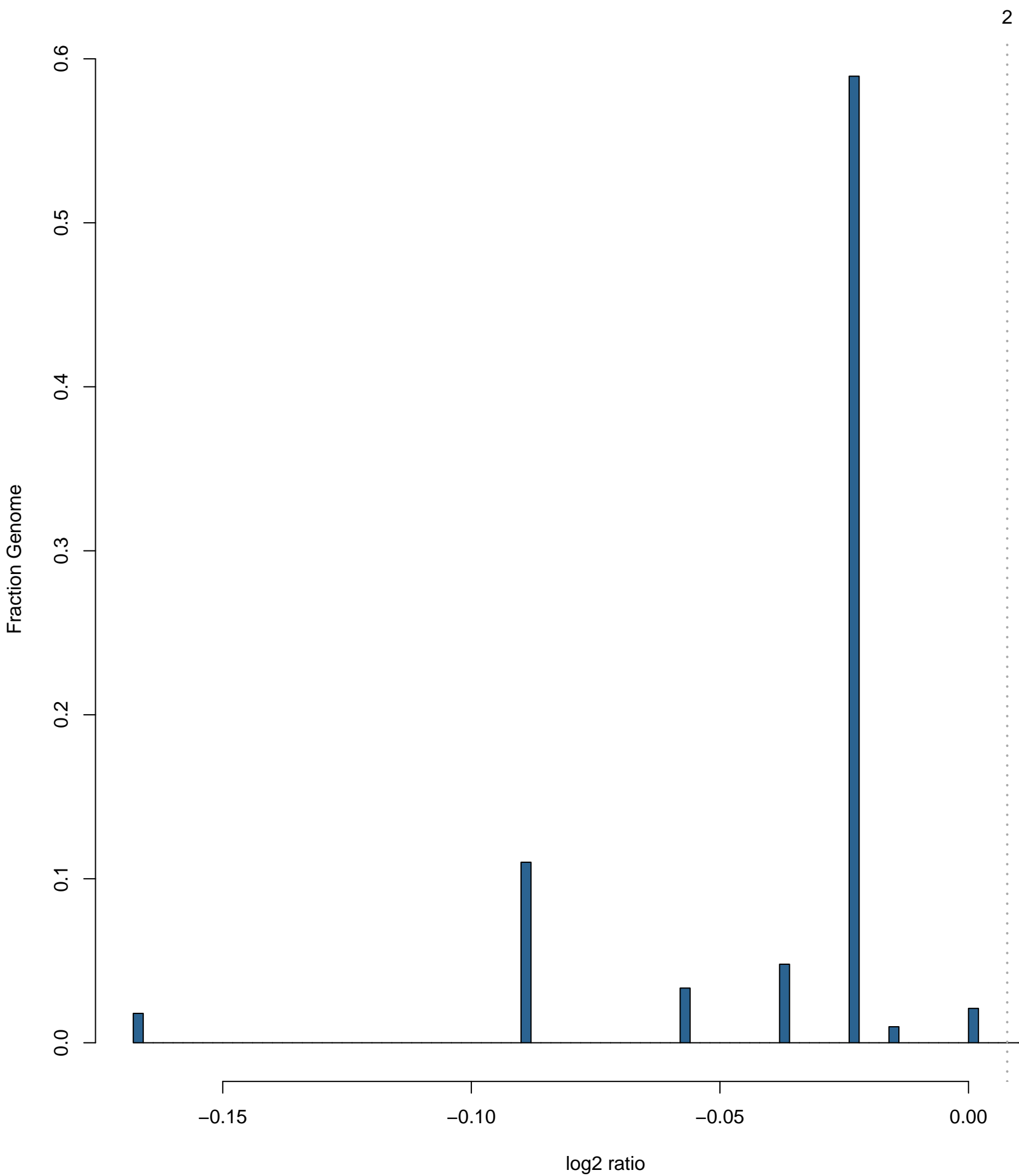
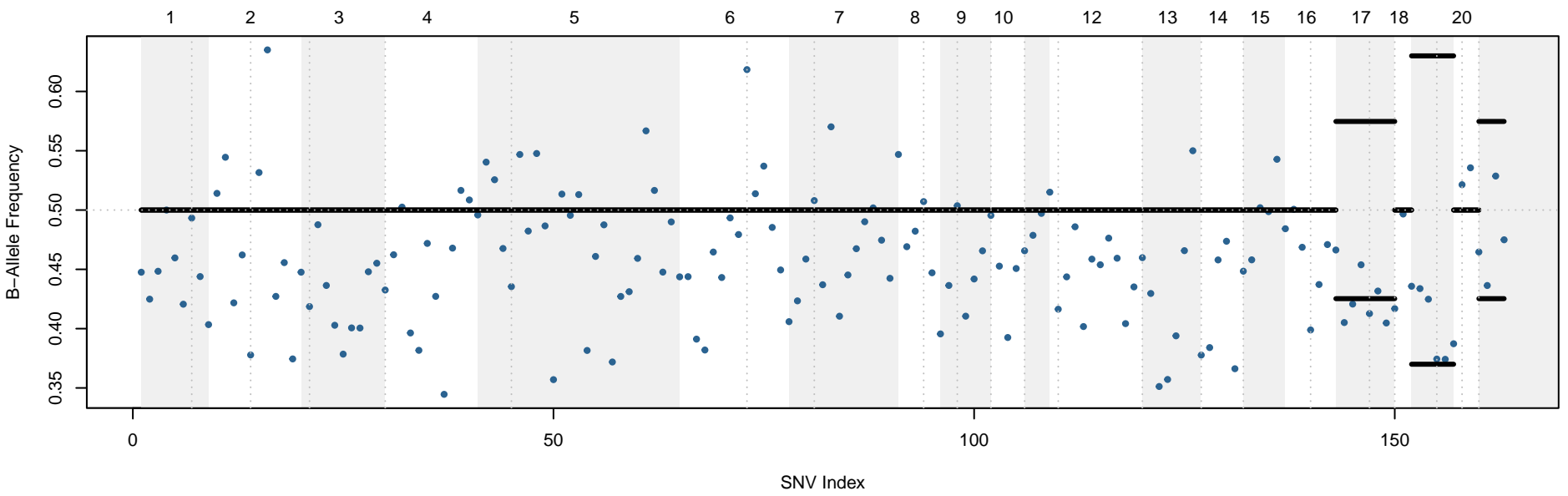


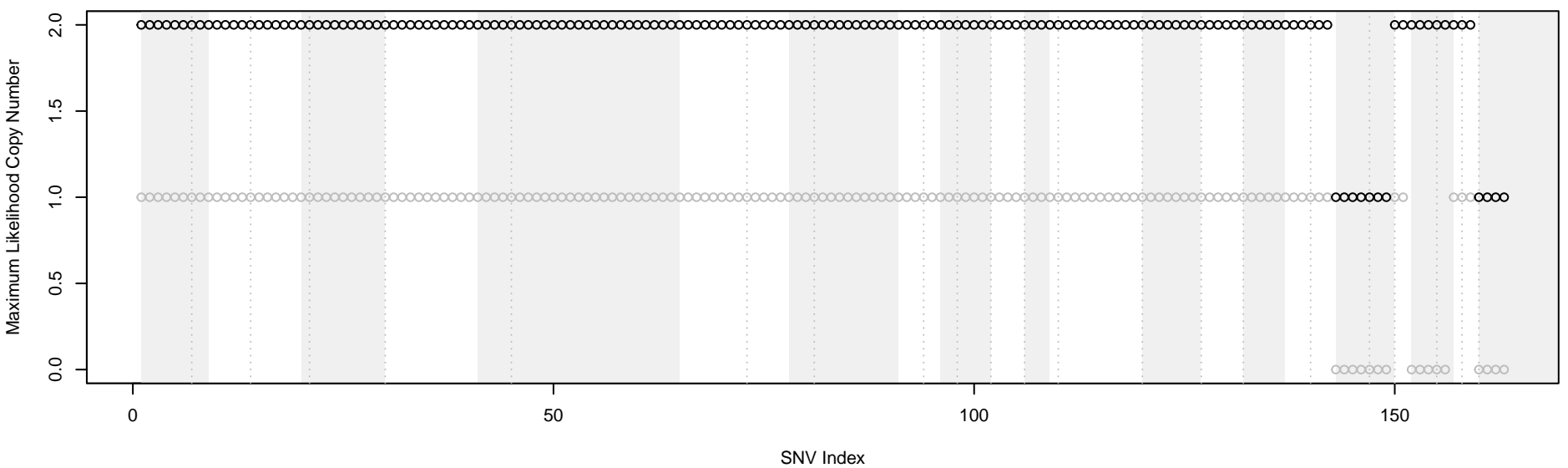
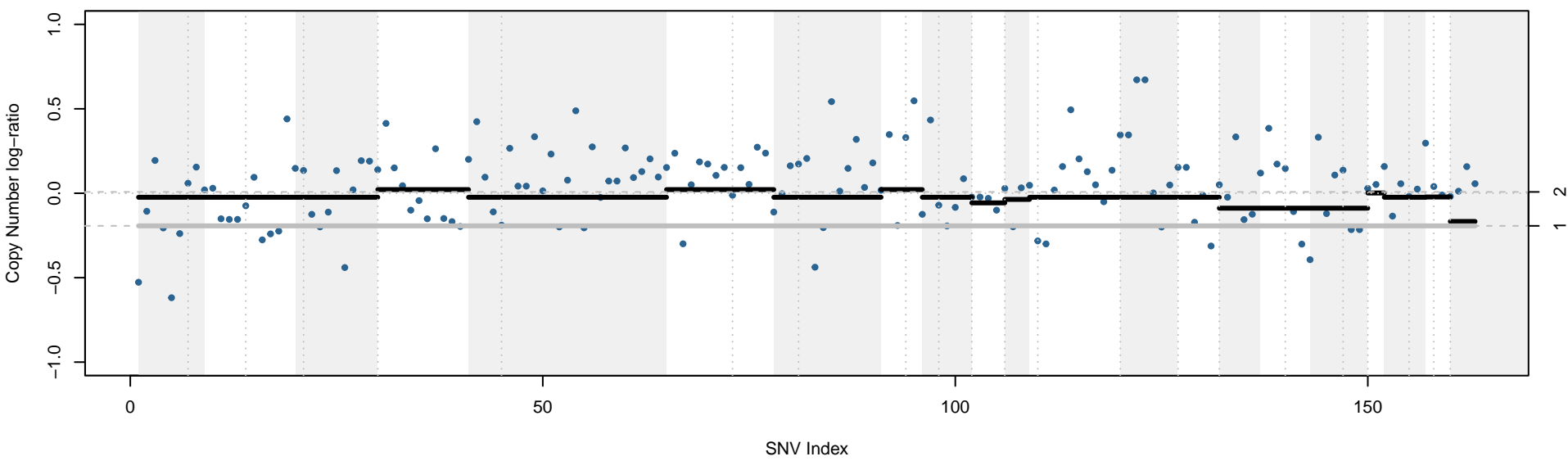
Purity: 0.26 Tumor ploidy: 1.959

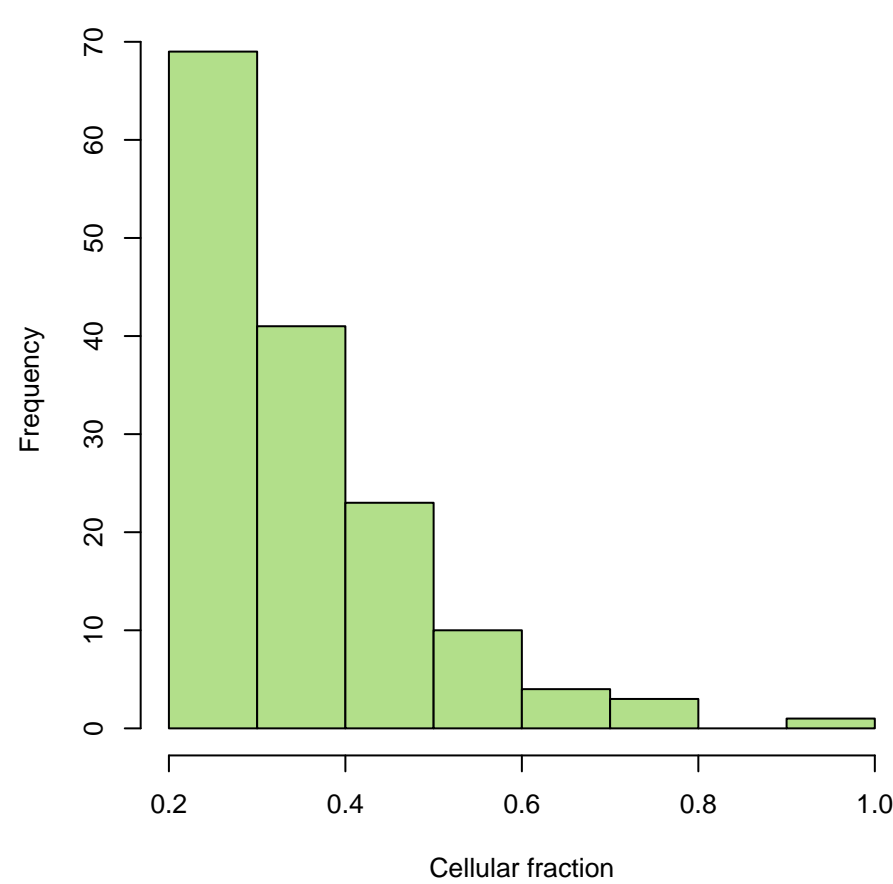
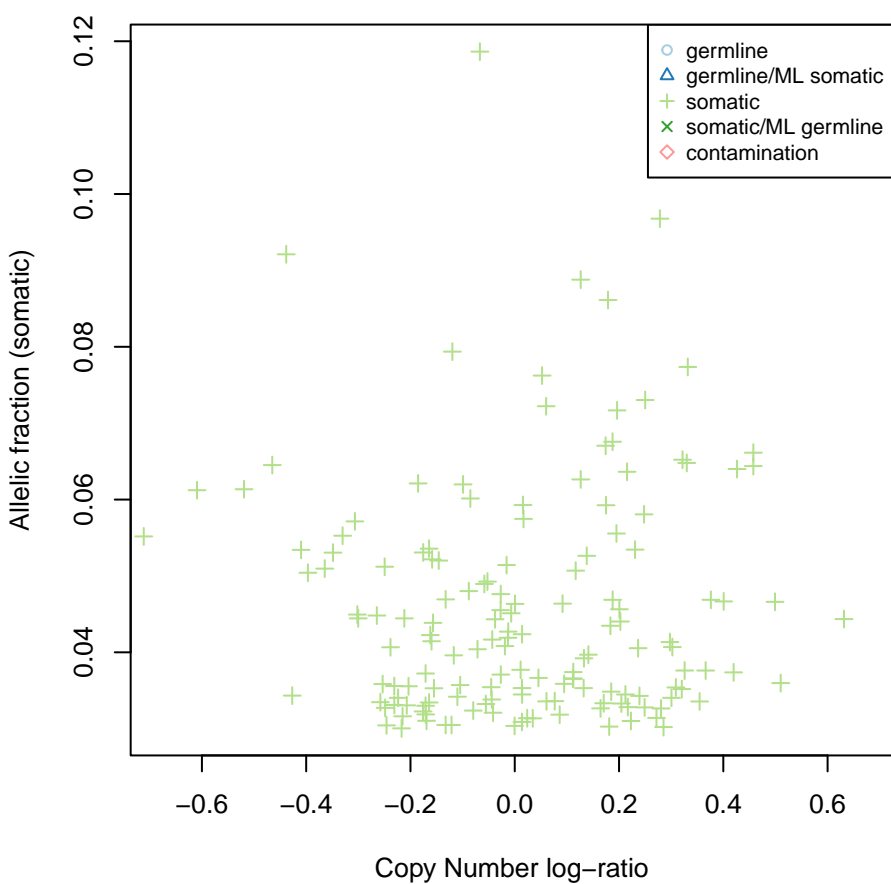
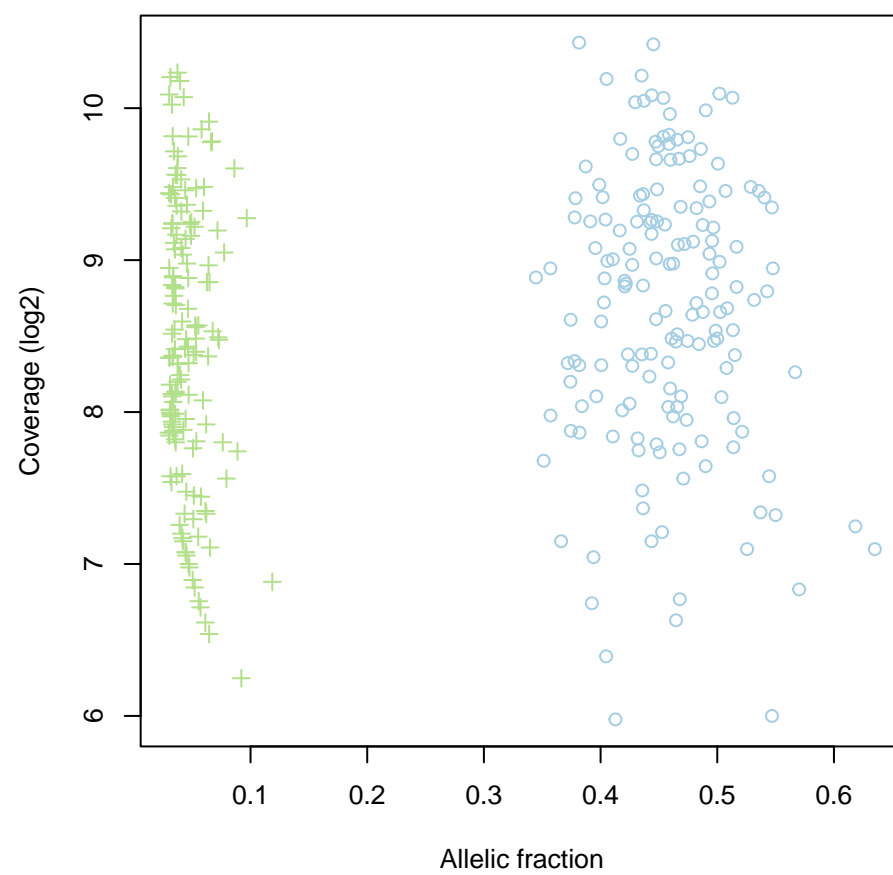
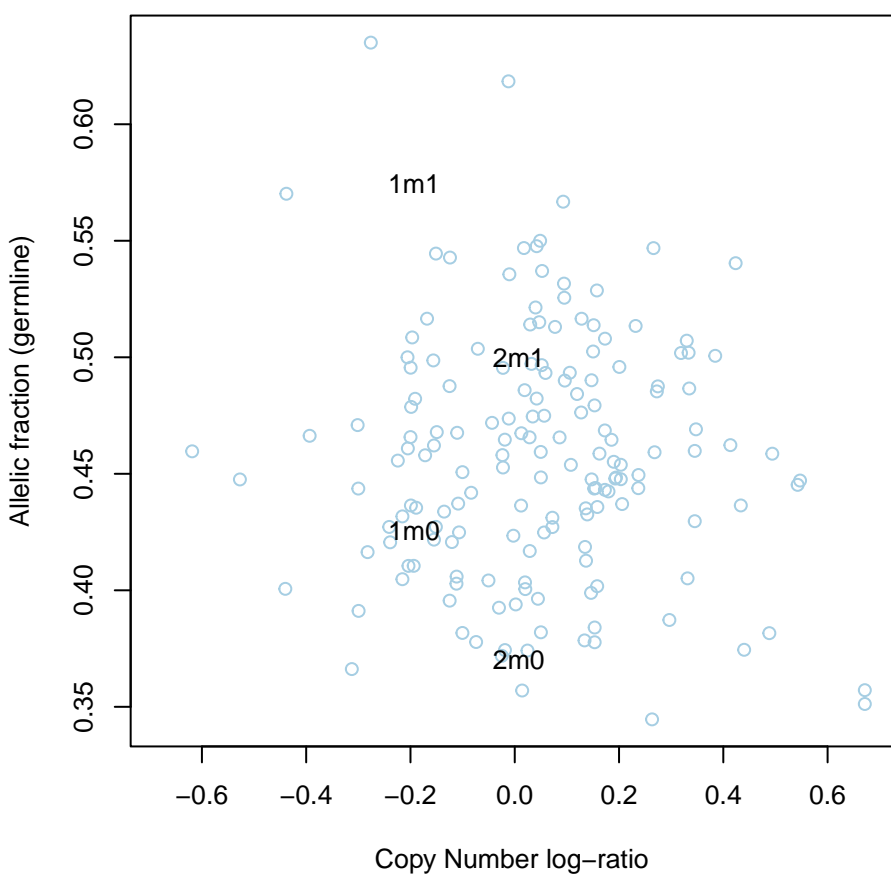


Purity: 0.26 Tumor ploidy: 1.959 SNV log-likelihood: 110.64 GoF: 87.4% Mean coverage: 299;455

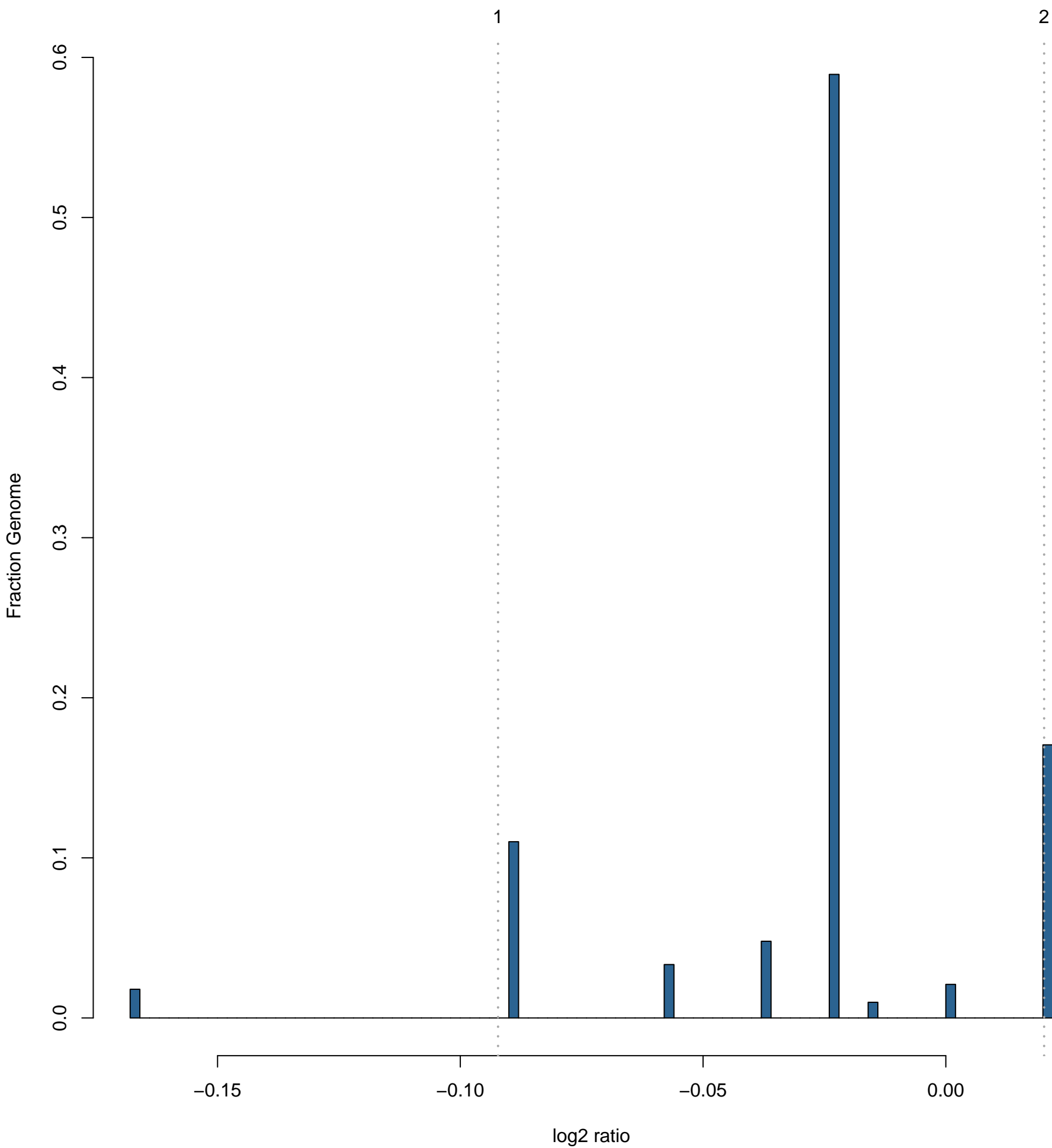


SCNA-fit log-likelihood: -13620.04

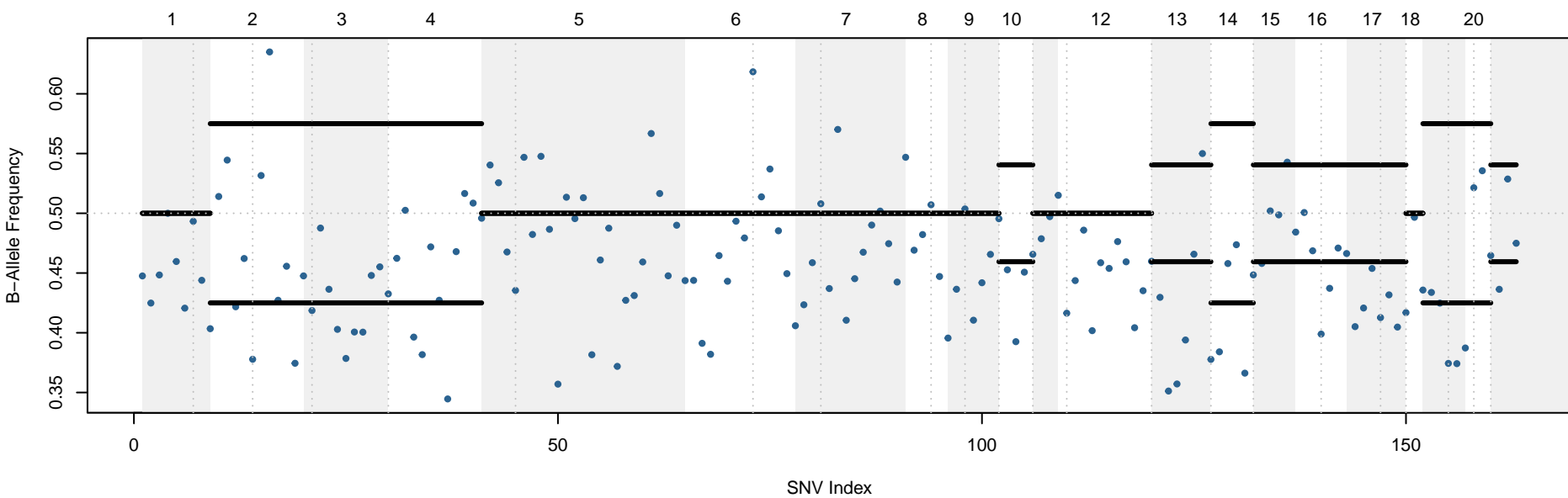




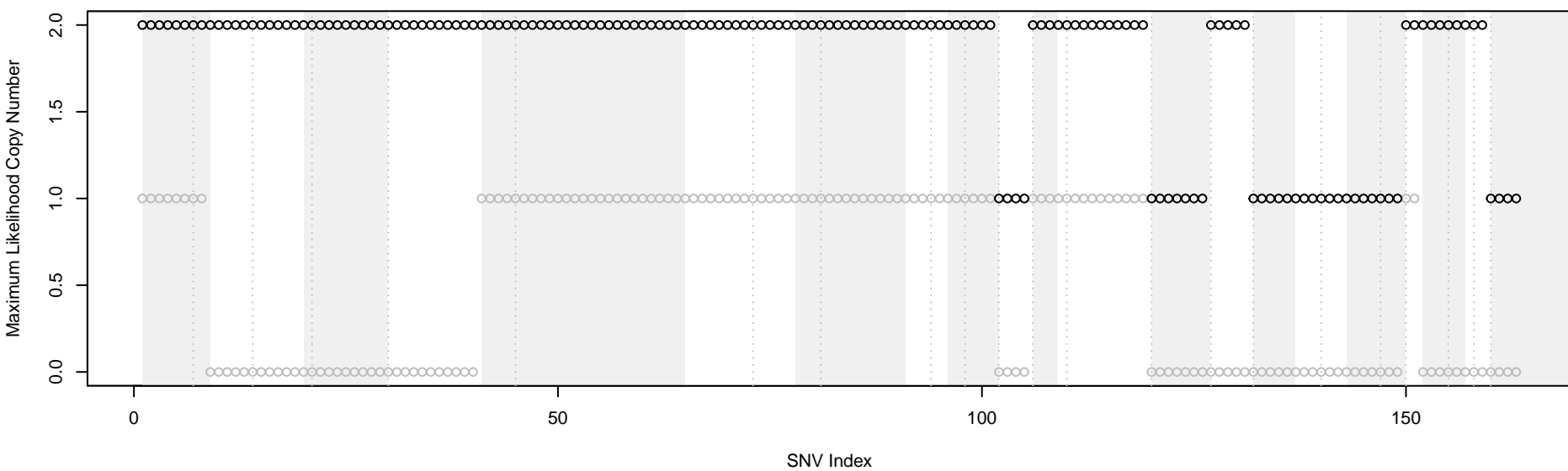
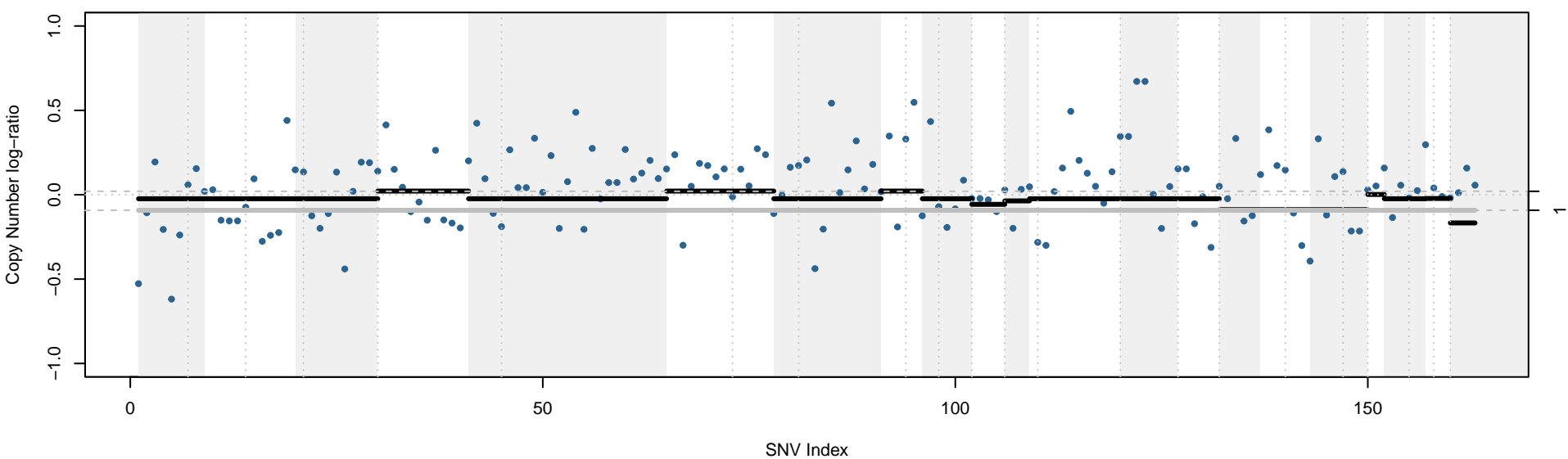
Purity: 0.15 Tumor ploidy: 1.814

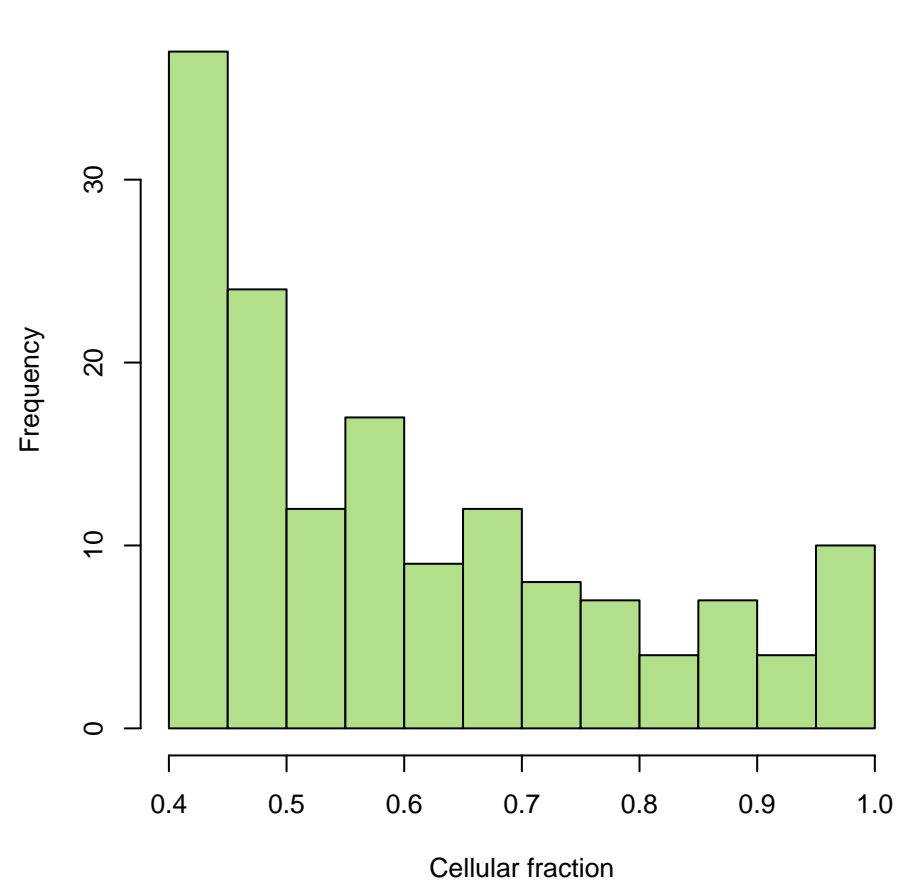
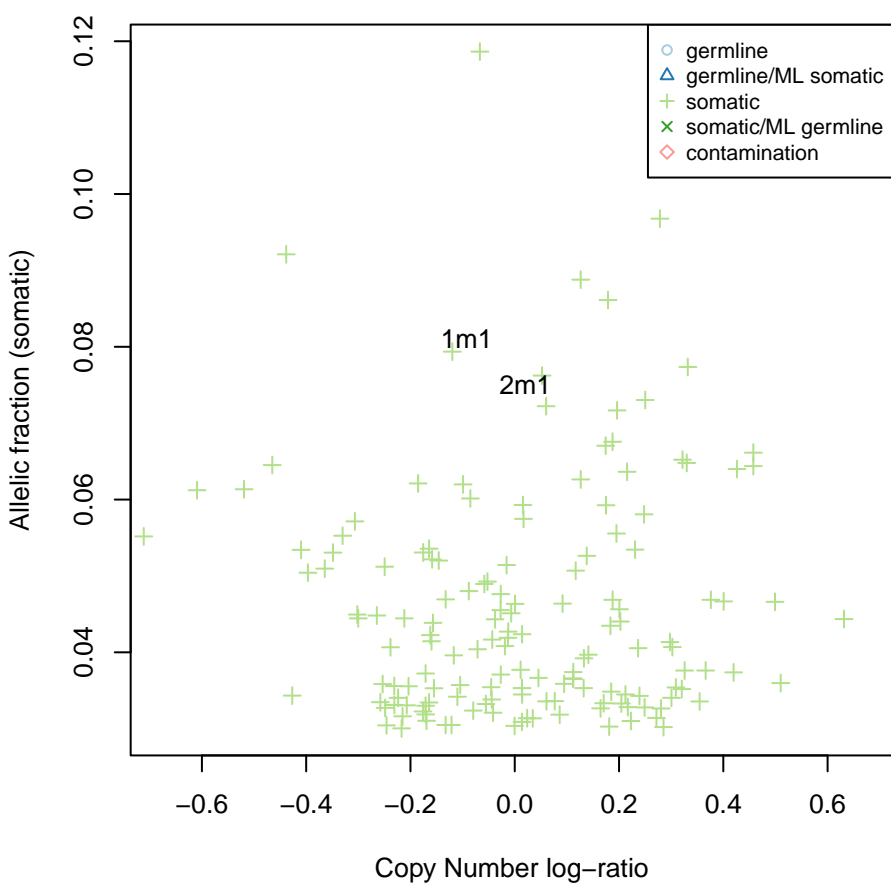
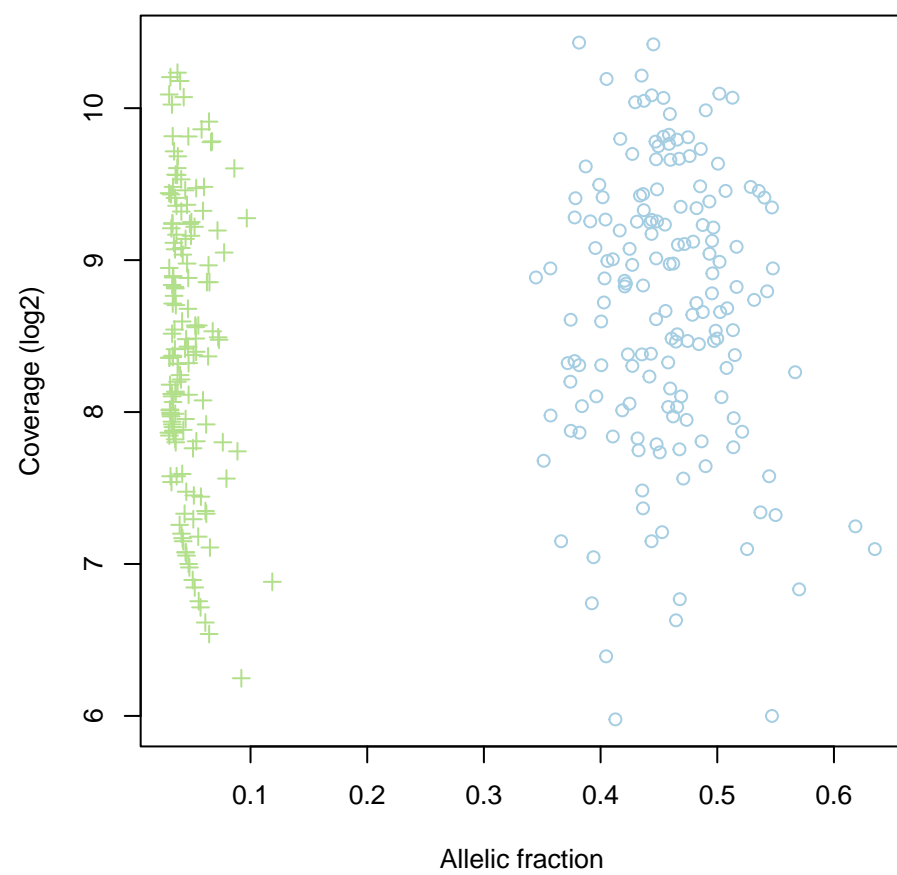
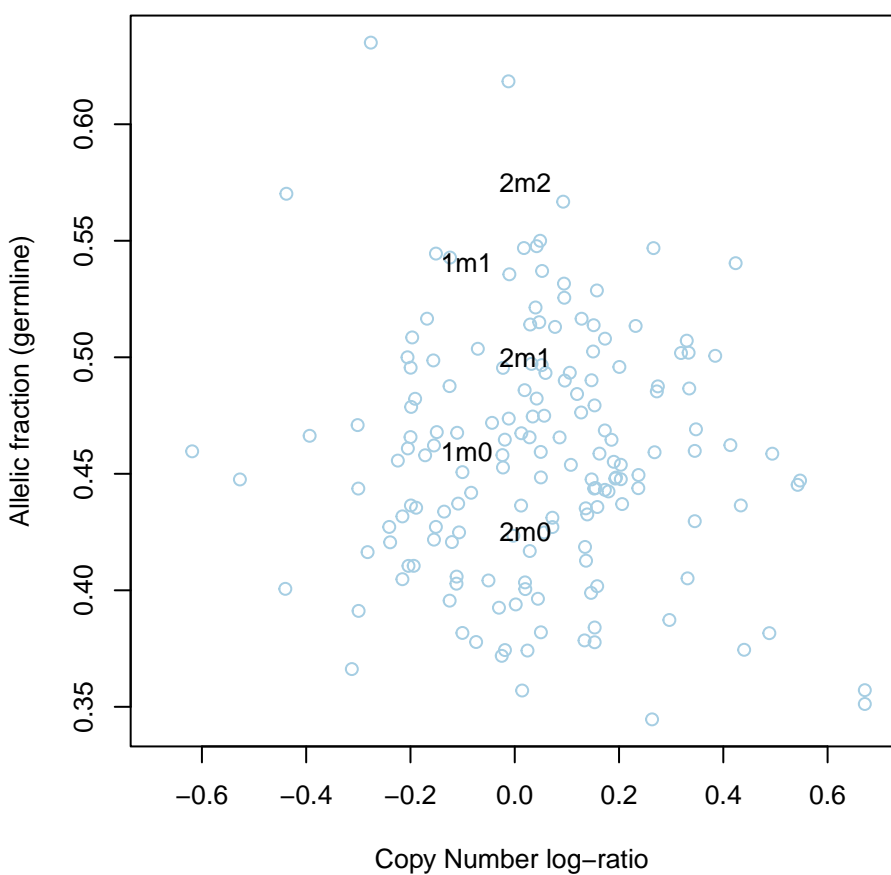


Purity: 0.15 Tumor ploidy: 1.814 SNV log-likelihood: 157.02 GoF: 96.3% Mean coverage: 299;455

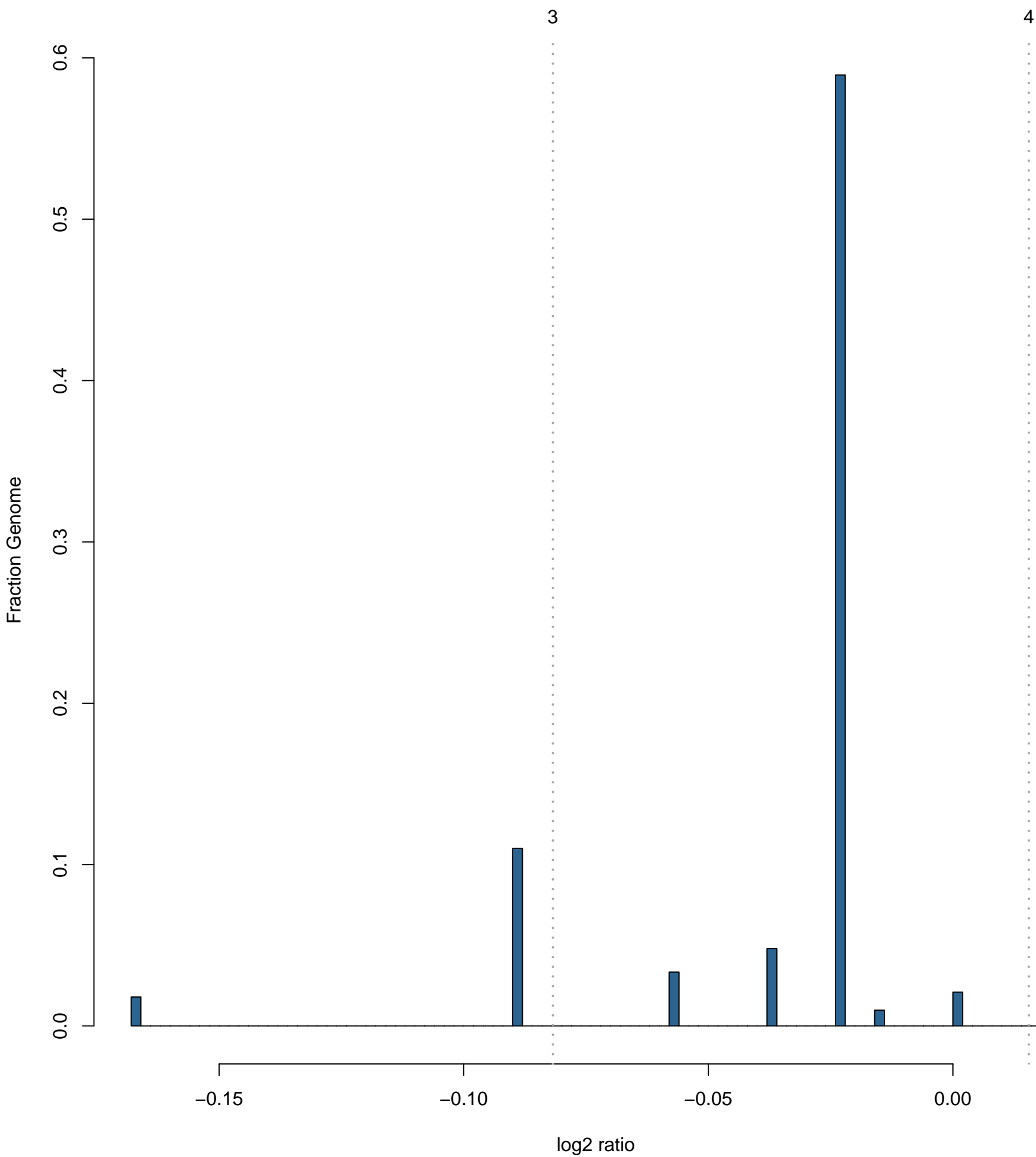


SCNA-fit log-likelihood: -13614.5

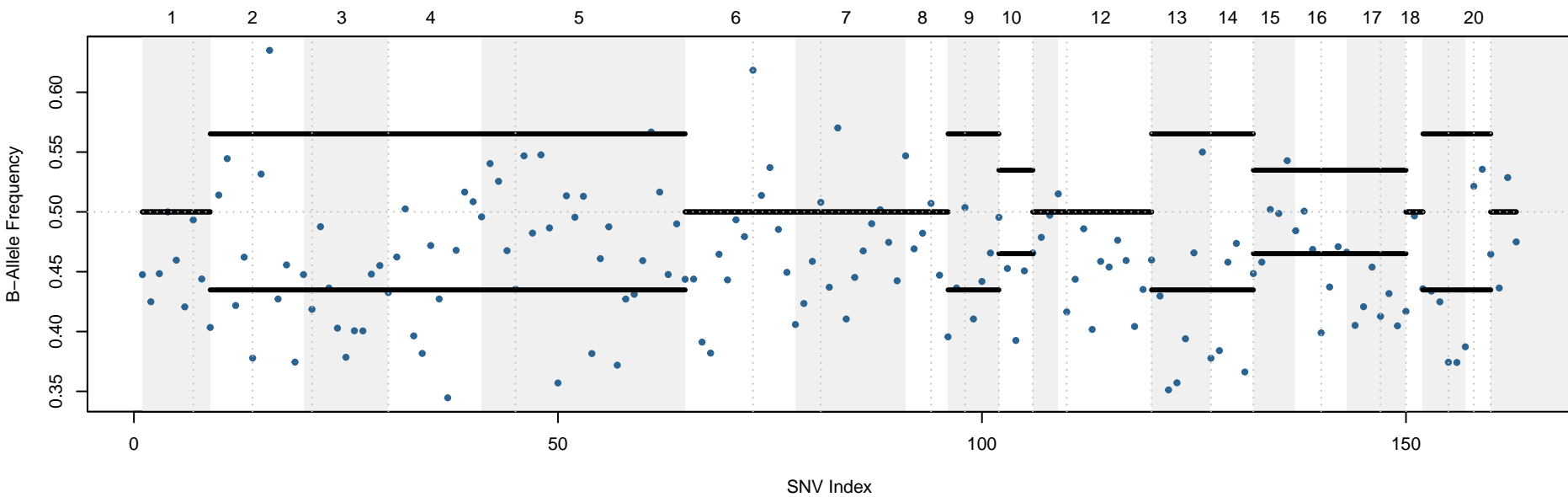




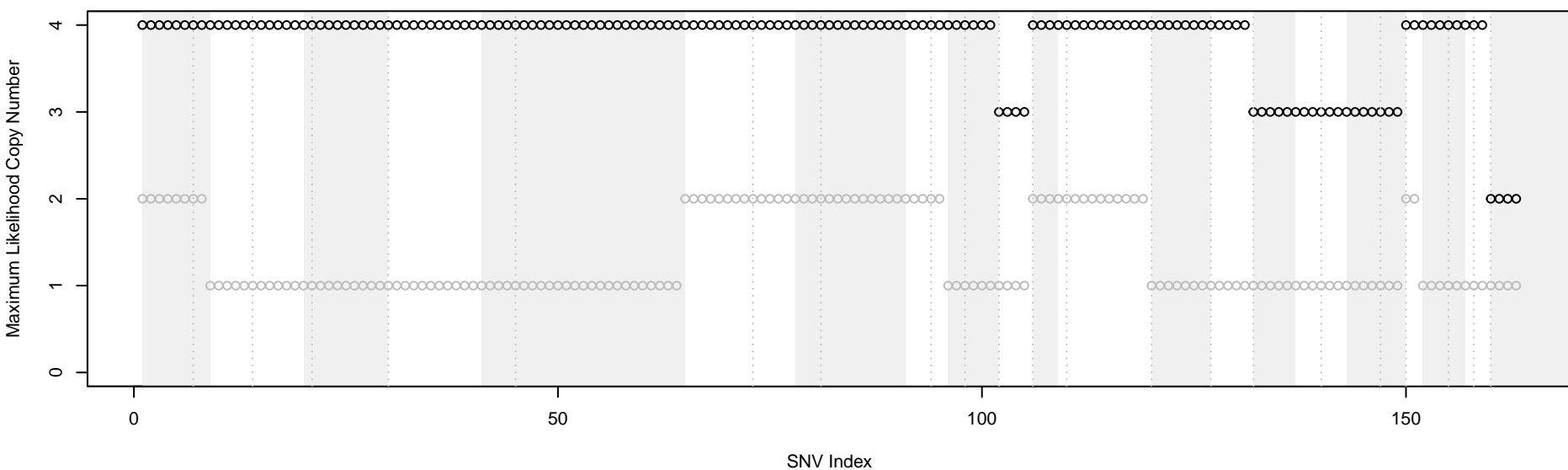
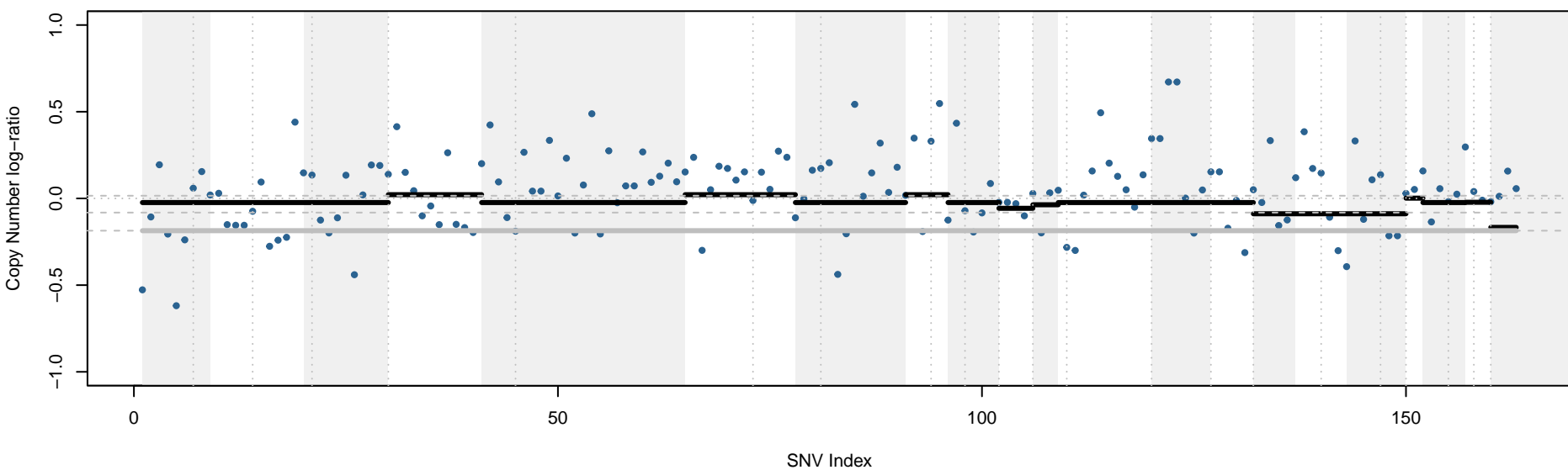
Purity: 0.15 Tumor ploidy: 3.836



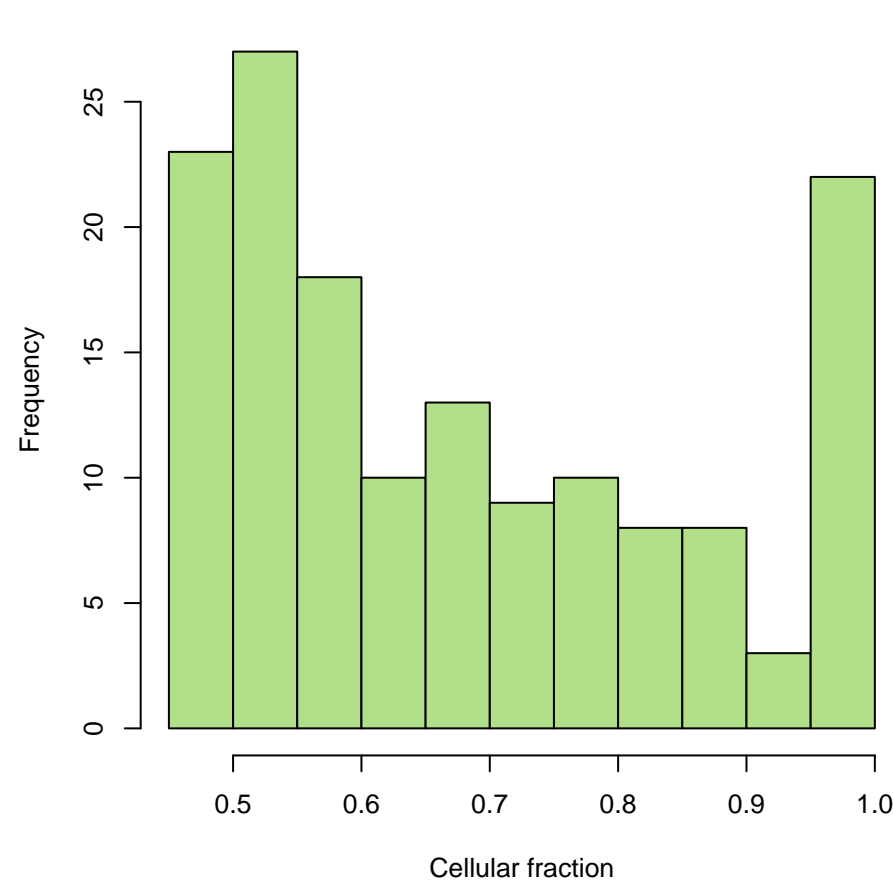
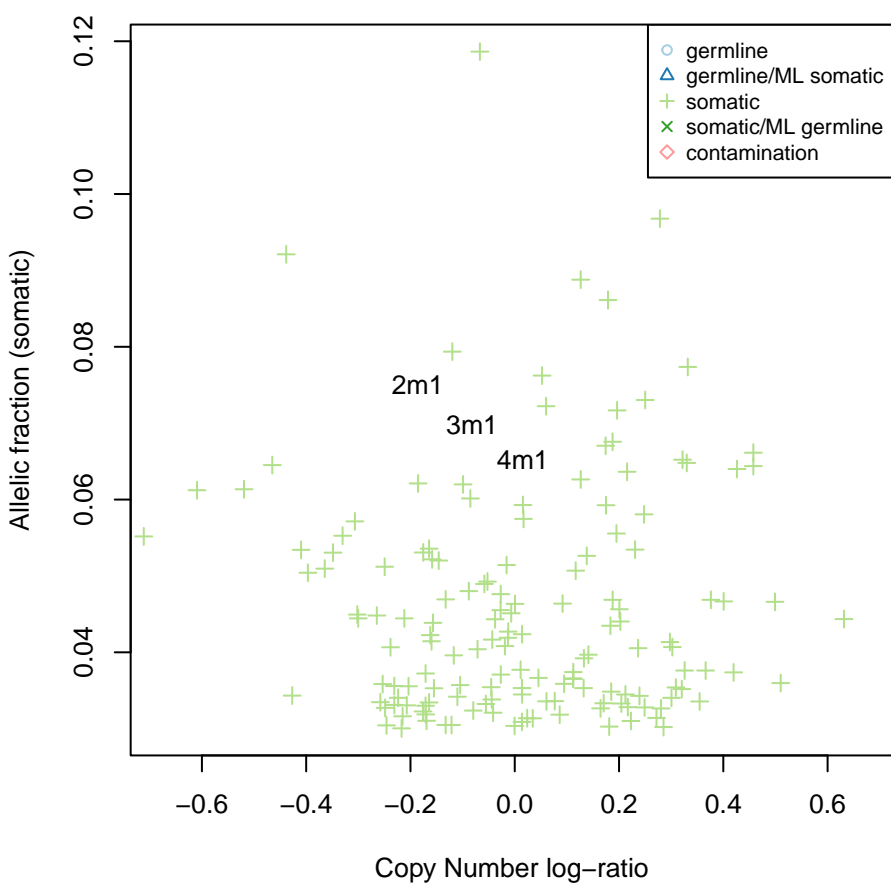
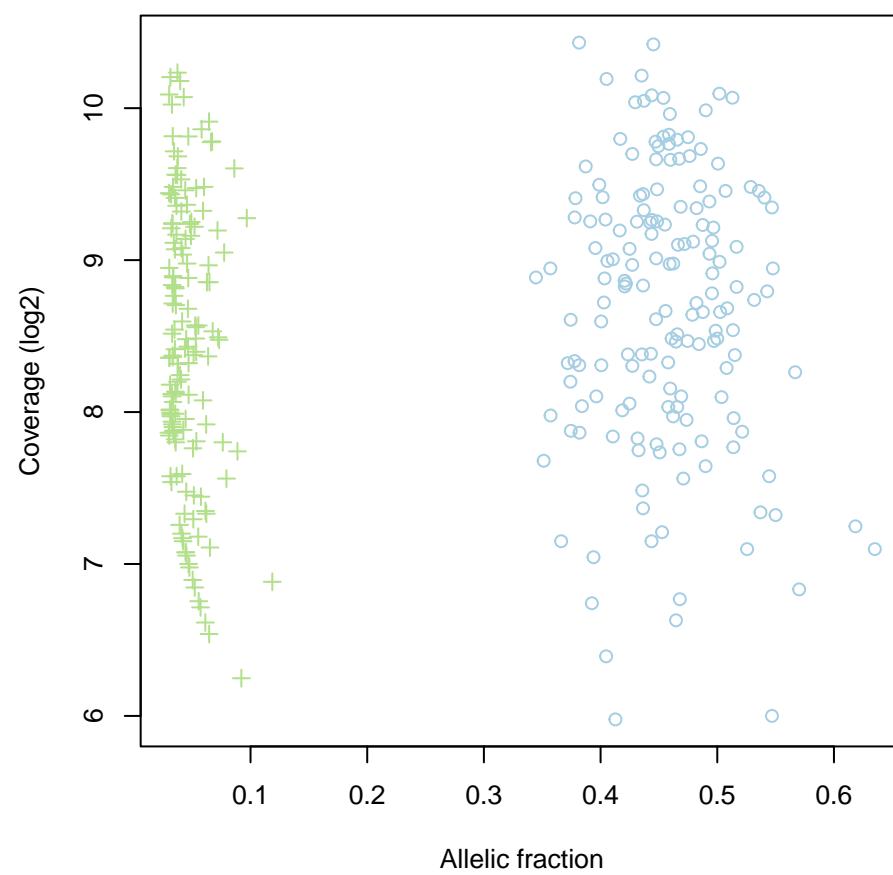
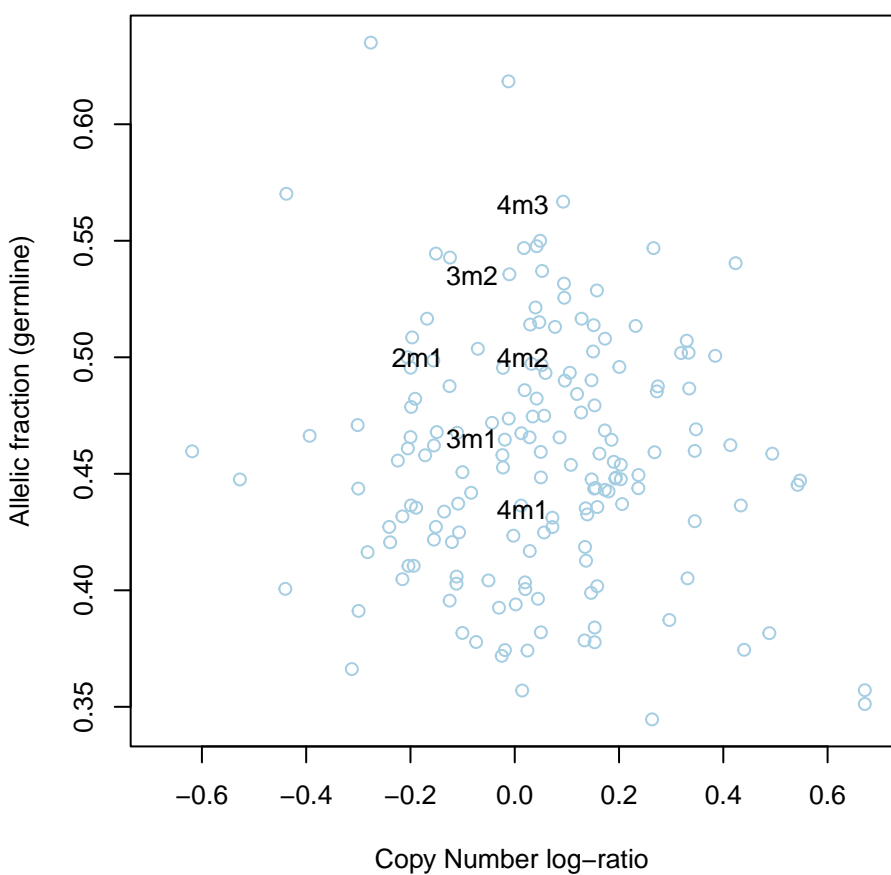
Purity: 0.15 Tumor ploidy: 3.836 SNV log-likelihood: -49.75 GoF: 97.1% Mean coverage: 299;455



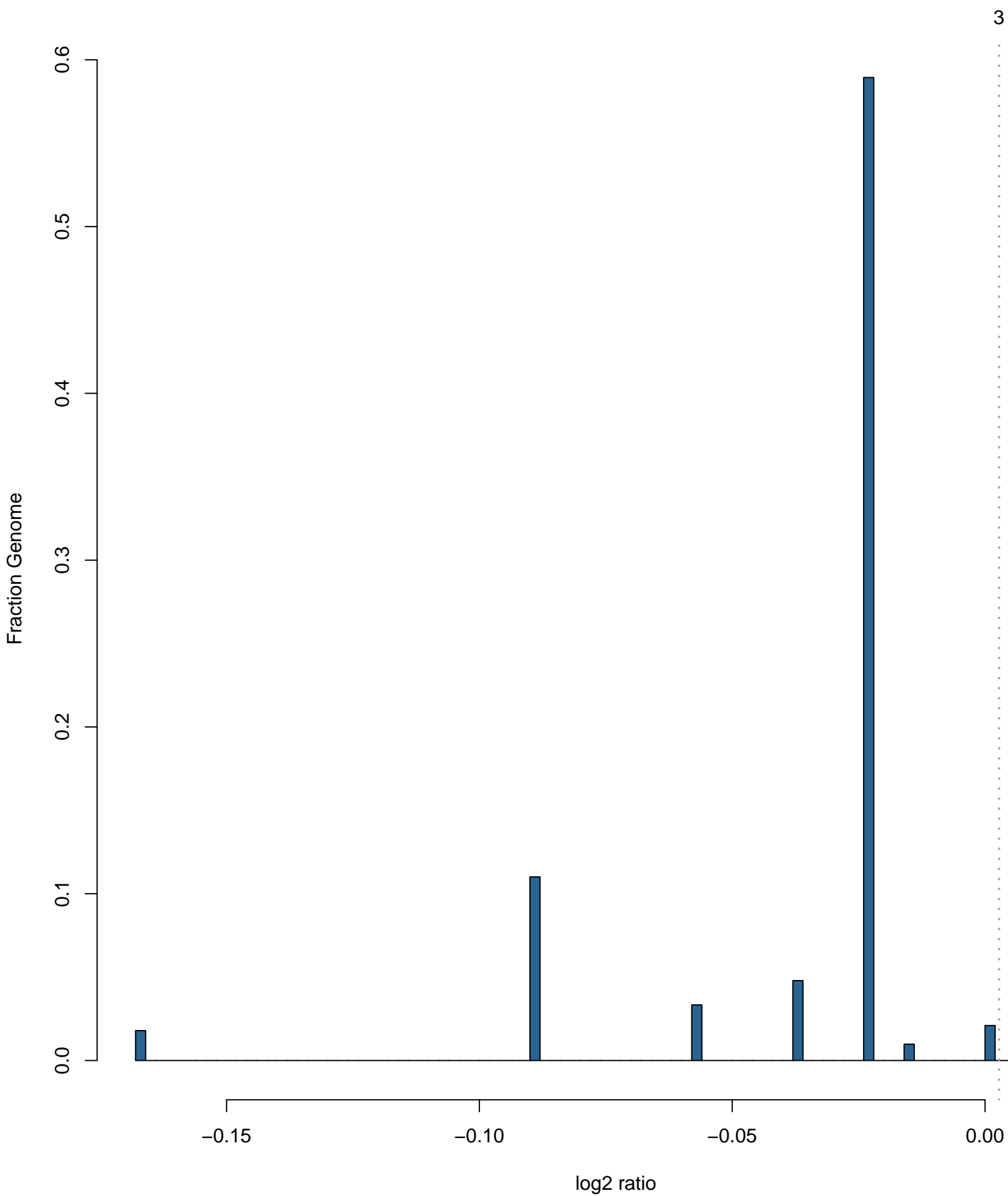
SCNA-fit log-likelihood: -13588.85



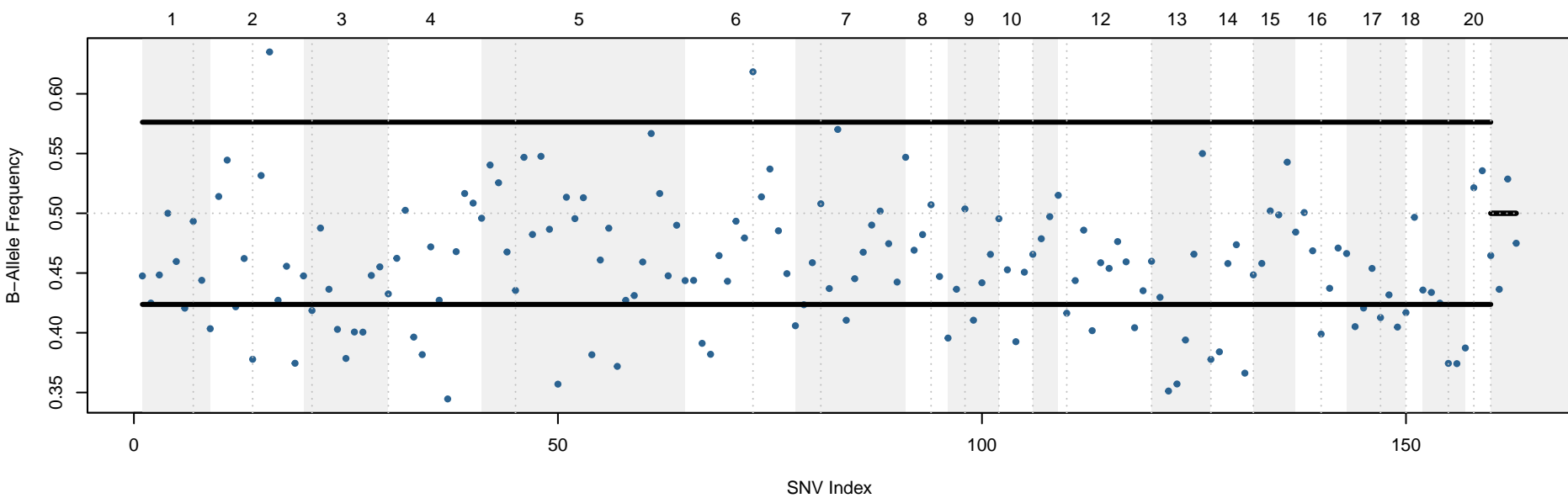




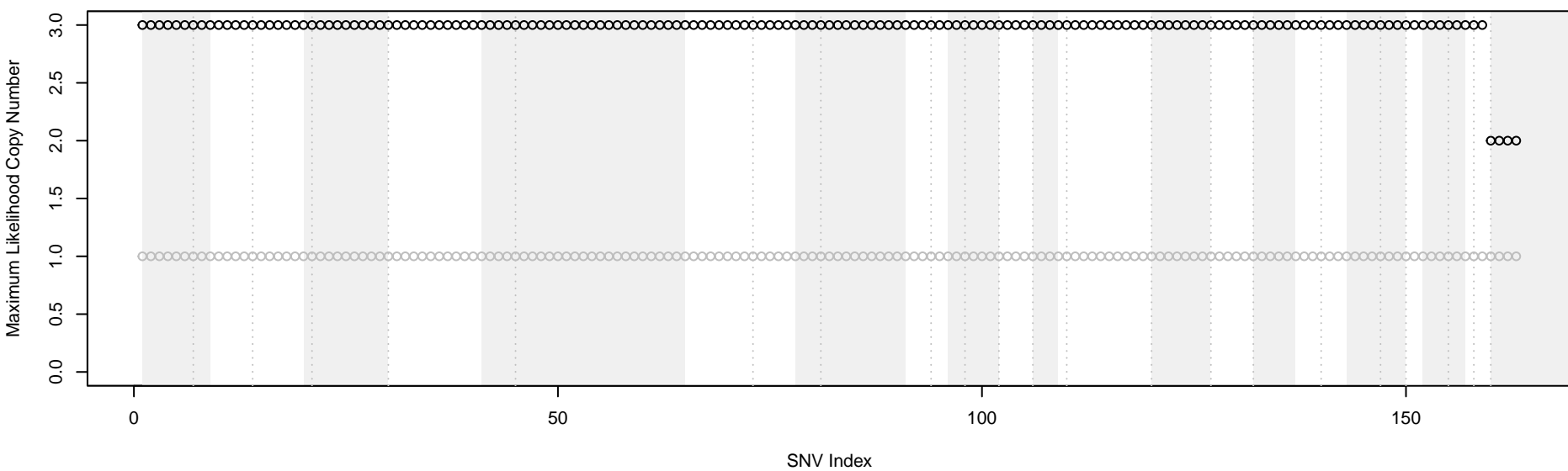
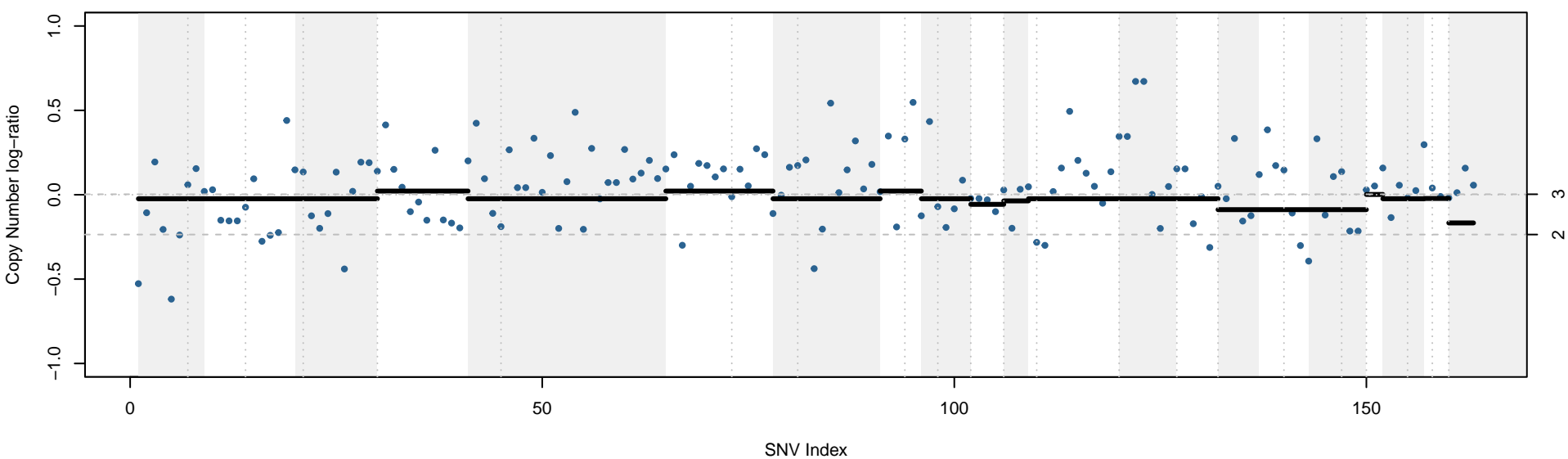
Purity: 0.36 Tumor ploidy: 2.987

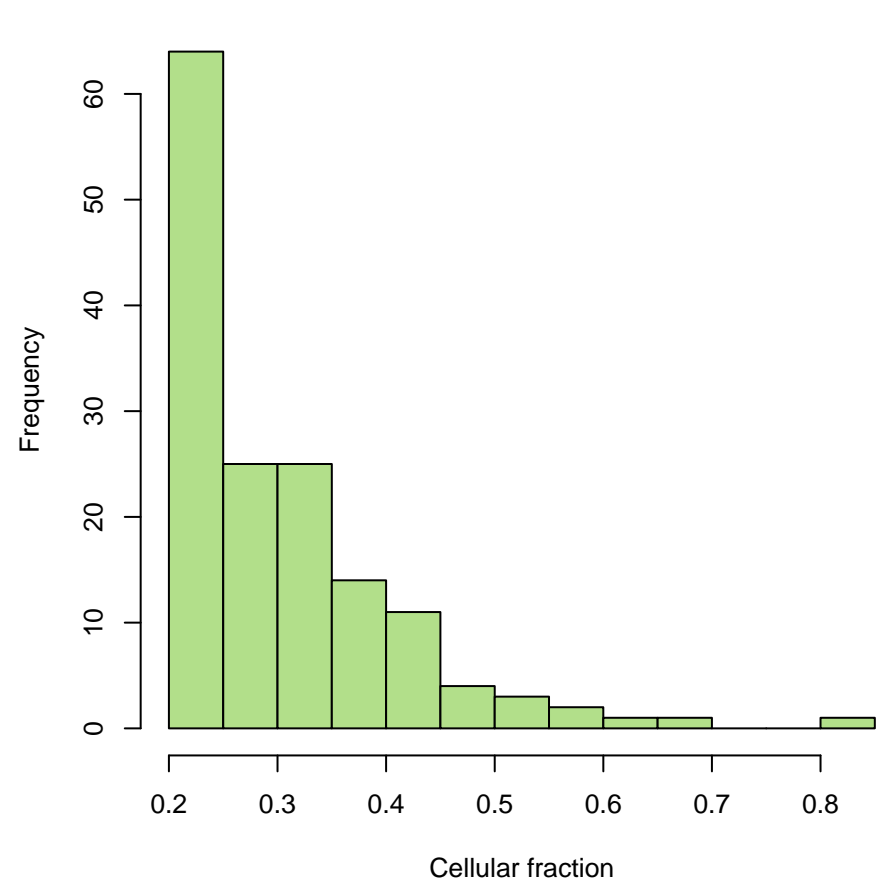
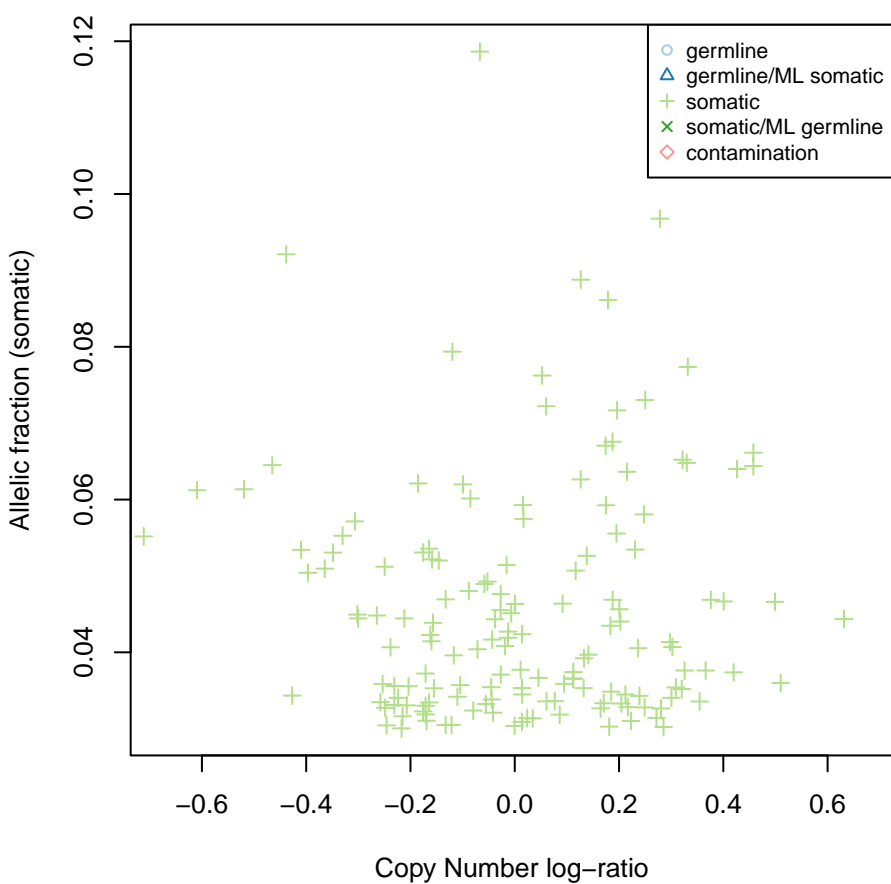
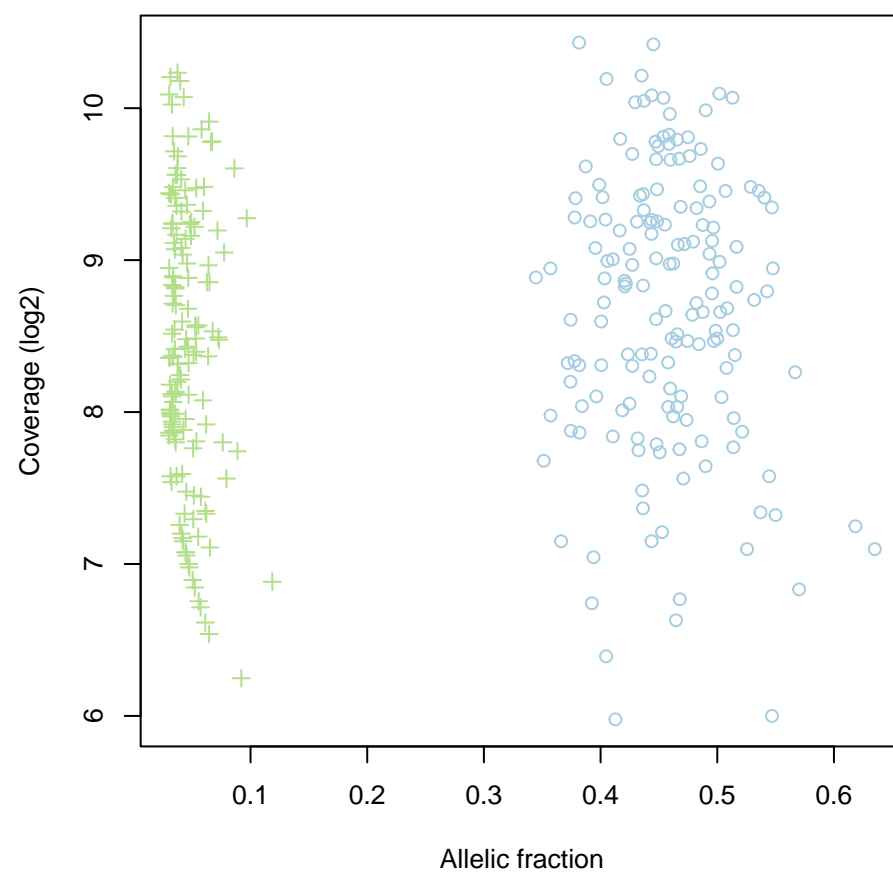
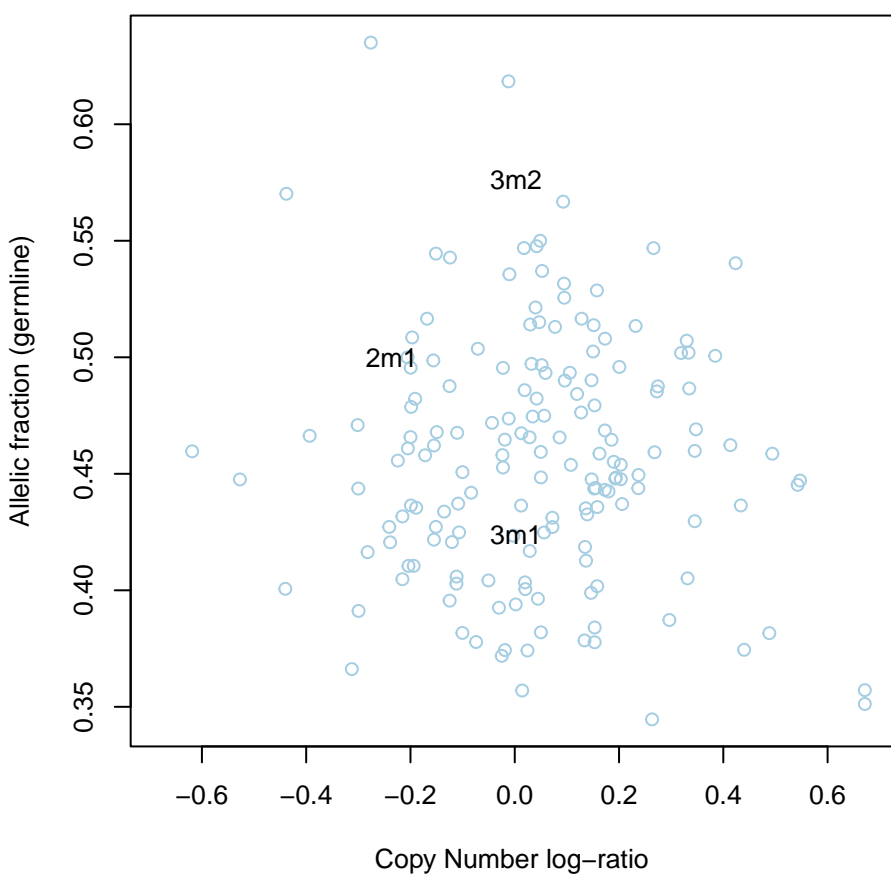


Purity: 0.36 Tumor ploidy: 2.987 SNV log-likelihood: -56.02 GoF: 83.7% Mean coverage: 299;455

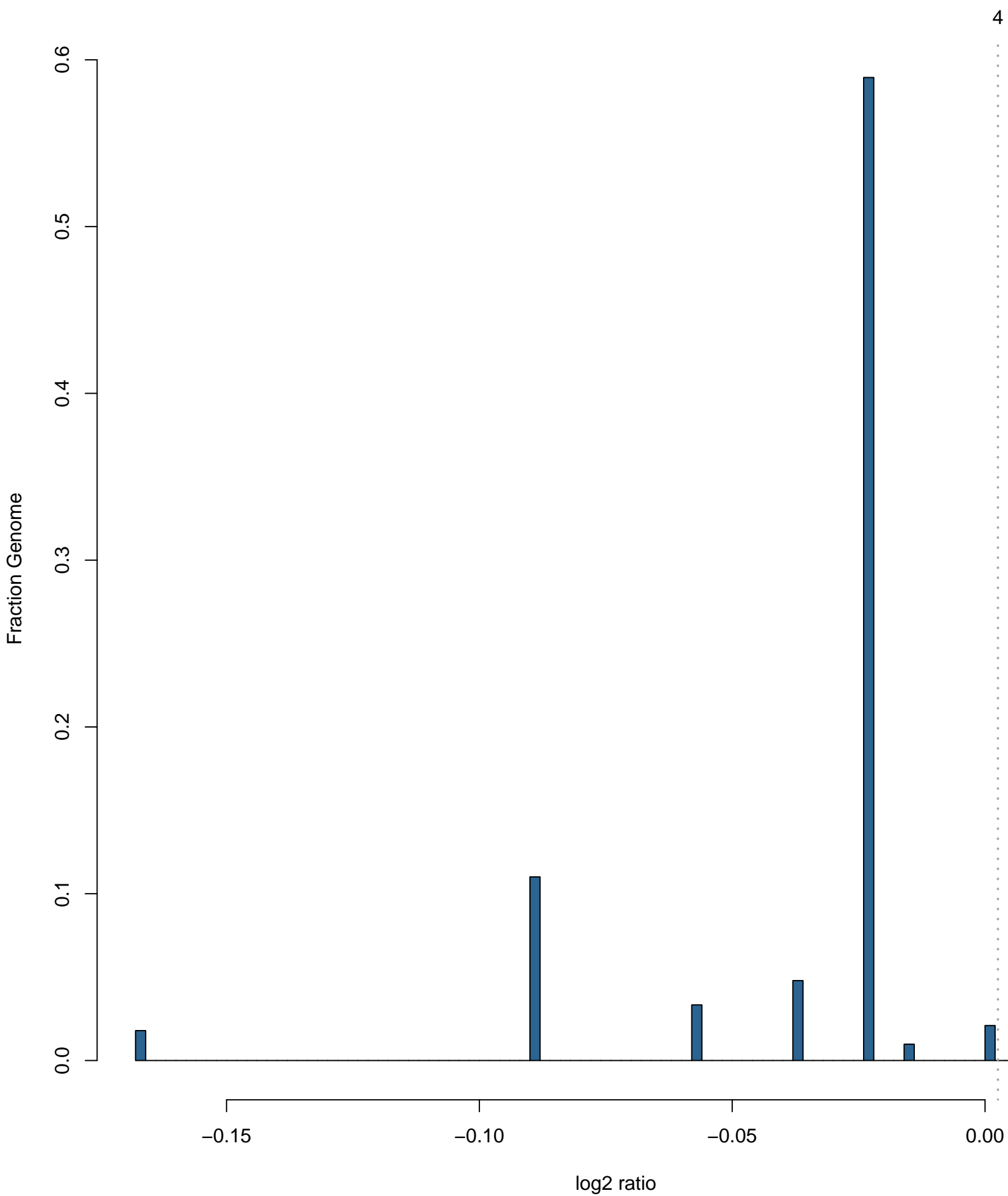


SCNA-fit log-likelihood: -13620

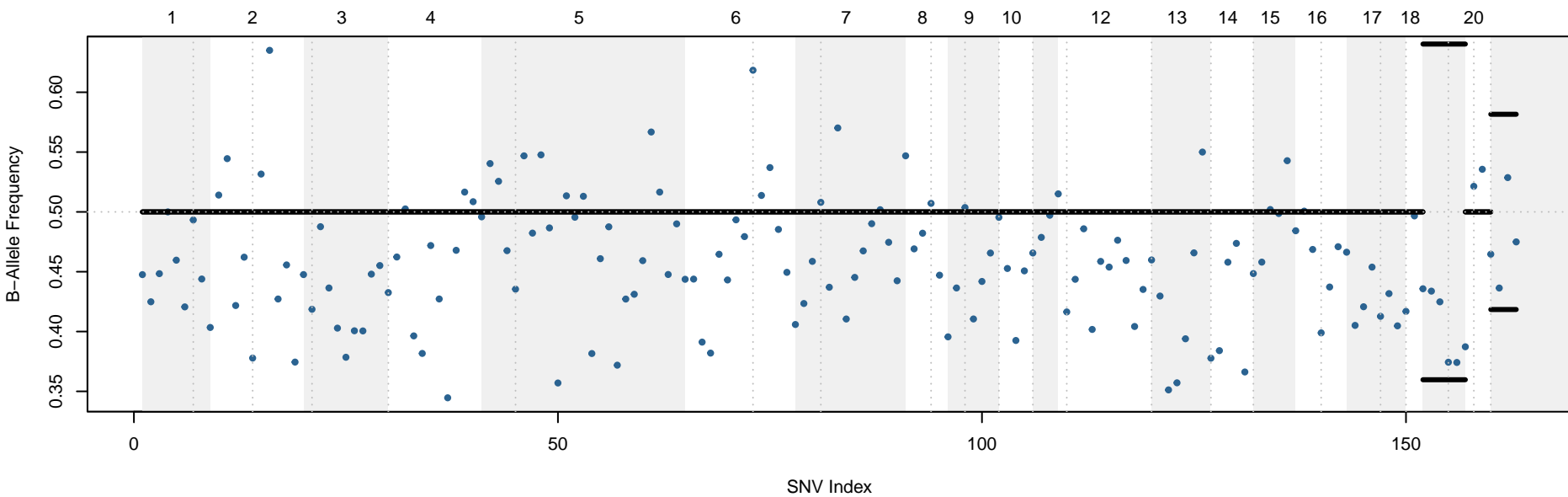




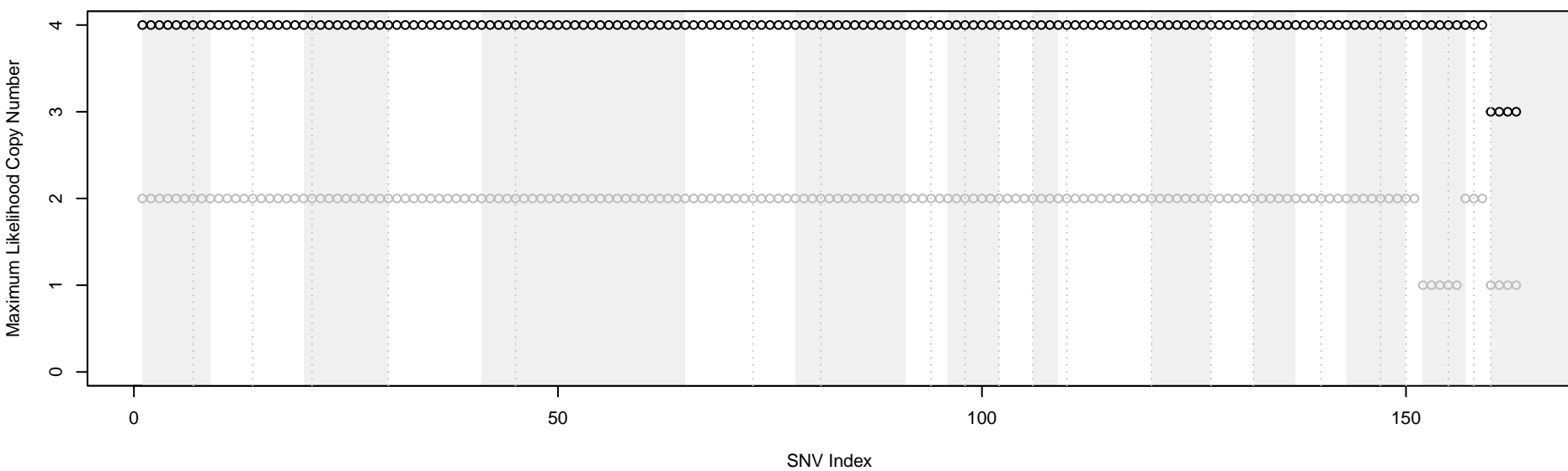
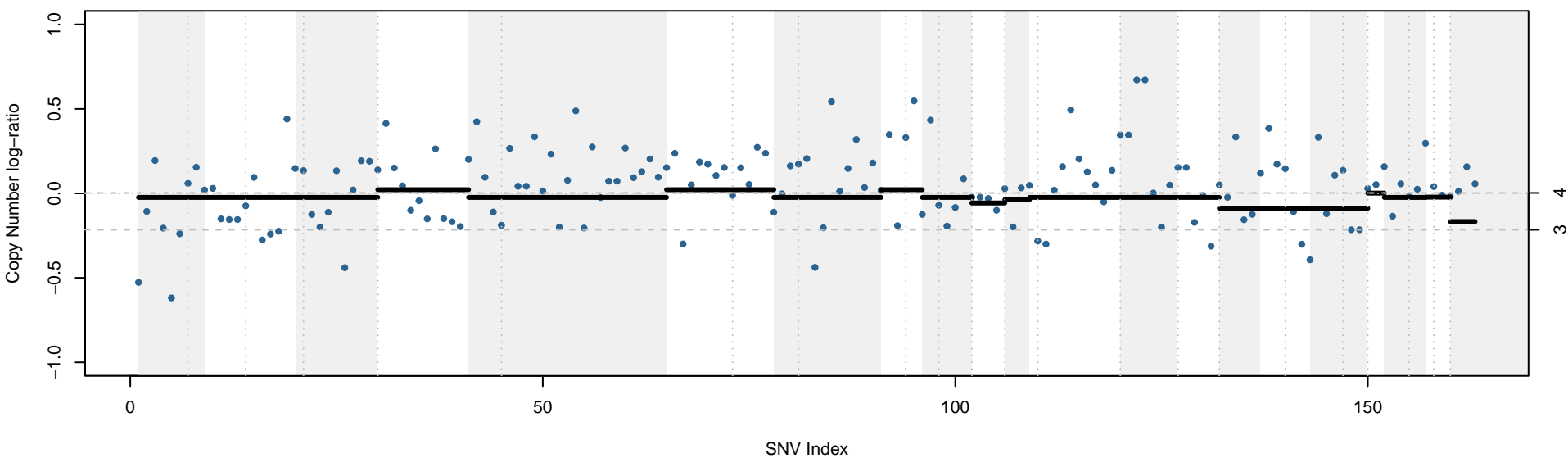
Purity: 0.39 Tumor ploidy: 3.987

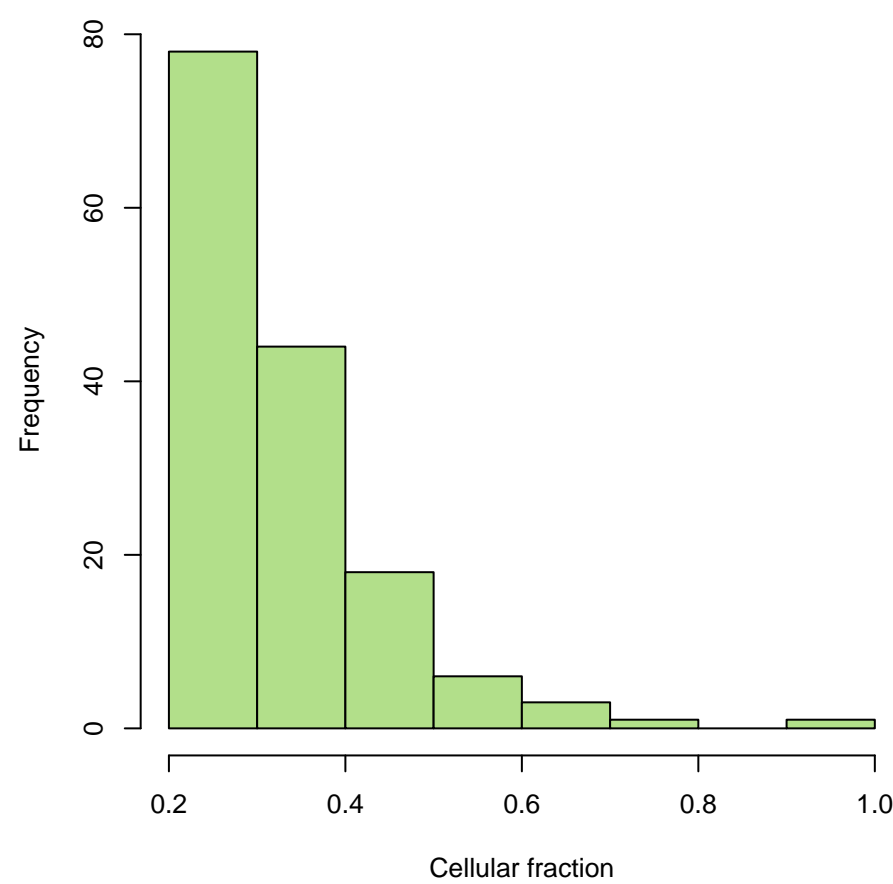
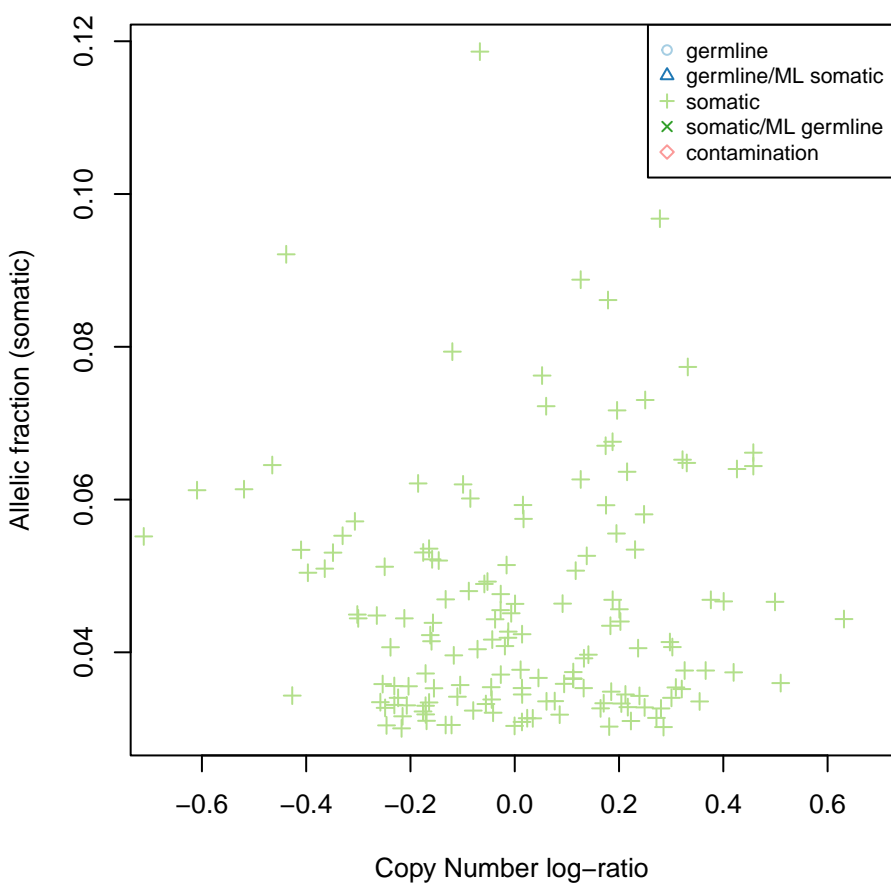
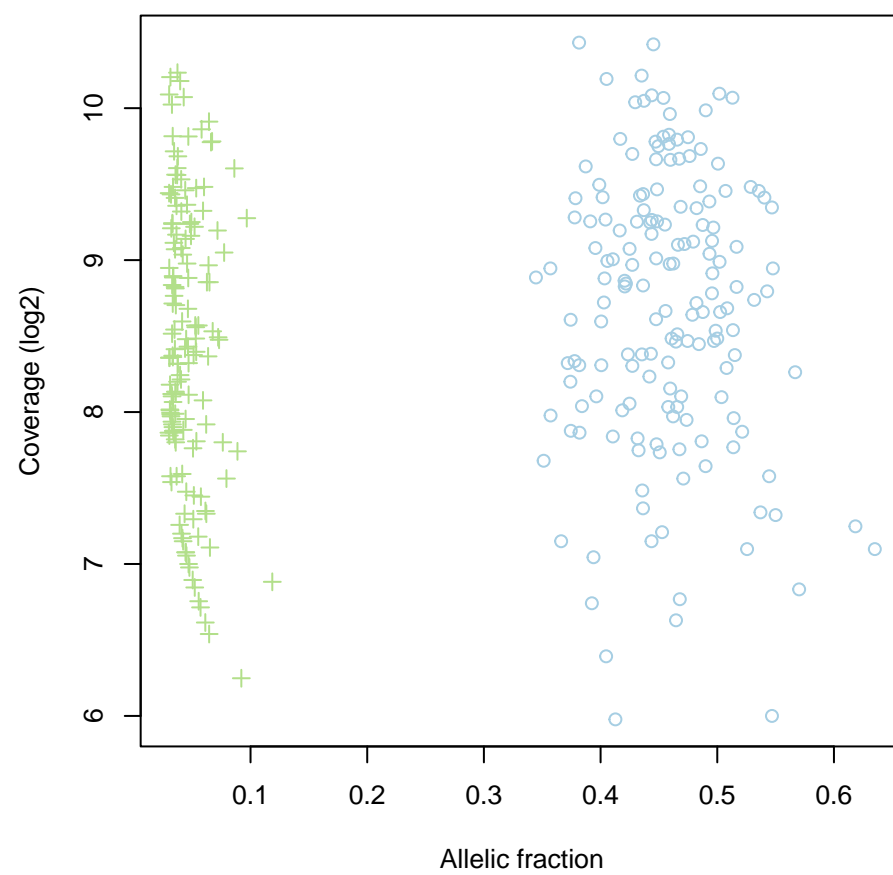
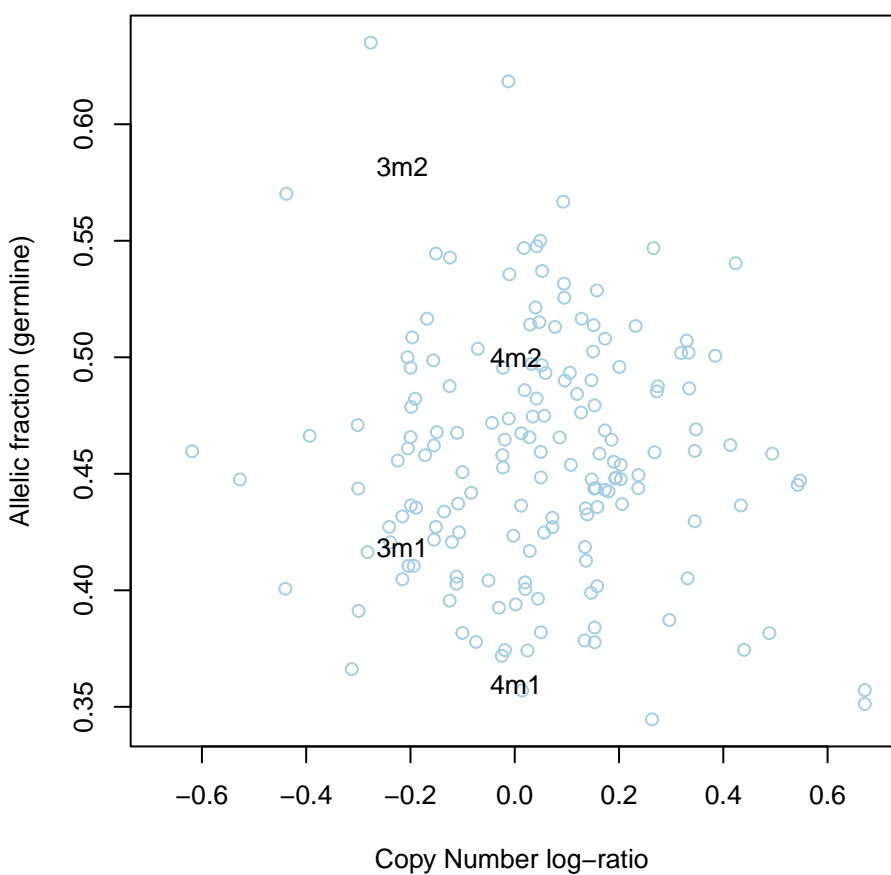


Purity: 0.39 Tumor ploidy: 3.987 SNV log-likelihood: -114.23 GoF: 85.1% Mean coverage: 299,455

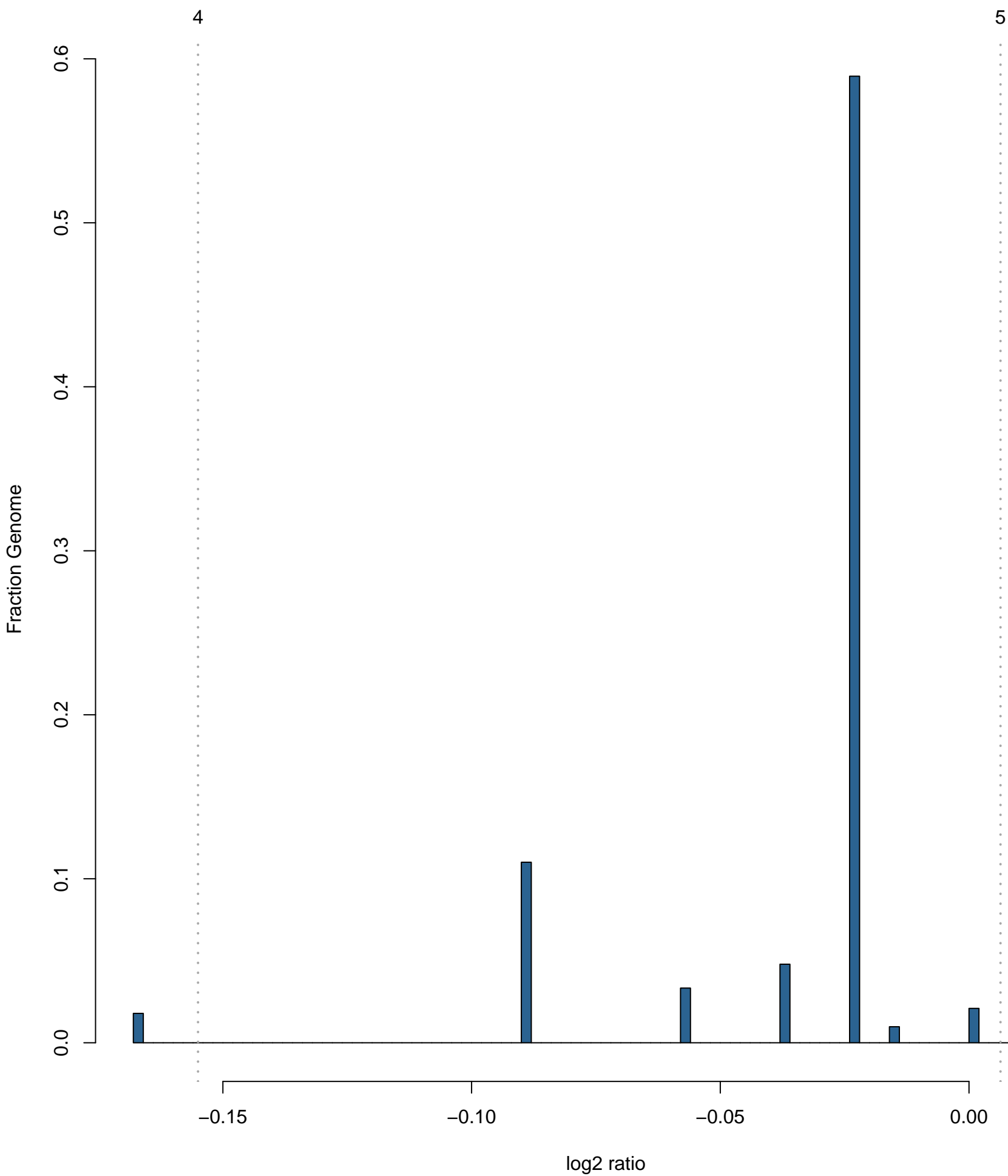


SCNA-fit log-likelihood: -13617.18



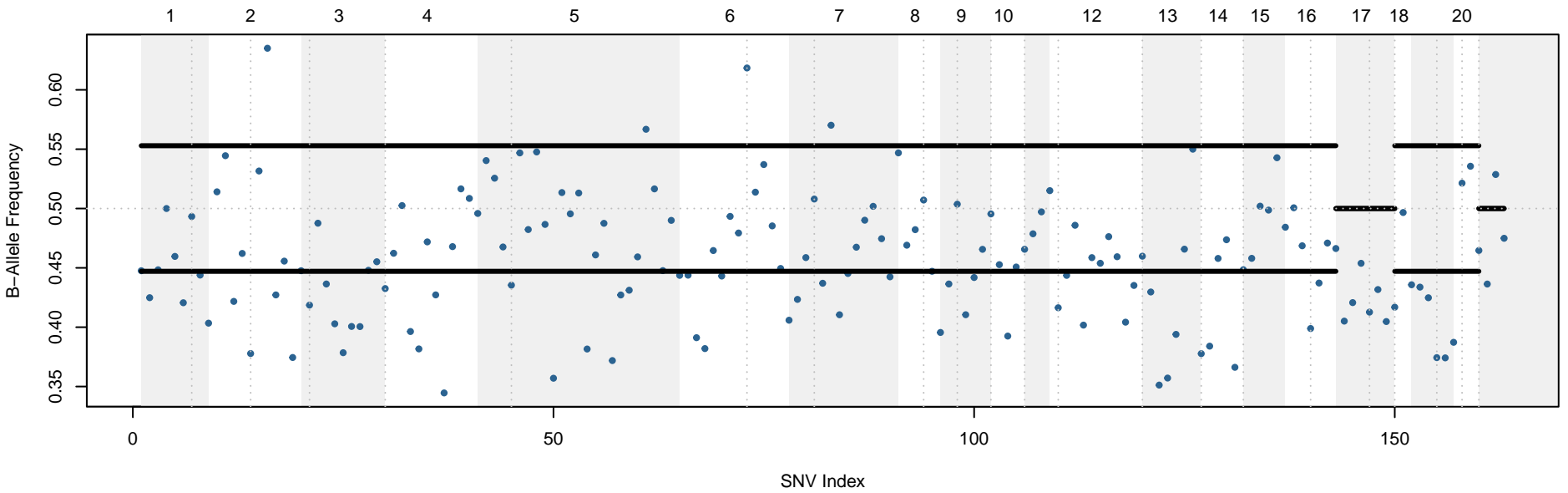


Purity: 0.31 Tumor ploidy: 4.959

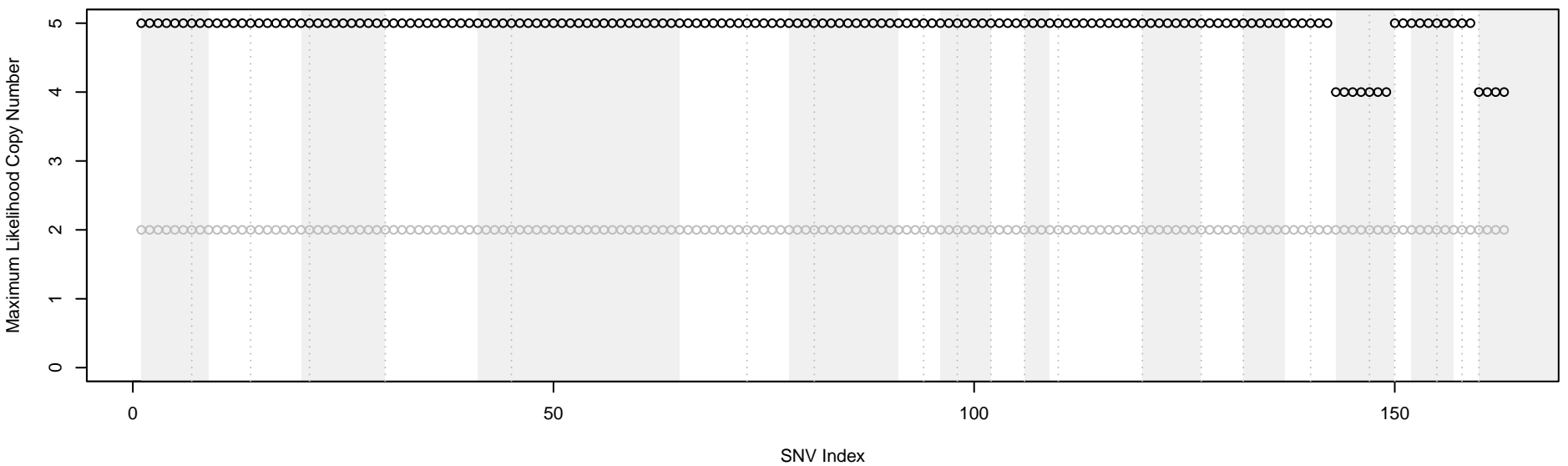
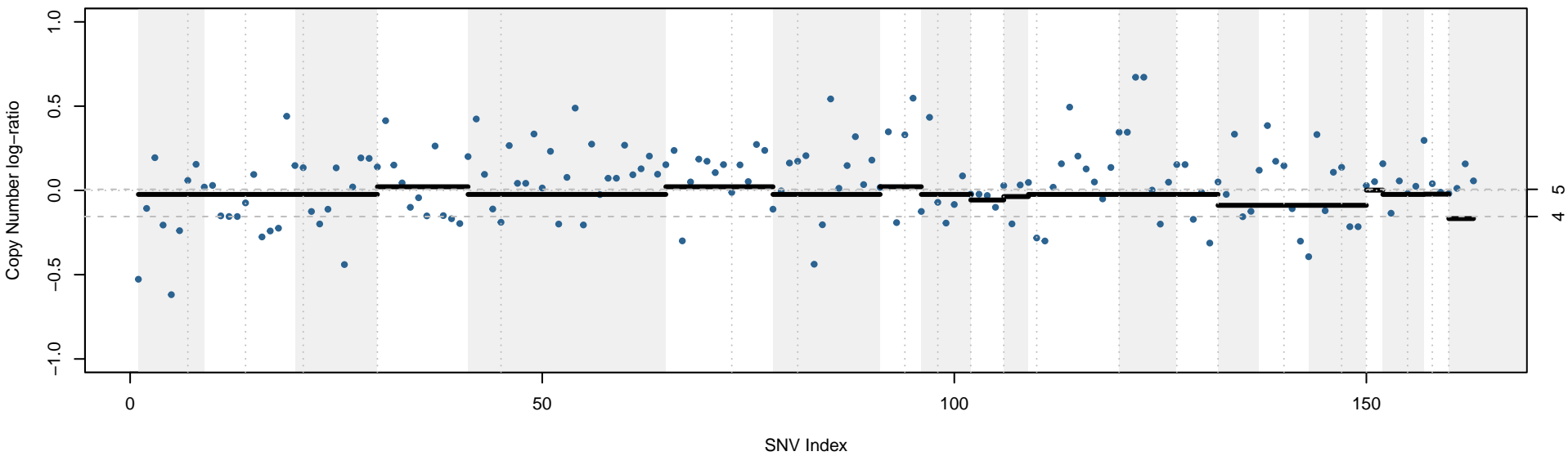


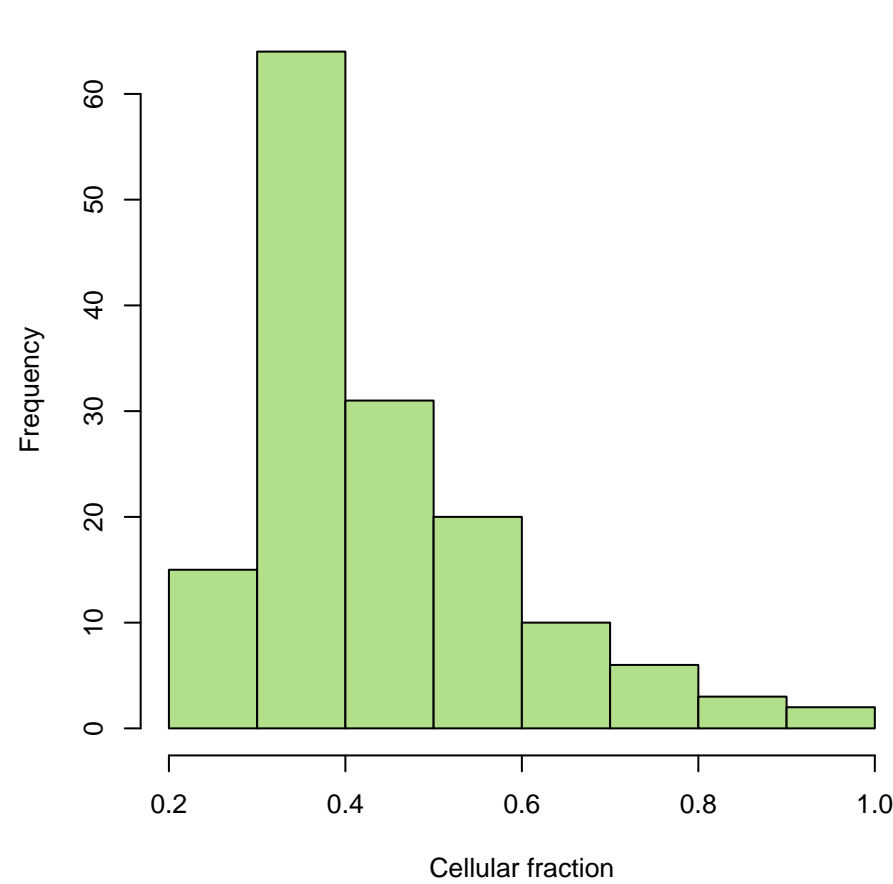
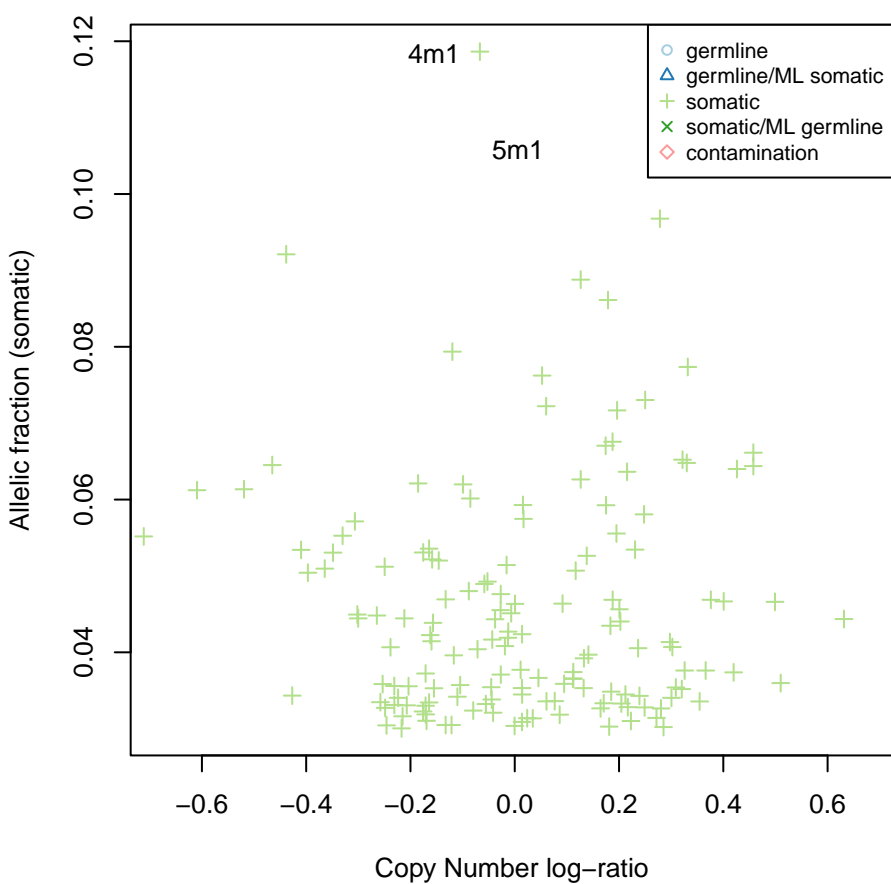
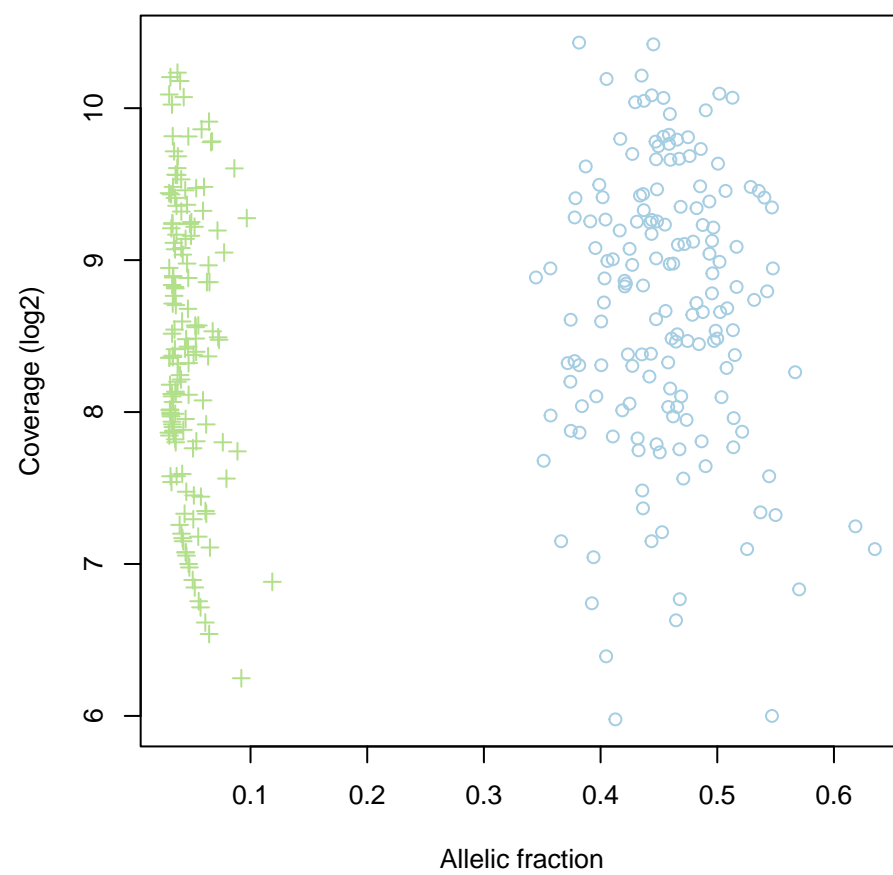
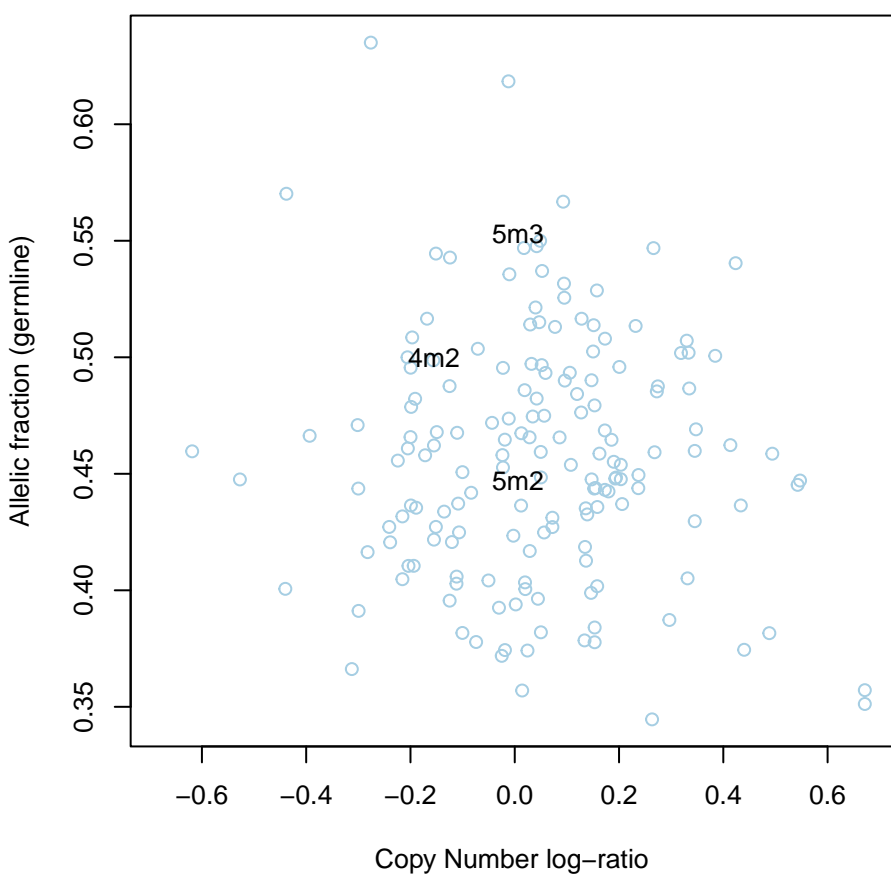


Purity: 0.31 Tumor ploidy: 4.959 SNV log-likelihood: -142.55 GoF: 93.8% Mean coverage: 299,455

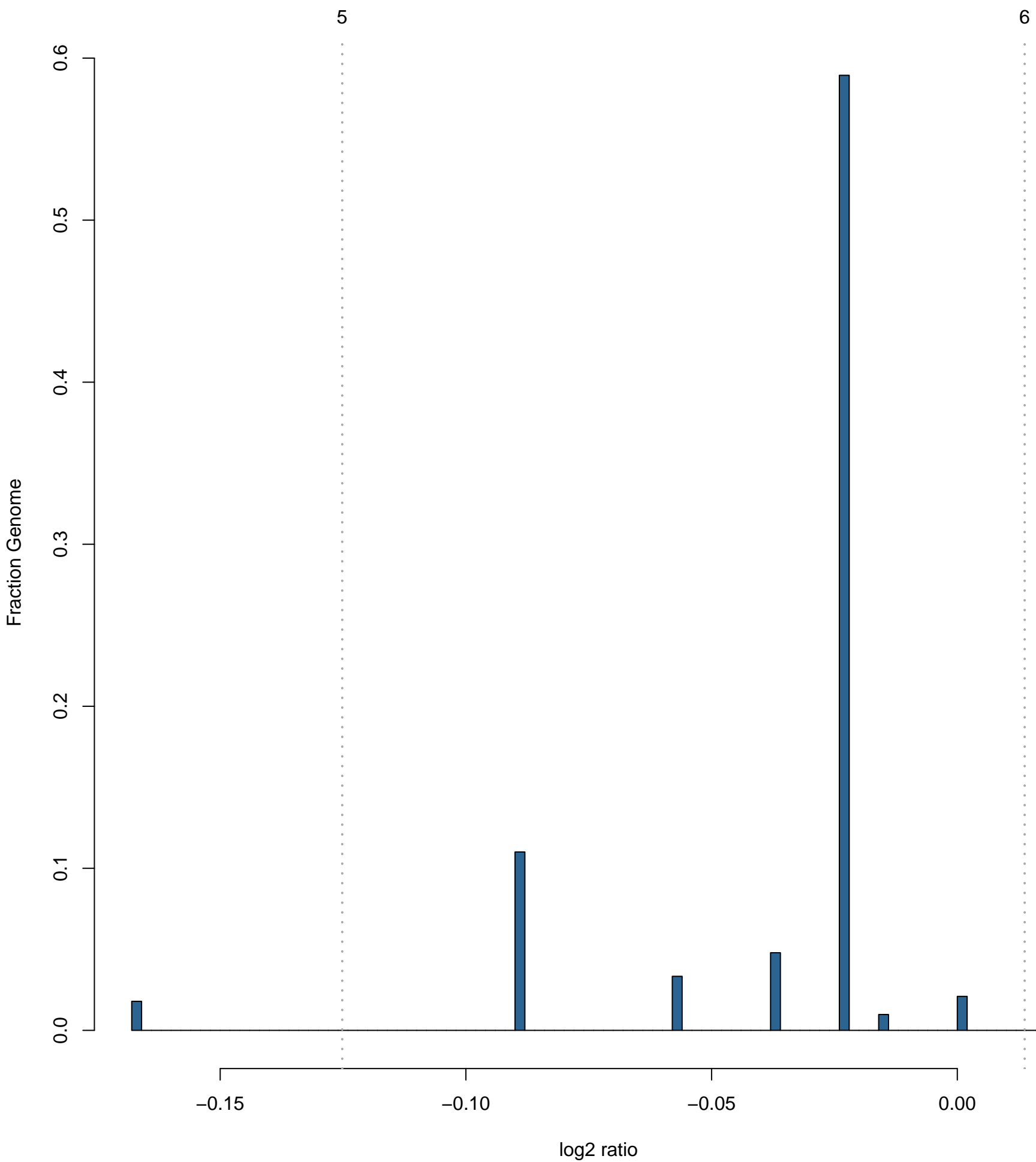


SCNA-fit log-likelihood: -13597.73

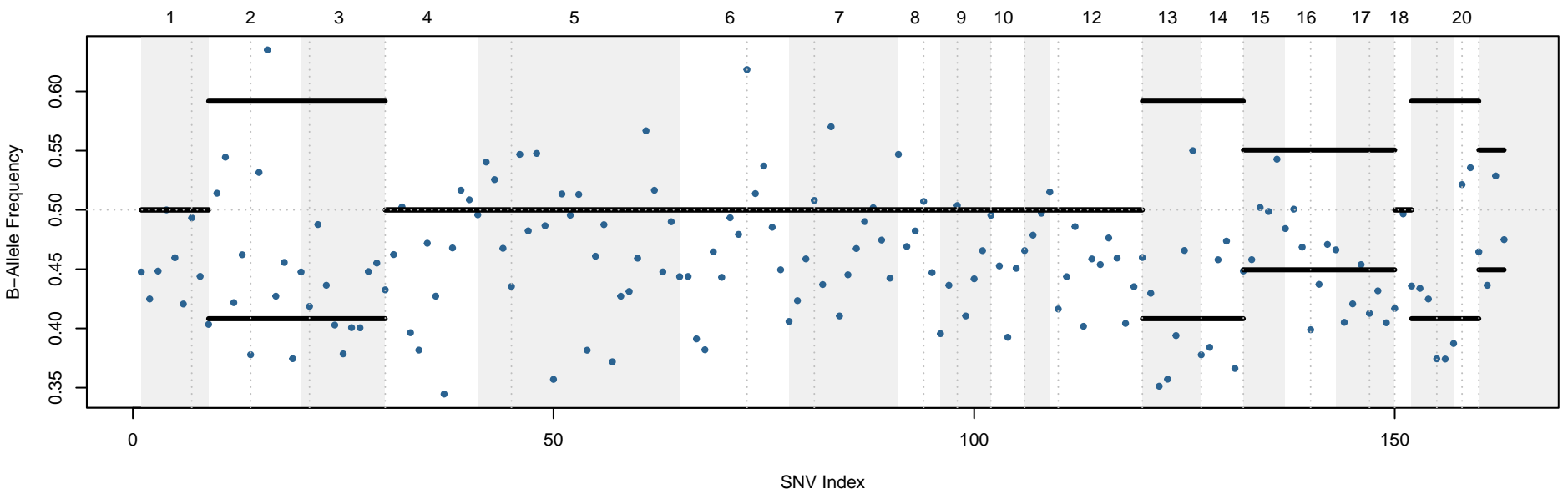




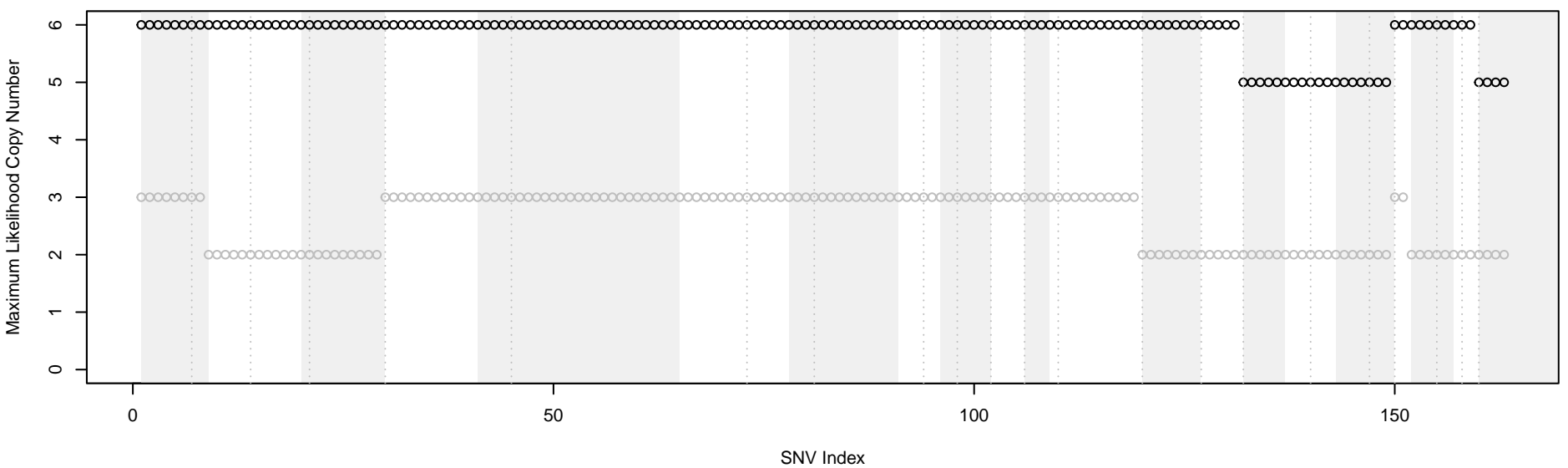
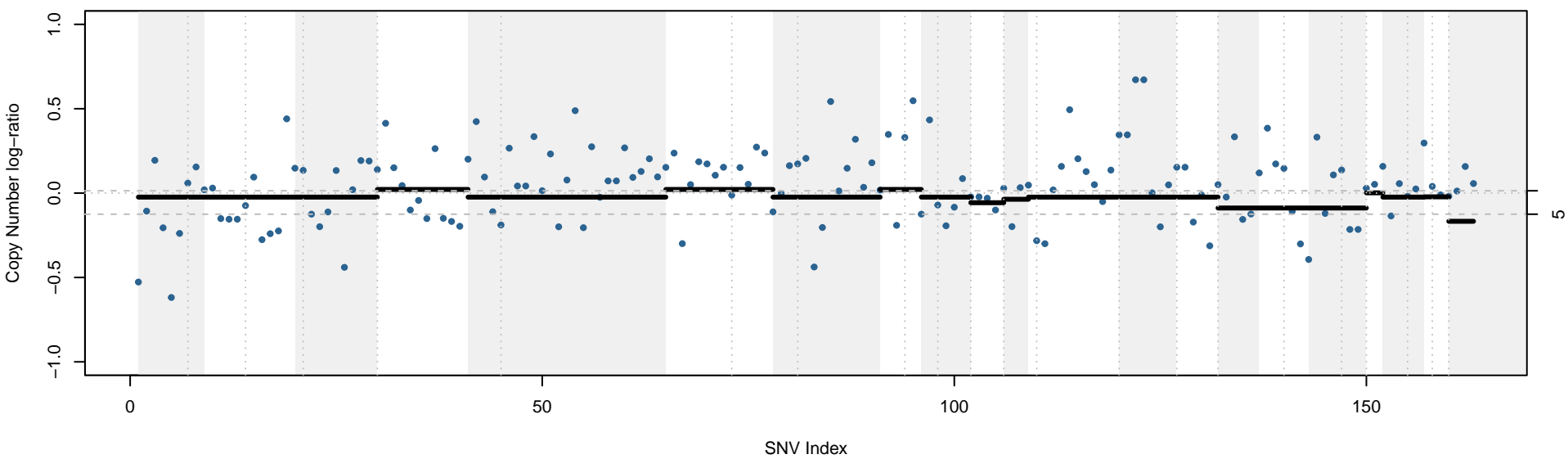
Purity: 0.29 Tumor ploidy: 5.897

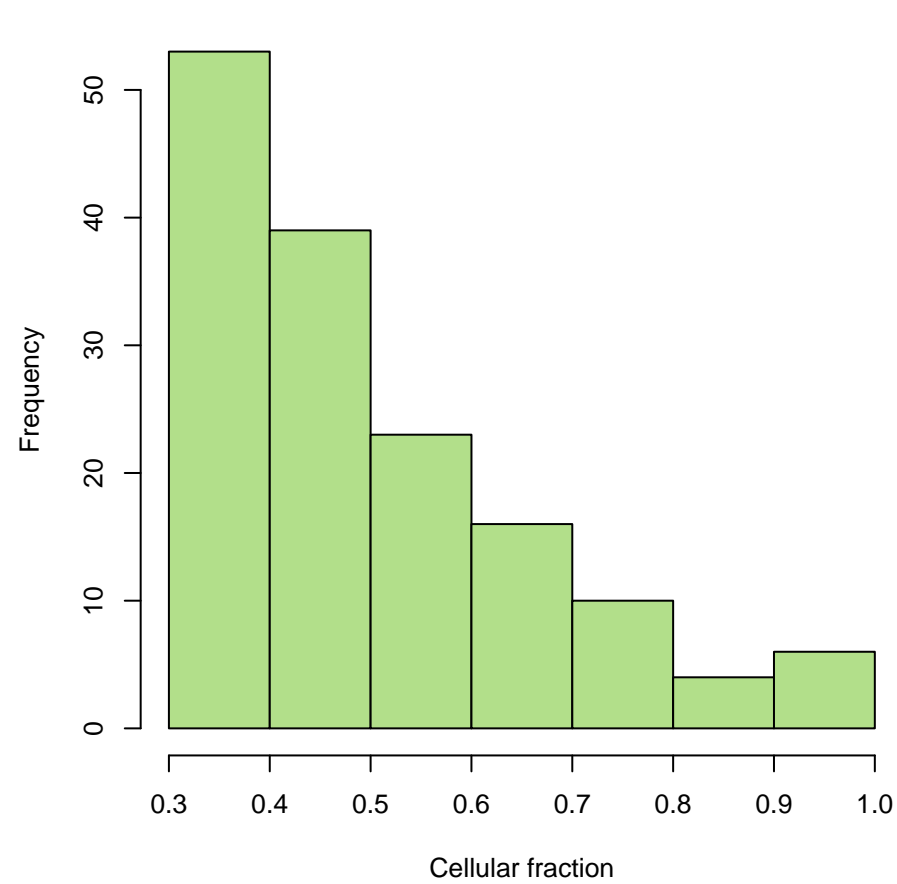
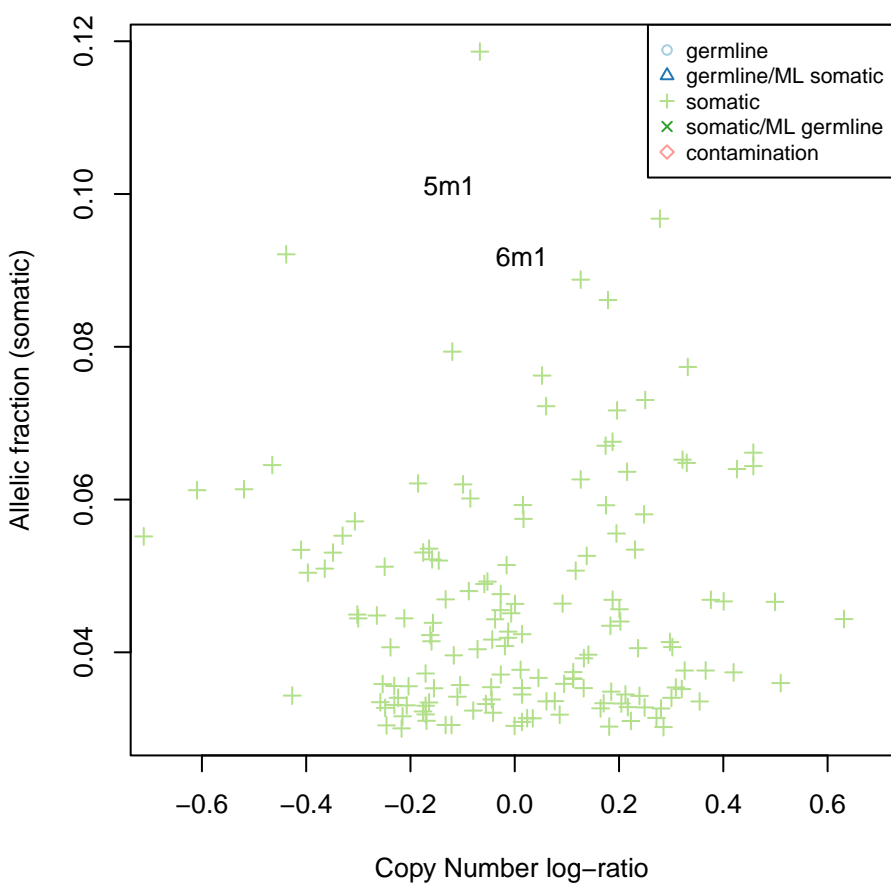
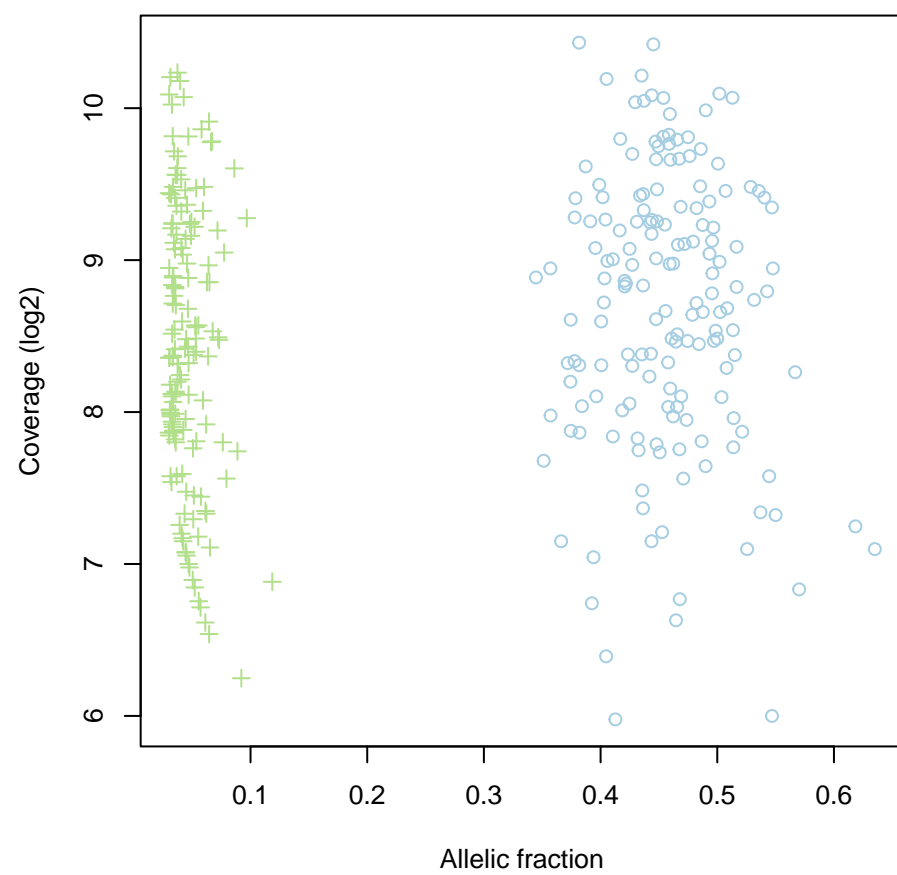
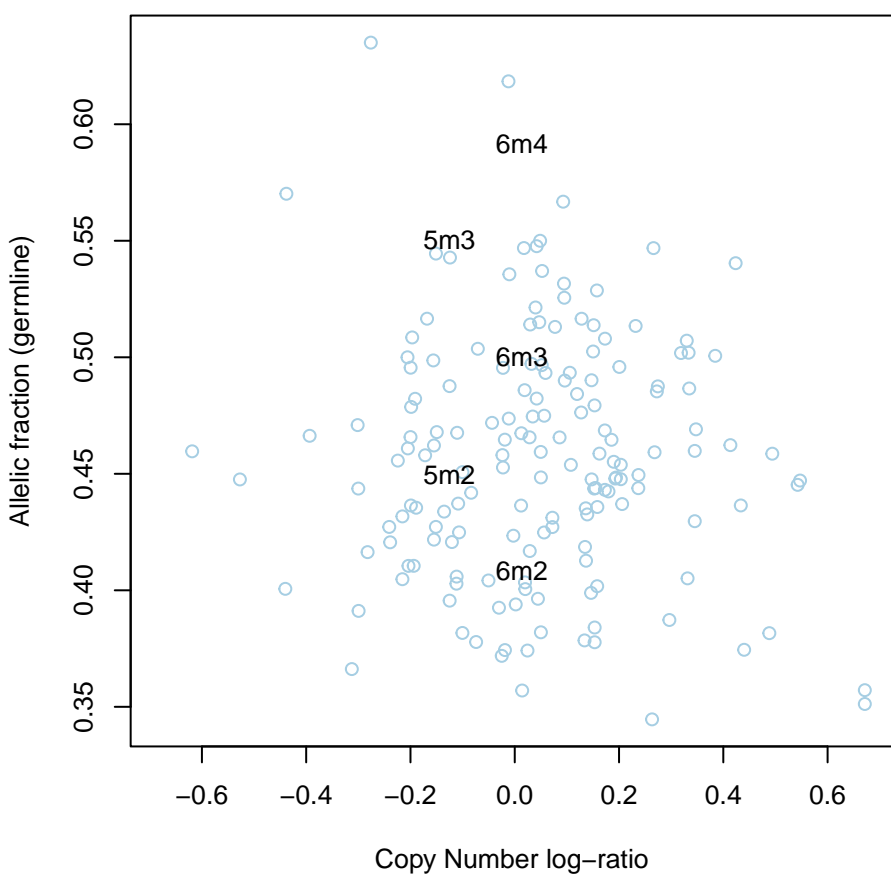


Purity: 0.29 Tumor ploidy: 5.897 SNV log-likelihood: -200.22 GoF: 94.5% Mean coverage: 299;455

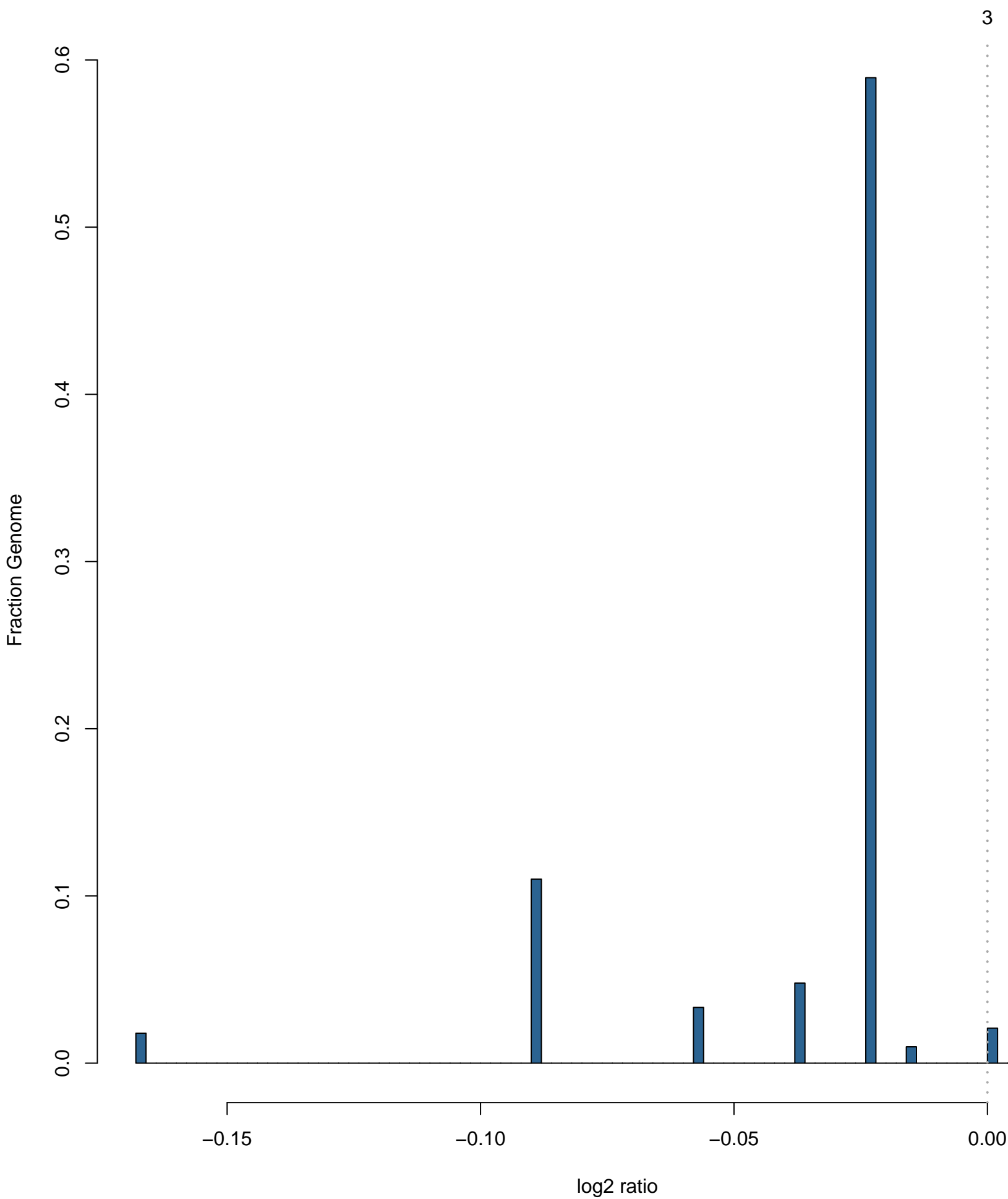


SCNA-fit log-likelihood: -13609.6

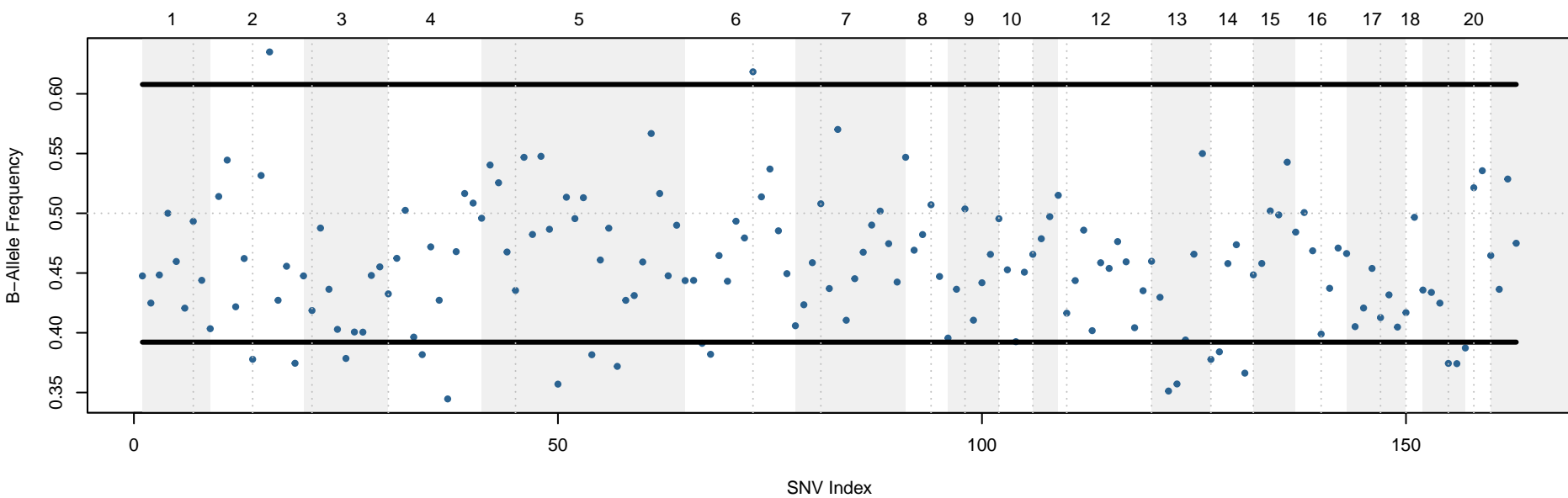




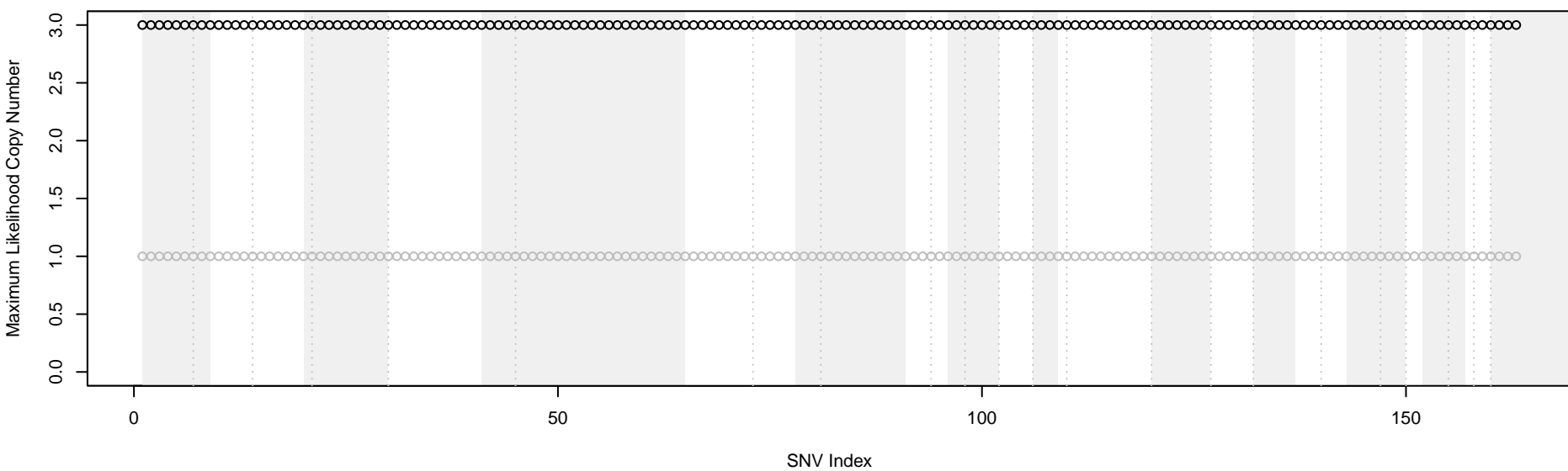
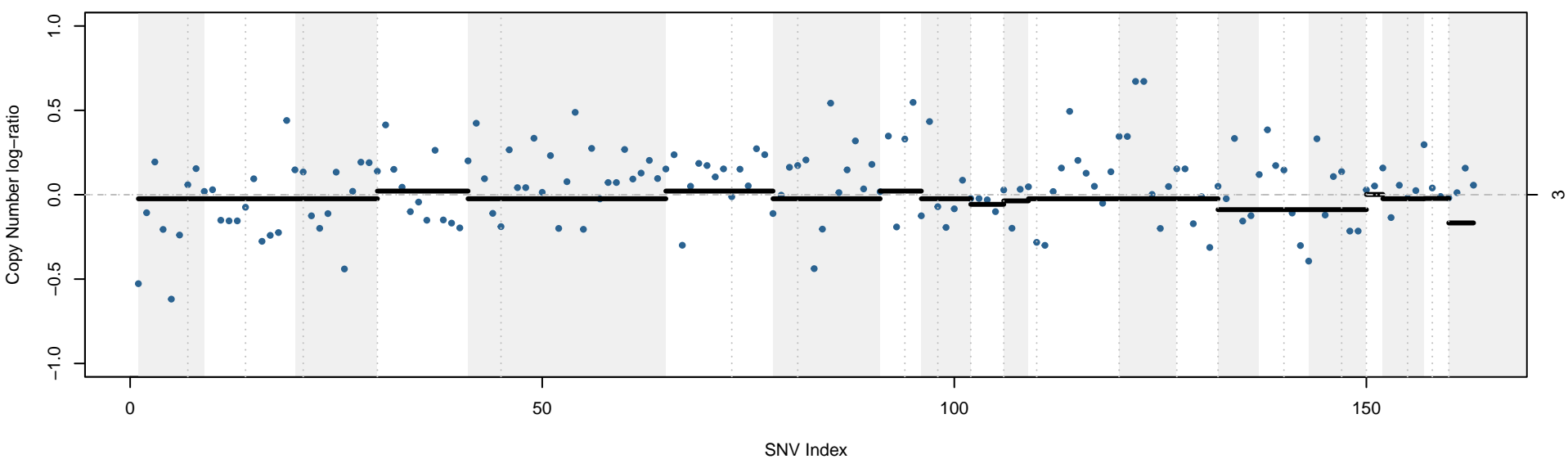
Purity: 0.55 Tumor ploidy: 3

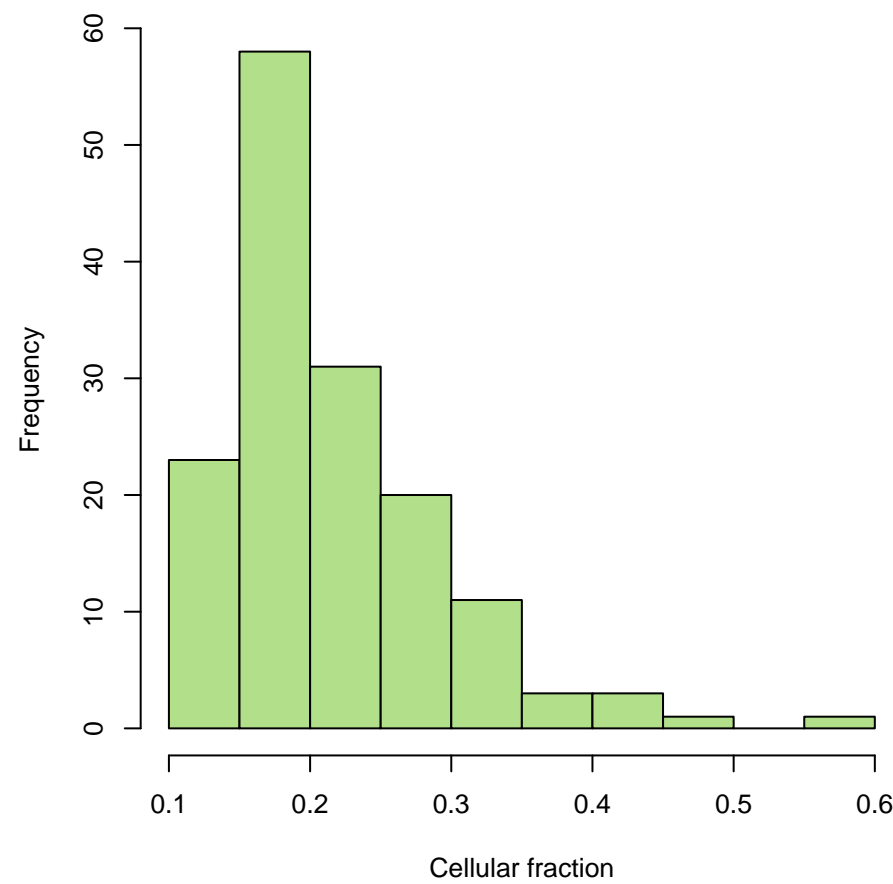
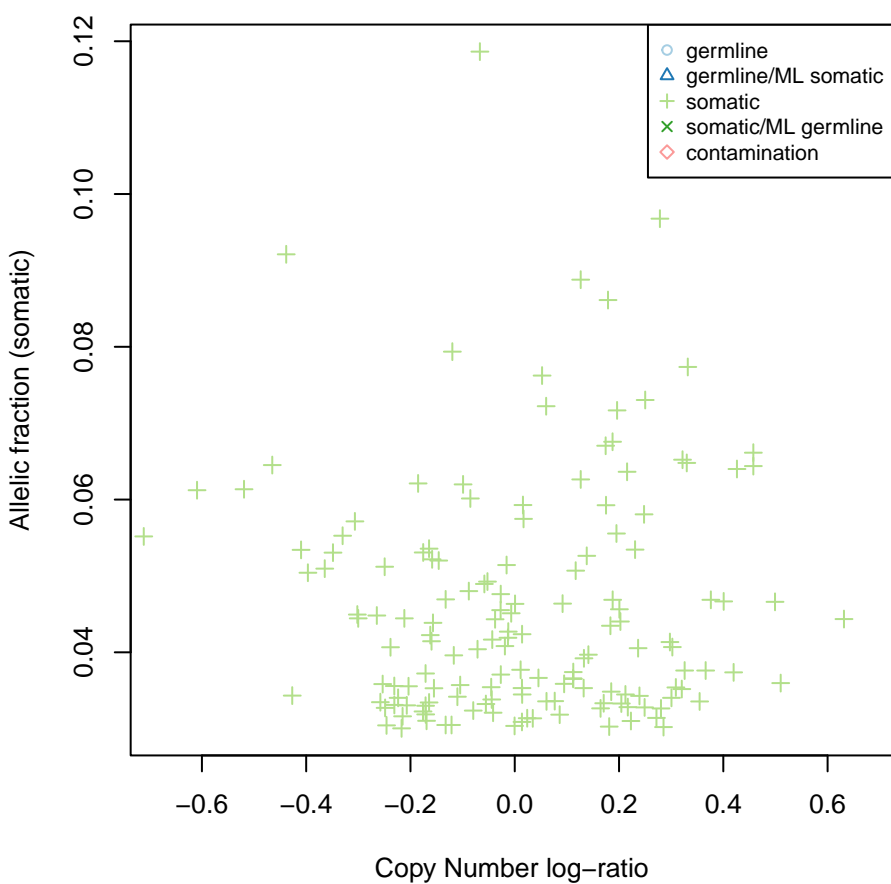
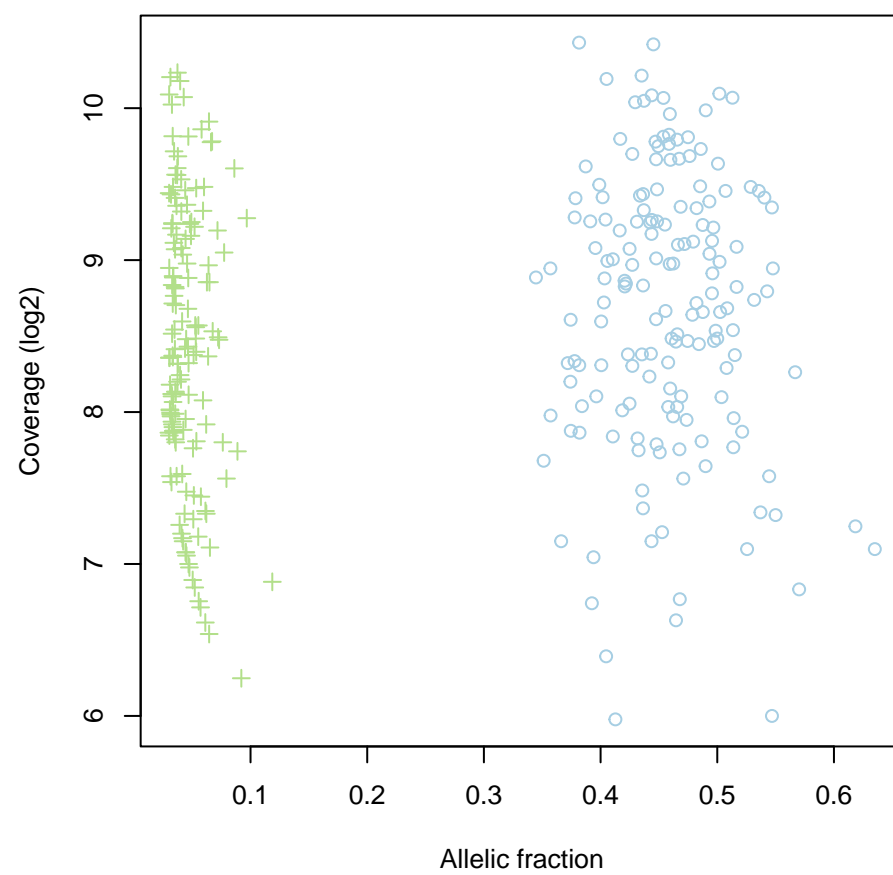
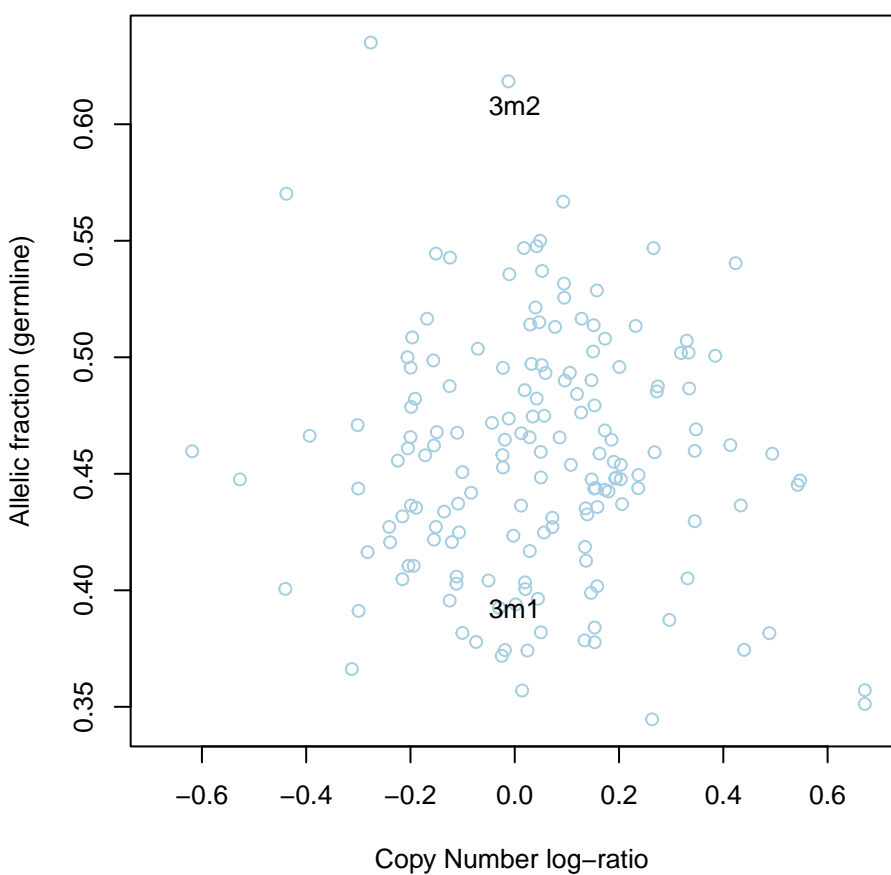


Purity: 0.55 Tumor ploidy: 3 SNV log-likelihood: -523.38 GoF: 59.6% Mean coverage: 299;455



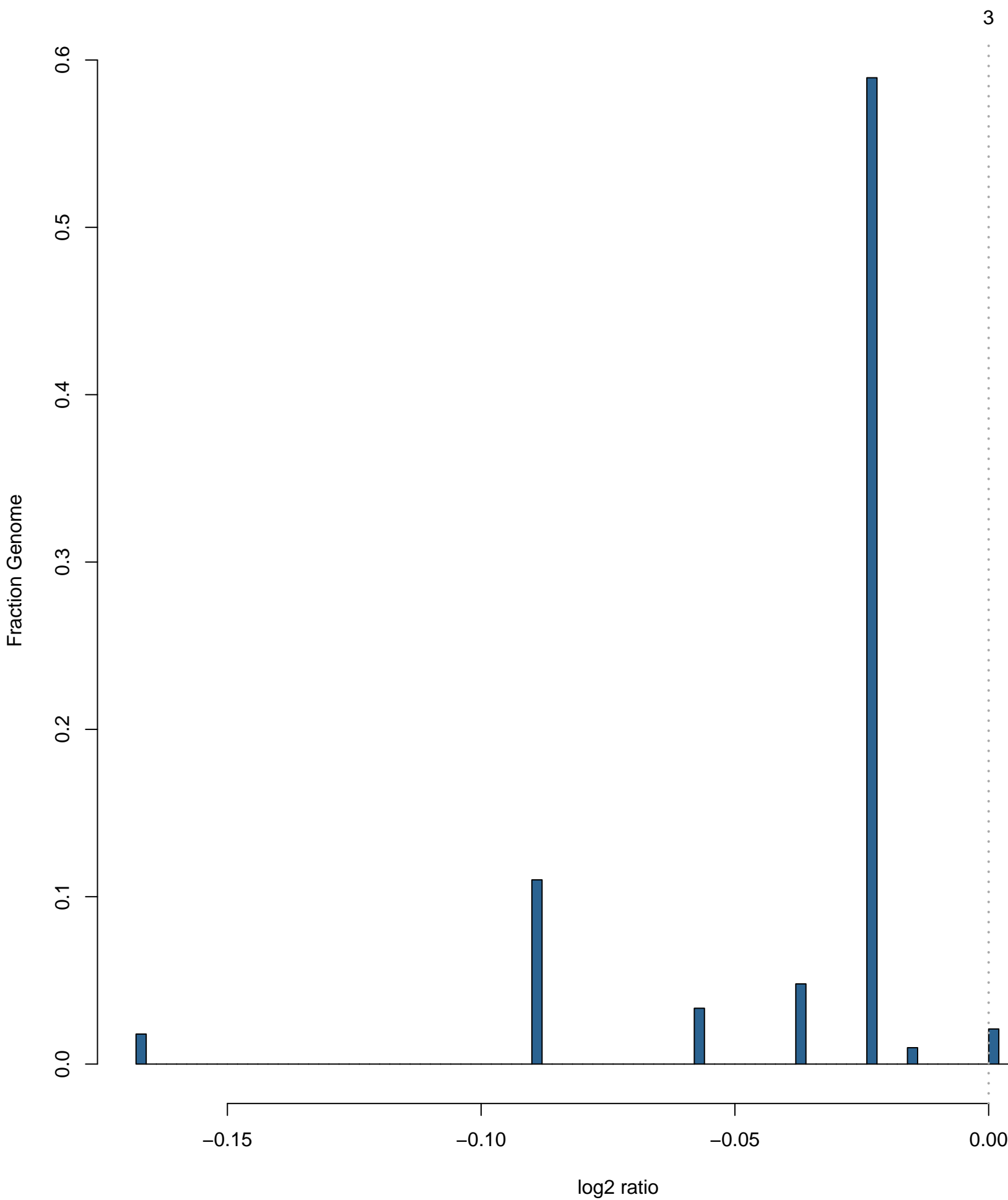
SCNA-fit log-likelihood: -13624.27



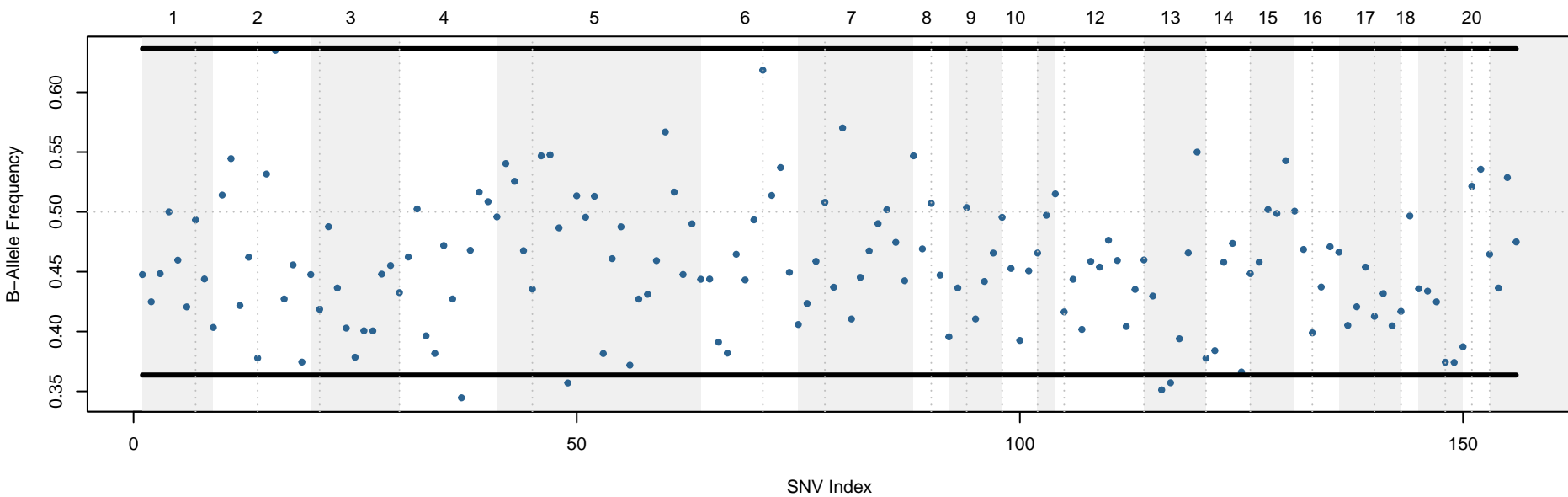




Purity: 0.75 Tumor ploidy: 3



Purity: 0.75 Tumor ploidy: 3 SNV log-likelihood: -1243.47 GoF: 28% Mean coverage: 299;455



SCNA-fit log-likelihood: -13620.35

