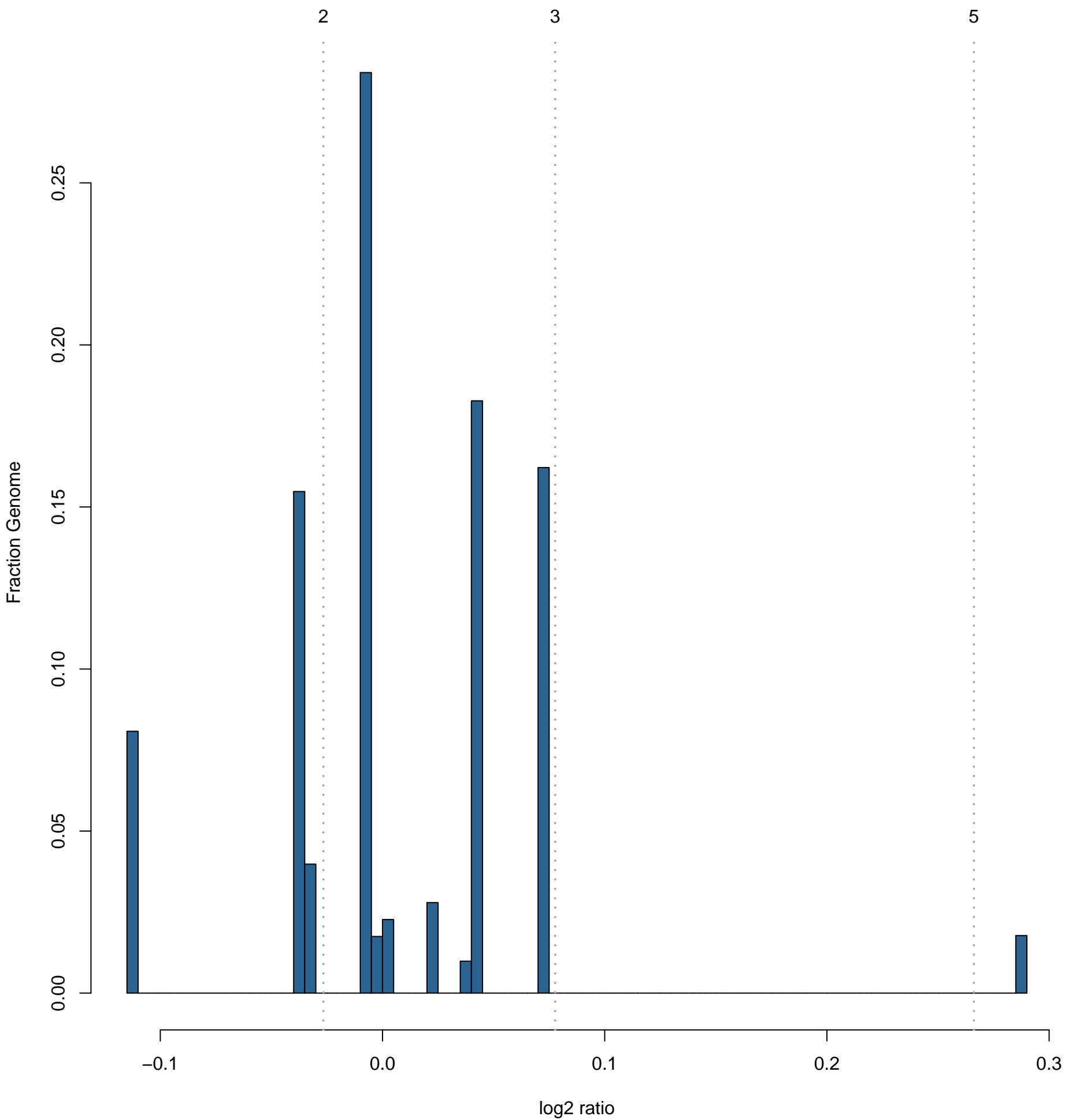
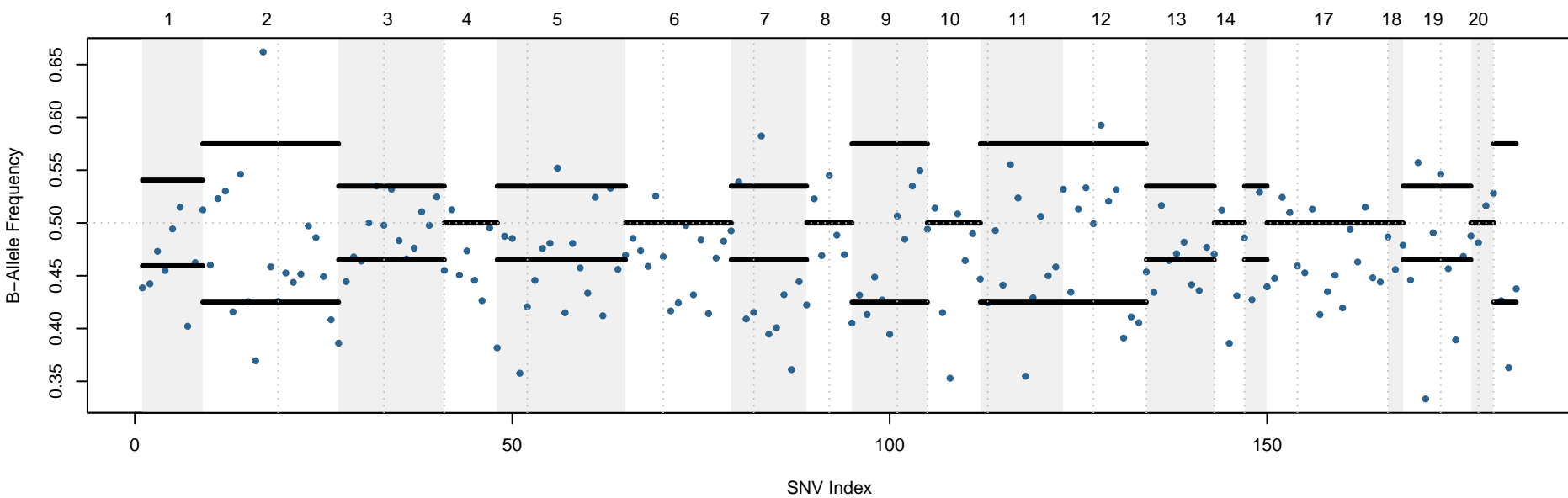


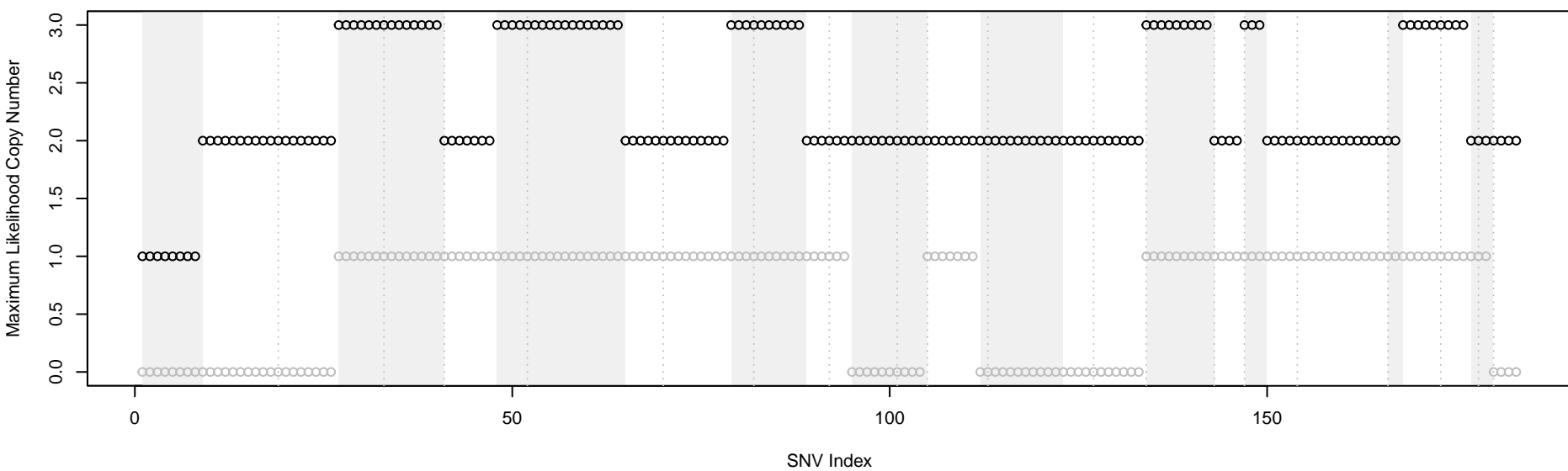
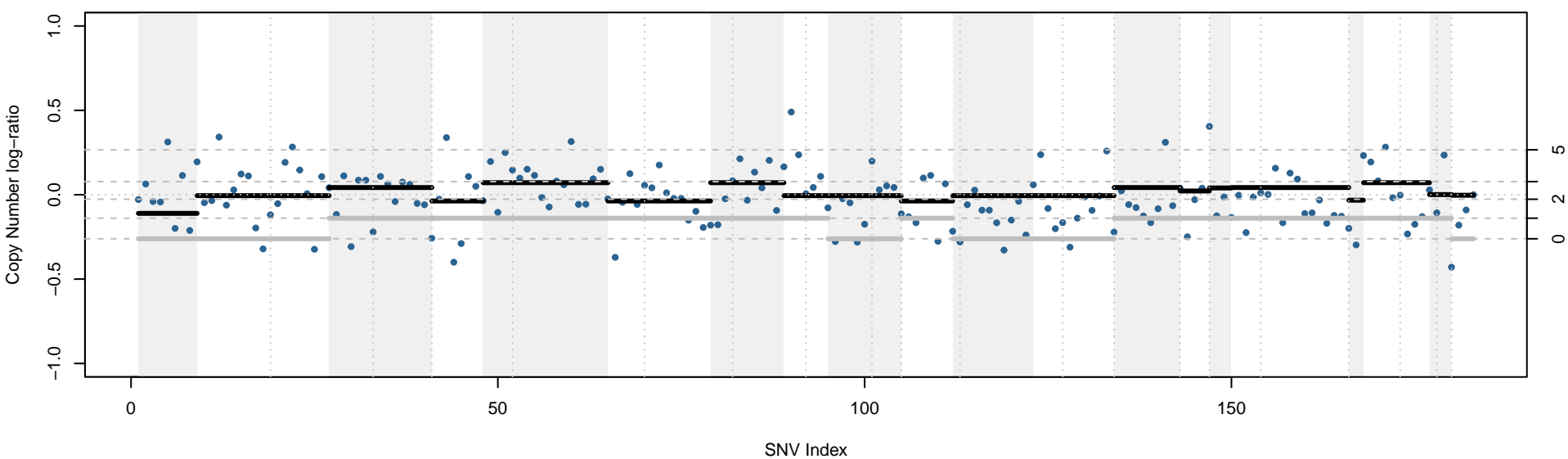
Purity: 0.15 Tumor ploidy: 2.248

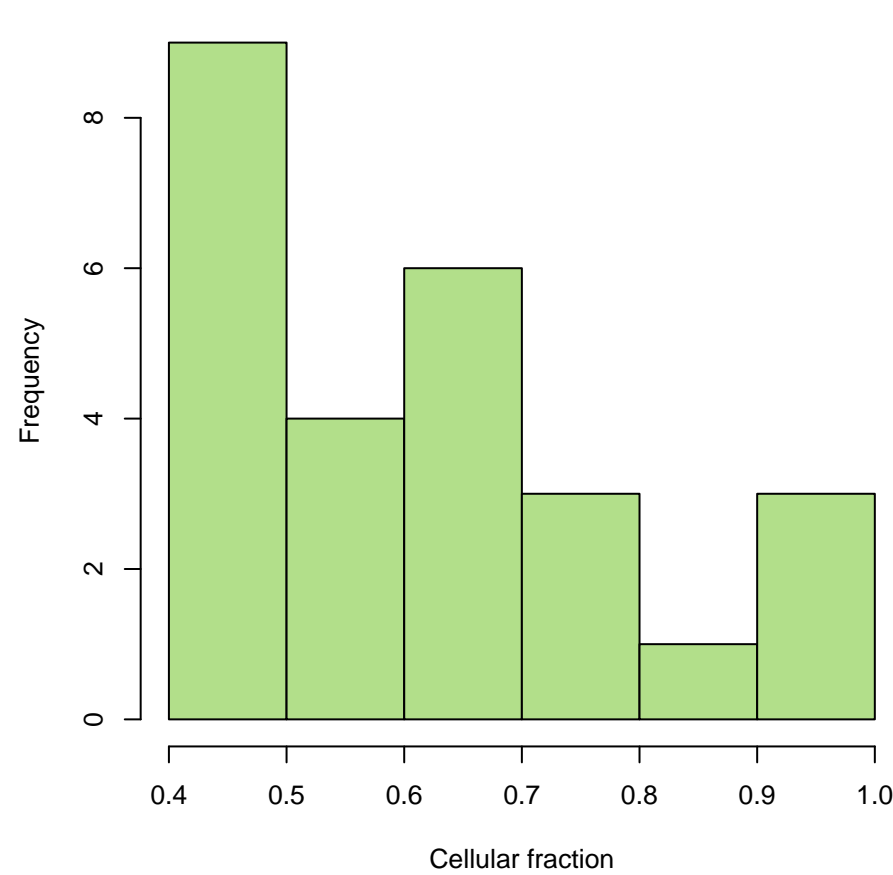
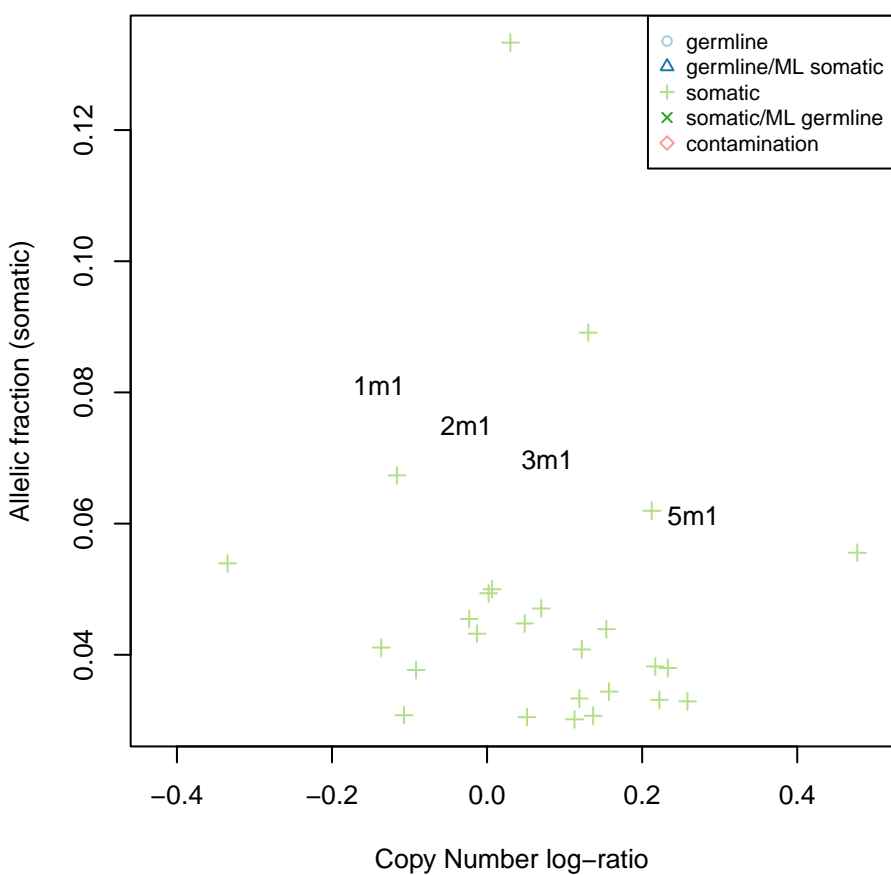
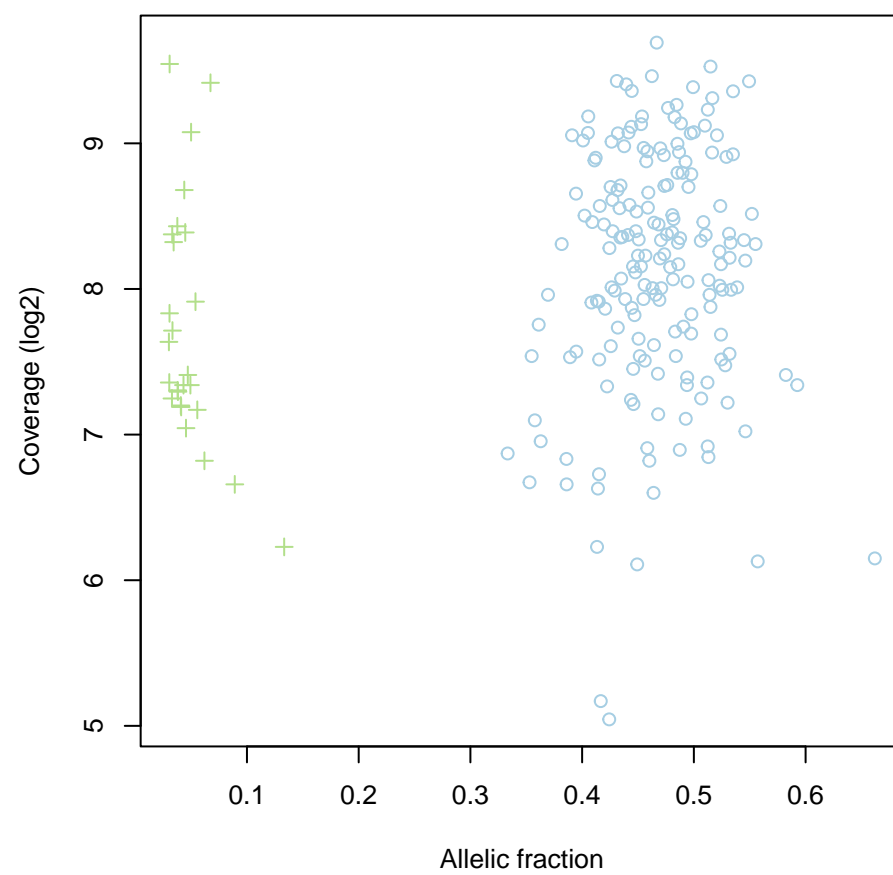
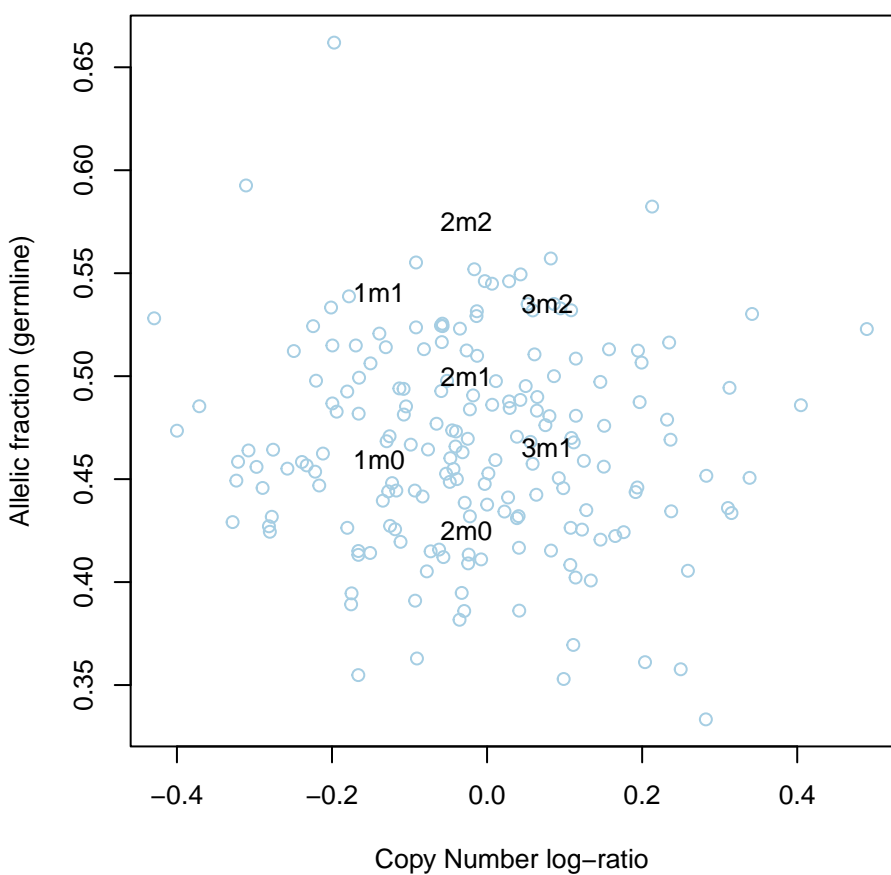


Purity: 0.15 Tumor ploidy: 2.248 SNV log-likelihood: 49.18 GoF: 96.3% Mean coverage: 317;651

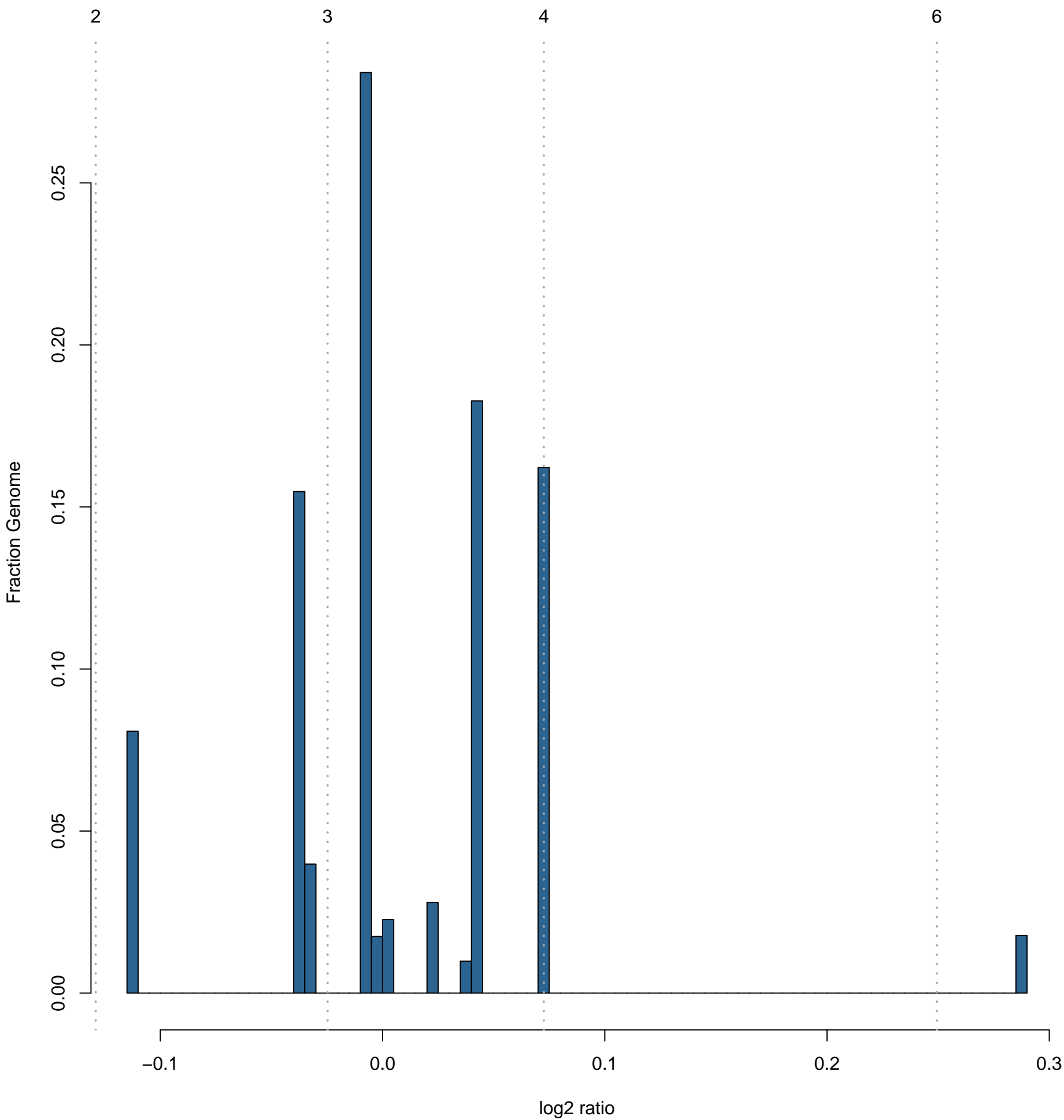


SCNA-fit log-likelihood: -9467.42

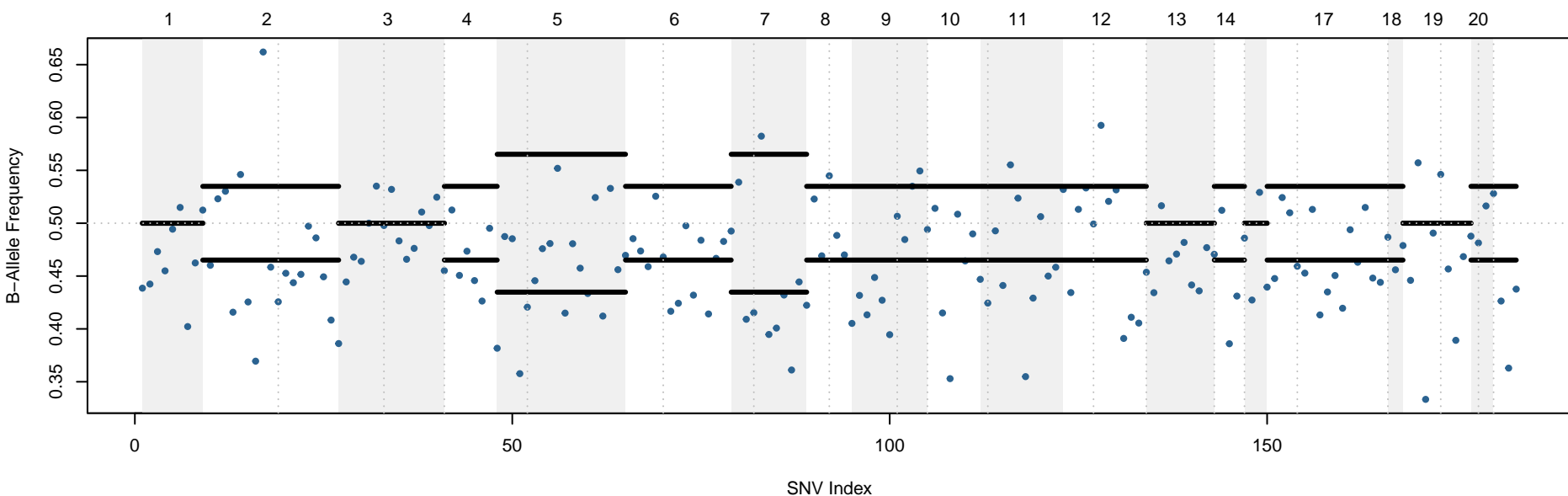




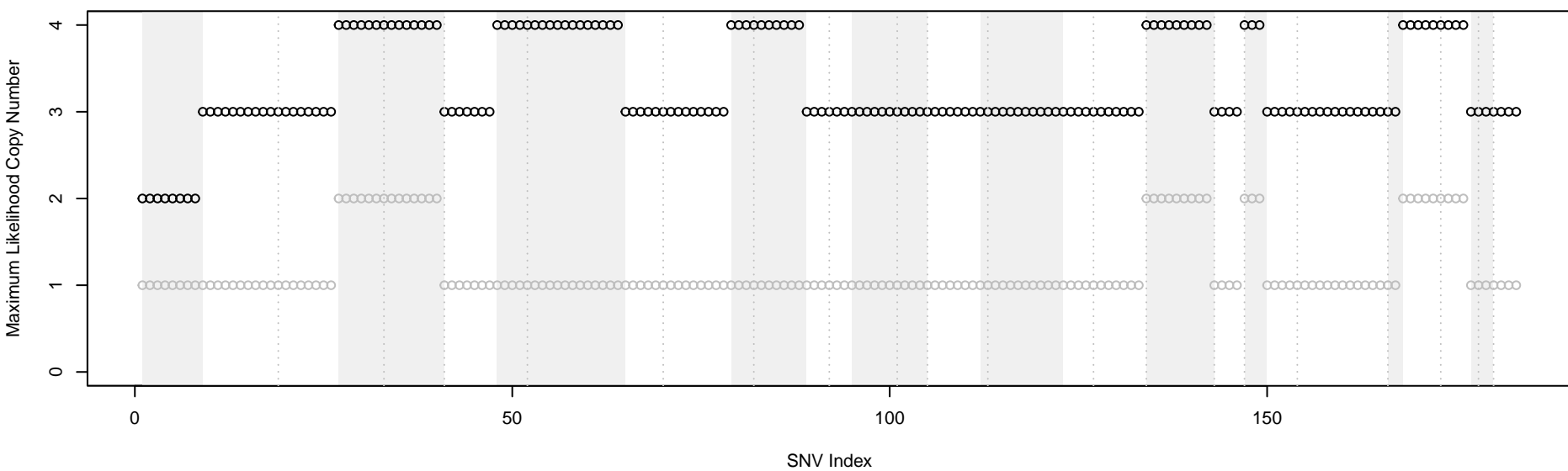
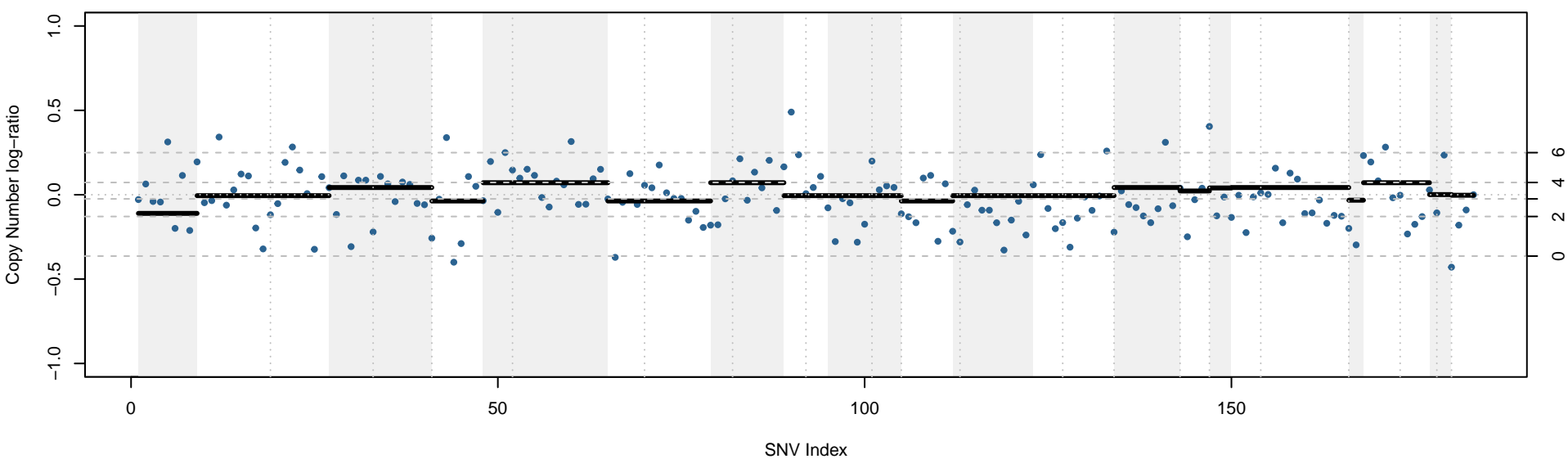
Purity: 0.15 Tumor ploidy: 3.248

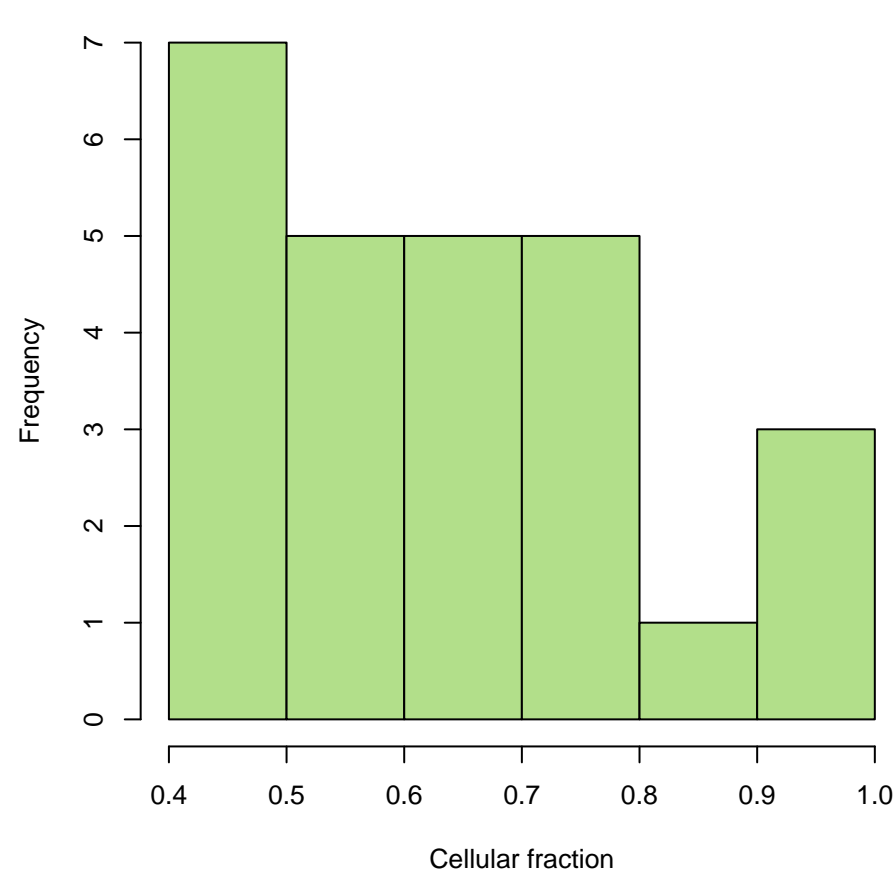
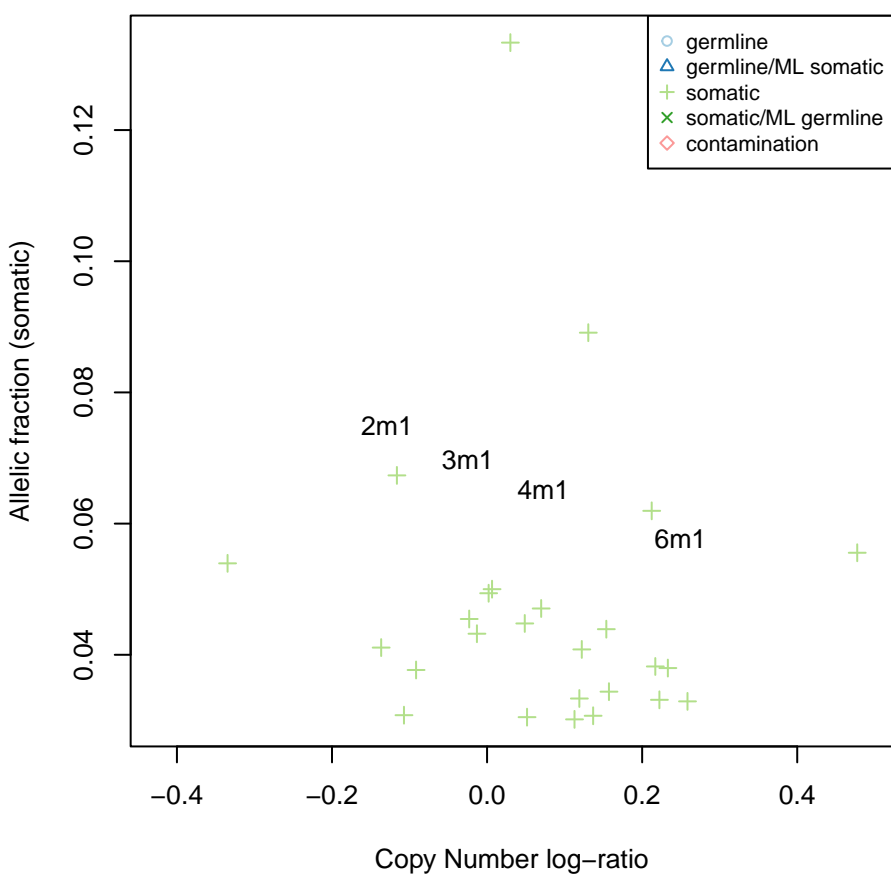
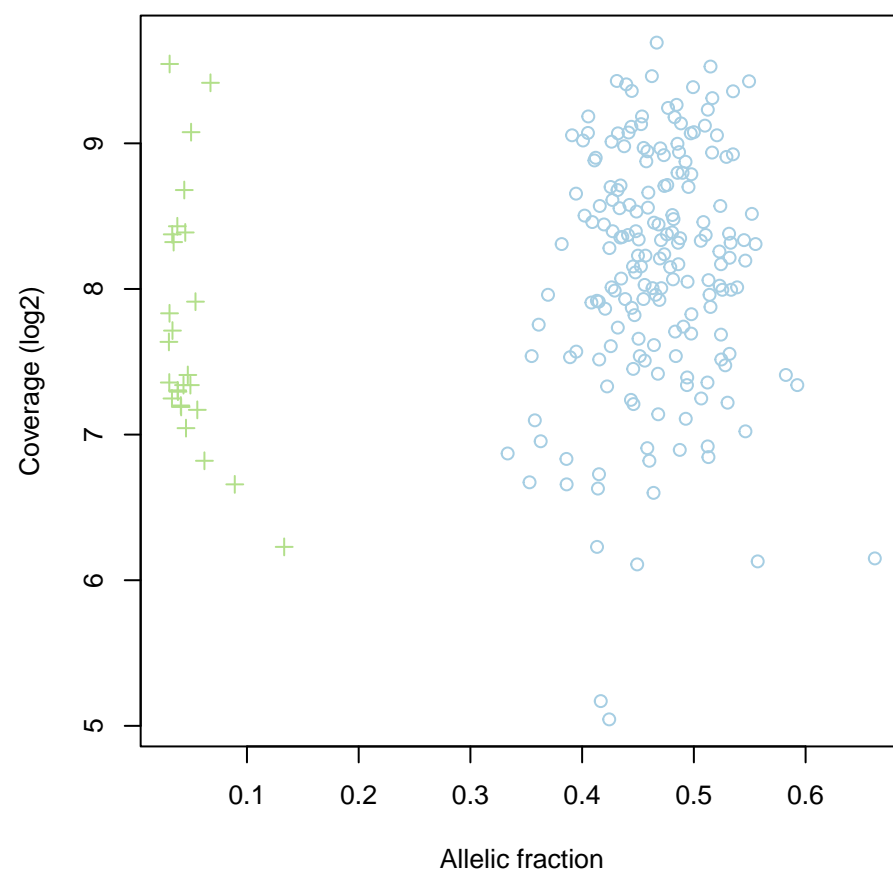
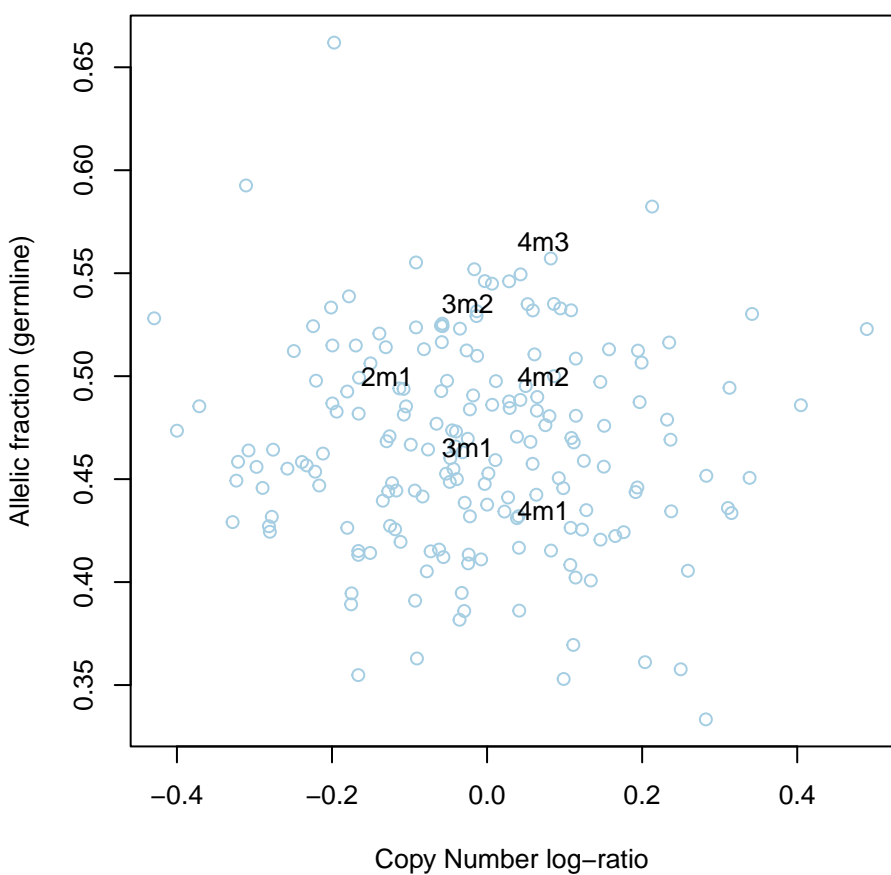


Purity: 0.15 Tumor ploidy: 3.248 SNV log-likelihood: -11.12 GoF: 96.6% Mean coverage: 317;651

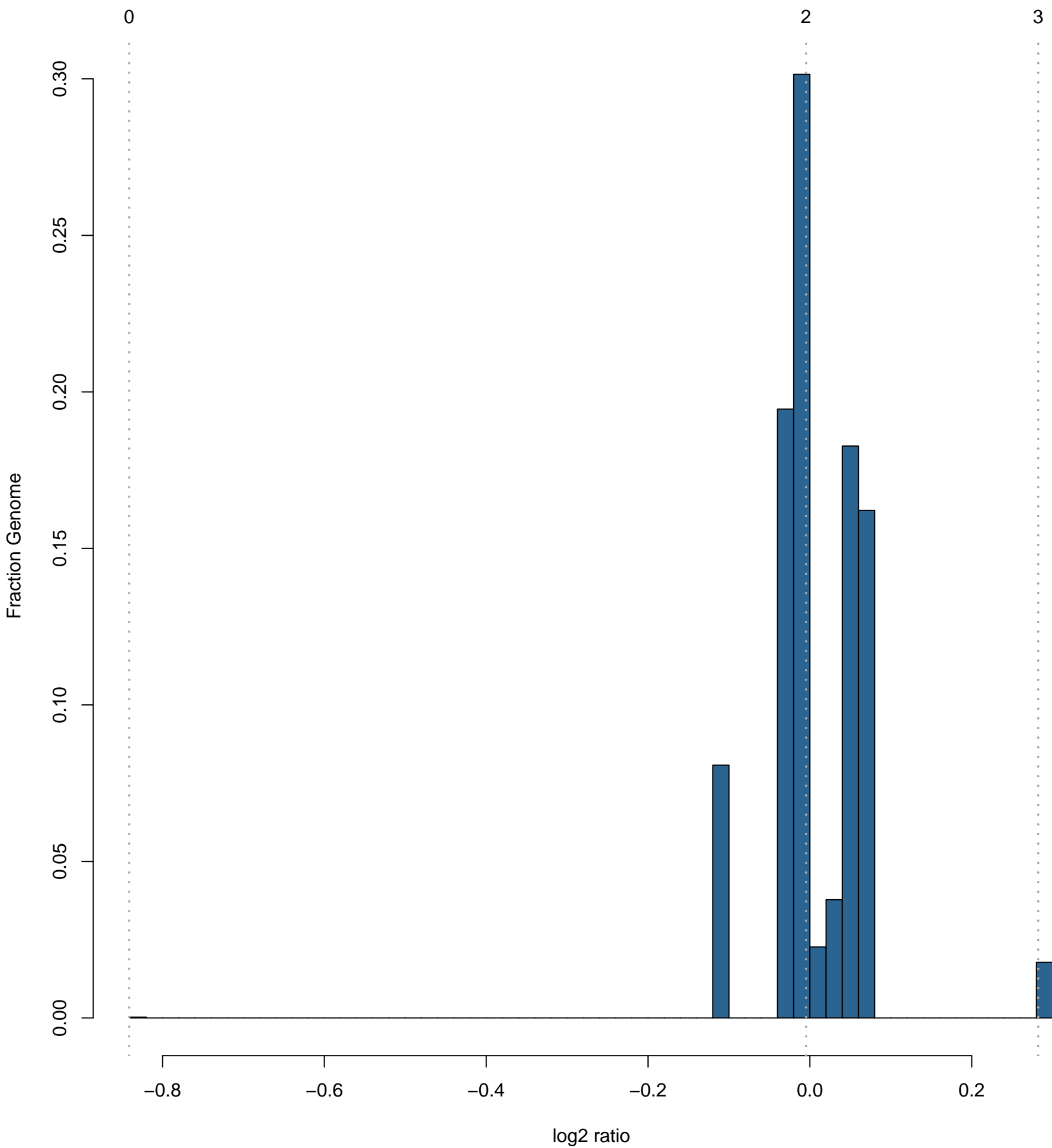


SCNA-fit log-likelihood: -9459.91

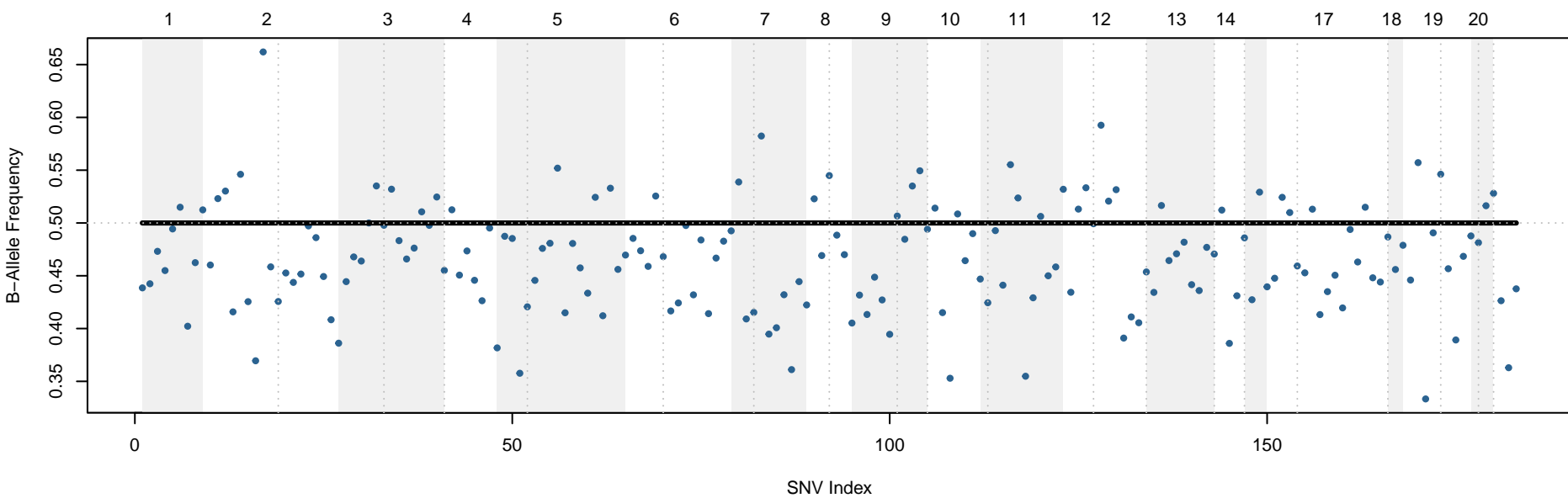




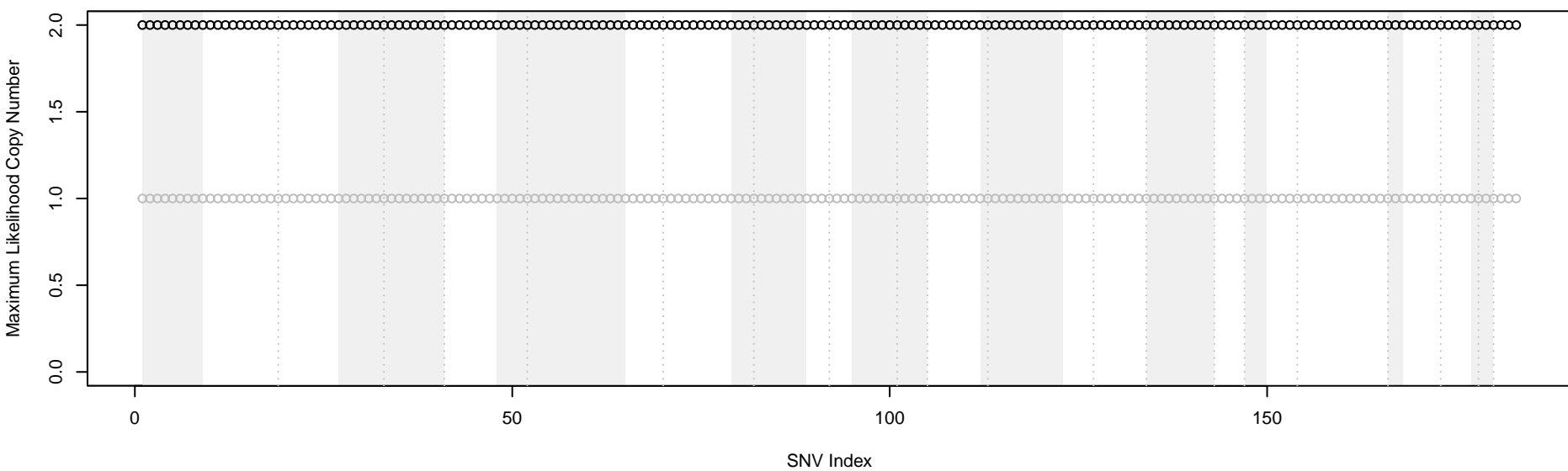
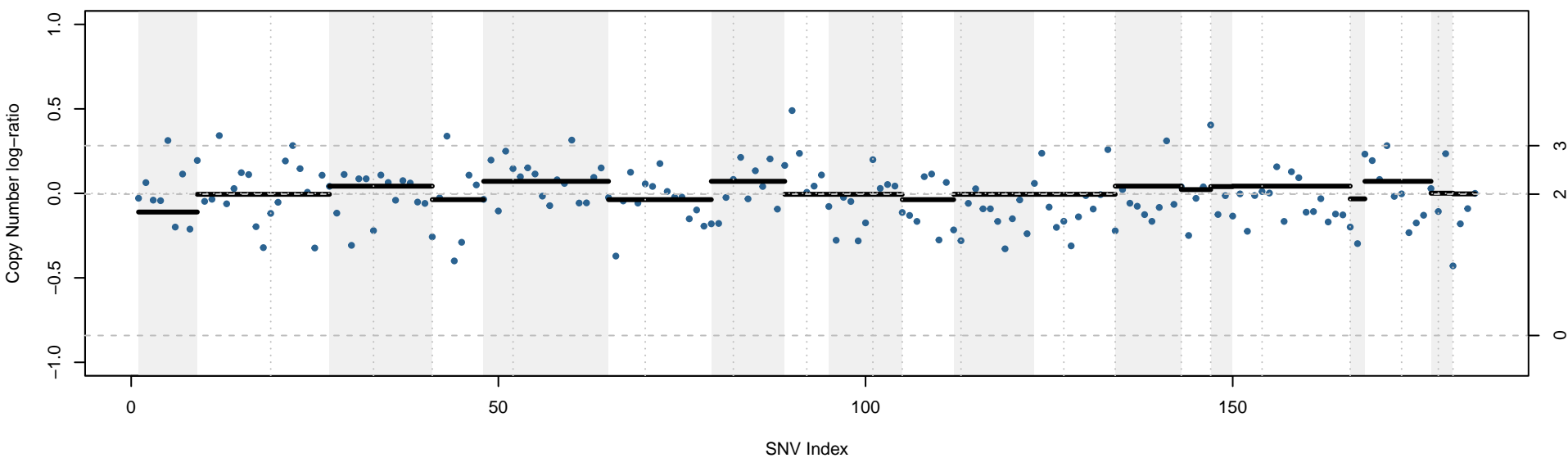
Purity: 0.44 Tumor ploidy: 2.015



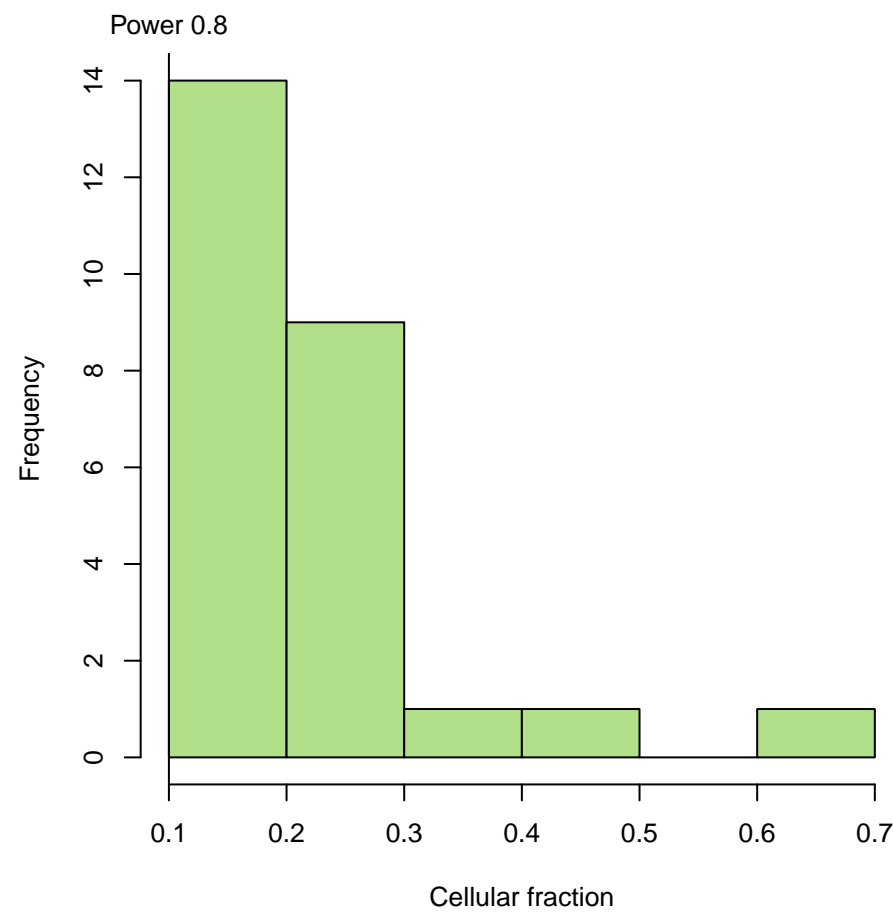
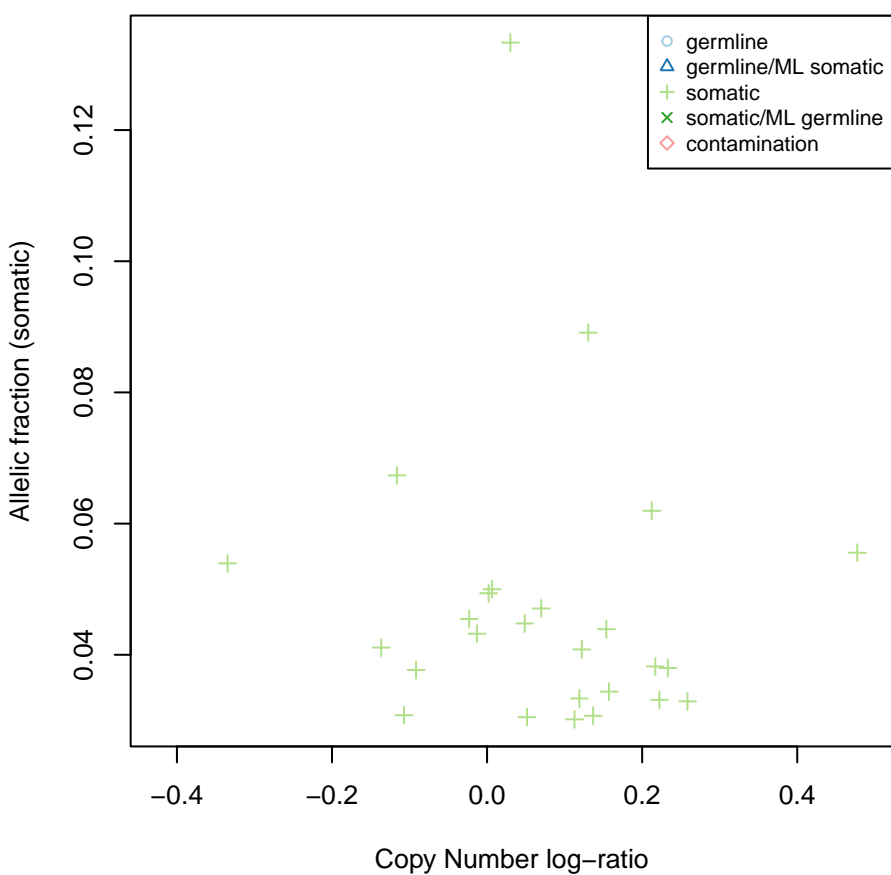
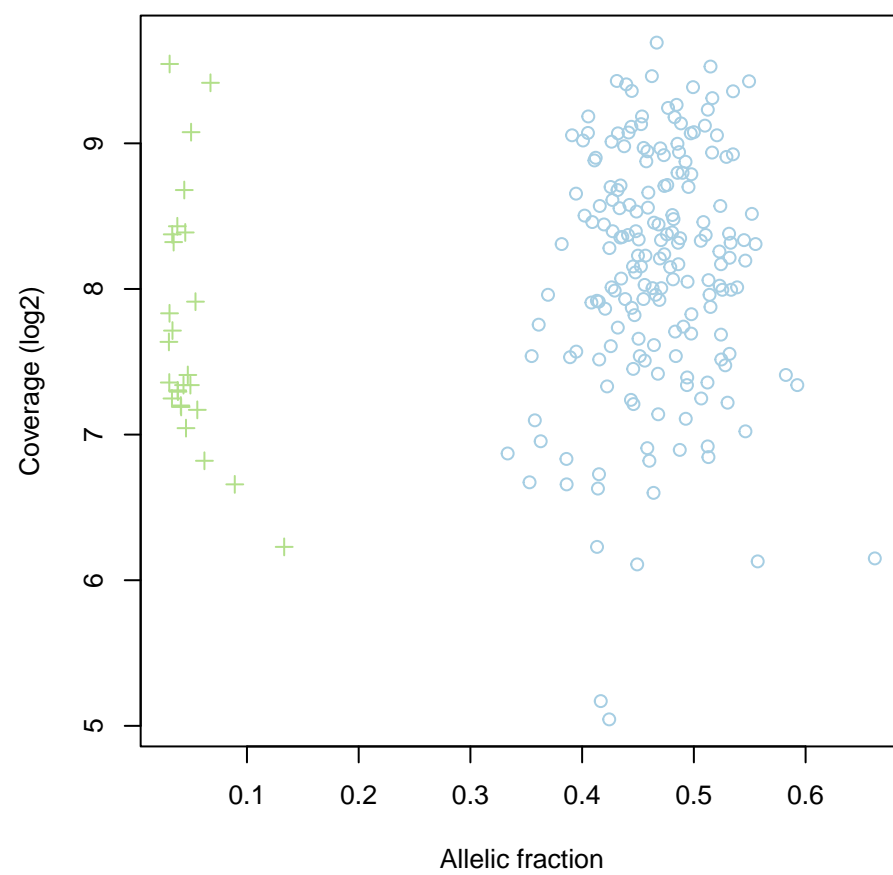
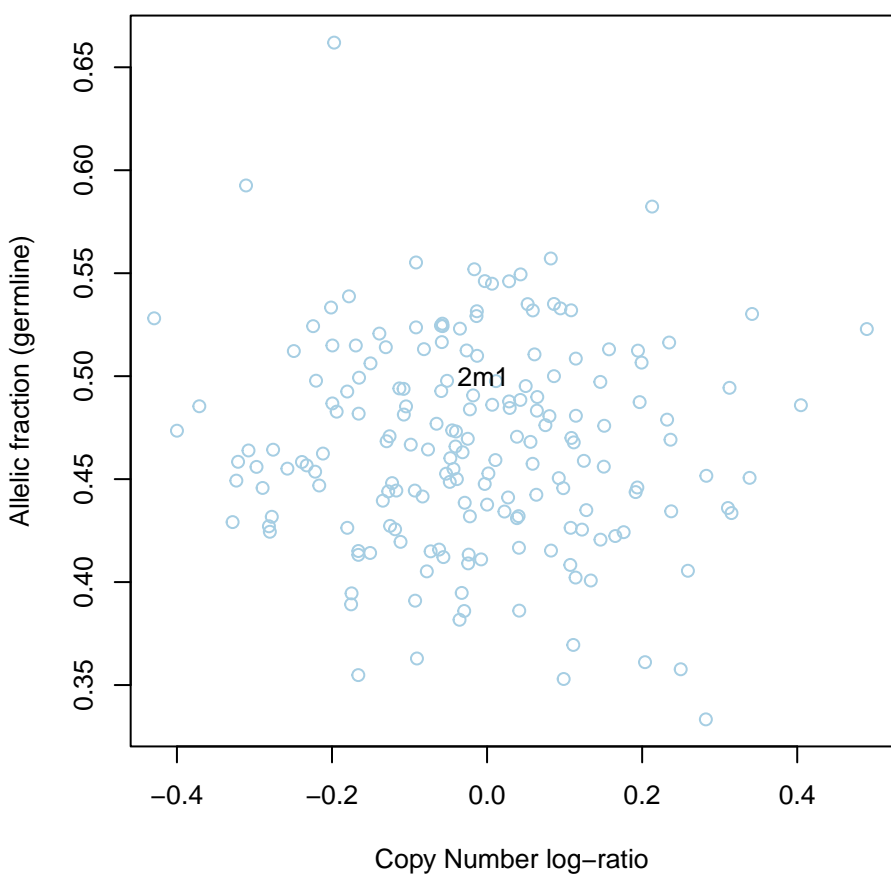
Purity: 0.44 Tumor ploidy: 2.015 SNV log-likelihood: 12.24 GoF: 84.4% Mean coverage: 317;651



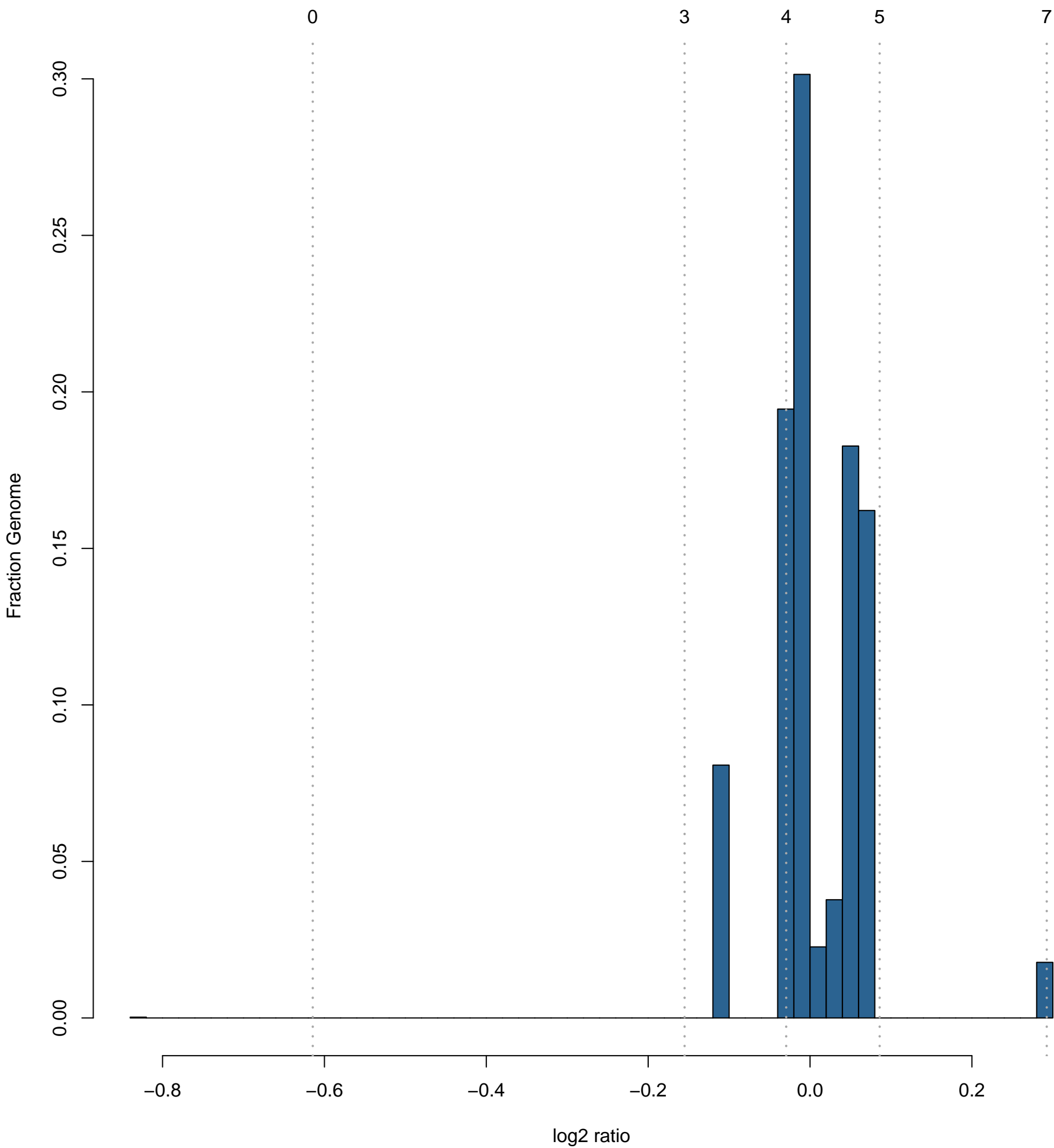
SCNA-fit log-likelihood: -9601.72



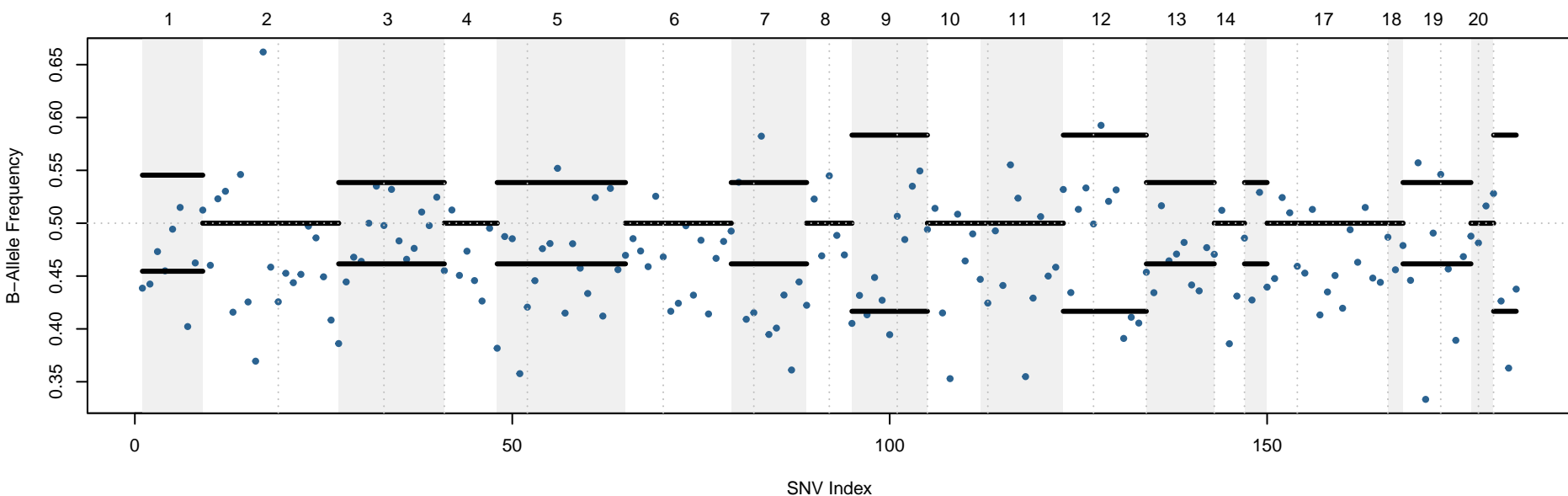




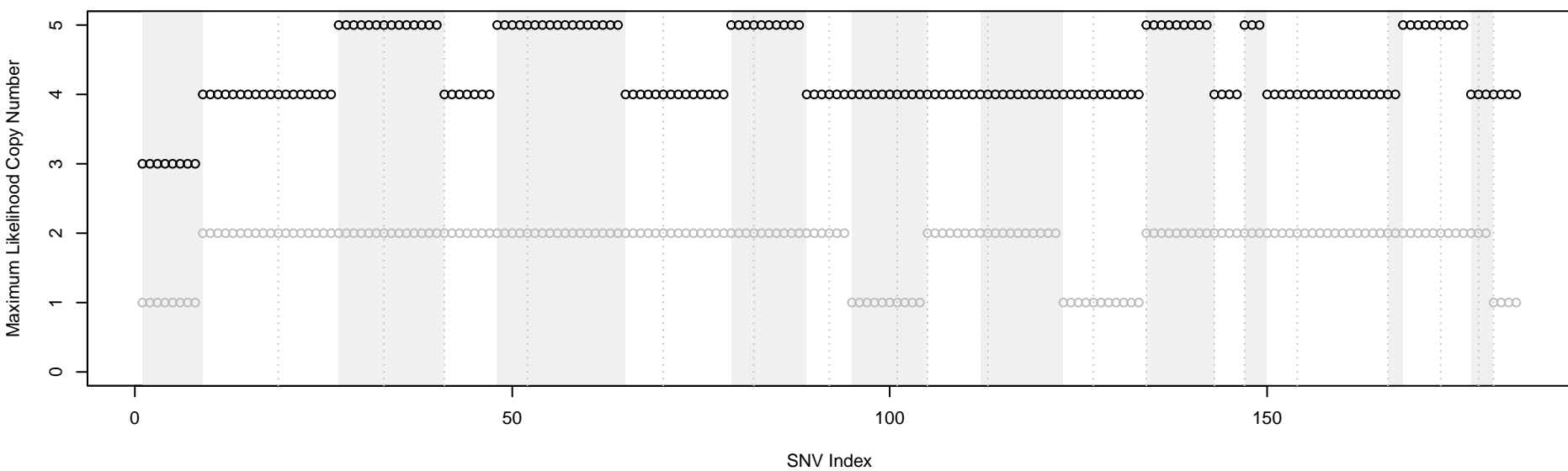
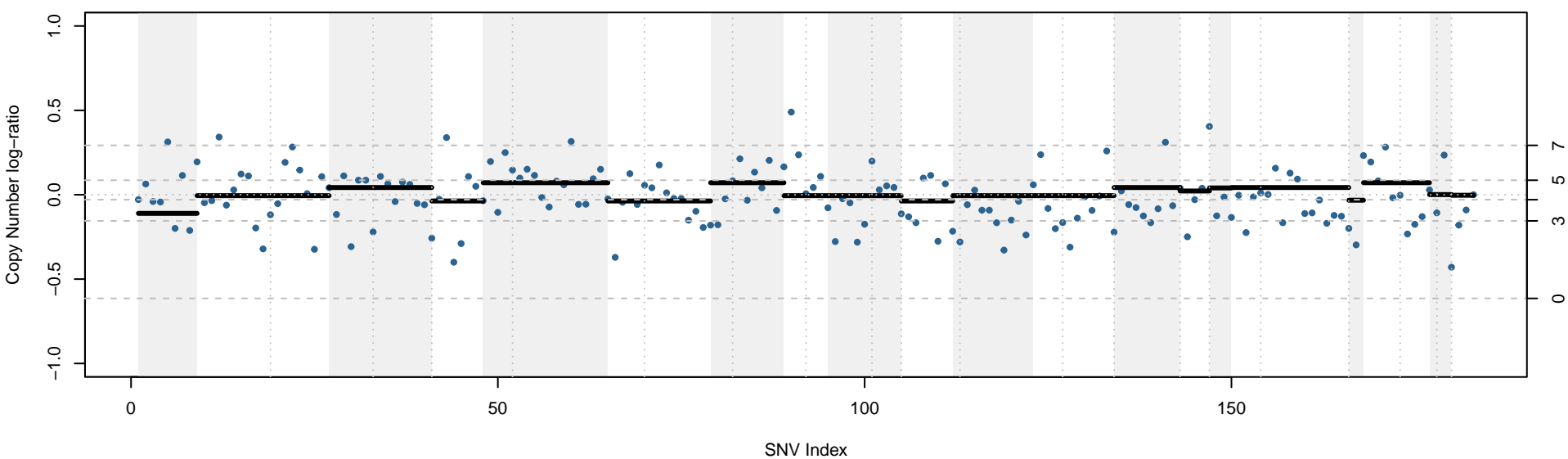
Purity: 0.2 Tumor ploidy: 4.248

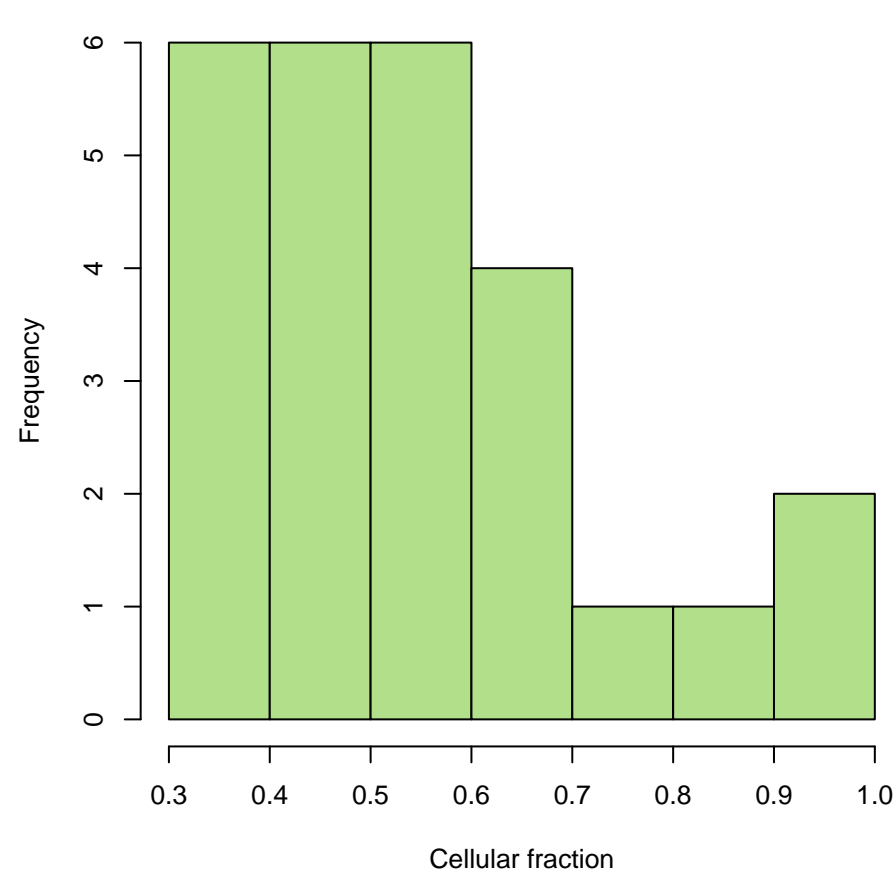
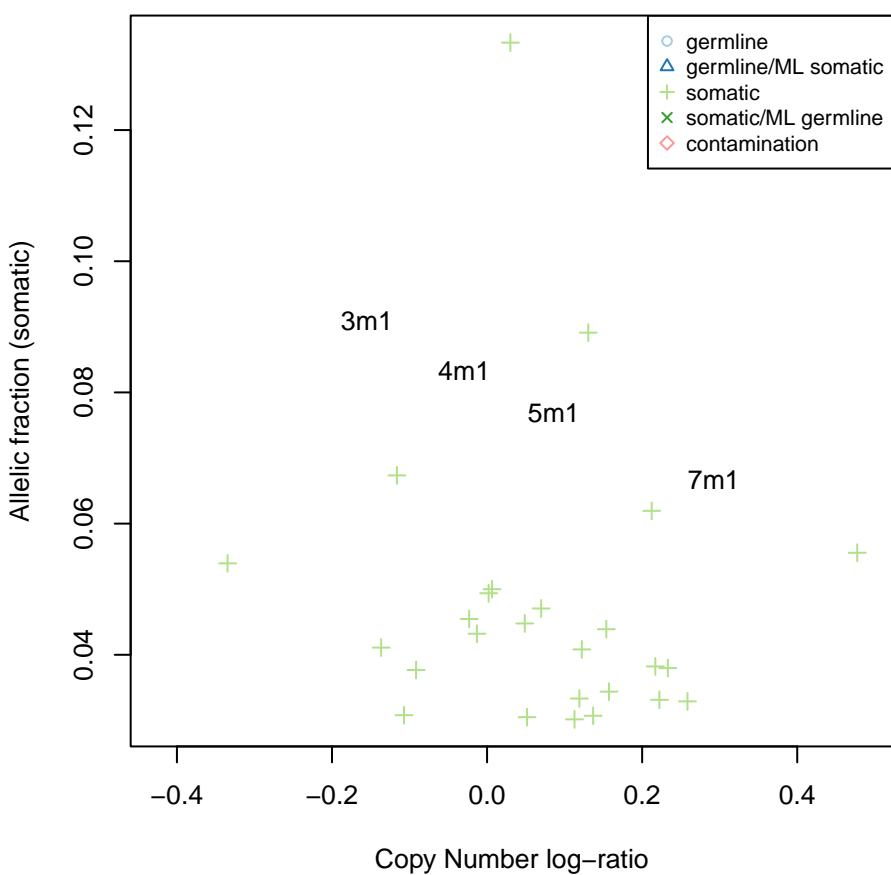
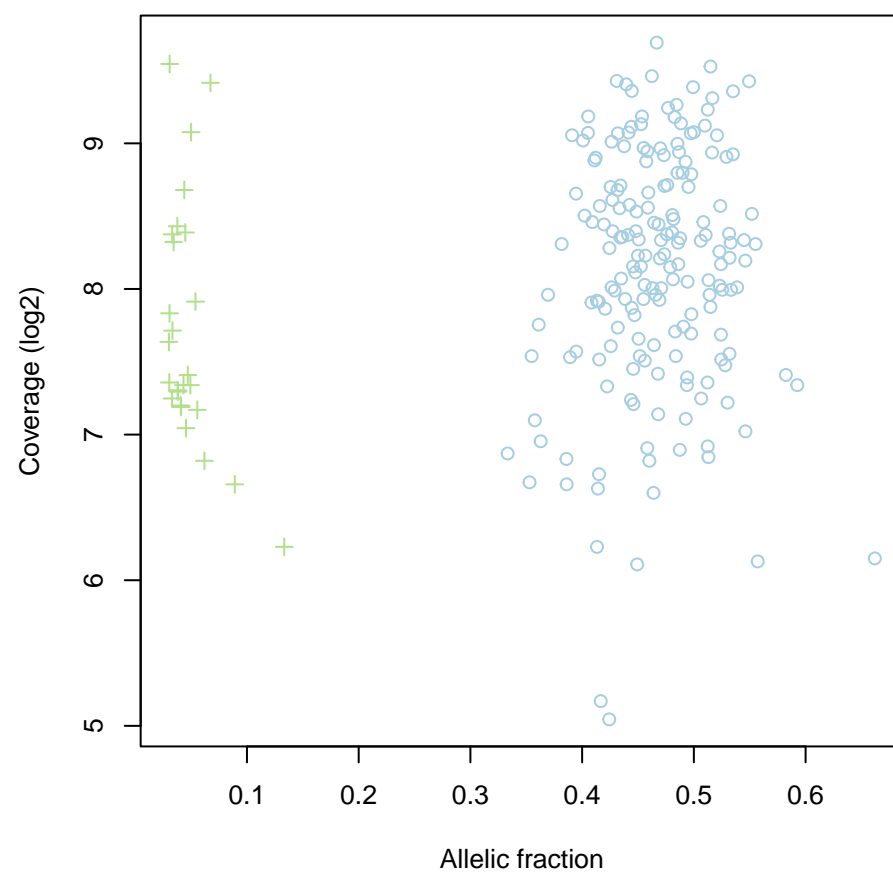
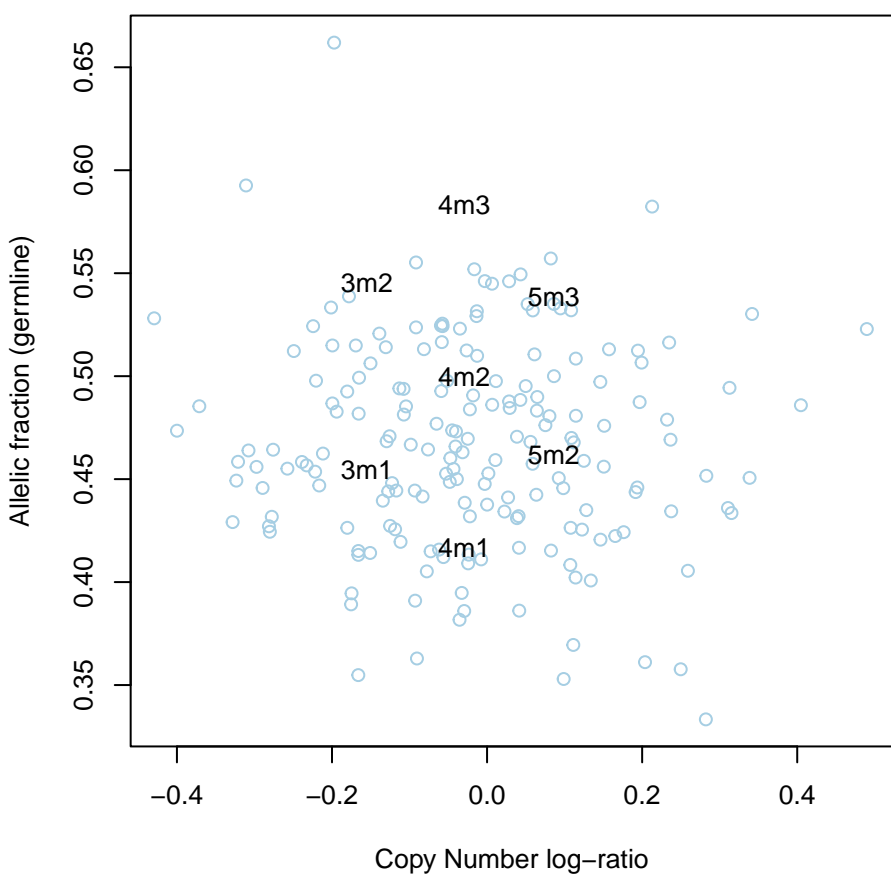


Purity: 0.2 Tumor ploidy: 4.248 SNV log-likelihood: -66.45 GoF: 95.3% Mean coverage: 317;651

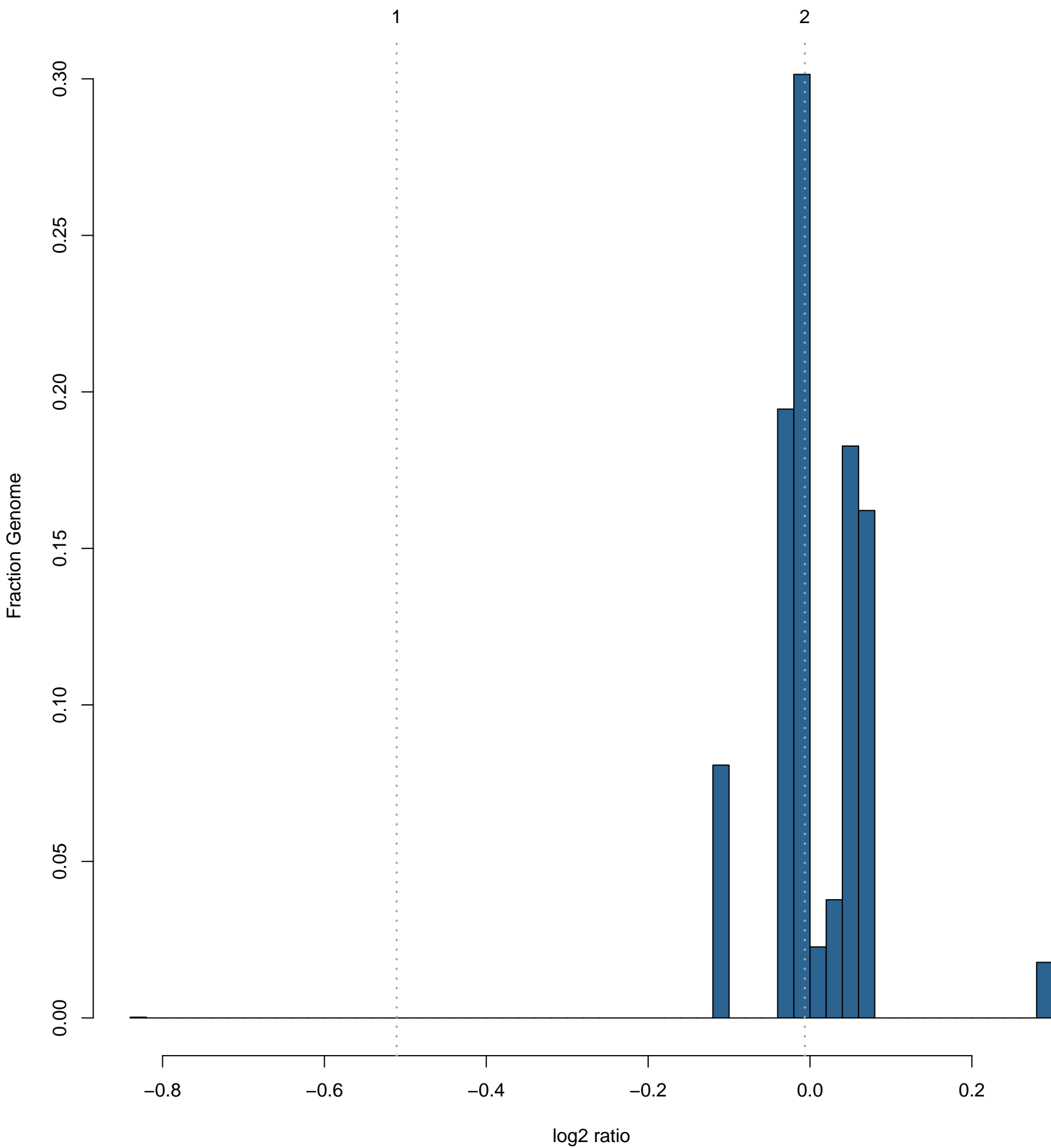


SCNA-fit log-likelihood: -9482.72

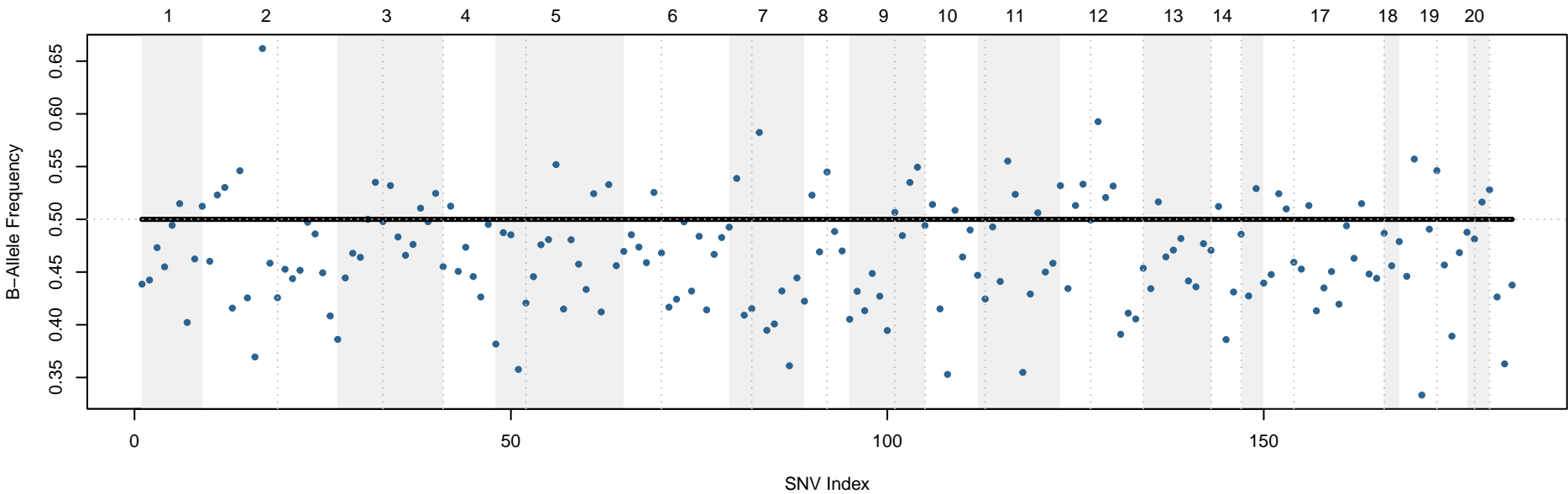




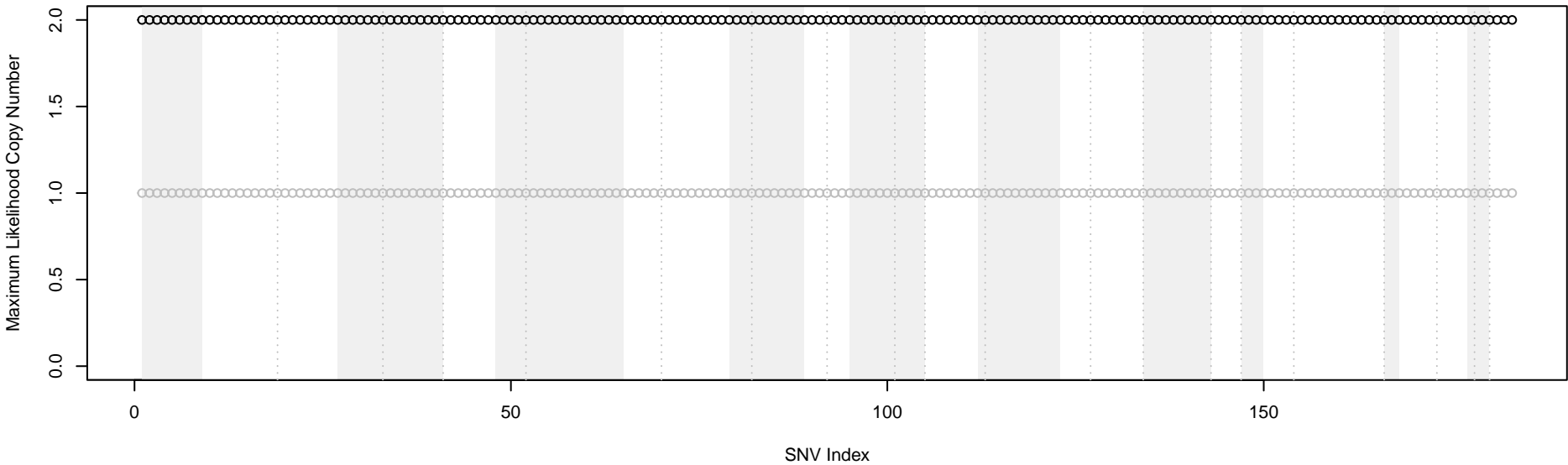
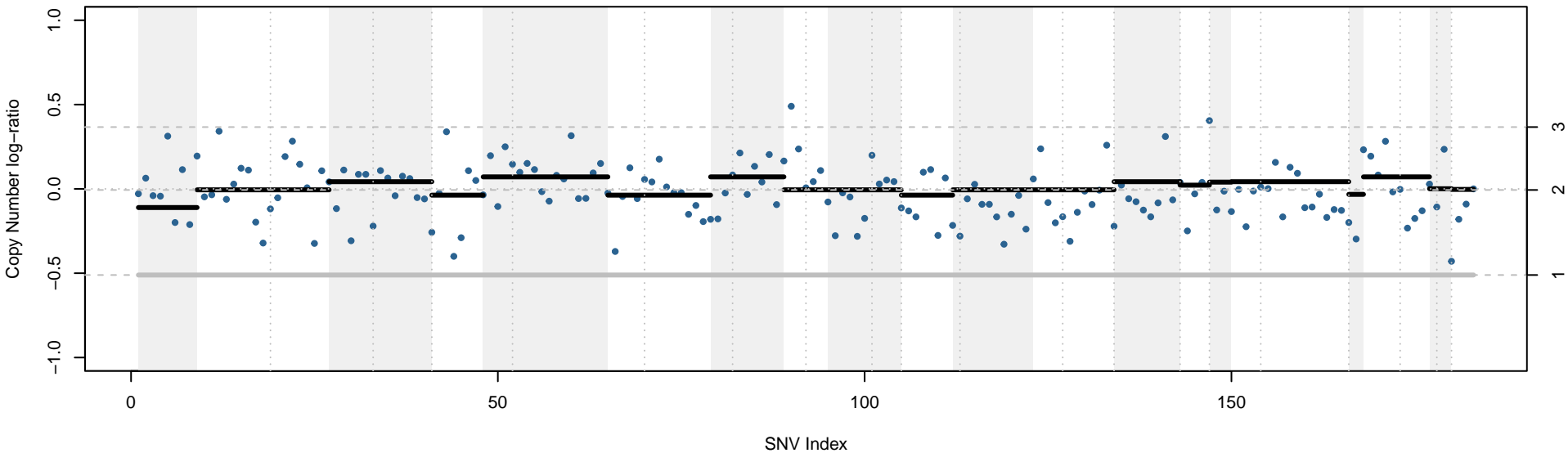
Purity: 0.59 Tumor ploidy: 2.015

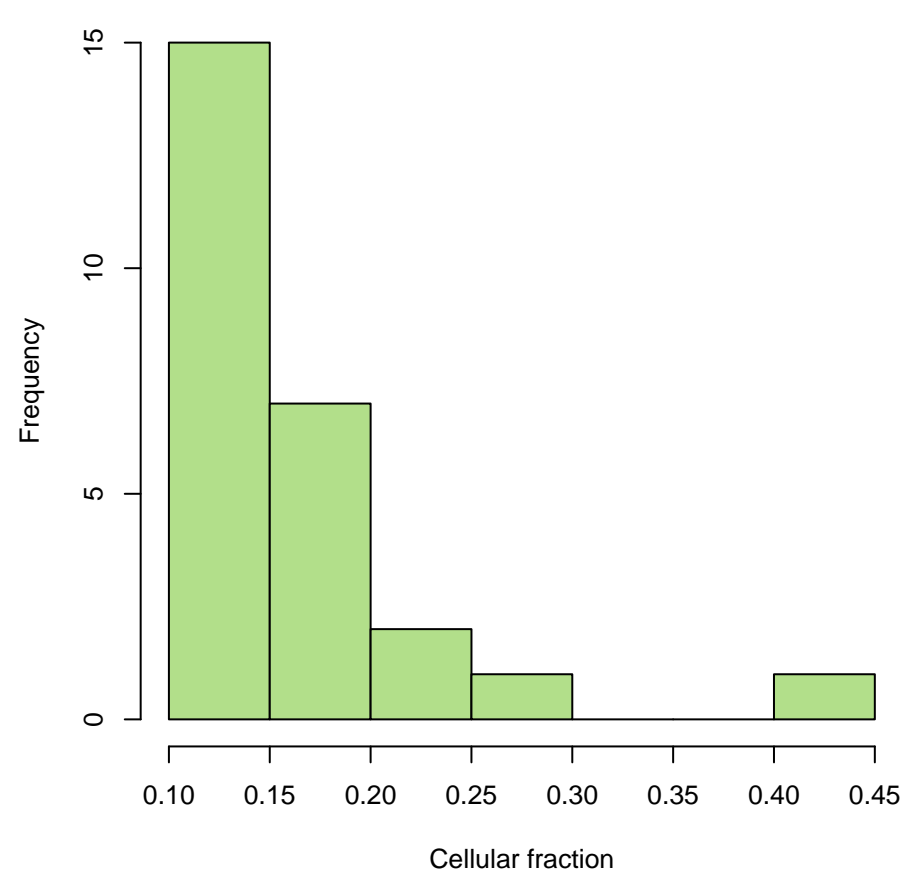
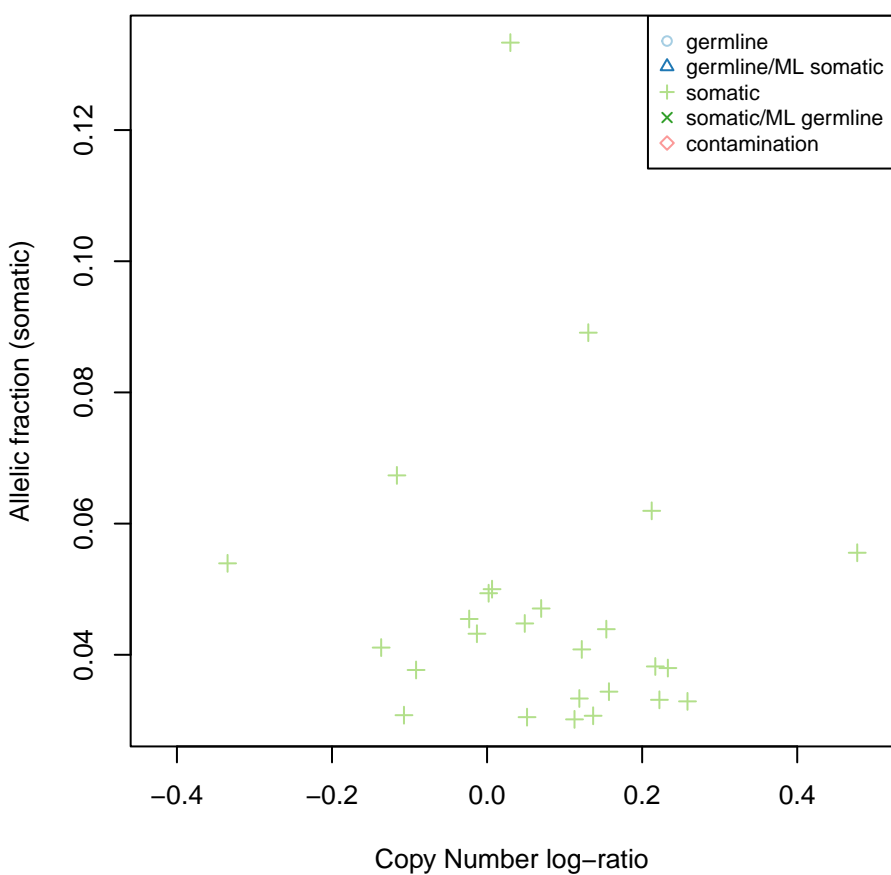
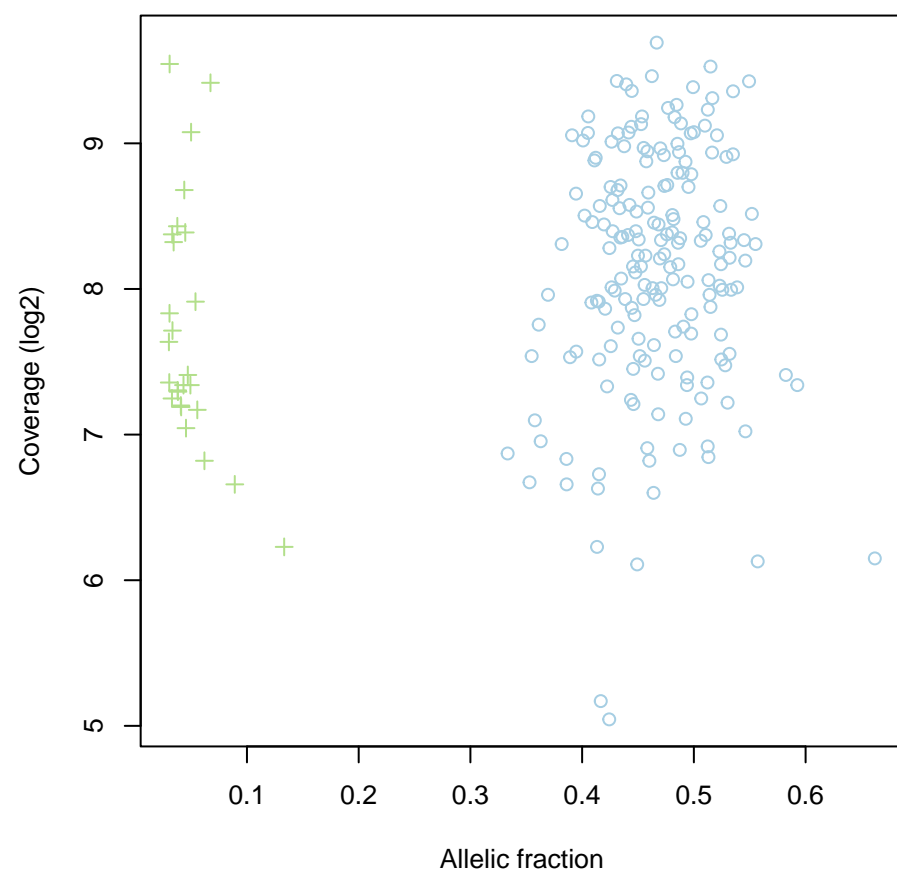
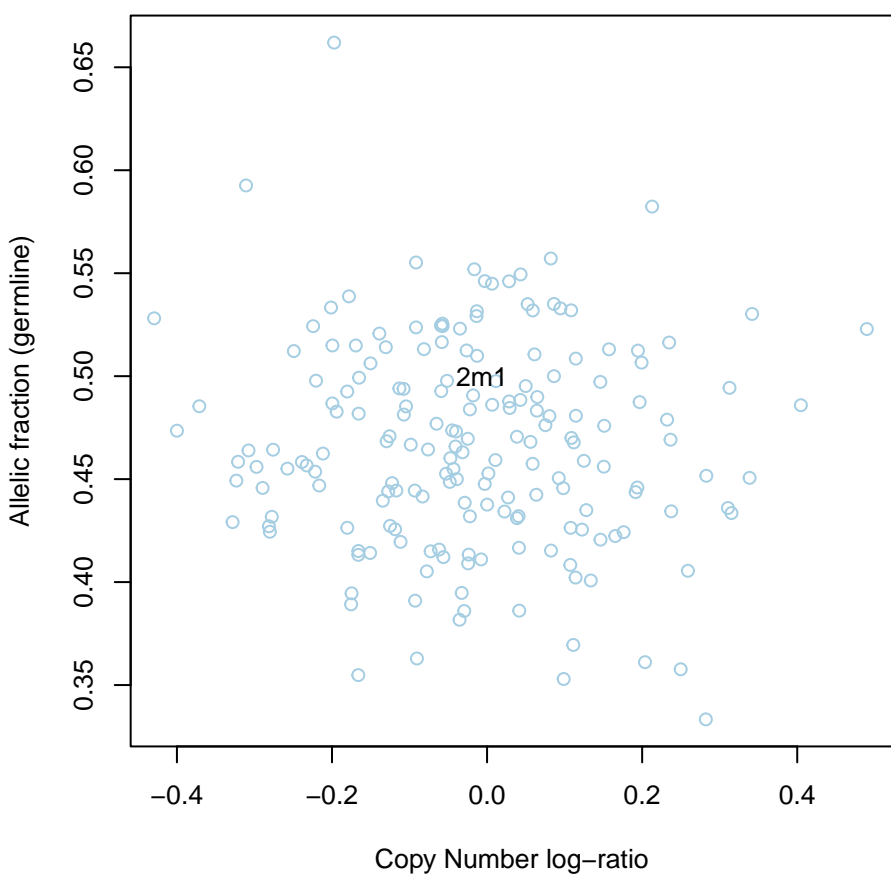


Purity: 0.59 Tumor ploidy: 2.015 SNV log-likelihood: -38.74 GoF: 75% Mean coverage: 317;651

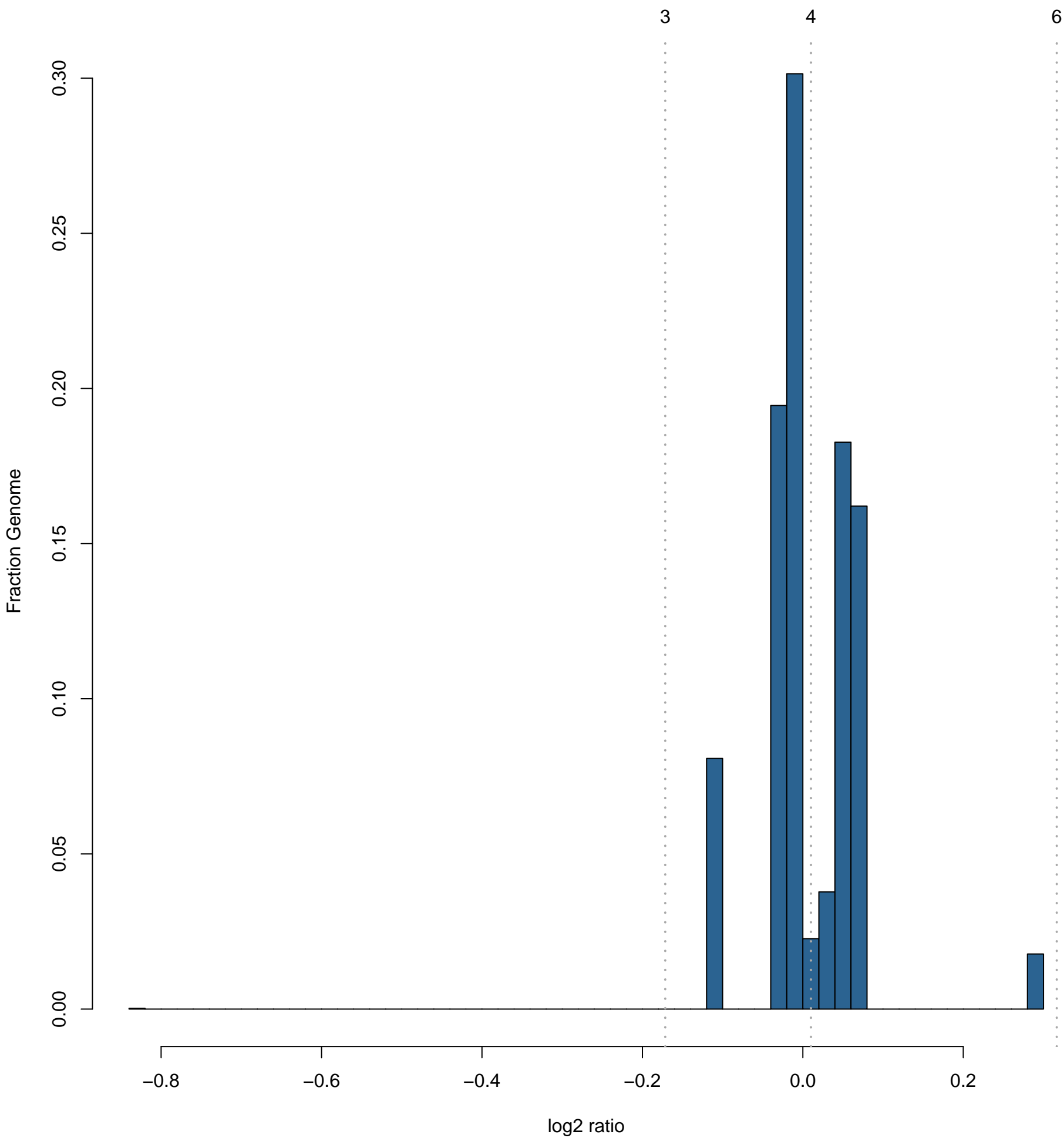


SCNA-fit log-likelihood: -9612.43



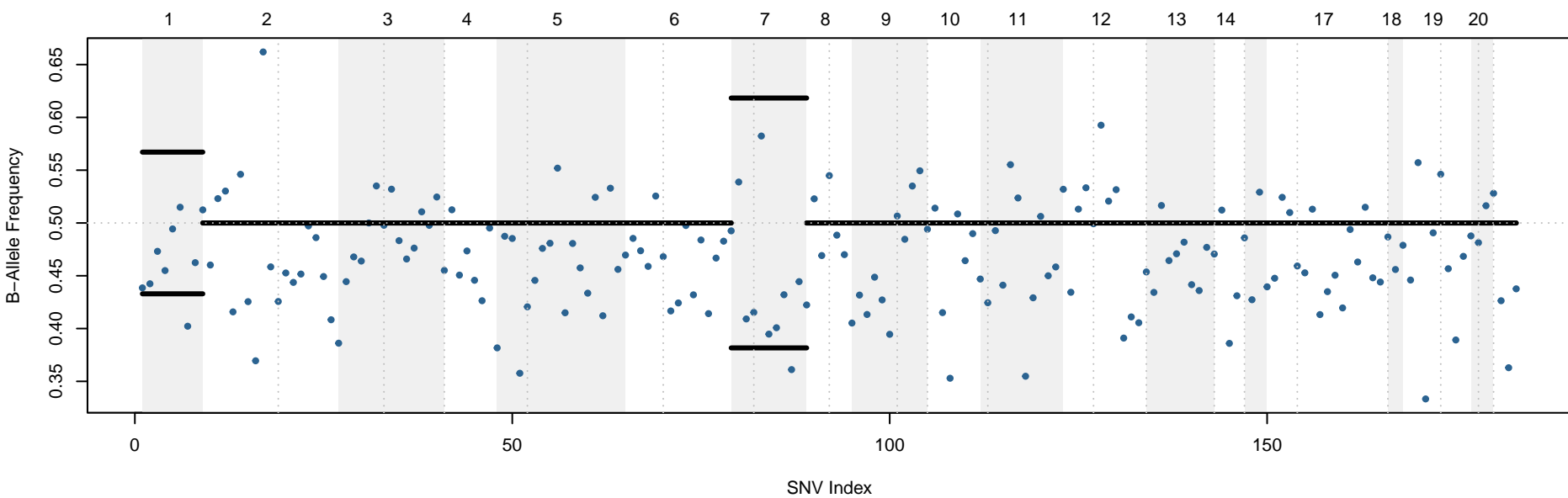


Purity: 0.31 Tumor ploidy: 3.941

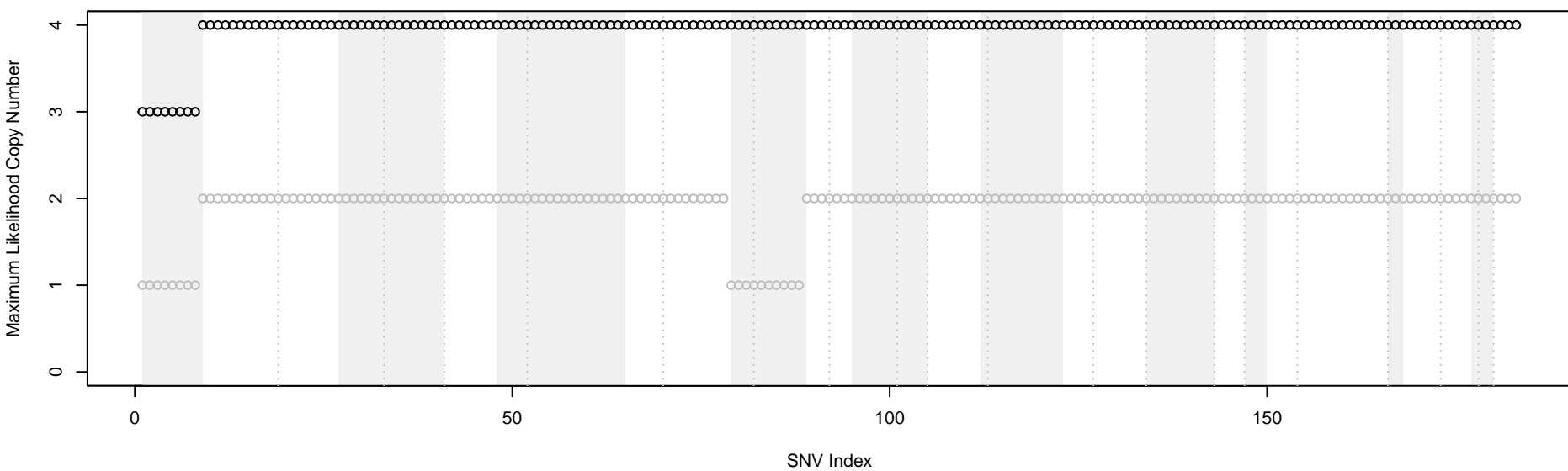
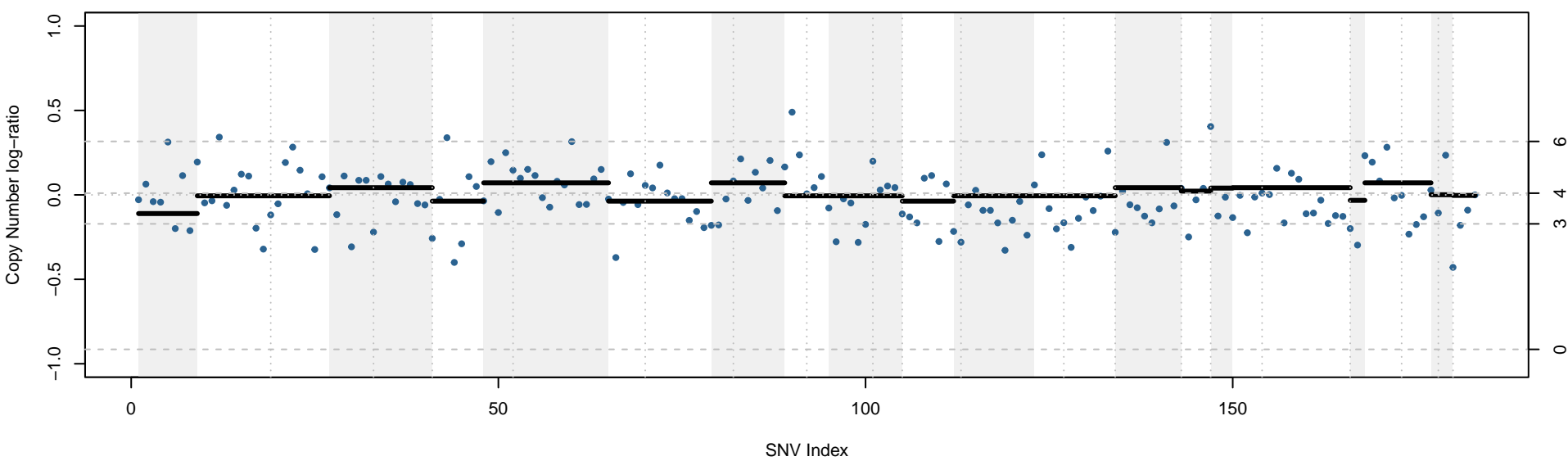


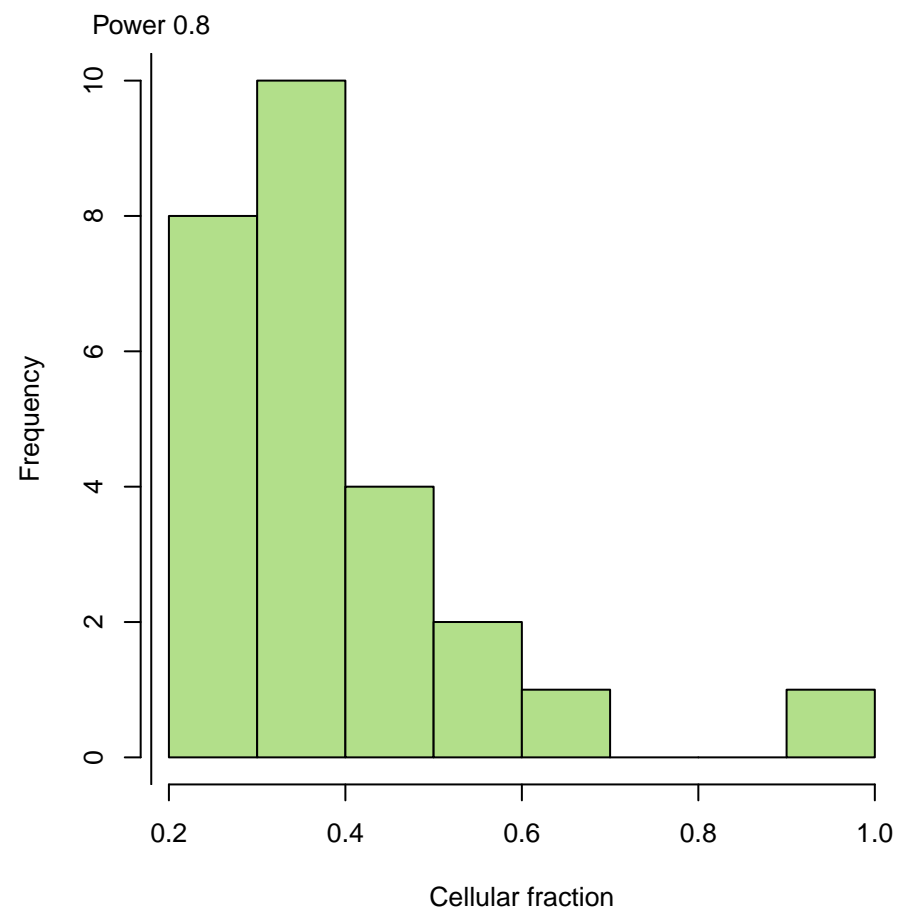
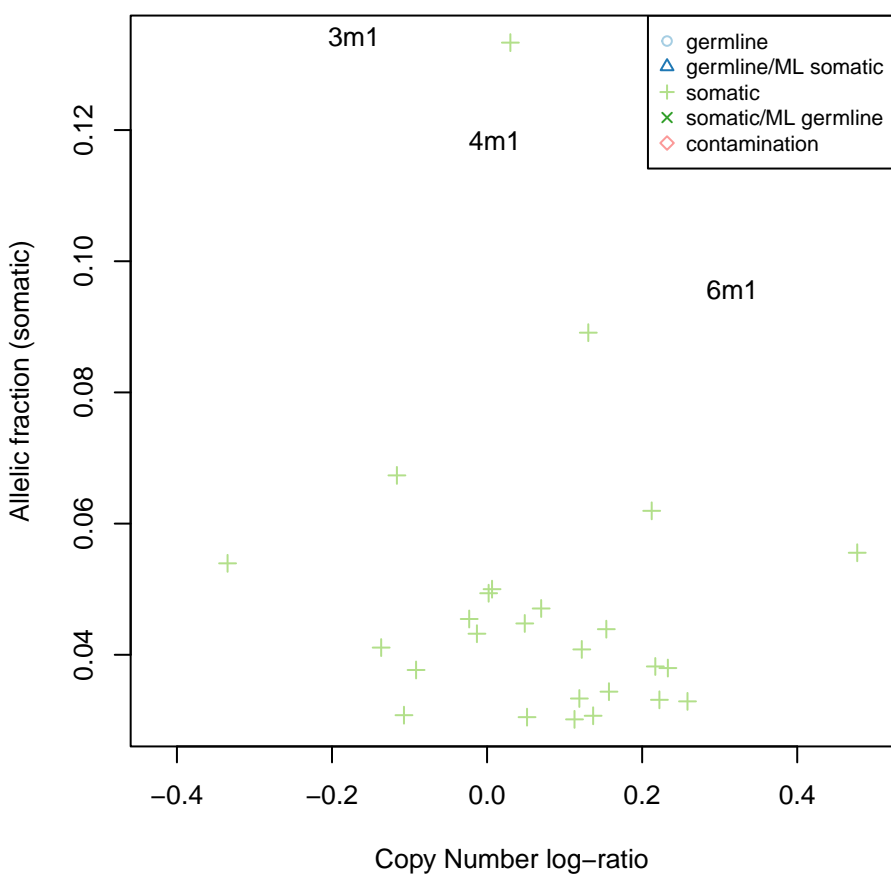
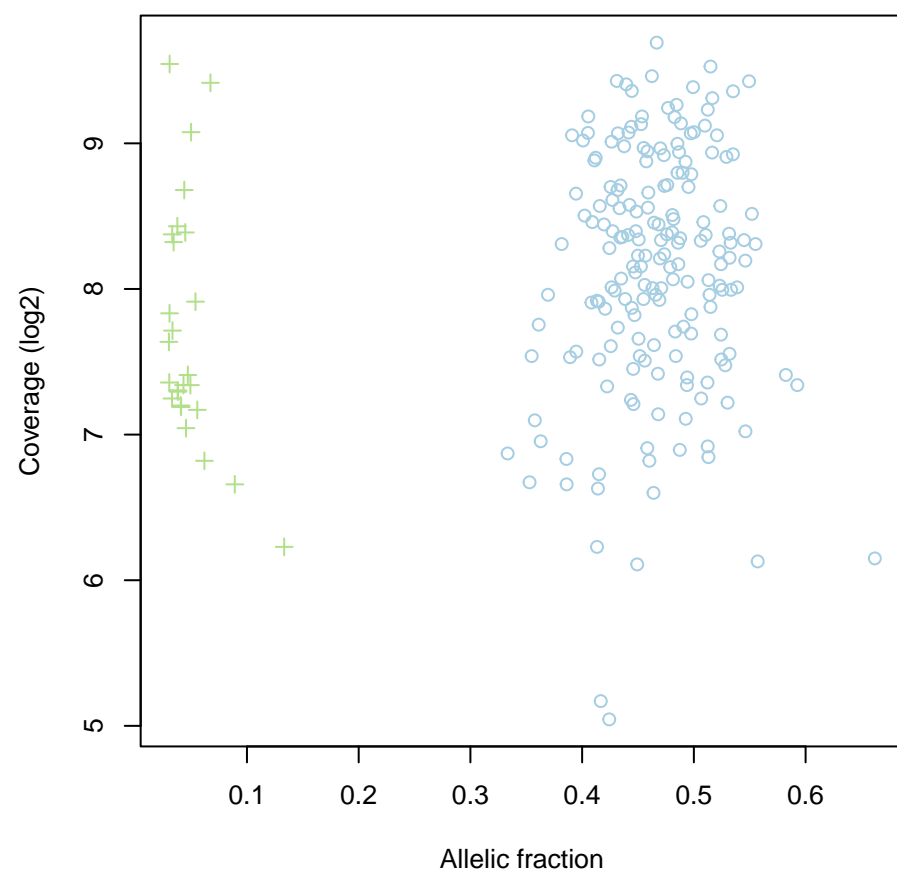
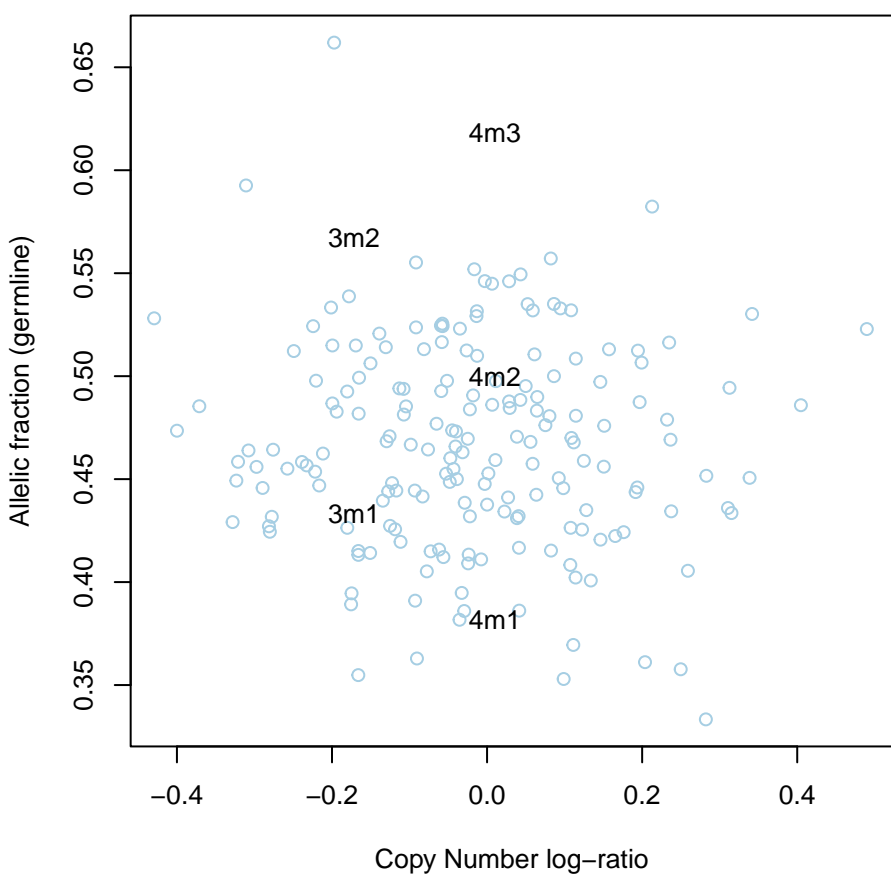


Purity: 0.31 Tumor ploidy: 3.941 SNV log-likelihood: -71.74 GoF: 92.2% Mean coverage: 317;651

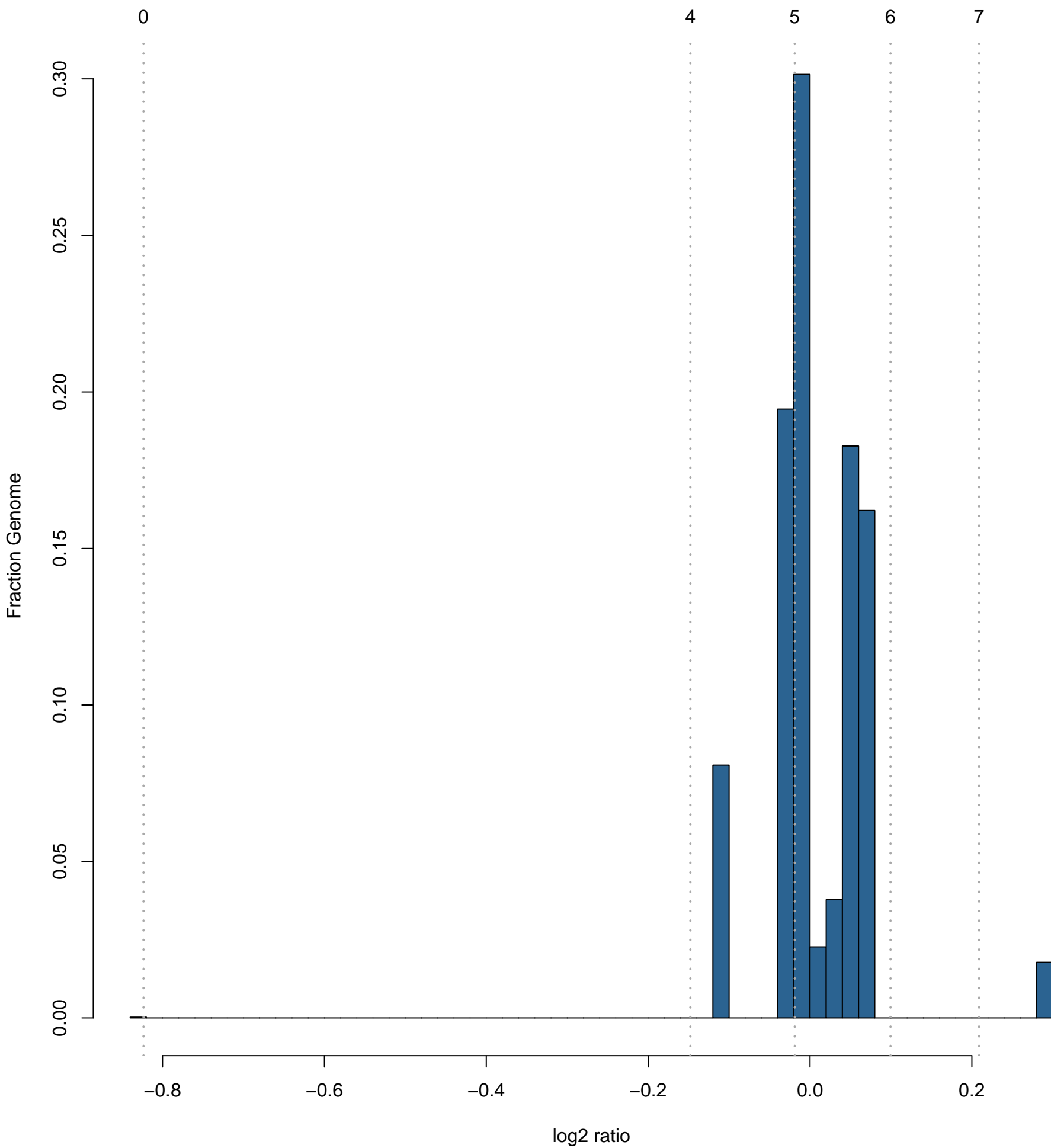


SCNA-fit log-likelihood: -9541.56

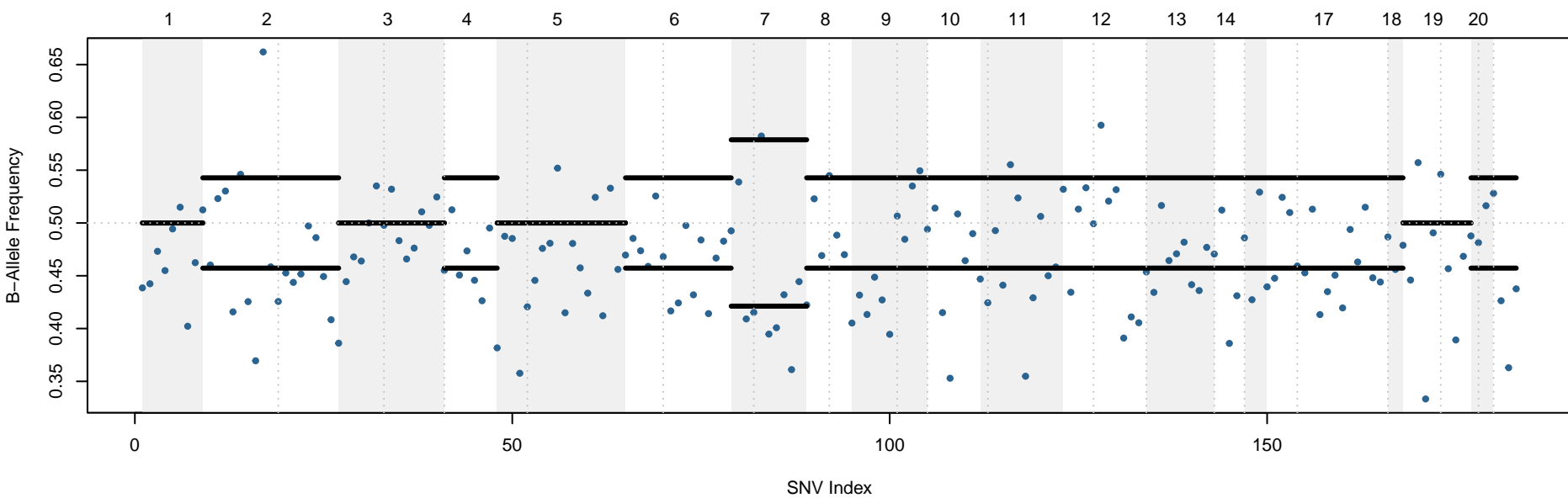




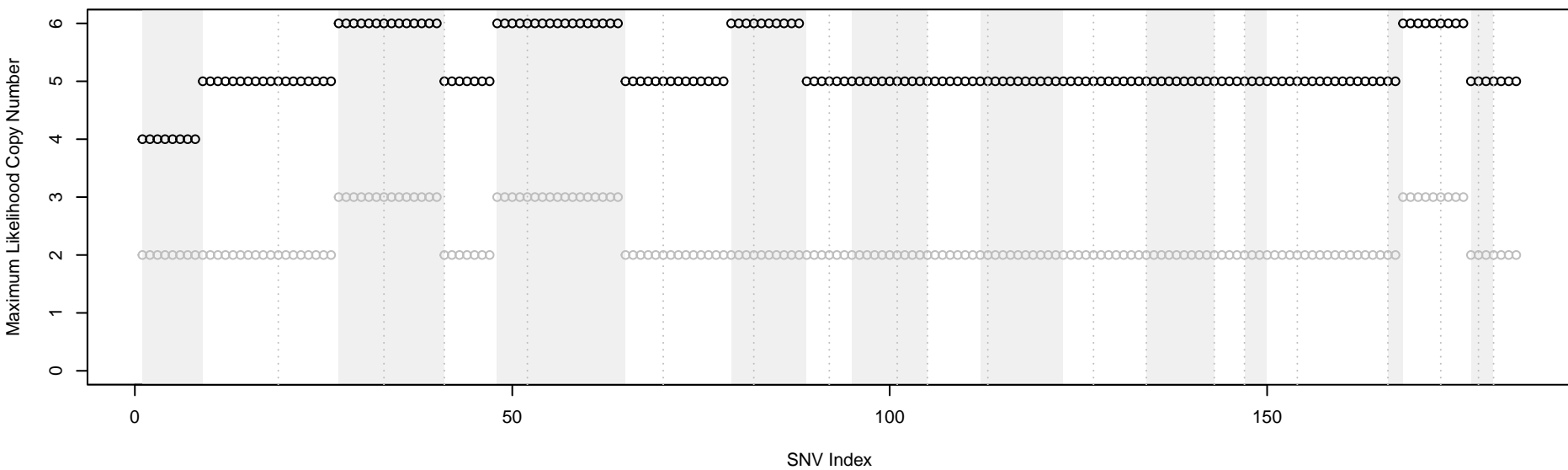
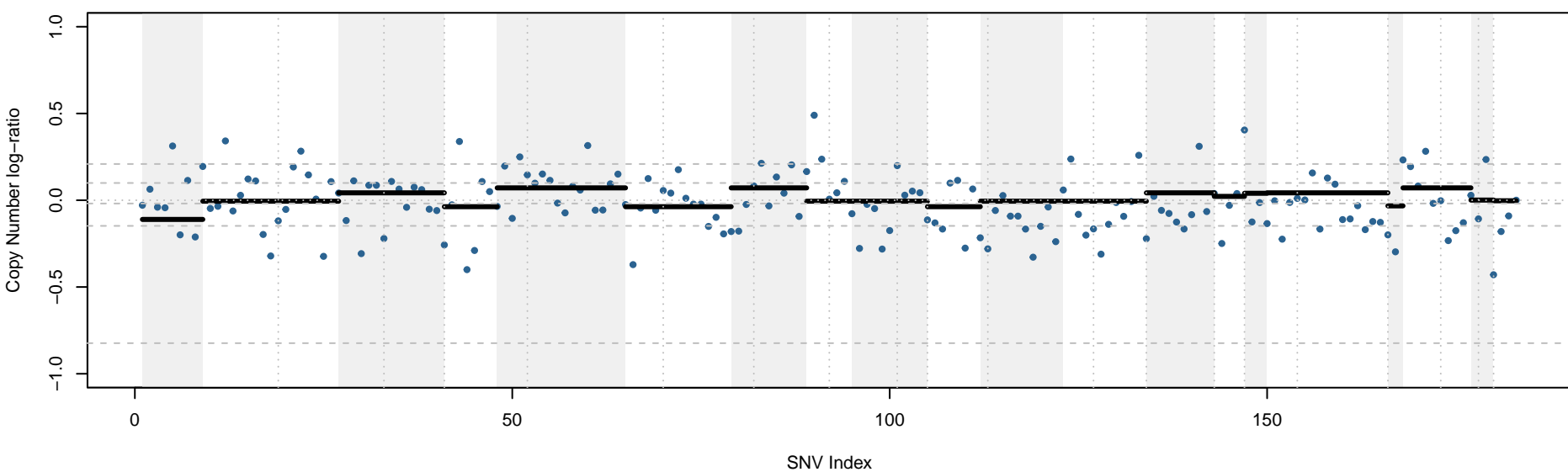
Purity: 0.23 Tumor ploidy: 5.154

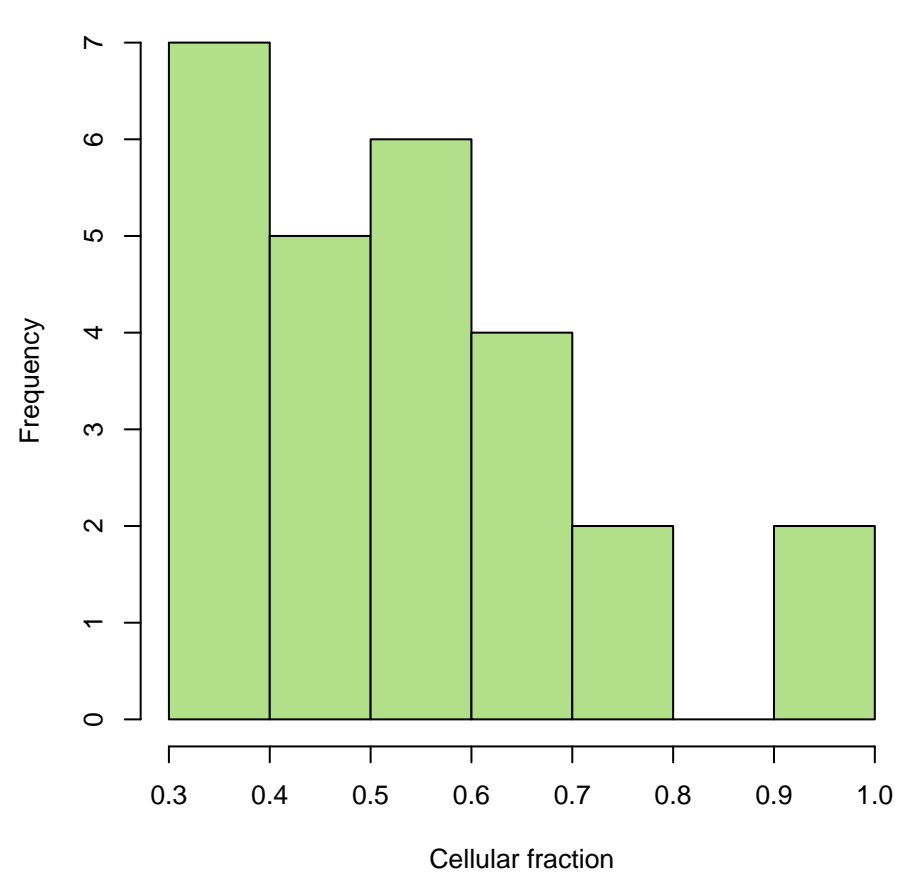
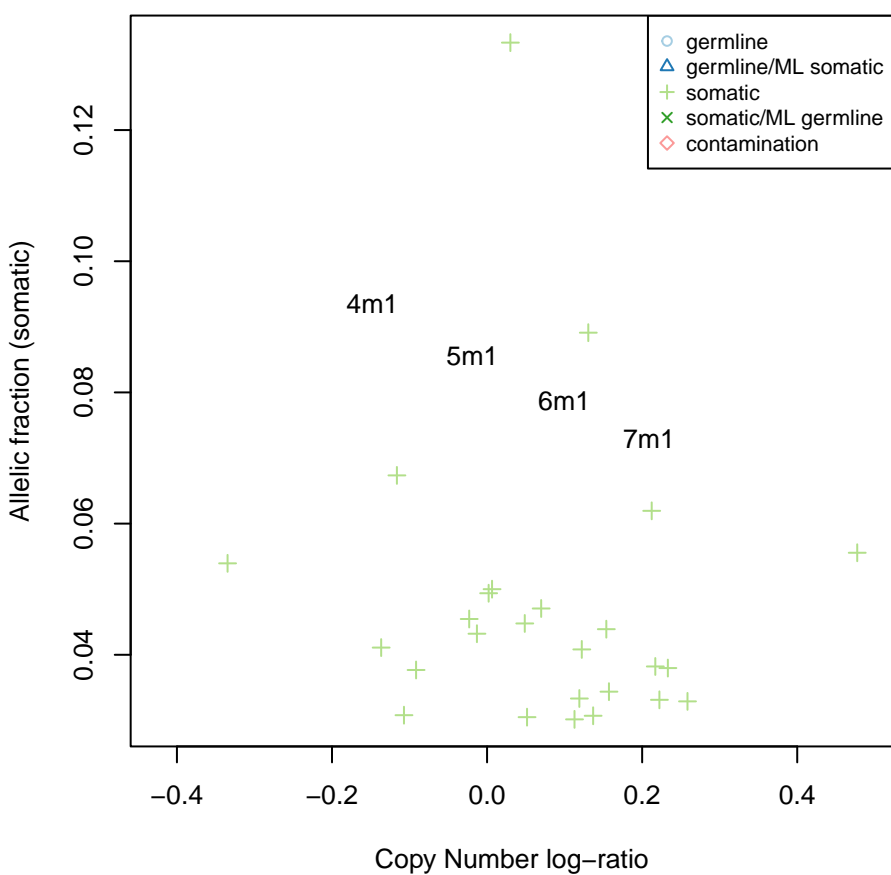
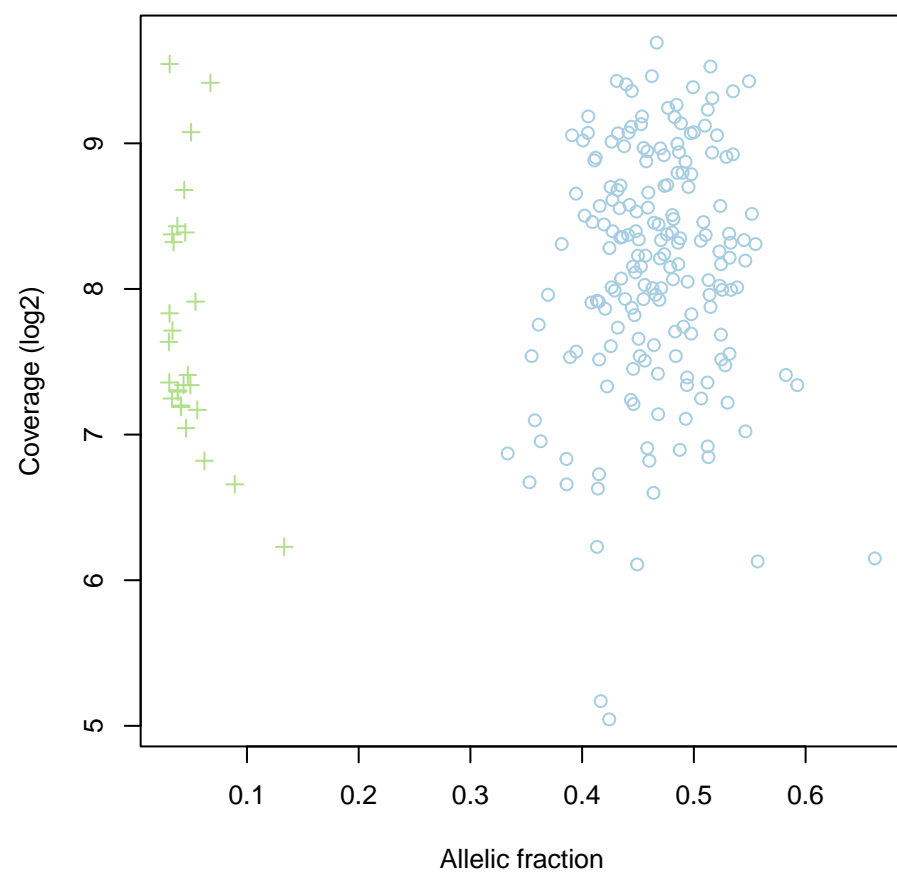
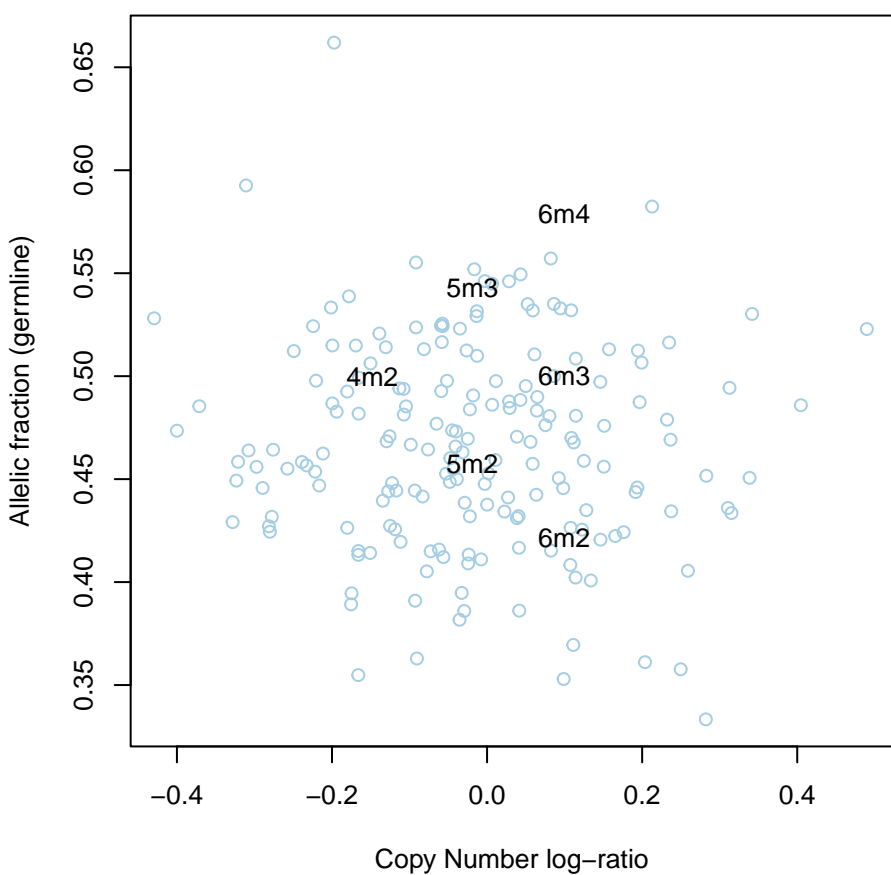


Purity: 0.23 Tumor ploidy: 5.154 SNV log-likelihood: -98.16 GoF: 96.2% Mean coverage: 317;651

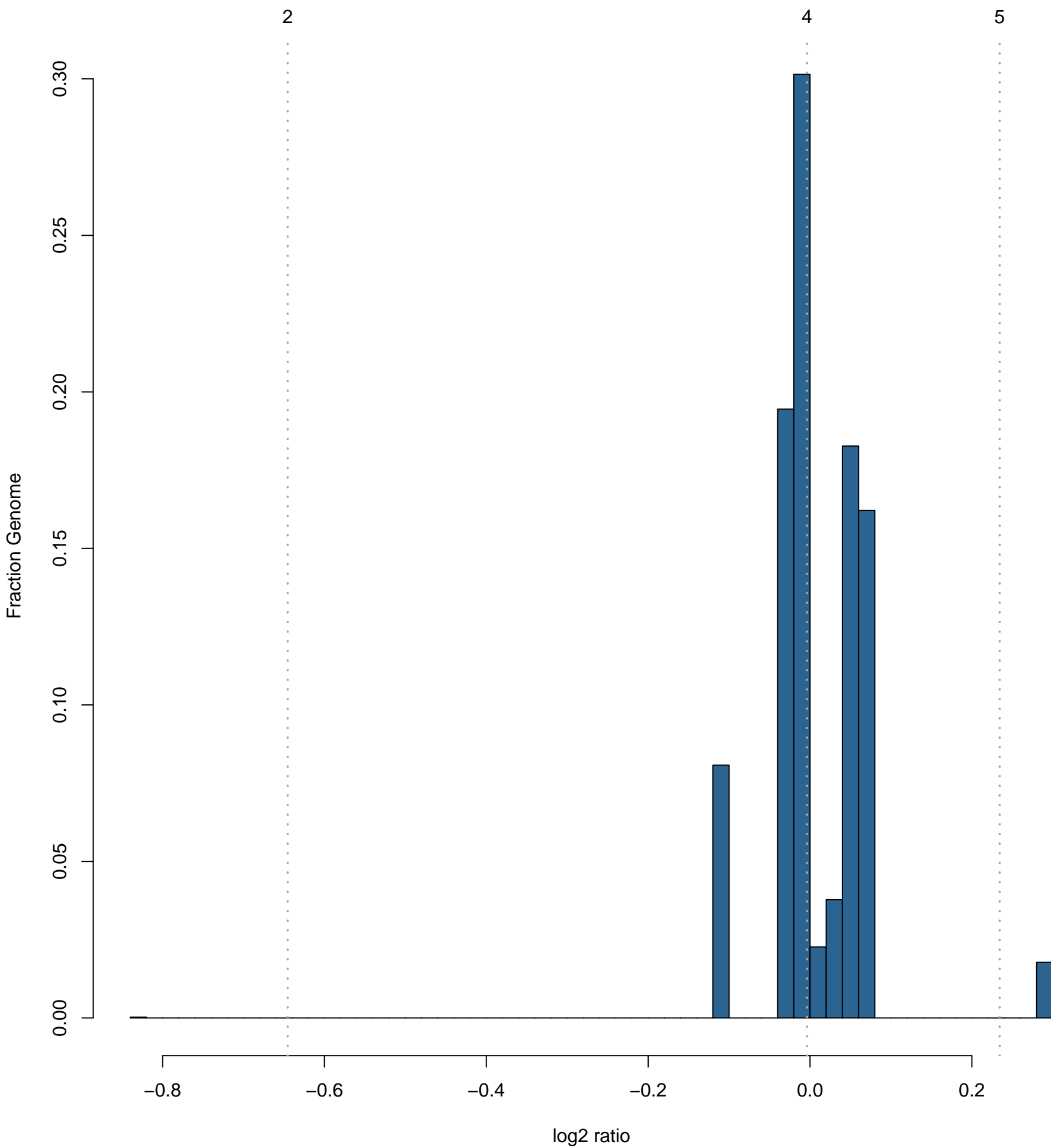


SCNA-fit log-likelihood: -9491.94

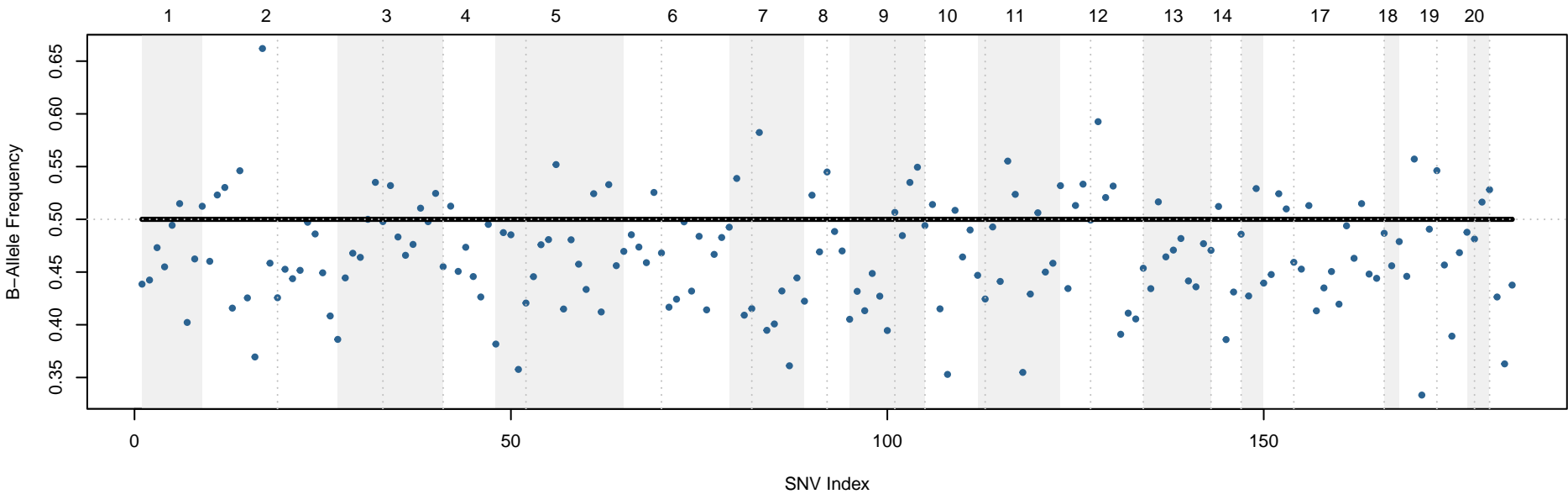




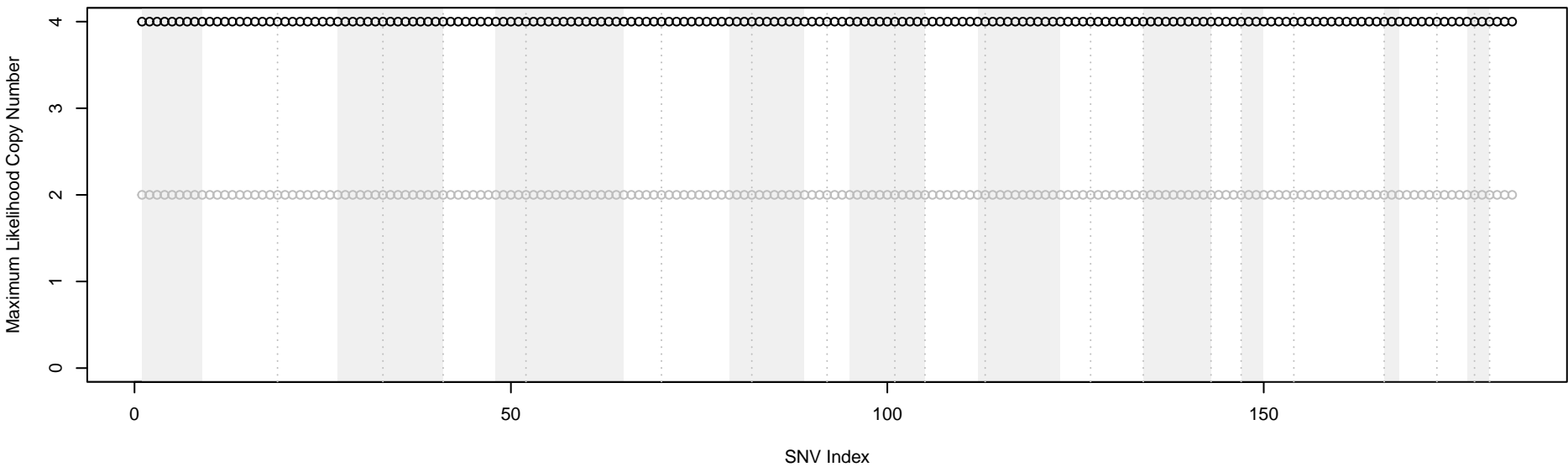
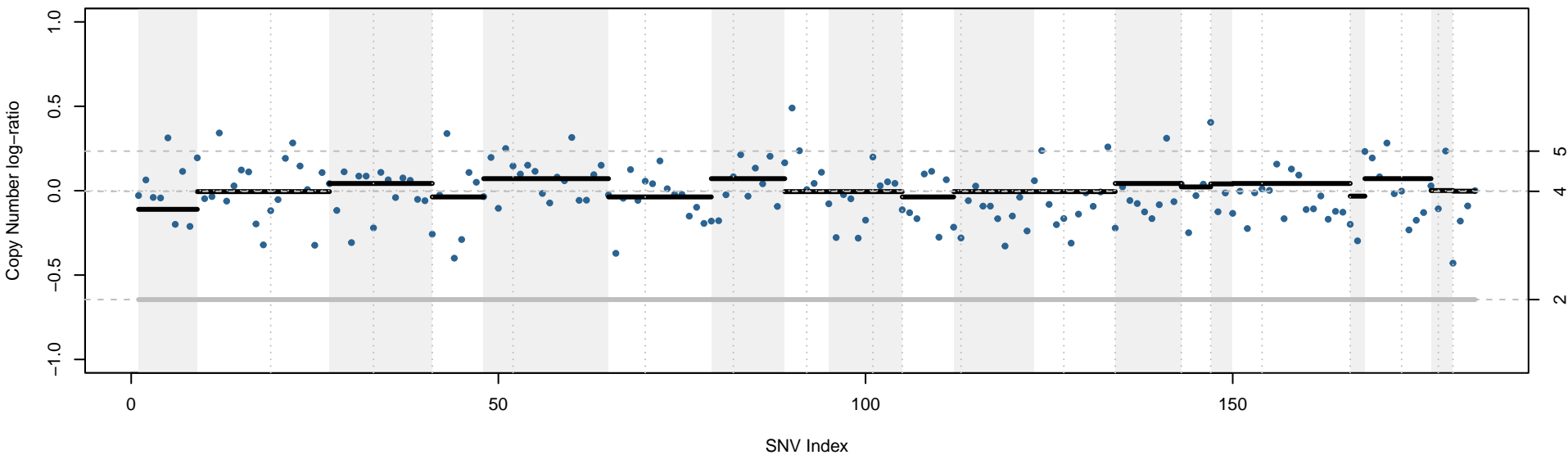
Purity: 0.56 Tumor ploidy: 4.015

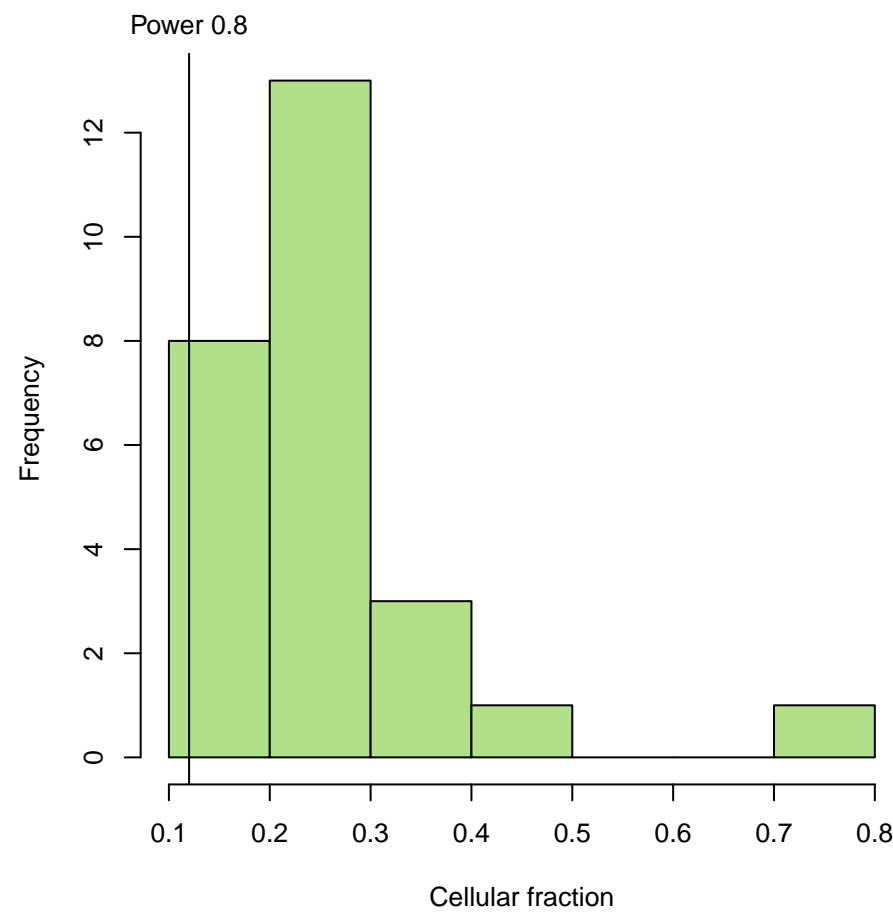
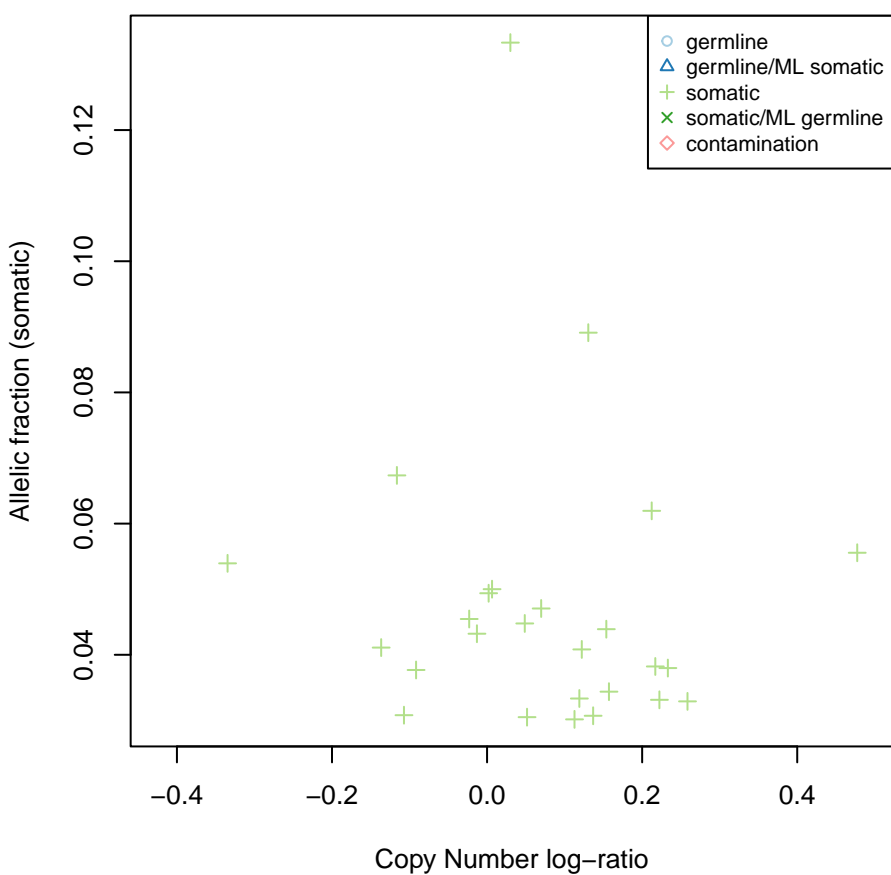
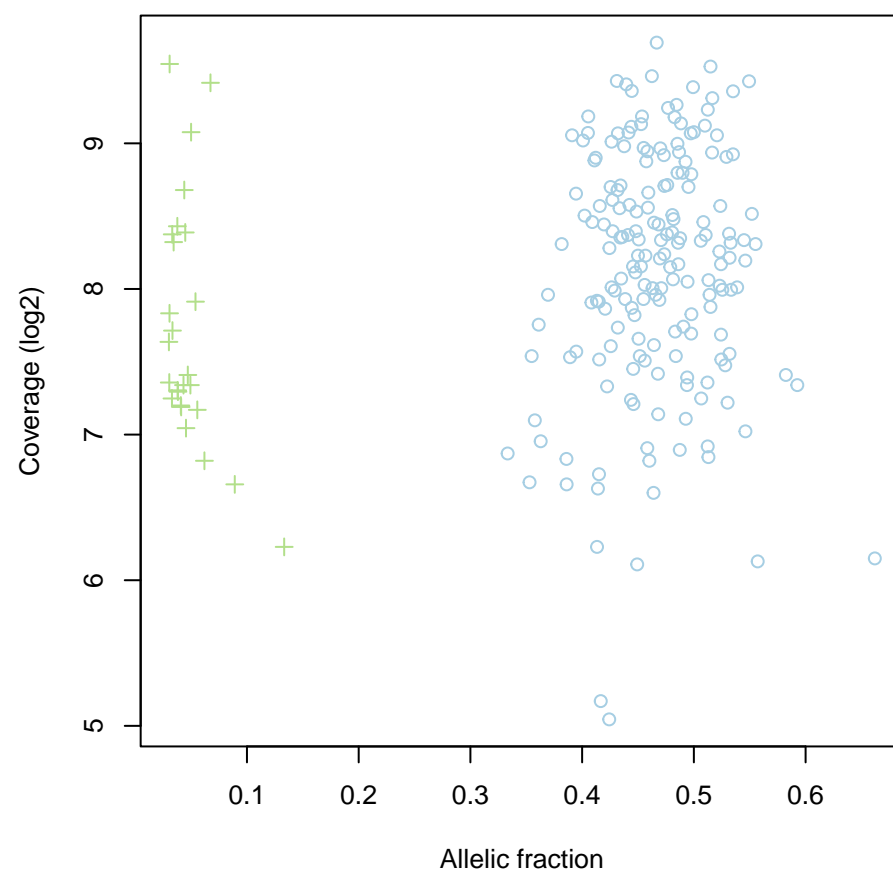
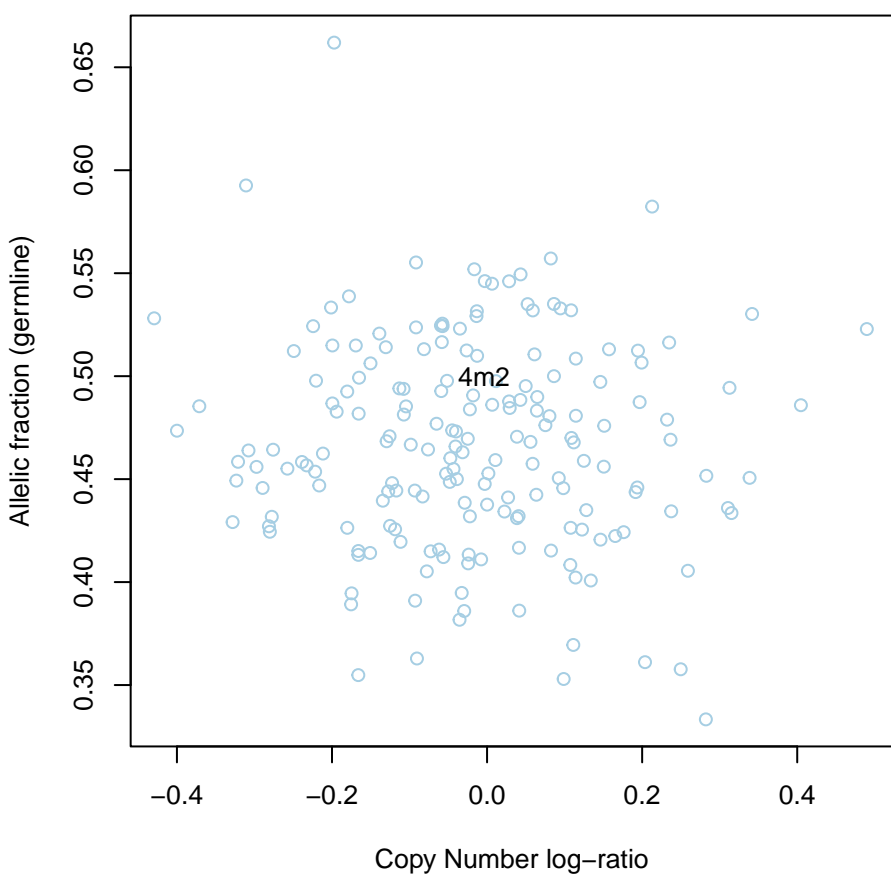


Purity: 0.56 Tumor ploidy: 4.015 SNV log-likelihood: -80.86 GoF: 88% Mean coverage: 317;651



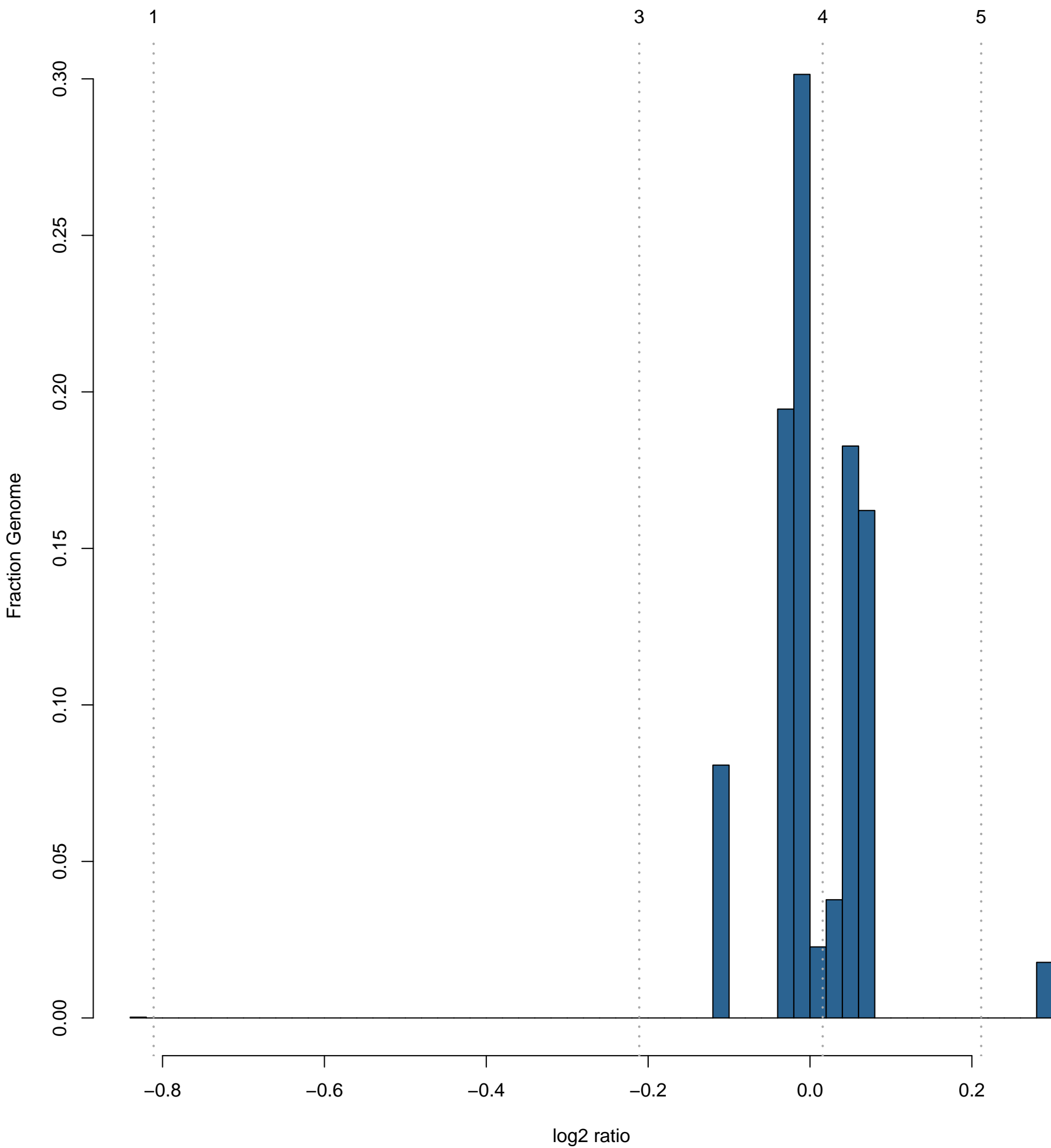
SCNA-fit log-likelihood: -9604.64



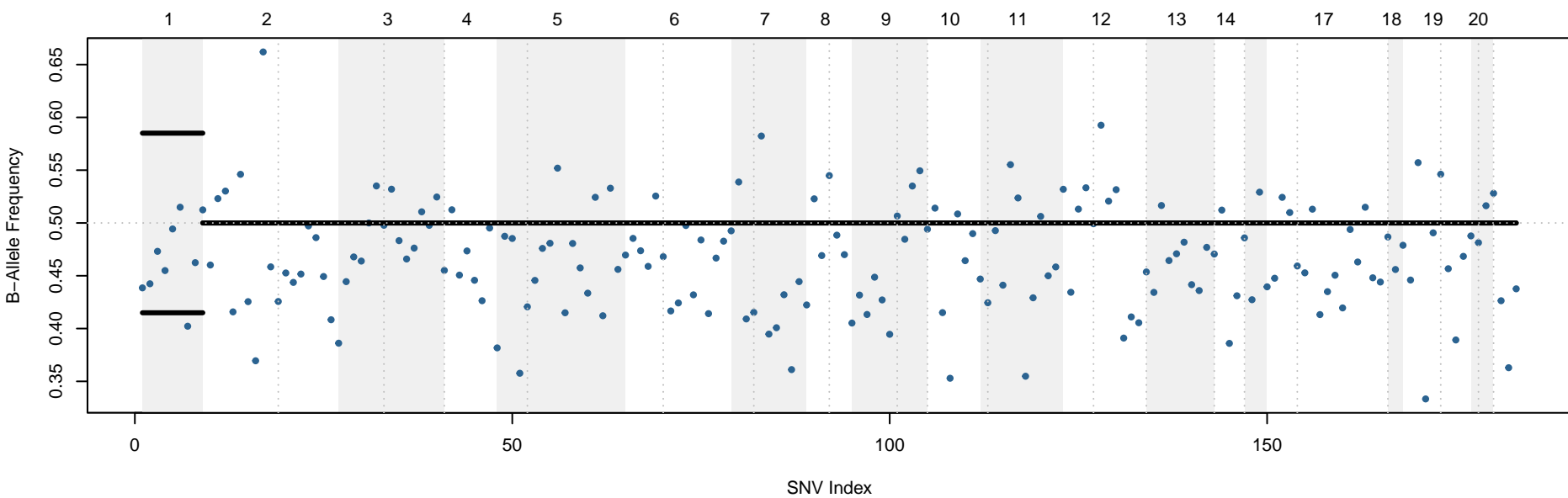




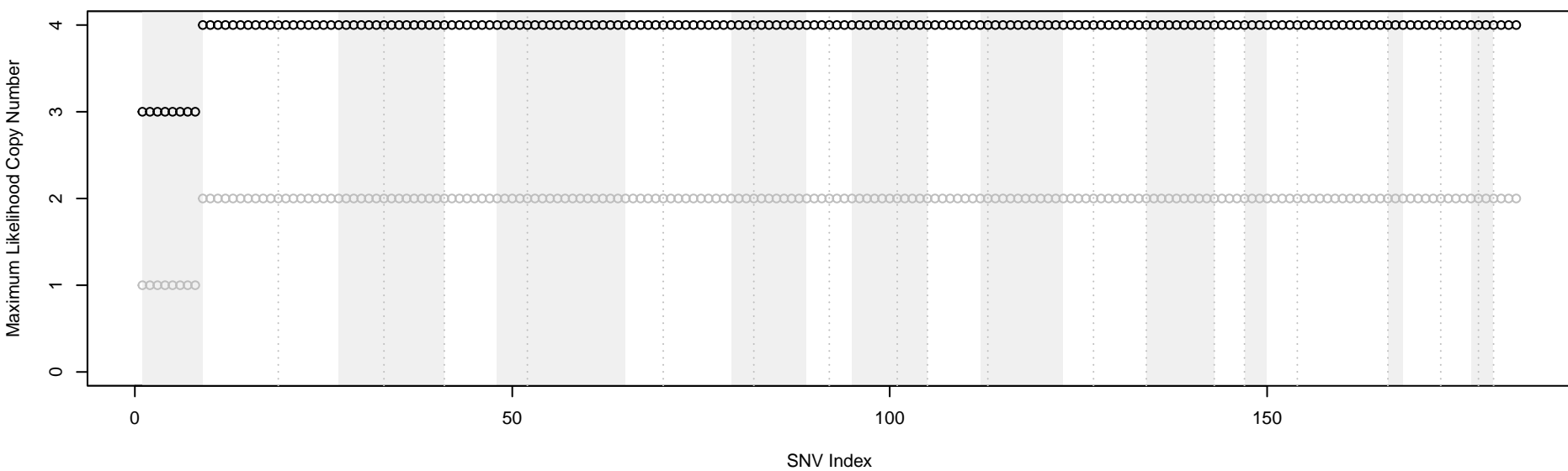
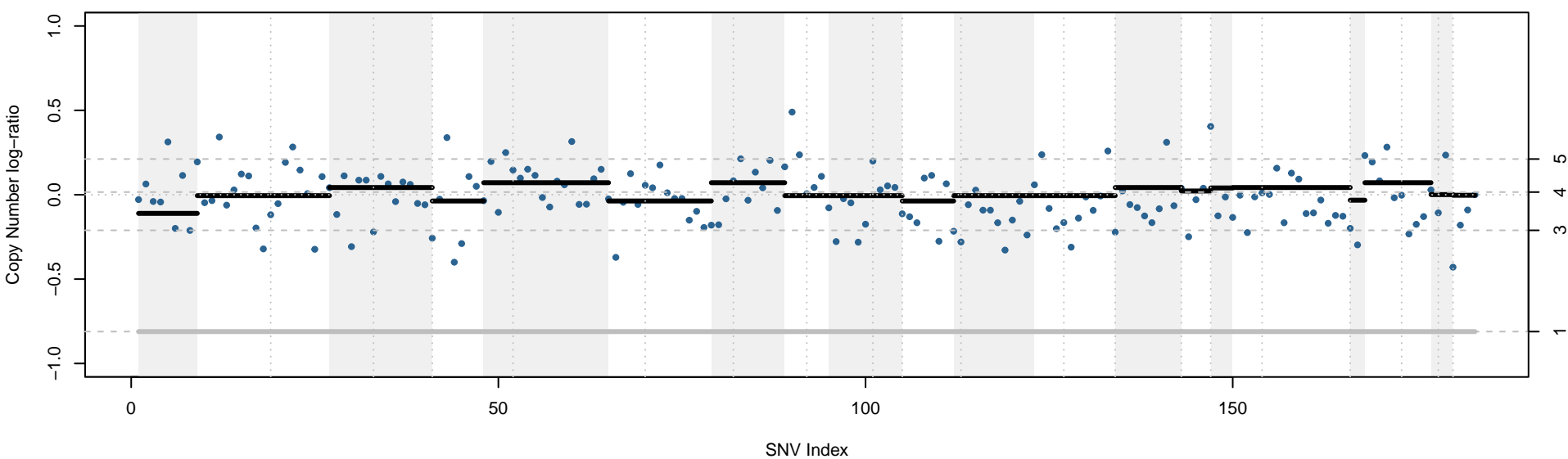
Purity: 0.41 Tumor ploidy: 3.926

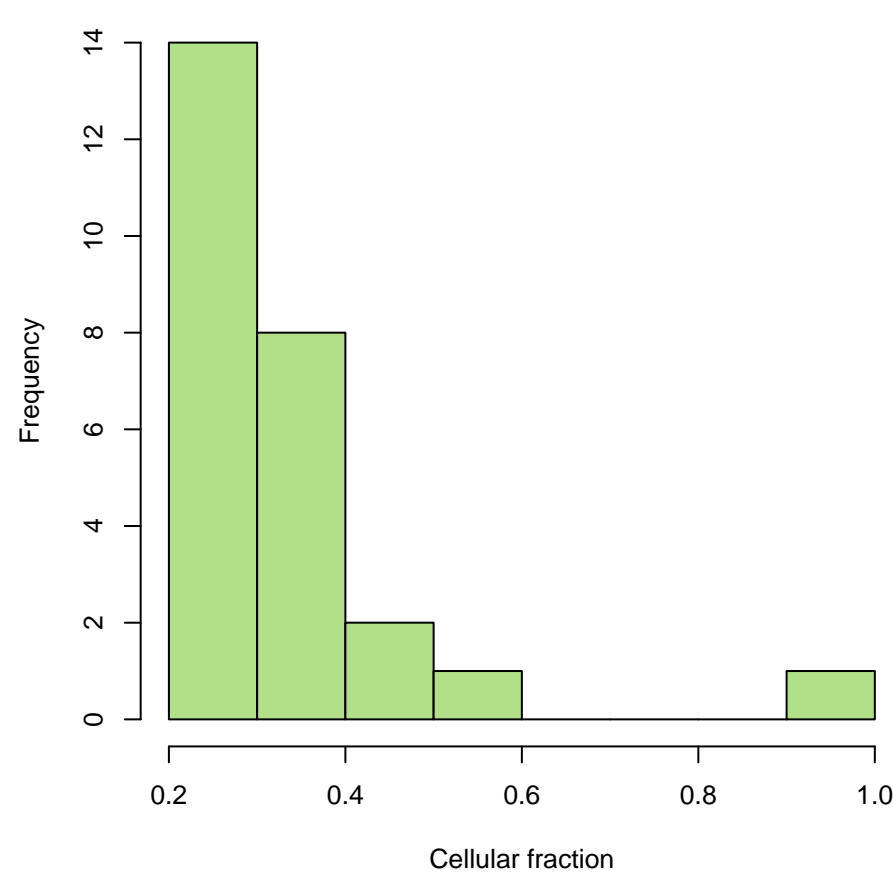
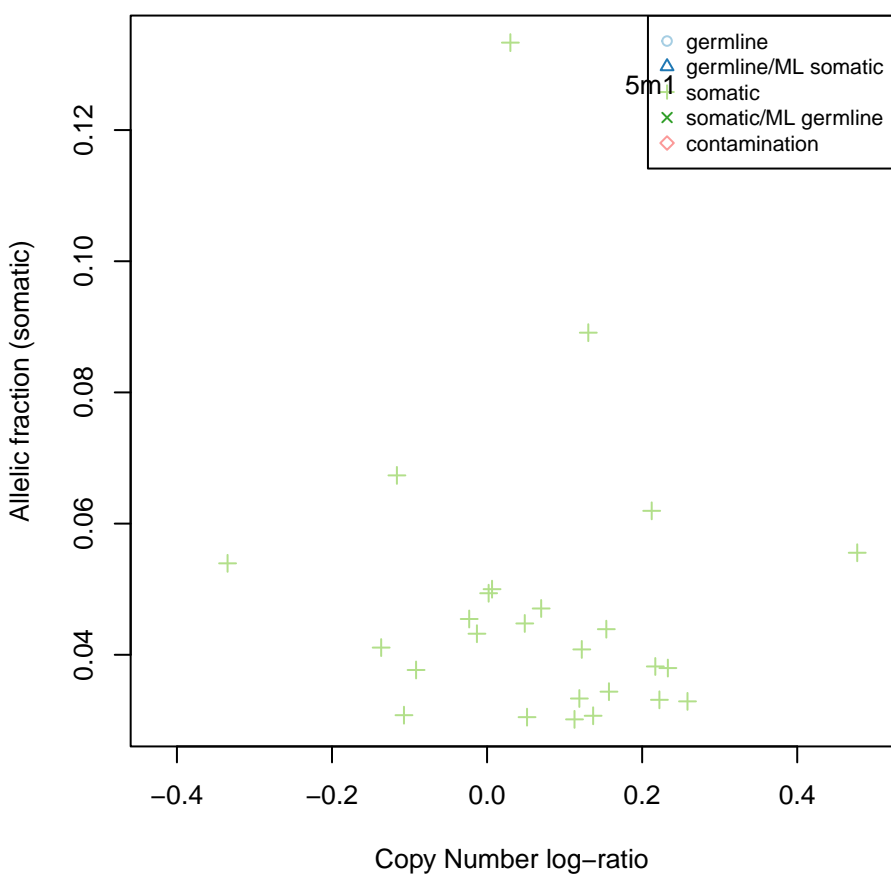
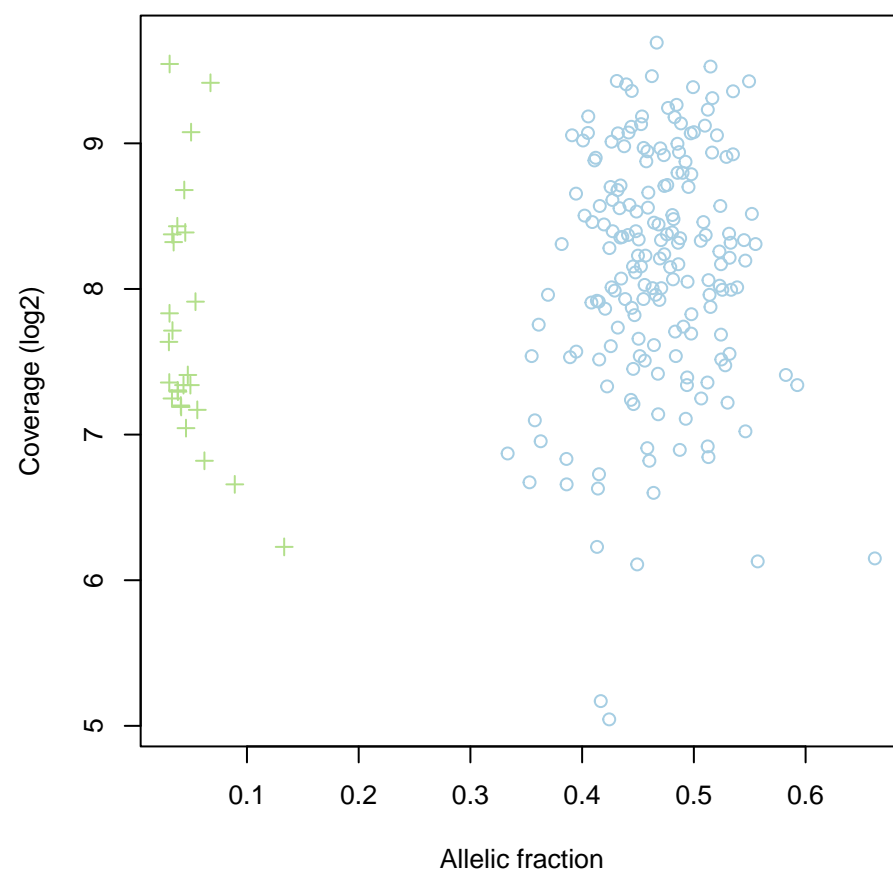
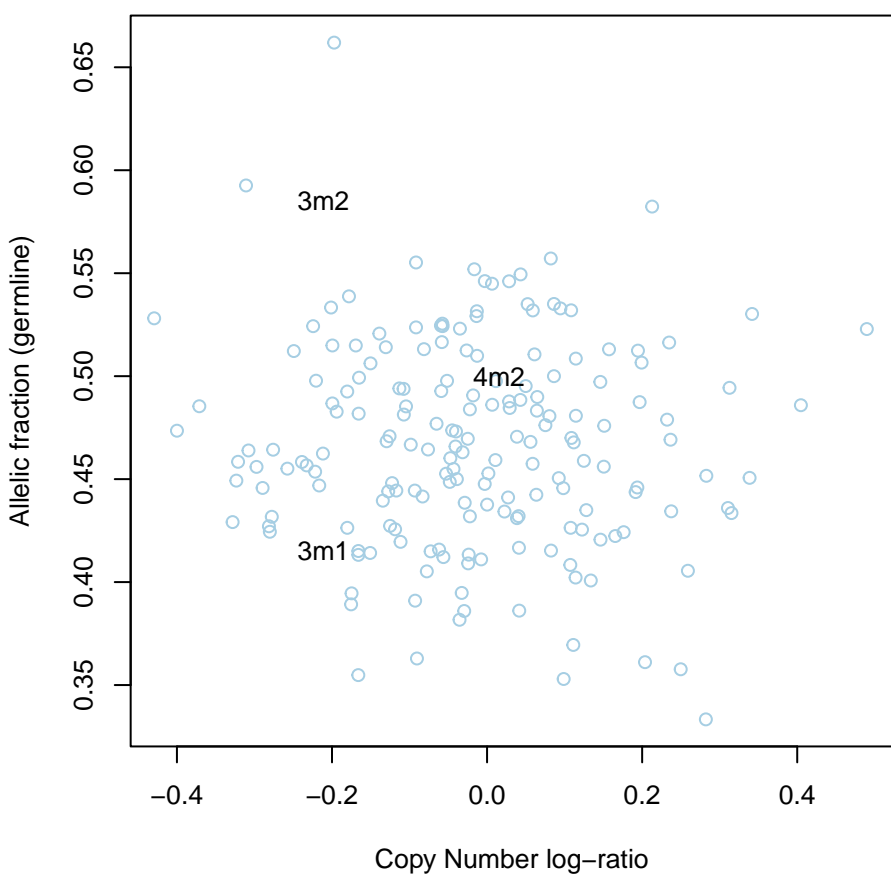


Purity: 0.41 Tumor ploidy: 3.926 SNV log-likelihood: -80.83 GoF: 90.1% Mean coverage: 317;651

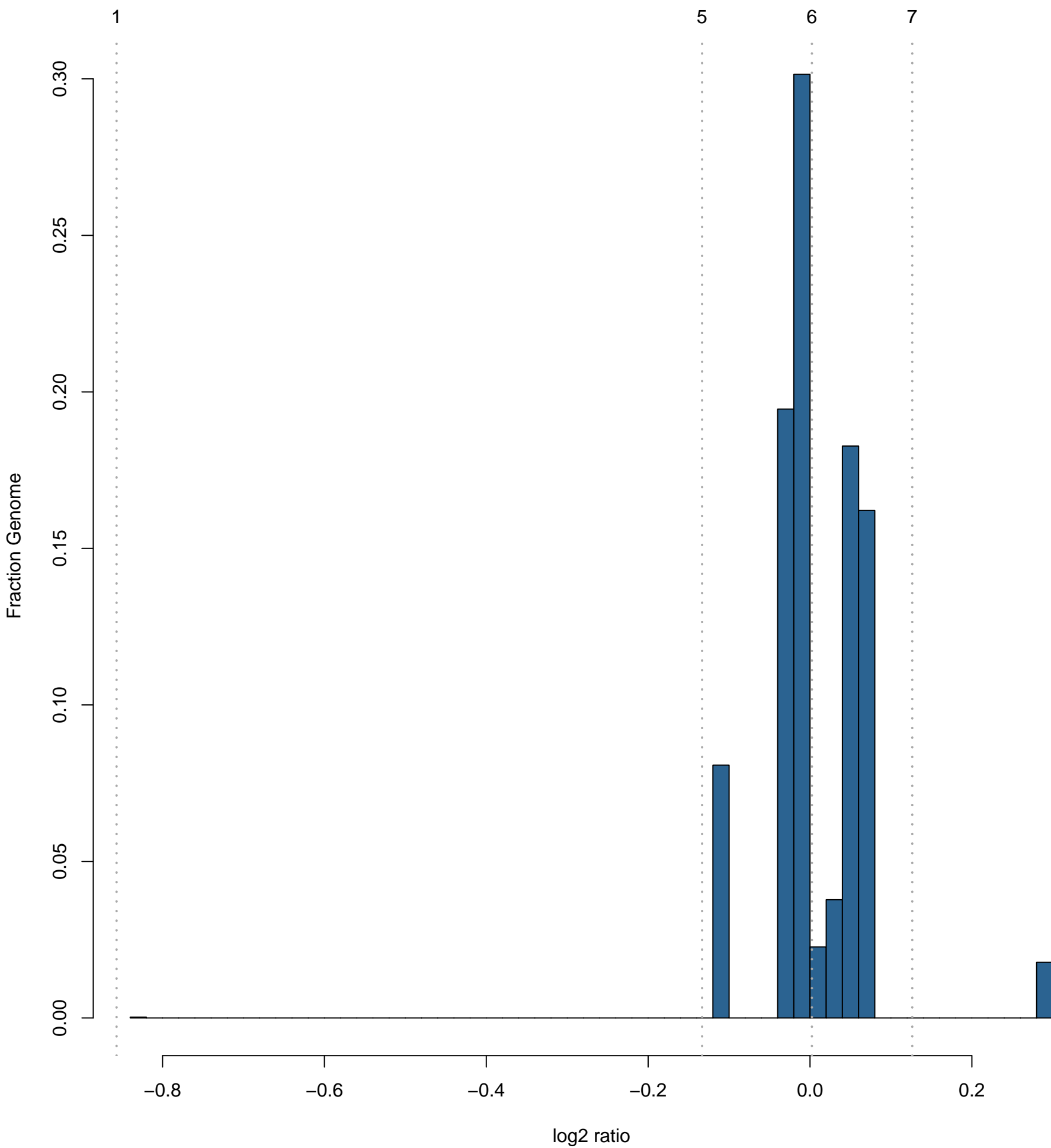


SCNA-fit log-likelihood: -9584.84

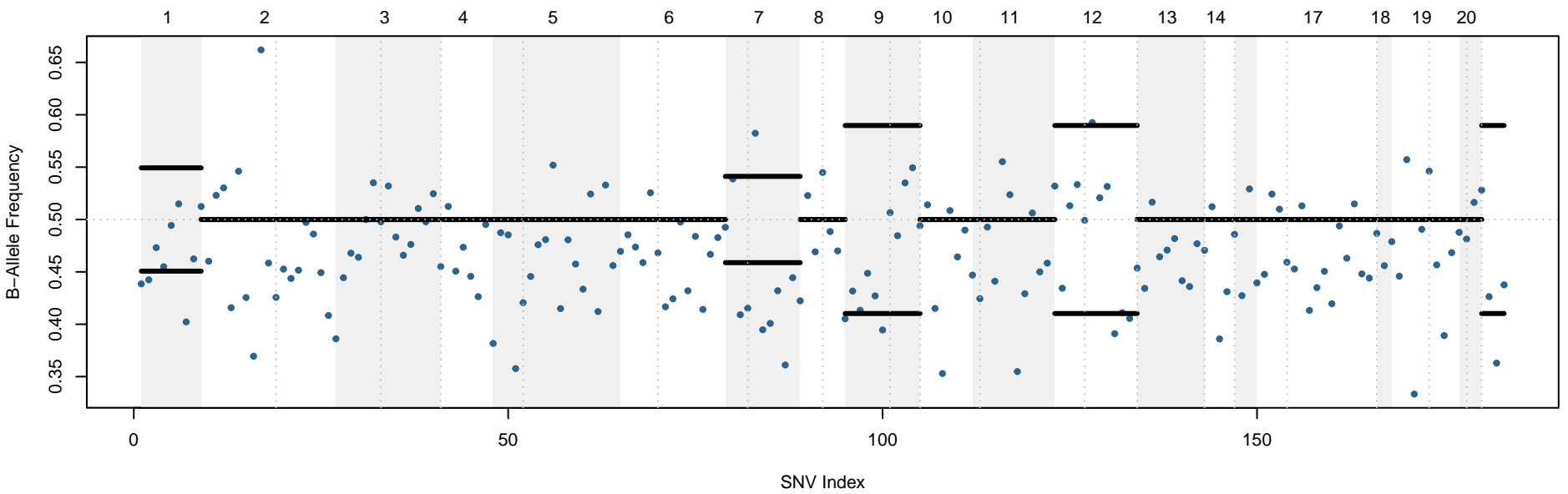




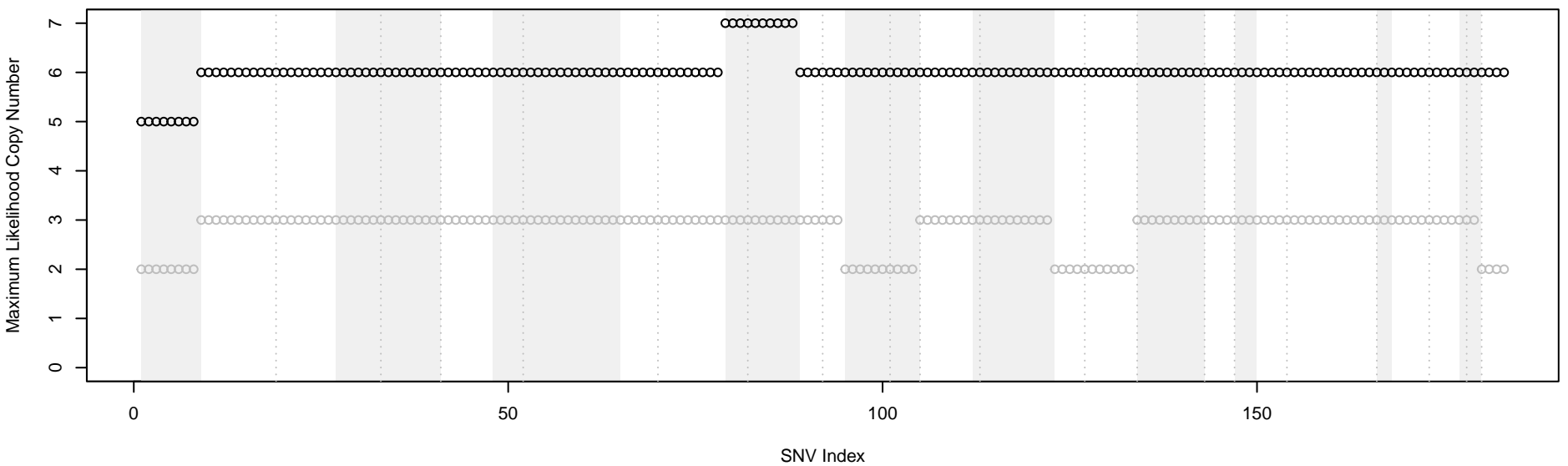
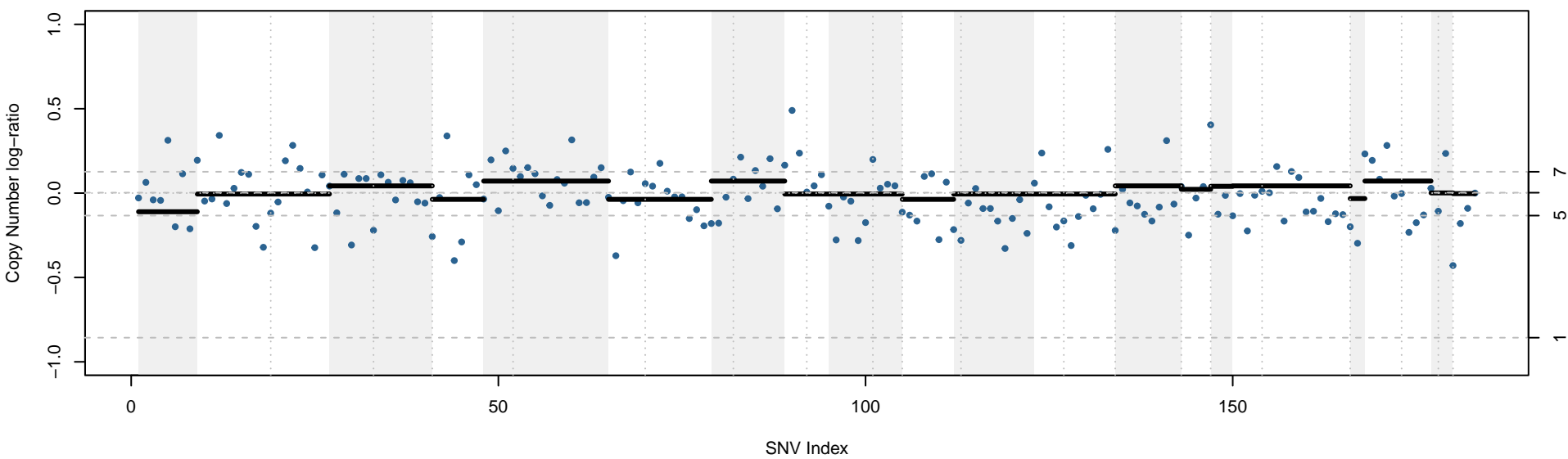
Purity: 0.28 Tumor ploidy: 5.982

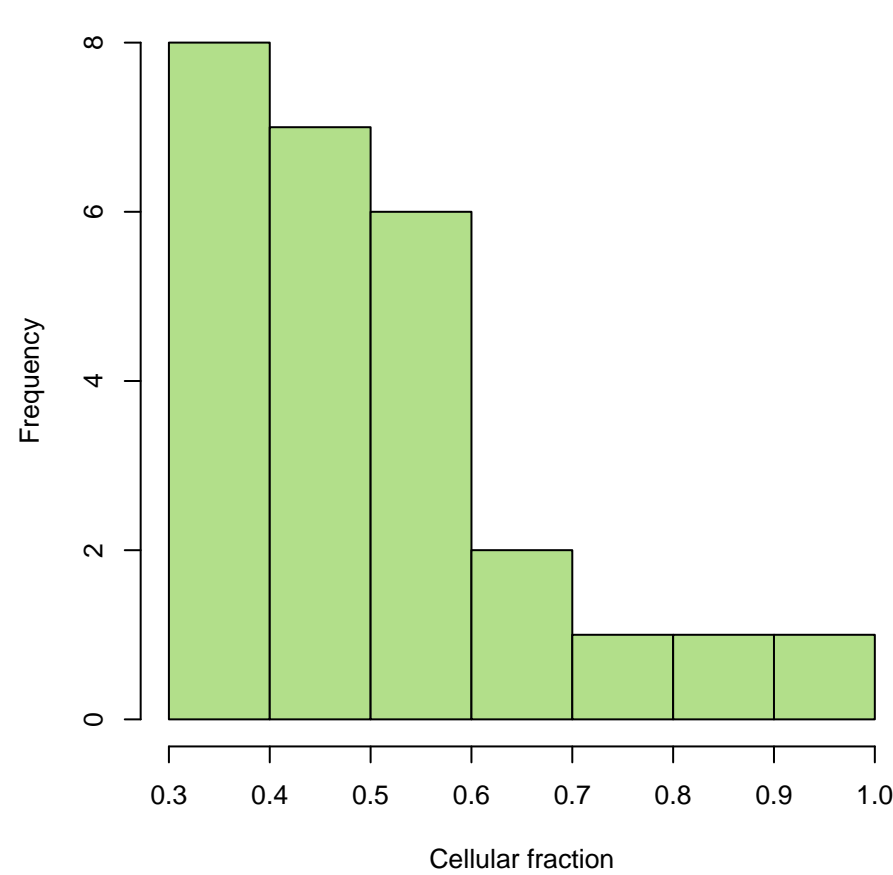
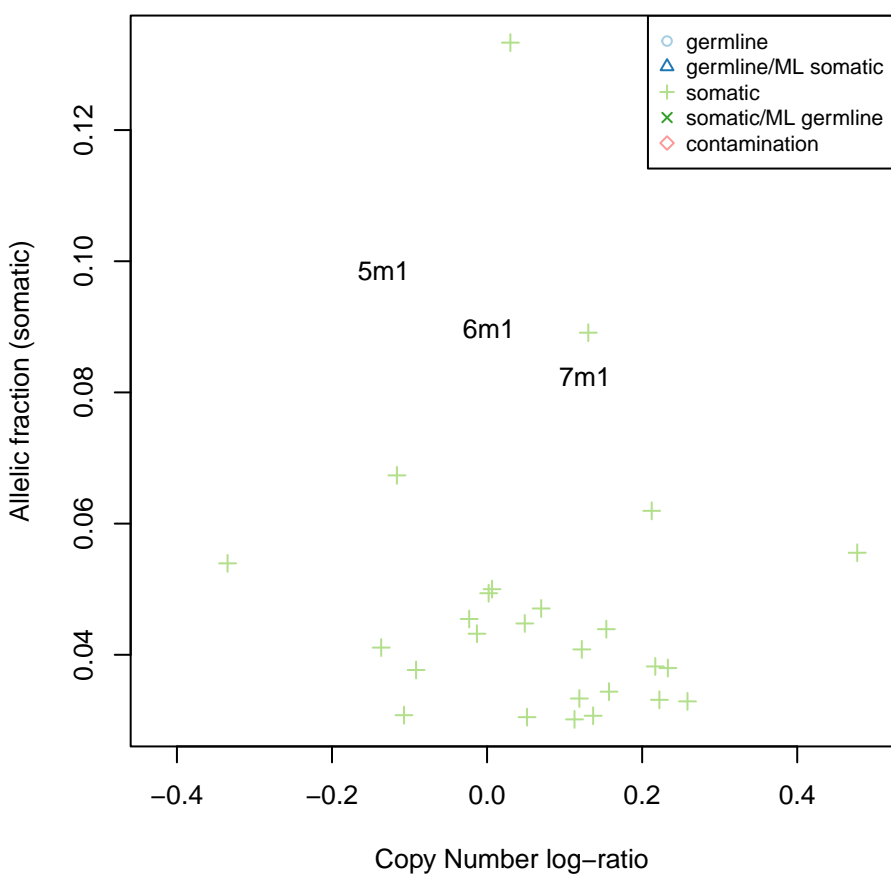
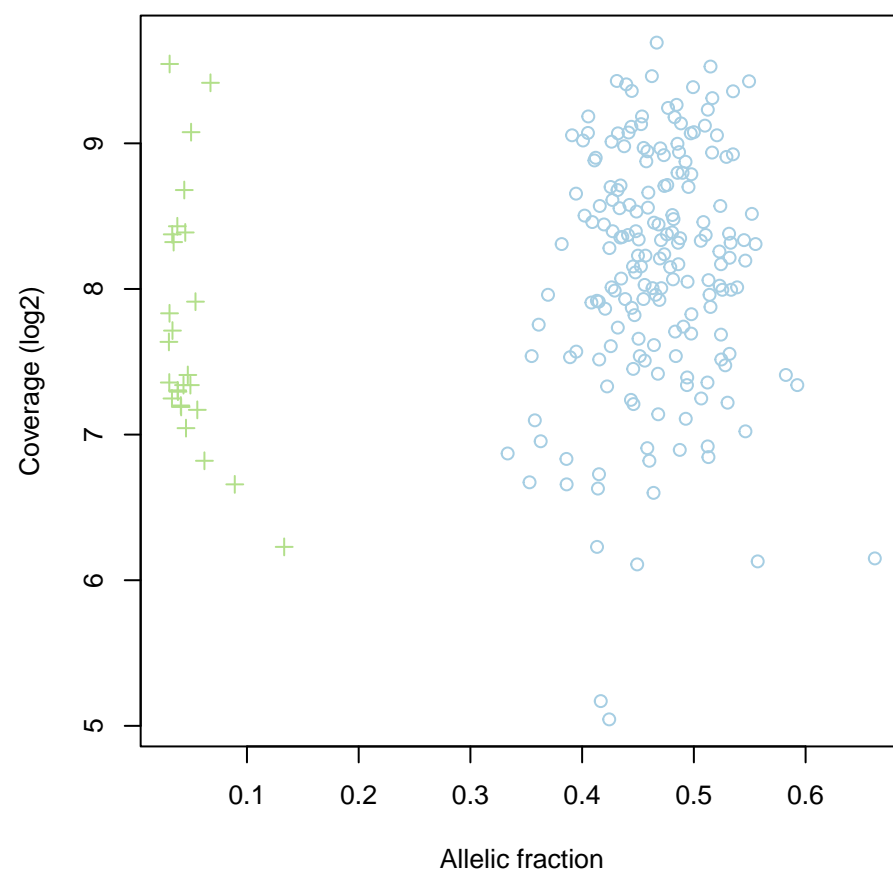
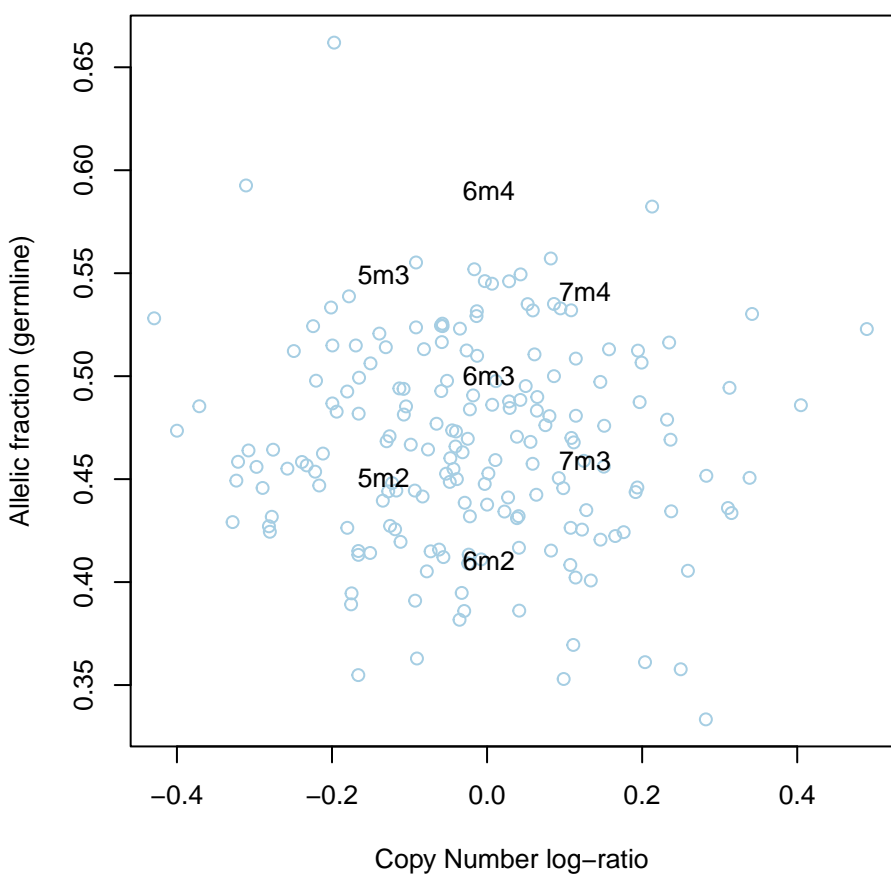


Purity: 0.28 Tumor ploidy: 5.982 SNV log-likelihood: -126.23 GoF: 94% Mean coverage: 317;651

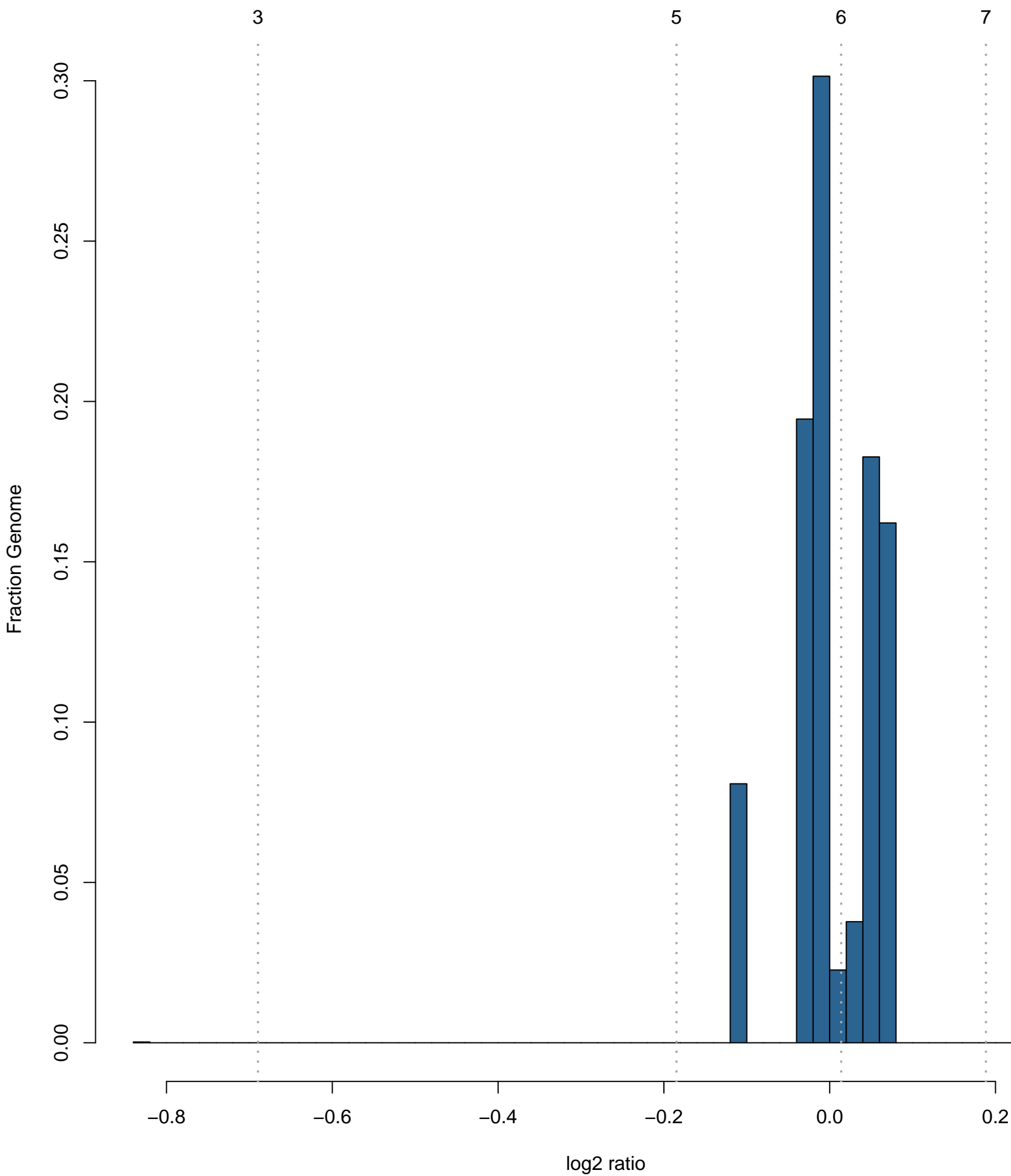


SCNA-fit log-likelihood: -9546.72

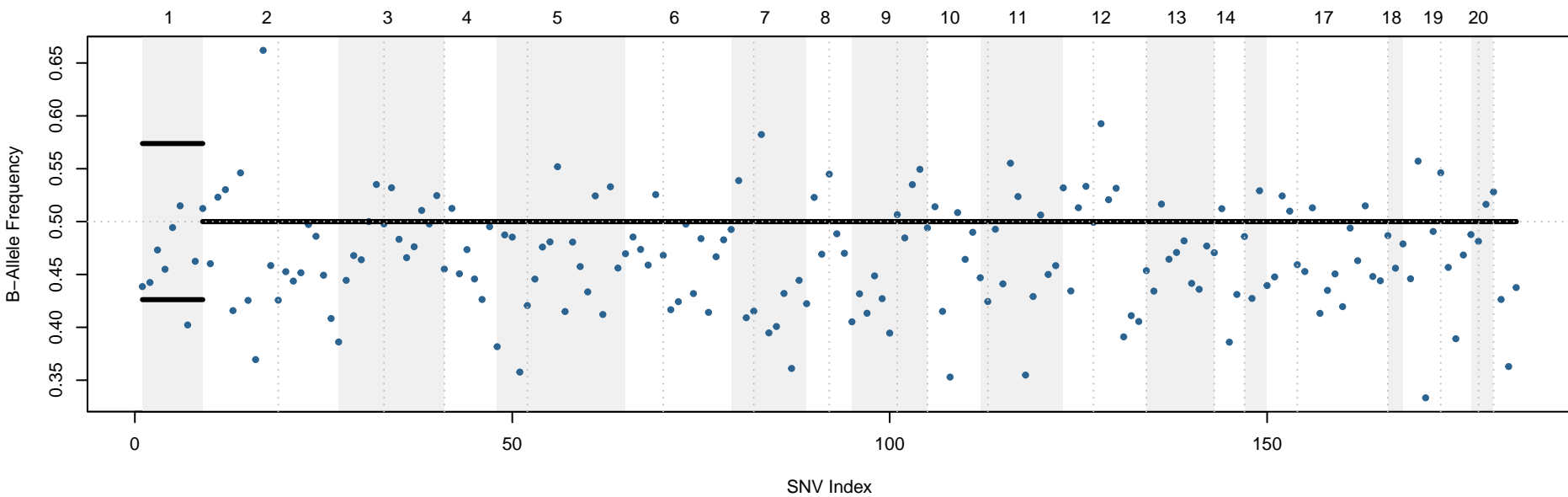




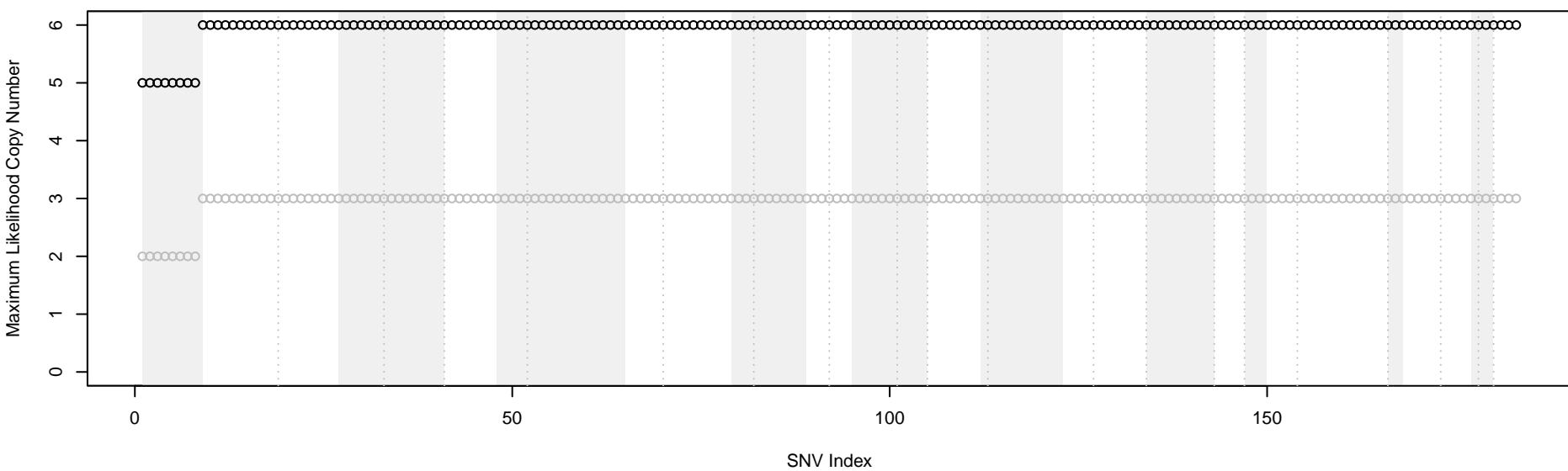
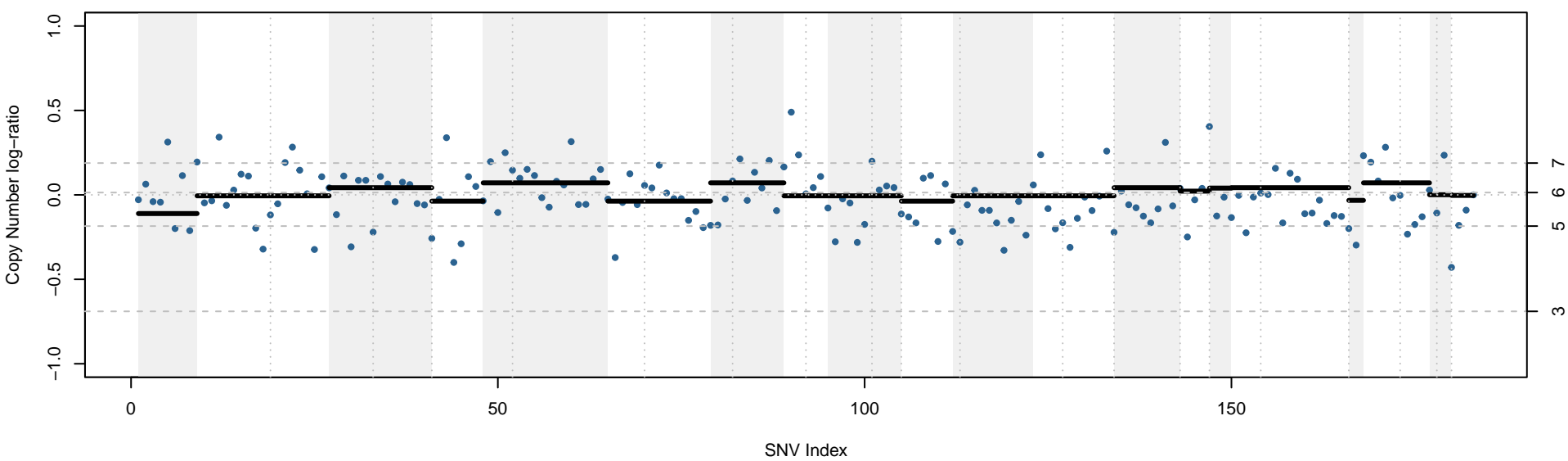
Purity: 0.53 Tumor ploidy: 5.926



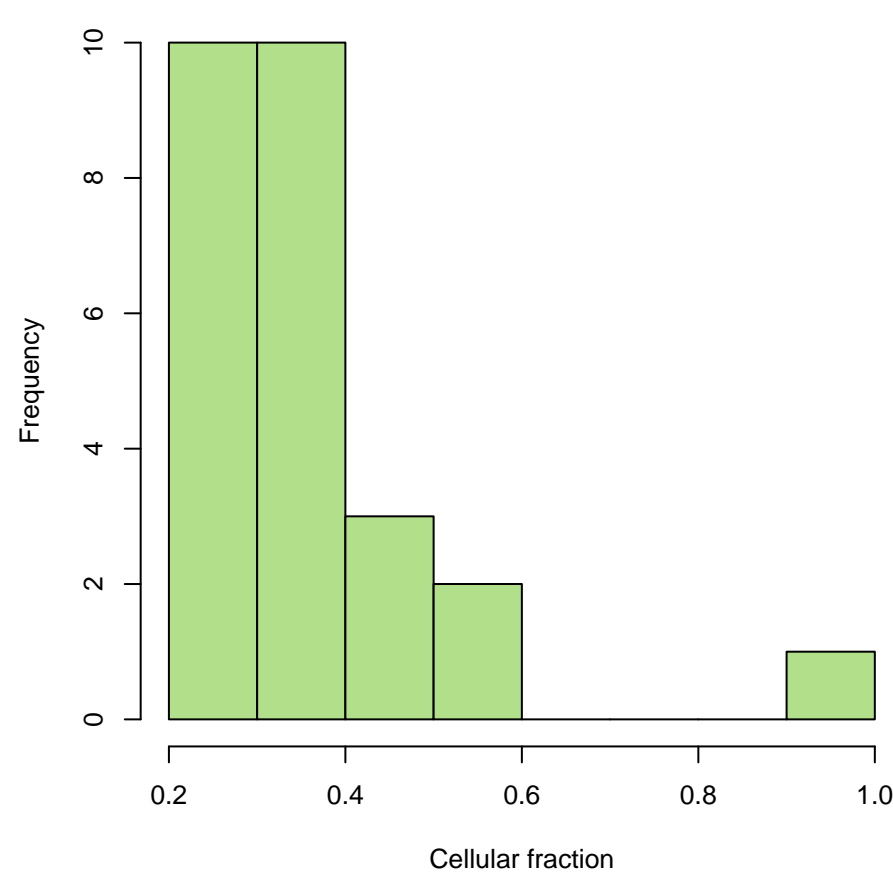
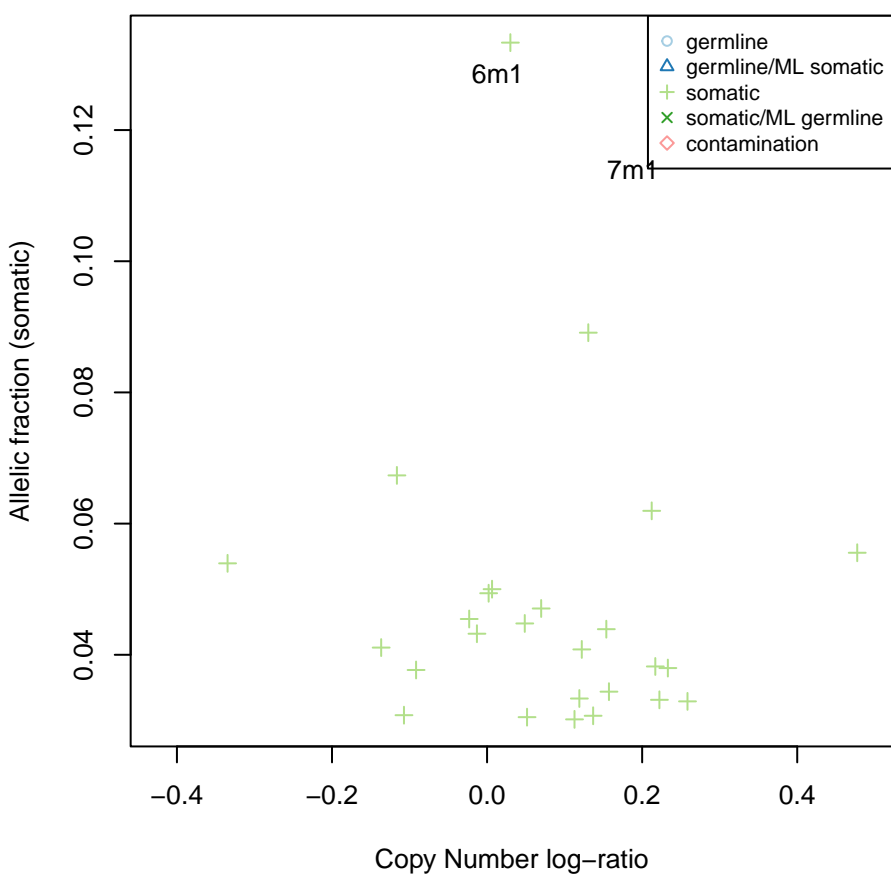
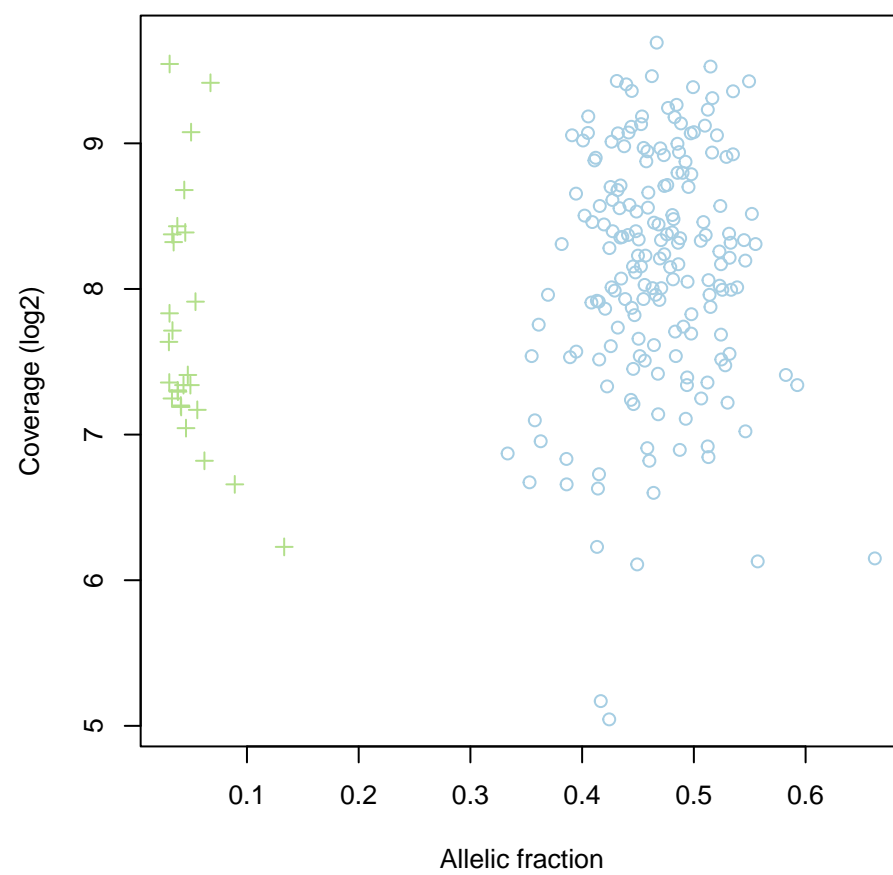
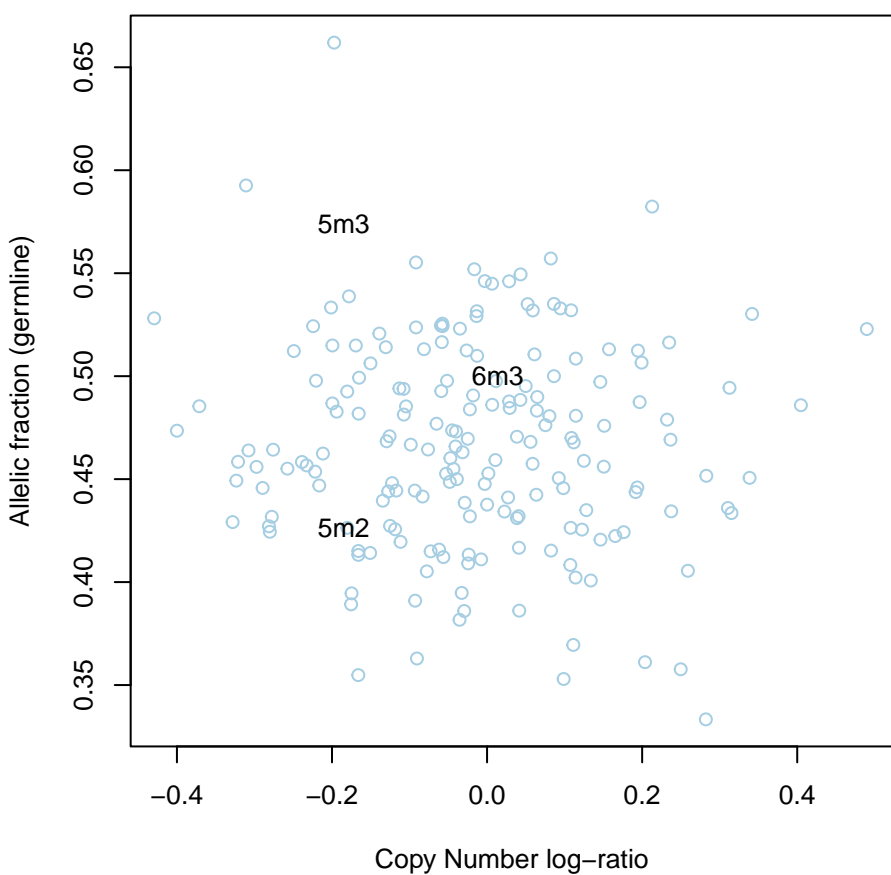
Purity: 0.53 Tumor ploidy: 5.926 SNV log-likelihood: -146.1 GoF: 91.2% Mean coverage: 317;651



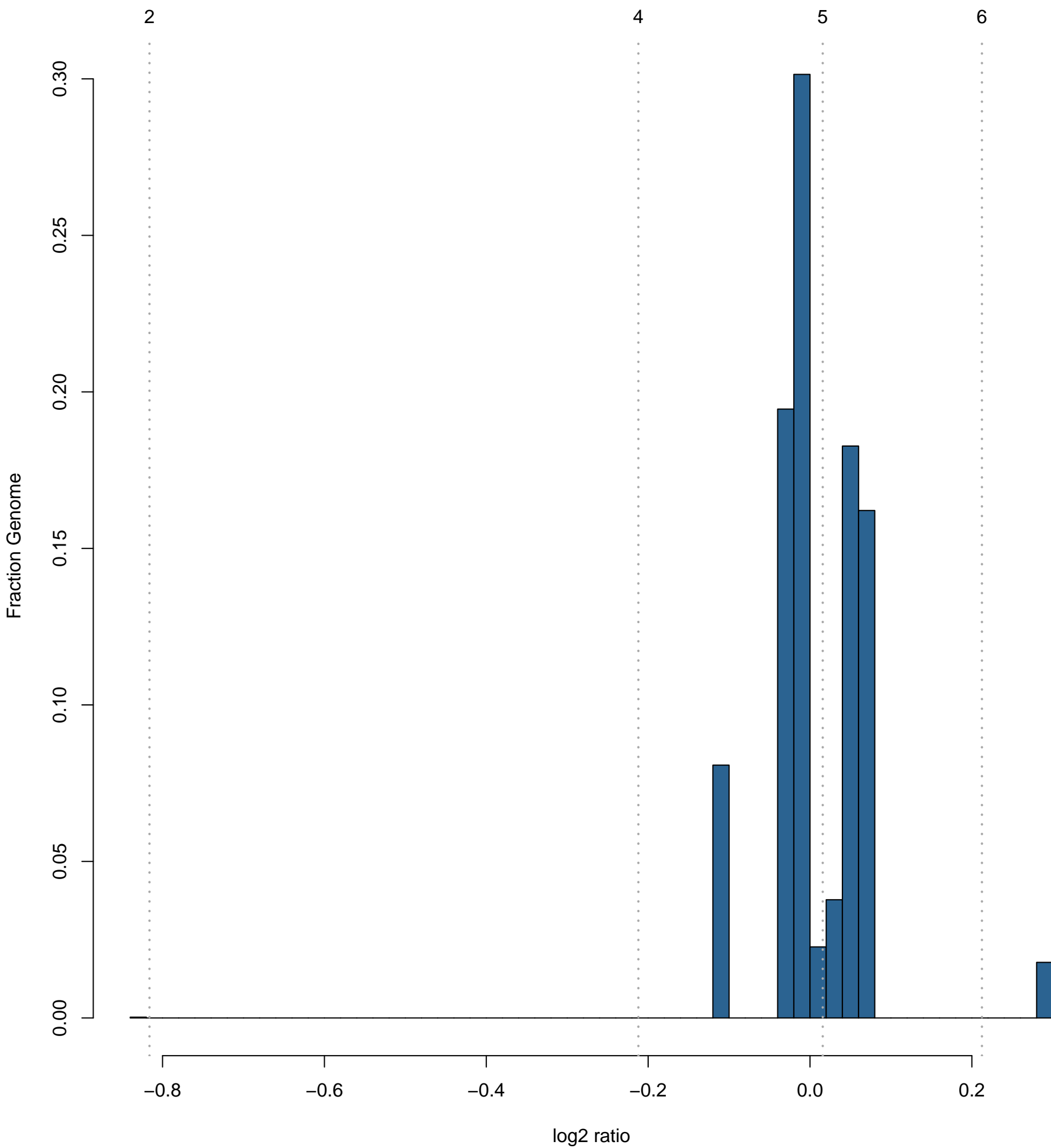
SCNA-fit log-likelihood: -9563.1



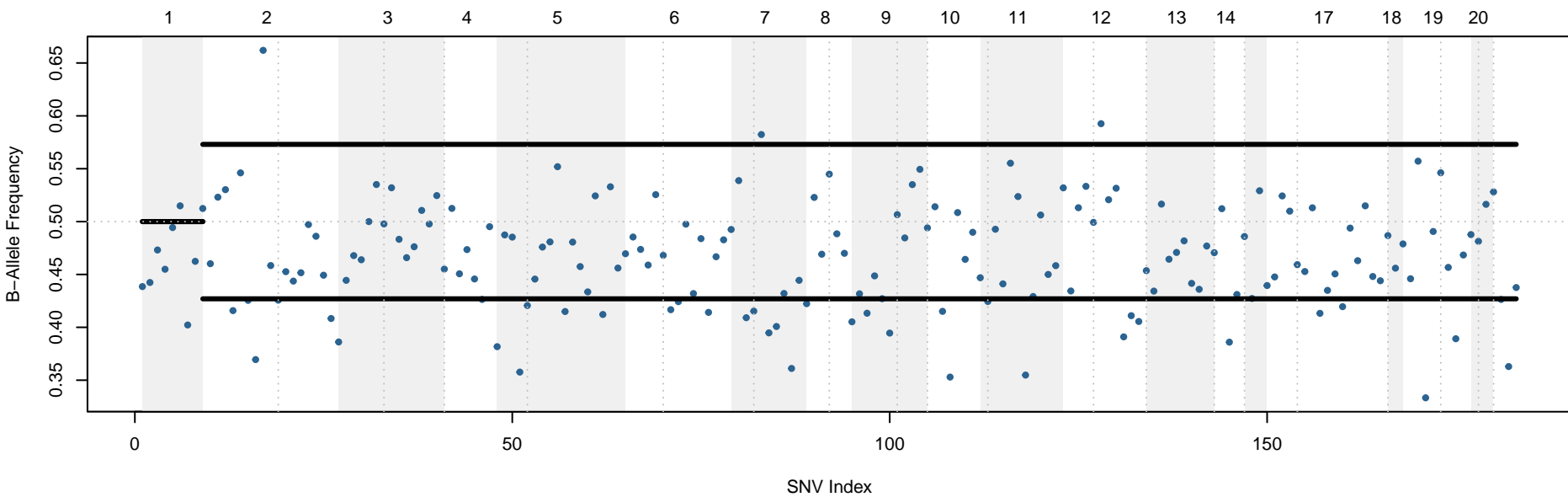




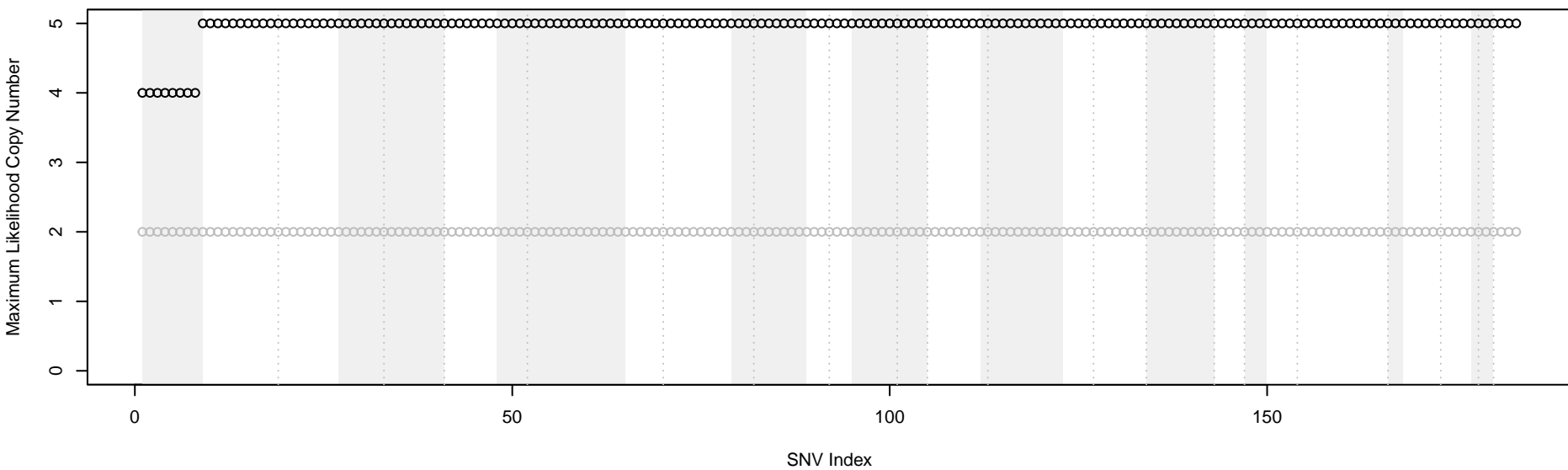
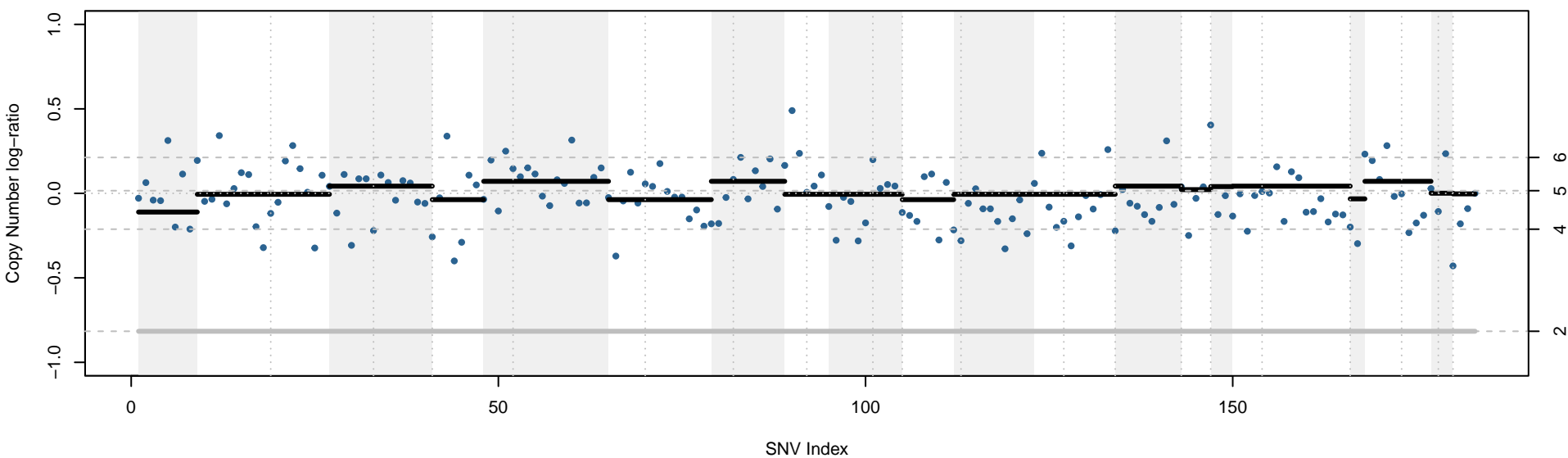
Purity: 0.52 Tumor ploidy: 4.926

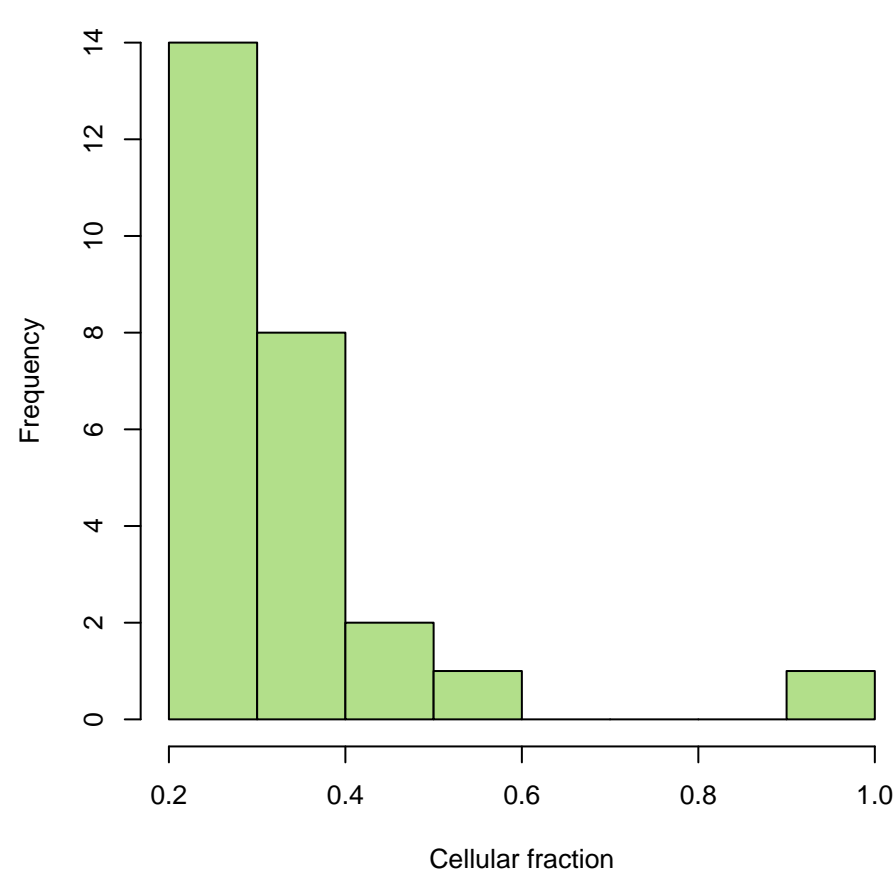
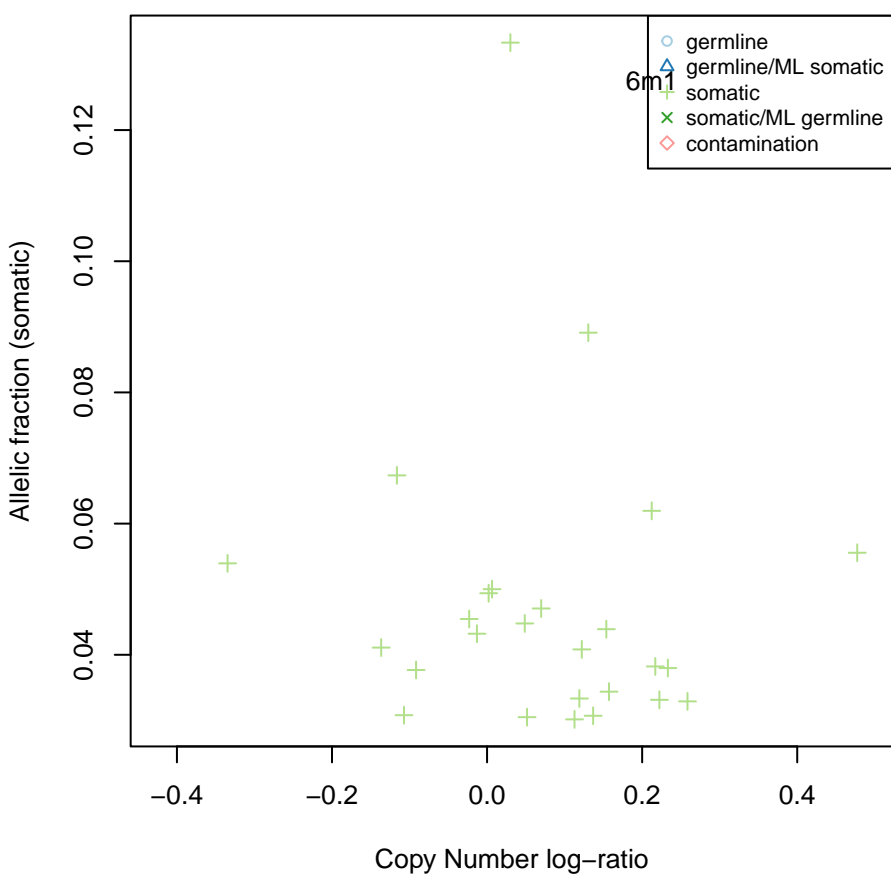
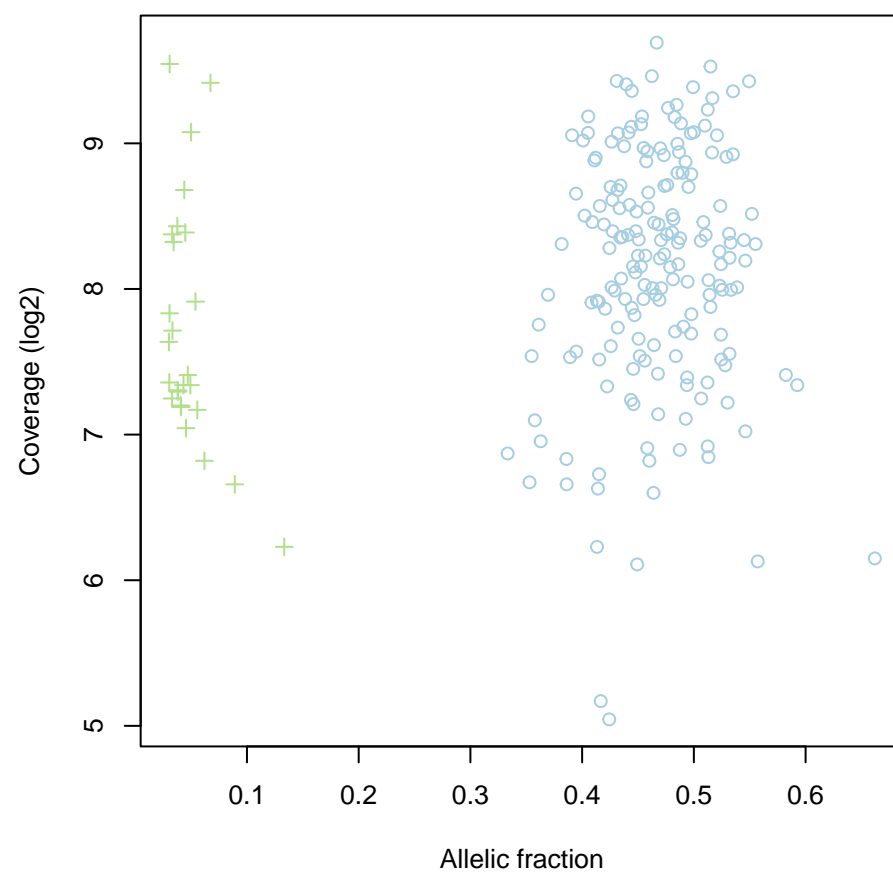
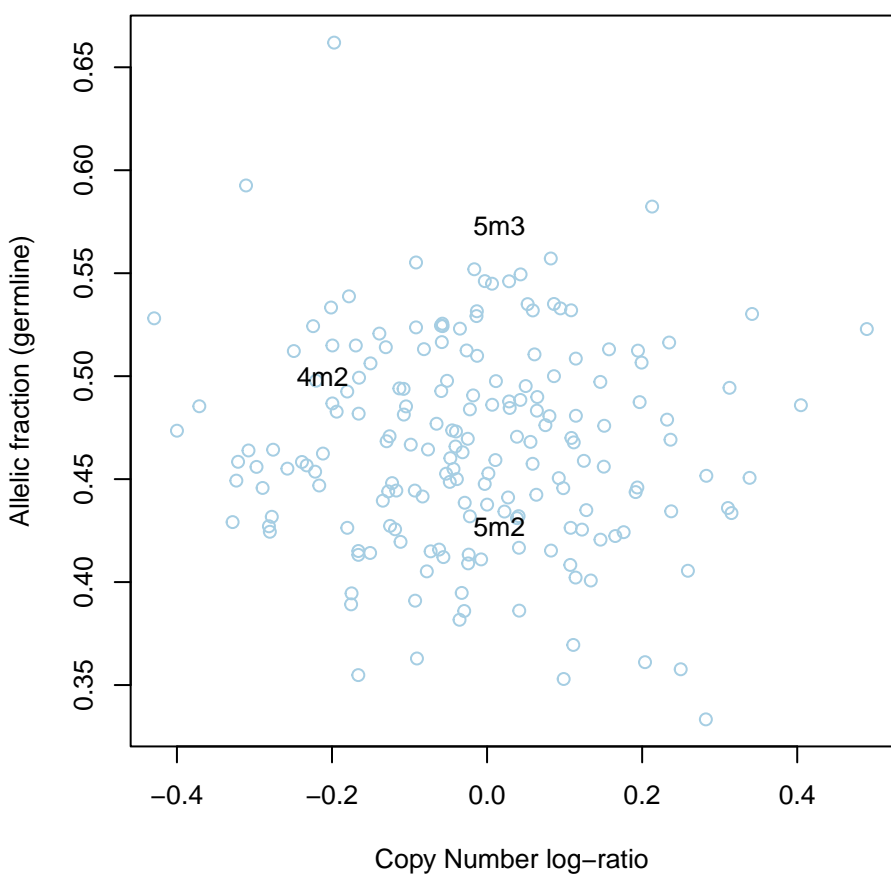


Purity: 0.52 Tumor ploidy: 4.926 SNV log-likelihood: -171.69 GoF: 92.7% Mean coverage: 317,651

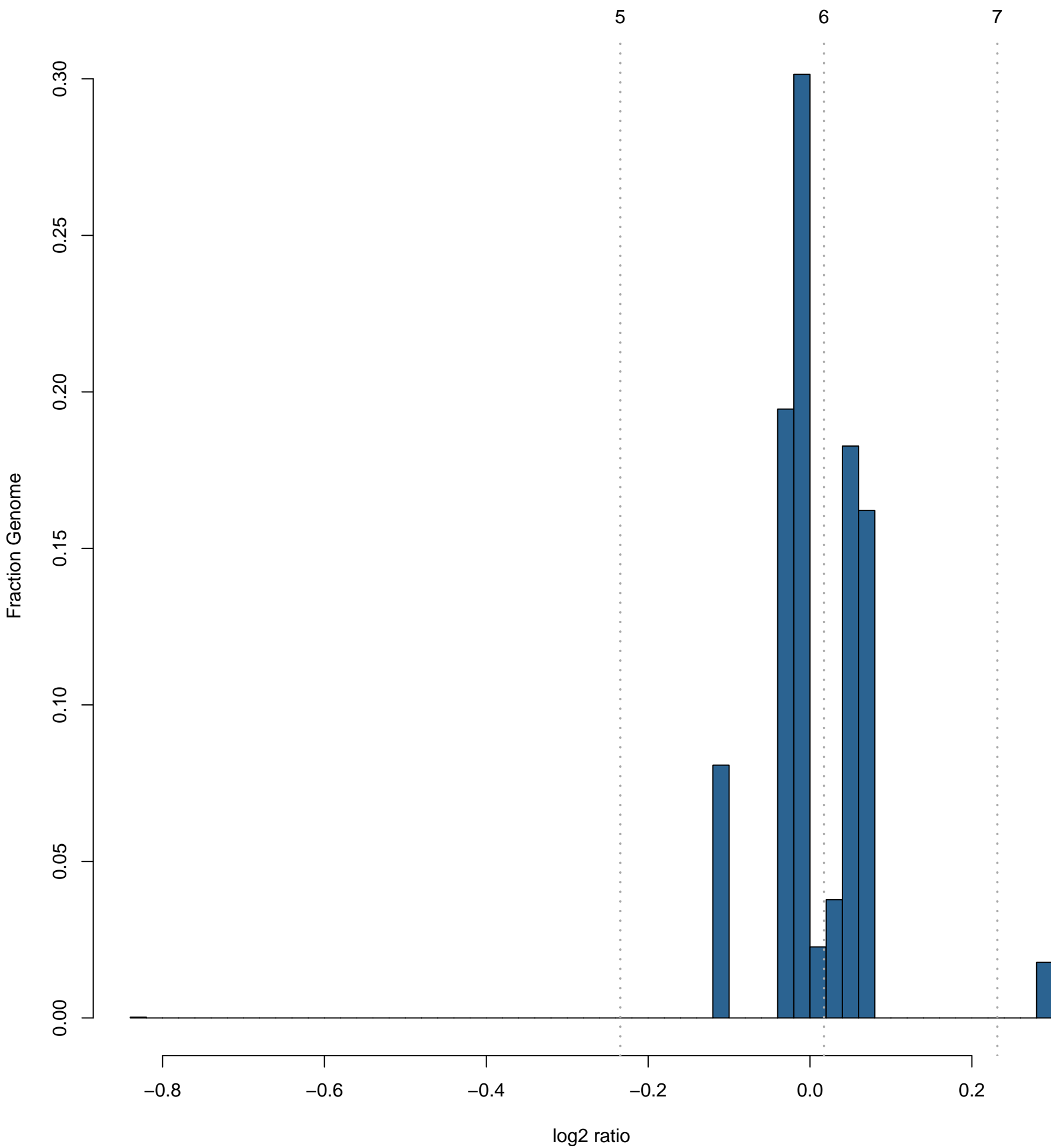


SCNA-fit log-likelihood: -9585.76

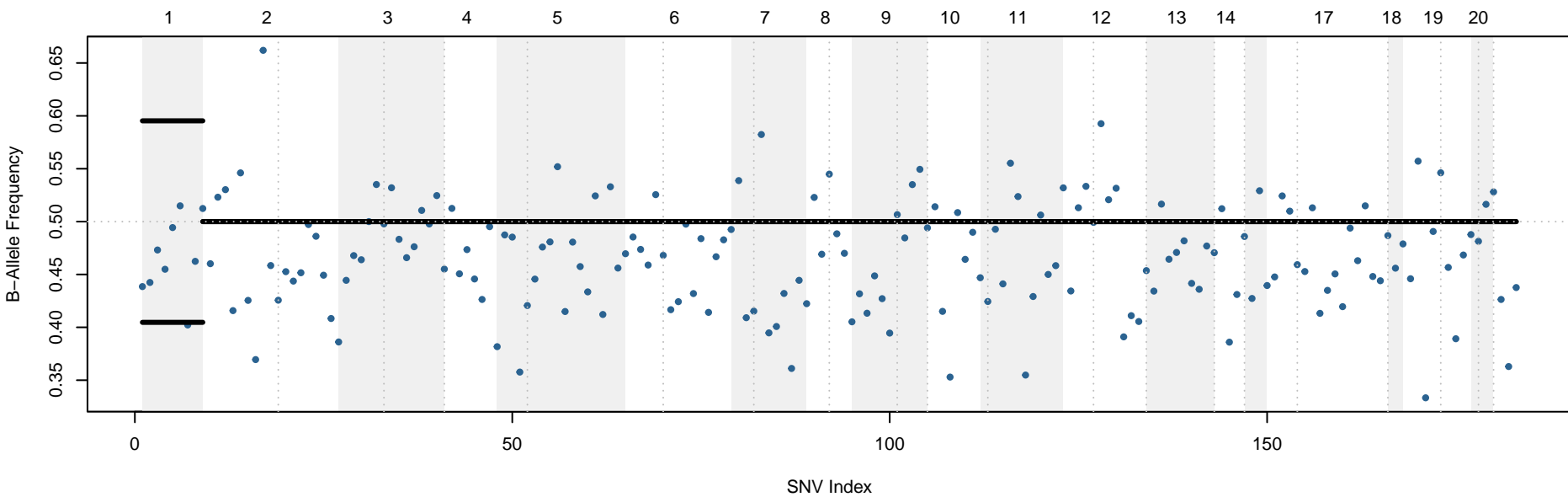




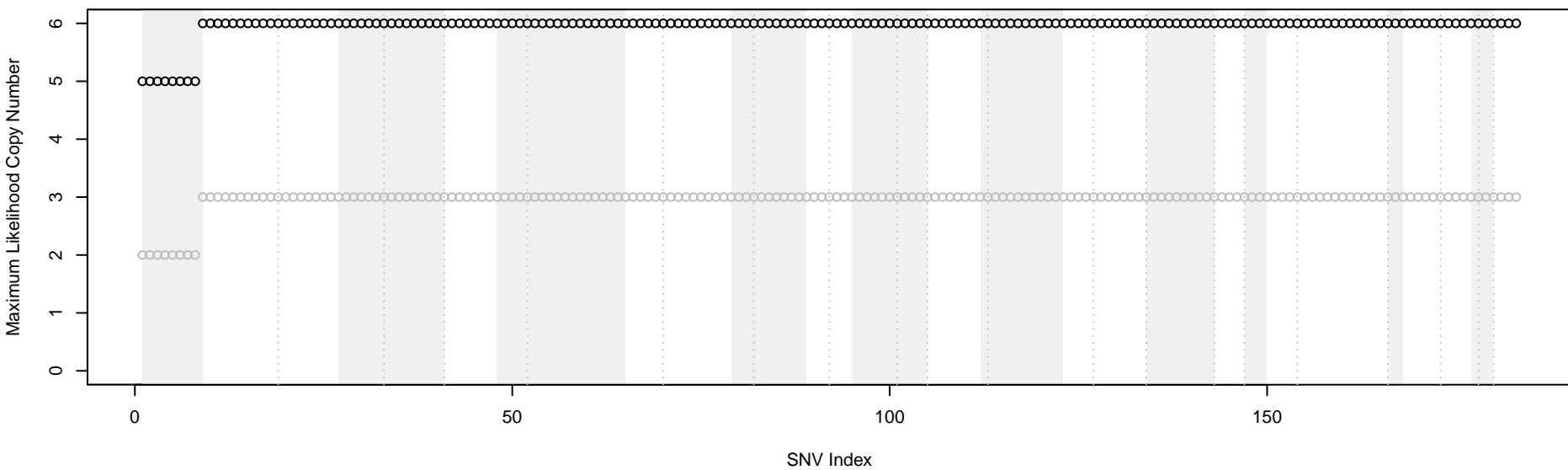
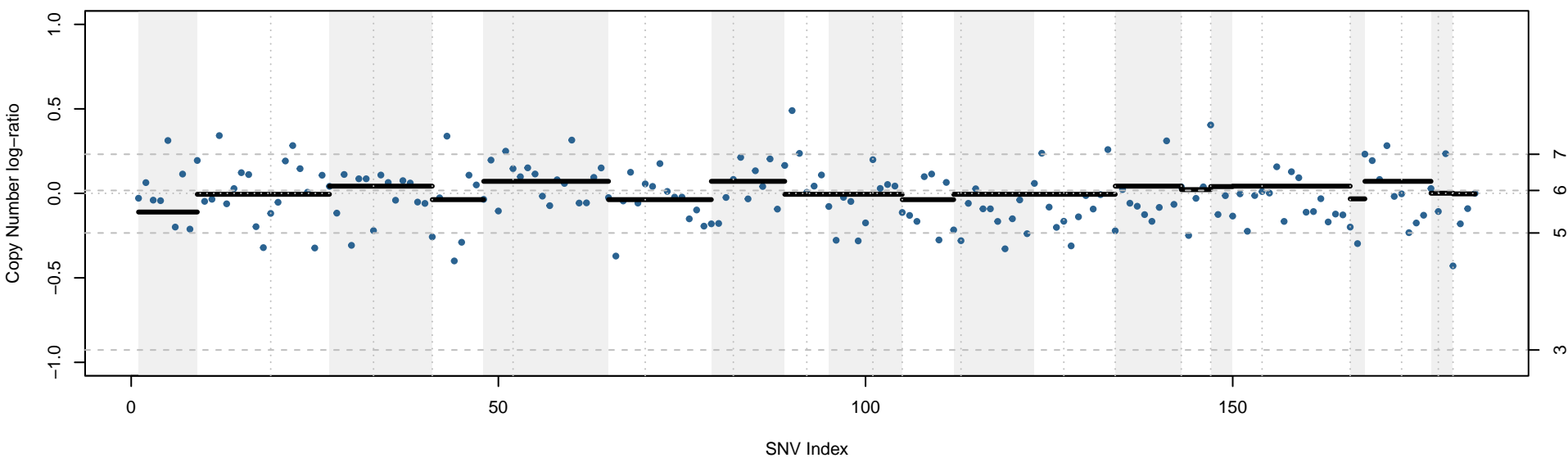
Purity: 0.89 Tumor ploidy: 5.926

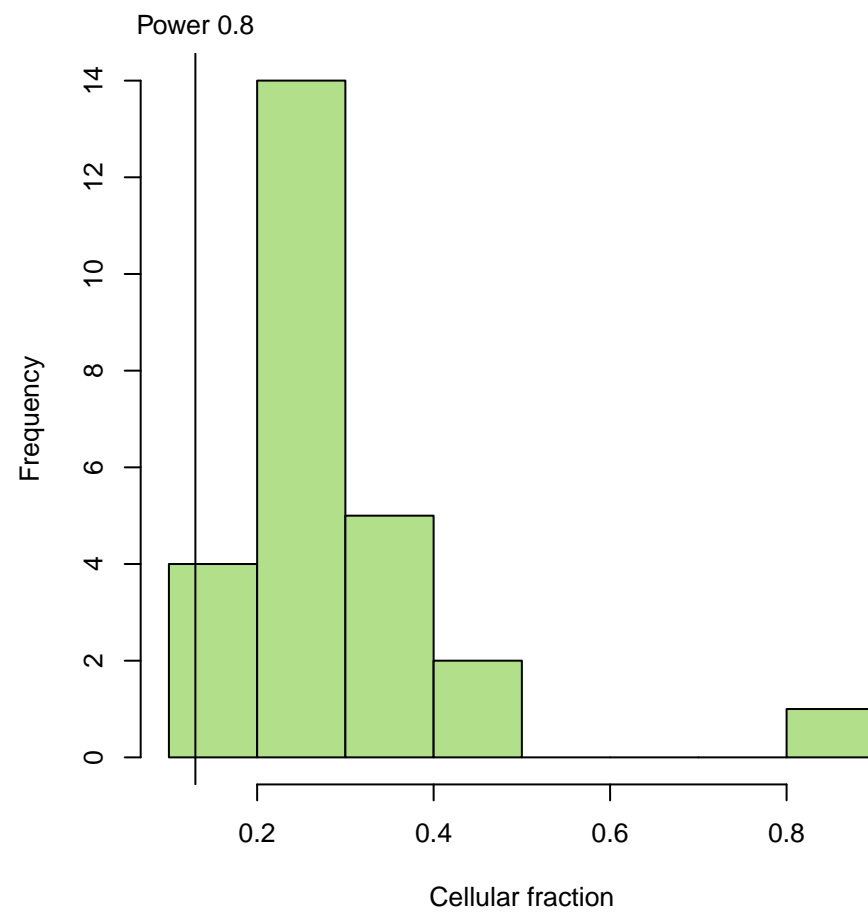
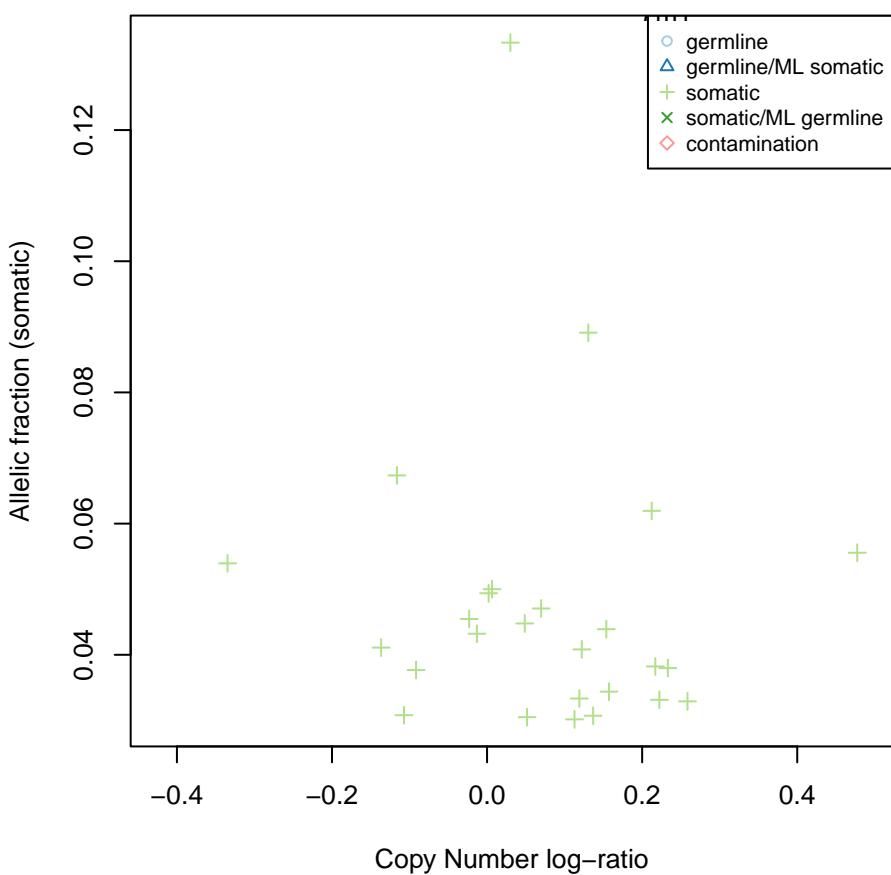
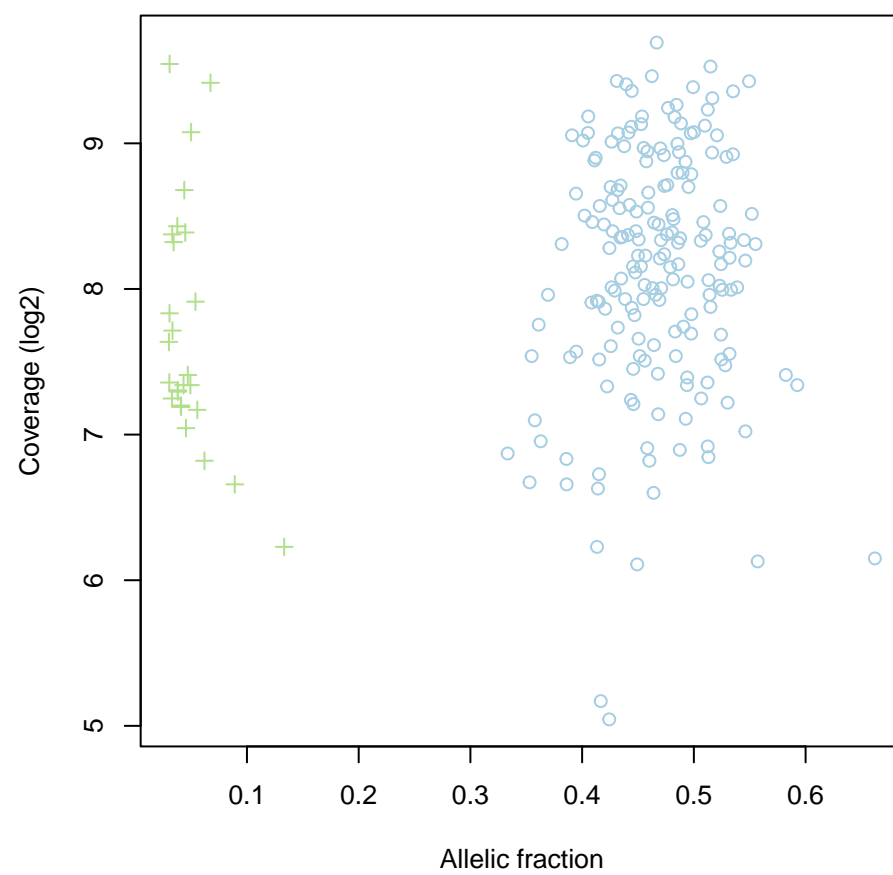
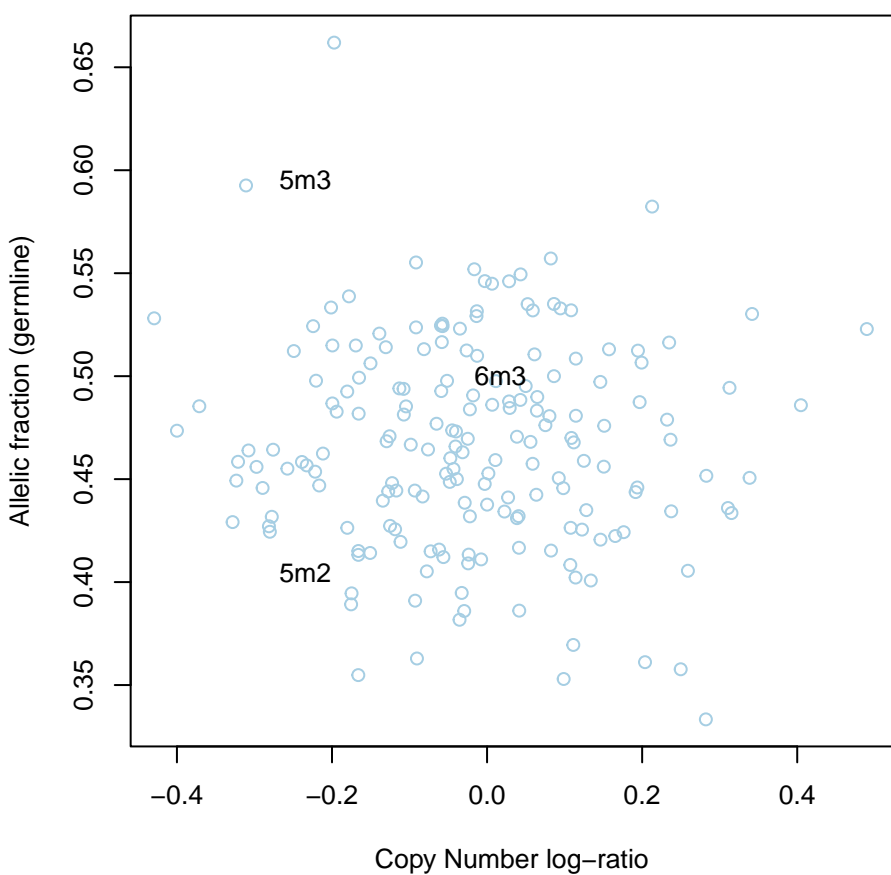


Purity: 0.89 Tumor ploidy: 5.926 SNV log-likelihood: -158.24 GoF: 89.1% Mean coverage: 317,651

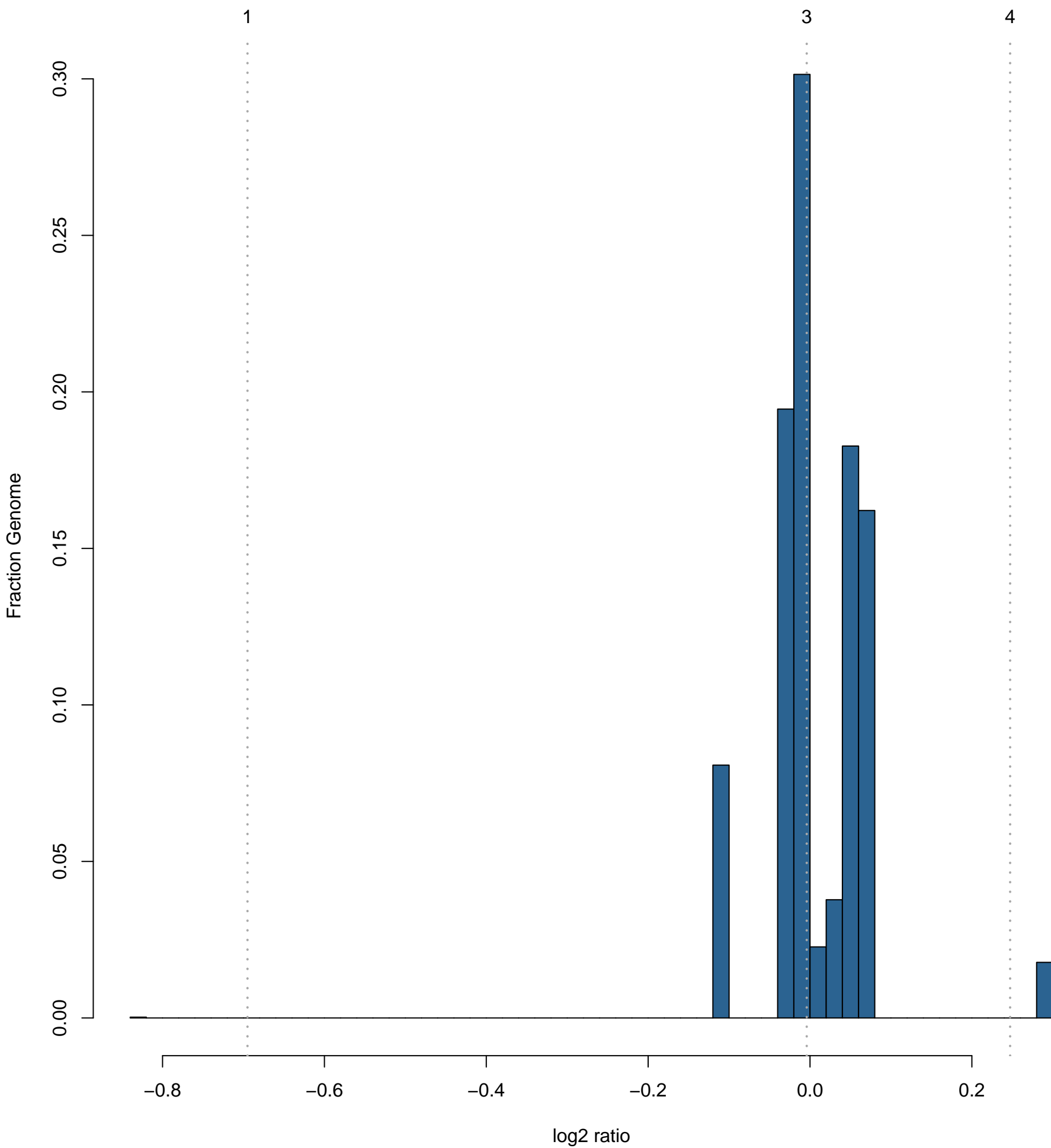


SCNA-fit log-likelihood: -9613.71



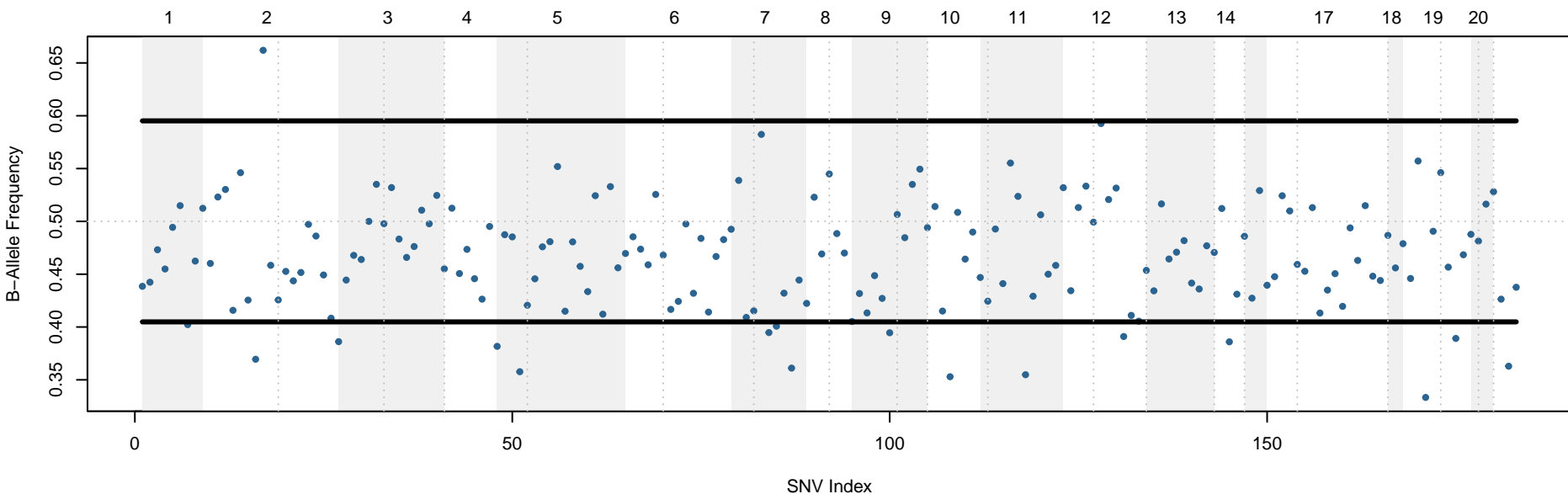


Purity: 0.47 Tumor ploidy: 3.015

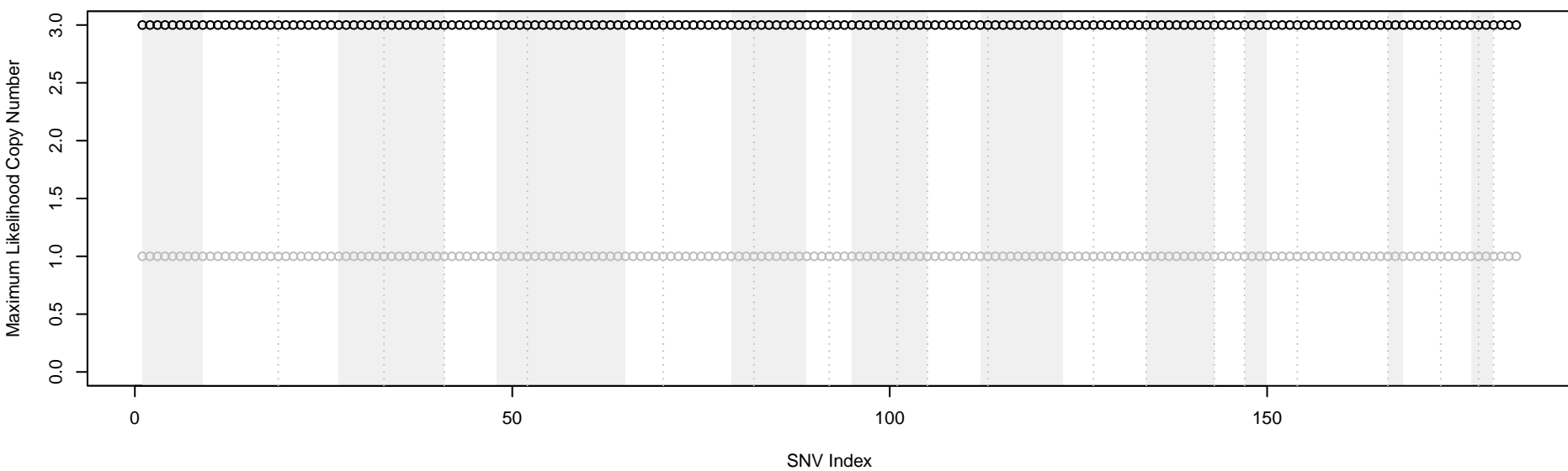
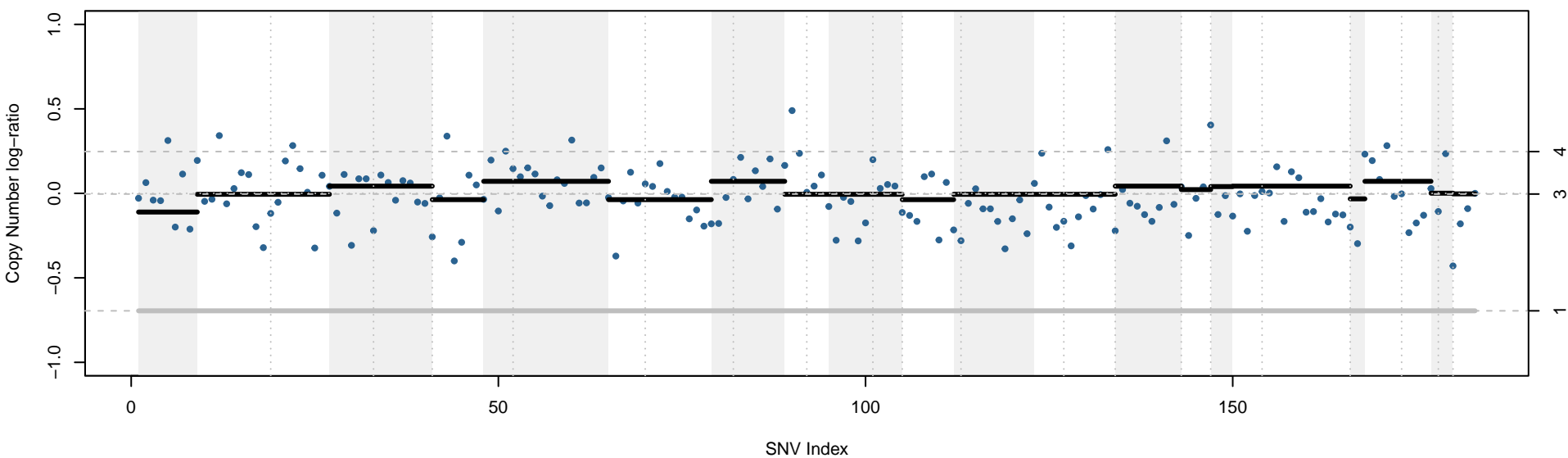


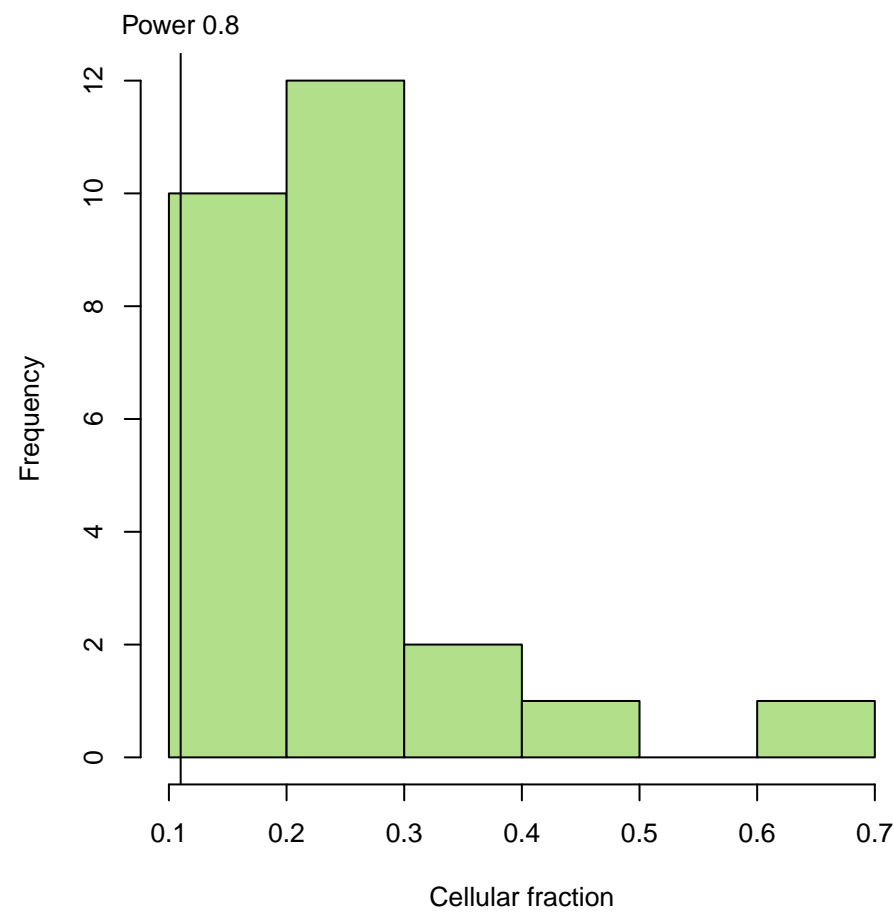
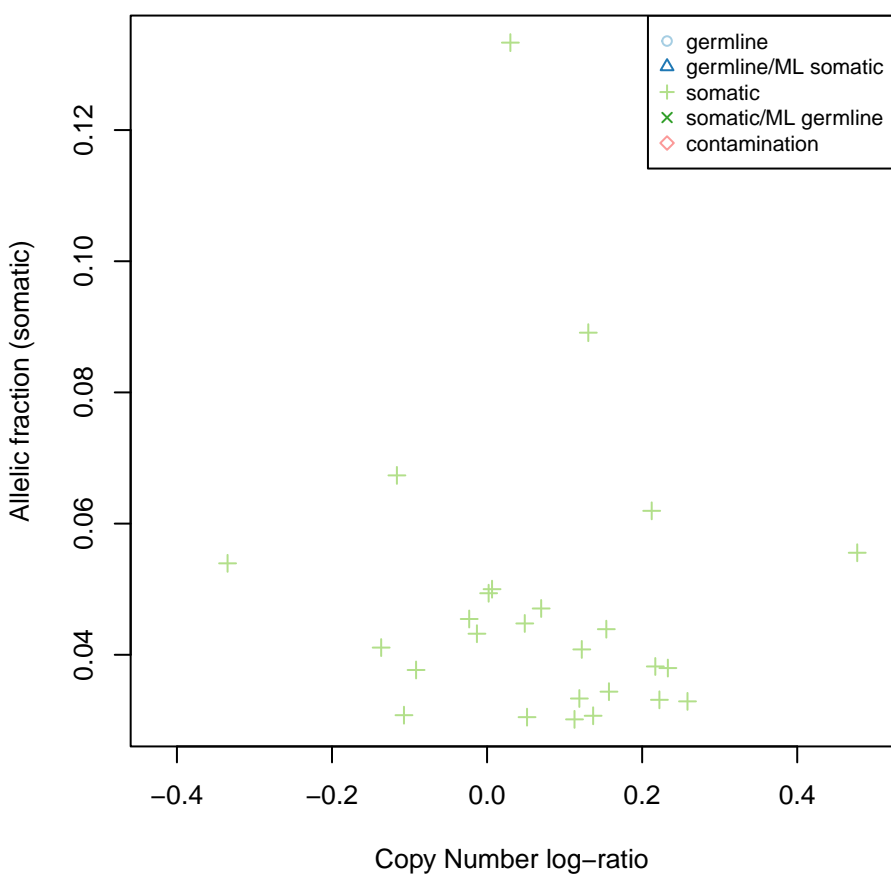
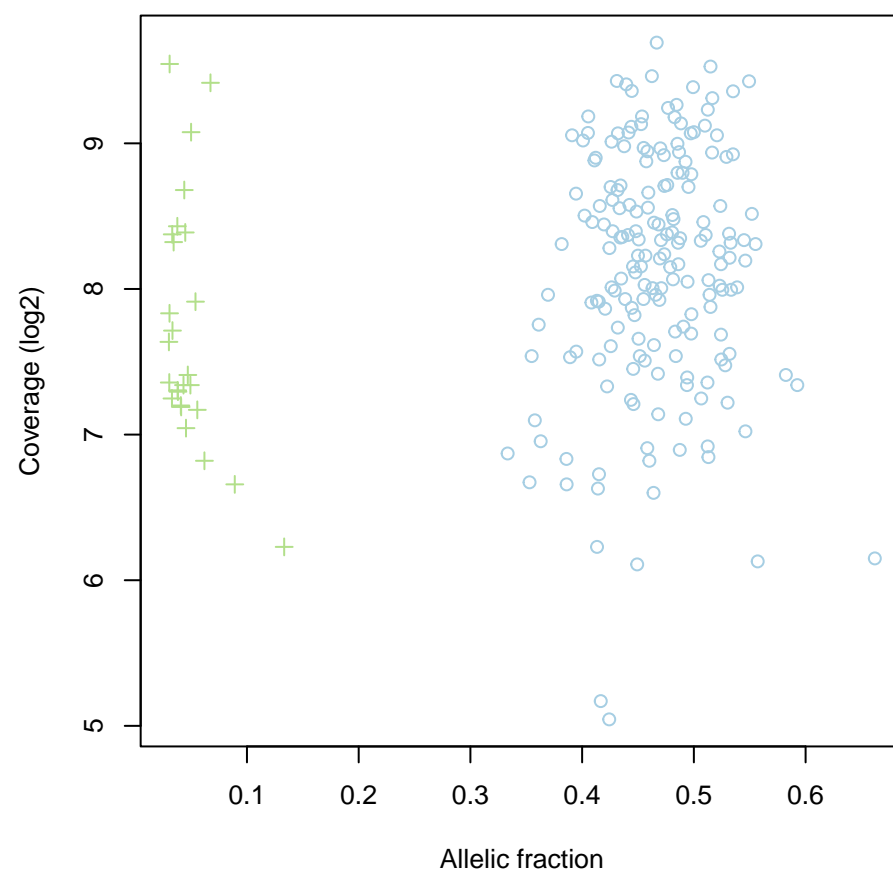
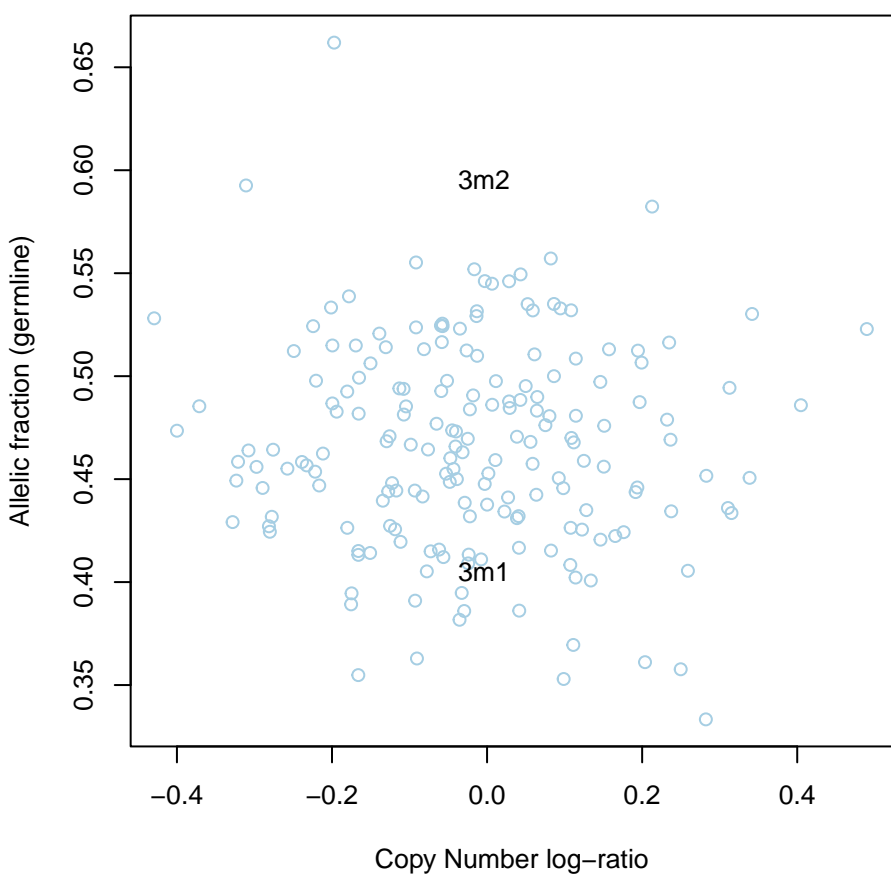


Purity: 0.47 Tumor ploidy: 3.015 SNV log-likelihood: -279.73 GoF: 85.7% Mean coverage: 317,651

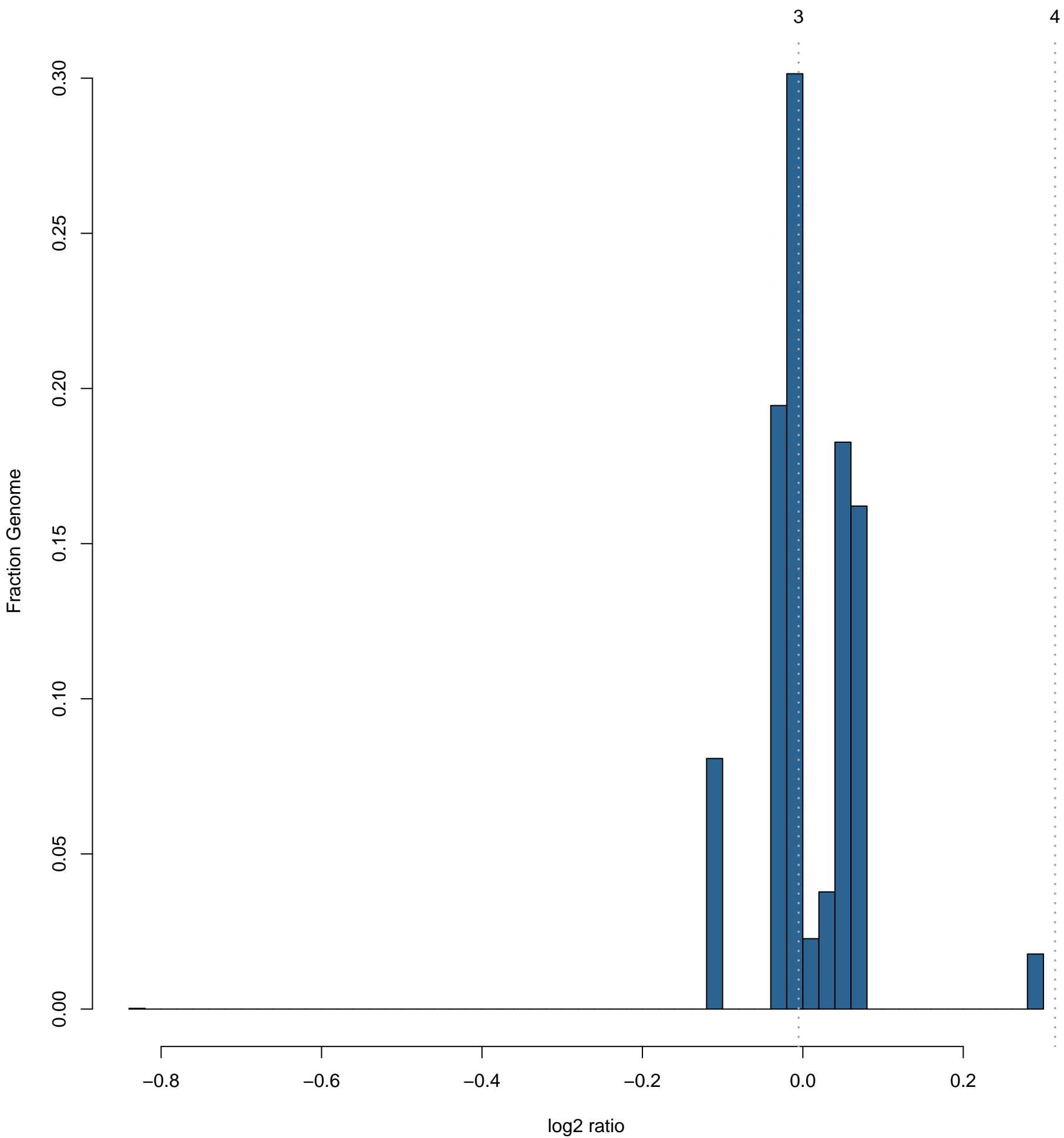


SCNA-fit log-likelihood: -9603.23

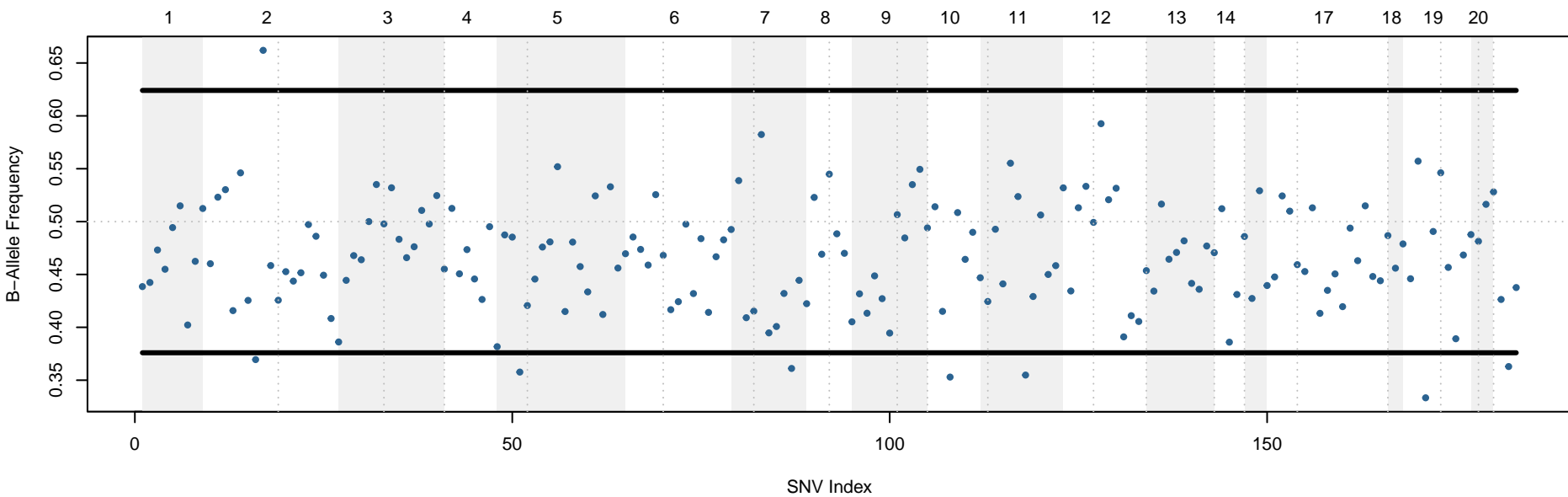




Purity: 0.66 Tumor ploidy: 3.015



Purity: 0.66 Tumor ploidy: 3.015 SNV log-likelihood: -685.21 GoF: 71.6% Mean coverage: 317,651



SCNA-fit log-likelihood: -9603.6

