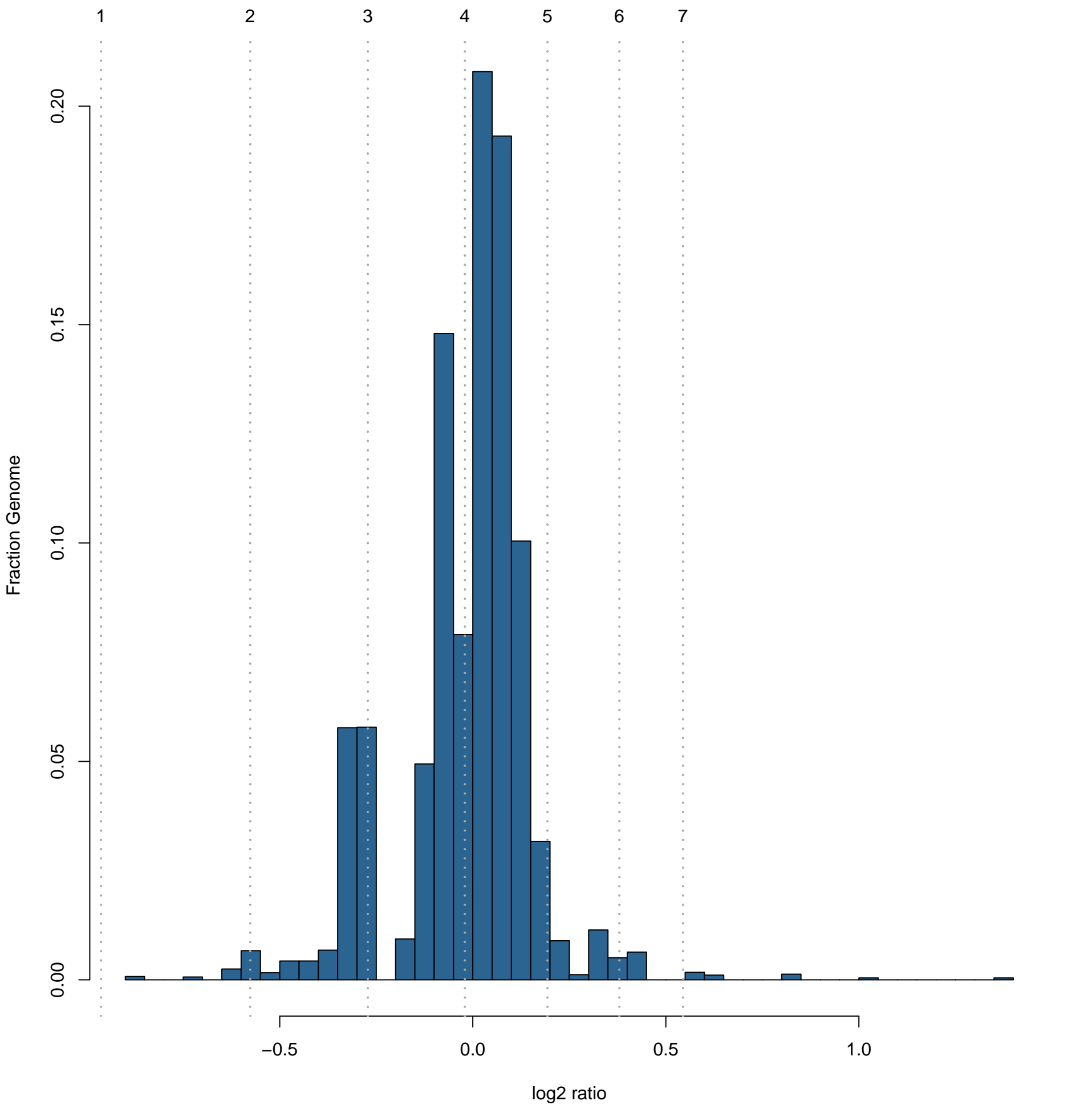
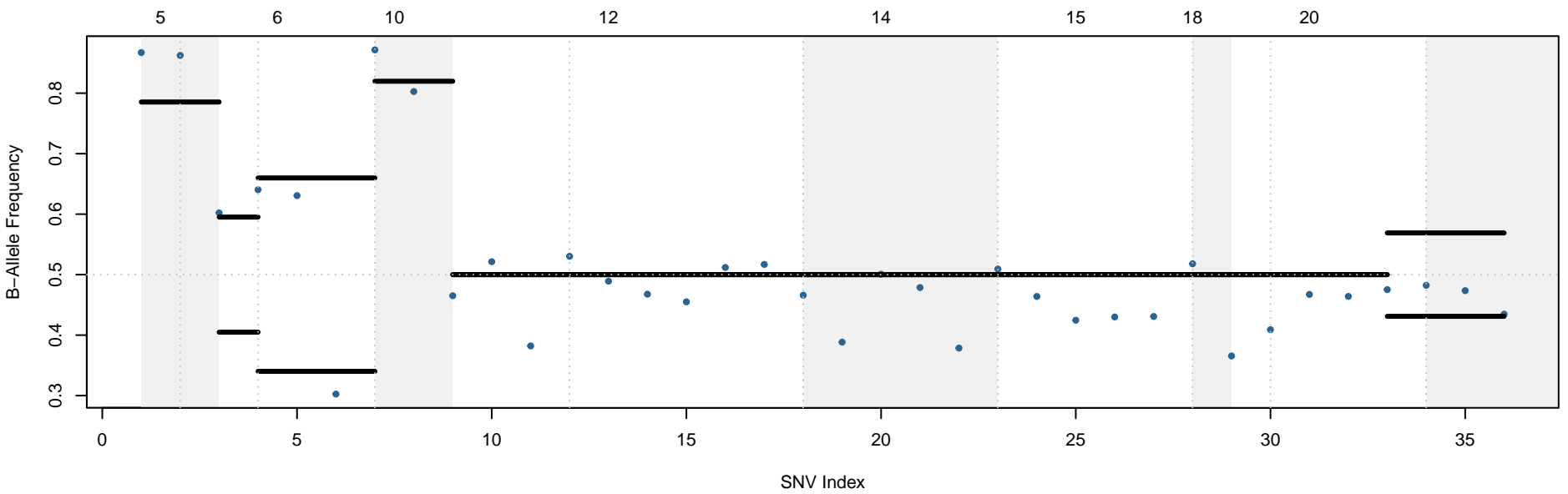


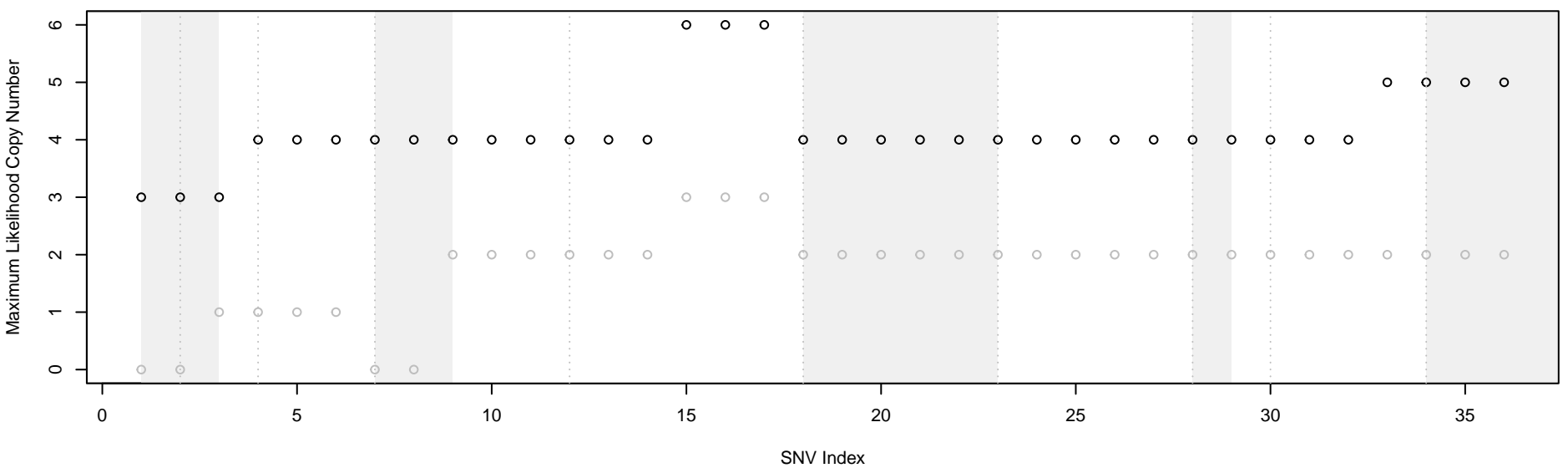
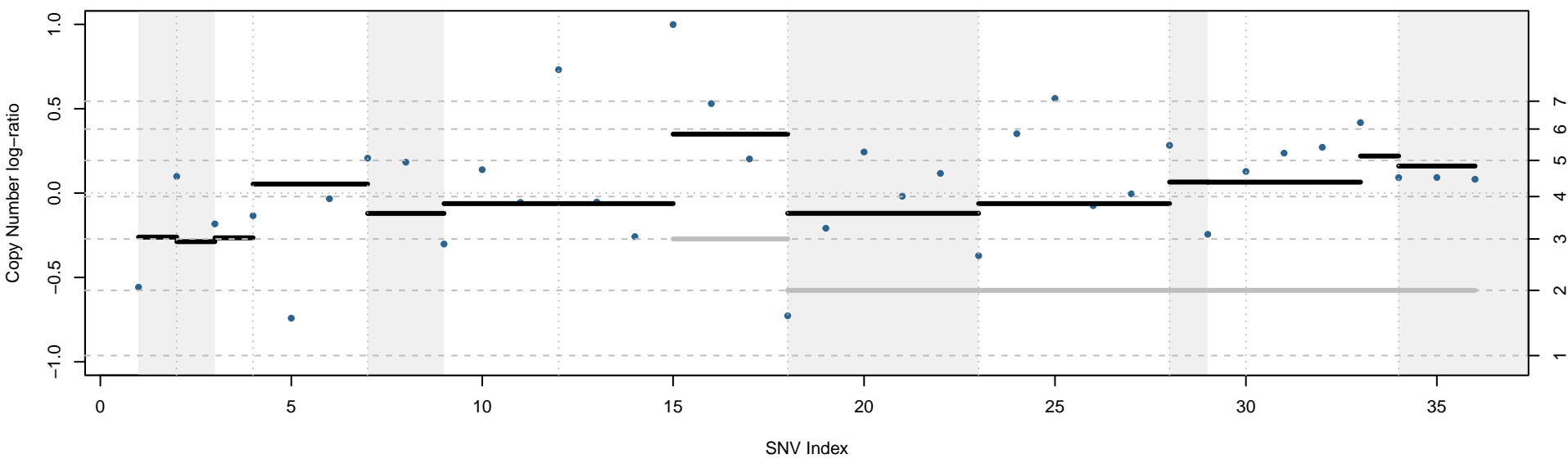
Purity: 0.47 Tumor ploidy: 4.09

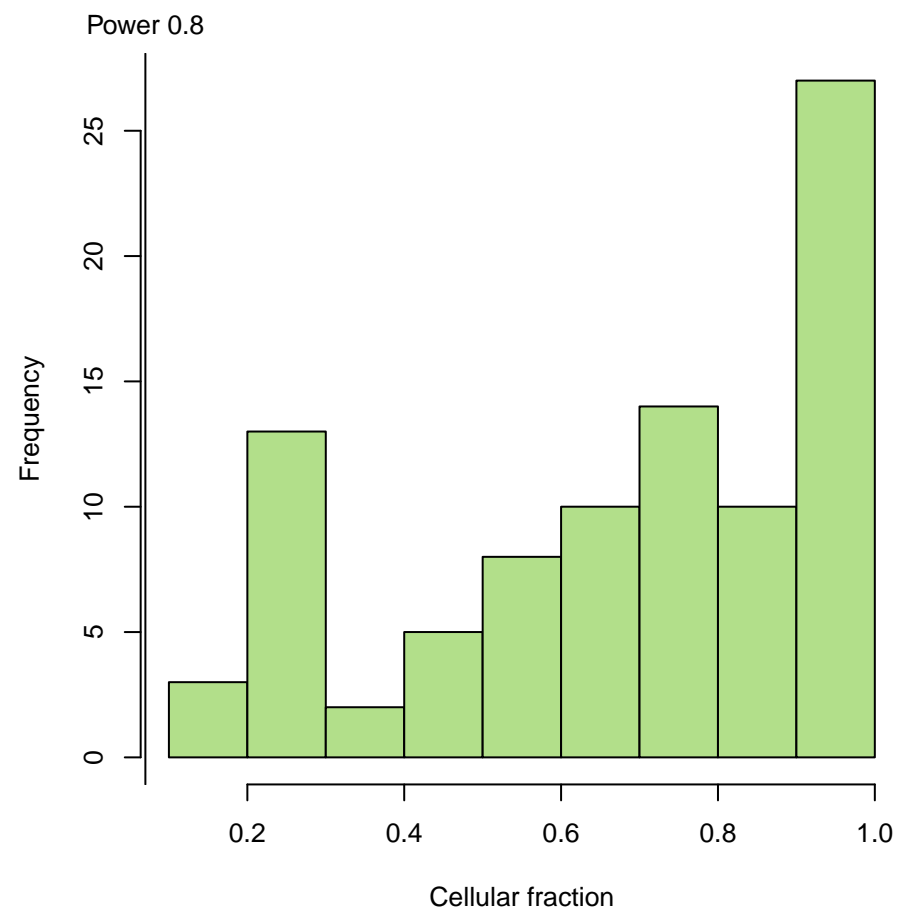
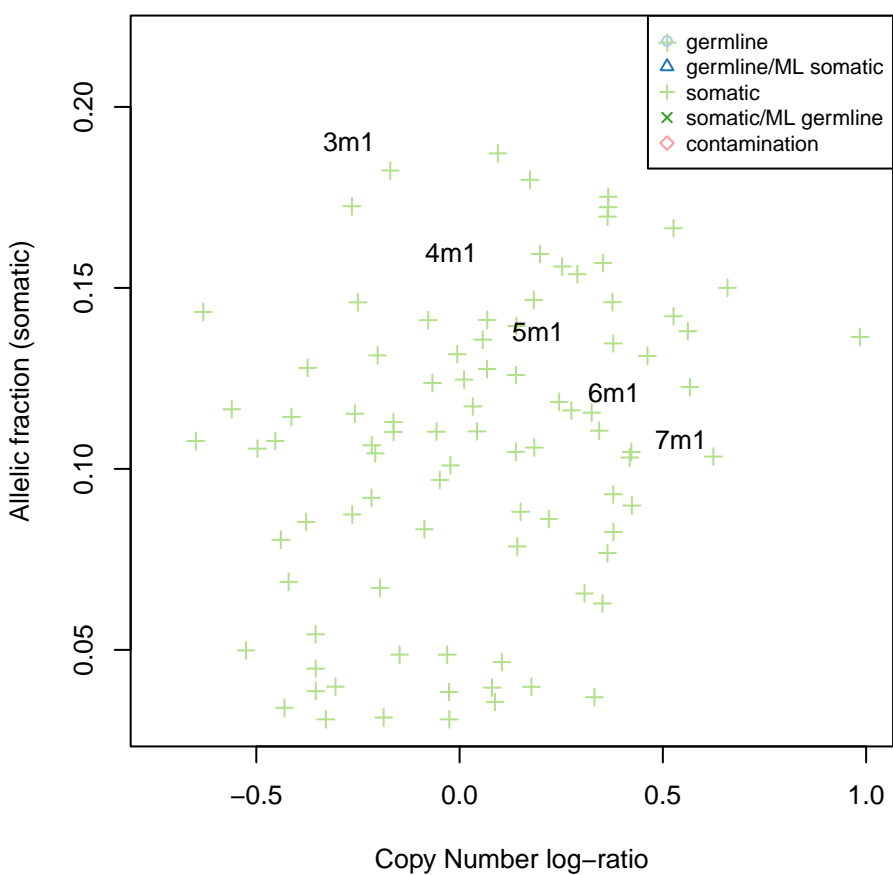
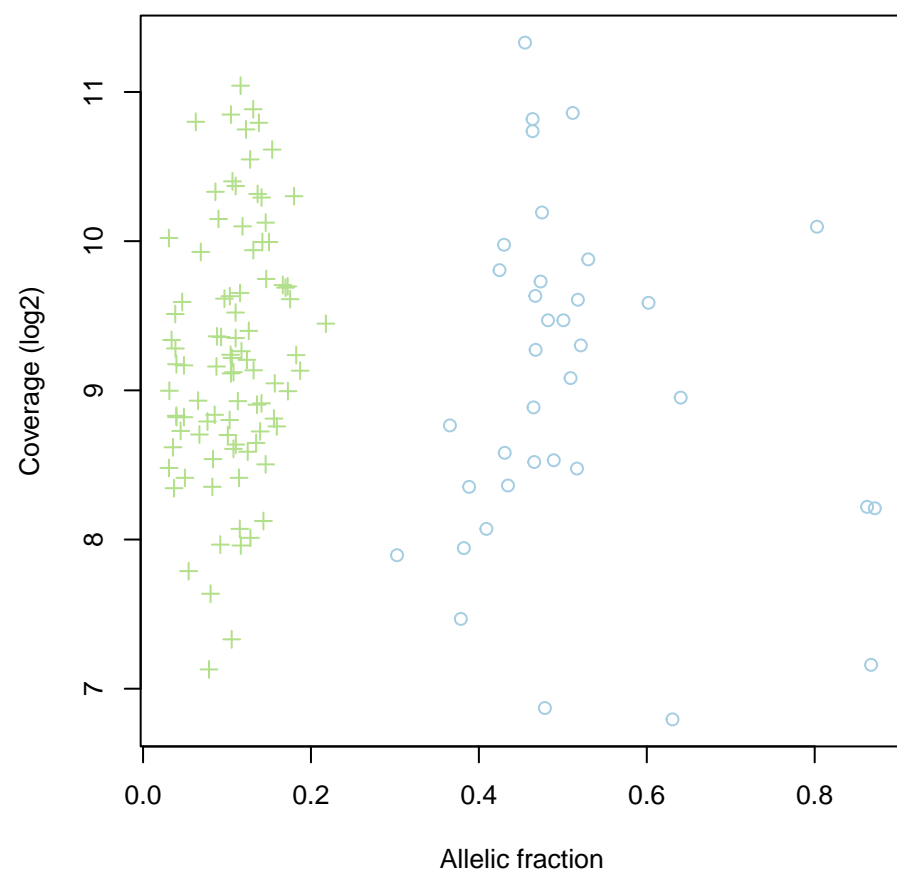
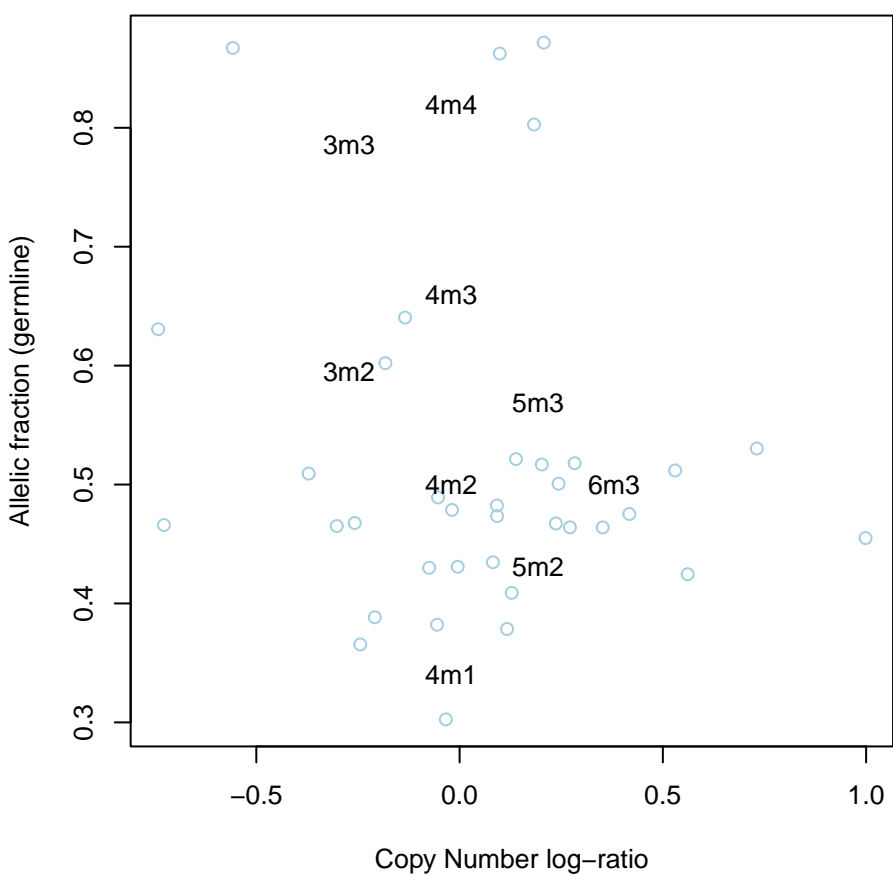


Purity: 0.47 Tumor ploidy: 4.09 SNV log-likelihood: -129.86 GoF: 89.2% Mean coverage: 350;709

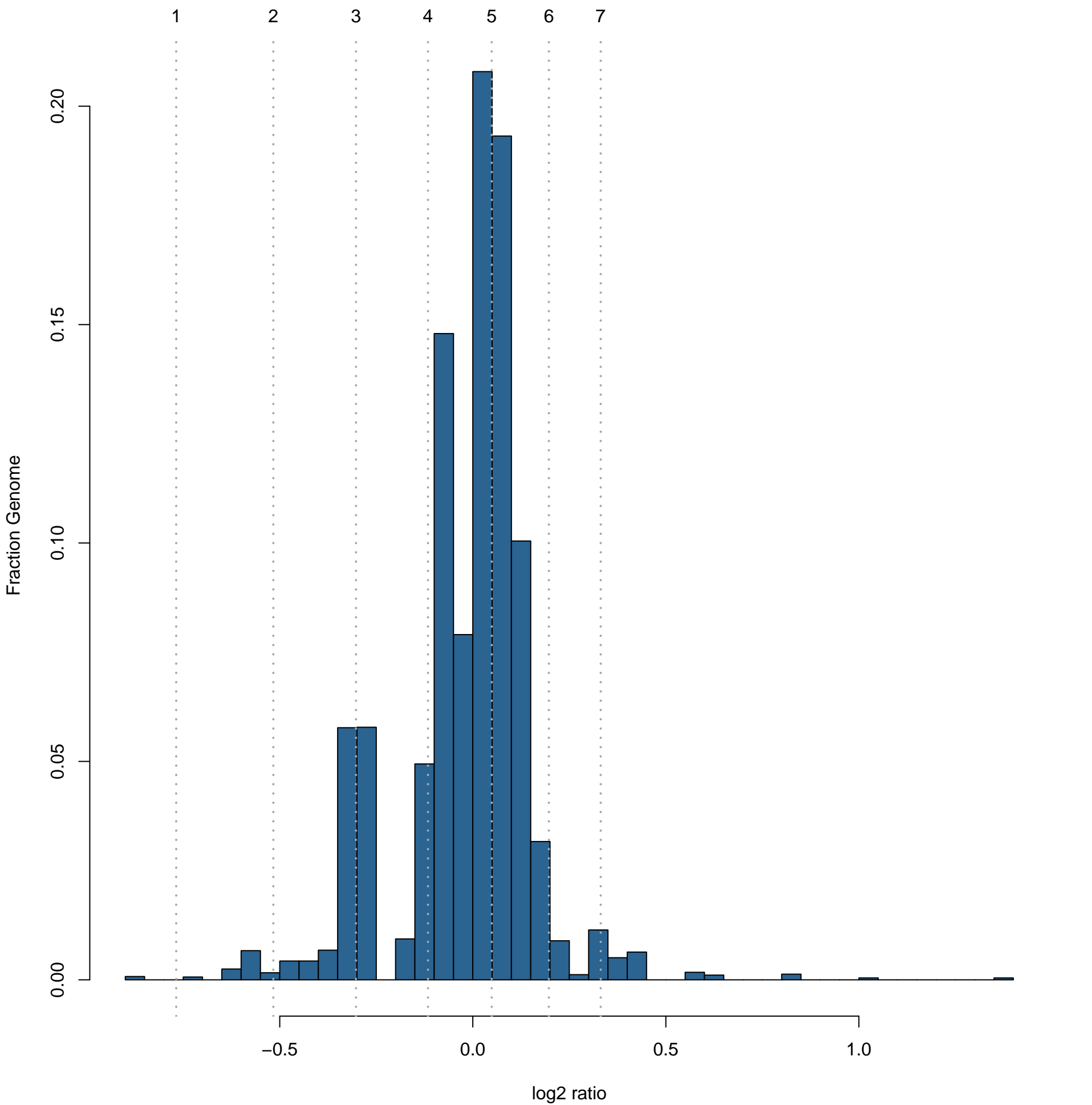


SCNA-fit log-likelihood: -4009.59

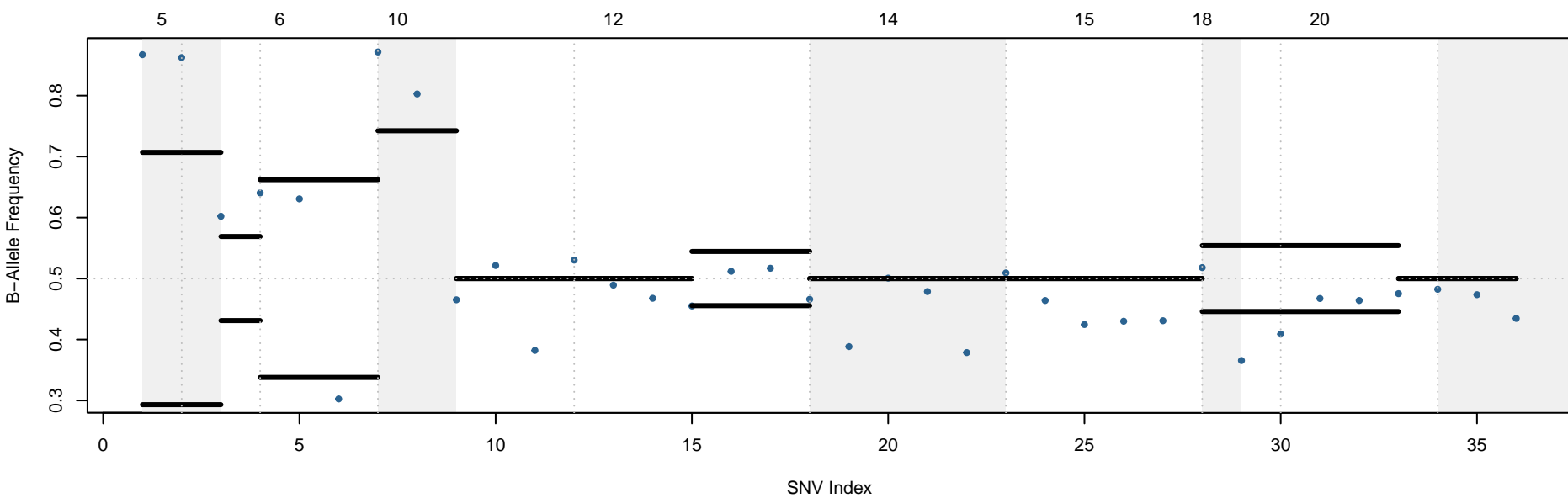




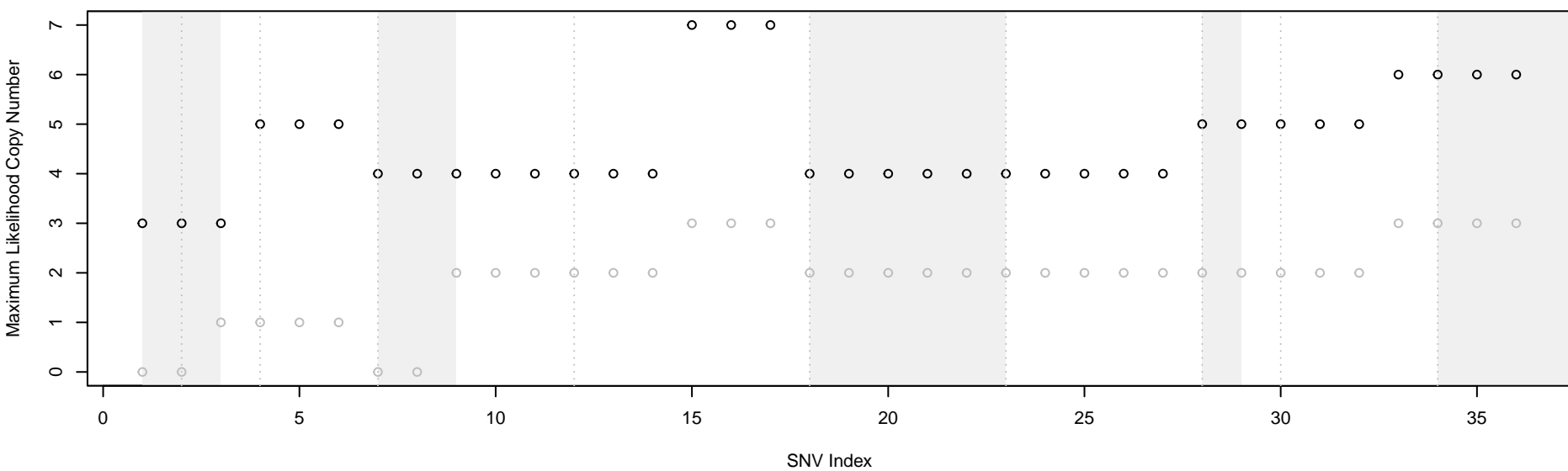
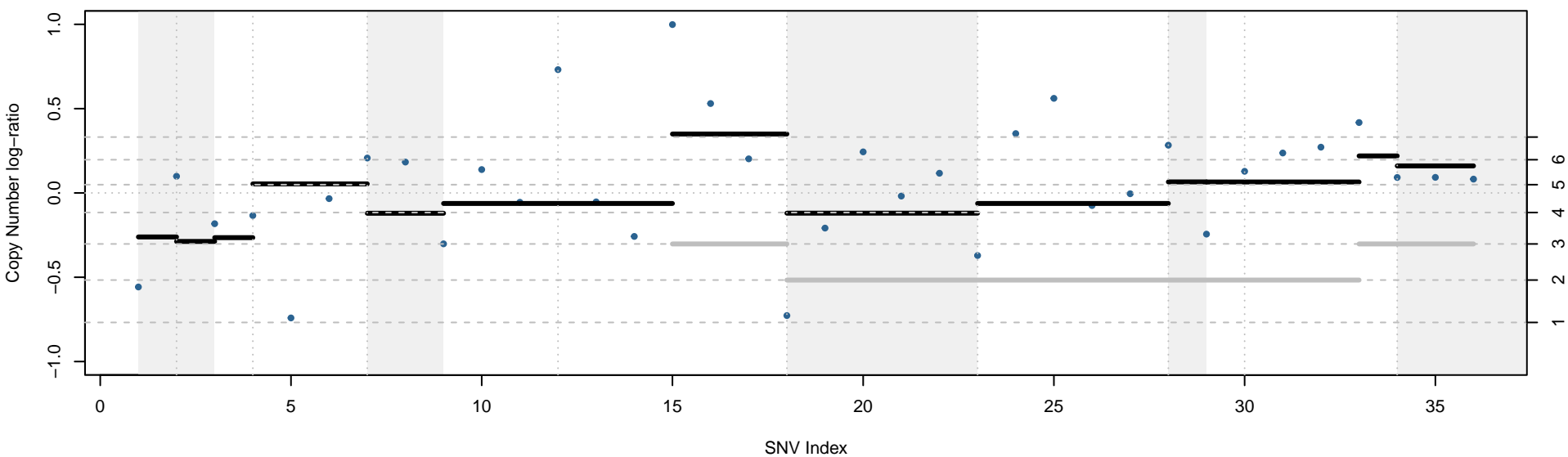
Purity: 0.32 Tumor ploidy: 4.691

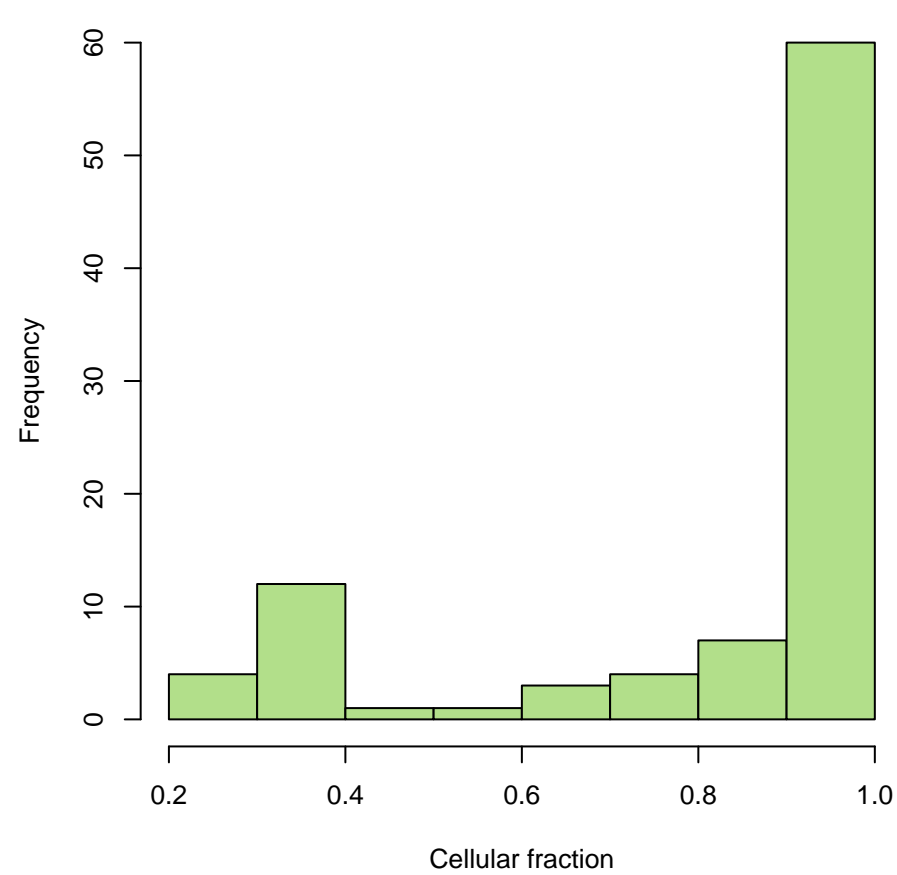
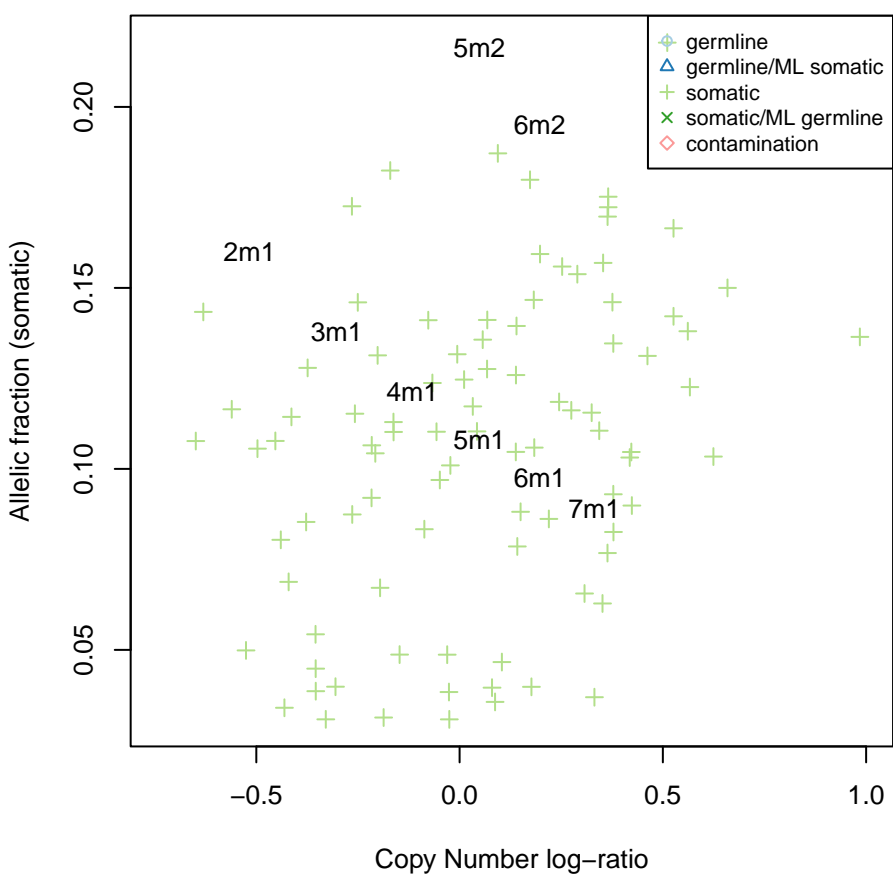
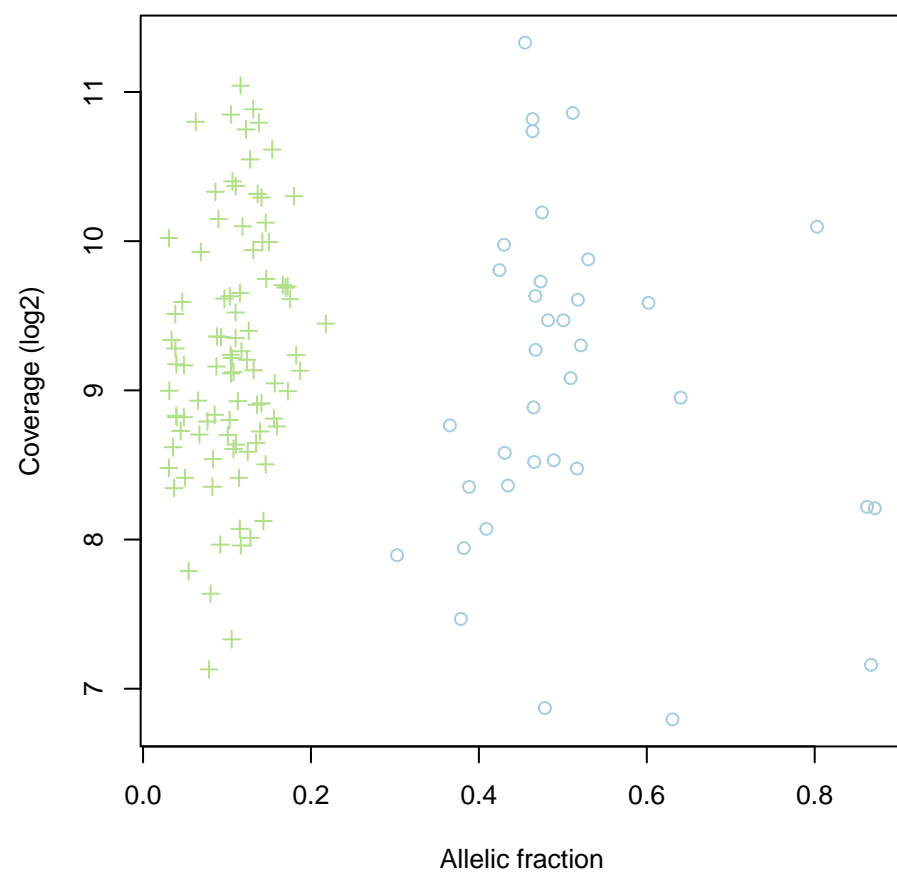
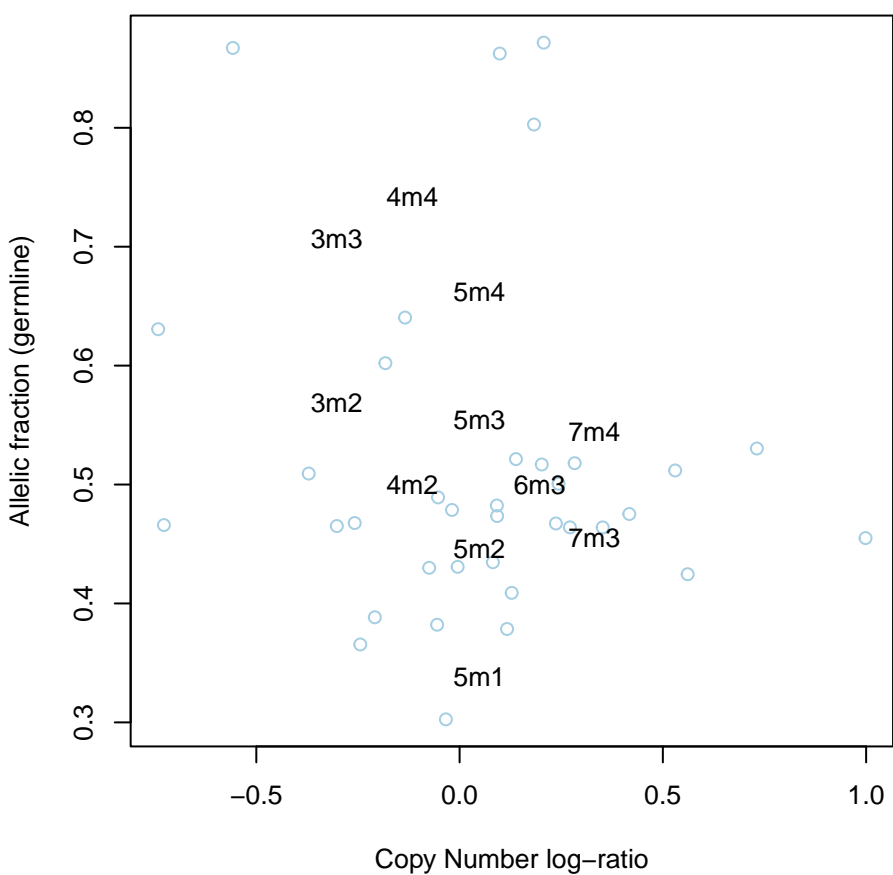


Purity: 0.32 Tumor ploidy: 4.691 SNV log-likelihood: -144.81 GoF: 93.3% Mean coverage: 350,709

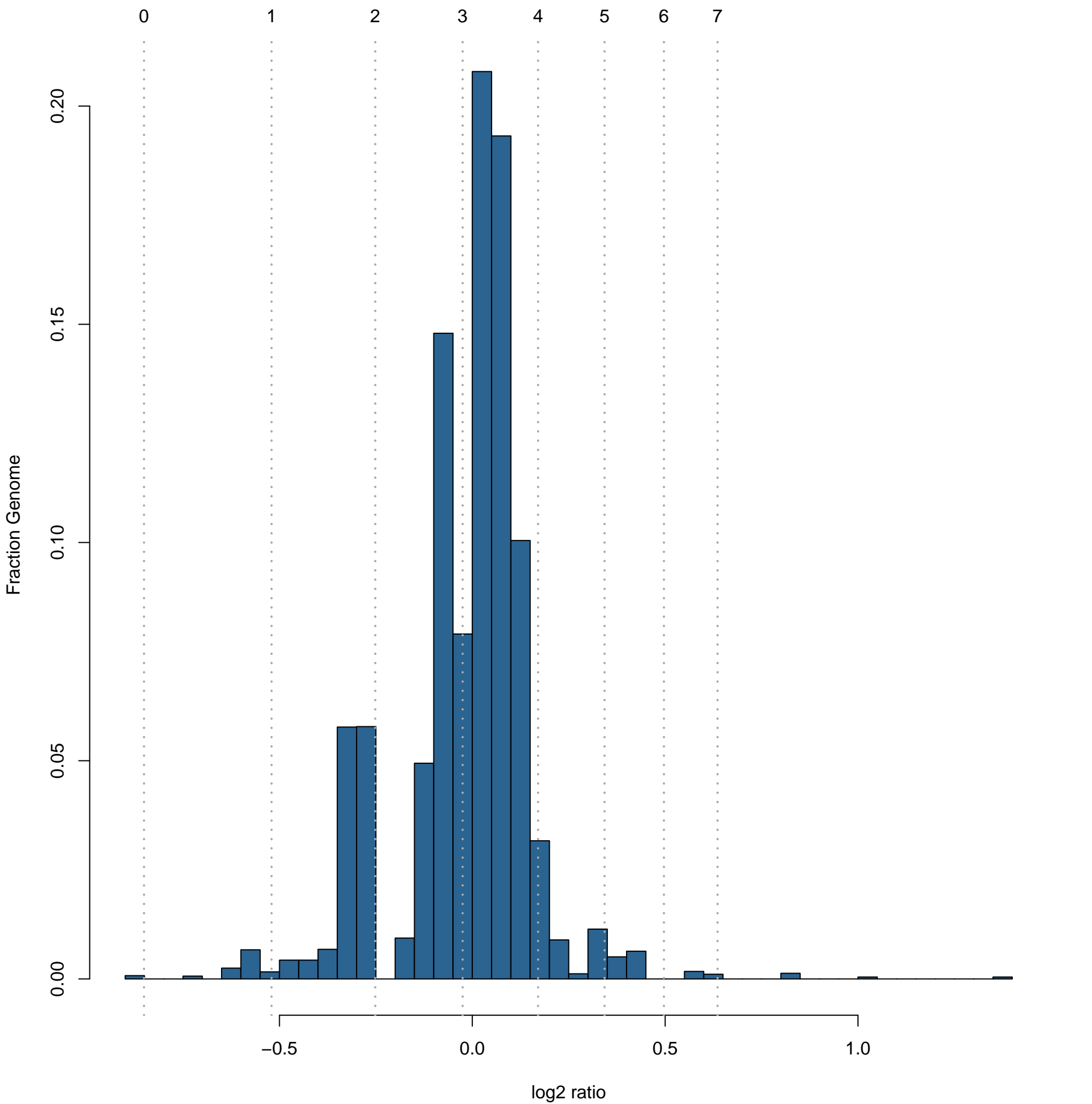


SCNA-fit log-likelihood: -3980.7

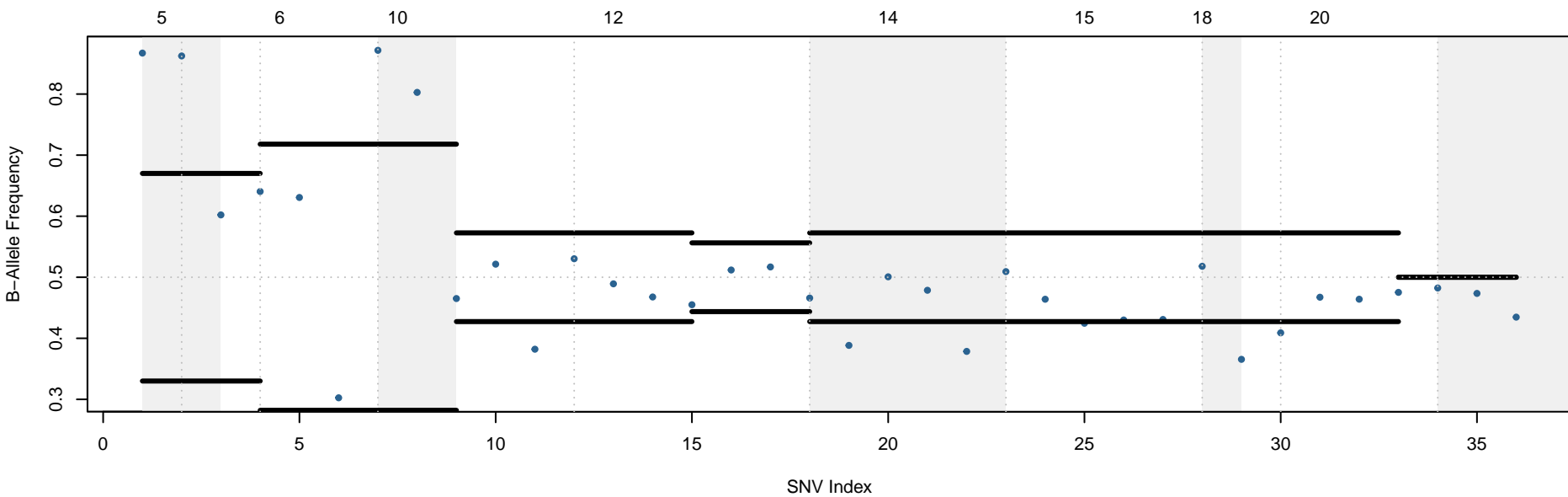




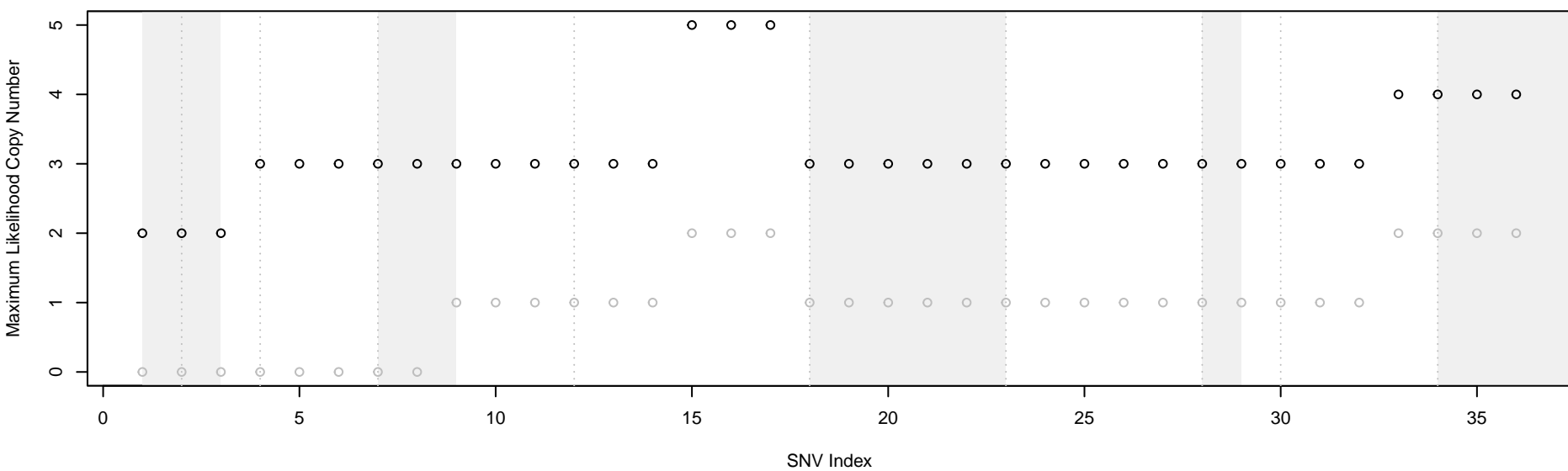
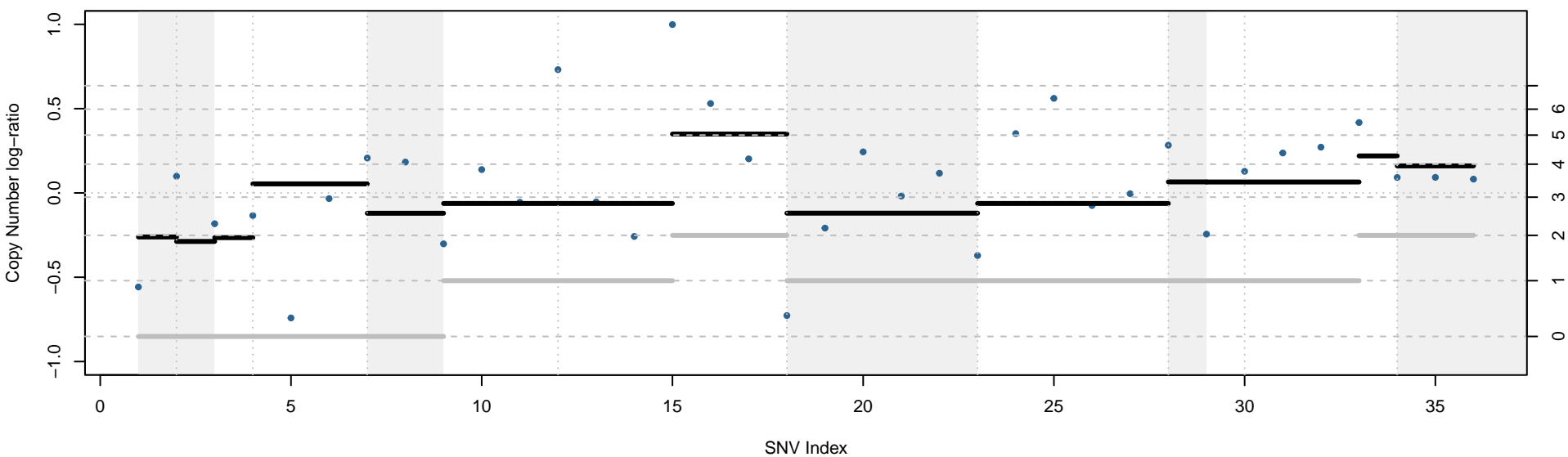
Purity: 0.34 Tumor ploidy: 3.121



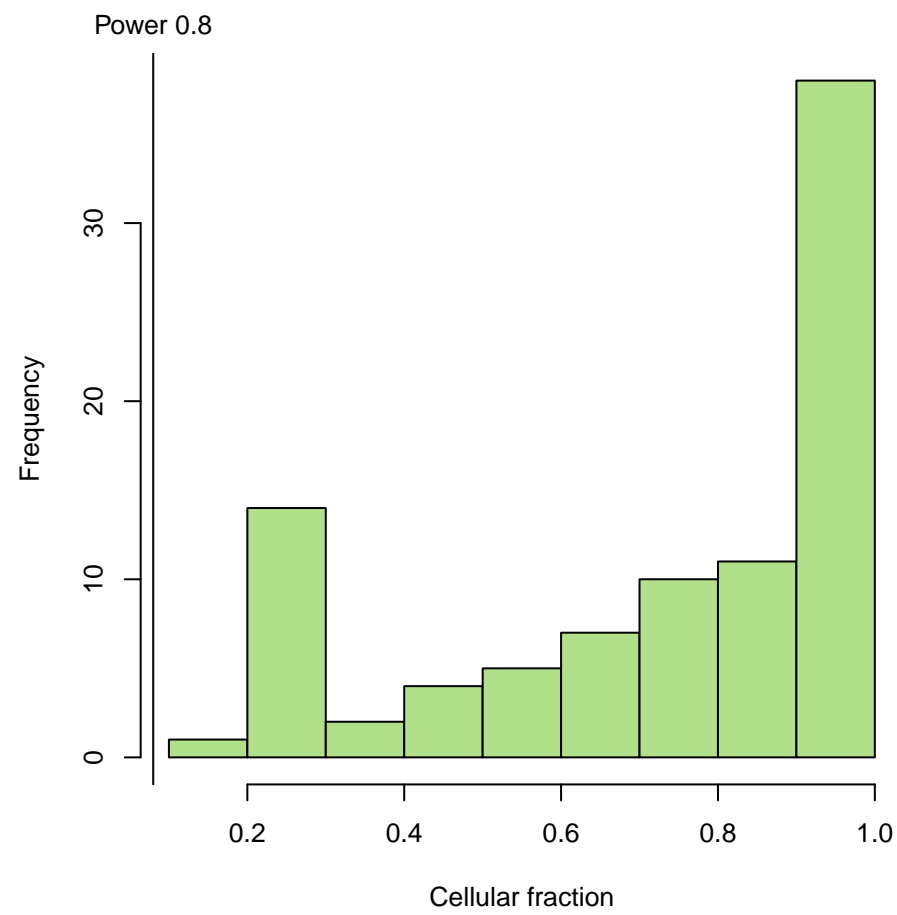
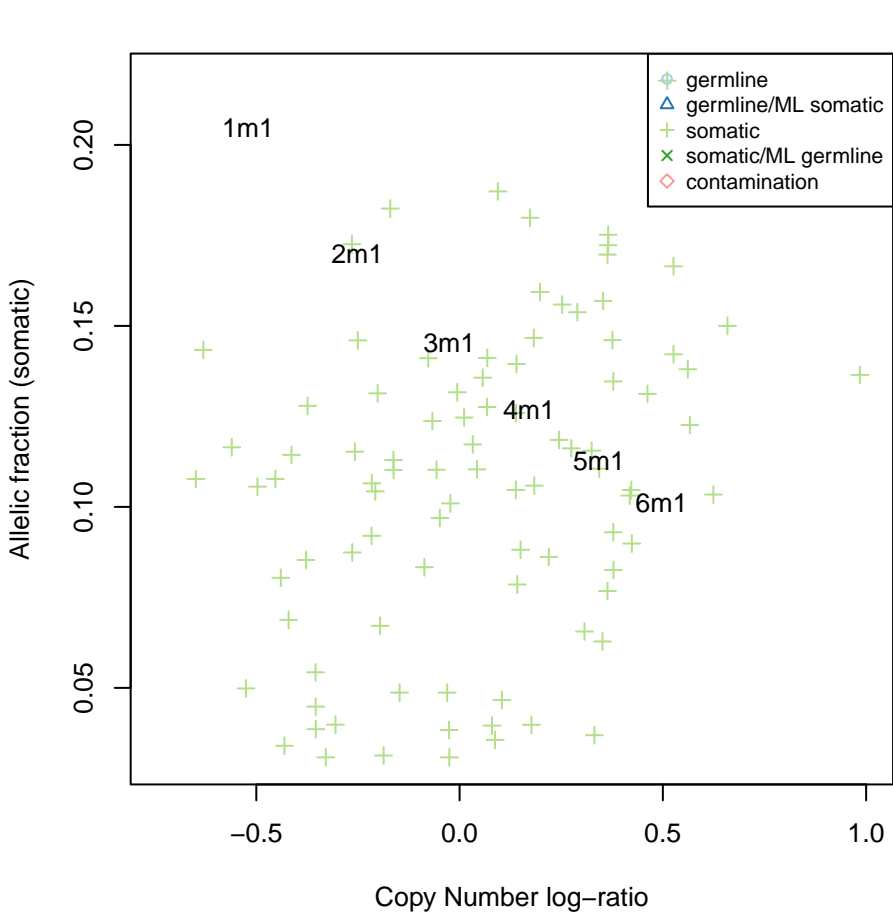
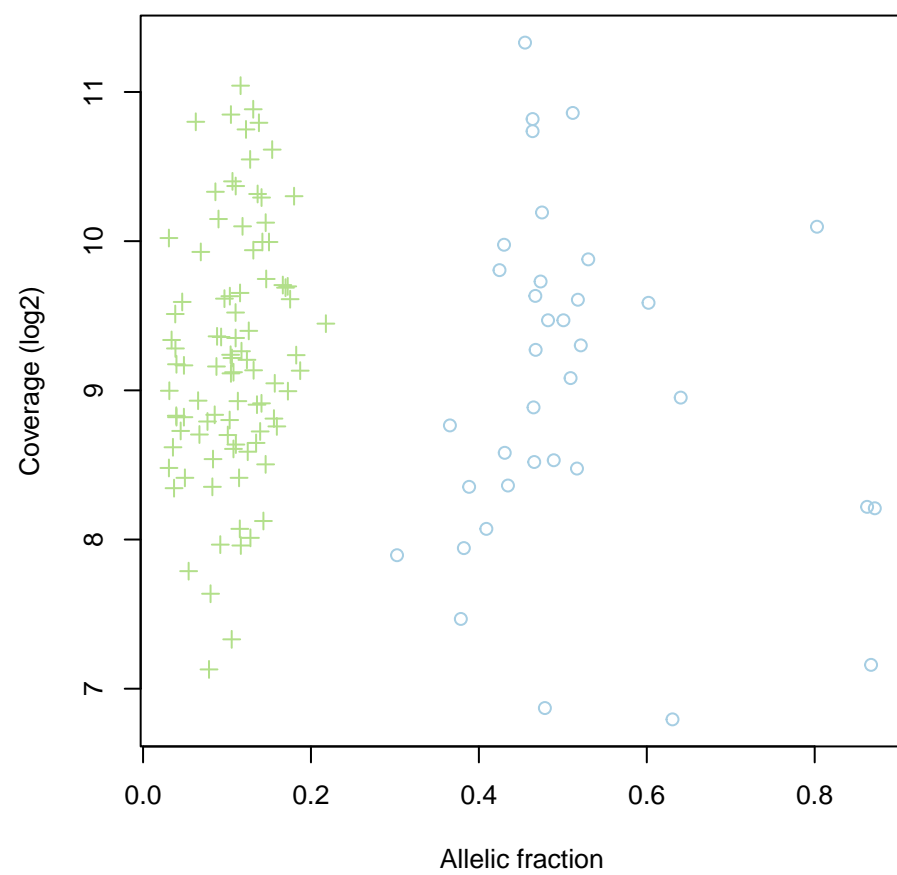
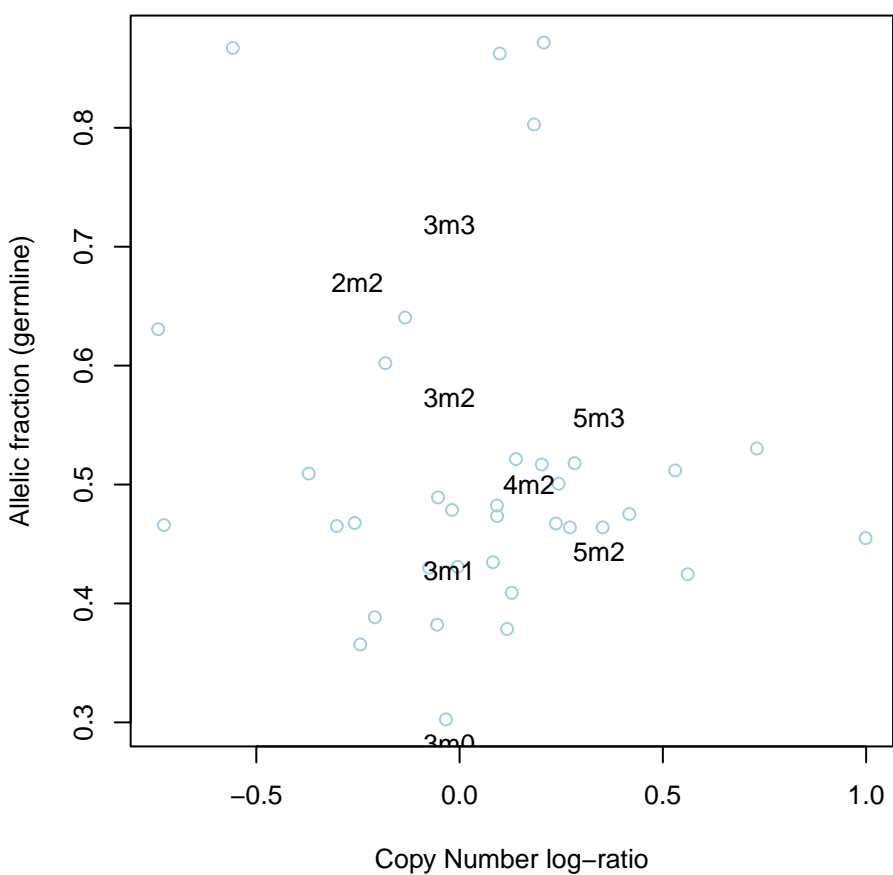
Purity: 0.34 Tumor ploidy: 3.121 SNV log-likelihood: -142.38 GoF: 90% Mean coverage: 350;709



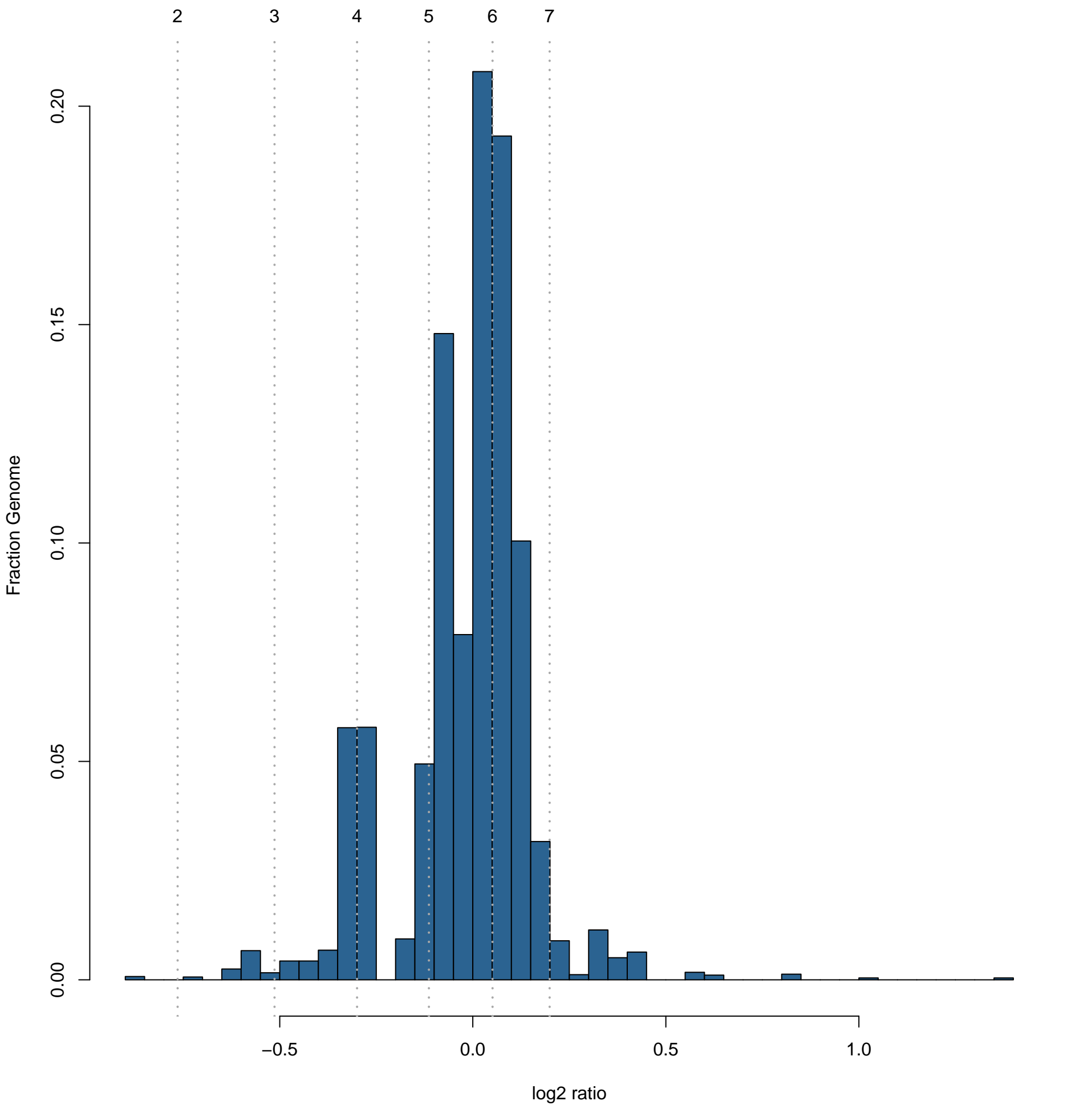
SCNA-fit log-likelihood: -3992.64



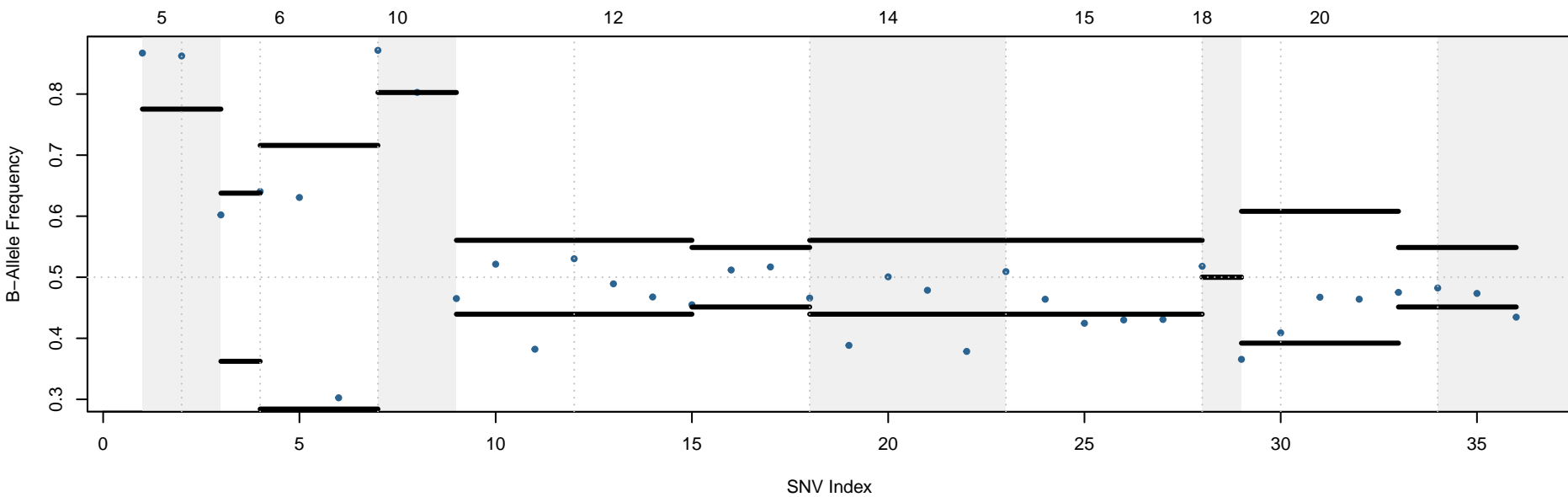




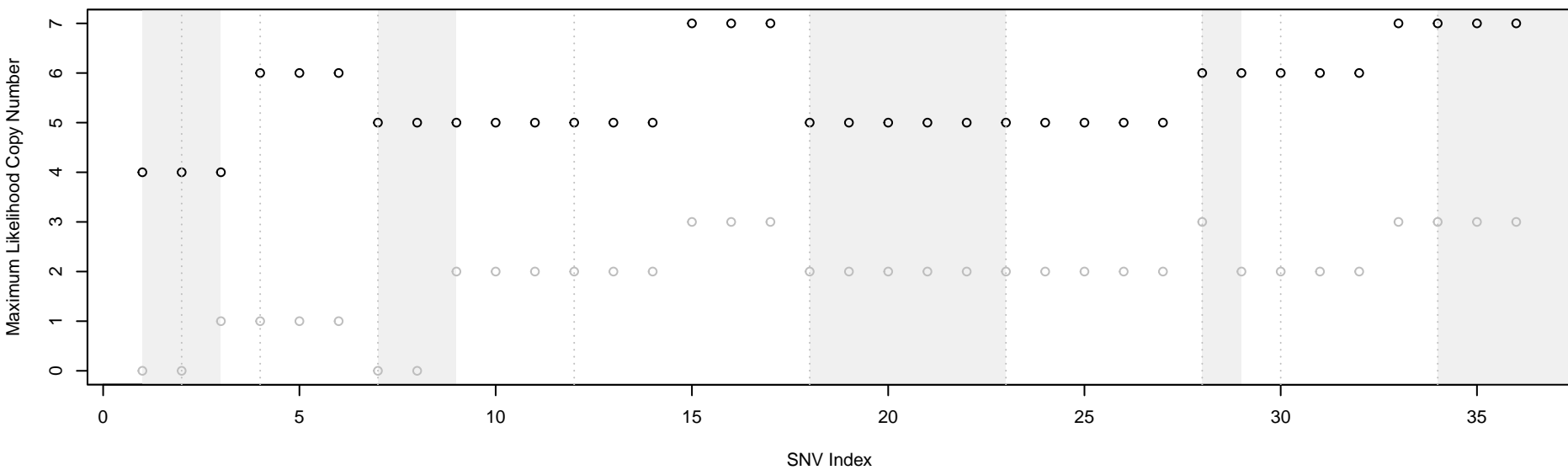
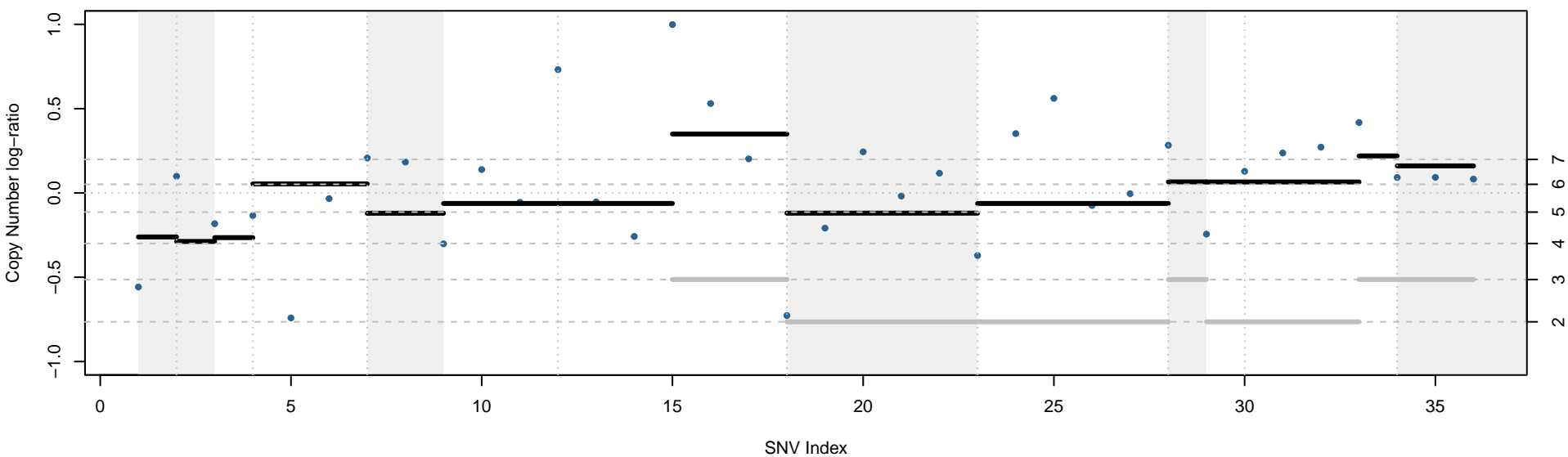
Purity: 0.38 Tumor ploidy: 5.677

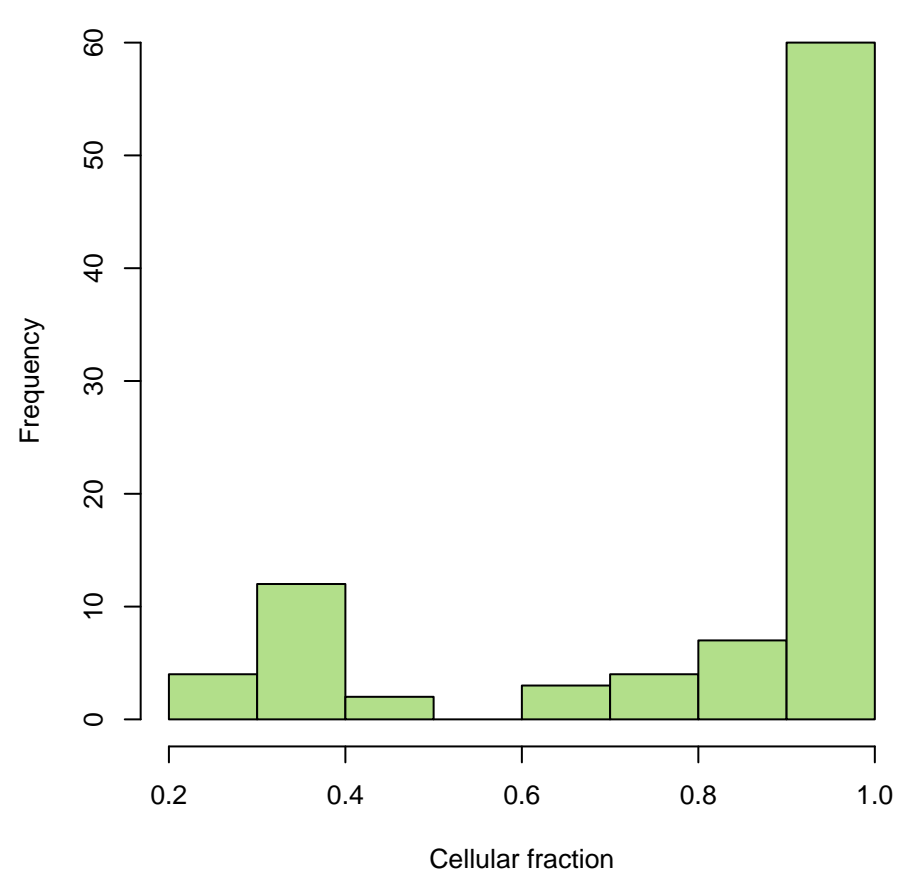
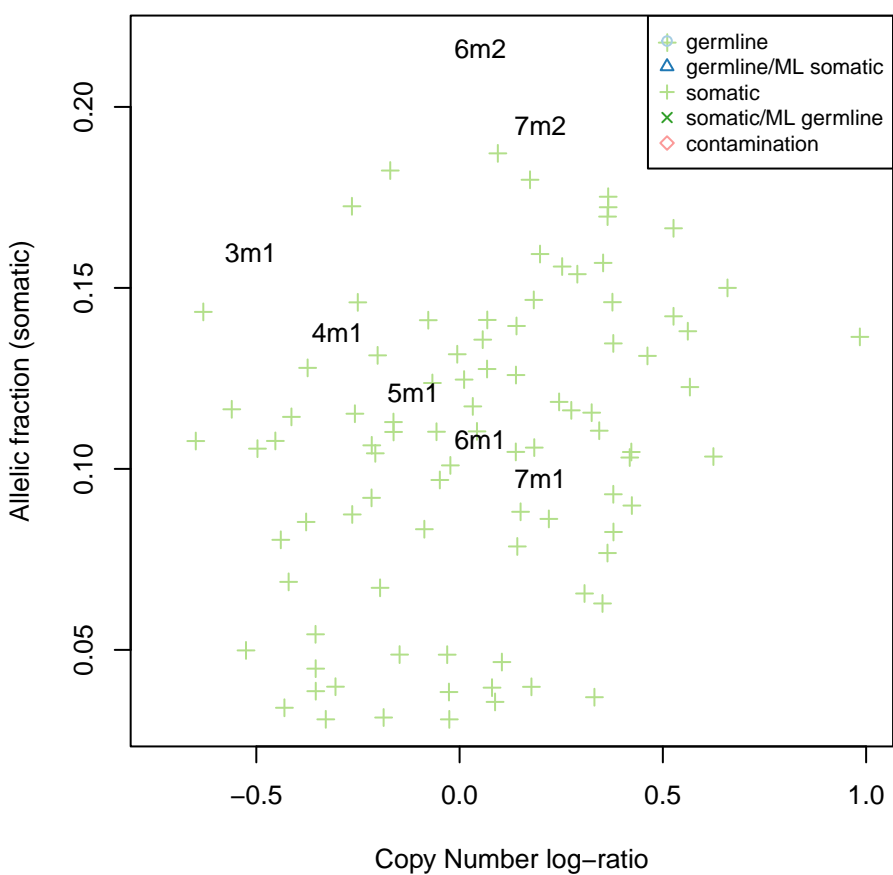
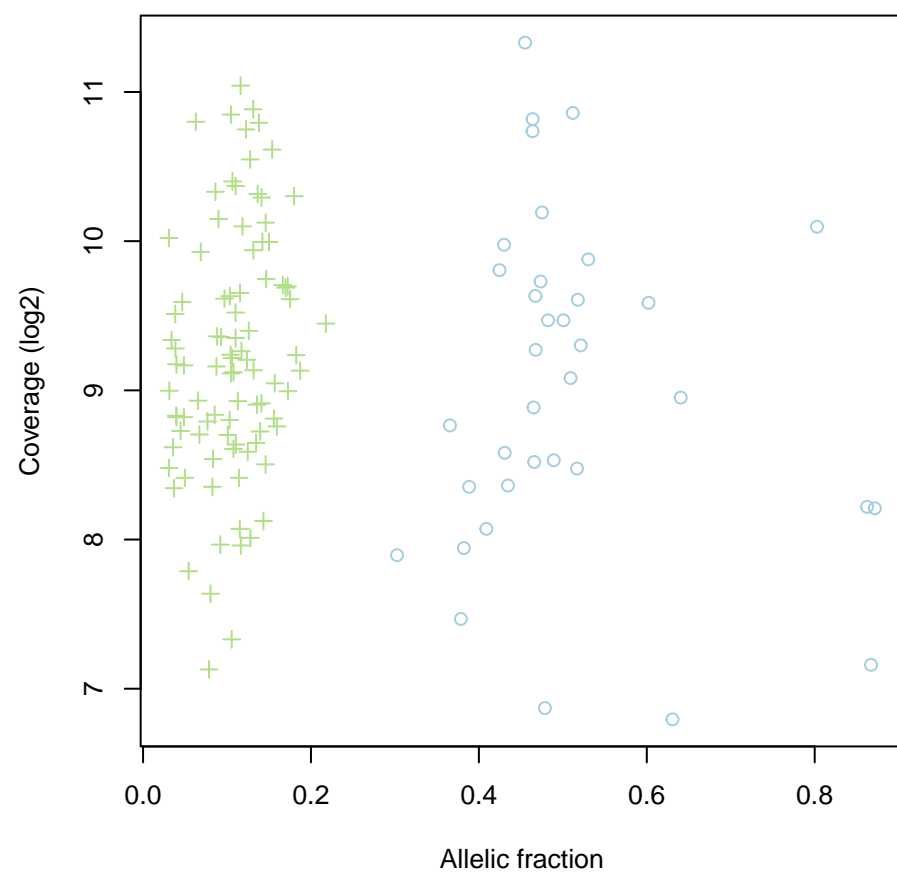
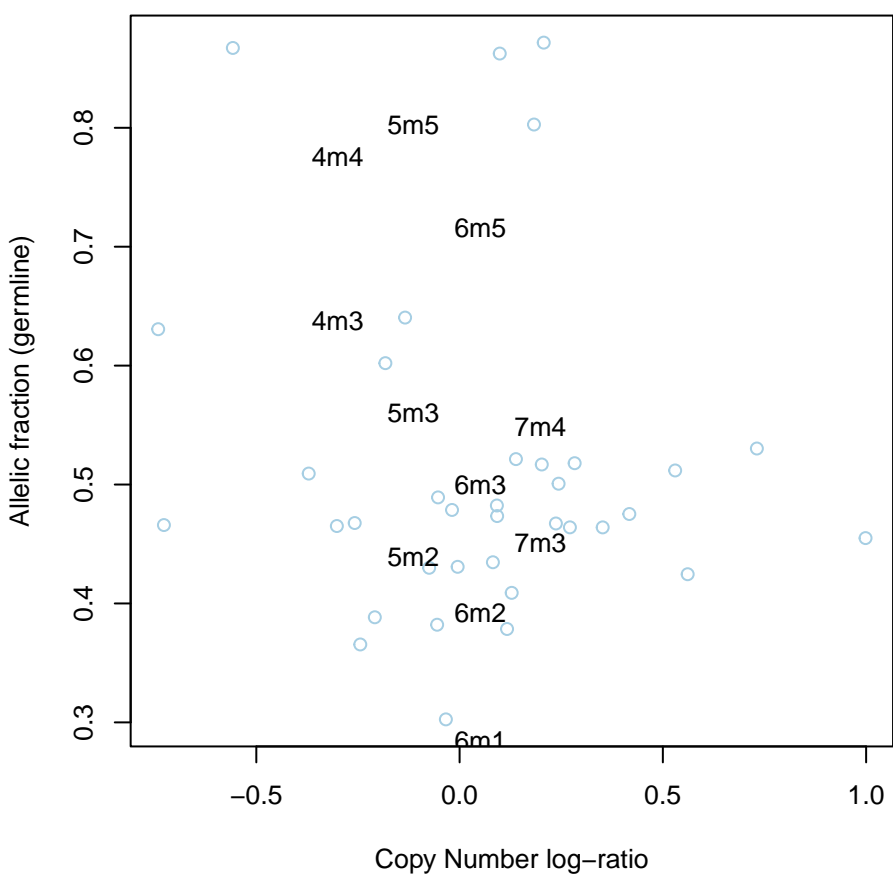


Purity: 0.38 Tumor ploidy: 5.677 SNV log-likelihood: -130.92 GoF: 94.7% Mean coverage: 350;709

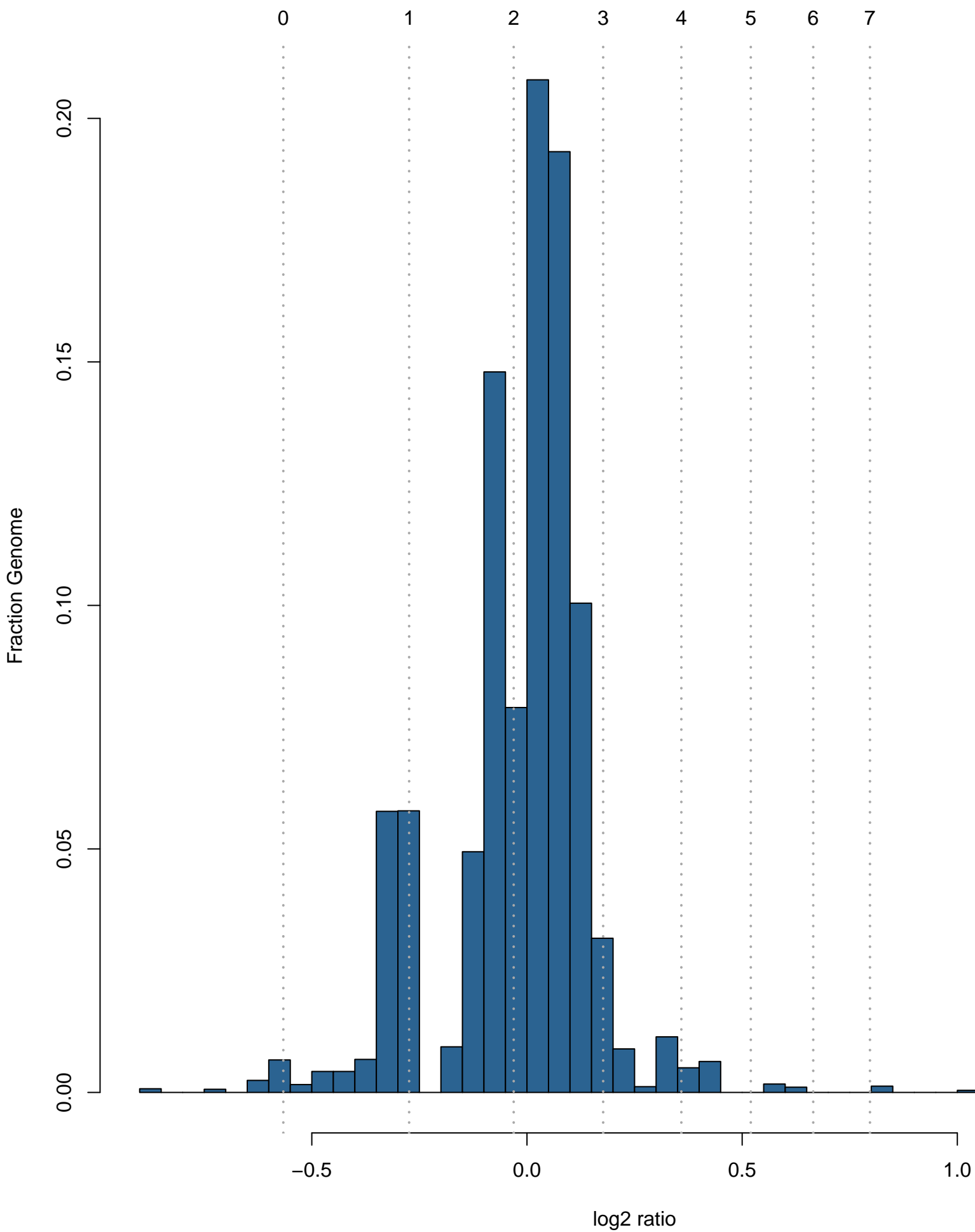


SCNA-fit log-likelihood: -4059.94

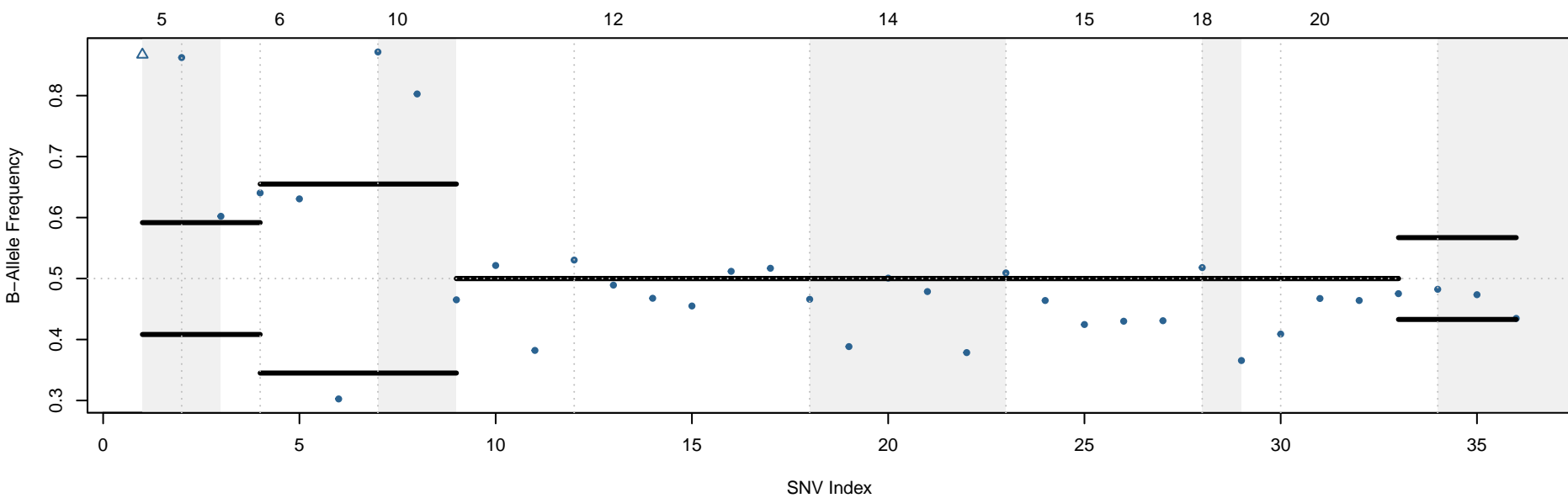




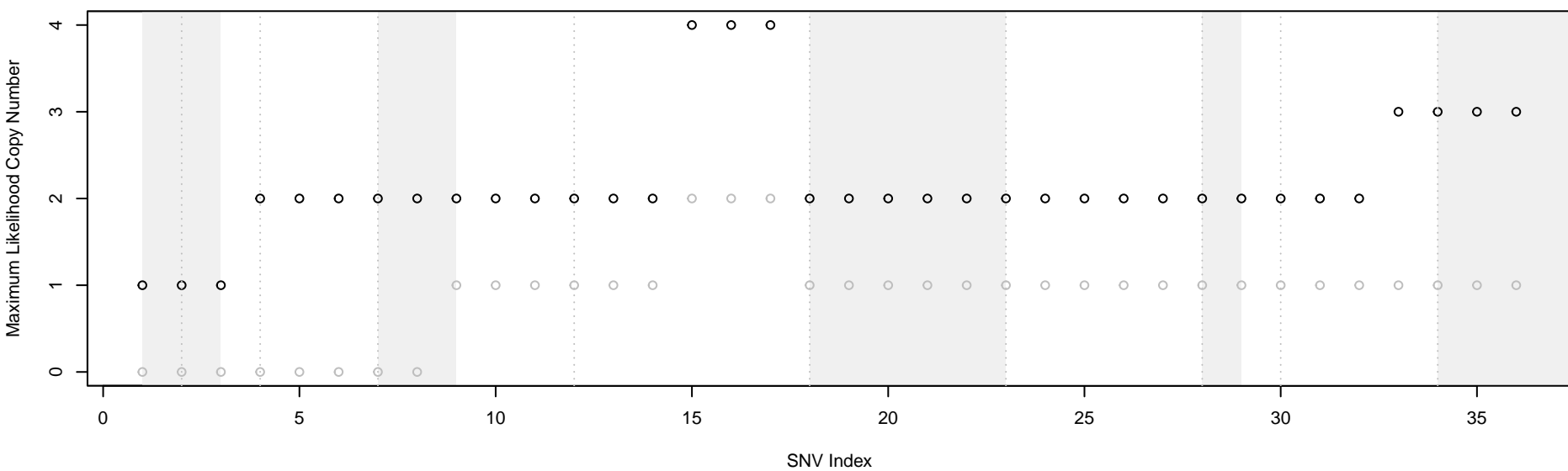
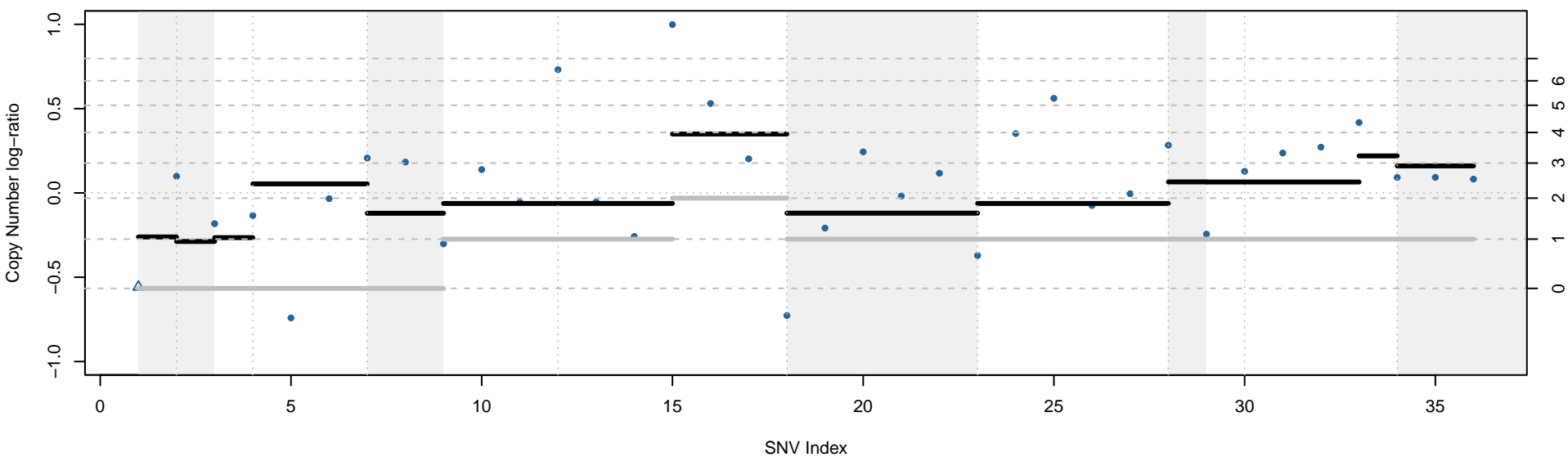
Purity: 0.31 Tumor ploidy: 2.14

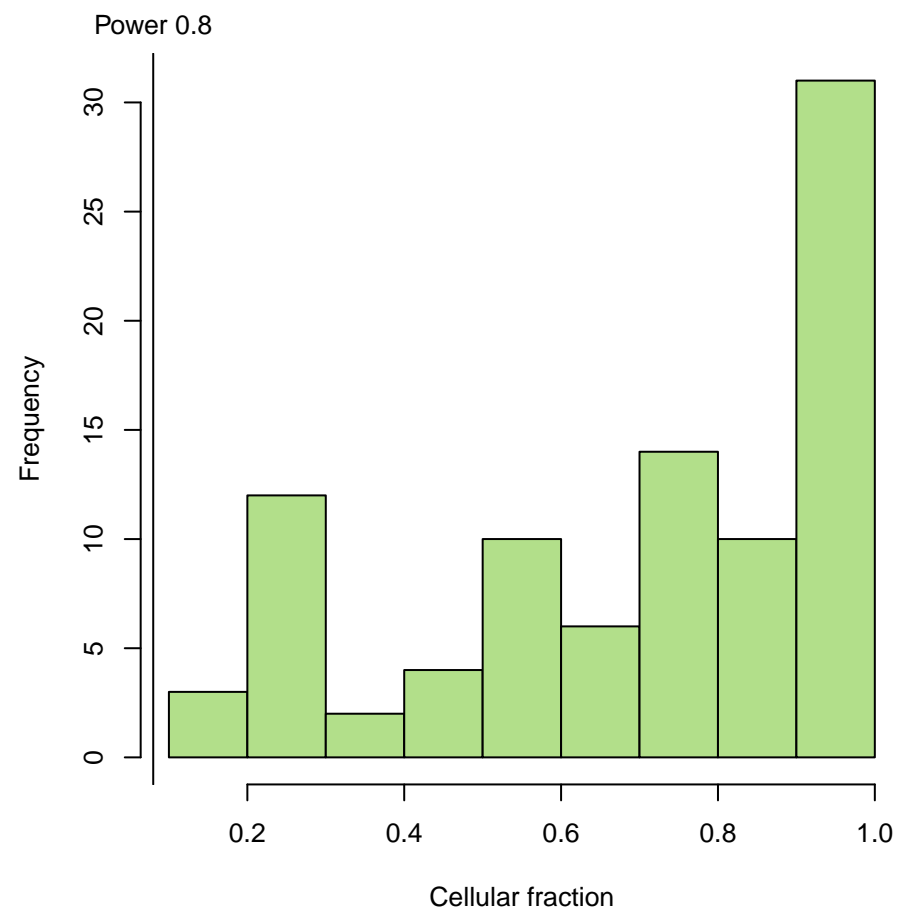
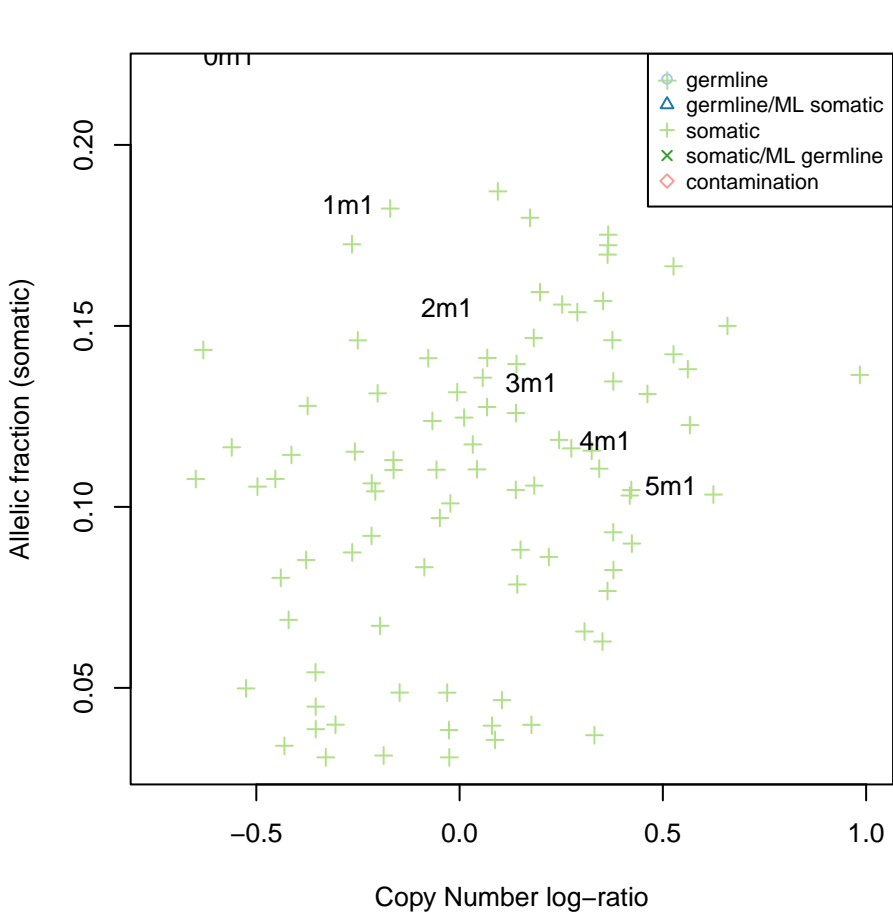
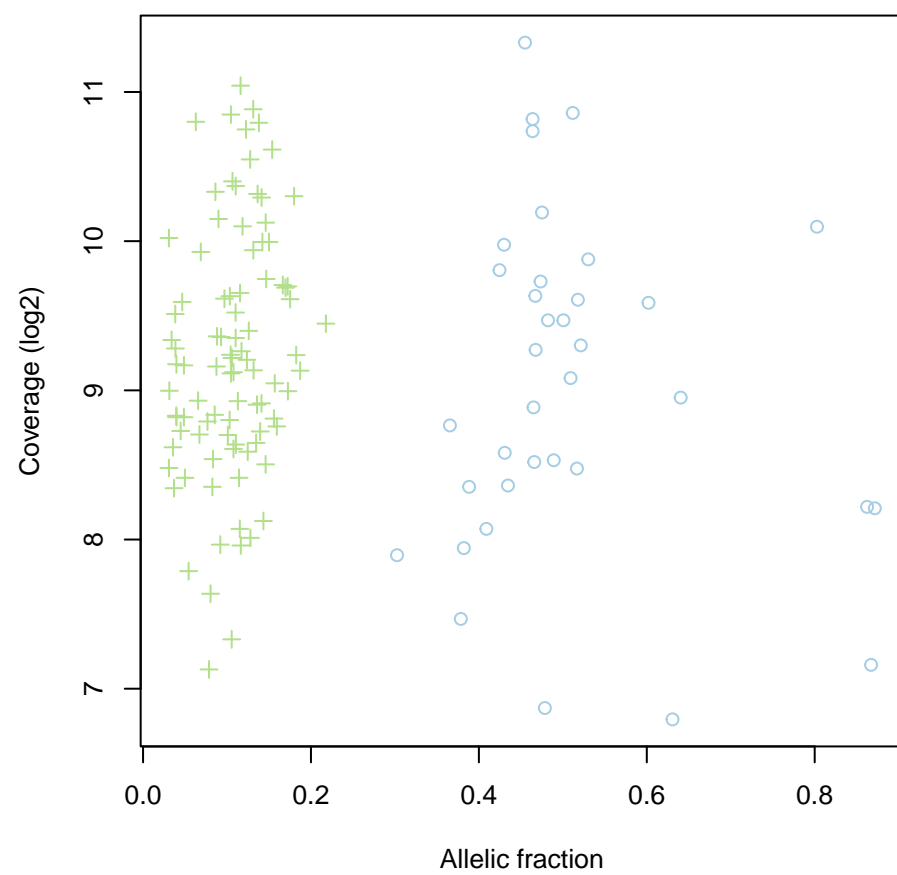
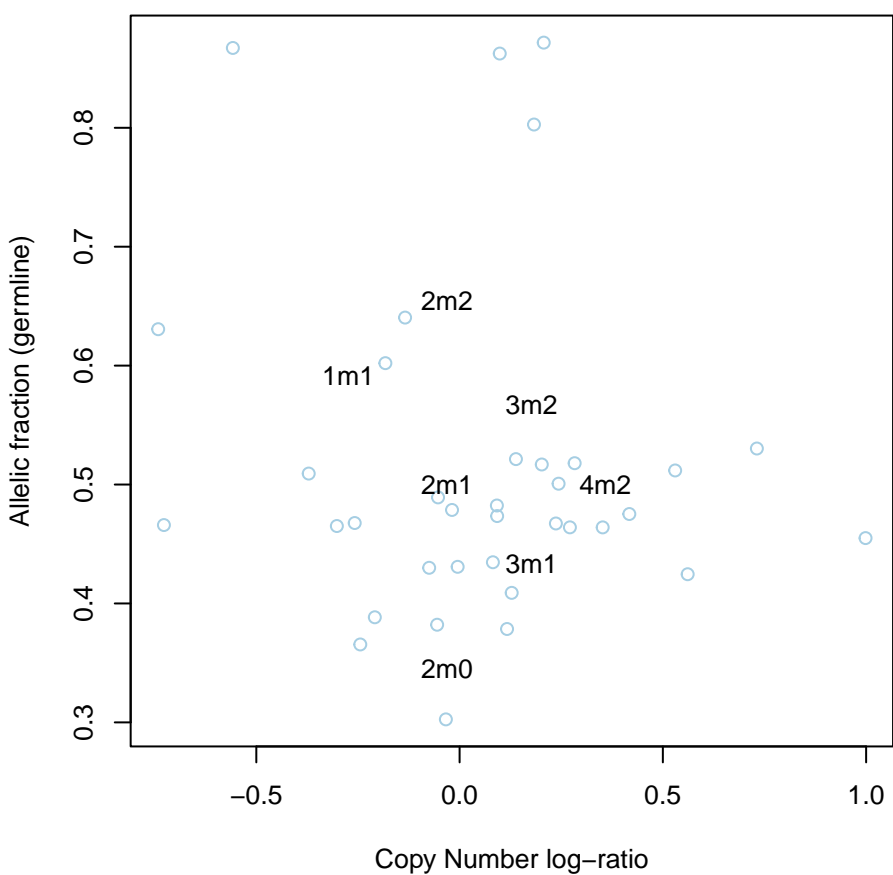


Purity: 0.31 Tumor ploidy: 2.14 SNV log-likelihood: -162.89 GoF: 87.3% Mean coverage: 350;709

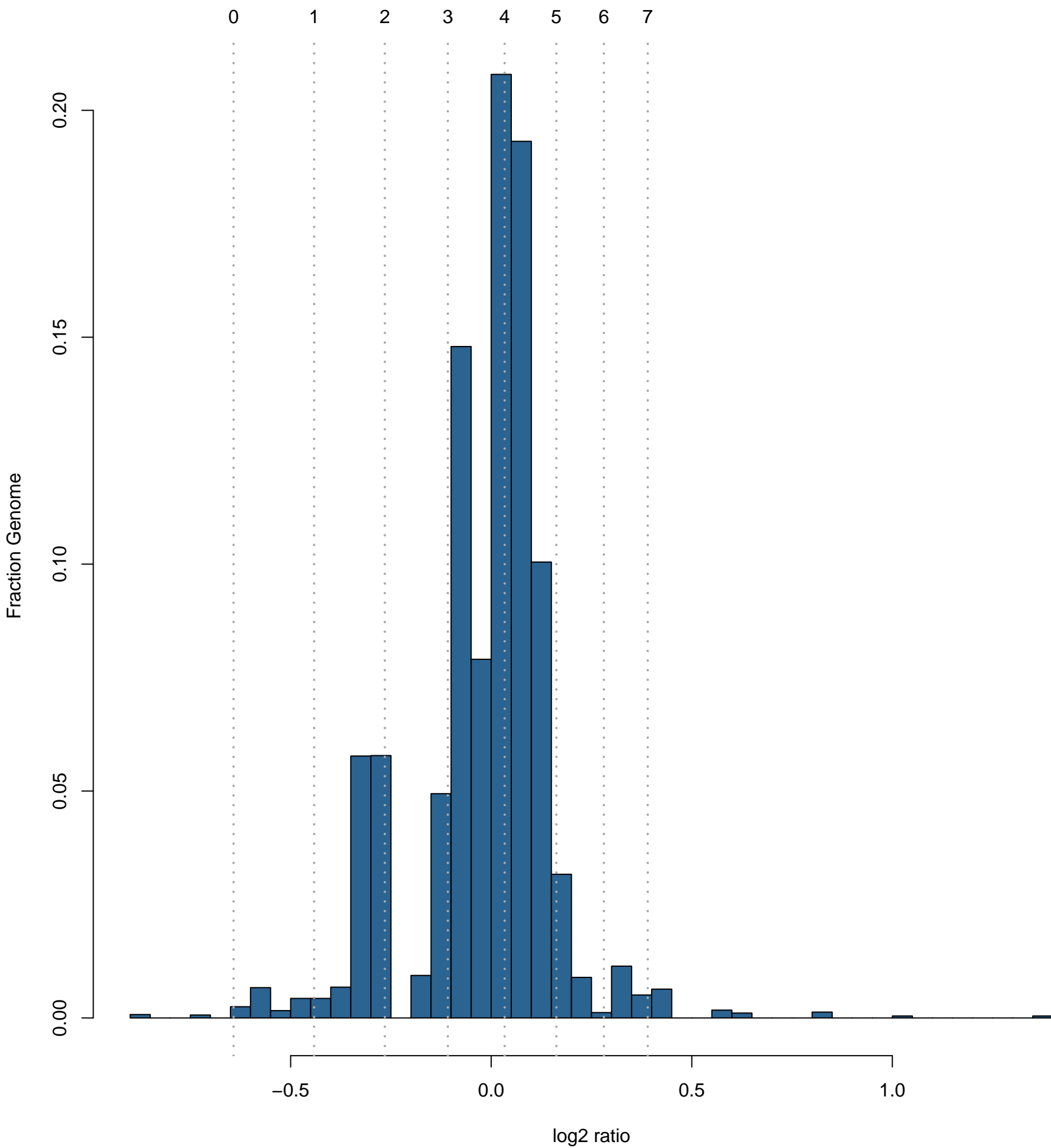


SCNA-fit log-likelihood: -4014.18



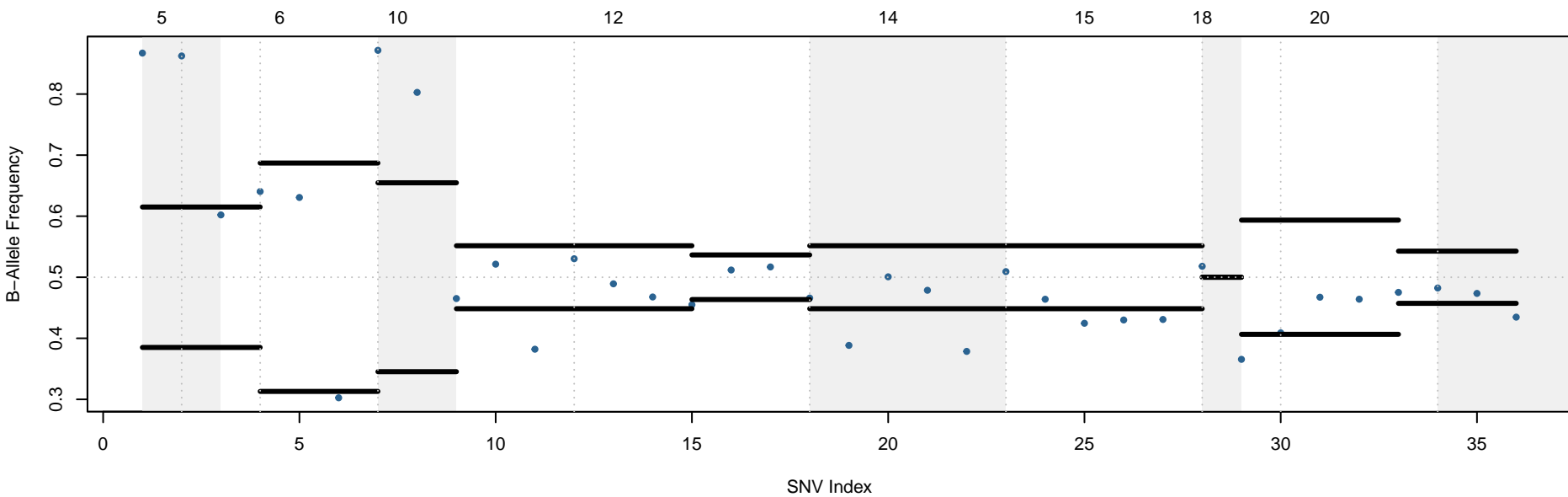


Purity: 0.23 Tumor ploidy: 3.755

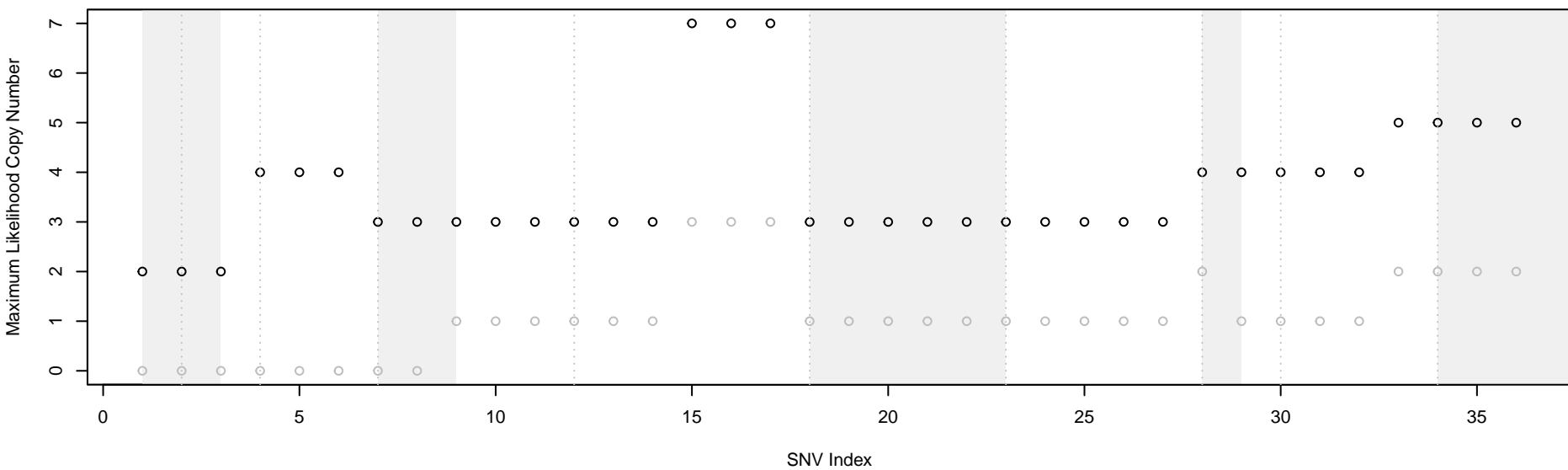
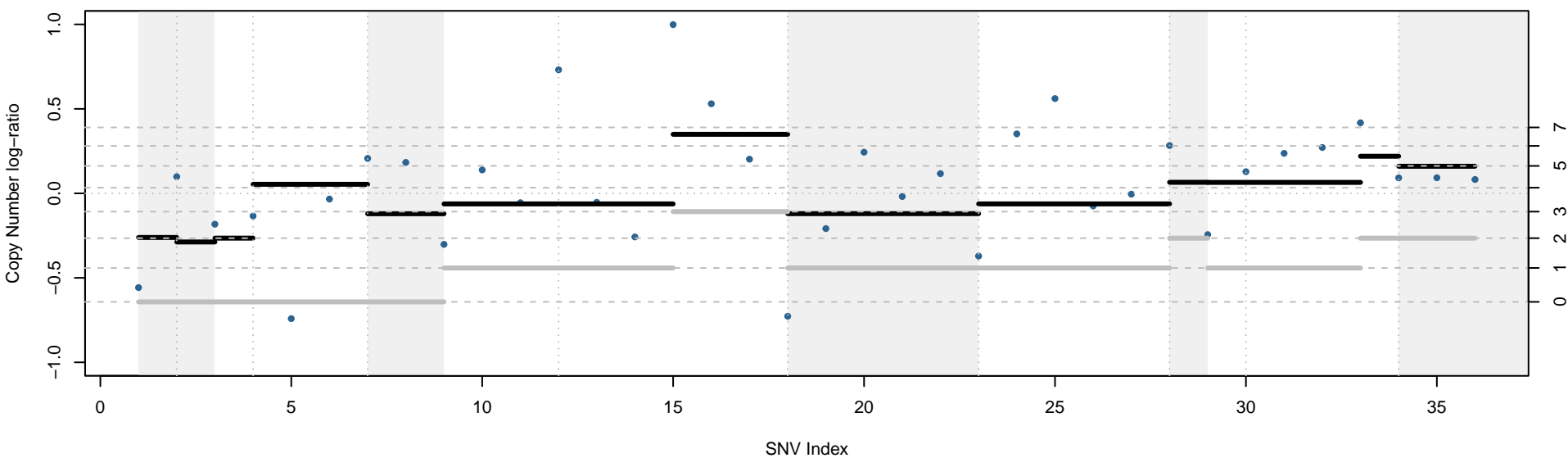


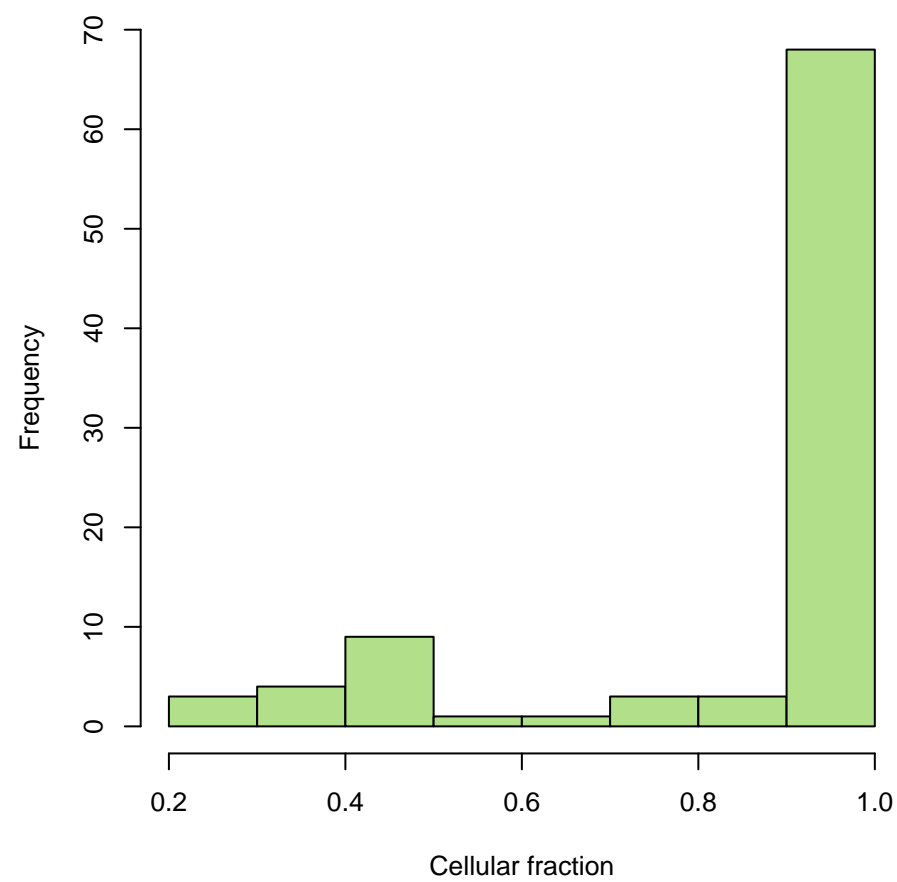
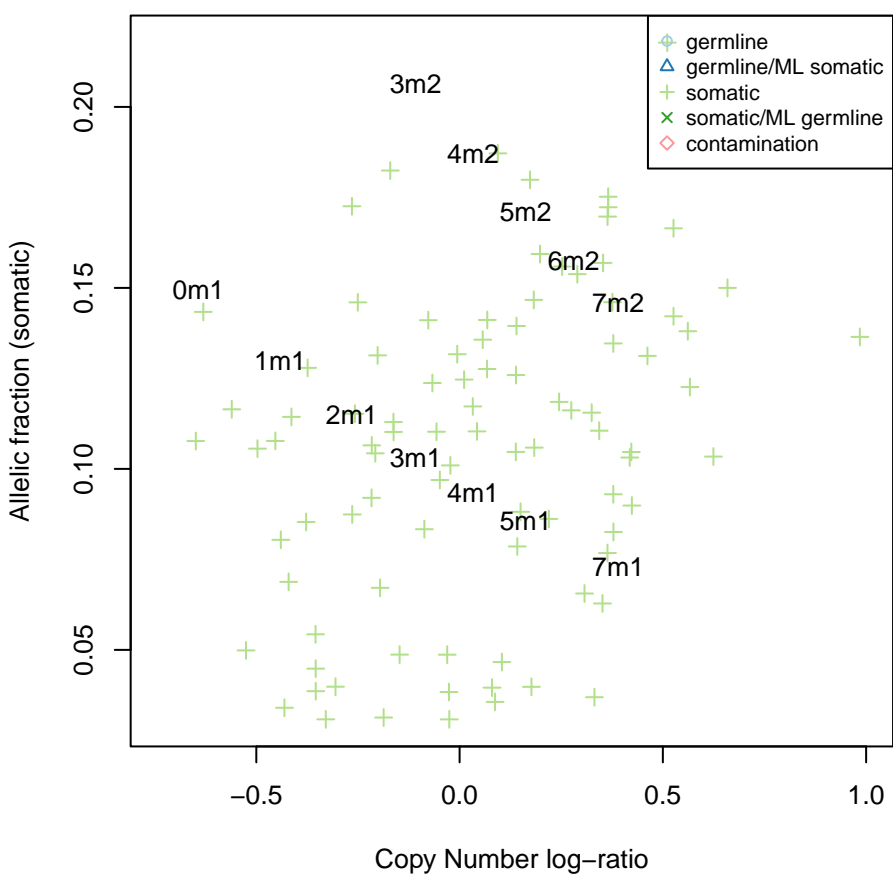
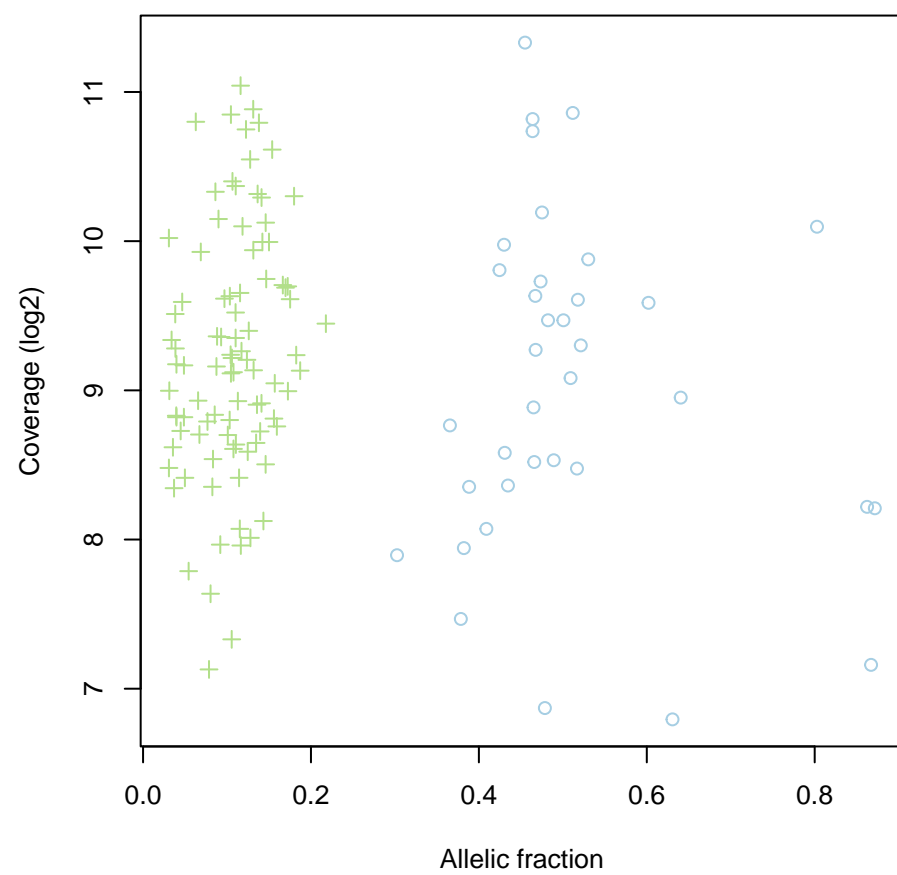
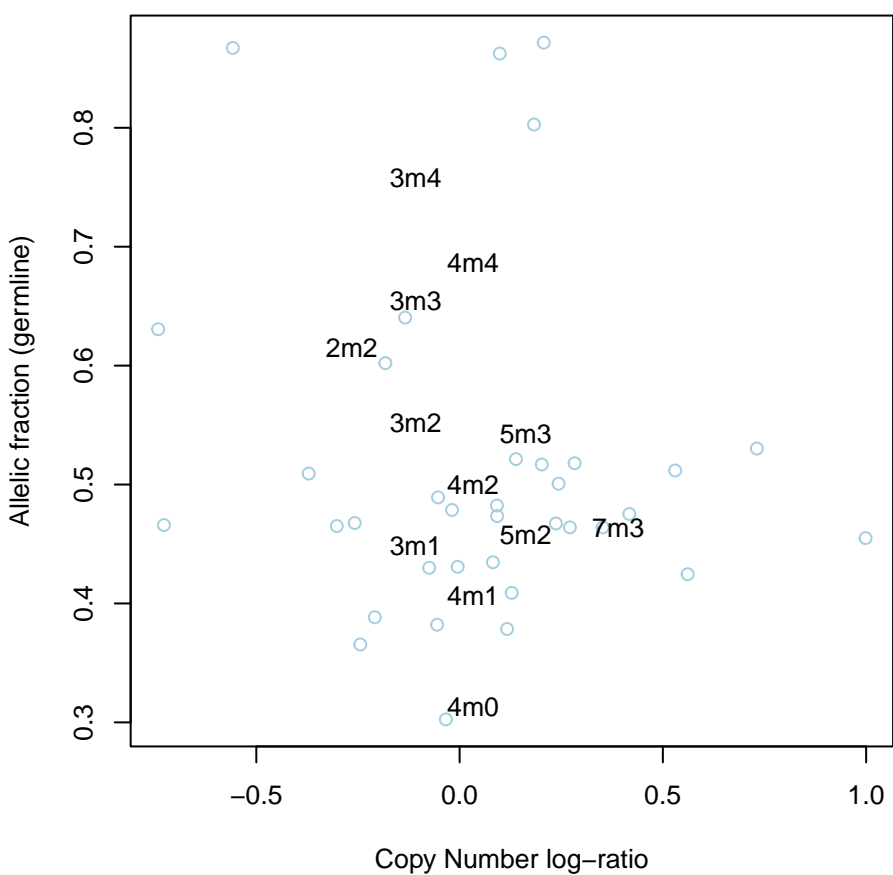


Purity: 0.23 Tumor ploidy: 3.755 SNV log-likelihood: -179.61 GoF: 93.1% Mean coverage: 350;709

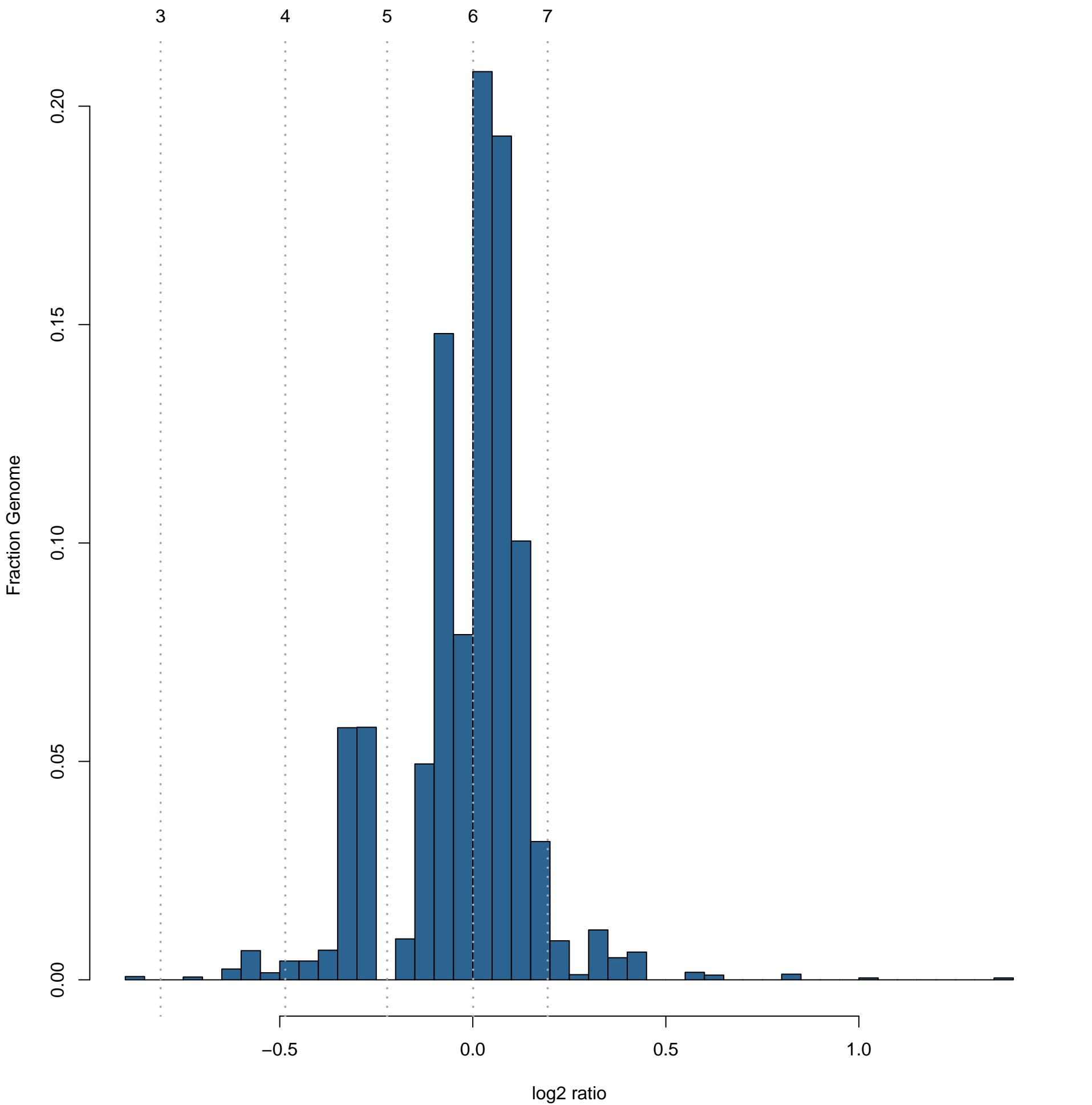


SCNA-fit log-likelihood: -3971.28

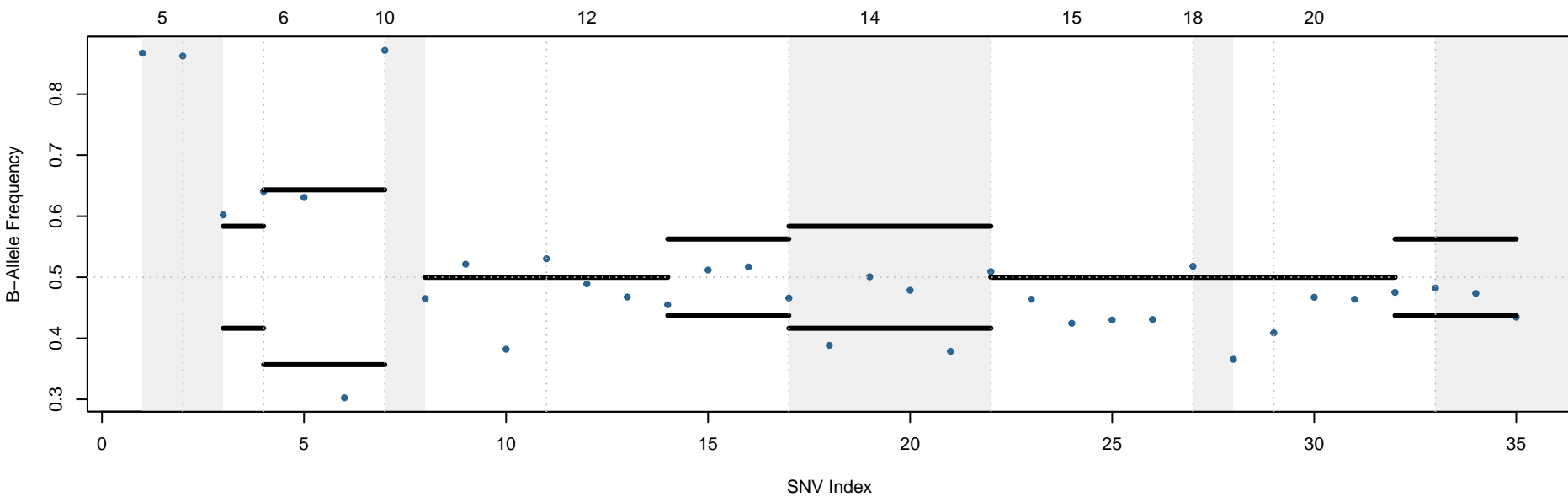




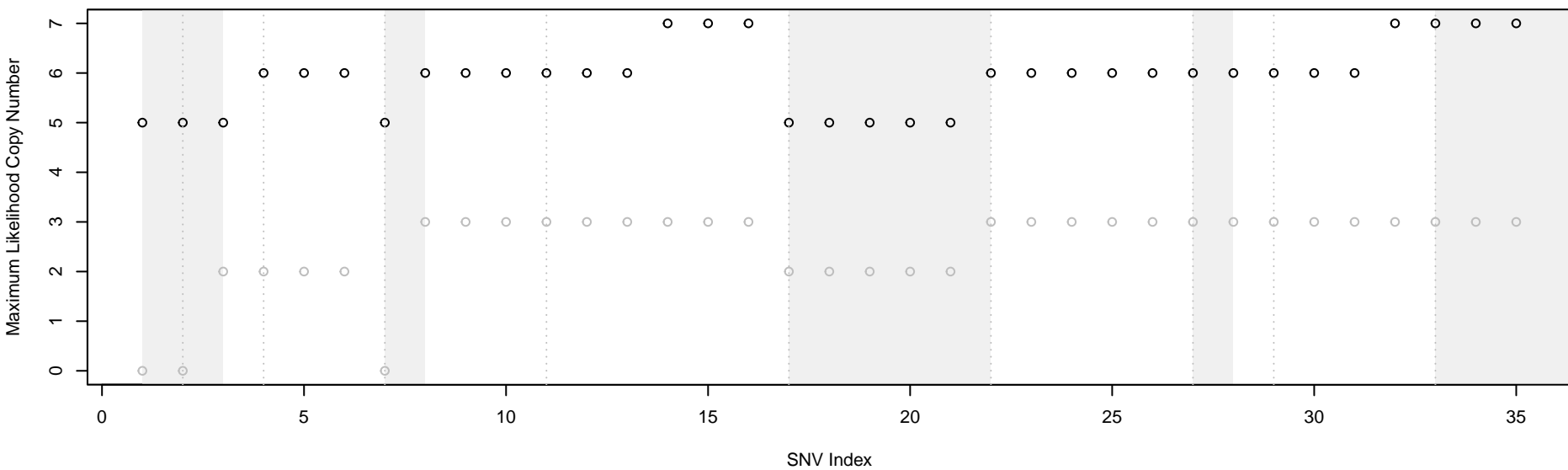
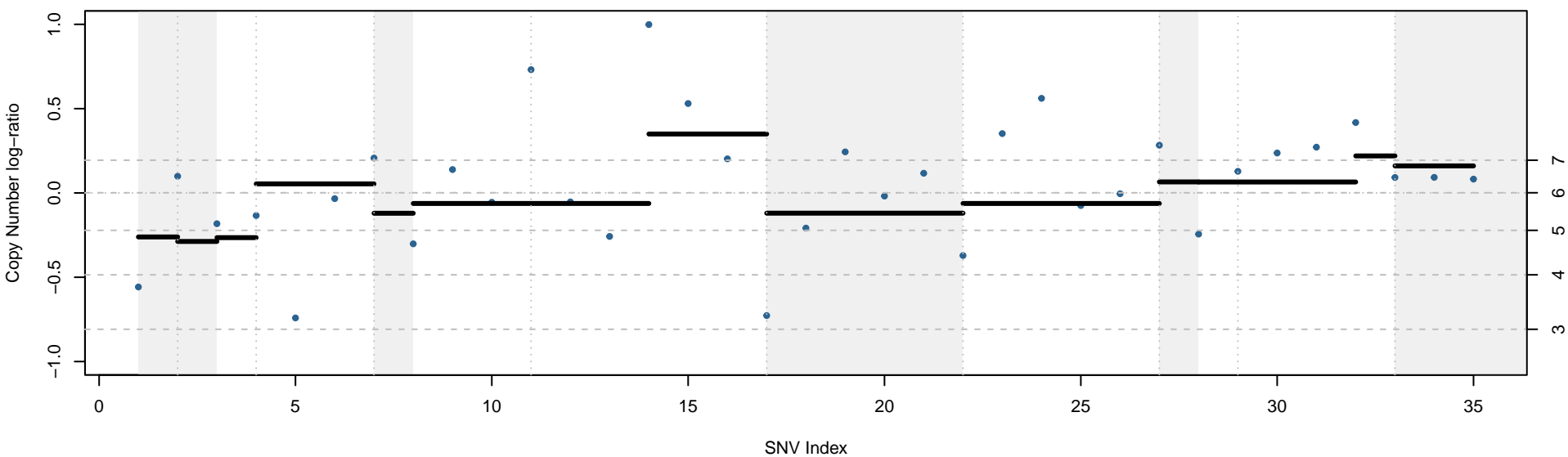
Purity: 0.67 Tumor ploidy: 5.995

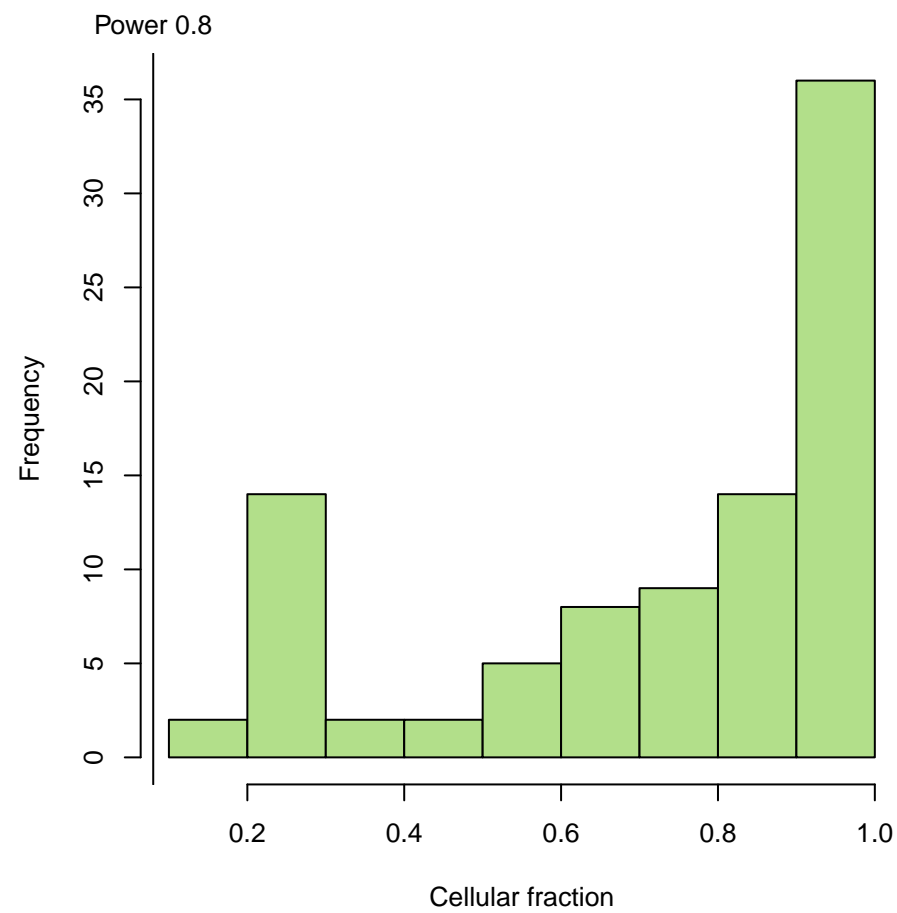
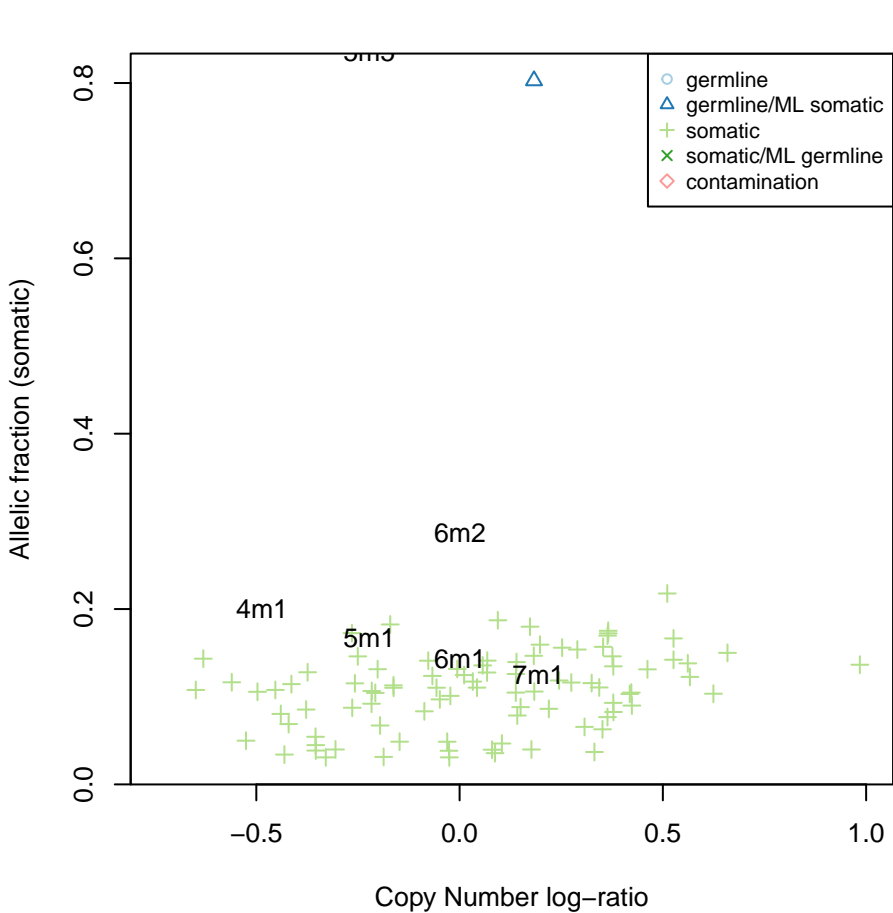
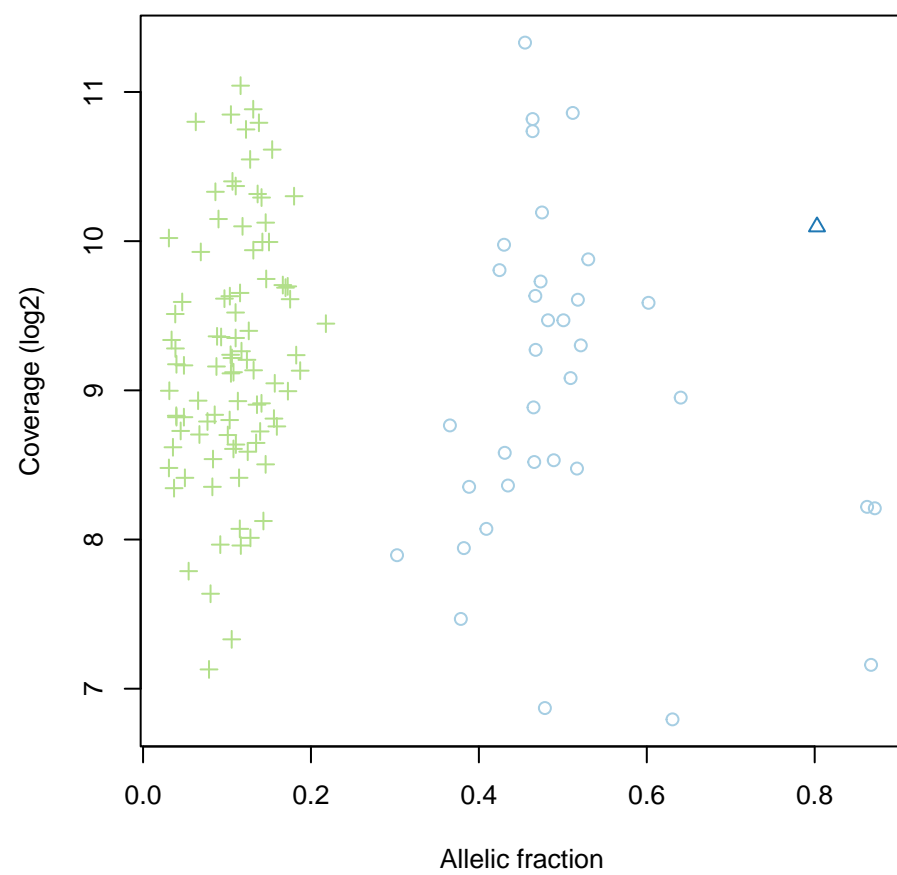
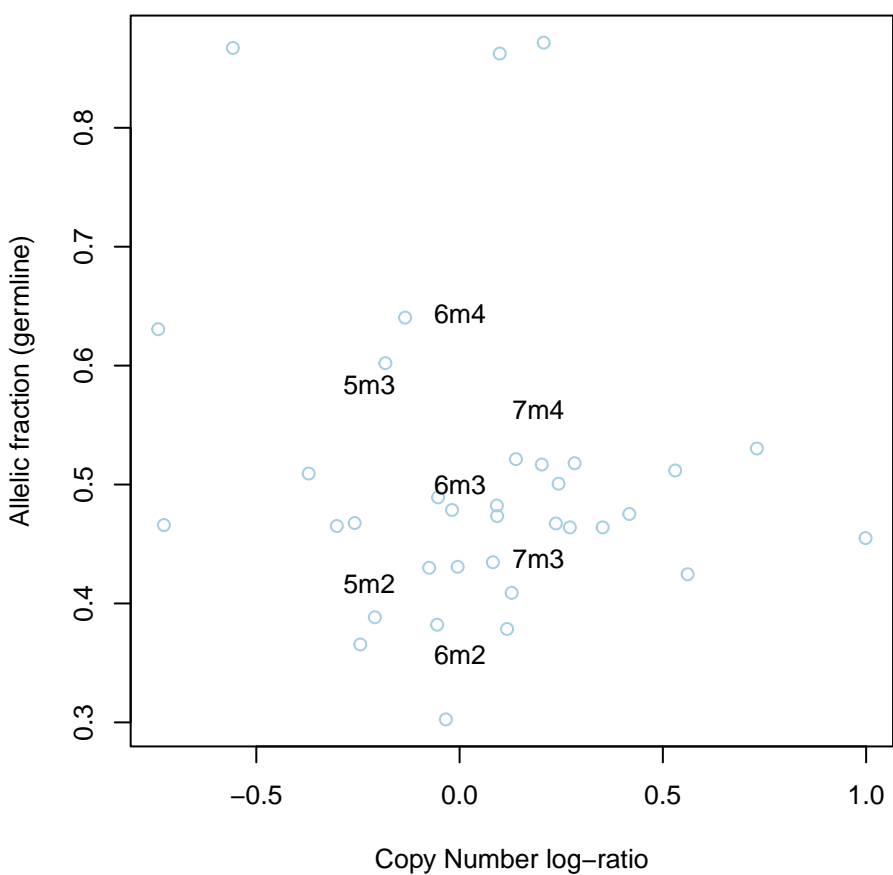


Purity: 0.67 Tumor ploidy: 5.995 SNV log-likelihood: -146.32 GoF: 92.2% Mean coverage: 350,709

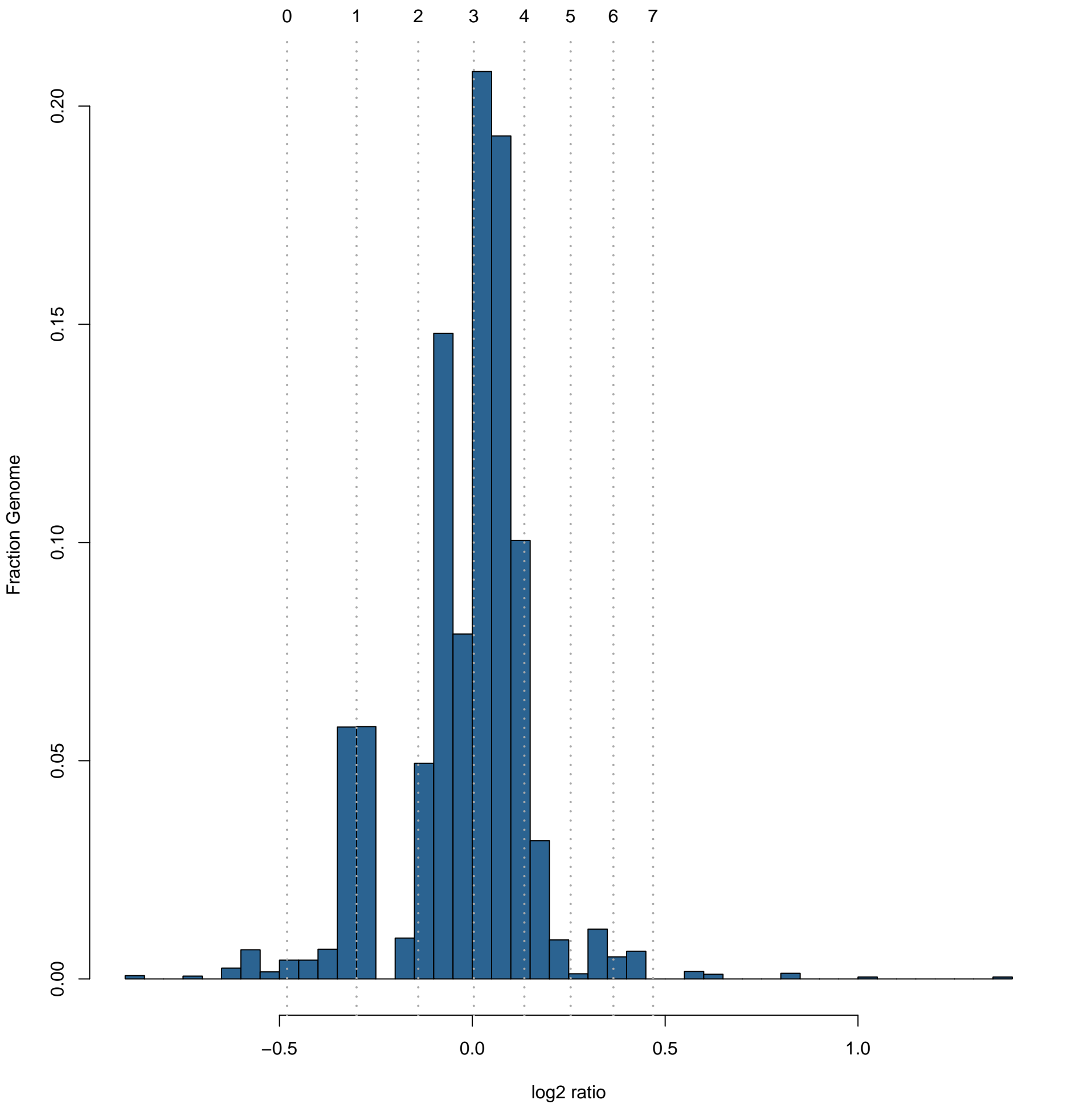


SCNA-fit log-likelihood: -4075.47

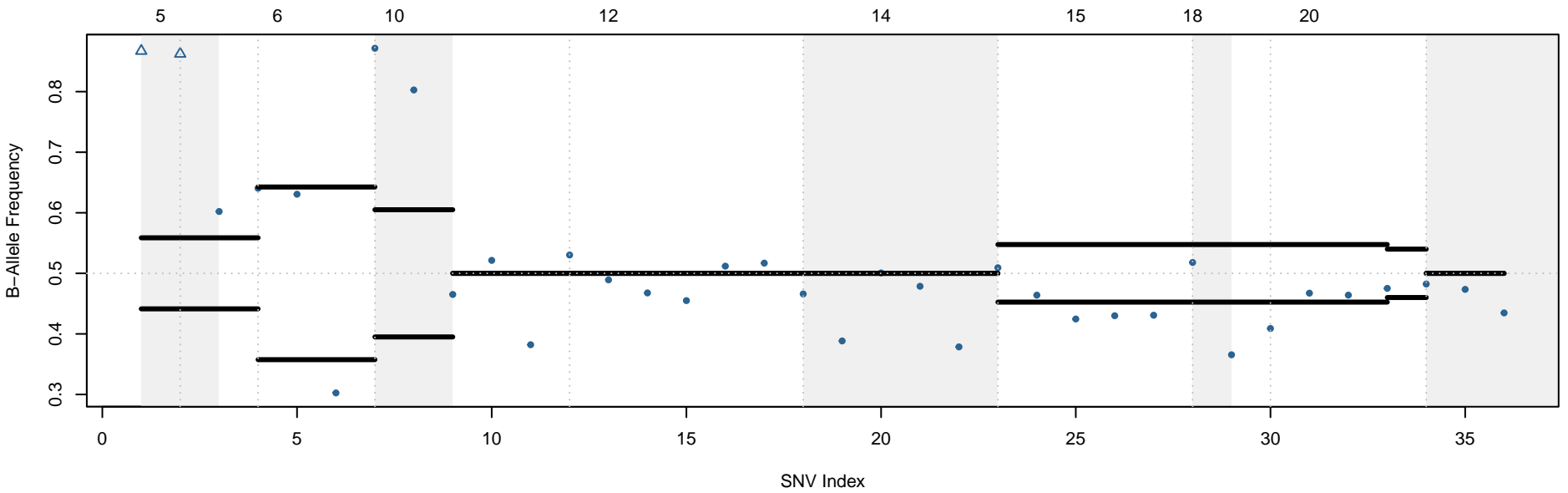




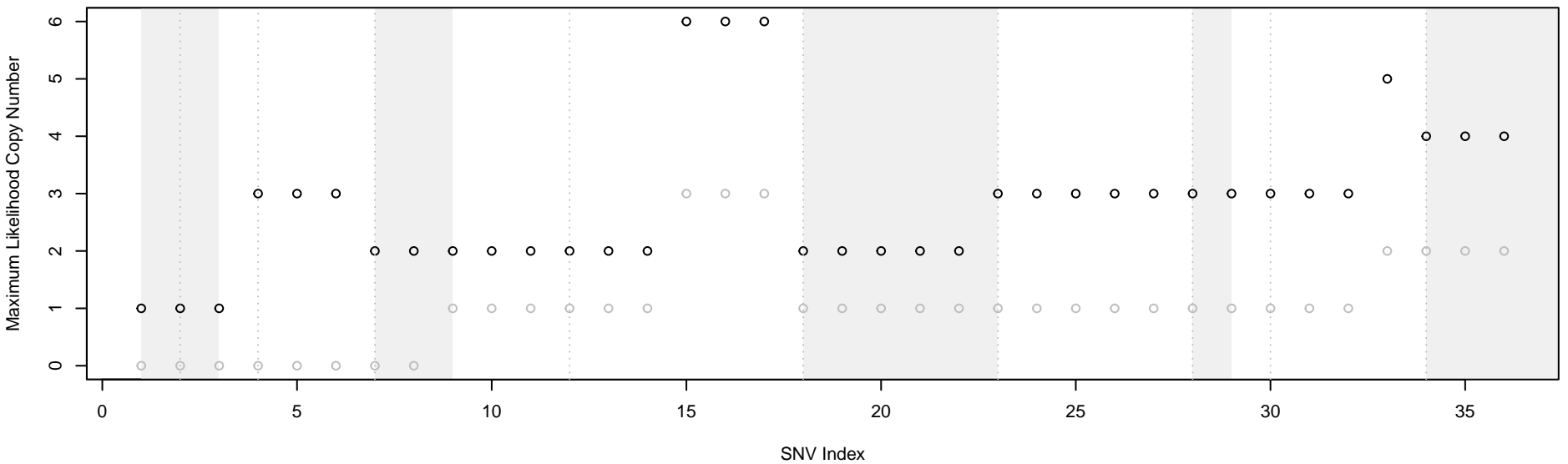
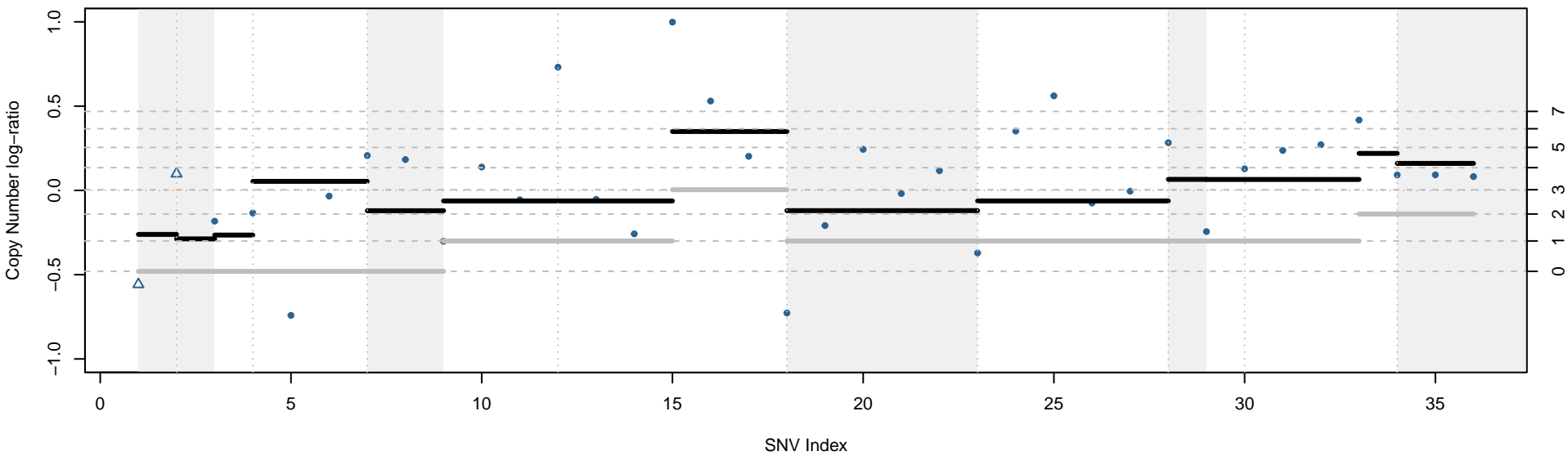
Purity: 0.21 Tumor ploidy: 2.971

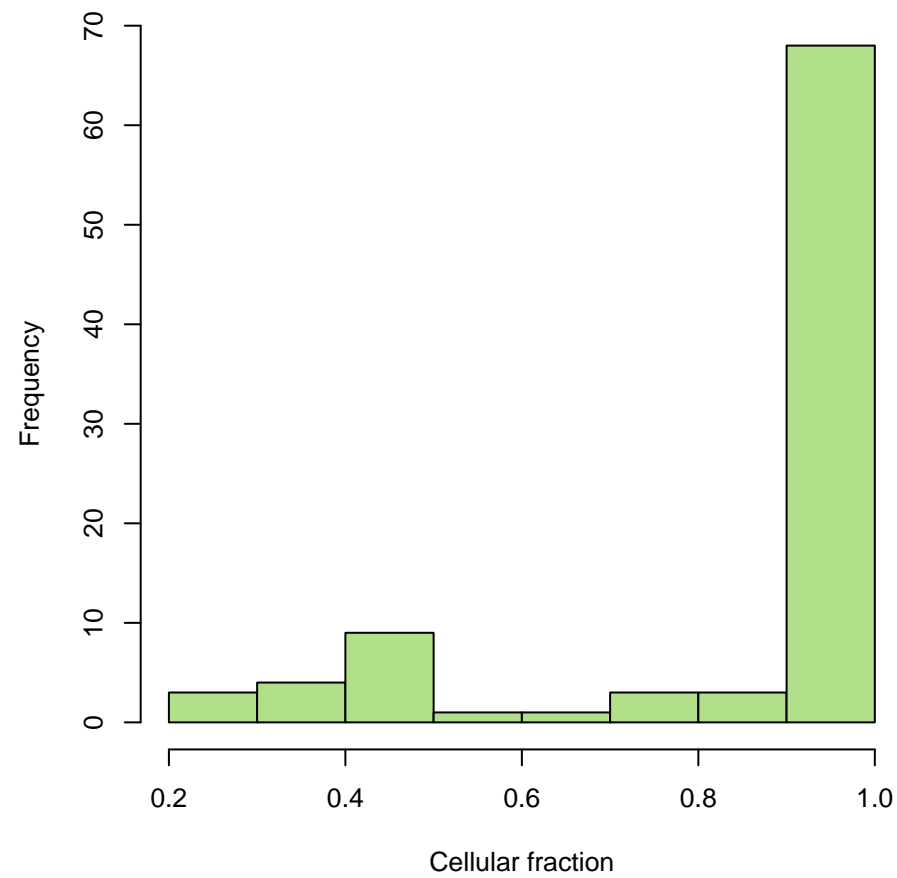
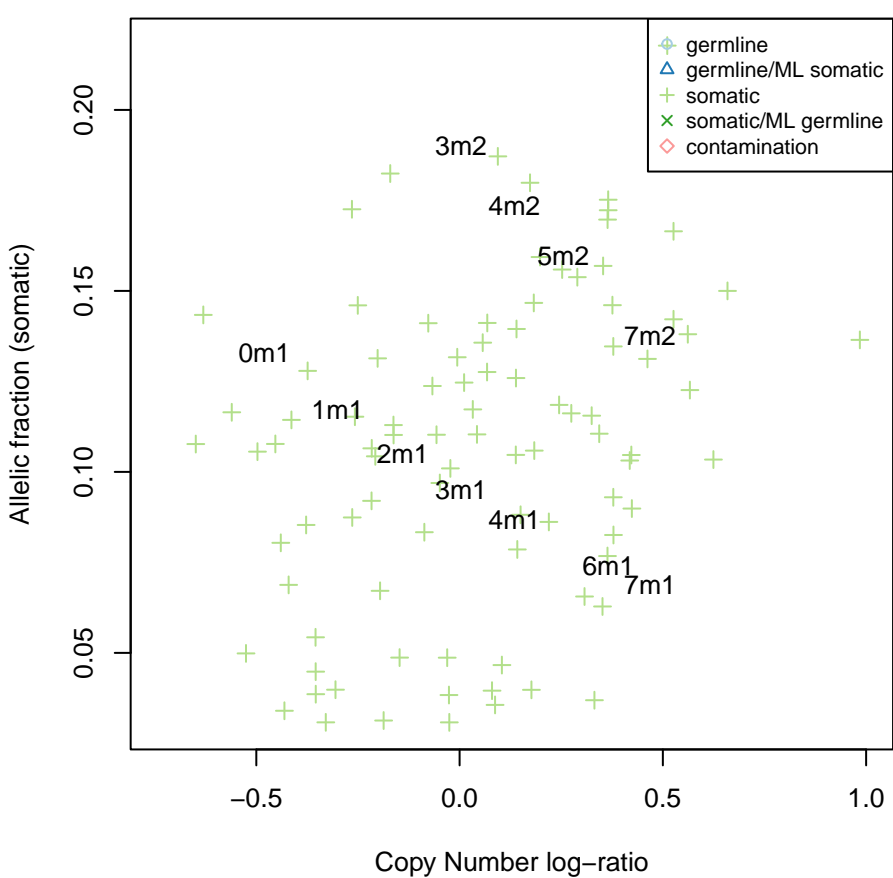
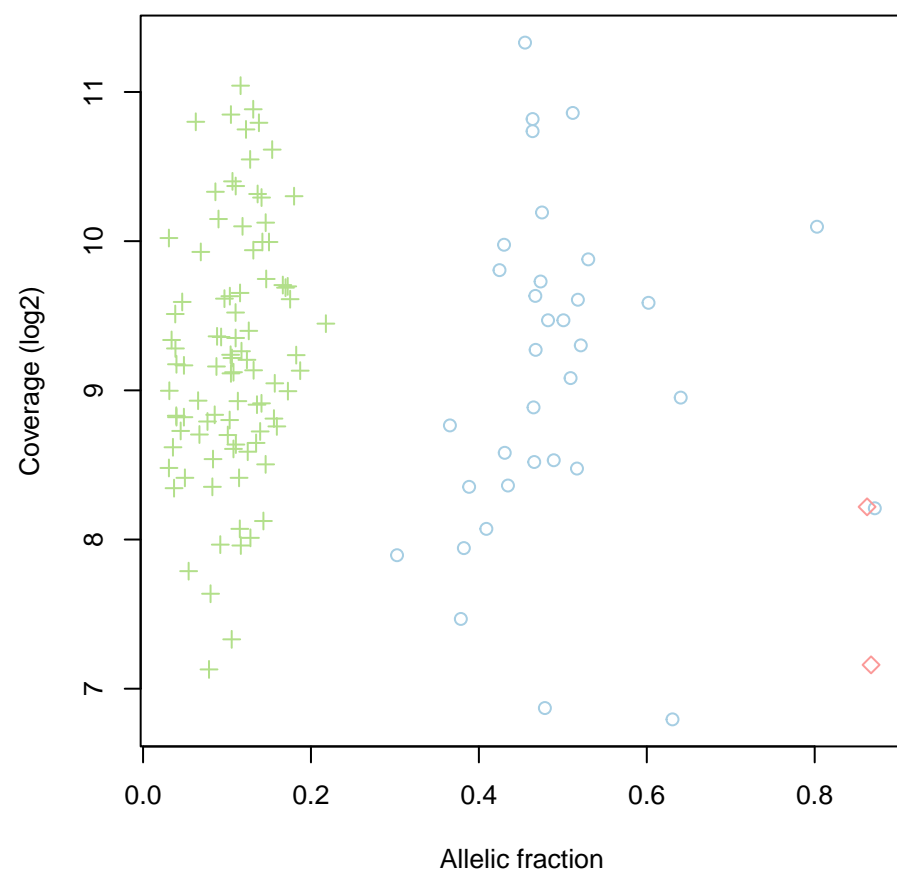
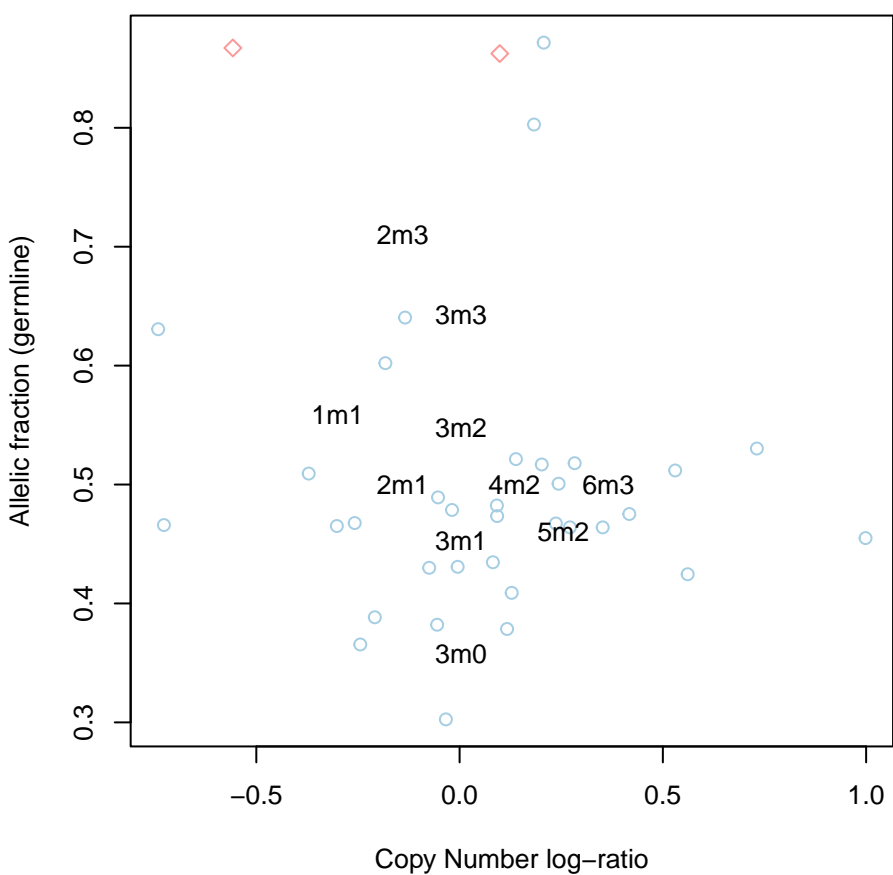


Purity: 0.21 Tumor ploidy: 2.971 SNV log-likelihood: -180.78 GoF: 95.5% Mean coverage: 350;709



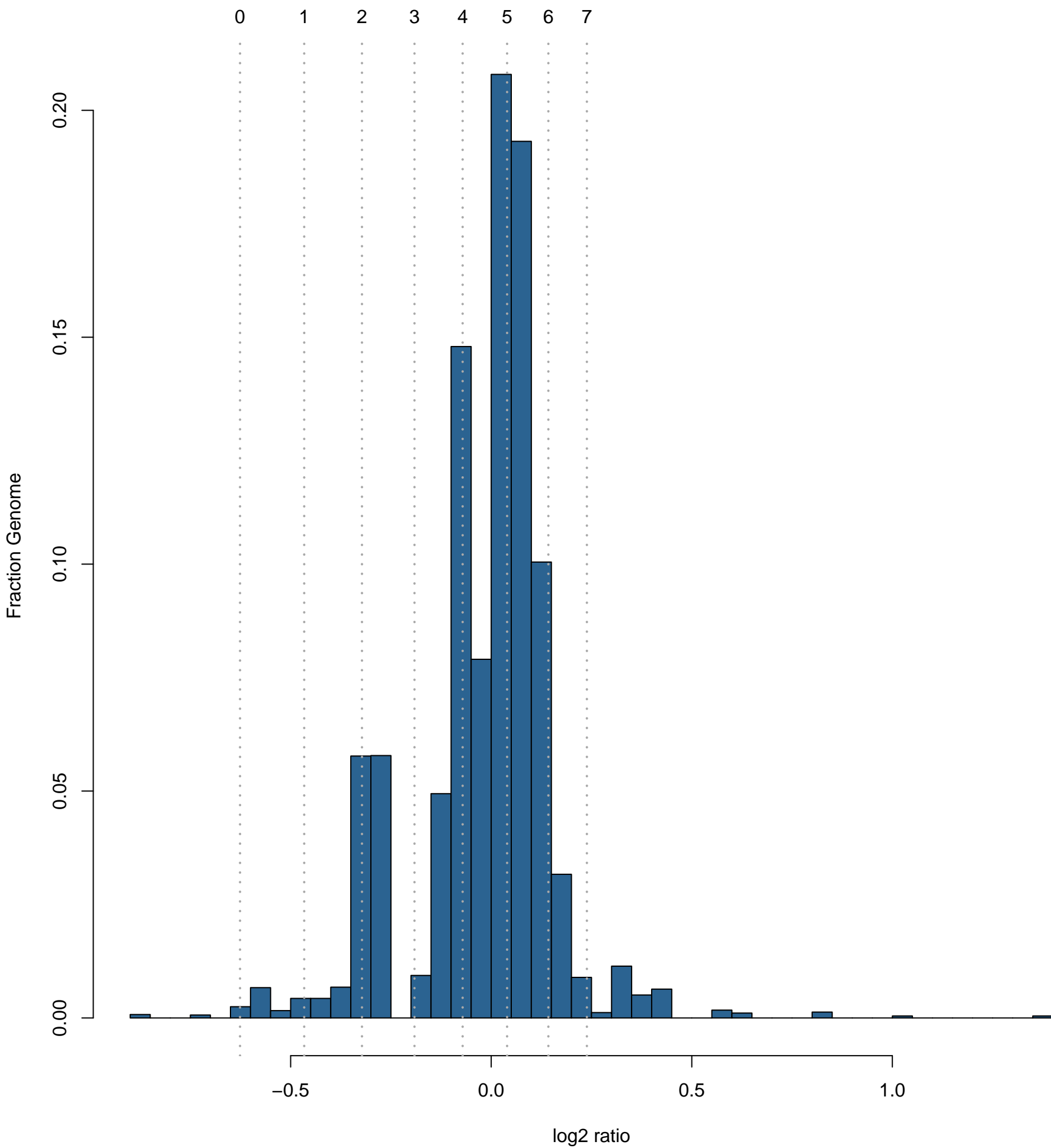
SCNA-fit log-likelihood: -3973.19



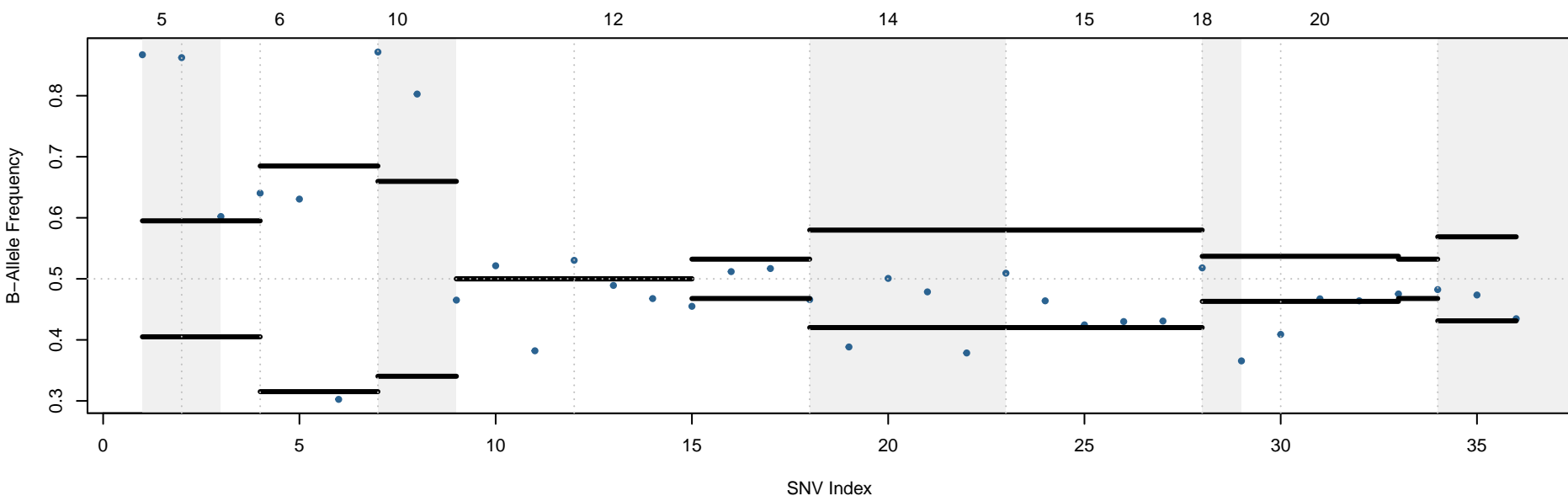




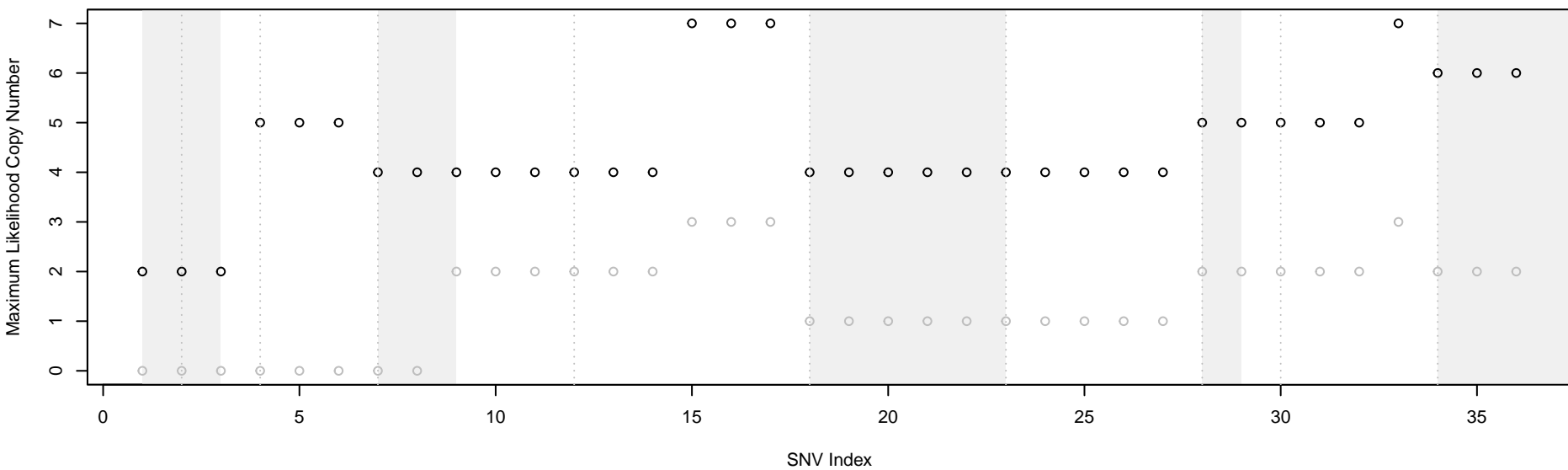
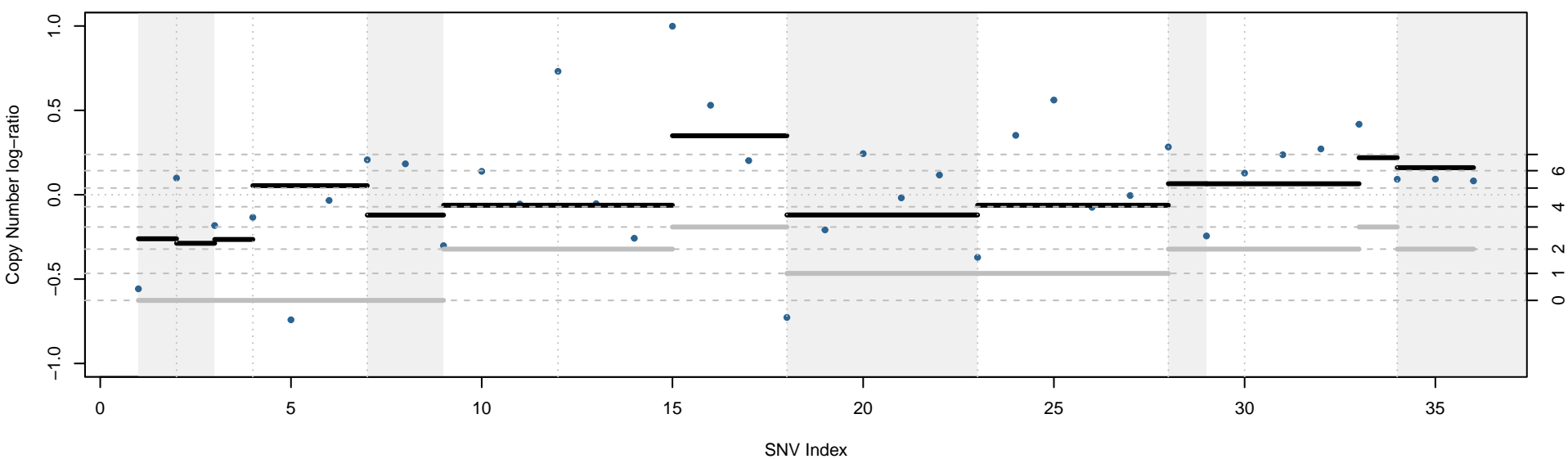
Purity: 0.19 Tumor ploidy: 4.634

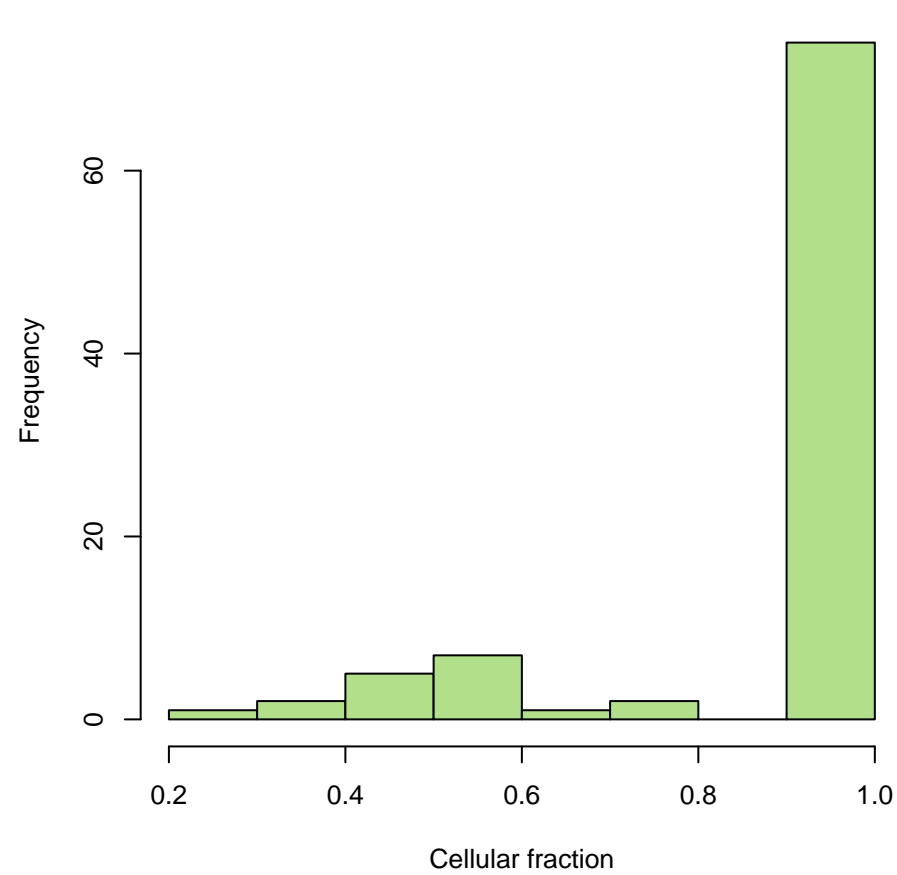
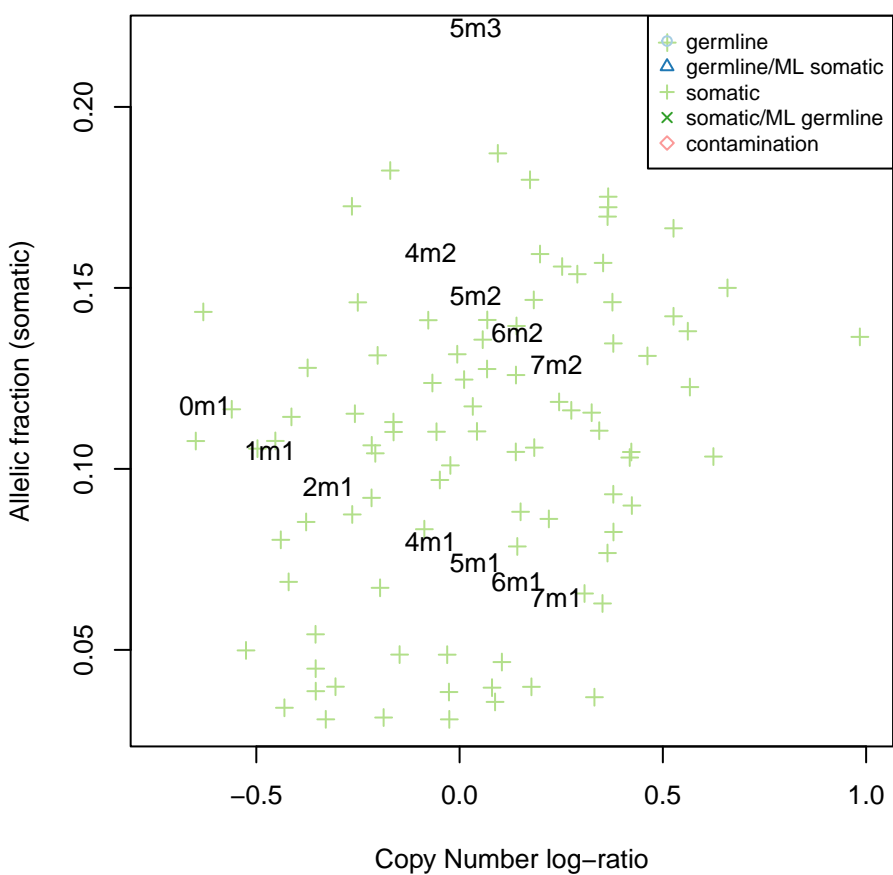
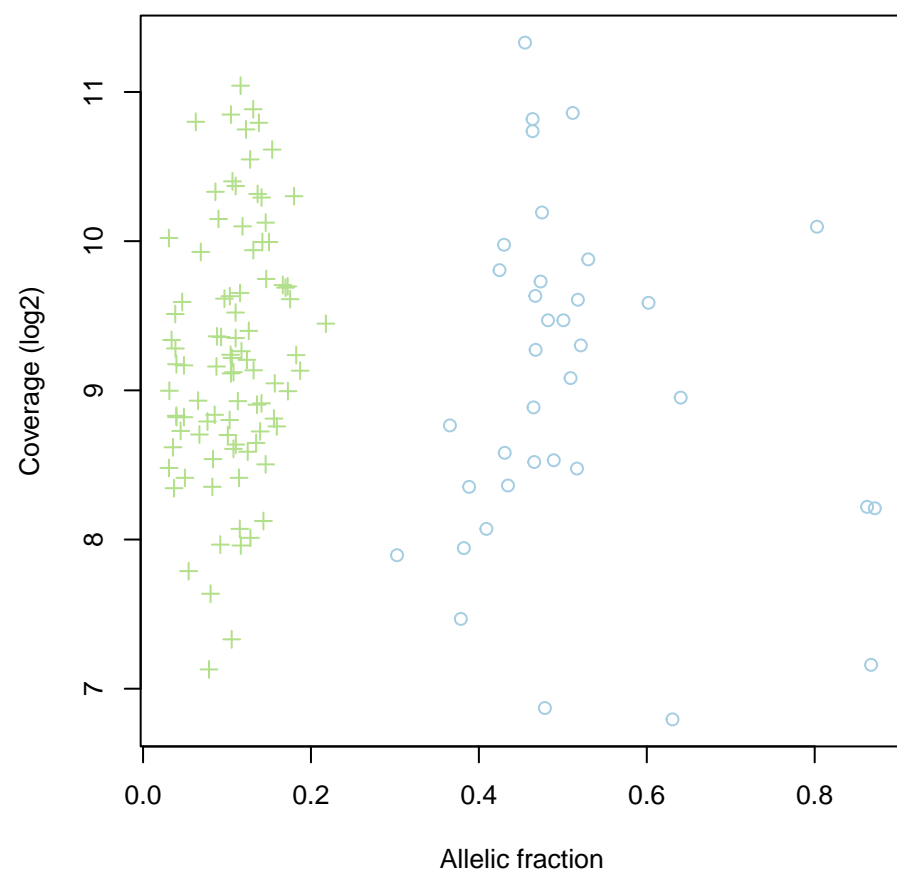
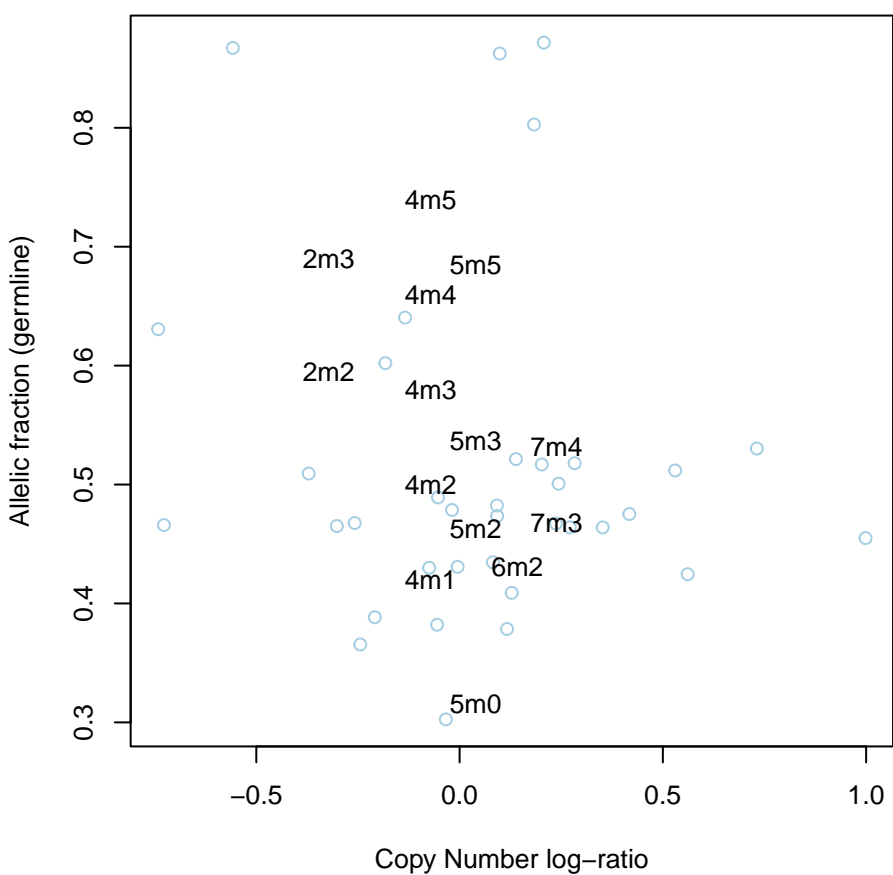


Purity: 0.19 Tumor ploidy: 4.634 SNV log-likelihood: -212.31 GoF: 94.4% Mean coverage: 350;709

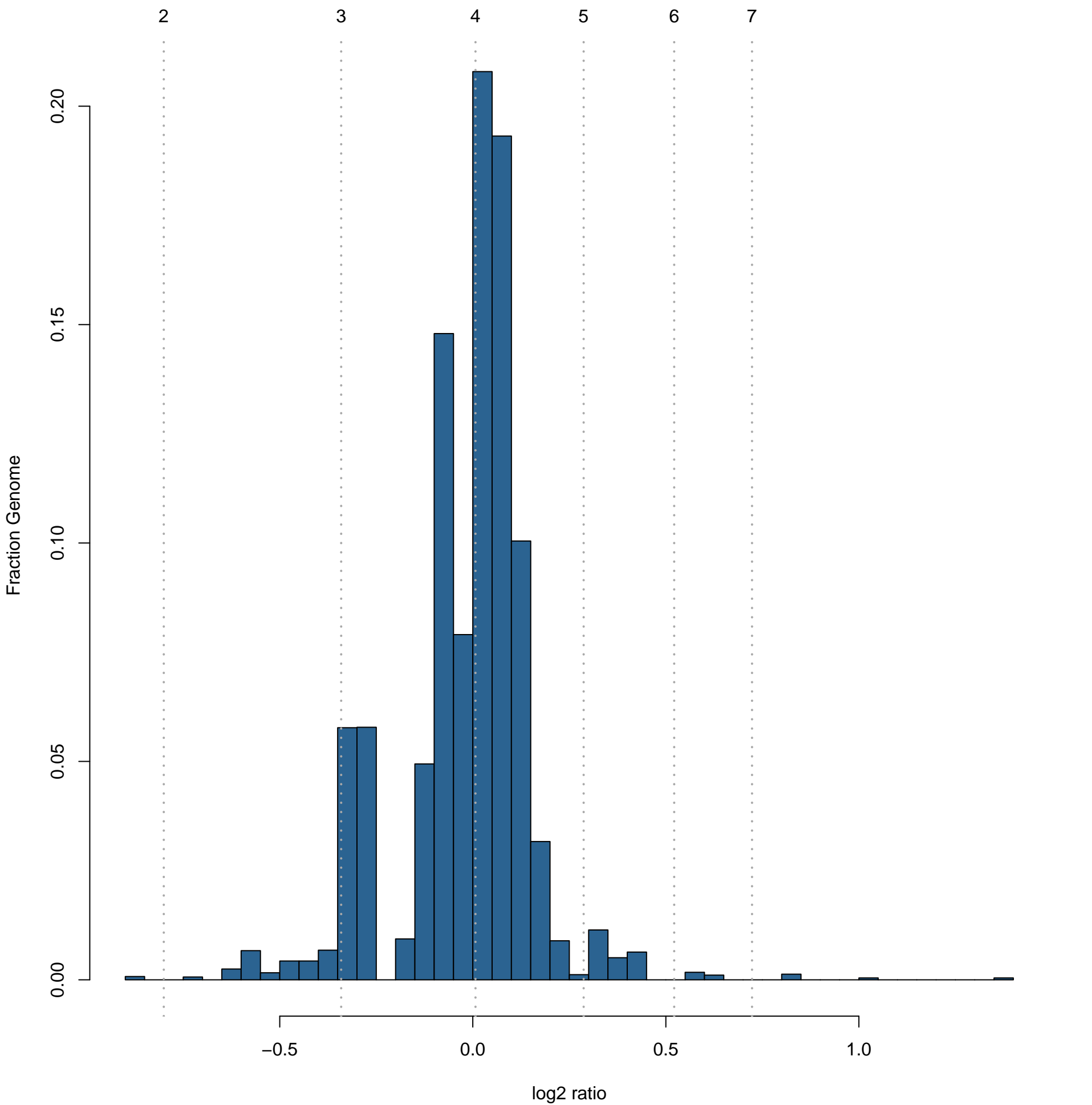


SCNA-fit log-likelihood: -4026.24

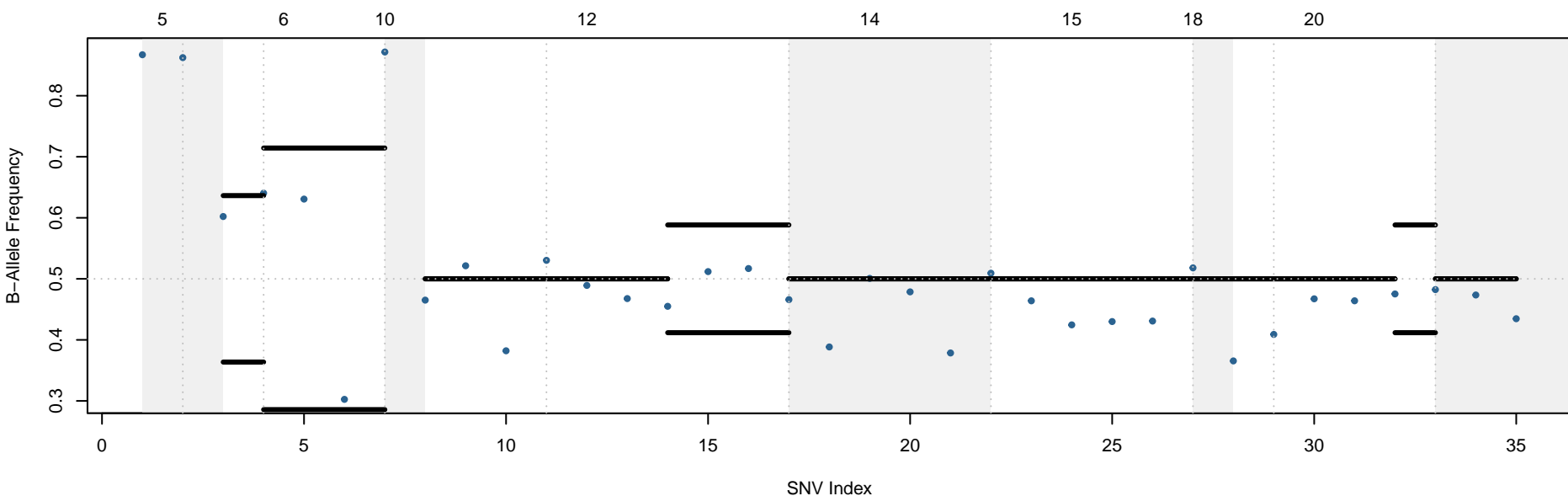




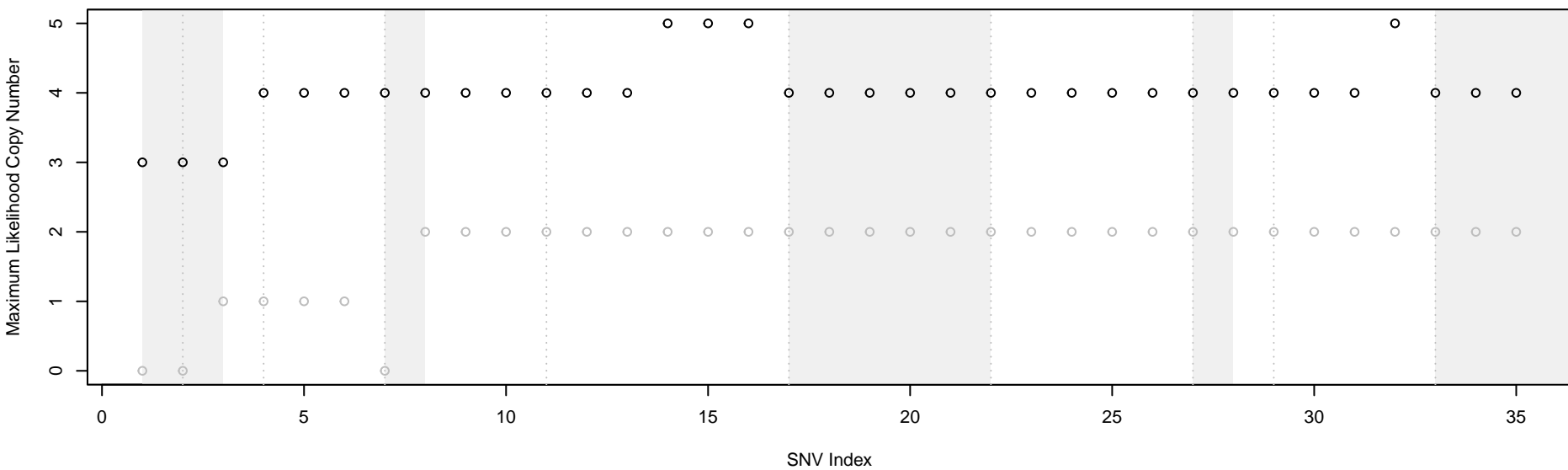
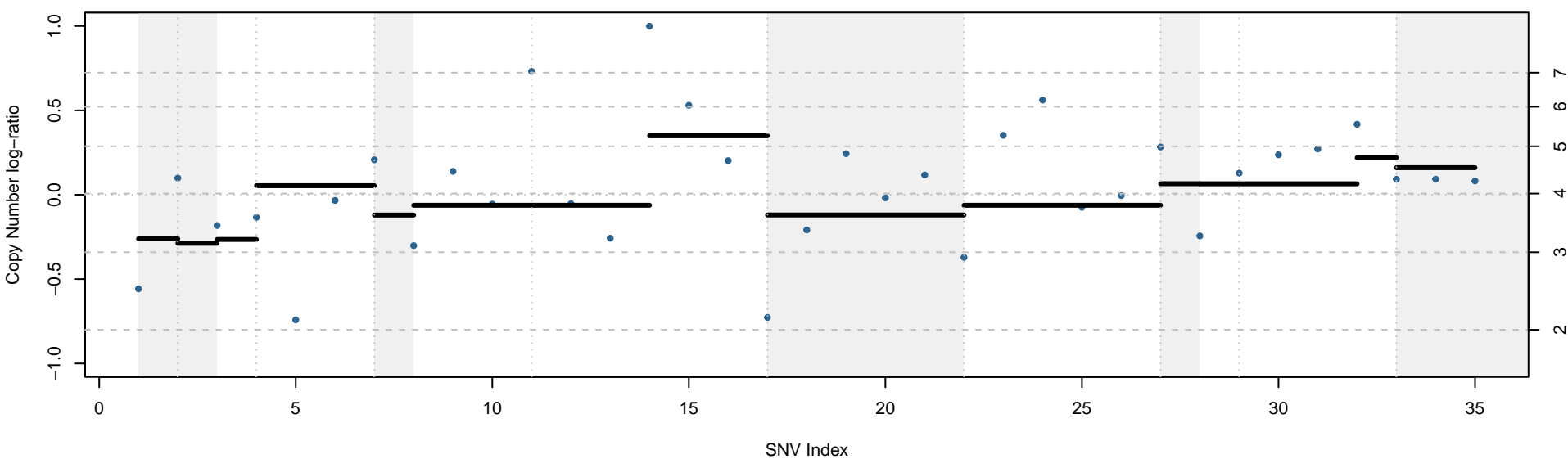
Purity: 0.75 Tumor ploidy: 3.977

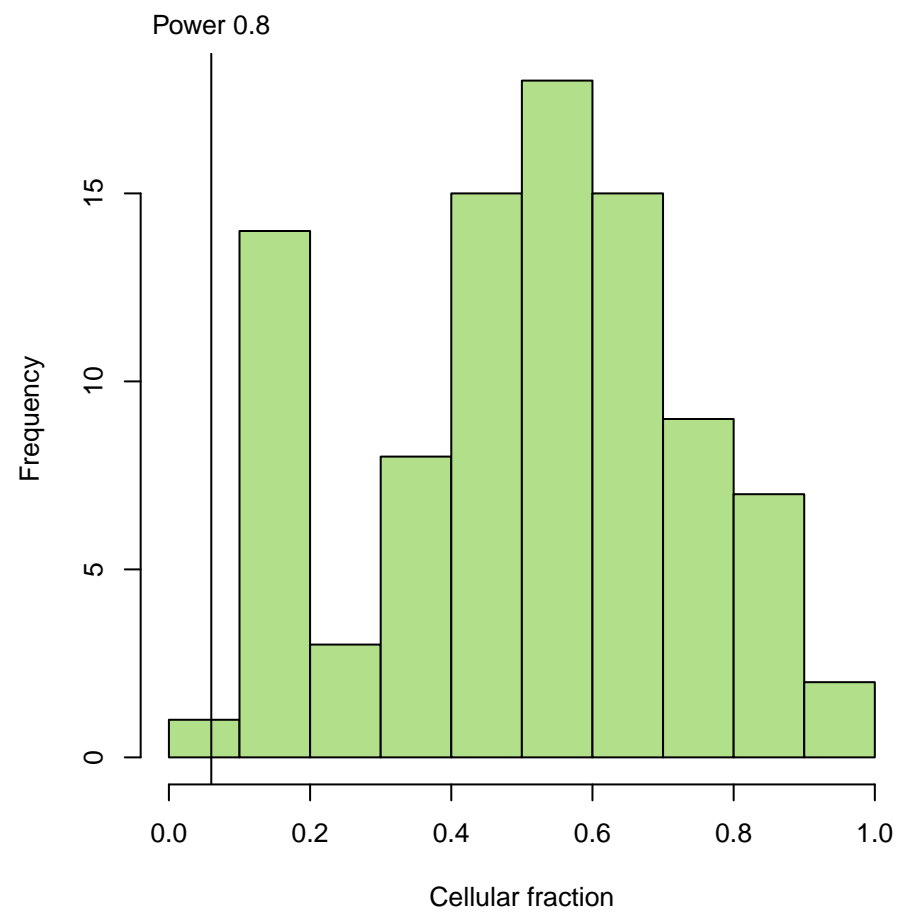
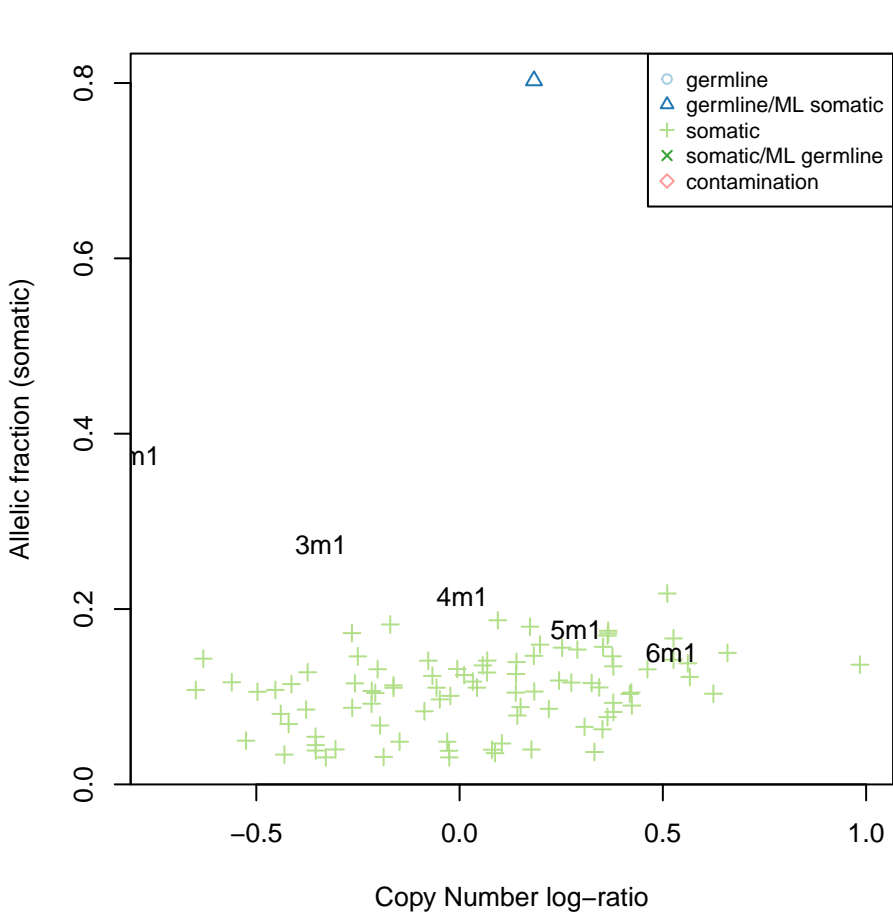
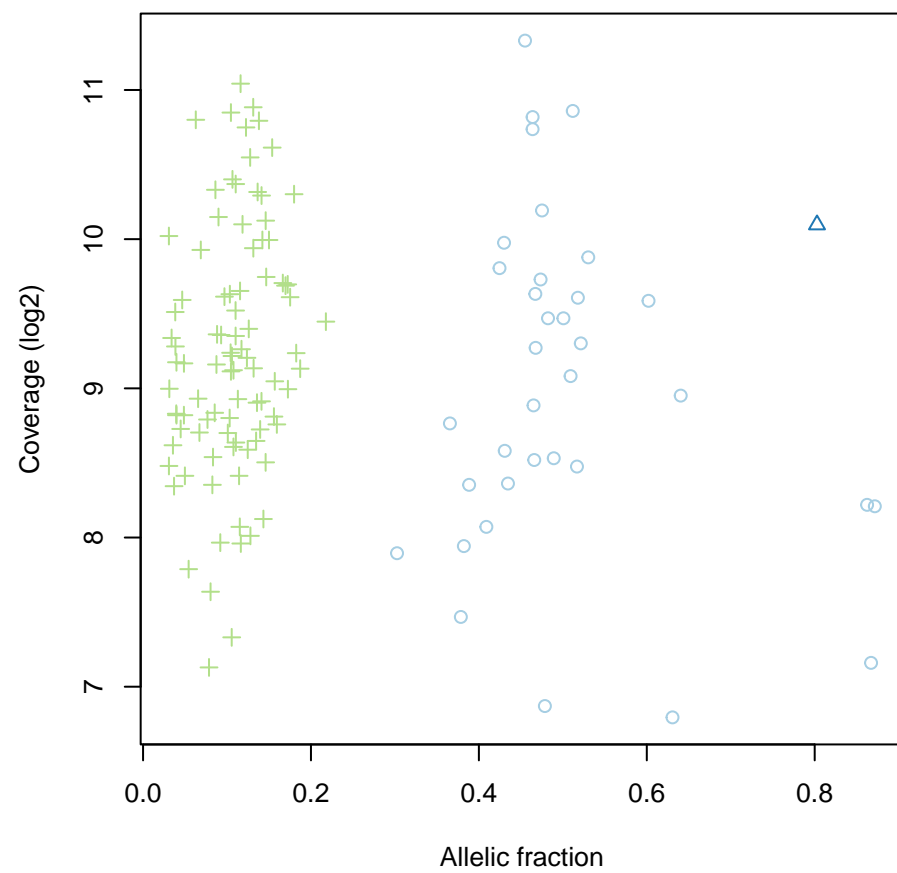
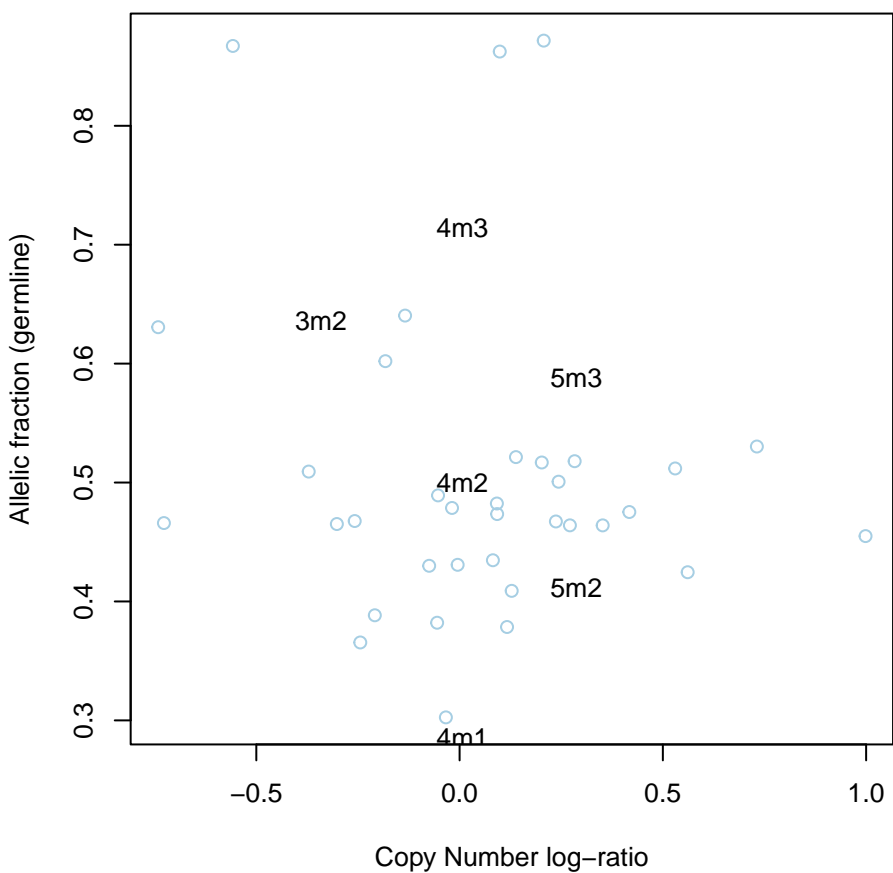


Purity: 0.75 Tumor ploidy: 3.977 SNV log-likelihood: -272.75 GoF: 70.2% Mean coverage: 350;709

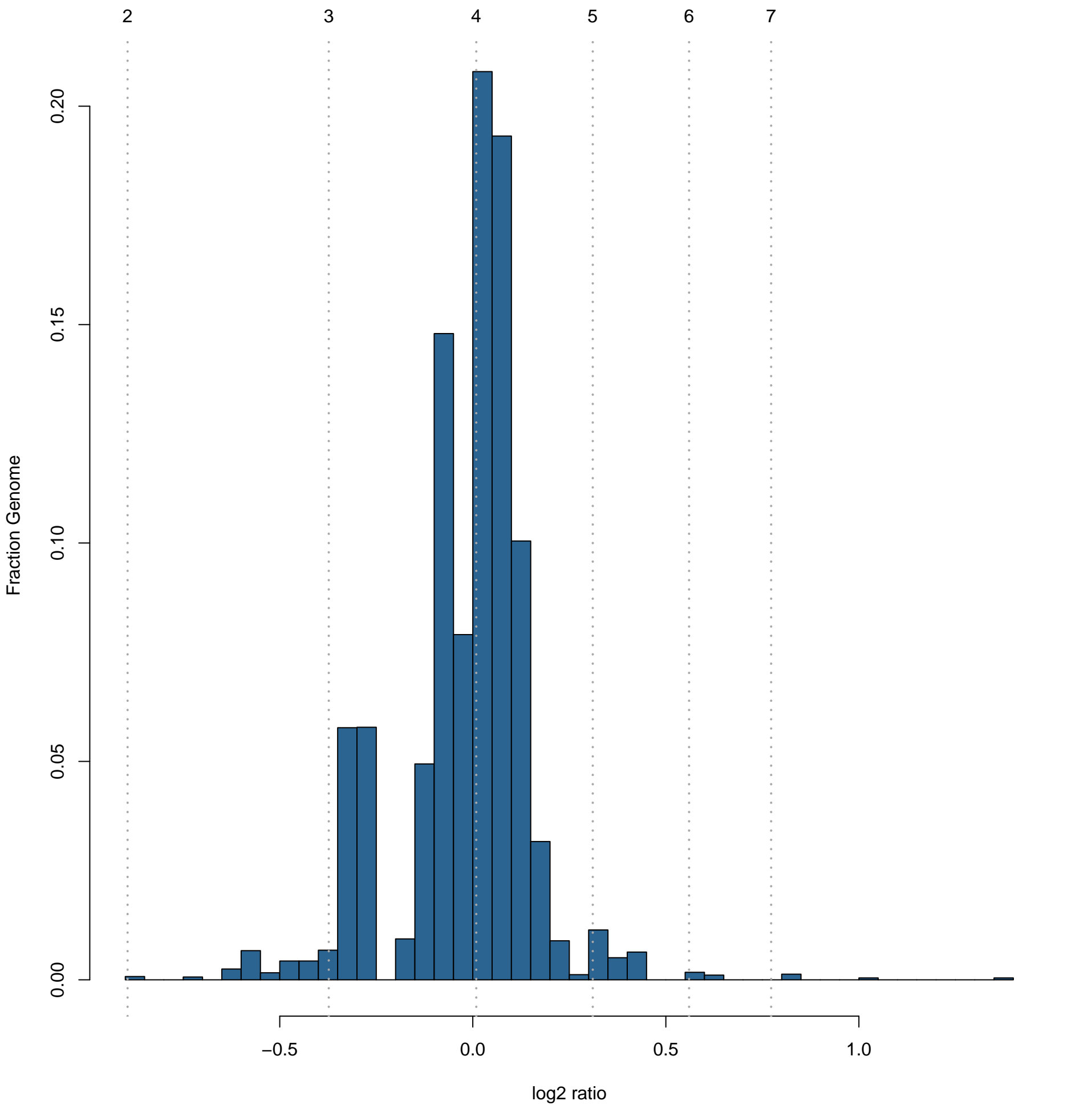


SCNA-fit log-likelihood: -4063.09

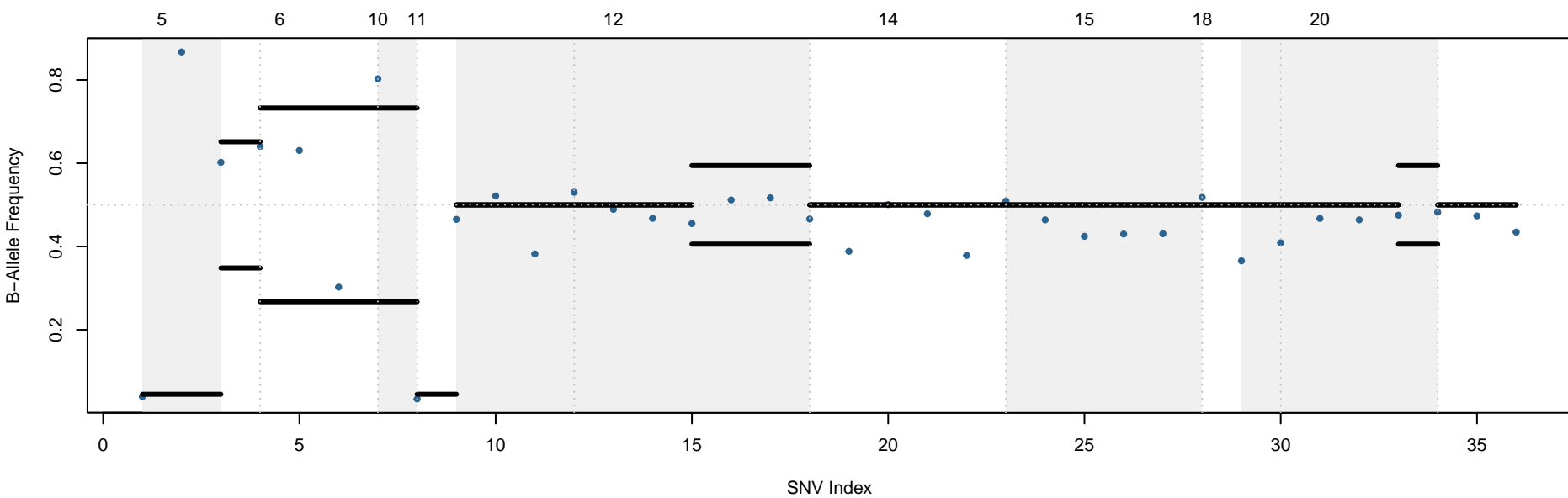




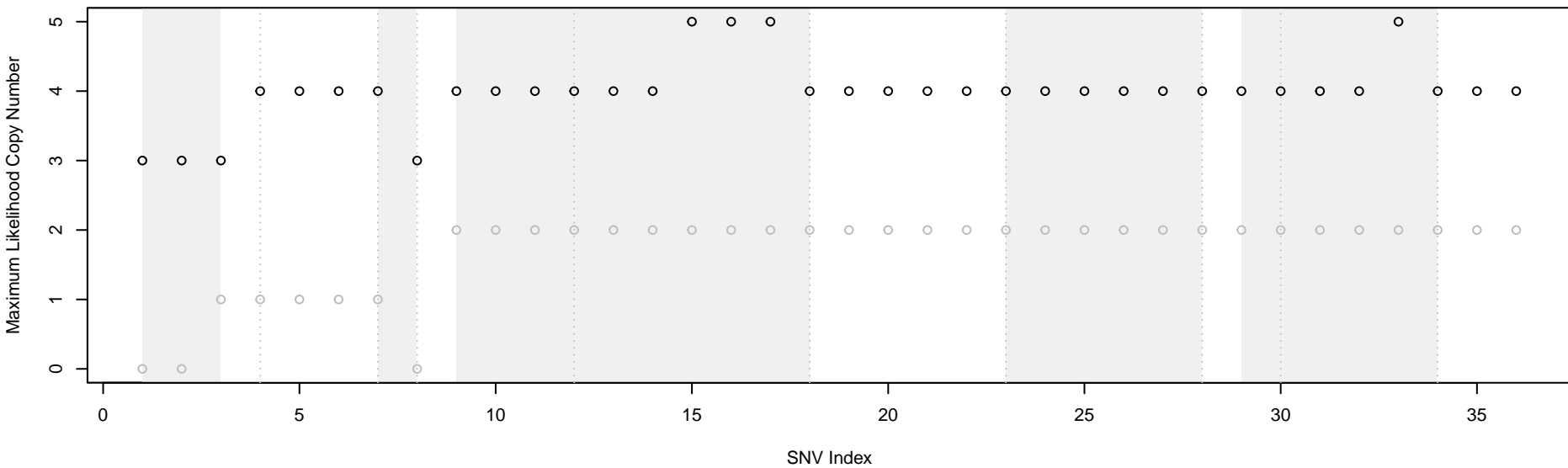
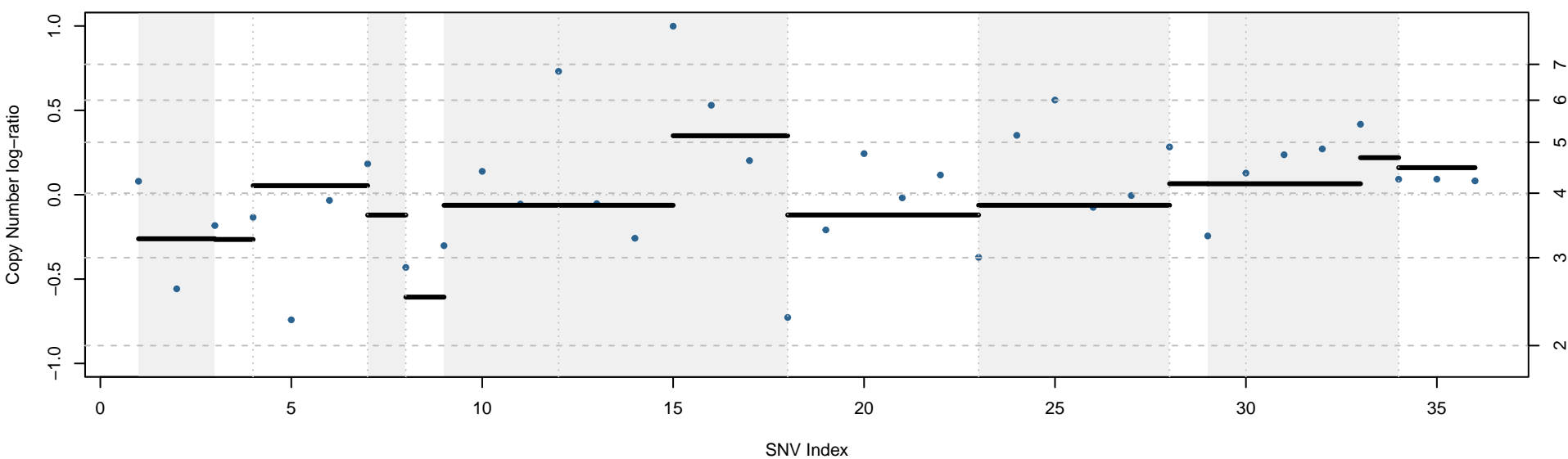
Purity: 0.87 Tumor ploidy: 3.973



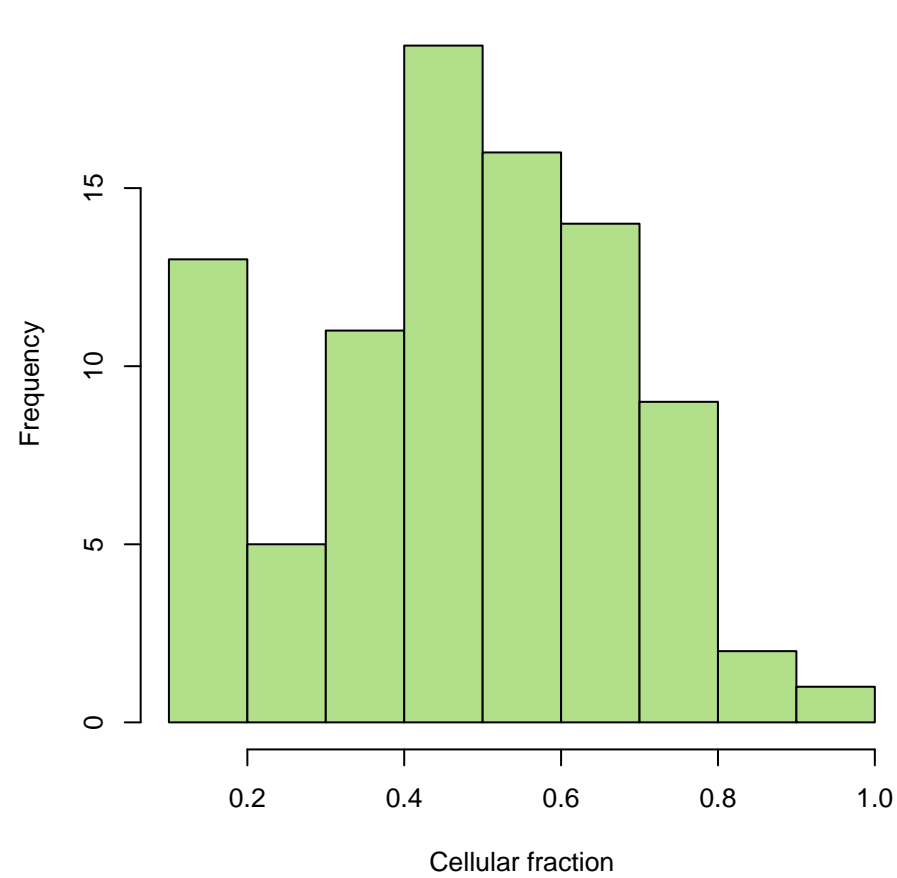
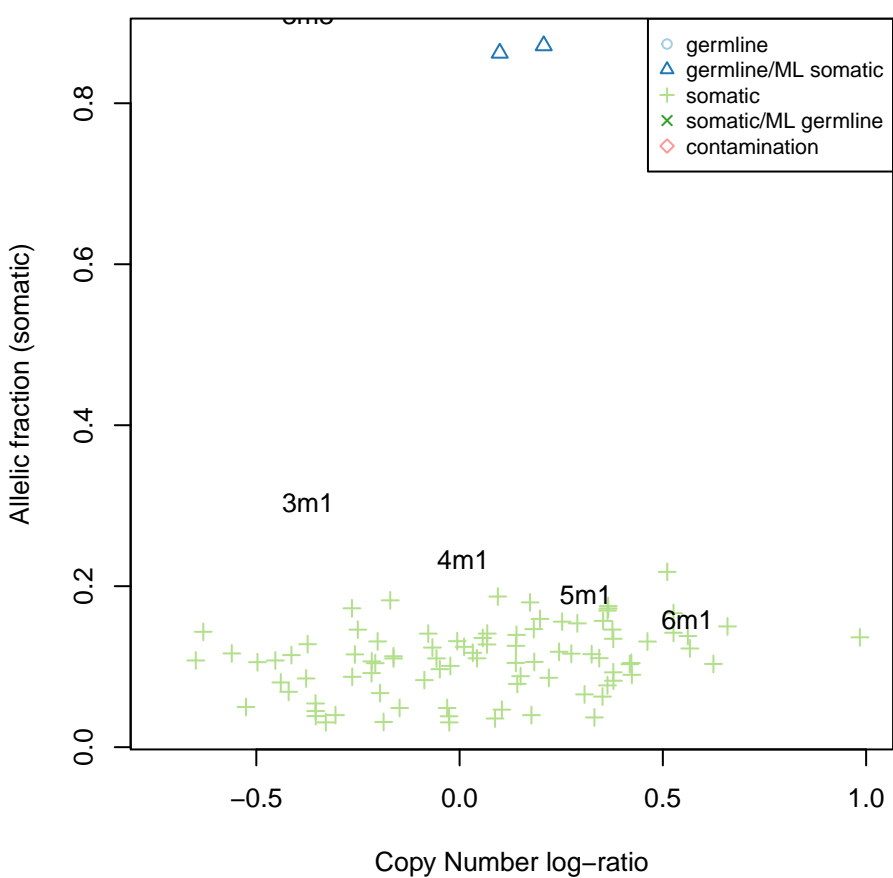
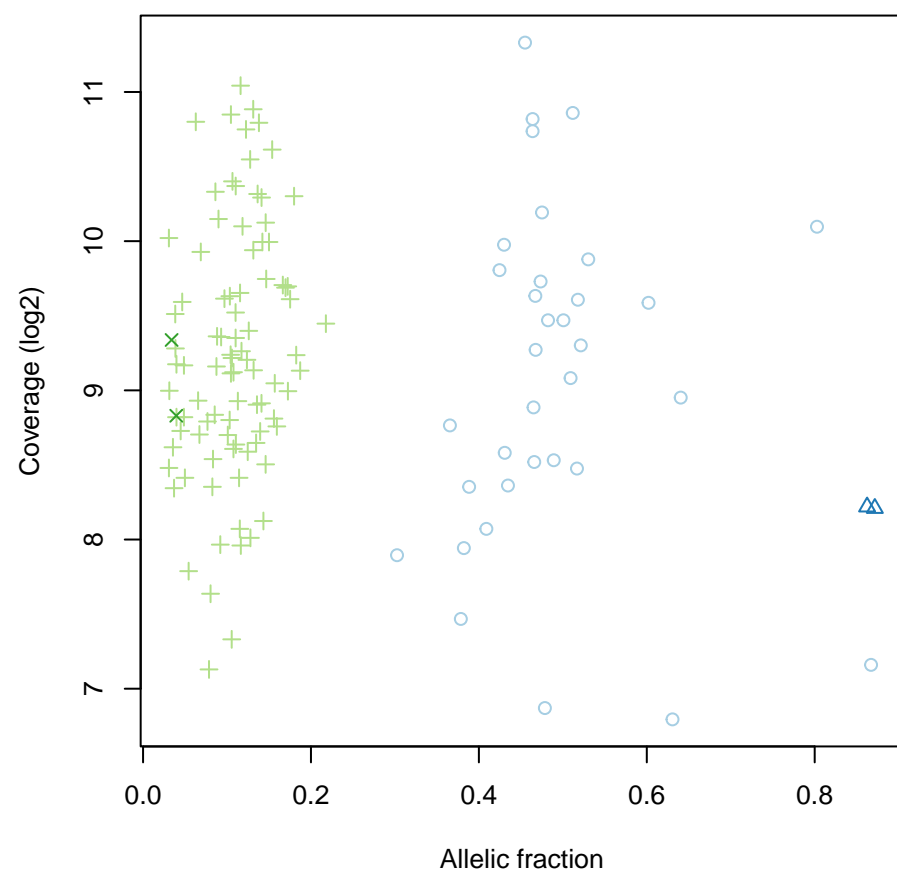
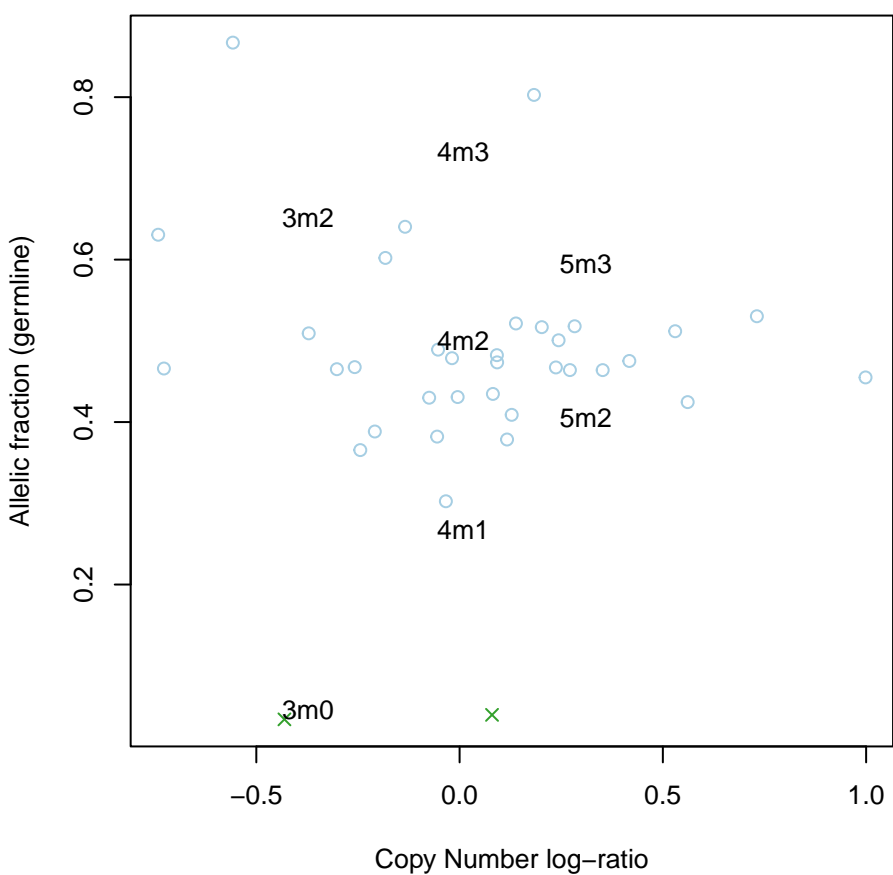
Purity: 0.87 Tumor ploidy: 3.973 SNV log-likelihood: -324.18 GoF: 66.6% Mean coverage: 350;709



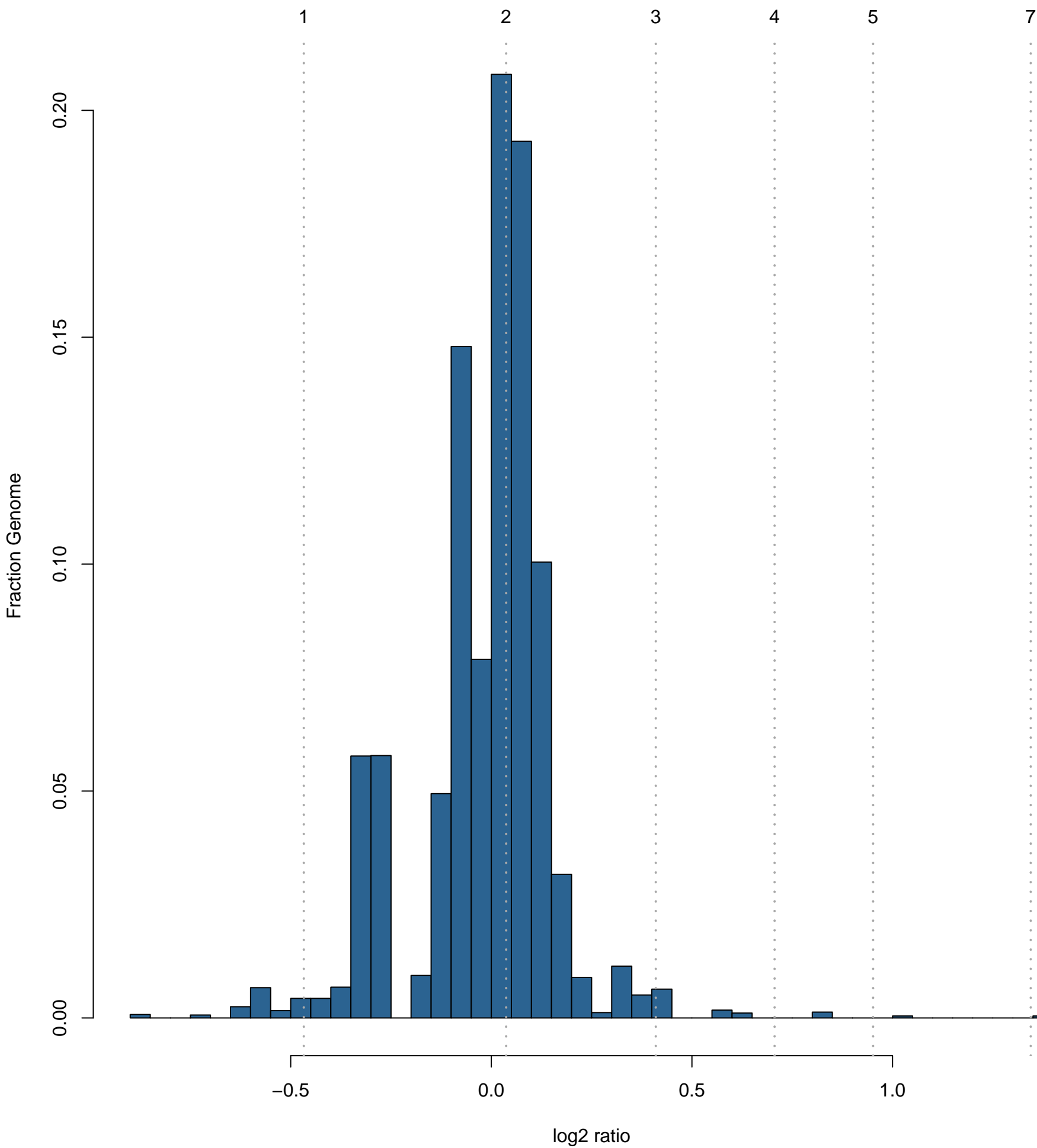
SCNA-fit log-likelihood: -4079.72



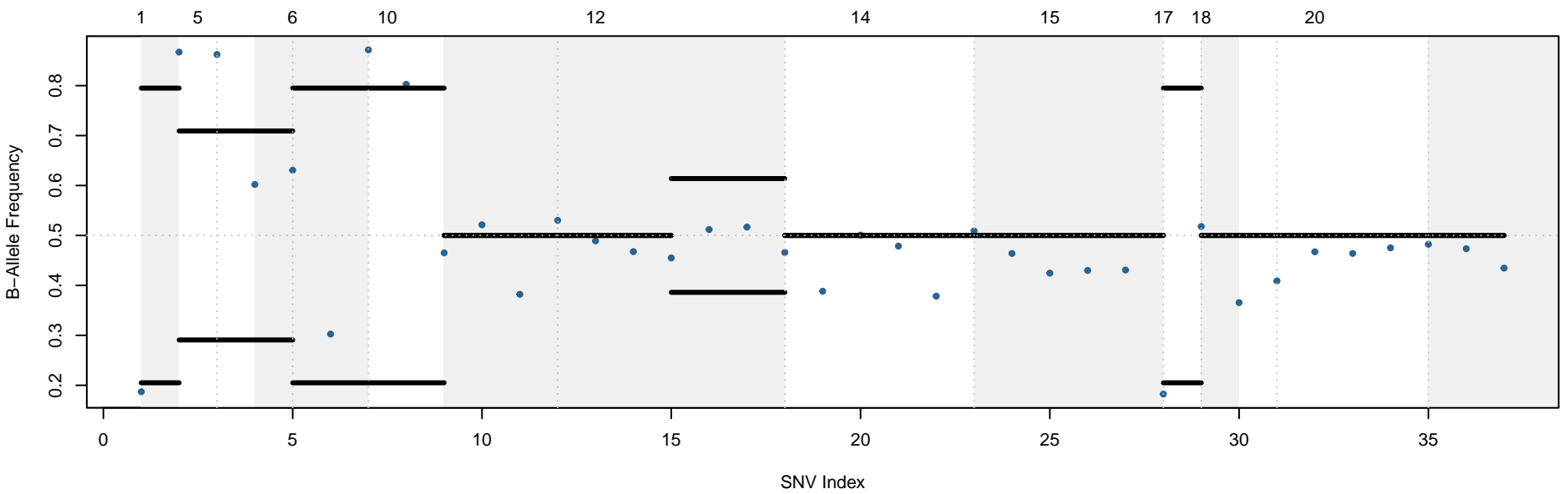




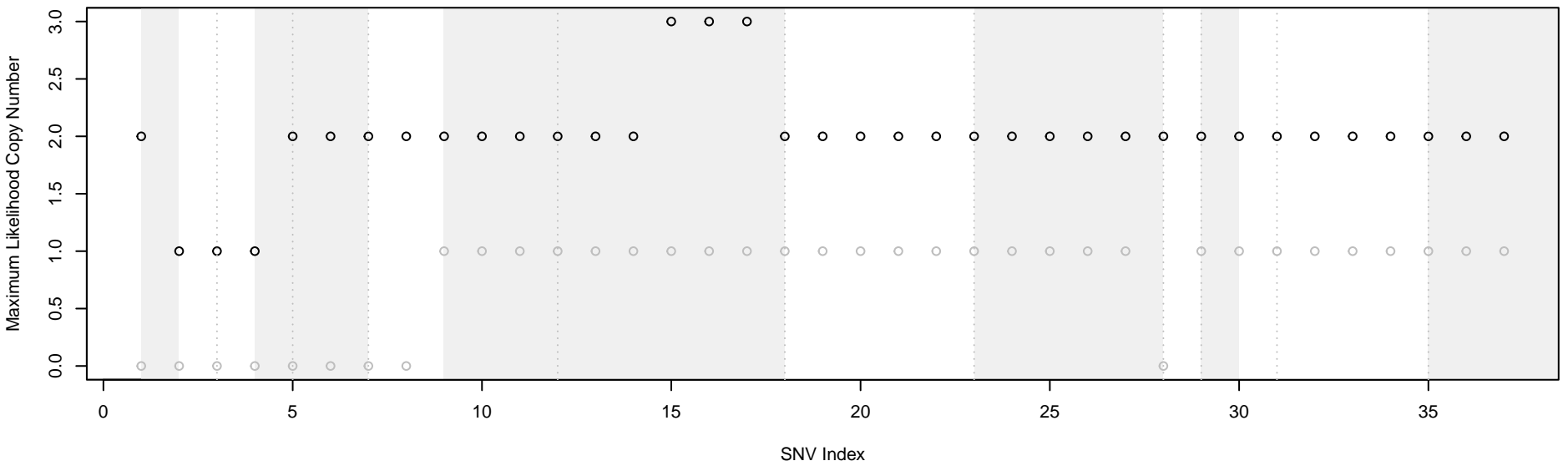
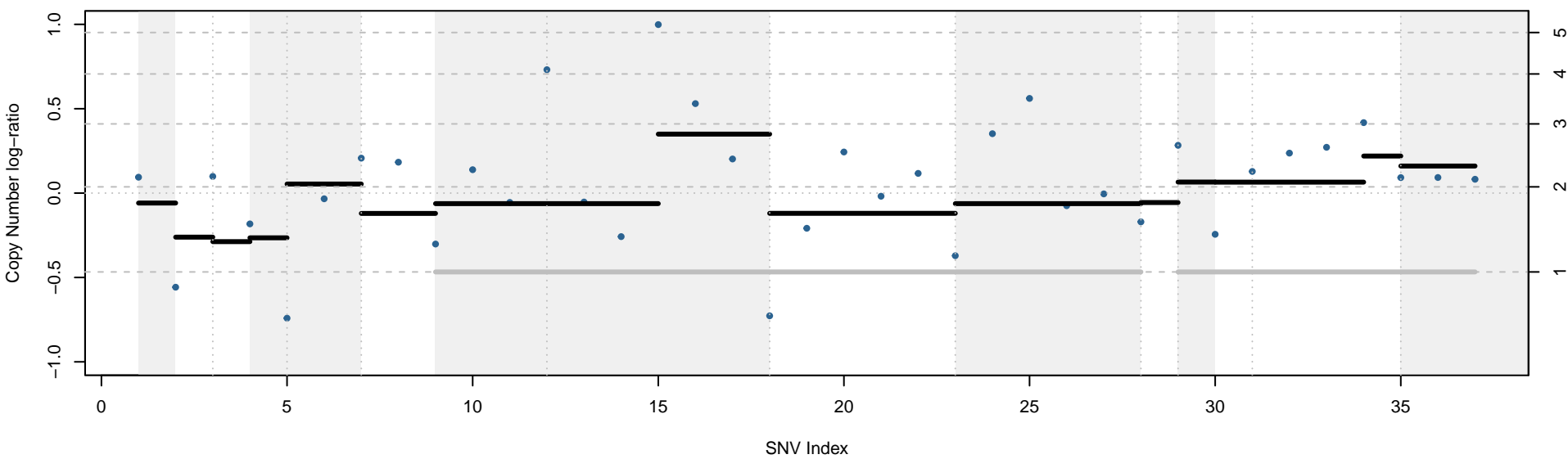
Purity: 0.59 Tumor ploidy: 1.914

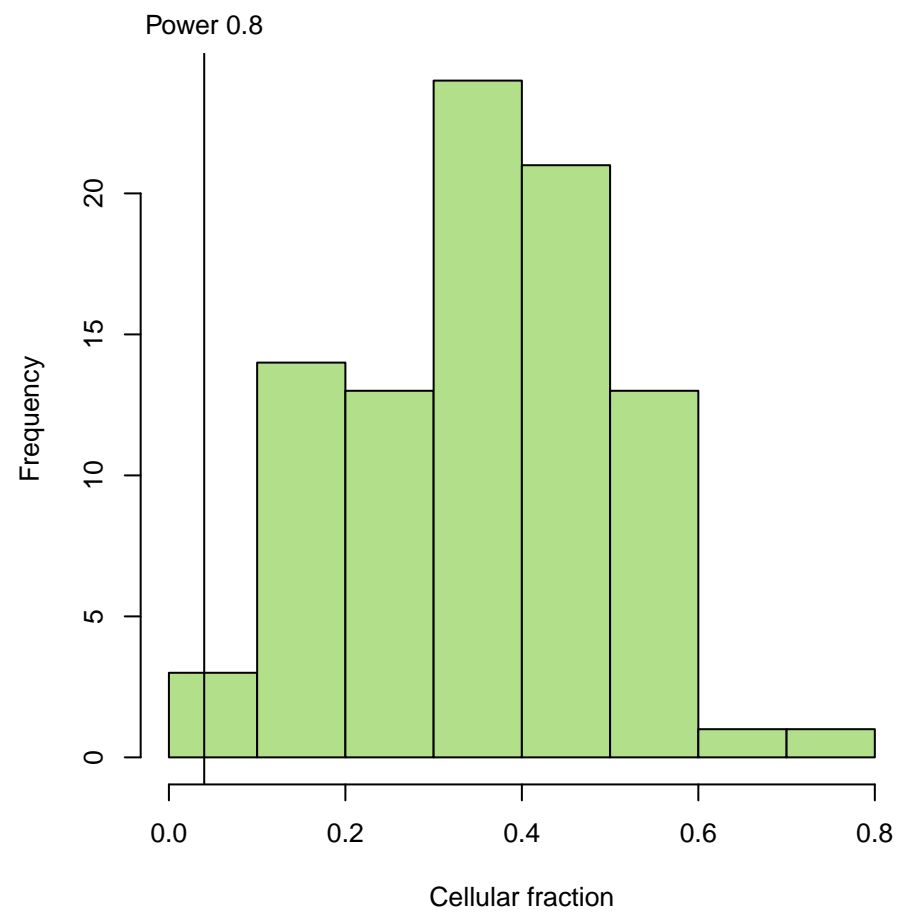
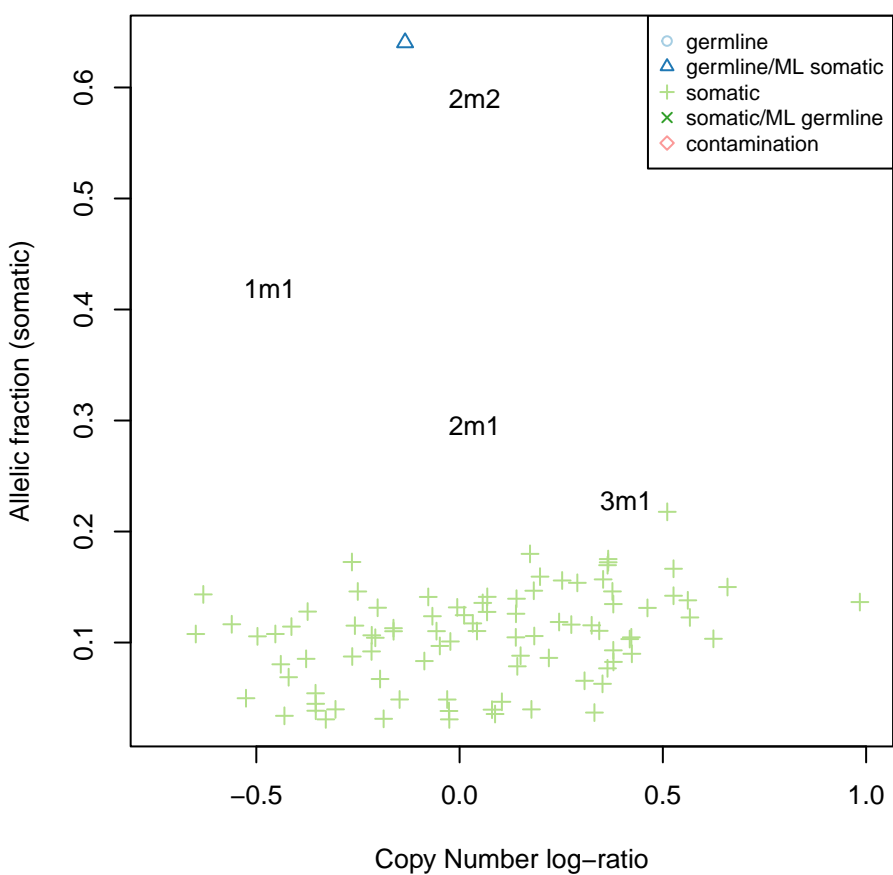
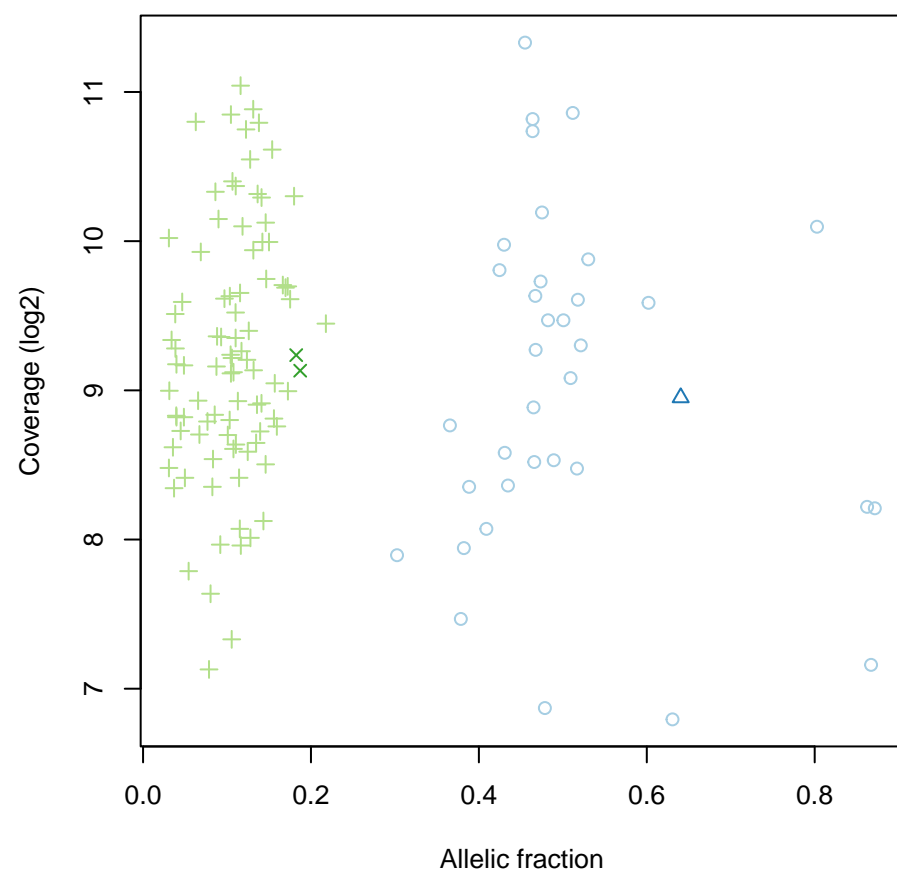
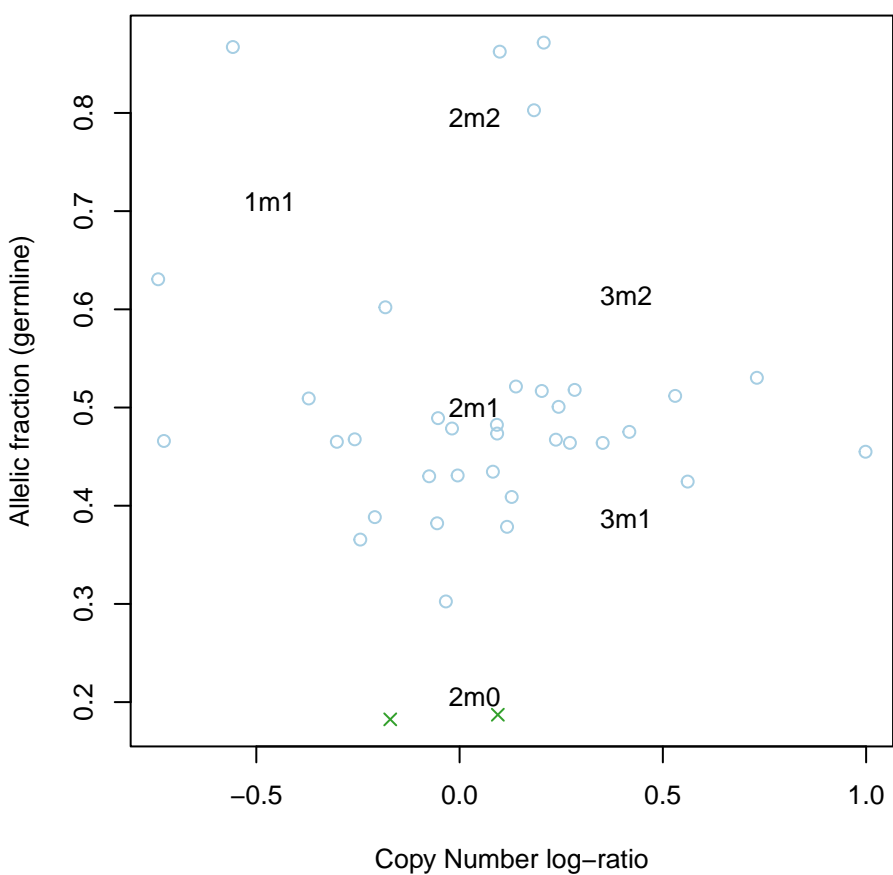


Purity: 0.59 Tumor ploidy: 1.914 SNV log-likelihood: -281.09 GoF: 22% Mean coverage: 350;709

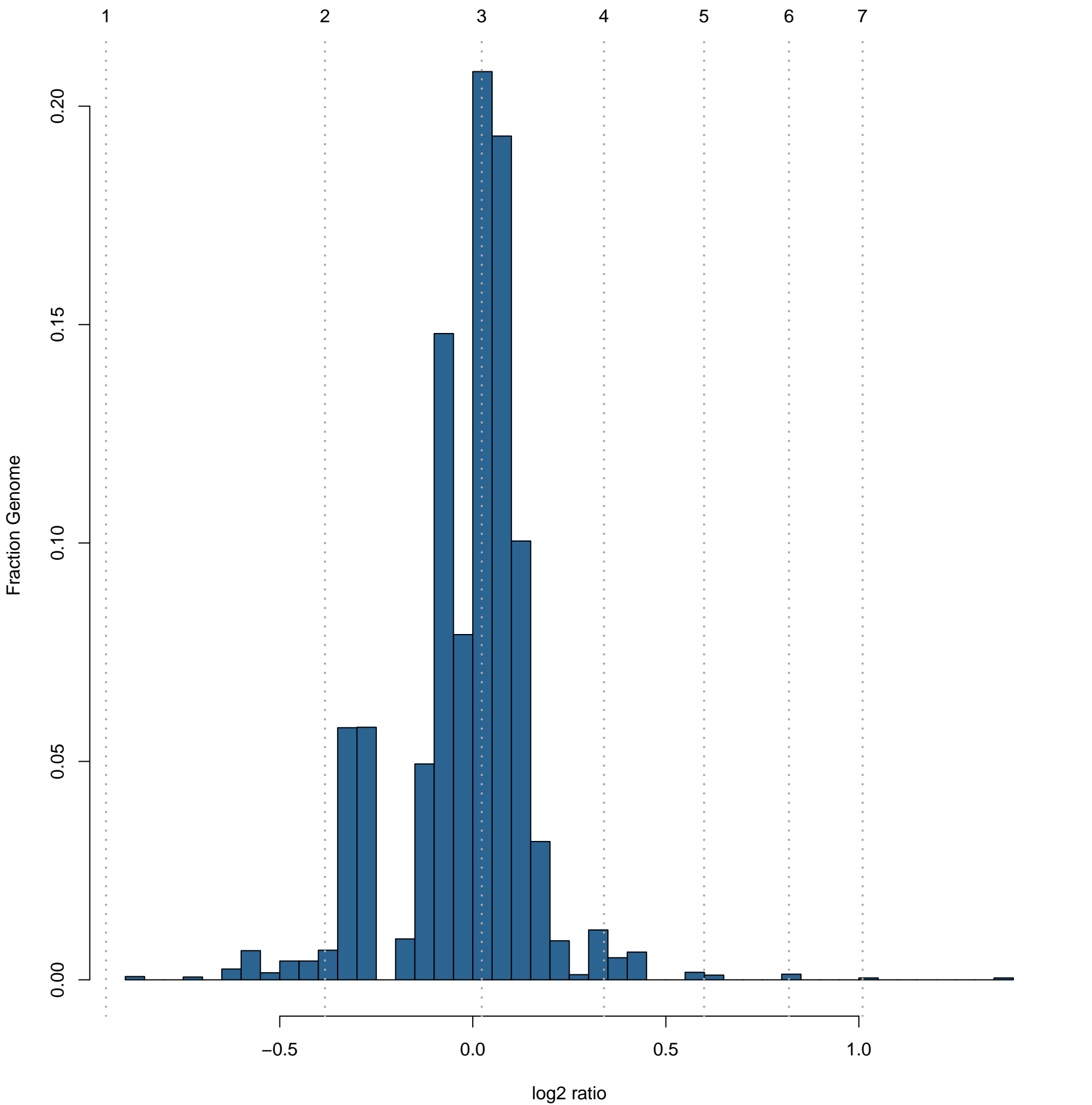


SCNA-fit log-likelihood: -4184.52

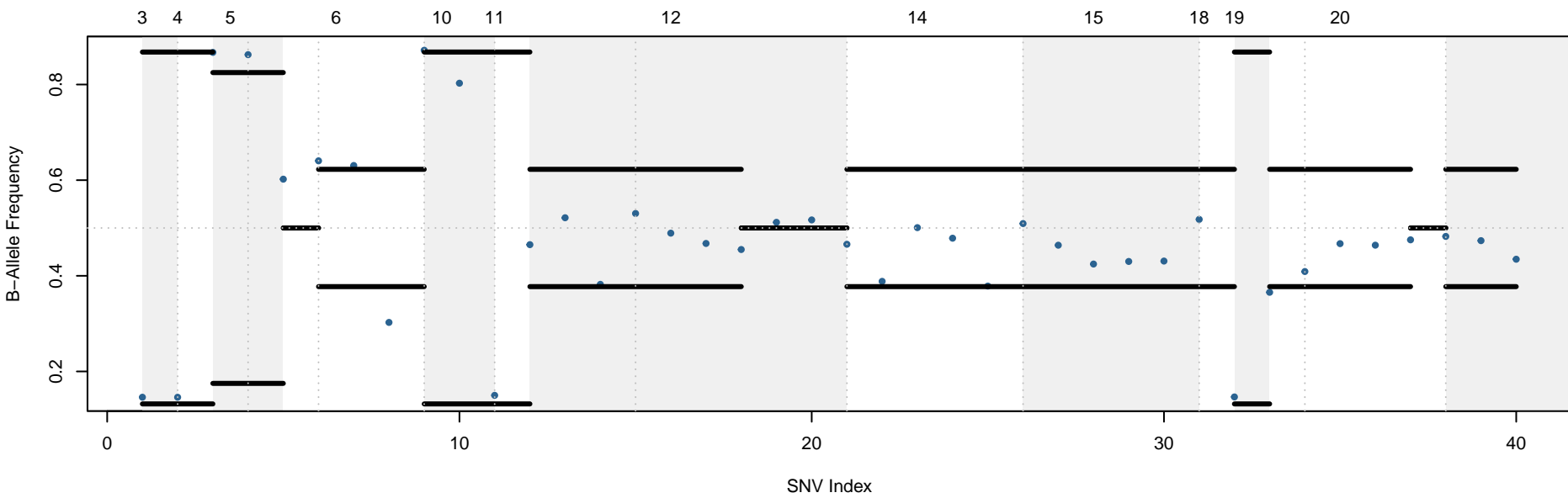




Purity: 0.65 Tumor ploidy: 2.935



Purity: 0.65 Tumor ploidy: 2.935 SNV log-likelihood: -332.76 GoF: 55.9% Mean coverage: 350;709



SCNA-fit log-likelihood: -4091.07

