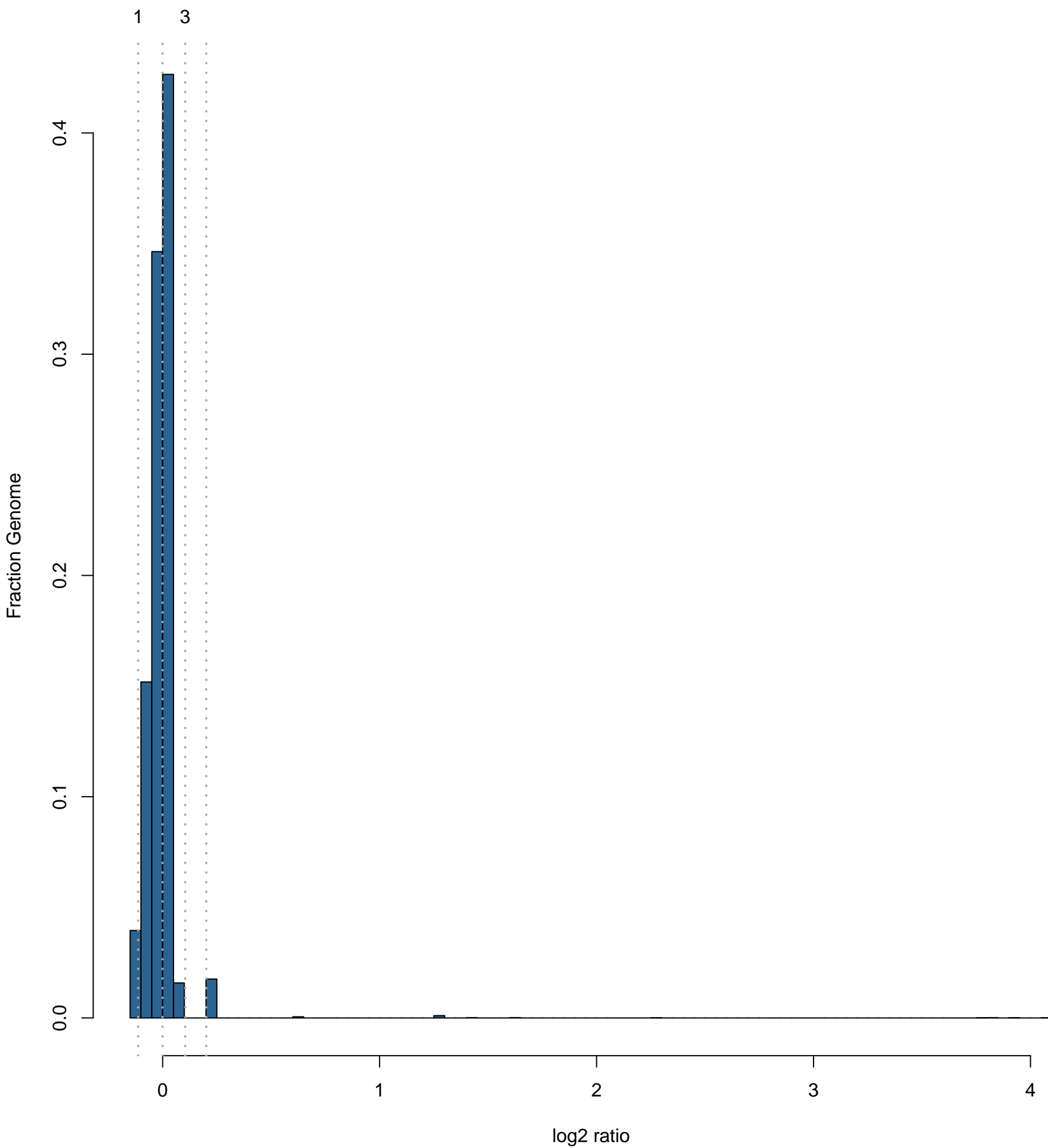
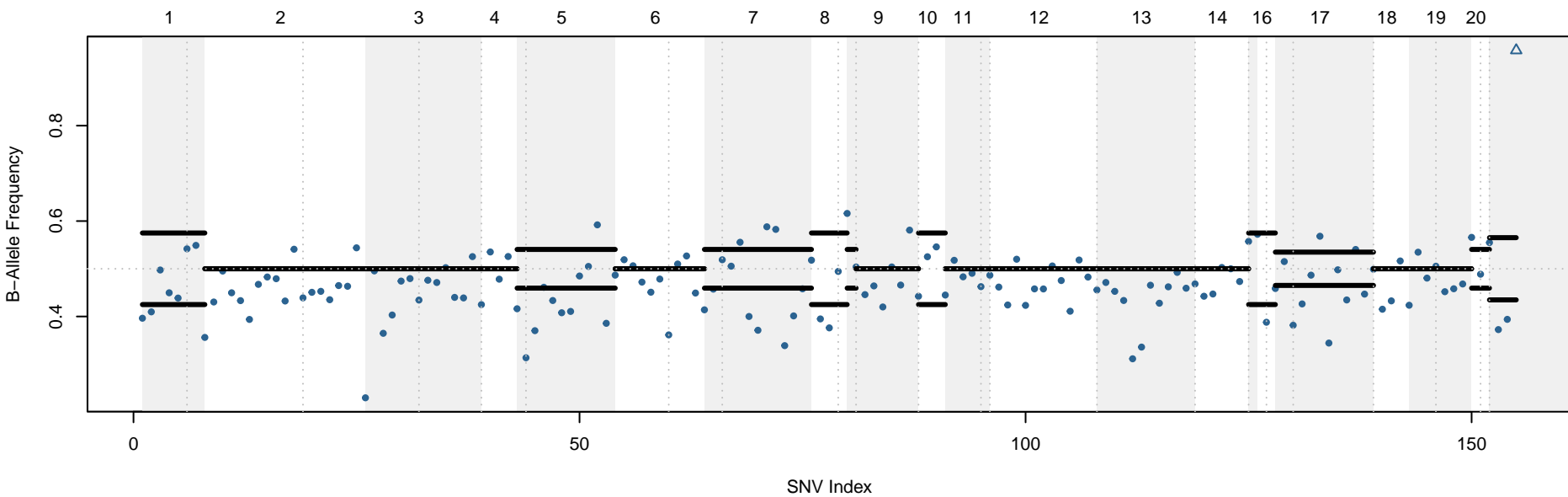


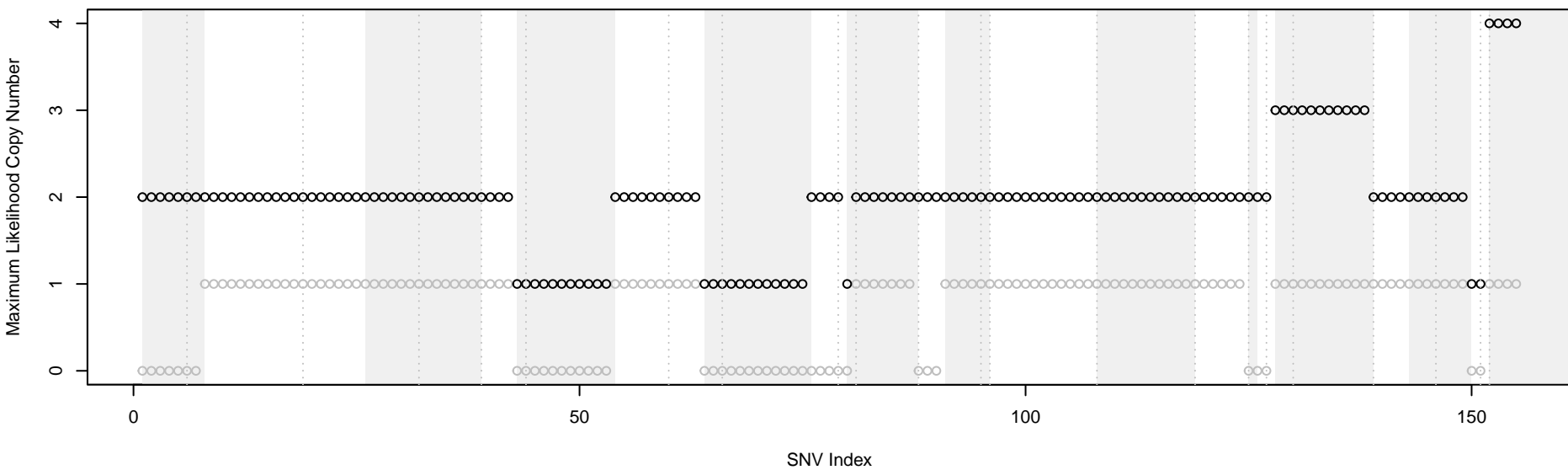
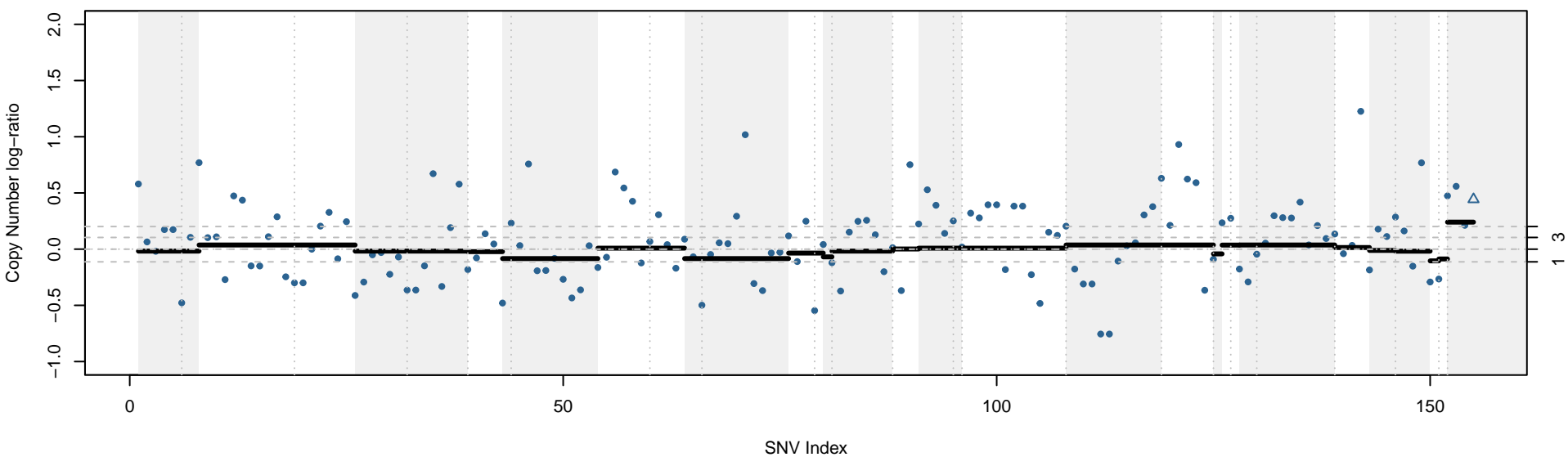
Purity: 0.15 Tumor ploidy: 2.004

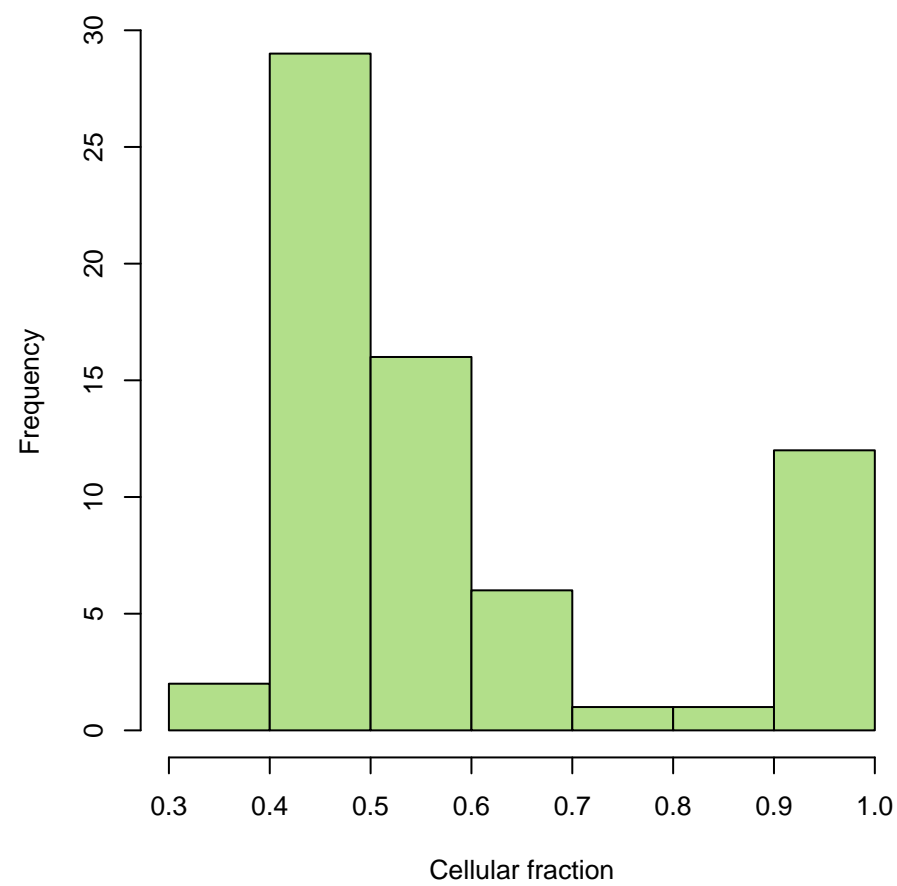
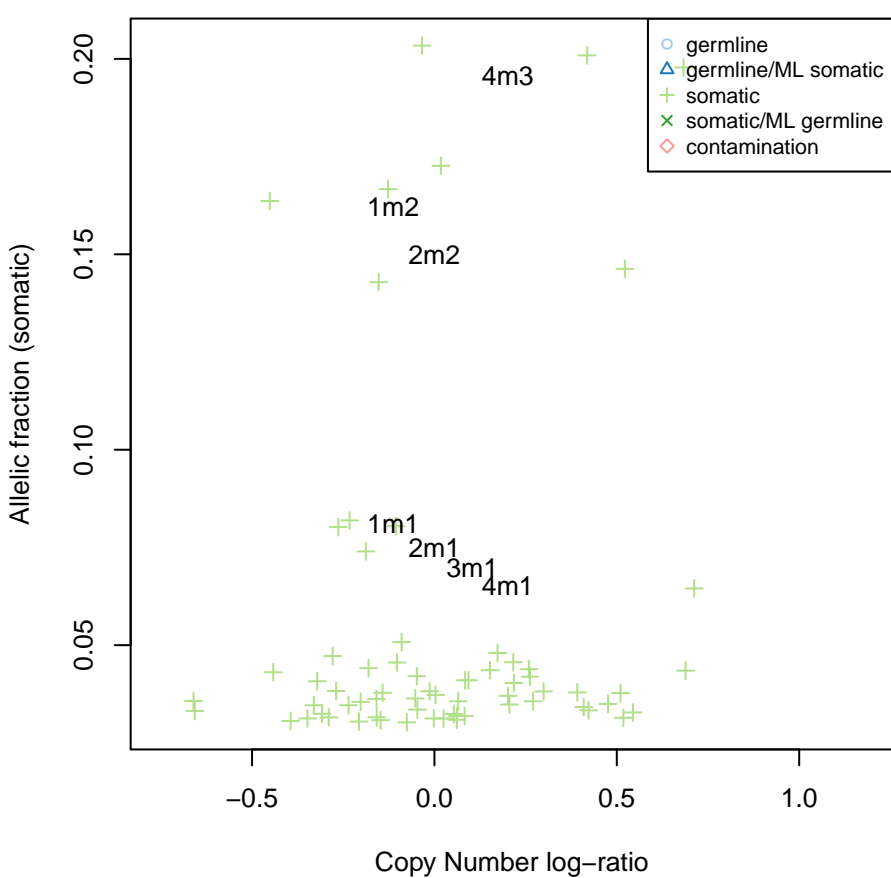
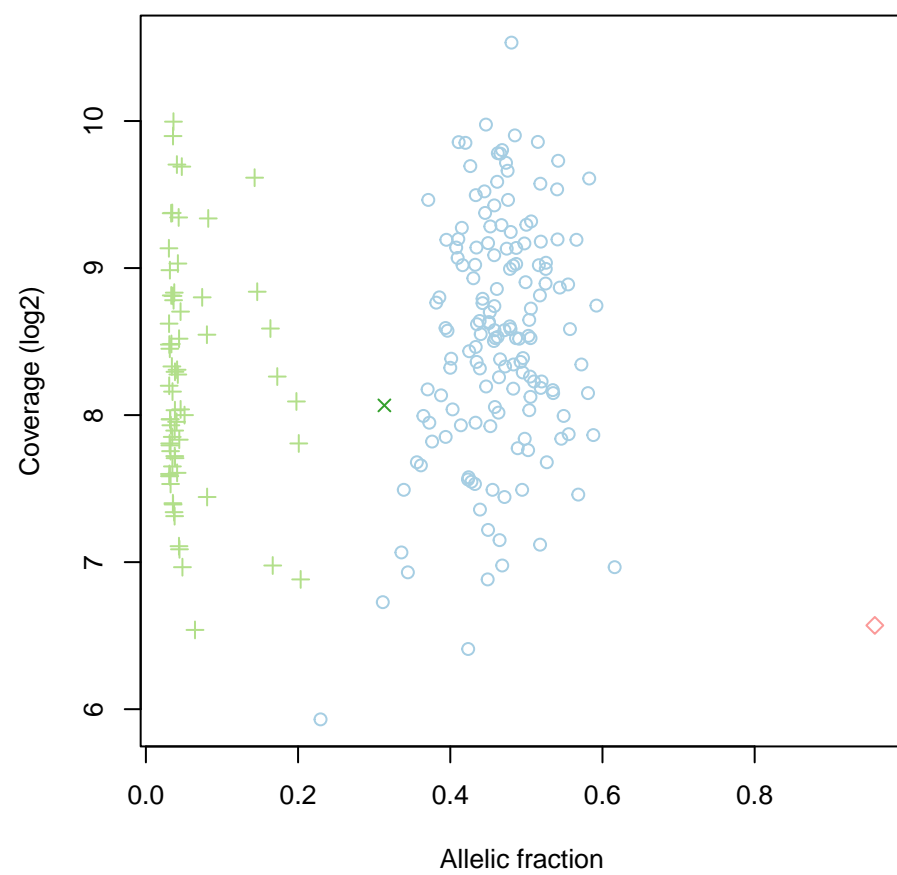
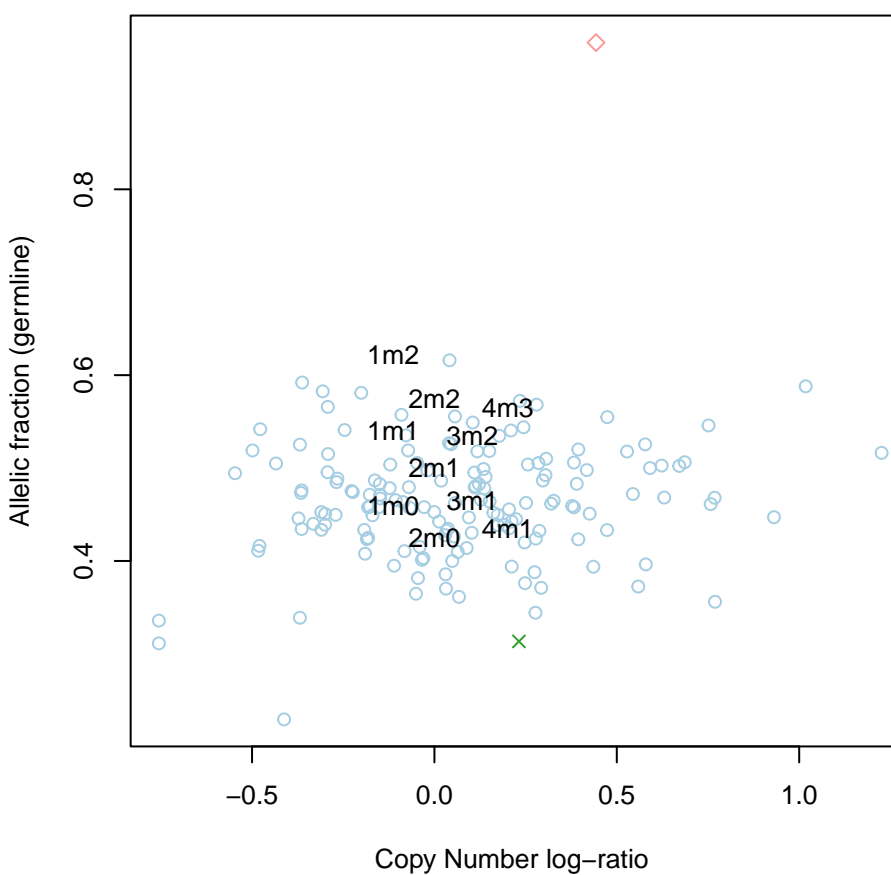


Purity: 0.15 Tumor ploidy: 2.004 SNV log-likelihood: 27.92 GoF: 94.4% Mean coverage: 401;62

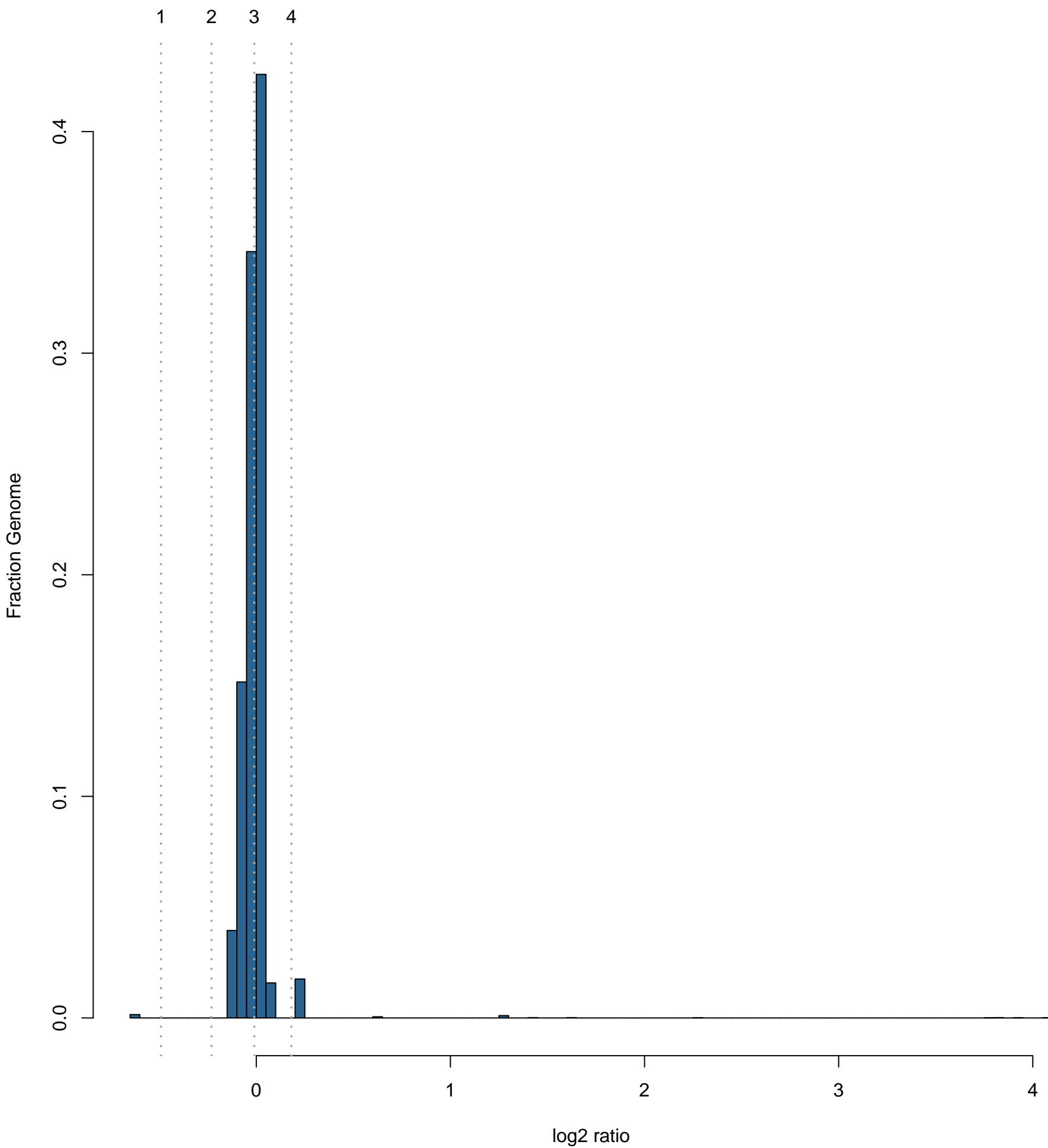


SCNA-fit log-likelihood: -15056.88

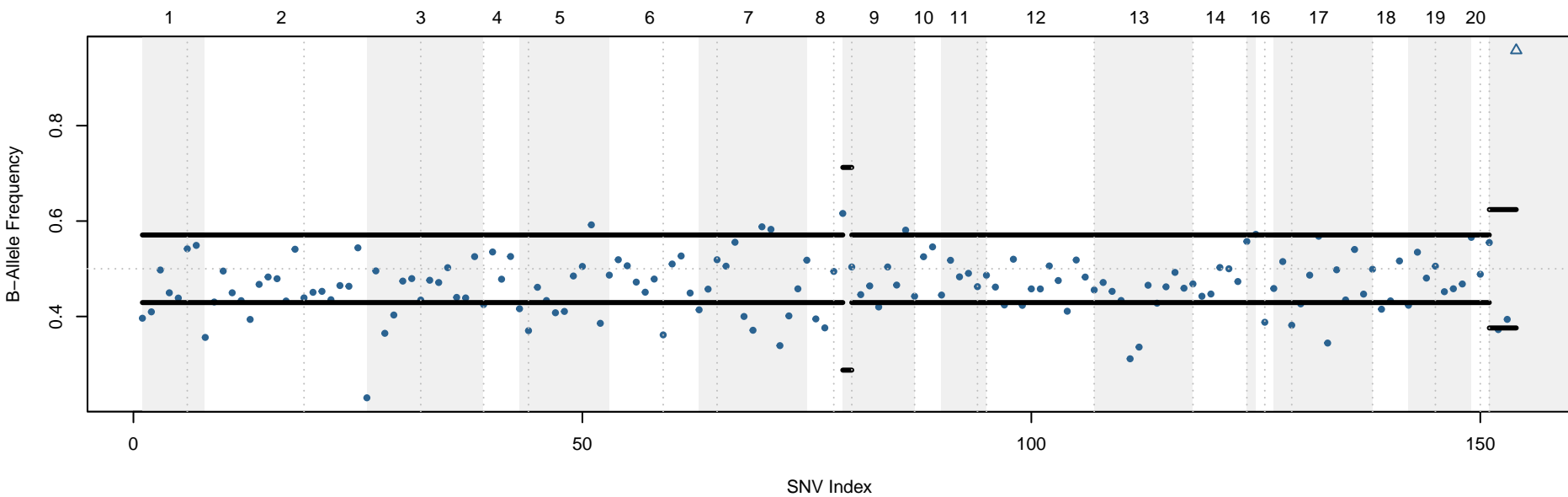




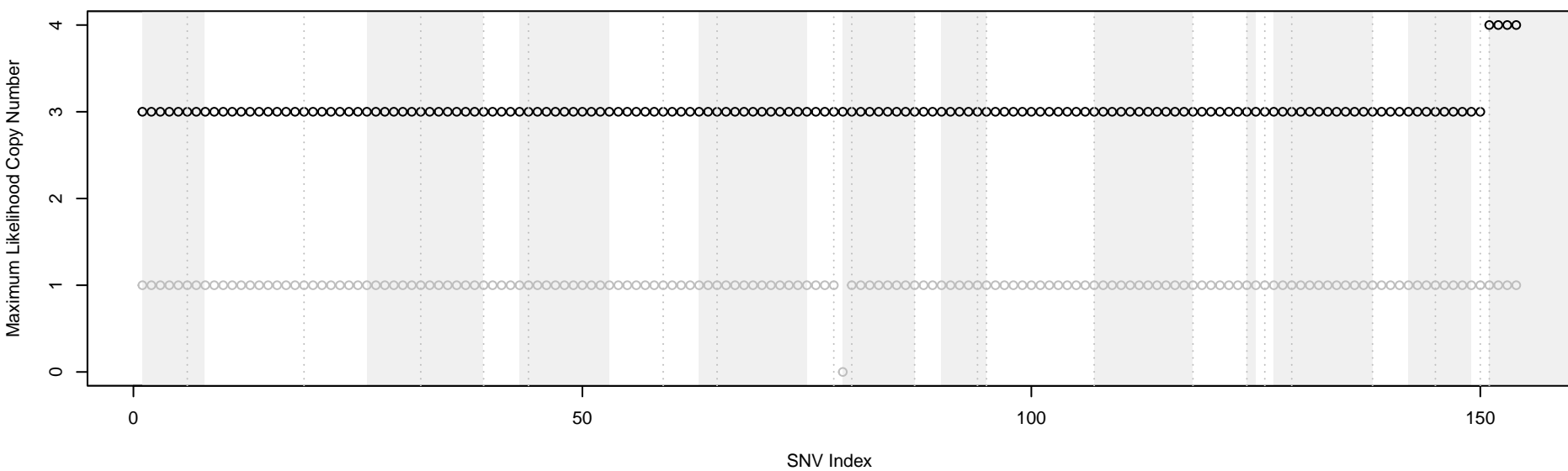
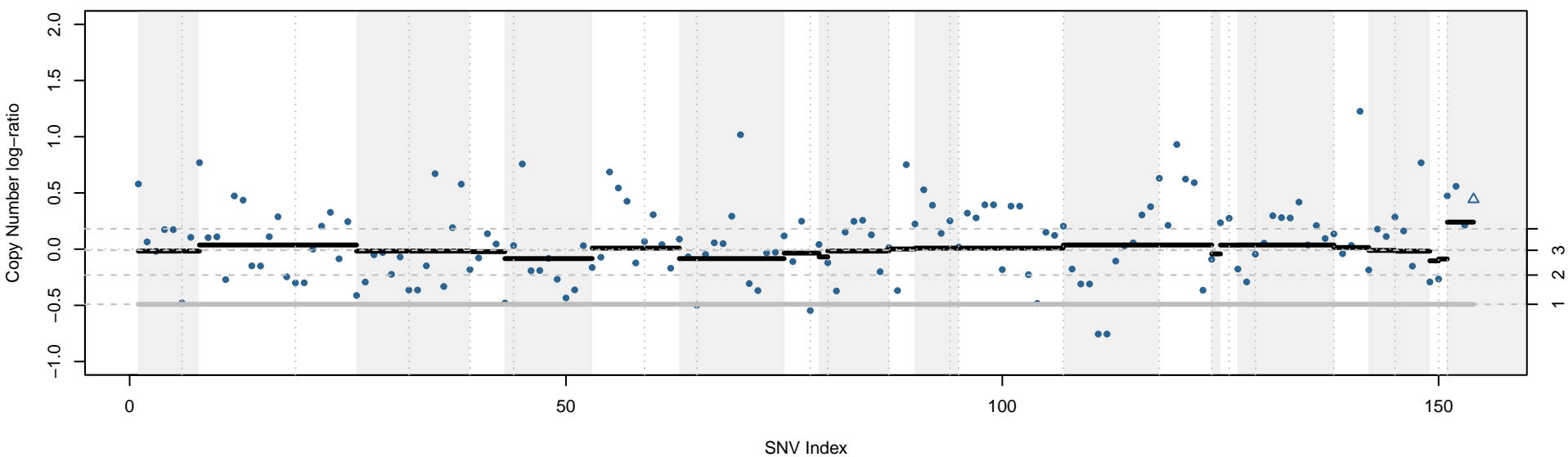
Purity: 0.33 Tumor ploidy: 3.05

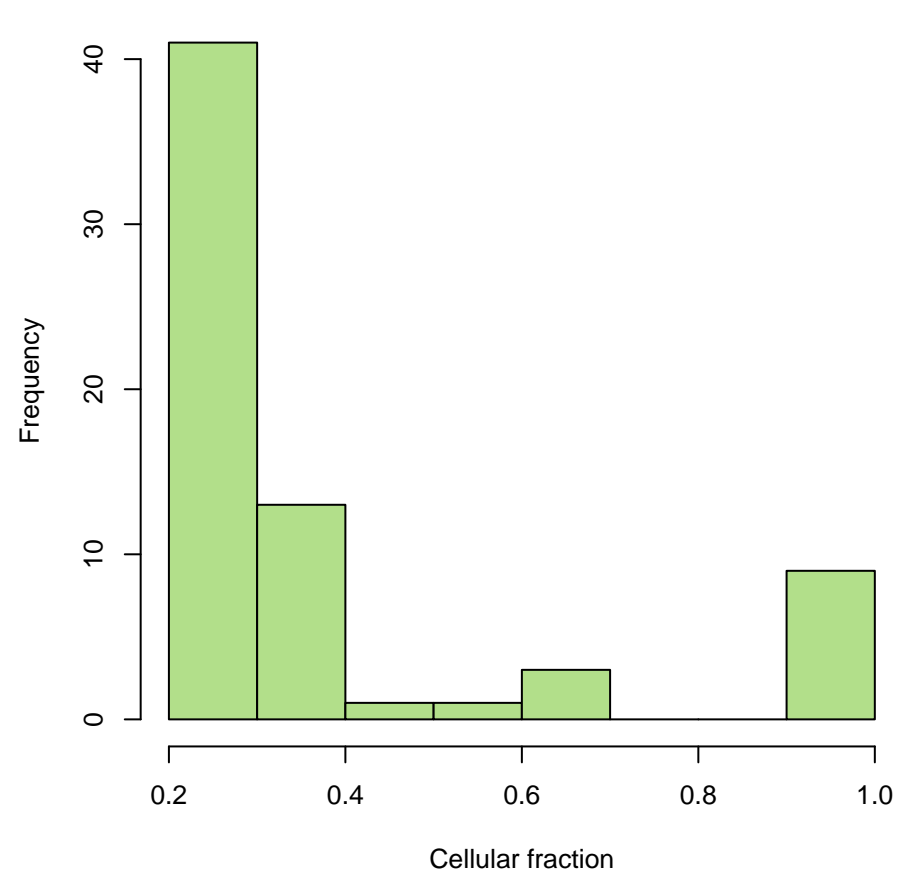
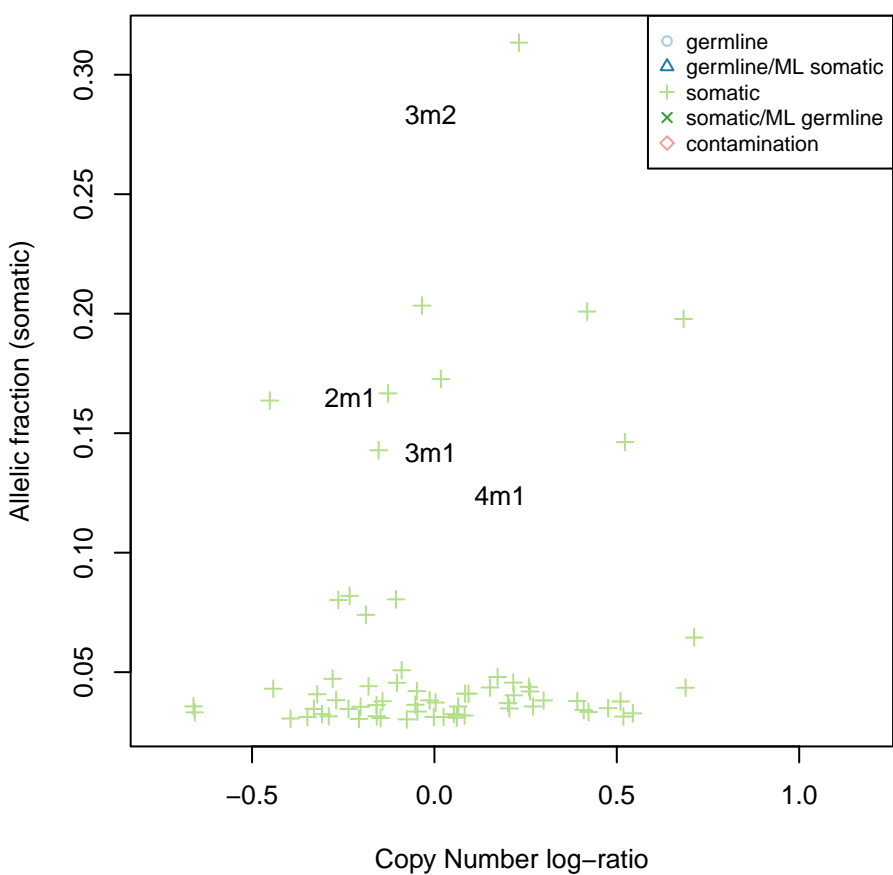
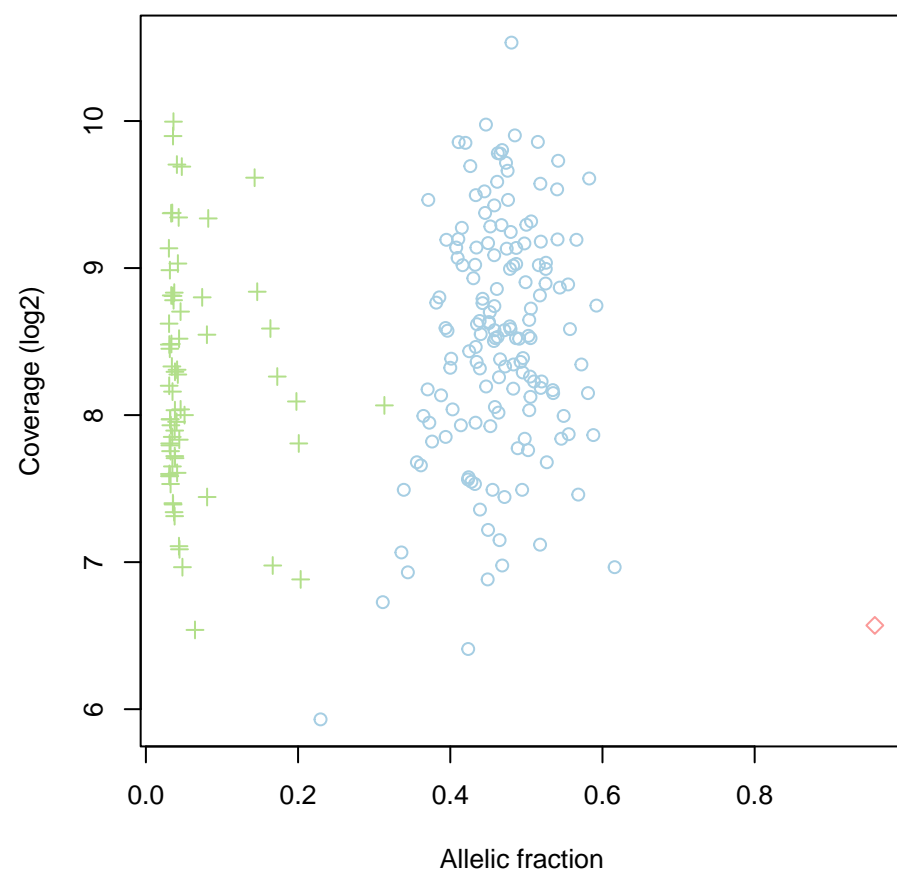
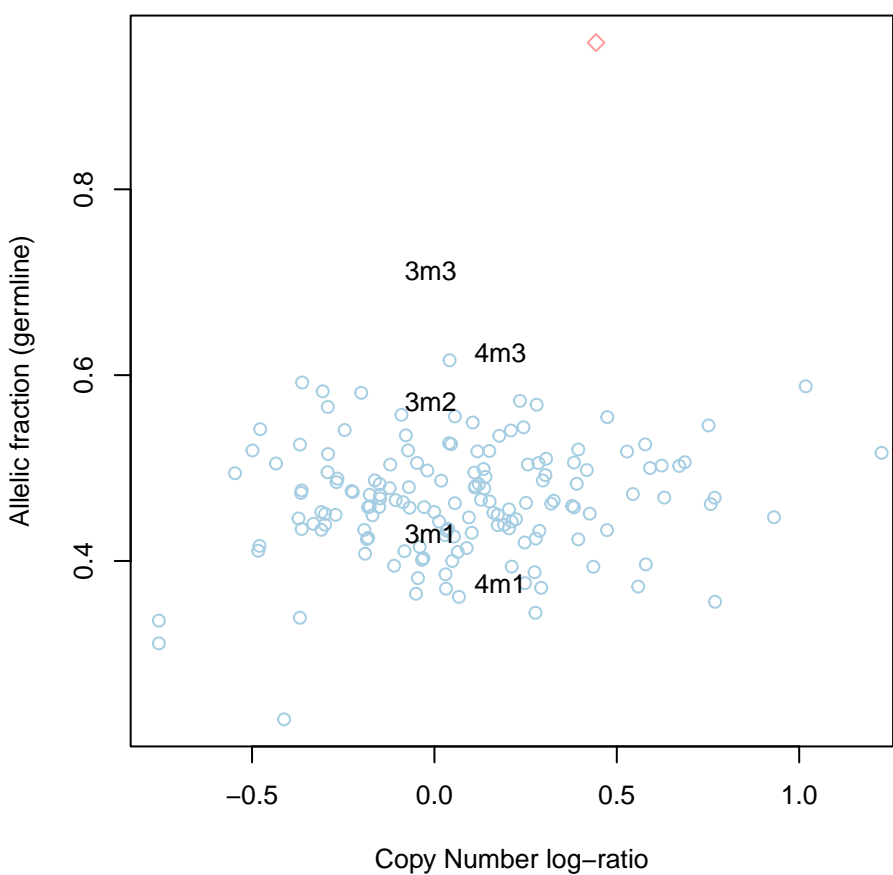


Purity: 0.33 Tumor ploidy: 3.05 SNV log-likelihood: -82.71 GoF: 89.4% Mean coverage: 401;62

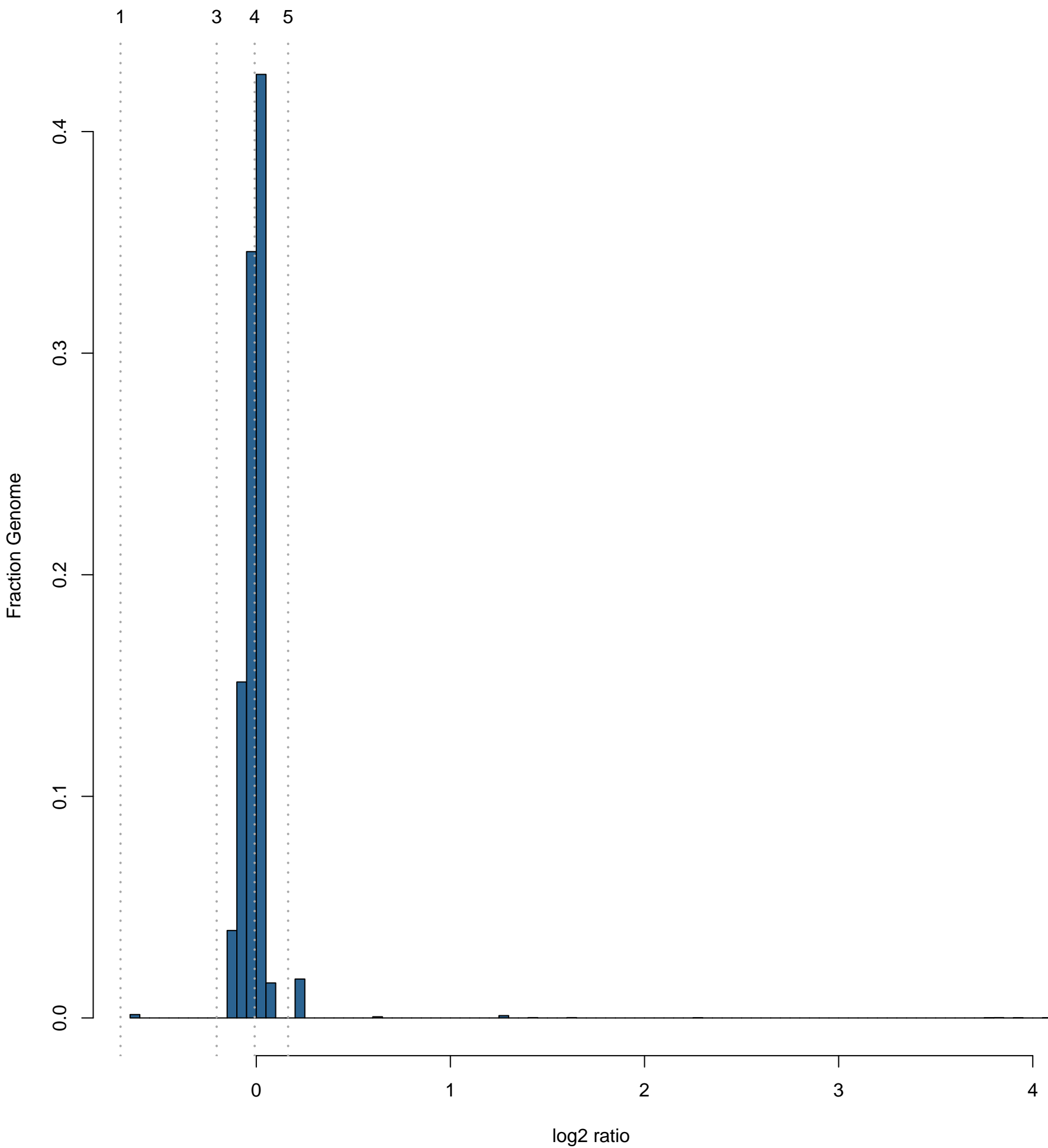


SCNA-fit log-likelihood: -15096.14

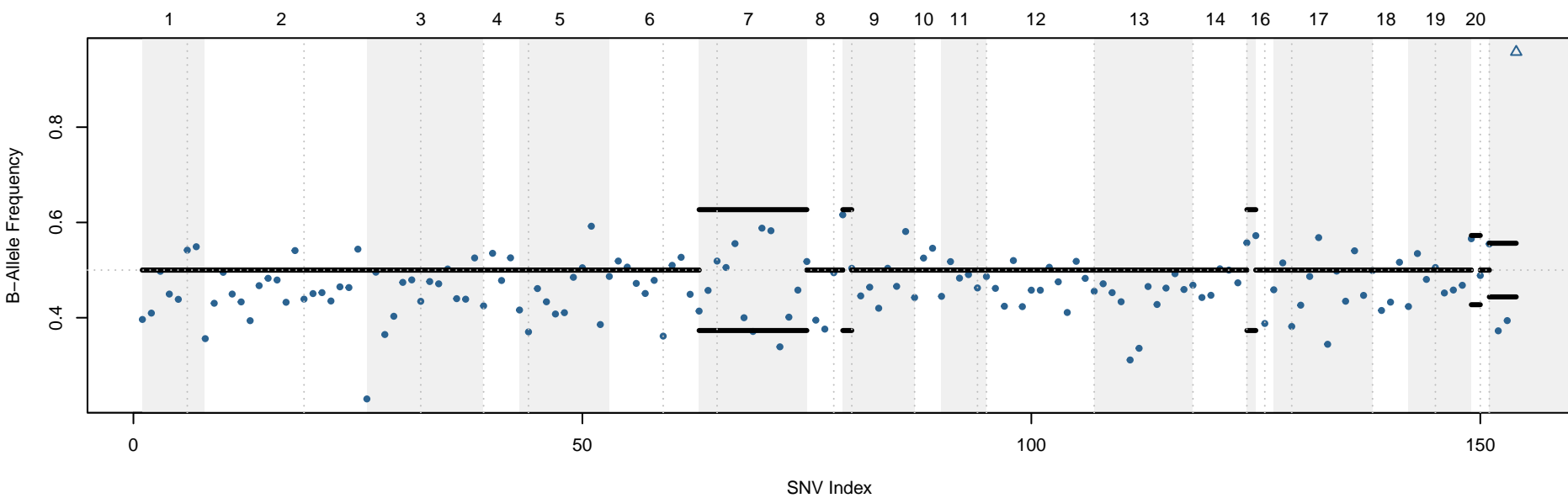




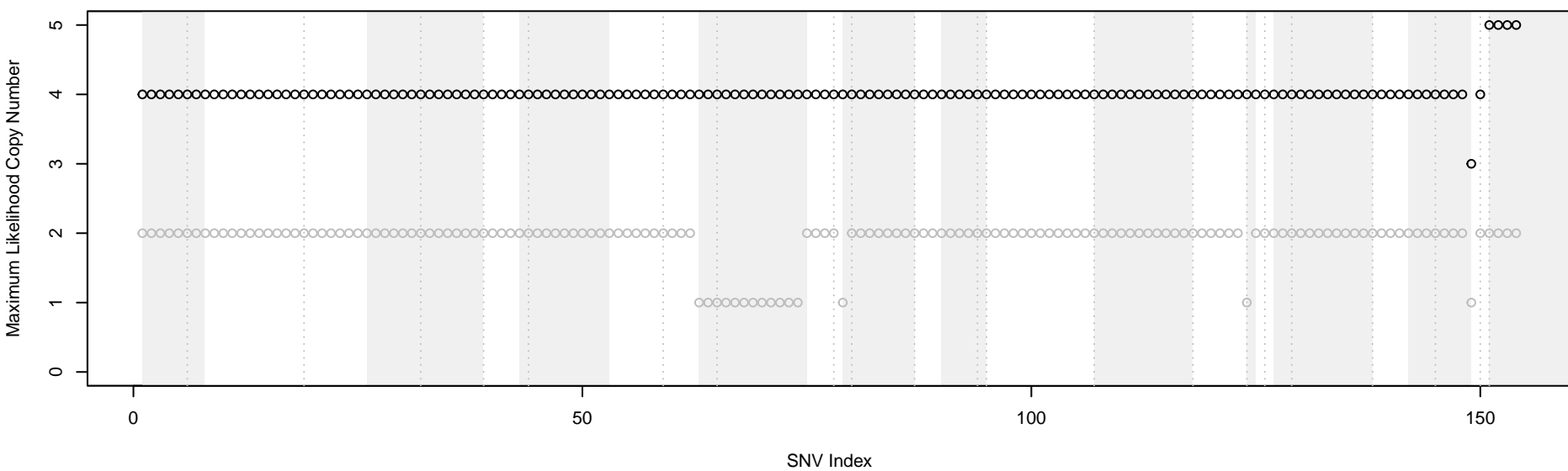
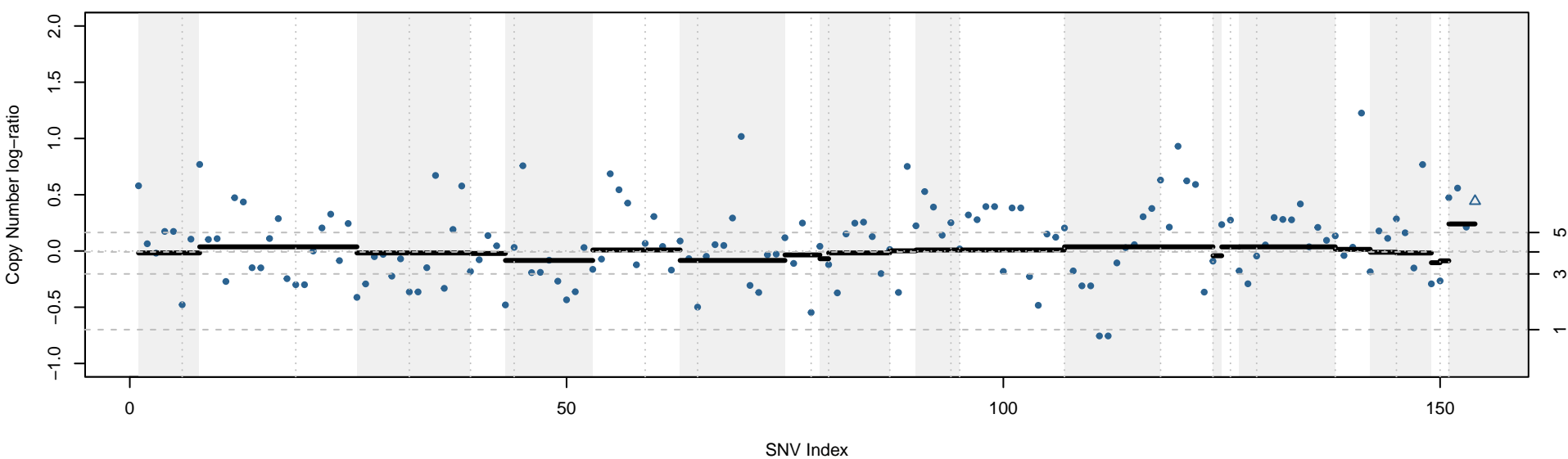
Purity: 0.34 Tumor ploidy: 4.045



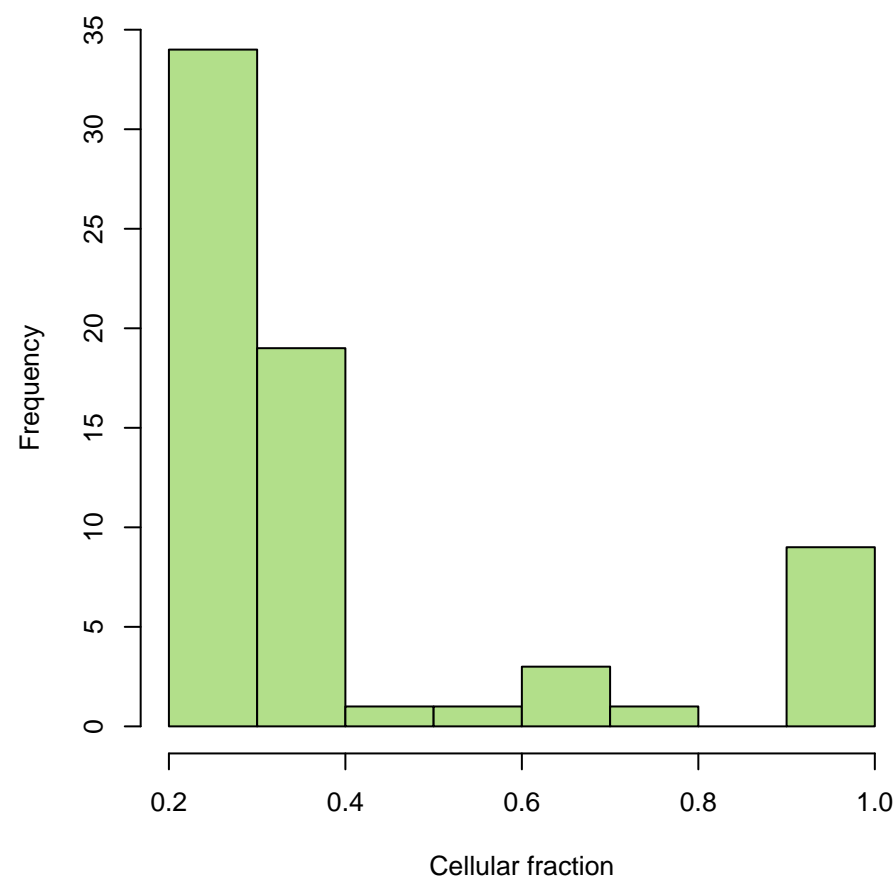
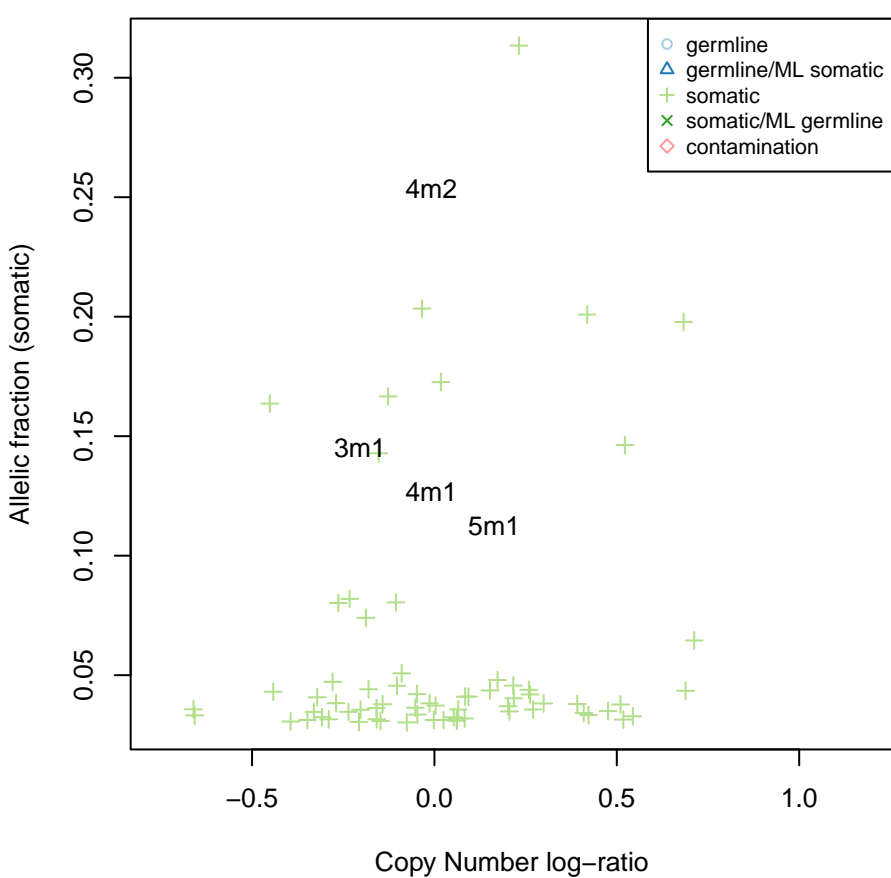
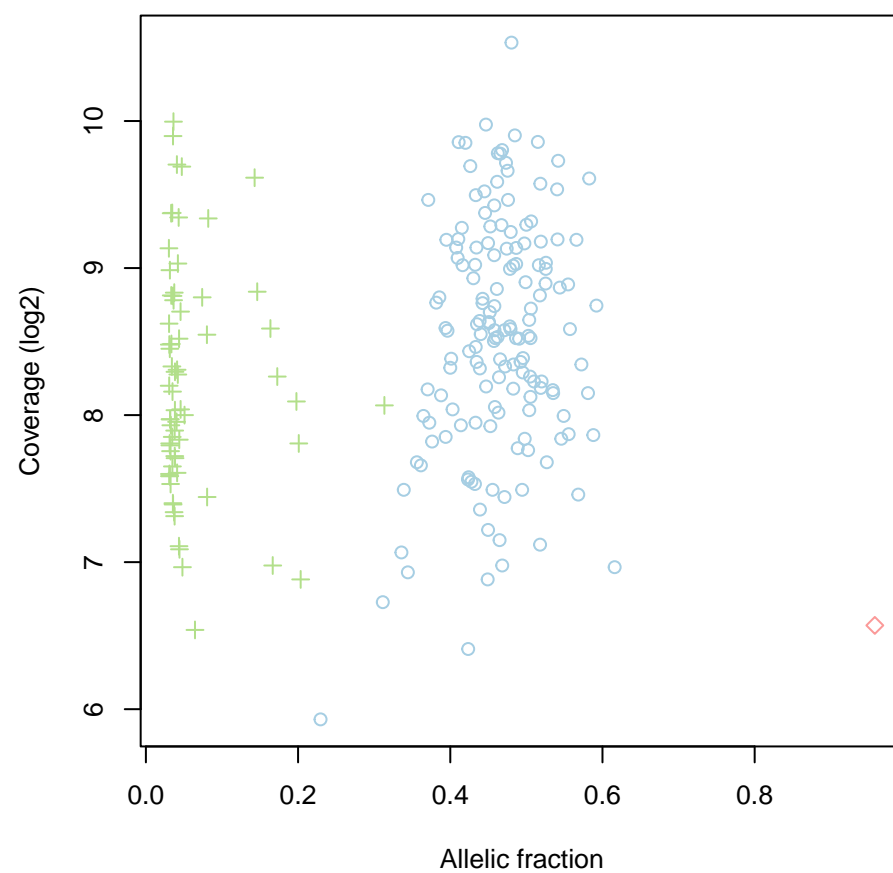
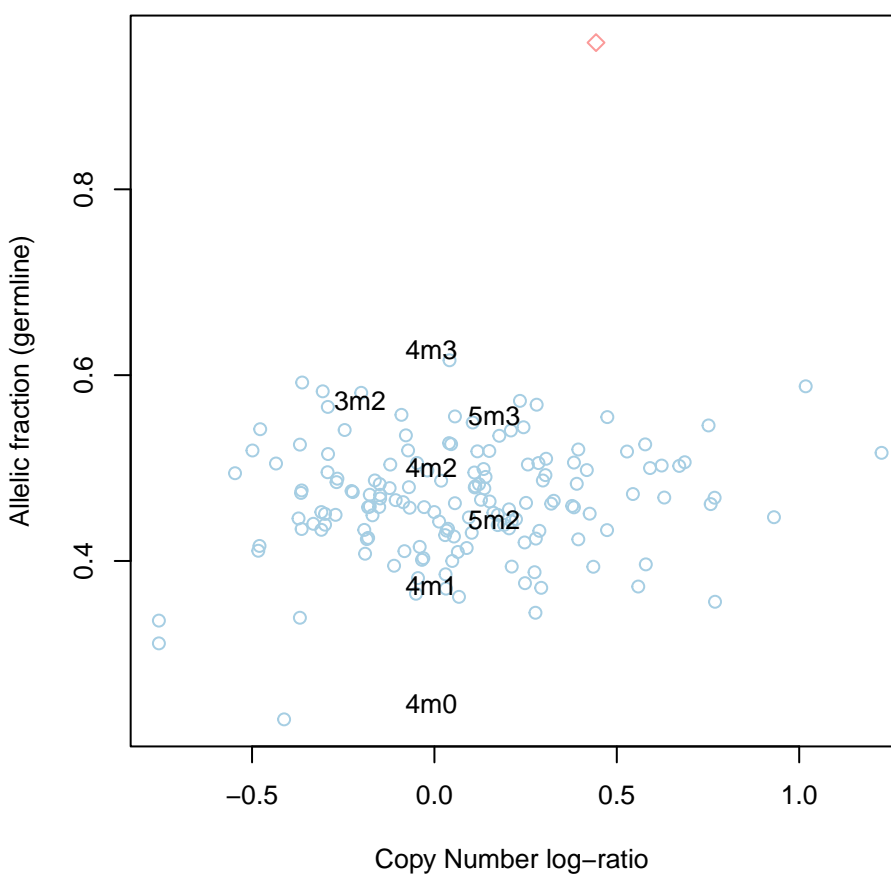
Purity: 0.34 Tumor ploidy: 4.045 SNV log-likelihood: -94.39 GoF: 90% Mean coverage: 401;62



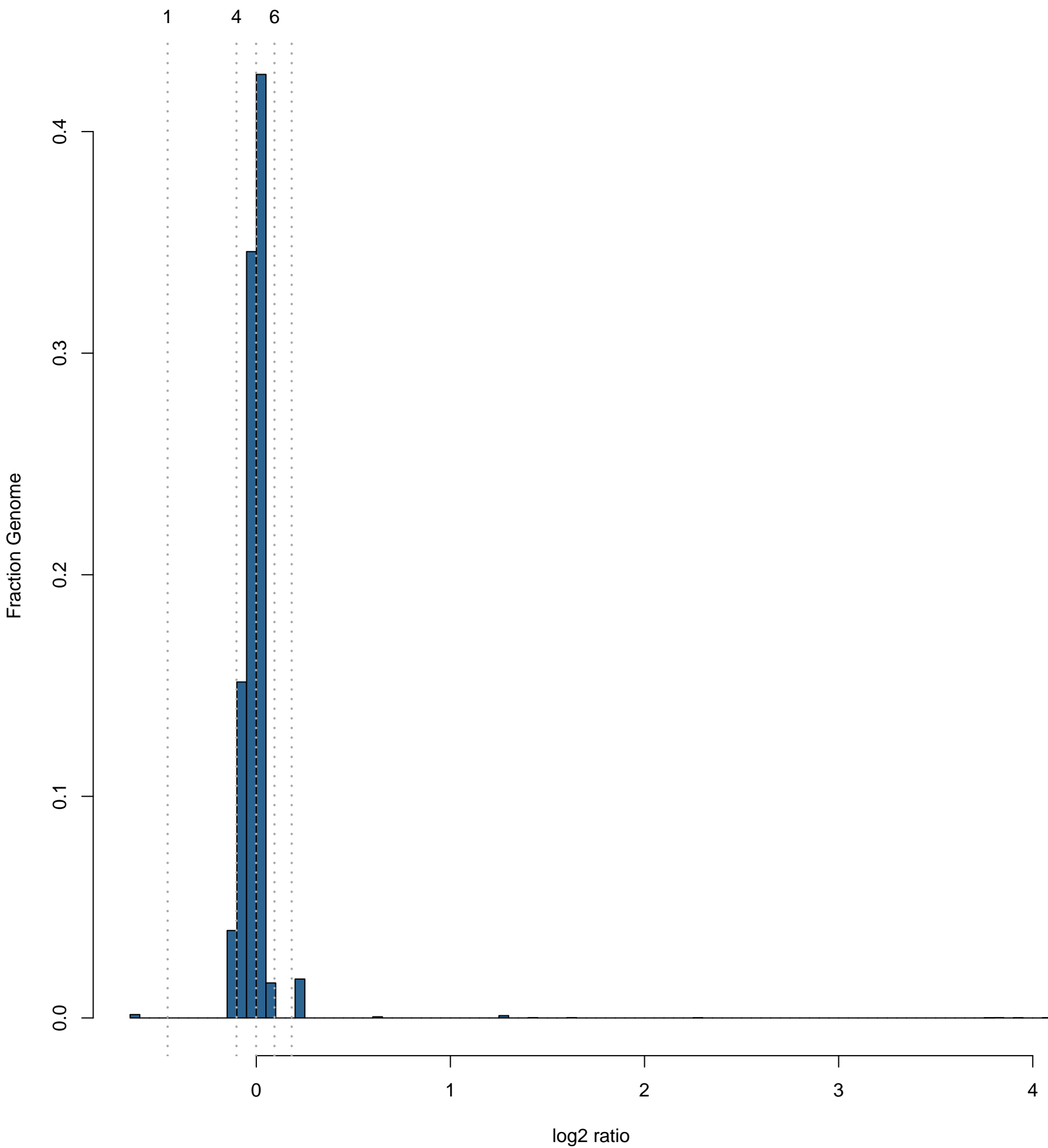
SCNA-fit log-likelihood: -15092.47



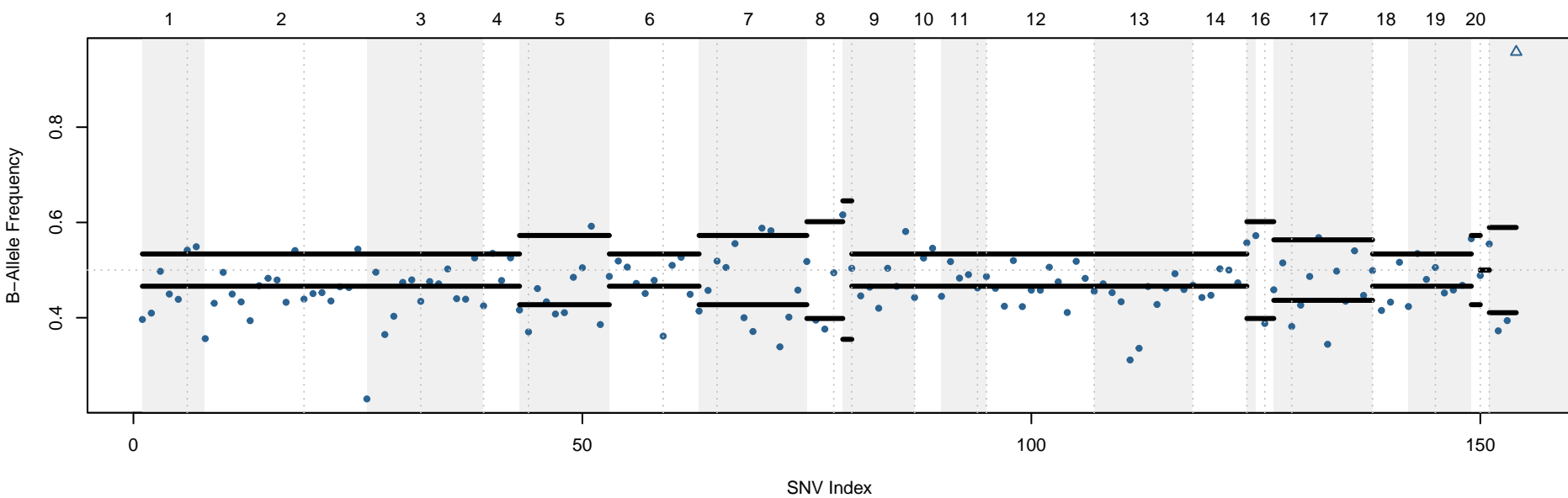




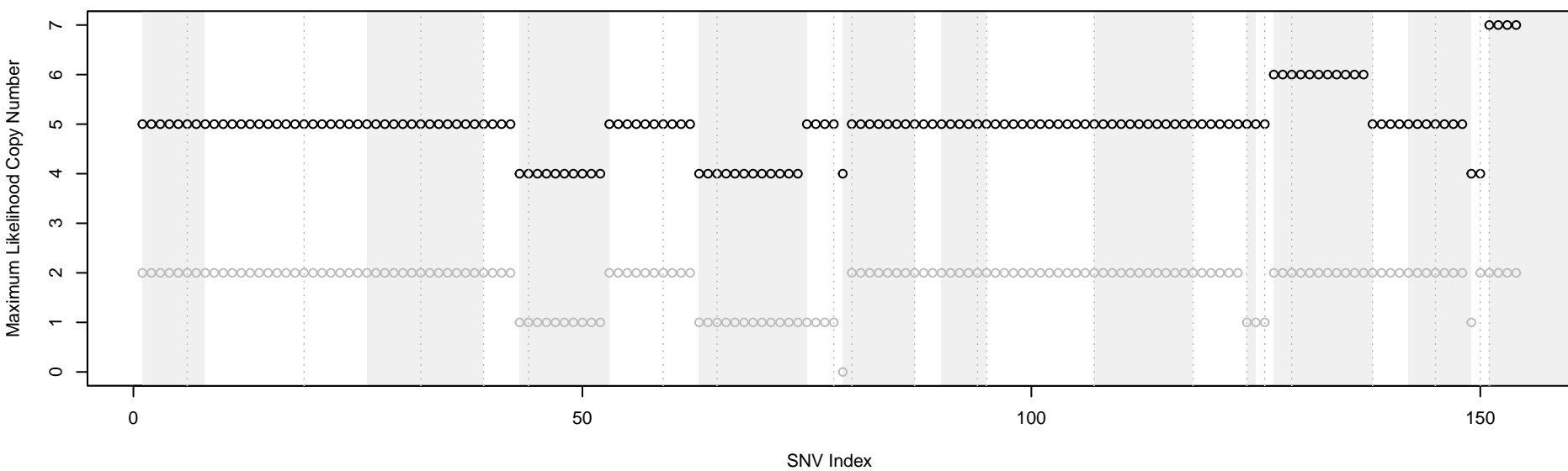
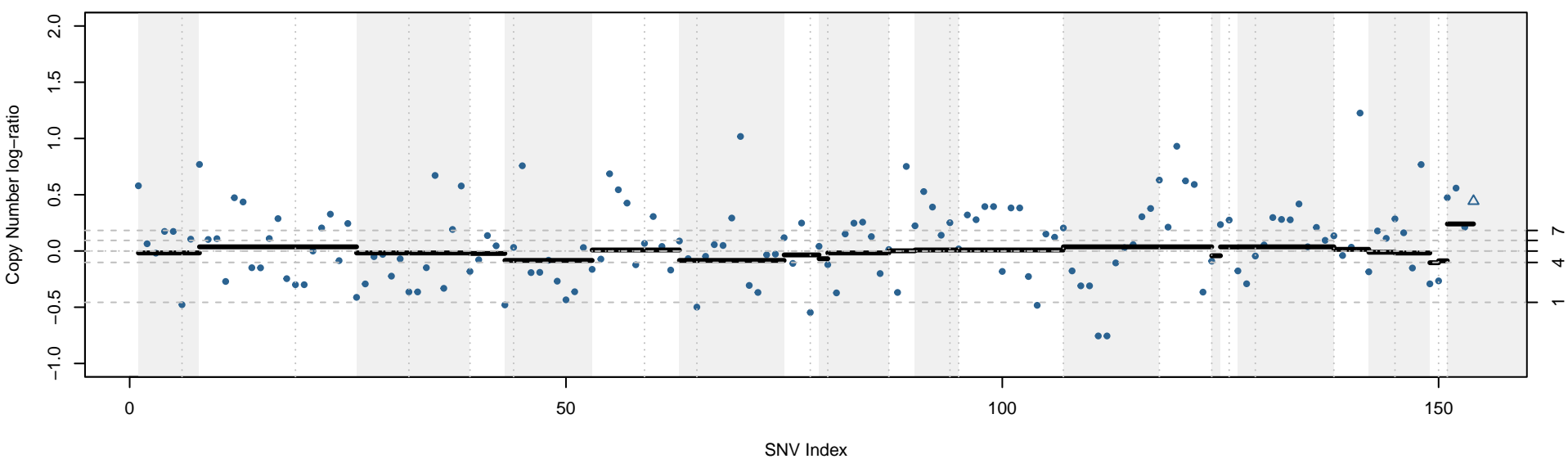
Purity: 0.17 Tumor ploidy: 5.008

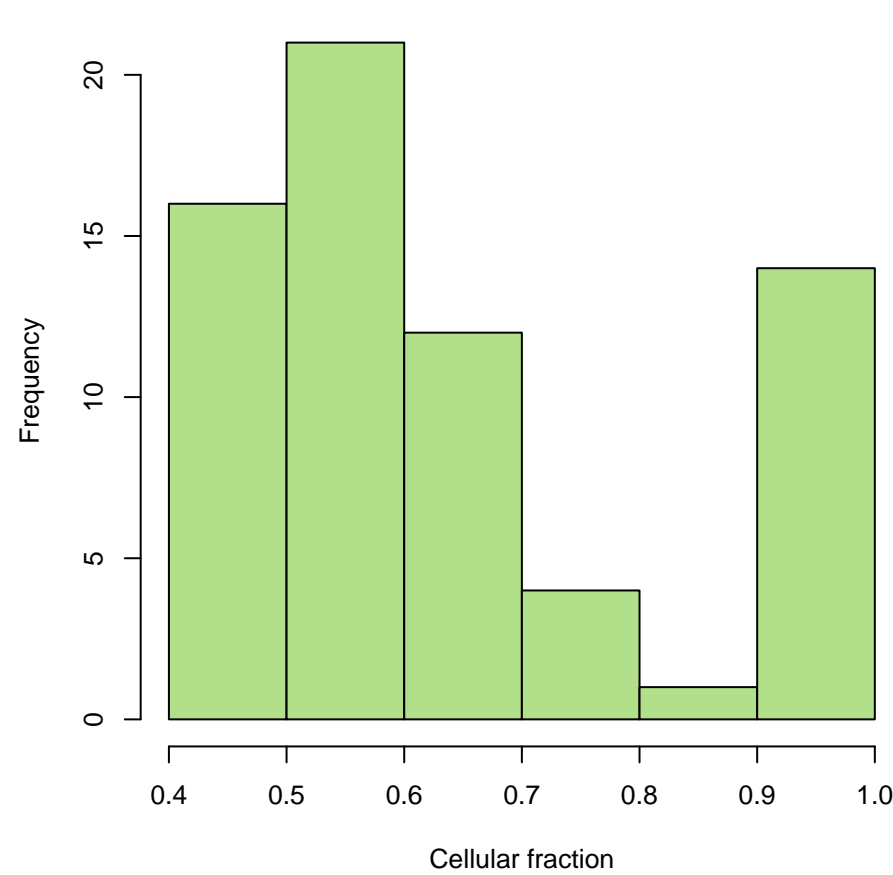
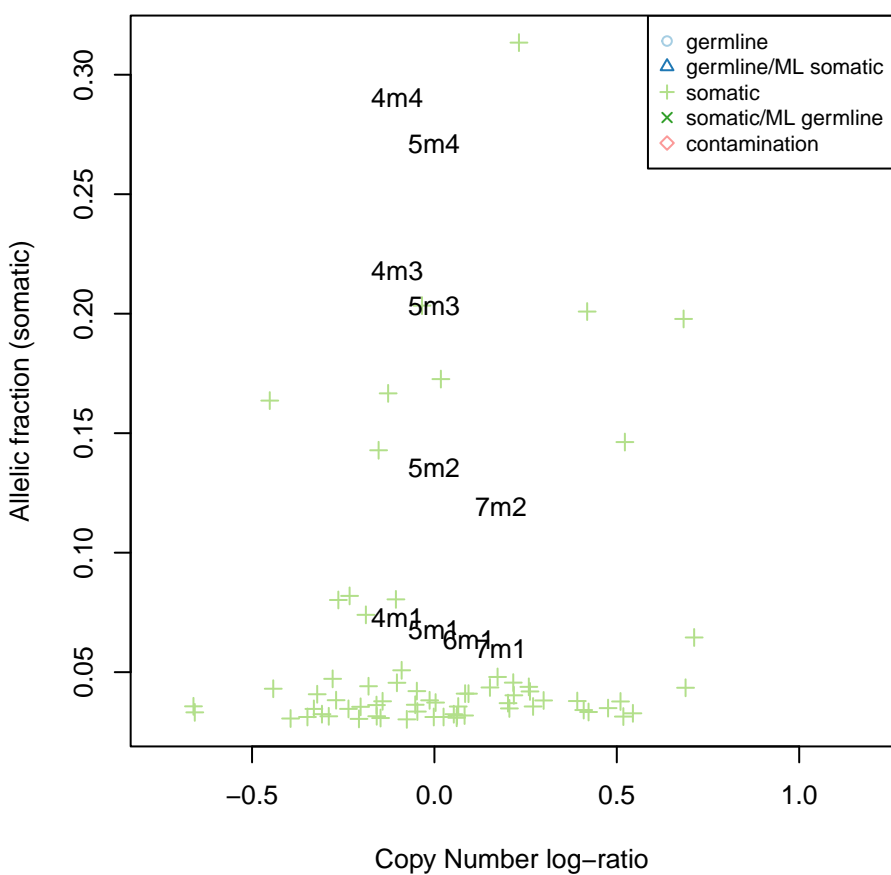
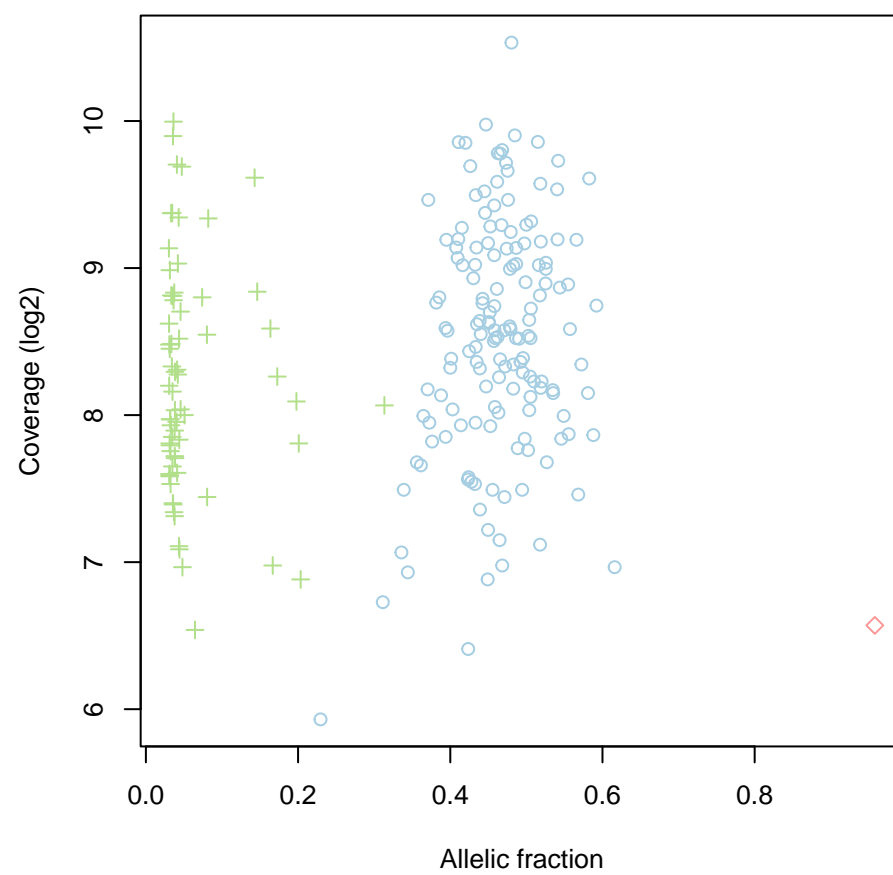
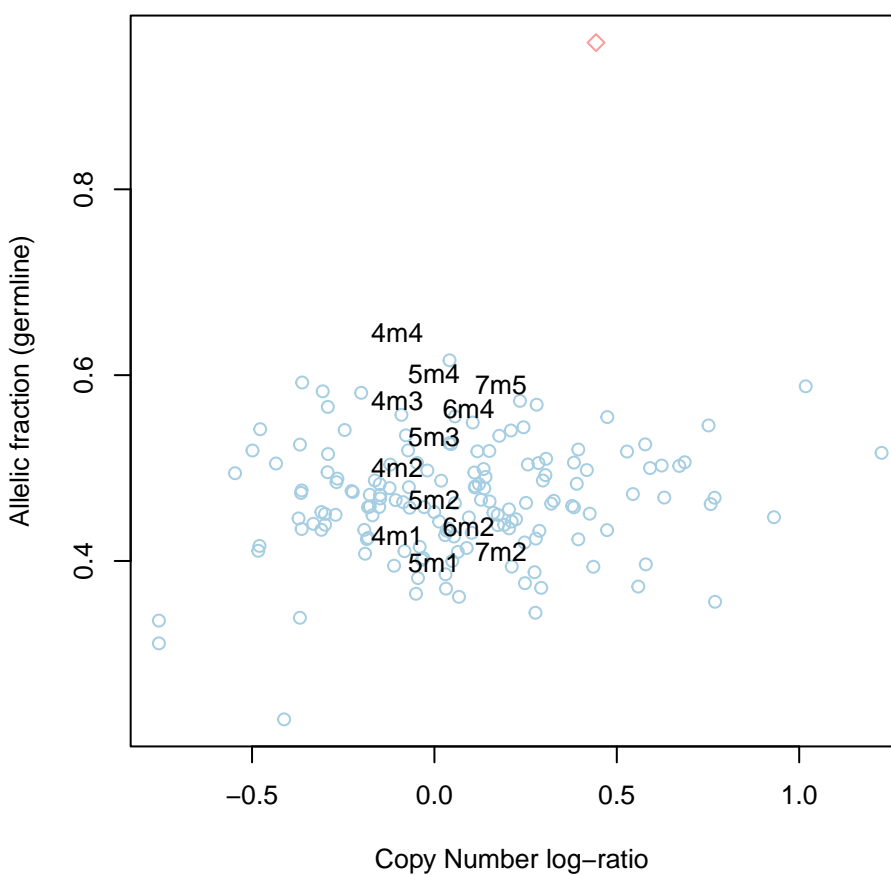


Purity: 0.17 Tumor ploidy: 5.008 SNV log-likelihood: -118.48 GoF: 96.8% Mean coverage: 401;62

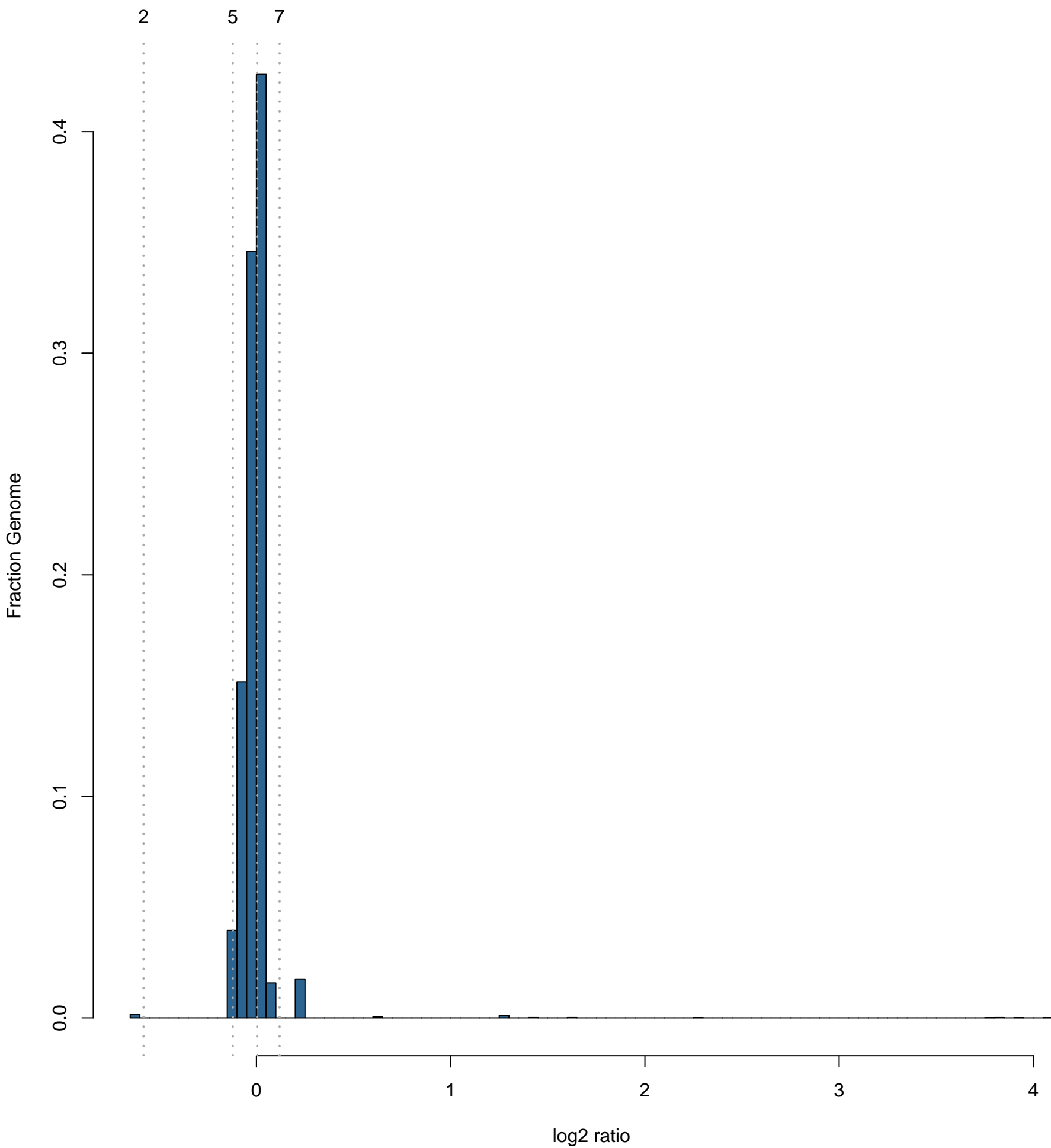


SCNA-fit log-likelihood: -15039

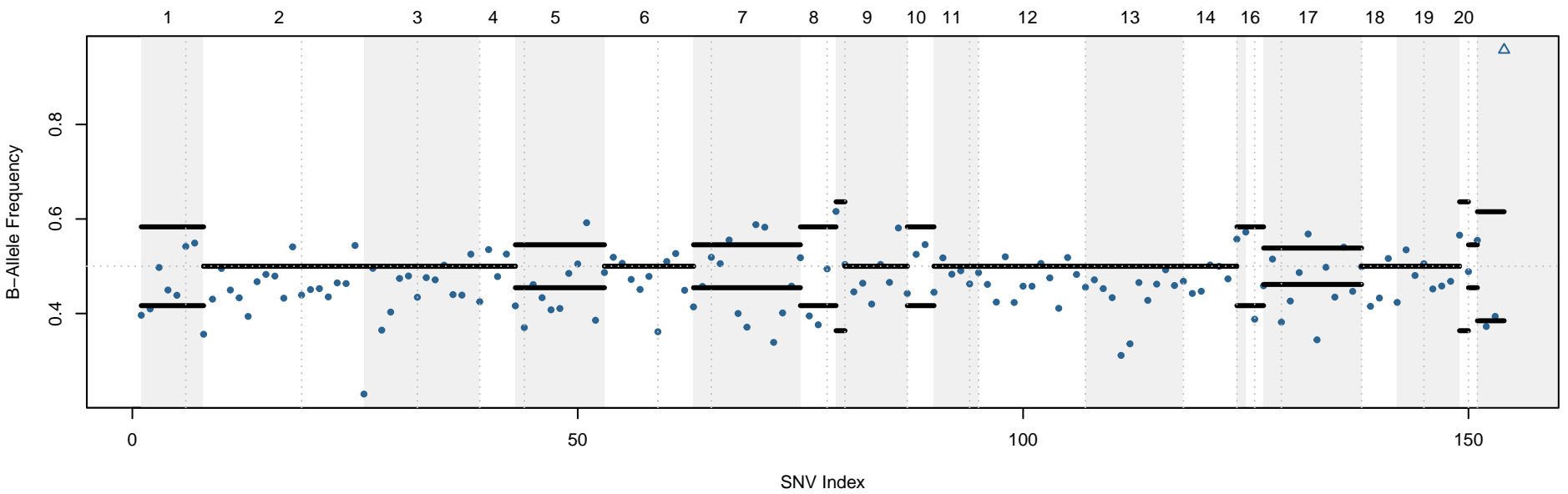




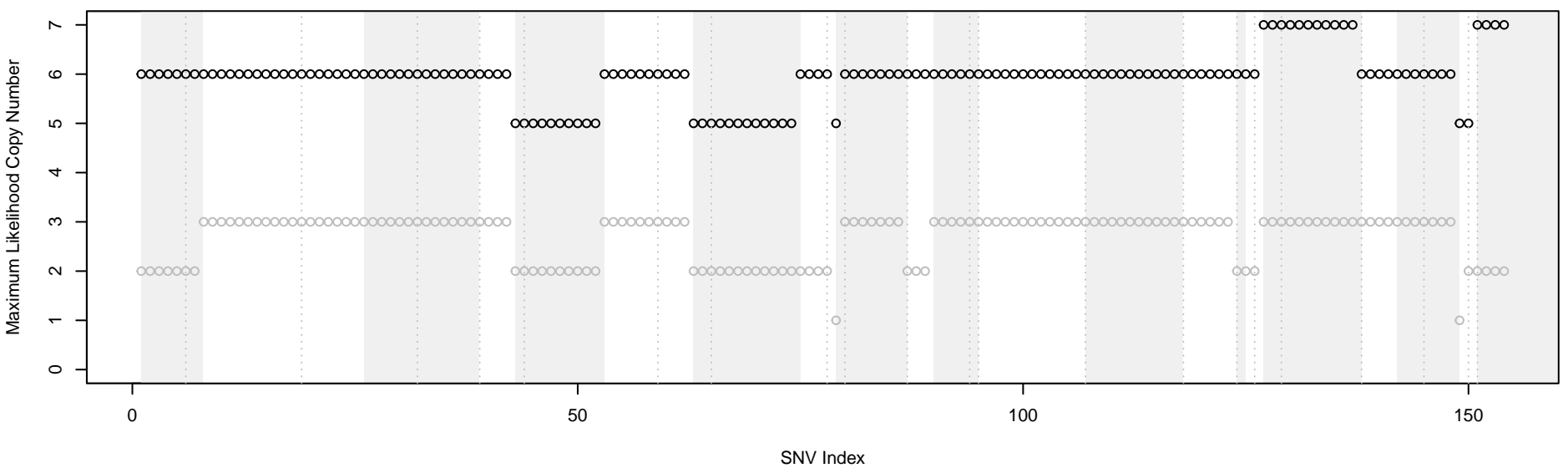
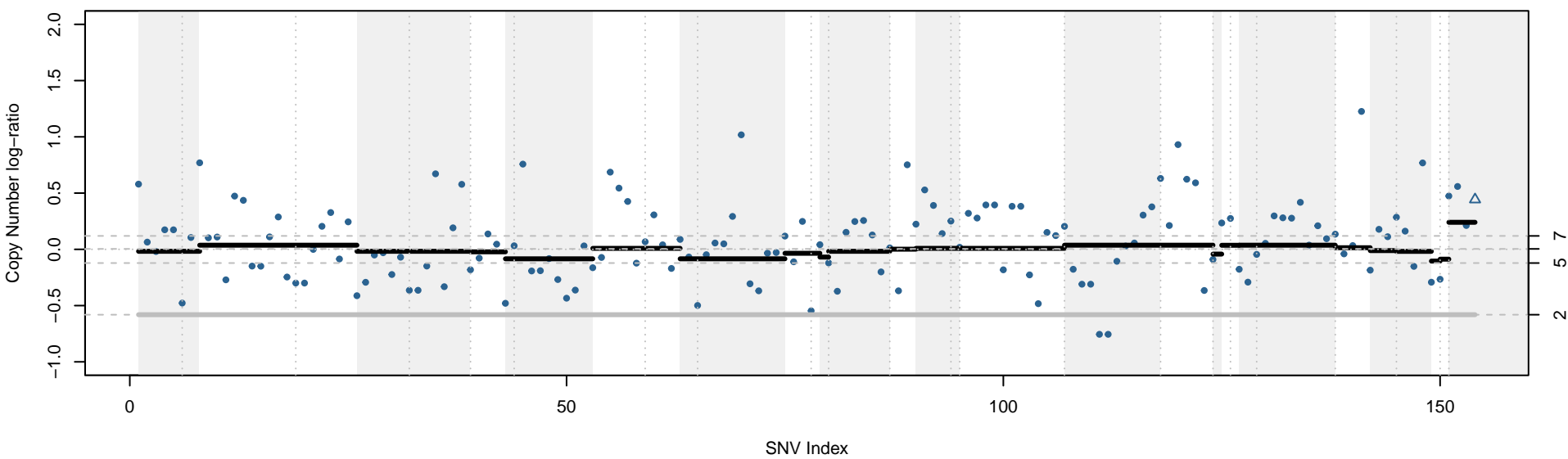
Purity: 0.25 Tumor ploidy: 5.968

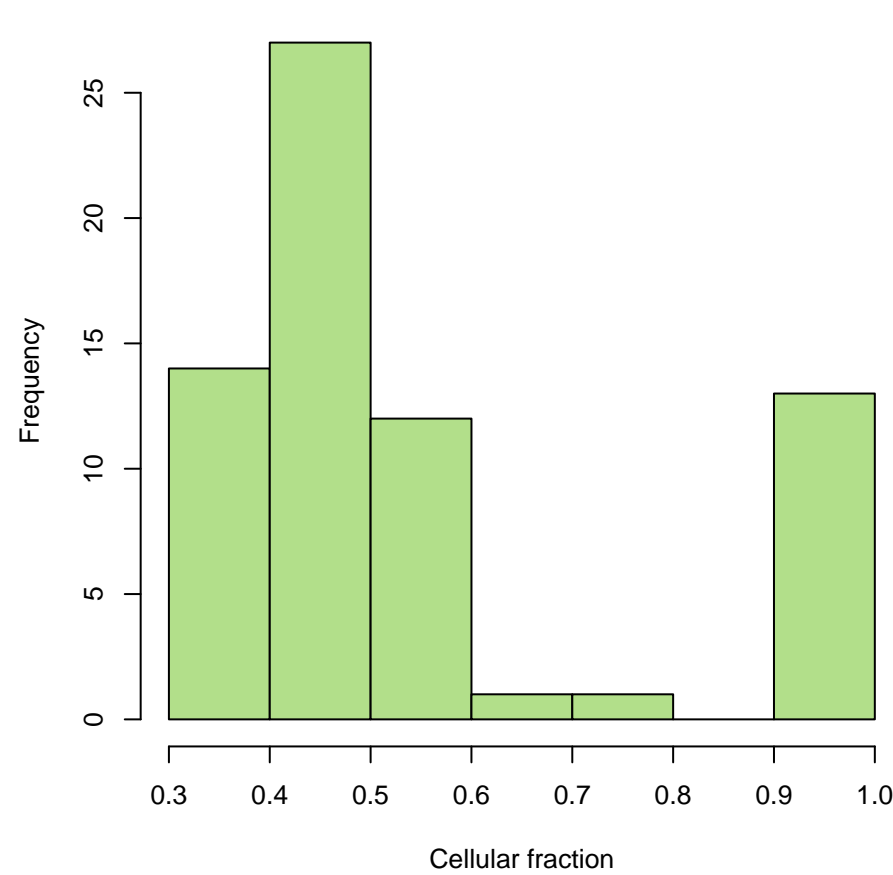
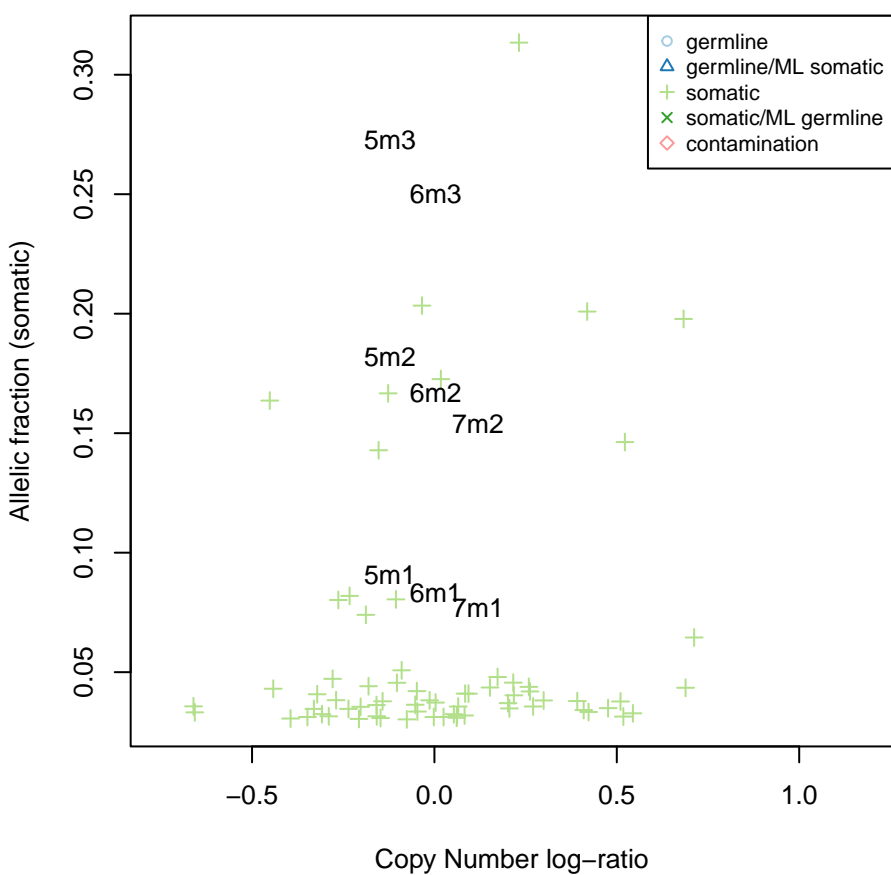
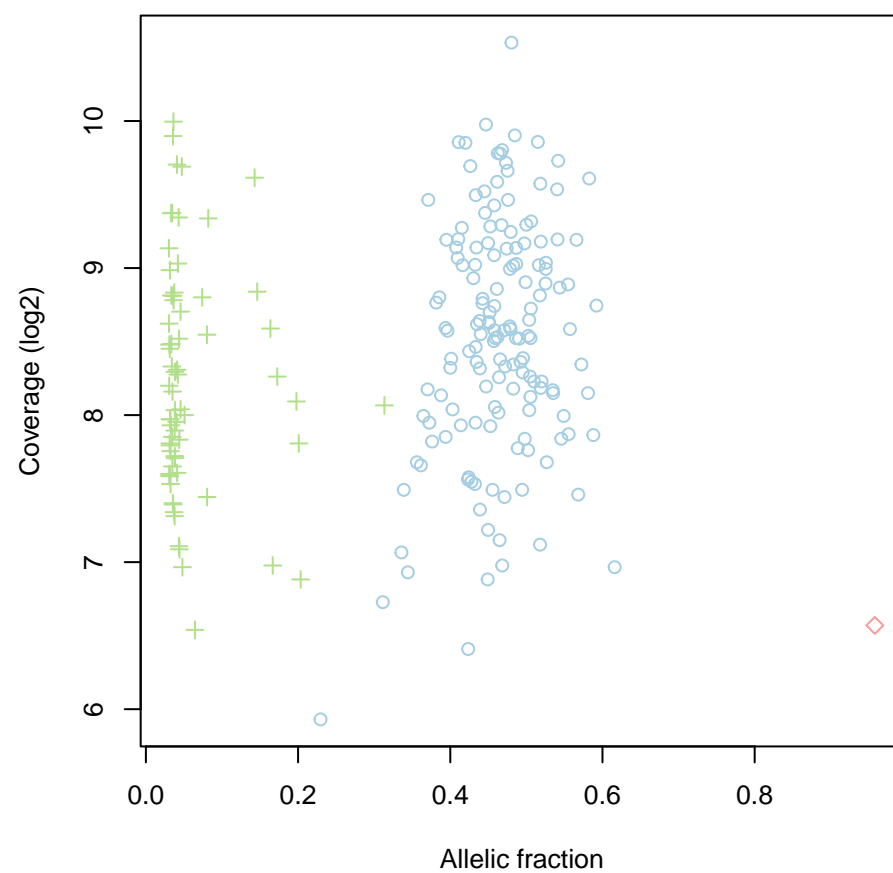
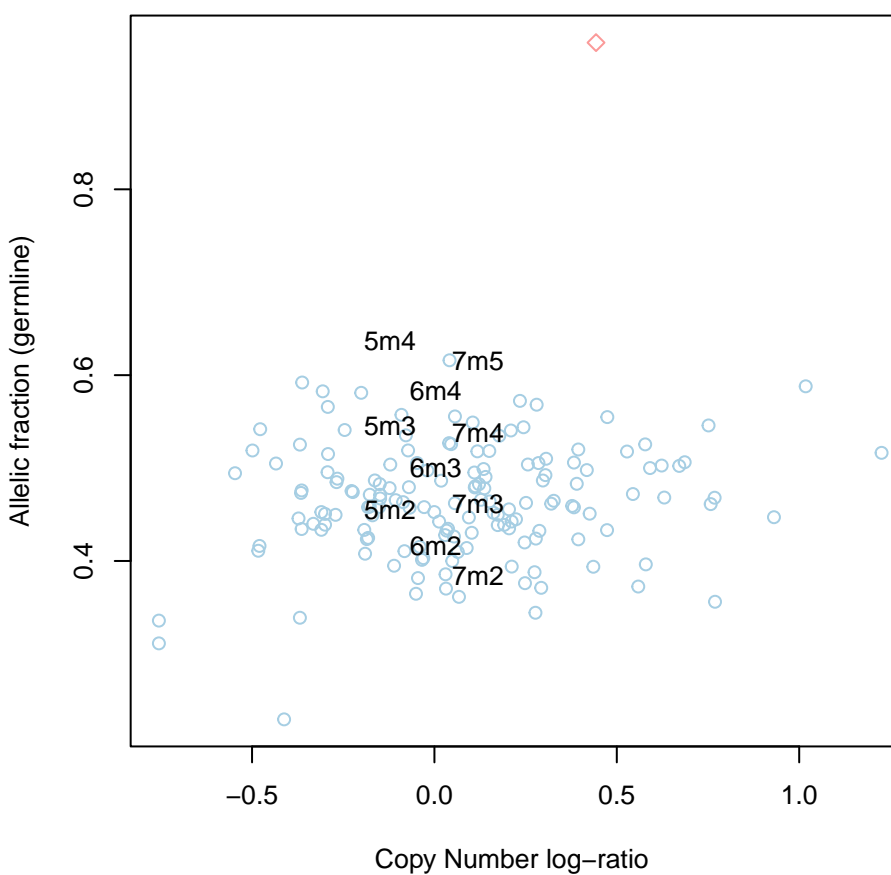


Purity: 0.25 Tumor ploidy: 5.968 SNV log-likelihood: -131.66 GoF: 94.2% Mean coverage: 401;62

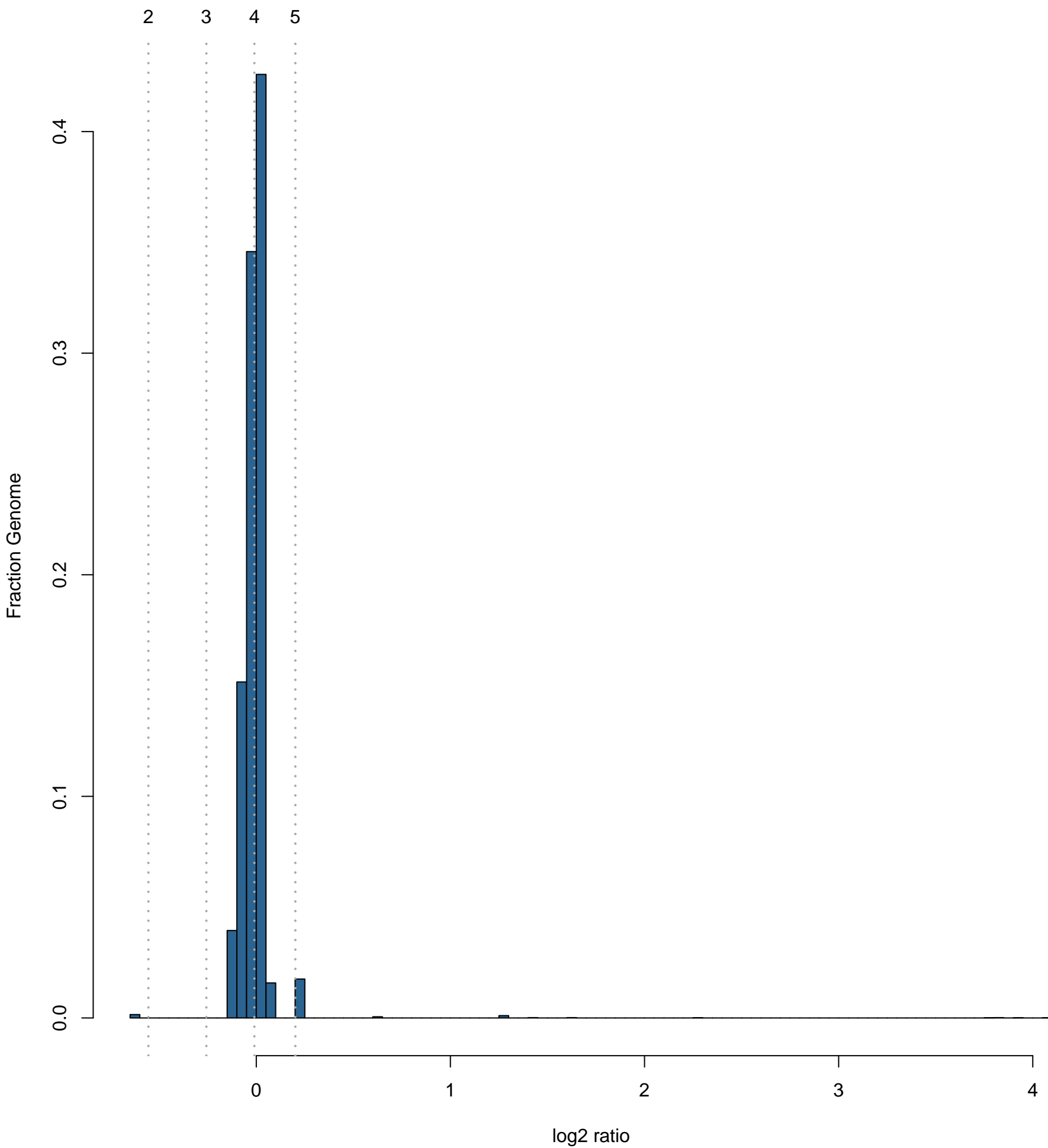


SCNA-fit log-likelihood: -15054.78



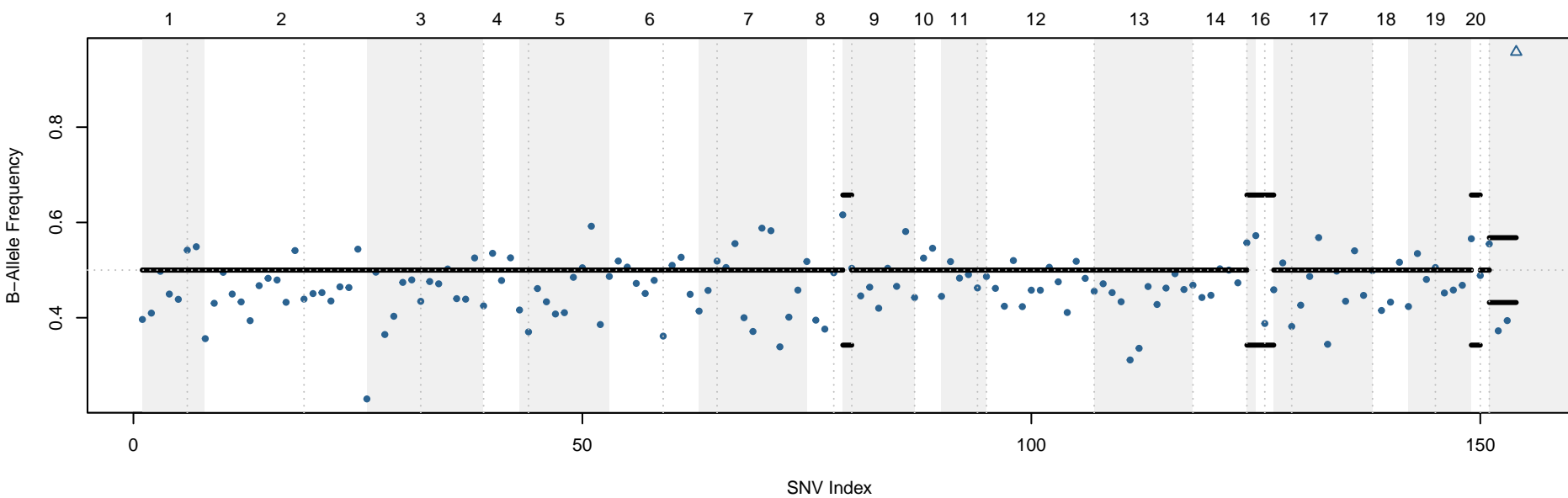


Purity: 0.46 Tumor ploidy: 4.043

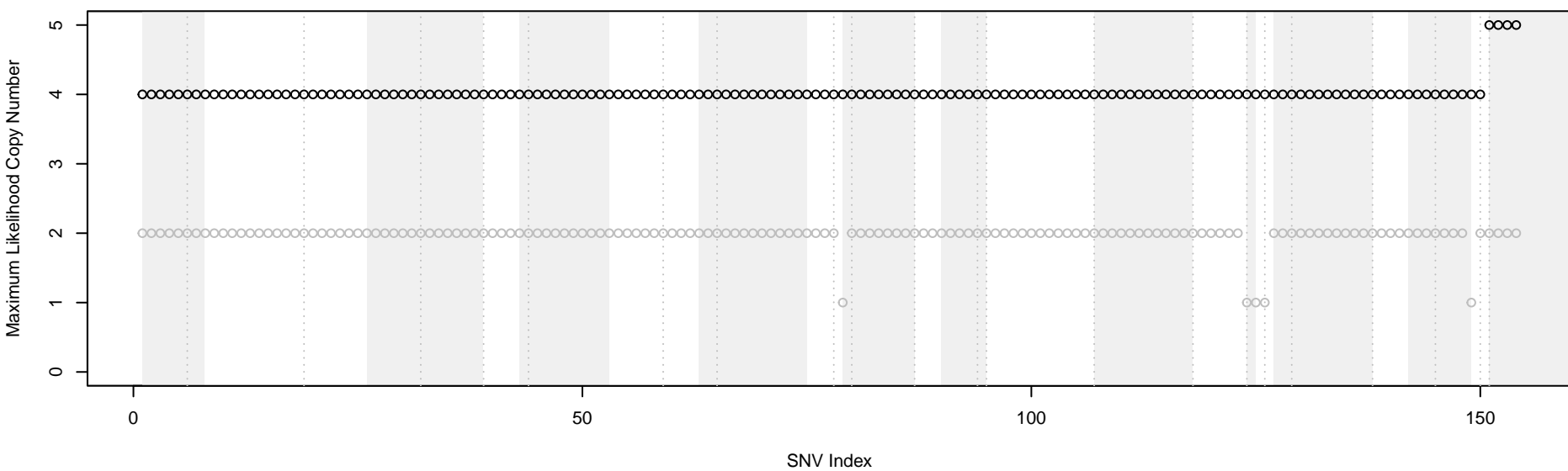
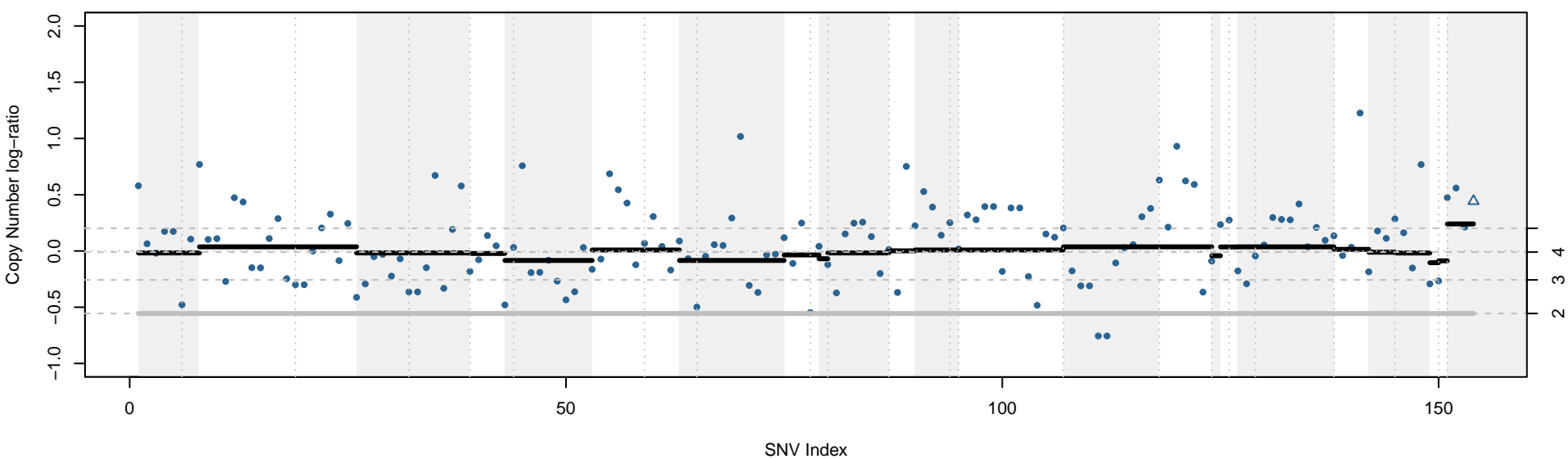


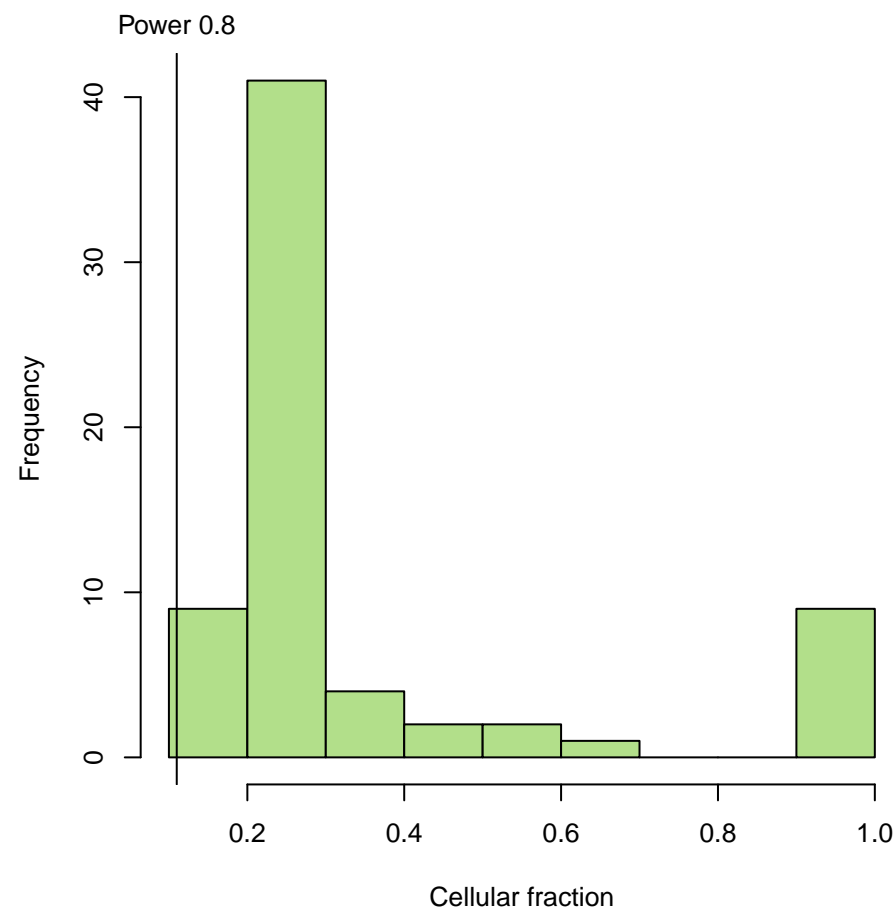
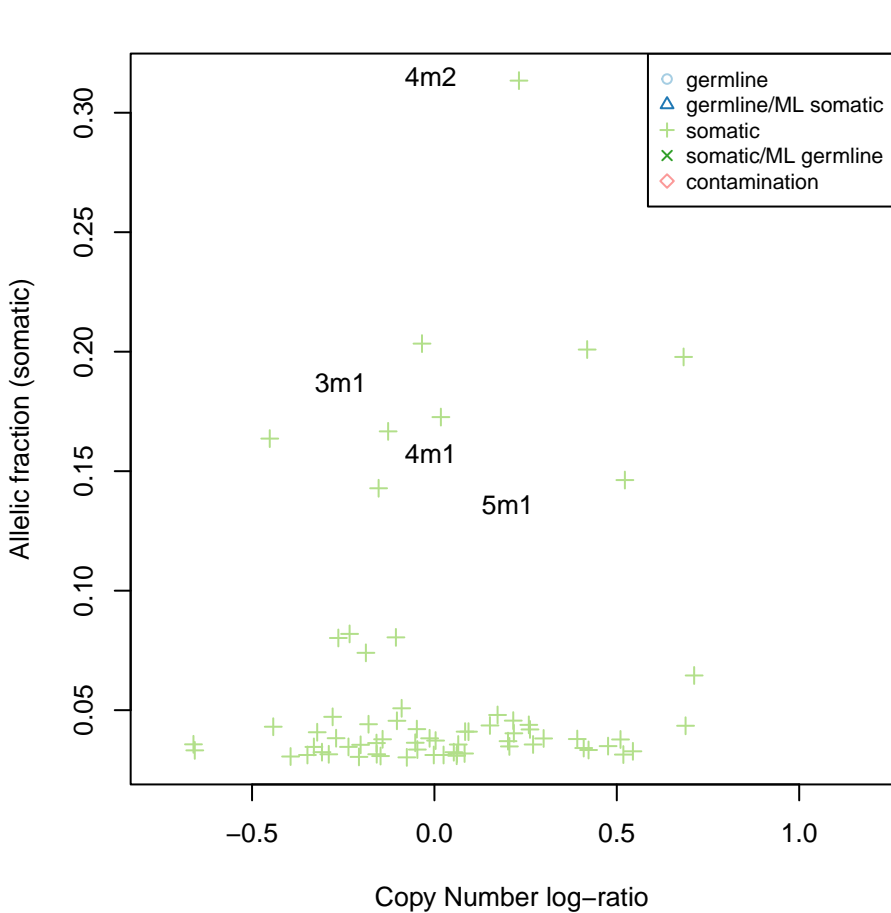
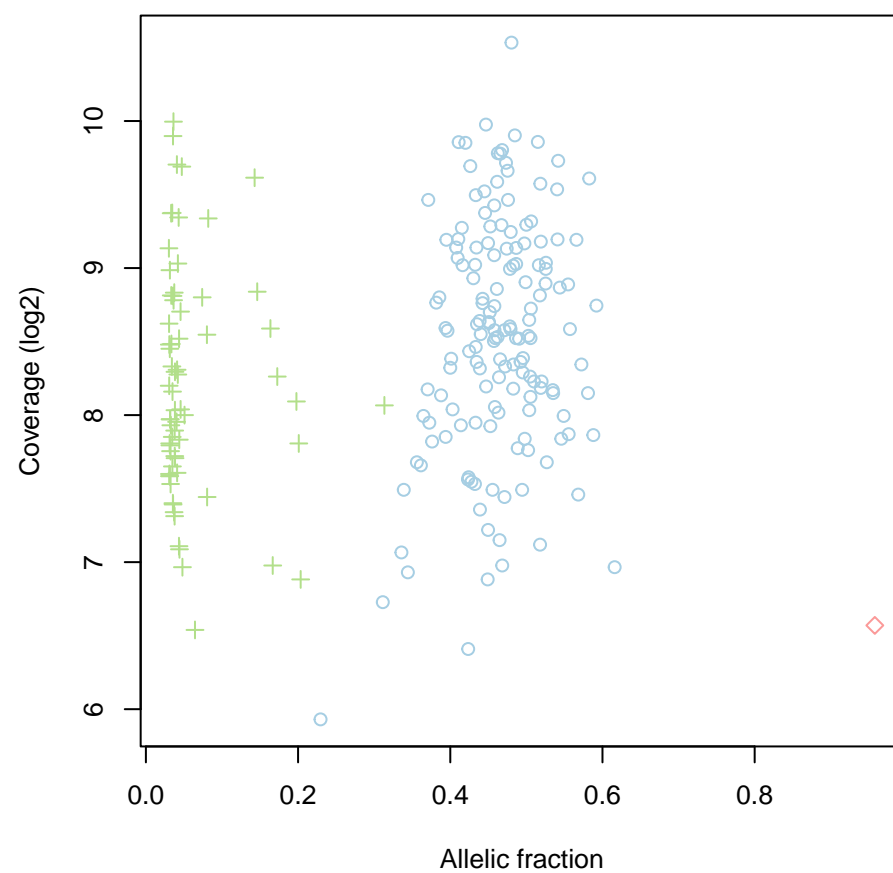
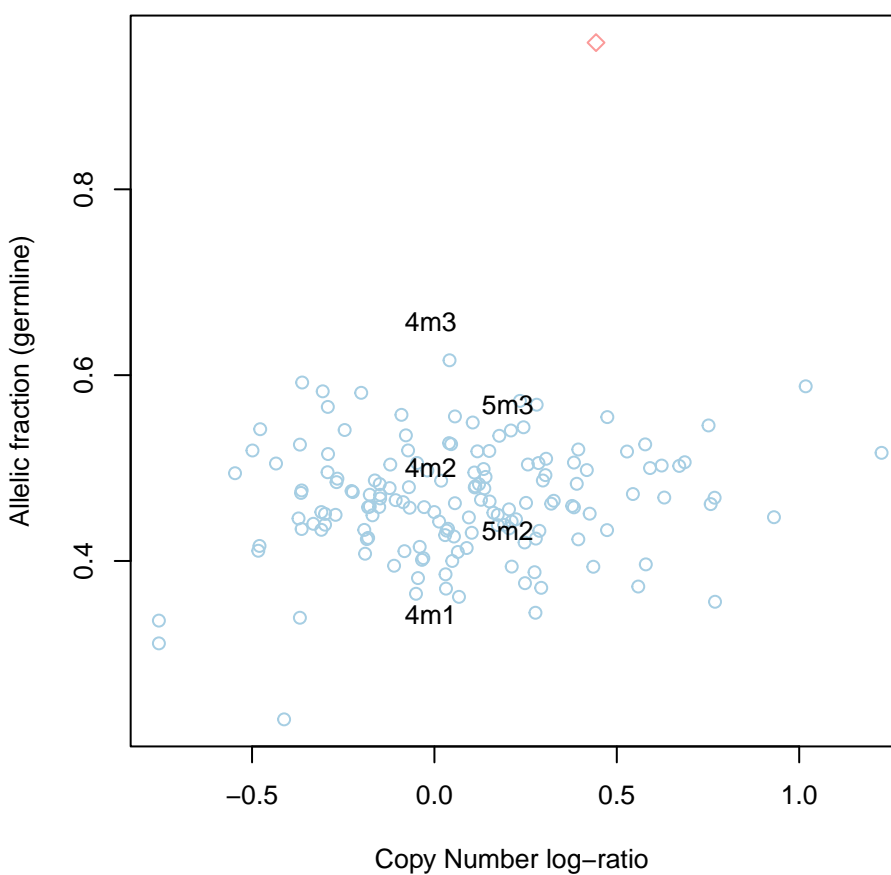


Purity: 0.46 Tumor ploidy: 4.043 SNV log-likelihood: -128.63 GoF: 84.7% Mean coverage: 401;62

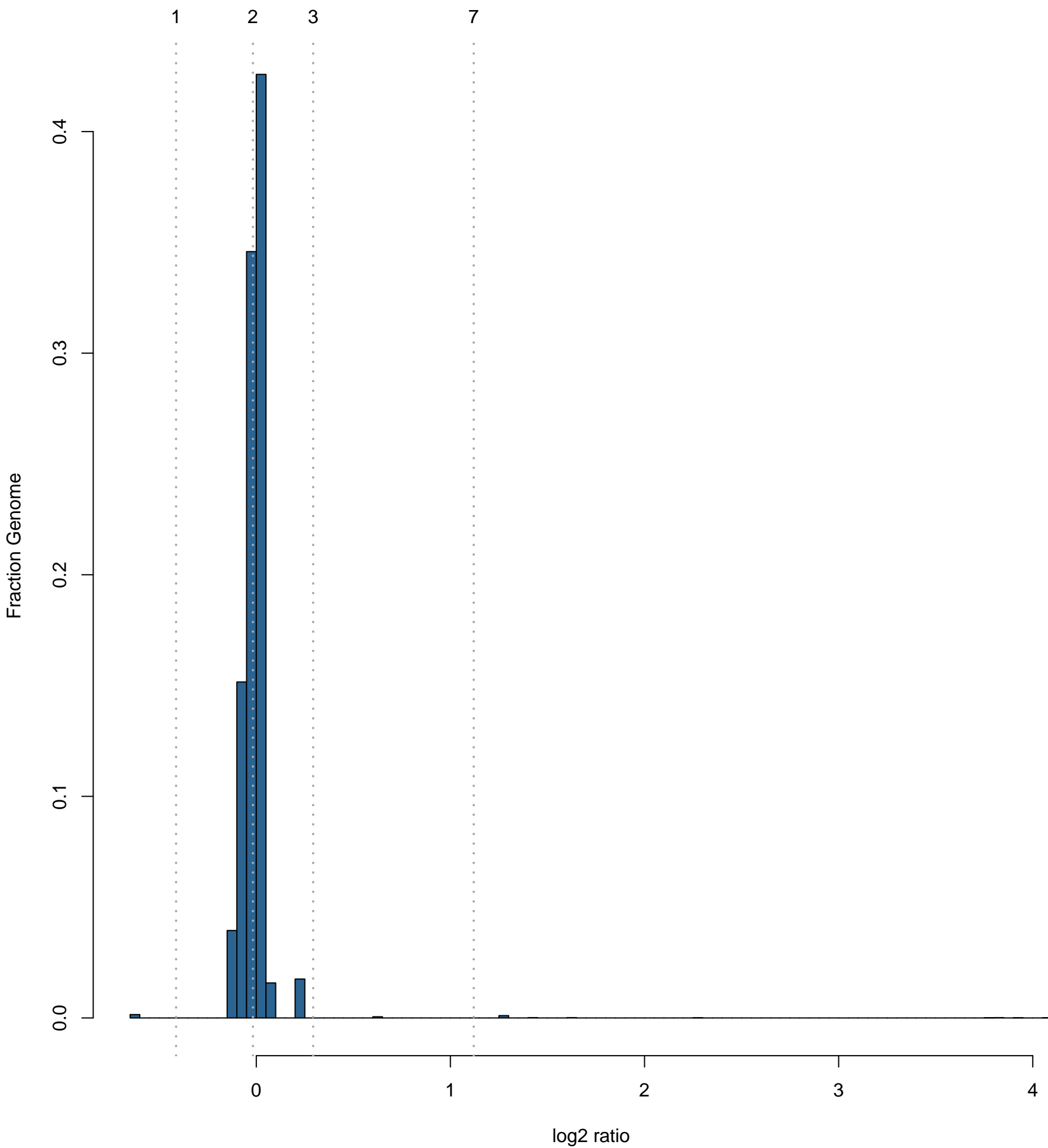


SCNA-fit log-likelihood: -15101.11

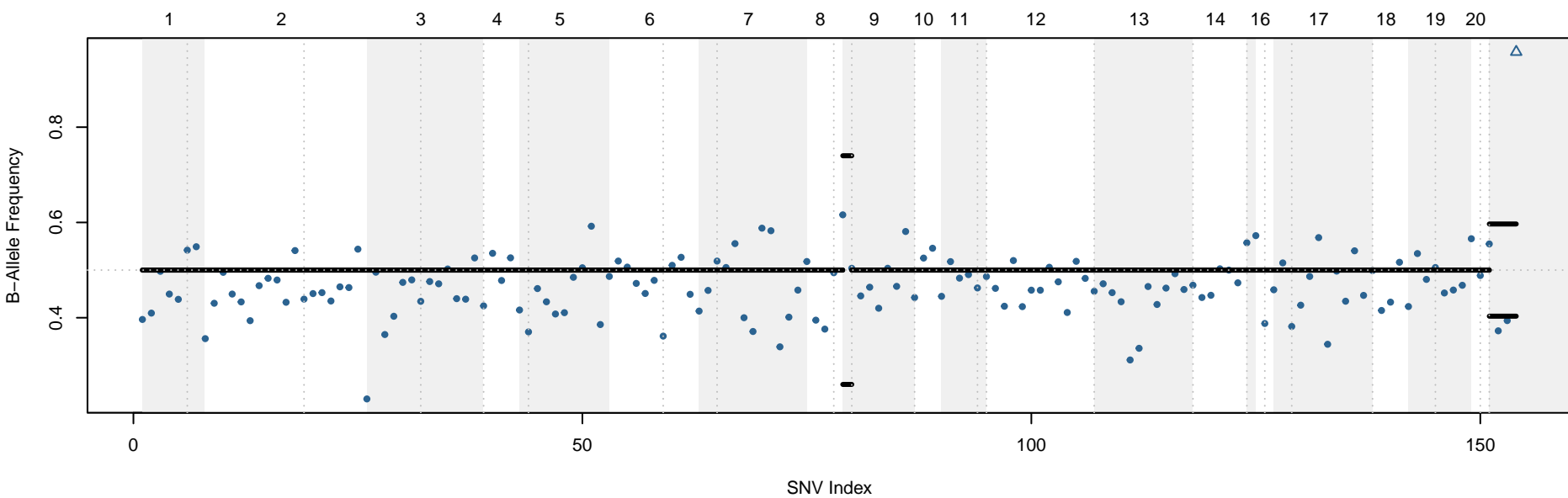




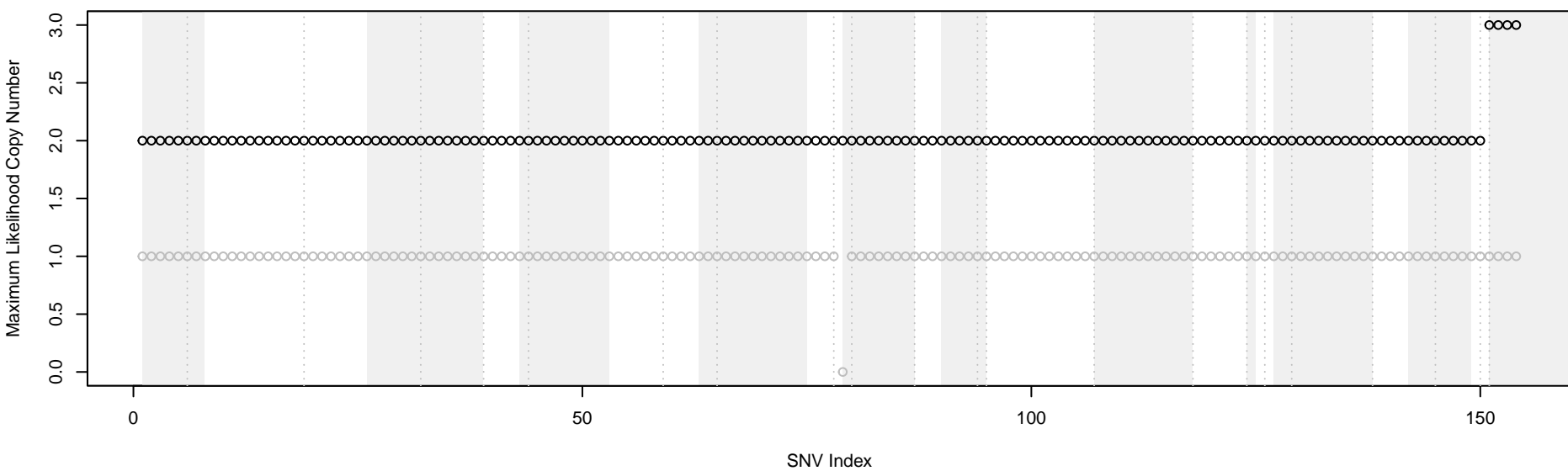
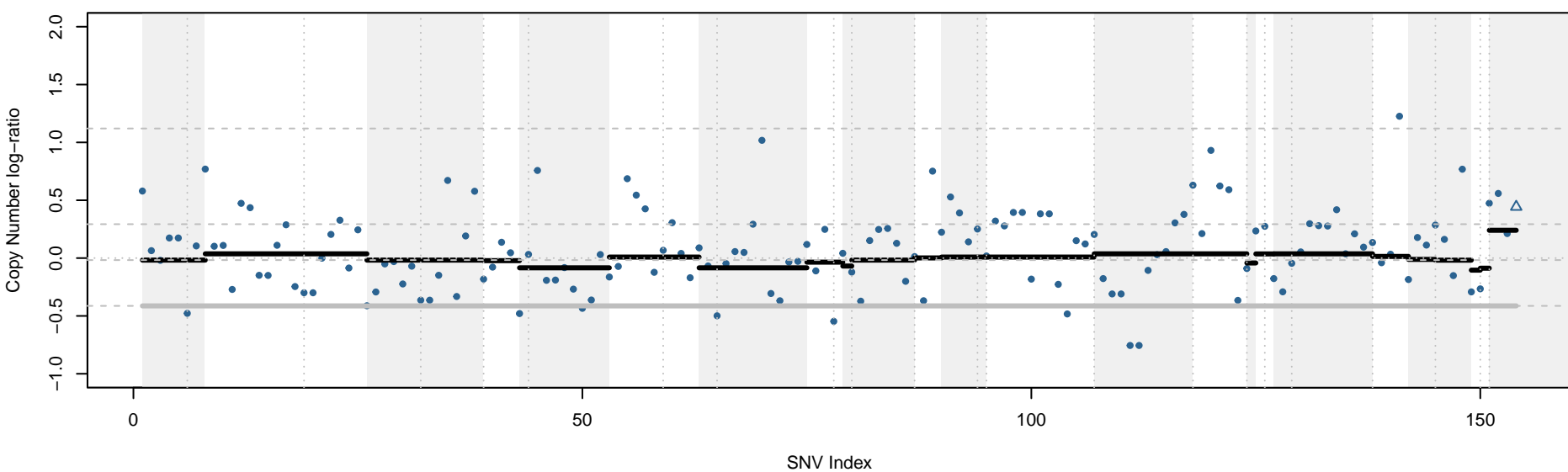
Purity: 0.48 Tumor ploidy: 2.05

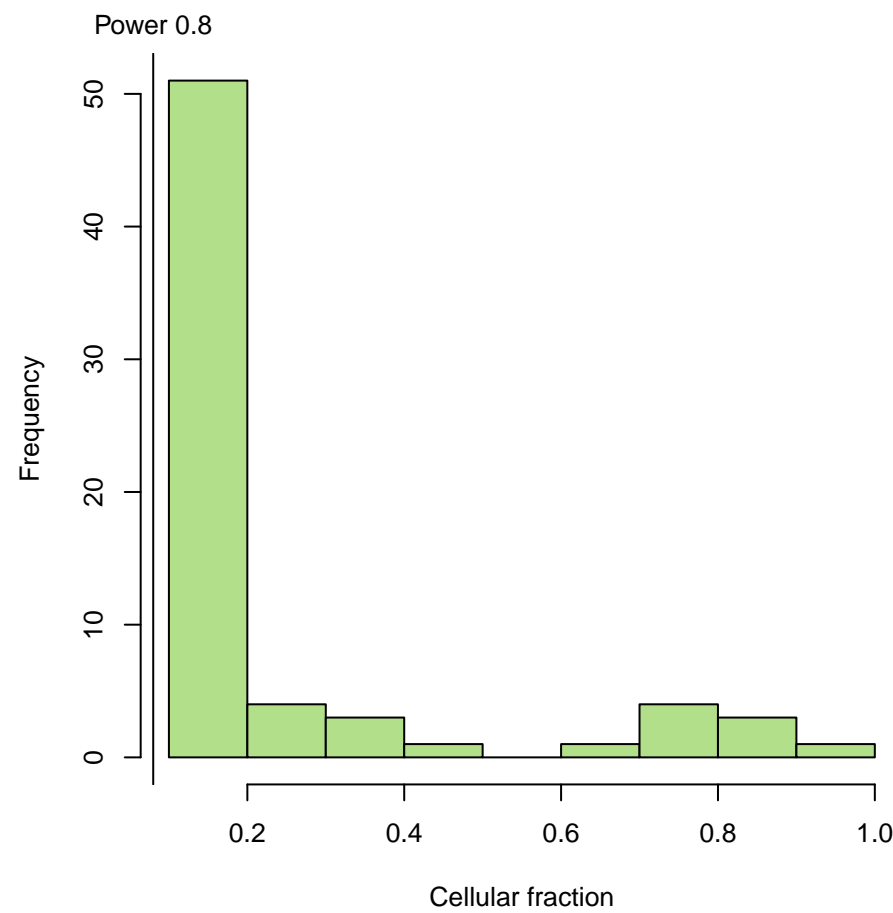
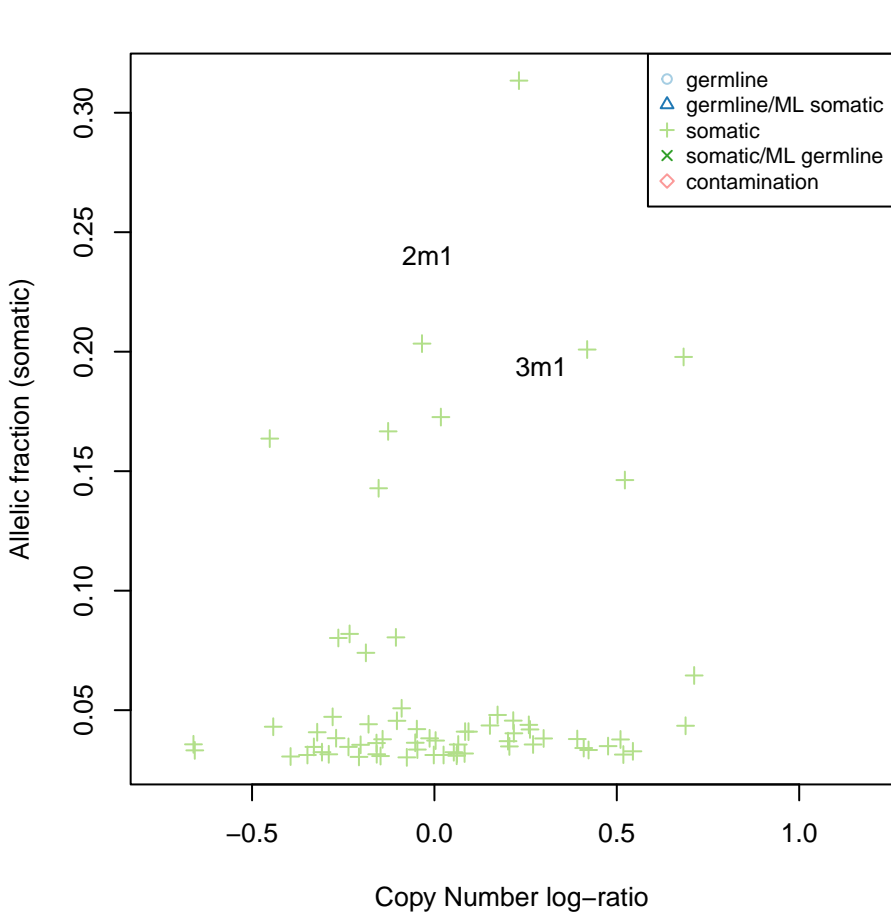
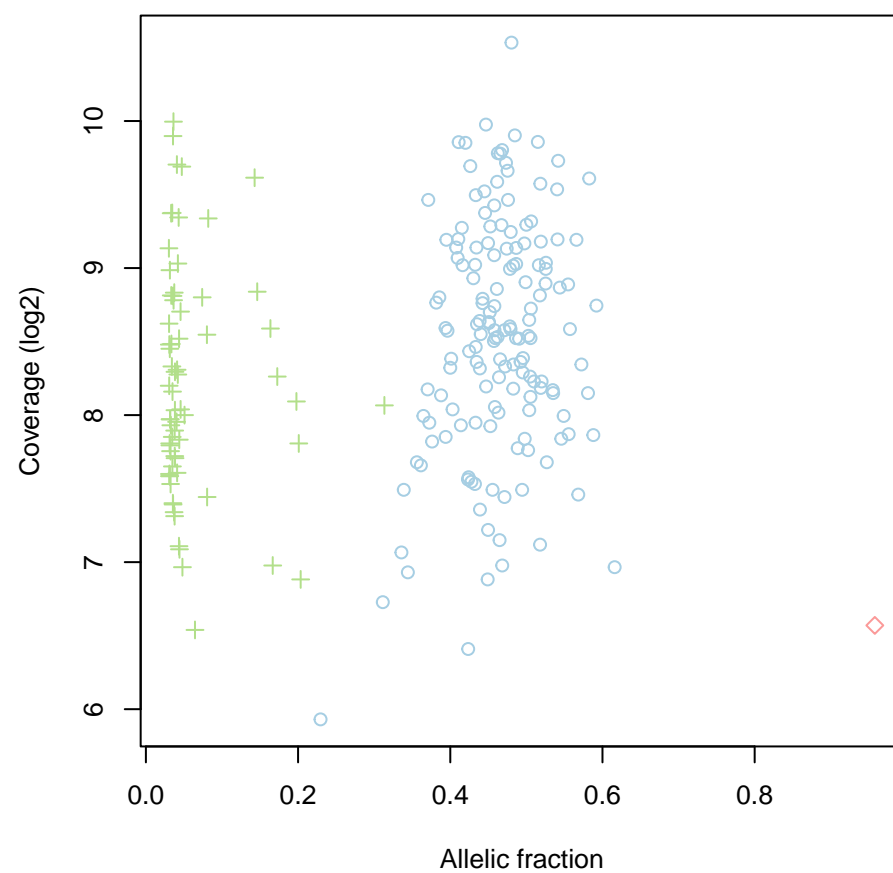
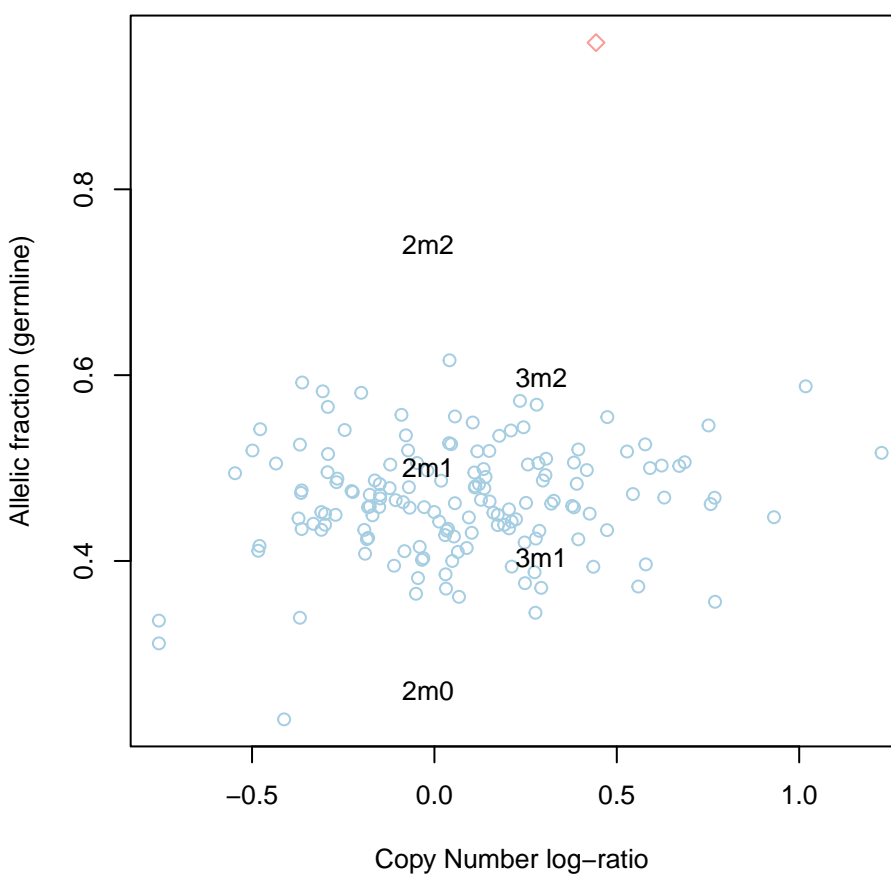


Purity: 0.48 Tumor ploidy: 2.05 SNV log-likelihood: -155.12 GoF: 68.1% Mean coverage: 401;62

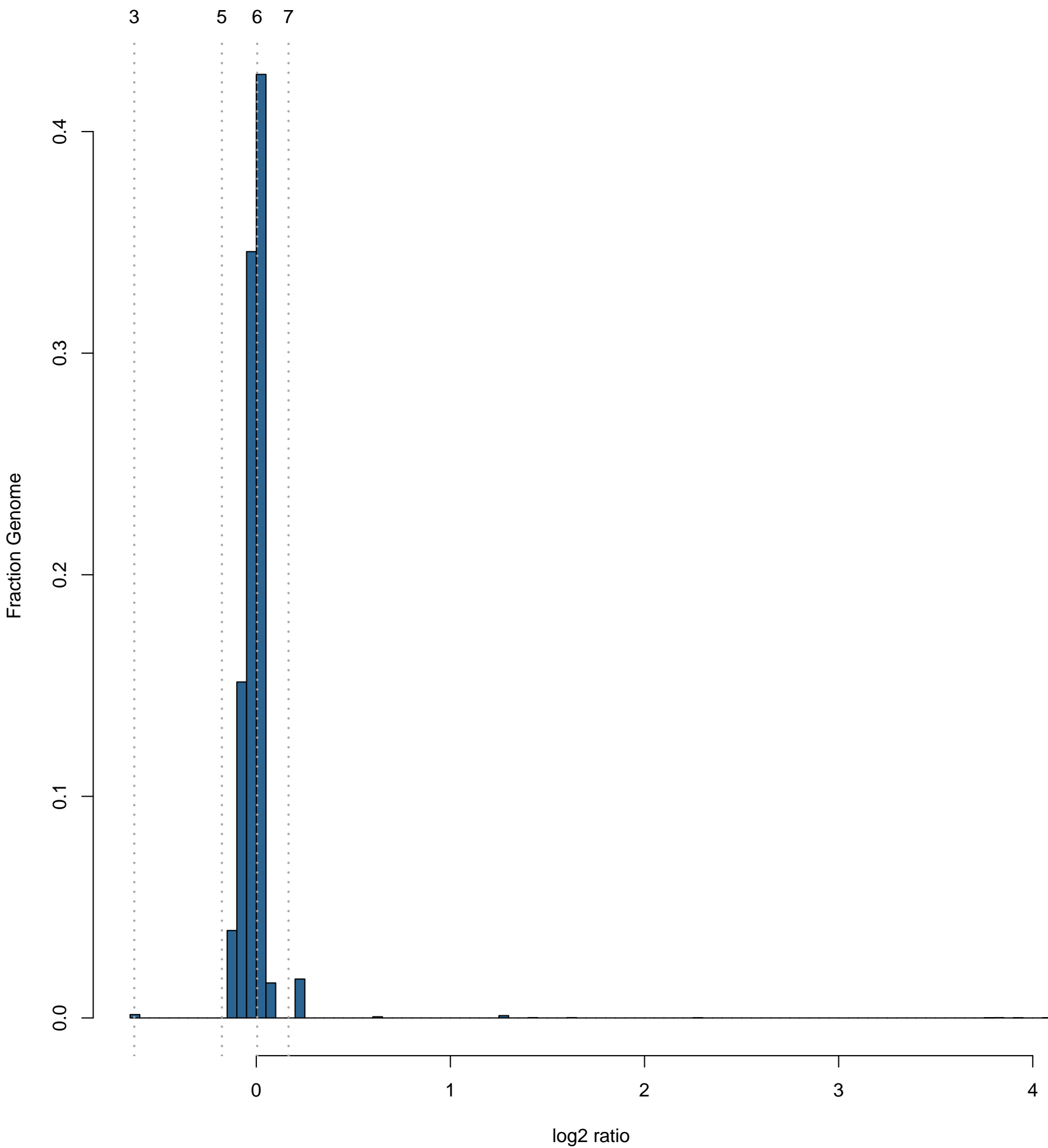


SCNA-fit log-likelihood: -15090.72

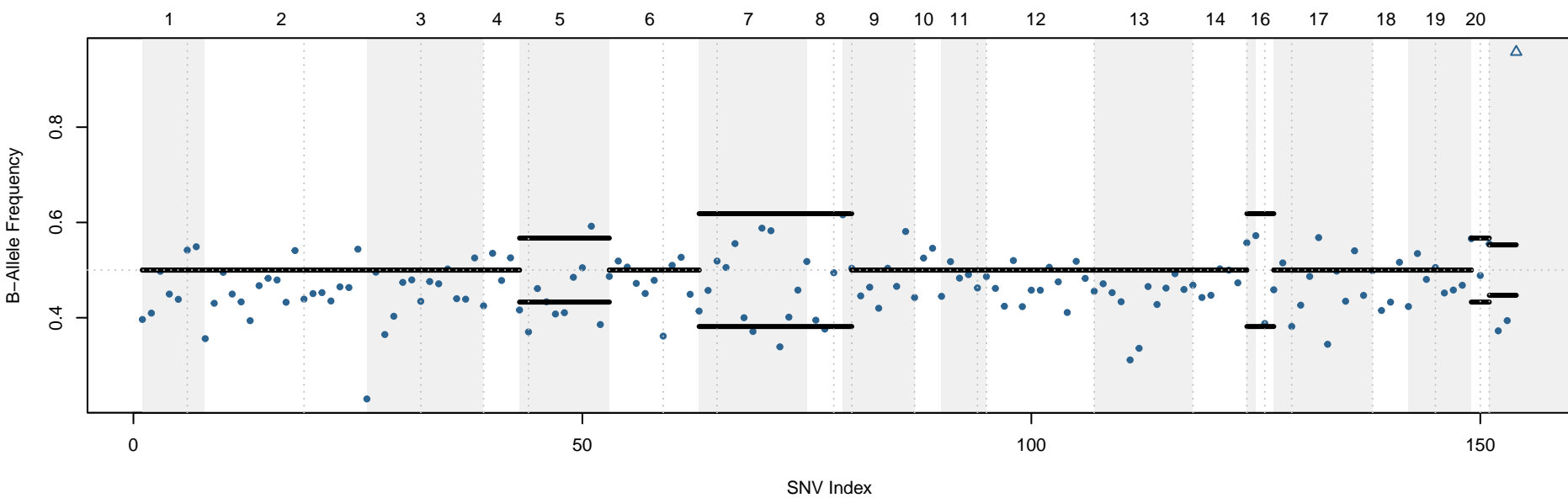




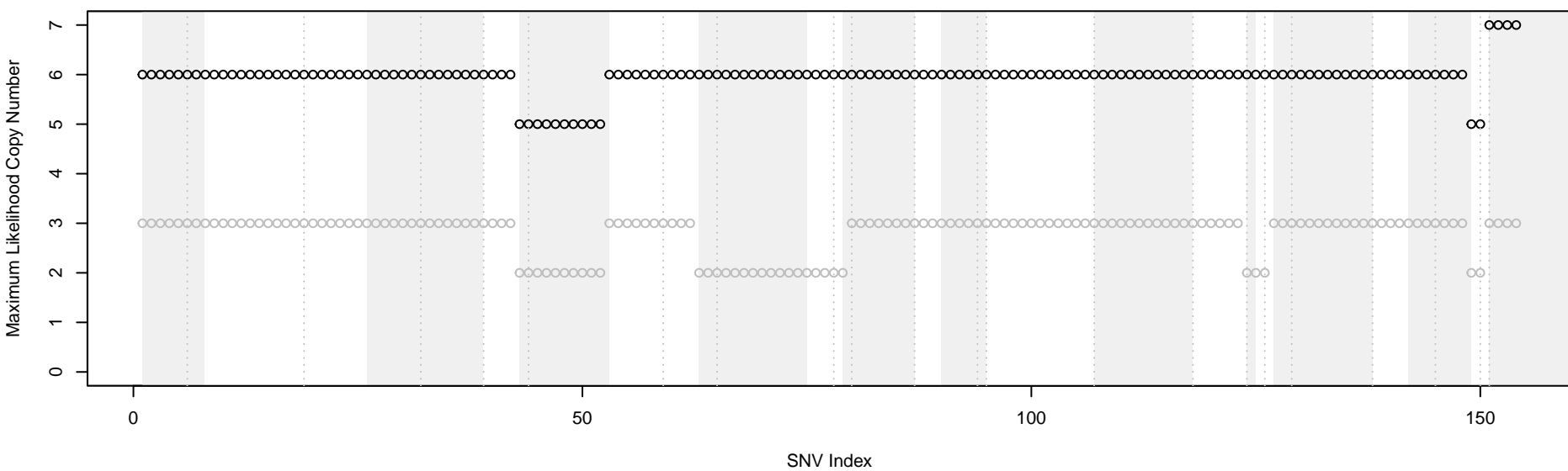
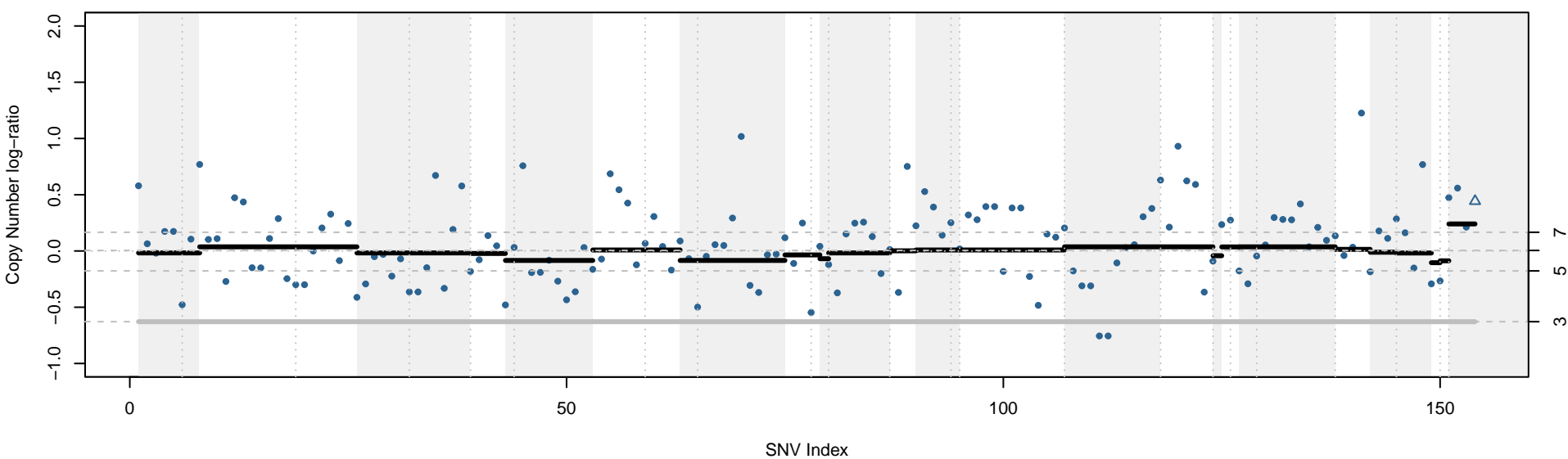
Purity: 0.45 Tumor ploidy: 5.972

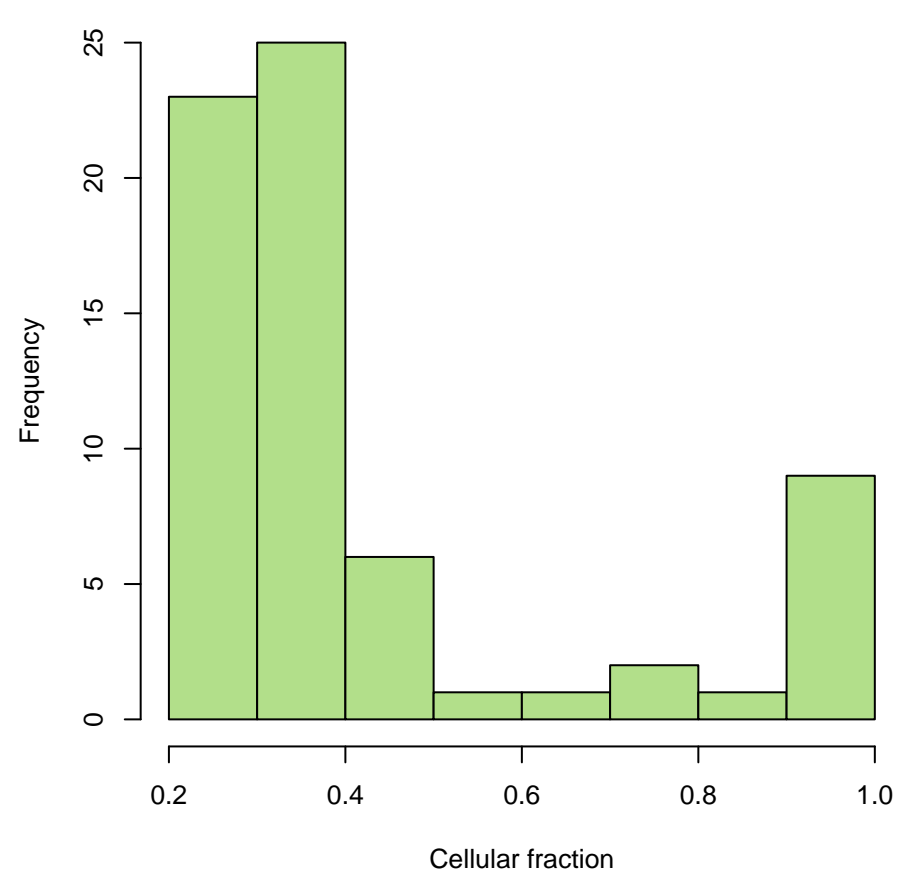
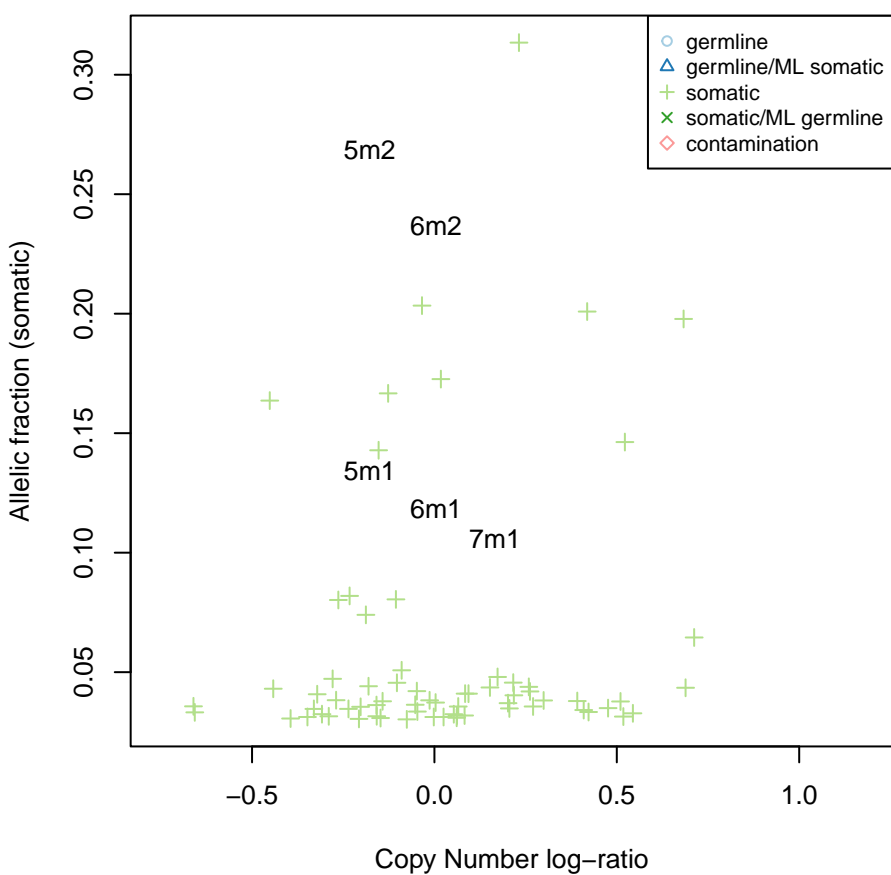
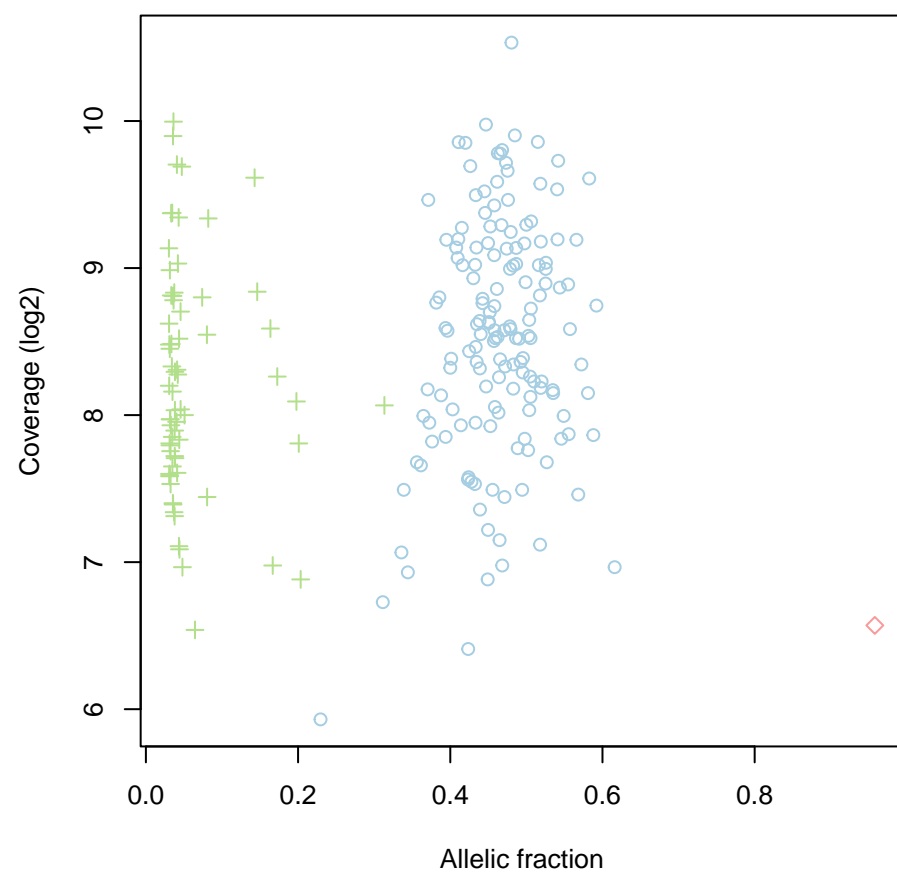
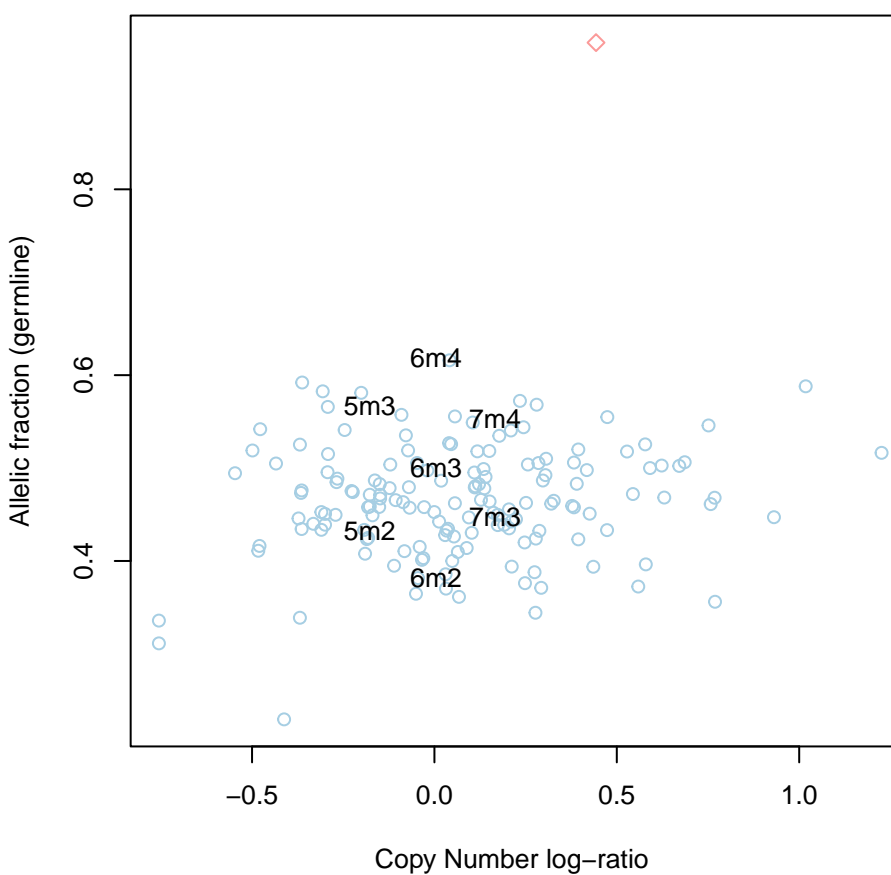


Purity: 0.45 Tumor ploidy: 5.972 SNV log-likelihood: -150.91 GoF: 90.3% Mean coverage: 401;62



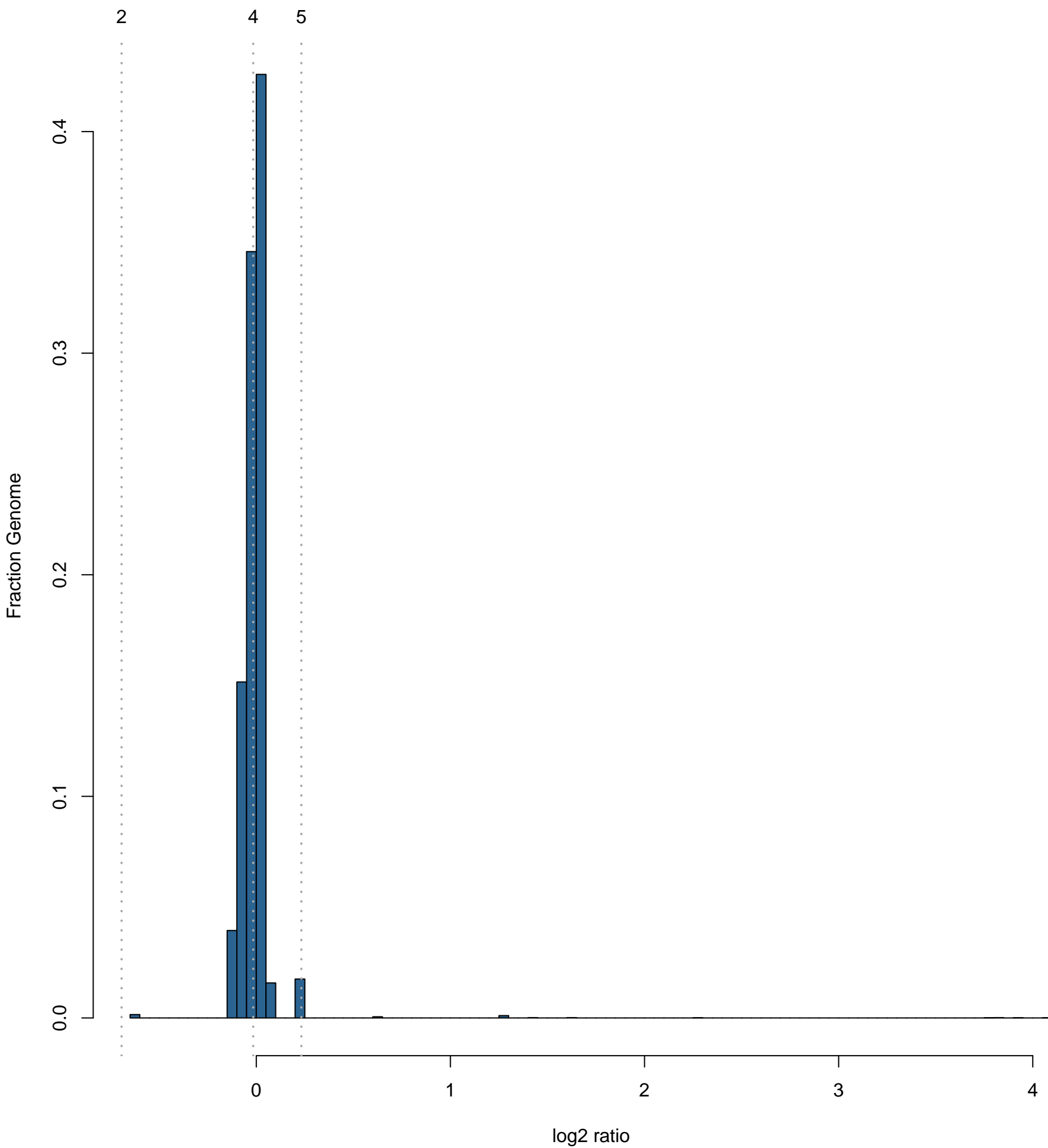
SCNA-fit log-likelihood: -15081.92



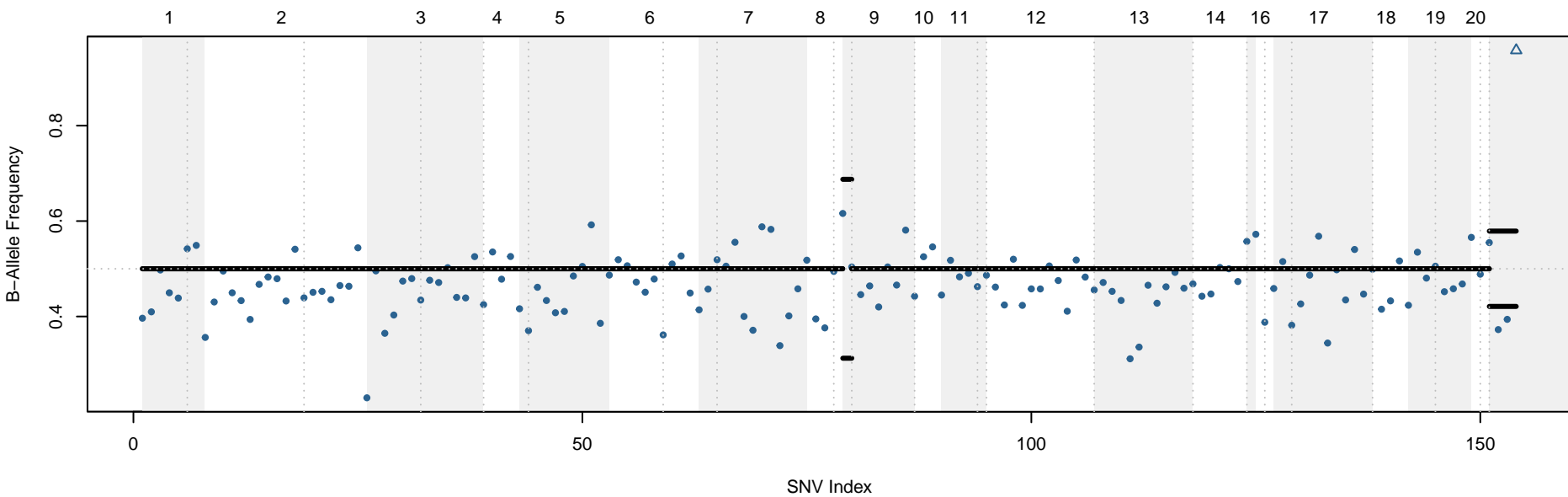




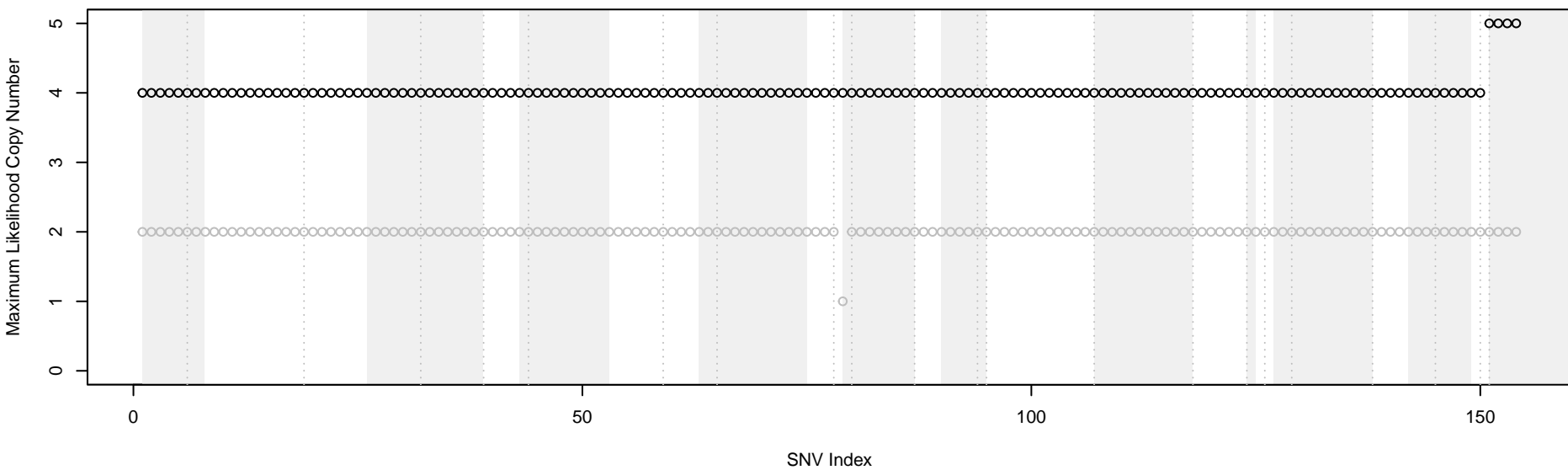
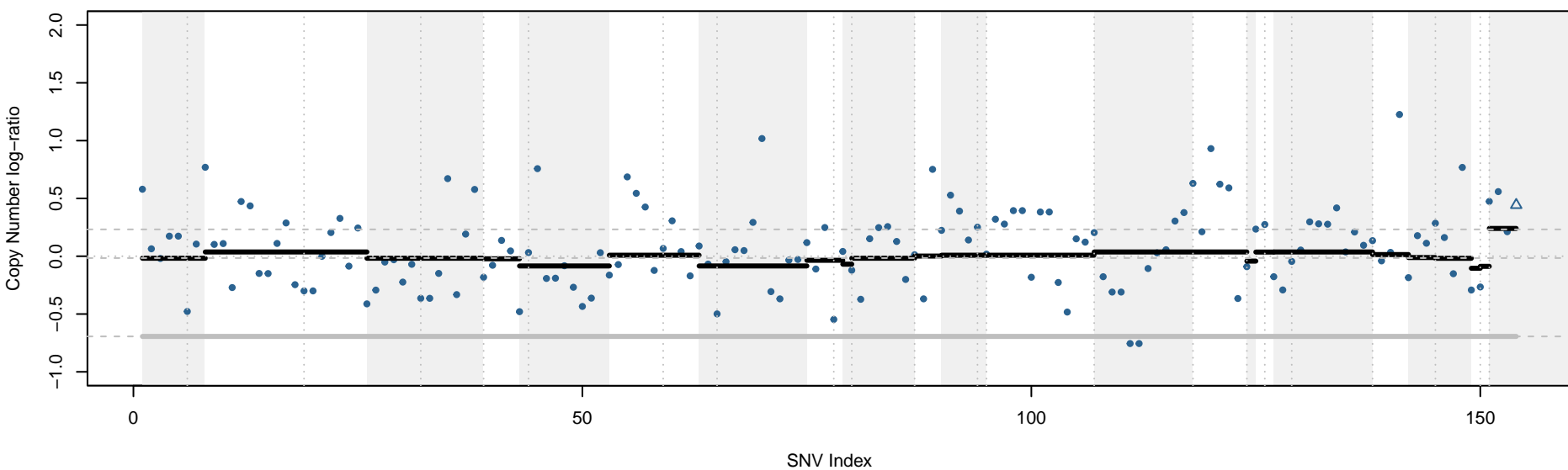
Purity: 0.6 Tumor ploidy: 4.058

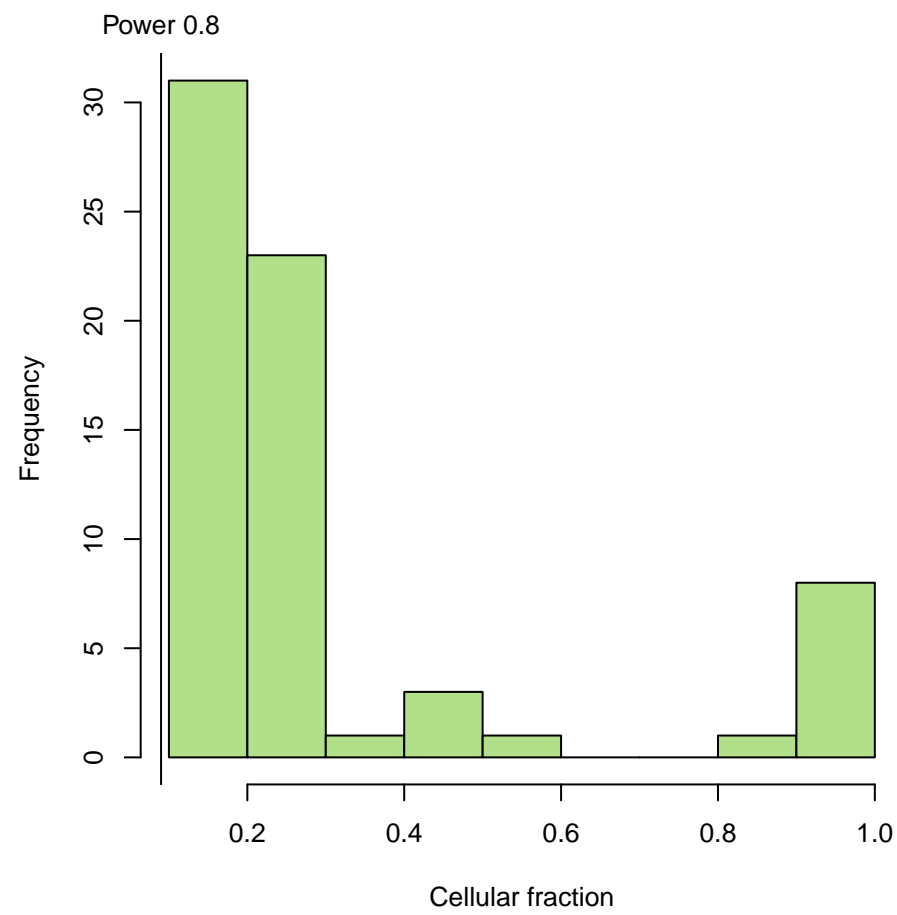
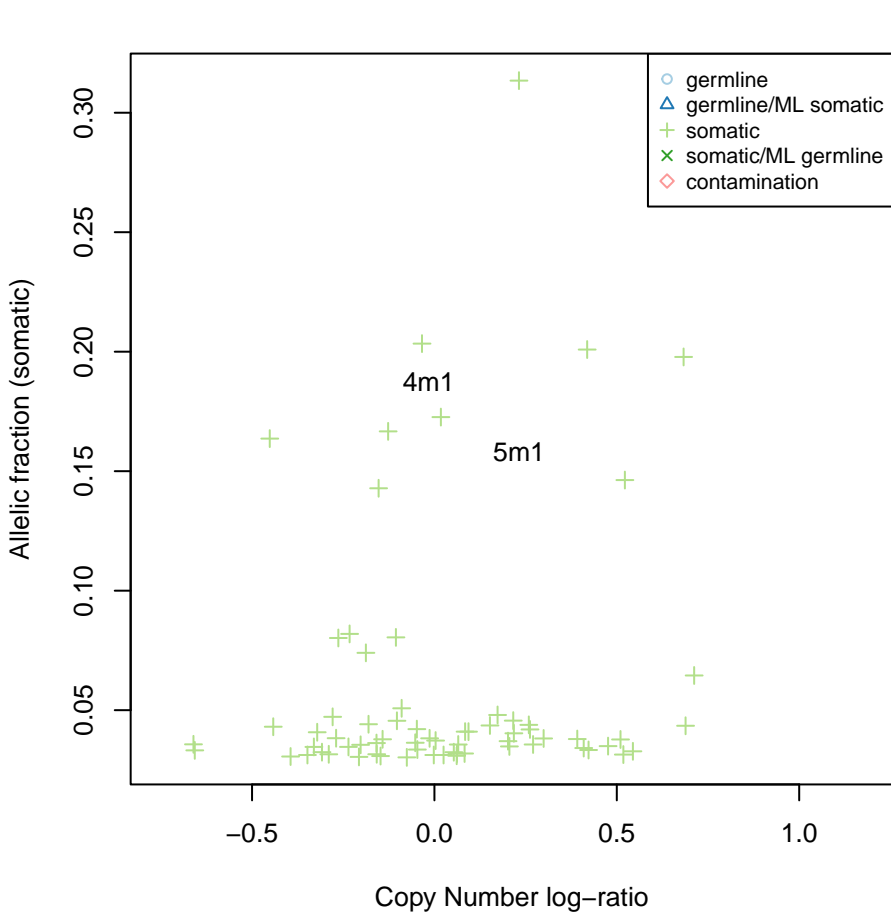
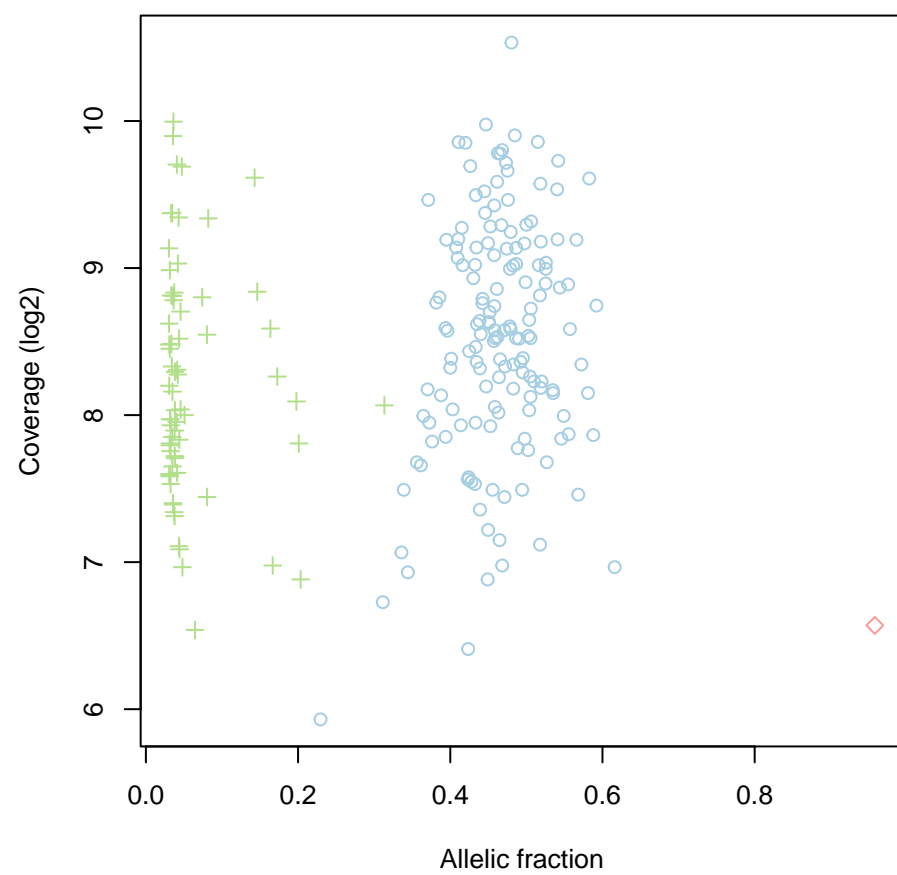
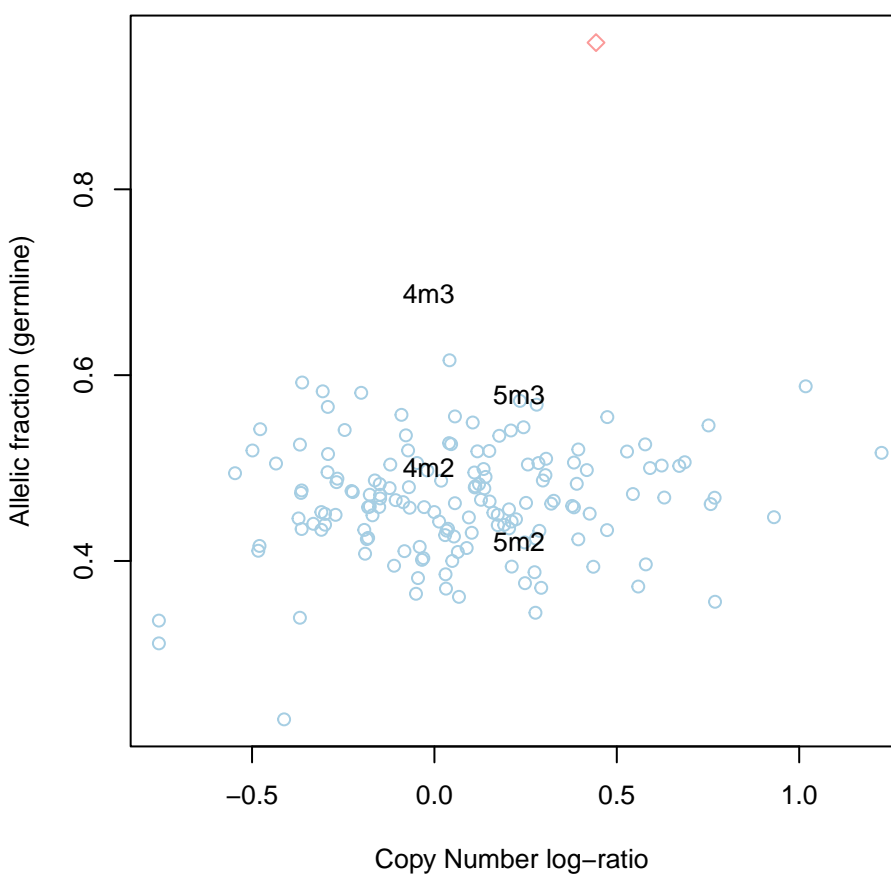


Purity: 0.6 Tumor ploidy: 4.058 SNV log-likelihood: -168.7 GoF: 79.5% Mean coverage: 401;62

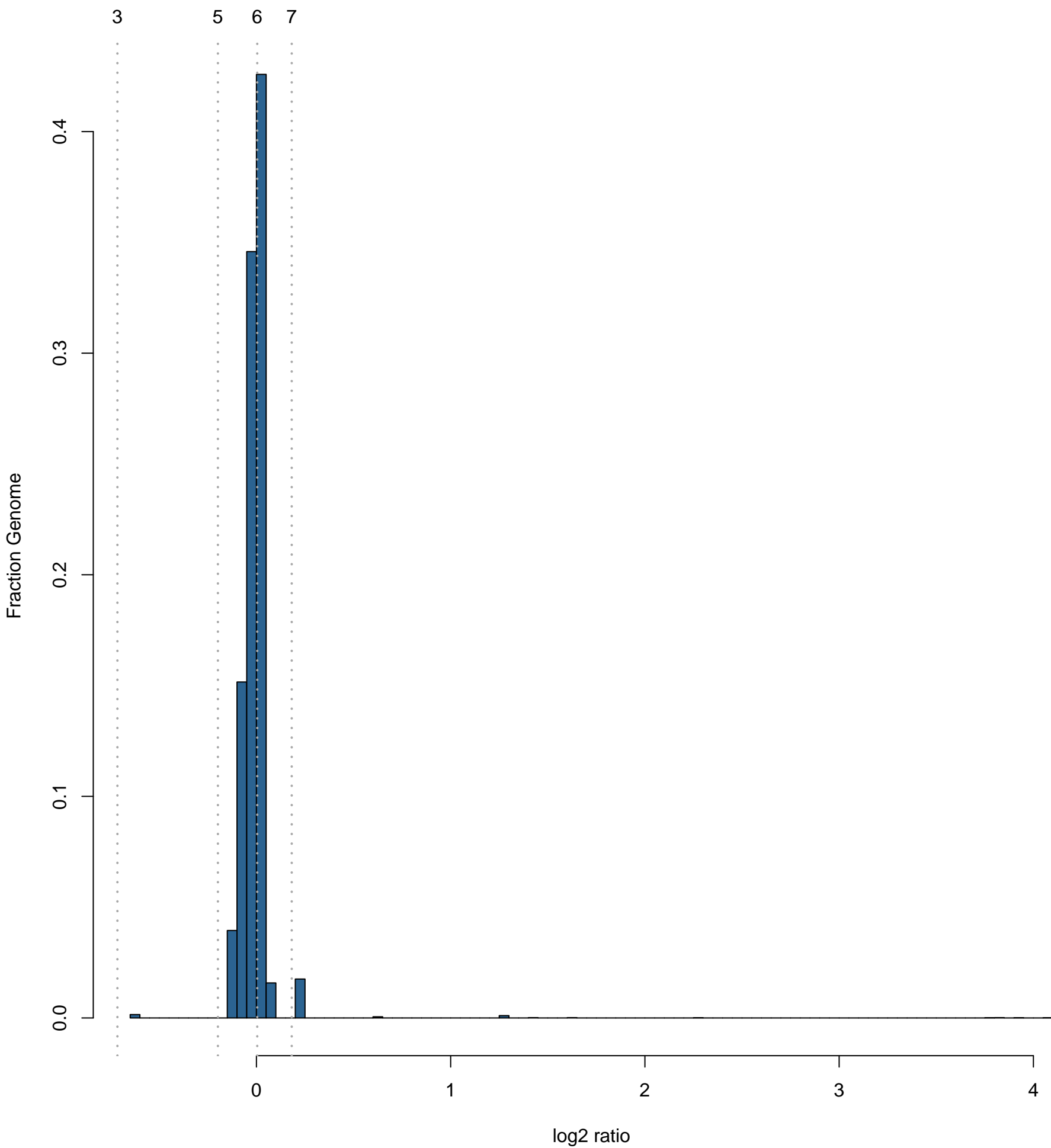


SCNA-fit log-likelihood: -15108.31

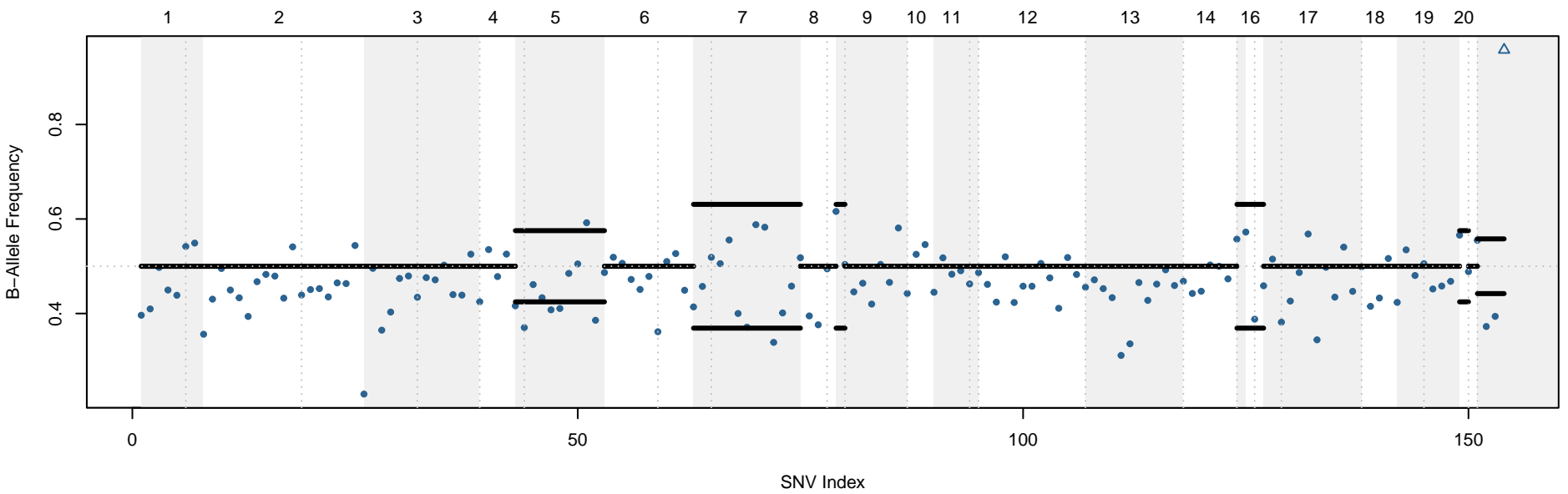




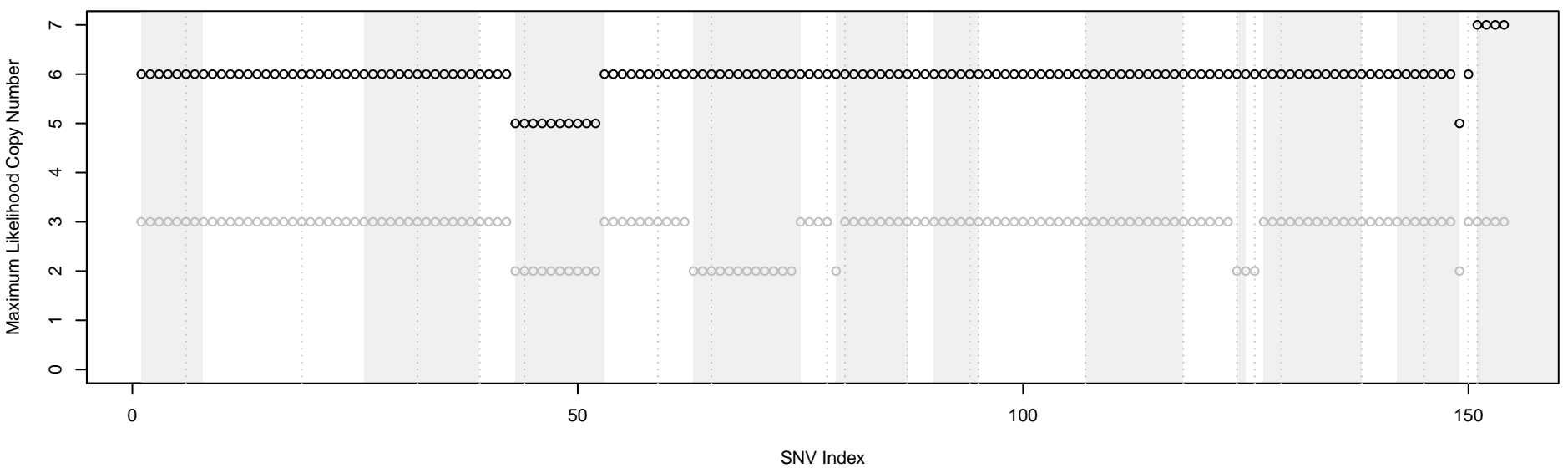
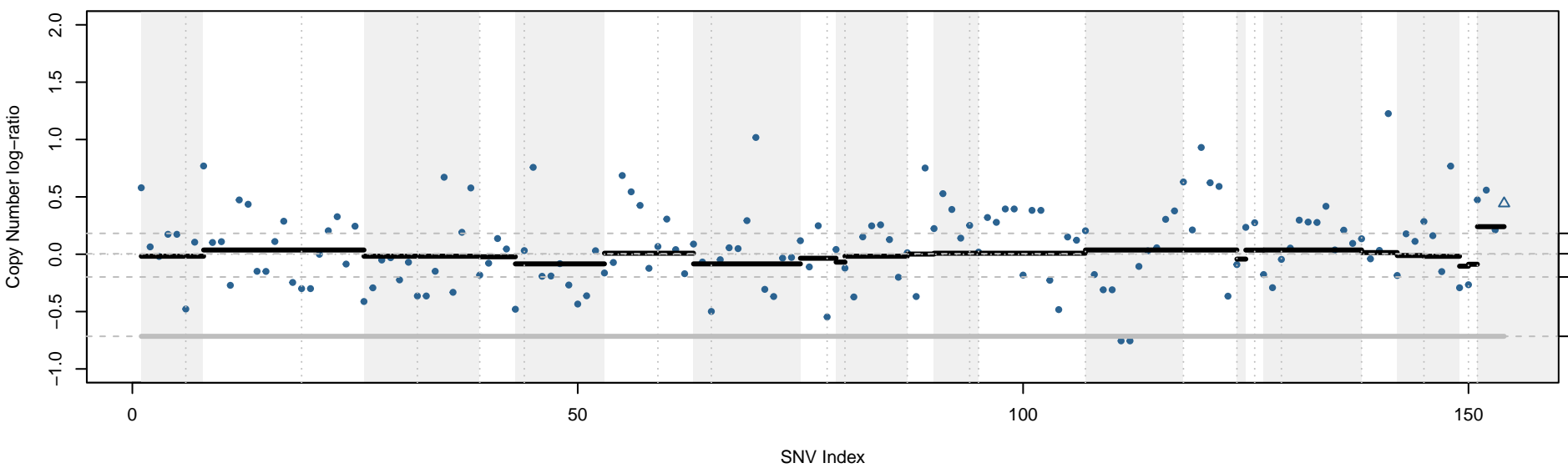
Purity: 0.55 Tumor ploidy: 5.978

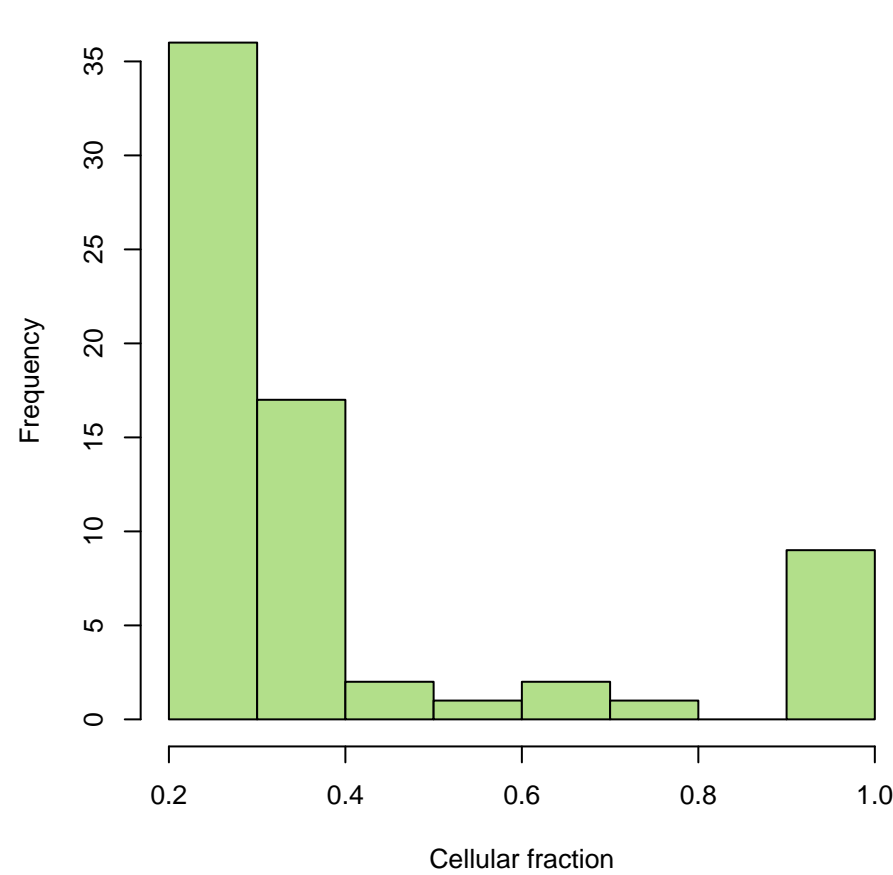
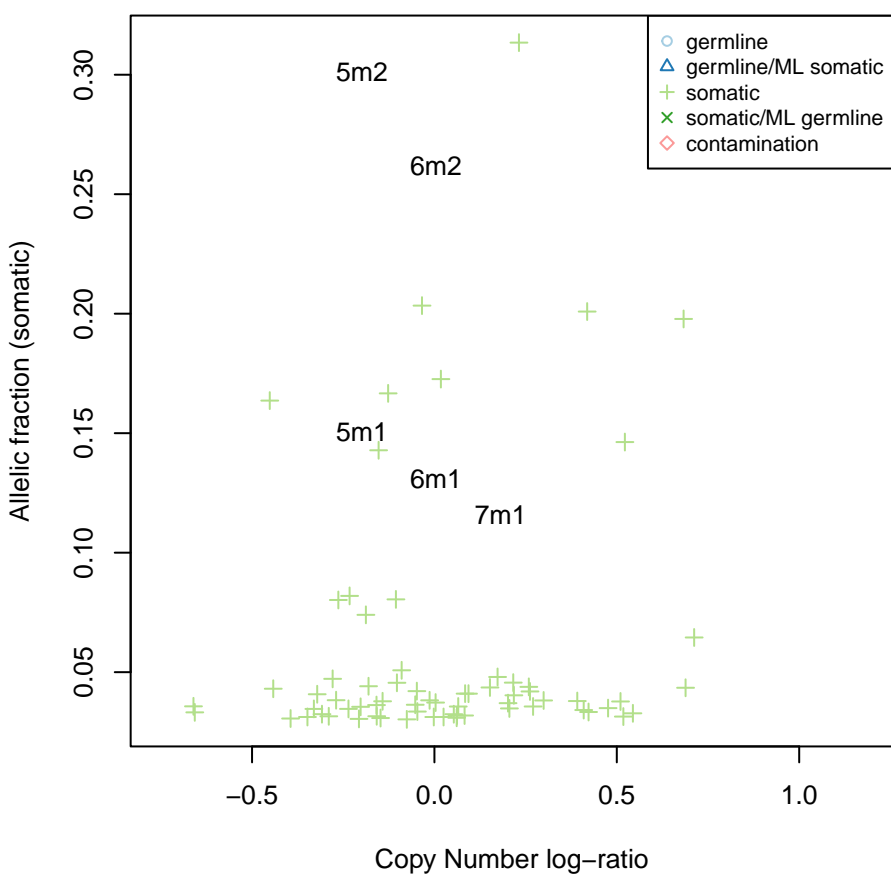
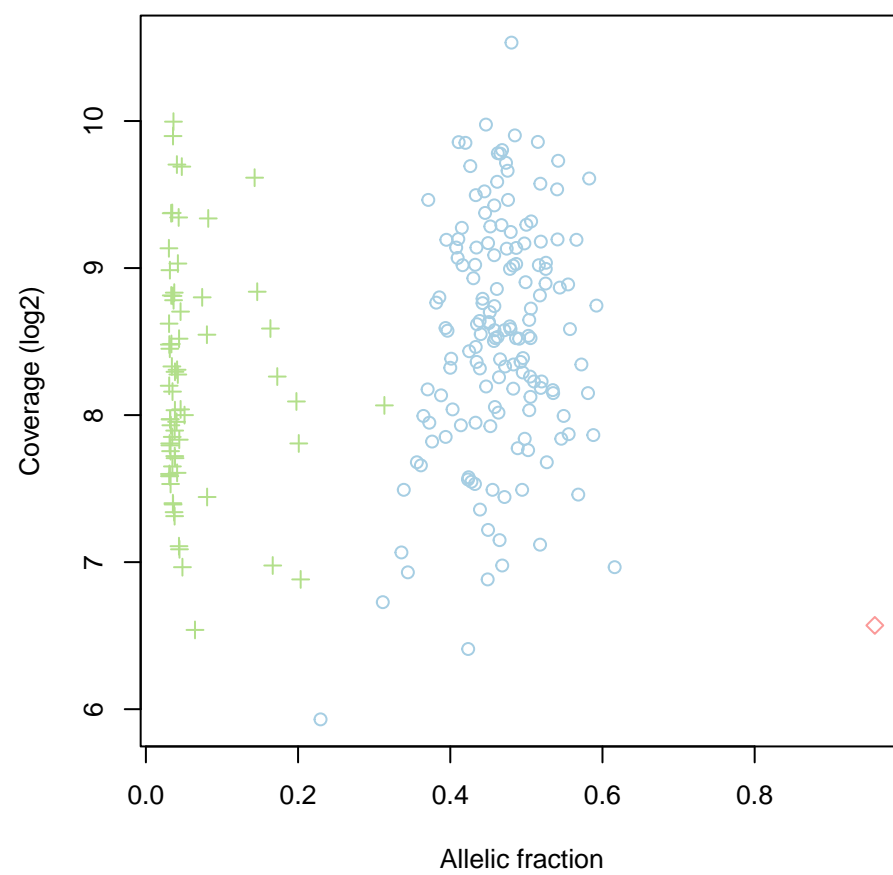
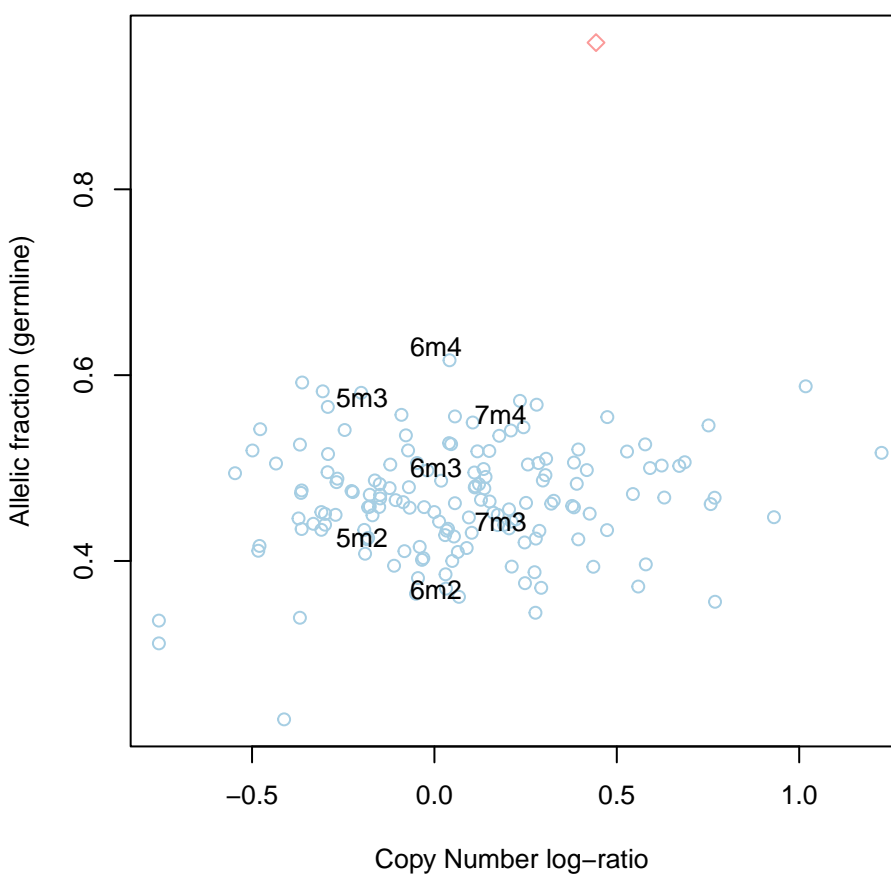


Purity: 0.55 Tumor ploidy: 5.978 SNV log-likelihood: -166.34 GoF: 88.8% Mean coverage: 401;62

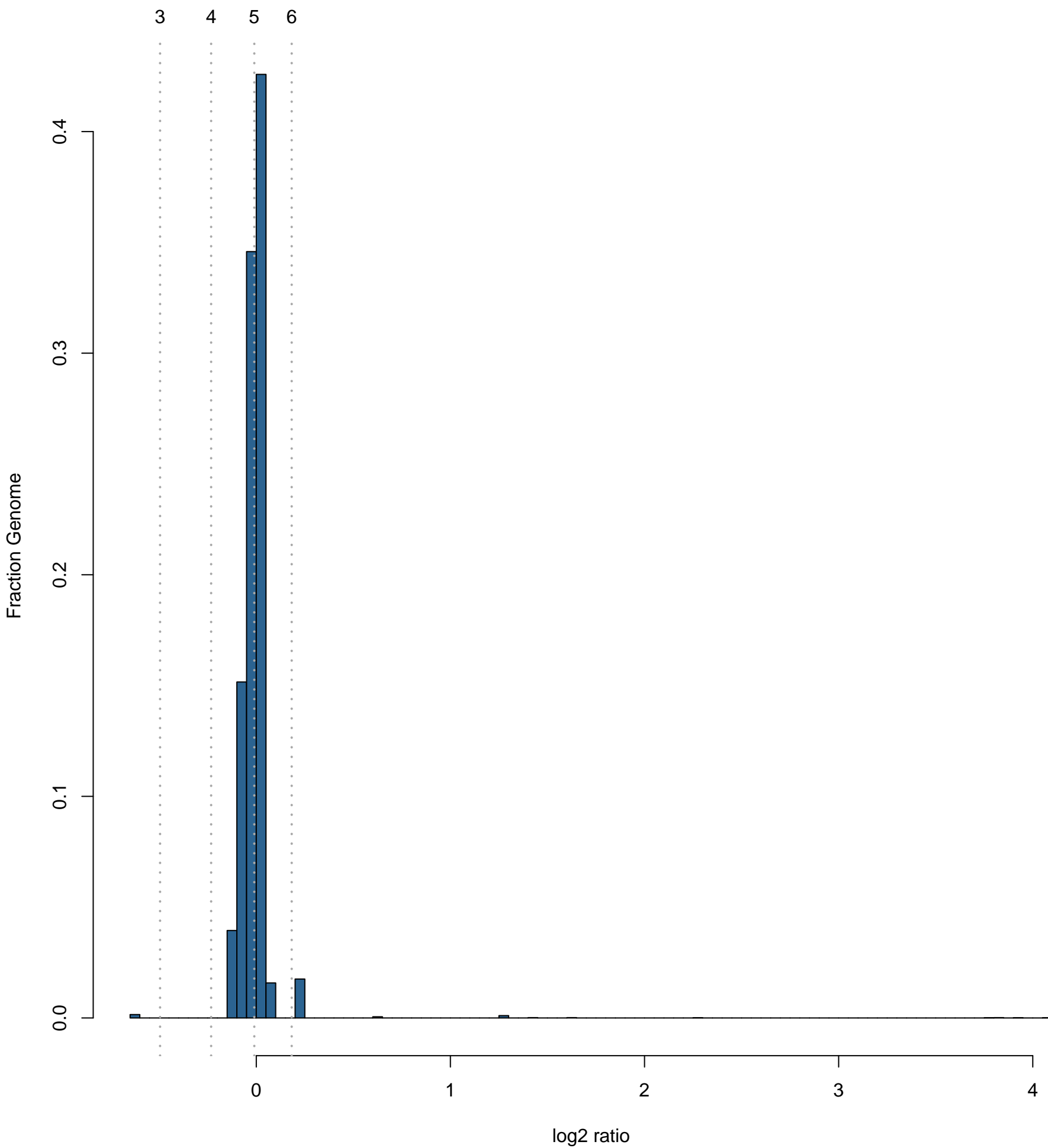


SCNA-fit log-likelihood: -15094.69

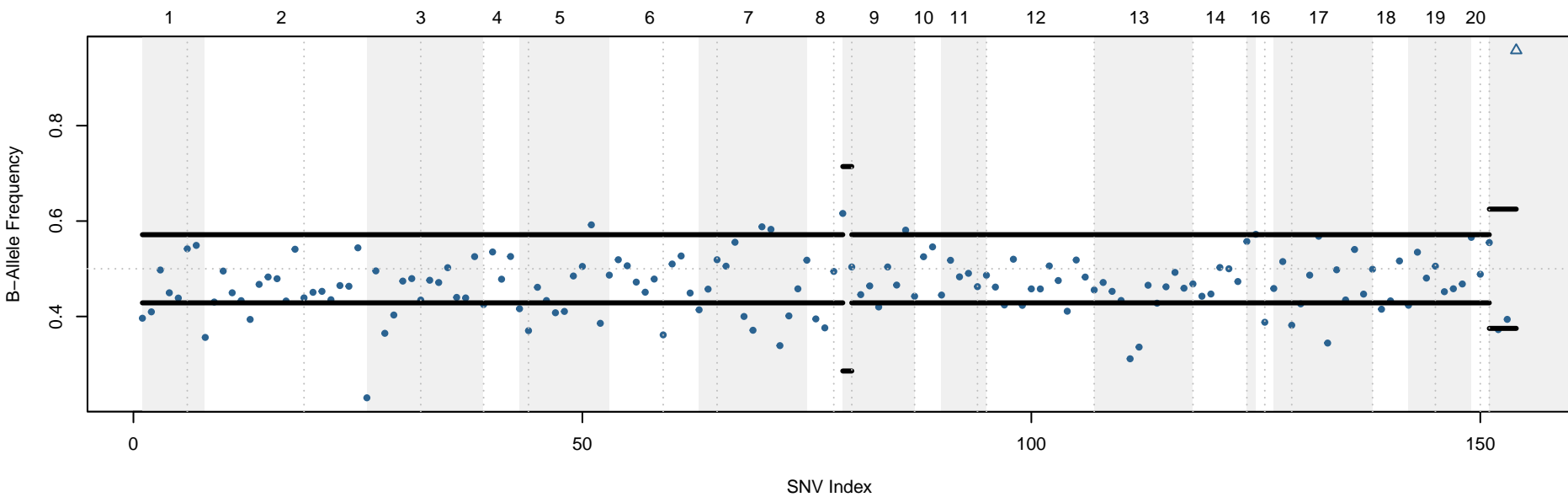




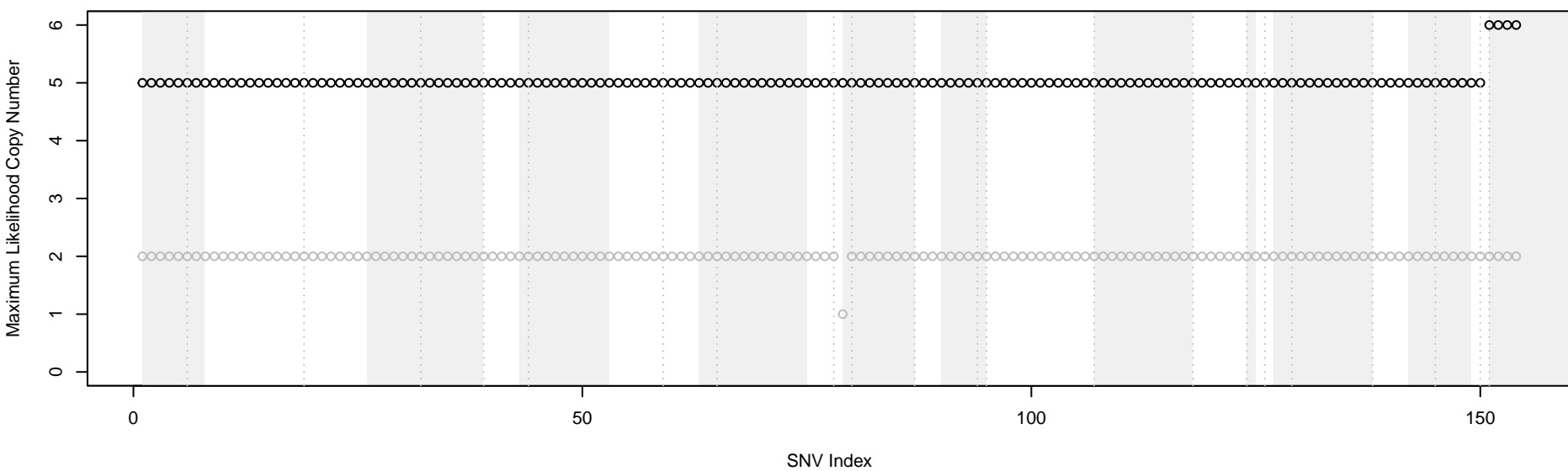
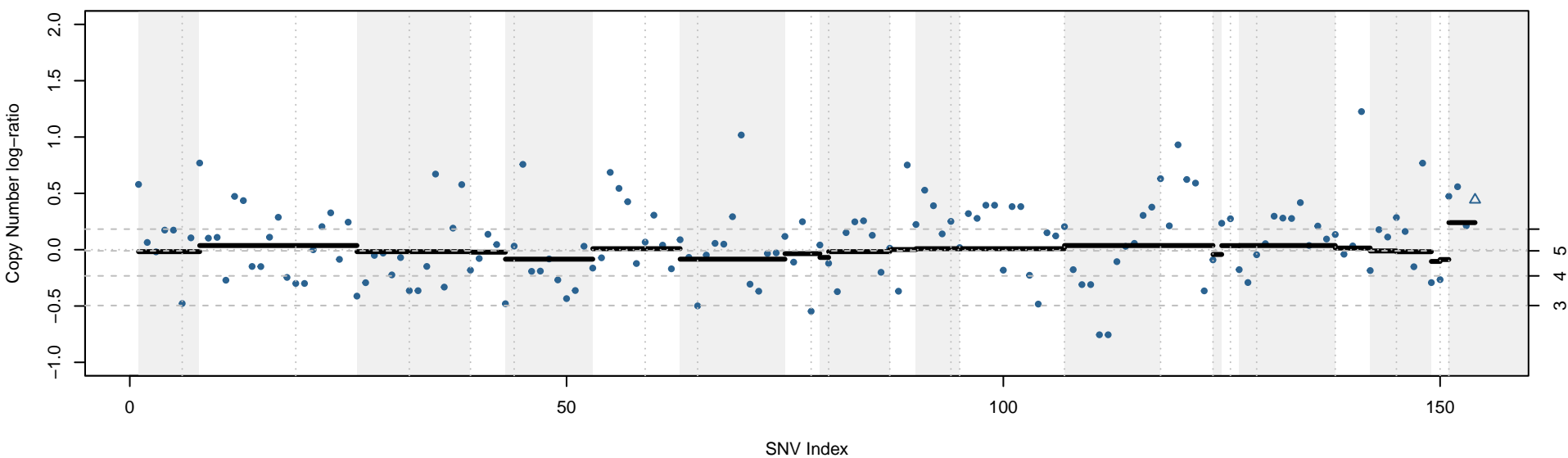
Purity: 0.5 Tumor ploidy: 5.05



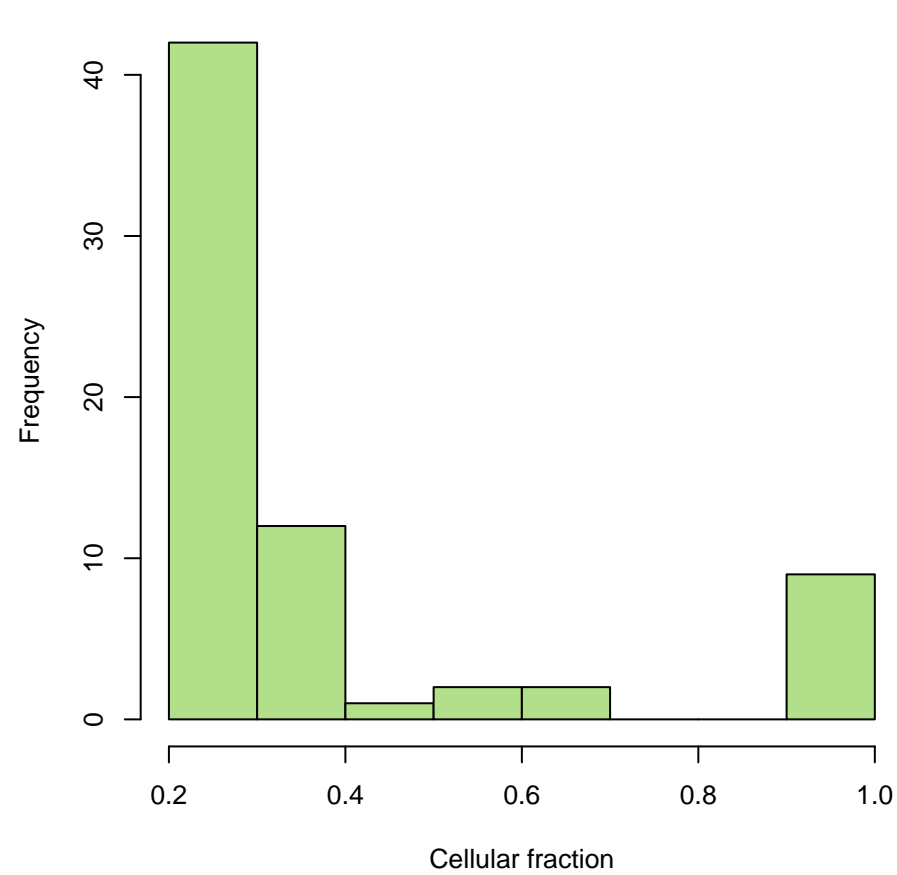
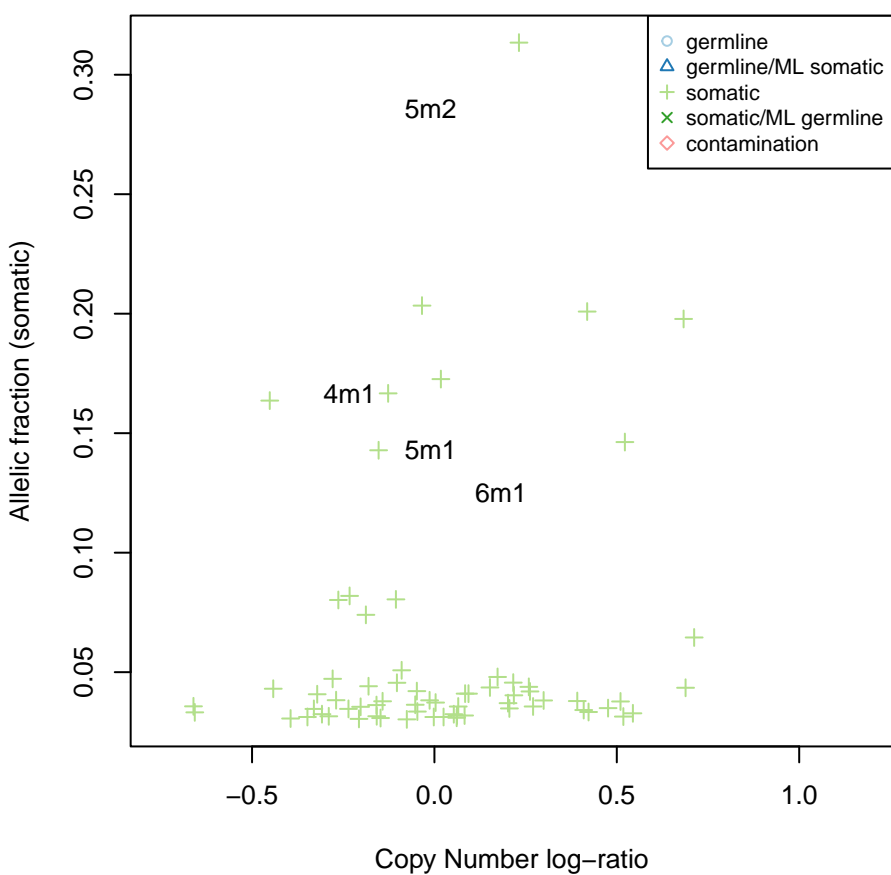
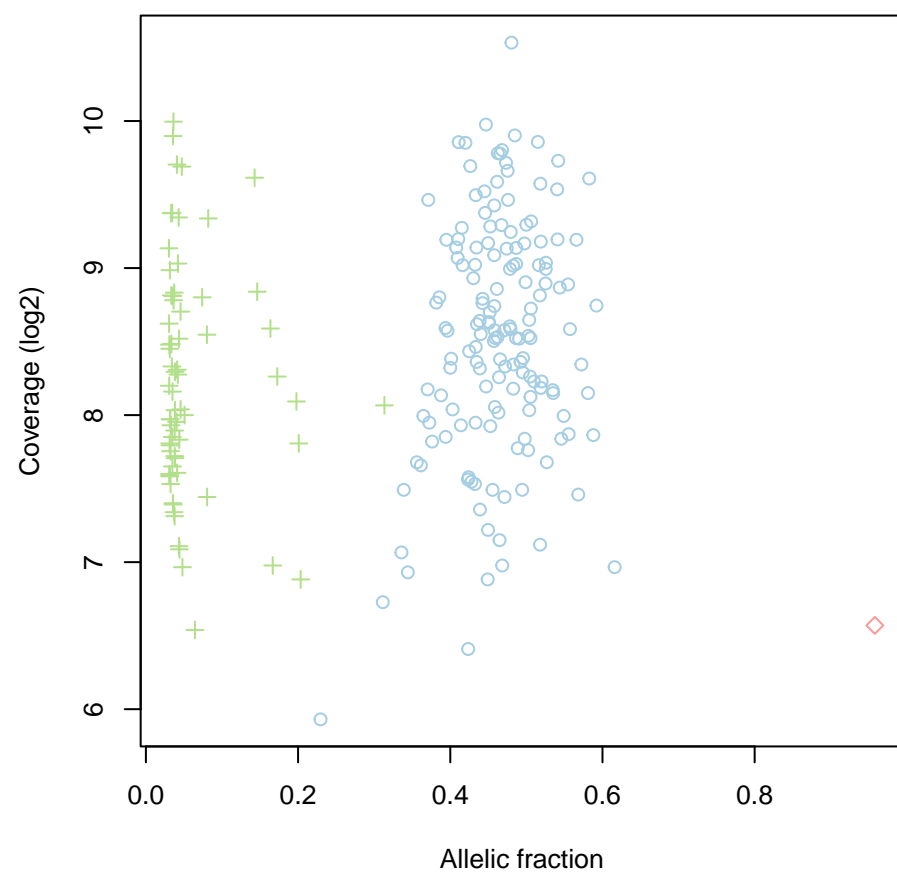
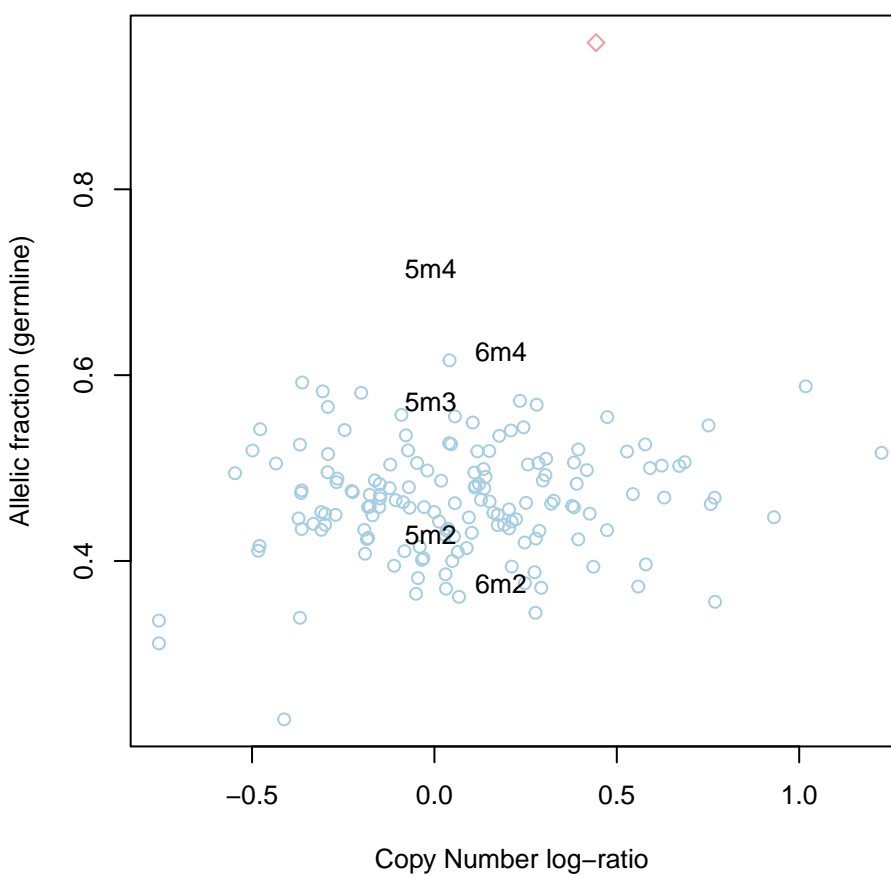
Purity: 0.5 Tumor ploidy: 5.05 SNV log-likelihood: -189.05 GoF: 89.2% Mean coverage: 401;62



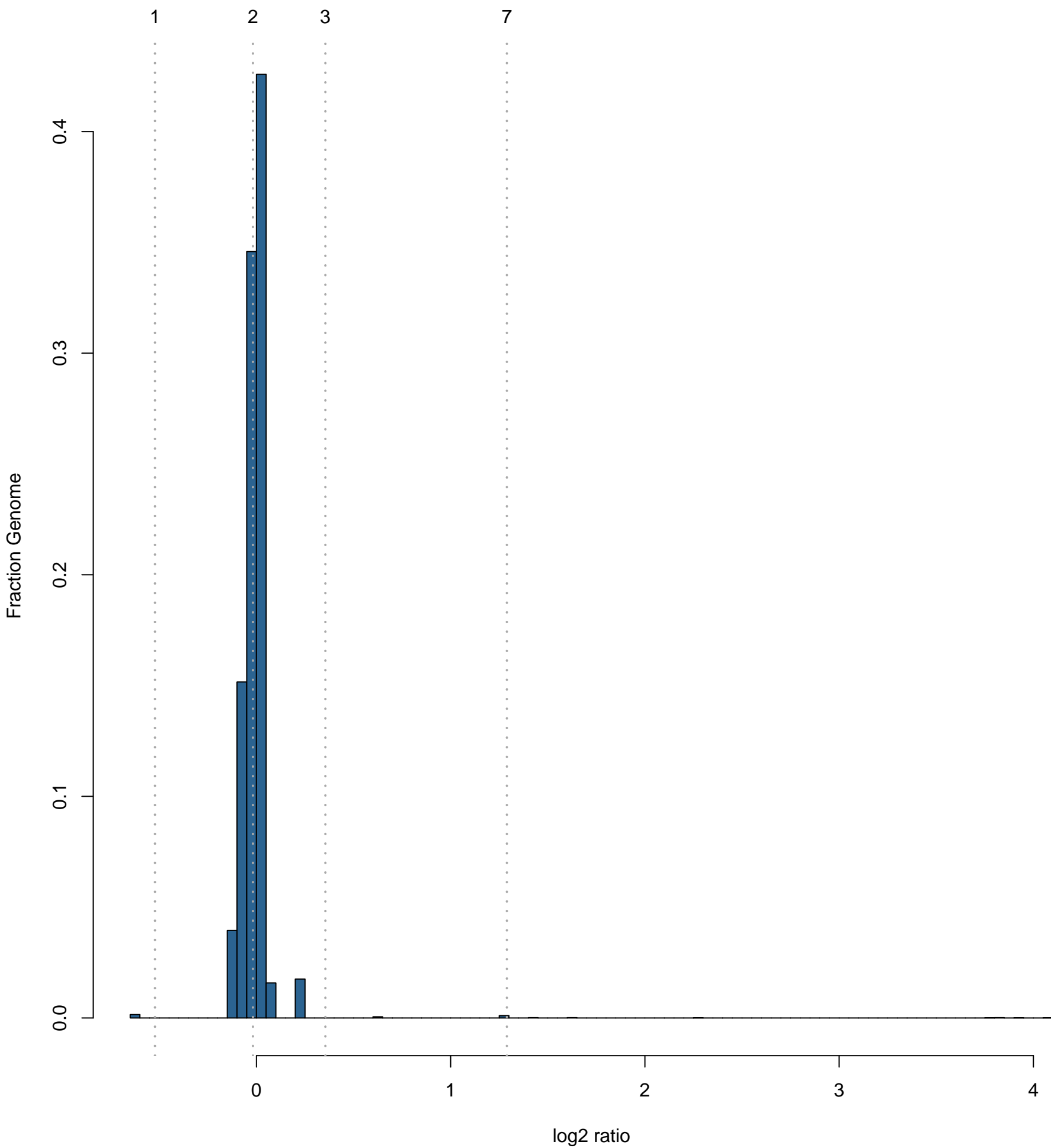
SCNA-fit log-likelihood: -15096.48



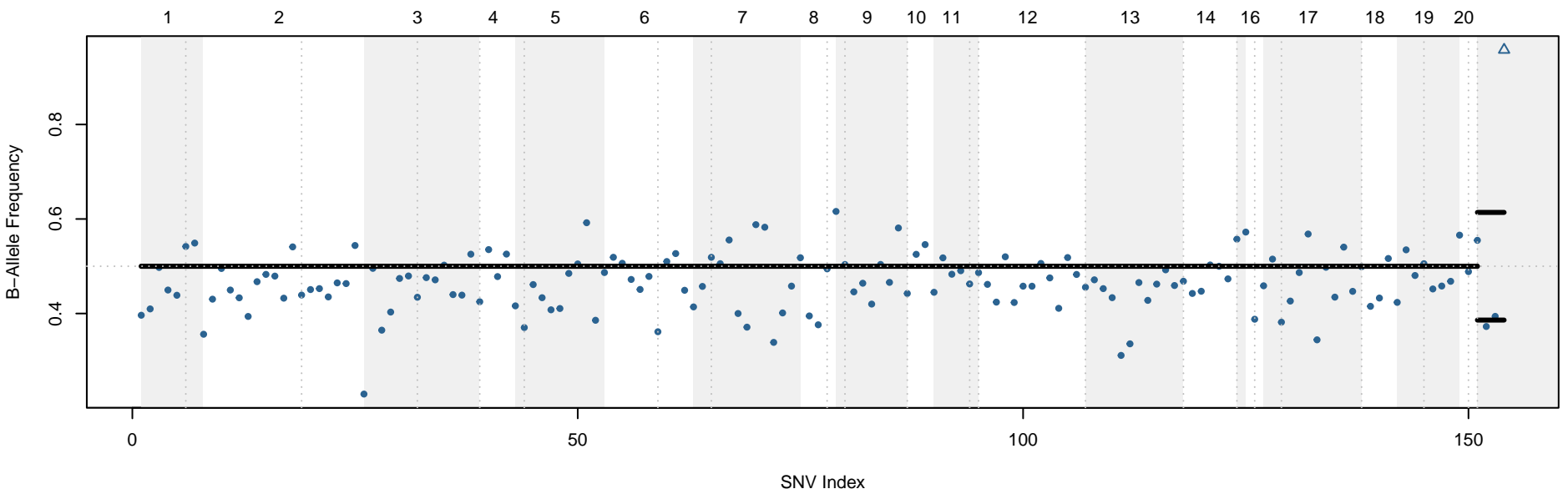




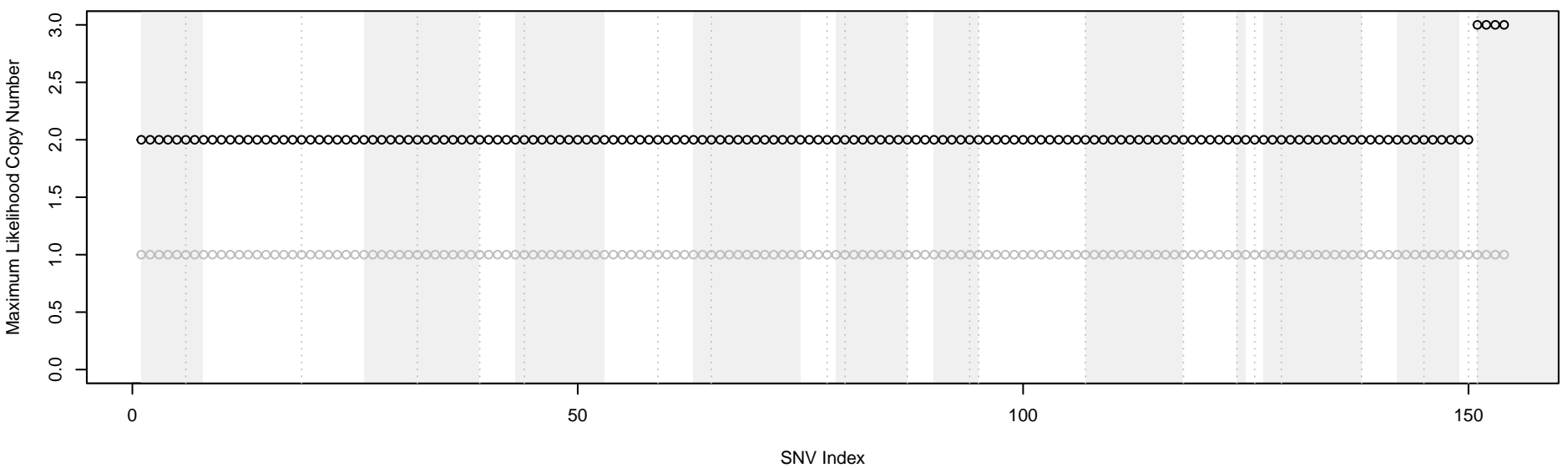
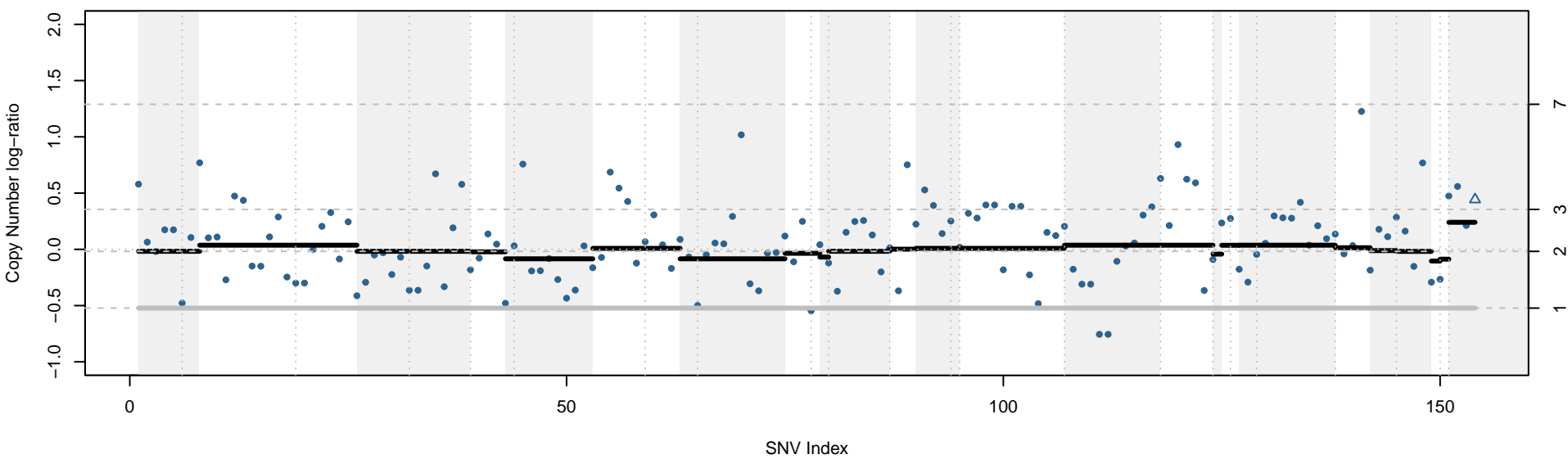
Purity: 0.59 Tumor ploidy: 2.043

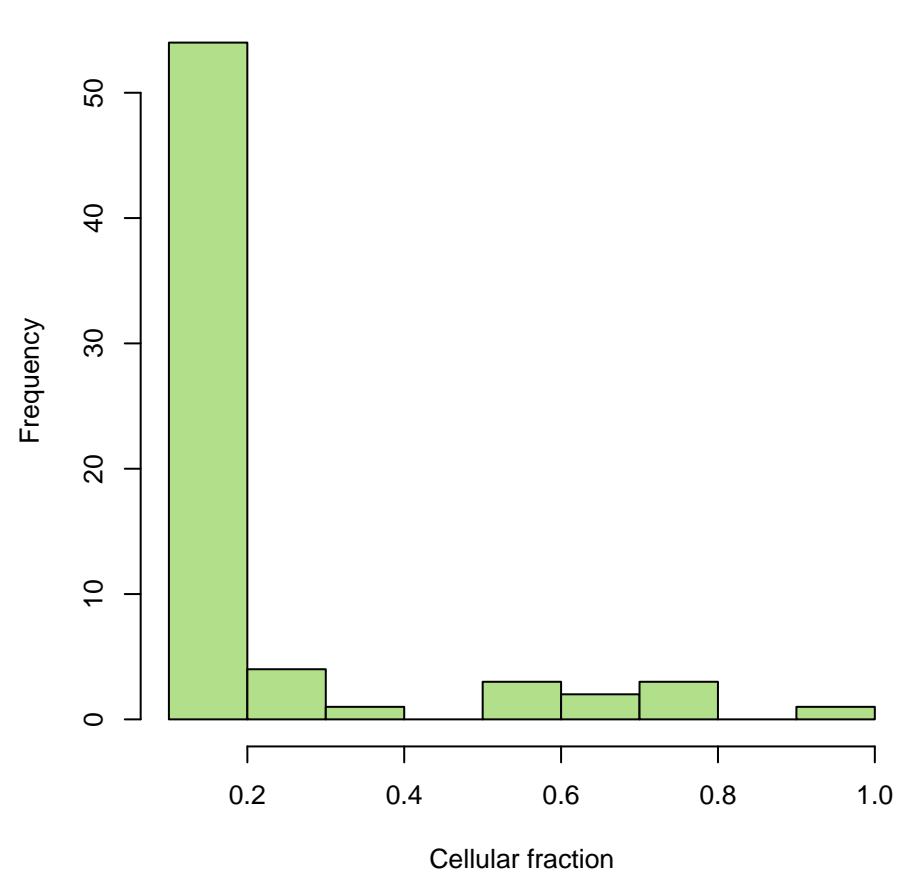
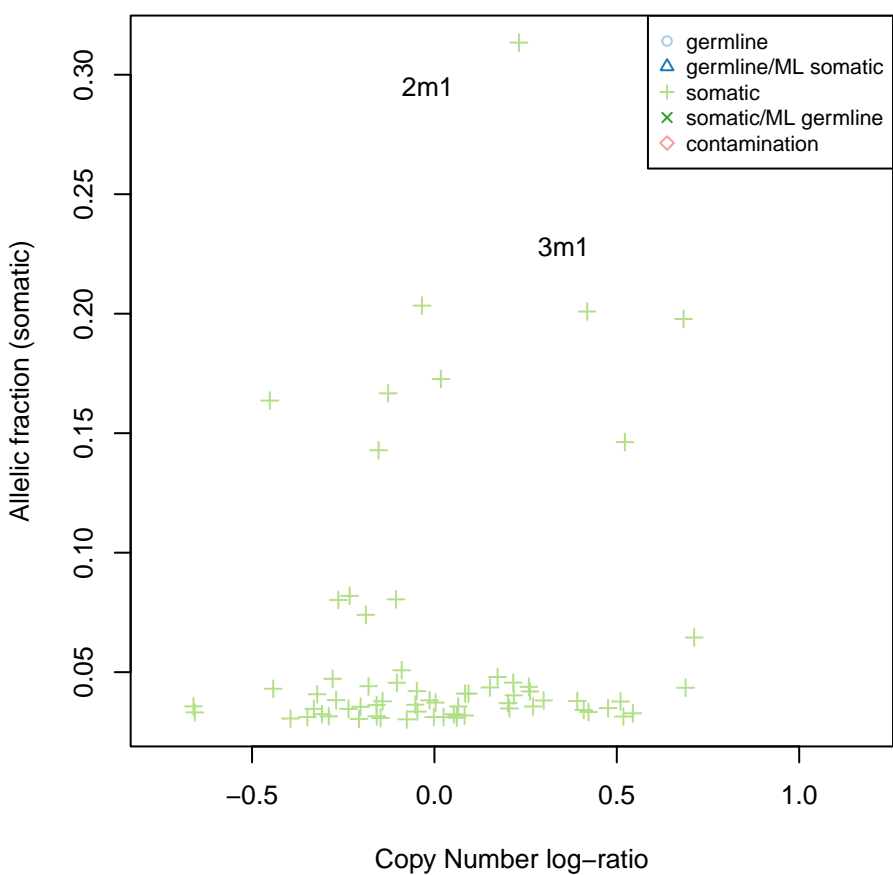
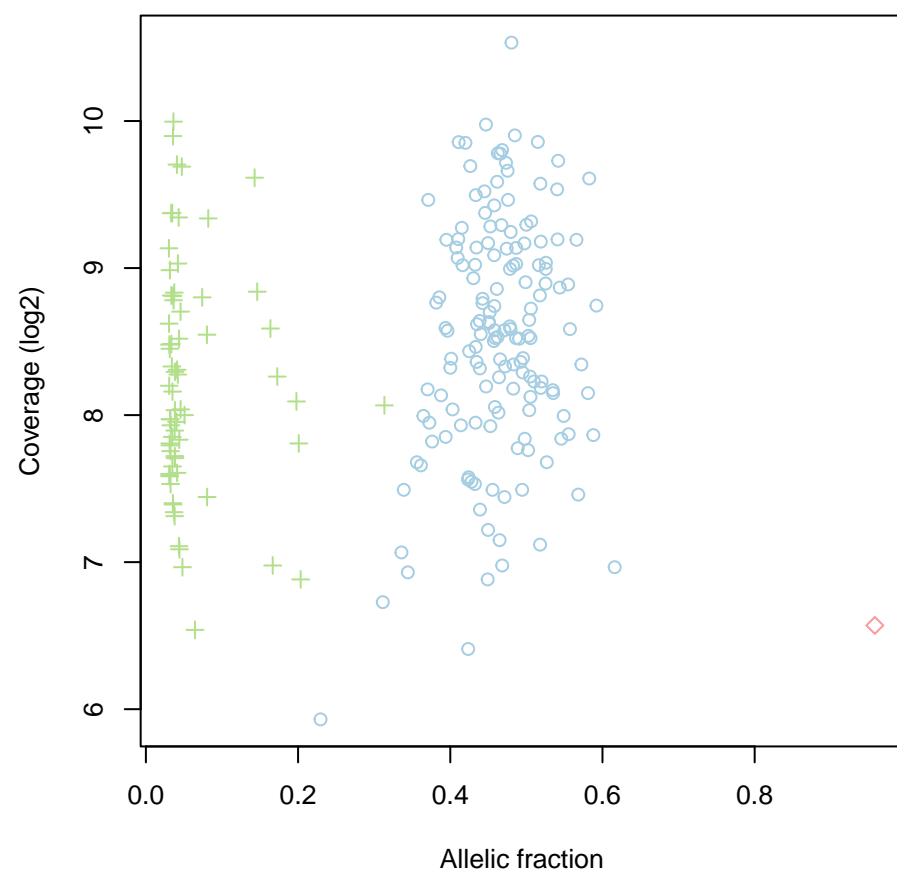
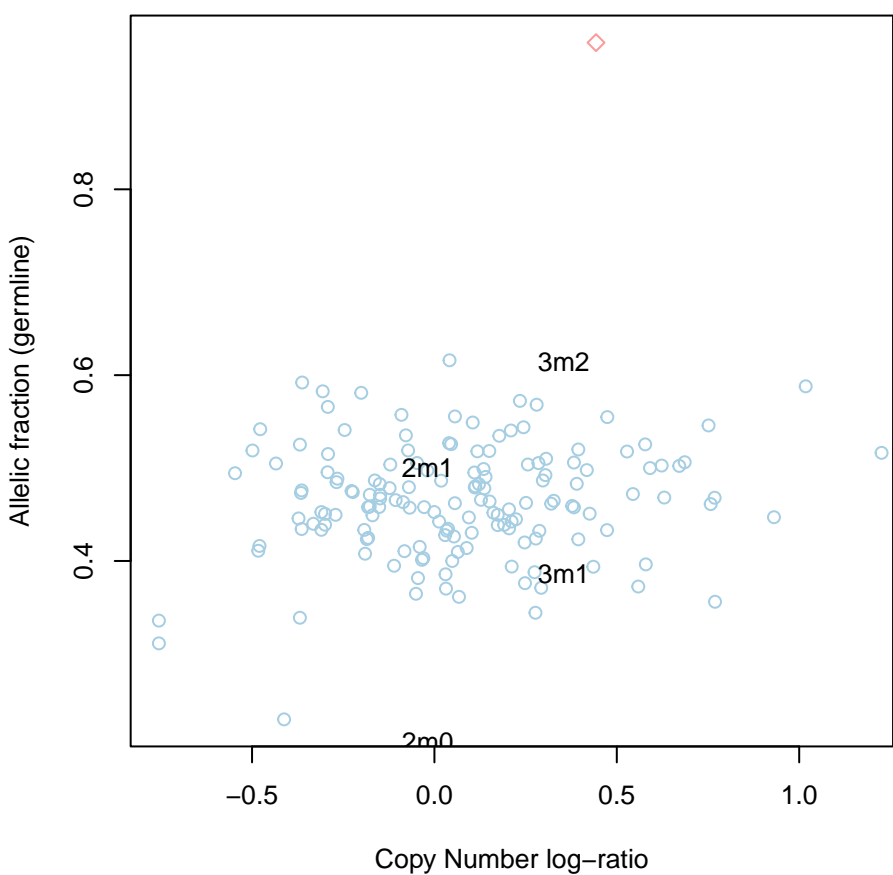


Purity: 0.59 Tumor ploidy: 2.043 SNV log-likelihood: -309.39 GoF: 51.2% Mean coverage: 401;62

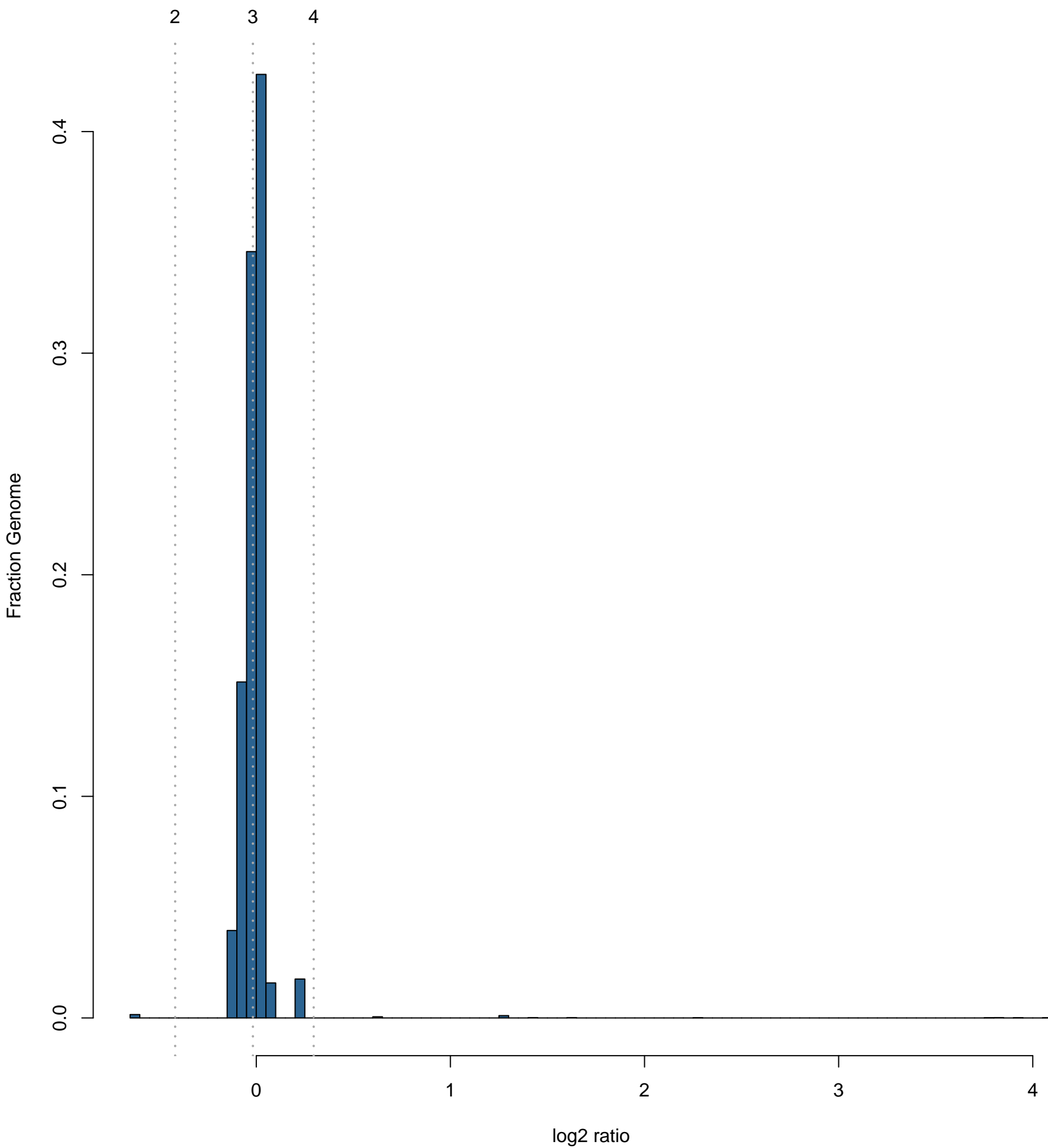


SCNA-fit log-likelihood: -15092.51

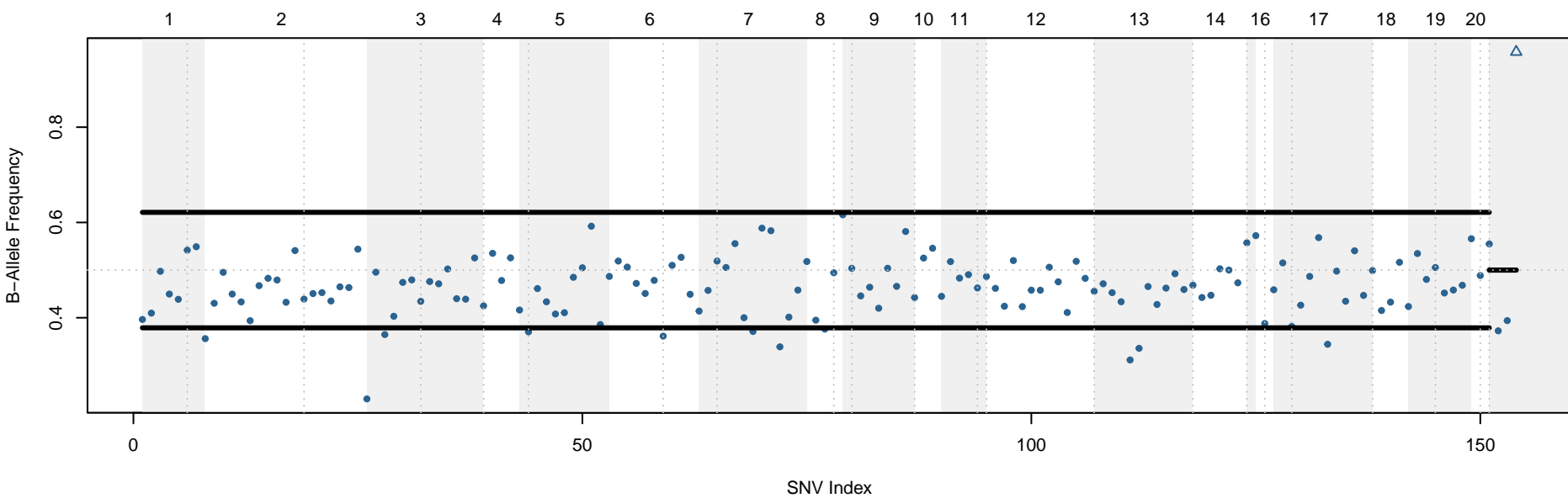




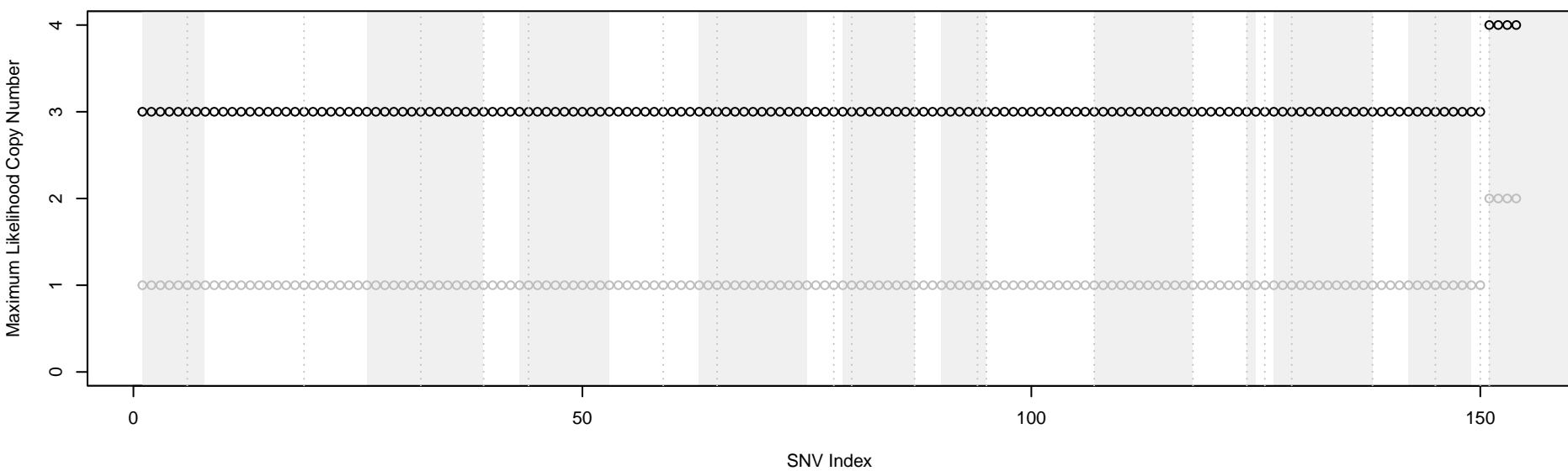
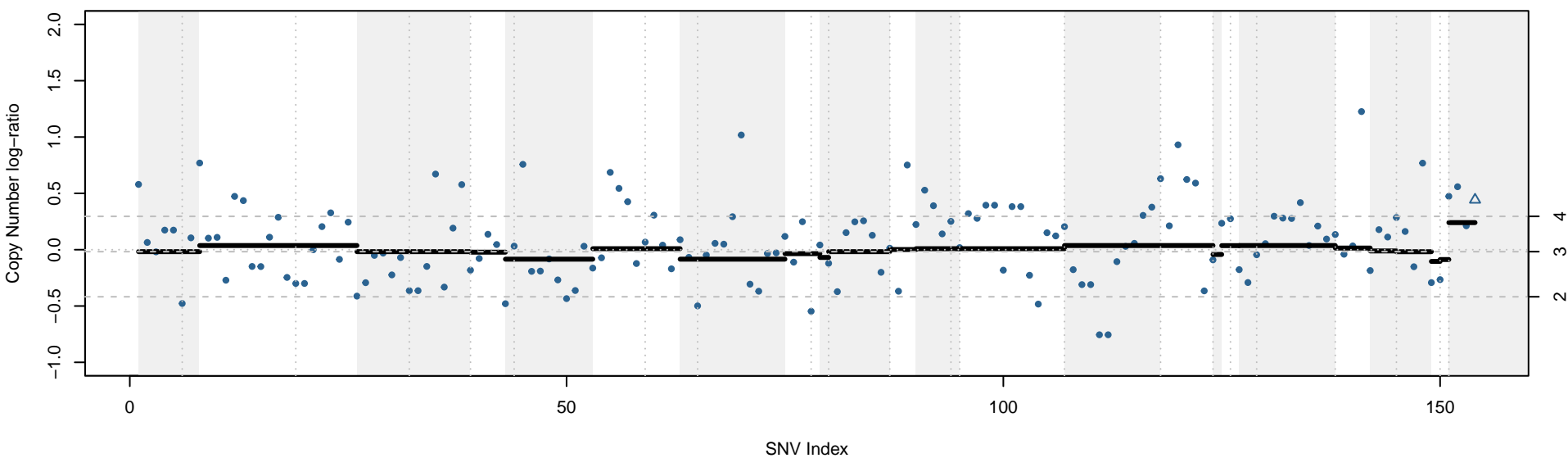
Purity: 0.64 Tumor ploidy: 3.05

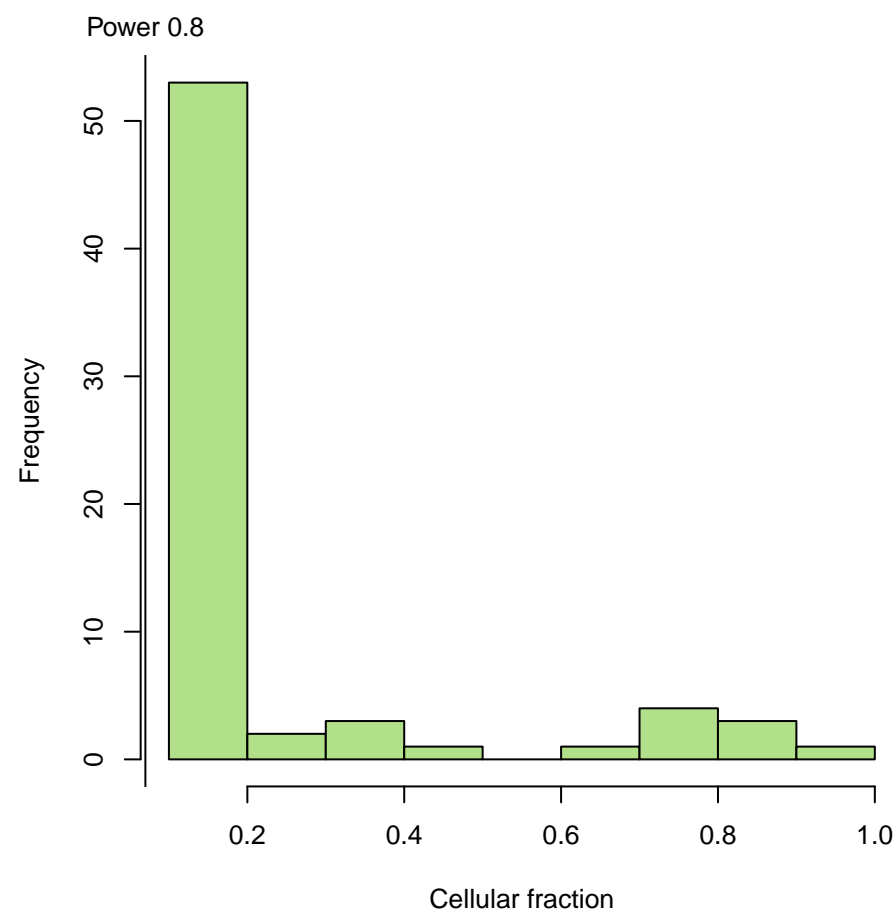
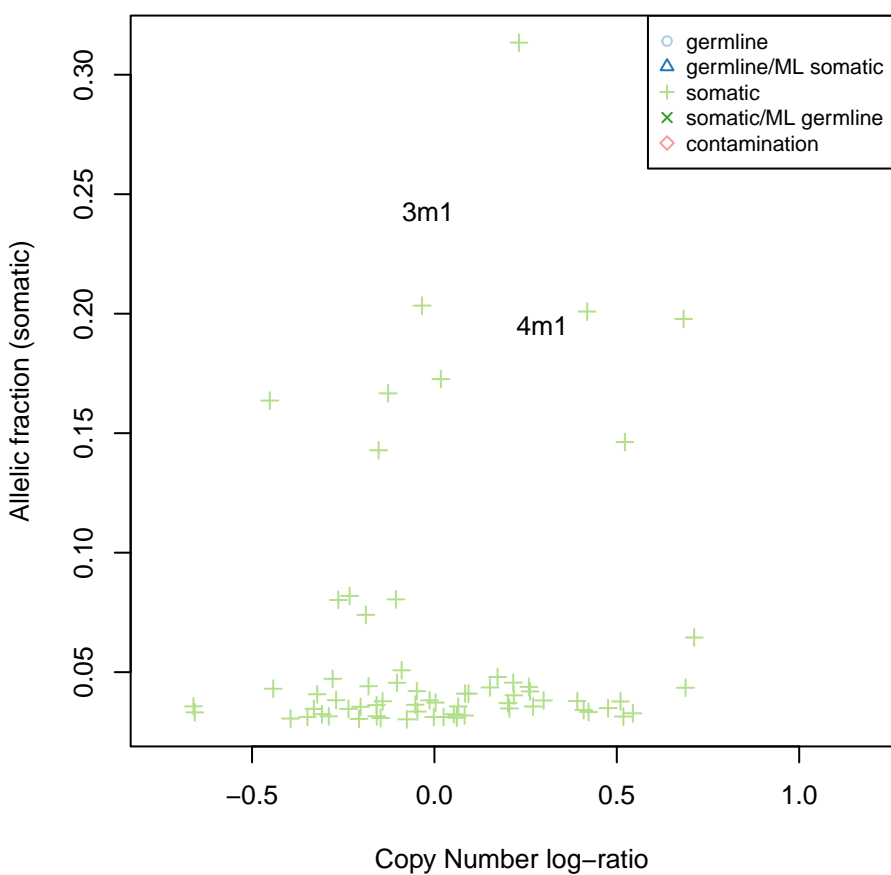
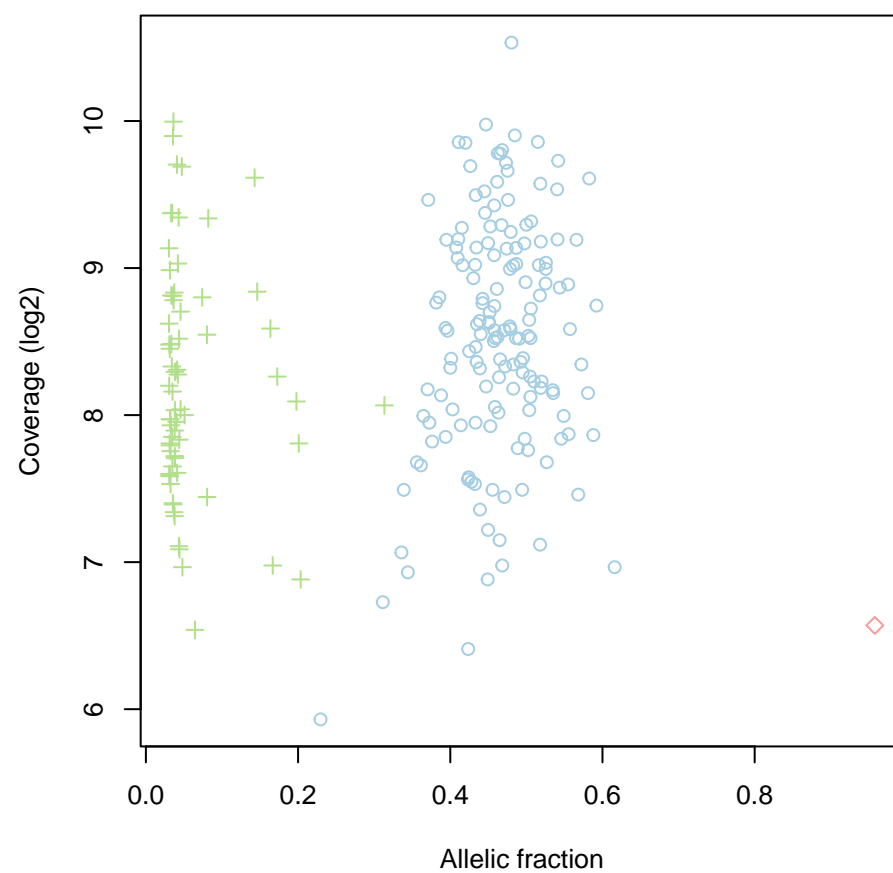
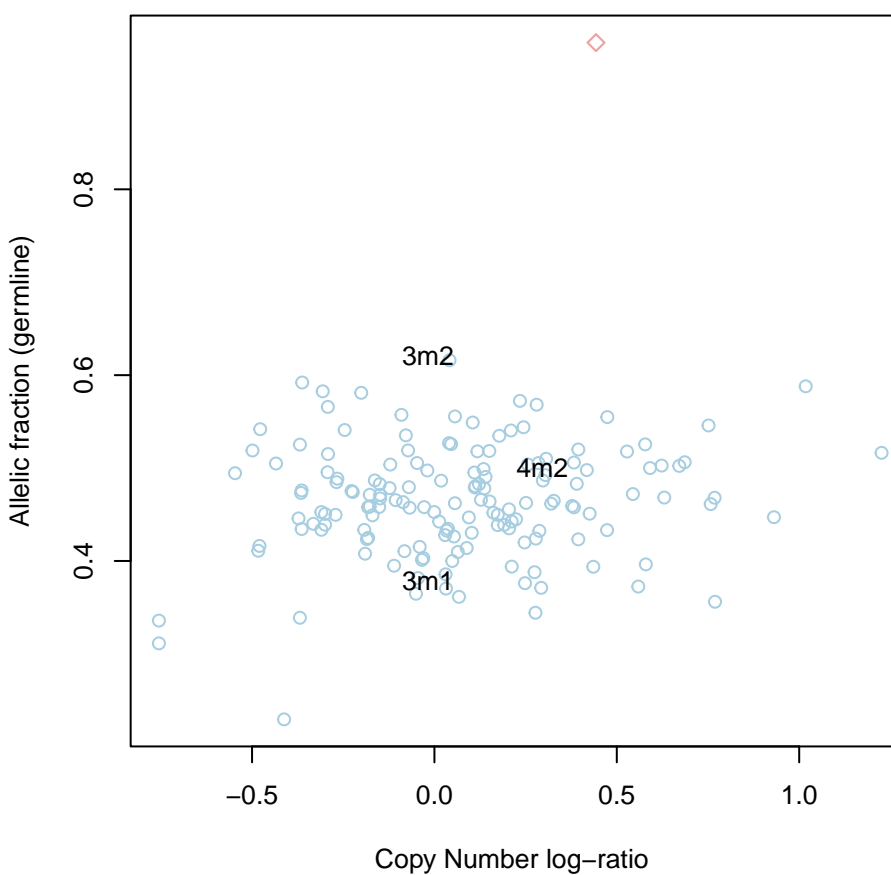


Purity: 0.64 Tumor ploidy: 3.05 SNV log-likelihood: -687.17 GoF: 61.2% Mean coverage: 401;62

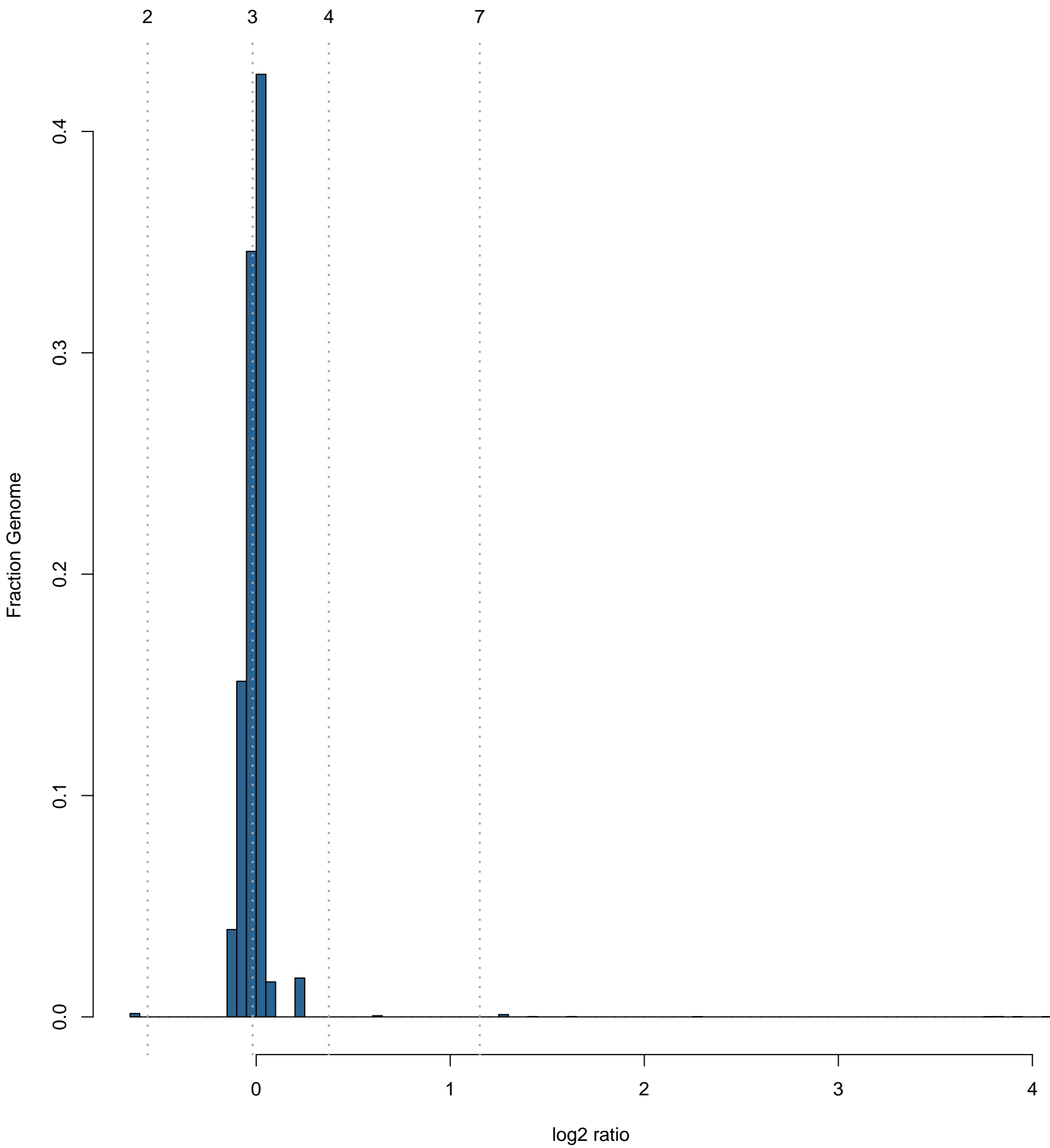


SCNA-fit log-likelihood: -15113.93



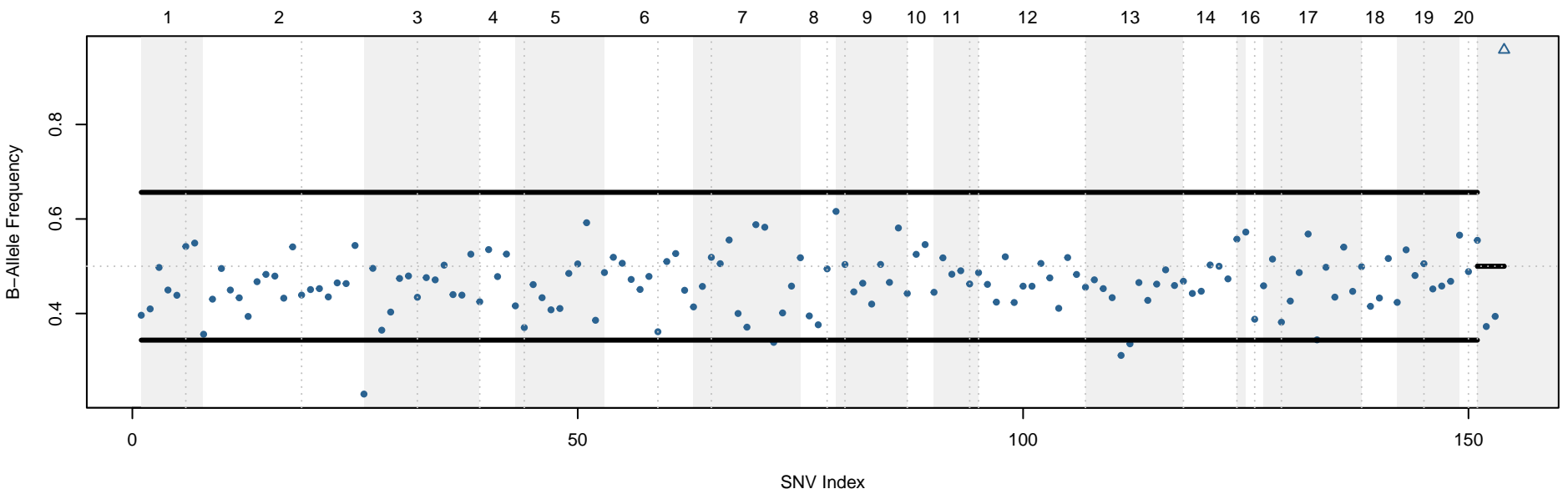


Purity: 0.91 Tumor ploidy: 3.041





Purity: 0.91 Tumor ploidy: 3.041 SNV log-likelihood: -1459.62 GoF: 27.9% Mean coverage: 401;62



SCNA-fit log-likelihood: -15096.7

