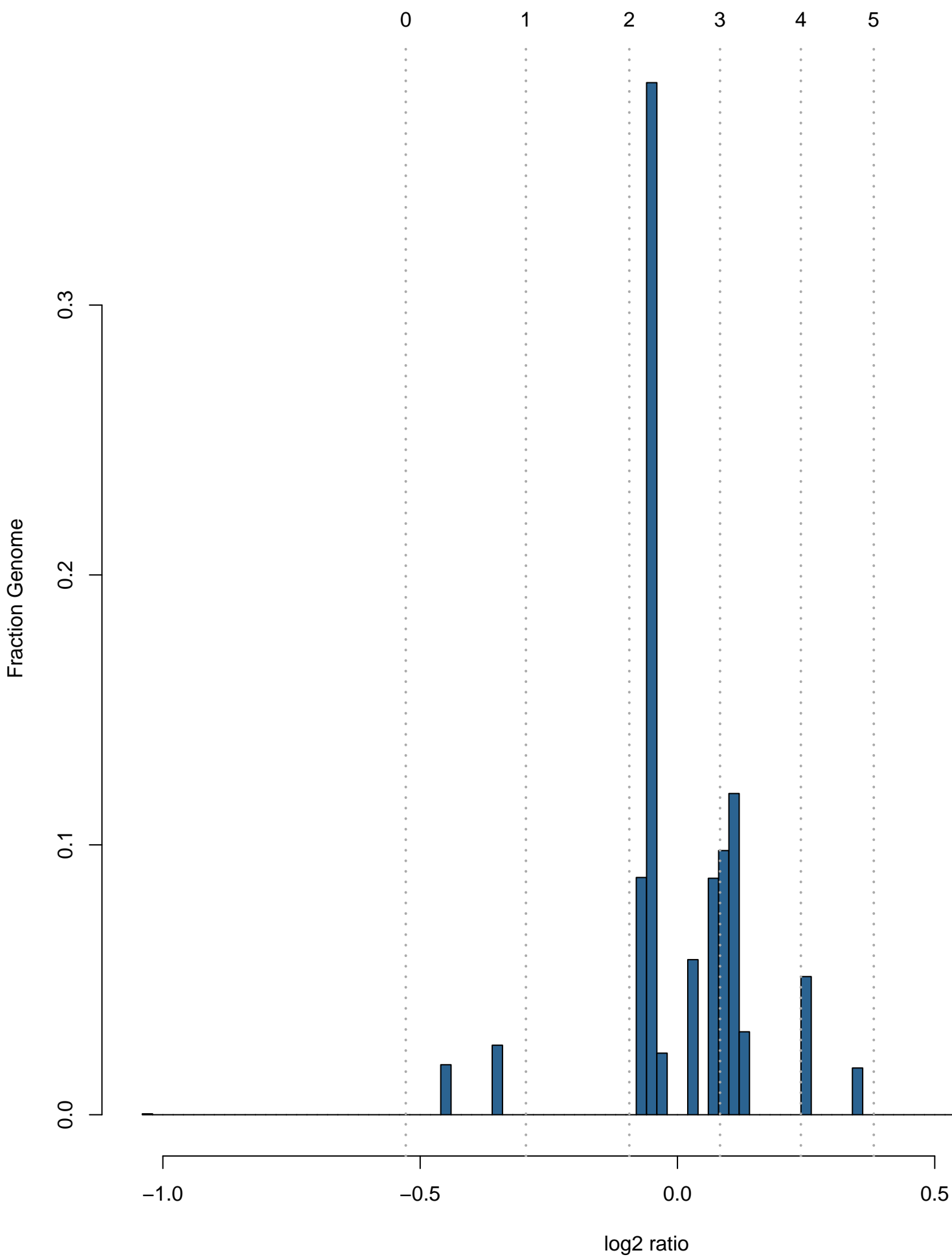
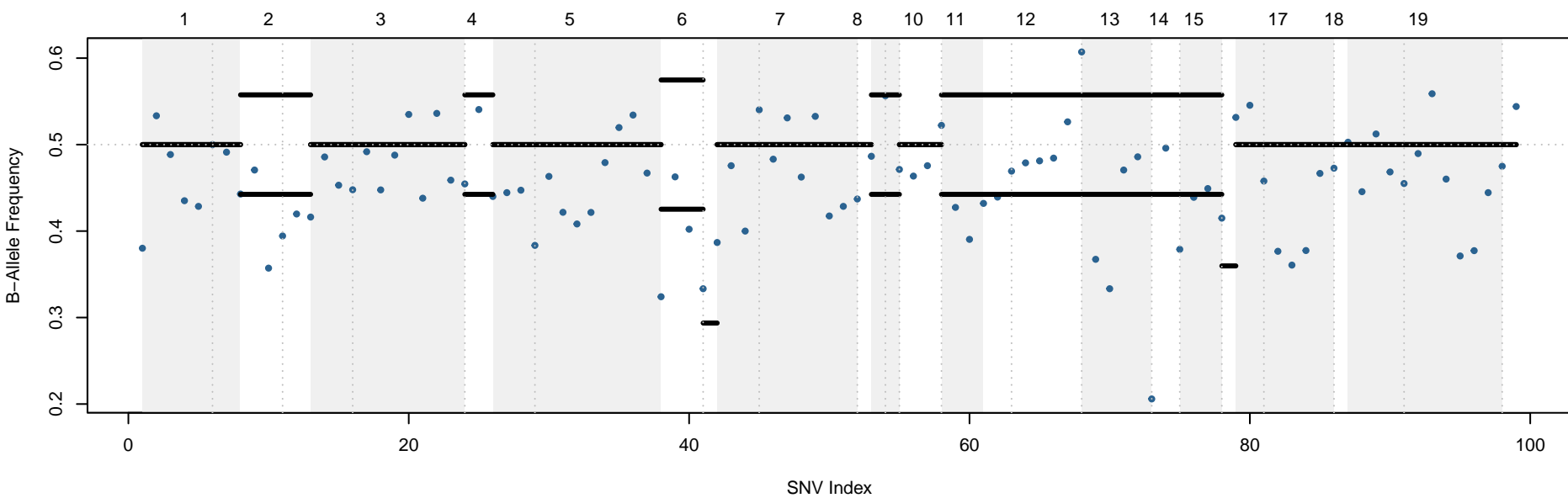


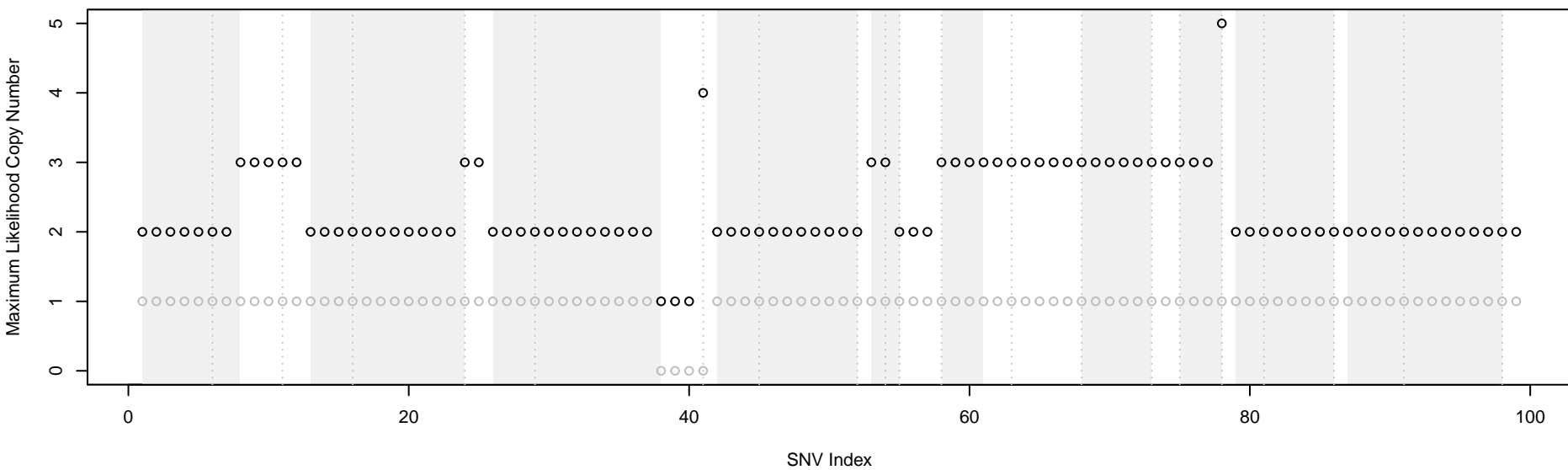
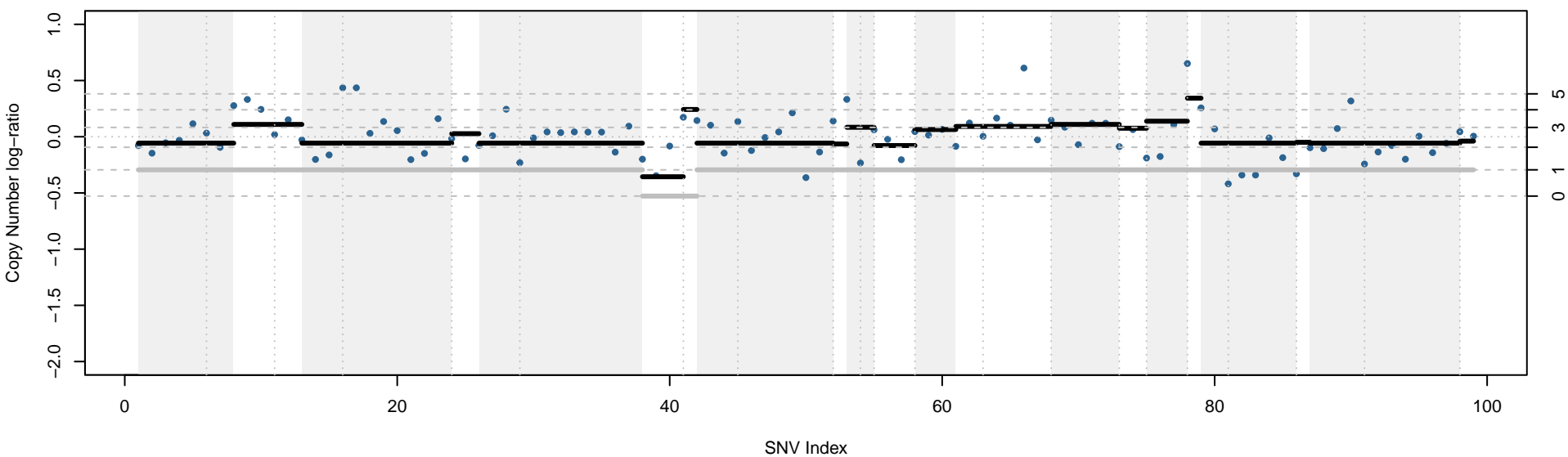
Purity: 0.26 Tumor ploidy: 2.515

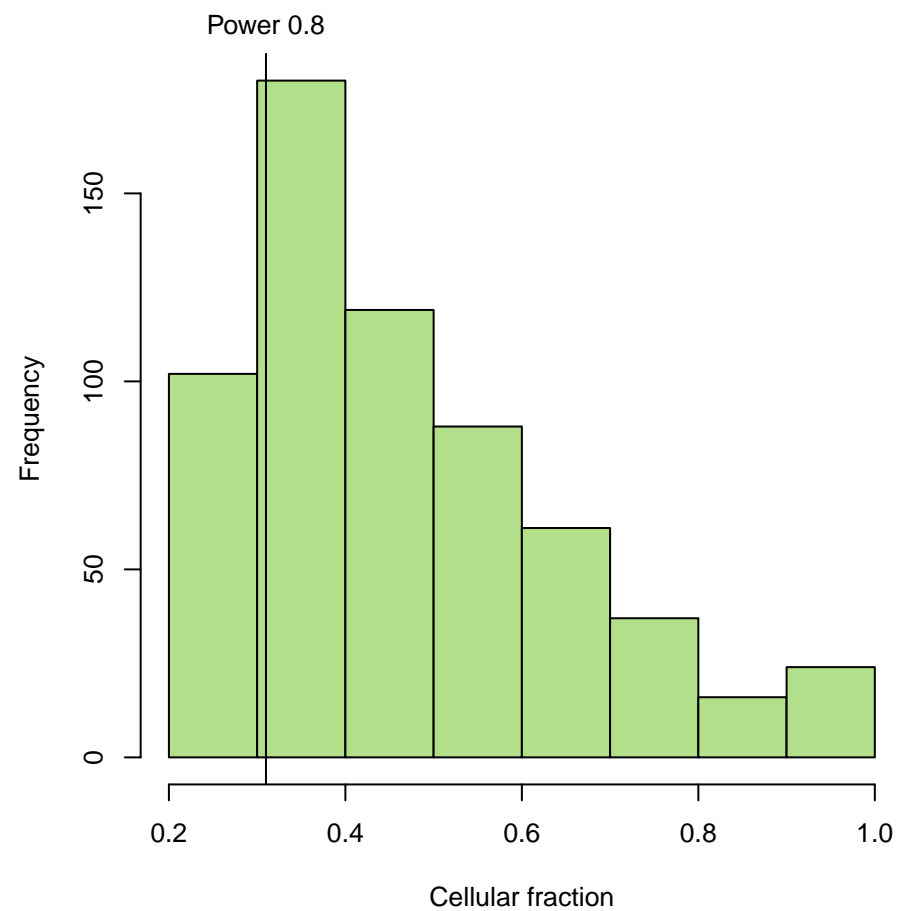
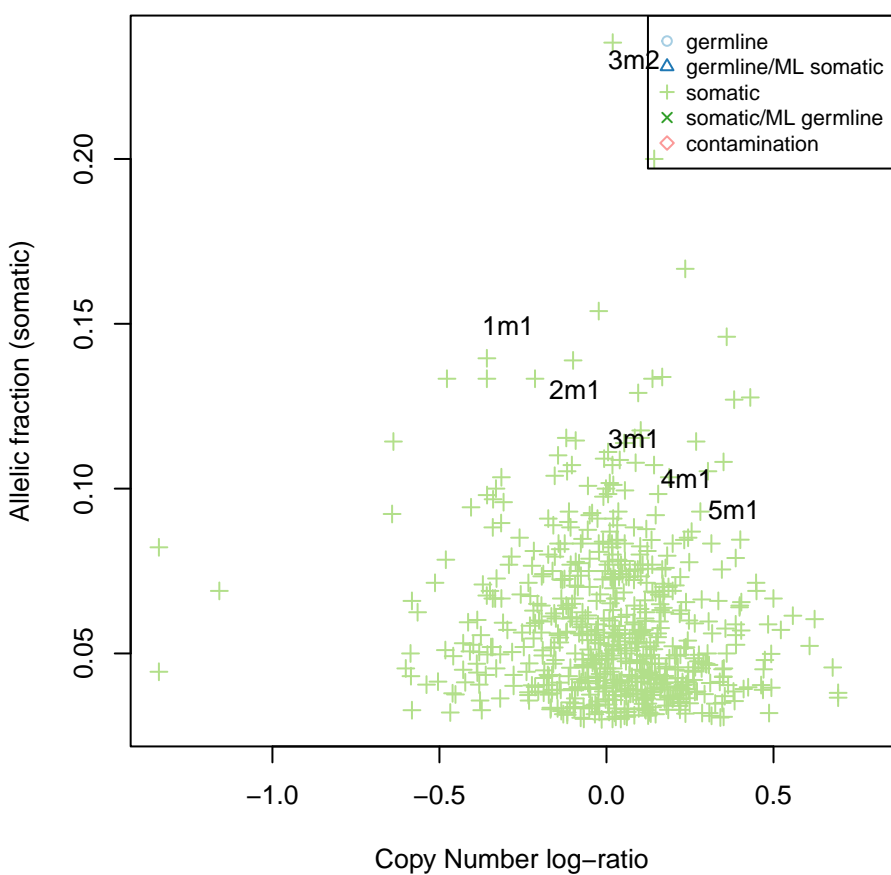
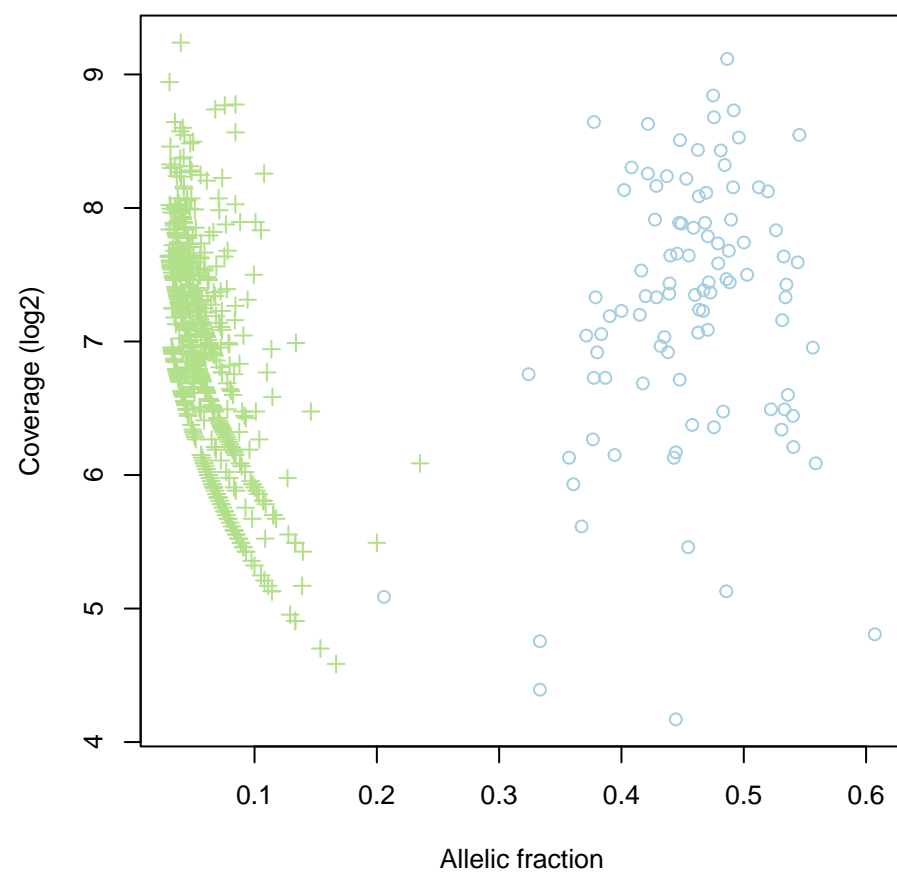
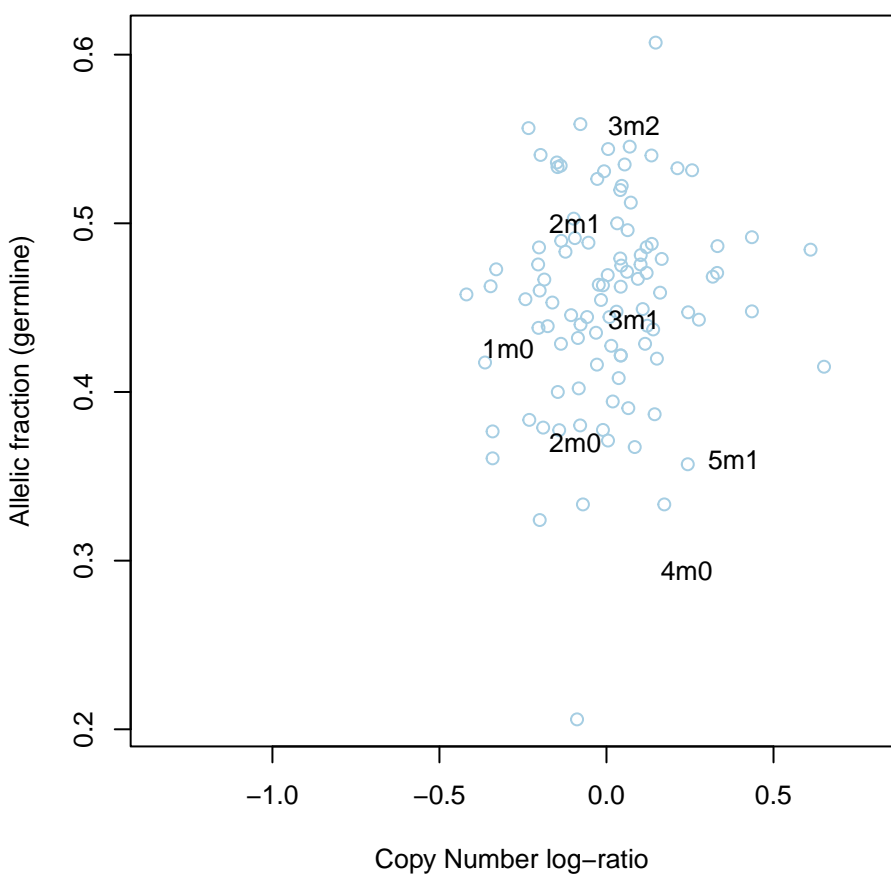


Purity: 0.26 Tumor ploidy: 2.515 SNV log-likelihood: 288.85 GoF: 88.8% Mean coverage: 363;148

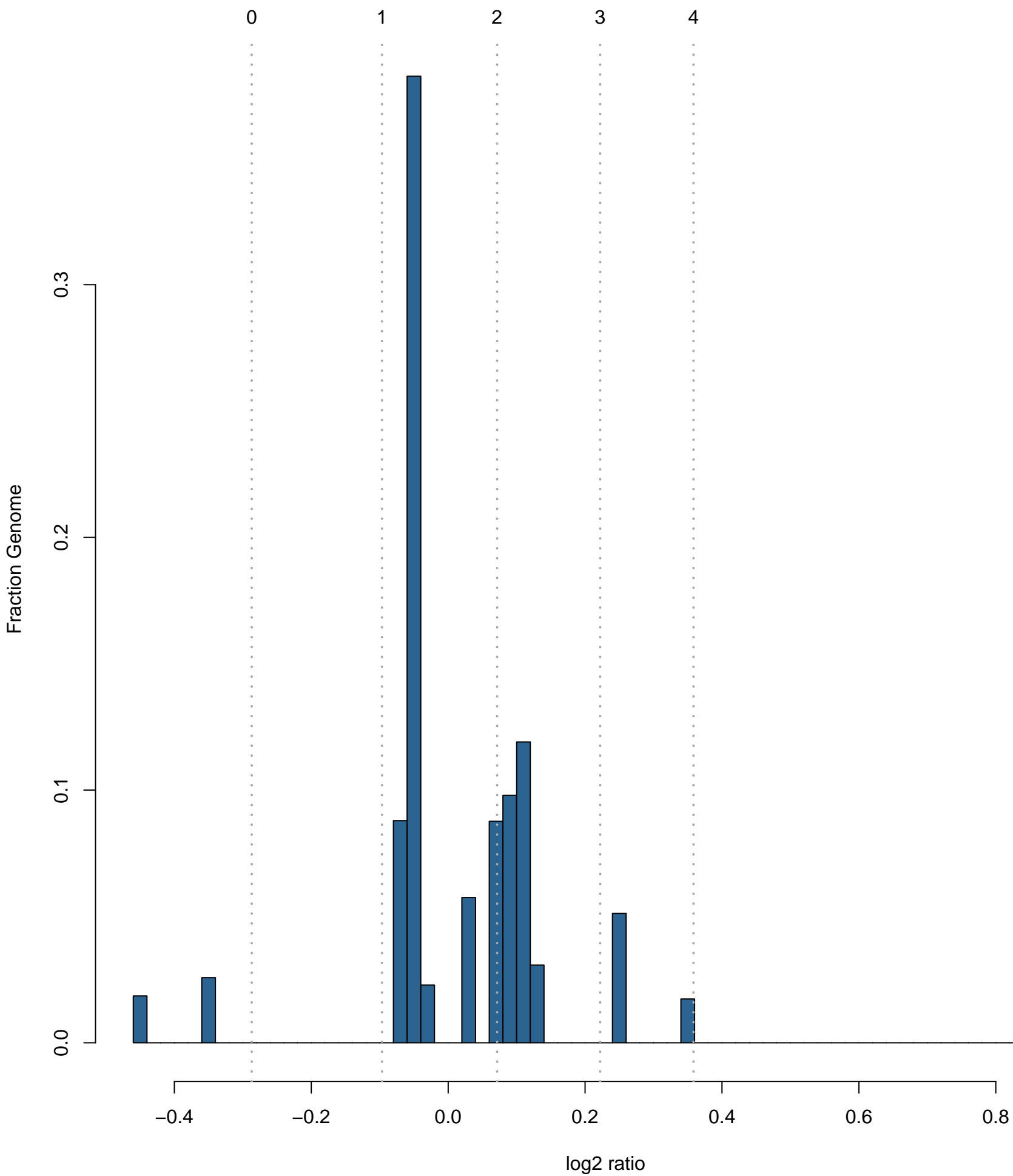


SCNA-fit log-likelihood: -19055.26

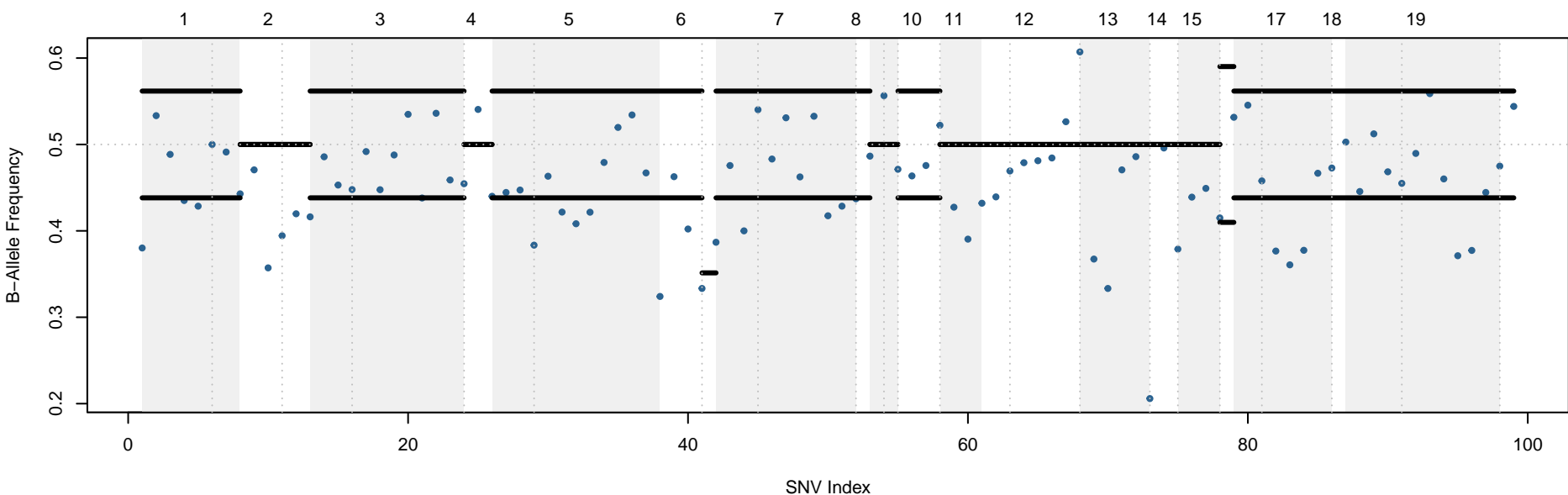




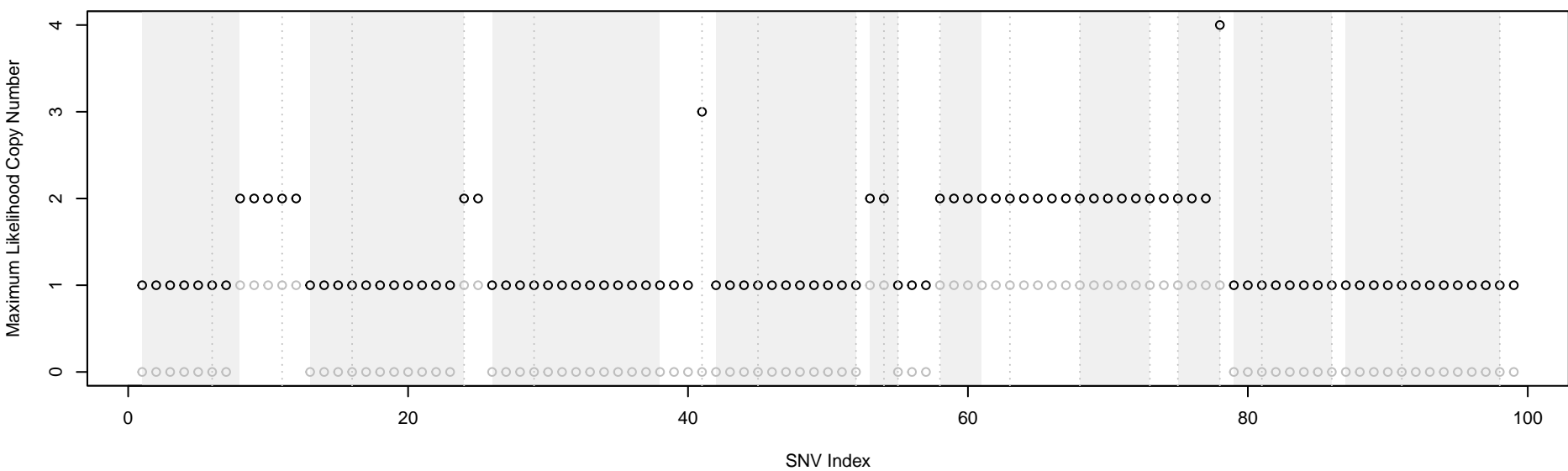
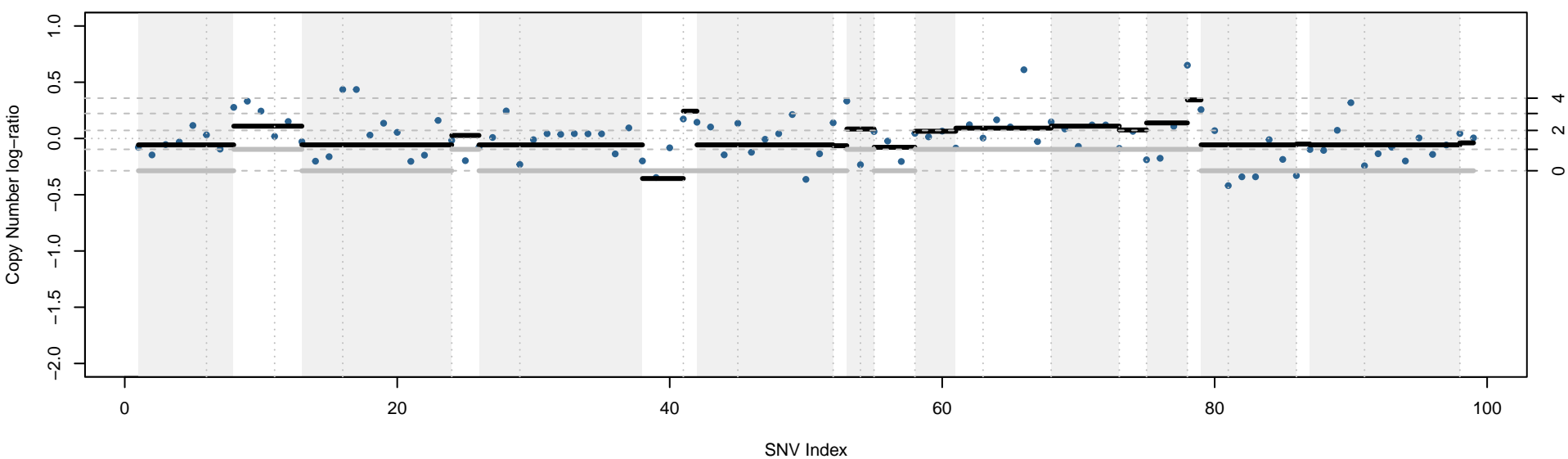
Purity: 0.22 Tumor ploidy: 1.56

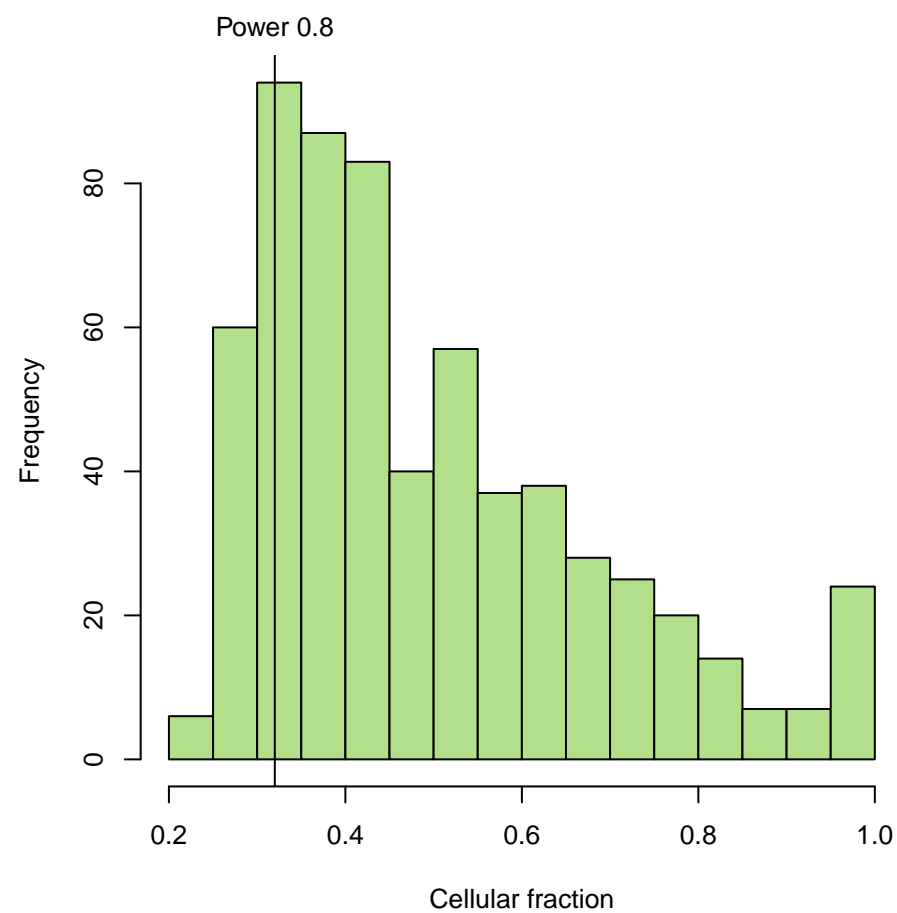
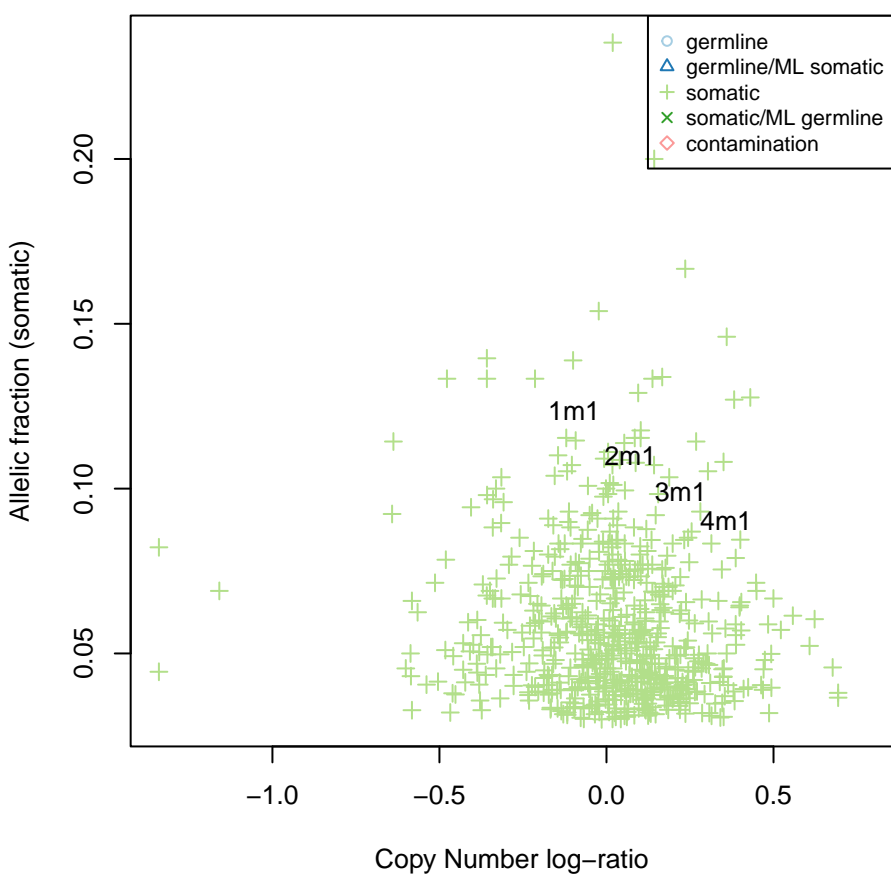
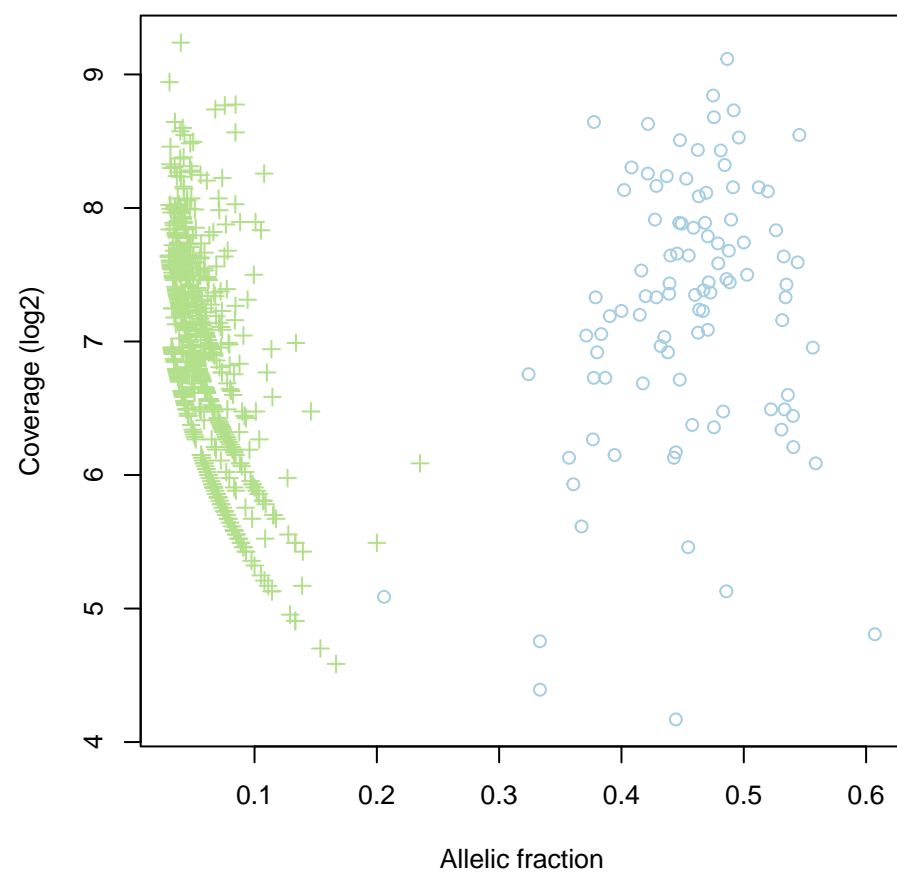
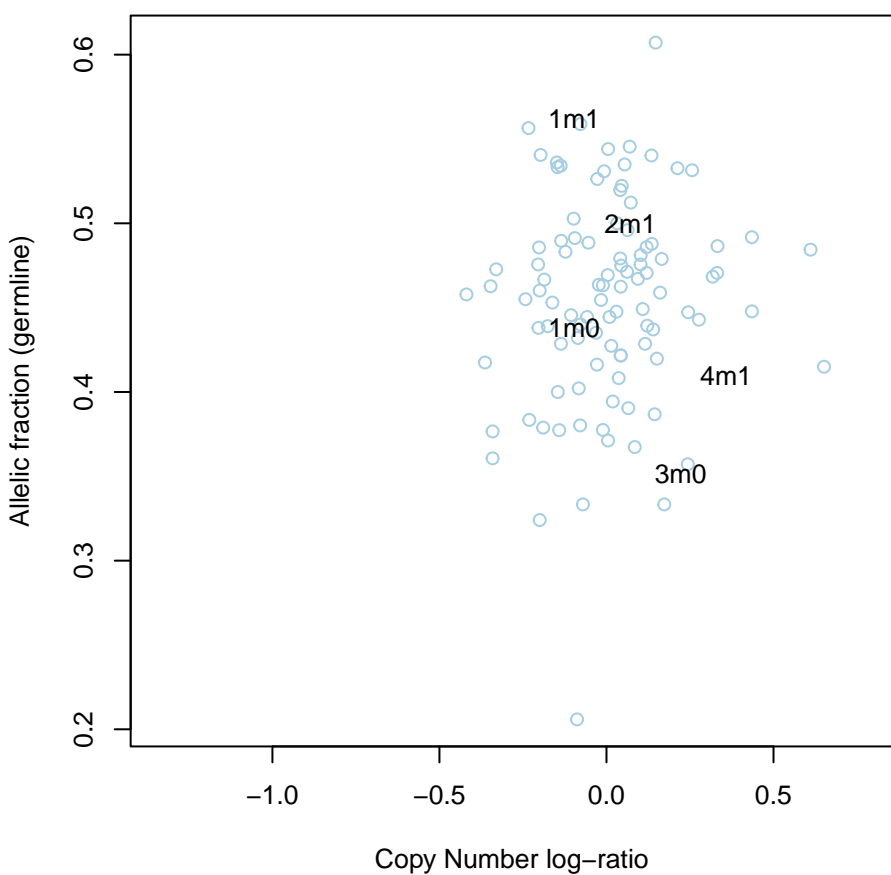


Purity: 0.22 Tumor ploidy: 1.56 SNV log-likelihood: 182.6 GoF: 90.6% Mean coverage: 363;148

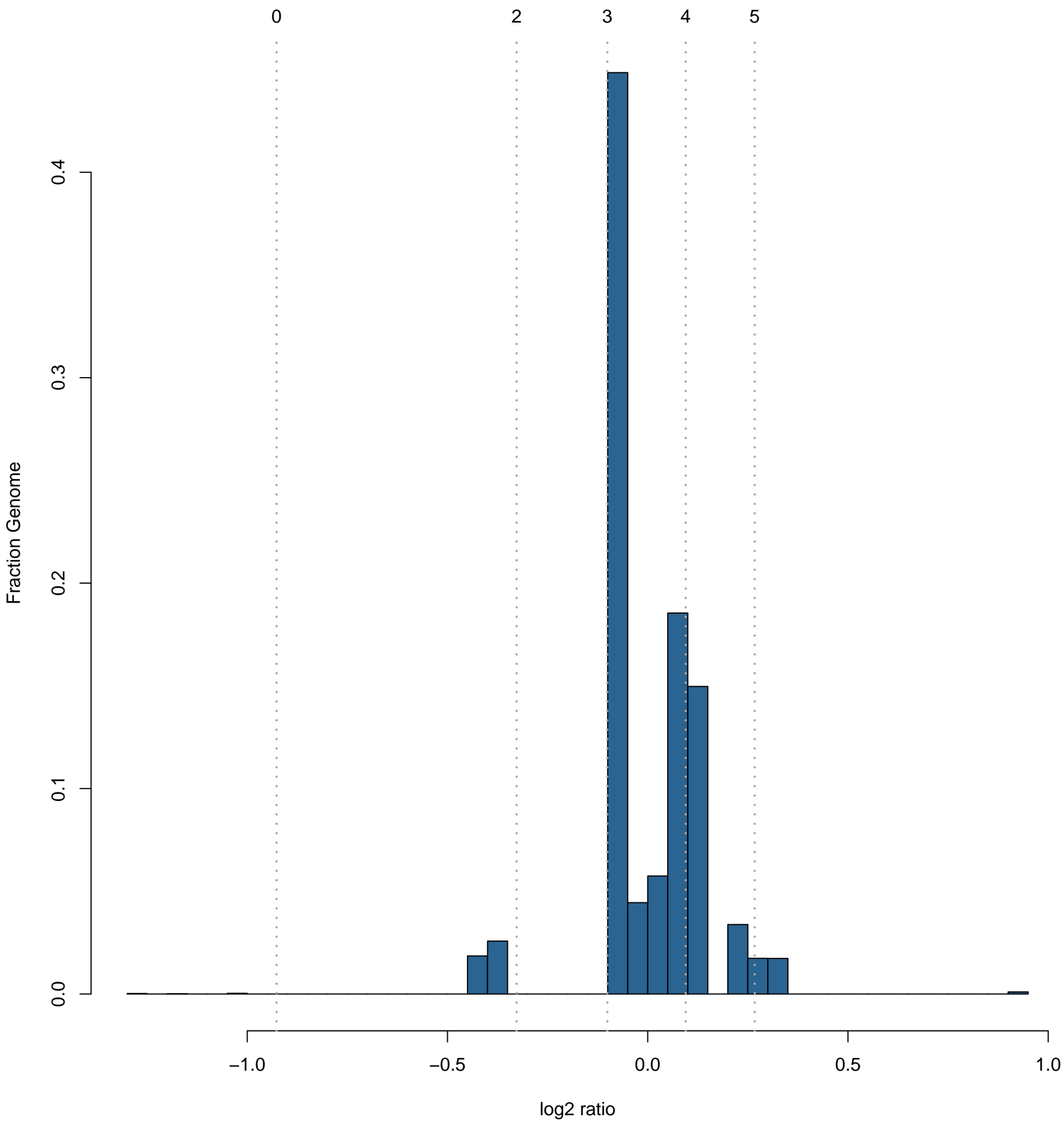


SCNA-fit log-likelihood: -19158.24

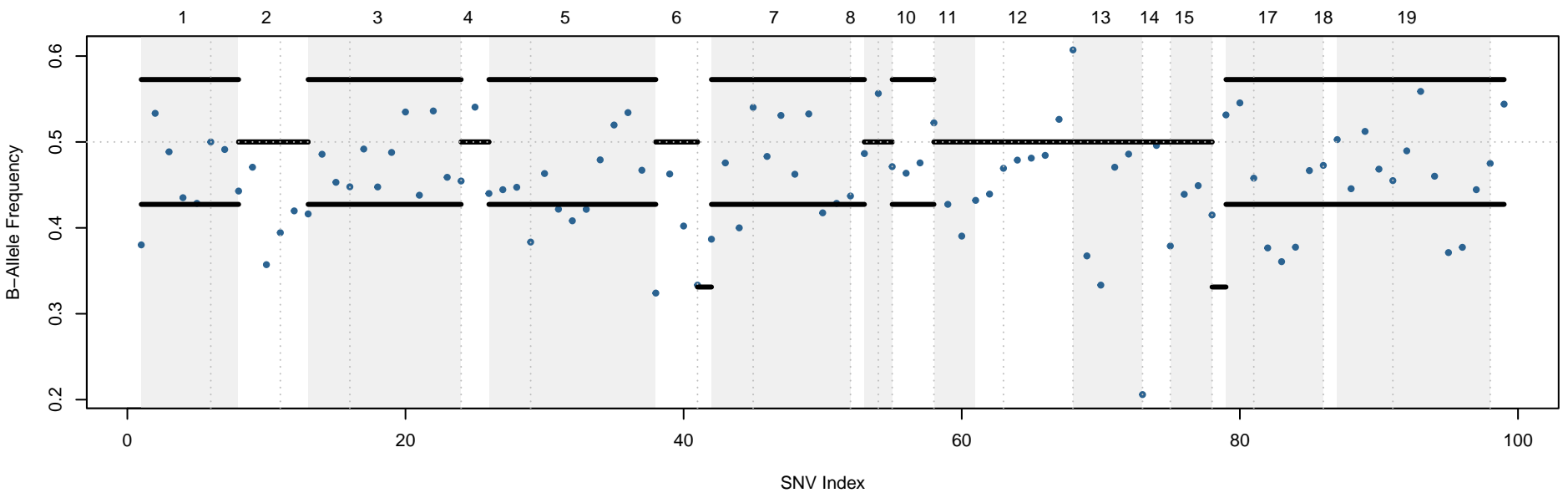




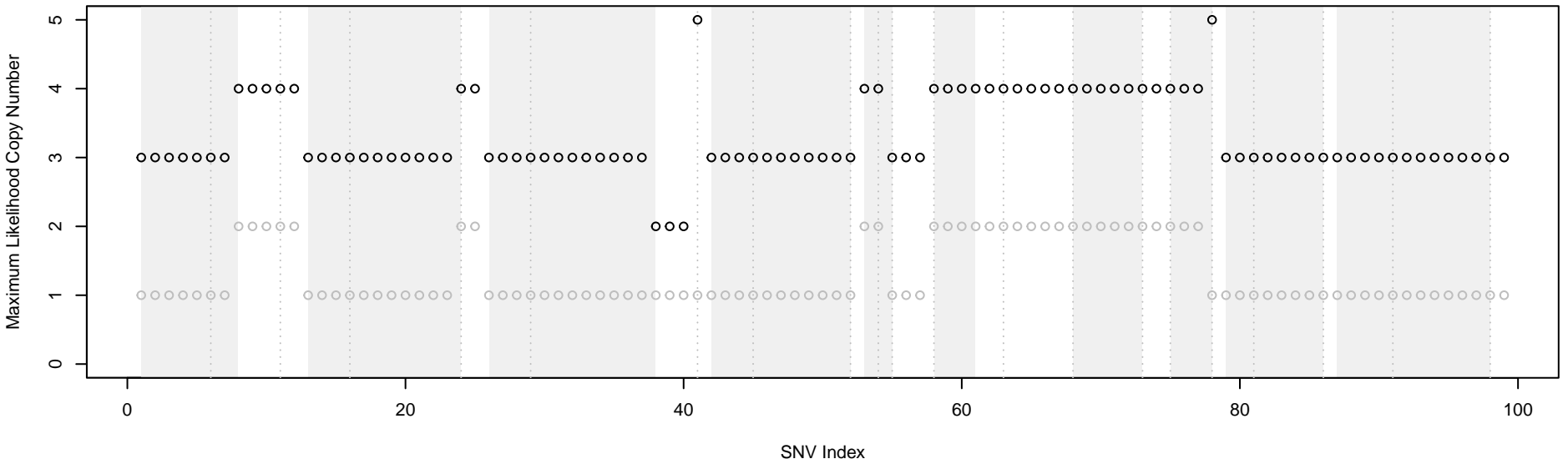
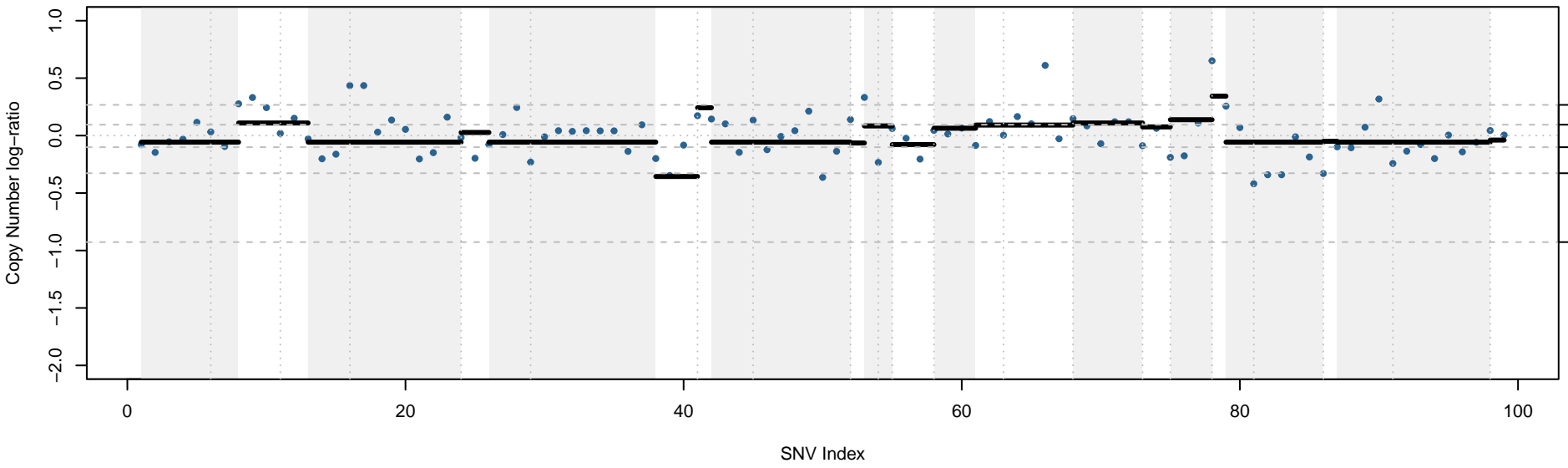
Purity: 0.34 Tumor ploidy: 3.499



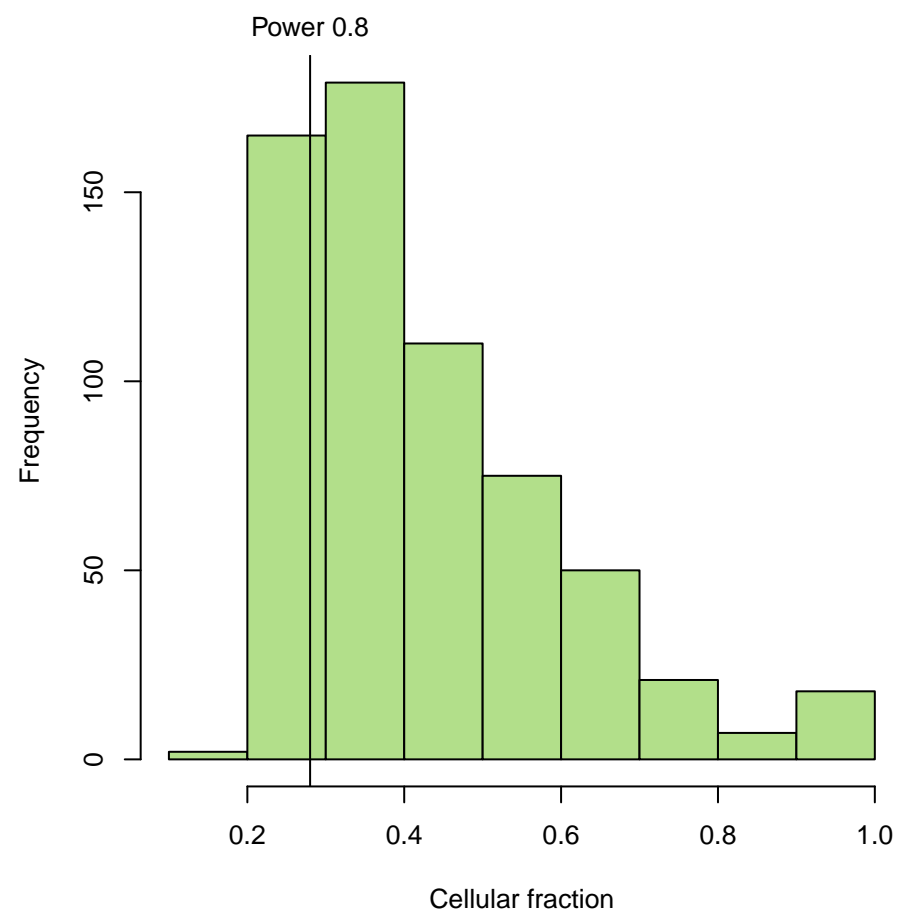
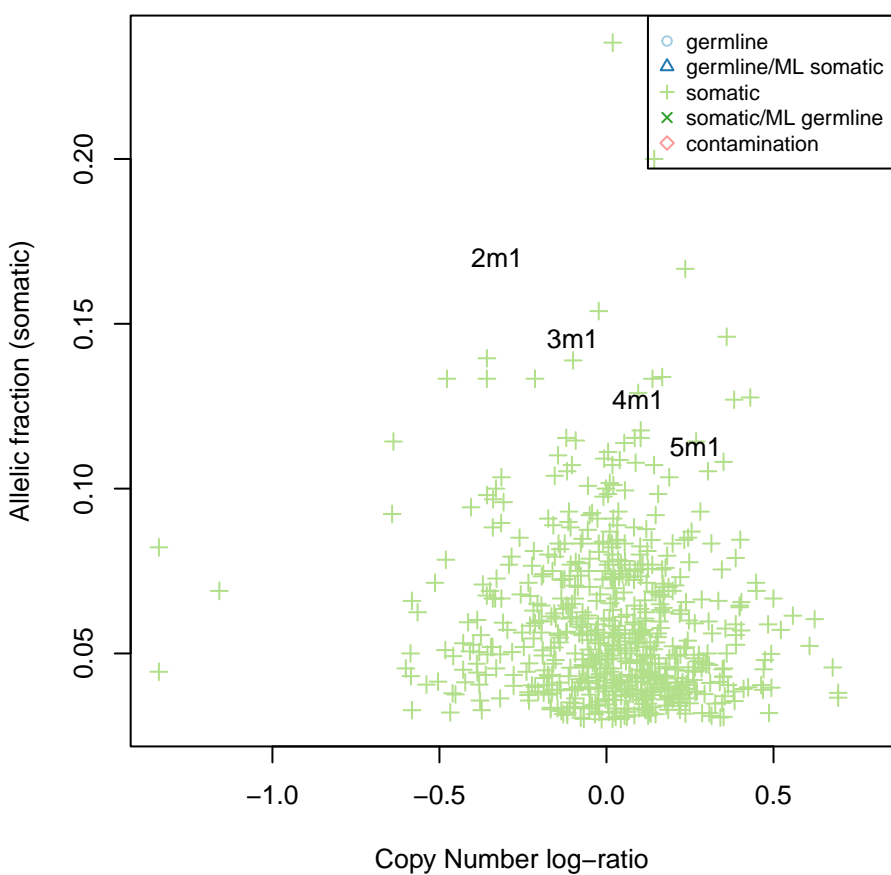
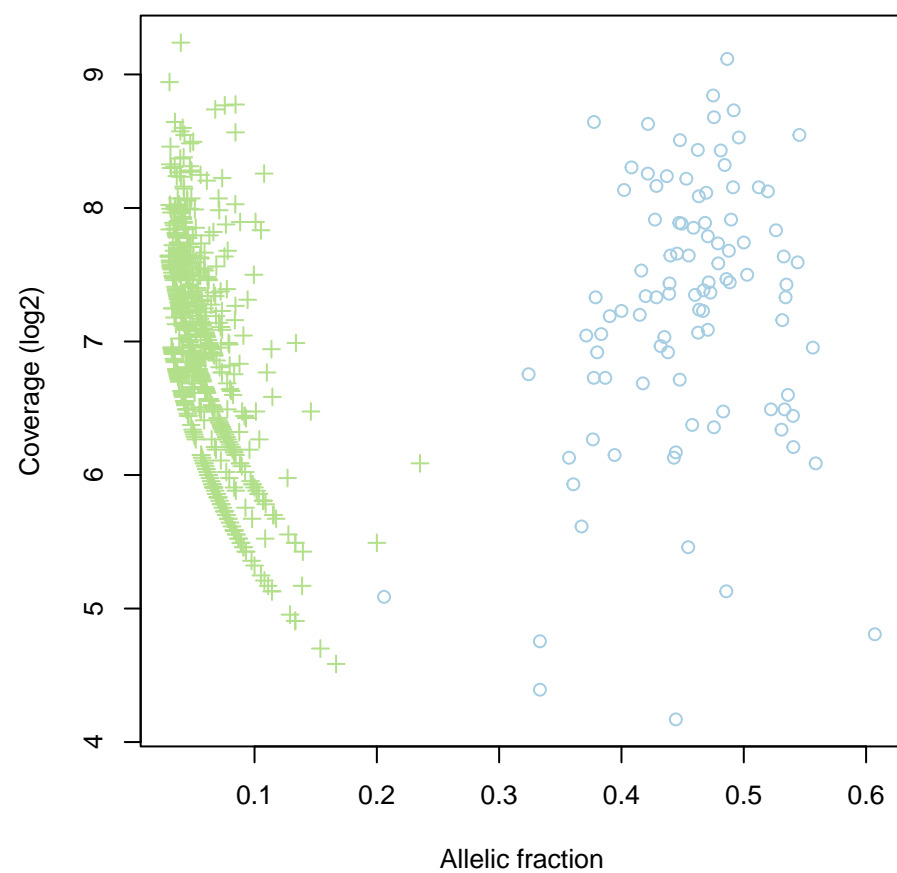
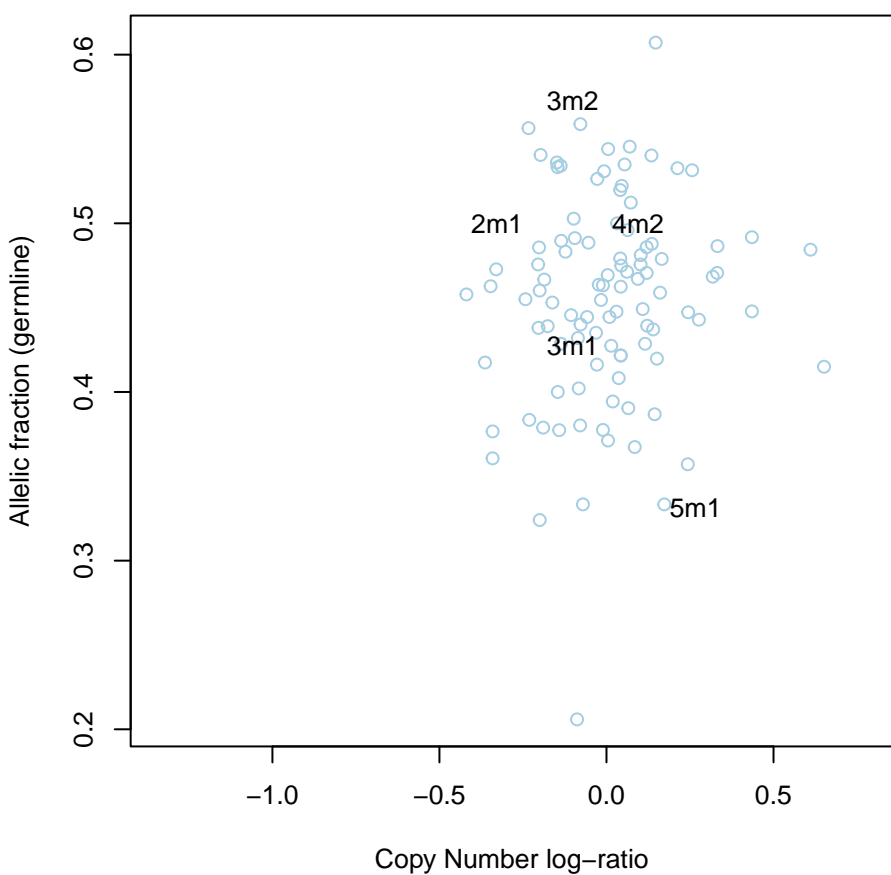
Purity: 0.34 Tumor ploidy: 3.499 SNV log-likelihood: -24.25 GoF: 84.3% Mean coverage: 363;148



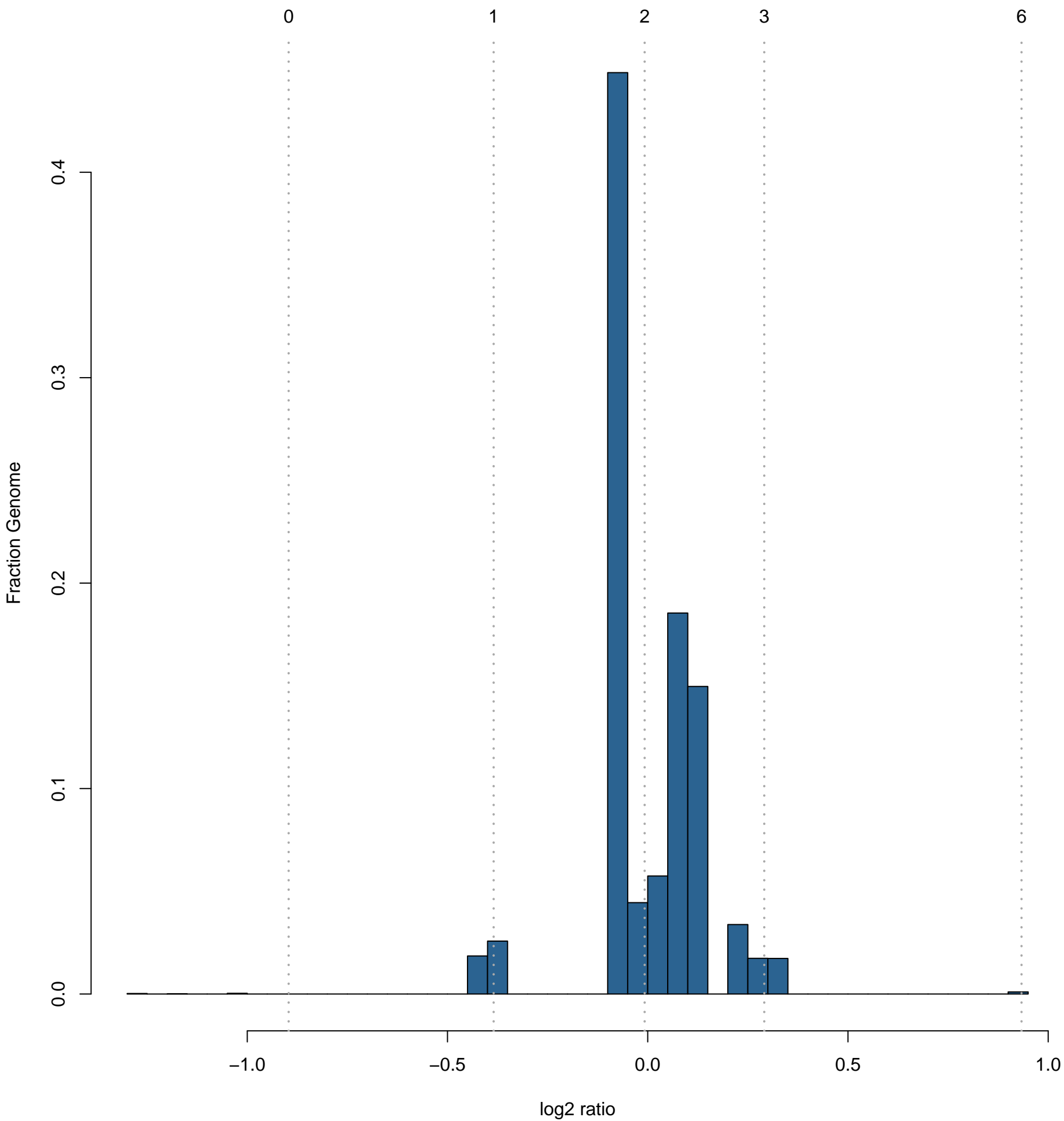
SCNA-fit log-likelihood: -19054.3



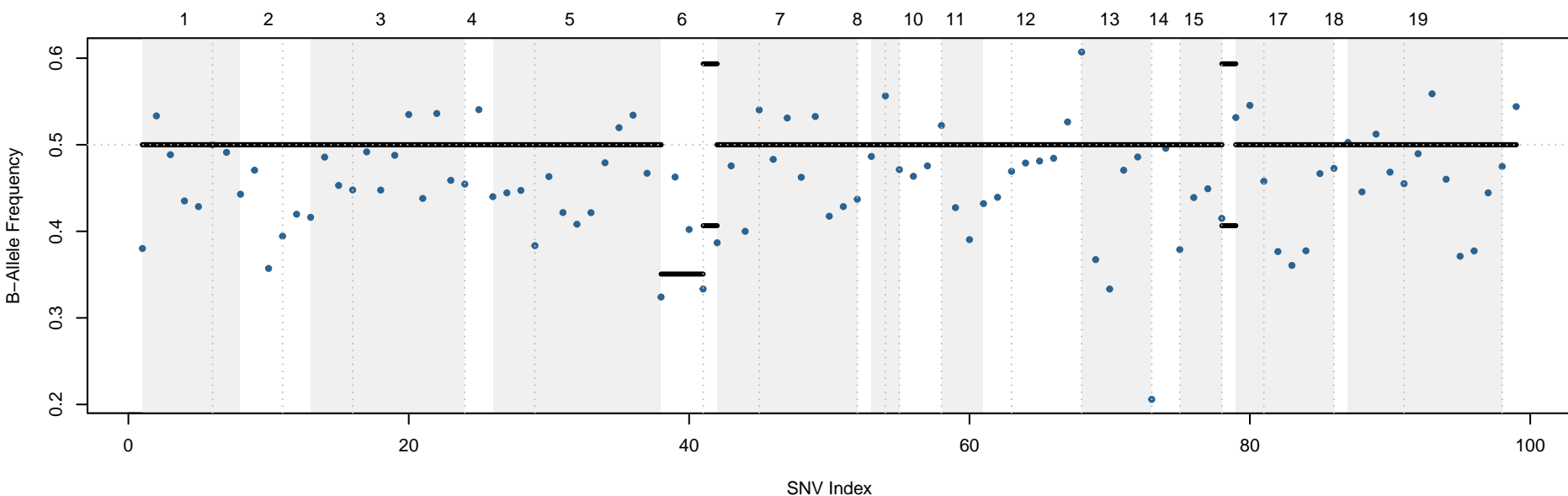




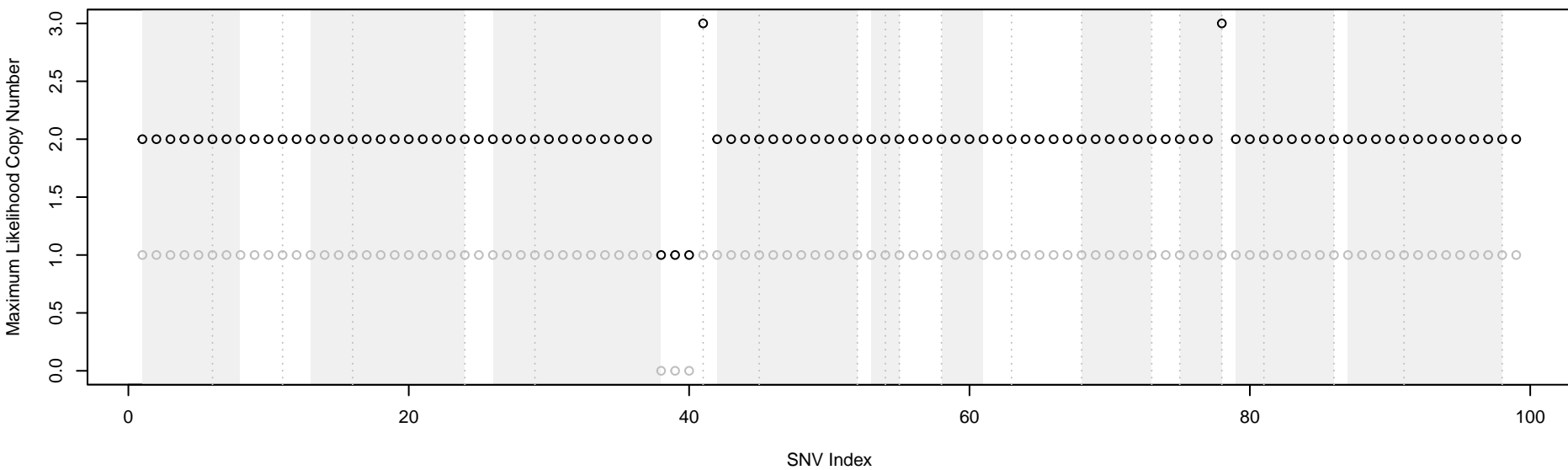
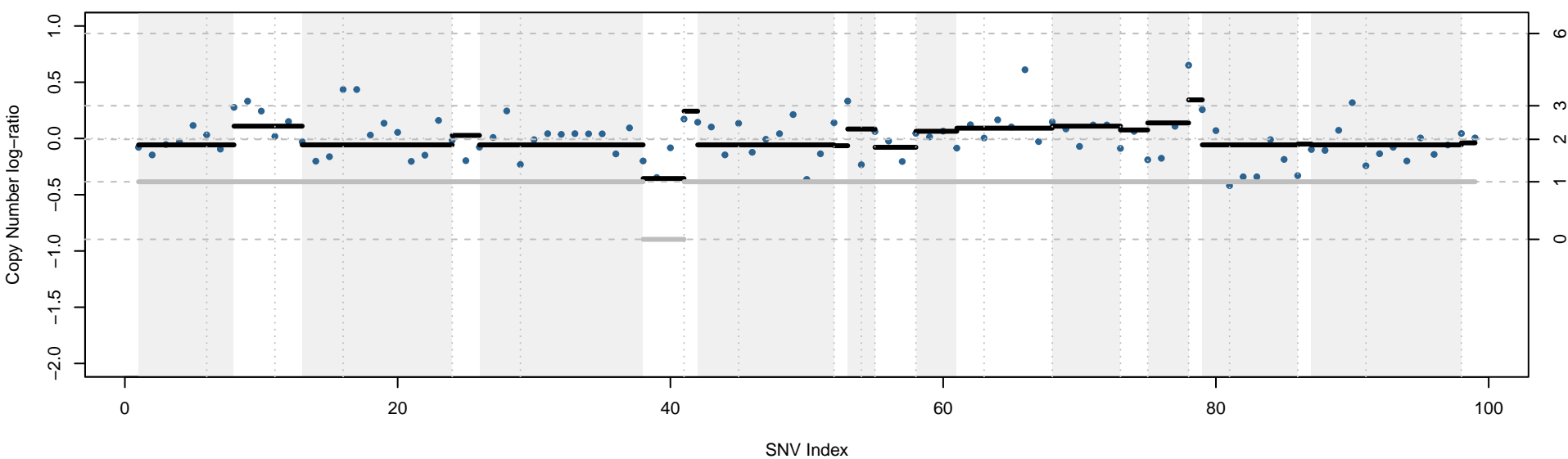
Purity: 0.46 Tumor ploidy: 2.023

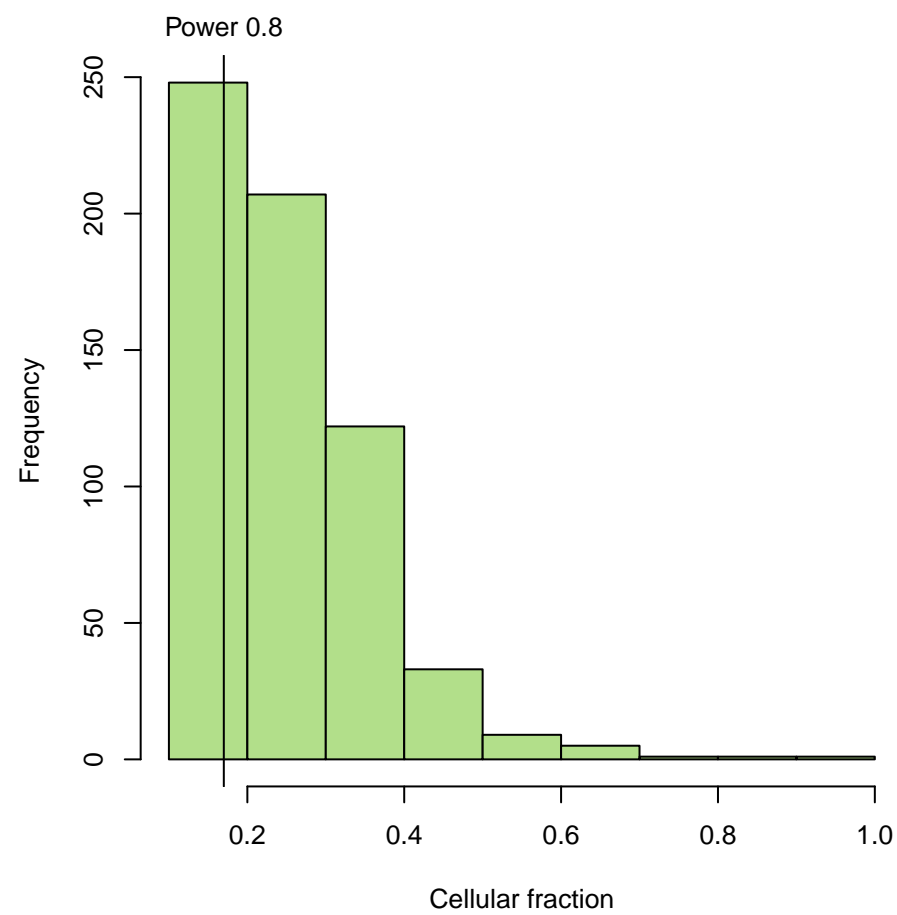
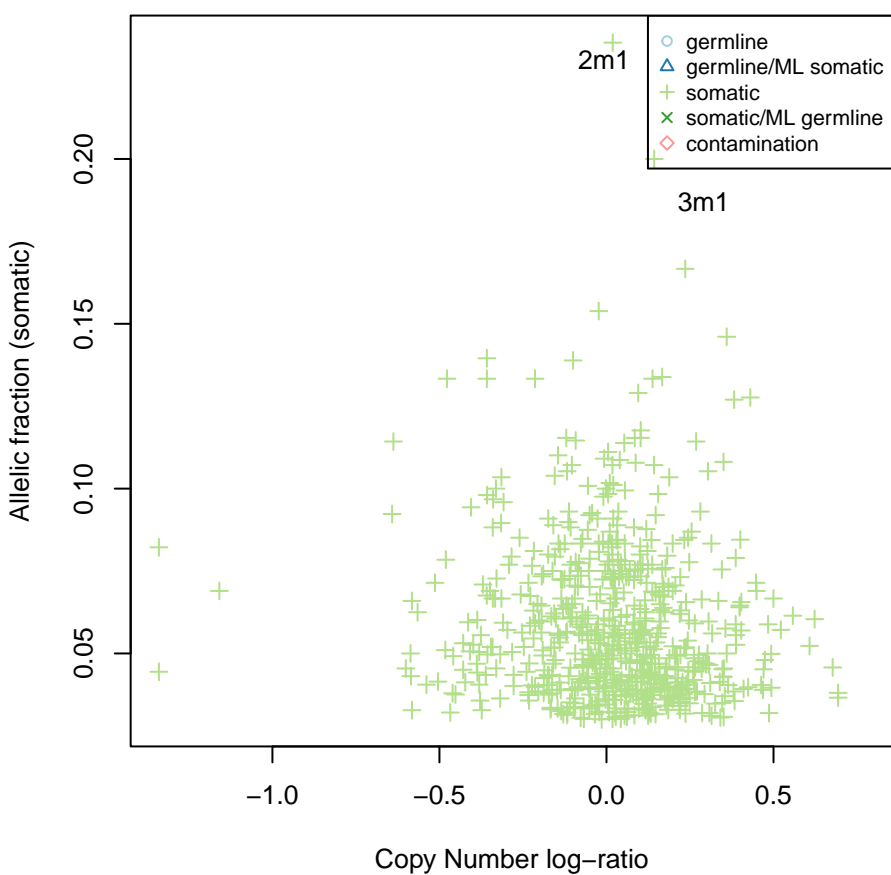
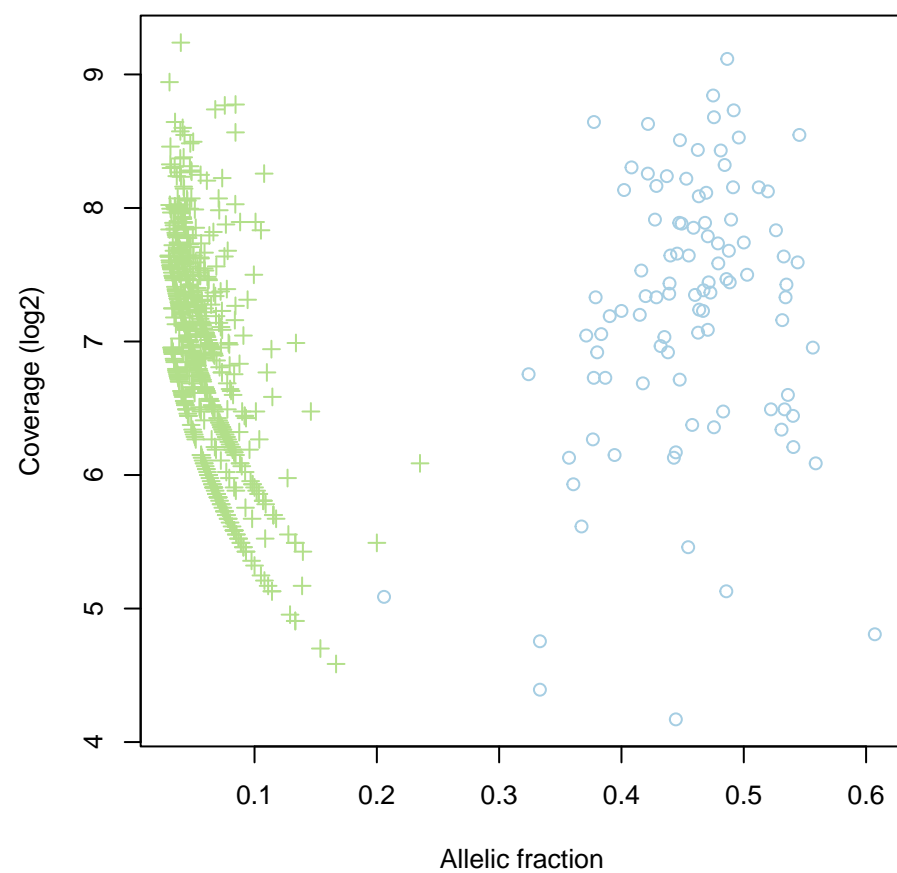
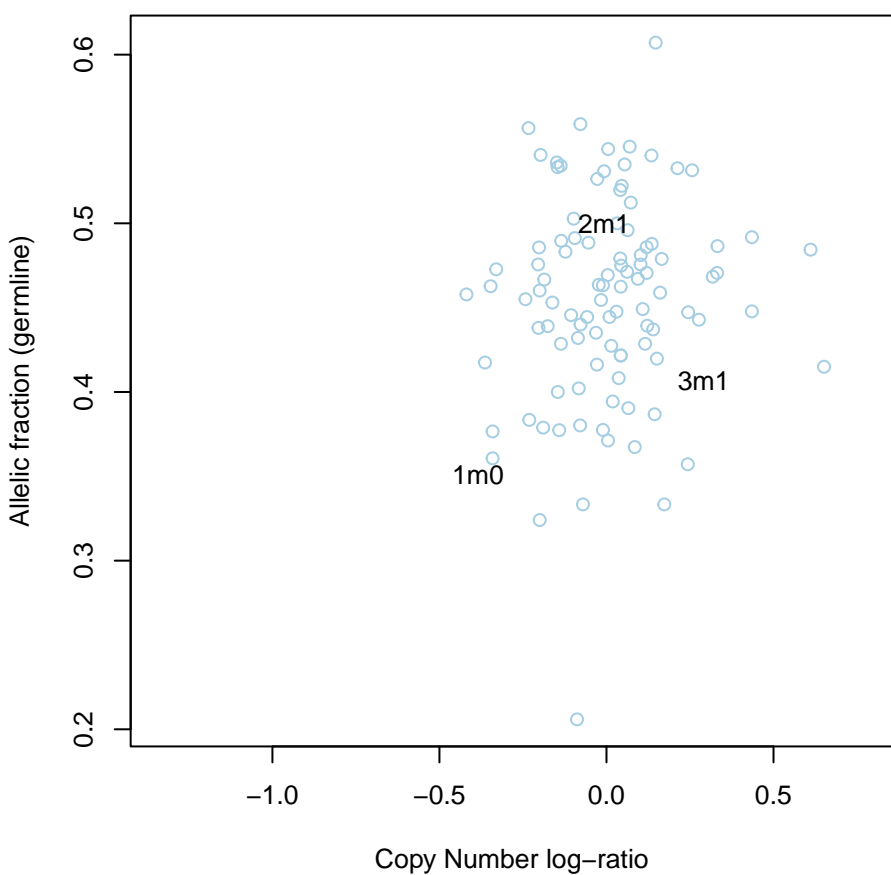


Purity: 0.46 Tumor ploidy: 2.023 SNV log-likelihood: 4.52 GoF: 33.7% Mean coverage: 363;148

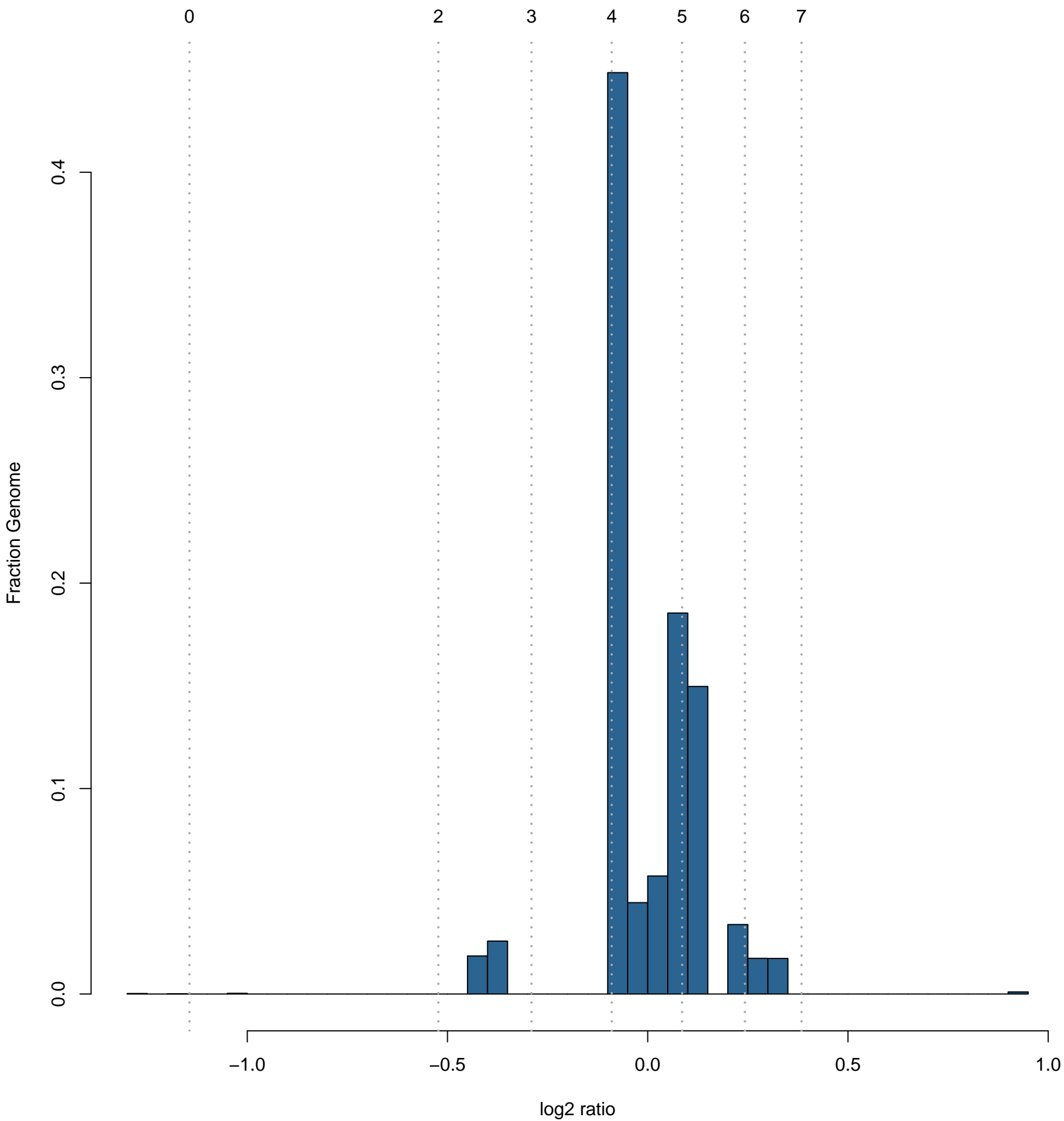


SCNA-fit log-likelihood: -19151.34

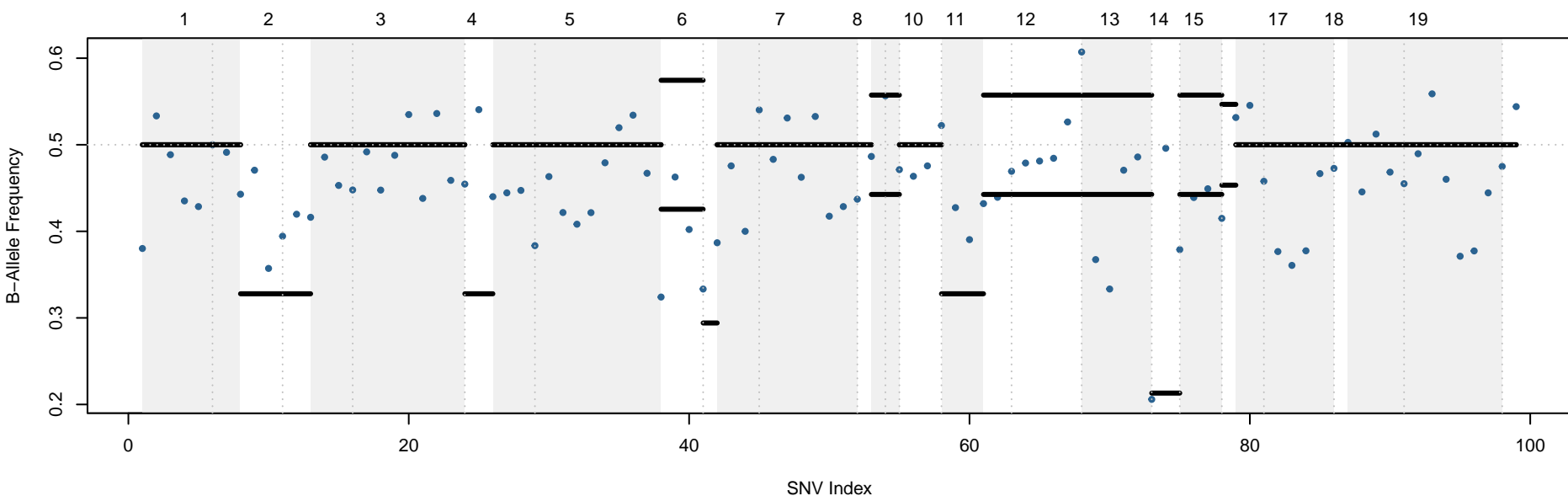




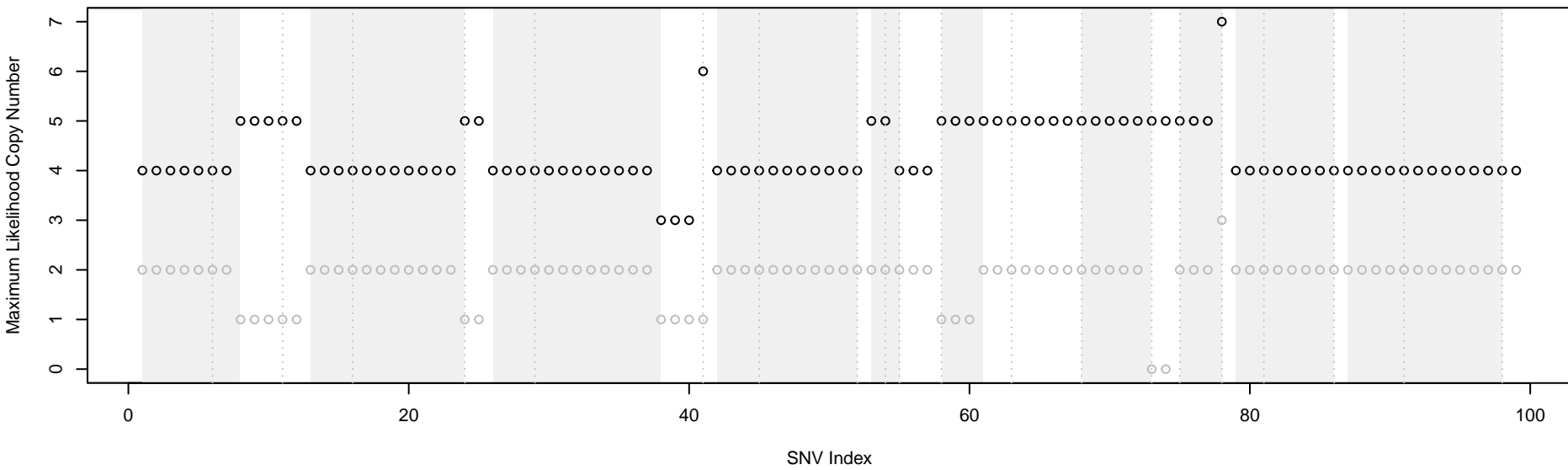
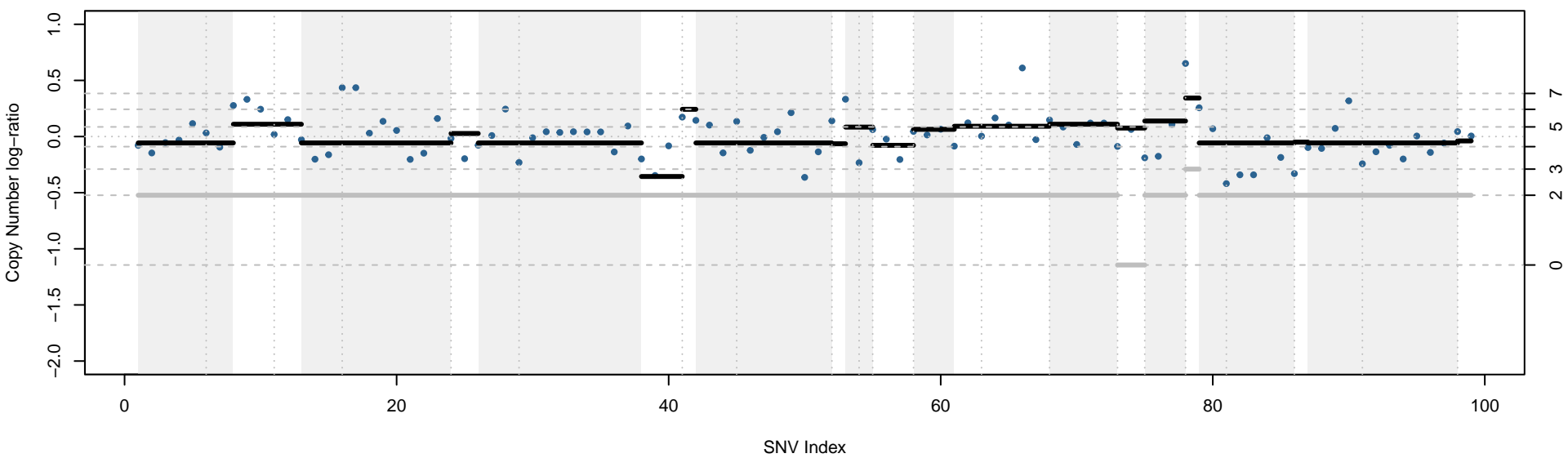
Purity: 0.35 Tumor ploidy: 4.496

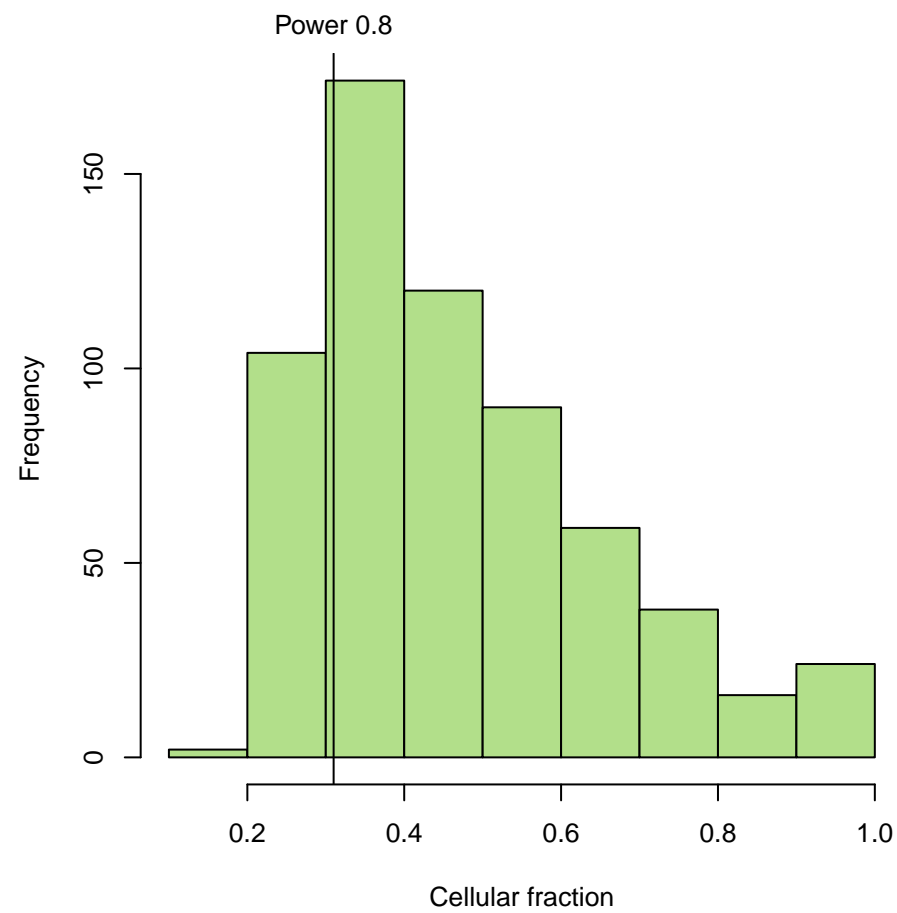
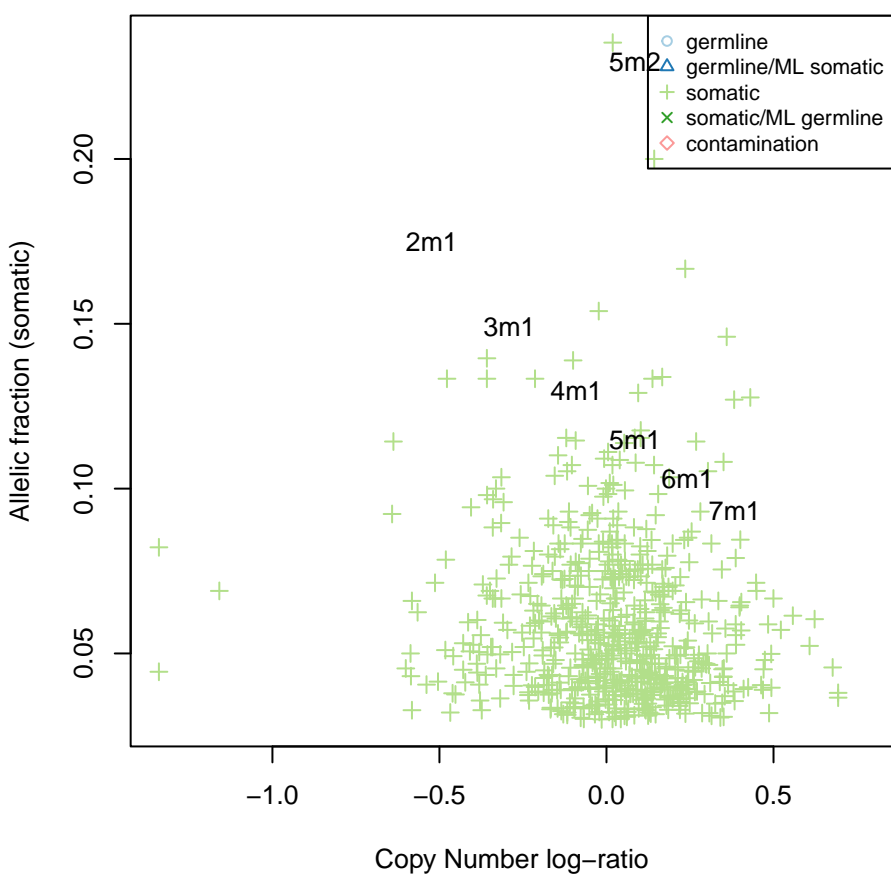
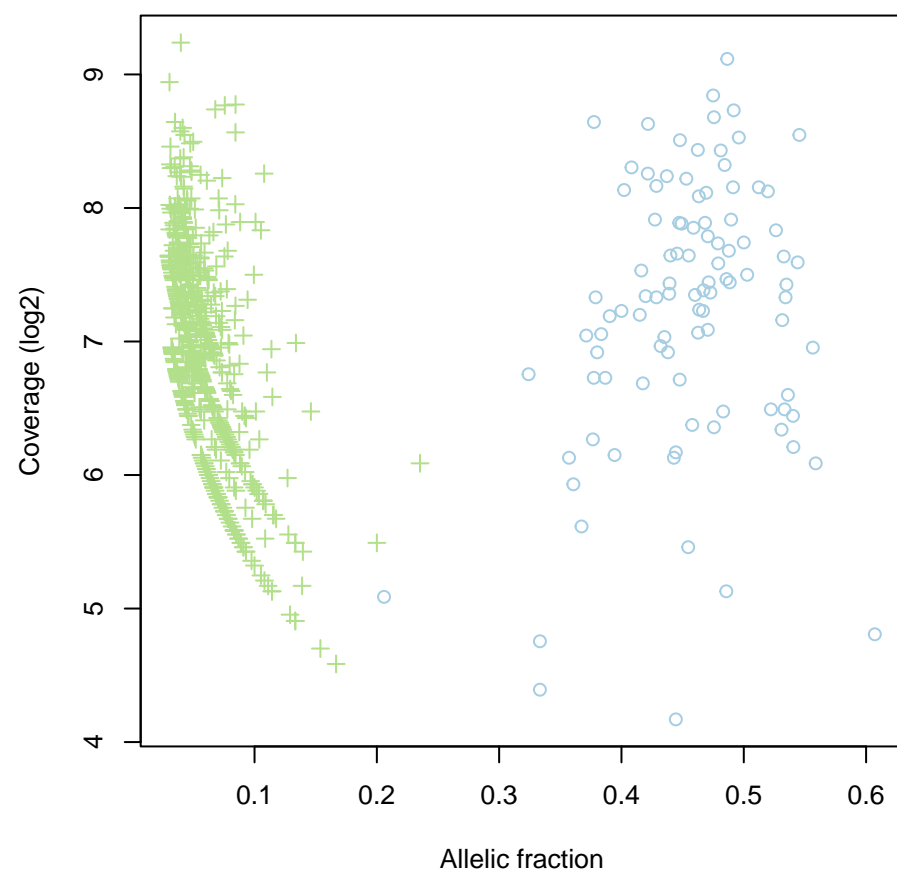
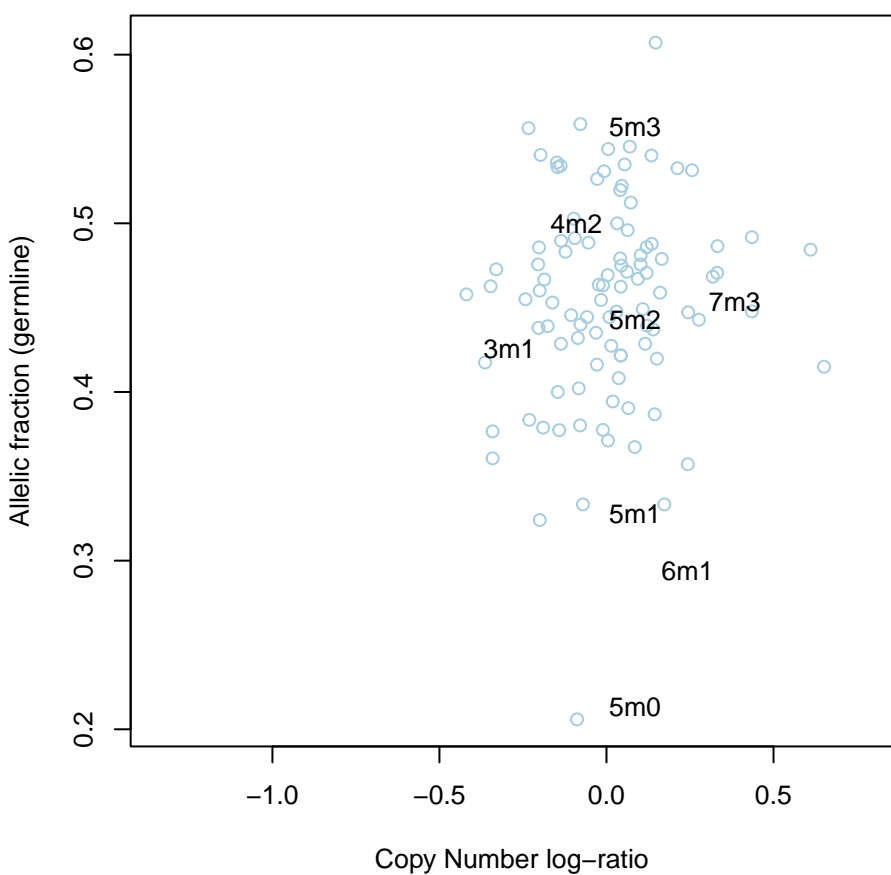


Purity: 0.35 Tumor ploidy: 4.496 SNV log-likelihood: -223.54 GoF: 88.3% Mean coverage: 363;148

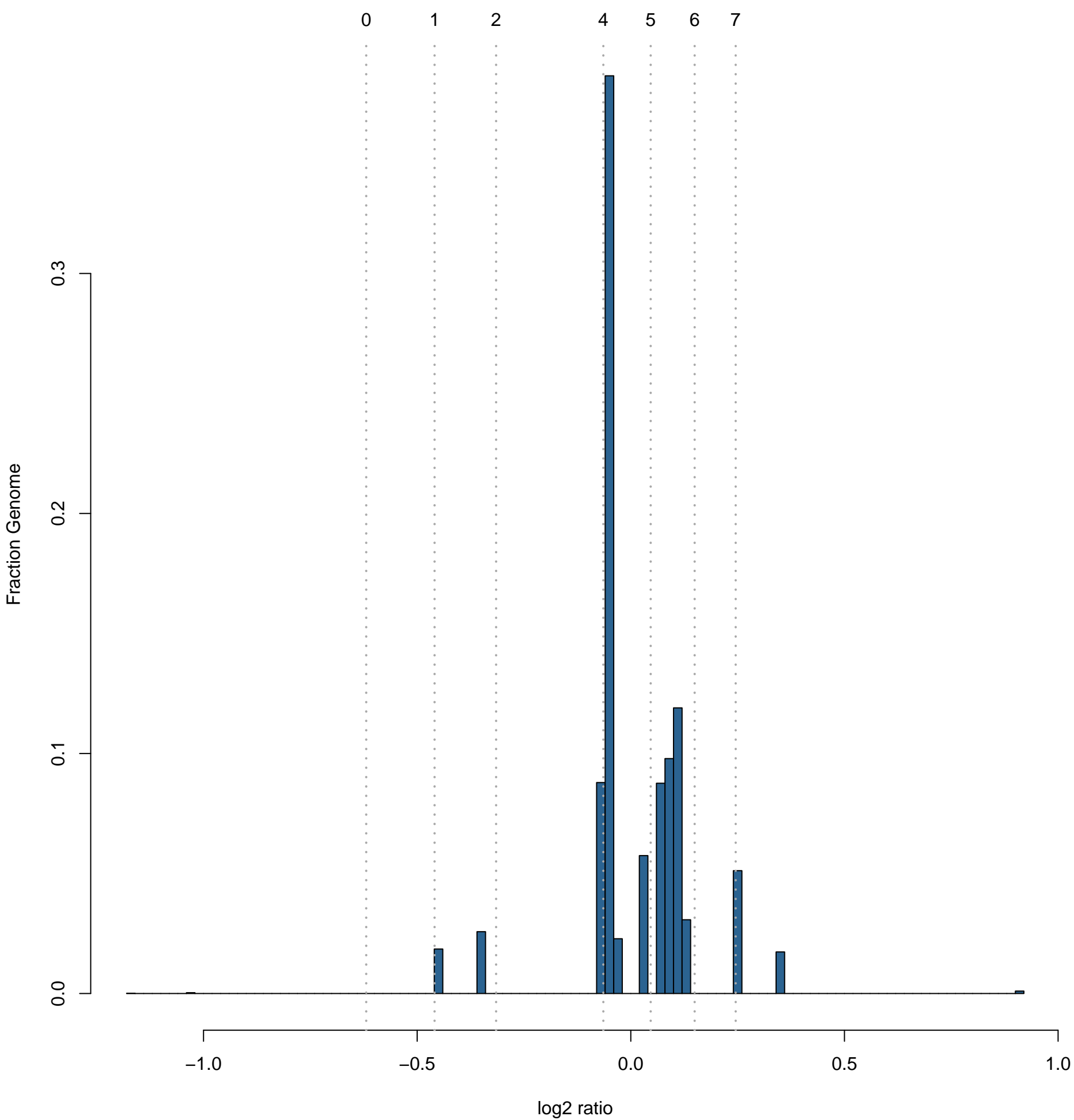


SCNA-fit log-likelihood: -19040.49



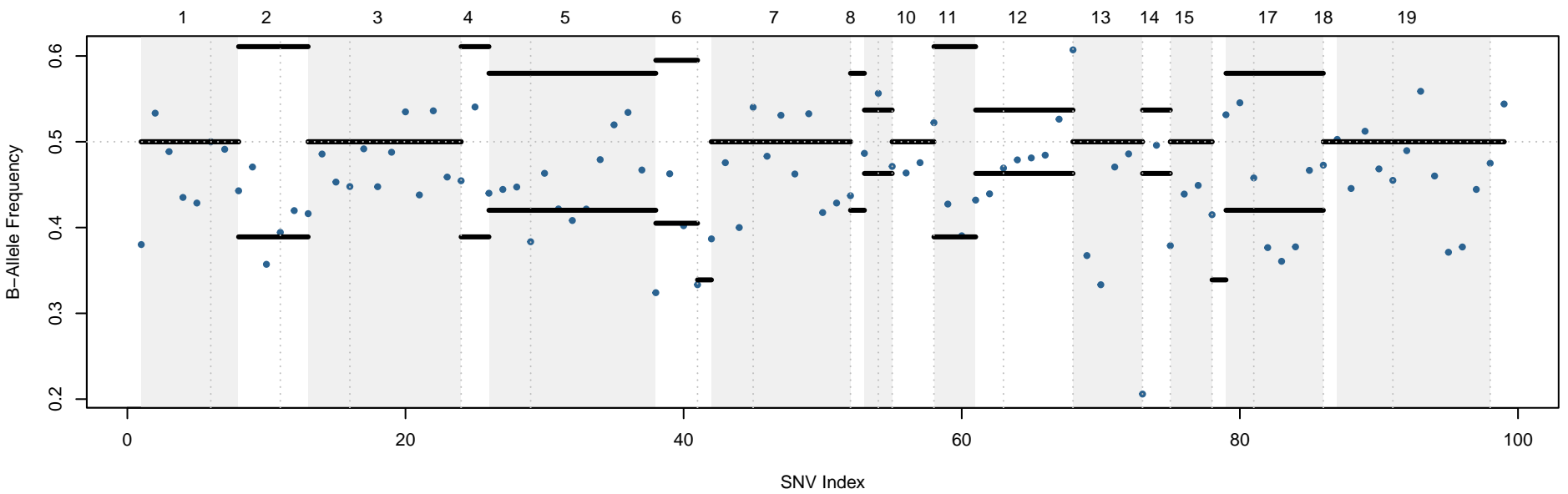


Purity: 0.19 Tumor ploidy: 4.571

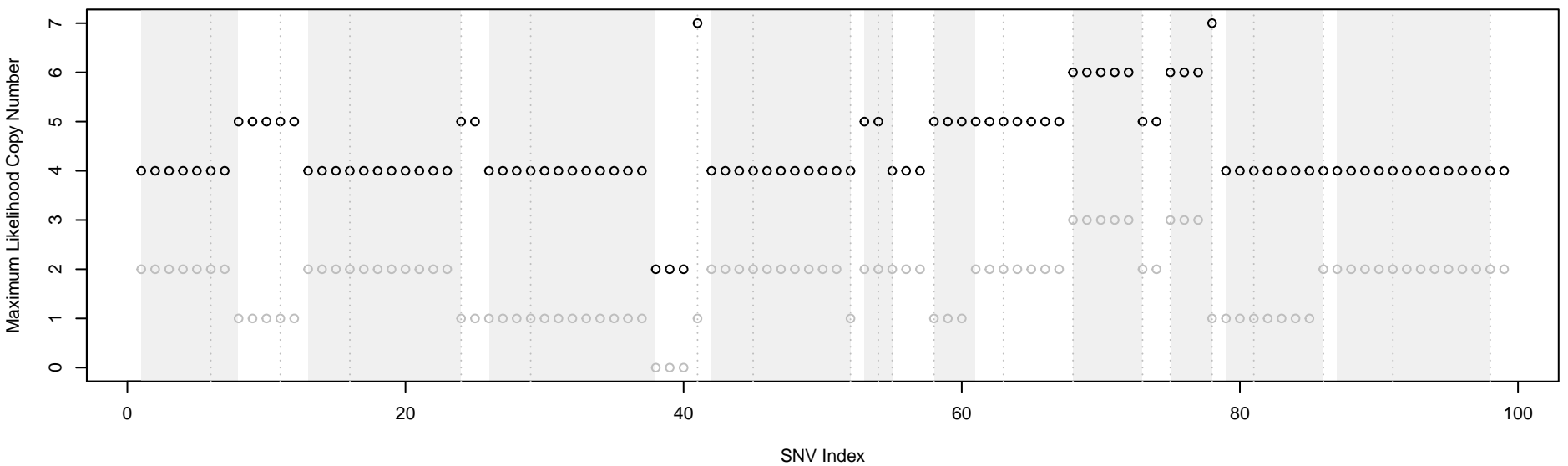
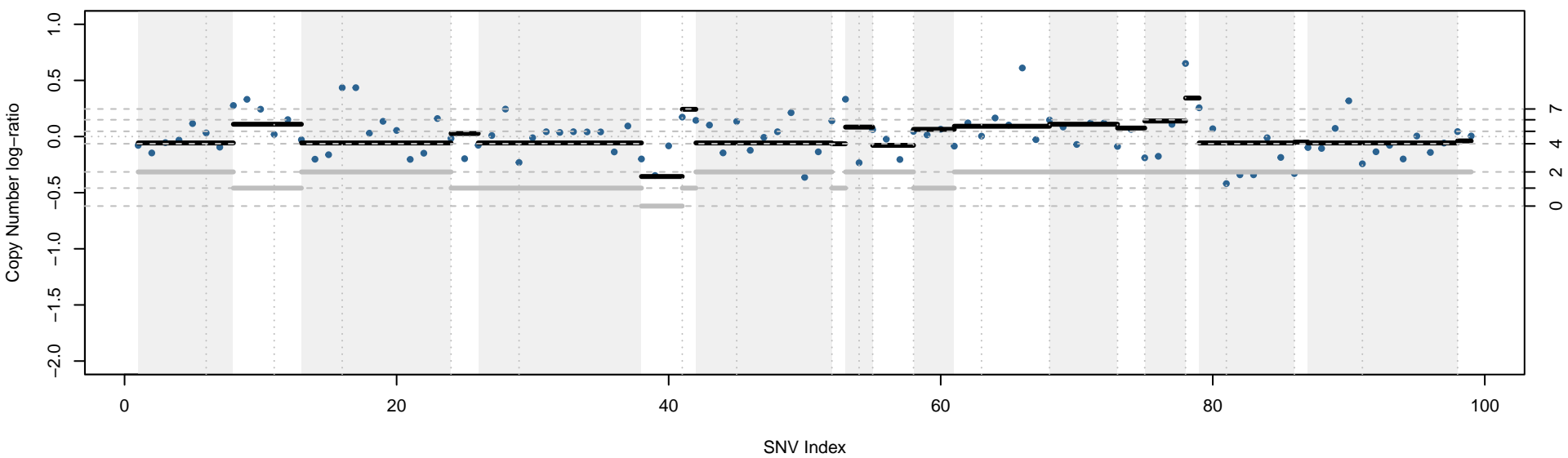


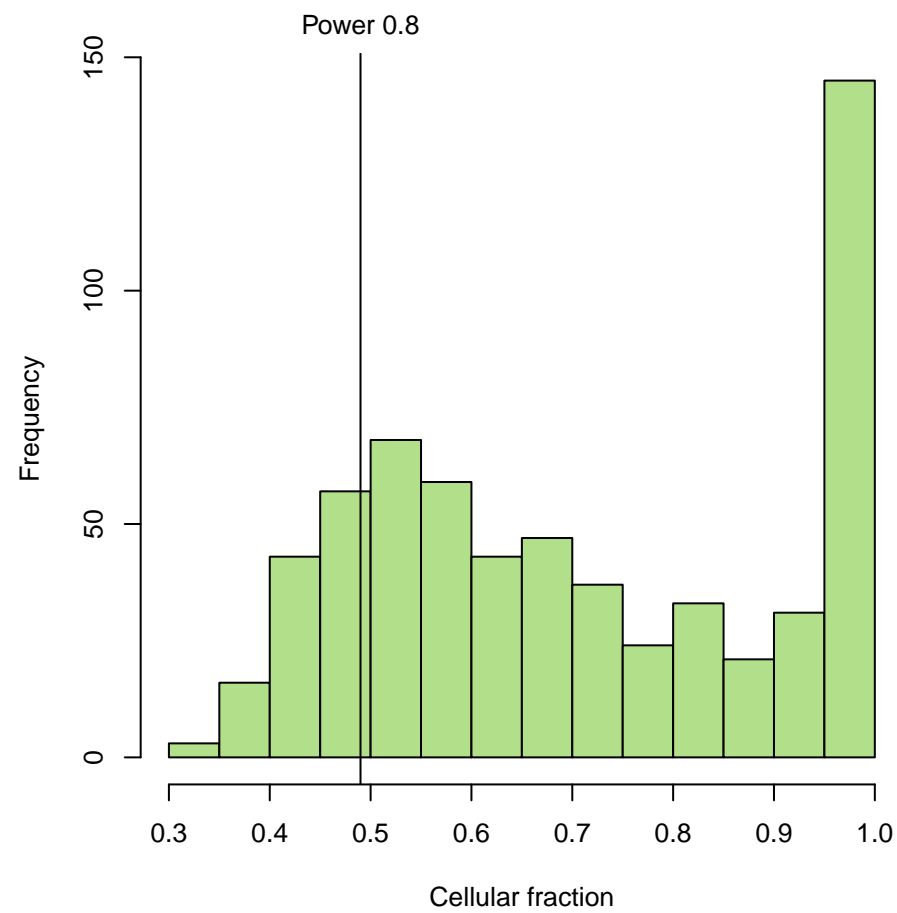
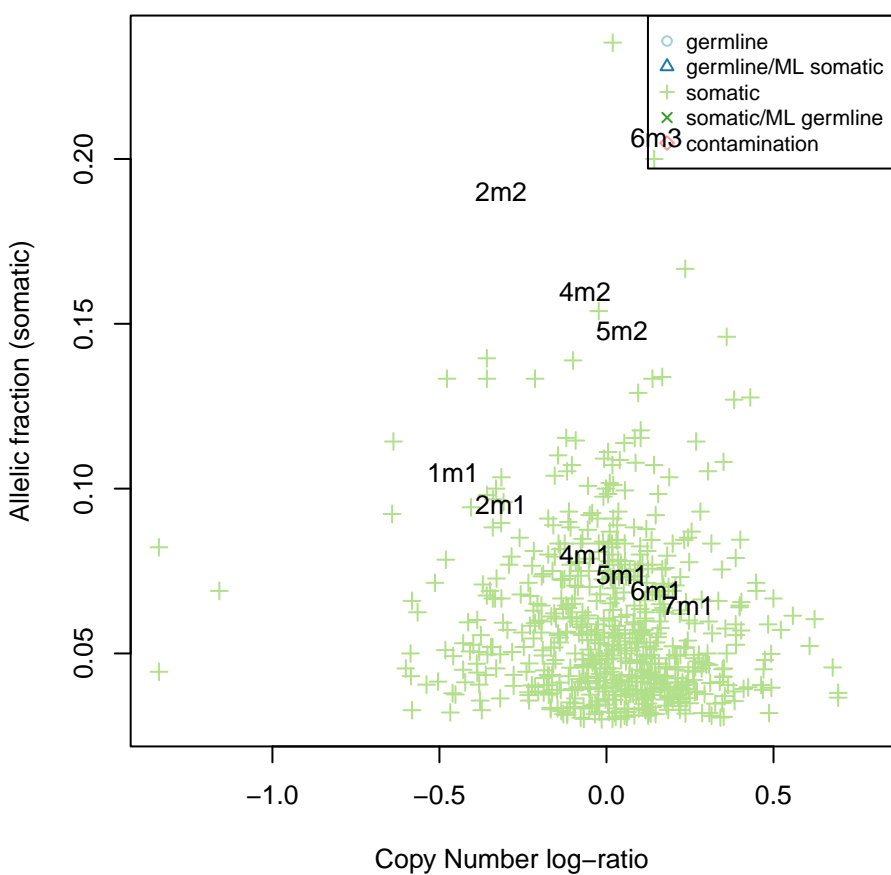
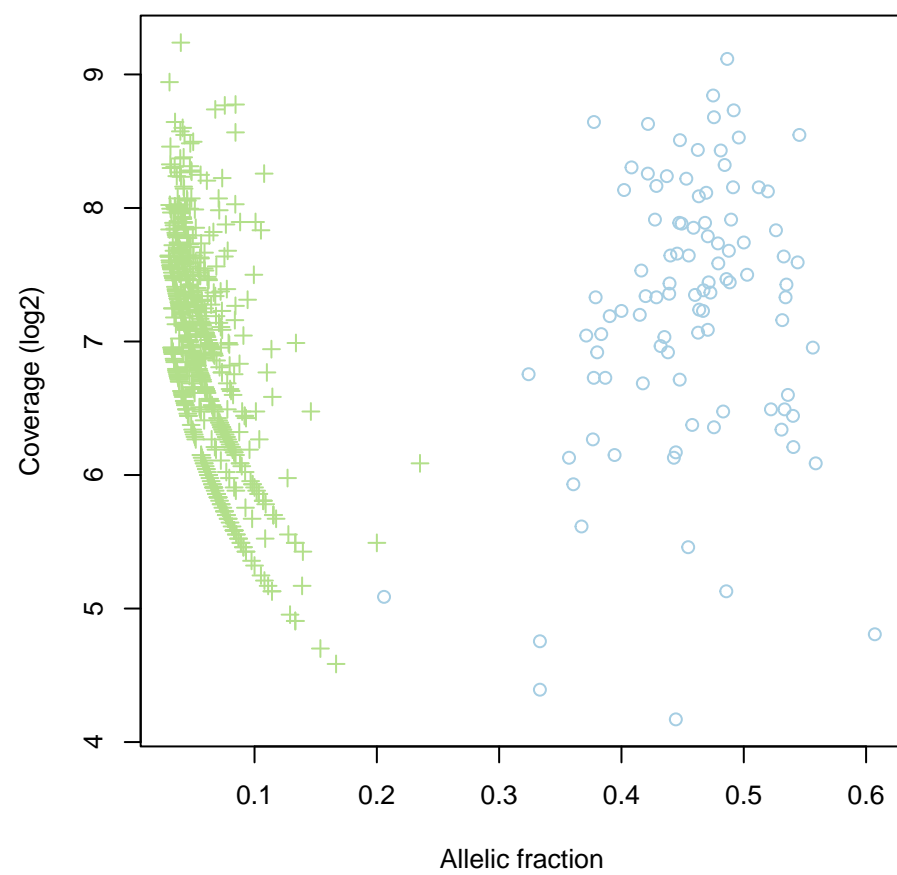
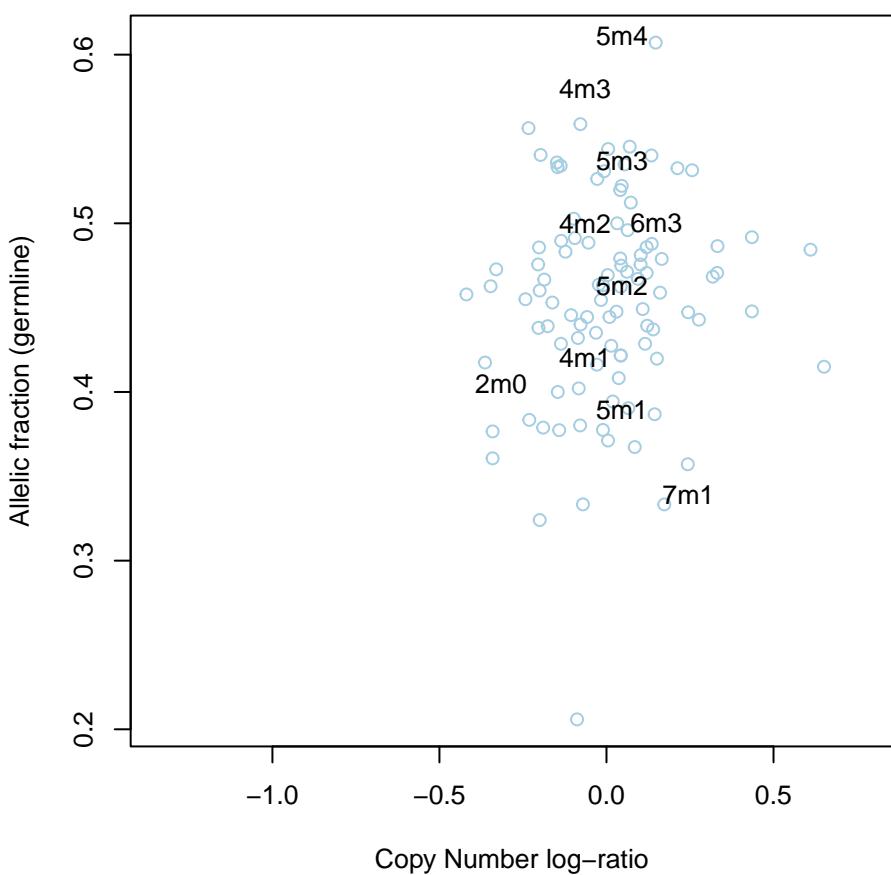


Purity: 0.19 Tumor ploidy: 4.571 SNV log-likelihood: -231.92 GoF: 96.9% Mean coverage: 363;148

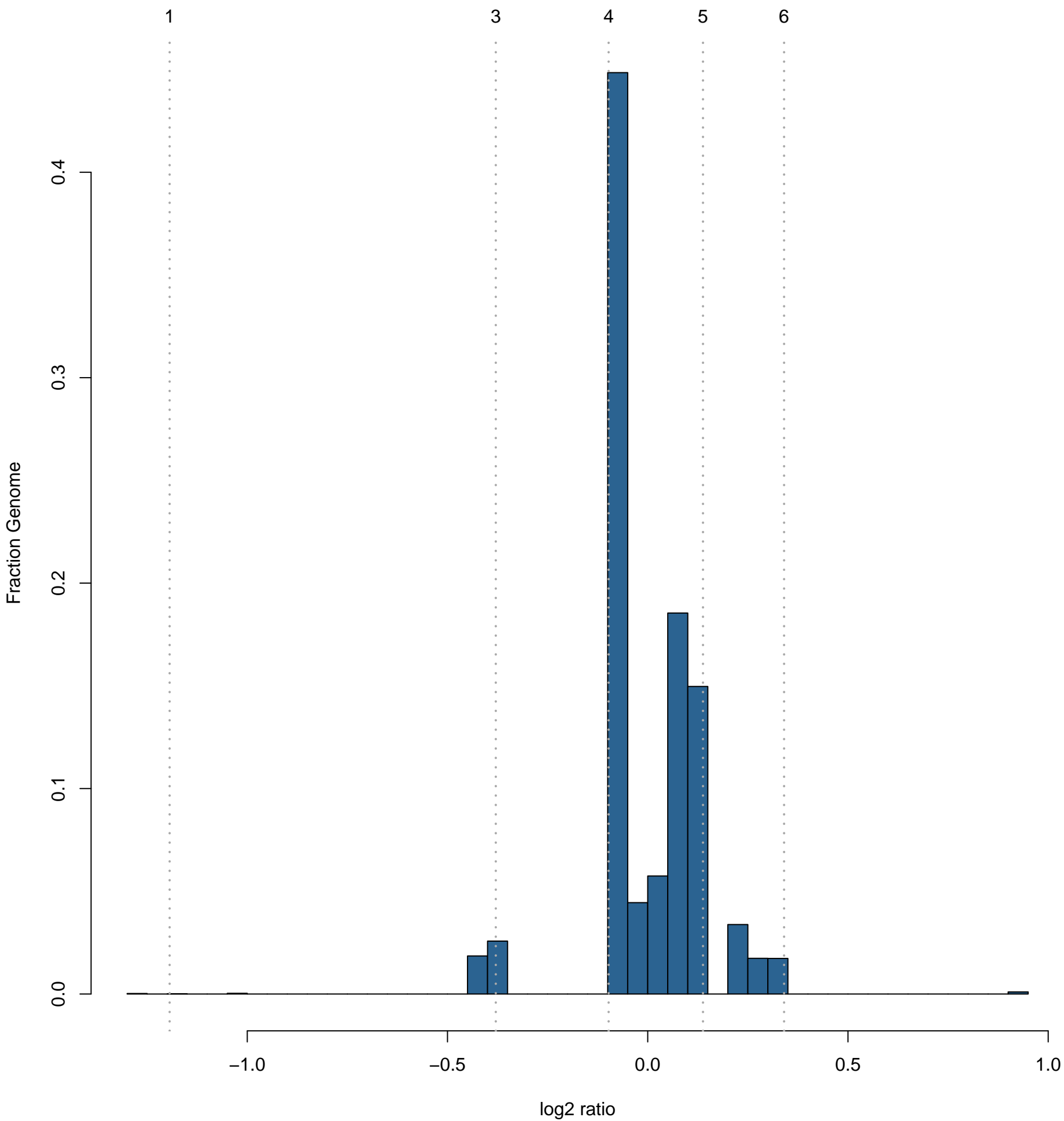


SCNA-fit log-likelihood: -19033.43

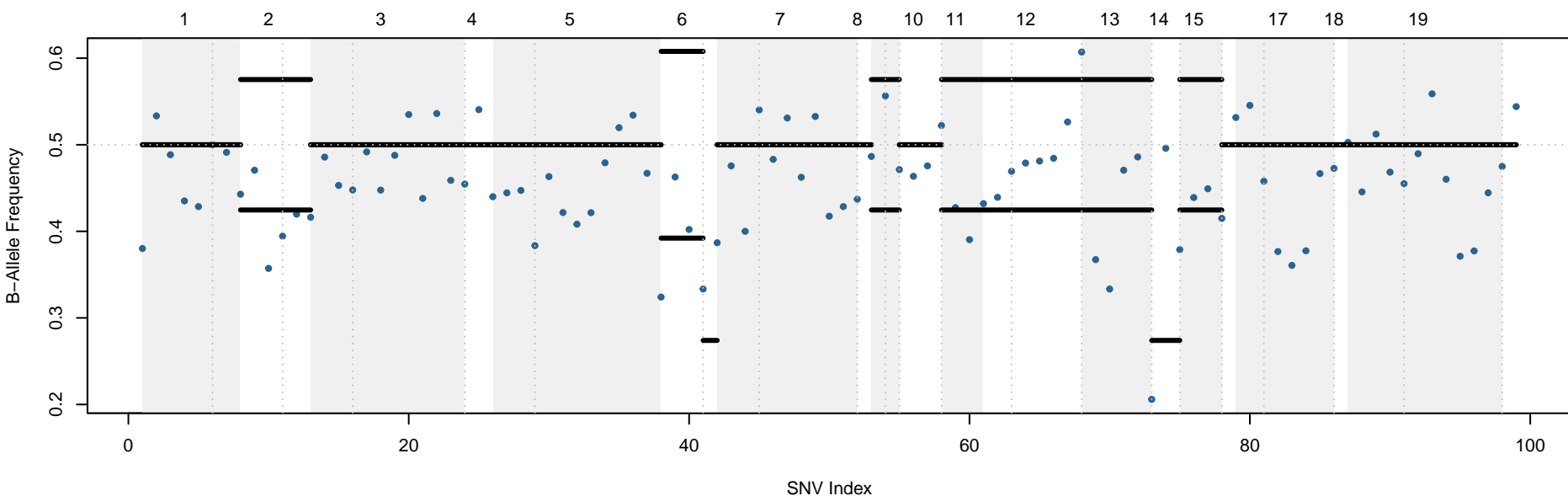




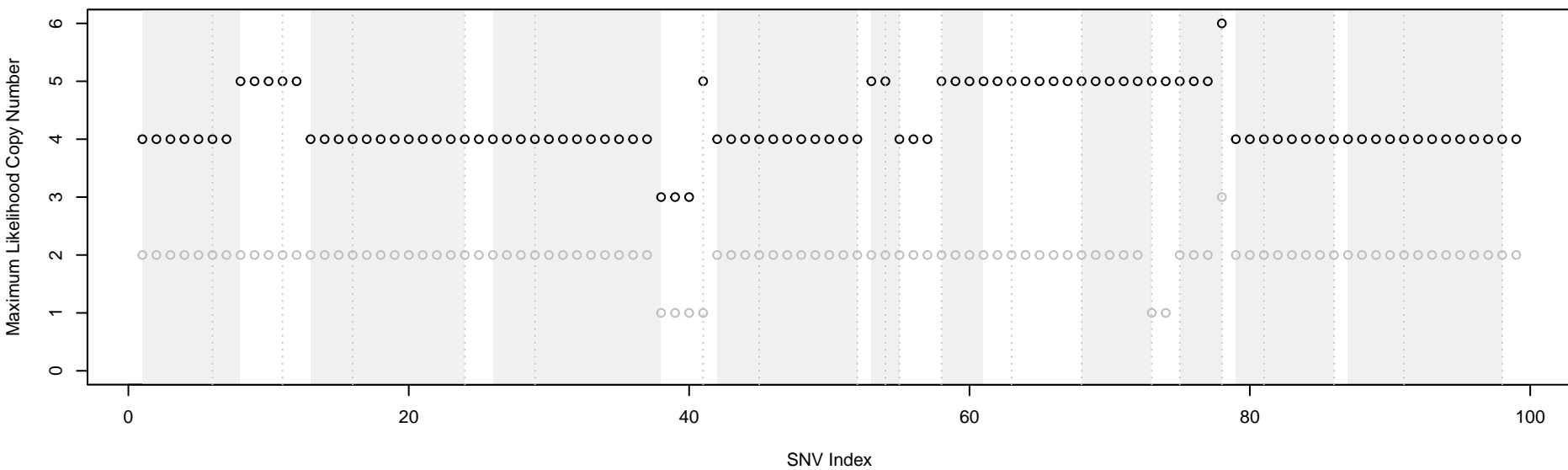
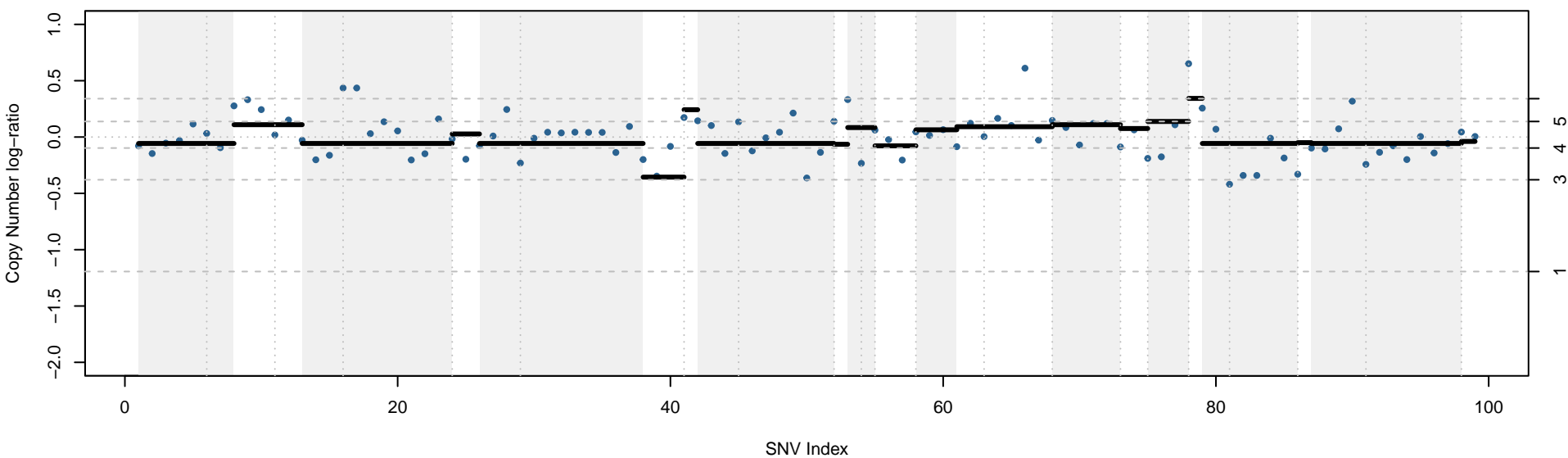
Purity: 0.55 Tumor ploidy: 4.395

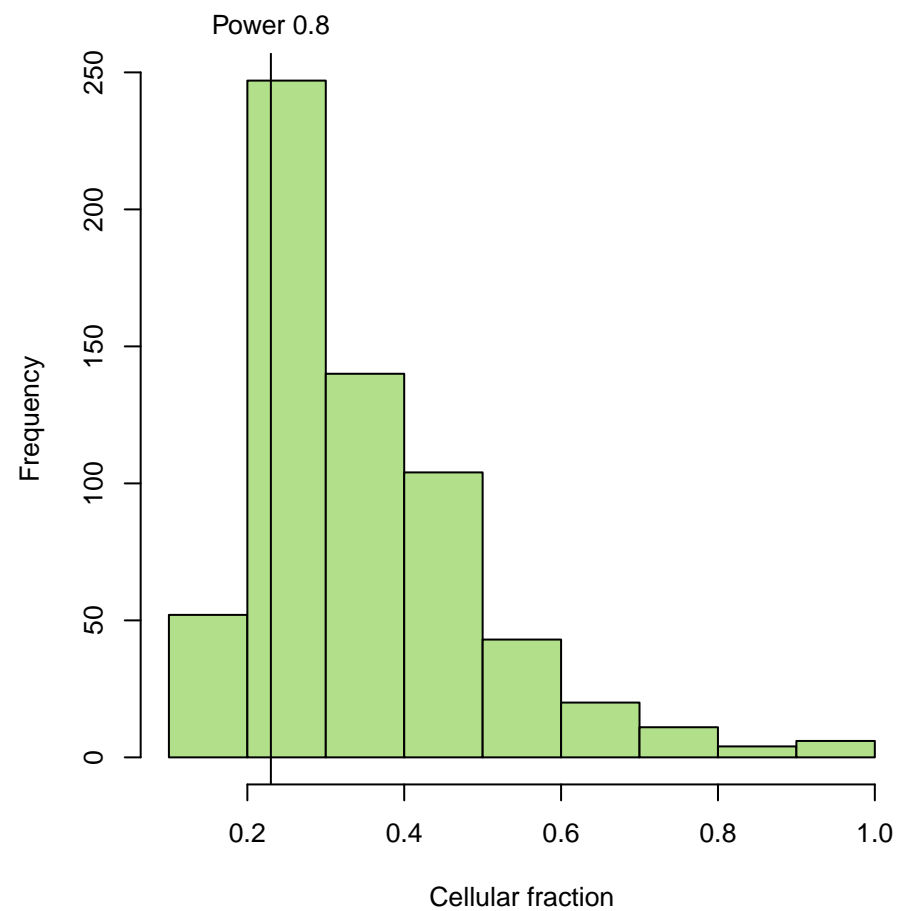
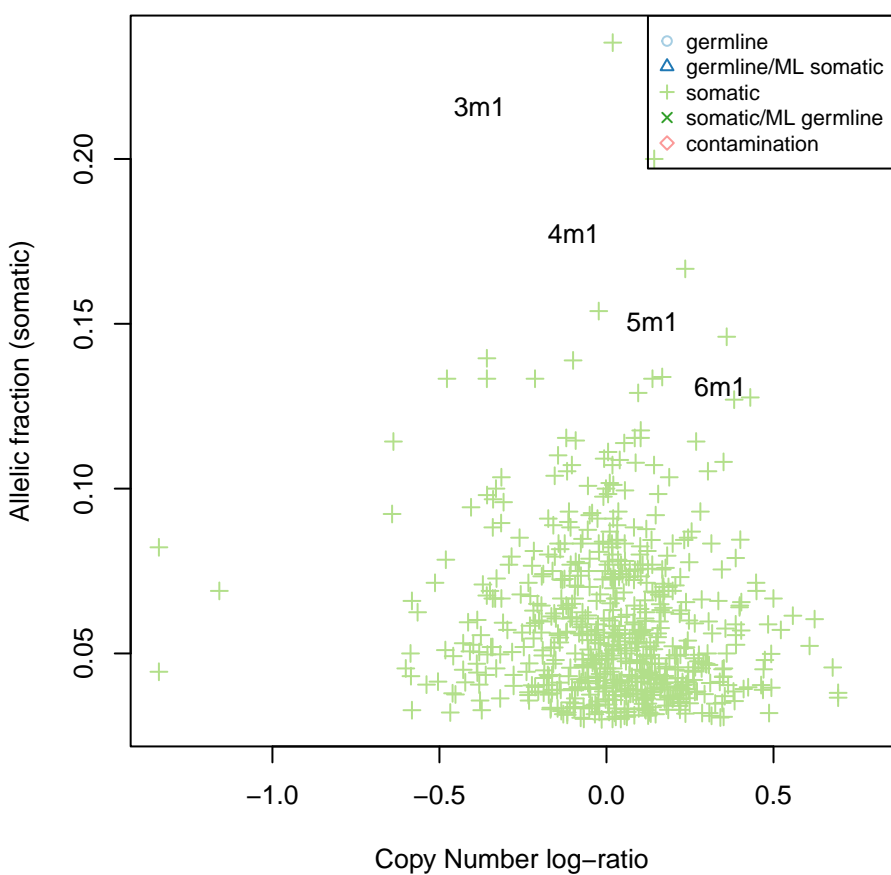
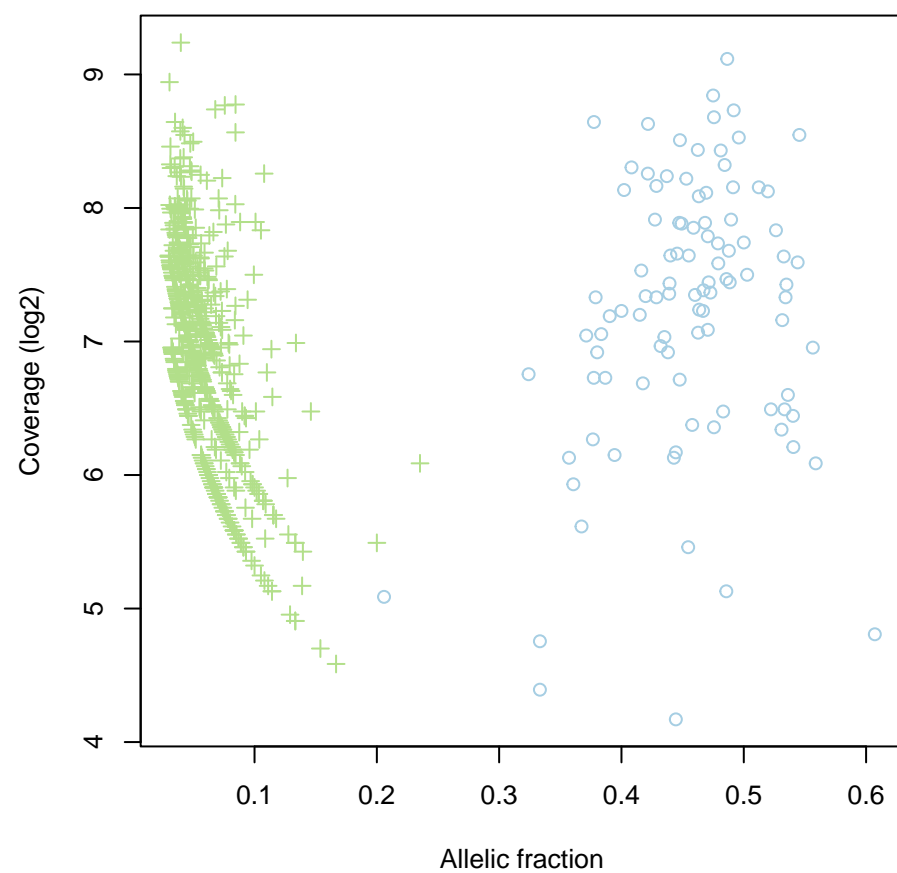
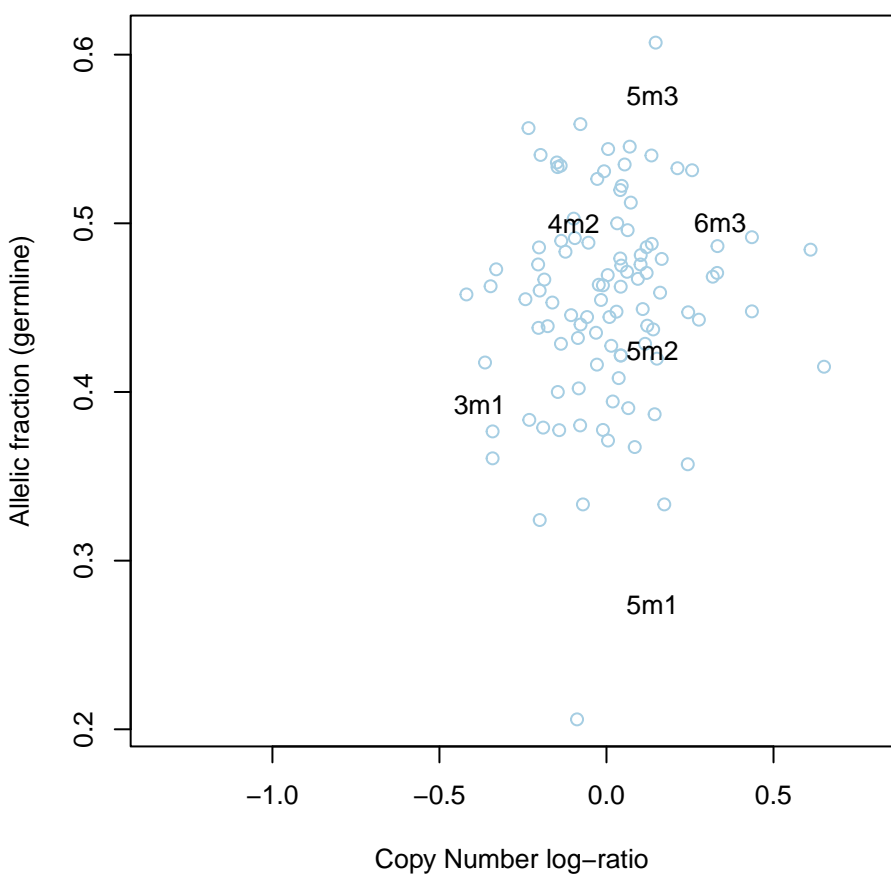


Purity: 0.55 Tumor ploidy: 4.395 SNV log-likelihood: -280.39 GoF: 72.1% Mean coverage: 363;148

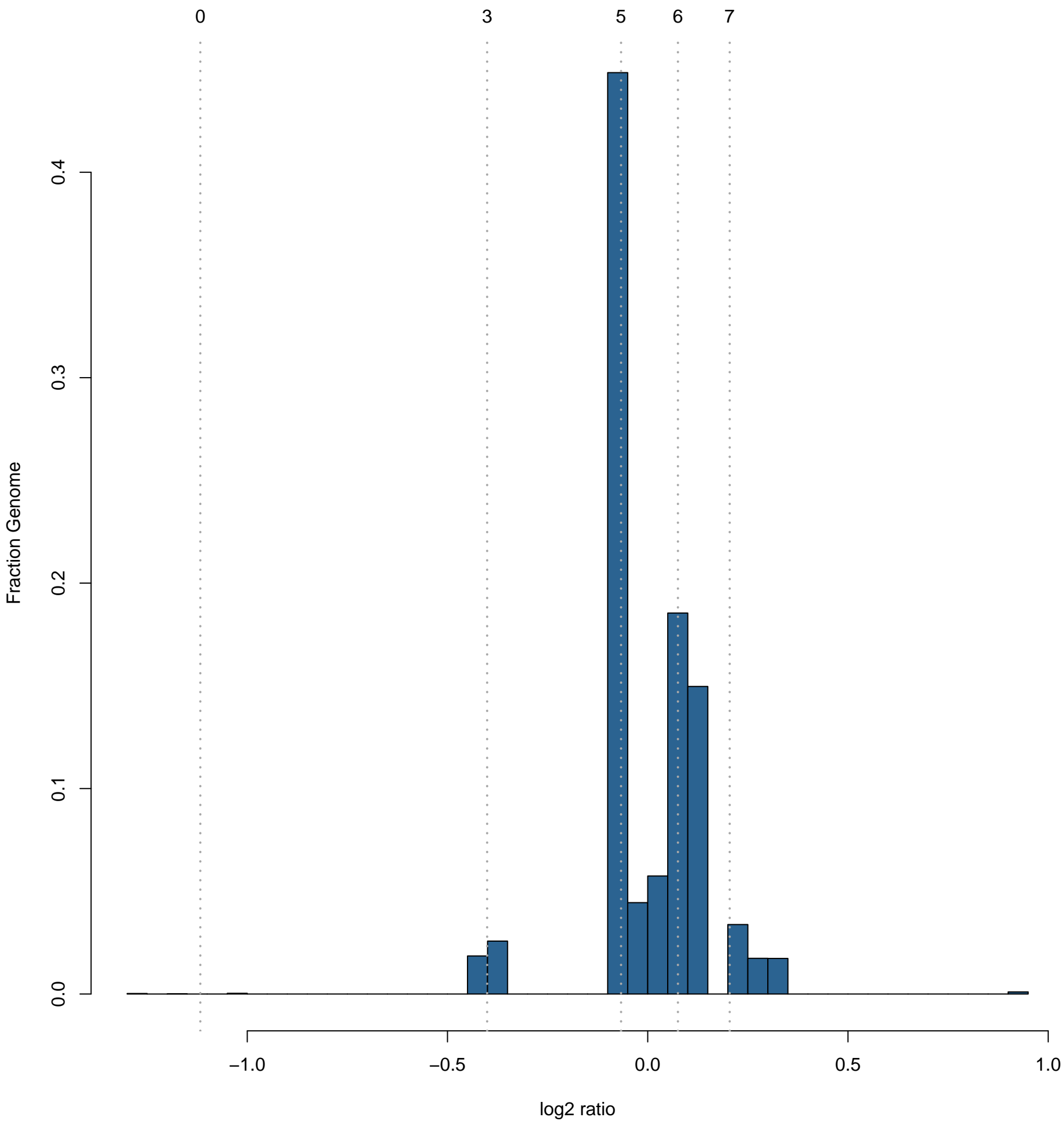


SCNA-fit log-likelihood: -19090.56

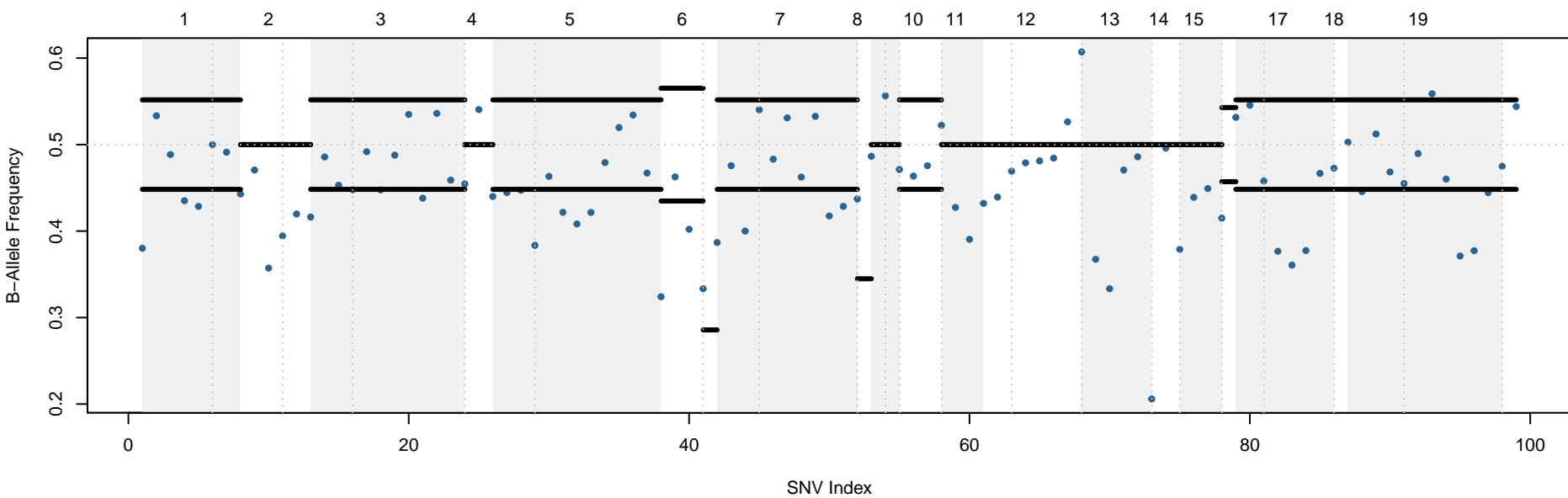




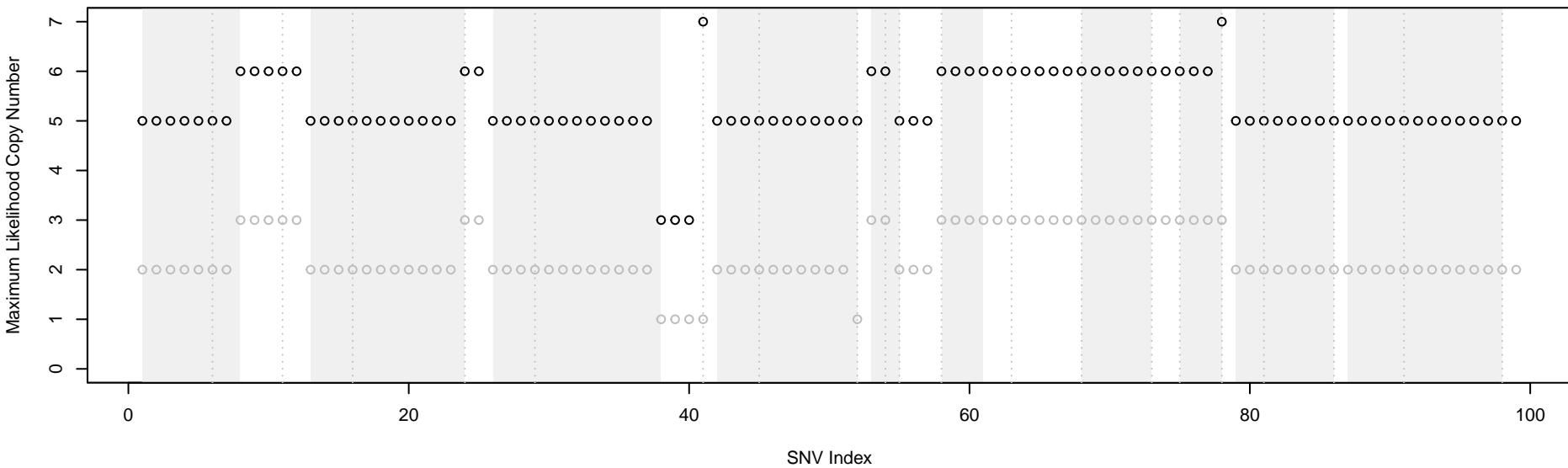
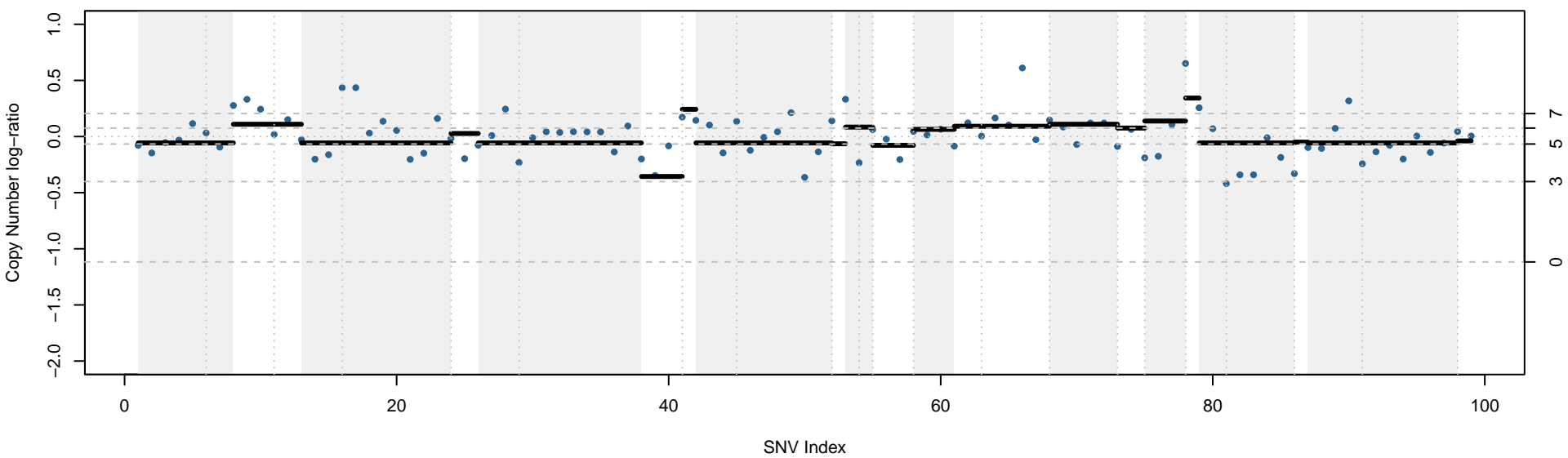
Purity: 0.3 Tumor ploidy: 5.457

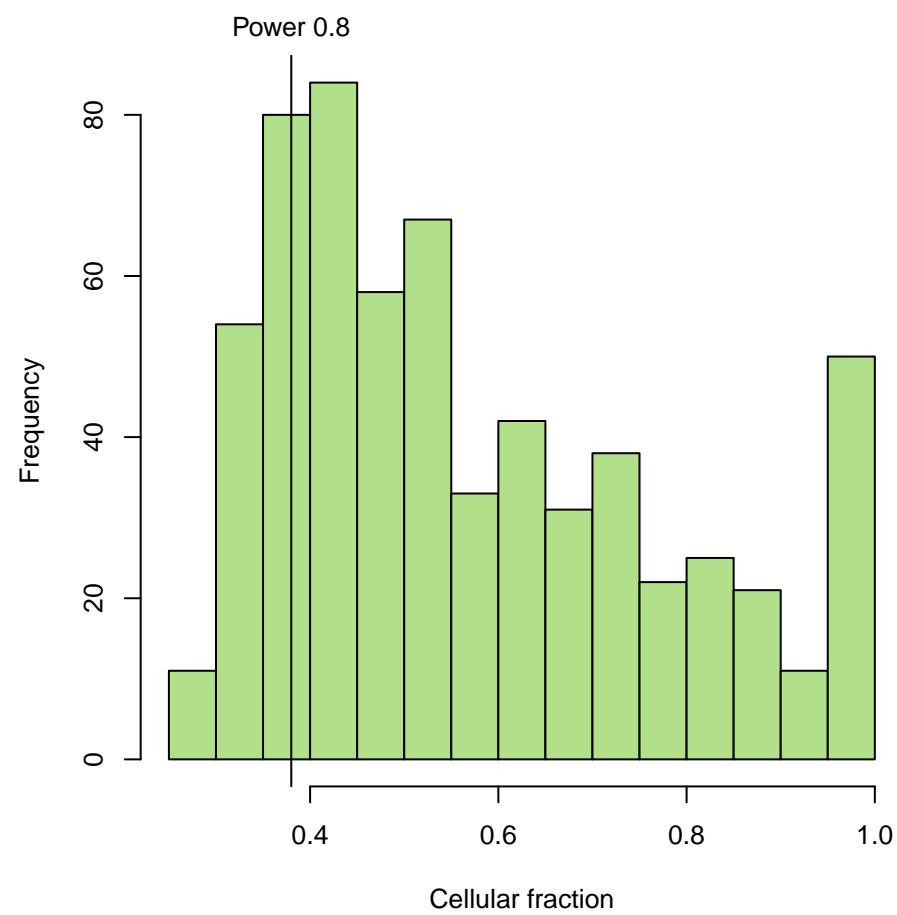
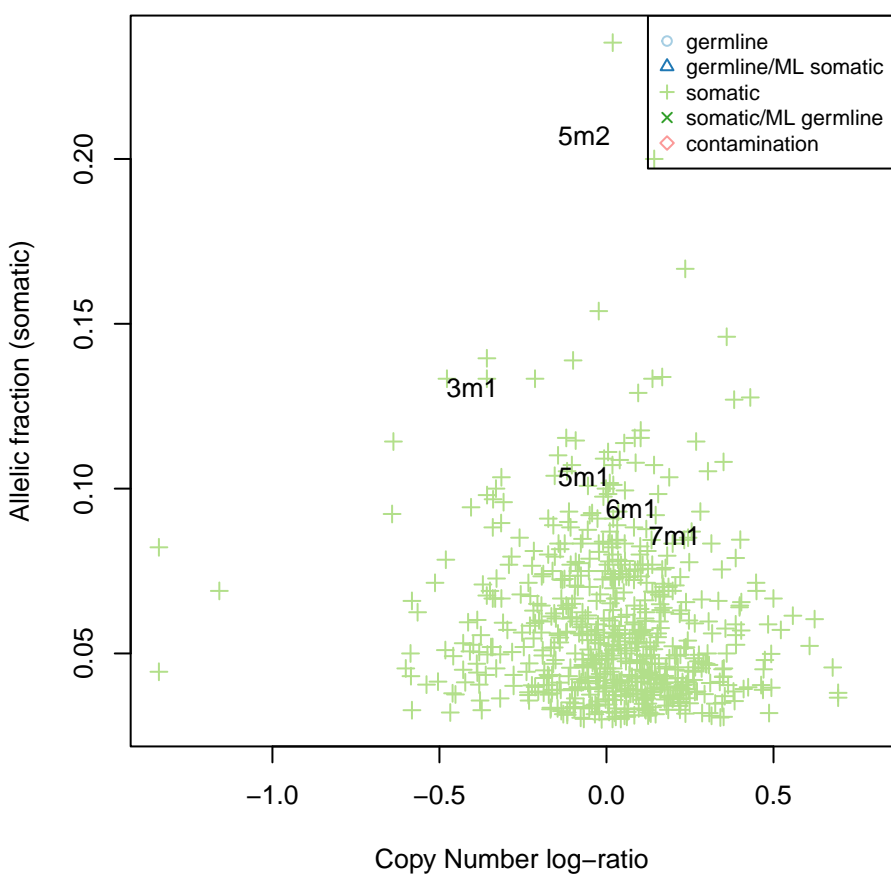
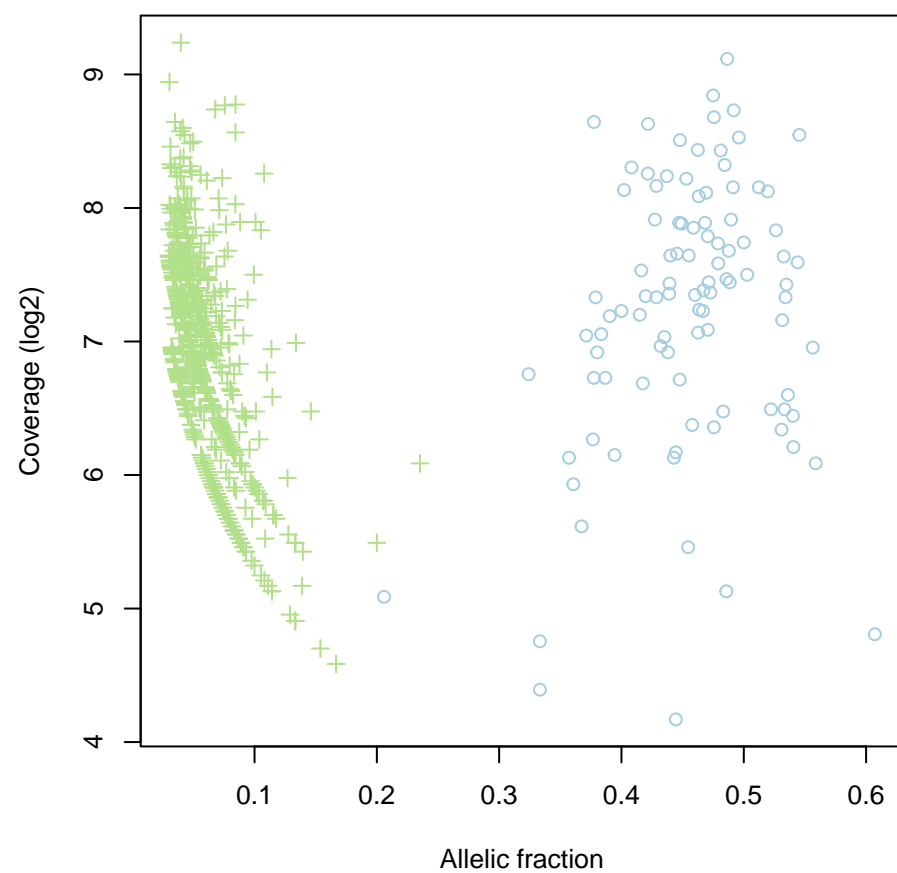
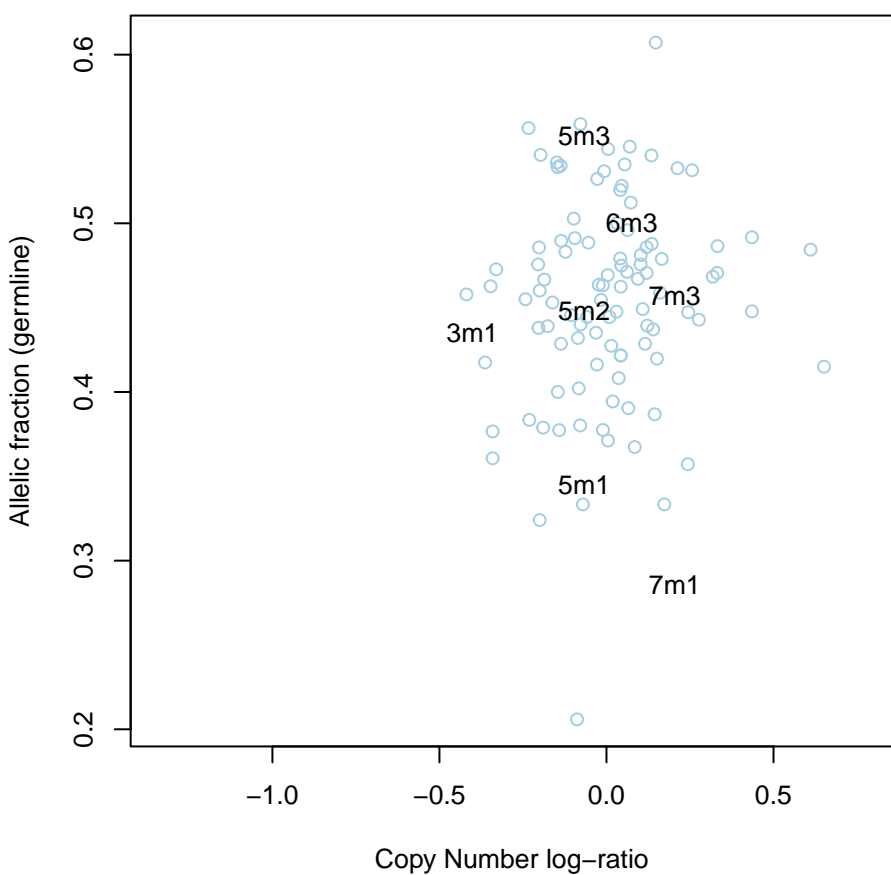


Purity: 0.3 Tumor ploidy: 5.457 SNV log-likelihood: -383.44 GoF: 94% Mean coverage: 363;148



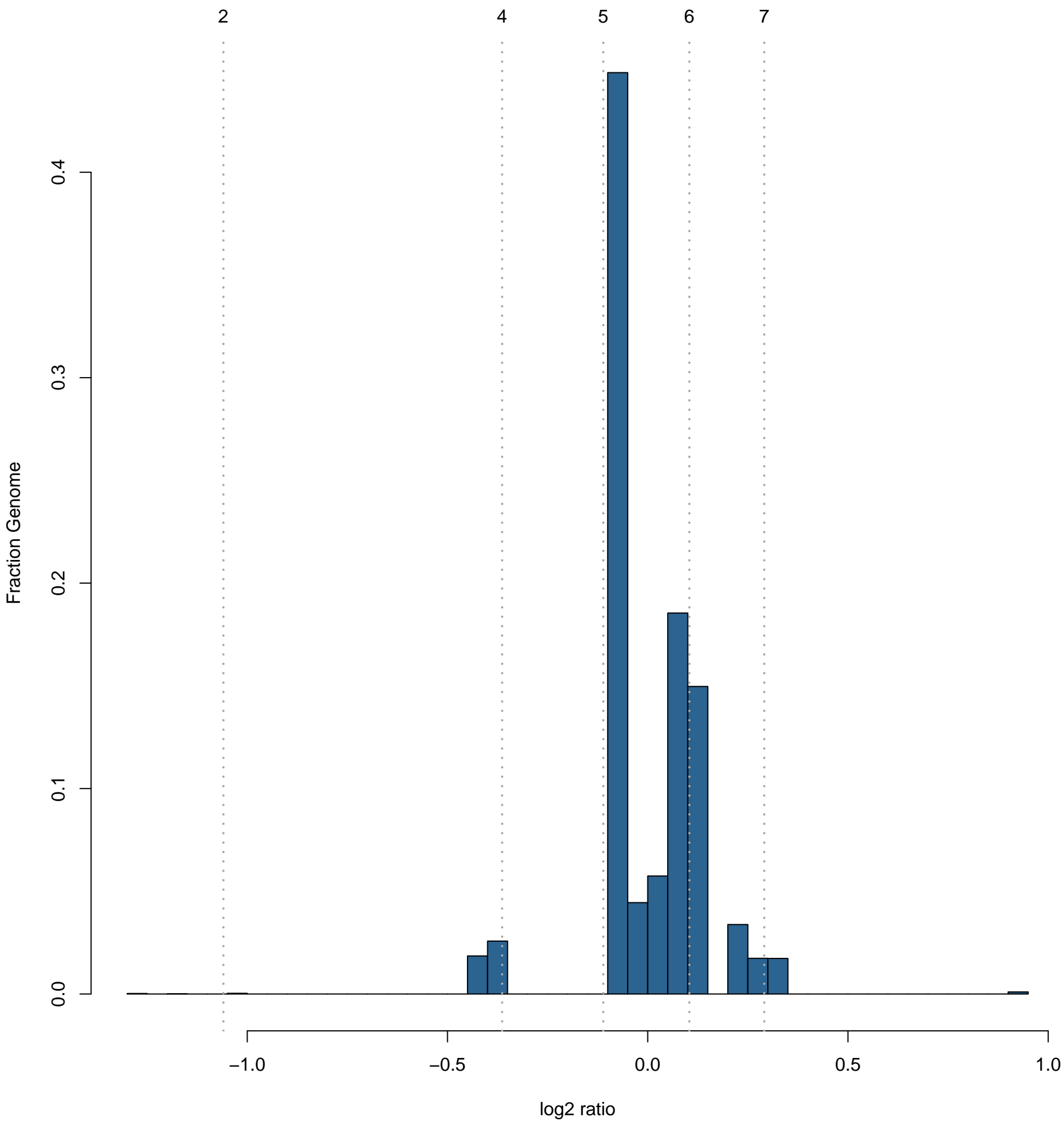
SCNA-fit log-likelihood: -19034.5



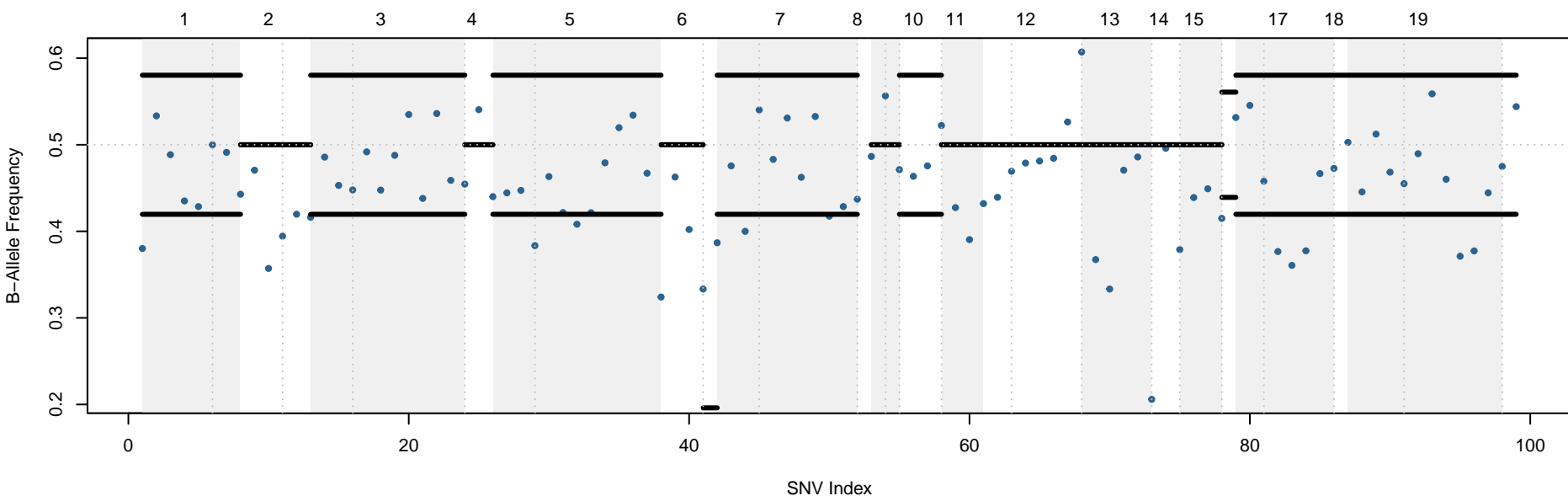




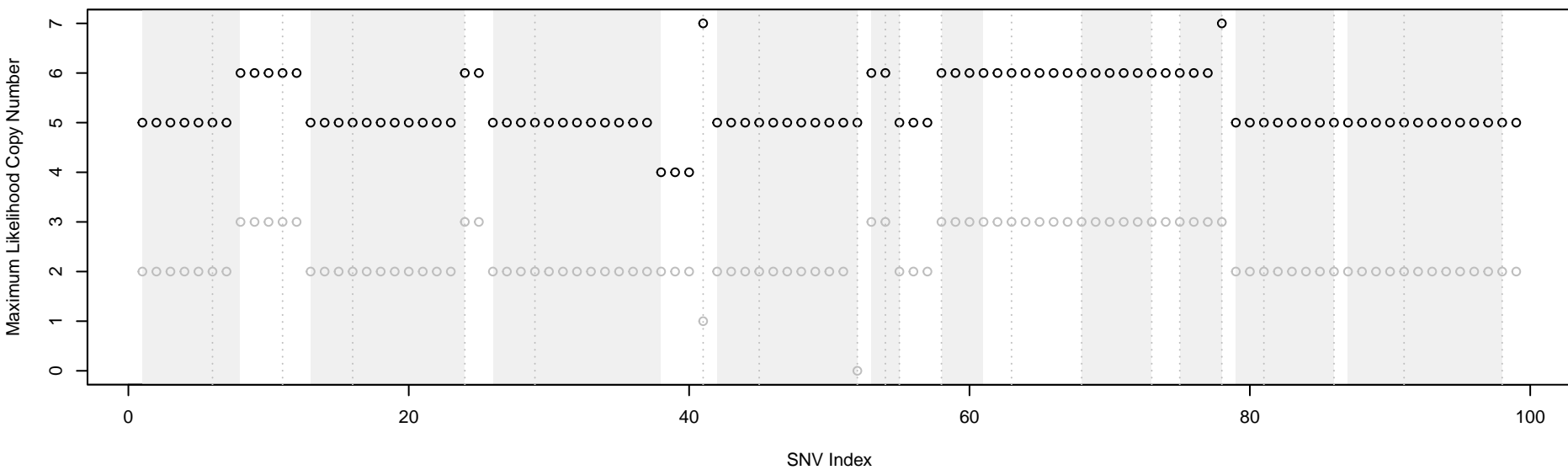
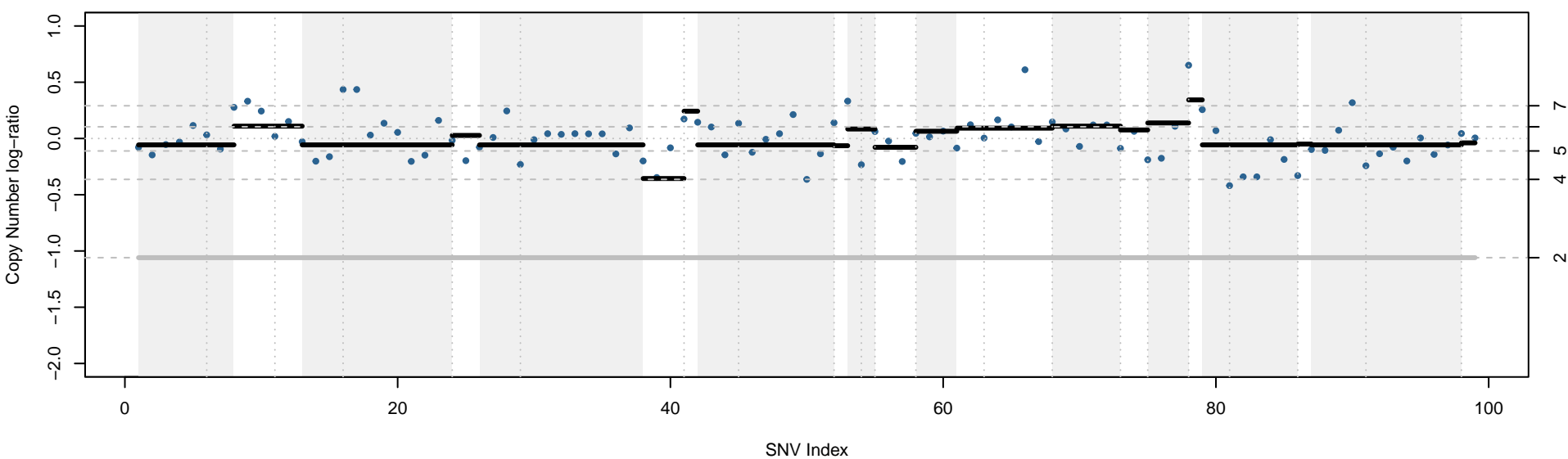
Purity: 0.62 Tumor ploidy: 5.498

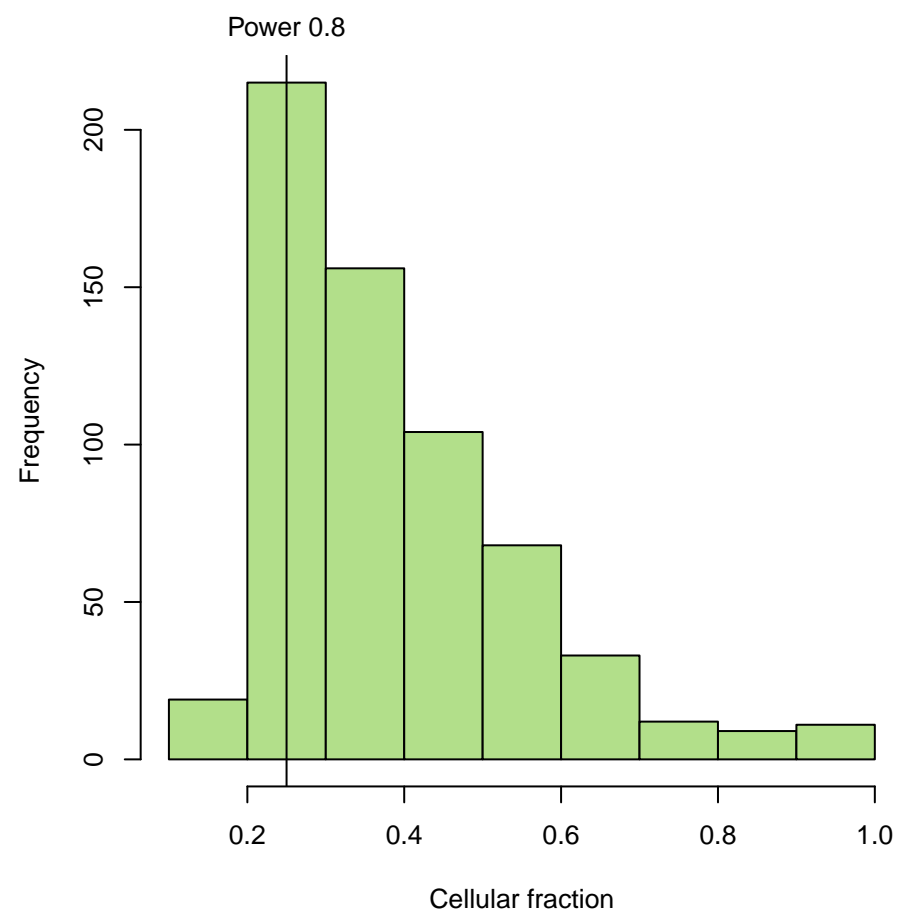
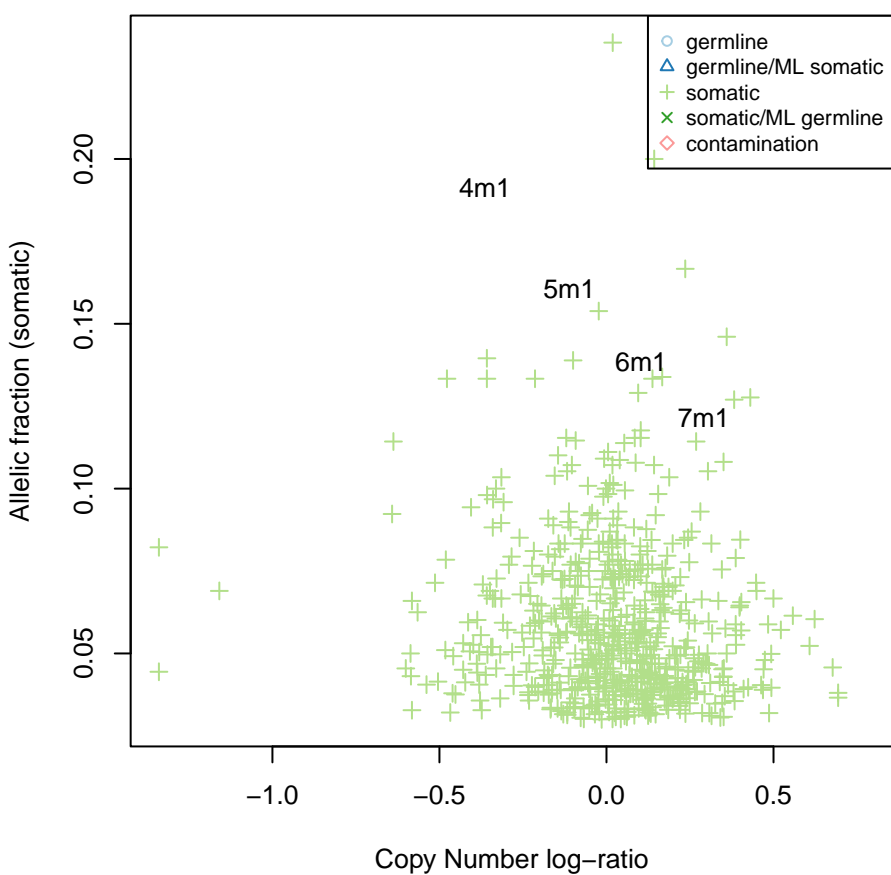
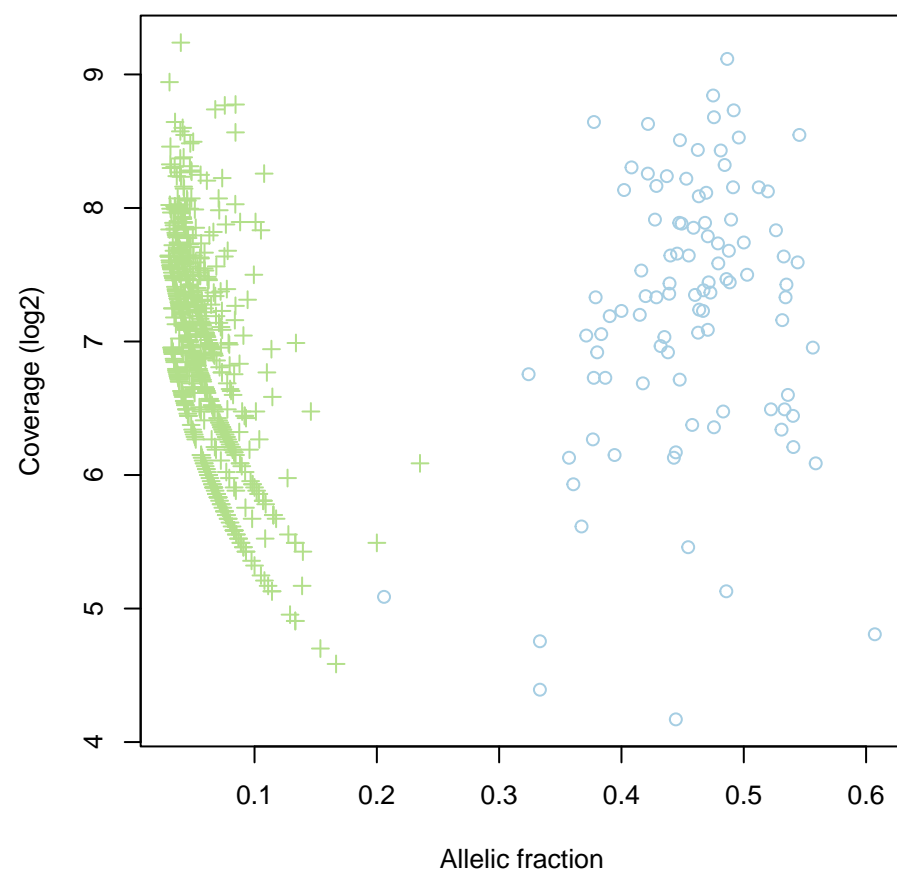
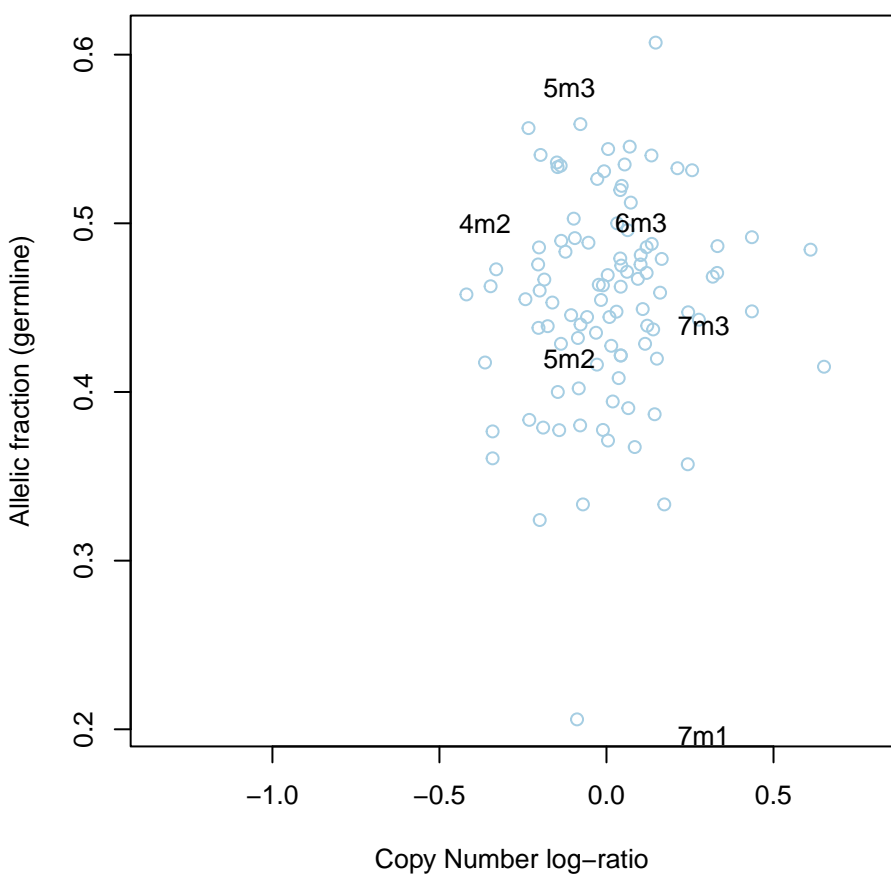


Purity: 0.62 Tumor ploidy: 5.498 SNV log-likelihood: -430.53 GoF: 79% Mean coverage: 363;148

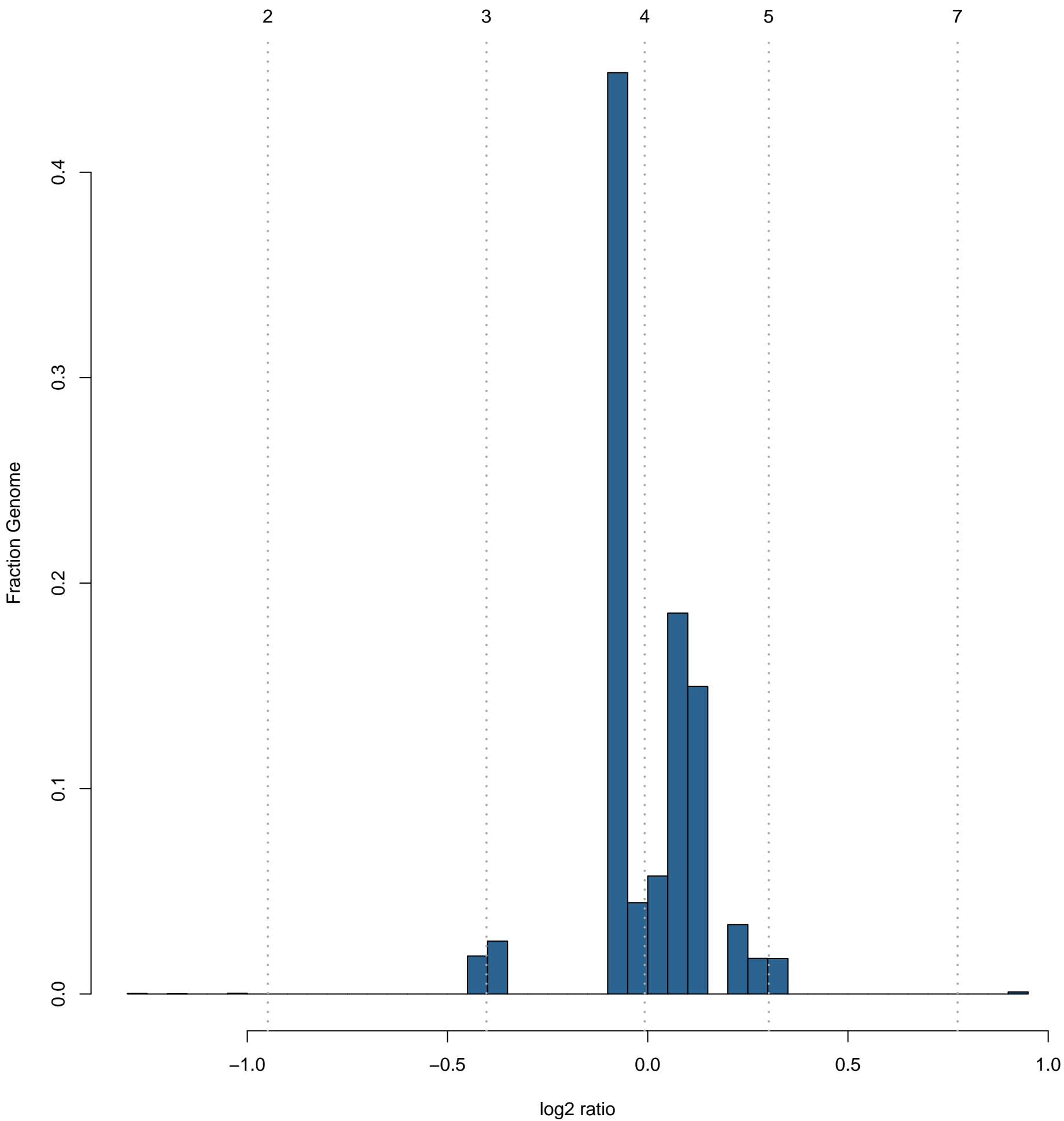


SCNA-fit log-likelihood: -19064.66

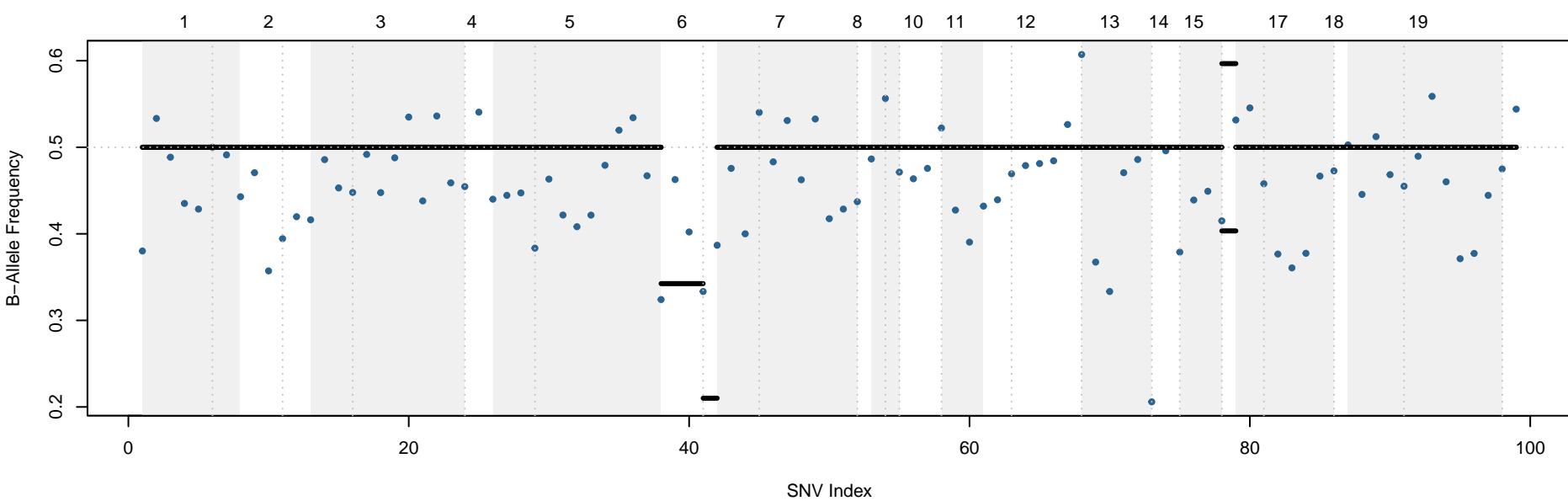




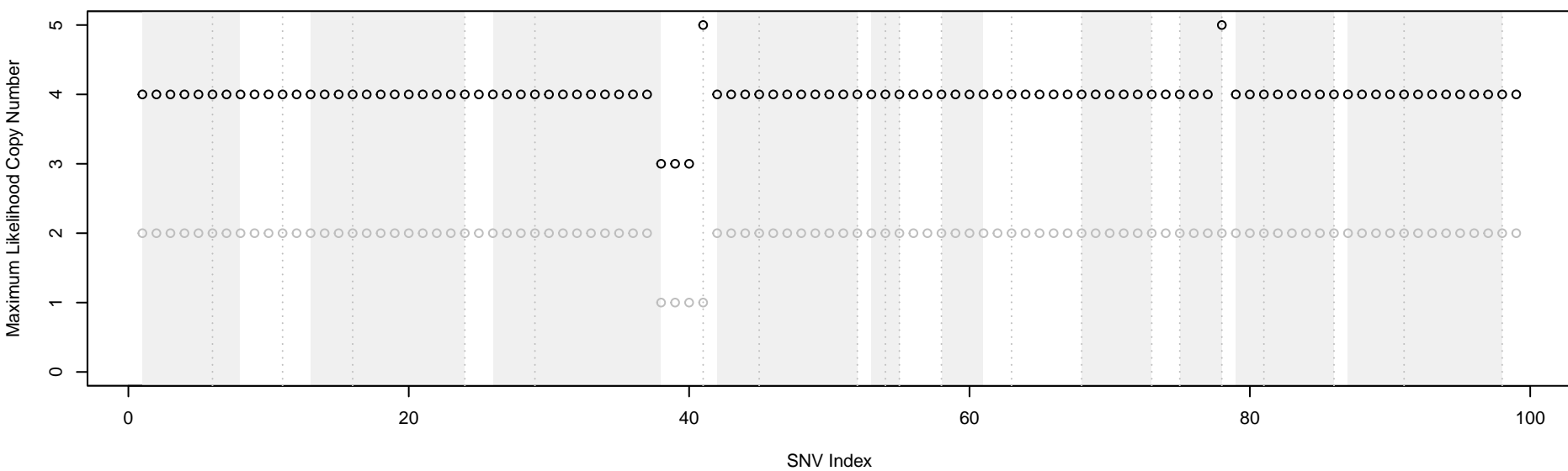
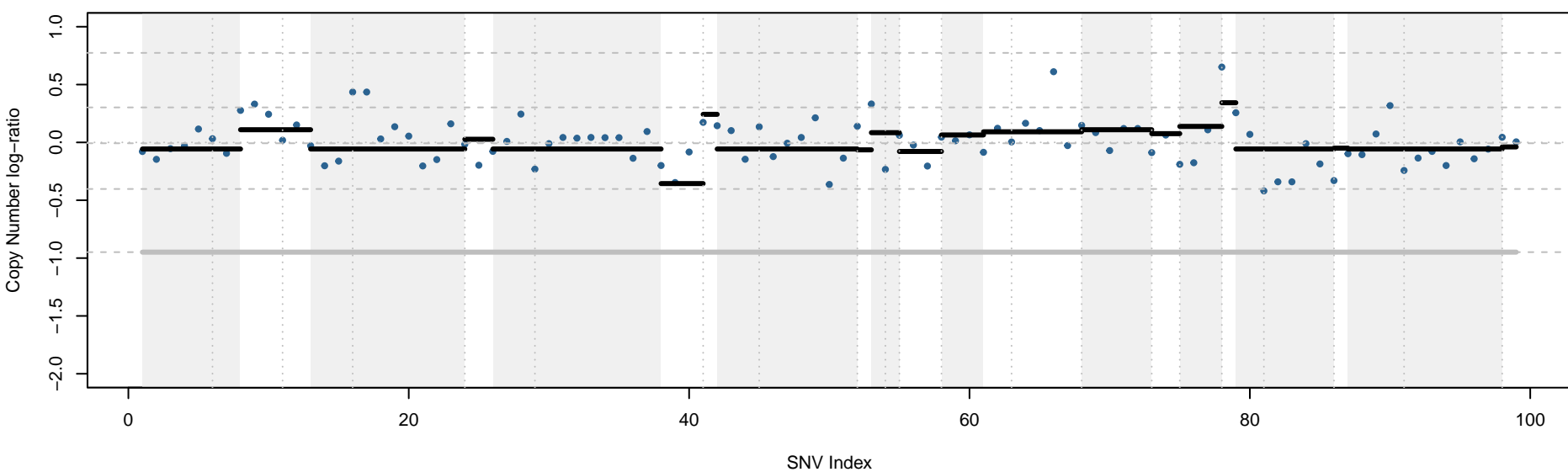
Purity: 0.92 Tumor ploidy: 4.022

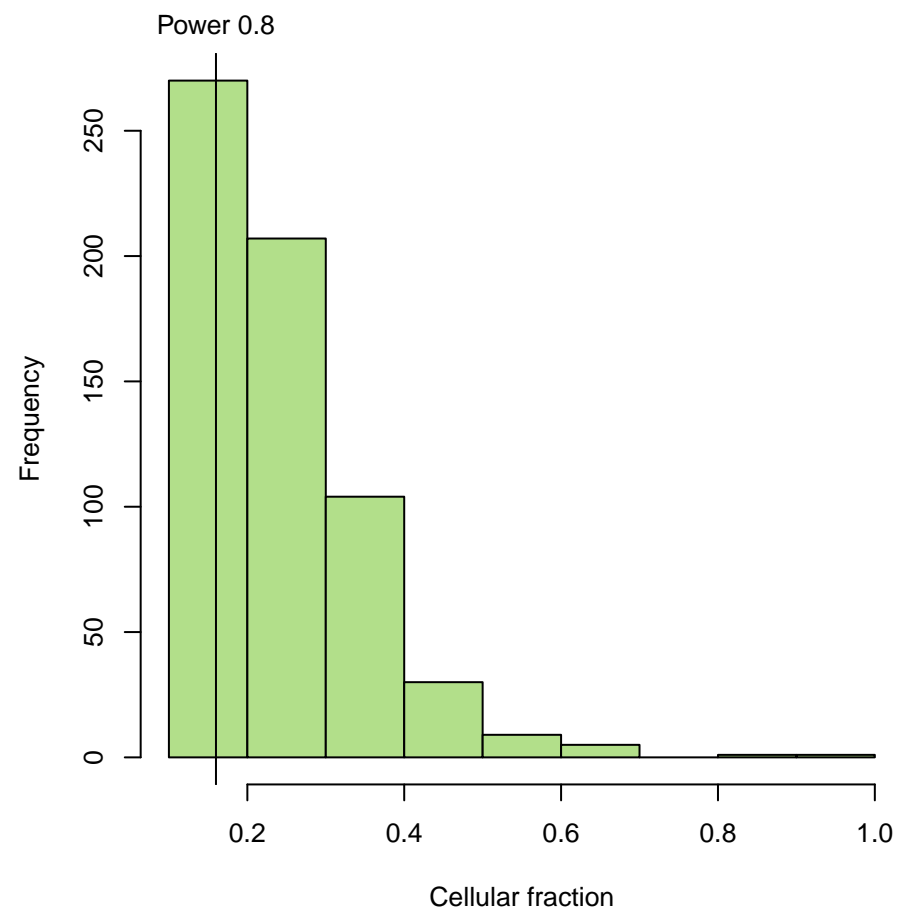
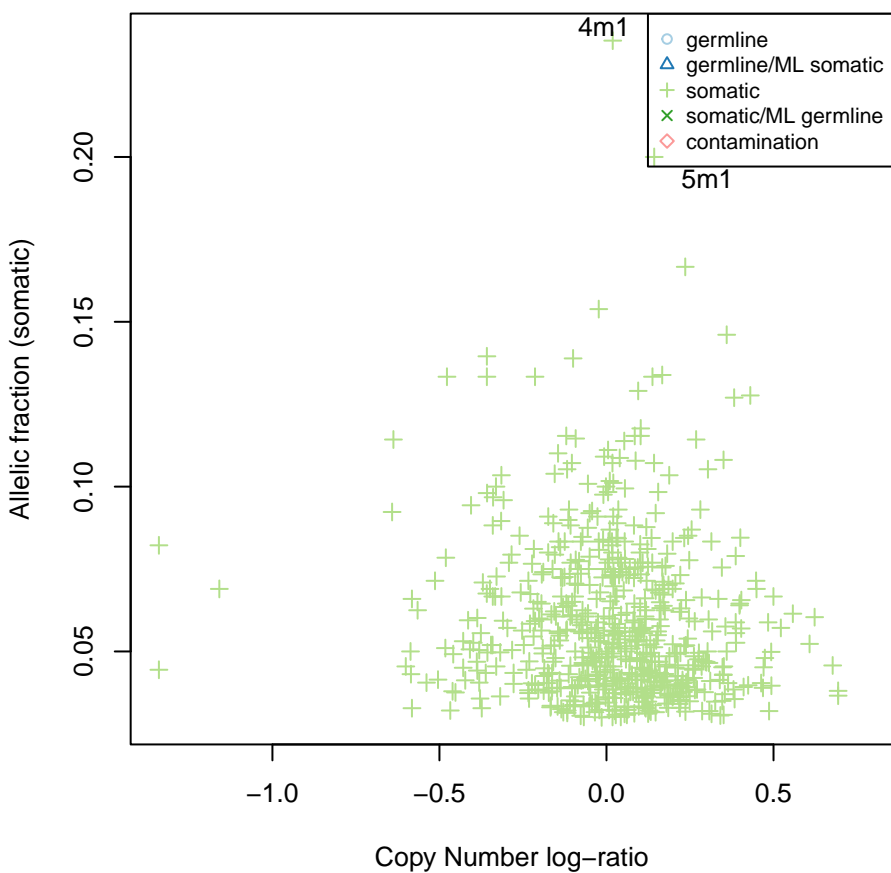
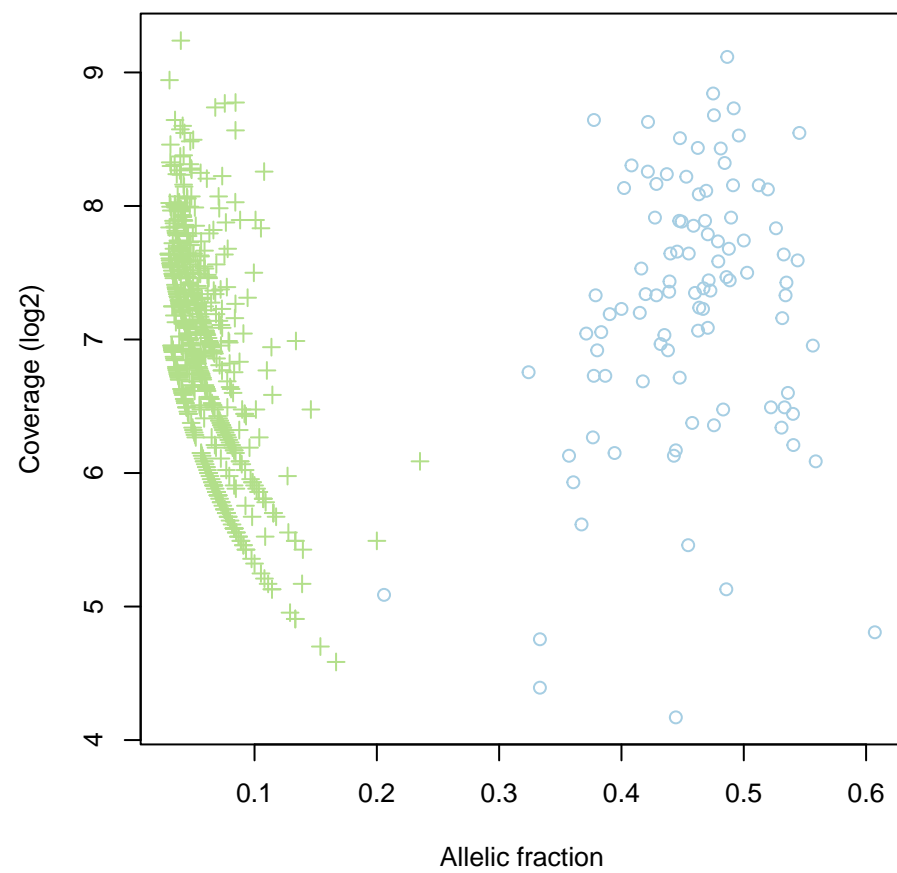
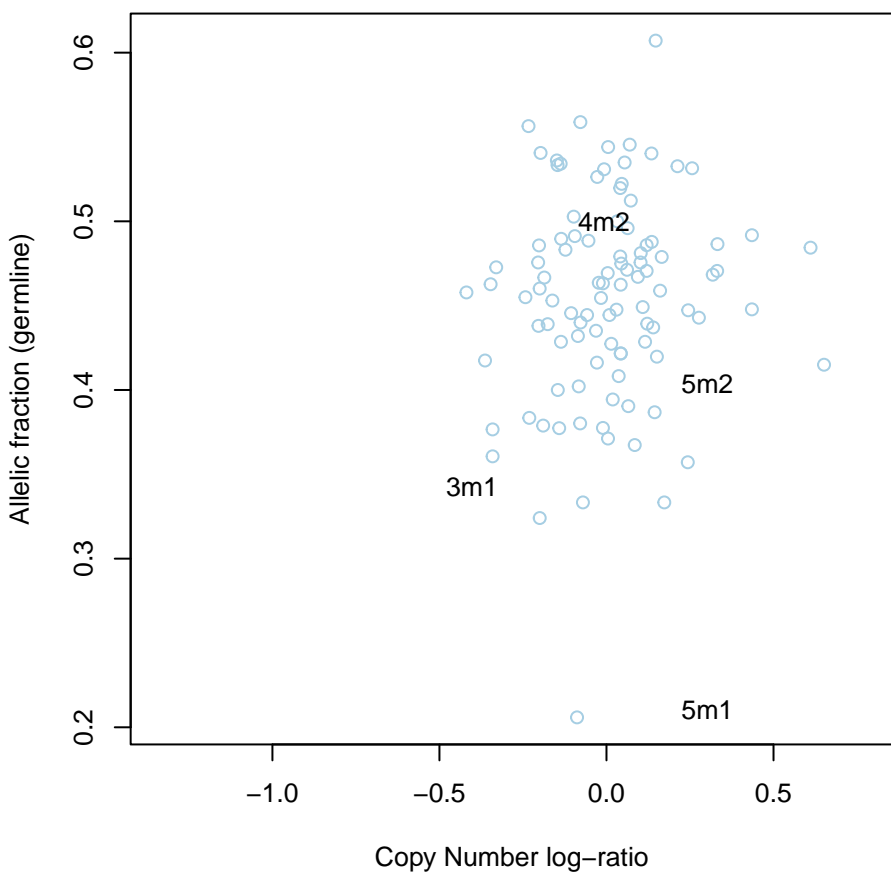


Purity: 0.92 Tumor ploidy: 4.022 SNV log-likelihood: -690.34 GoF: 26.3% Mean coverage: 363;148

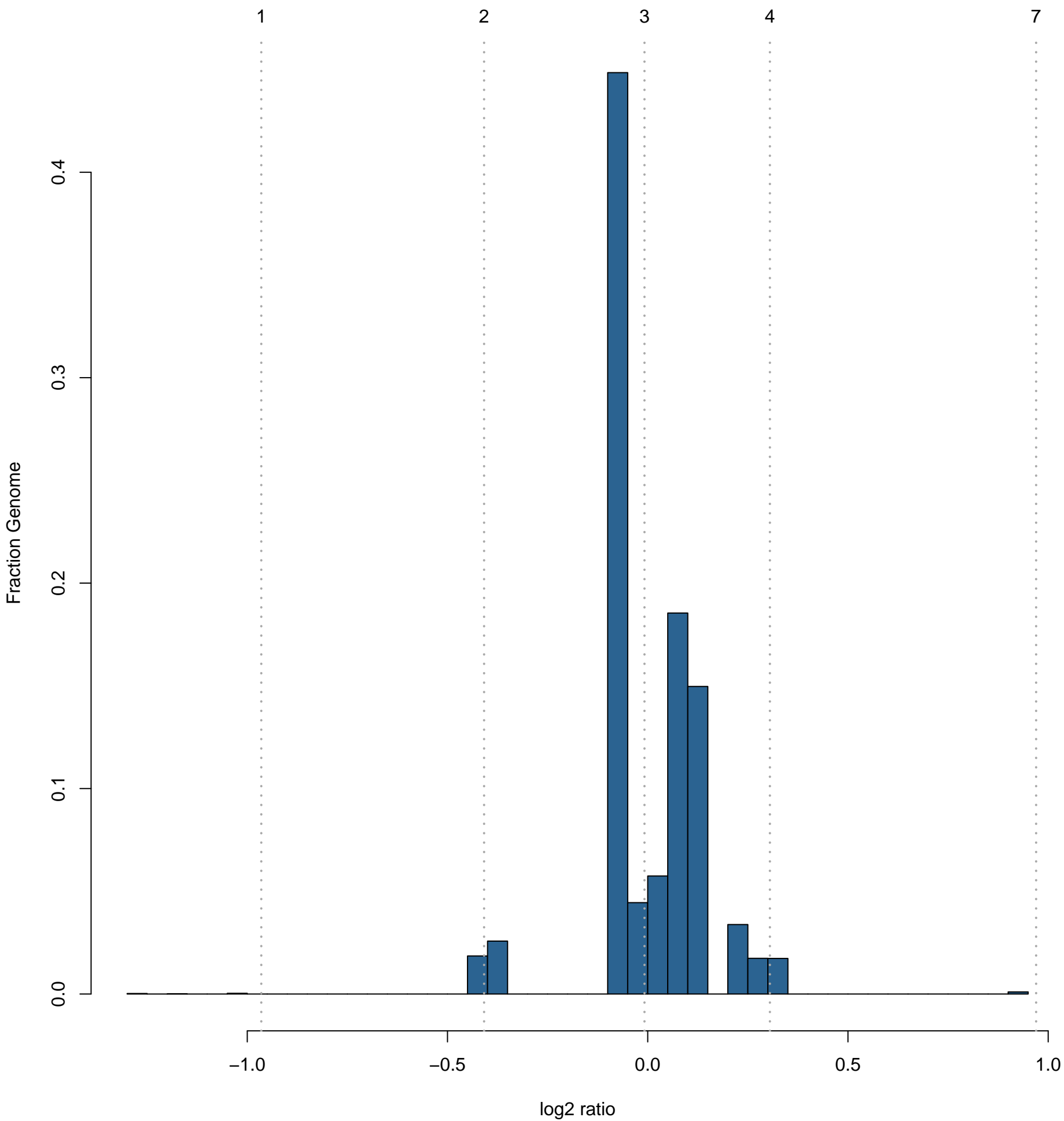


SCNA-fit log-likelihood: -19152.37

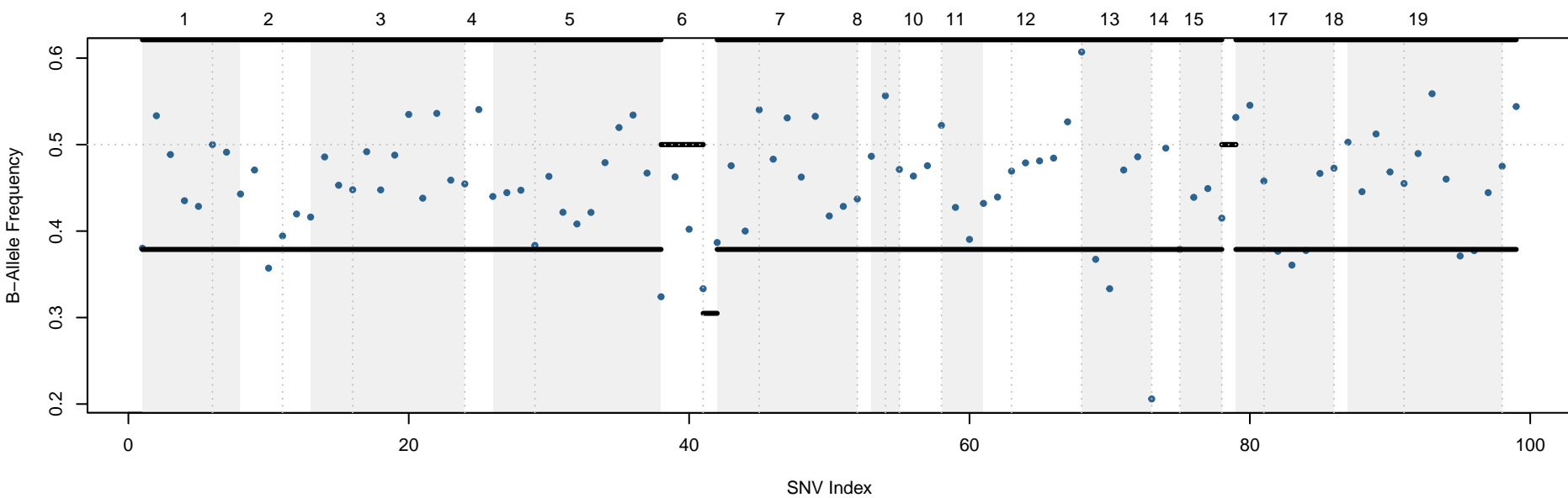




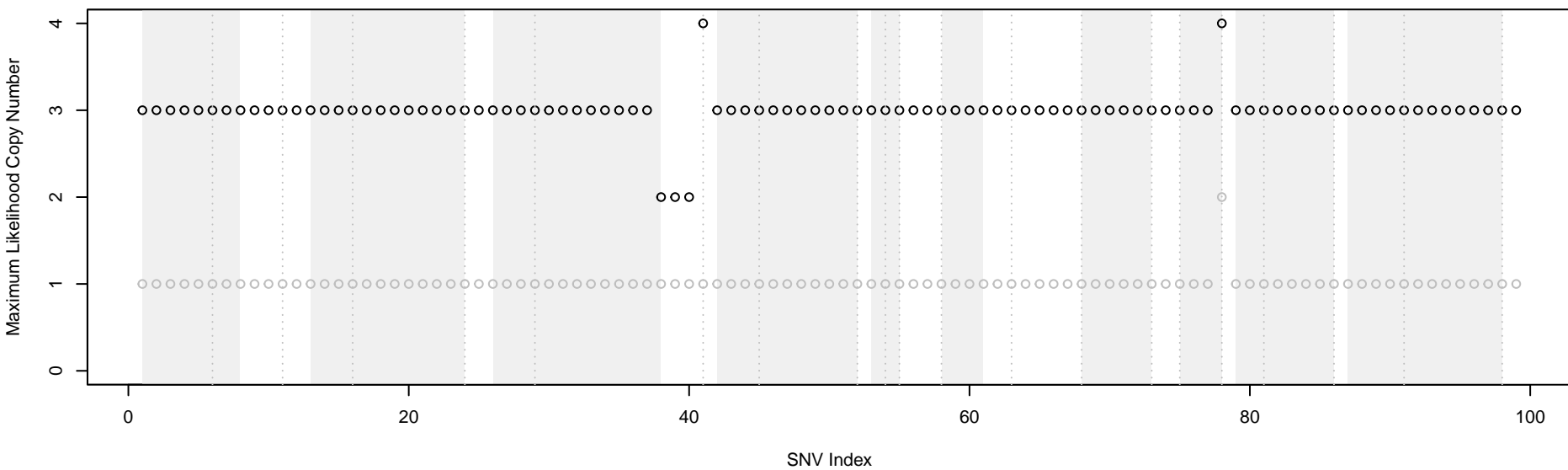
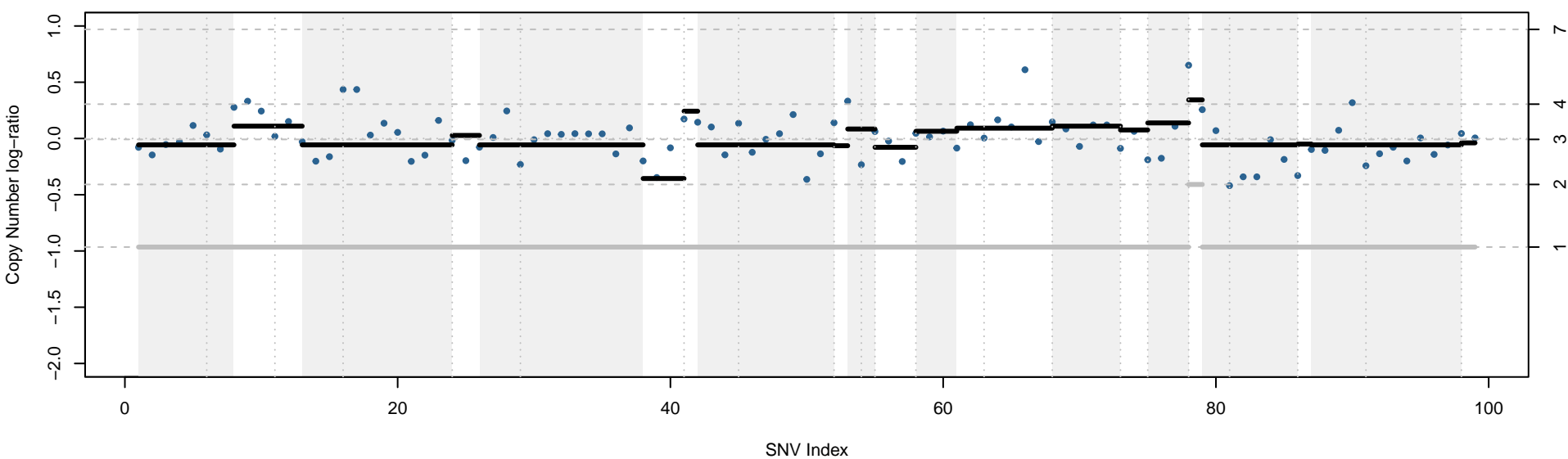
Purity: 0.64 Tumor ploidy: 3.023



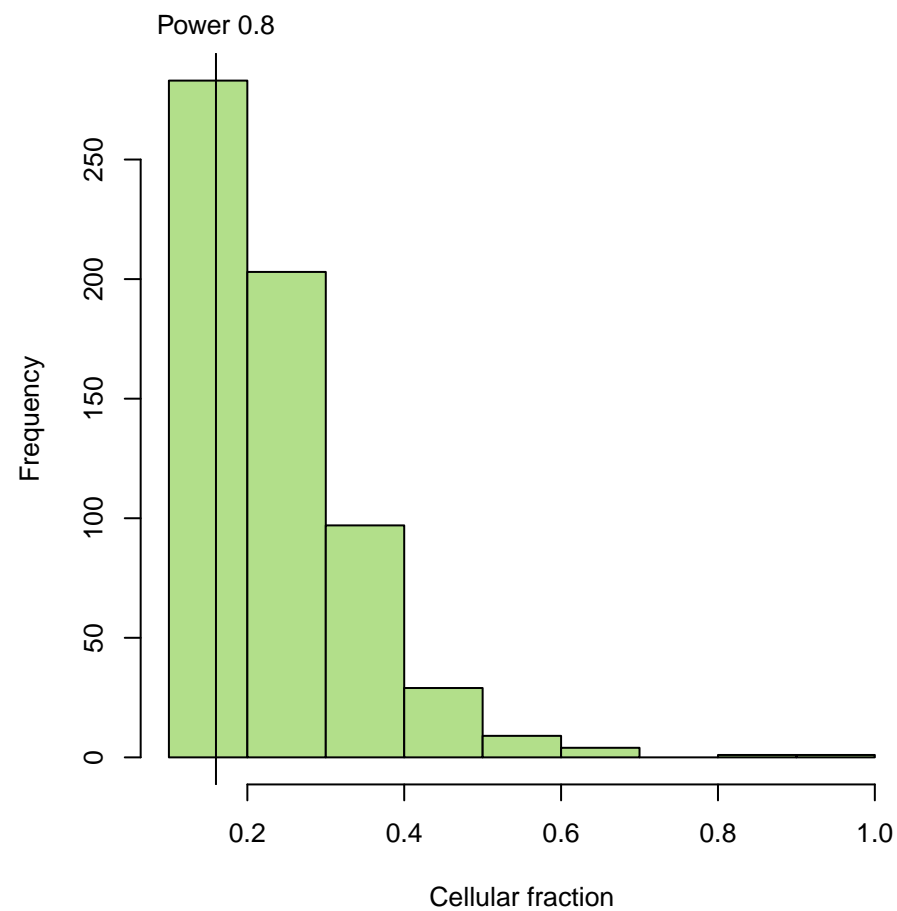
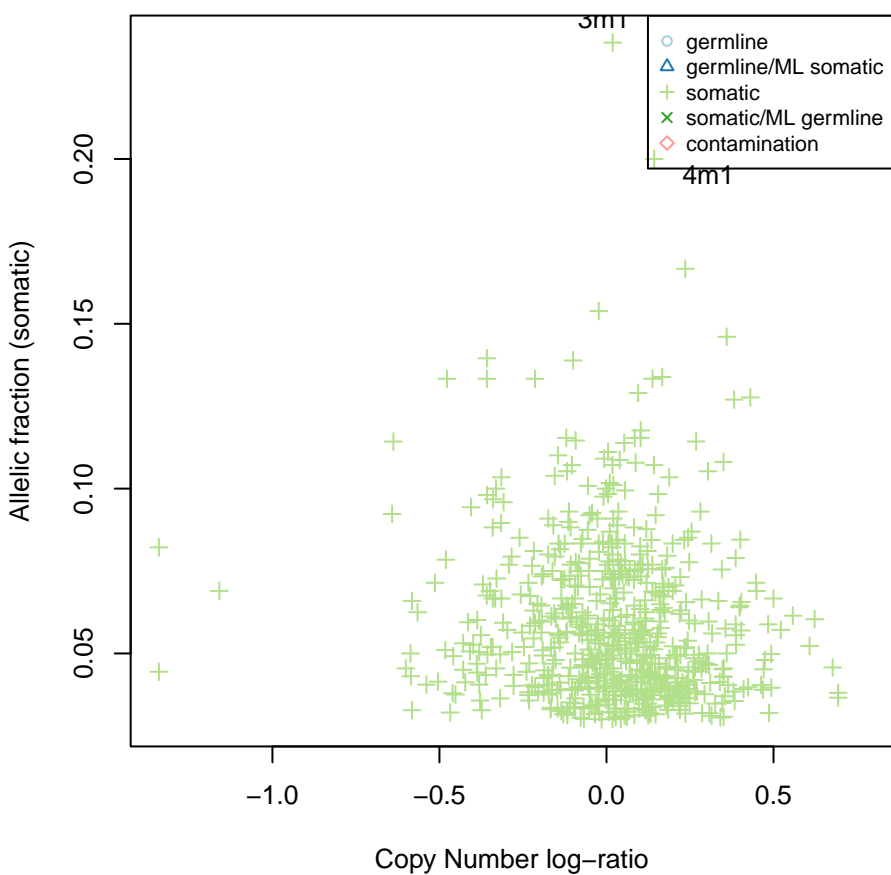
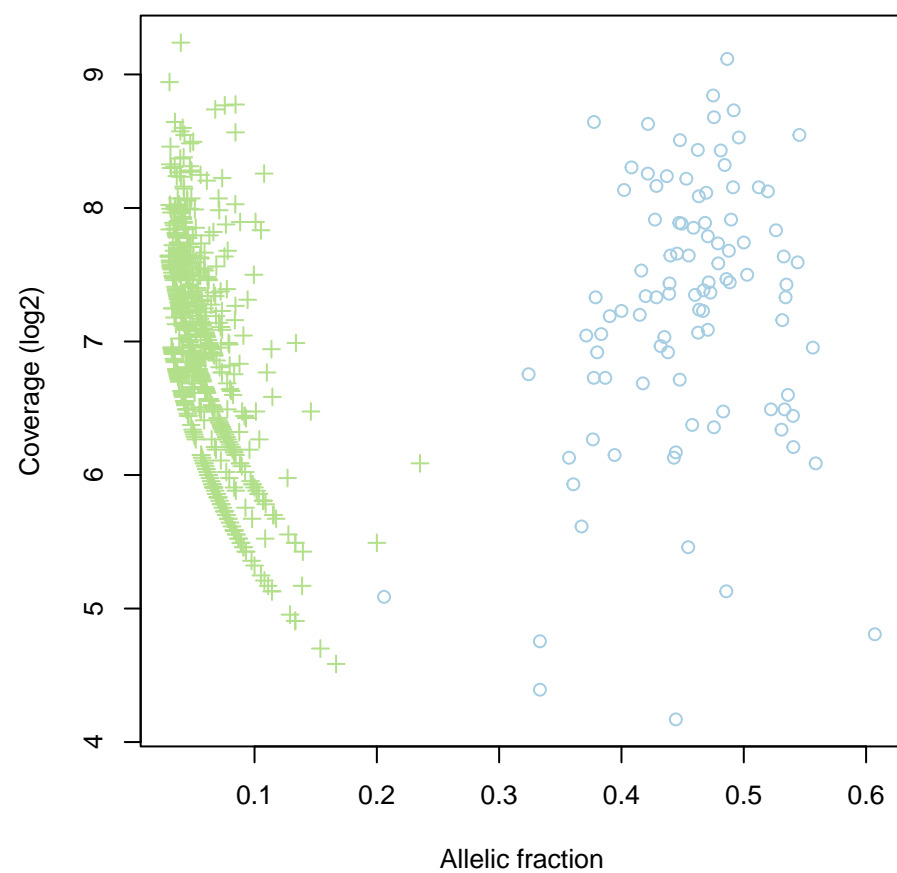
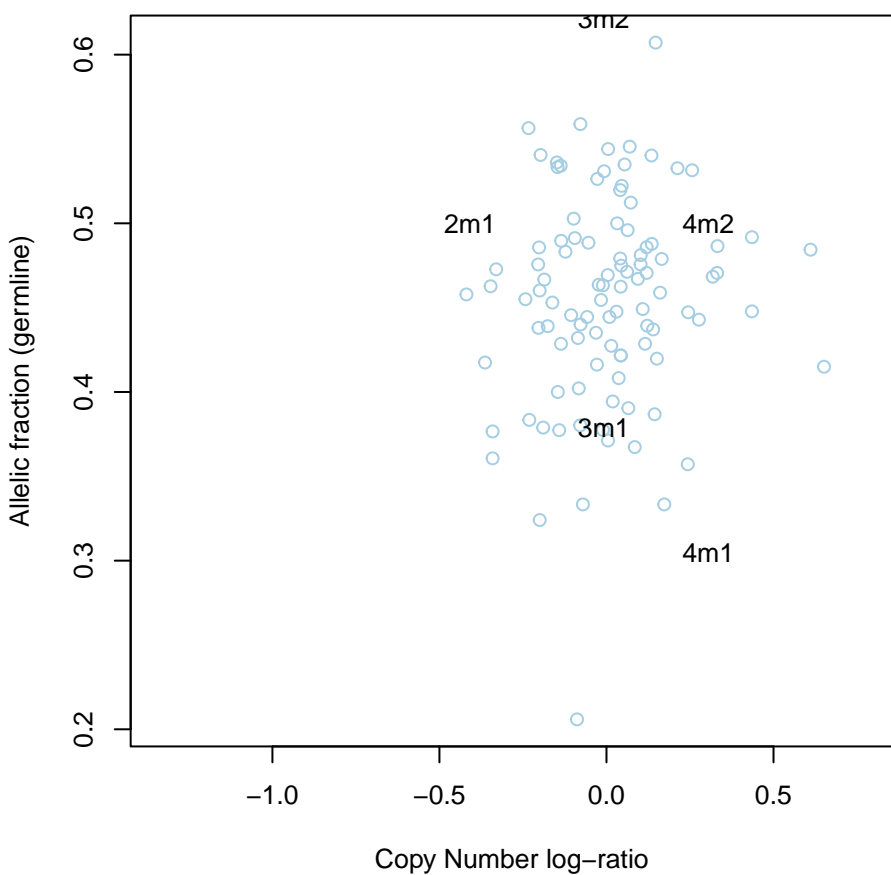
Purity: 0.64 Tumor ploidy: 3.023 SNV log-likelihood: -755.65 GoF: 23.3% Mean coverage: 363;148



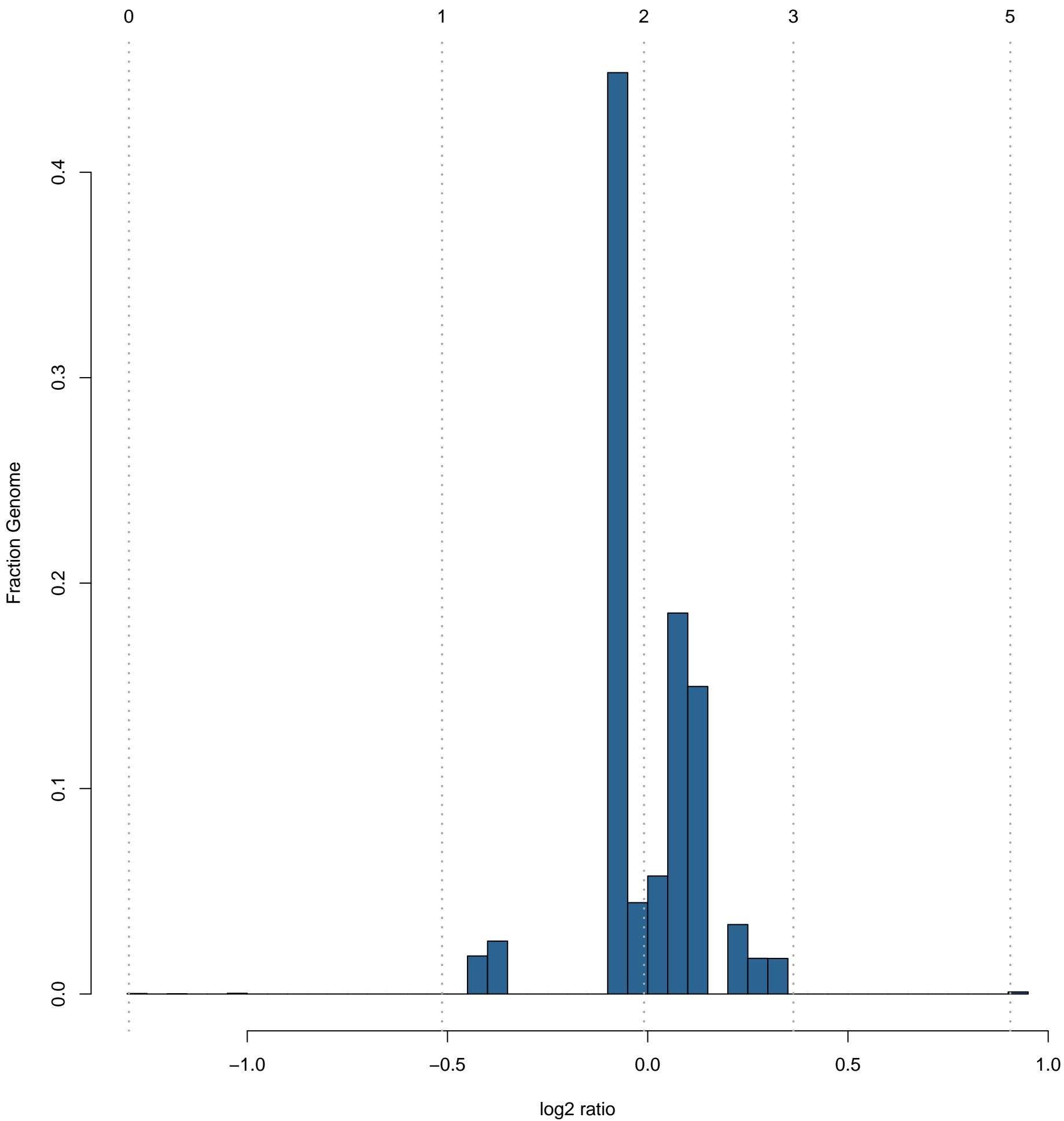
SCNA-fit log-likelihood: -19152.58



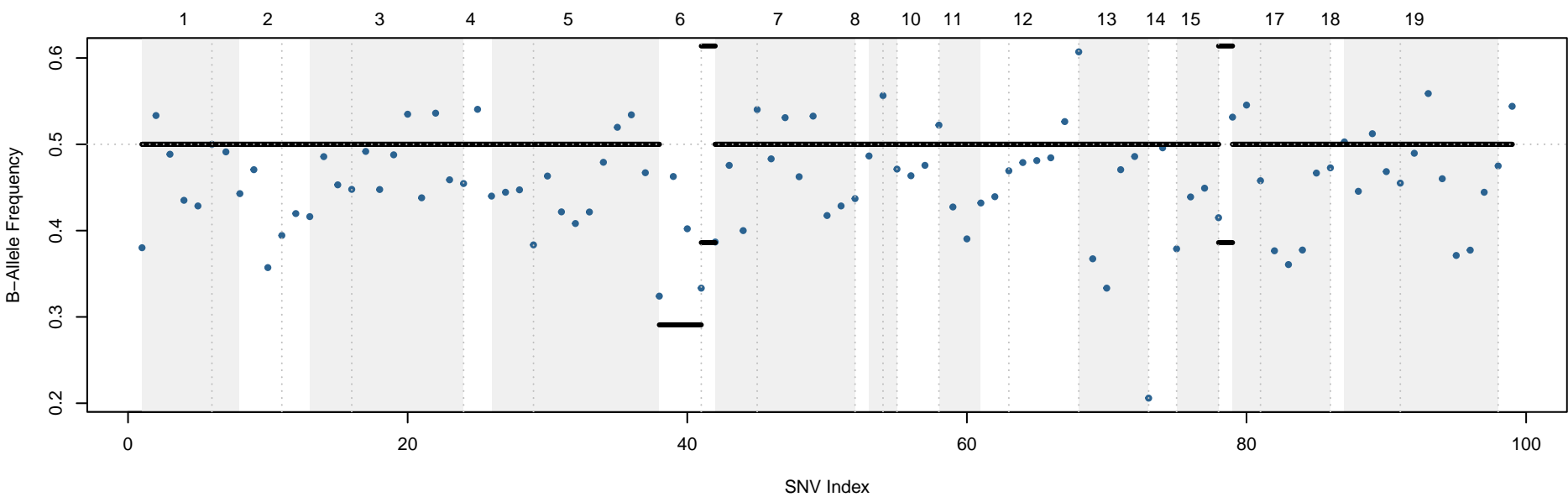




Purity: 0.59 Tumor ploidy: 2.022



Purity: 0.59 Tumor ploidy: 2.022 SNV log-likelihood: -831.06 GoF: 0% Mean coverage: 363;148



SCNA-fit log-likelihood: -19182.96

