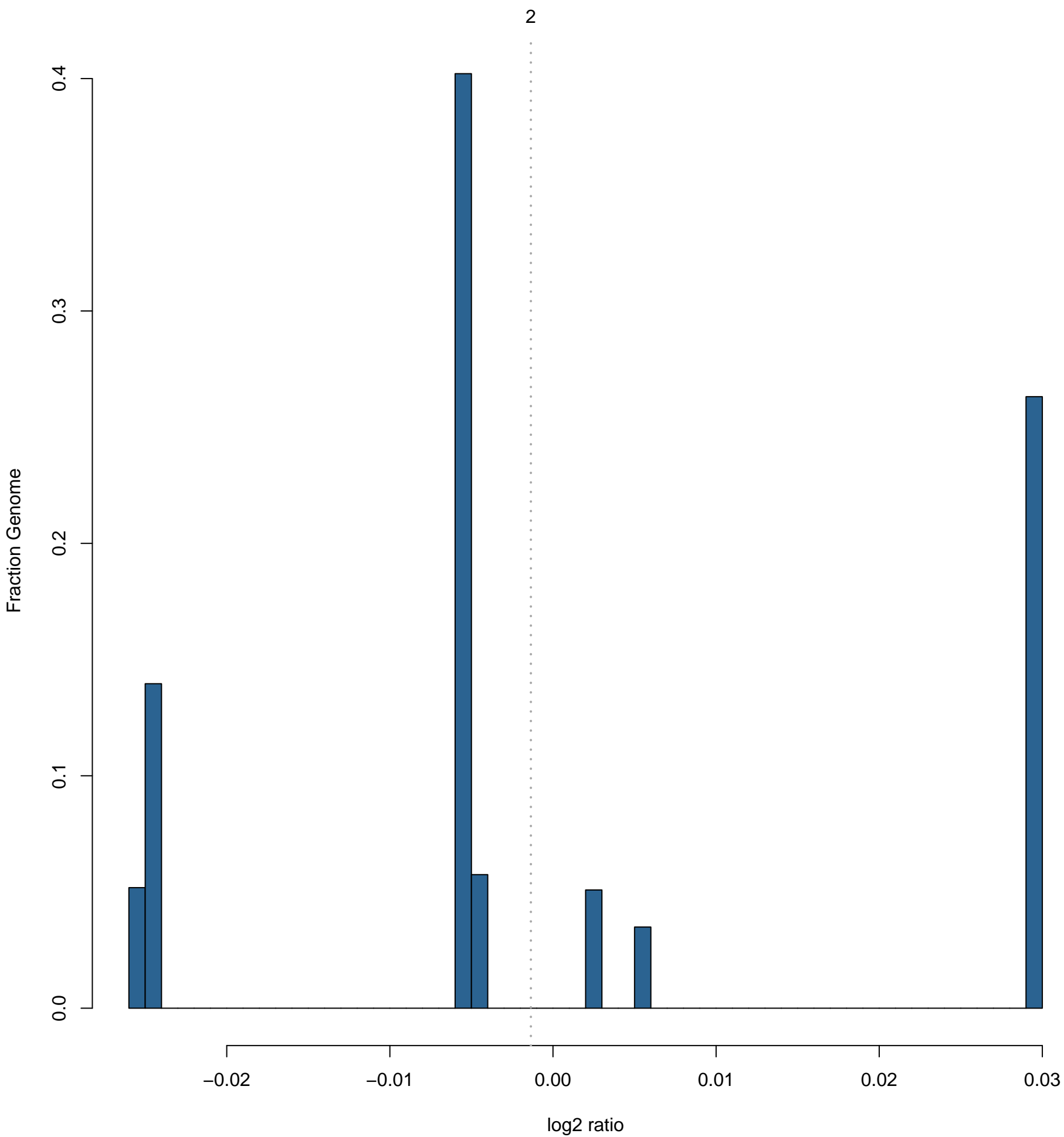
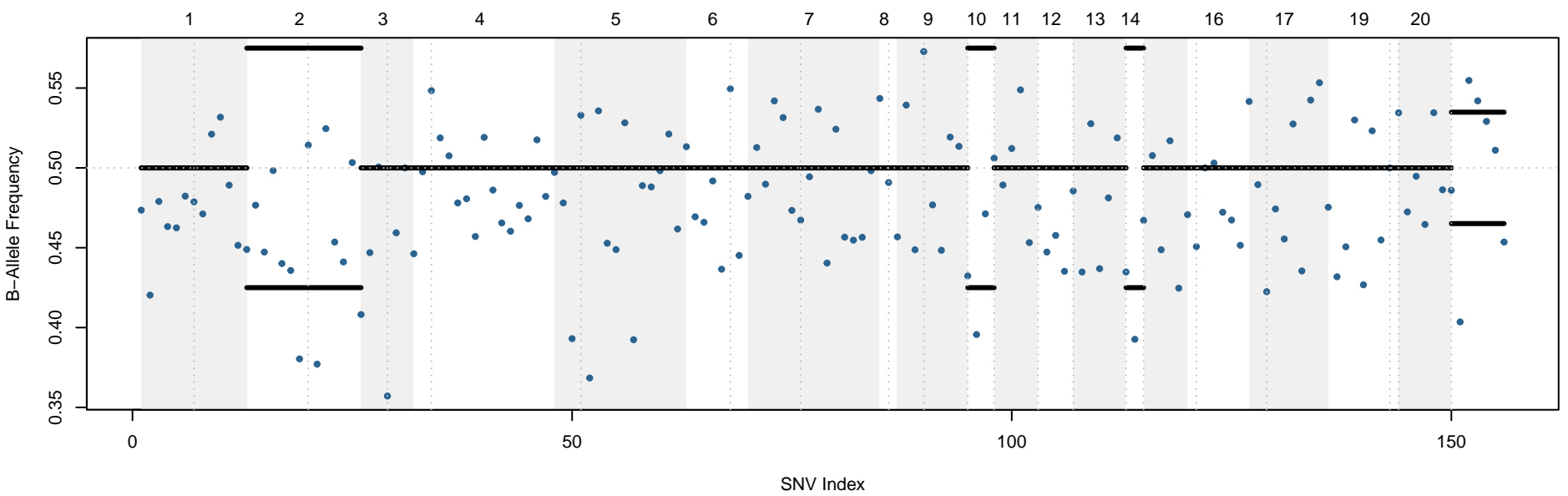


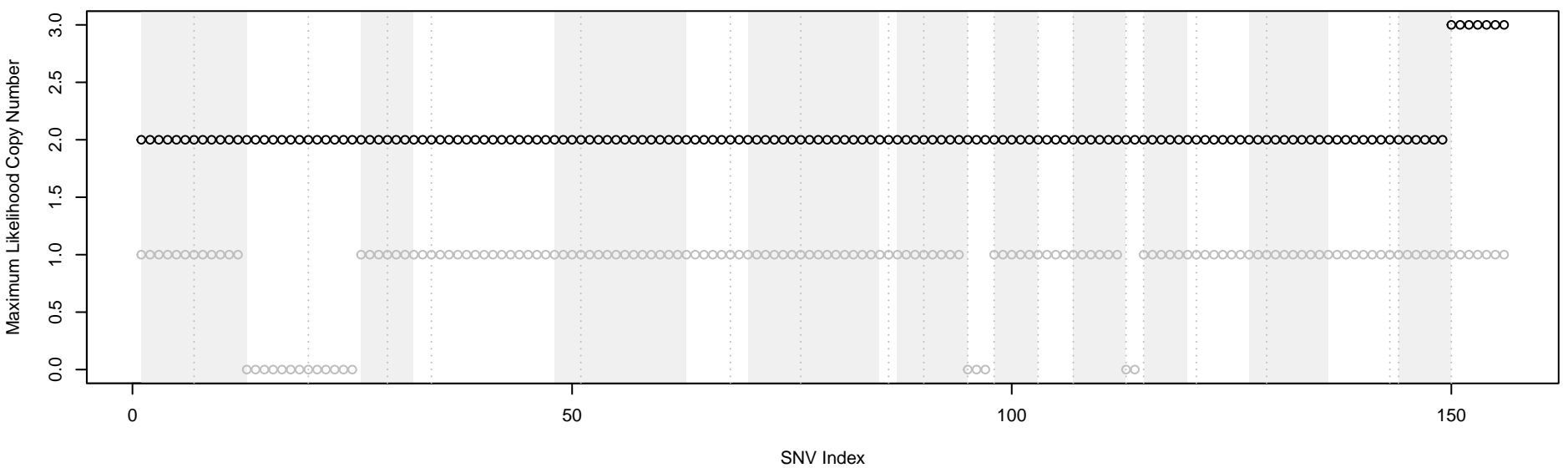
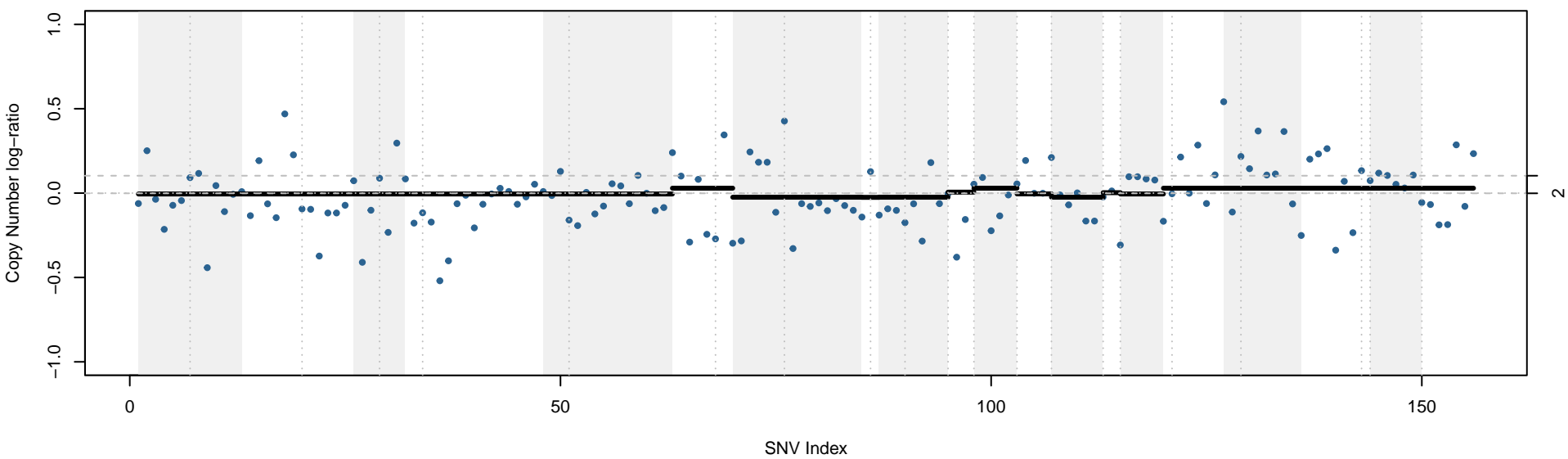
Purity: 0.15 Tumor ploidy: 2.013

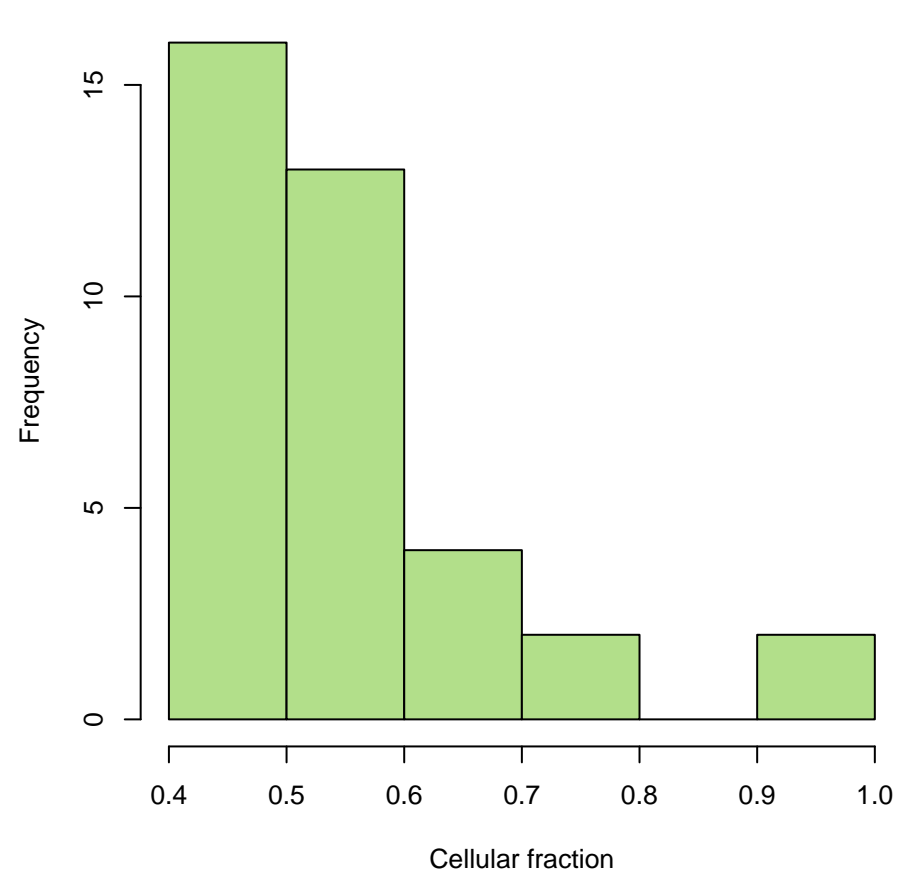
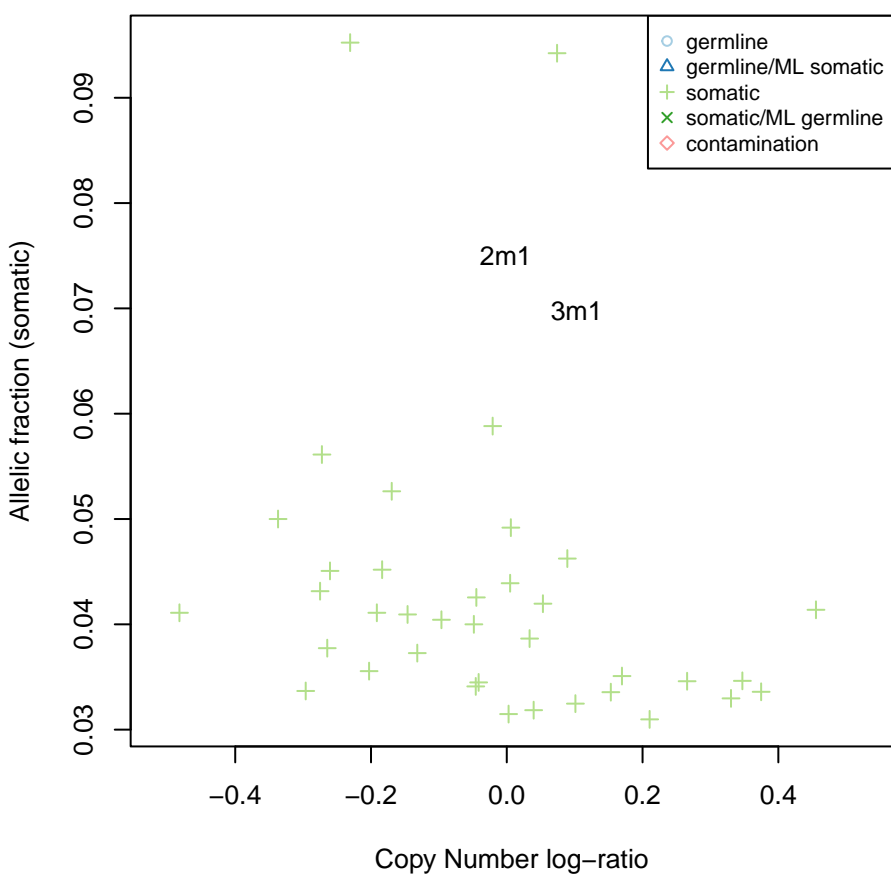
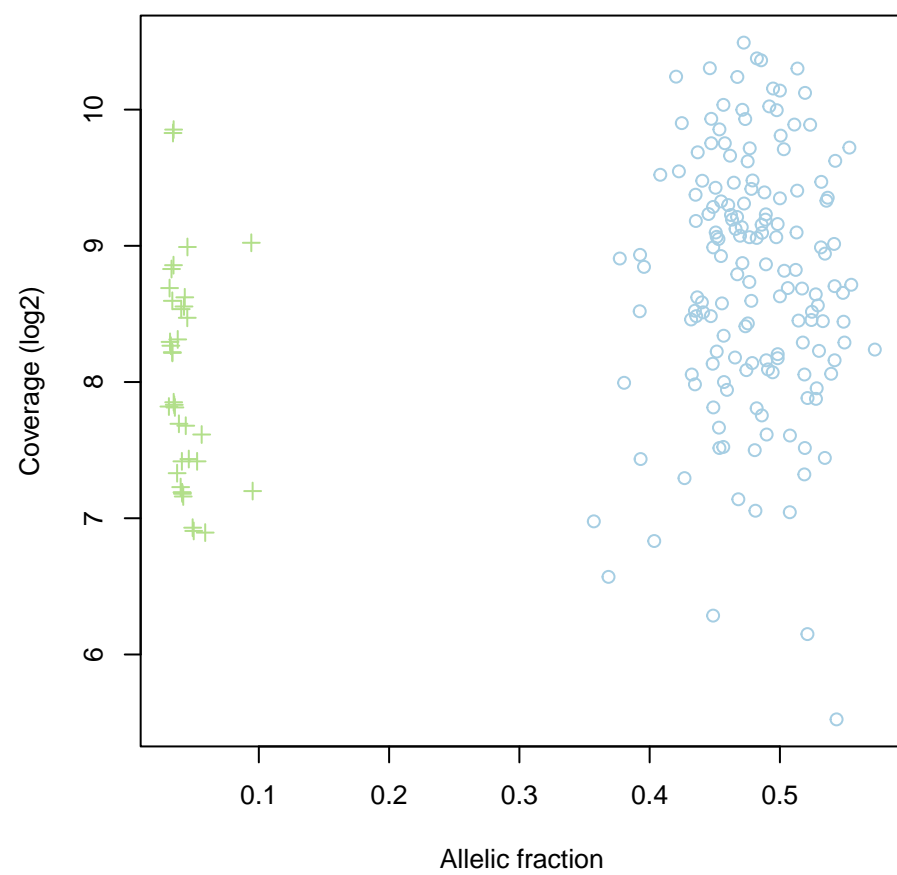
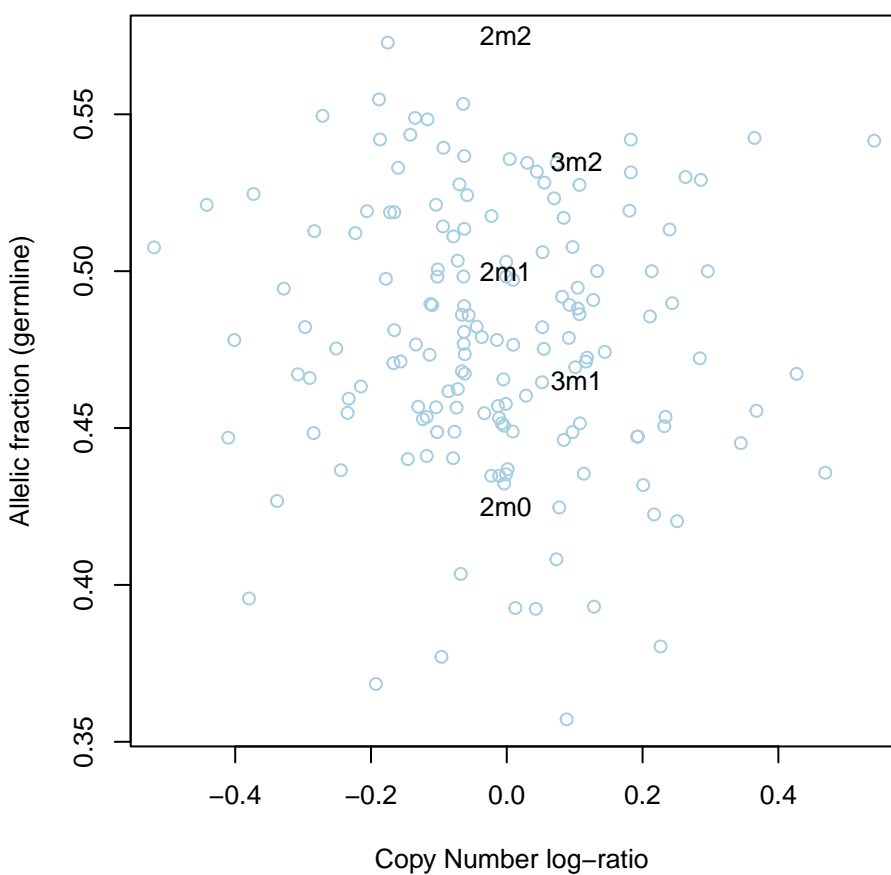


Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477

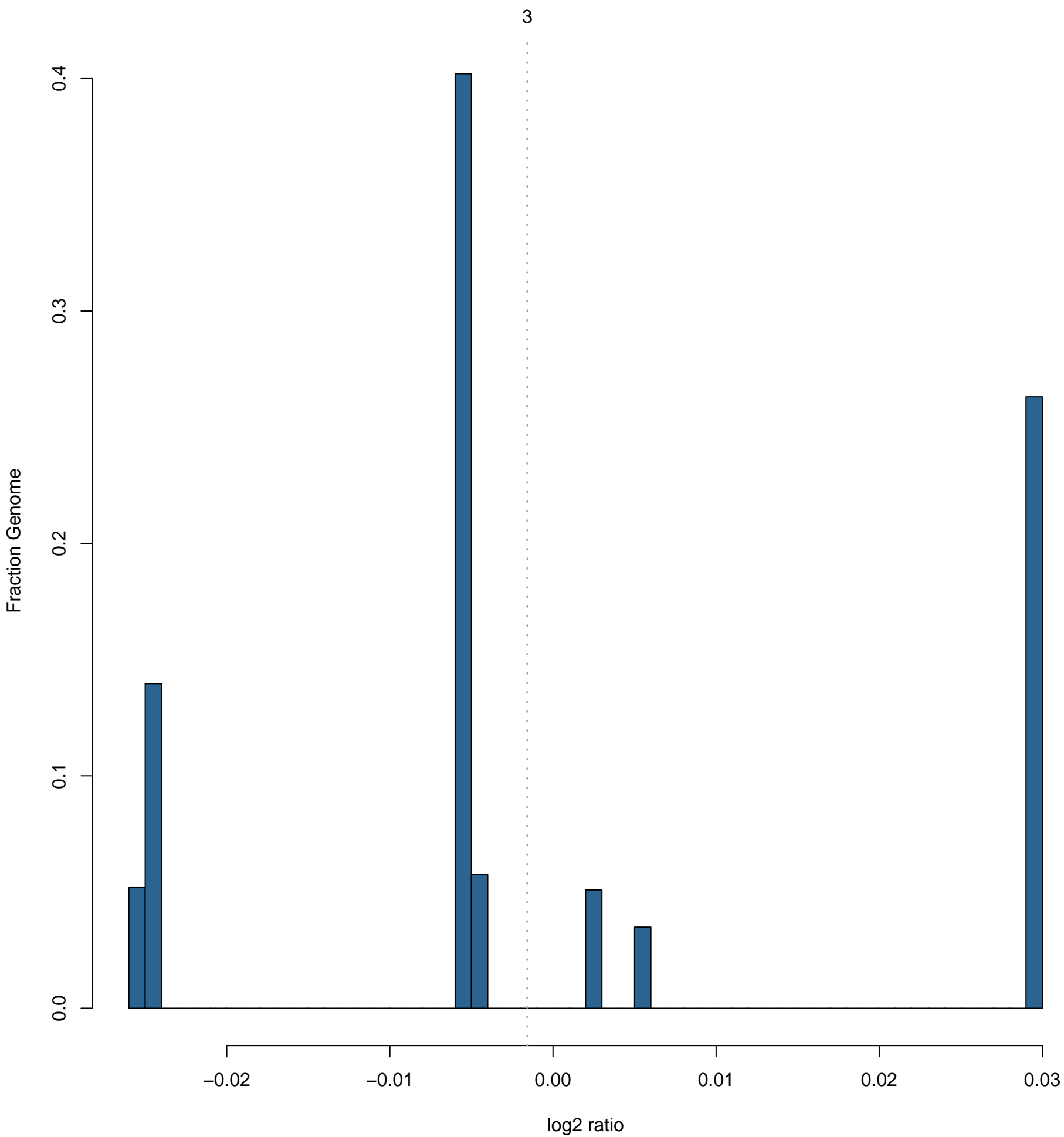


SCNA-fit log-likelihood: -7173.18

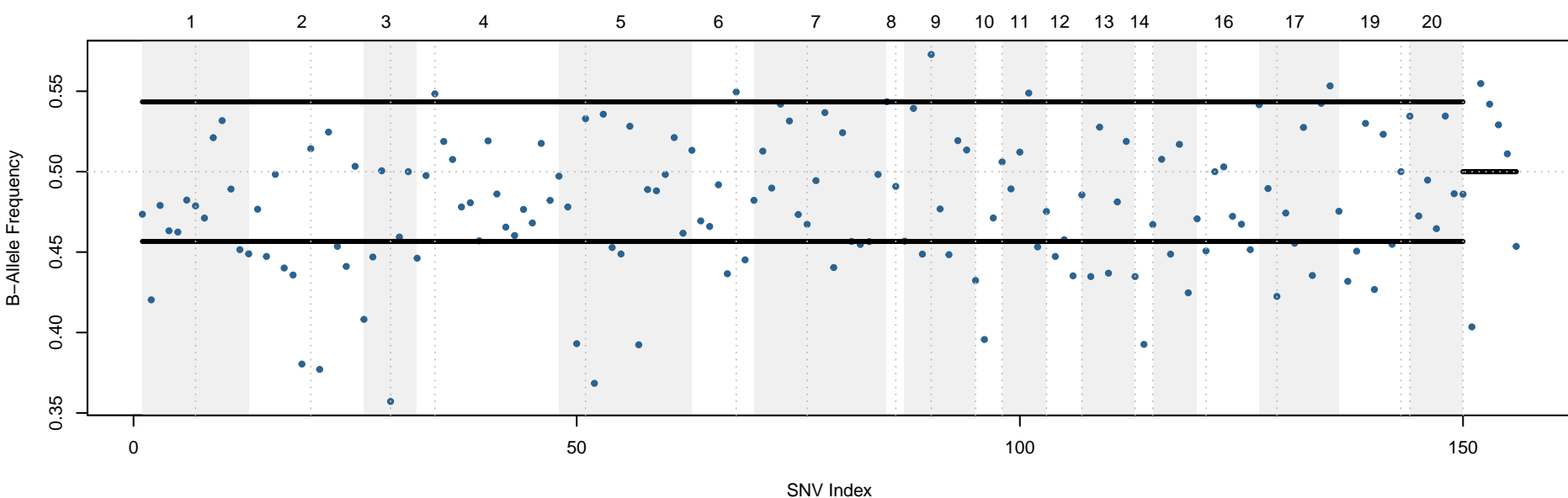




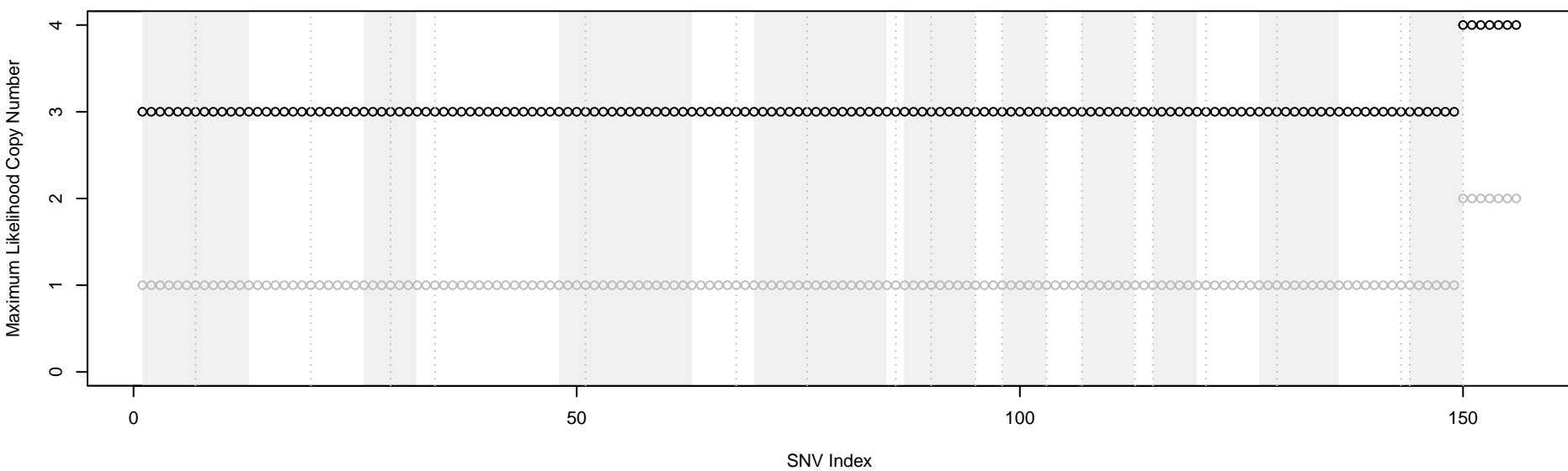
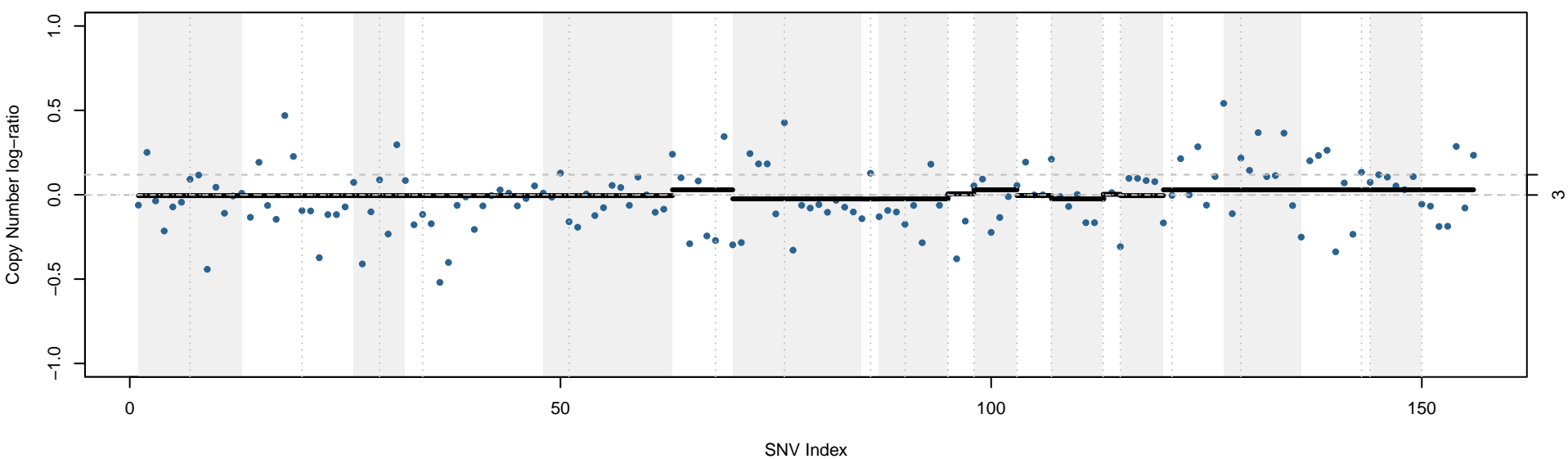
Purity: 0.19 Tumor ploidy: 3.013

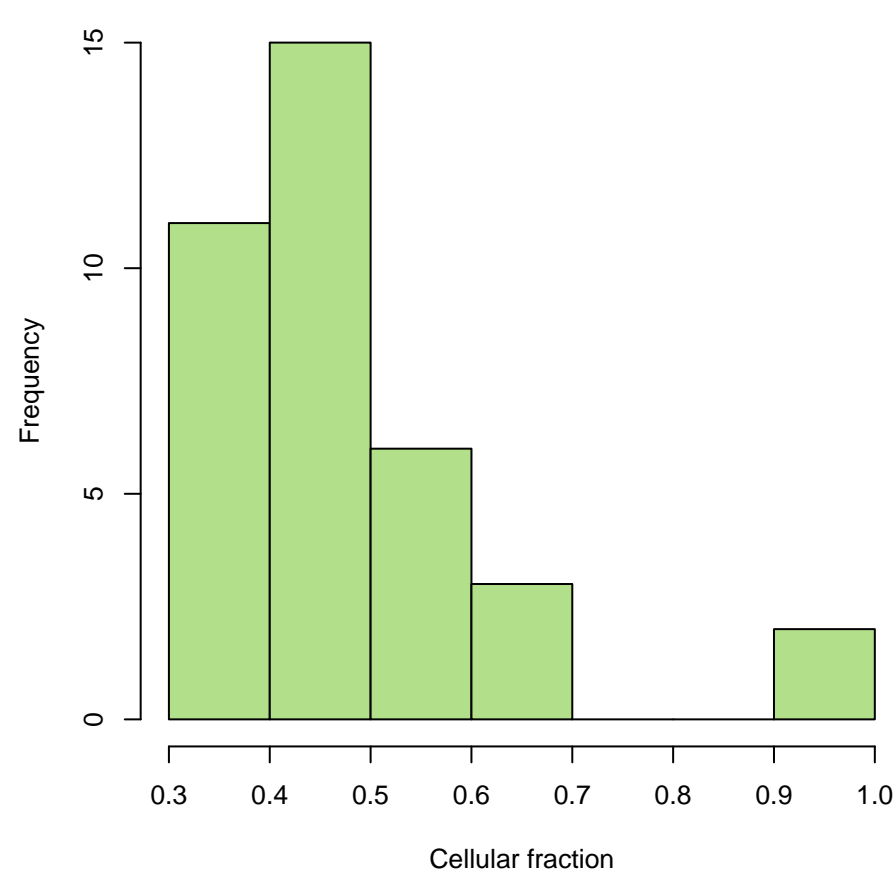
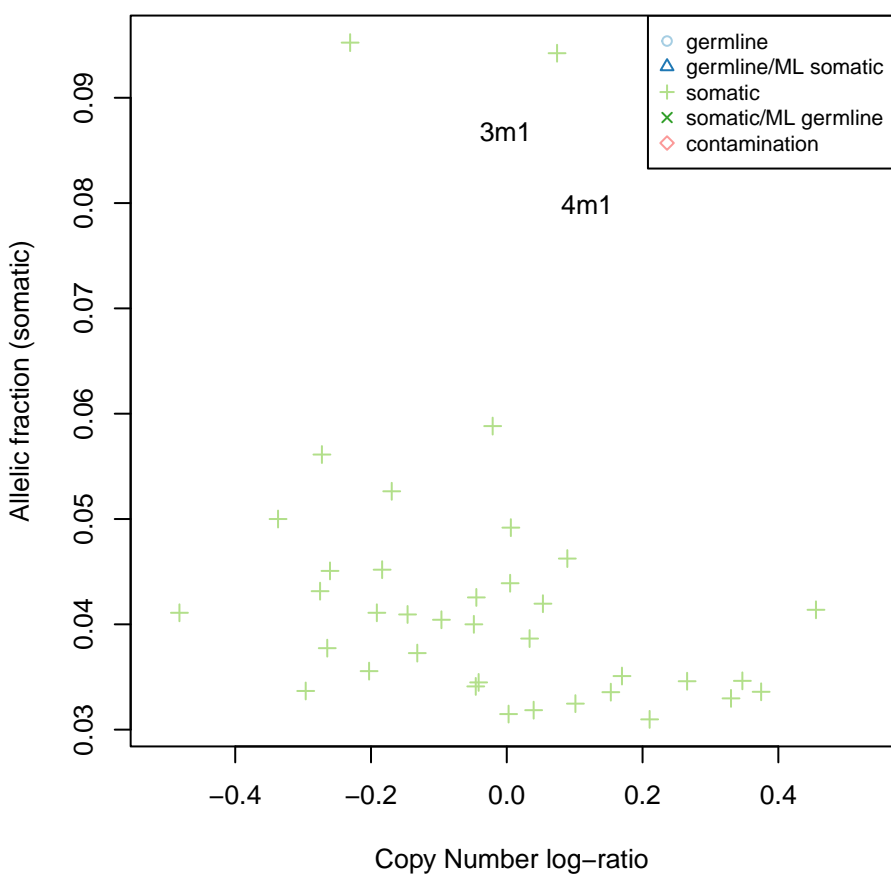
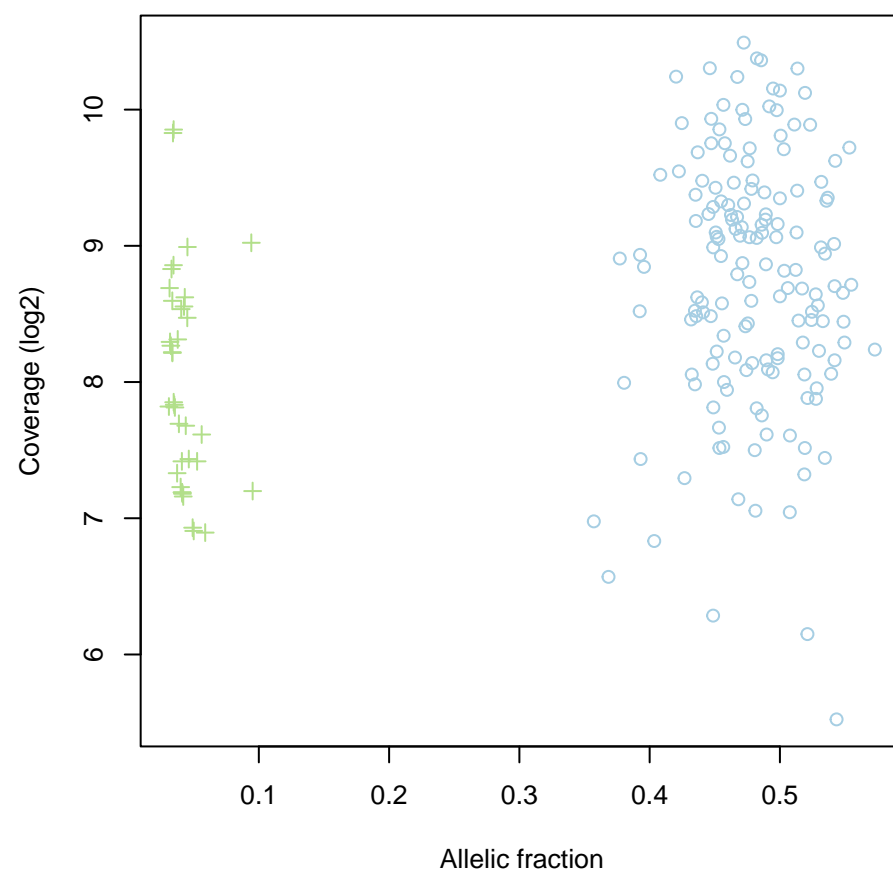
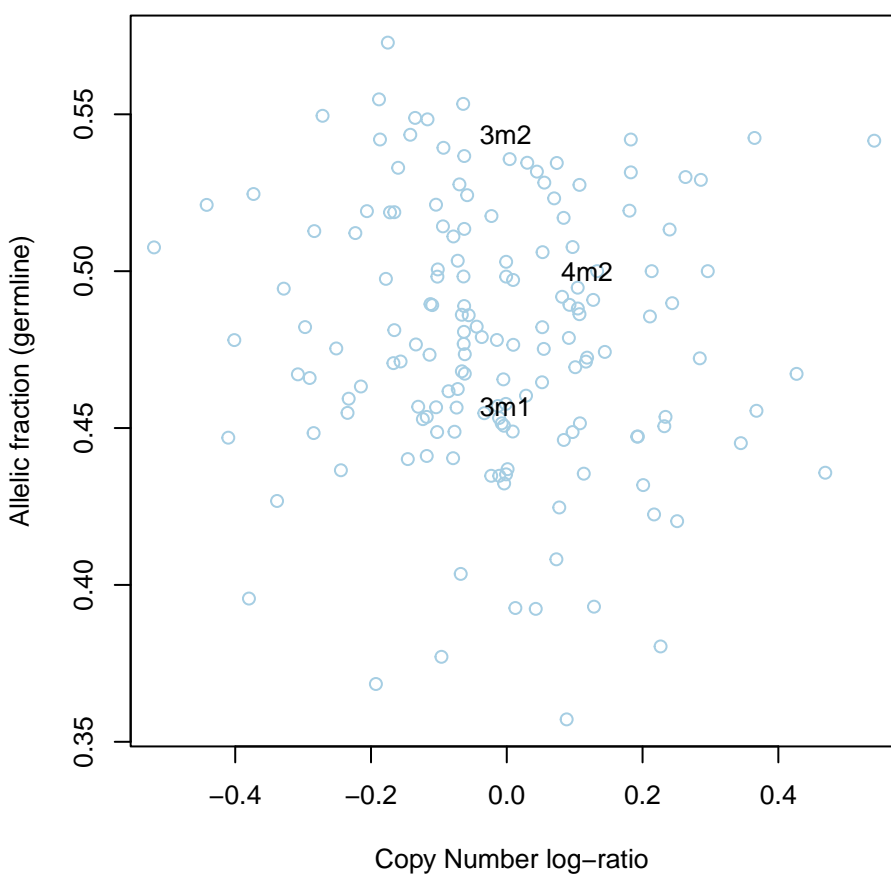


Purity: 0.19 Tumor ploidy: 3.013 SNV log-likelihood: 19.73 GoF: 97.3% Mean coverage: 410;477

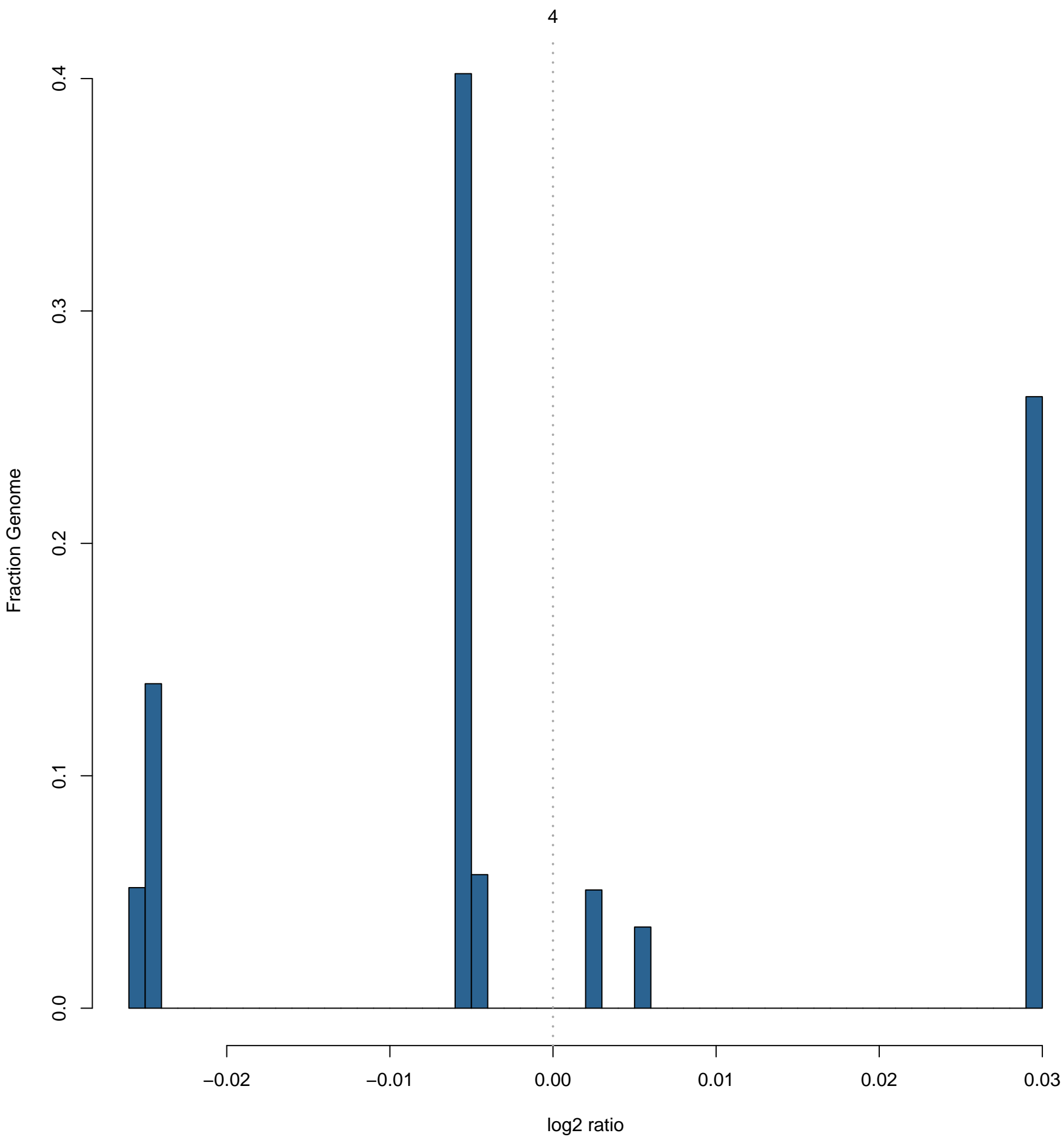


SCNA-fit log-likelihood: -7174.73

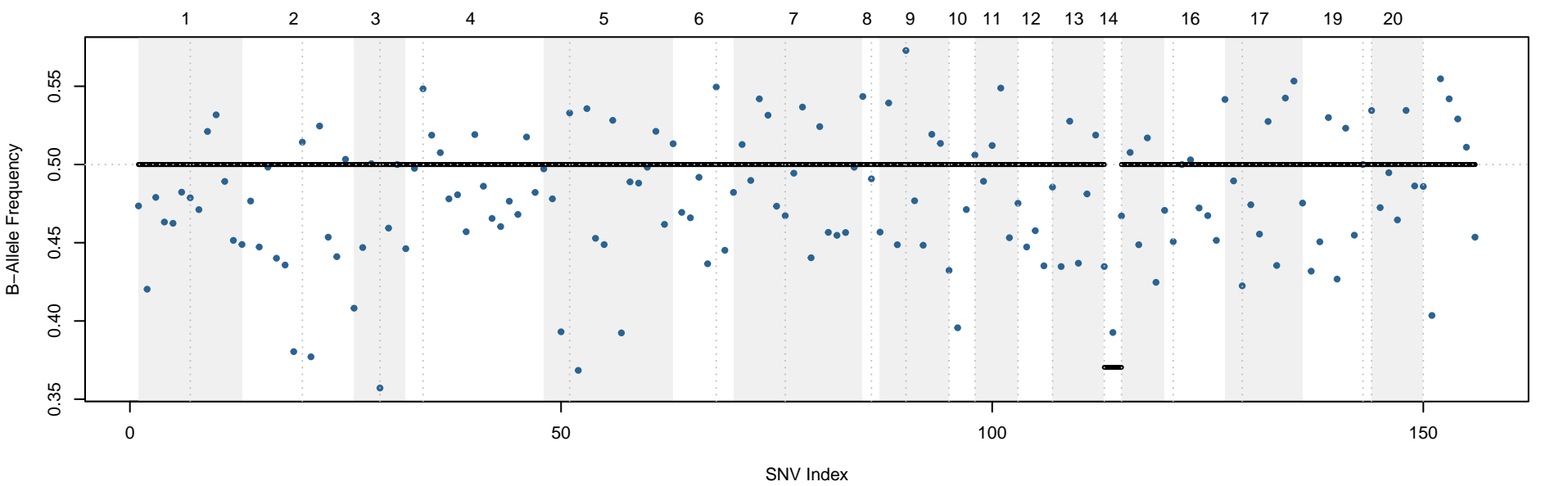




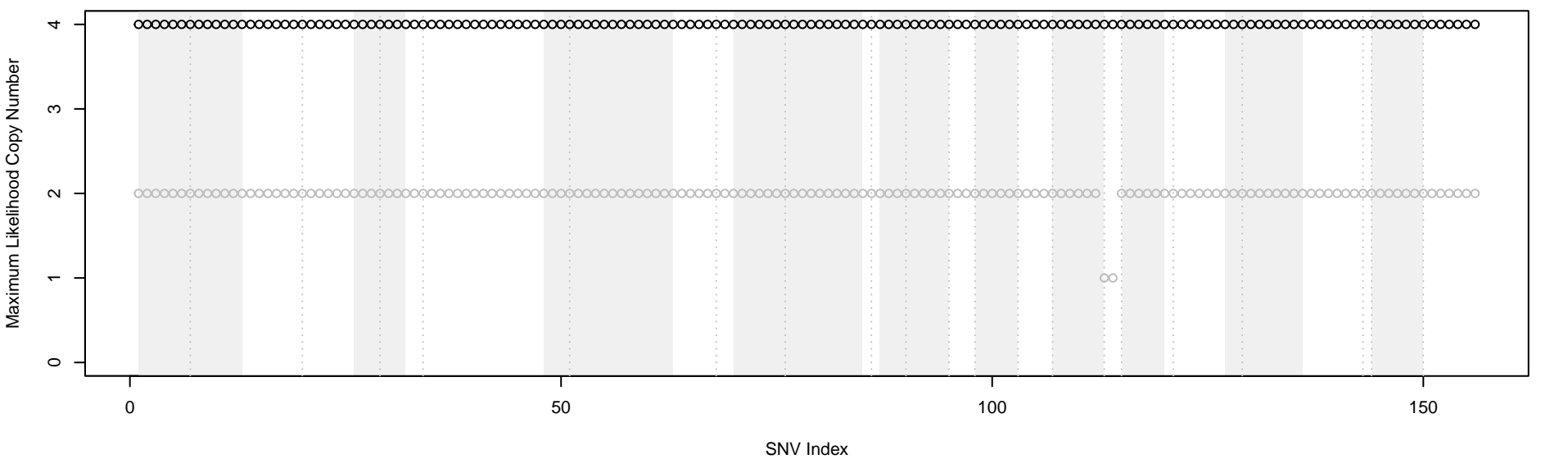
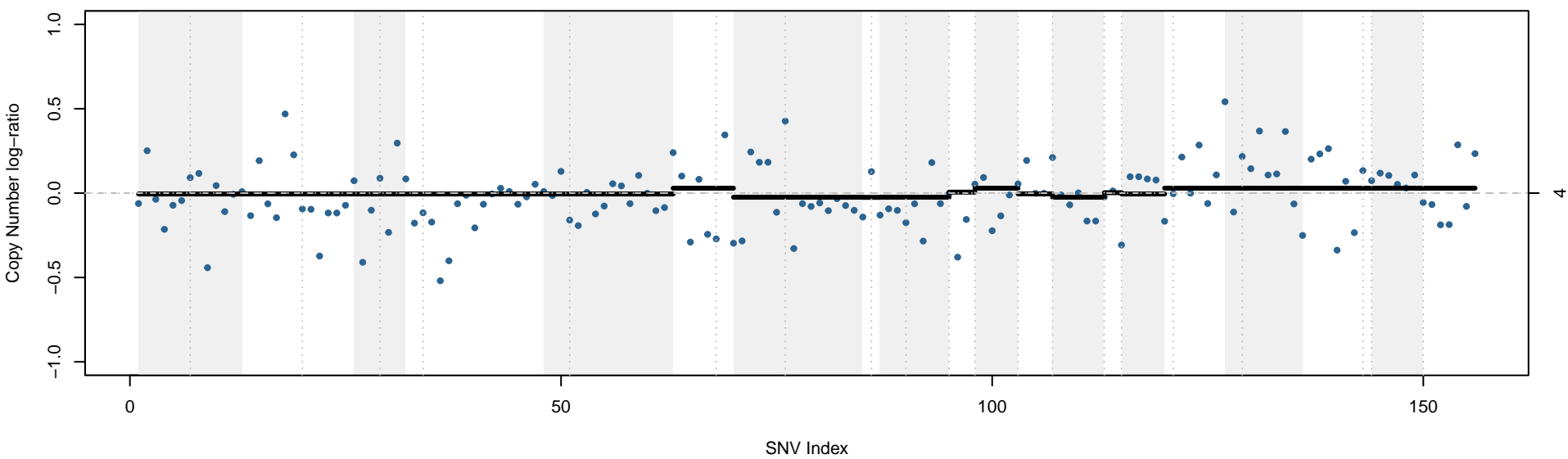
Purity: 0.35 Tumor ploidy: 4



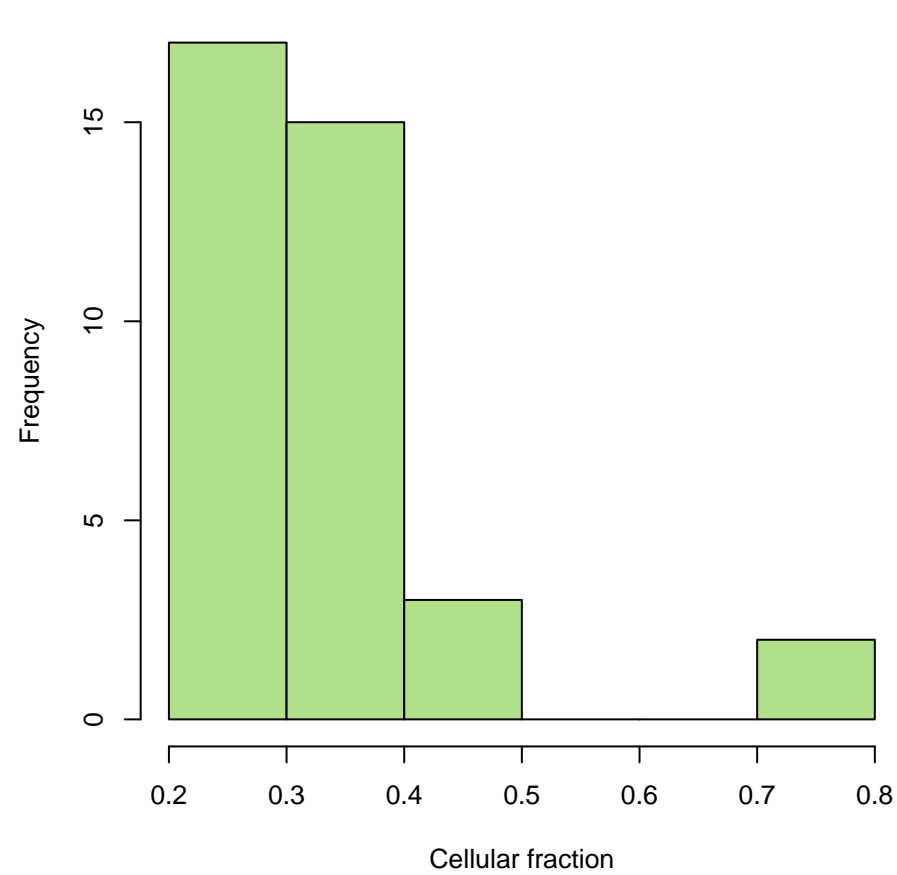
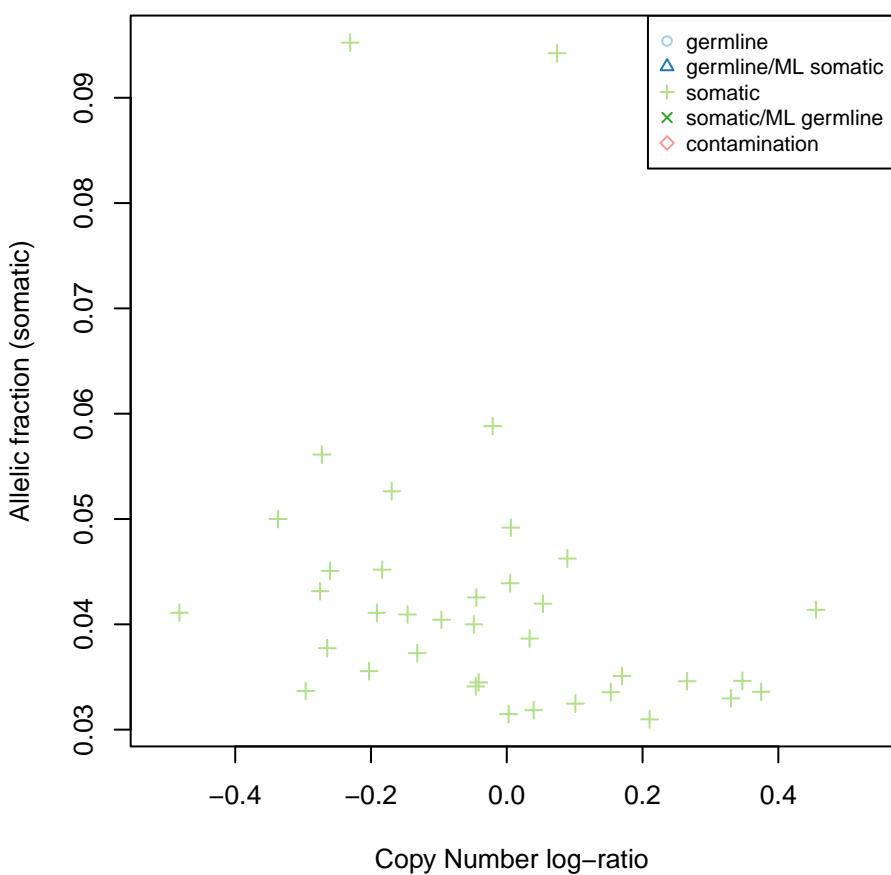
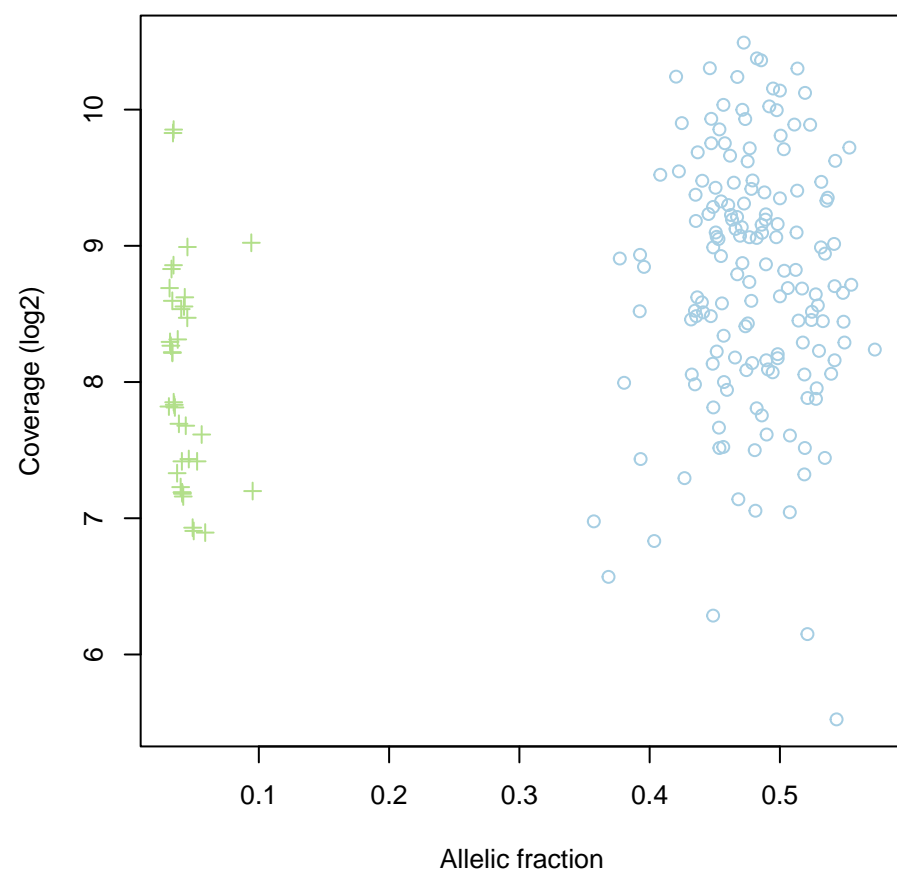
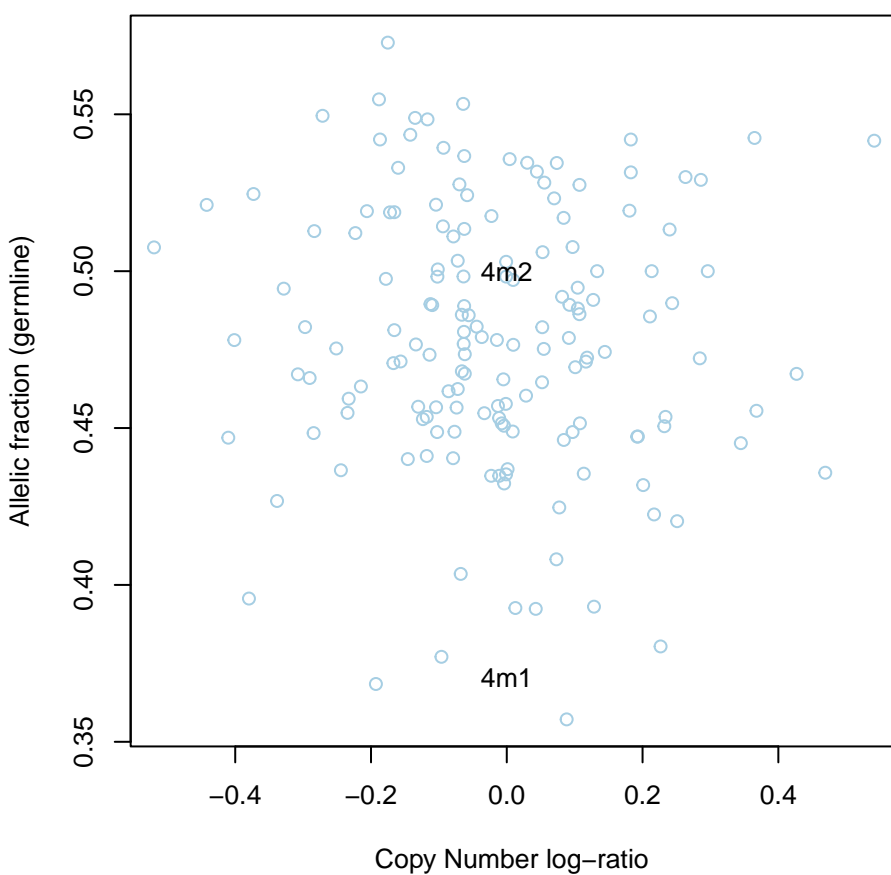
Purity: 0.35 Tumor ploidy: 4 SNV log-likelihood: 4.17 GoF: 92.6% Mean coverage: 410;477



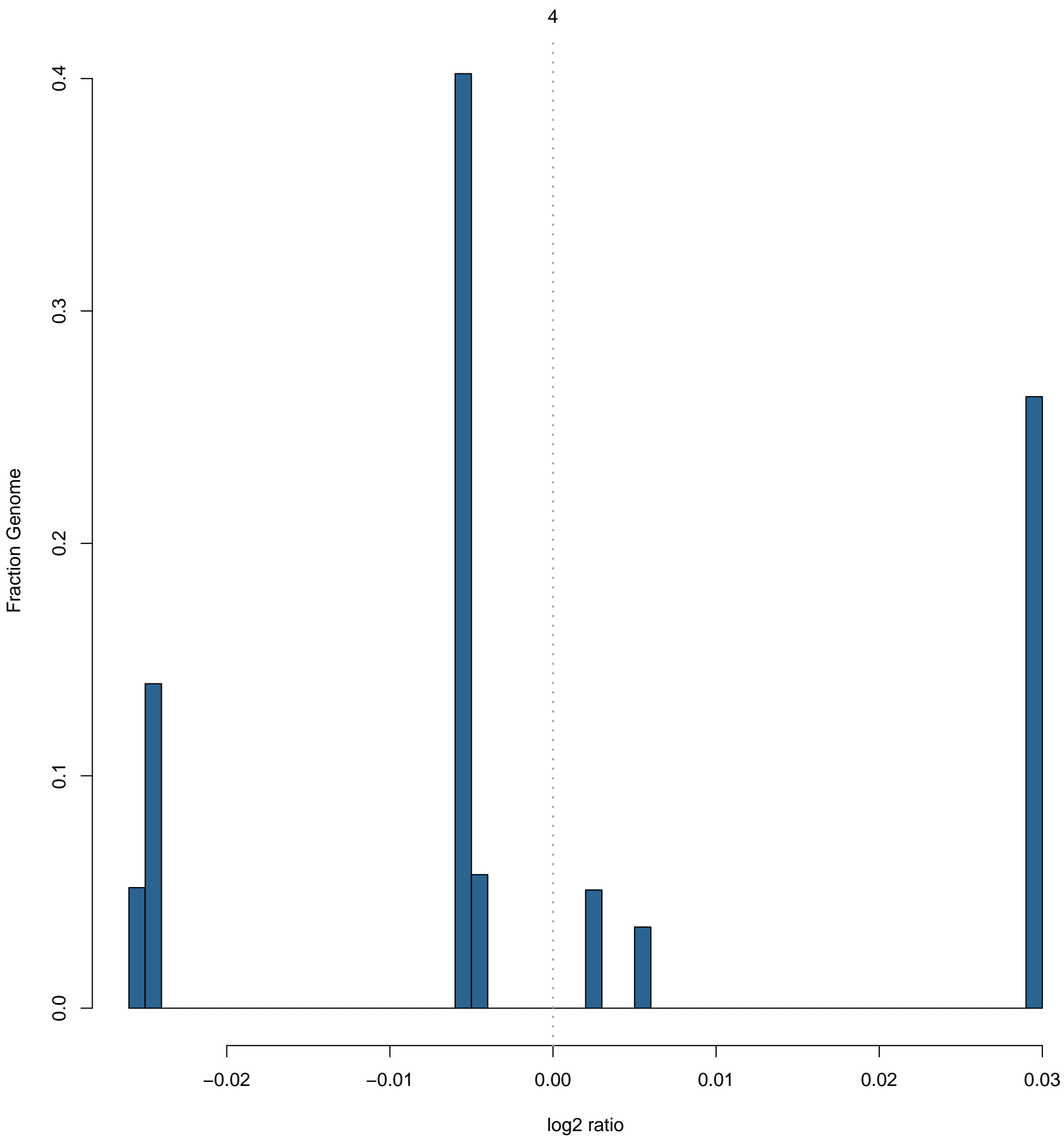
SCNA-fit log-likelihood: -7180.74



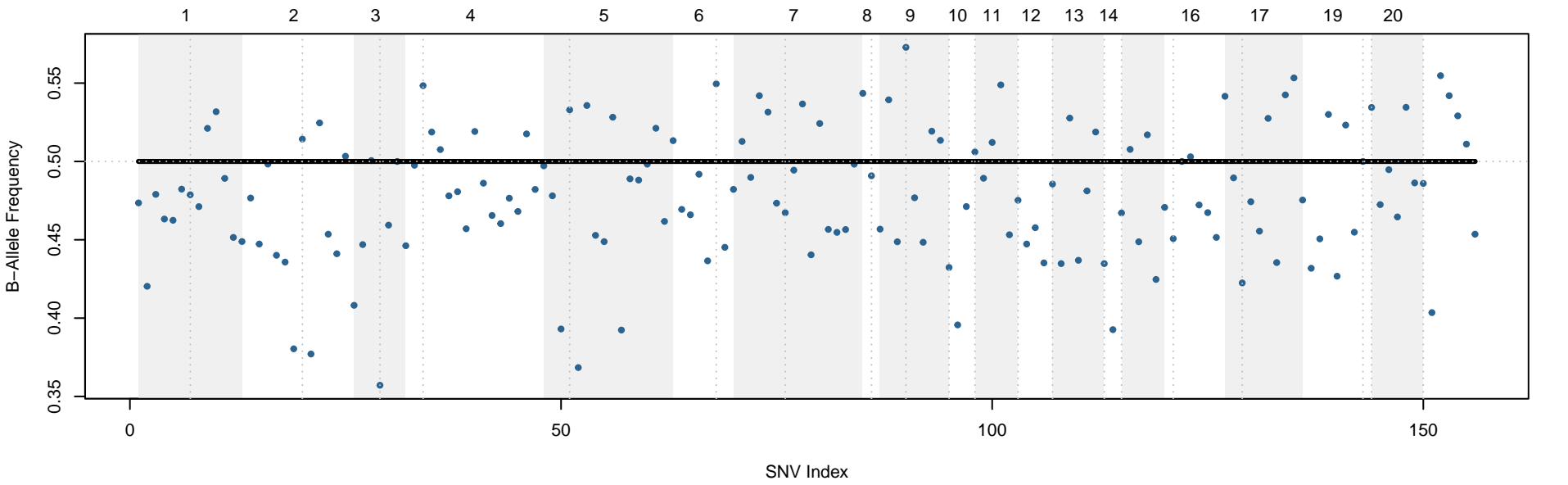




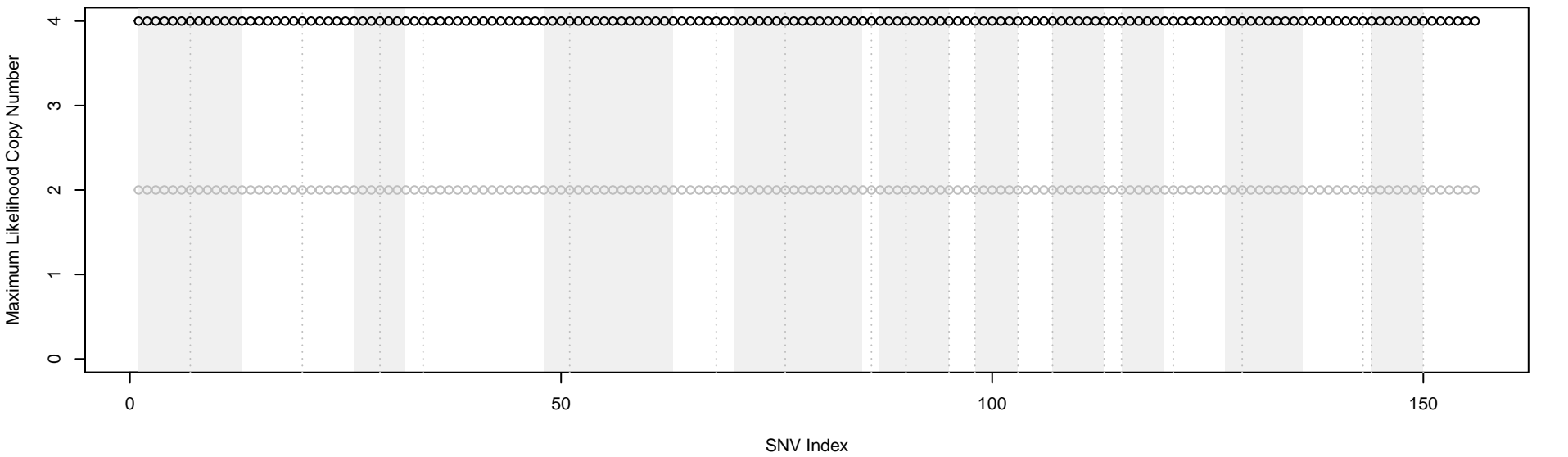
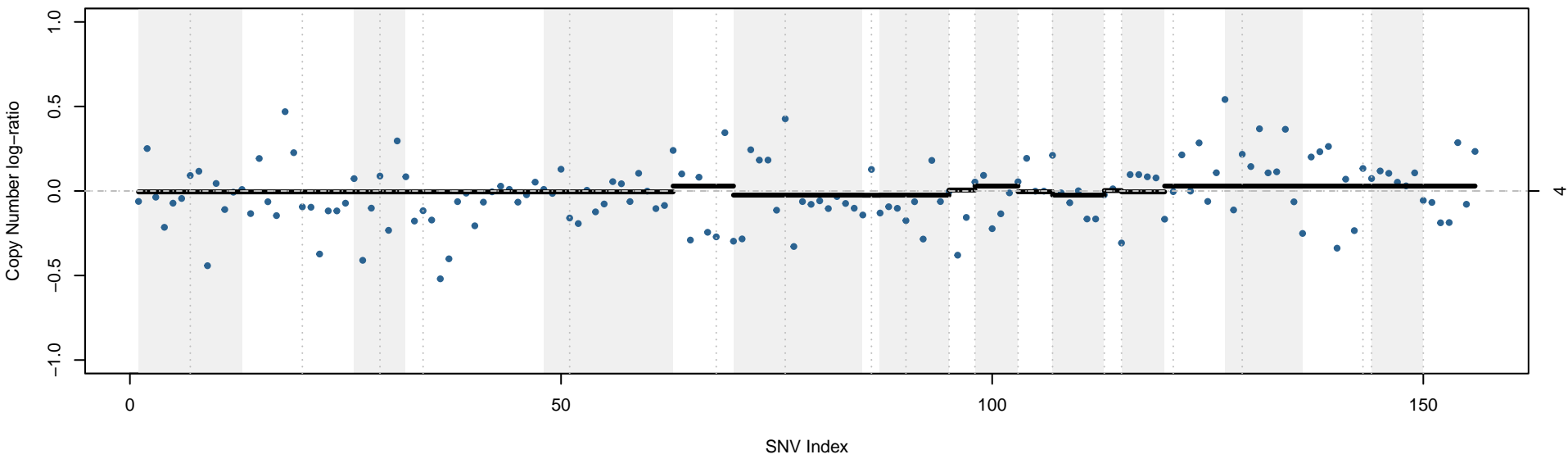
Purity: 0.54 Tumor ploidy: 4

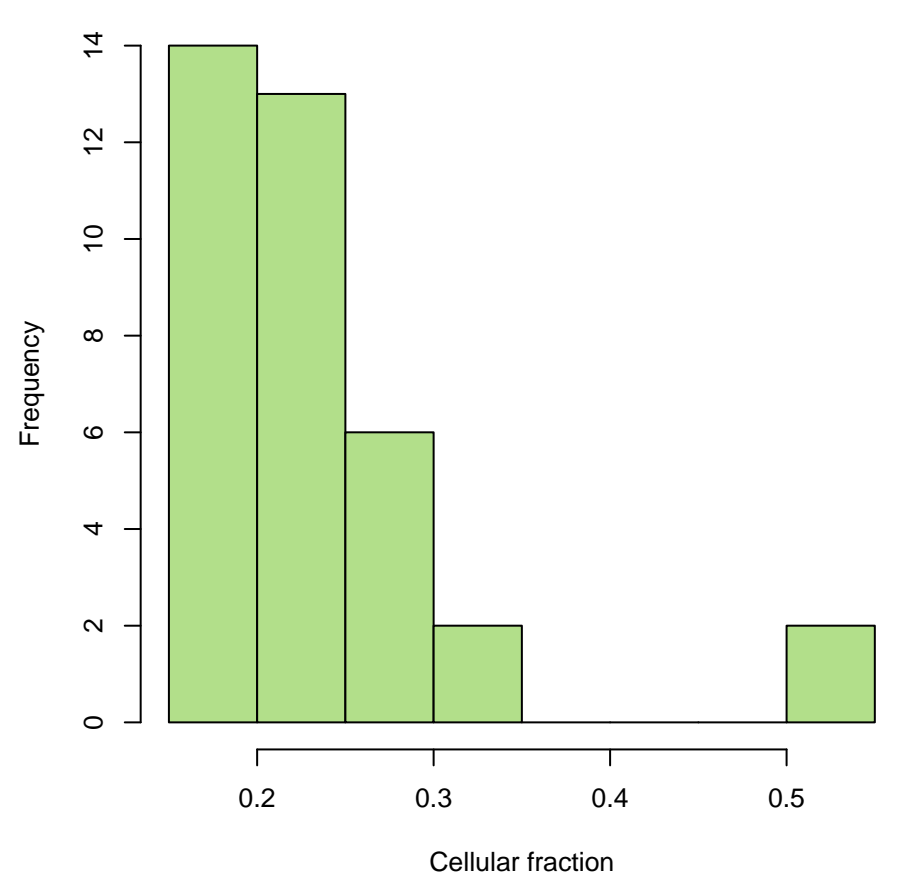
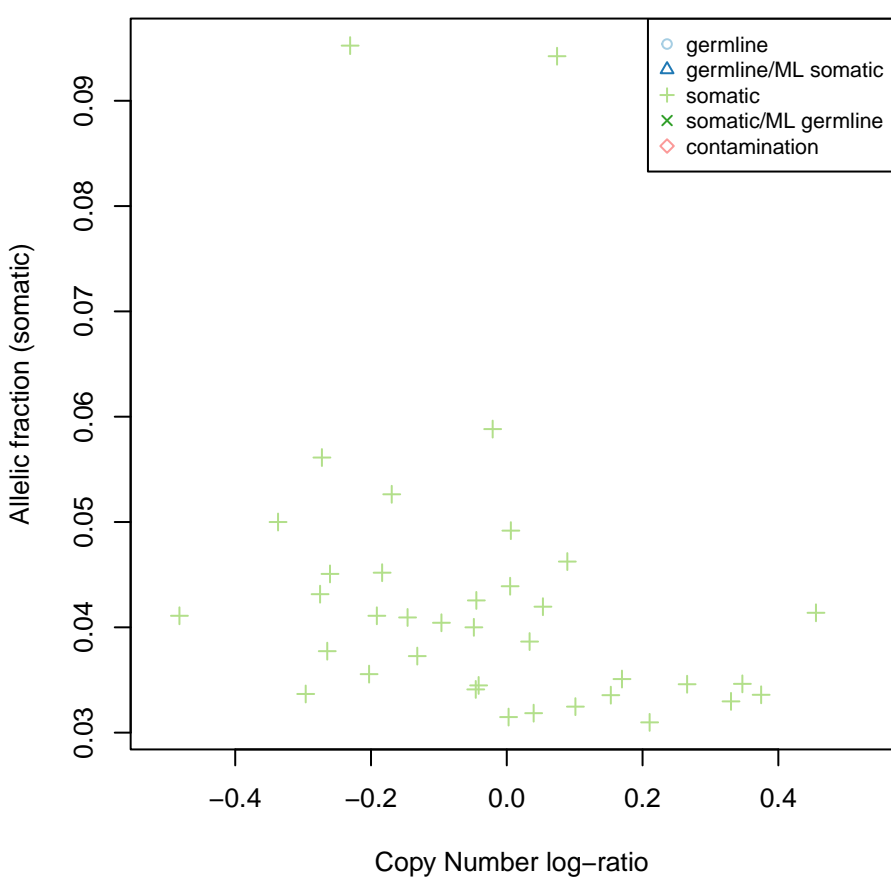
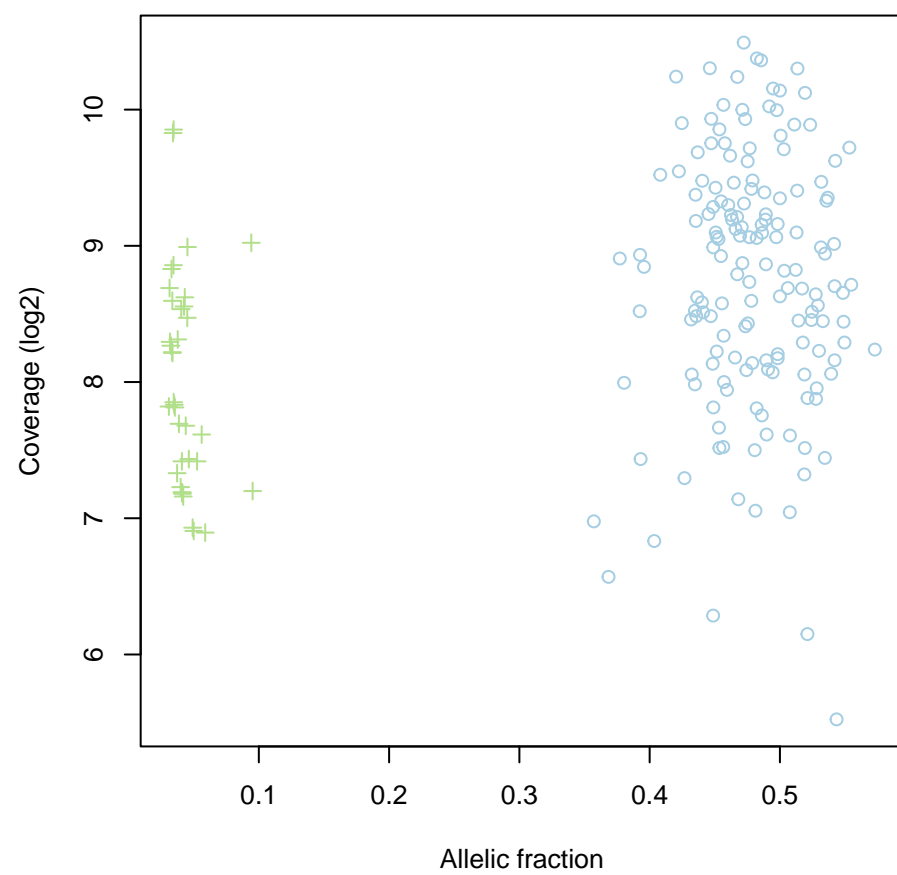
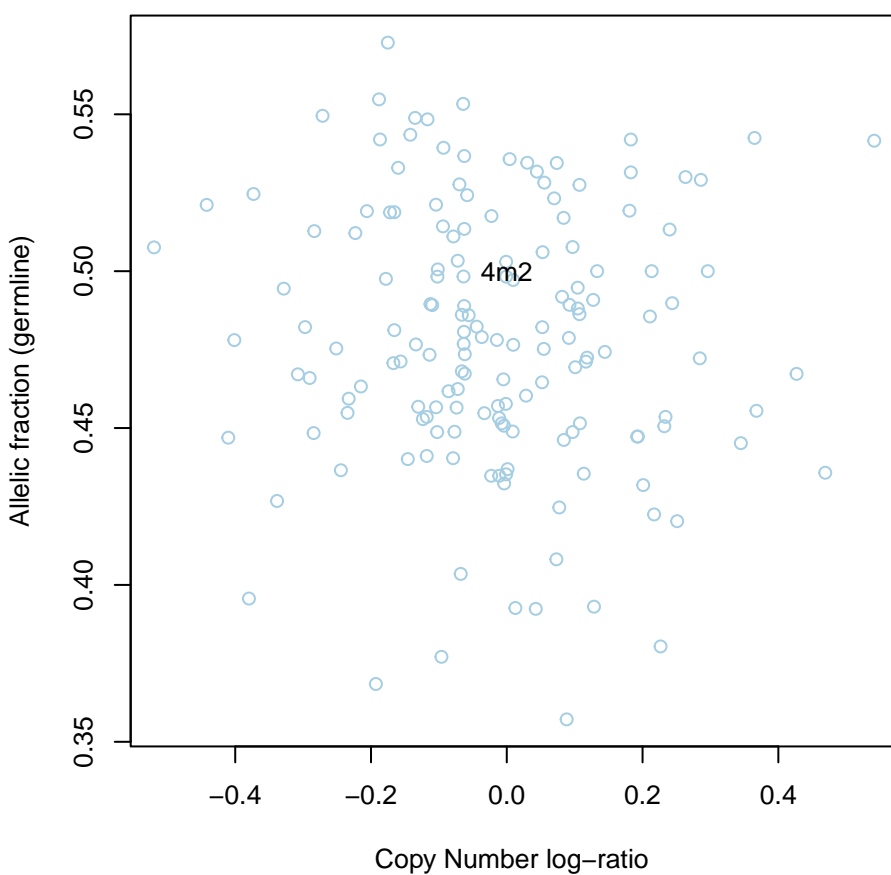


Purity: 0.54 Tumor ploidy: 4 SNV log-likelihood: -22.46 GoF: 87.7% Mean coverage: 410;477

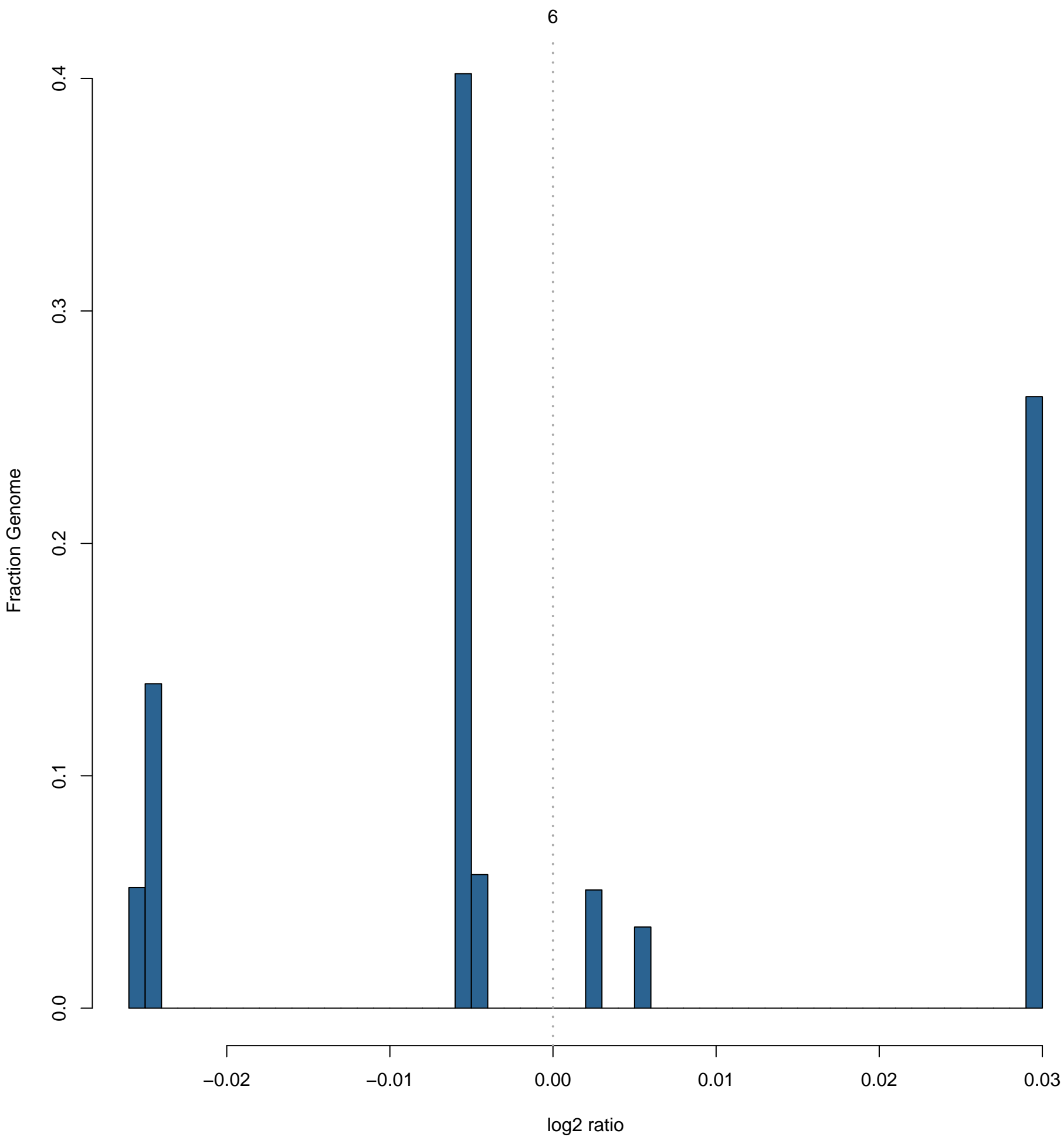


SCNA-fit log-likelihood: -7178.99

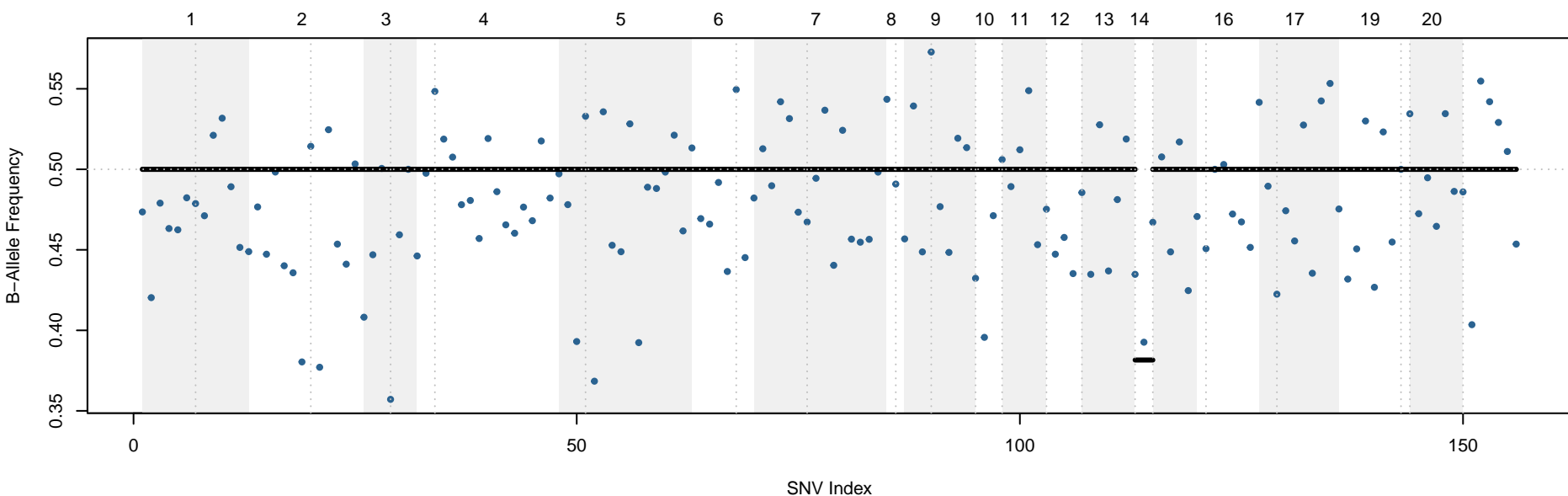




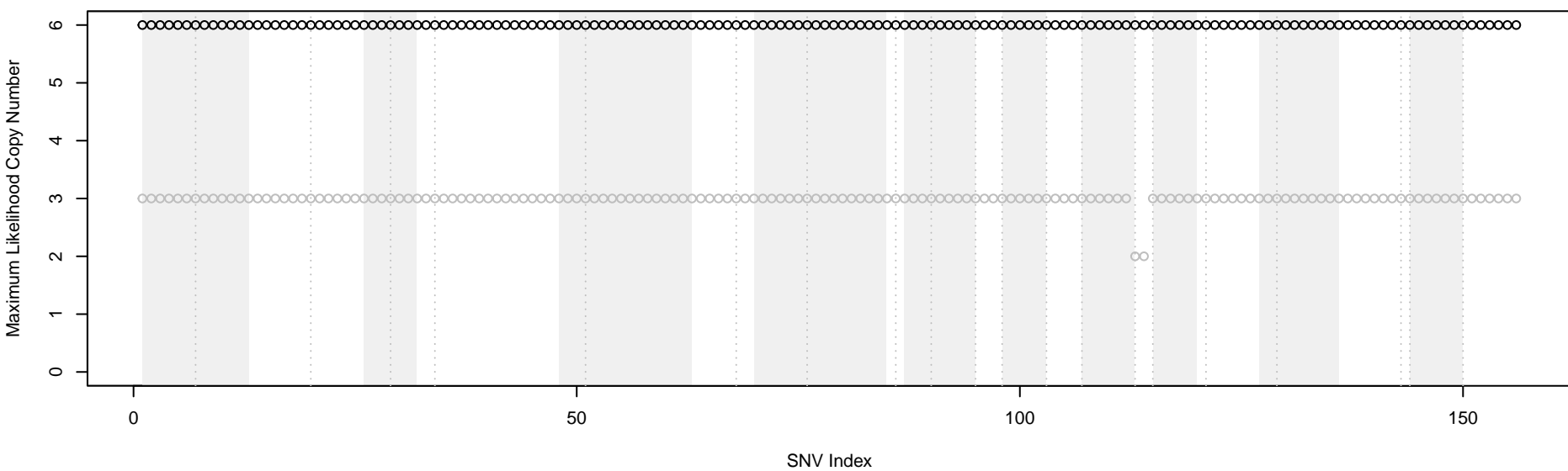
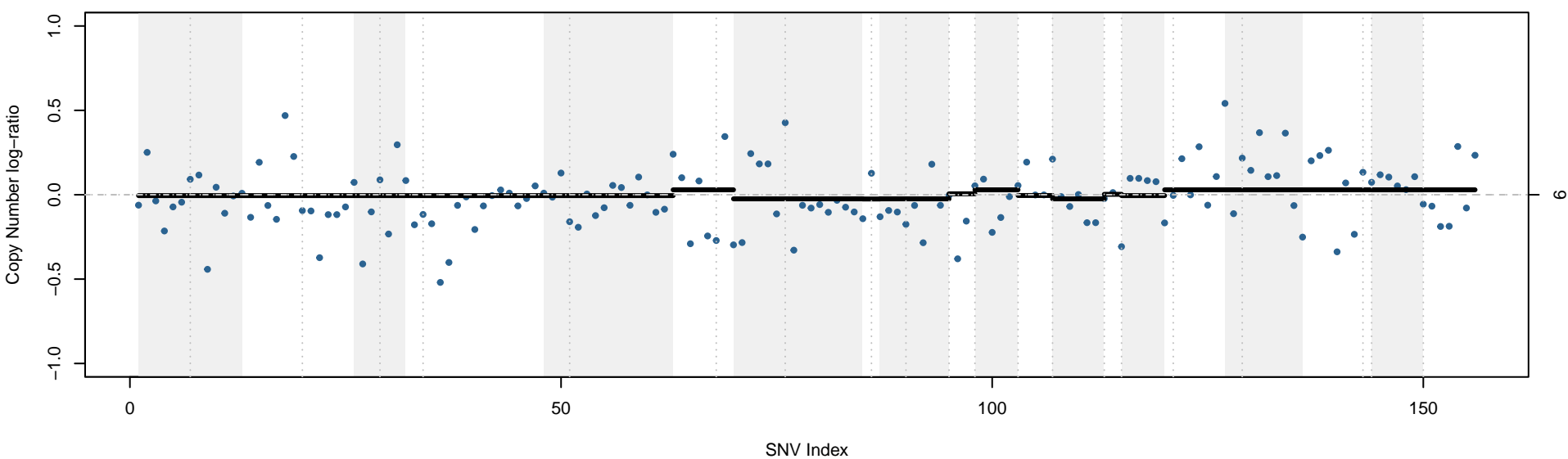
Purity: 0.45 Tumor ploidy: 6

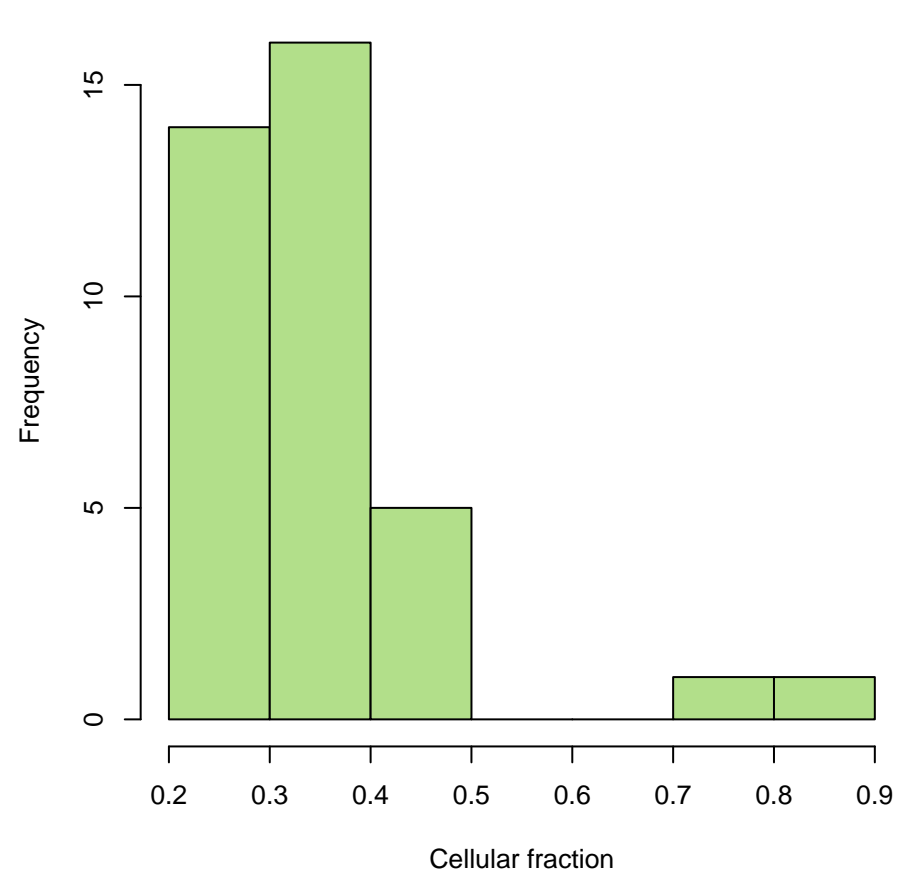
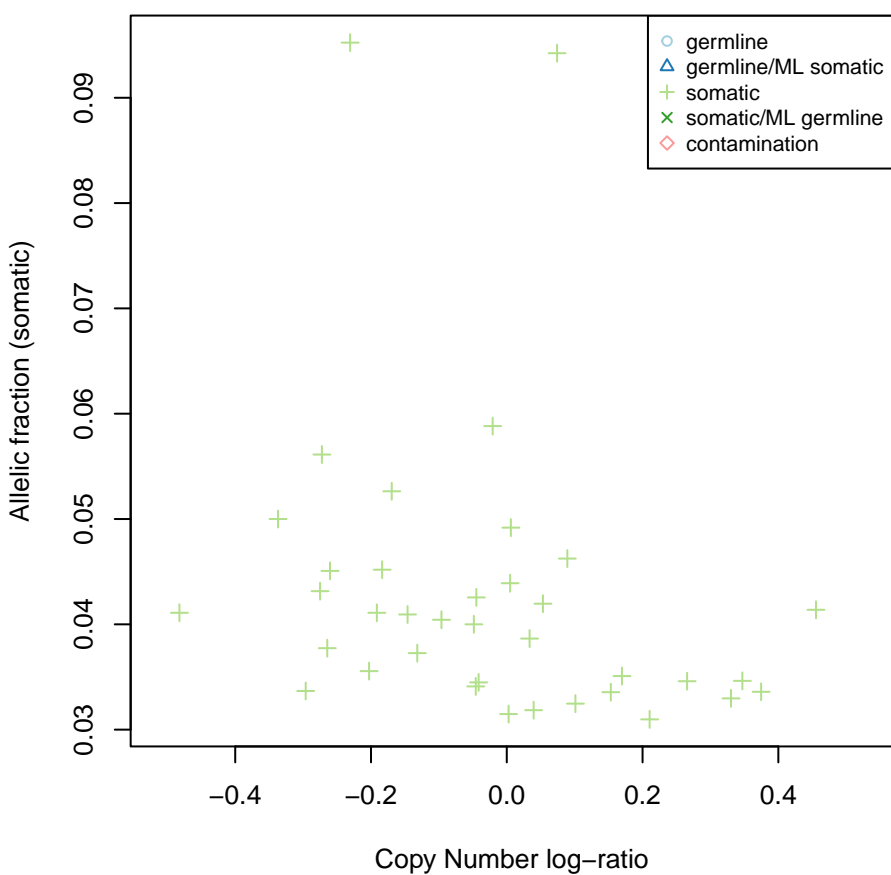
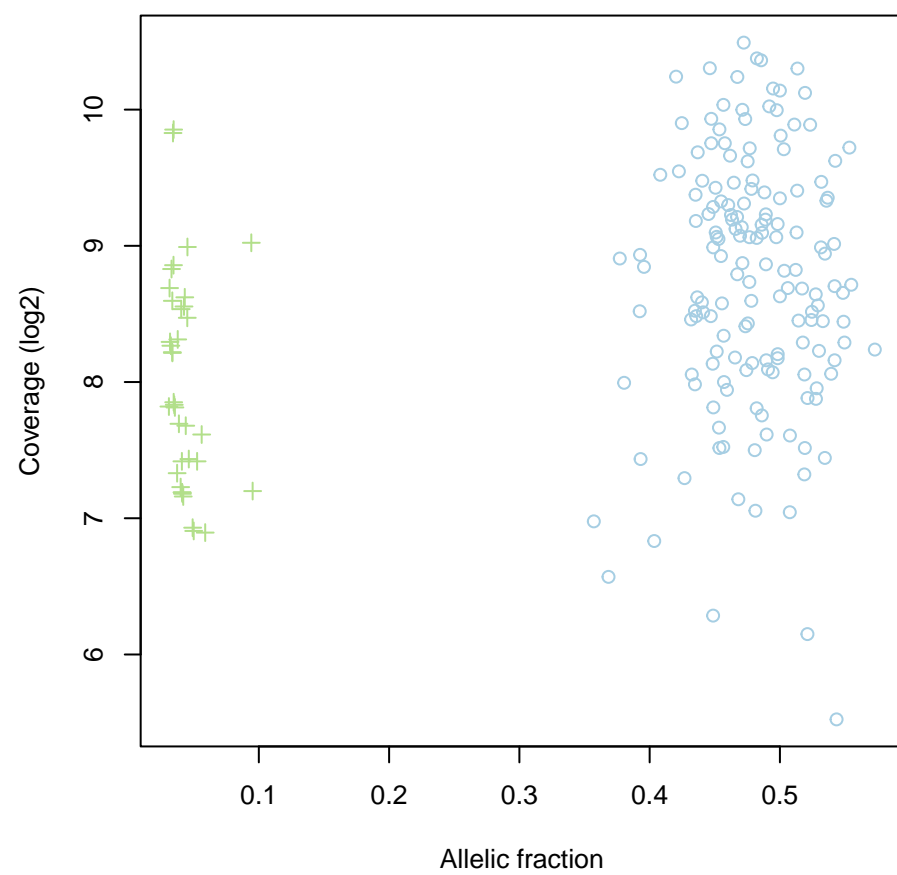
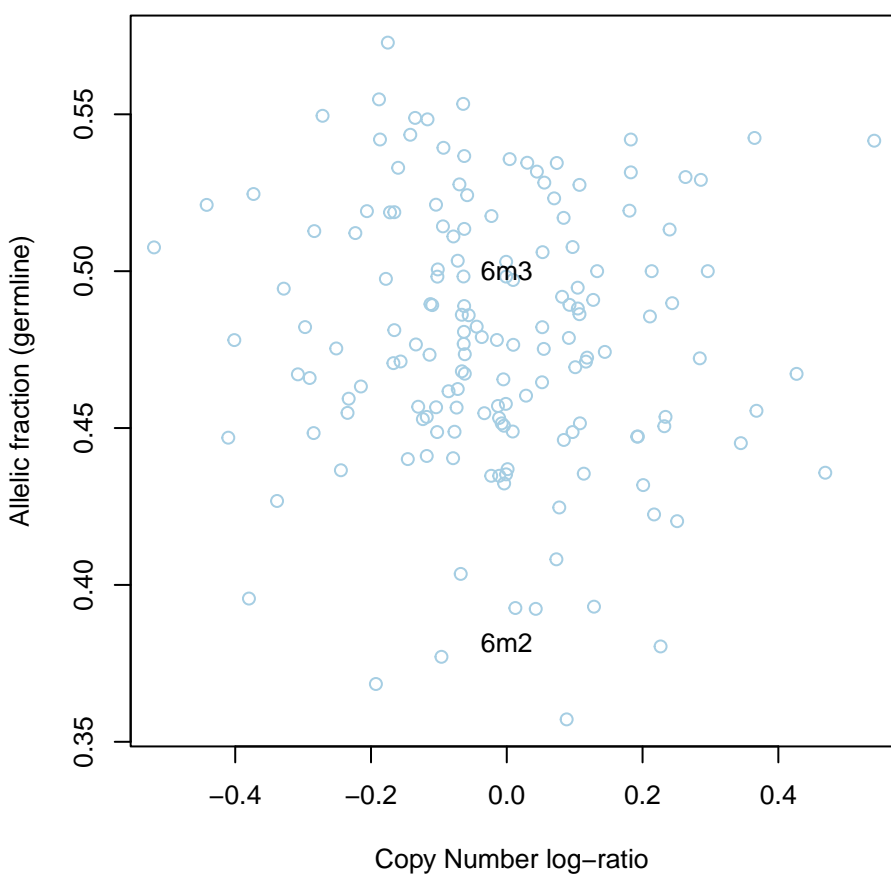


Purity: 0.45 Tumor ploidy: 6 SNV log-likelihood: -55.1 GoF: 93.4% Mean coverage: 410;477

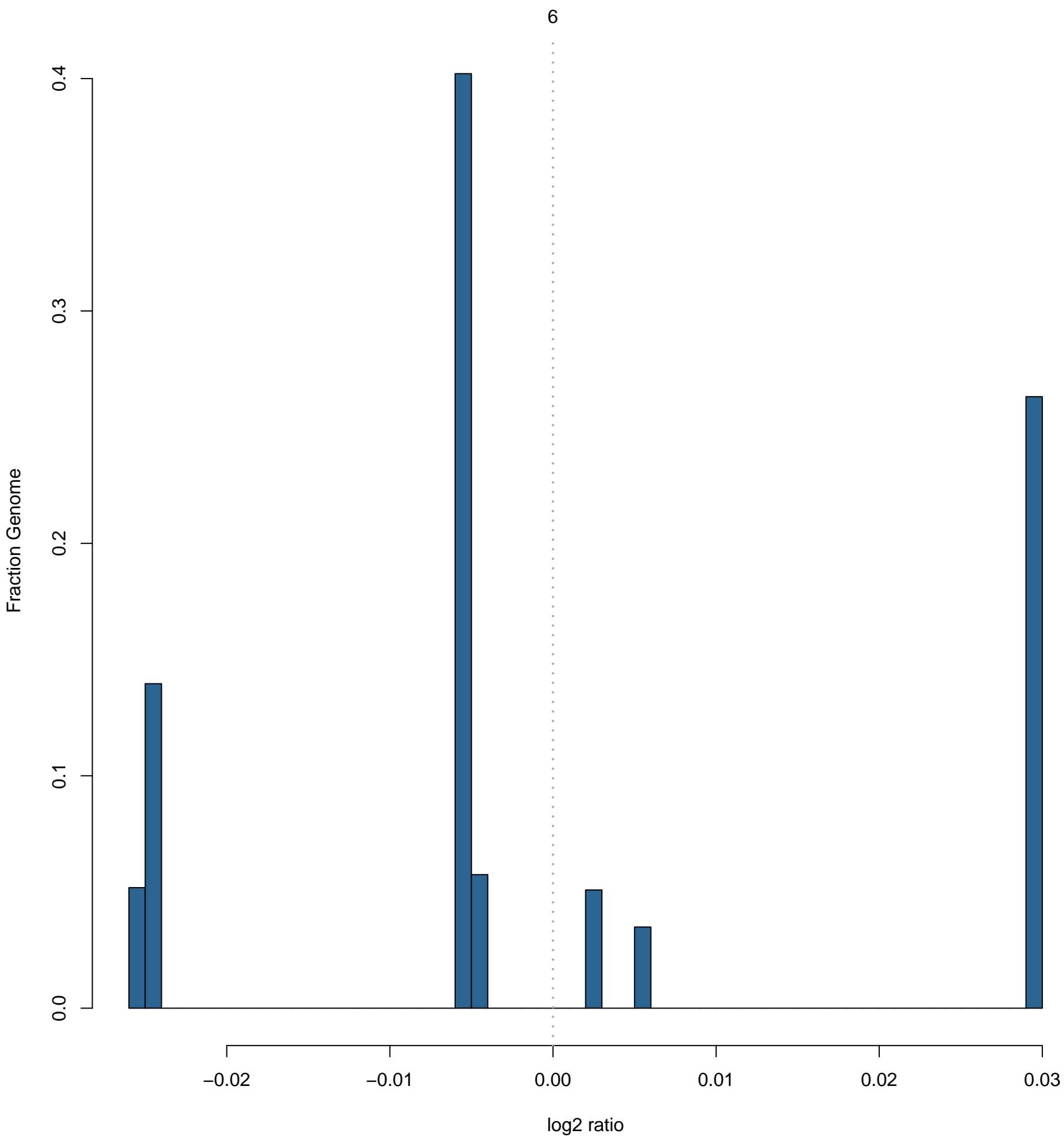


SCNA-fit log-likelihood: -7178.98



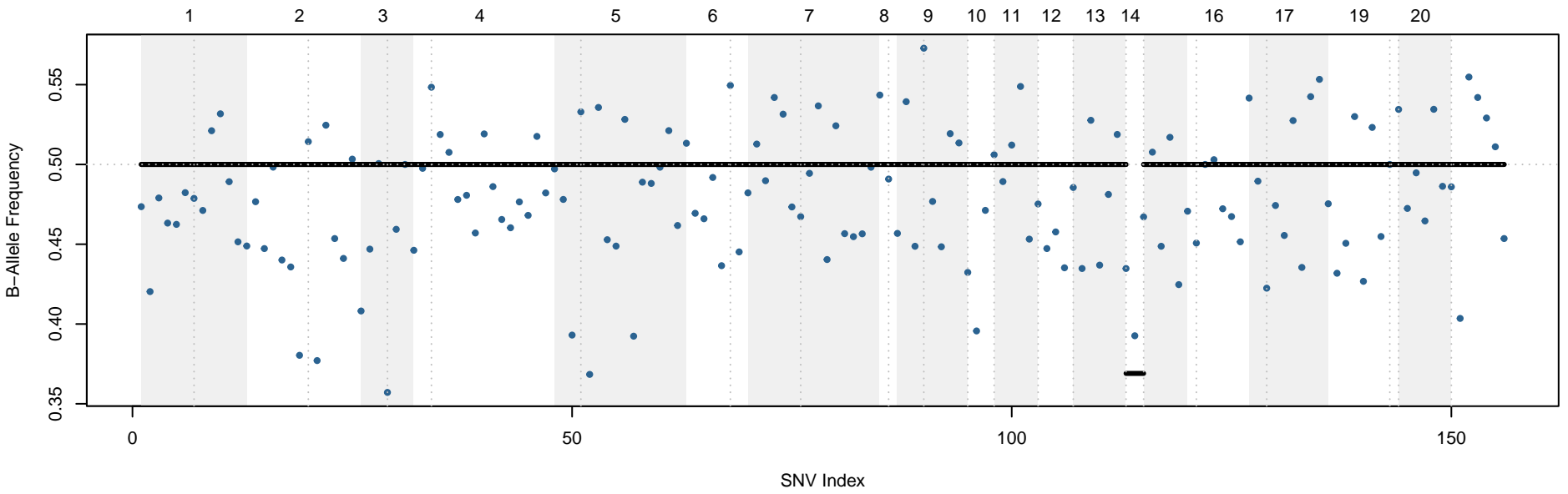


Purity: 0.55 Tumor ploidy: 6

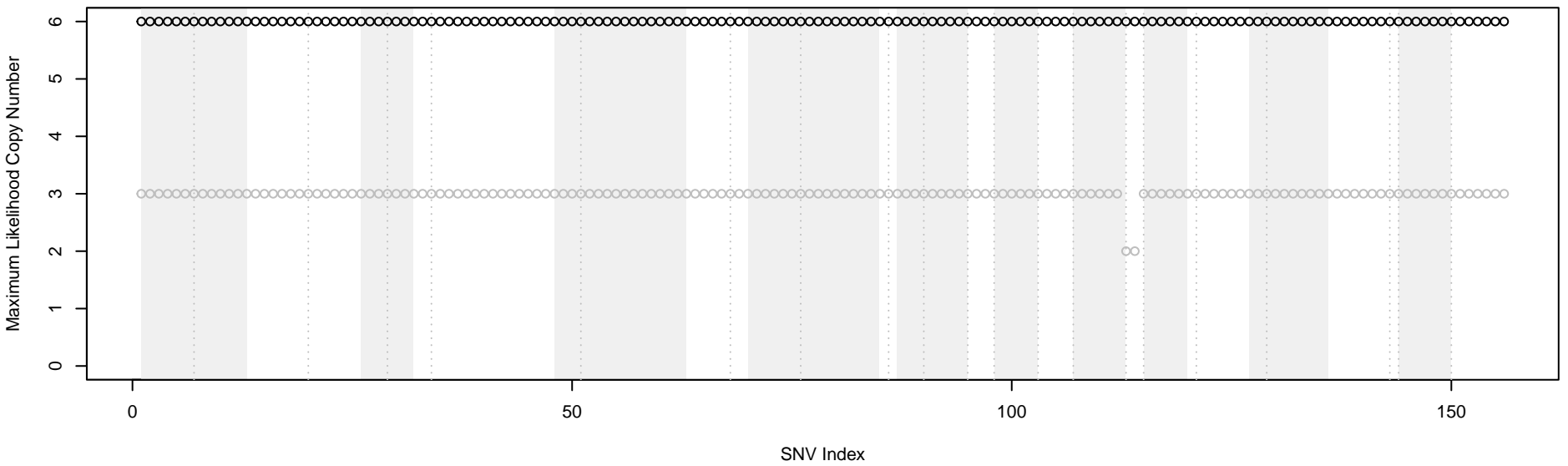
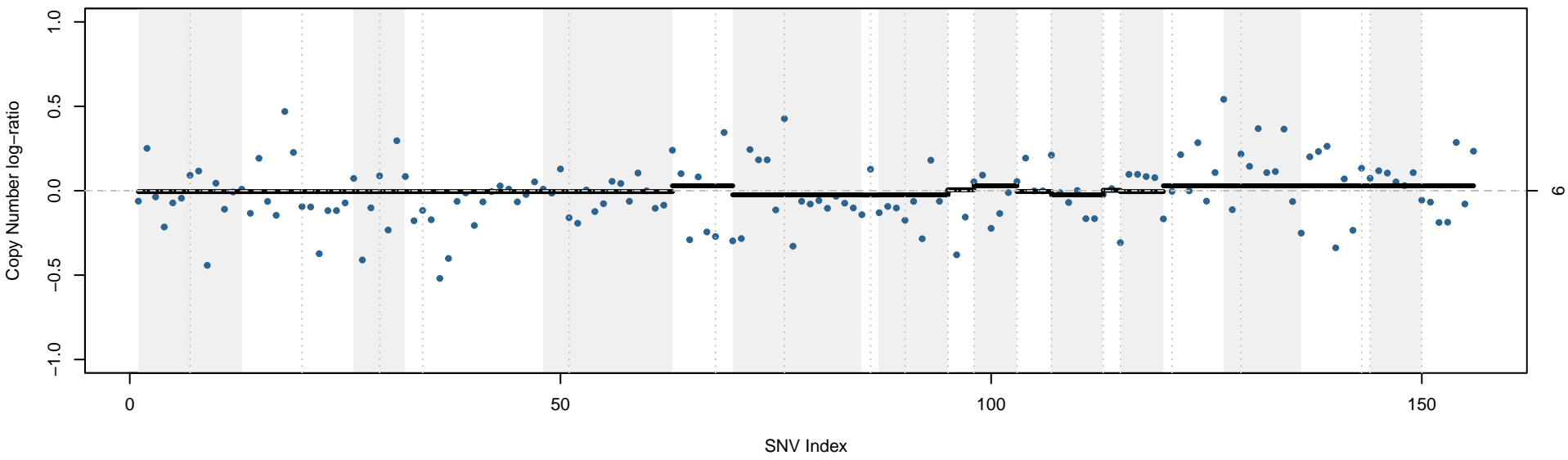


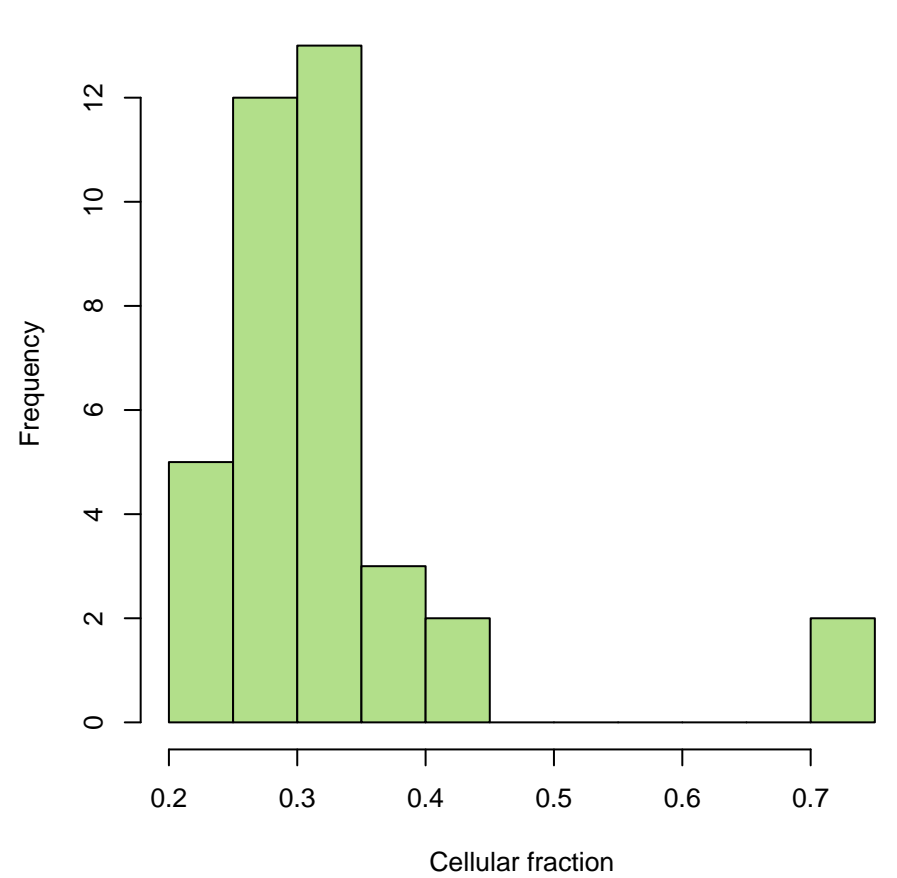
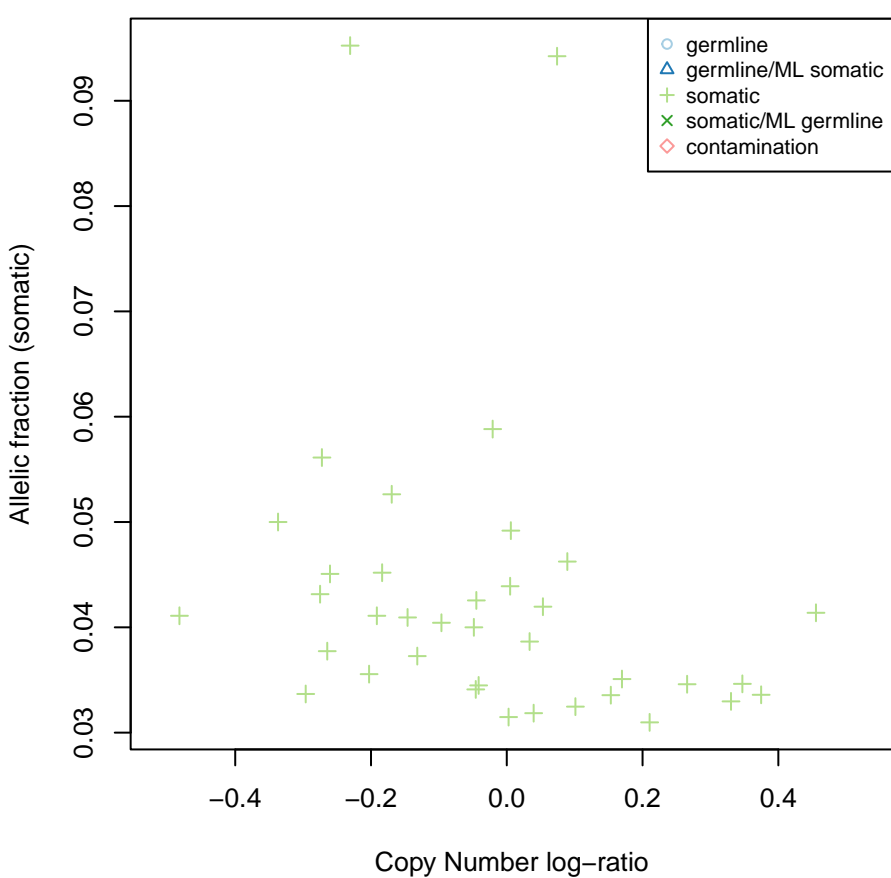
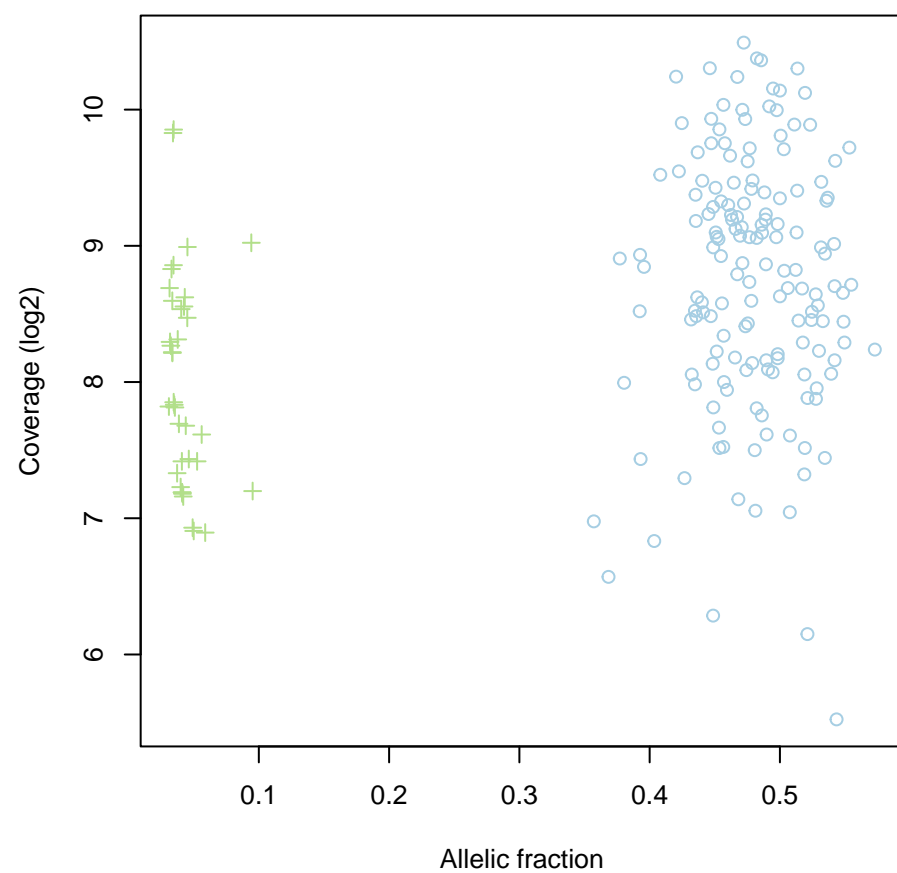
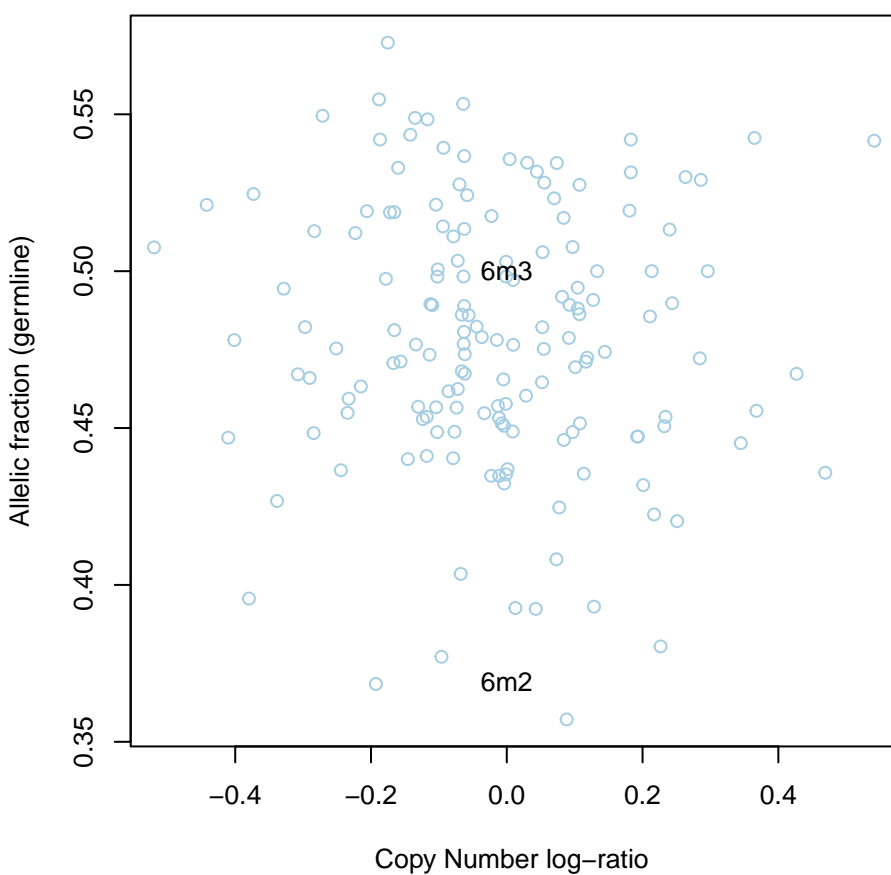


Purity: 0.55 Tumor ploidy: 6 SNV log-likelihood: -58.59 GoF: 92.4% Mean coverage: 410;477

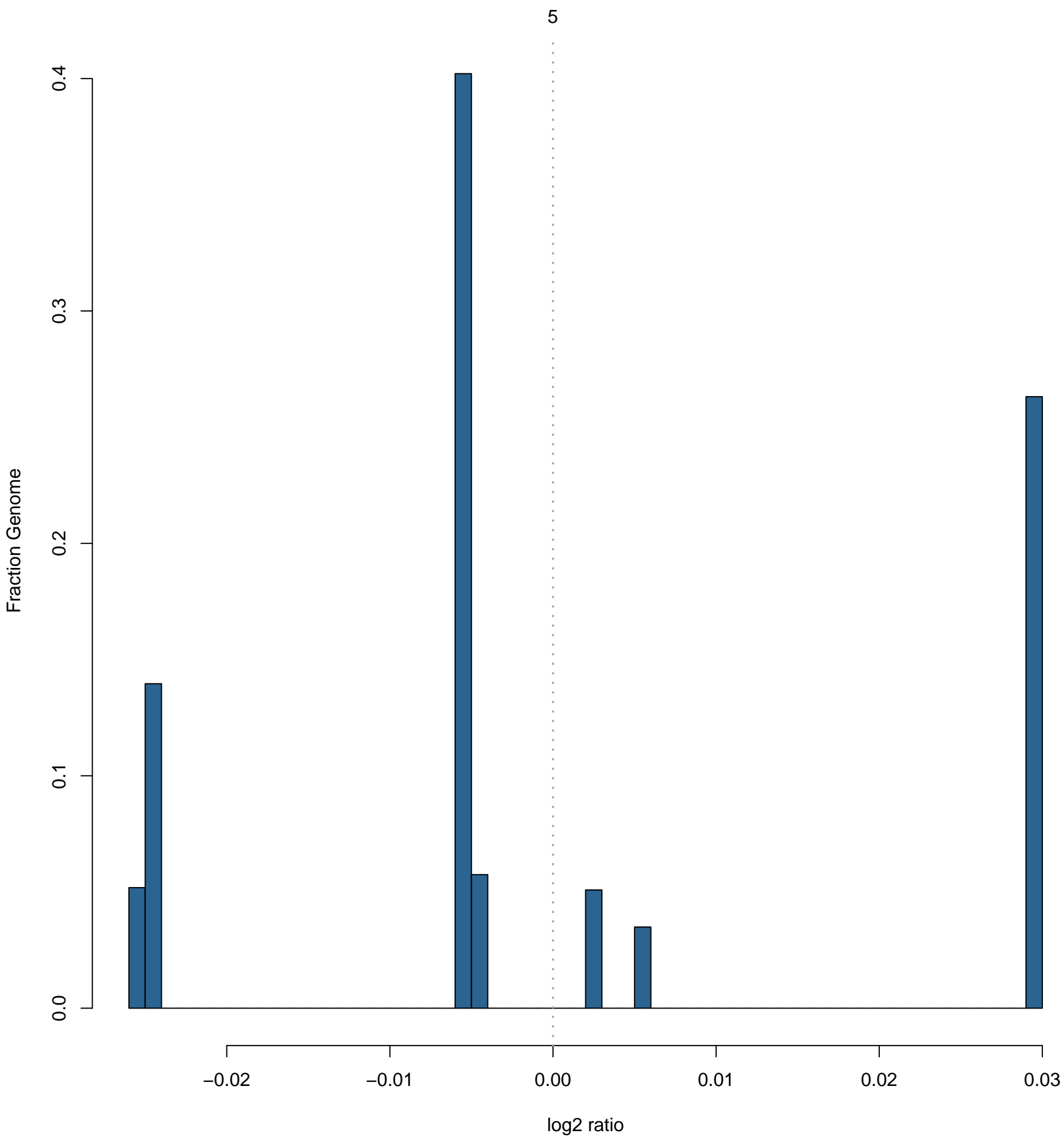


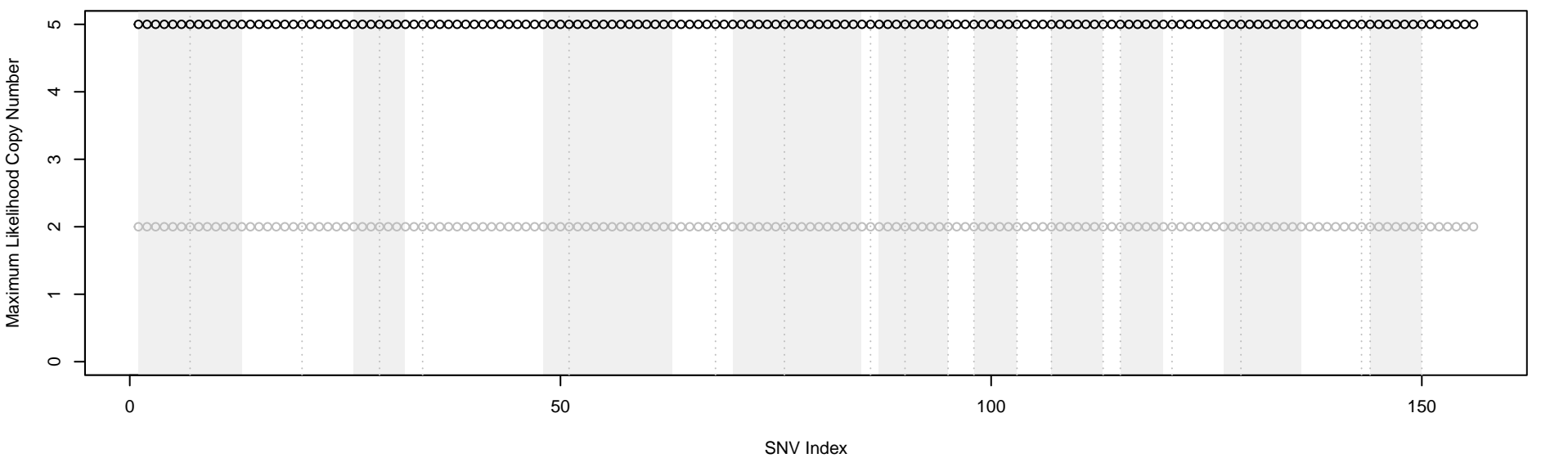
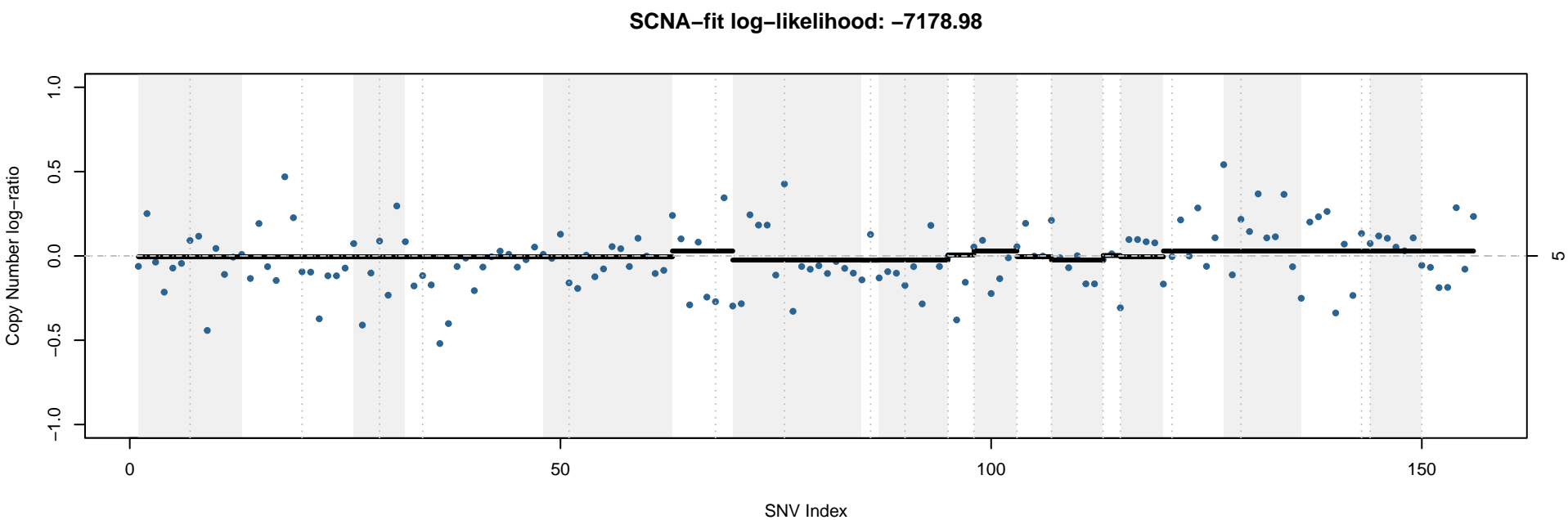
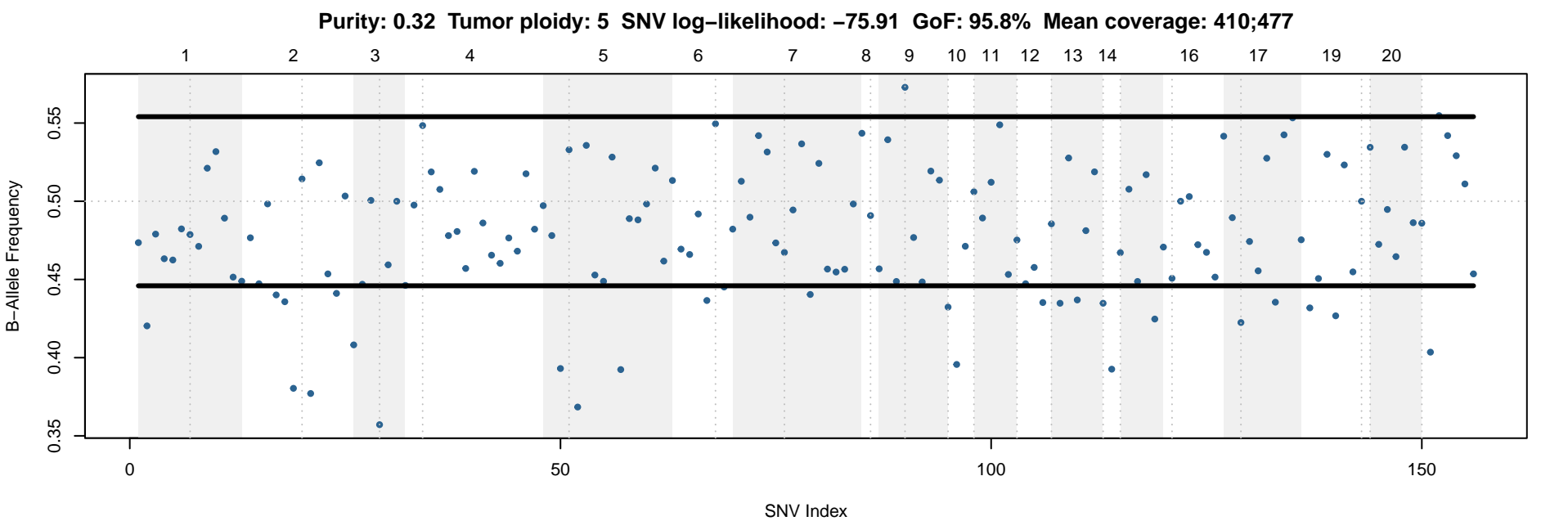
SCNA-fit log-likelihood: -7178.99

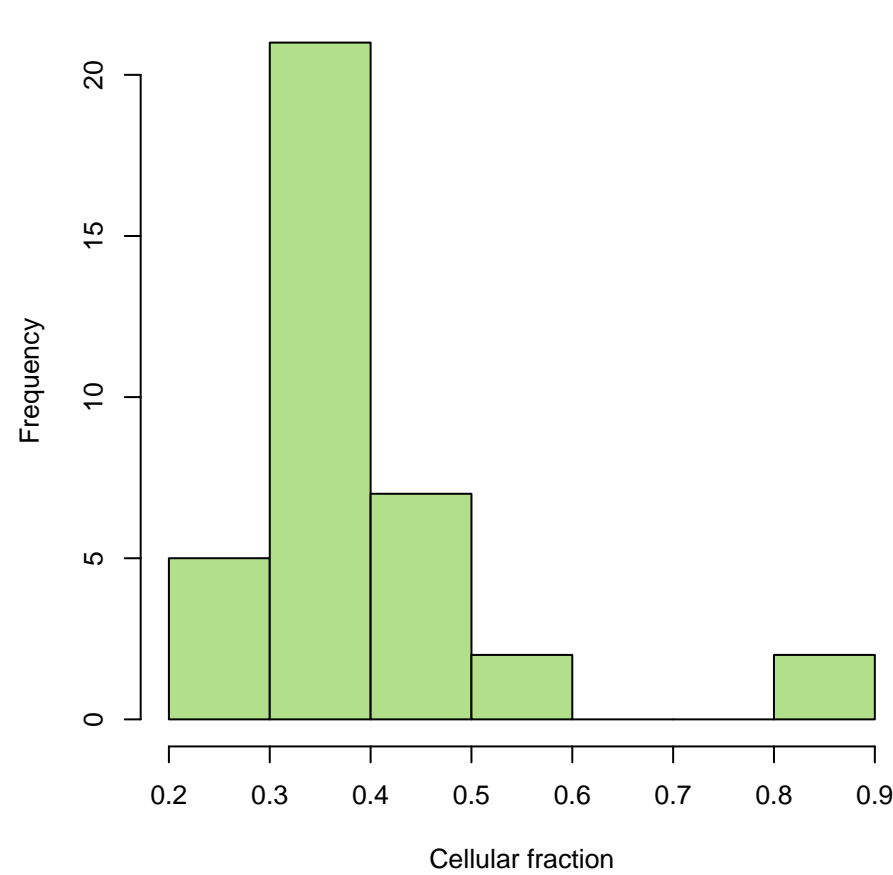
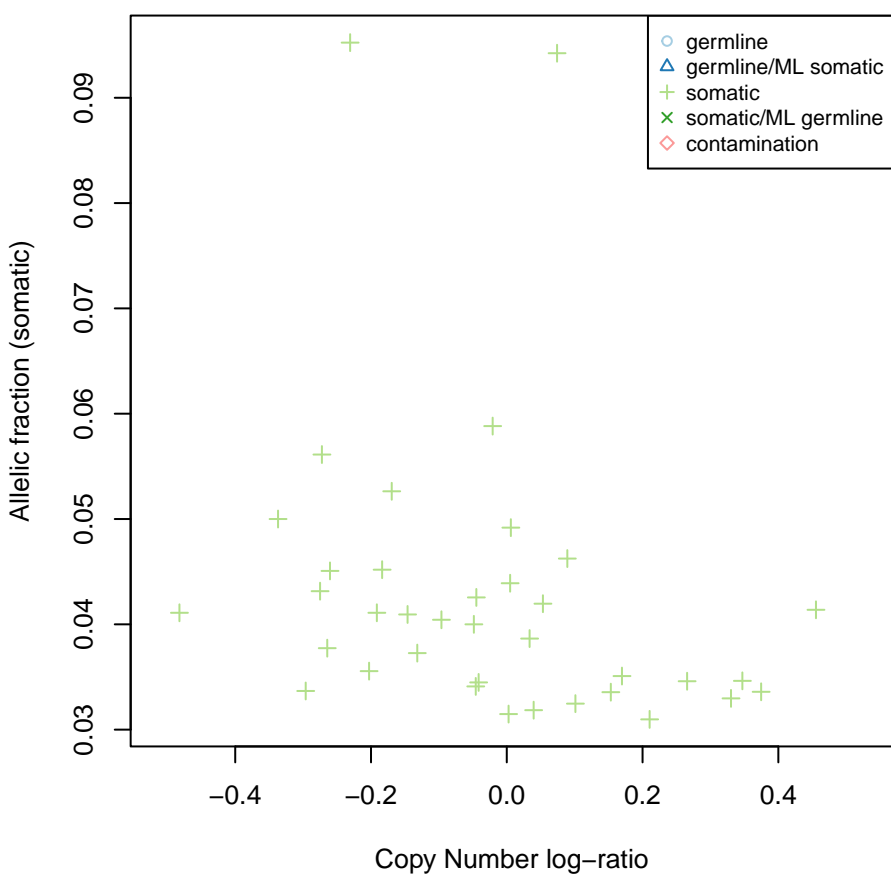
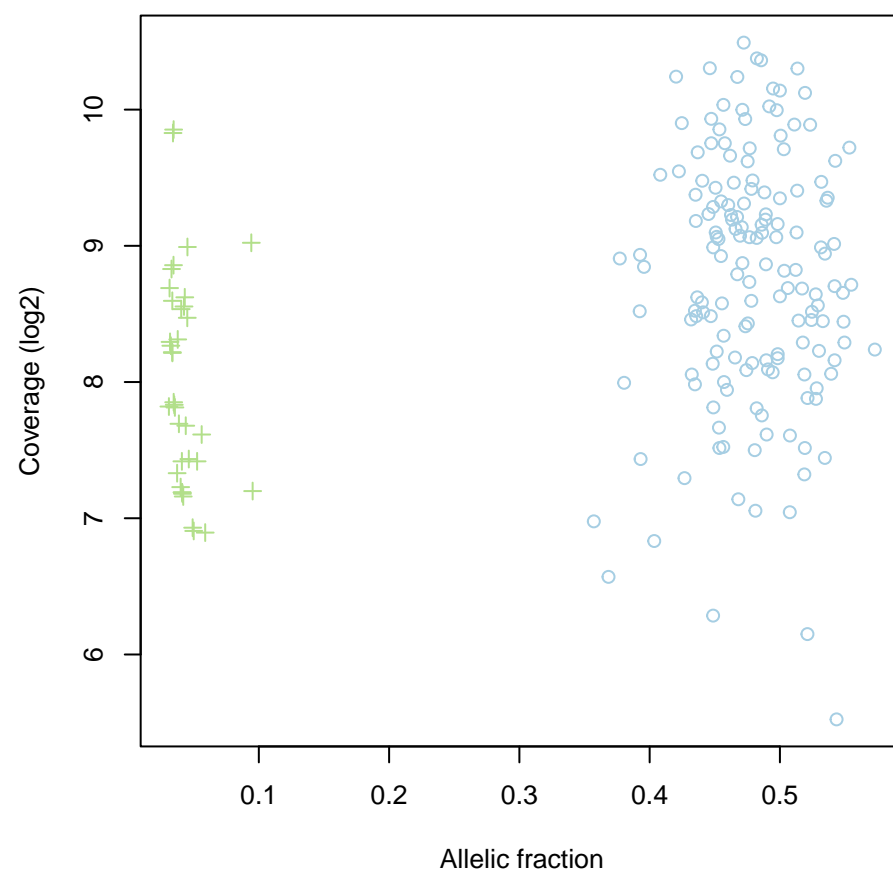
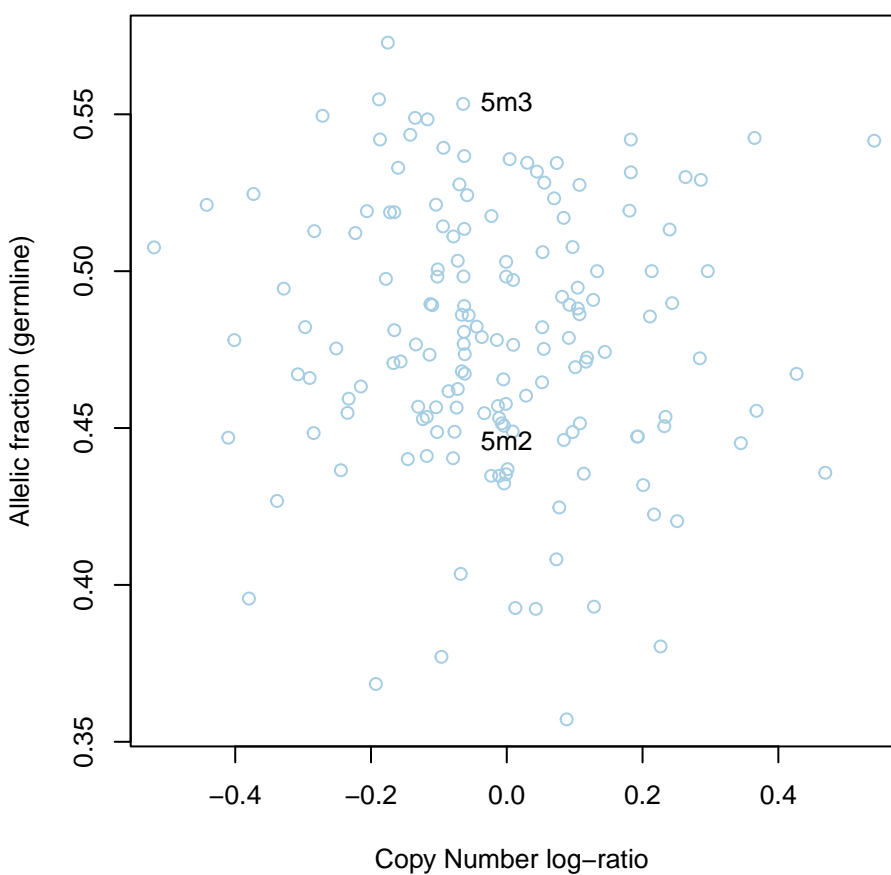




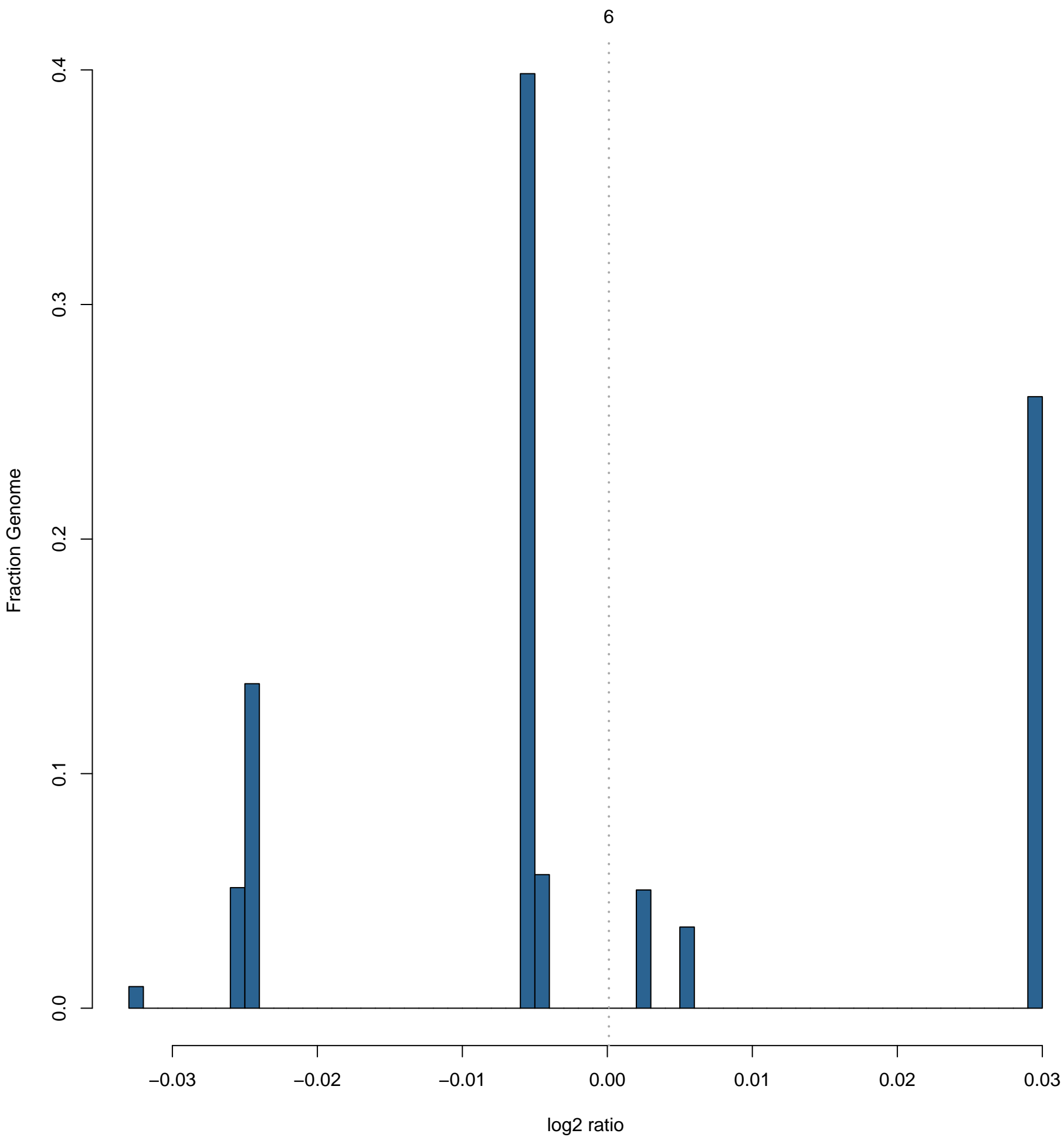
Purity: 0.32 Tumor ploidy: 5



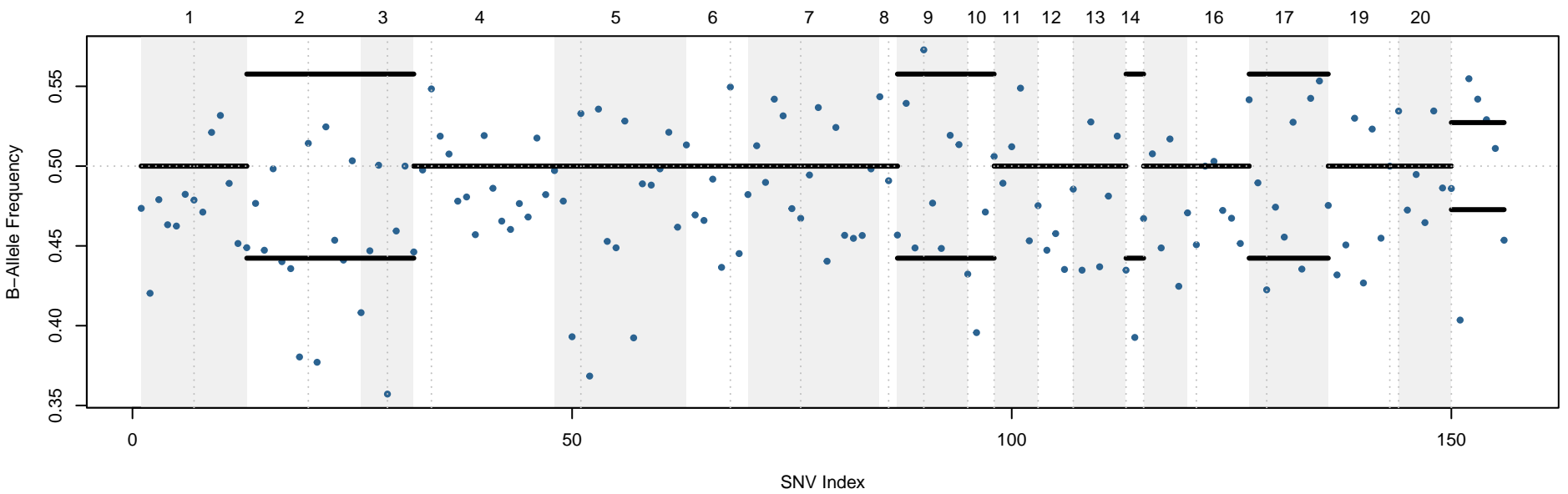




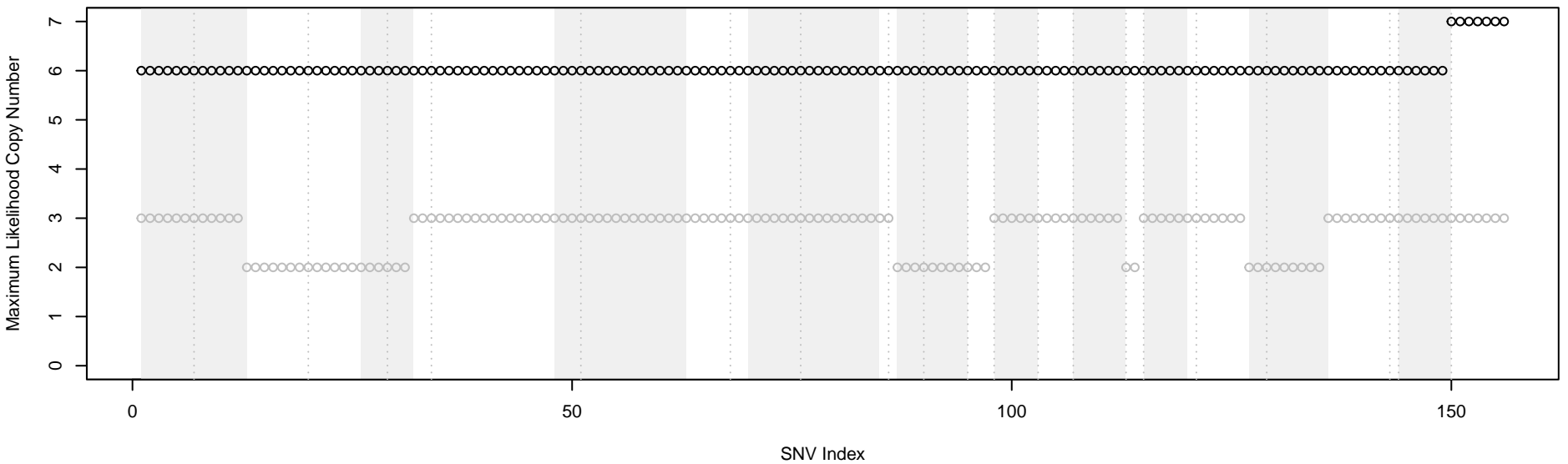
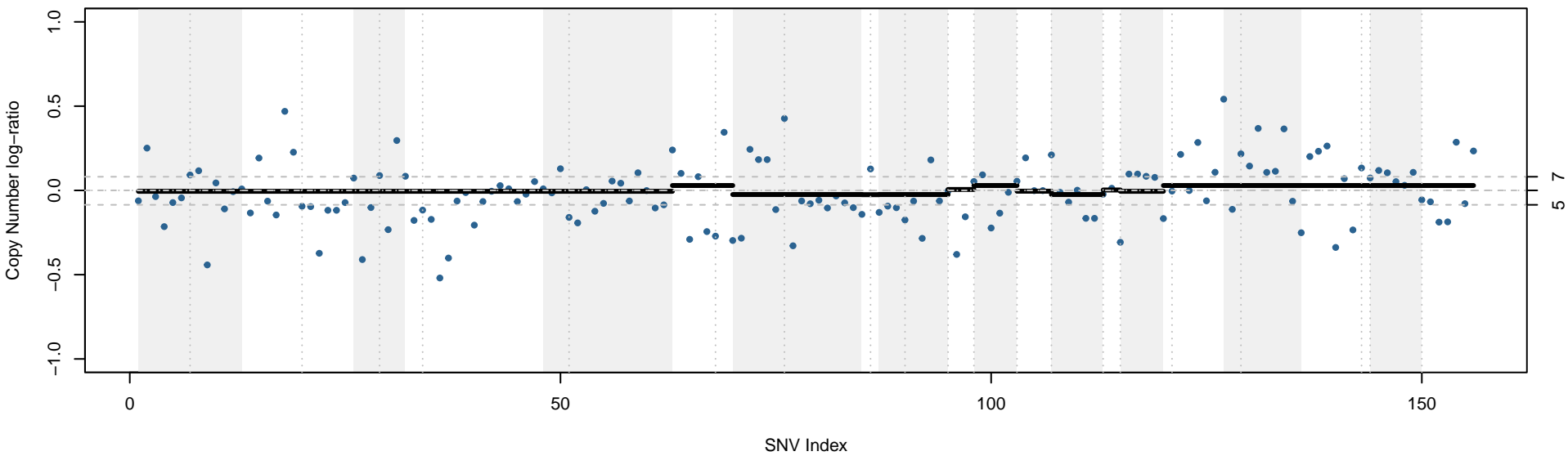
Purity: 0.15 Tumor ploidy: 5.999

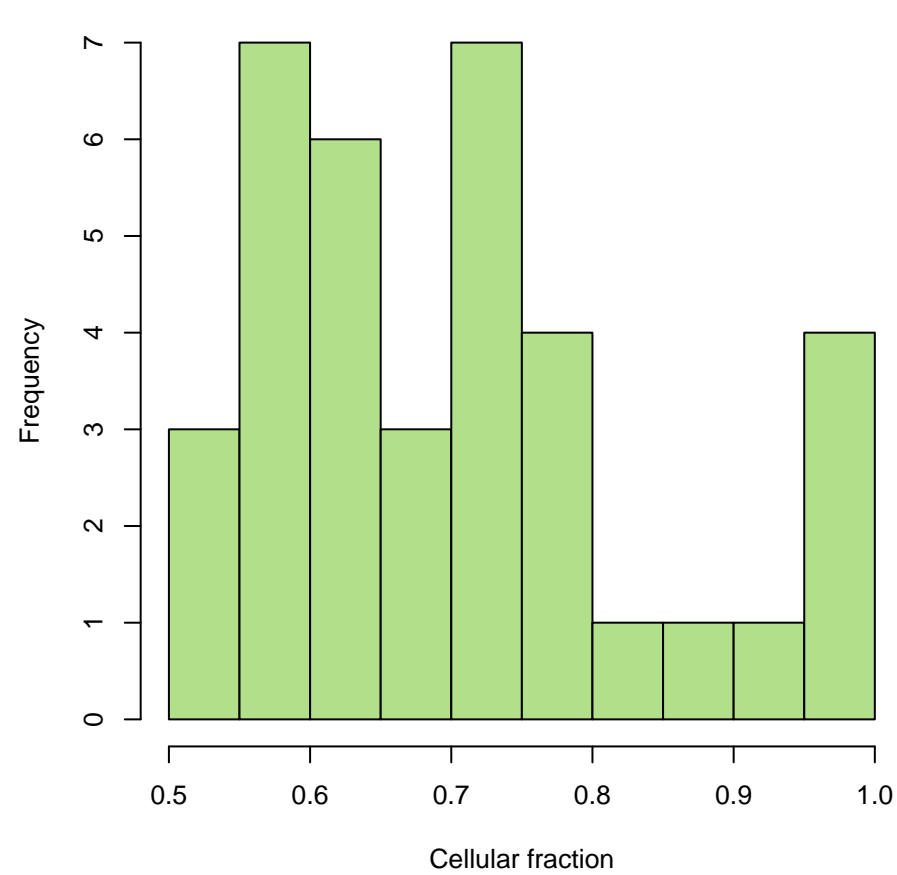
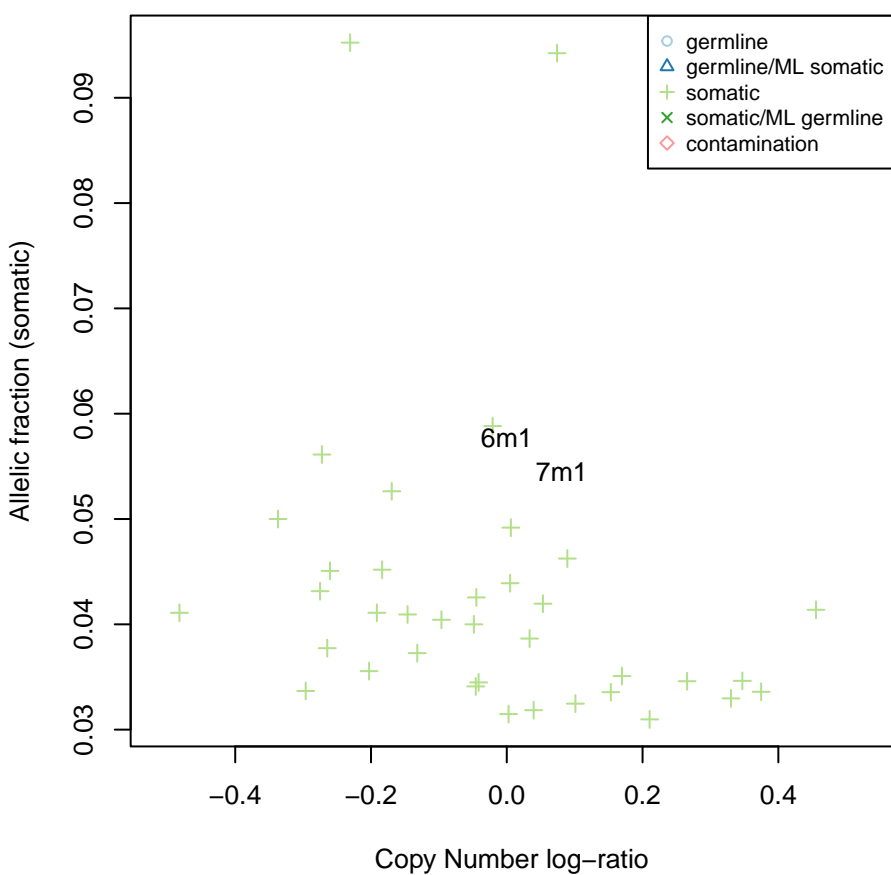
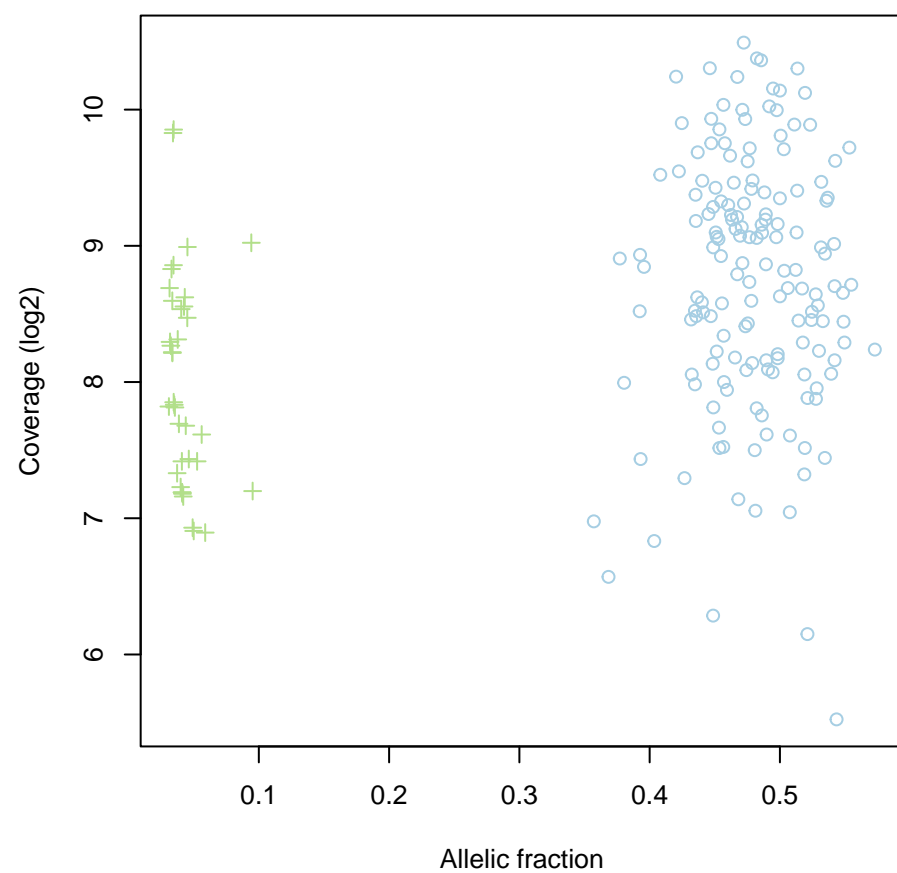
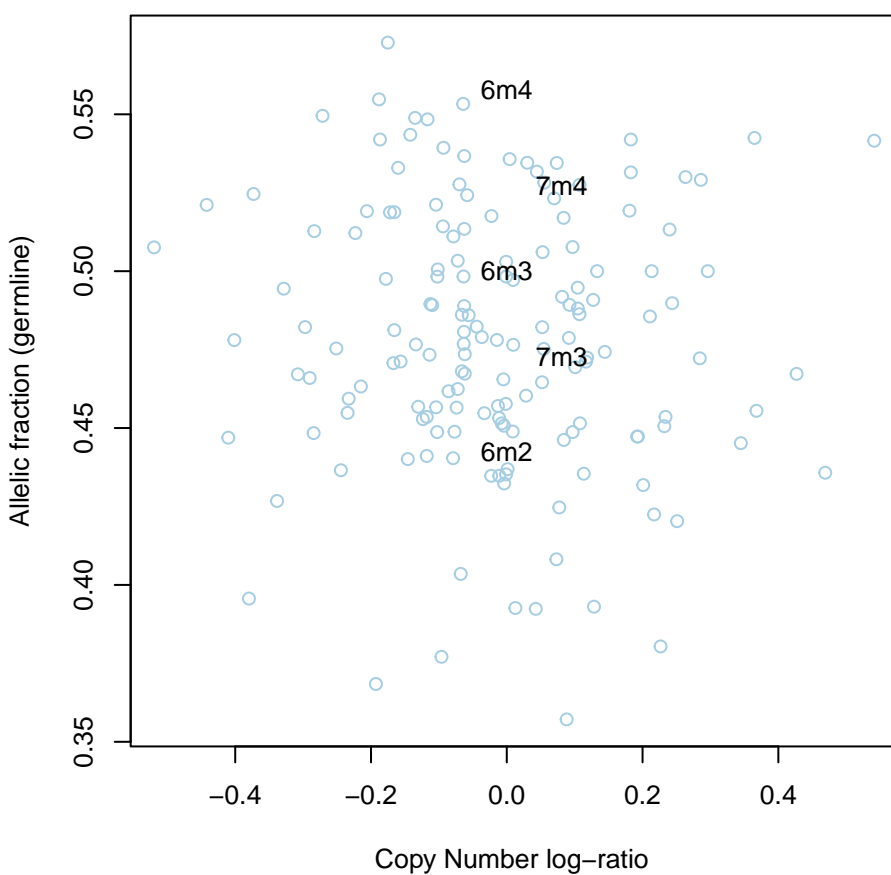


Purity: 0.15 Tumor ploidy: 5.999 SNV log-likelihood: -61.31 GoF: 97.2% Mean coverage: 410;477



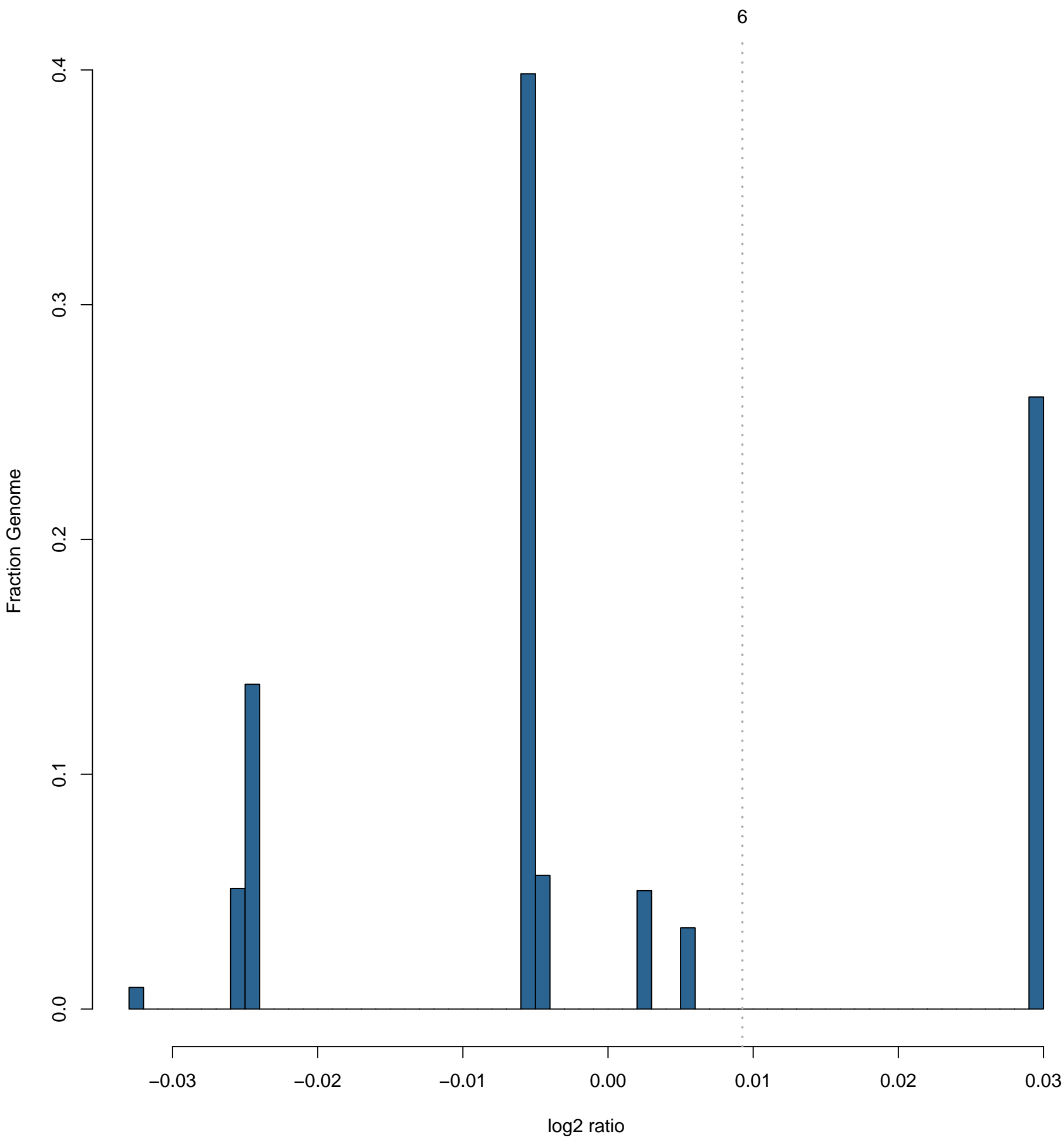
SCNA-fit log-likelihood: -7173.43



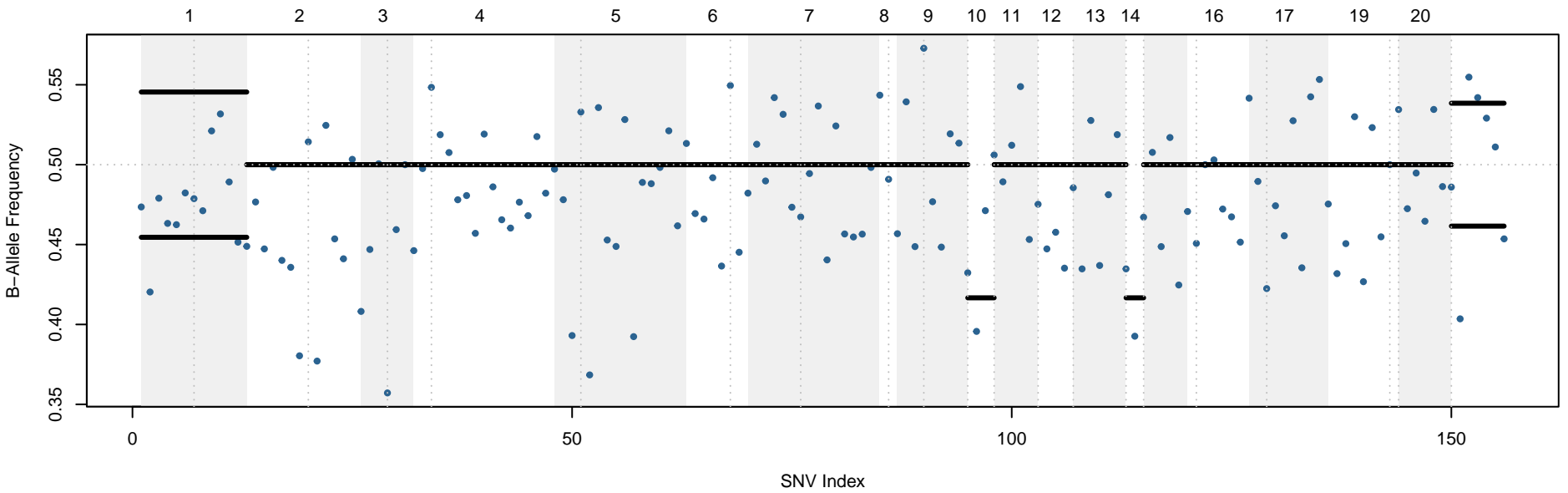




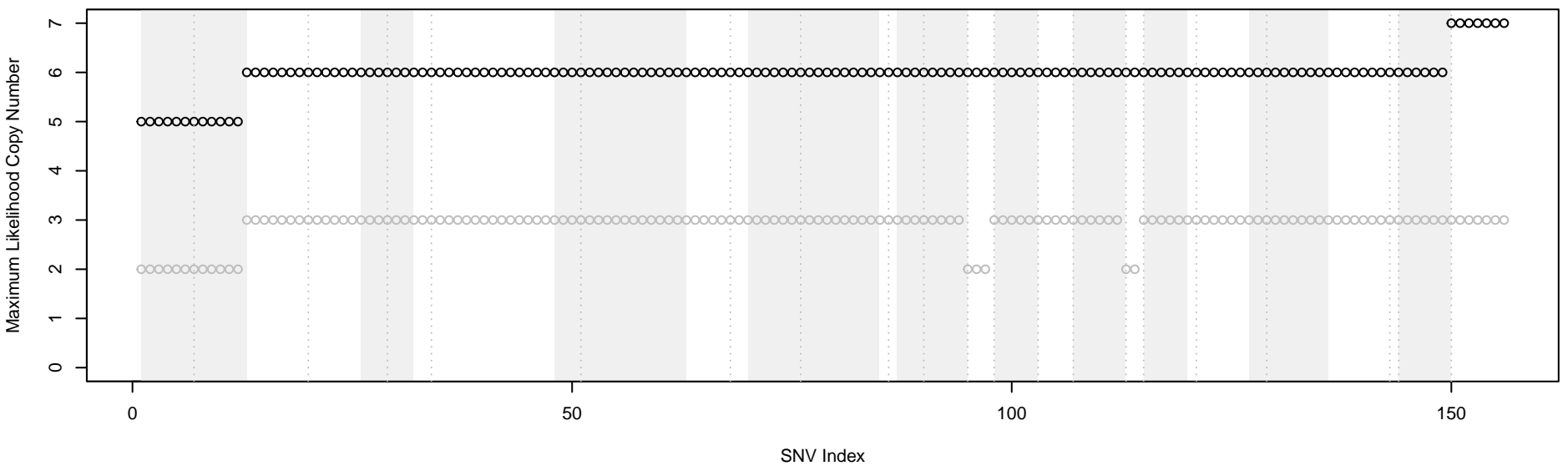
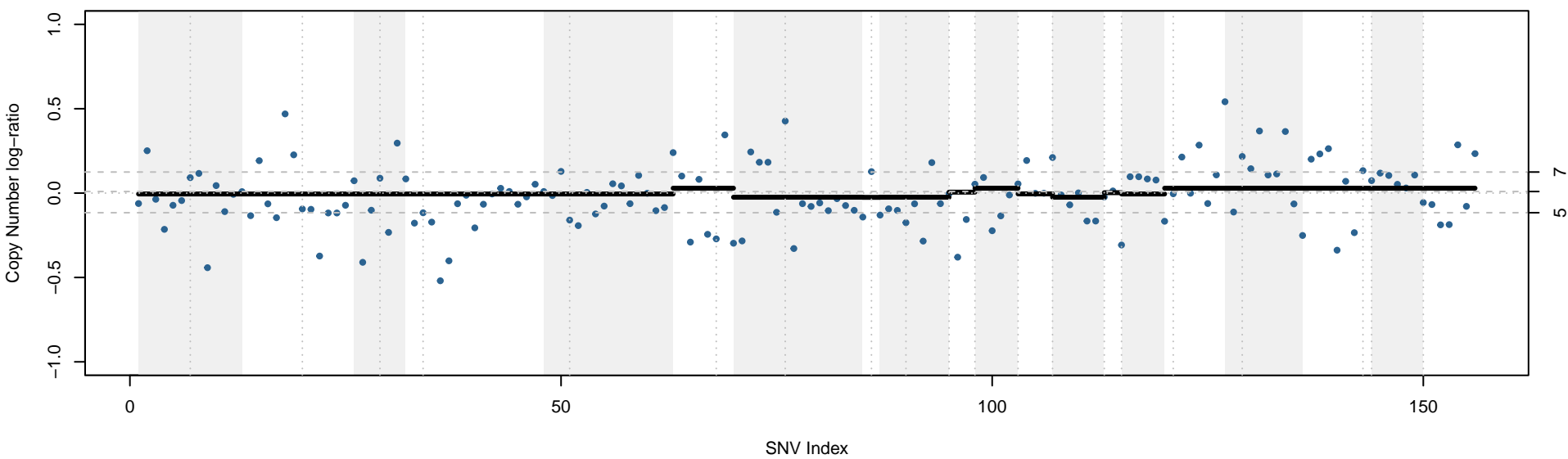
Purity: 0.25 Tumor ploidy: 5.923

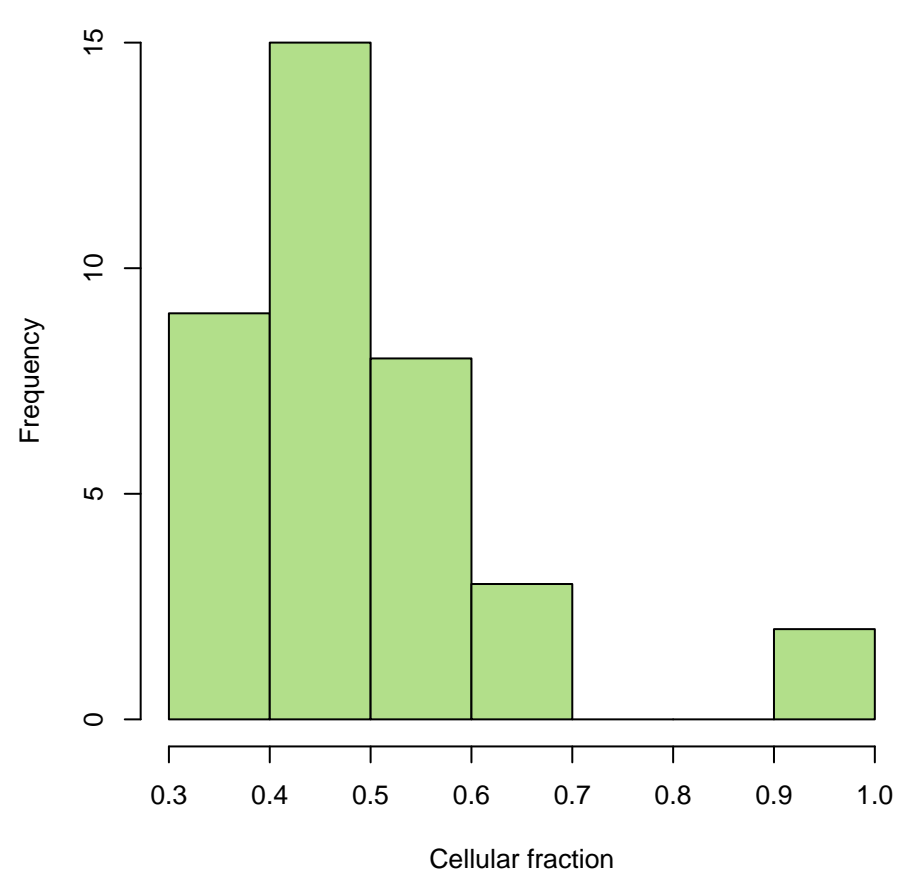
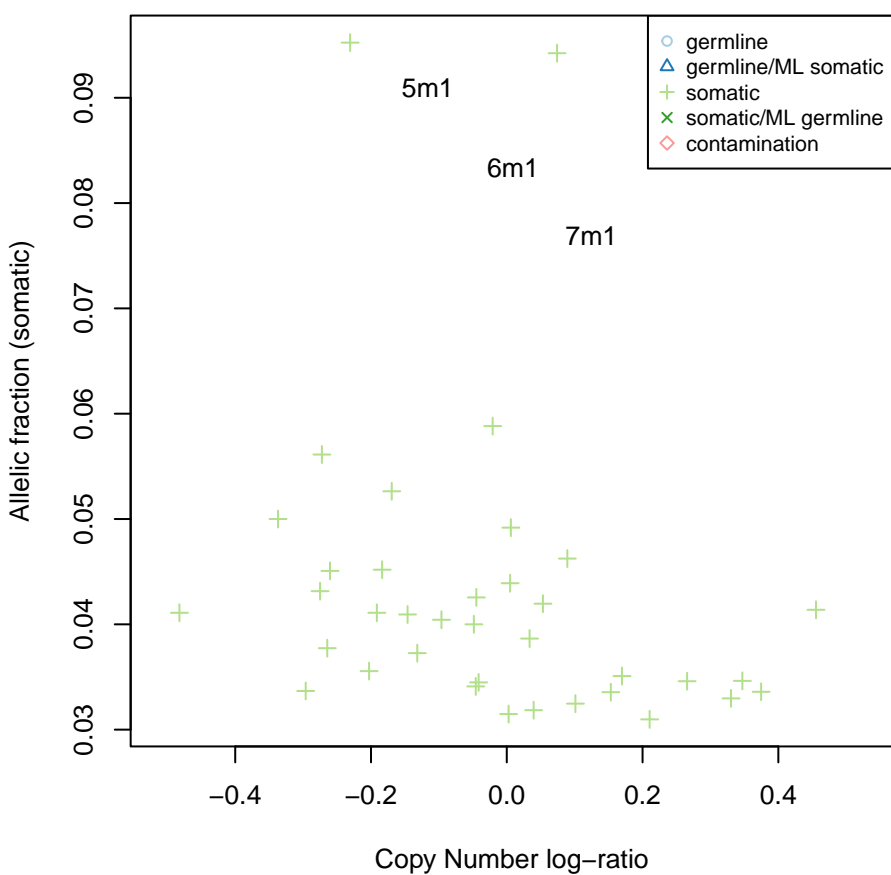
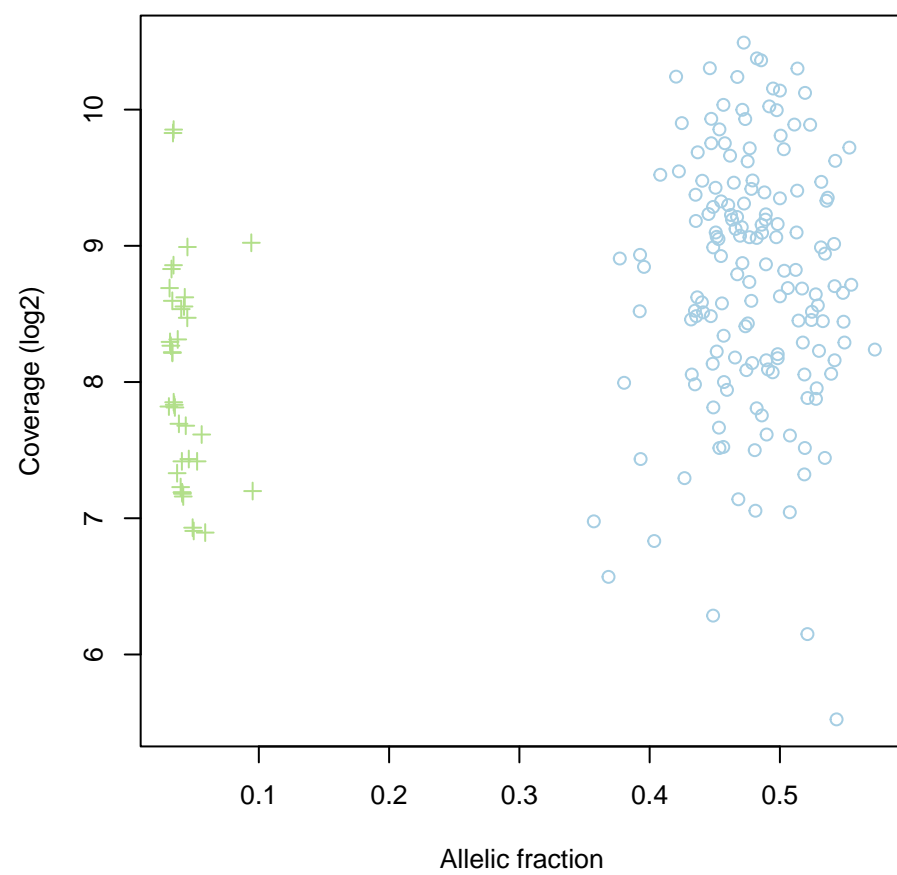
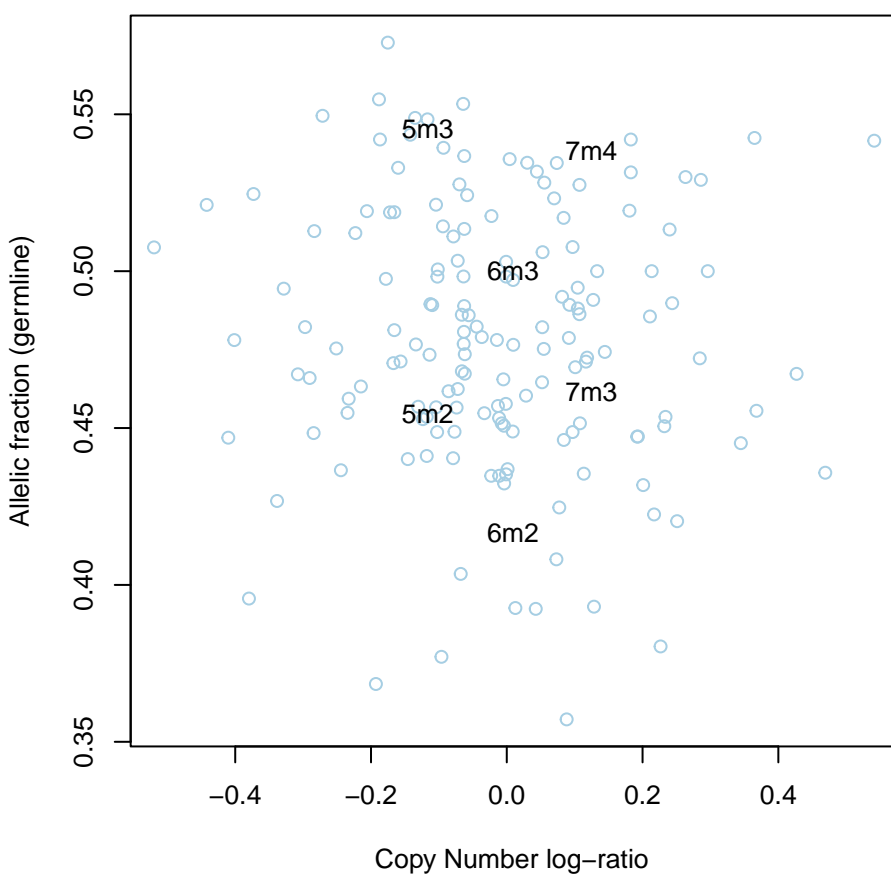


Purity: 0.25 Tumor ploidy: 5.923 SNV log-likelihood: -66.18 GoF: 95.7% Mean coverage: 410;477

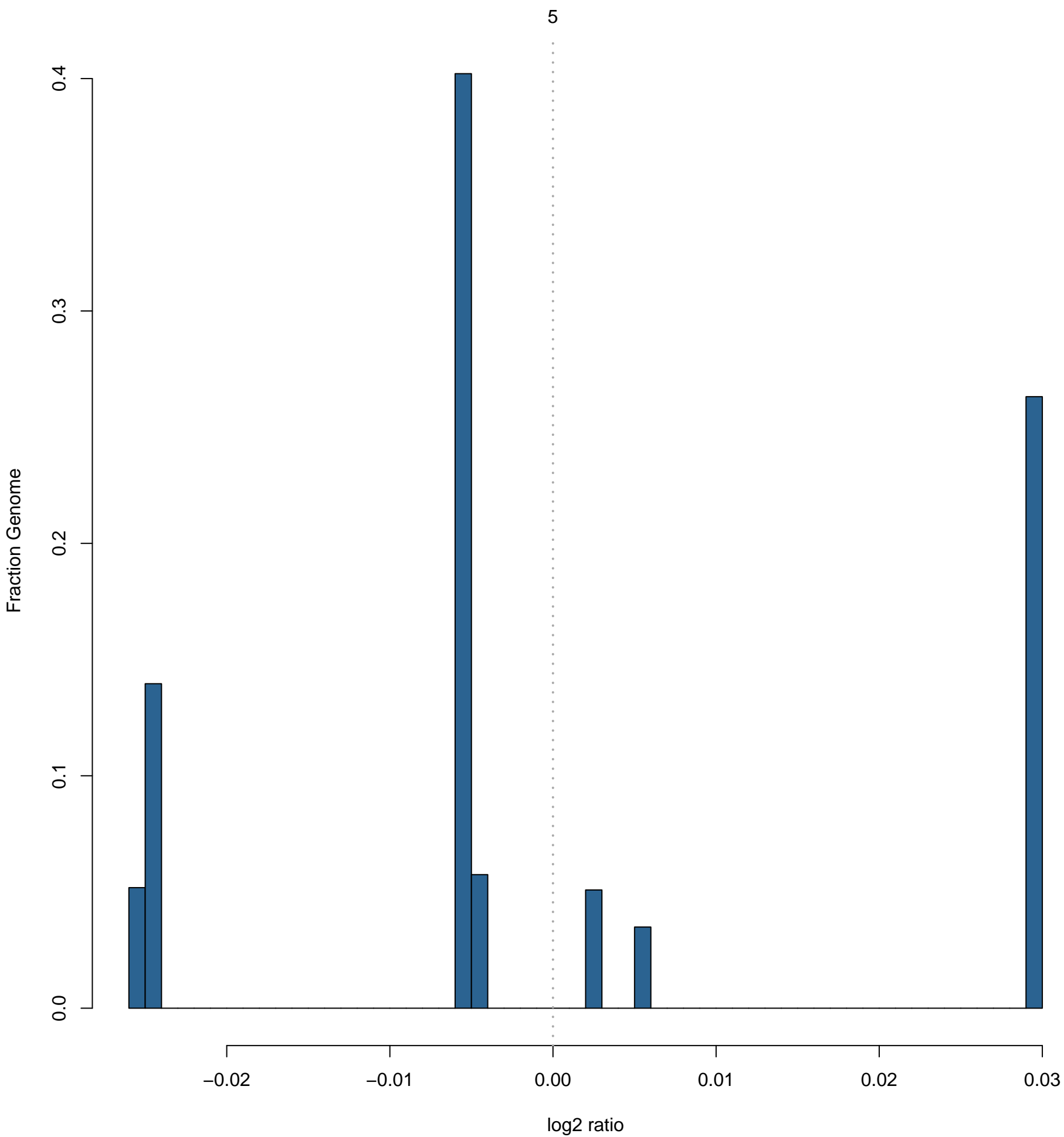


SCNA-fit log-likelihood: -7261.85

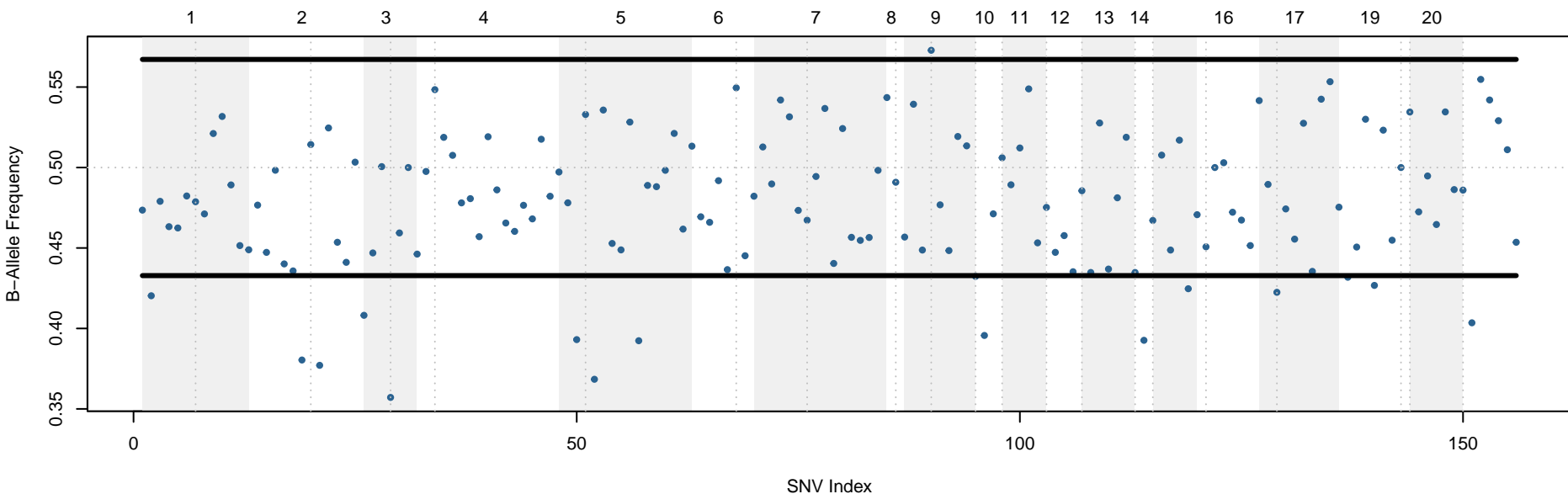




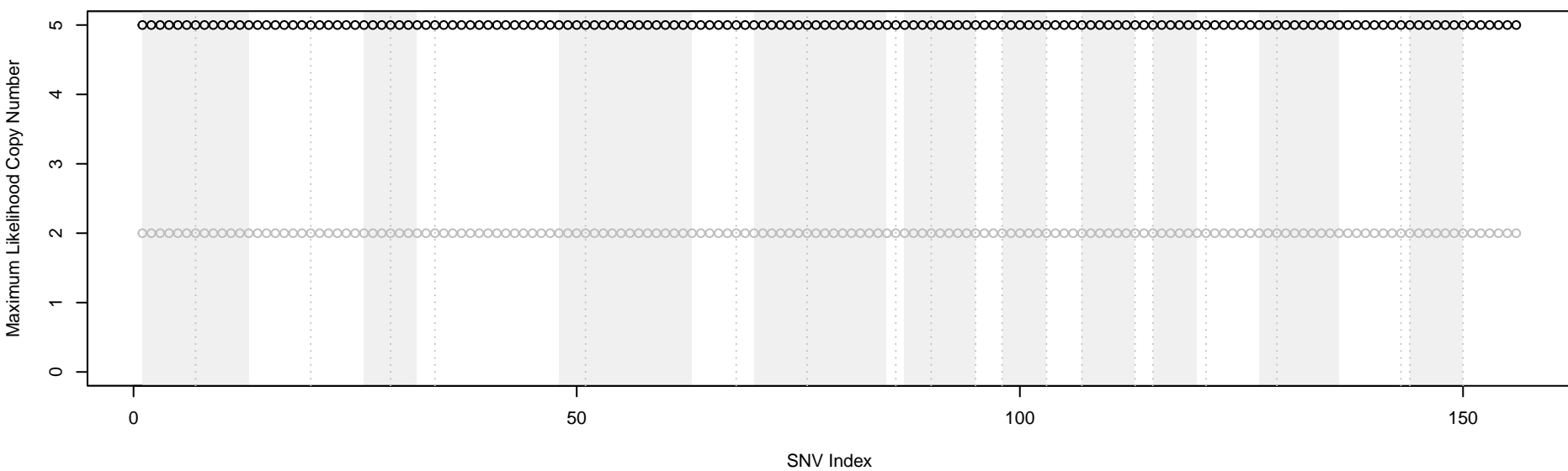
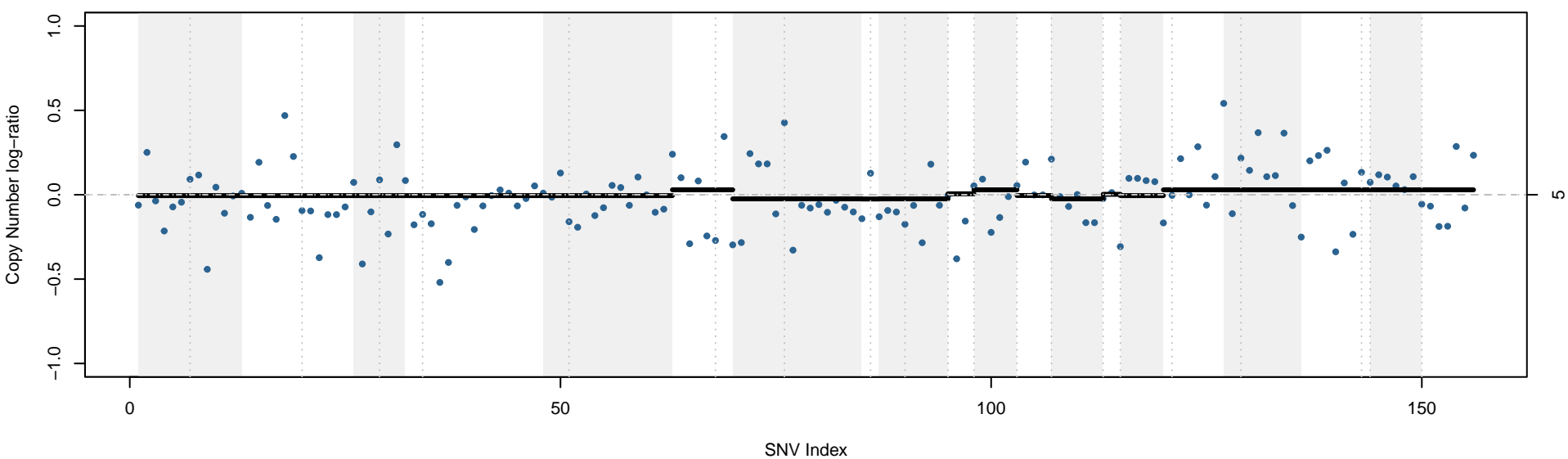
Purity: 0.45 Tumor ploidy: 5

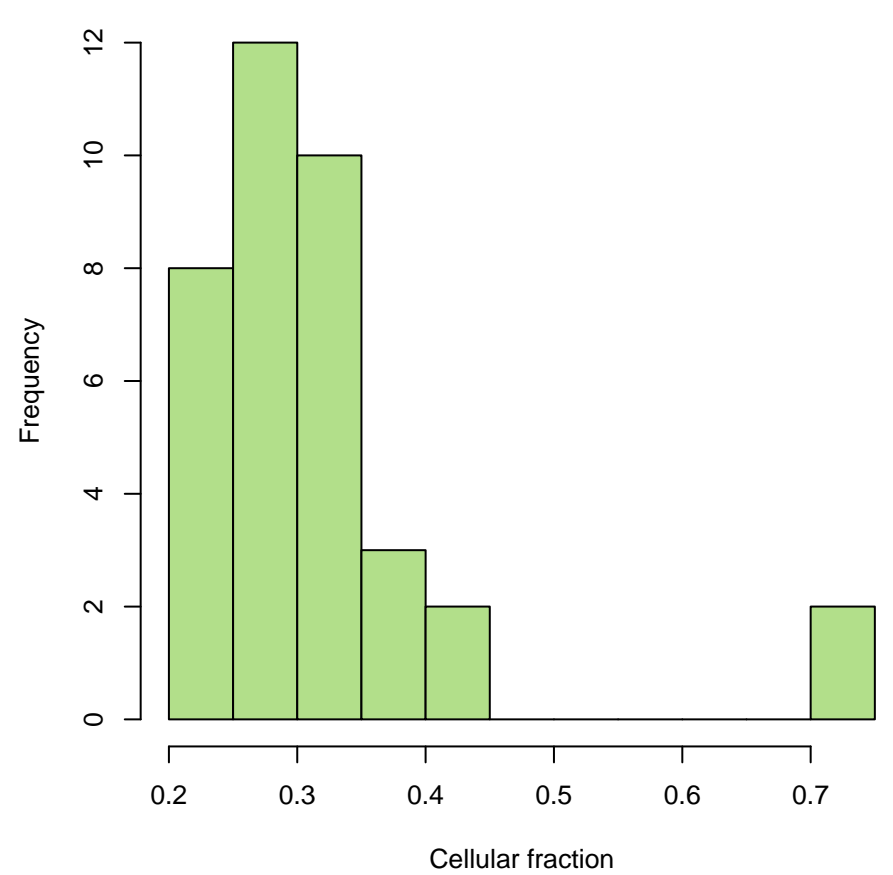
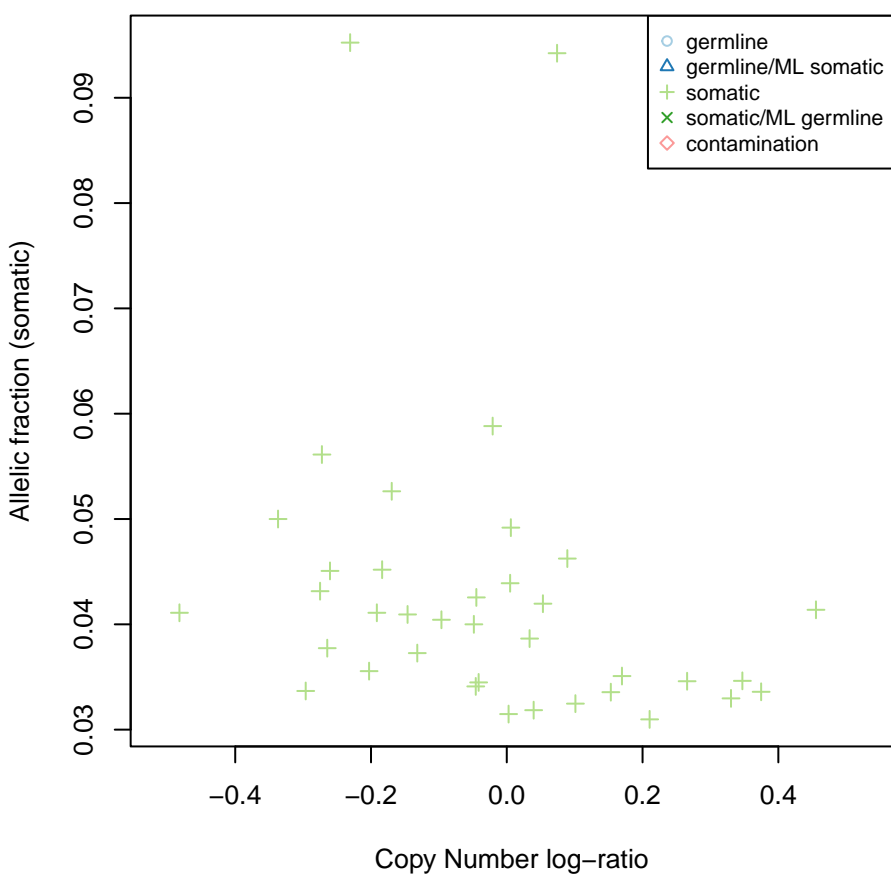
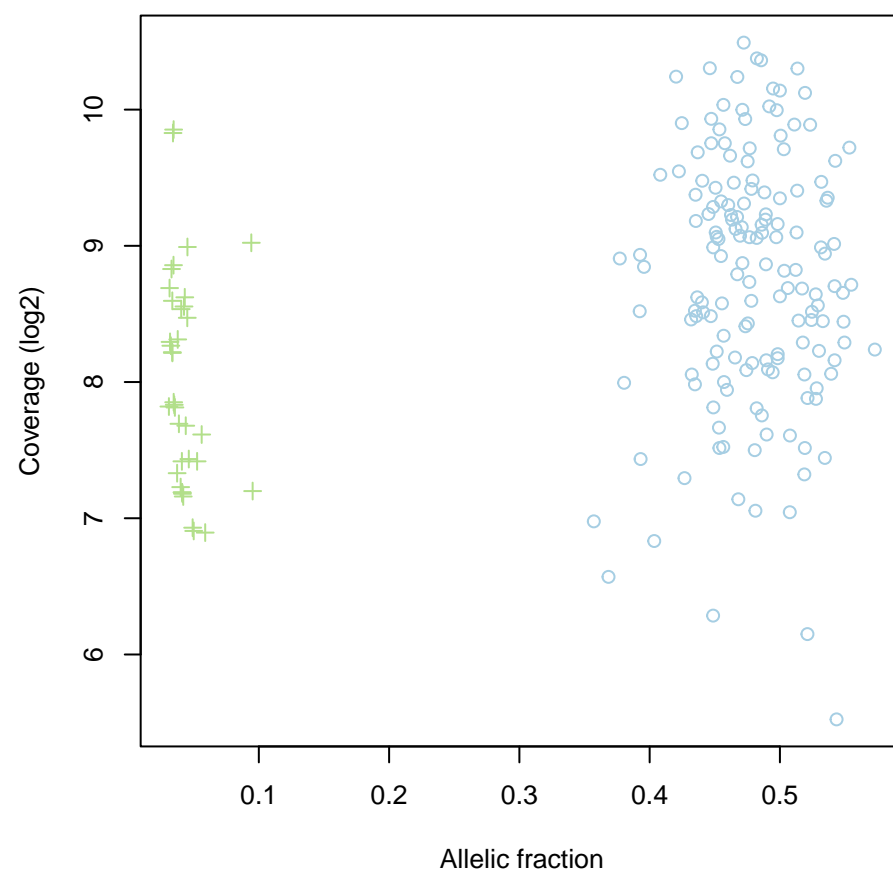
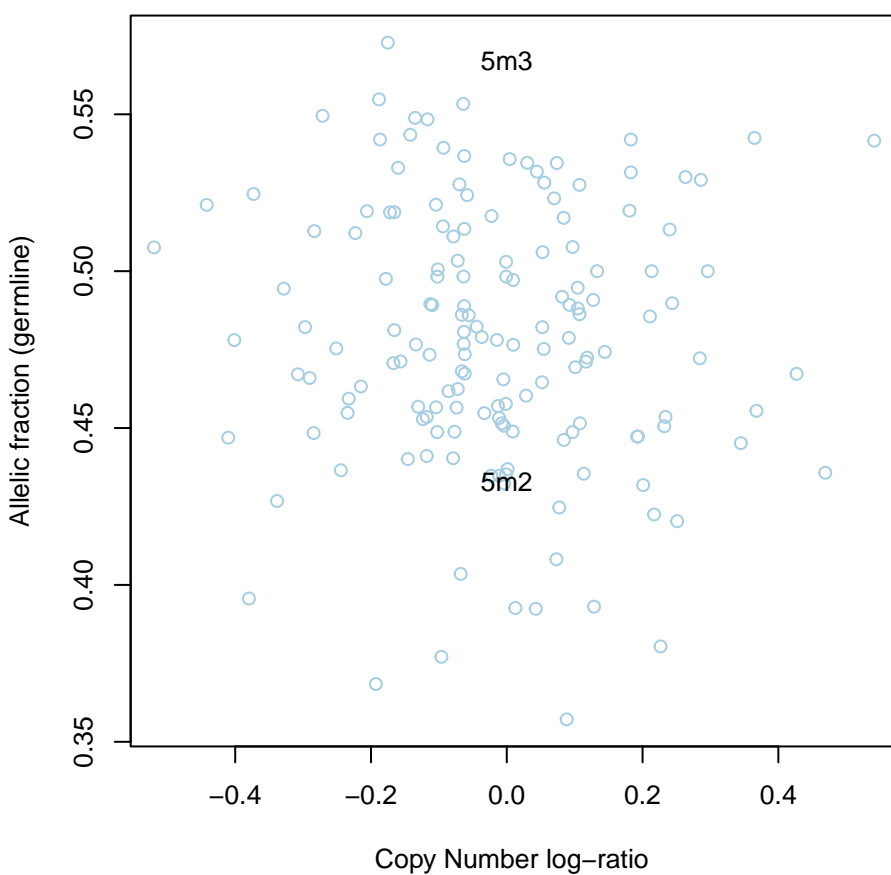


Purity: 0.45 Tumor ploidy: 5 SNV log-likelihood: -141.31 GoF: 92.6% Mean coverage: 410;477

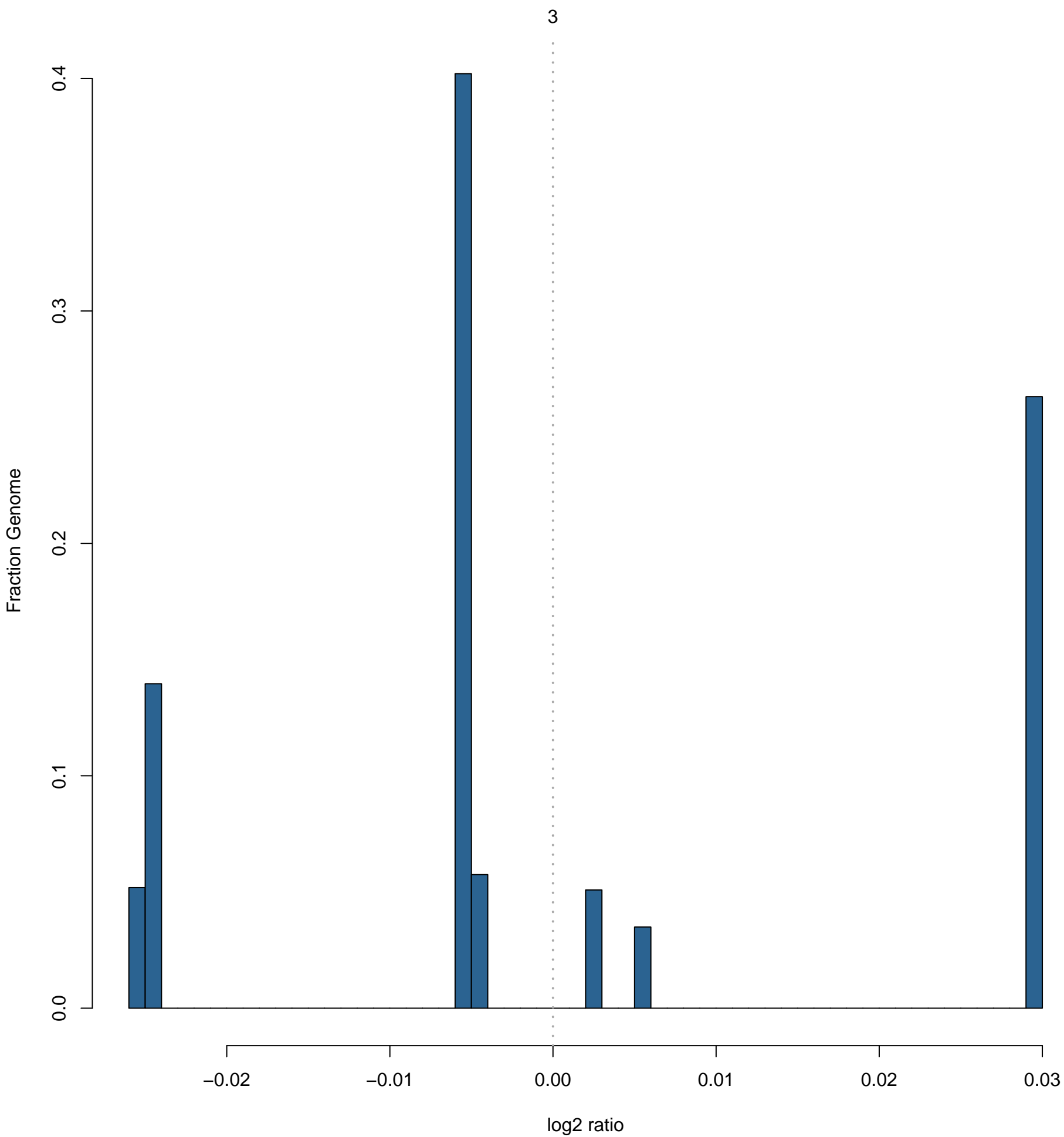


SCNA-fit log-likelihood: -7182.06

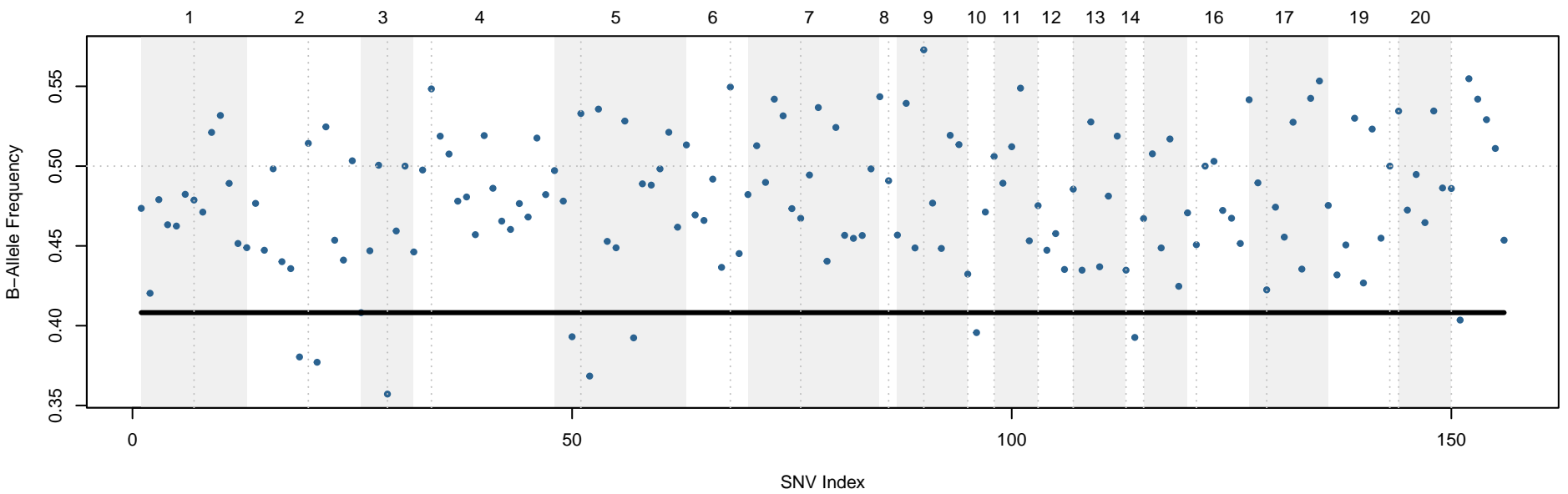




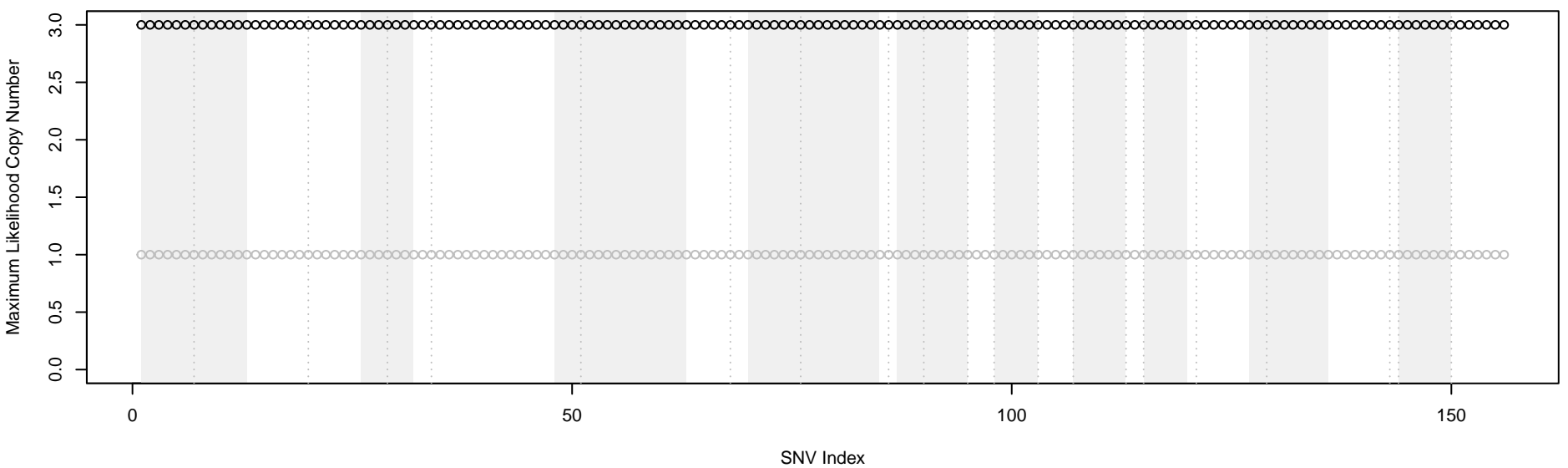
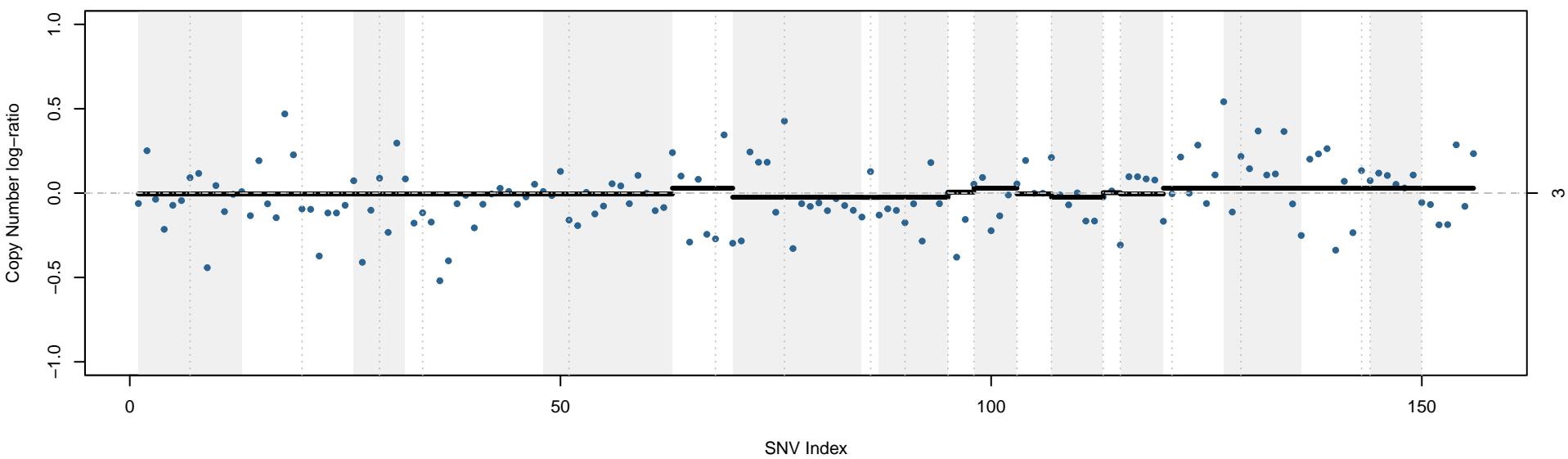
Purity: 0.45 Tumor ploidy: 3



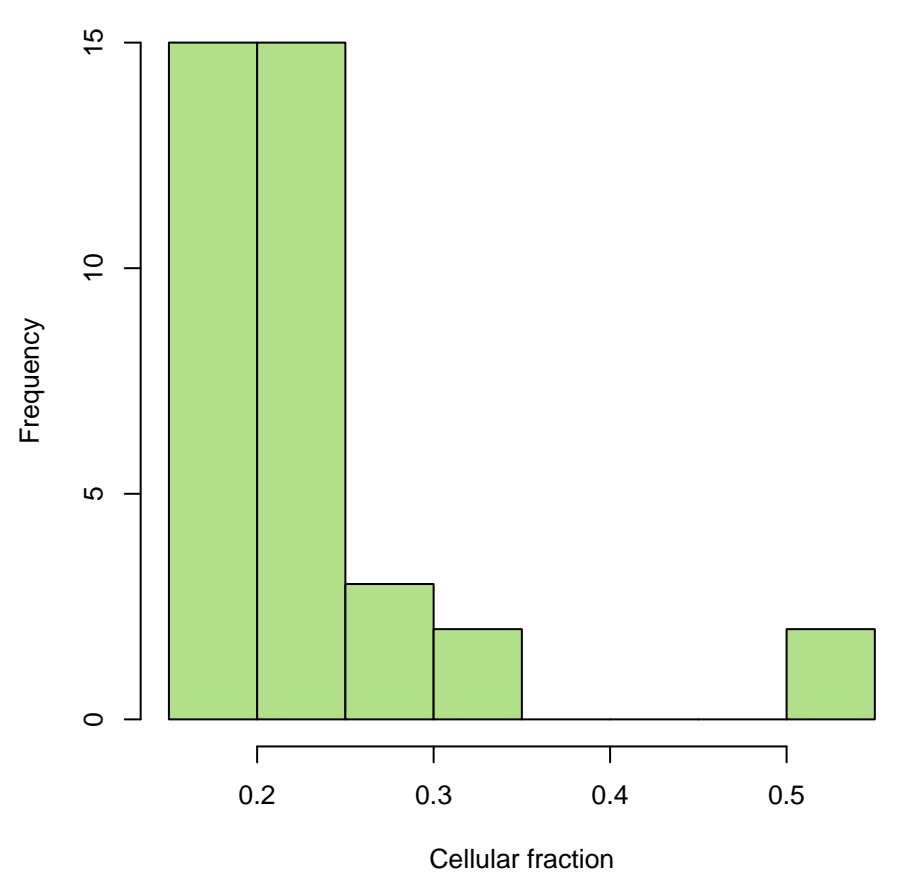
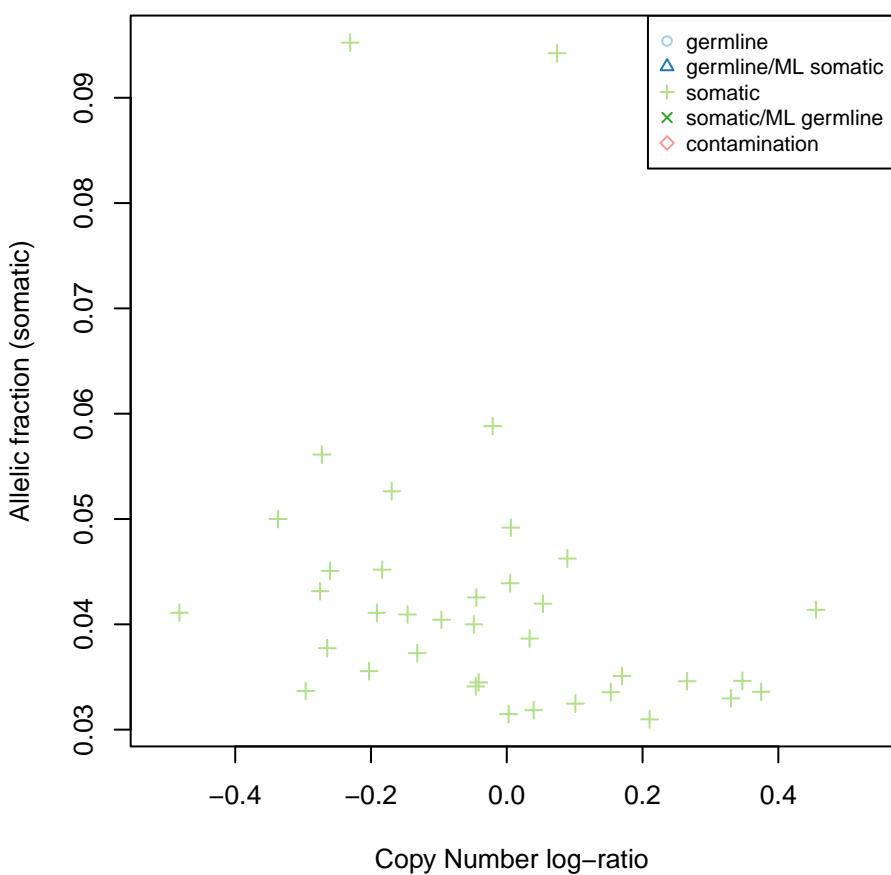
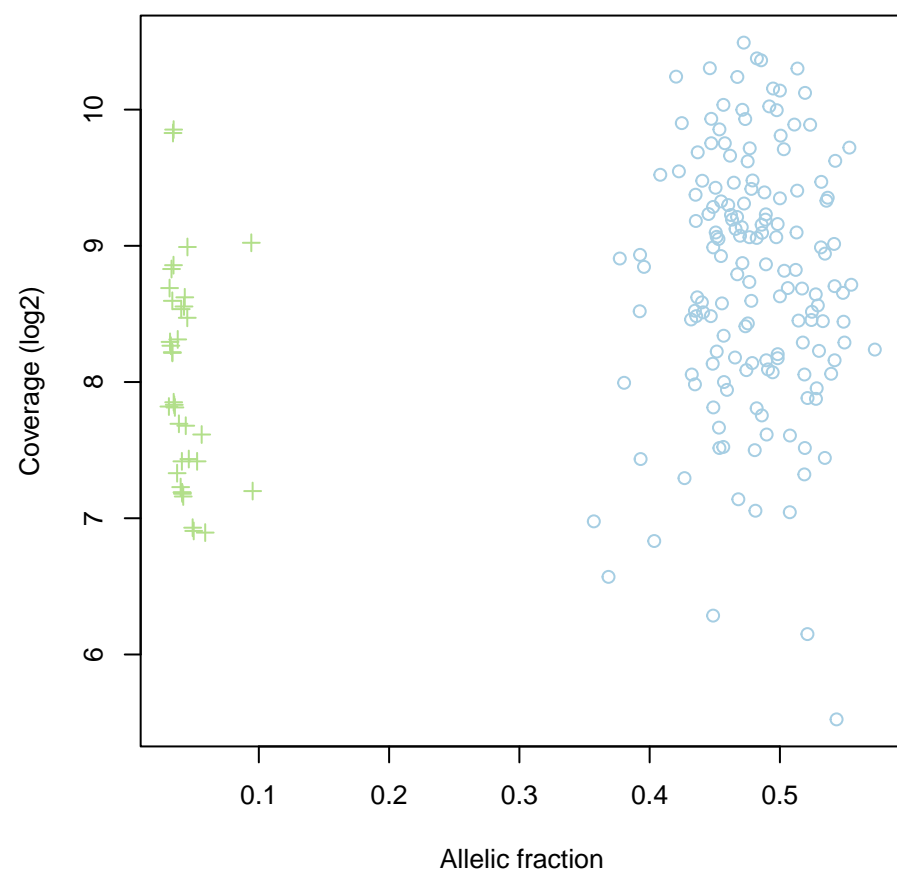
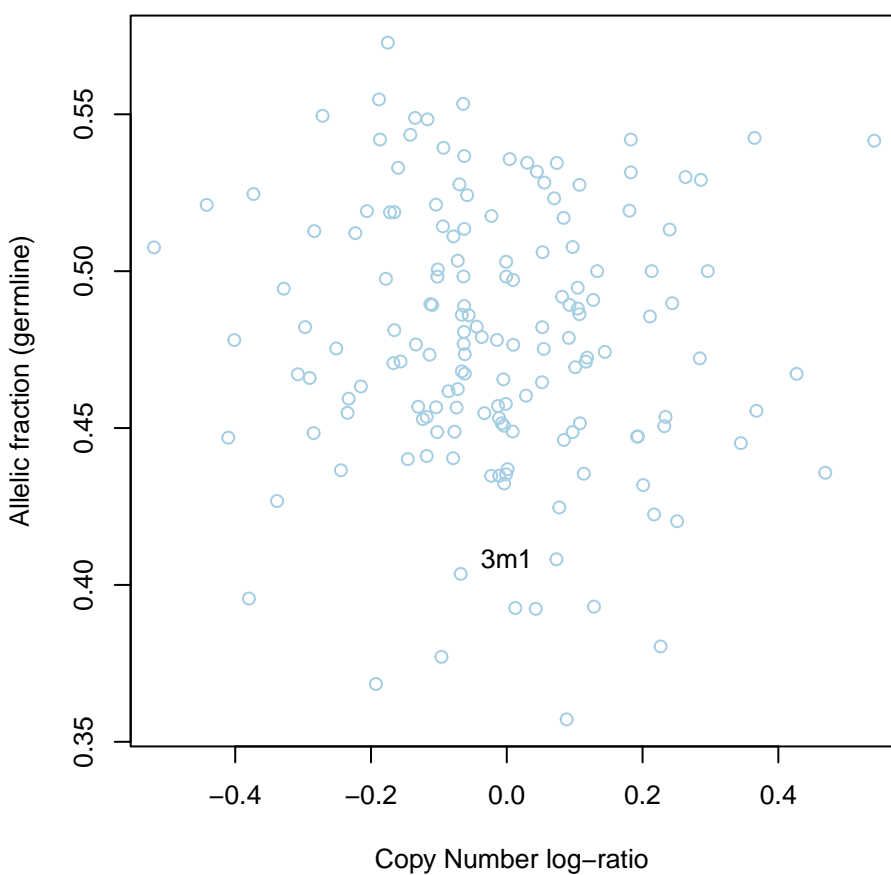
Purity: 0.45 Tumor ploidy: 3 SNV log-likelihood: -282.39 GoF: 82.7% Mean coverage: 410;477



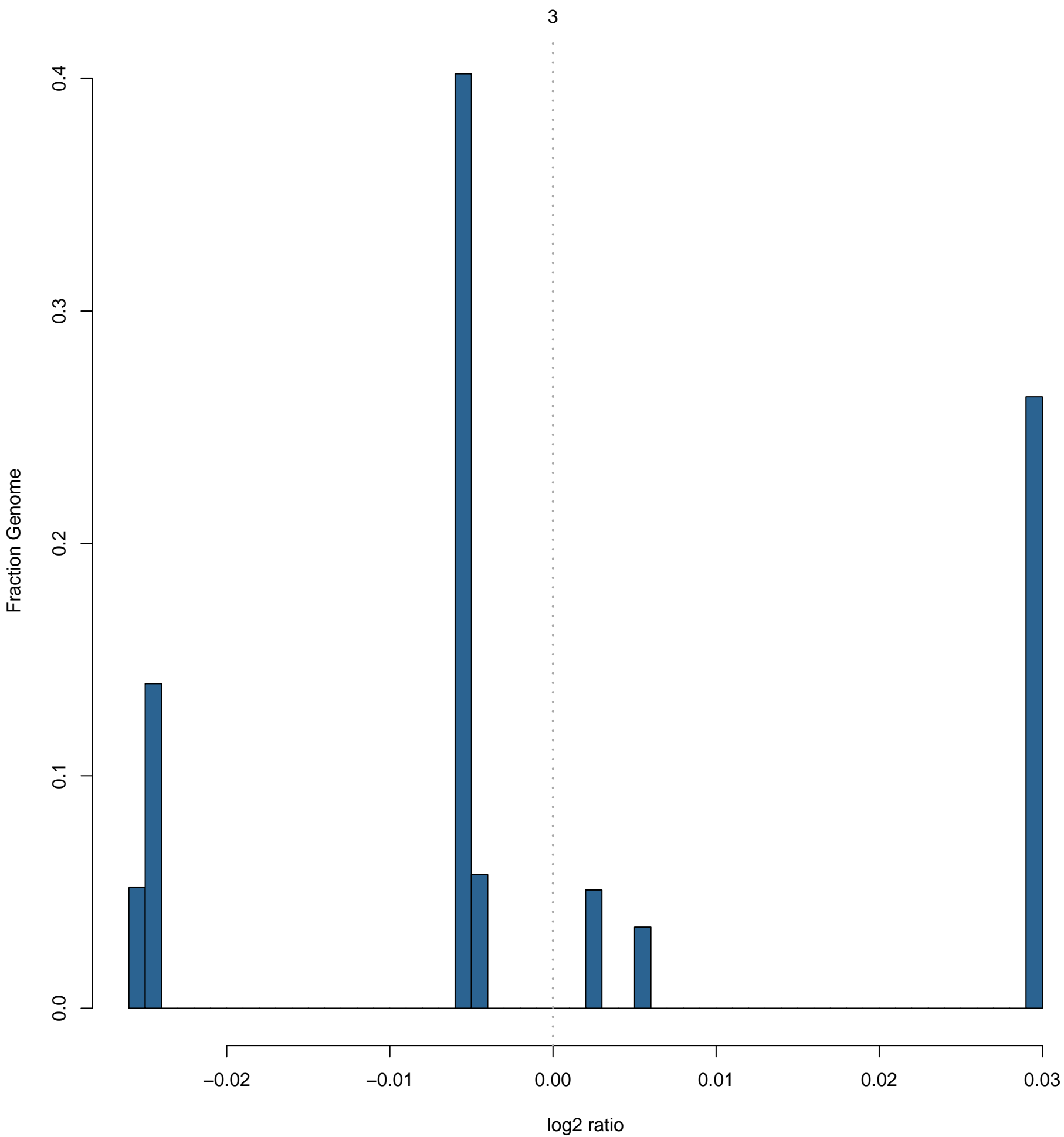
SCNA-fit log-likelihood: -7178.98



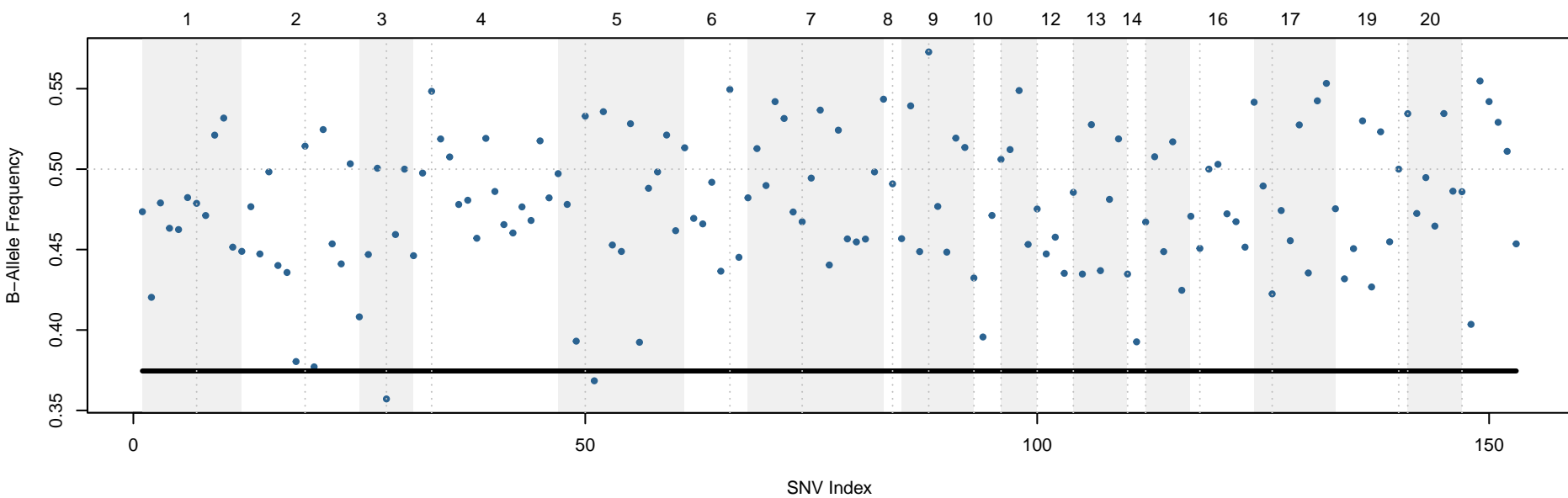




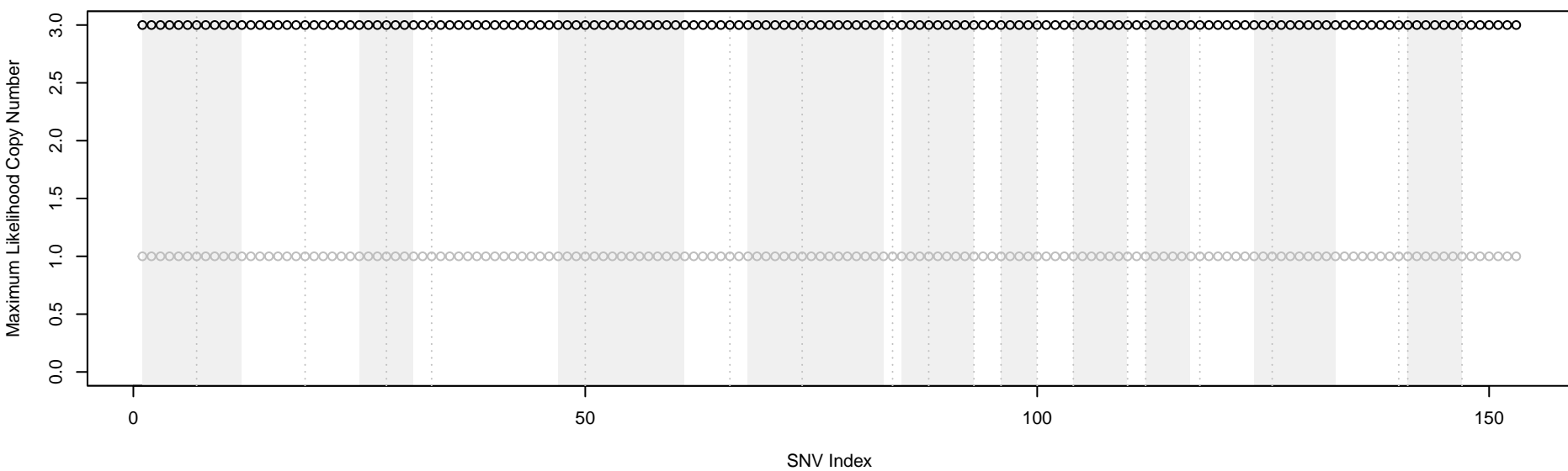
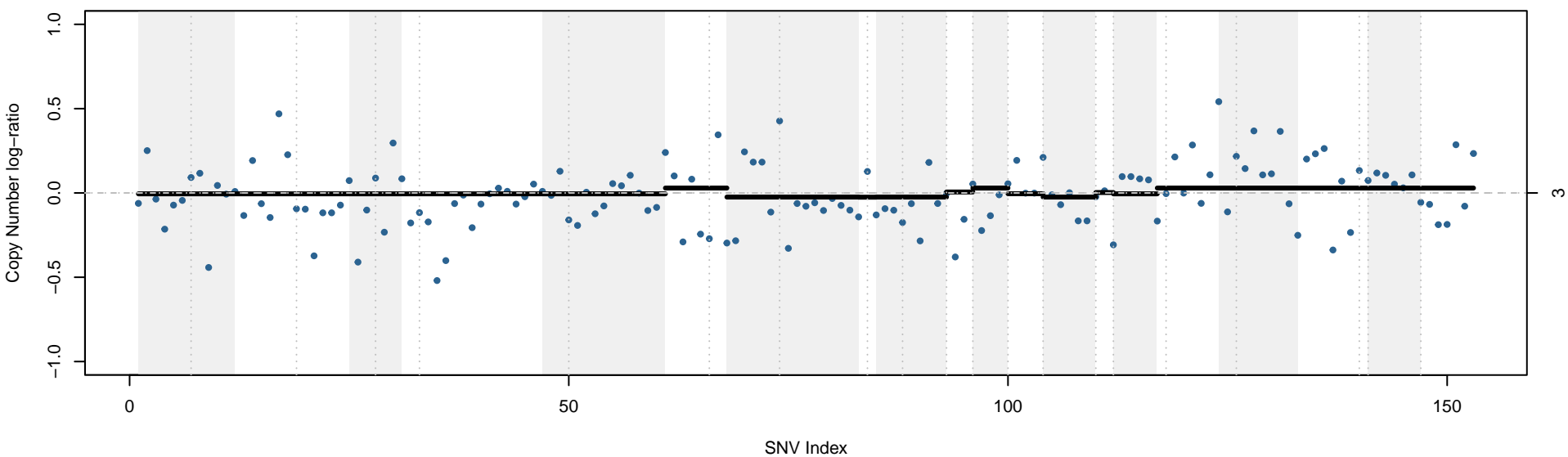
Purity: 0.67 Tumor ploidy: 3

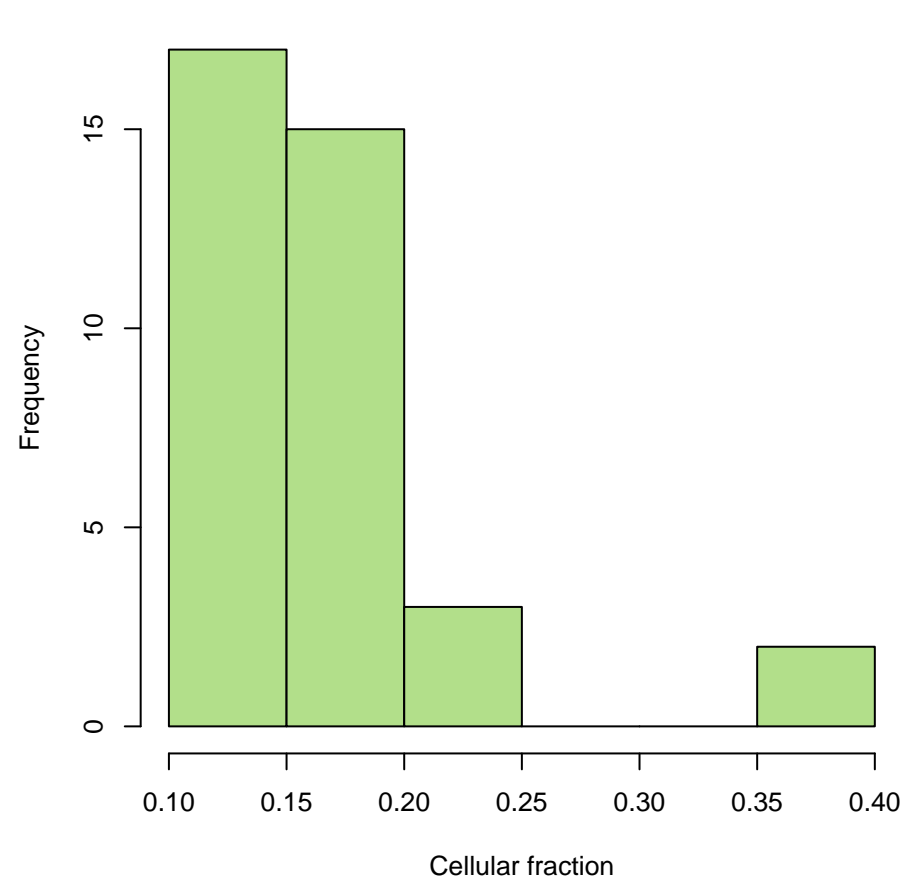
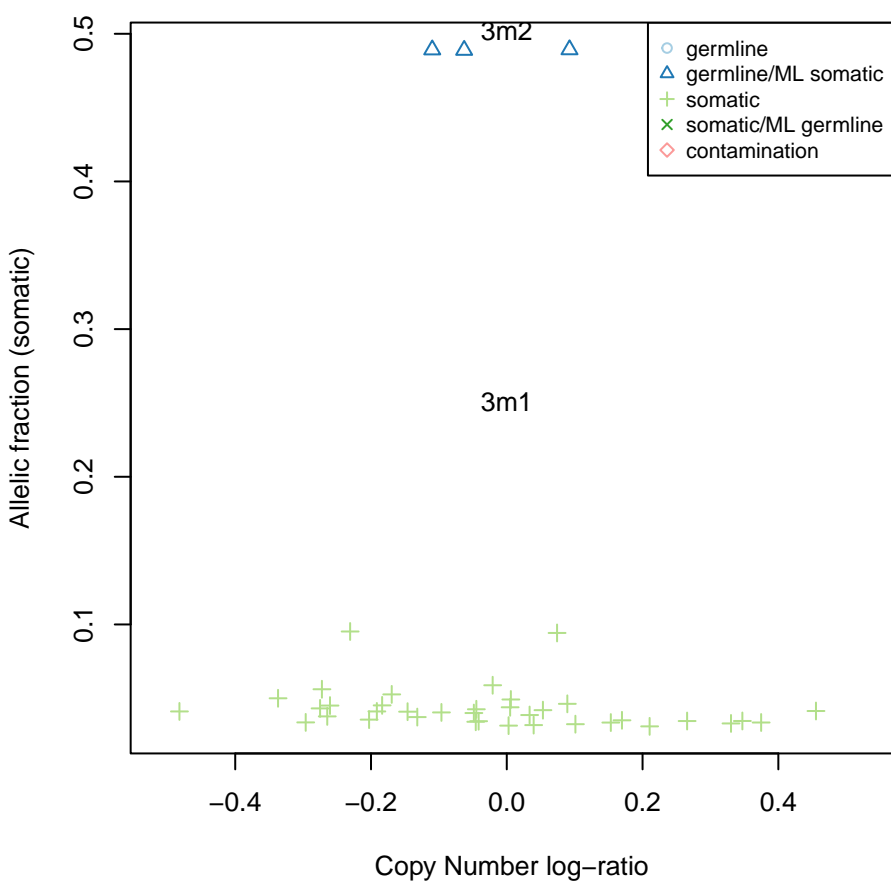
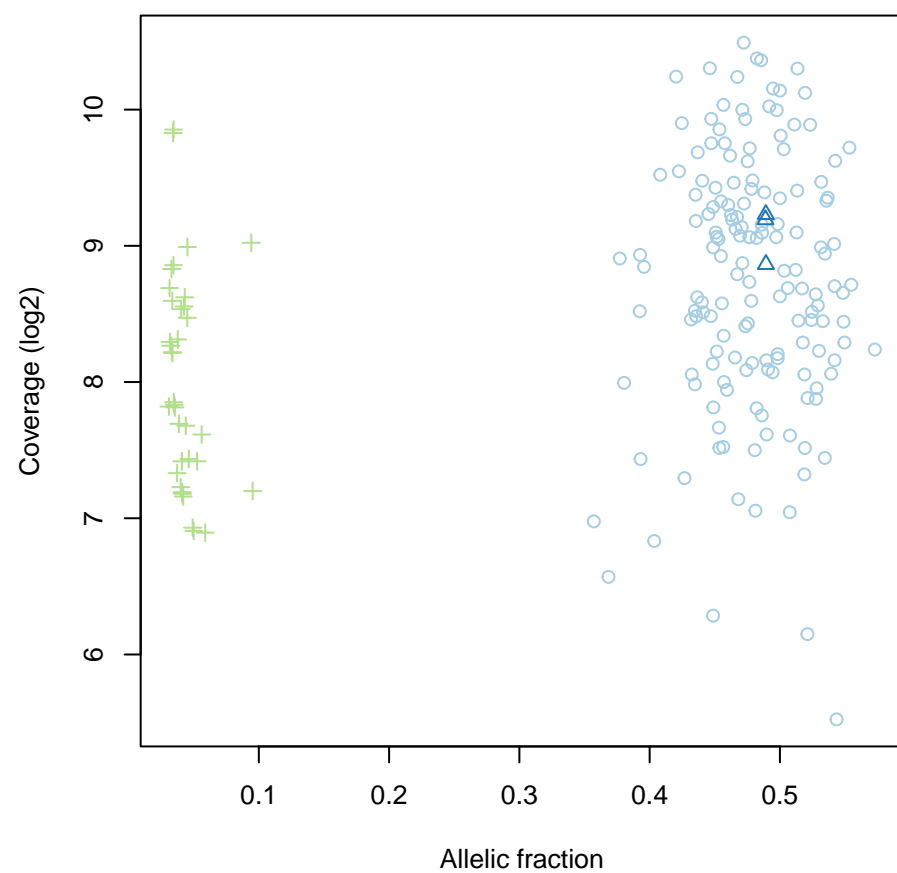
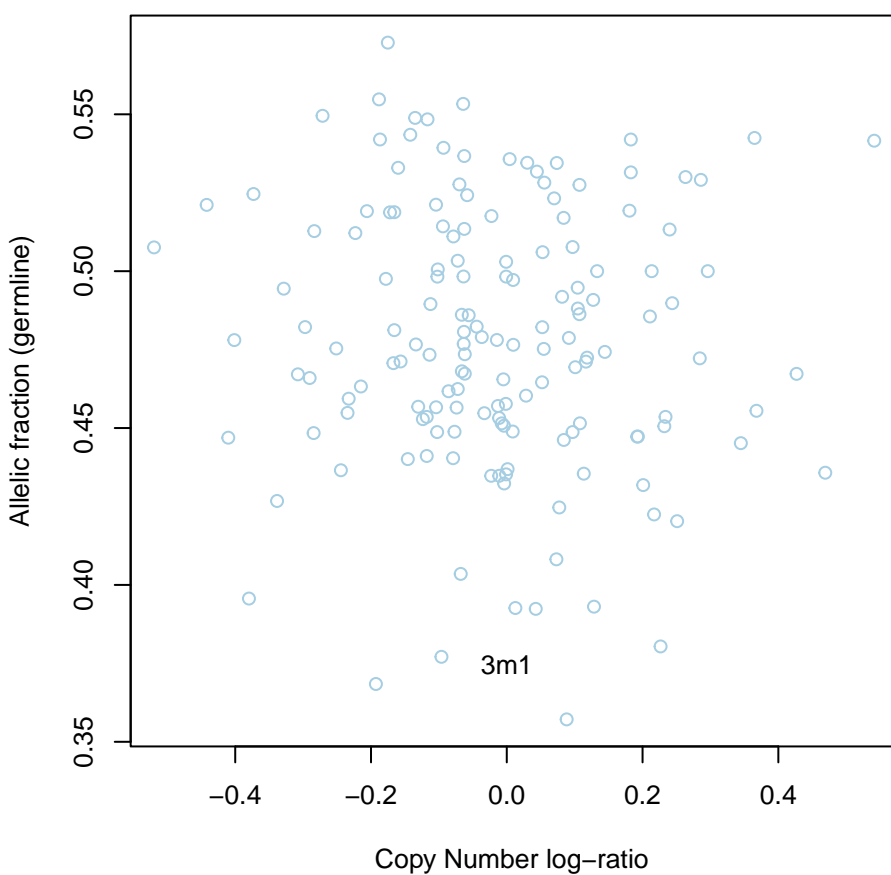


Purity: 0.67 Tumor ploidy: 3 SNV log-likelihood: -800.34 GoF: 62.2% Mean coverage: 410;477

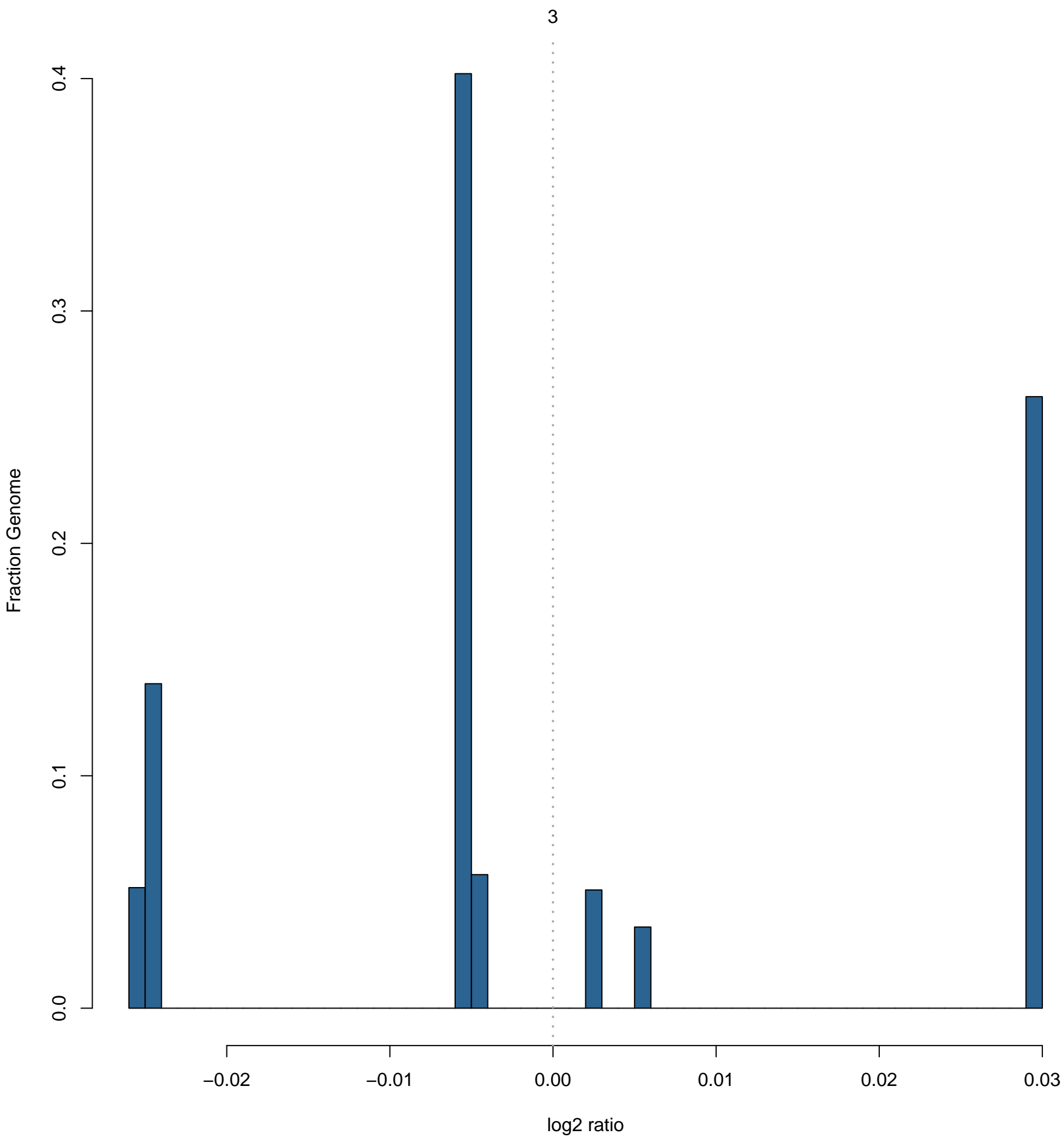


SCNA-fit log-likelihood: -7179.01

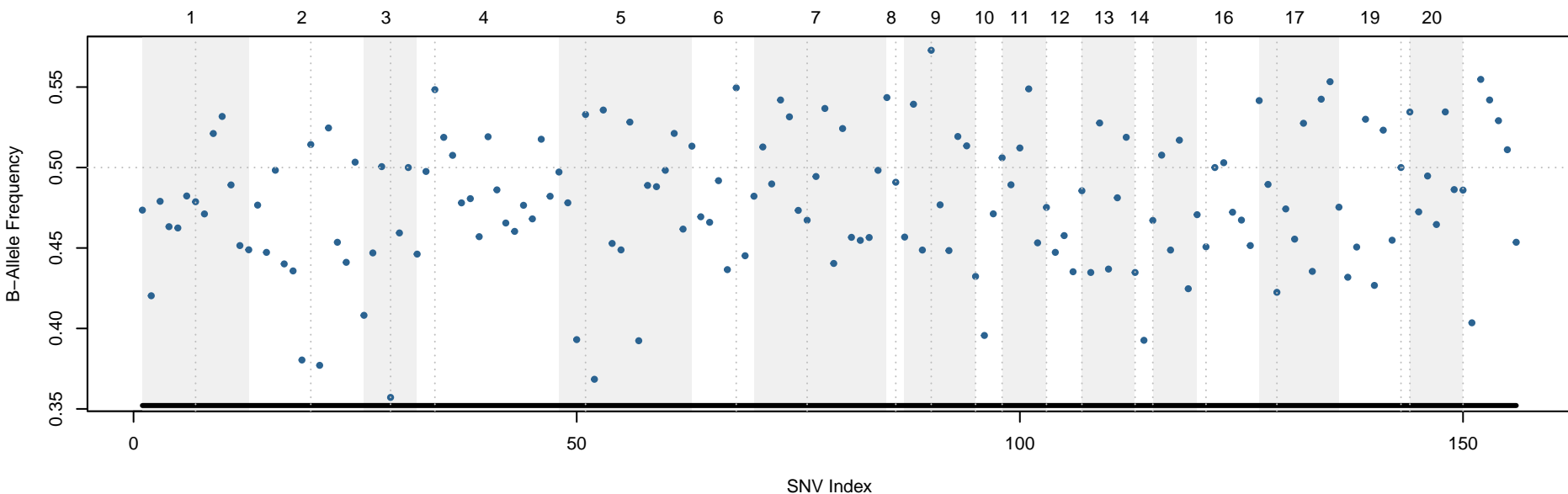




Purity: 0.84 Tumor ploidy: 3



Purity: 0.84 Tumor ploidy: 3 SNV log-likelihood: -1284.25 GoF: 42.6% Mean coverage: 410;477



SCNA-fit log-likelihood: -7179.04

