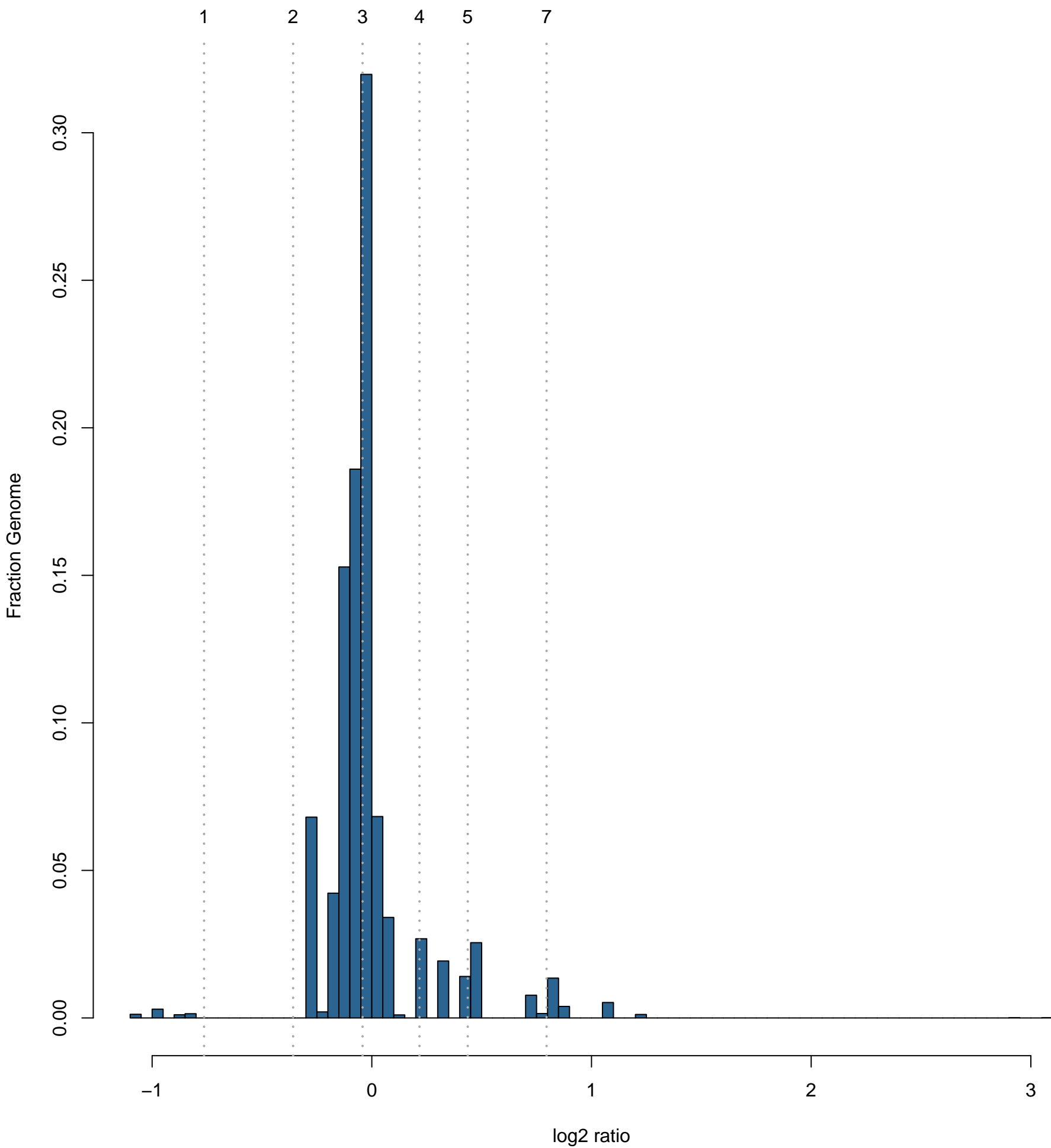
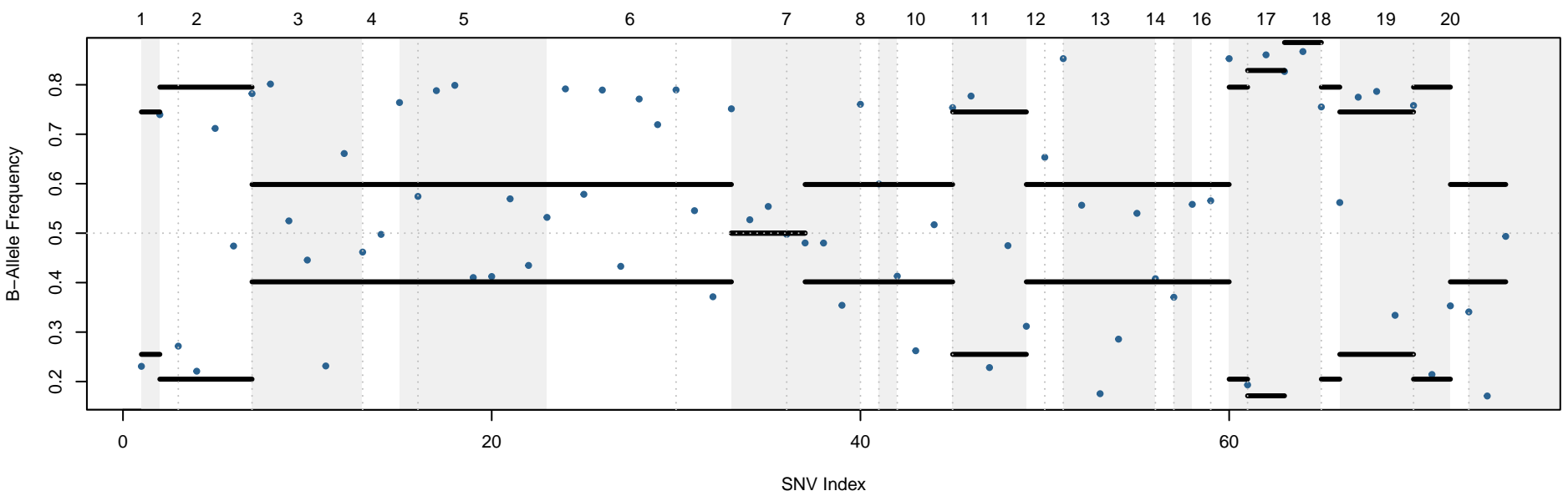


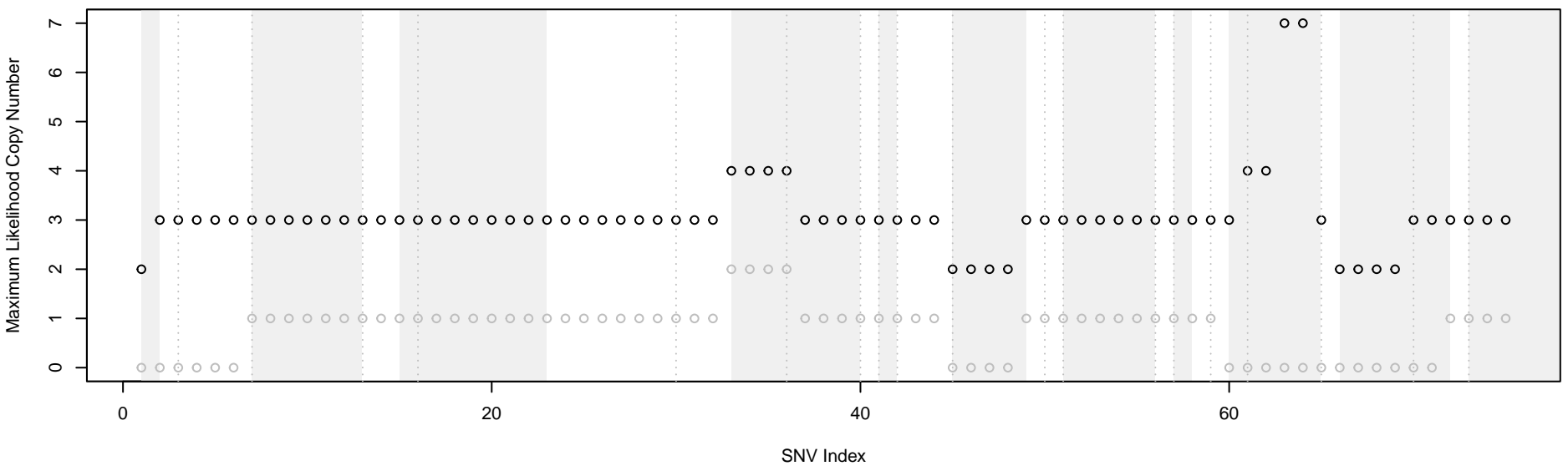
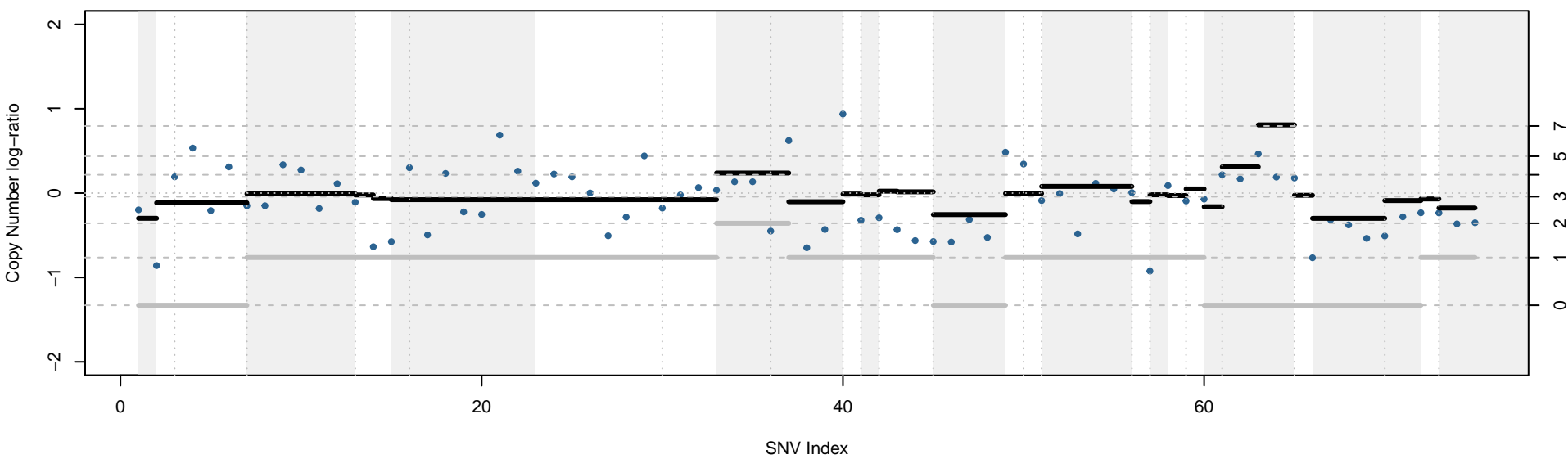
Purity: 0.49 Tumor ploidy: 3.151

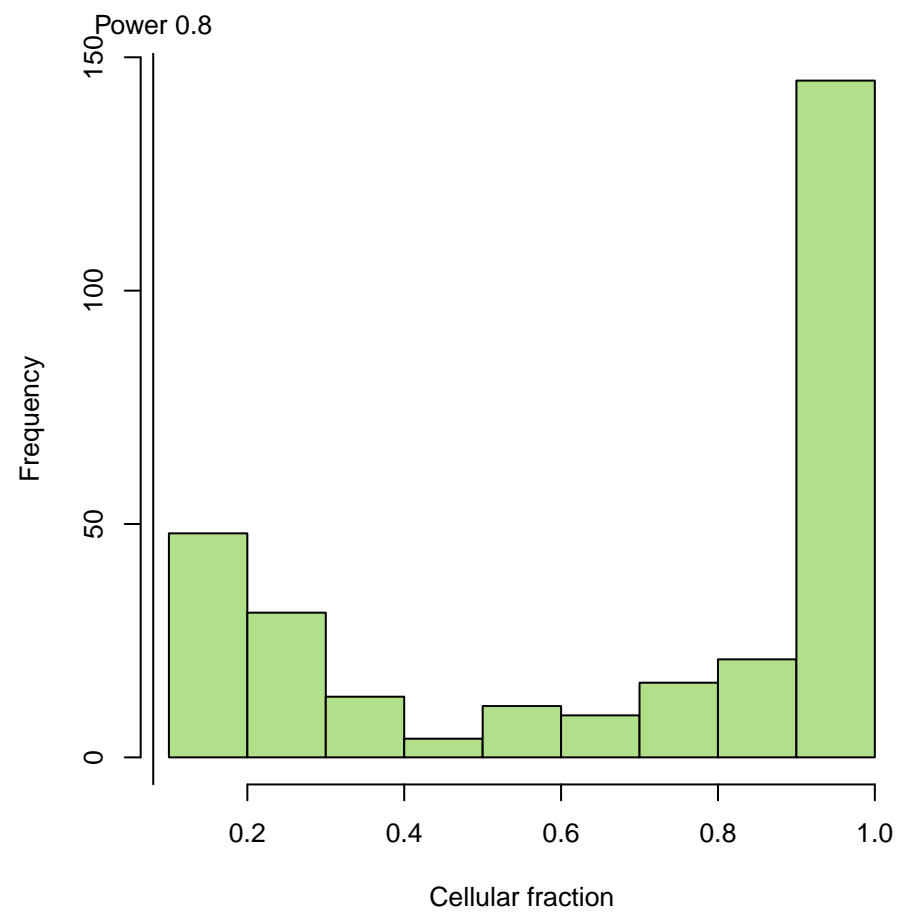
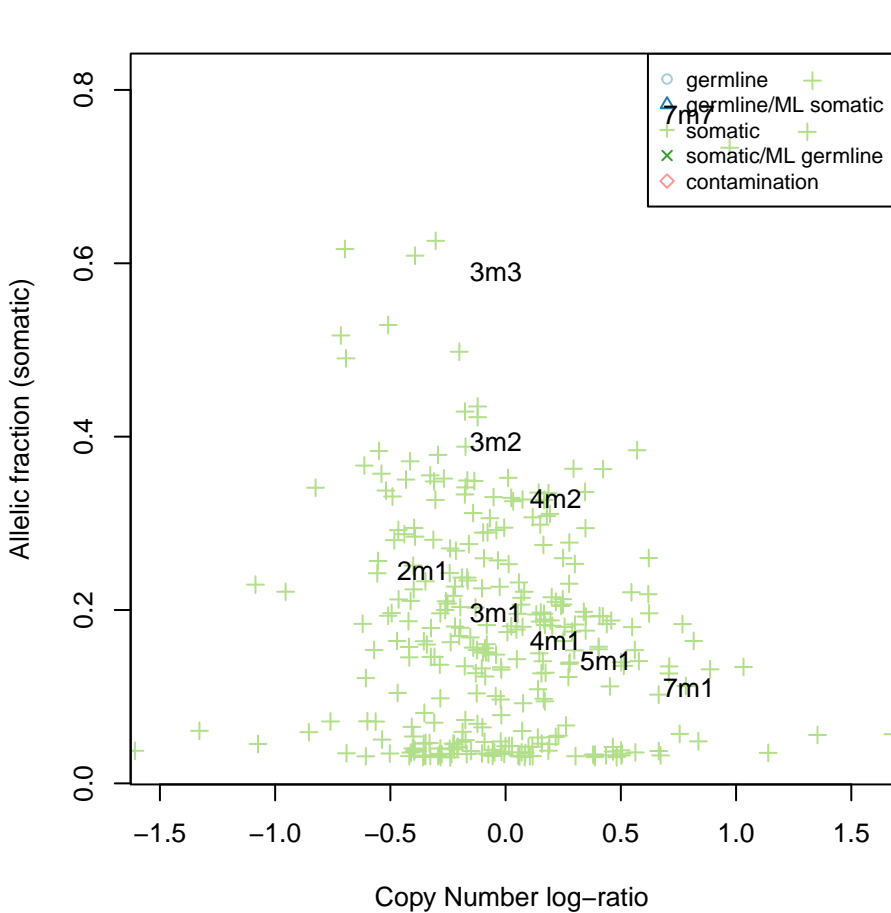
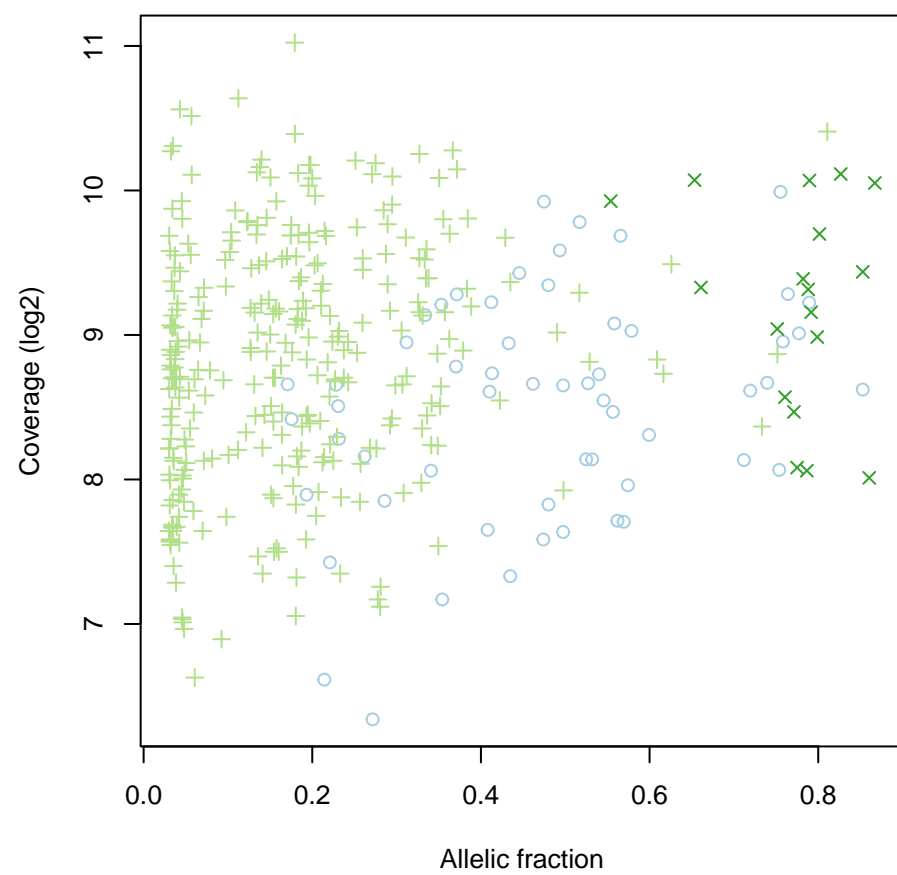
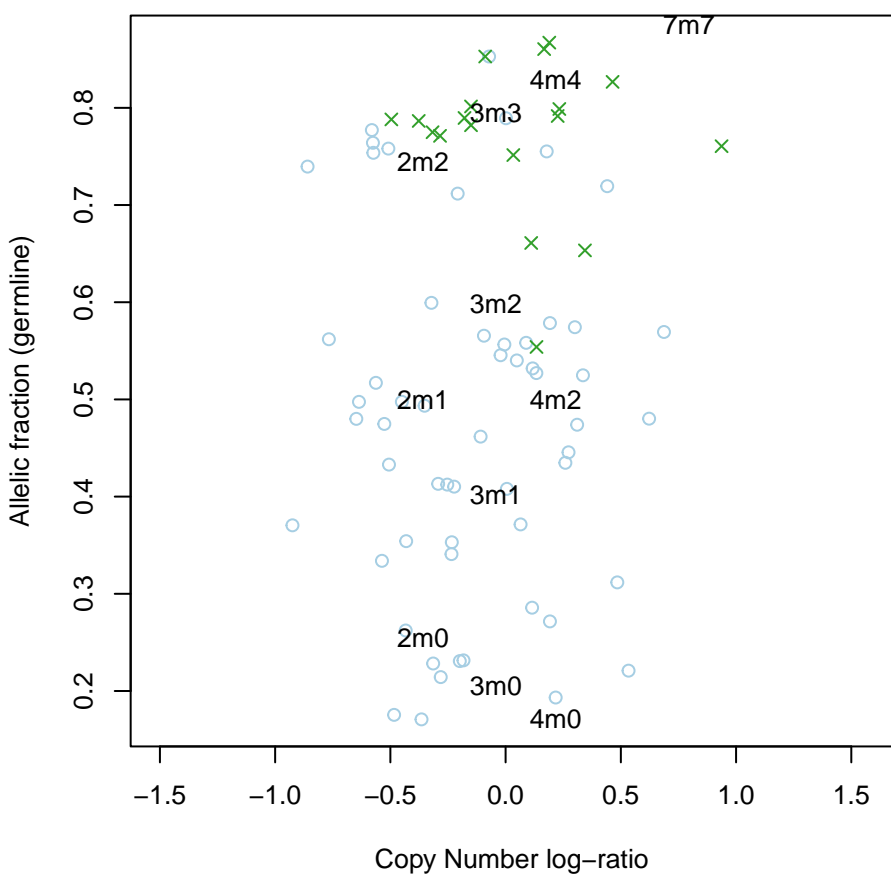


Purity: 0.49 Tumor ploidy: 3.151 SNV log-likelihood: -736.83 GoF: 79.1% Mean coverage: 522;32

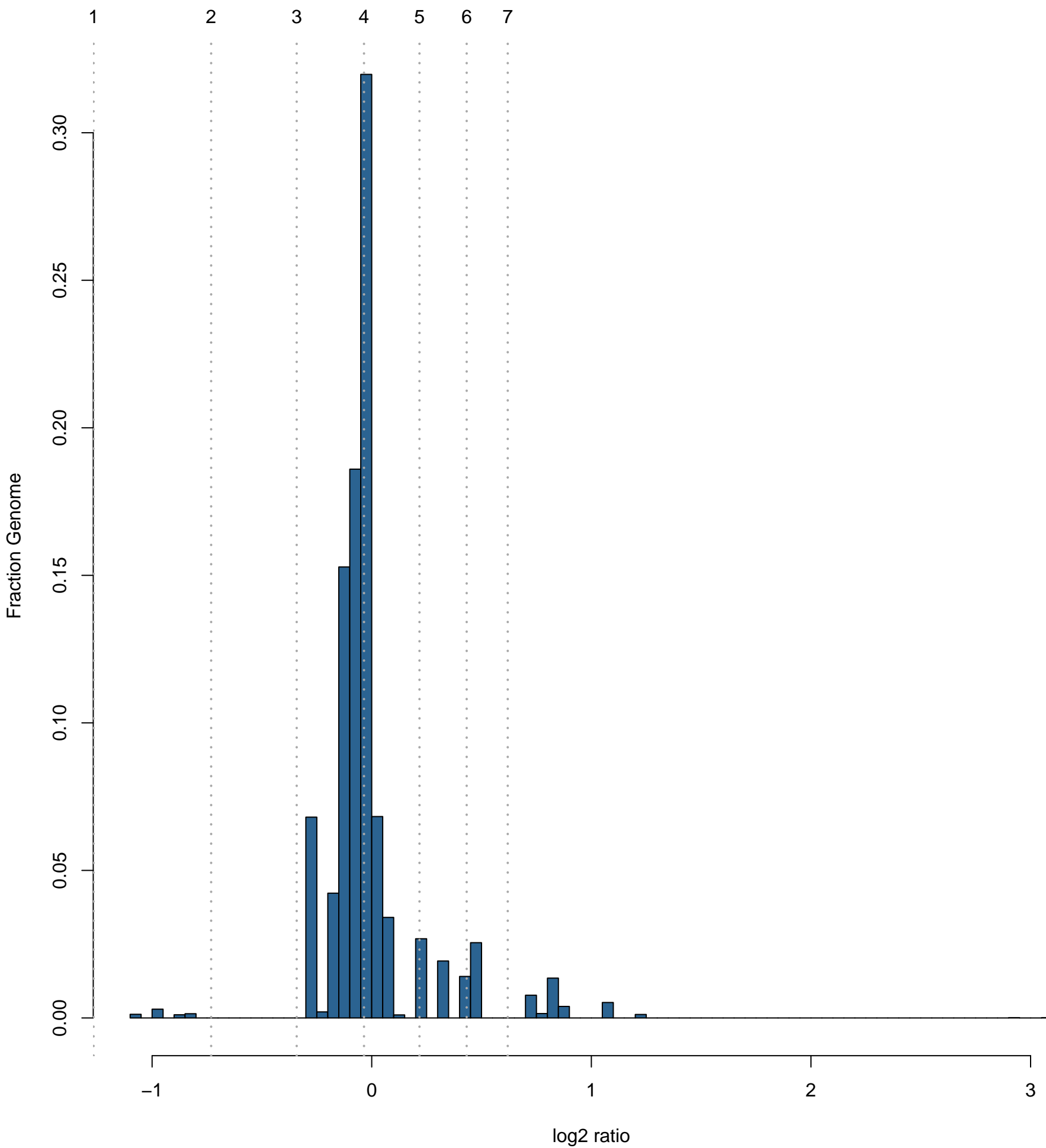


SCNA-fit log-likelihood: -16926.19

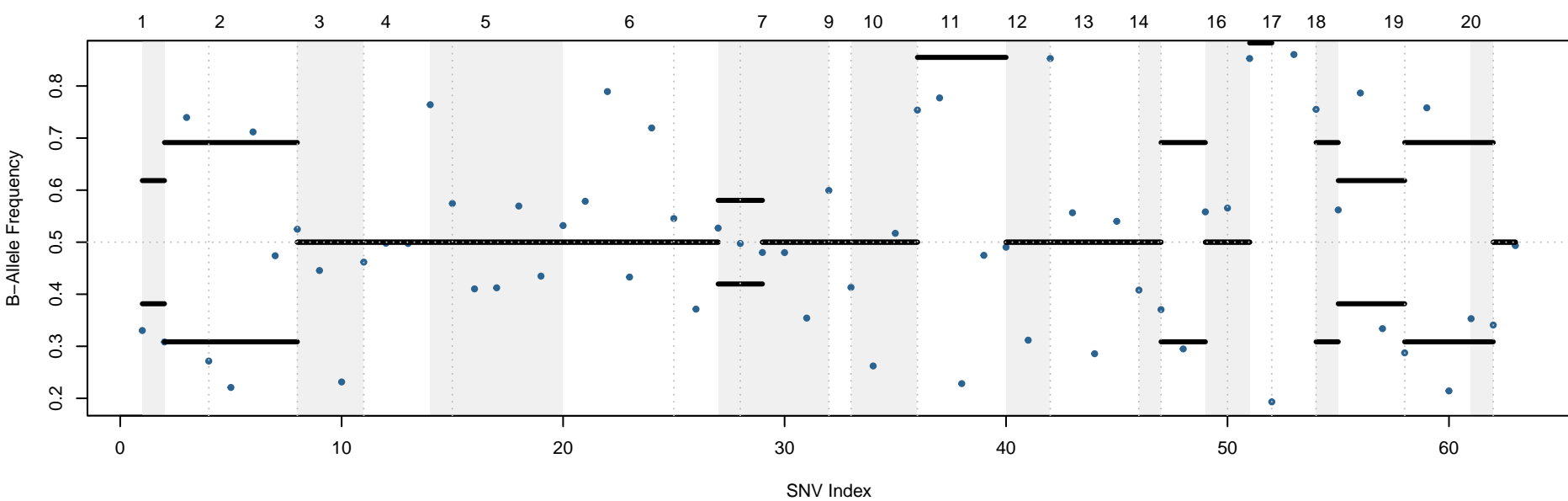




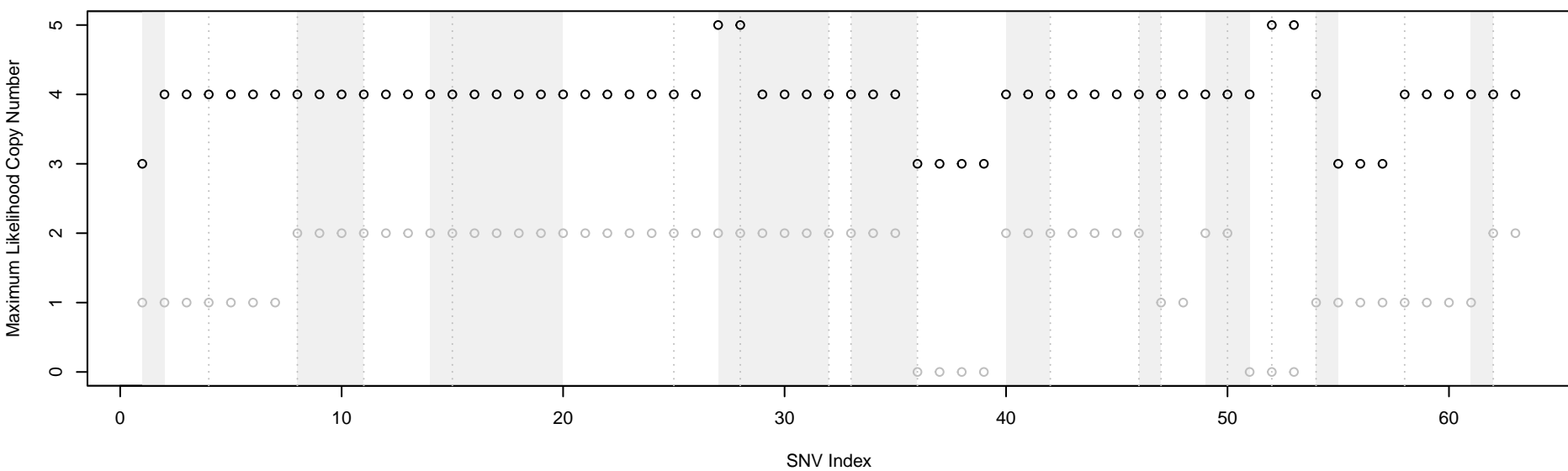
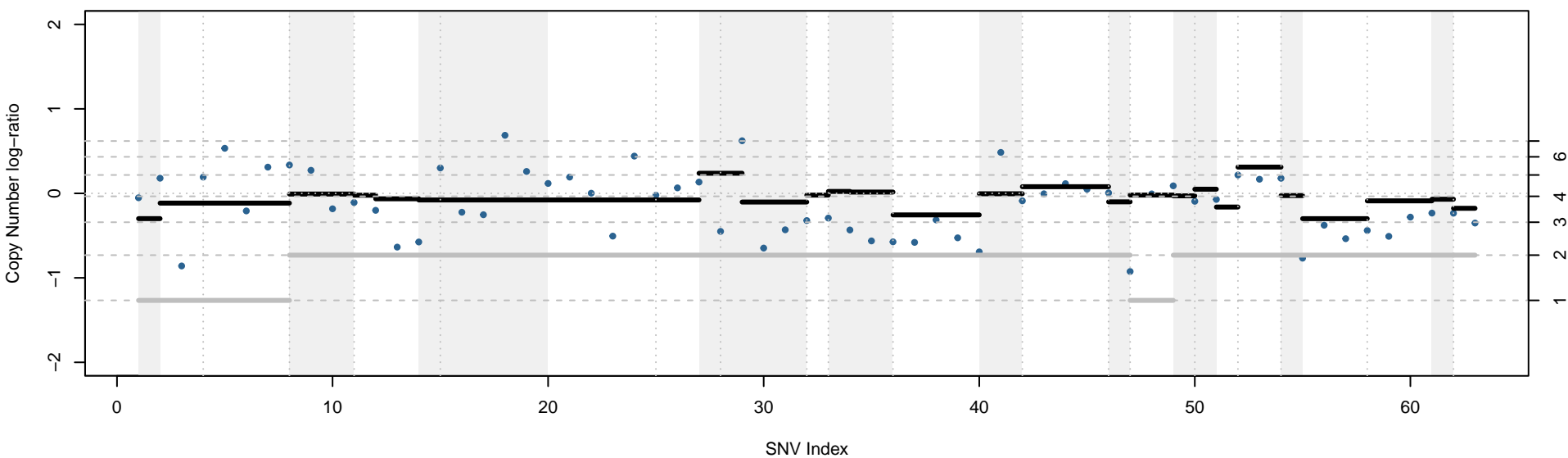
Purity: 0.62 Tumor ploidy: 4.129

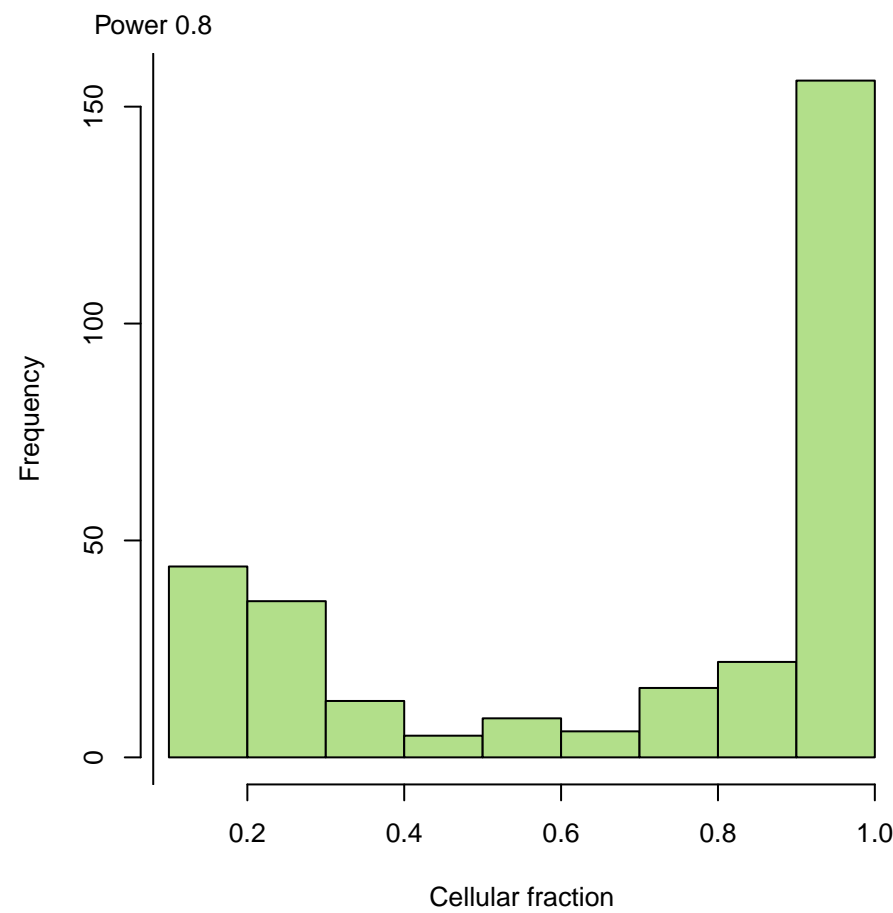
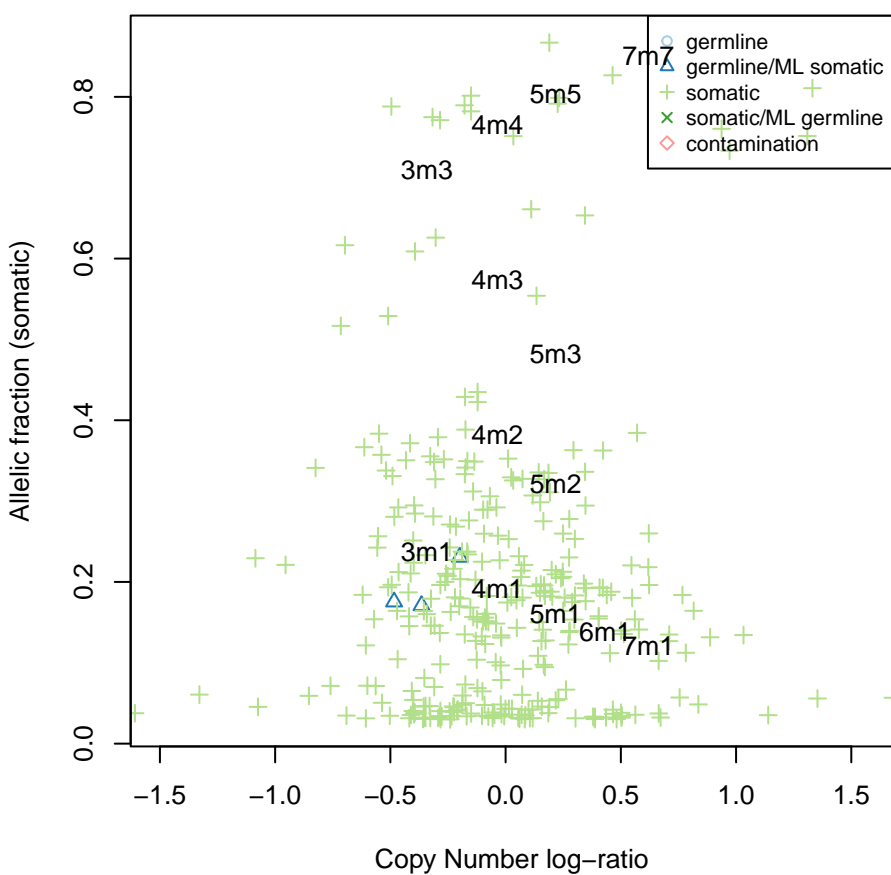
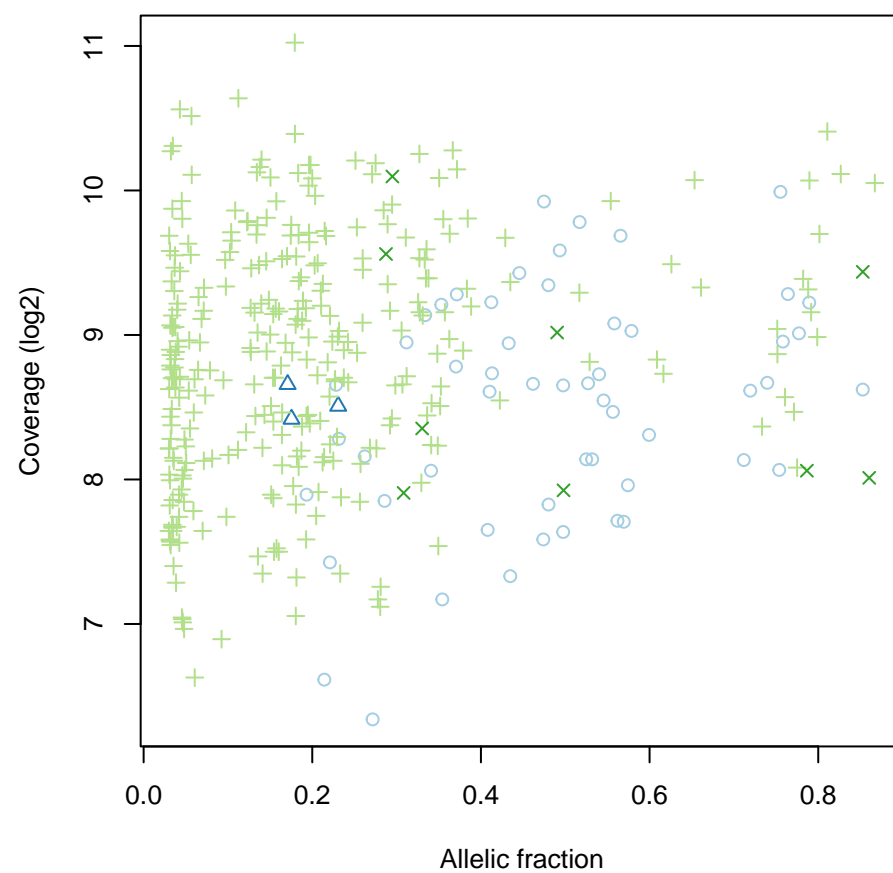
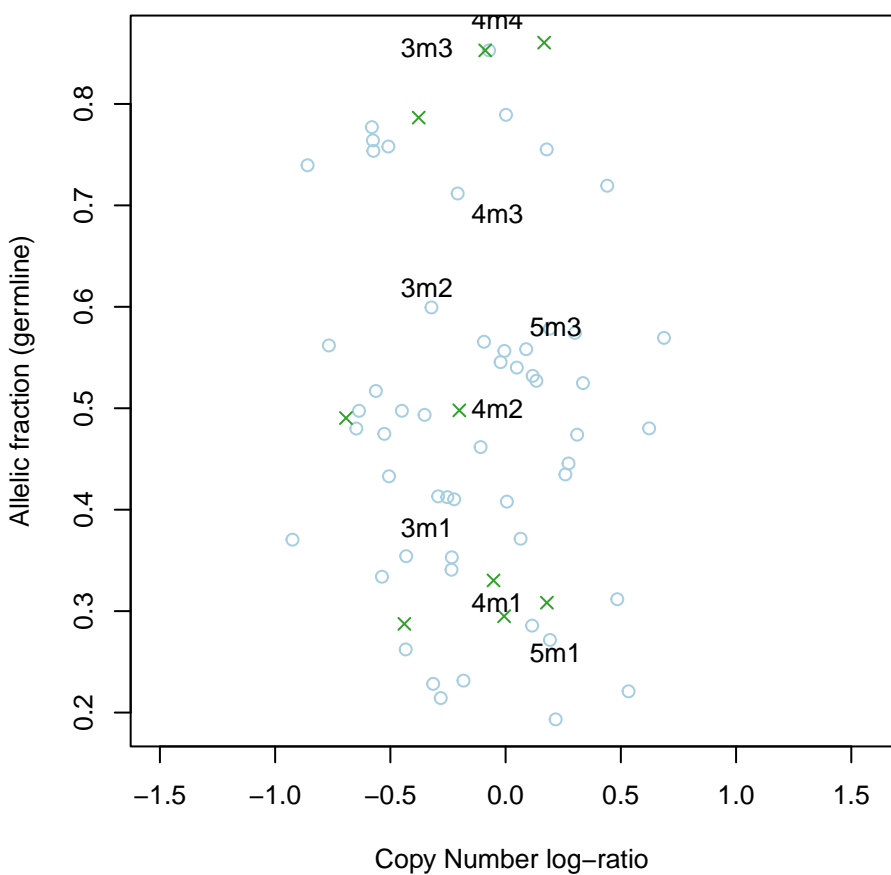


Purity: 0.62 Tumor ploidy: 4.129 SNV log-likelihood: -801.86 GoF: 79.7% Mean coverage: 522;32

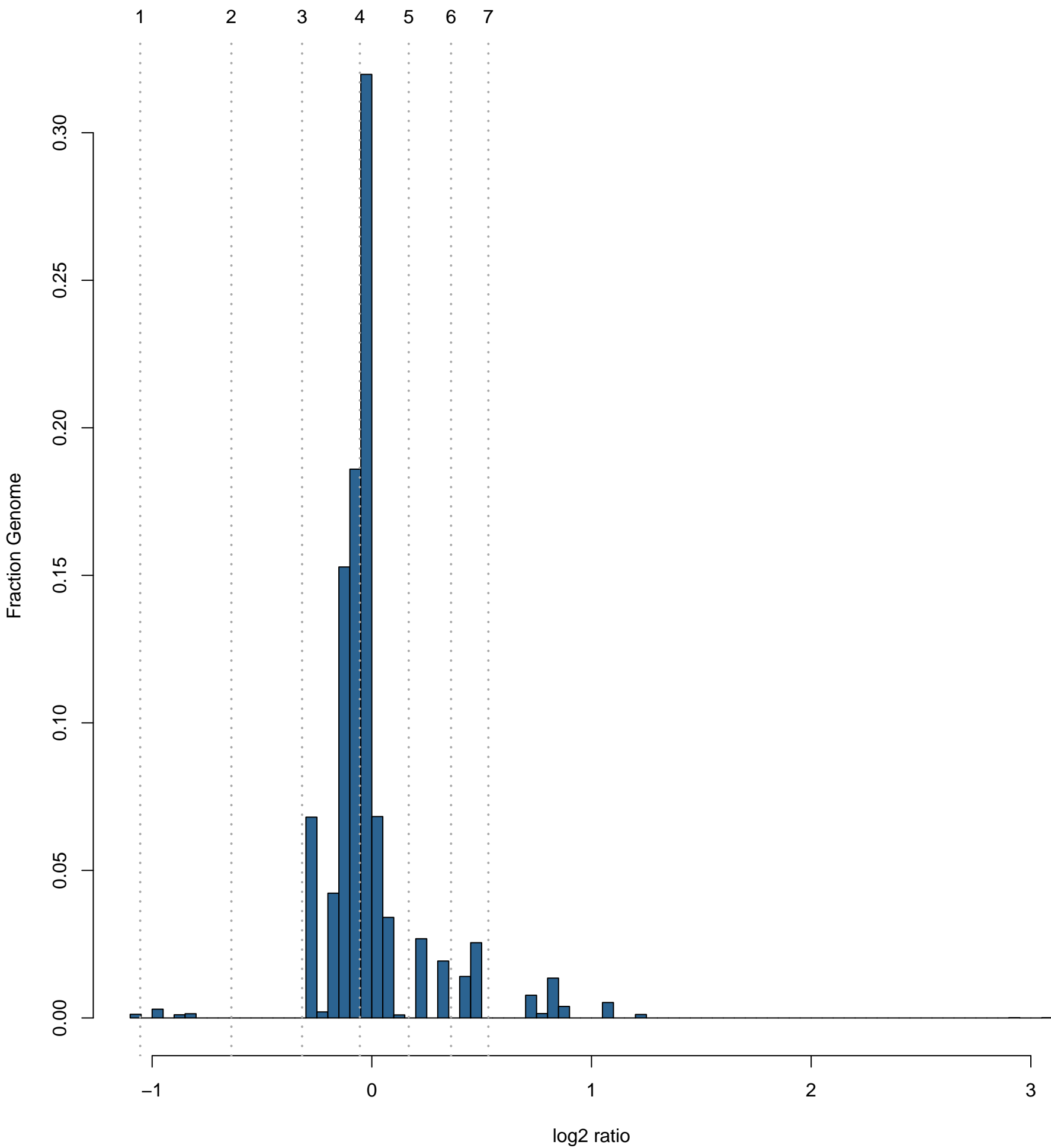


SCNA-fit log-likelihood: -16948.28

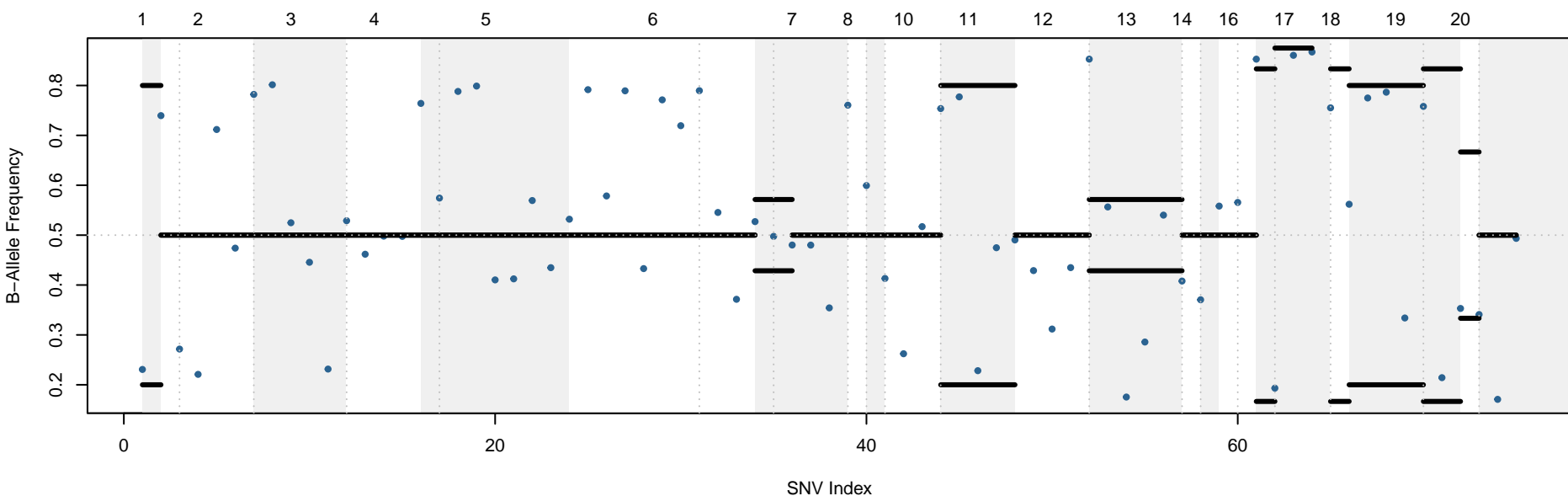




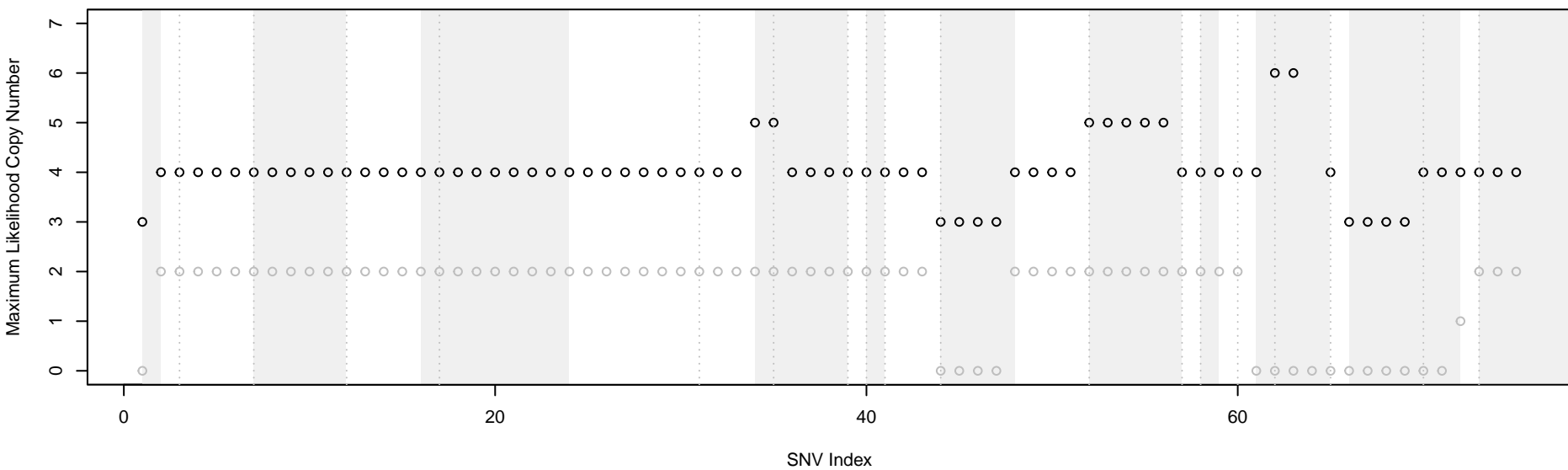
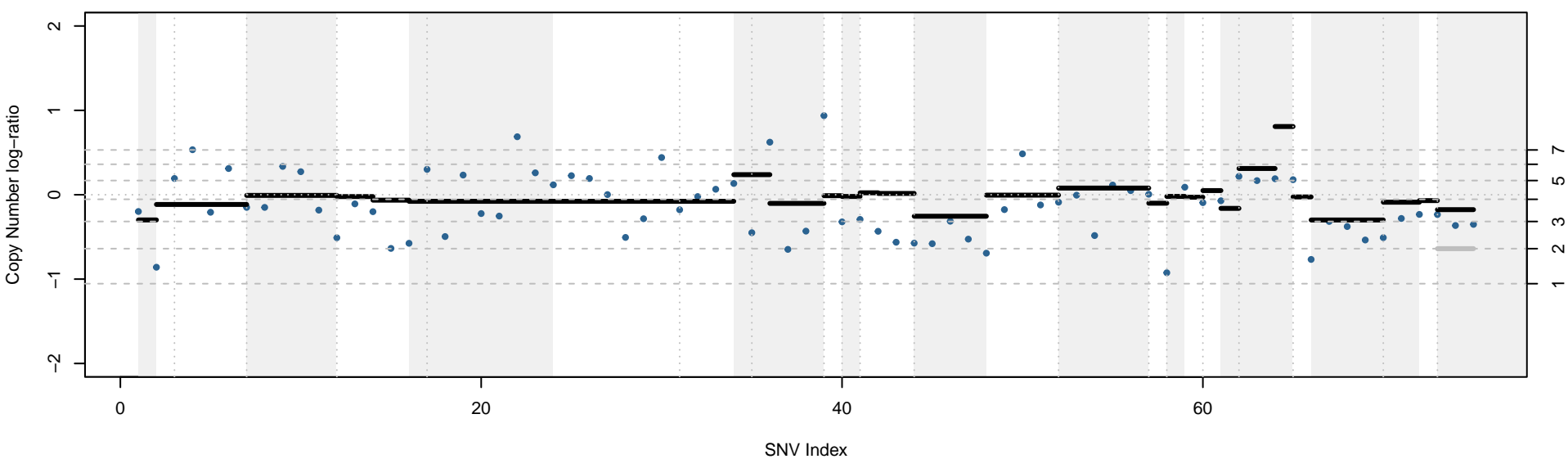
Purity: 0.5 Tumor ploidy: 4.23



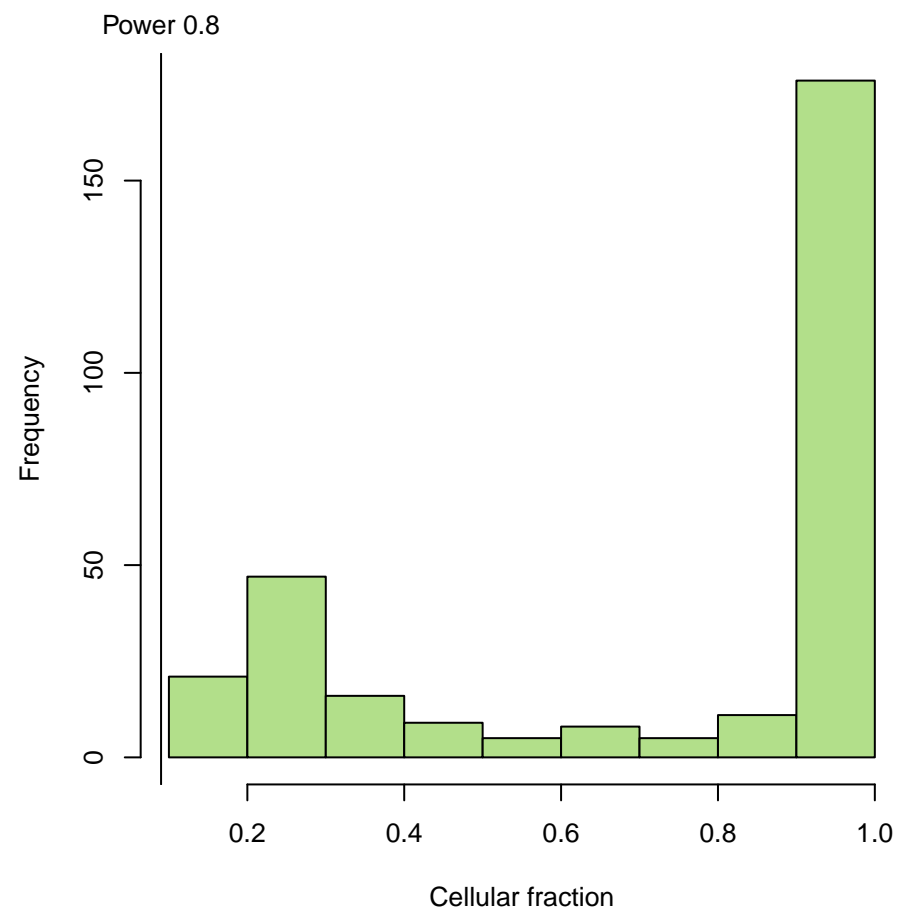
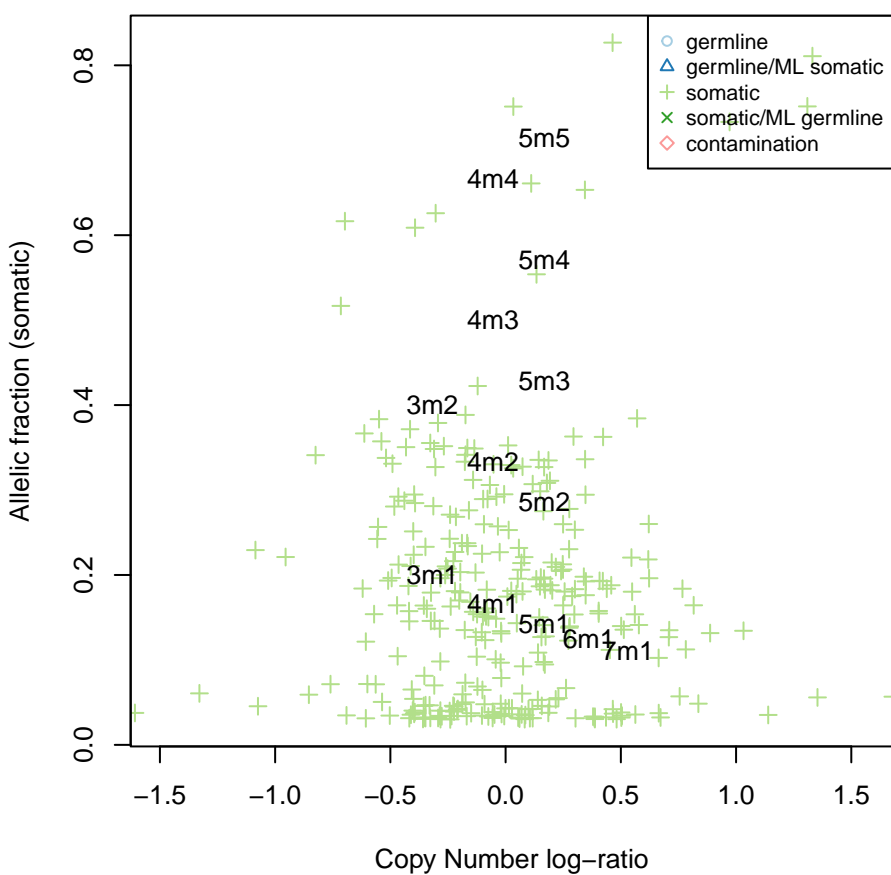
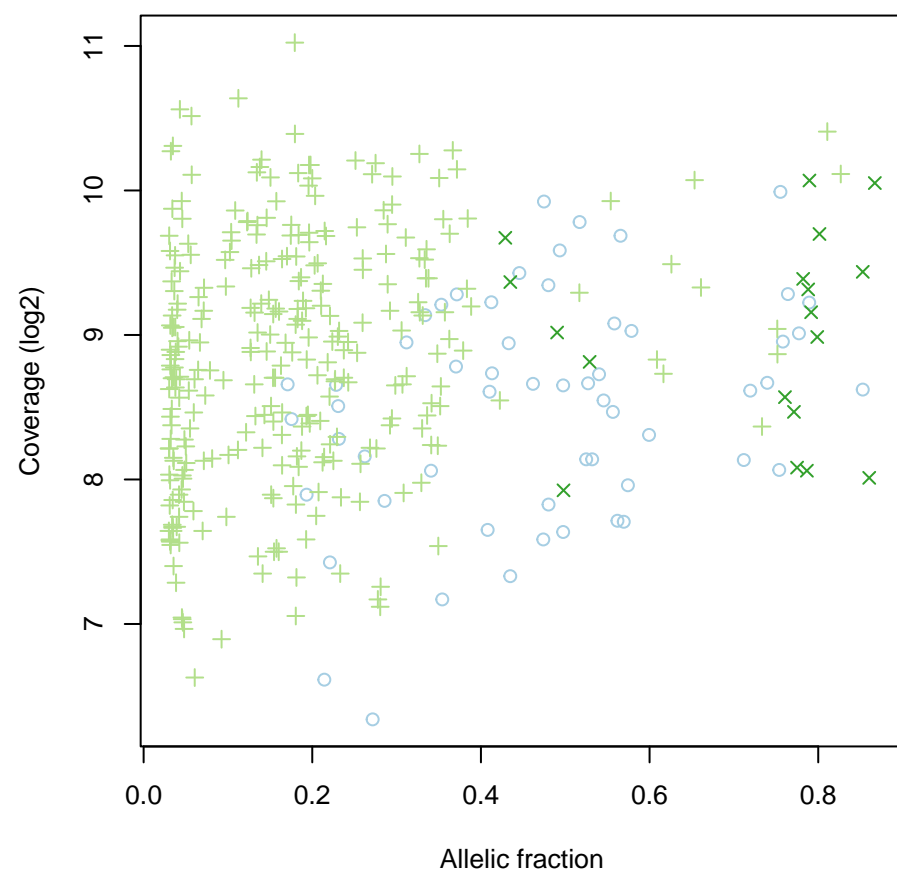
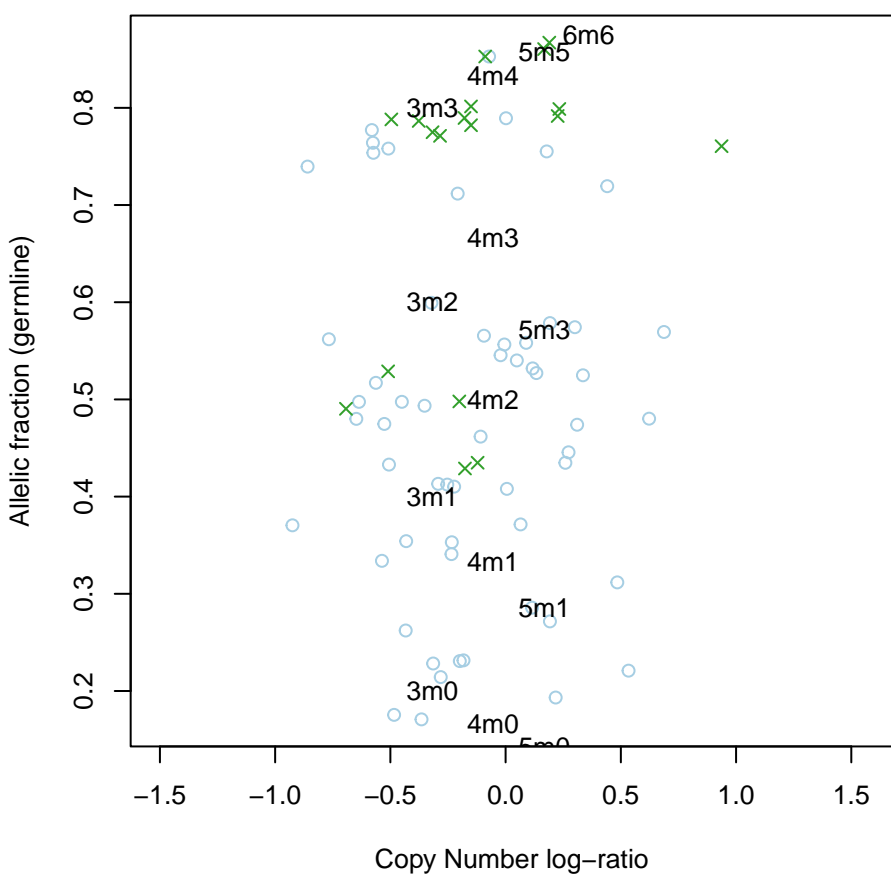
Purity: 0.5 Tumor ploidy: 4.23 SNV log-likelihood: -770.68 GoF: 84.3% Mean coverage: 522;32



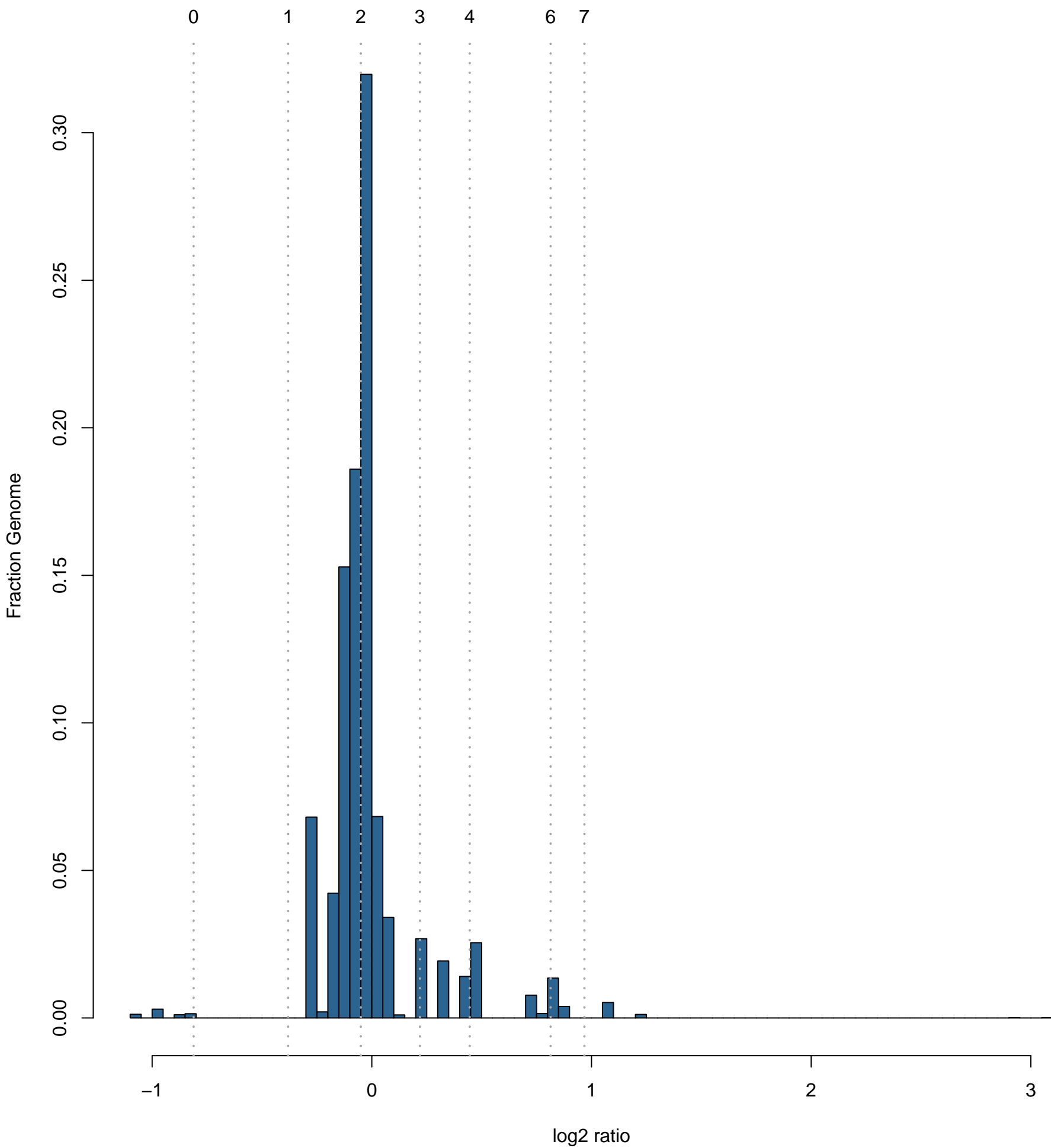
SCNA-fit log-likelihood: -17029.23



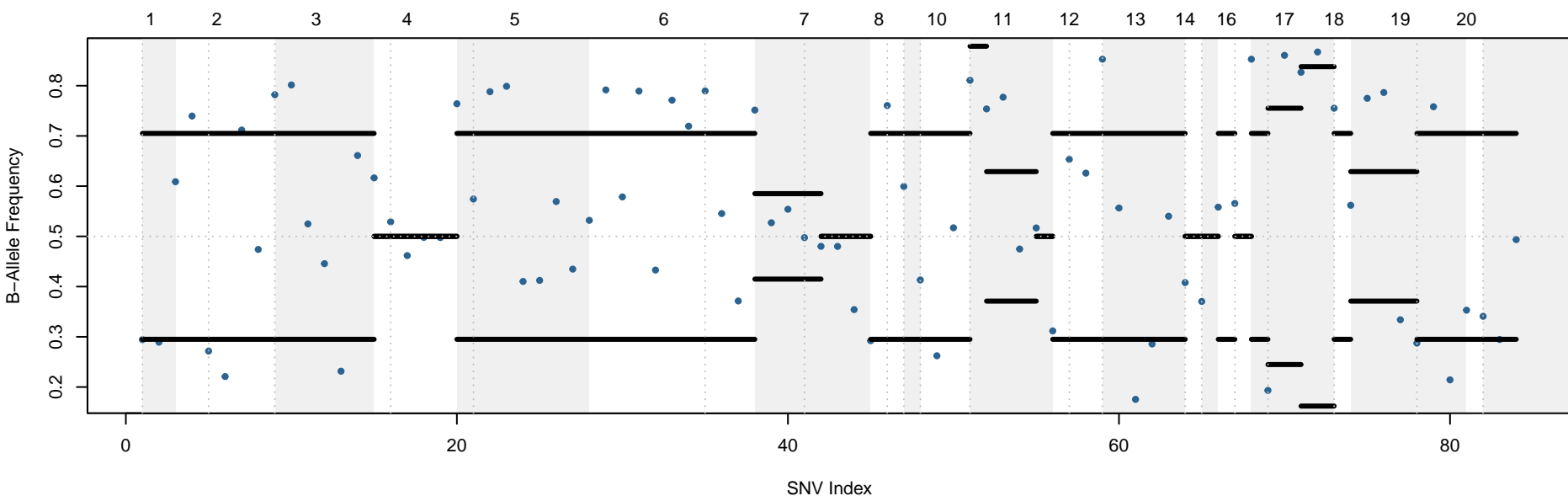




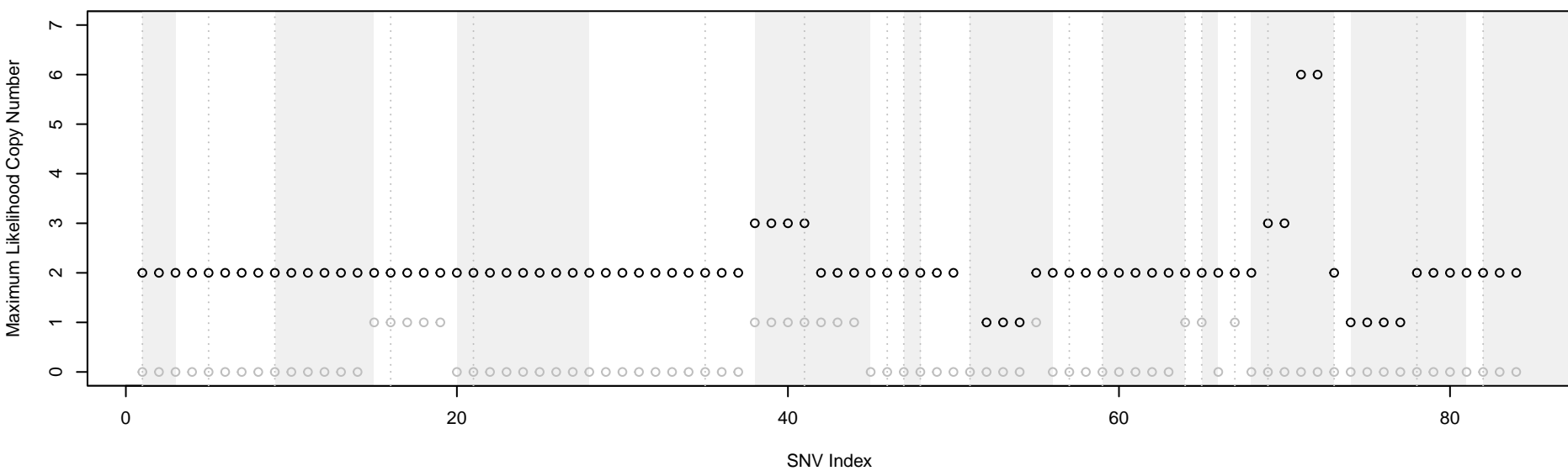
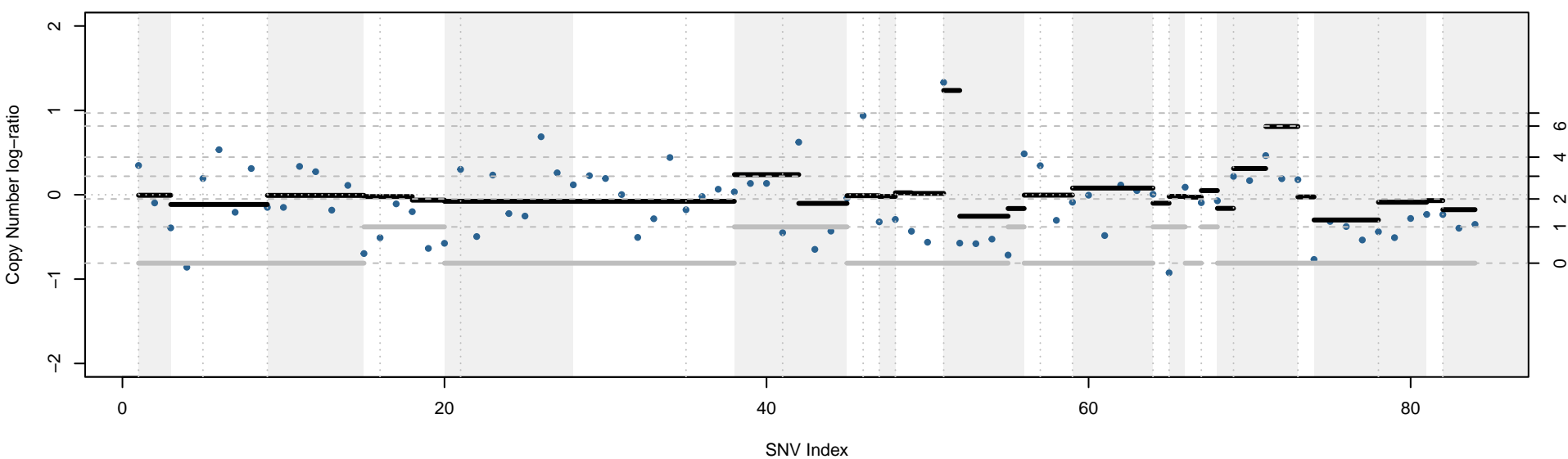
Purity: 0.41 Tumor ploidy: 2.172

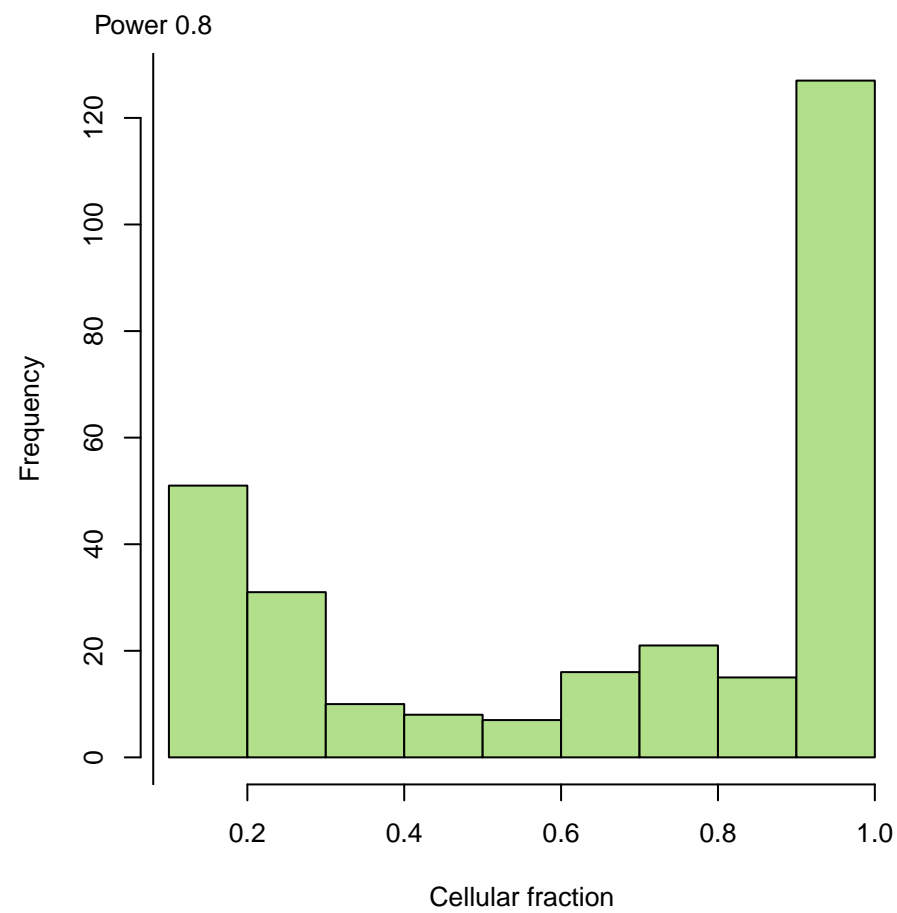
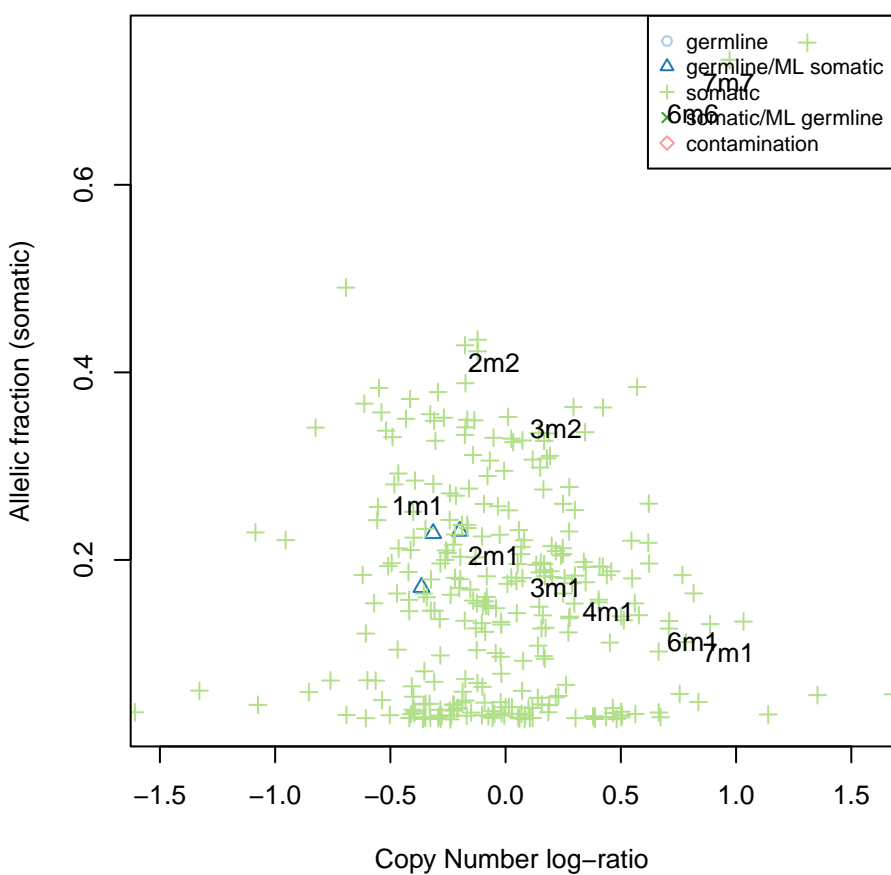
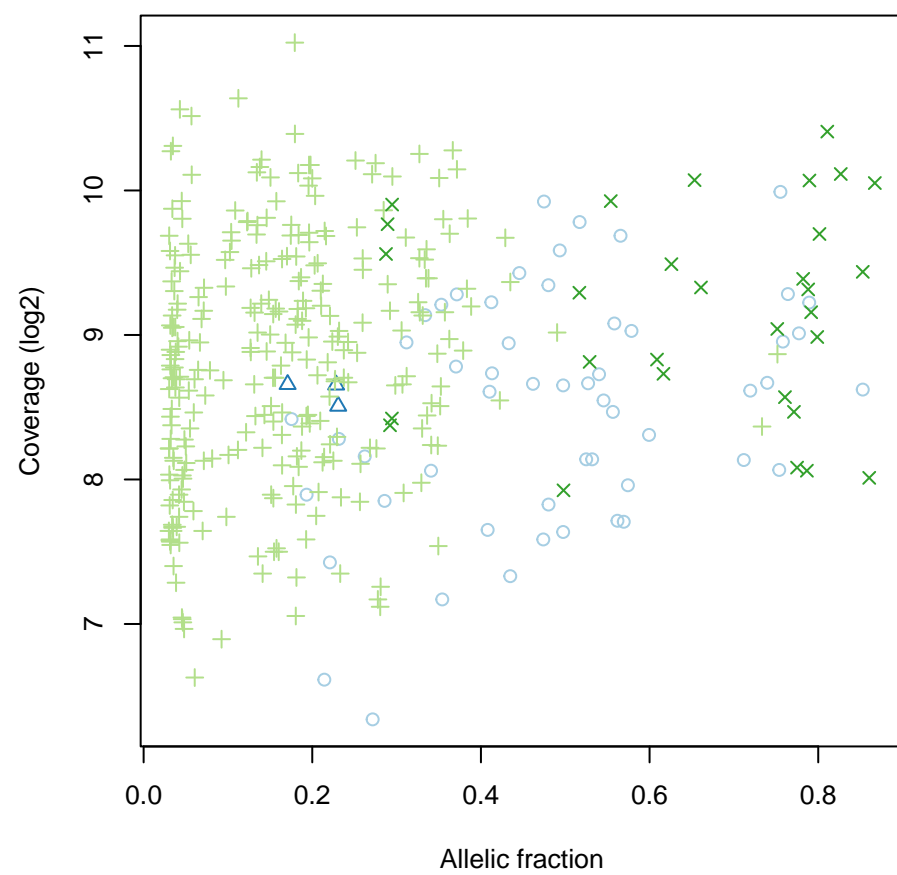
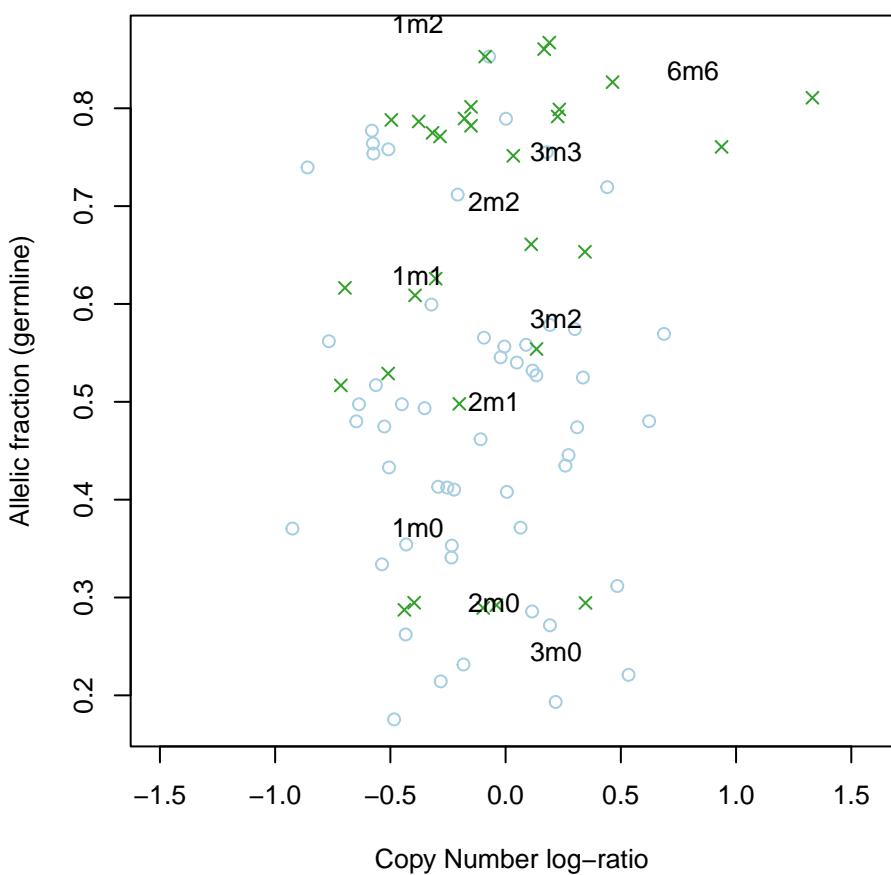


Purity: 0.41 Tumor ploidy: 2.172 SNV log-likelihood: -952.19 GoF: 75.8% Mean coverage: 522;32

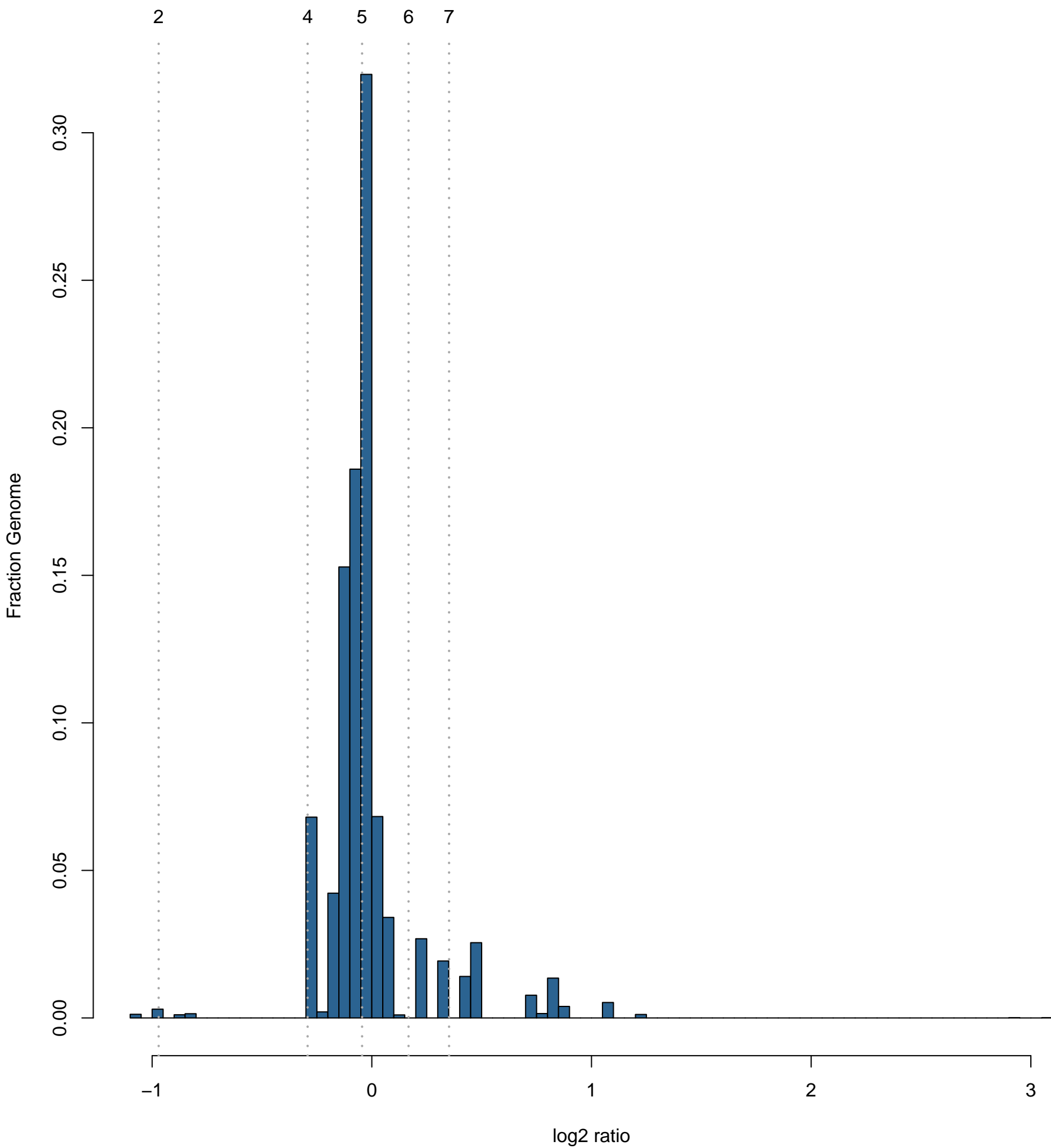


SCNA-fit log-likelihood: -16914.83

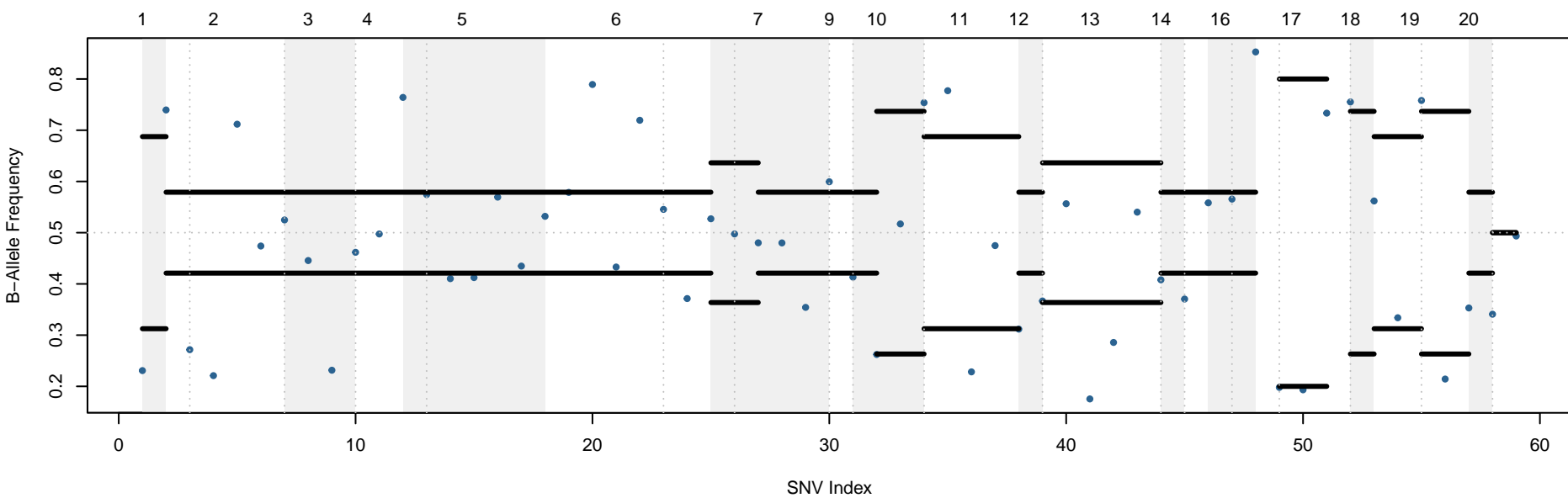




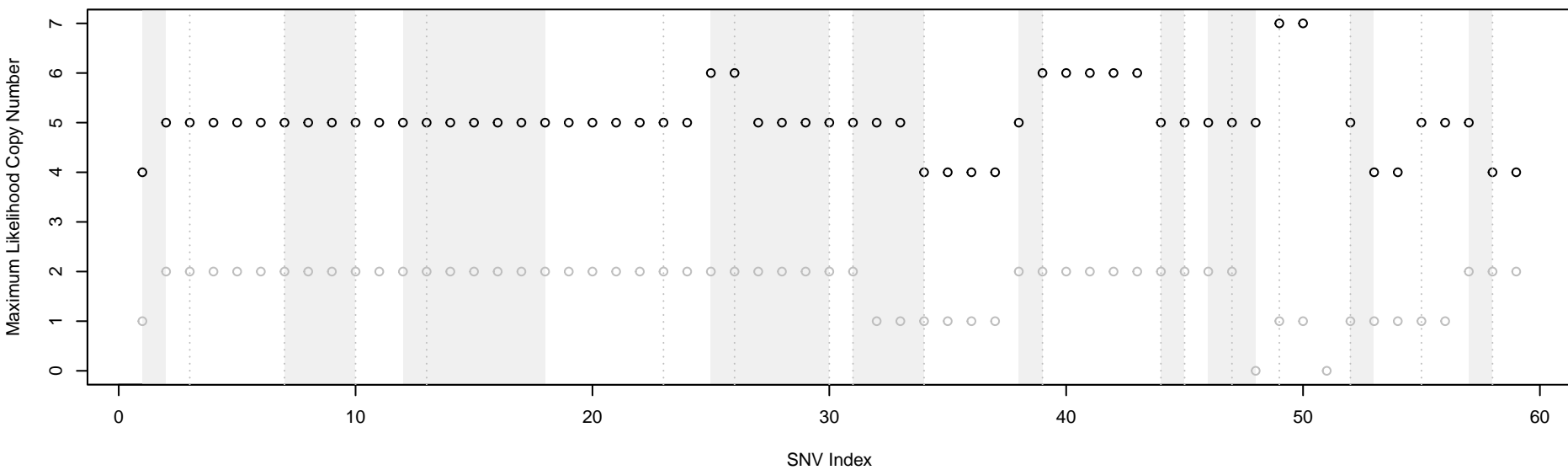
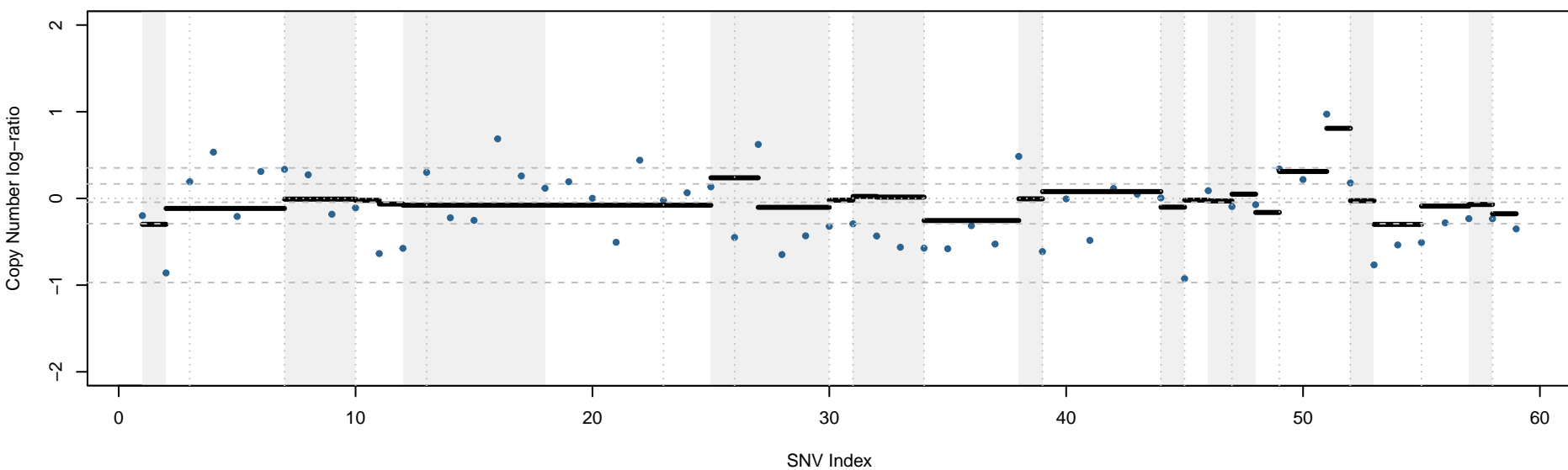
Purity: 0.6 Tumor ploidy: 5.198

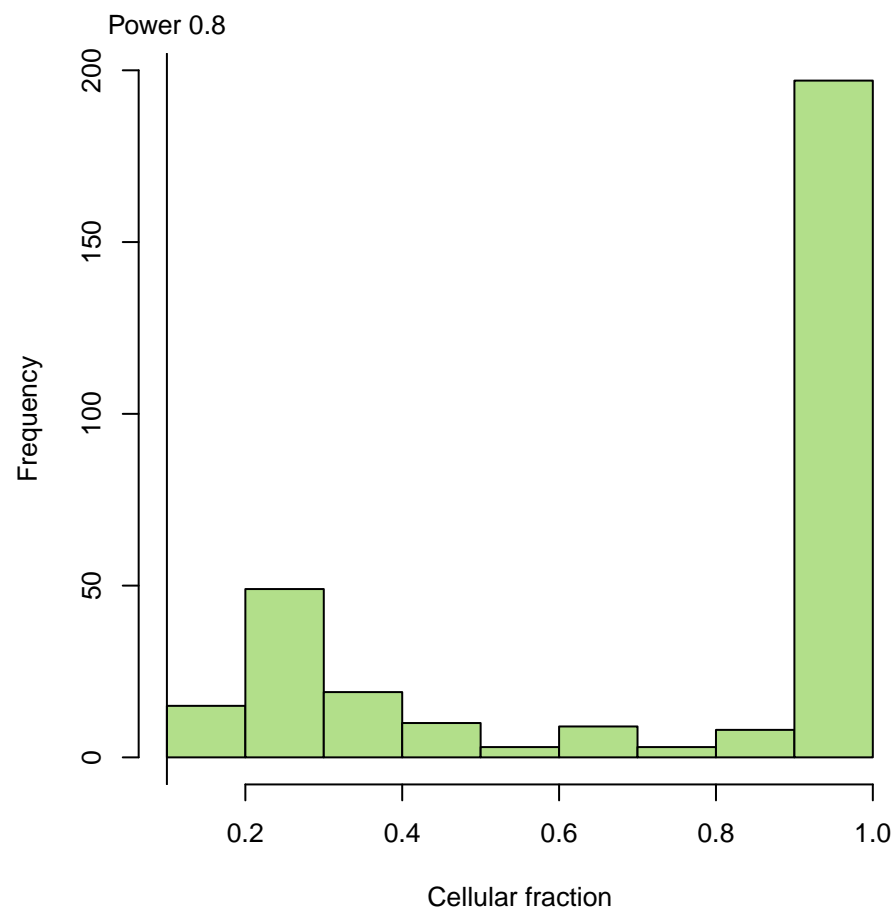
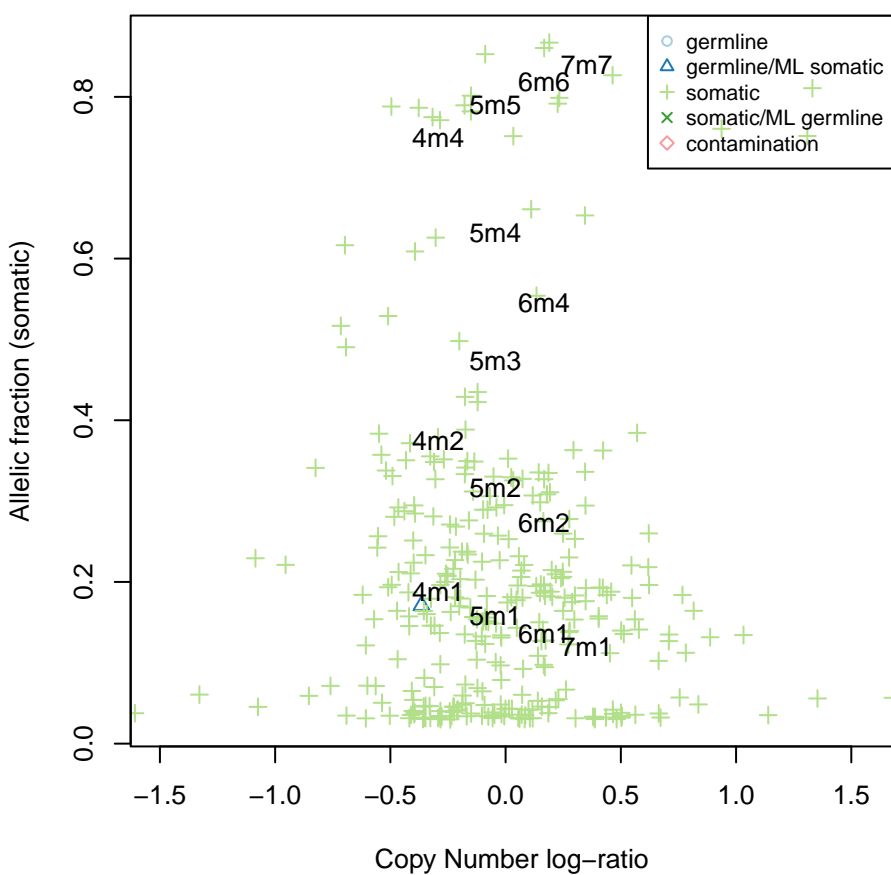
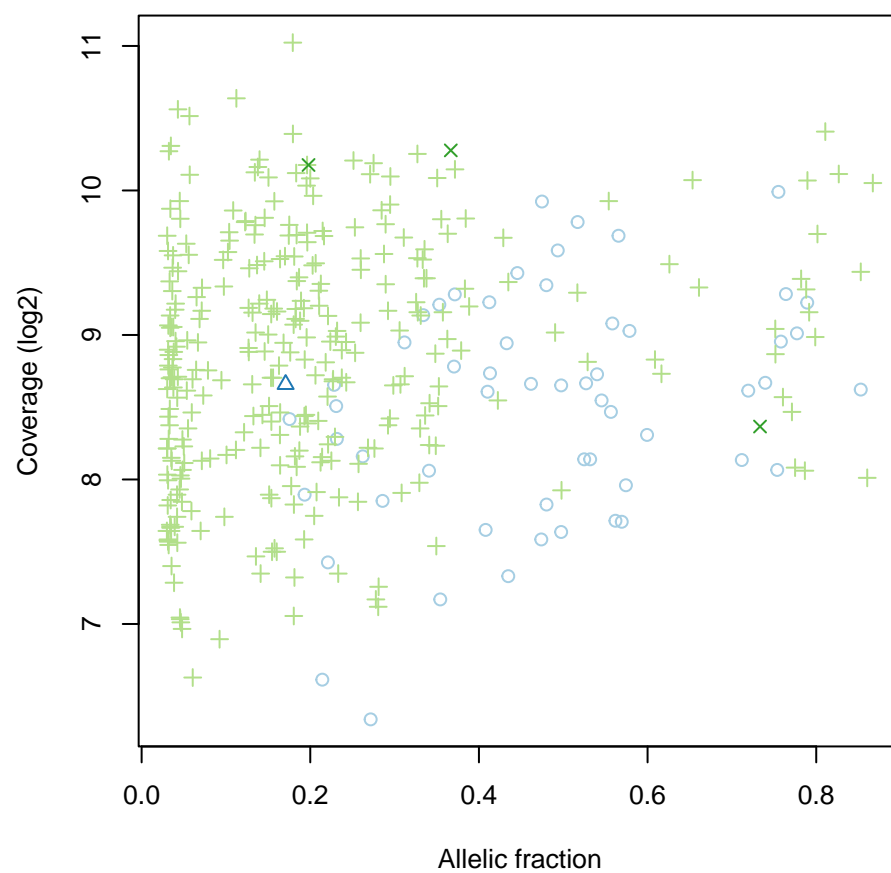
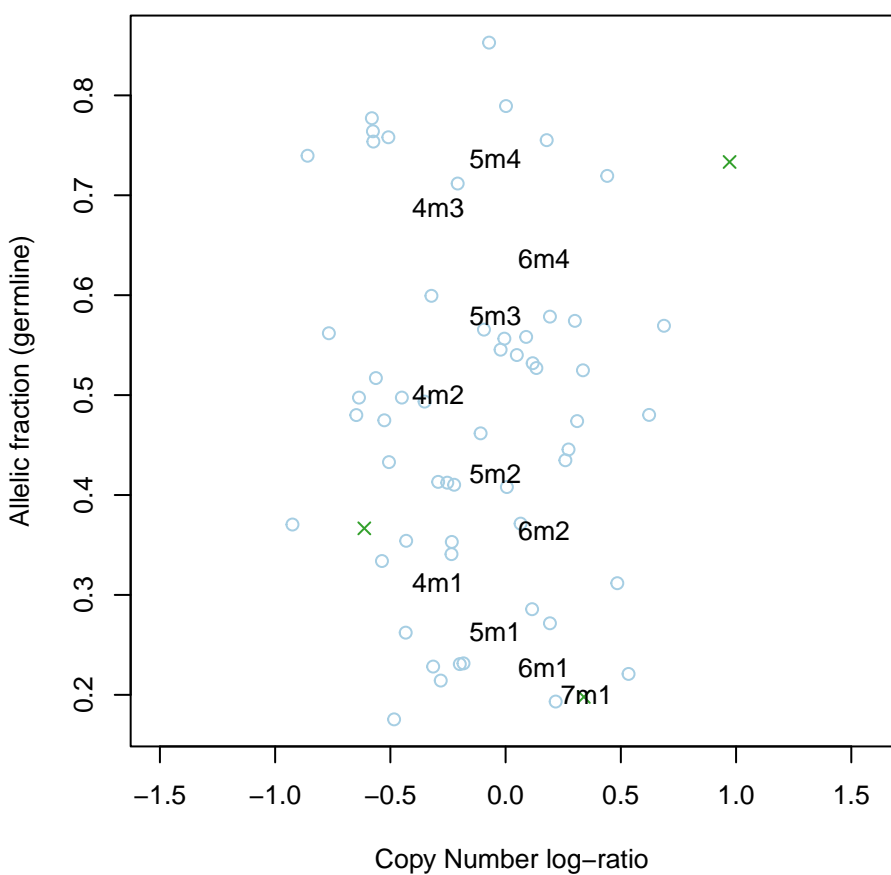


Purity: 0.6 Tumor ploidy: 5.198 SNV log-likelihood: -865.07 GoF: 84.9% Mean coverage: 522;32

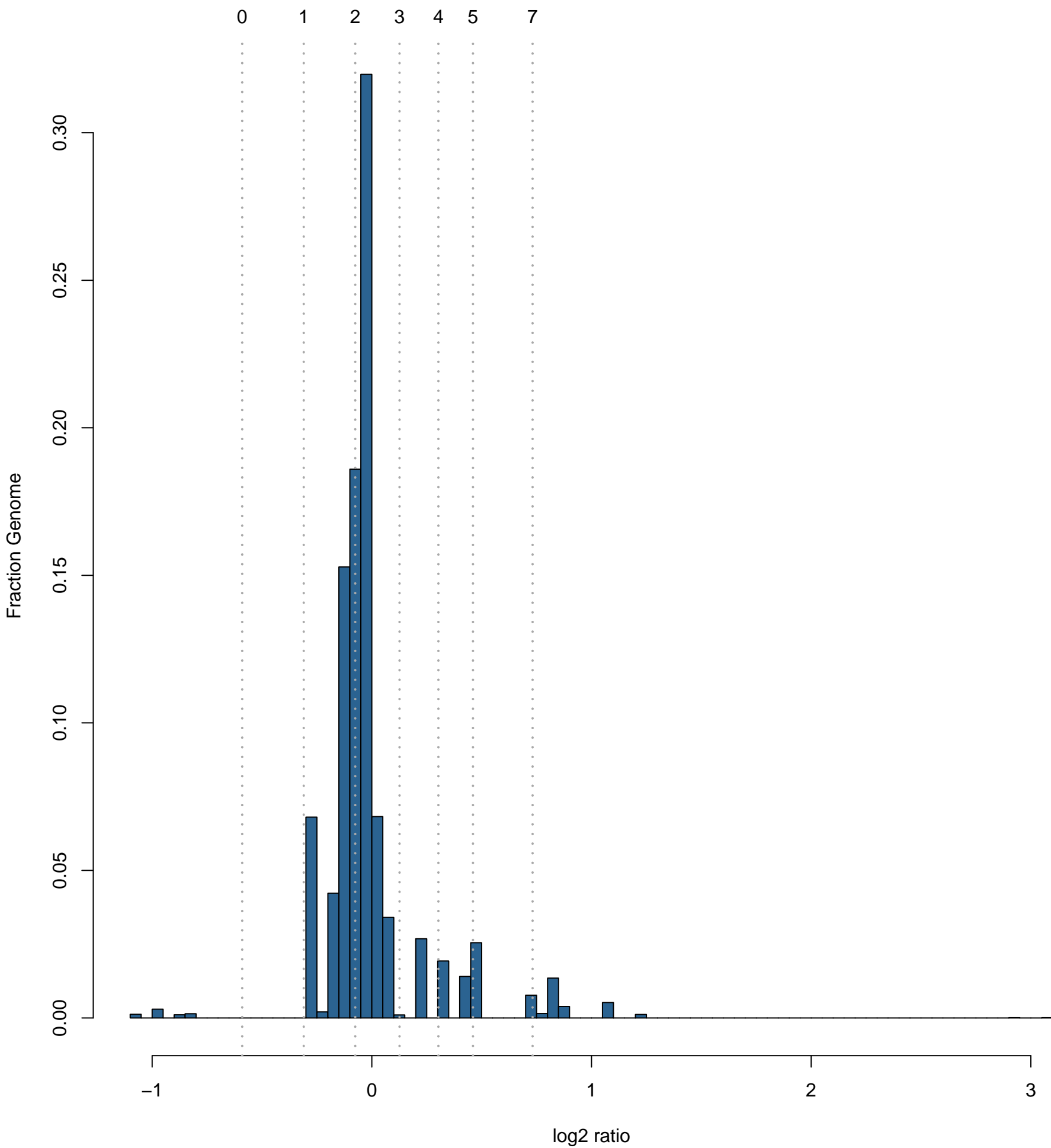


SCNA-fit log-likelihood: -17136.07



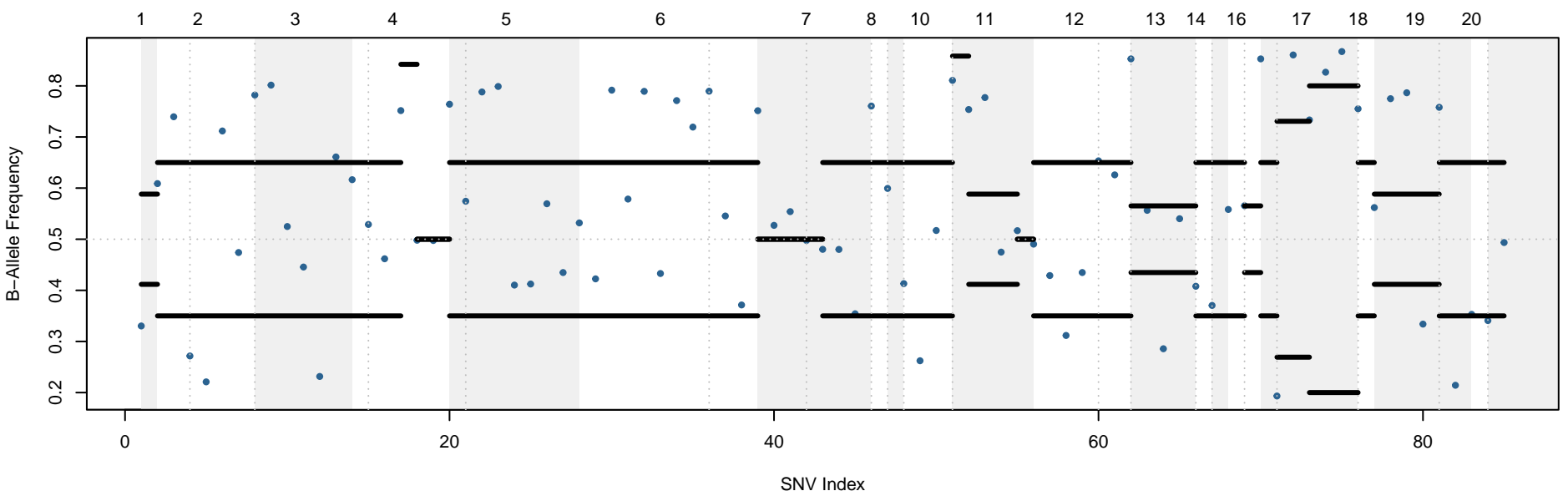


Purity: 0.3 Tumor ploidy: 2.358

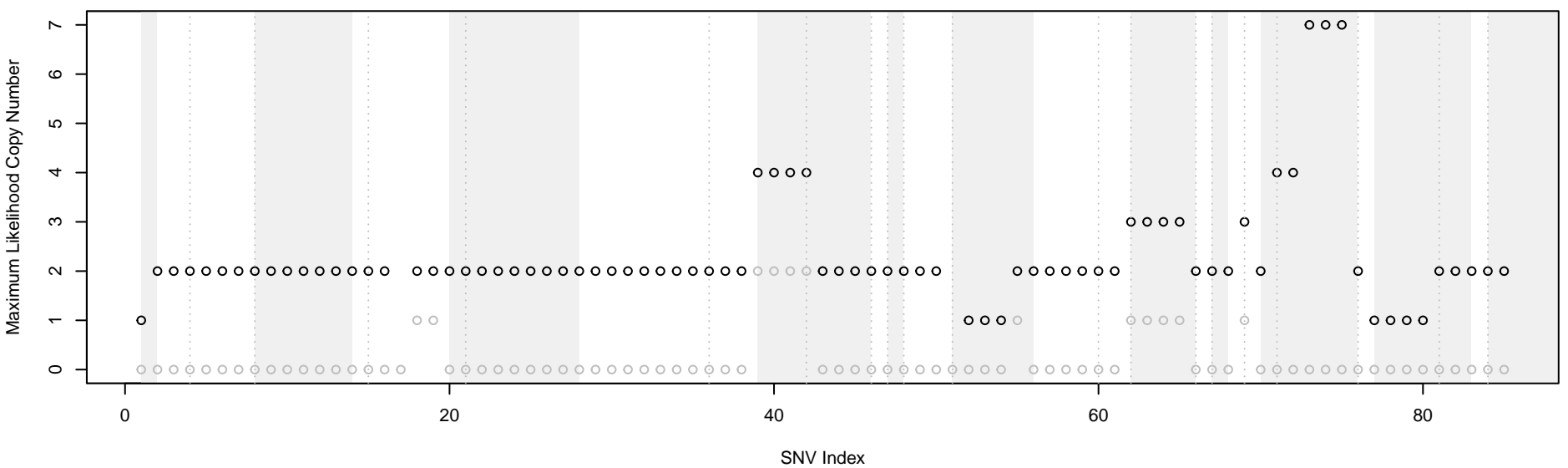
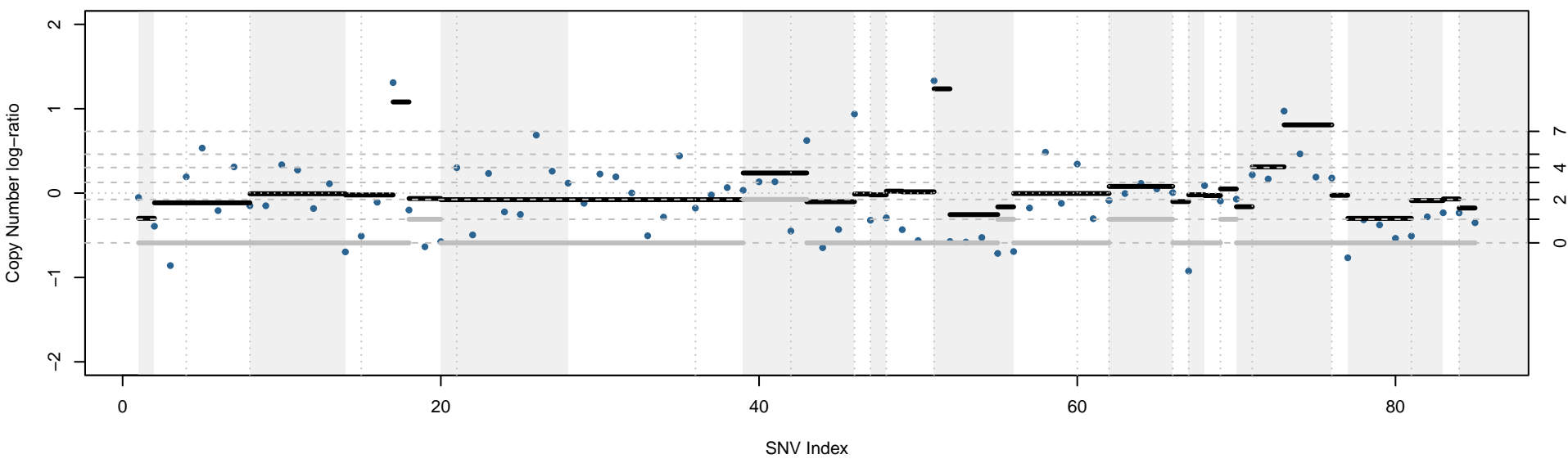




Purity: 0.3 Tumor ploidy: 2.358 SNV log-likelihood: -1071.31 GoF: 83.7% Mean coverage: 522;32

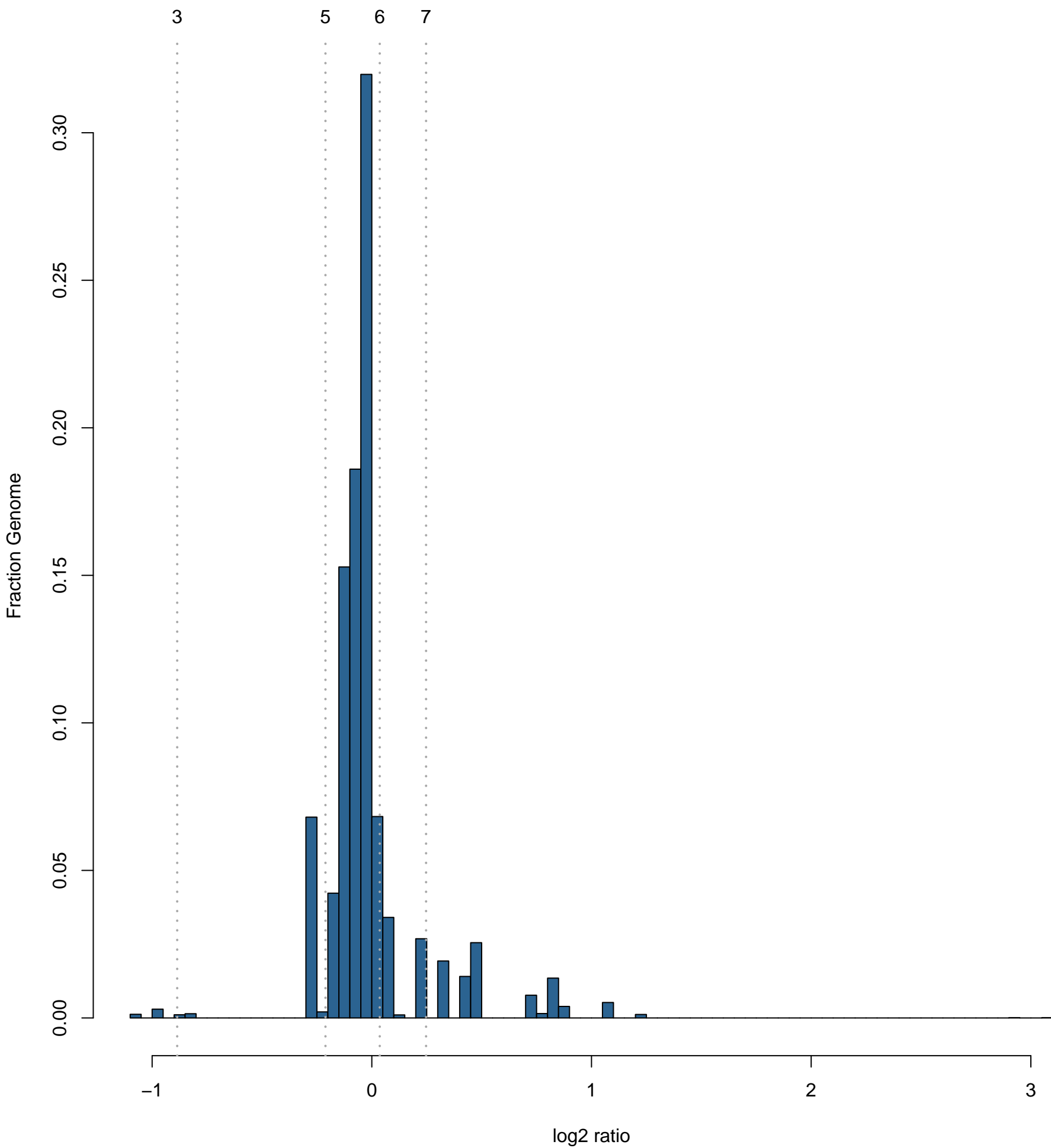


SCNA-fit log-likelihood: -16971.98

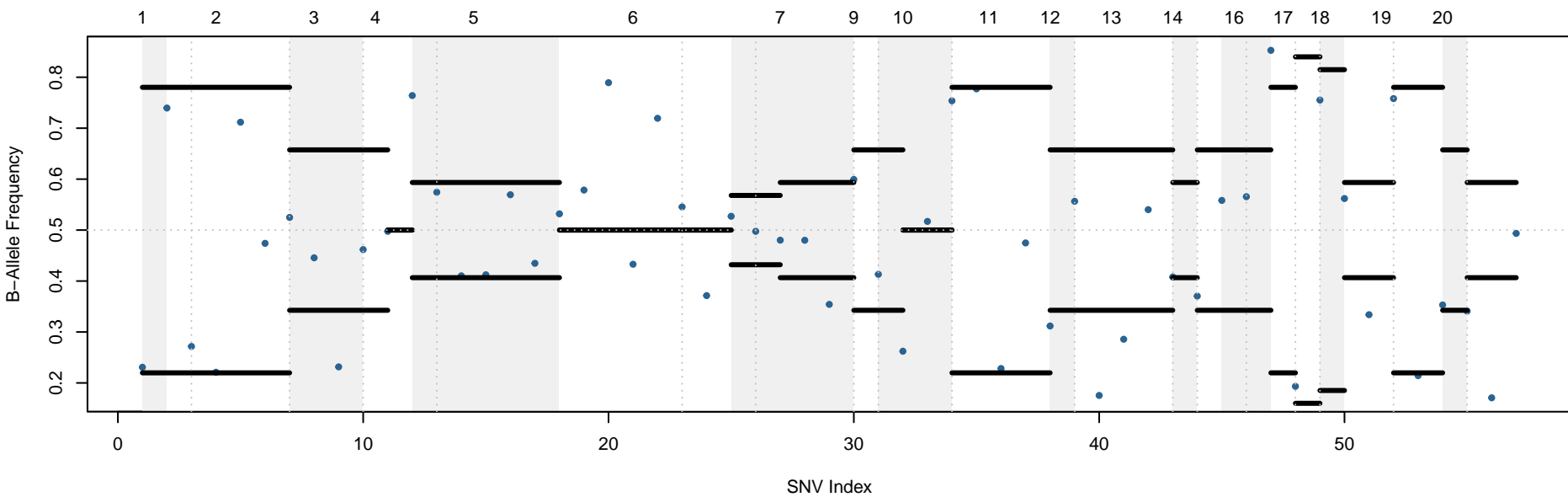




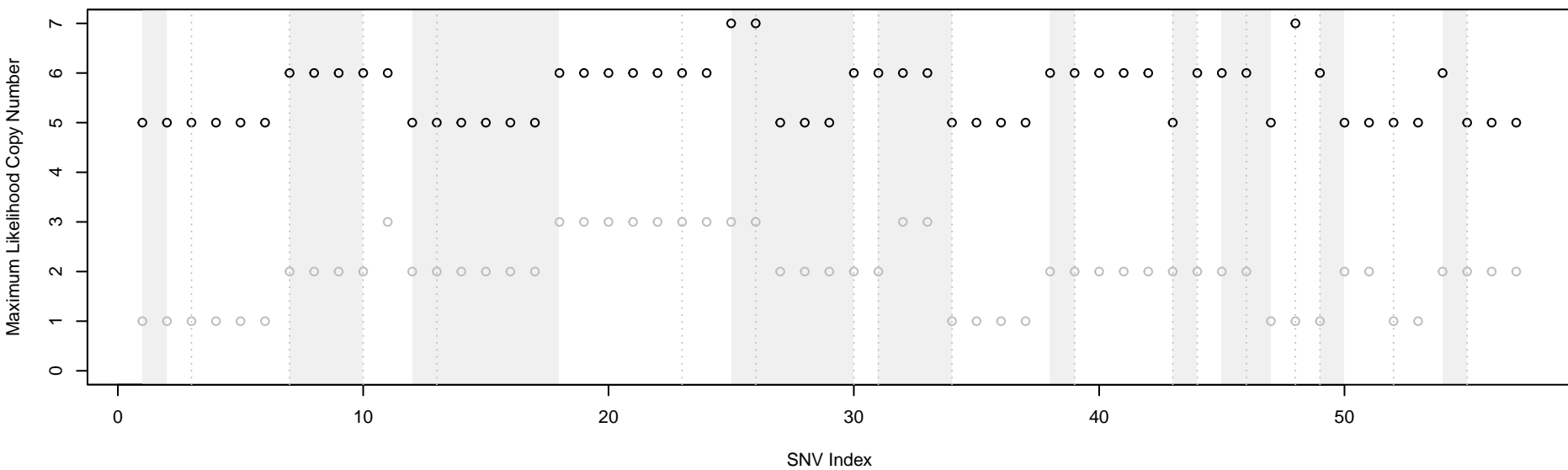
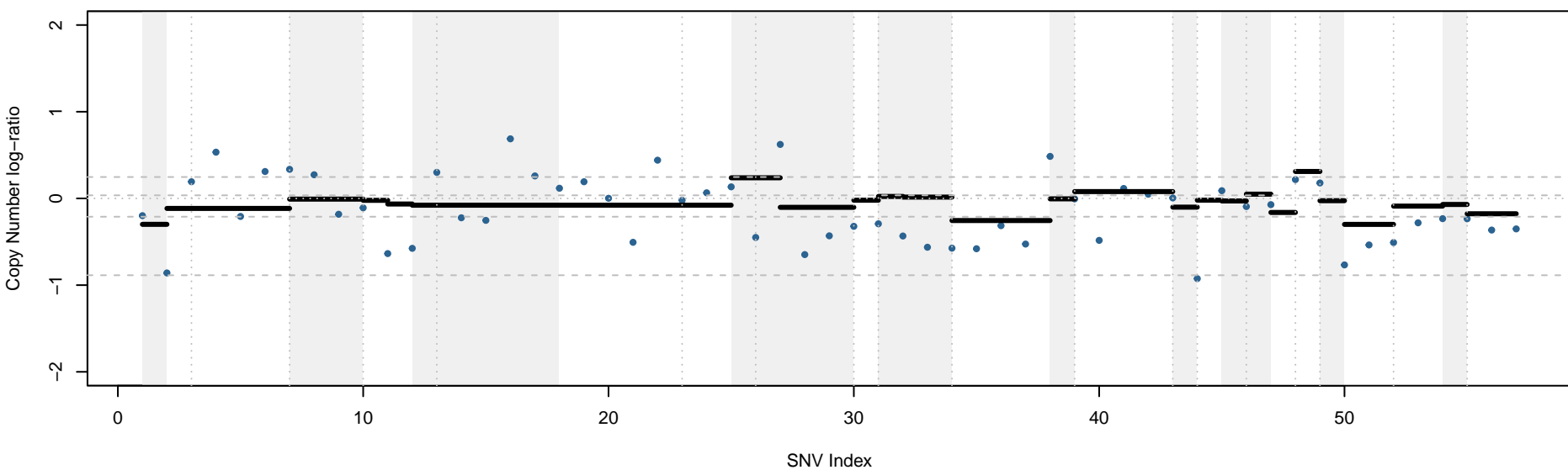
Purity: 0.85 Tumor ploidy: 5.843

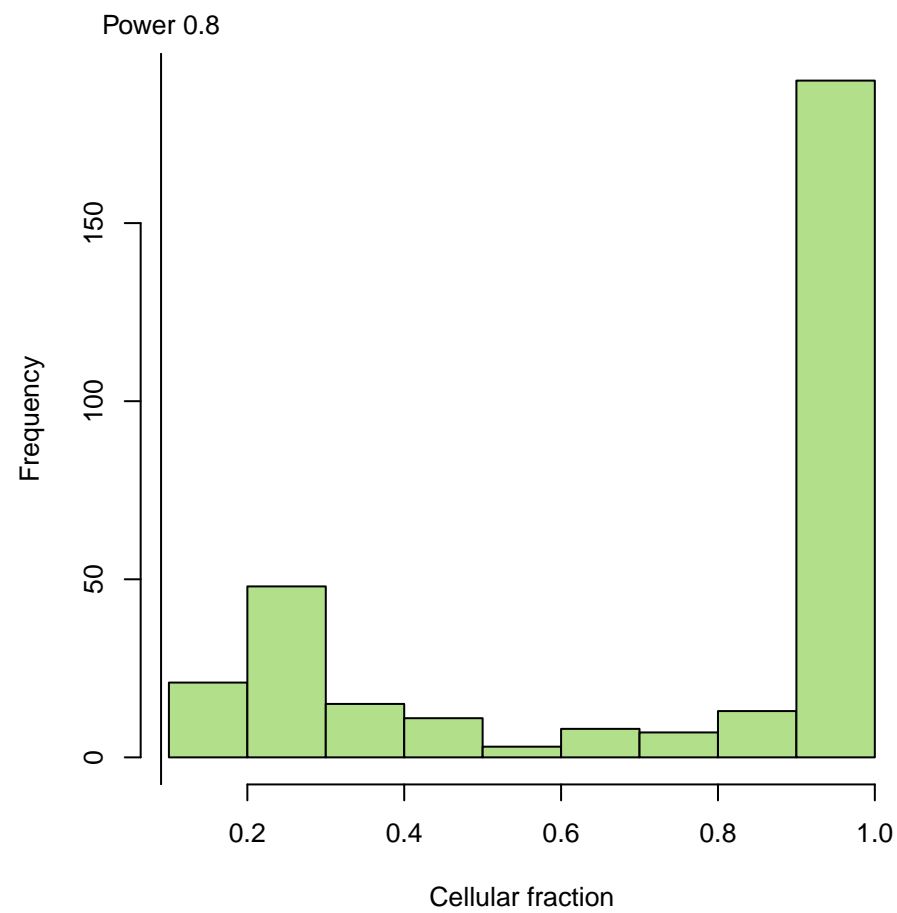
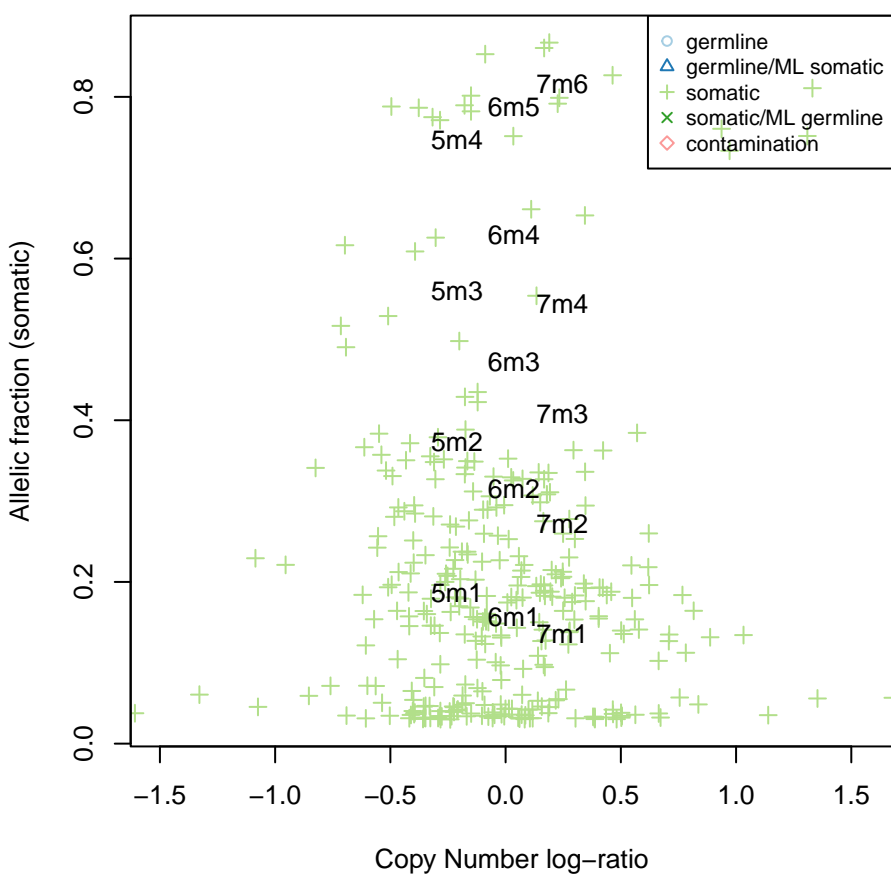
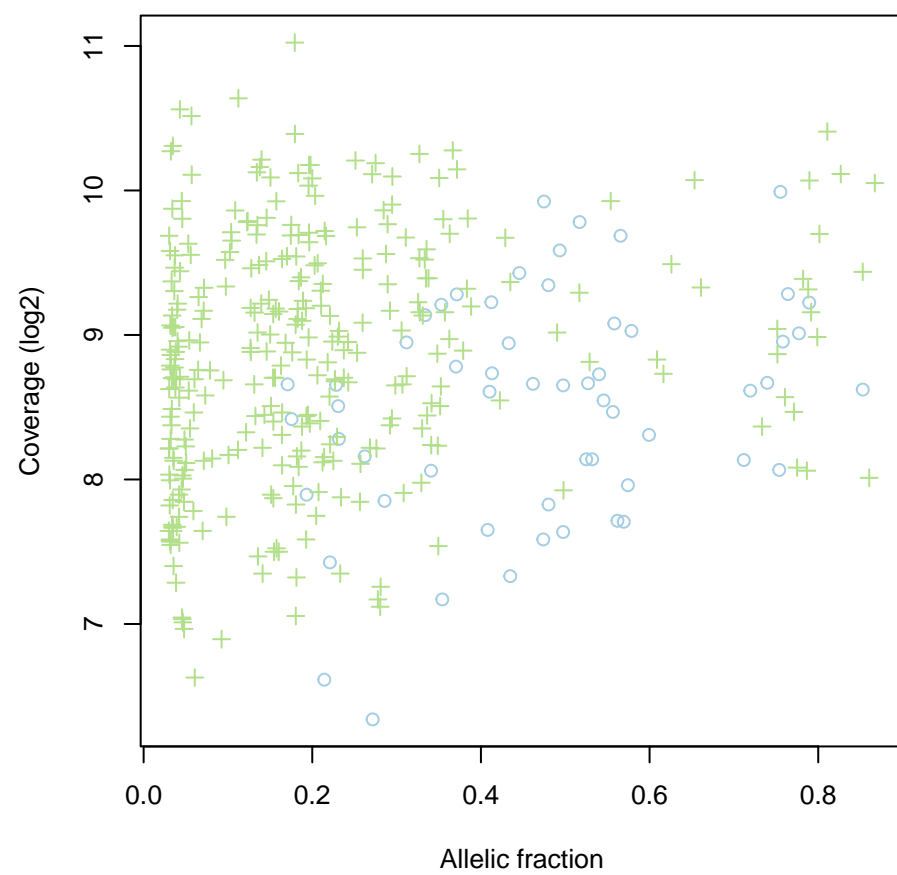
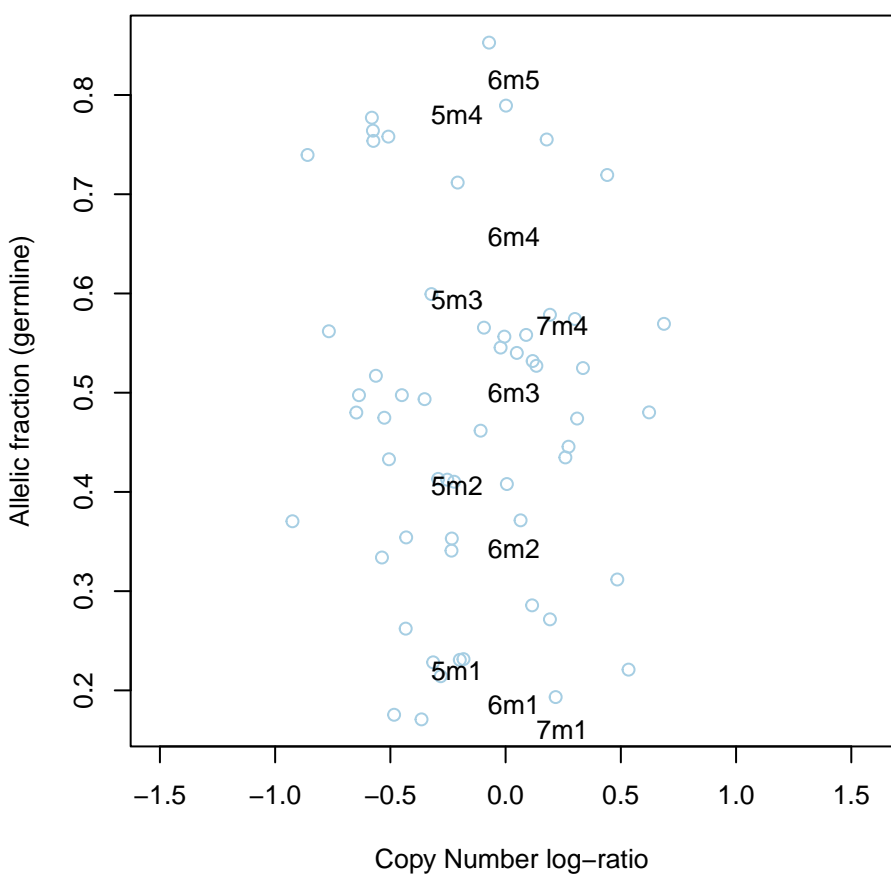


Purity: 0.85 Tumor ploidy: 5.843 SNV log-likelihood: -1003.55 GoF: 83.2% Mean coverage: 522;32

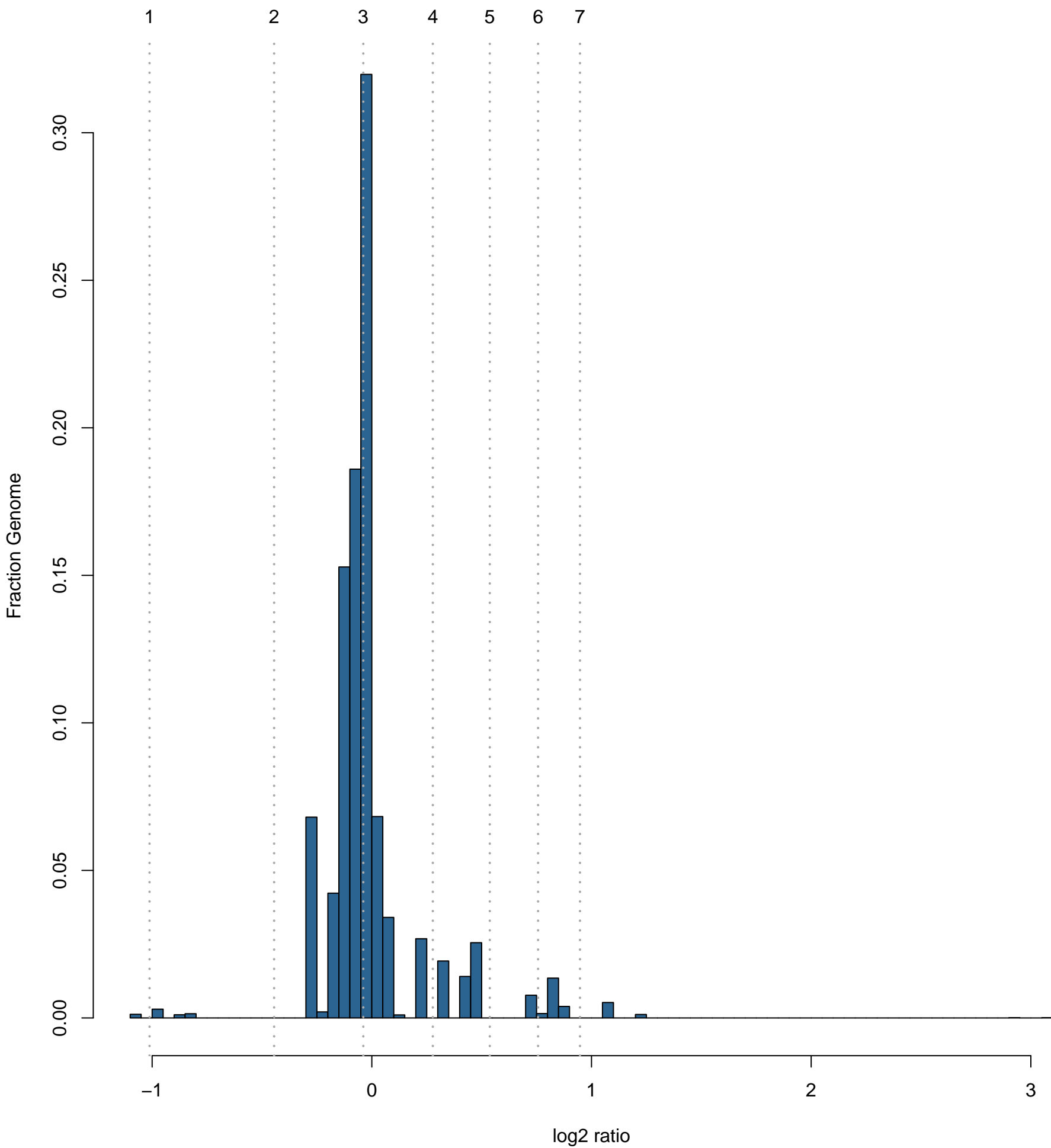


SCNA-fit log-likelihood: -17239.37

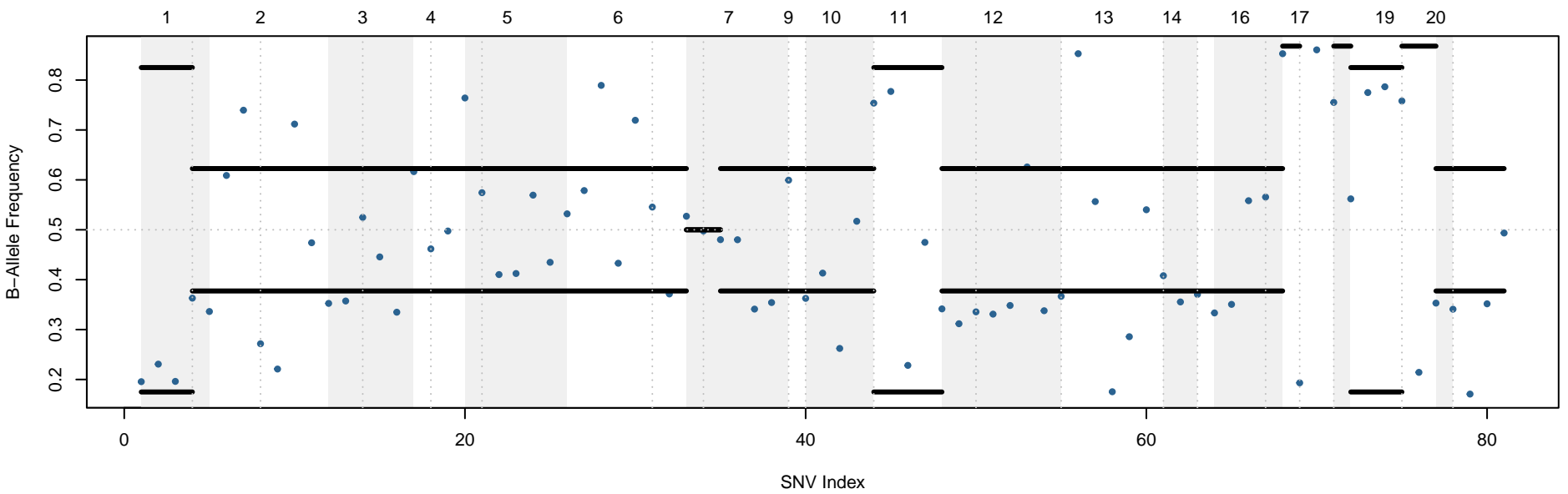




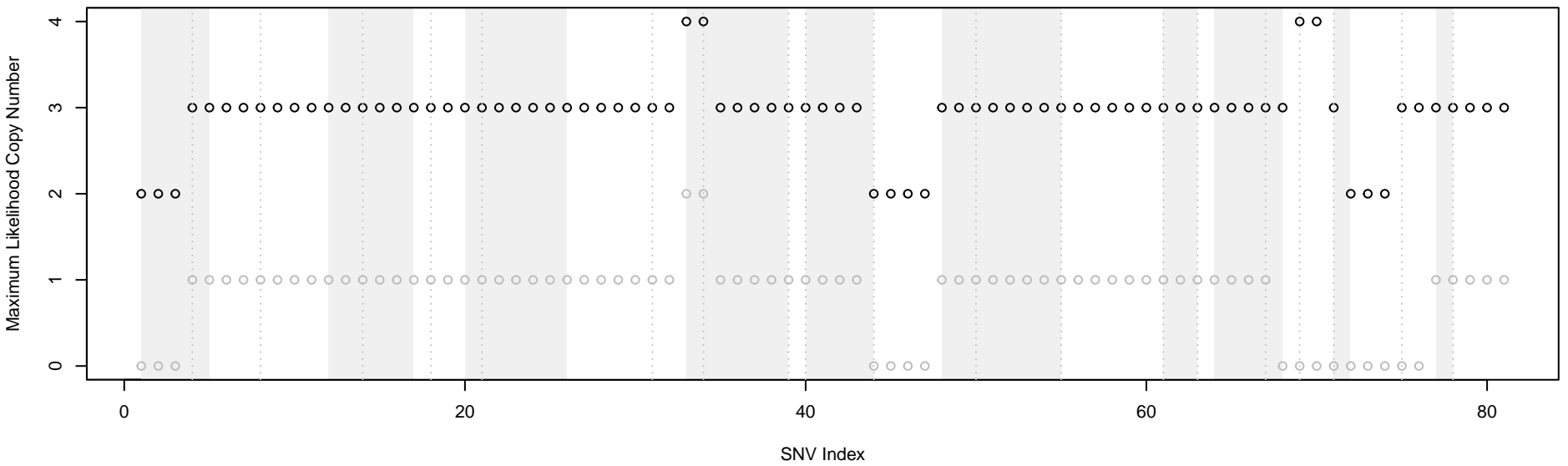
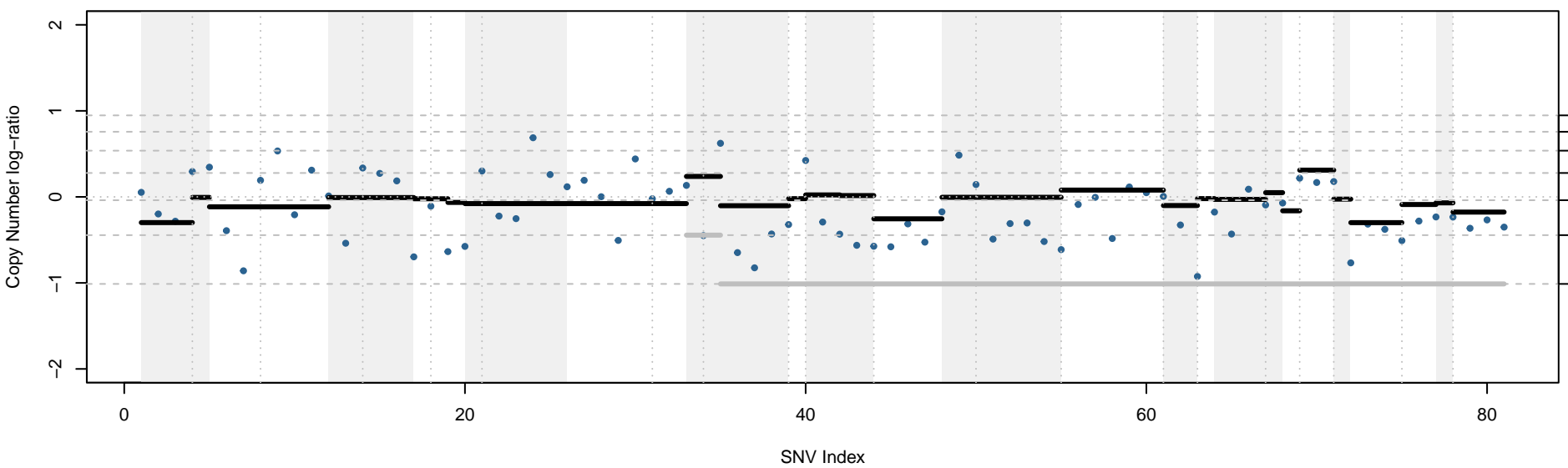
Purity: 0.65 Tumor ploidy: 3.111

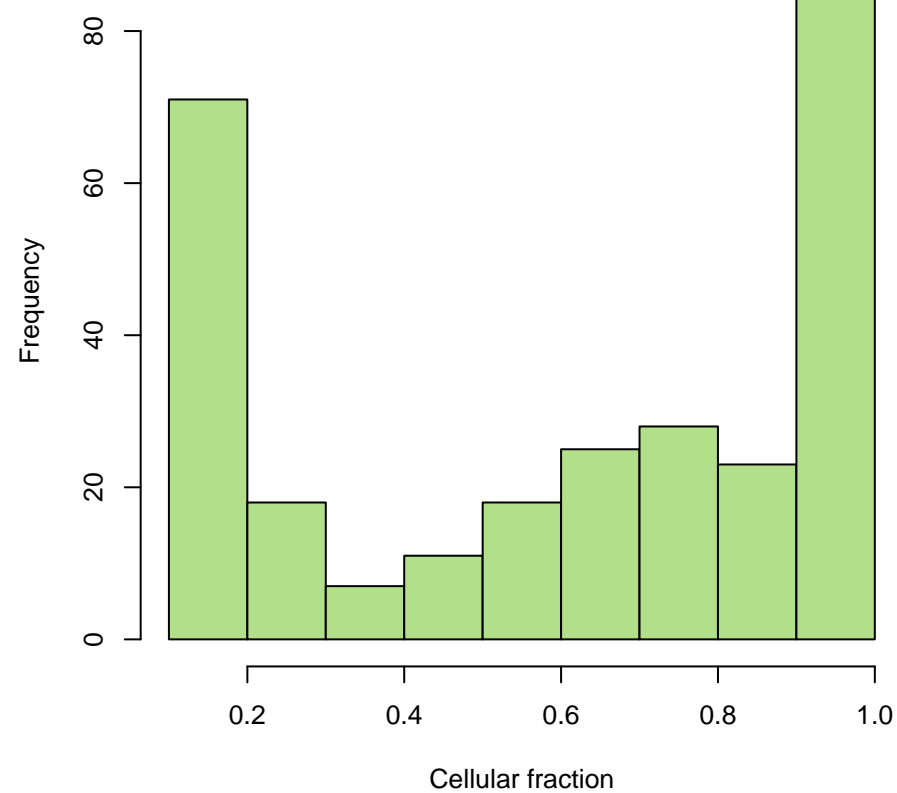
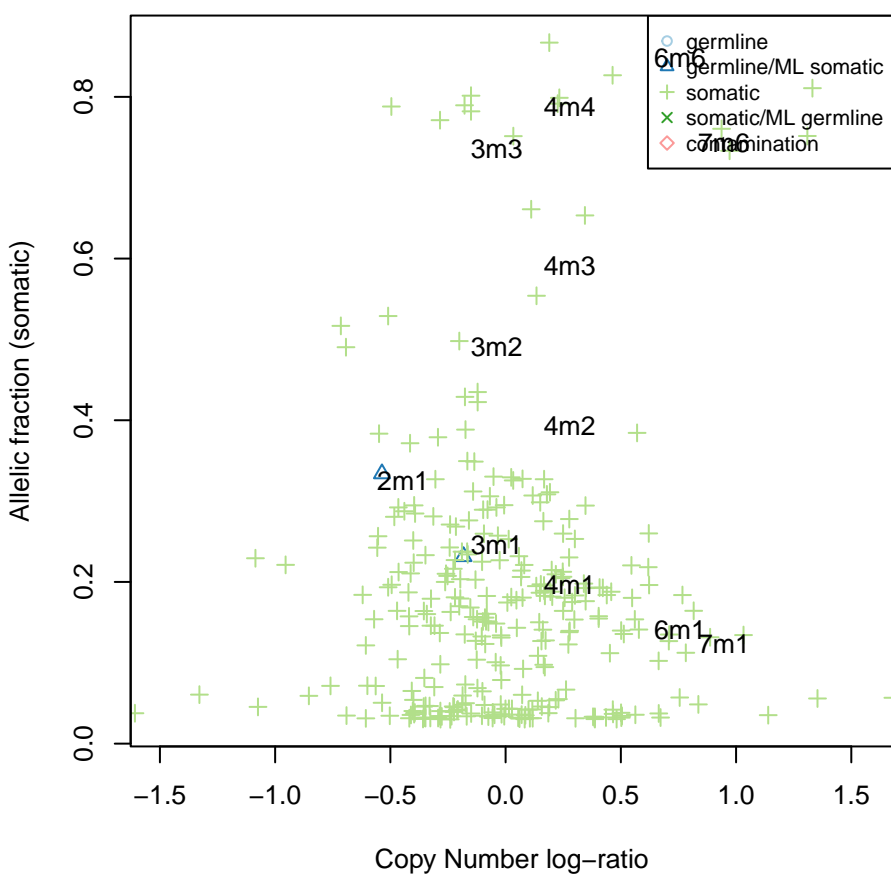
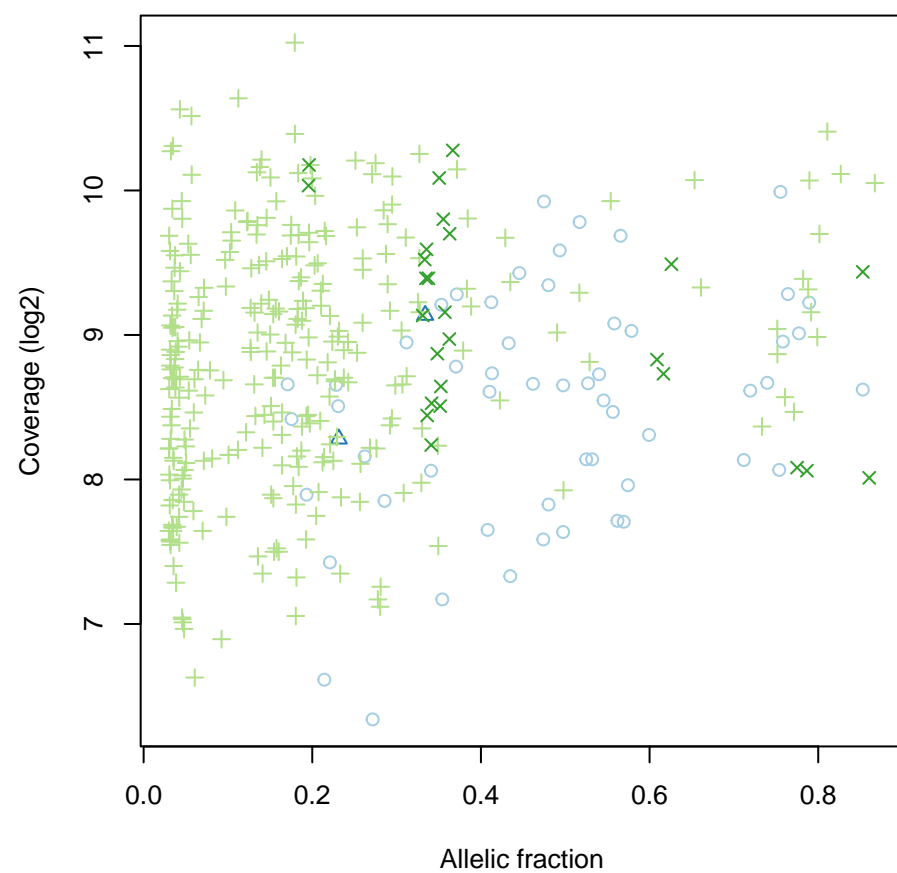
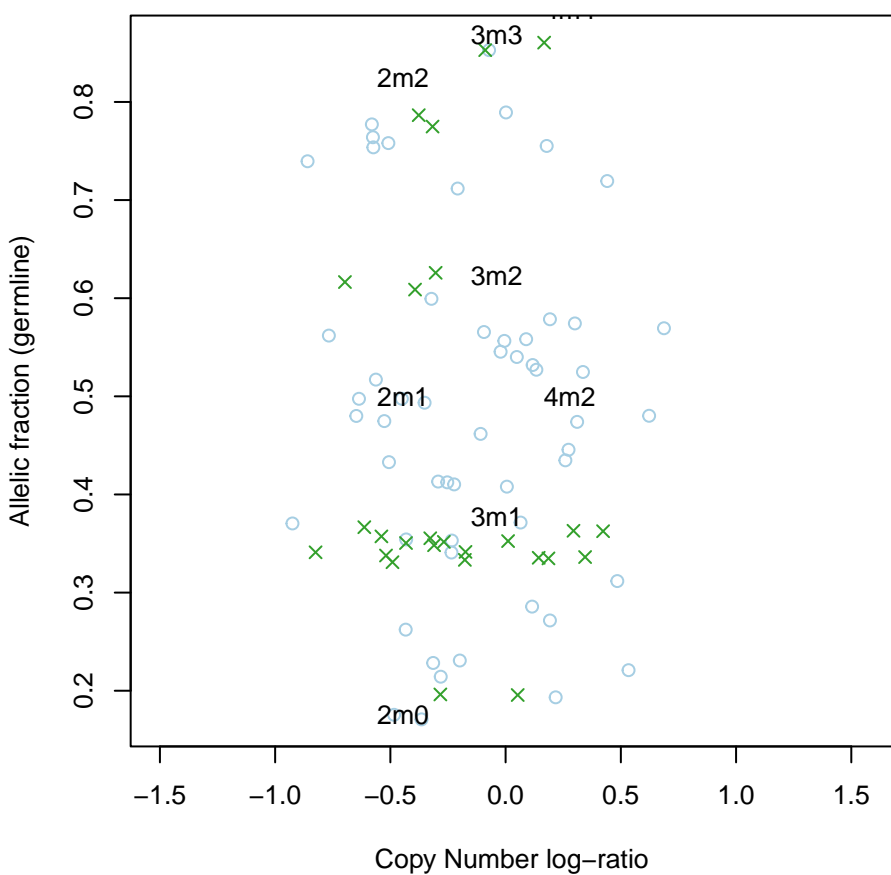


Purity: 0.65 Tumor ploidy: 3.111 SNV log-likelihood: -1177.37 GoF: 64.7% Mean coverage: 522;32



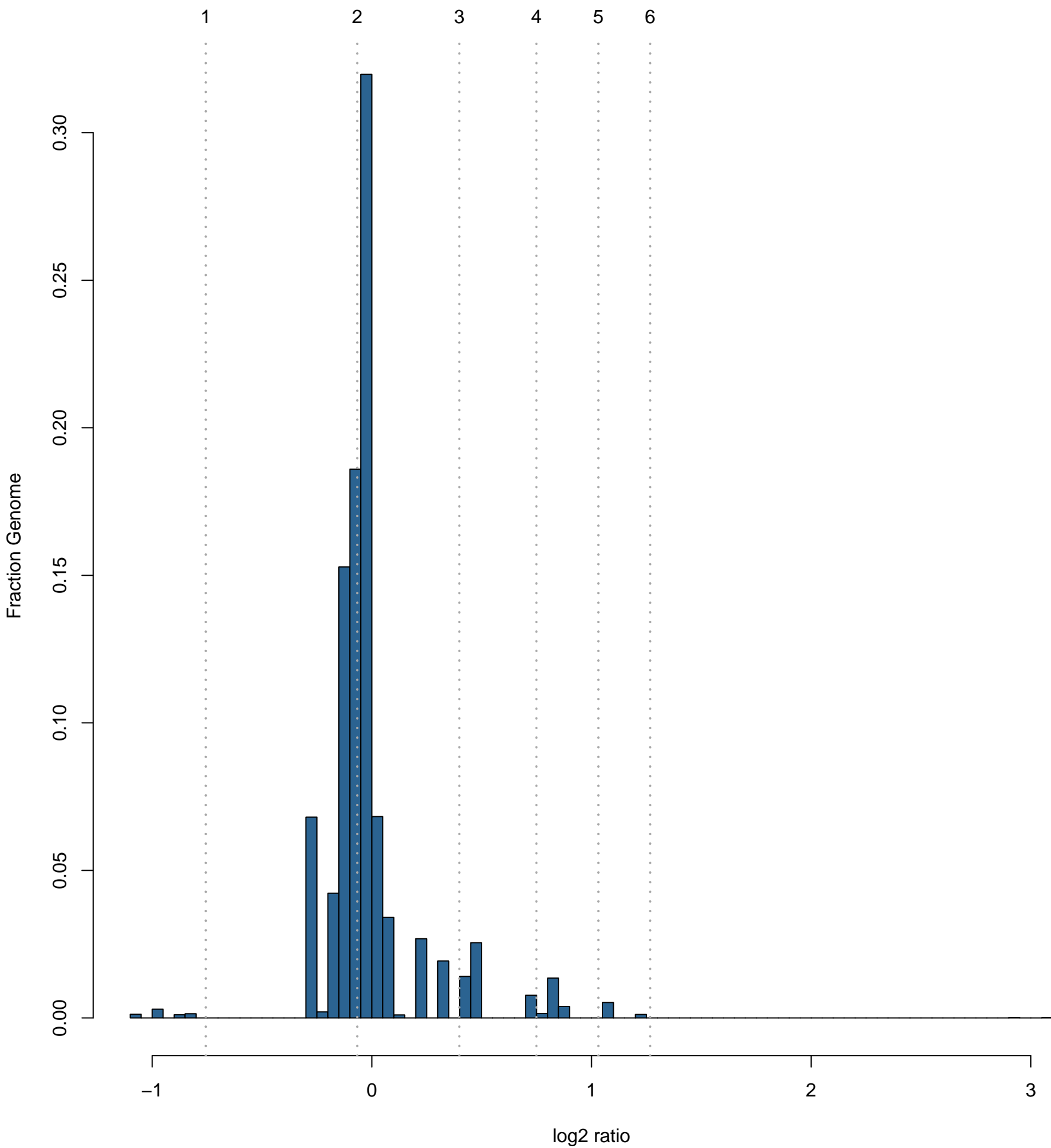
SCNA-fit log-likelihood: -16910.12



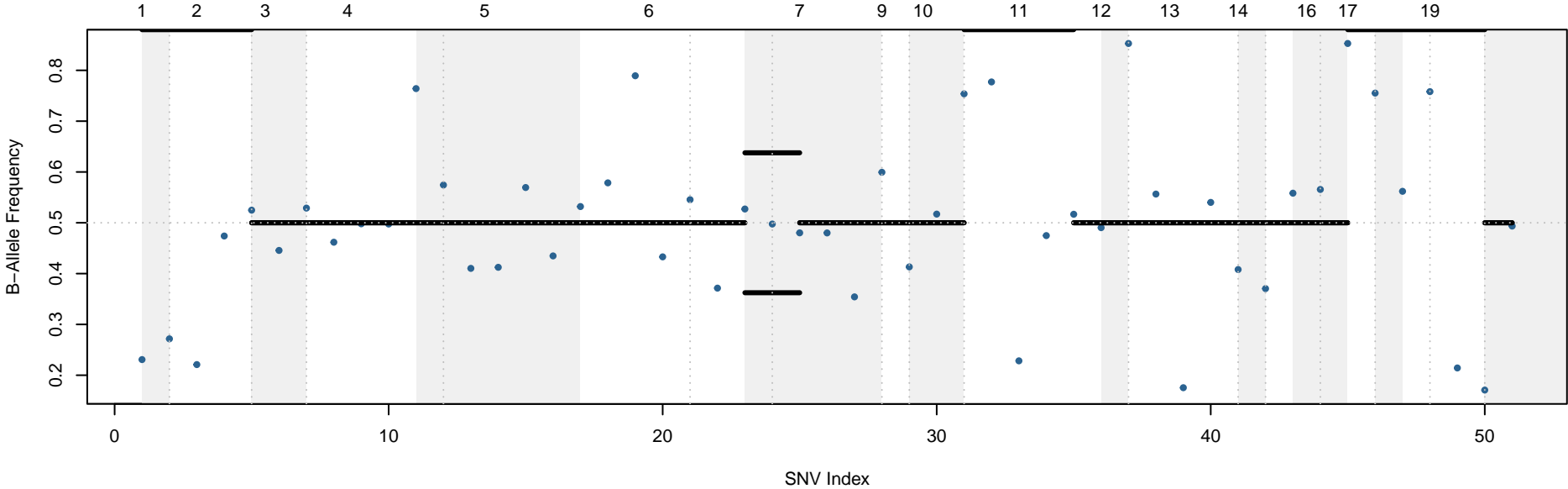




Purity: 0.76 Tumor ploidy: 2.124



Purity: 0.76 Tumor ploidy: 2.124 SNV log-likelihood: -1914.06 GoF: 0% Mean coverage: 522;32



SCNA-fit log-likelihood: -16979.54

