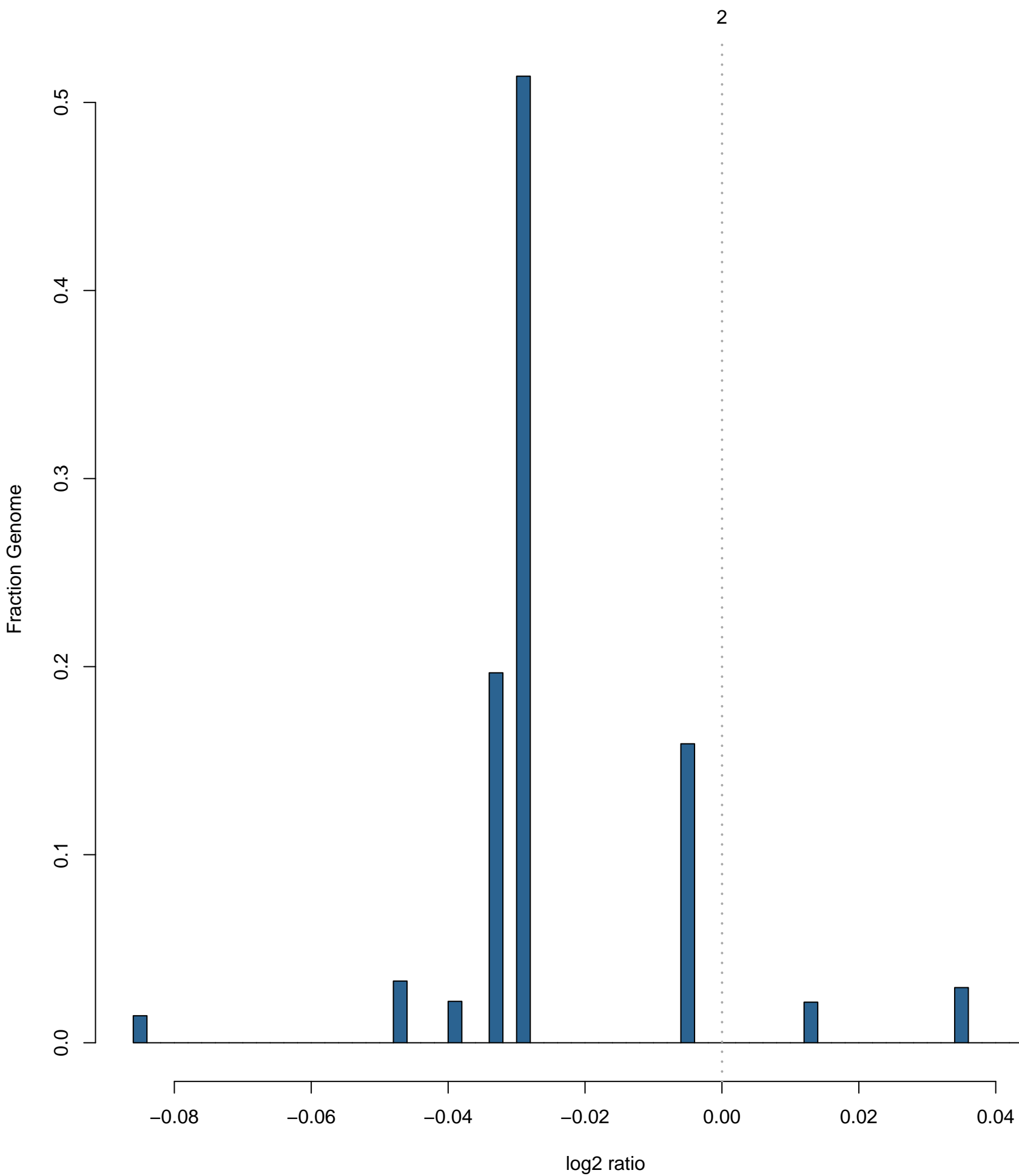
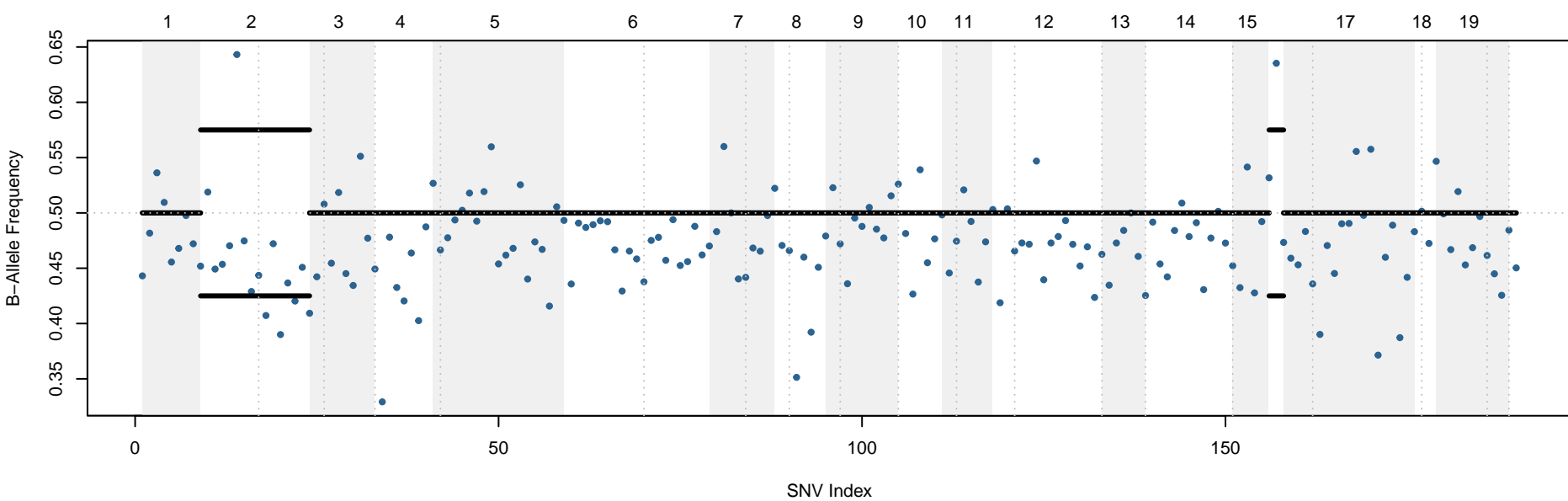


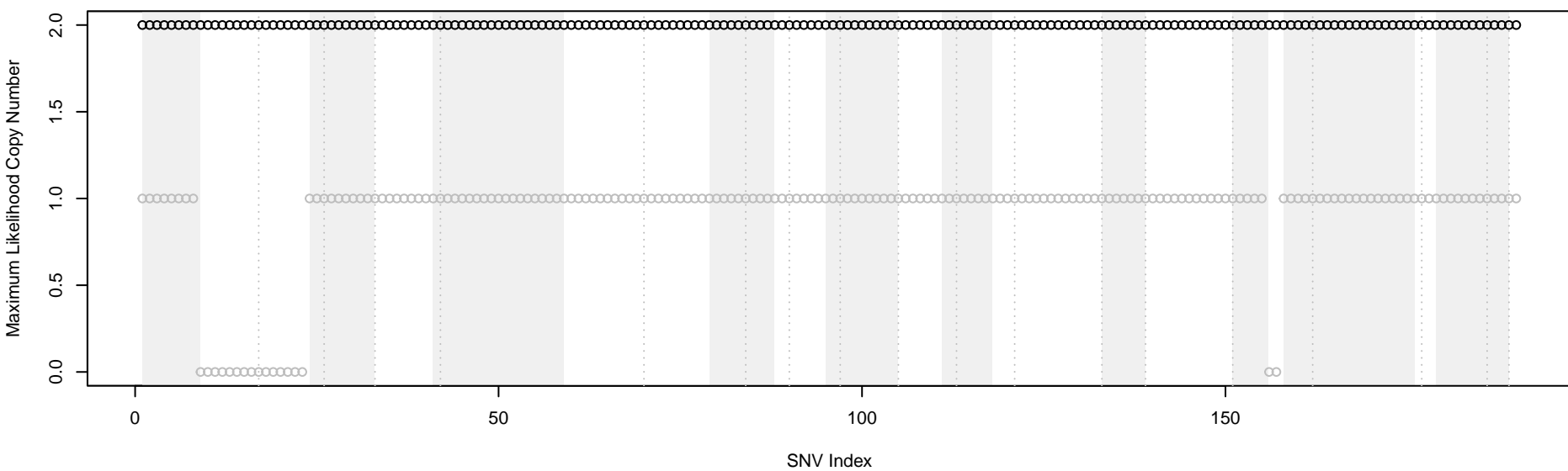
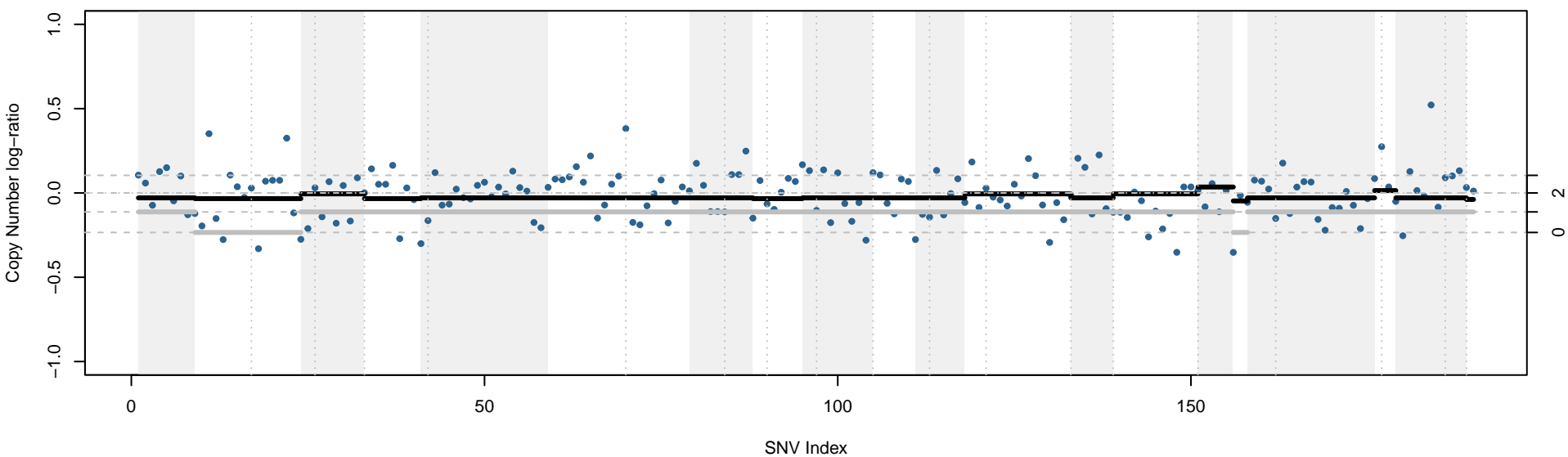
Purity: 0.15 Tumor ploidy: 2

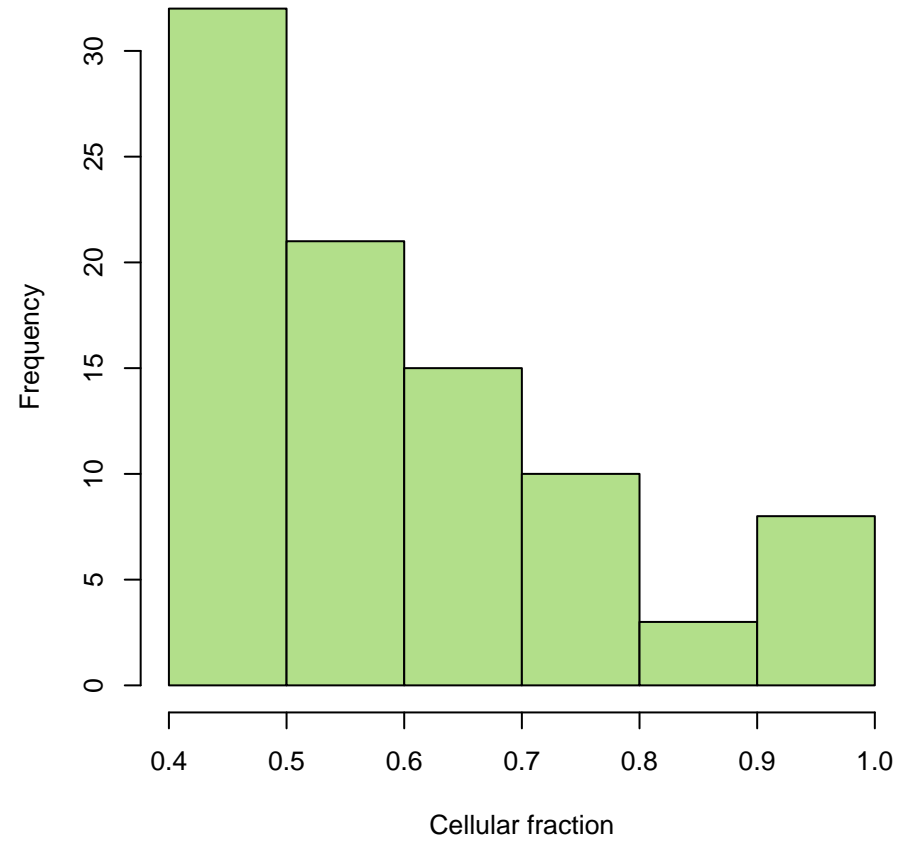
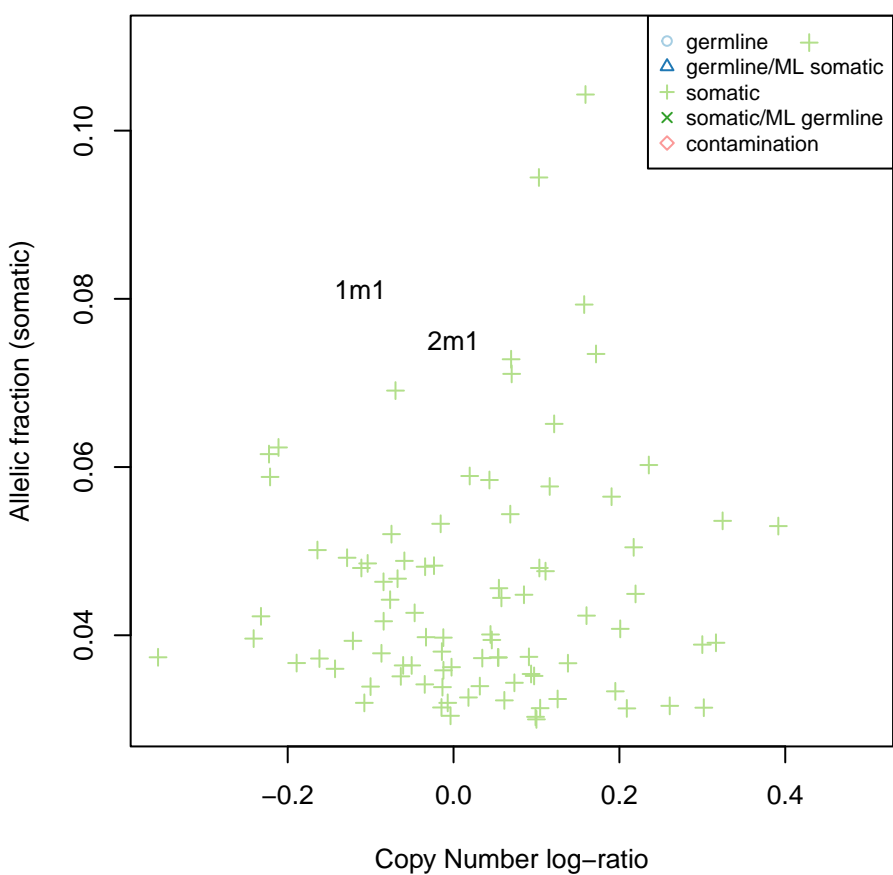
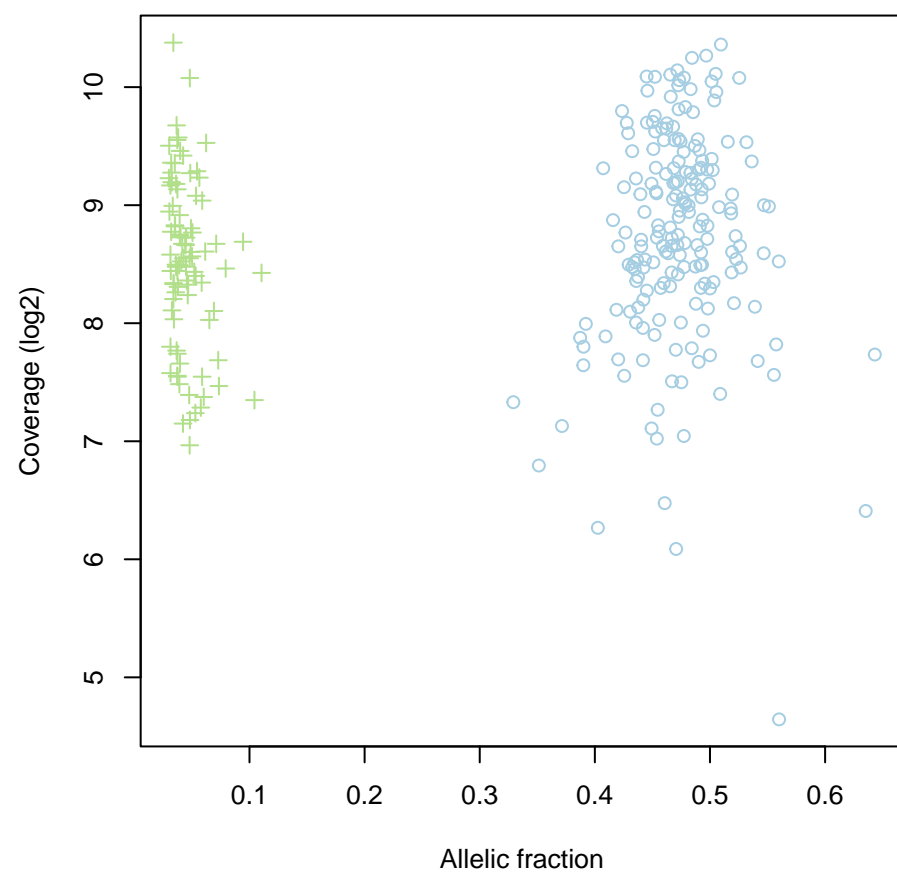
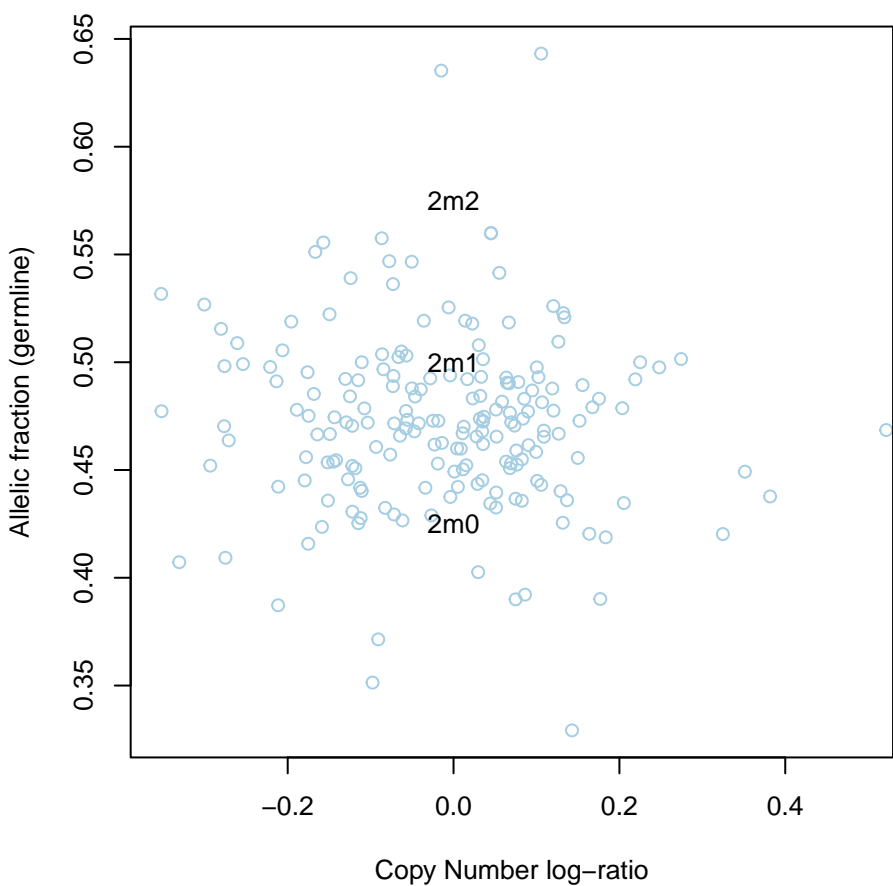


Purity: 0.15 Tumor ploidy: 2 SNV log-likelihood: 173.22 GoF: 96.2% Mean coverage: 476;548

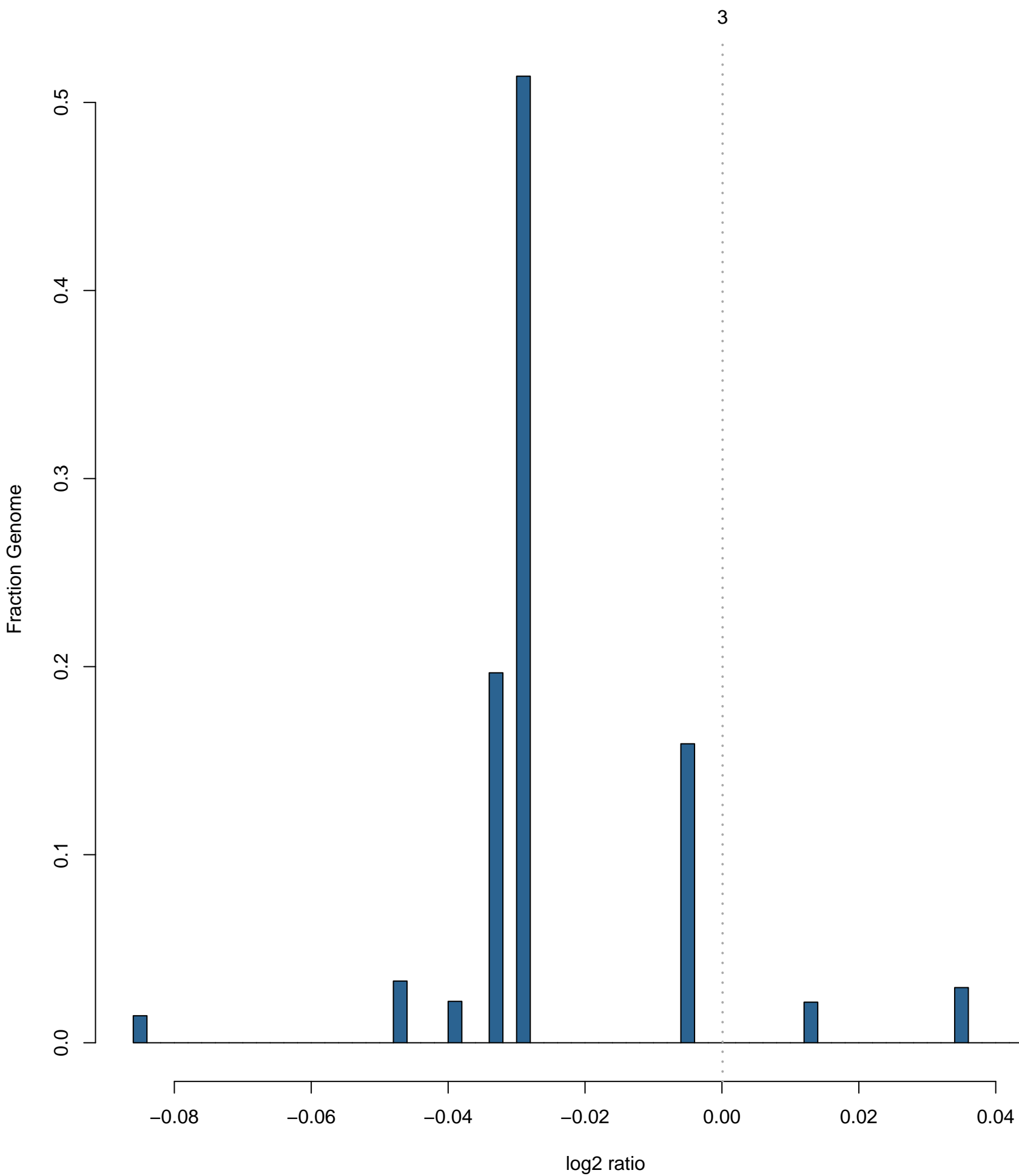


SCNA-fit log-likelihood: -14289.25

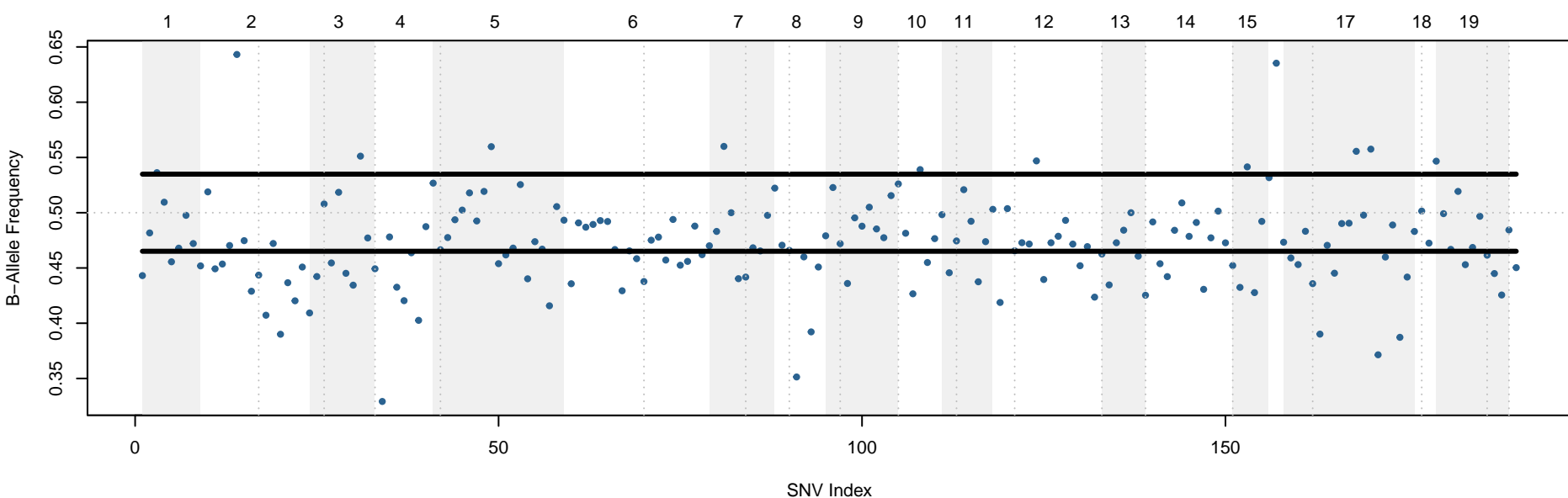




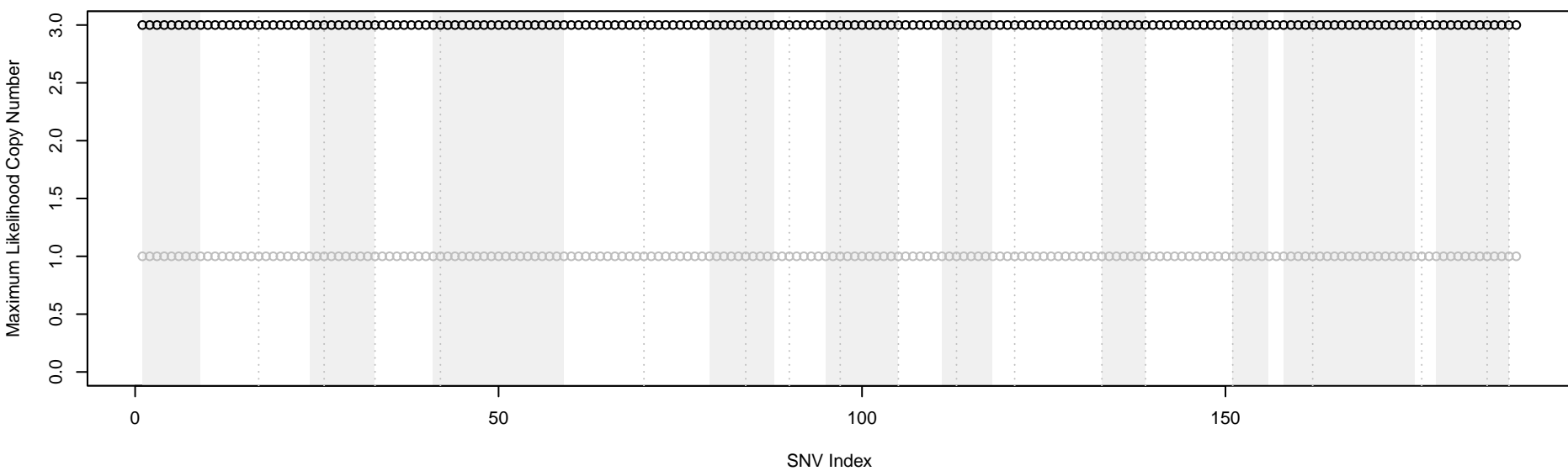
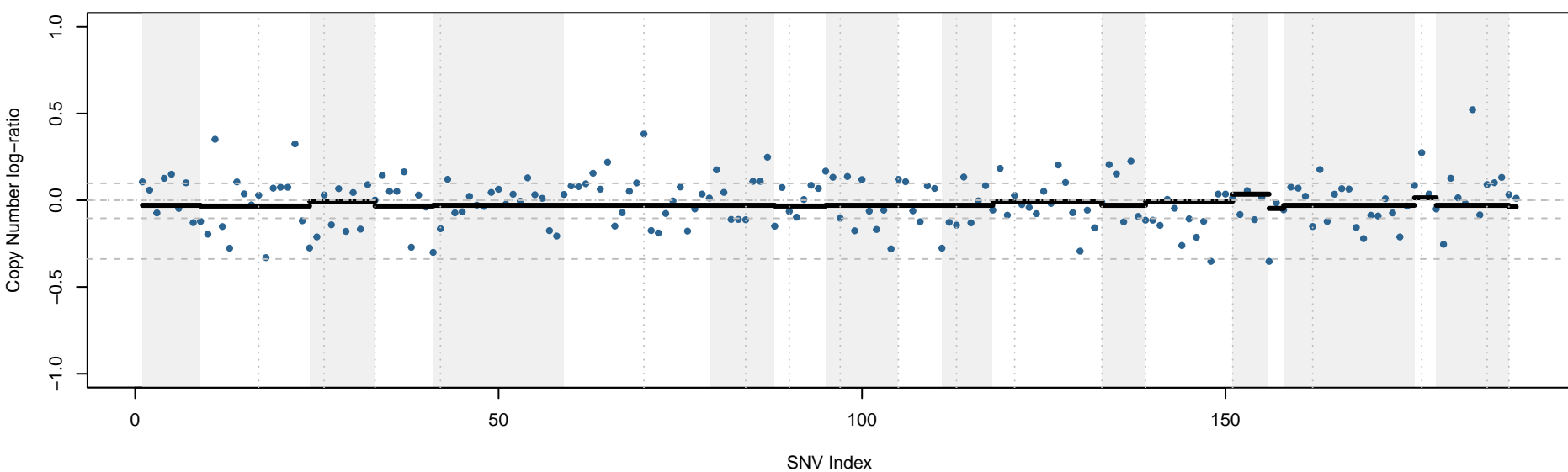
Purity: 0.15 Tumor ploidy: 2.999

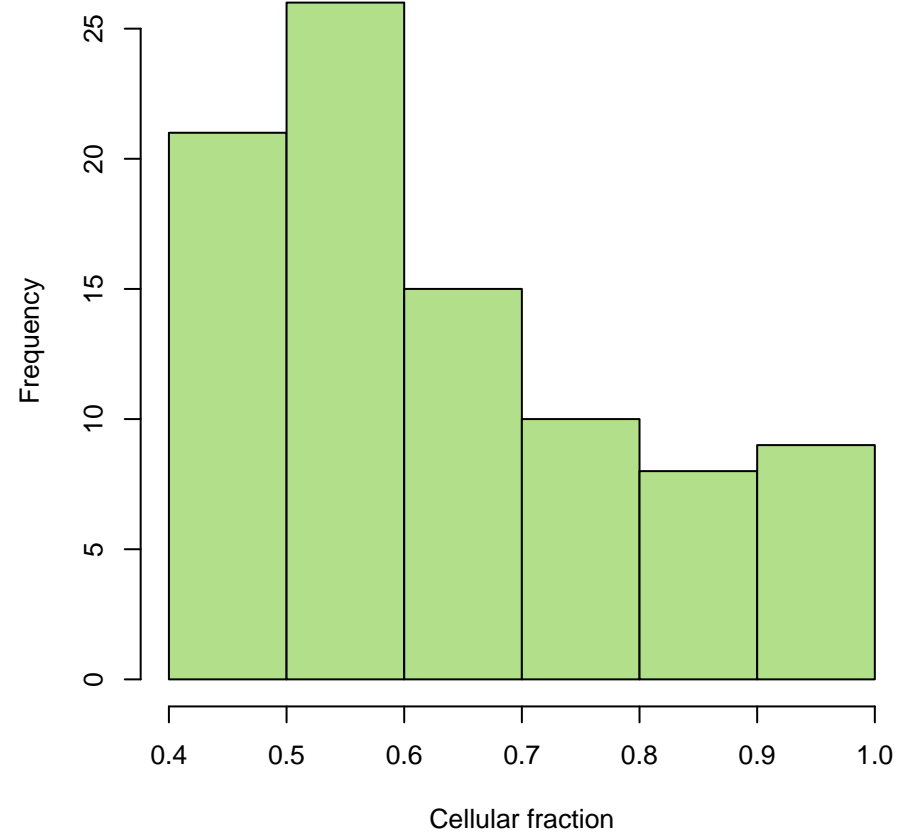
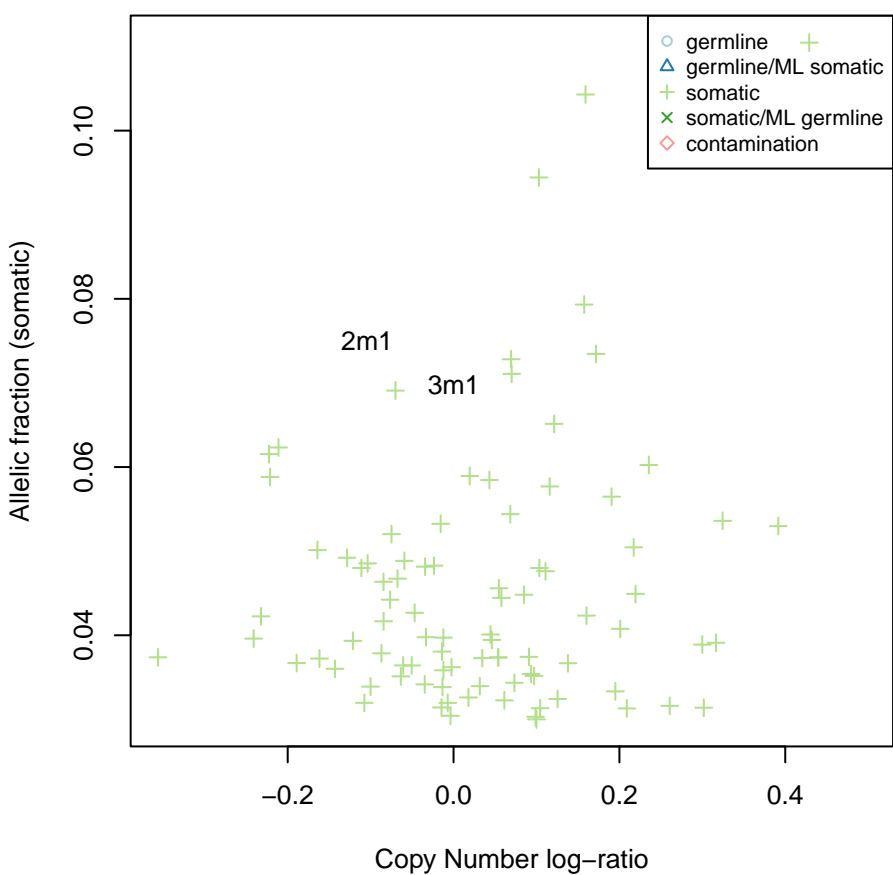
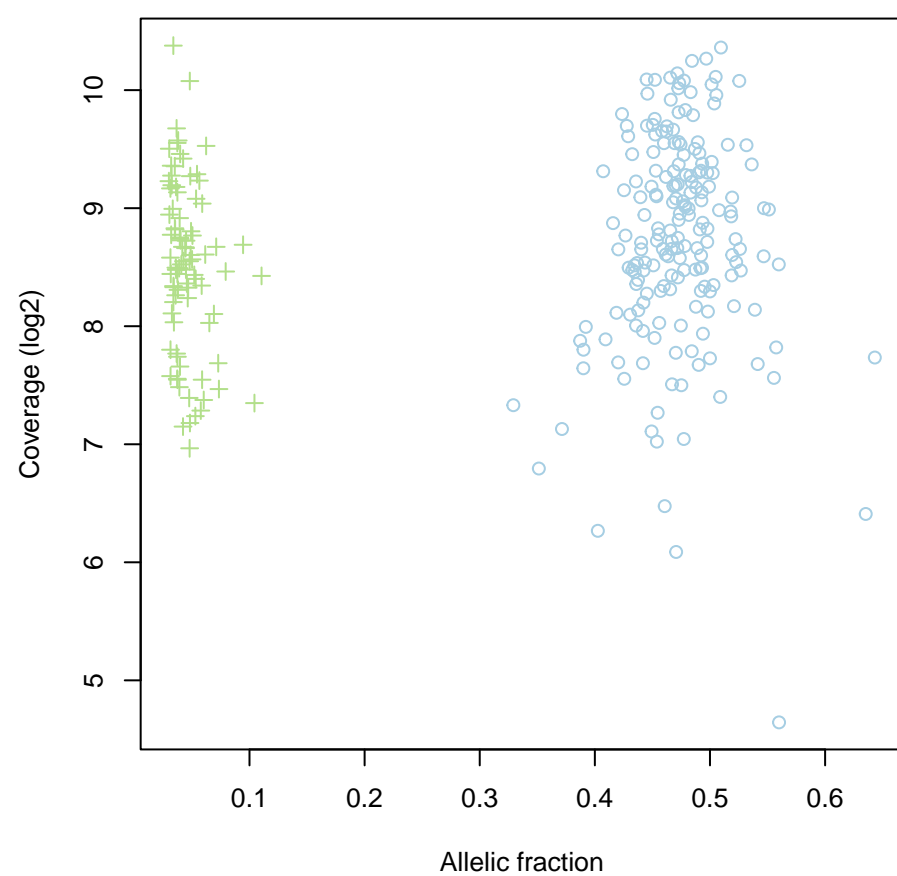
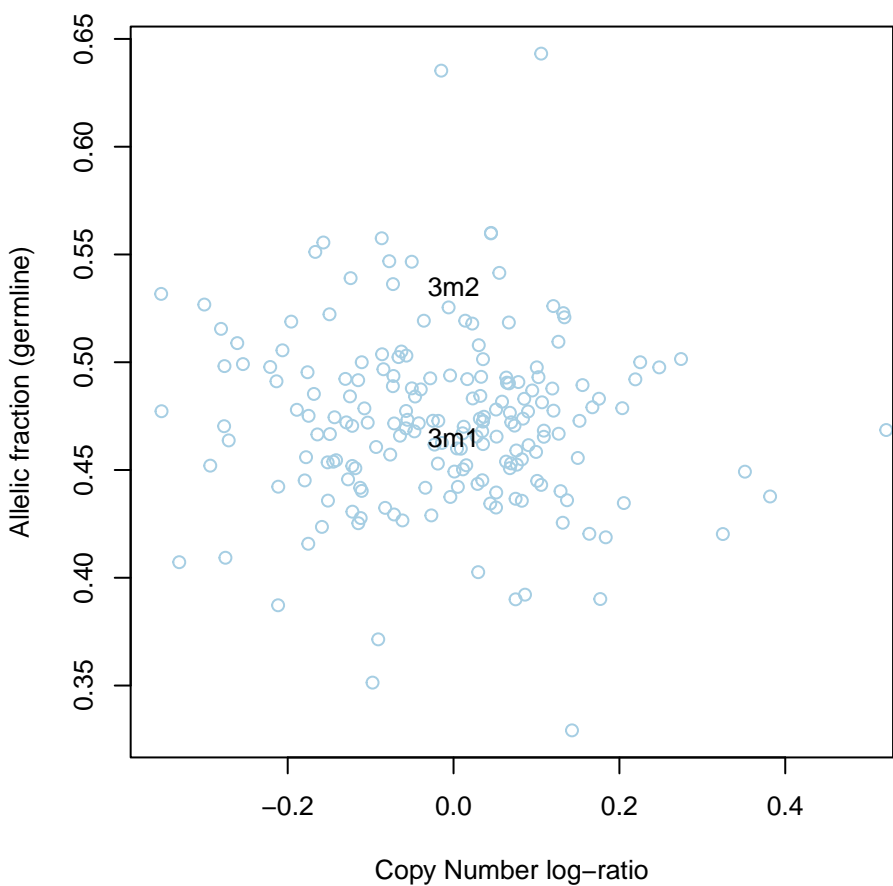


Purity: 0.15 Tumor ploidy: 2.999 SNV log-likelihood: 26.72 GoF: 97.9% Mean coverage: 476;548

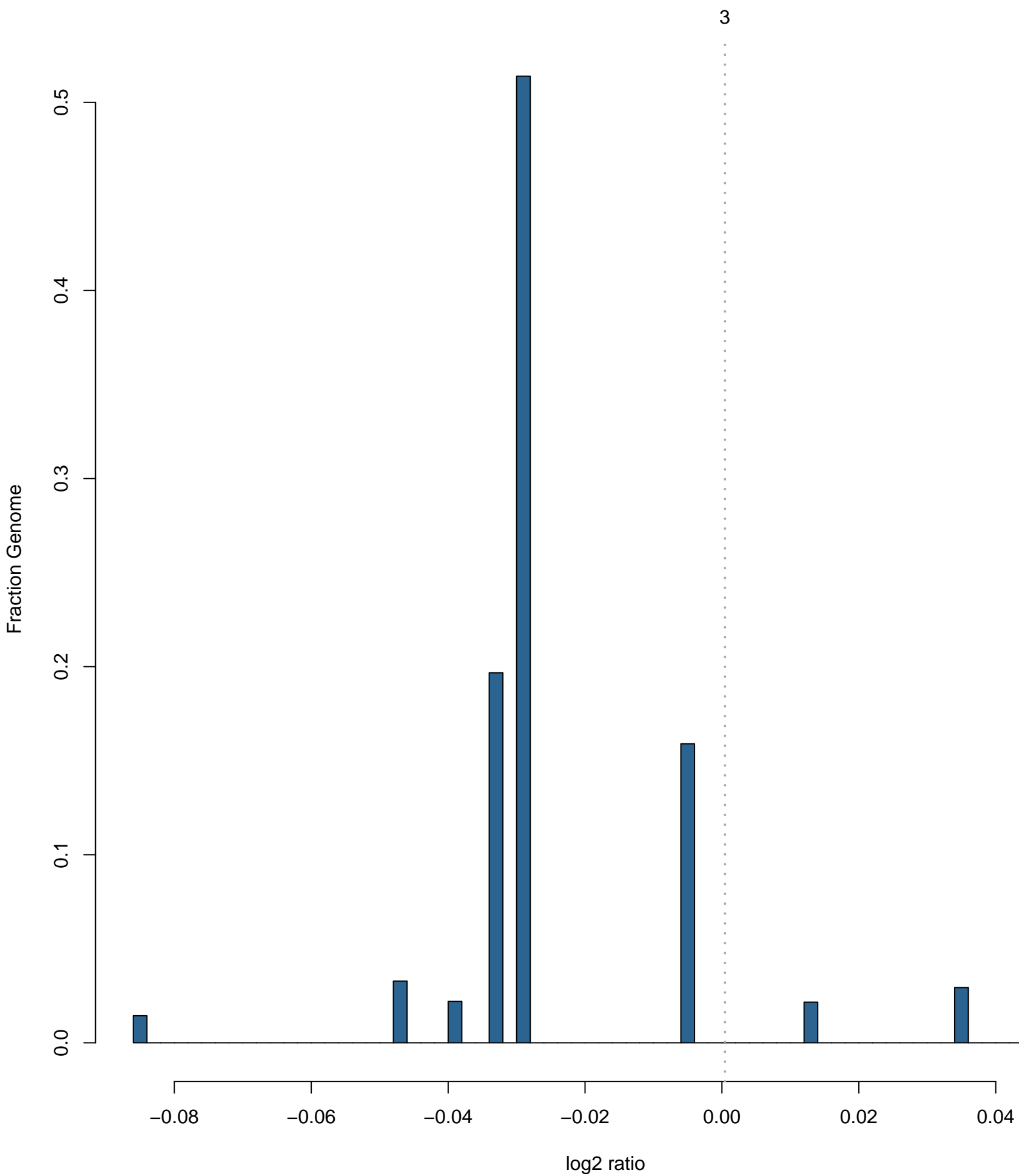


SCNA-fit log-likelihood: -14302.13

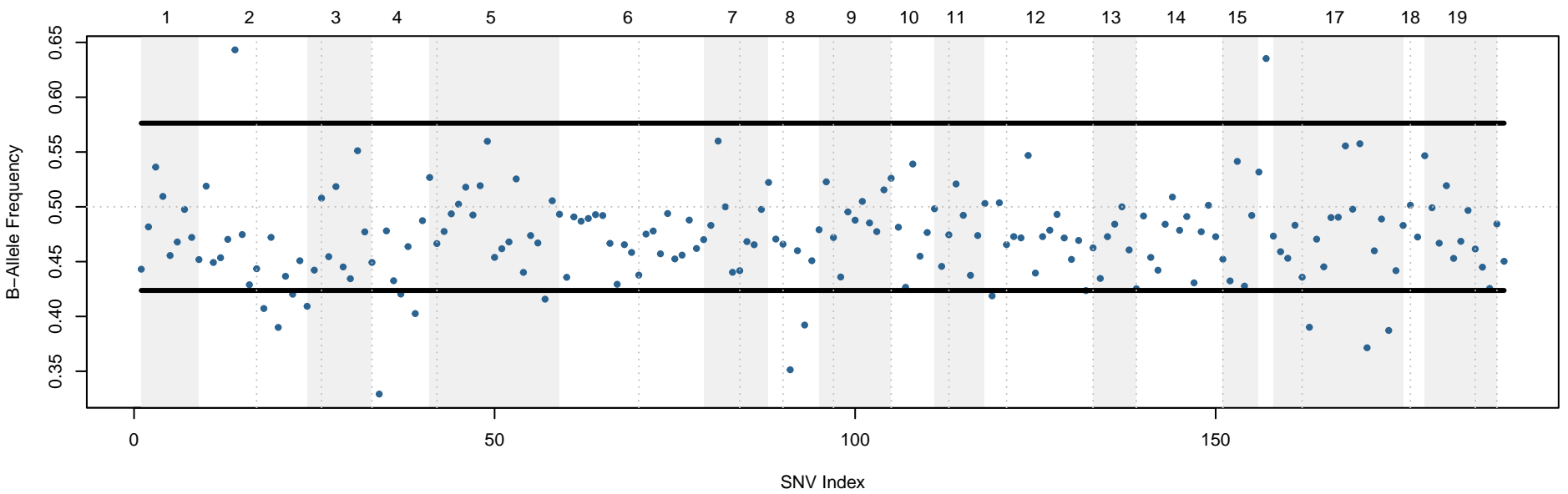




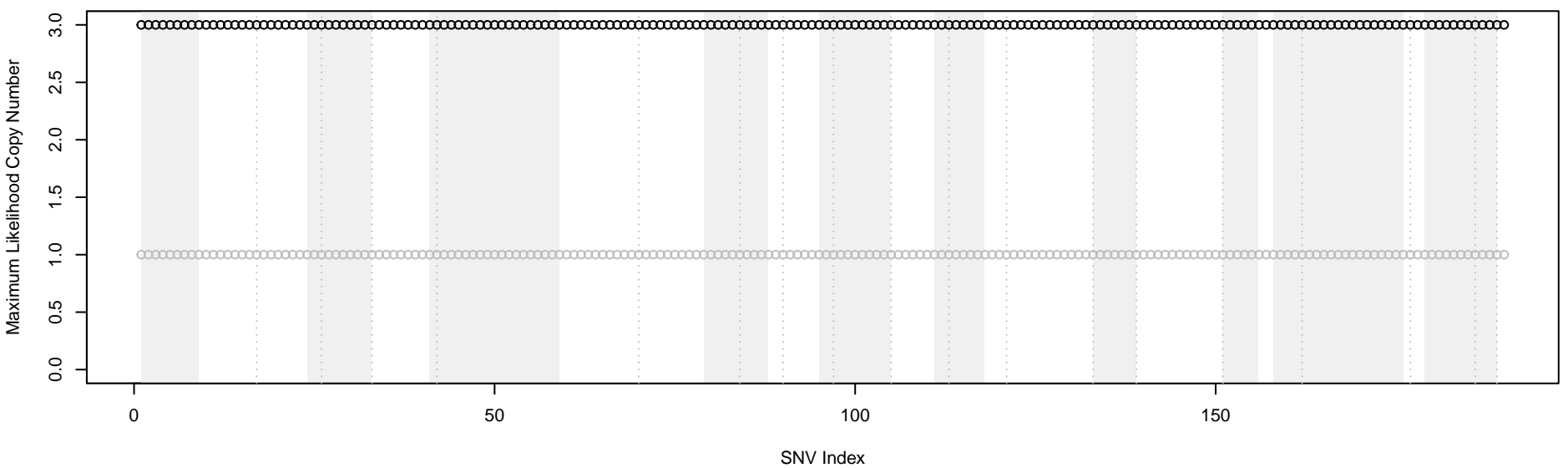
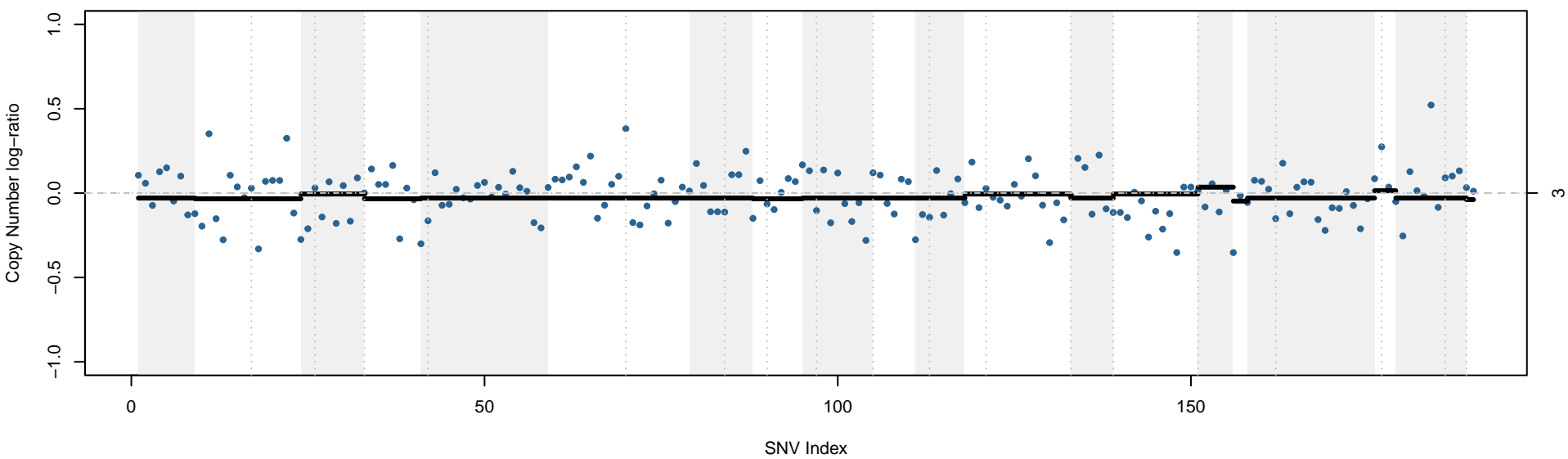
Purity: 0.36 Tumor ploidy: 2.998



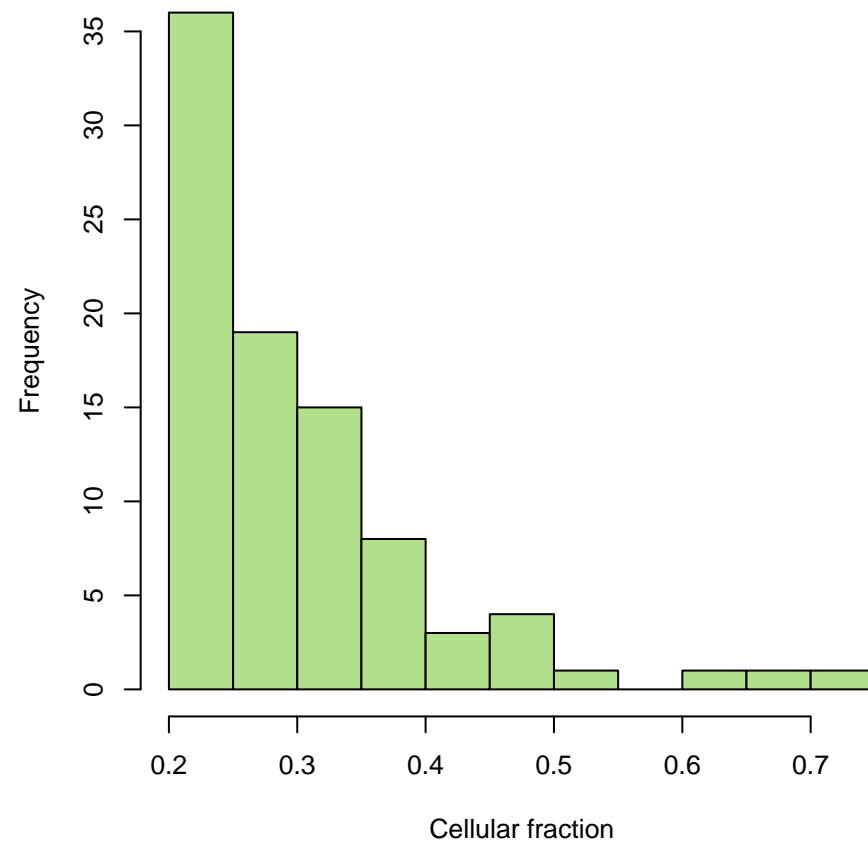
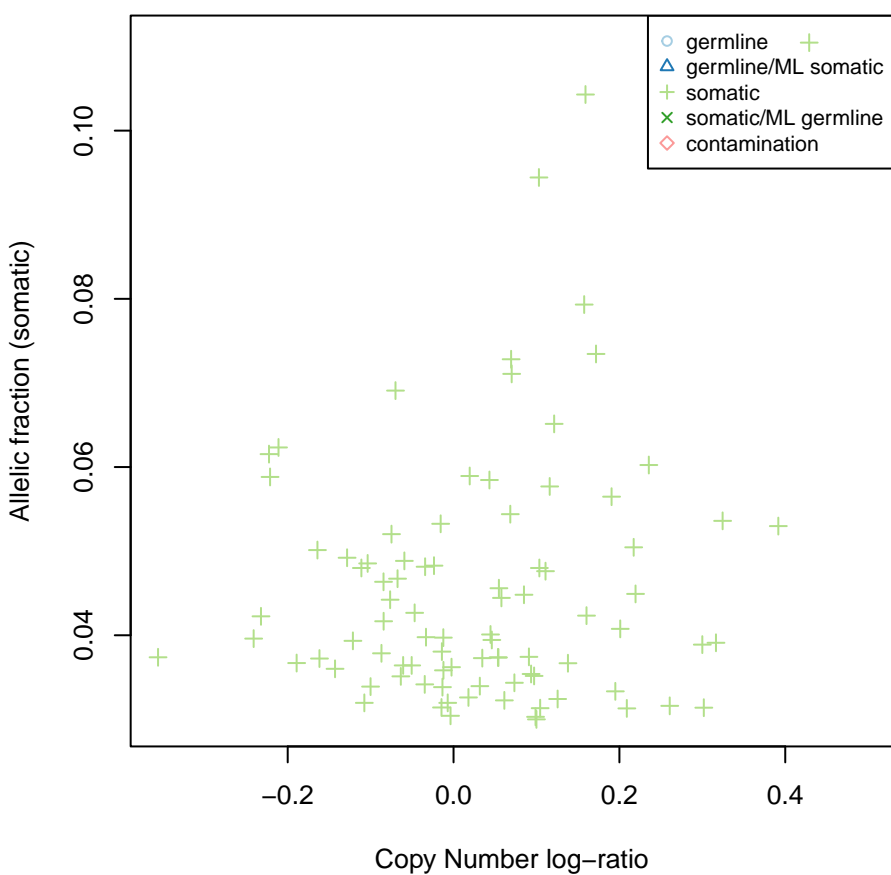
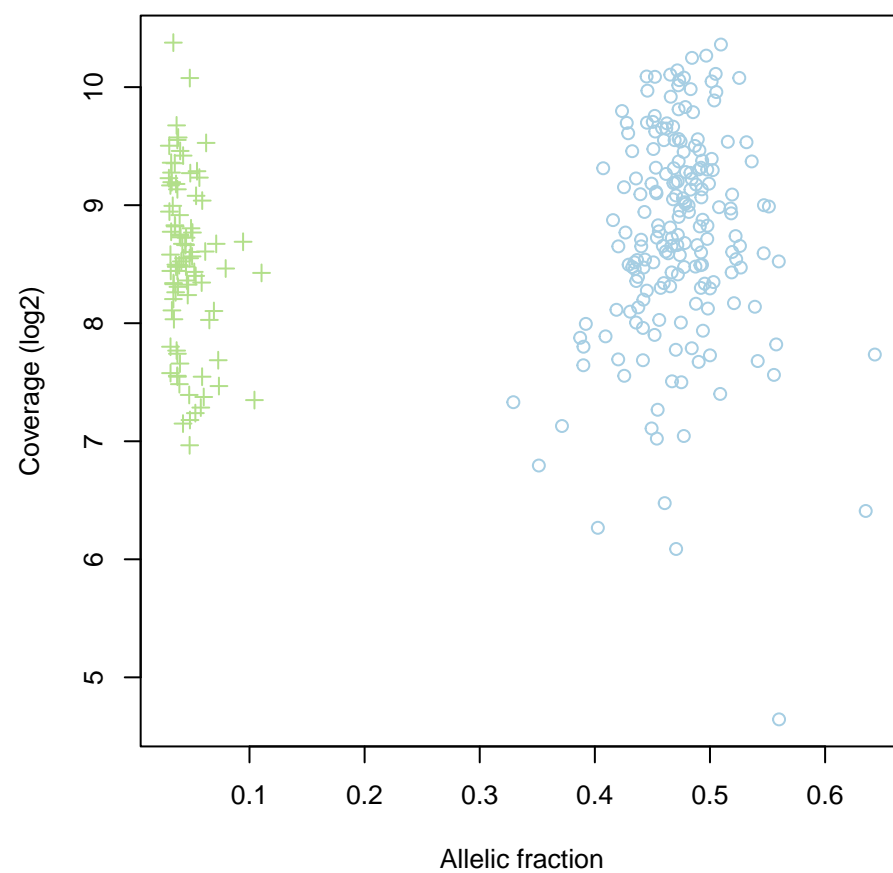
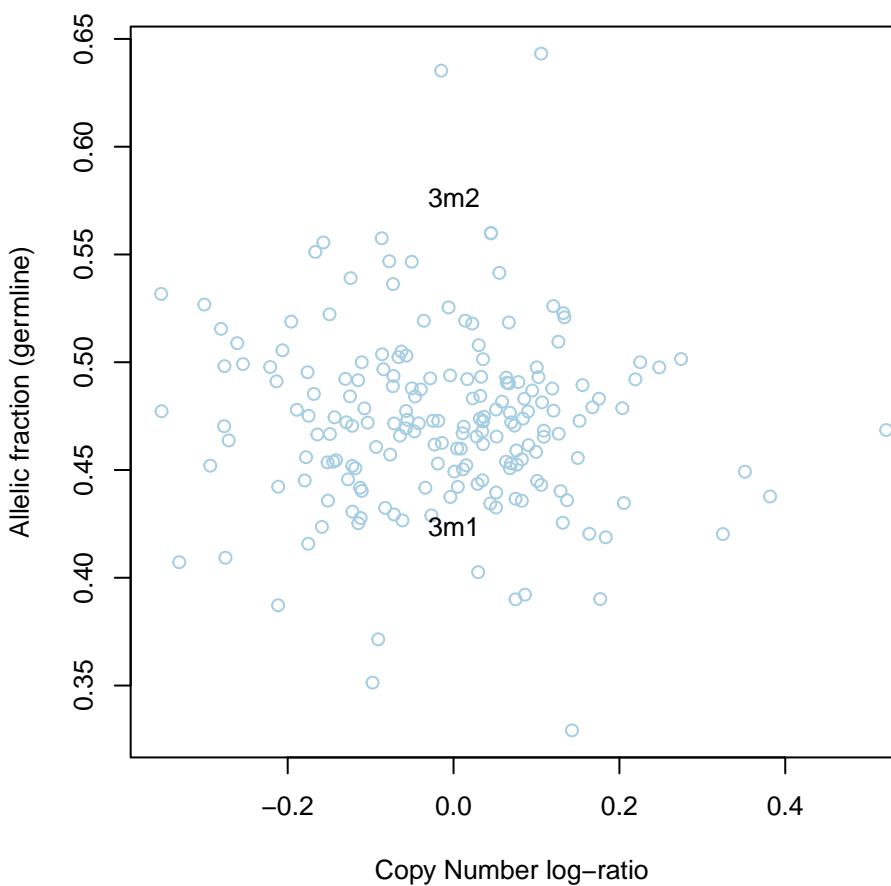
Purity: 0.36 Tumor ploidy: 2.998 SNV log-likelihood: -174.84 GoF: 86.3% Mean coverage: 476,548



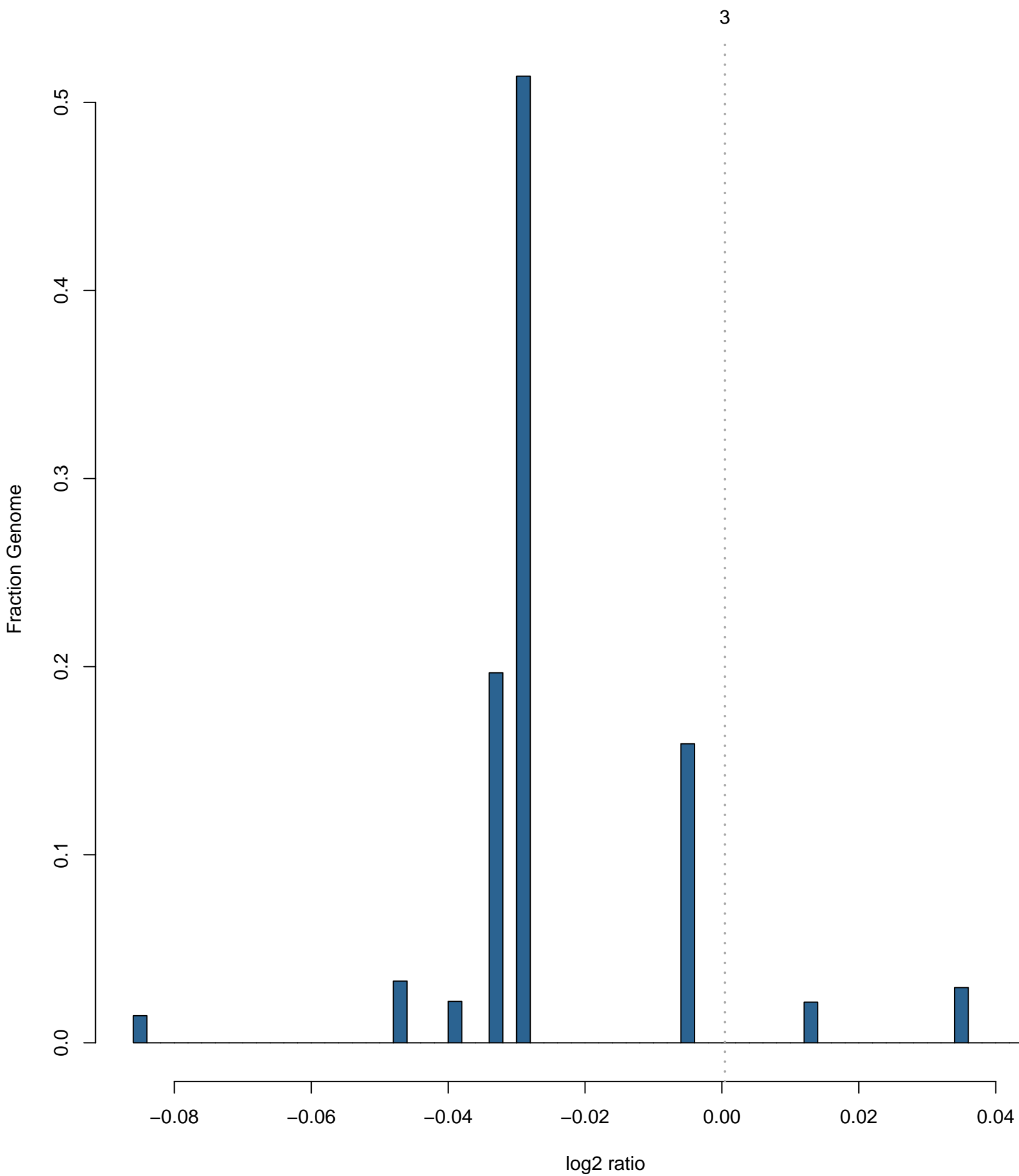
SCNA-fit log-likelihood: -14300.49



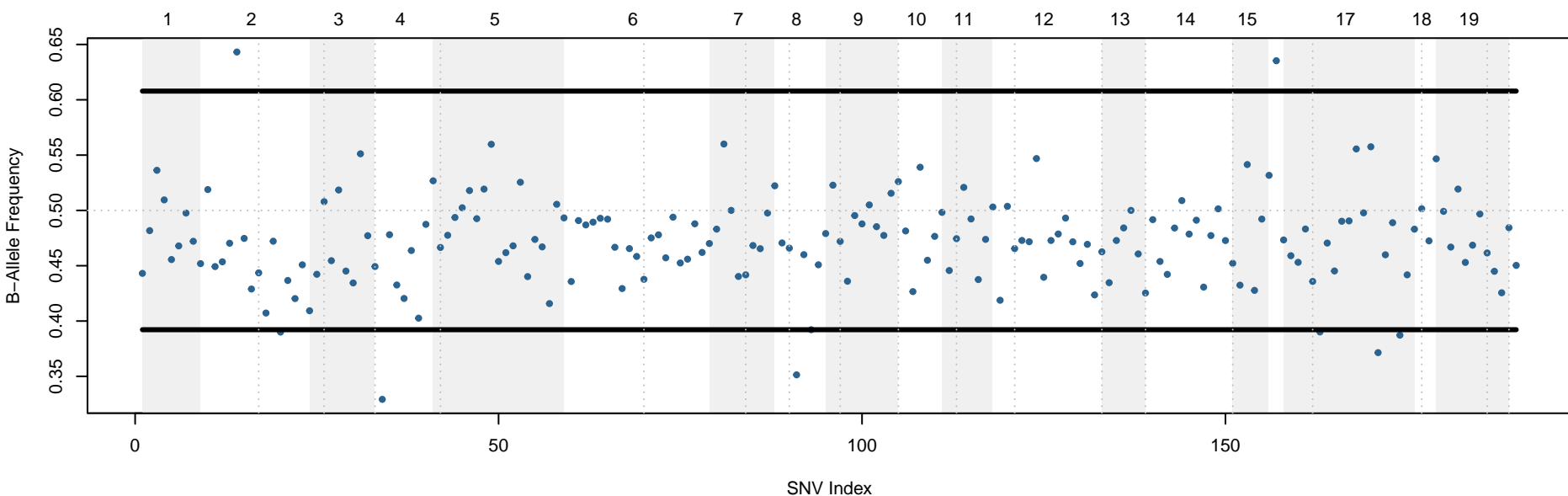




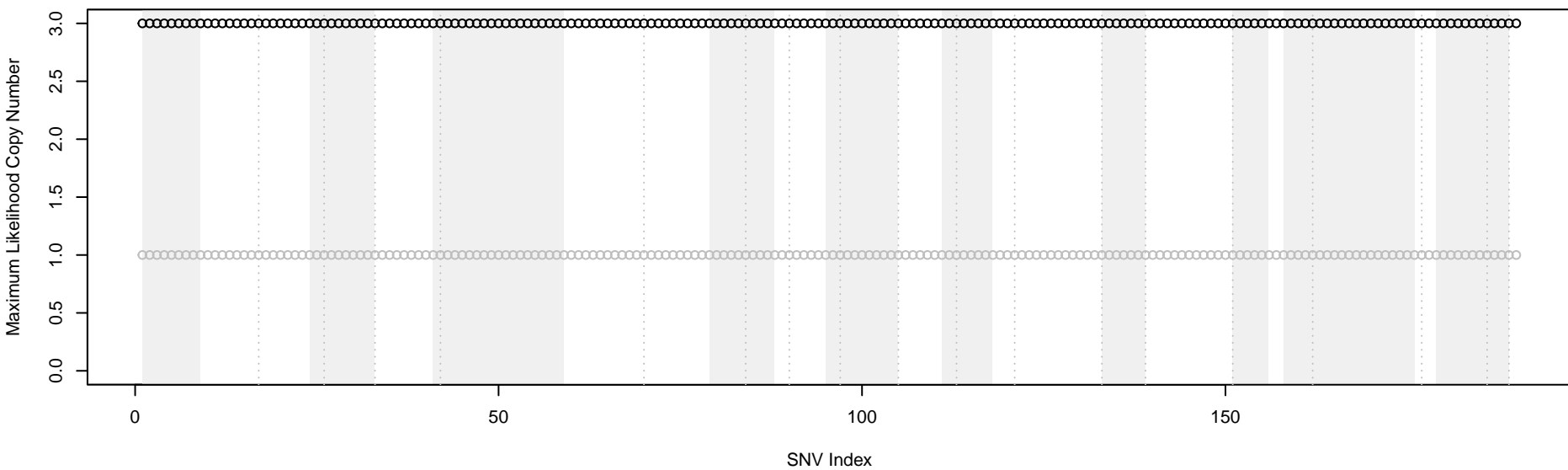
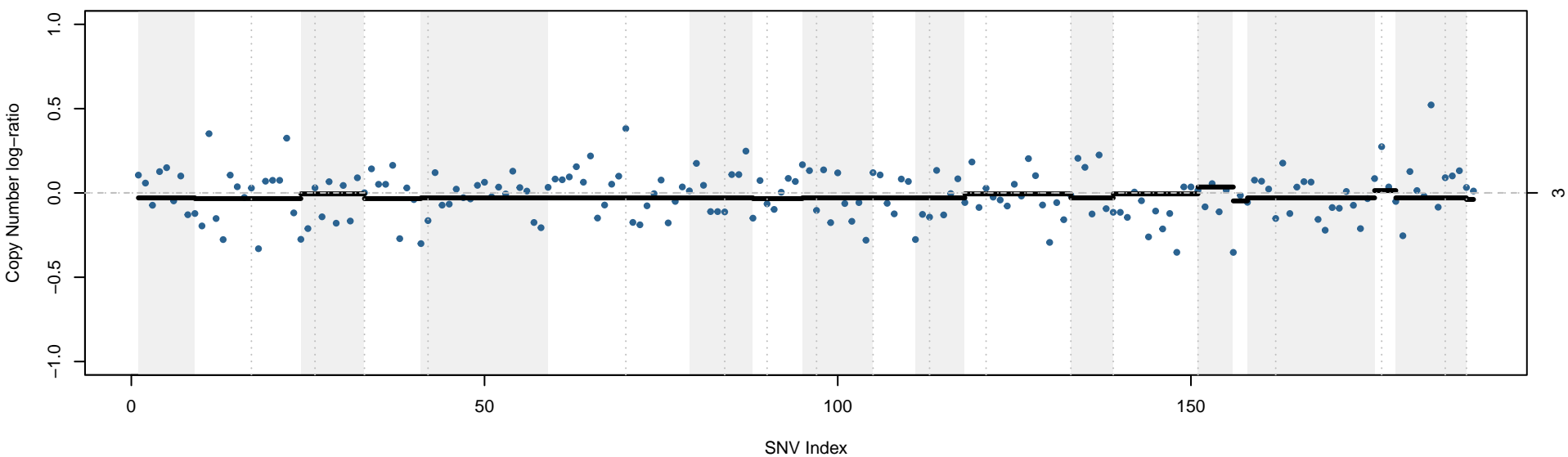
Purity: 0.55 Tumor ploidy: 2.999

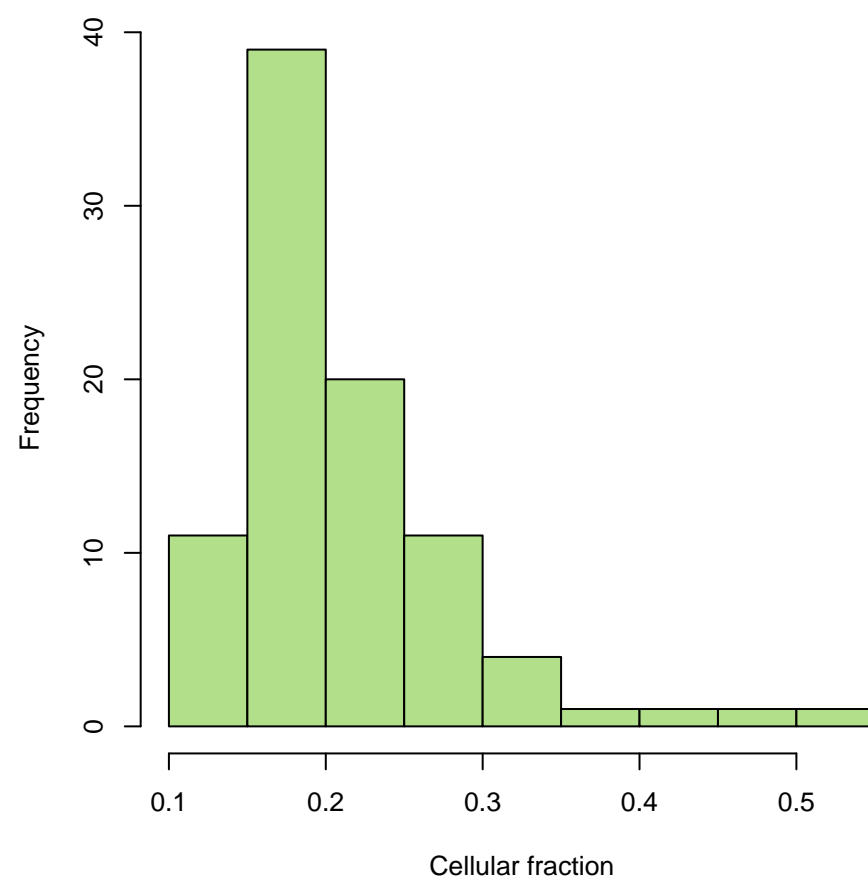
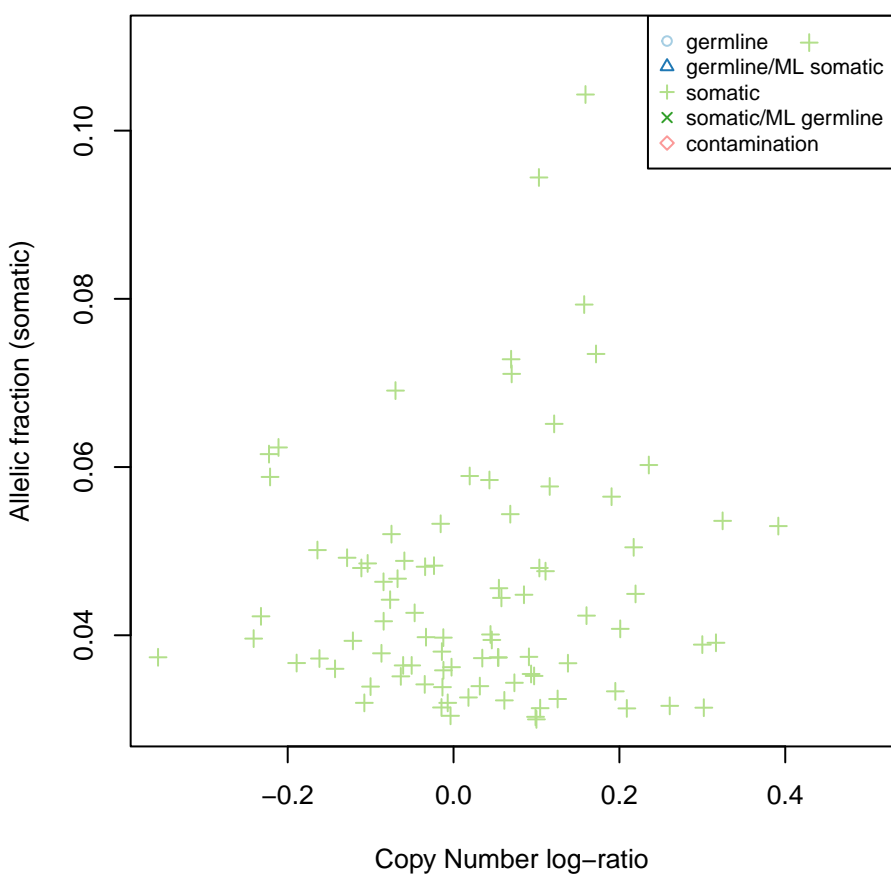
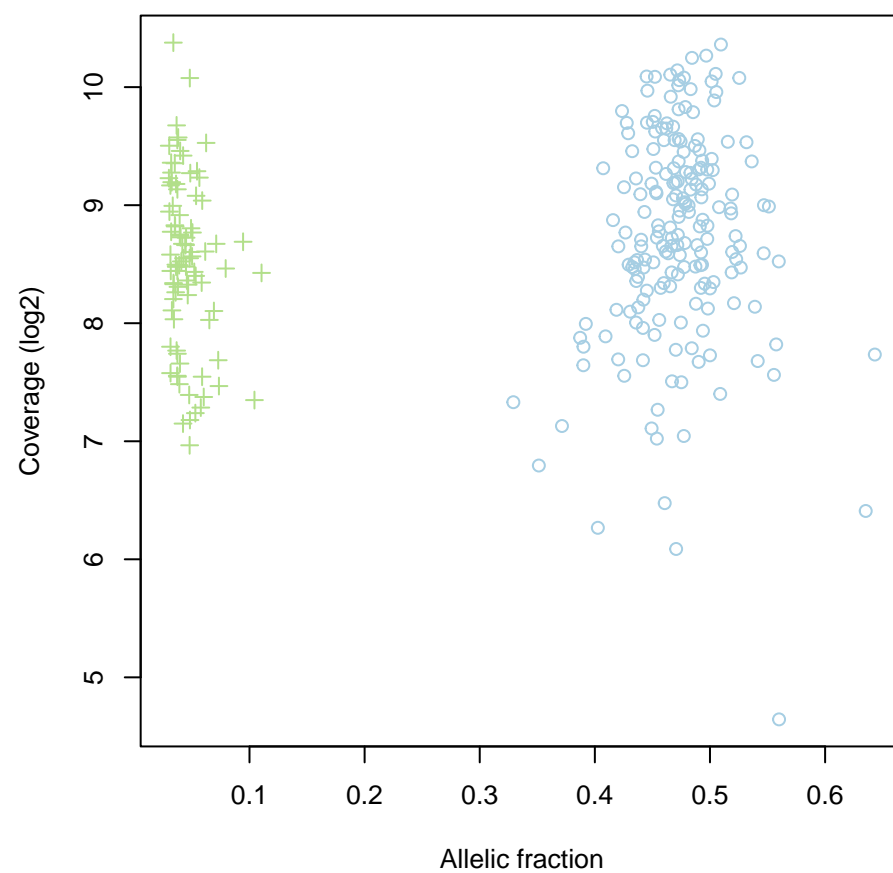
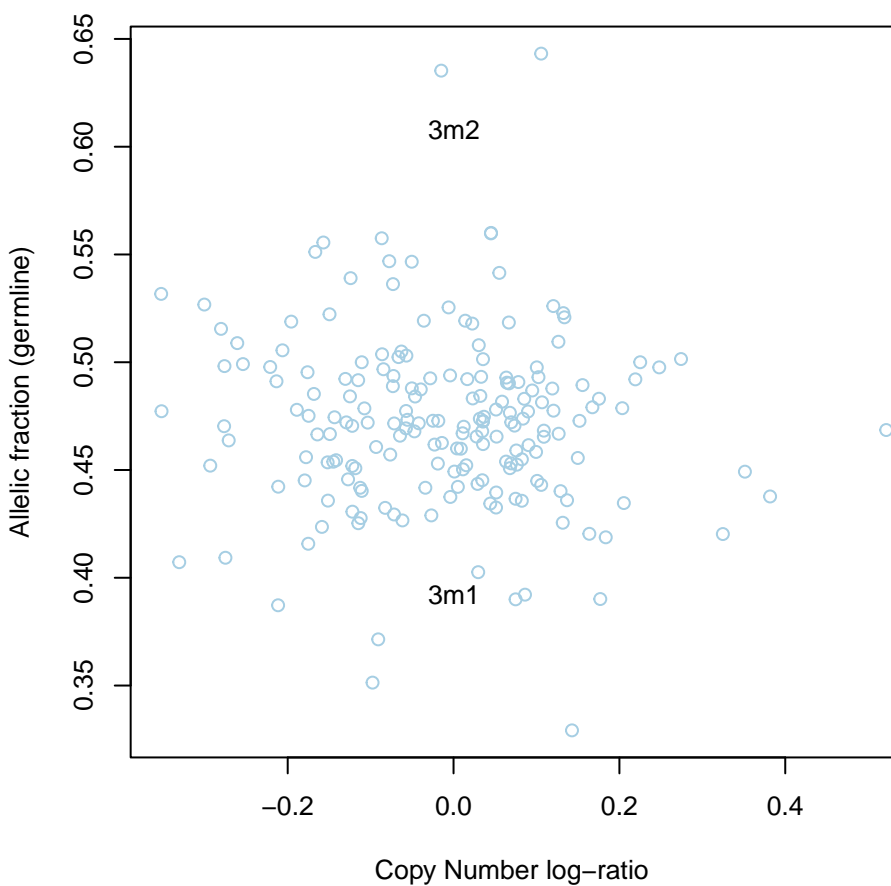


Purity: 0.55 Tumor ploidy: 2.999 SNV log-likelihood: -725.47 GoF: 66.2% Mean coverage: 476,548

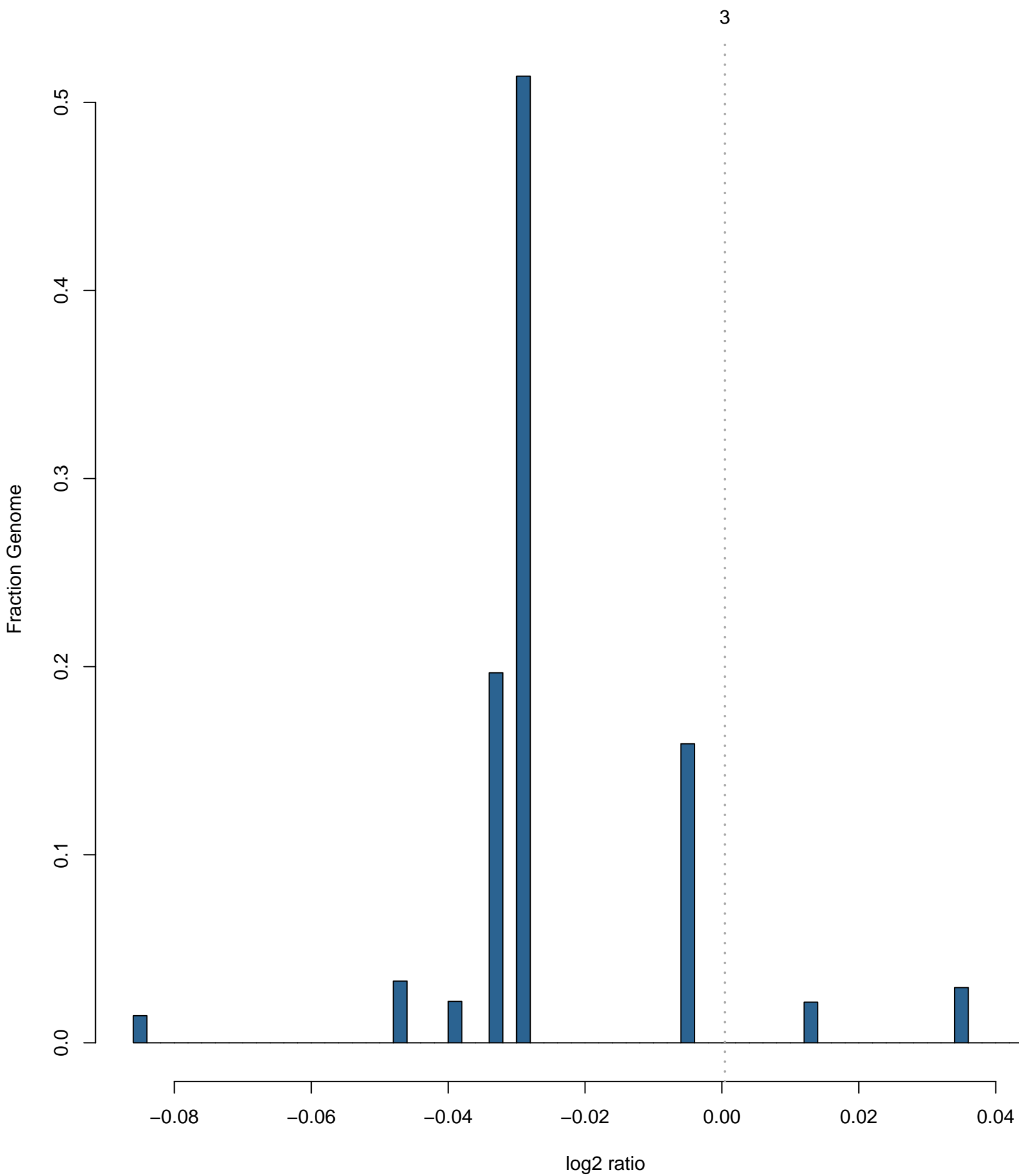


SCNA-fit log-likelihood: -14296.54

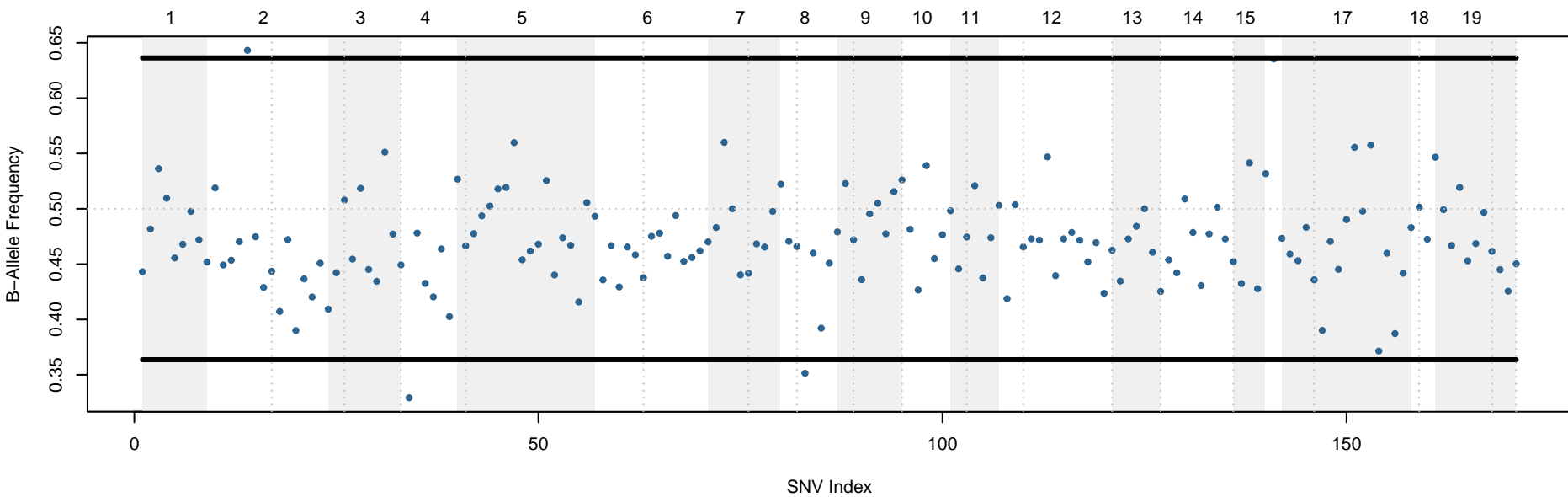




Purity: 0.75 Tumor ploidy: 2.999



Purity: 0.75 Tumor ploidy: 2.999 SNV log-likelihood: -1501.25 GoF: 42.4% Mean coverage: 476;548



SCNA-fit log-likelihood: -14293.42

