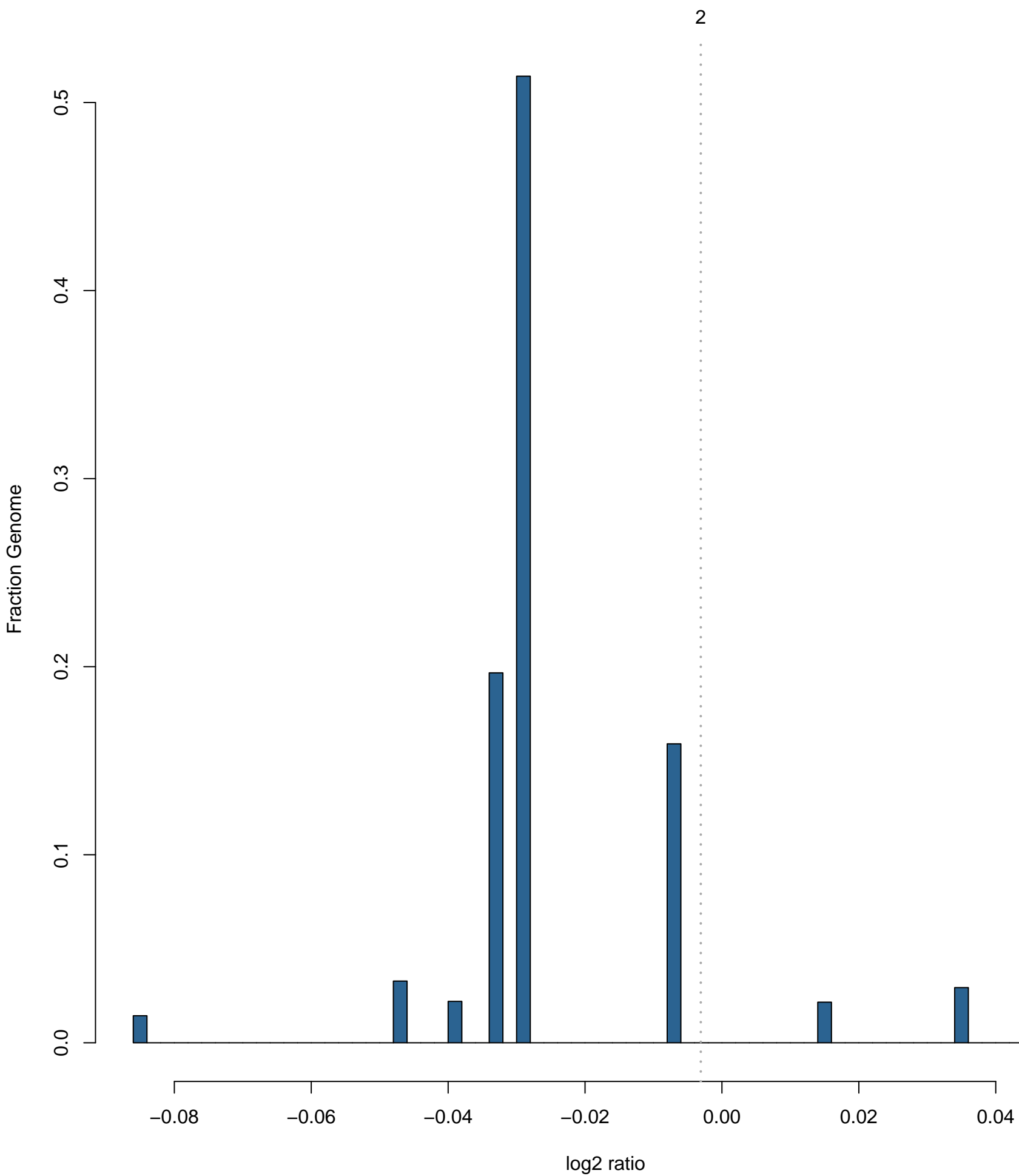
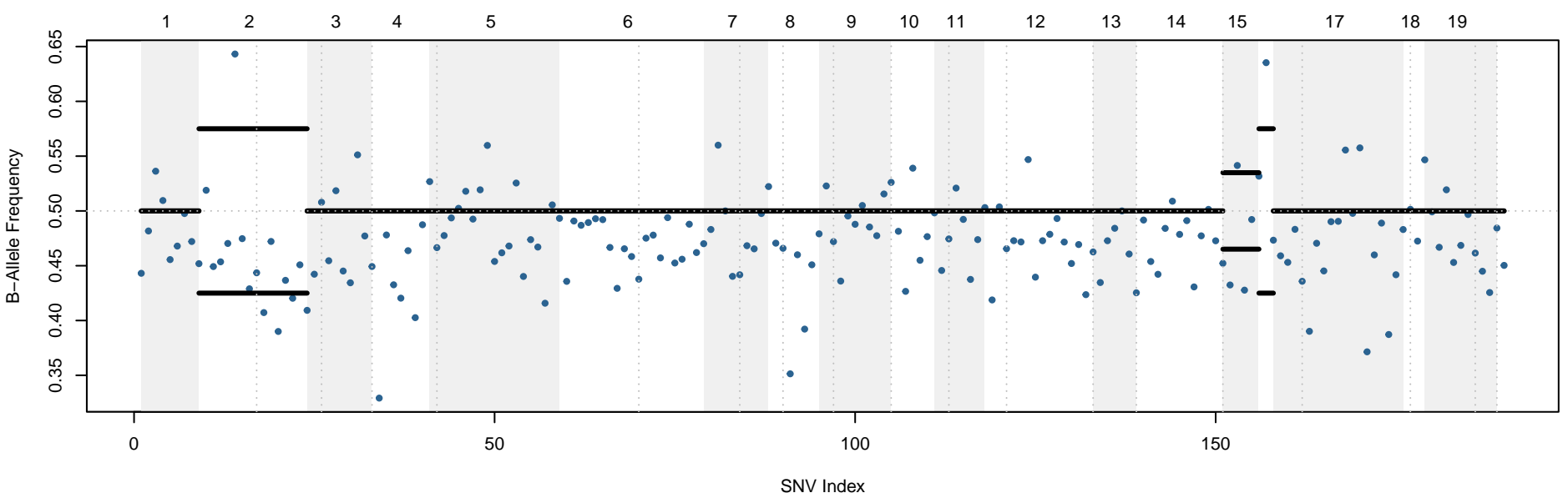


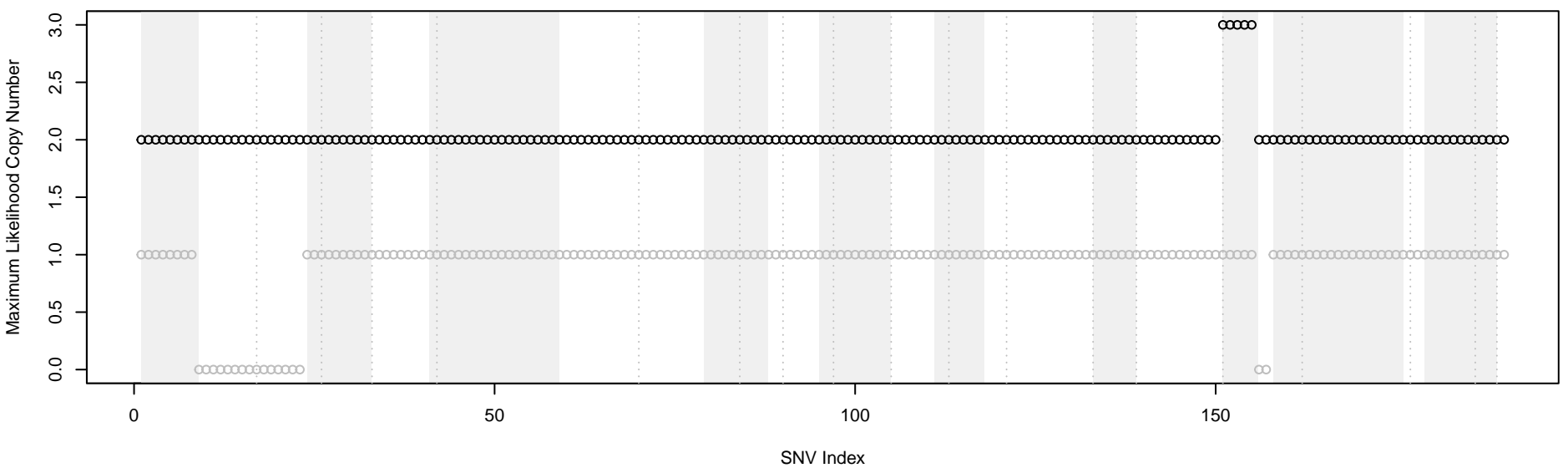
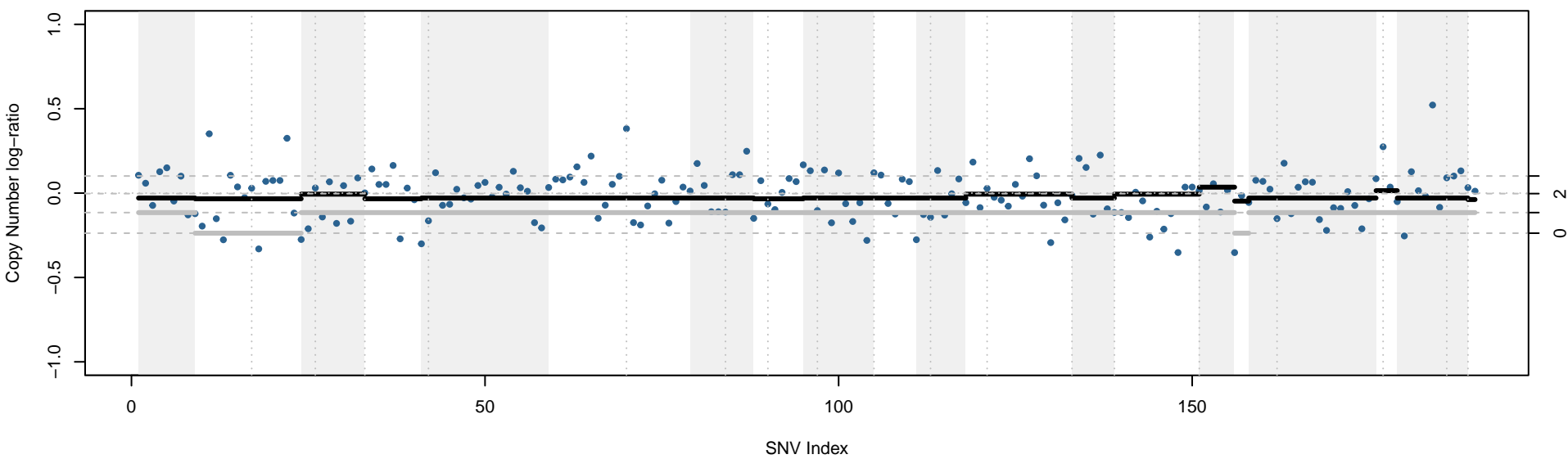
Purity: 0.15 Tumor ploidy: 2.029

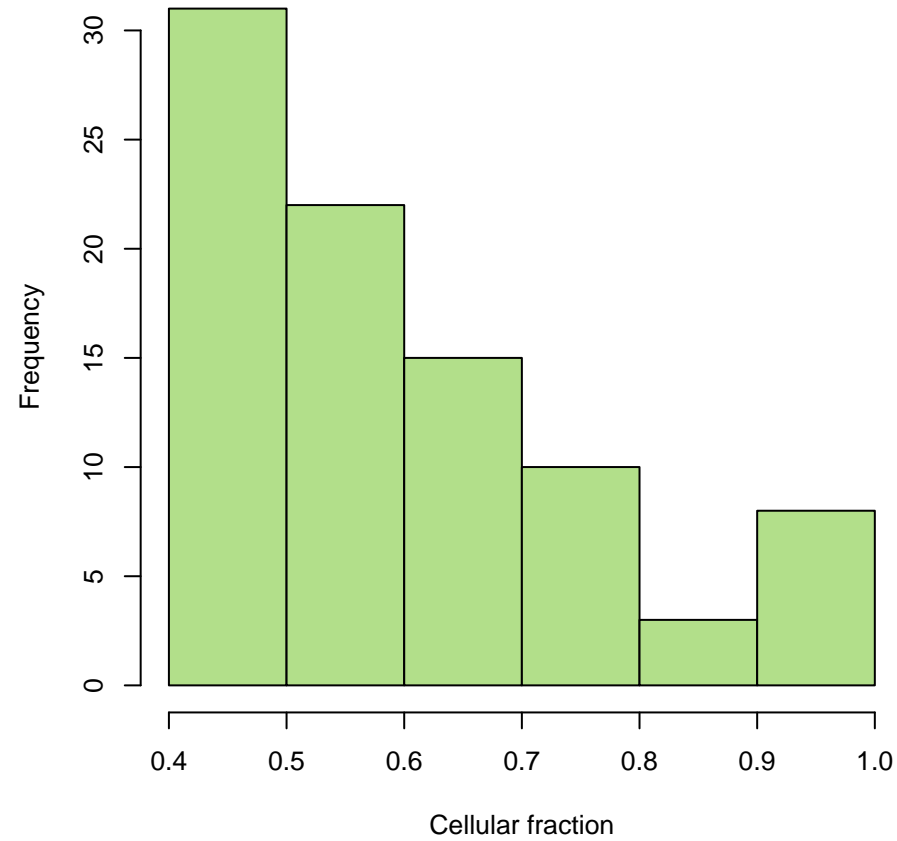
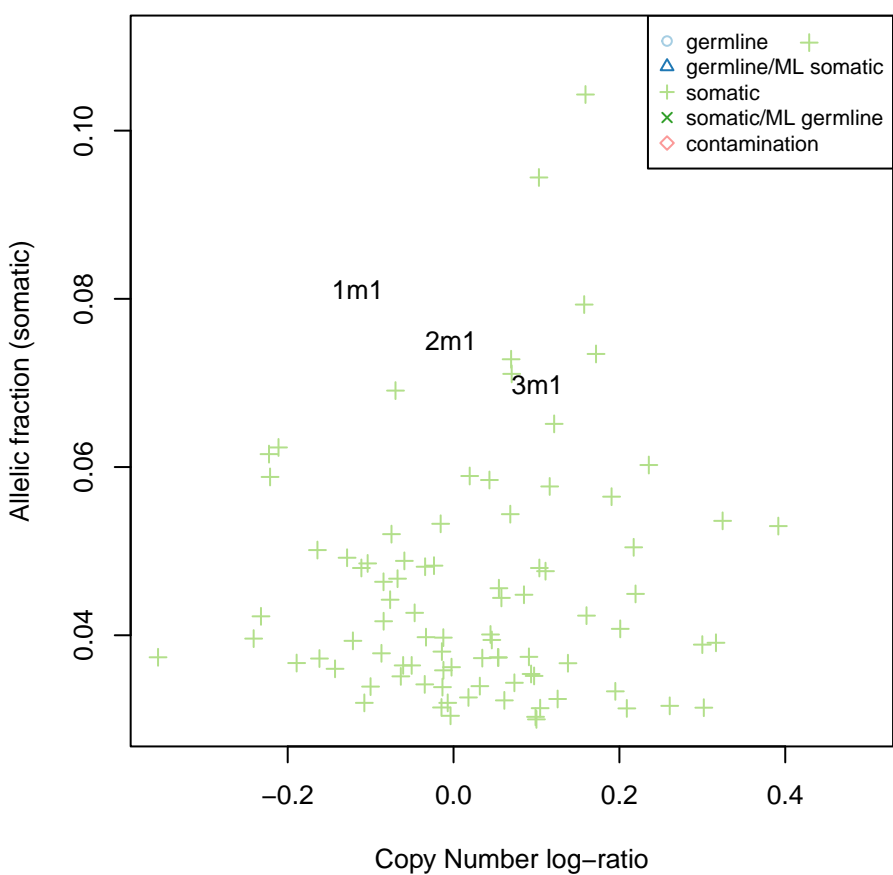
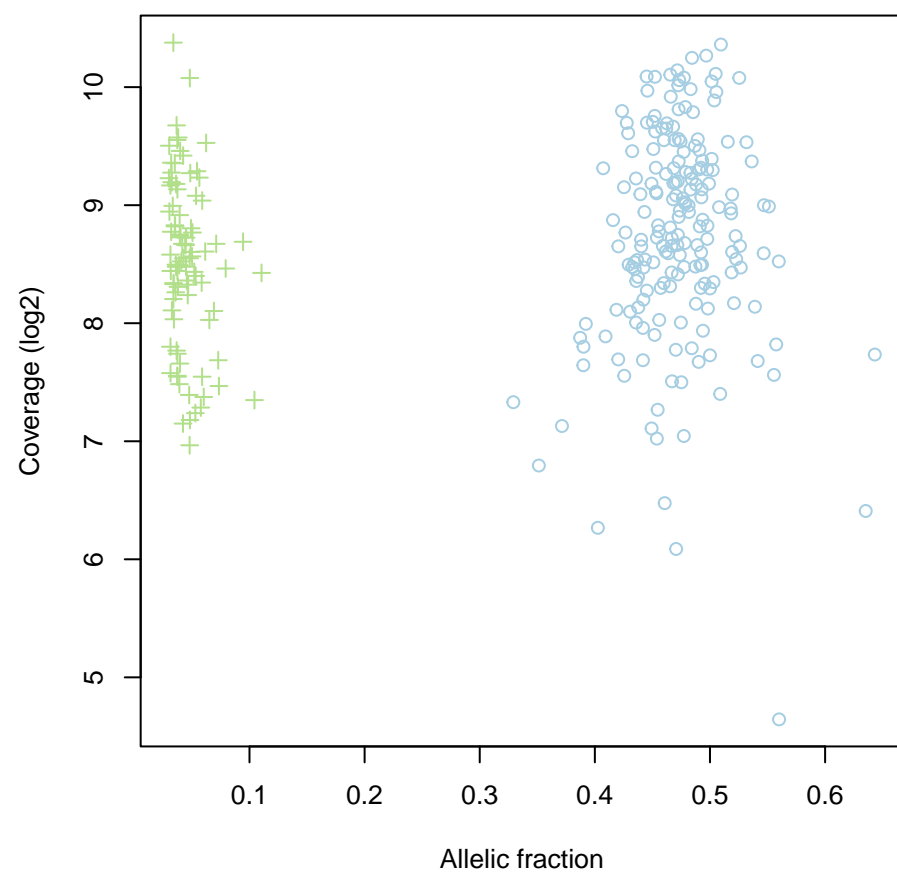
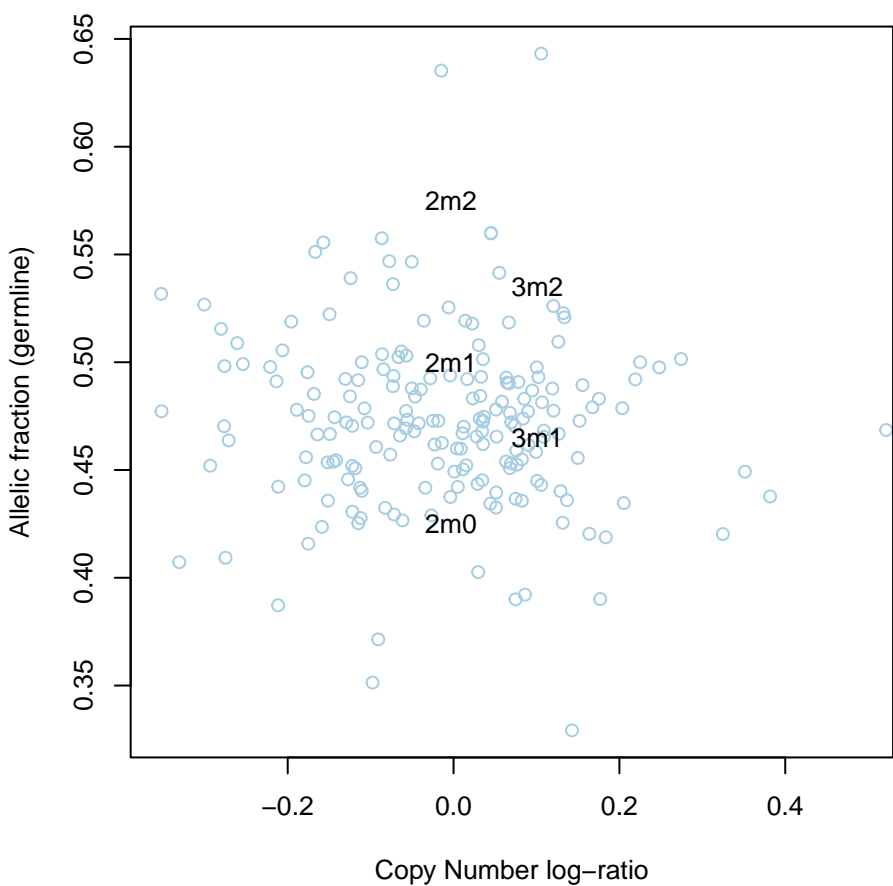


Purity: 0.15 Tumor ploidy: 2.029 SNV log-likelihood: 173.71 GoF: 96.3% Mean coverage: 476;548

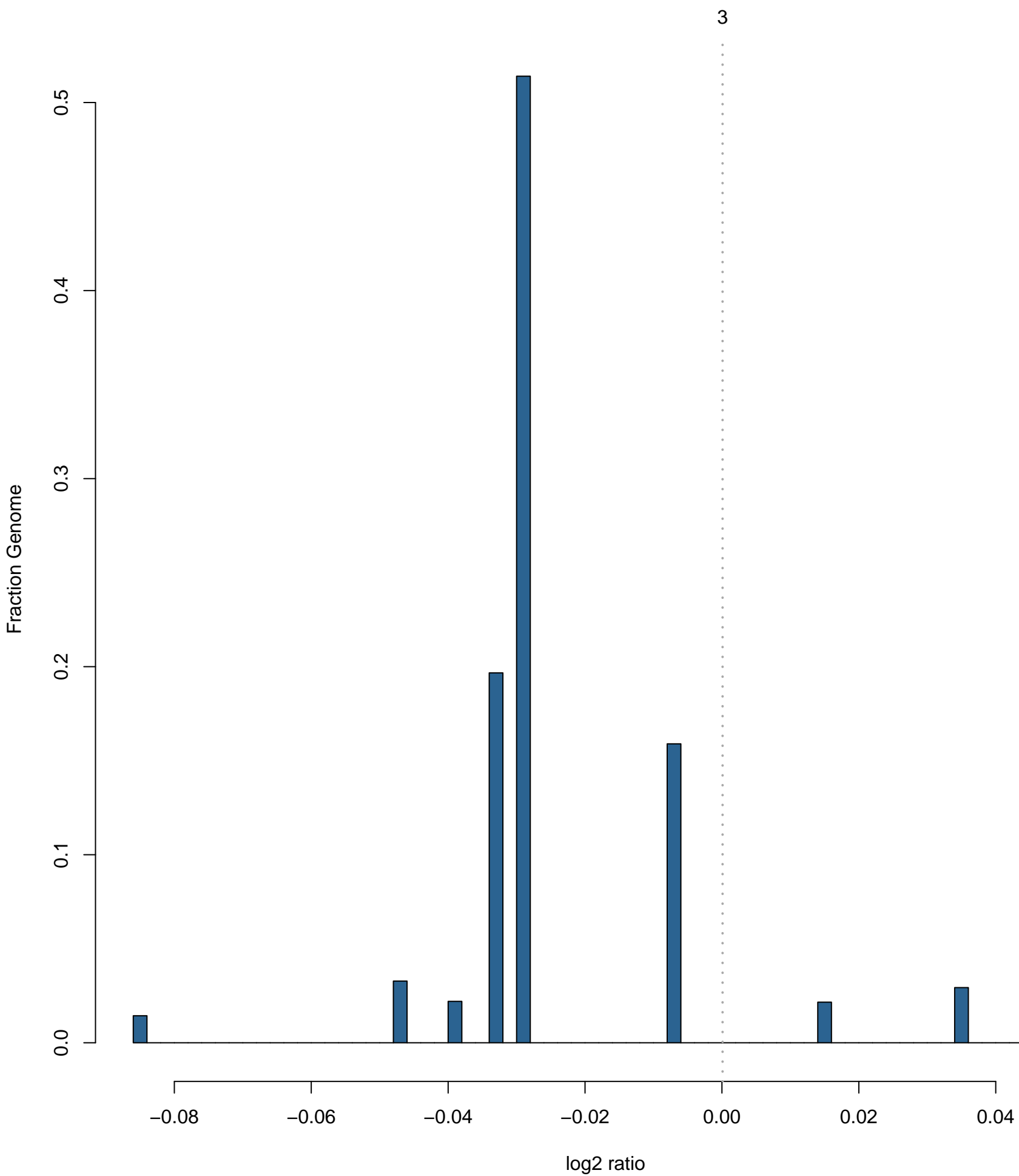


SCNA-fit log-likelihood: -14288.96

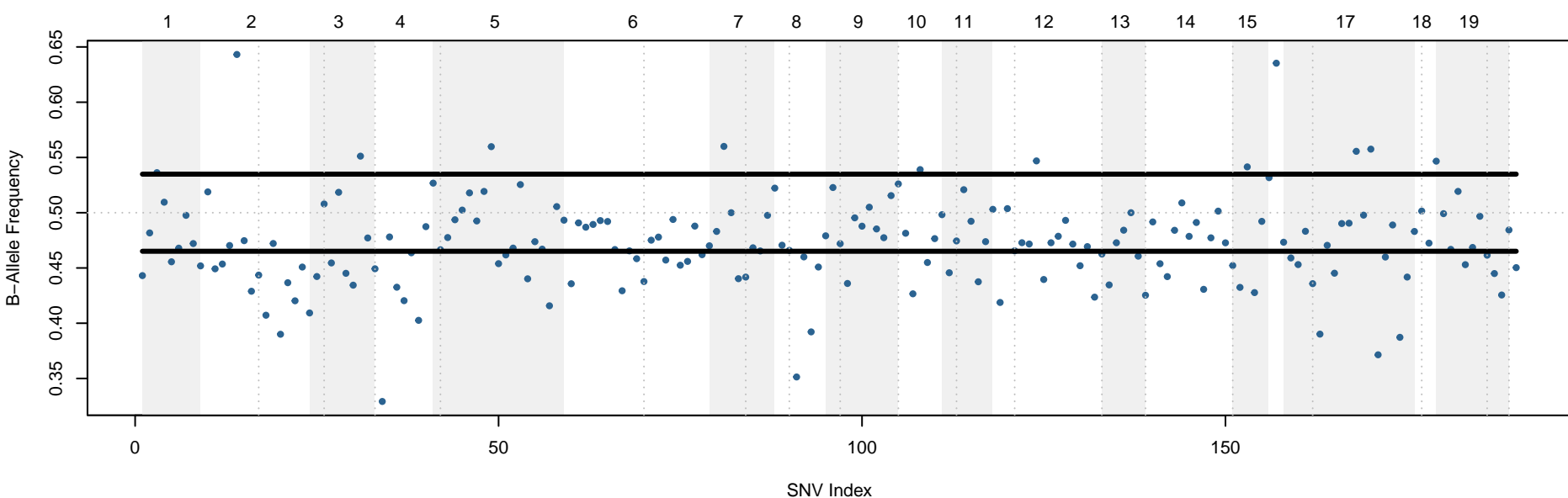




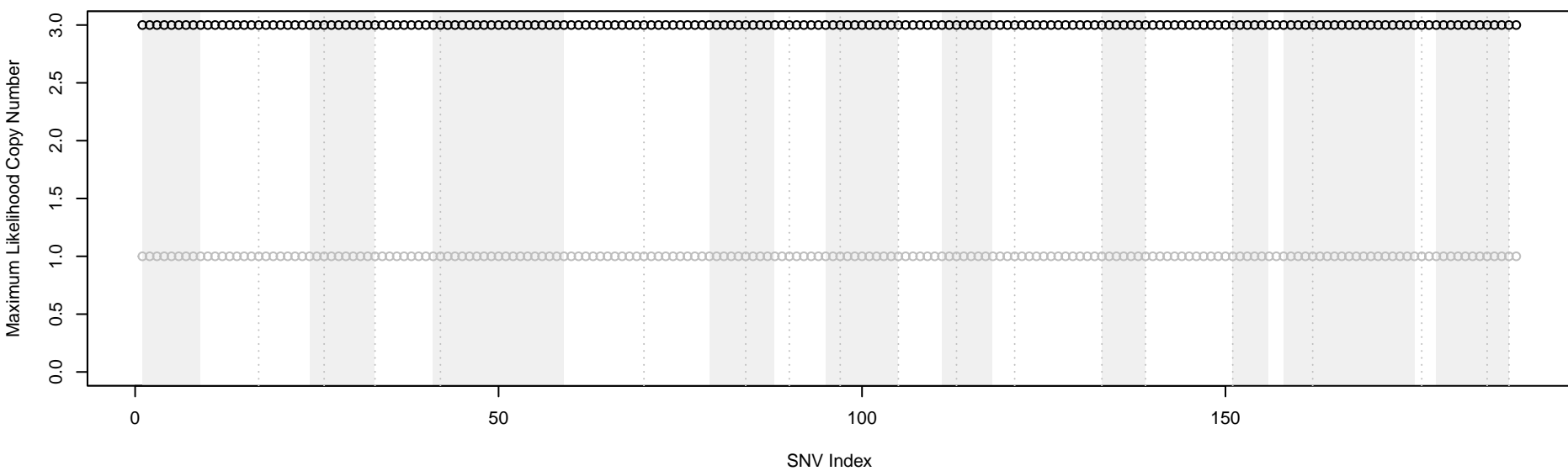
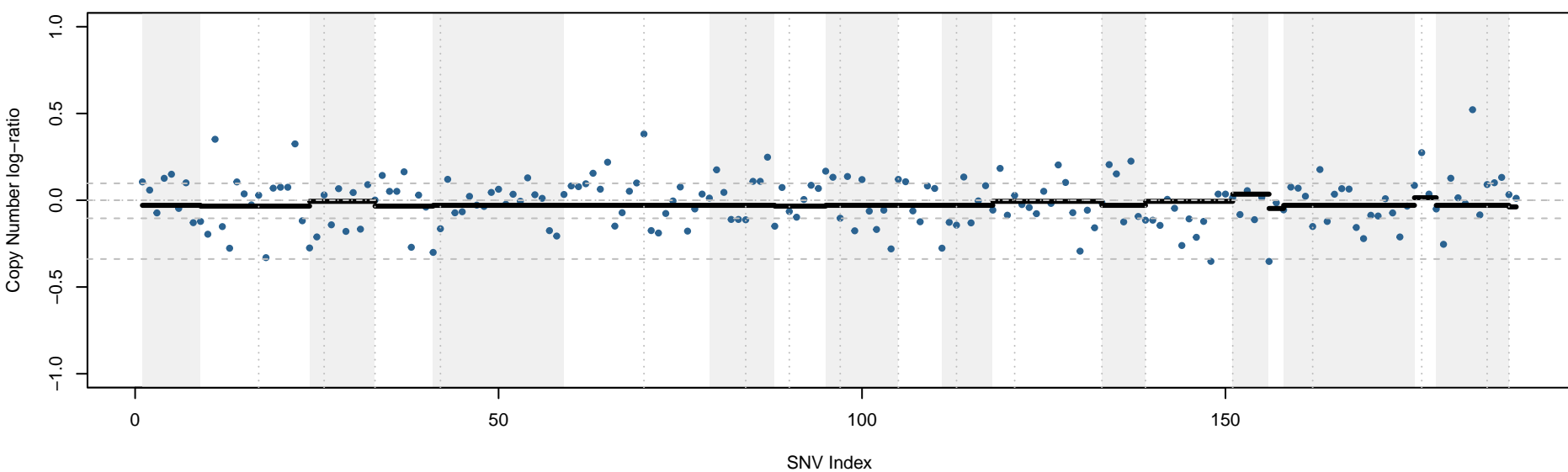
Purity: 0.15 Tumor ploidy: 2.999

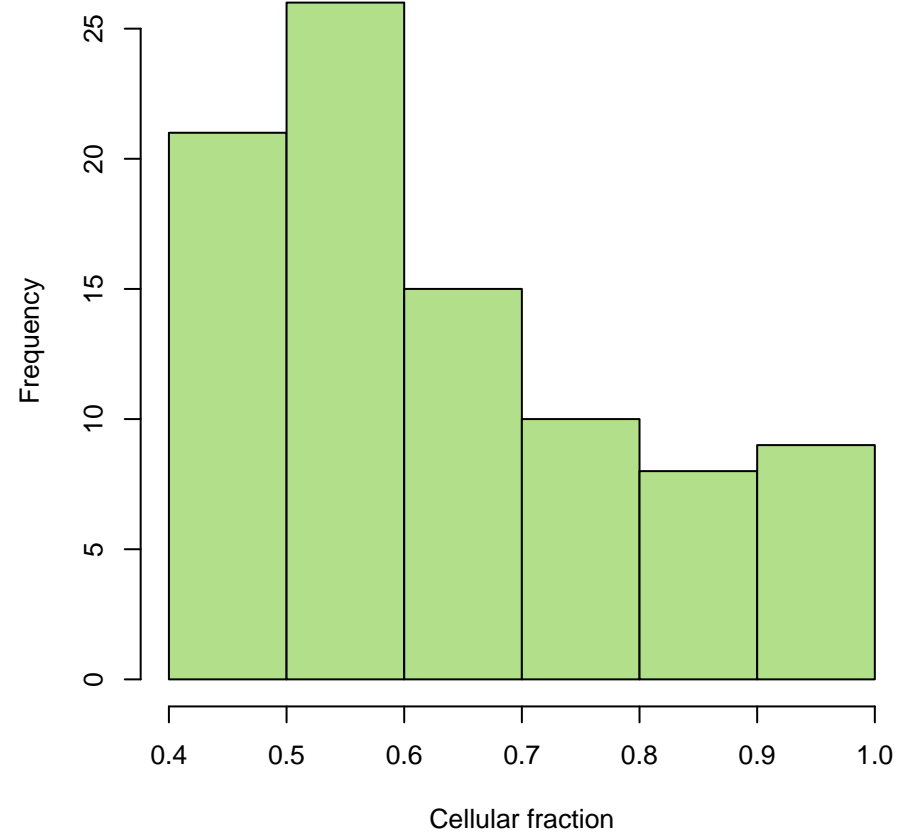
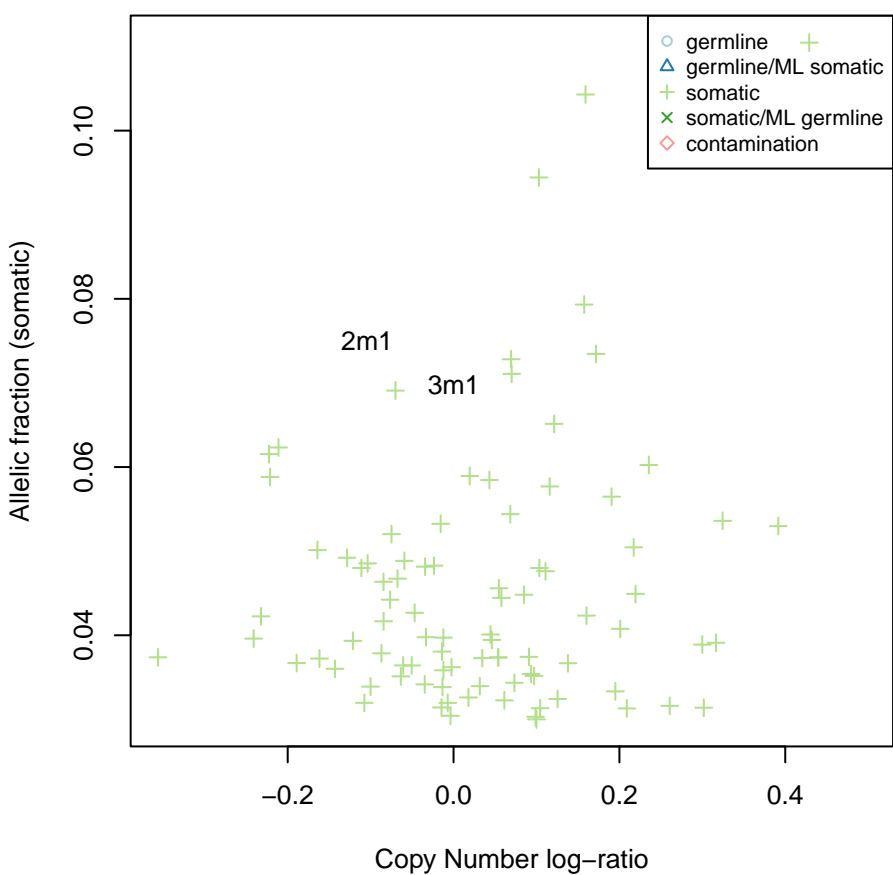
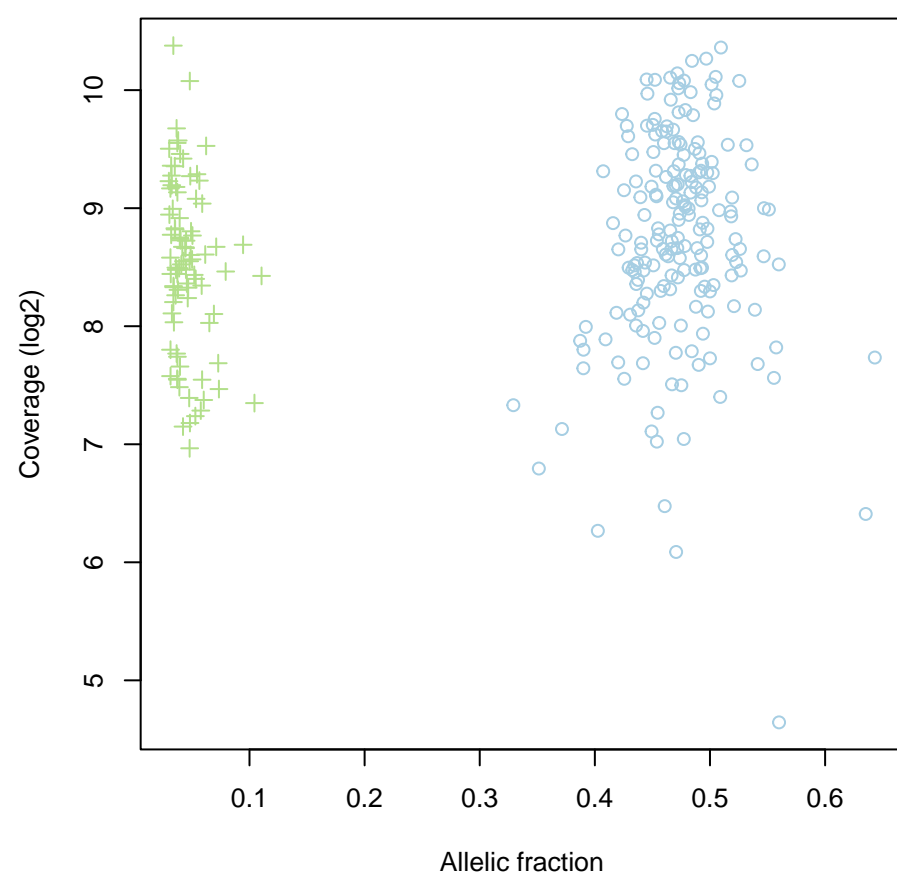
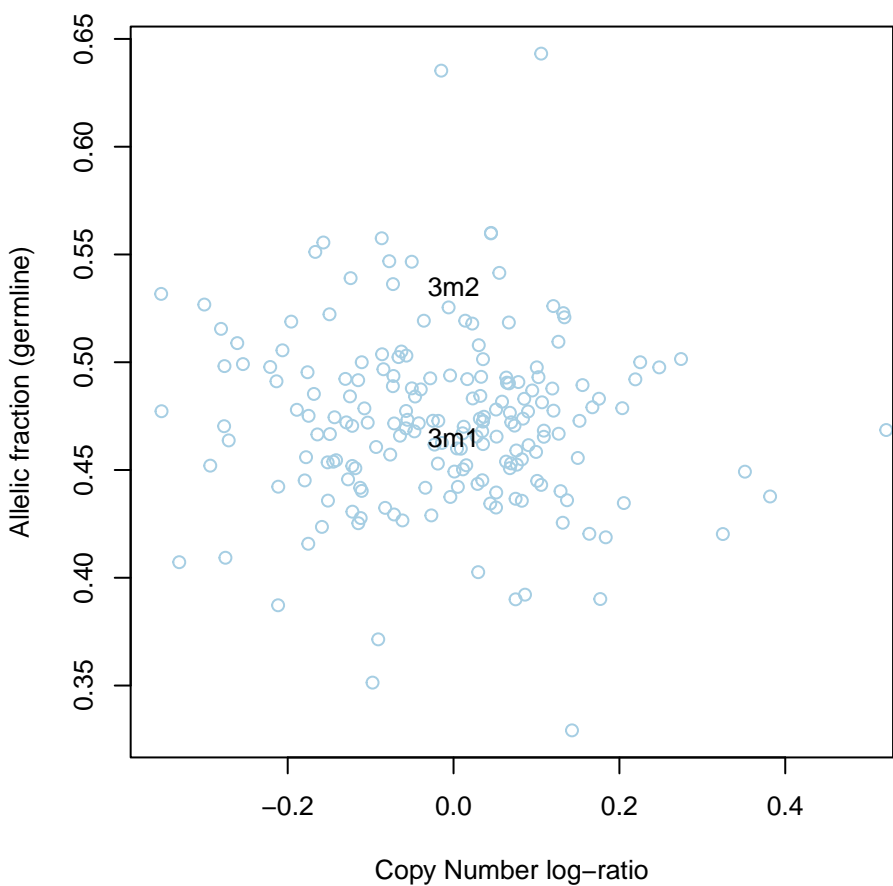


Purity: 0.15 Tumor ploidy: 2.999 SNV log-likelihood: 26.72 GoF: 97.9% Mean coverage: 476;548

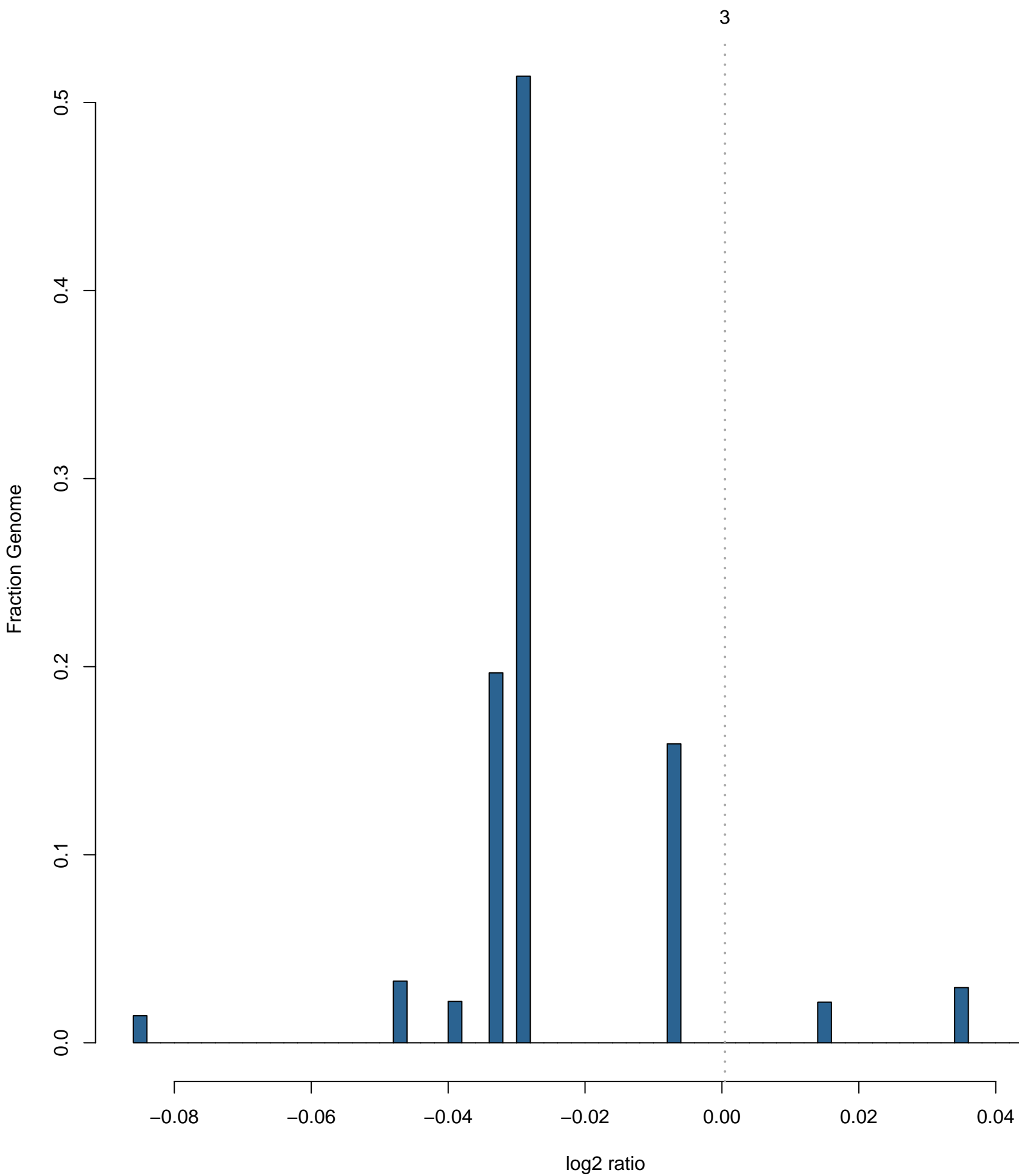


SCNA-fit log-likelihood: -14302.94

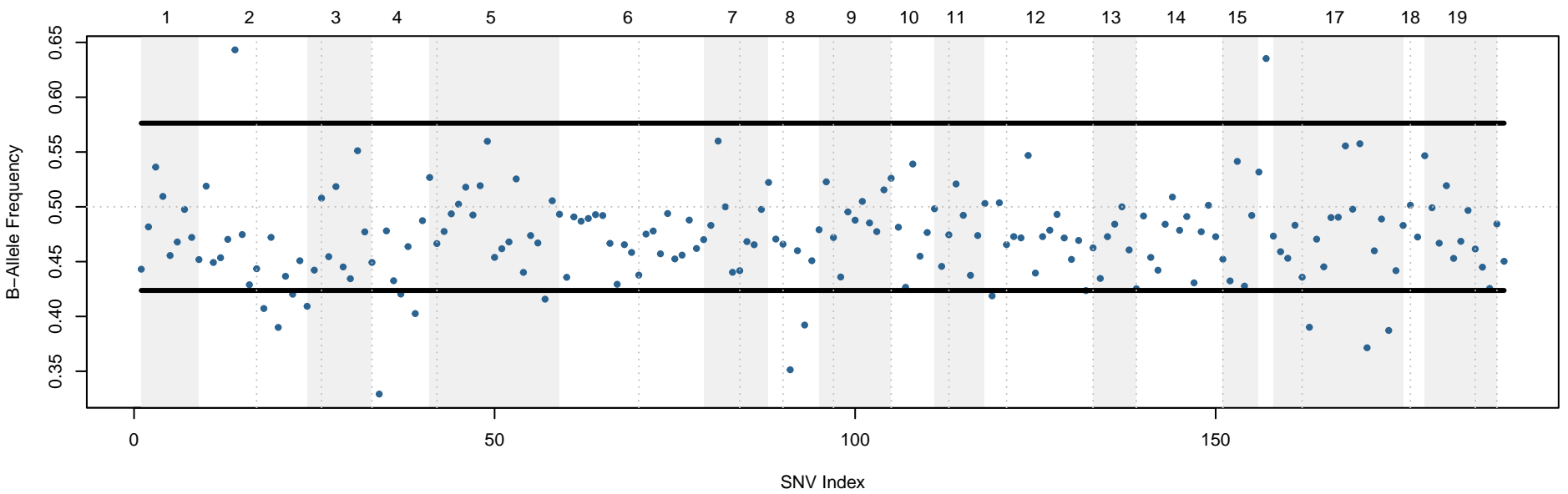




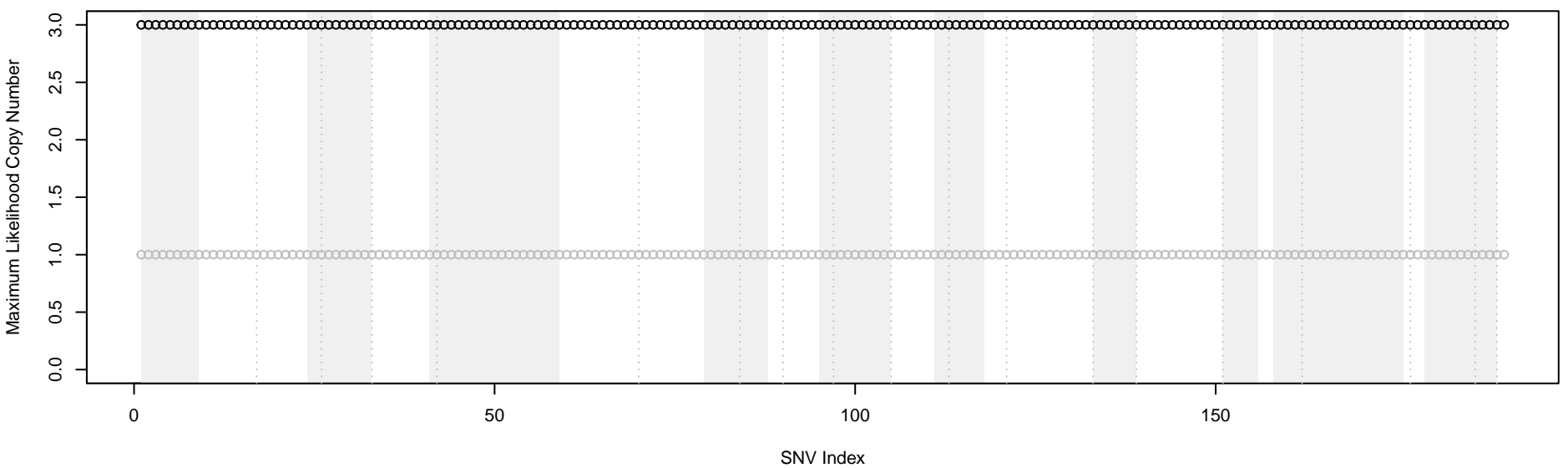
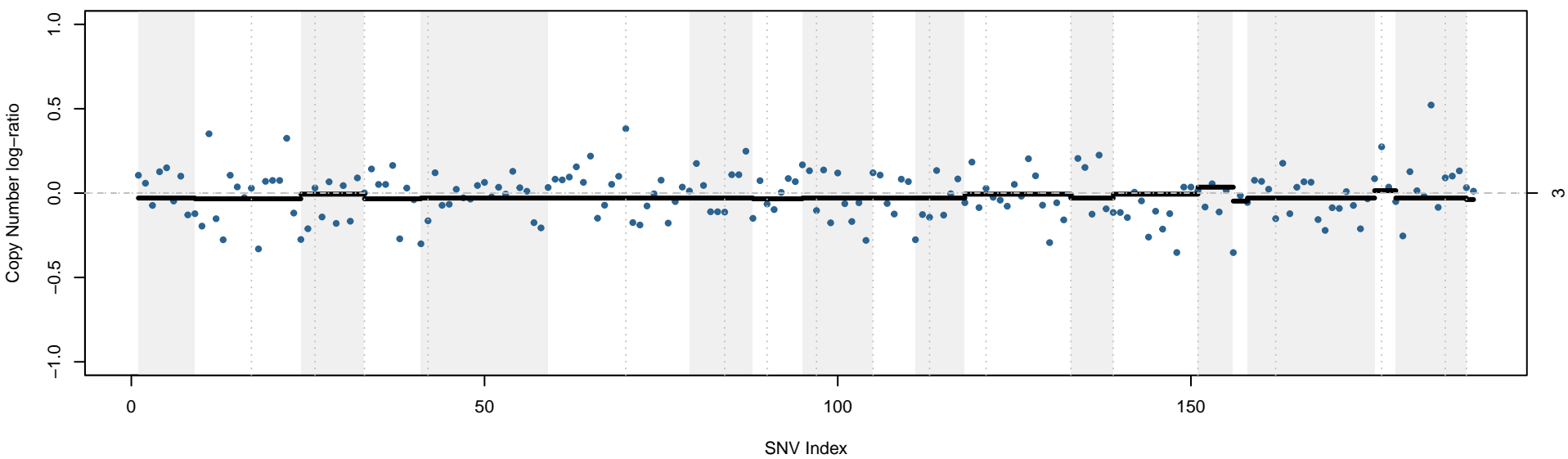
Purity: 0.36 Tumor ploidy: 2.998



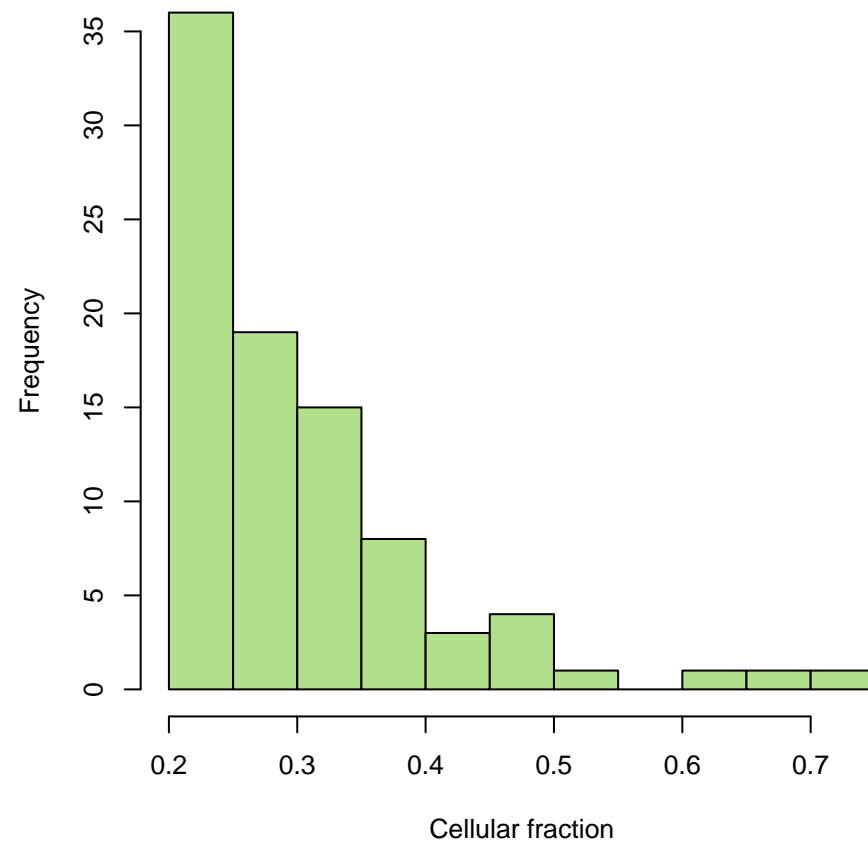
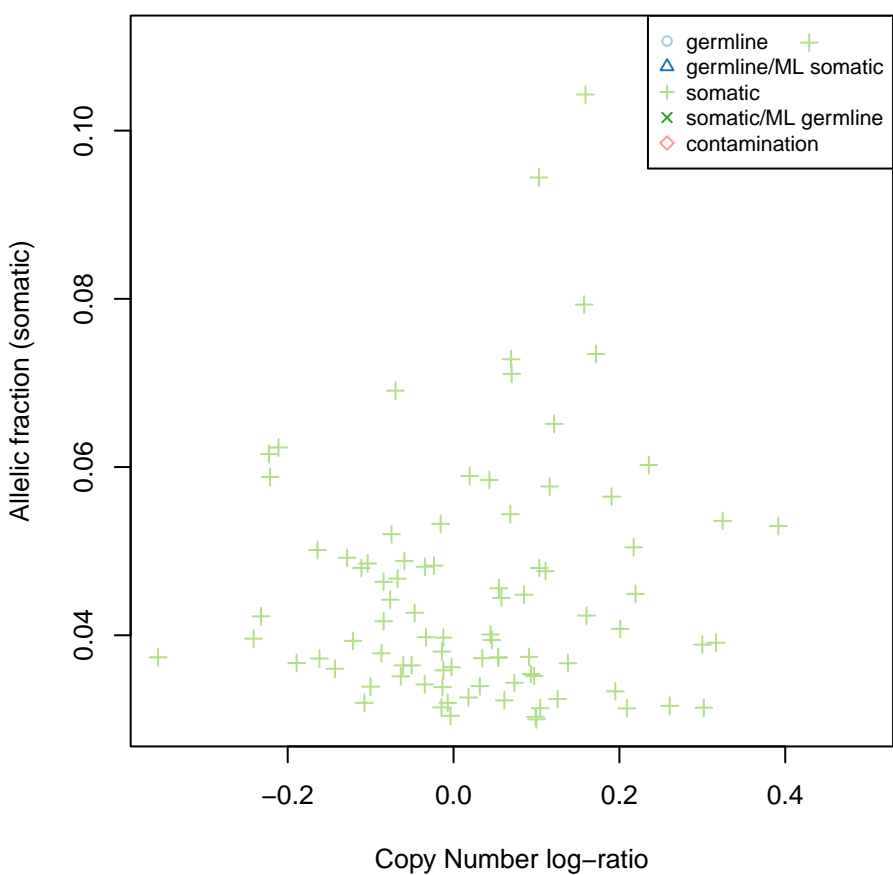
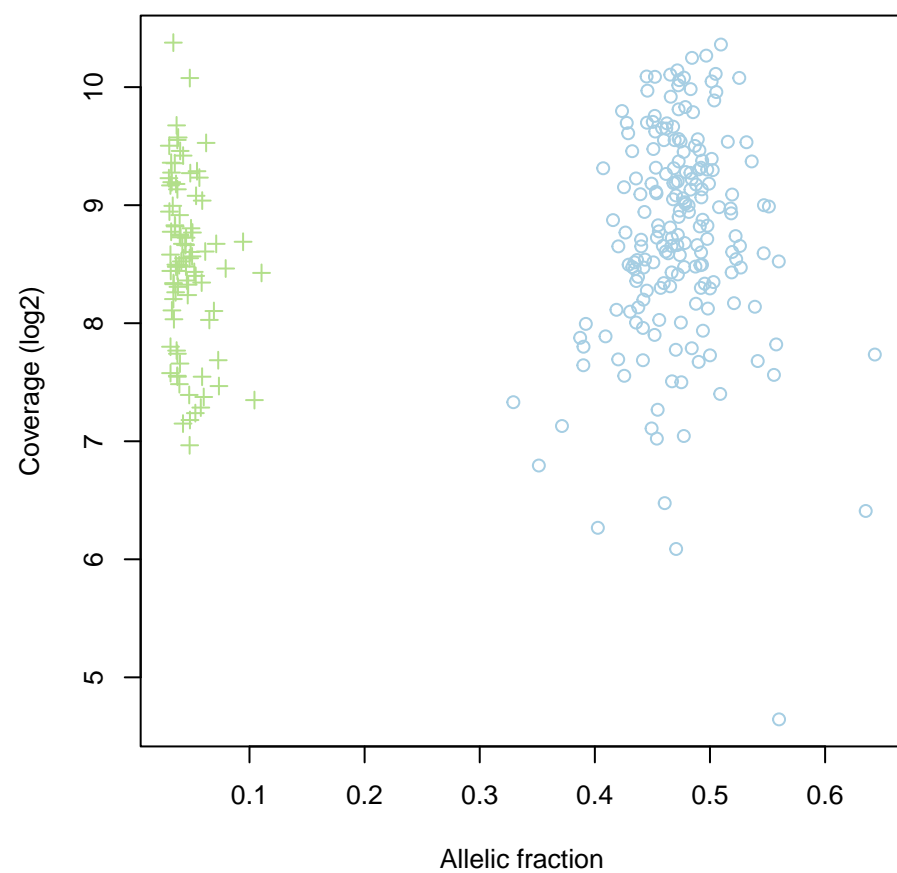
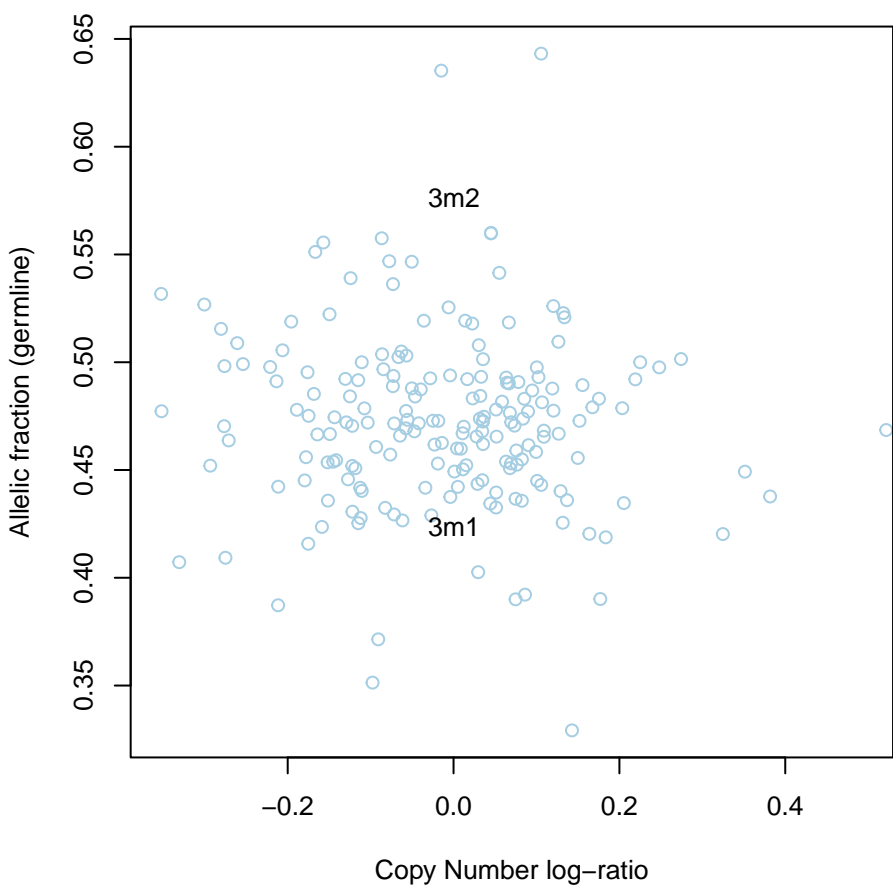
Purity: 0.36 Tumor ploidy: 2.998 SNV log-likelihood: -174.84 GoF: 86.3% Mean coverage: 476,548



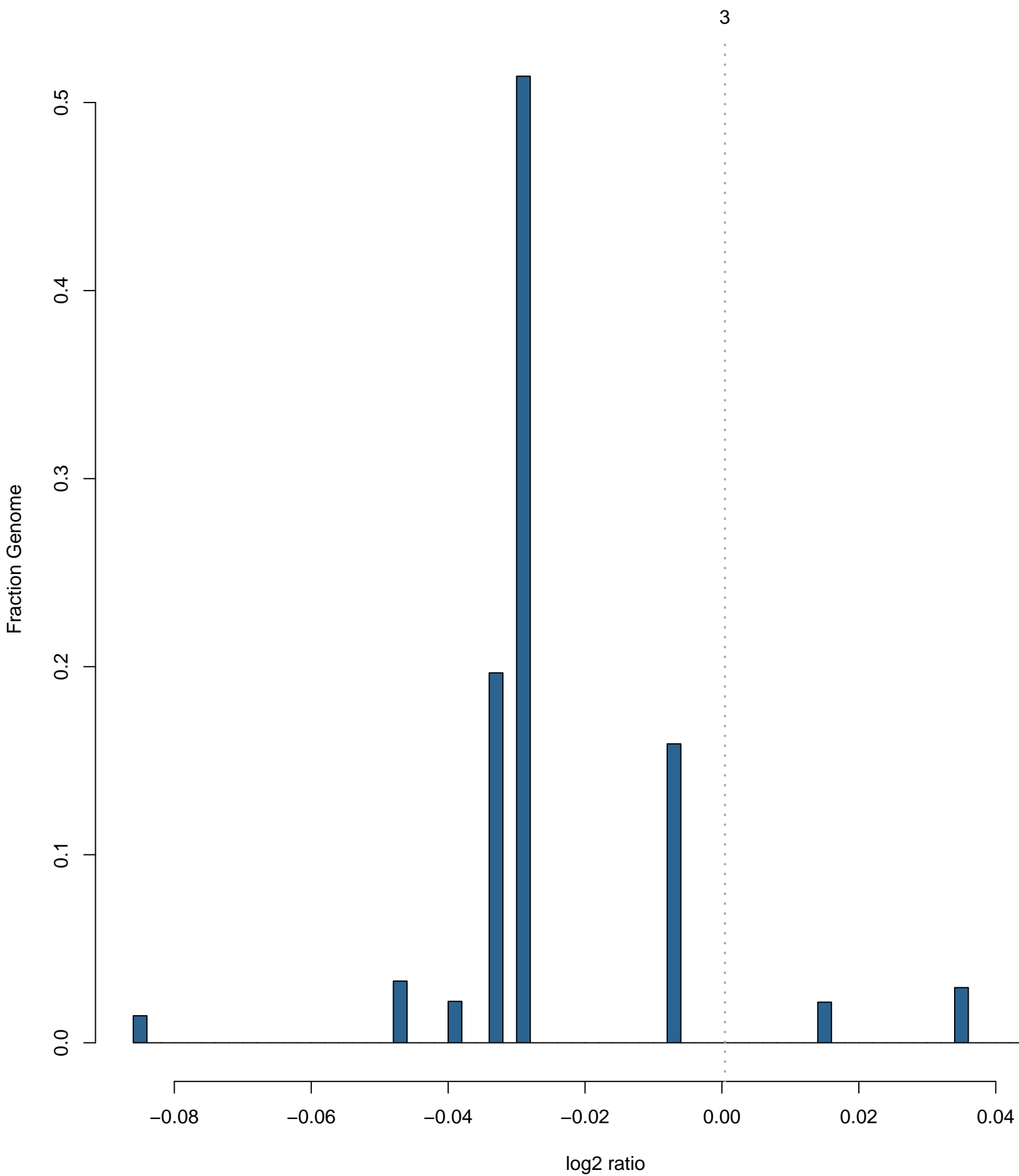
SCNA-fit log-likelihood: -14301.3



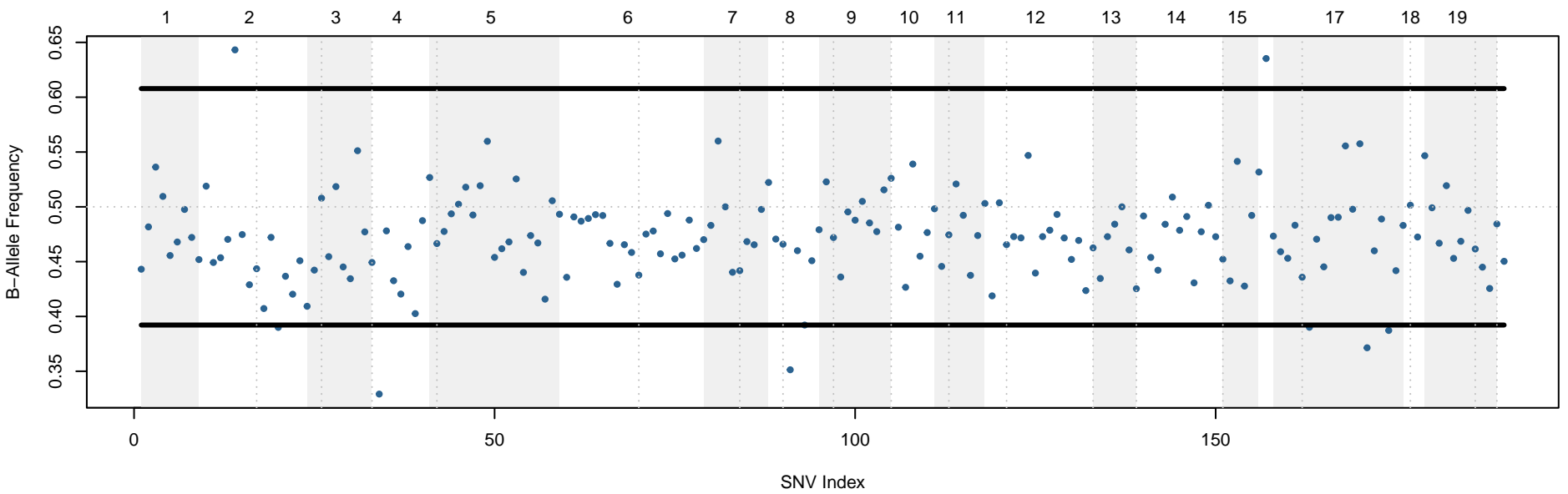




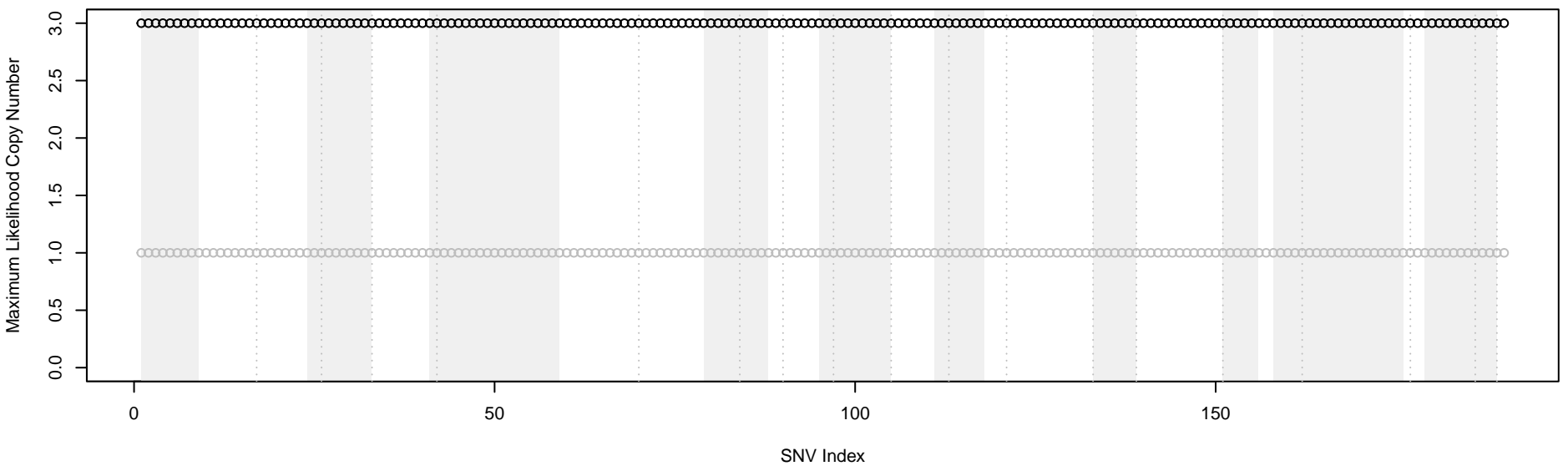
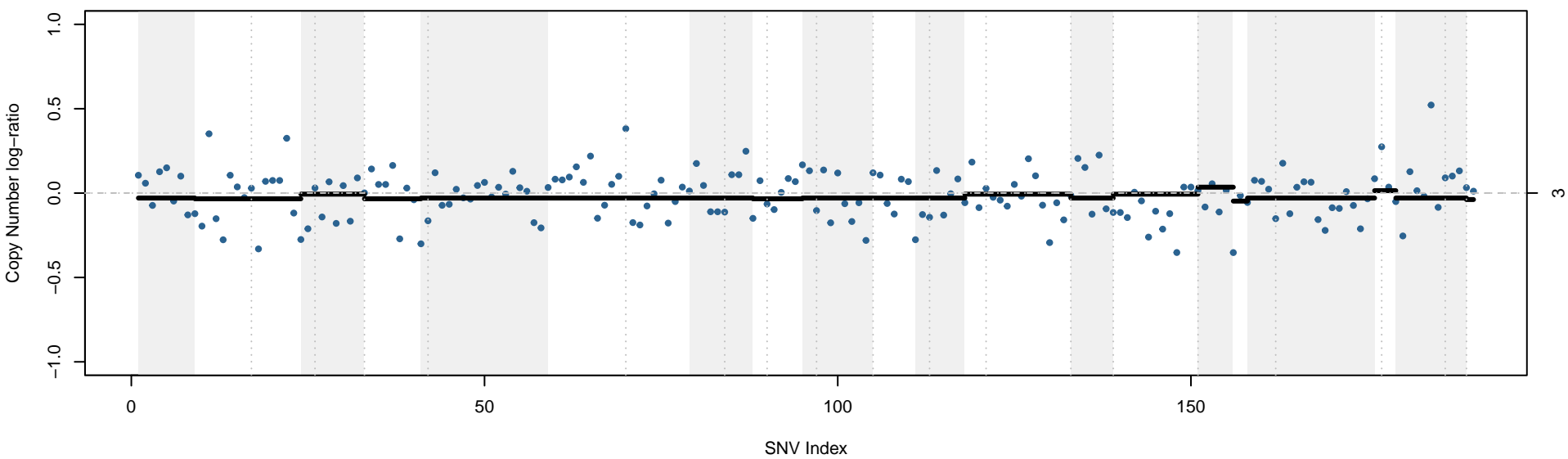
Purity: 0.55 Tumor ploidy: 2.999

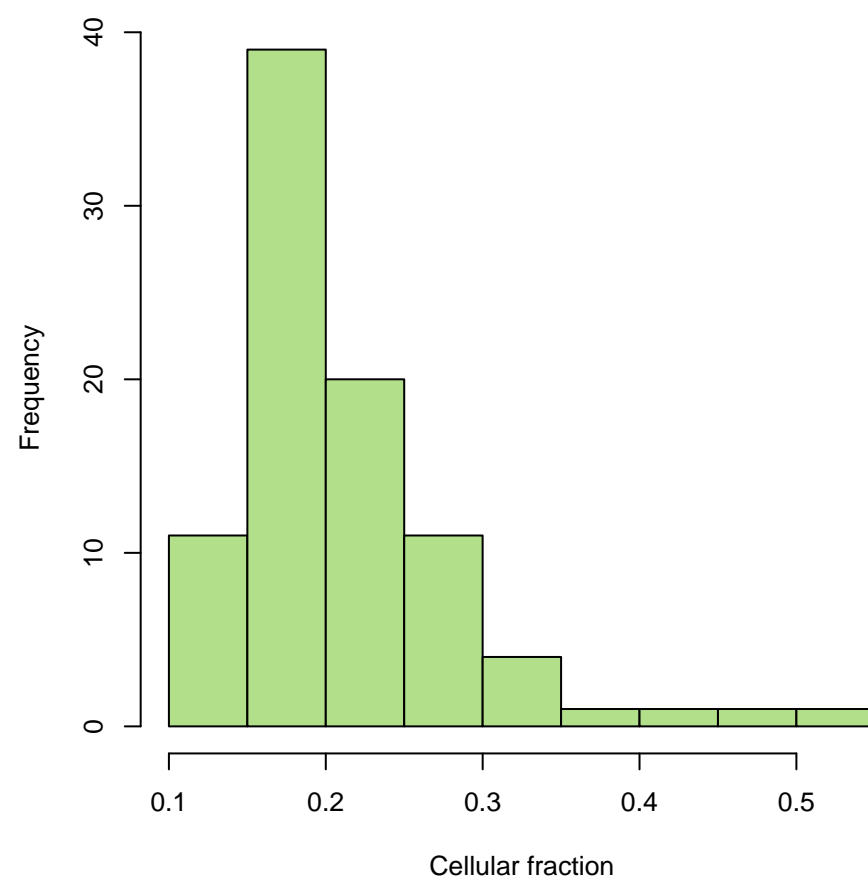
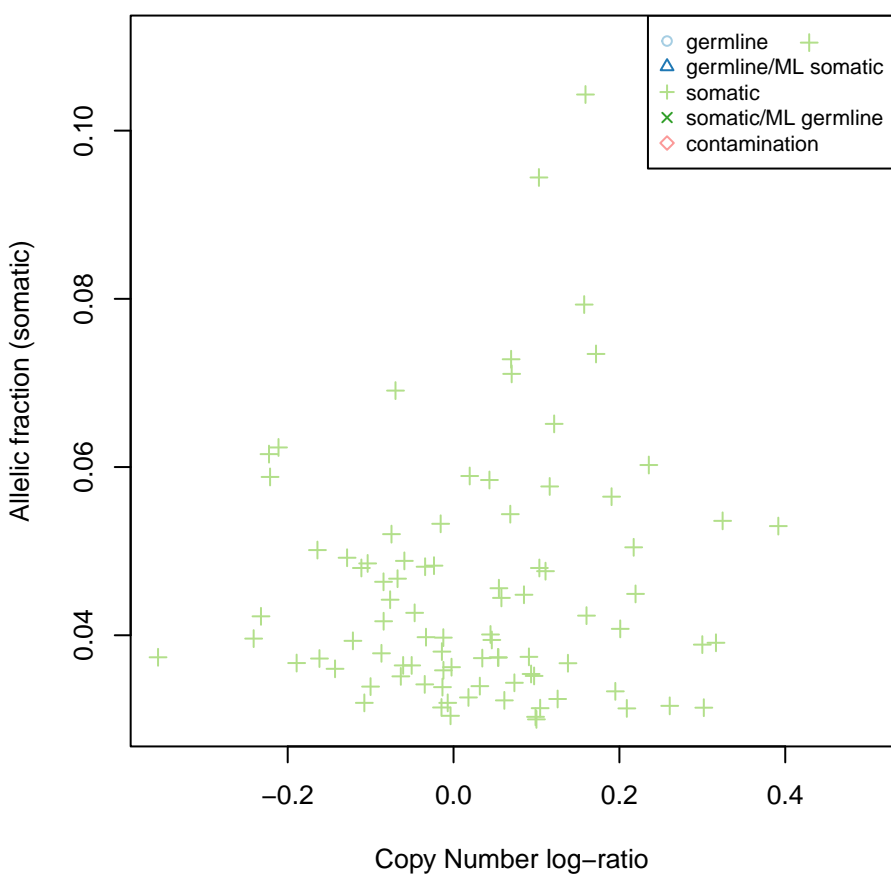
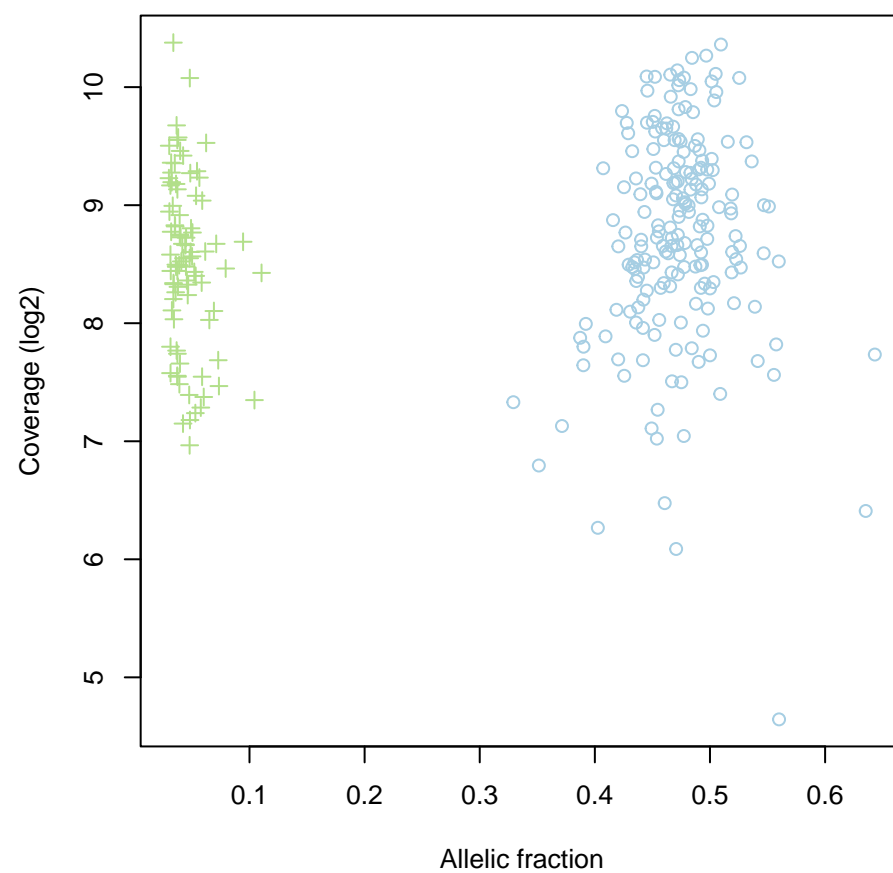
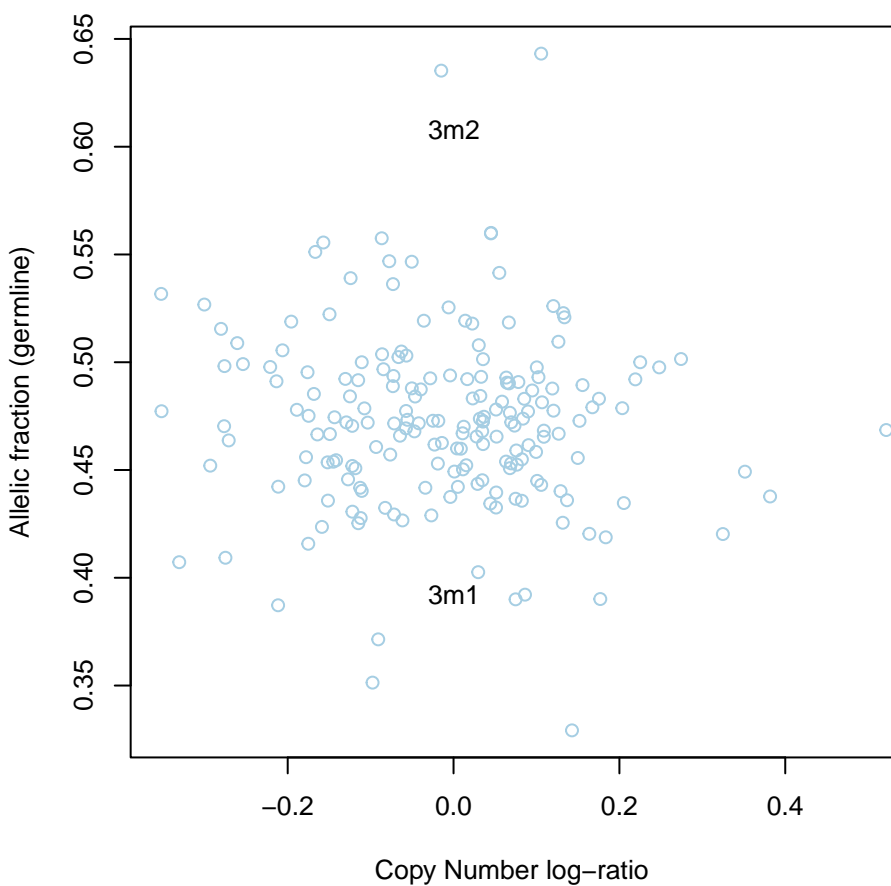


Purity: 0.55 Tumor ploidy: 2.999 SNV log-likelihood: -725.47 GoF: 66.2% Mean coverage: 476,548

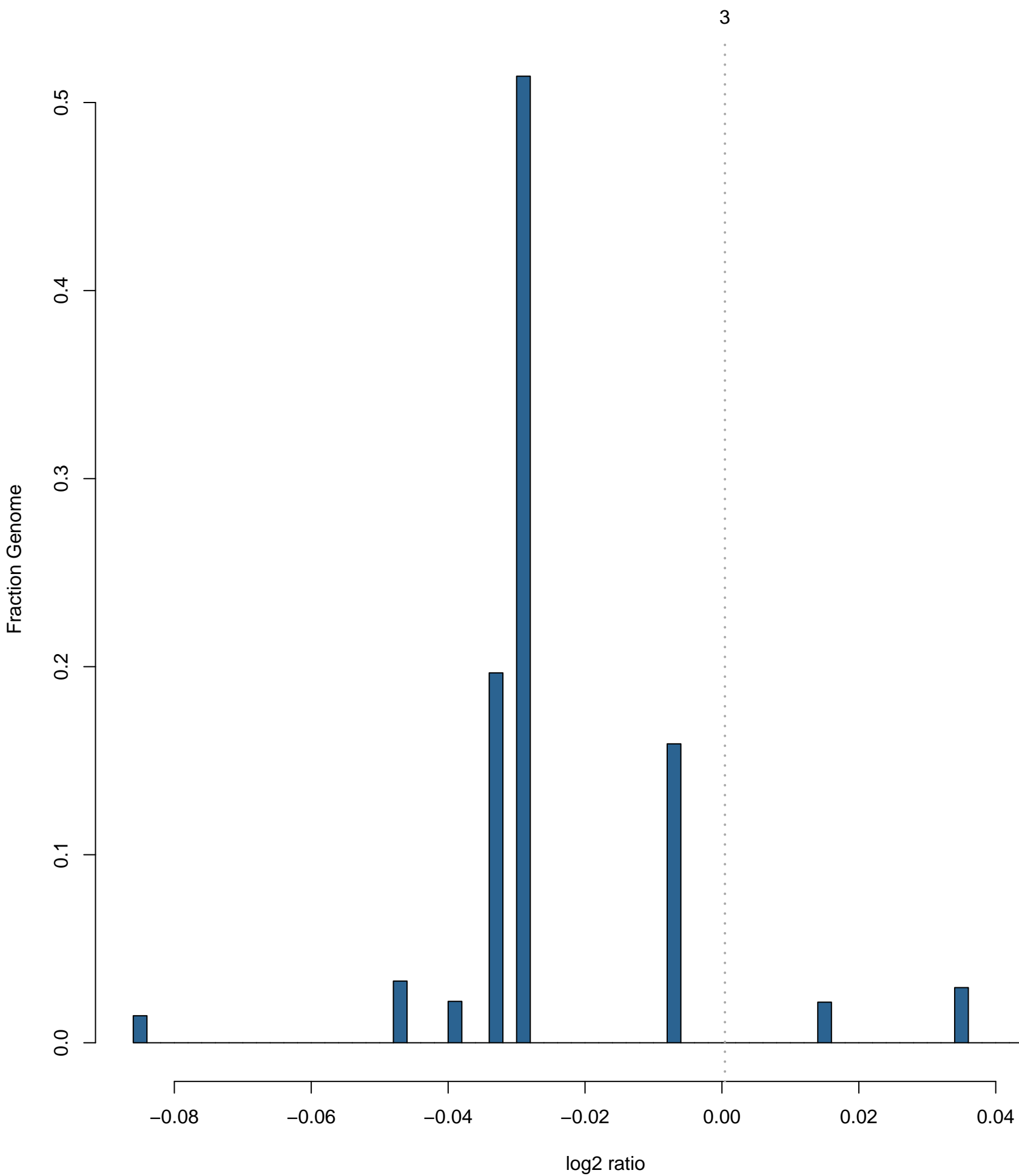


SCNA-fit log-likelihood: -14297.34

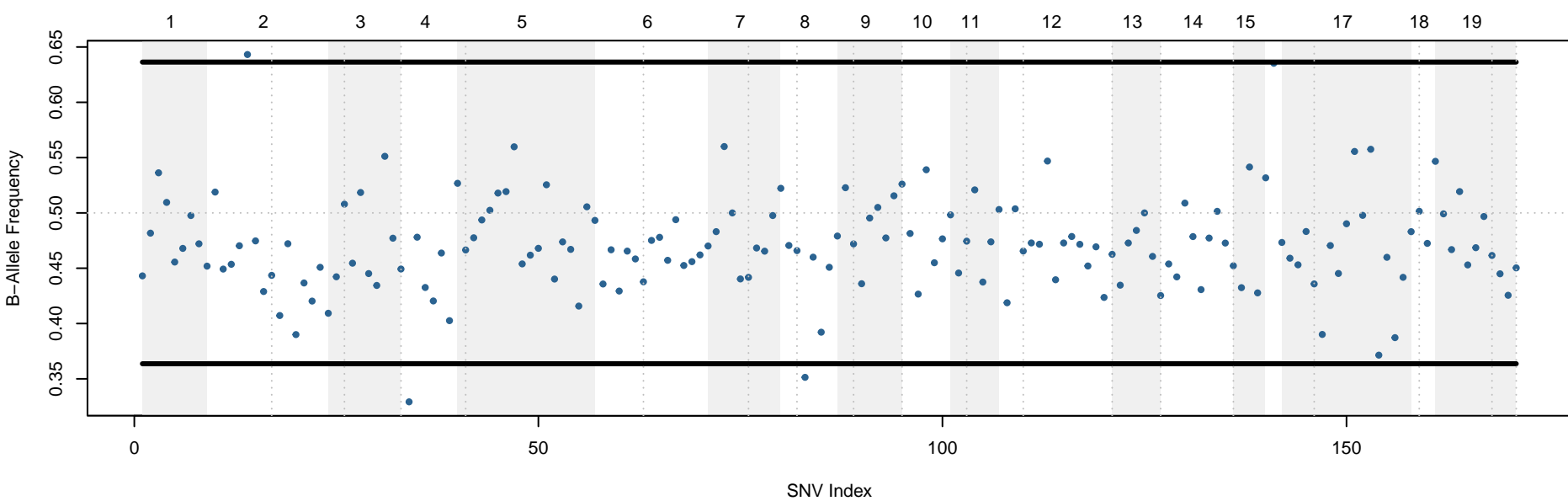




Purity: 0.75 Tumor ploidy: 2.999



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



SCNA-fit log-likelihood: -14294.22

