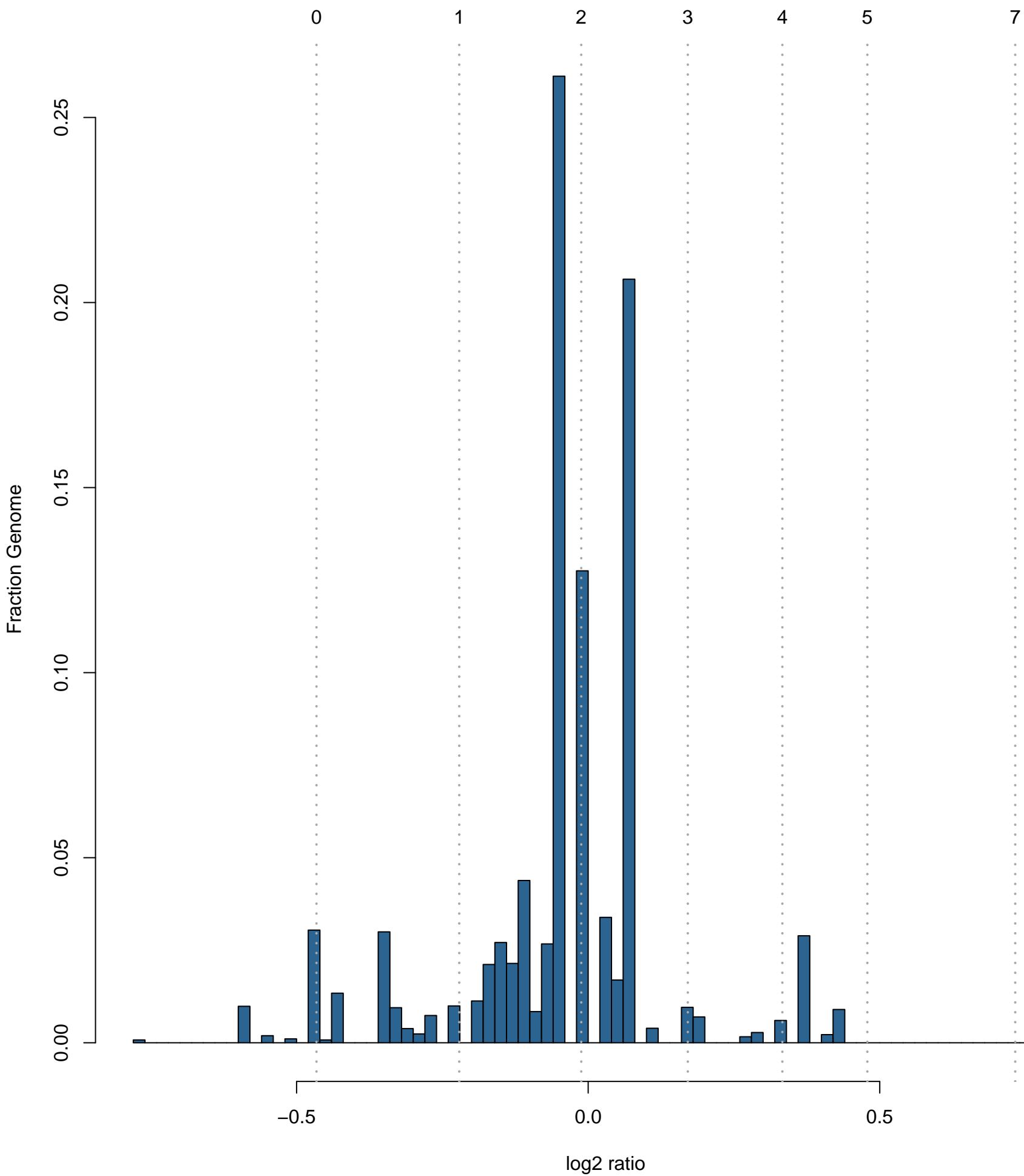
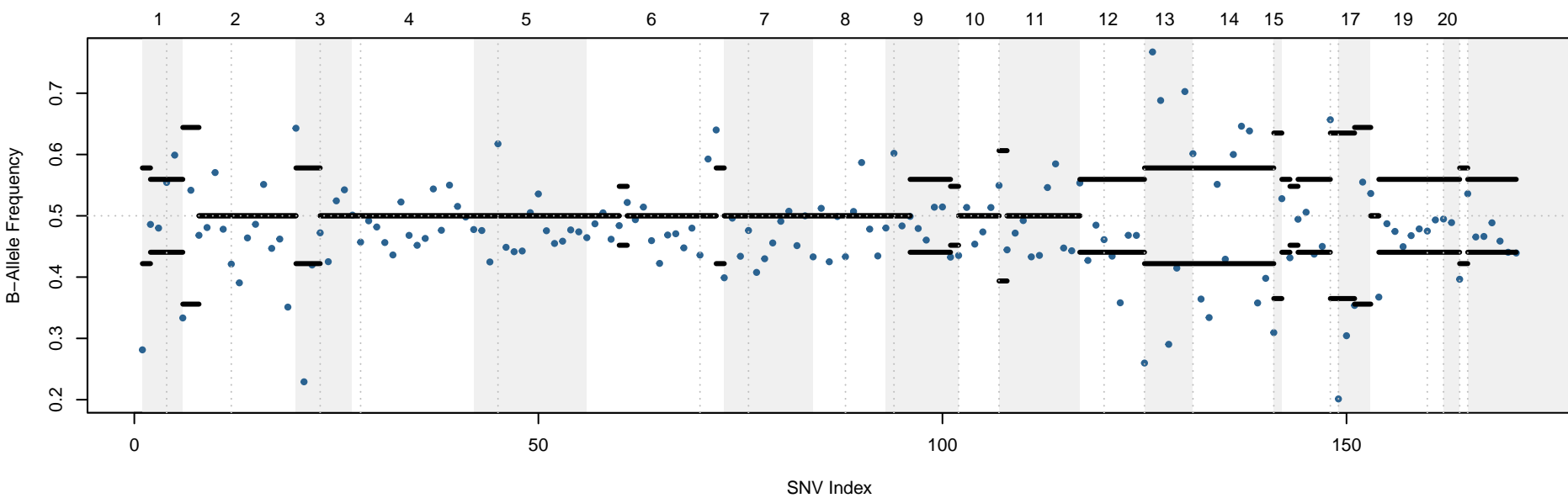


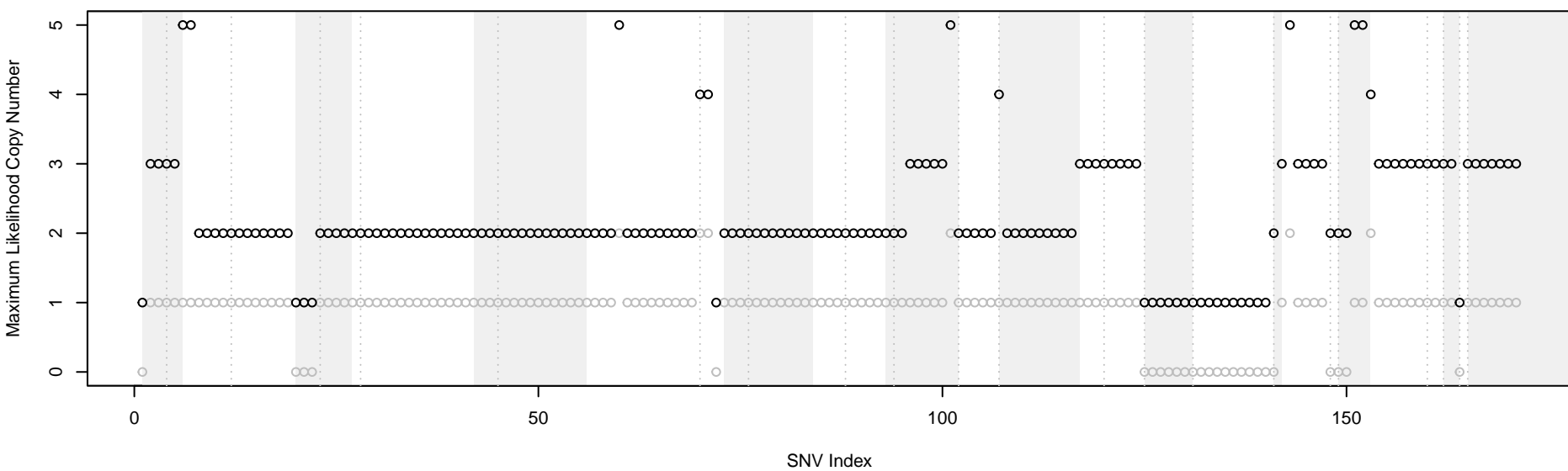
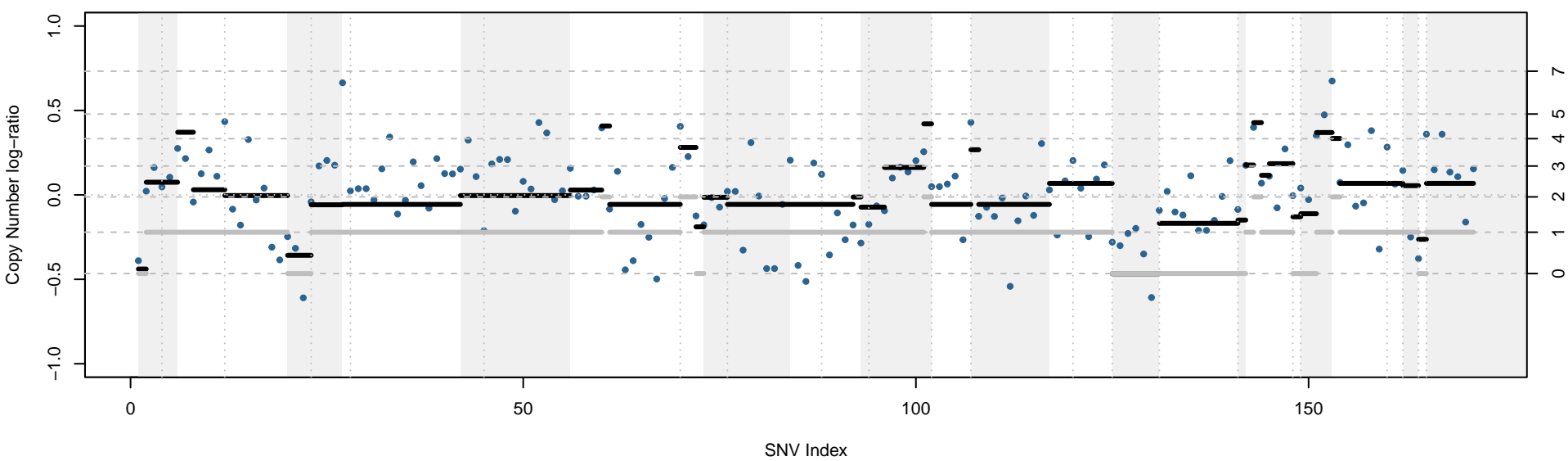
Purity: 0.27 Tumor ploidy: 2.061



Purity: 0.27 Tumor ploidy: 2.061 SNV log-likelihood: -72.2 GoF: 89.8% Mean coverage: 514,449

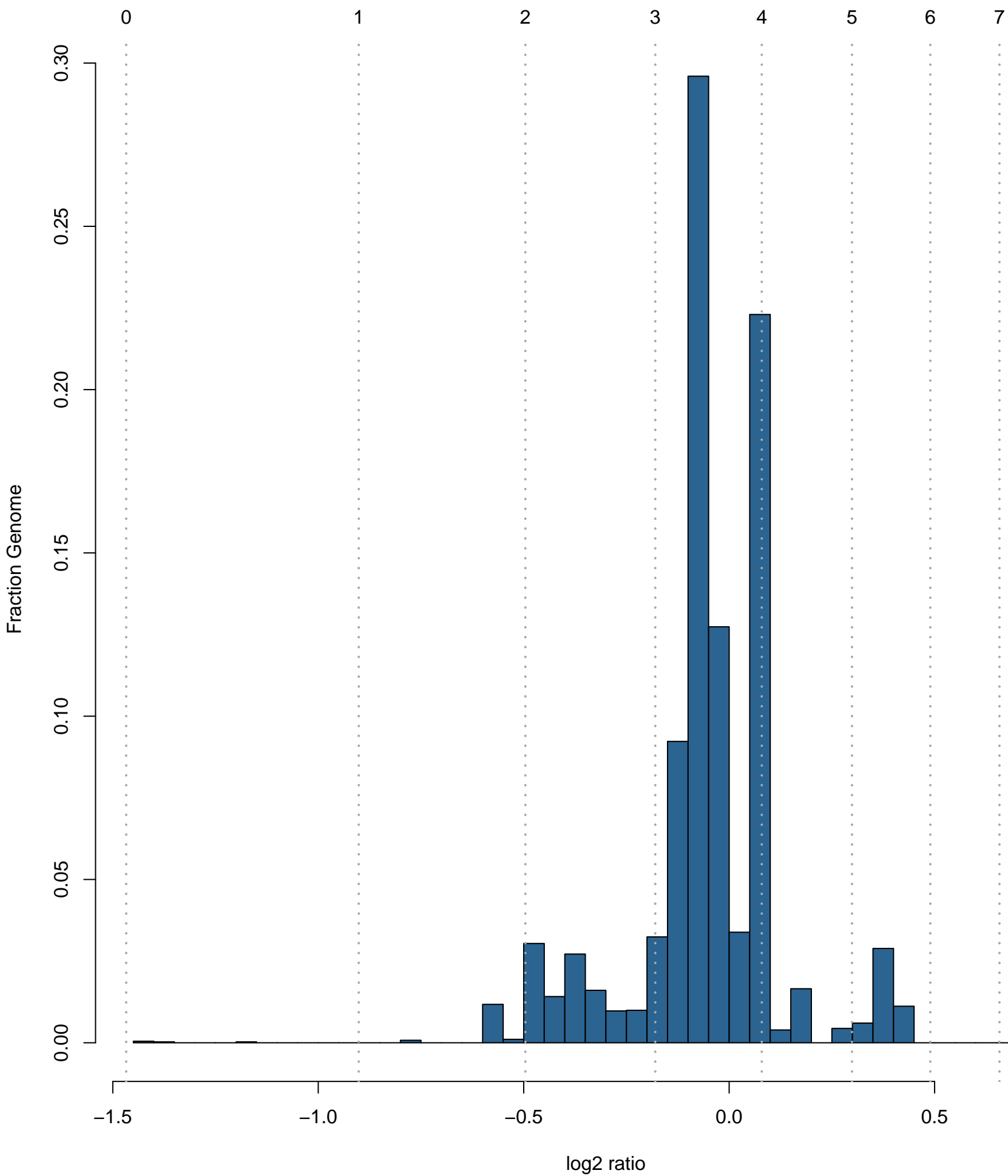


SCNA-fit log-likelihood: -4597.46

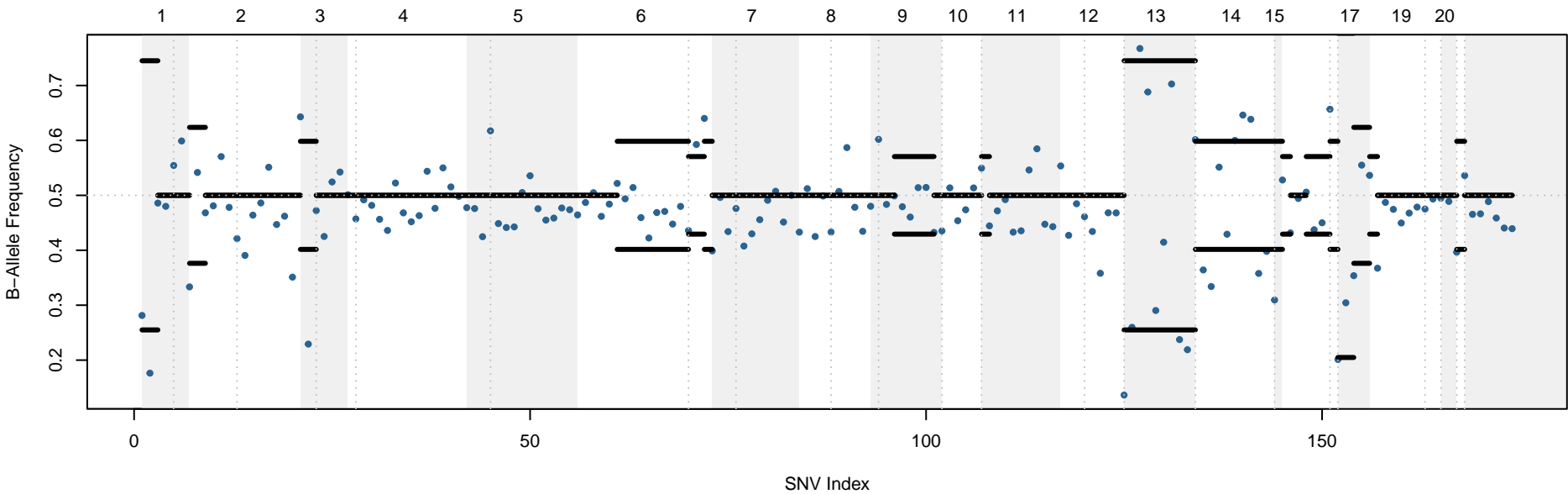




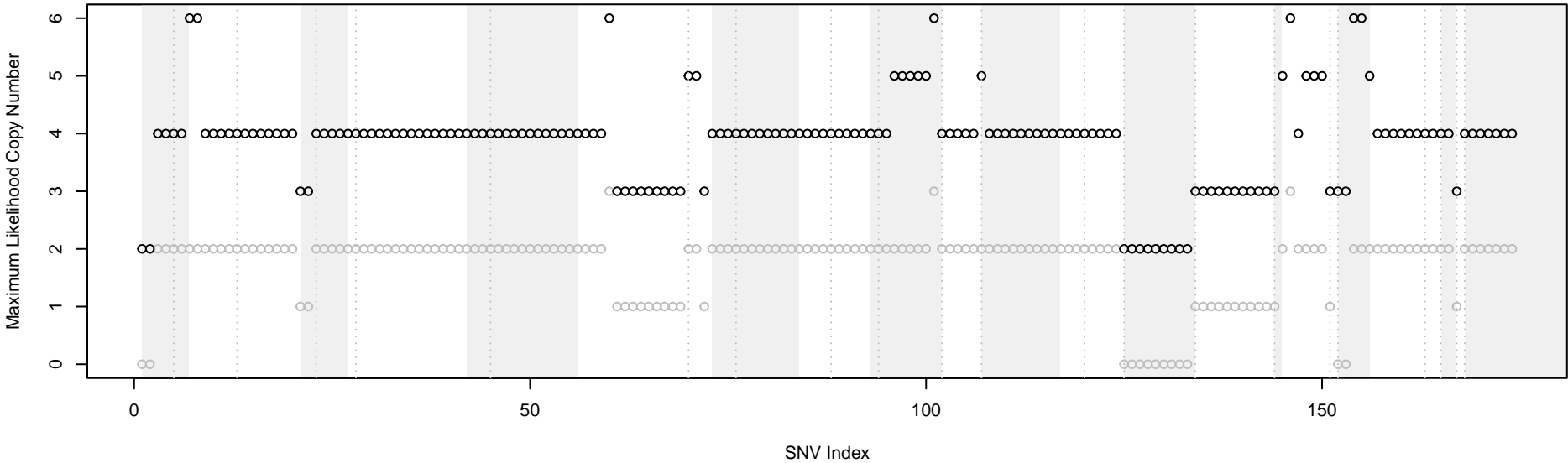
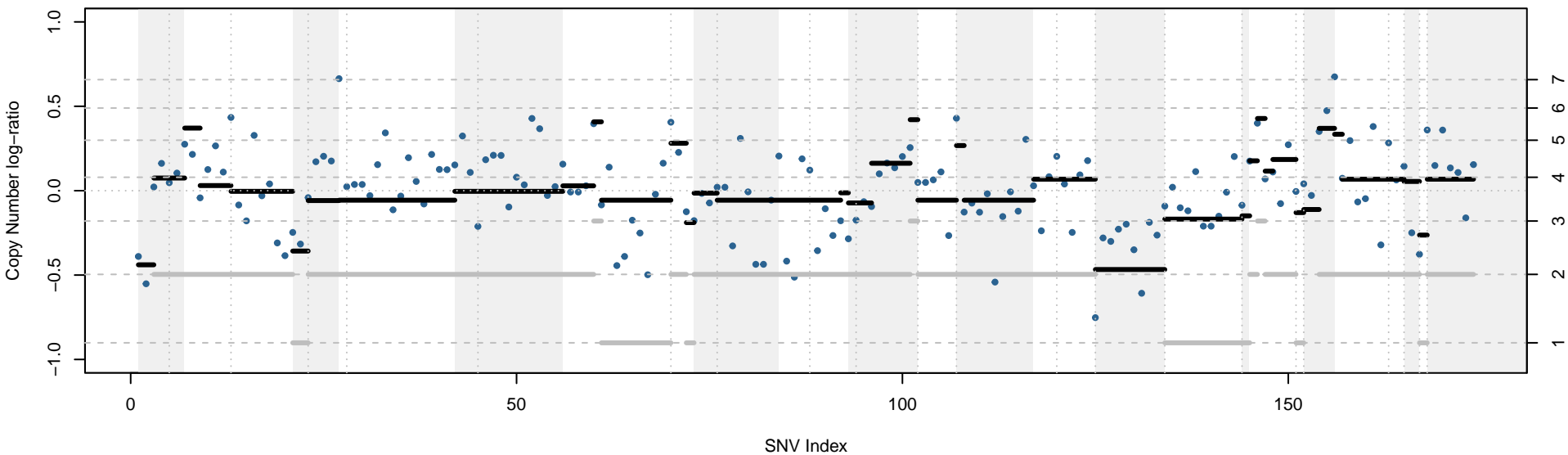
Purity: 0.49 Tumor ploidy: 3.675

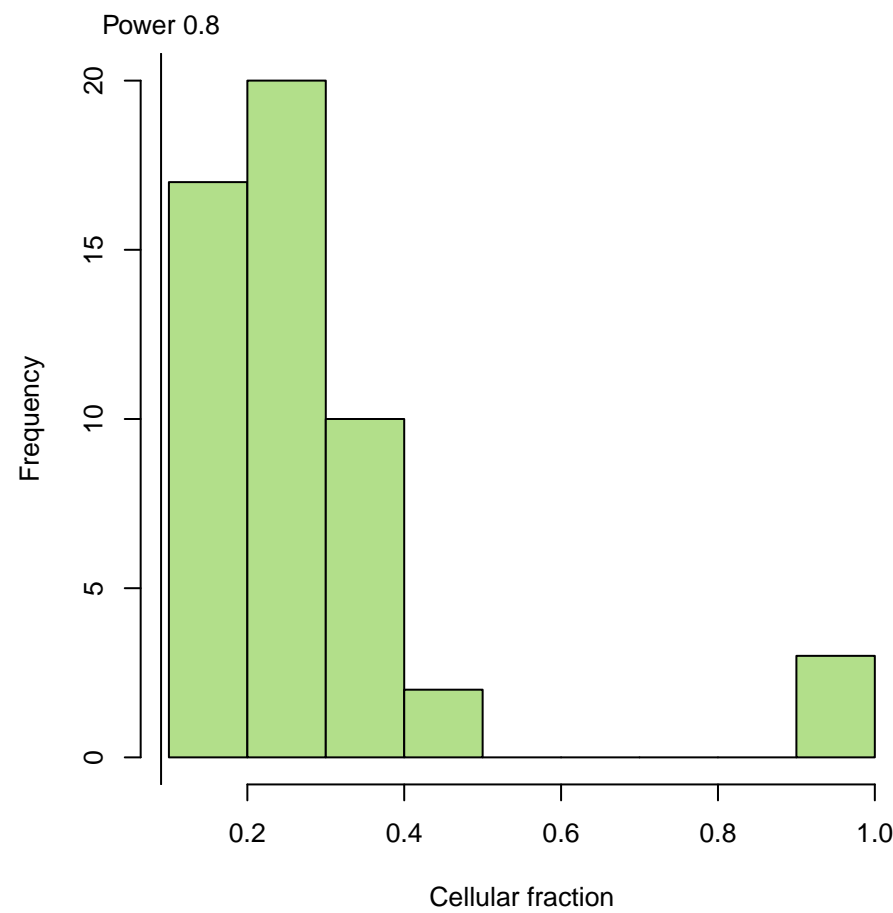
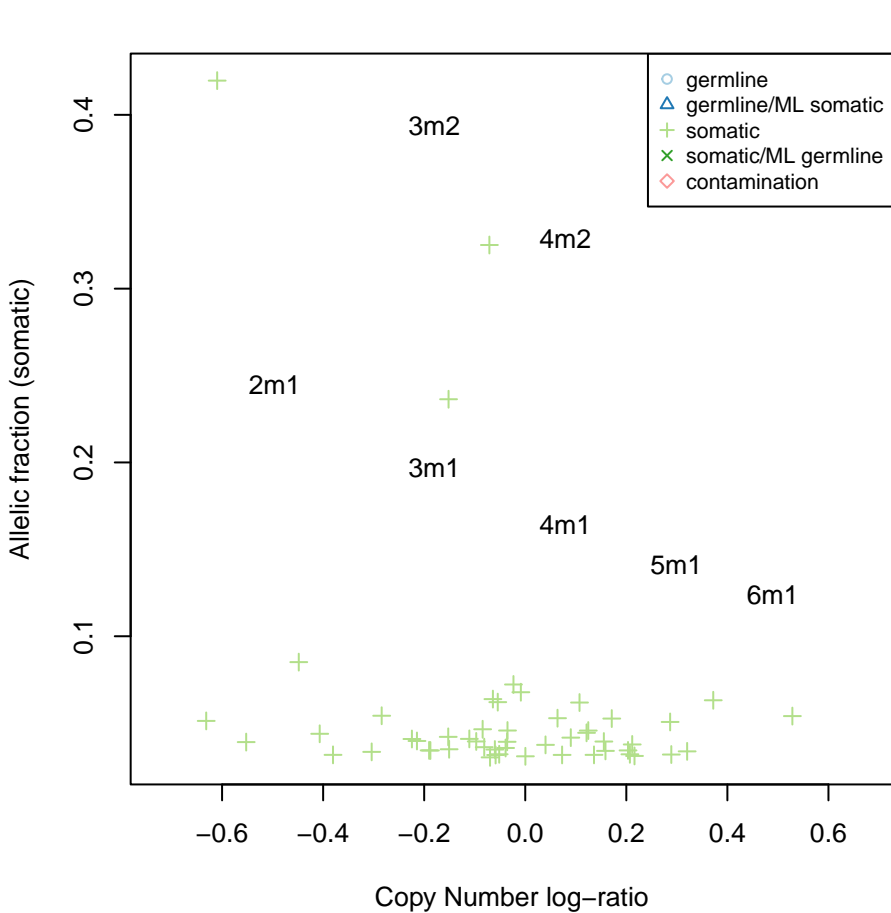
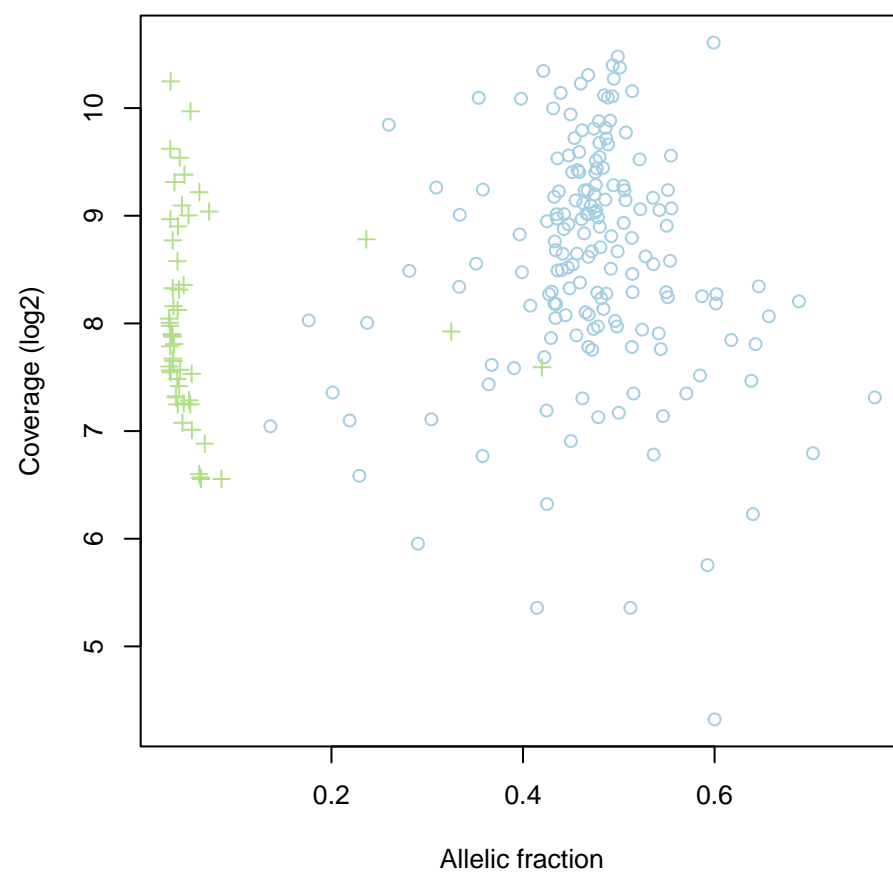
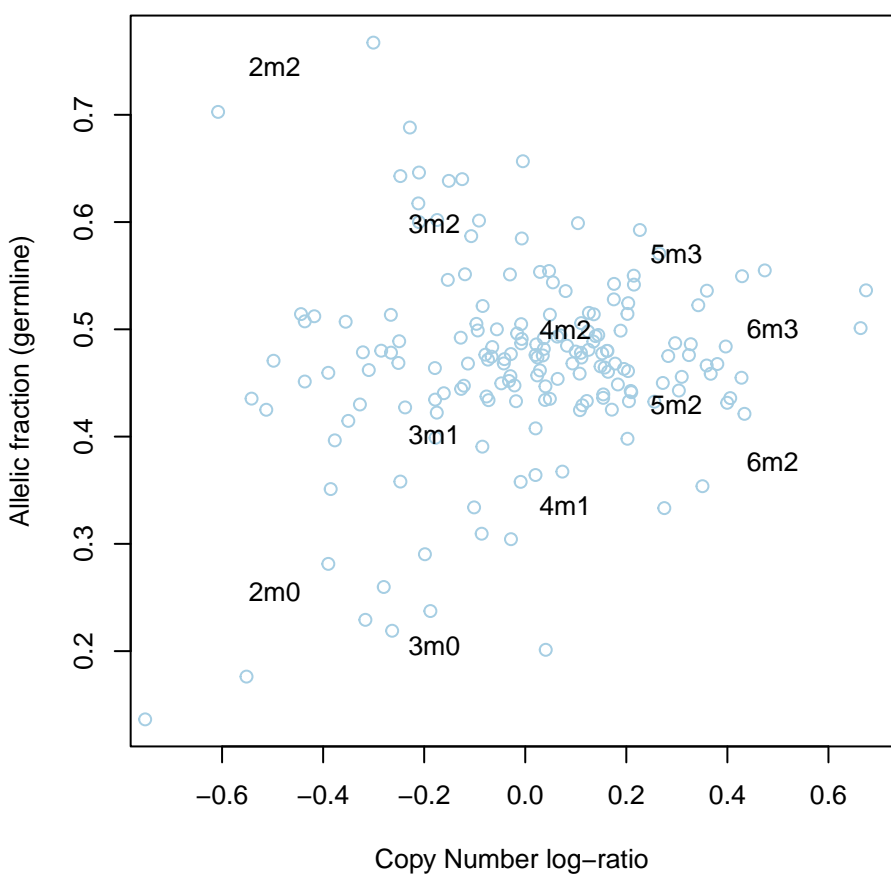


Purity: 0.49 Tumor ploidy: 3.675 SNV log-likelihood: -119 GoF: 84.6% Mean coverage: 514;449

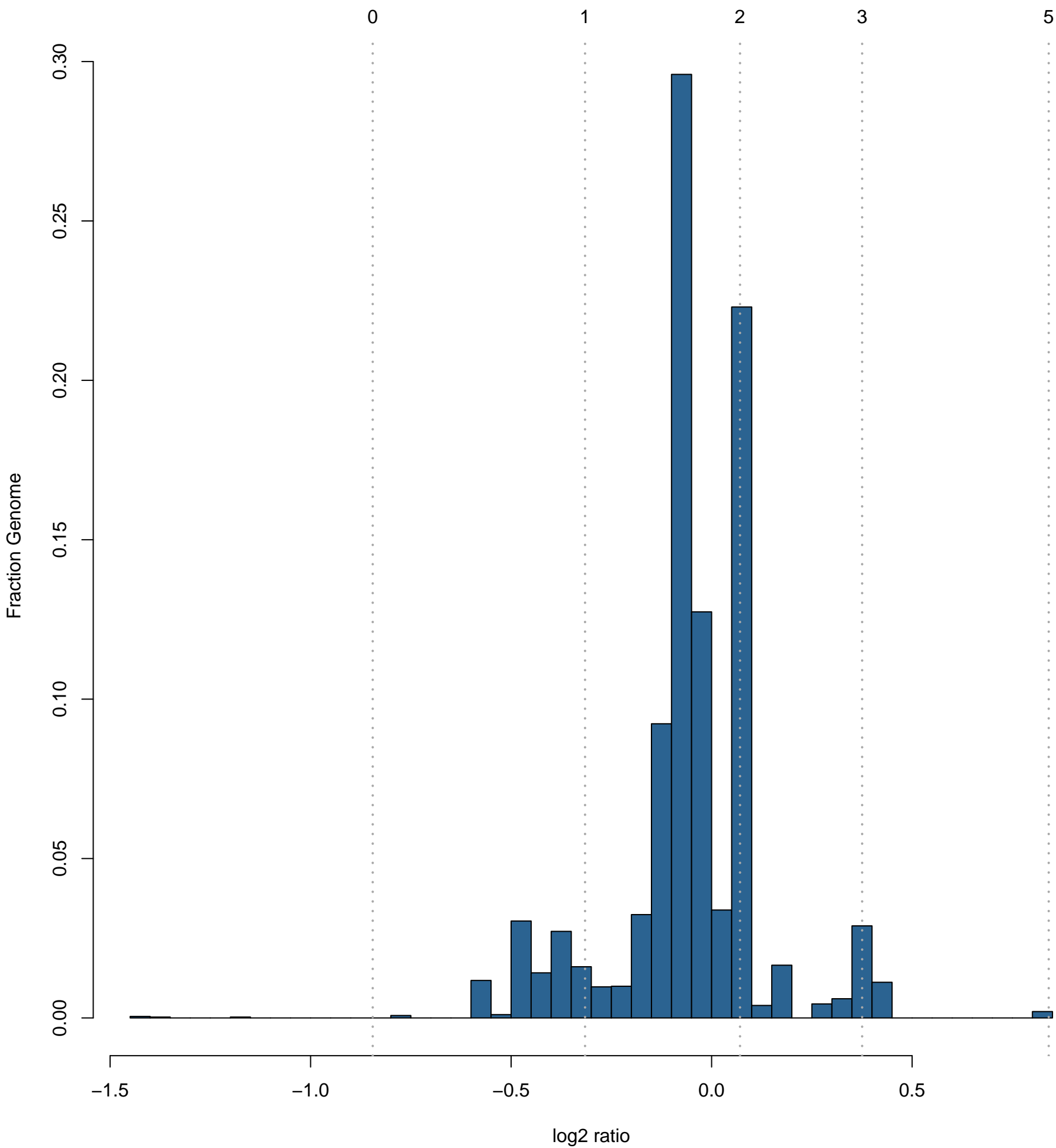


SCNA-fit log-likelihood: -4601.29

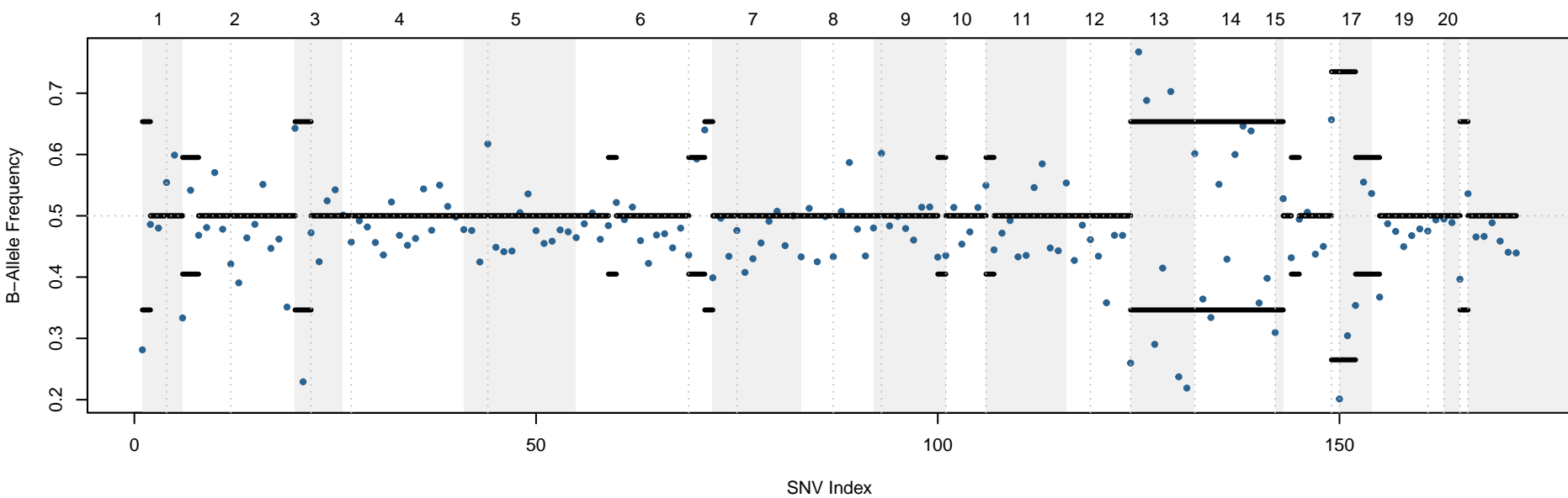




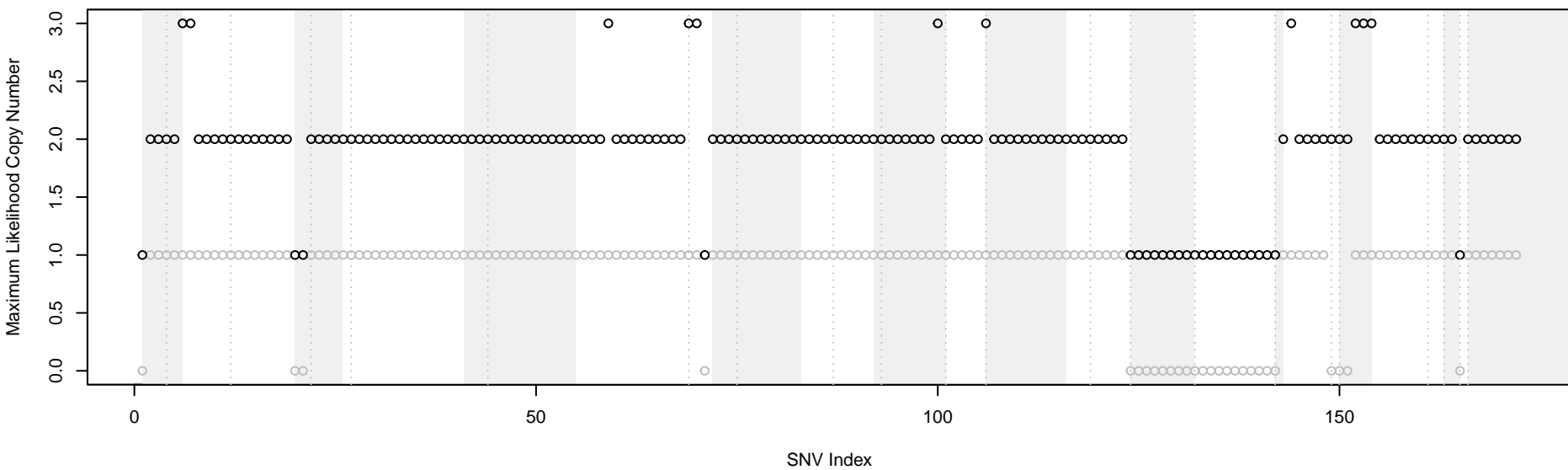
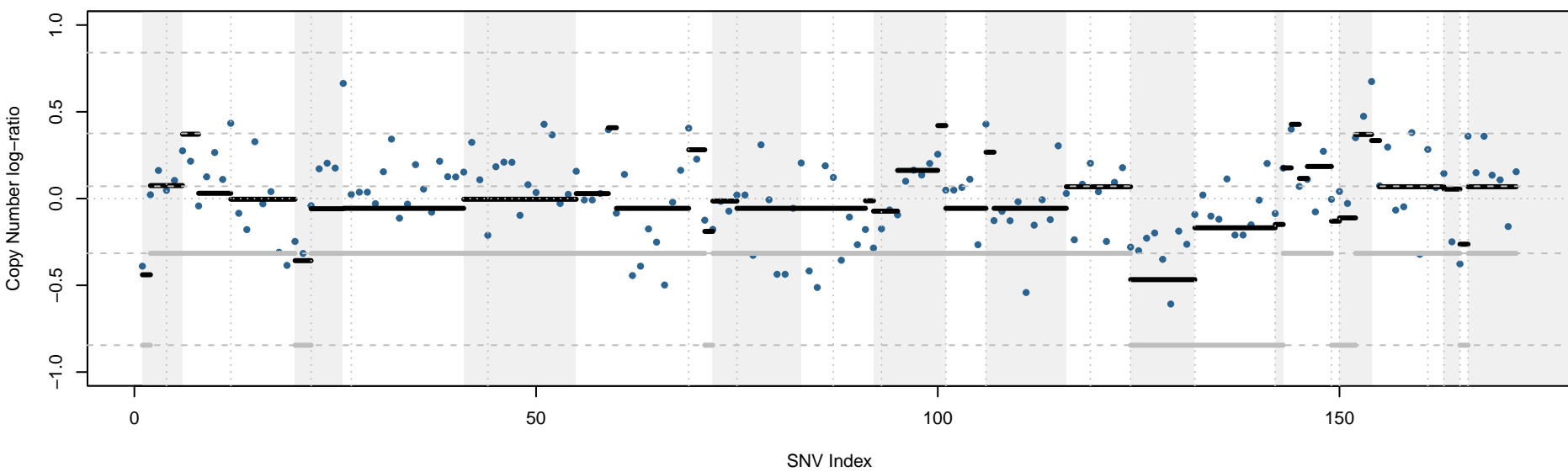
Purity: 0.47 Tumor ploidy: 1.796



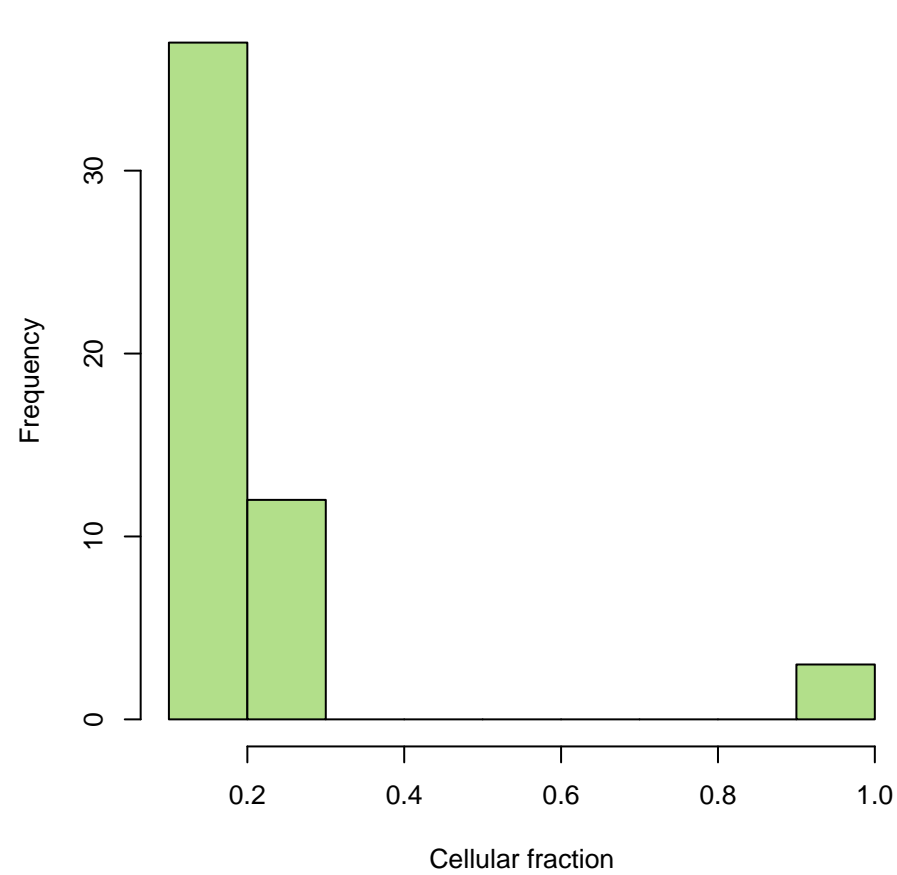
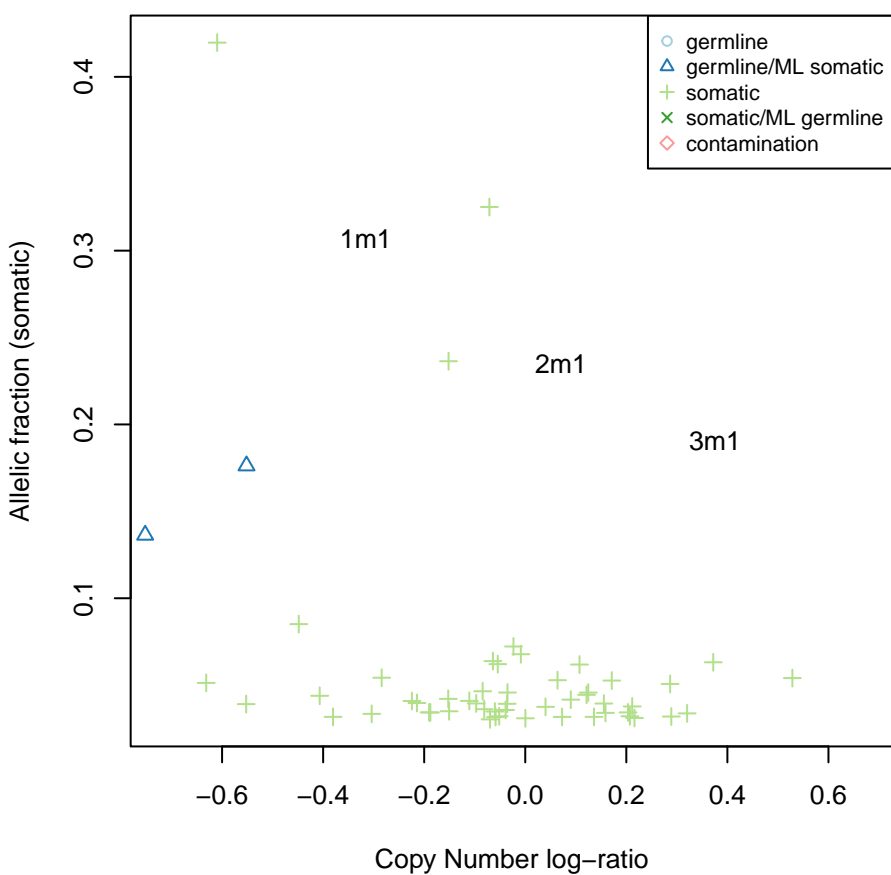
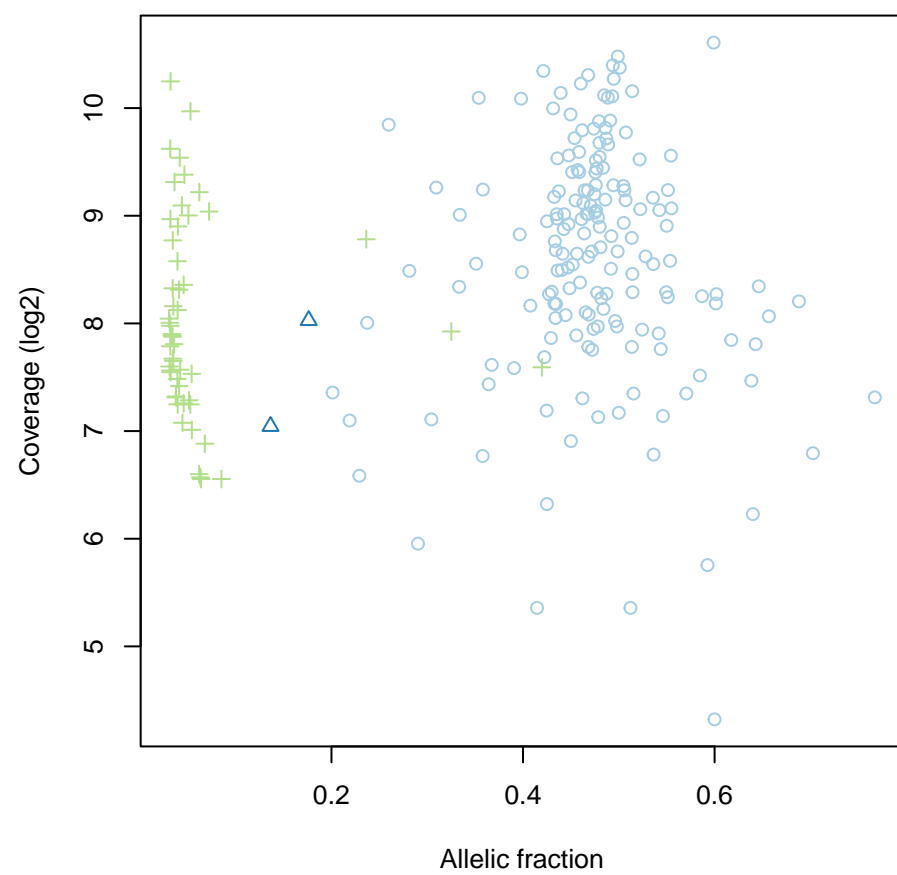
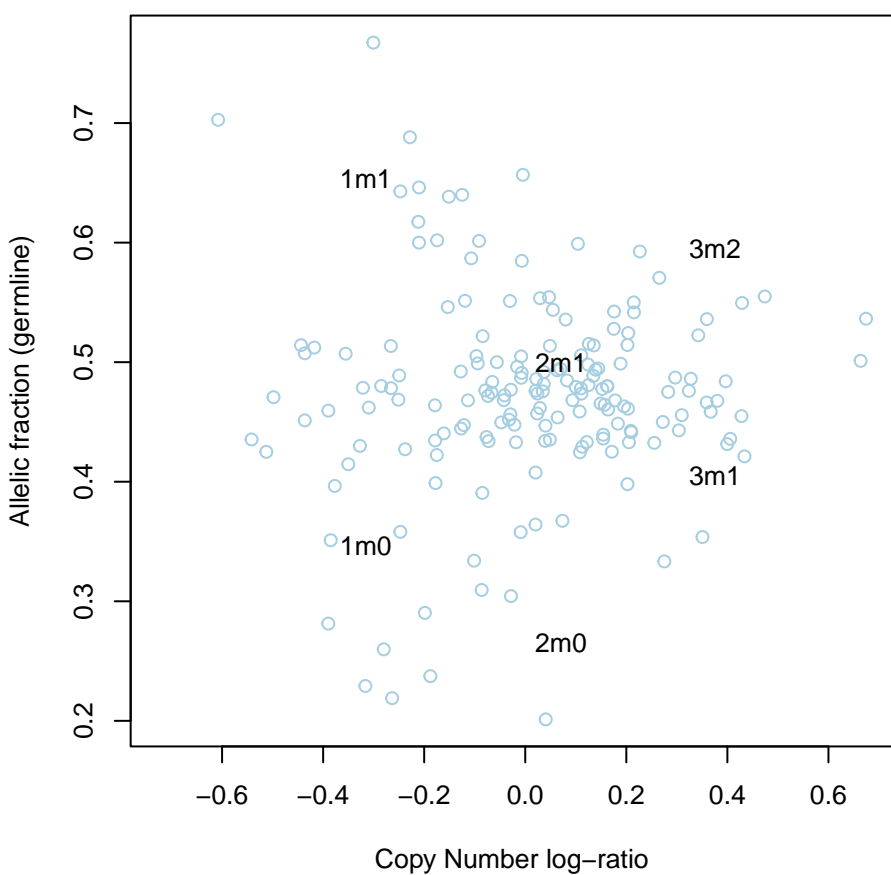
Purity: 0.47 Tumor ploidy: 1.796 SNV log-likelihood: -104.94 GoF: 71.2% Mean coverage: 514,449



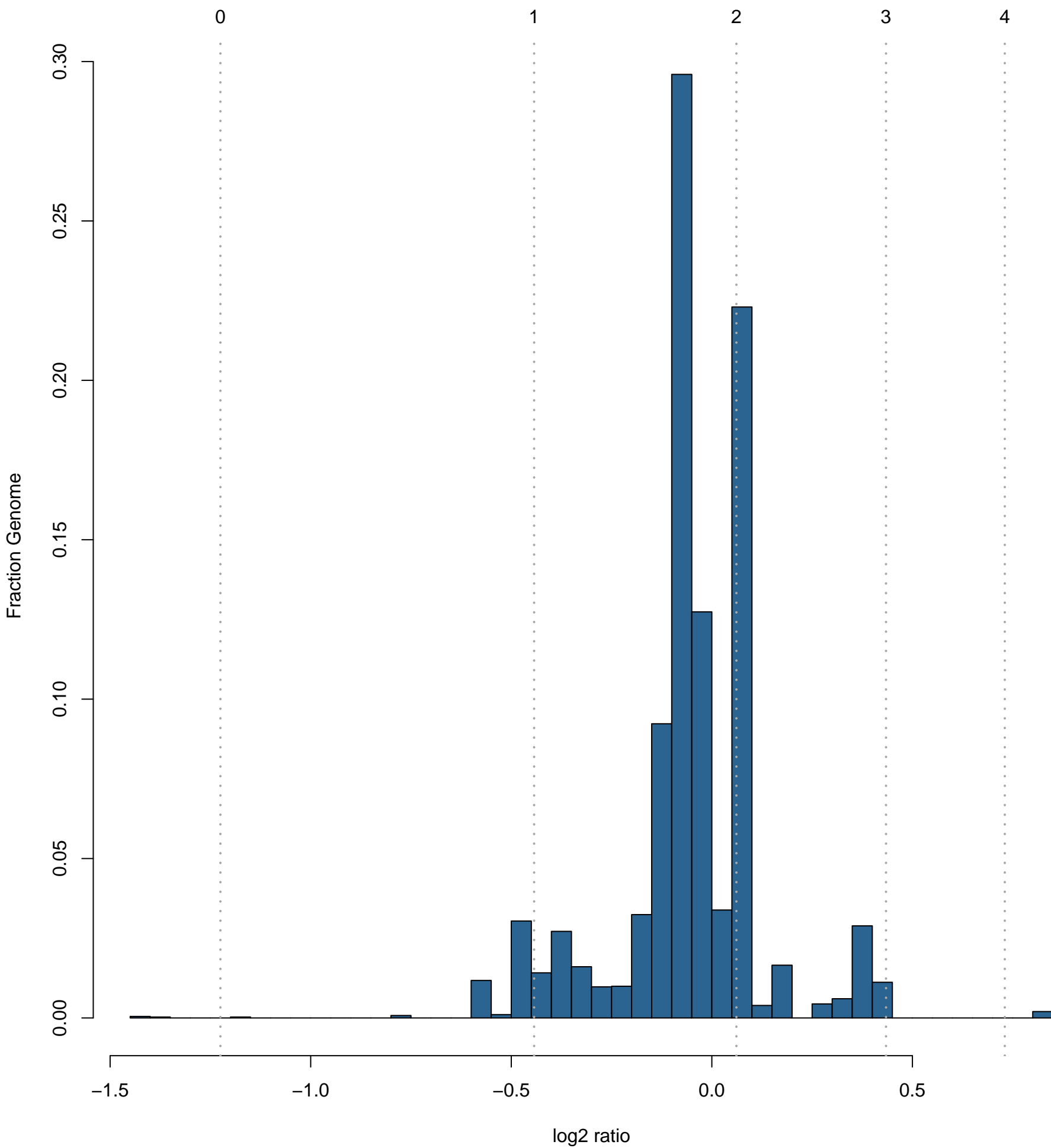
SCNA-fit log-likelihood: -4770.73



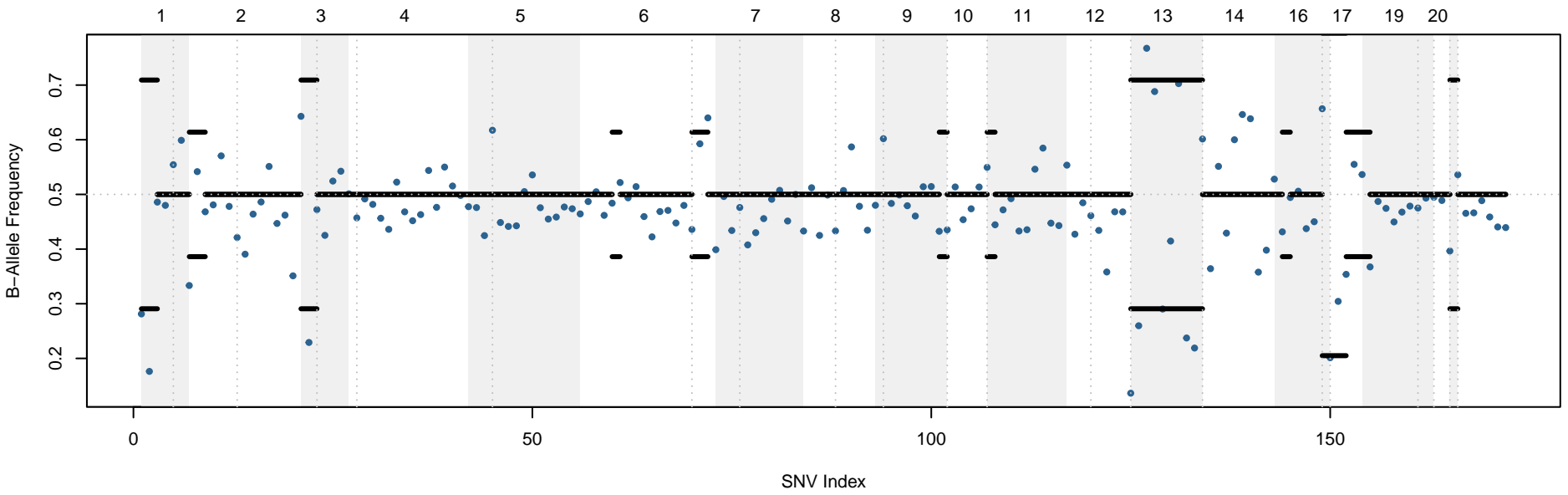




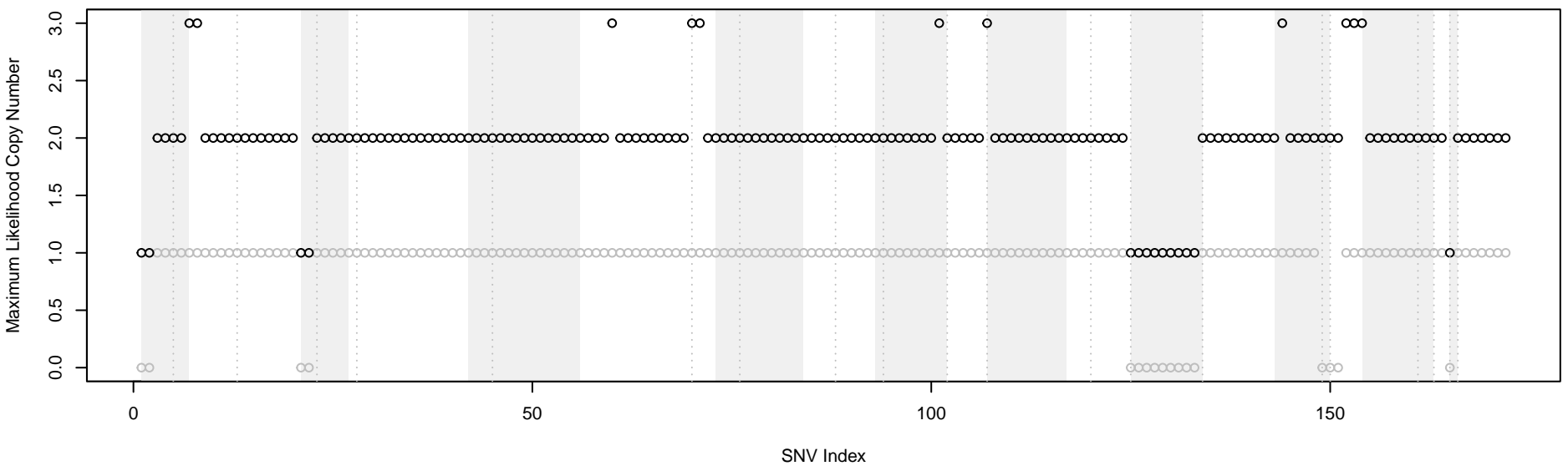
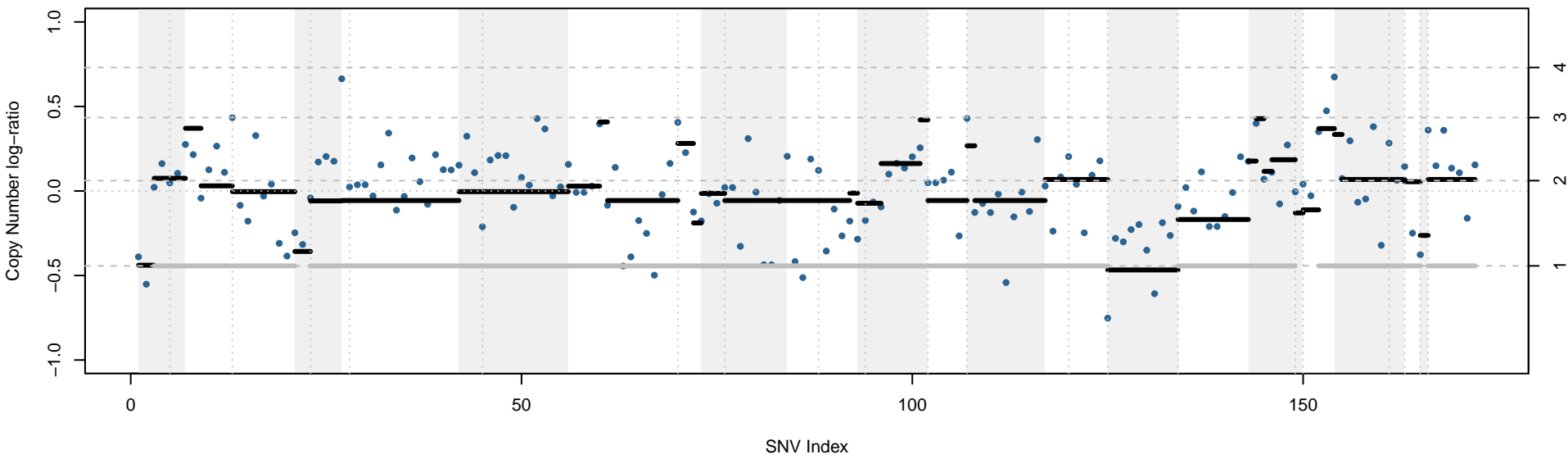
Purity: 0.59 Tumor ploidy: 1.859

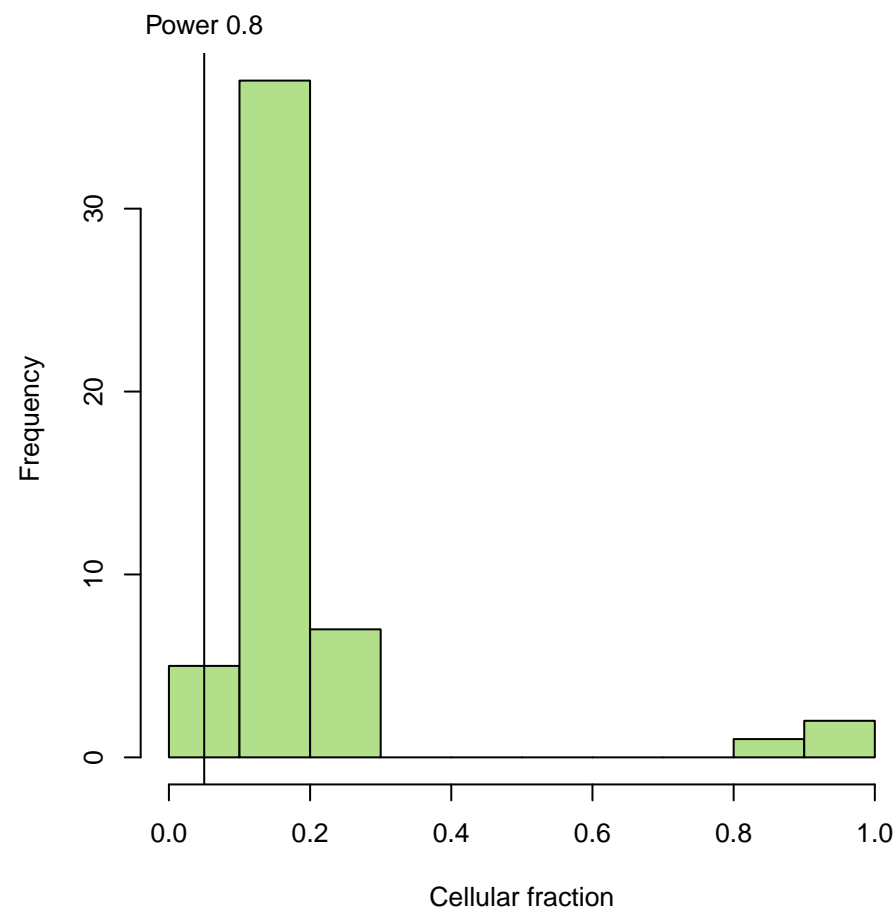
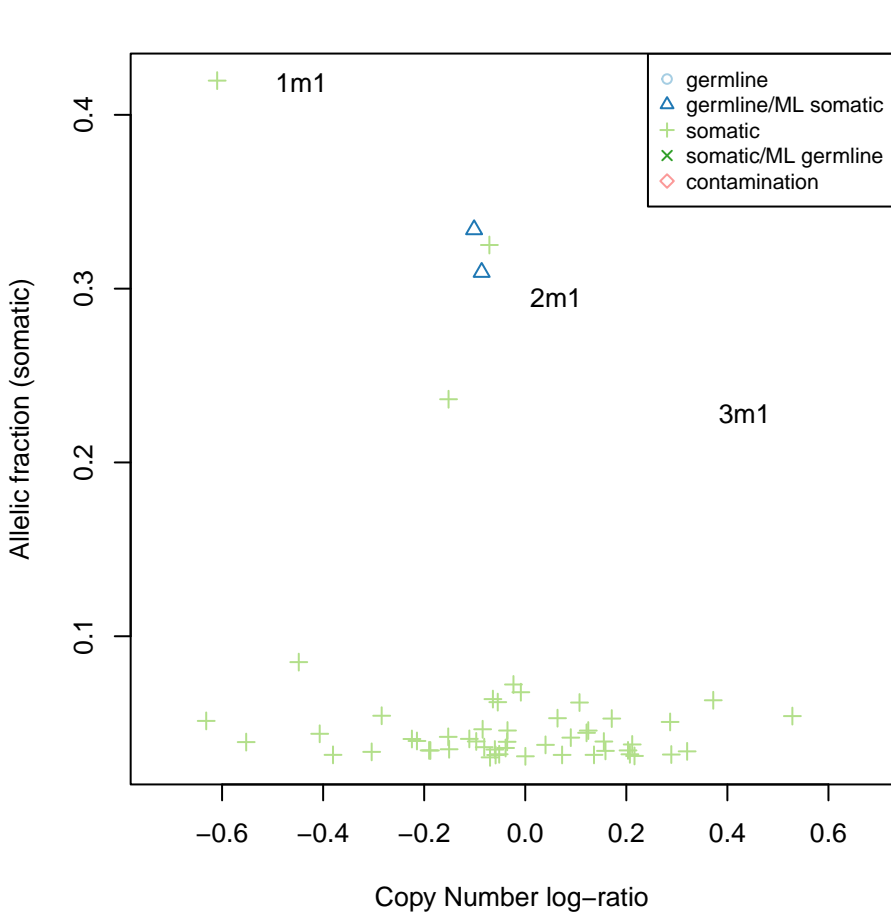
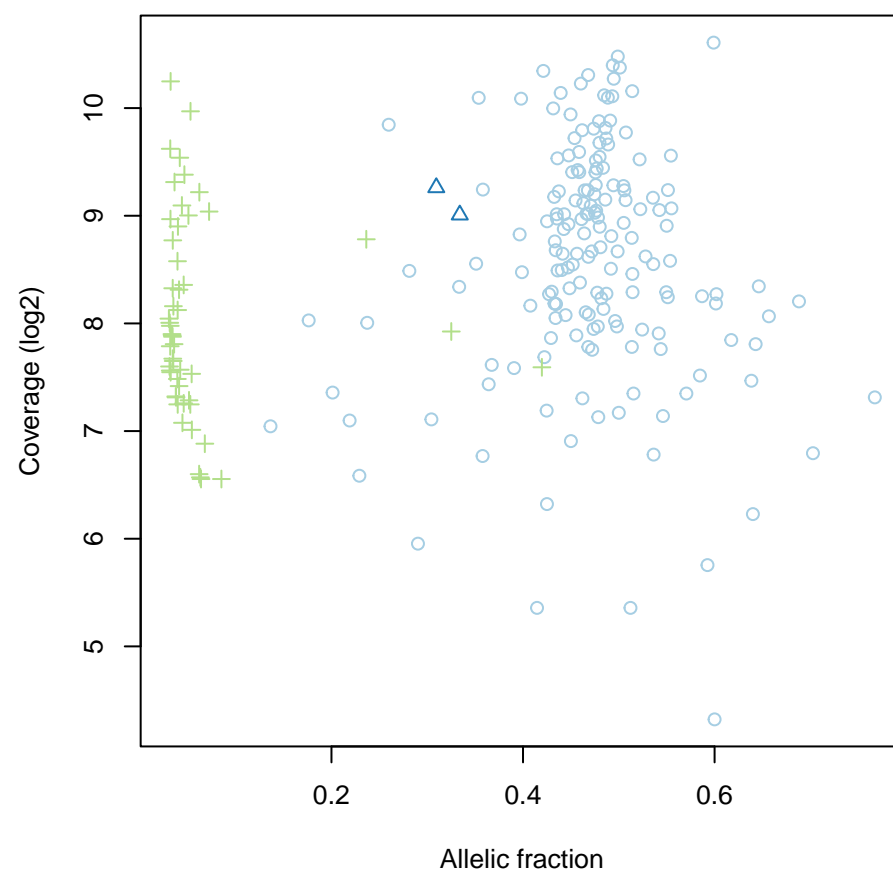
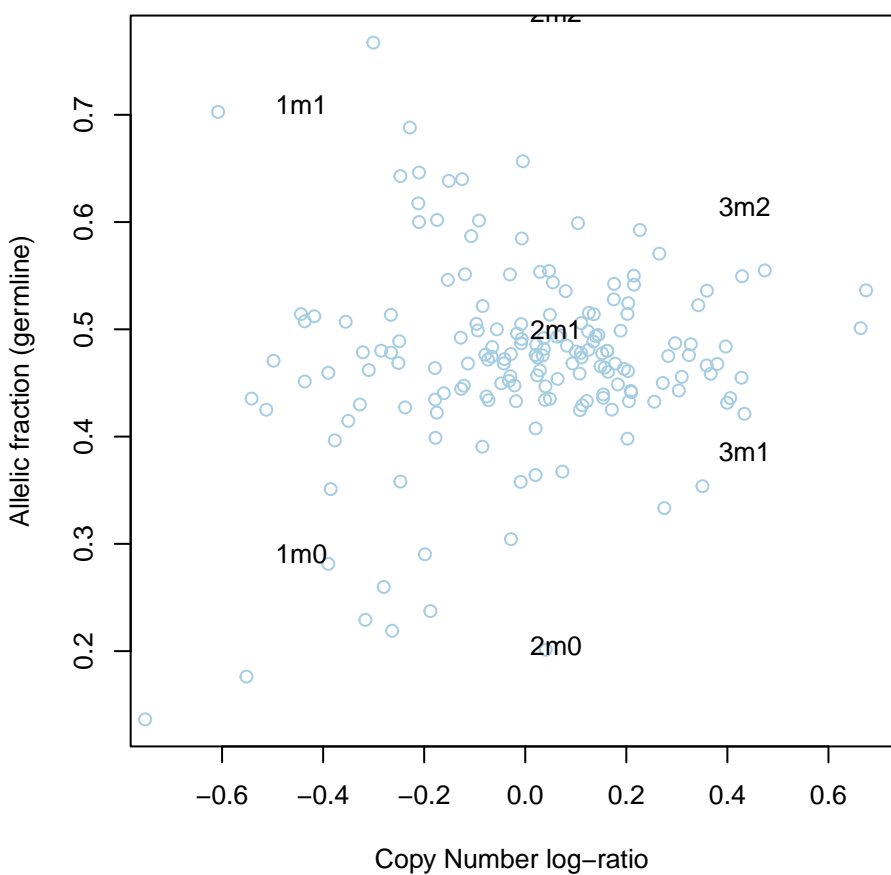


Purity: 0.59 Tumor ploidy: 1.859 SNV log-likelihood: -280.54 GoF: 54.3% Mean coverage: 514,449

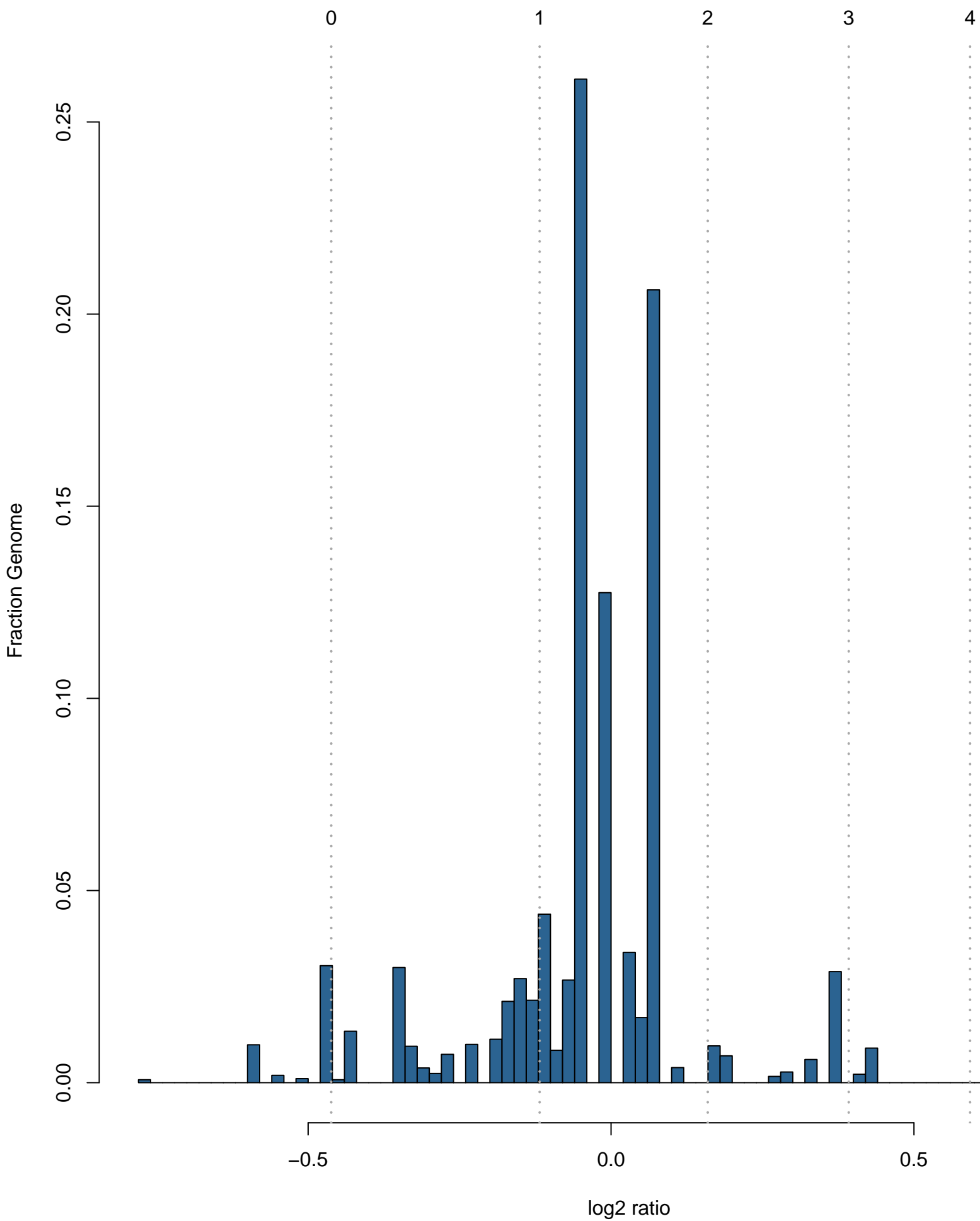


SCNA-fit log-likelihood: -4760.91

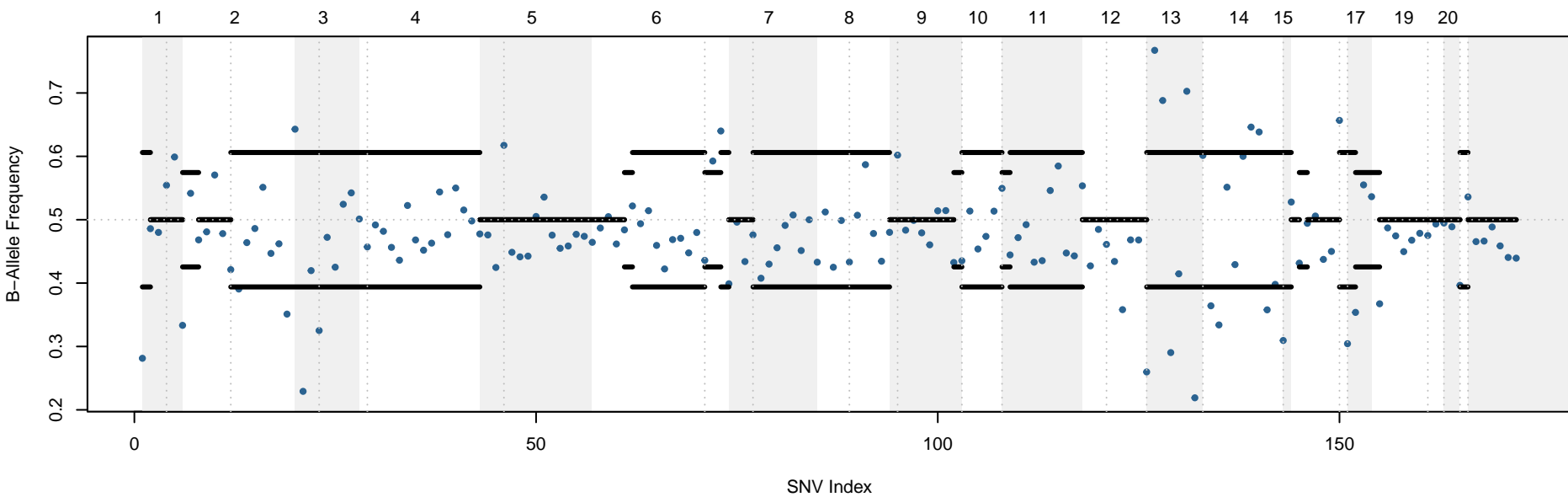




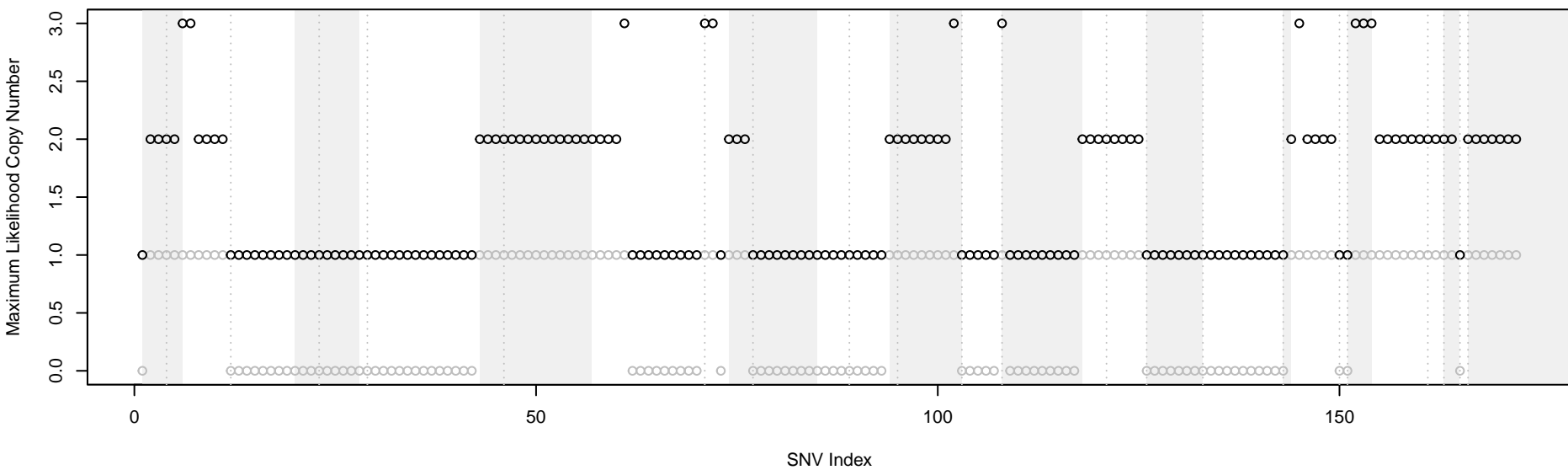
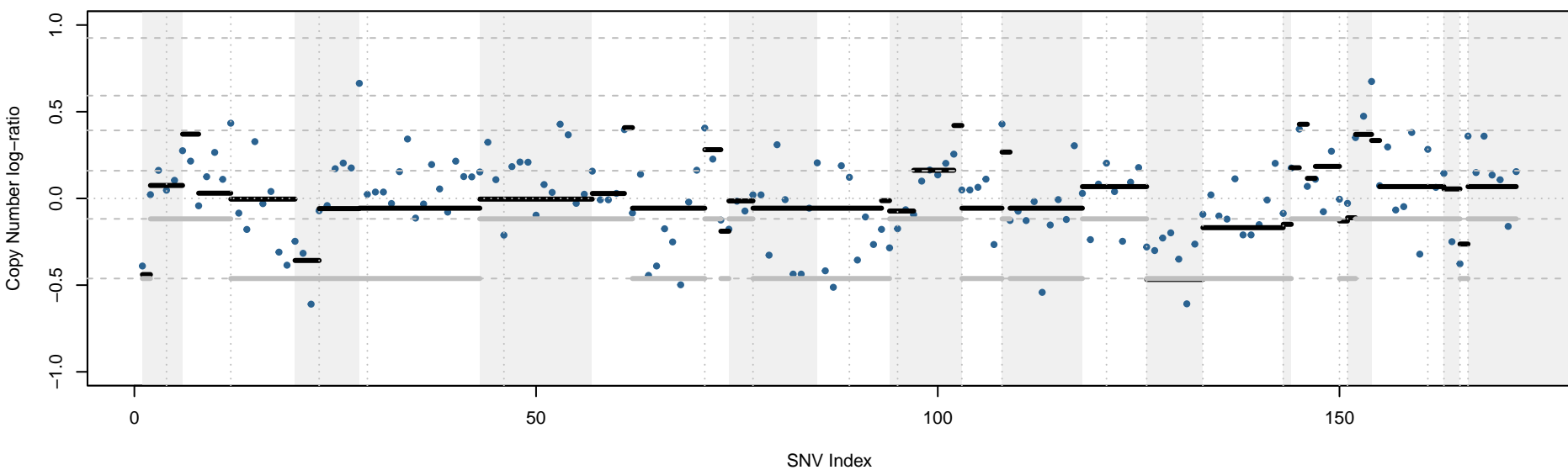
Purity: 0.35 Tumor ploidy: 1.401

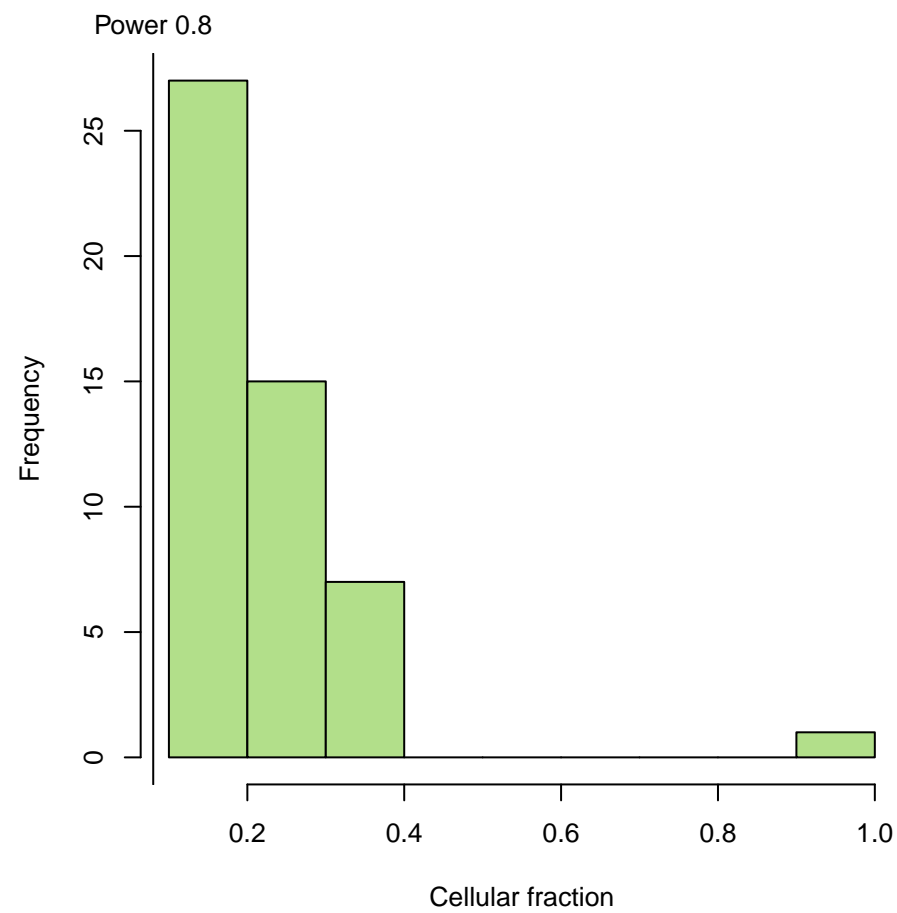
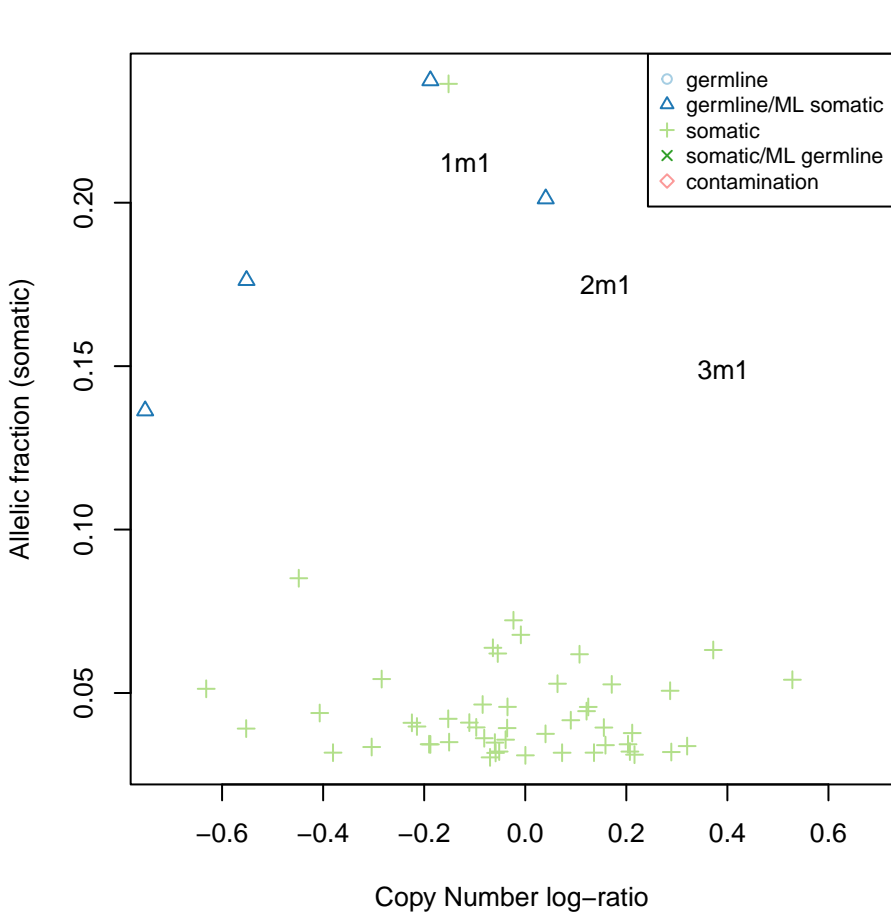
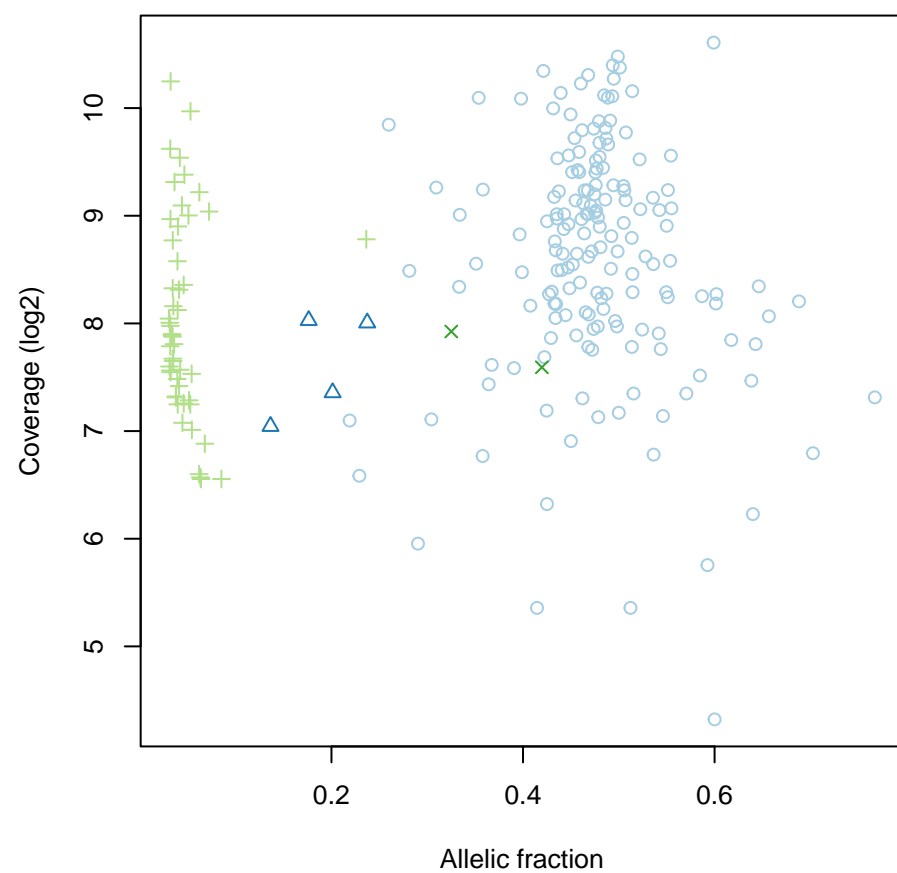
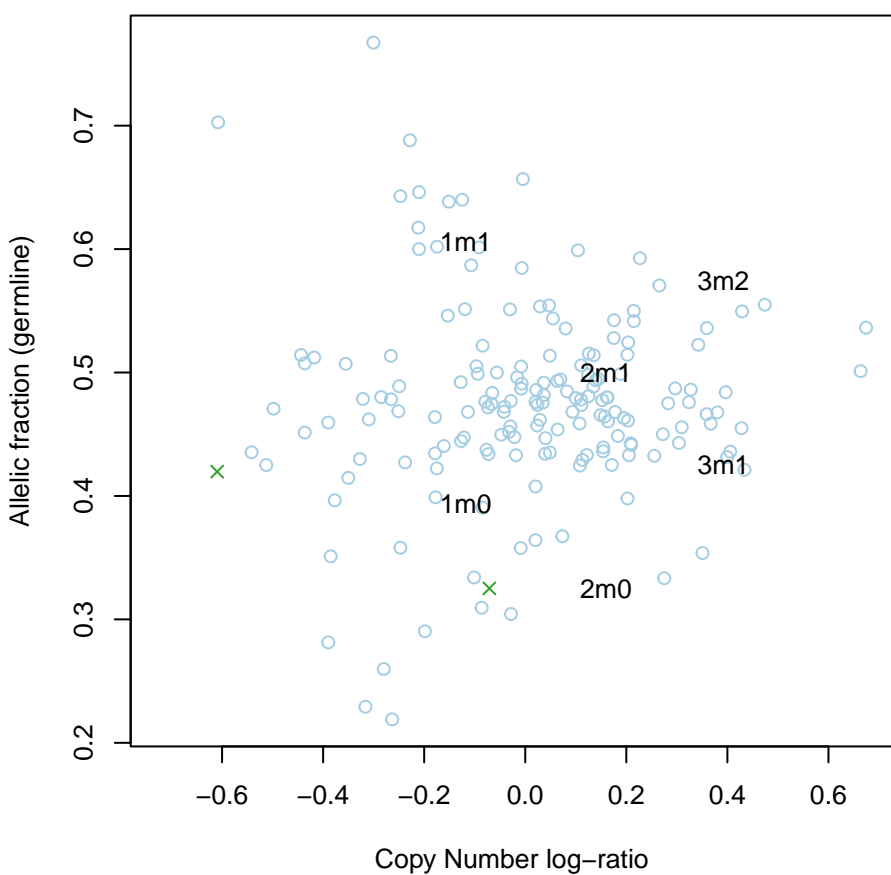


Purity: 0.35 Tumor ploidy: 1.401 SNV log-likelihood: -331.73 GoF: 79.1% Mean coverage: 514,449

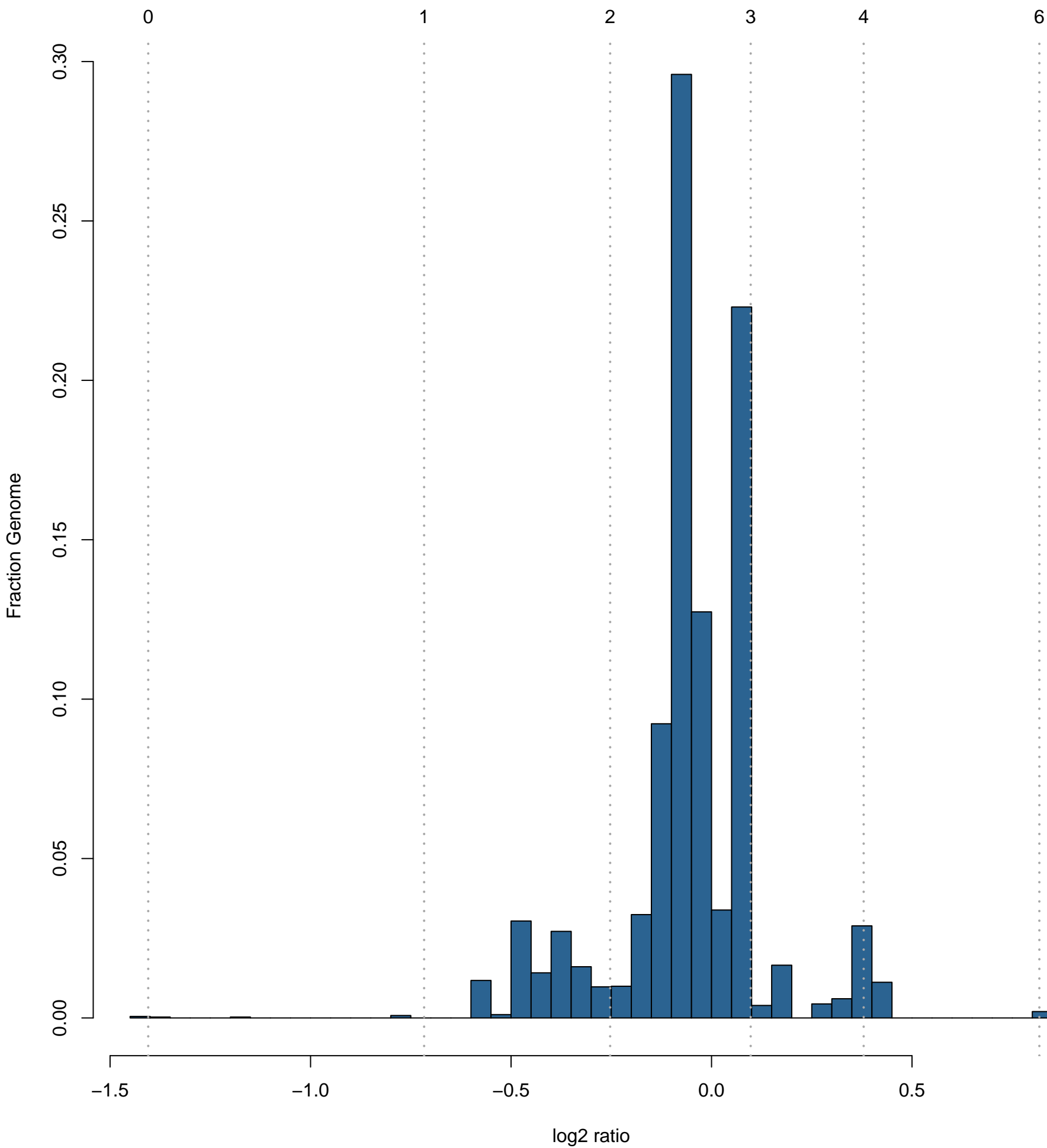


SCNA-fit log-likelihood: -4949.52



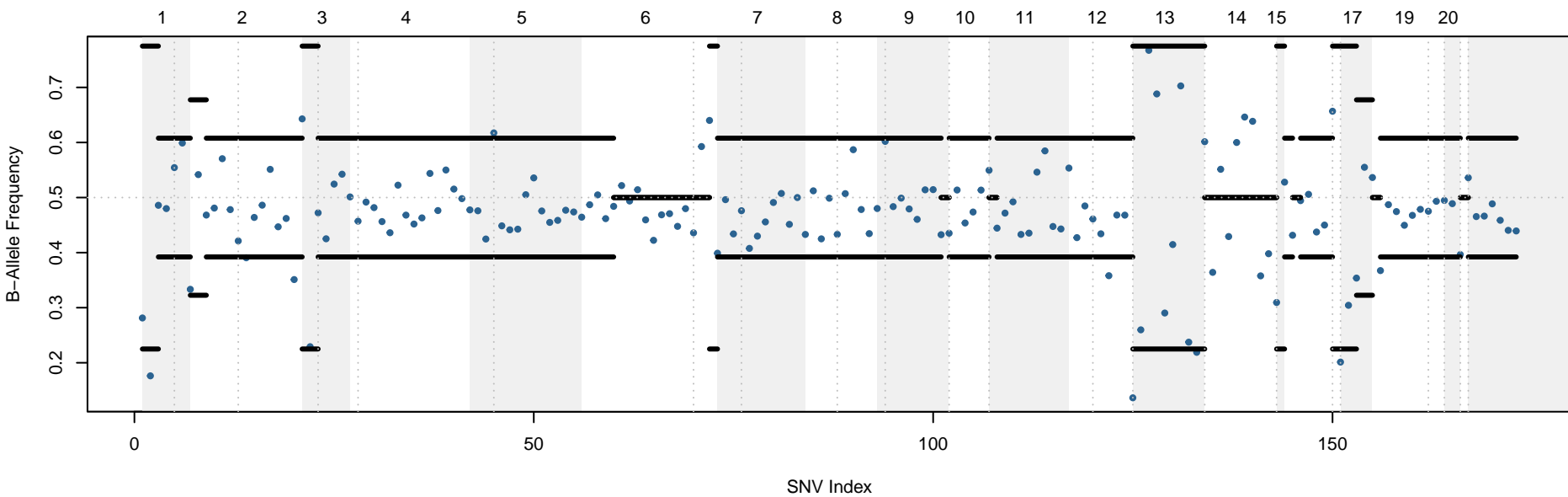


Purity: 0.55 Tumor ploidy: 2.697

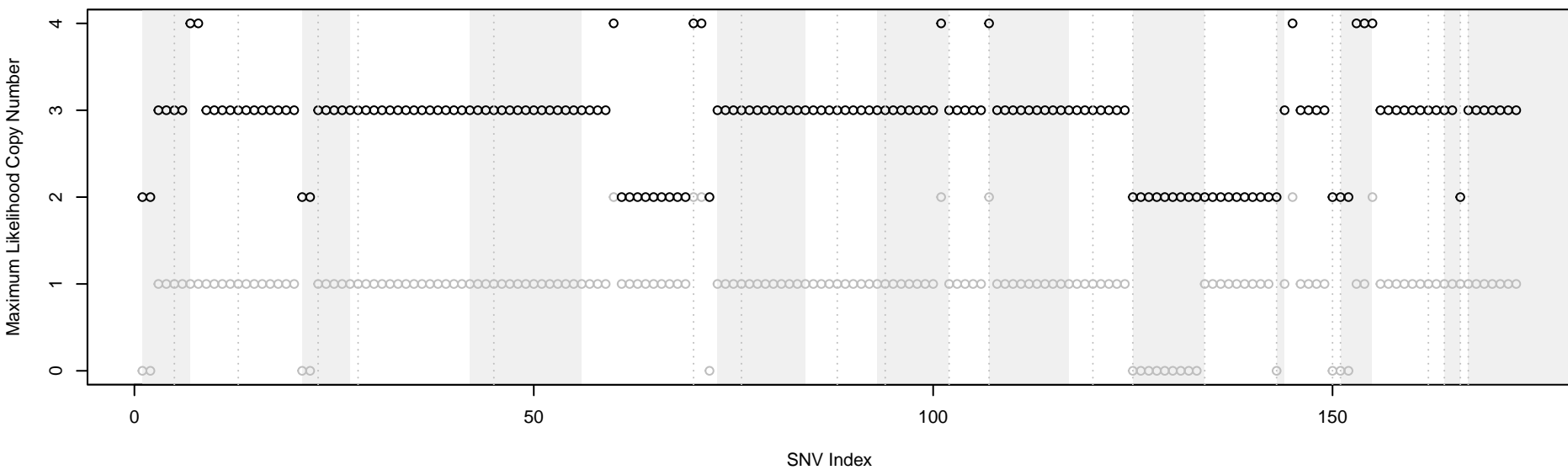
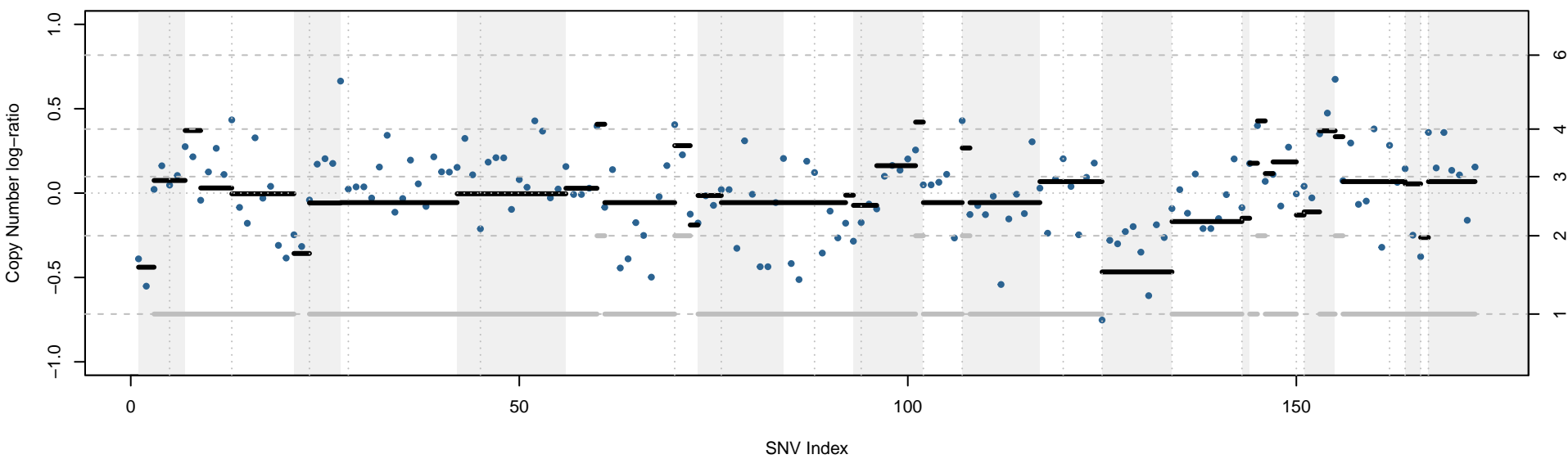


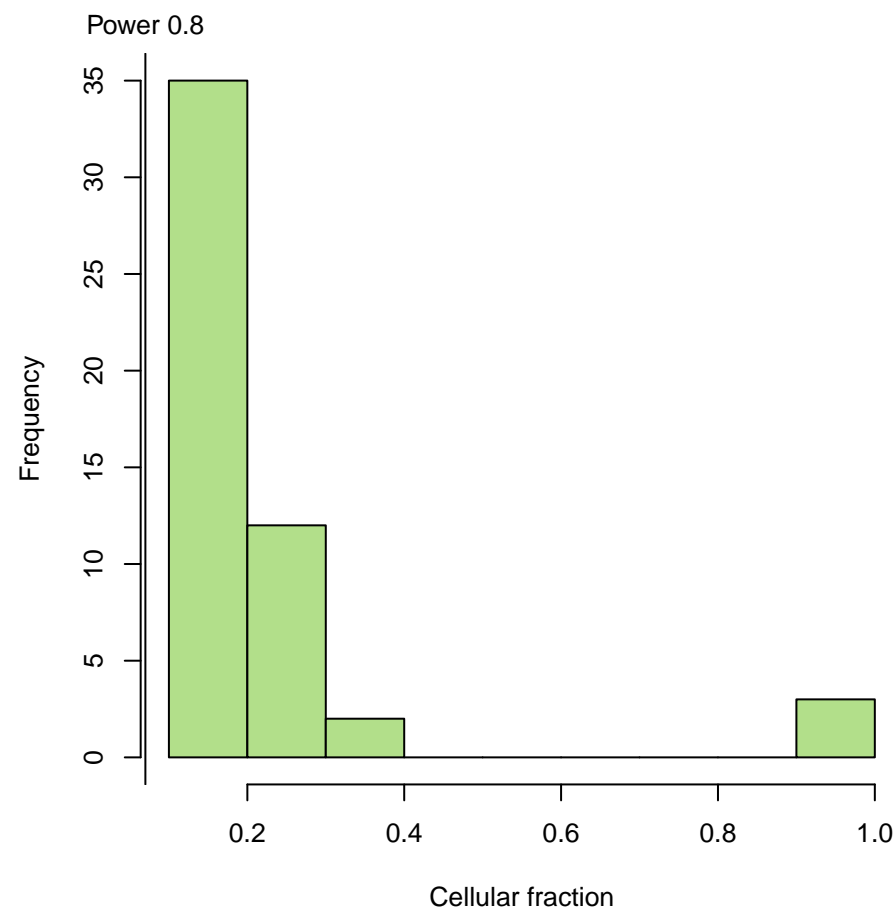
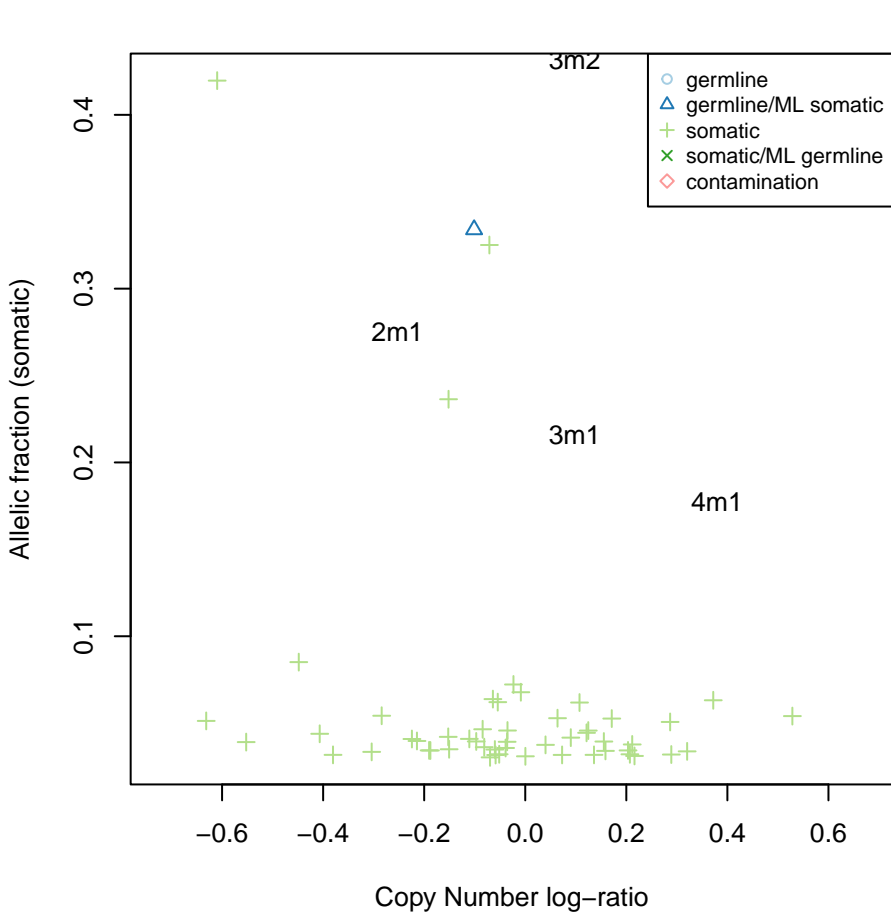
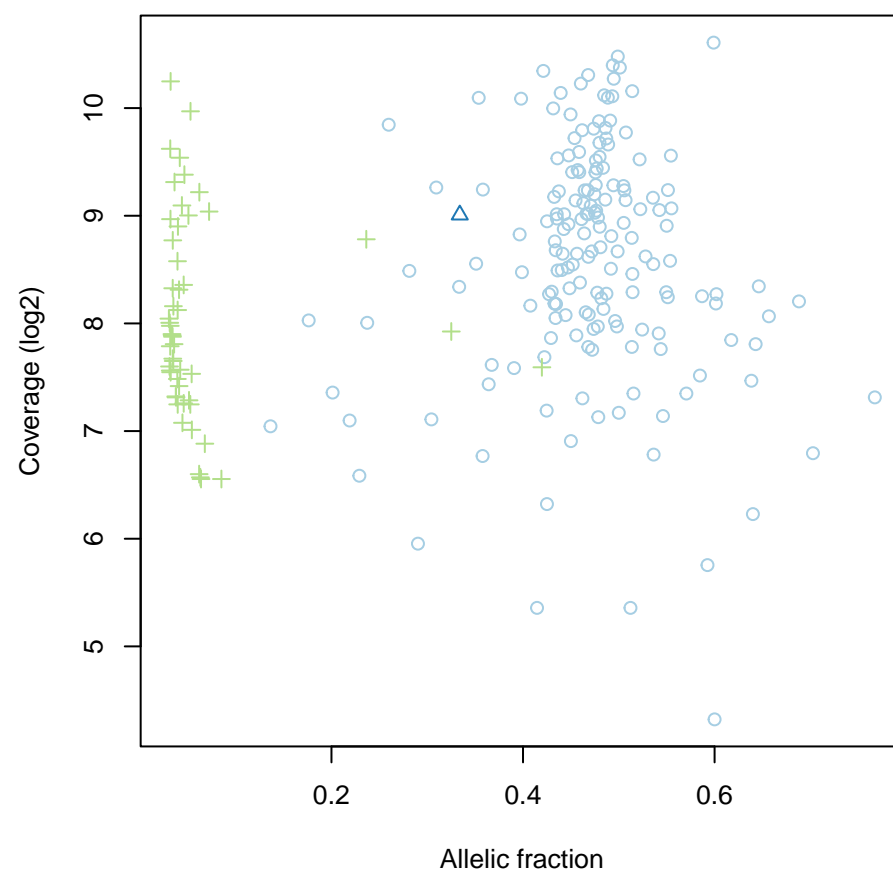
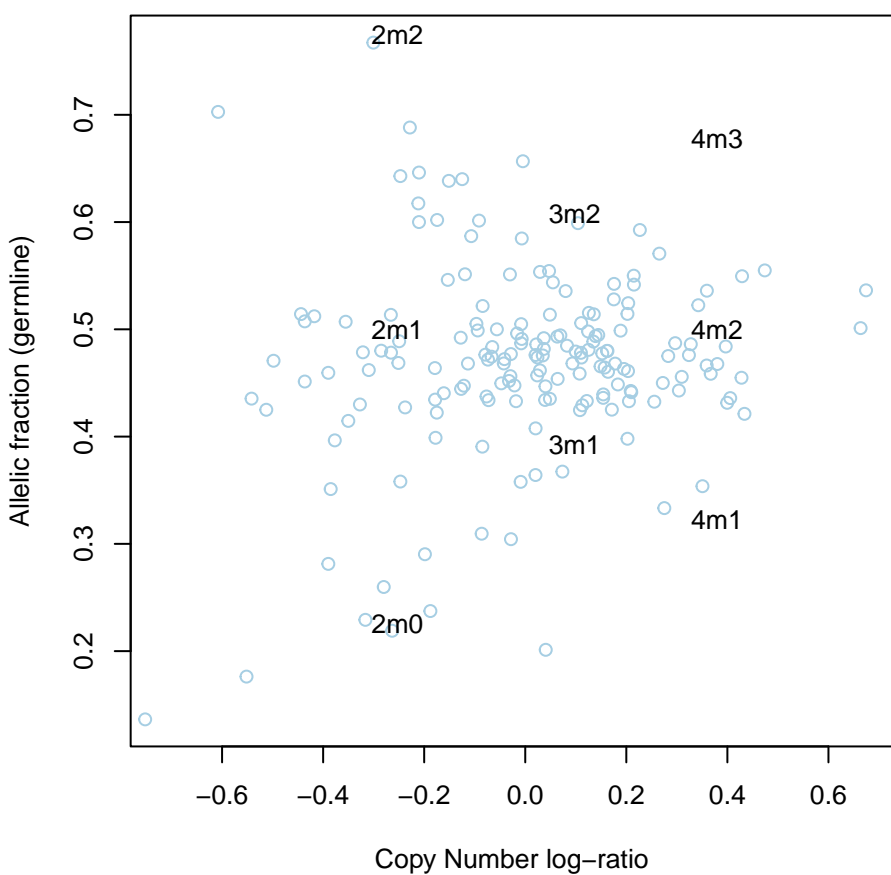


Purity: 0.55 Tumor ploidy: 2.697 SNV log-likelihood: -562.15 GoF: 67.5% Mean coverage: 514,449

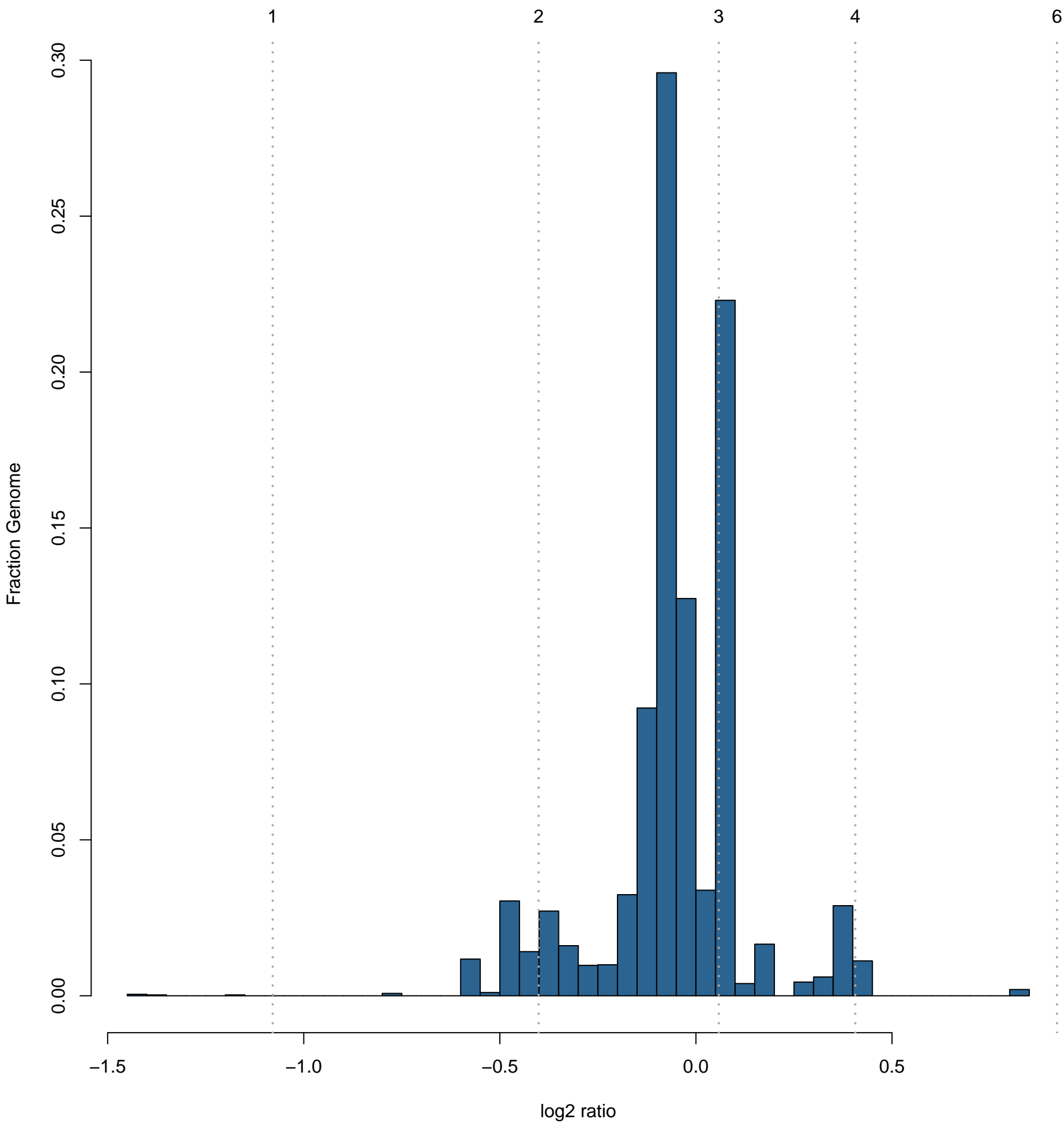


SCNA-fit log-likelihood: -4808.38

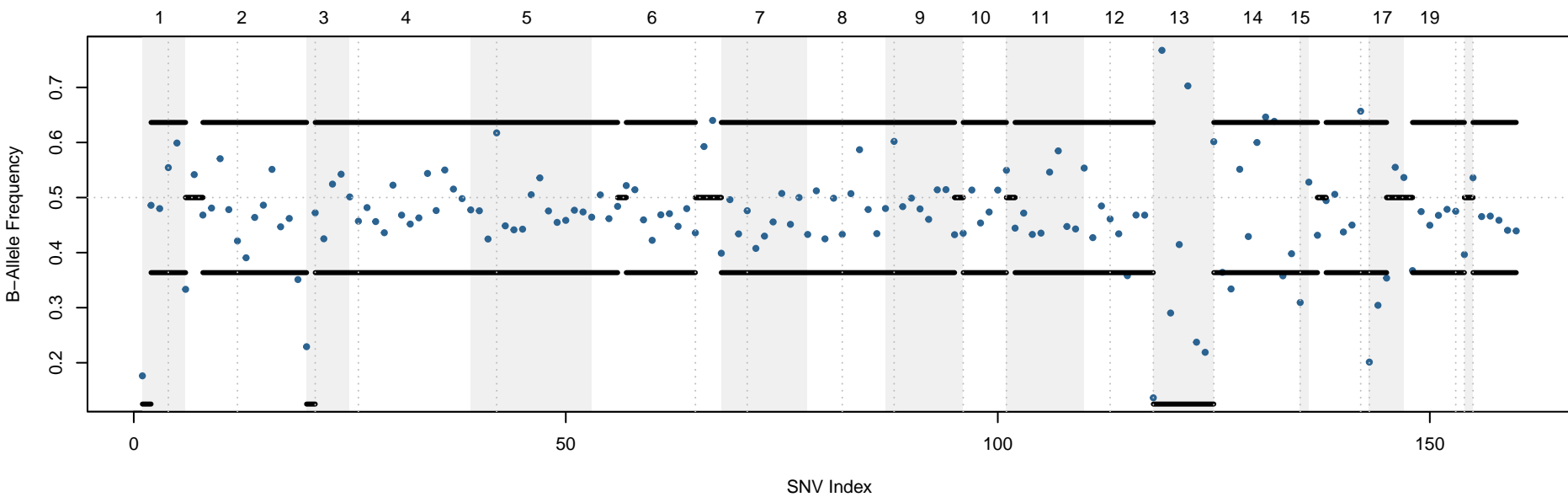




Purity: 0.75 Tumor ploidy: 2.855



Purity: 0.75 Tumor ploidy: 2.855 SNV log-likelihood: -1110.61 GoF: 50.3% Mean coverage: 514;449



SCNA-fit log-likelihood: -4719.38

