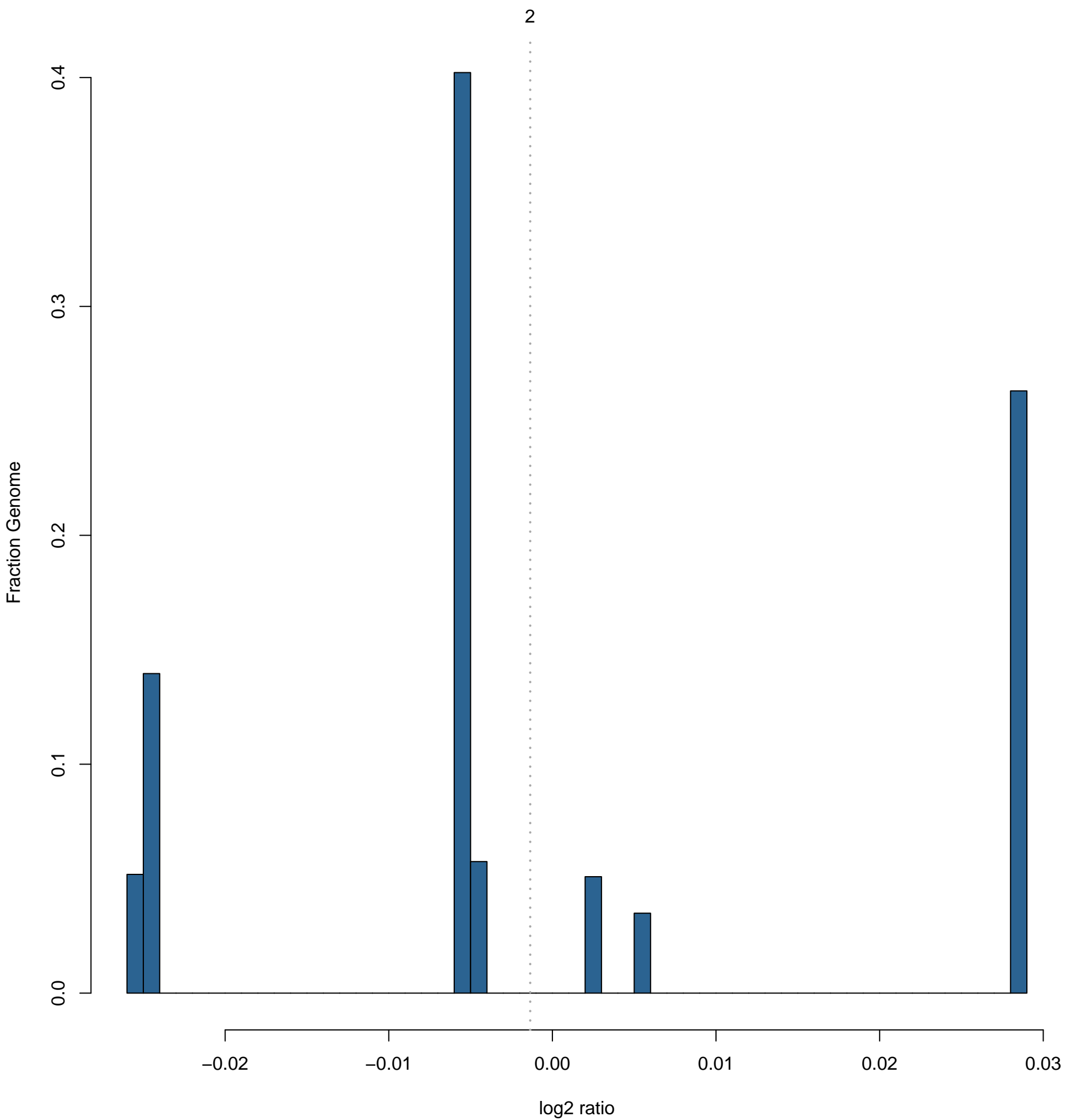
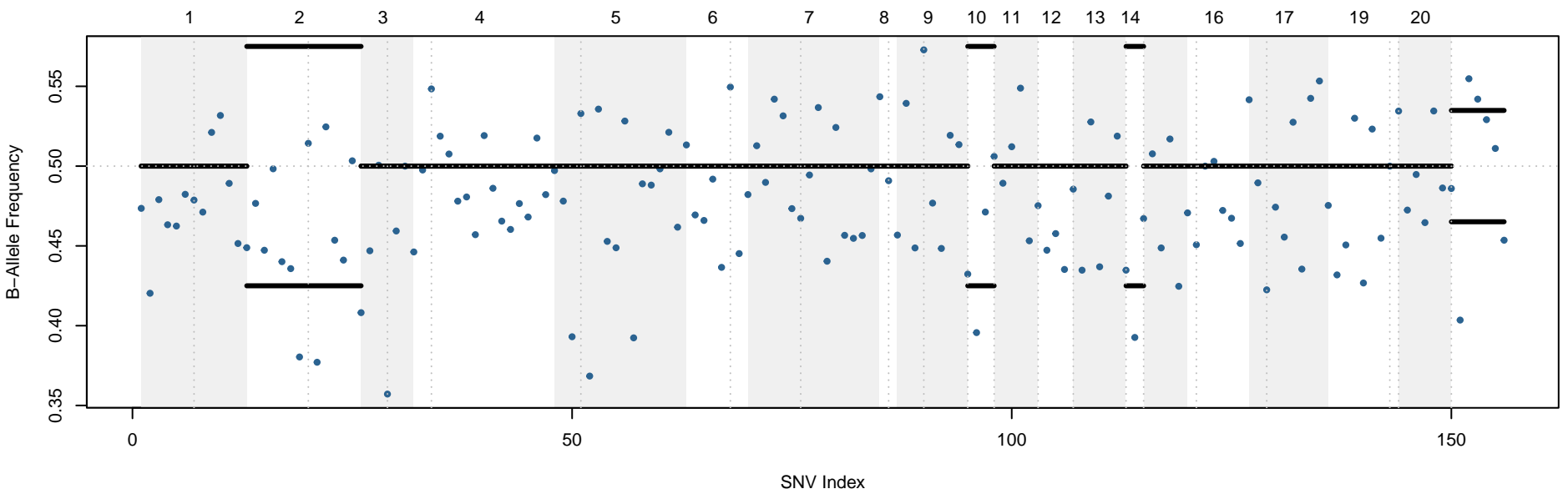


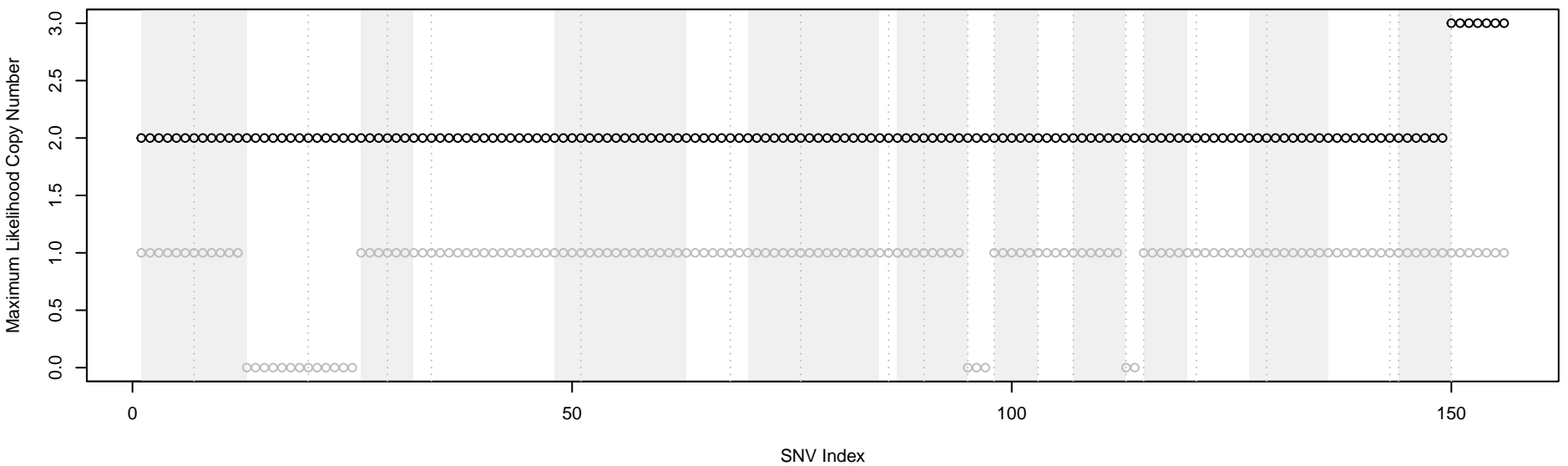
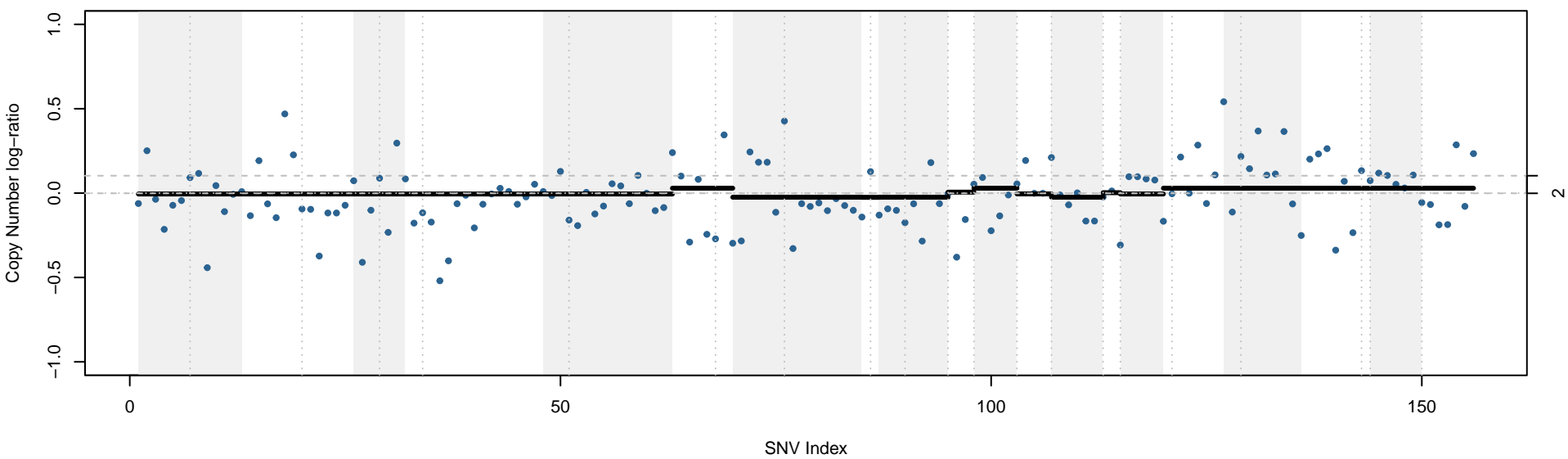
Purity: 0.15 Tumor ploidy: 2.013

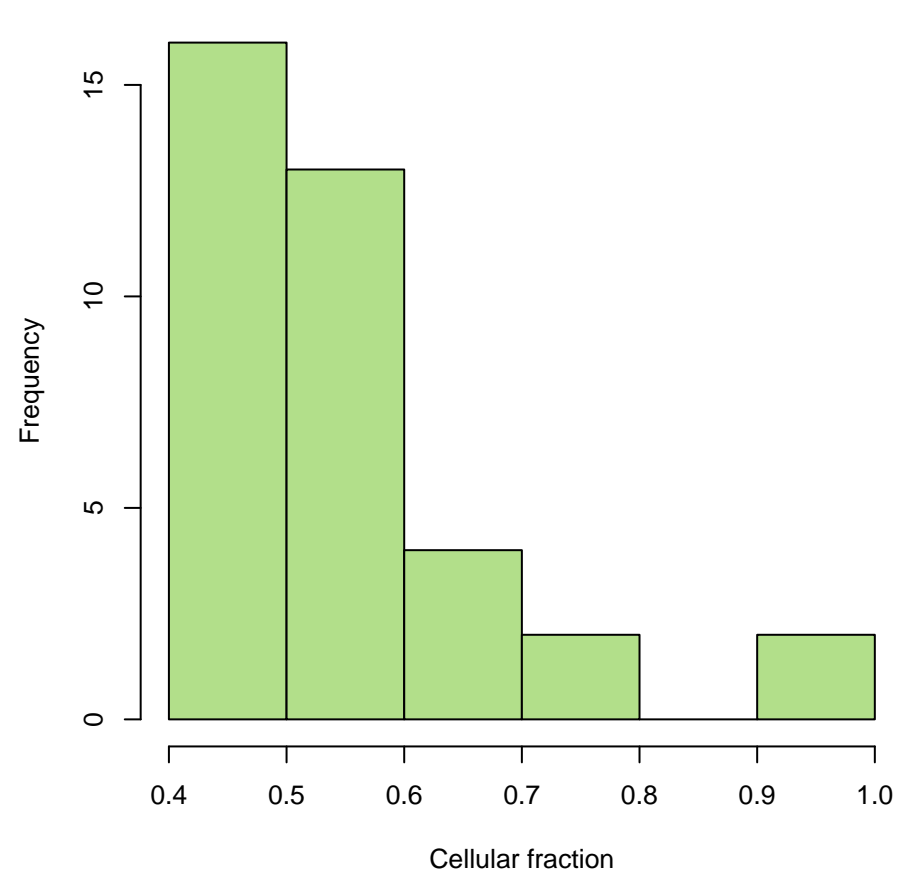
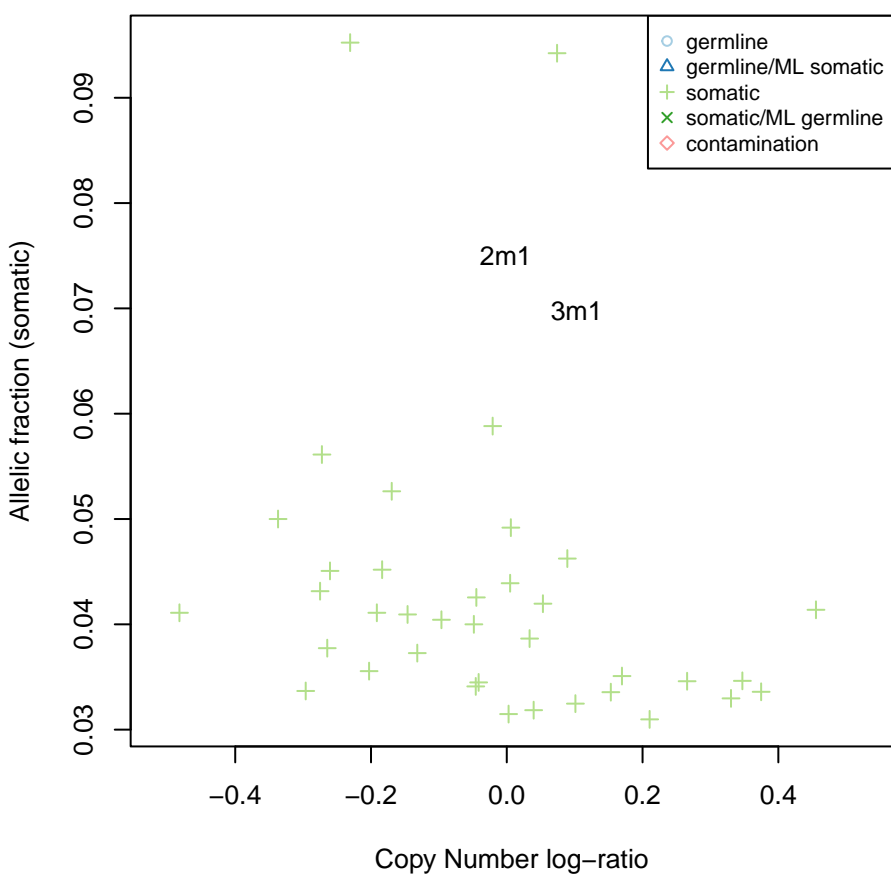
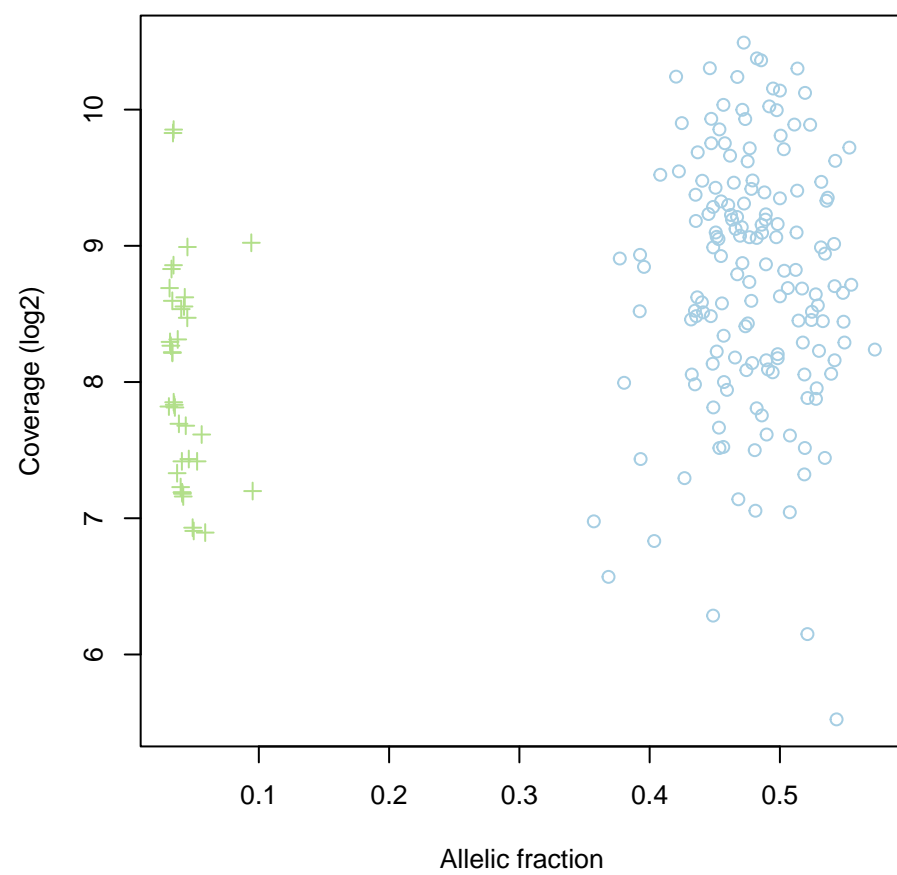
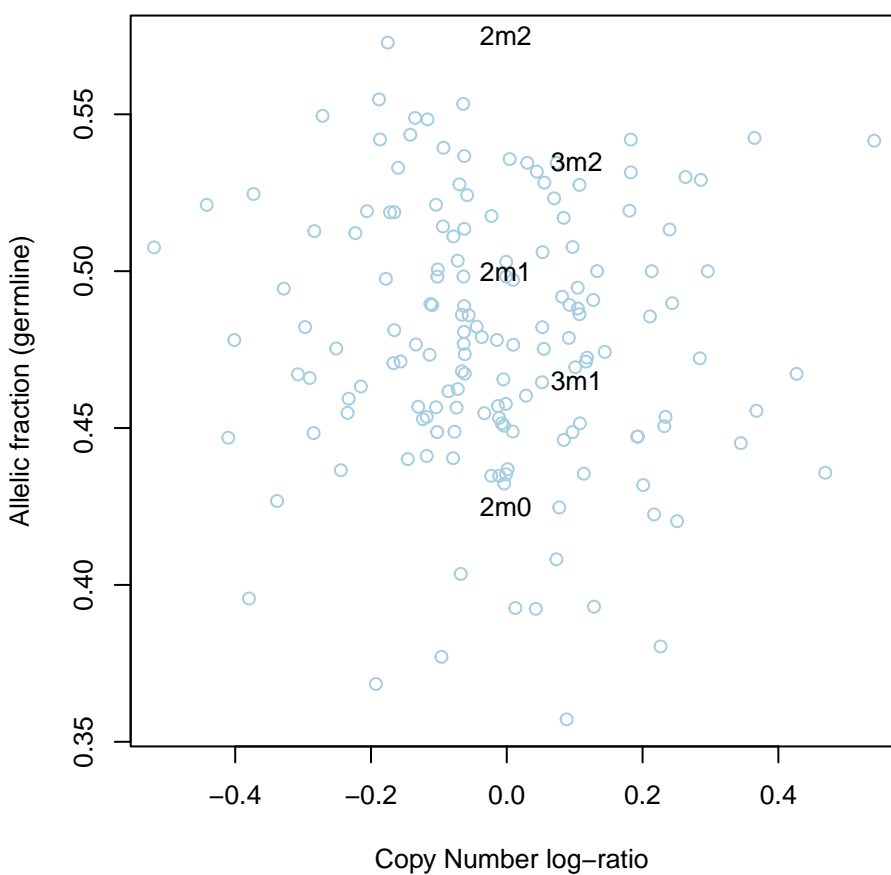


Purity: 0.15 Tumor ploidy: 2.013 SNV log-likelihood: 100.67 GoF: 96.2% Mean coverage: 410;477

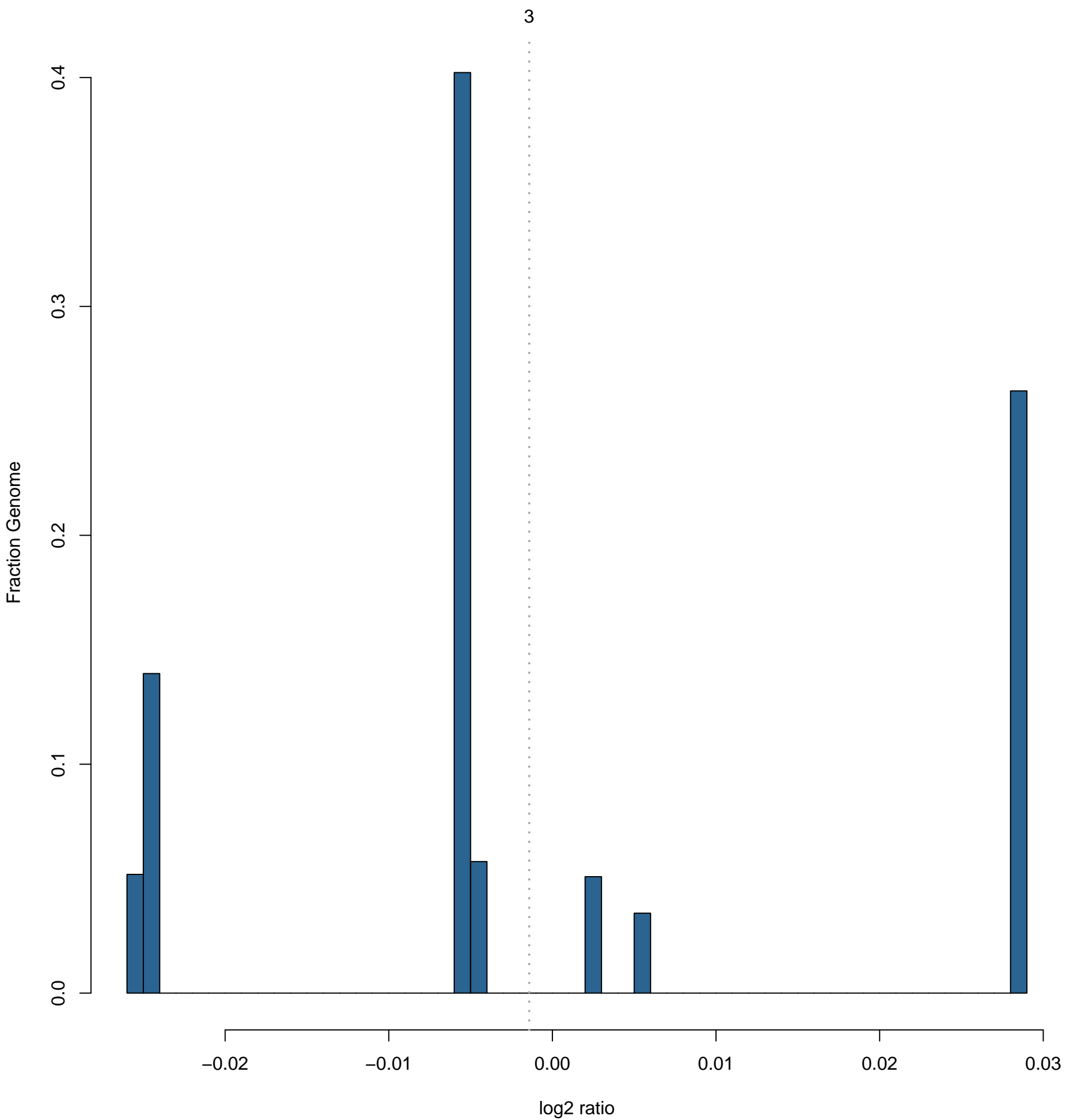


SCNA-fit log-likelihood: -7173.05

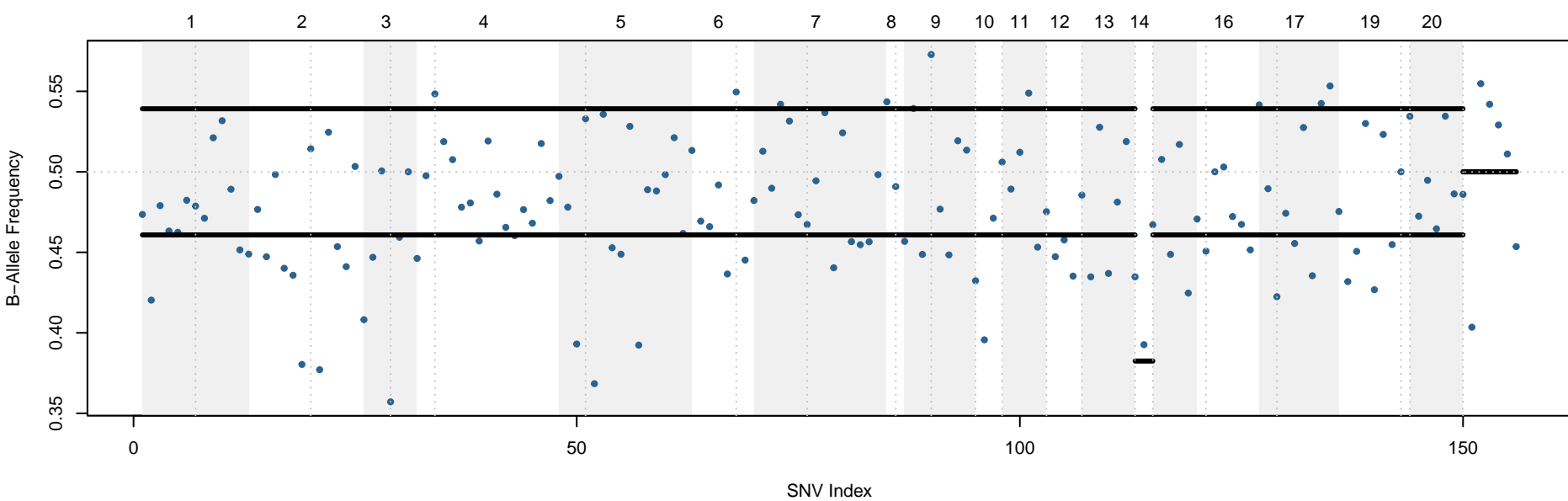




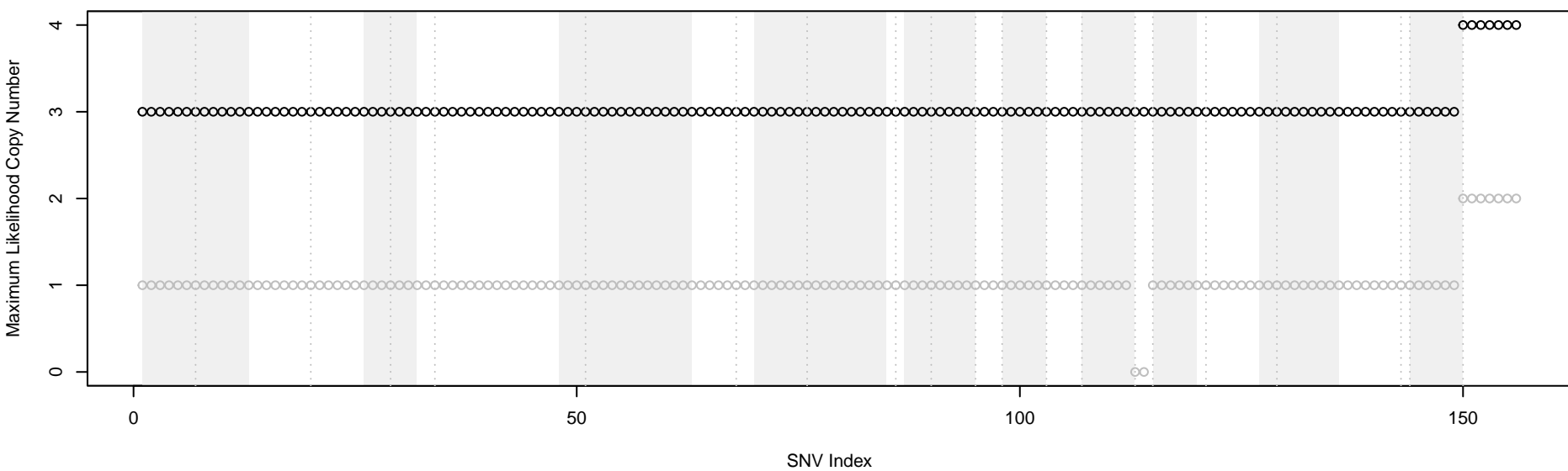
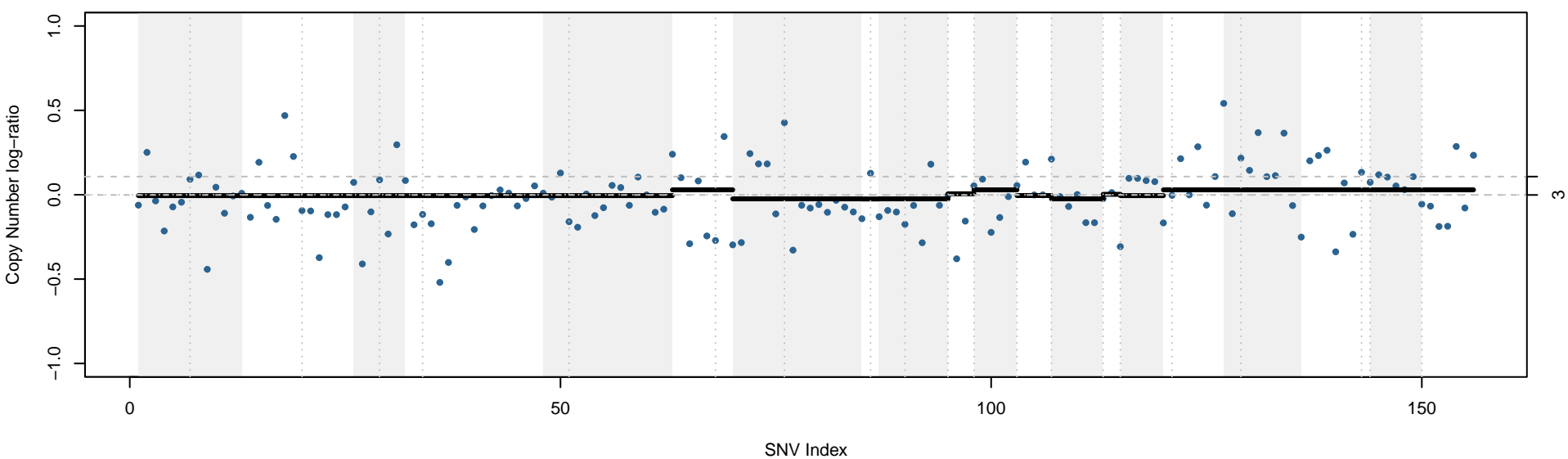
Purity: 0.17 Tumor ploidy: 3.013

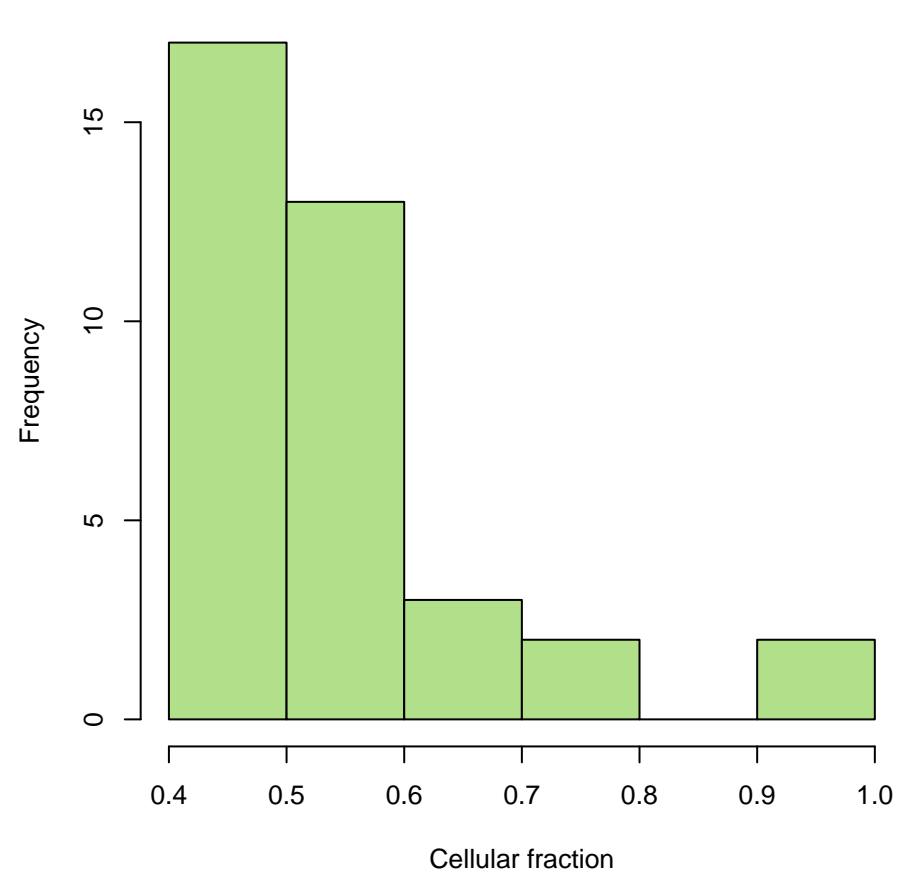
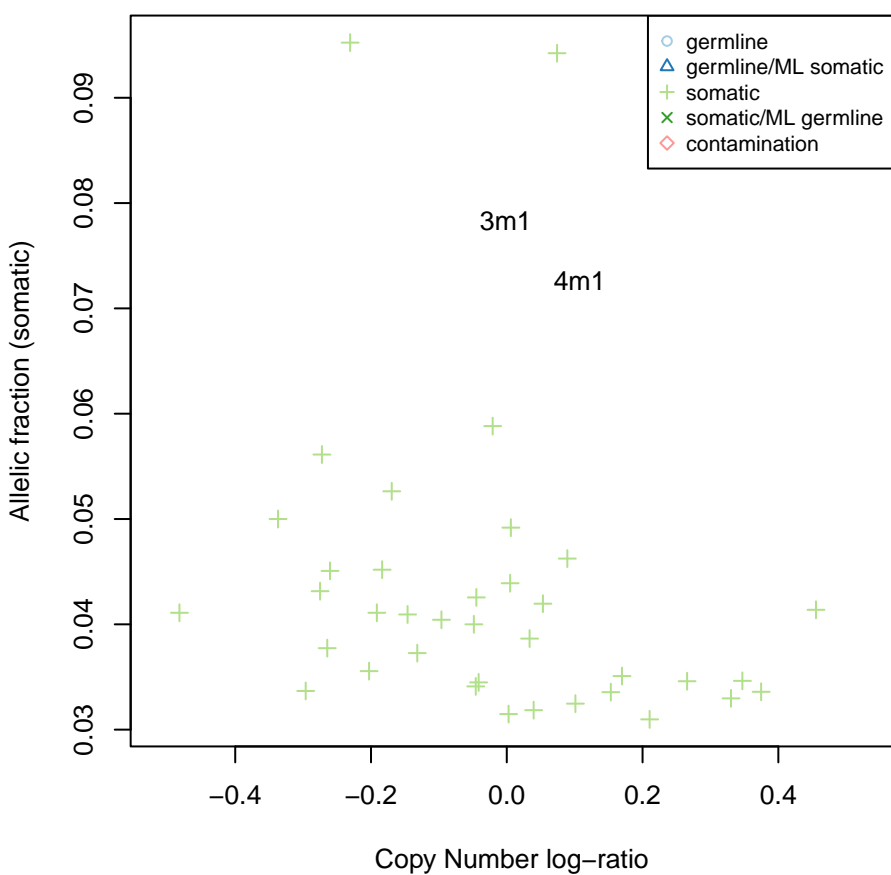
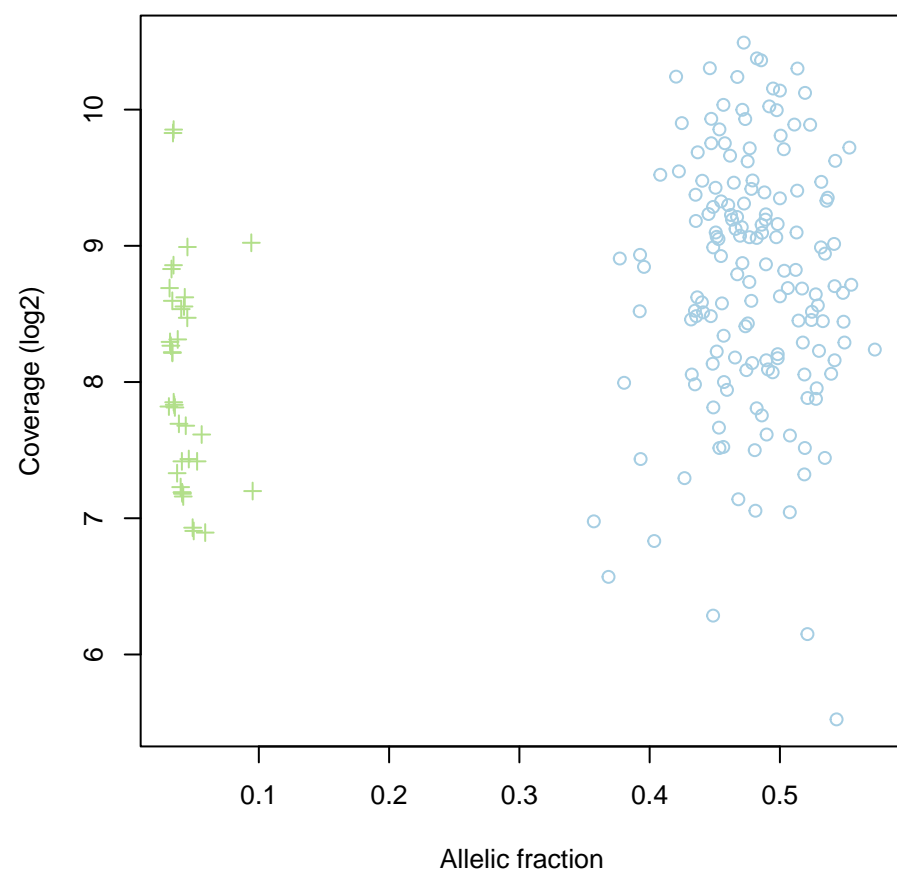
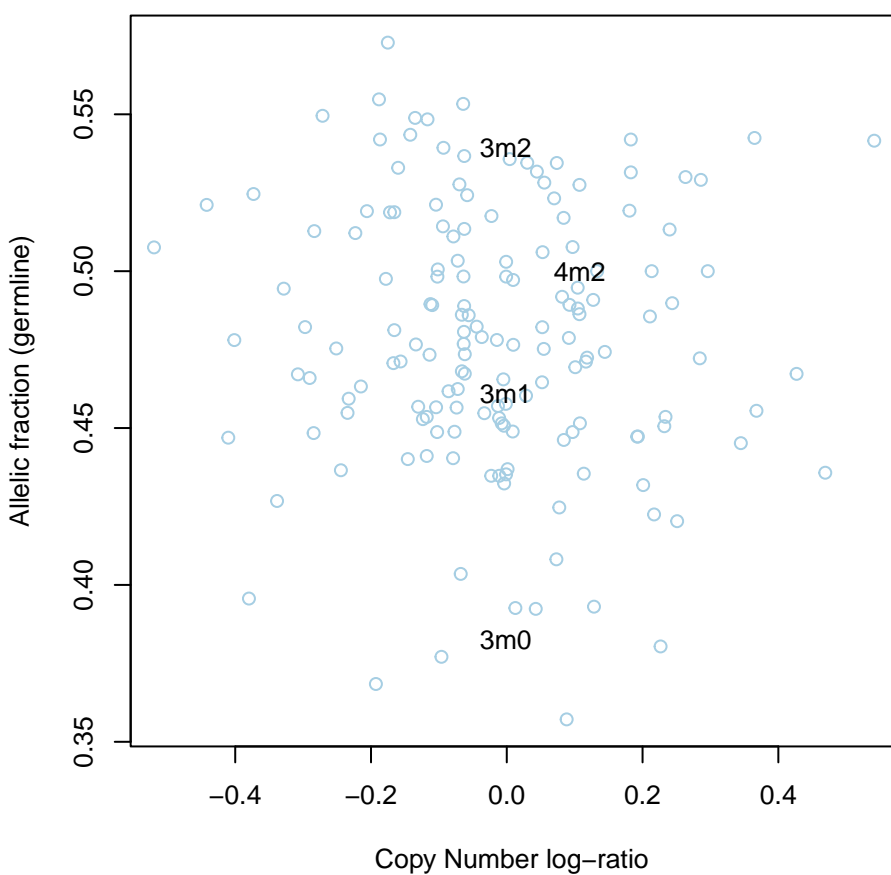


Purity: 0.17 Tumor ploidy: 3.013 SNV log-likelihood: 19.38 GoF: 97.8% Mean coverage: 410;477

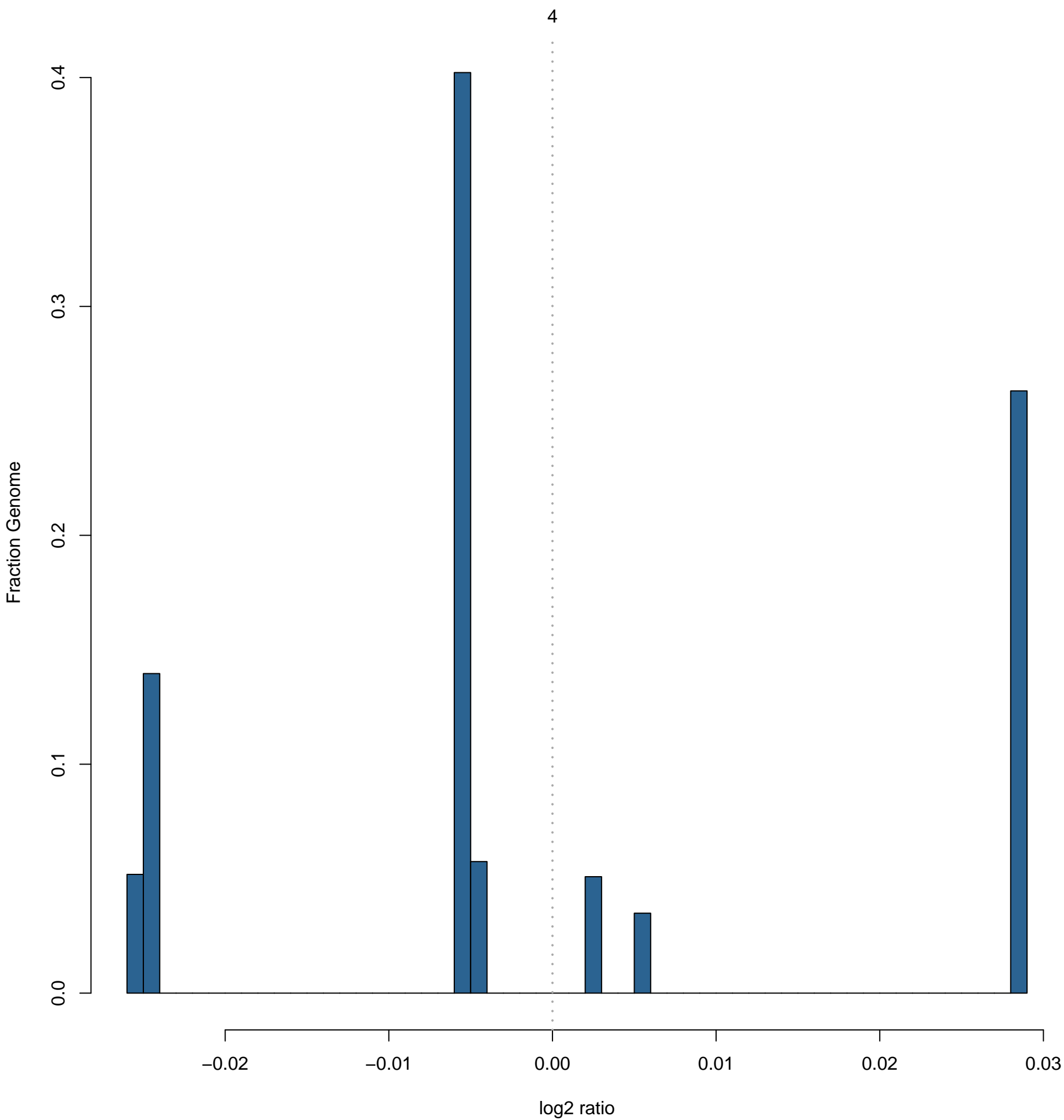


SCNA-fit log-likelihood: -7173.43

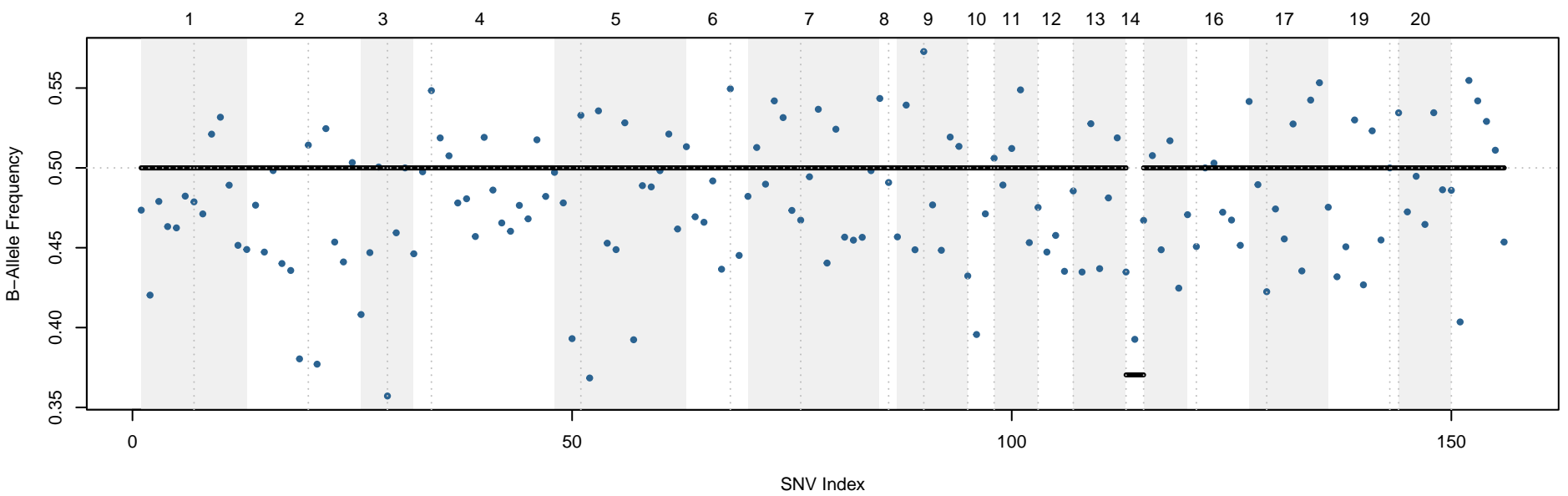




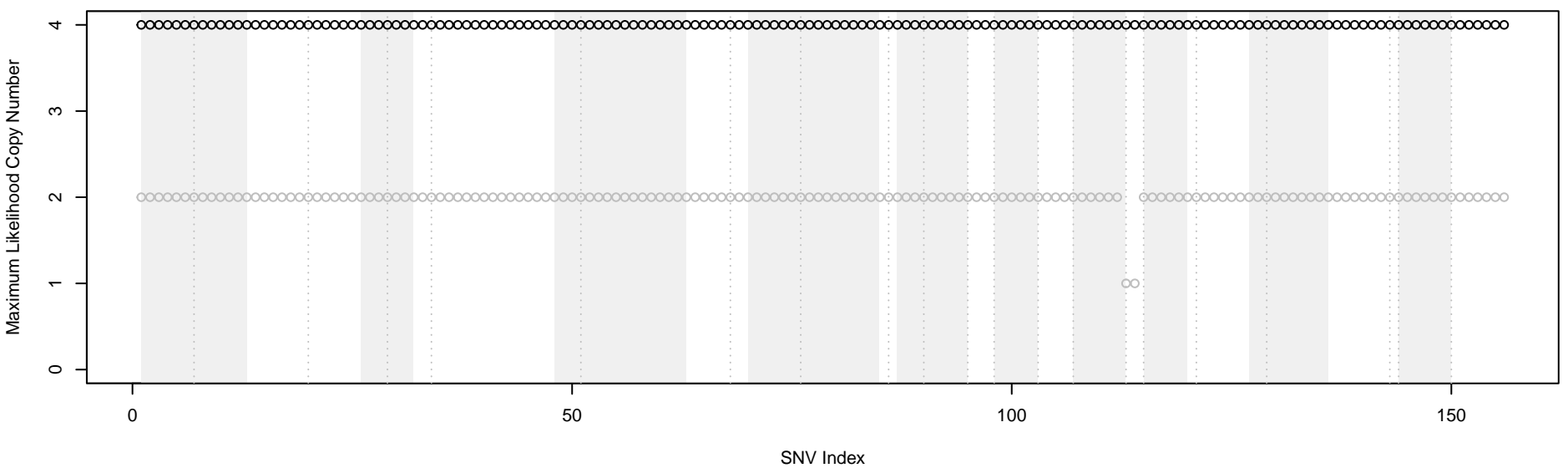
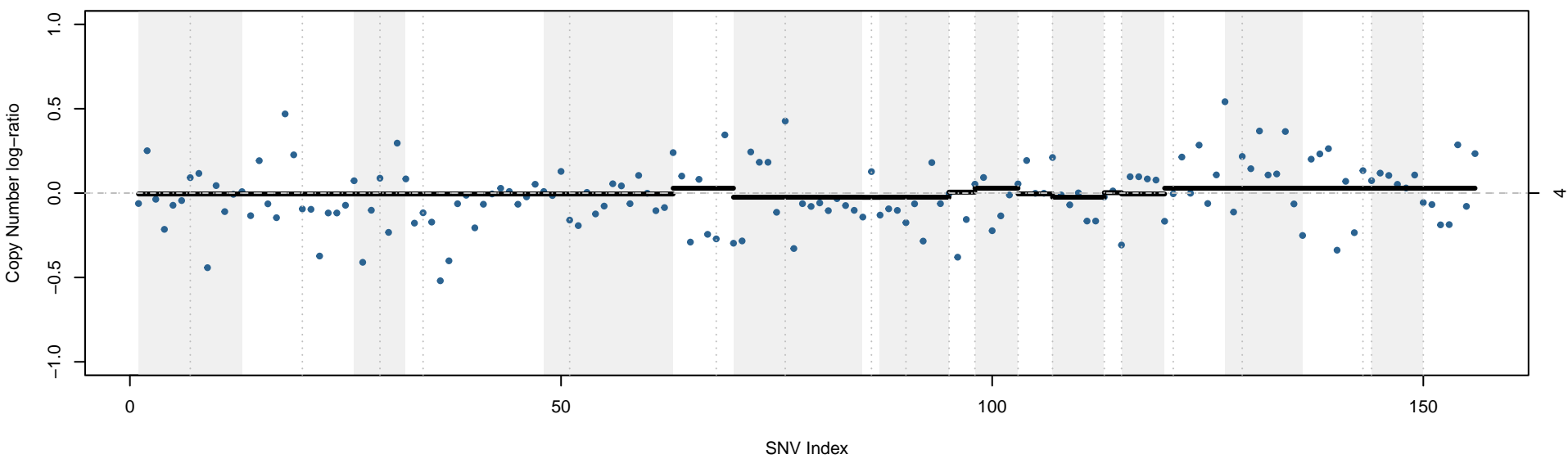
Purity: 0.35 Tumor ploidy: 4



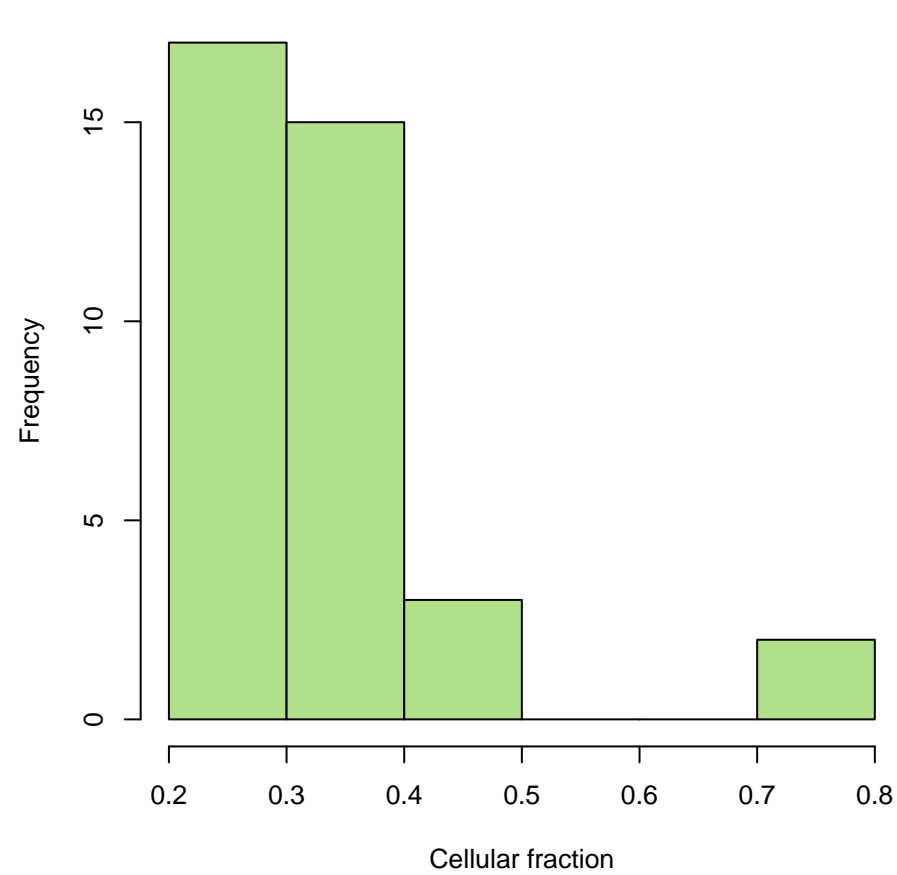
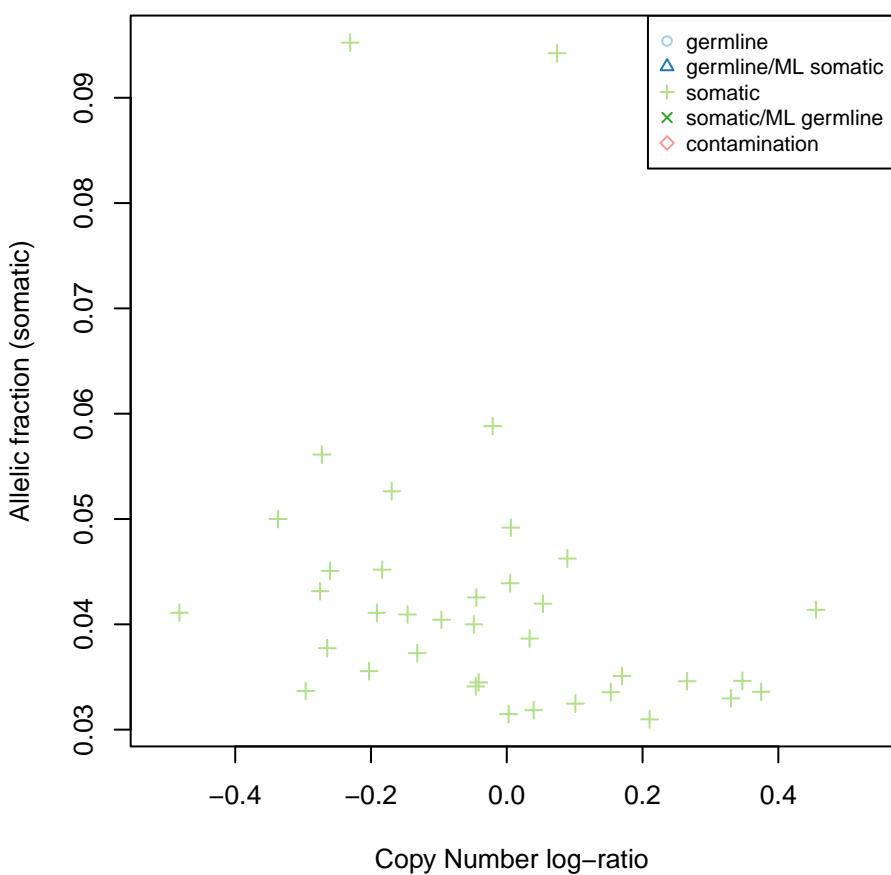
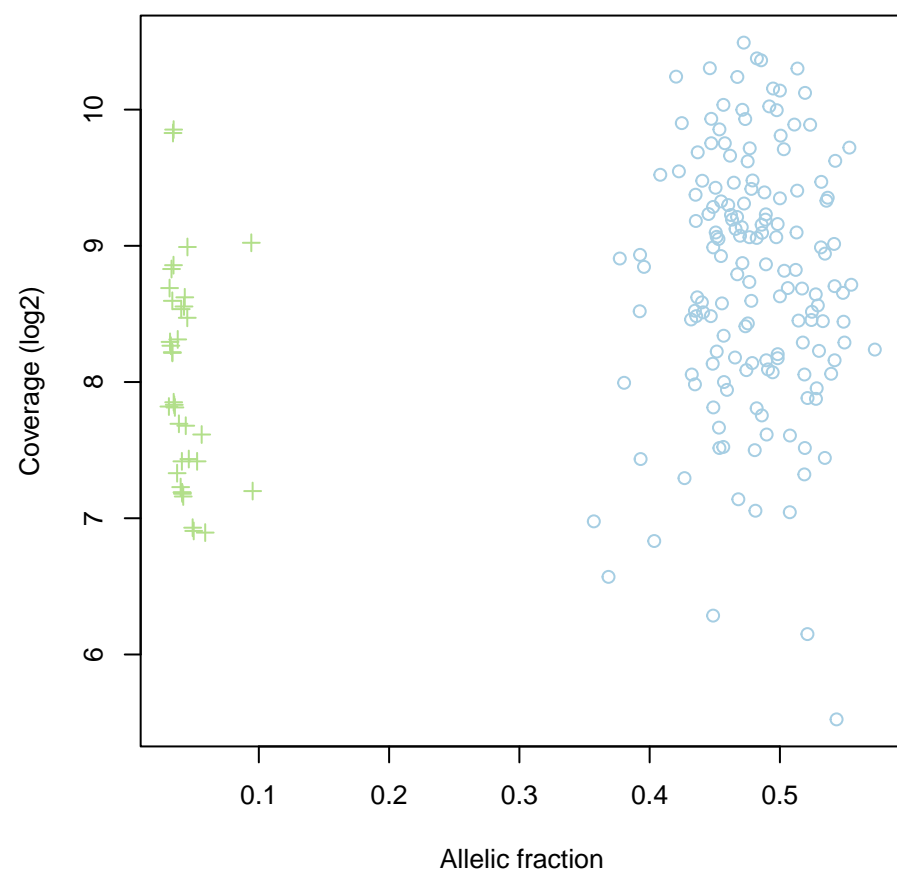
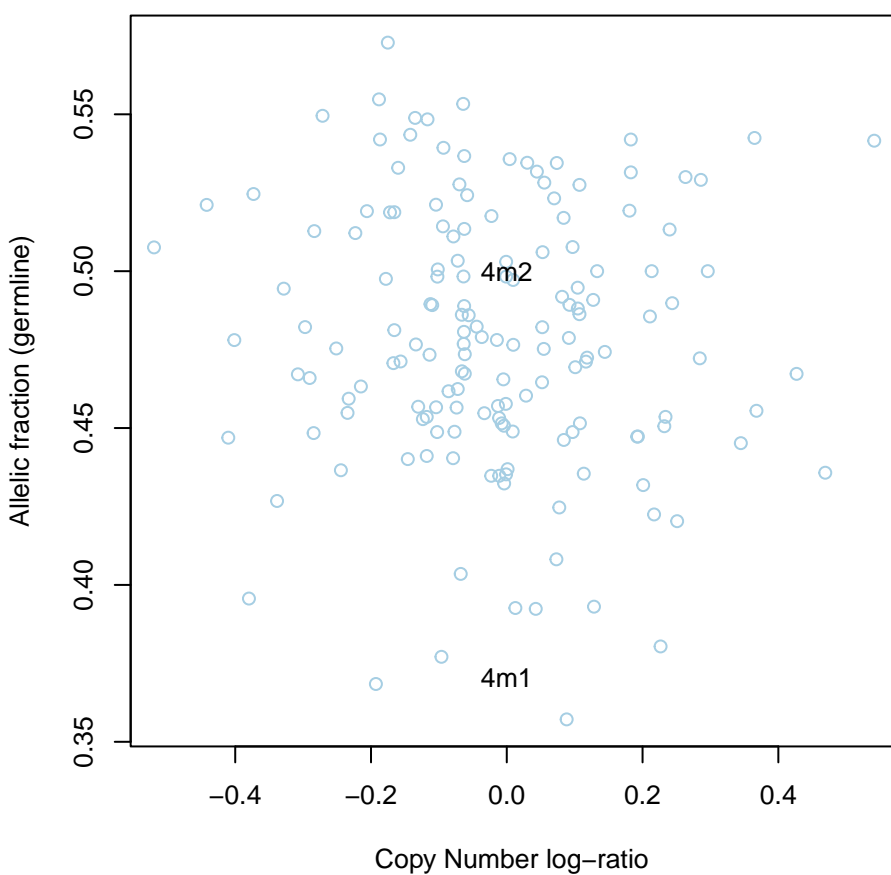
Purity: 0.35 Tumor ploidy: 4 SNV log-likelihood: 4.17 GoF: 92.6% Mean coverage: 410;477



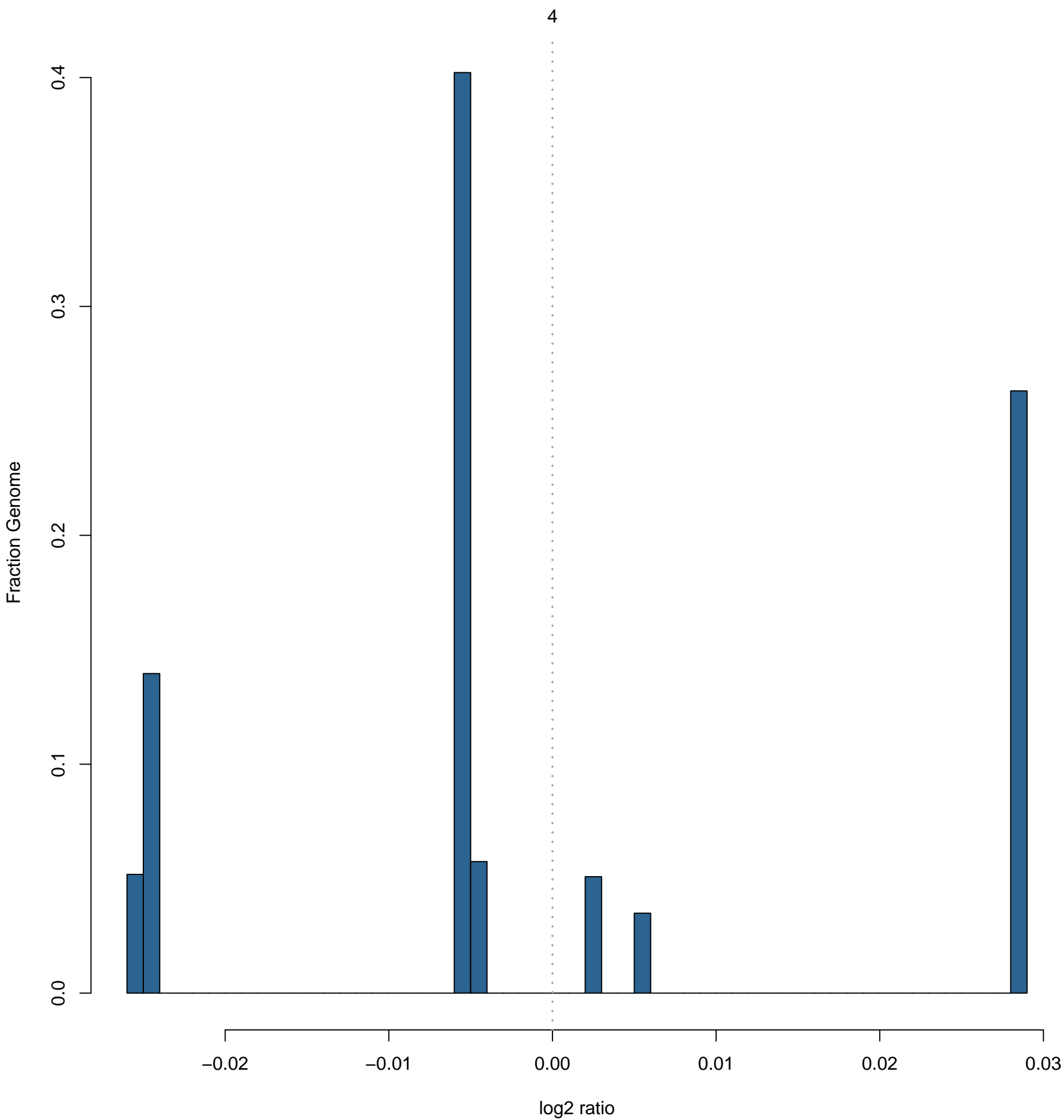
SCNA-fit log-likelihood: -7180.62



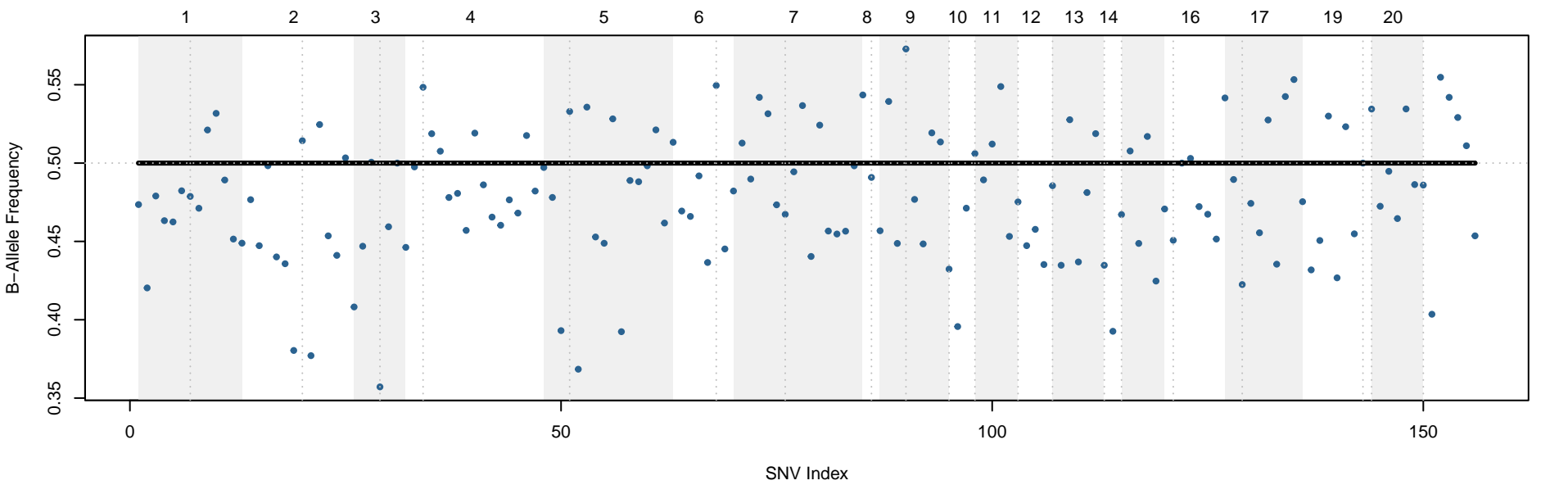




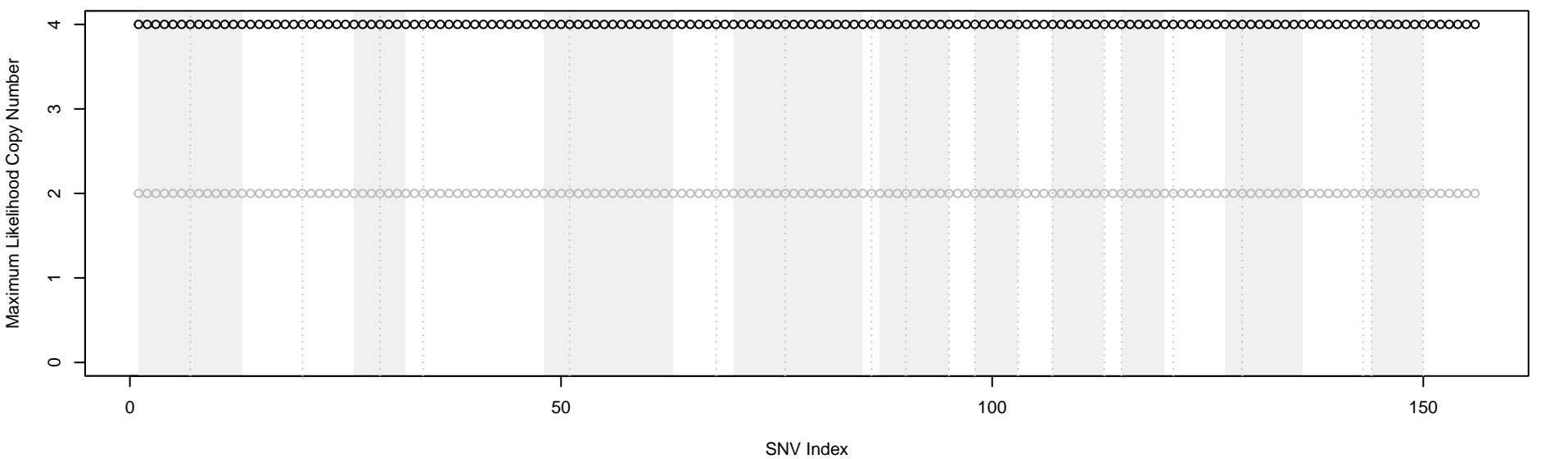
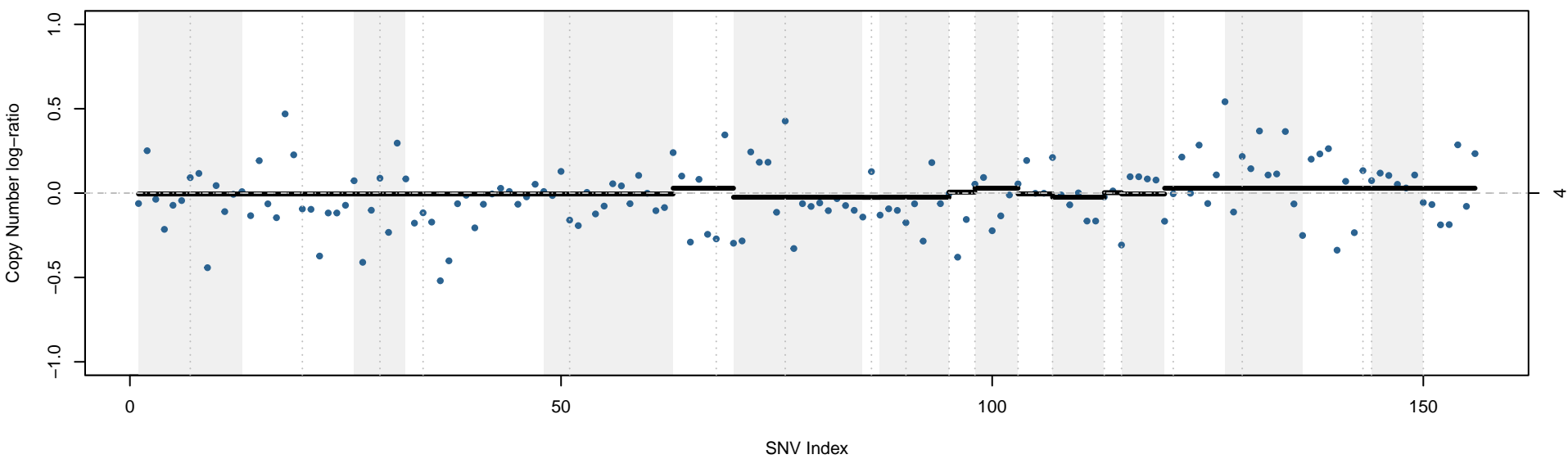
Purity: 0.54 Tumor ploidy: 4

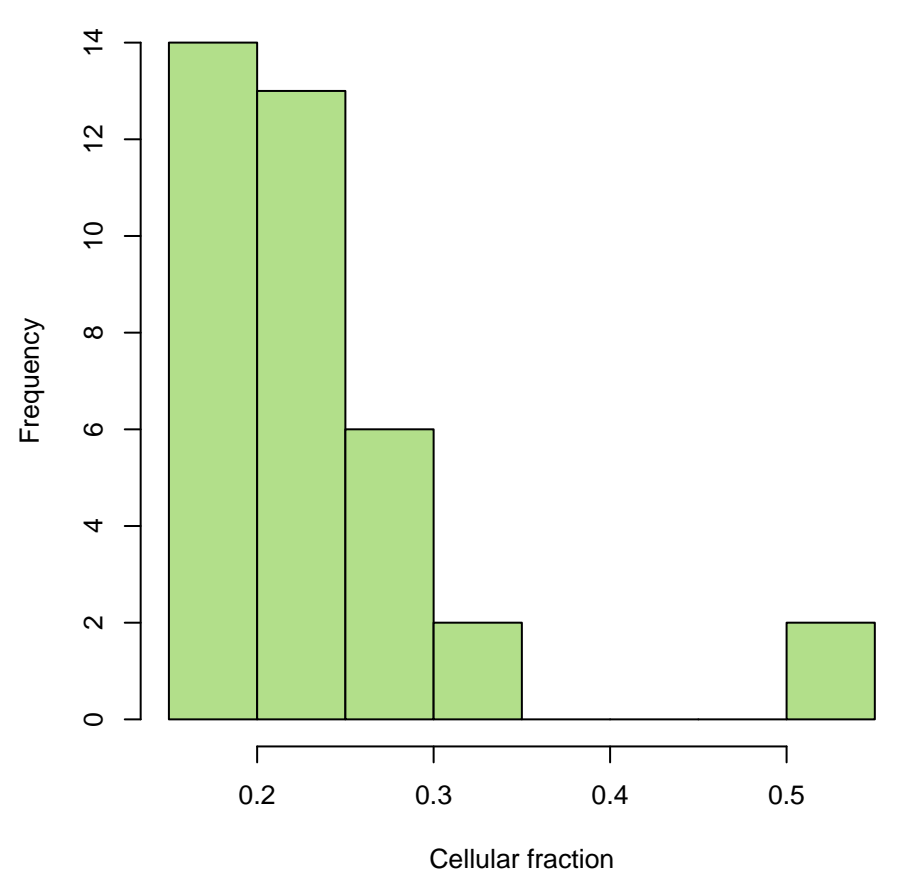
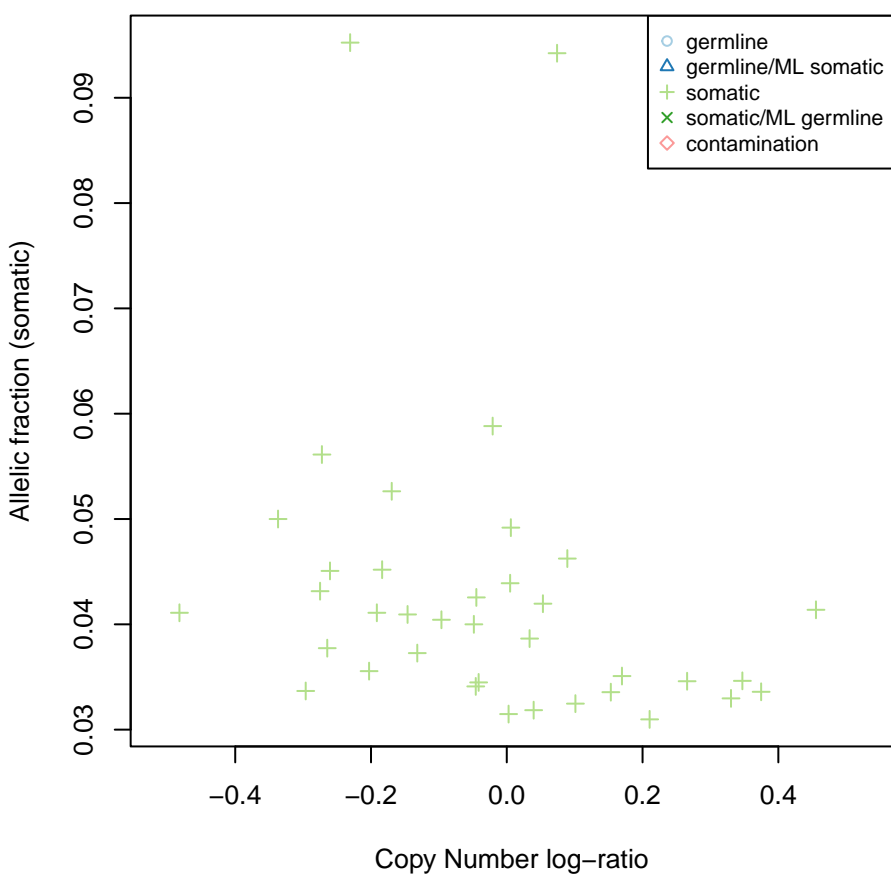
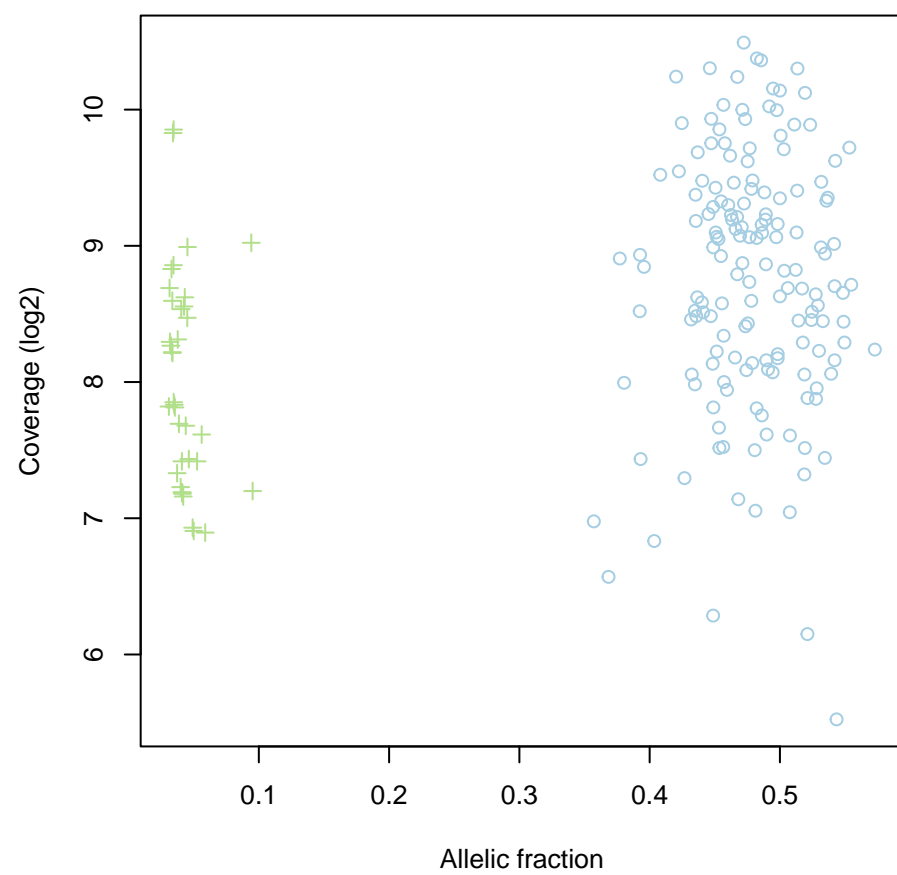
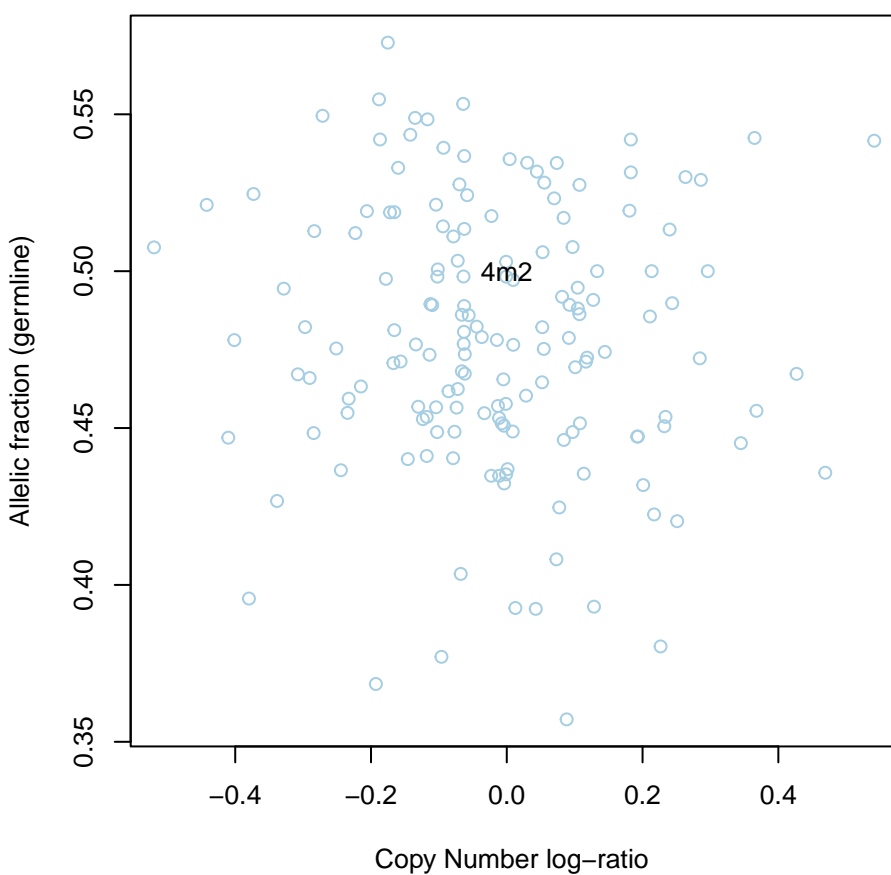


Purity: 0.54 Tumor ploidy: 4 SNV log-likelihood: -22.46 GoF: 87.7% Mean coverage: 410;477

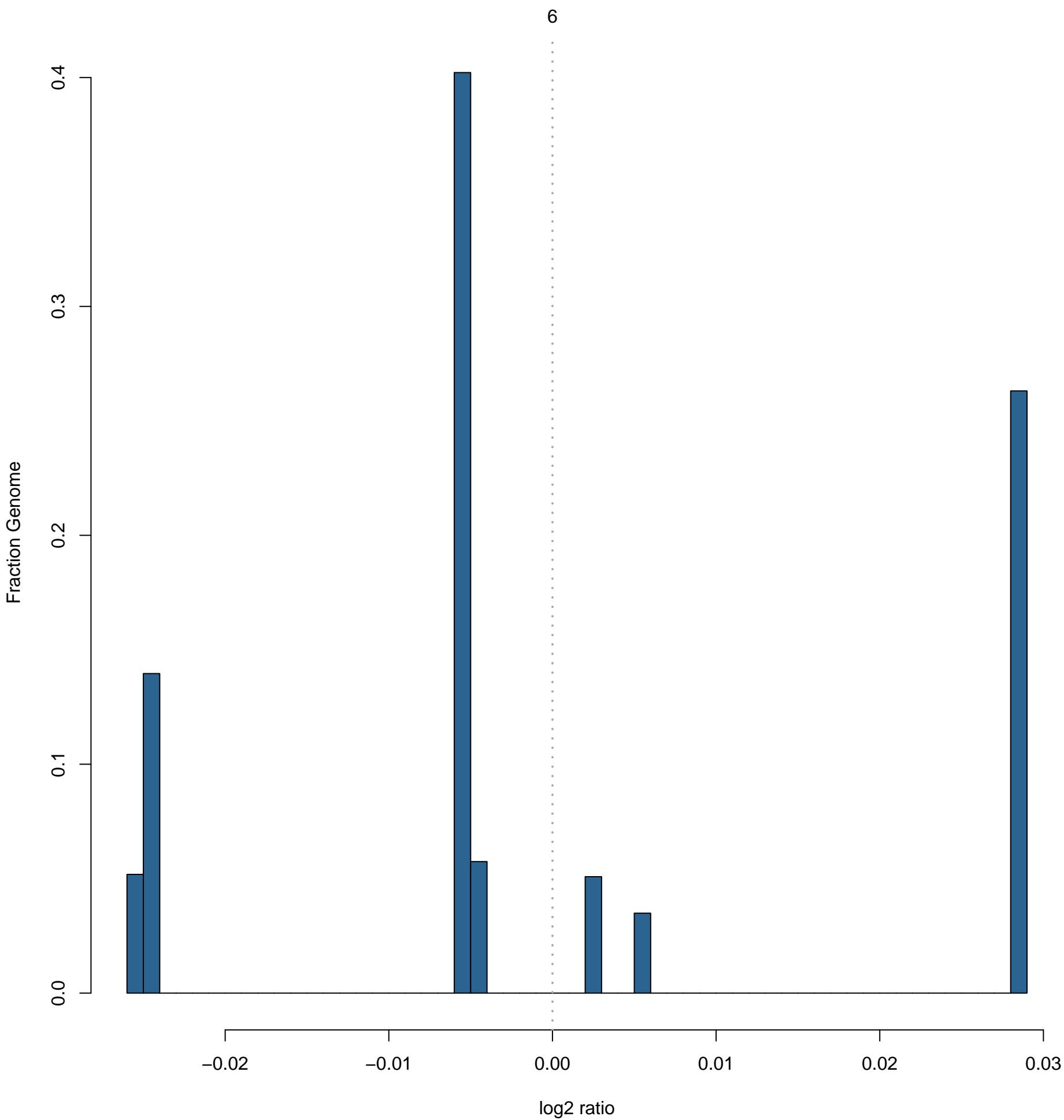


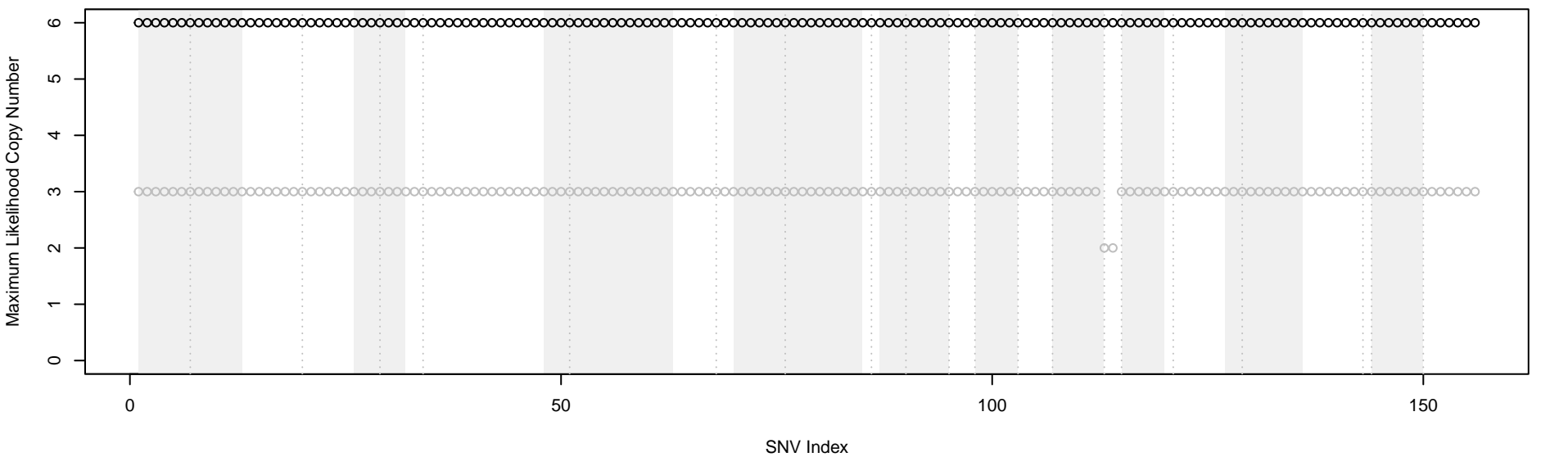
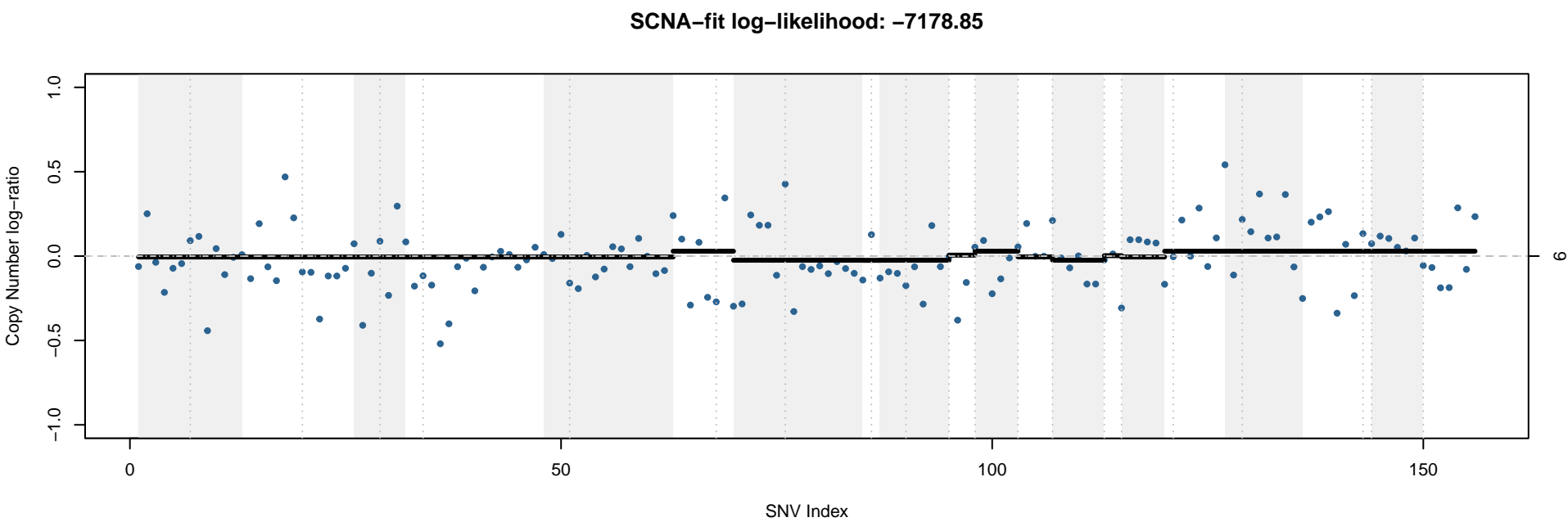
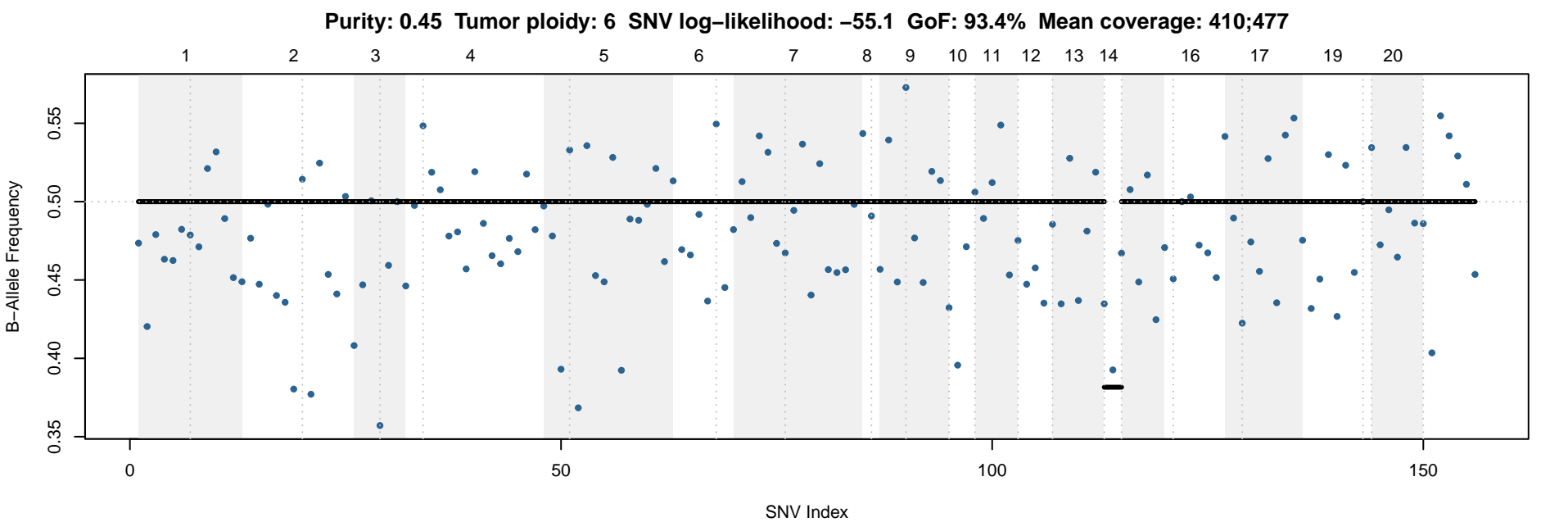
SCNA-fit log-likelihood: -7178.86

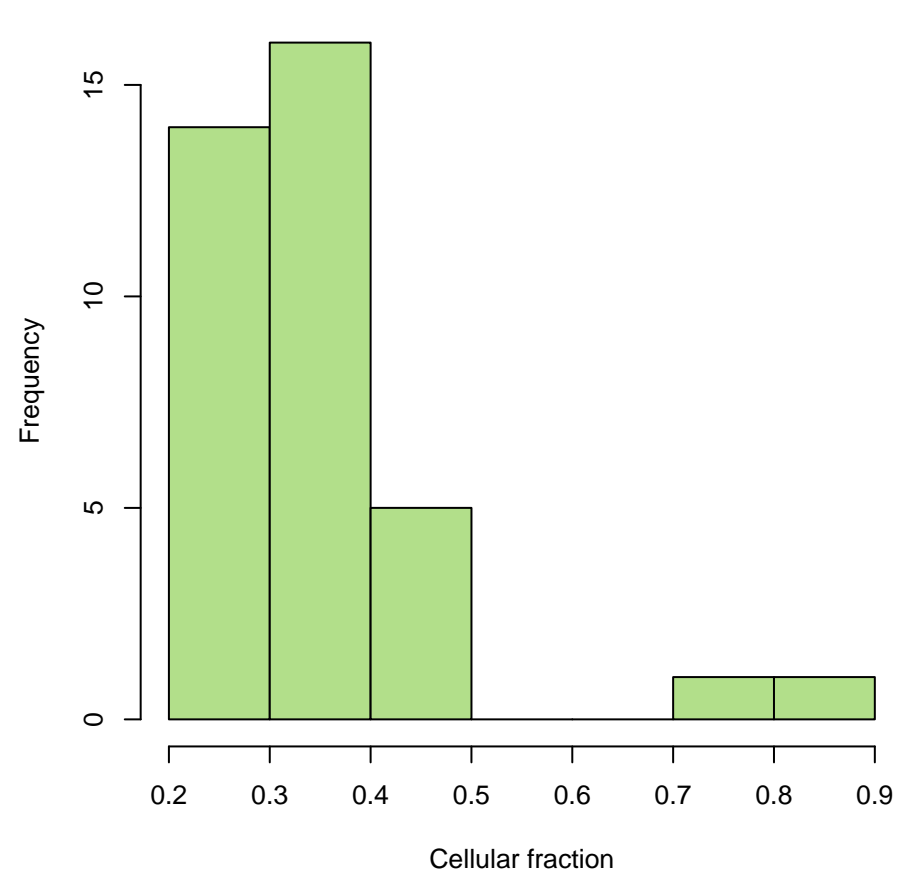
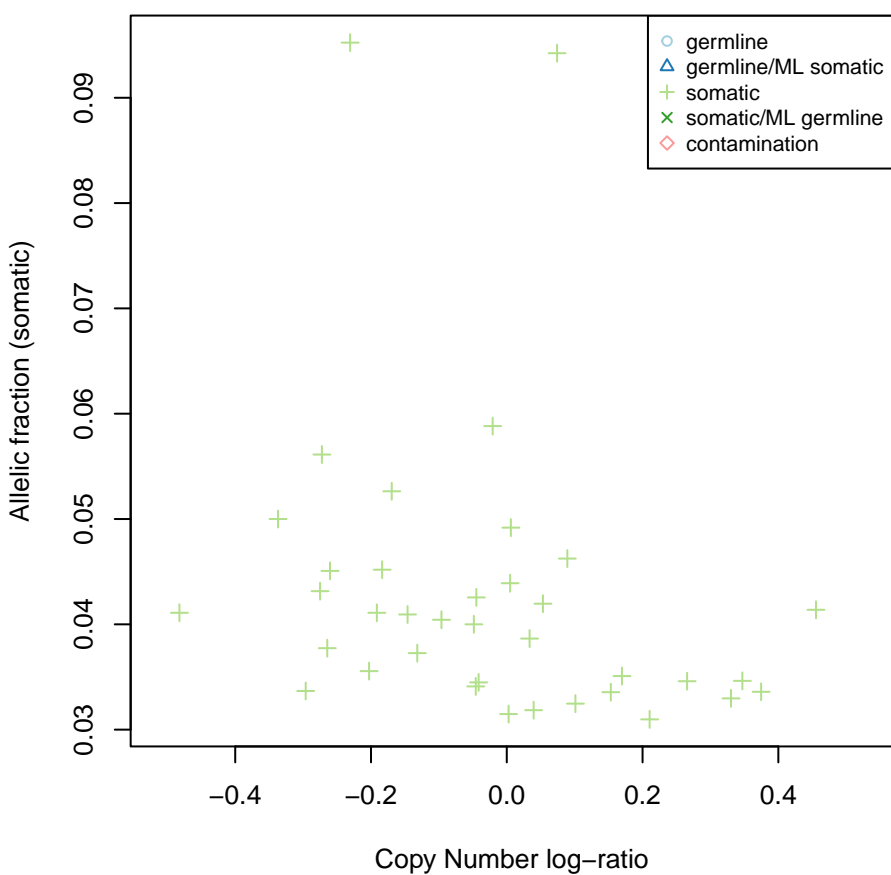
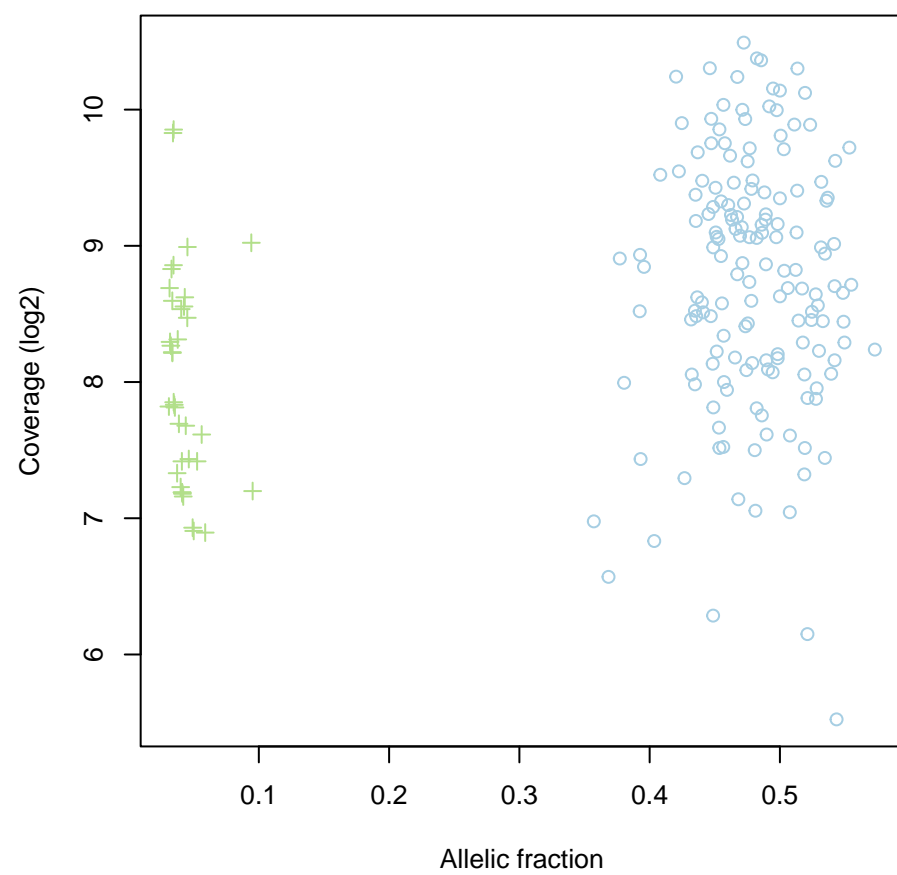
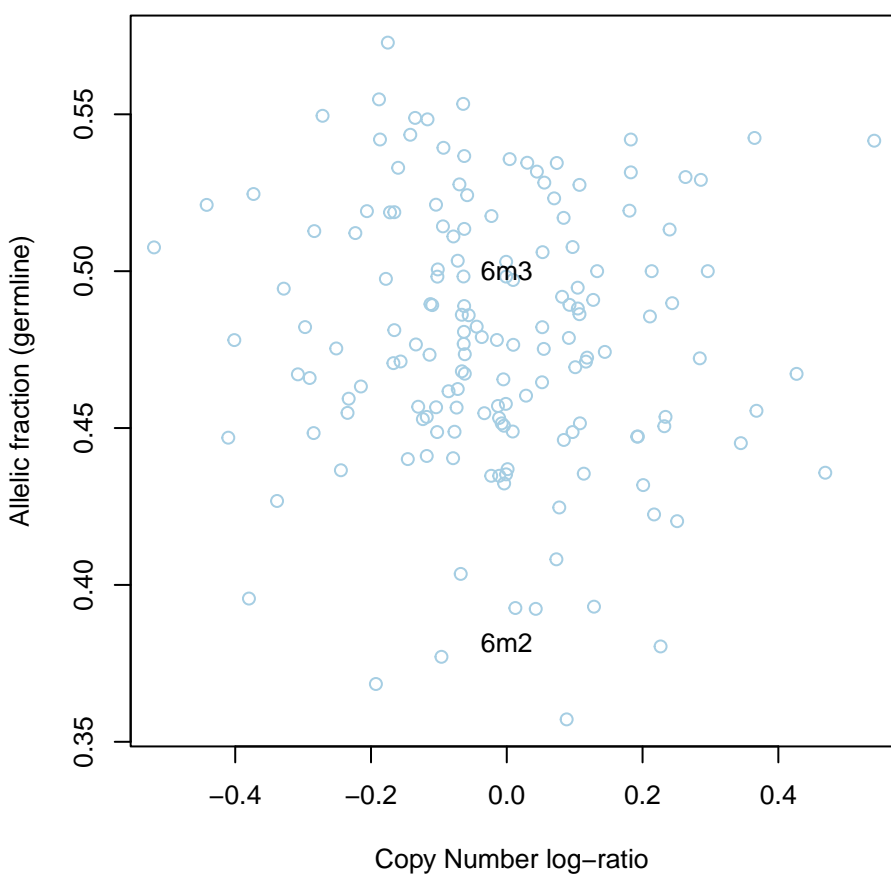




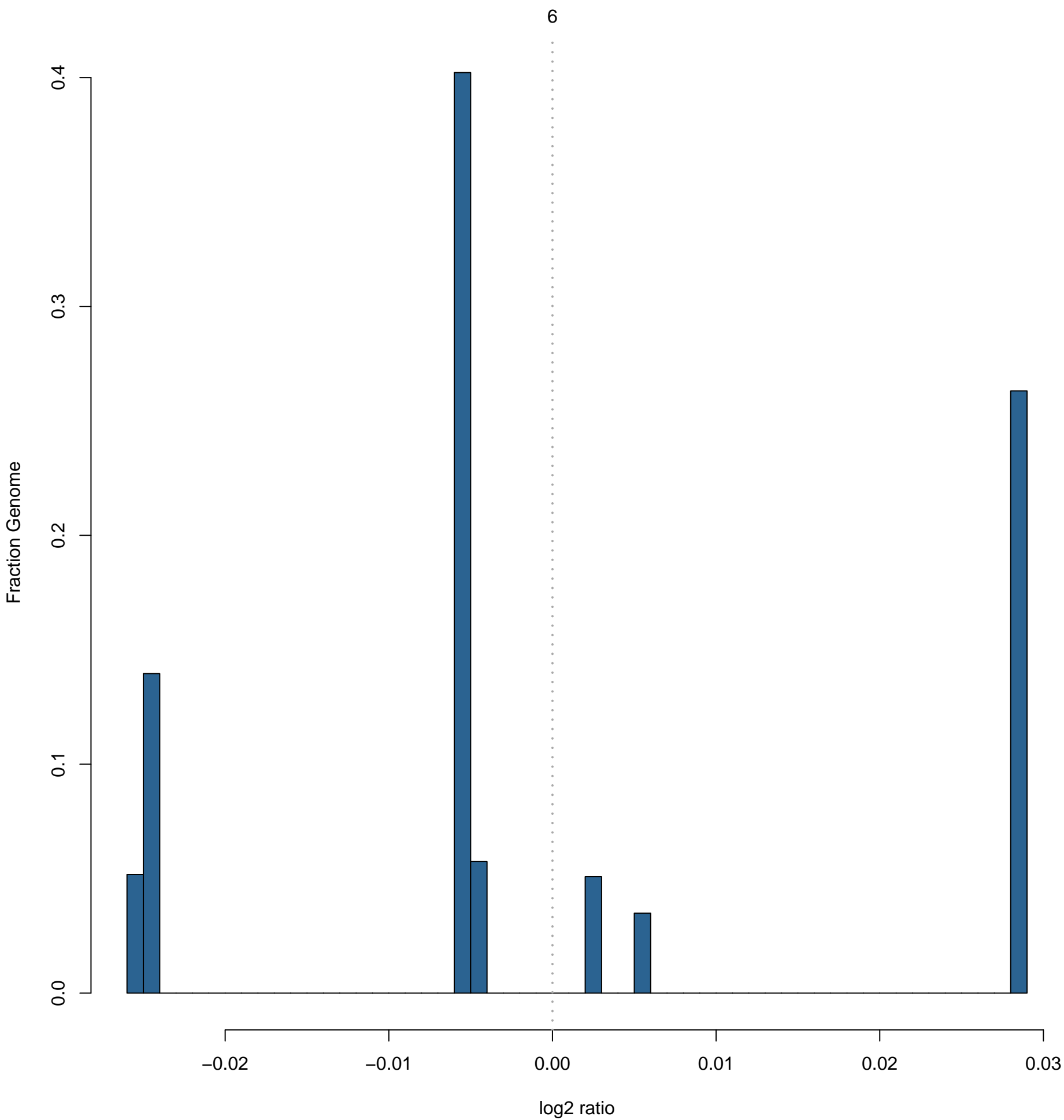
Purity: 0.45 Tumor ploidy: 6





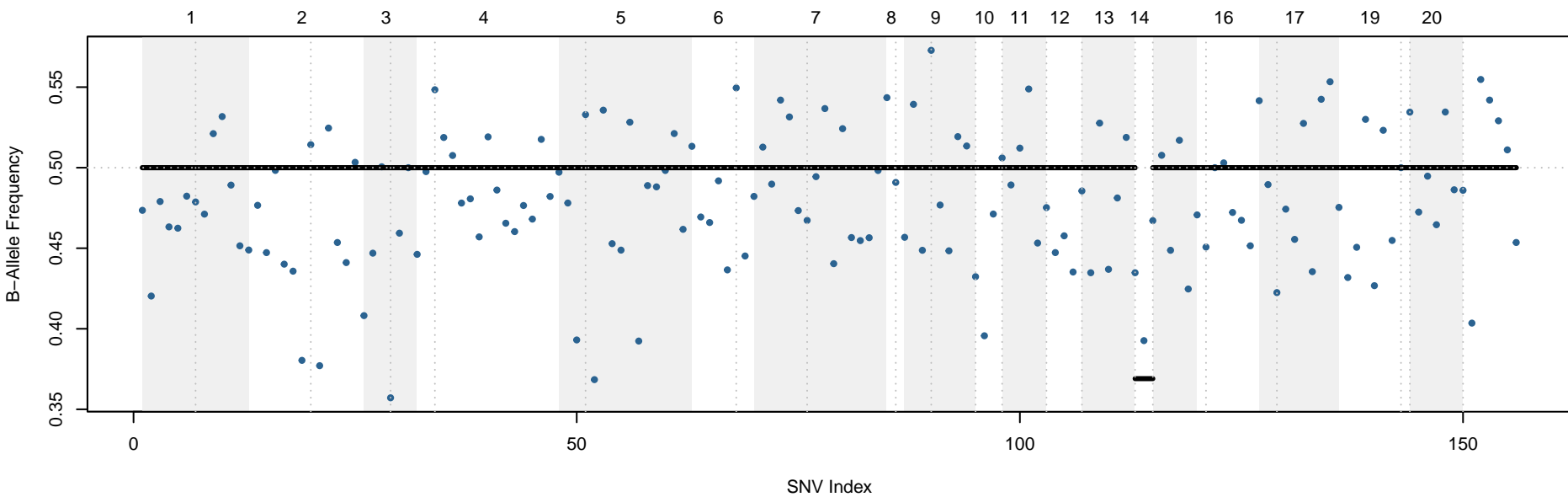


Purity: 0.55 Tumor ploidy: 6

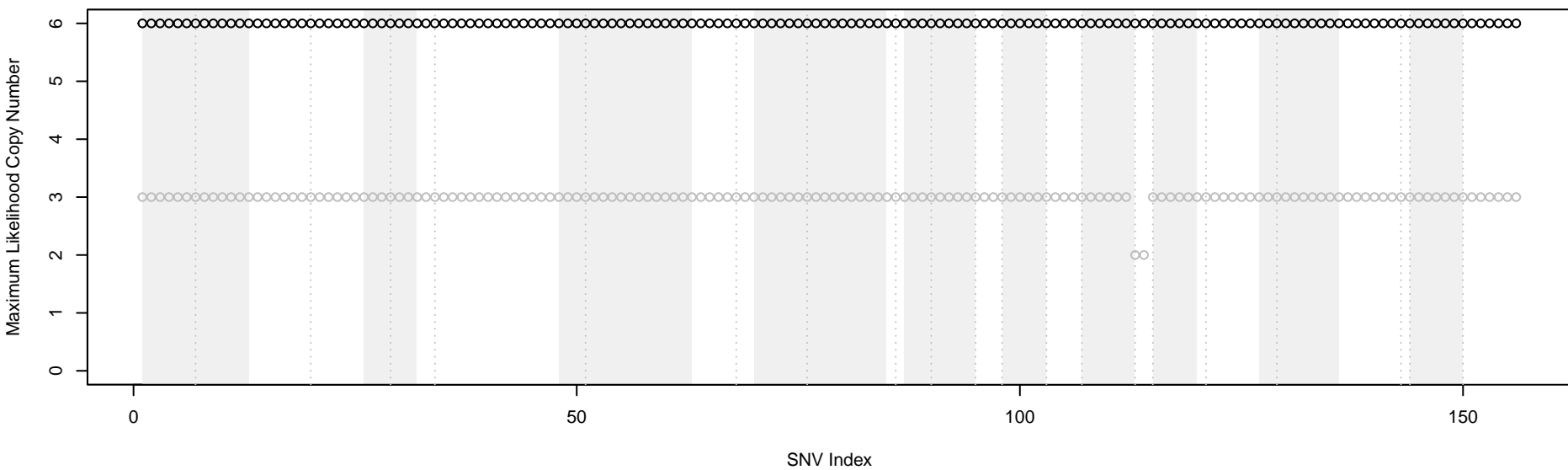
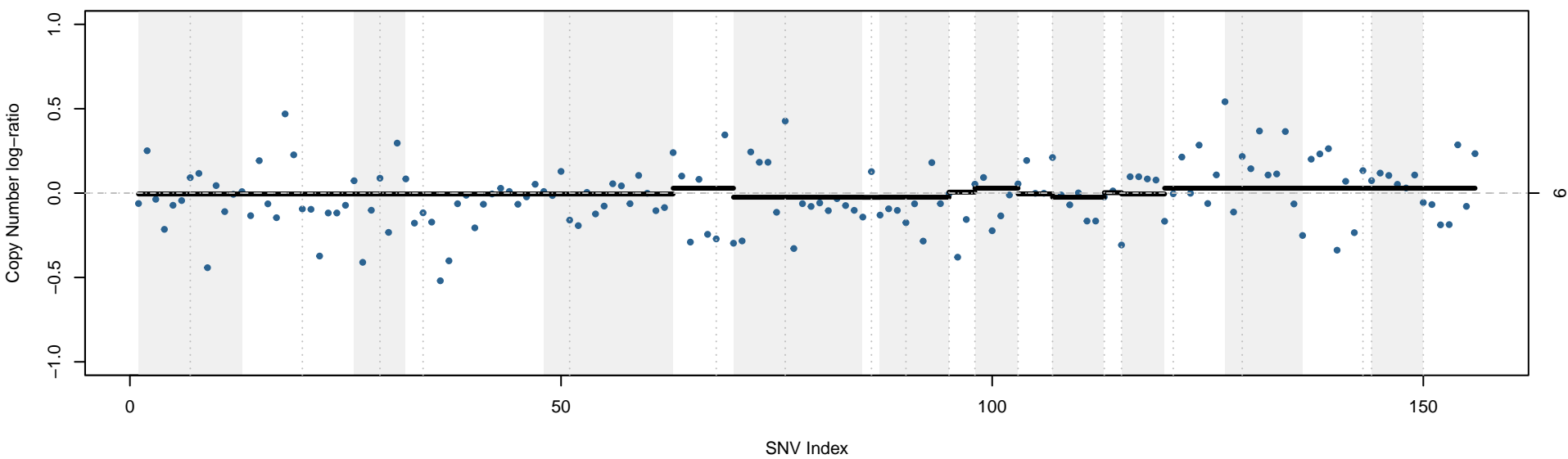


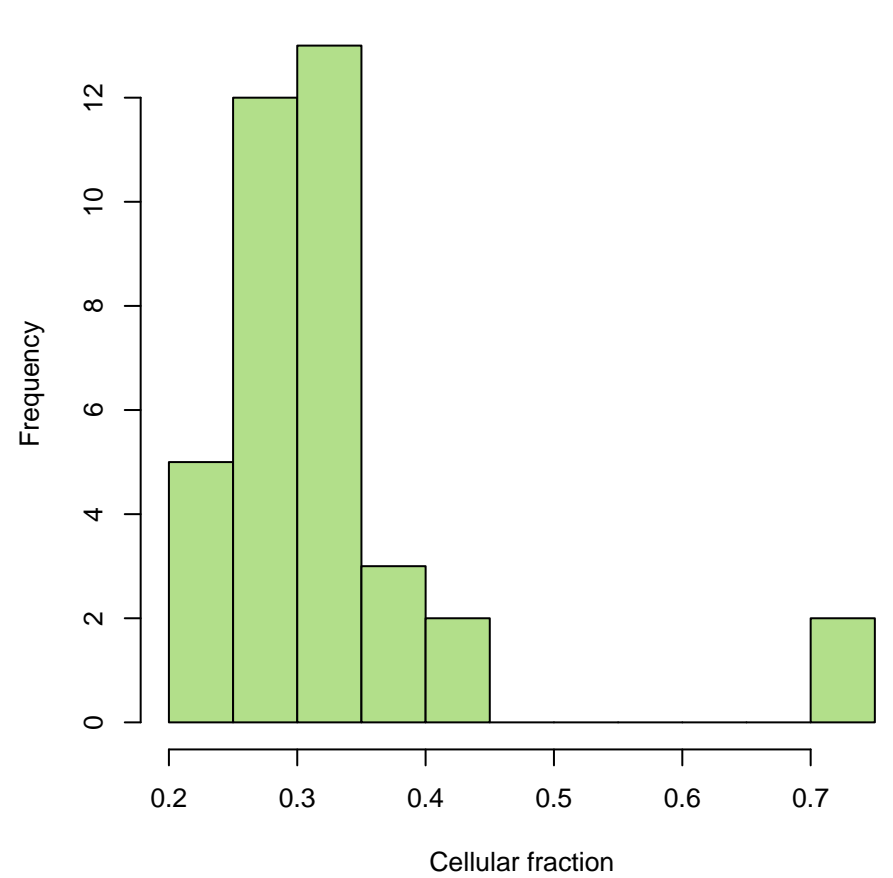
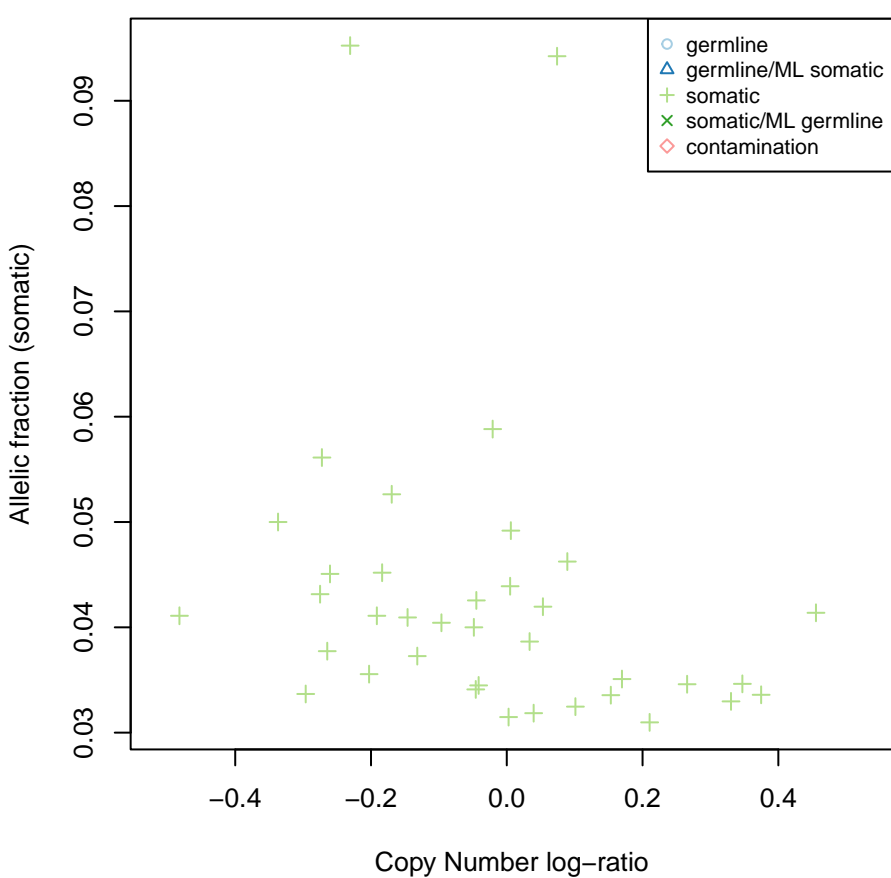
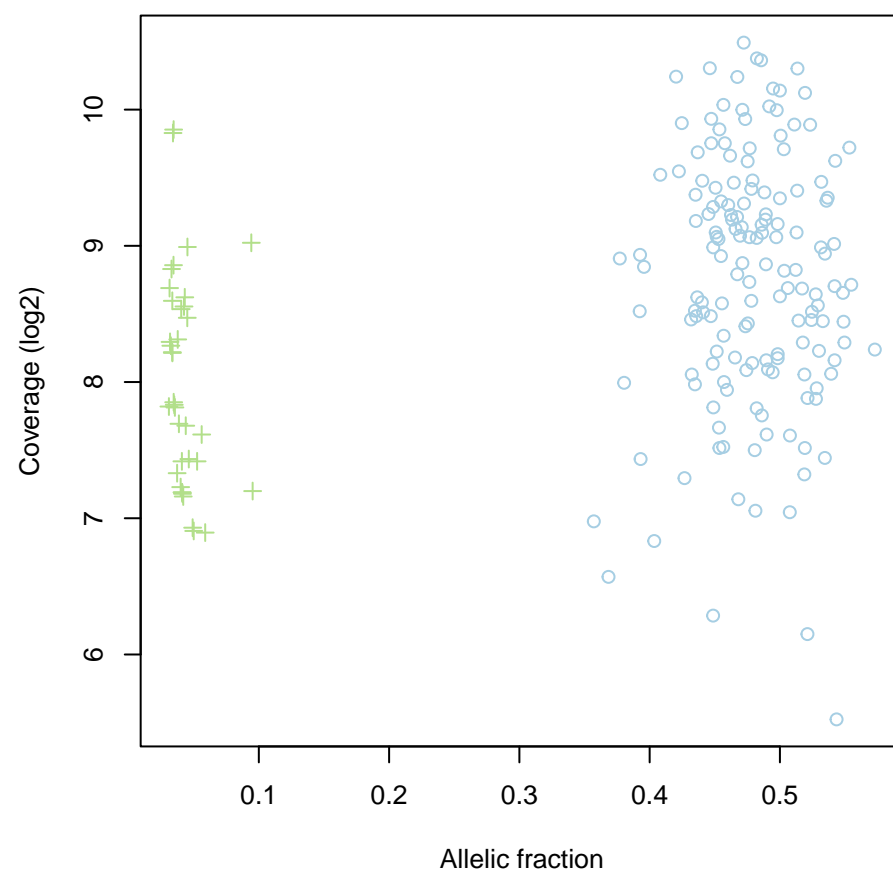
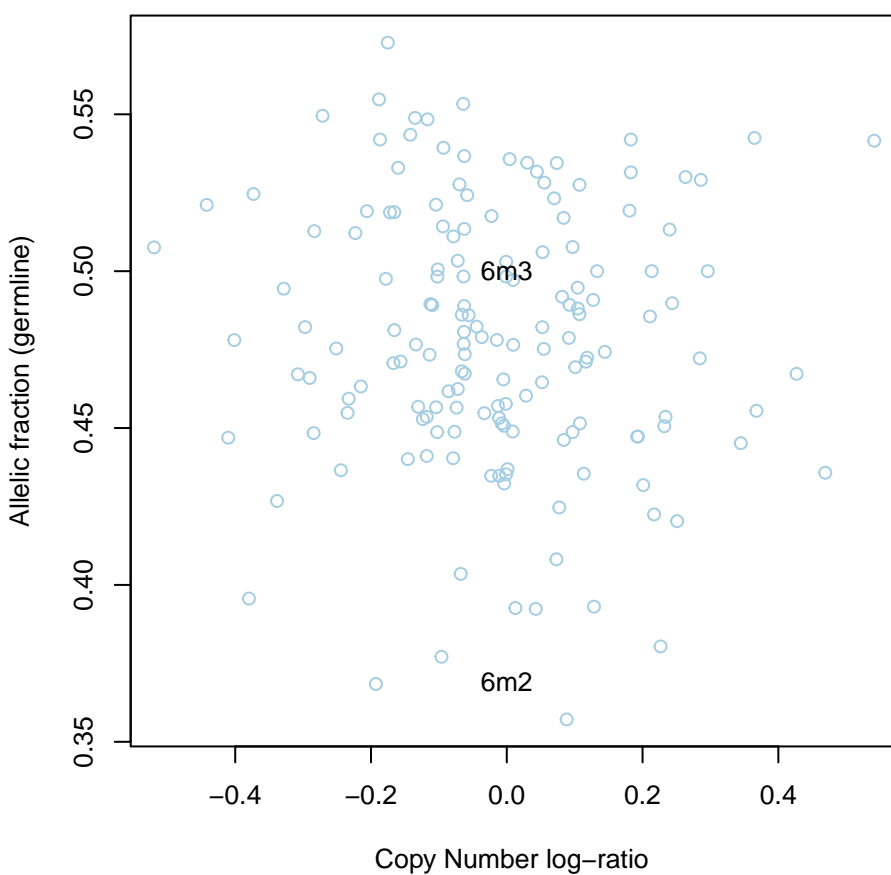


Purity: 0.55 Tumor ploidy: 6 SNV log-likelihood: -58.59 GoF: 92.4% Mean coverage: 410;477

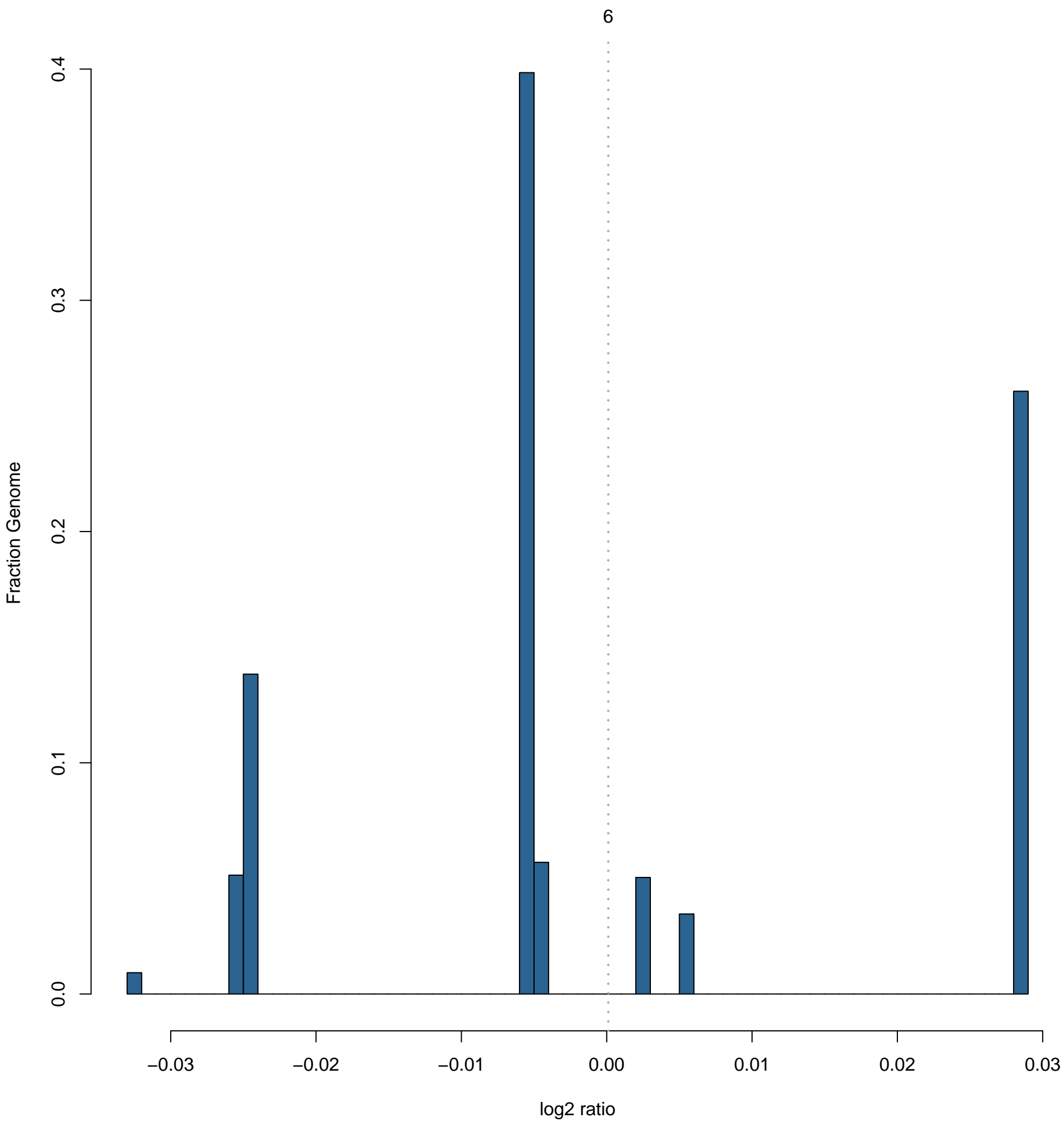


SCNA-fit log-likelihood: -7178.86

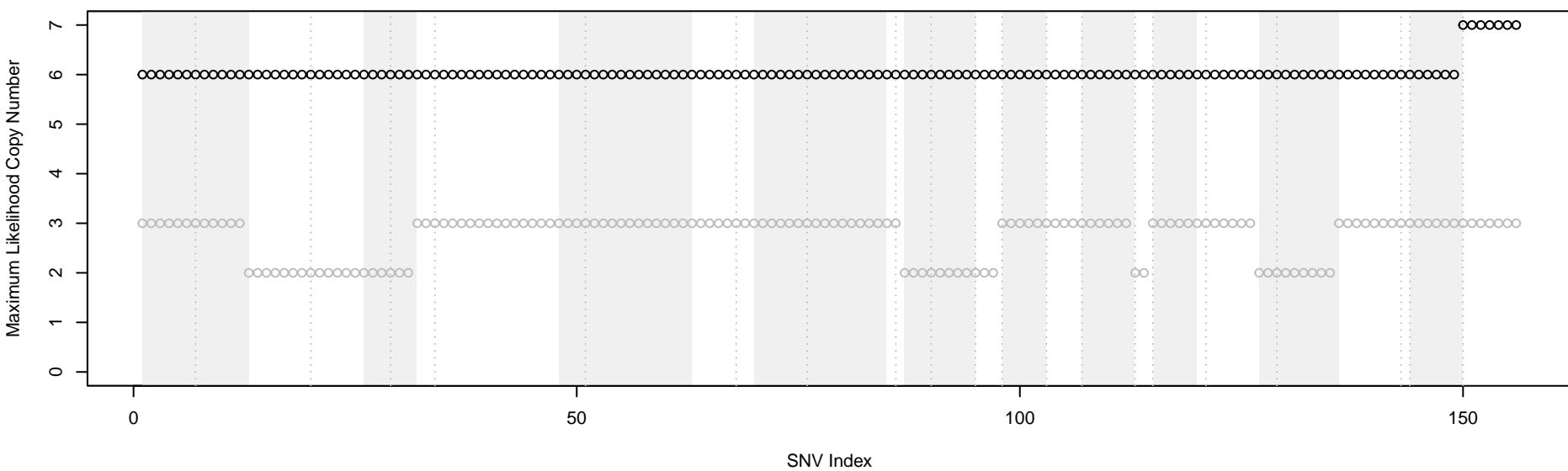
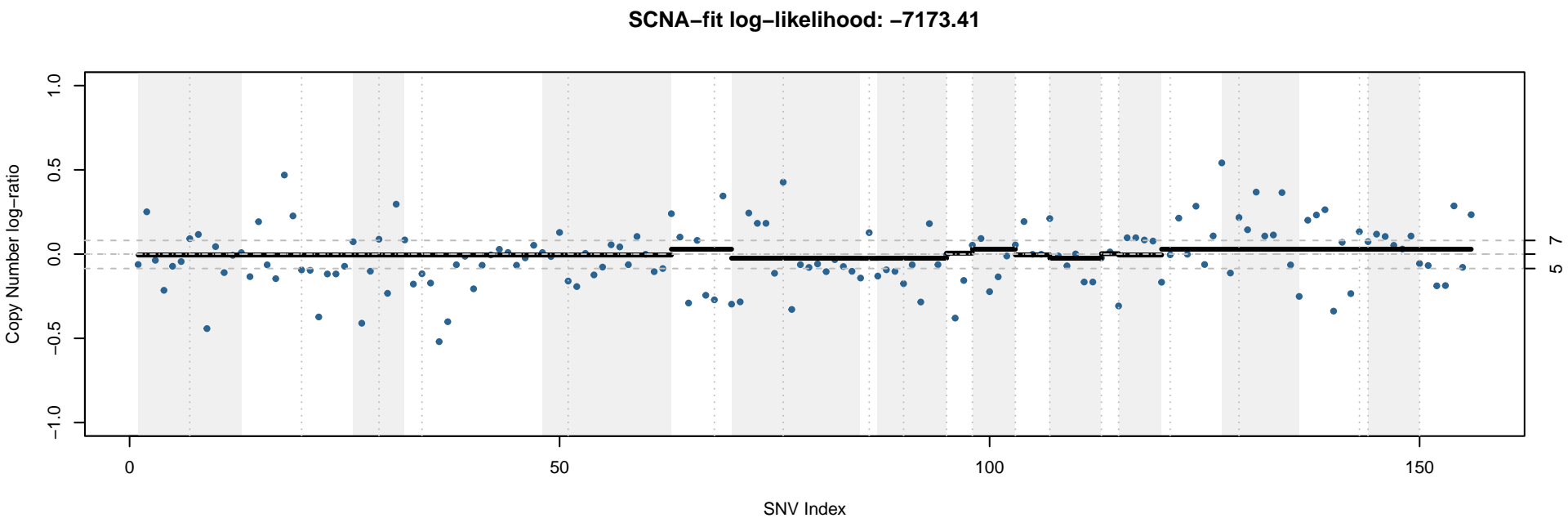


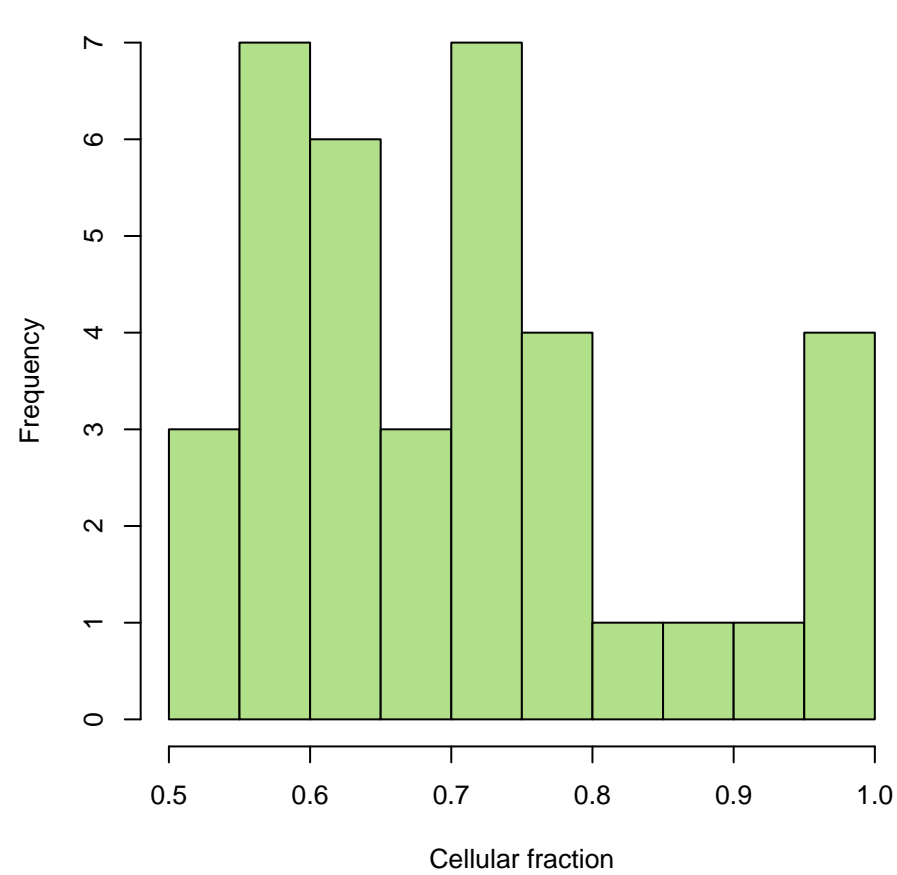
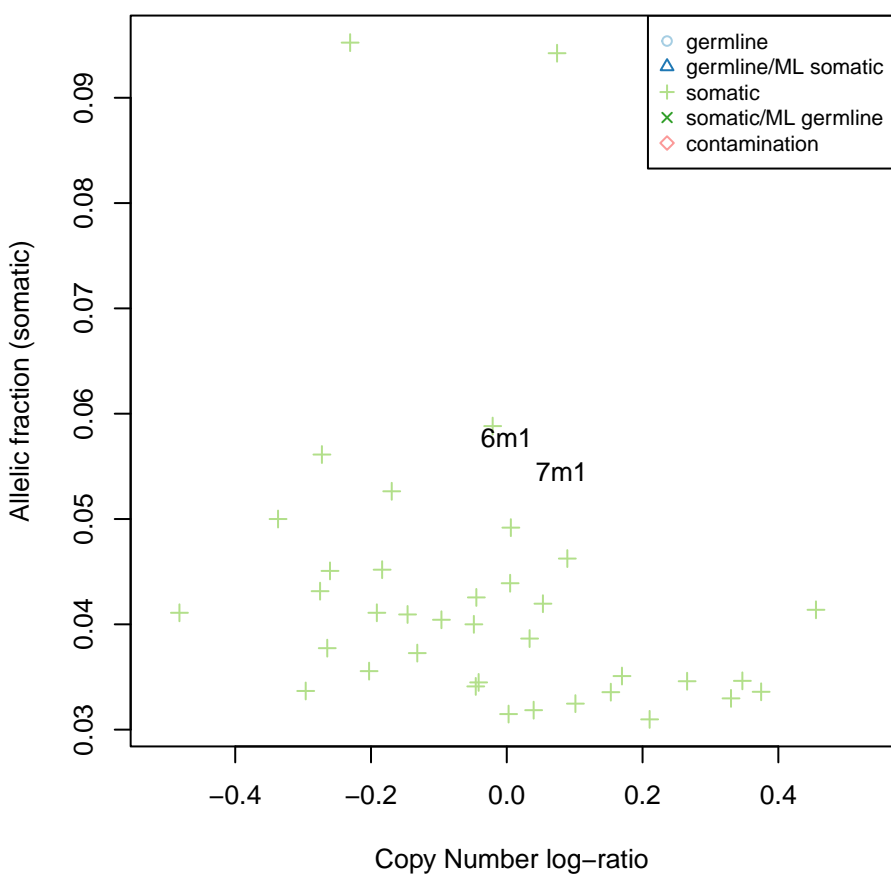
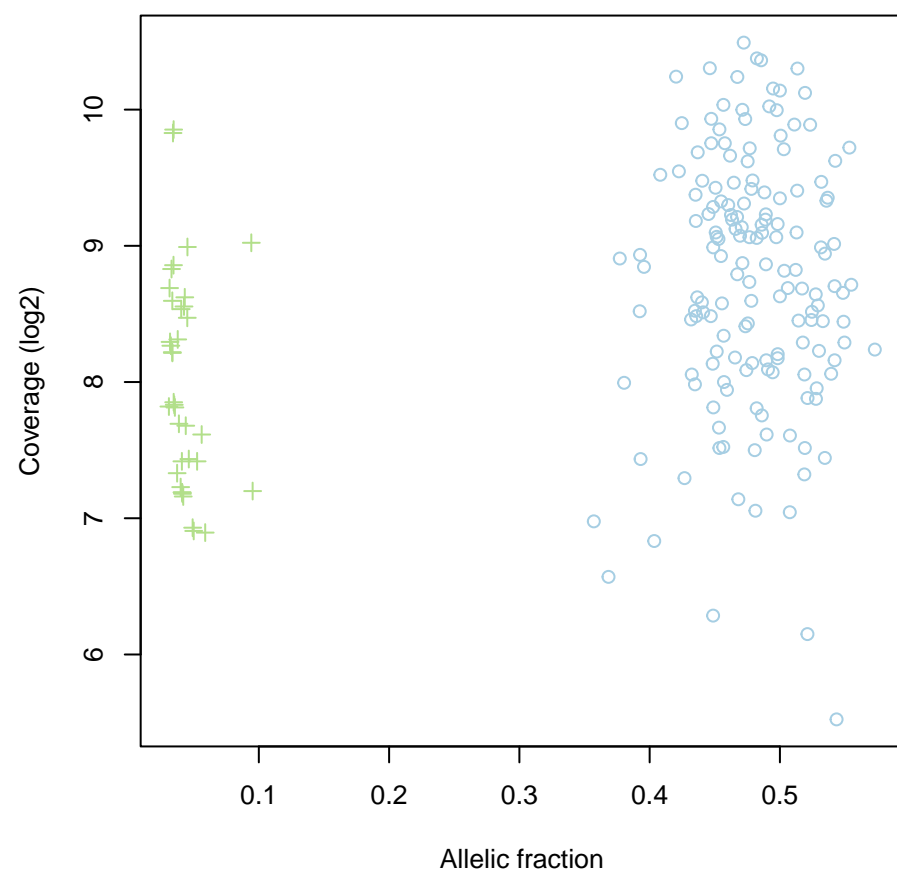
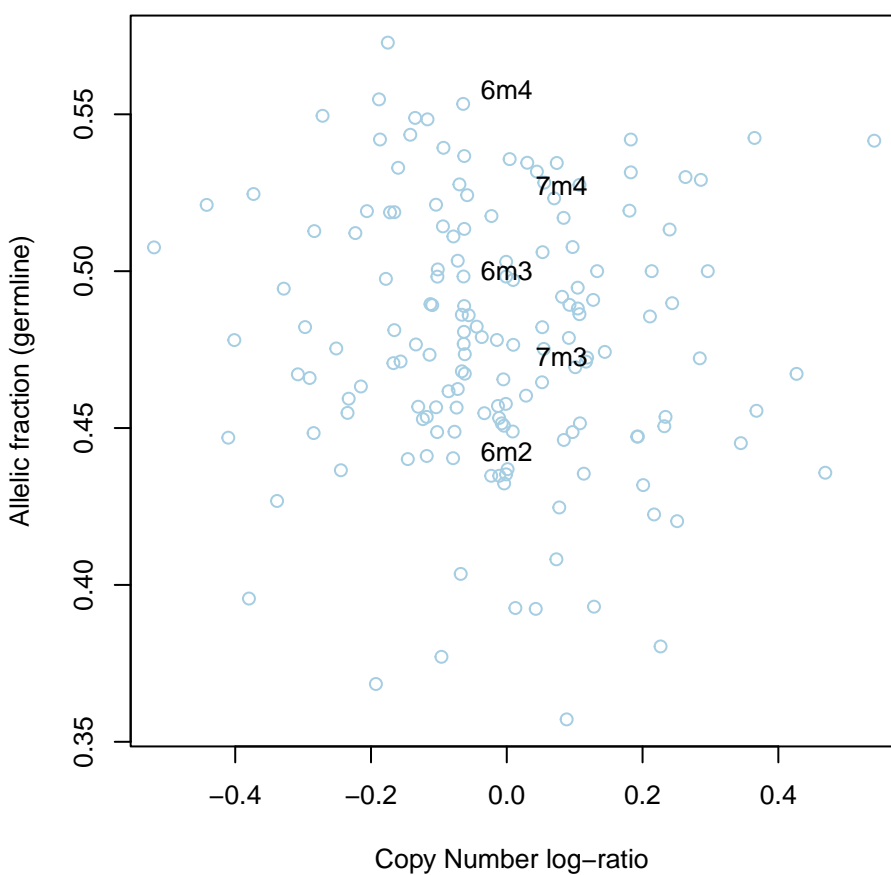


Purity: 0.15 Tumor ploidy: 5.999

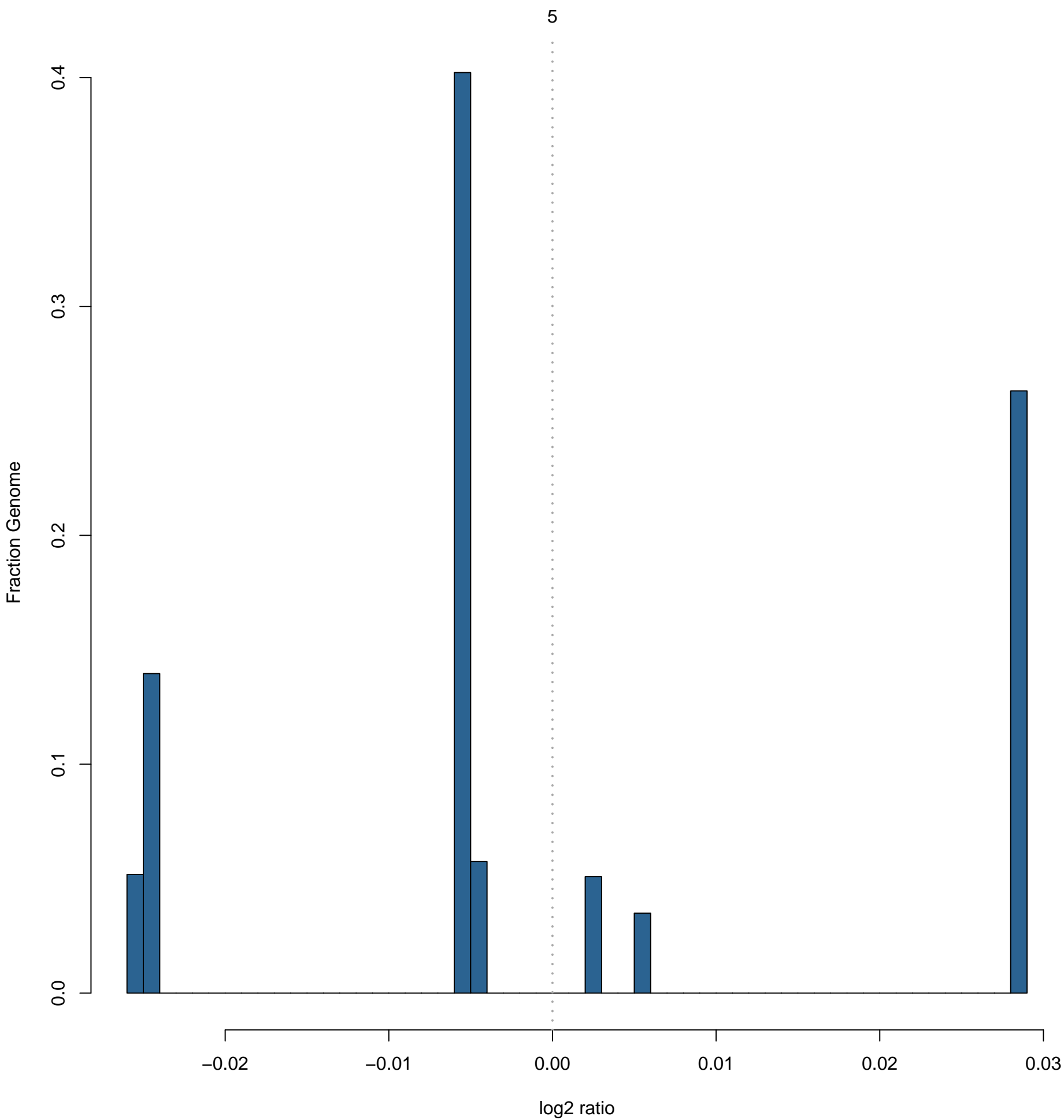


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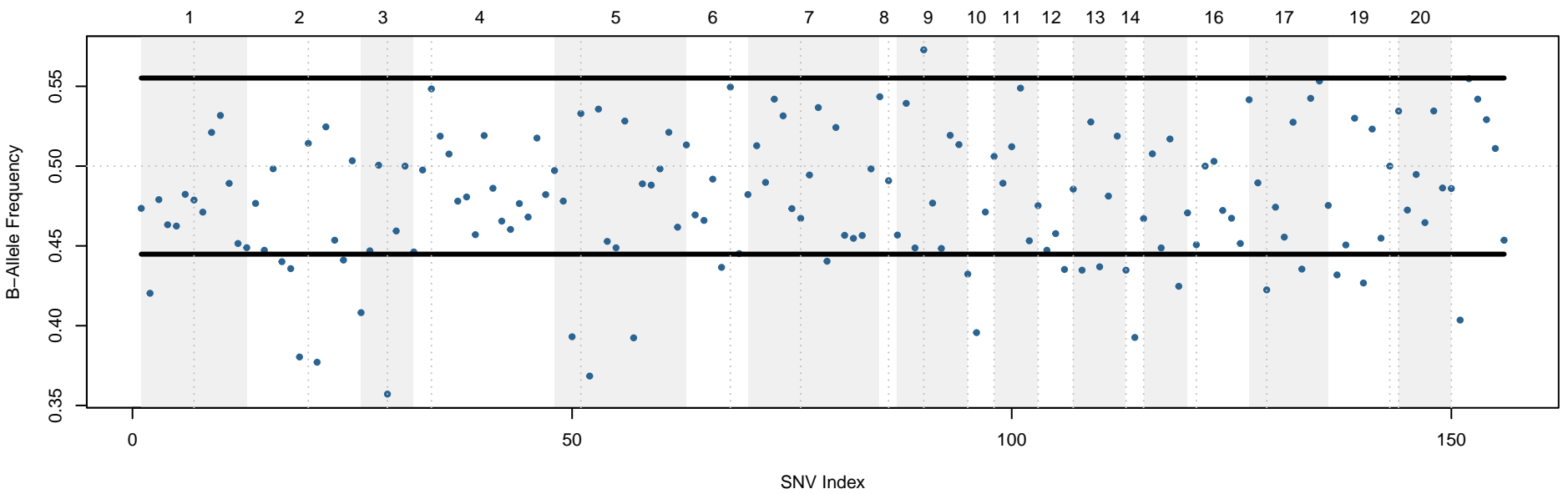




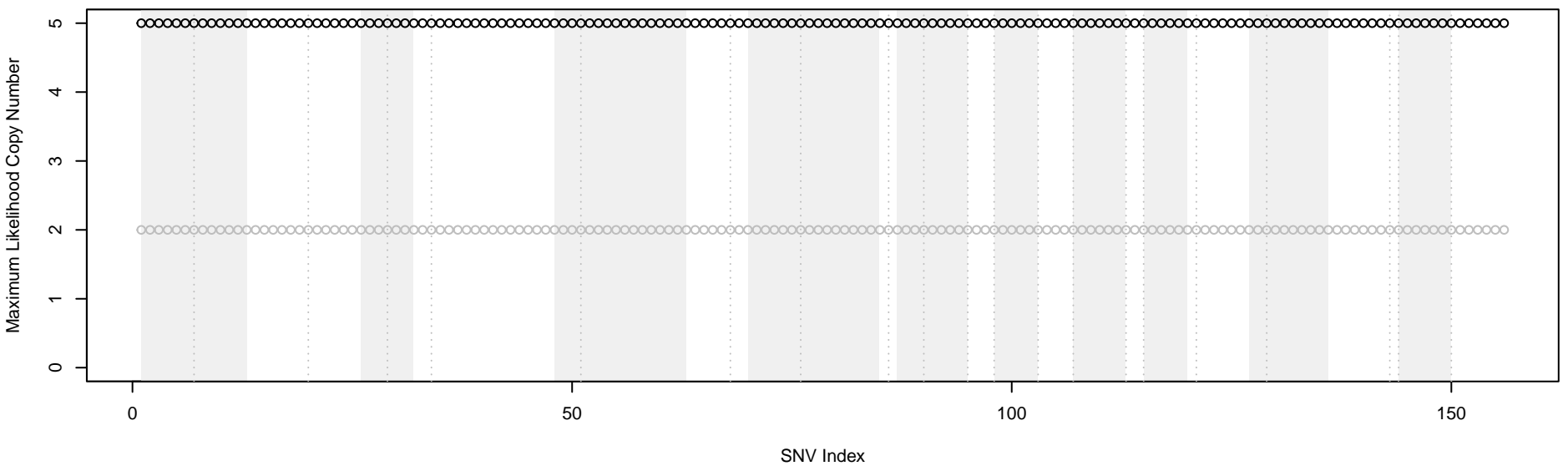
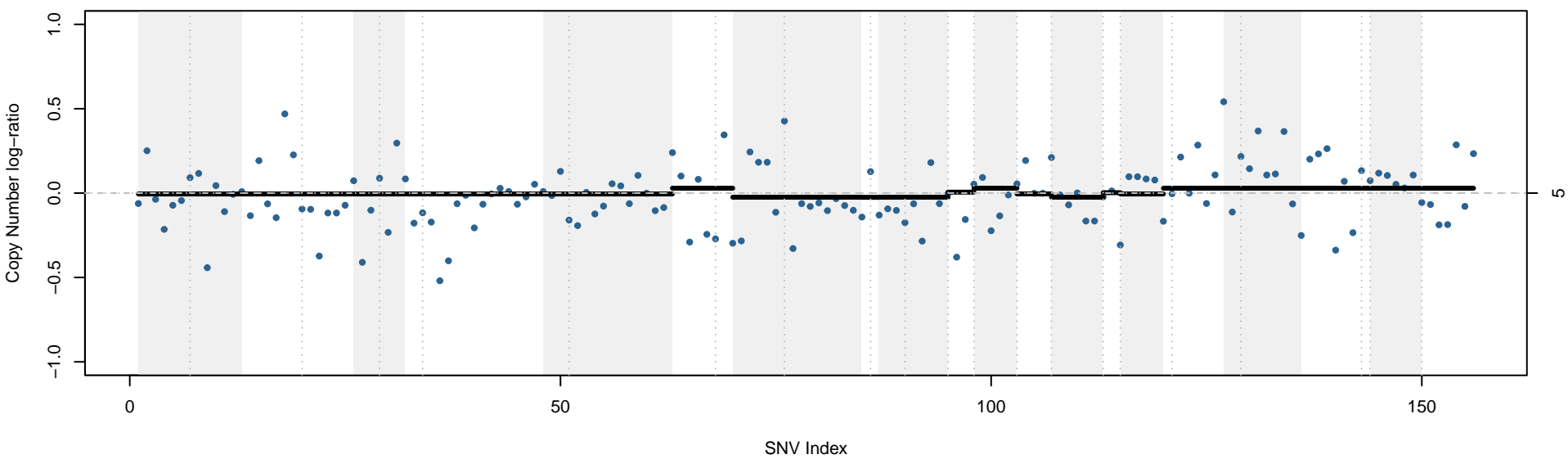
Purity: 0.33 Tumor ploidy: 5

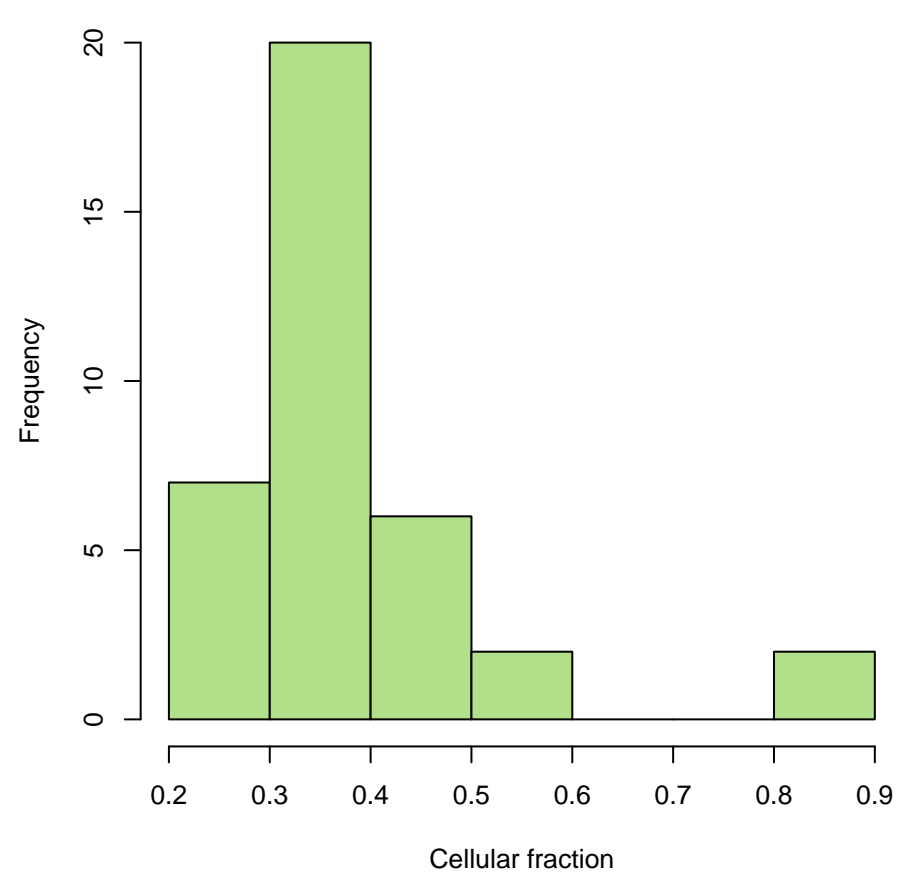
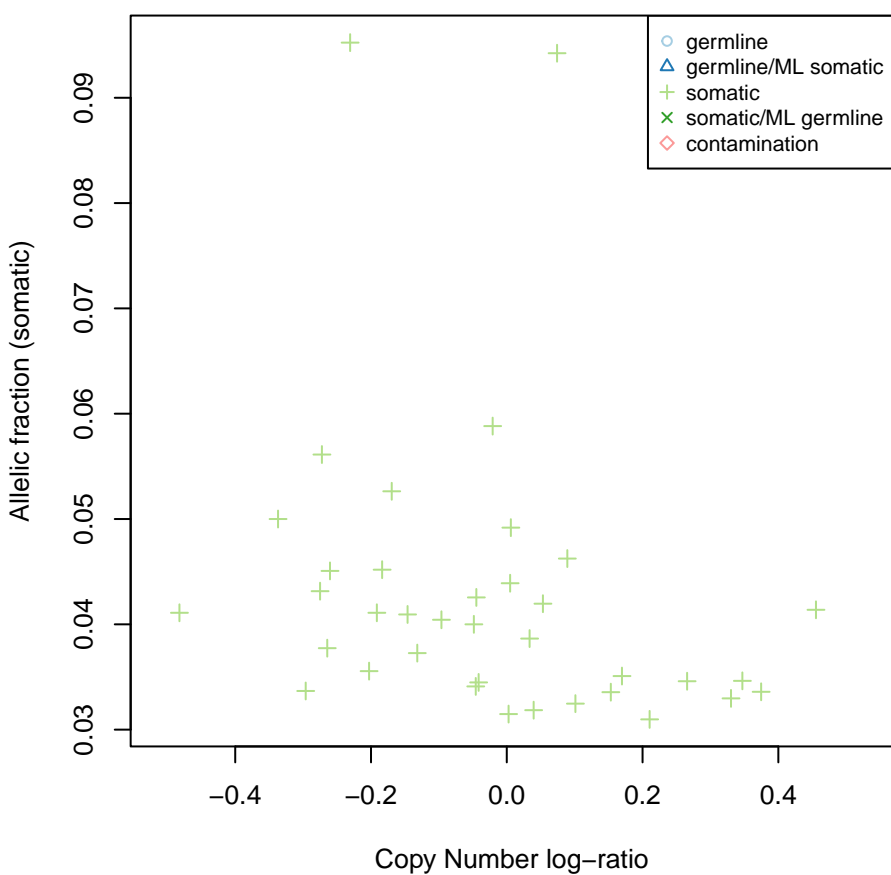
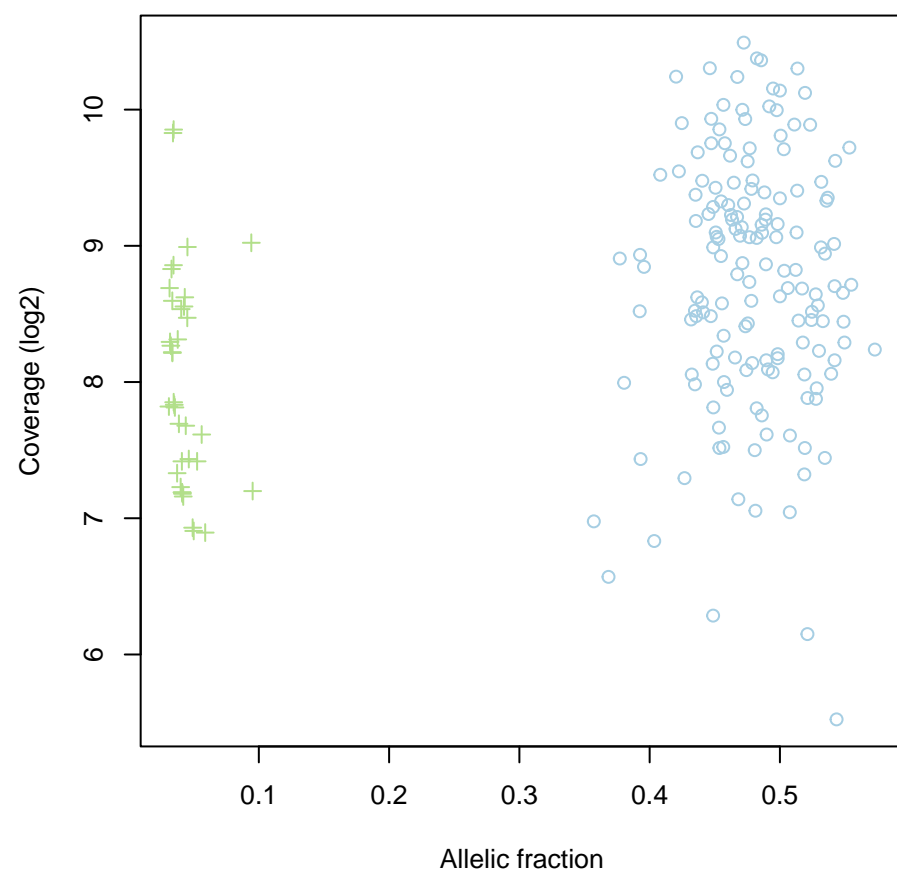
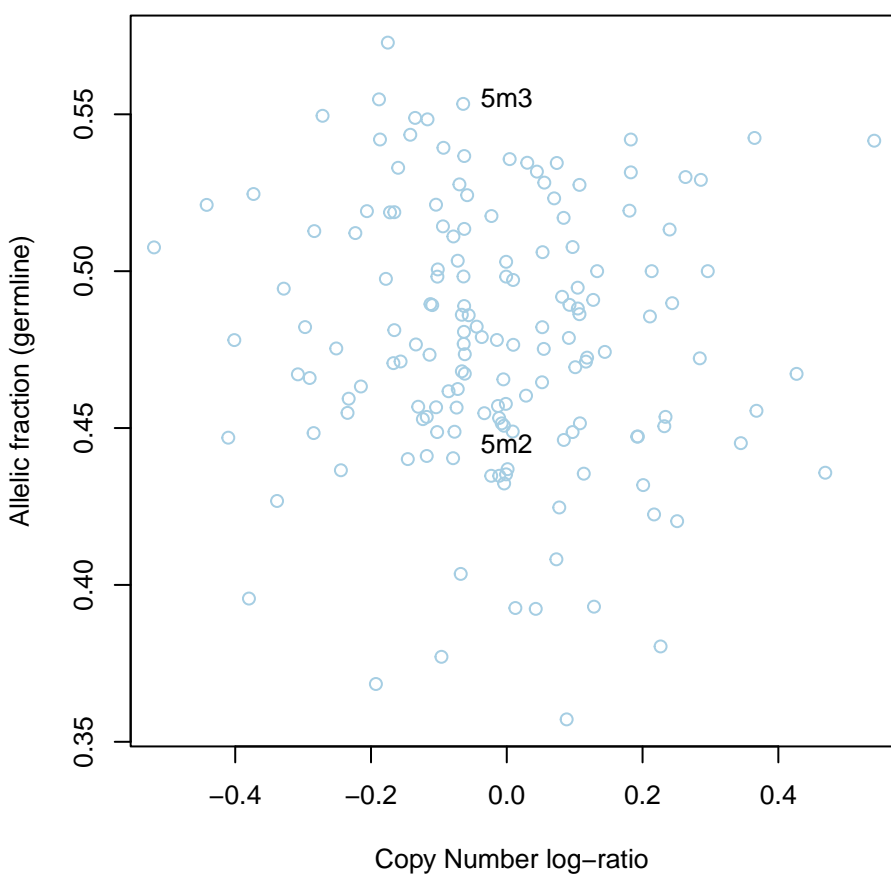


Purity: 0.33 Tumor ploidy: 5 SNV log-likelihood: -80.53 GoF: 95.6% Mean coverage: 410;477



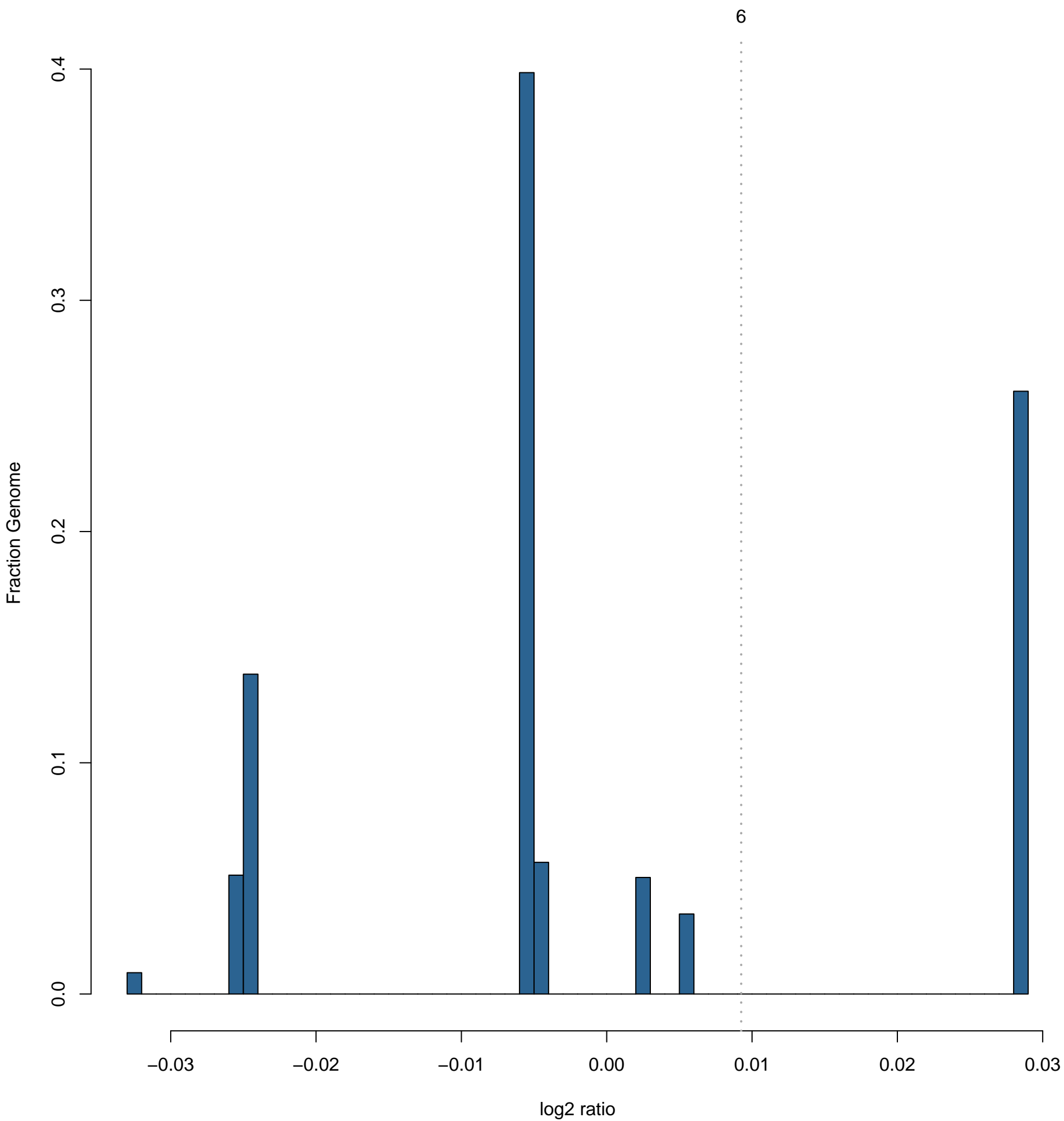
SCNA-fit log-likelihood: -7178.85



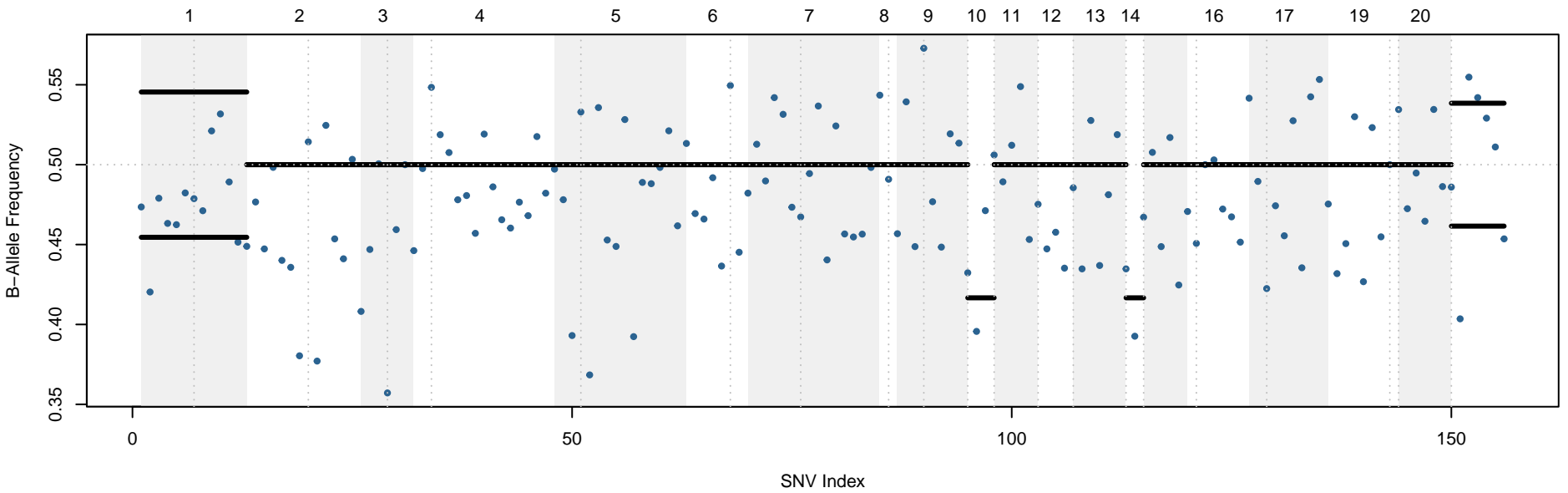




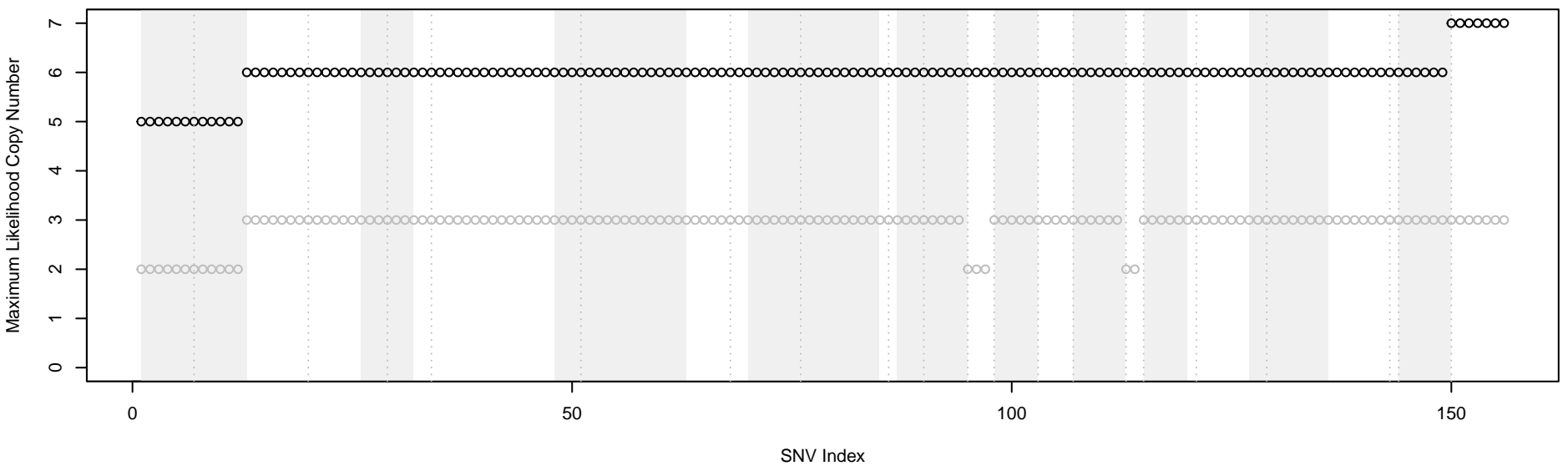
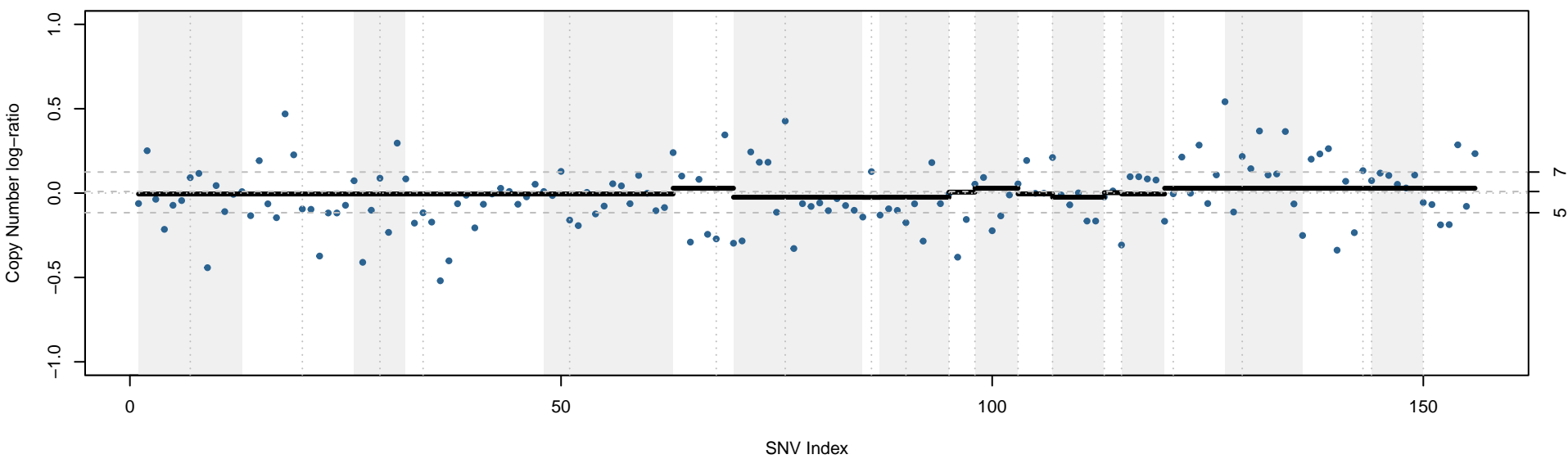
Purity: 0.25 Tumor ploidy: 5.923

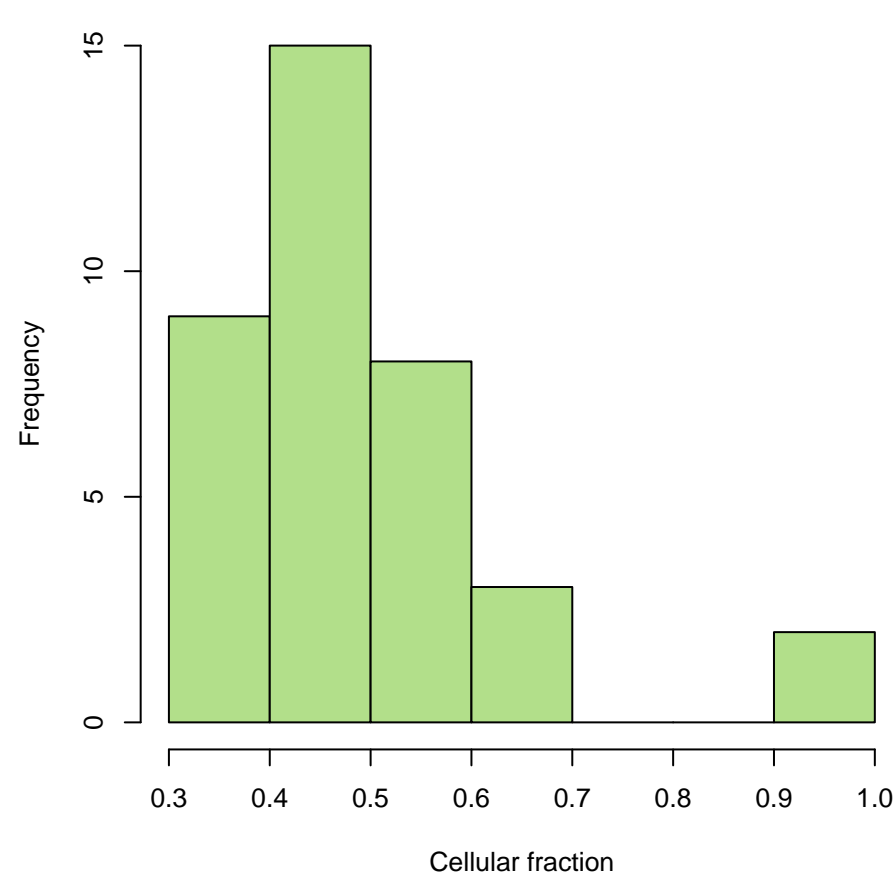
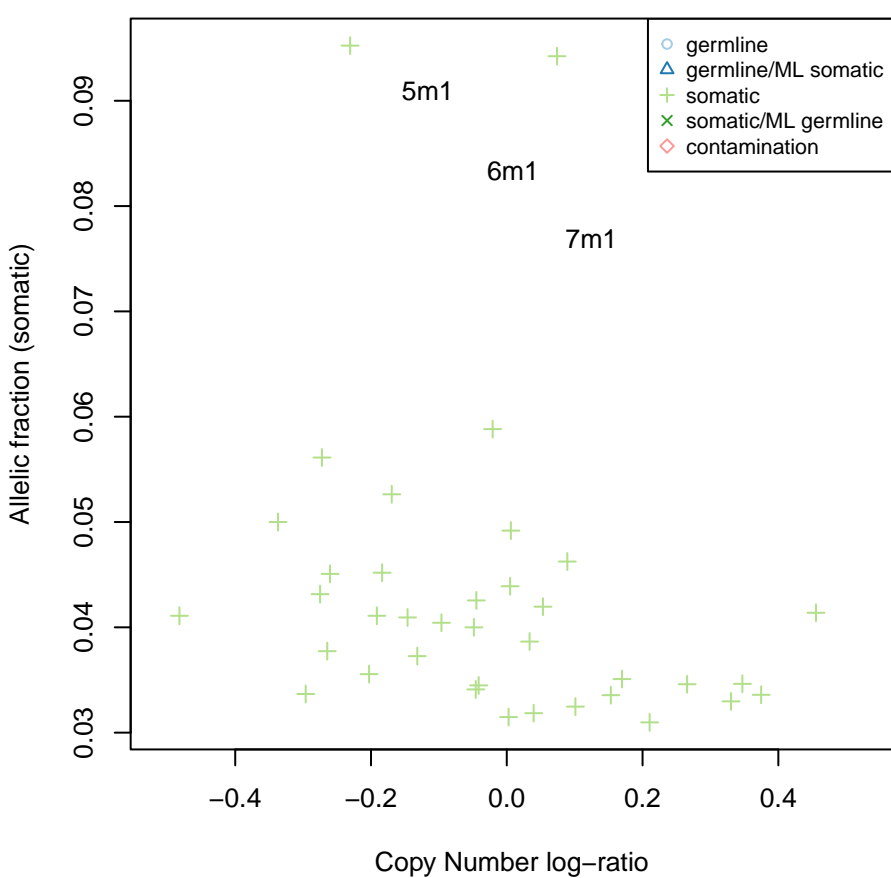
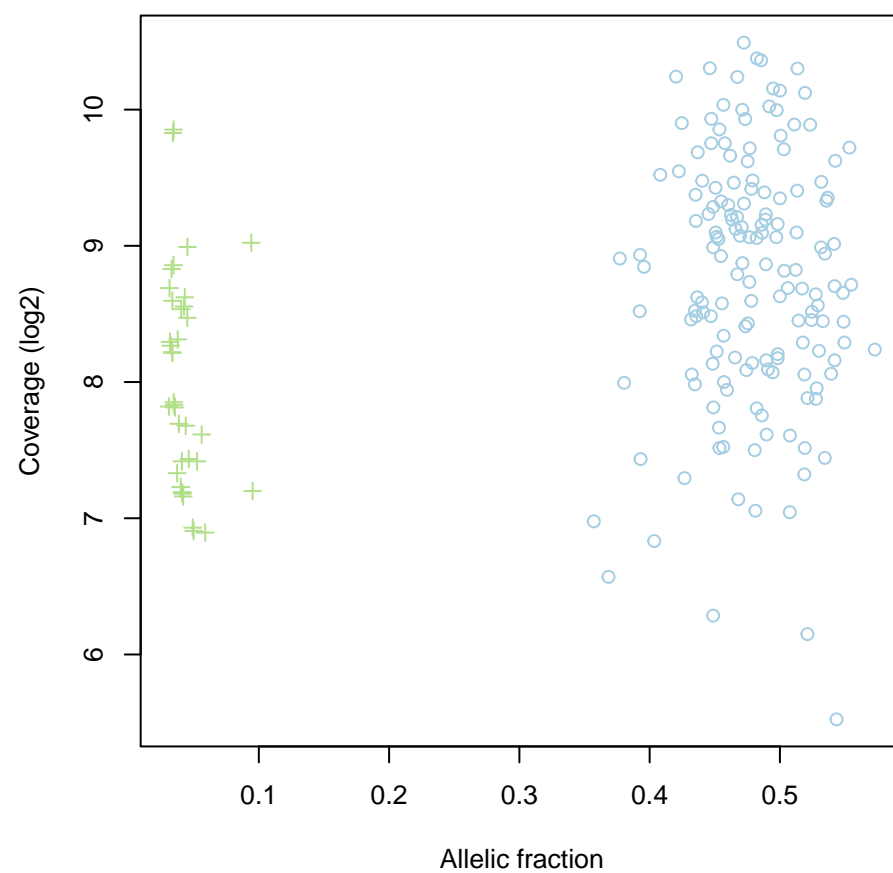
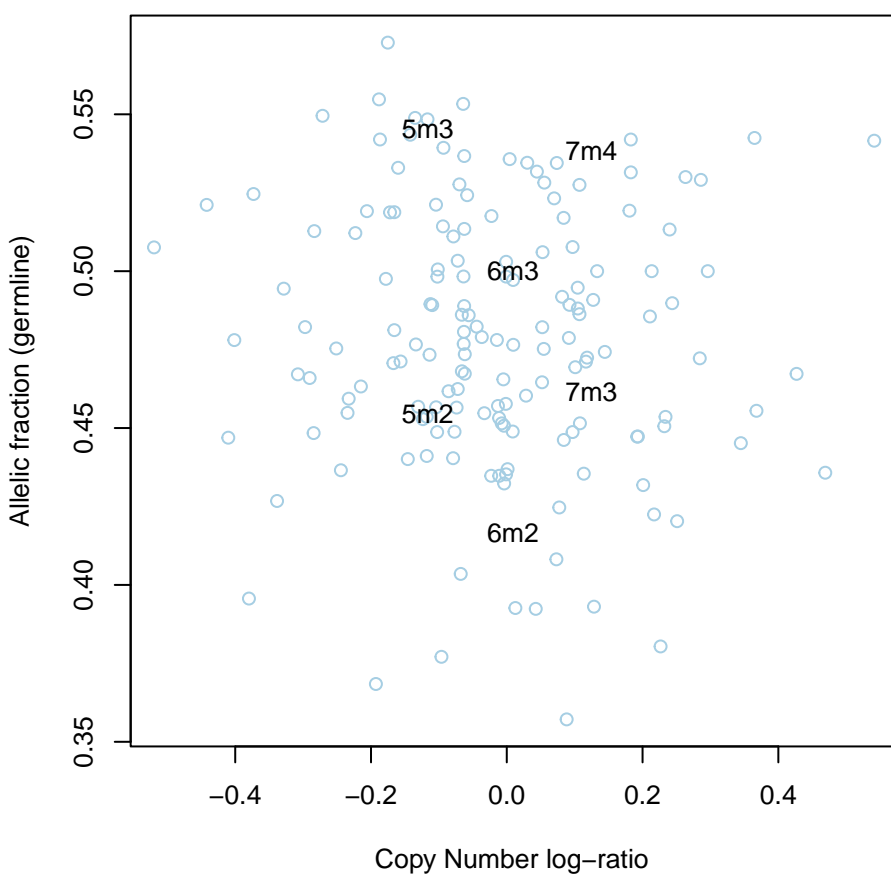


Purity: 0.25 Tumor ploidy: 5.923 SNV log-likelihood: -66.18 GoF: 95.7% Mean coverage: 410;477

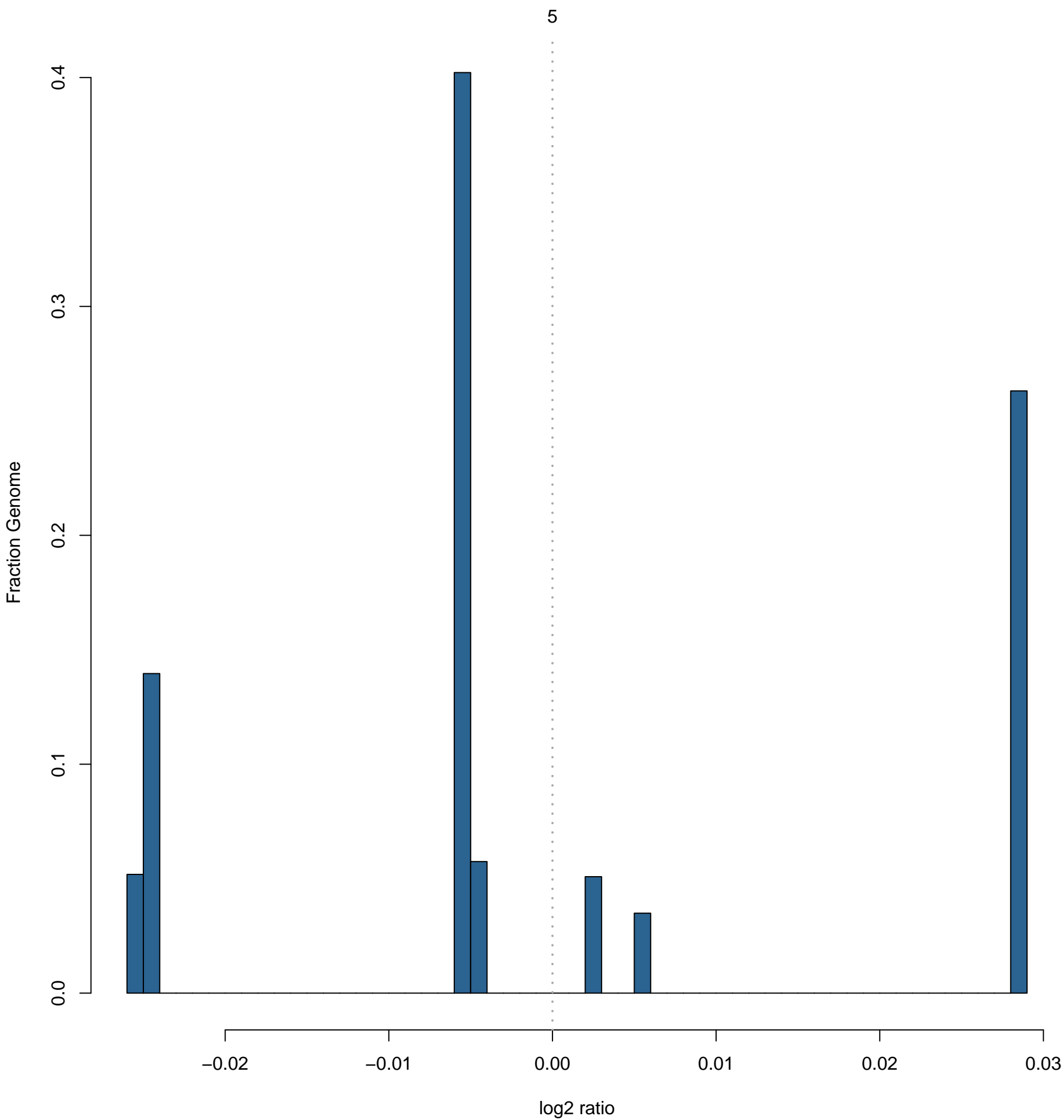


SCNA-fit log-likelihood: -7261.63

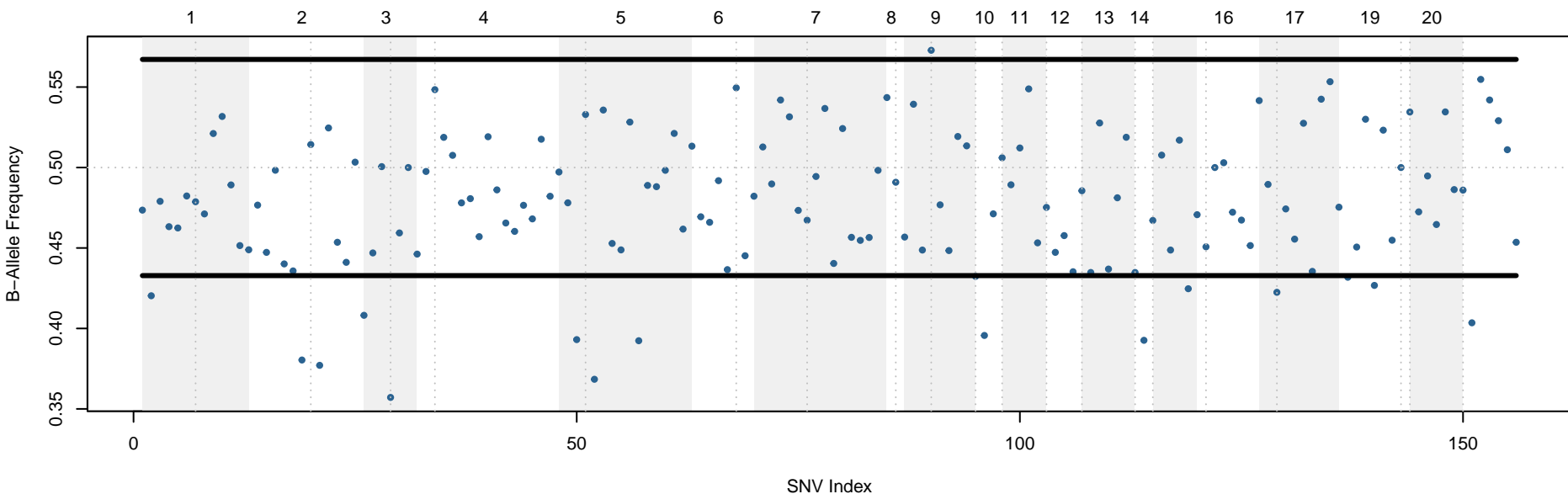




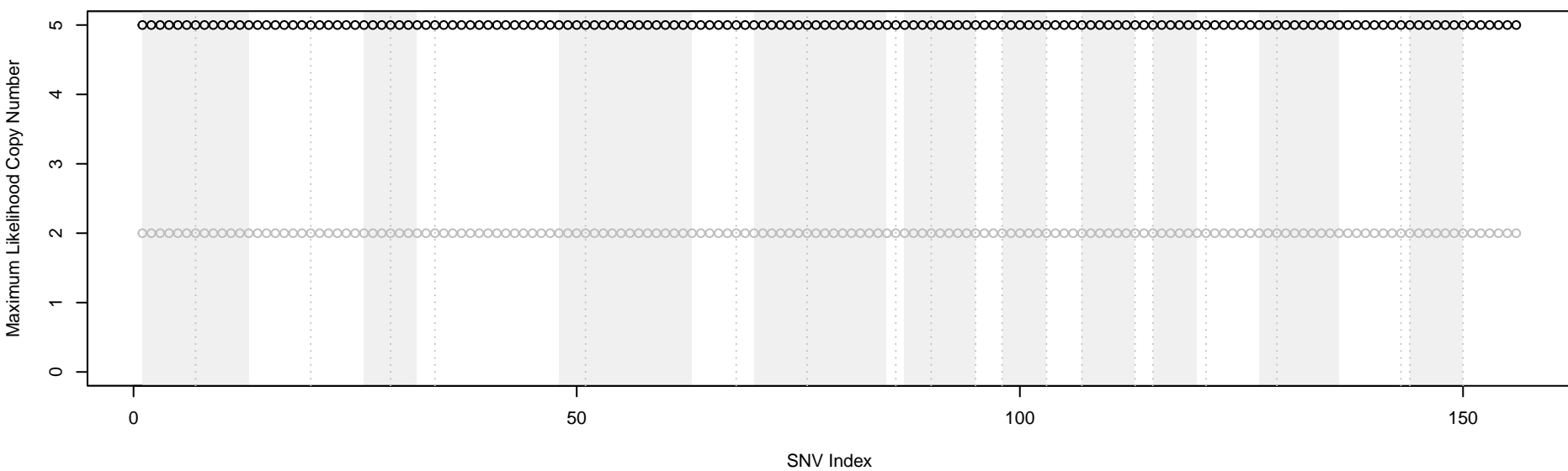
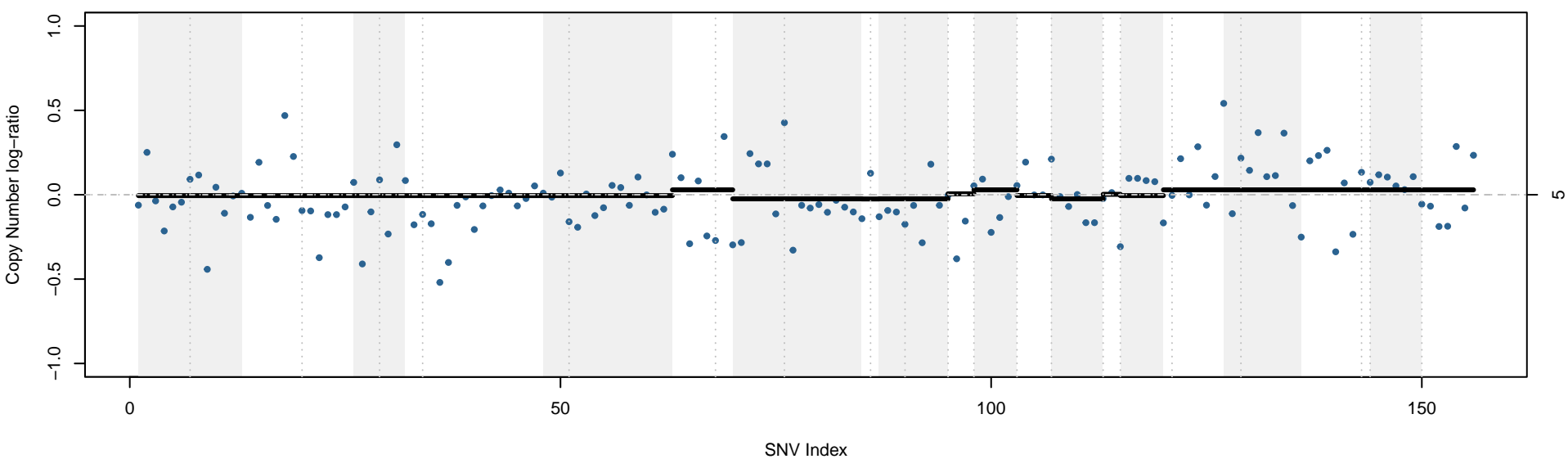
Purity: 0.45 Tumor ploidy: 5

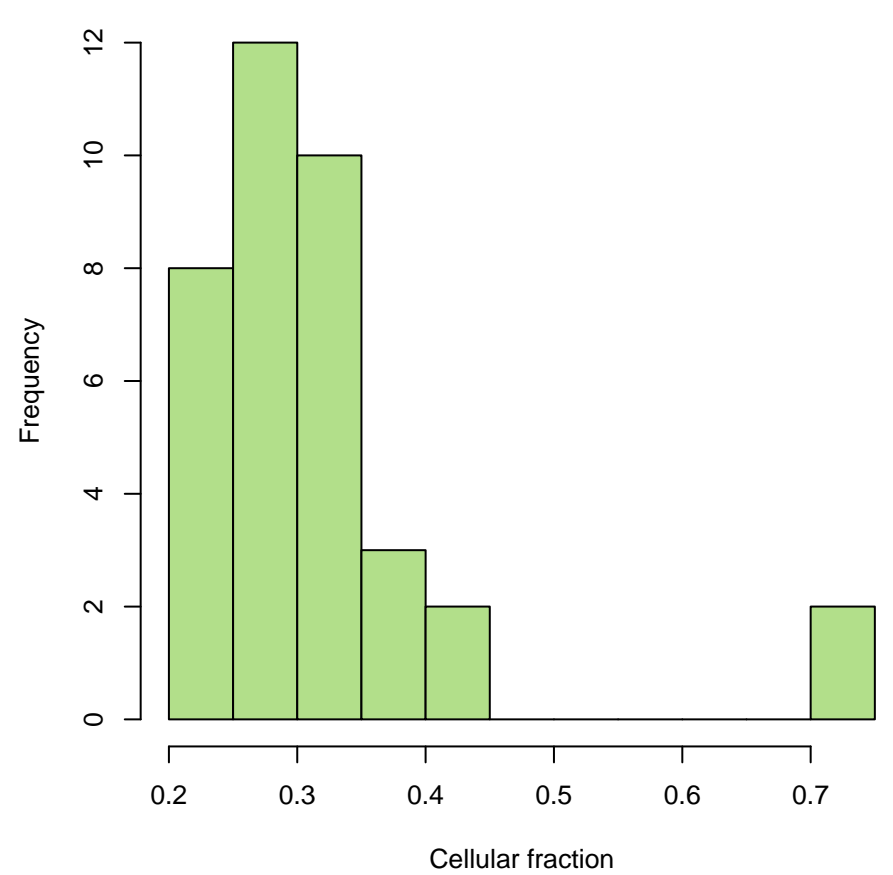
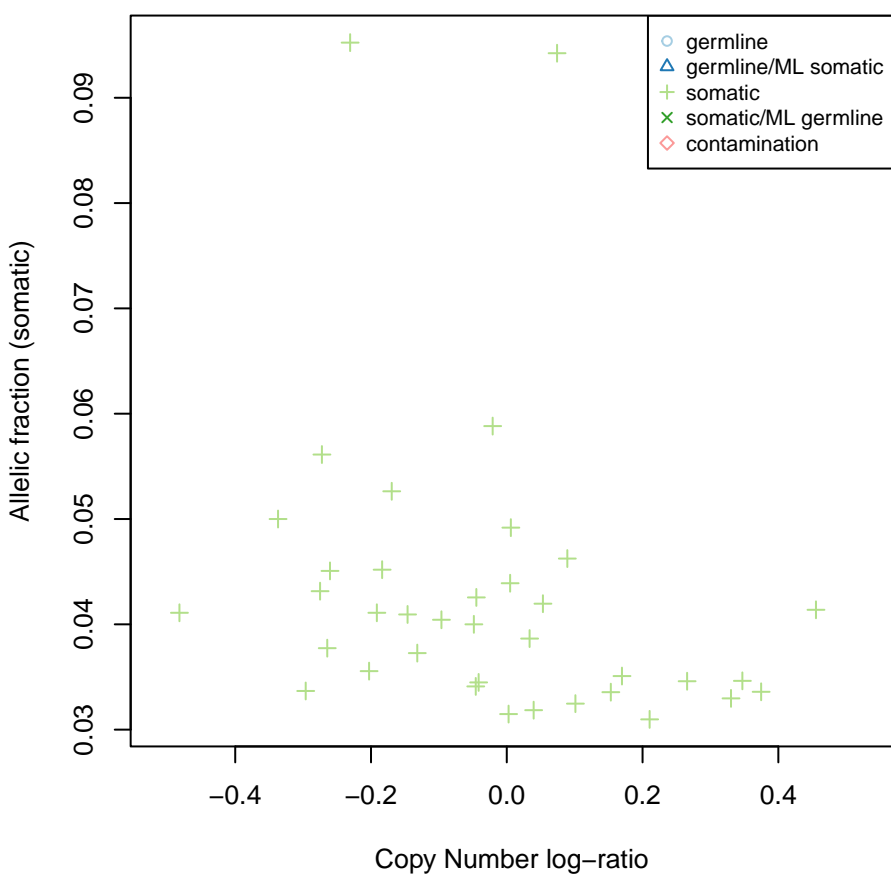
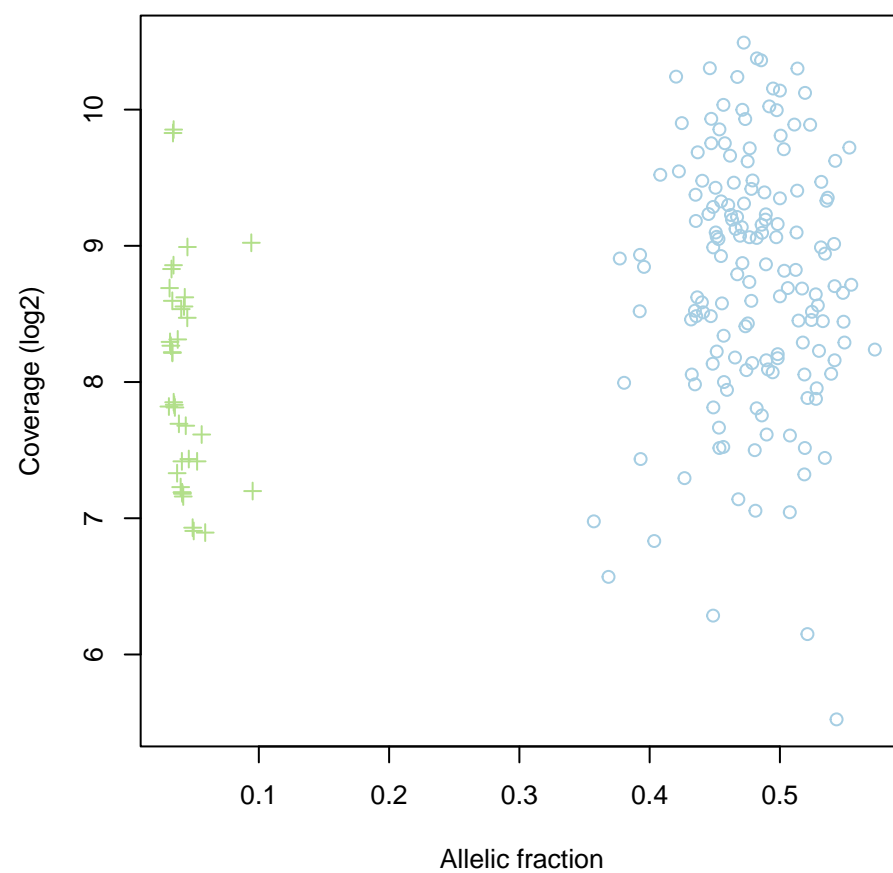
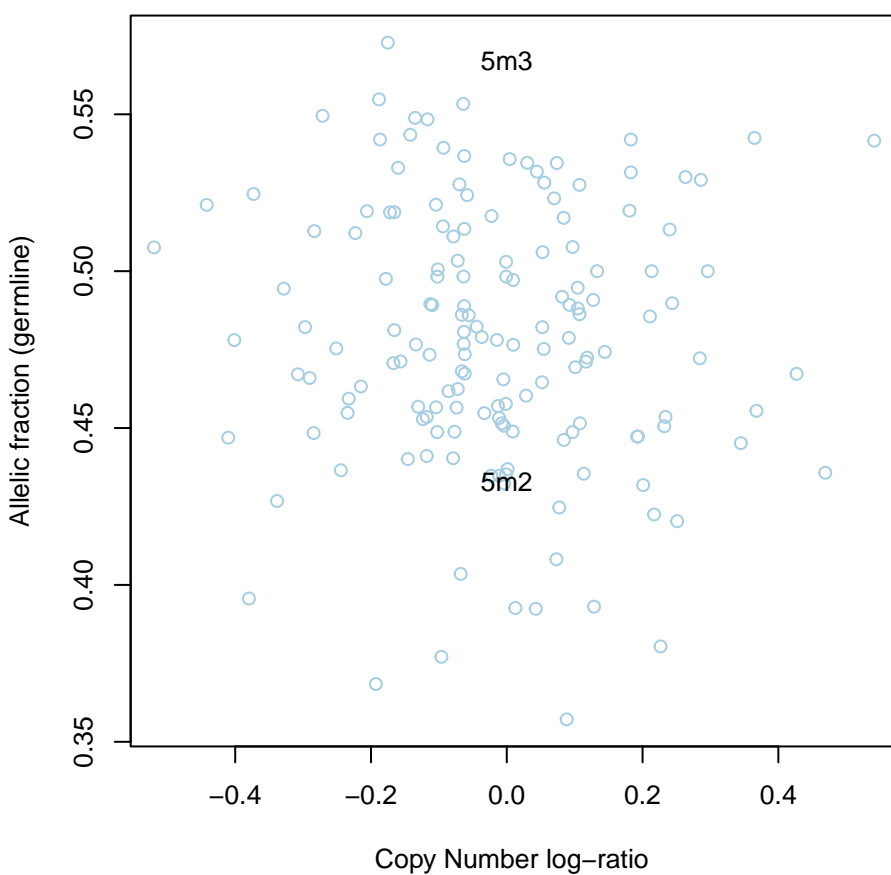


Purity: 0.45 Tumor ploidy: 5 SNV log-likelihood: -141.31 GoF: 92.6% Mean coverage: 410;477

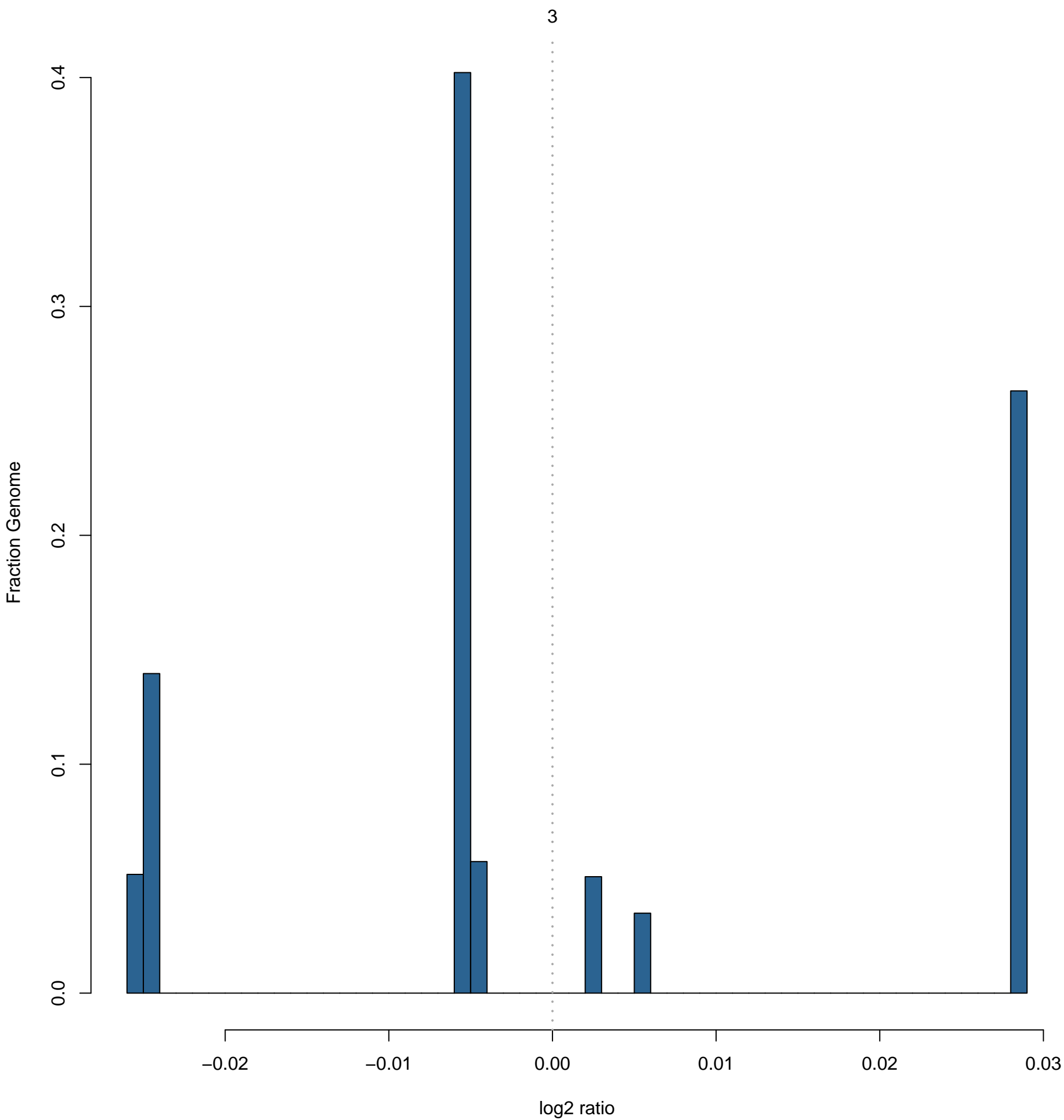


SCNA-fit log-likelihood: -7181.94

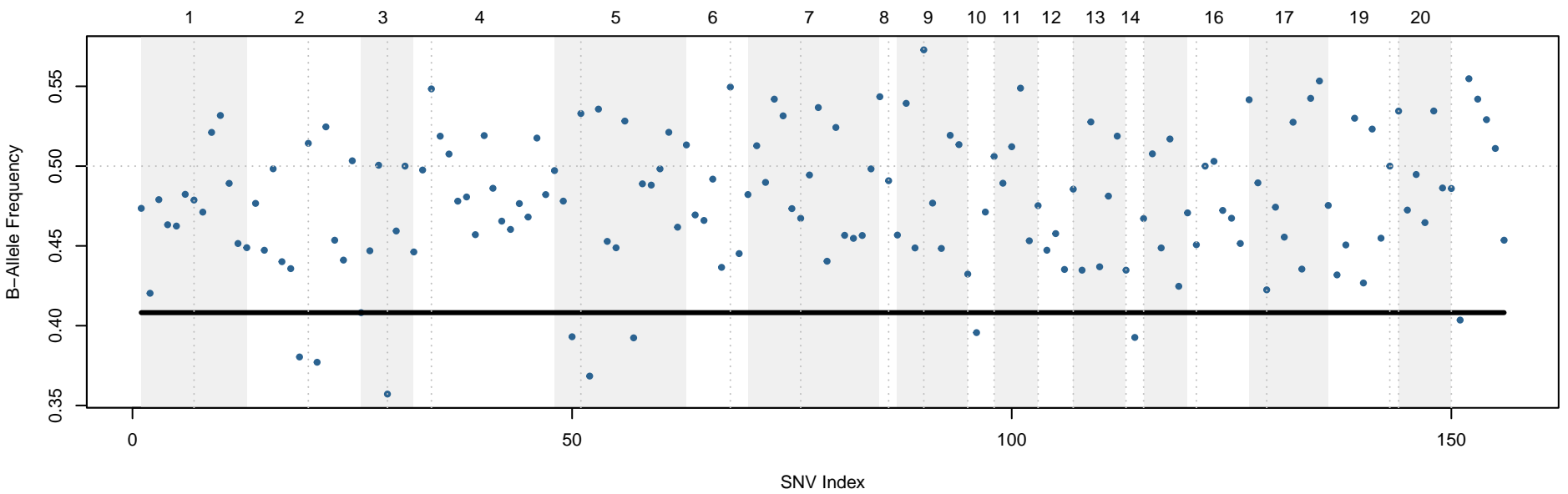




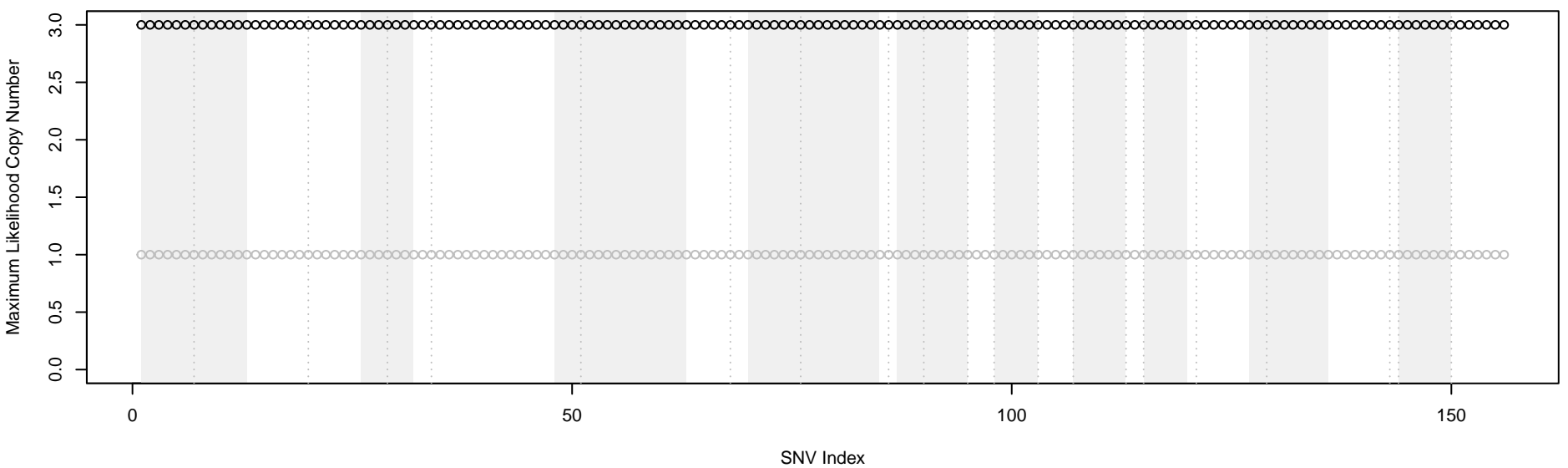
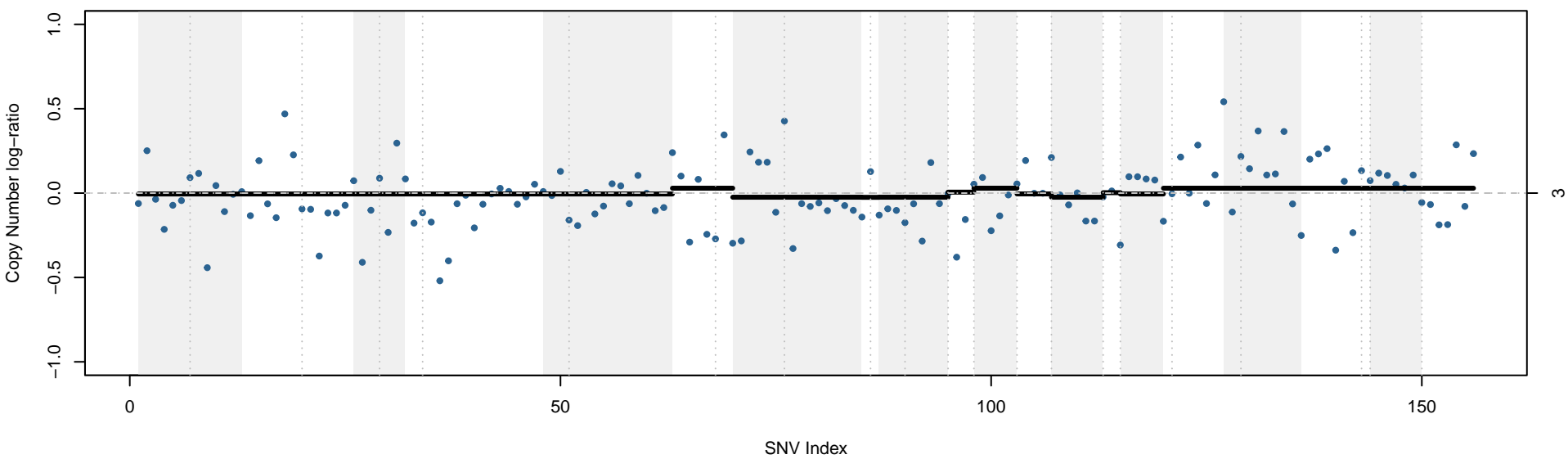
Purity: 0.45 Tumor ploidy: 3



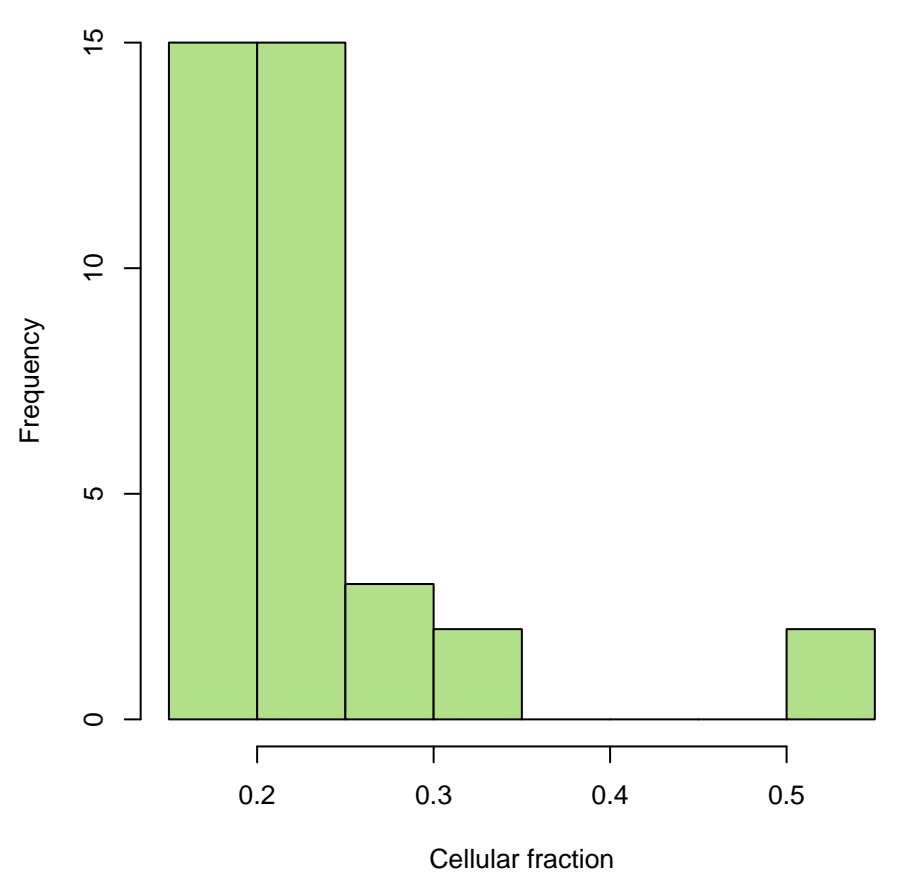
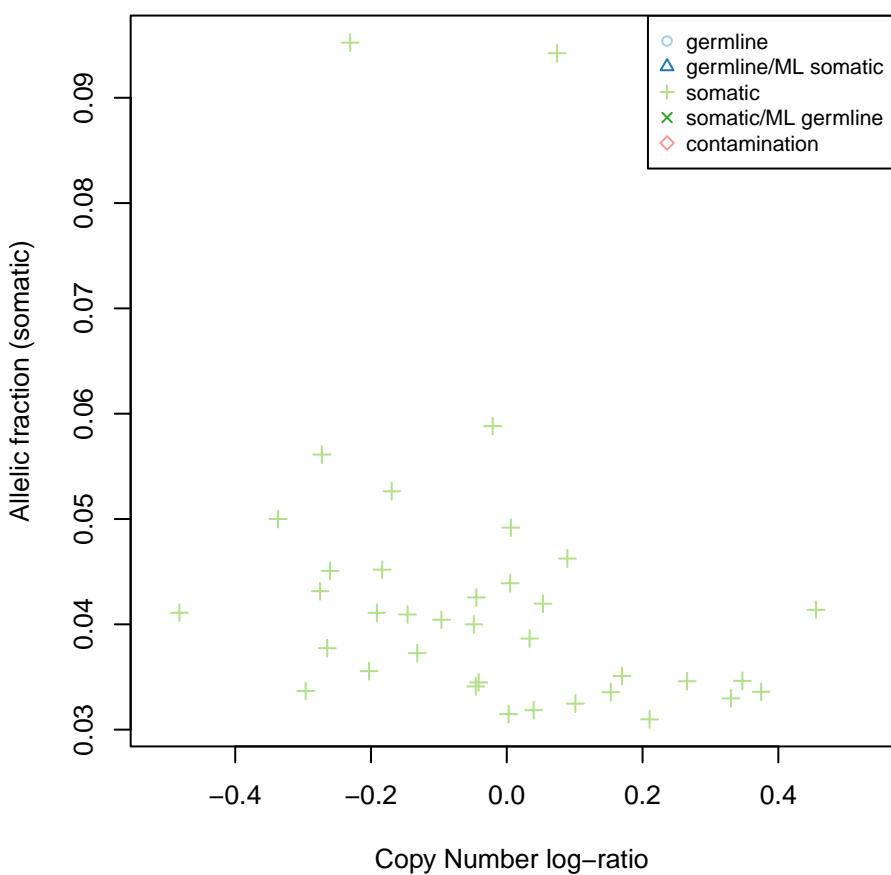
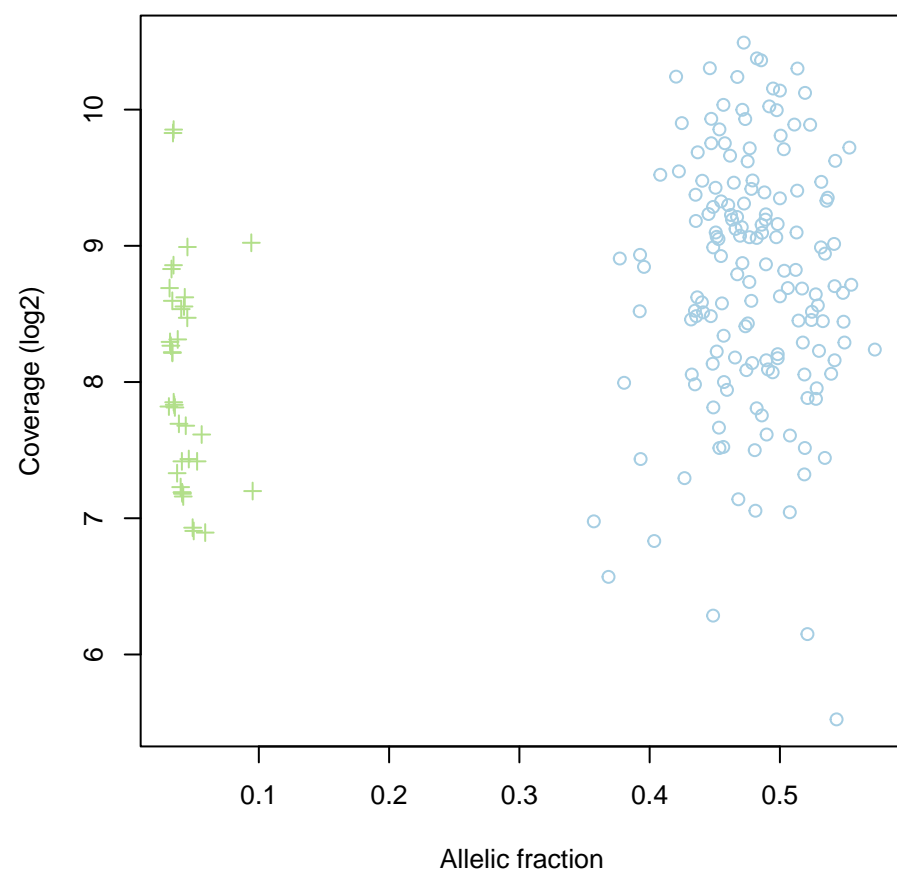
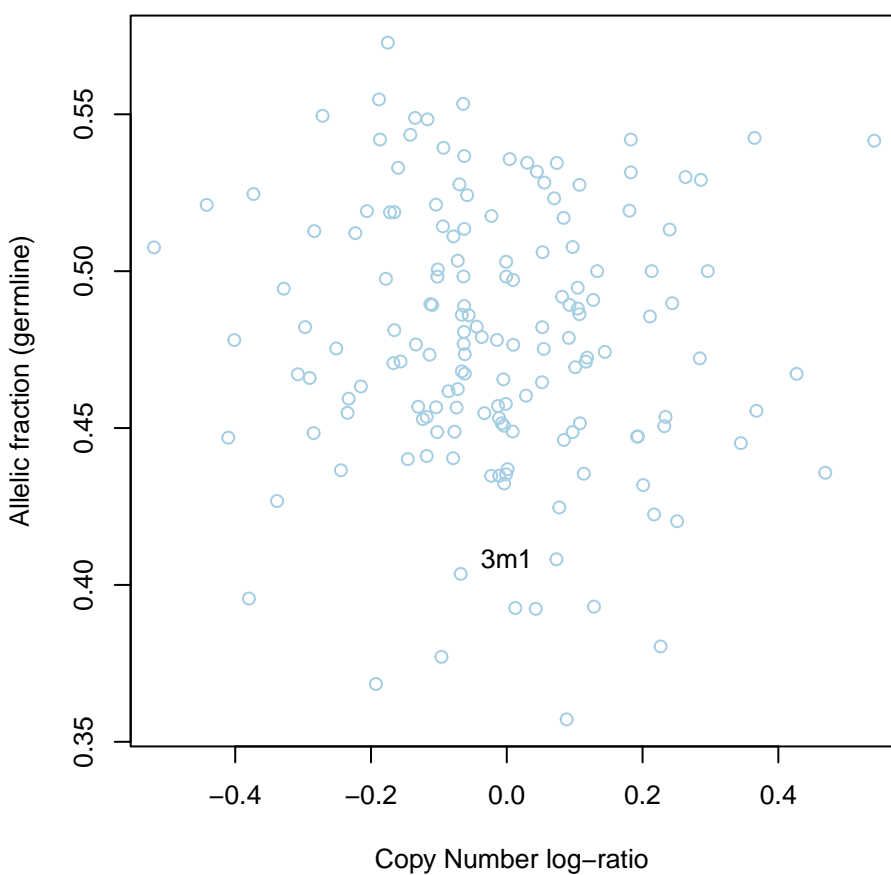
Purity: 0.45 Tumor ploidy: 3 SNV log-likelihood: -282.39 GoF: 82.7% Mean coverage: 410;477



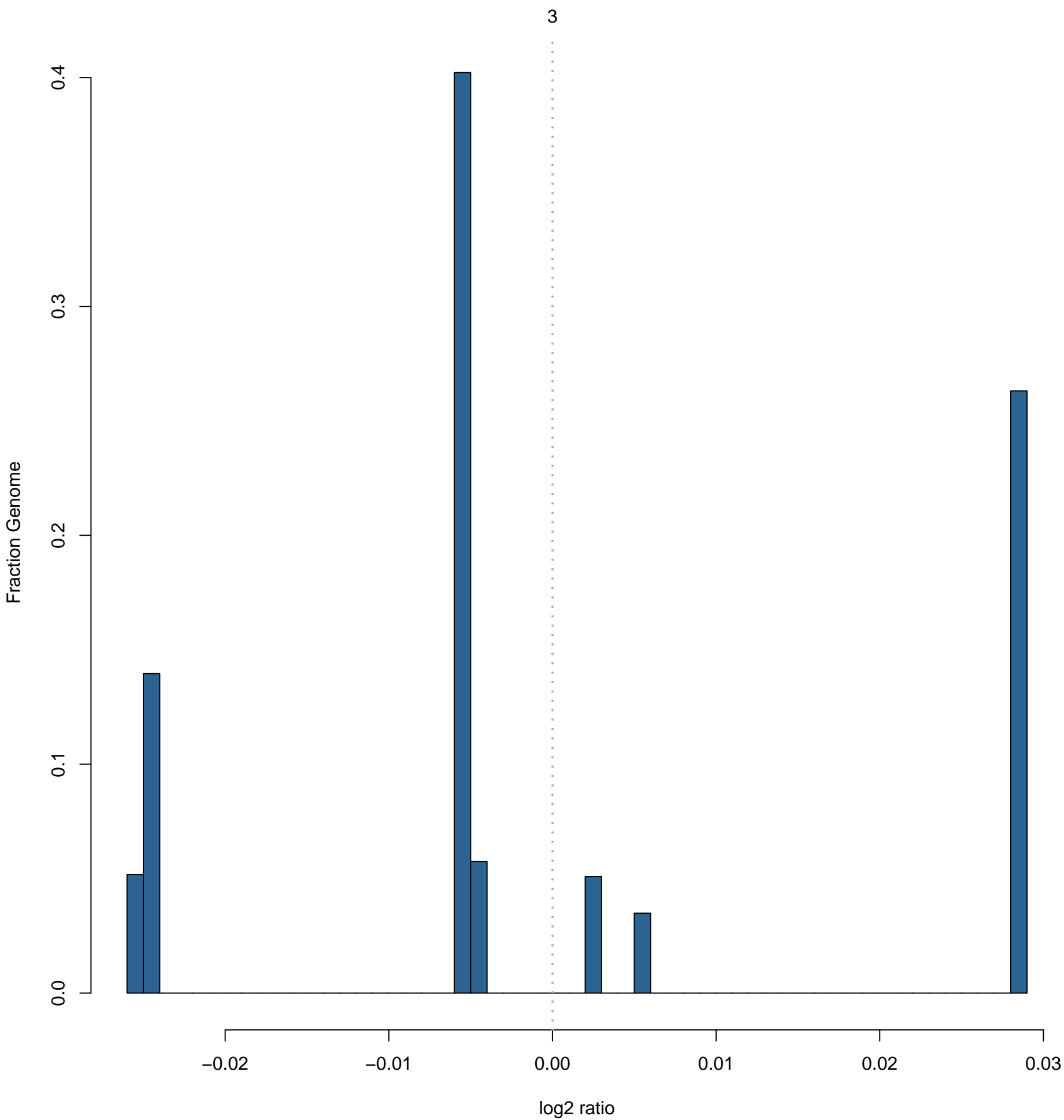
SCNA-fit log-likelihood: -7178.85



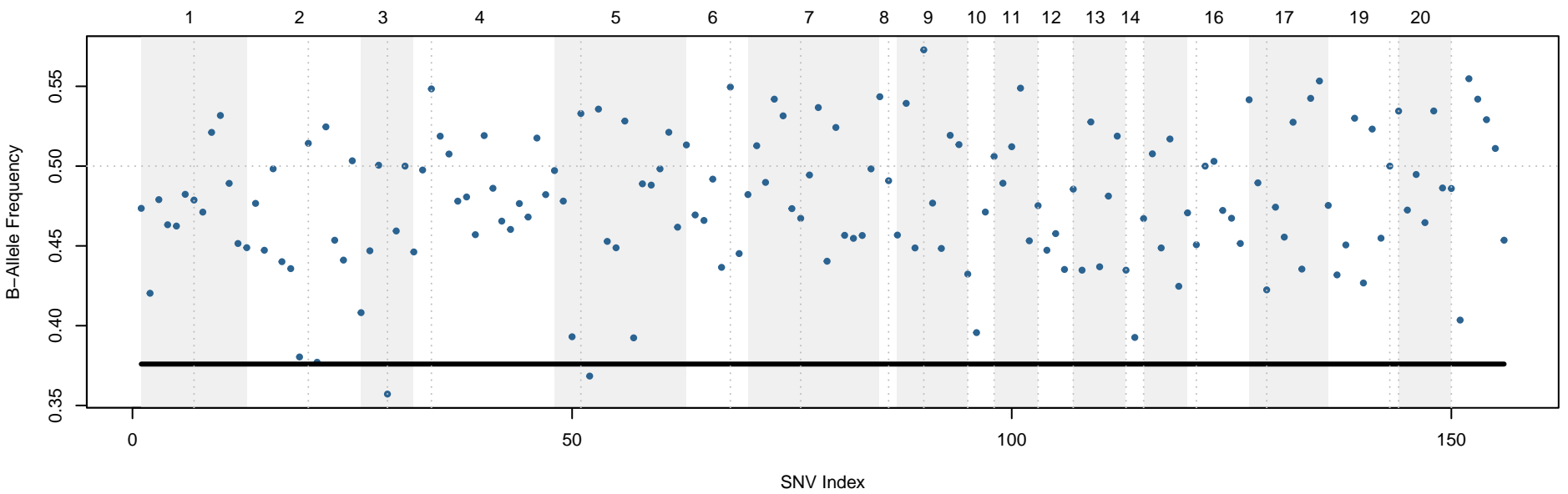




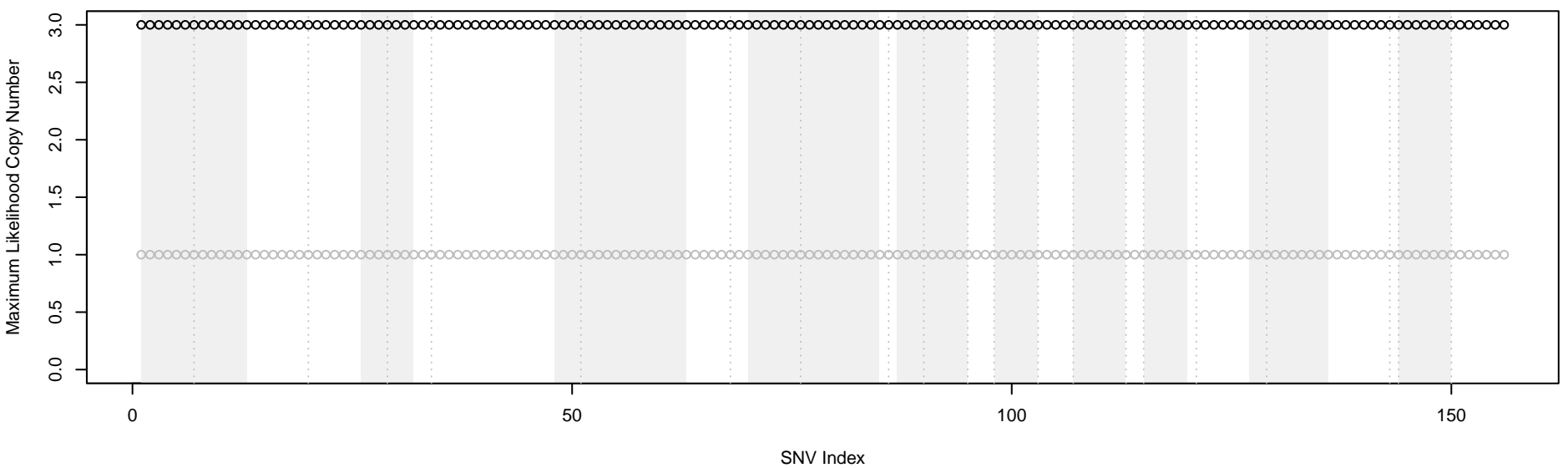
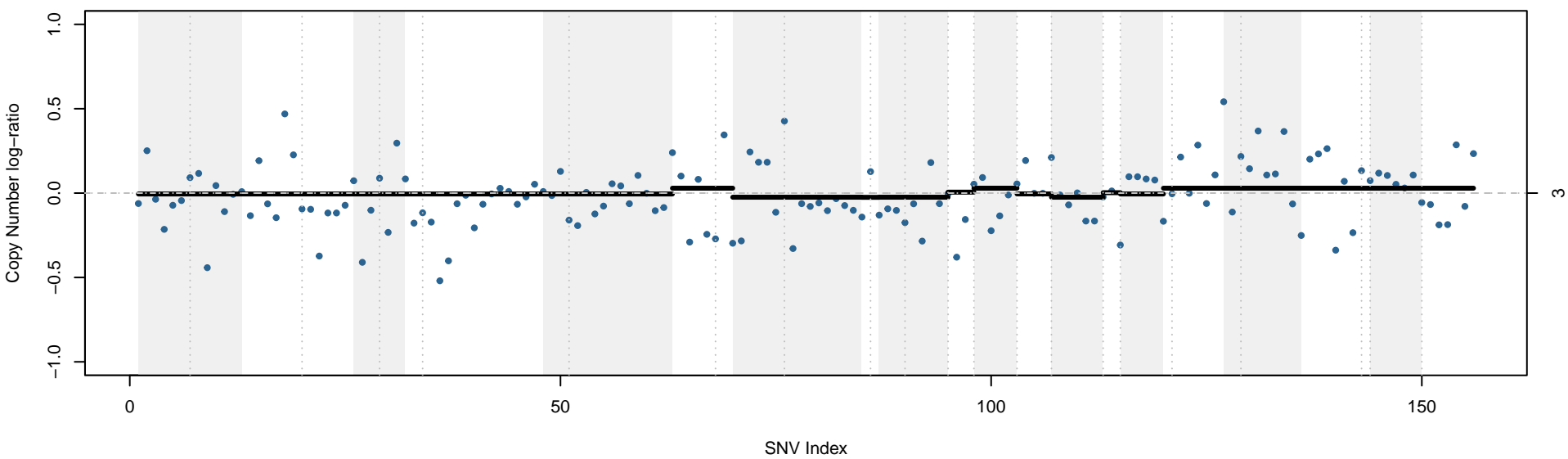
Purity: 0.66 Tumor ploidy: 3

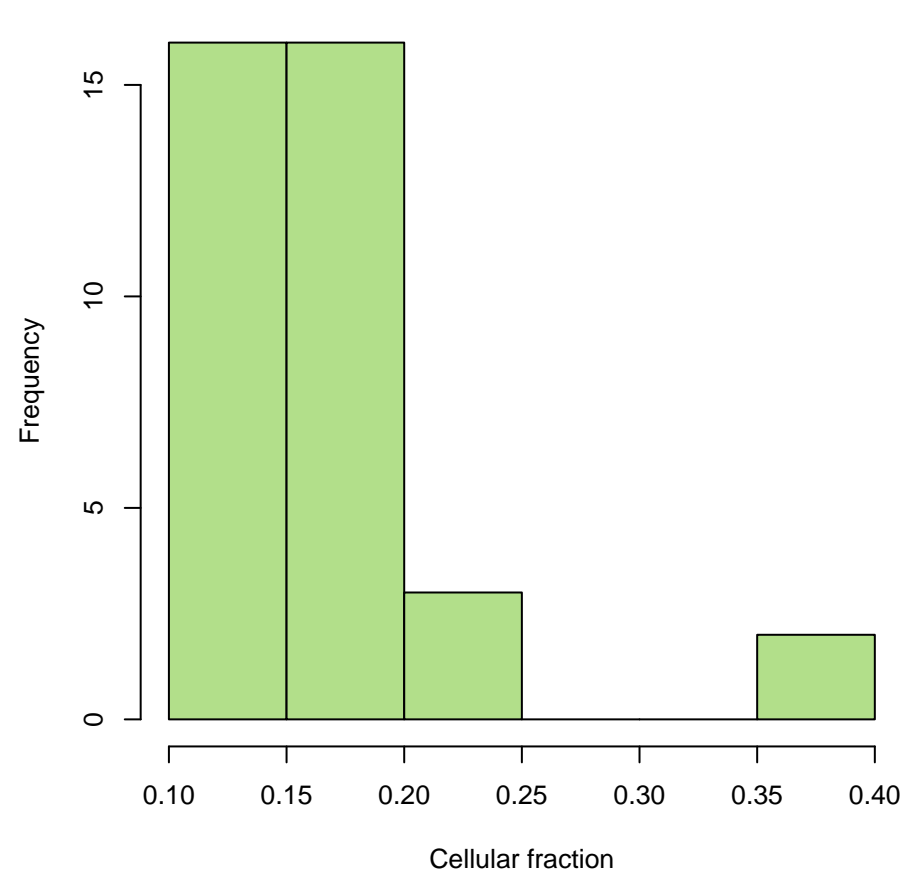
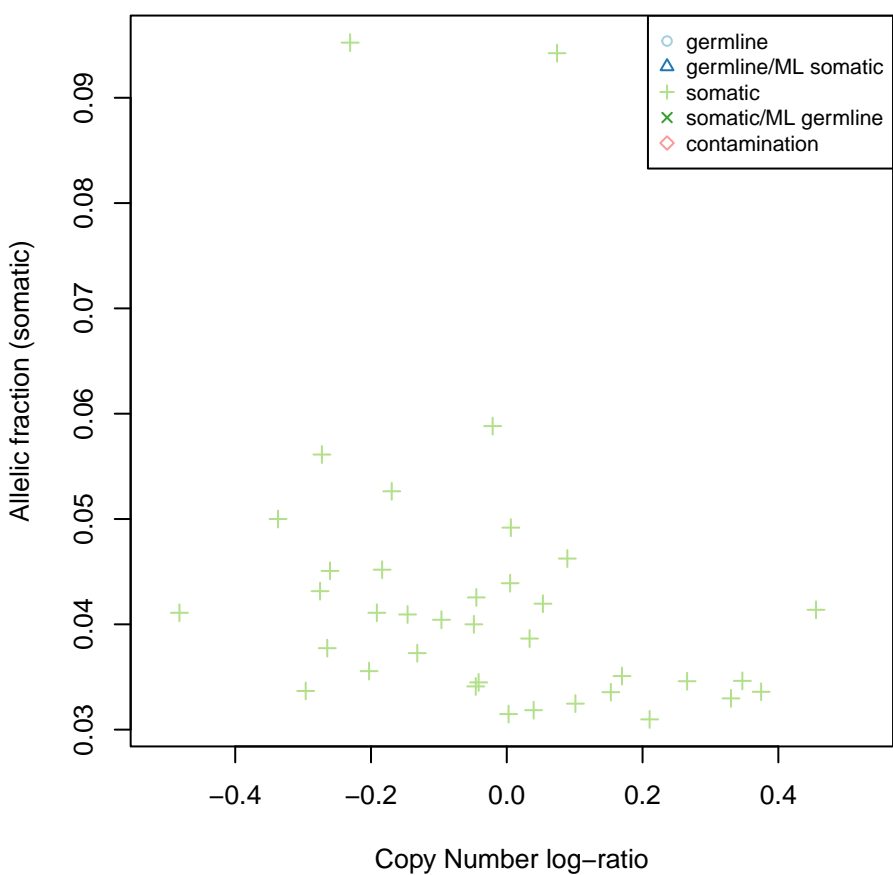
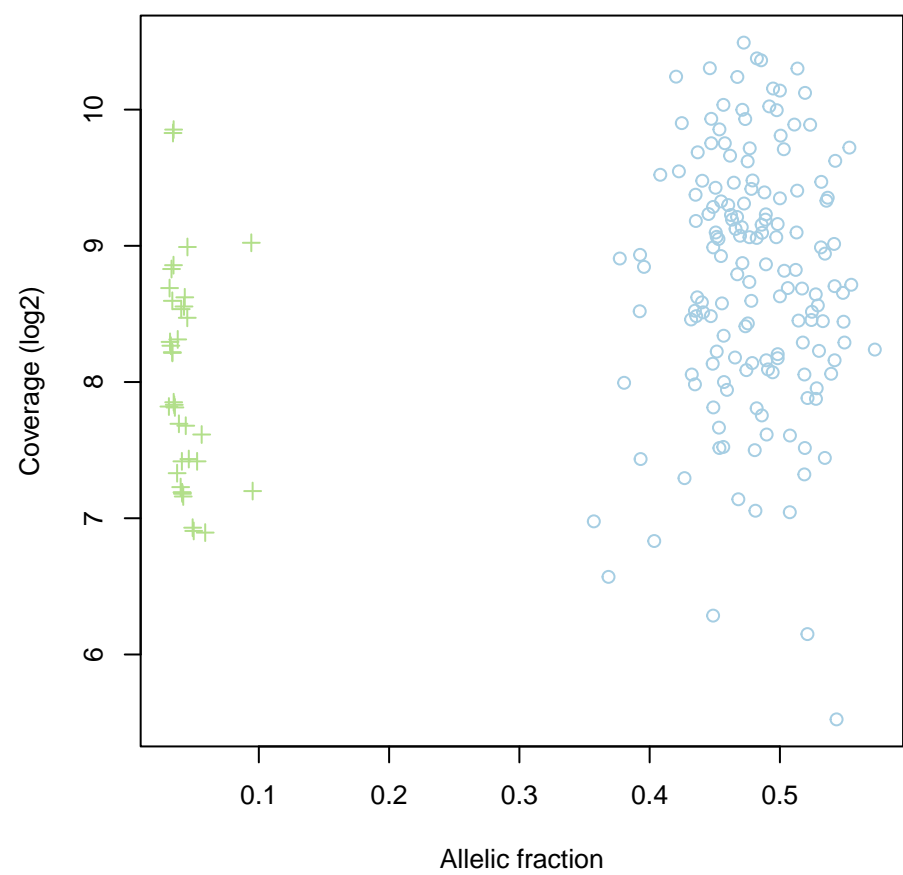
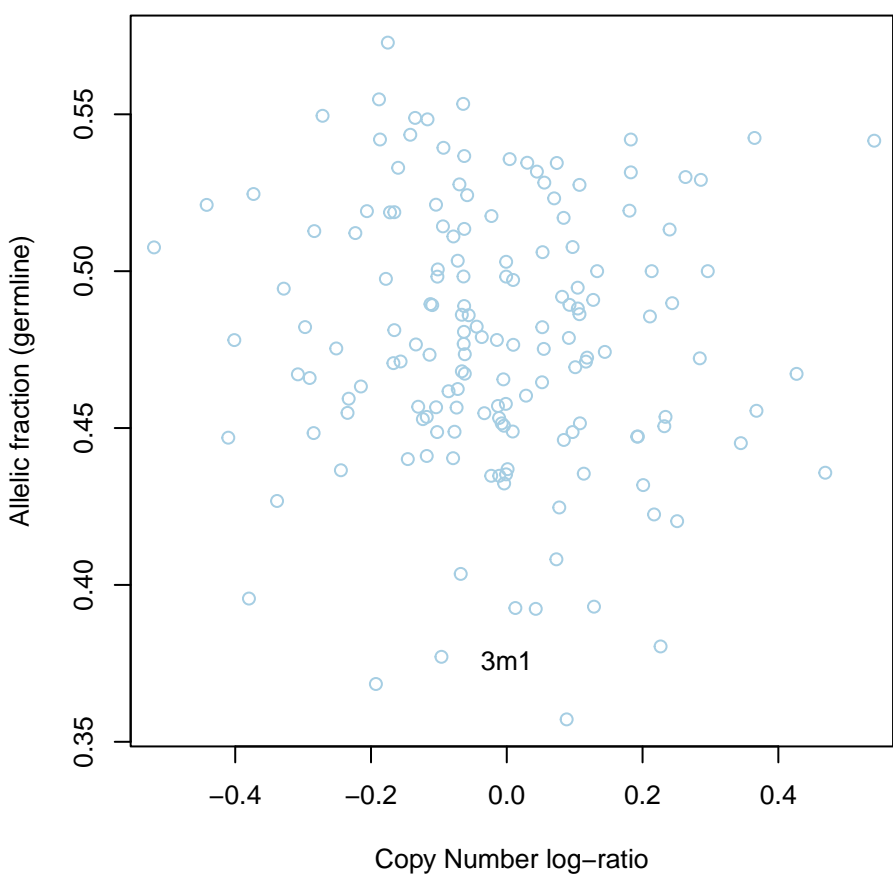


Purity: 0.66 Tumor ploidy: 3 SNV log-likelihood: -773.82 GoF: 62.7% Mean coverage: 410;477

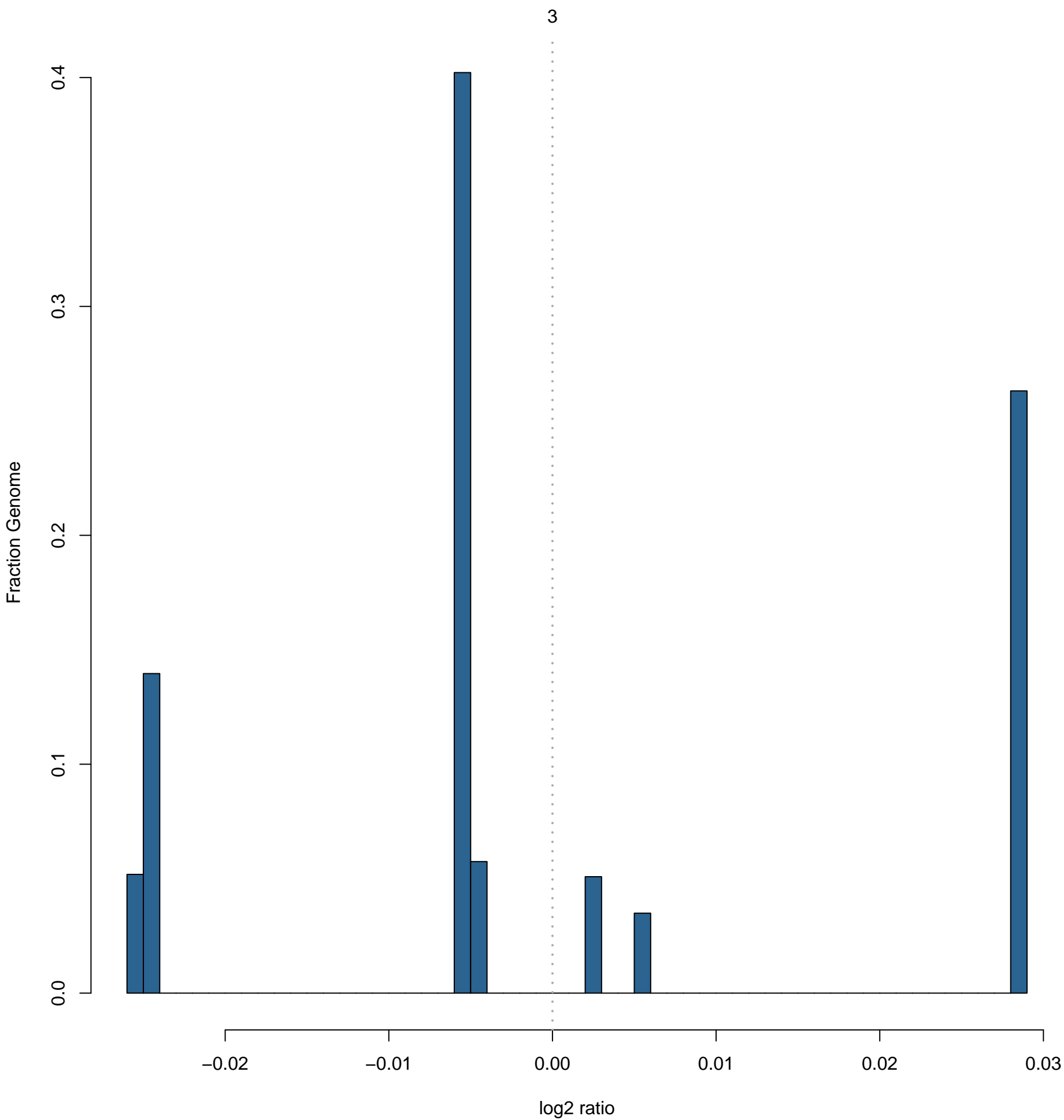


SCNA-fit log-likelihood: -7178.88

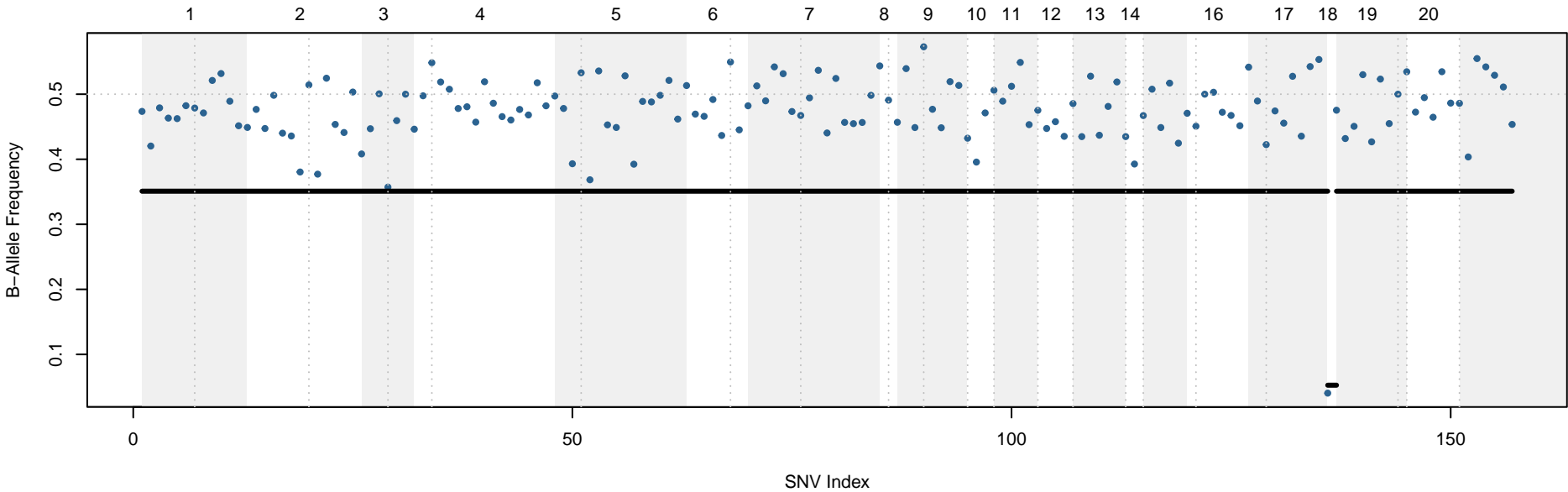




Purity: 0.85 Tumor ploidy: 3



Purity: 0.85 Tumor ploidy: 3 SNV log-likelihood: -1314.23 GoF: 42.3% Mean coverage: 410;477



SCNA-fit log-likelihood: -7178.92

