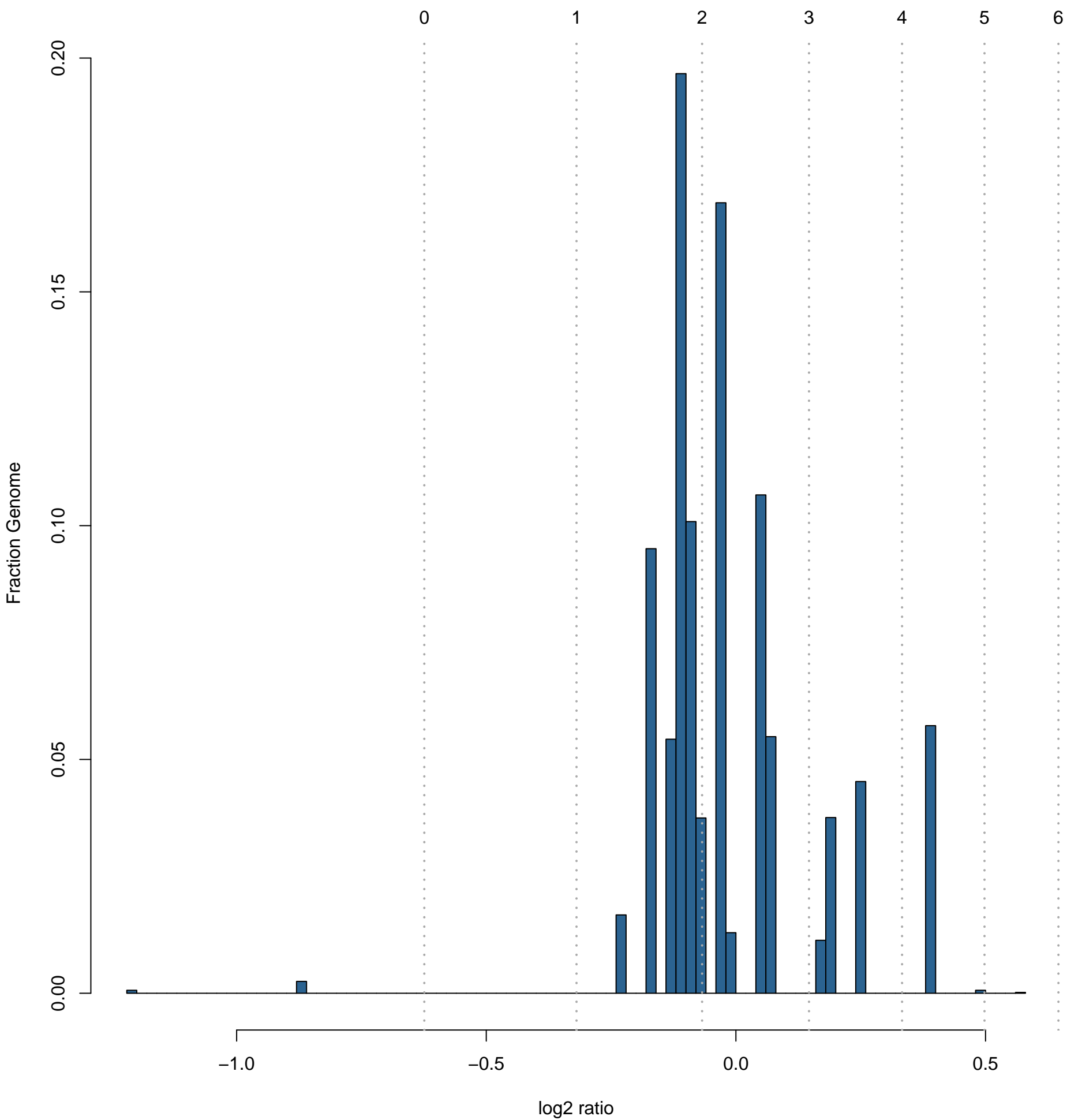
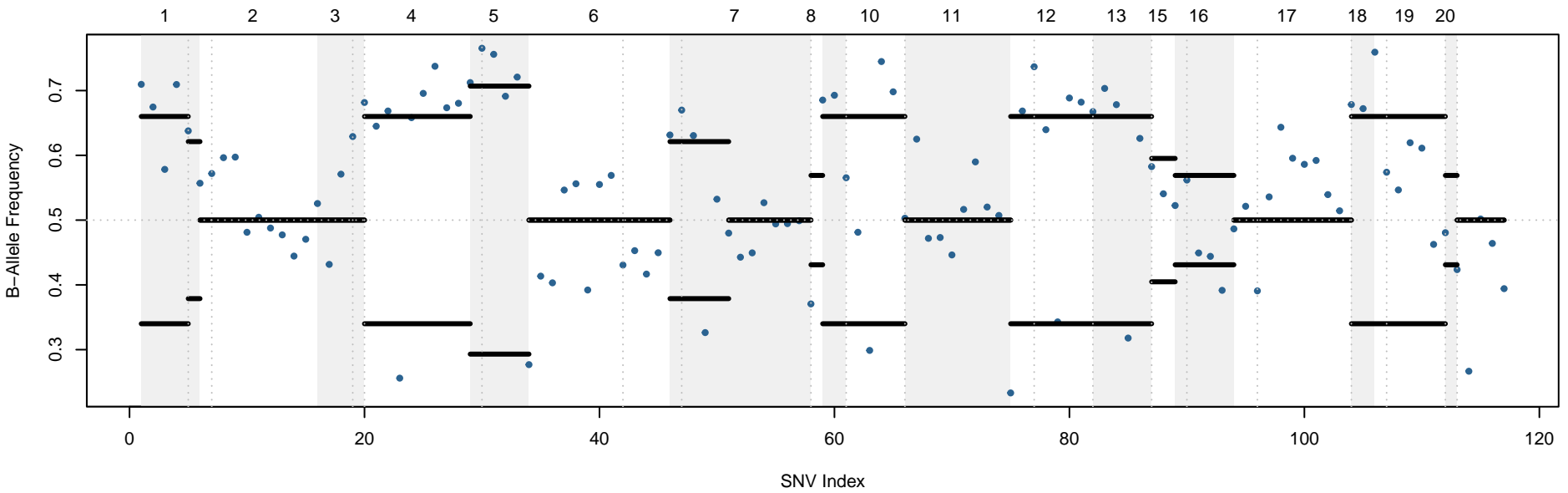


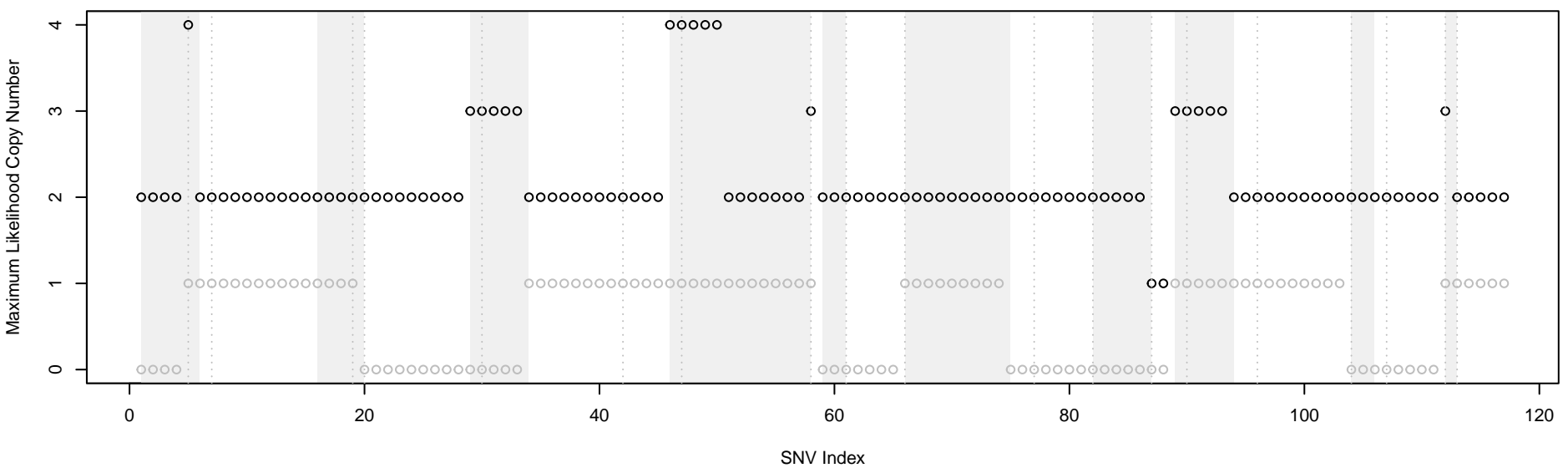
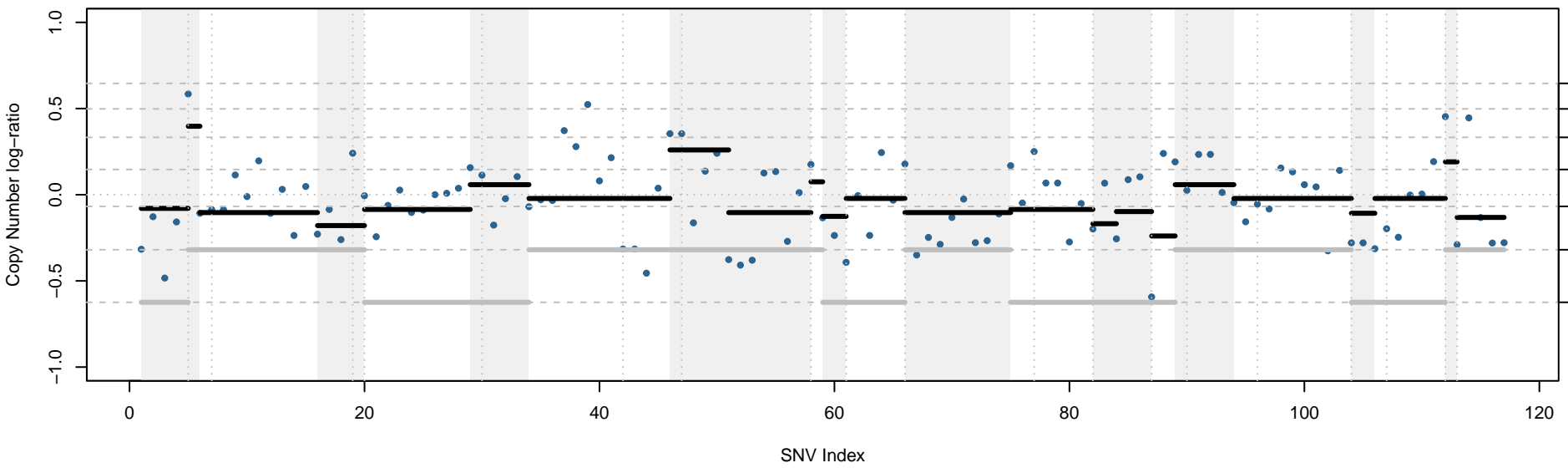
Purity: 0.32 Tumor ploidy: 2.3

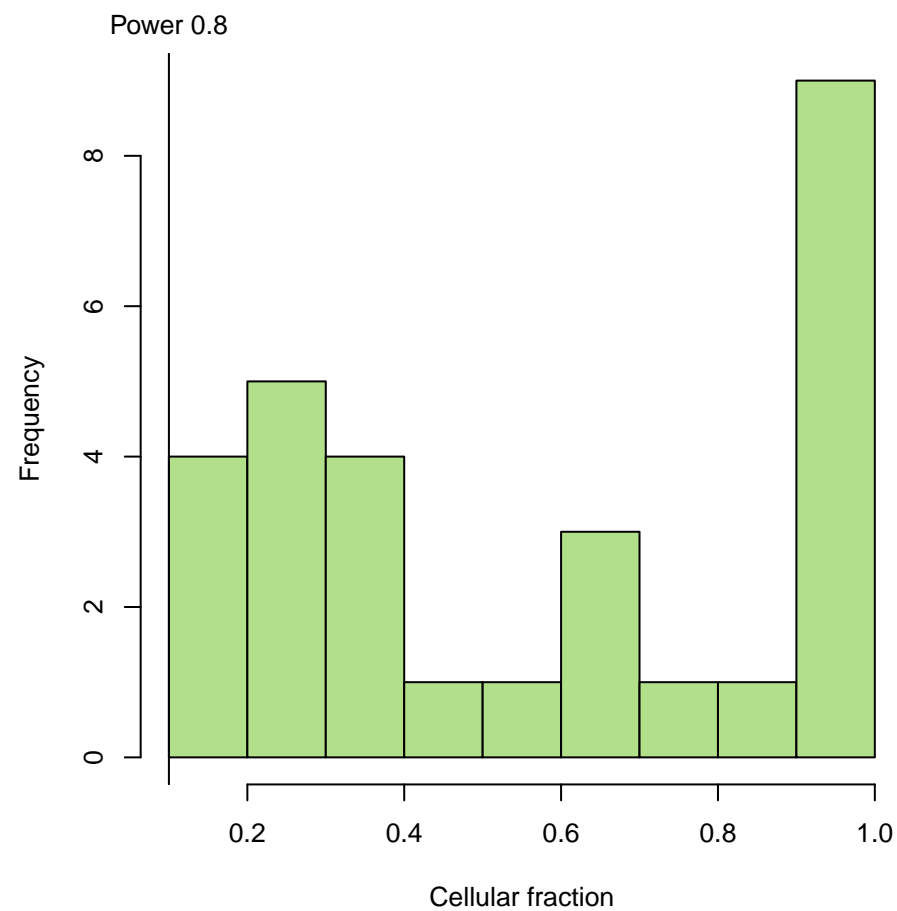
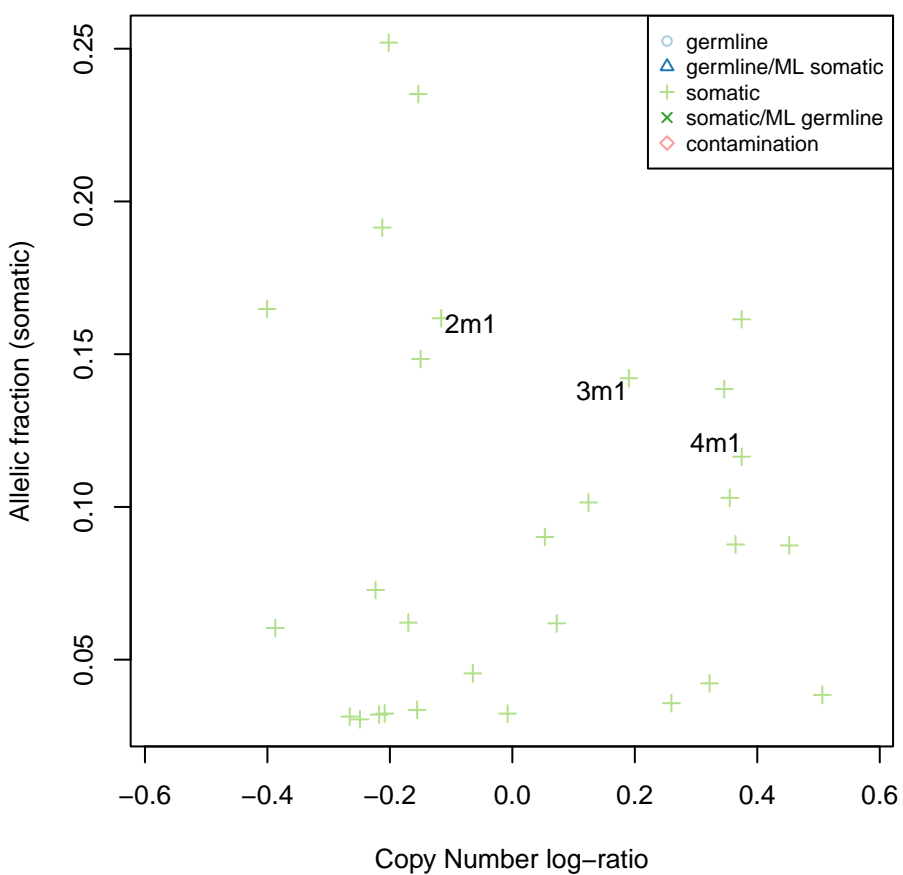
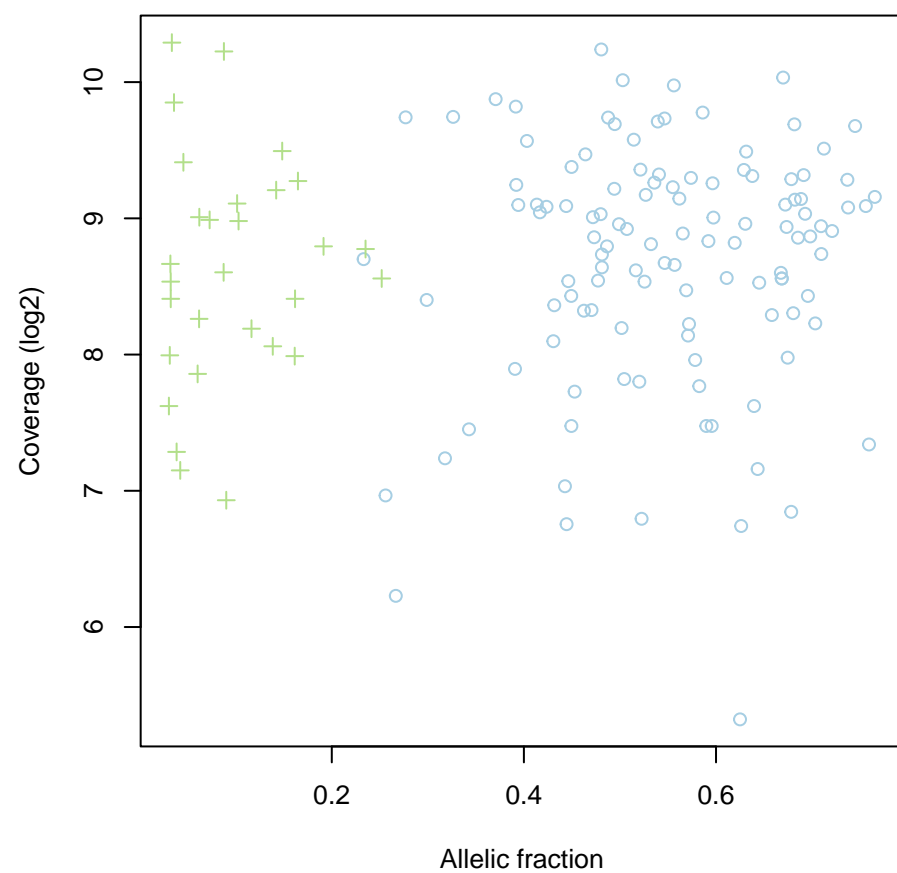
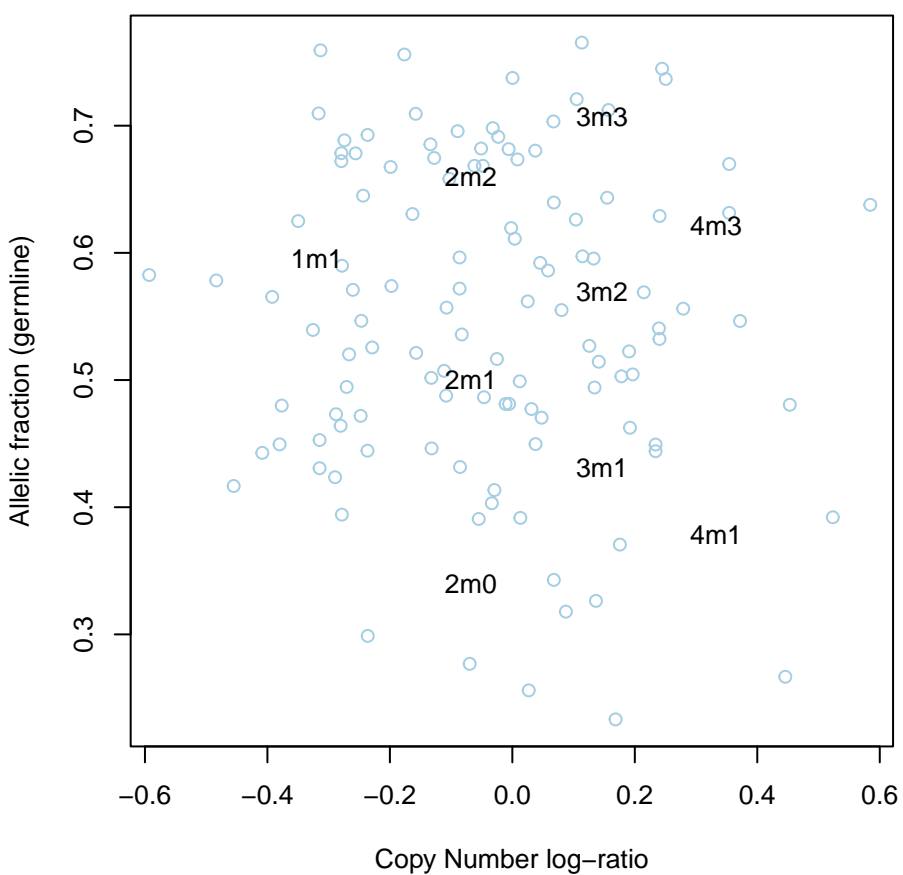


Purity: 0.32 Tumor ploidy: 2.3 SNV log-likelihood: -124.43 GoF: 89.4% Mean coverage: 477;413

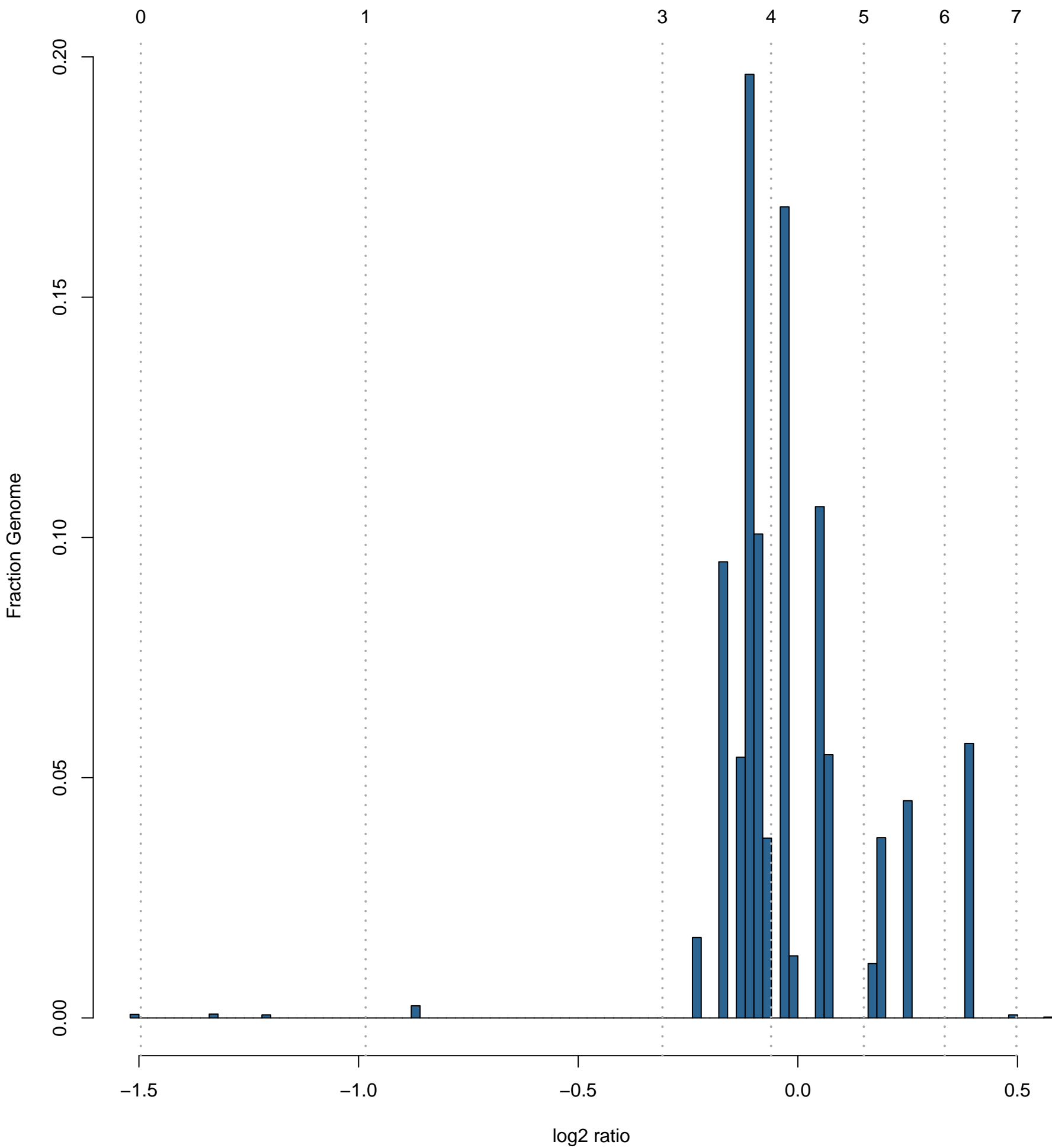


SCNA-fit log-likelihood: -6916.42

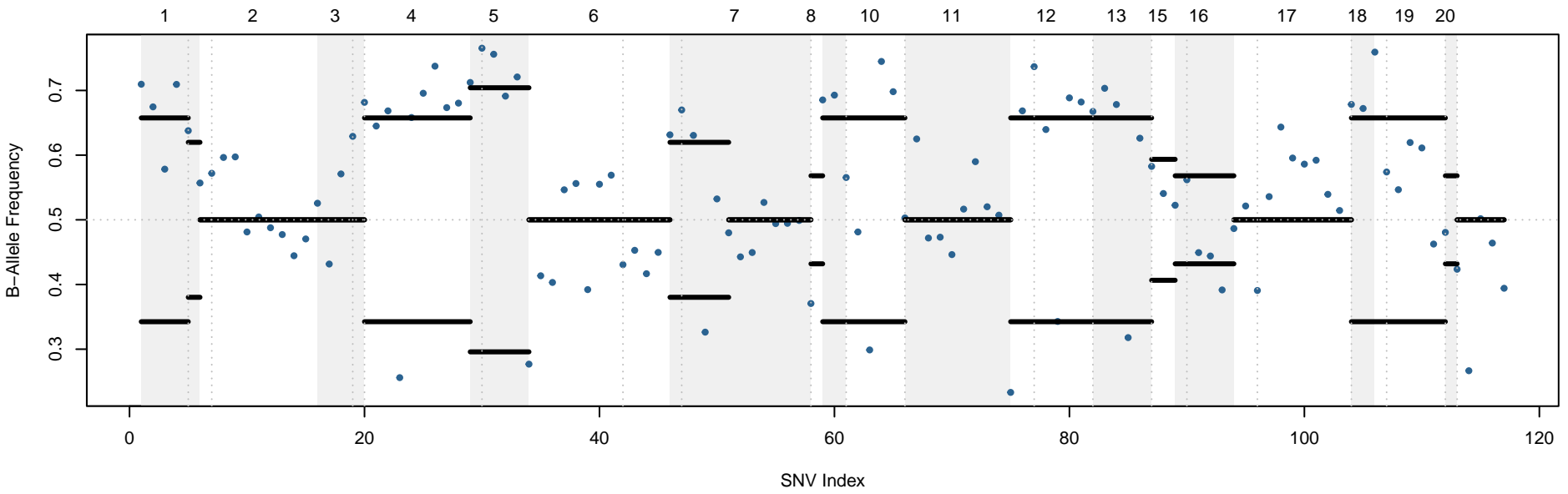




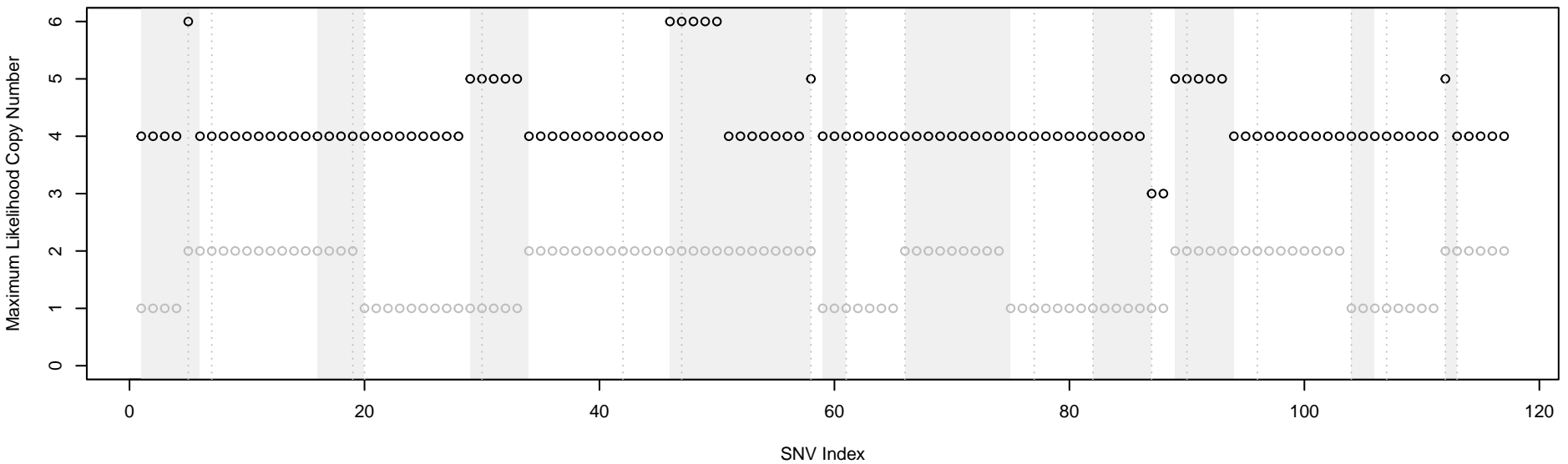
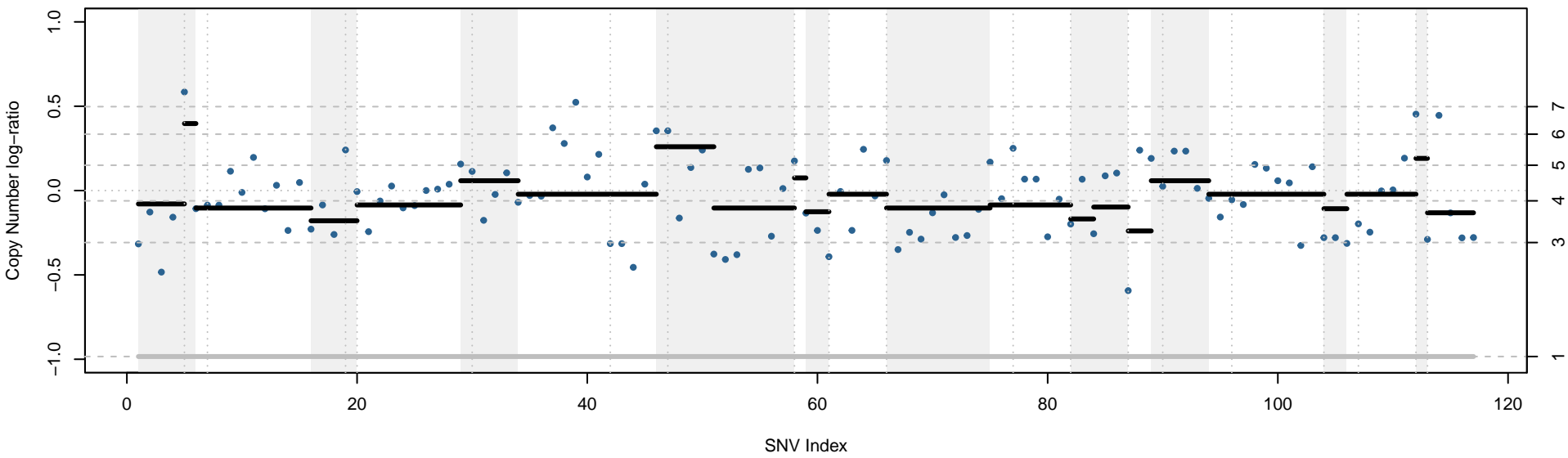
Purity: 0.46 Tumor ploidy: 4.274

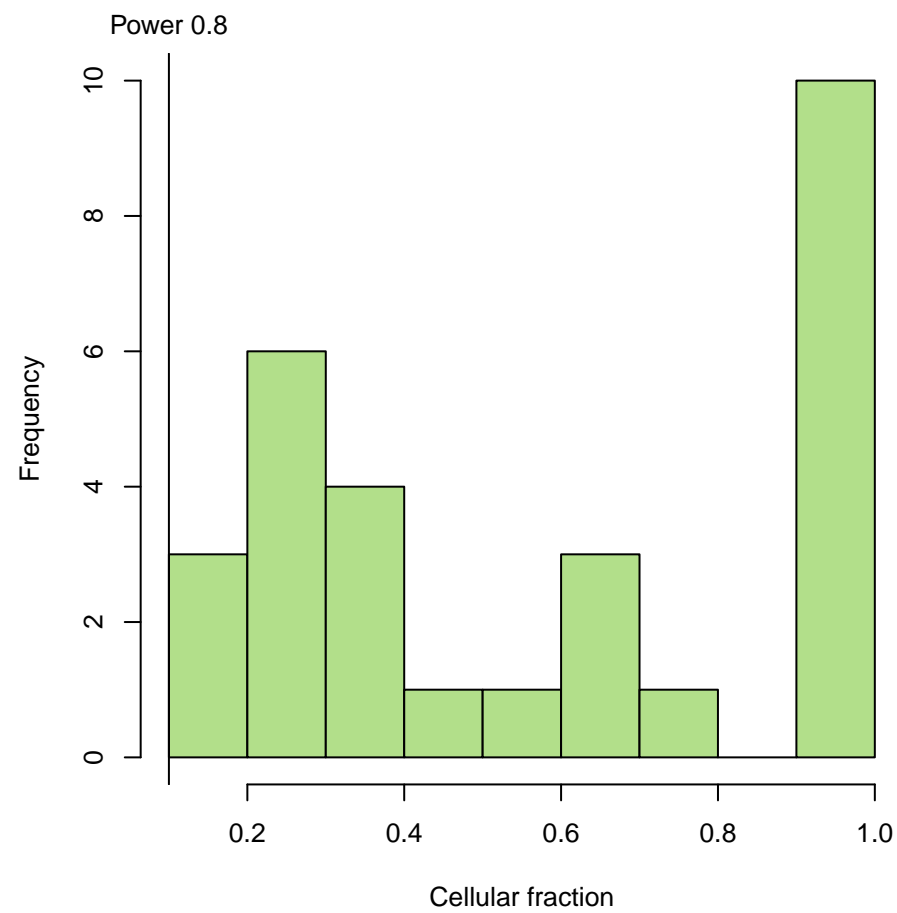
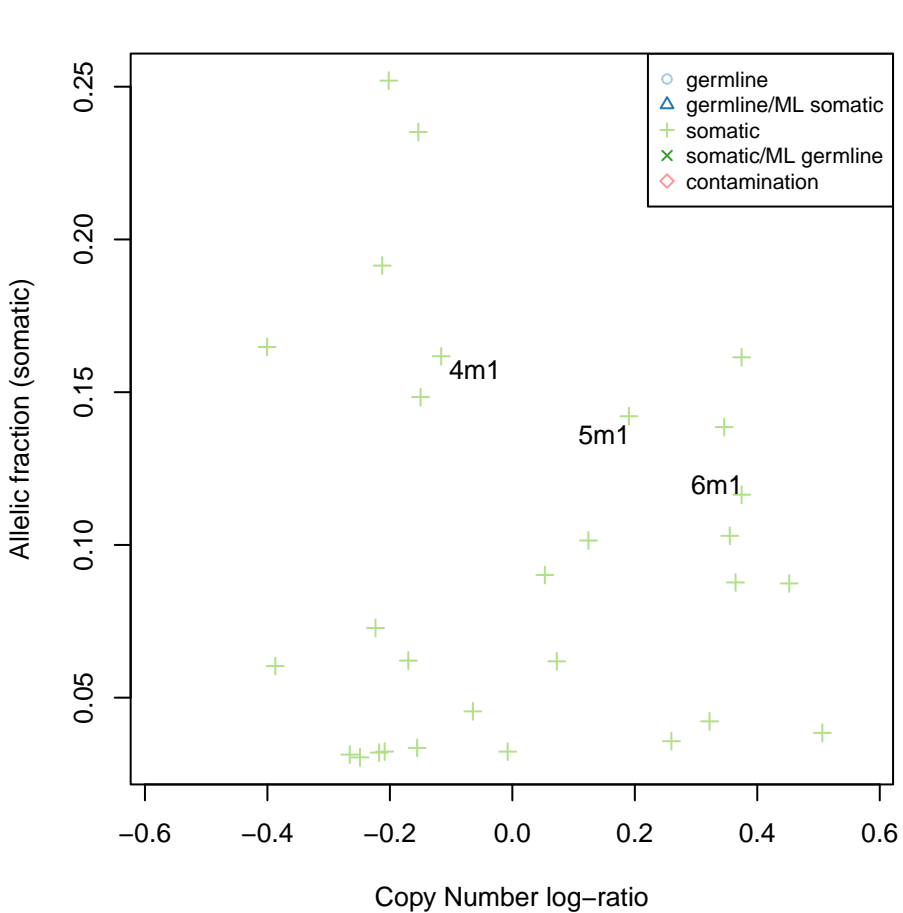
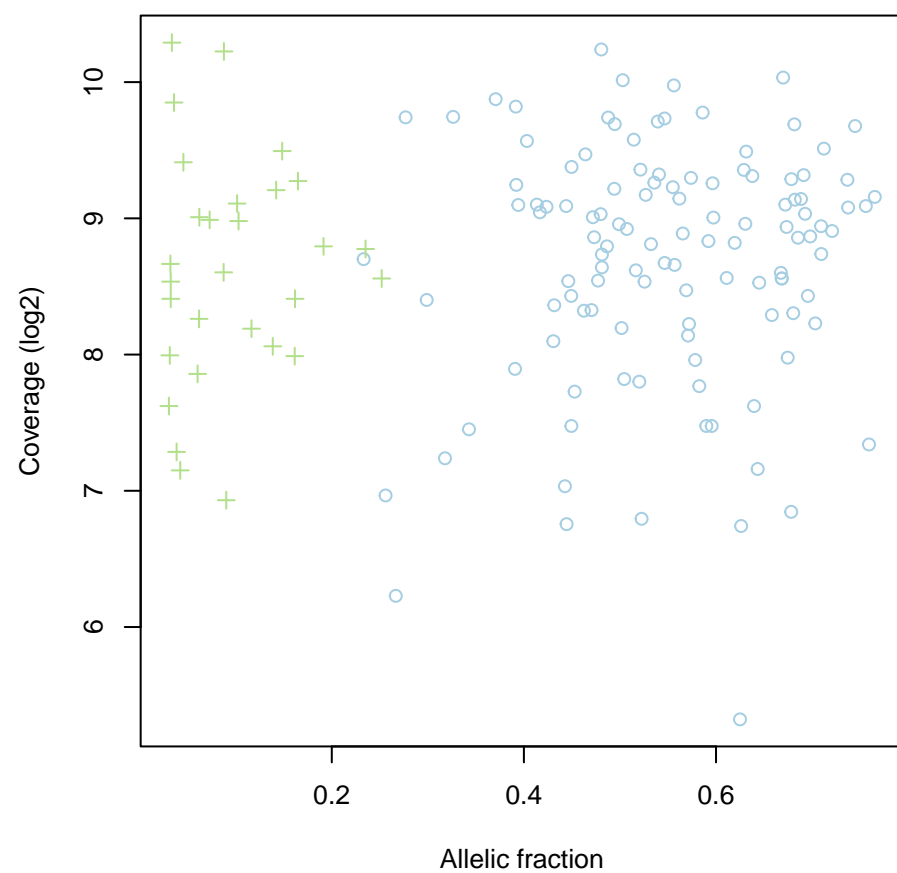
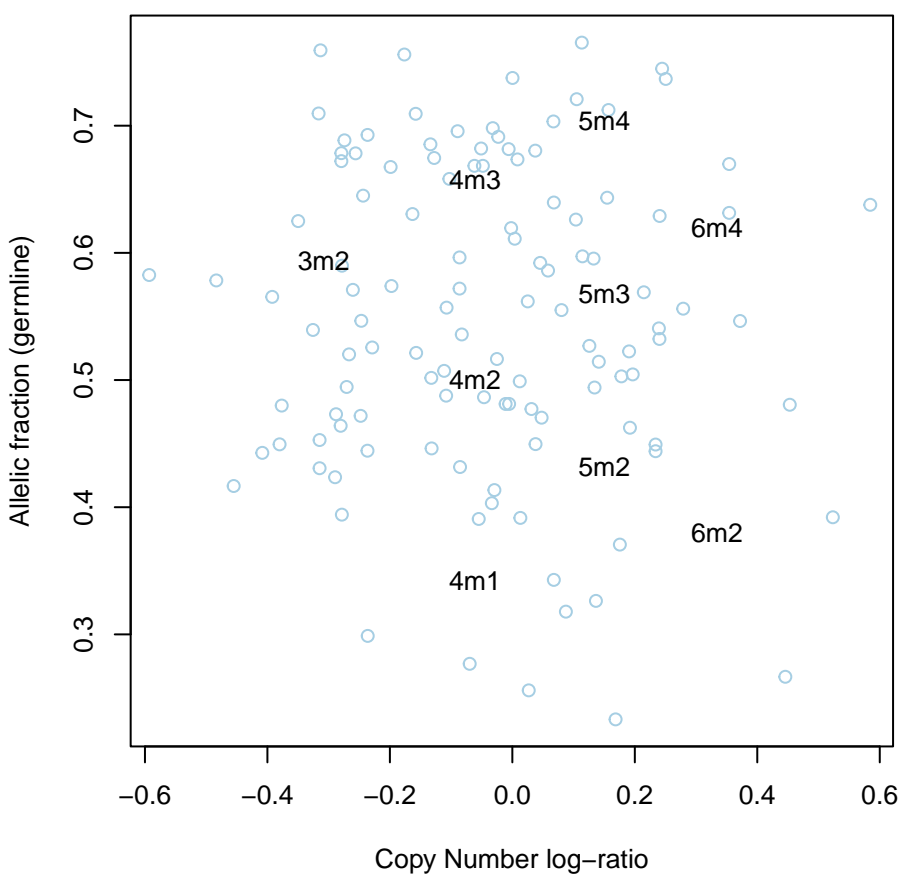


Purity: 0.46 Tumor ploidy: 4.274 SNV log-likelihood: -199.68 GoF: 89.4% Mean coverage: 477,413

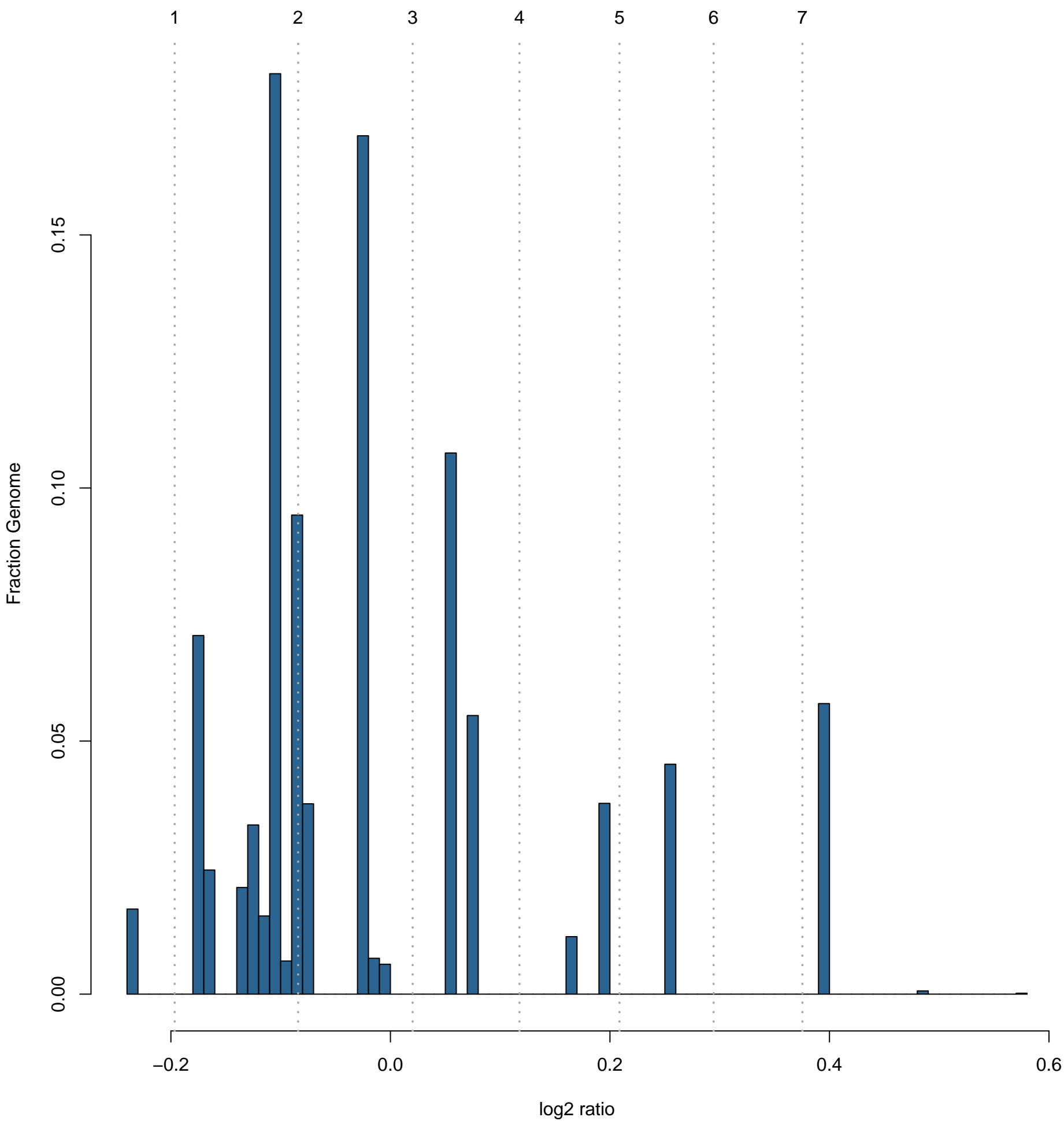


SCNA-fit log-likelihood: -6877.97

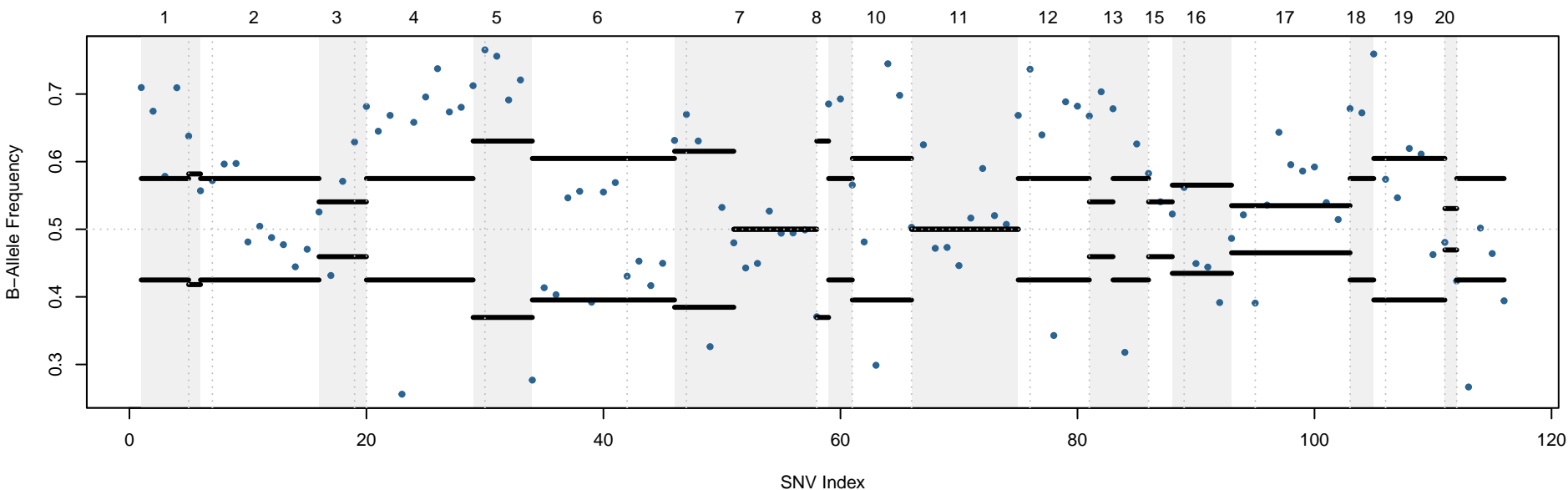




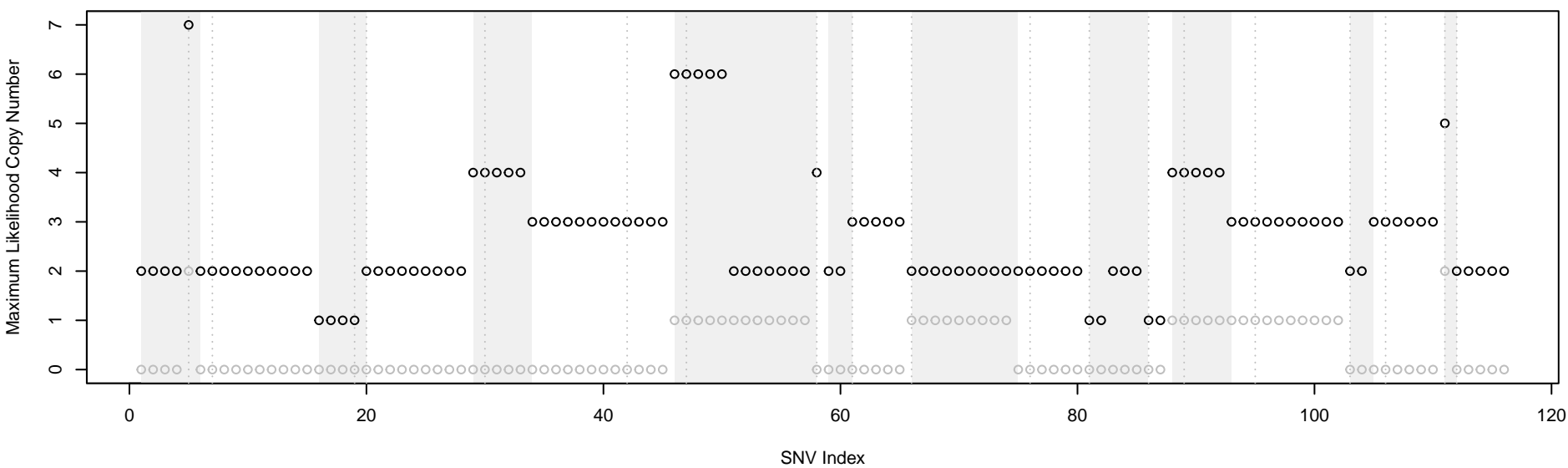
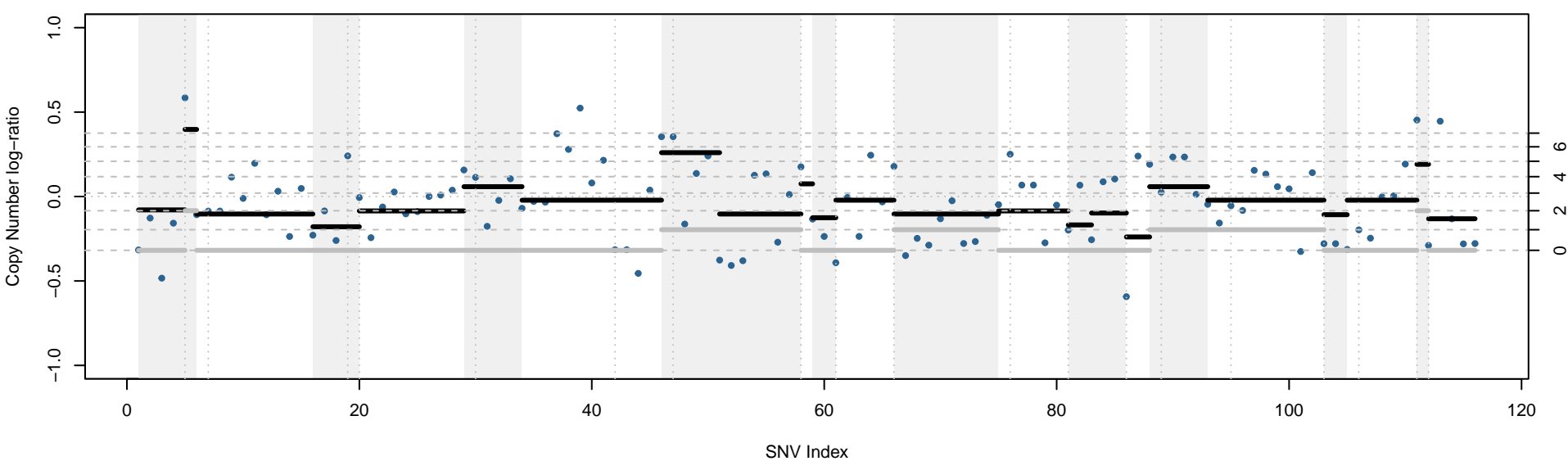
Purity: 0.15 Tumor ploidy: 2.8

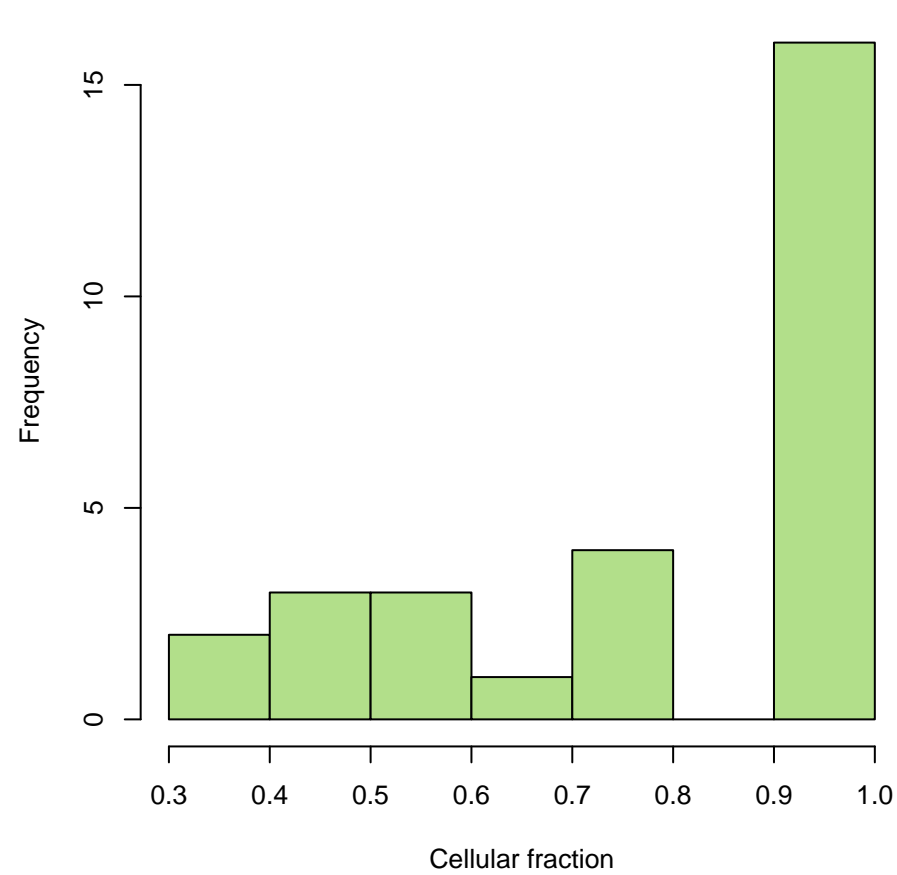
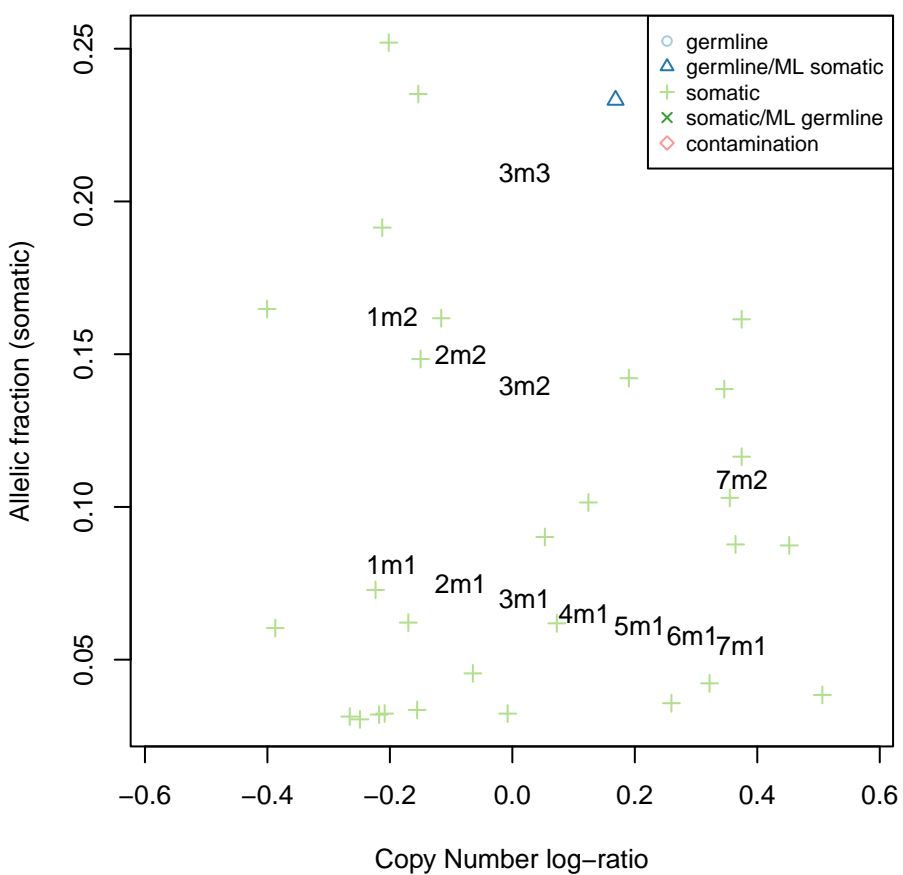
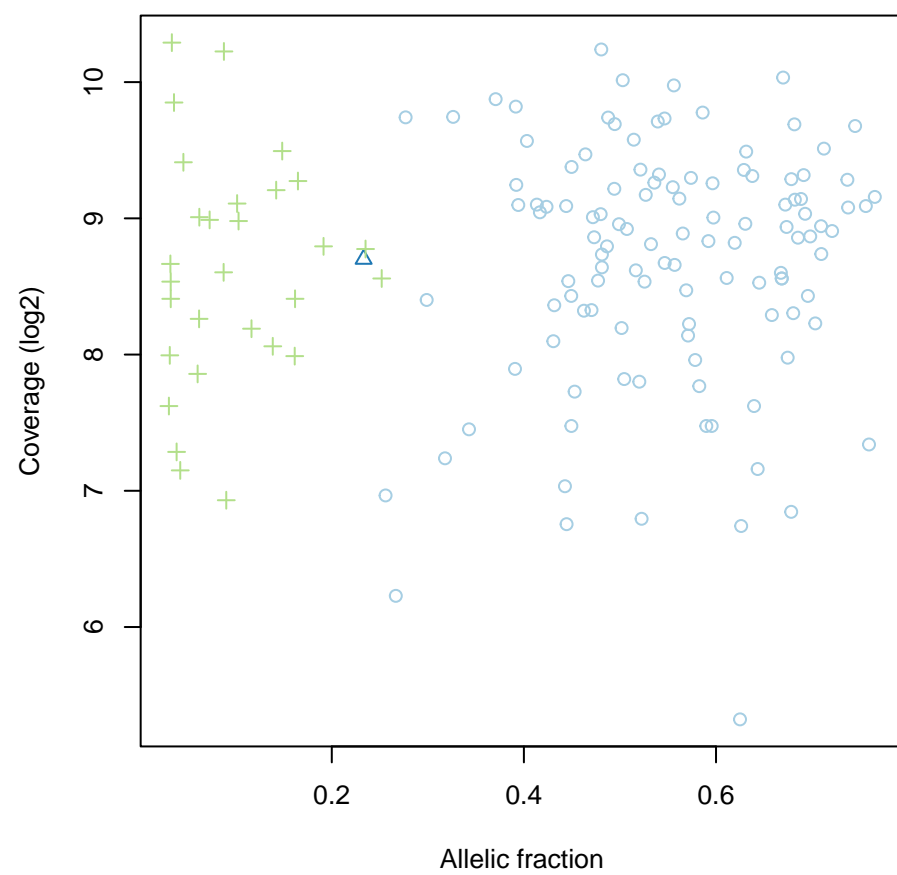
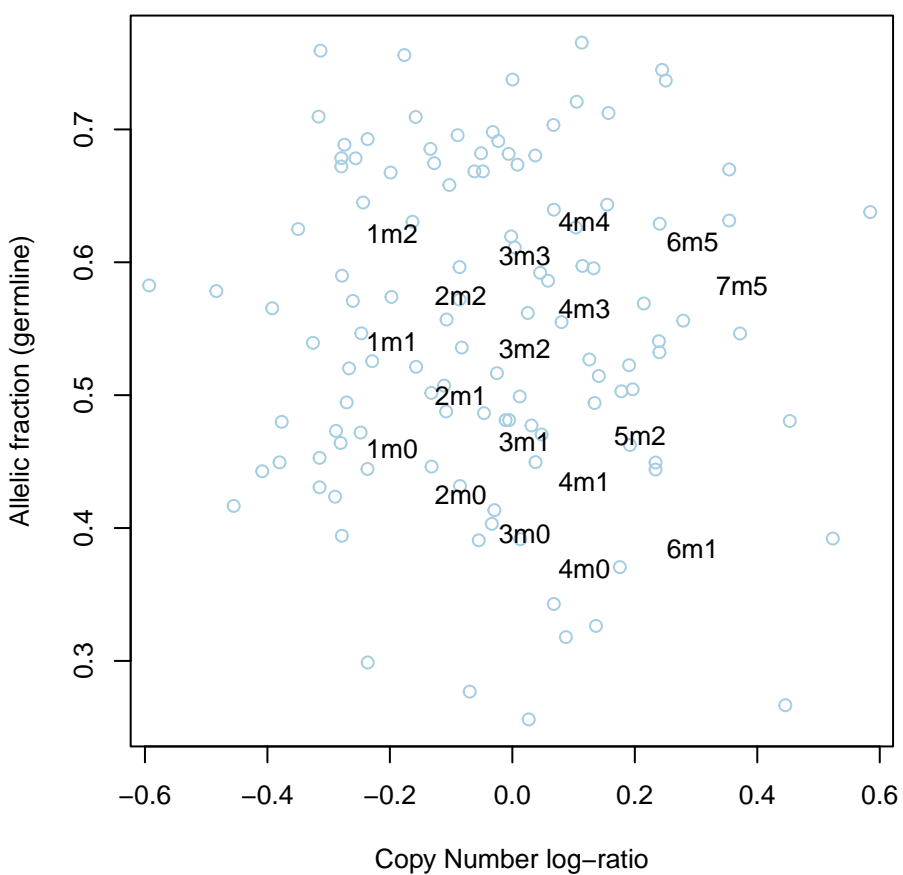


Purity: 0.15 Tumor ploidy: 2.8 SNV log-likelihood: -307.12 GoF: 88% Mean coverage: 477;413

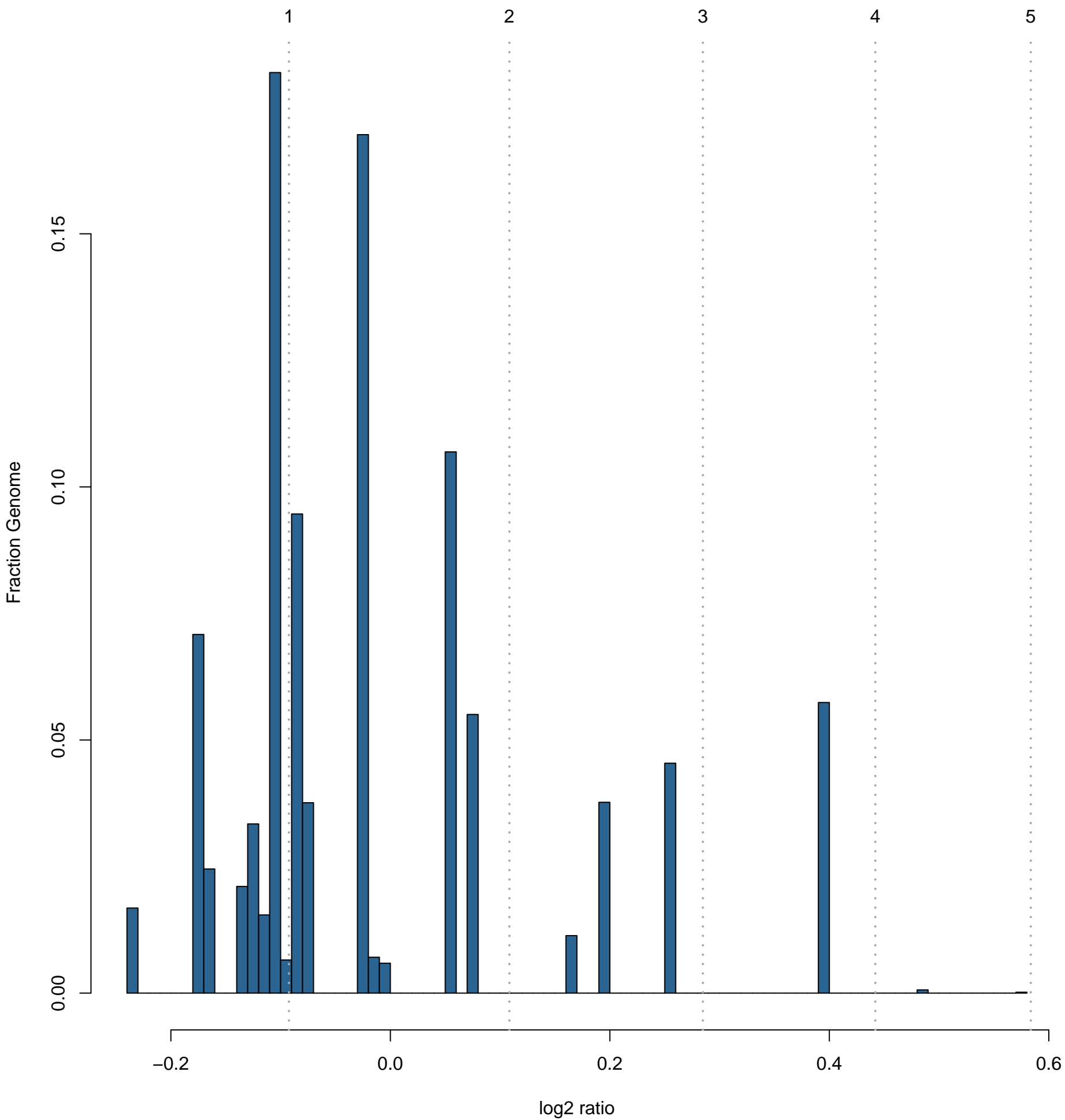


SCNA-fit log-likelihood: -6856.8

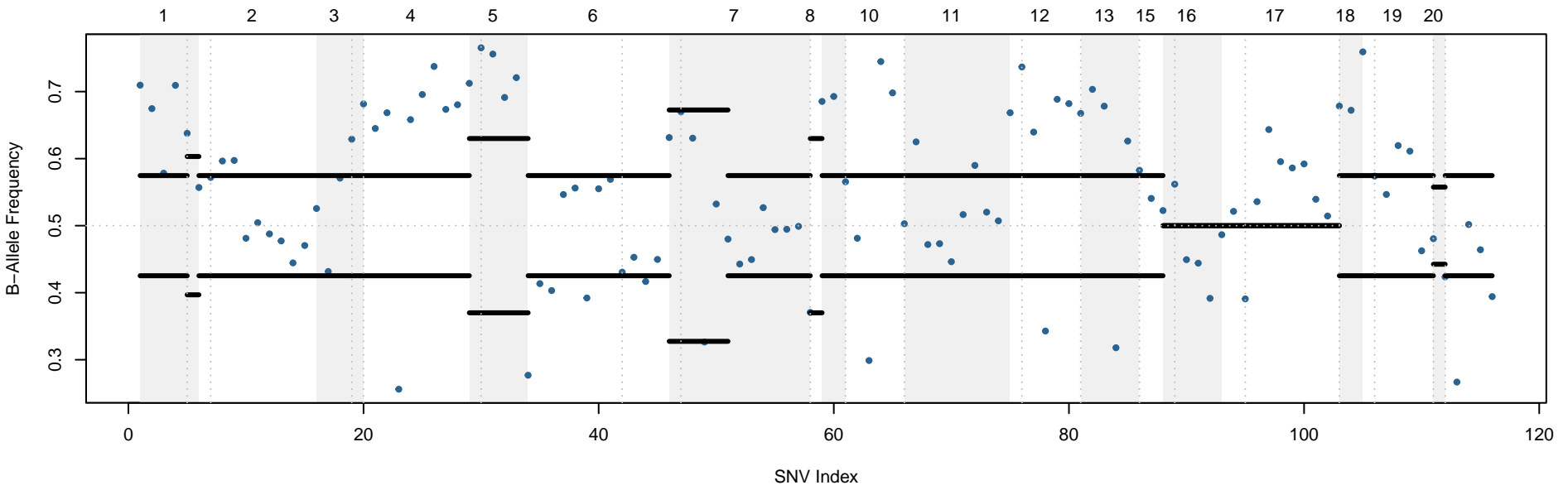




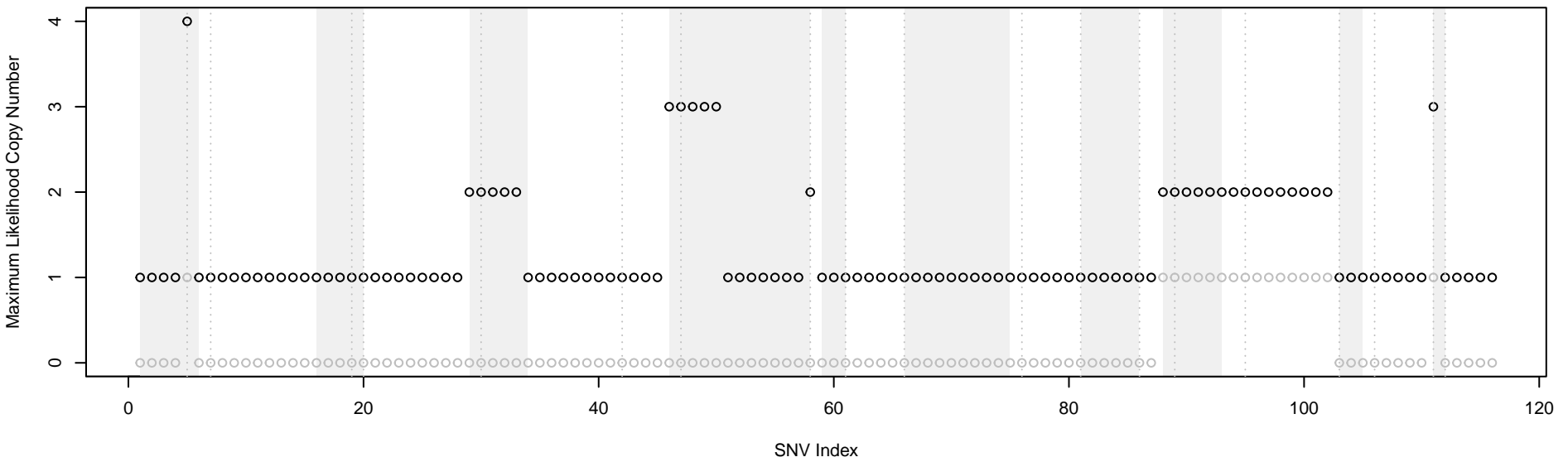
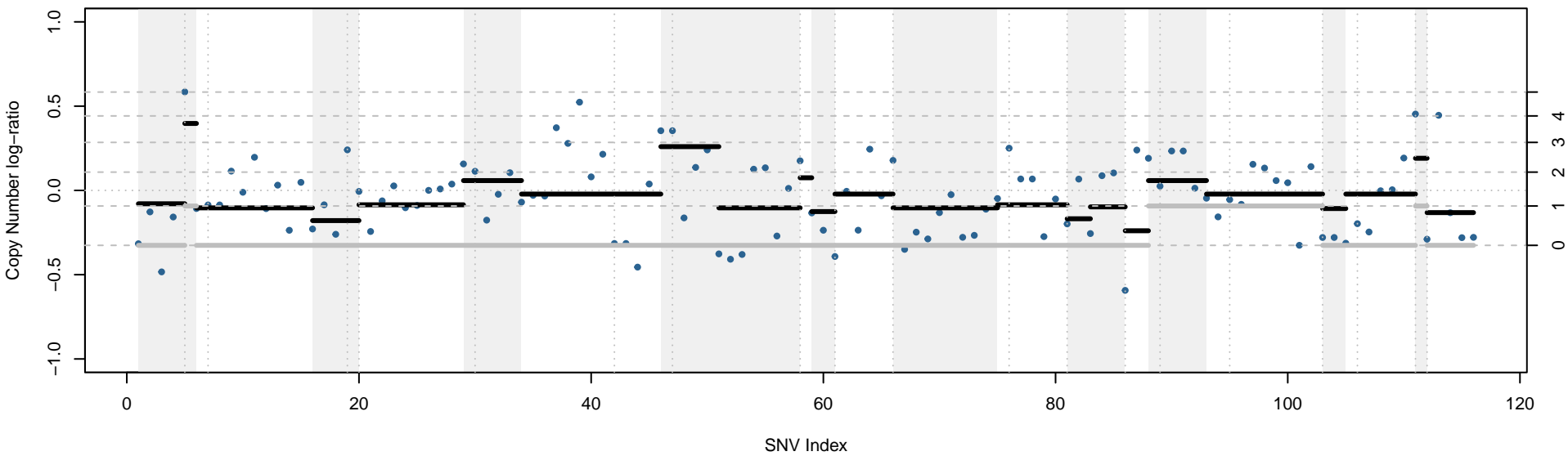
Purity: 0.26 Tumor ploidy: 1.443

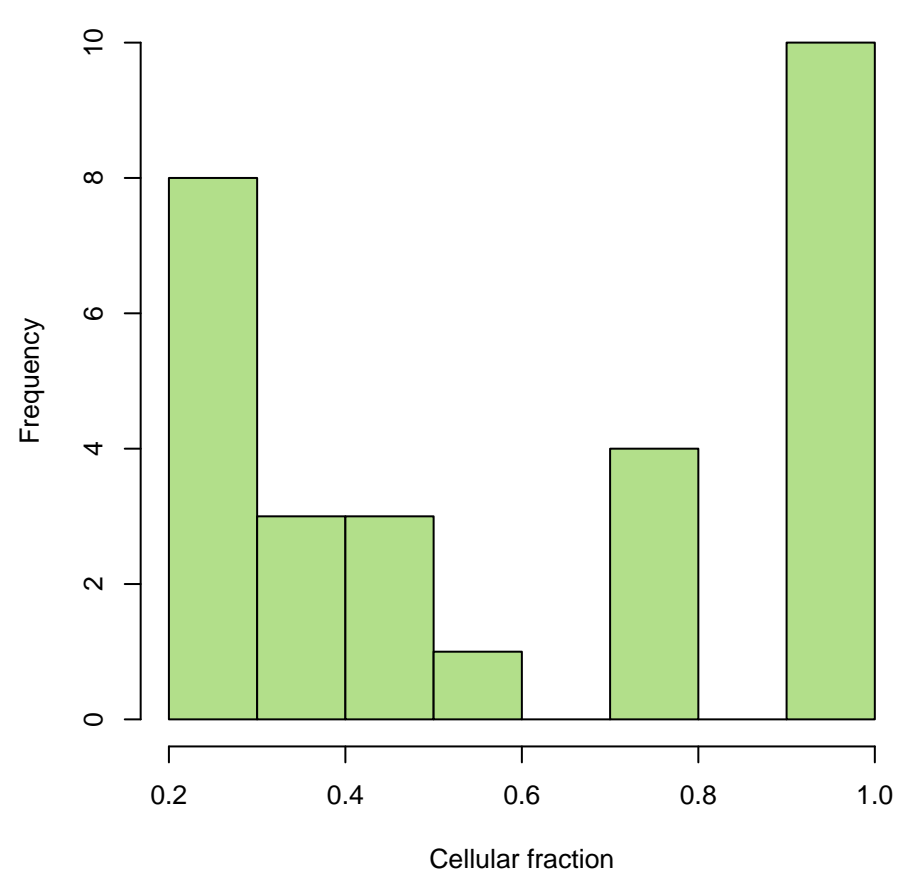
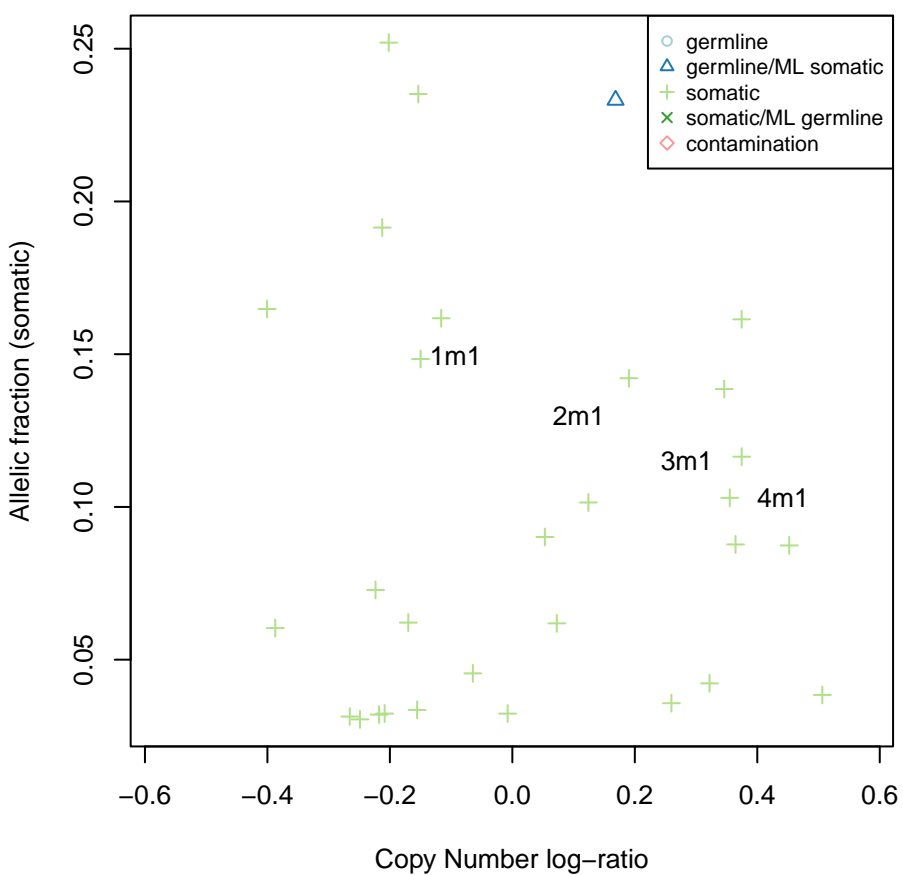
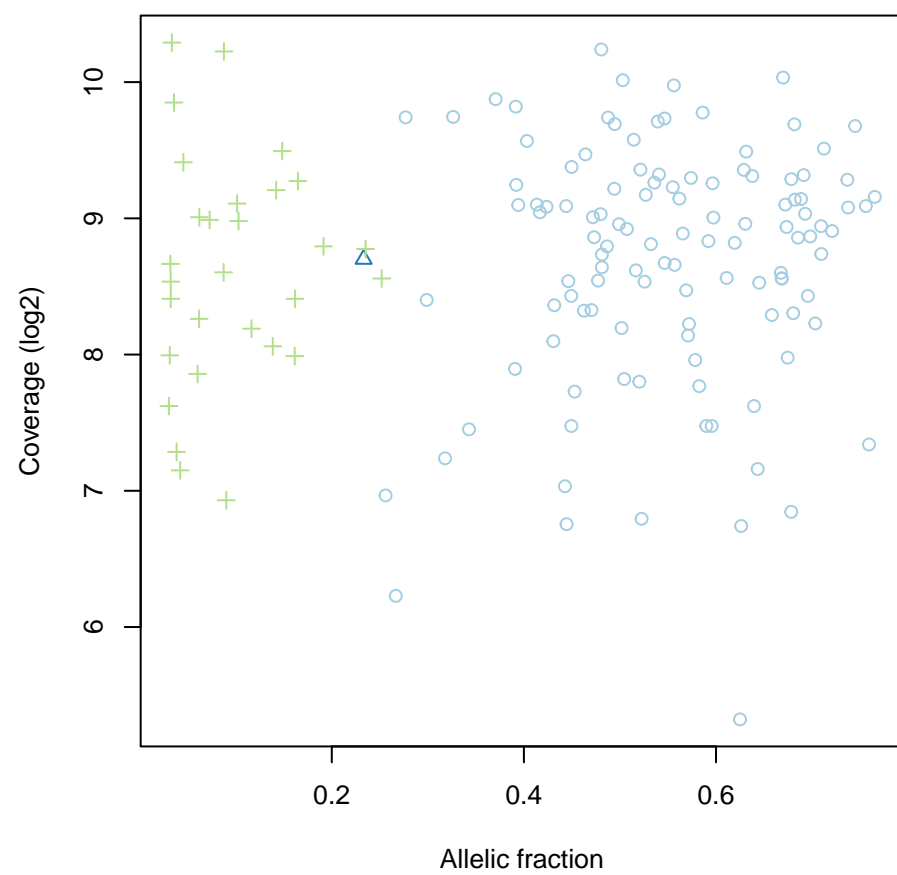
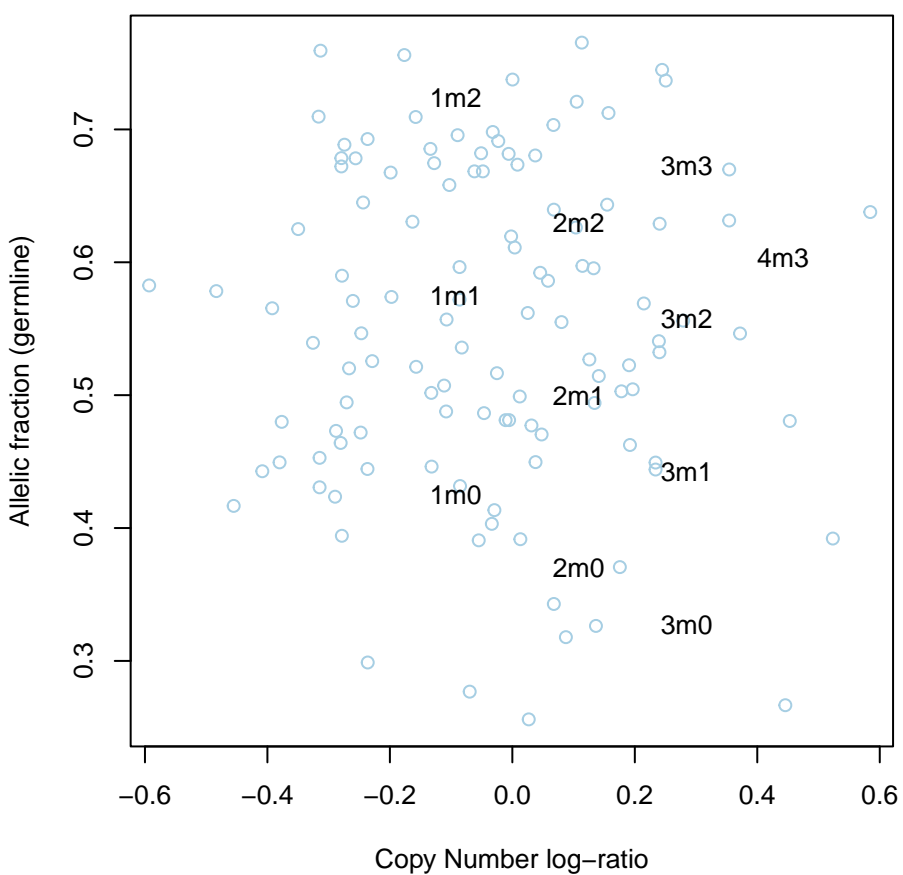


Purity: 0.26 Tumor ploidy: 1.443 SNV log-likelihood: -370.87 GoF: 85.1% Mean coverage: 477;413

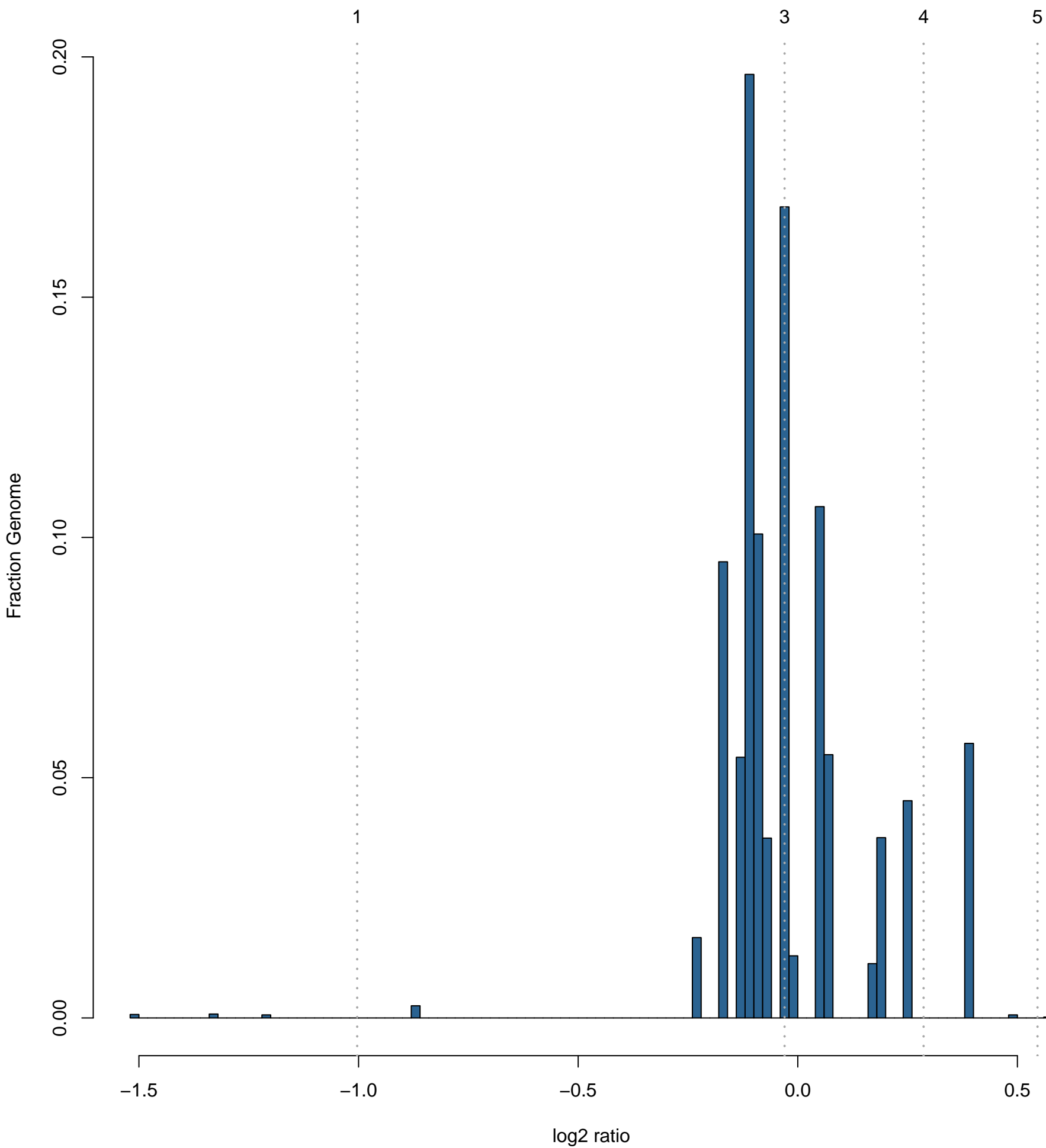


SCNA-fit log-likelihood: -6911.05

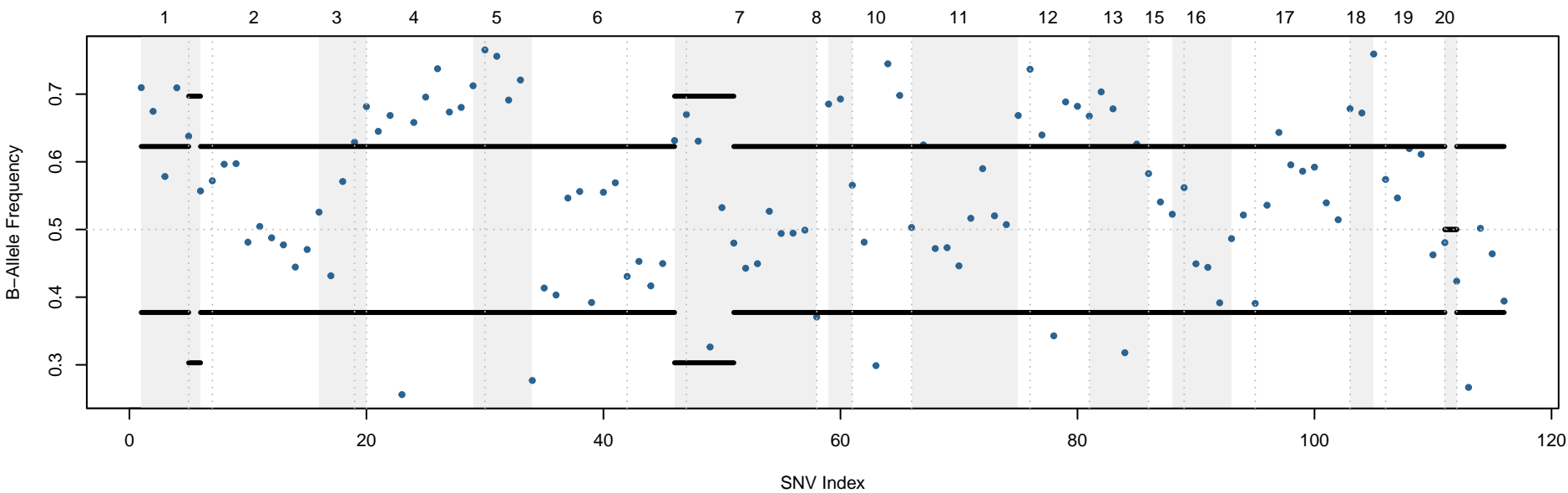




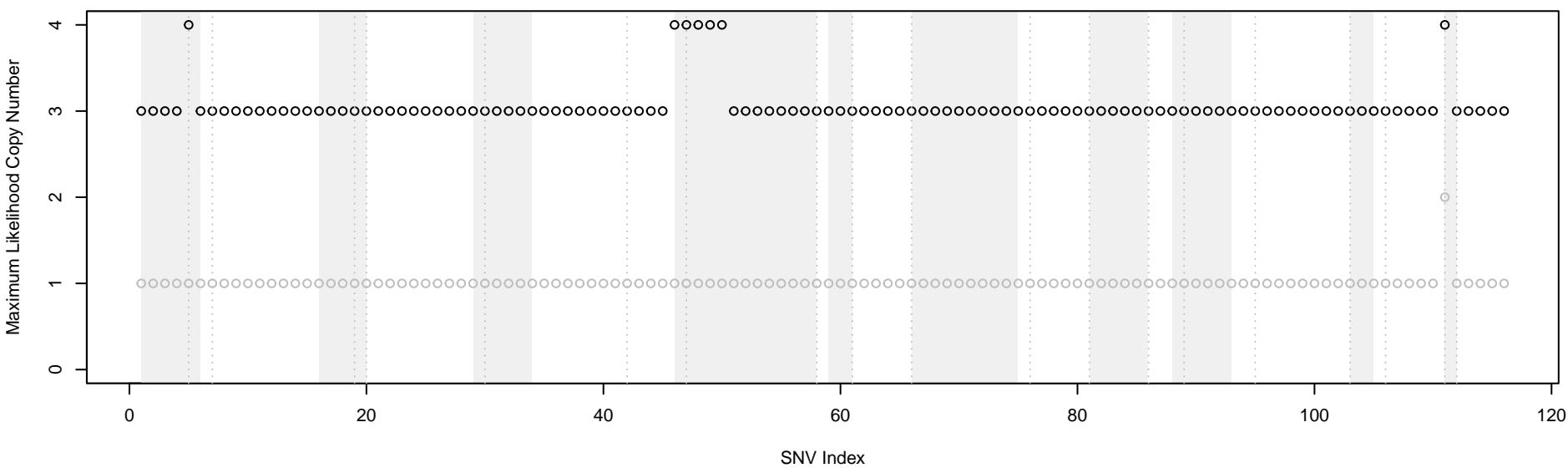
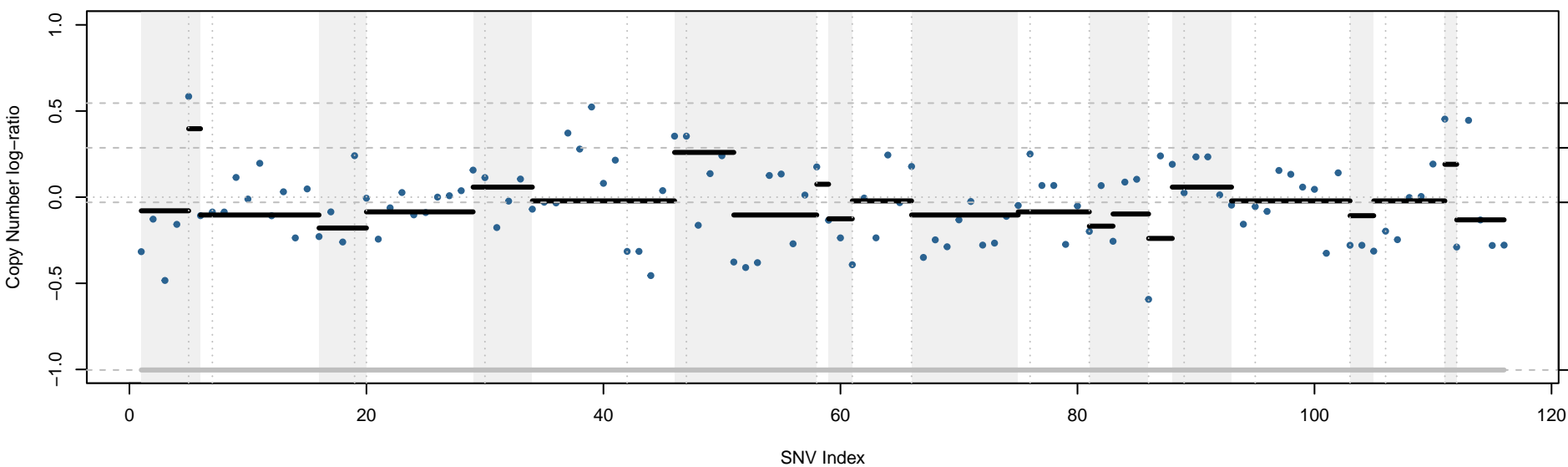
Purity: 0.65 Tumor ploidy: 3.086

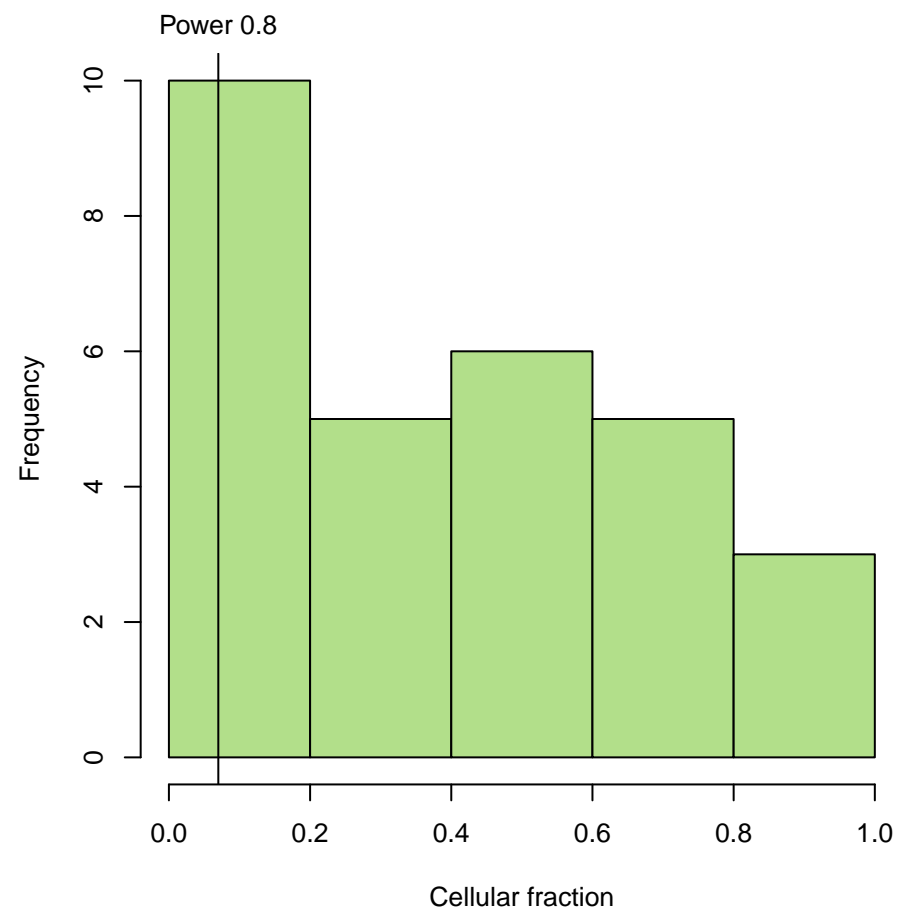
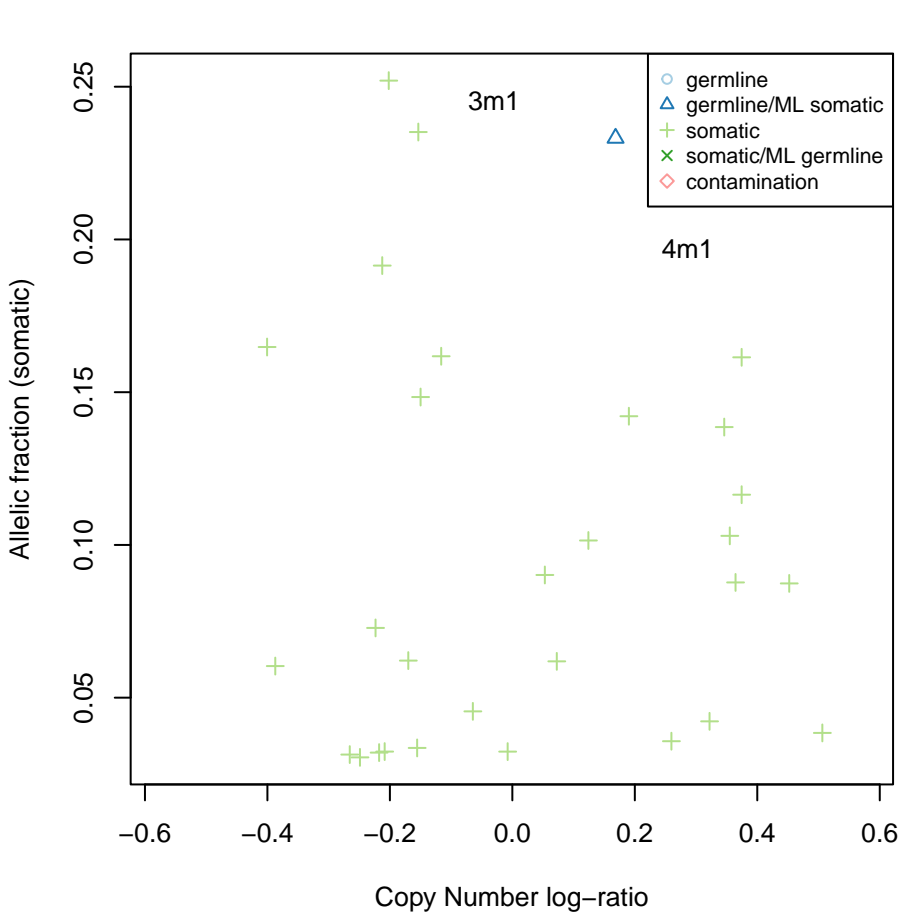
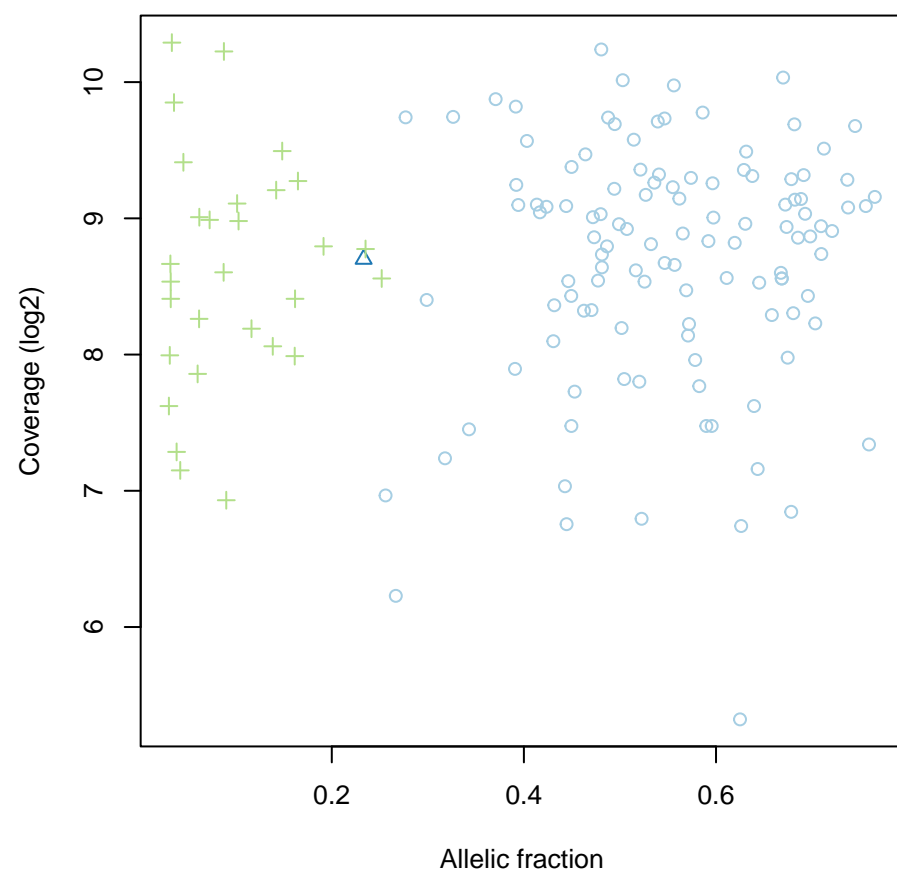
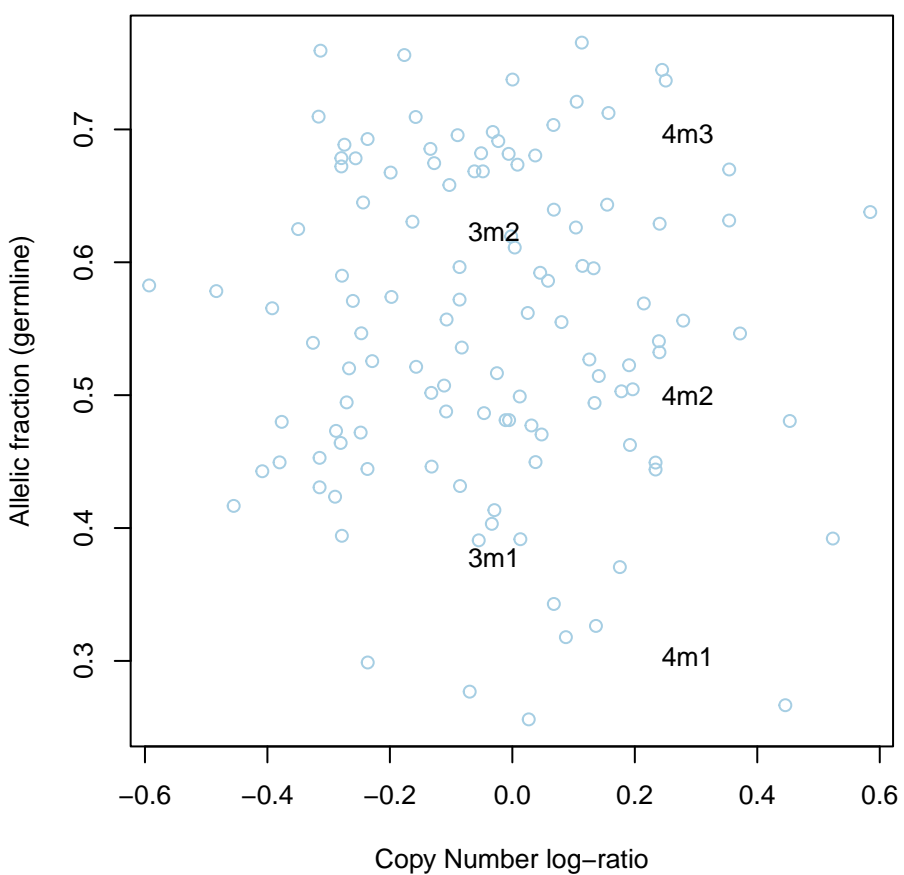


Purity: 0.65 Tumor ploidy: 3.086 SNV log-likelihood: -391.45 GoF: 76.5% Mean coverage: 477;413

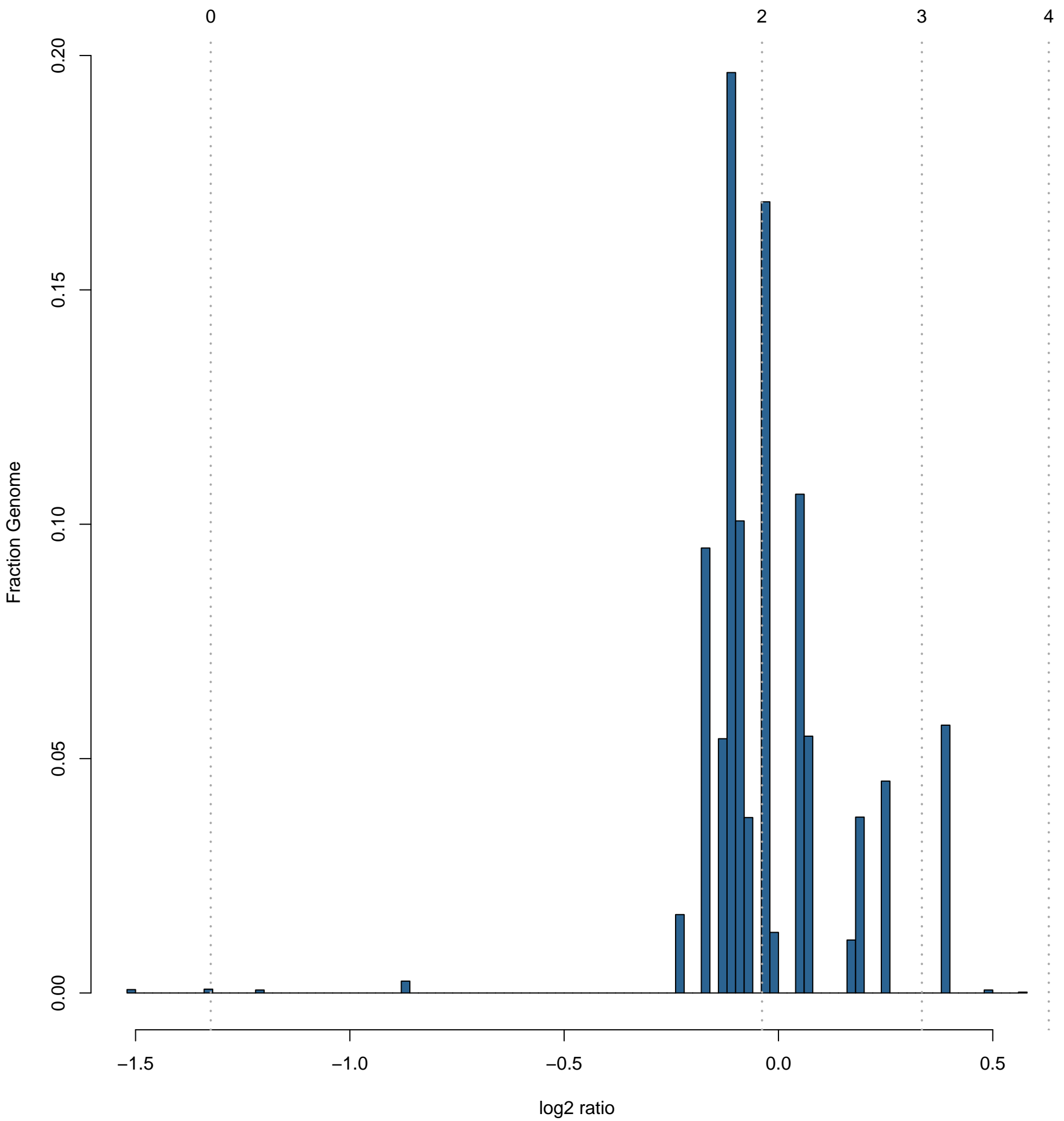


SCNA-fit log-likelihood: -6966.45

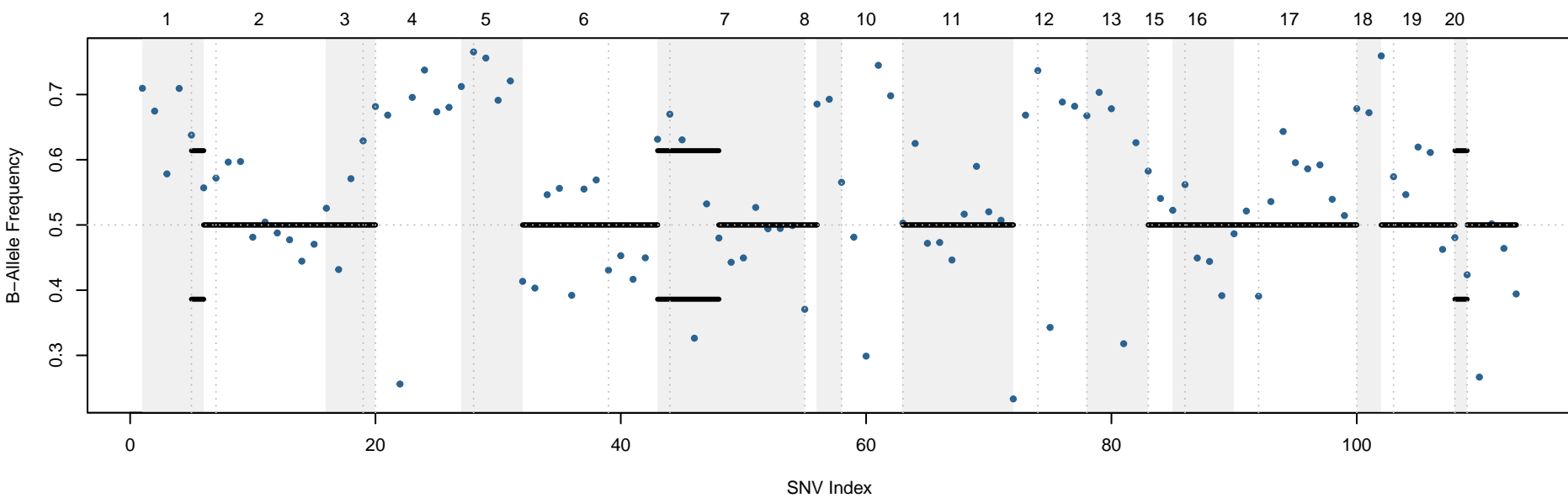




Purity: 0.59 Tumor ploidy: 2.091



Purity: 0.59 Tumor ploidy: 2.091 SNV log-likelihood: -428.4 GoF: 68.4% Mean coverage: 477;413



SCNA-fit log-likelihood: -6965.1

