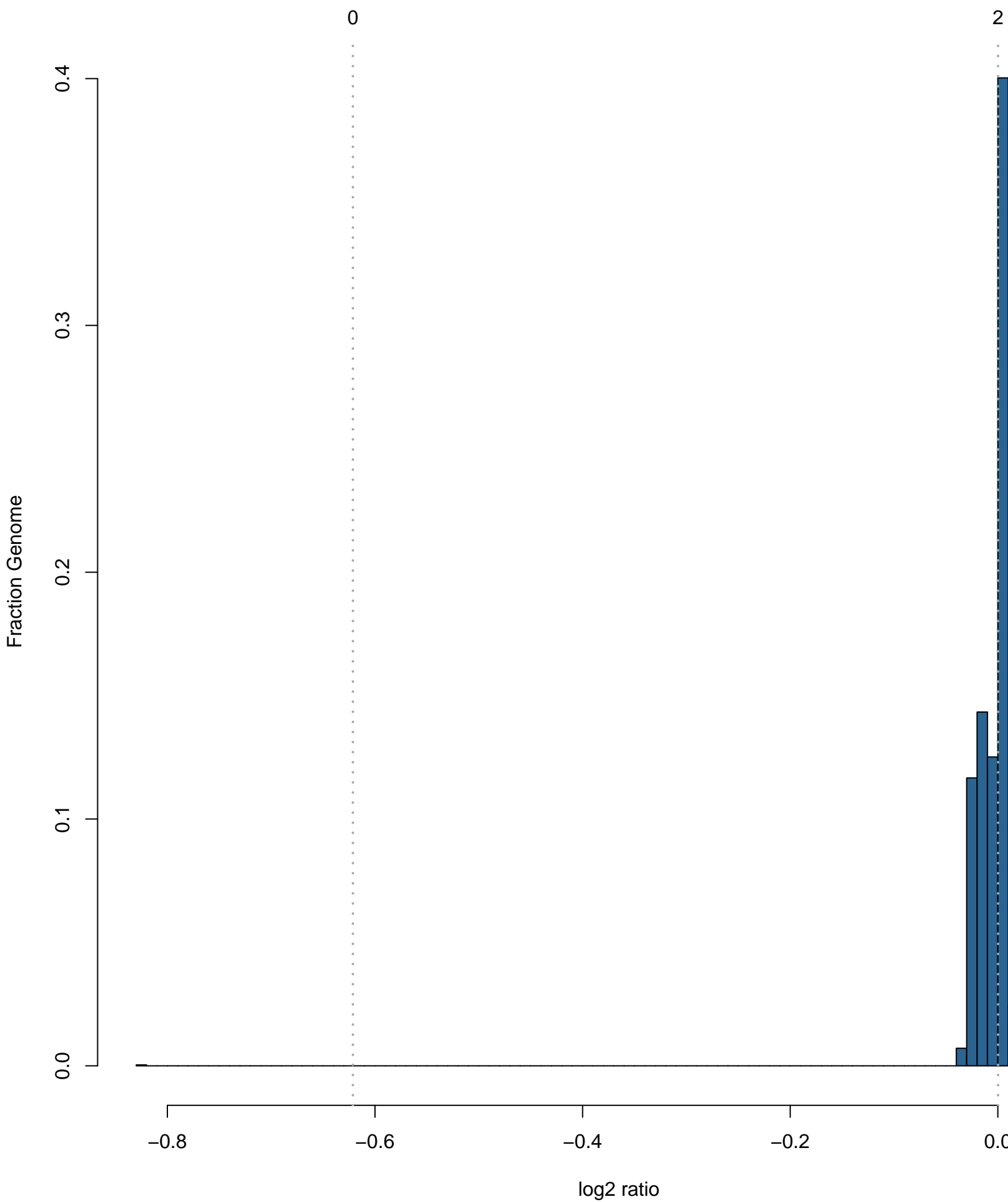
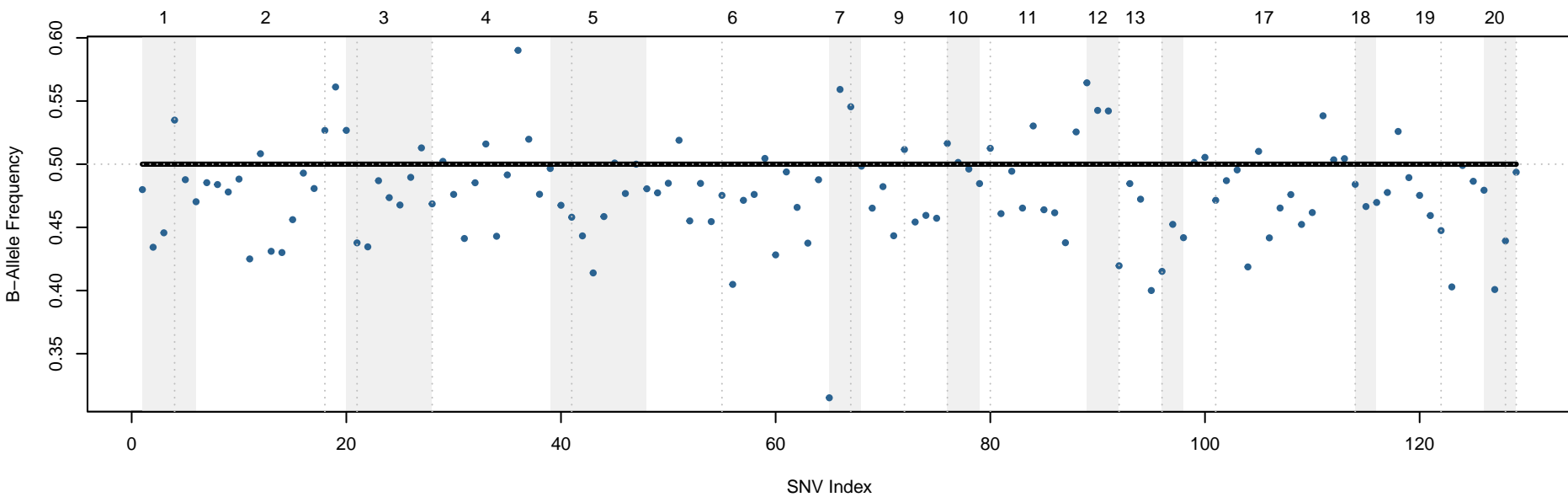


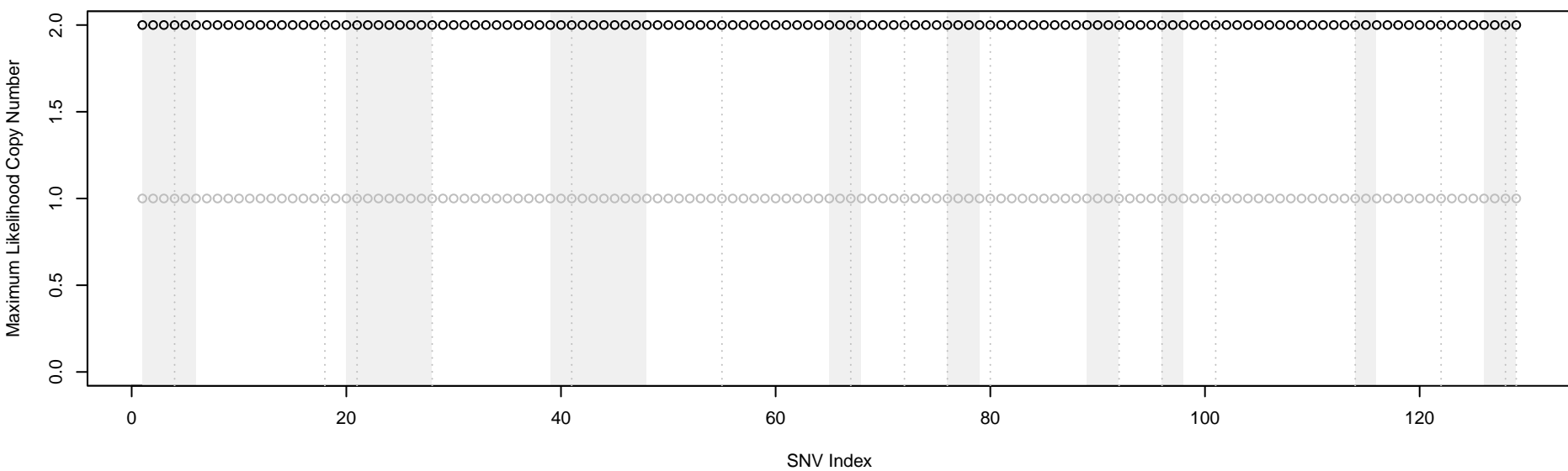
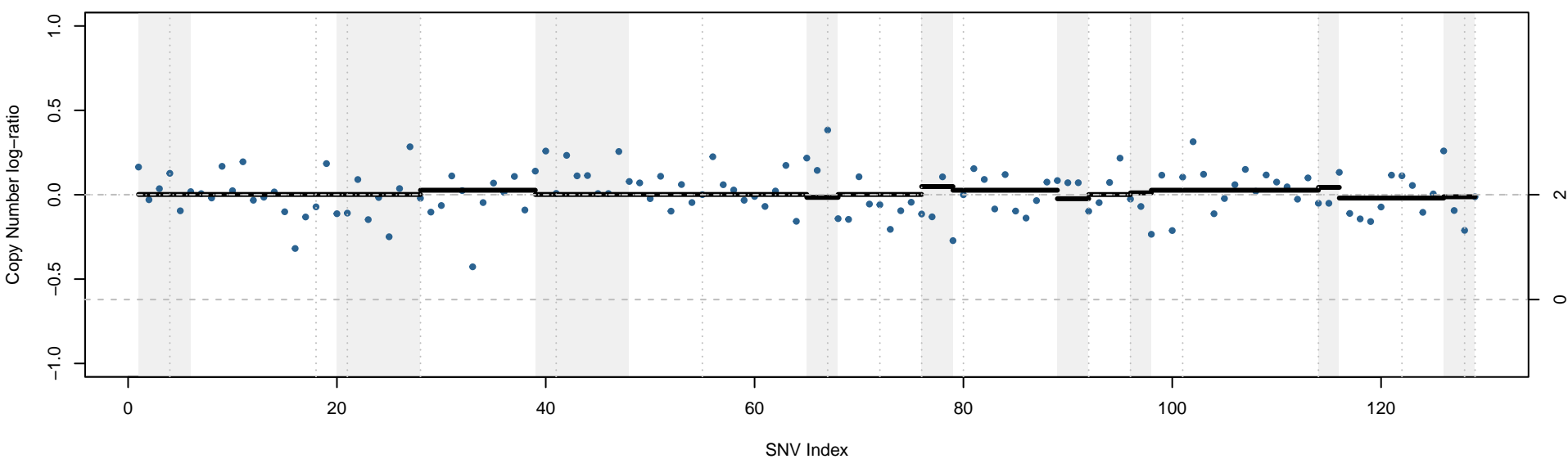
Purity: 0.35 Tumor ploidy: 1.999

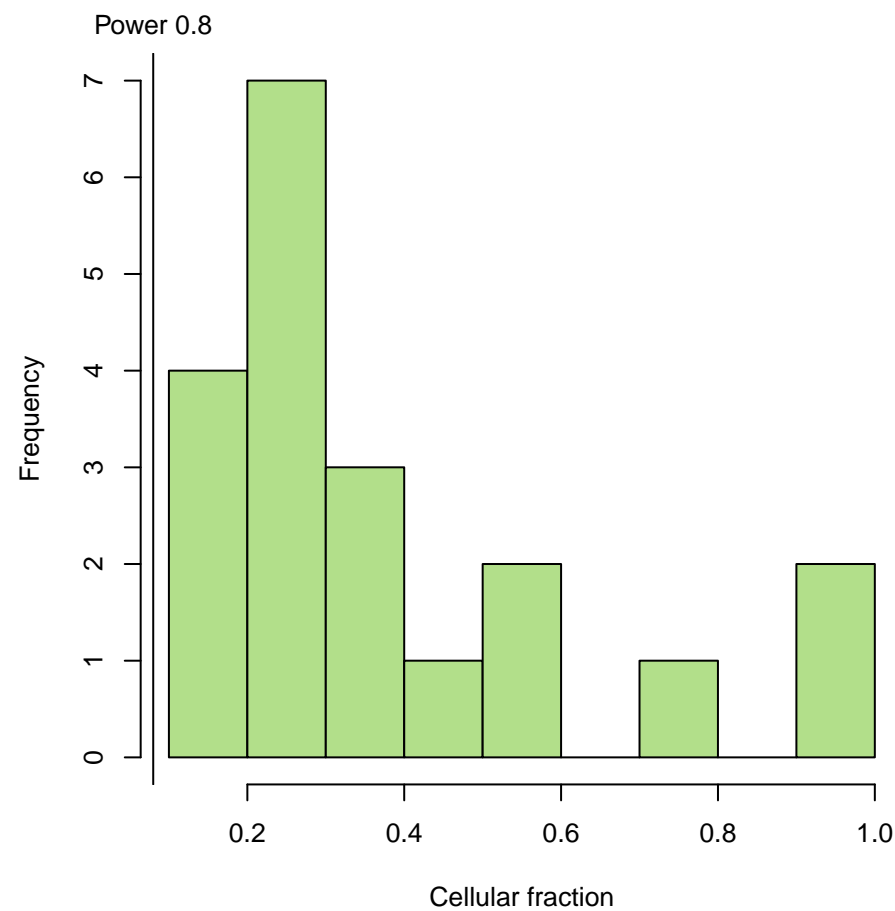
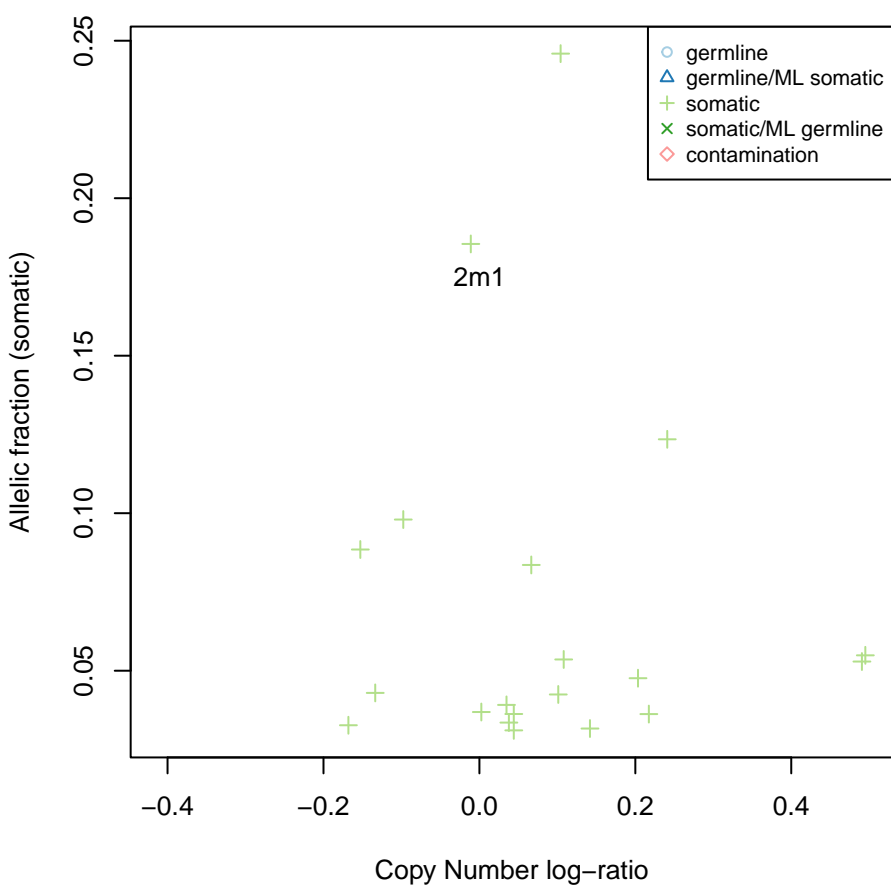
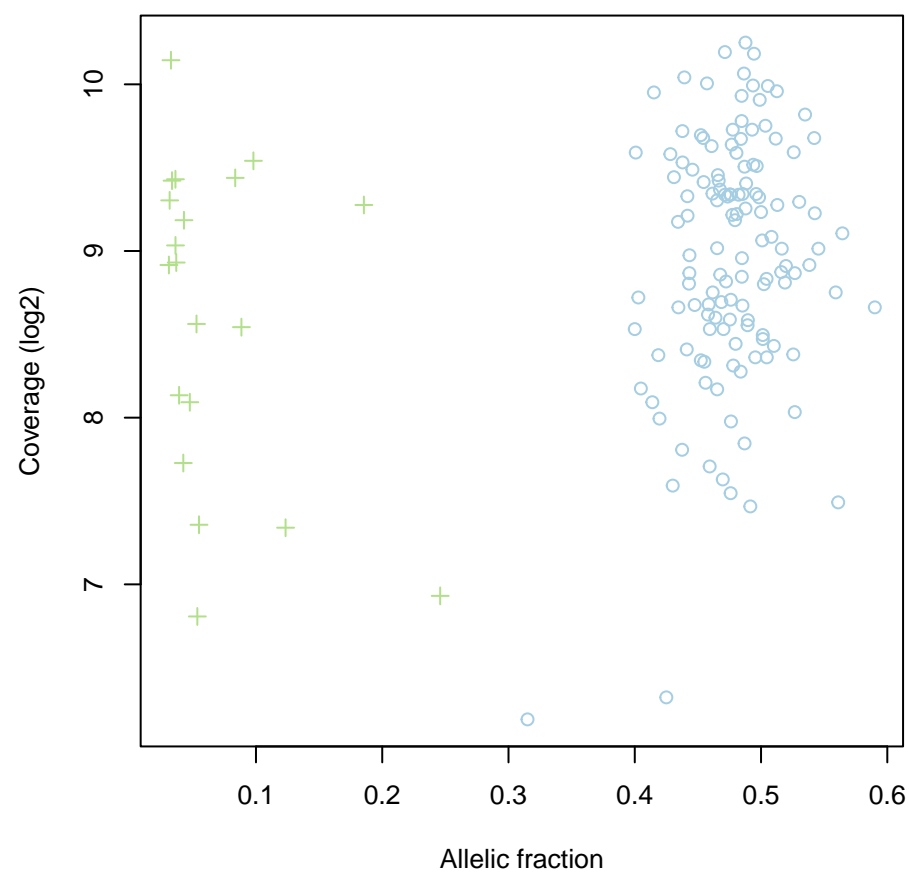
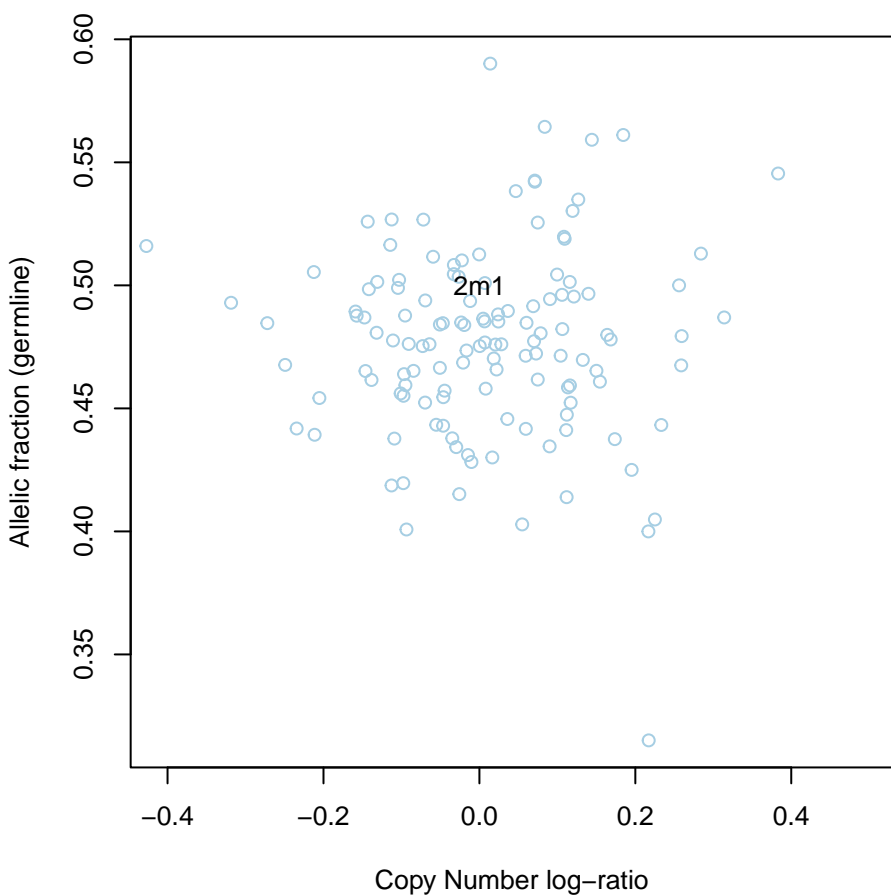


Purity: 0.35 Tumor ploidy: 1.999 SNV log-likelihood: 89.77 GoF: 91.9% Mean coverage: 549,494

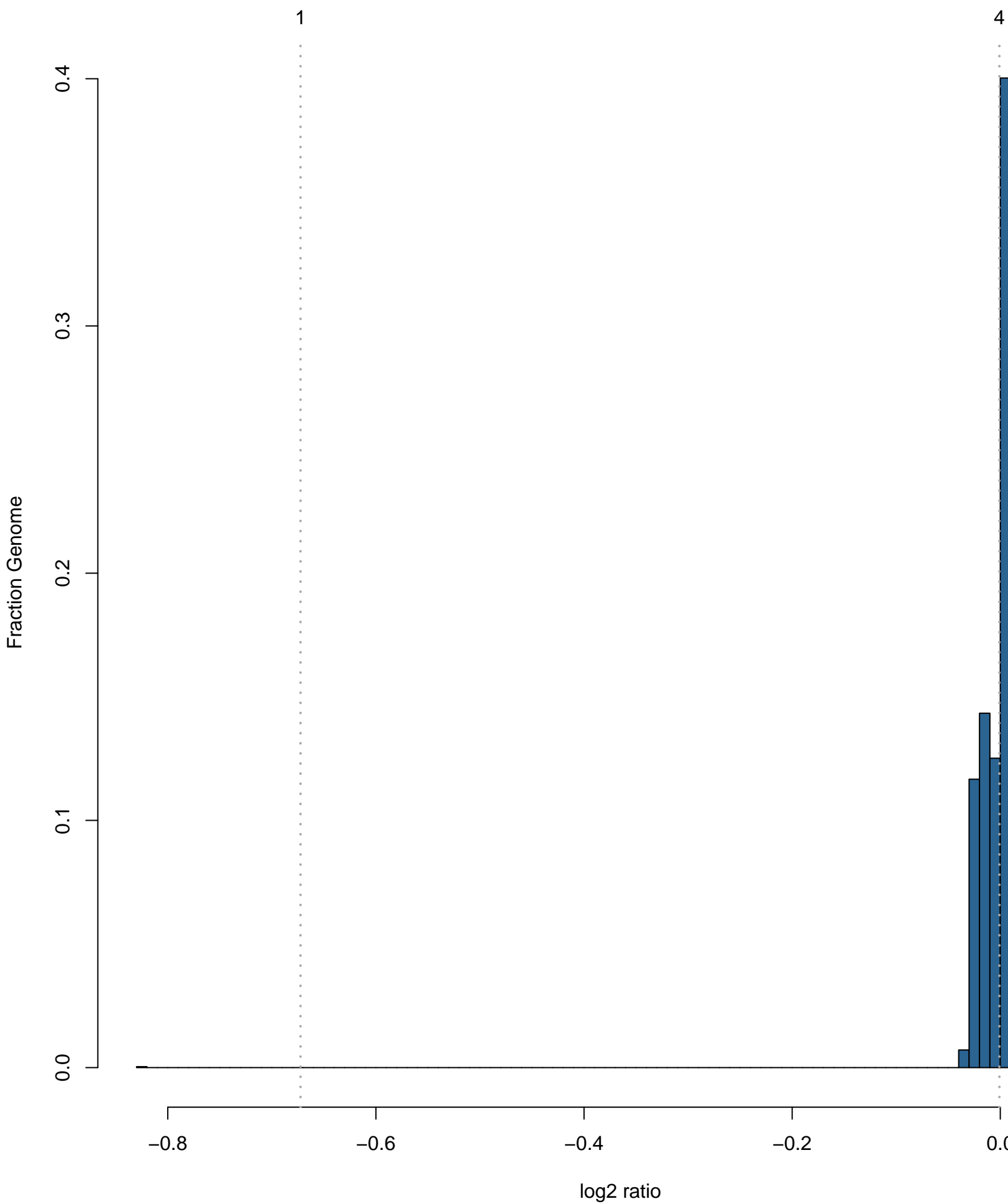


SCNA-fit log-likelihood: -4217.63

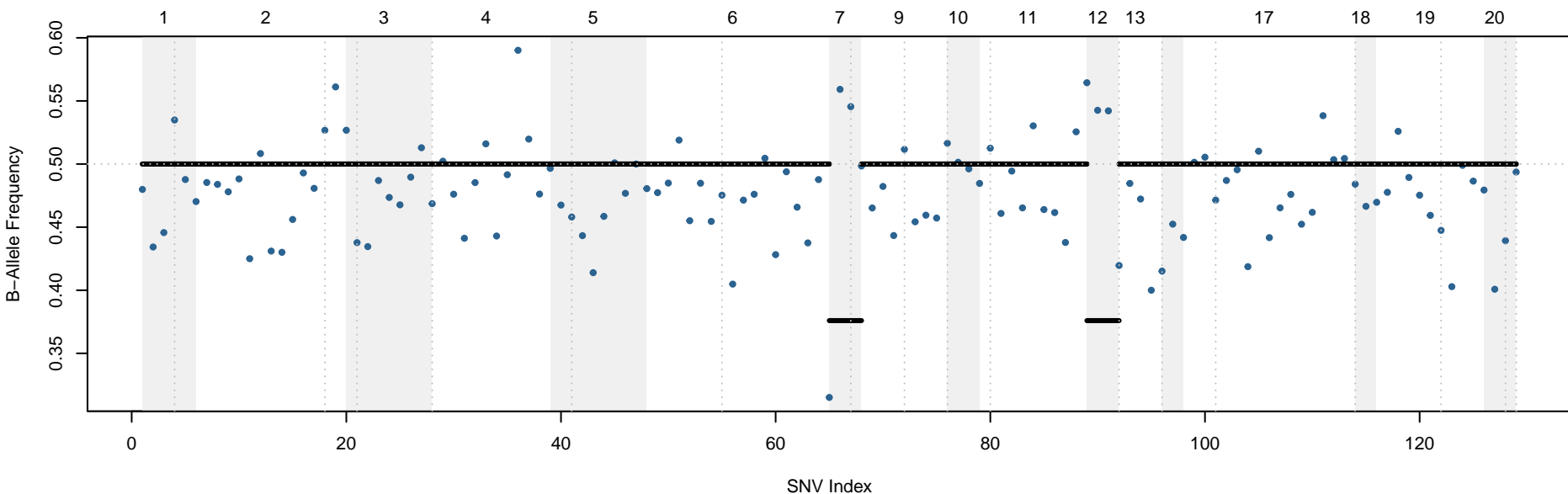




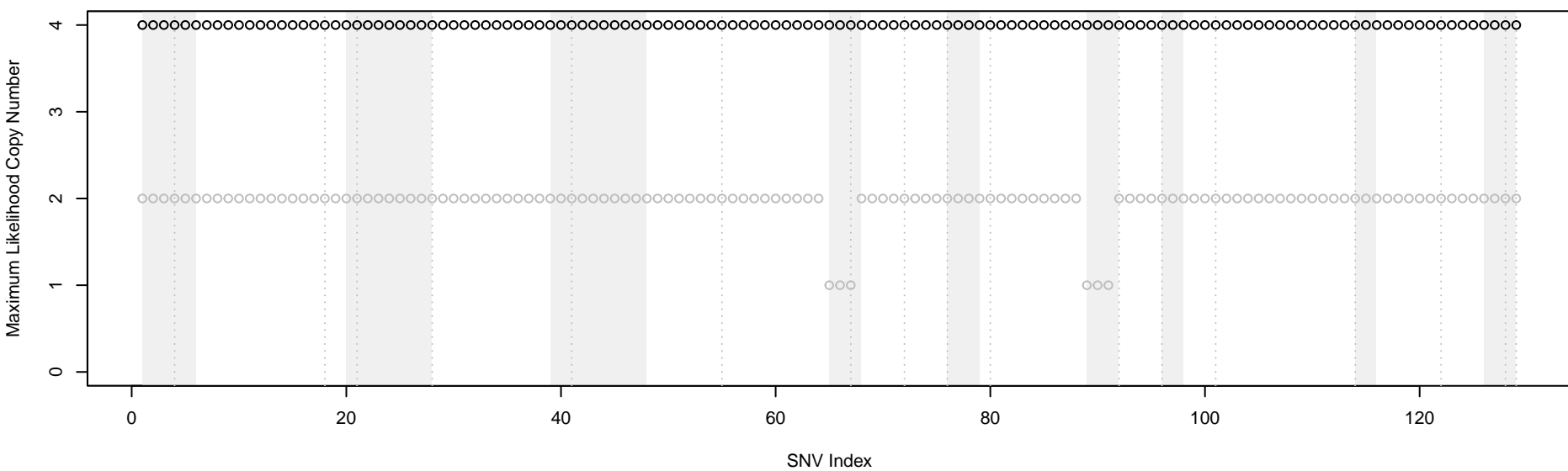
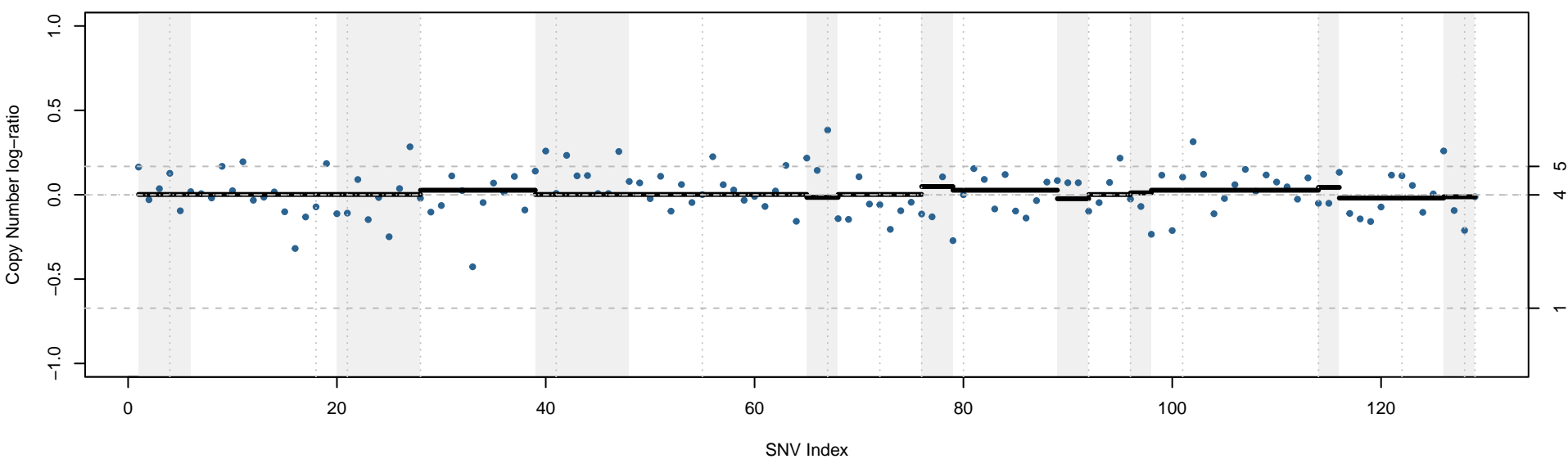
Purity: 0.33 Tumor ploidy: 4.005

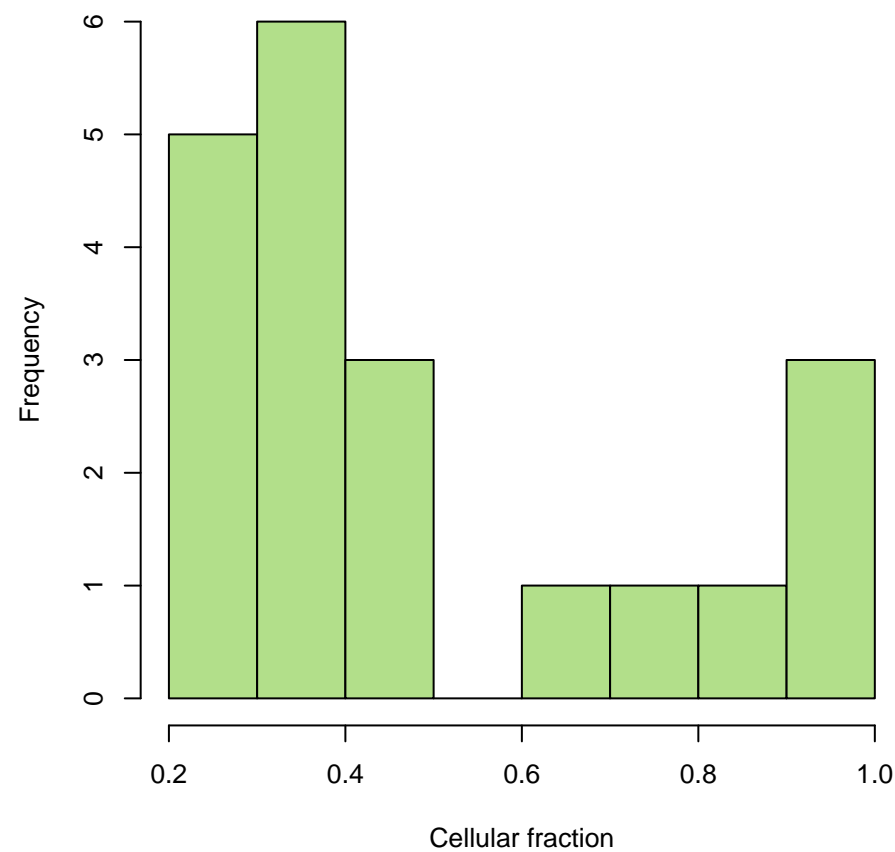
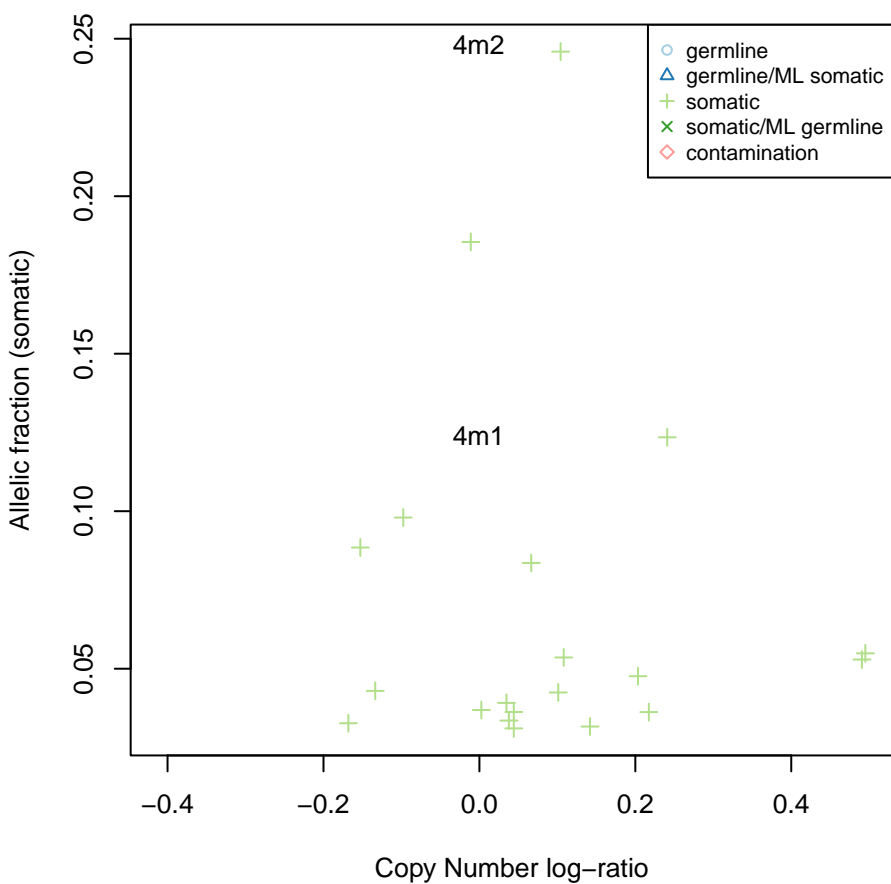
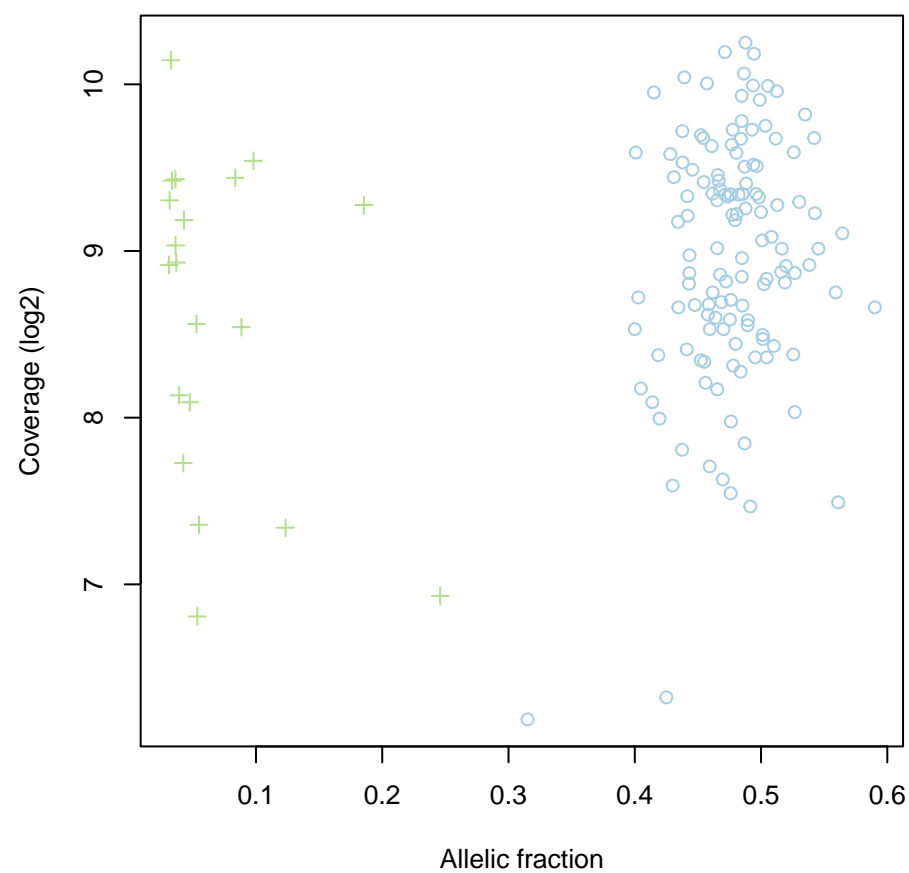
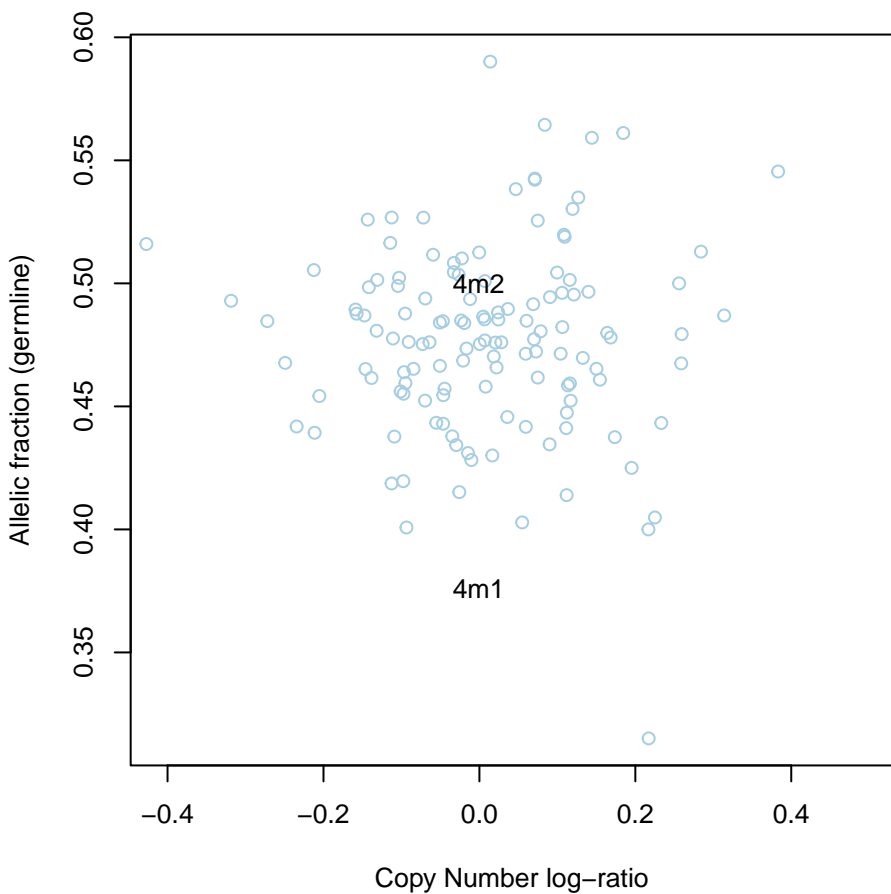


Purity: 0.33 Tumor ploidy: 4.005 SNV log-likelihood: 28.5 GoF: 95.5% Mean coverage: 549;494

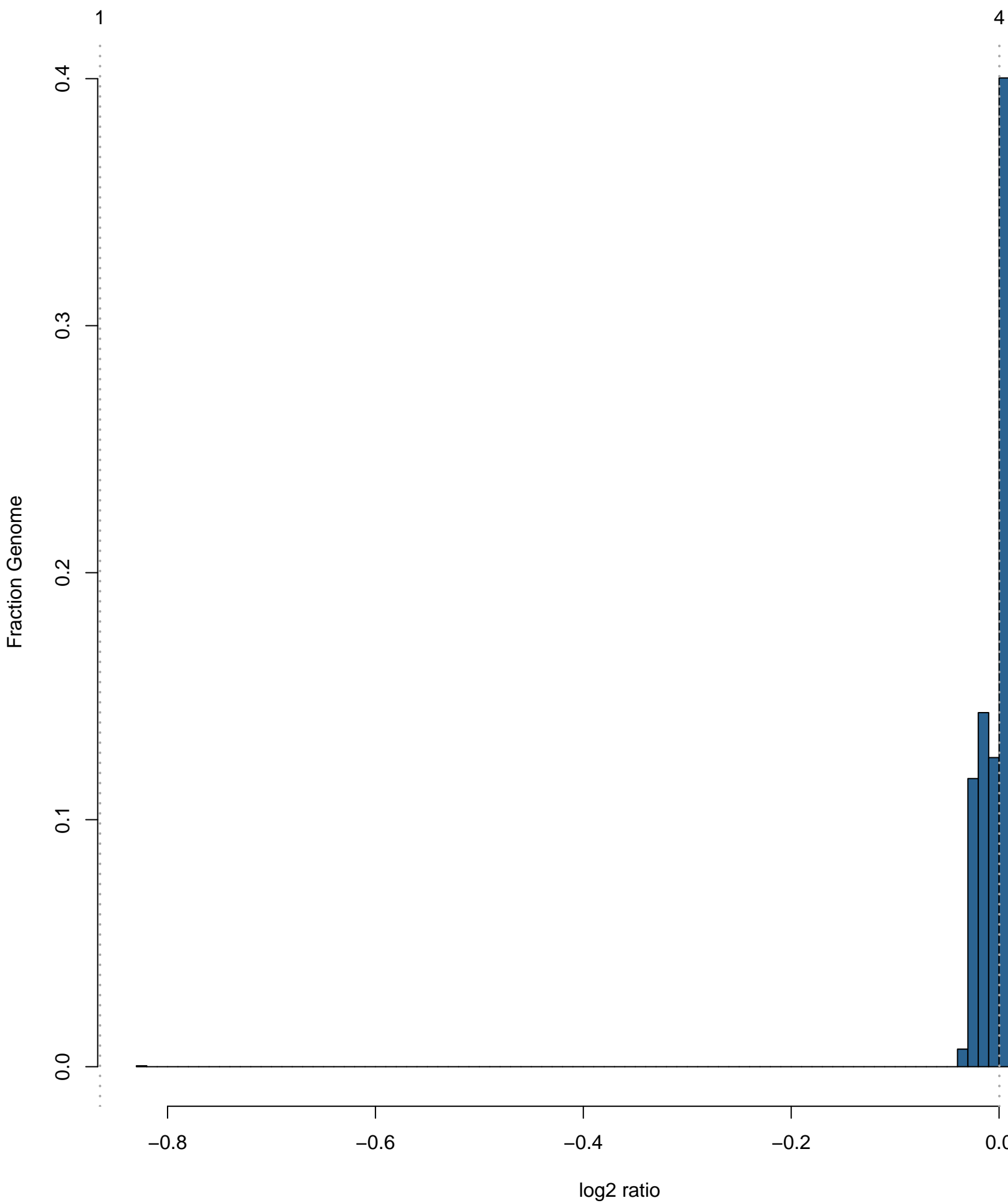


SCNA-fit log-likelihood: -4216.22

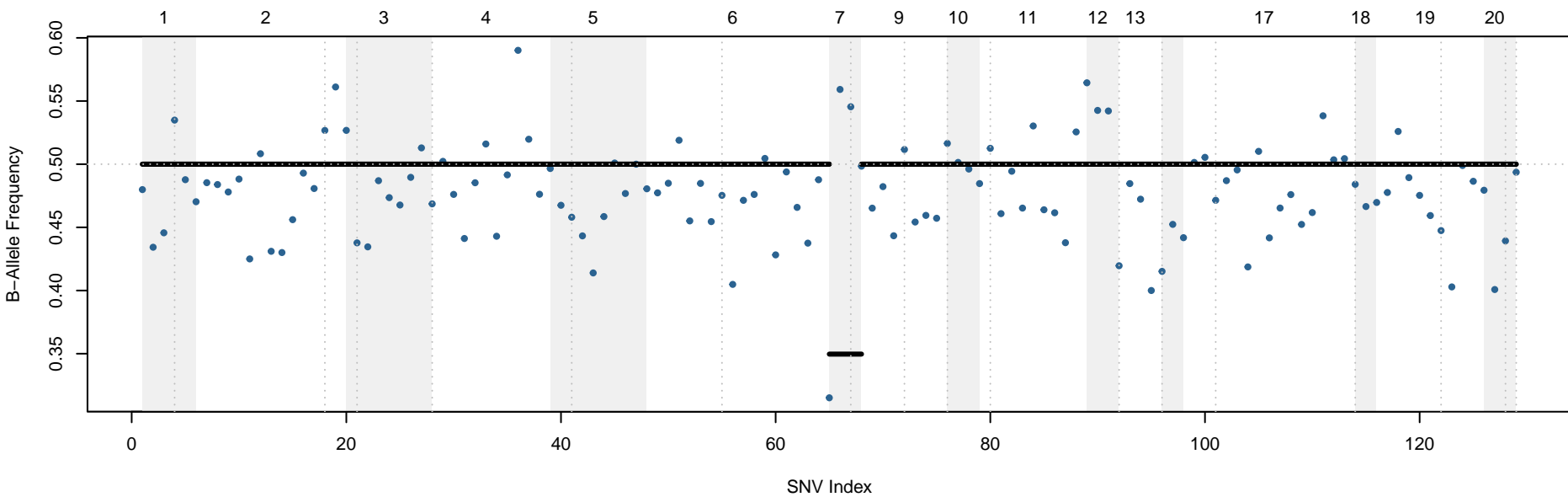




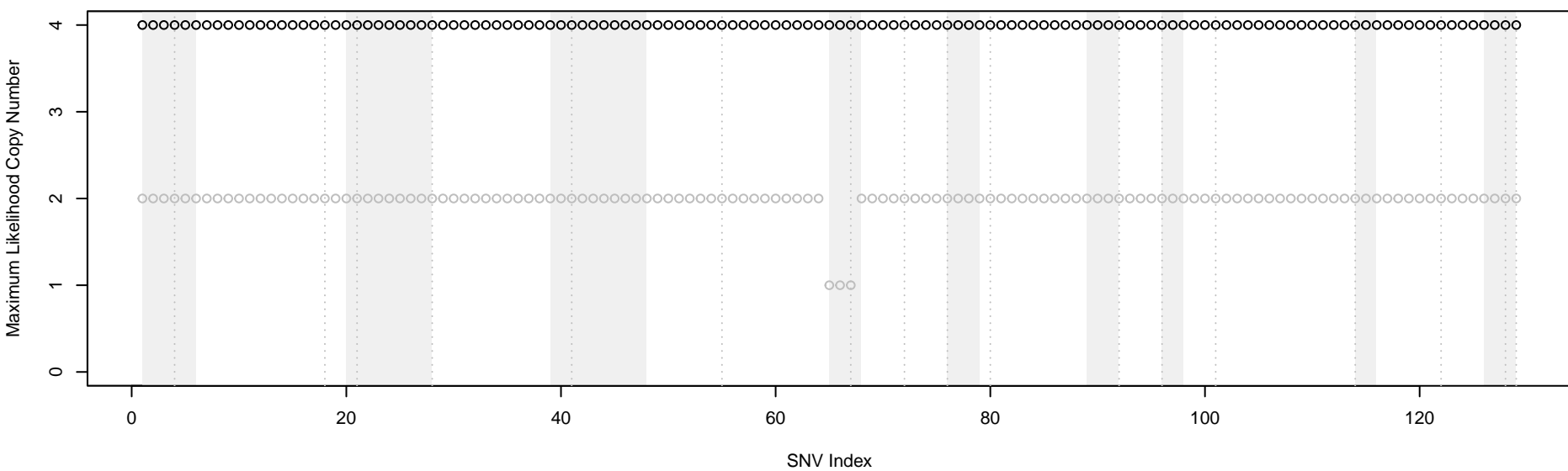
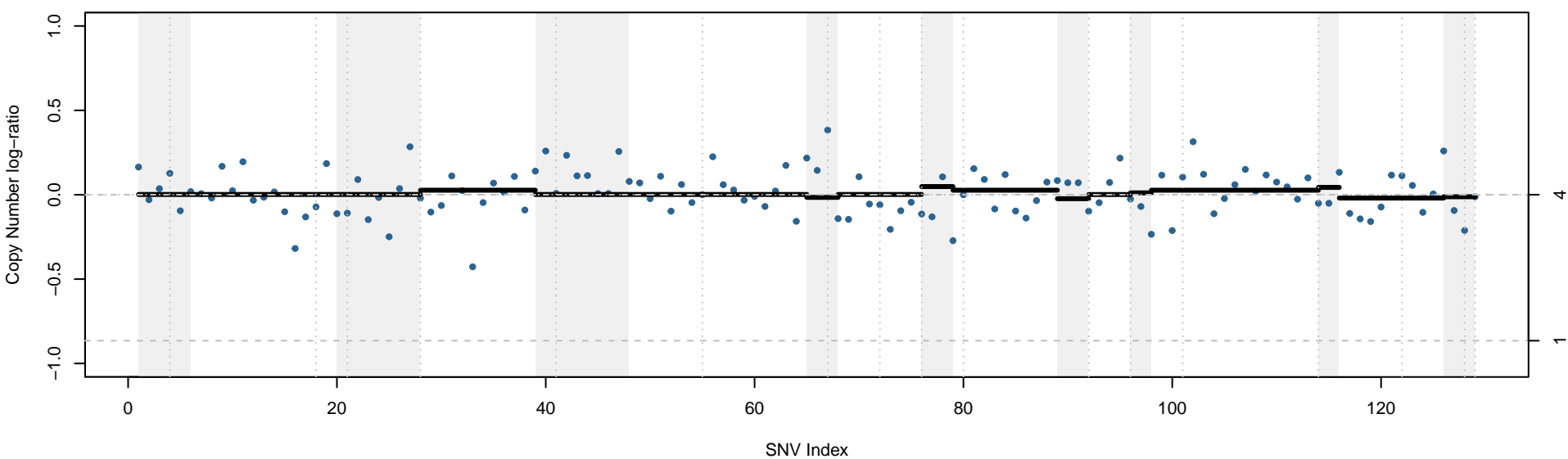
Purity: 0.43 Tumor ploidy: 3.999



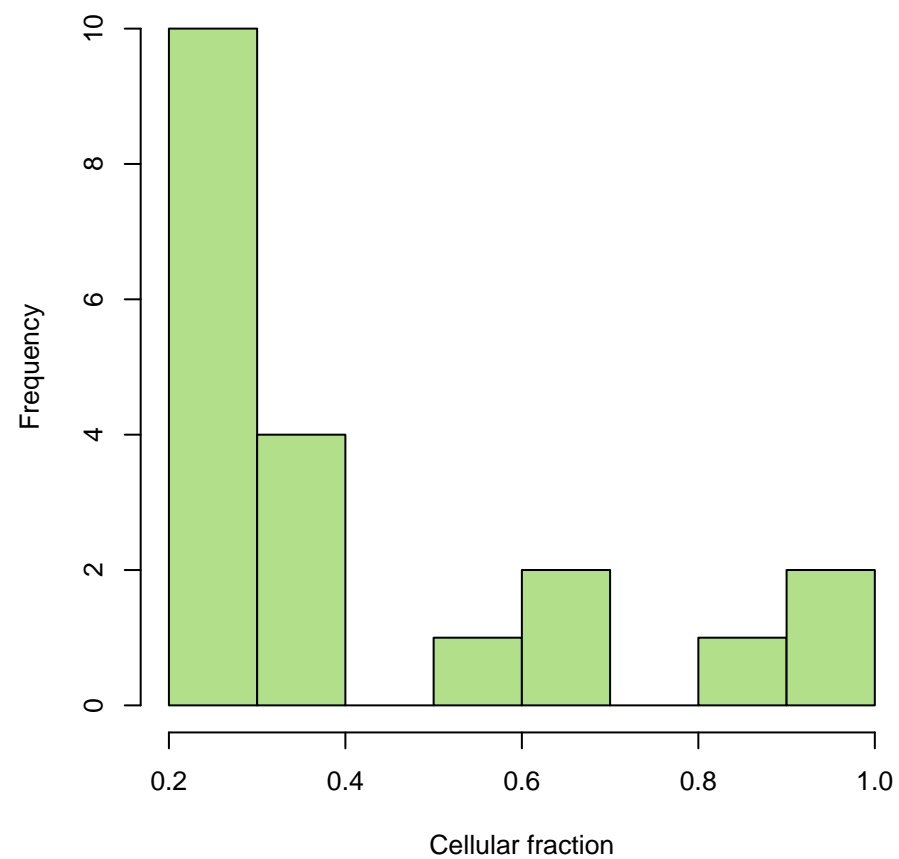
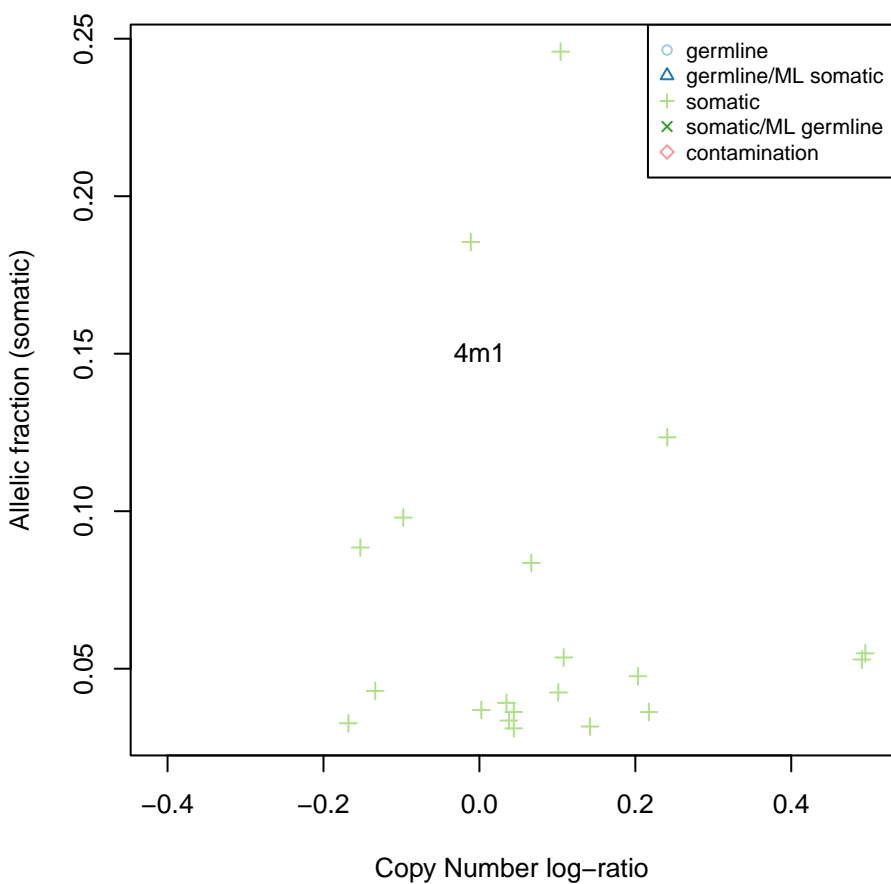
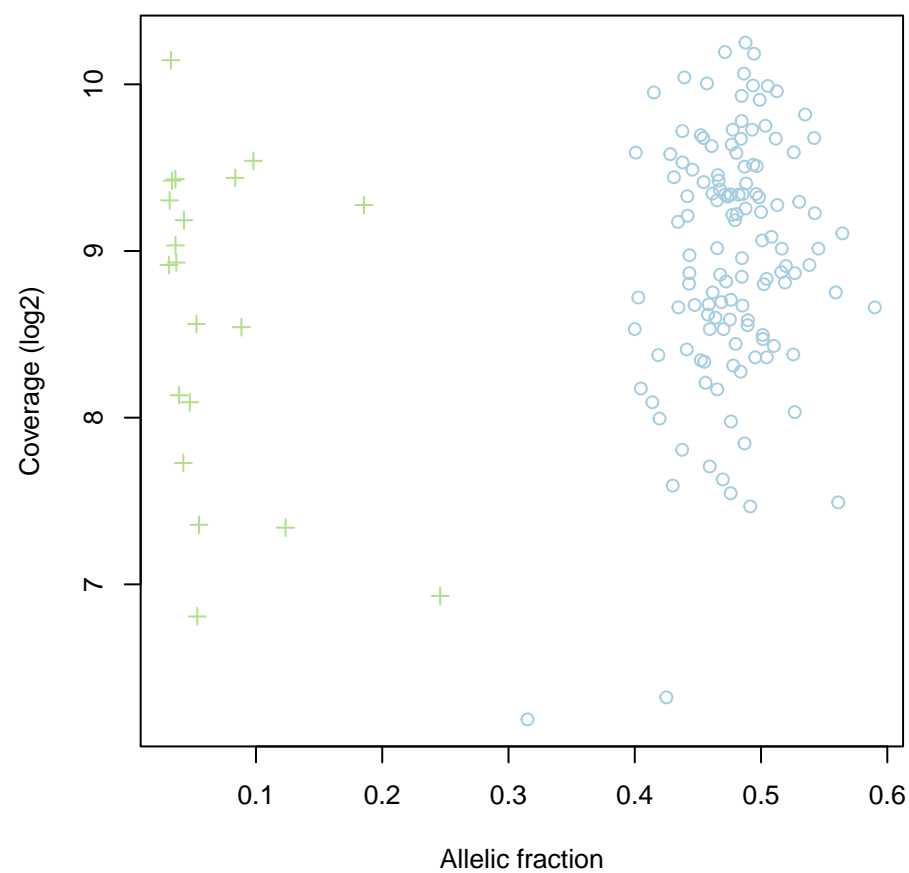
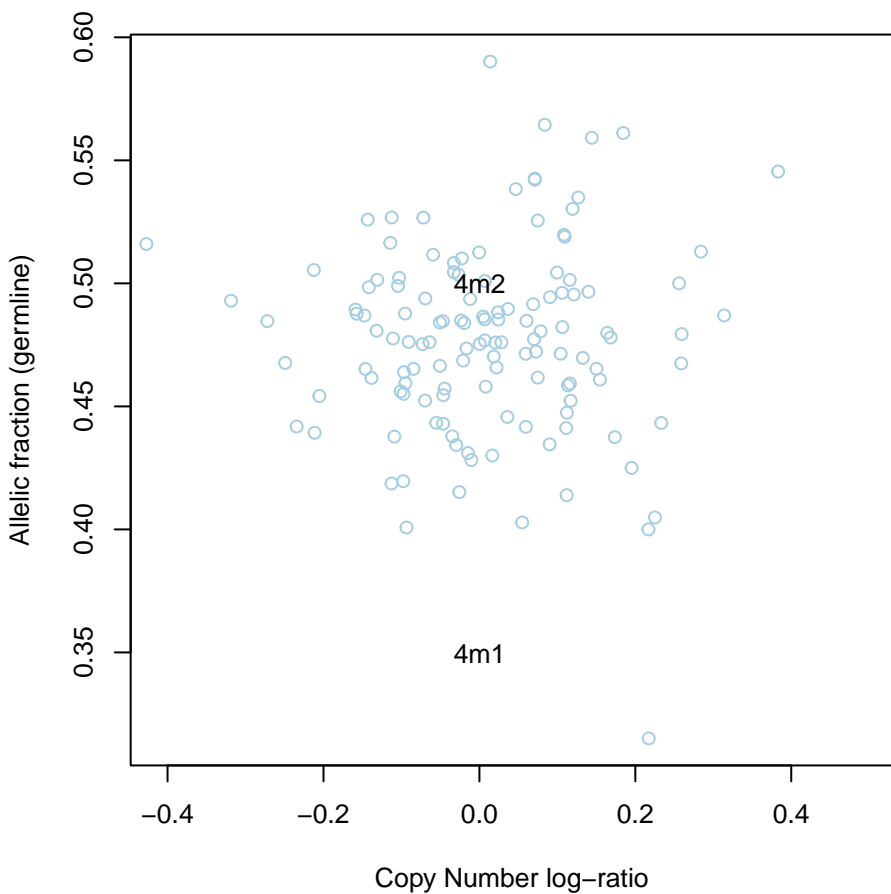
Purity: 0.43 Tumor ploidy: 3.999 SNV log-likelihood: 13.7 GoF: 94% Mean coverage: 549;494



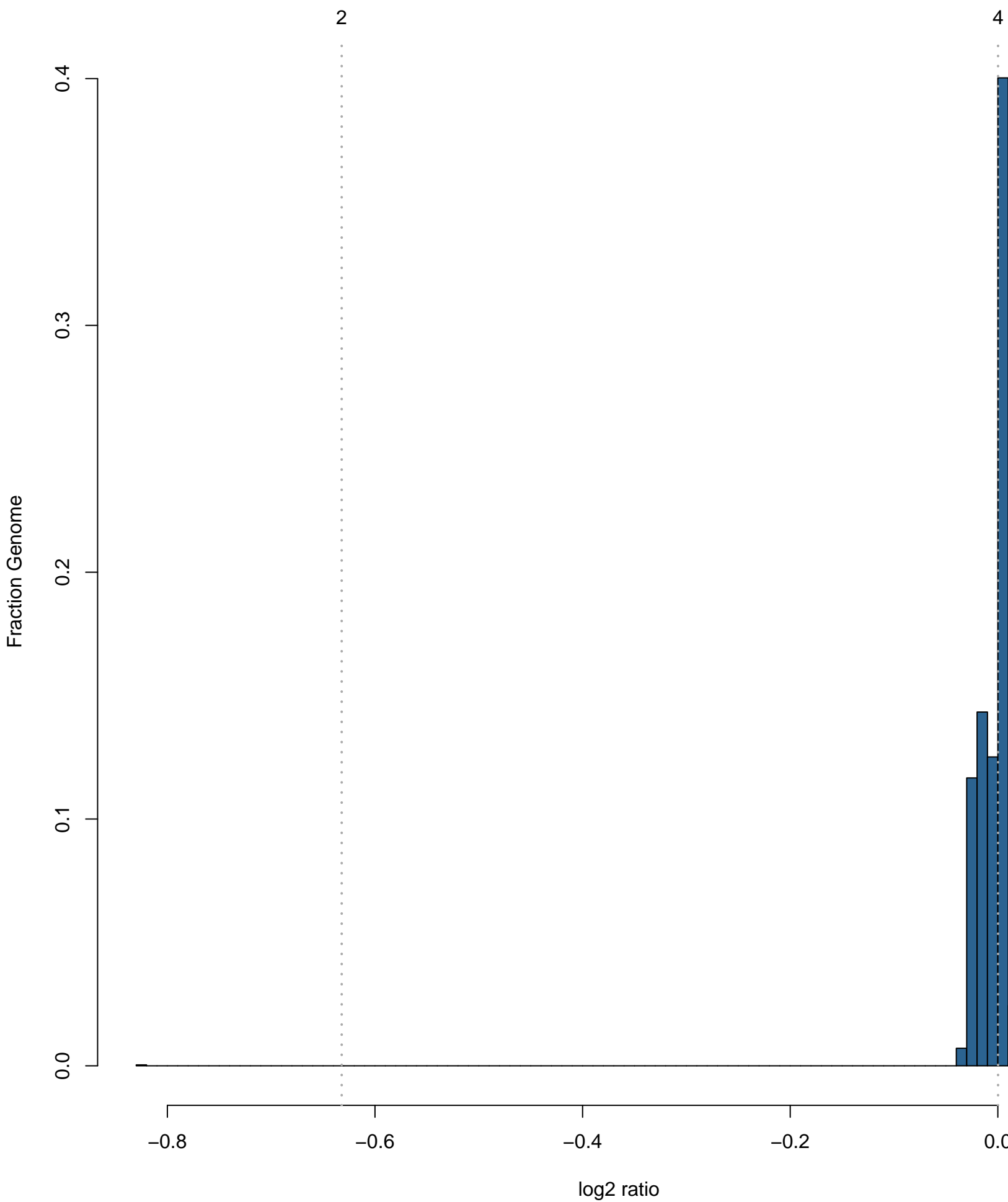
SCNA-fit log-likelihood: -4216.78



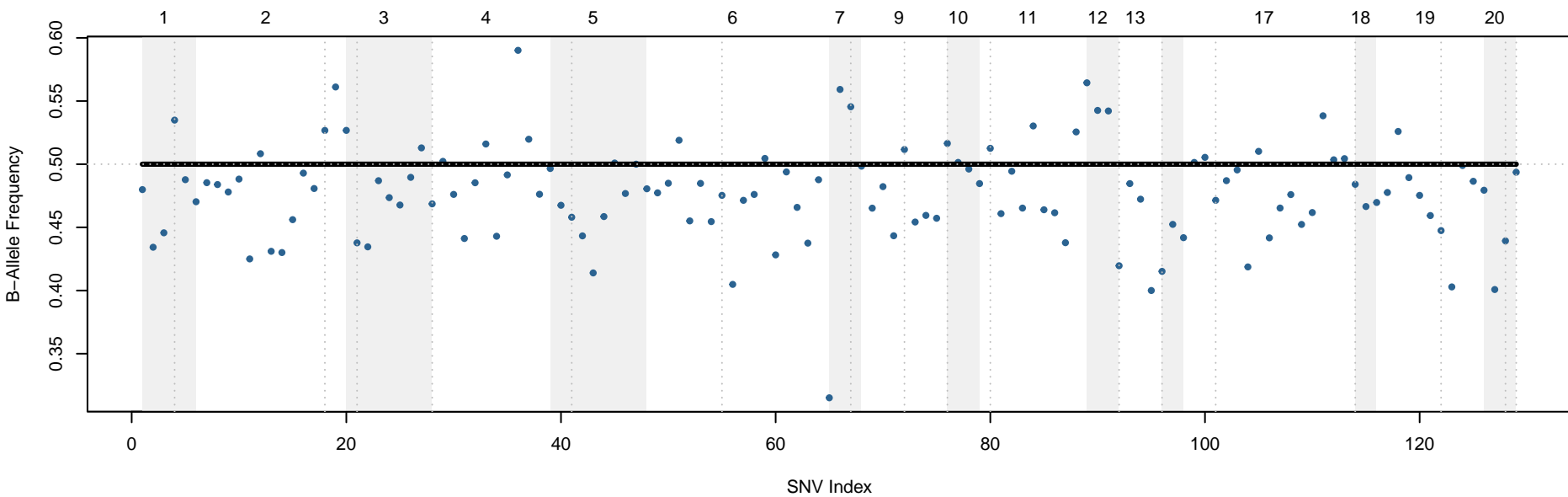




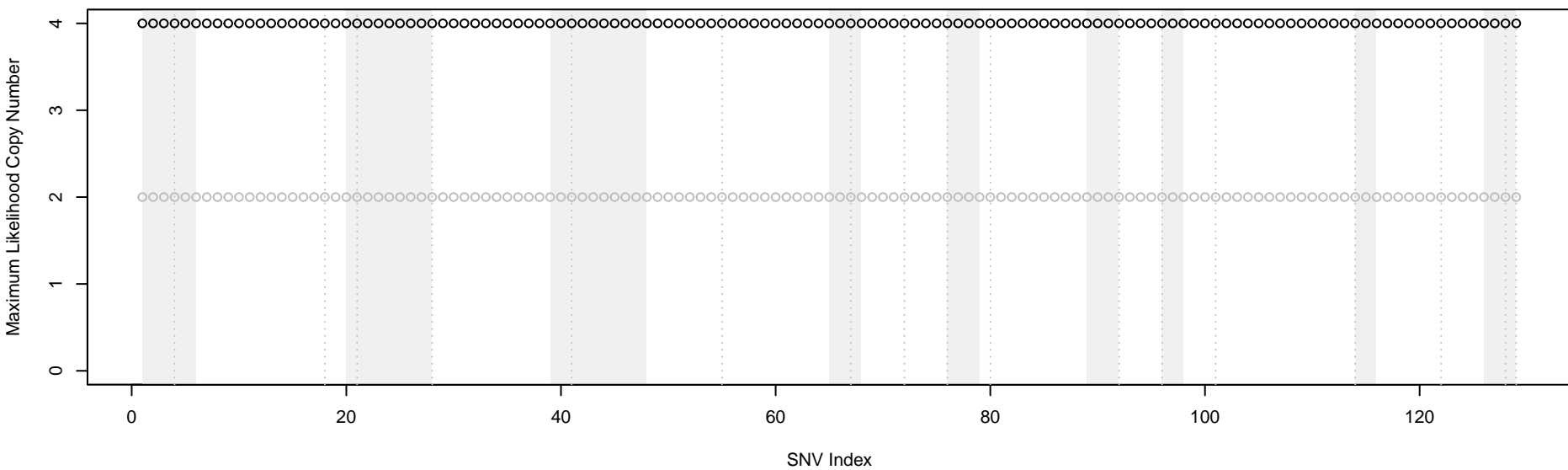
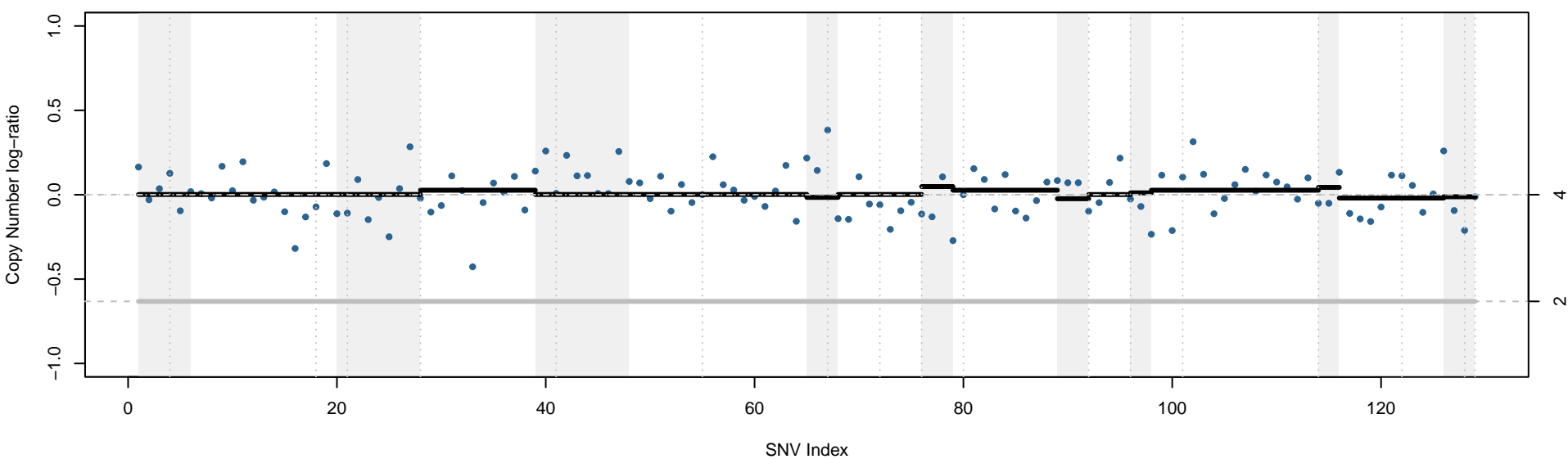
Purity: 0.55 Tumor ploidy: 3.999

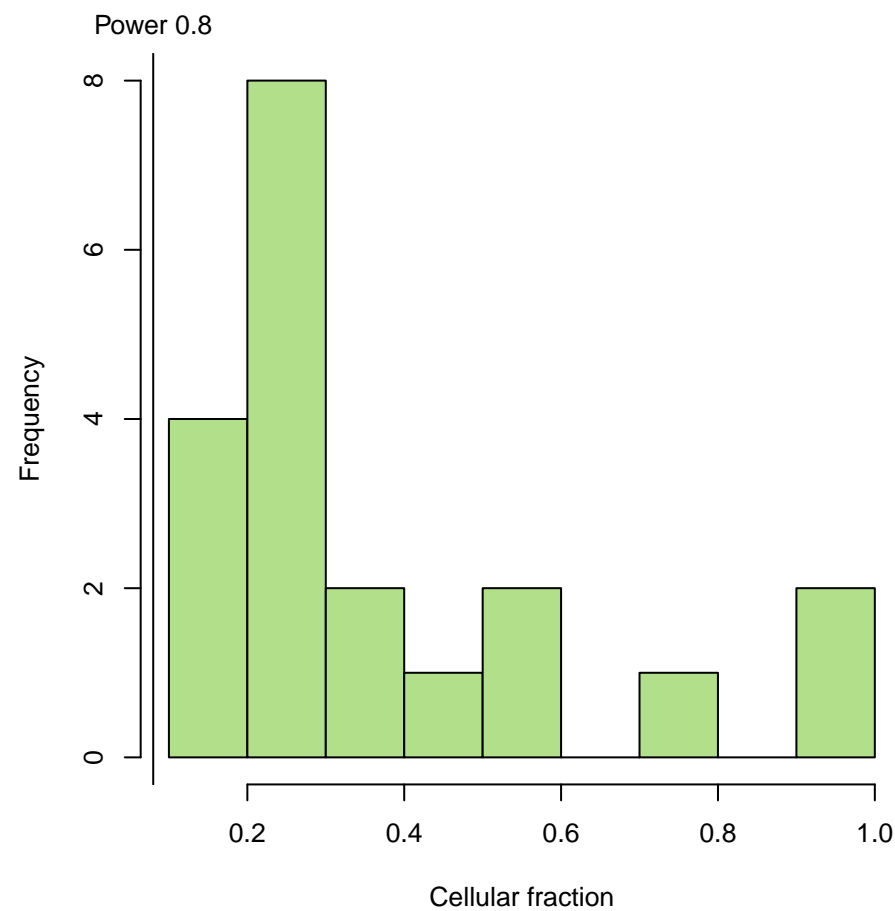
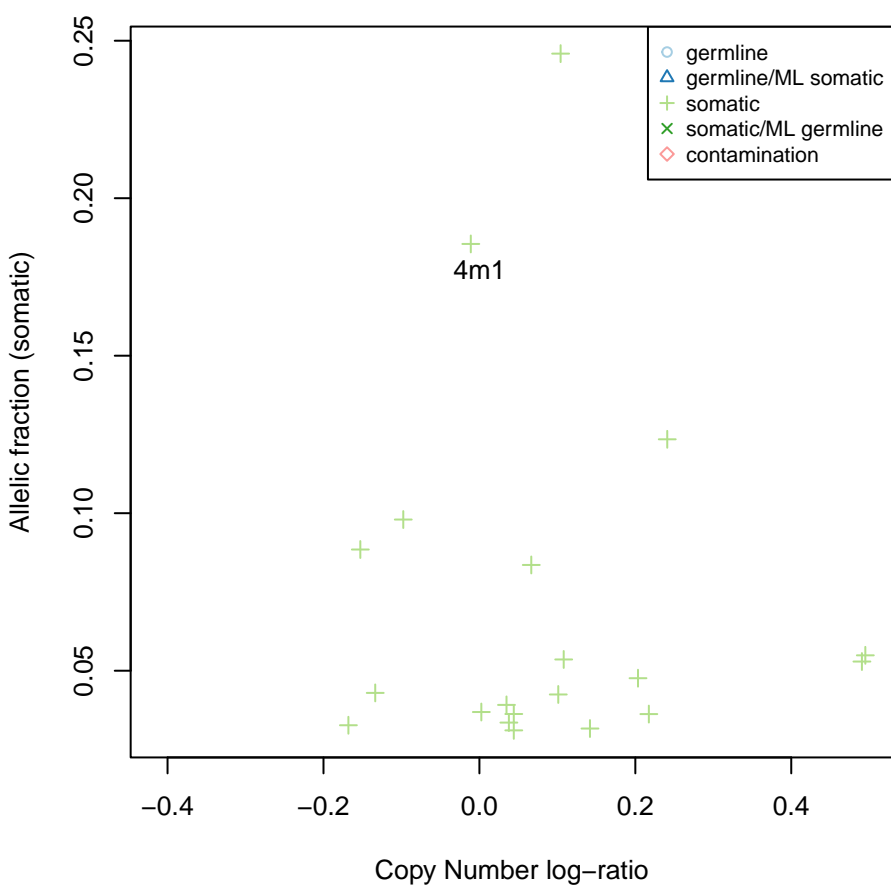
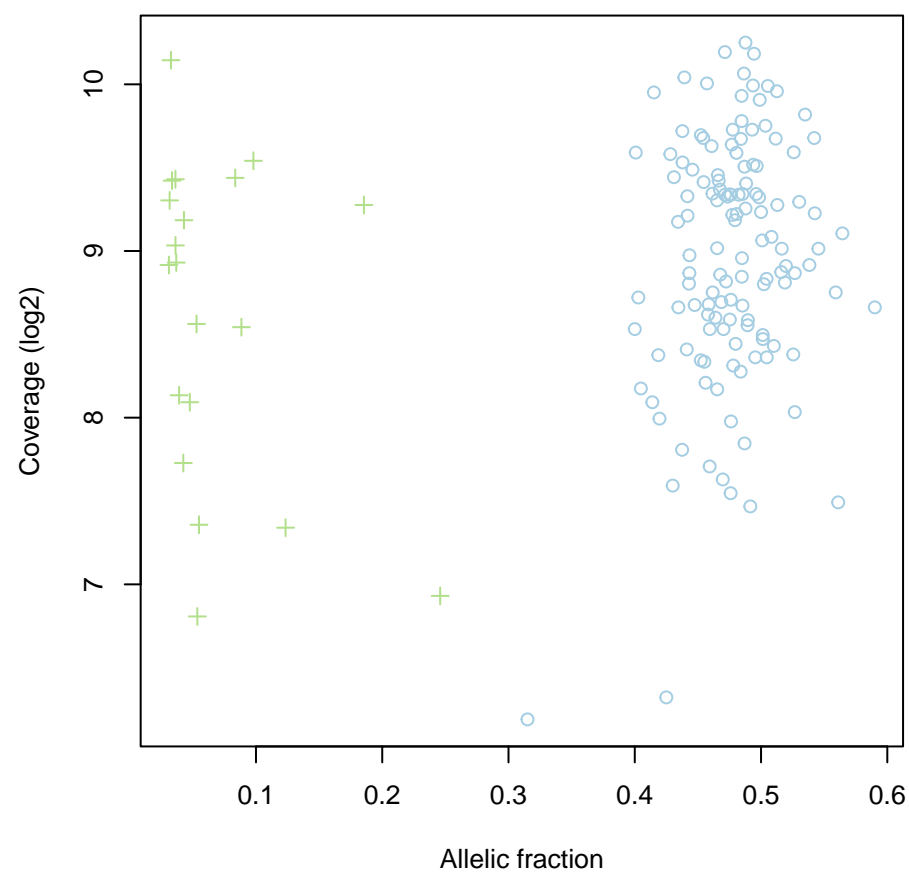
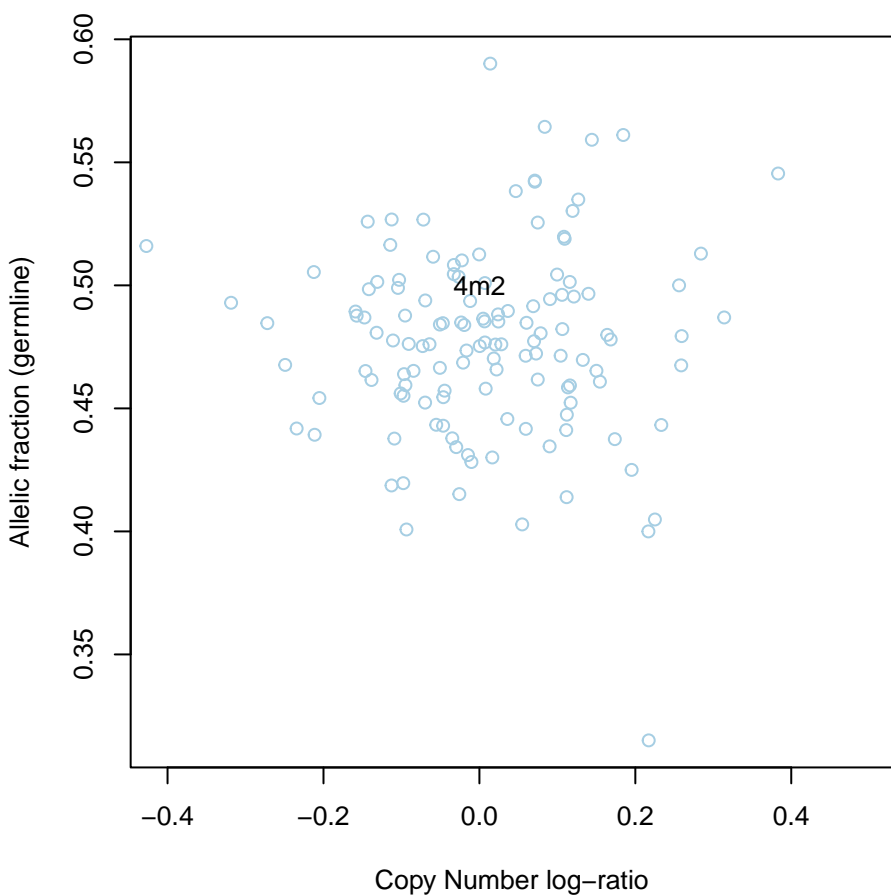


Purity: 0.55 Tumor ploidy: 3.999 SNV log-likelihood: 7.18 GoF: 91.7% Mean coverage: 549;494

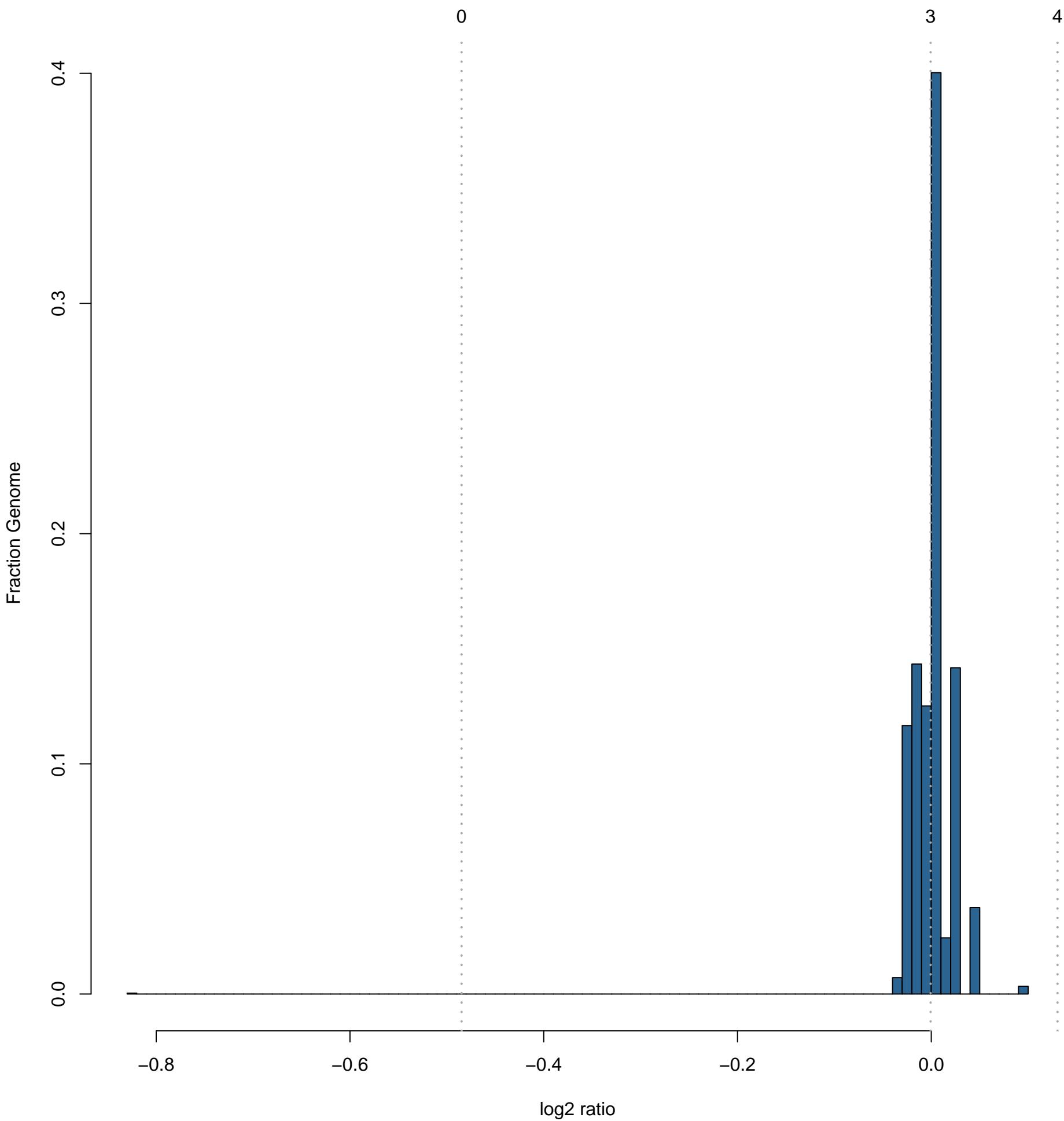


SCNA-fit log-likelihood: -4217.56

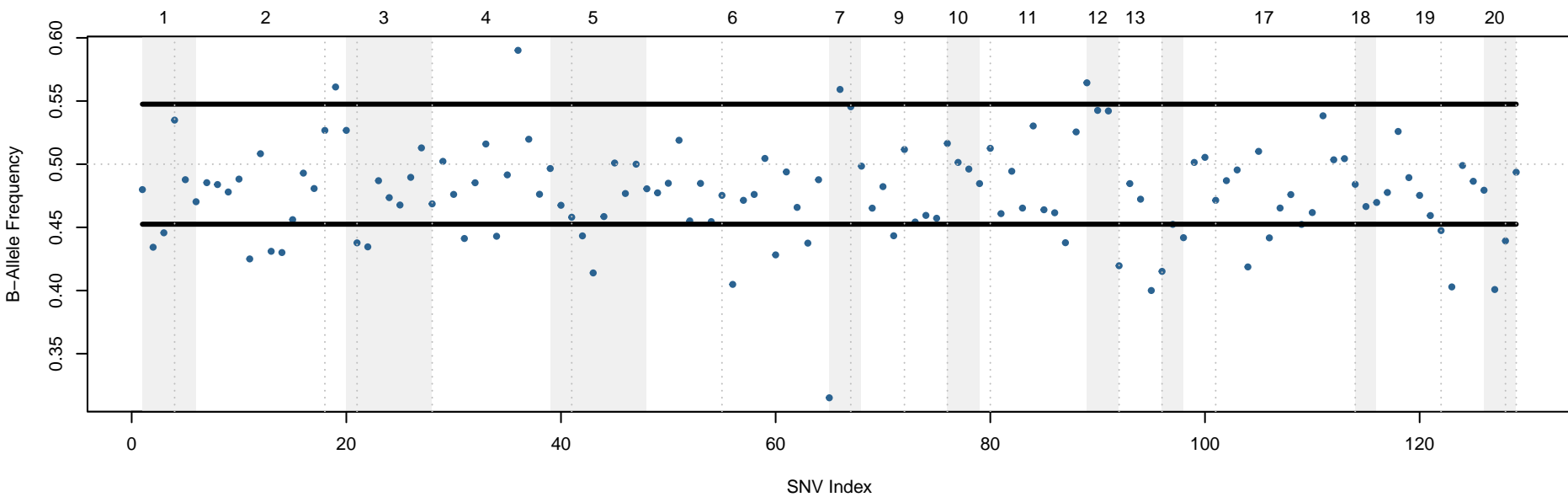




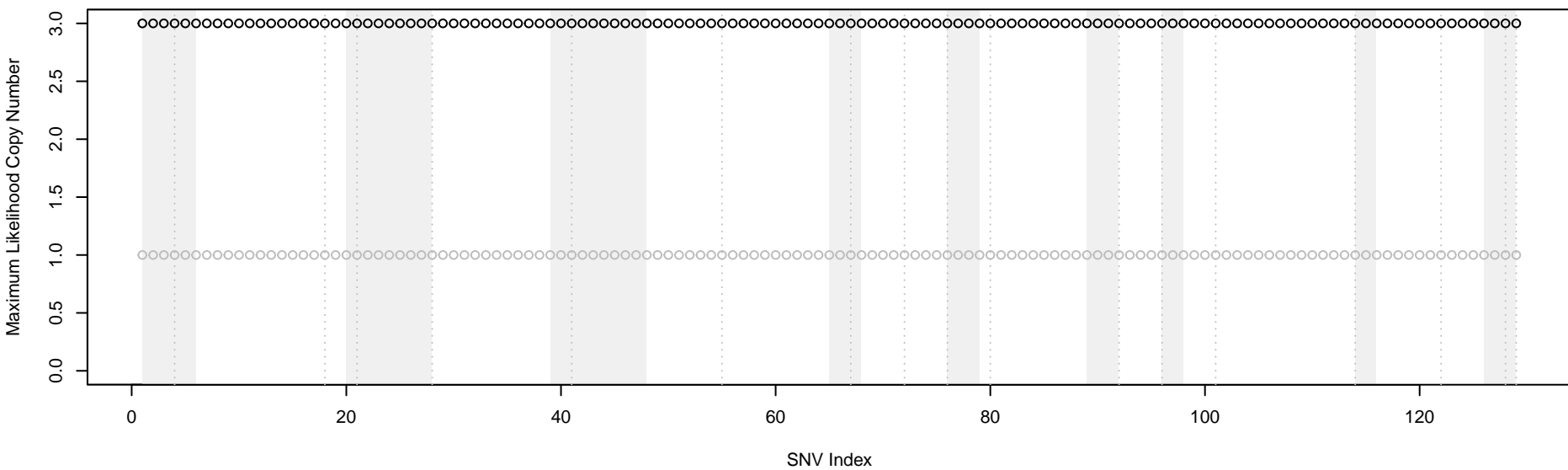
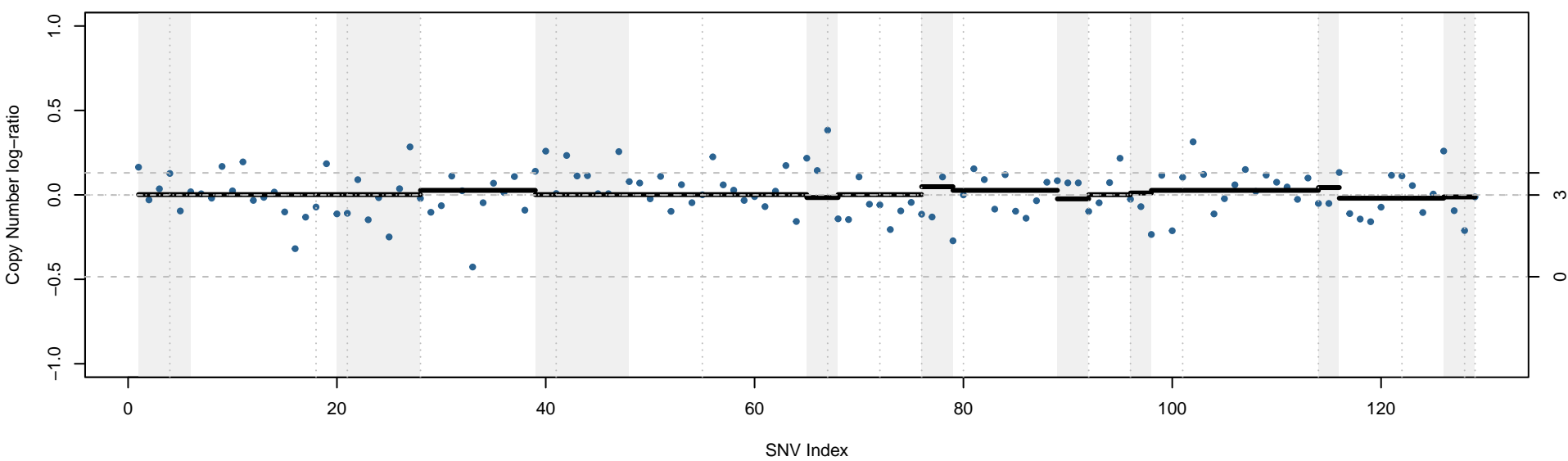
Purity: 0.21 Tumor ploidy: 3.005

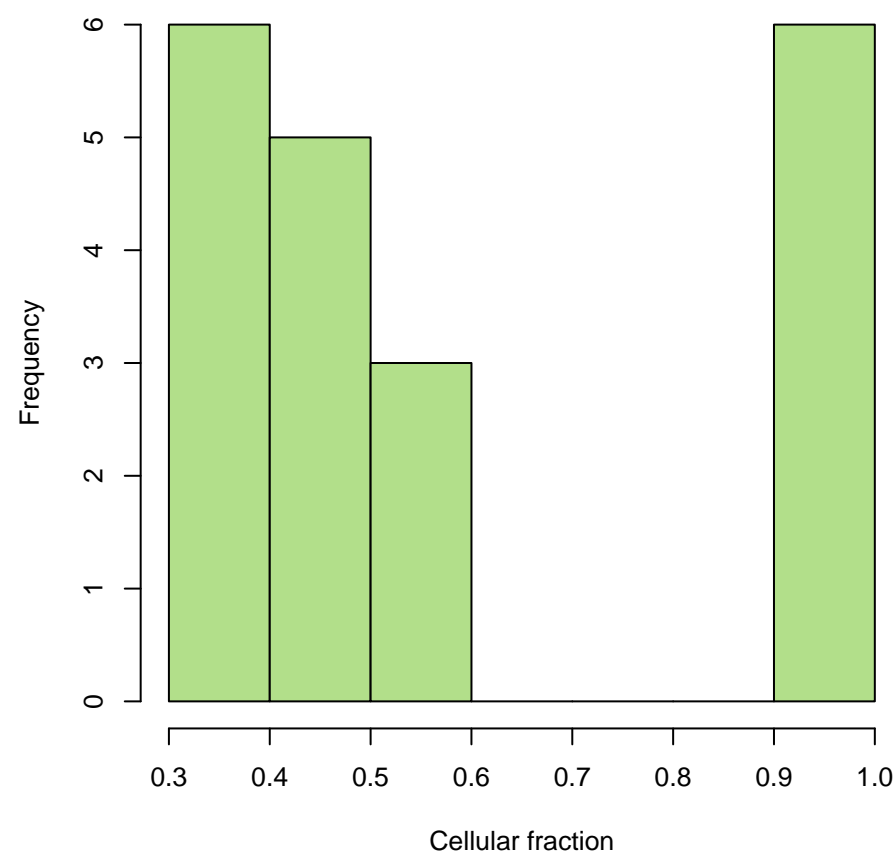
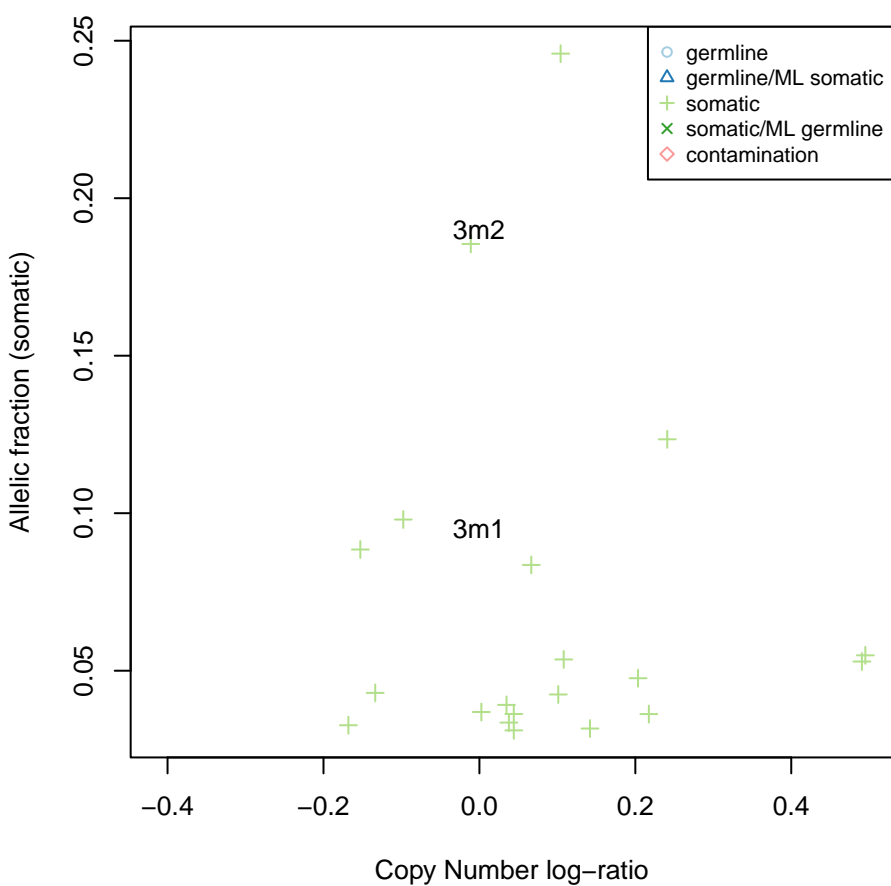
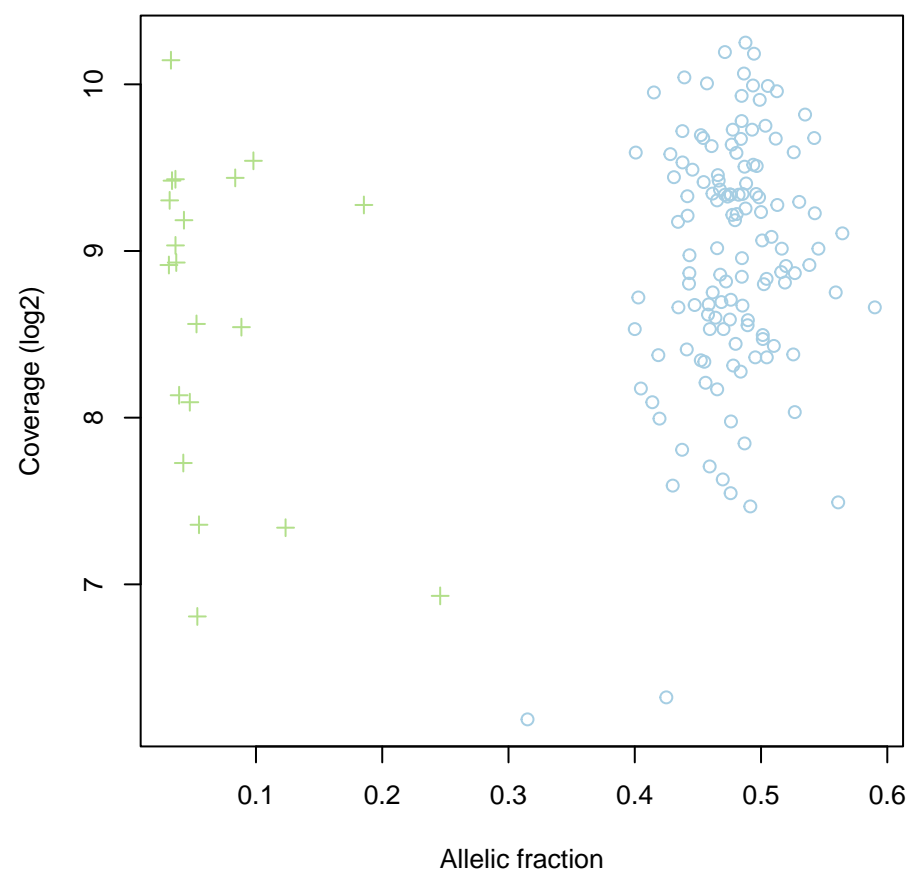
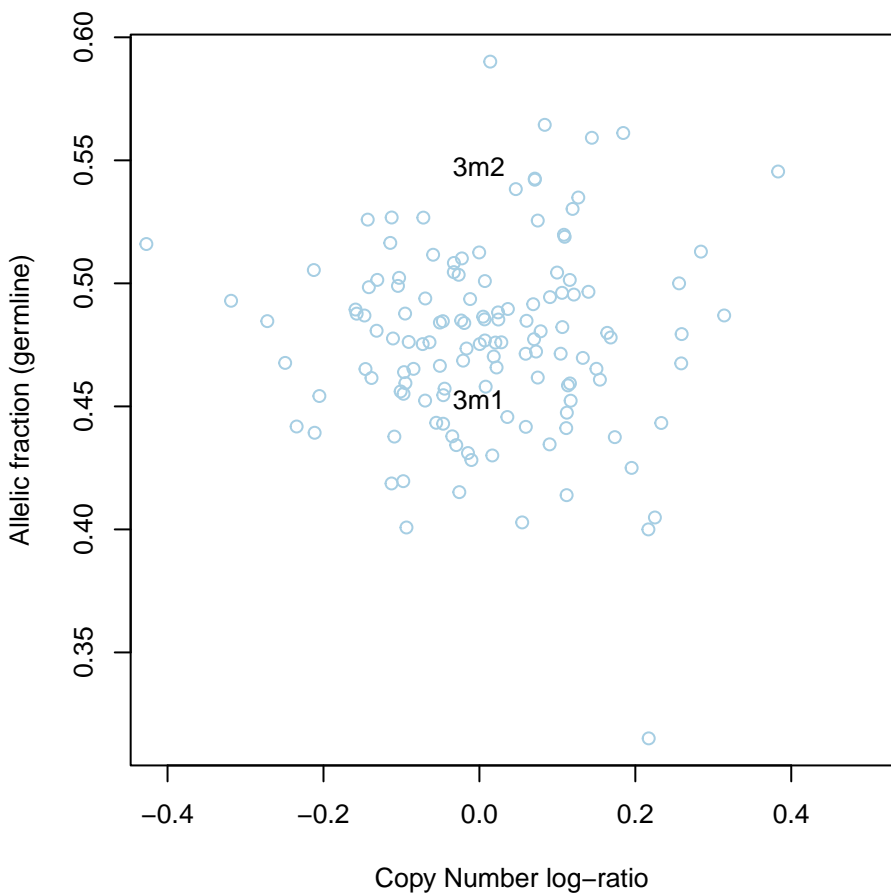


Purity: 0.21 Tumor ploidy: 3.005 SNV log-likelihood: -0.41 GoF: 97.1% Mean coverage: 549;494

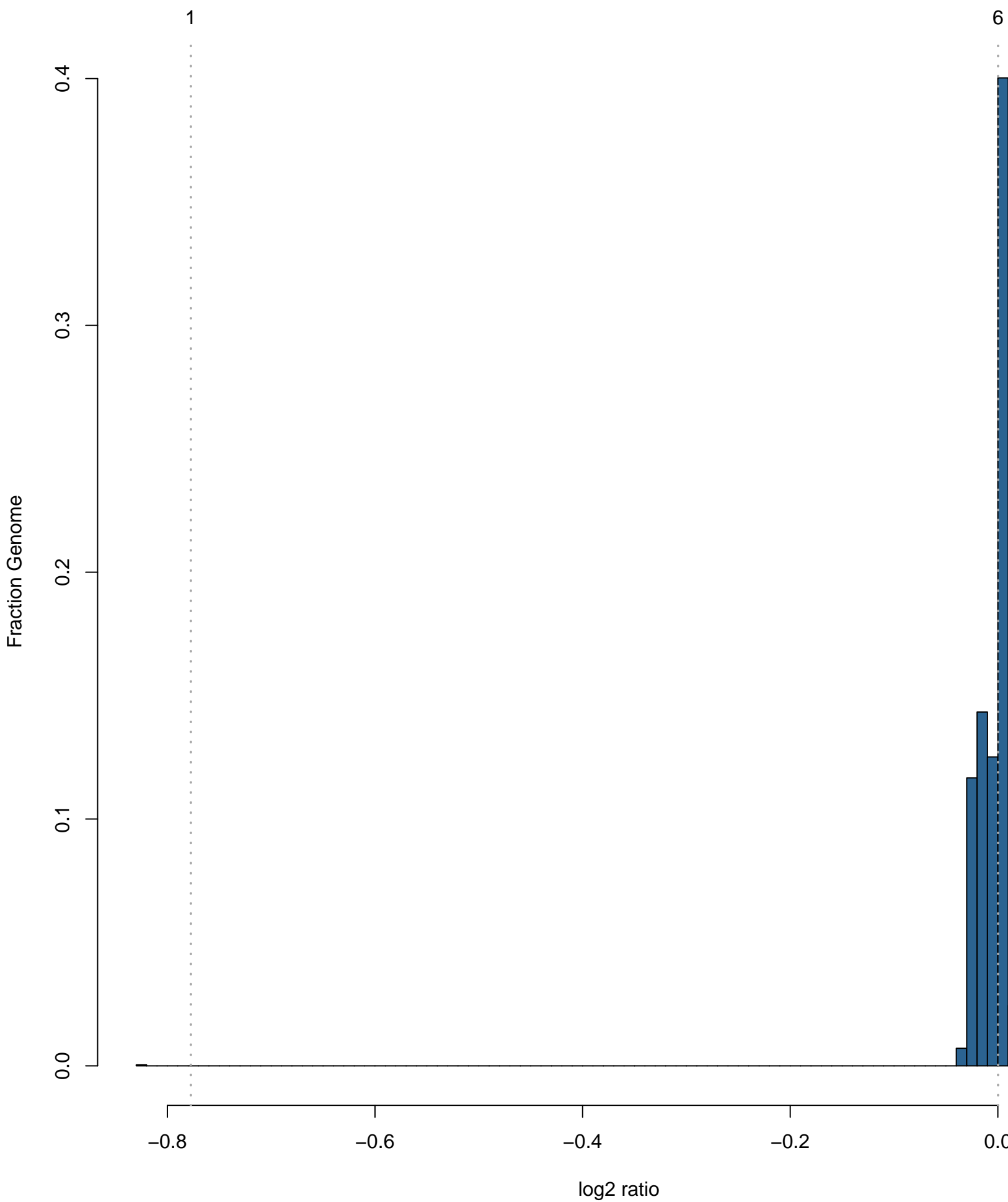


SCNA-fit log-likelihood: -4217.52



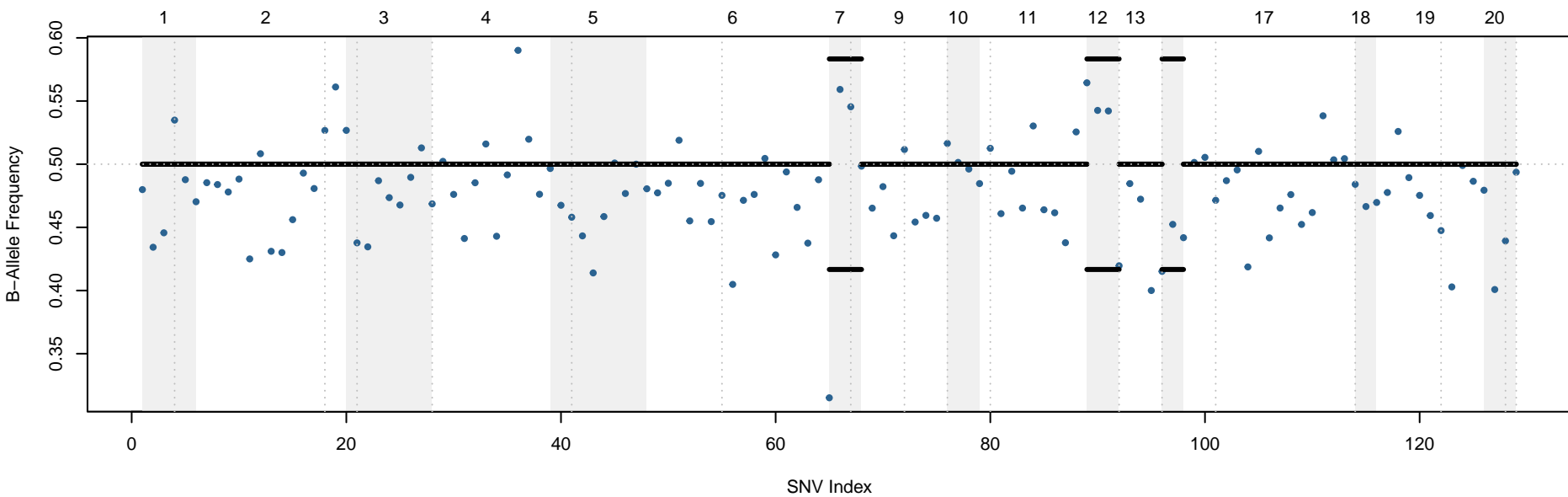


Purity: 0.25 Tumor ploidy: 5.998

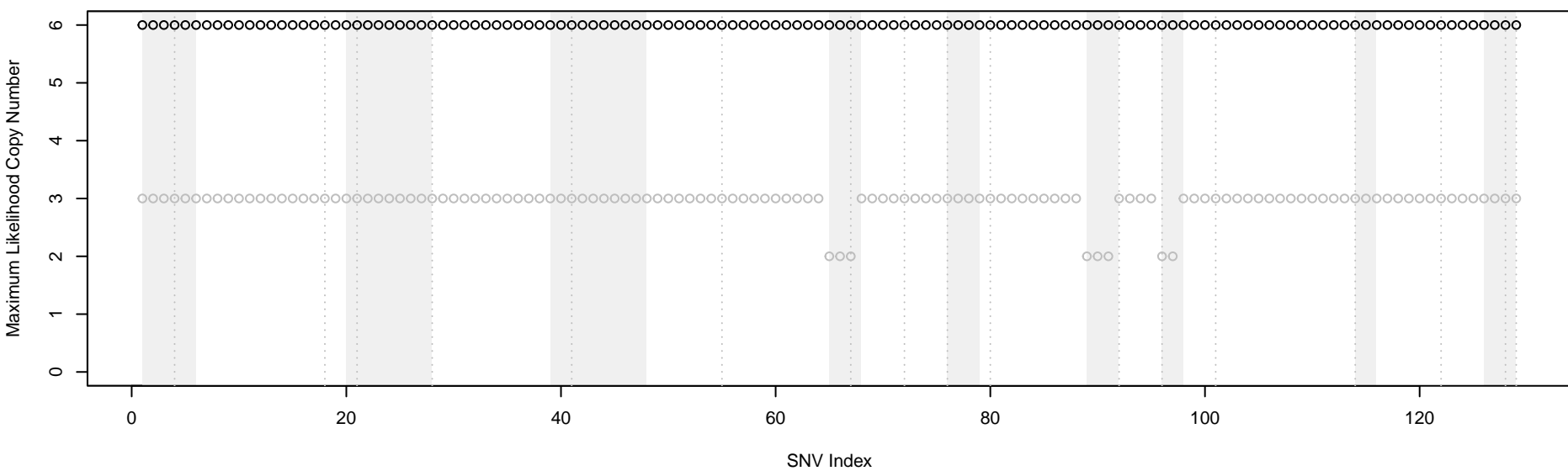
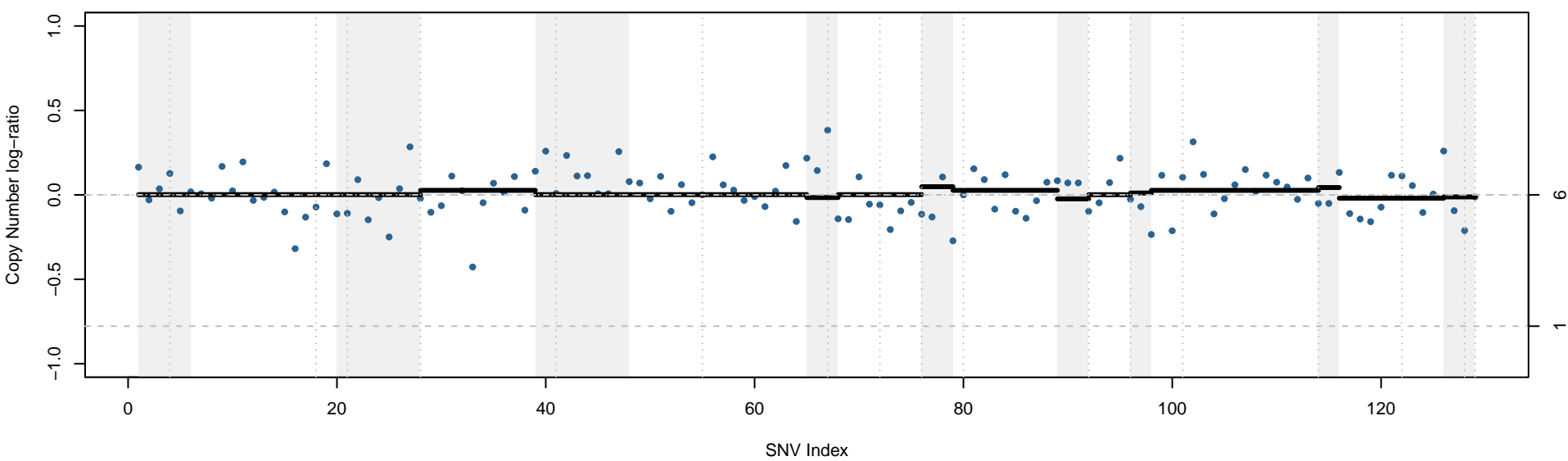


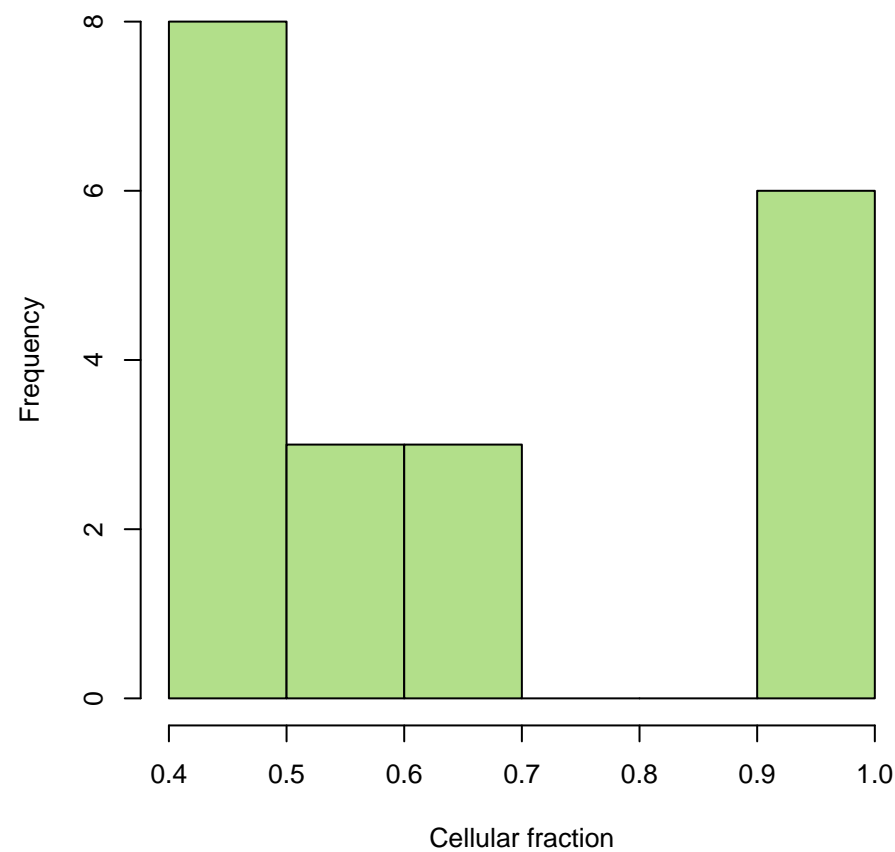
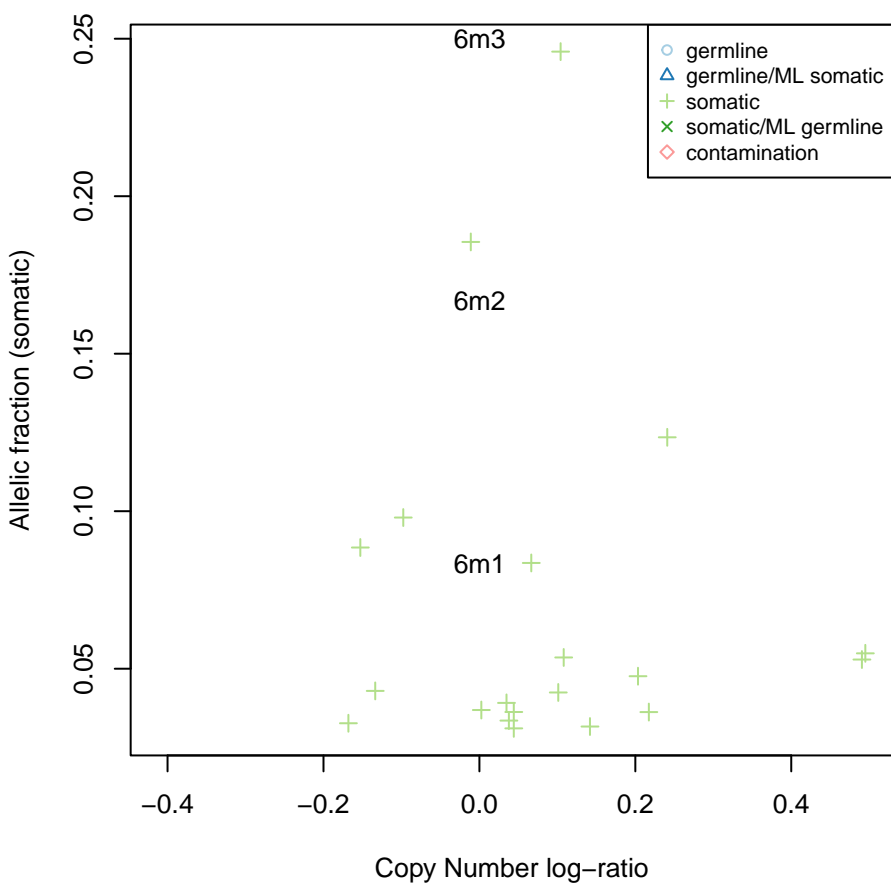
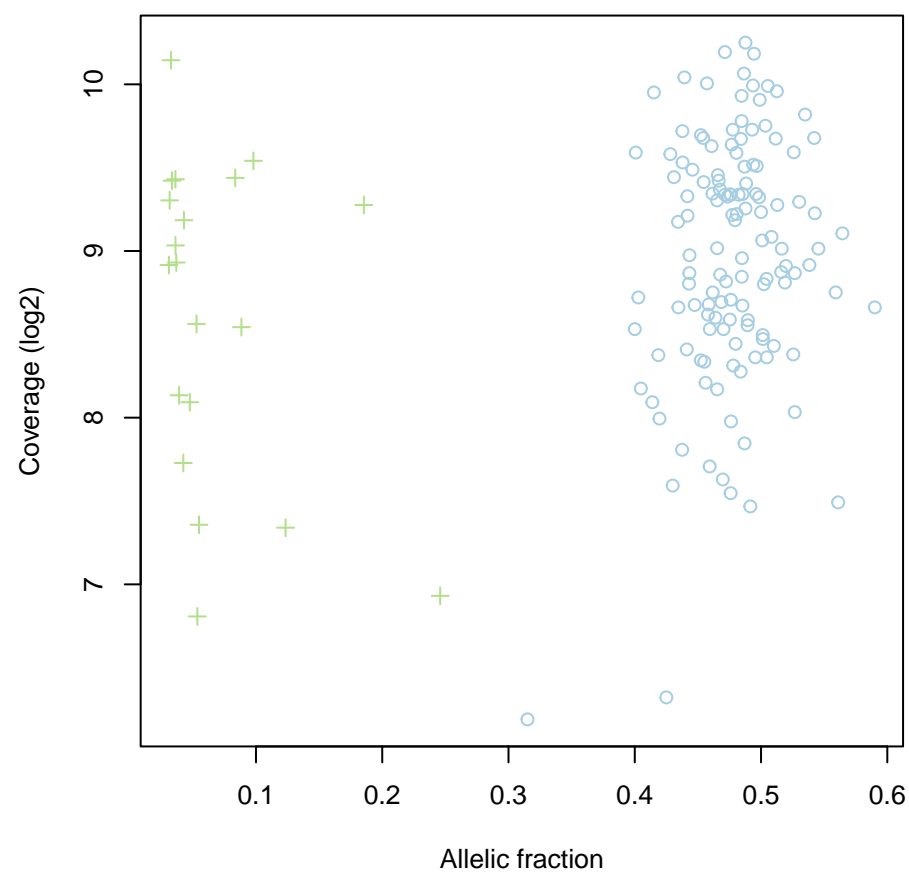
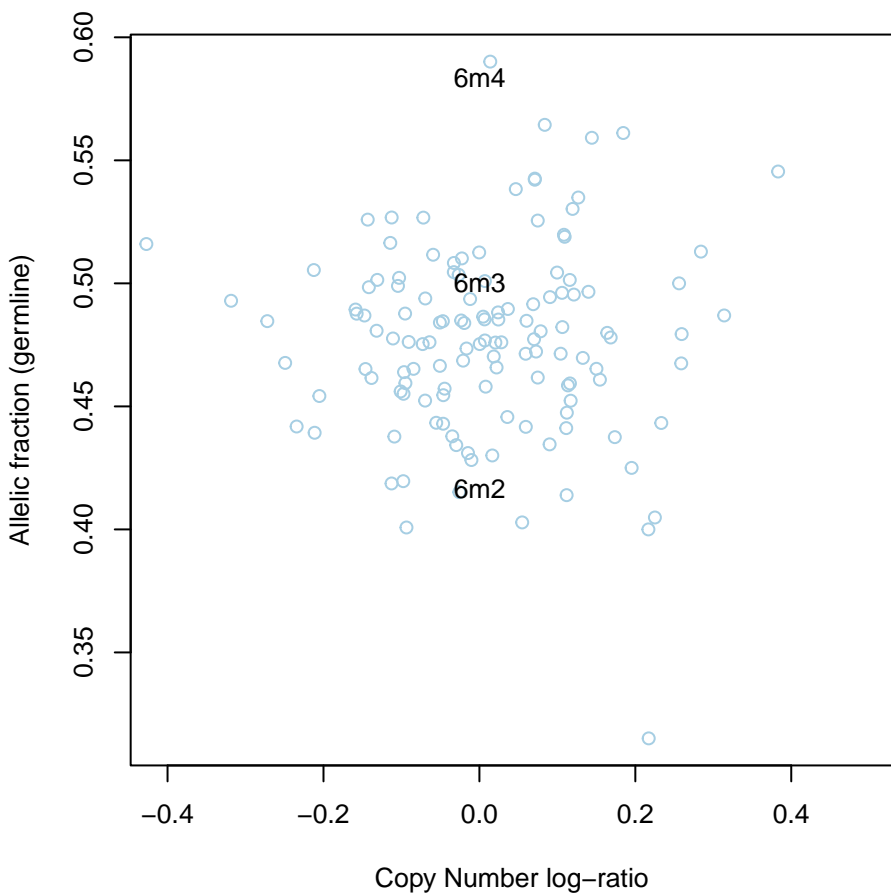


Purity: 0.25 Tumor ploidy: 5.998 SNV log-likelihood: -17.97 GoF: 96.9% Mean coverage: 549;494

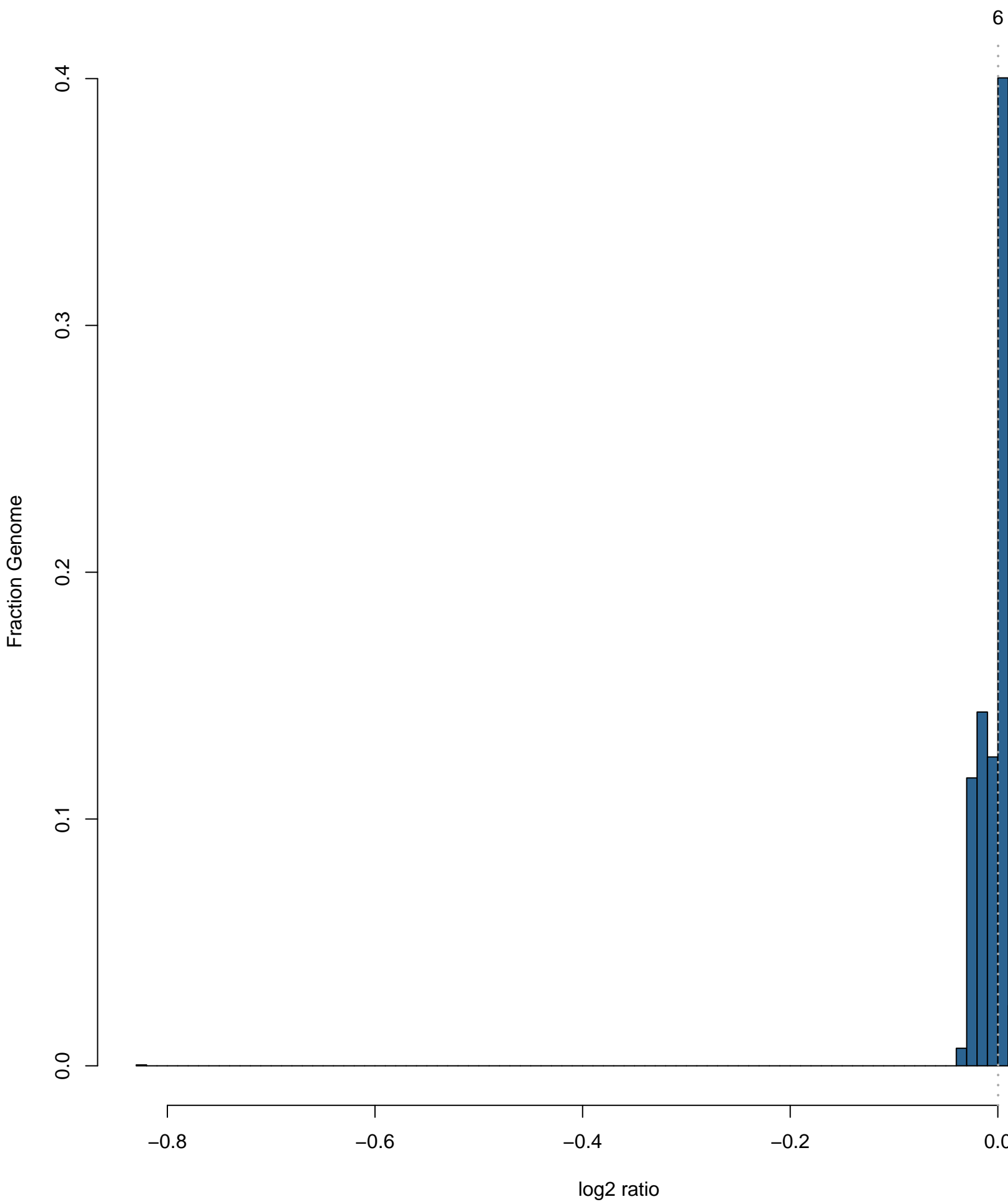


SCNA-fit log-likelihood: -4216.78

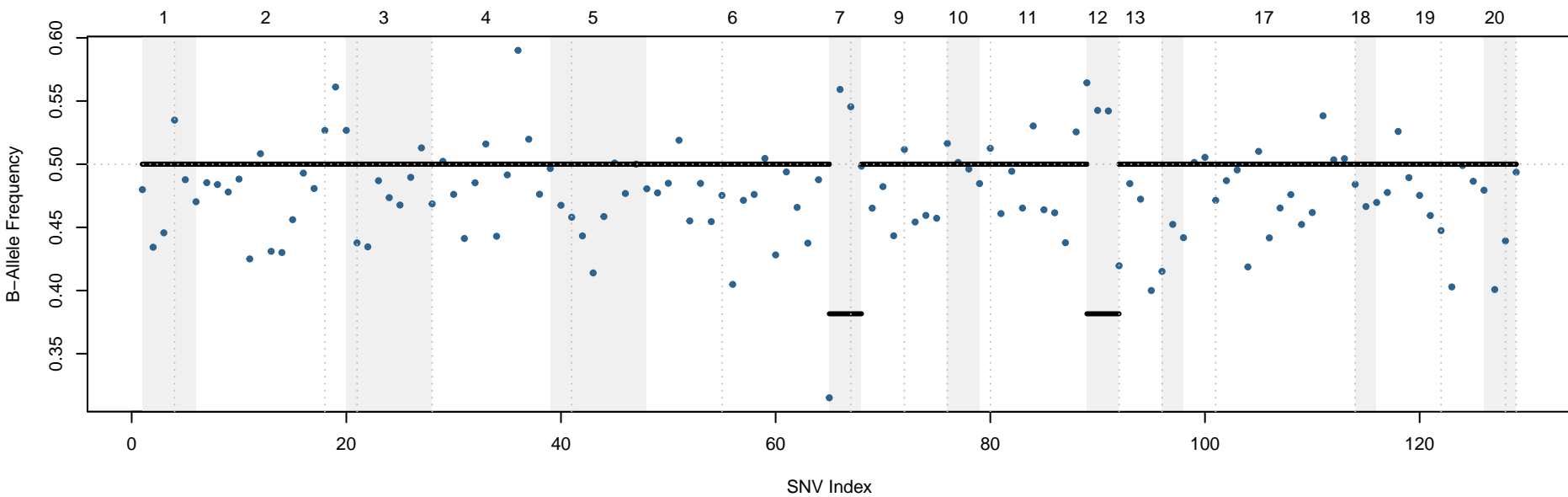




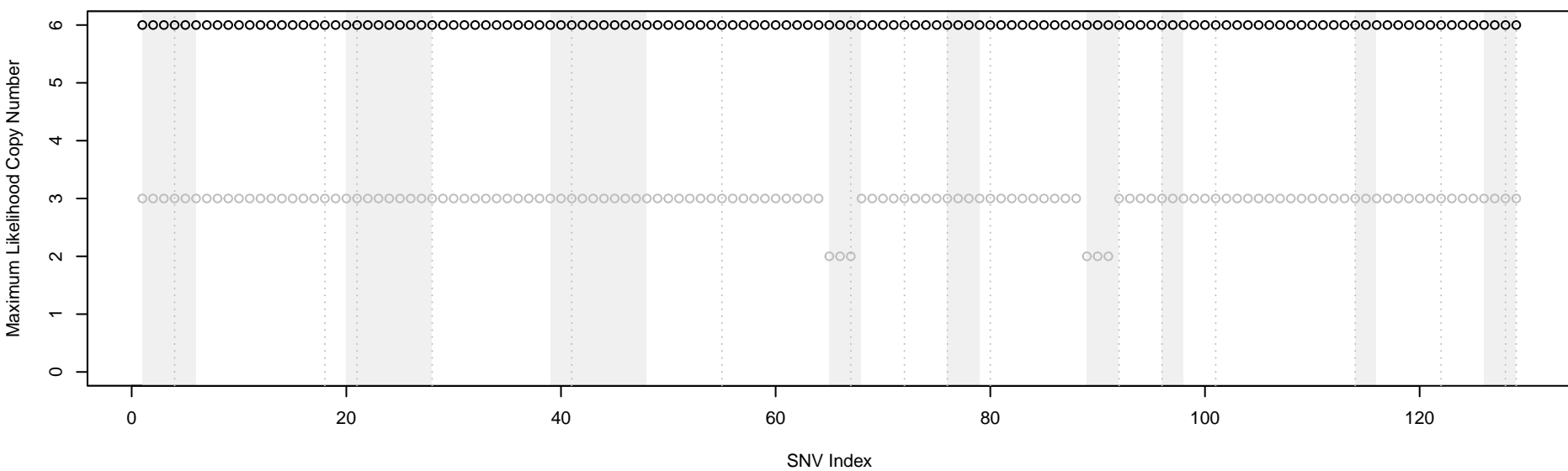
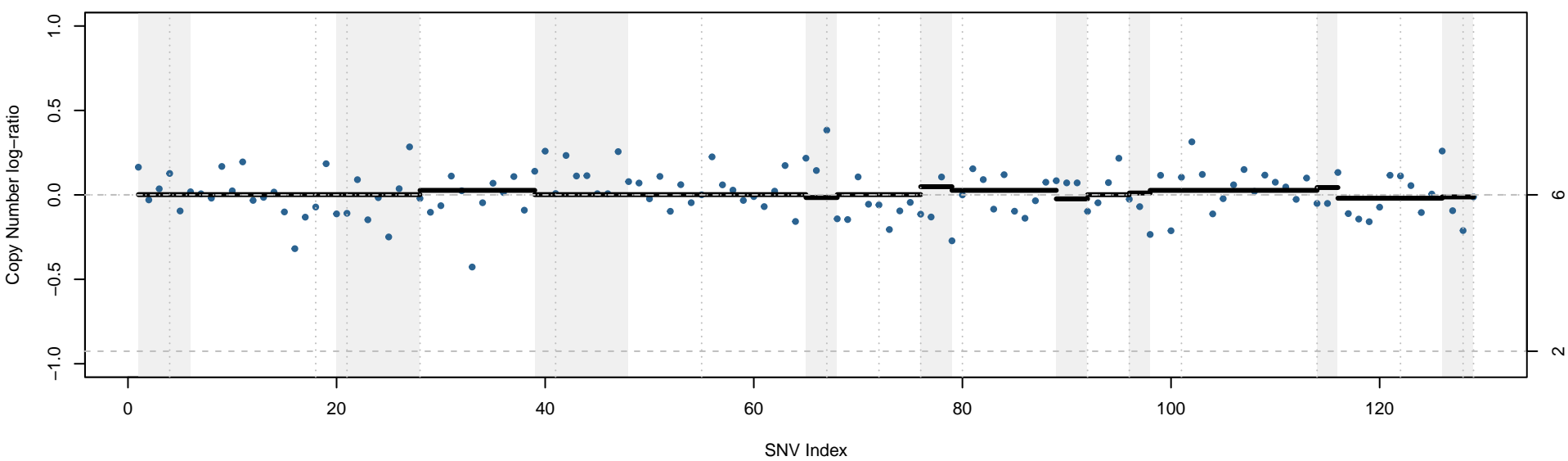
Purity: 0.45 Tumor ploidy: 5.998

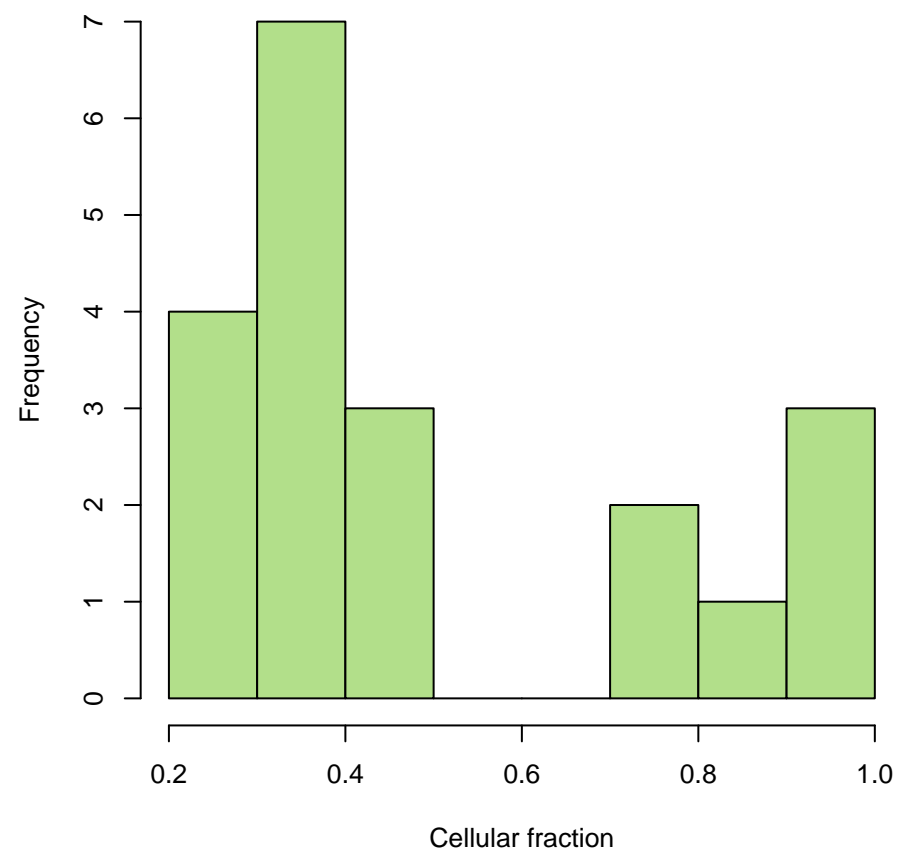
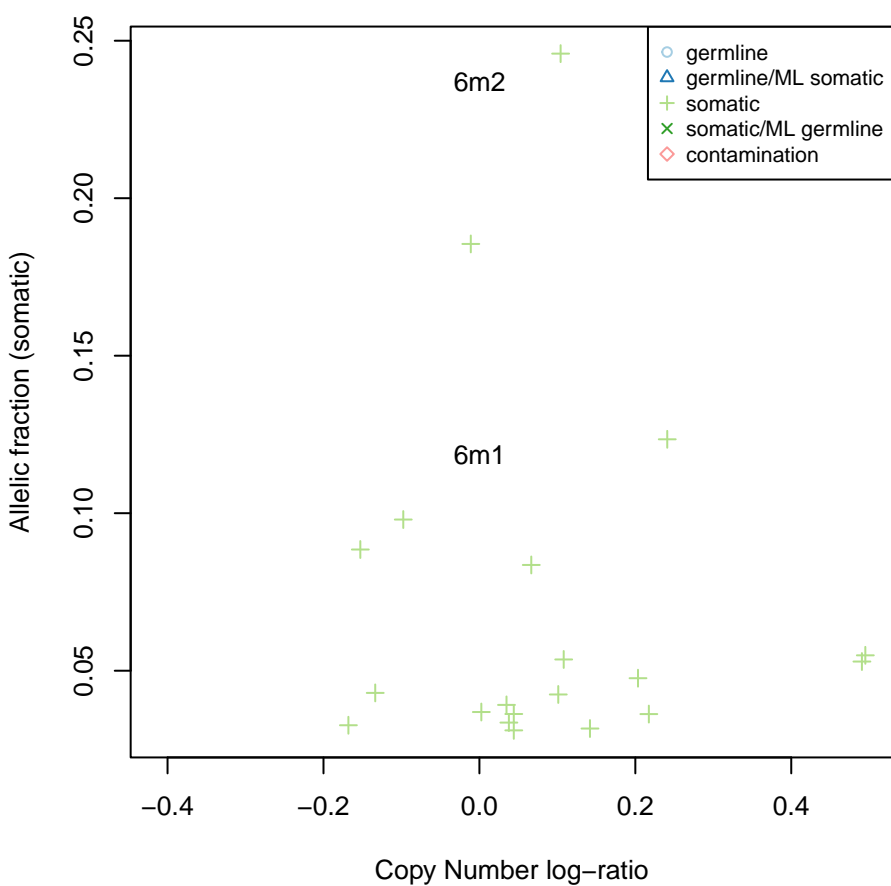
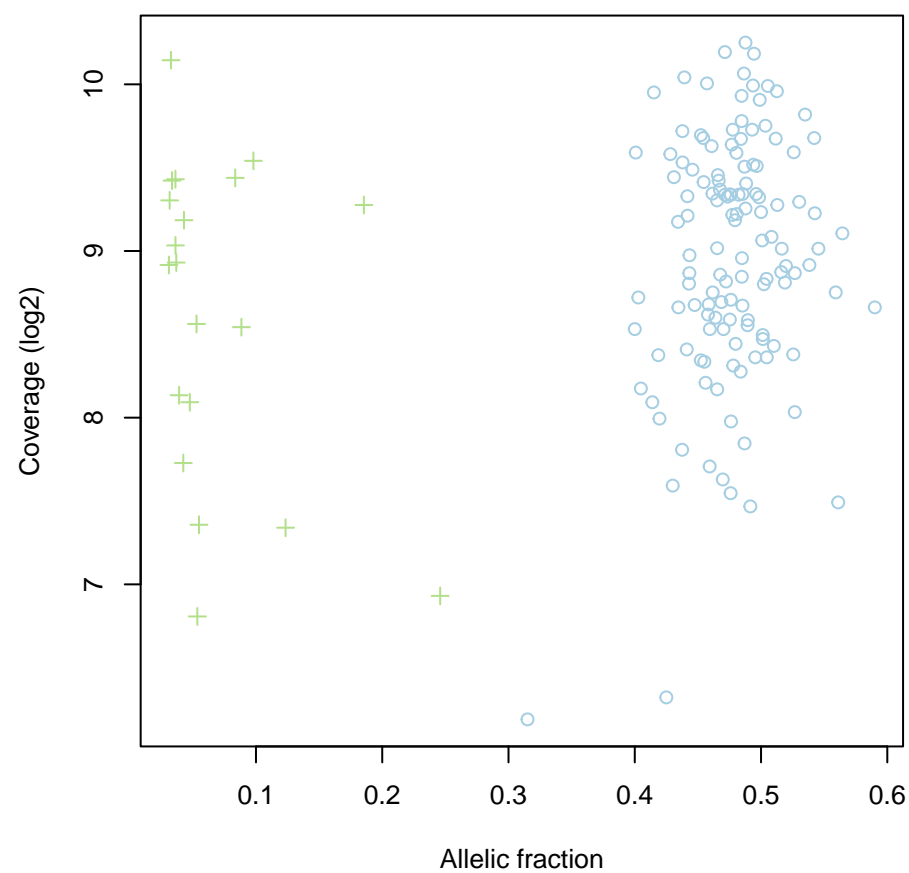
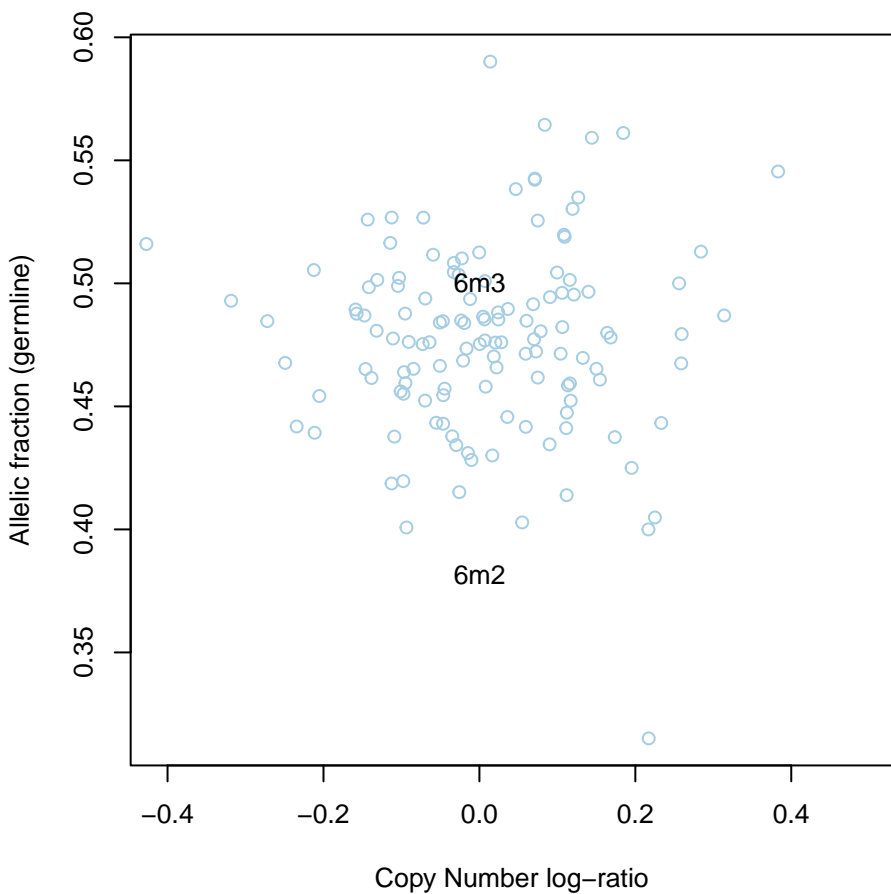


Purity: 0.45 Tumor ploidy: 5.998 SNV log-likelihood: -18.89 GoF: 95.8% Mean coverage: 549;494

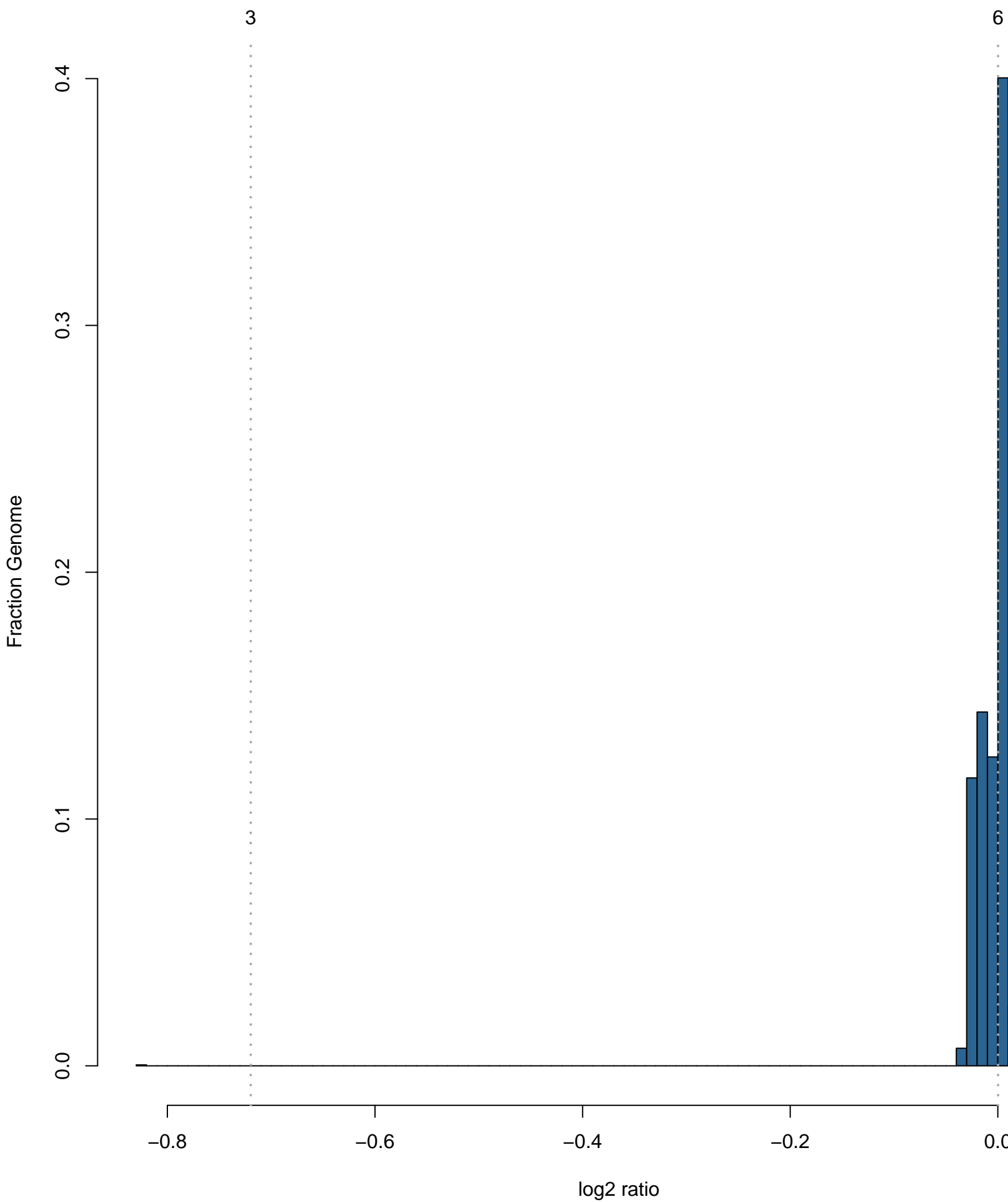


SCNA-fit log-likelihood: -4216.99

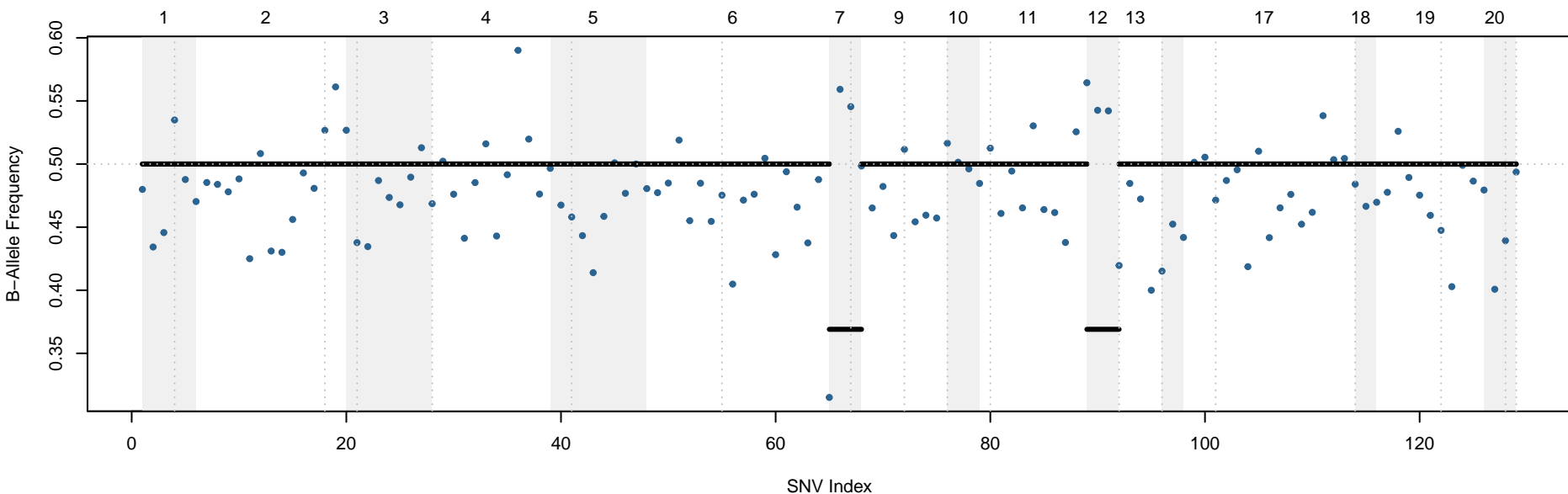




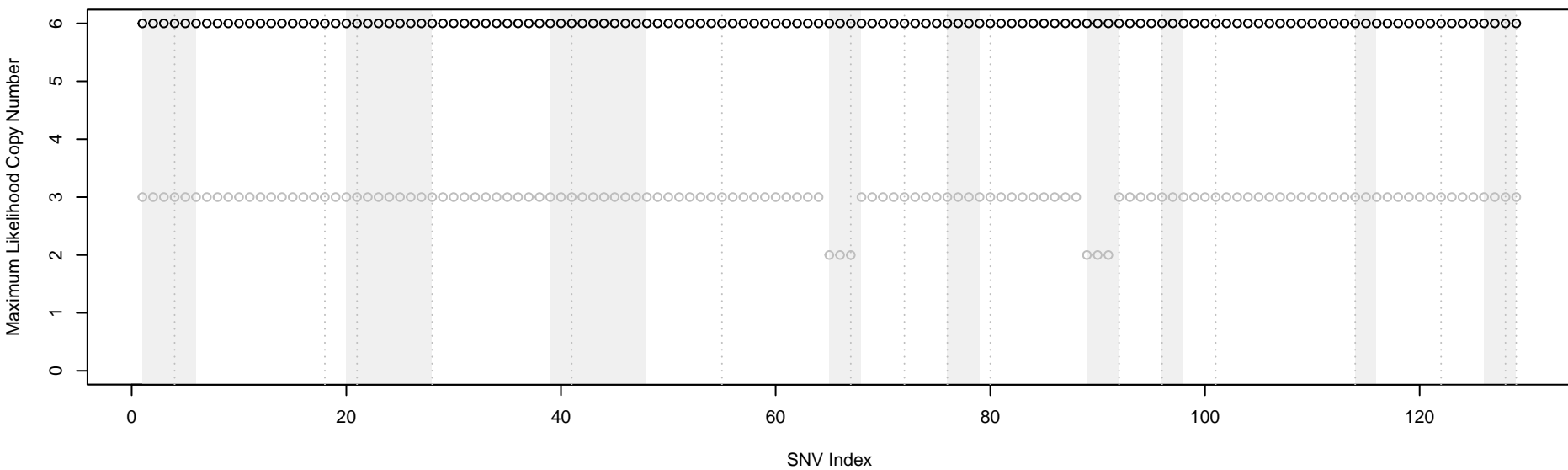
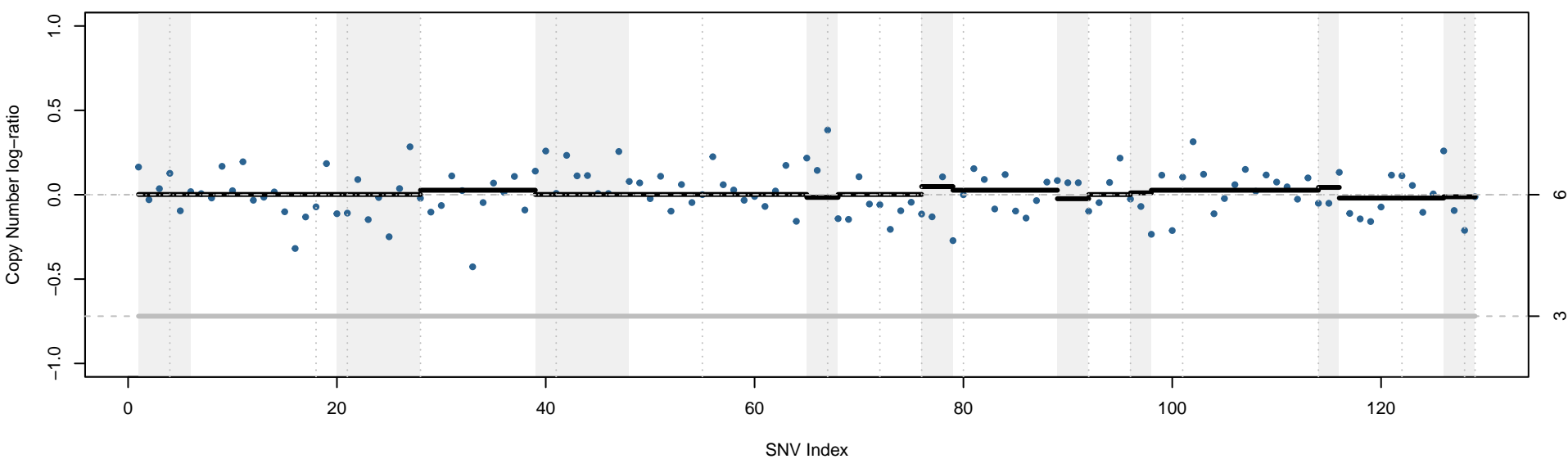
Purity: 0.55 Tumor ploidy: 5.999

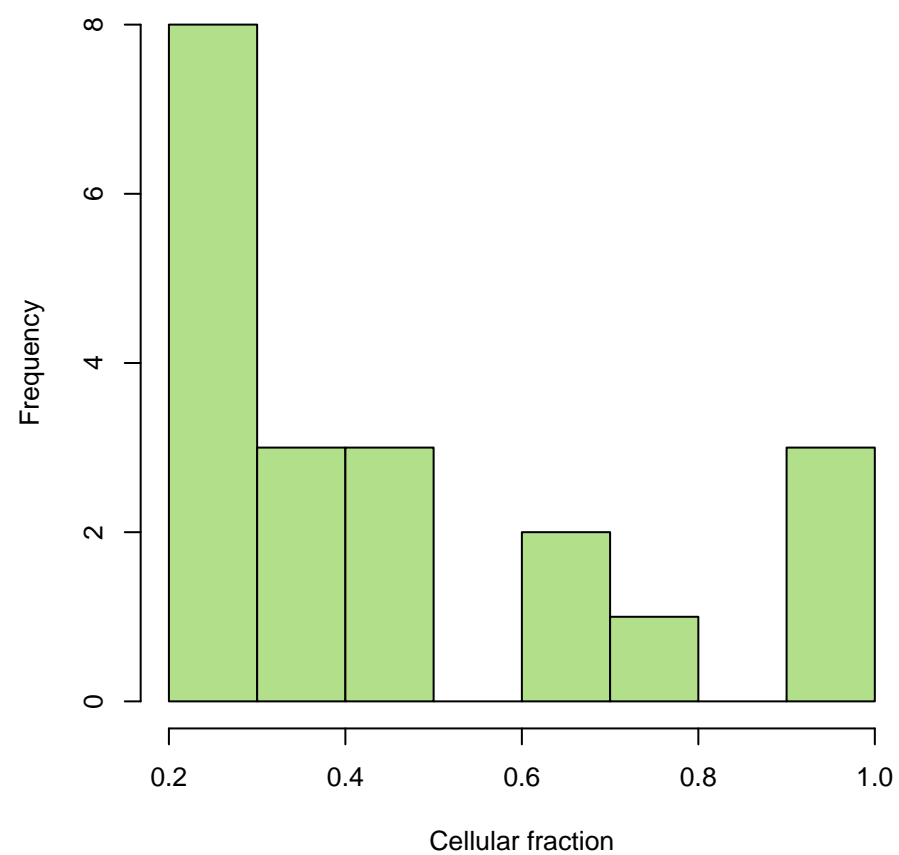
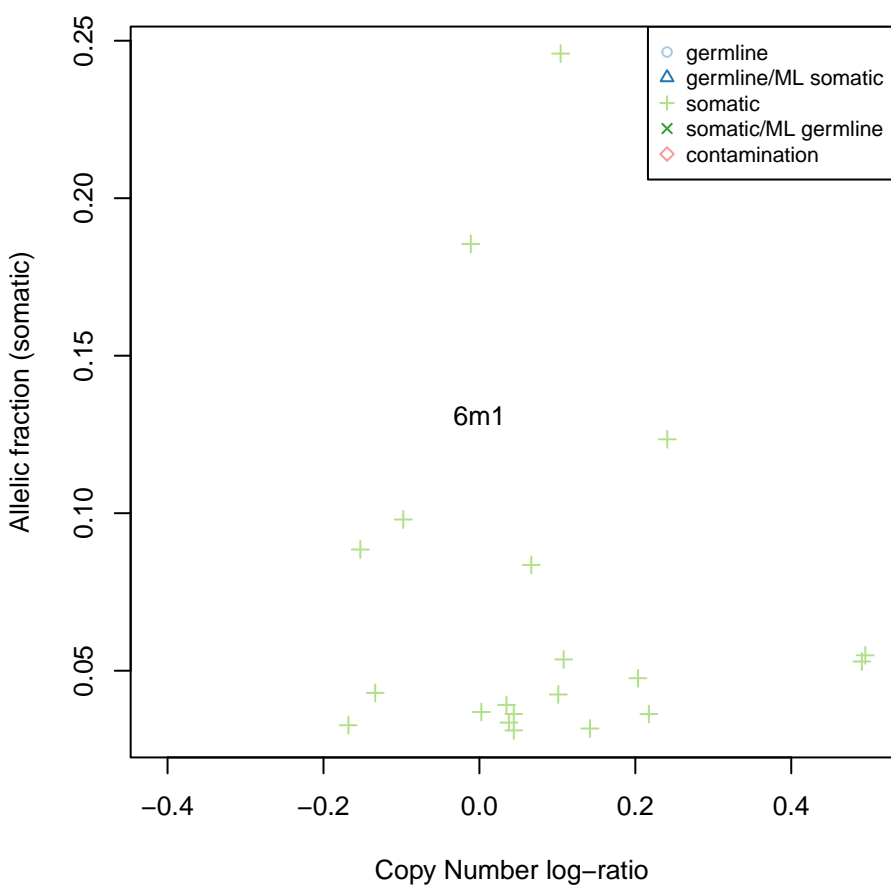
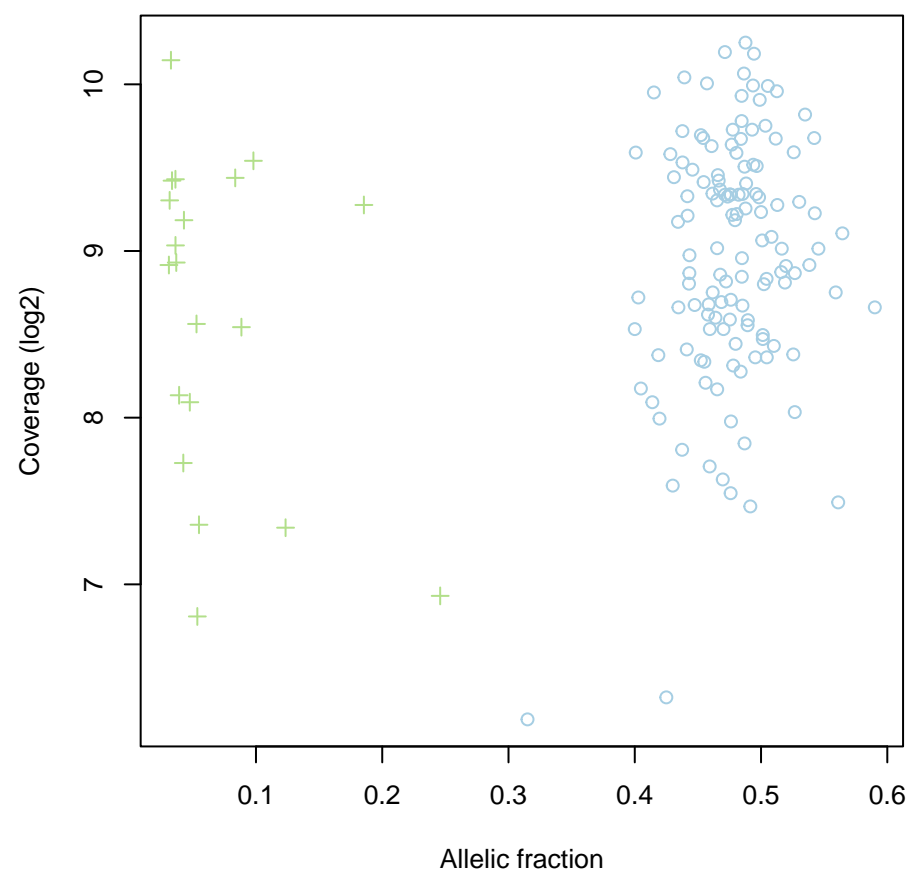
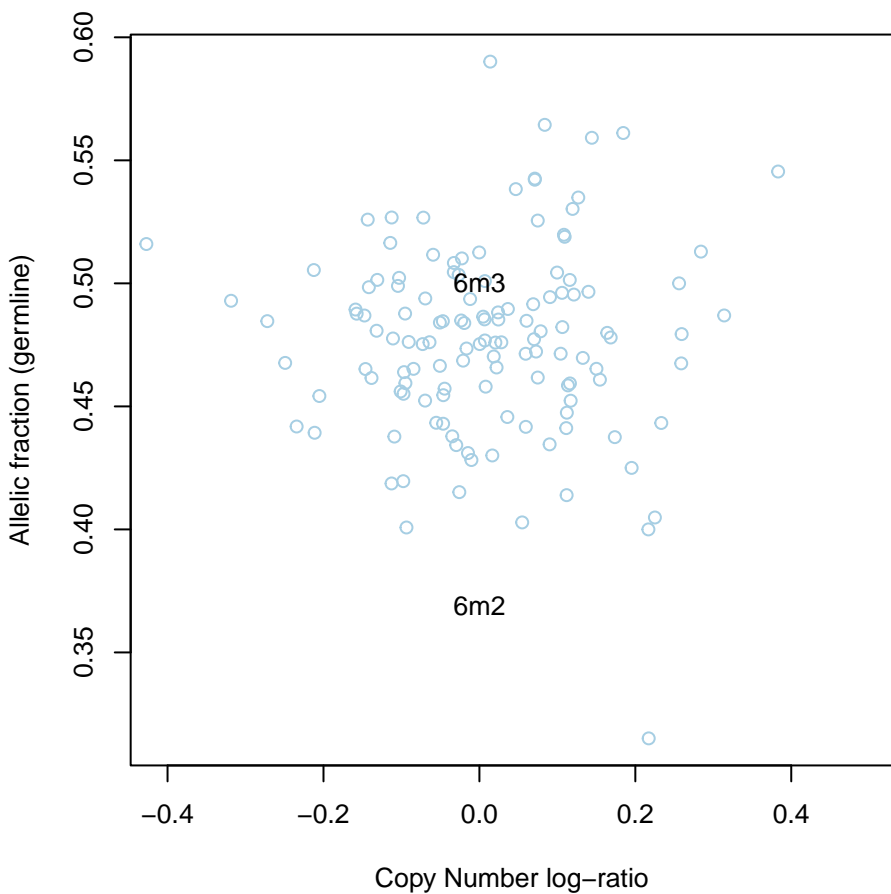


Purity: 0.55 Tumor ploidy: 5.999 SNV log-likelihood: -28.09 GoF: 95.1% Mean coverage: 549;494



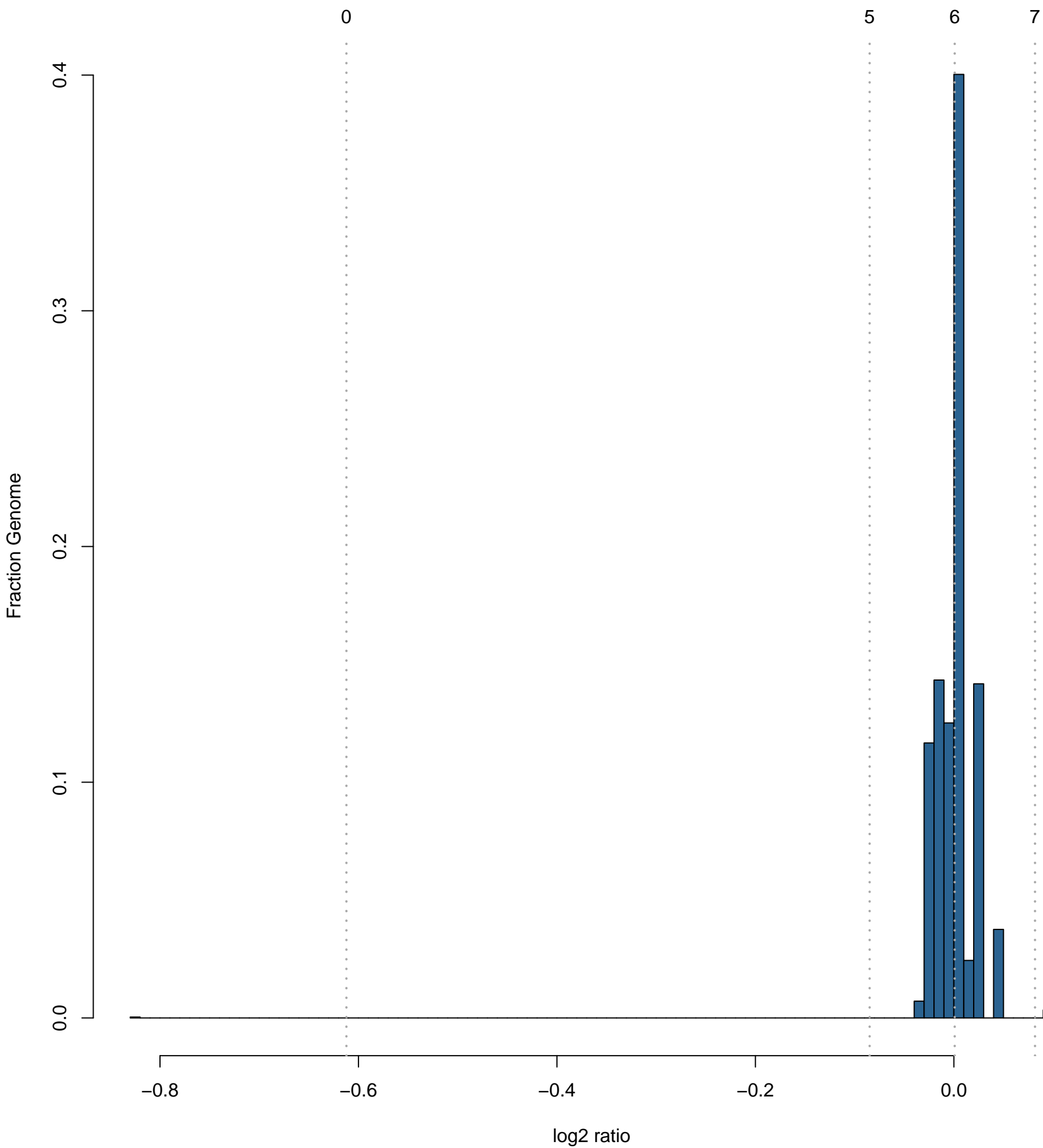
SCNA-fit log-likelihood: -4216.99



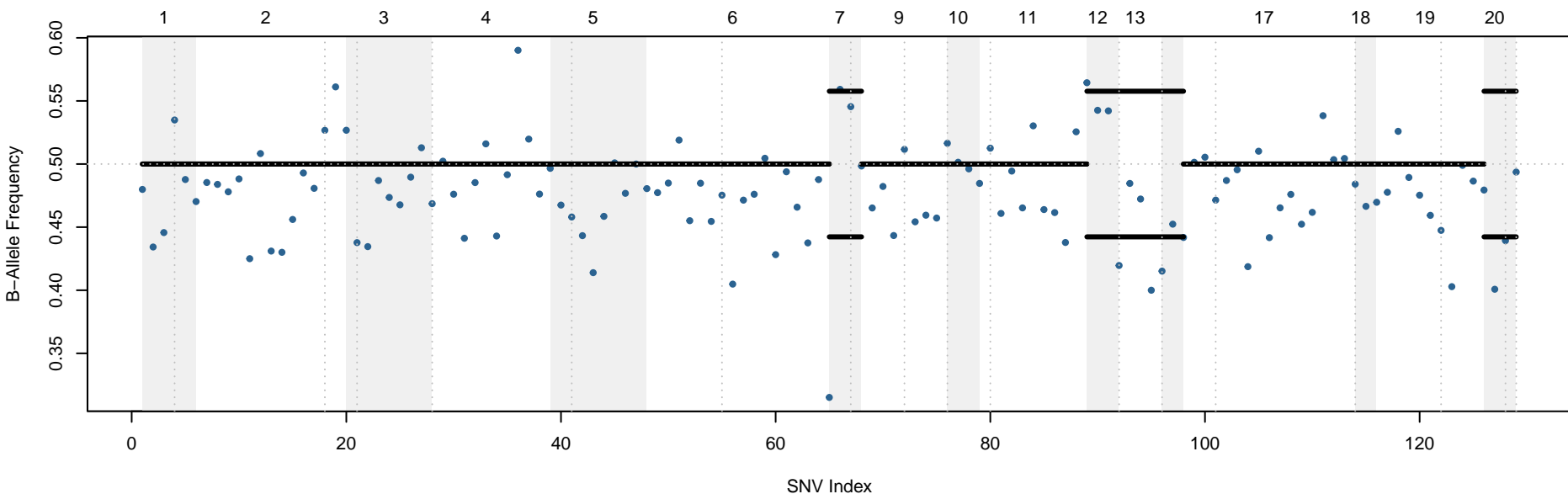




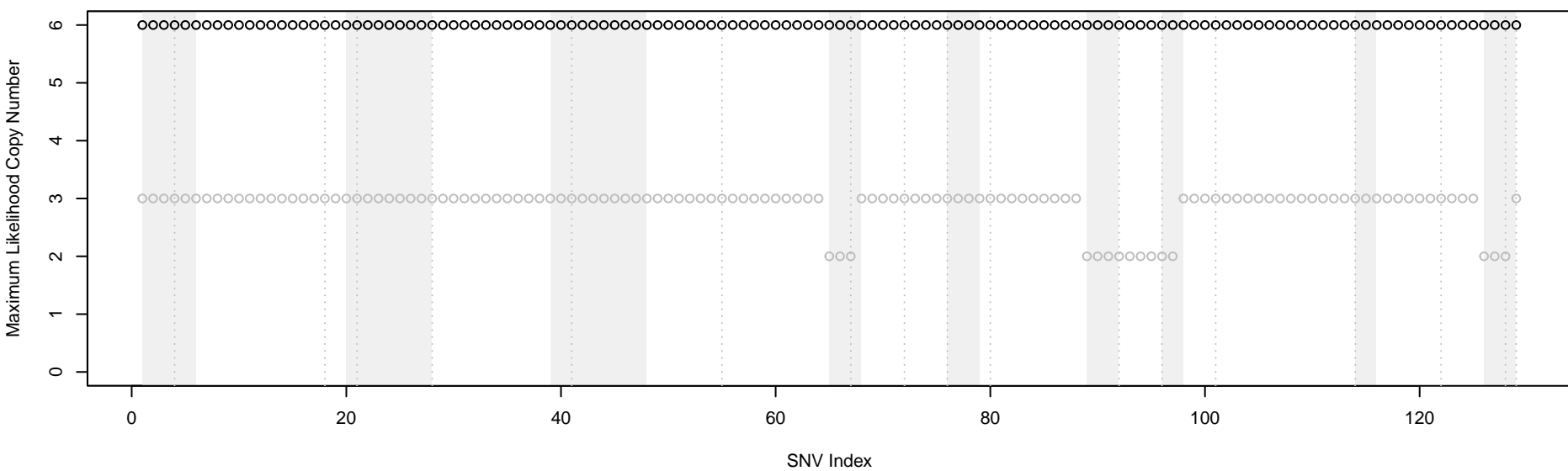
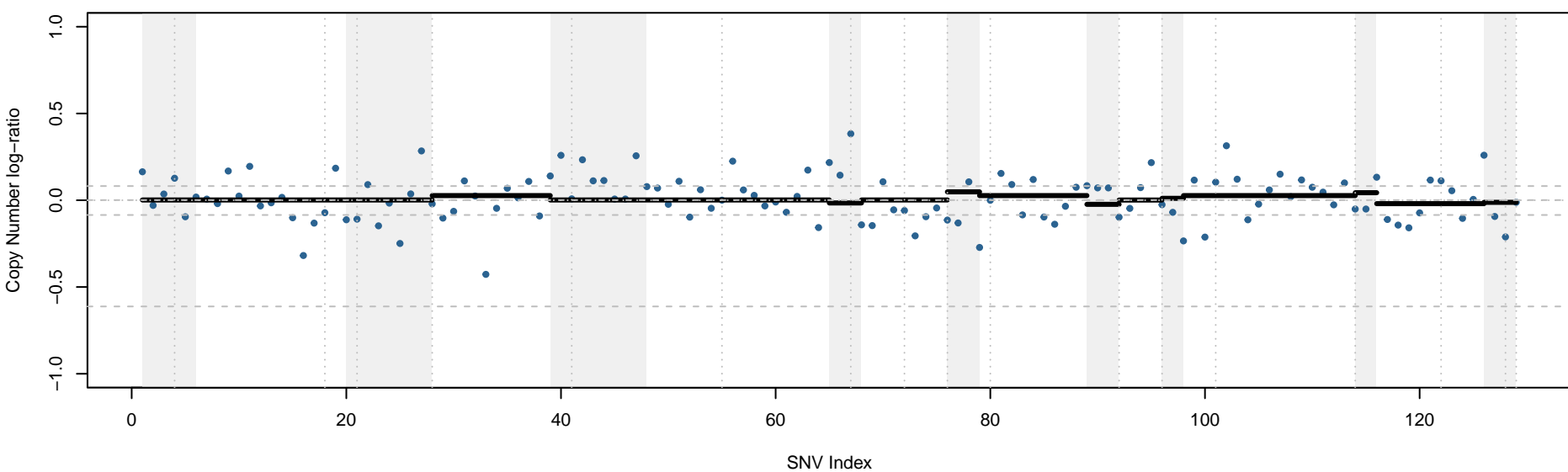
Purity: 0.15 Tumor ploidy: 5.99

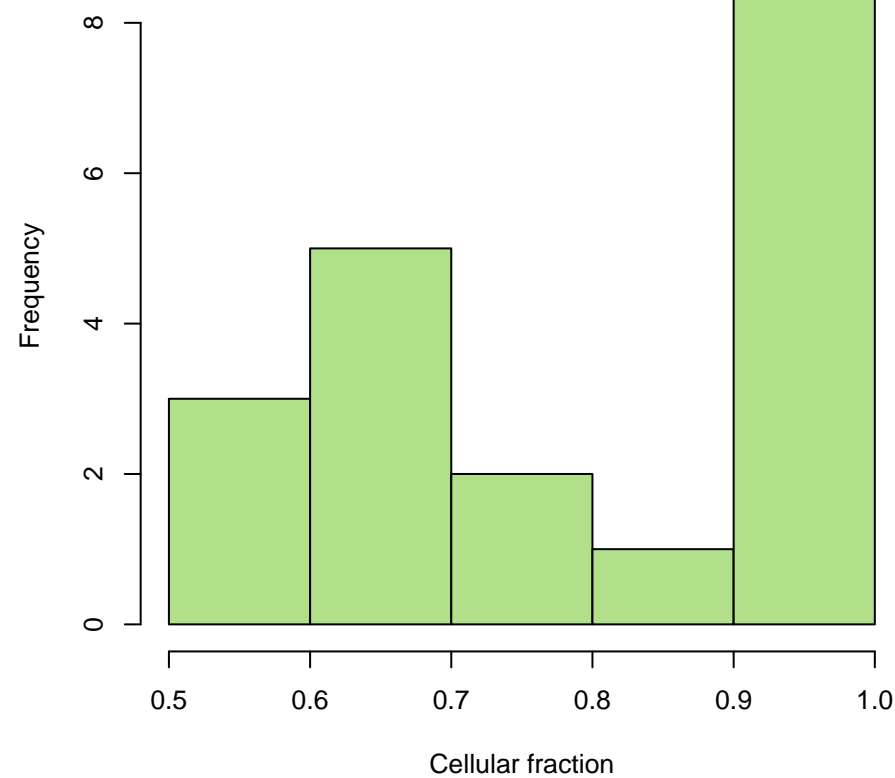
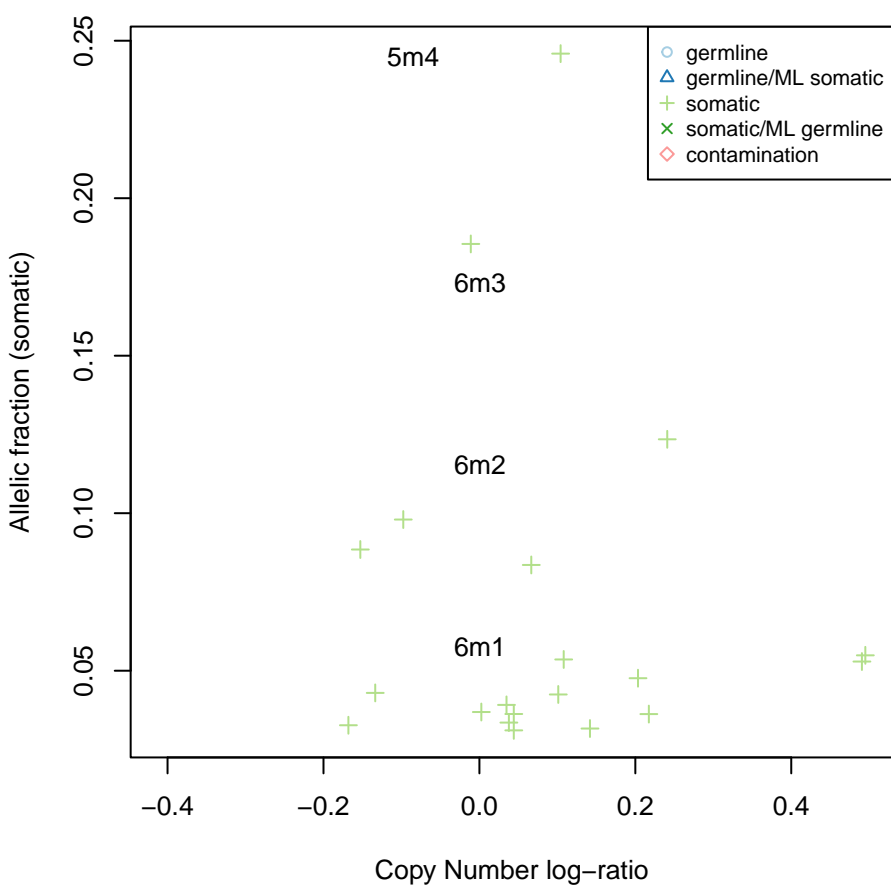
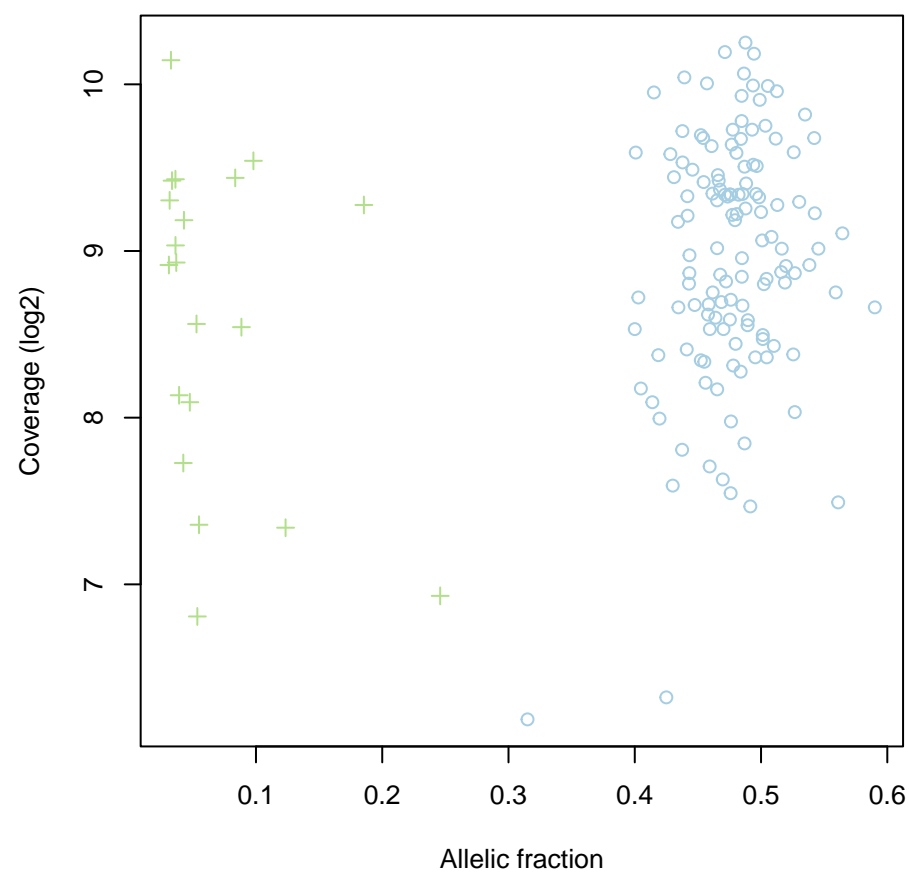
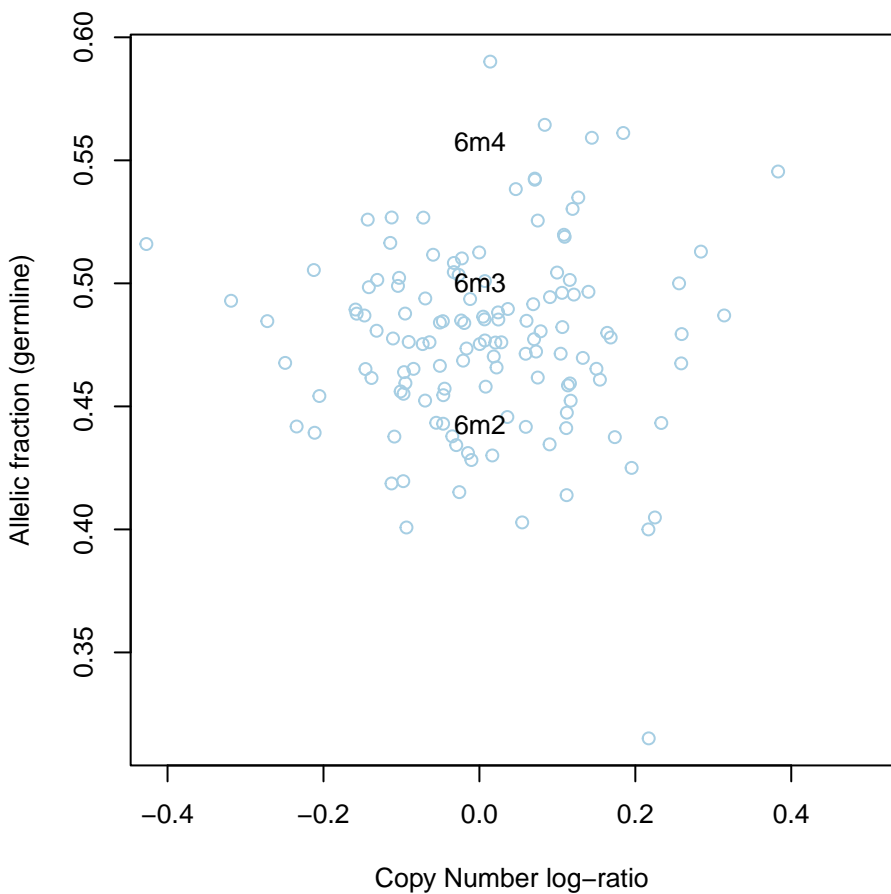


Purity: 0.15 Tumor ploidy: 5.99 SNV log-likelihood: -16.78 GoF: 97.4% Mean coverage: 549;494

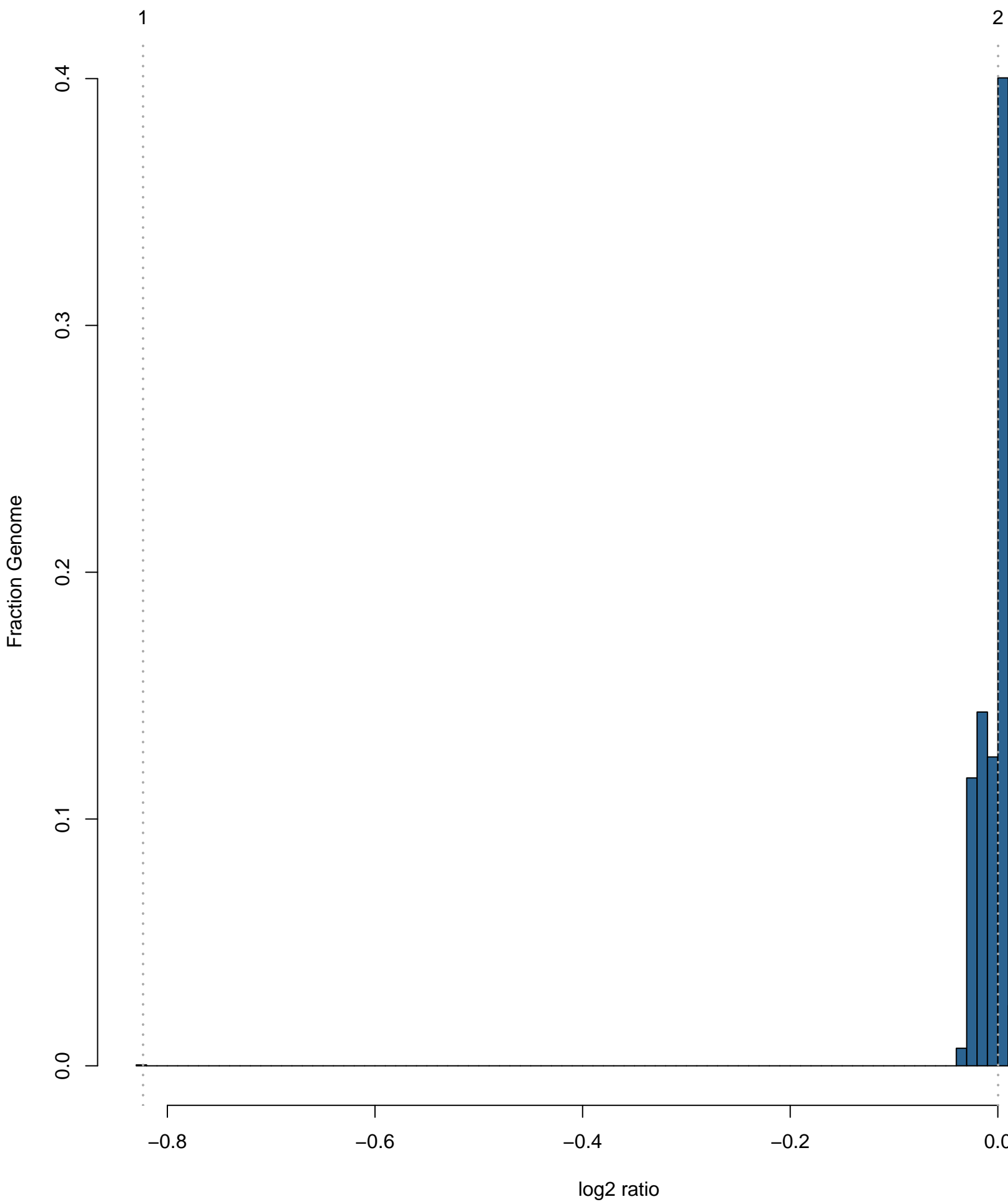


SCNA-fit log-likelihood: -4216.33

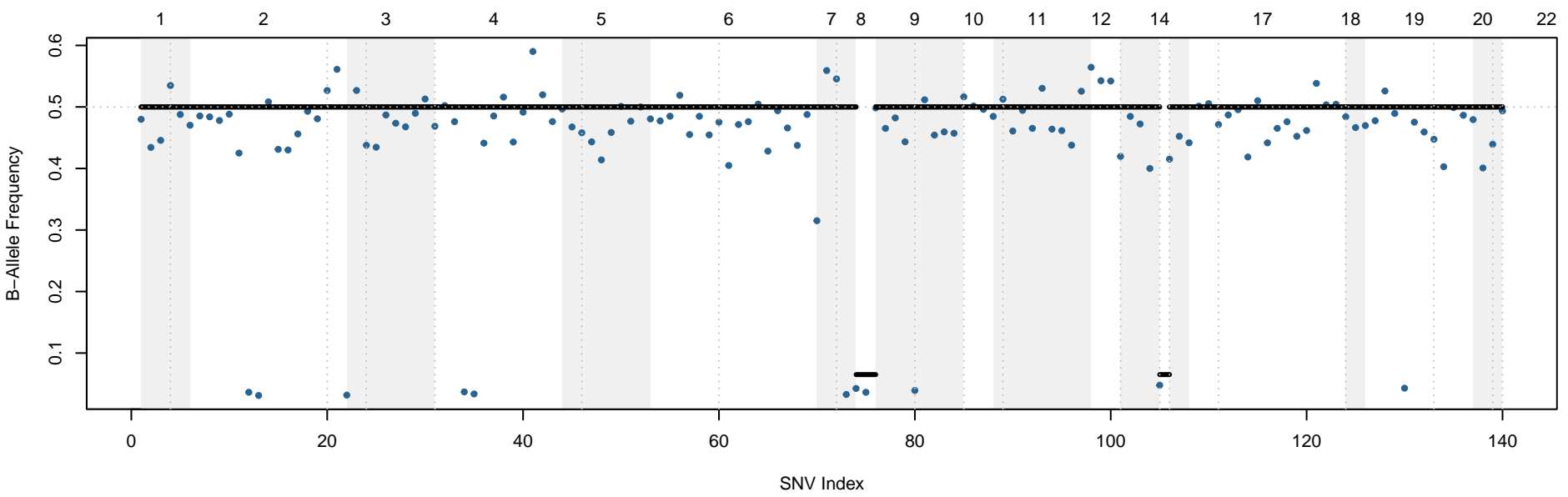




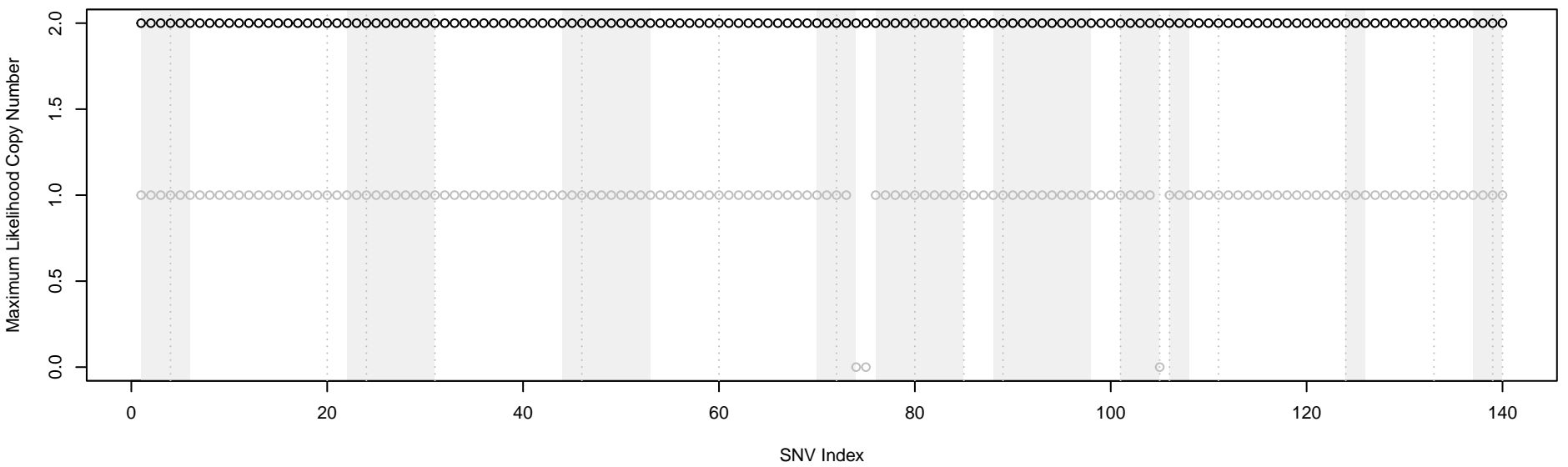
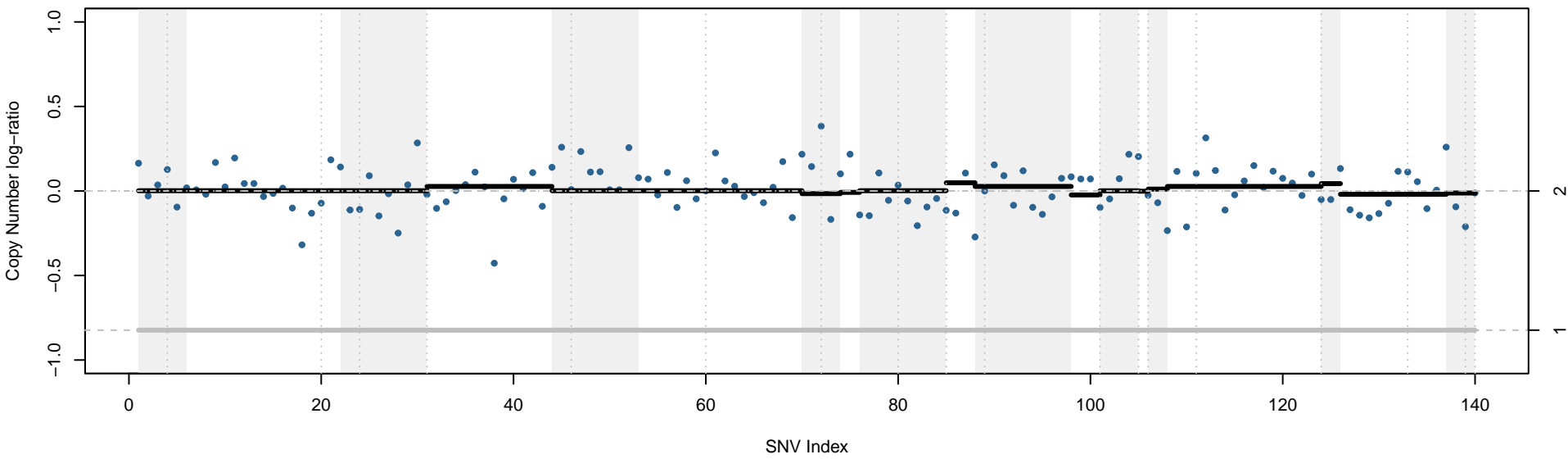
Purity: 0.87 Tumor ploidy: 2

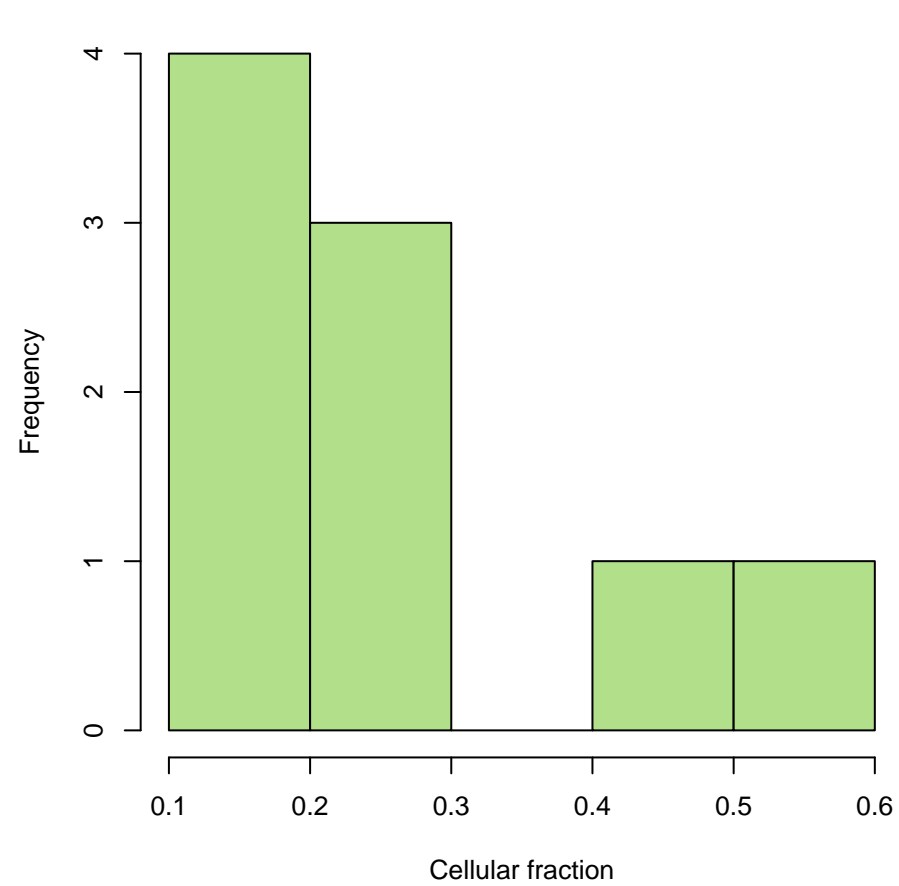
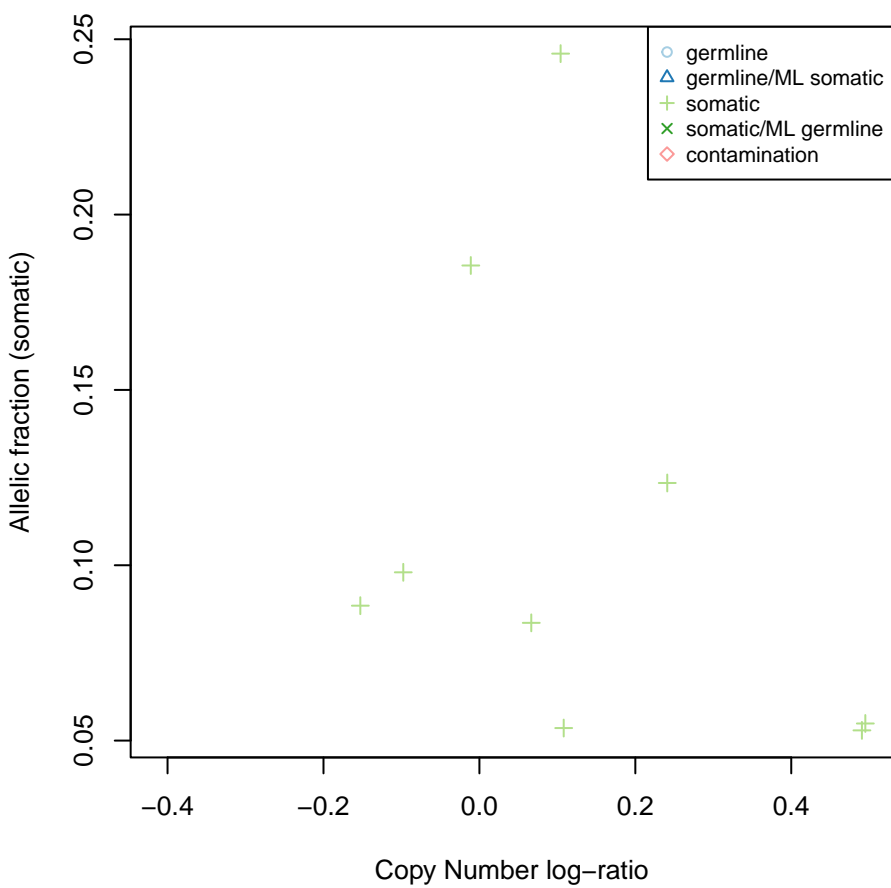
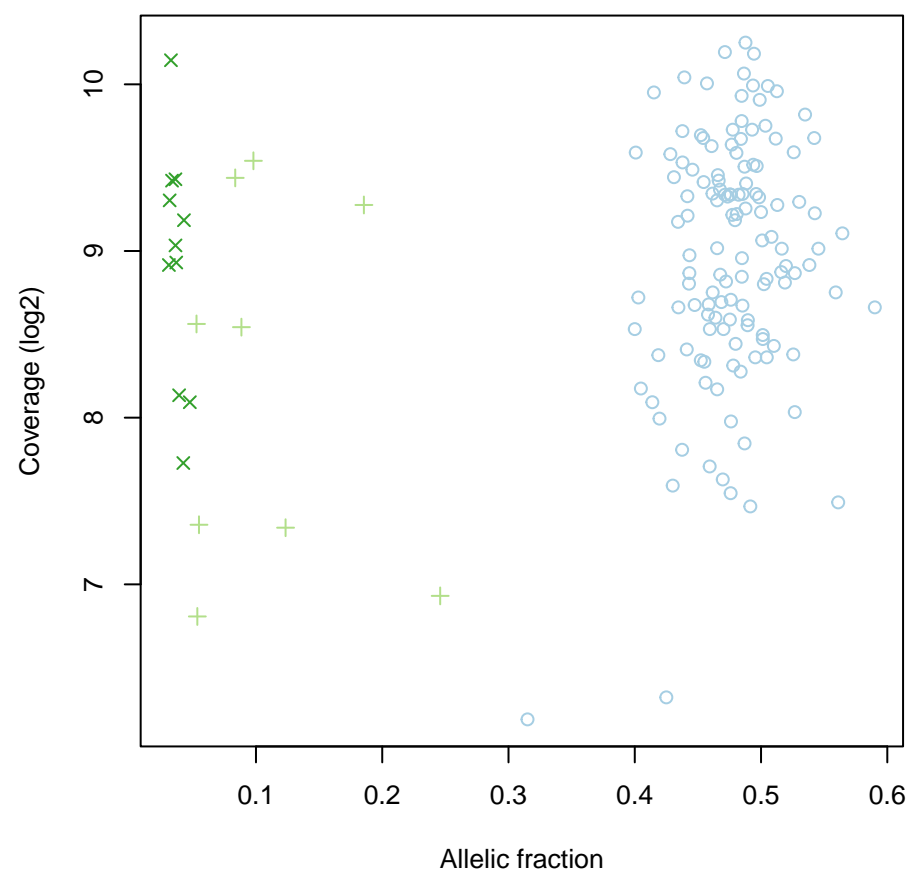
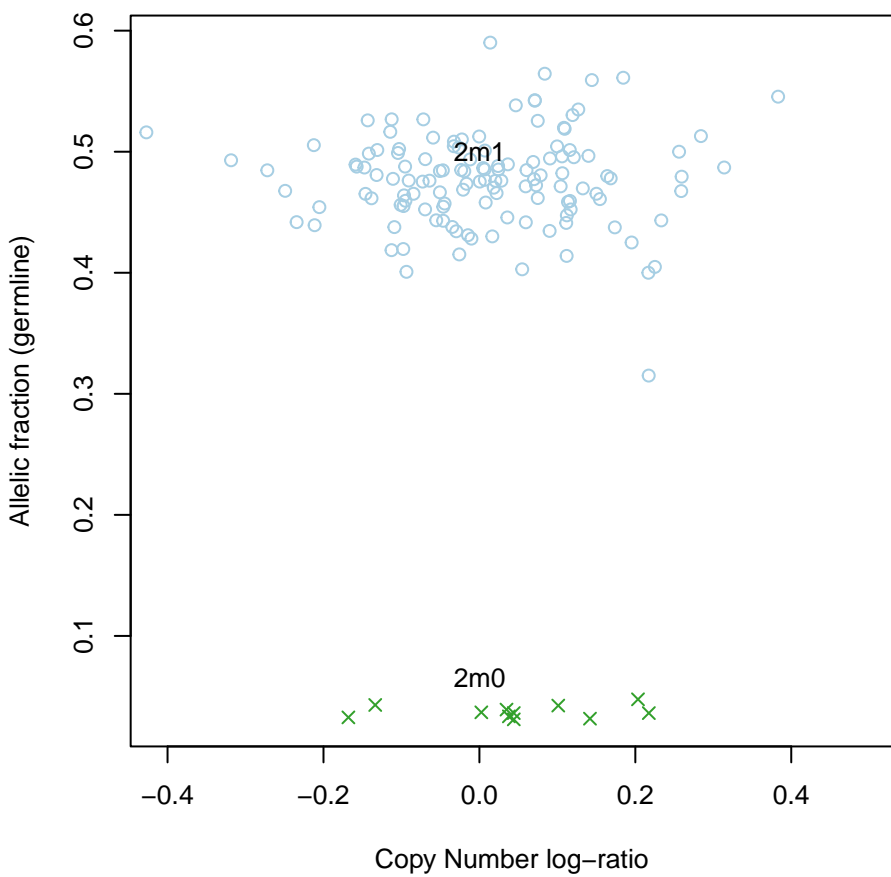


Purity: 0.87 Tumor ploidy: 2 SNV log-likelihood: -56.62 GoF: 80.2% Mean coverage: 549;494

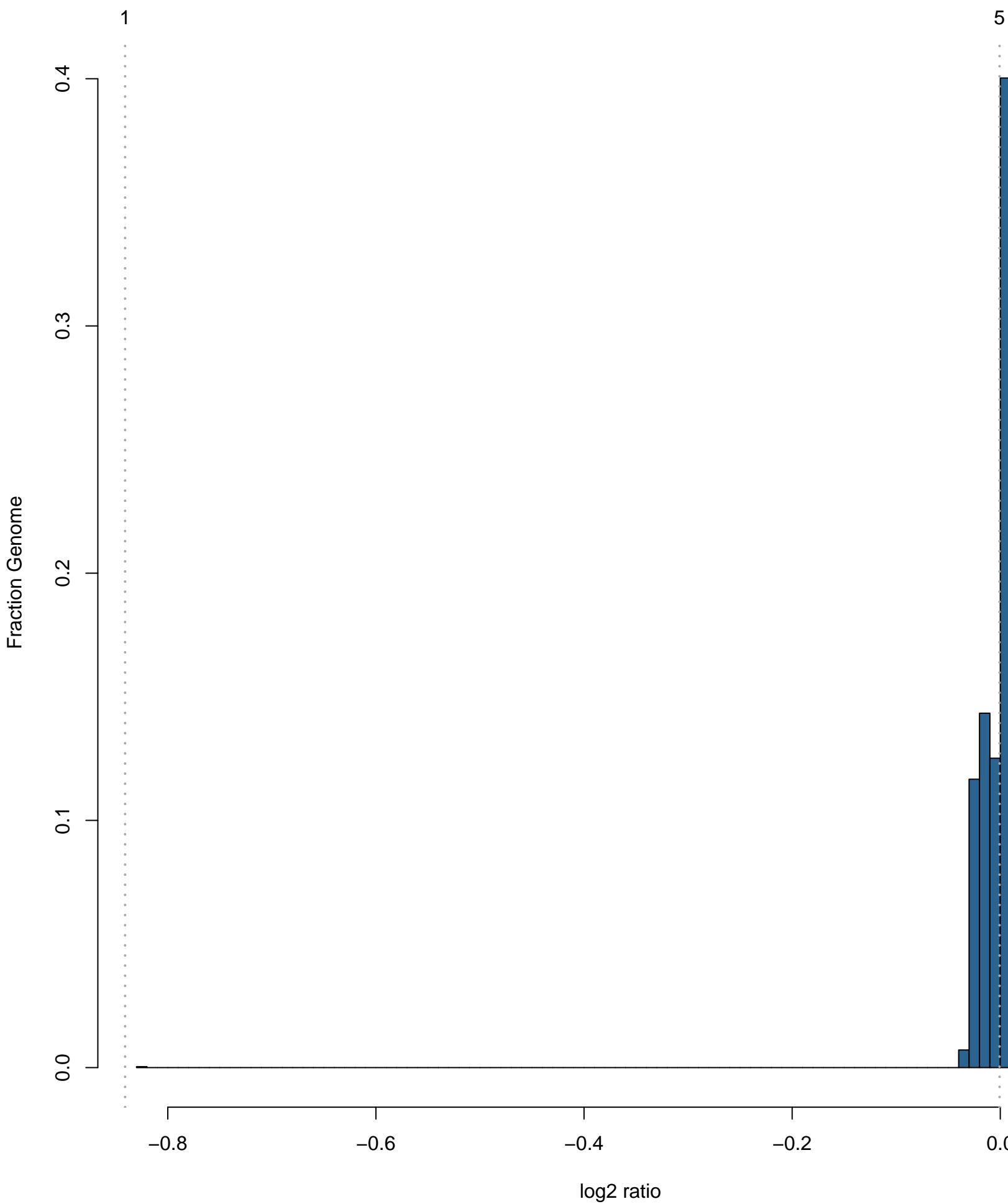


SCNA-fit log-likelihood: -4216.81

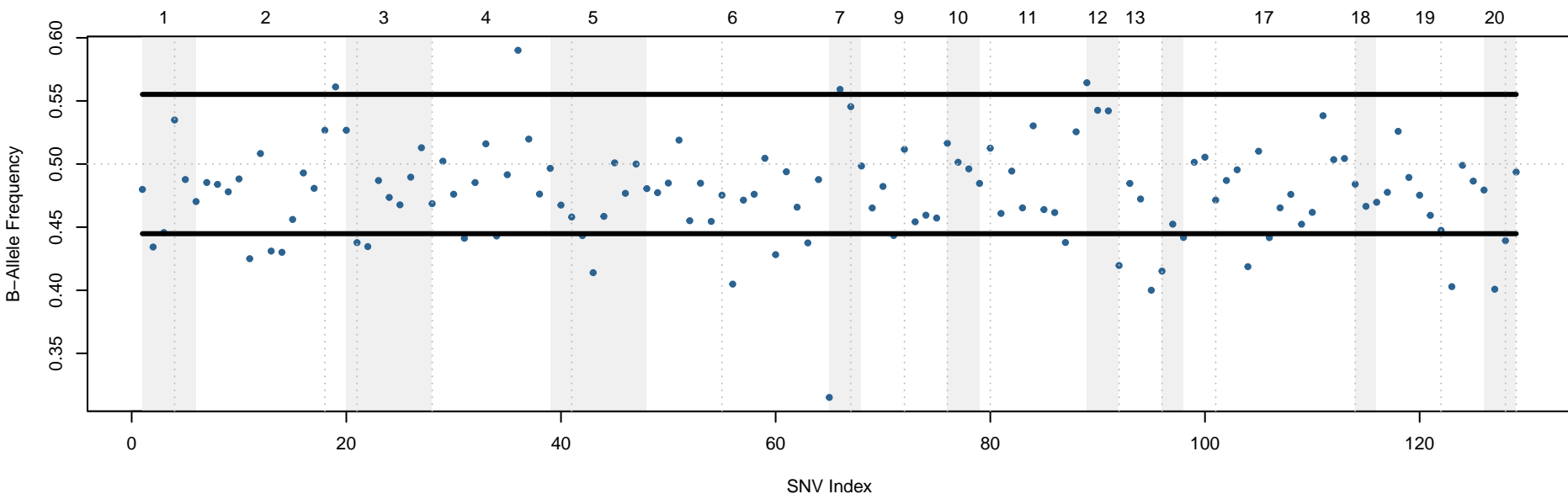




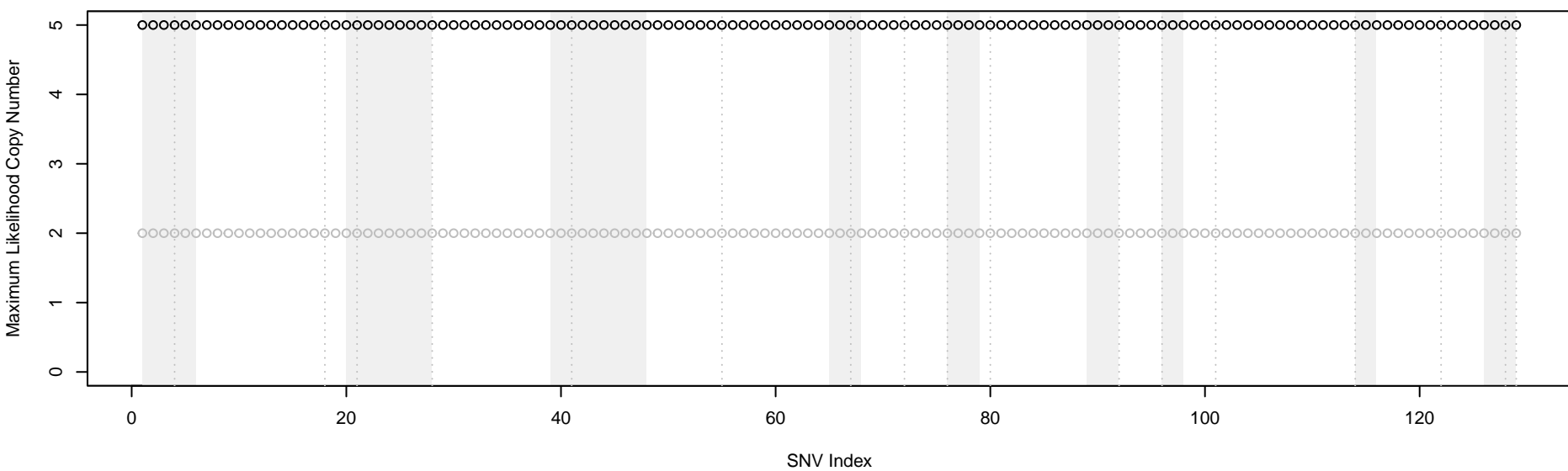
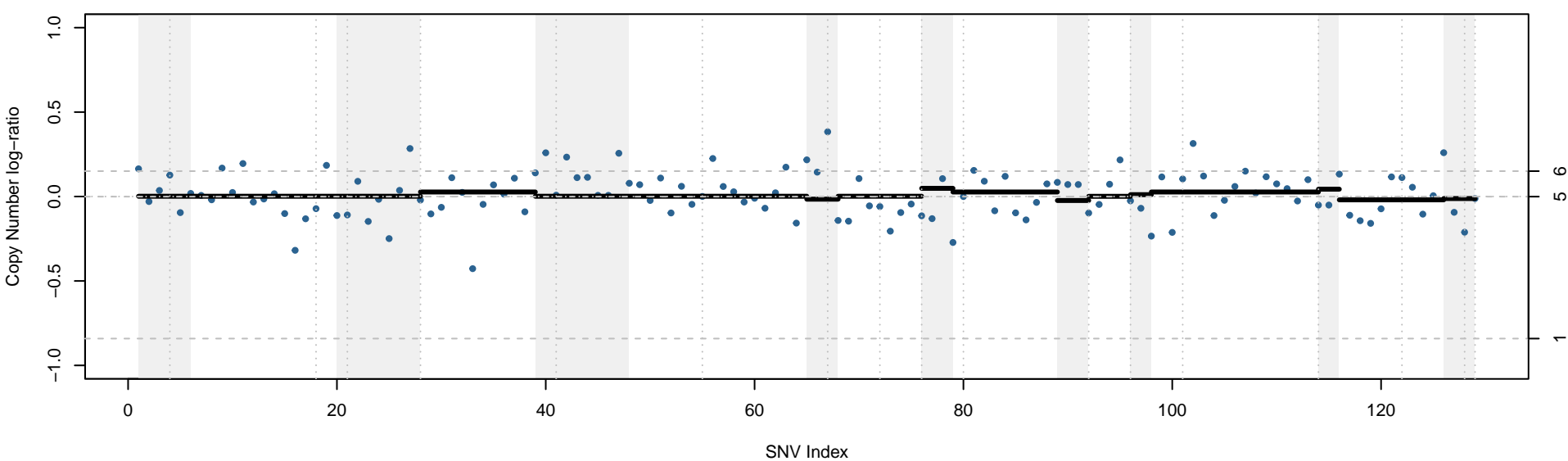
Purity: 0.33 Tumor ploidy: 5.004



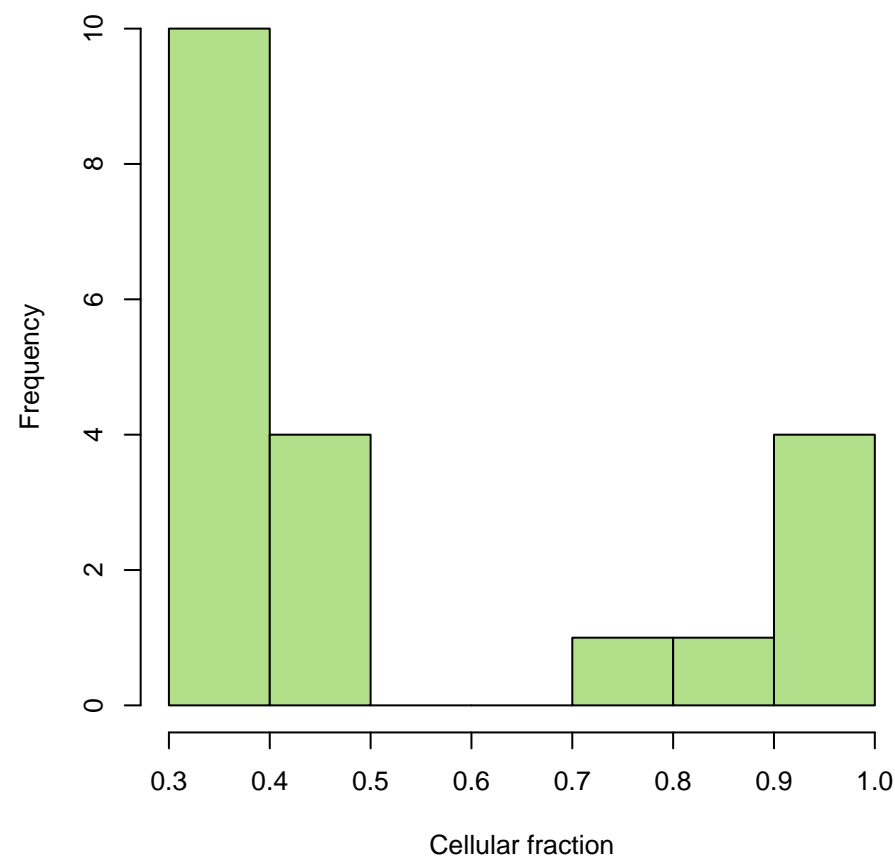
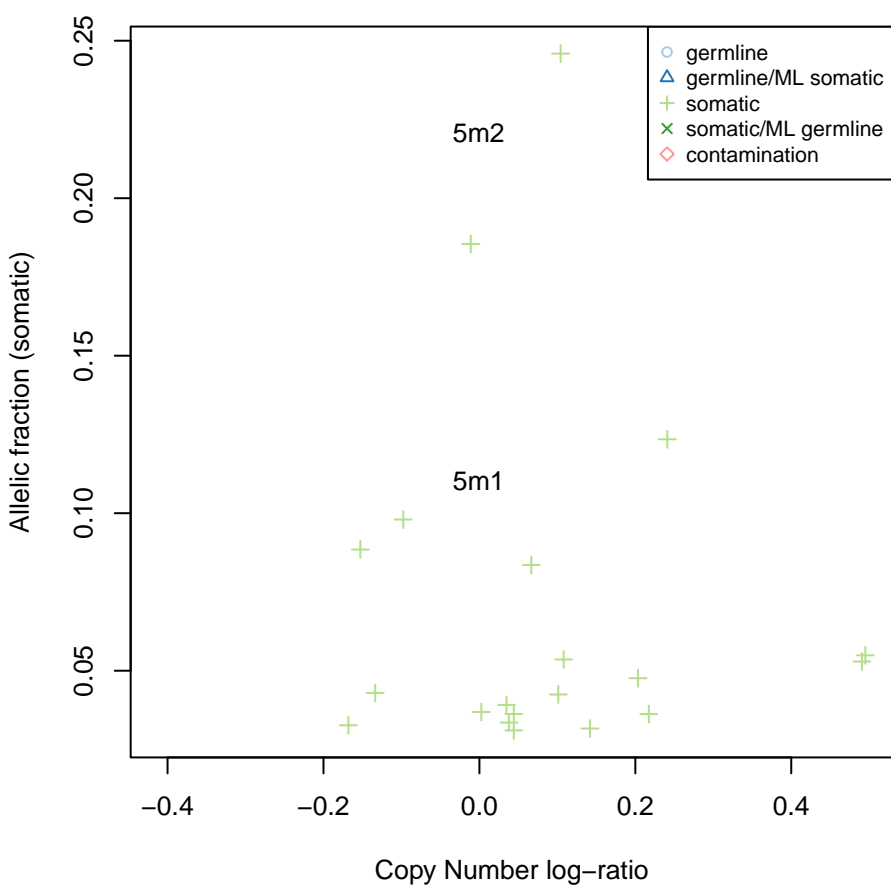
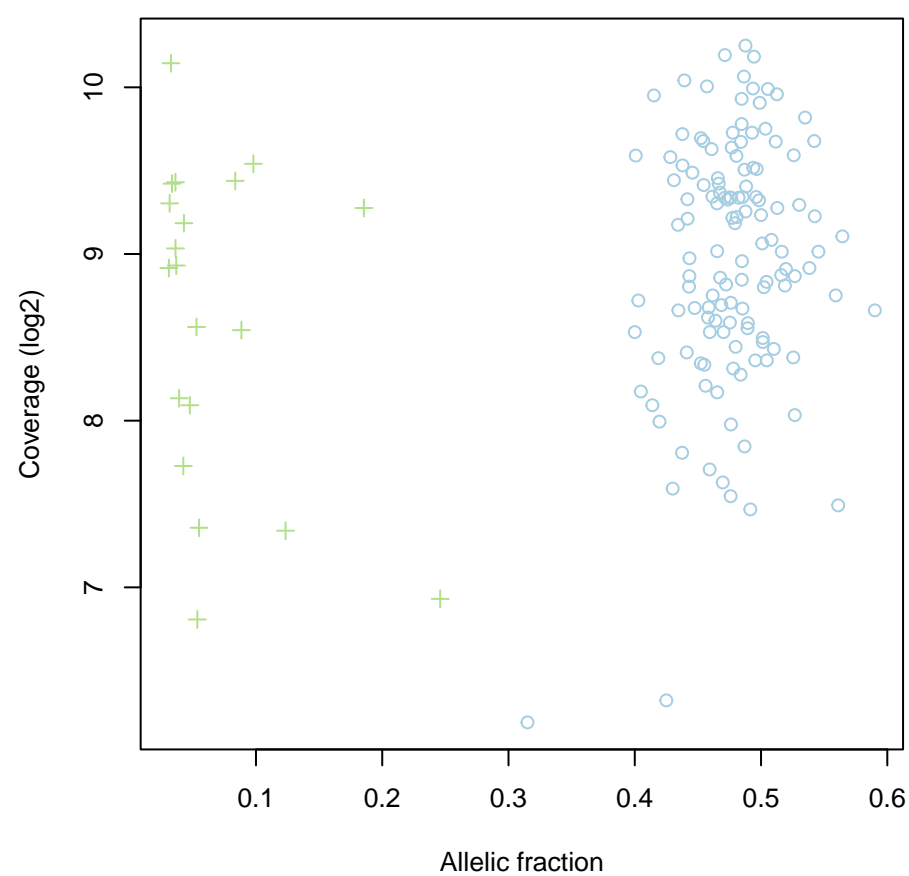
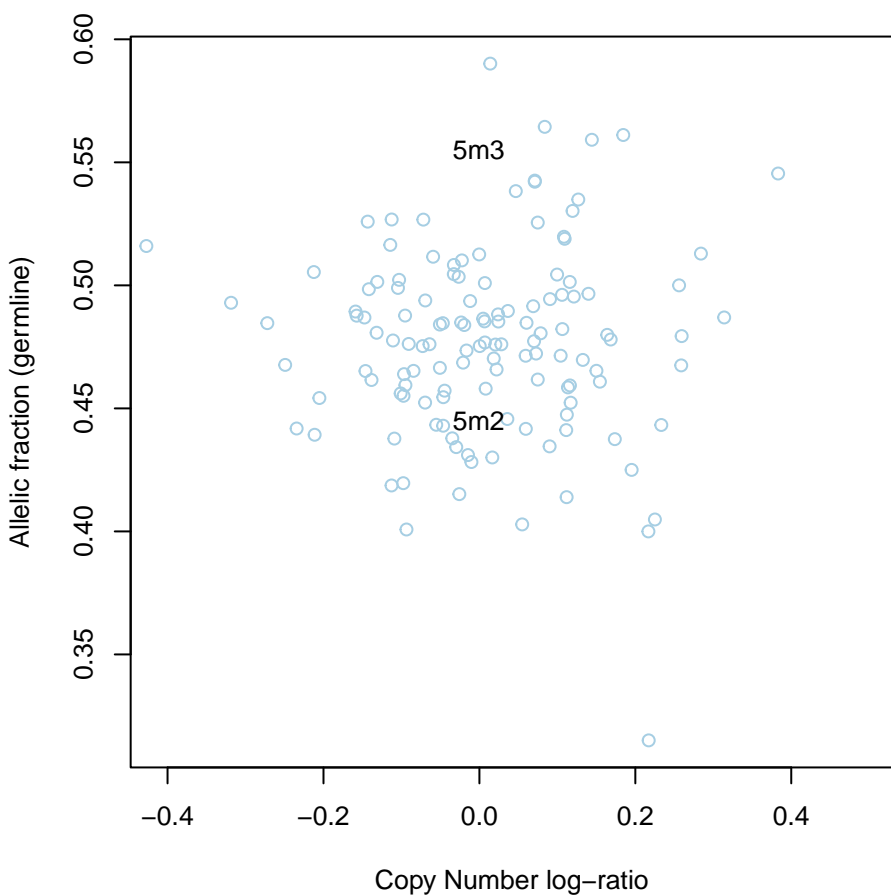
Purity: 0.33 Tumor ploidy: 5.004 SNV log-likelihood: -88.67 GoF: 95.9% Mean coverage: 549;494



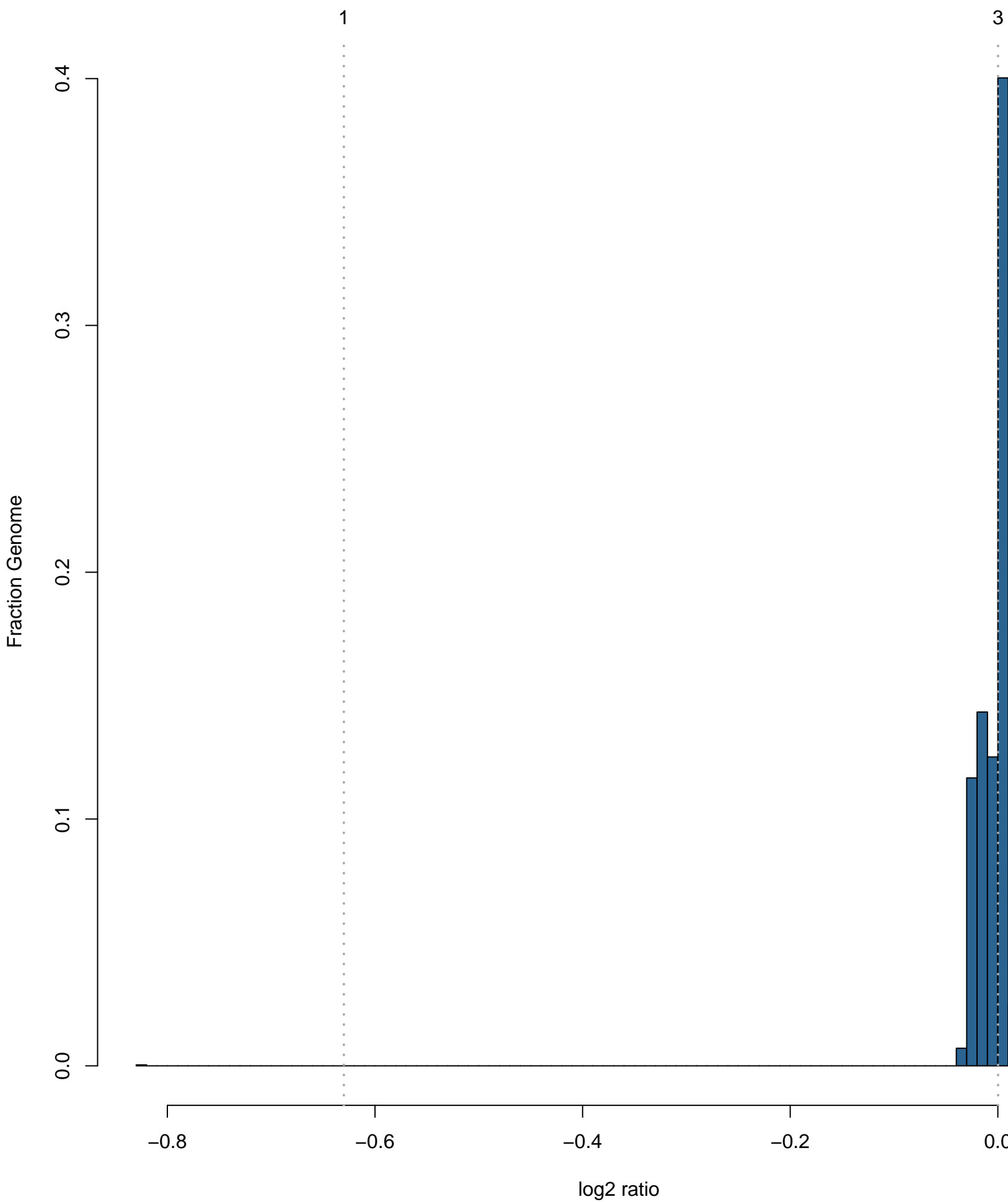
SCNA-fit log-likelihood: -4215.31



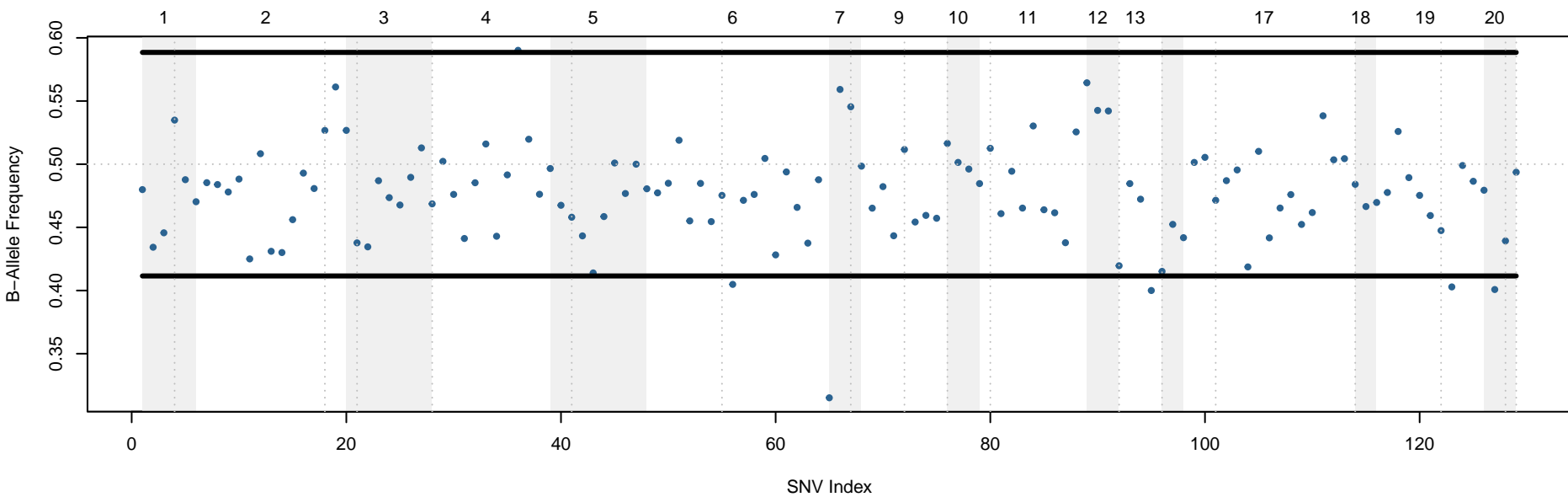




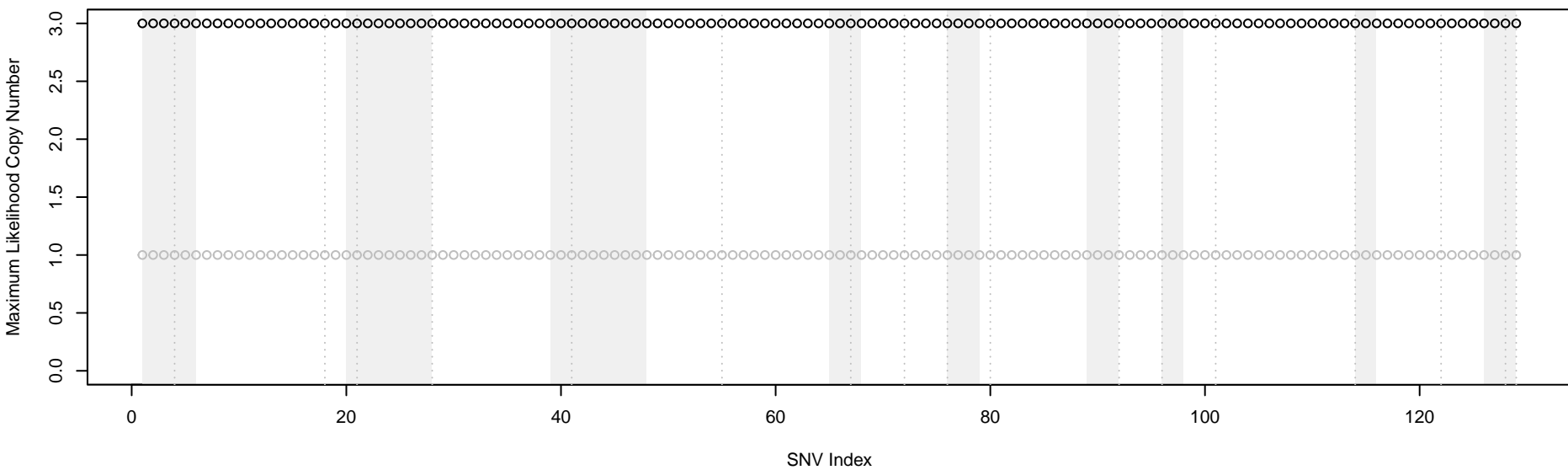
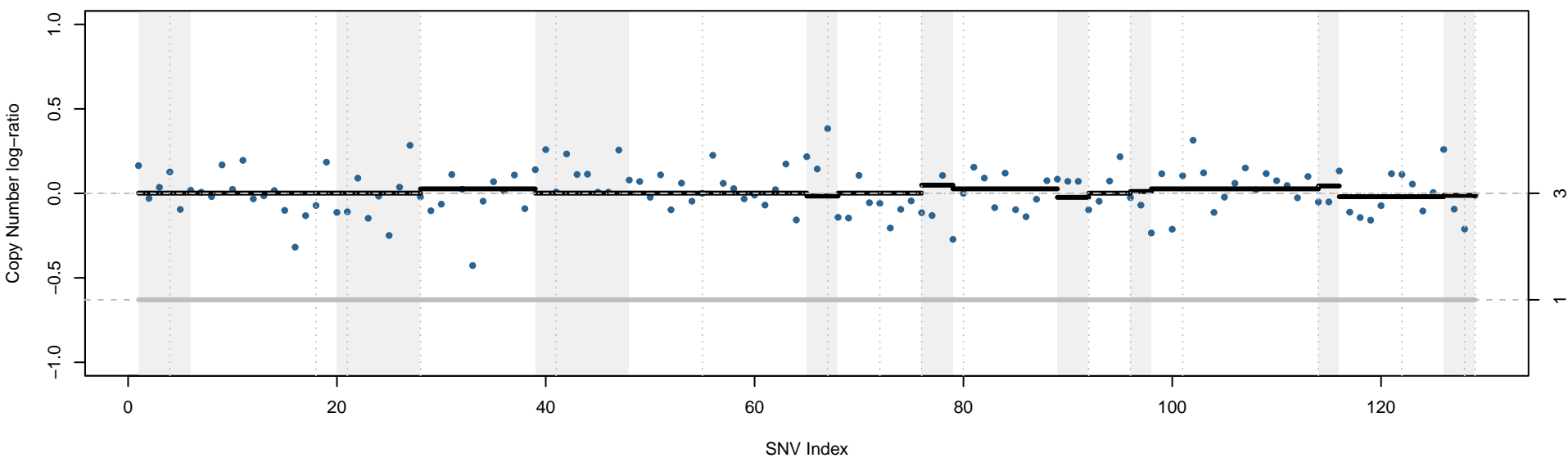
Purity: 0.43 Tumor ploidy: 2.999

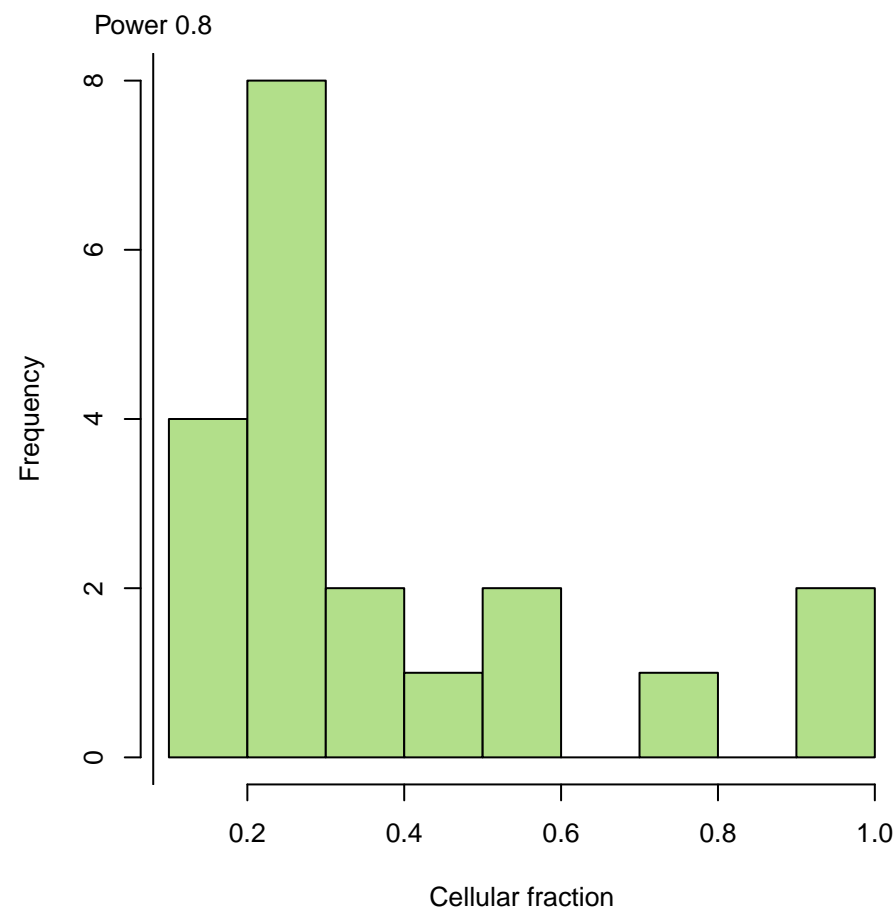
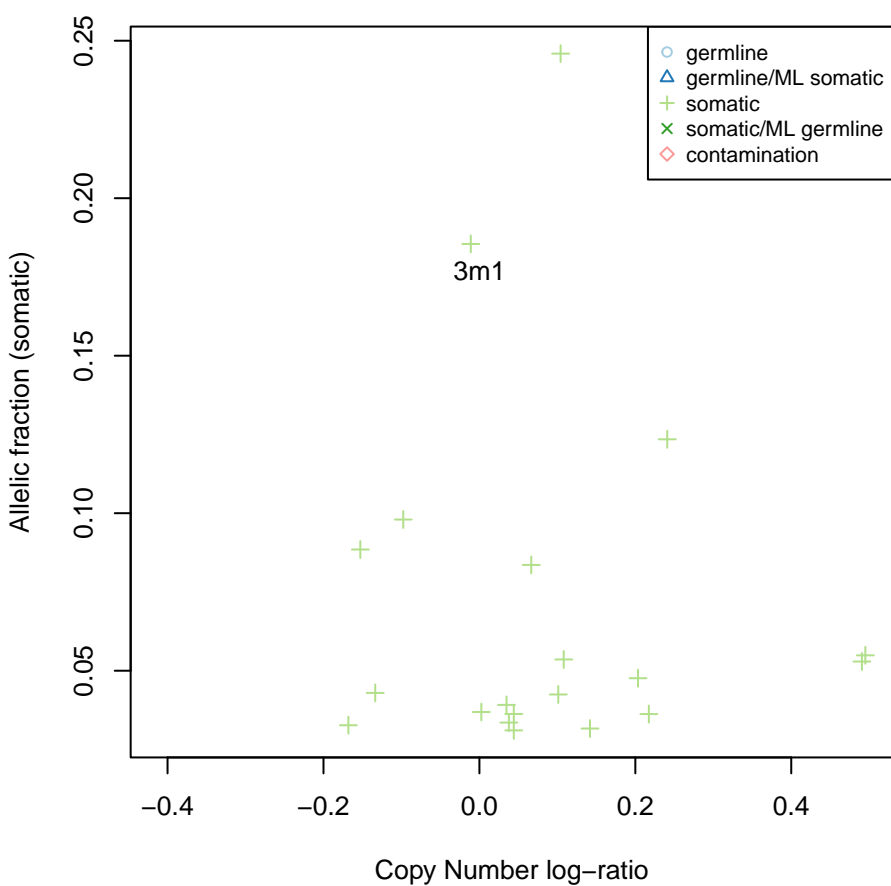
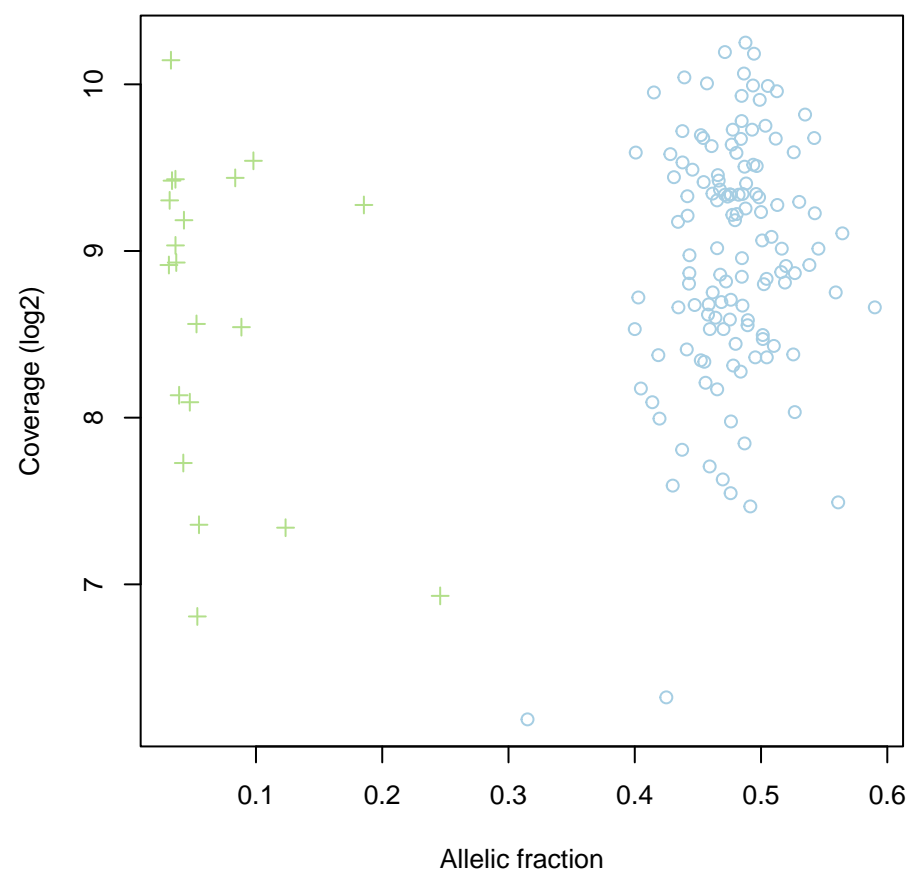
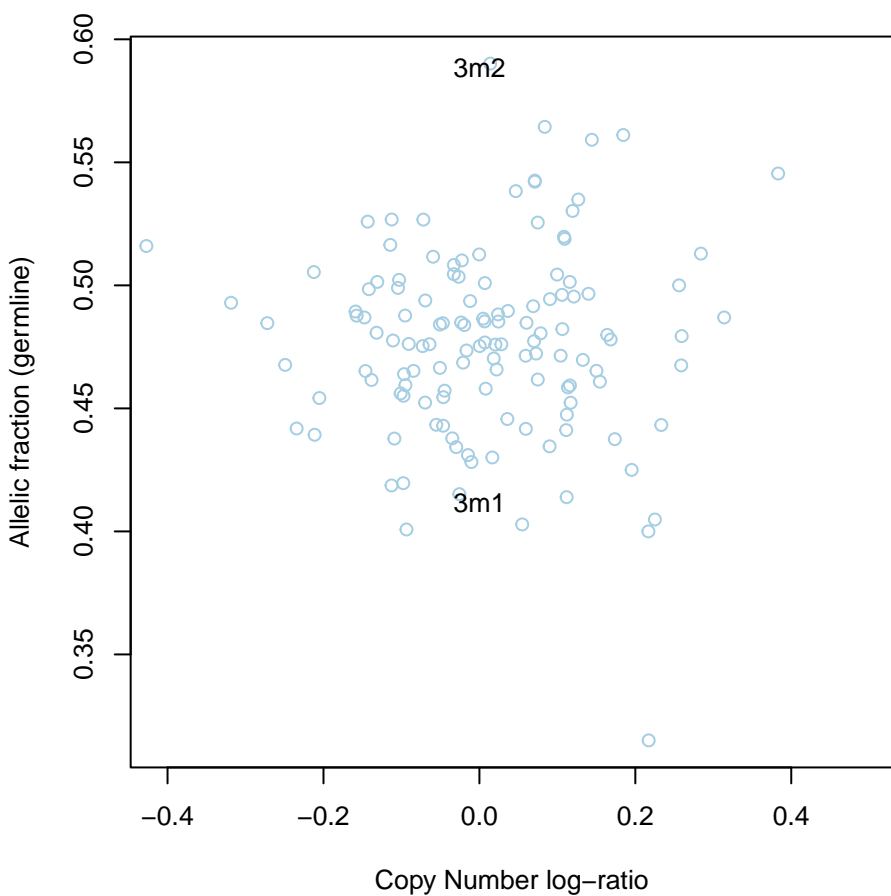


Purity: 0.43 Tumor ploidy: 2.999 SNV log-likelihood: -261.32 GoF: 86.2% Mean coverage: 549,494

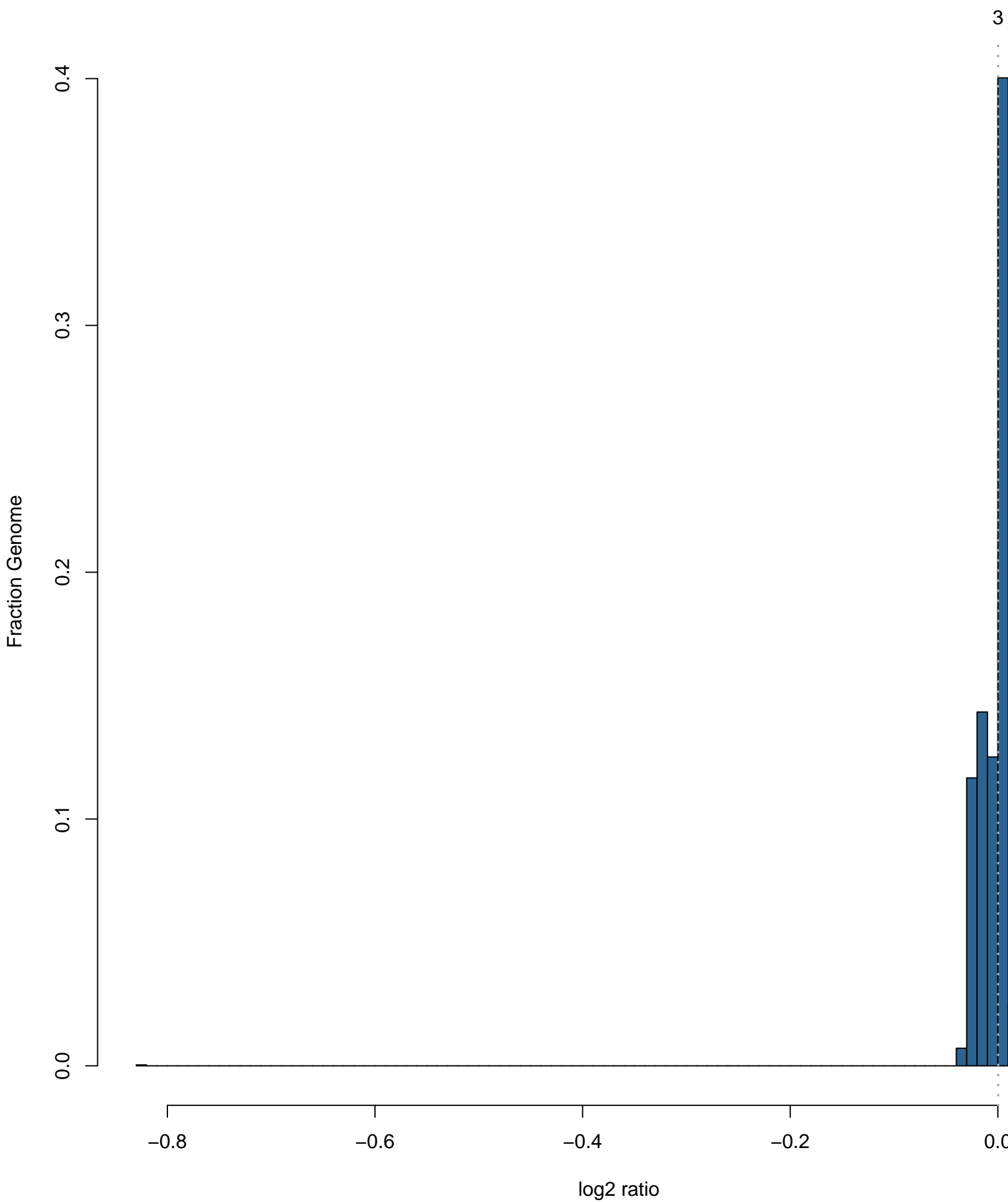


SCNA-fit log-likelihood: -4217.56

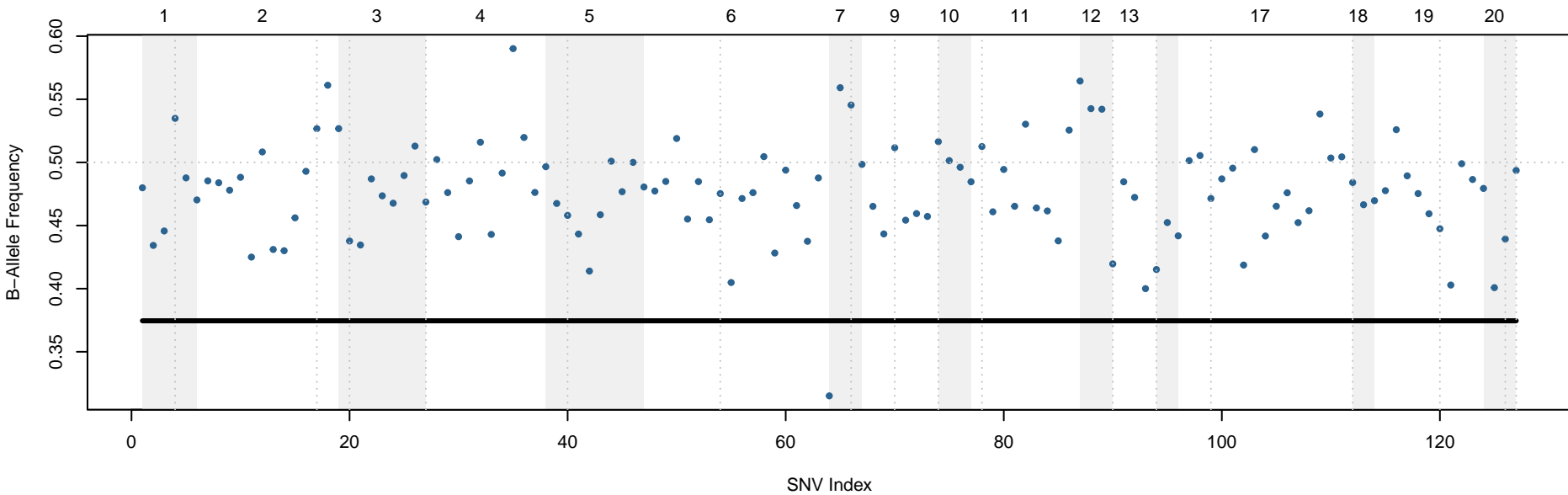




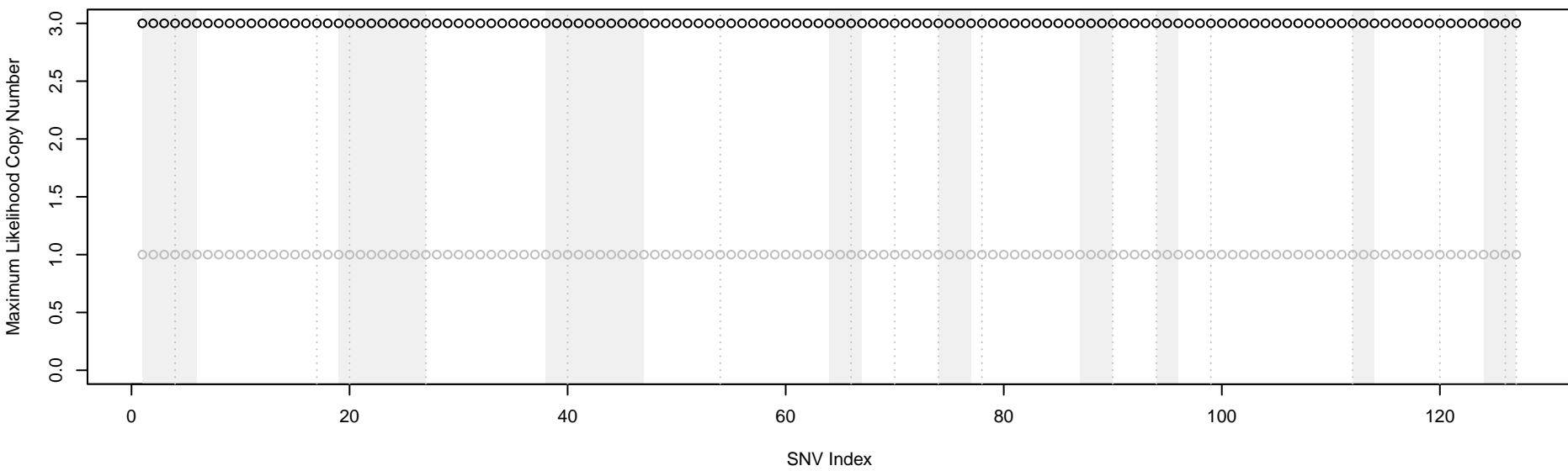
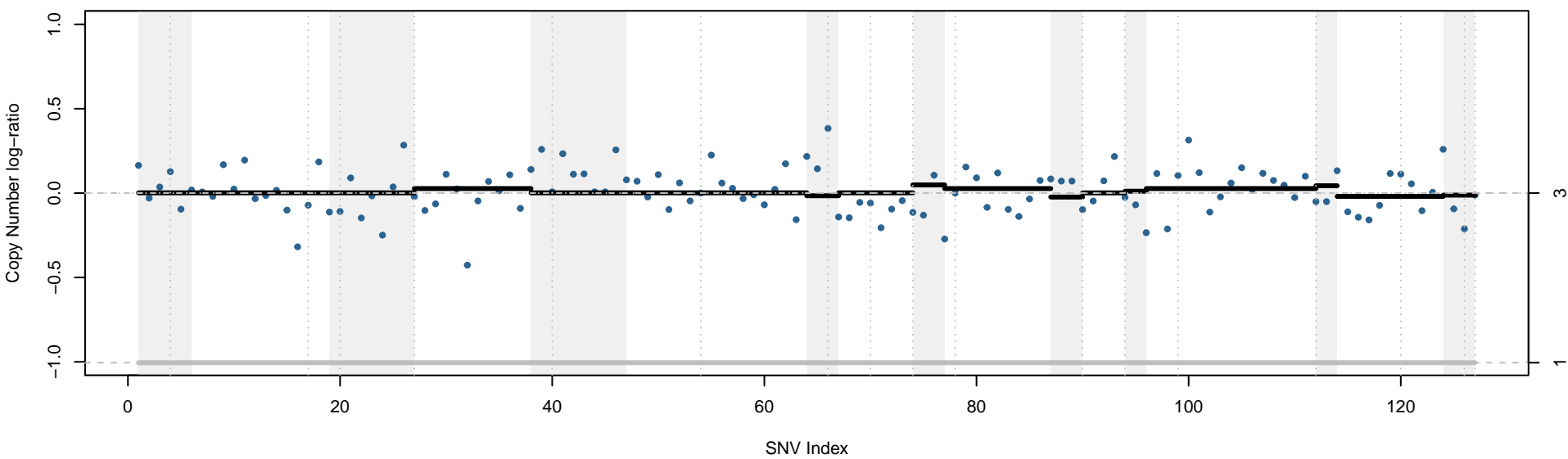
Purity: 0.67 Tumor ploidy: 2.999

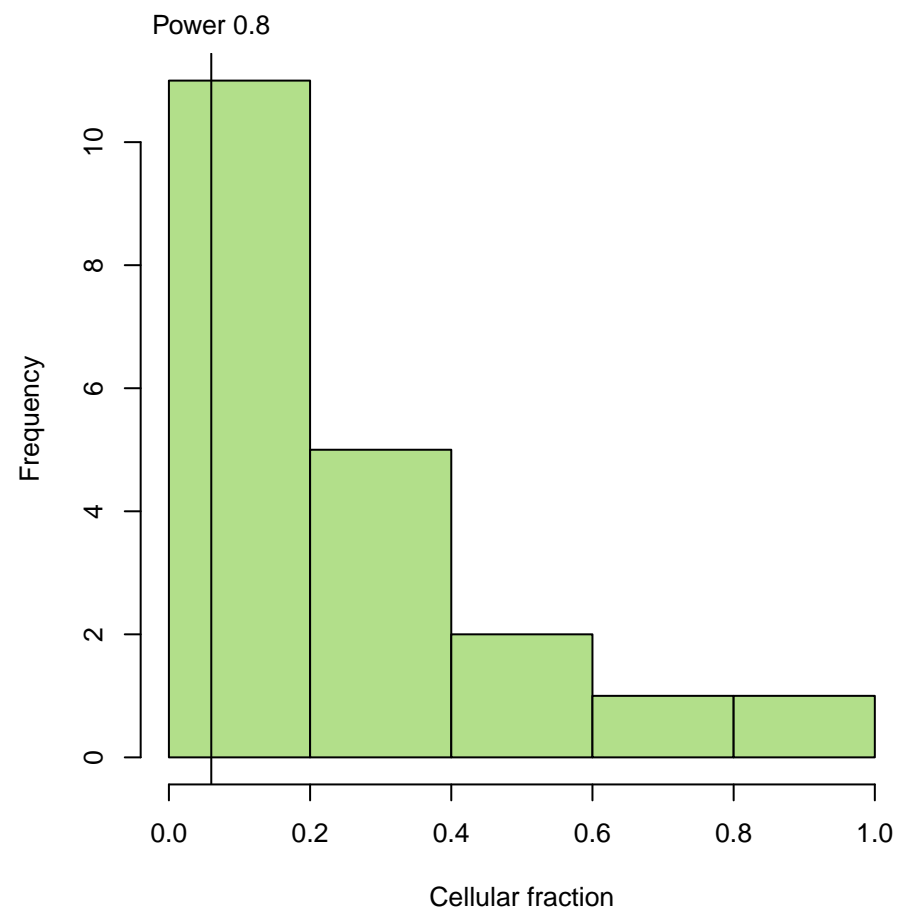
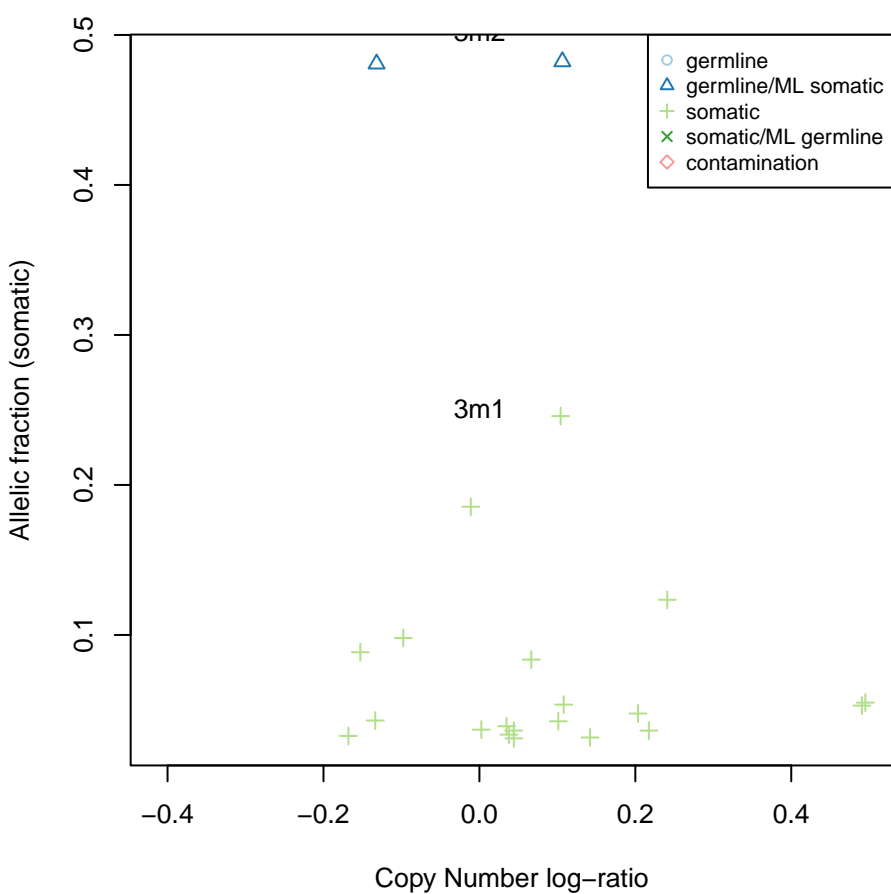
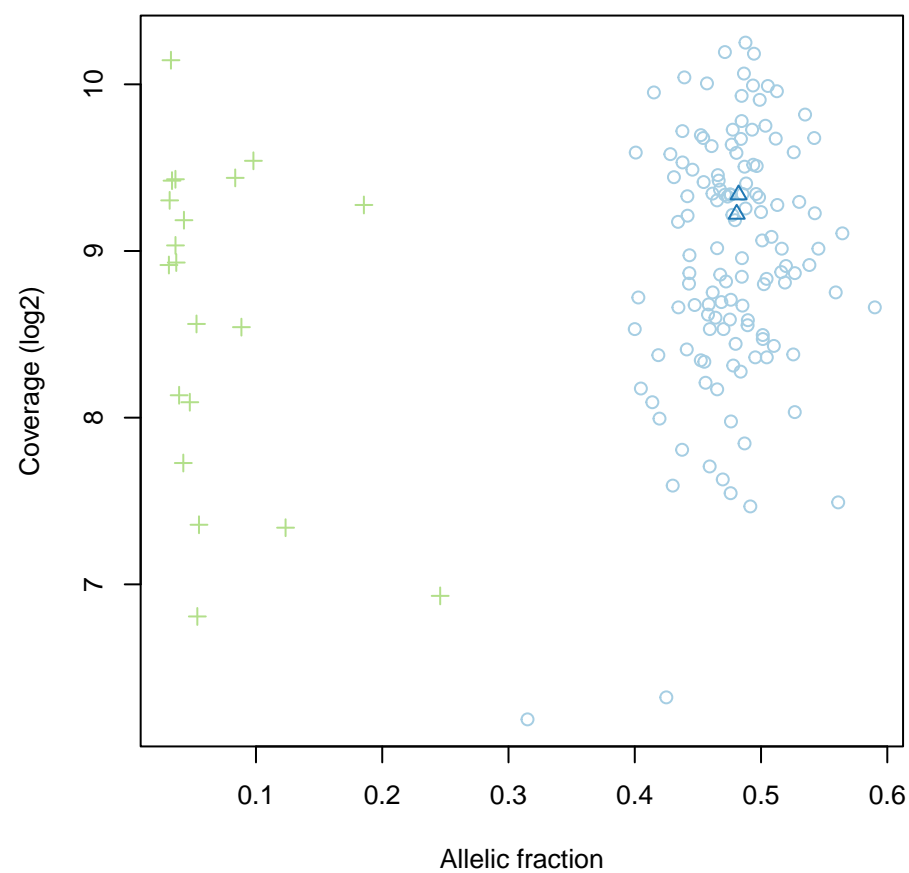
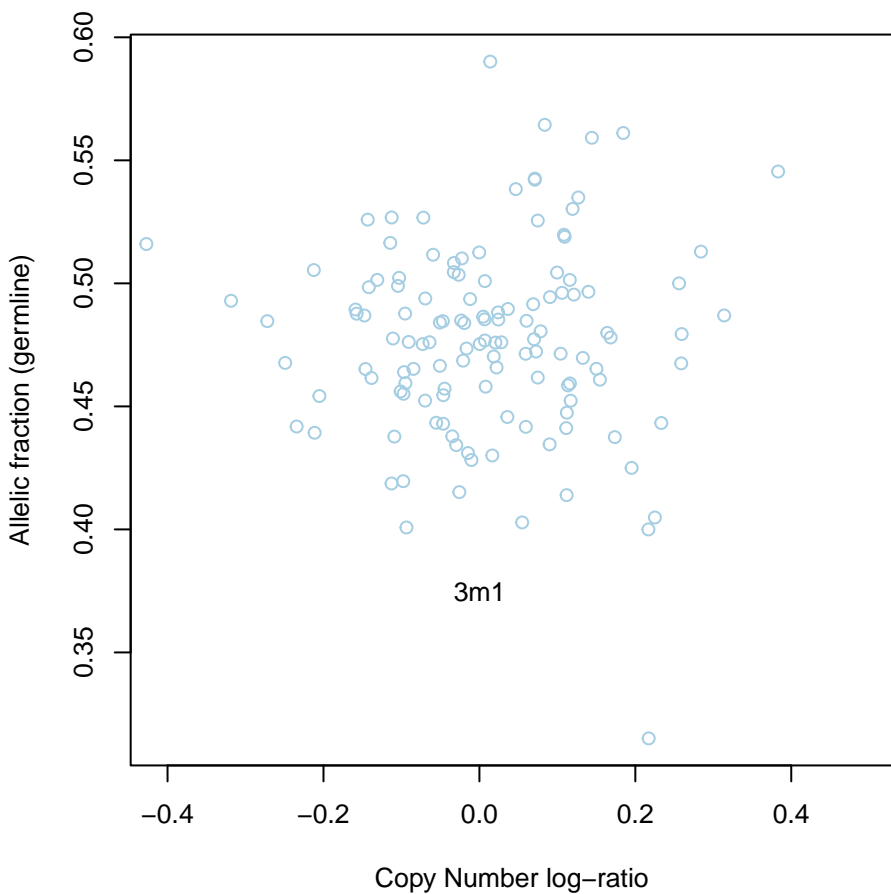


Purity: 0.67 Tumor ploidy: 2.999 SNV log-likelihood: -741.92 GoF: 67.3% Mean coverage: 549,494

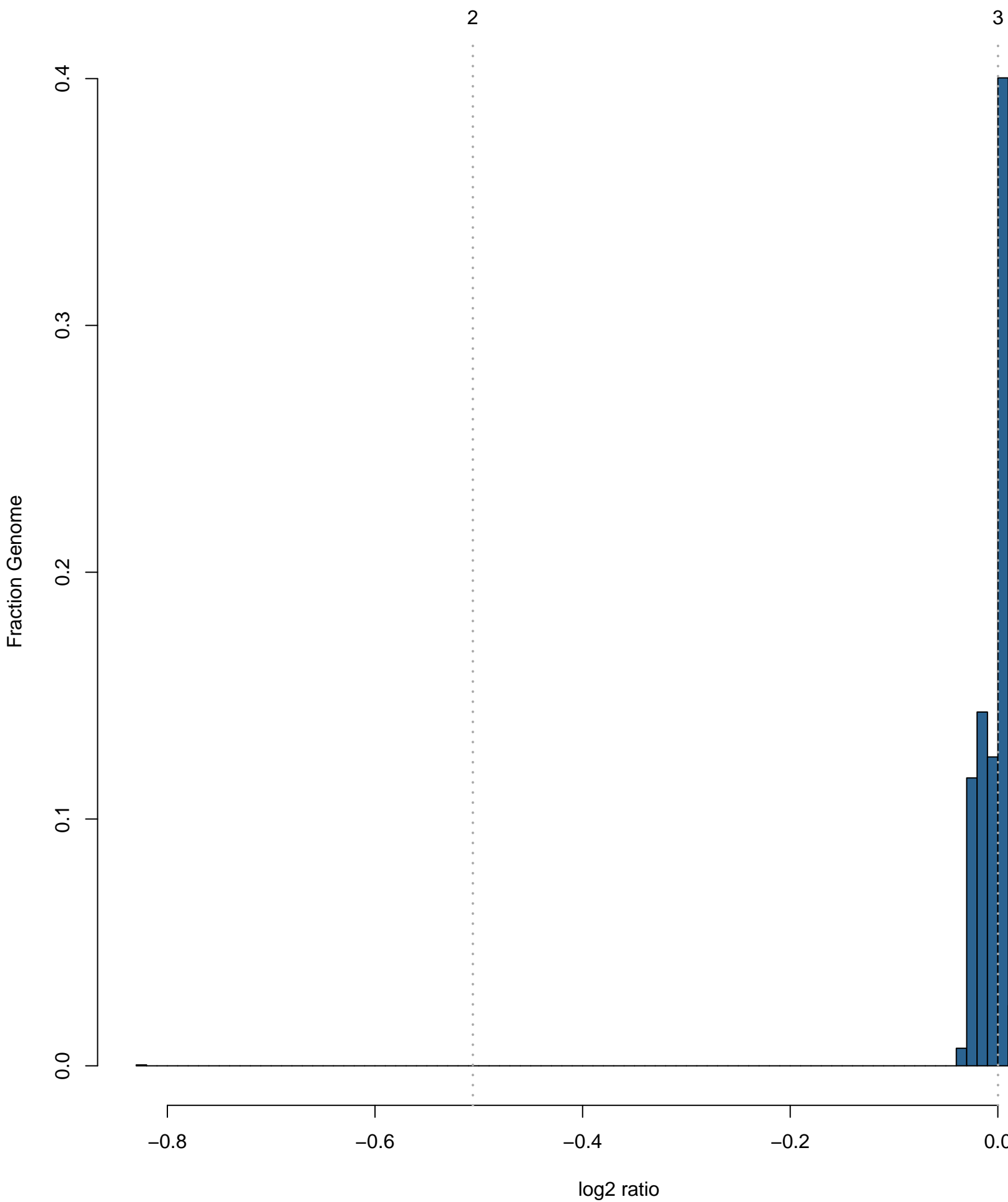


SCNA-fit log-likelihood: -4217.53



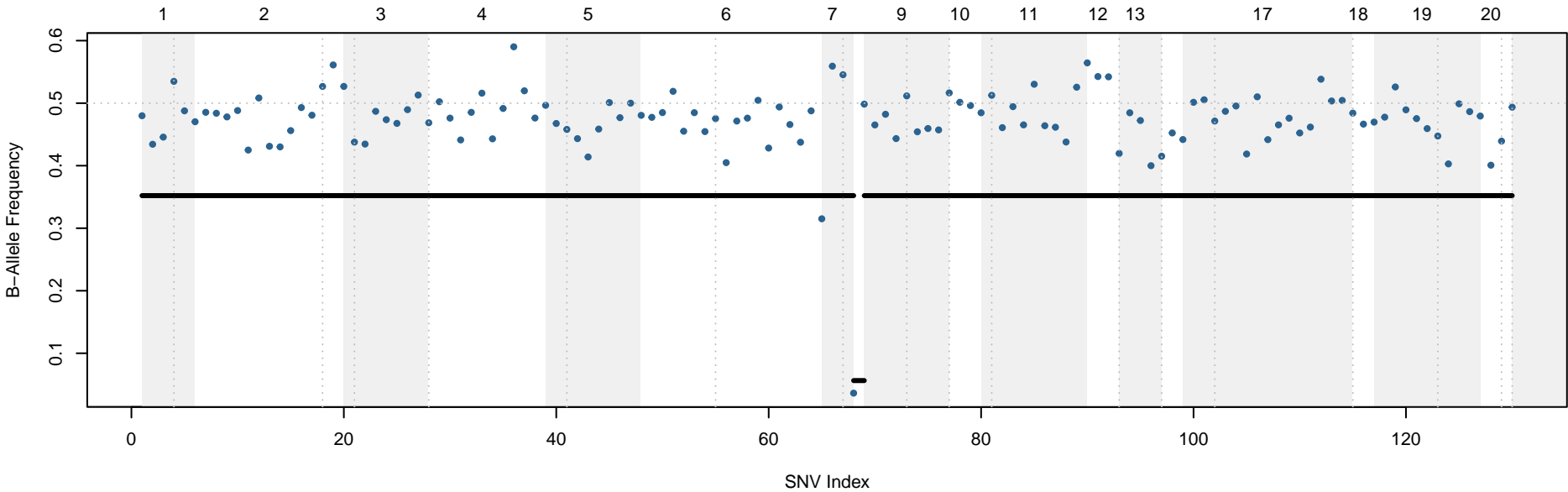


Purity: 0.84 Tumor ploidy: 3





Purity: 0.84 Tumor ploidy: 3 SNV log-likelihood: -1164.86 GoF: 51.5% Mean coverage: 549;494



SCNA-fit log-likelihood: -4219.04

