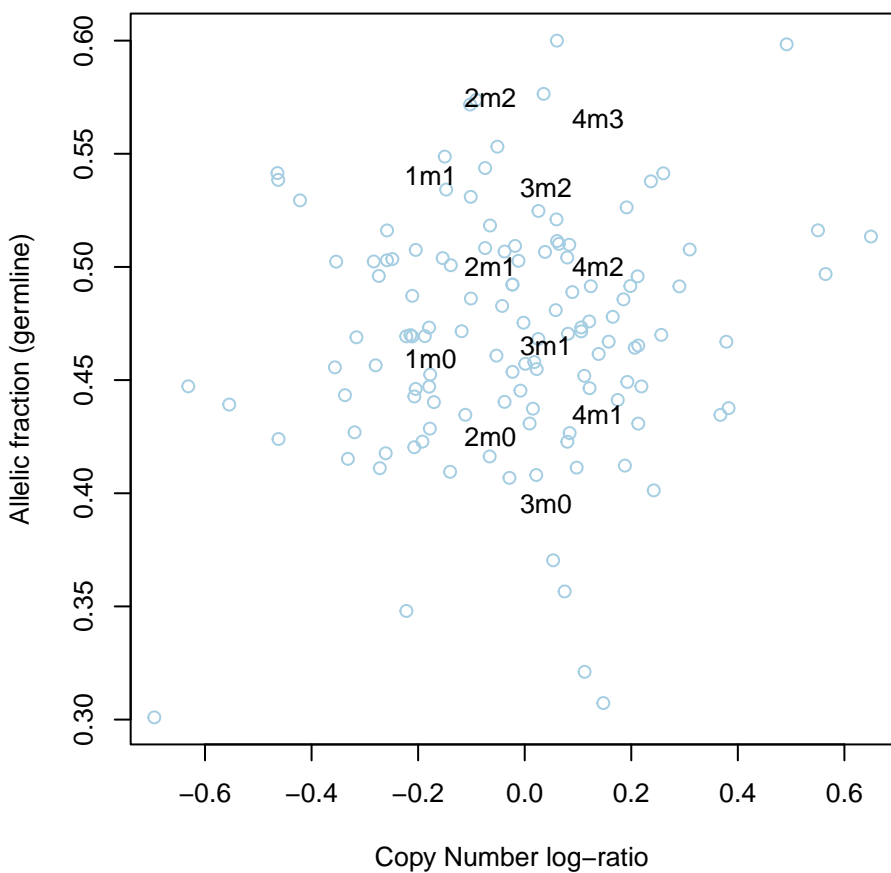
Purity: 0.15 Tumor ploidy: 2.607 SNV log-likelihood: 19.3 GoF: 96.2% Mean coverage: 577;448



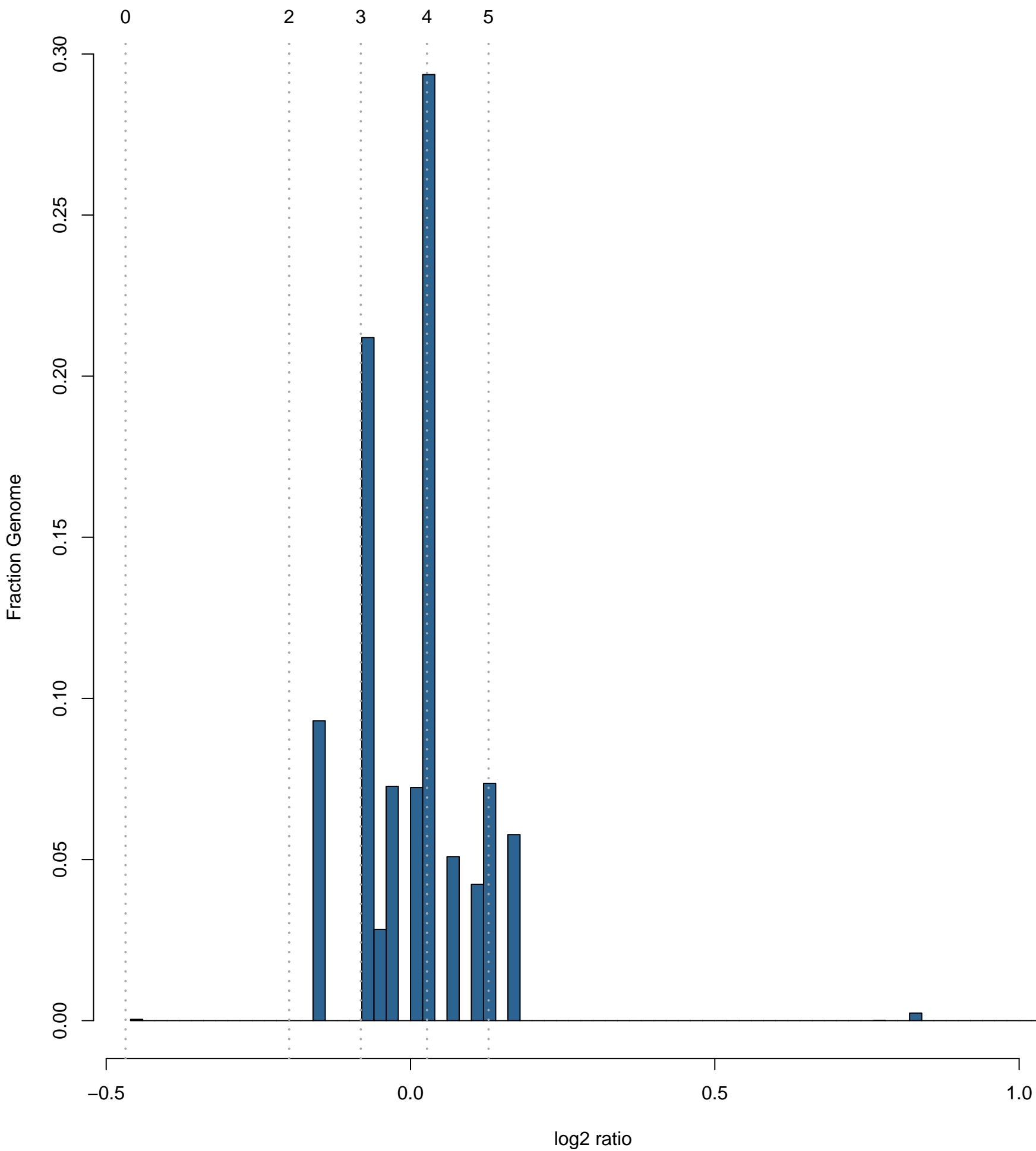
SCNA-fit log-likelihood: -14500.07



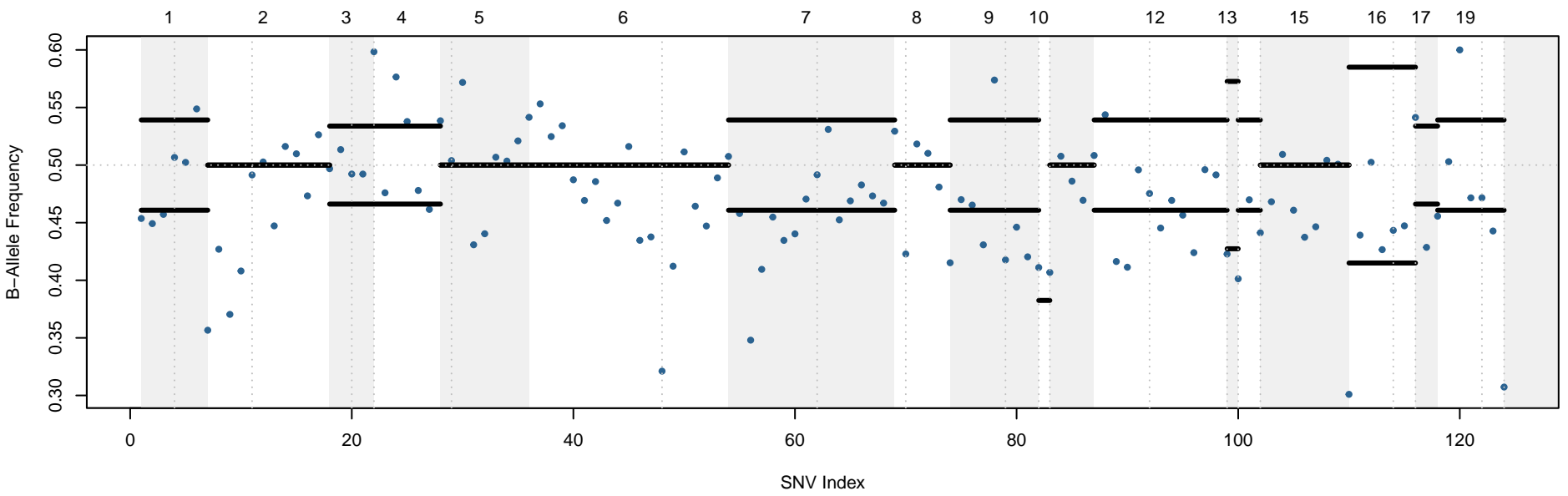




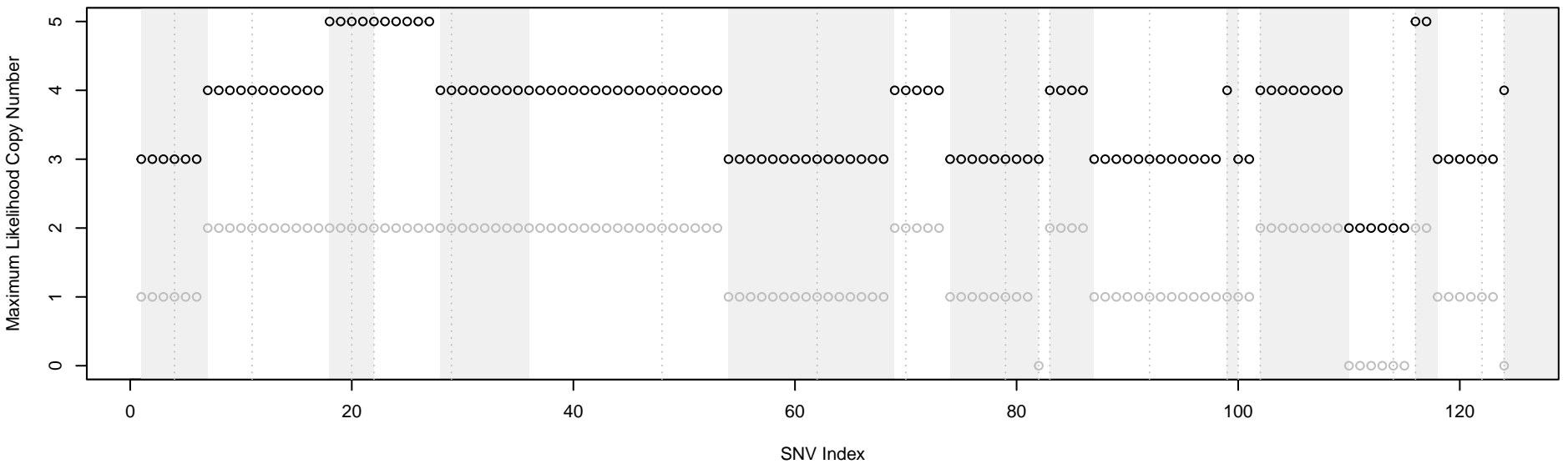
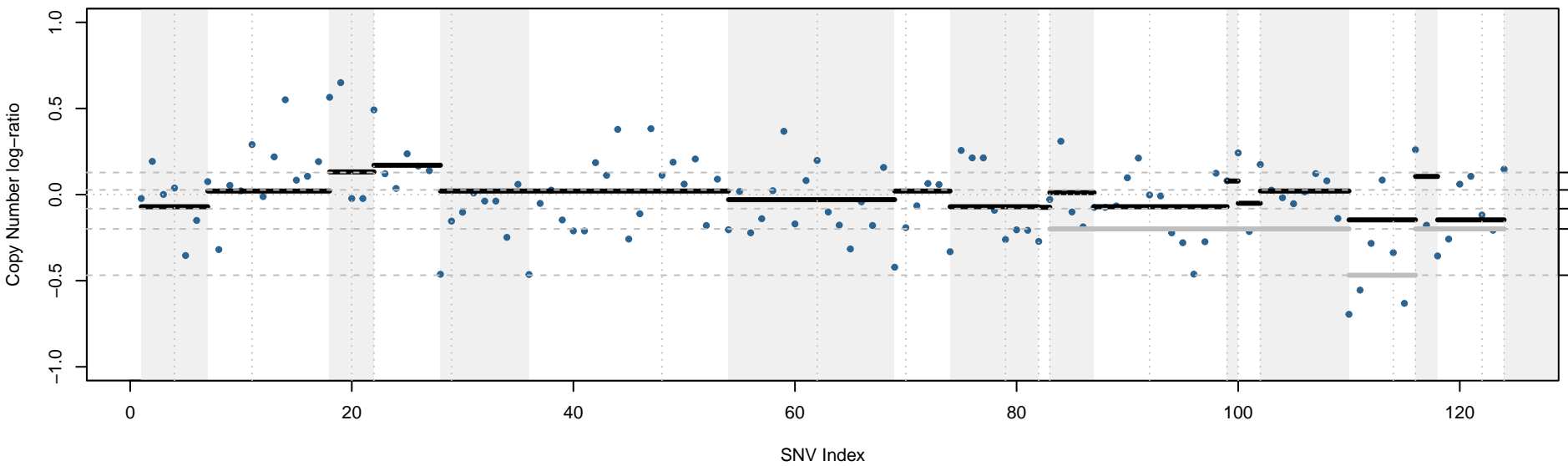
Purity: 0.17 Tumor ploidy: 3.743

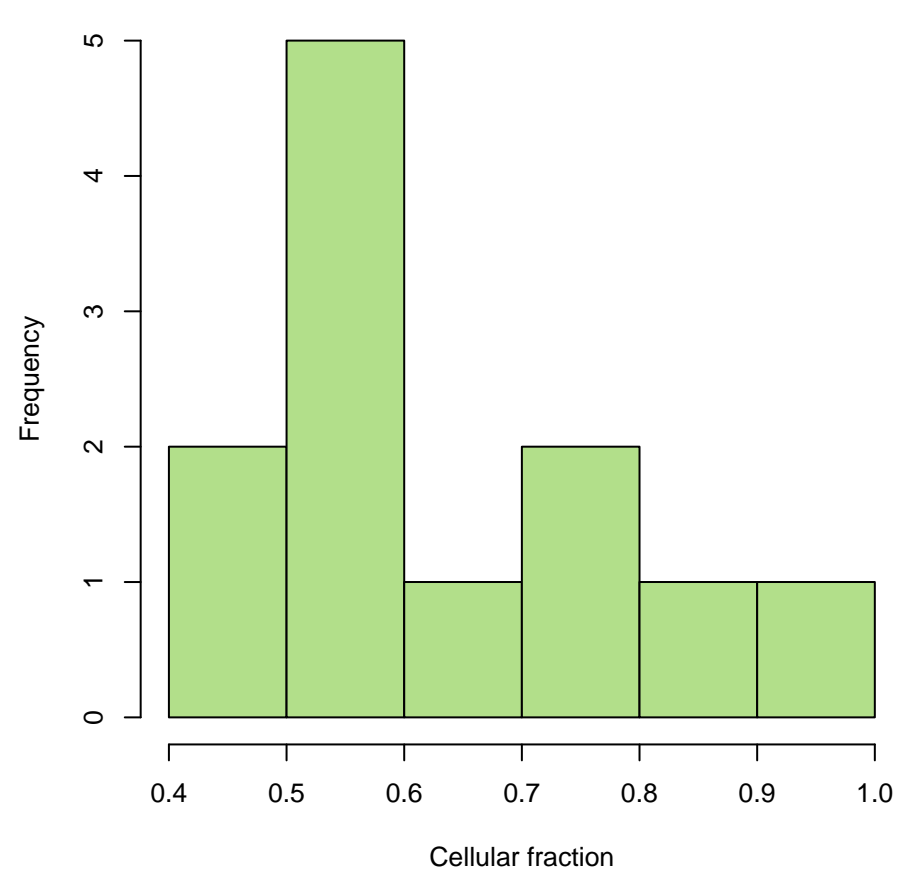
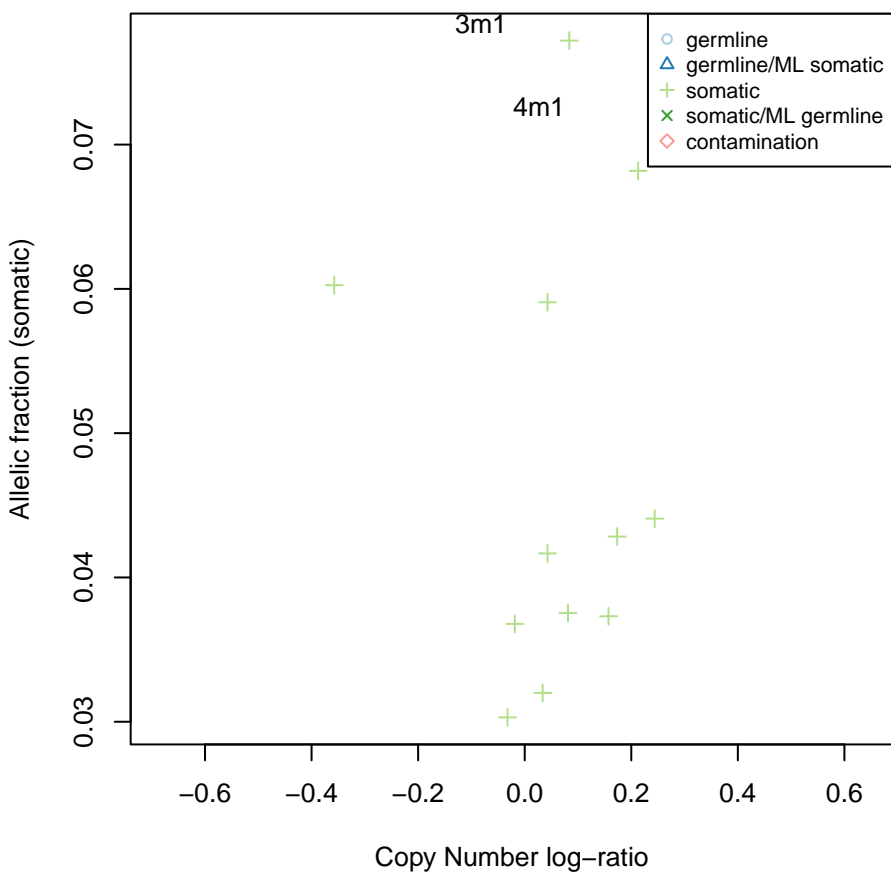
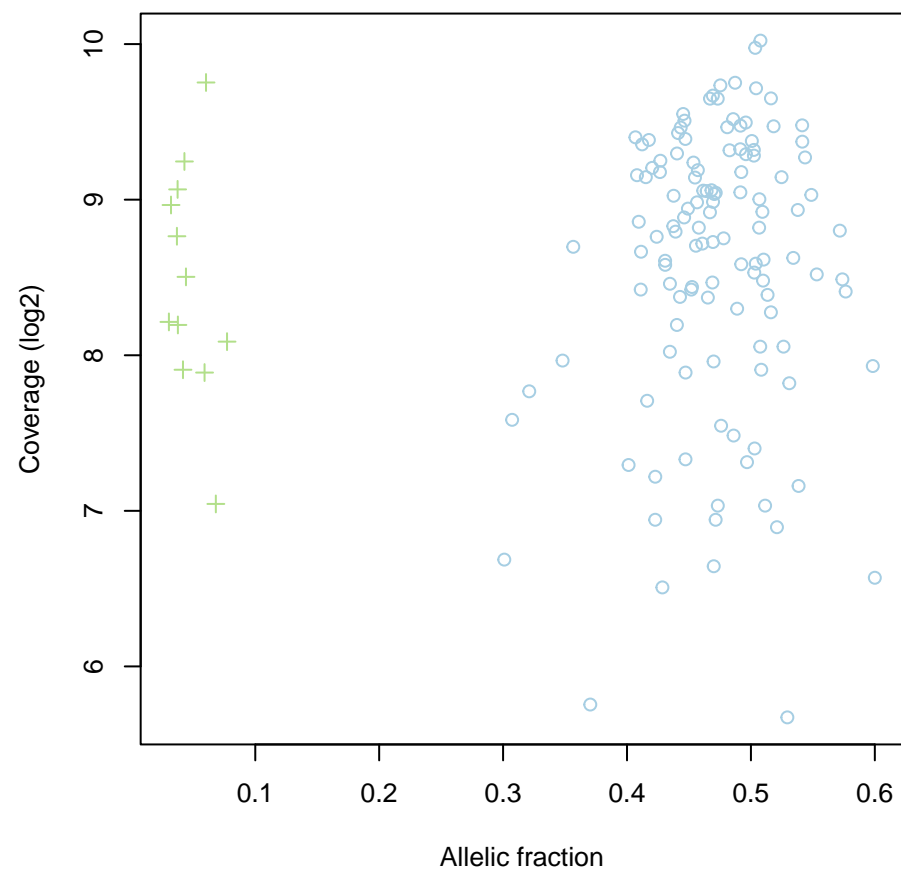
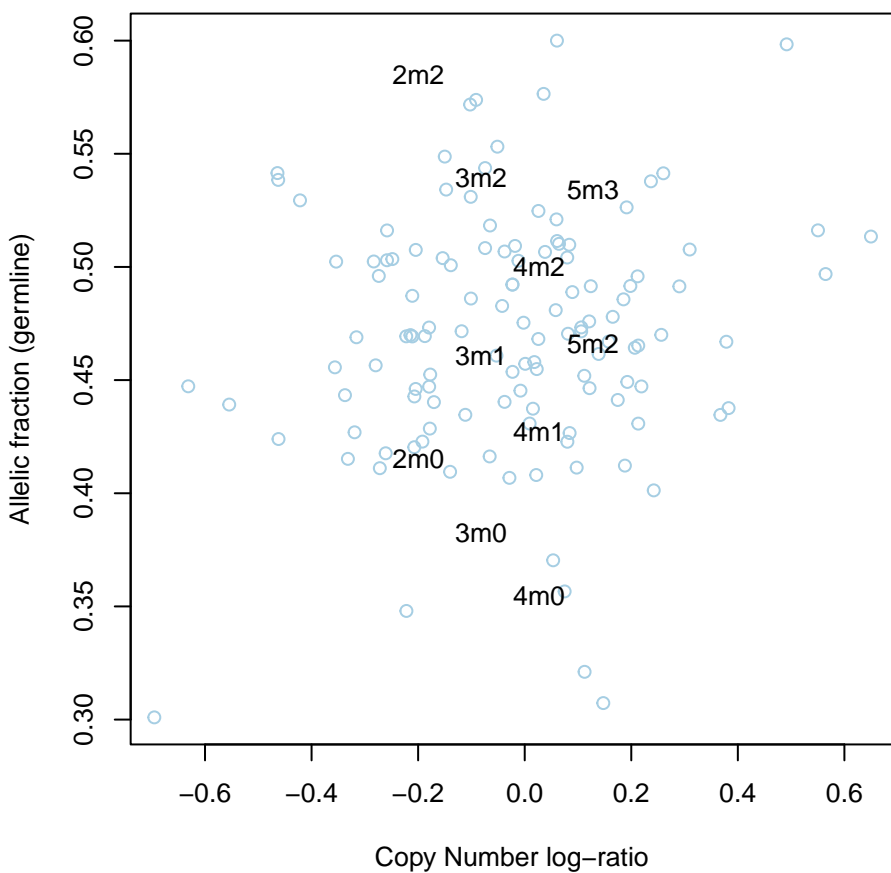


Purity: 0.17 Tumor ploidy: 3.743 SNV log-likelihood: -27.75 GoF: 96.3% Mean coverage: 577;448

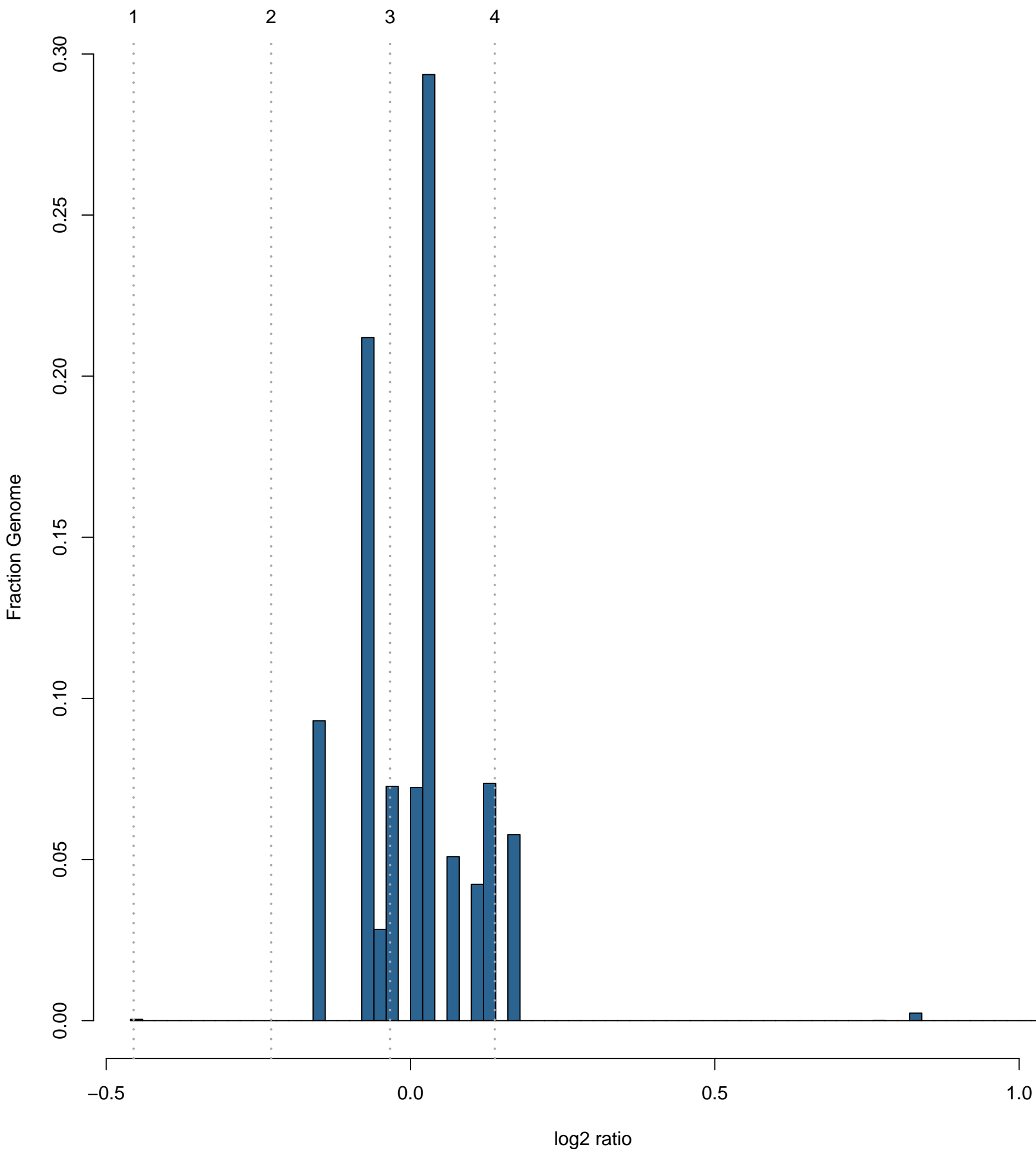


SCNA-fit log-likelihood: -14501.47

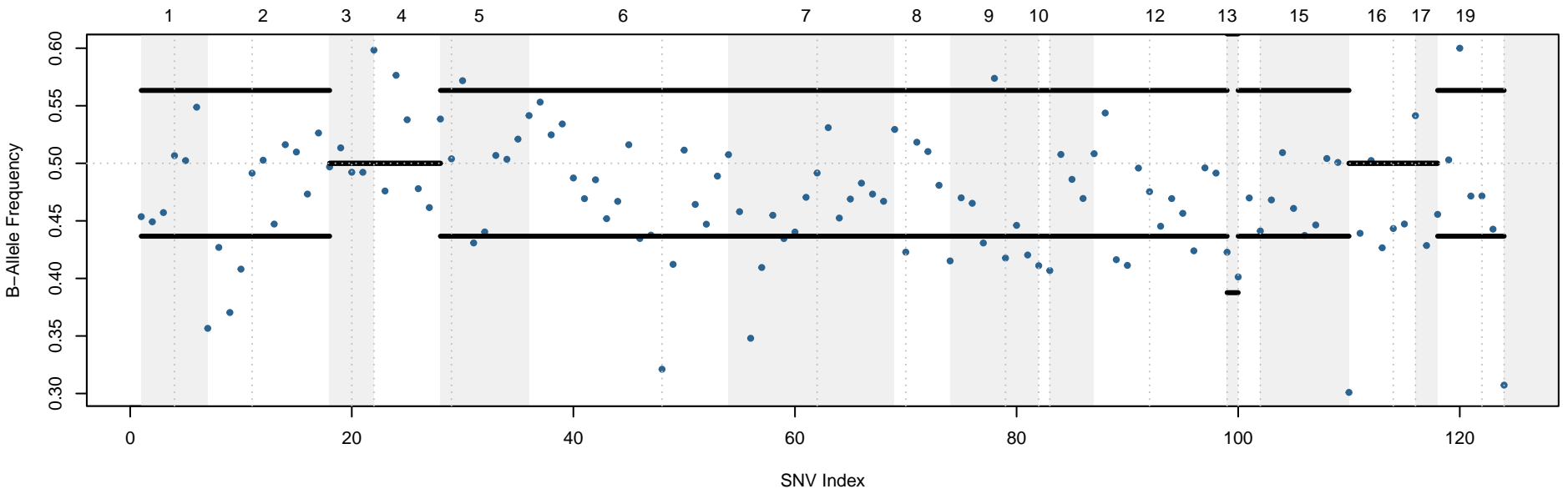




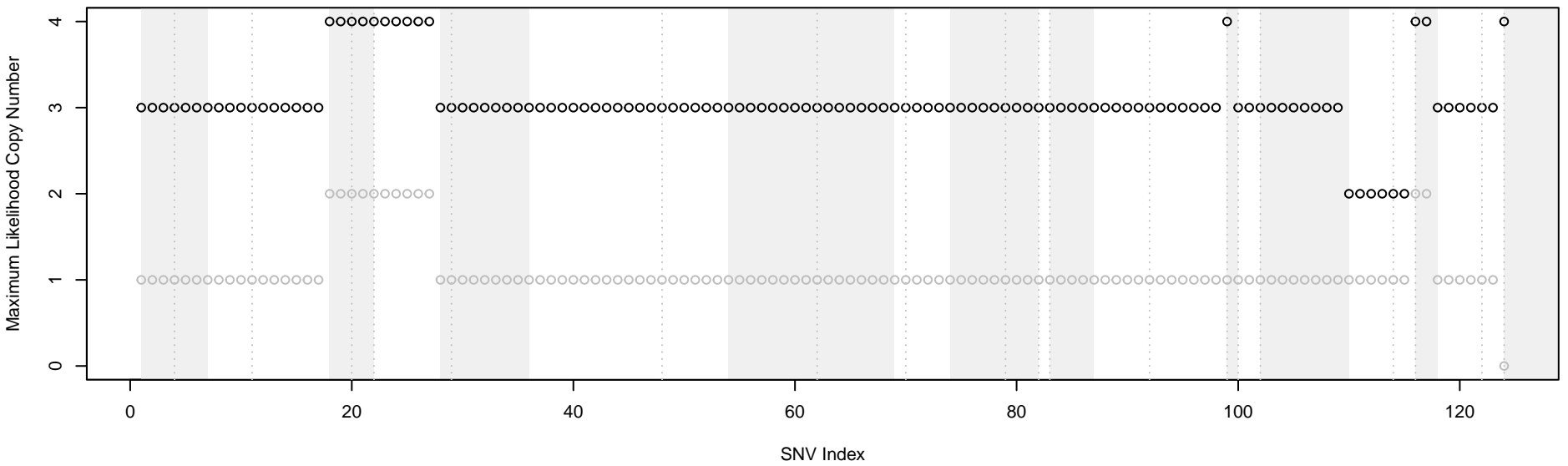
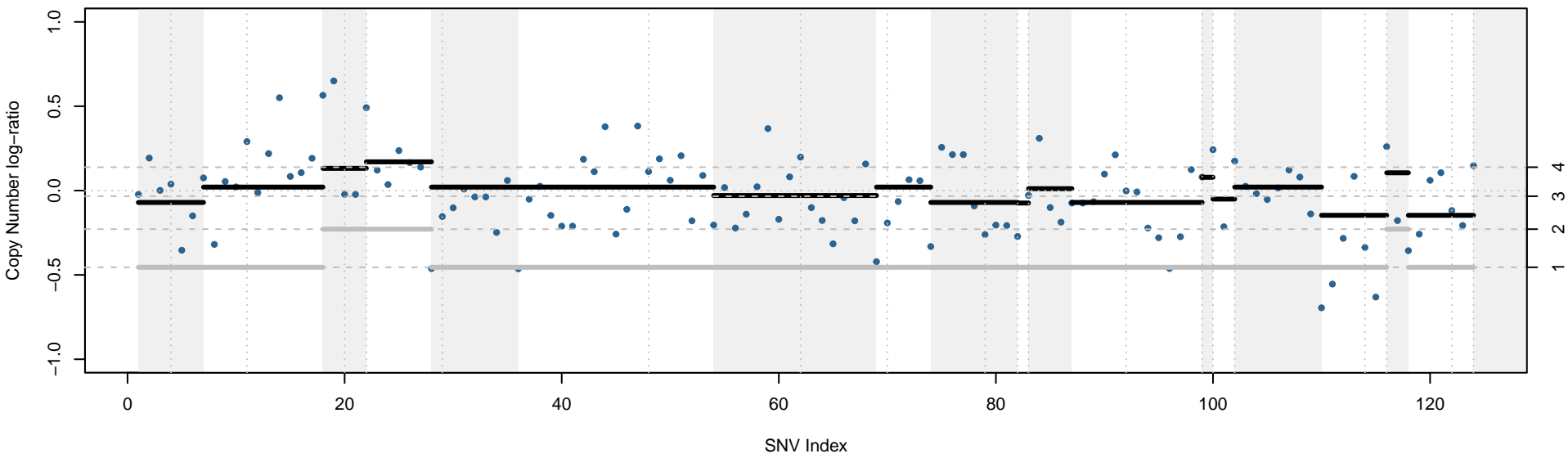
Purity: 0.29 Tumor ploidy: 3.186

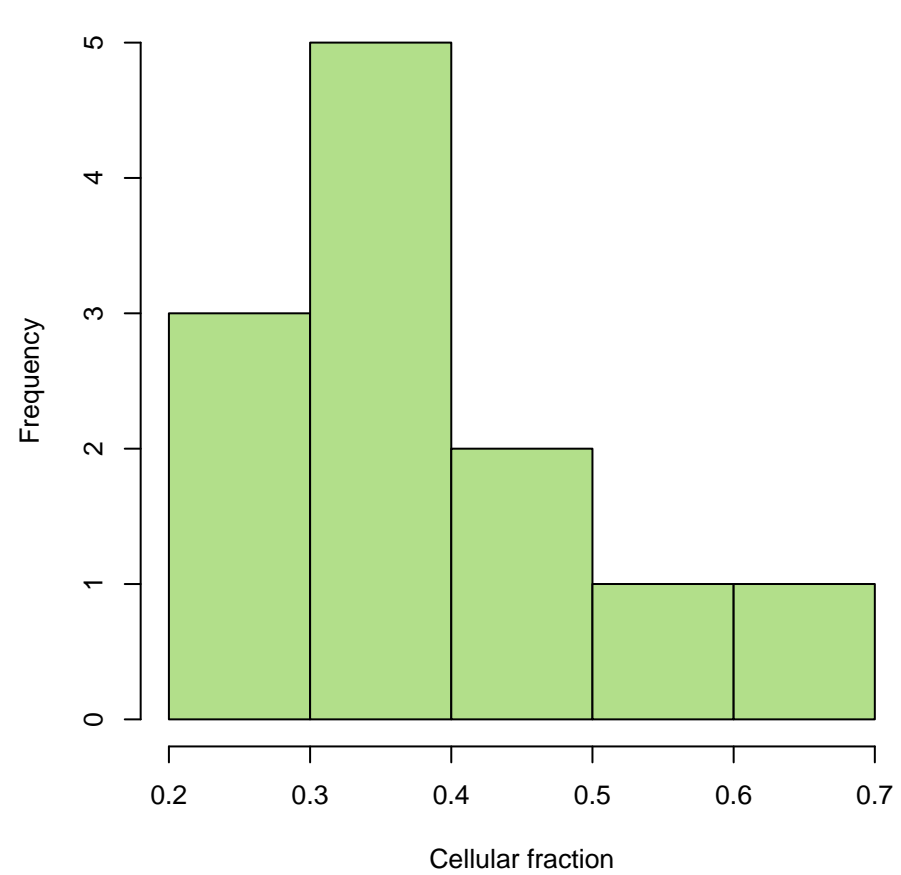
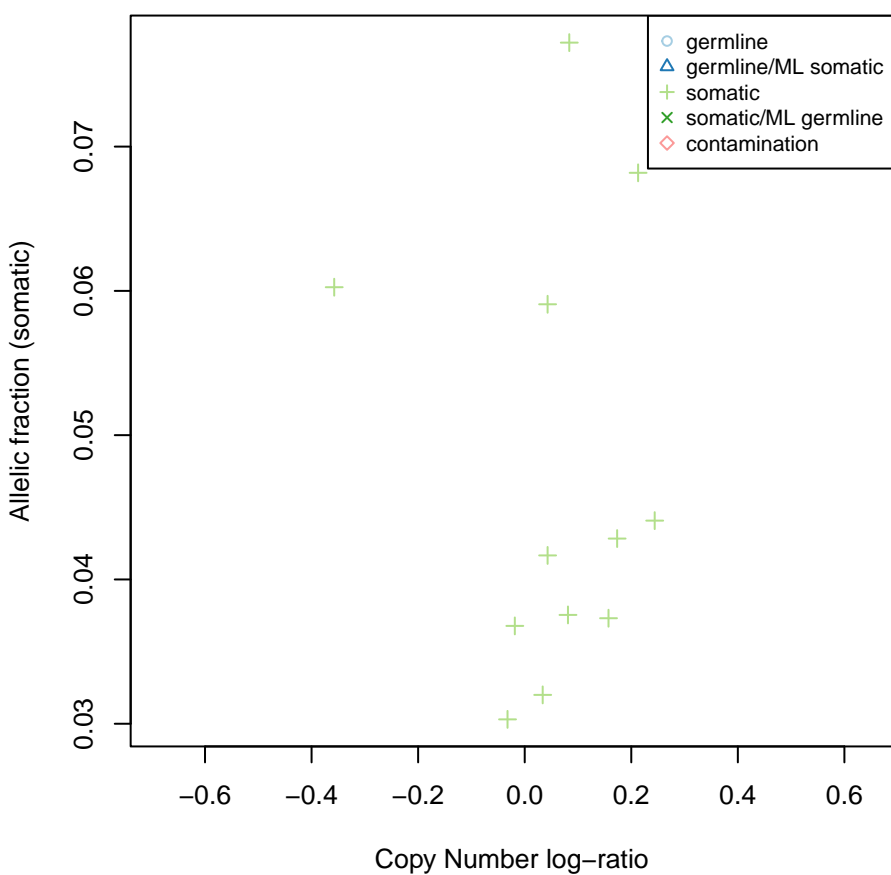
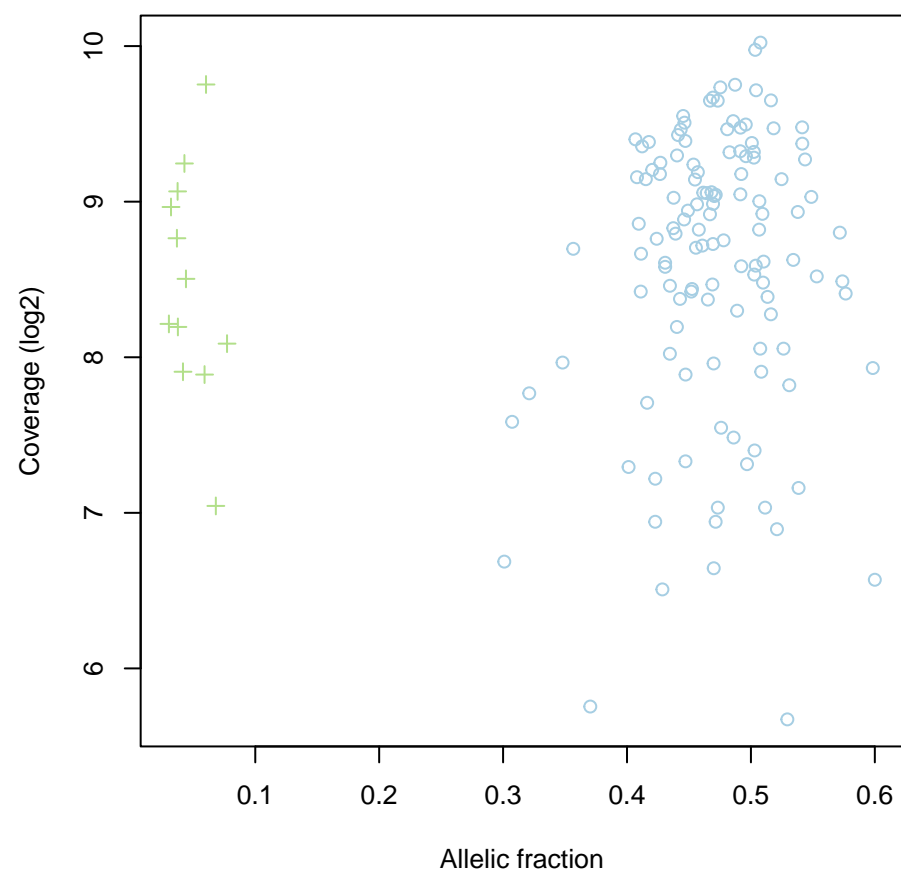
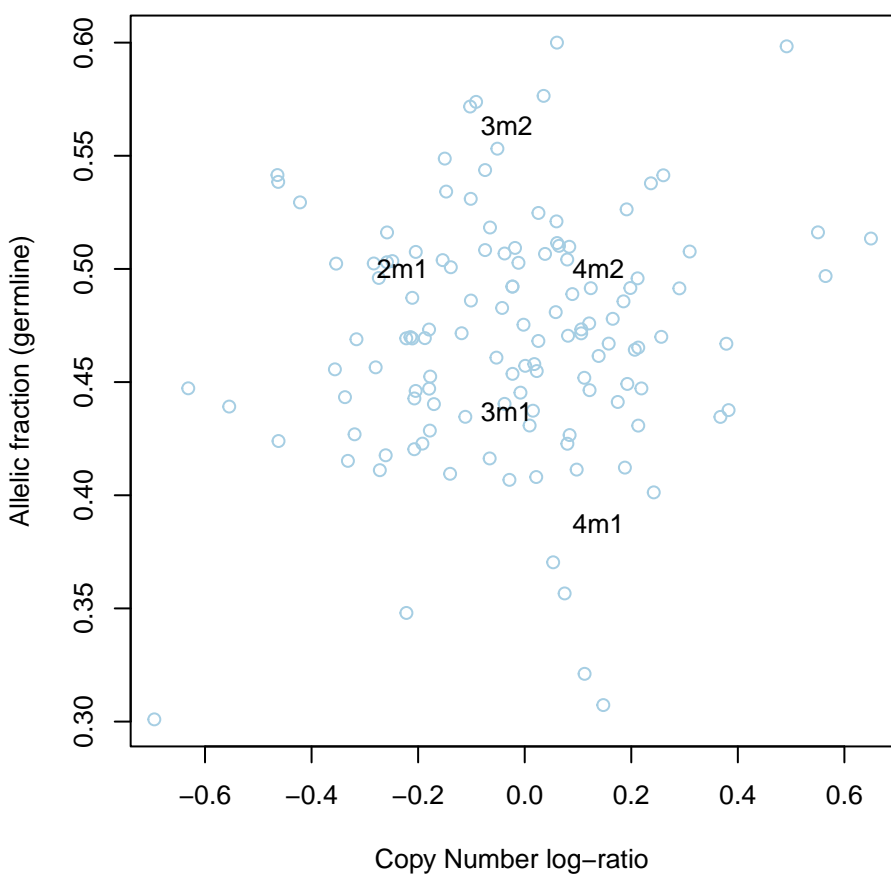


Purity: 0.29 Tumor ploidy: 3.186 SNV log-likelihood: -38.98 GoF: 94.2% Mean coverage: 577;448

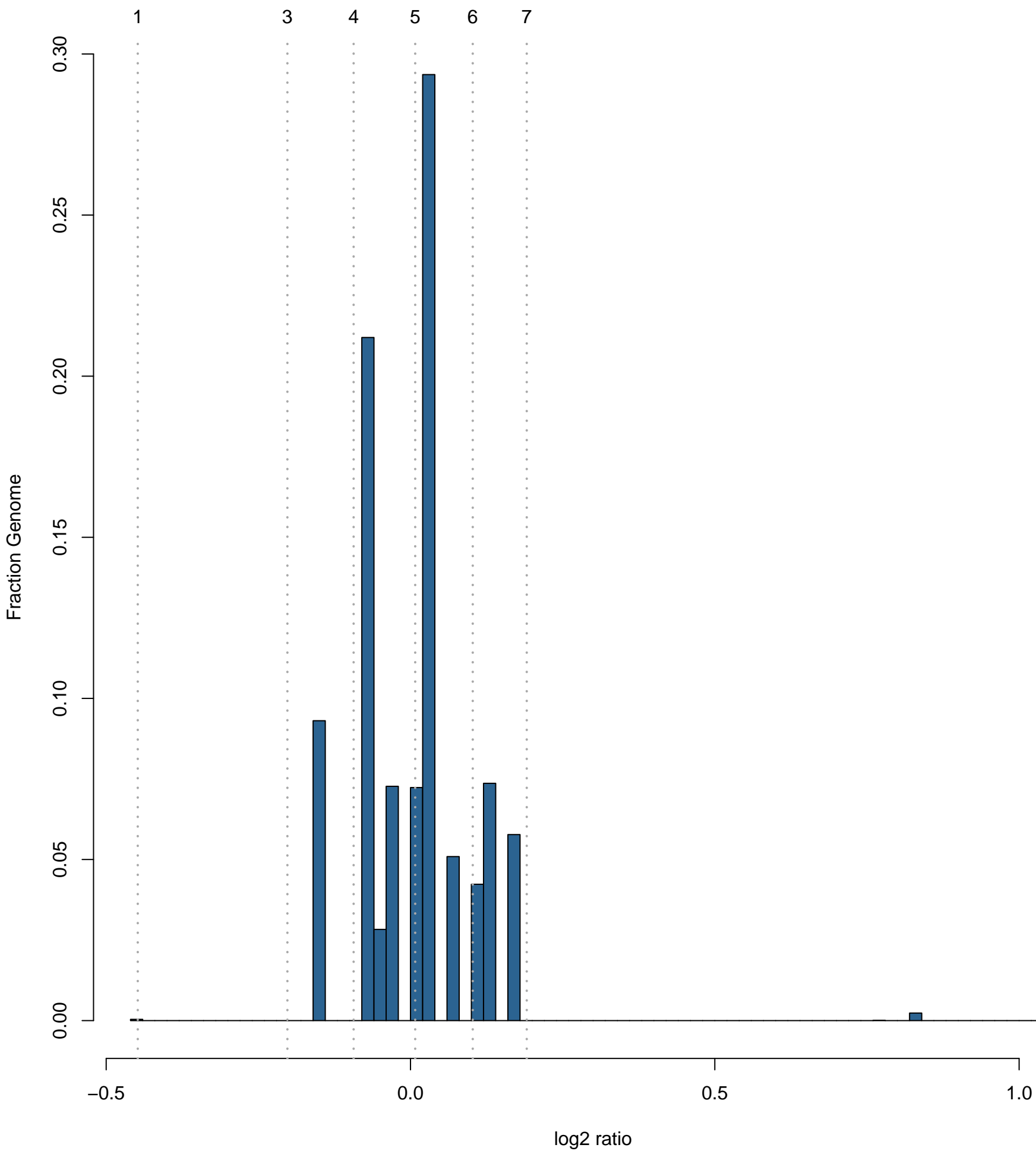


SCNA-fit log-likelihood: -14564.12



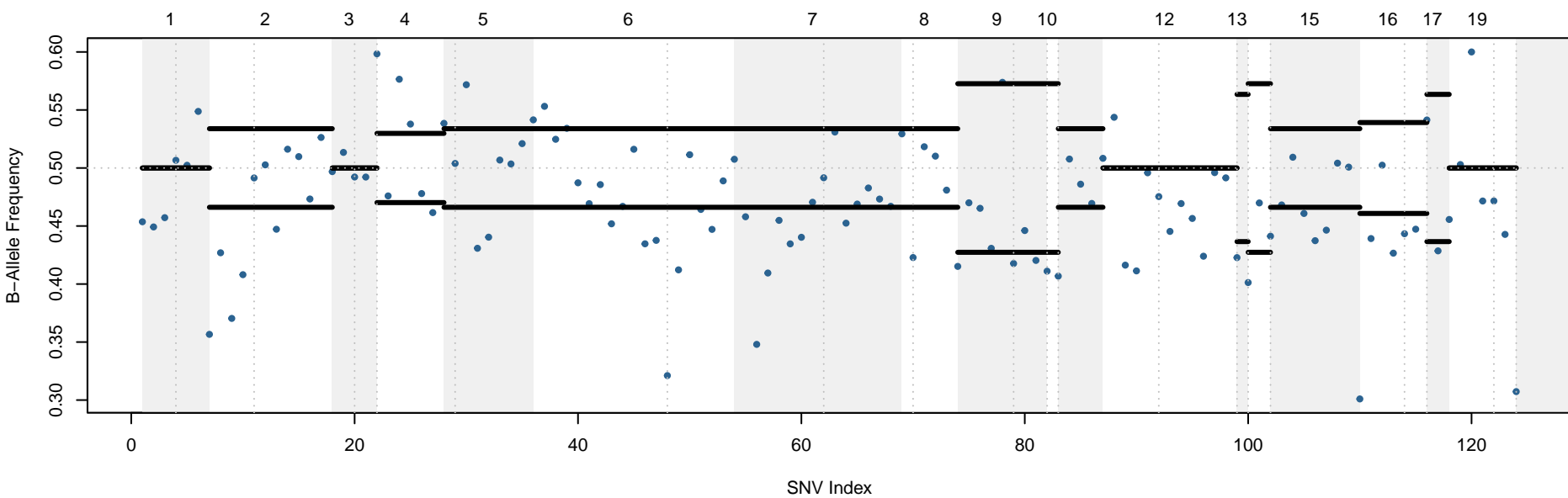


Purity: 0.17 Tumor ploidy: 4.921

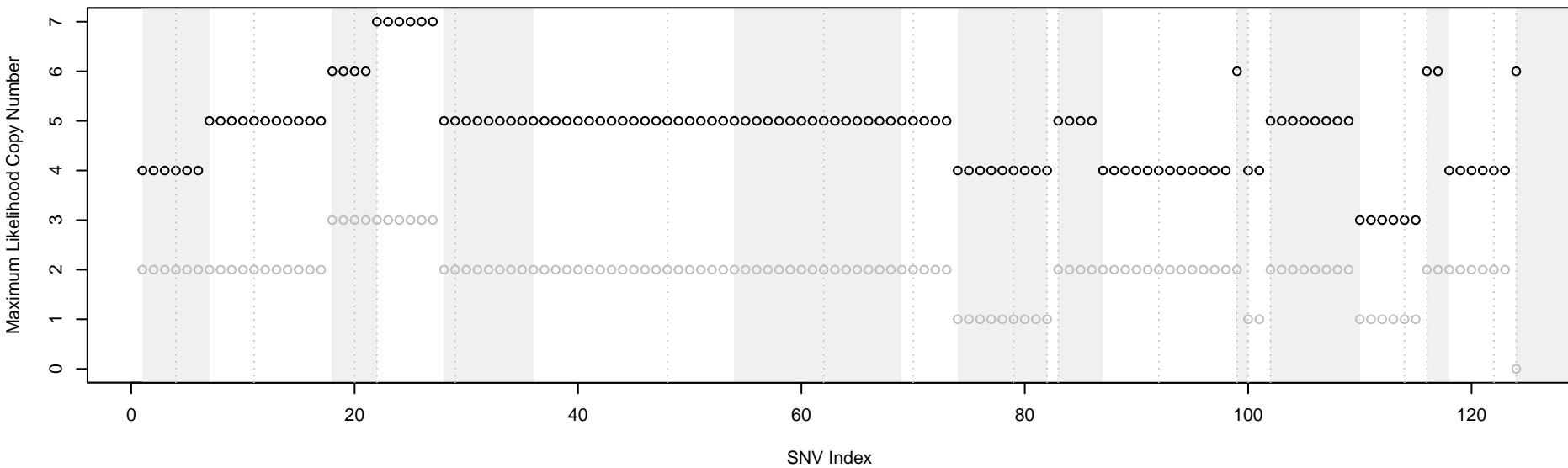
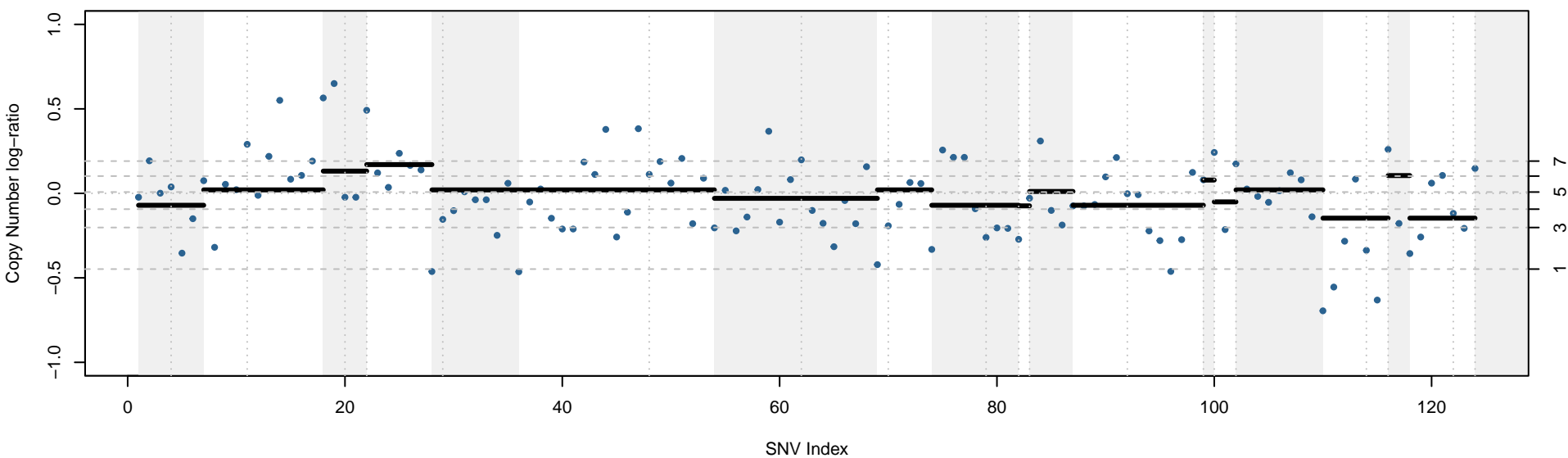


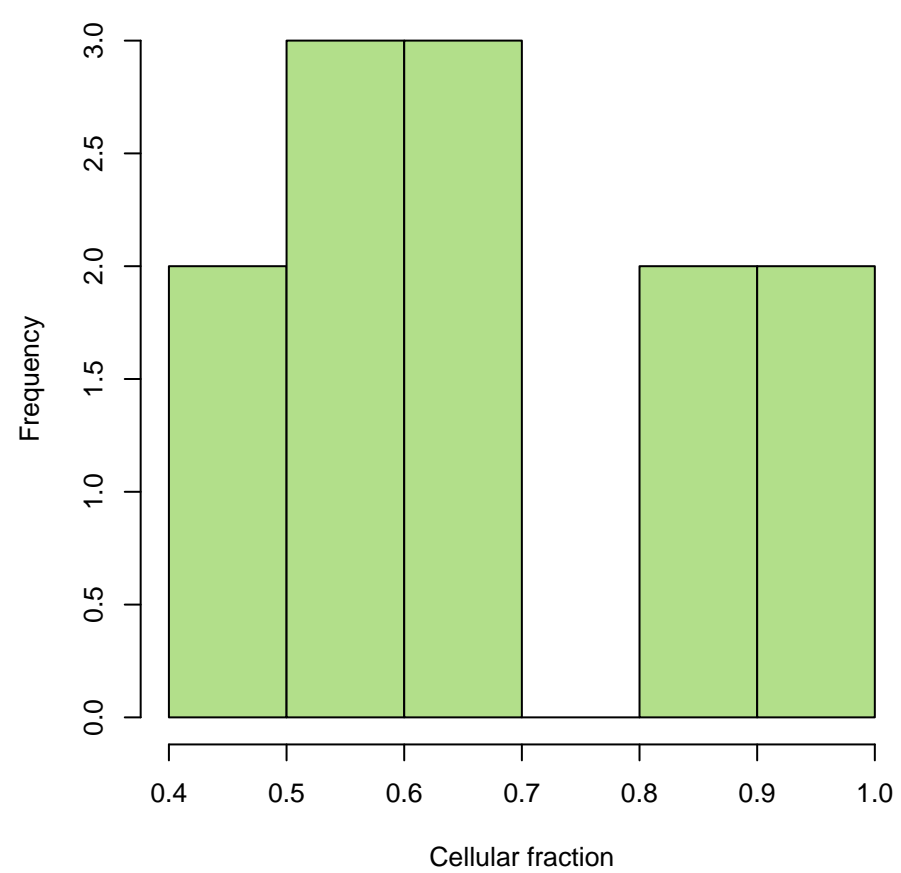
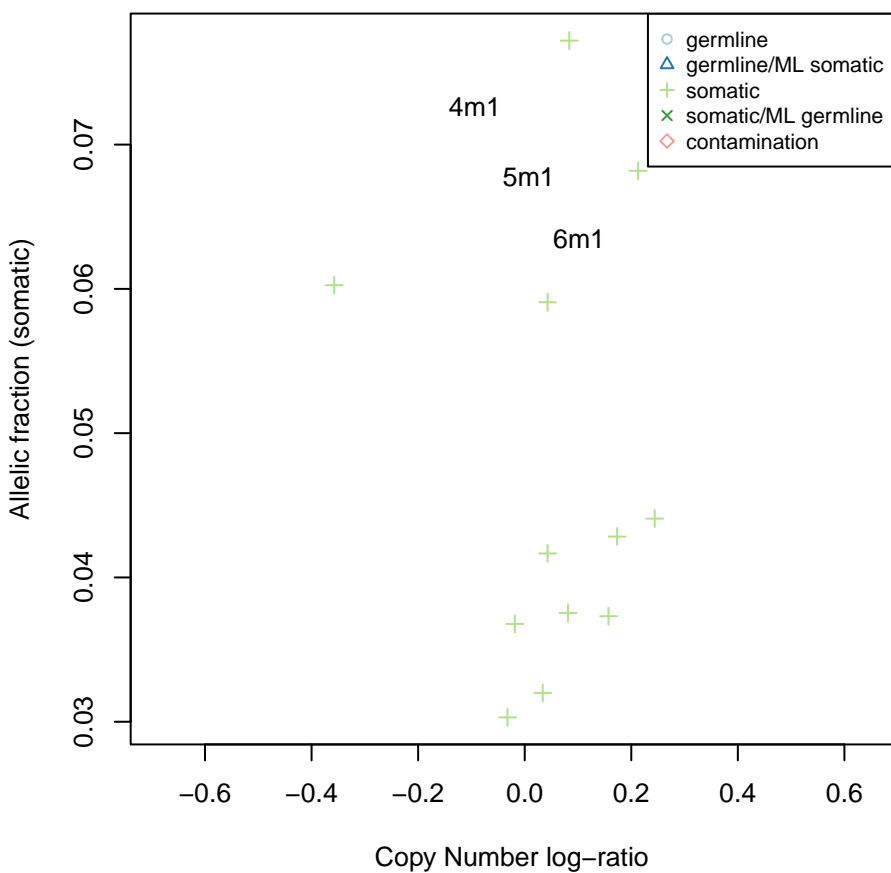
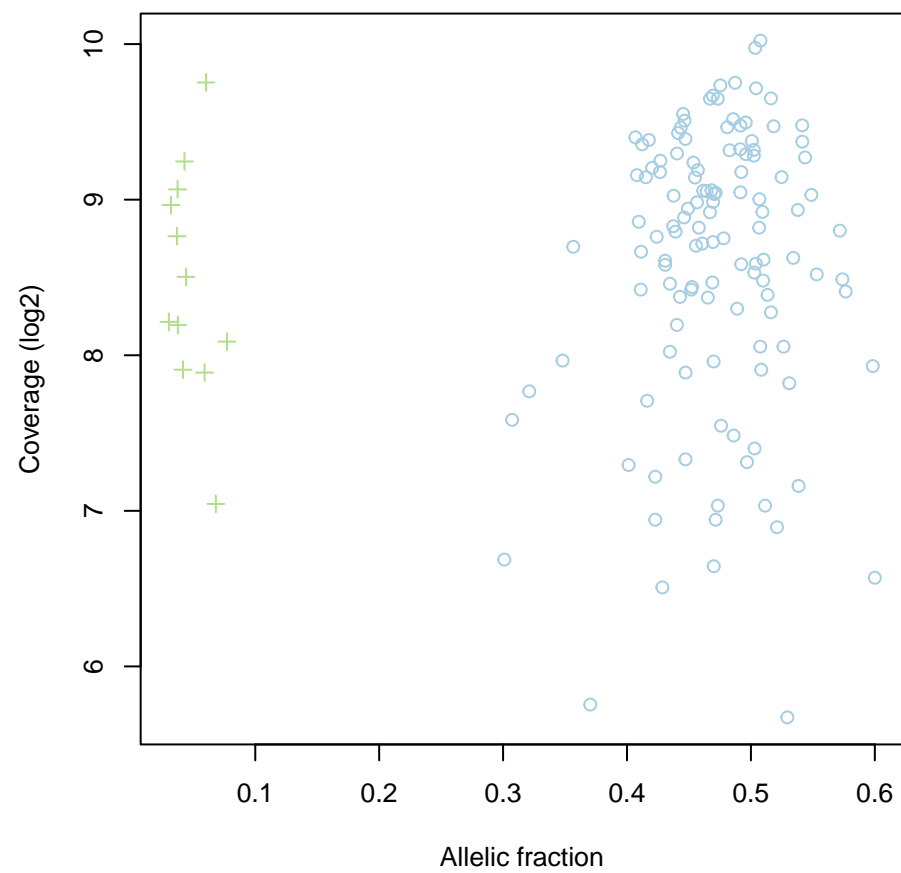
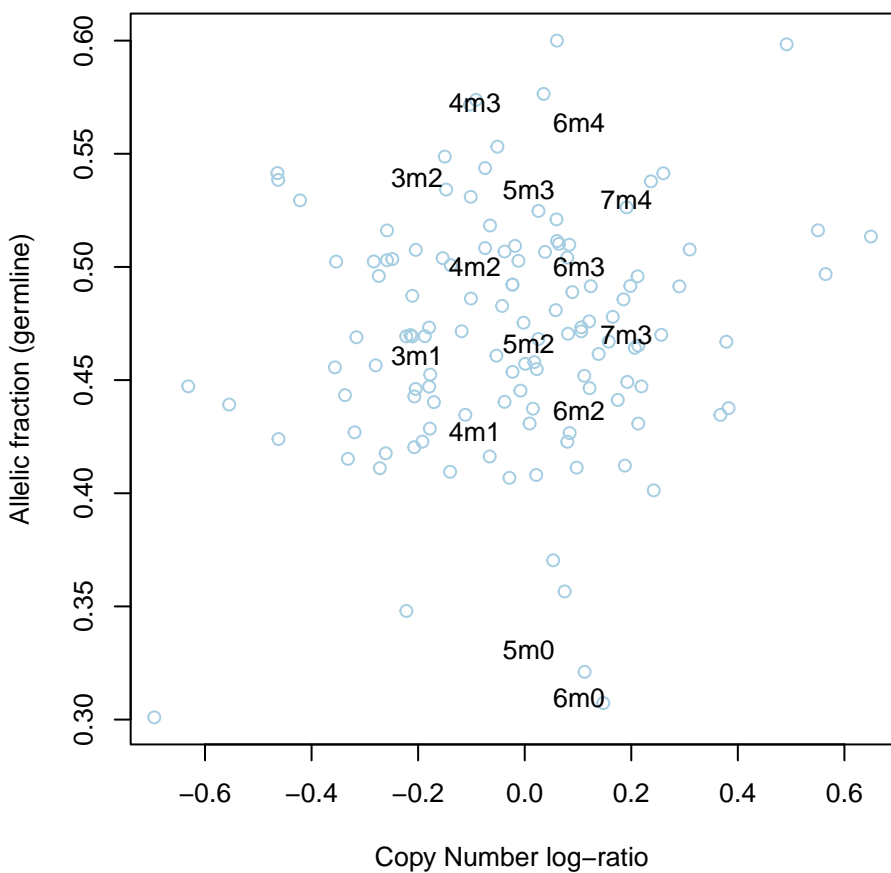


Purity: 0.17 Tumor ploidy: 4.921 SNV log-likelihood: -47.87 GoF: 96.9% Mean coverage: 577;448

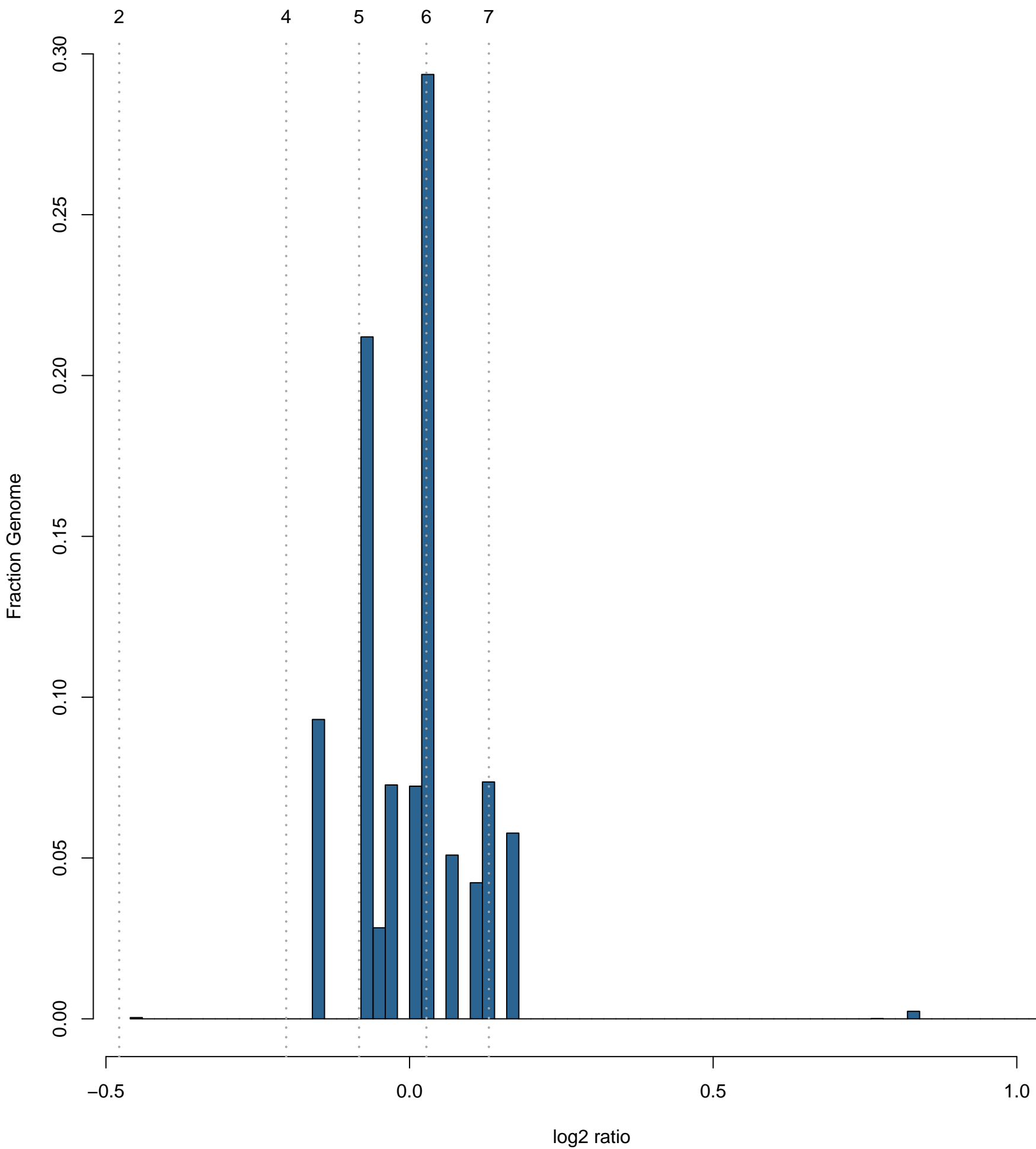


SCNA-fit log-likelihood: -14498.1

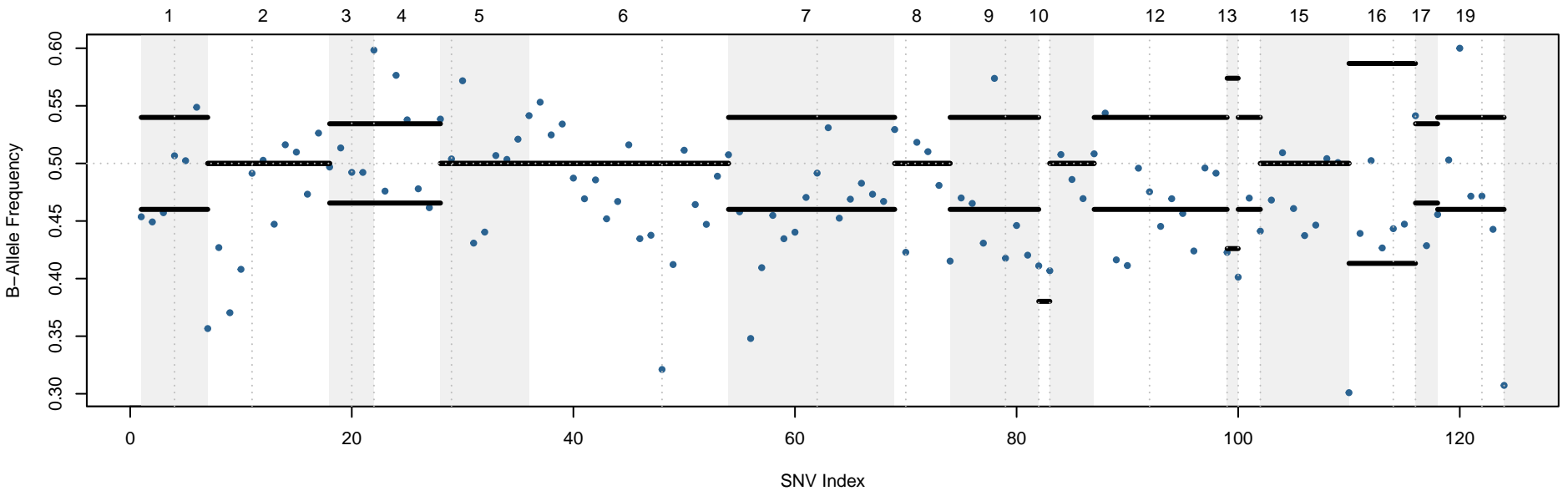




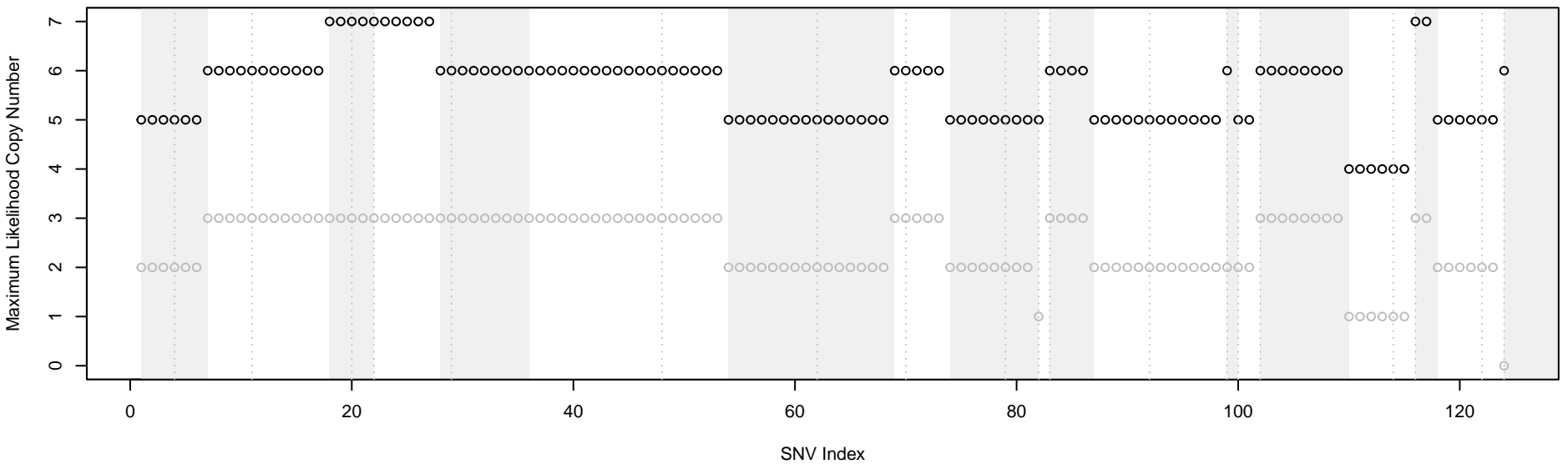
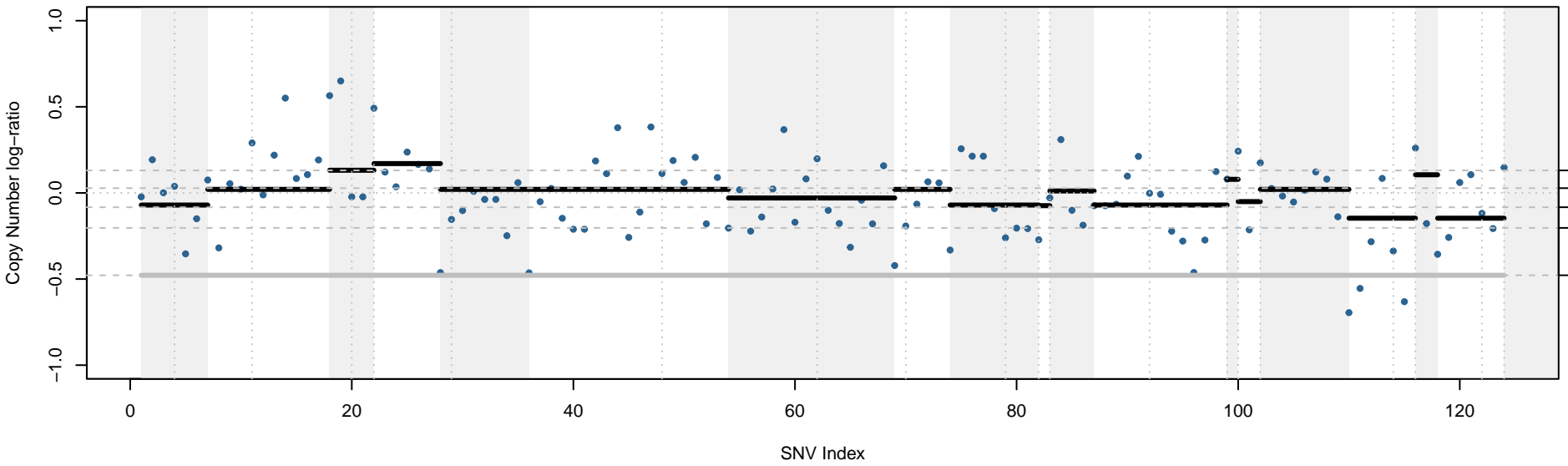
Purity: 0.21 Tumor ploidy: 5.743

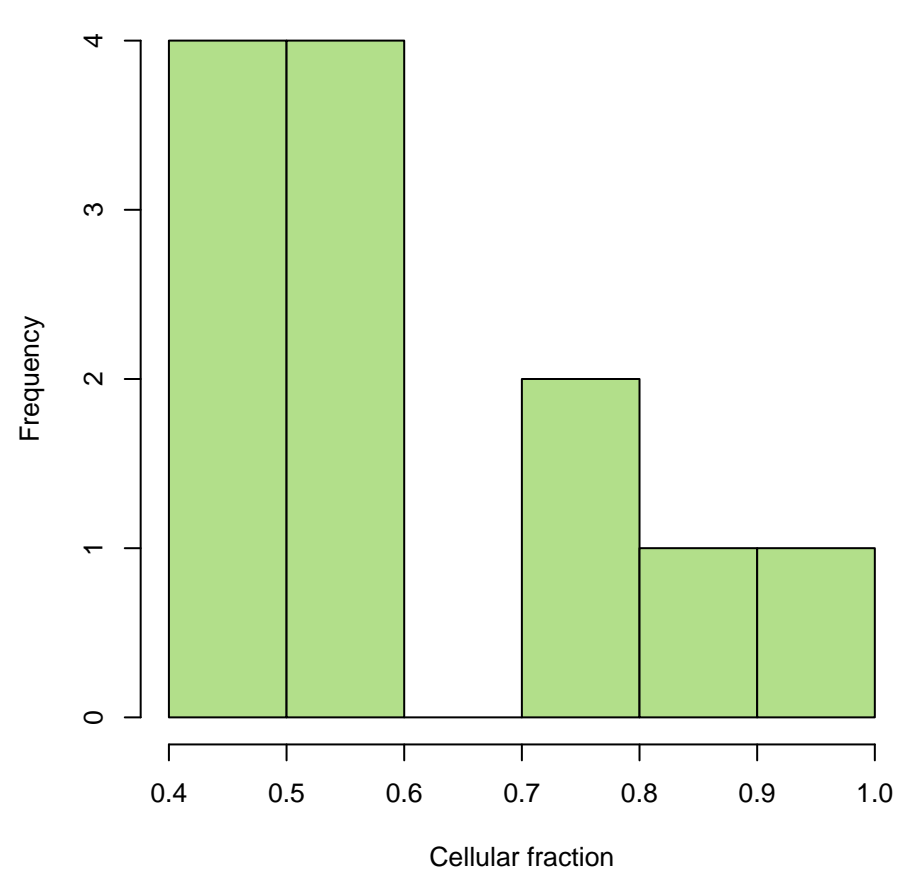
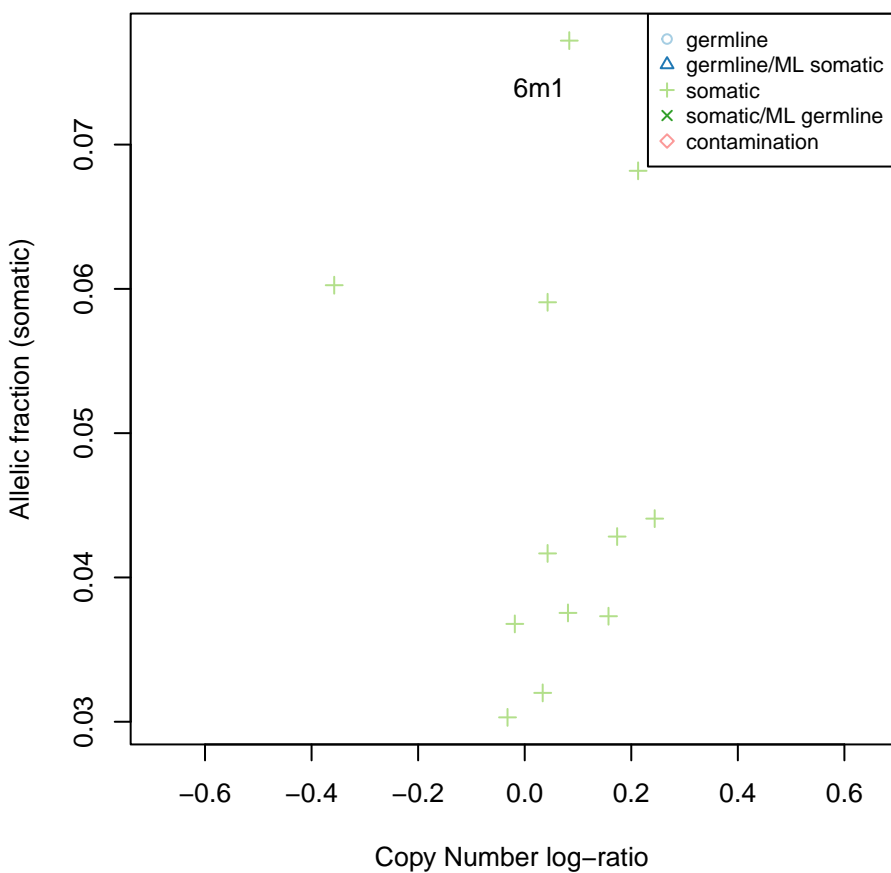
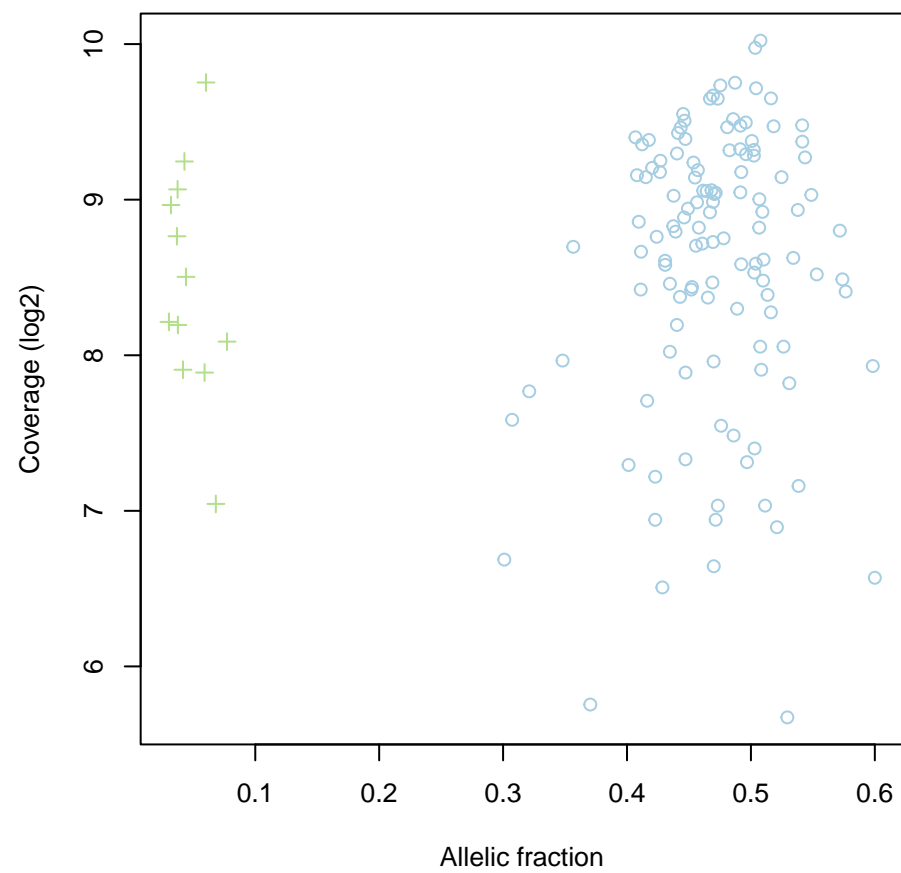
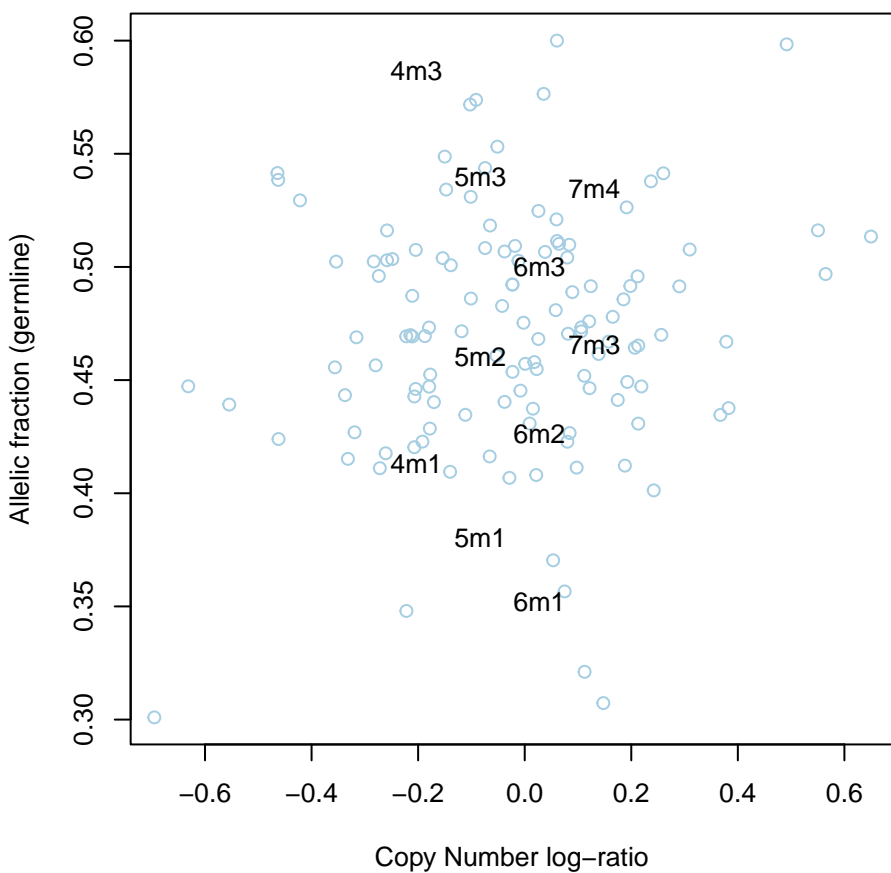


Purity: 0.21 Tumor ploidy: 5.743 SNV log-likelihood: -79.78 GoF: 96.3% Mean coverage: 577;448

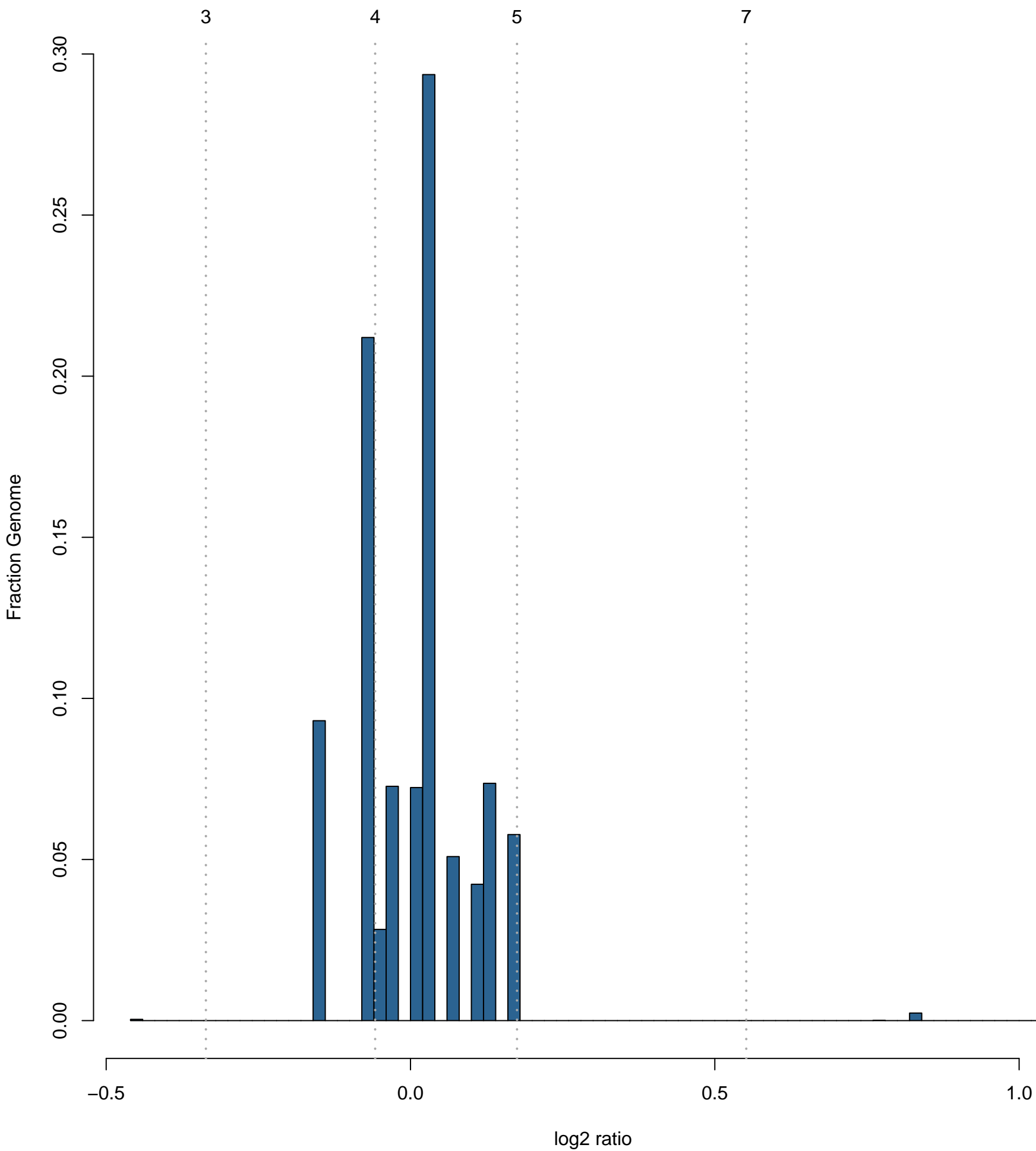


SCNA-fit log-likelihood: -14502.56

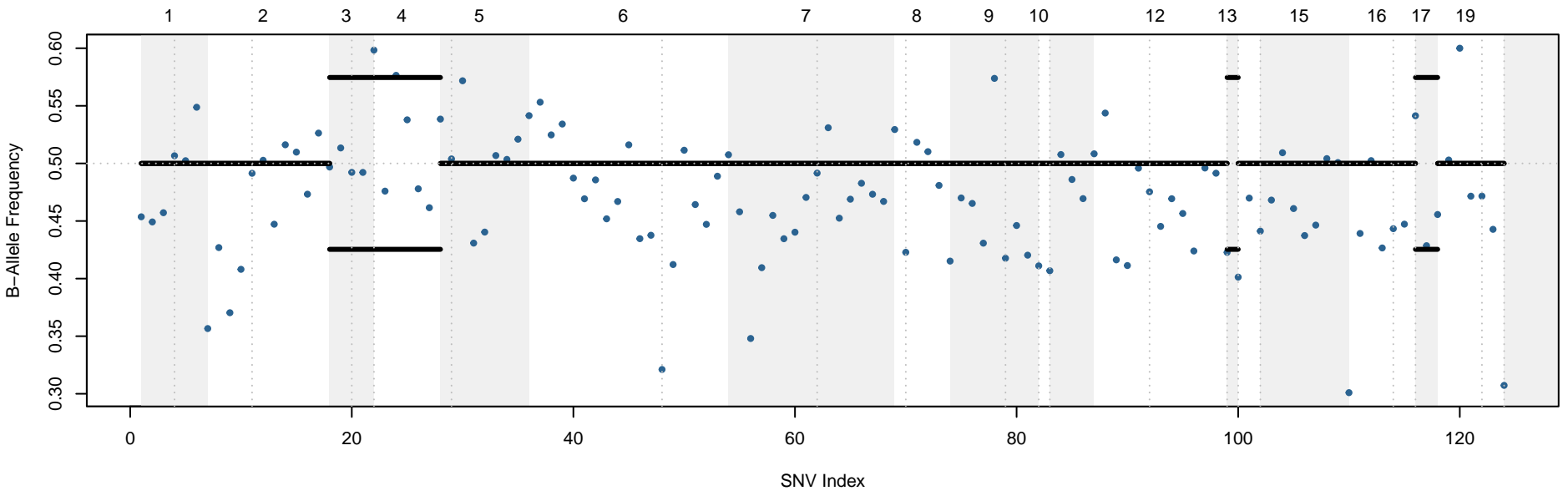




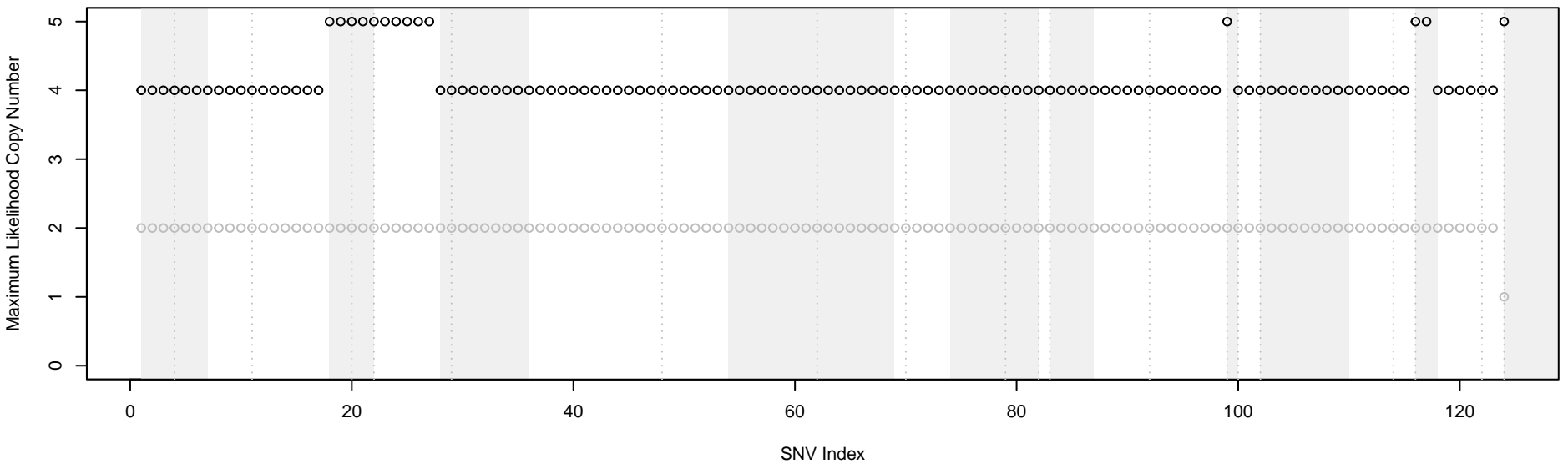
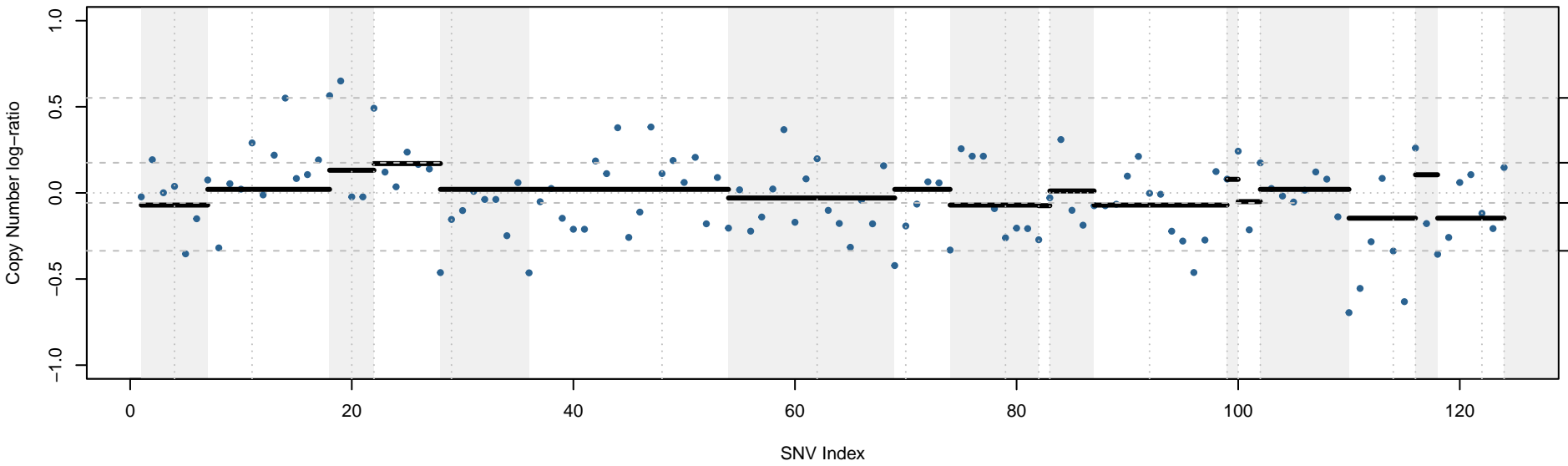
Purity: 0.54 Tumor ploidy: 4.234

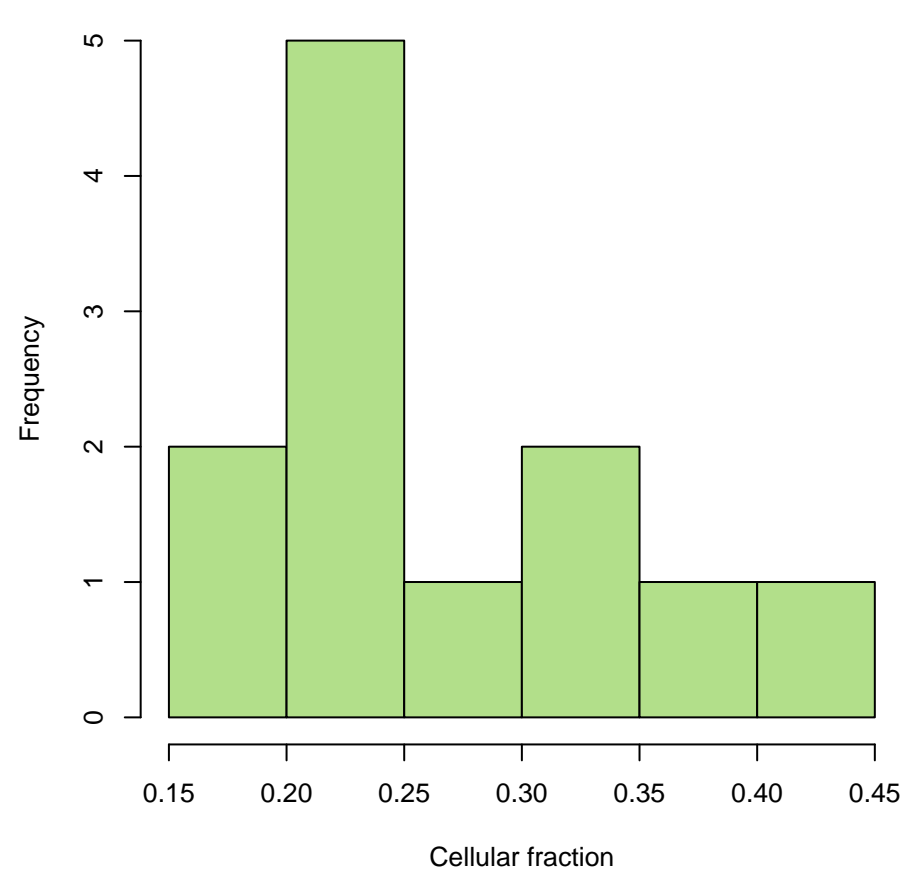
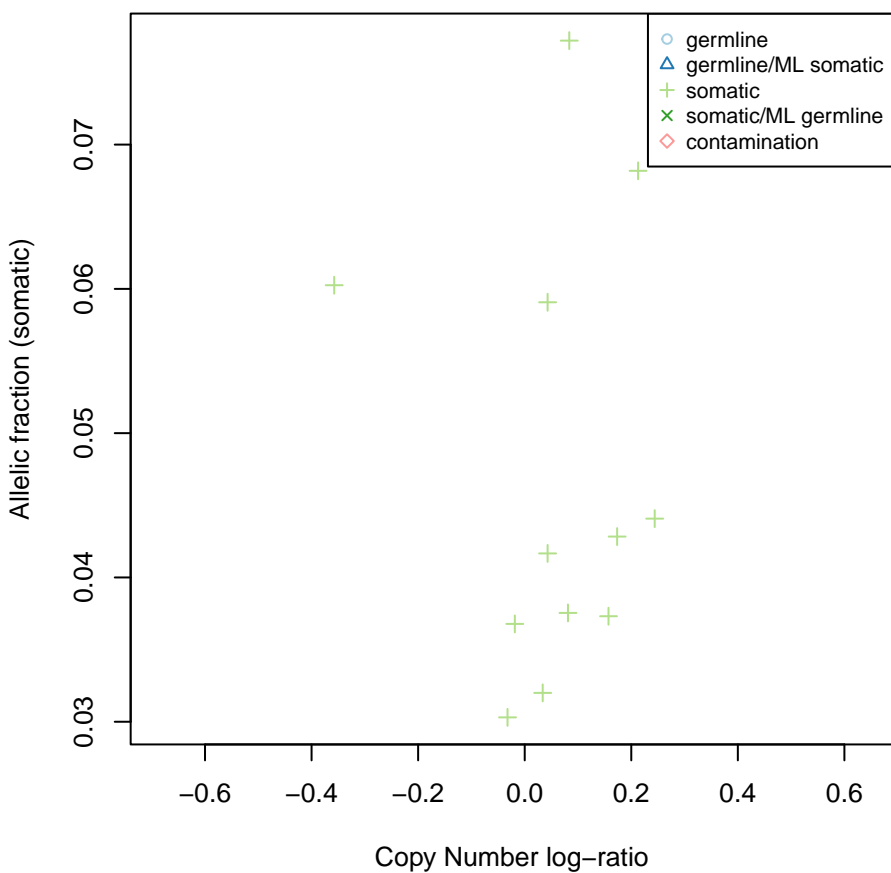
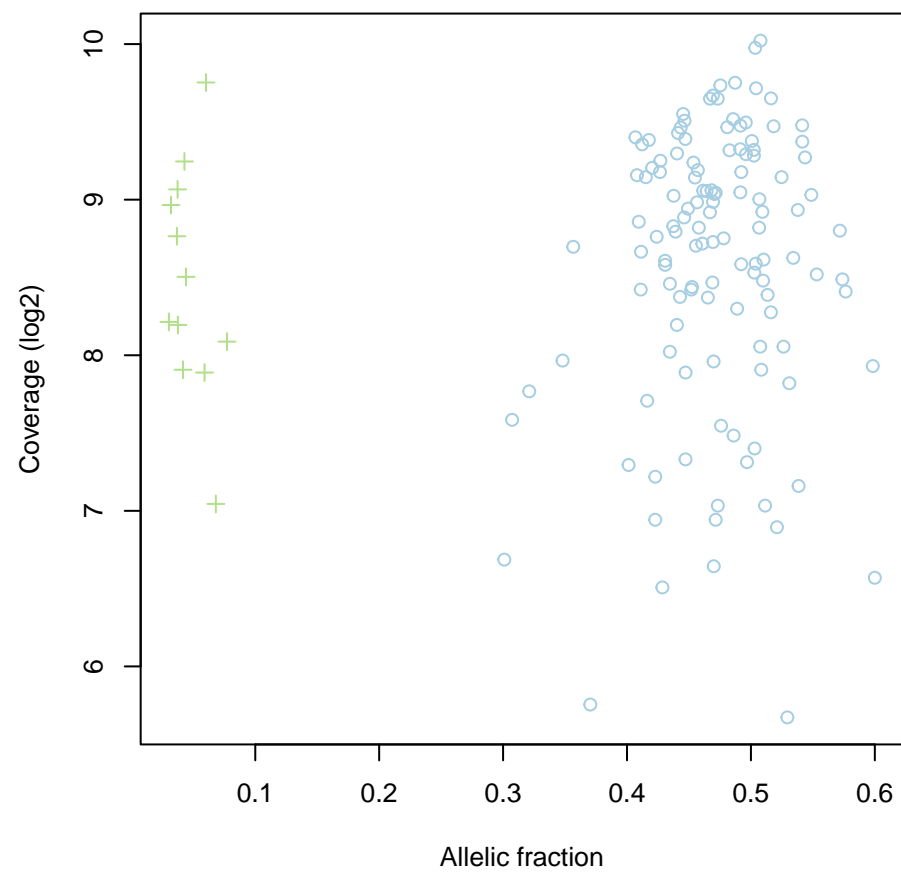
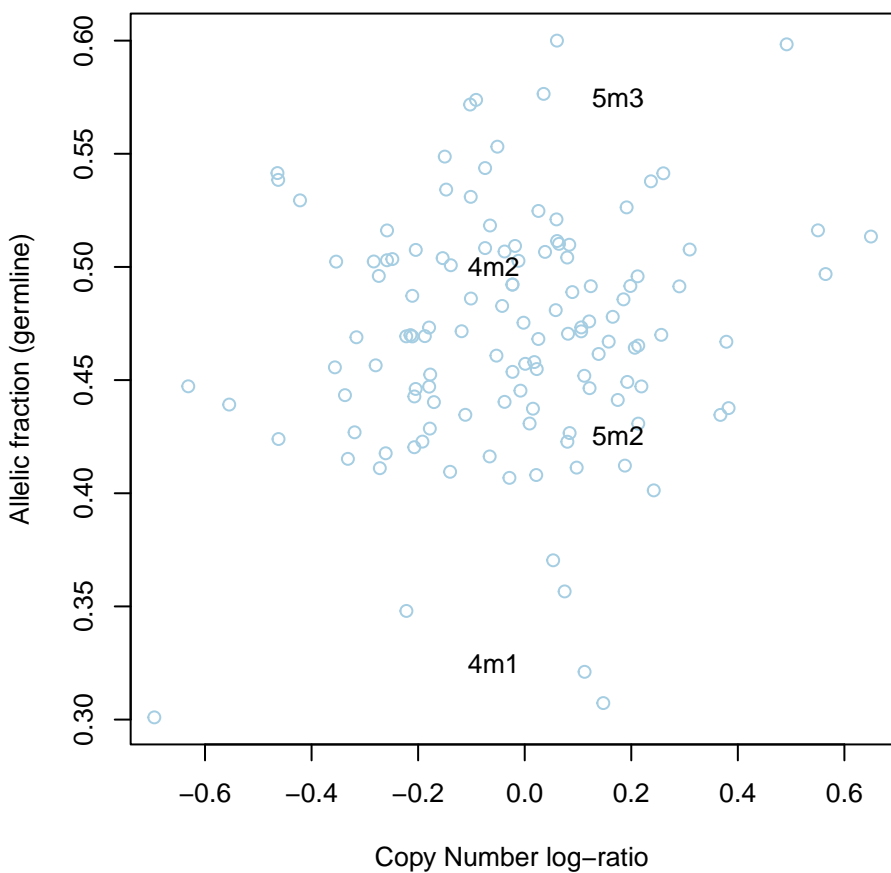


Purity: 0.54 Tumor ploidy: 4.234 SNV log-likelihood: -66.27 GoF: 91.1% Mean coverage: 577;448



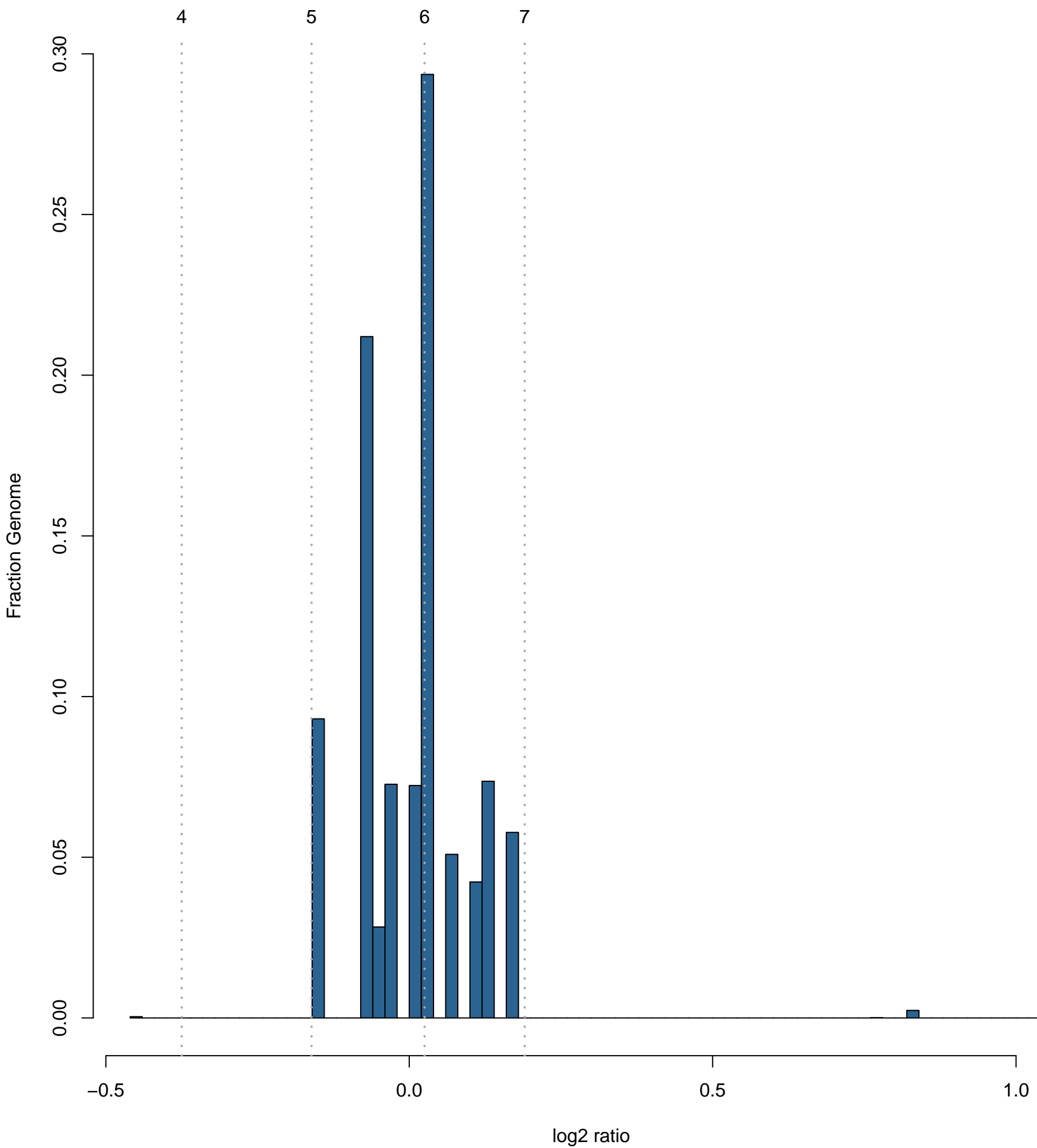
SCNA-fit log-likelihood: -14637.07



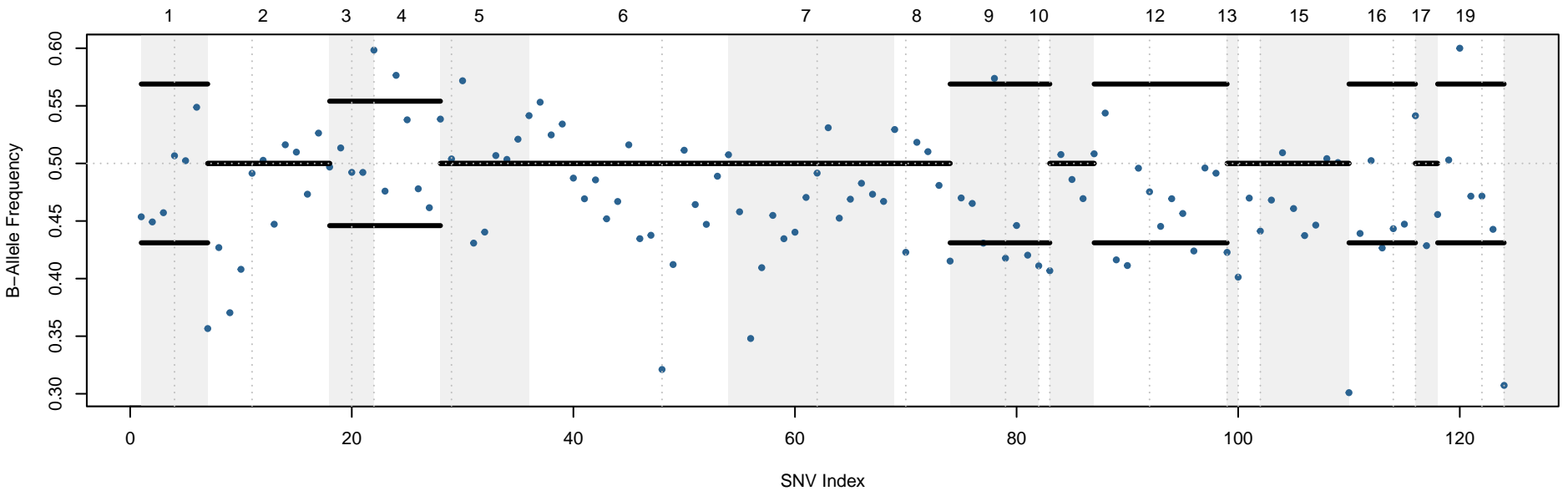




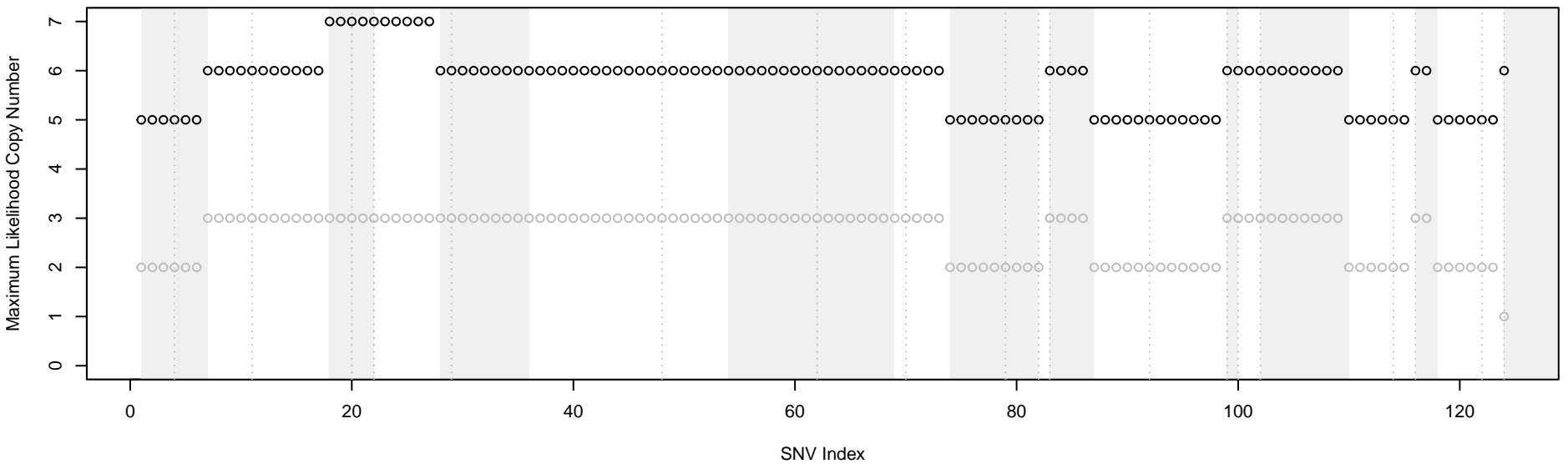
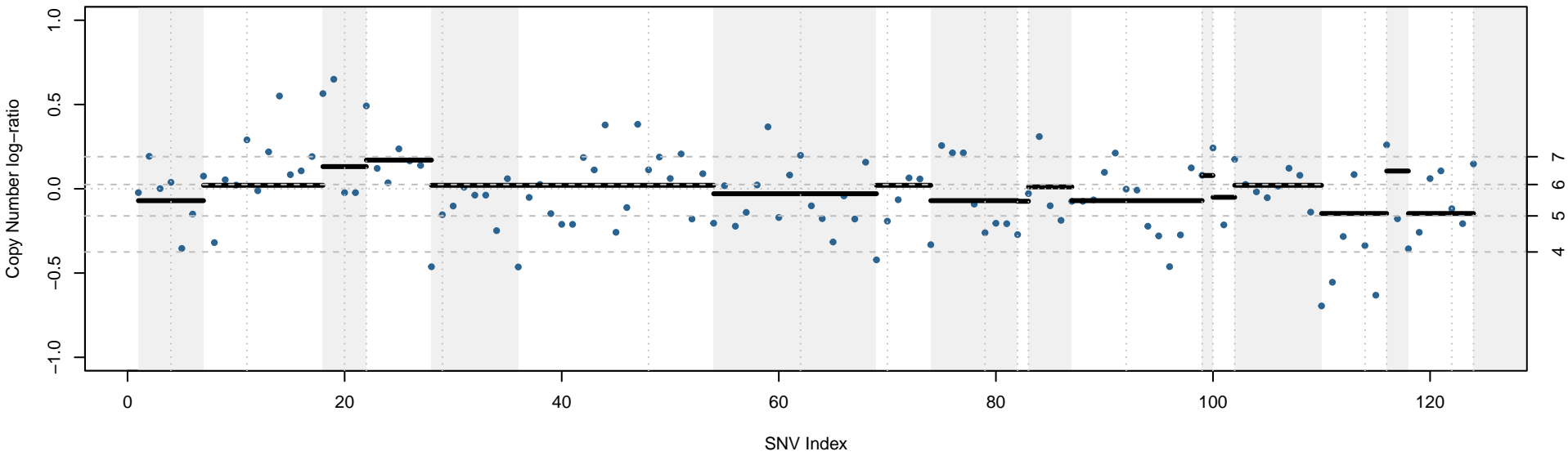
Purity: 0.47 Tumor ploidy: 5.857

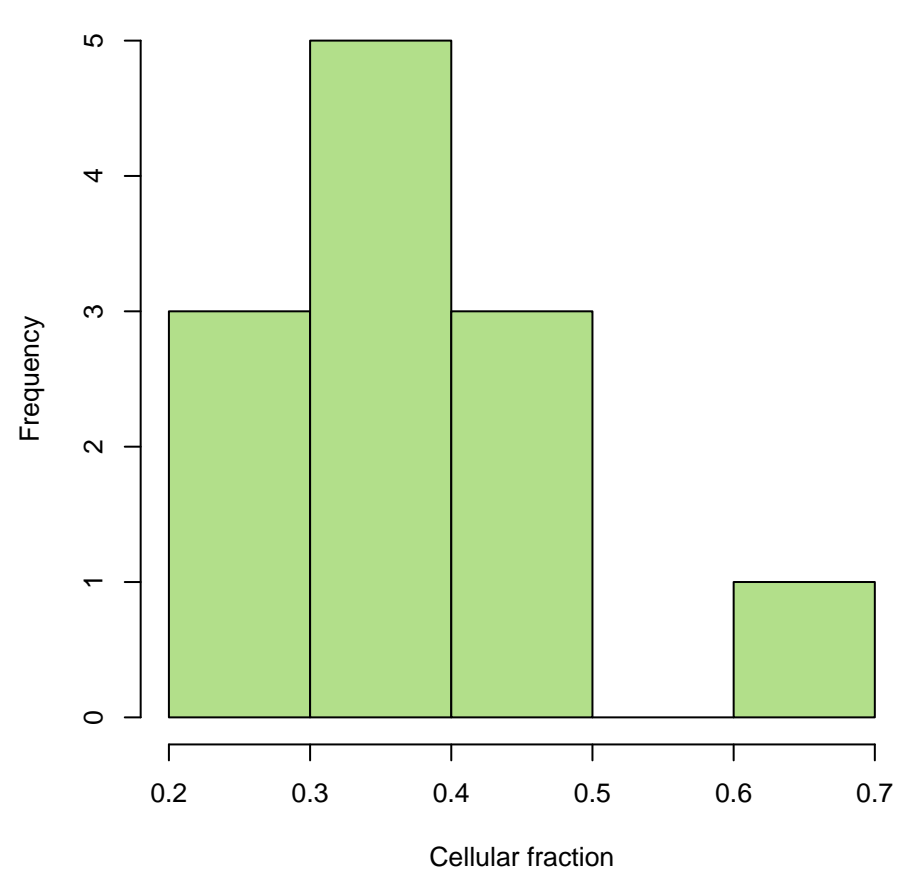
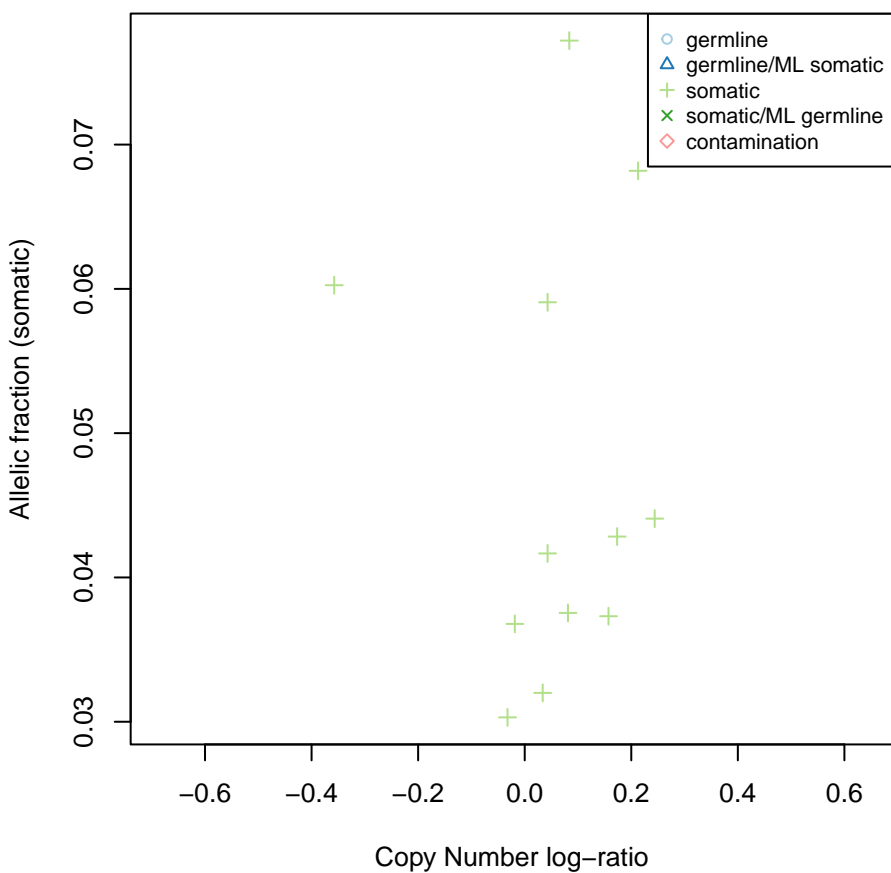
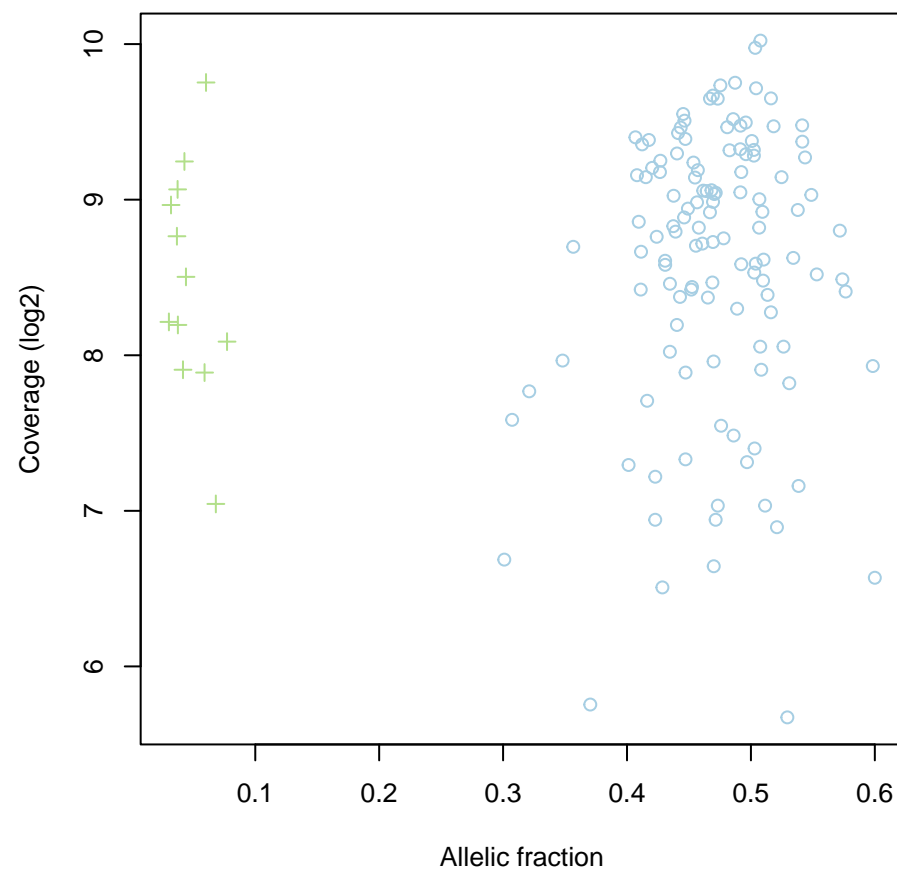
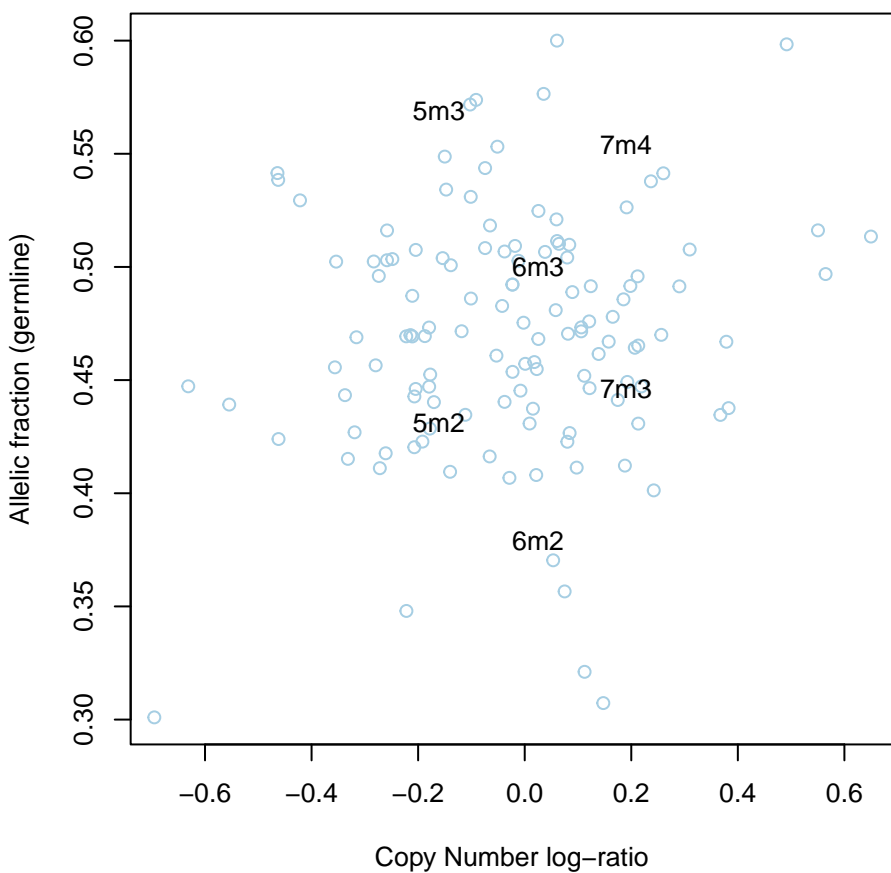


Purity: 0.47 Tumor ploidy: 5.857 SNV log-likelihood: -93.26 GoF: 94.4% Mean coverage: 577;448

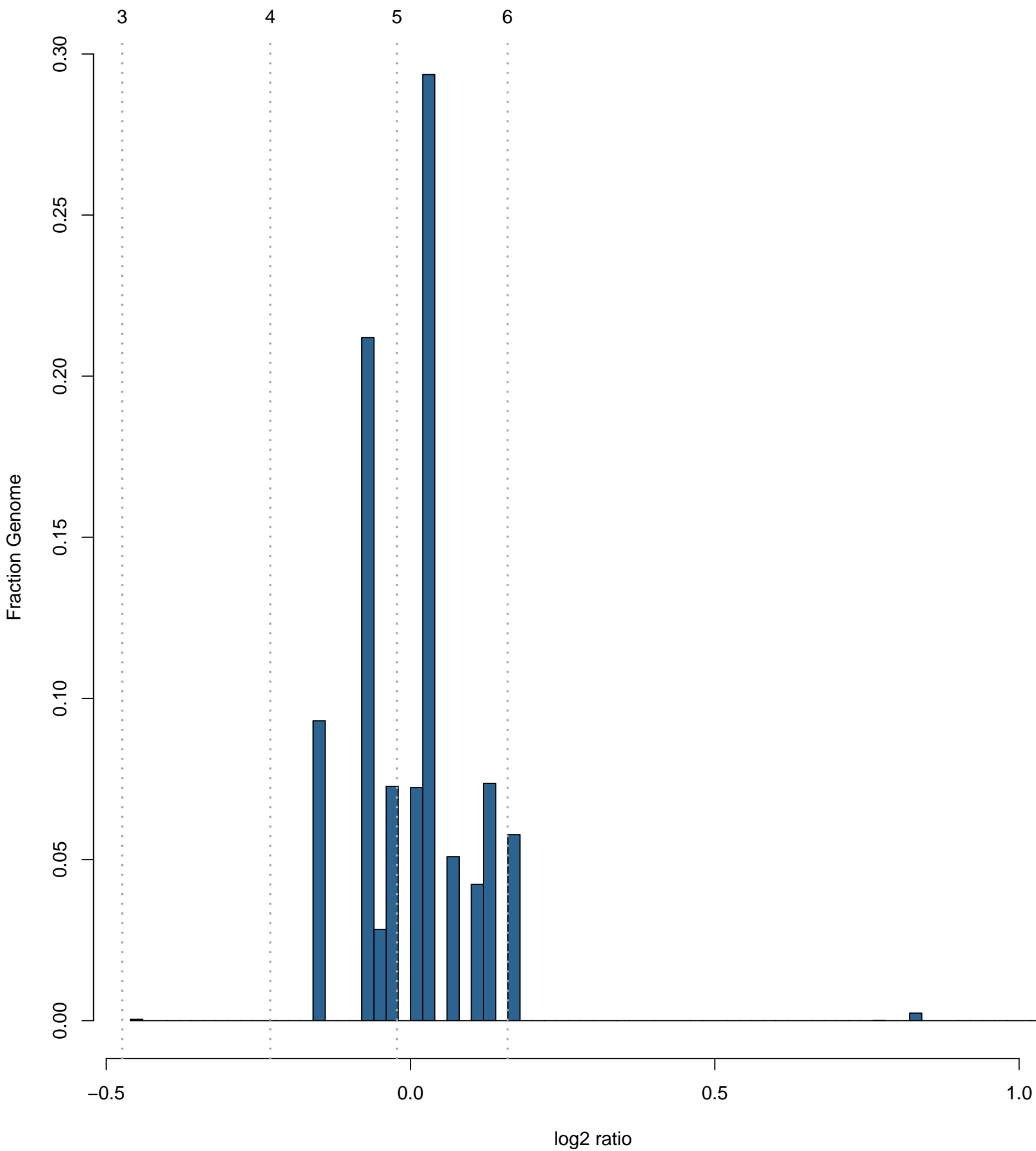


SCNA-fit log-likelihood: -14588.54

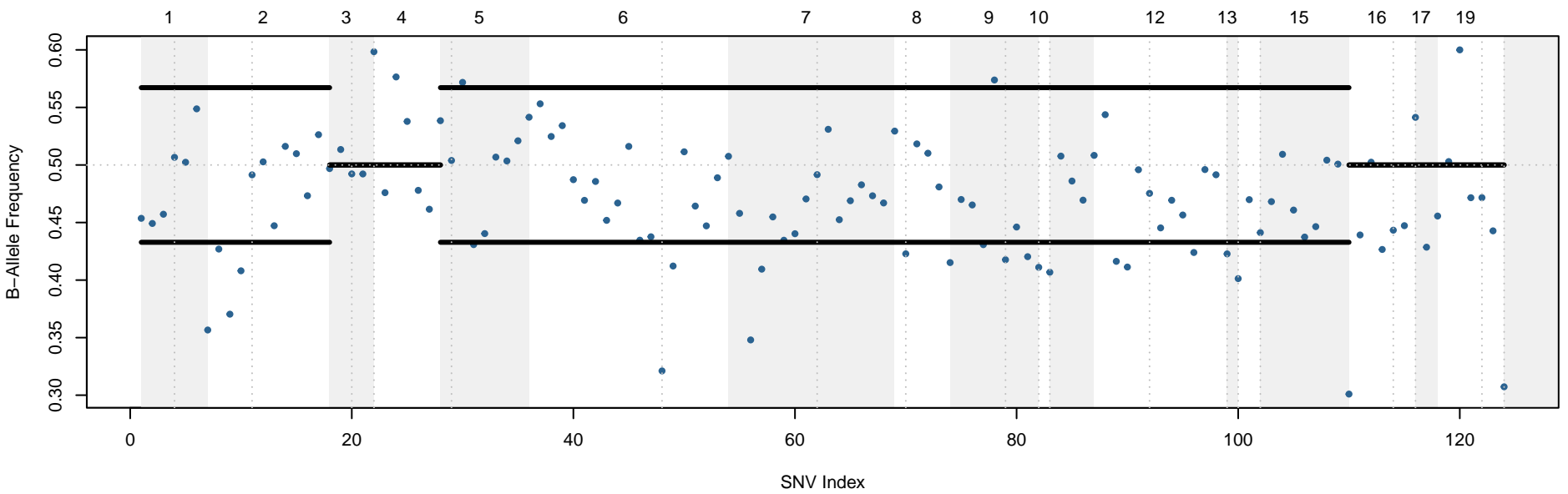




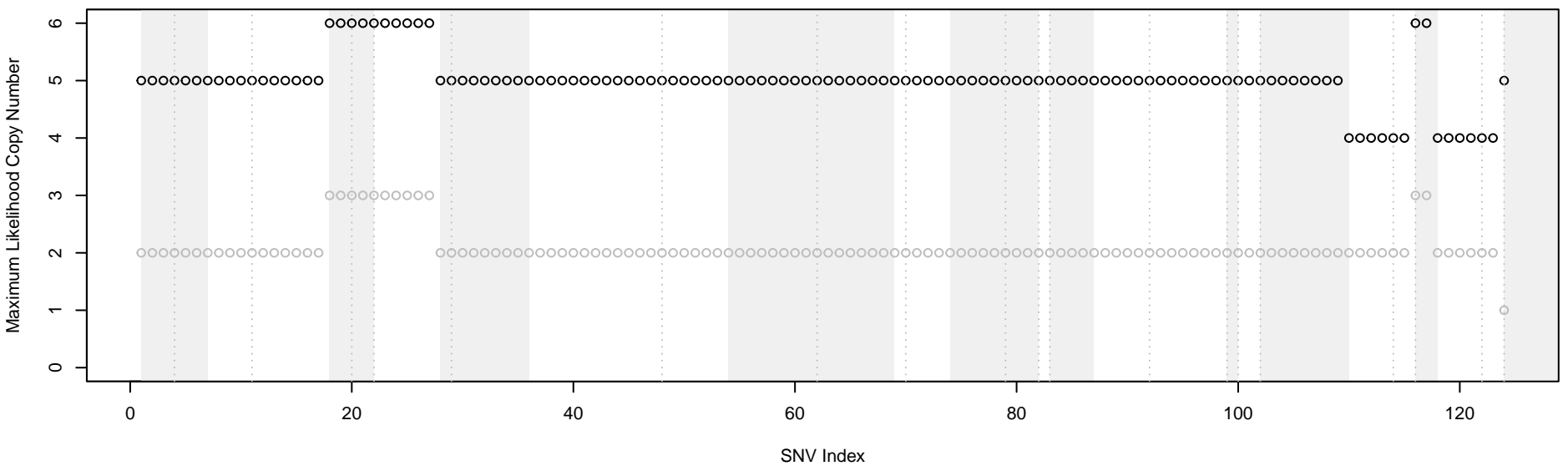
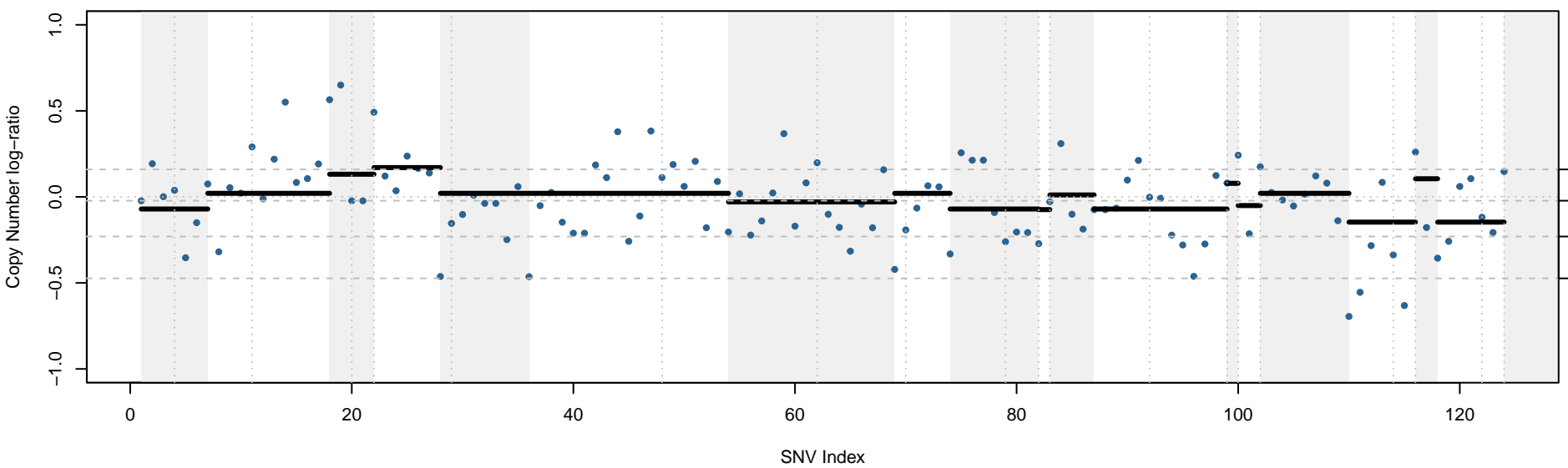
Purity: 0.45 Tumor ploidy: 5.116

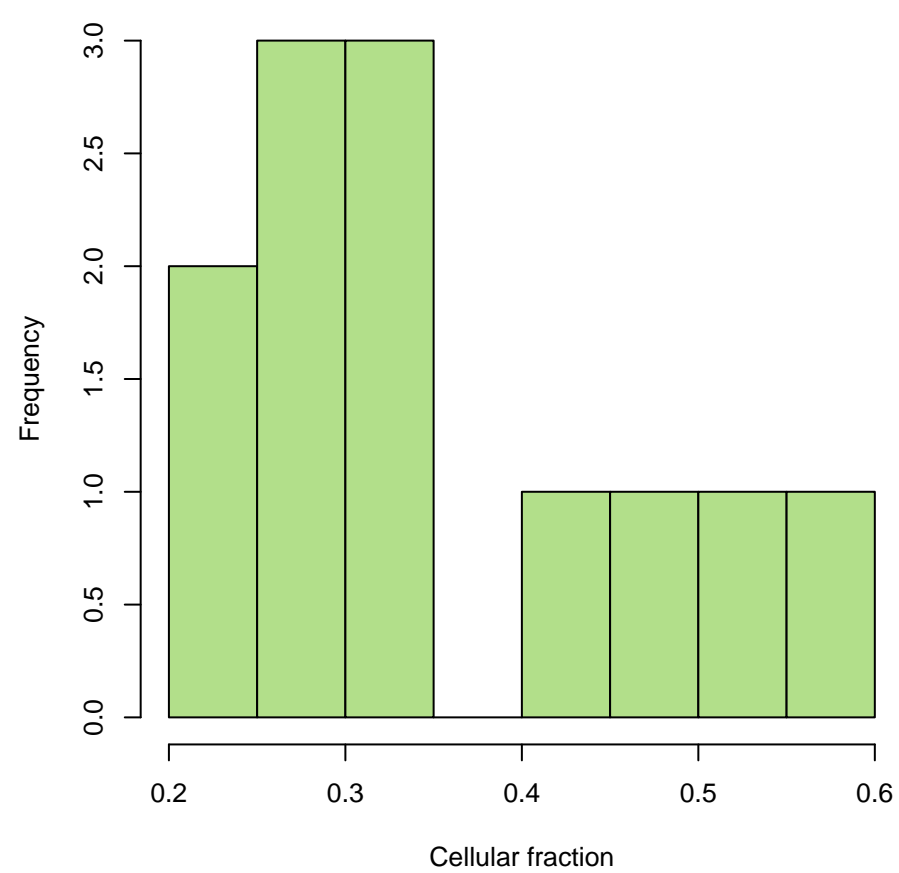
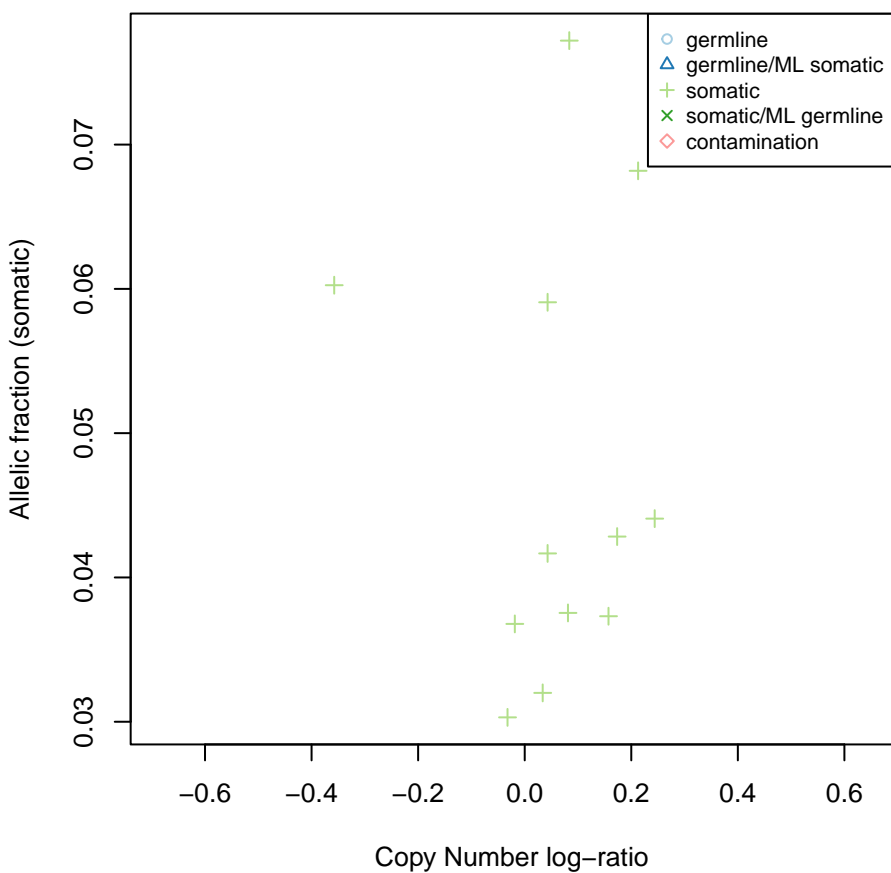
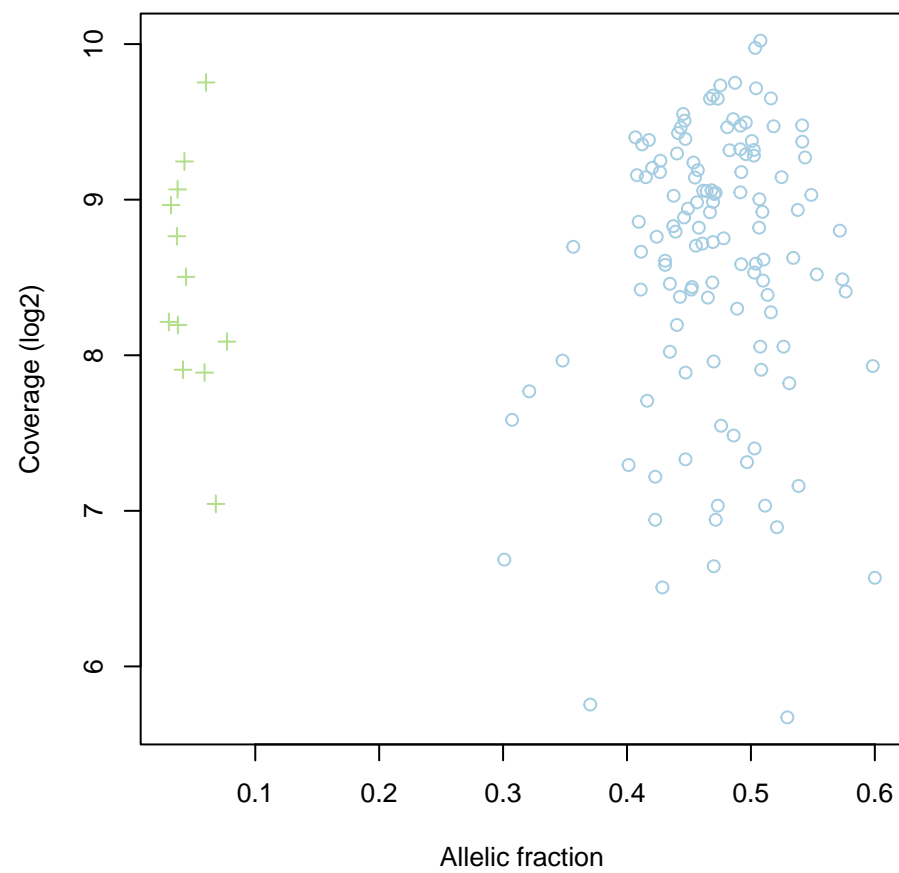
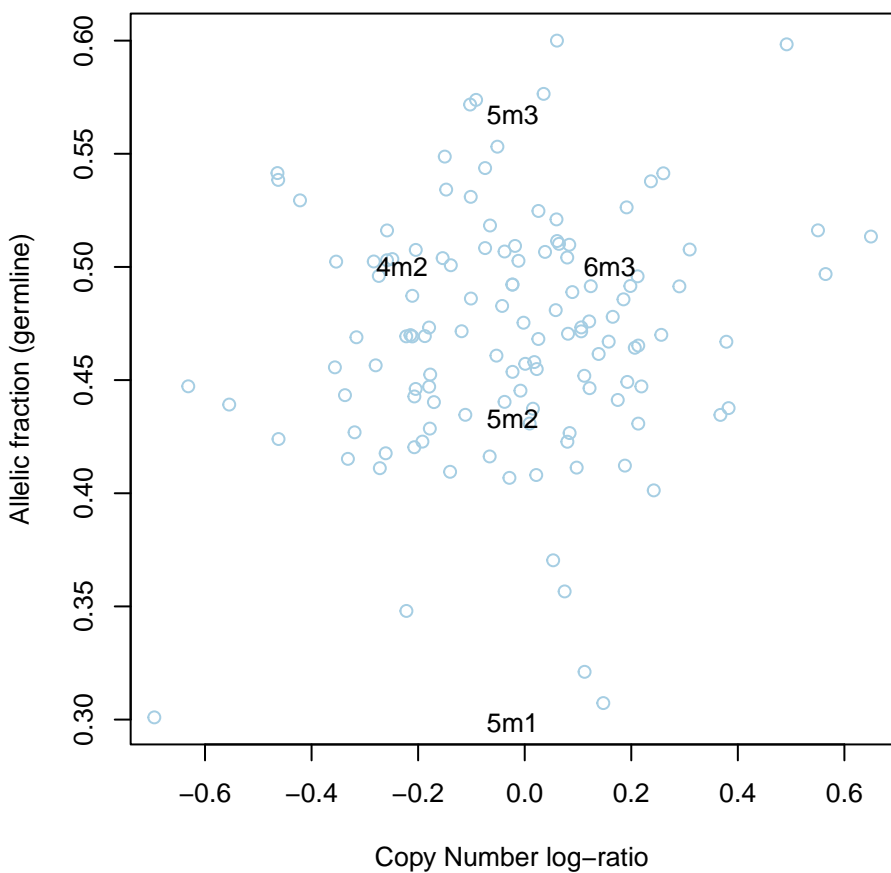


Purity: 0.45 Tumor ploidy: 5.116 SNV log-likelihood: -106.78 GoF: 93.5% Mean coverage: 577,448

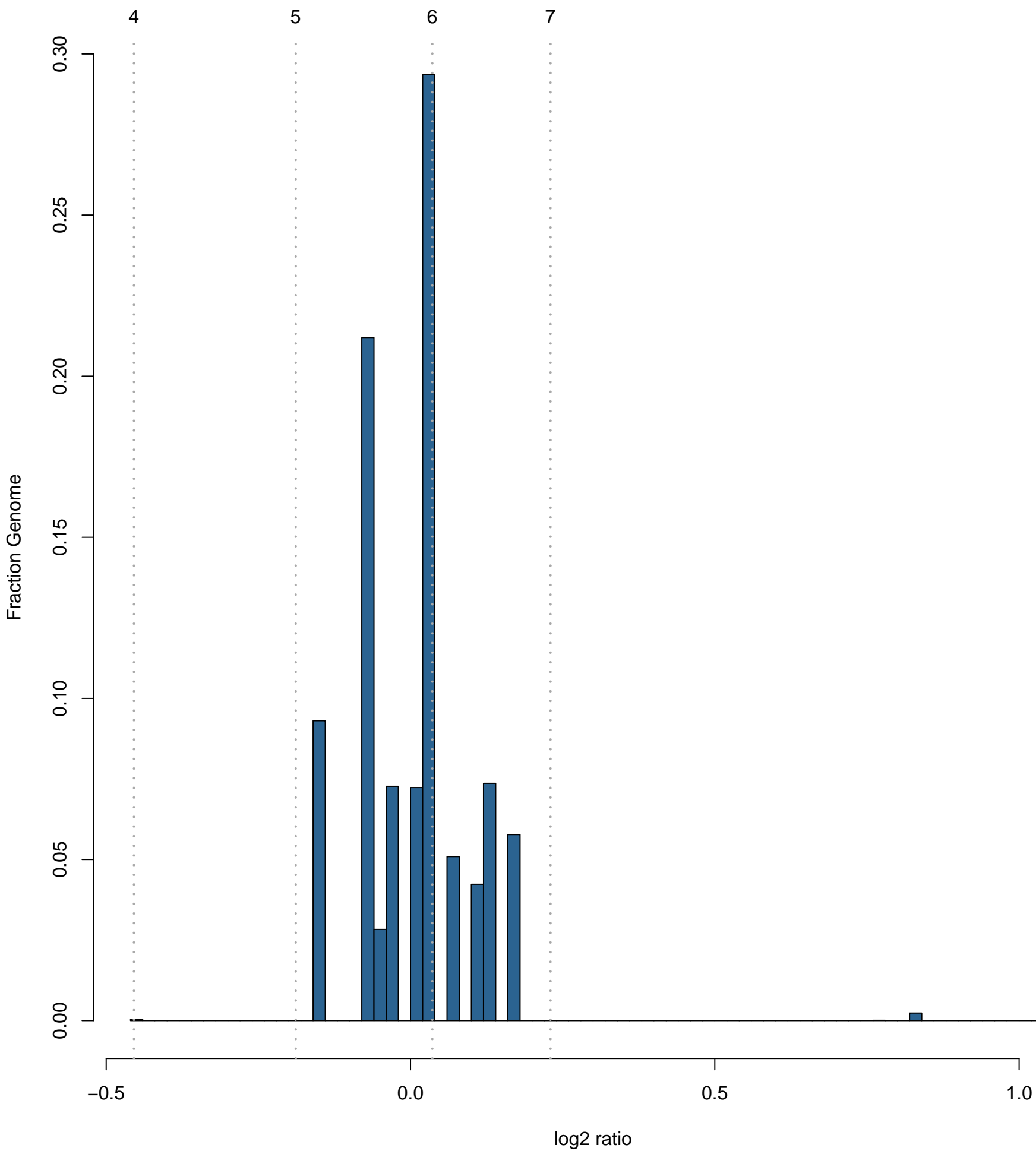


SCNA-fit log-likelihood: -14573.45

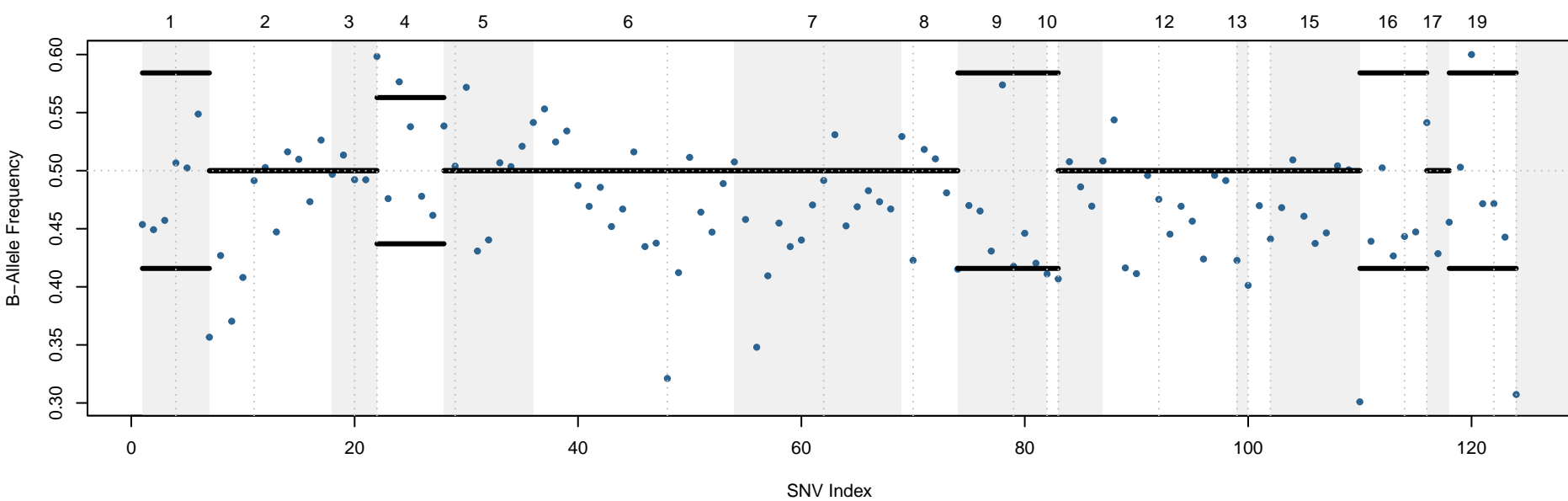




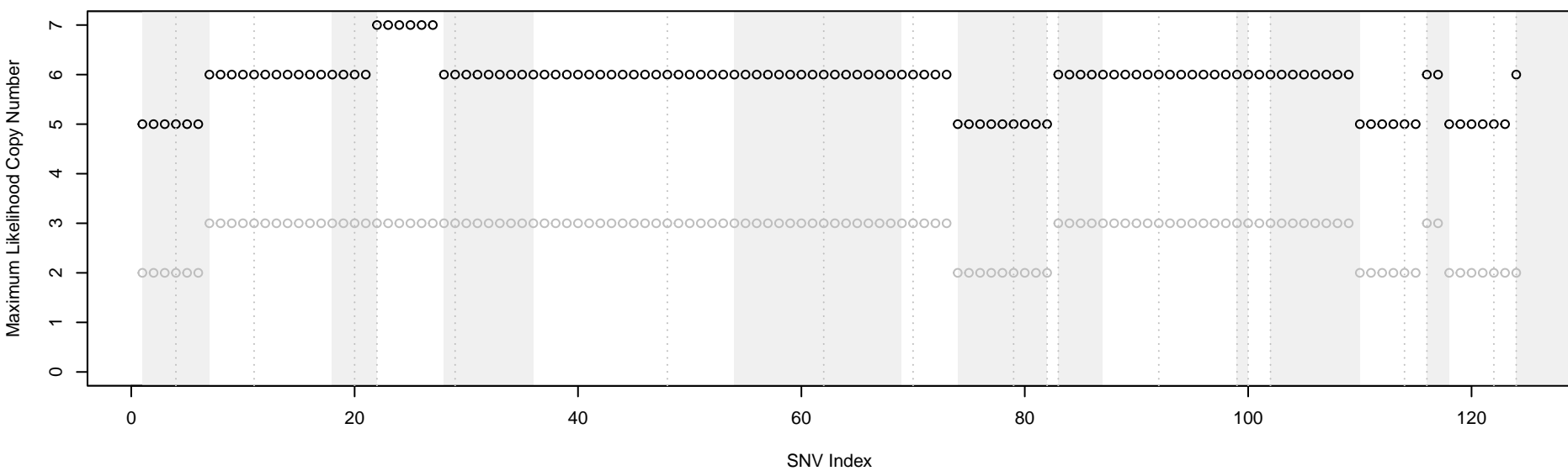
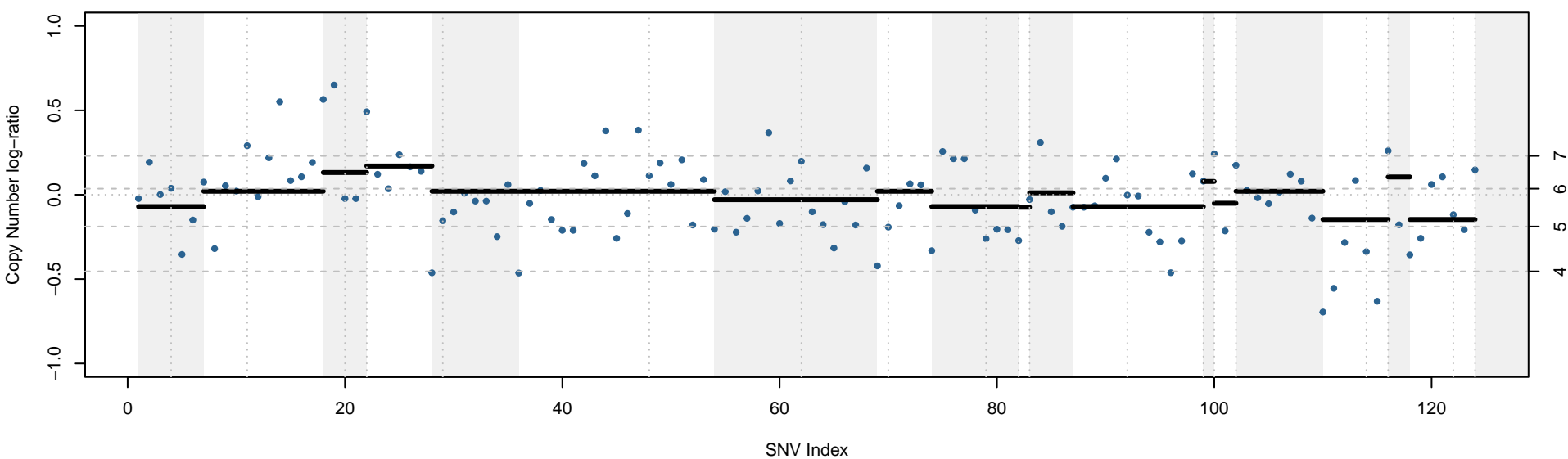
Purity: 0.68 Tumor ploidy: 5.829



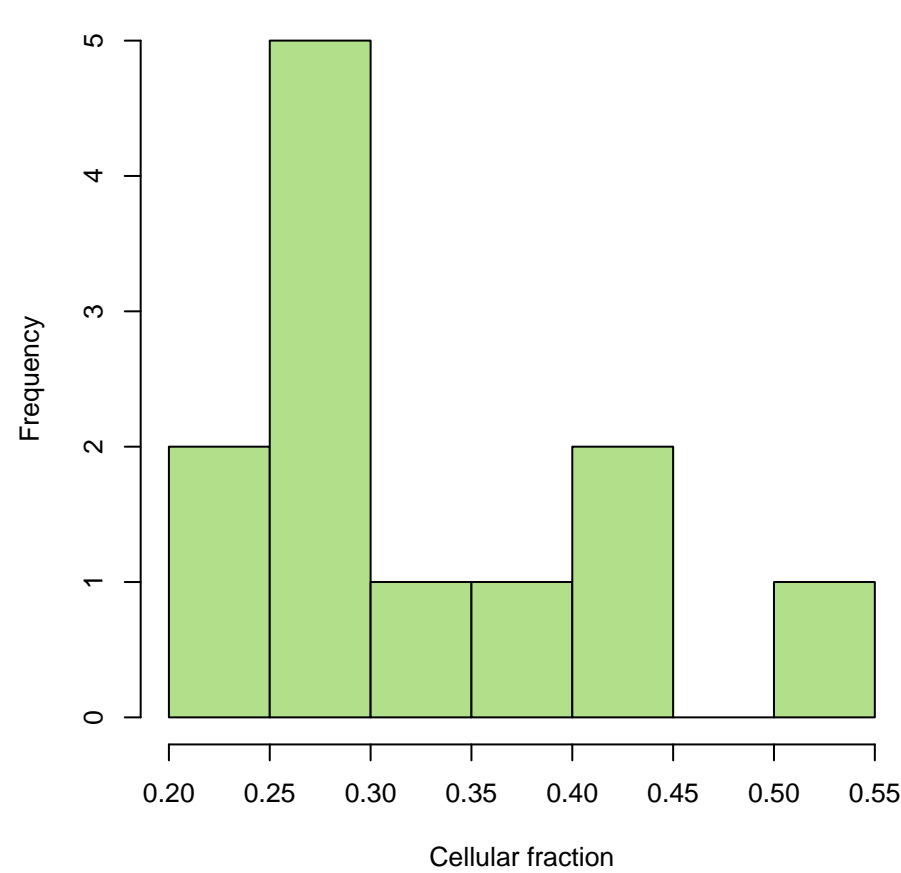
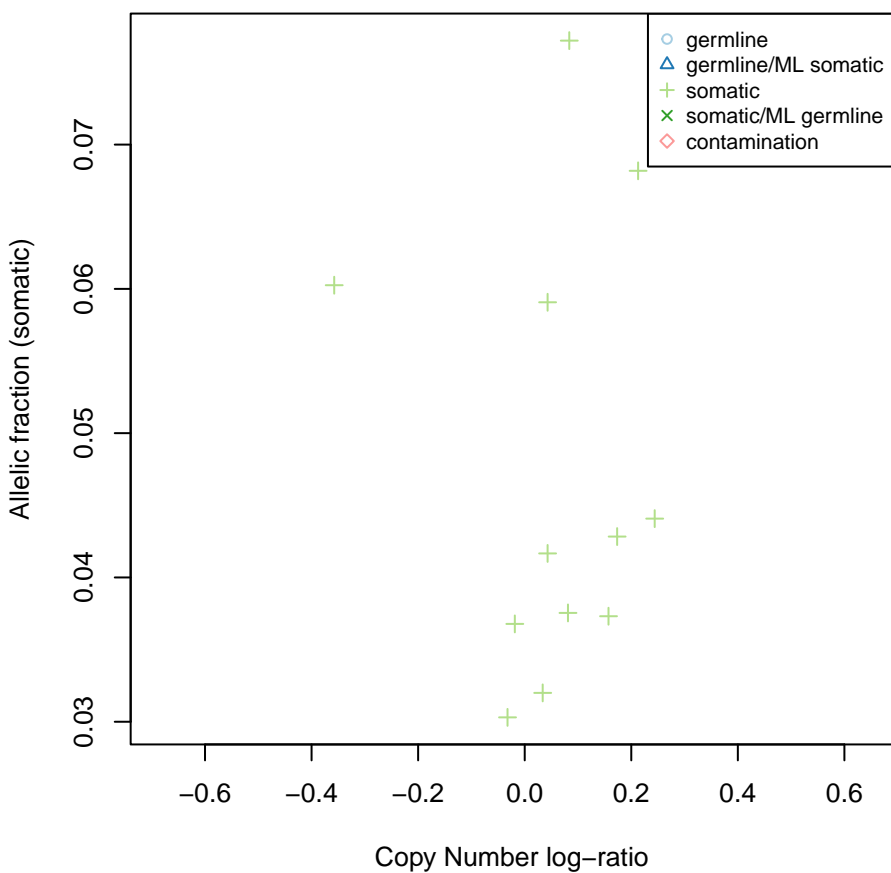
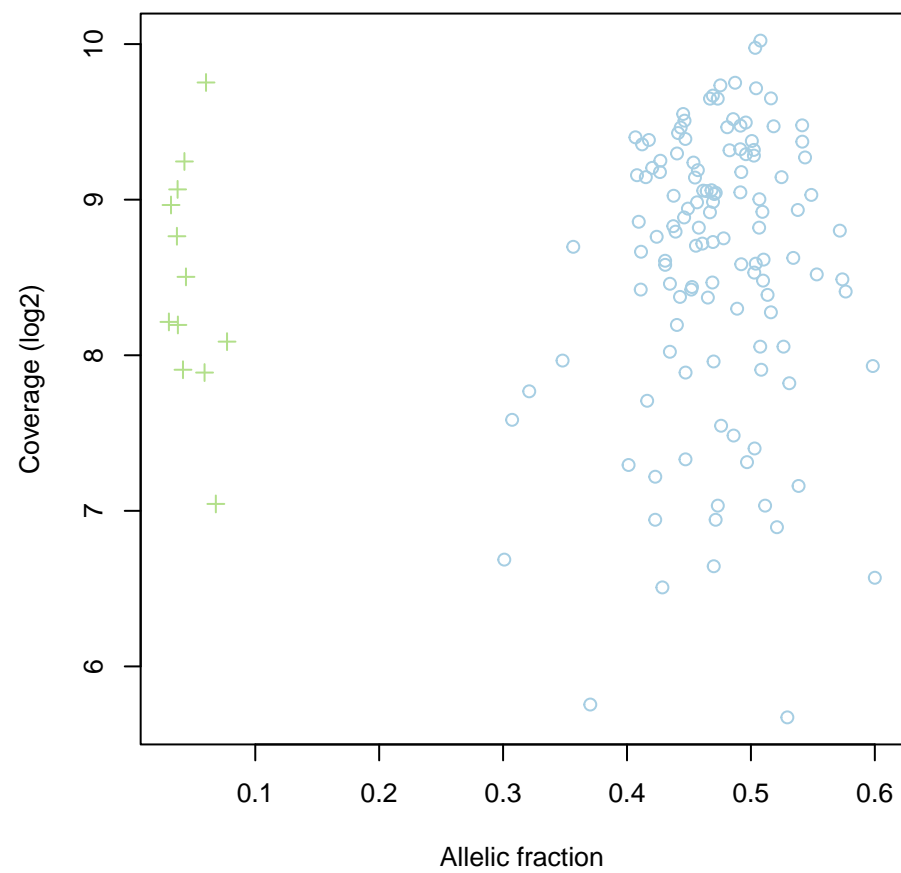
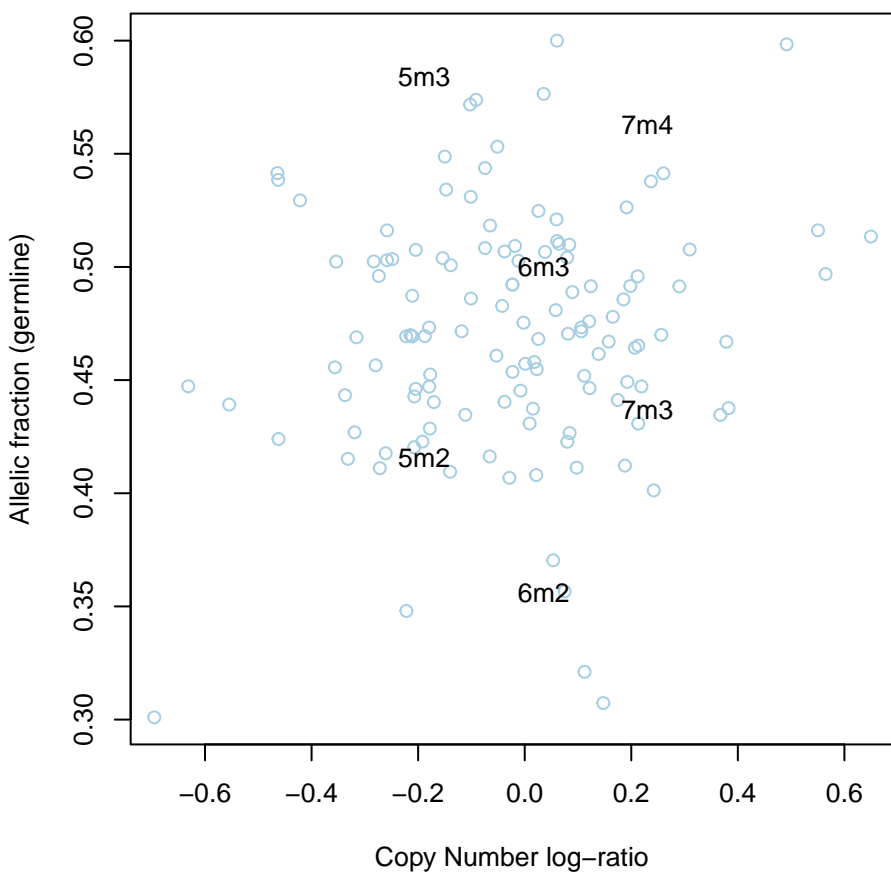
Purity: 0.68 Tumor ploidy: 5.829 SNV log-likelihood: -91.25 GoF: 93.2% Mean coverage: 577;448



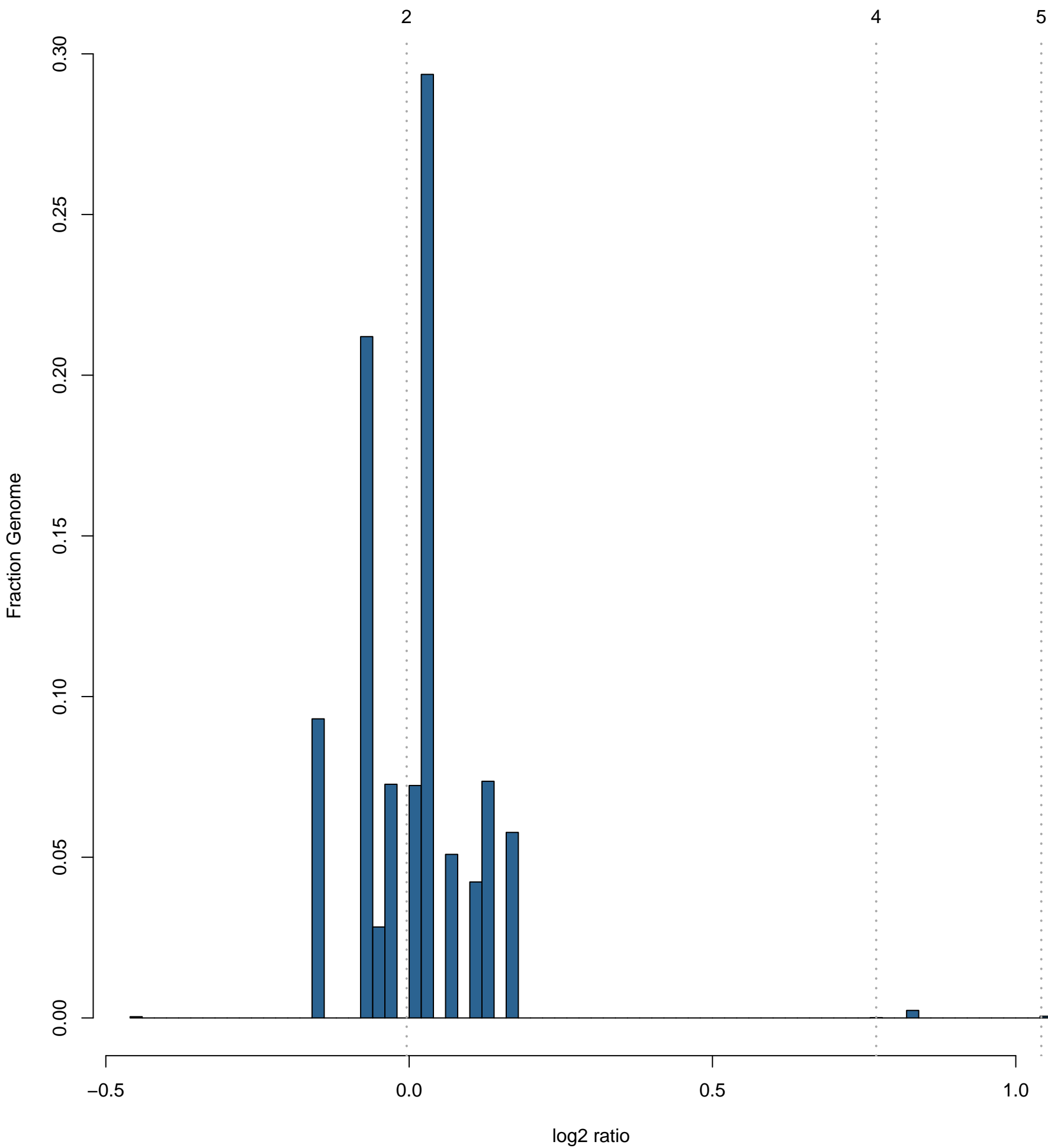
SCNA-fit log-likelihood: -14656.18



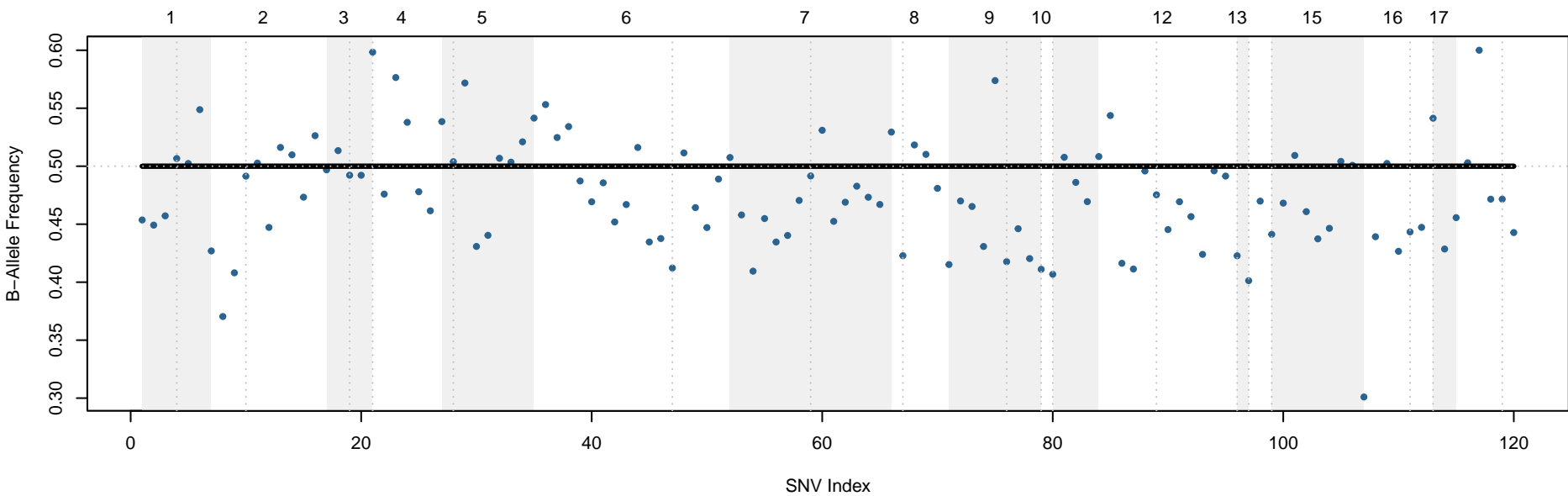




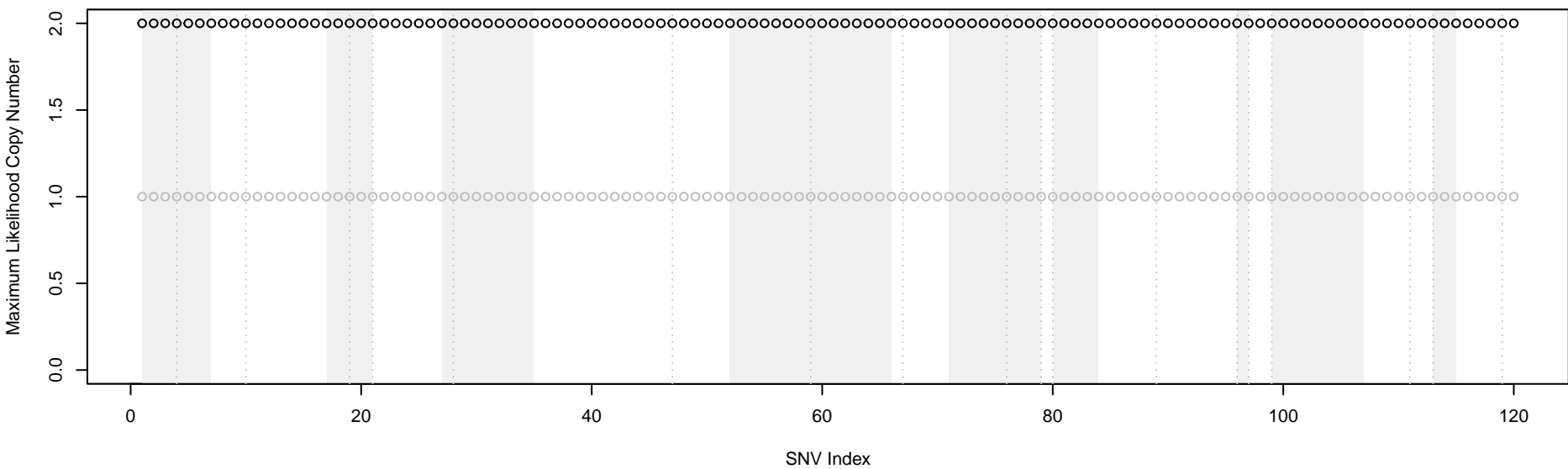
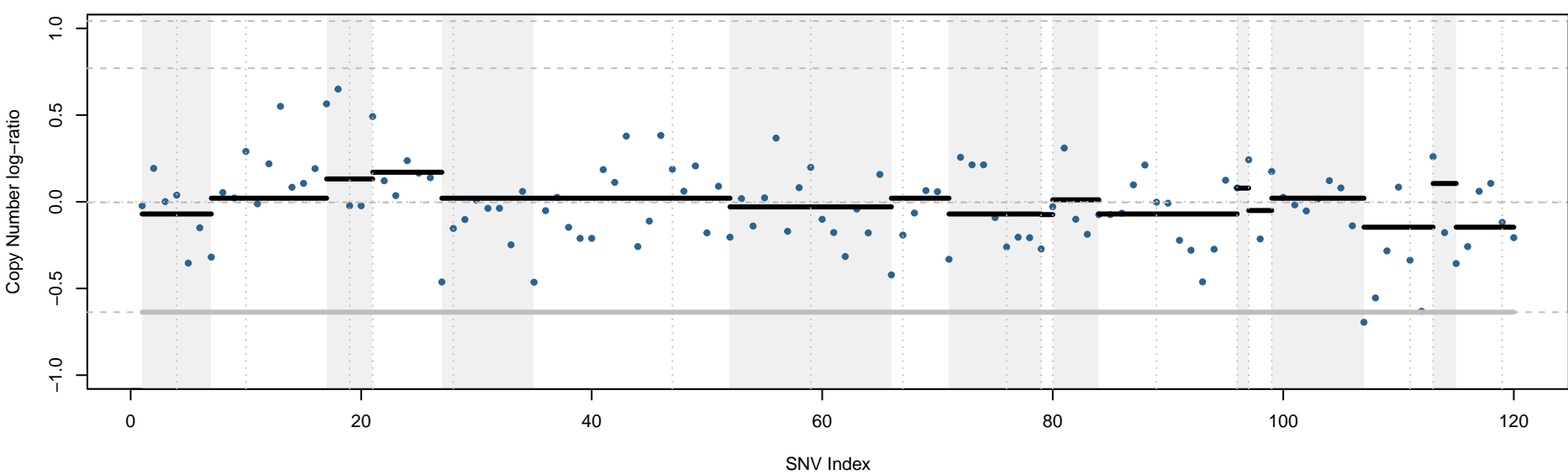
Purity: 0.71 Tumor ploidy: 2.008

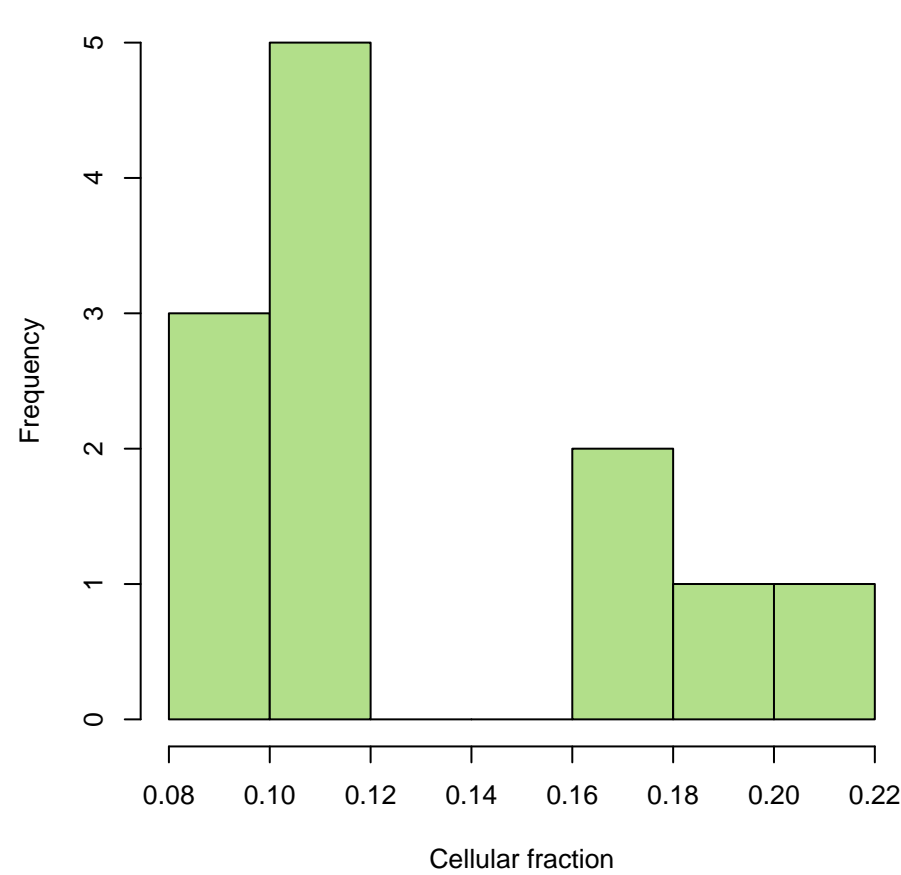
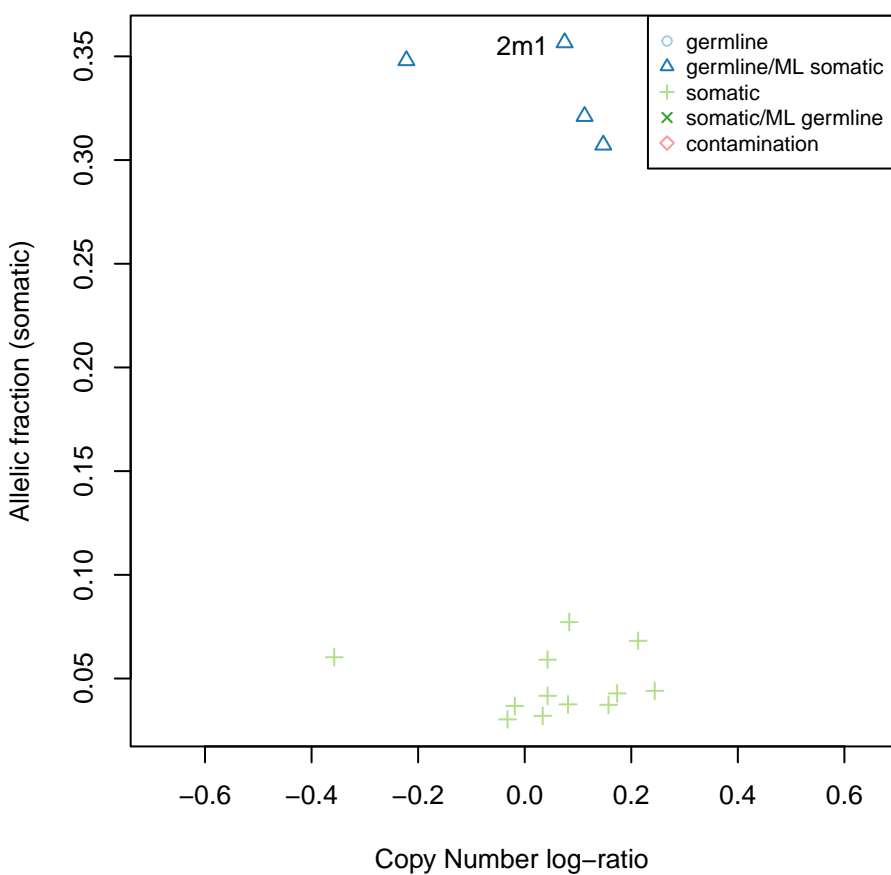
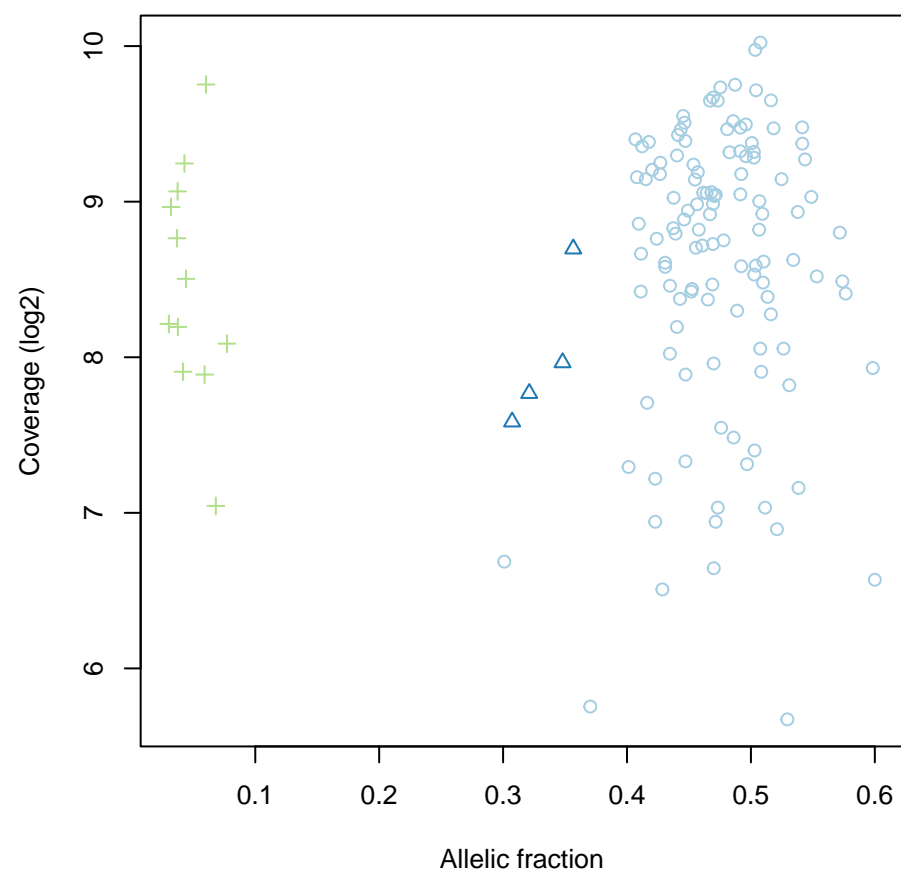
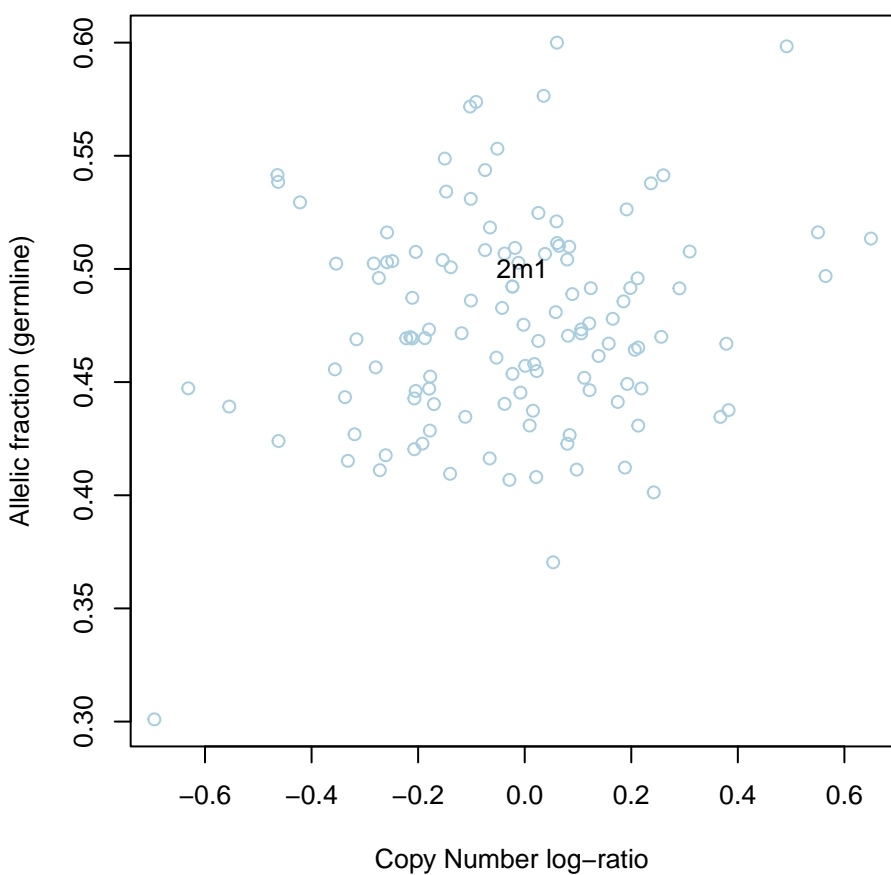


Purity: 0.71 Tumor ploidy: 2.008 SNV log-likelihood: -73.38 GoF: 73.5% Mean coverage: 577;448

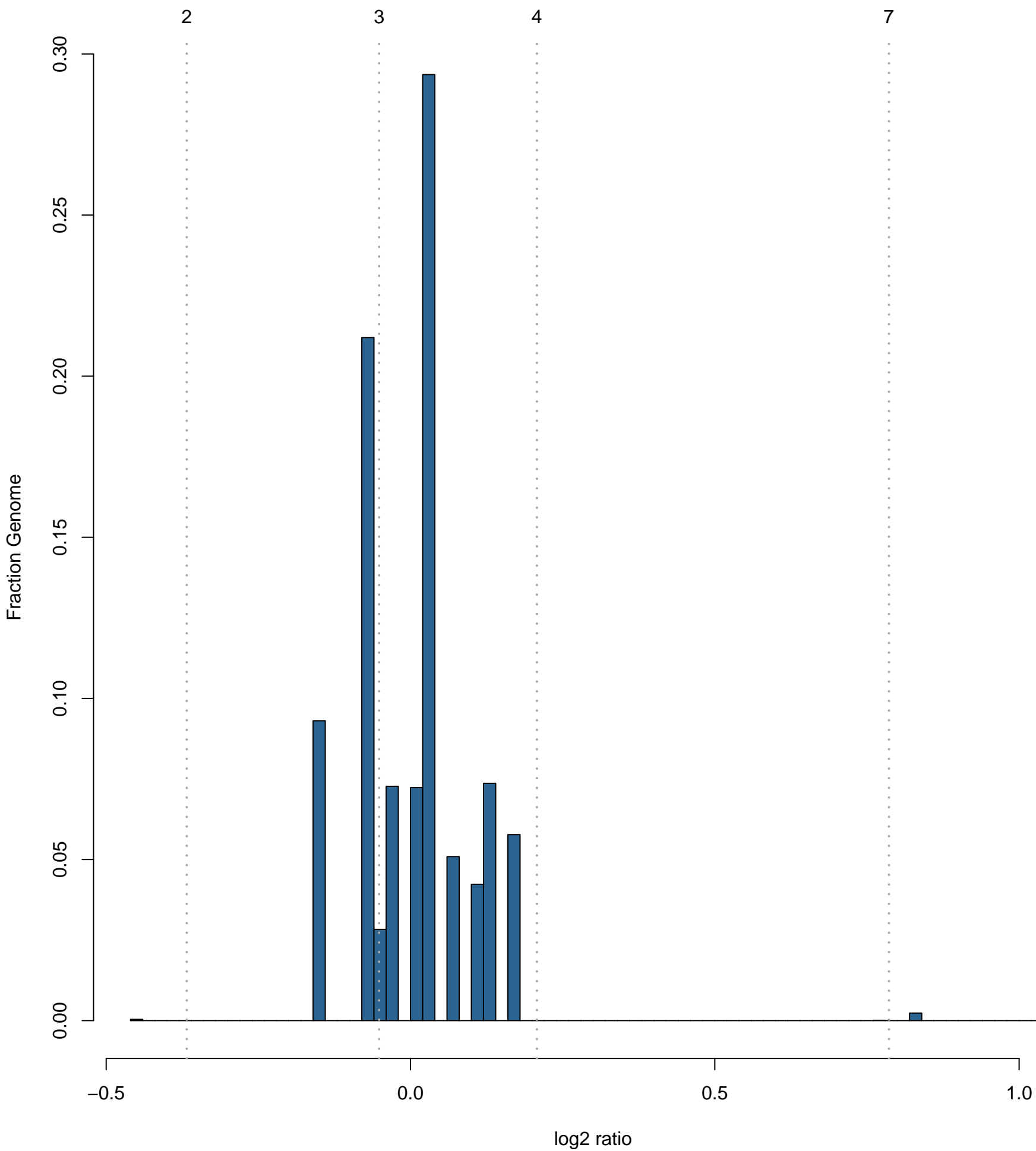


SCNA-fit log-likelihood: -14735.1

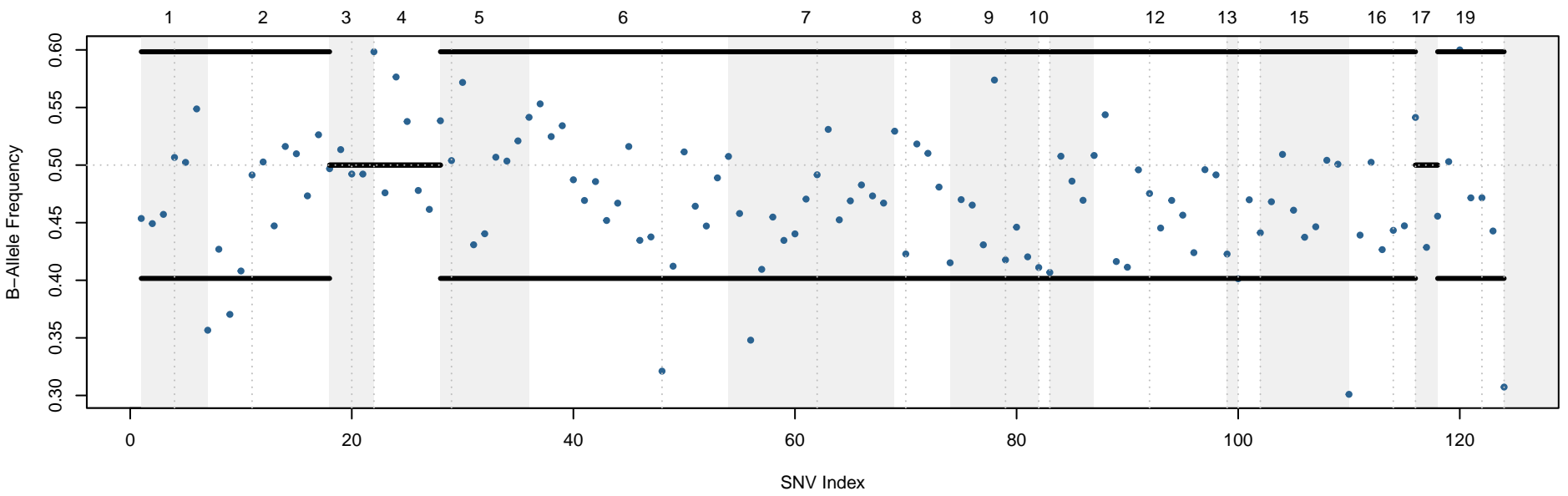




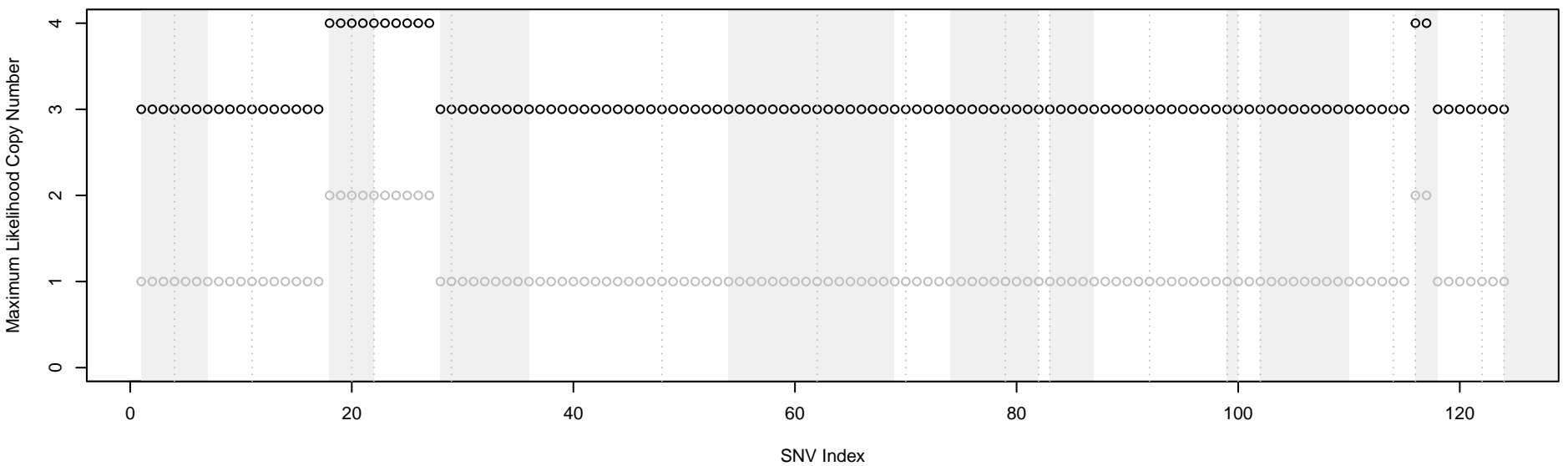
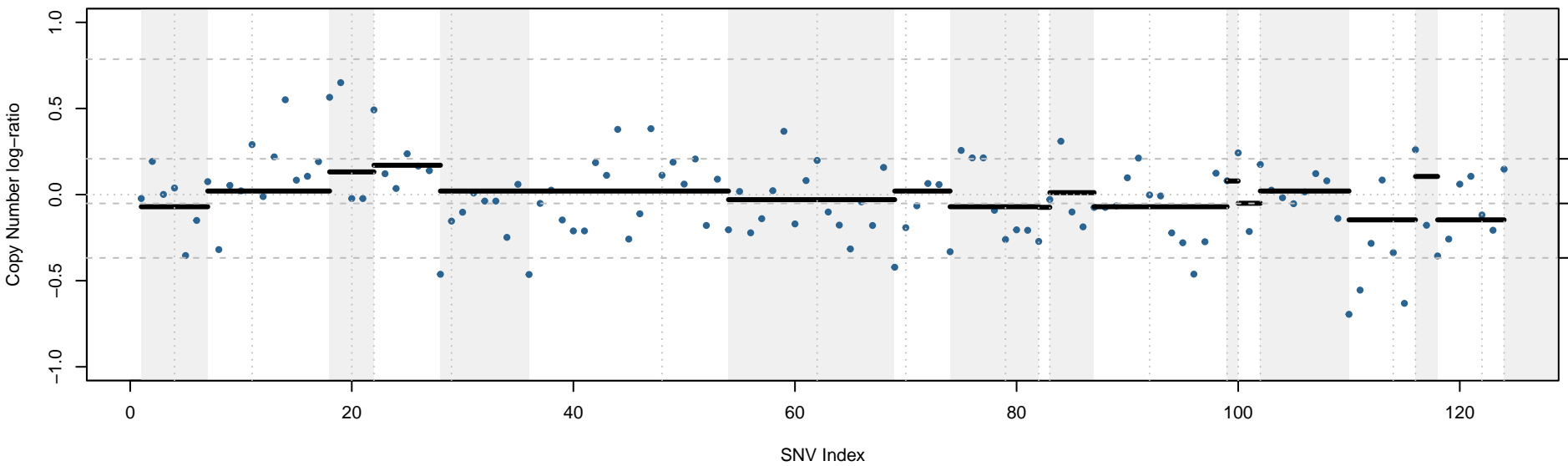
Purity: 0.49 Tumor ploidy: 3.185

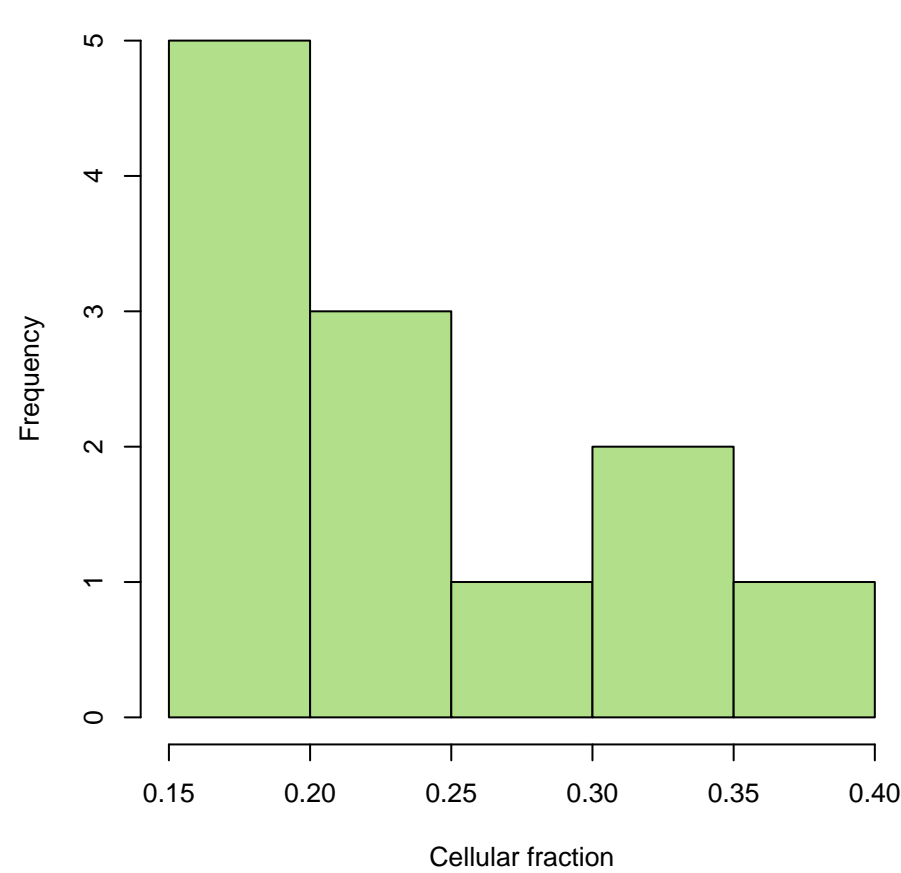
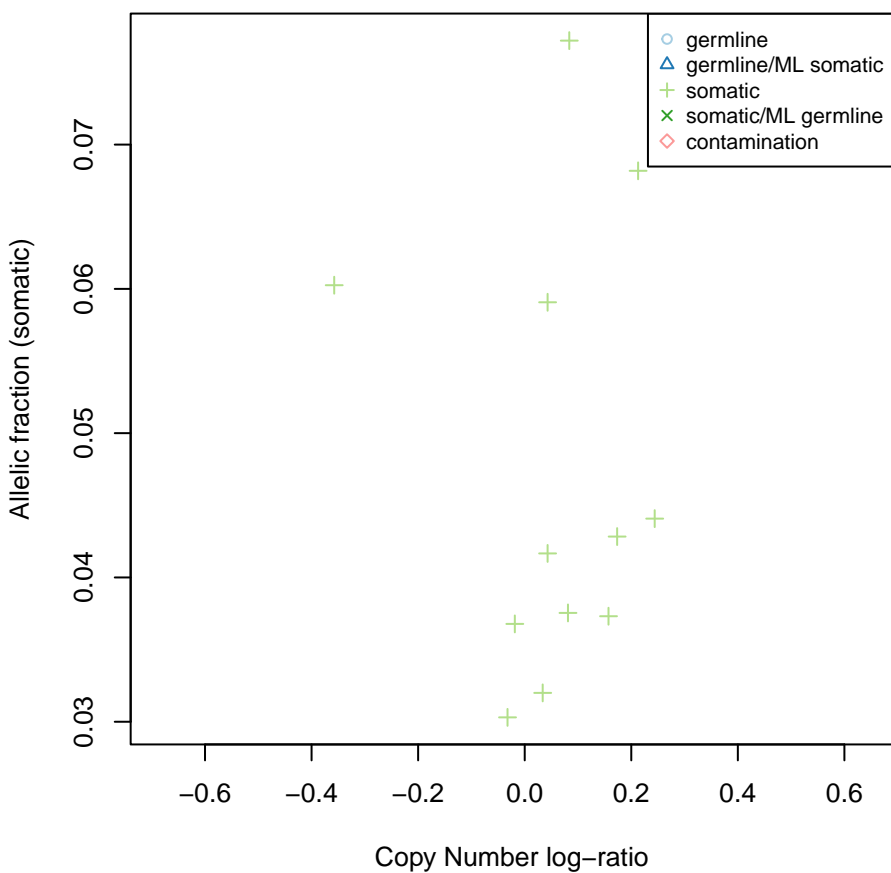
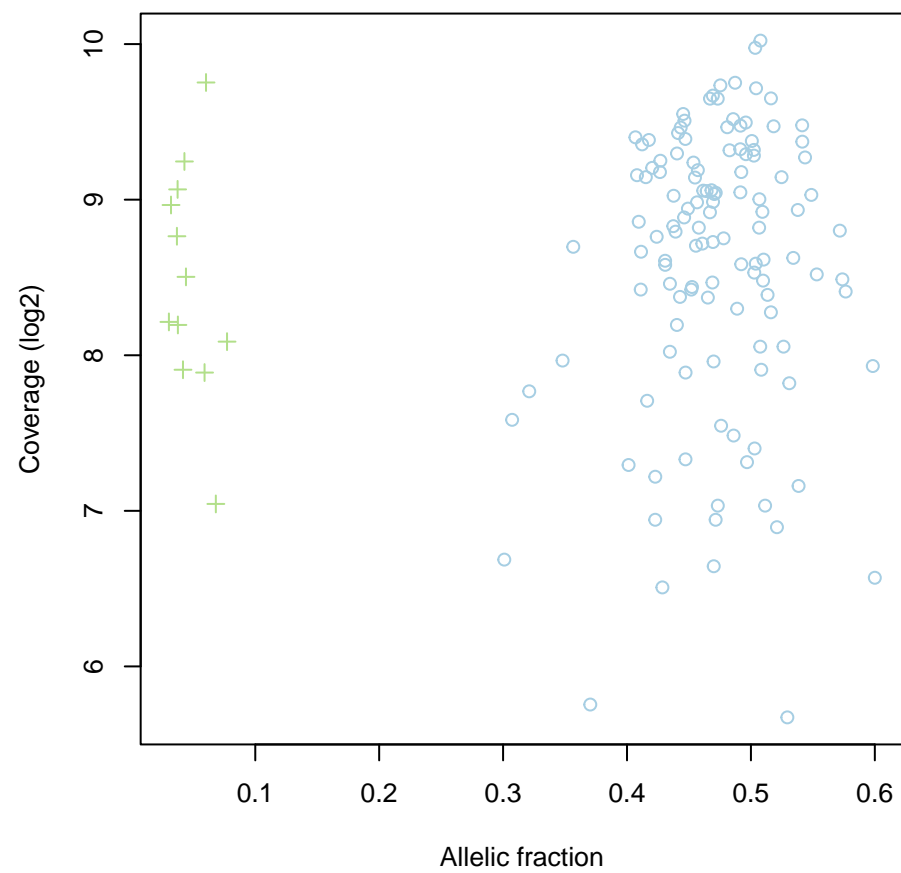
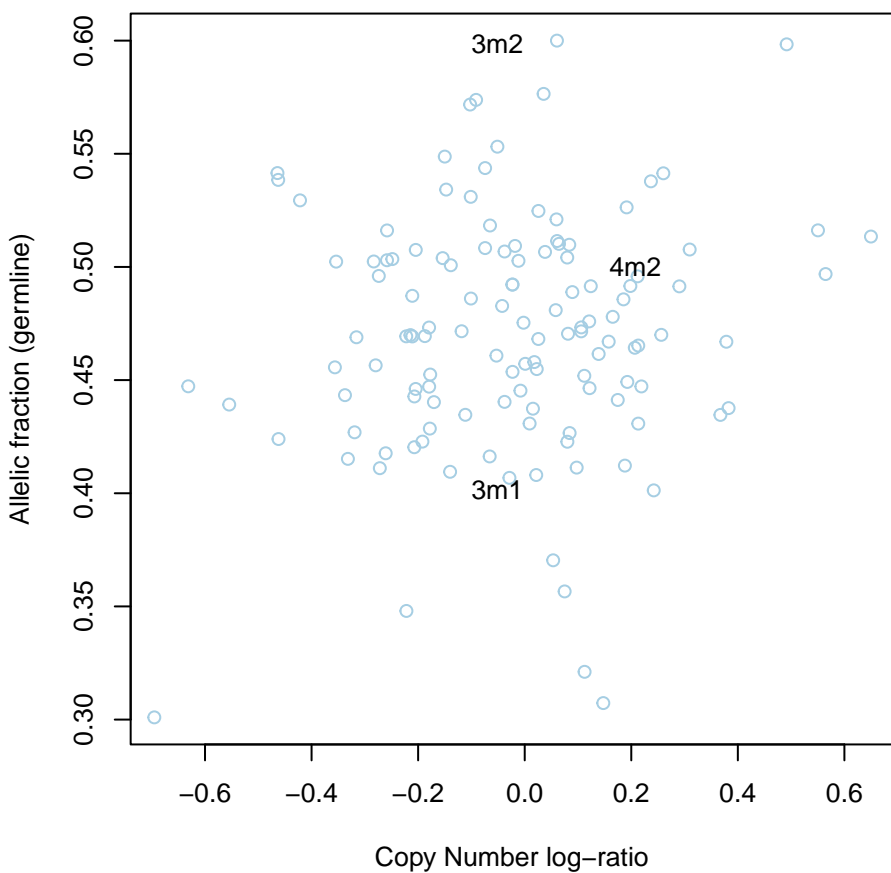


Purity: 0.49 Tumor ploidy: 3.185 SNV log-likelihood: -220.96 GoF: 85.7% Mean coverage: 577,448

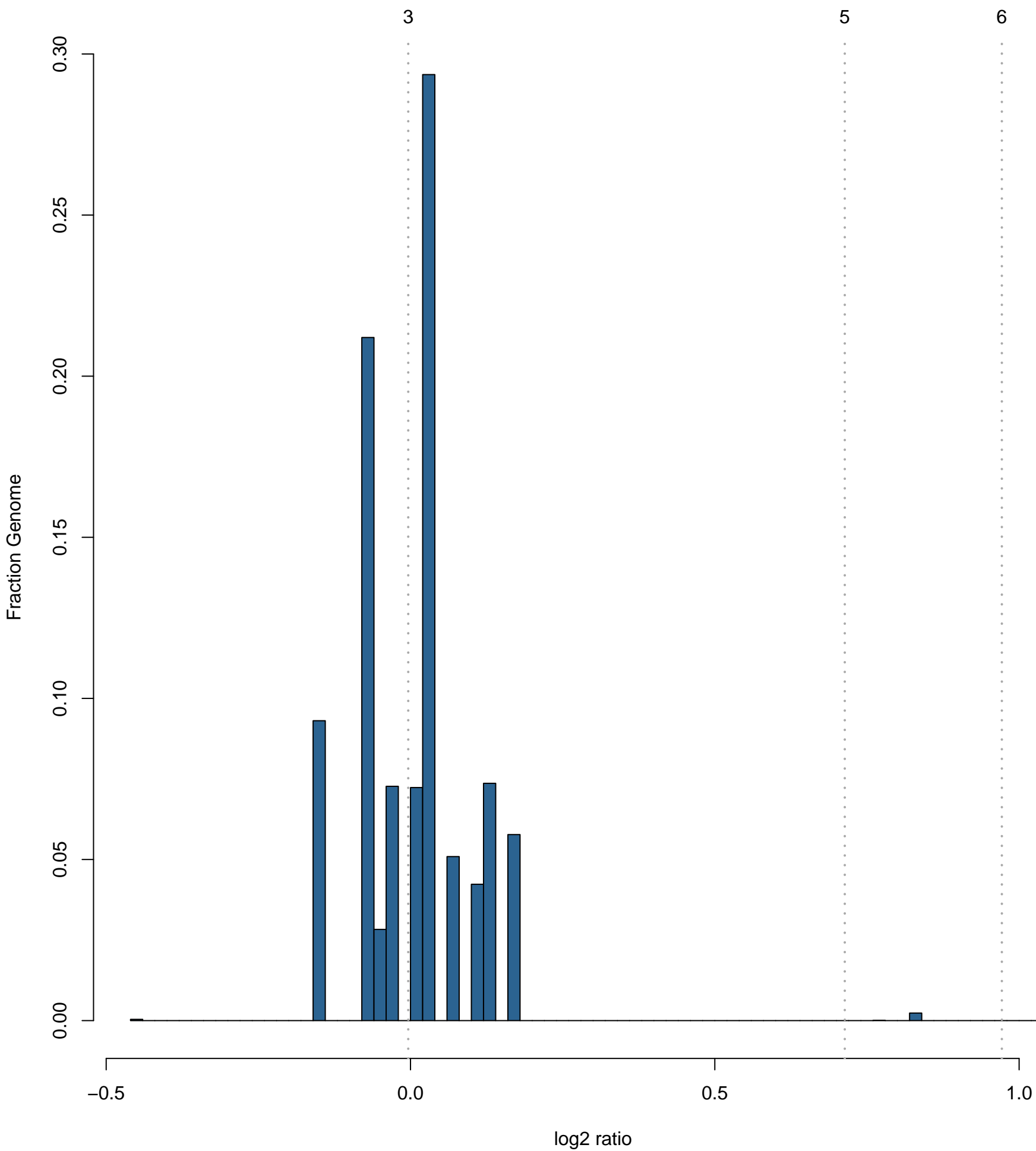


SCNA-fit log-likelihood: -14633.06



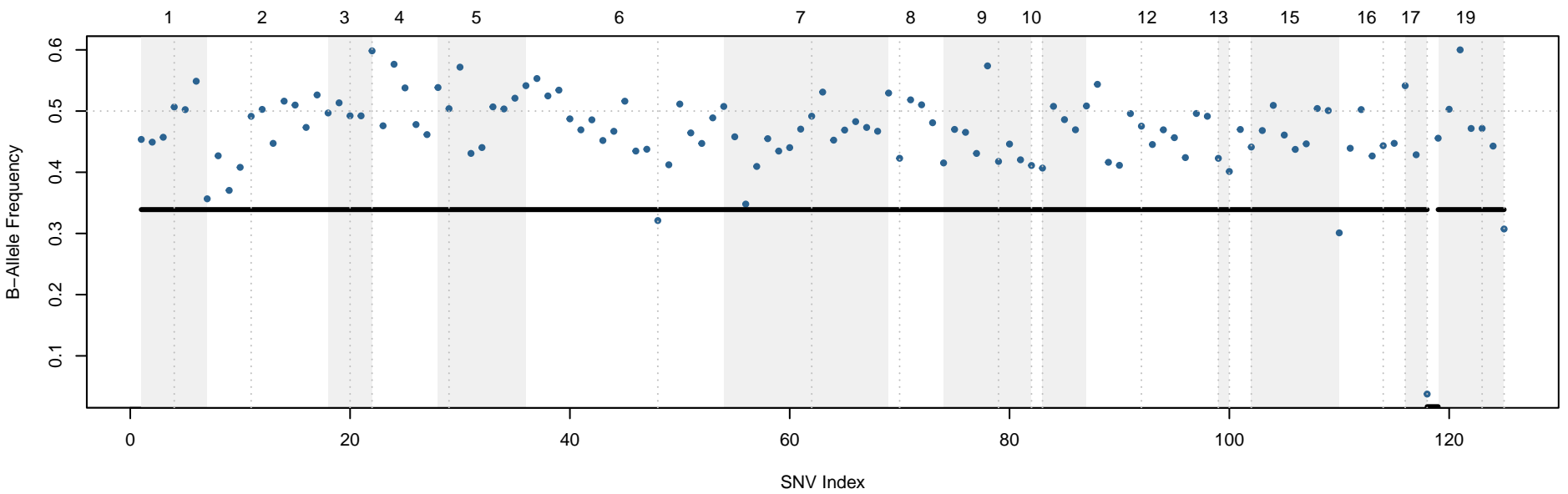


Purity: 0.95 Tumor ploidy: 3.008





Purity: 0.95 Tumor ploidy: 3.008 SNV log-likelihood: -1084.1 GoF: 50.6% Mean coverage: 577,448



SCNA-fit log-likelihood: -14736.18

