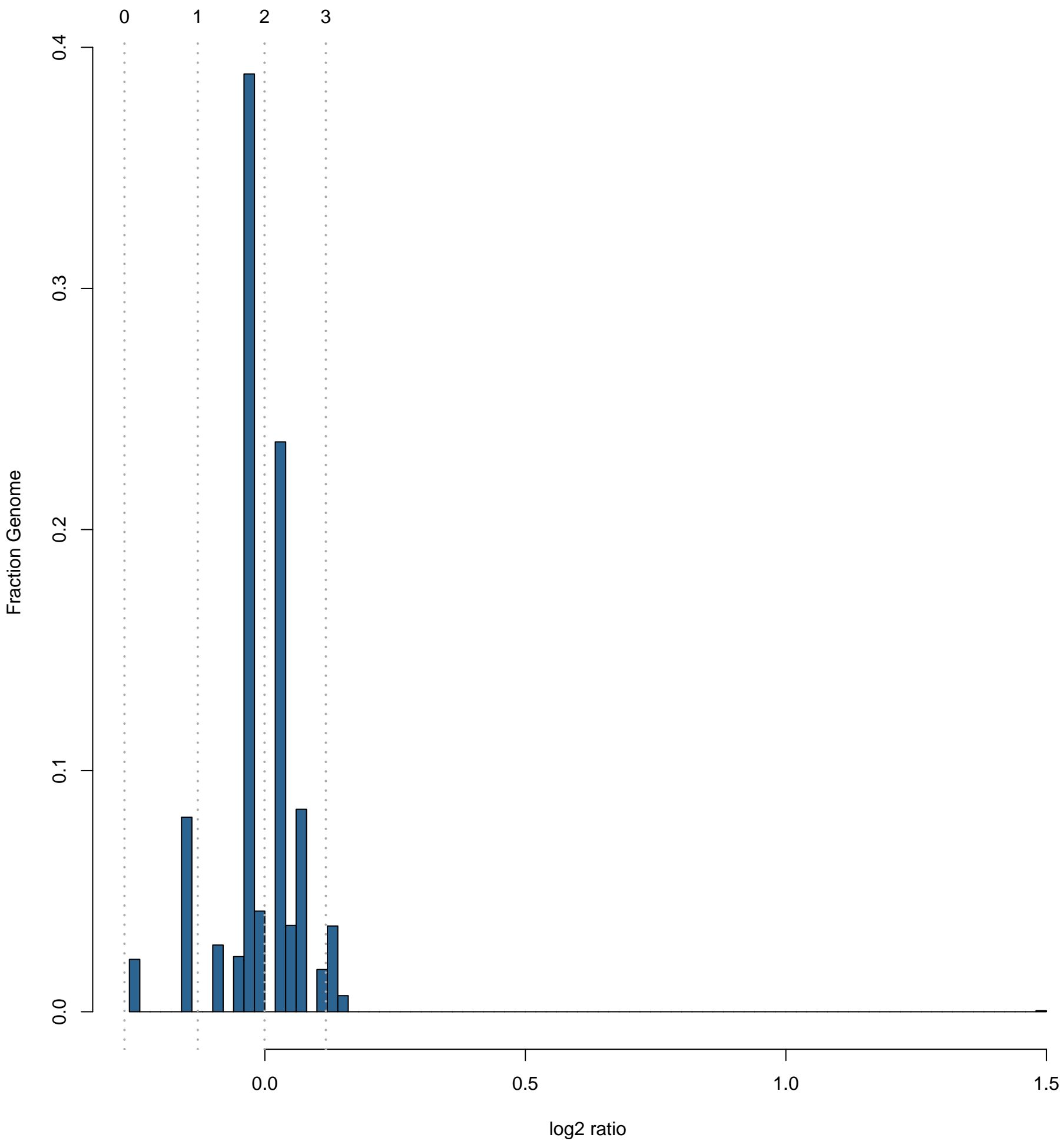
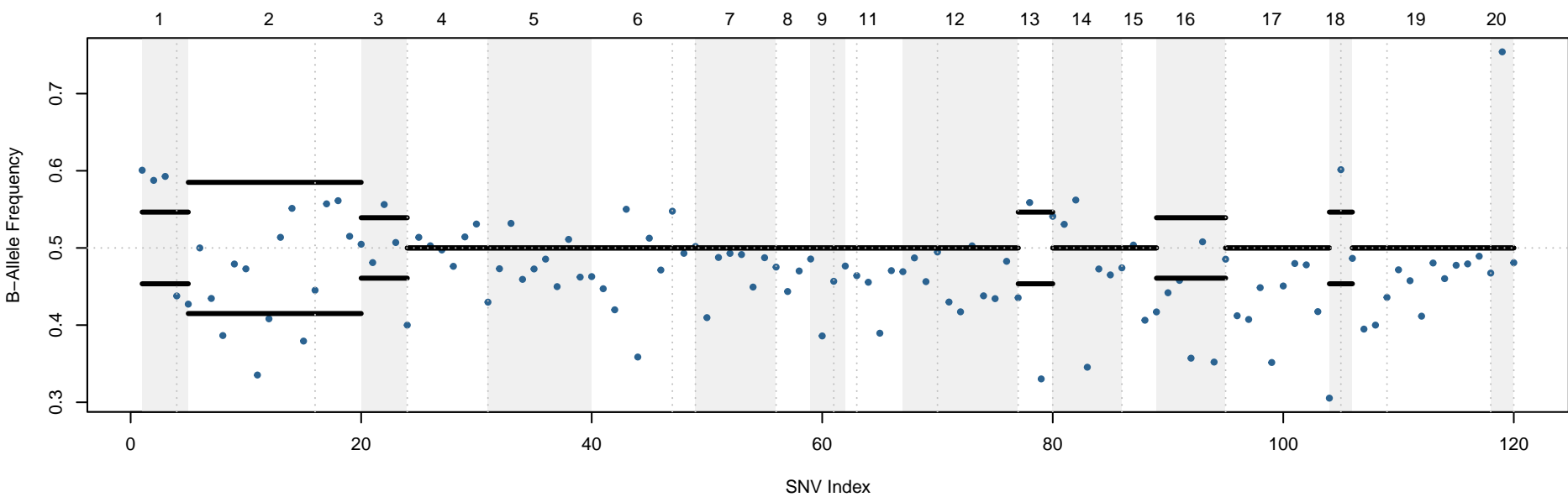


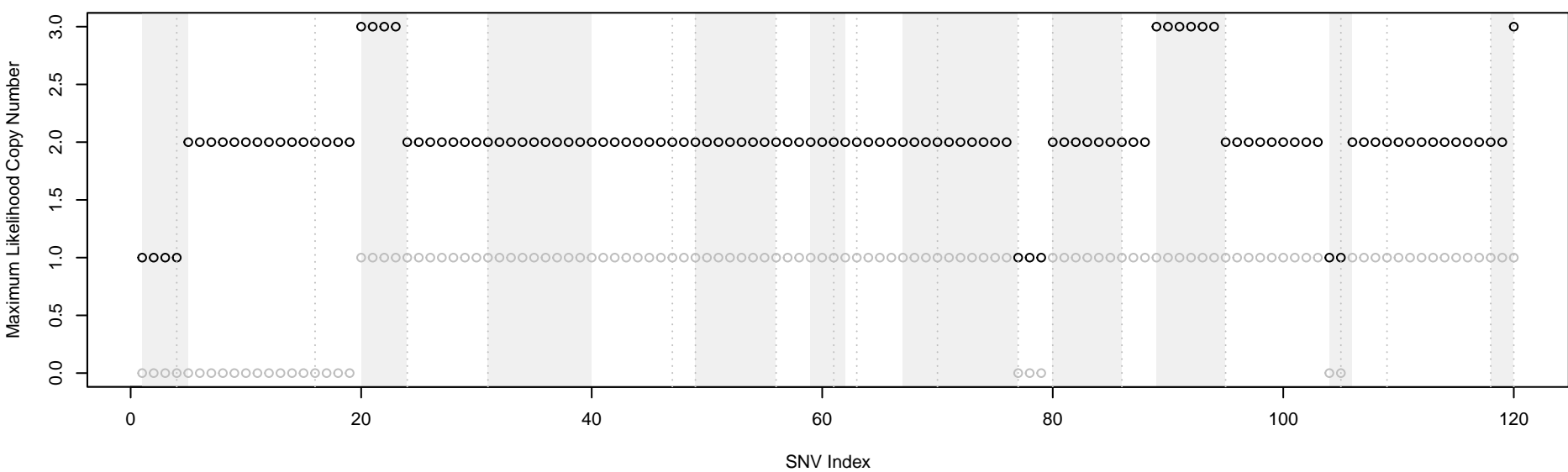
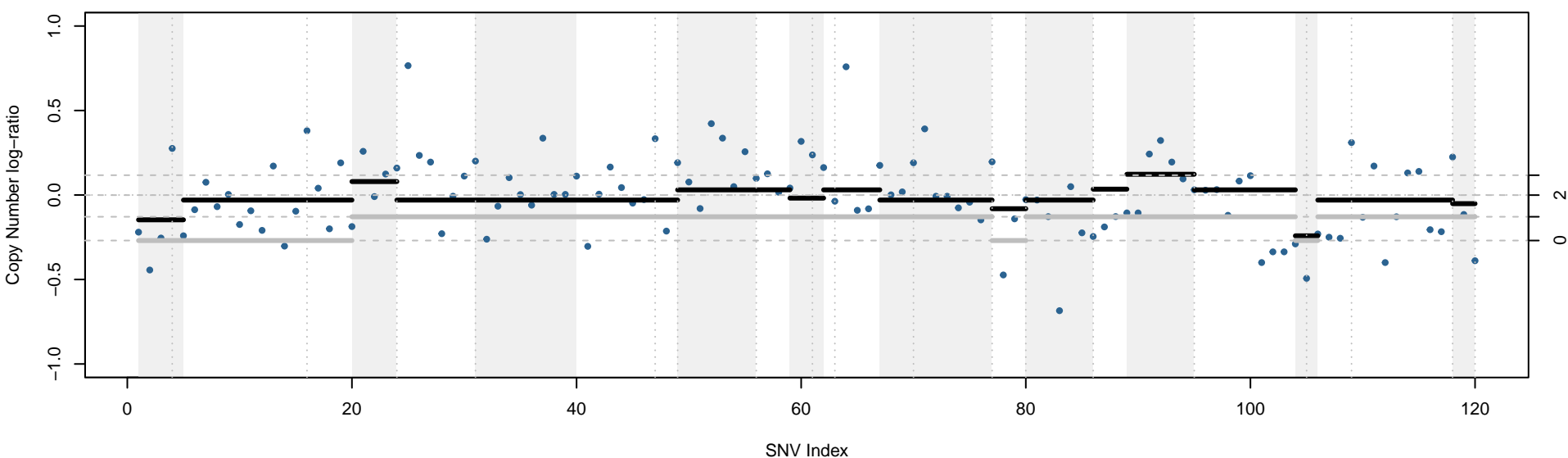
Purity: 0.17 Tumor ploidy: 2.005

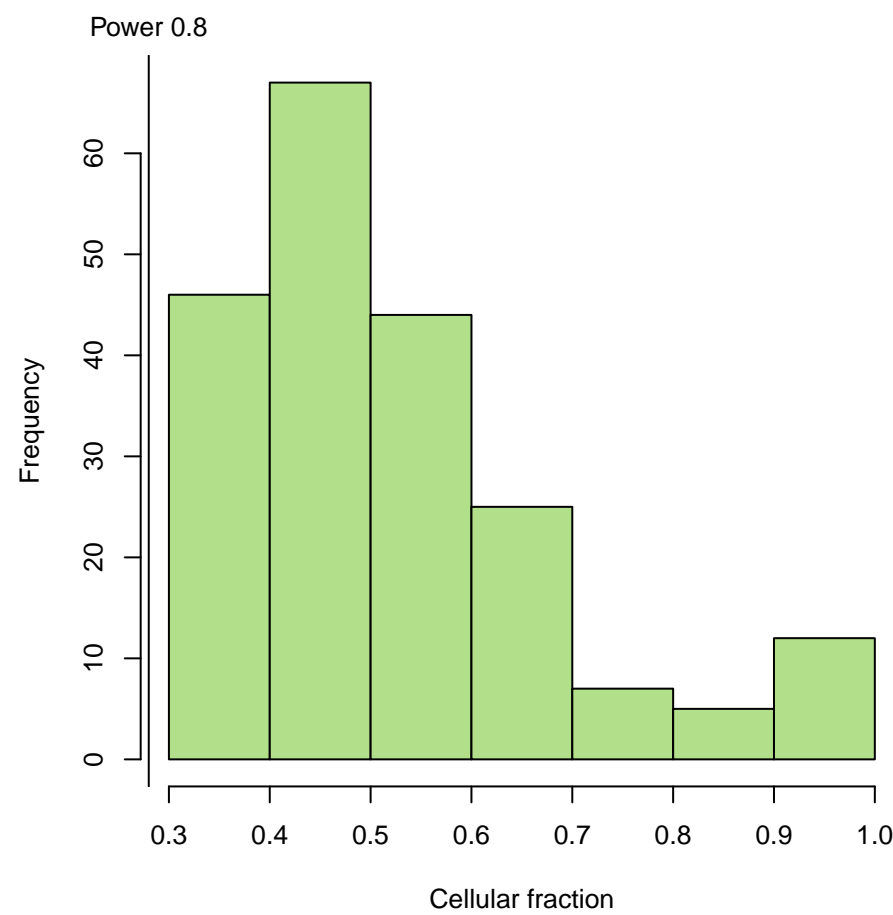
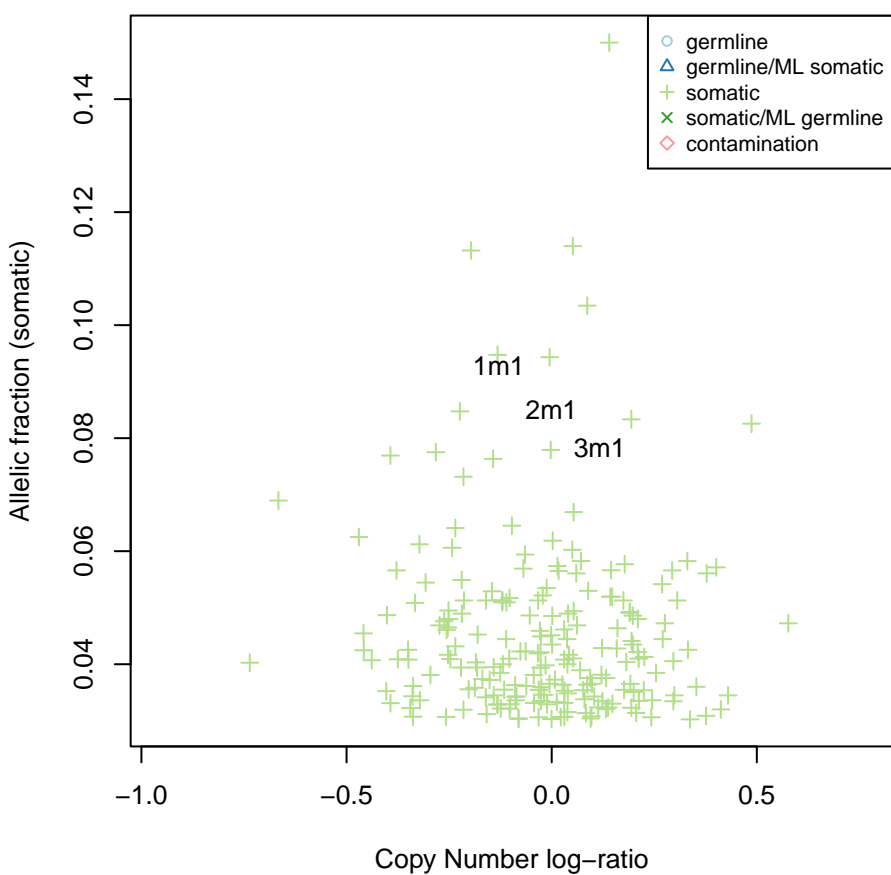
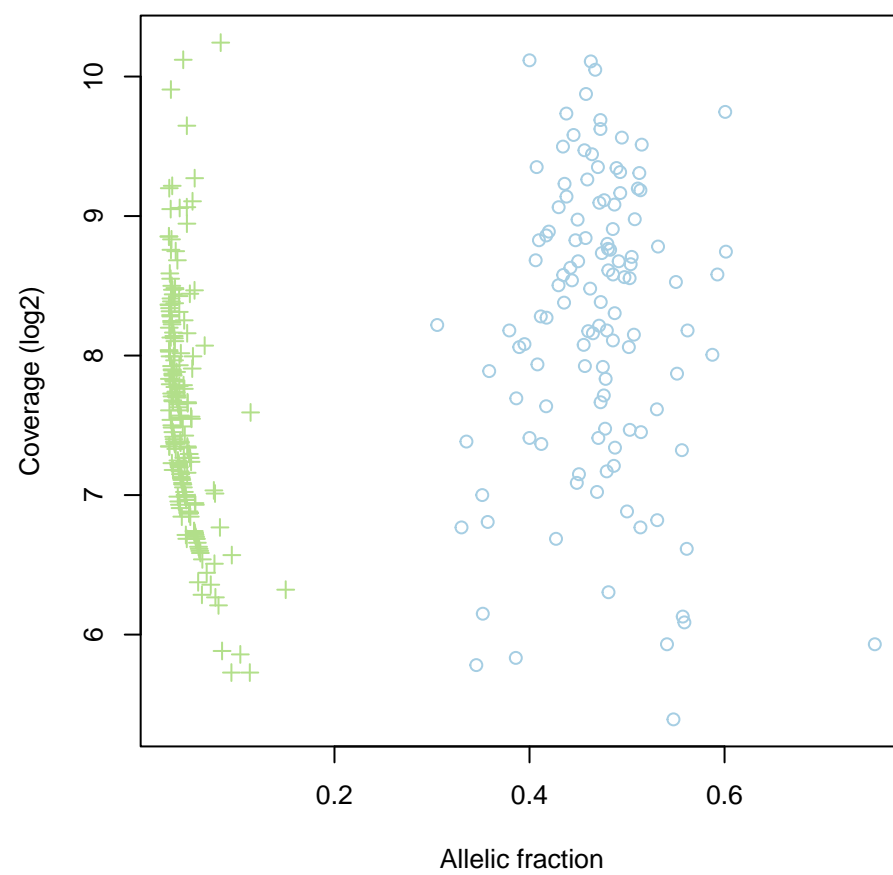
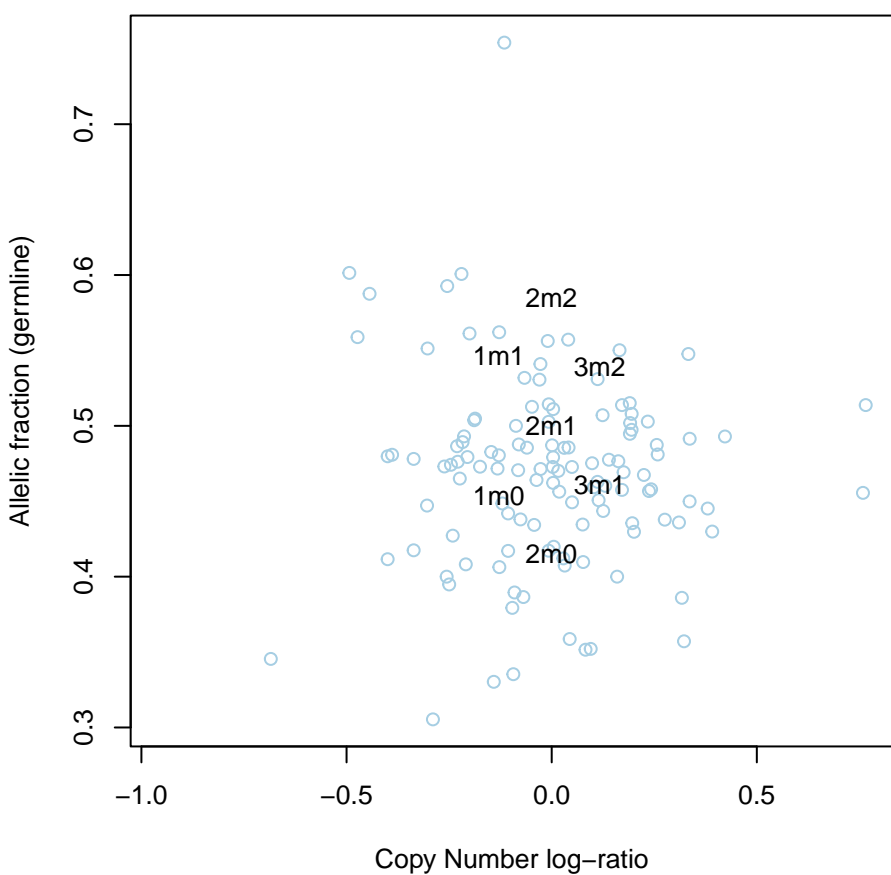


Purity: 0.17 Tumor ploidy: 2.005 SNV log-likelihood: 158.74 GoF: 94.3% Mean coverage: 406;281

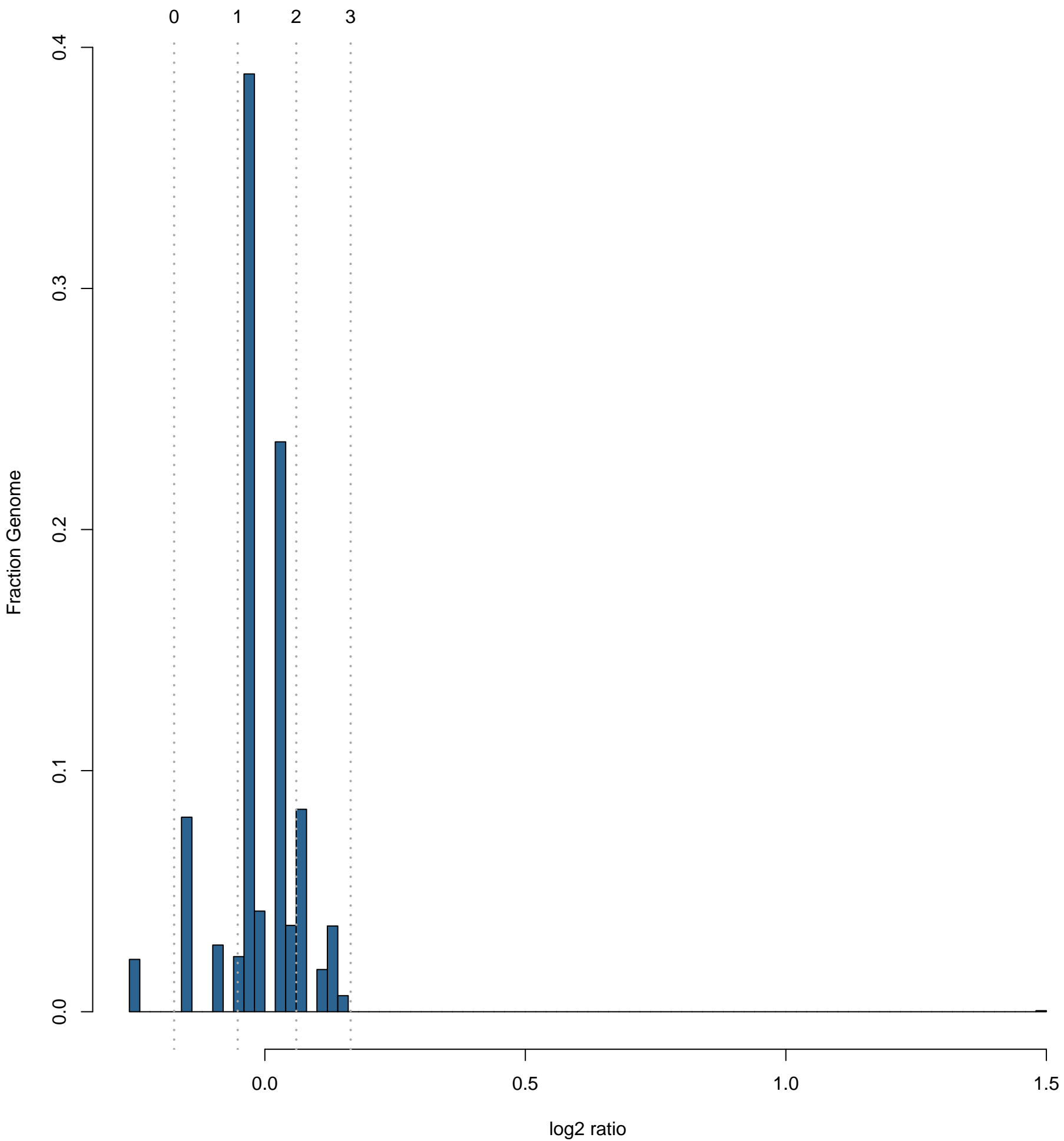


SCNA-fit log-likelihood: -15960.35

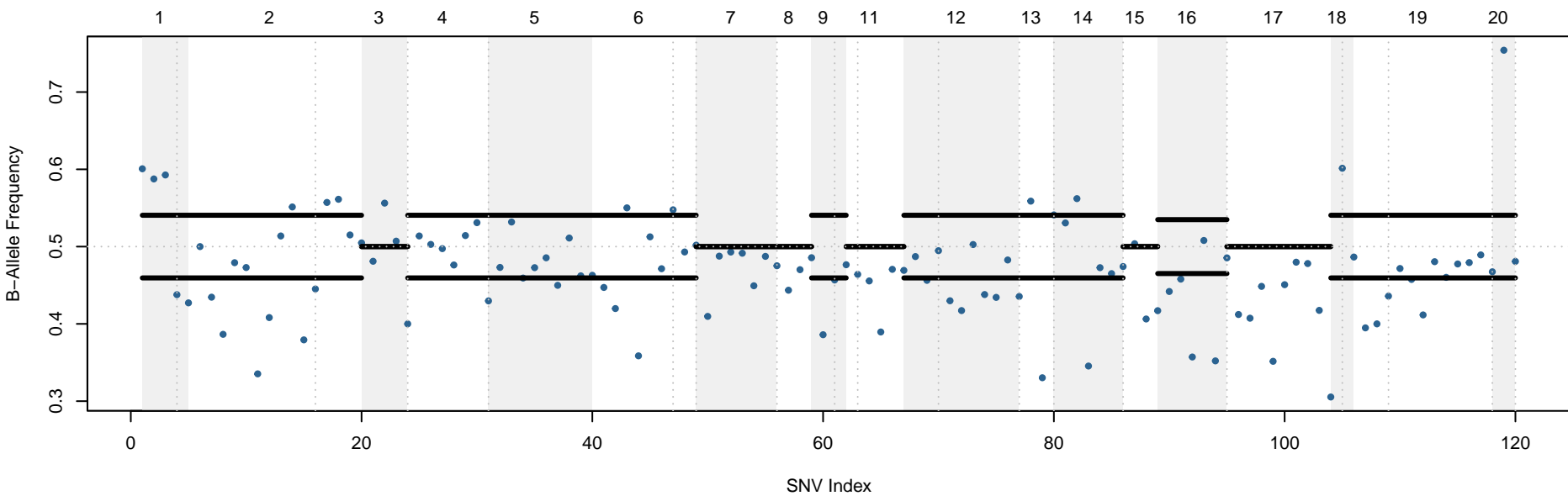




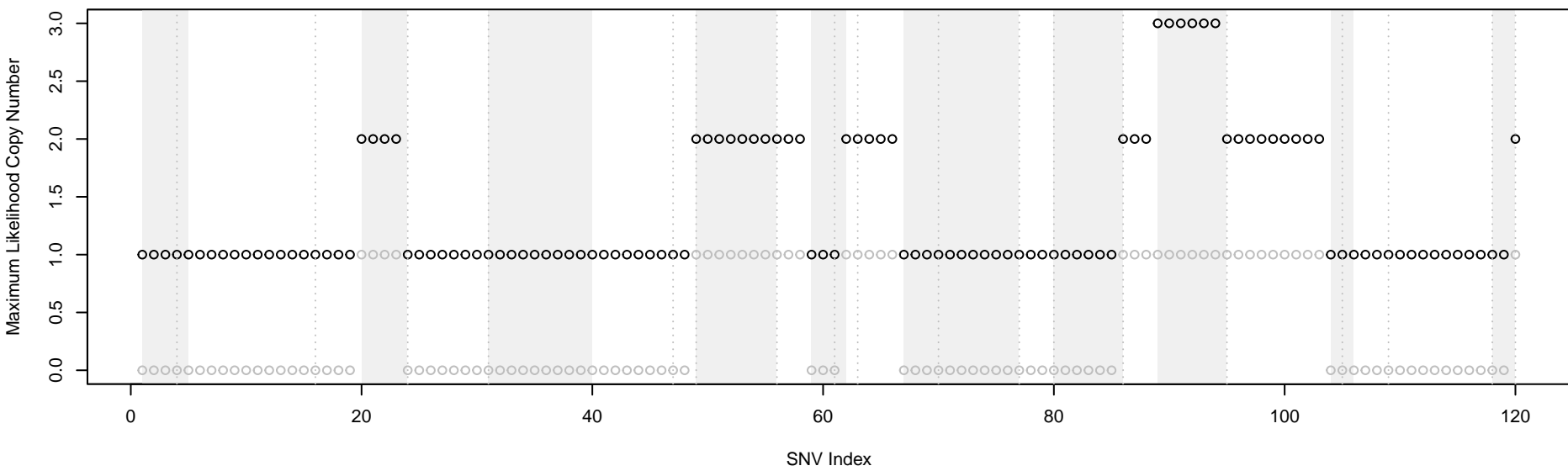
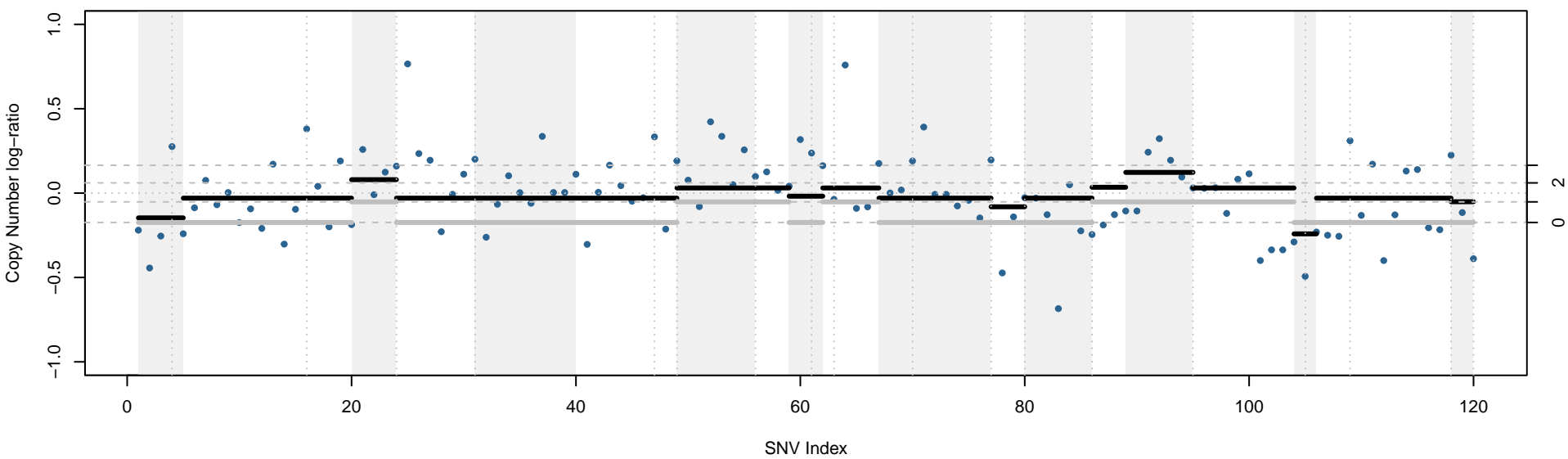
Purity: 0.15 Tumor ploidy: 1.454

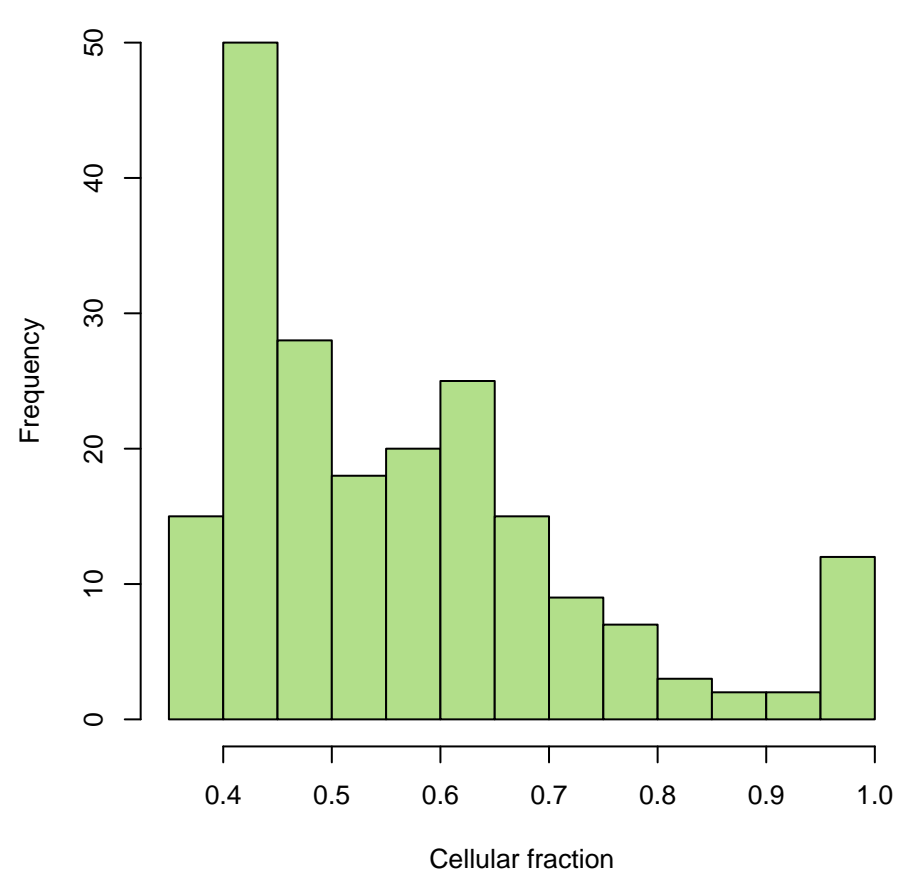
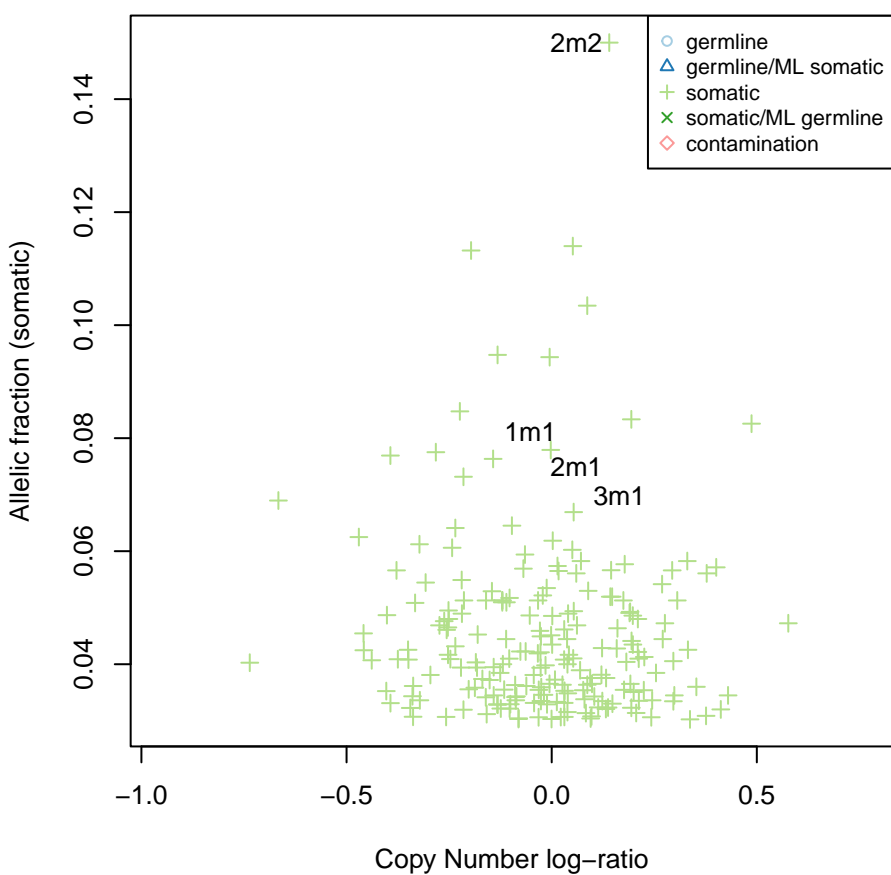
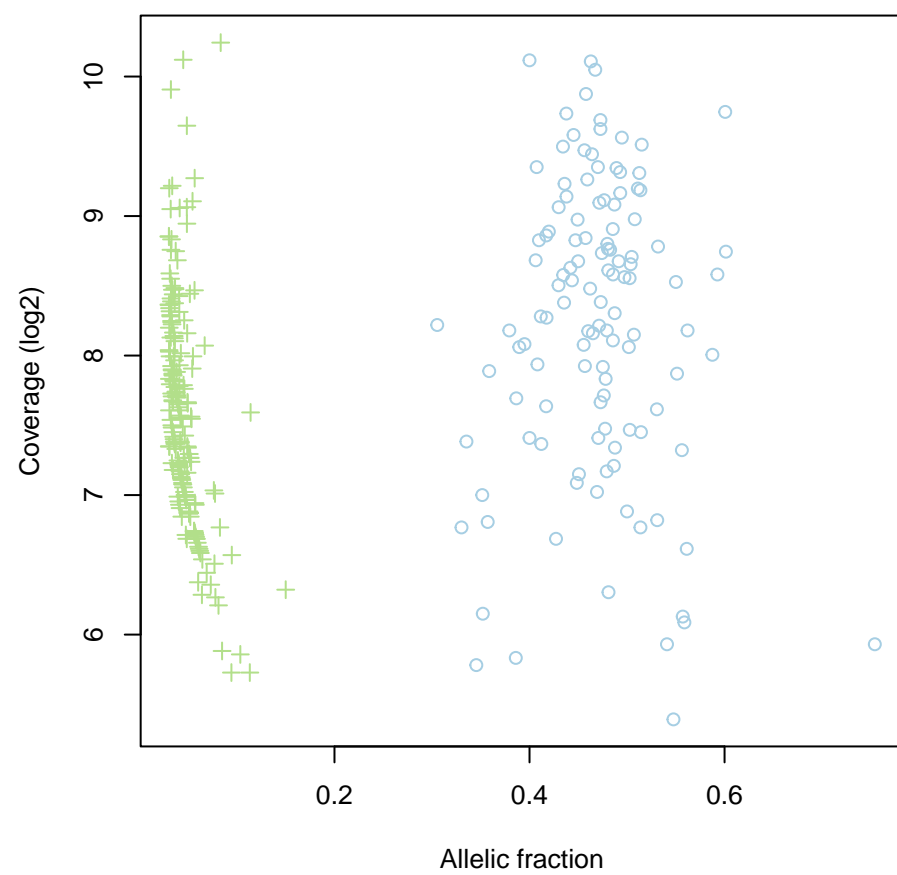
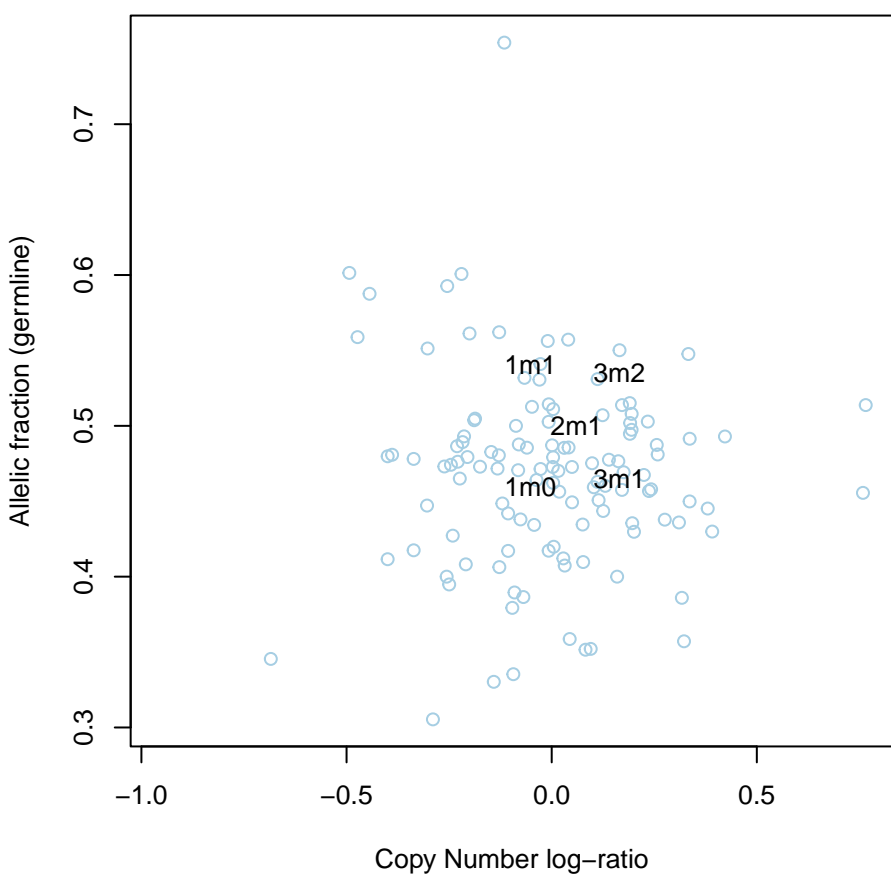


Purity: 0.15 Tumor ploidy: 1.454 SNV log-likelihood: 44.34 GoF: 95.7% Mean coverage: 406;281

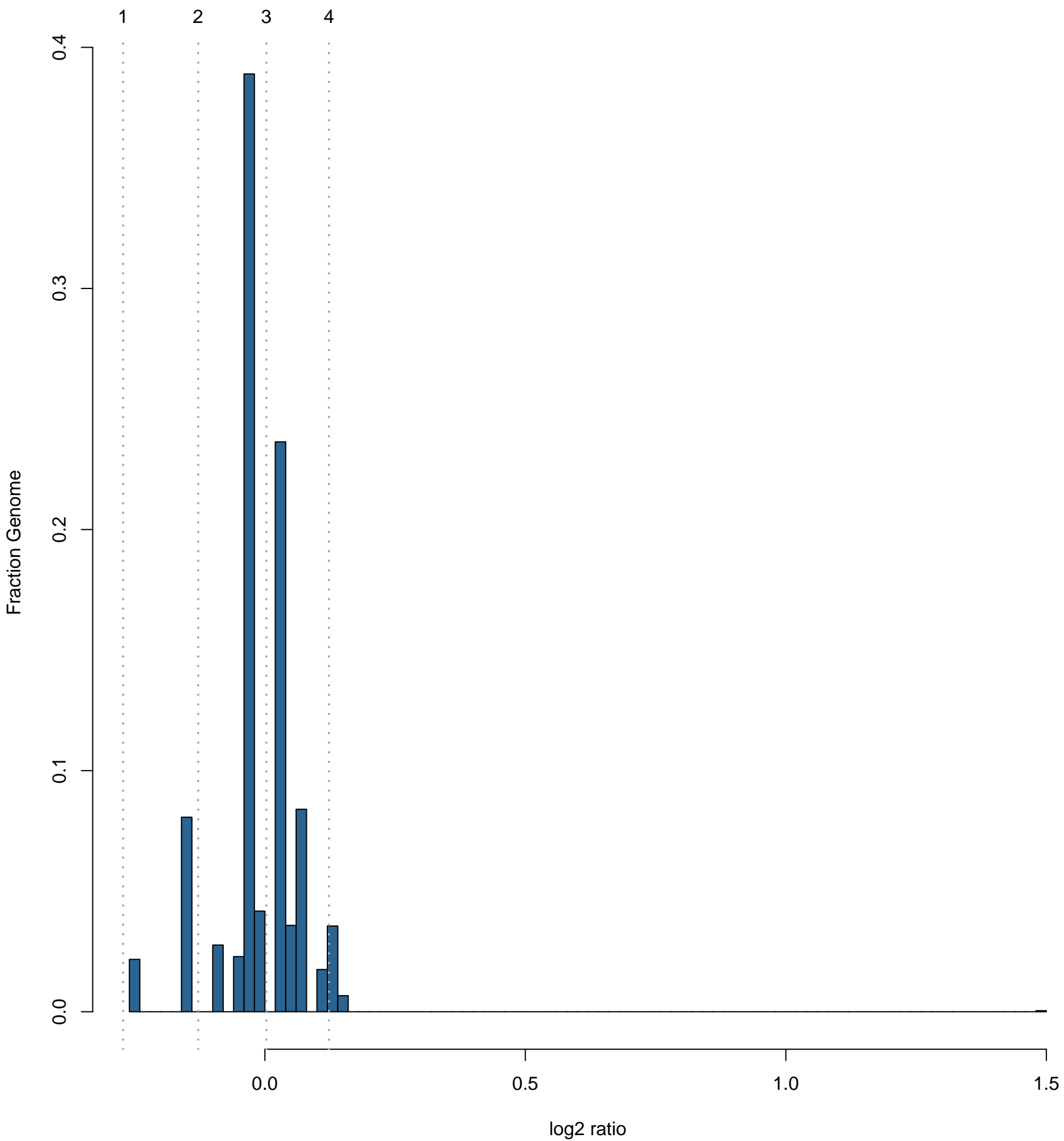


SCNA-fit log-likelihood: -15991.2

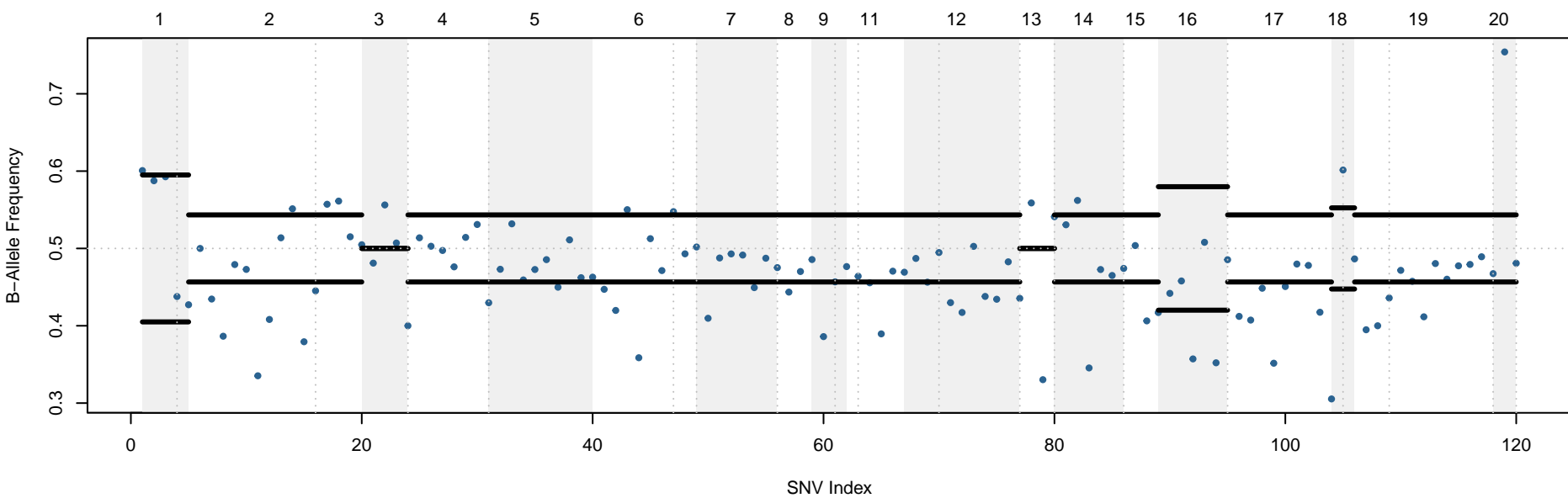




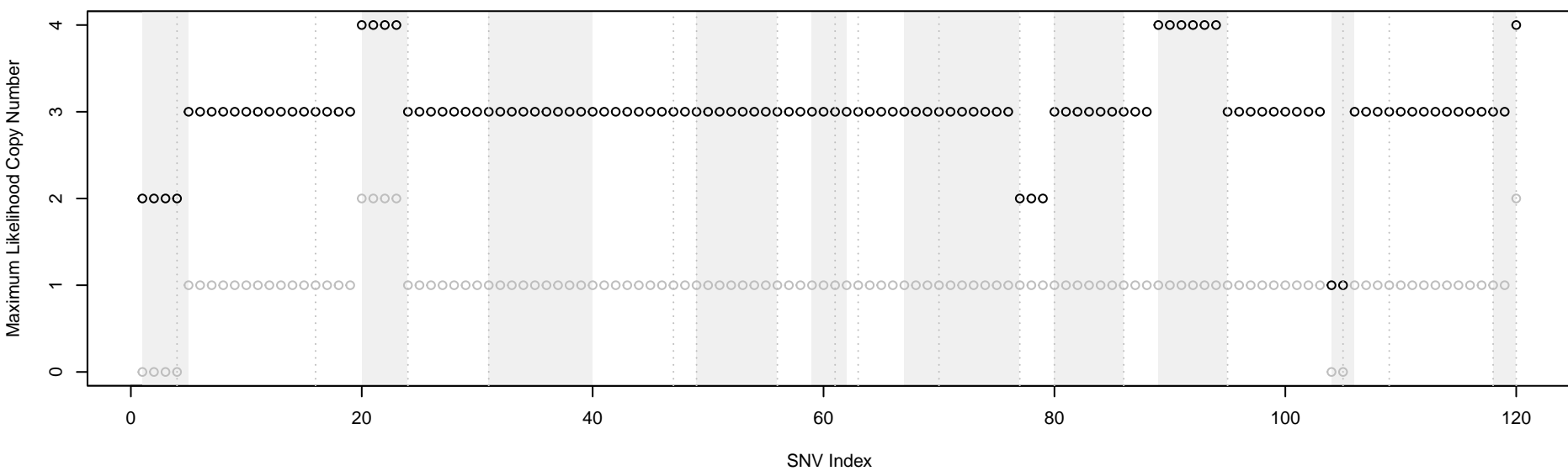
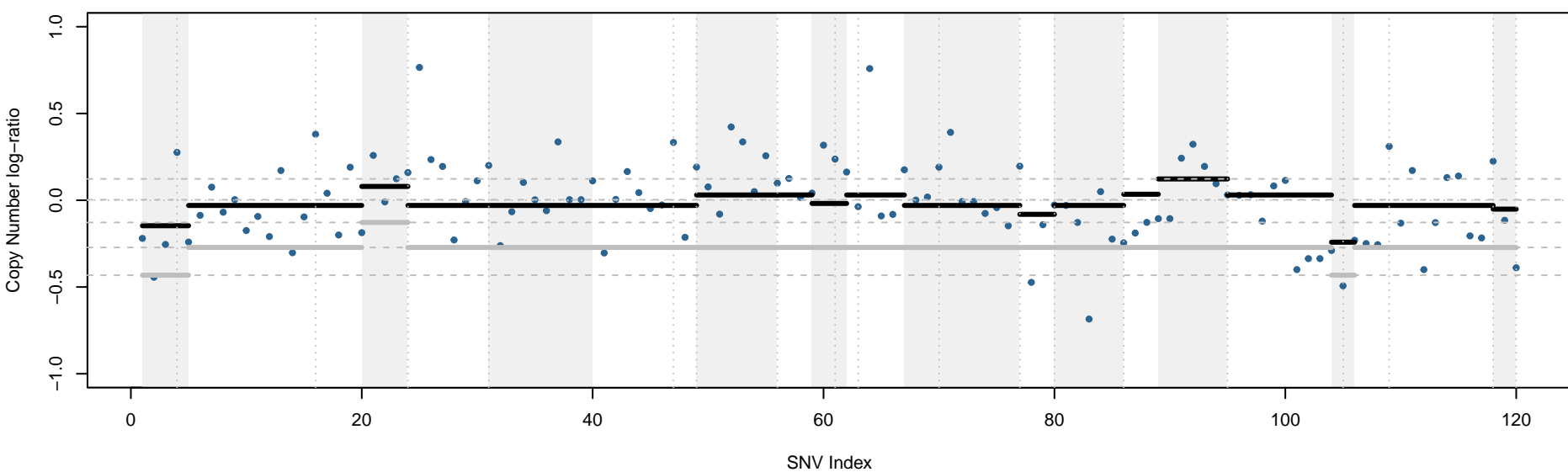
Purity: 0.19 Tumor ploidy: 2.976



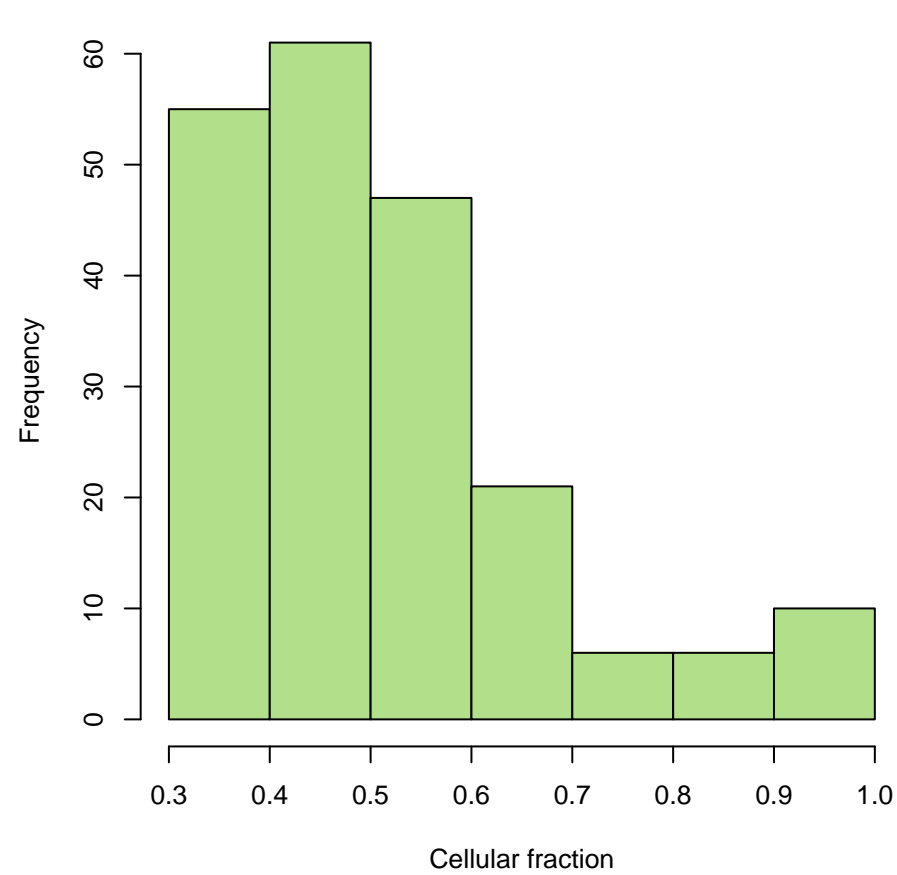
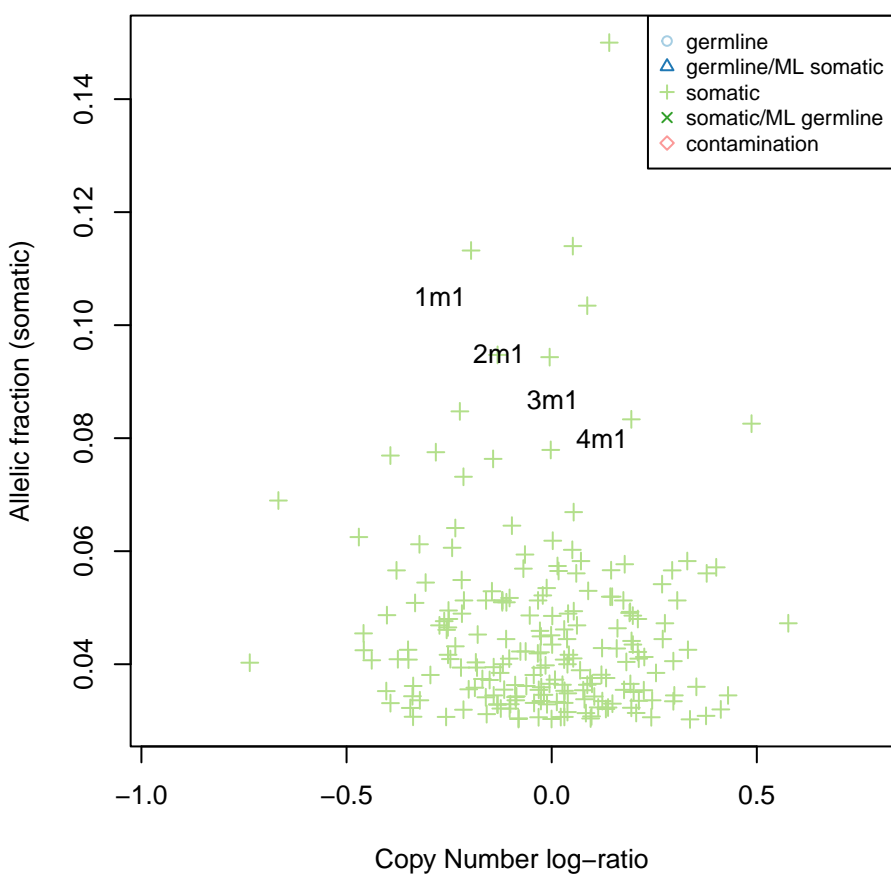
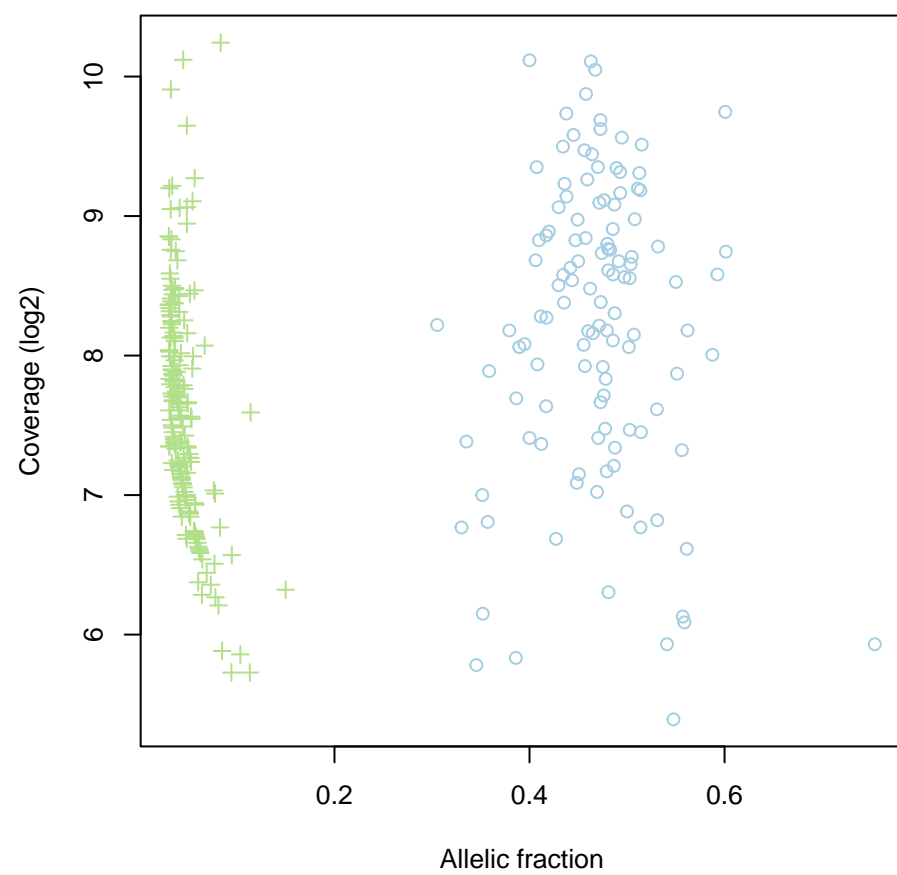
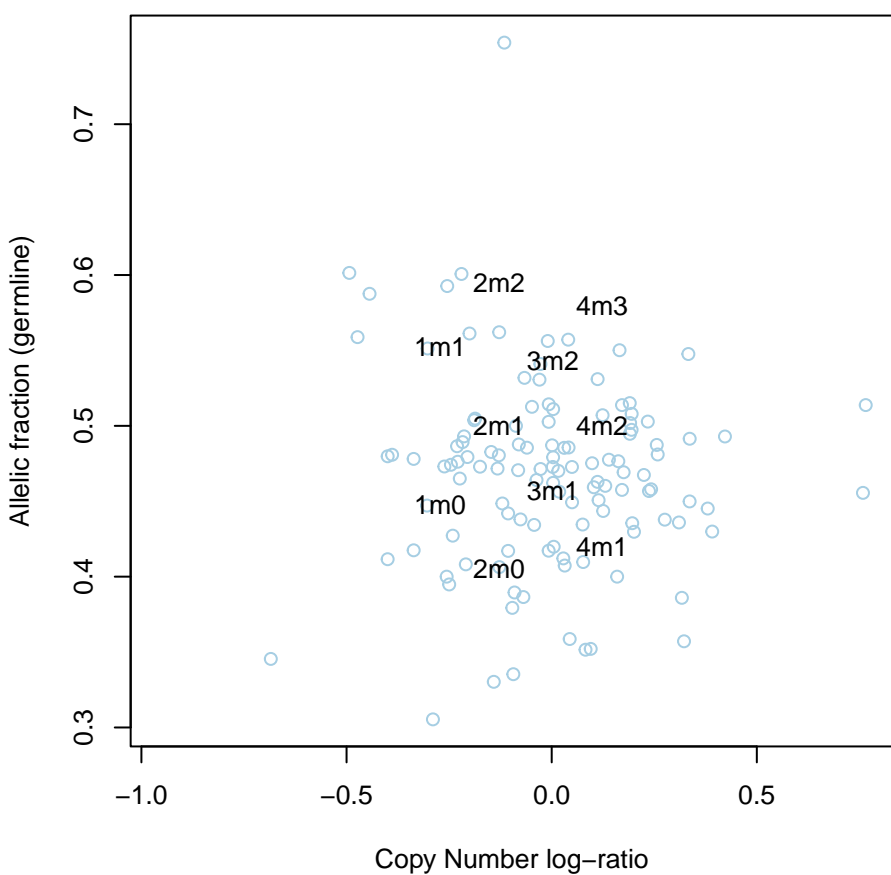
Purity: 0.19 Tumor ploidy: 2.976 SNV log-likelihood: 14.38 GoF: 95.1% Mean coverage: 406;281



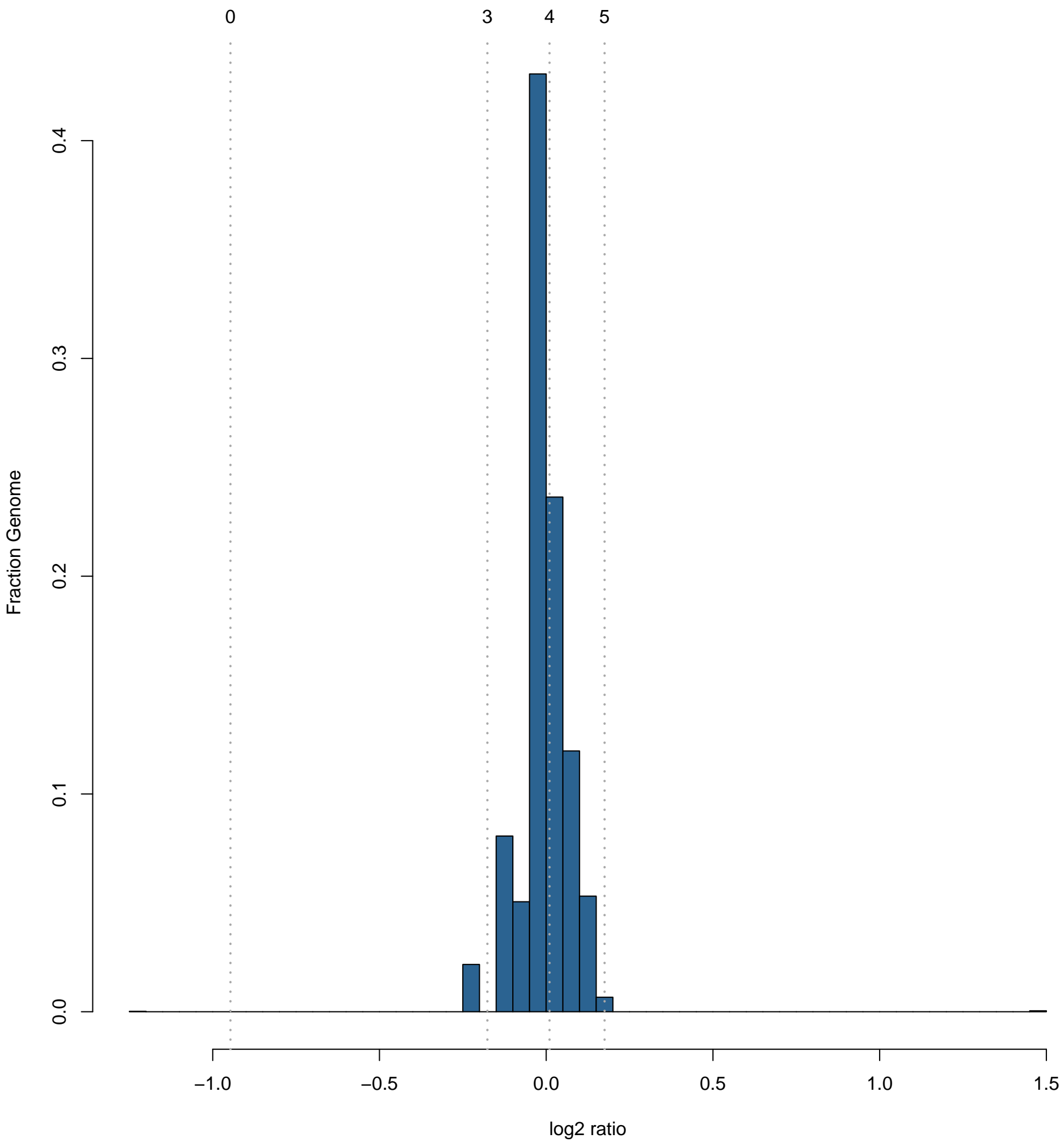
SCNA-fit log-likelihood: -15953.58



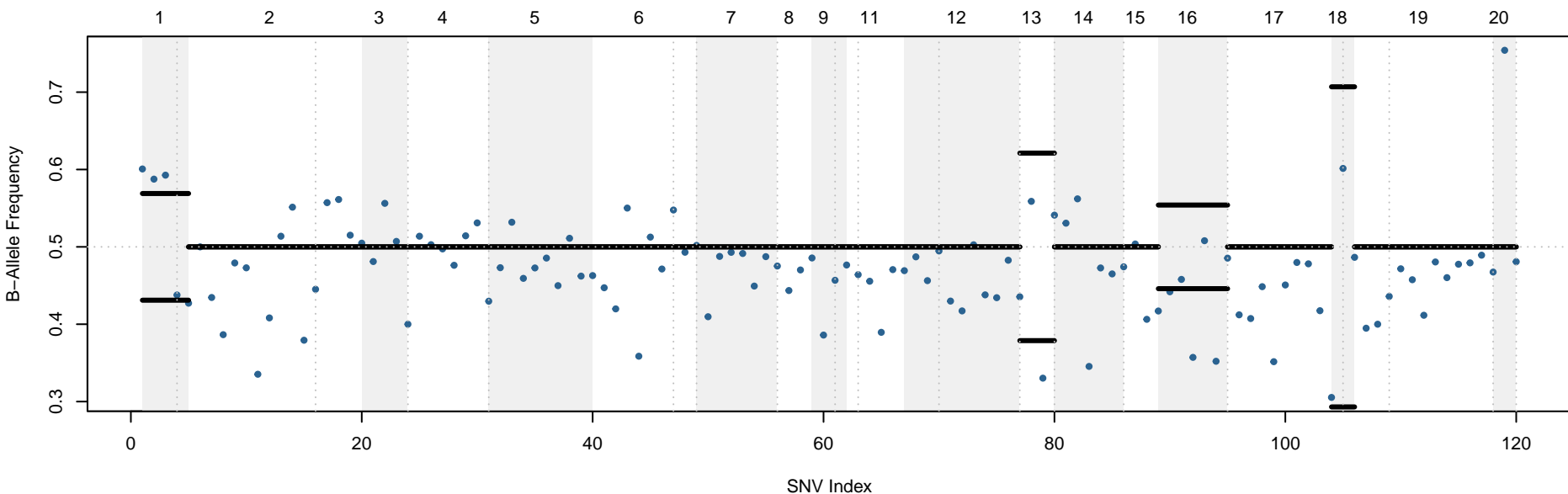




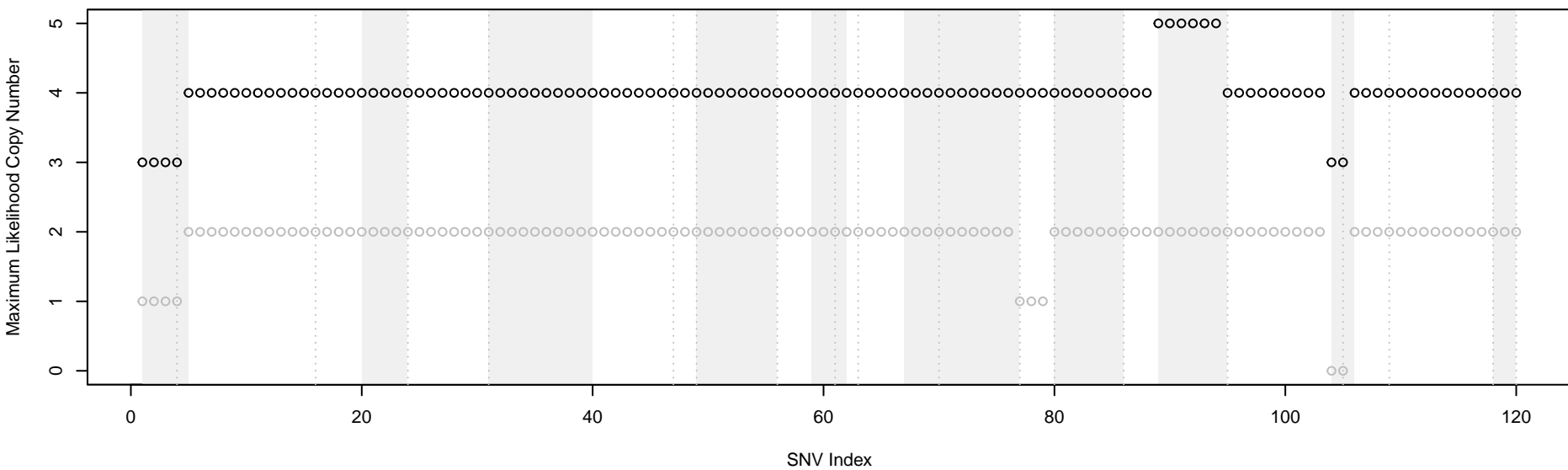
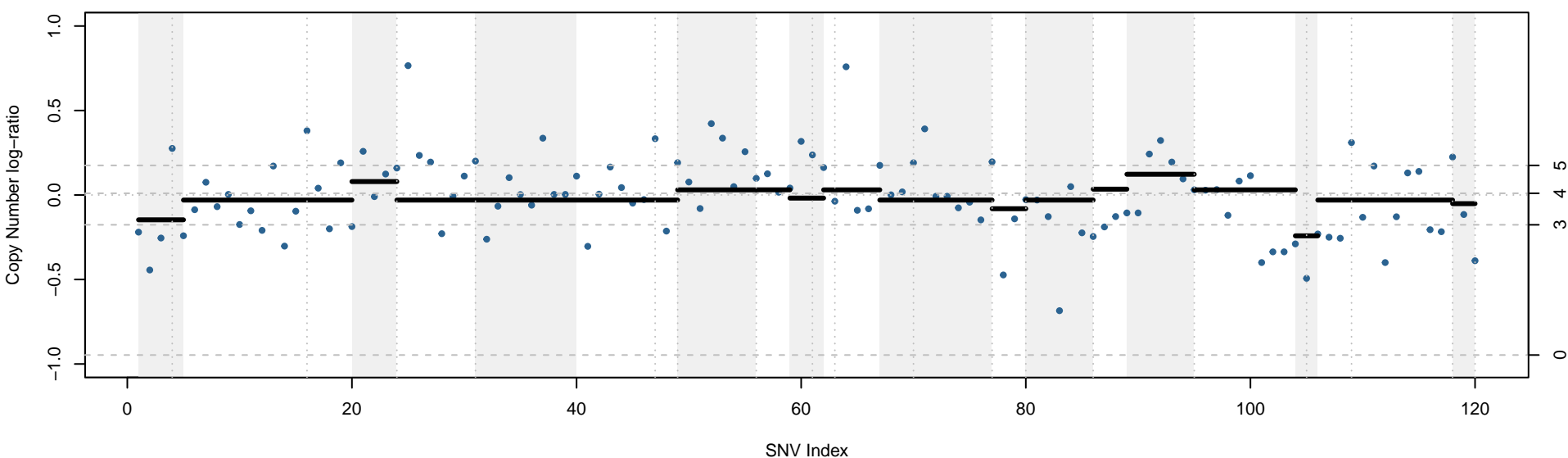
Purity: 0.32 Tumor ploidy: 3.943

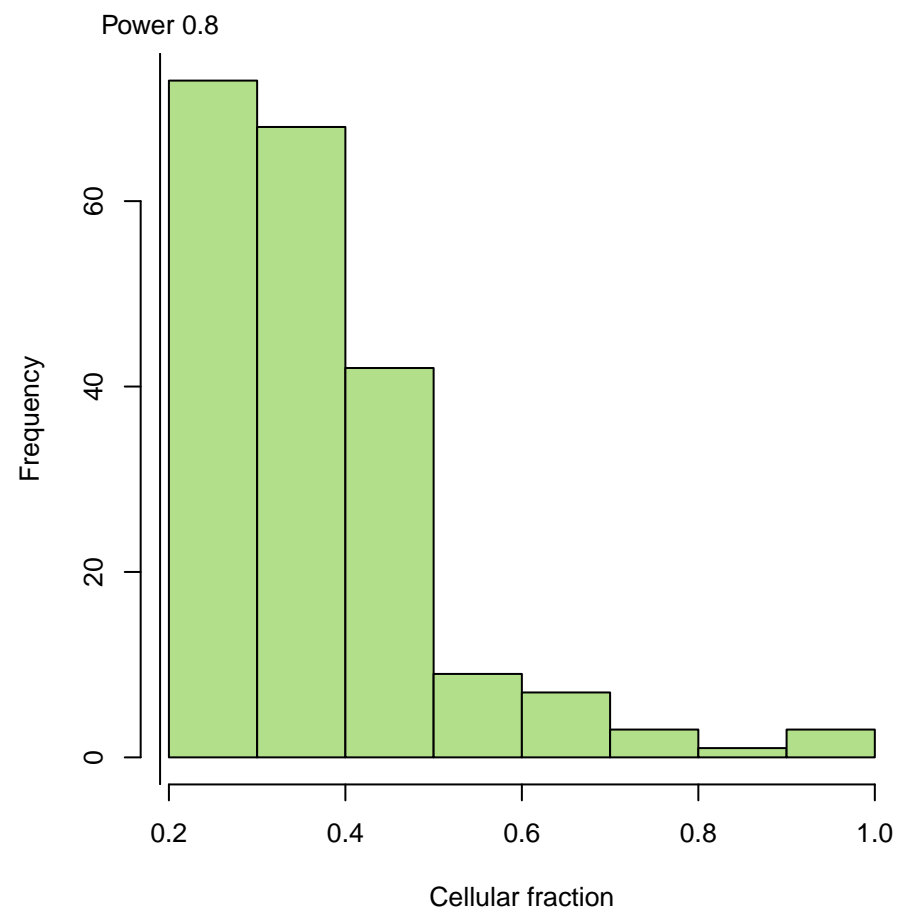
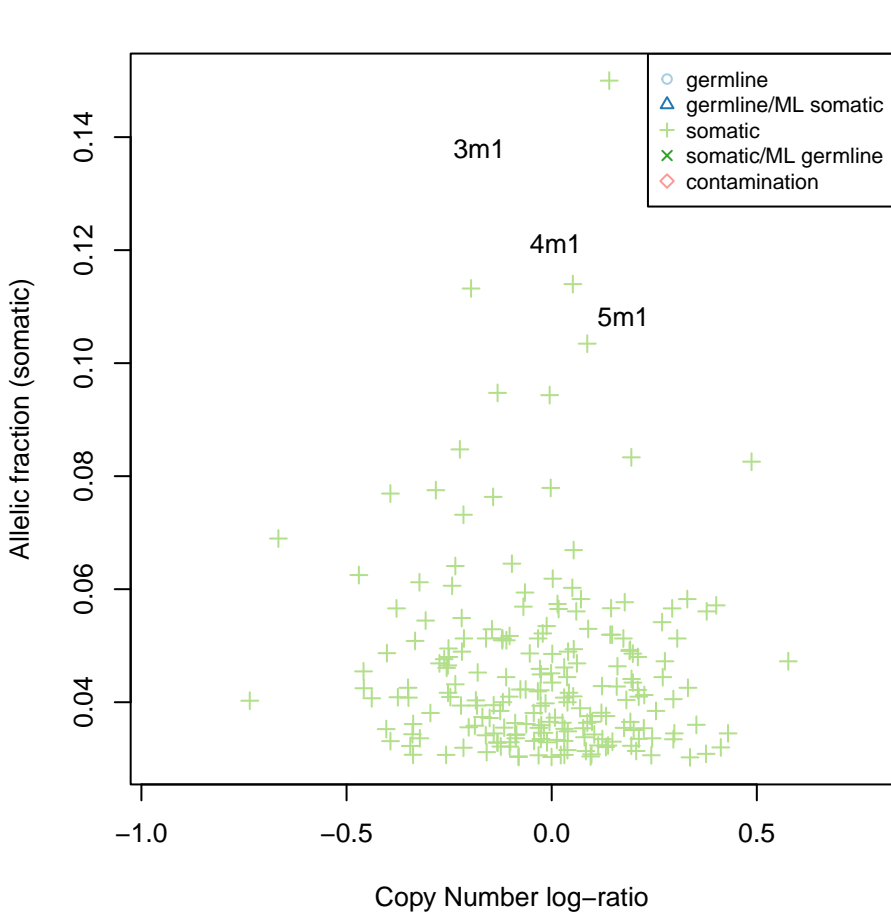
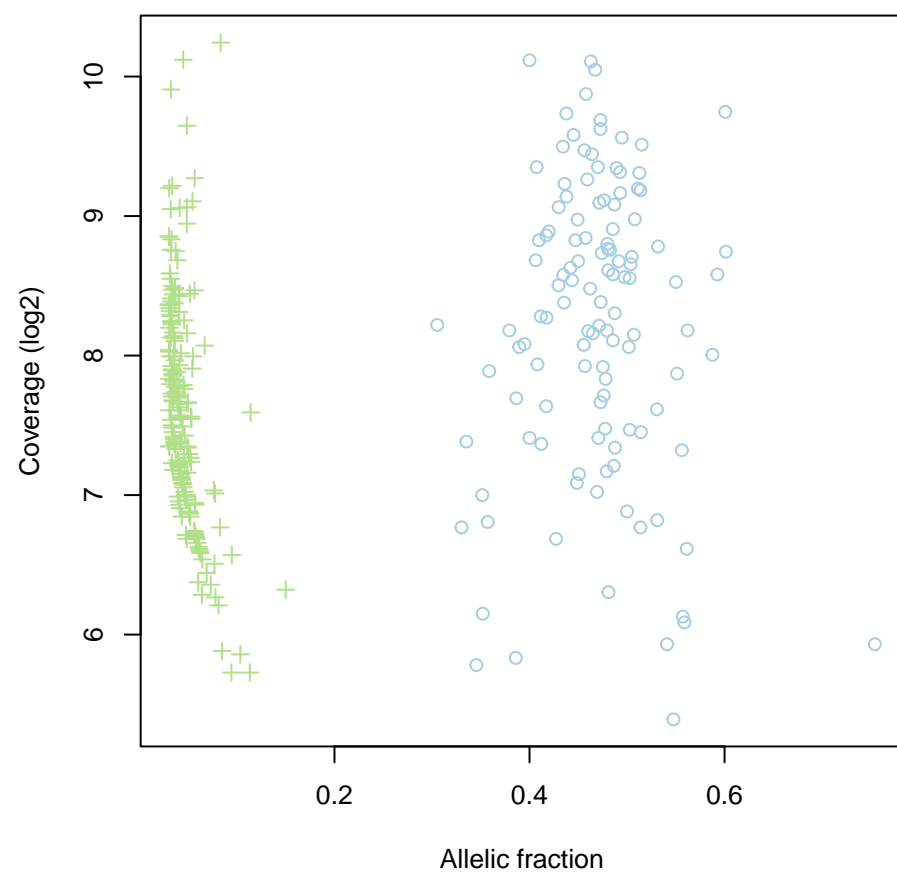
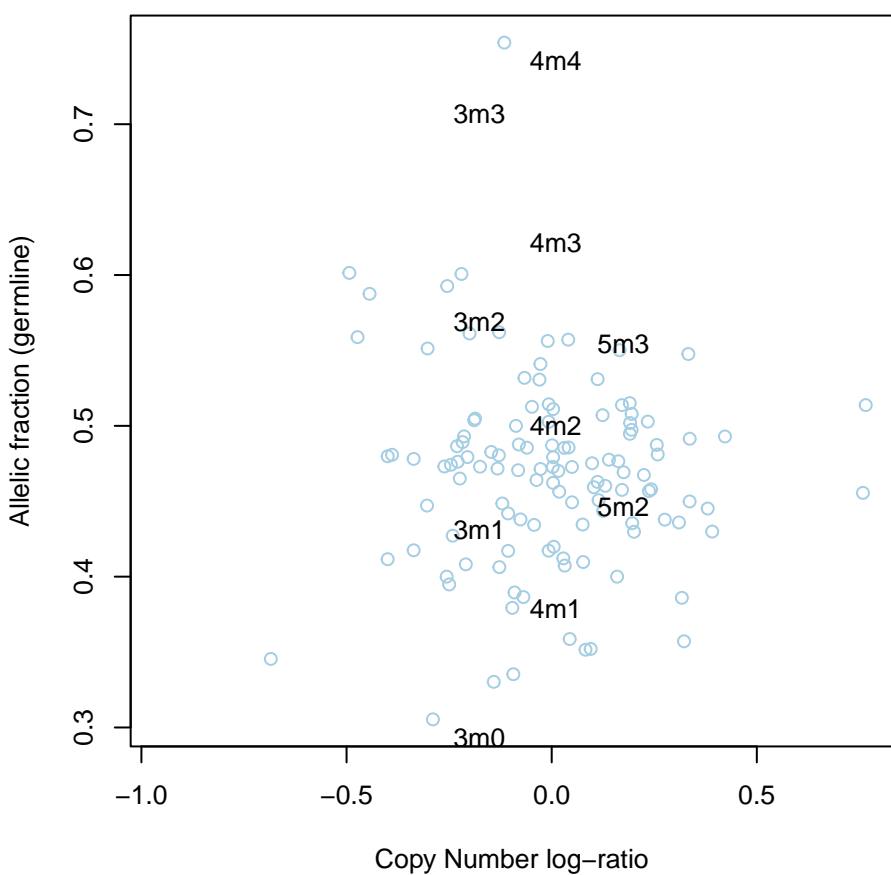


Purity: 0.32 Tumor ploidy: 3.943 SNV log-likelihood: 3.08 GoF: 88.3% Mean coverage: 406;281

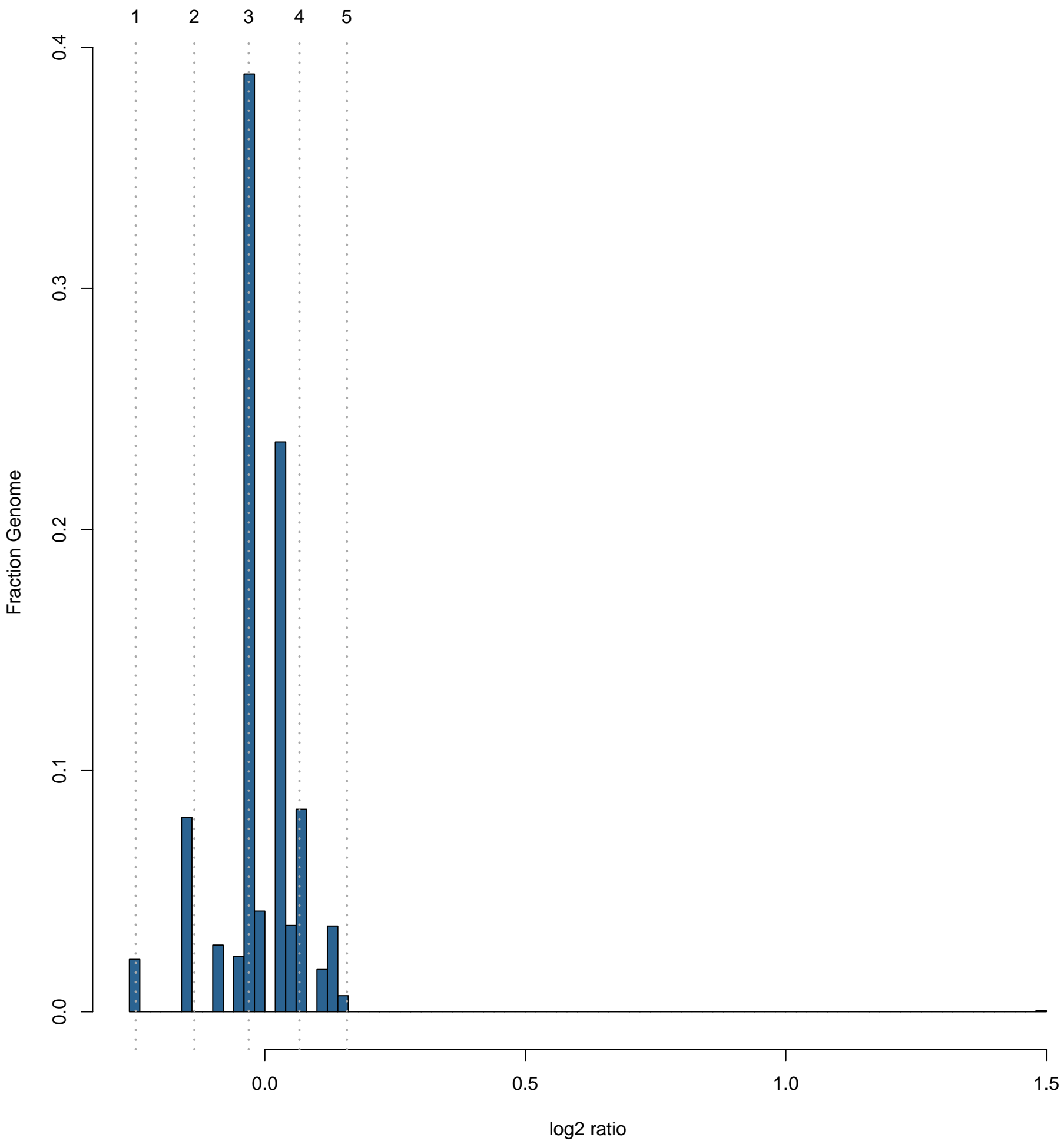


SCNA-fit log-likelihood: -15980.35

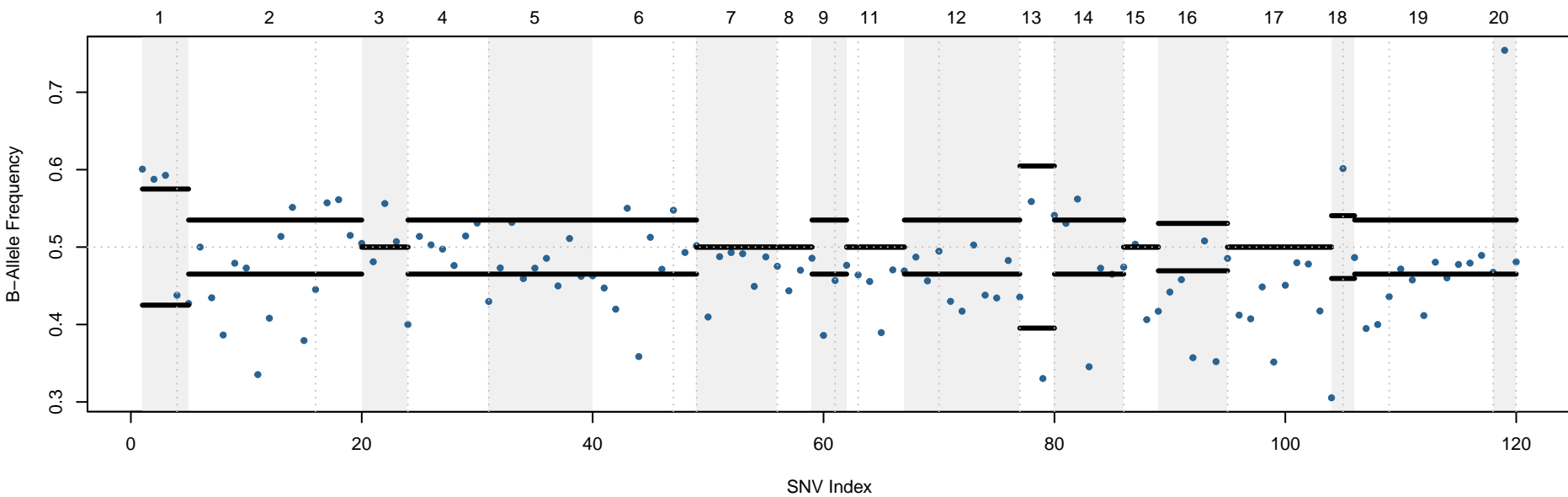




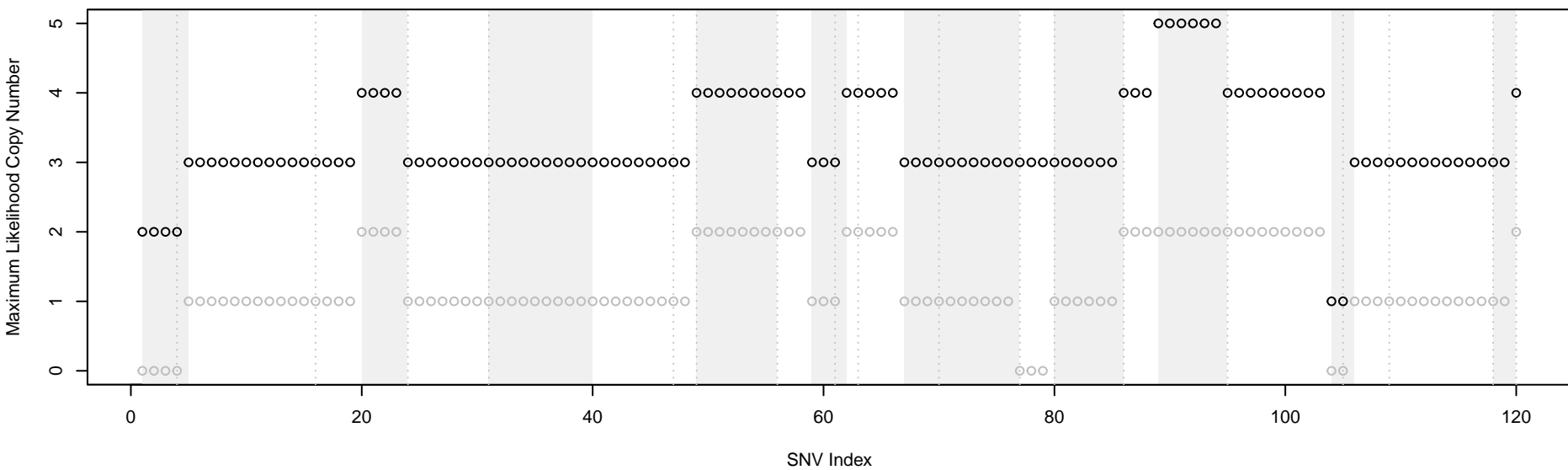
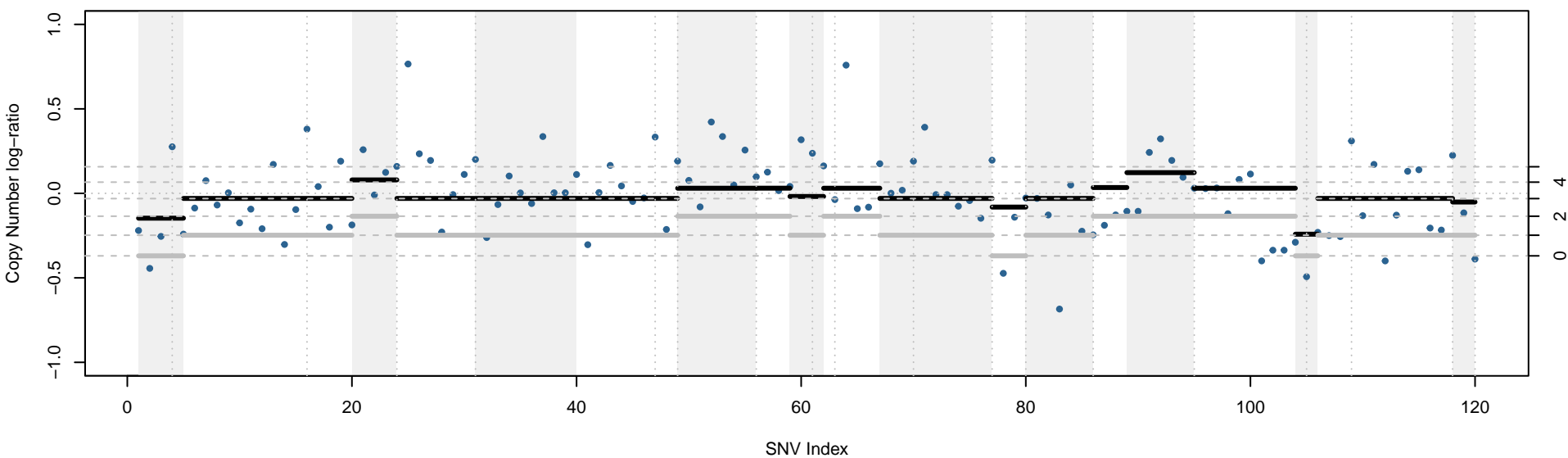
Purity: 0.15 Tumor ploidy: 3.311

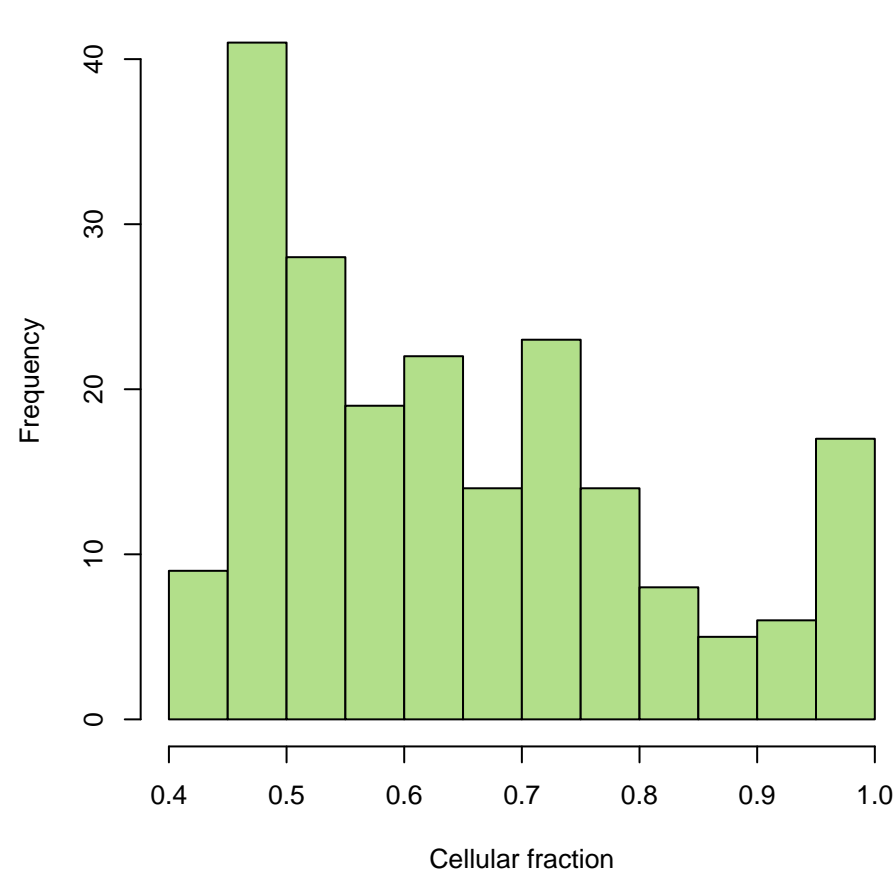
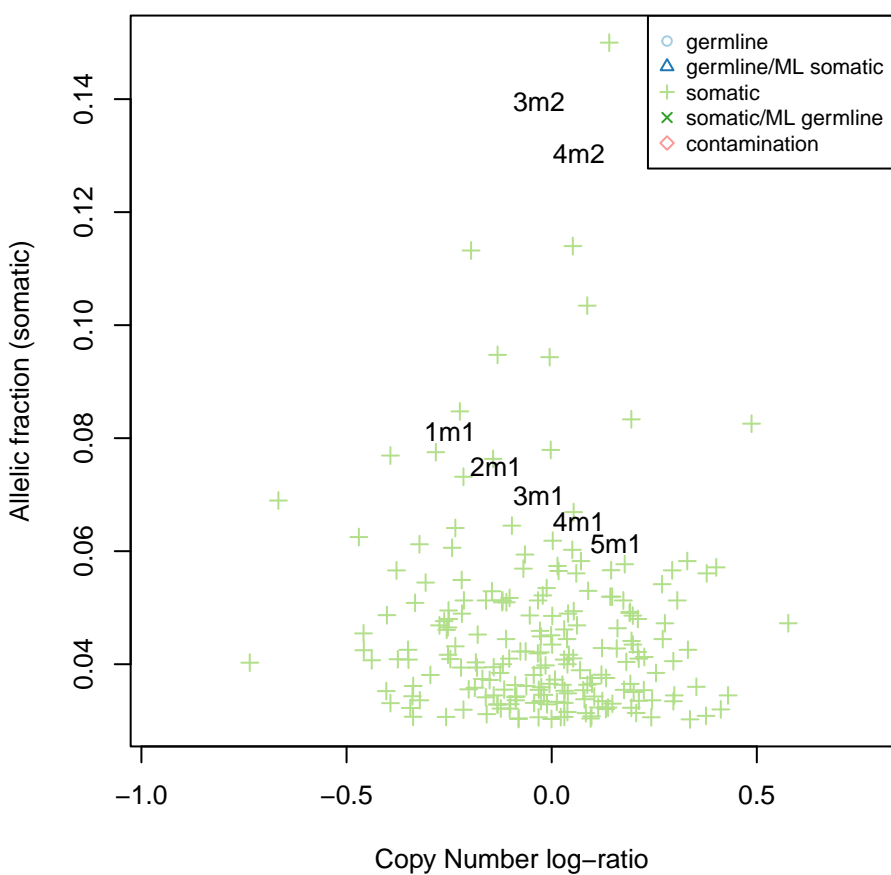
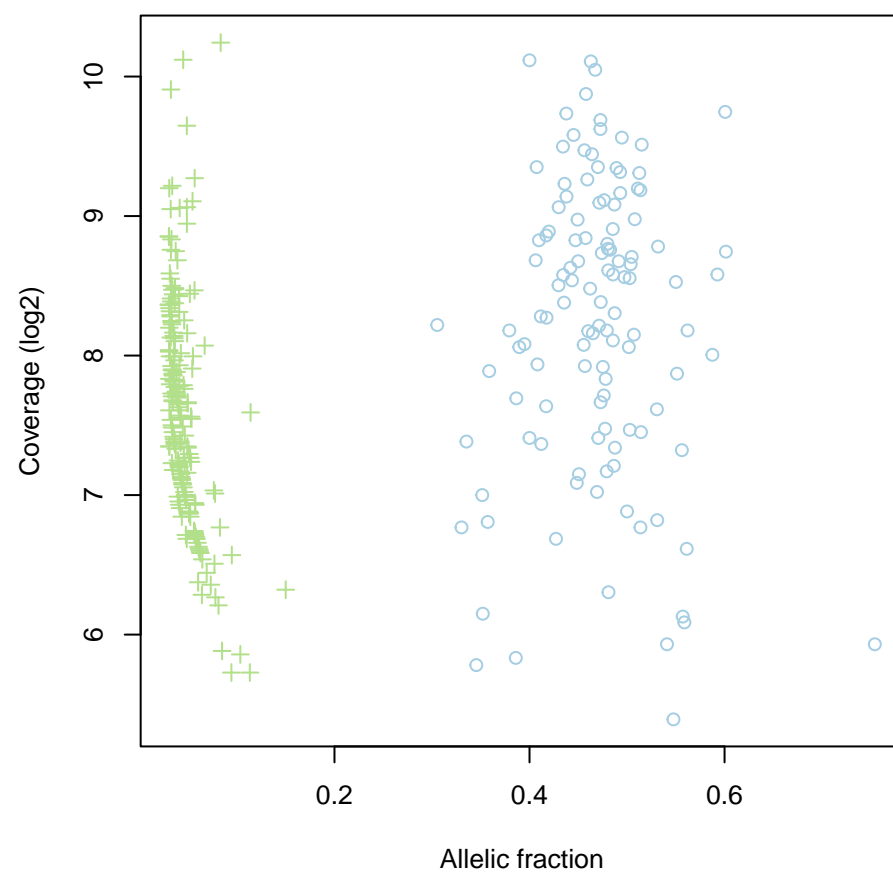
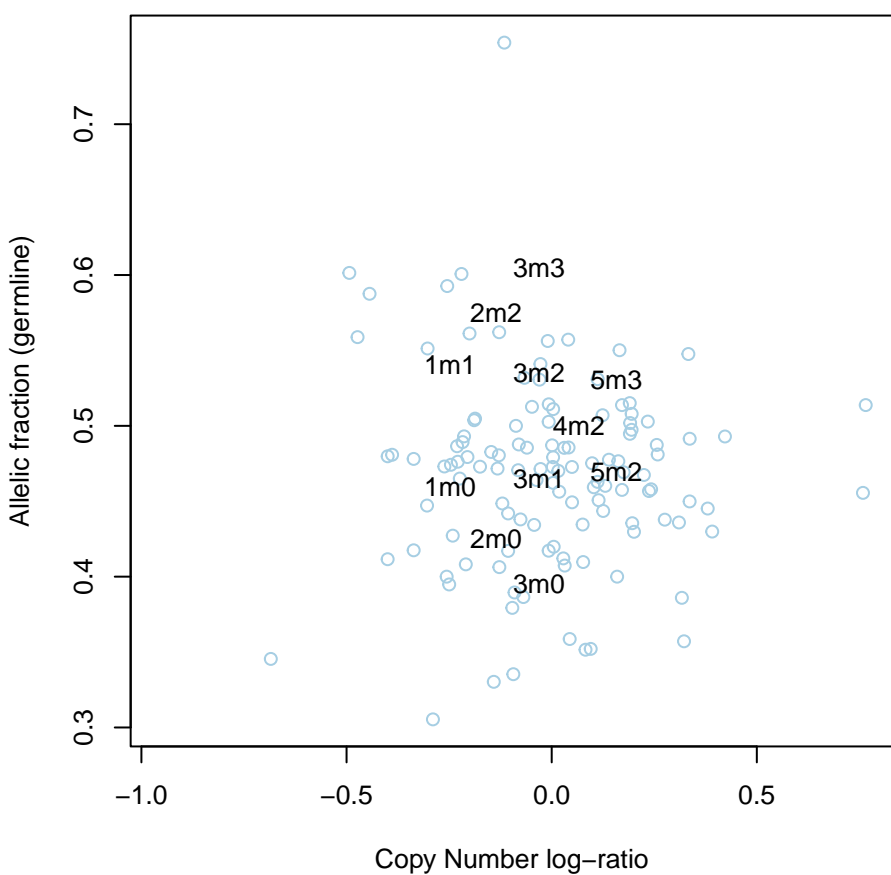


Purity: 0.15 Tumor ploidy: 3.311 SNV log-likelihood: -20.69 GoF: 96.7% Mean coverage: 406;281

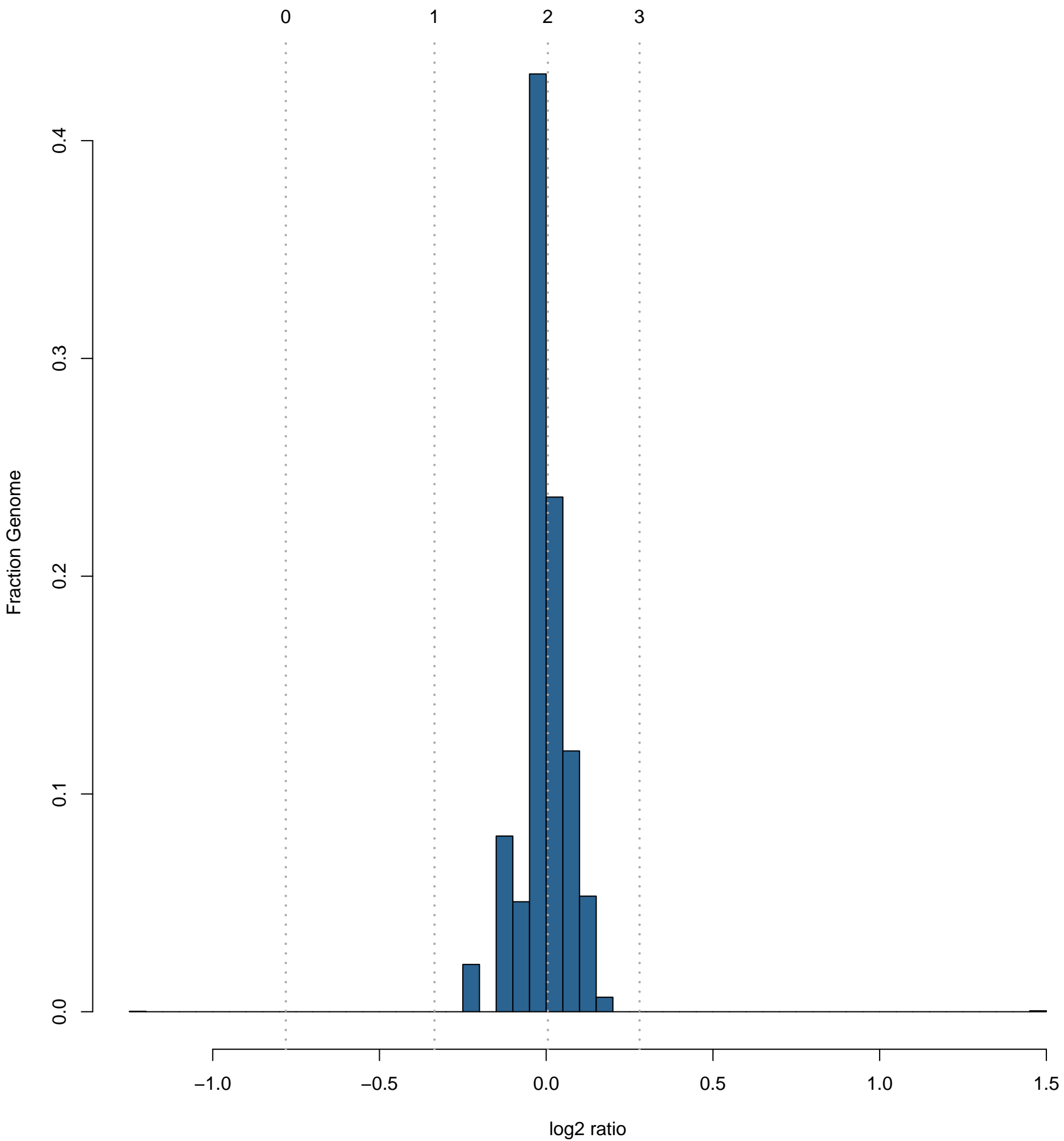


SCNA-fit log-likelihood: -15932.8



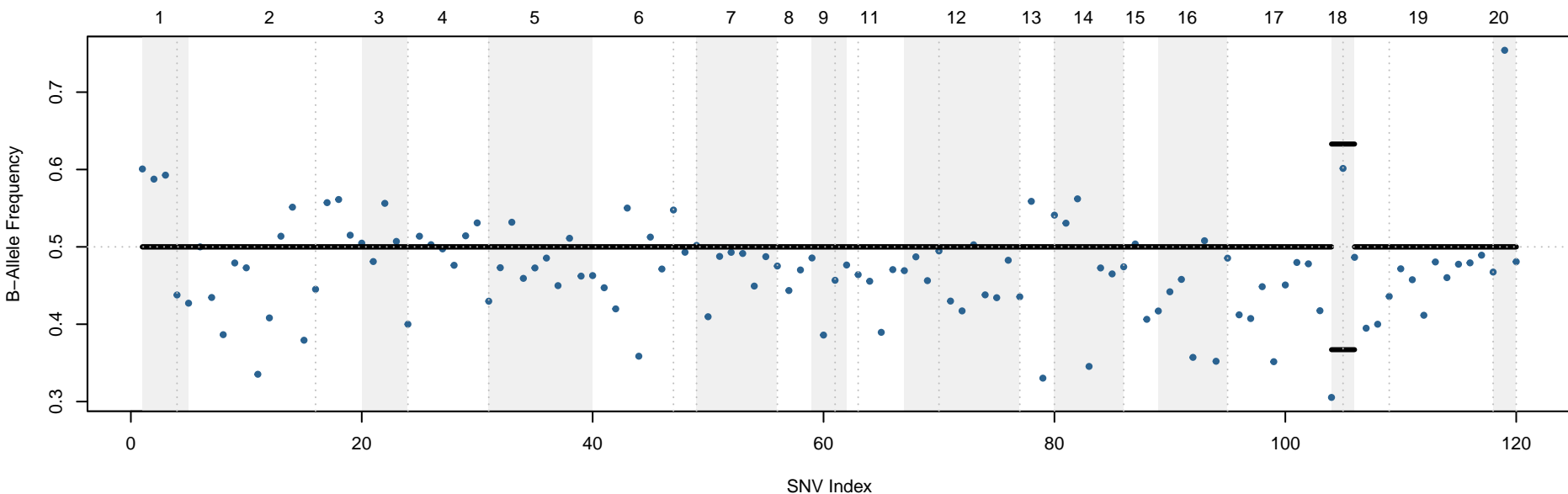


Purity: 0.42 Tumor ploidy: 1.983

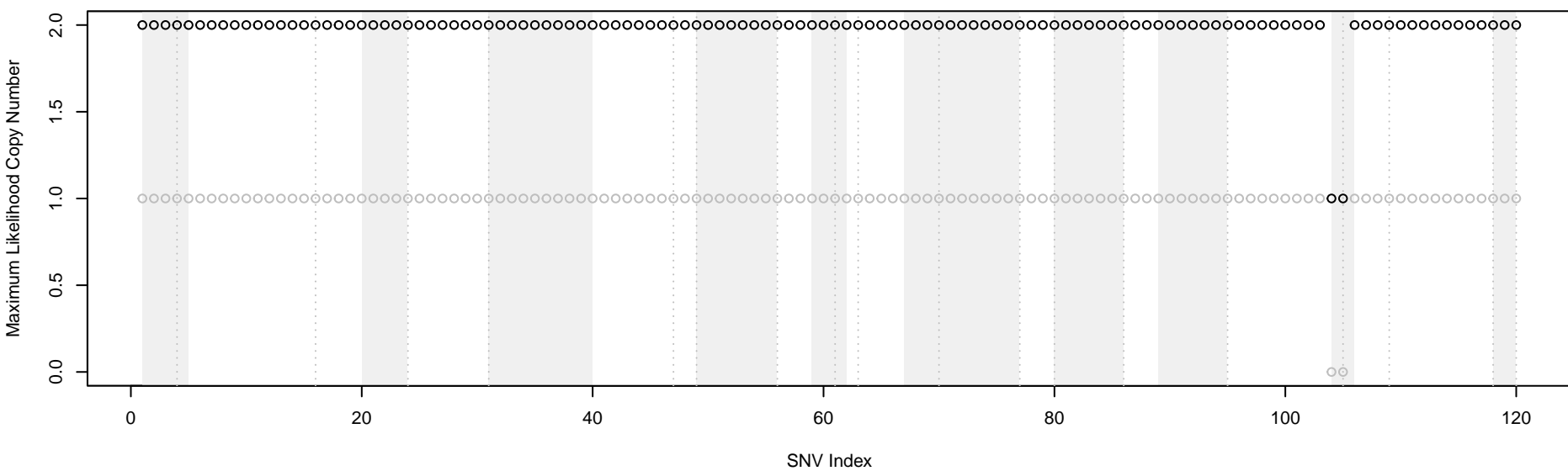
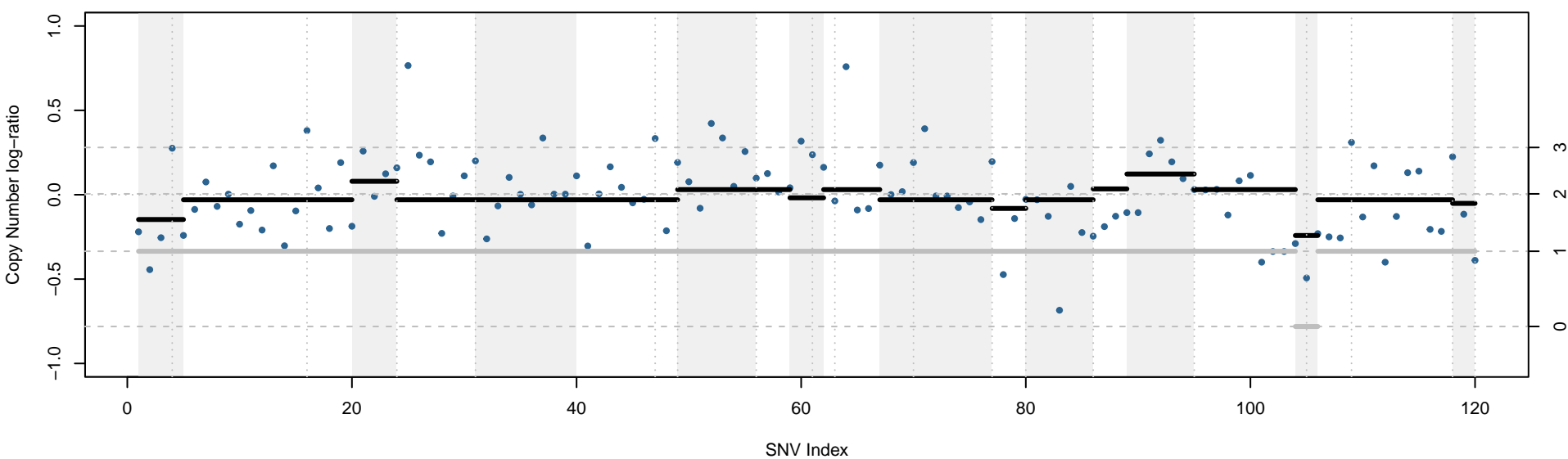


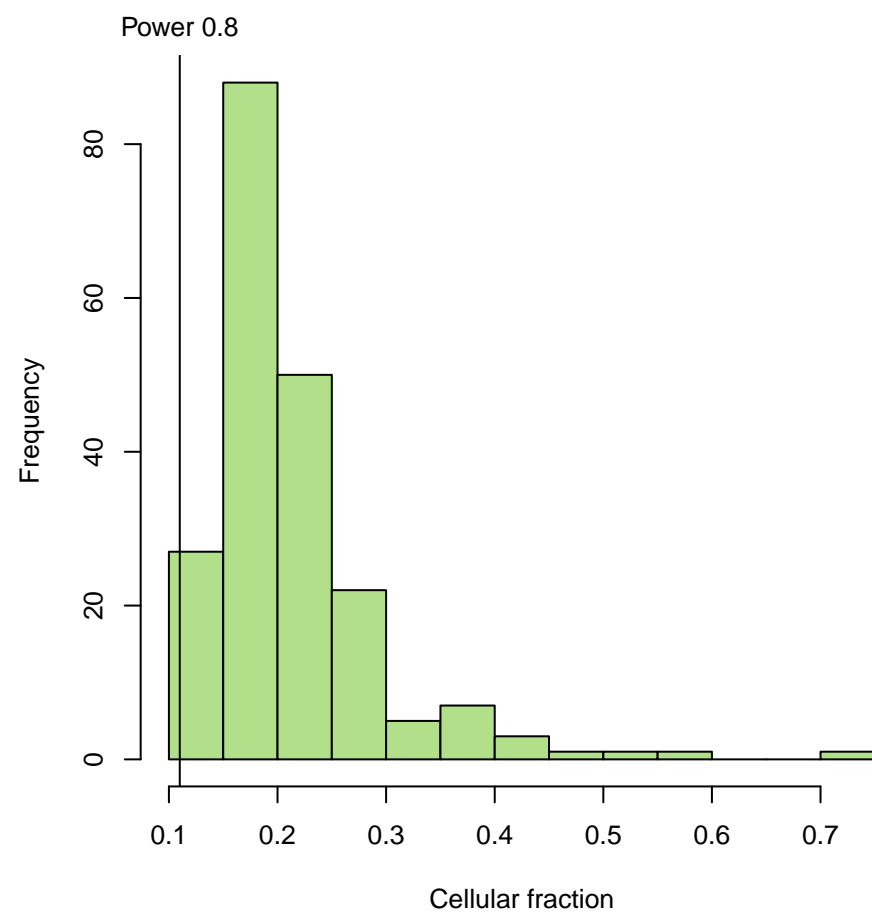
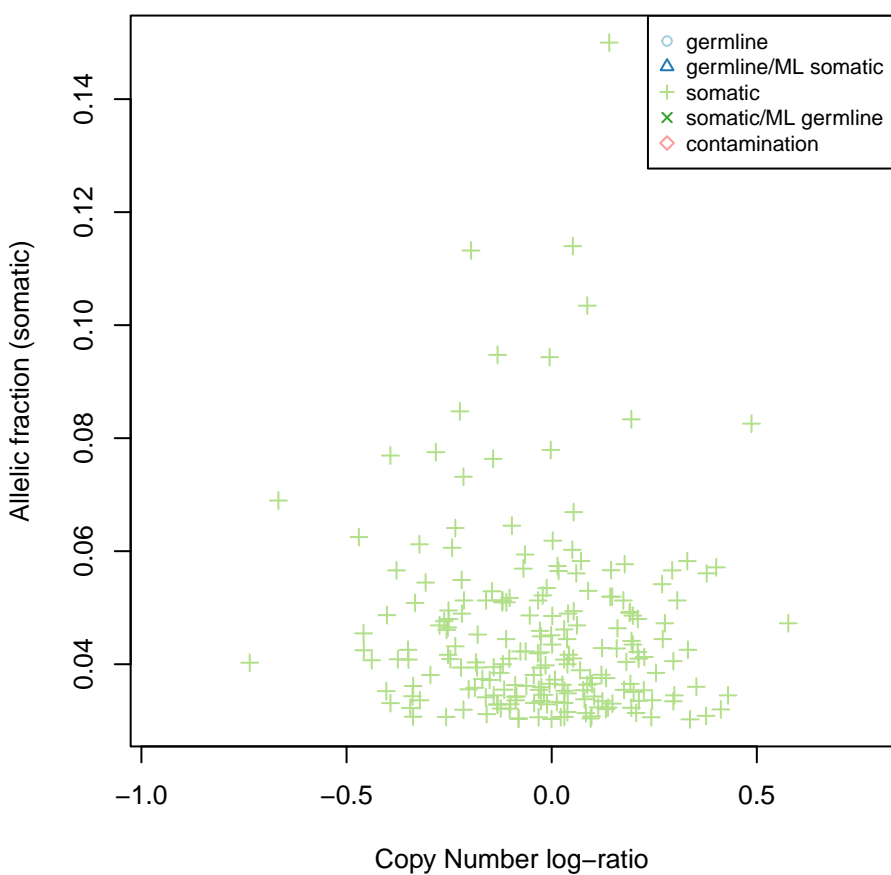
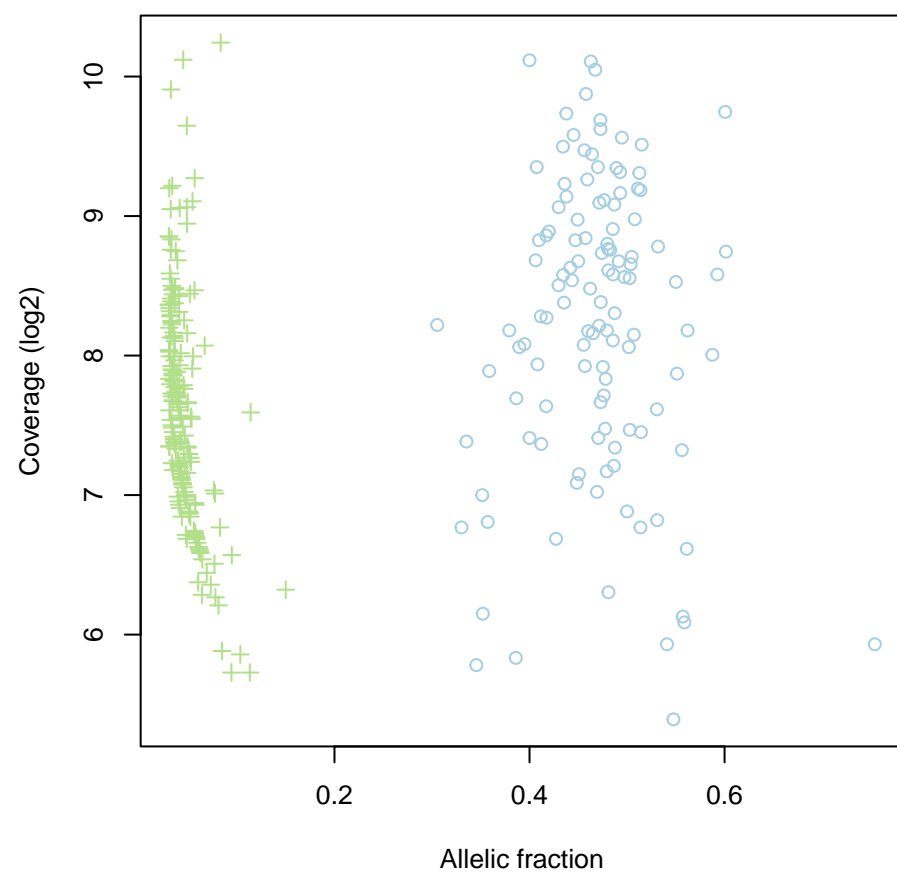
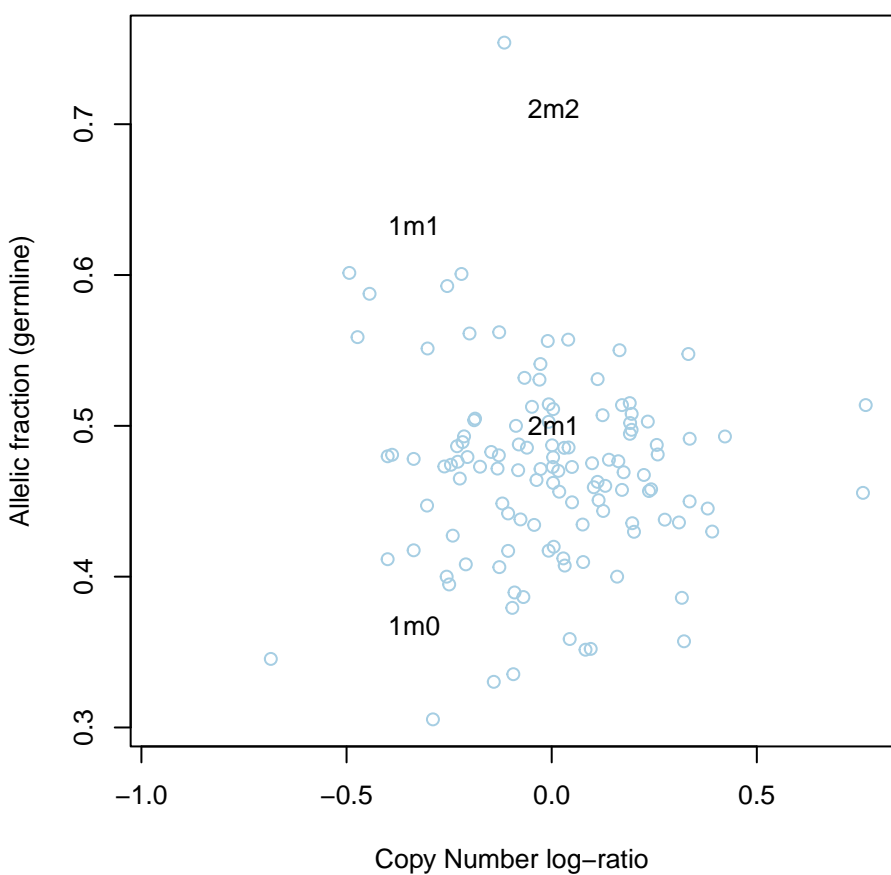


Purity: 0.42 Tumor ploidy: 1.983 SNV log-likelihood: -38.2 GoF: 54% Mean coverage: 406;281

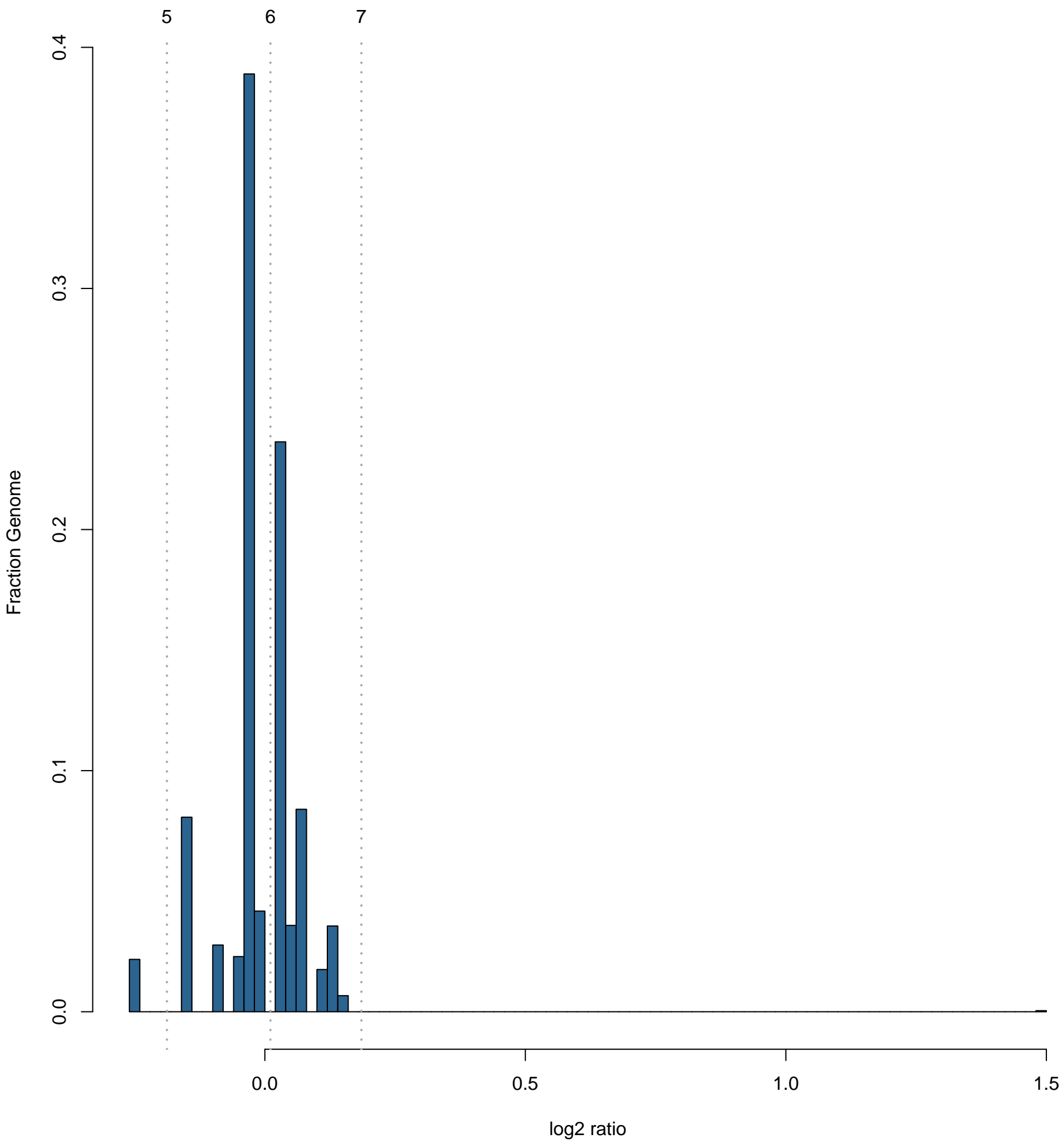


SCNA-fit log-likelihood: -16067.8

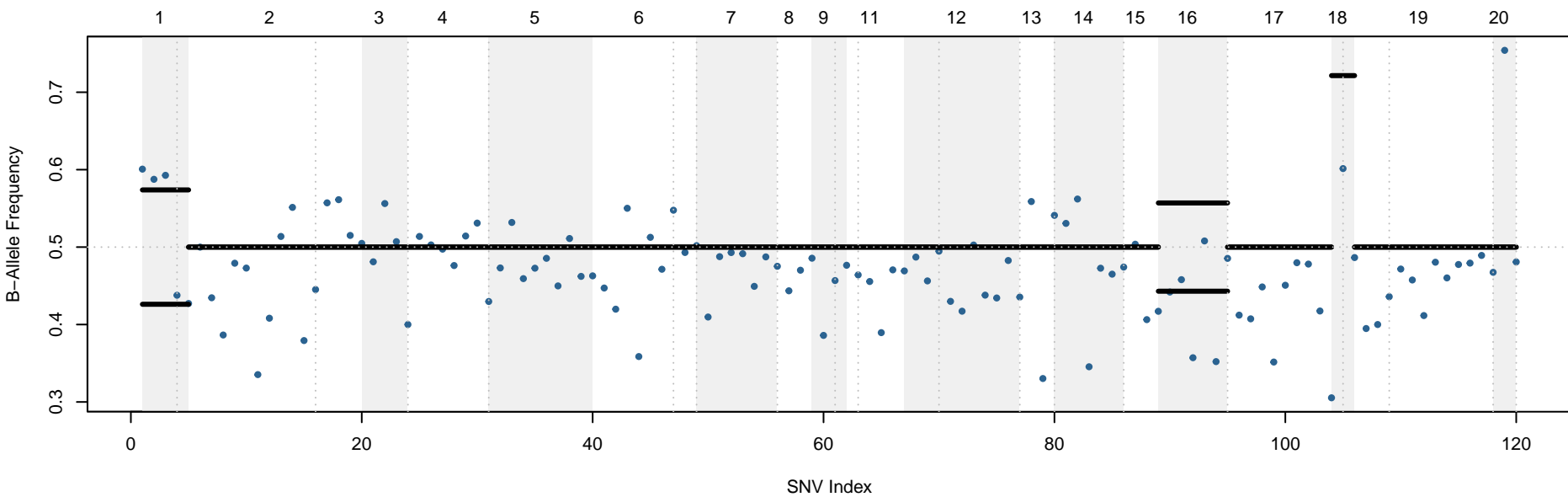




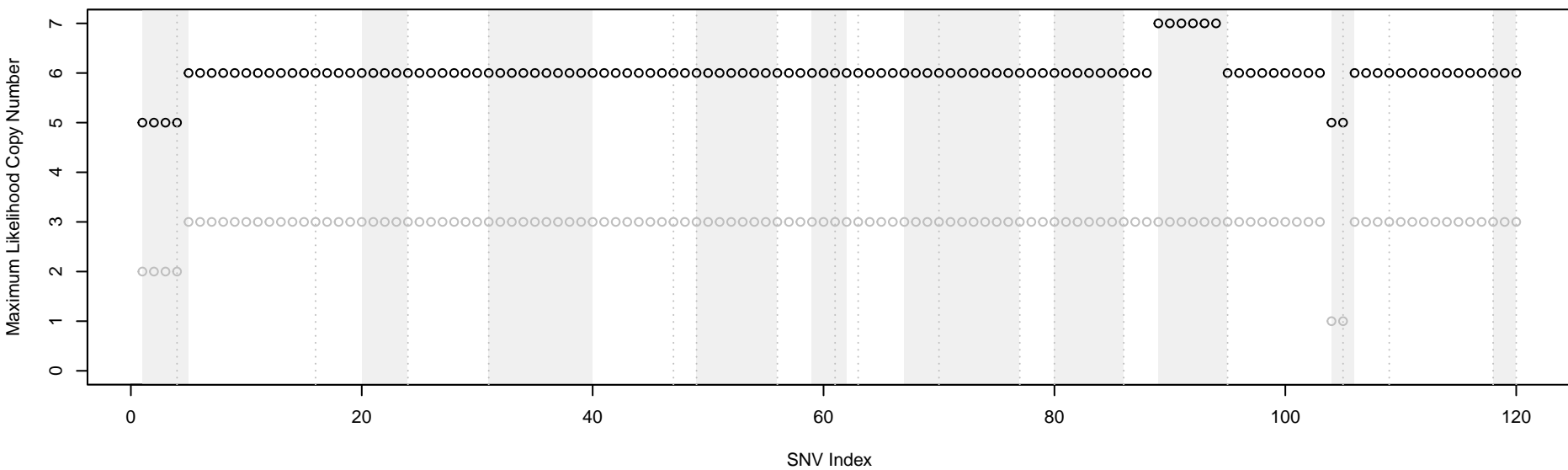
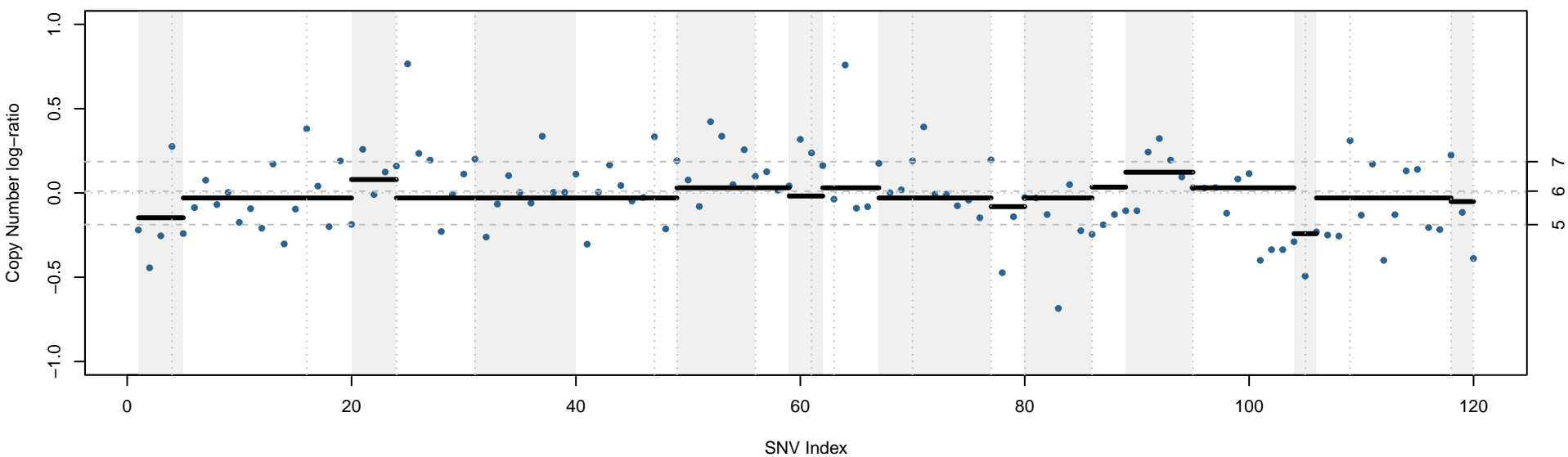
Purity: 0.53 Tumor ploidy: 5.942

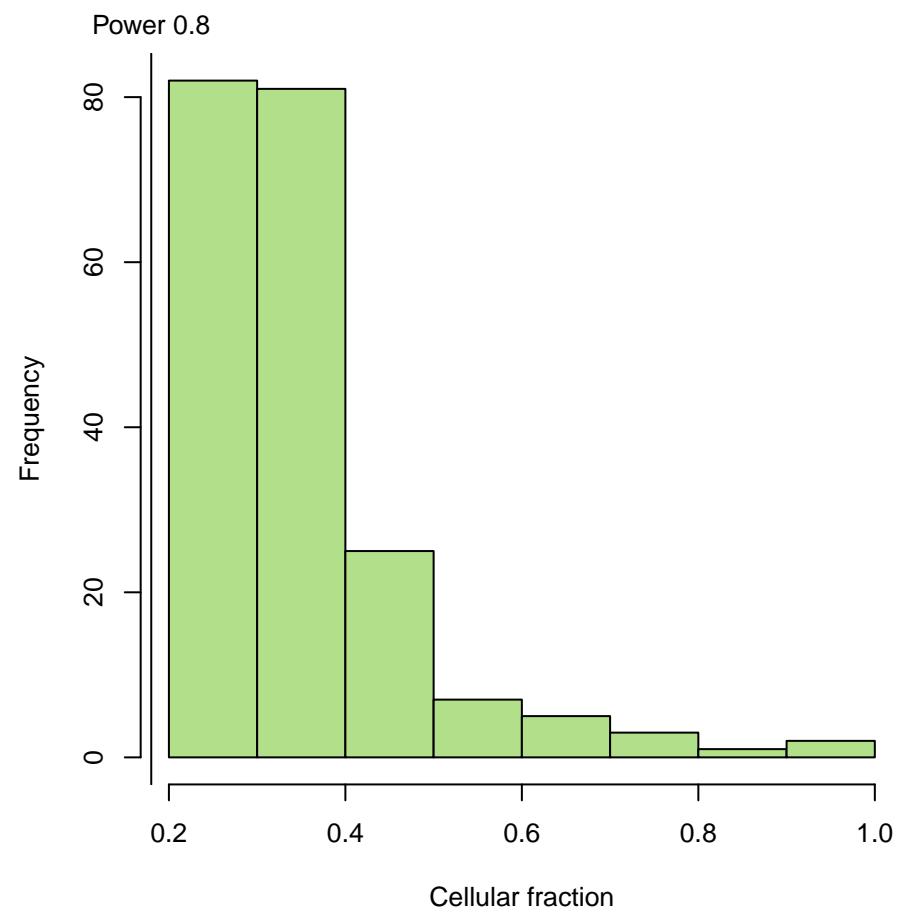
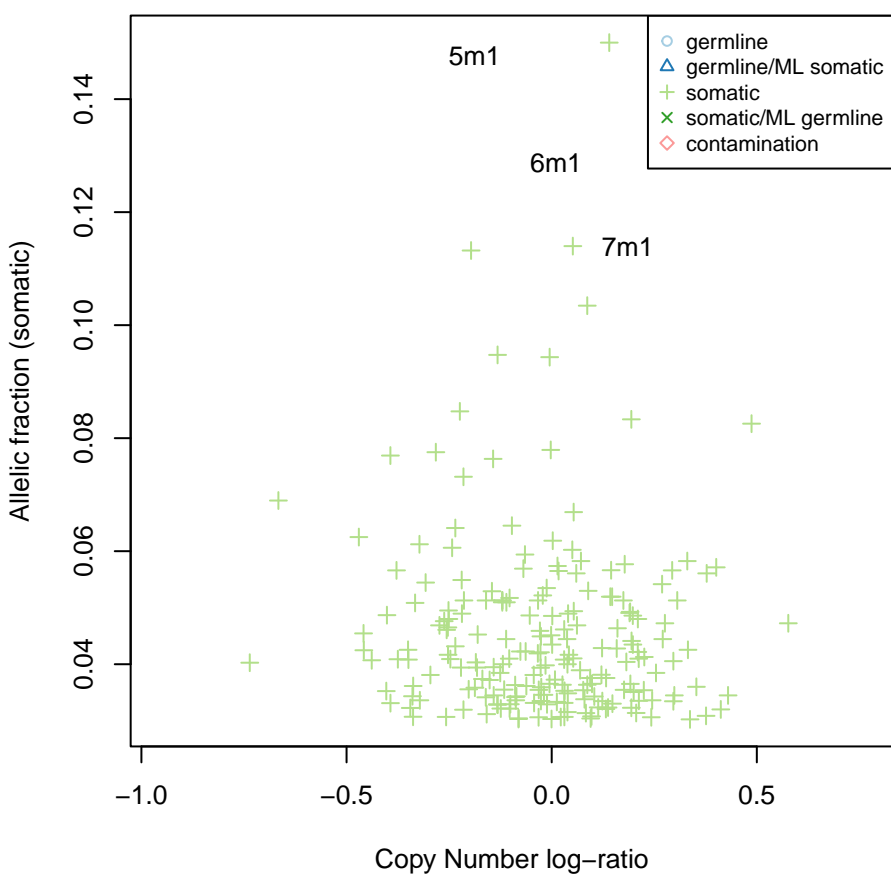
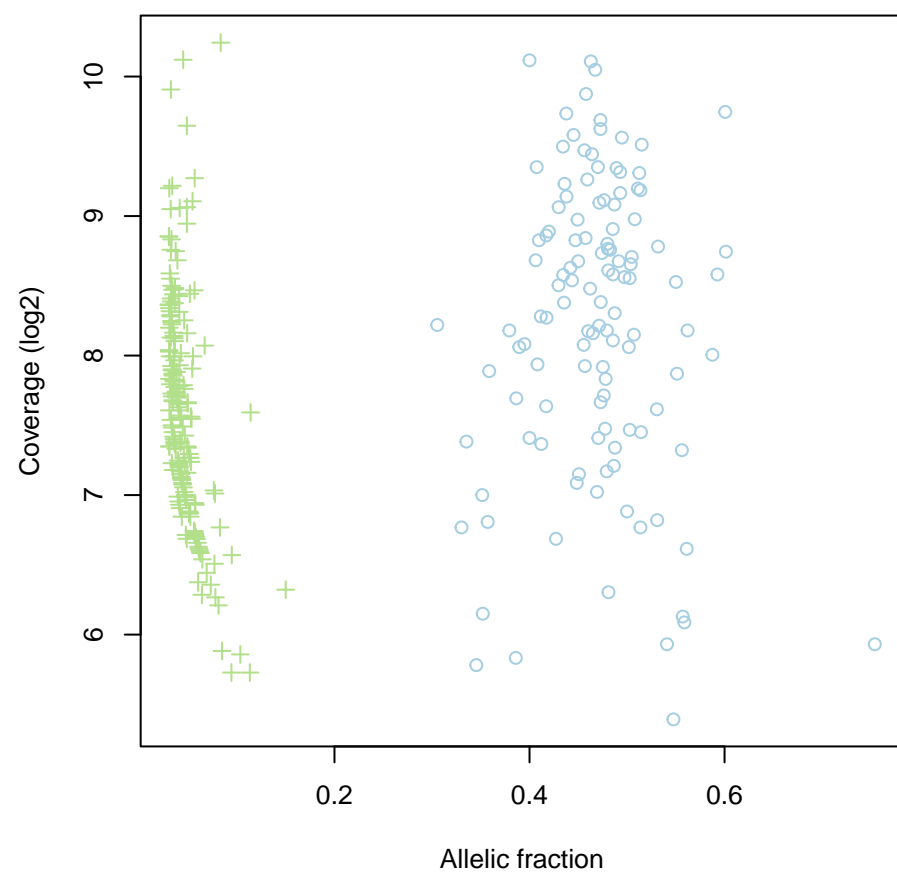
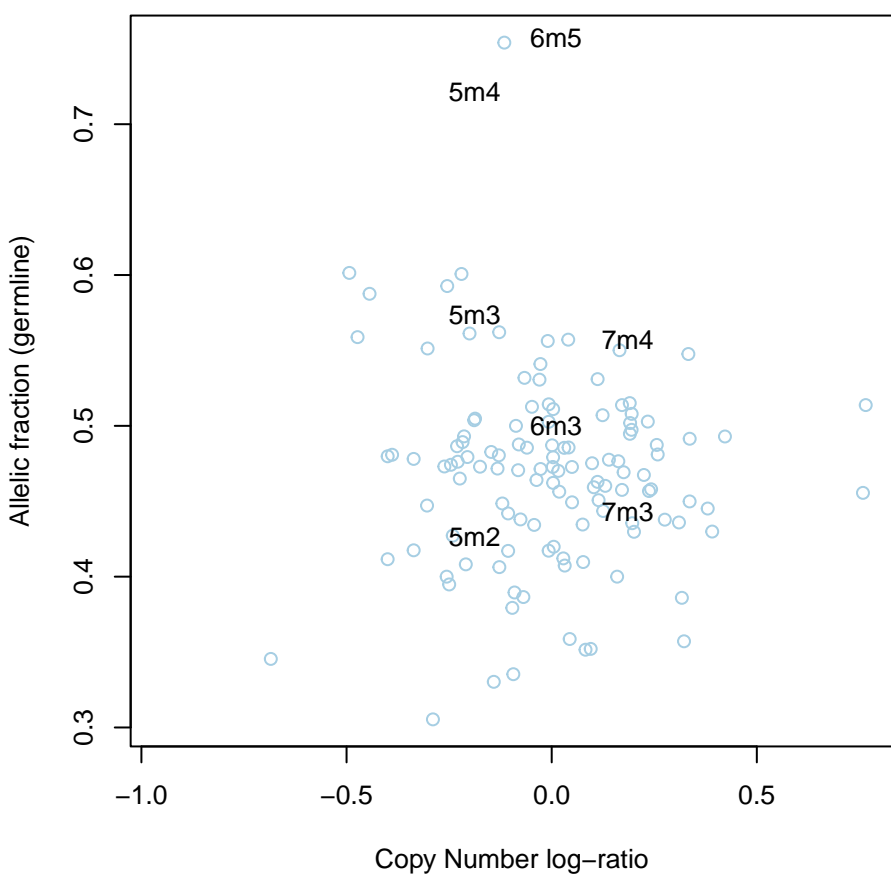


Purity: 0.53 Tumor ploidy: 5.942 SNV log-likelihood: -117.38 GoF: 86.2% Mean coverage: 406;281

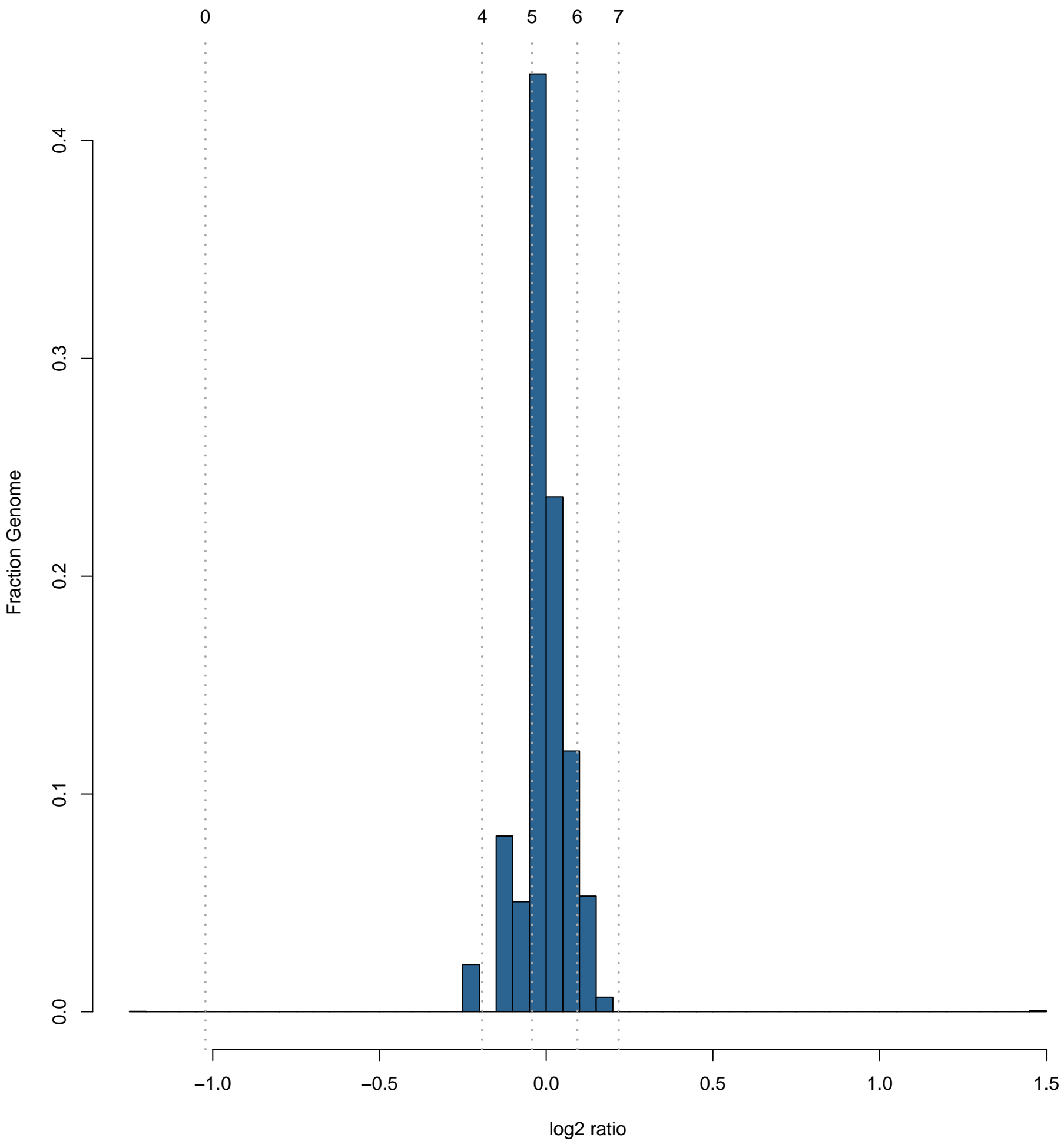


SCNA-fit log-likelihood: -15983.72

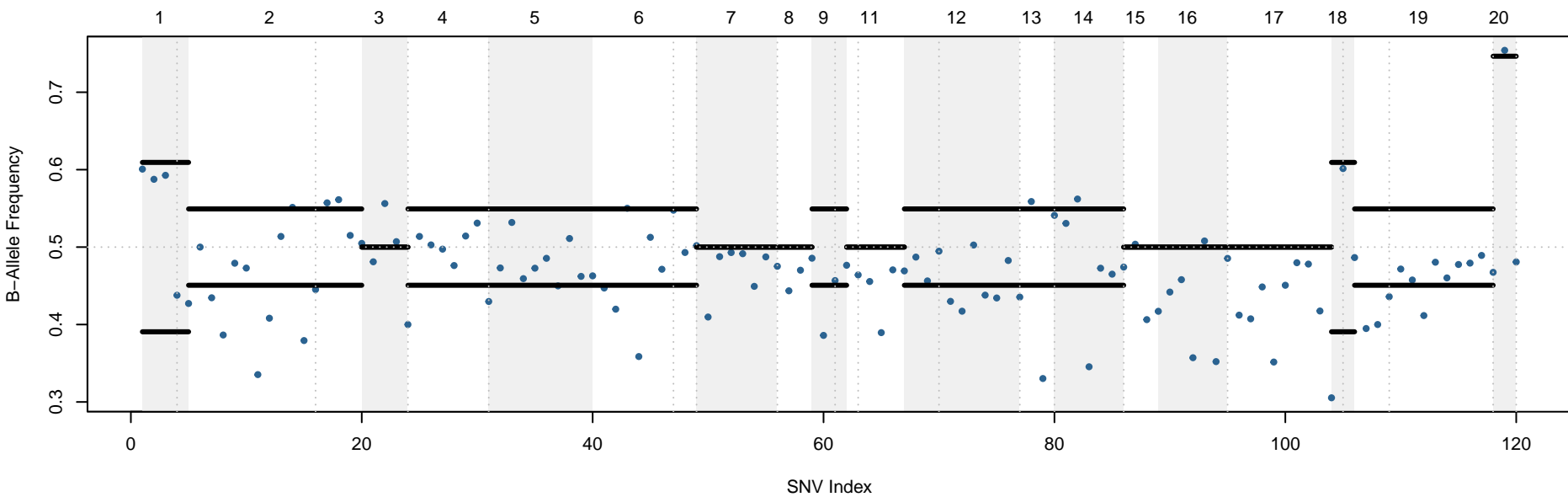




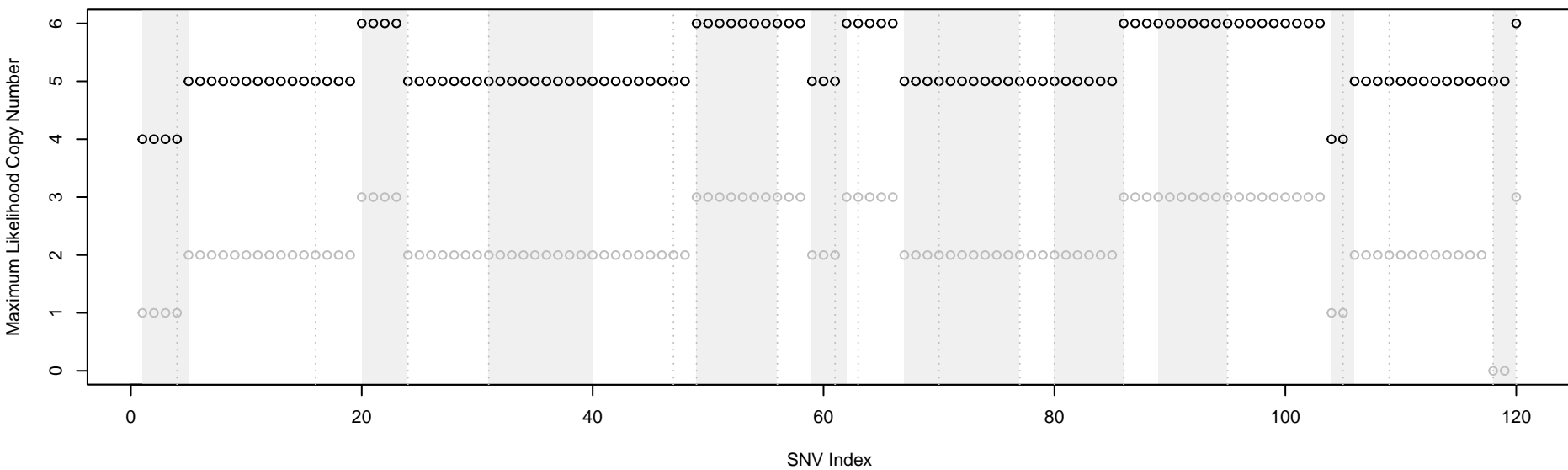
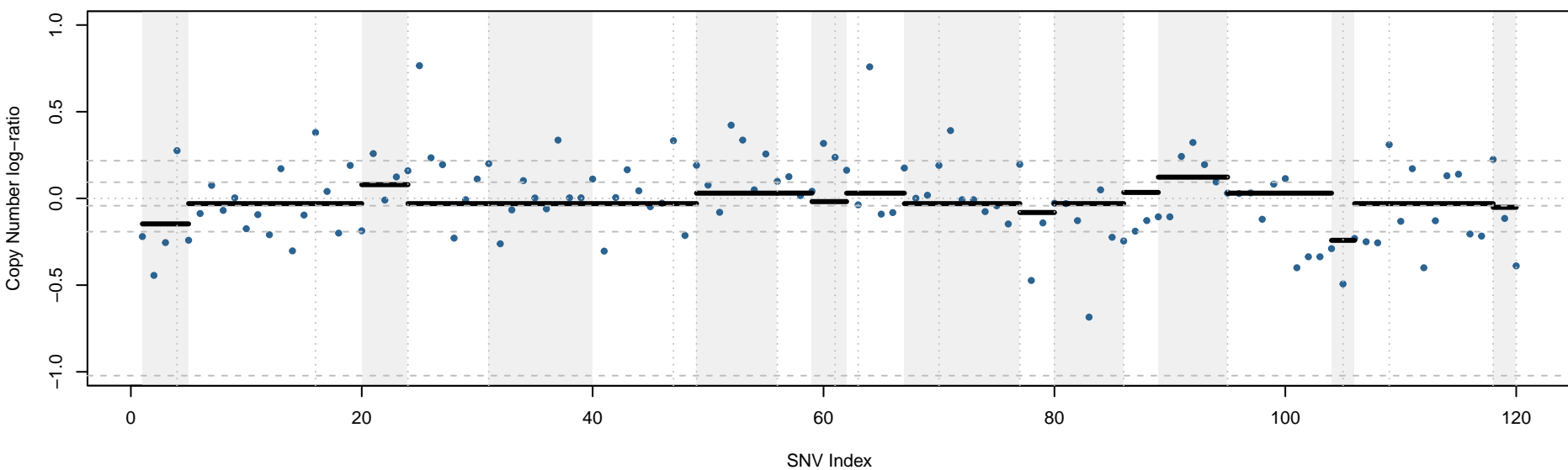
Purity: 0.28 Tumor ploidy: 5.301

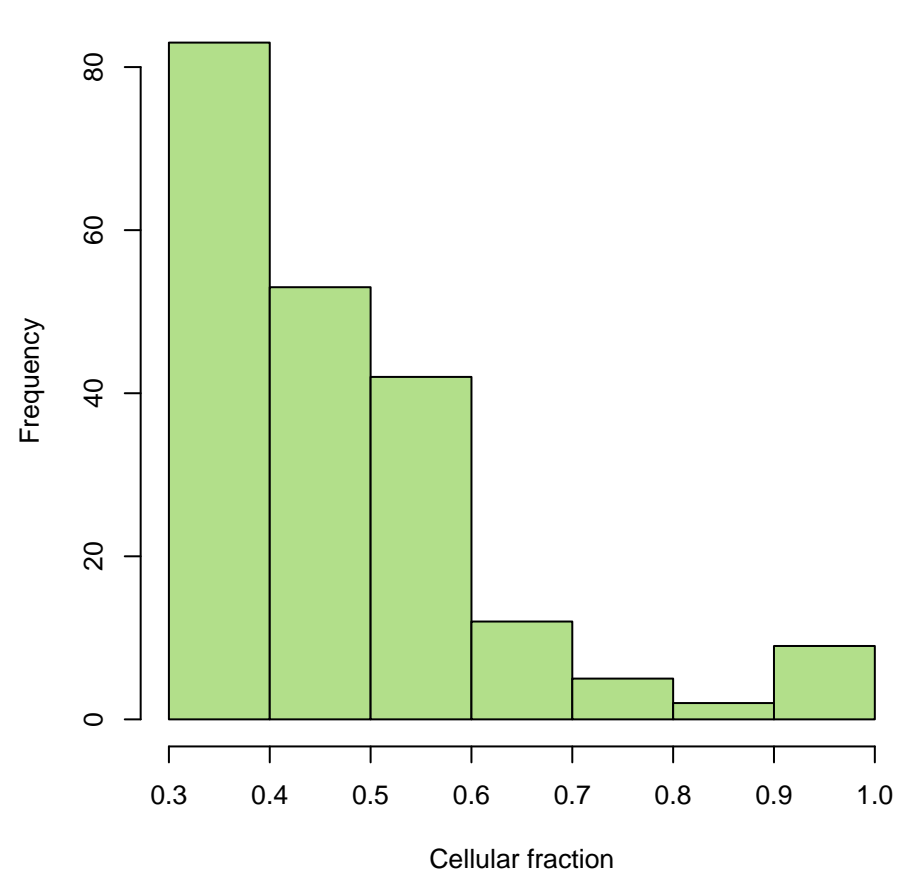
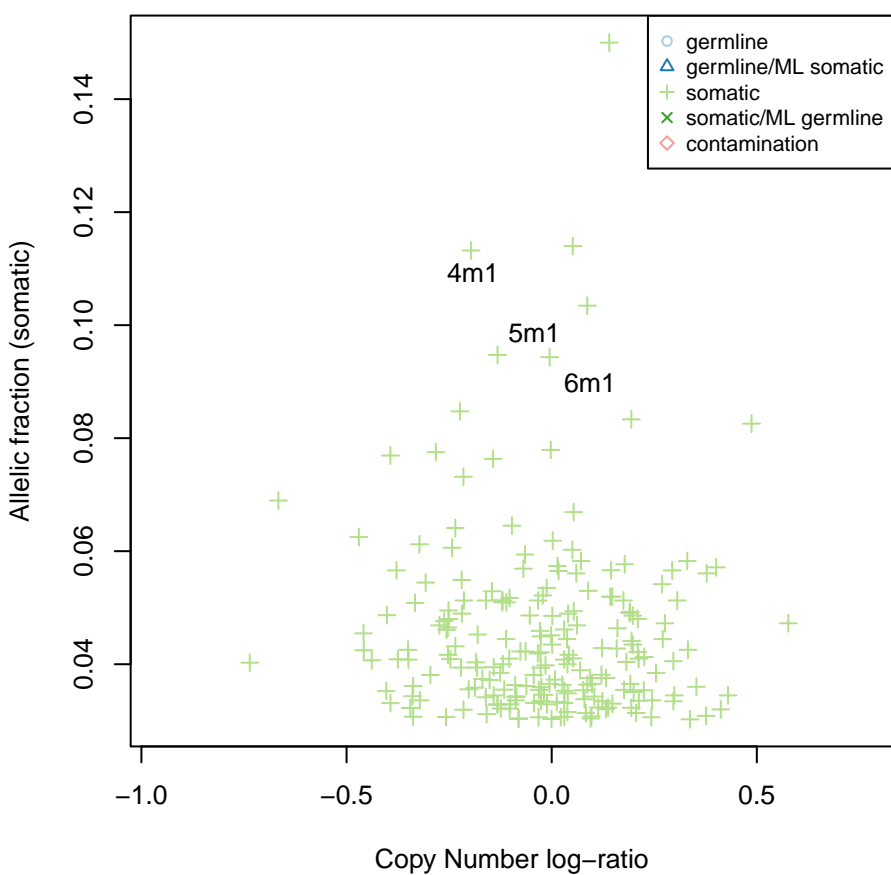
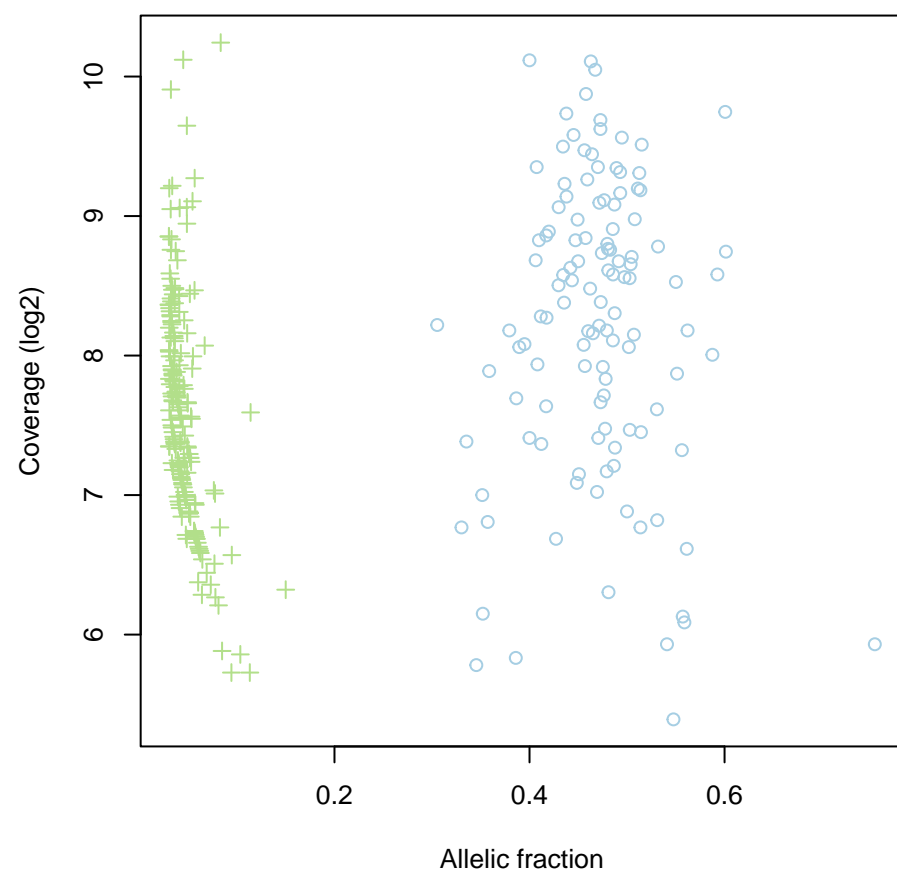
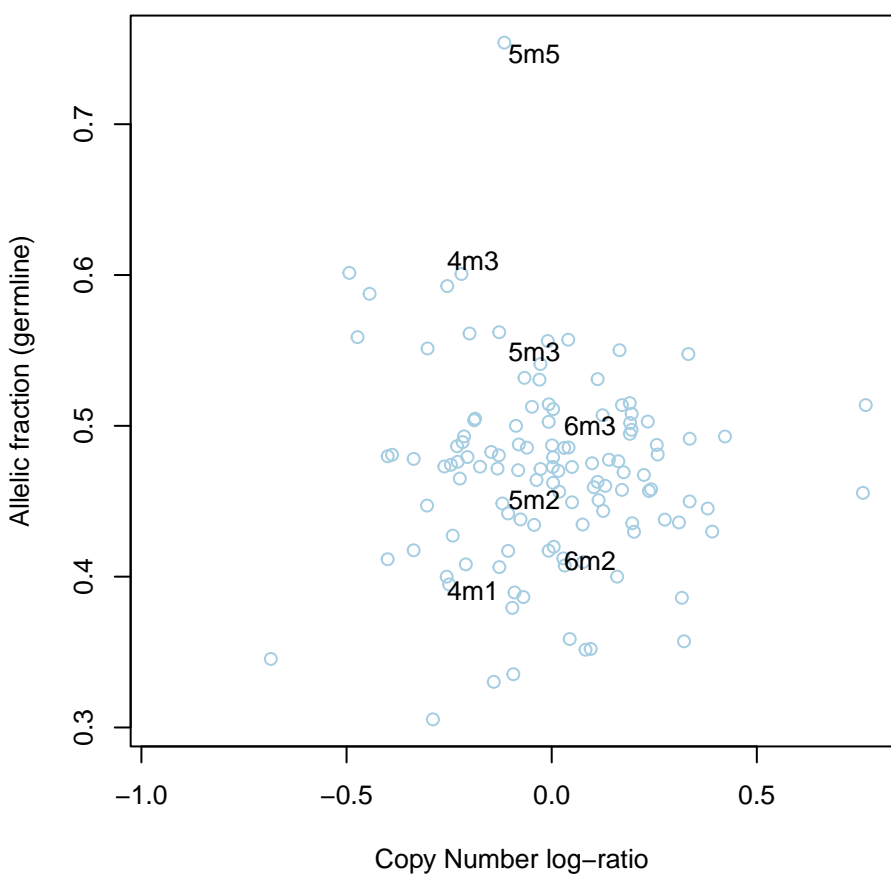


Purity: 0.28 Tumor ploidy: 5.301 SNV log-likelihood: -131.26 GoF: 94% Mean coverage: 406;281



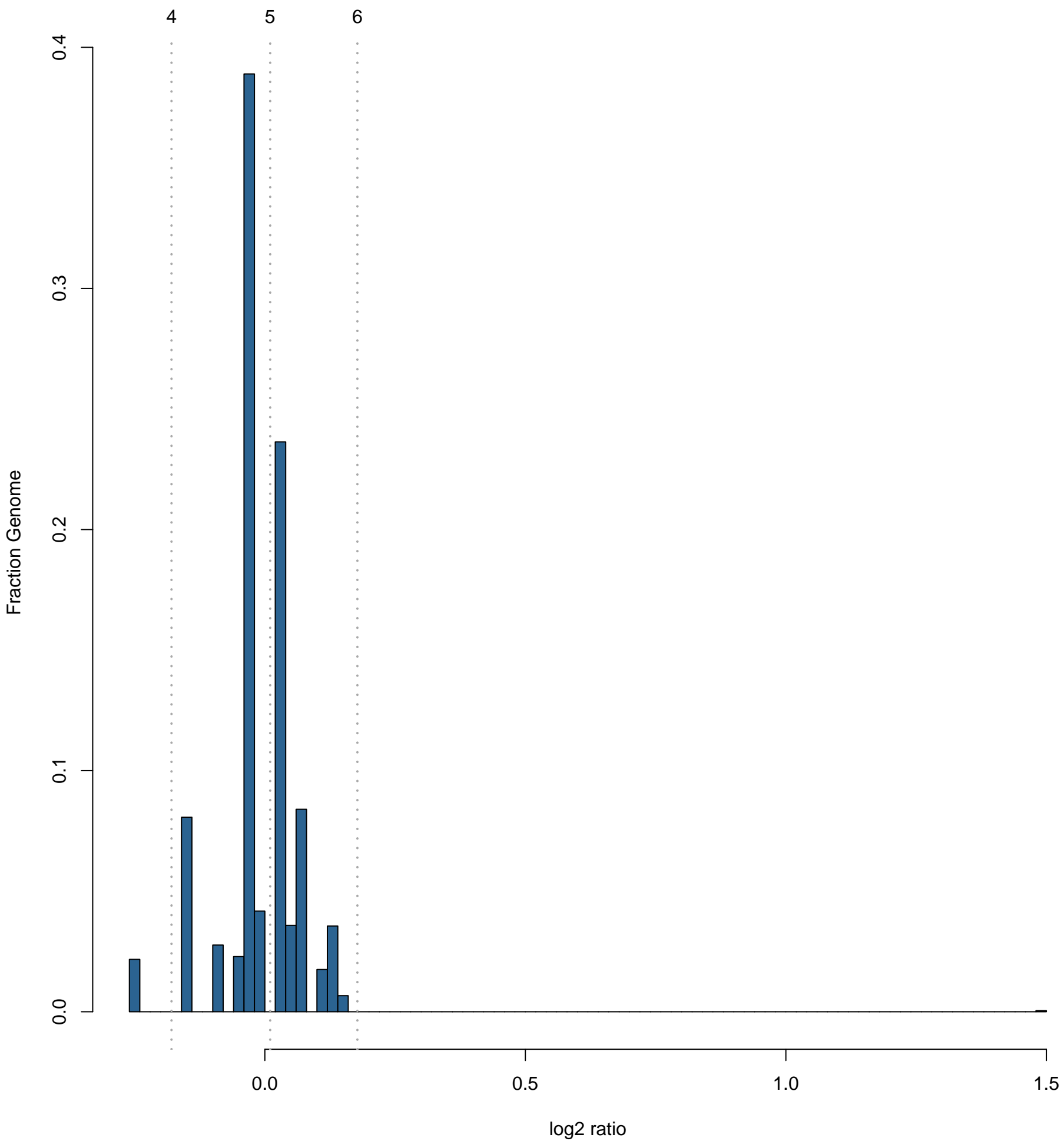
SCNA-fit log-likelihood: -15965.16



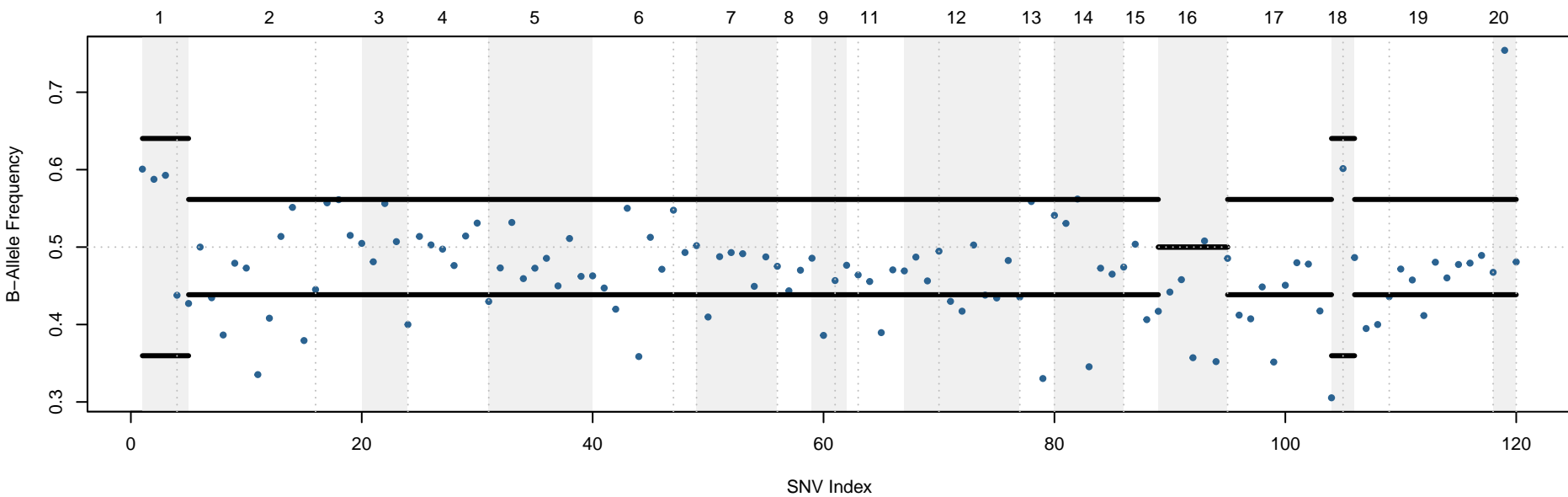




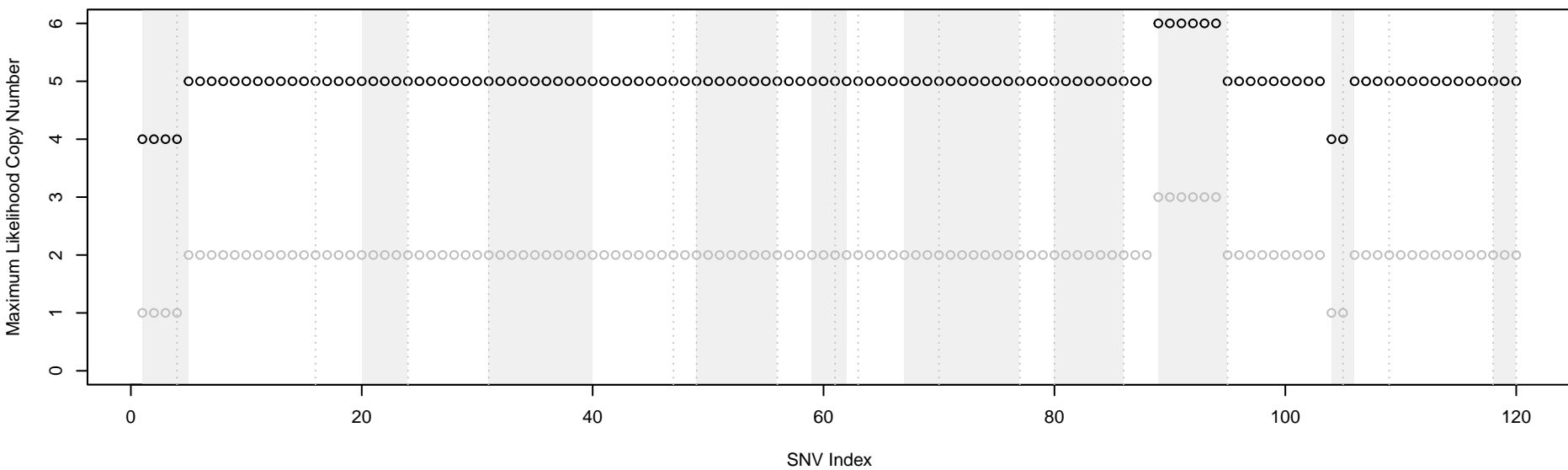
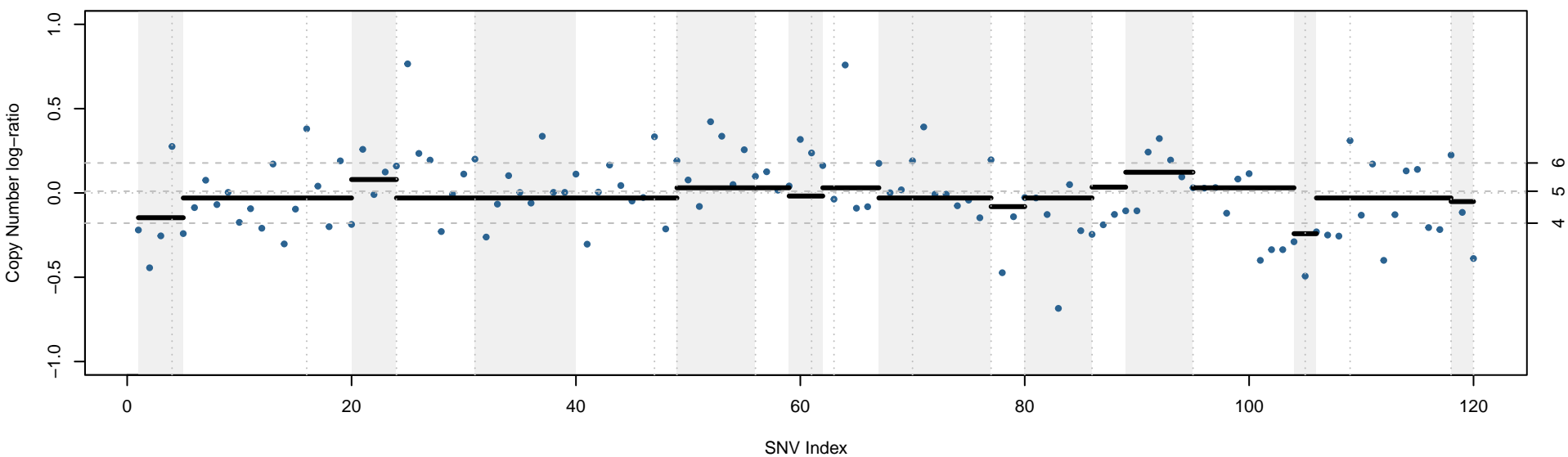
Purity: 0.39 Tumor ploidy: 4.943

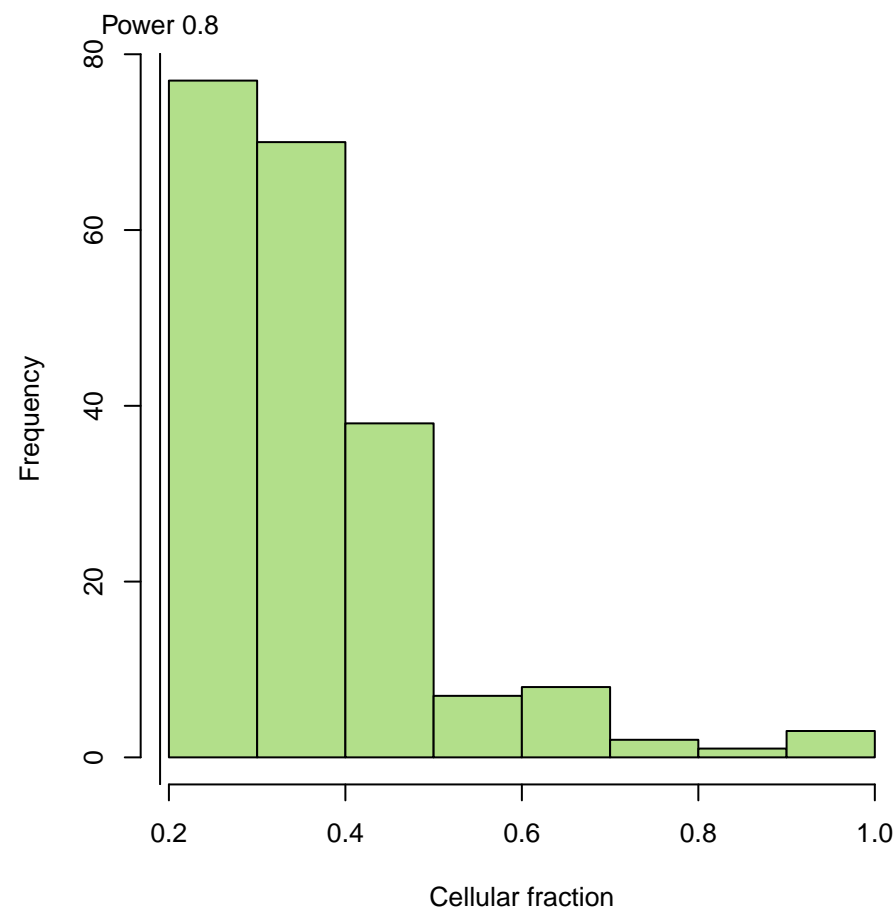
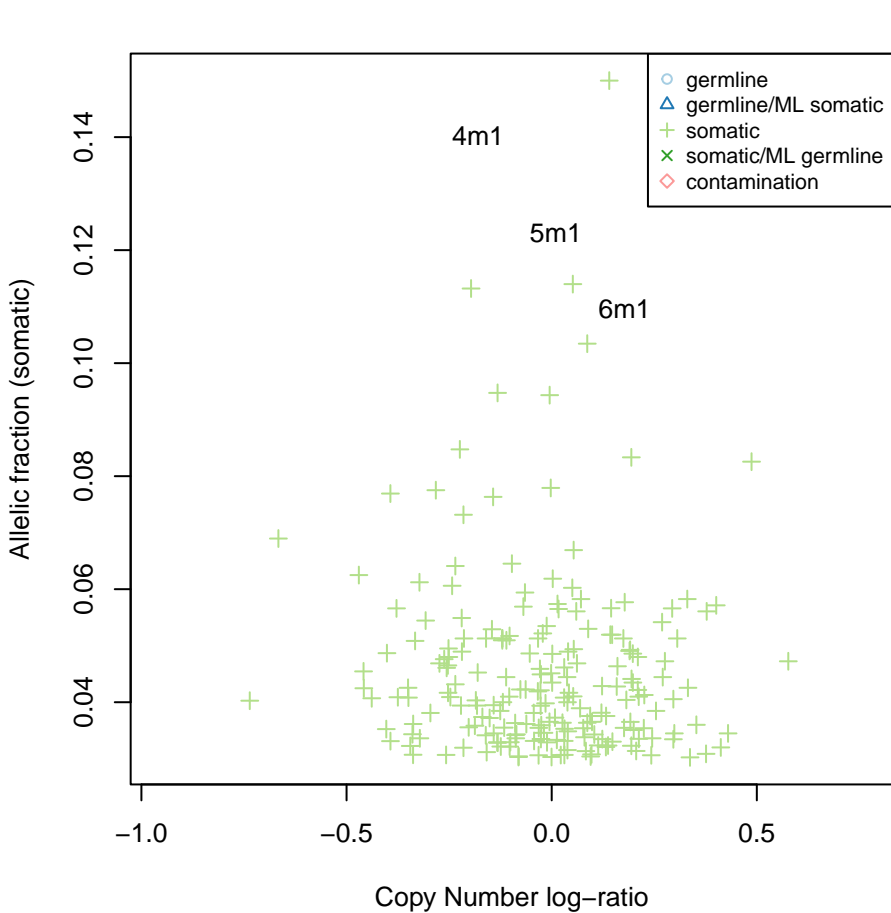
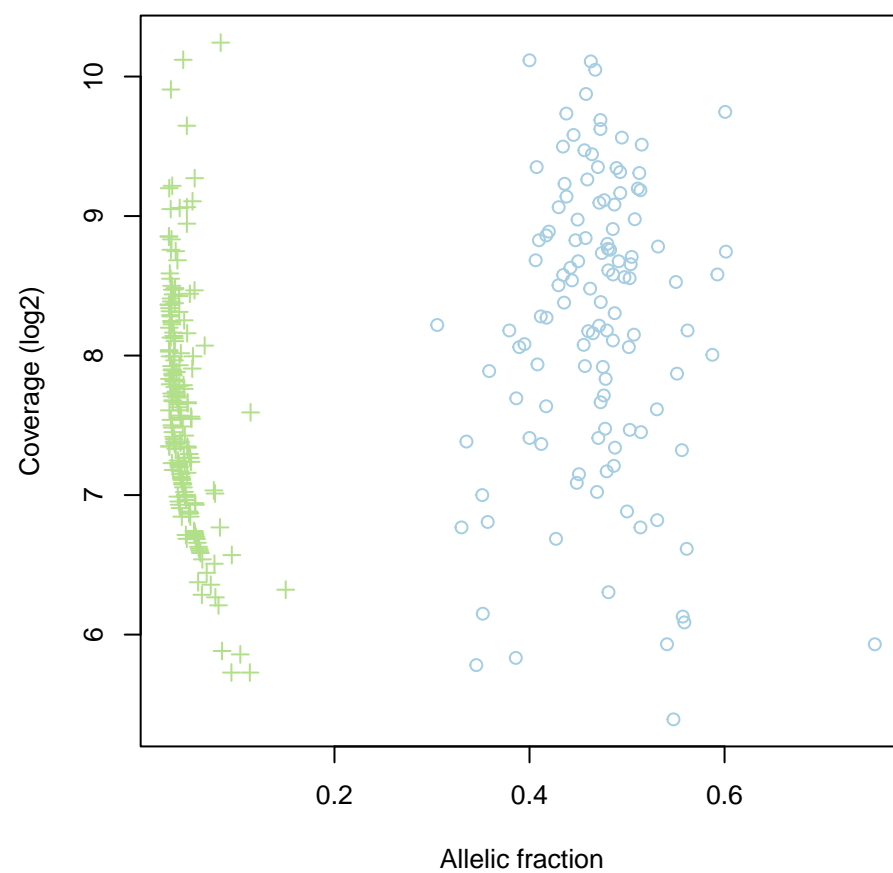
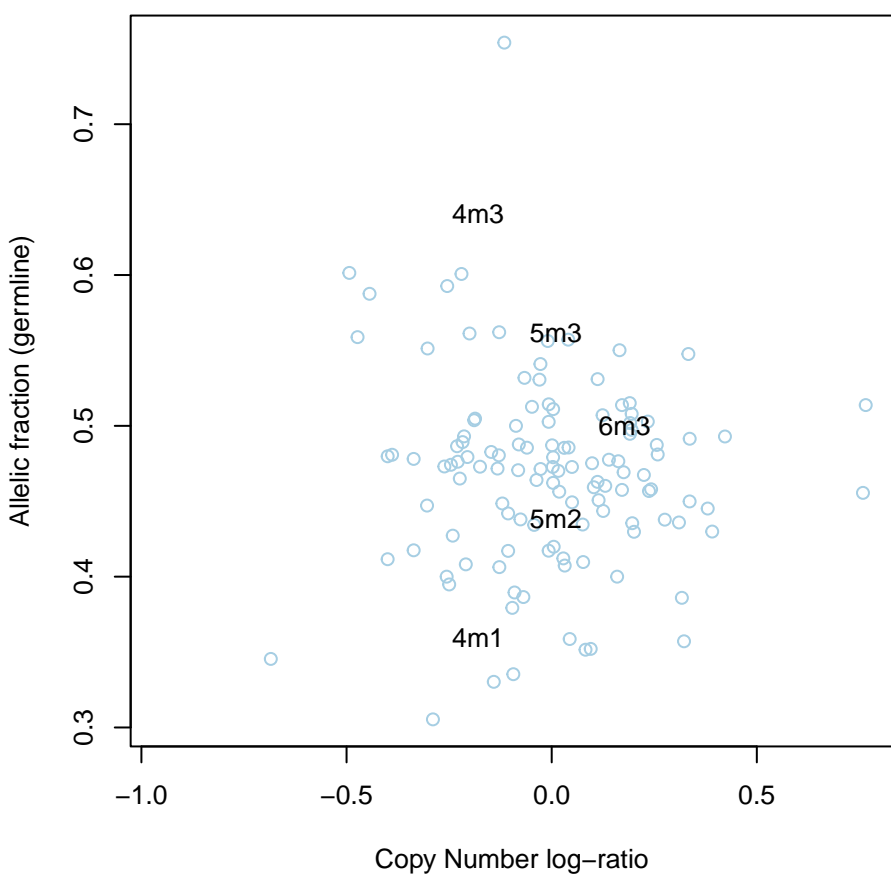


Purity: 0.39 Tumor ploidy: 4.943 SNV log-likelihood: -138.44 GoF: 88.1% Mean coverage: 406;281

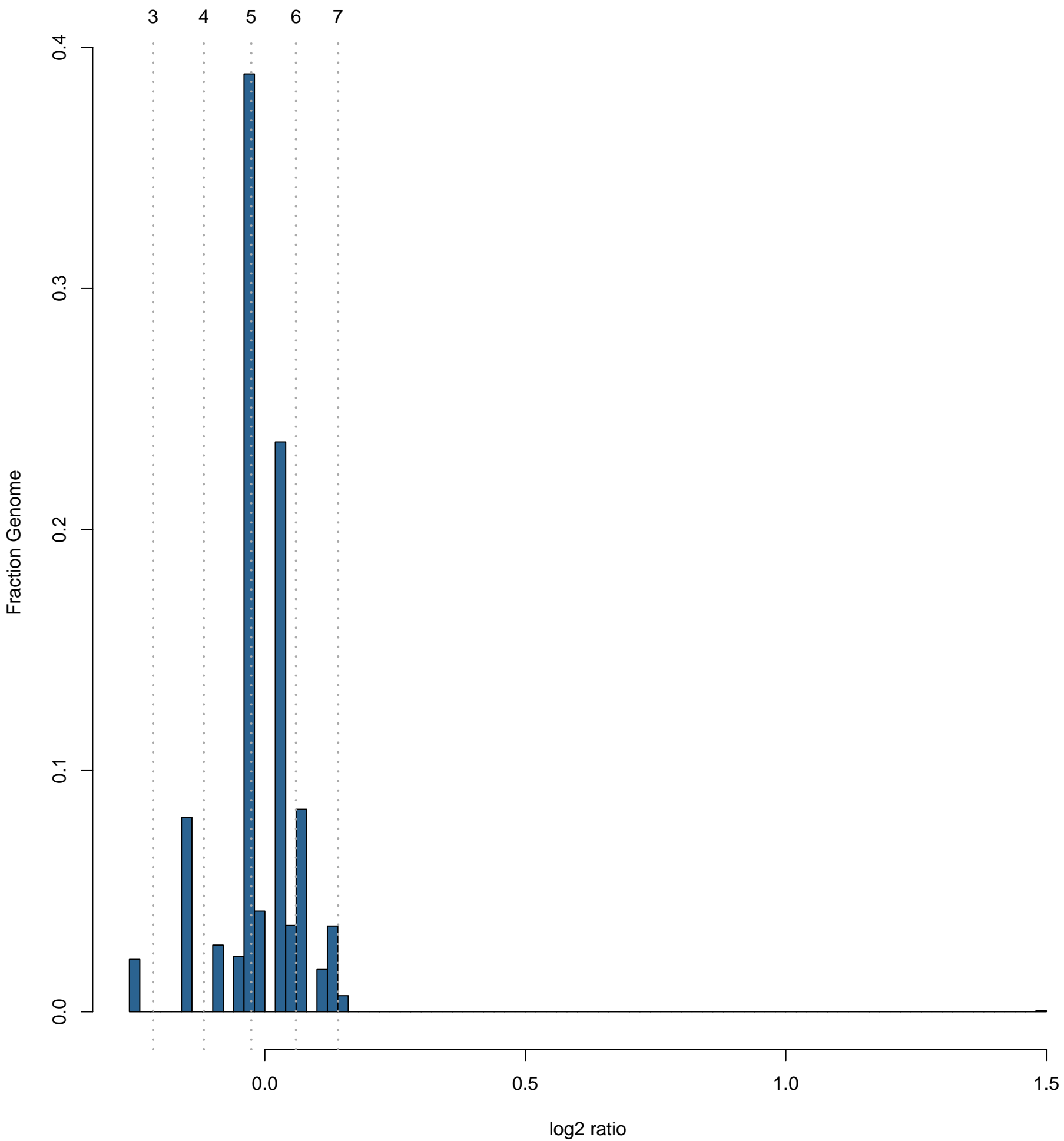


SCNA-fit log-likelihood: -15980.84

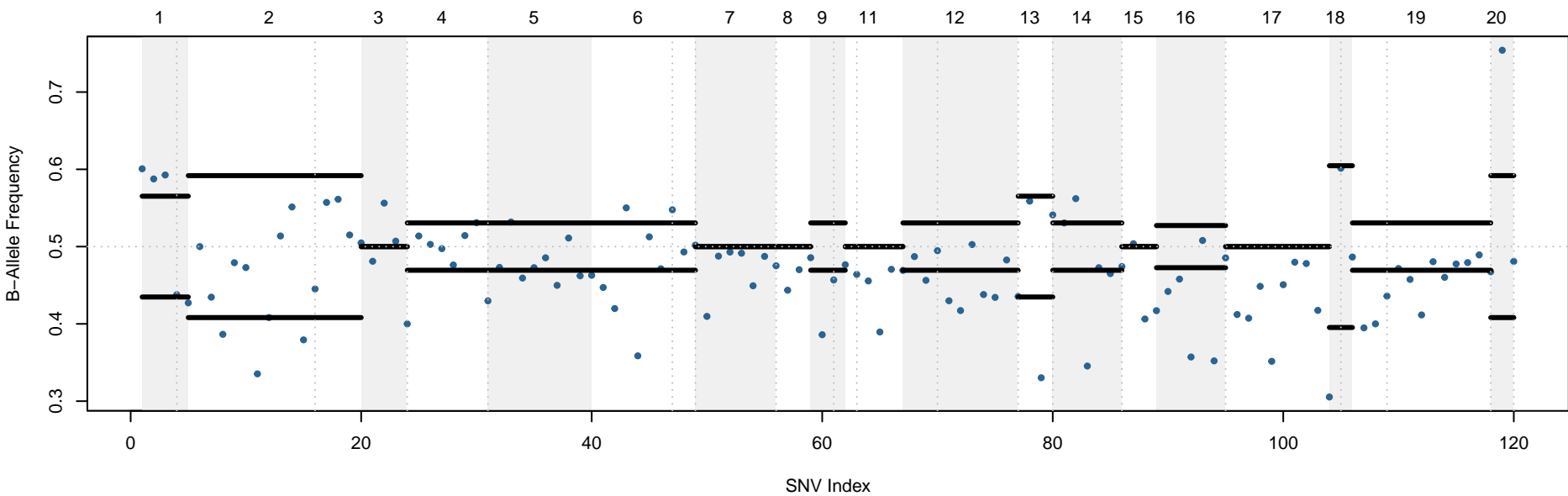




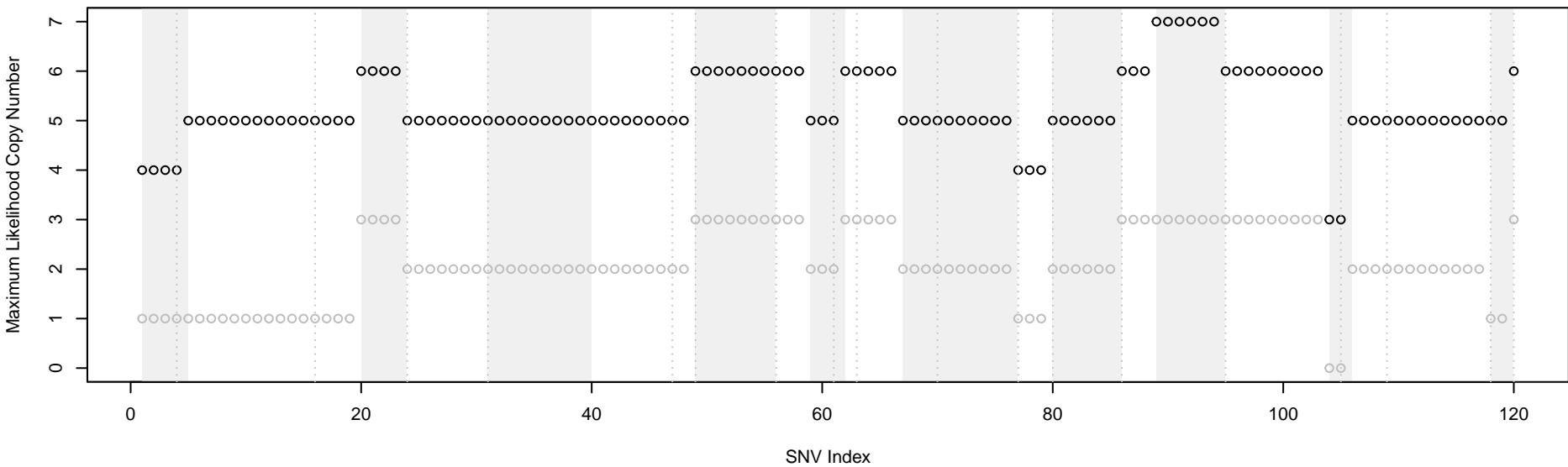
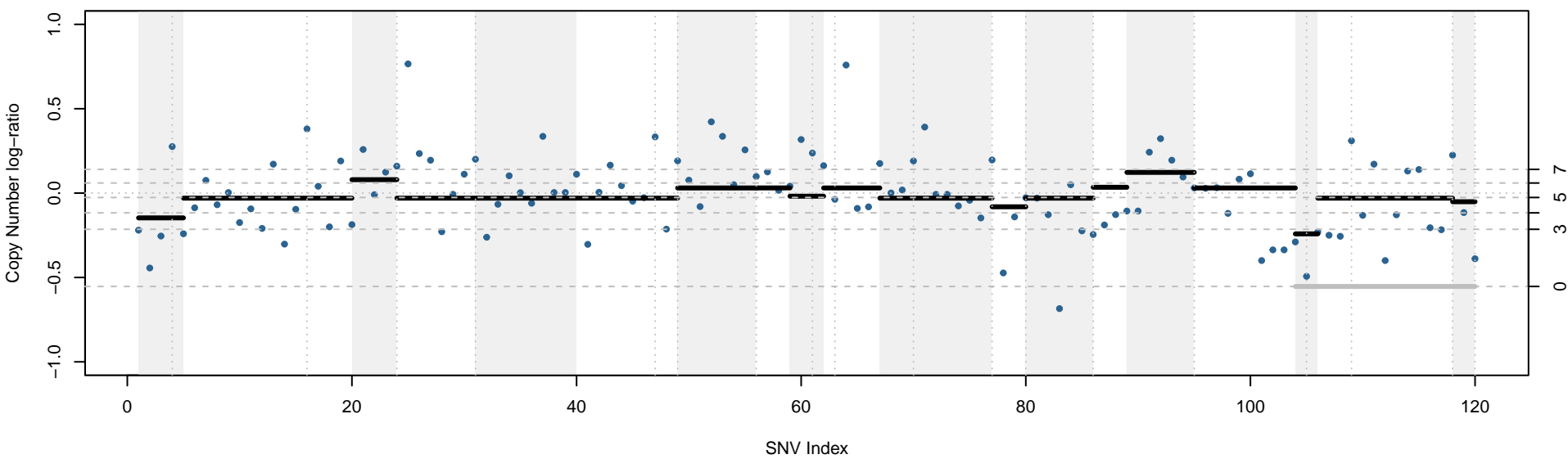
Purity: 0.15 Tumor ploidy: 5.298

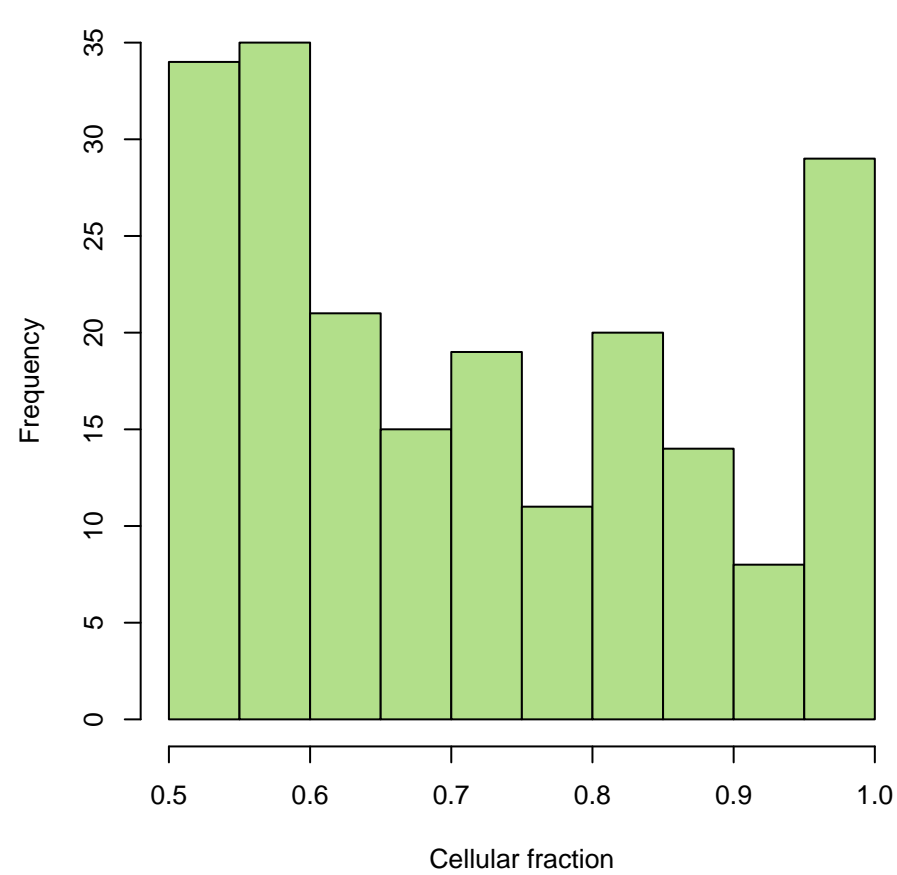
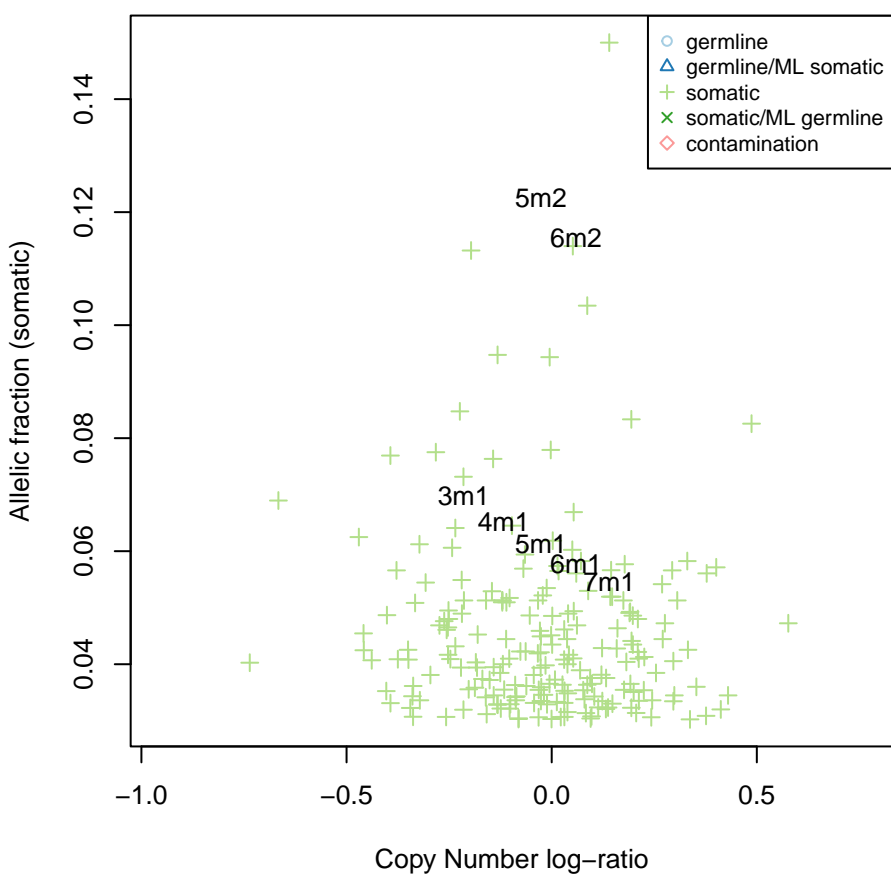
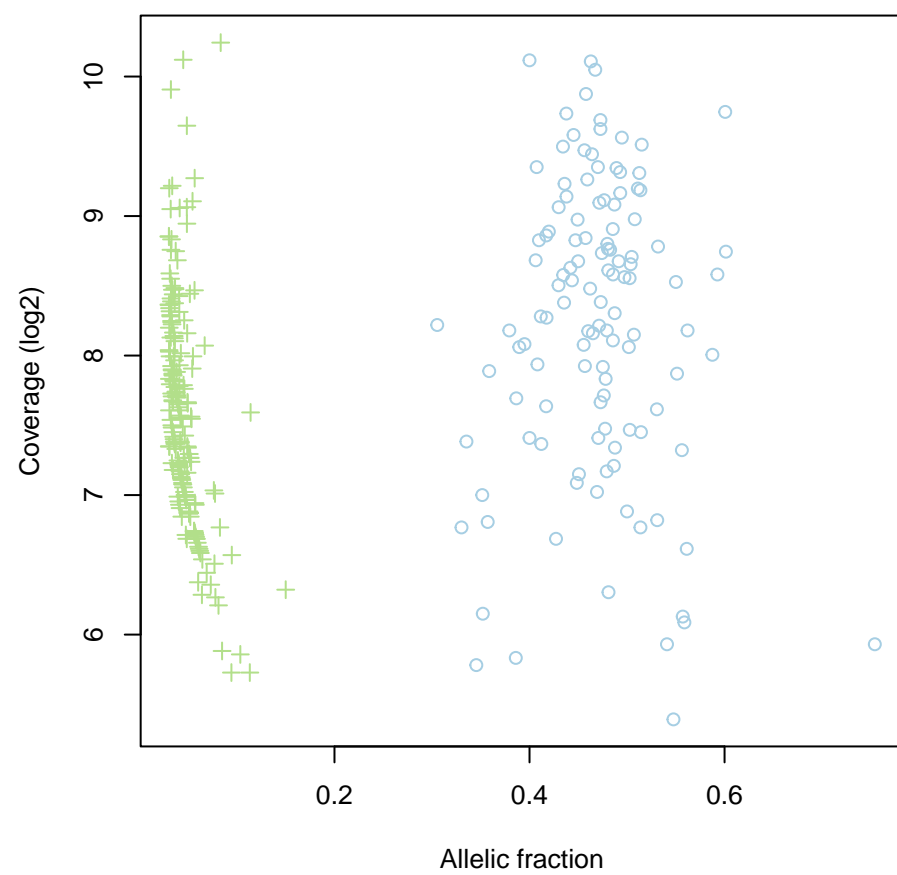
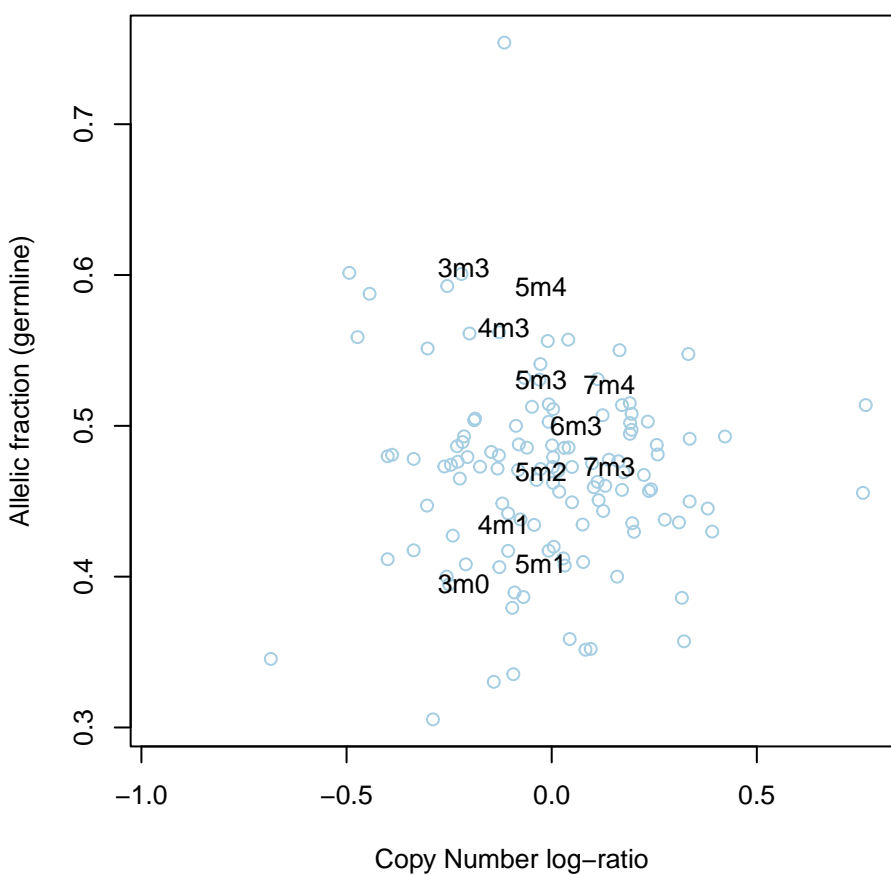


Purity: 0.15 Tumor ploidy: 5.298 SNV log-likelihood: -158.66 GoF: 97.3% Mean coverage: 406;281

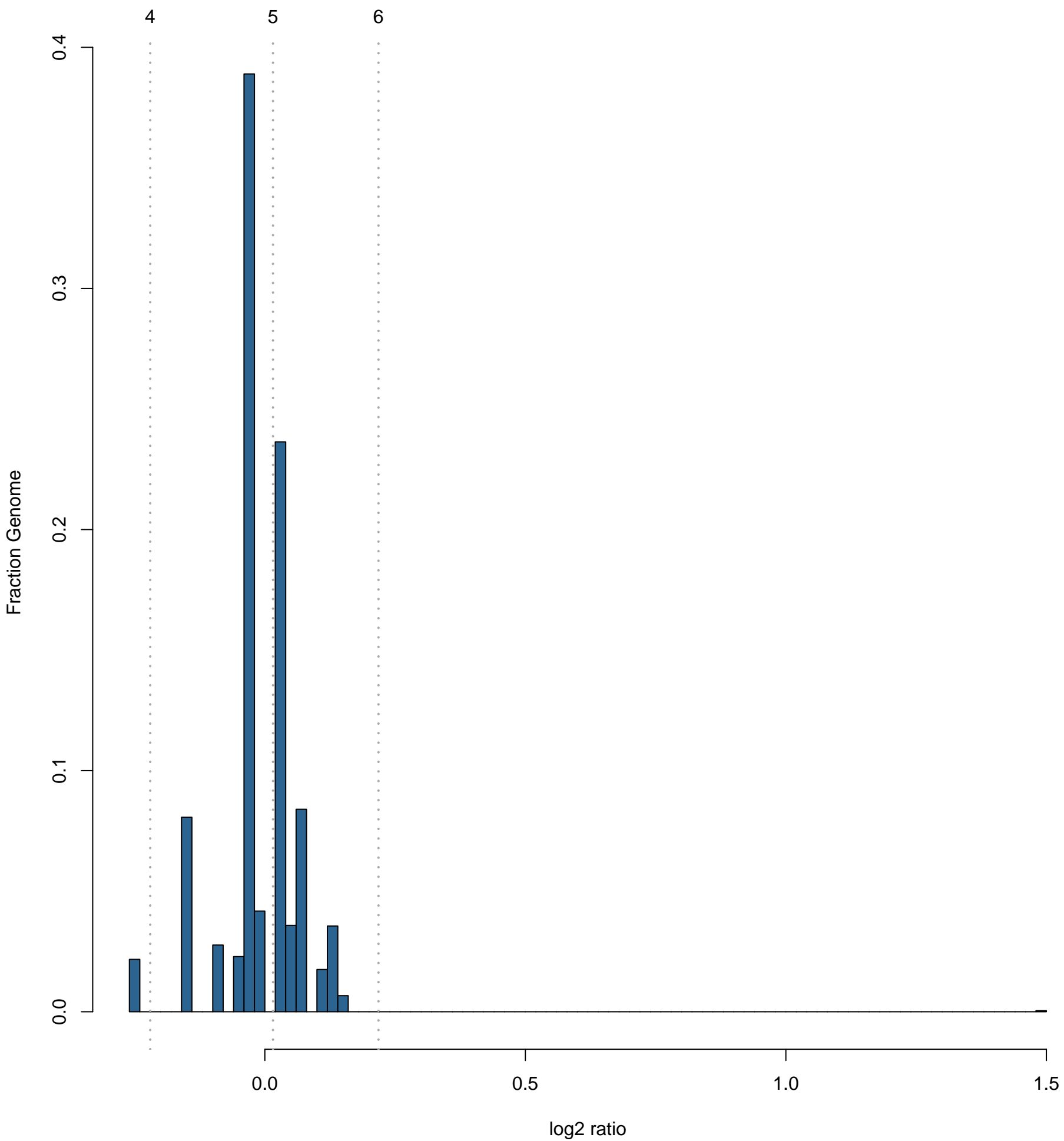


SCNA-fit log-likelihood: -15930.7

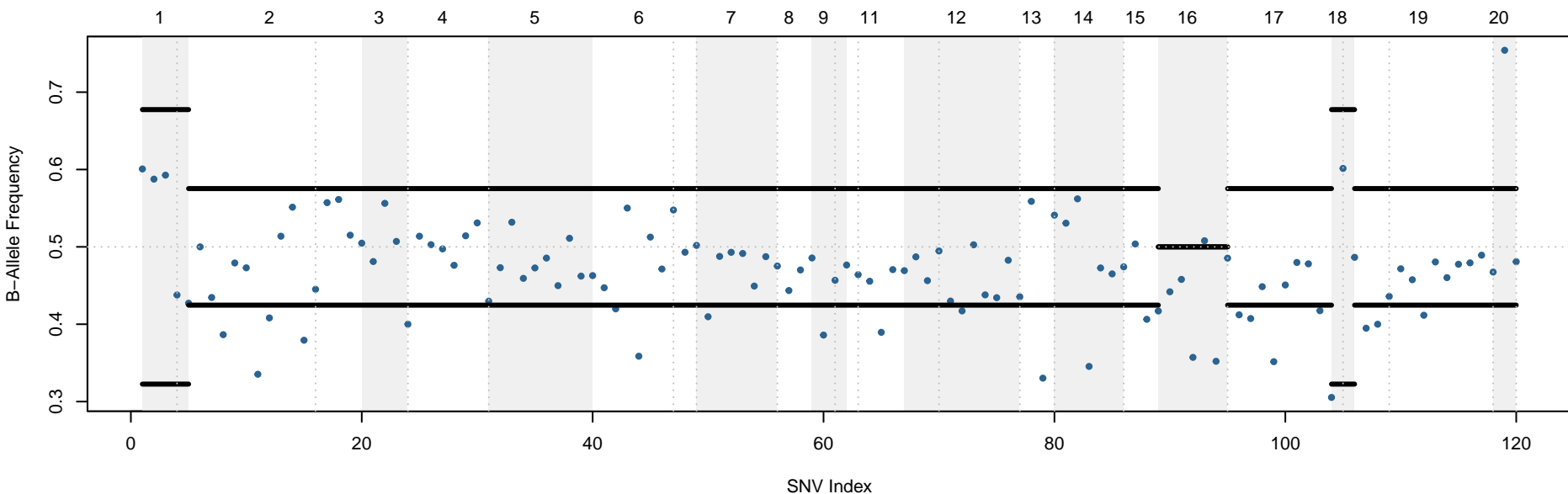




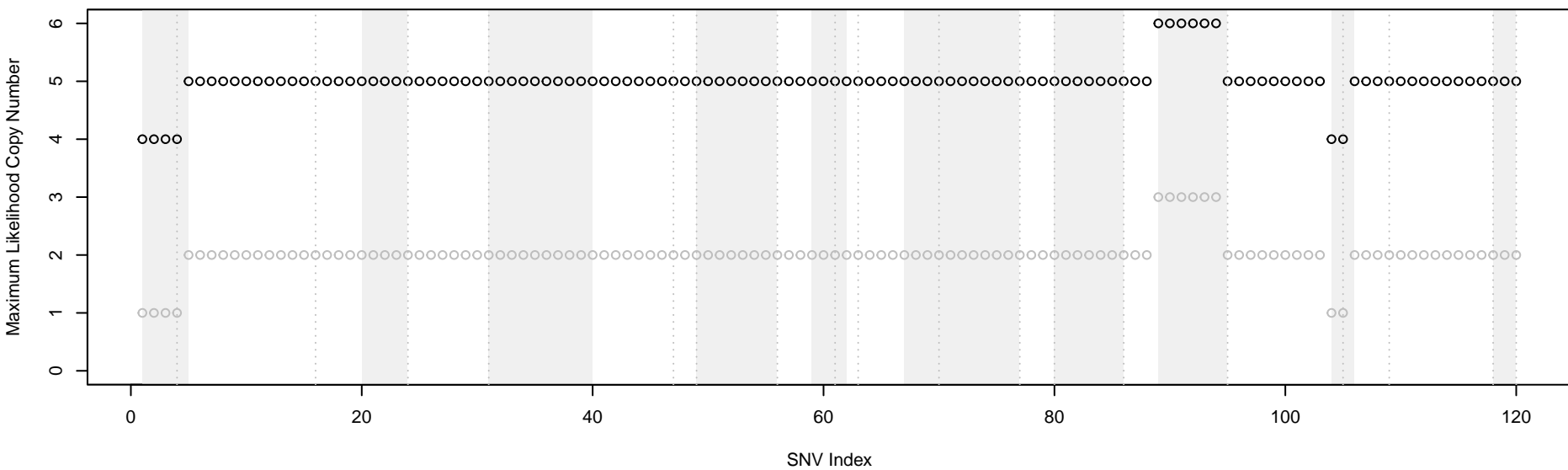
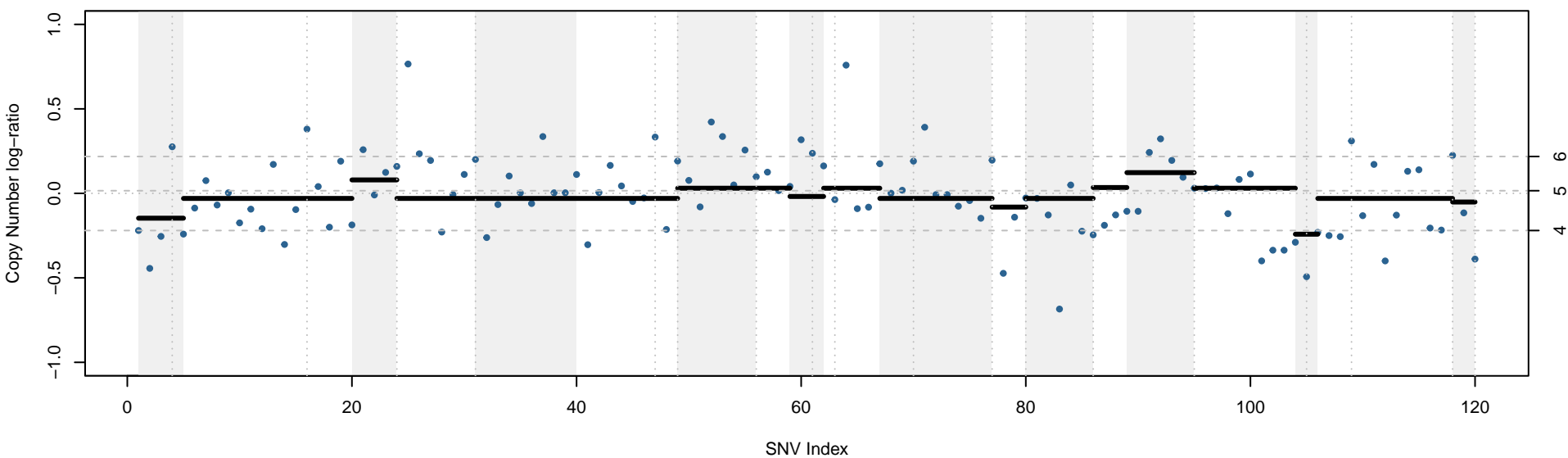
Purity: 0.55 Tumor ploidy: 4.928



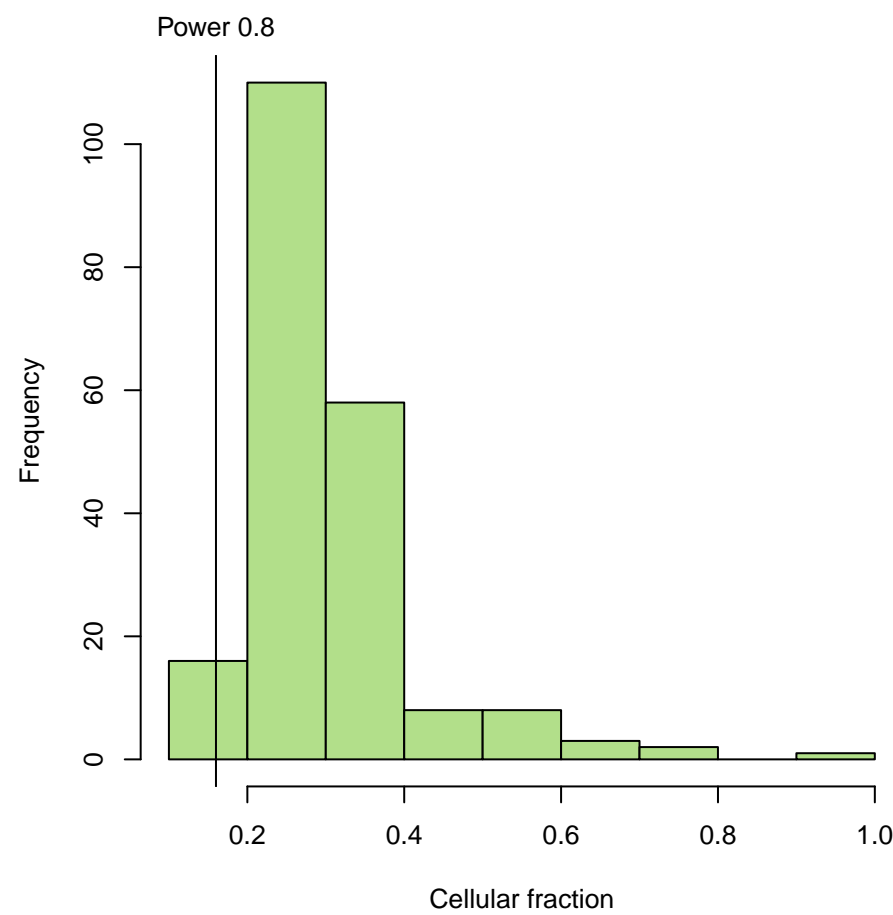
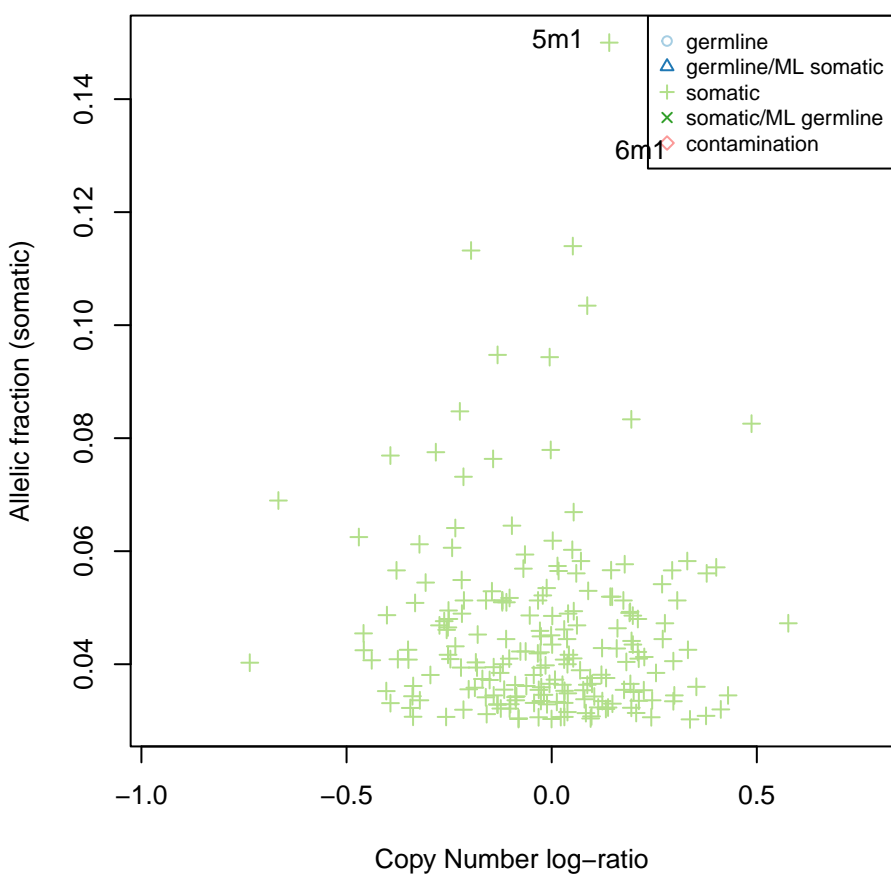
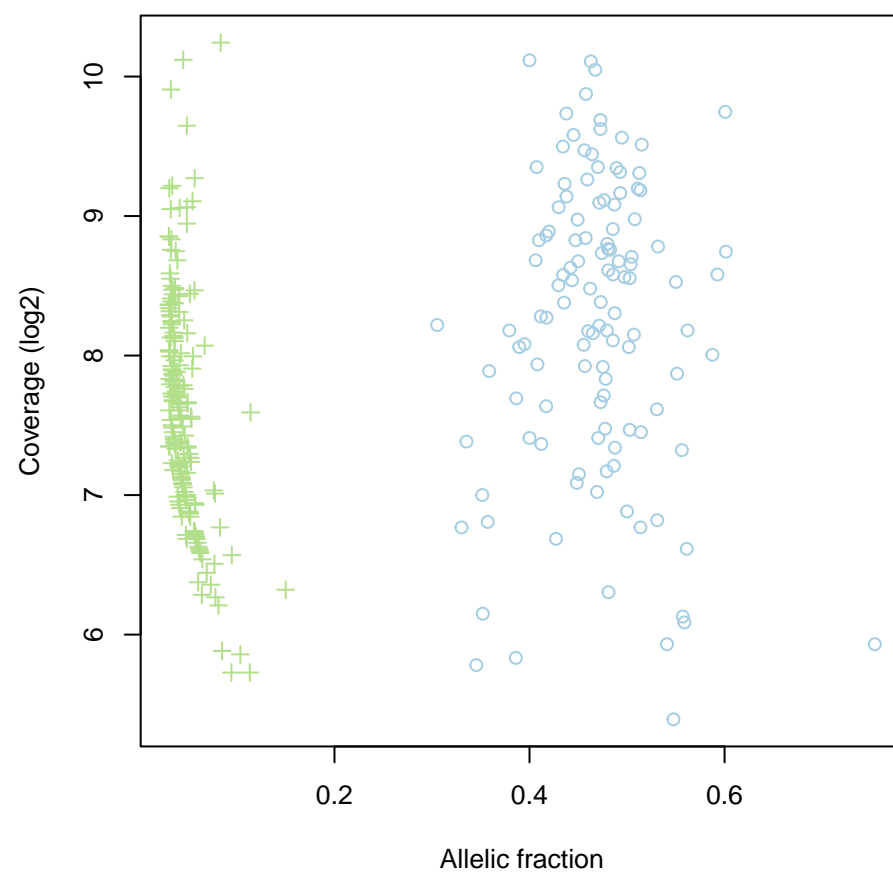
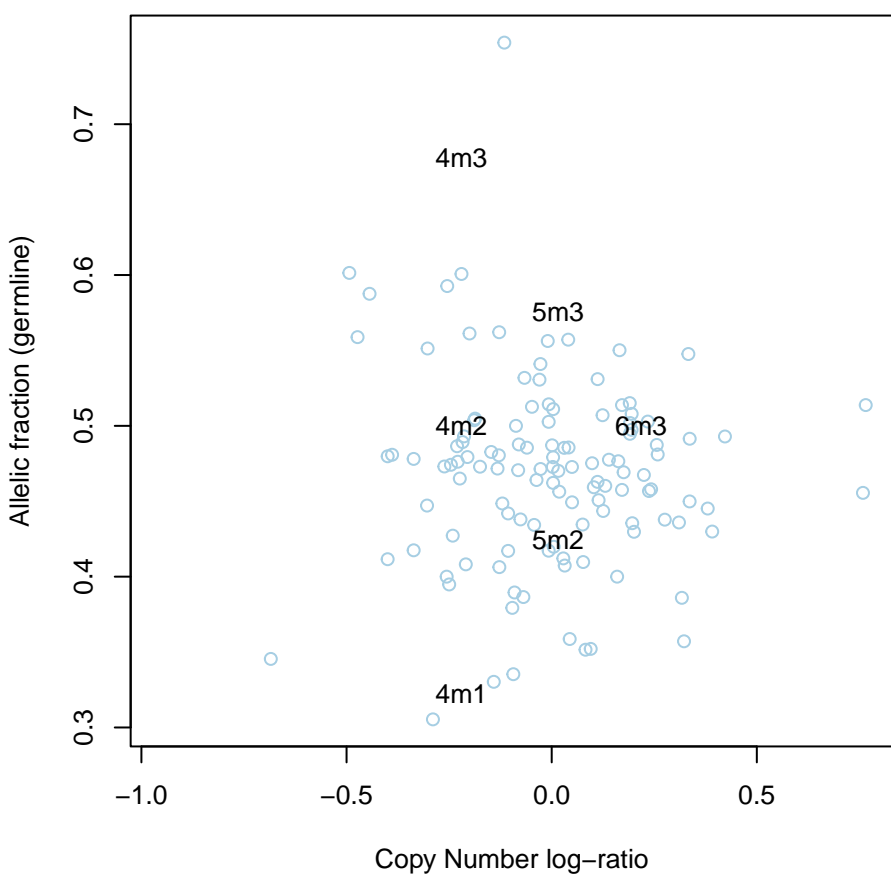
Purity: 0.55 Tumor ploidy: 4.928 SNV log-likelihood: -229.51 GoF: 79.3% Mean coverage: 406,281



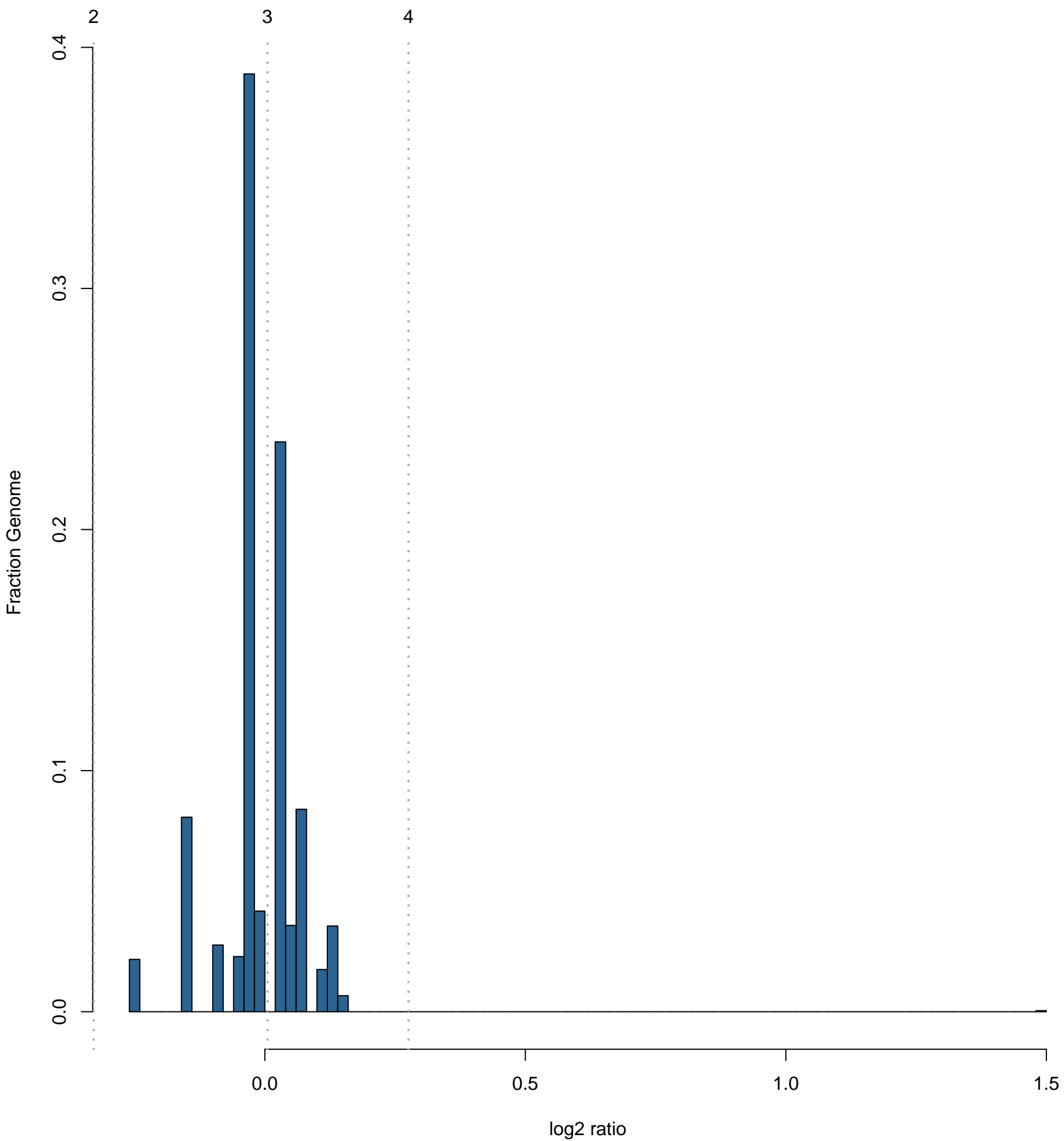
SCNA-fit log-likelihood: -16003.8



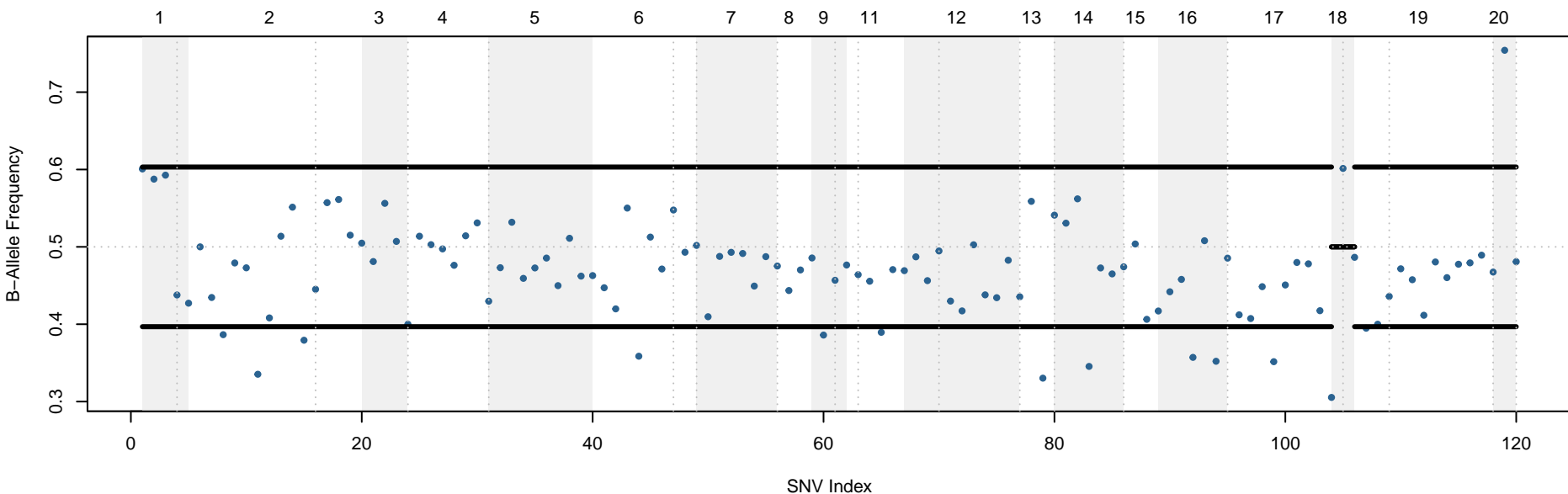




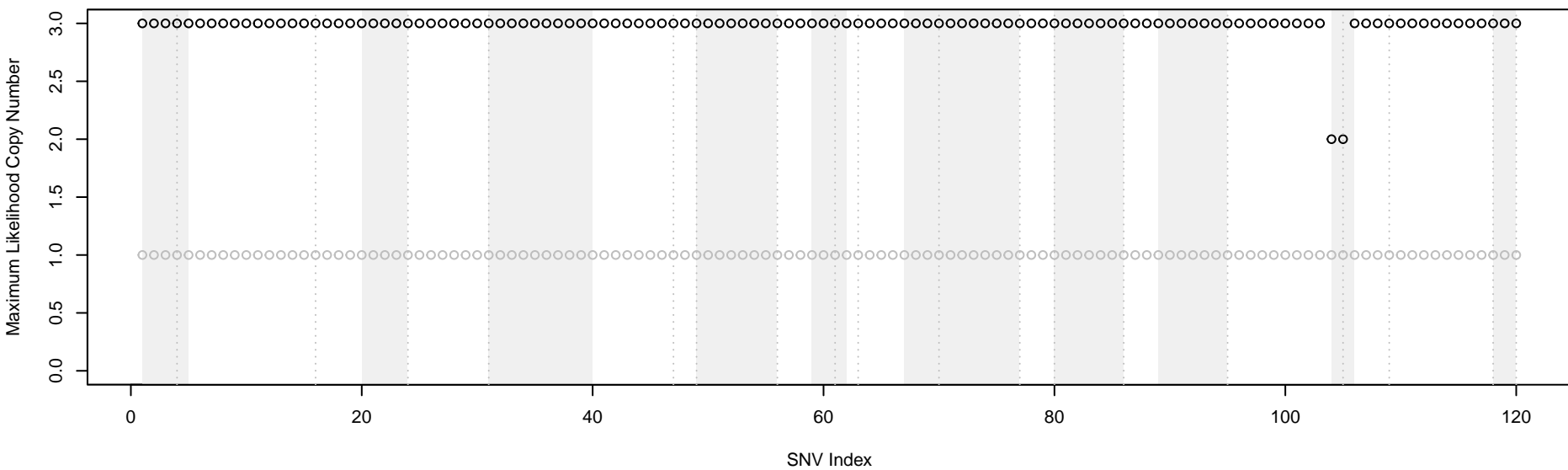
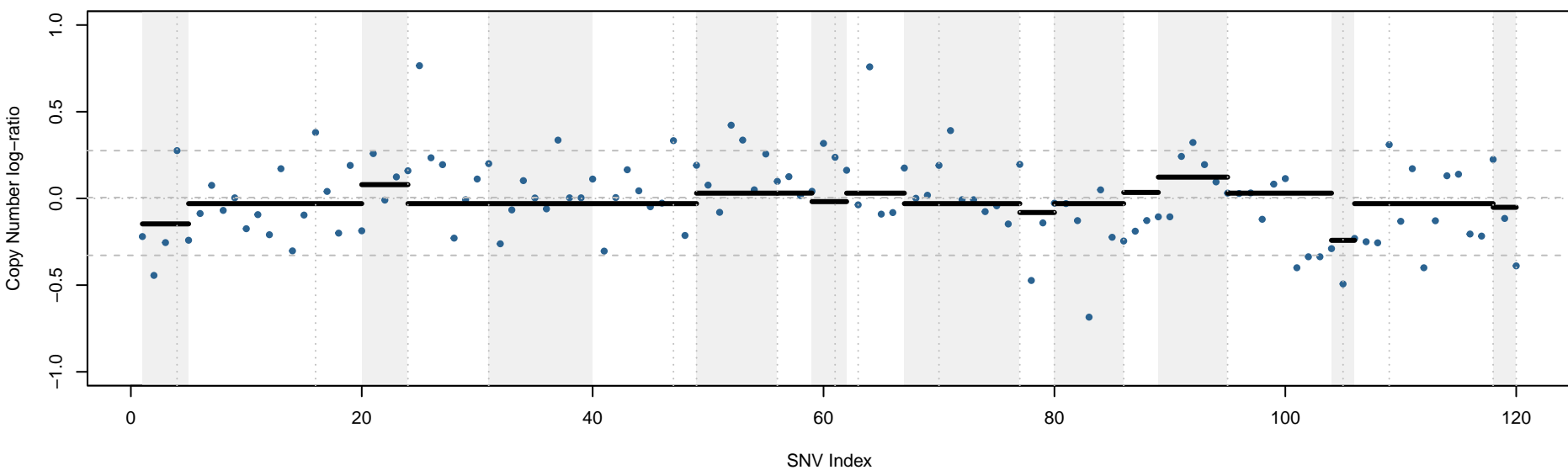
Purity: 0.52 Tumor ploidy: 2.983

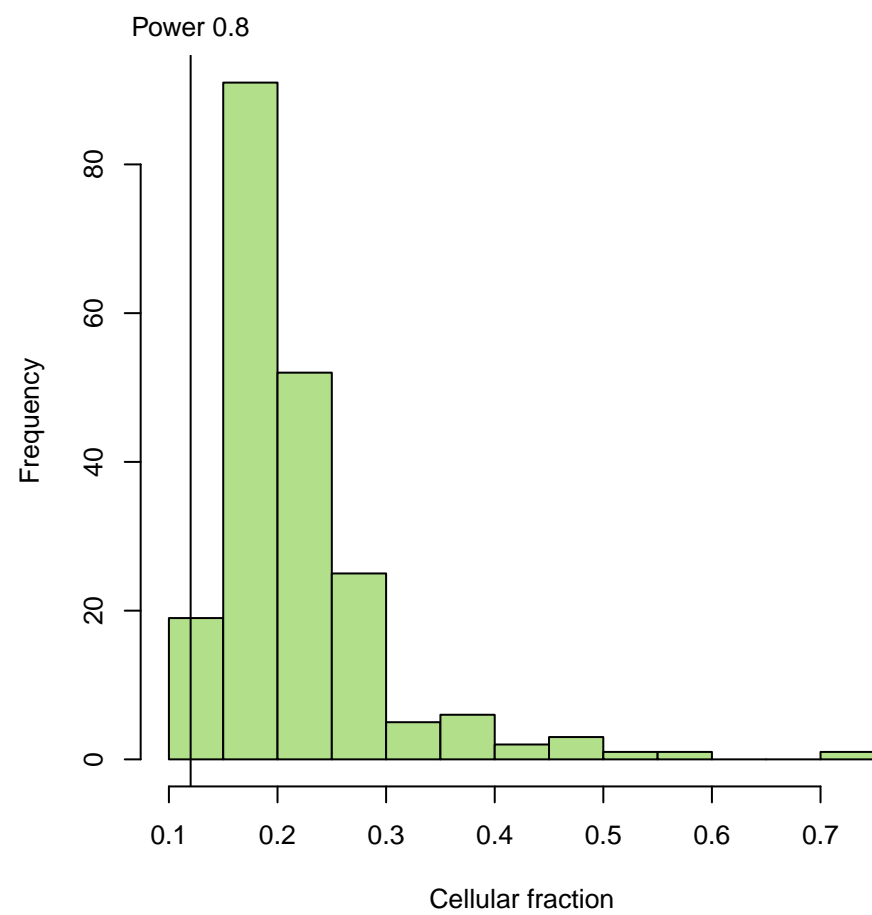
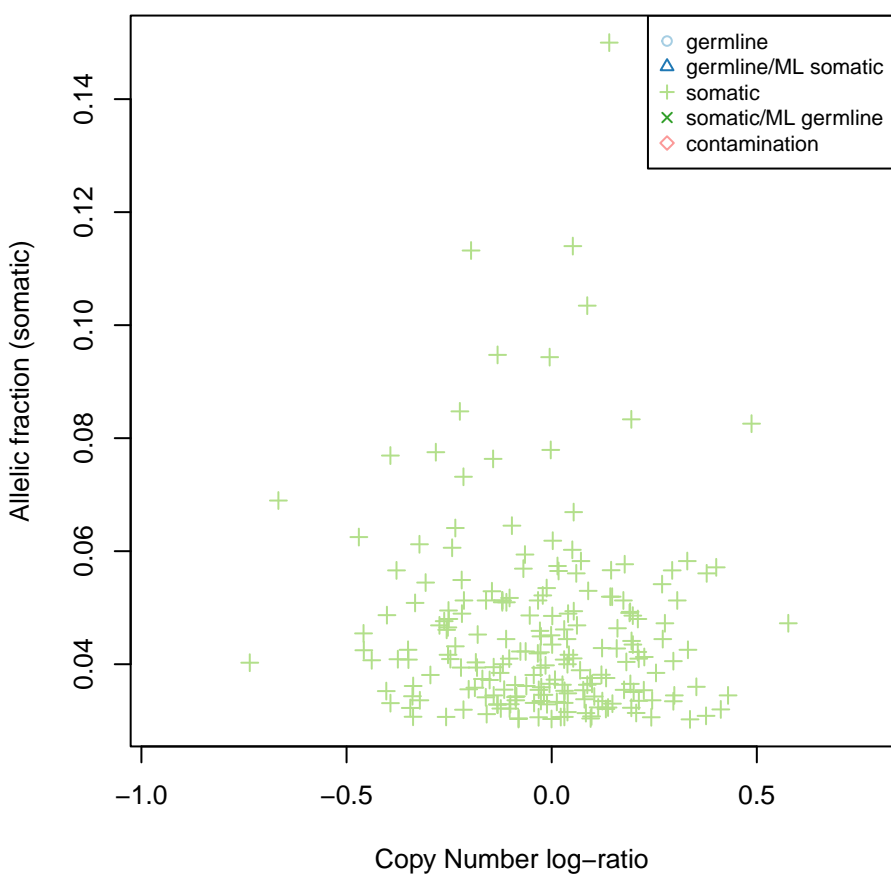
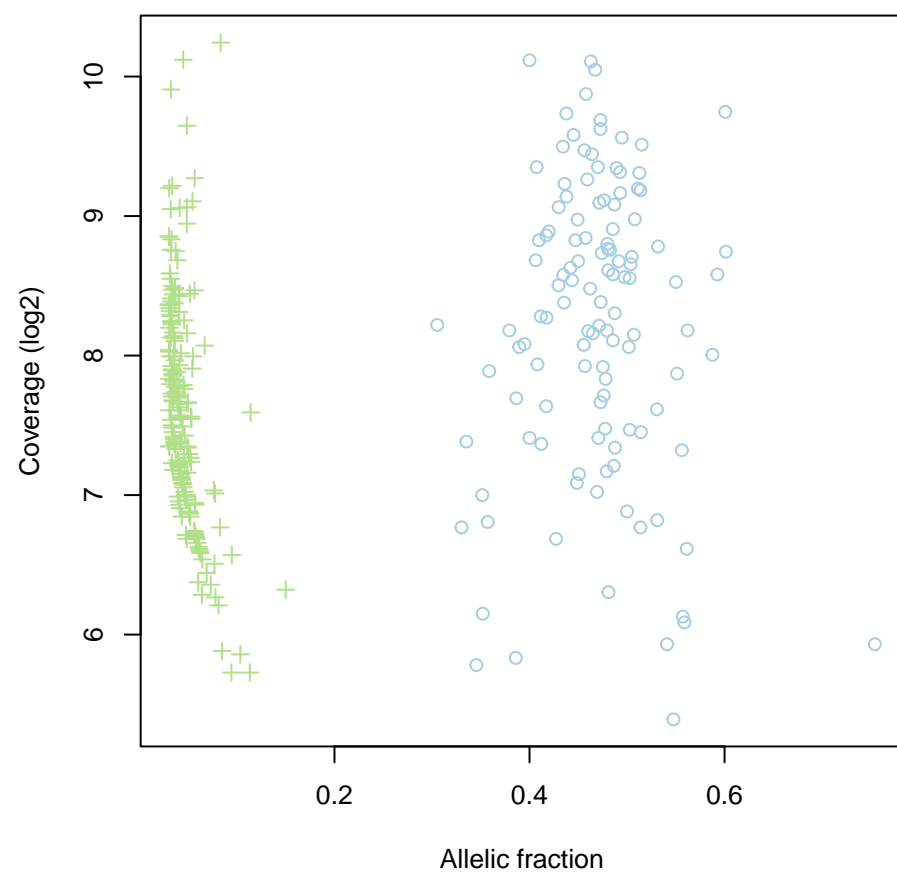
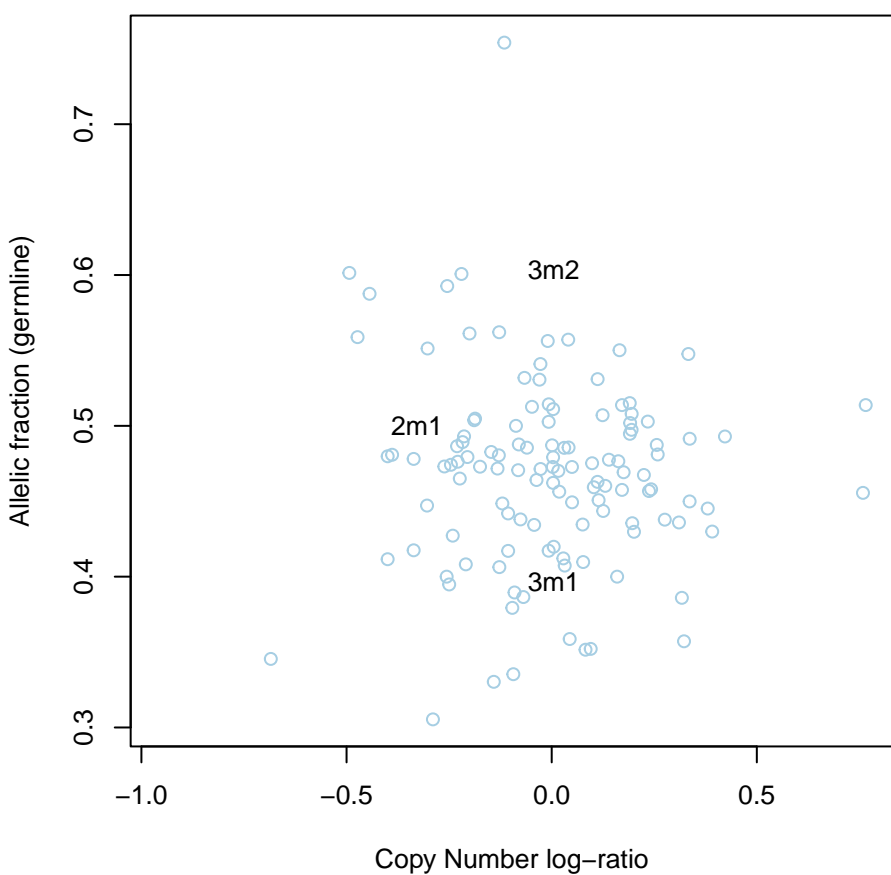


Purity: 0.52 Tumor ploidy: 2.983 SNV log-likelihood: -424.91 GoF: 54.2% Mean coverage: 406;281

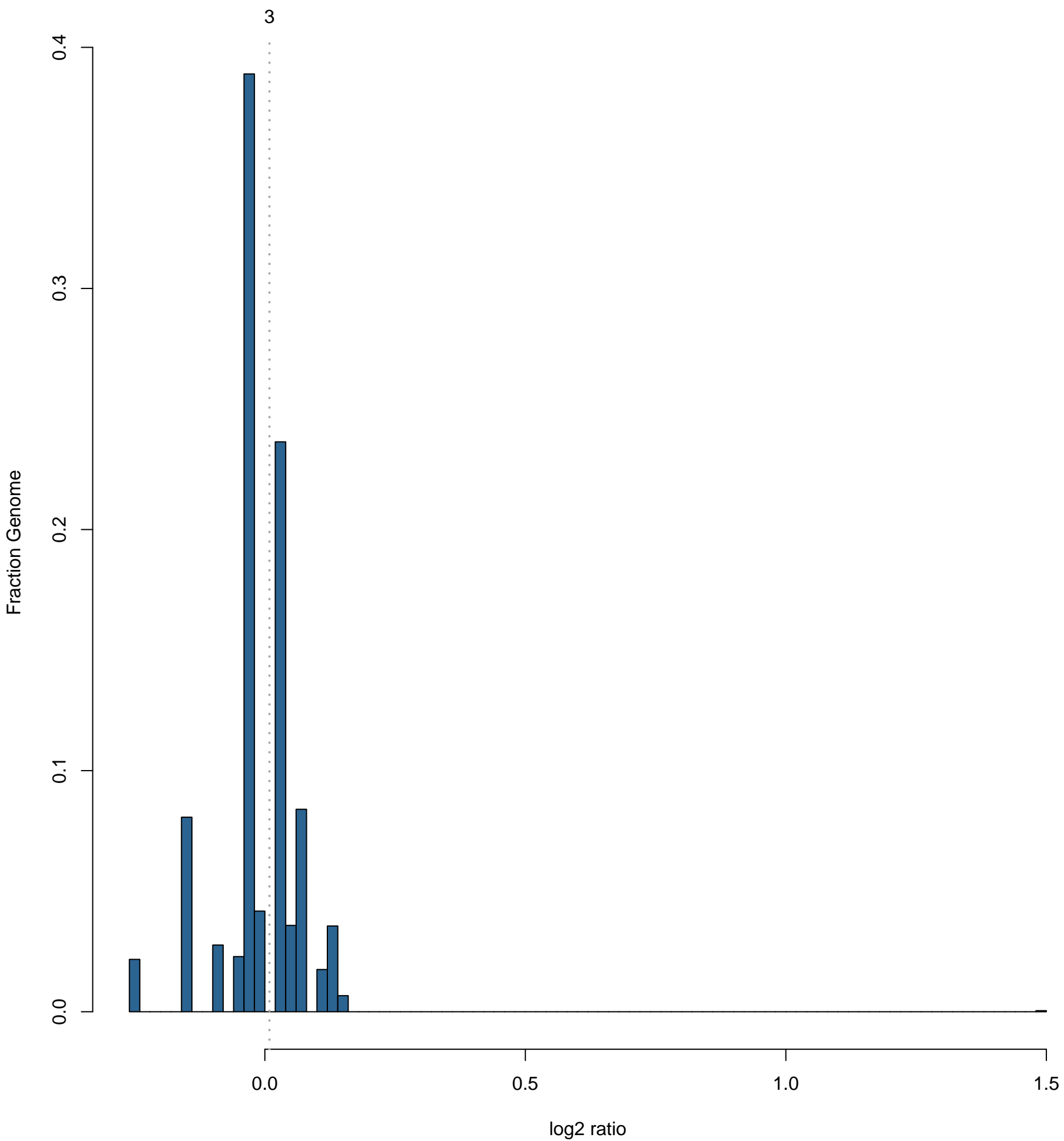


SCNA-fit log-likelihood: -16066.47

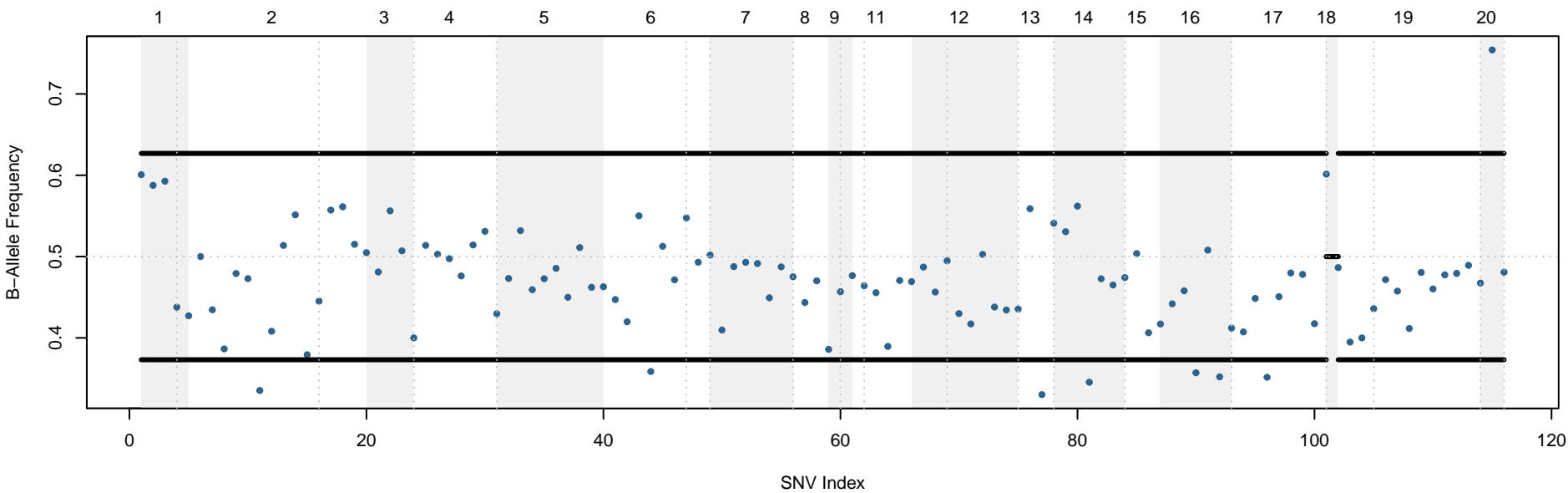




Purity: 0.68 Tumor ploidy: 2.976



Purity: 0.68 Tumor ploidy: 2.976 SNV log-likelihood: -930.94 GoF: 23.9% Mean coverage: 406,281



SCNA-fit log-likelihood: -16088.24

