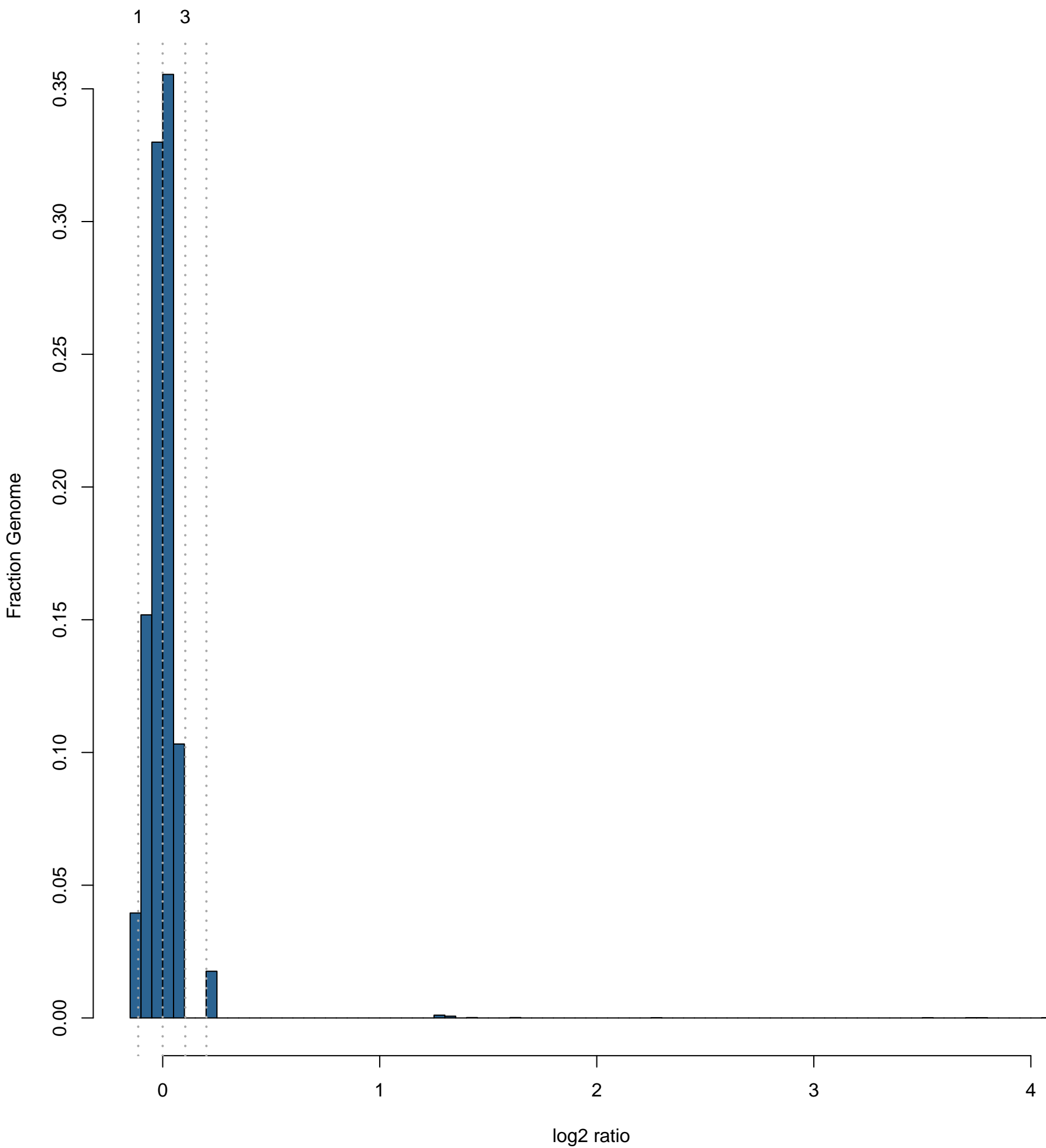
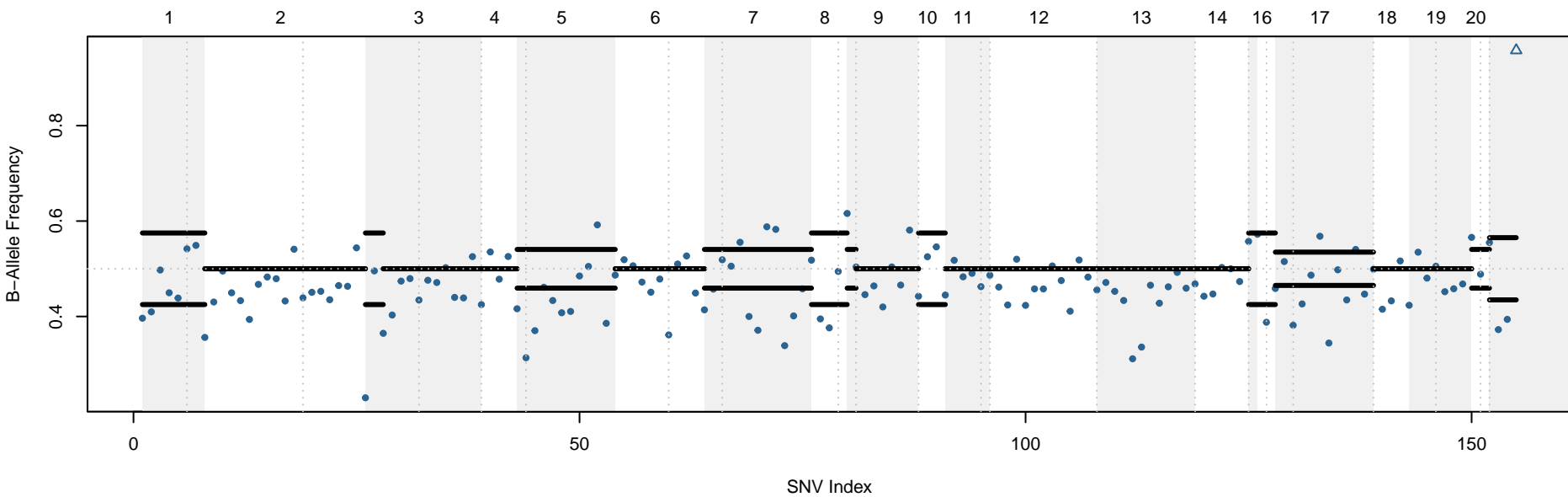


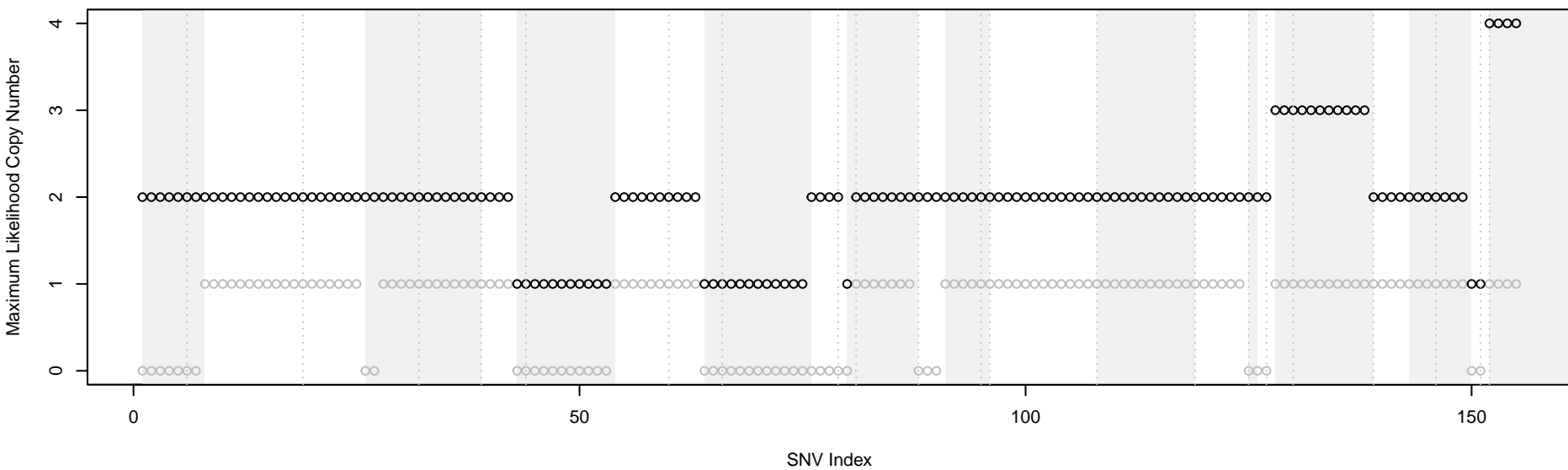
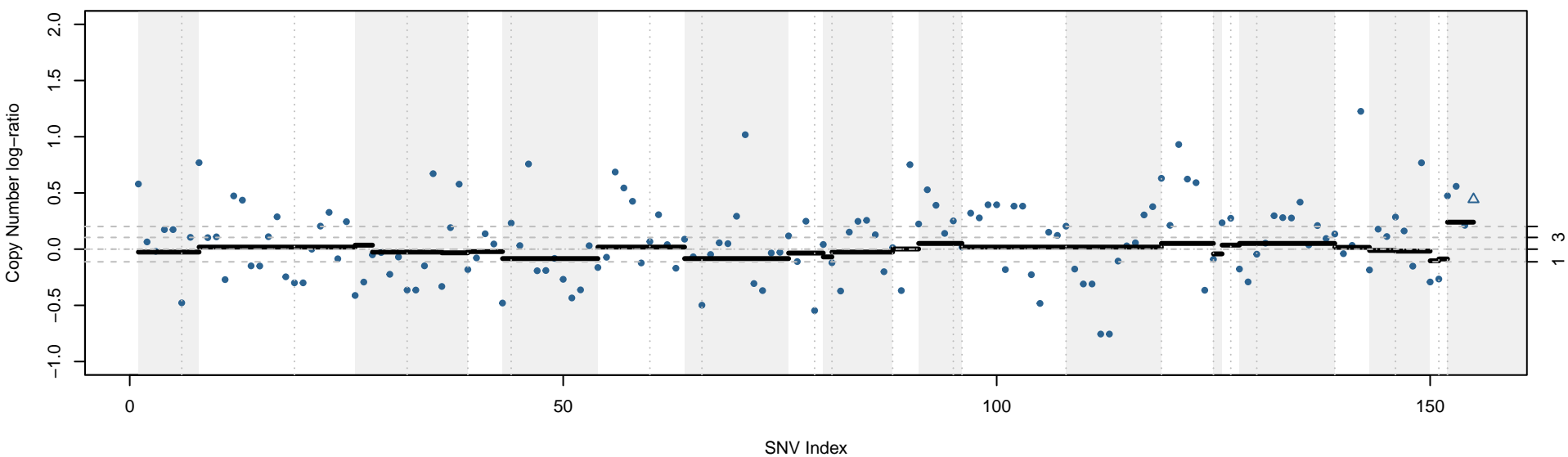
Purity: 0.15 Tumor ploidy: 2.002

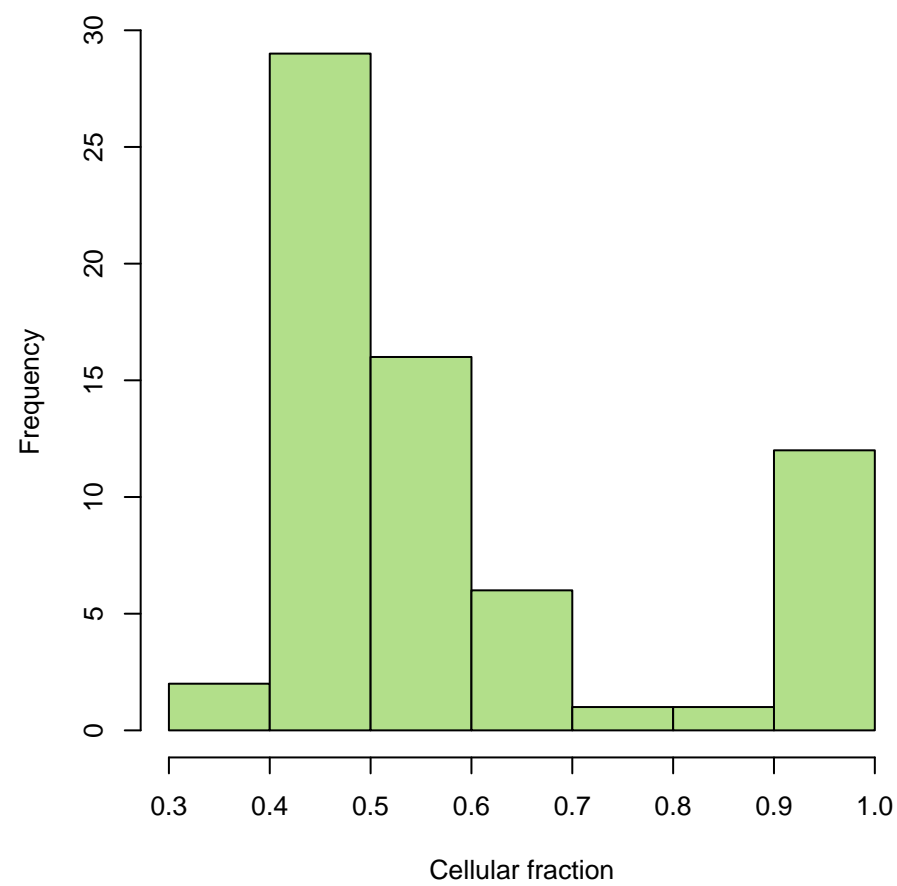
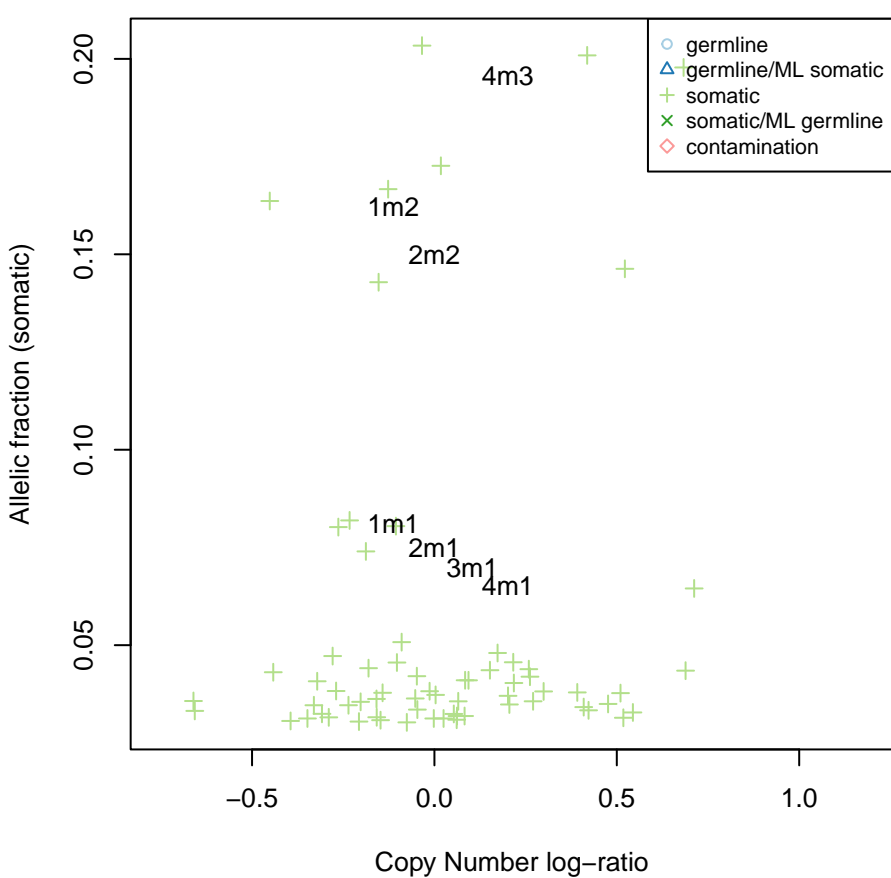
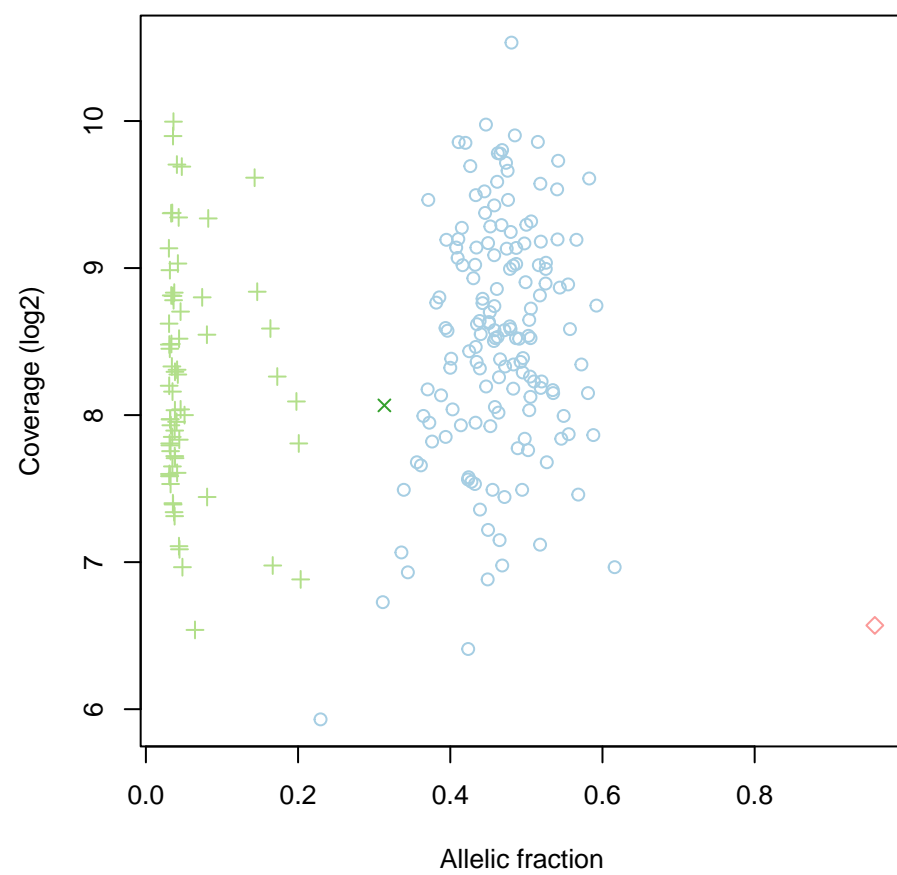
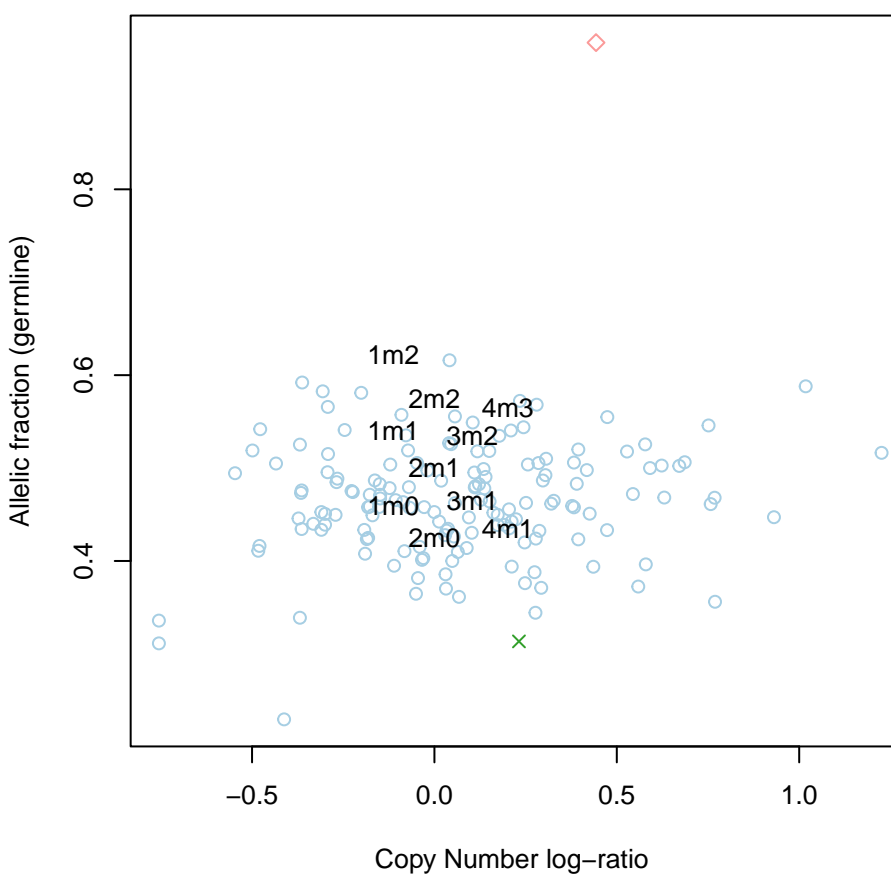


Purity: 0.15 Tumor ploidy: 2.002 SNV log-likelihood: 35.93 GoF: 94.7% Mean coverage: 401;62

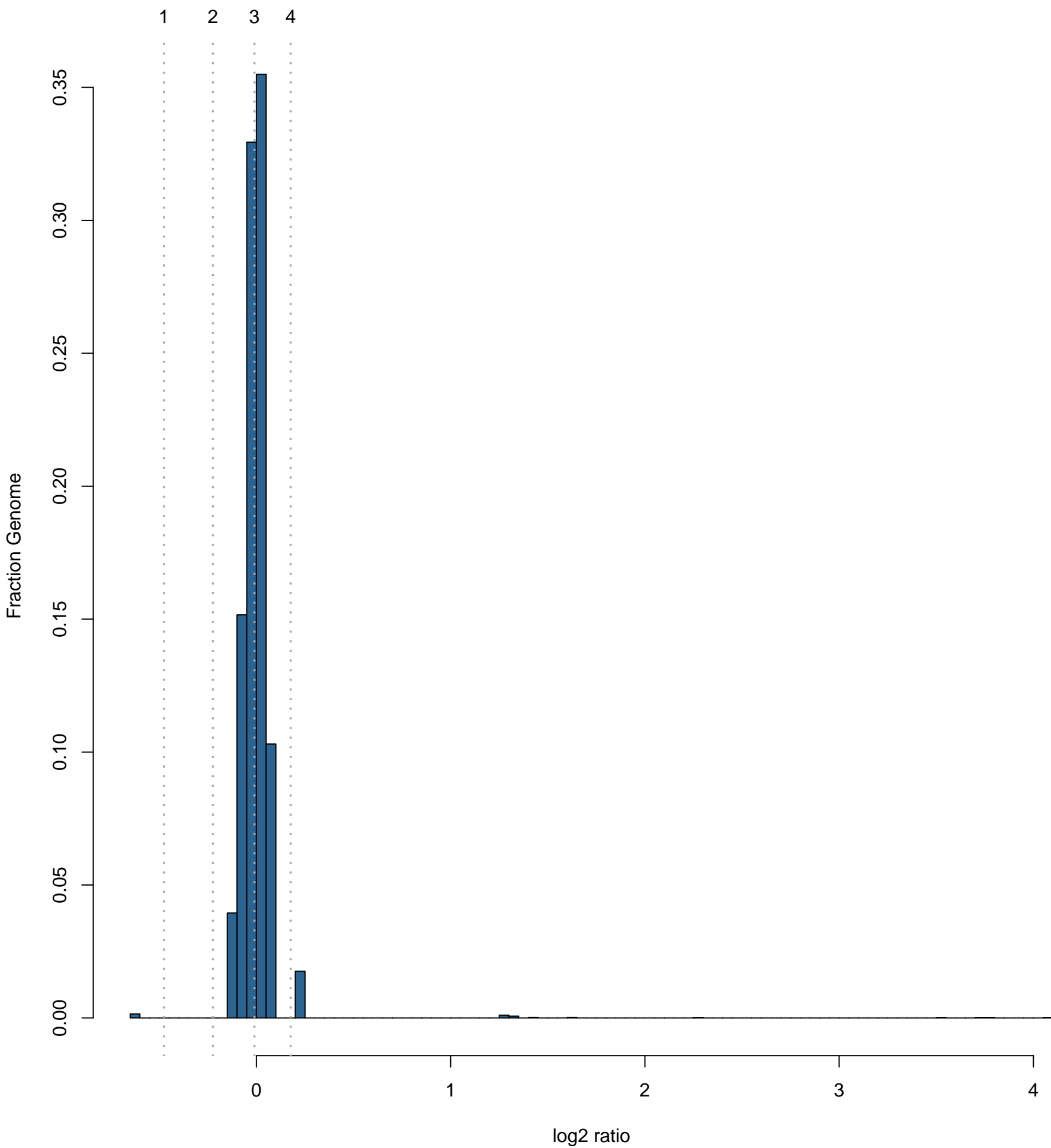


SCNA-fit log-likelihood: -15014.72

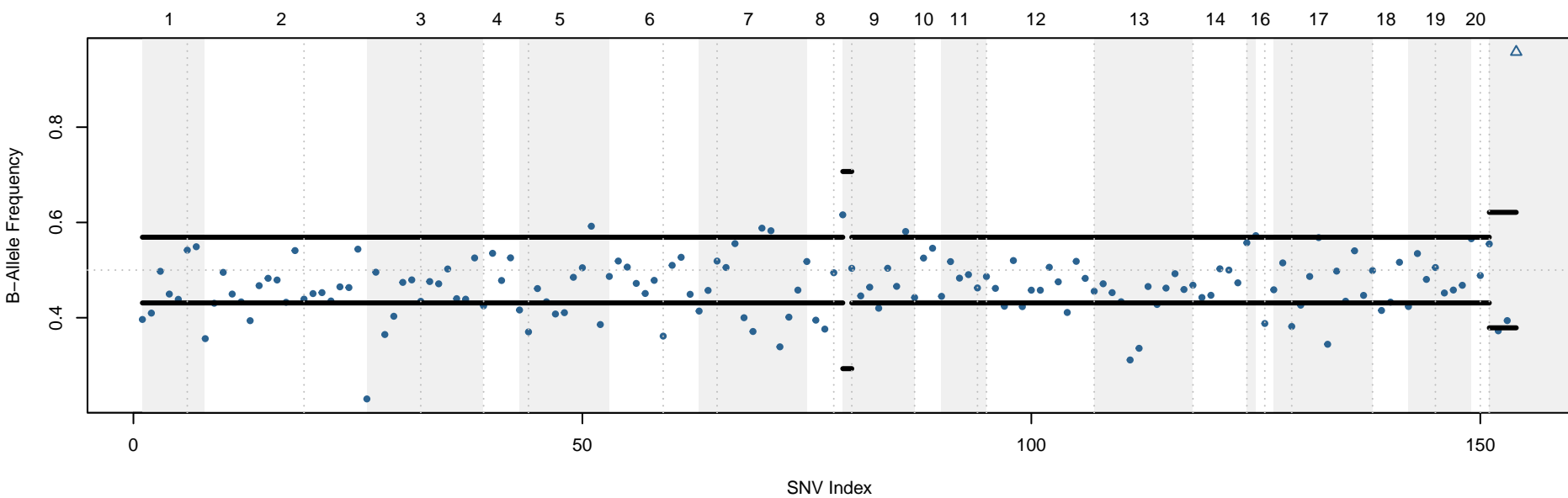




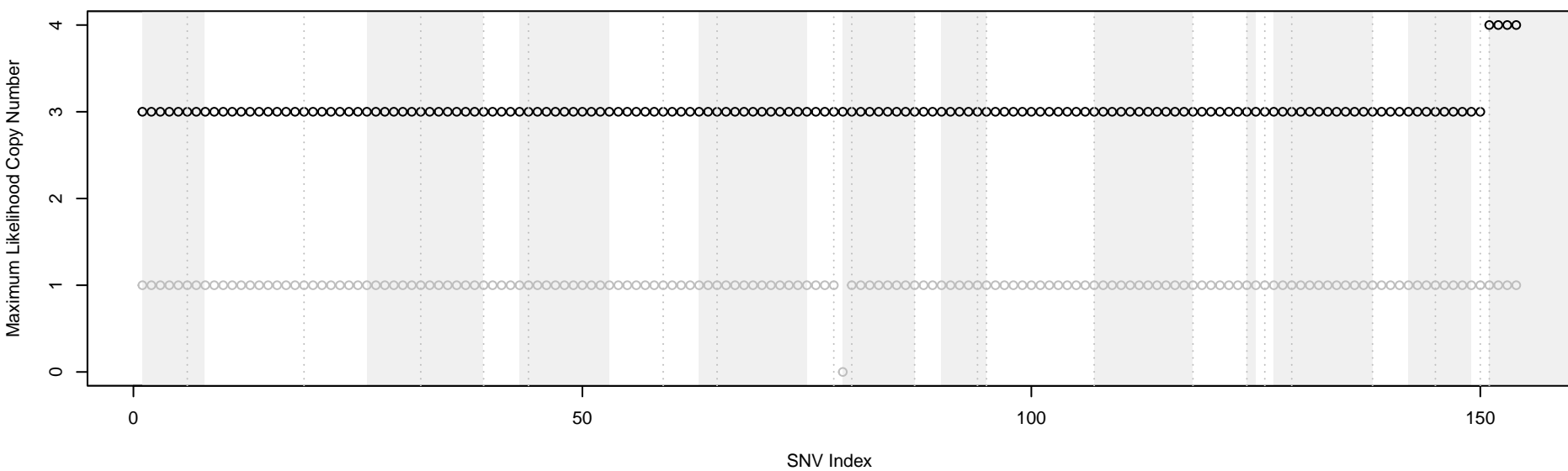
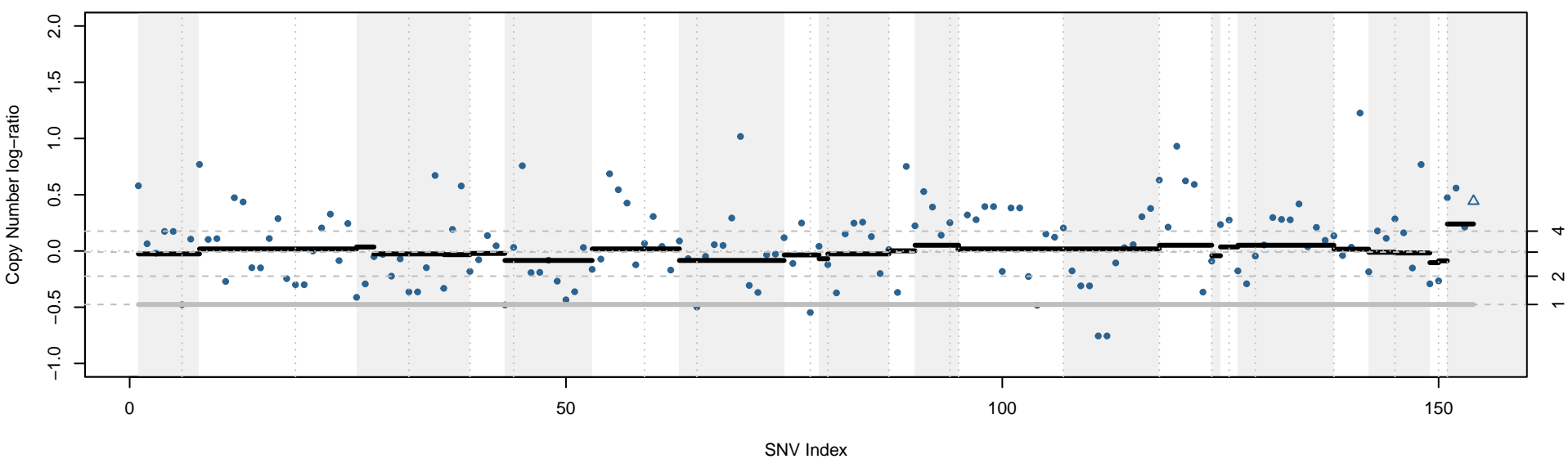
Purity: 0.32 Tumor ploidy: 3.051

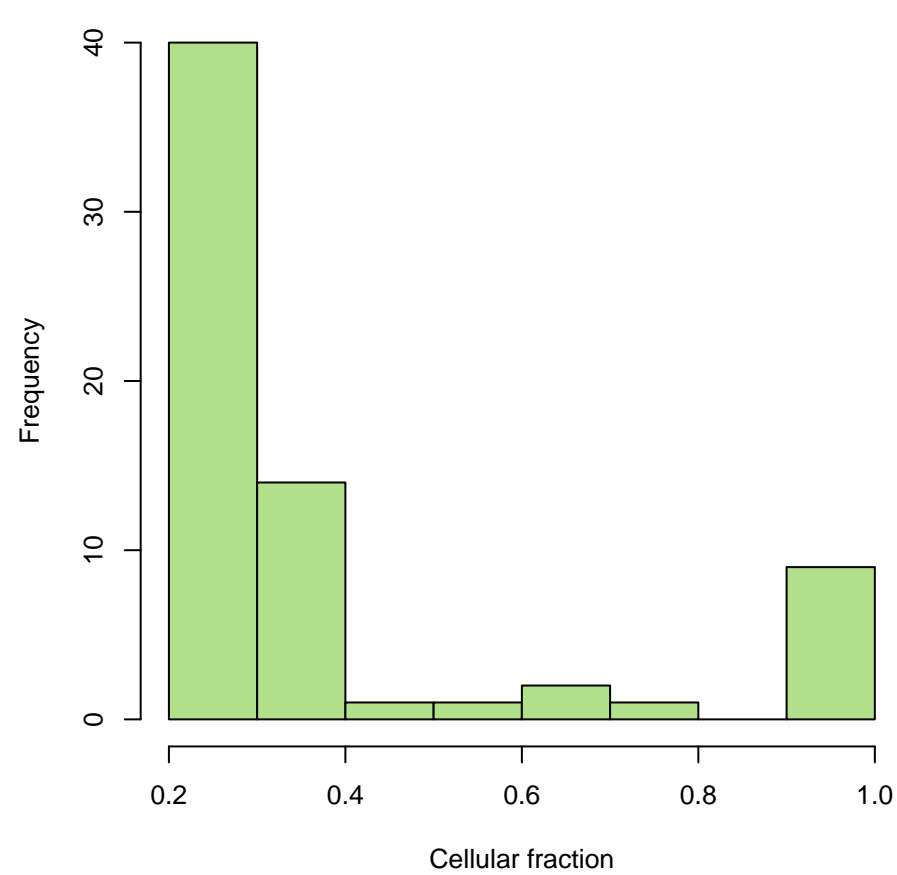
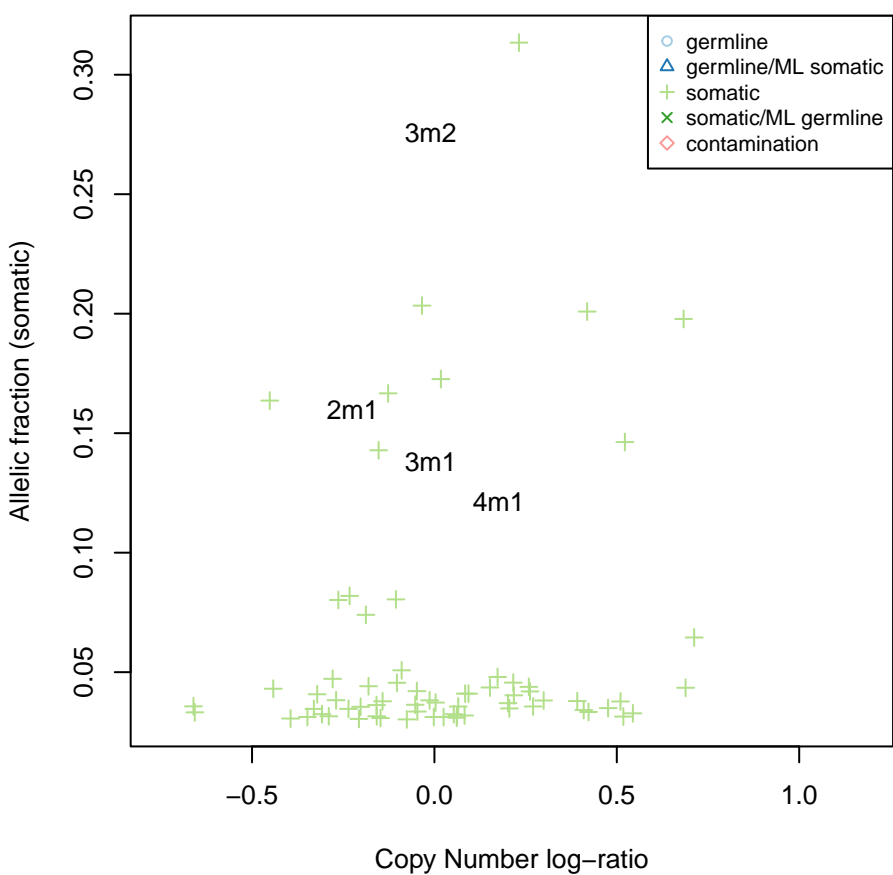
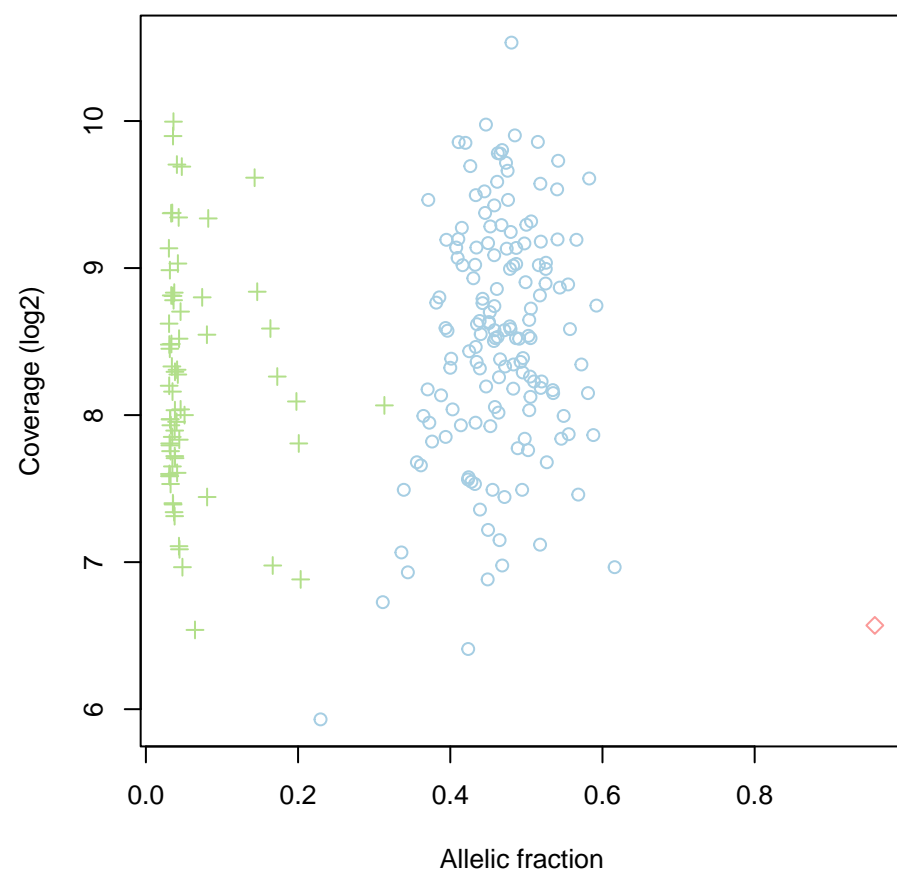
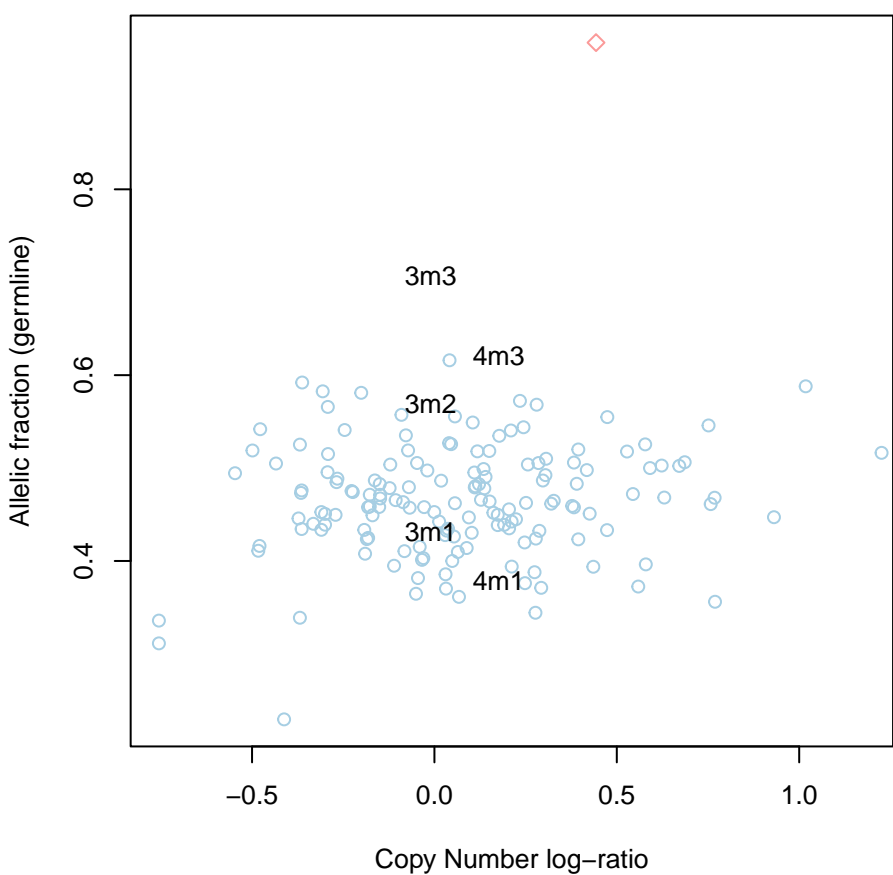


Purity: 0.32 Tumor ploidy: 3.051 SNV log-likelihood: -72.07 GoF: 90.1% Mean coverage: 401;62

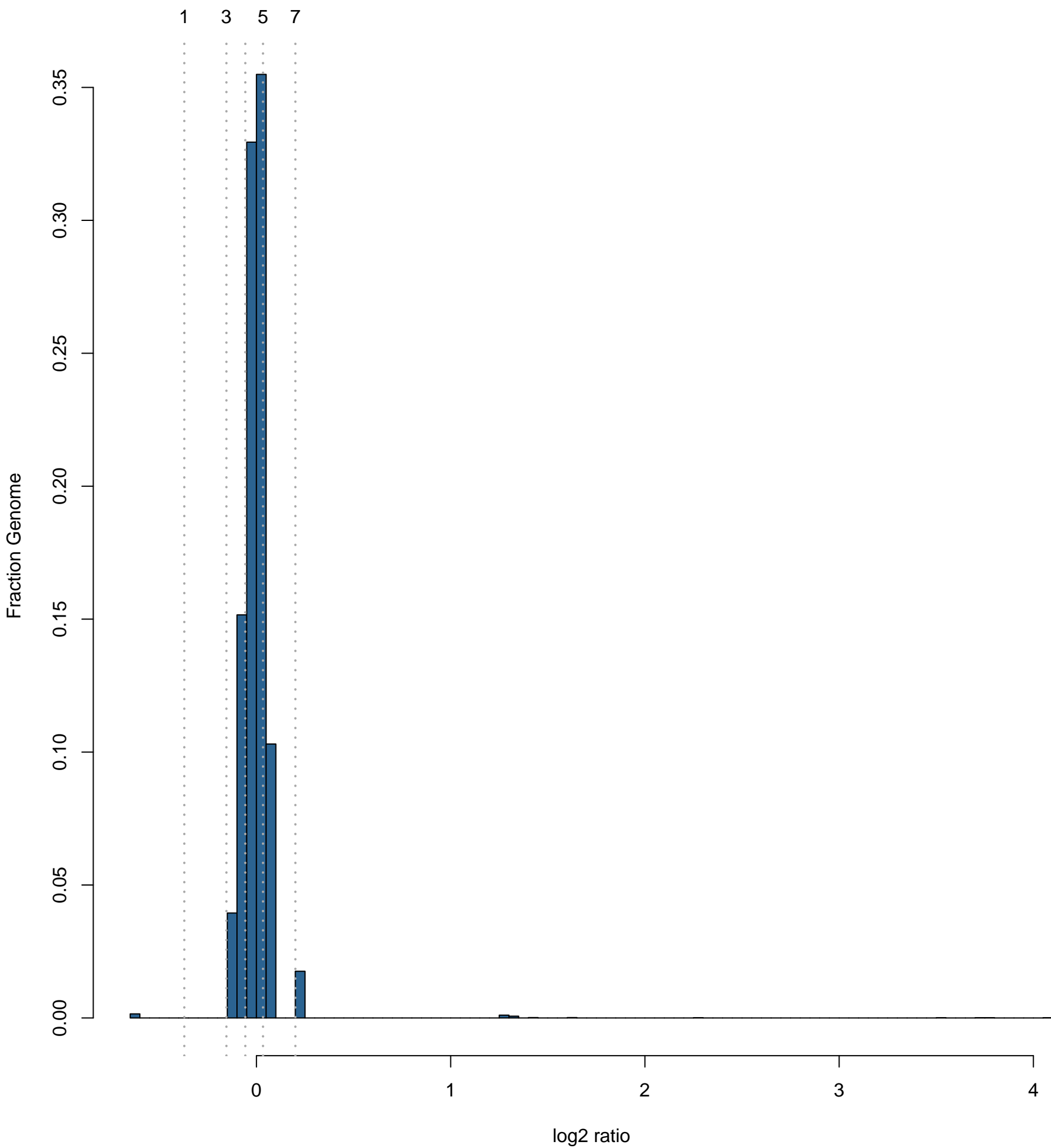


SCNA-fit log-likelihood: -15051.4

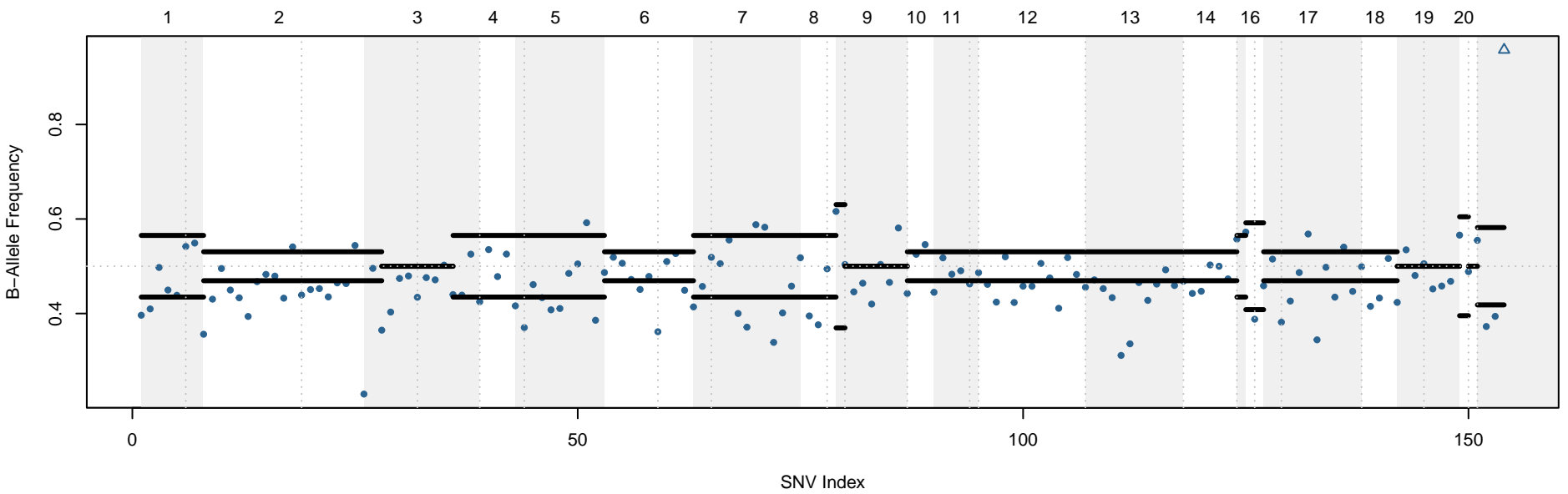




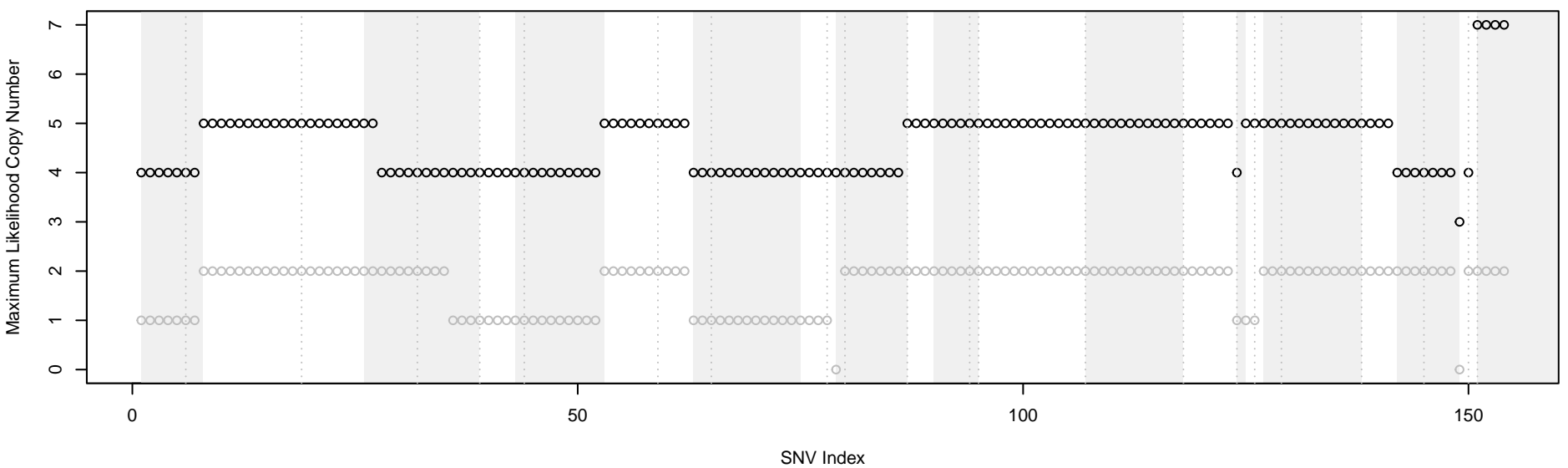
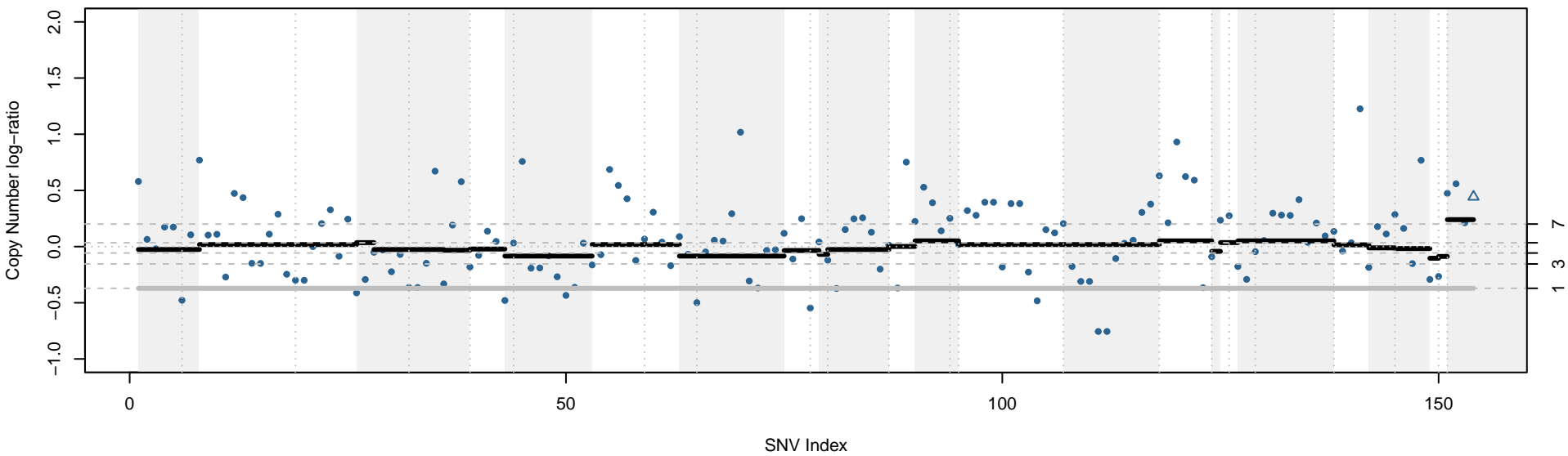
Purity: 0.15 Tumor ploidy: 4.621

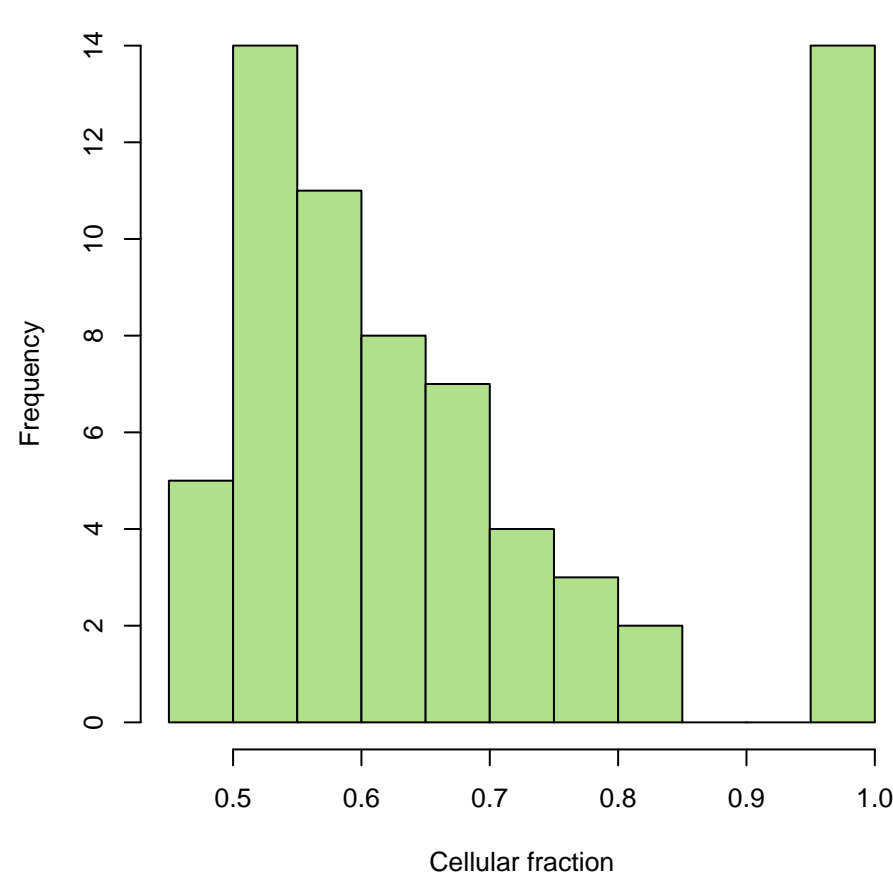
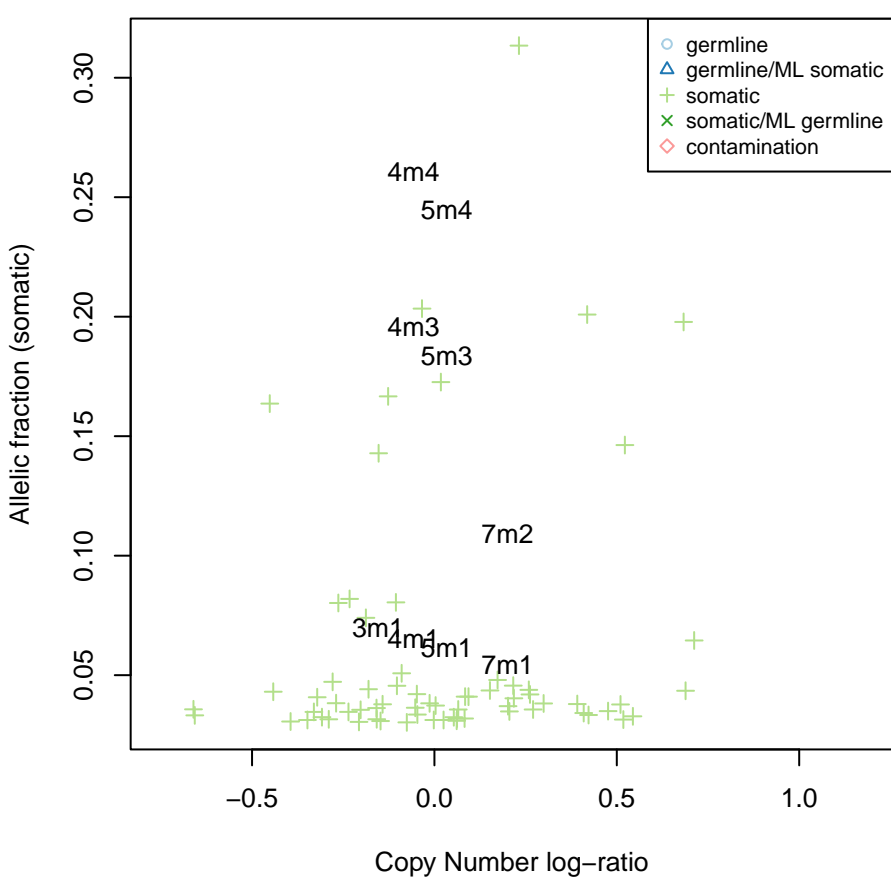
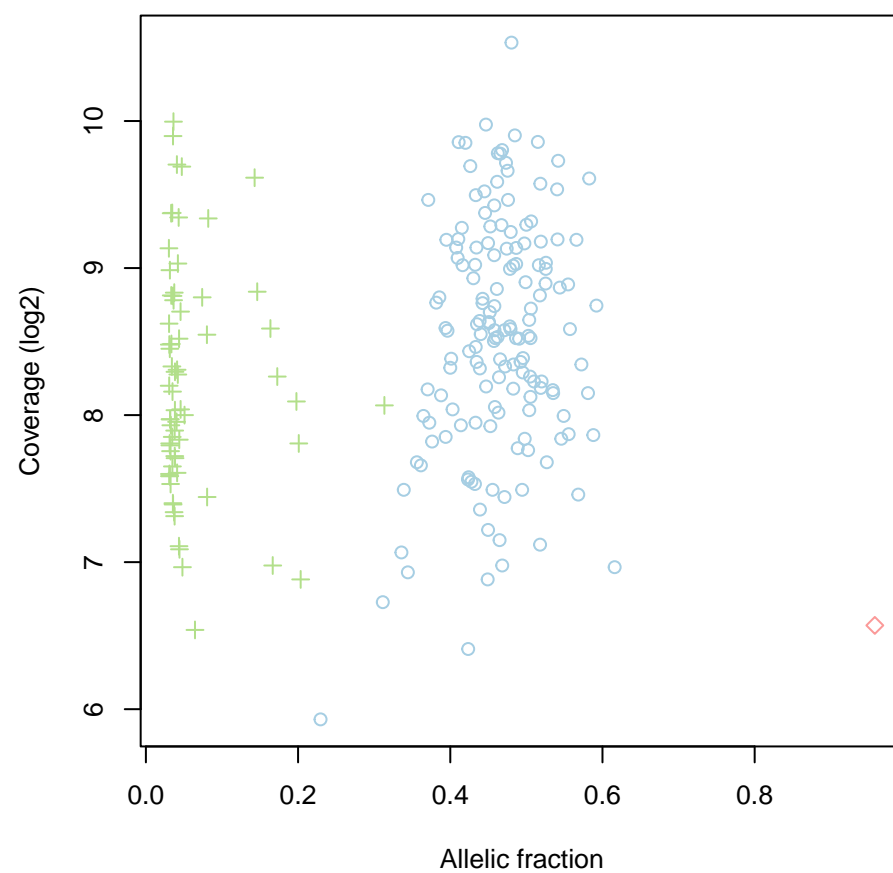
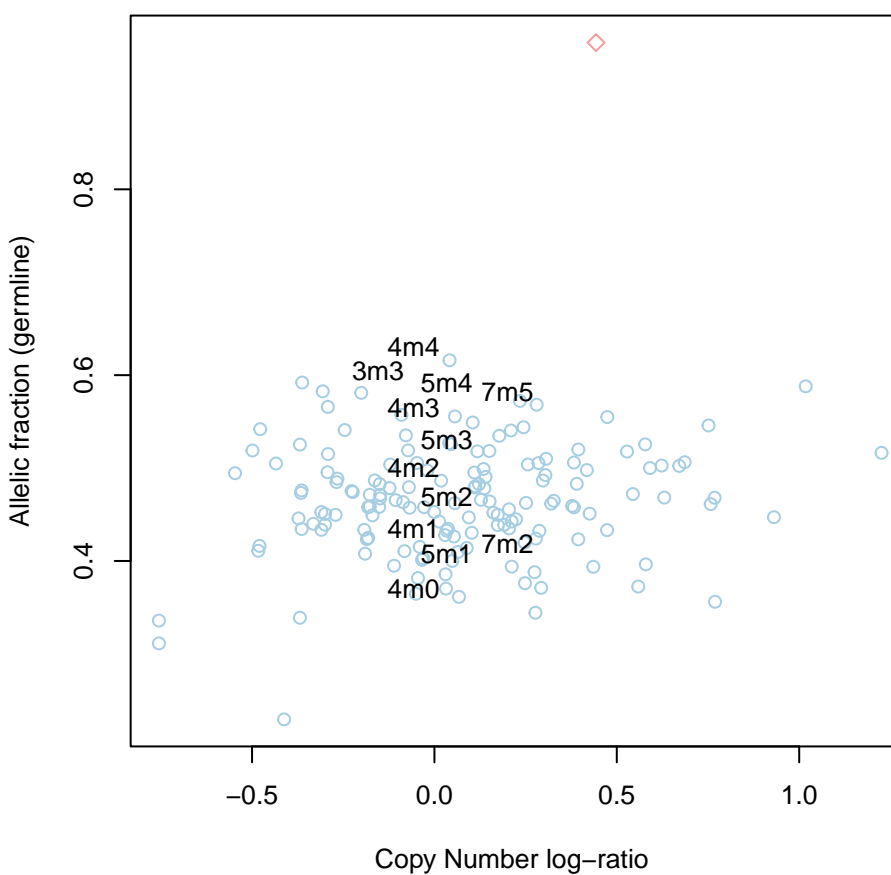


Purity: 0.15 Tumor ploidy: 4.621 SNV log-likelihood: -100.69 GoF: 96.8% Mean coverage: 401;62

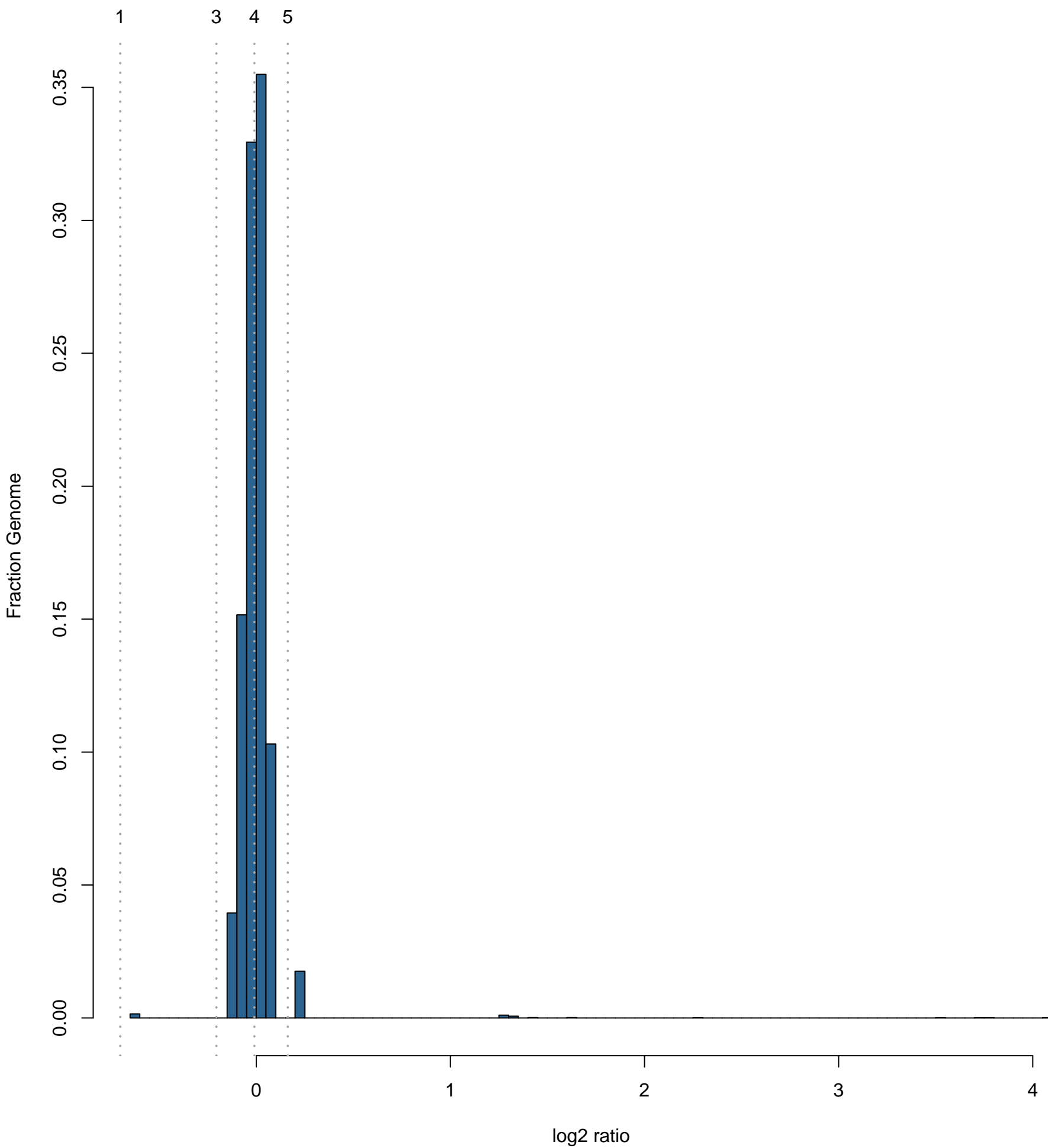


SCNA-fit log-likelihood: -14994.45

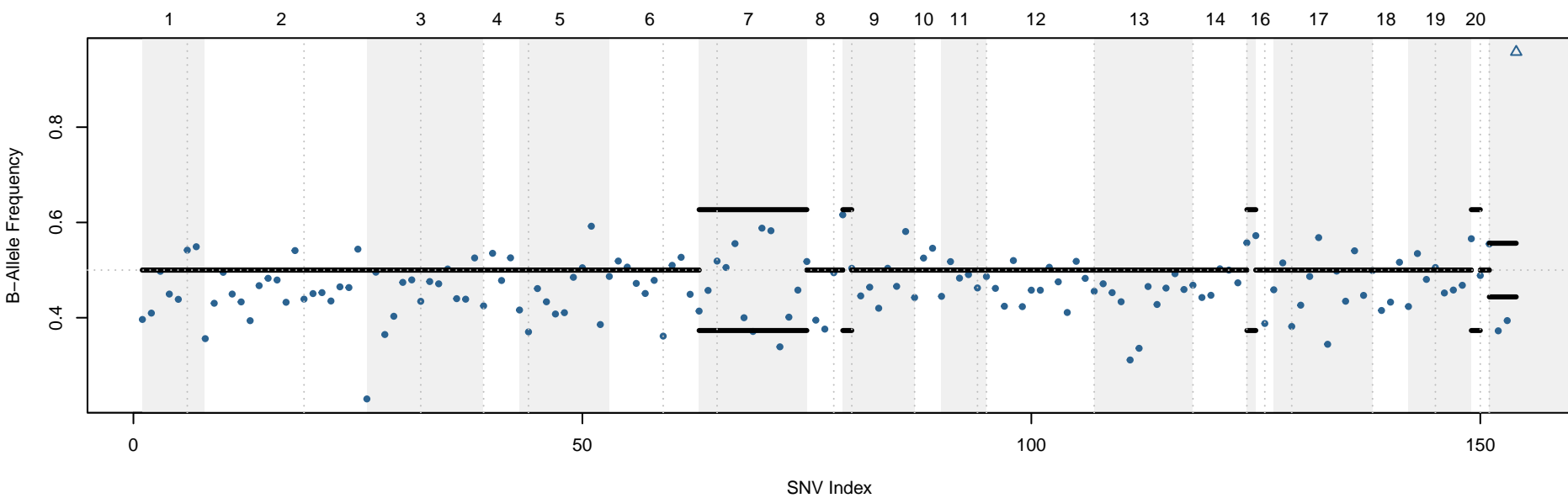




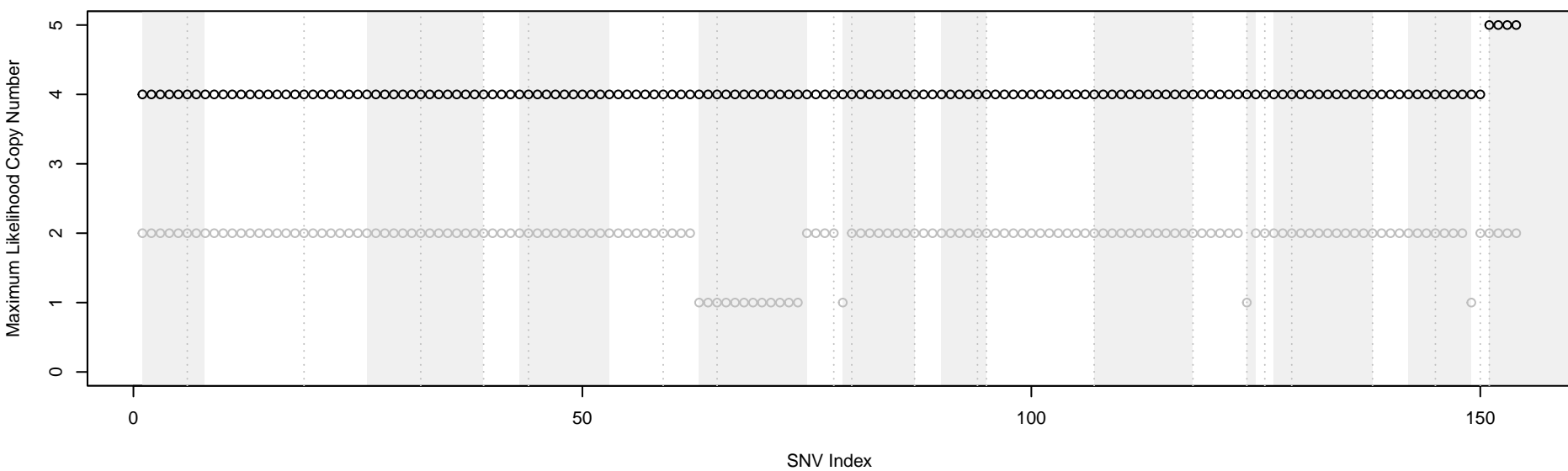
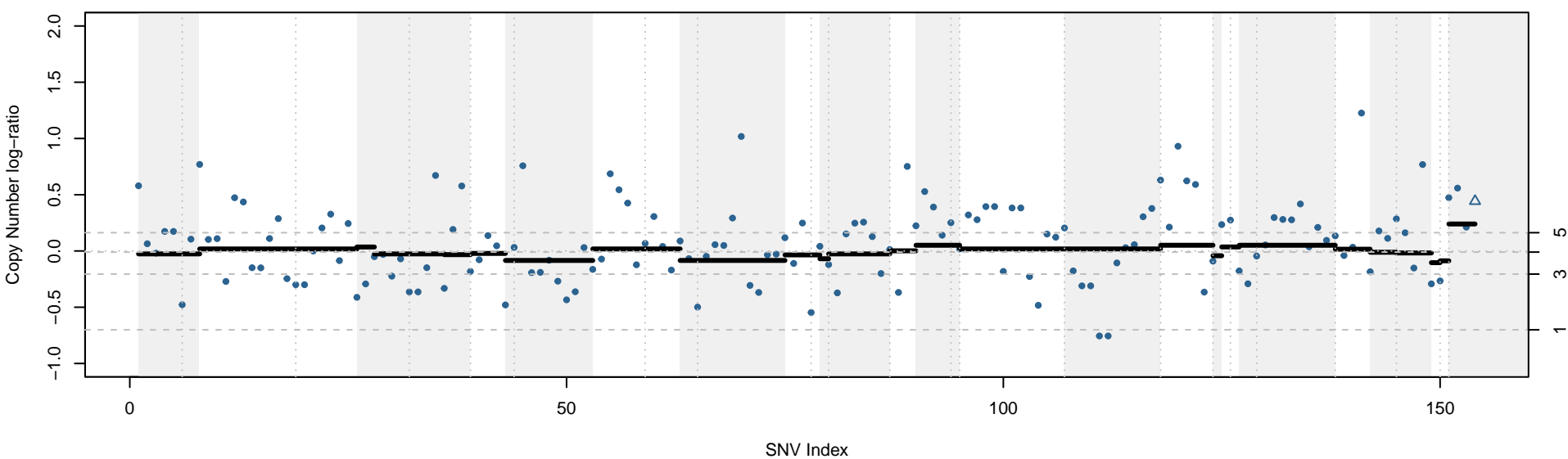
Purity: 0.34 Tumor ploidy: 4.054

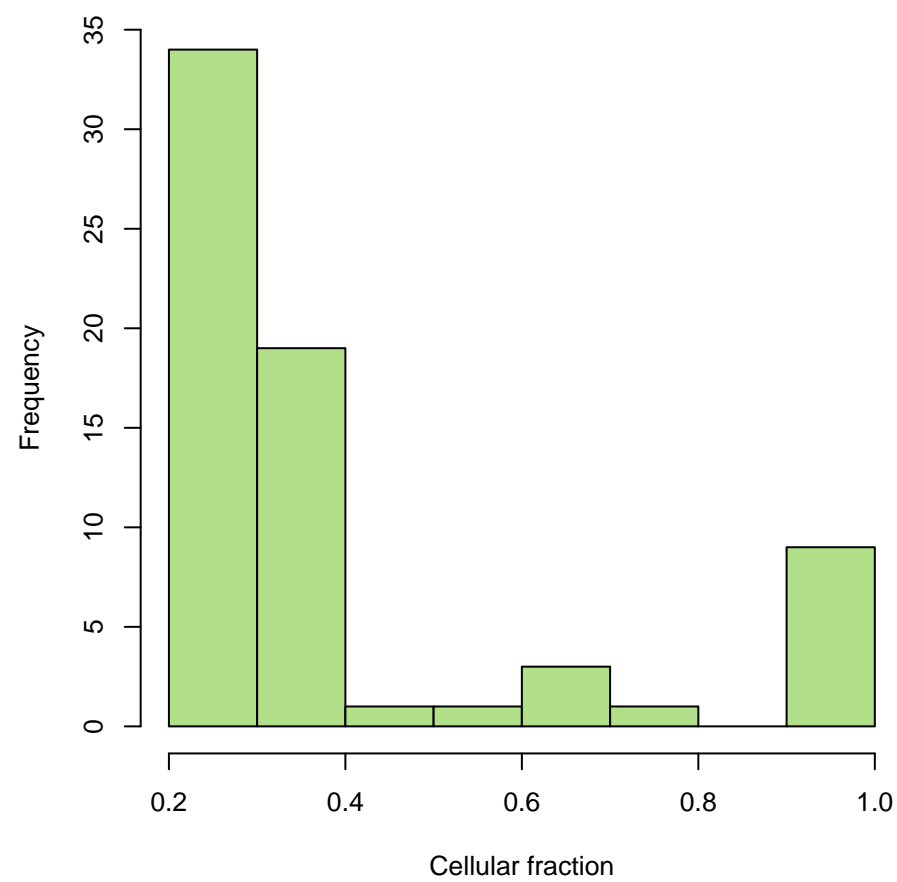
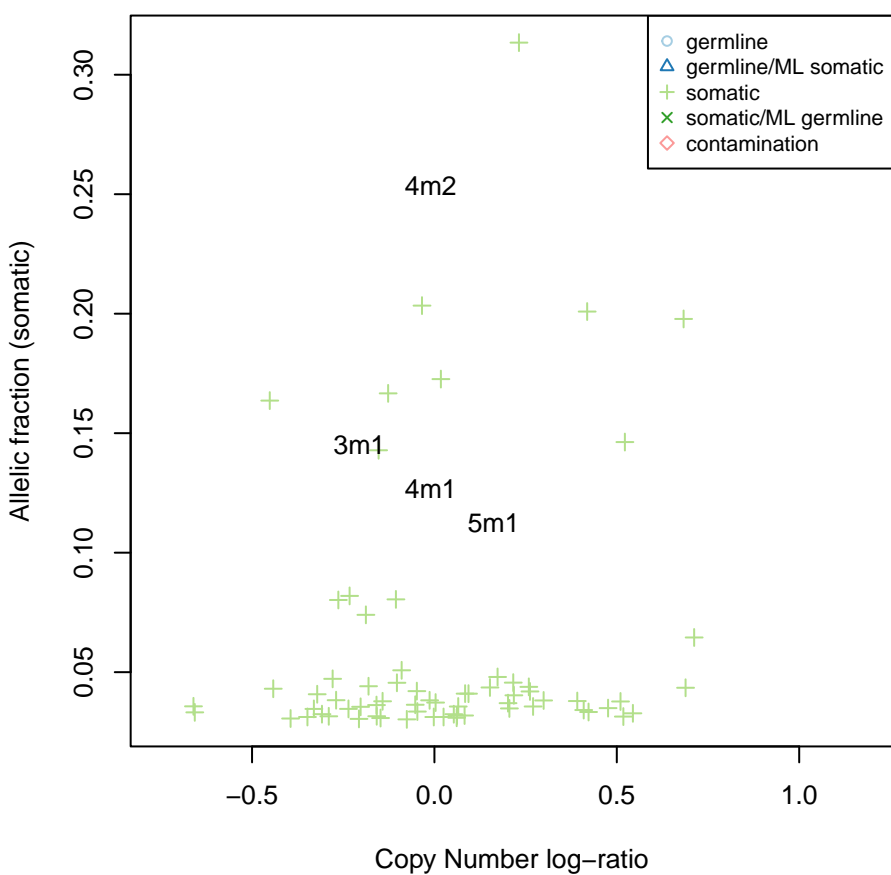
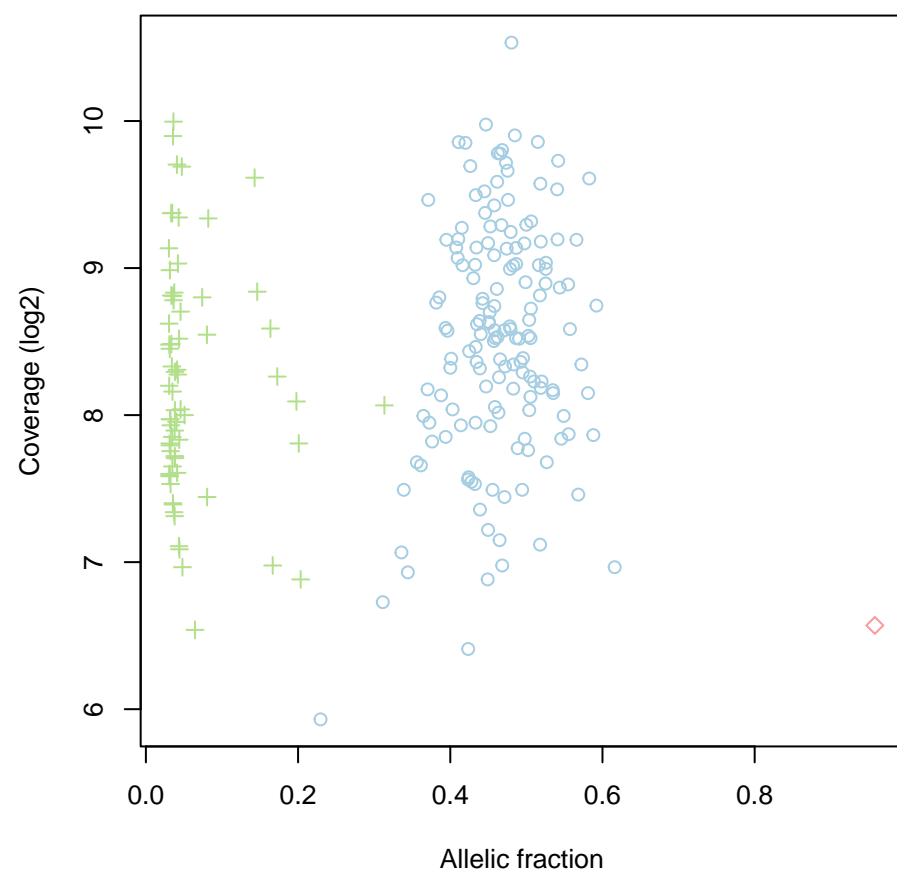
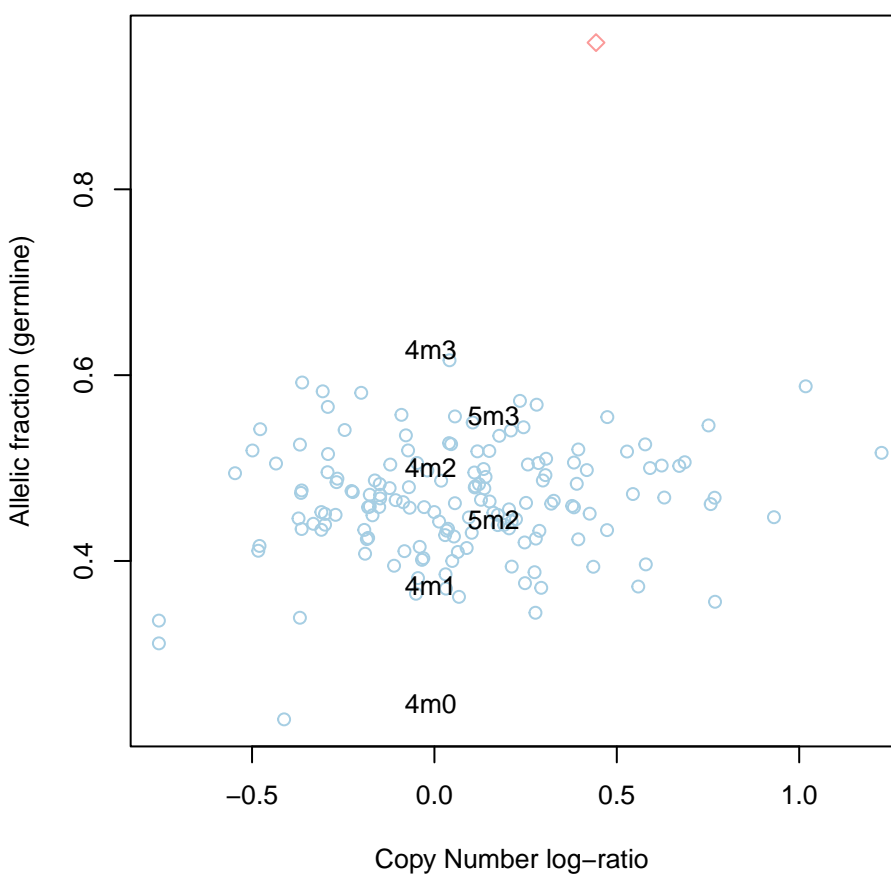


Purity: 0.34 Tumor ploidy: 4.054 SNV log-likelihood: -89.78 GoF: 89.9% Mean coverage: 401;62

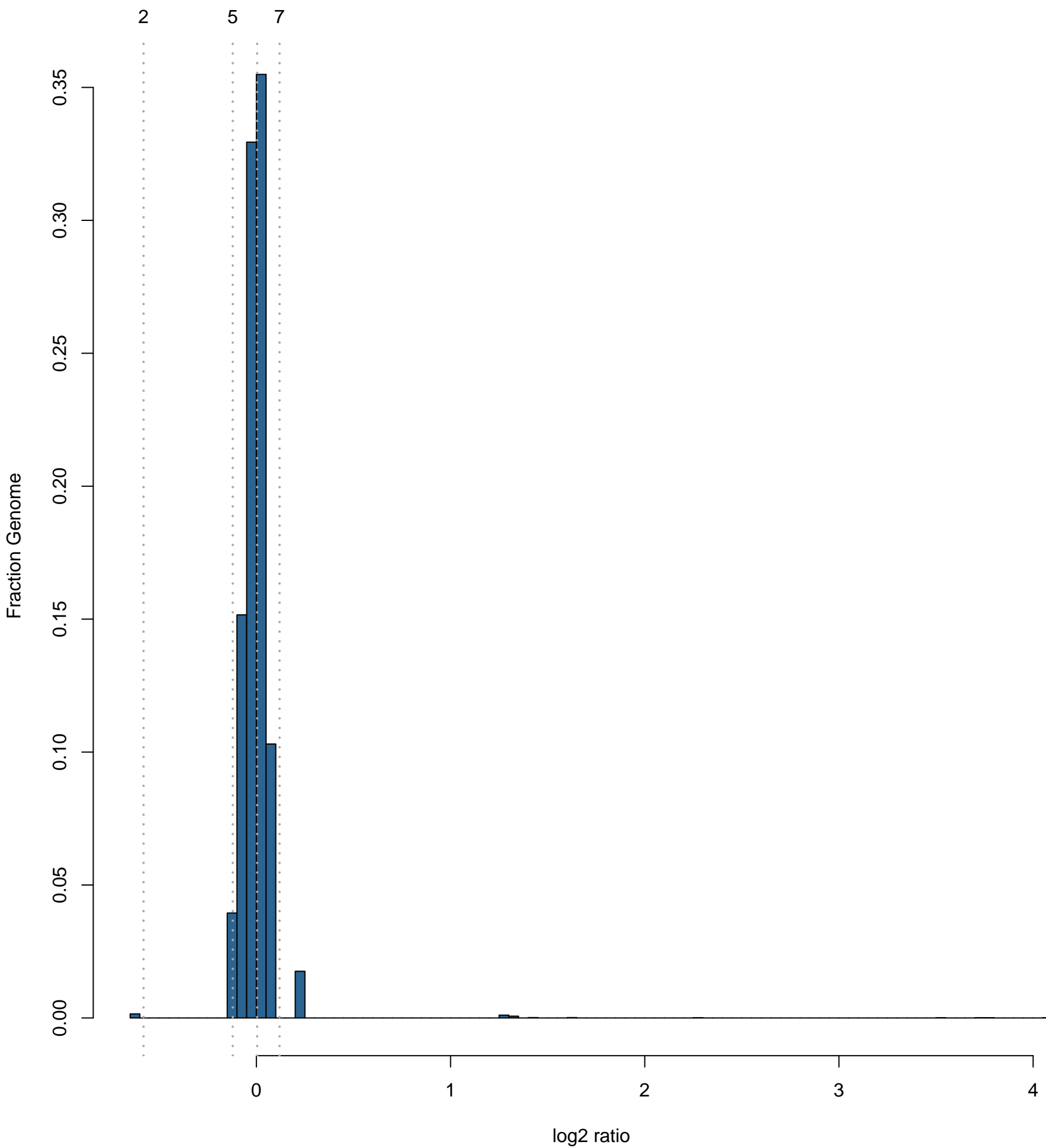


SCNA-fit log-likelihood: -15048.63

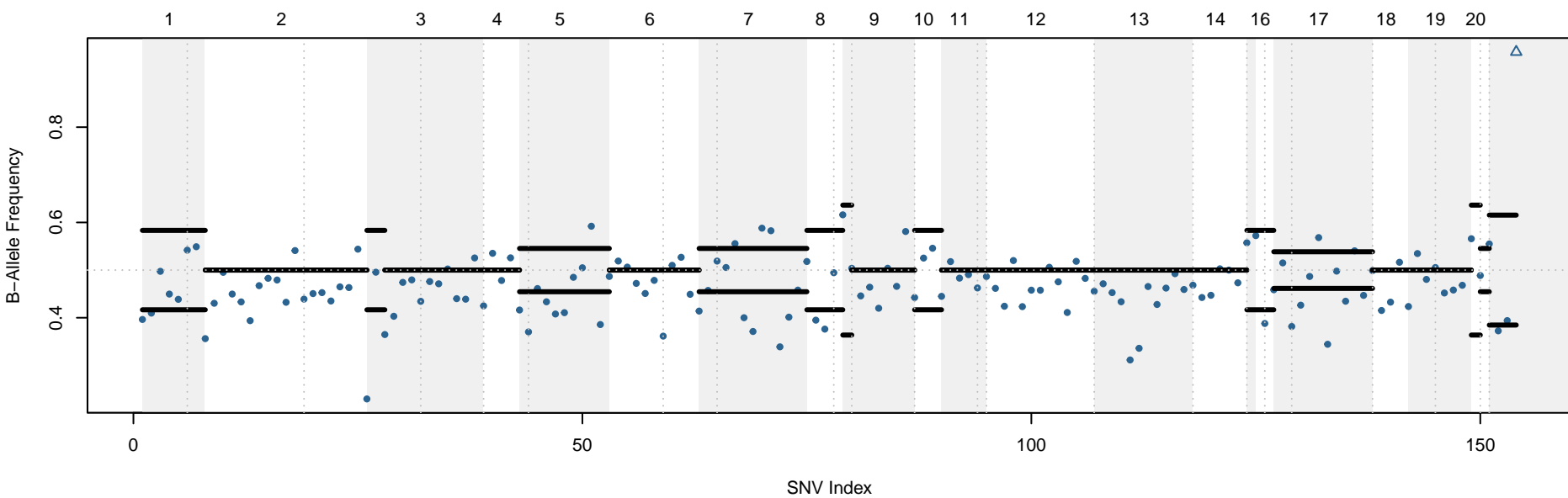




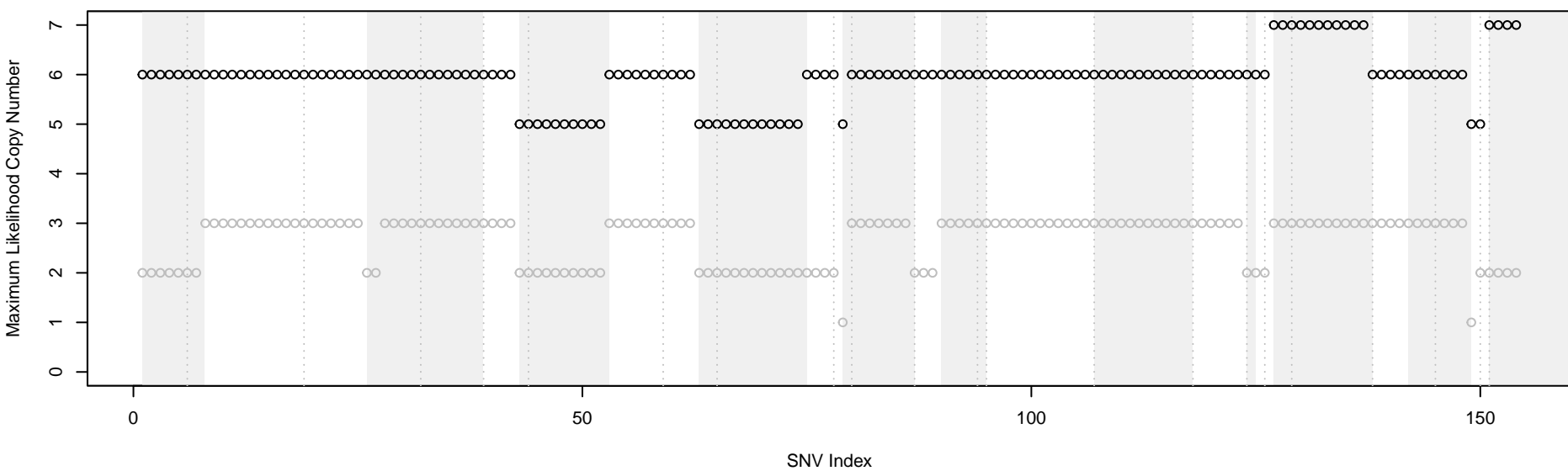
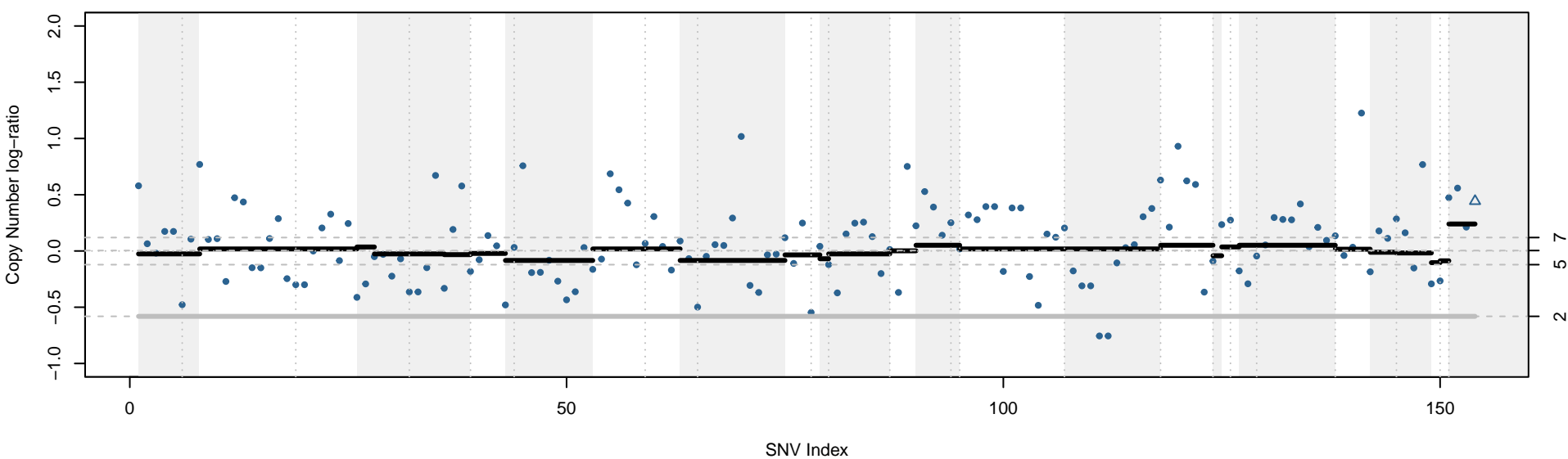
Purity: 0.25 Tumor ploidy: 5.967

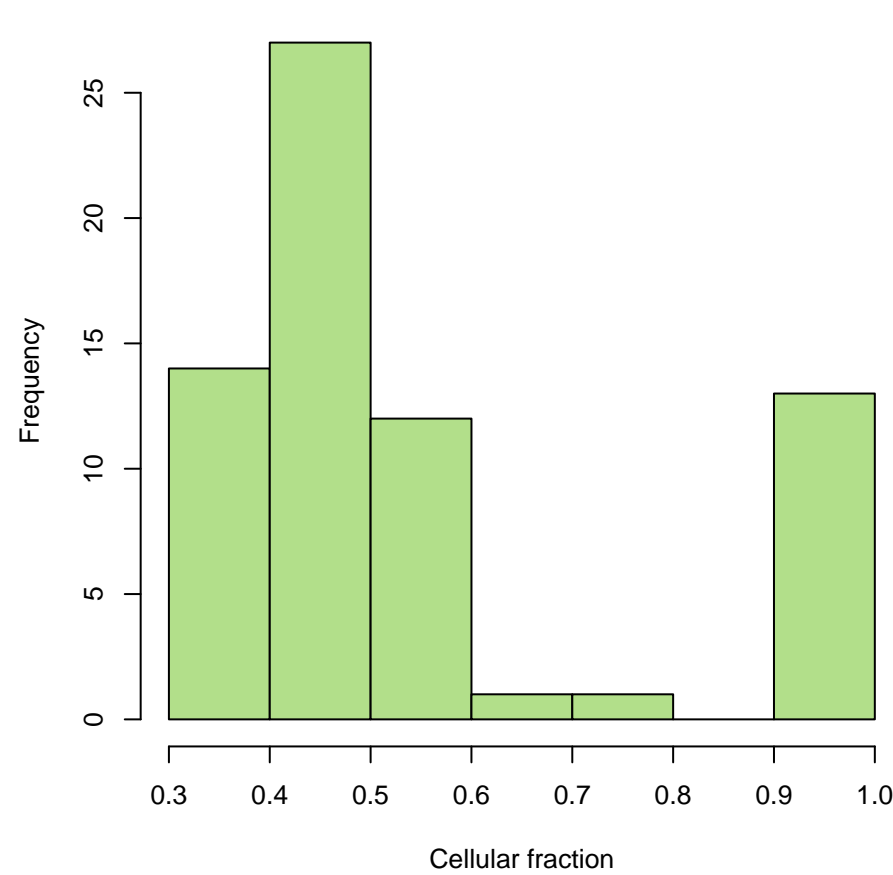
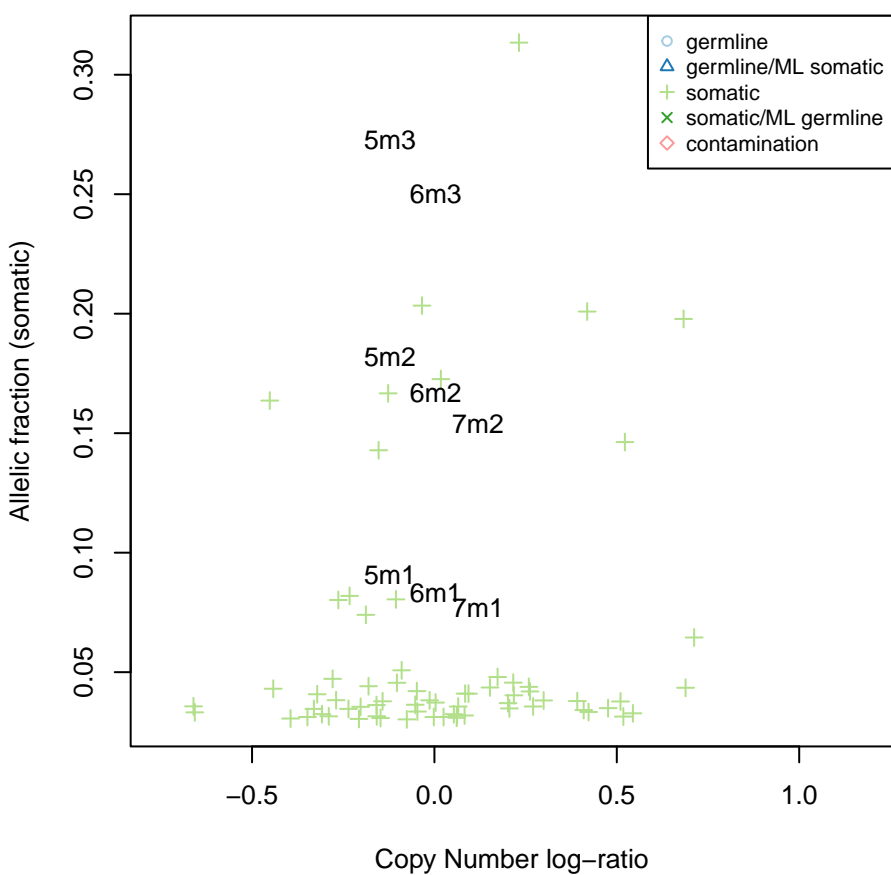
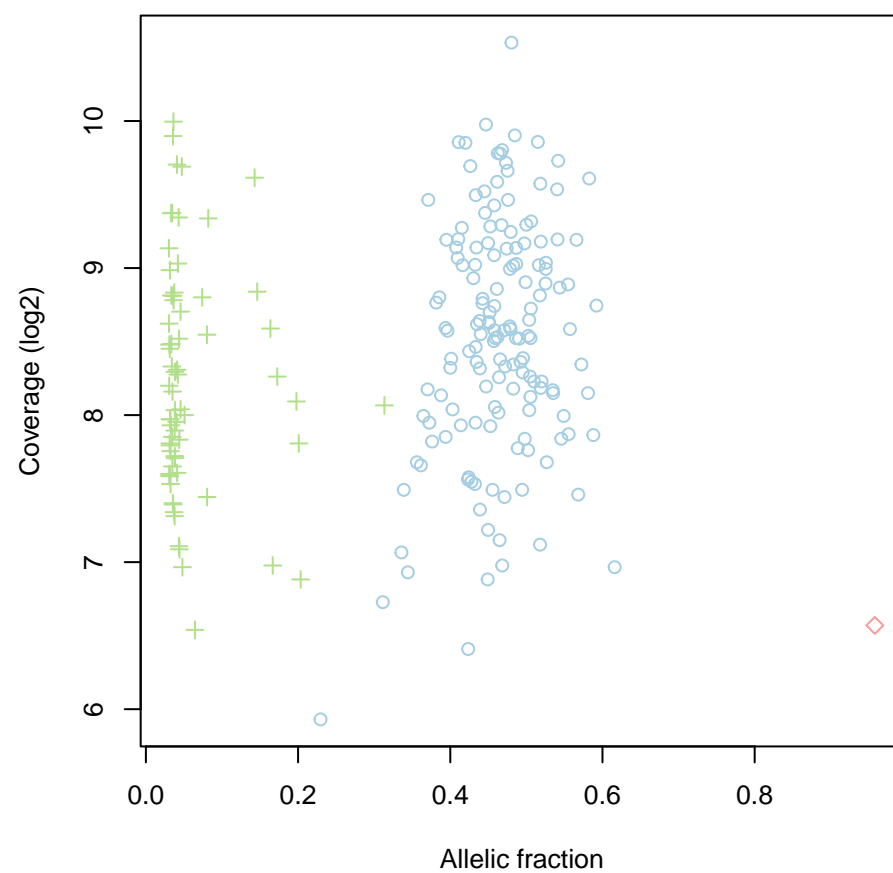
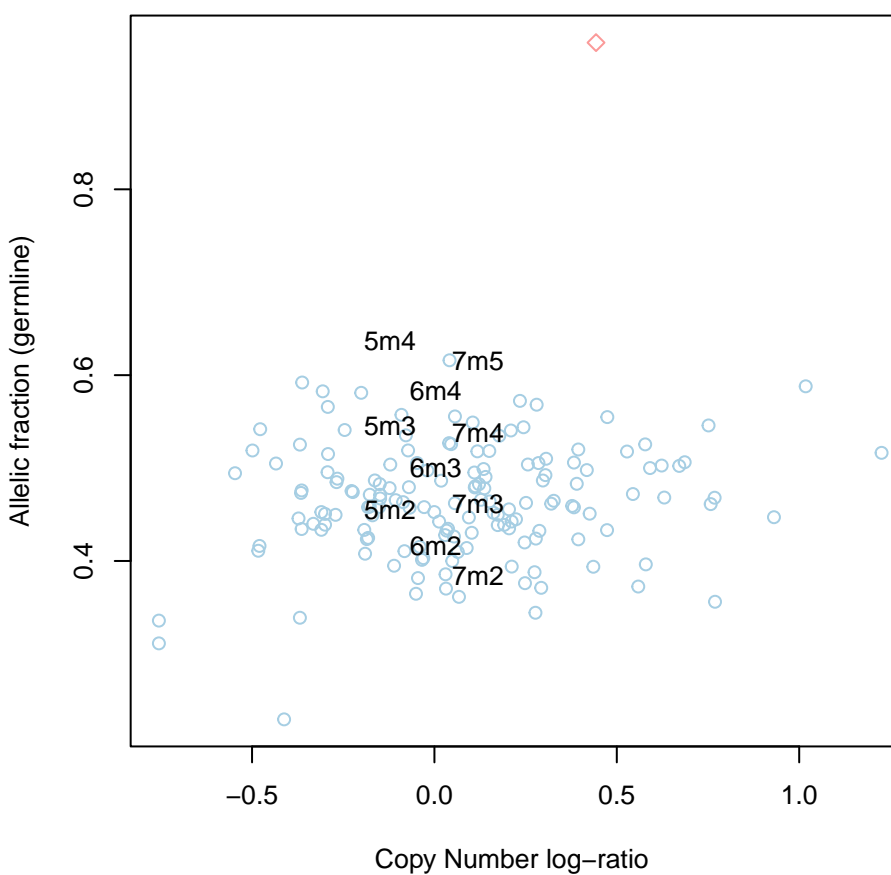


Purity: 0.25 Tumor ploidy: 5.967 SNV log-likelihood: -127.77 GoF: 94.6% Mean coverage: 401;62

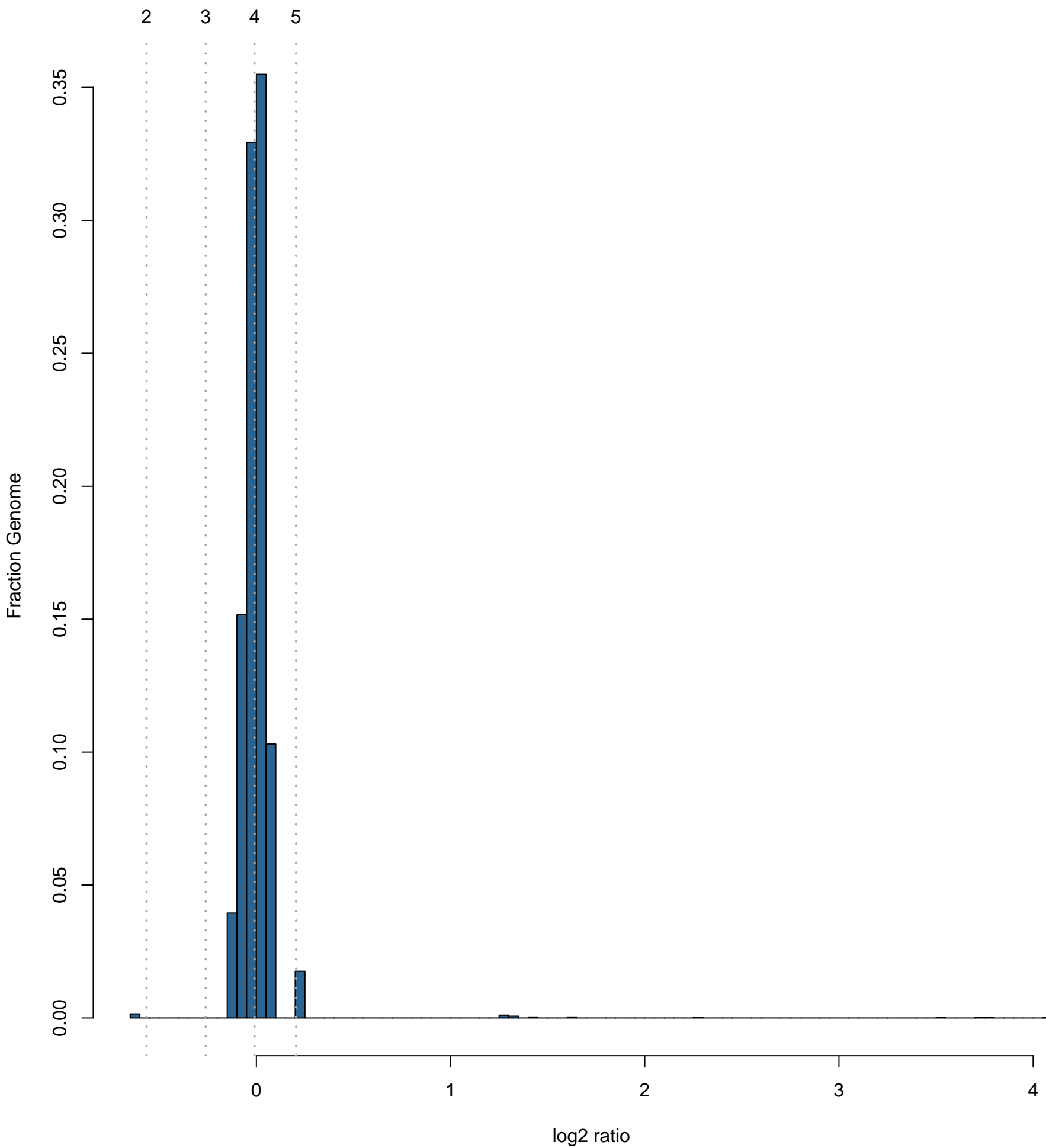


SCNA-fit log-likelihood: -15012.86

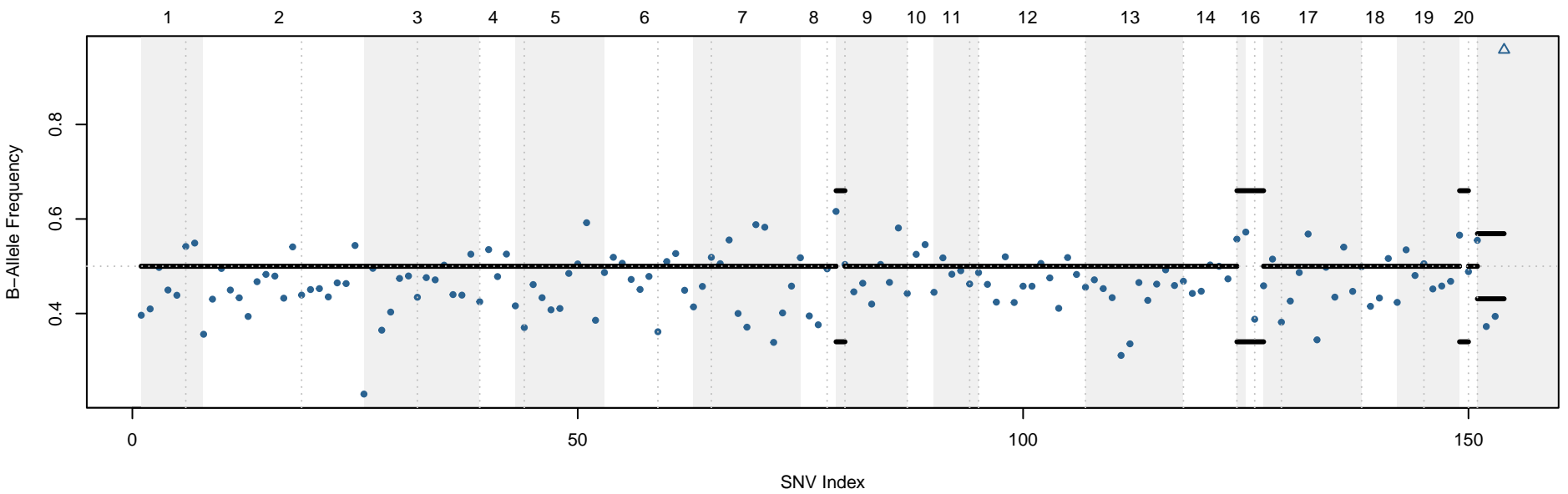




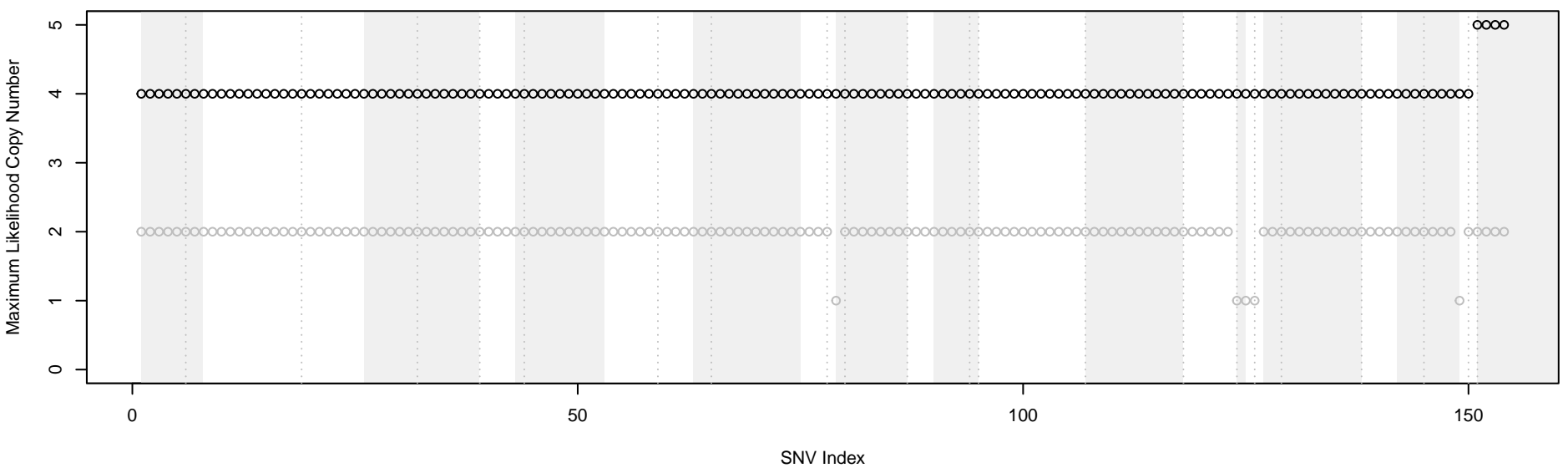
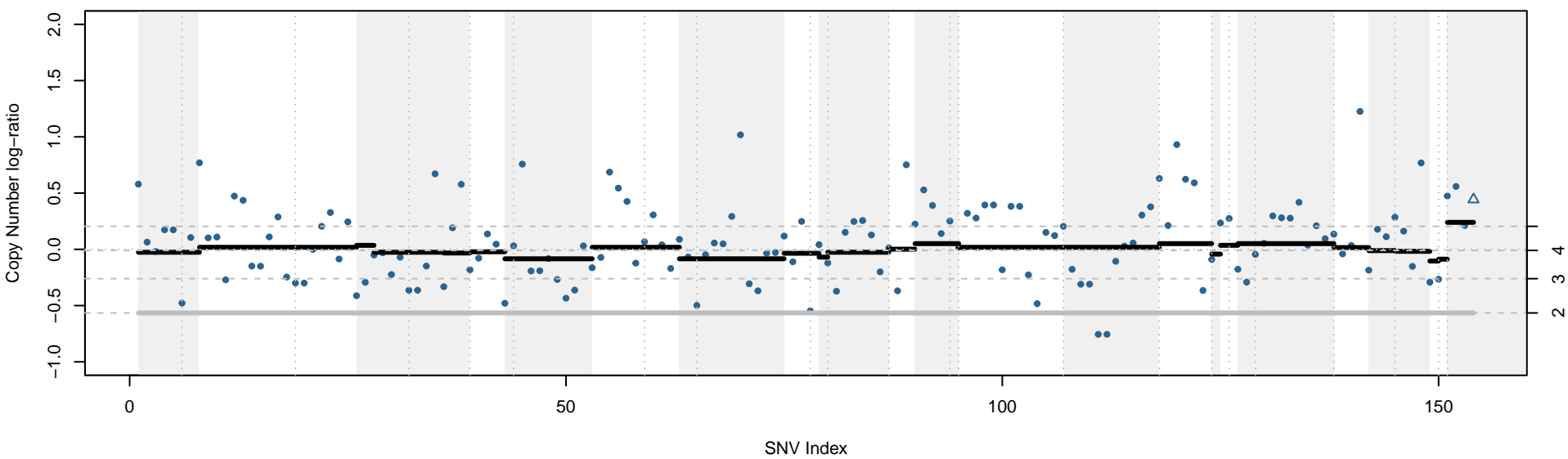
Purity: 0.47 Tumor ploidy: 4.042

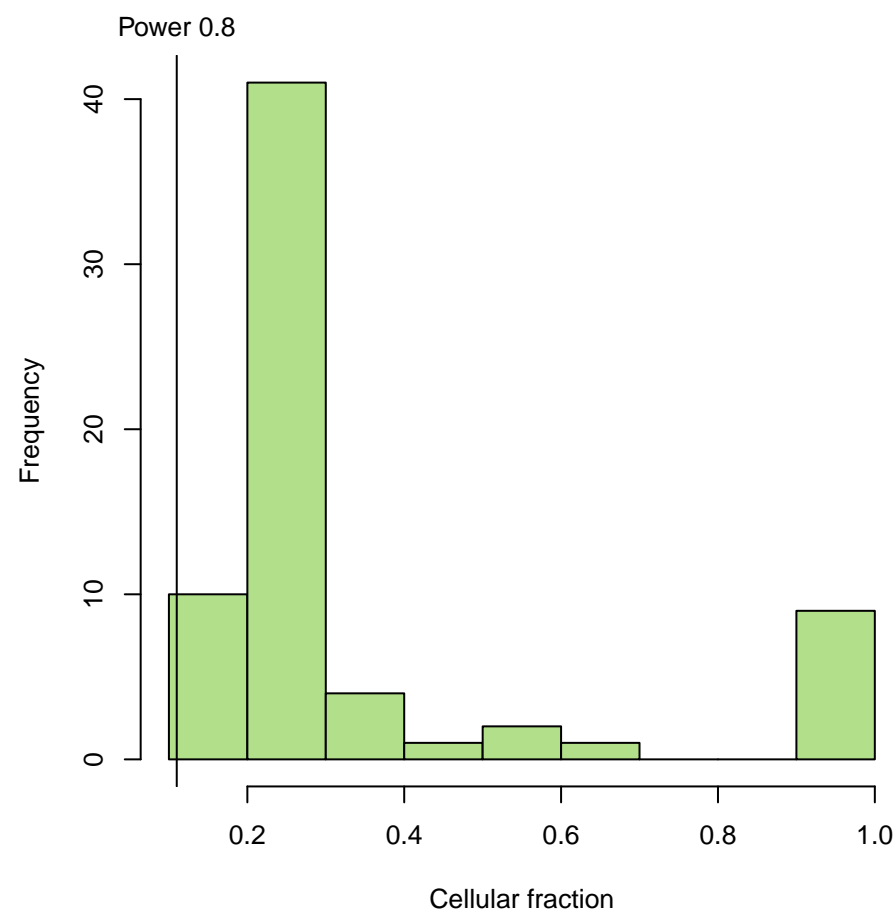
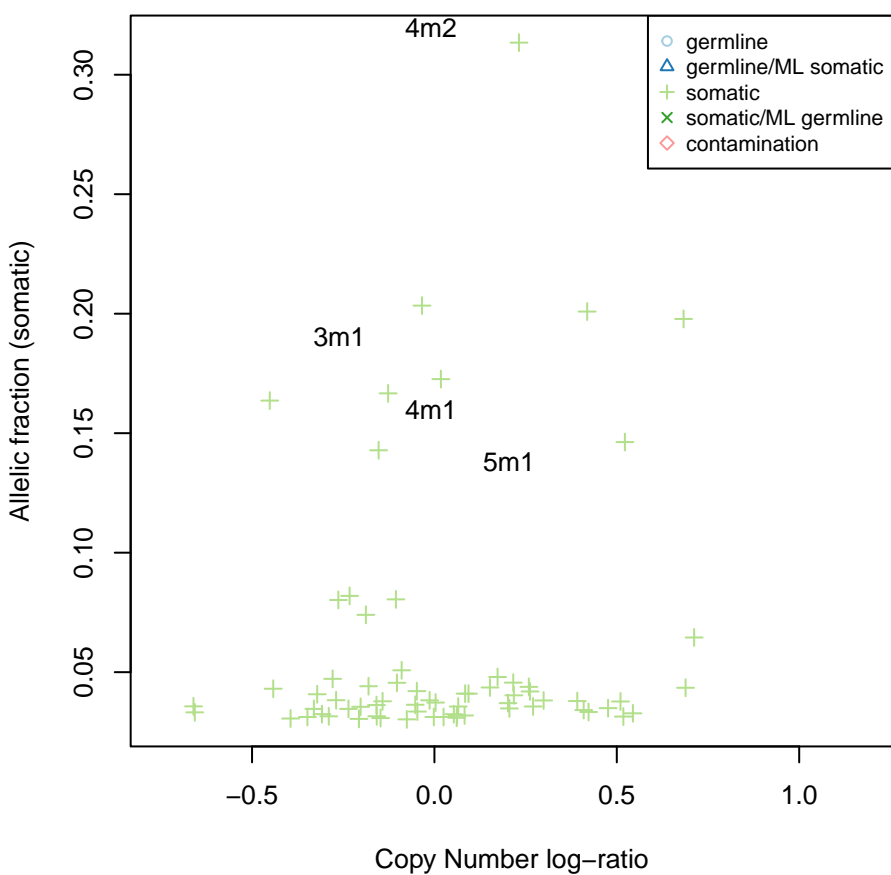
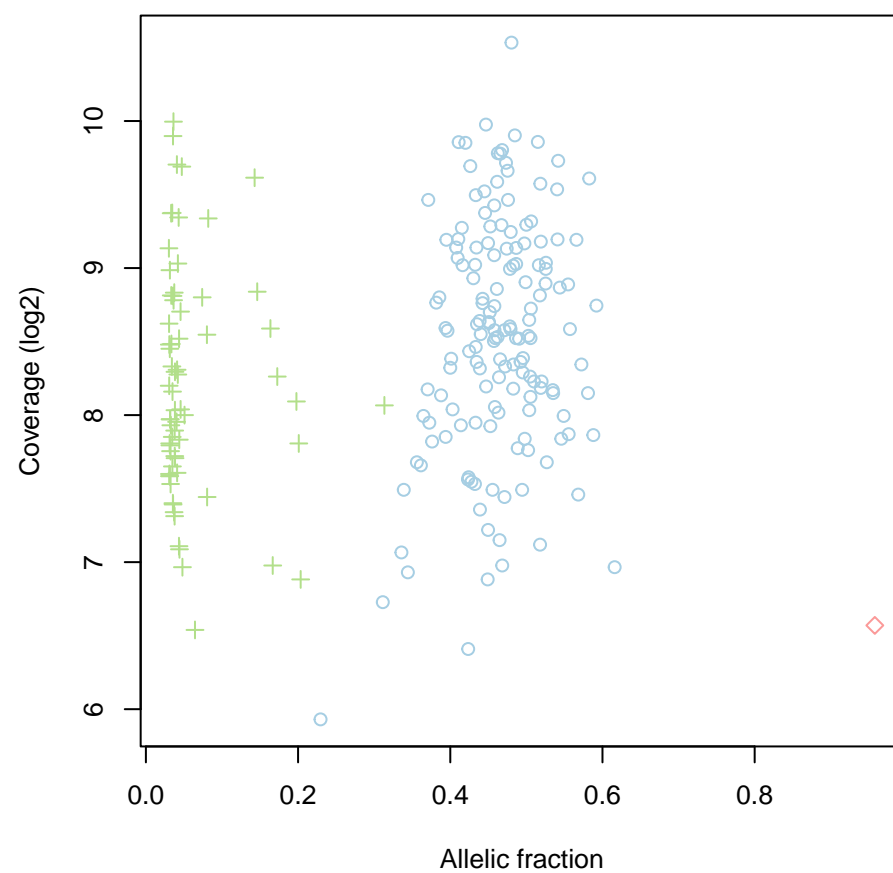
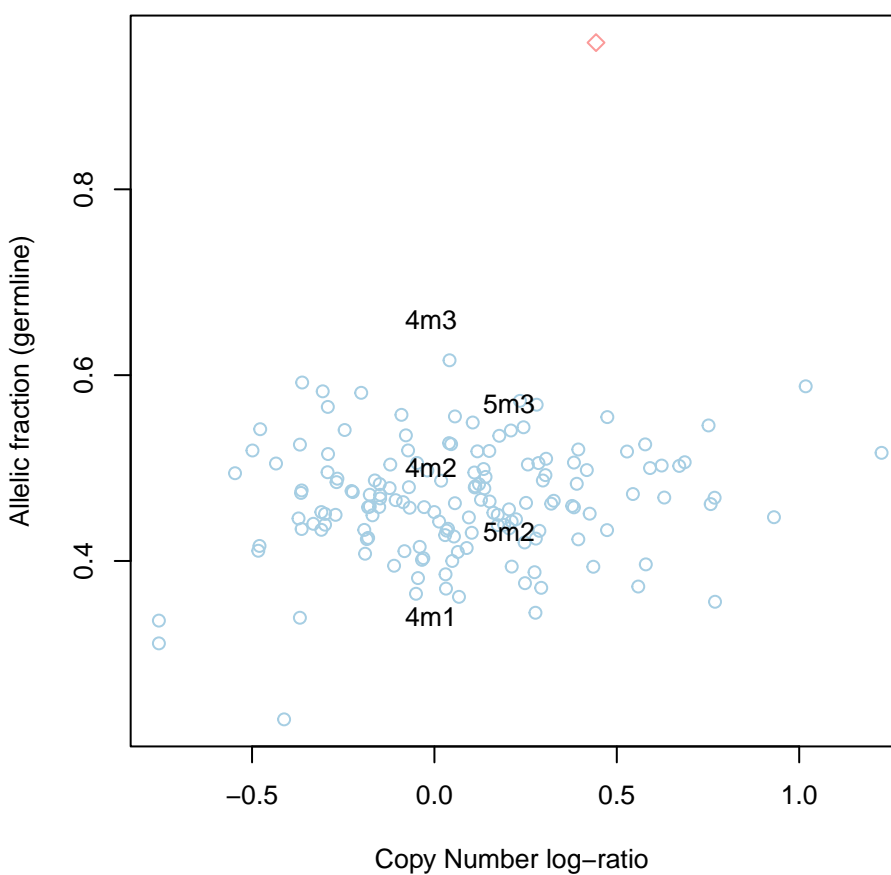


Purity: 0.47 Tumor ploidy: 4.042 SNV log-likelihood: -130.81 GoF: 84.3% Mean coverage: 401;62

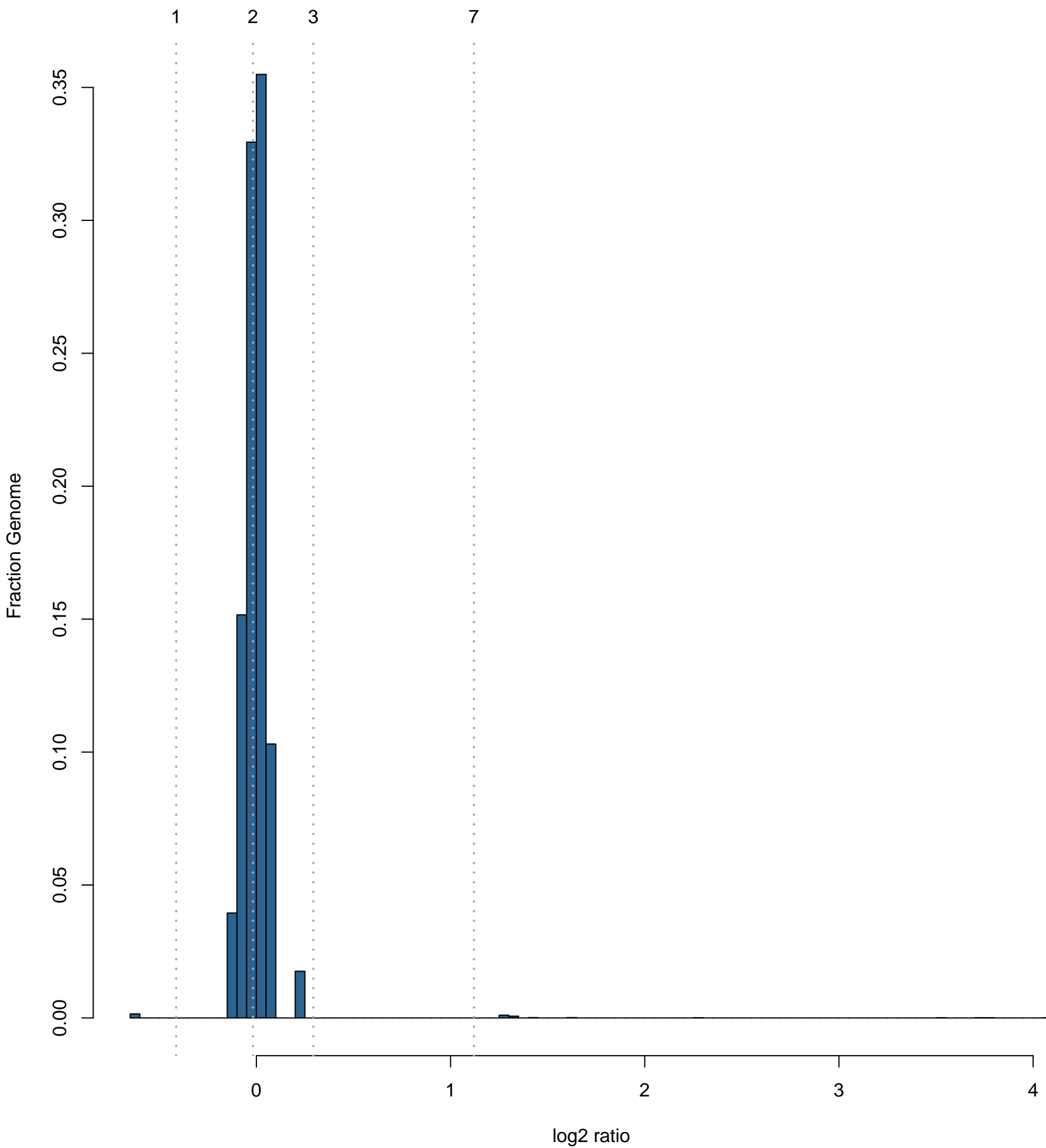


SCNA-fit log-likelihood: -15057.63

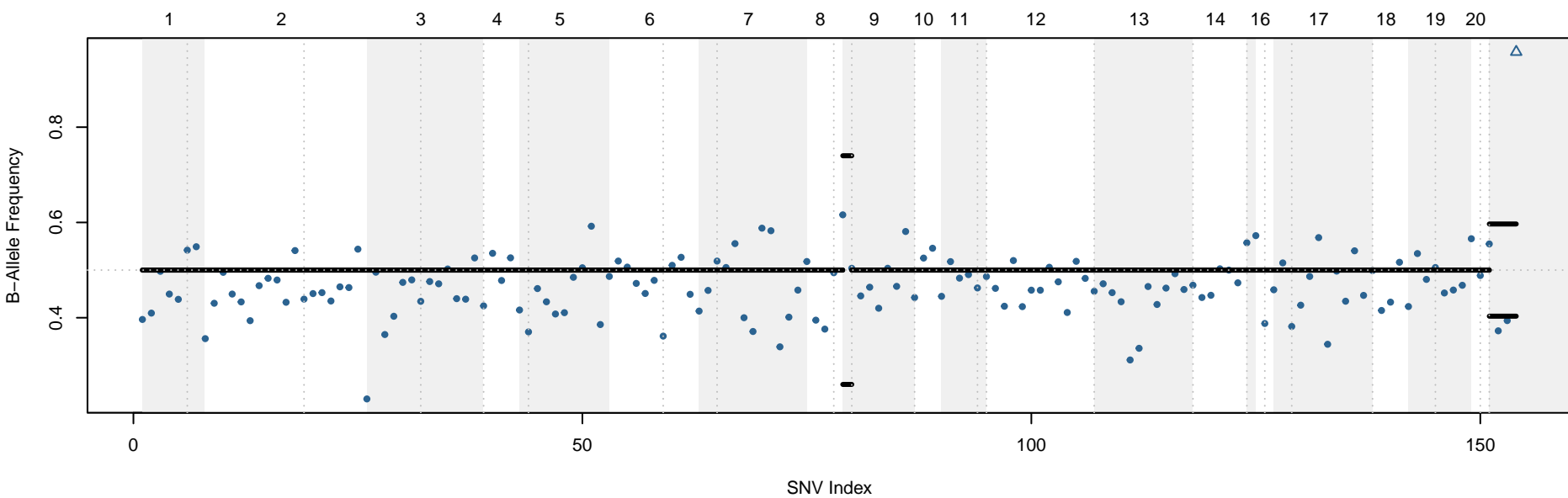




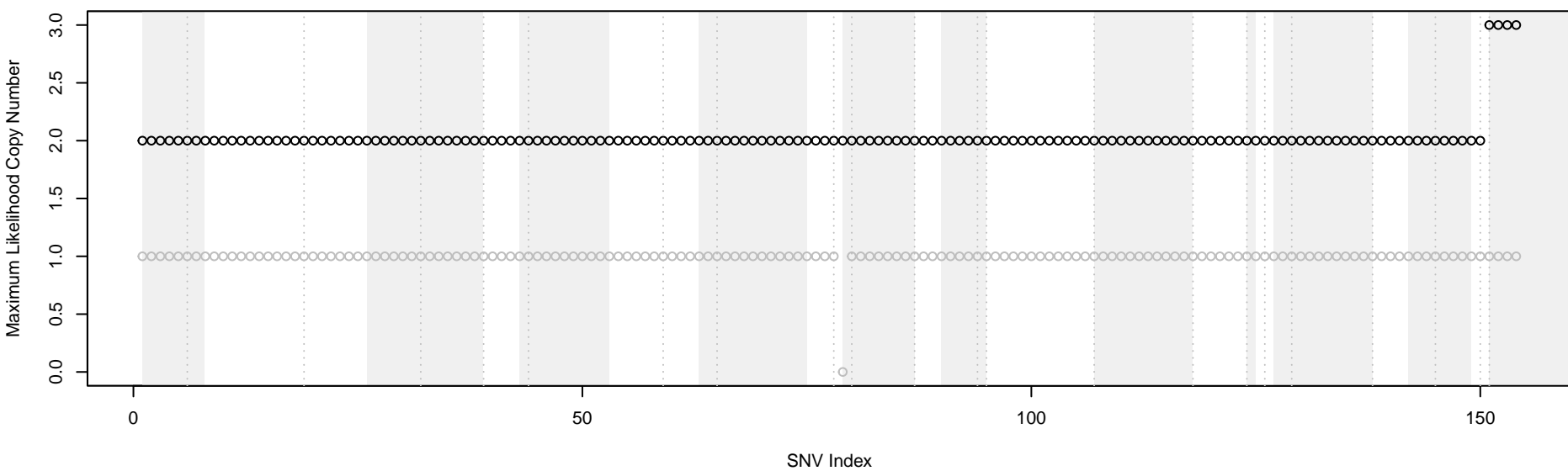
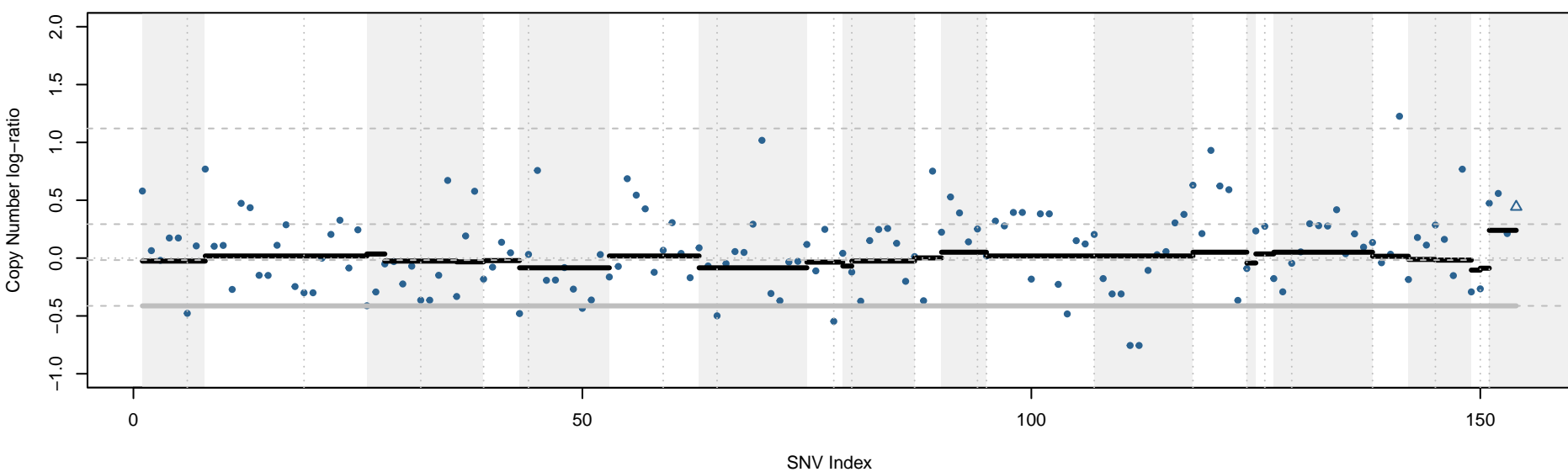
Purity: 0.48 Tumor ploidy: 2.05

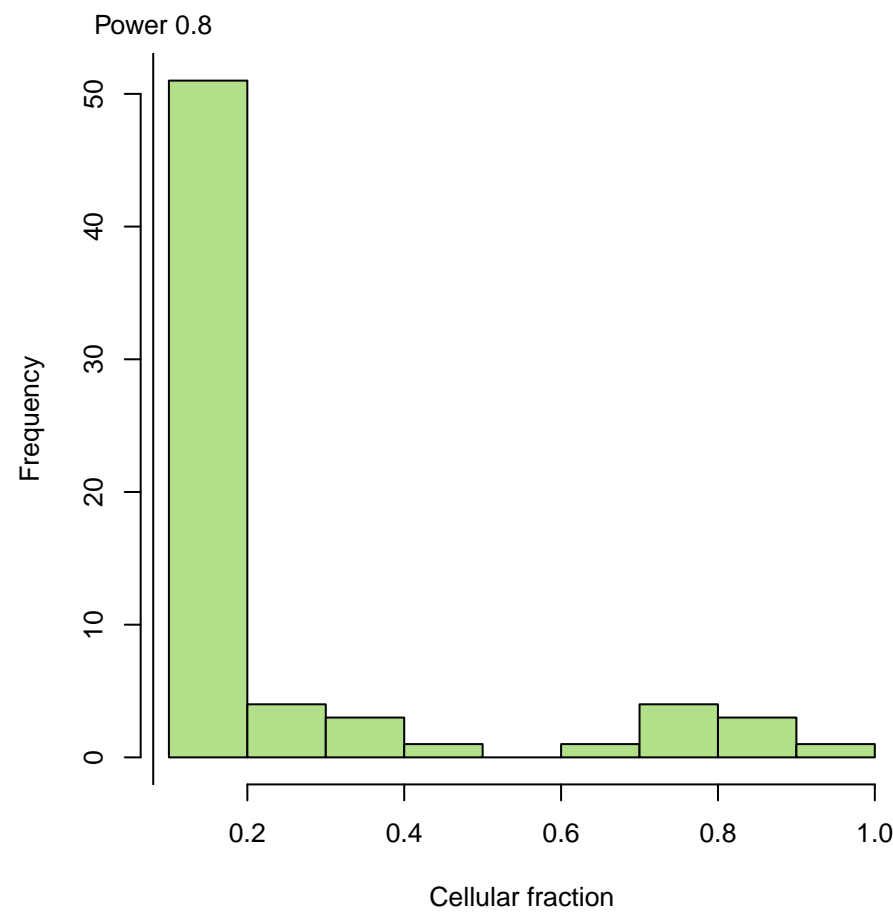
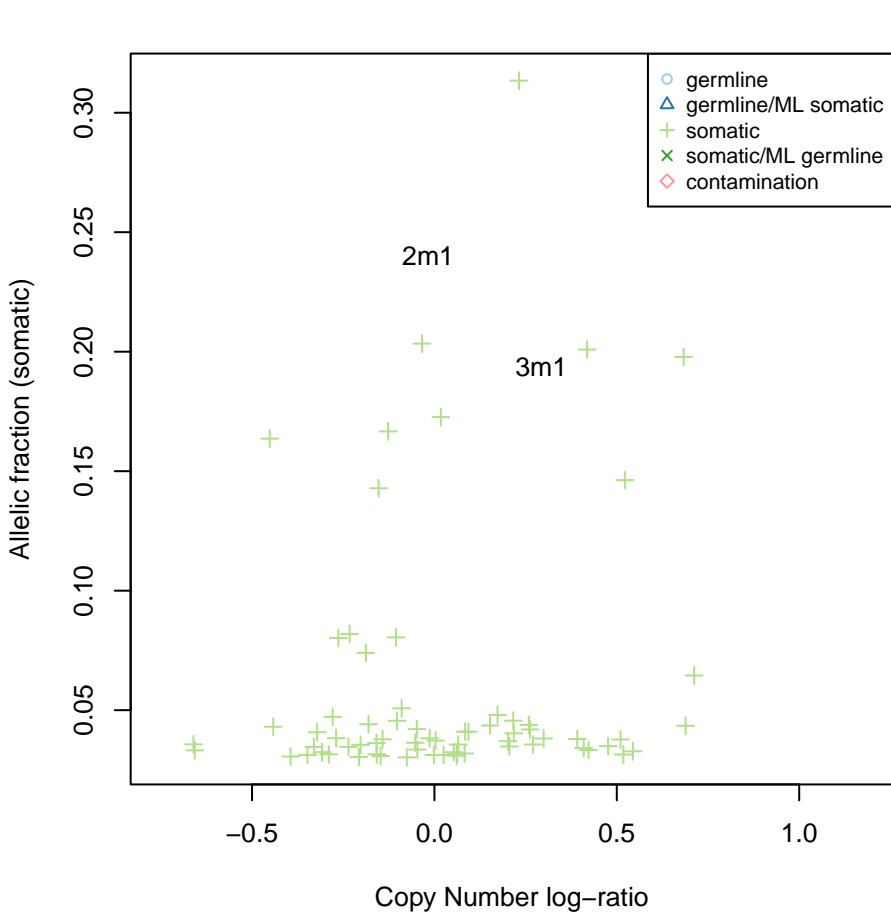
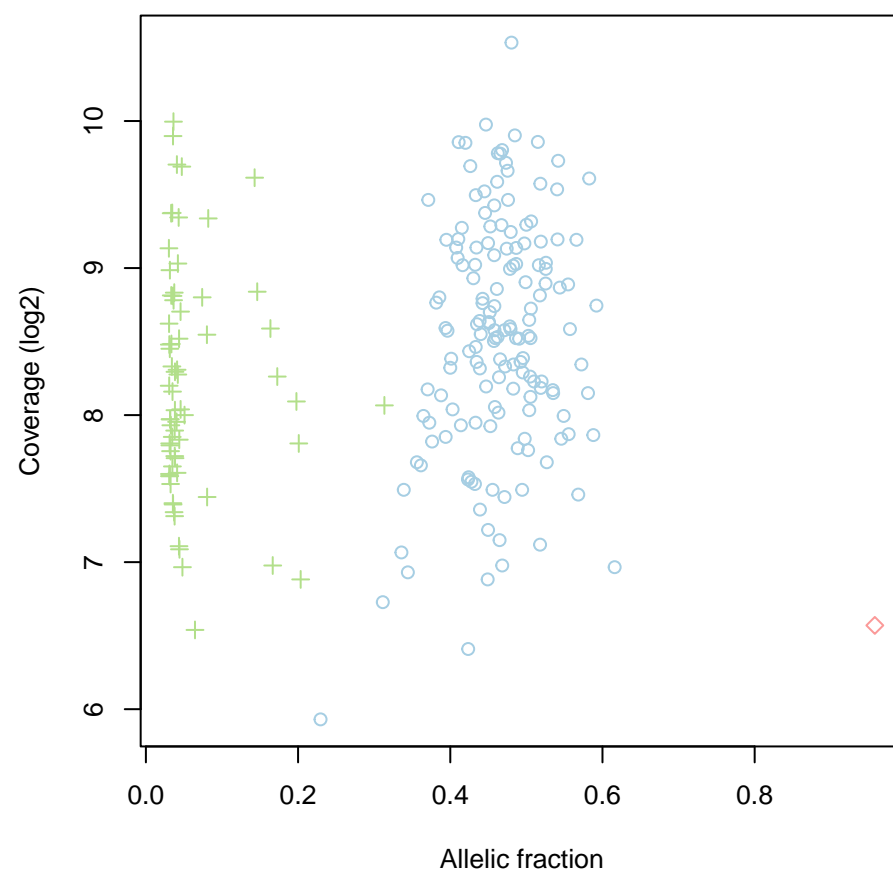
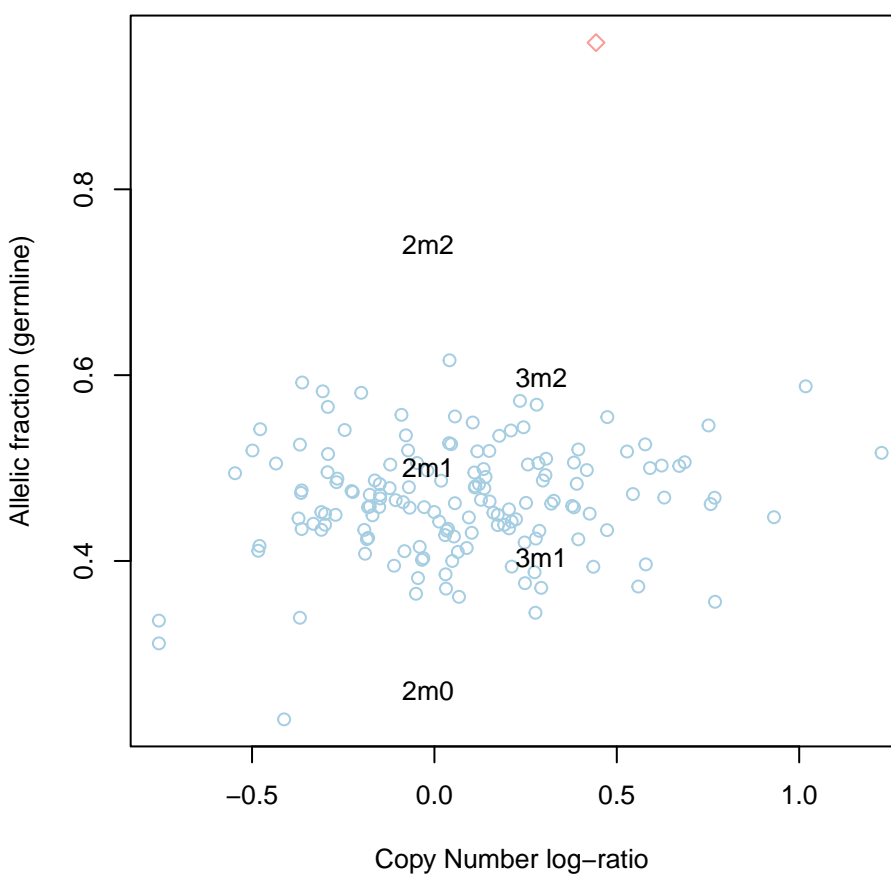


Purity: 0.48 Tumor ploidy: 2.05 SNV log-likelihood: -155.12 GoF: 68.1% Mean coverage: 401;62

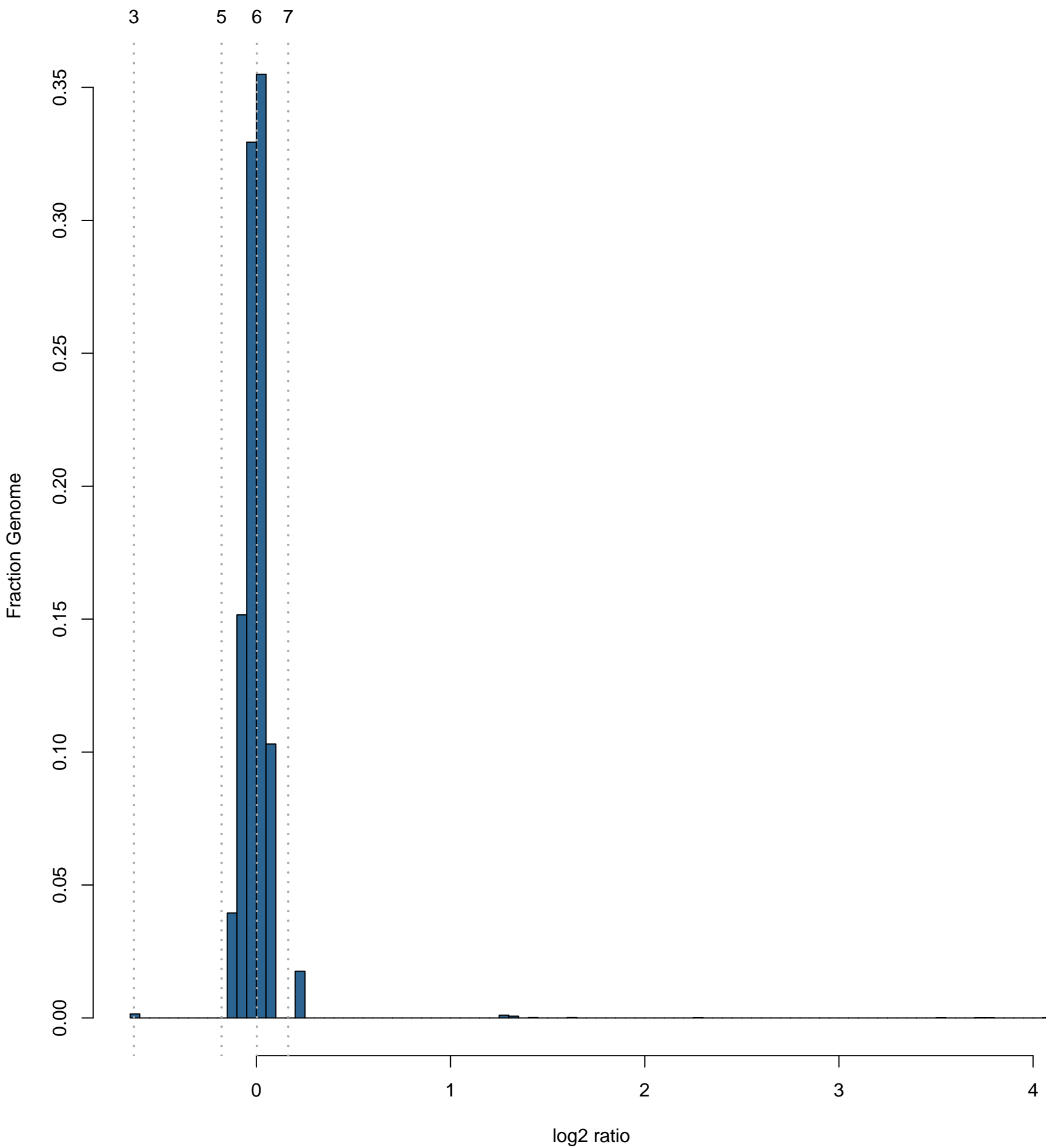


SCNA-fit log-likelihood: -15045.96

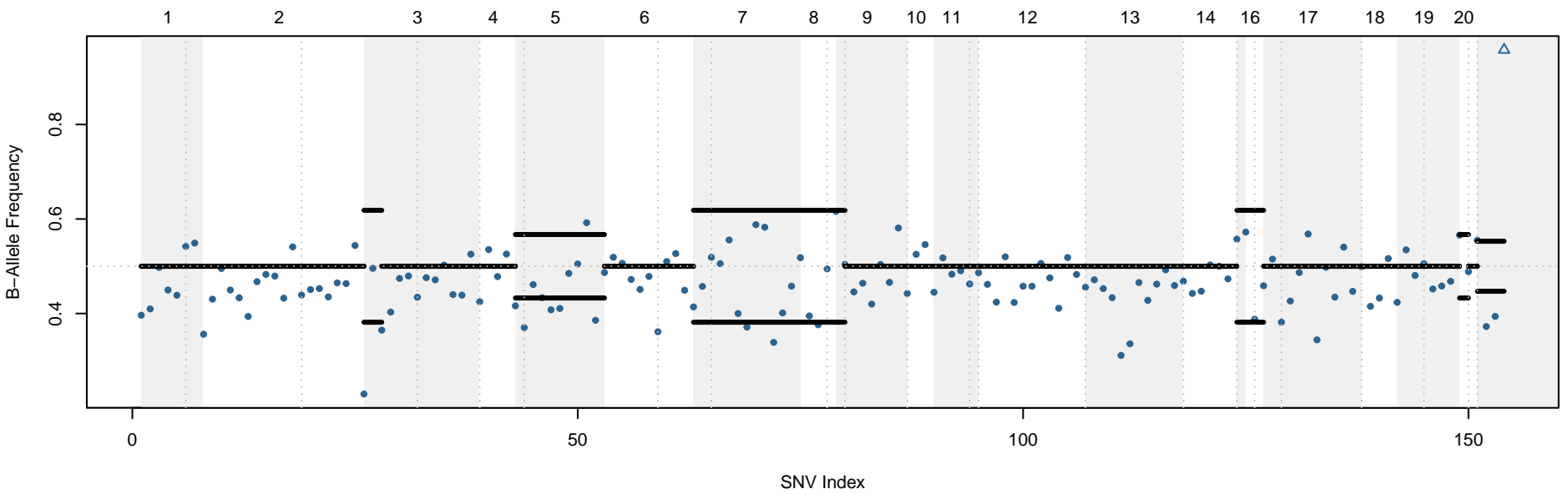




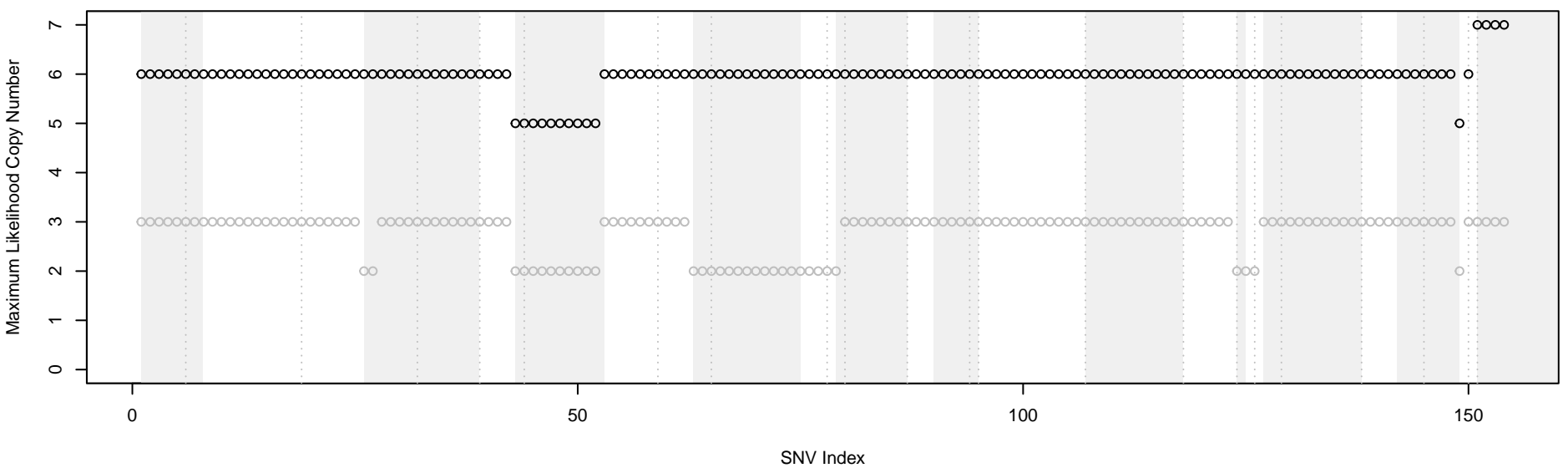
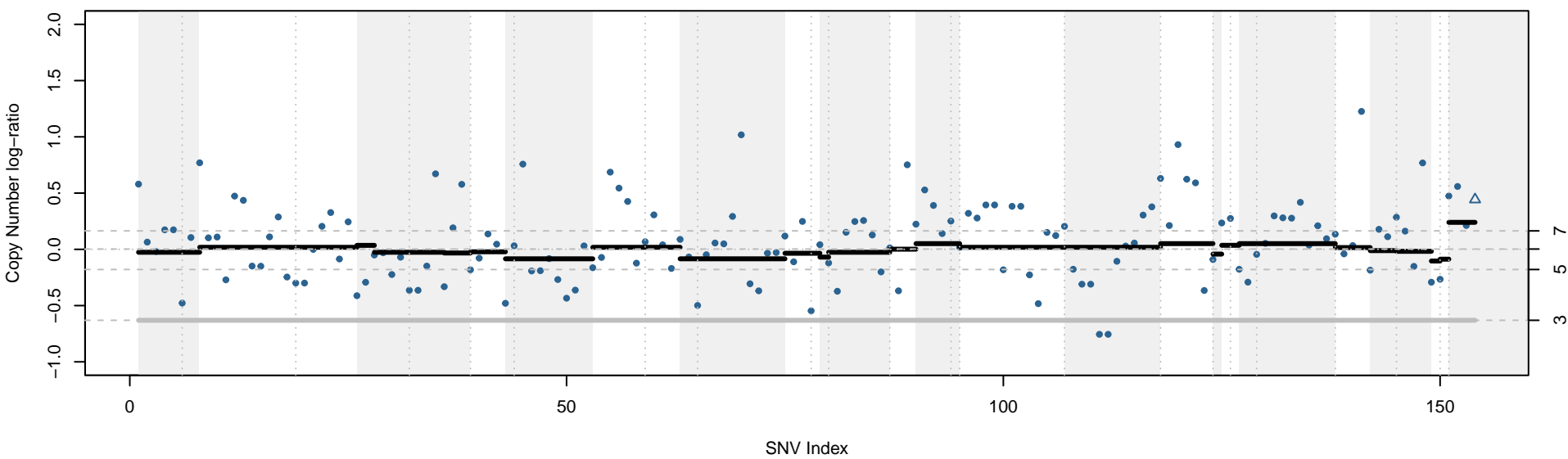
Purity: 0.45 Tumor ploidy: 5.985

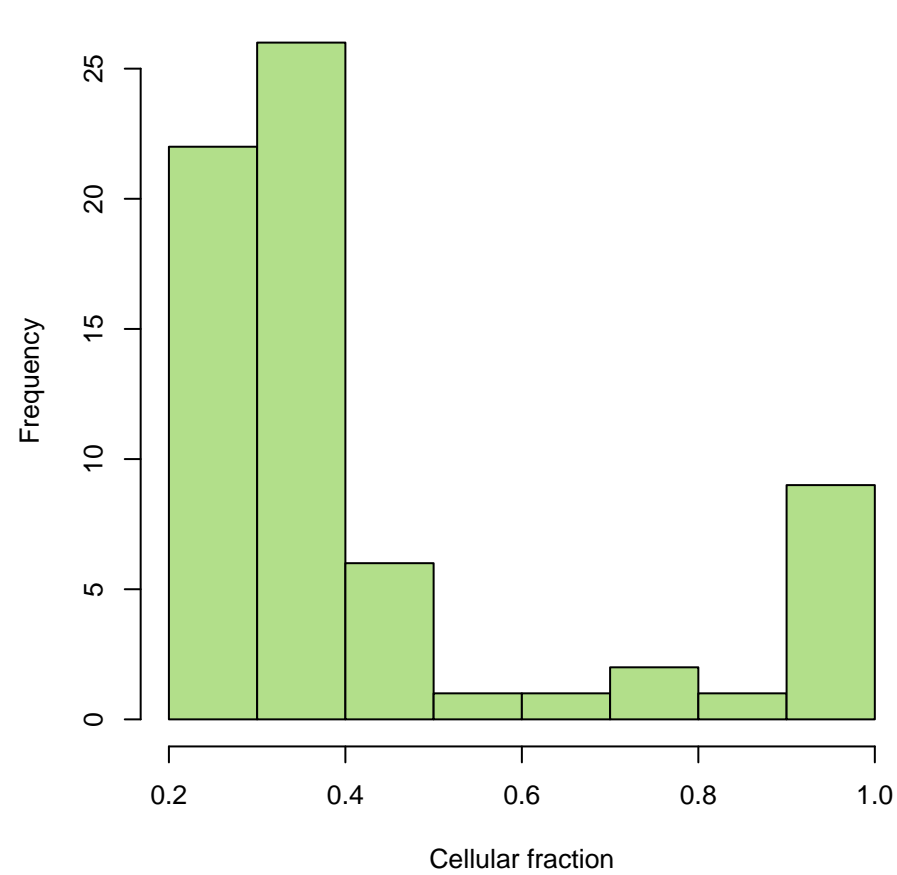
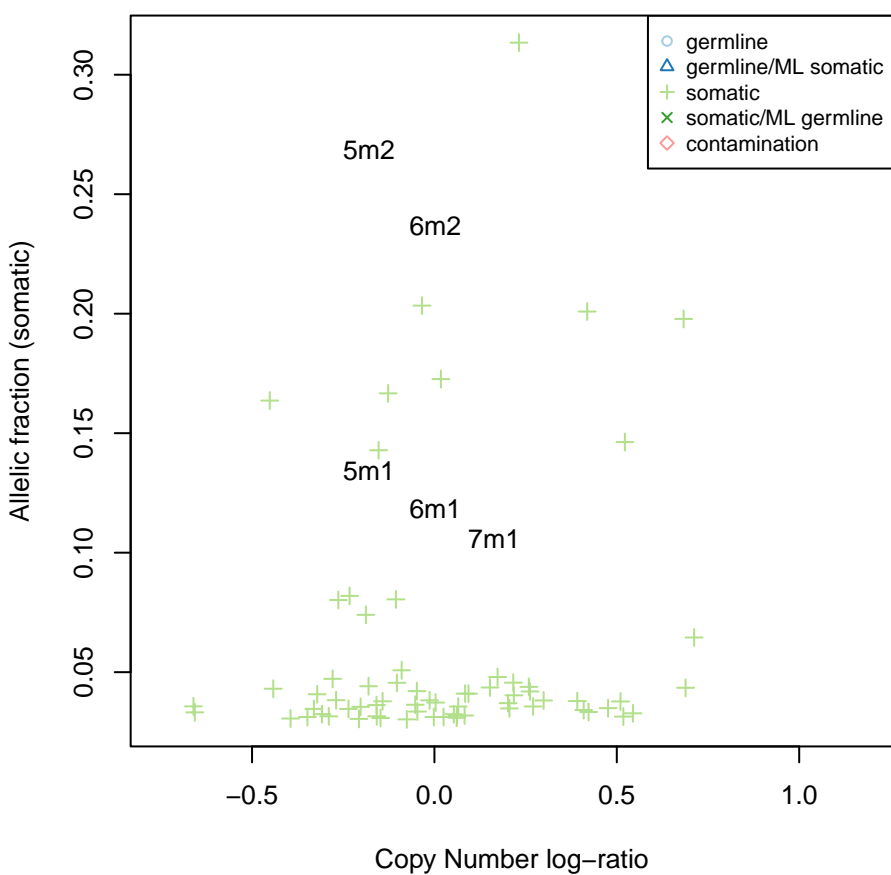
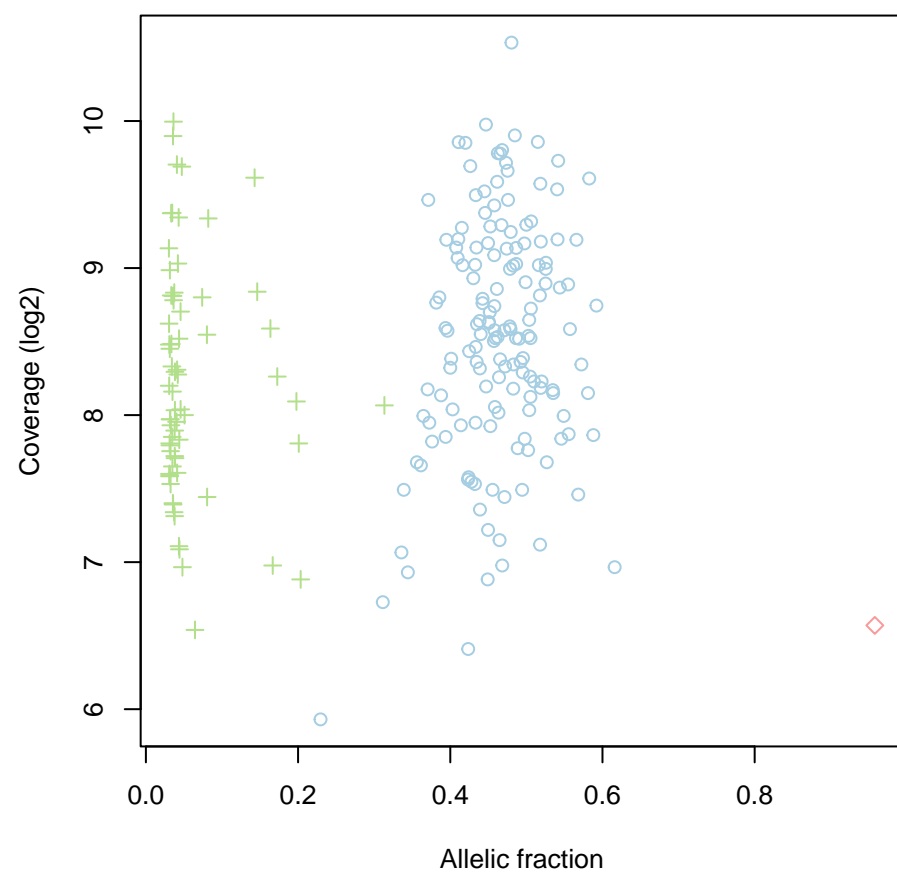
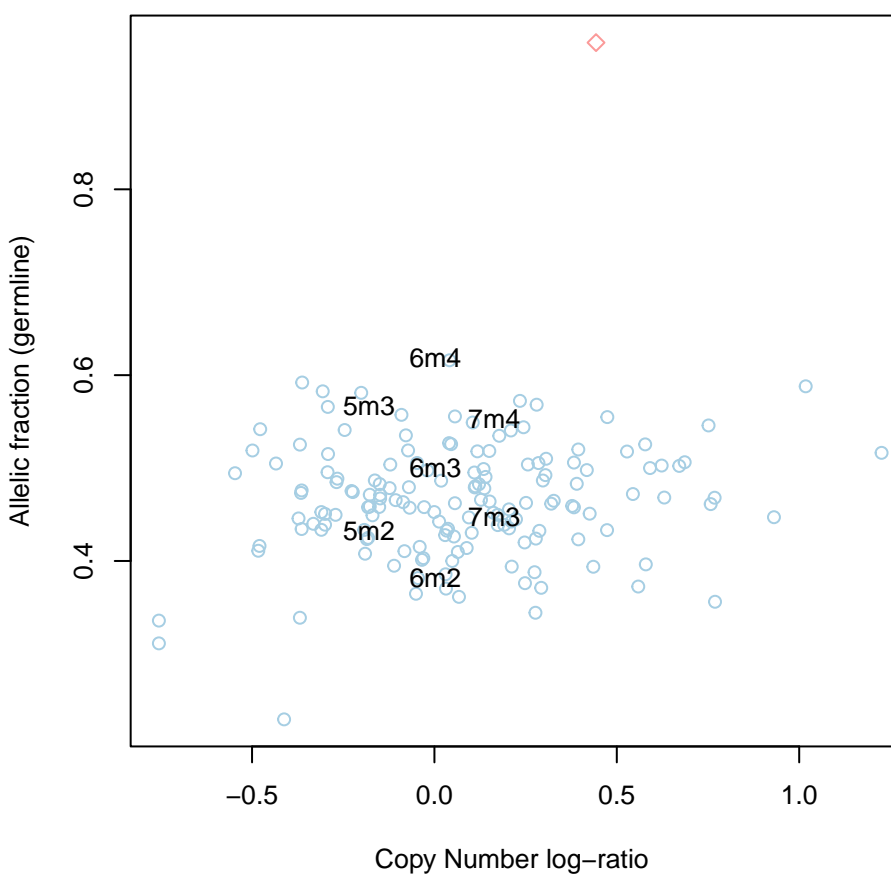


Purity: 0.45 Tumor ploidy: 5.985 SNV log-likelihood: -148.96 GoF: 90.9% Mean coverage: 401;62

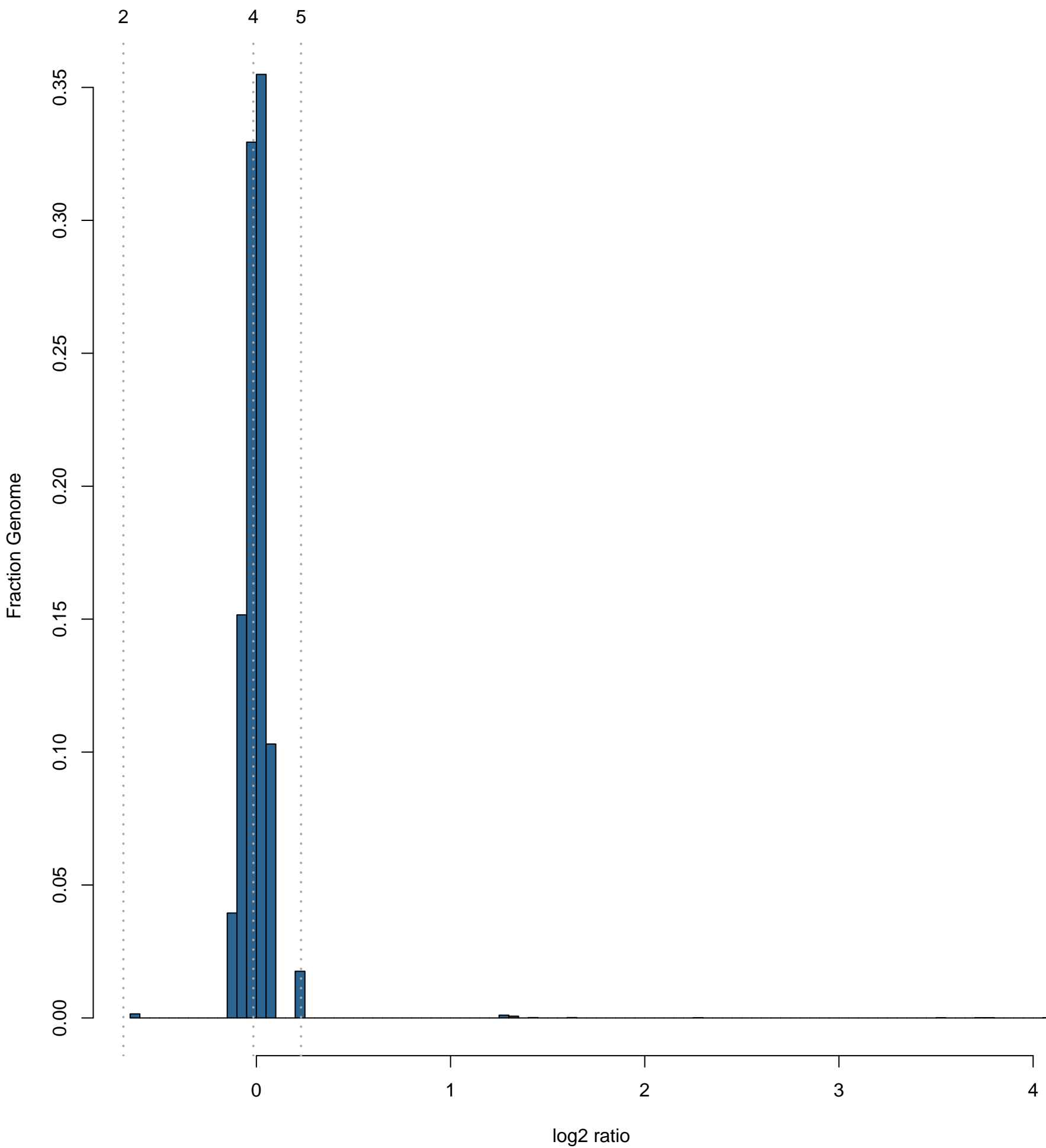


SCNA-fit log-likelihood: -15038.51

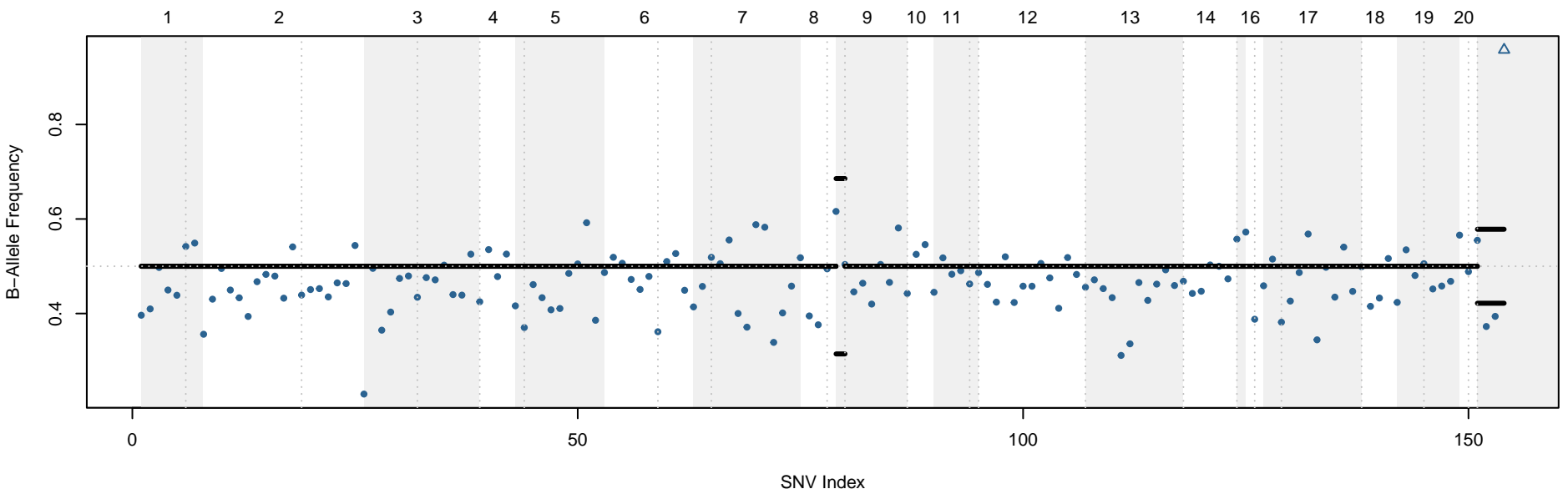




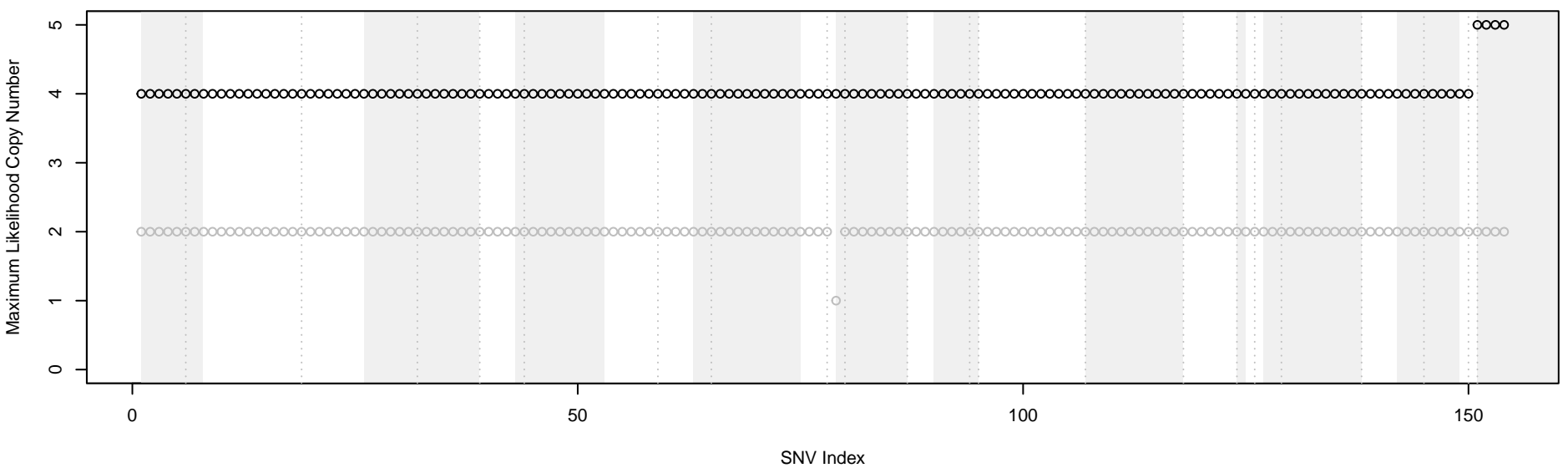
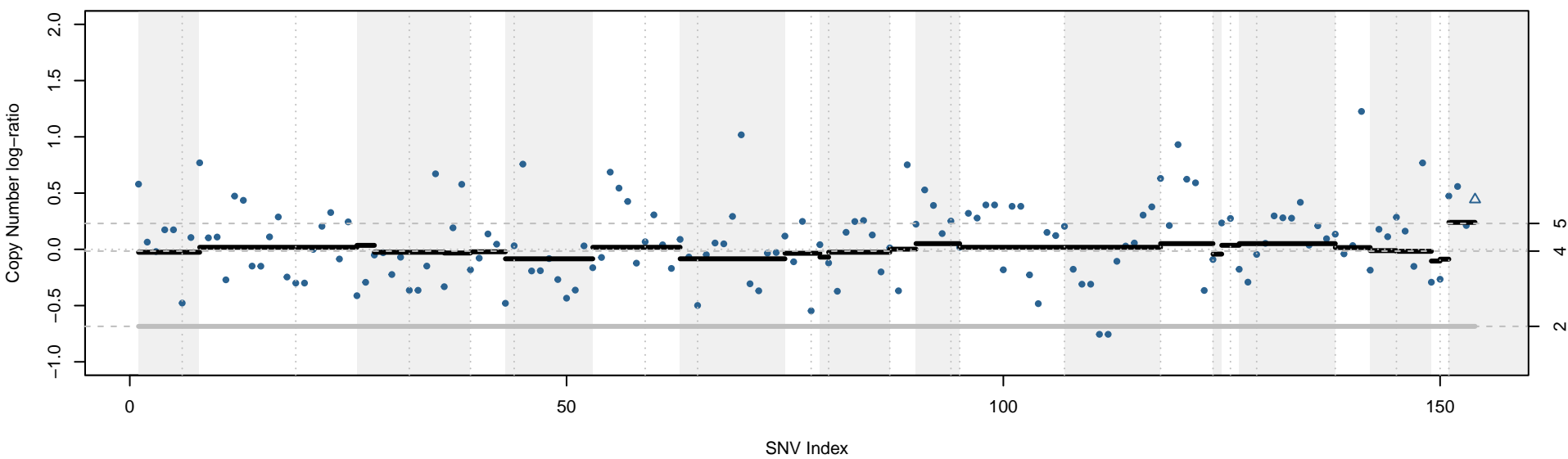
Purity: 0.59 Tumor ploidy: 4.058

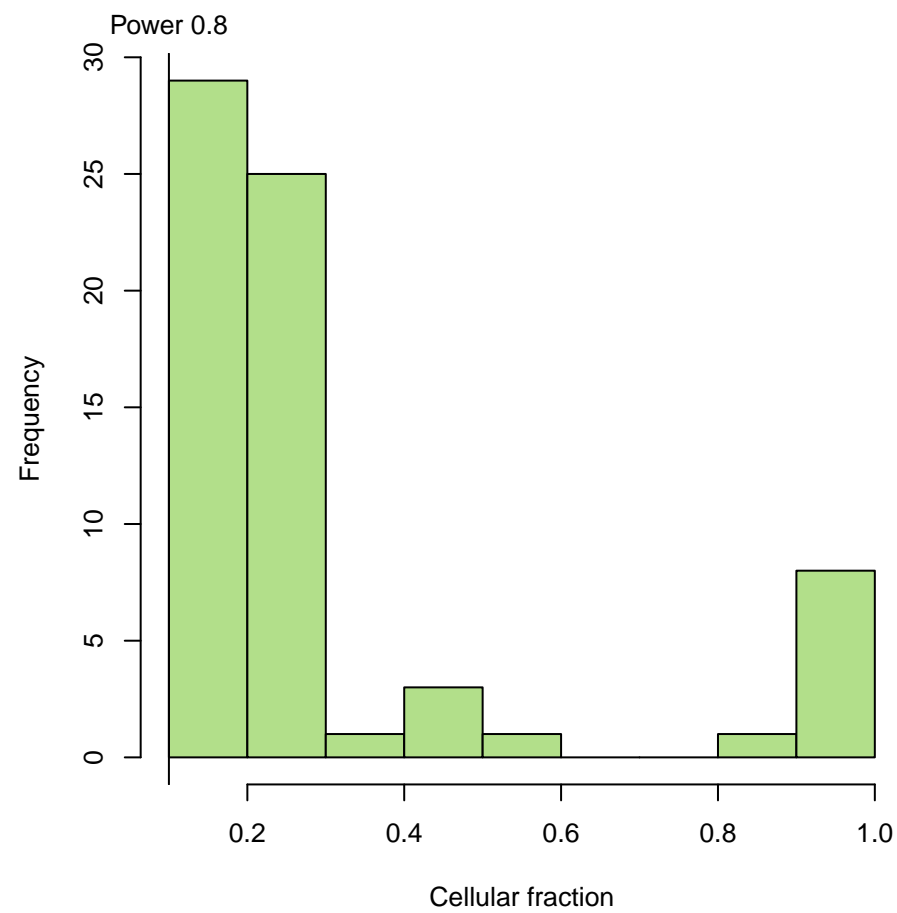
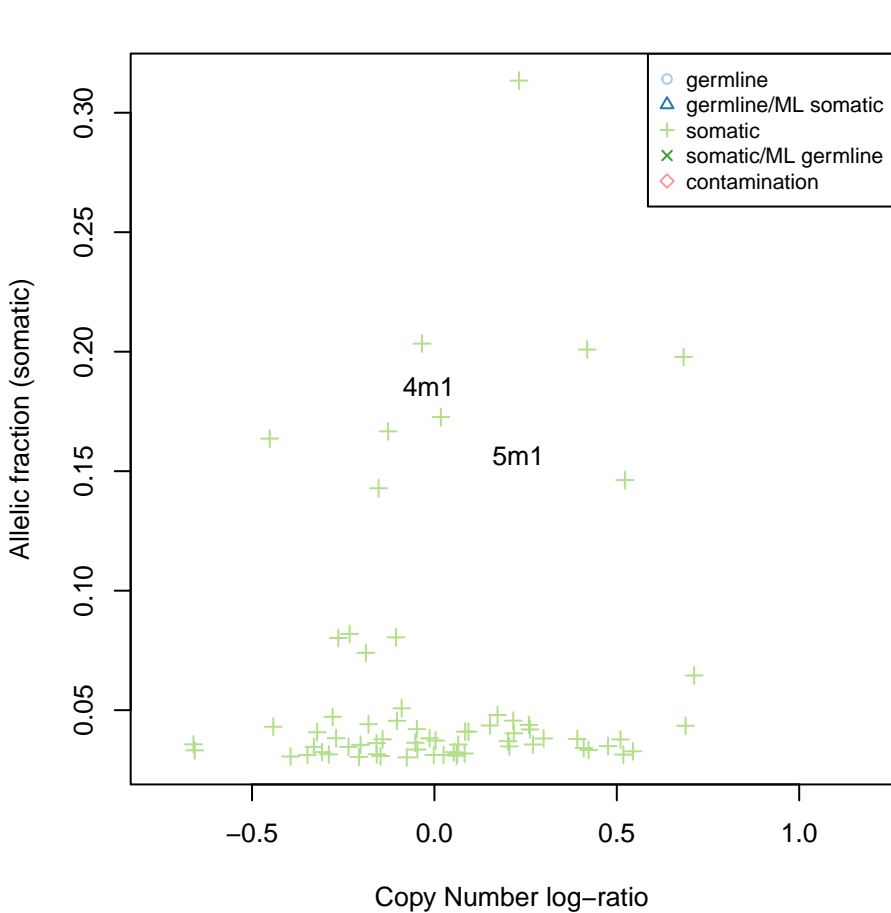
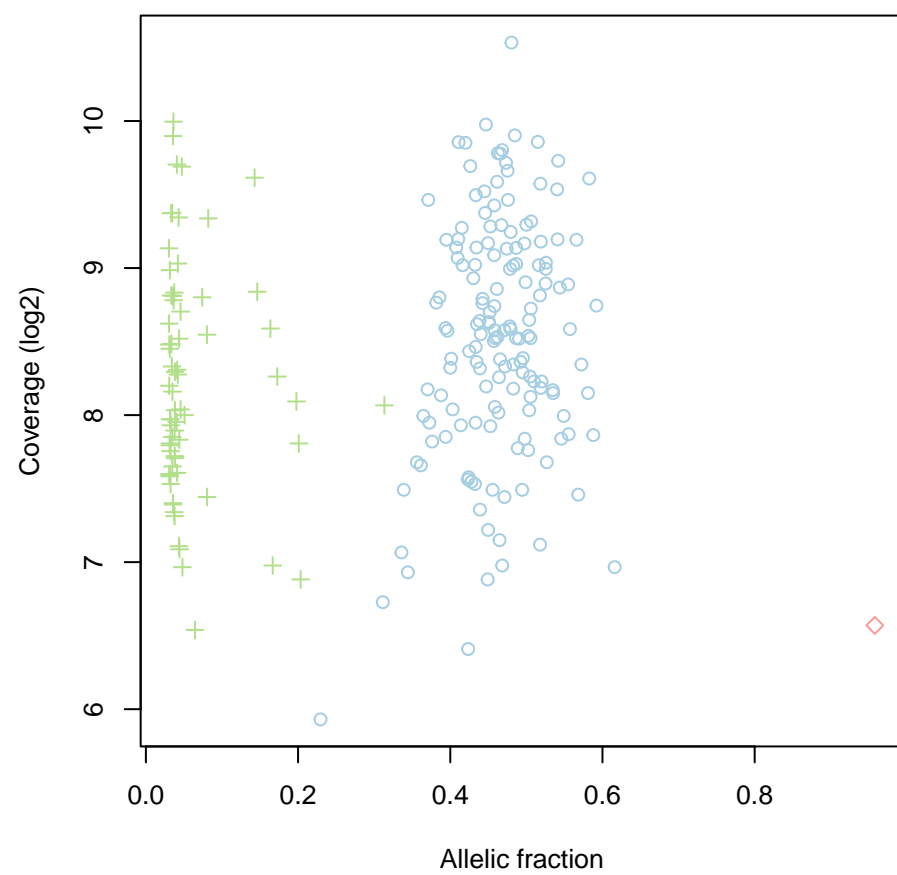
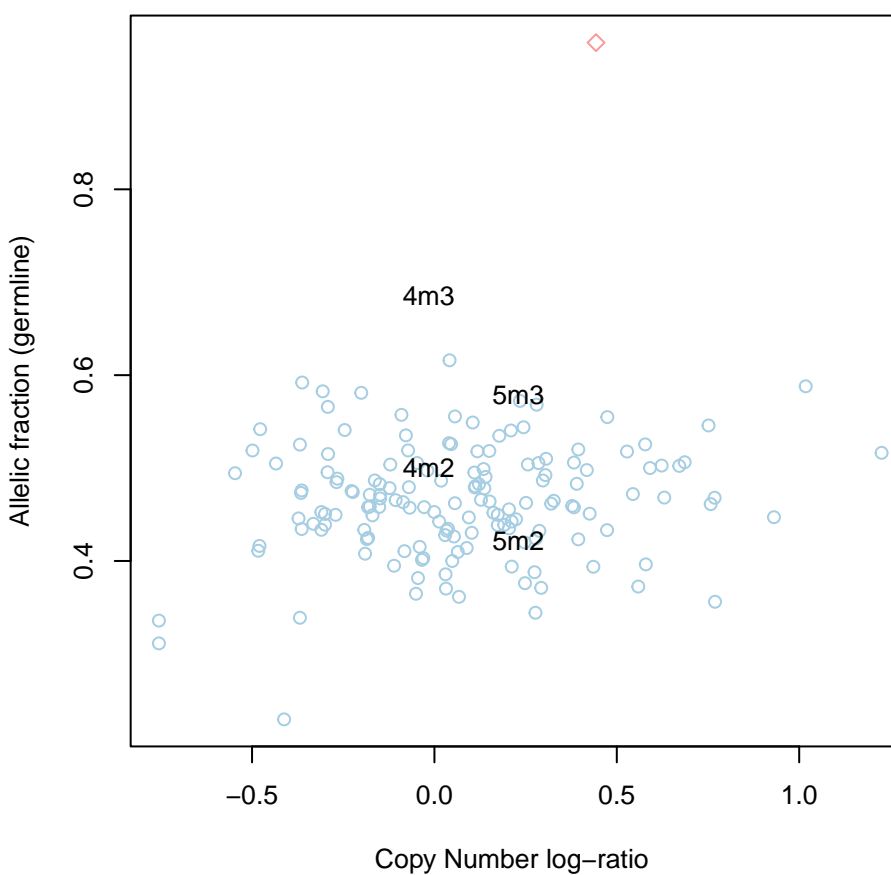


Purity: 0.59 Tumor ploidy: 4.058 SNV log-likelihood: -165.54 GoF: 79.9% Mean coverage: 401;62

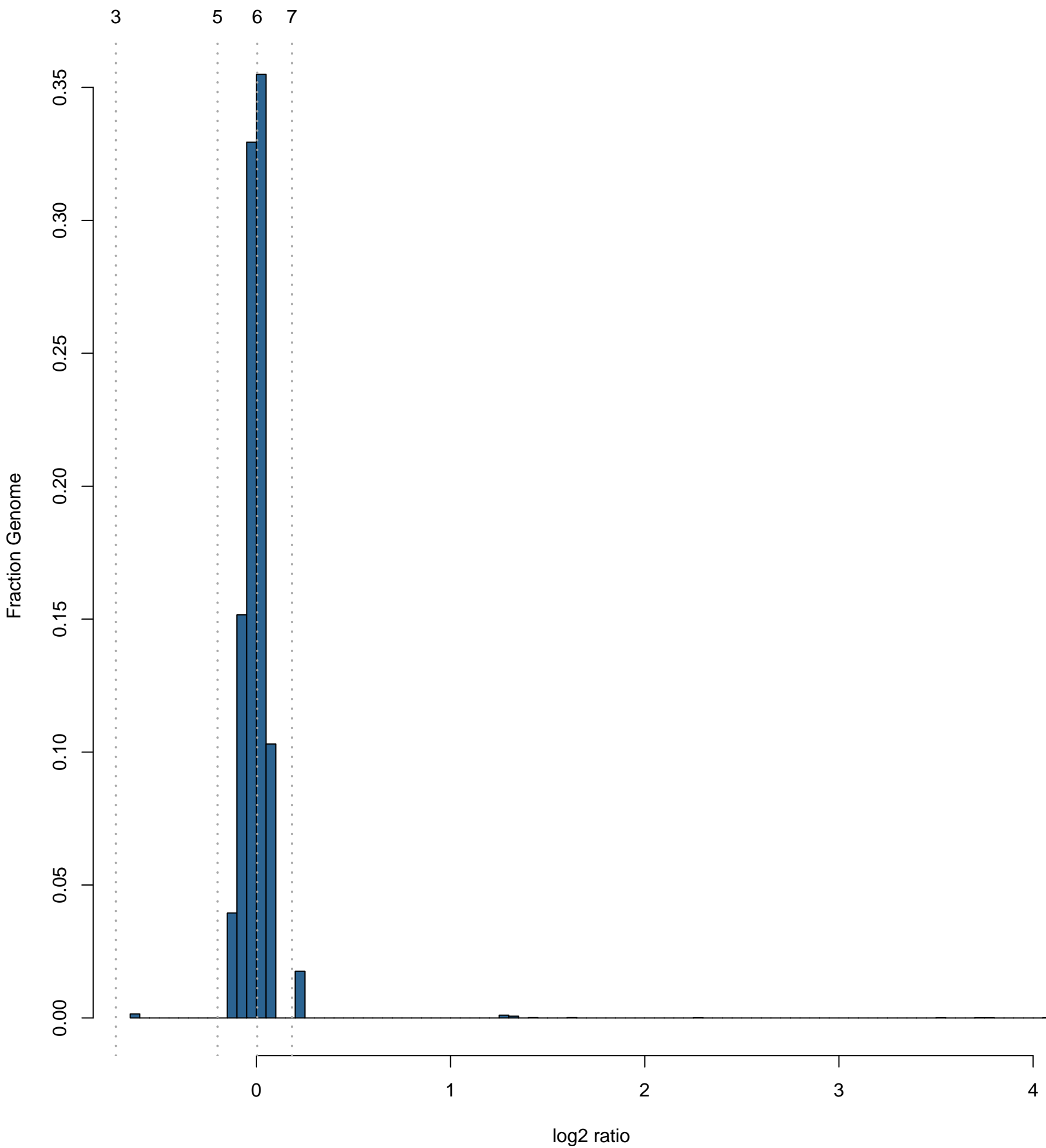


SCNA-fit log-likelihood: -15063.42

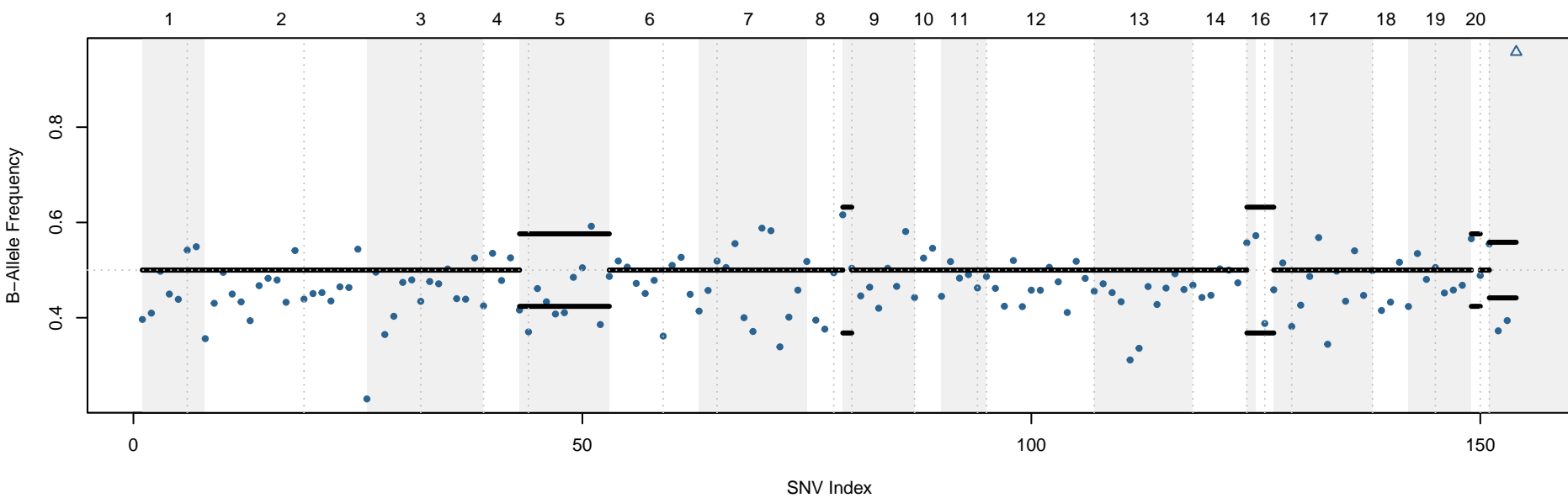




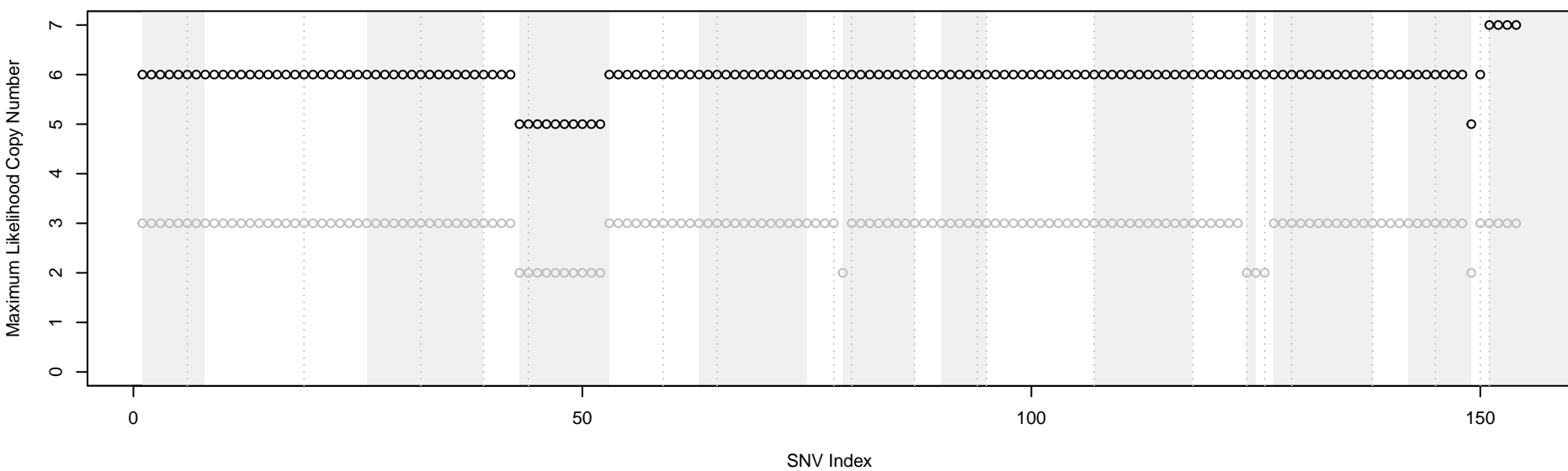
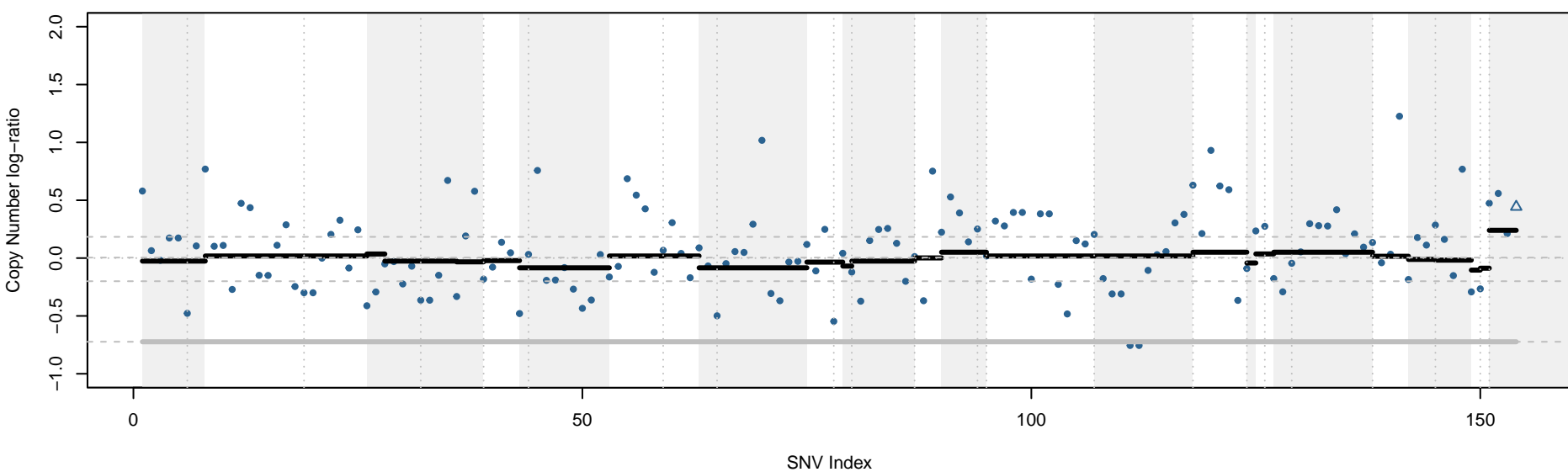
Purity: 0.56 Tumor ploidy: 5.976

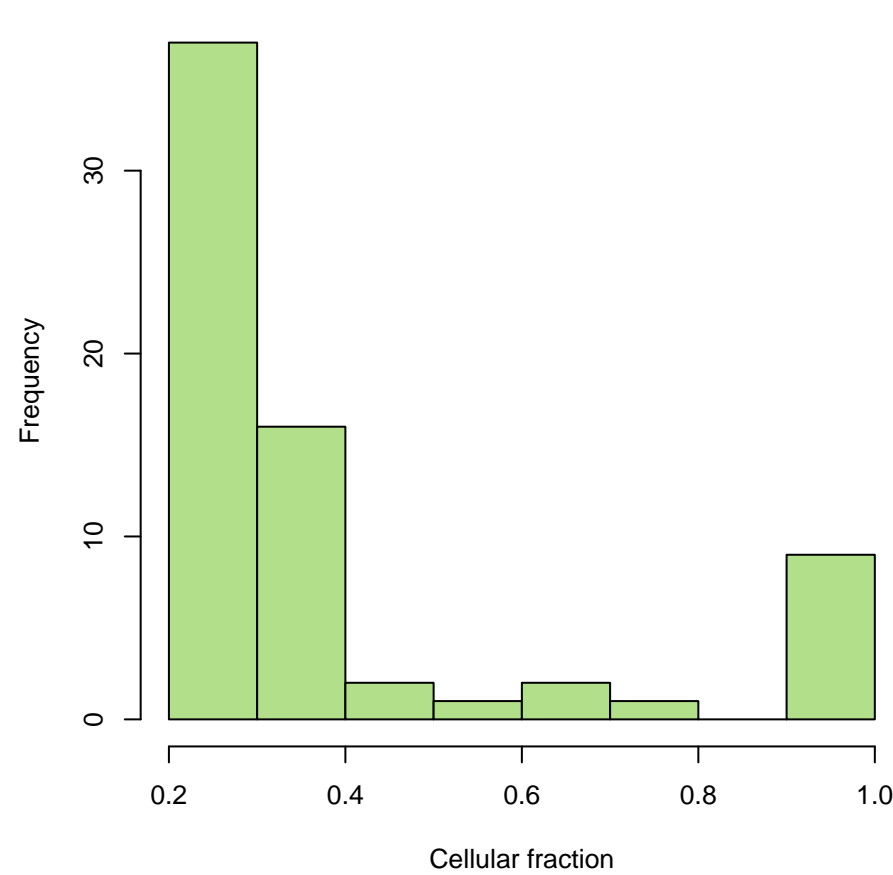
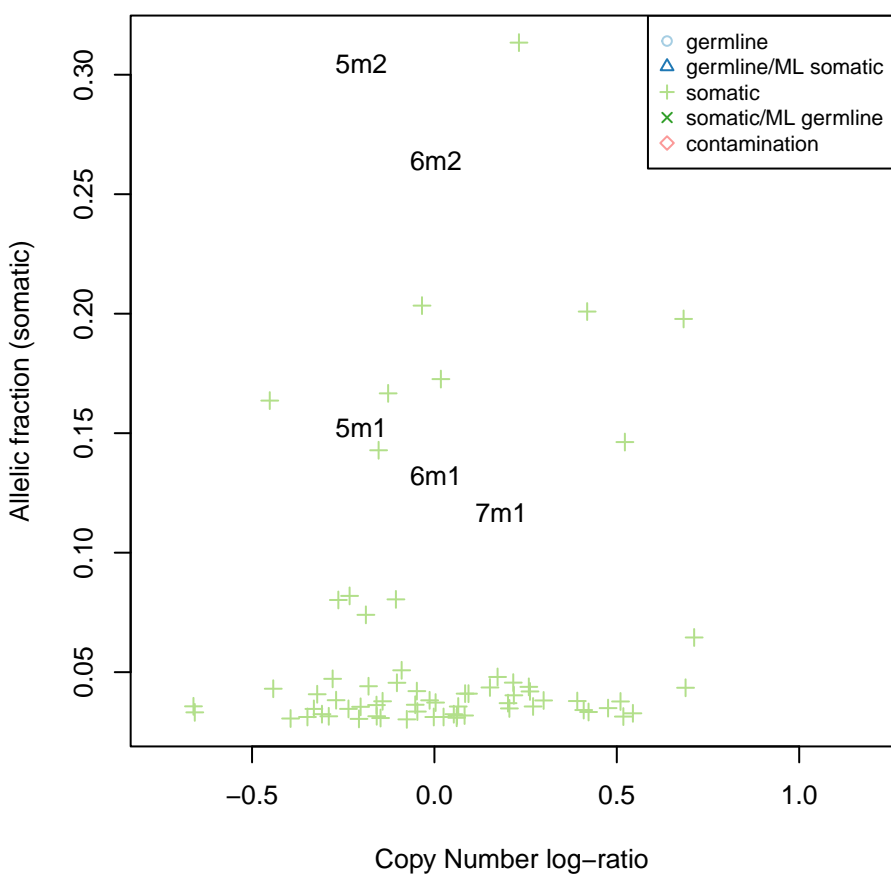
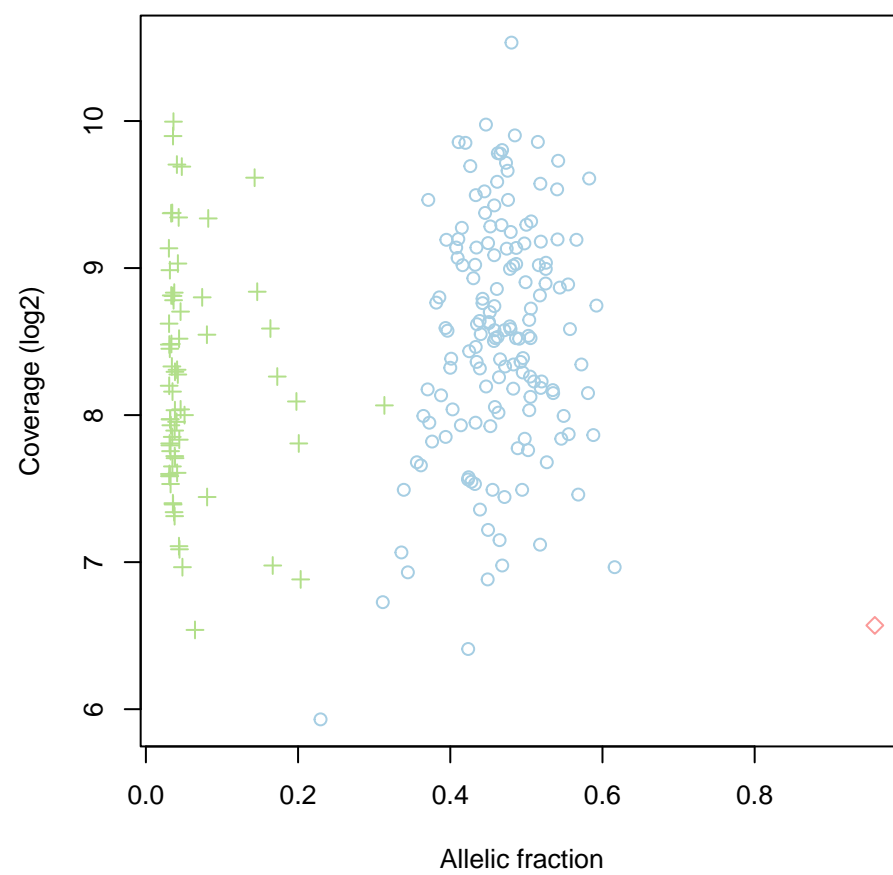
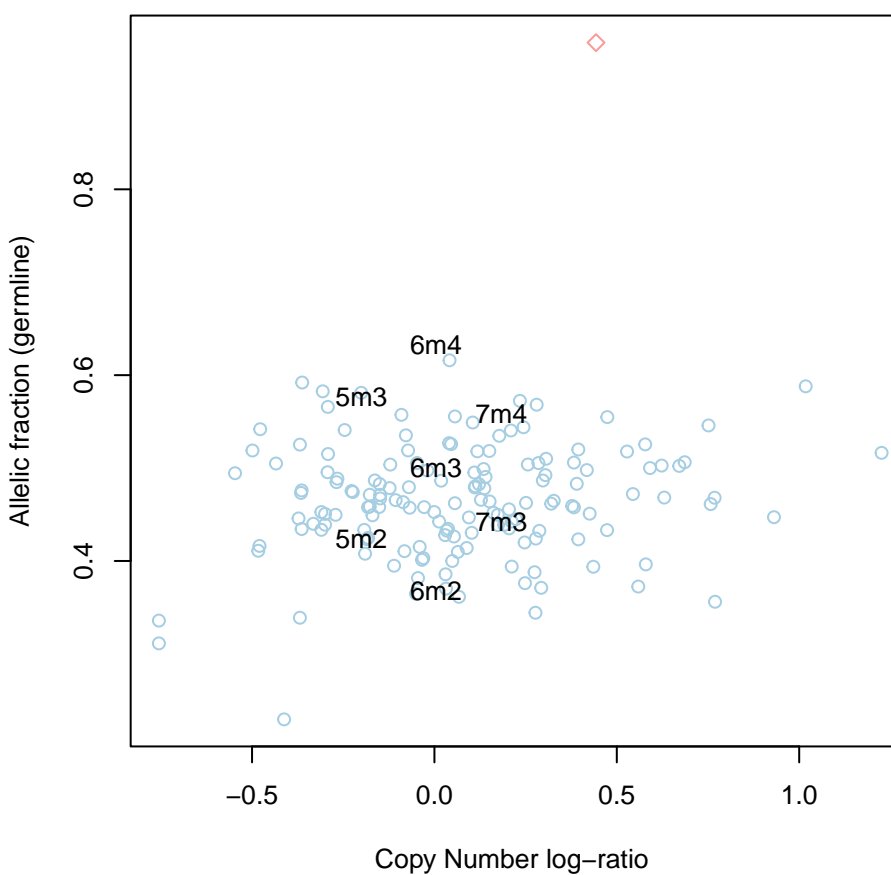


Purity: 0.56 Tumor ploidy: 5.976 SNV log-likelihood: -167.37 GoF: 88.3% Mean coverage: 401;62

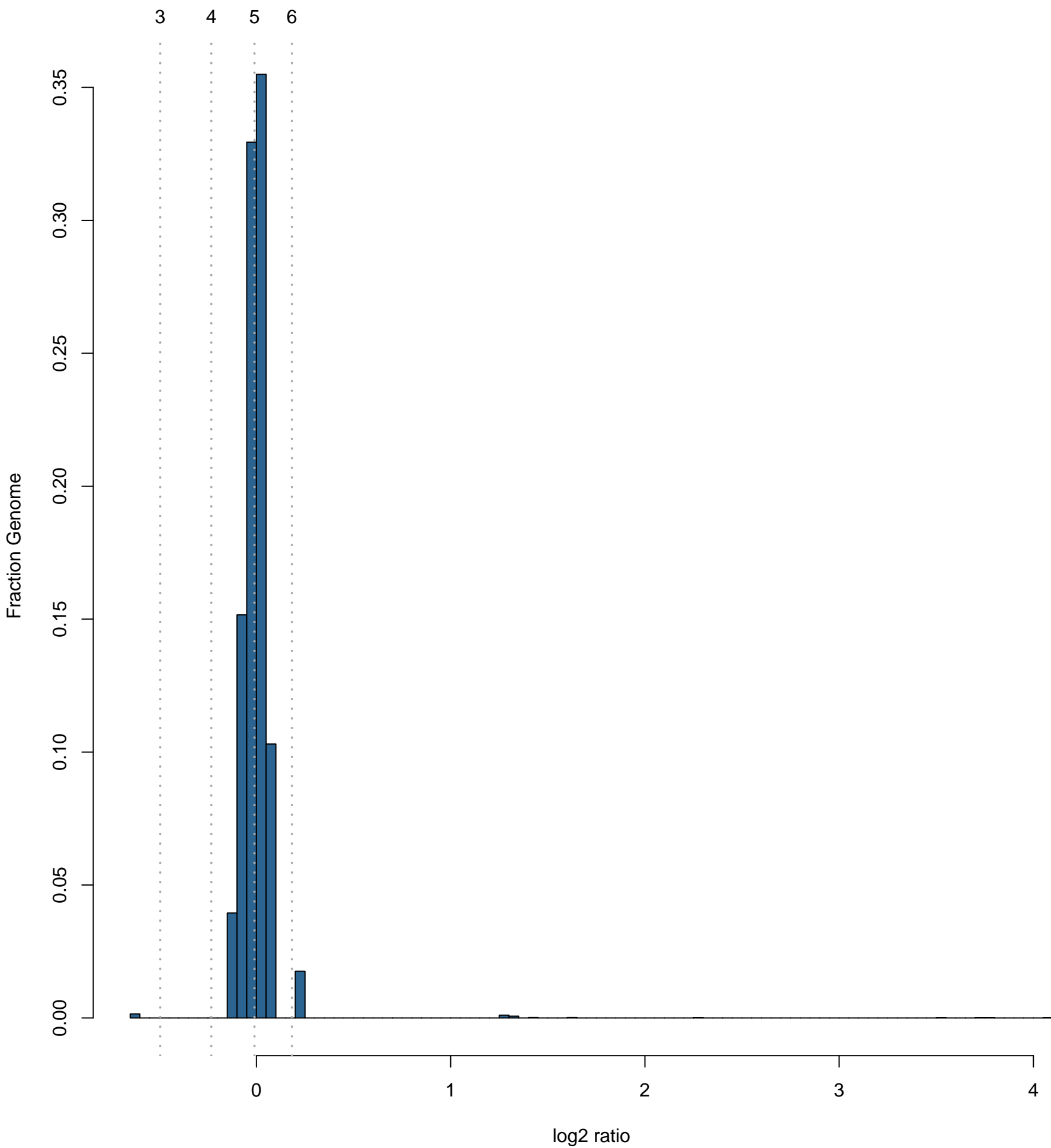


SCNA-fit log-likelihood: -15051.83

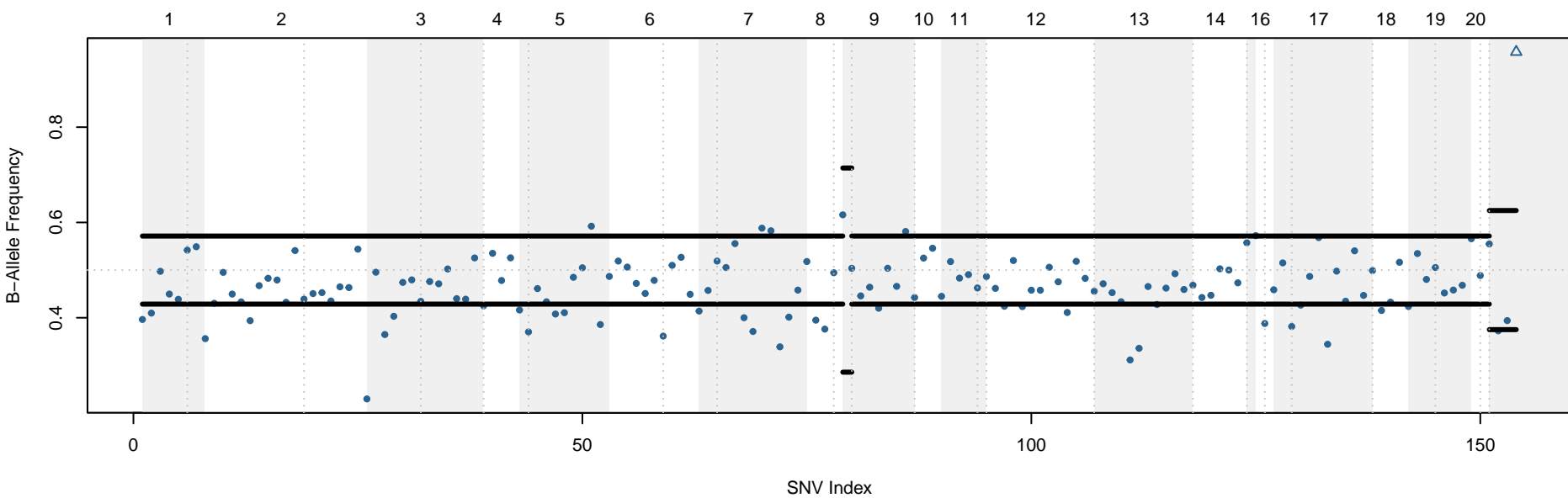




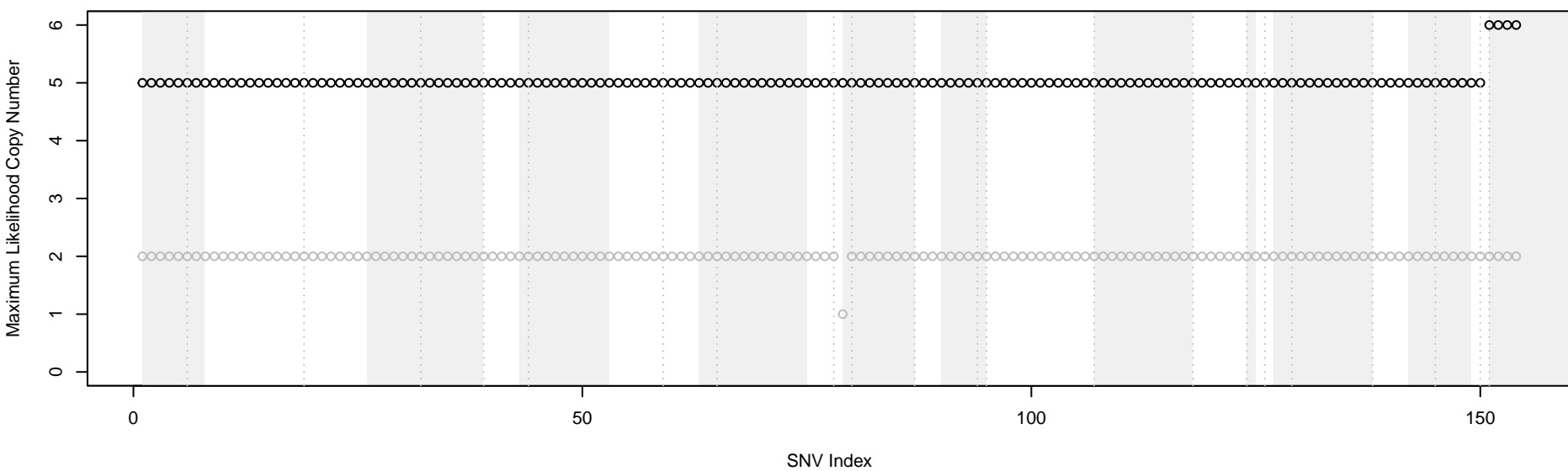
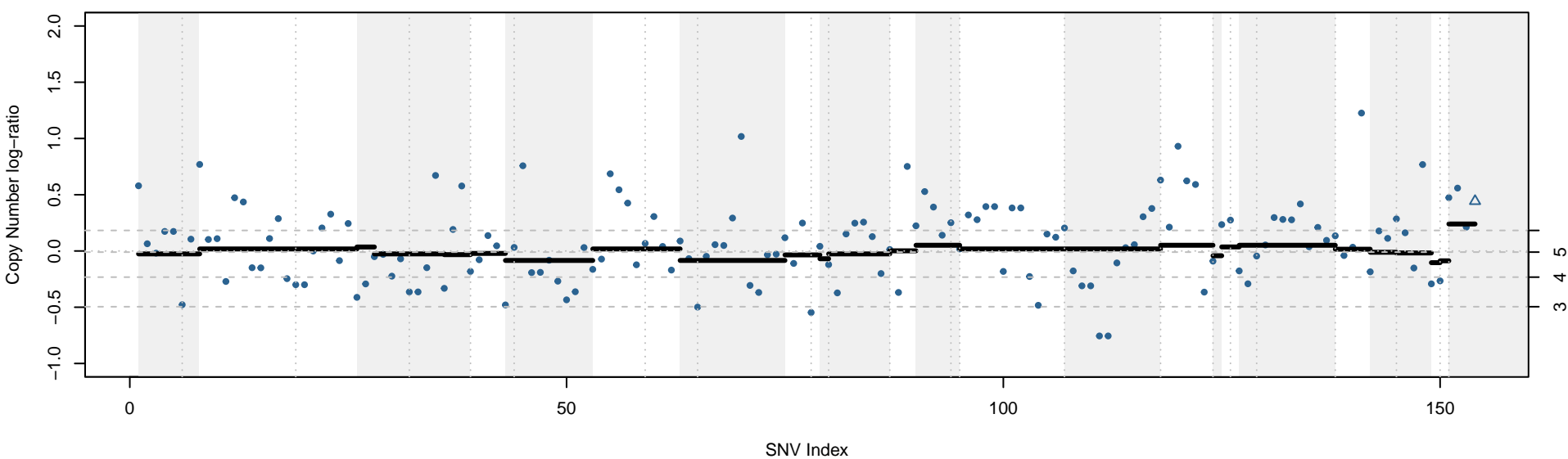
Purity: 0.5 Tumor ploidy: 5.049

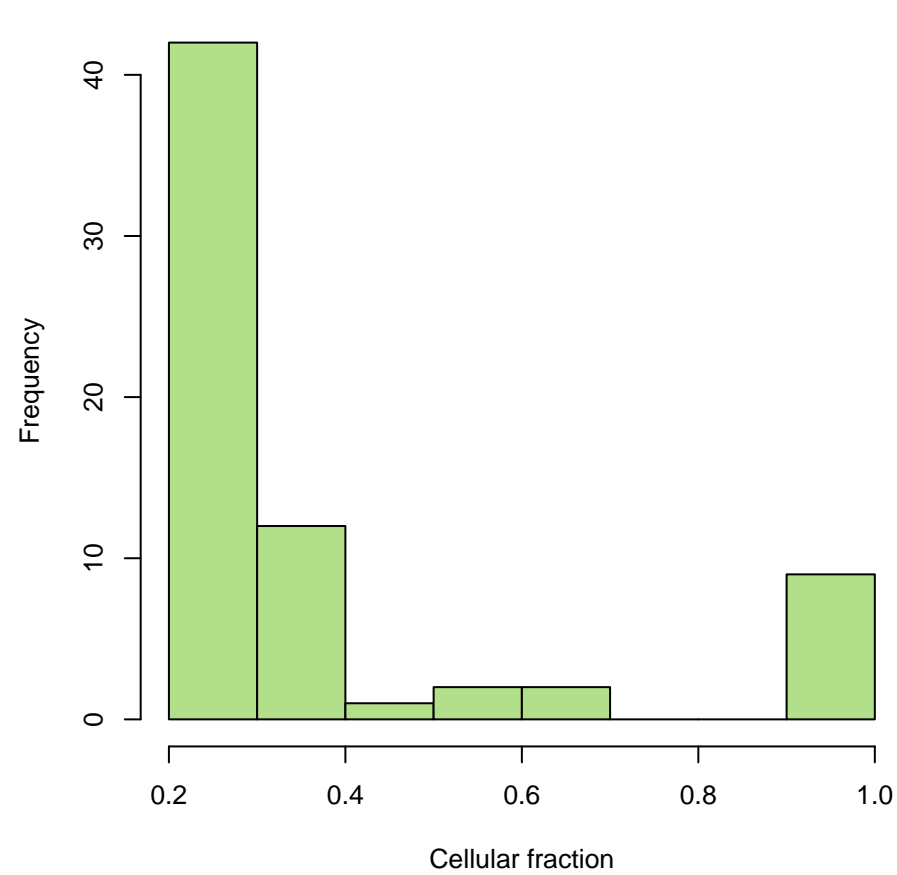
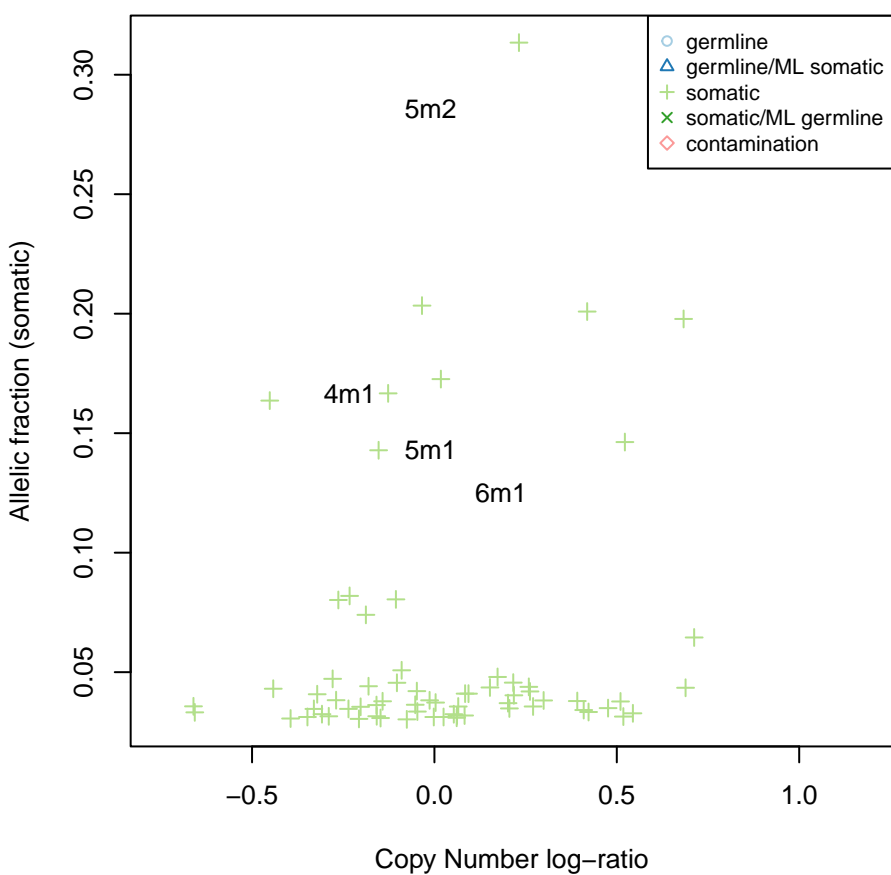
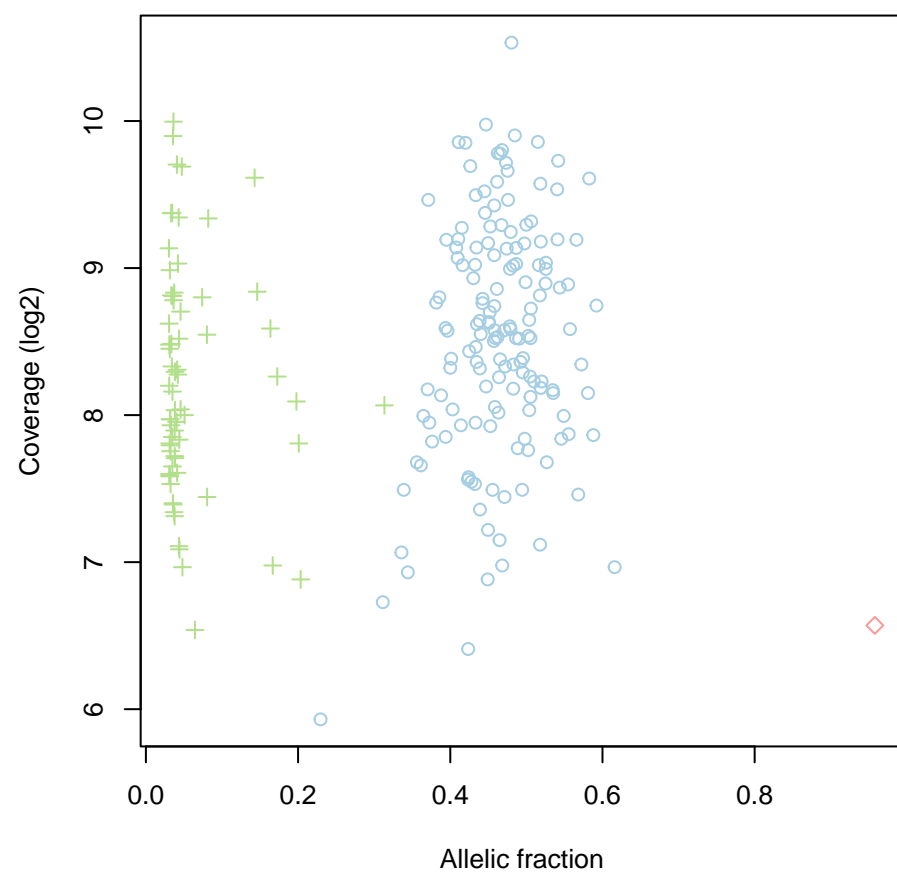
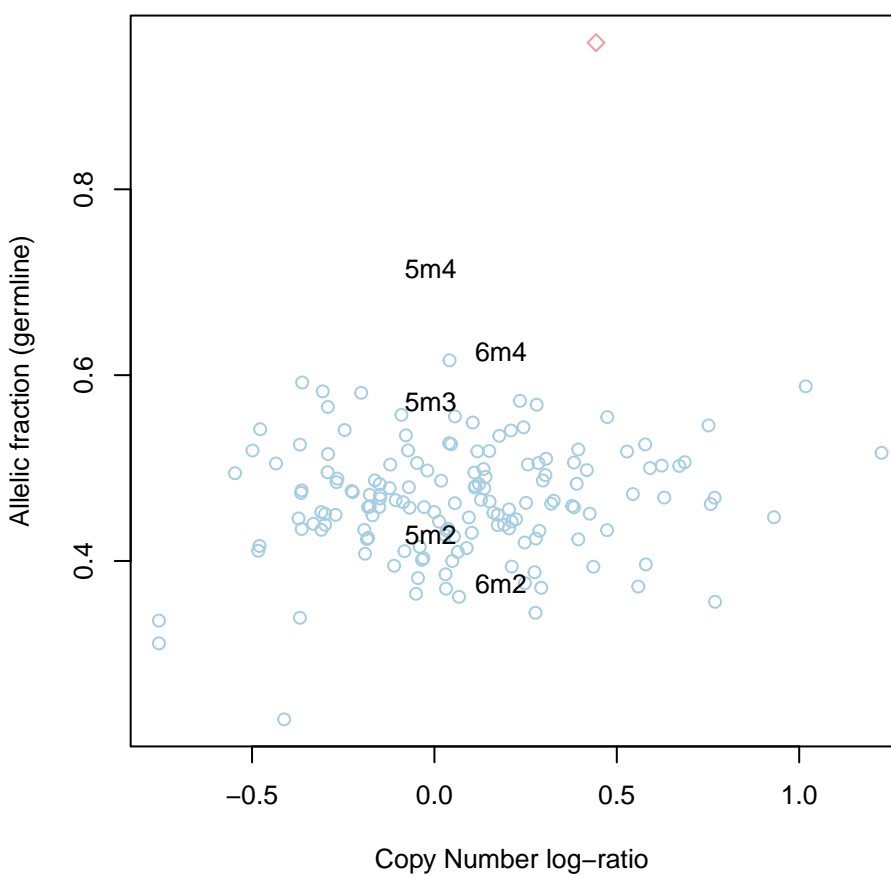


Purity: 0.5 Tumor ploidy: 5.049 SNV log-likelihood: -189.05 GoF: 89.2% Mean coverage: 401;62

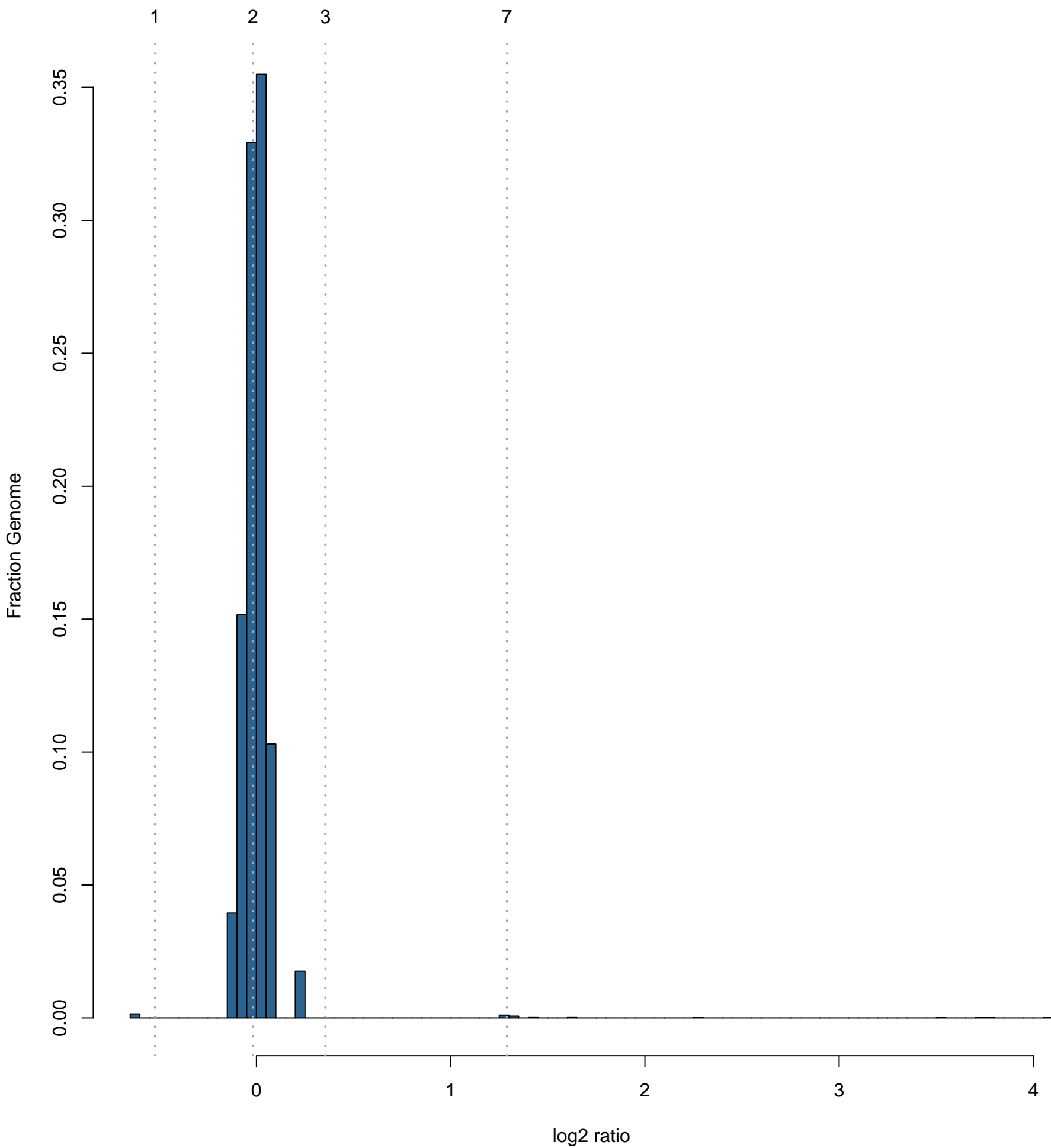


SCNA-fit log-likelihood: -15052.26

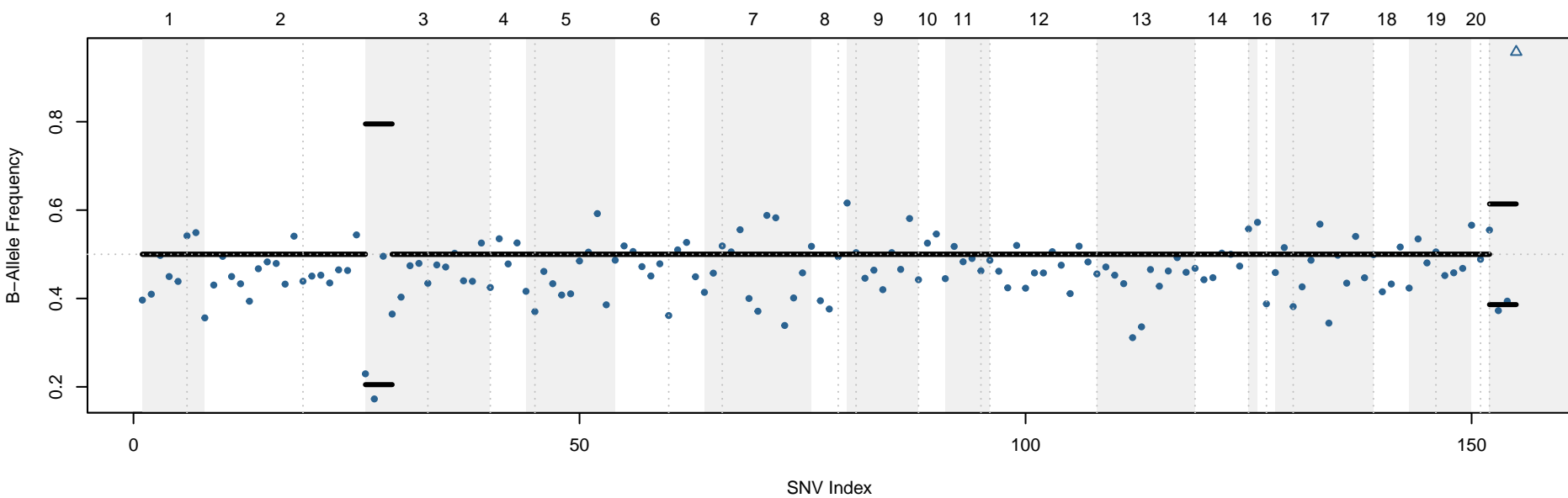




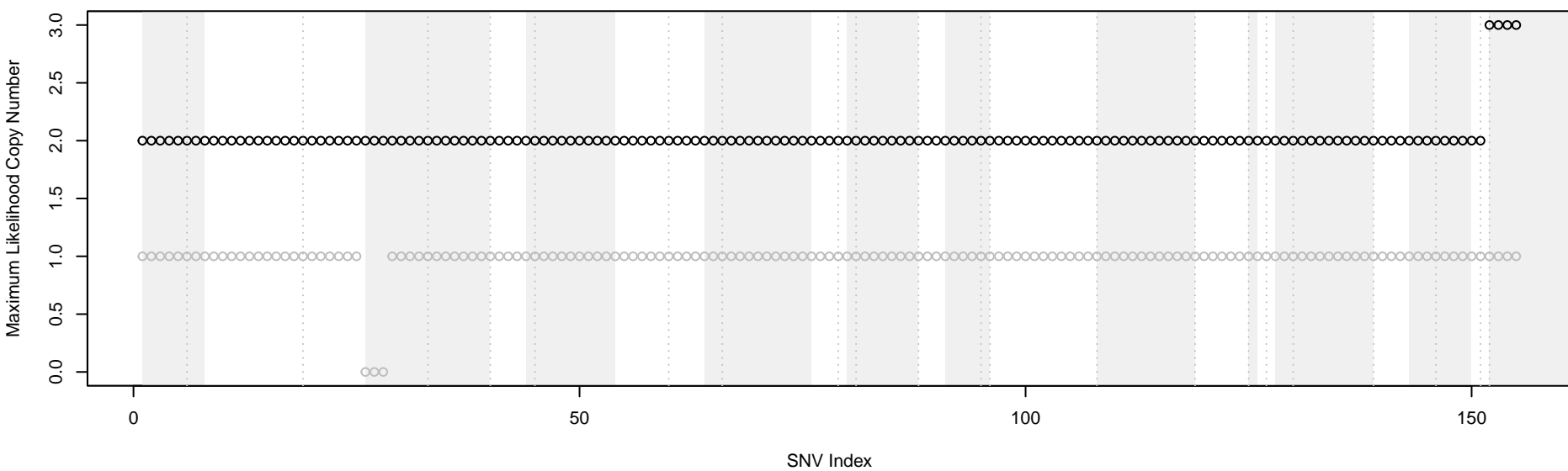
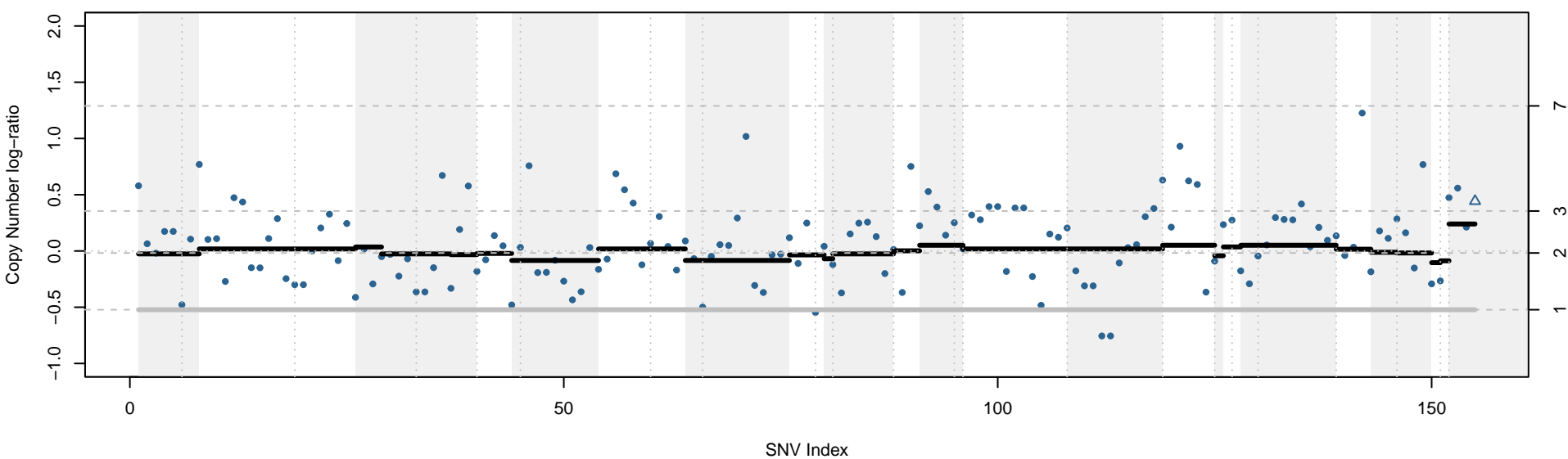
Purity: 0.59 Tumor ploidy: 2.042

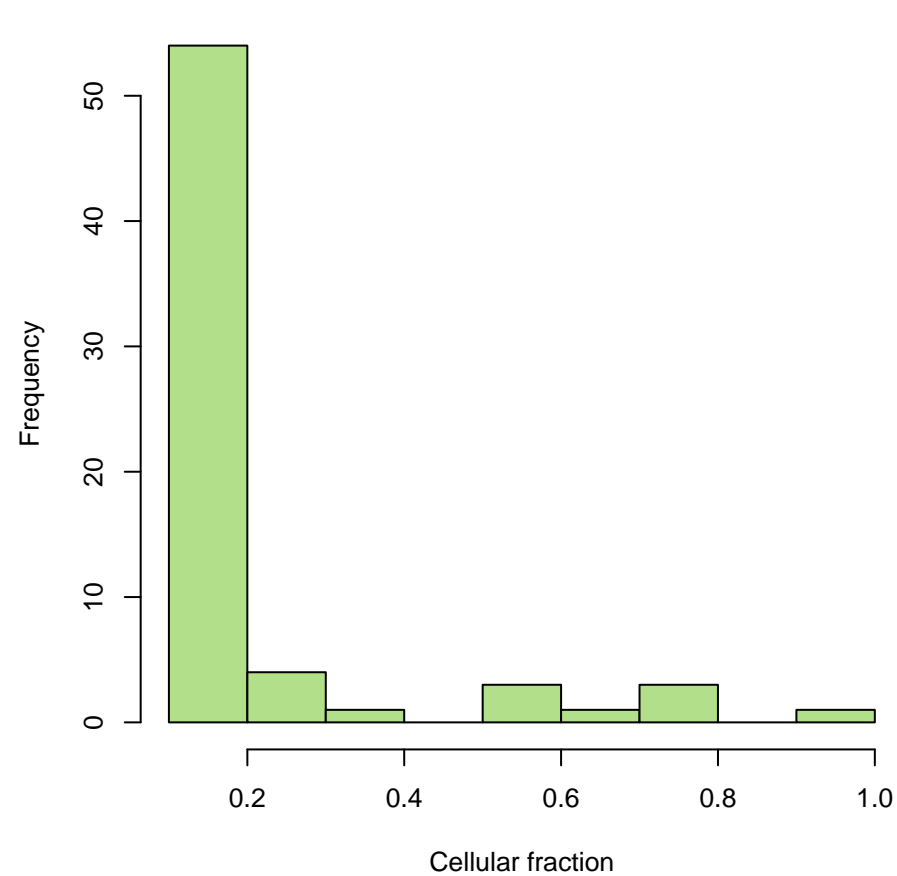
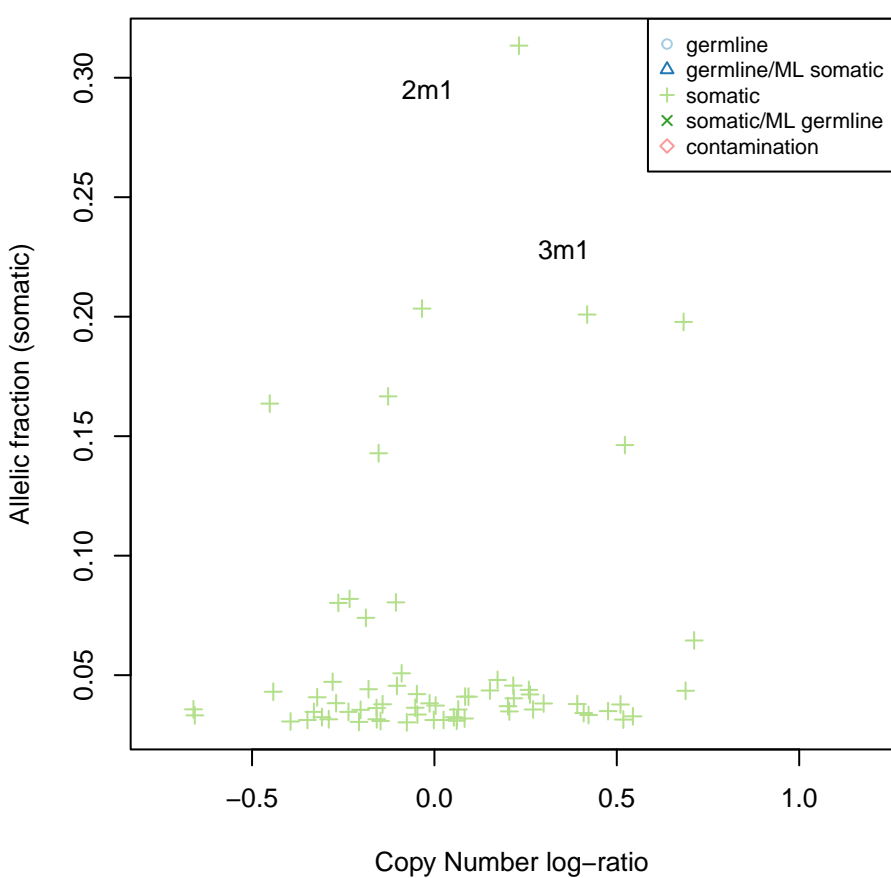
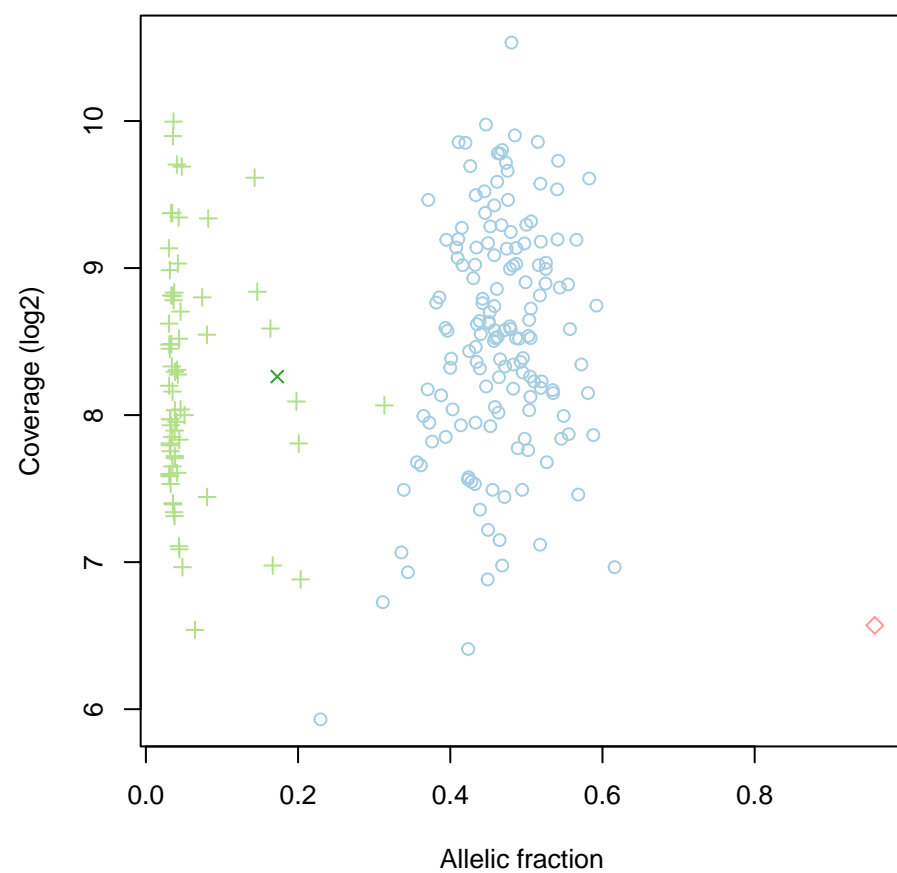
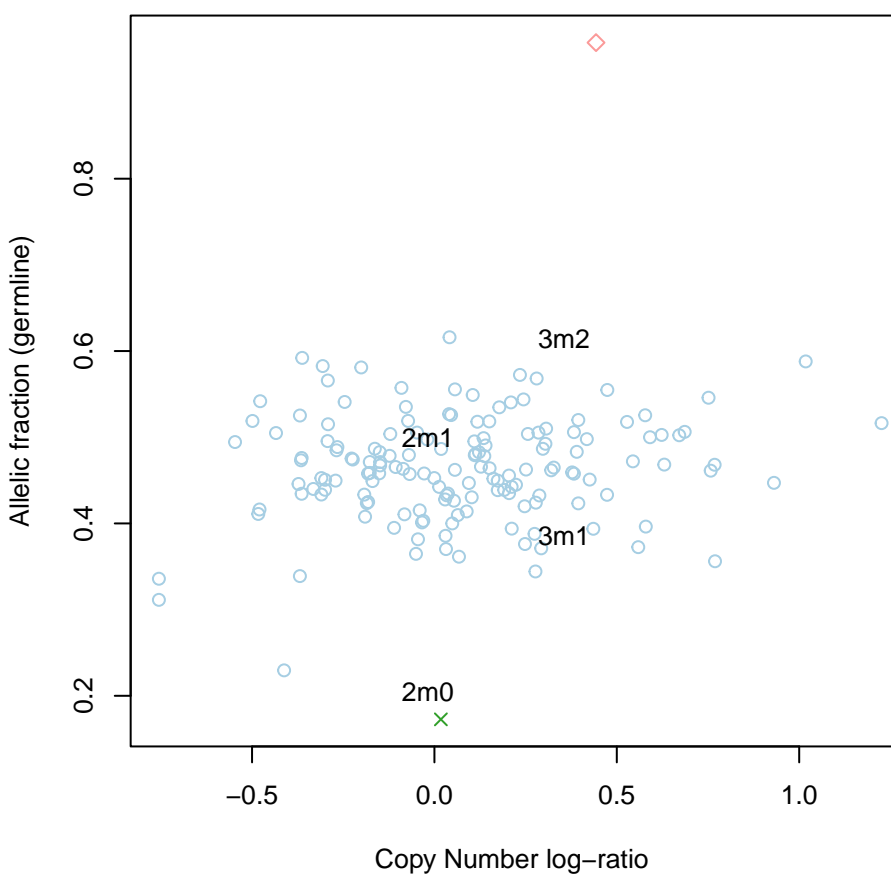


Purity: 0.59 Tumor ploidy: 2.042 SNV log-likelihood: -307.26 GoF: 51.3% Mean coverage: 401;62

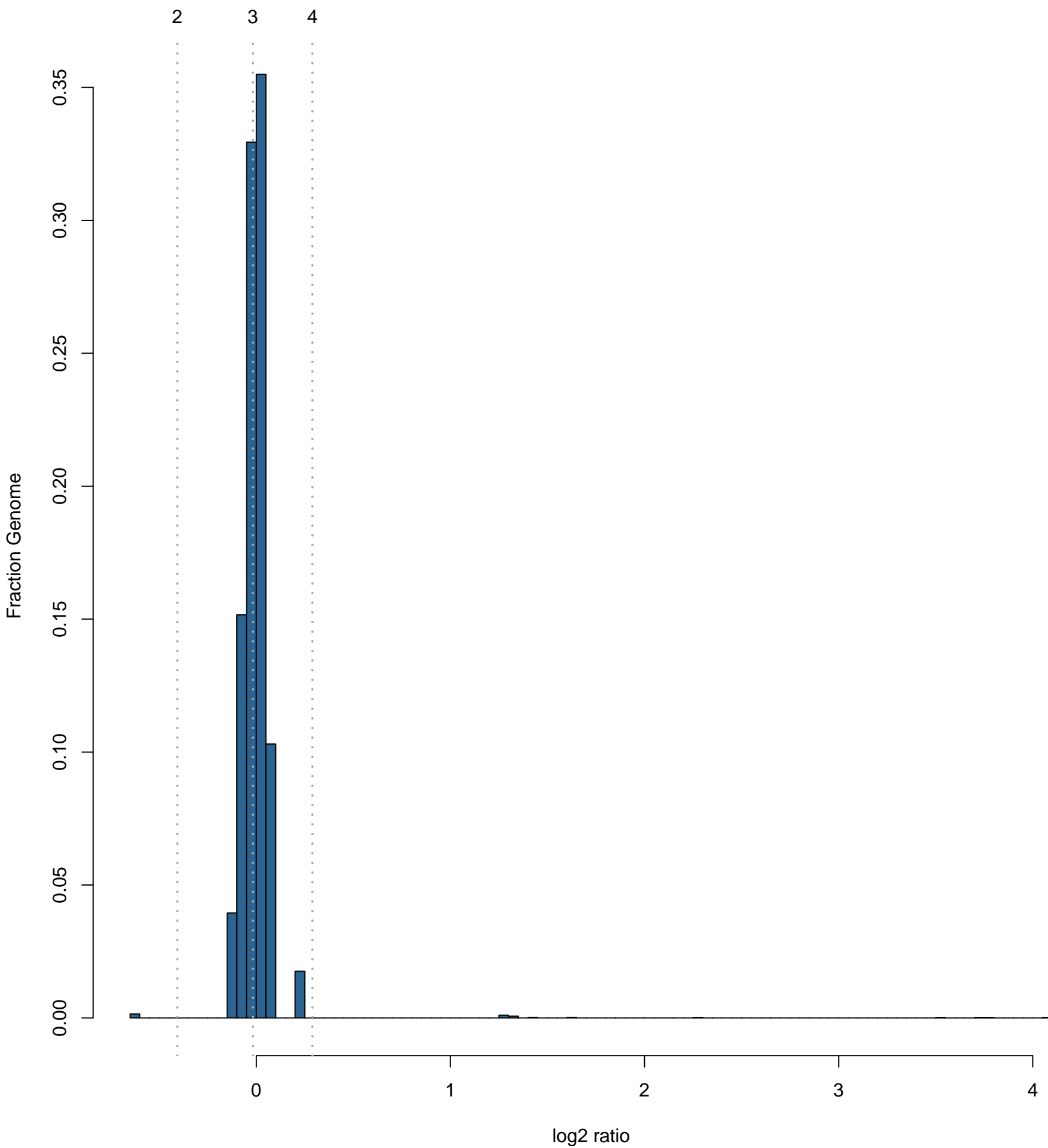


SCNA-fit log-likelihood: -15047.58

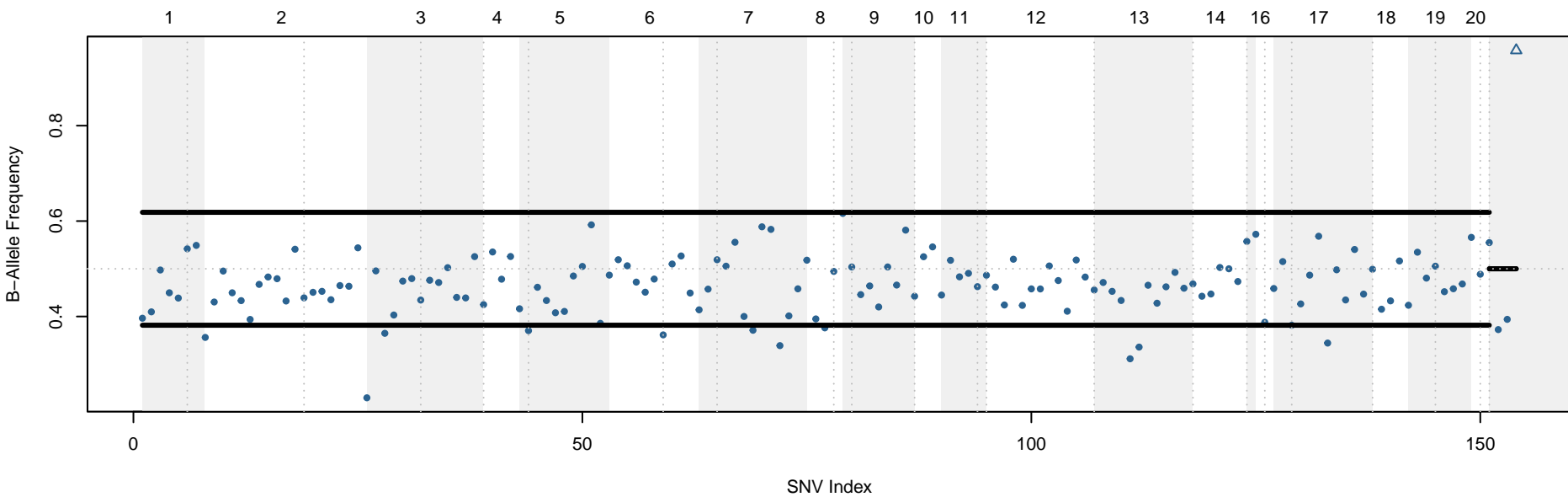




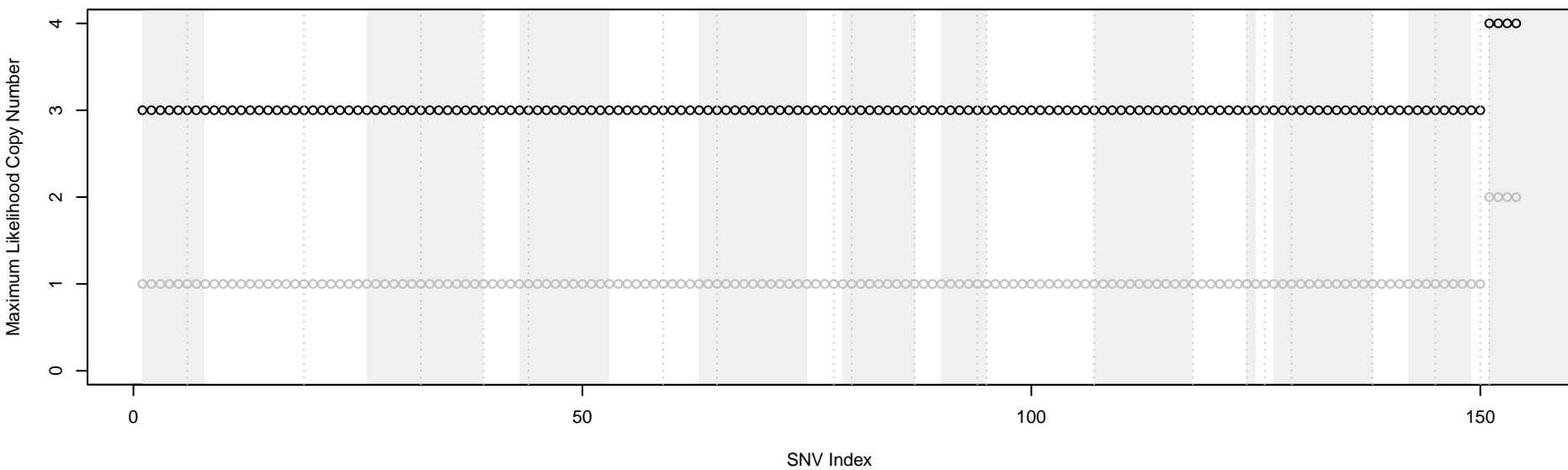
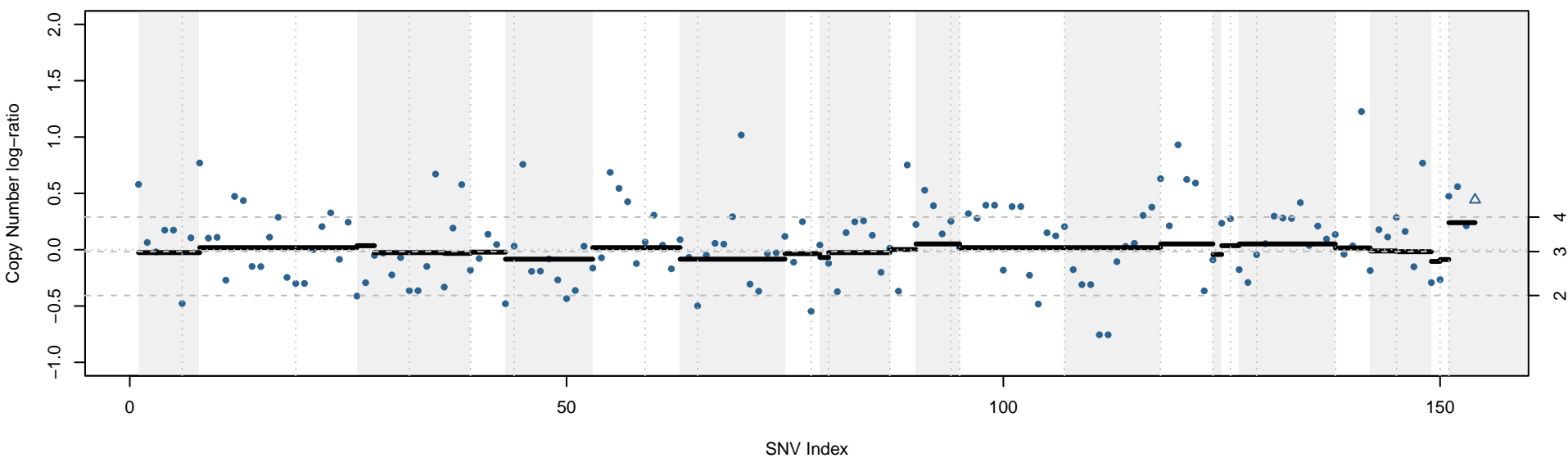
Purity: 0.62 Tumor ploidy: 3.05

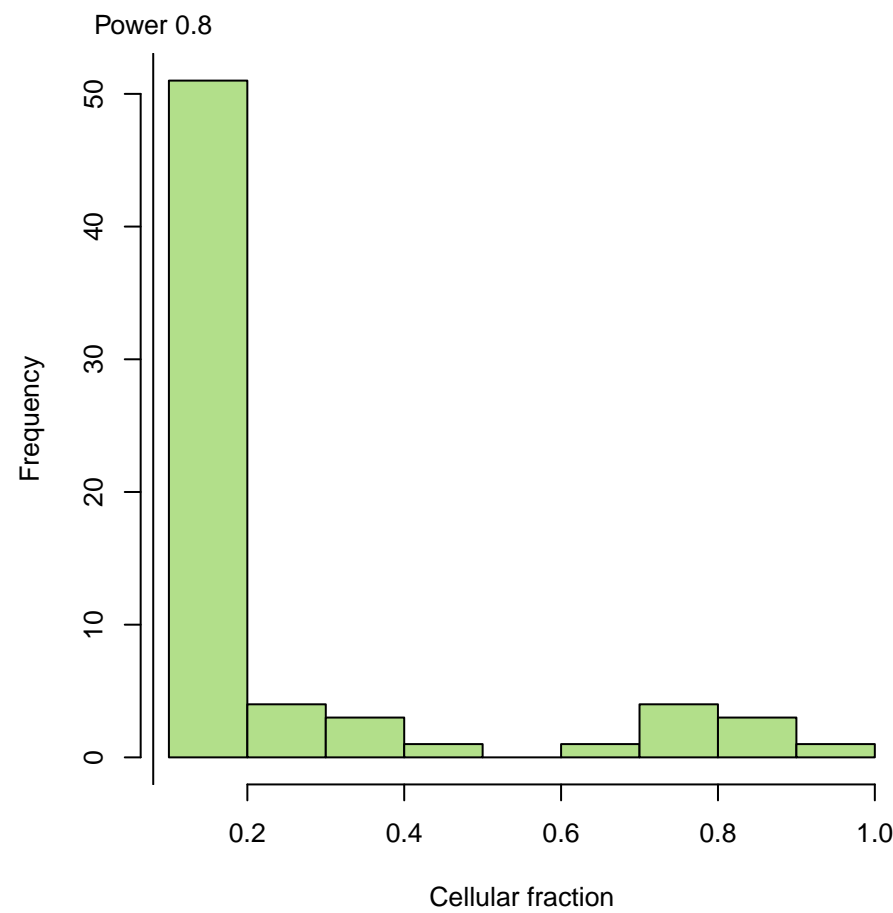
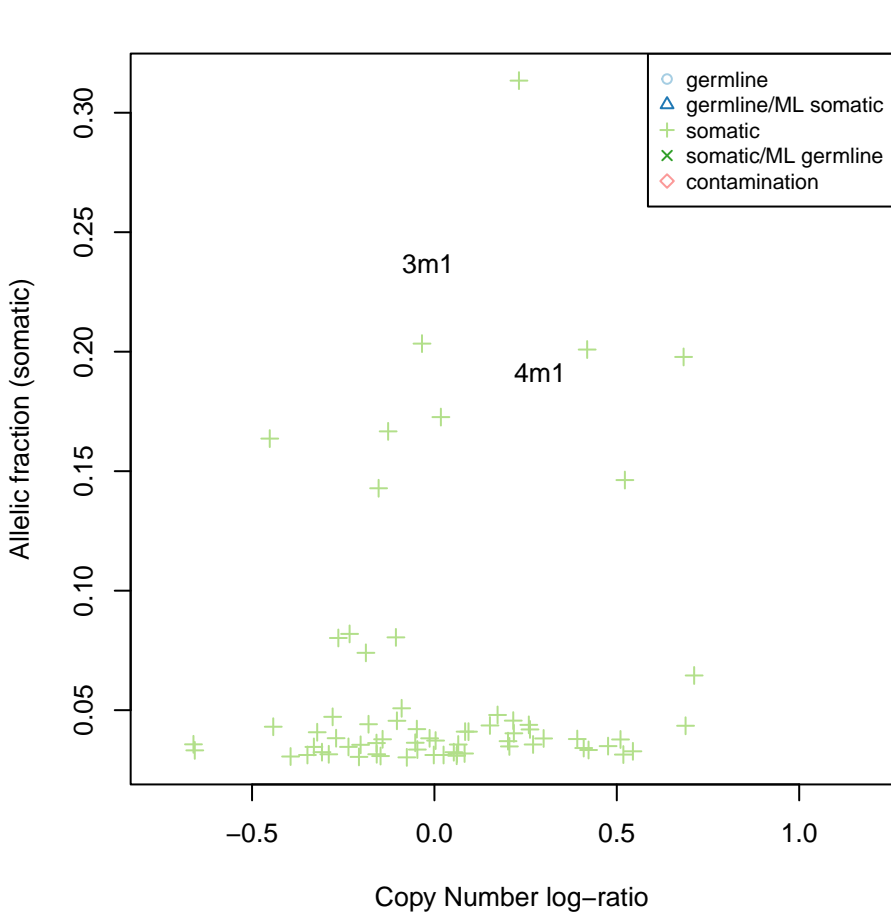
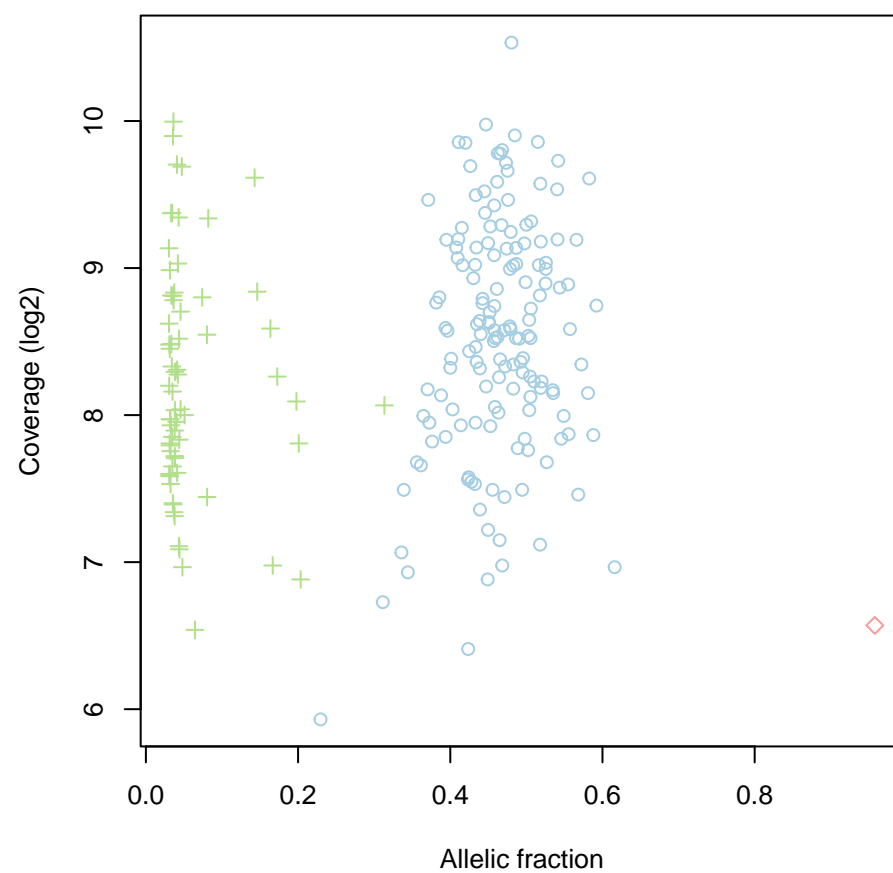
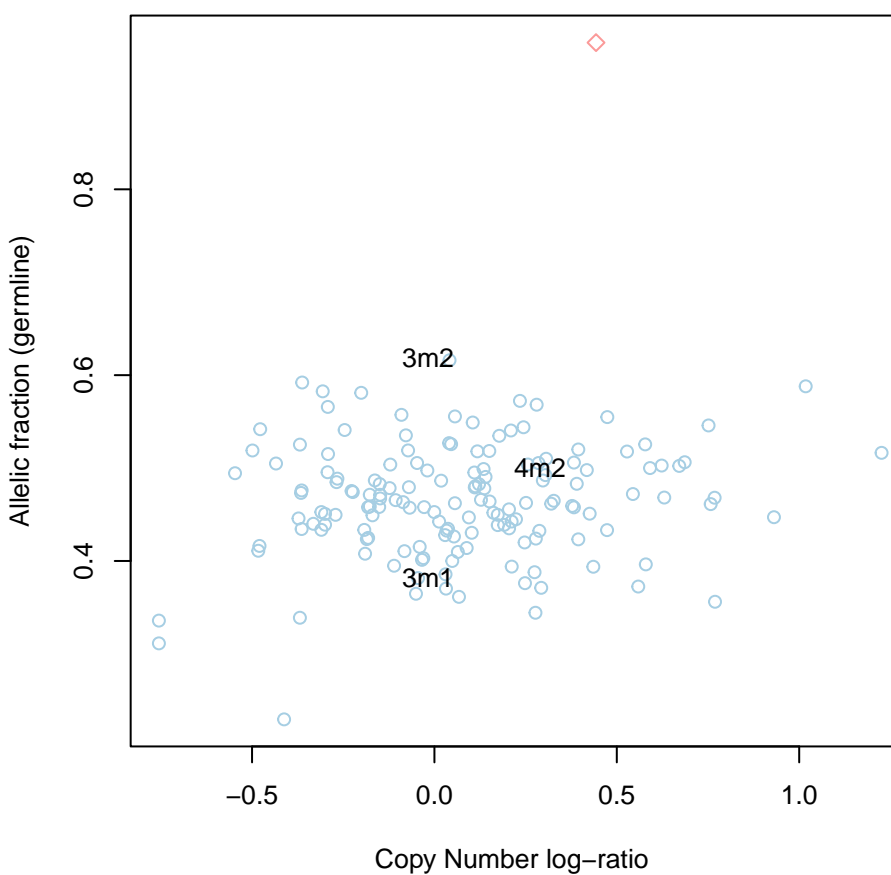


Purity: 0.62 Tumor ploidy: 3.05 SNV log-likelihood: -635.4 GoF: 63.4% Mean coverage: 401;62

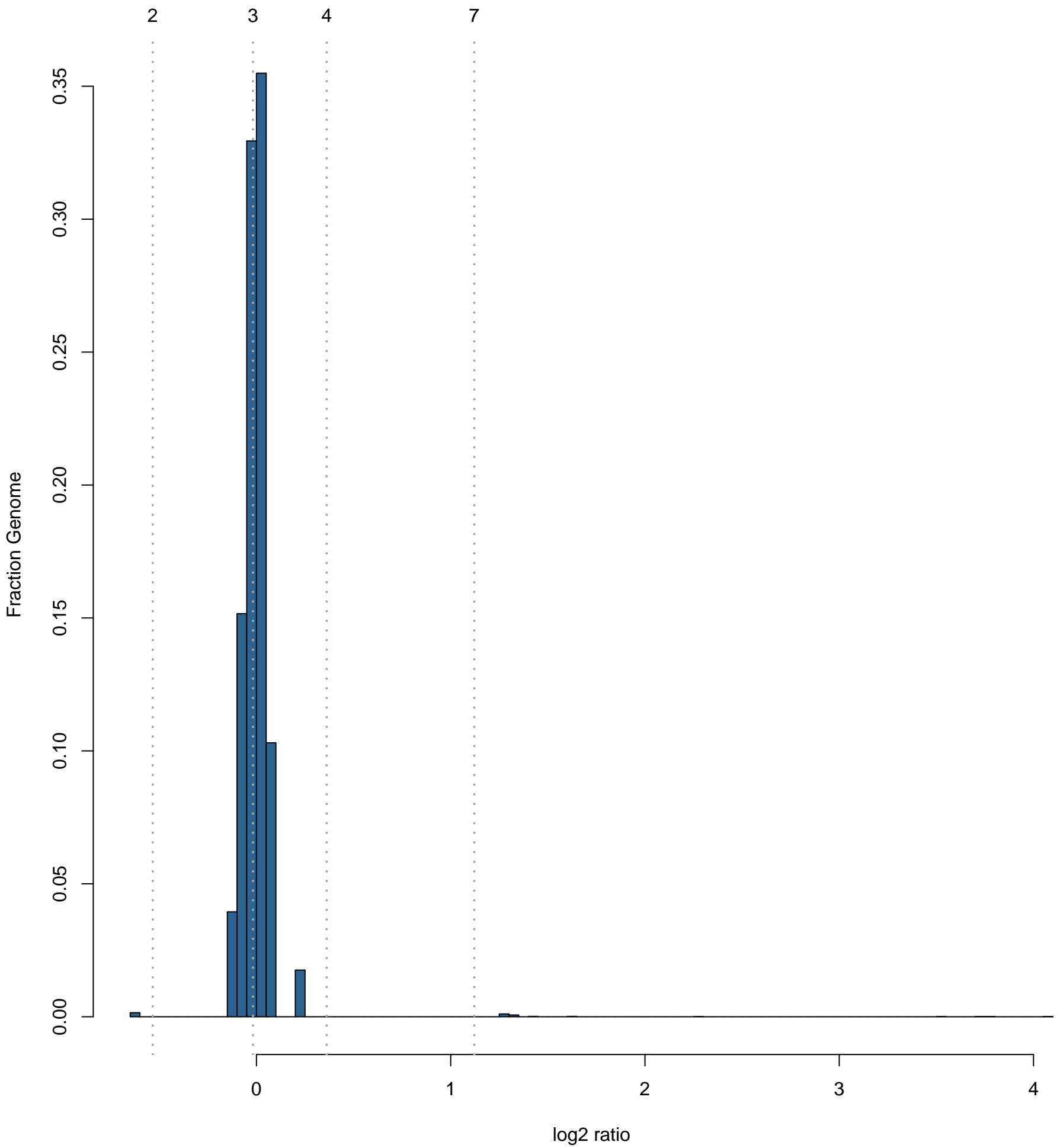


SCNA-fit log-likelihood: -15068.55

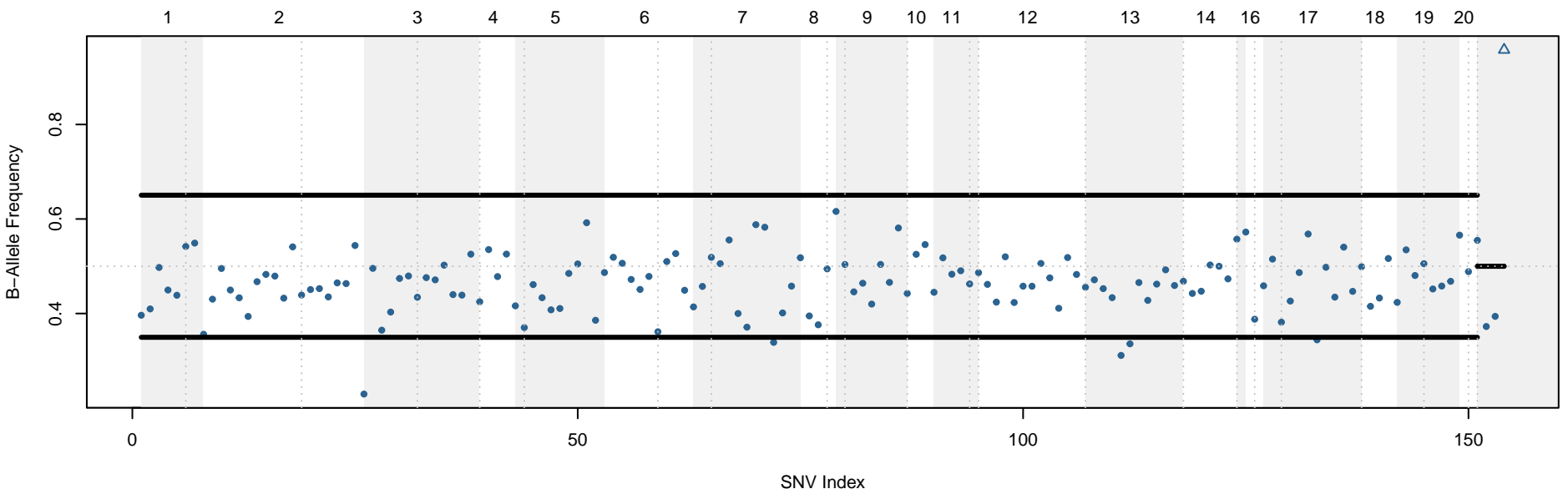




Purity: 0.86 Tumor ploidy: 3.042



Purity: 0.86 Tumor ploidy: 3.042 SNV log-likelihood: -1309.13 GoF: 34.4% Mean coverage: 401;62



SCNA-fit log-likelihood: -15052.71

