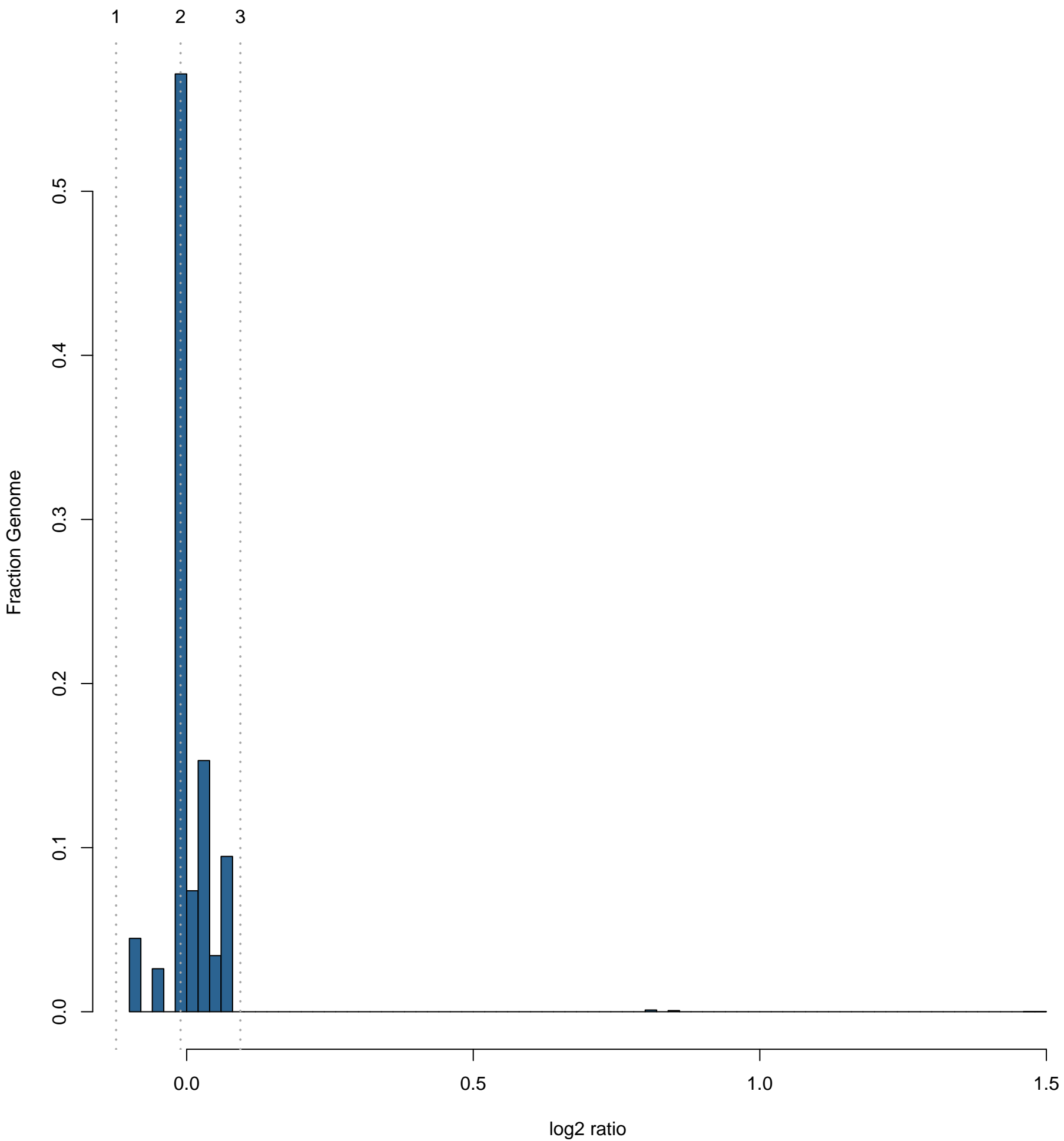
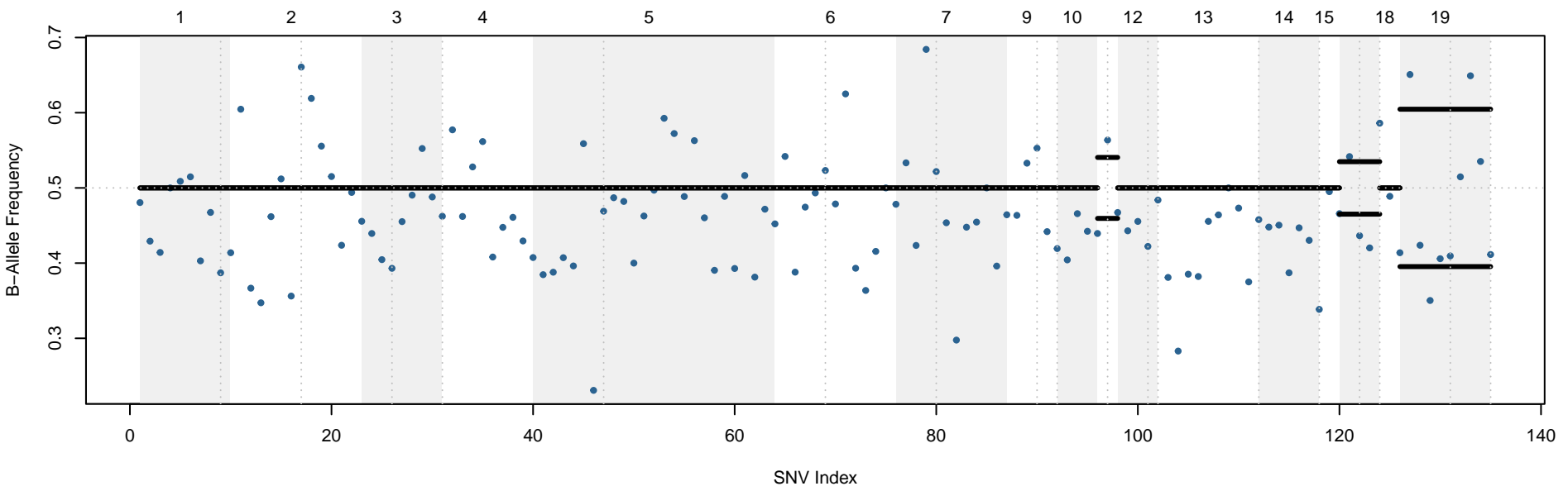


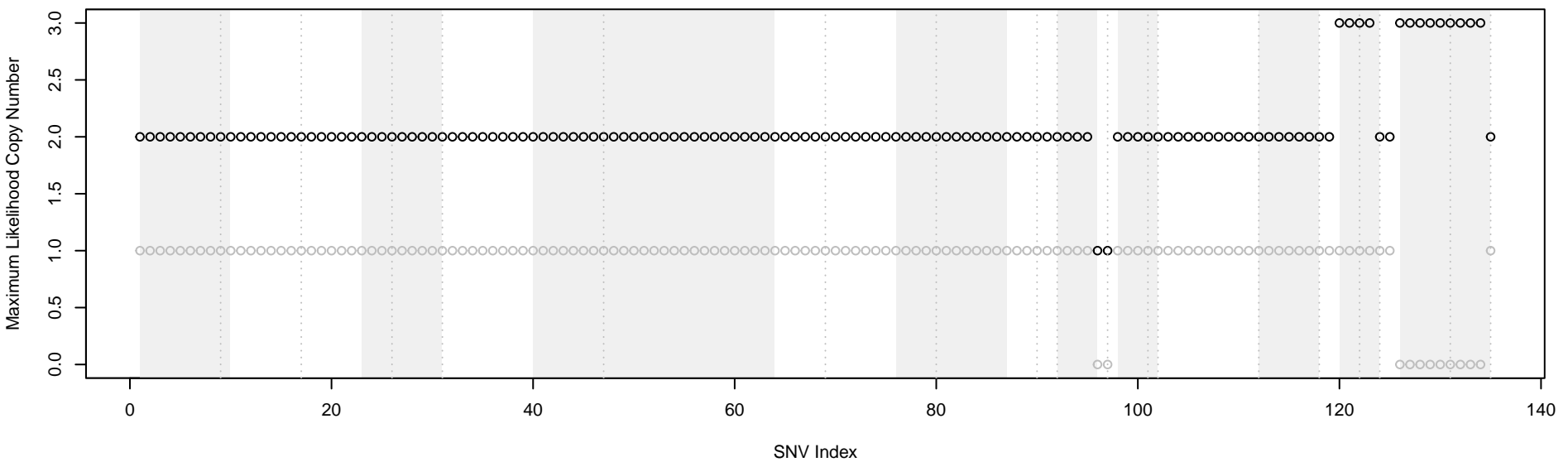
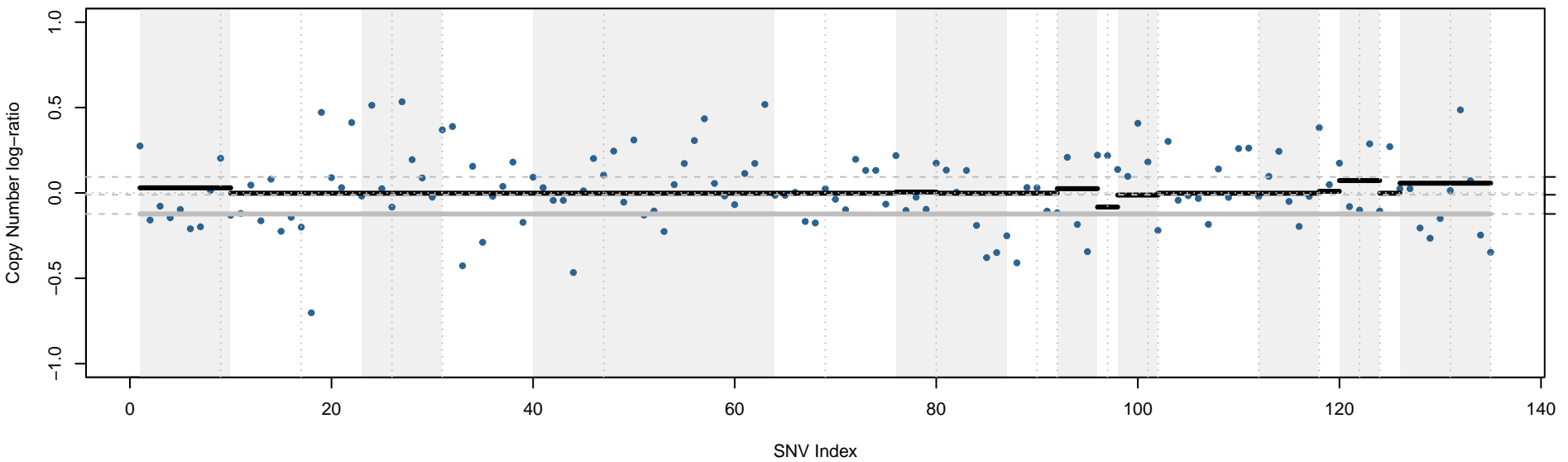
Purity: 0.15 Tumor ploidy: 2.099

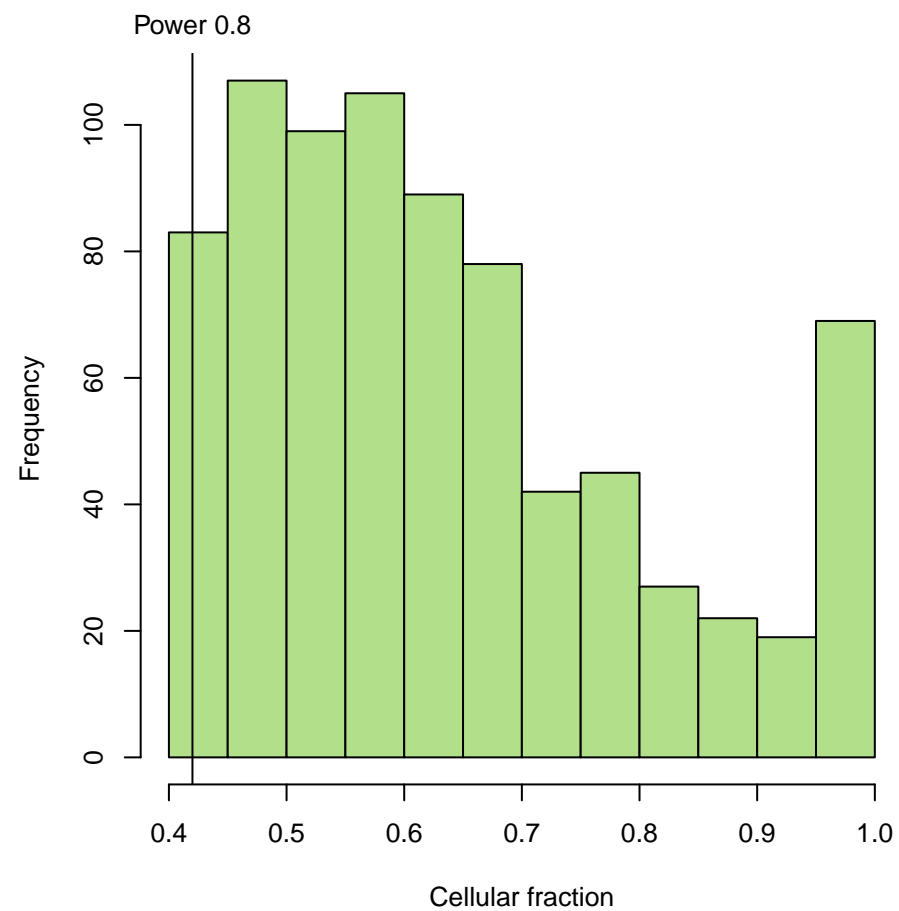
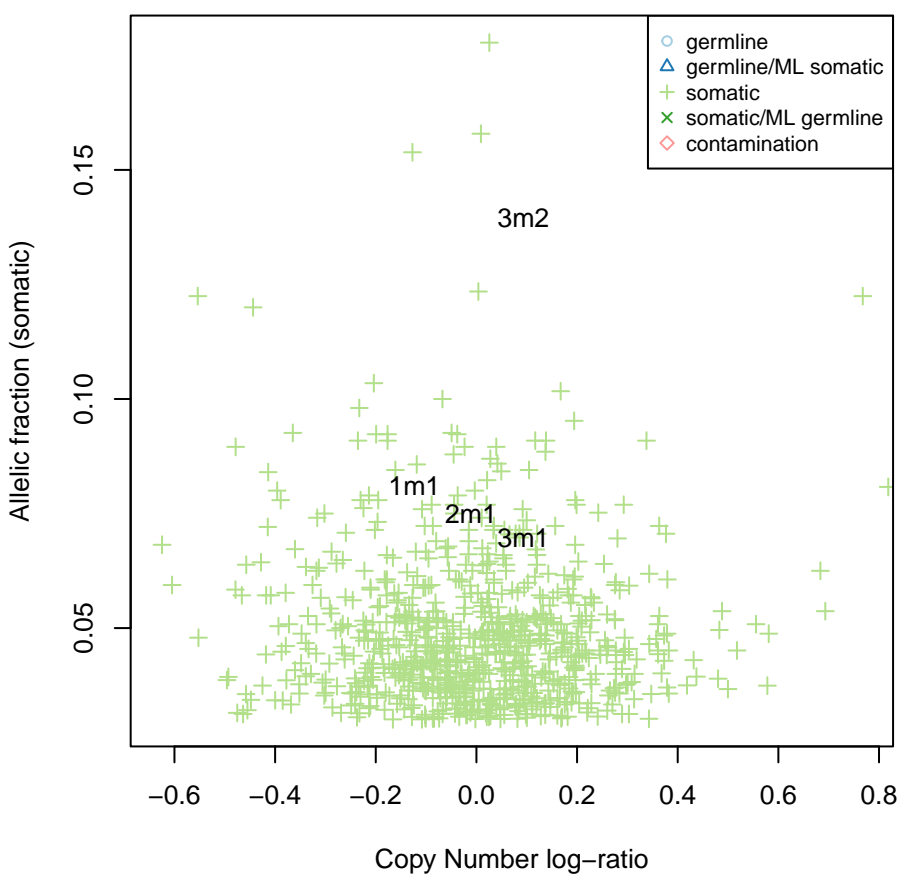
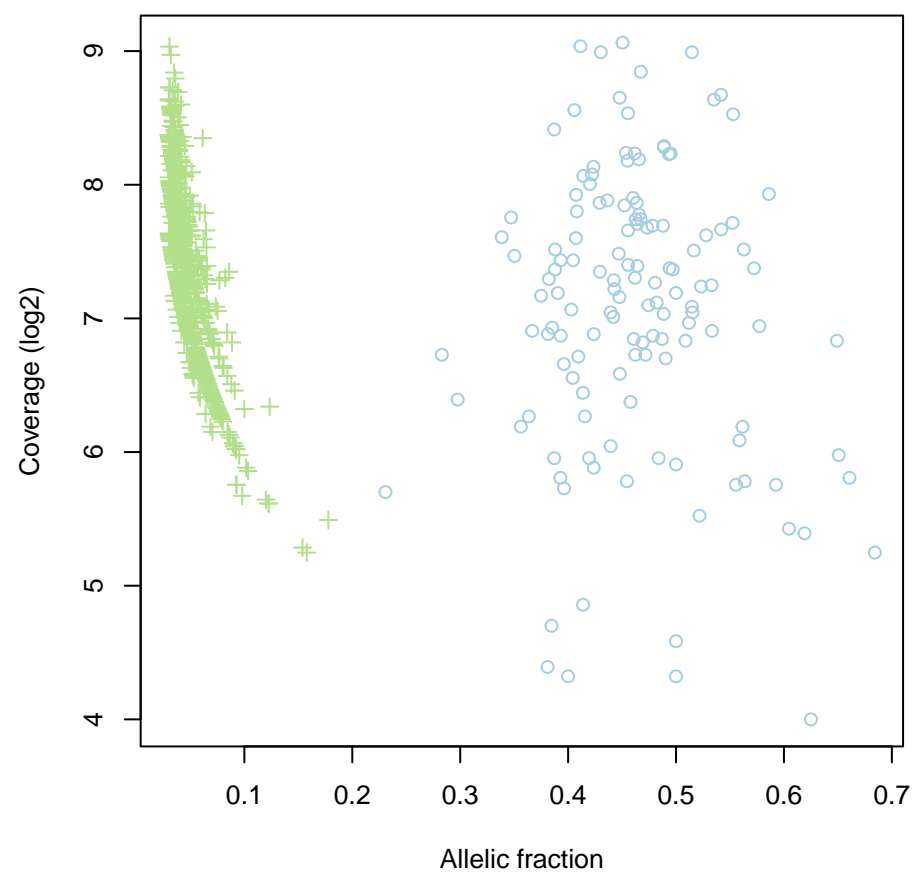
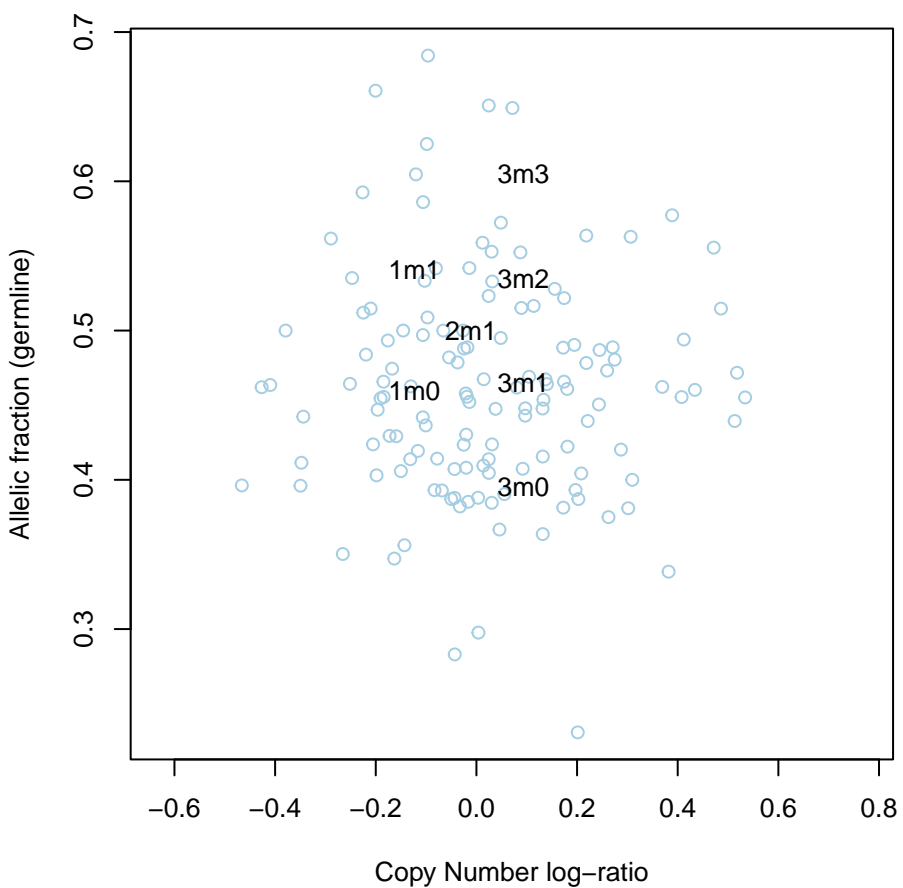


Purity: 0.15 Tumor ploidy: 2.099 SNV log-likelihood: 521.08 GoF: 96.1% Mean coverage: 176;149

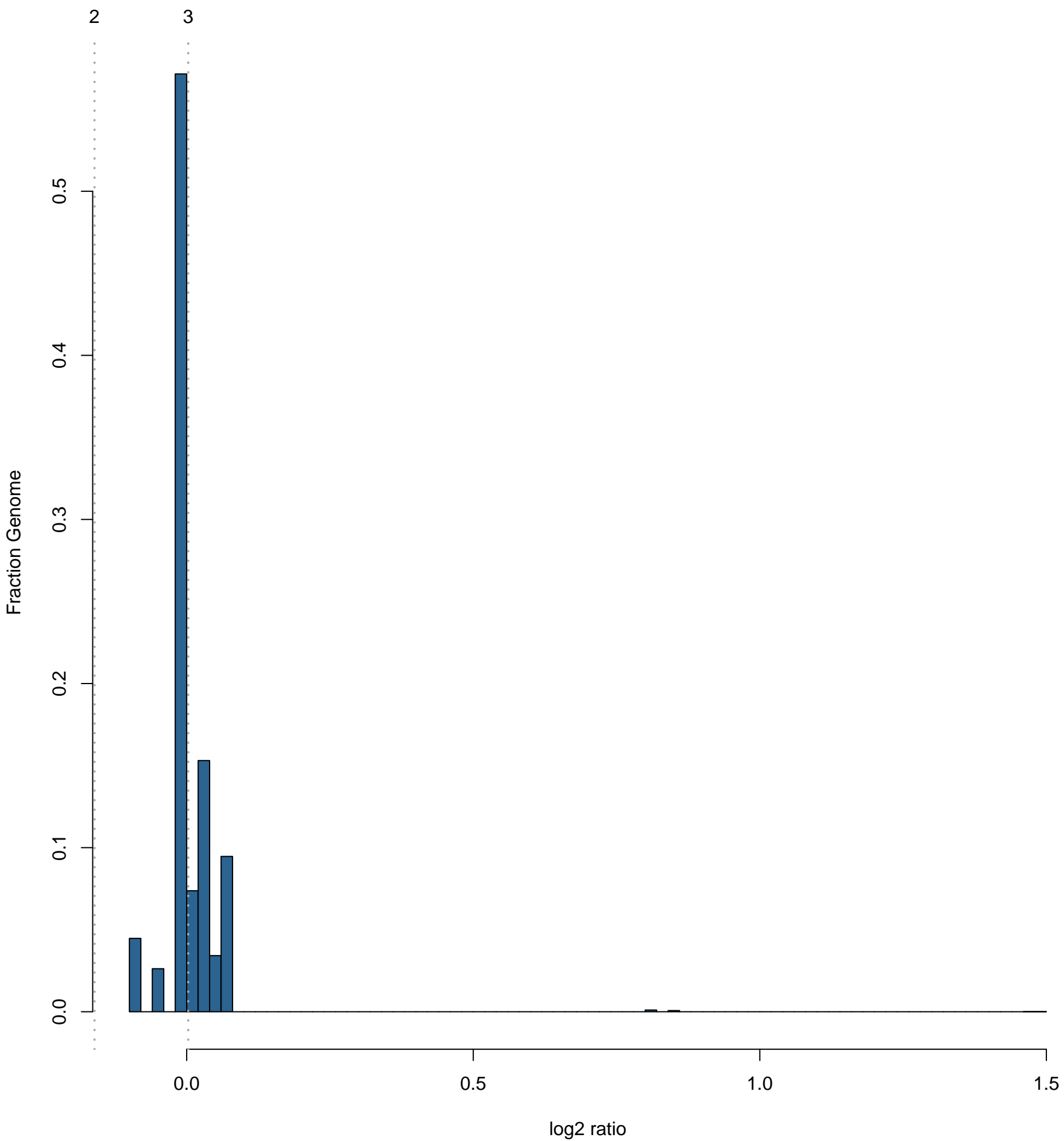


SCNA-fit log-likelihood: -14131.4

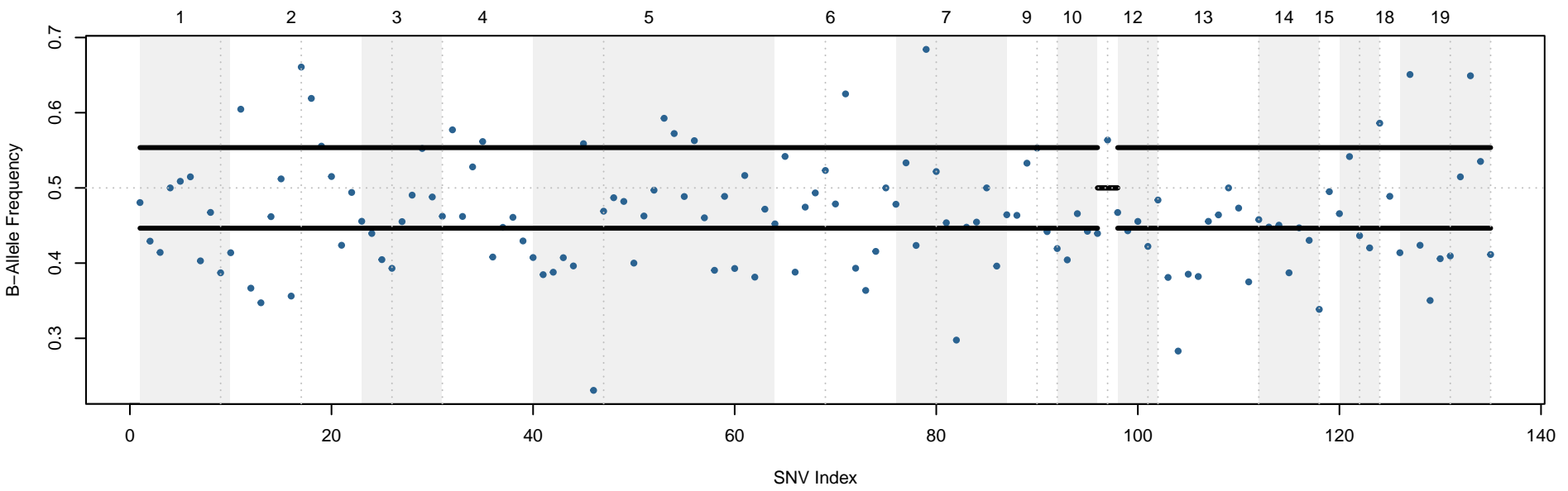




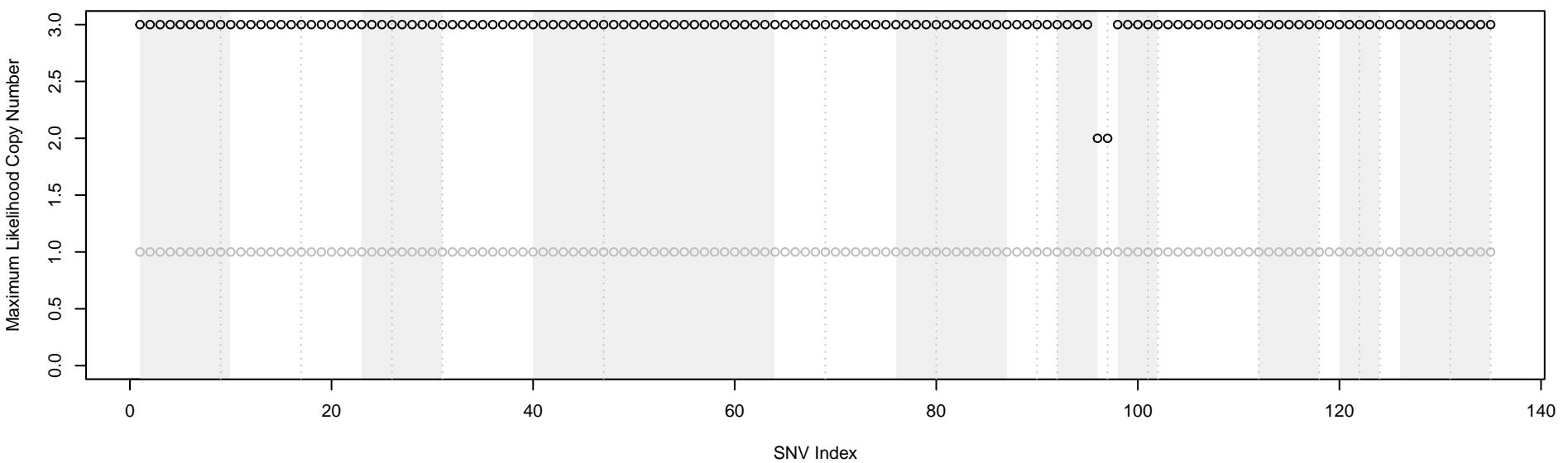
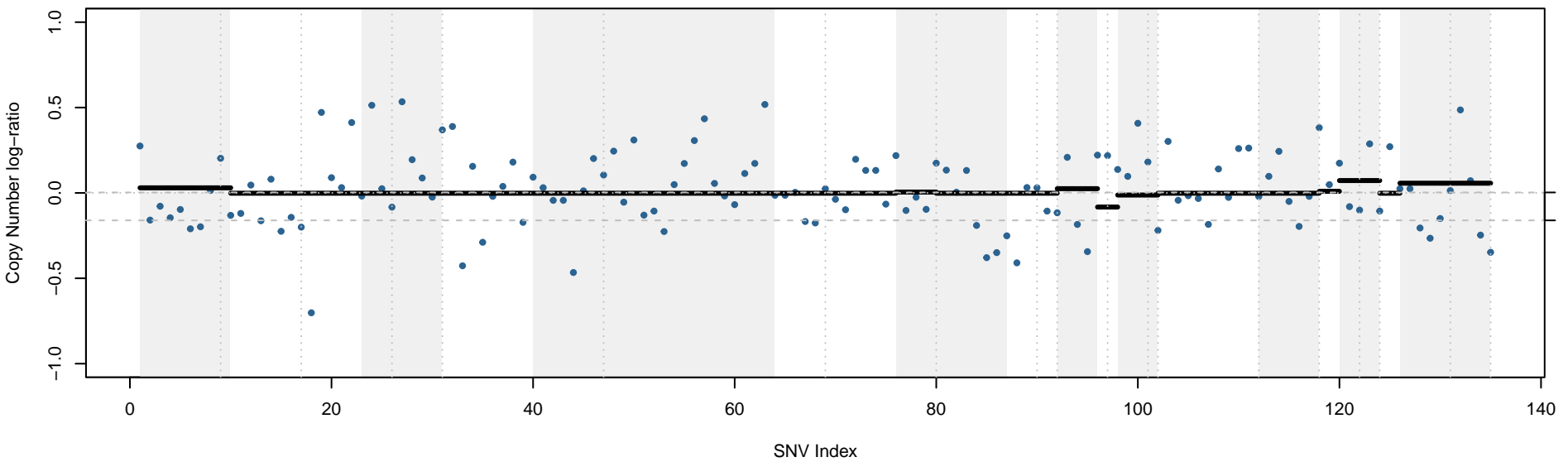
Purity: 0.24 Tumor ploidy: 2.983

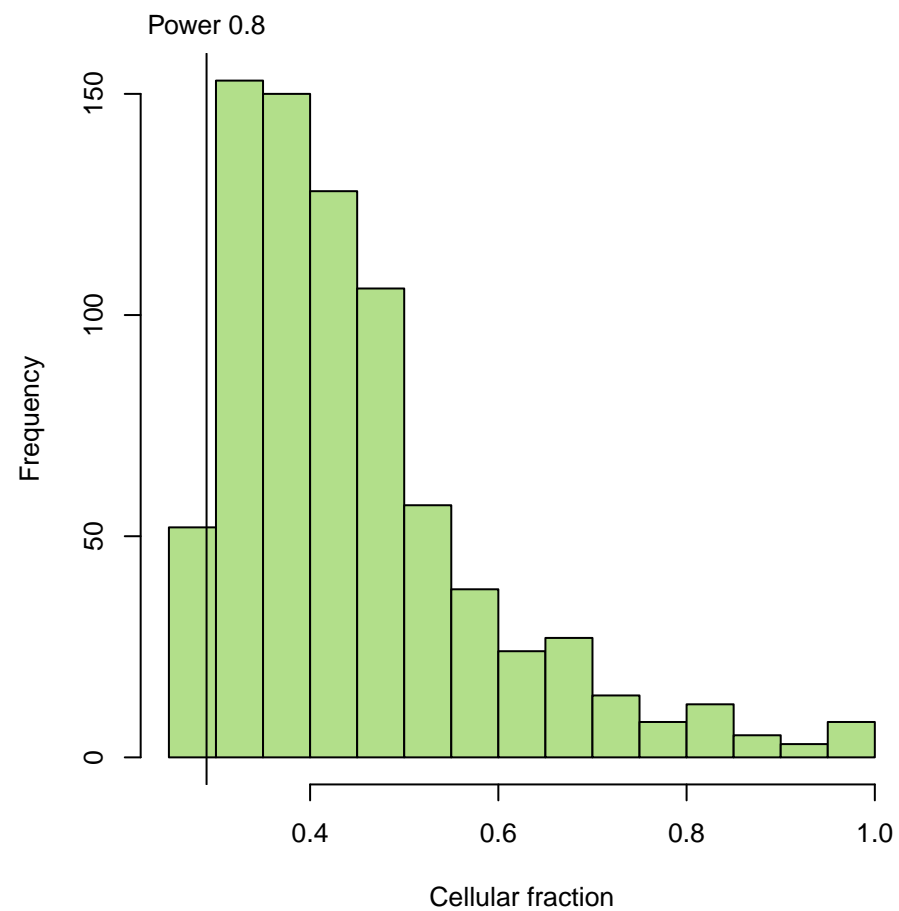
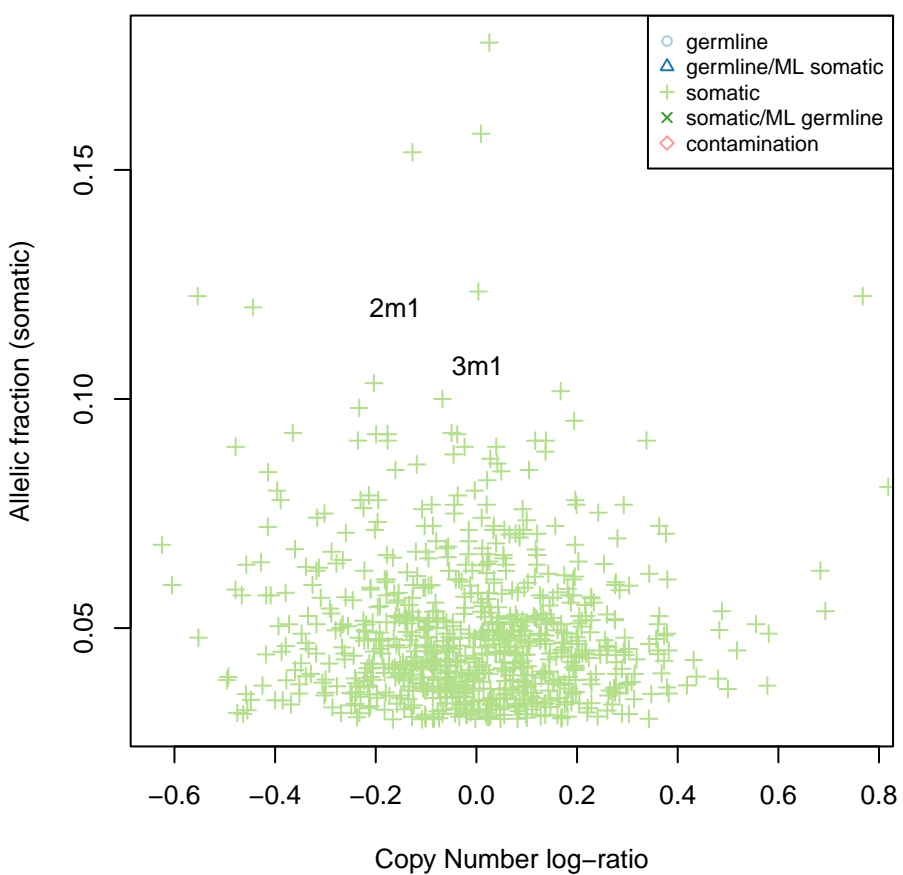
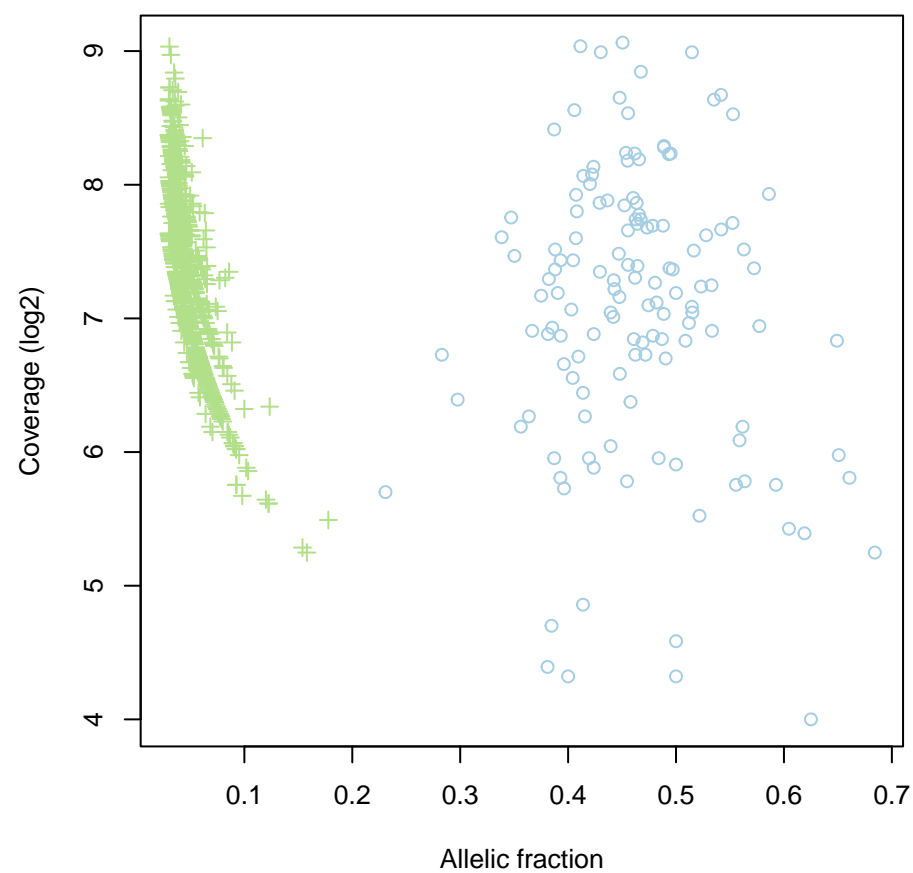
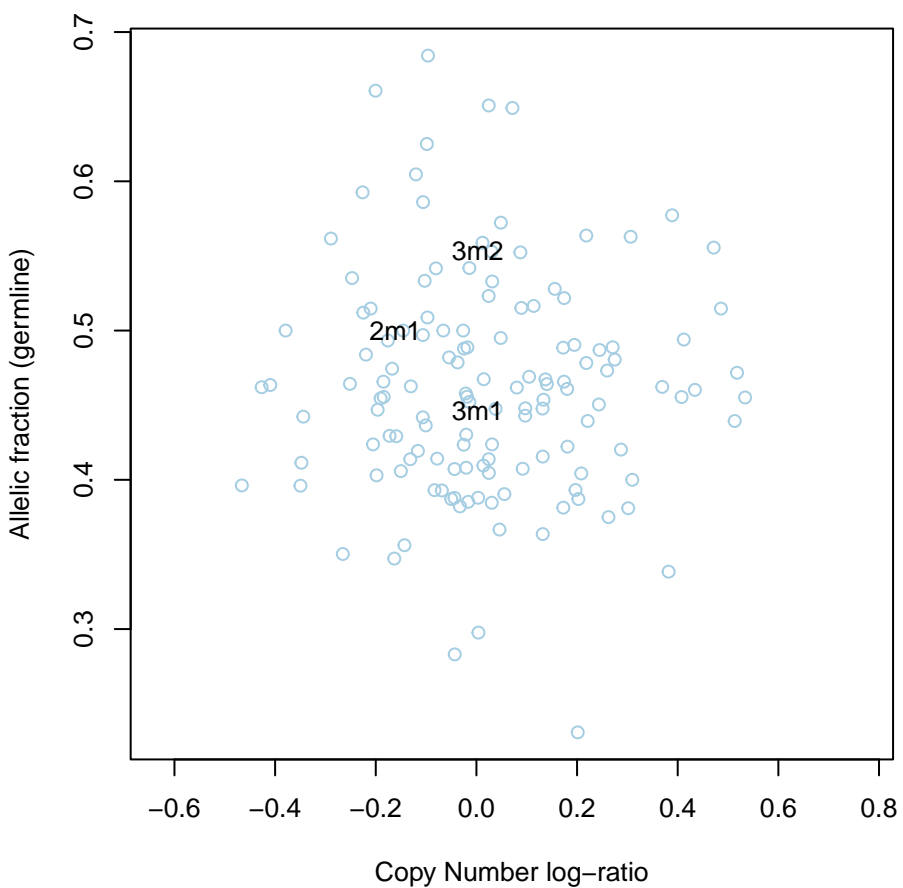


Purity: 0.24 Tumor ploidy: 2.983 SNV log-likelihood: 242.21 GoF: 91.5% Mean coverage: 176;149

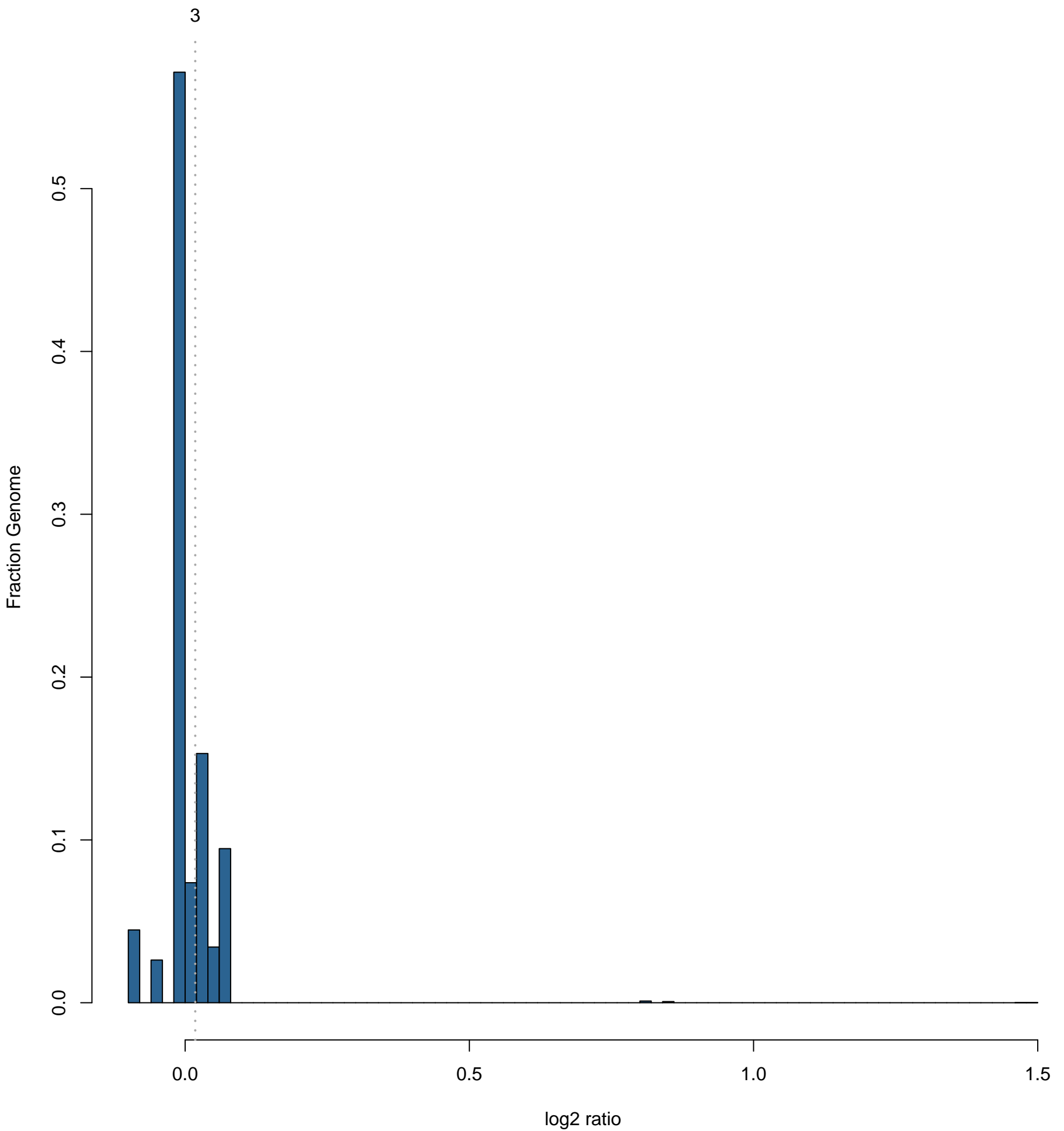


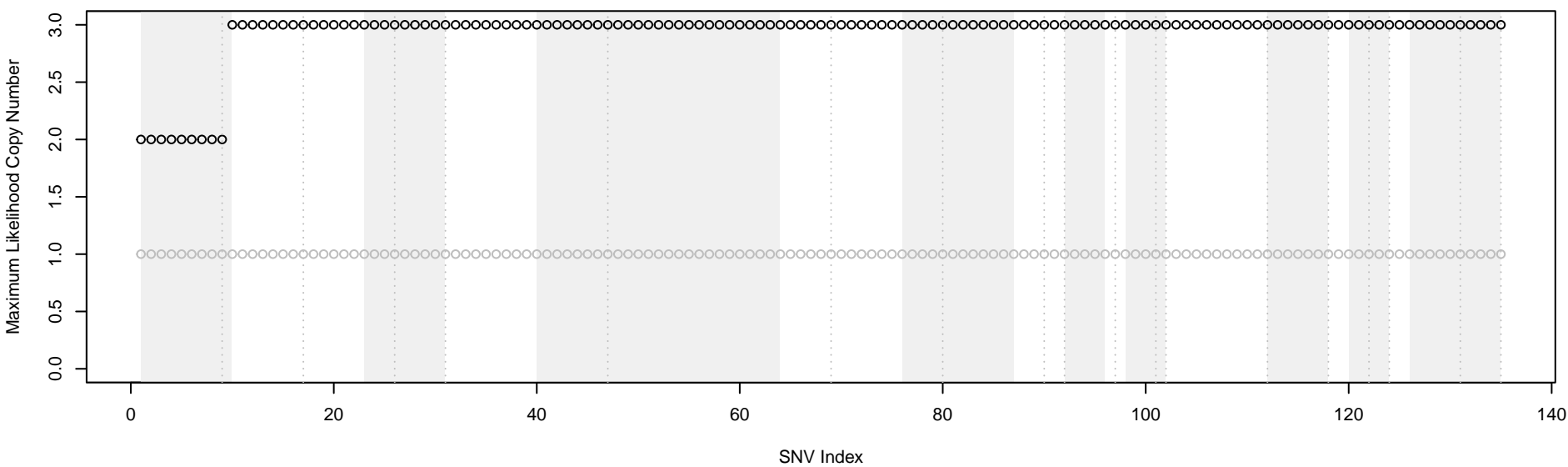
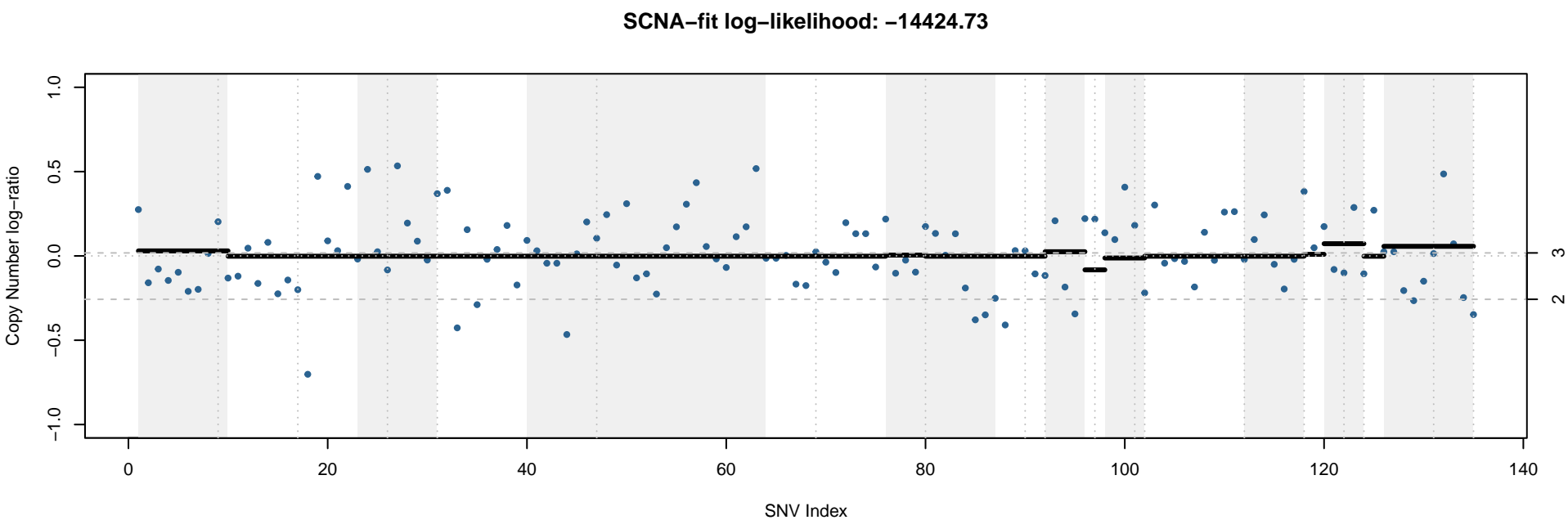
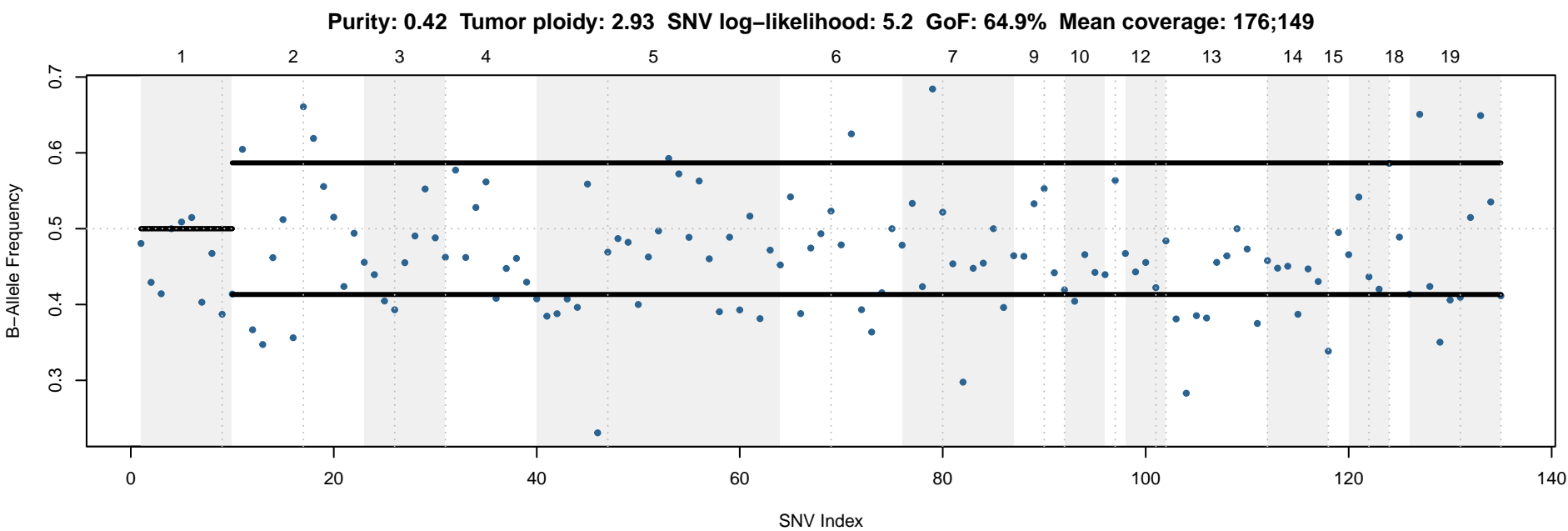
SCNA-fit log-likelihood: -14148.23



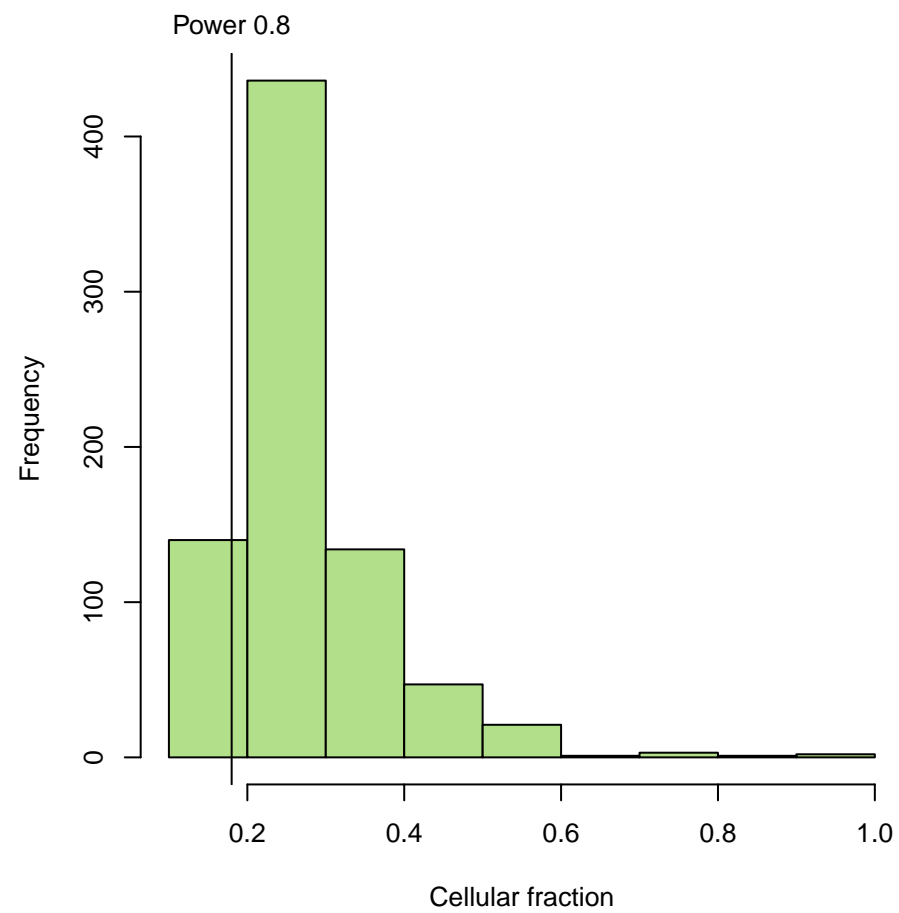
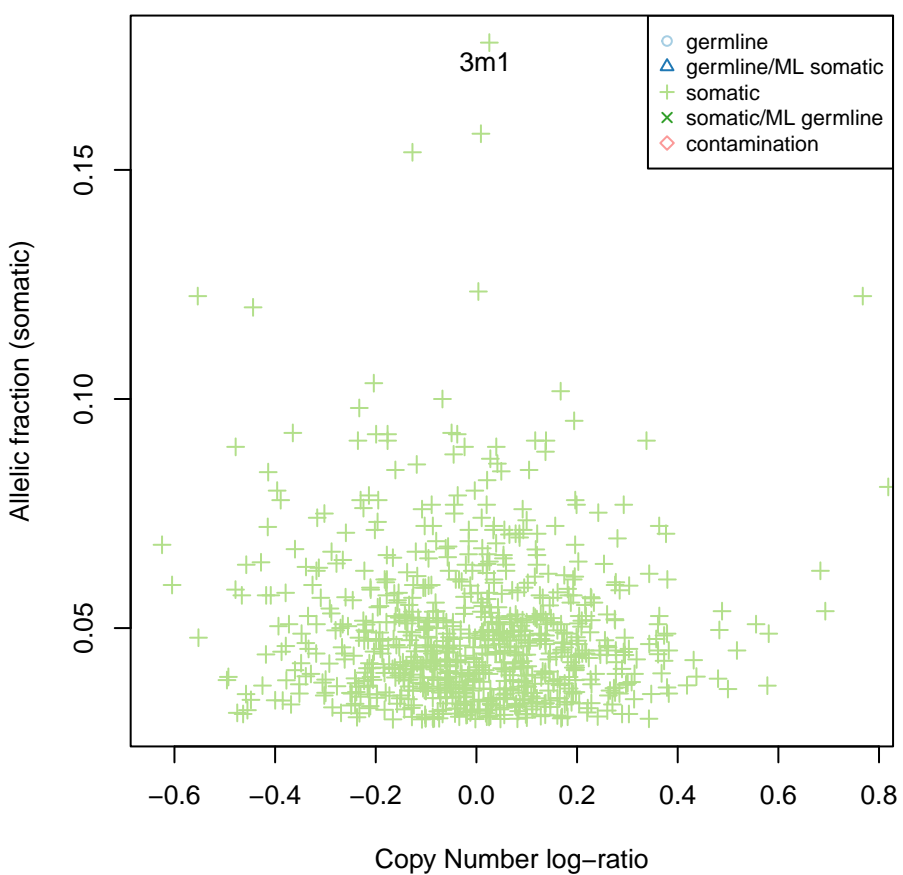
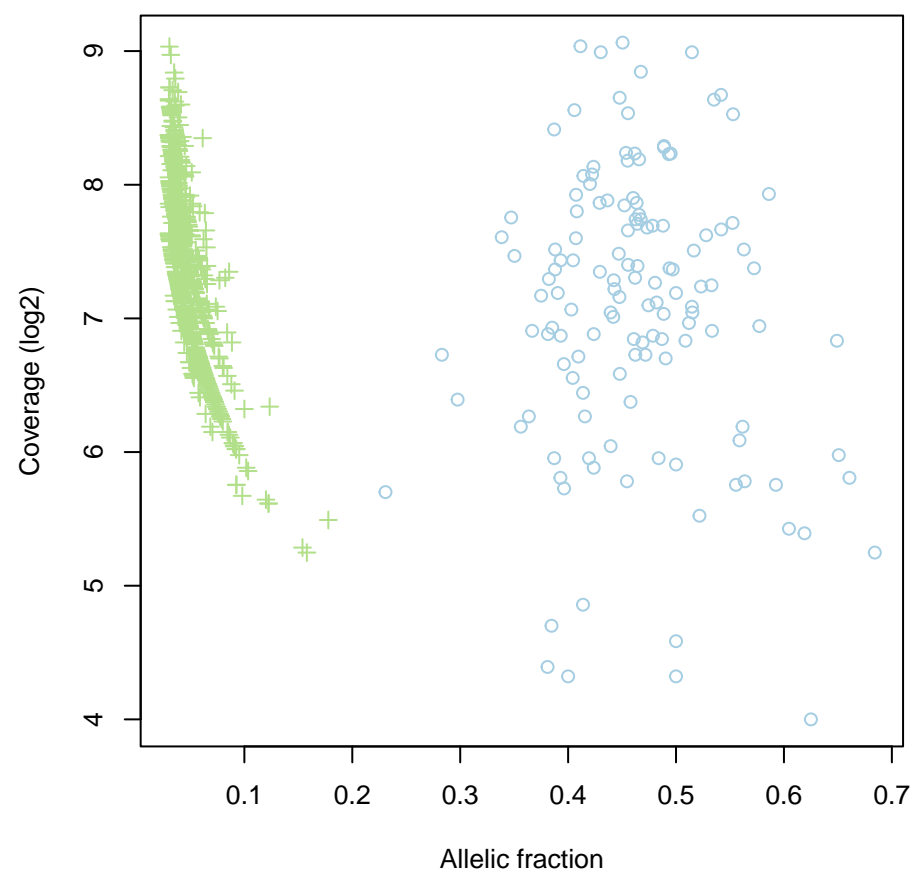
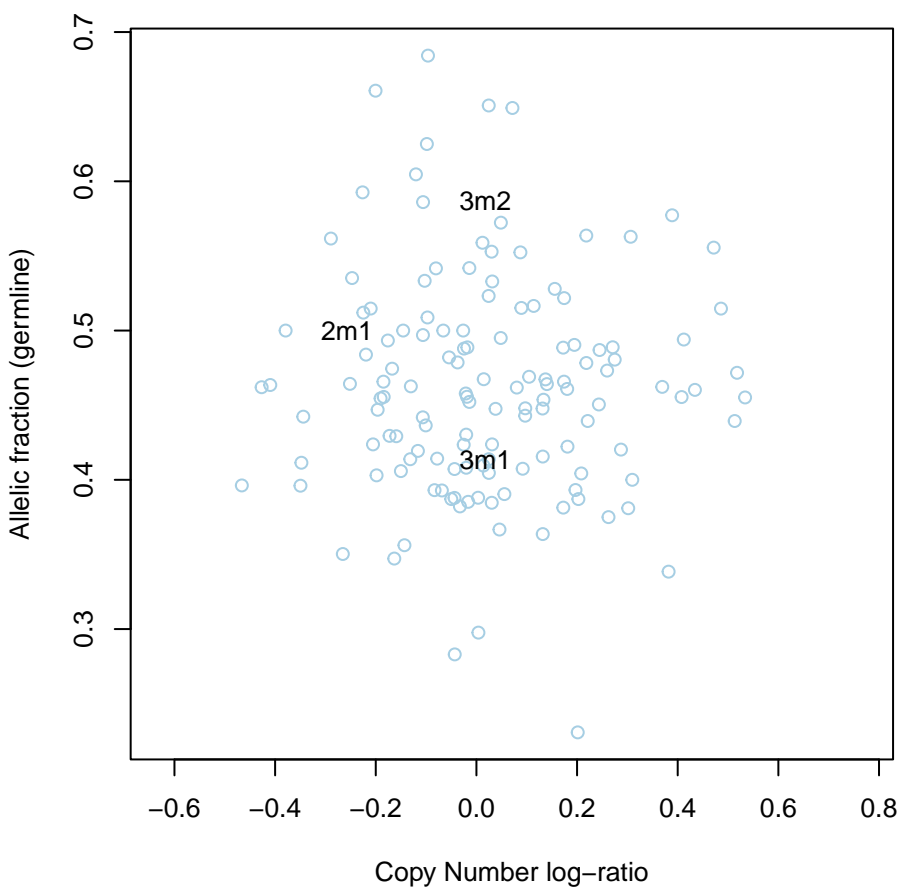


Purity: 0.42 Tumor ploidy: 2.93

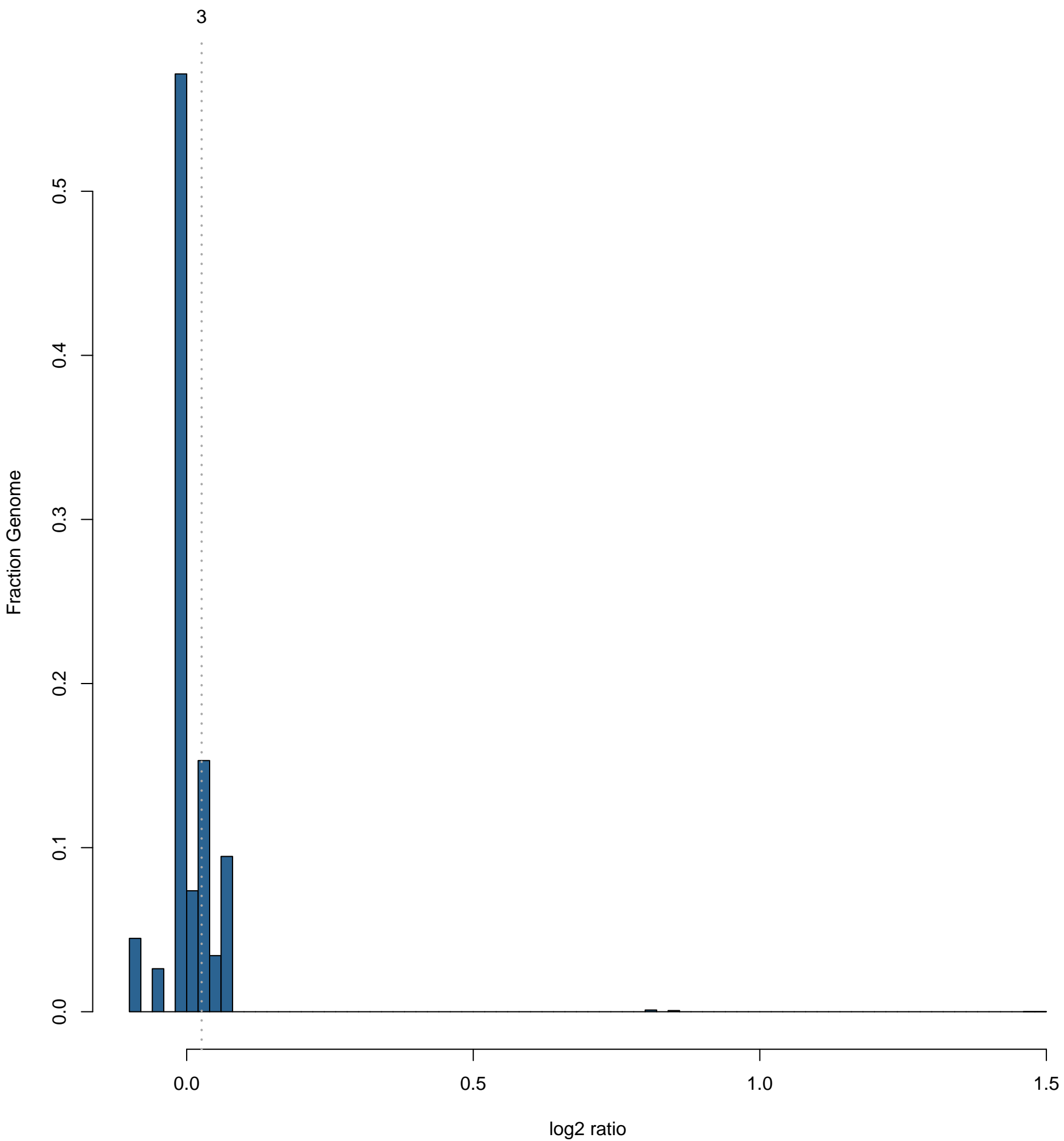




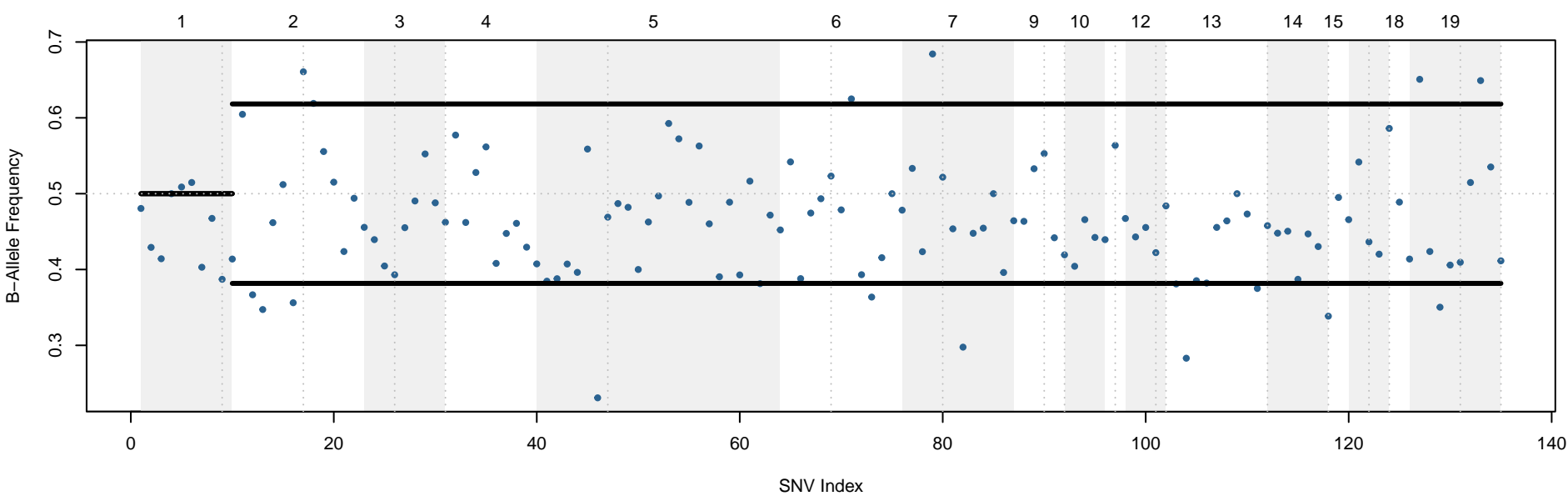




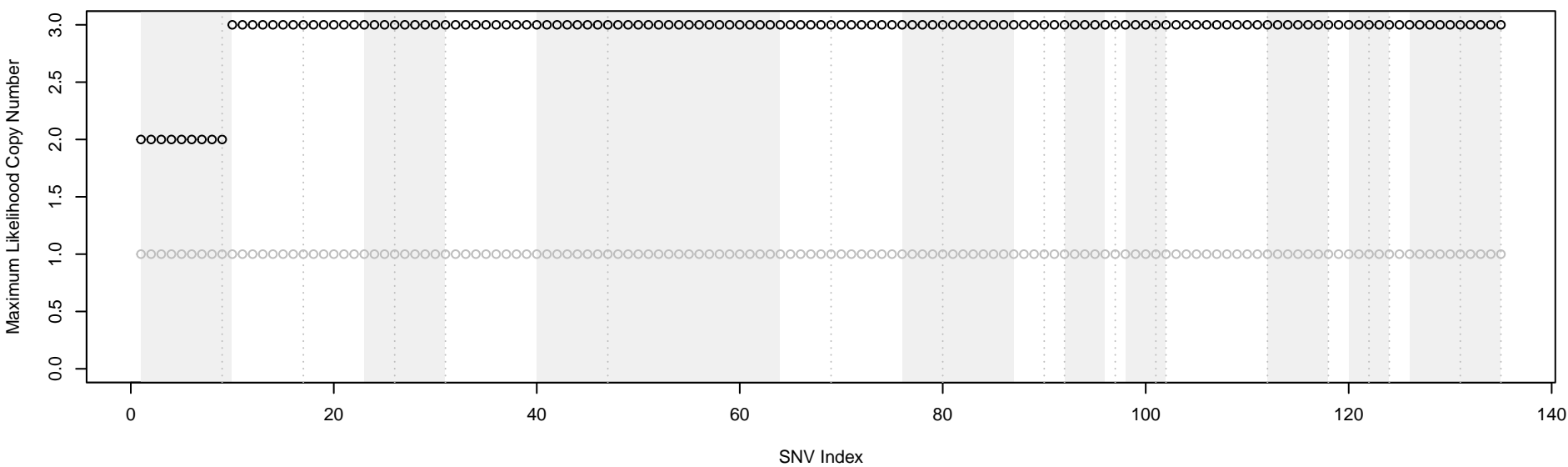
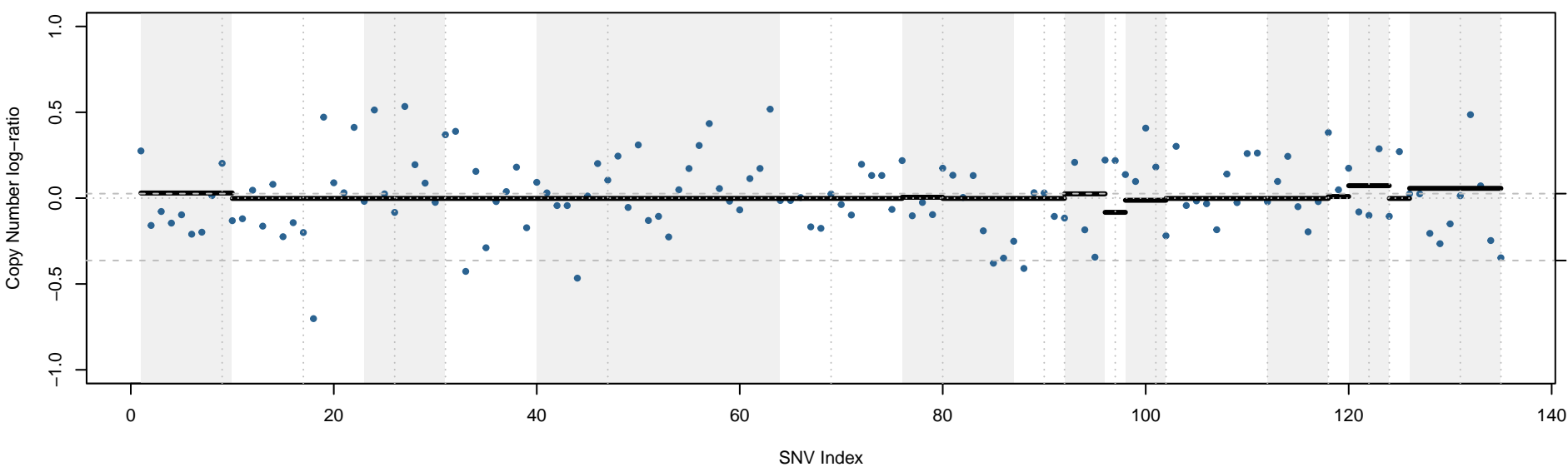
Purity: 0.62 Tumor ploidy: 2.924

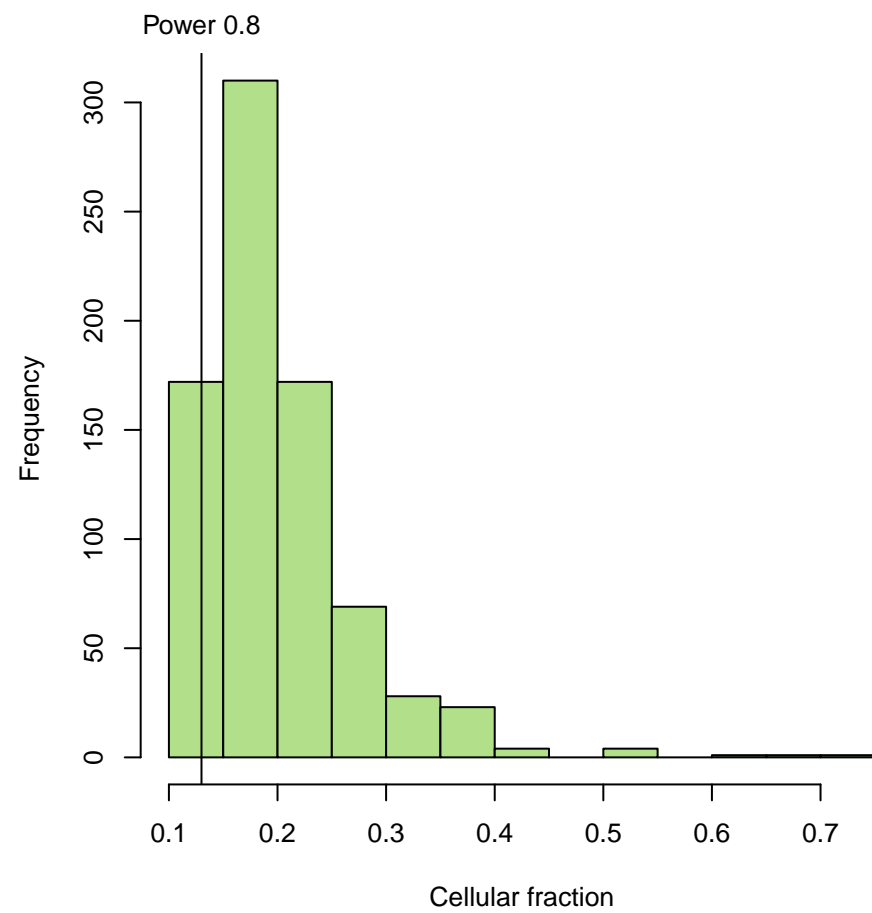
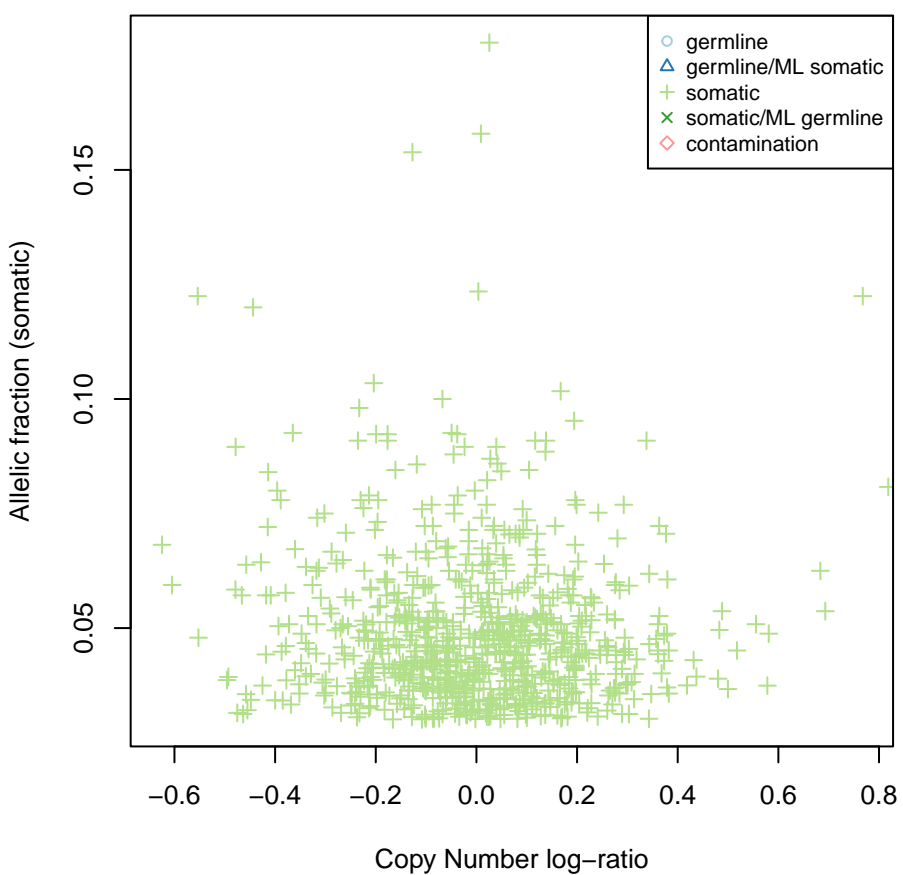
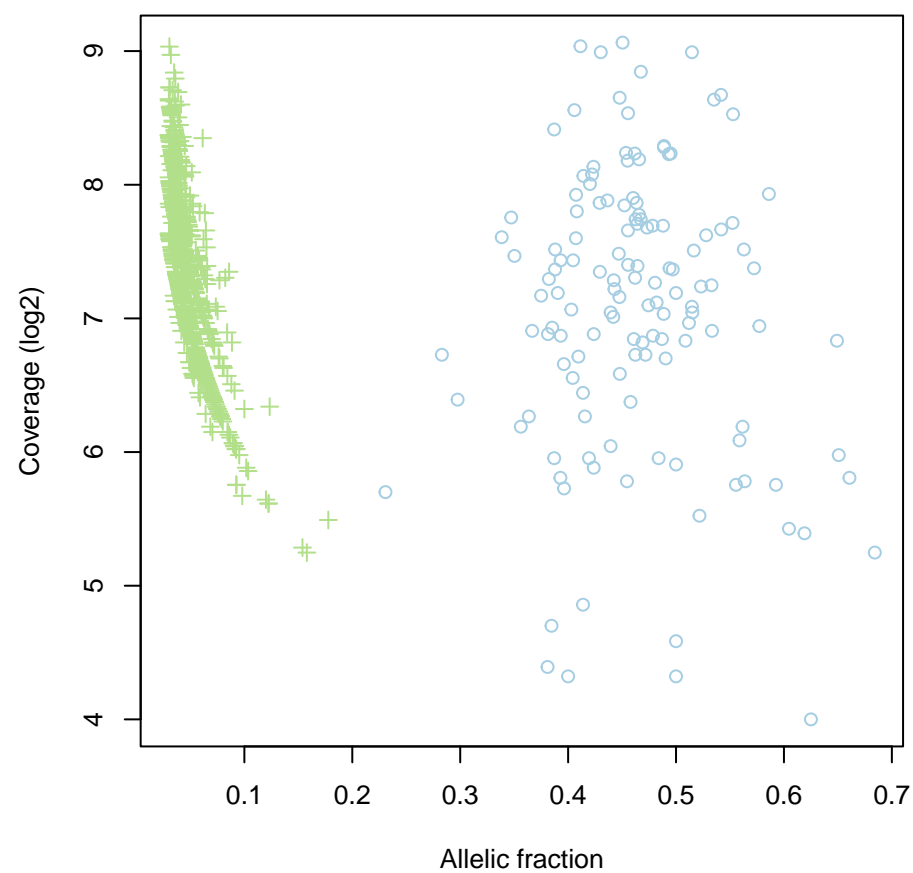
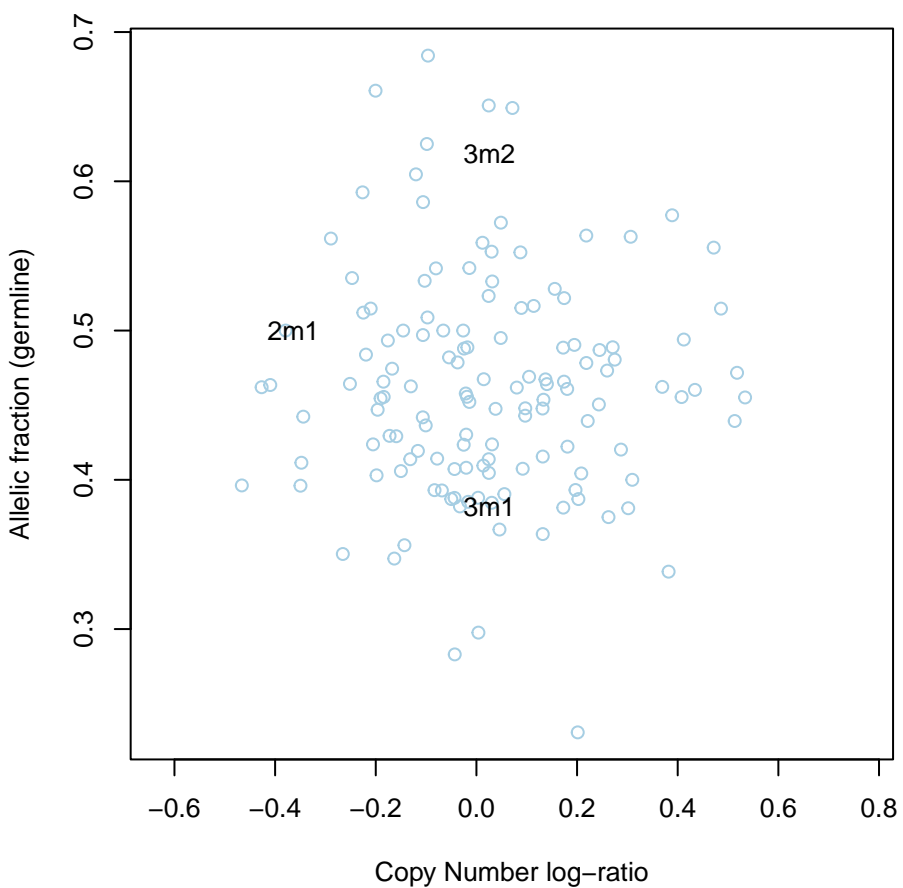


Purity: 0.62 Tumor ploidy: 2.924 SNV log-likelihood: -1190.31 GoF: 19.8% Mean coverage: 176;149

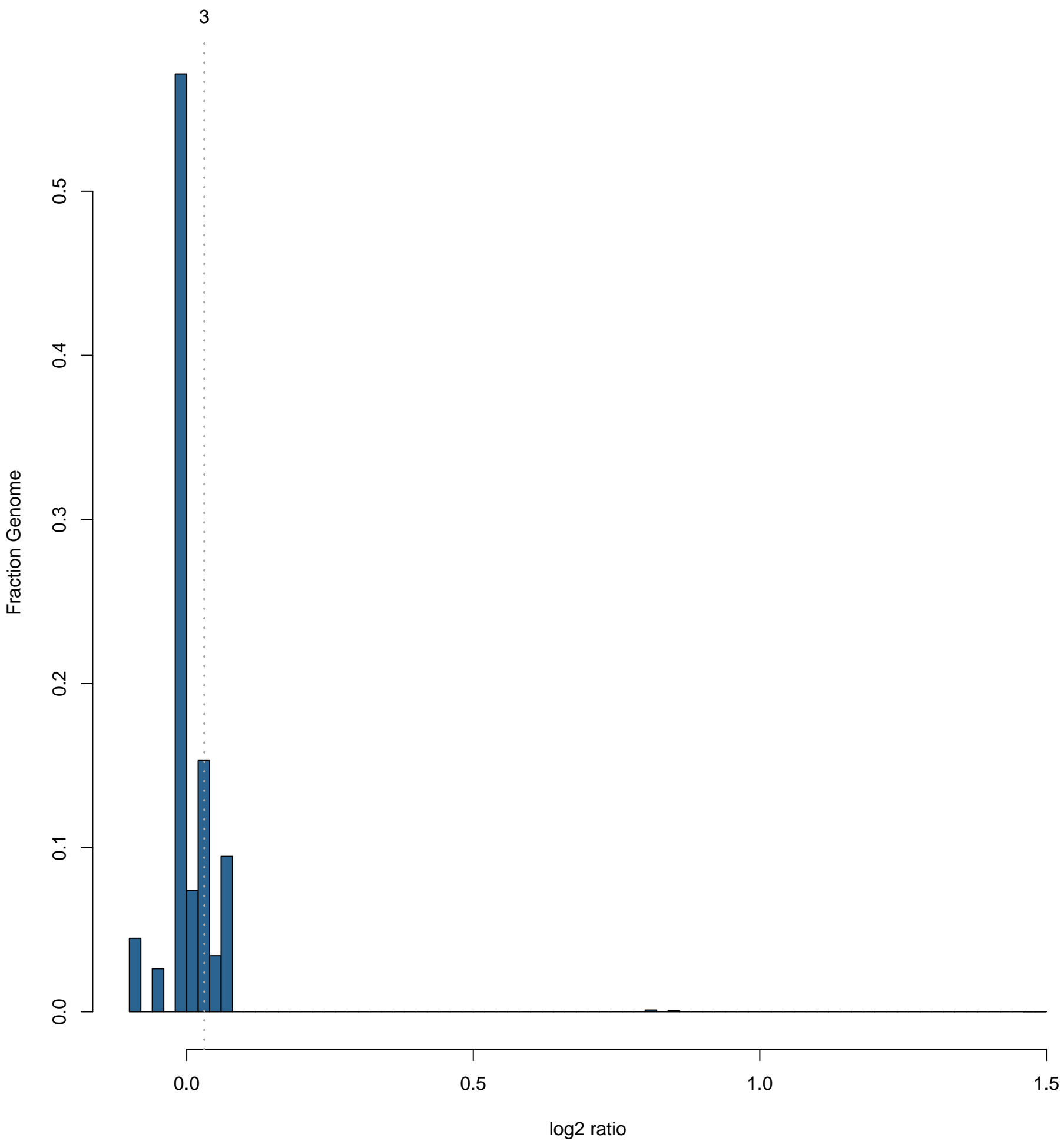


SCNA-fit log-likelihood: -14670.53

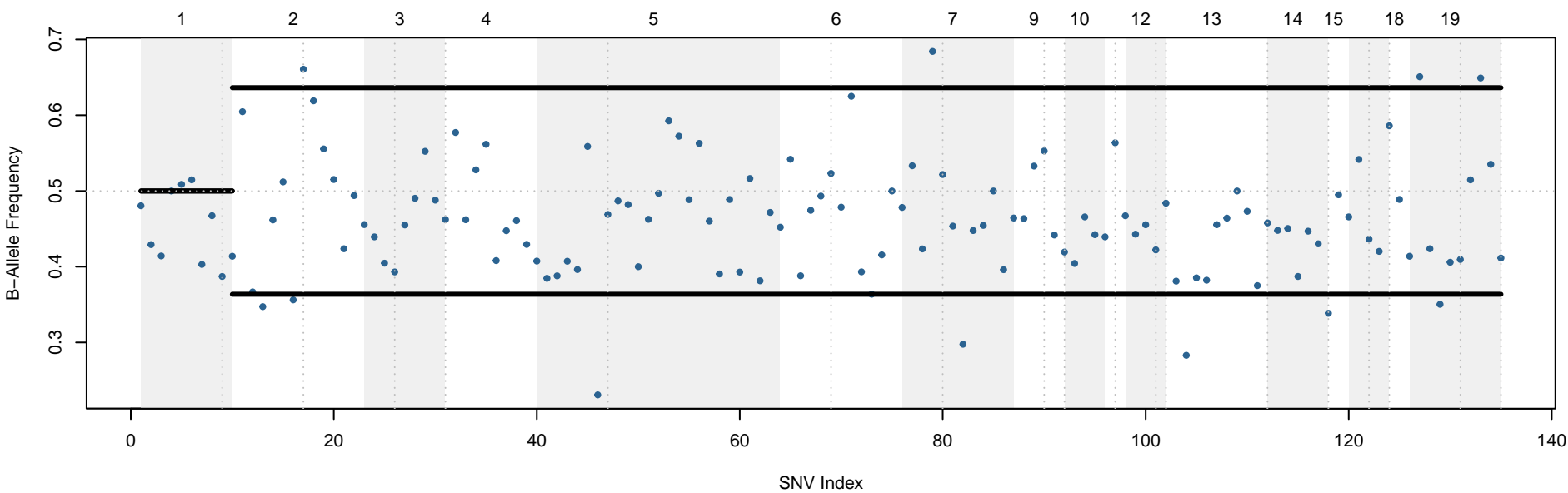




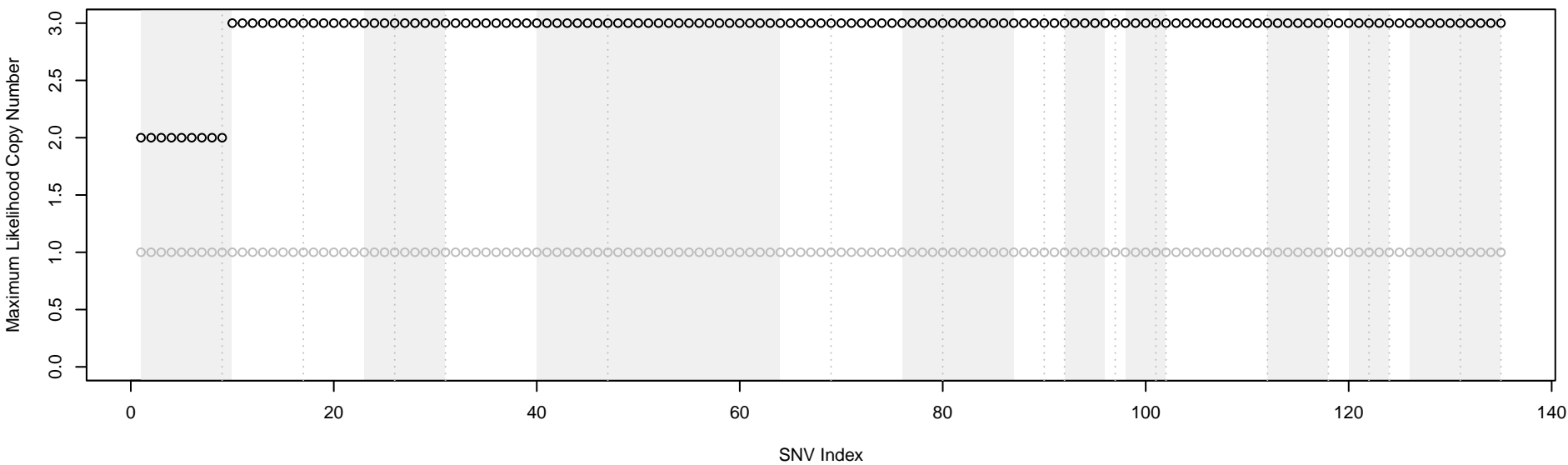
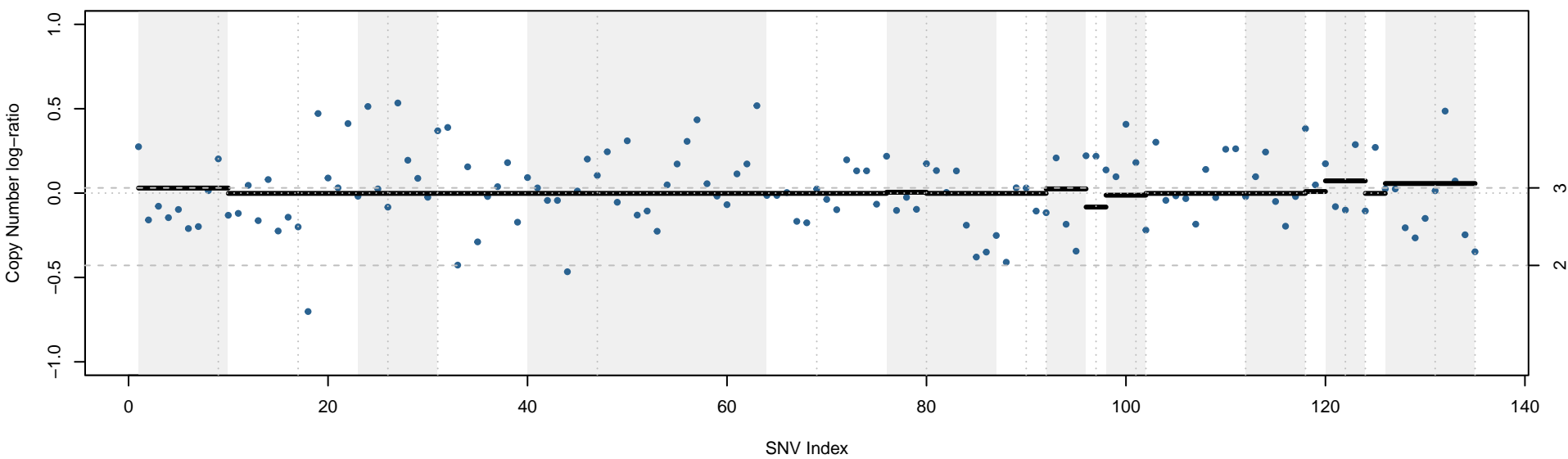
Purity: 0.75 Tumor ploidy: 2.923

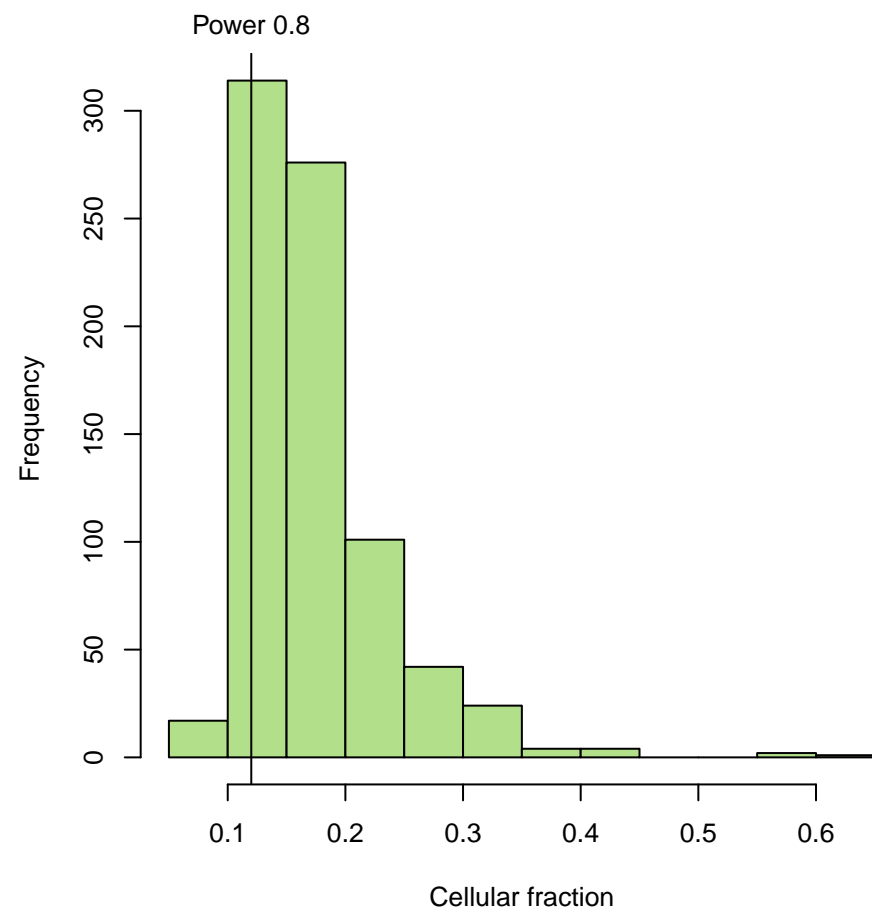
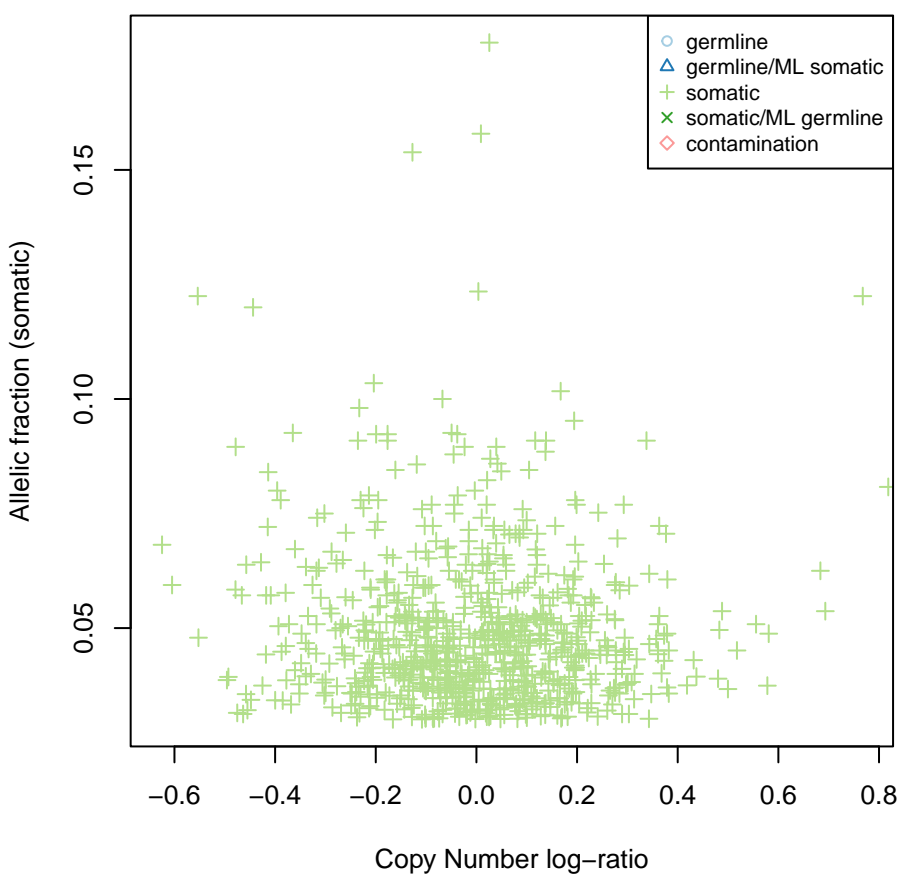
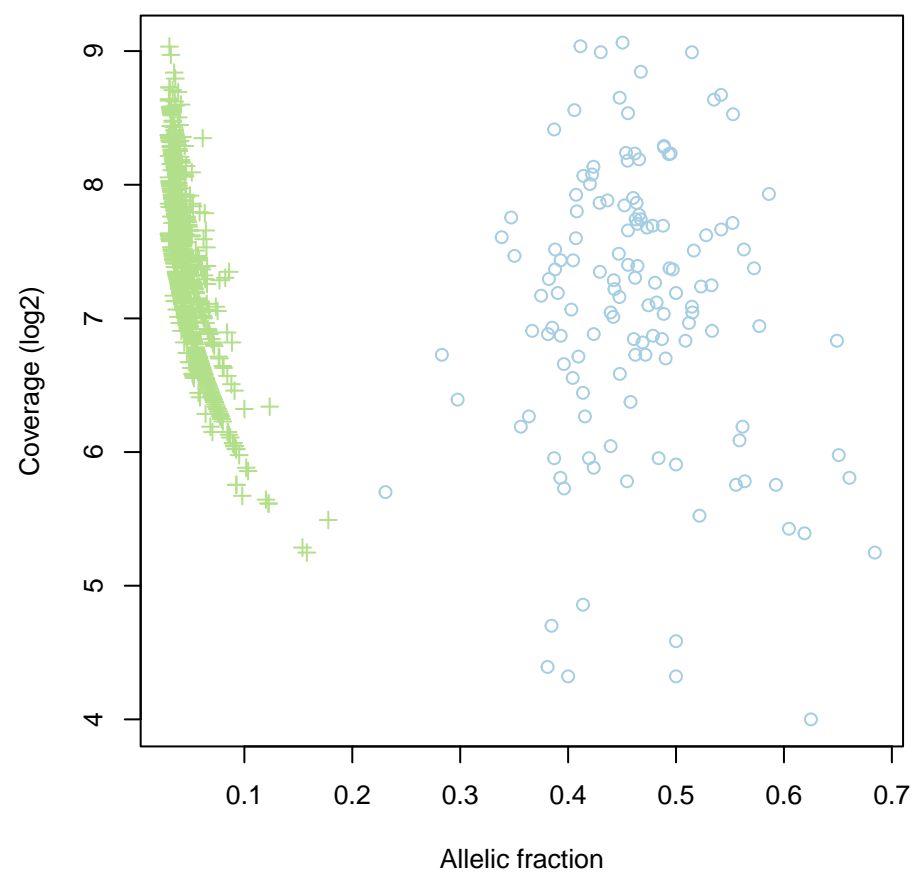
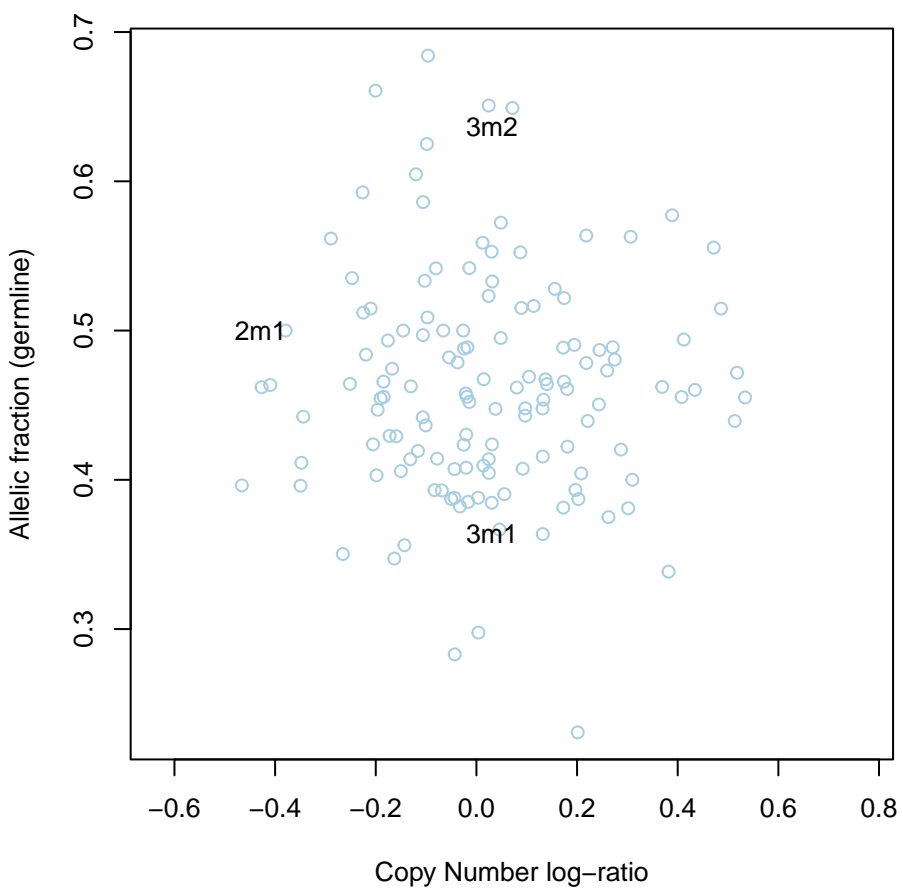


Purity: 0.75 Tumor ploidy: 2.923 SNV log-likelihood: -2184.91 GoF: 0% Mean coverage: 176;149

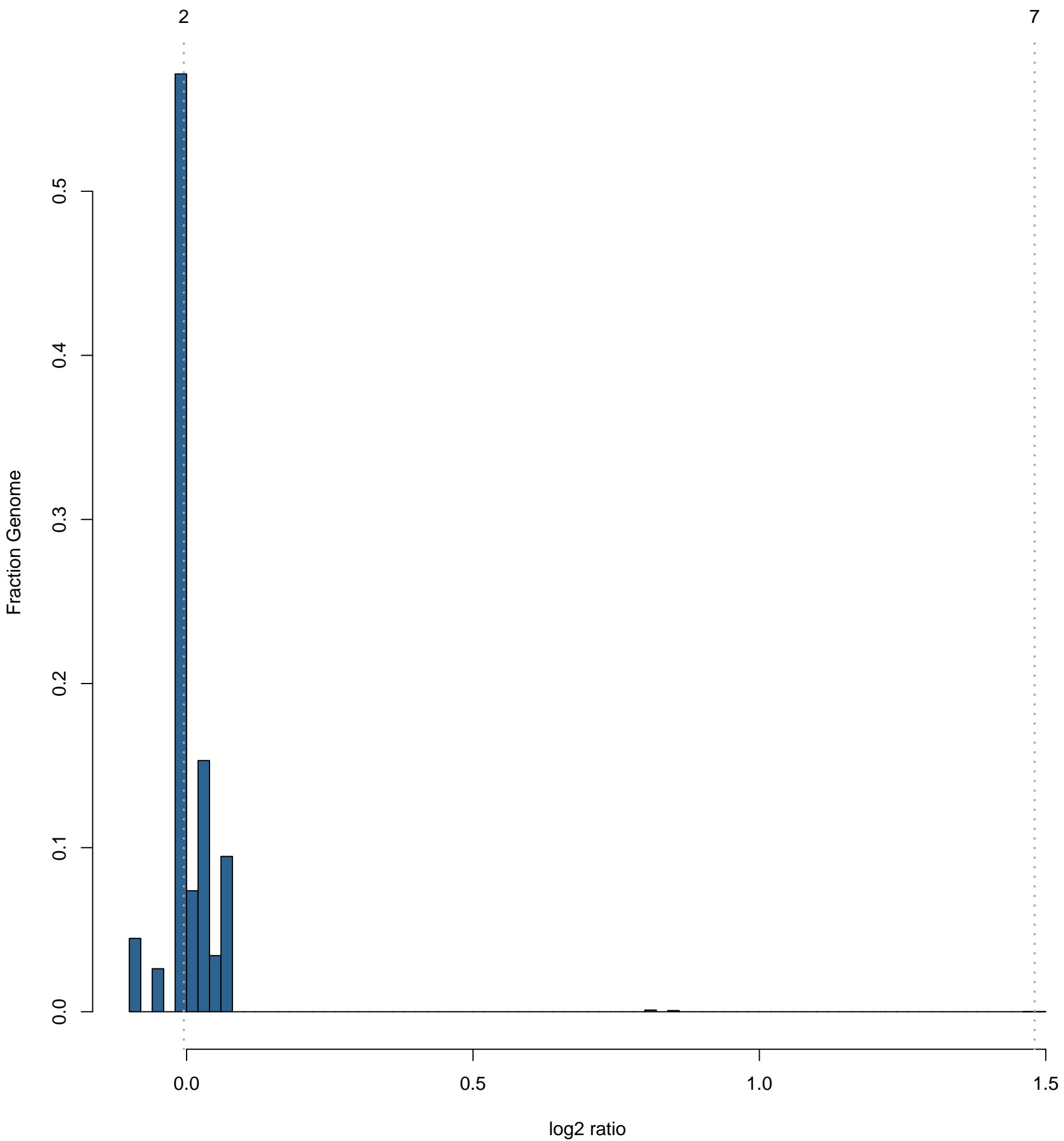


SCNA-fit log-likelihood: -14859.73



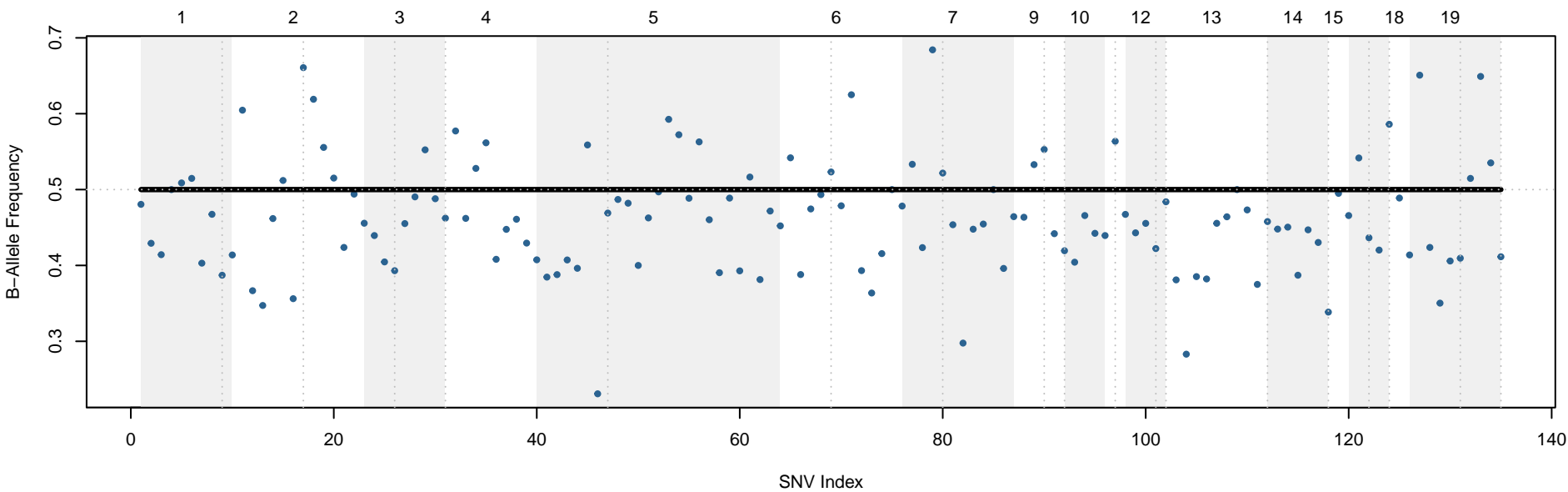


Purity: 0.72 Tumor ploidy: 2.009





Purity: 0.72 Tumor ploidy: 2.009 SNV log-likelihood: -3644.73 GoF: 0% Mean coverage: 176;149



SCNA-fit log-likelihood: -14155.63

