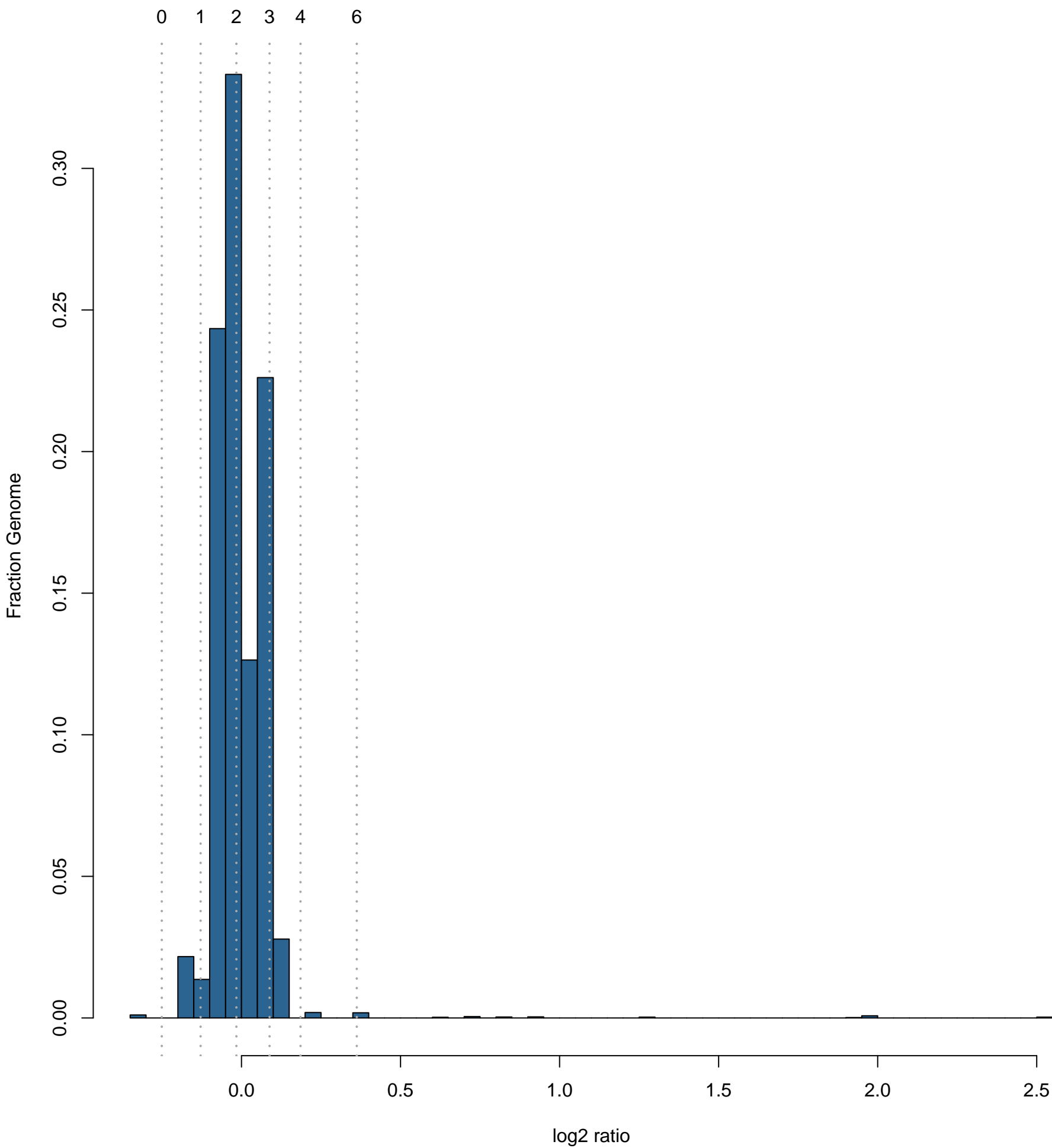
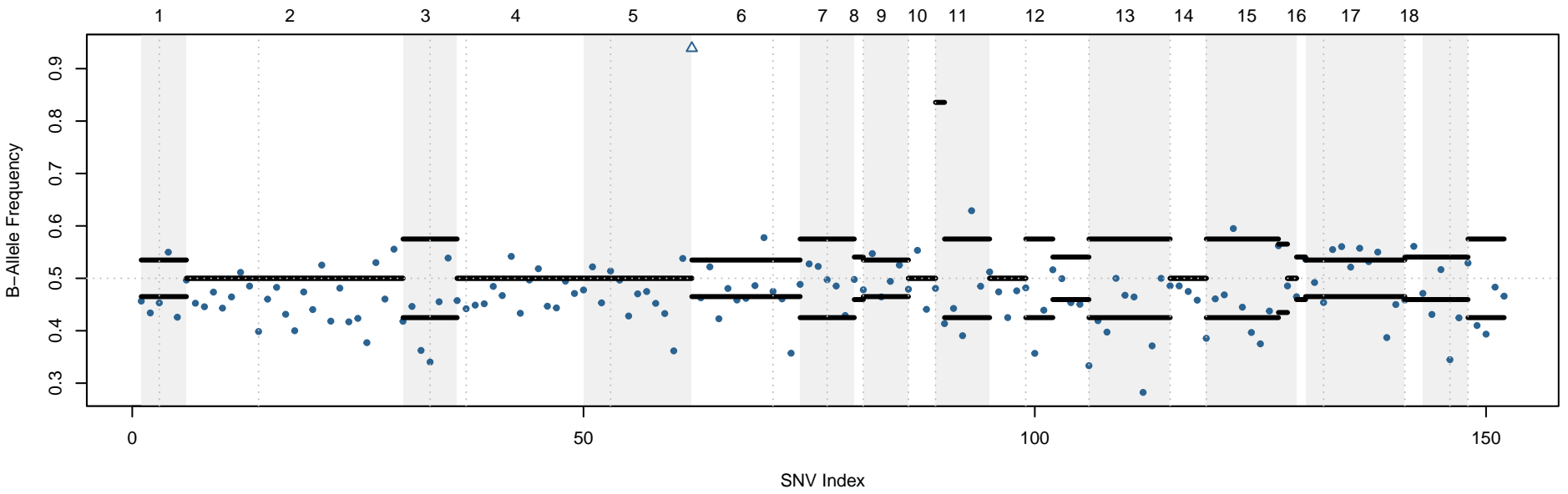


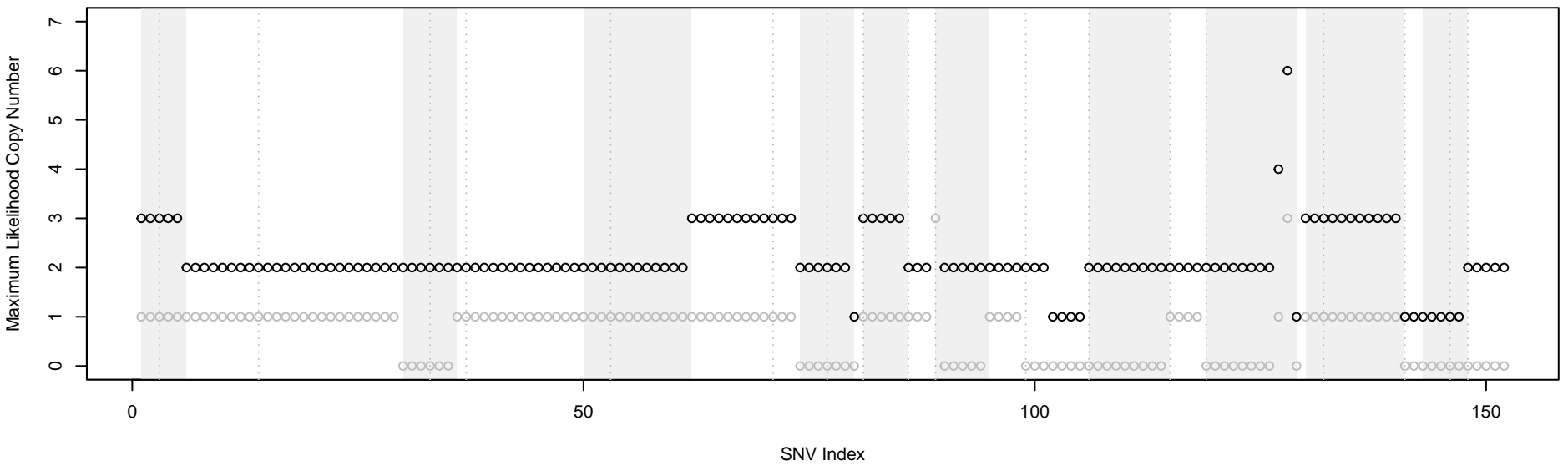
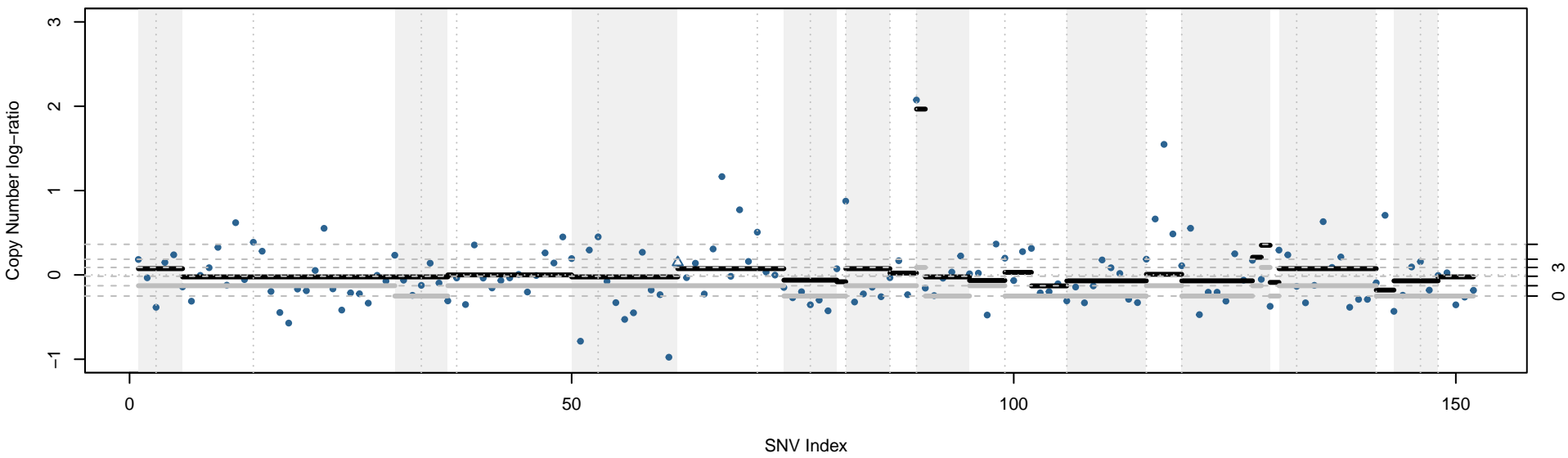
Purity: 0.15 Tumor ploidy: 2.149

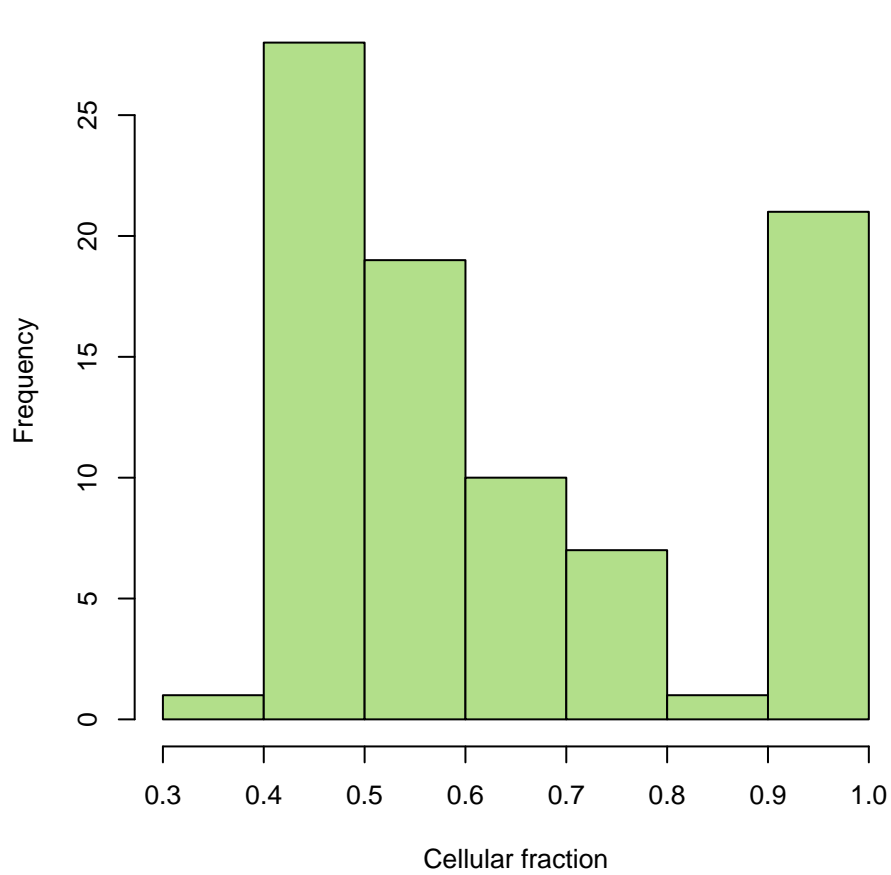
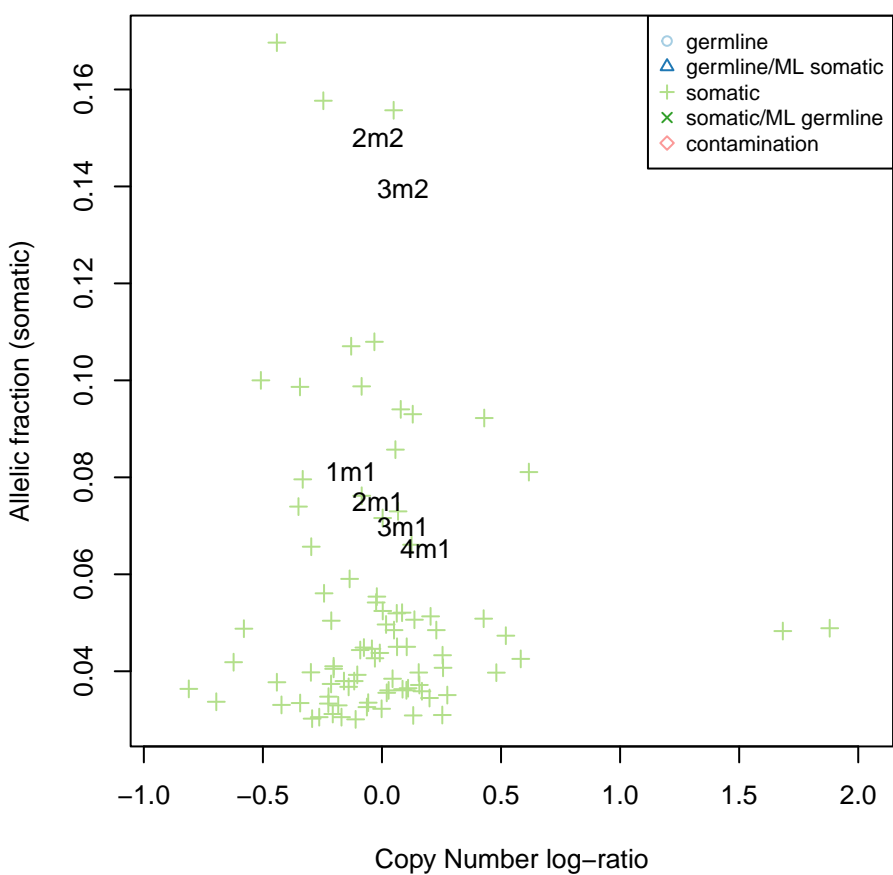
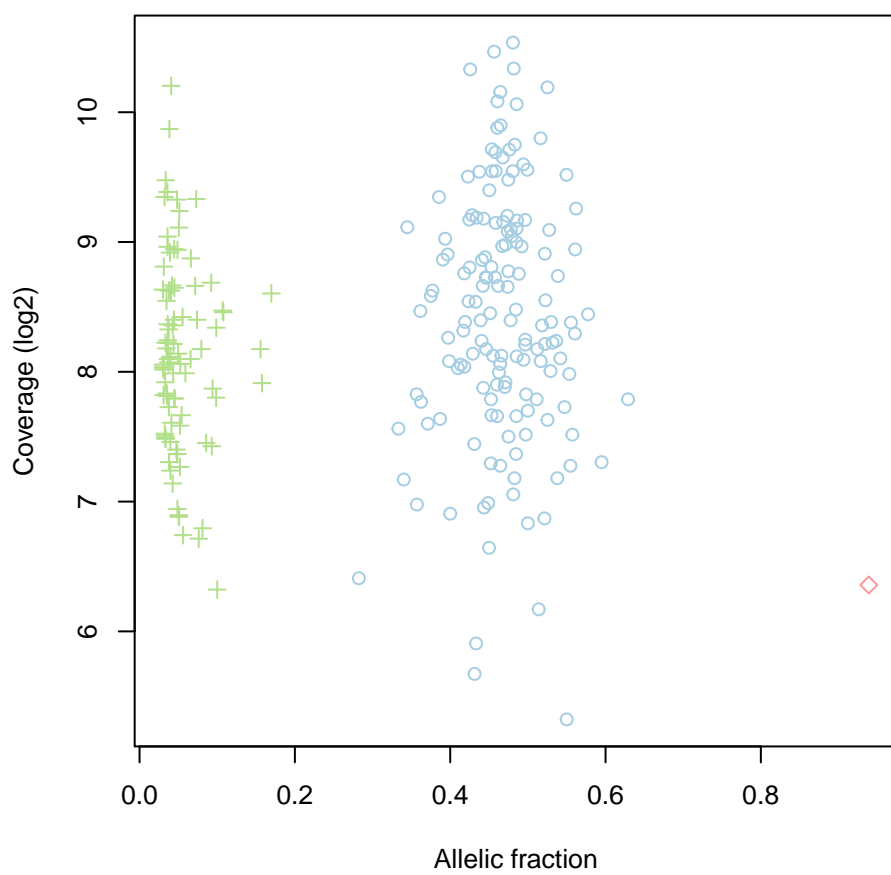
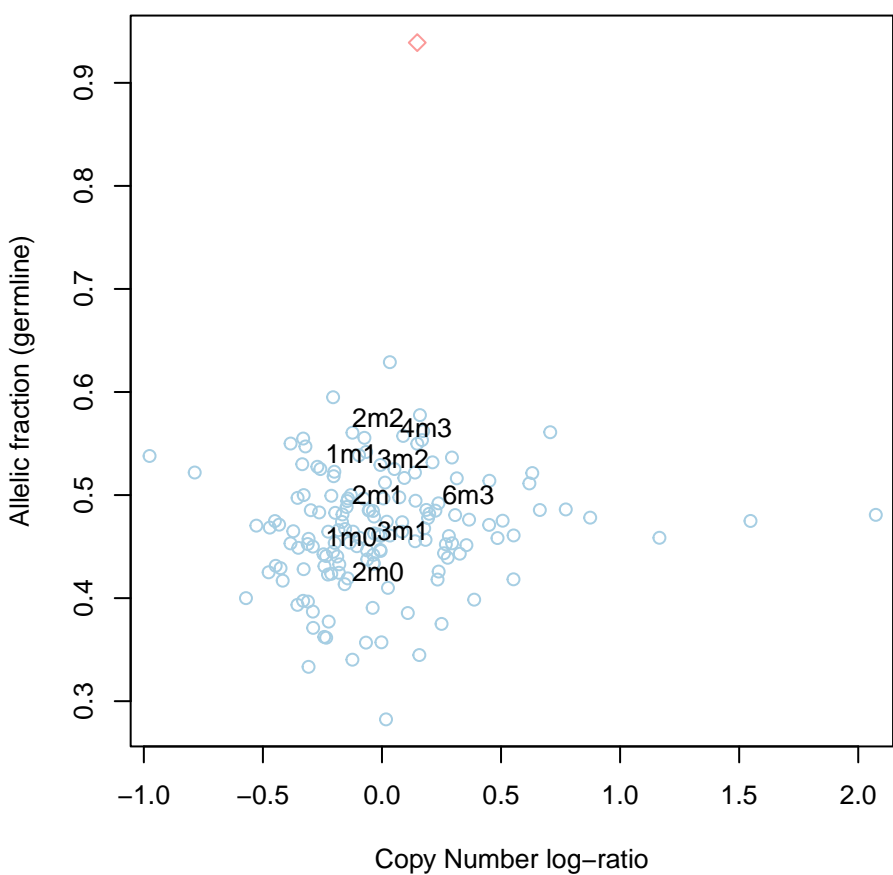


Purity: 0.15 Tumor ploidy: 2.149 SNV log-likelihood: -16.21 GoF: 95.1% Mean coverage: 108;387

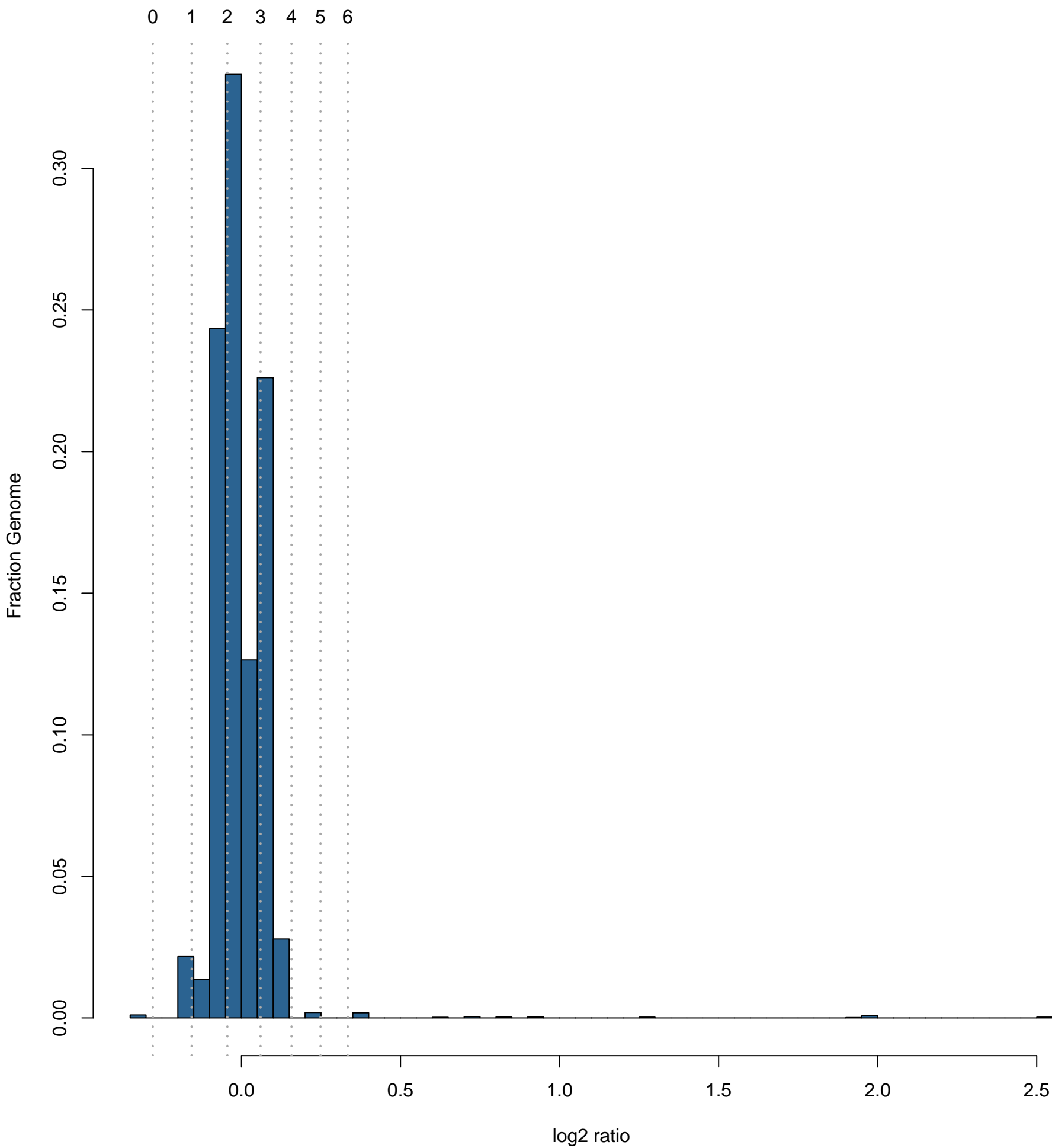


SCNA-fit log-likelihood: -15714.62

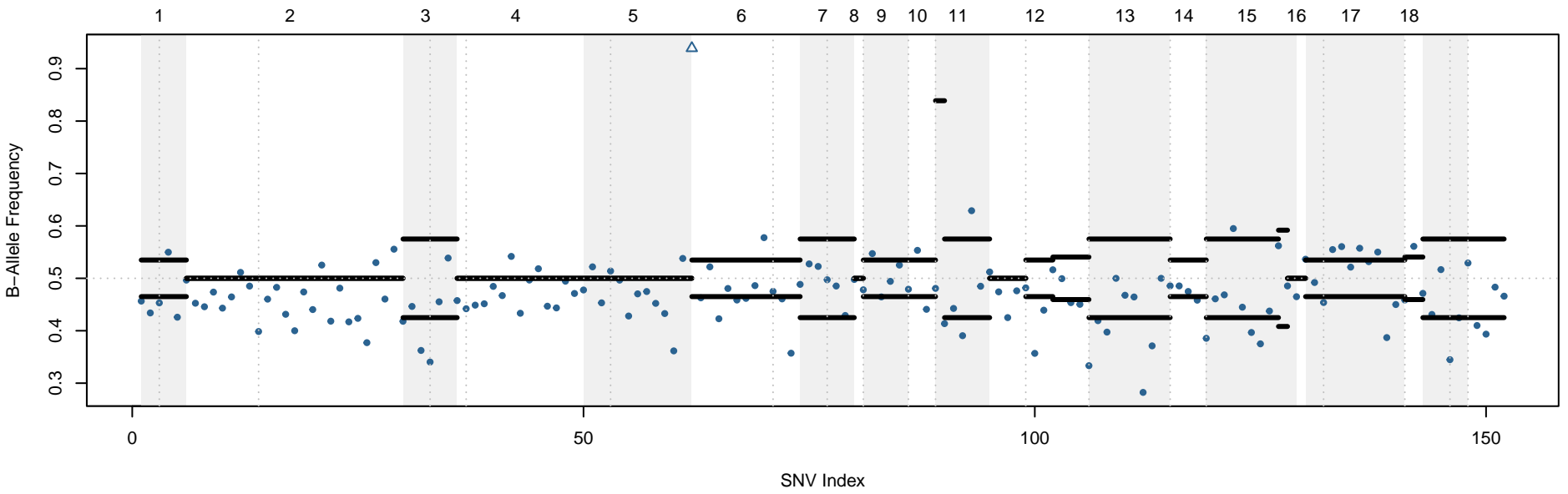




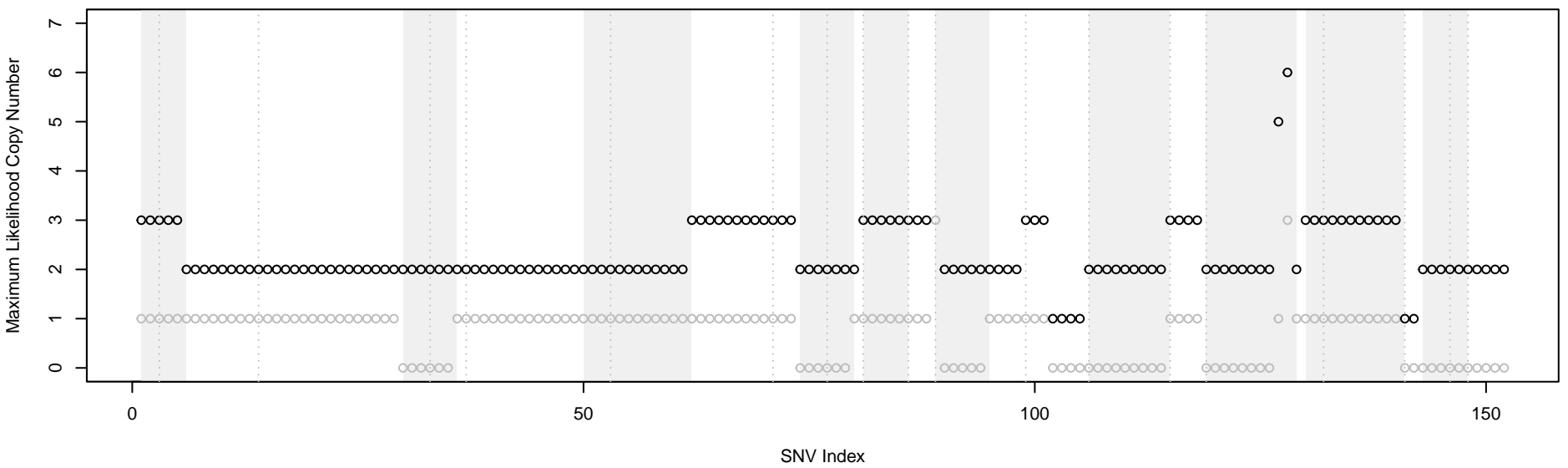
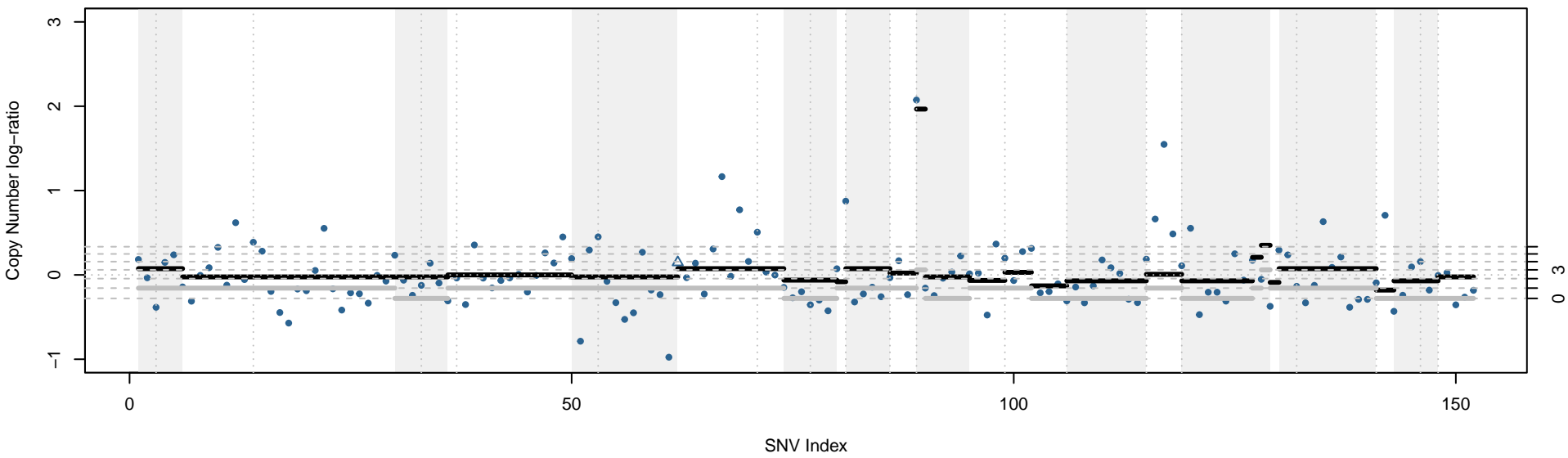
Purity: 0.15 Tumor ploidy: 2.416

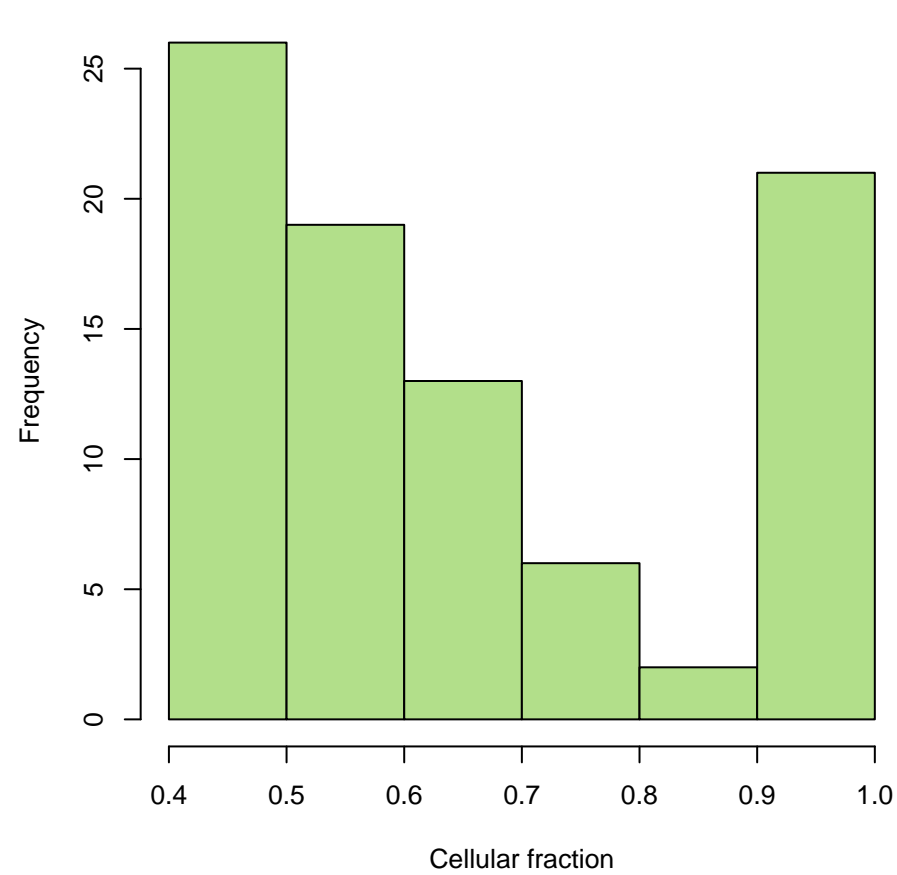
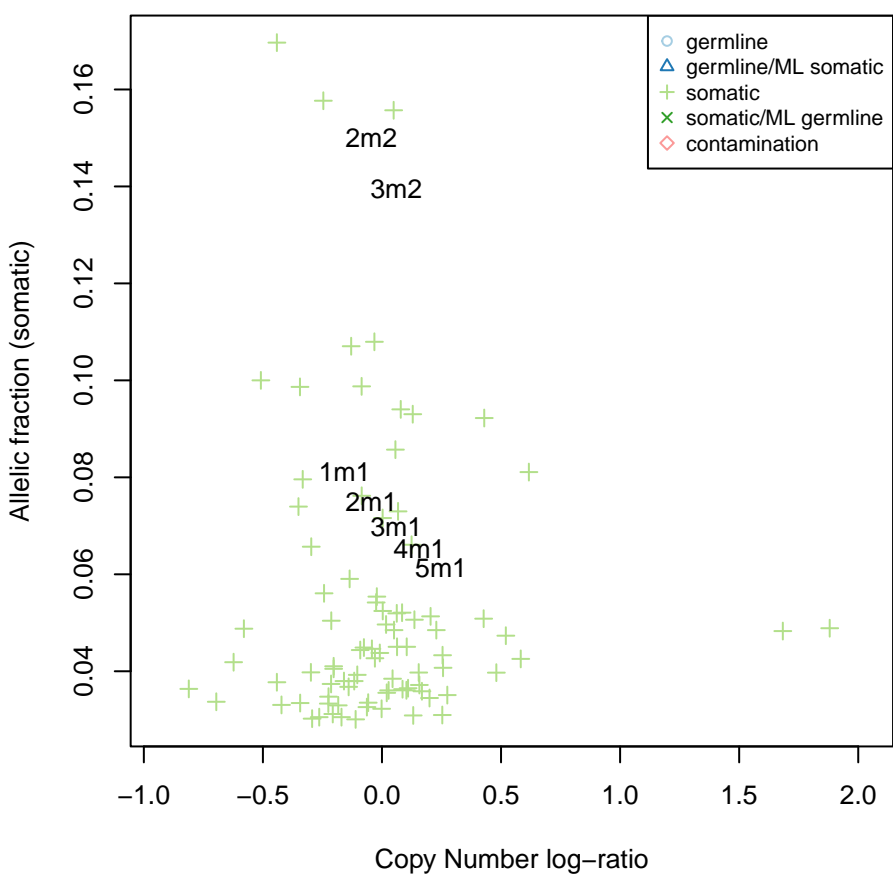
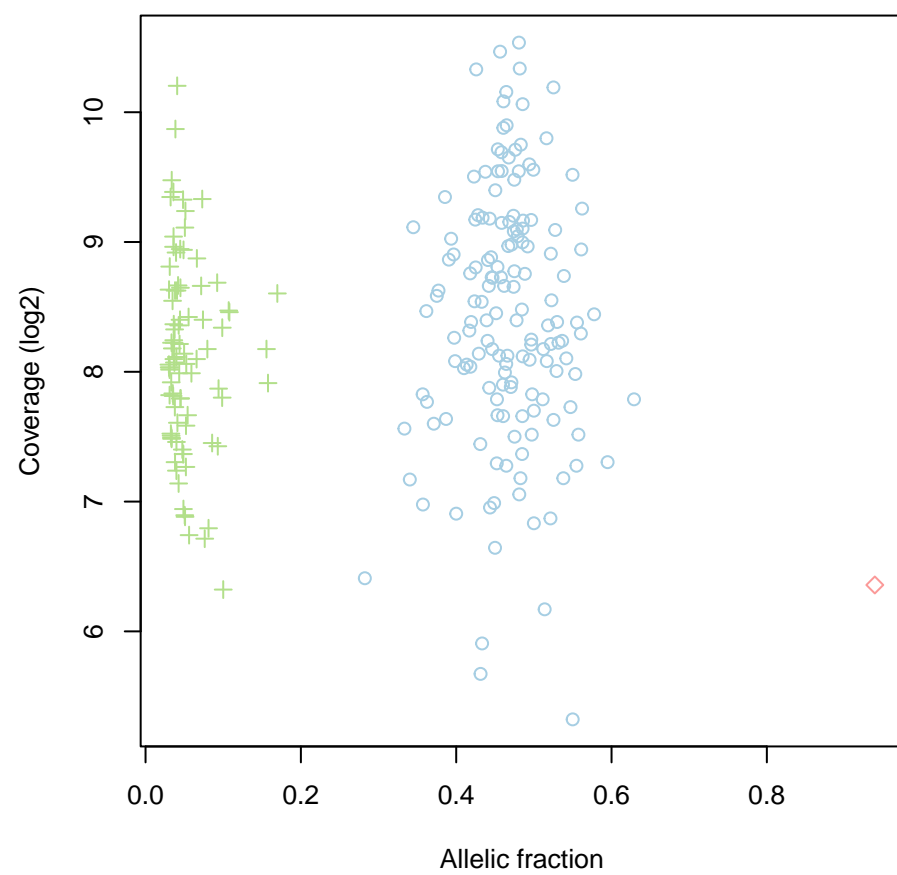
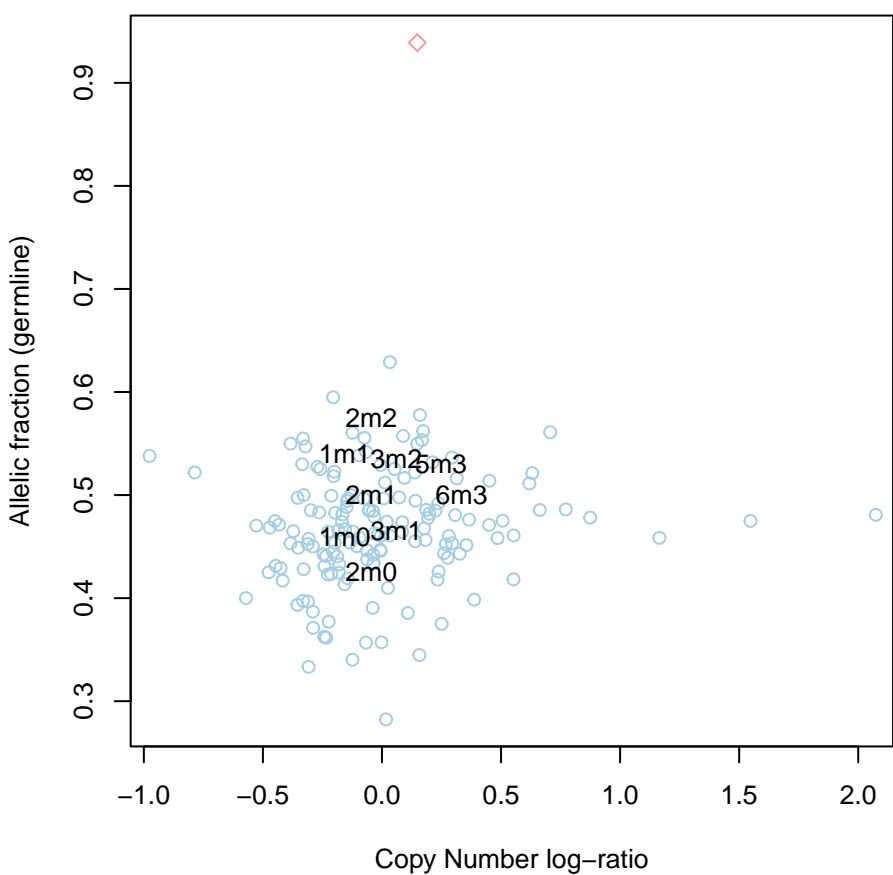


Purity: 0.15 Tumor ploidy: 2.416 SNV log-likelihood: -32.93 GoF: 95.6% Mean coverage: 108;387

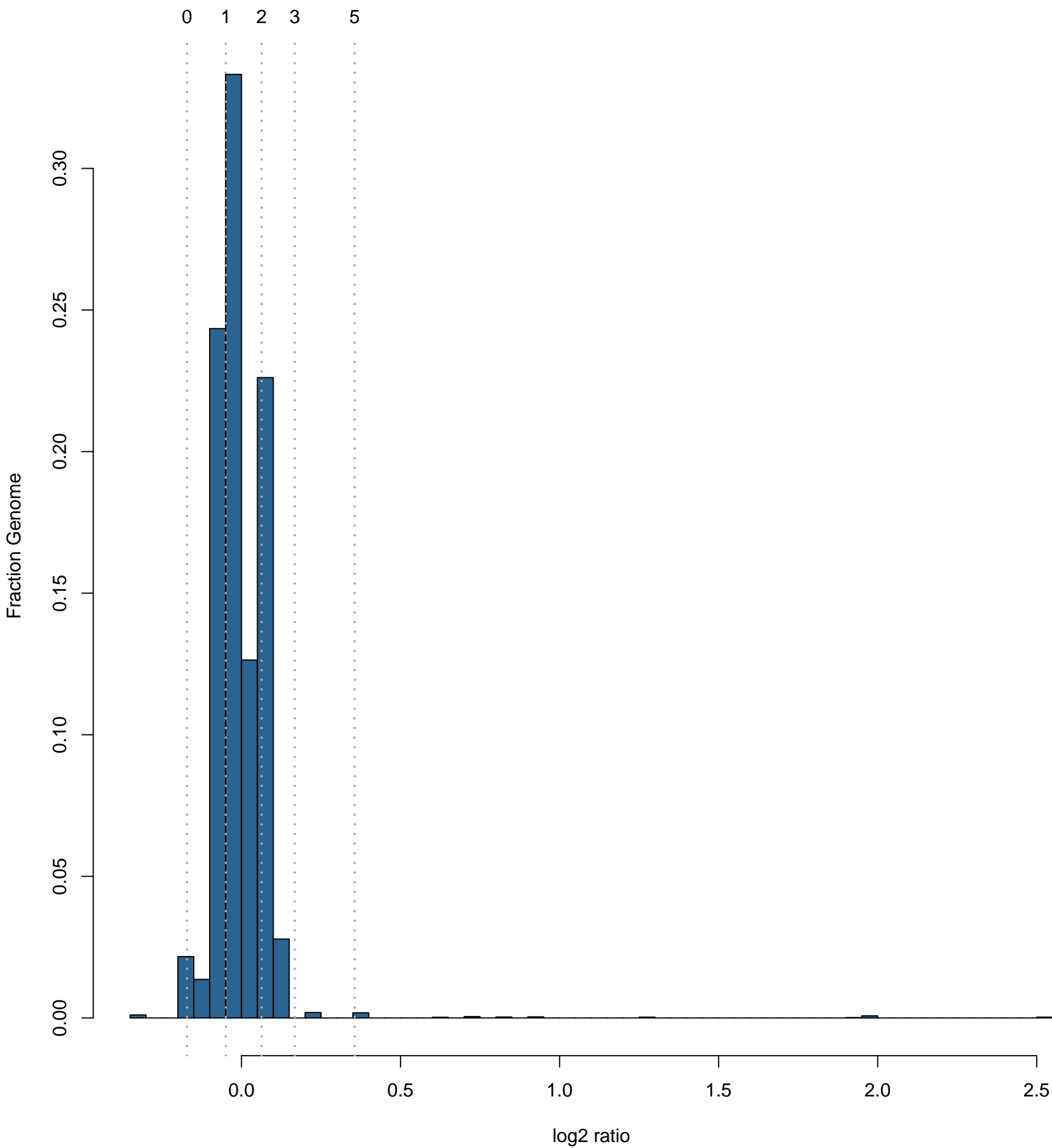


SCNA-fit log-likelihood: -15713.34

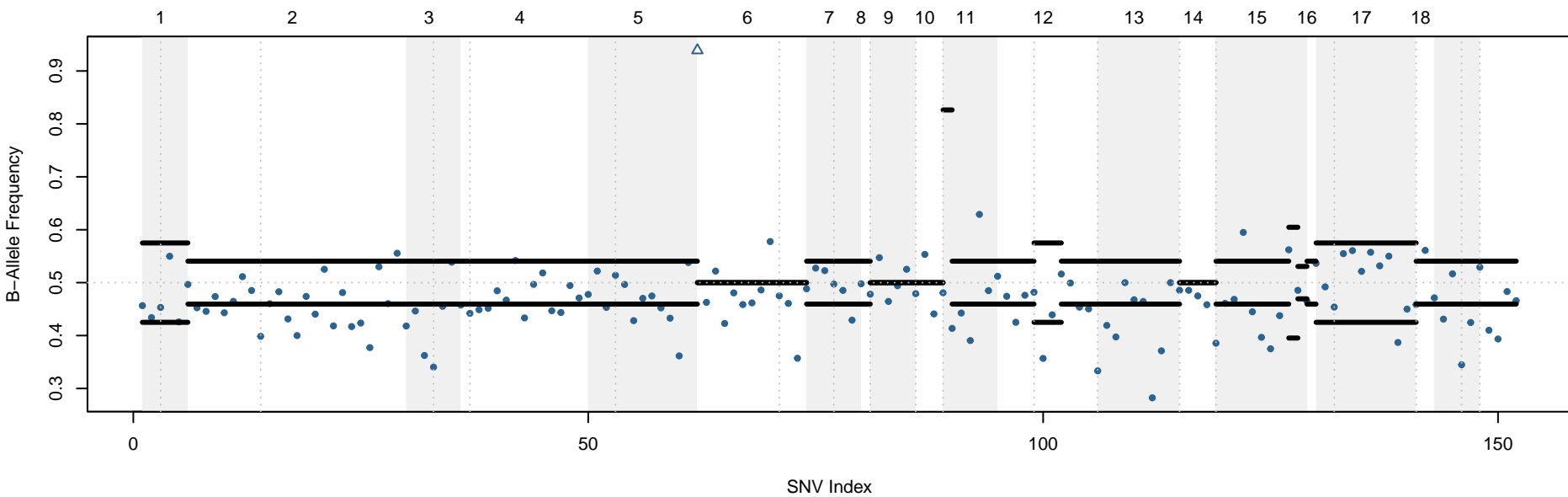




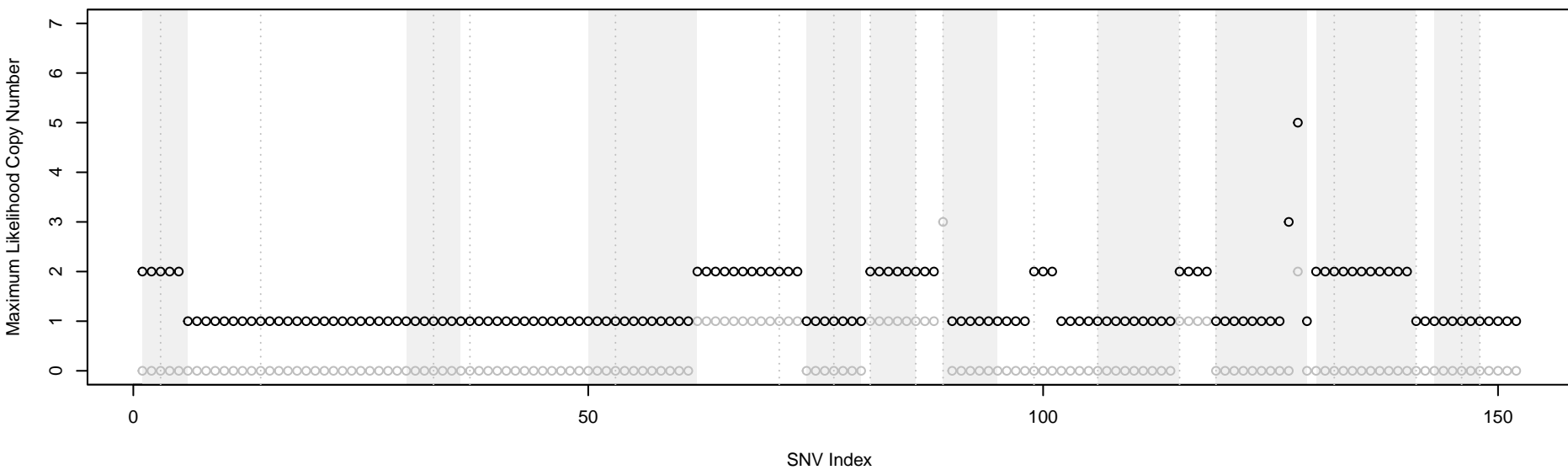
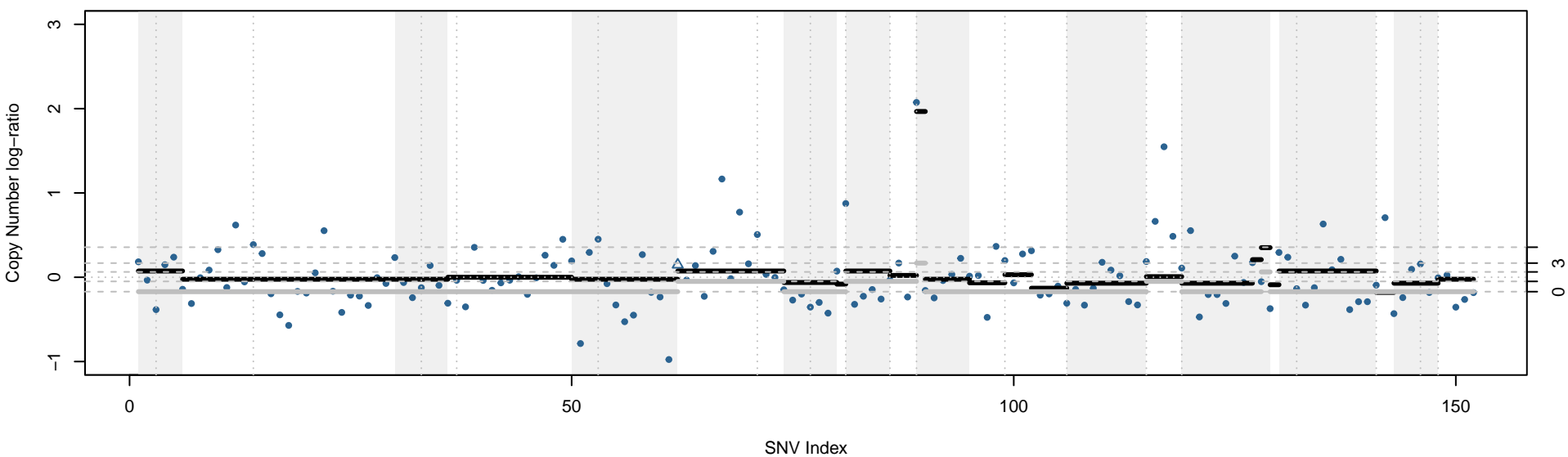
Purity: 0.15 Tumor ploidy: 1.428

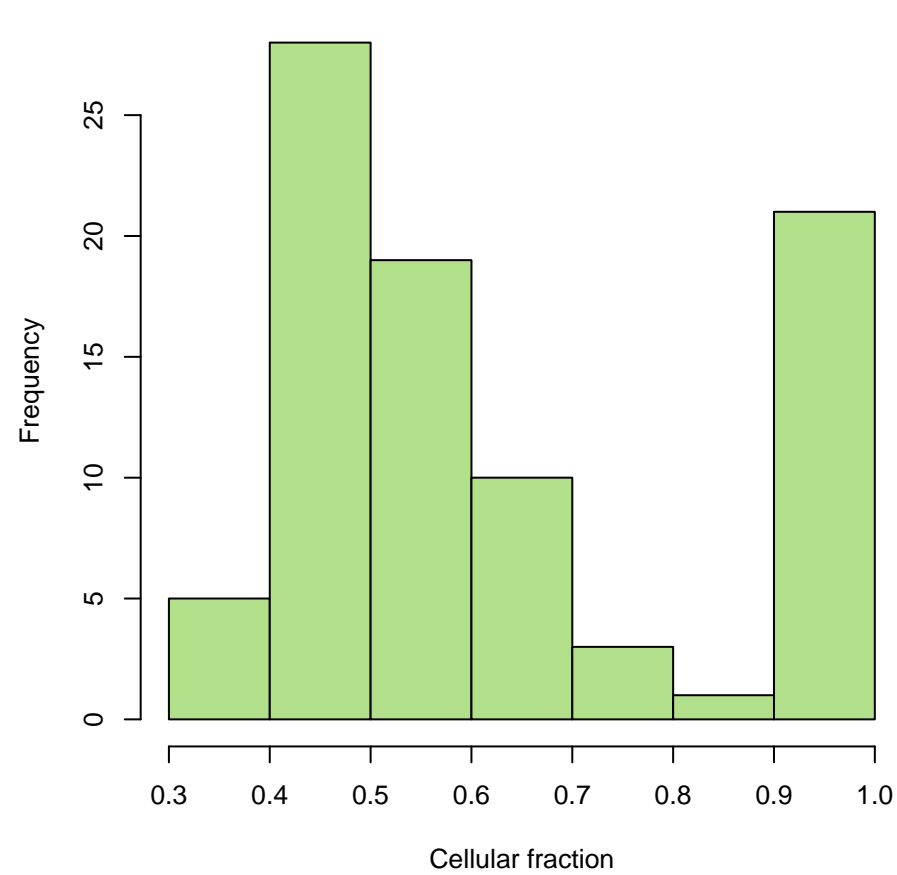
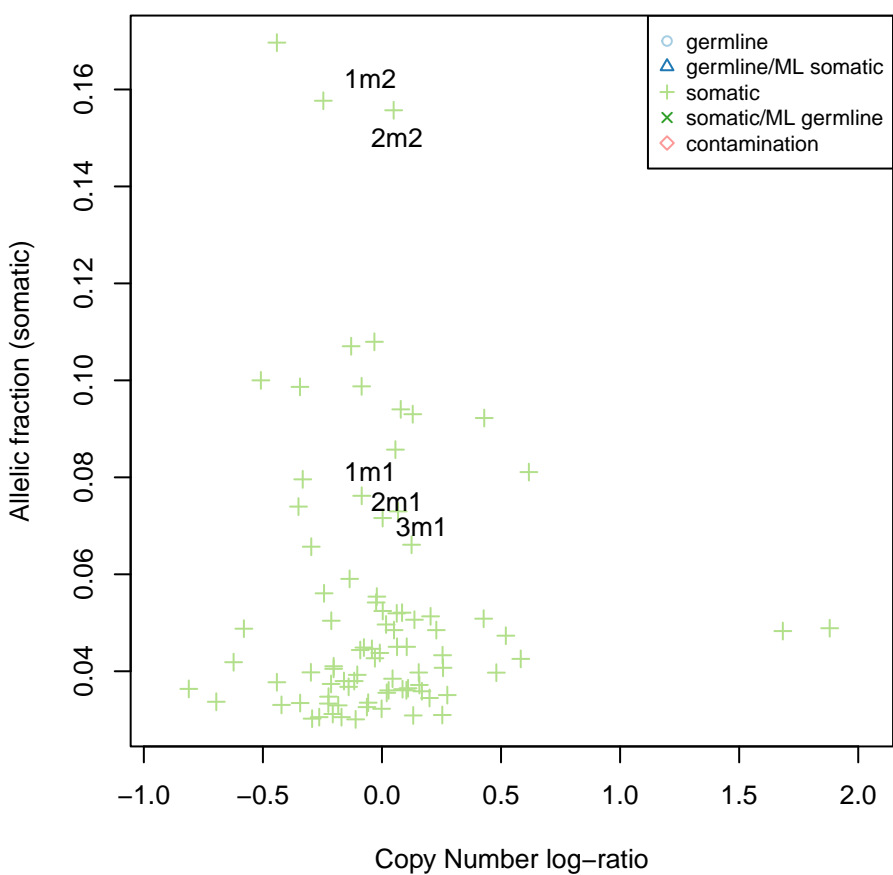
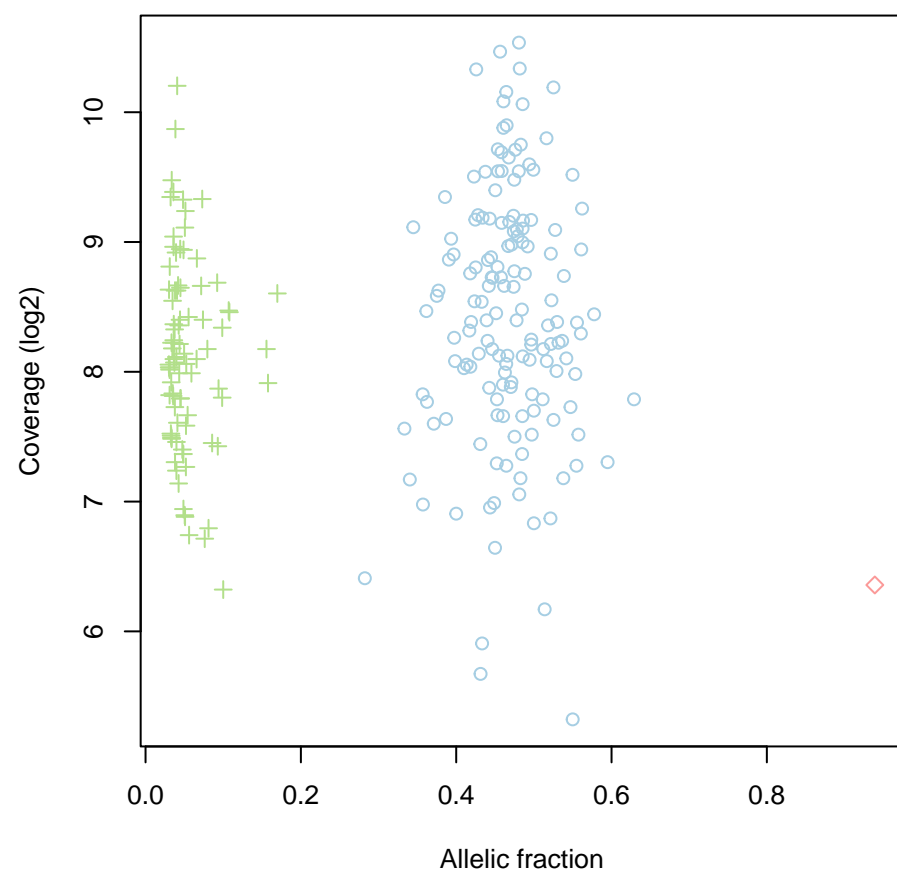
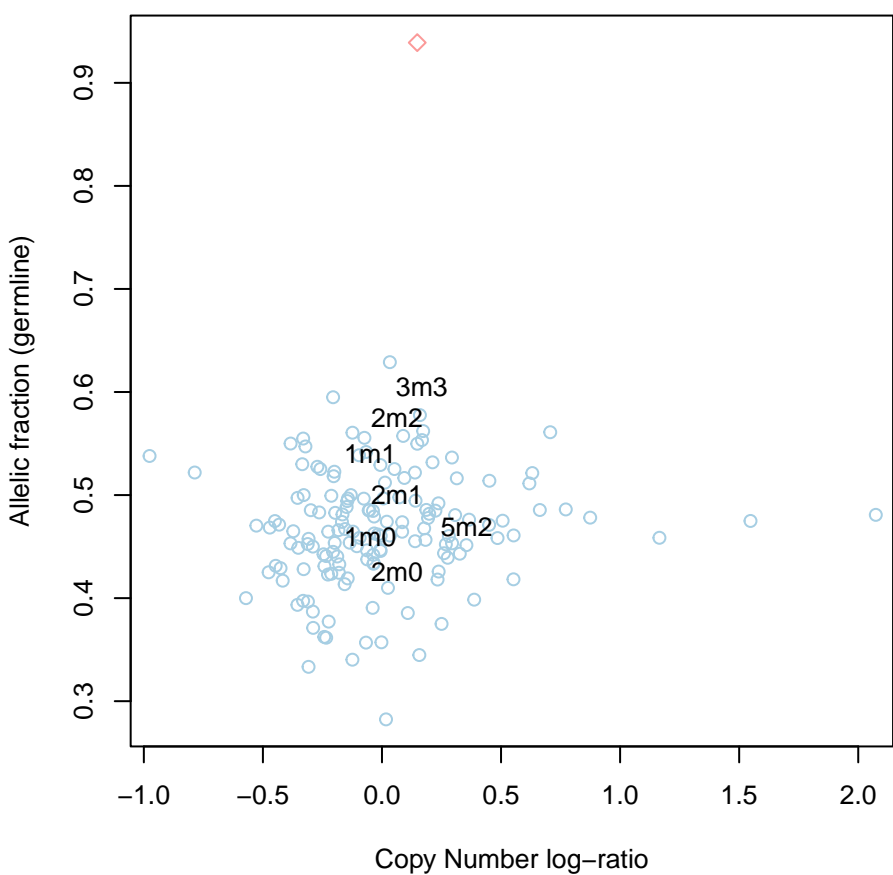


Purity: 0.15 Tumor ploidy: 1.428 SNV log-likelihood: -93.07 GoF: 95.5% Mean coverage: 108;387



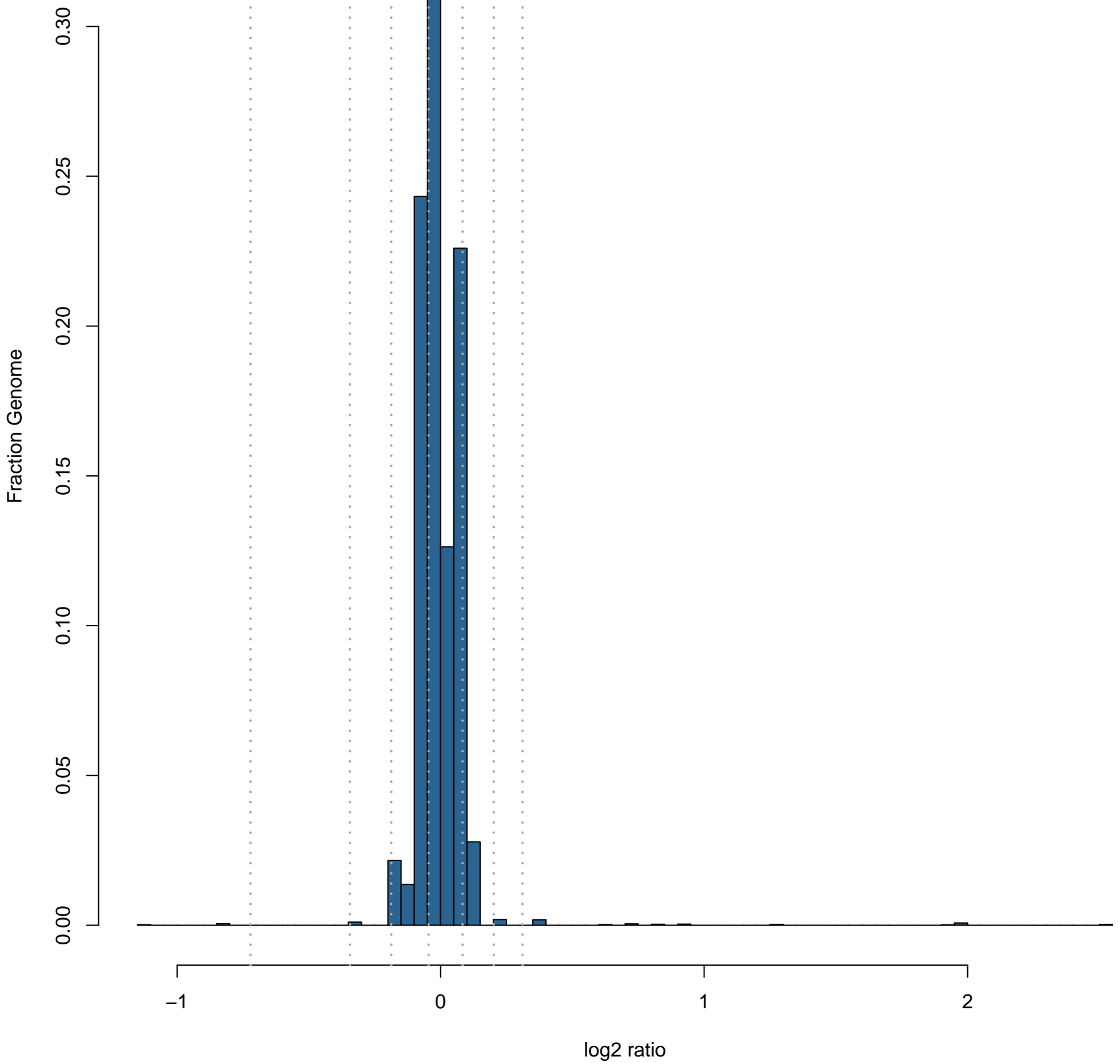
SCNA-fit log-likelihood: -15735.17



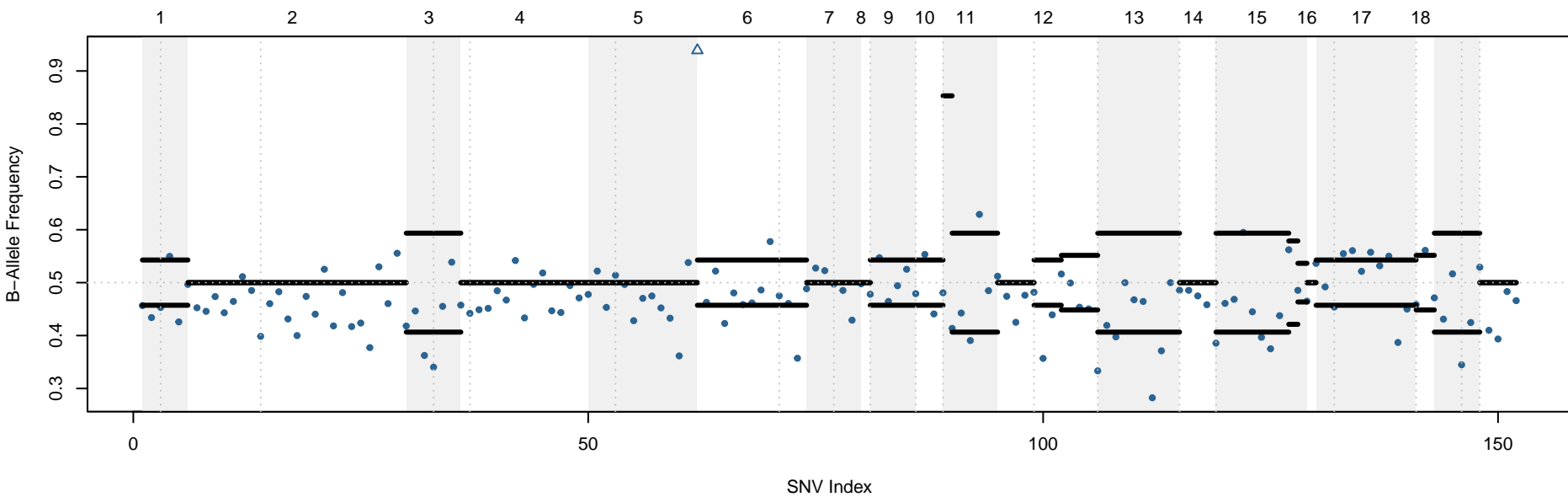


Purity: 0.23 Tumor ploidy: 4.344

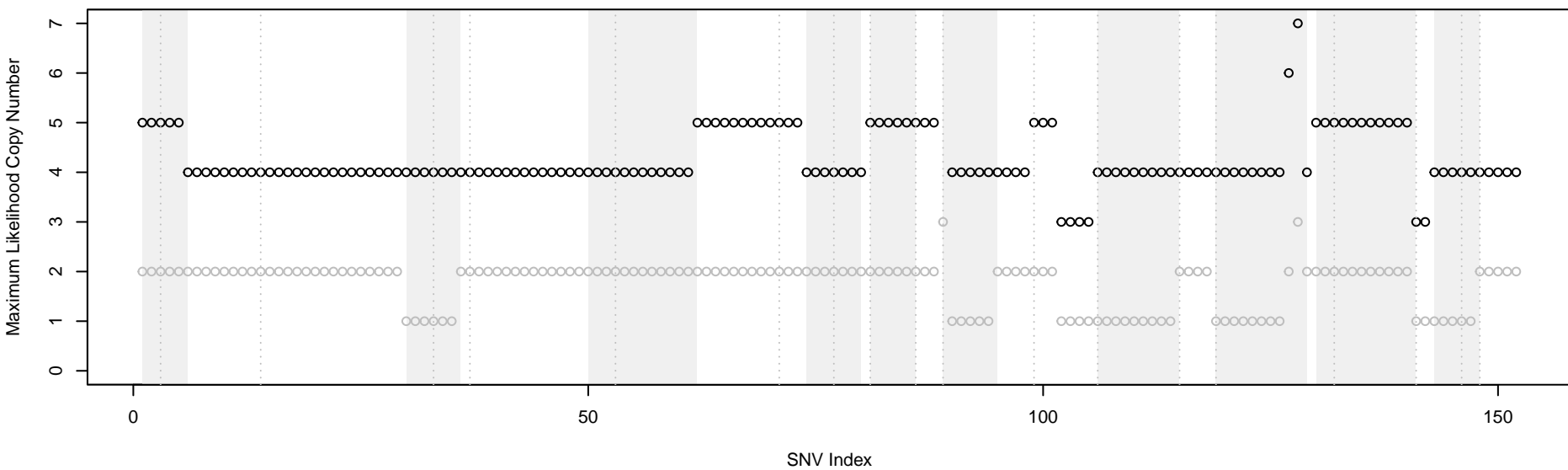
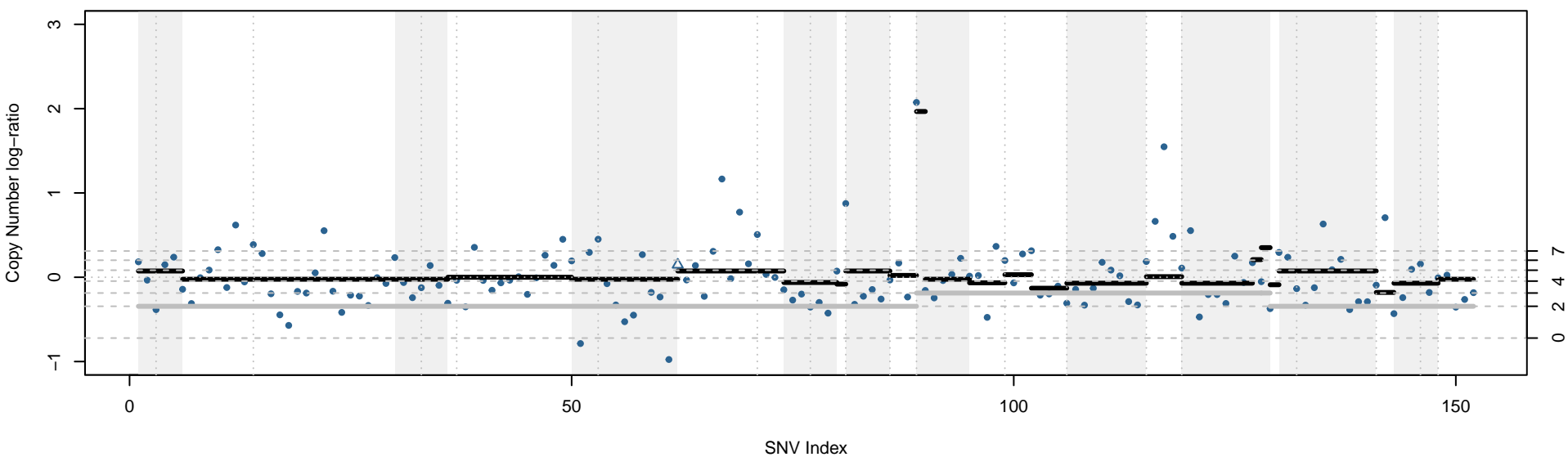
0 2 3 4 5 6 7

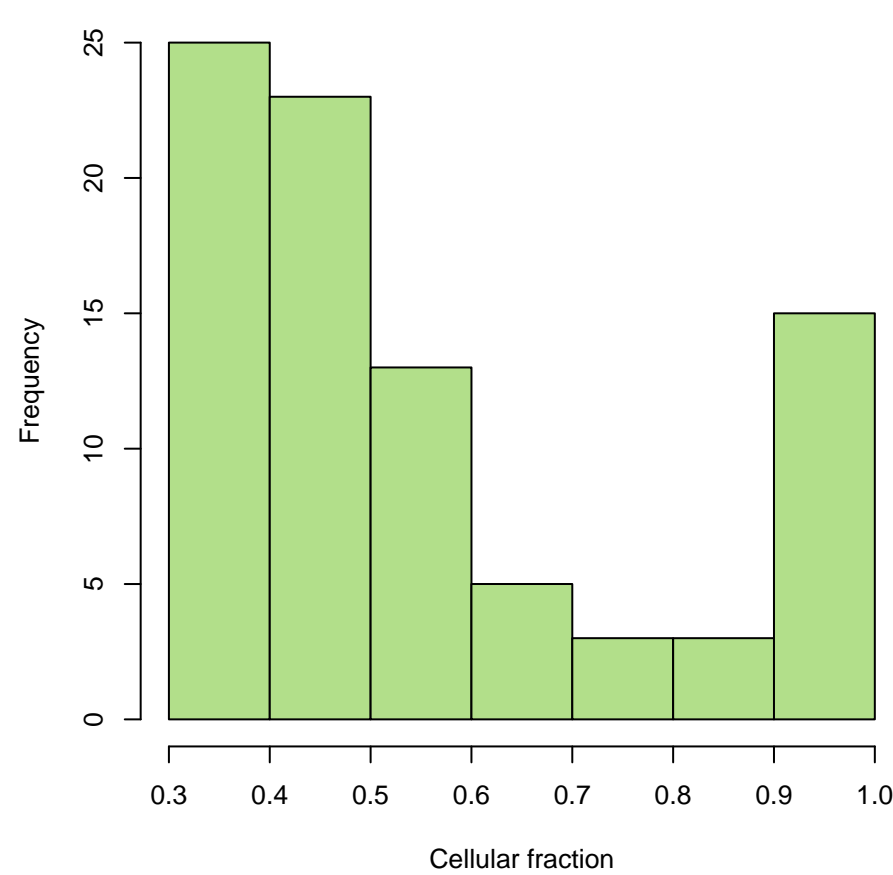
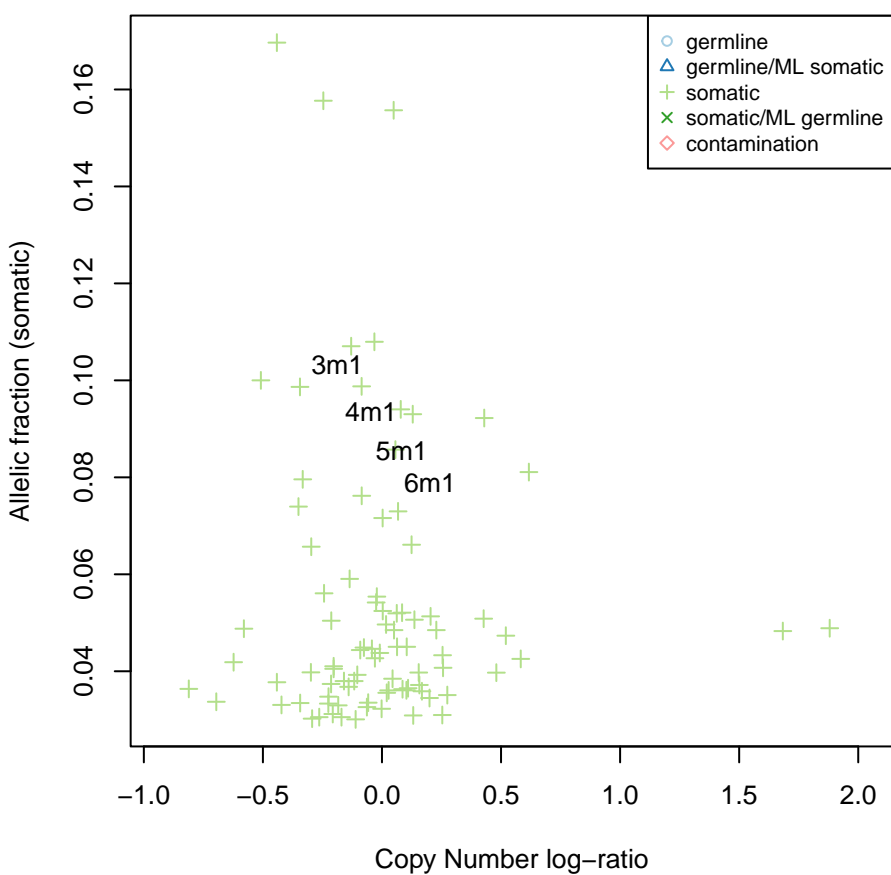
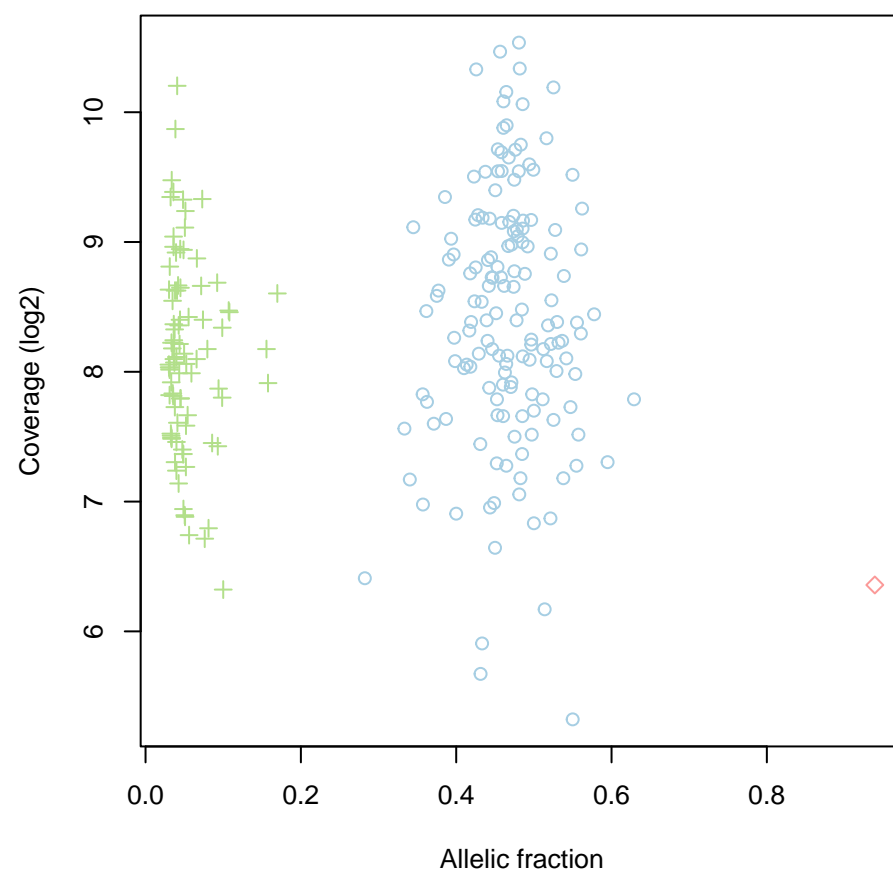
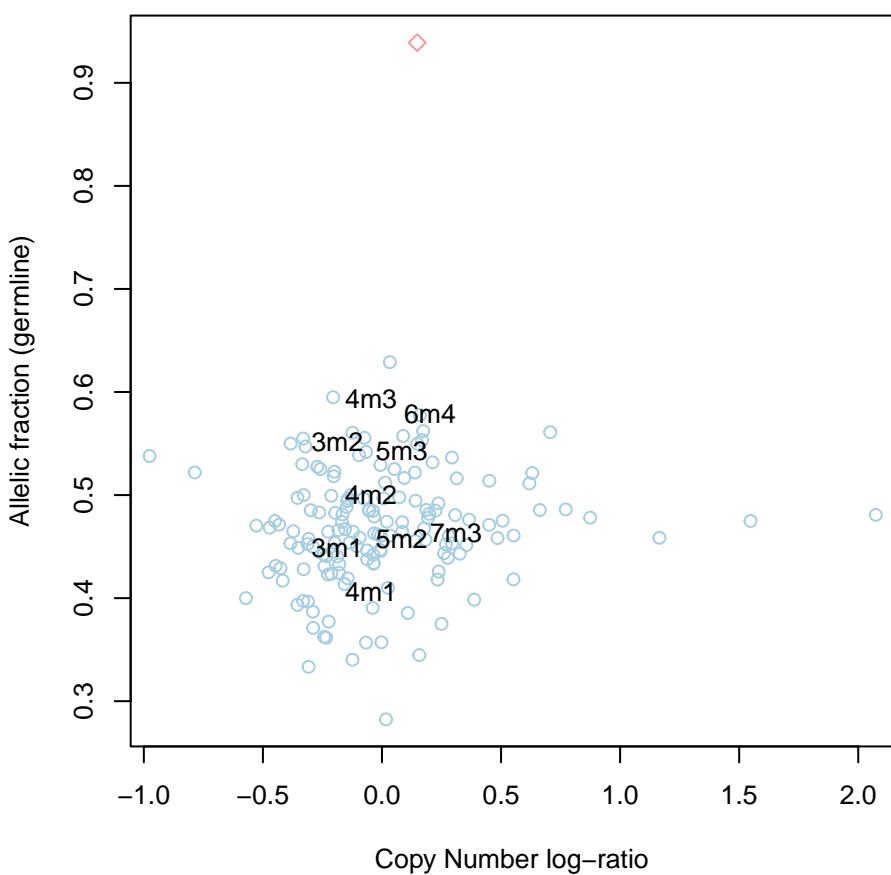


Purity: 0.23 Tumor ploidy: 4.344 SNV log-likelihood: -126.63 GoF: 94.6% Mean coverage: 108,387

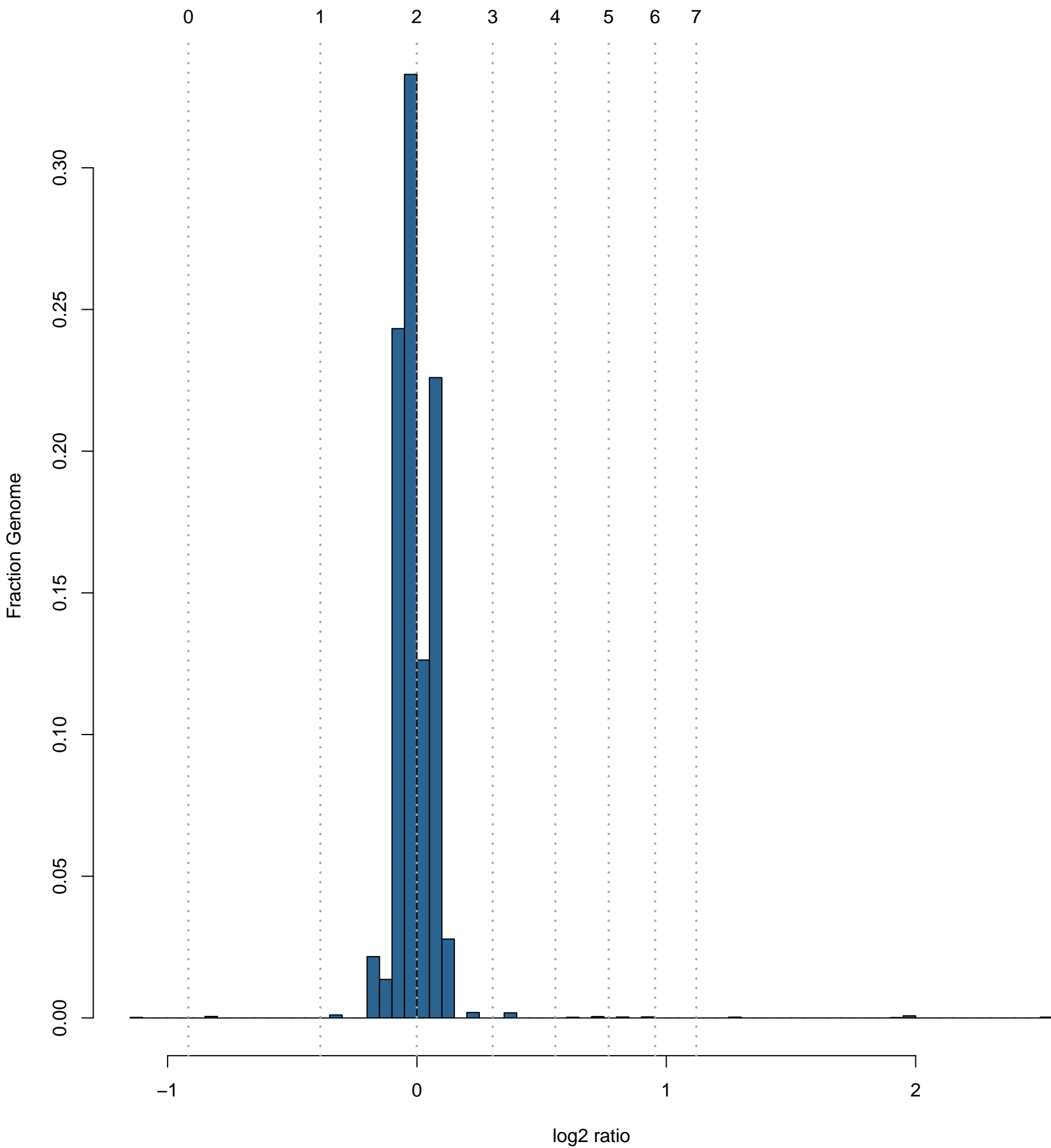


SCNA-fit log-likelihood: -15709.82

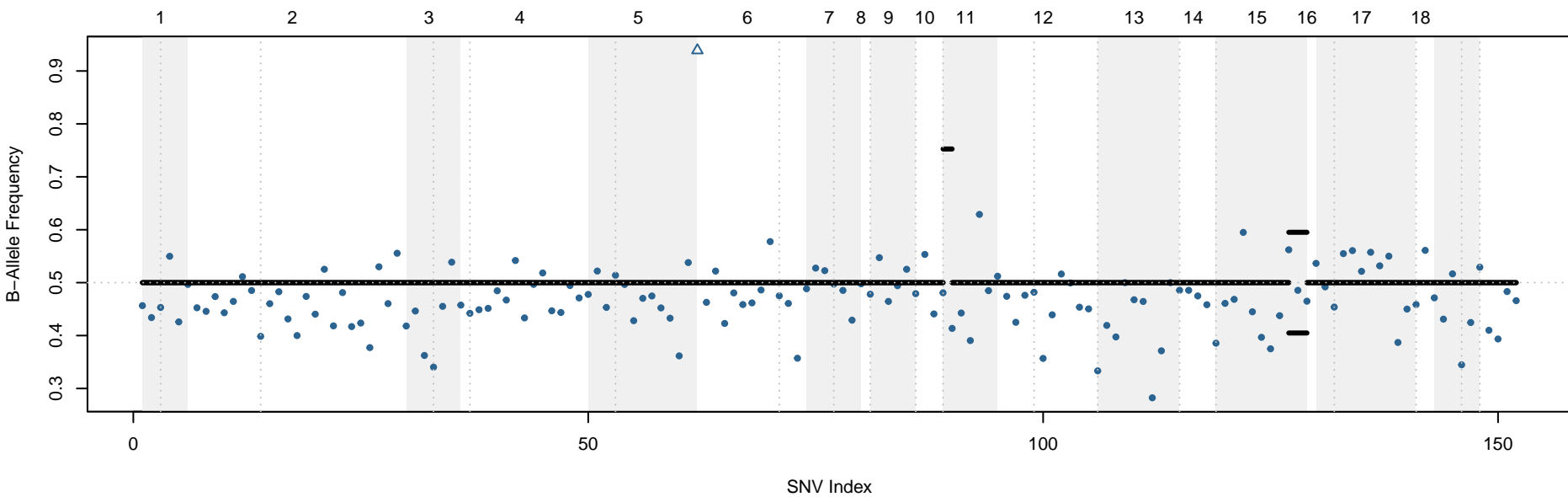




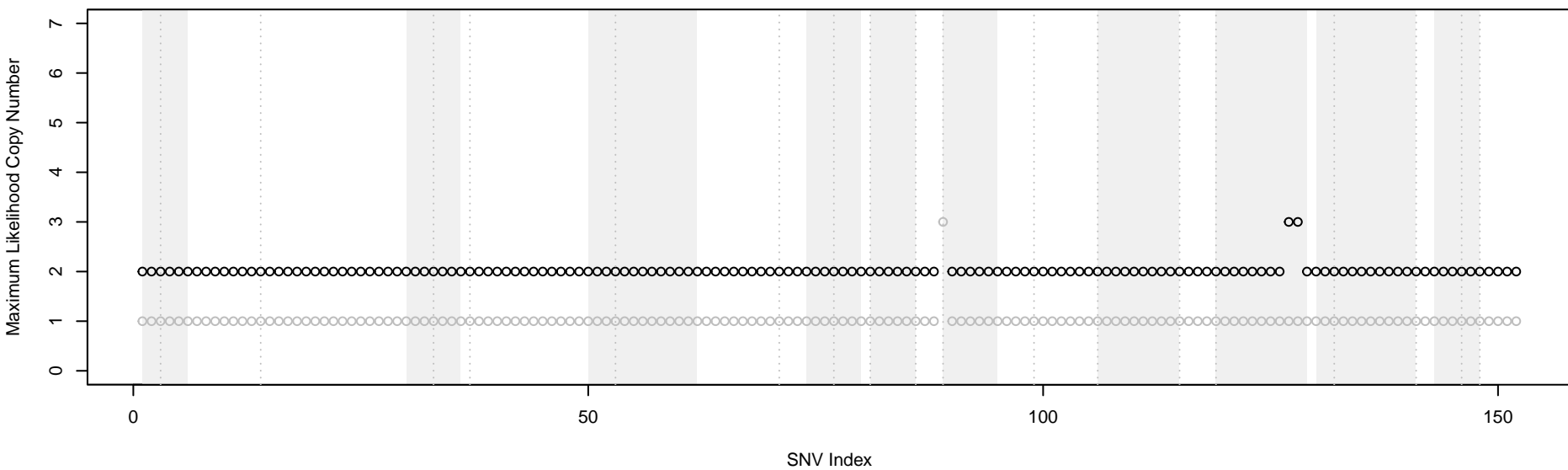
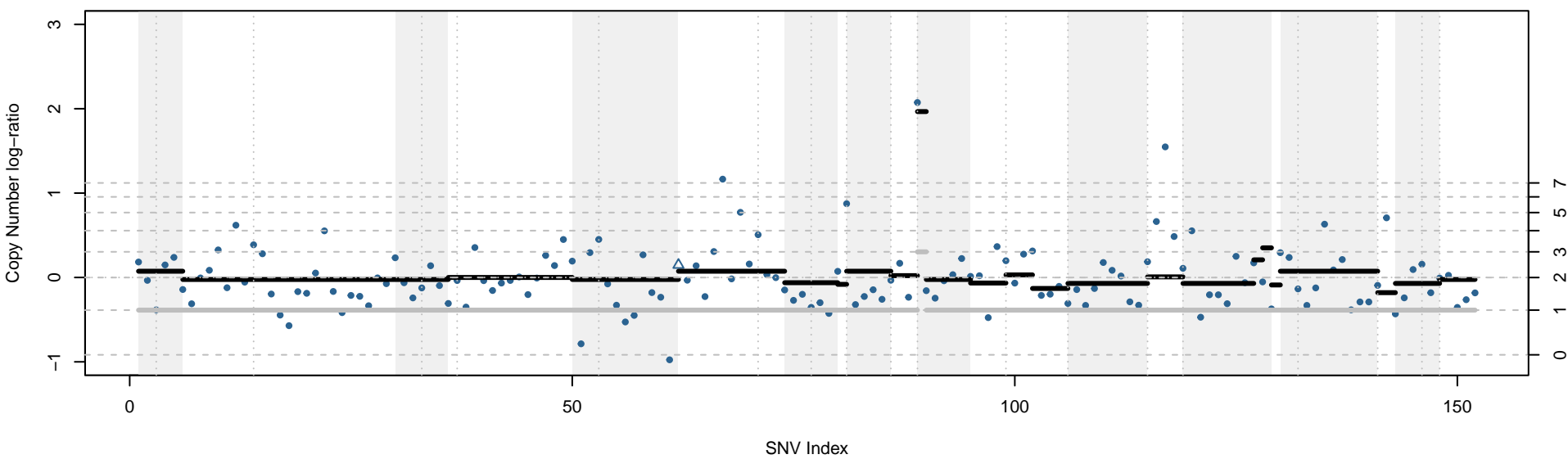
Purity: 0.47 Tumor ploidy: 2.002

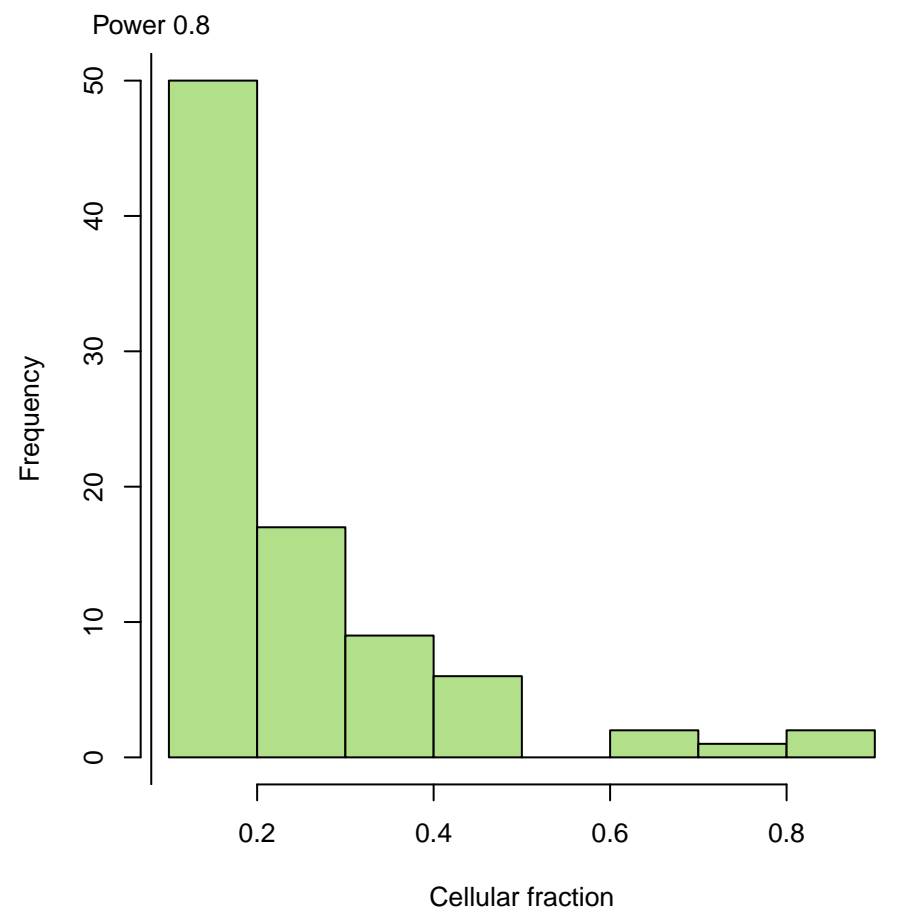
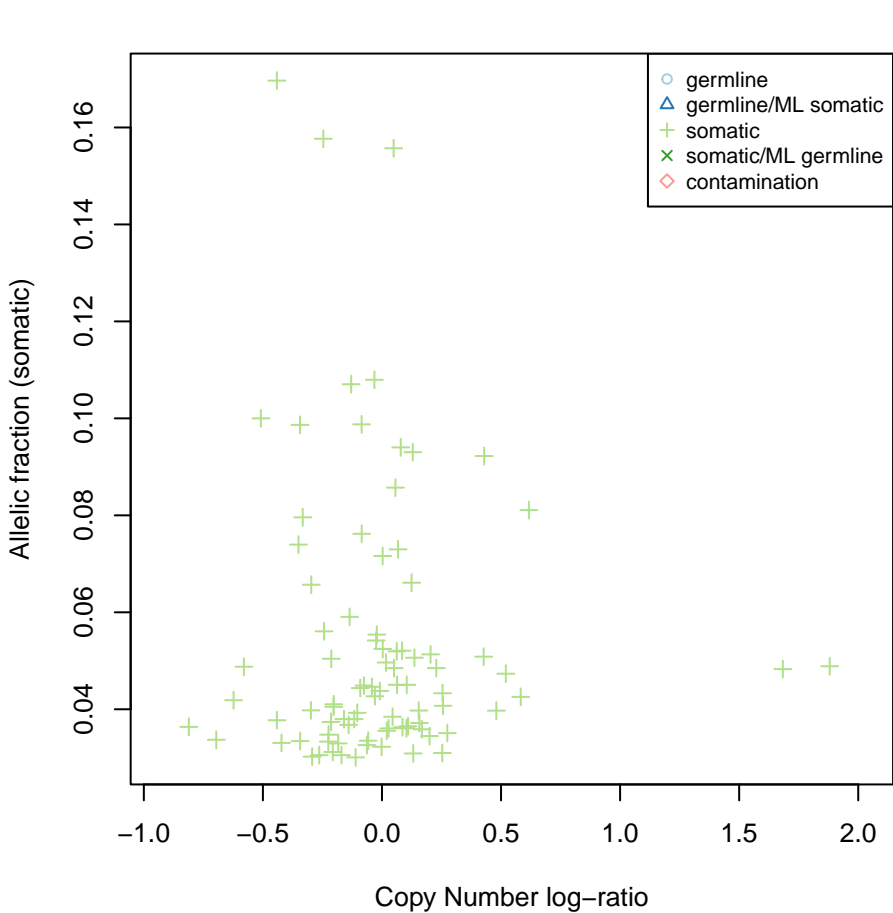
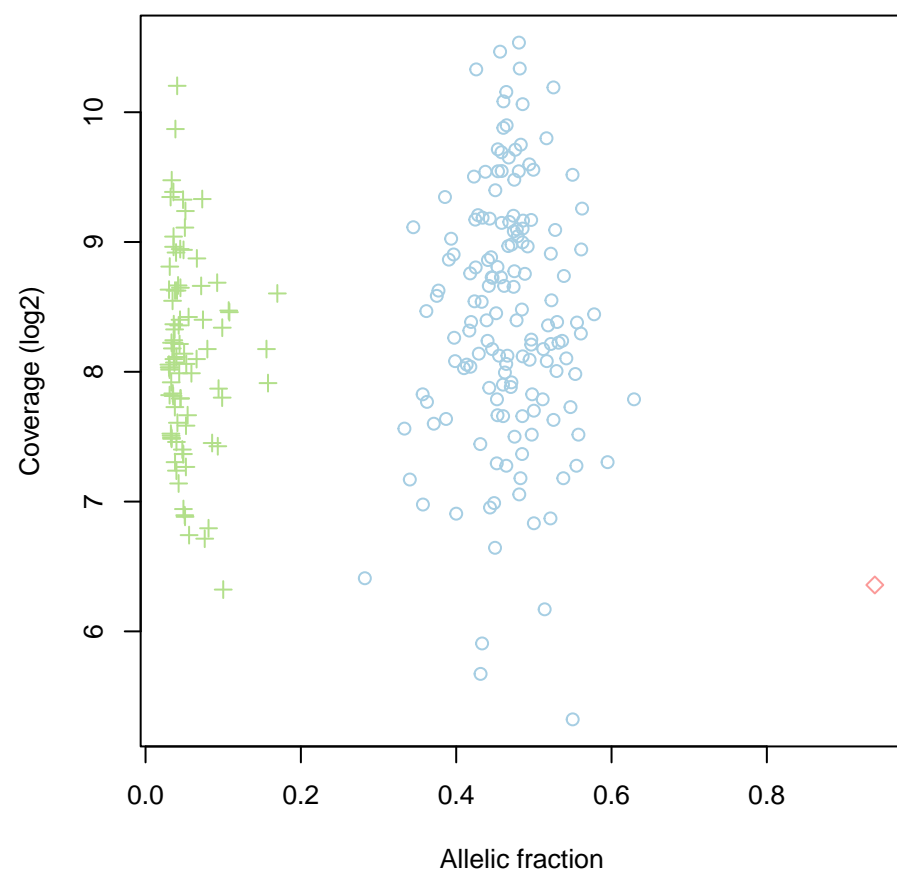
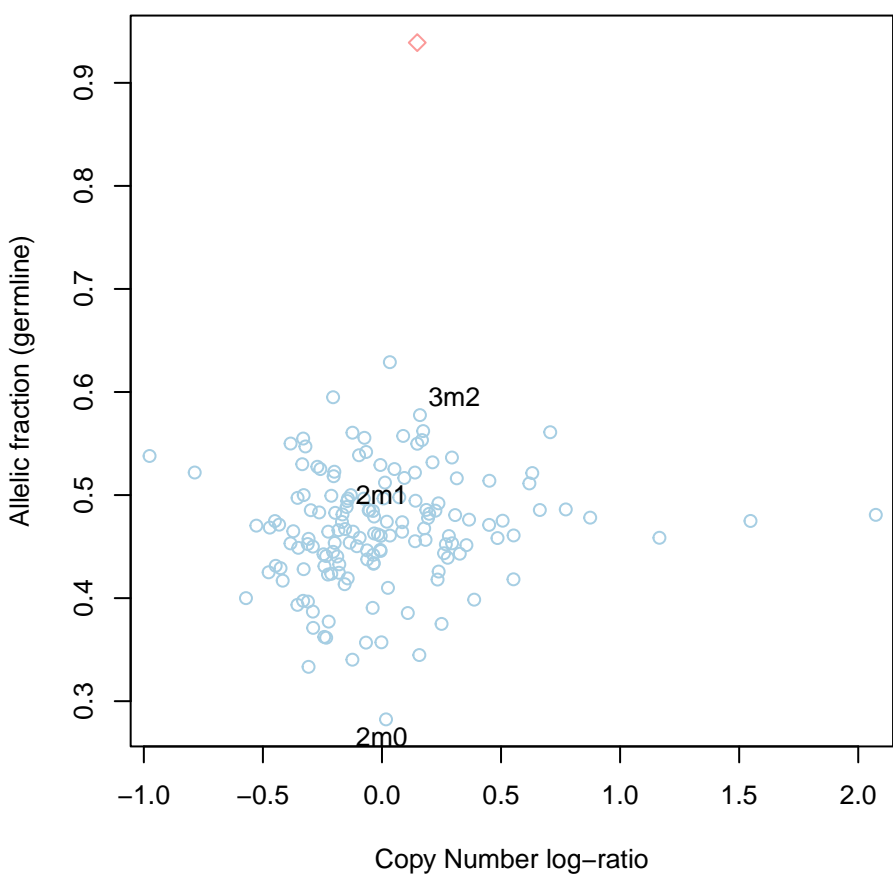


Purity: 0.47 Tumor ploidy: 2.002 SNV log-likelihood: -141.56 GoF: 65.3% Mean coverage: 108,387

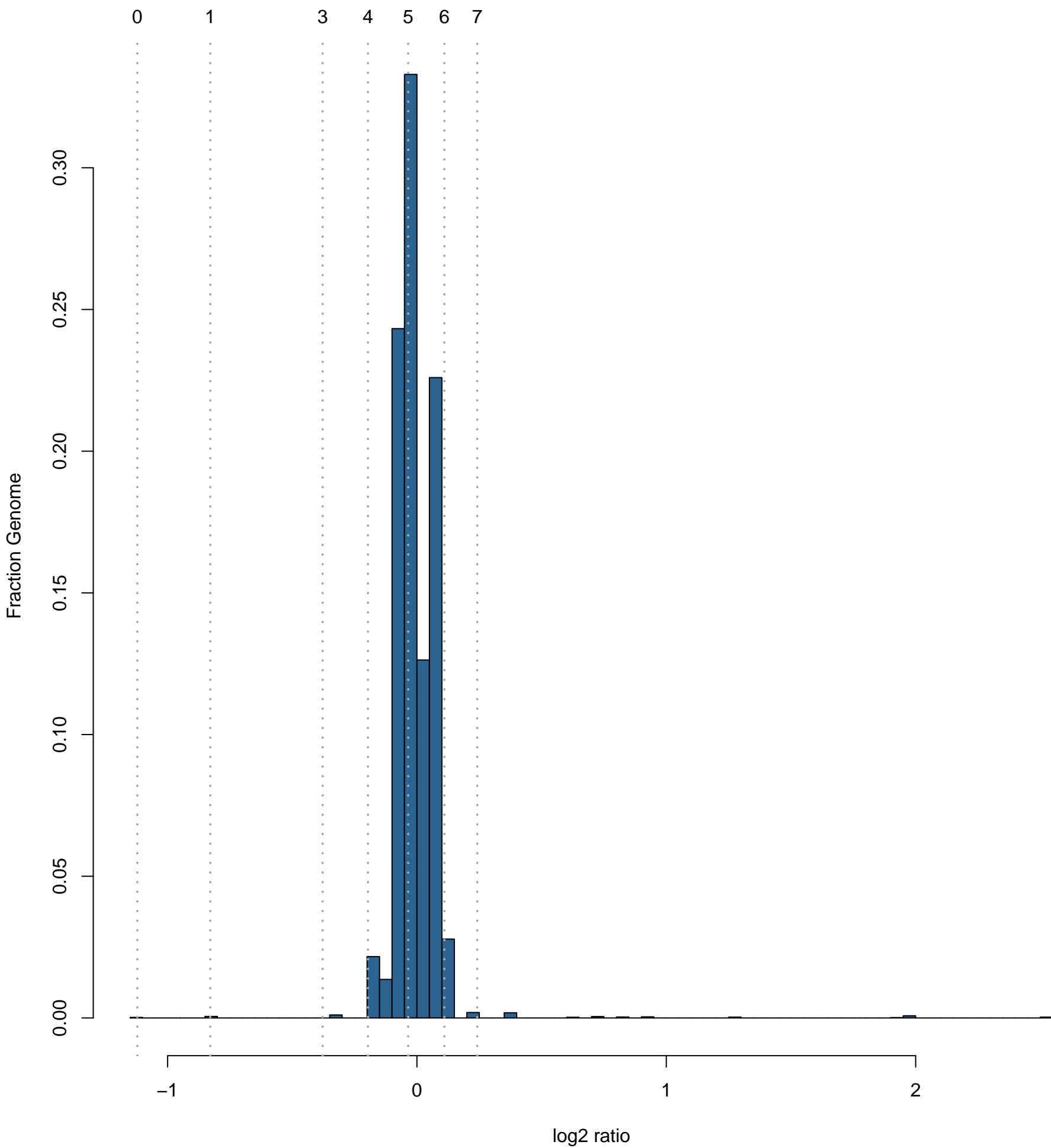


SCNA-fit log-likelihood: -15809.57

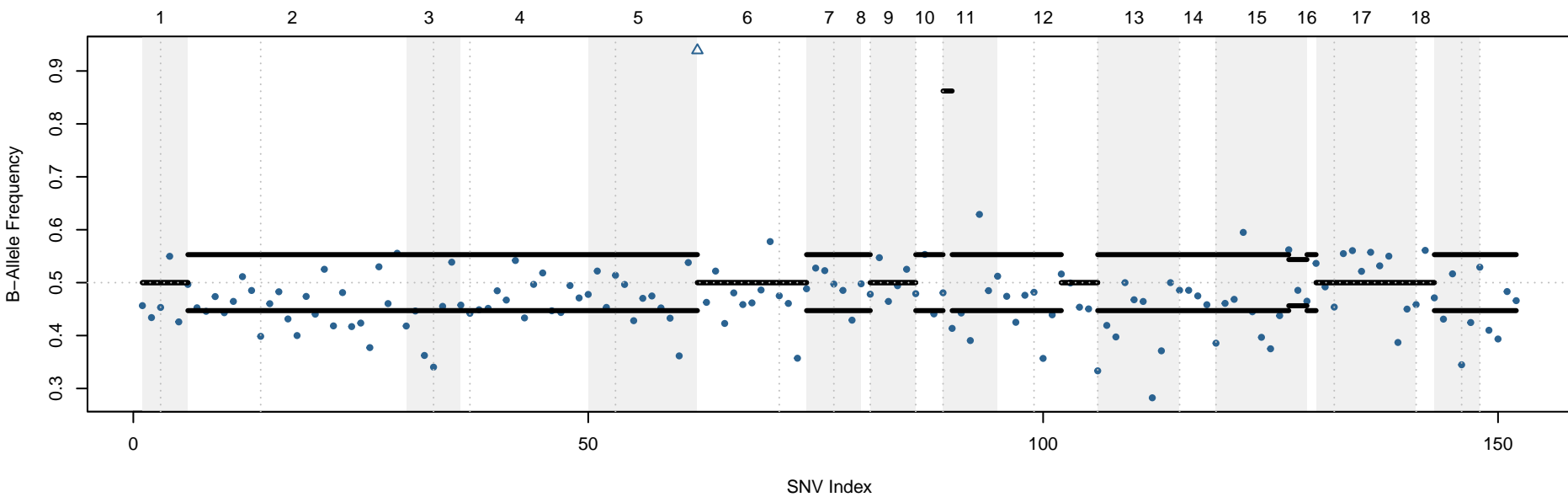




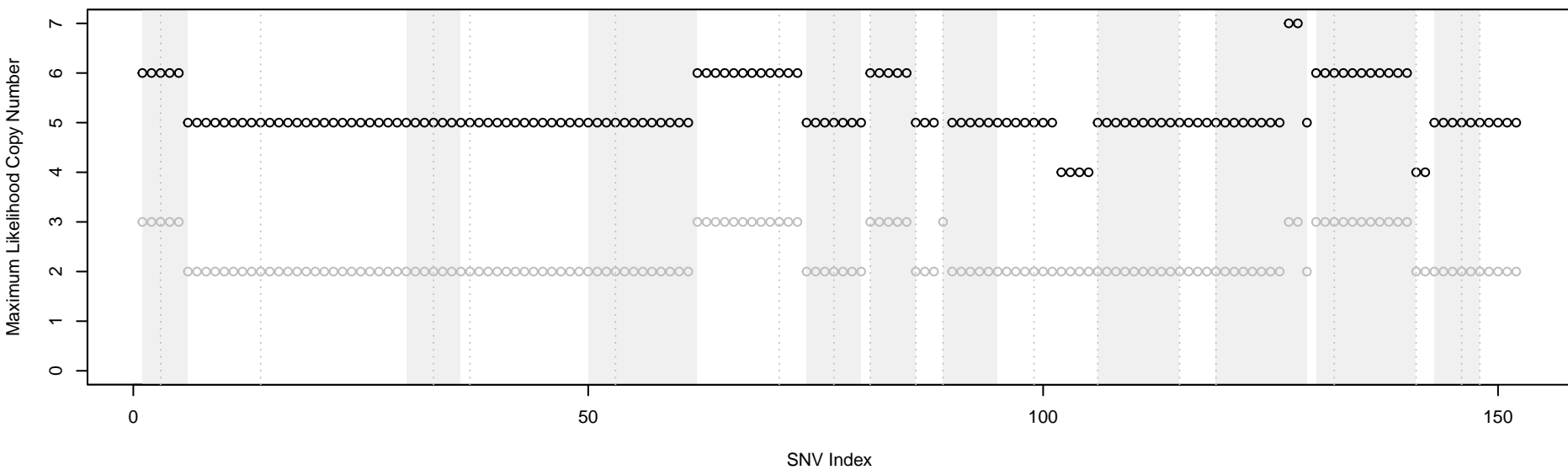
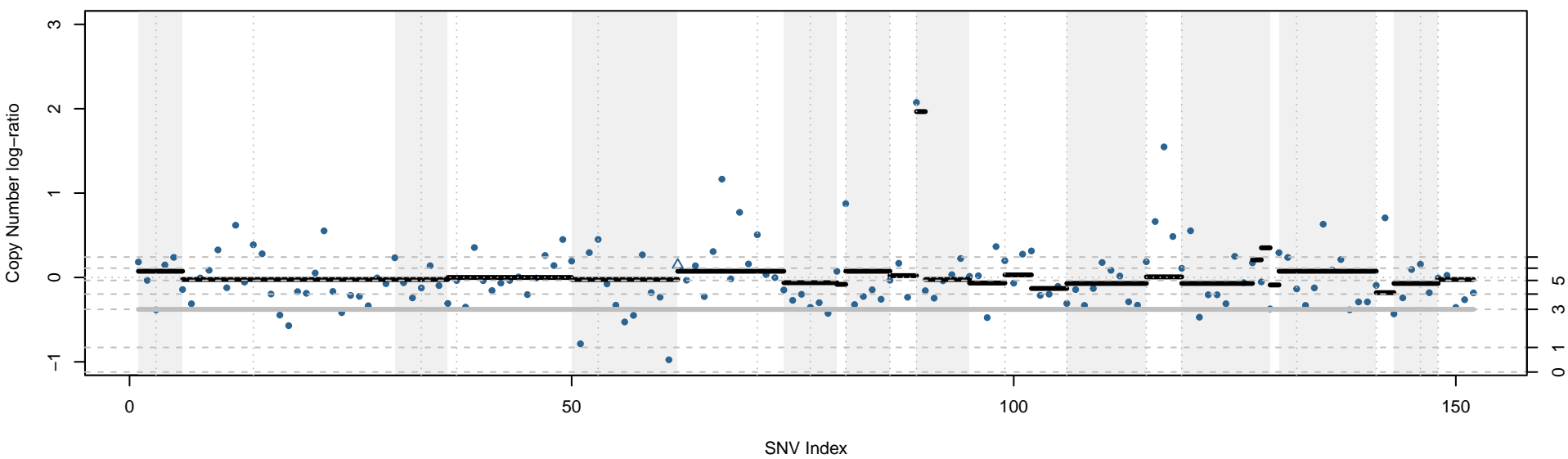
Purity: 0.31 Tumor ploidy: 5.233

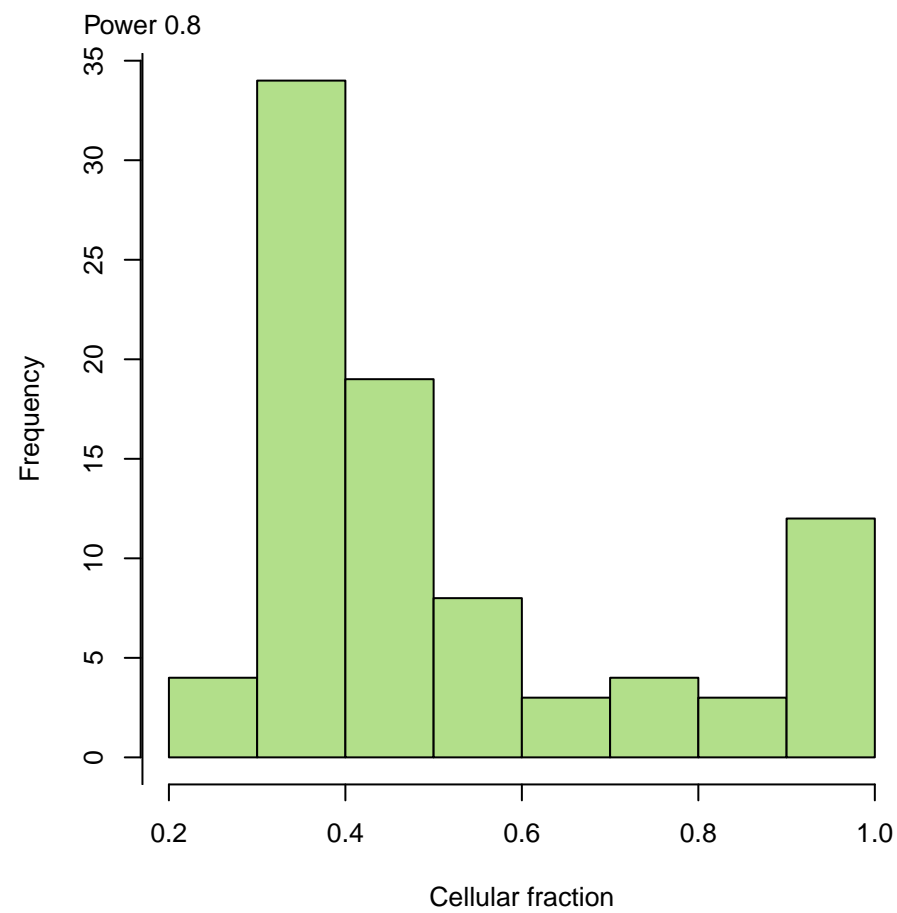
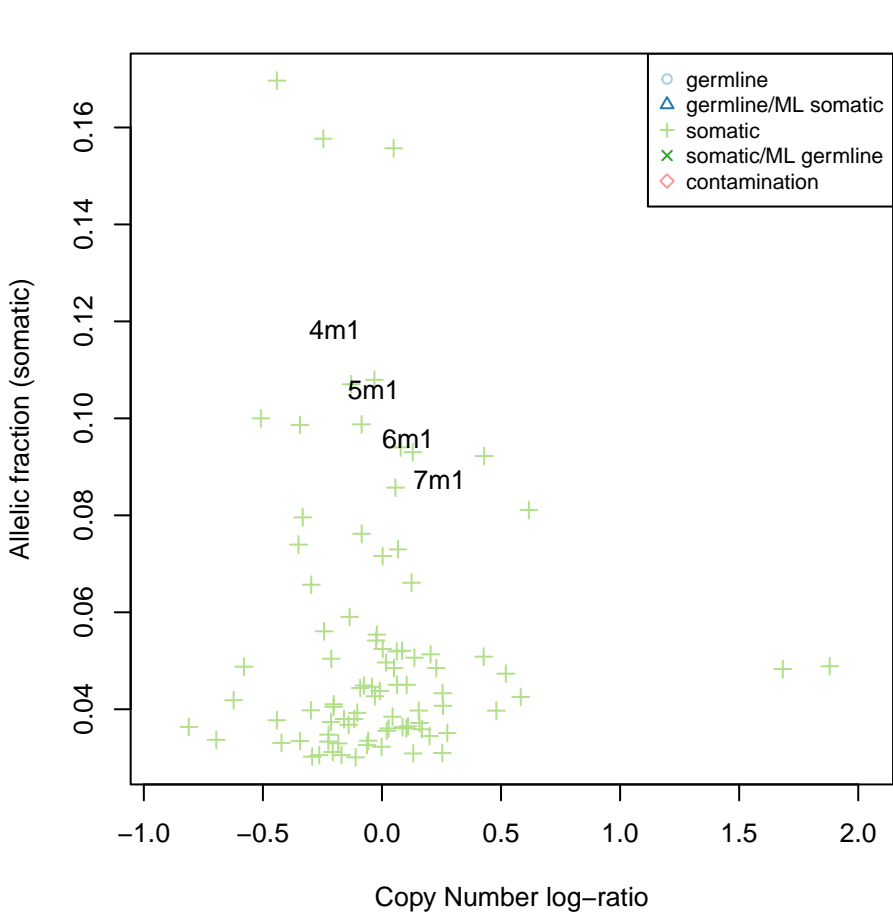
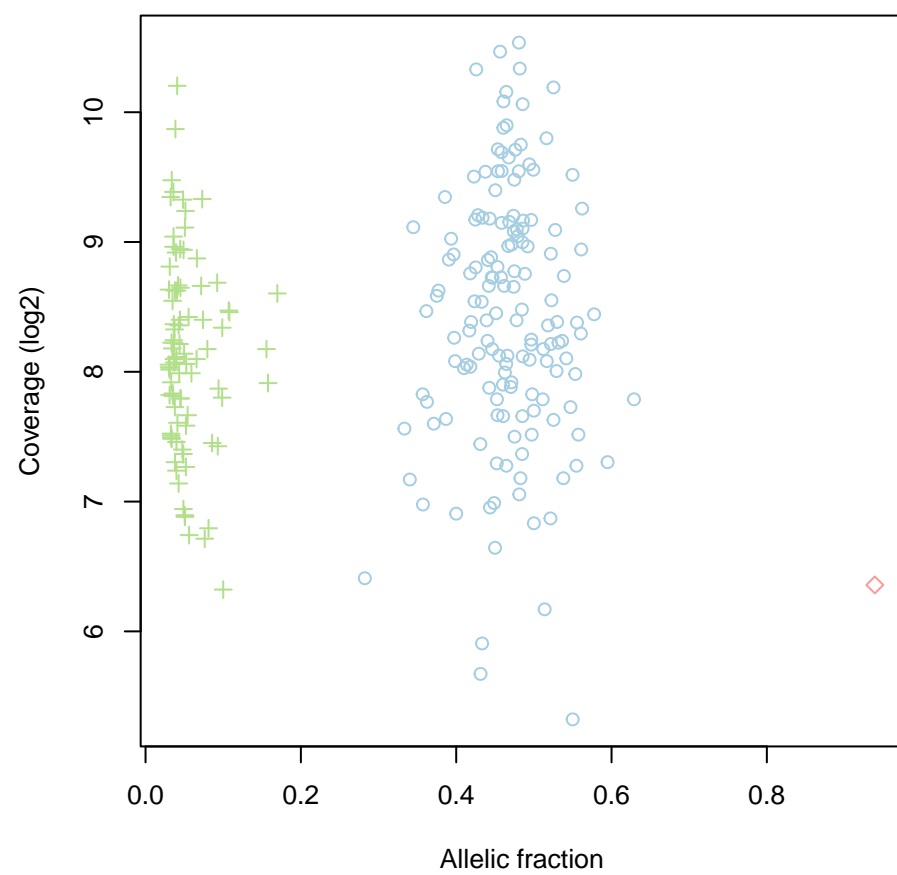
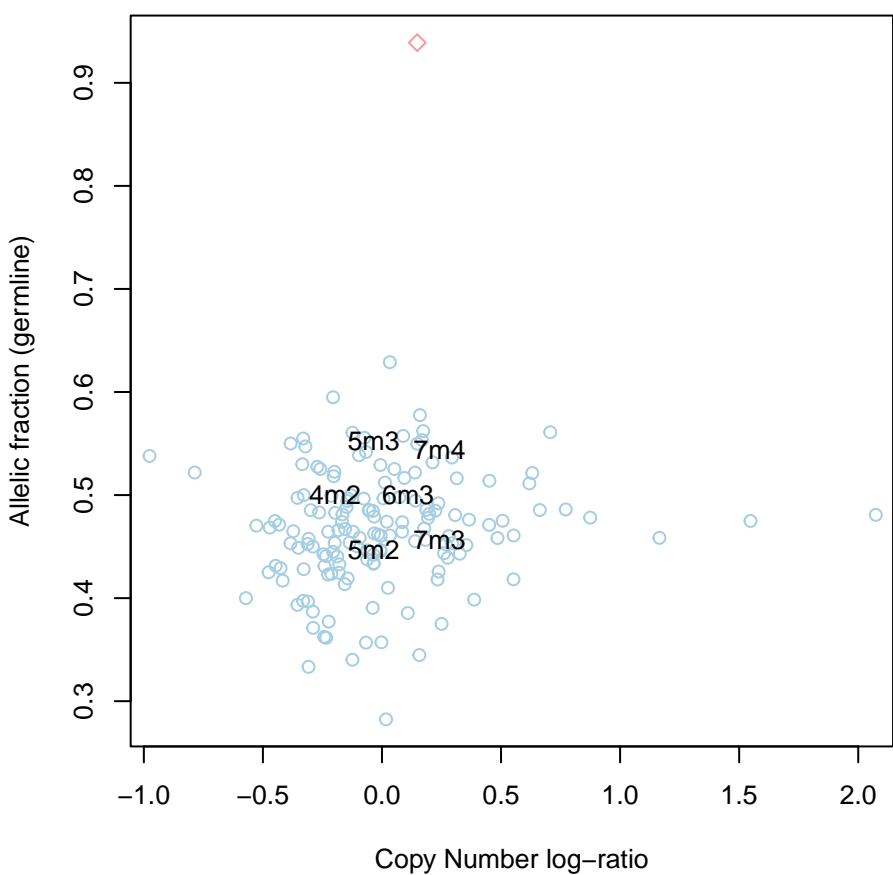


Purity: 0.31 Tumor ploidy: 5.233 SNV log-likelihood: -187.48 GoF: 93.4% Mean coverage: 108,387

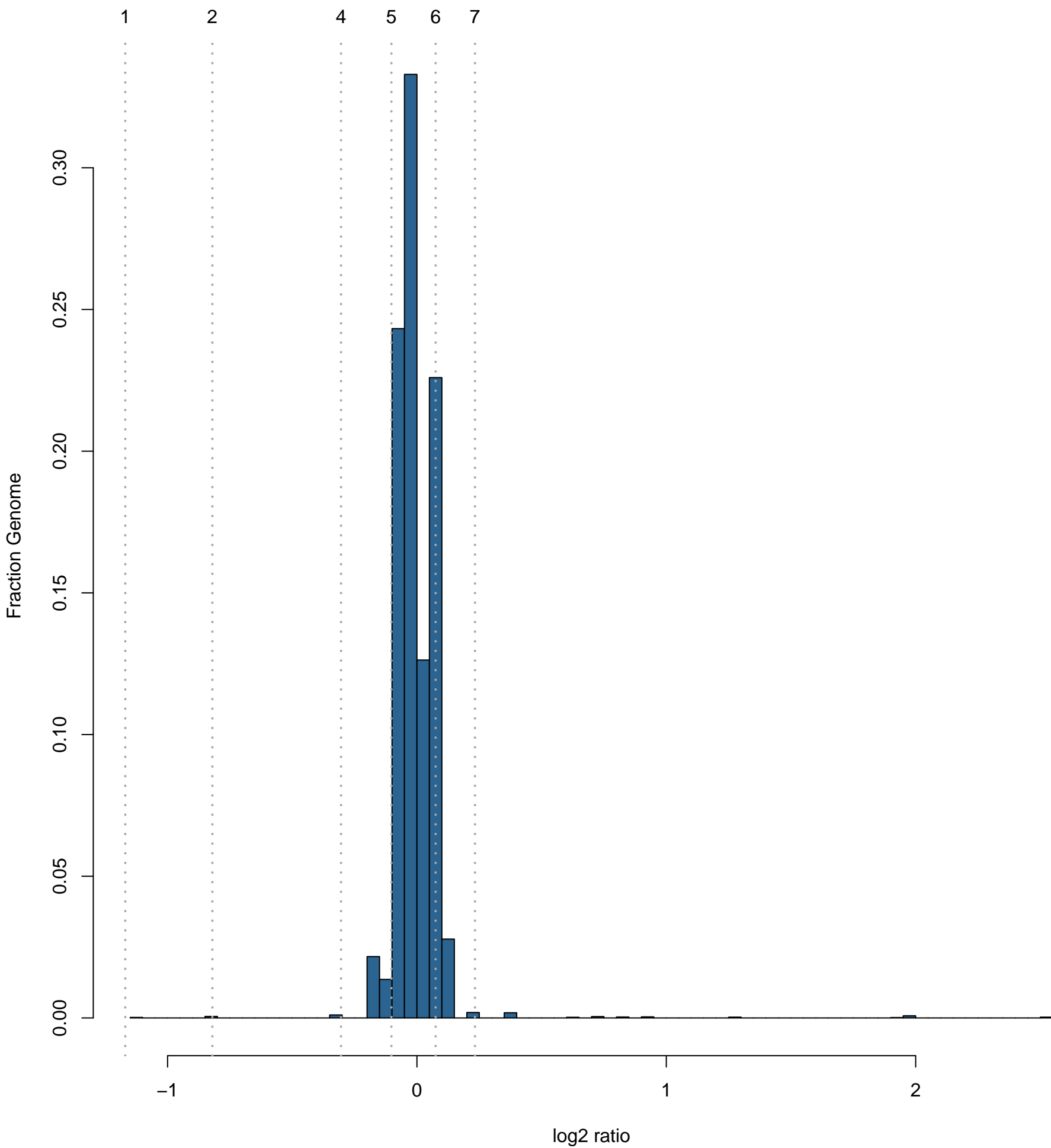


SCNA-fit log-likelihood: -15719.68

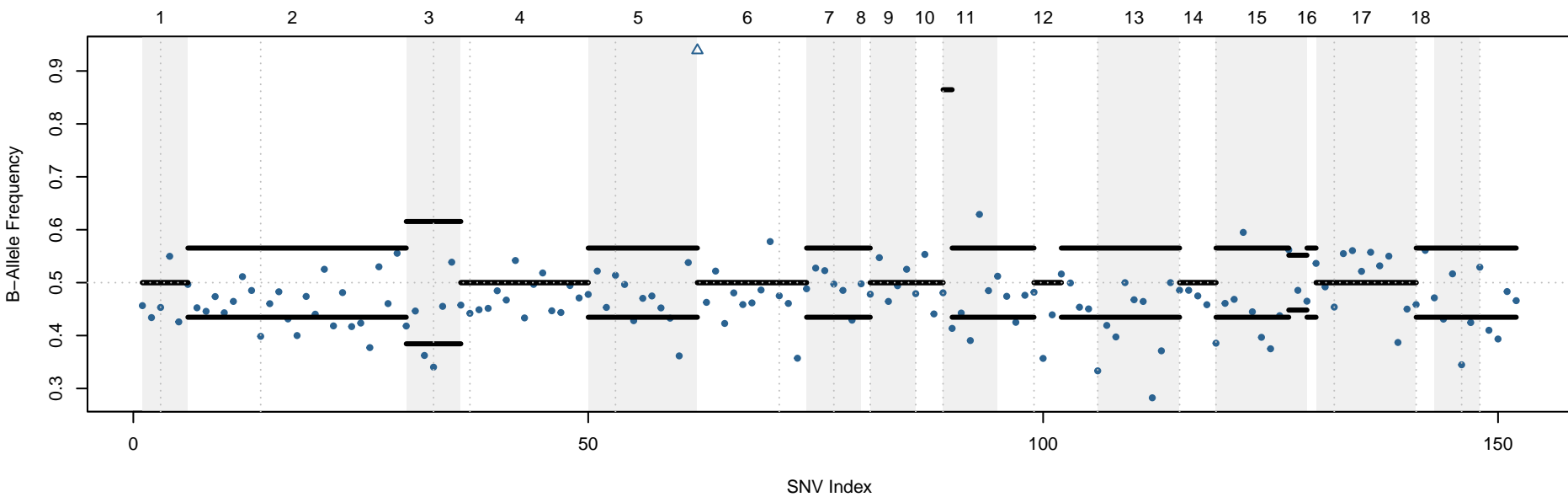




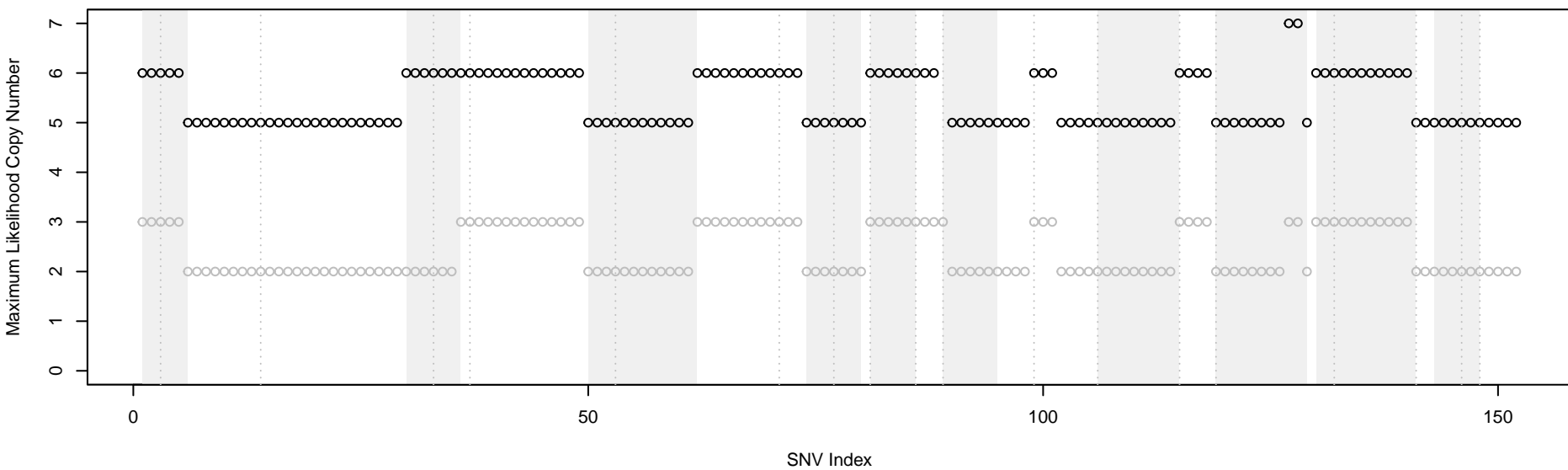
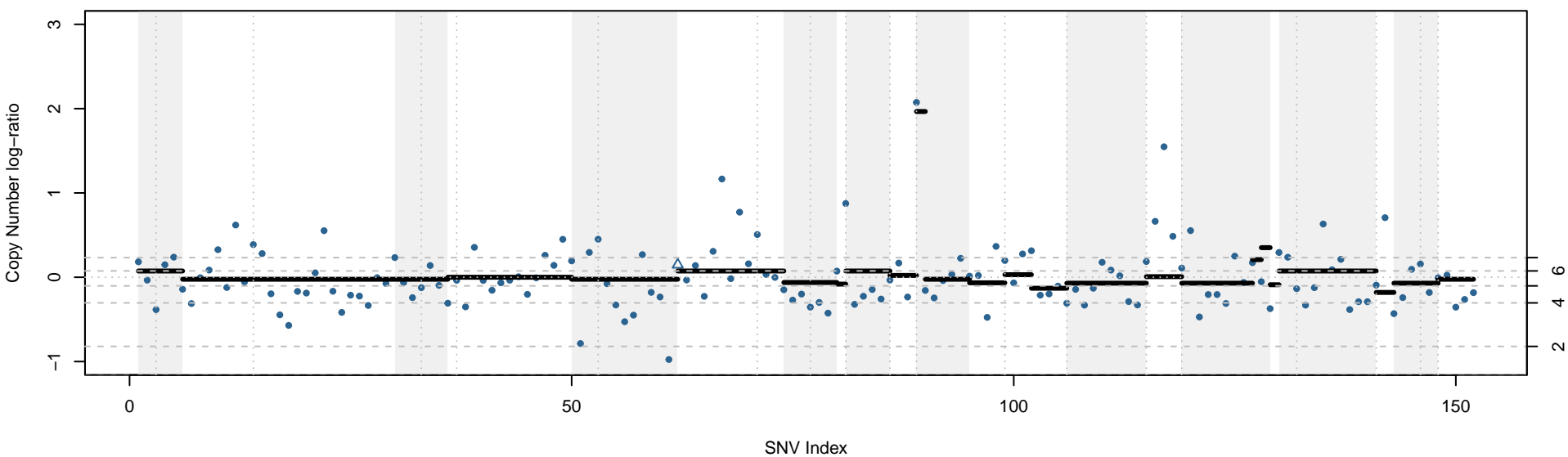
Purity: 0.43 Tumor ploidy: 5.562

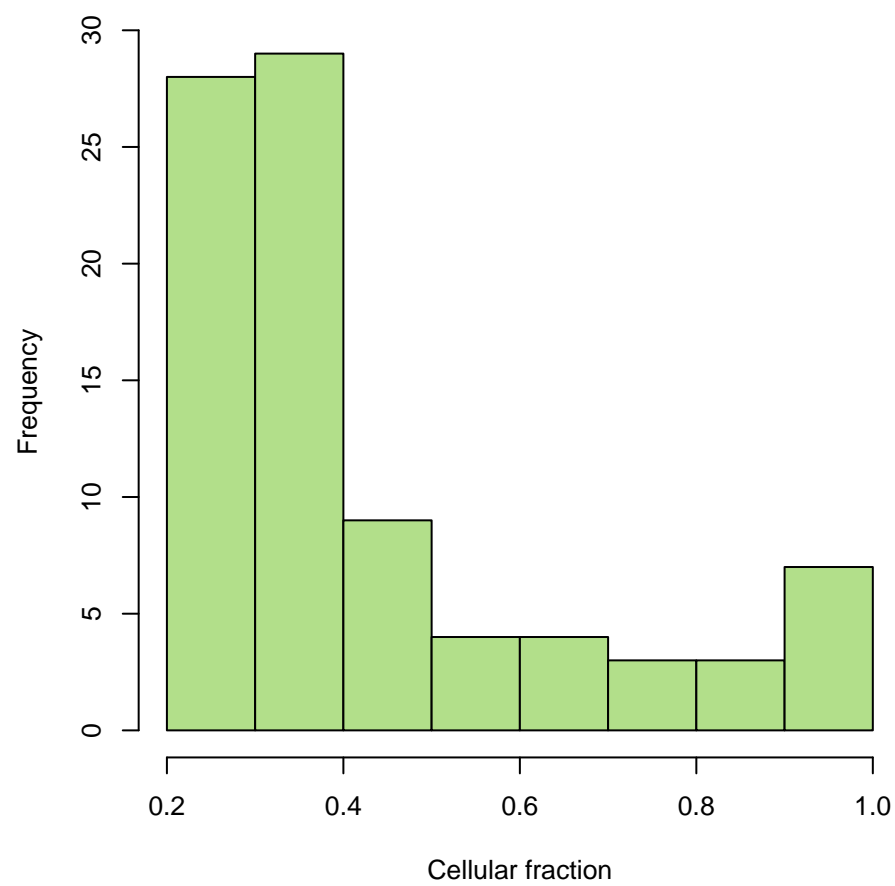
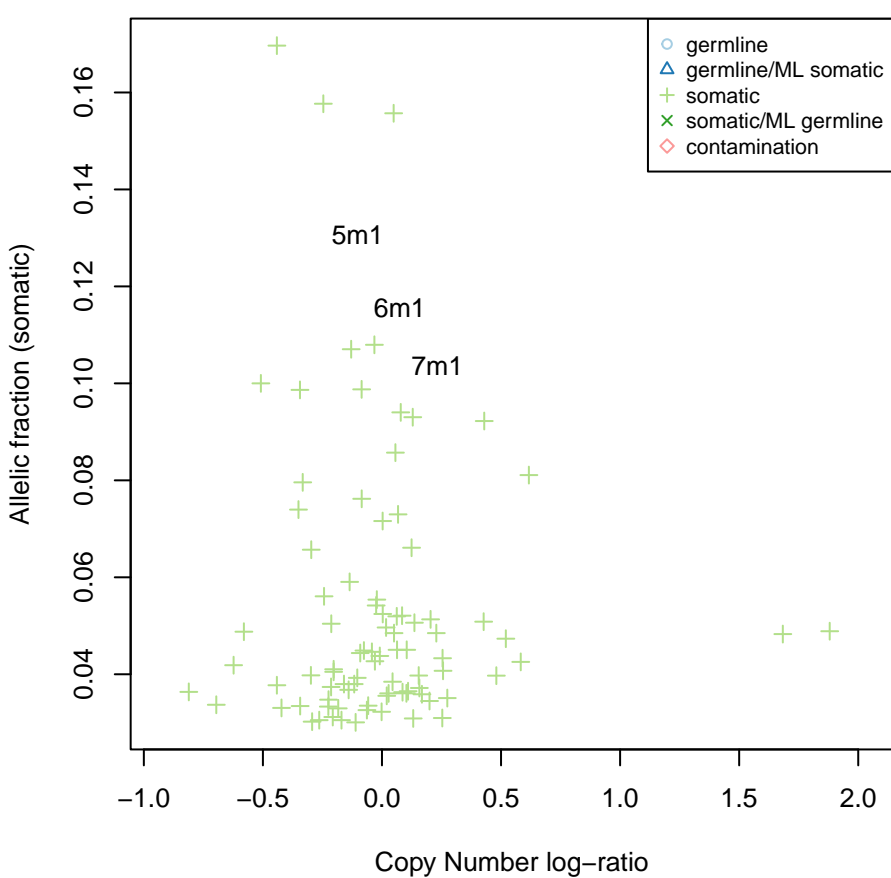
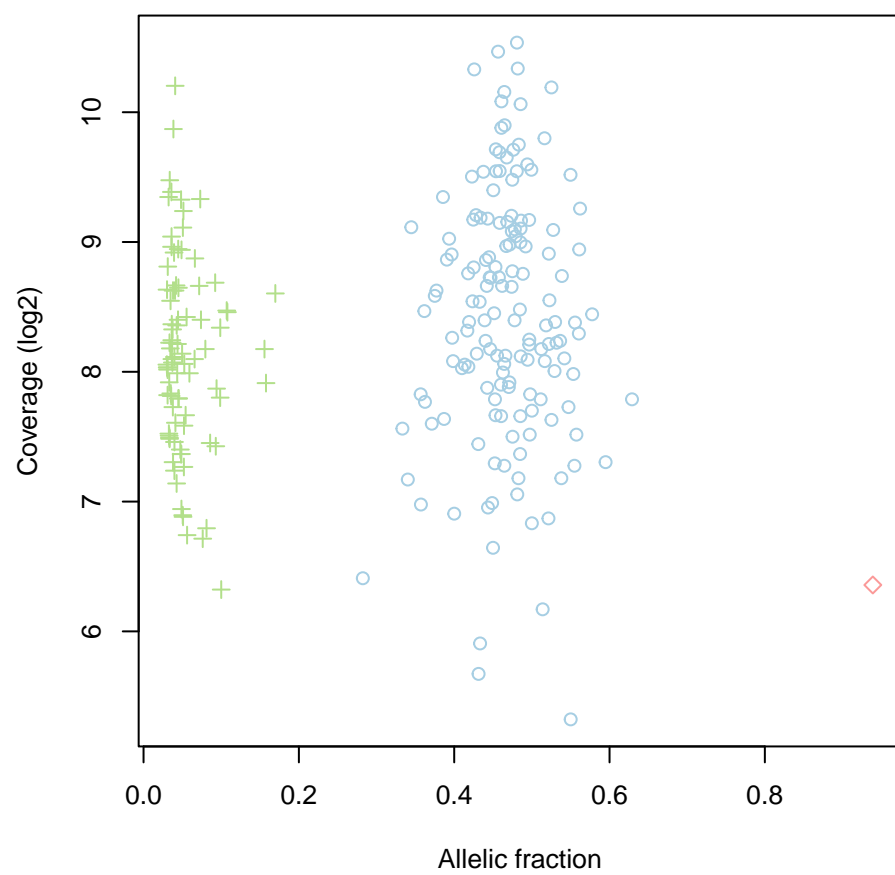
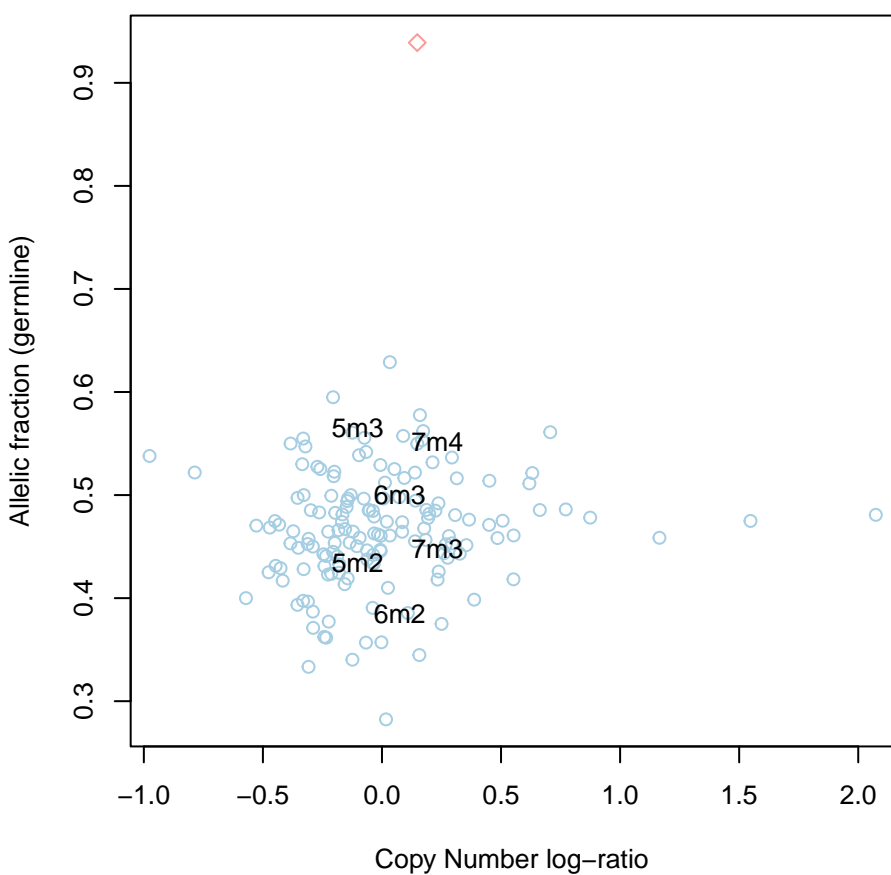


Purity: 0.43 Tumor ploidy: 5.562 SNV log-likelihood: -185.19 GoF: 90.9% Mean coverage: 108,387

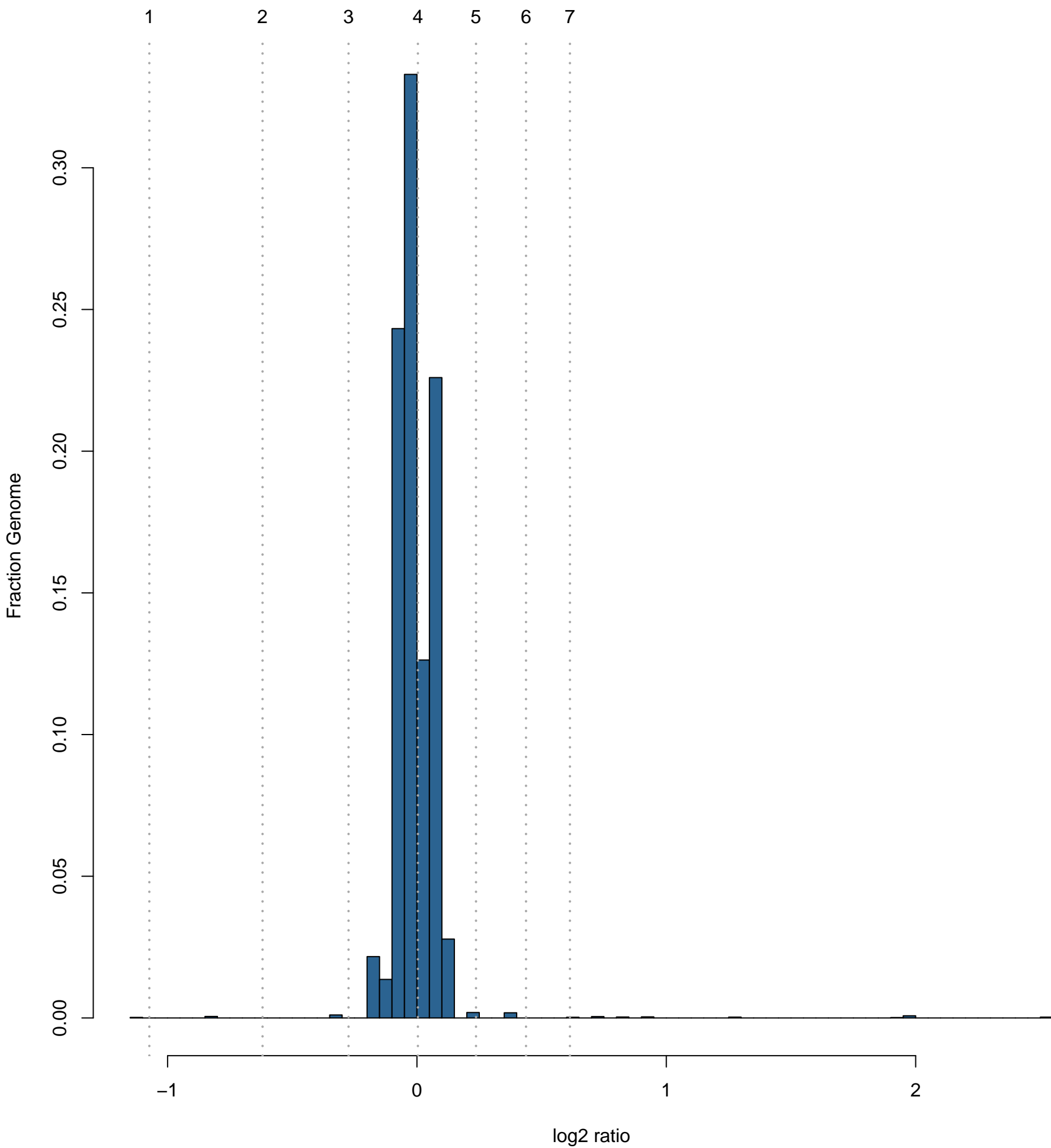


SCNA-fit log-likelihood: -15778.78

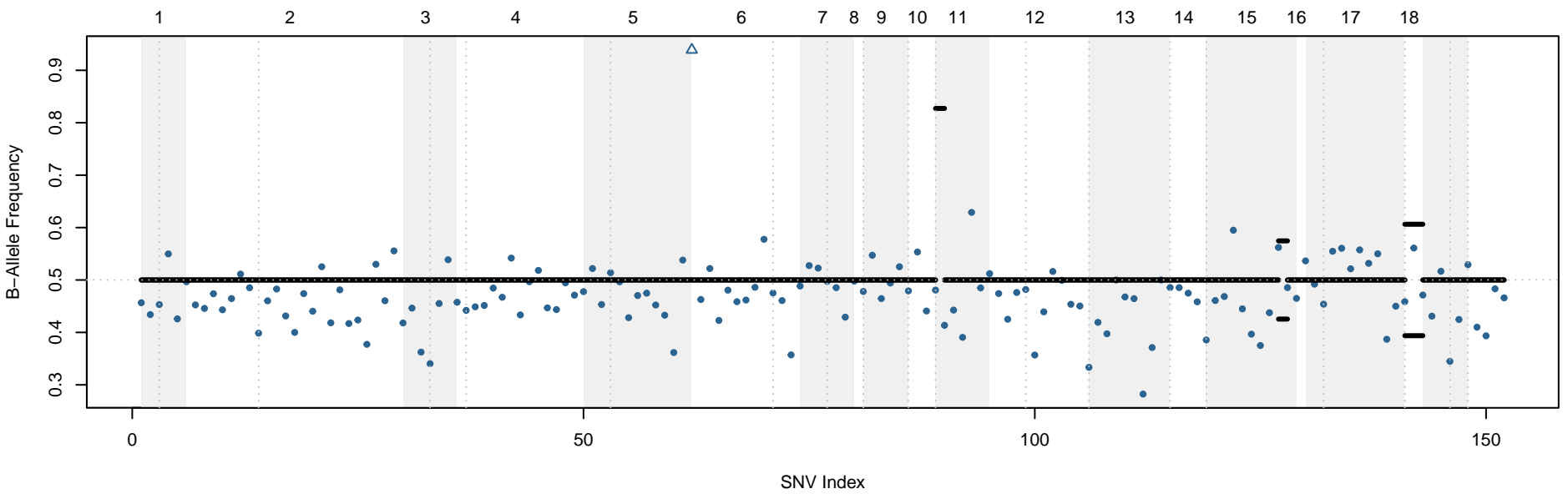




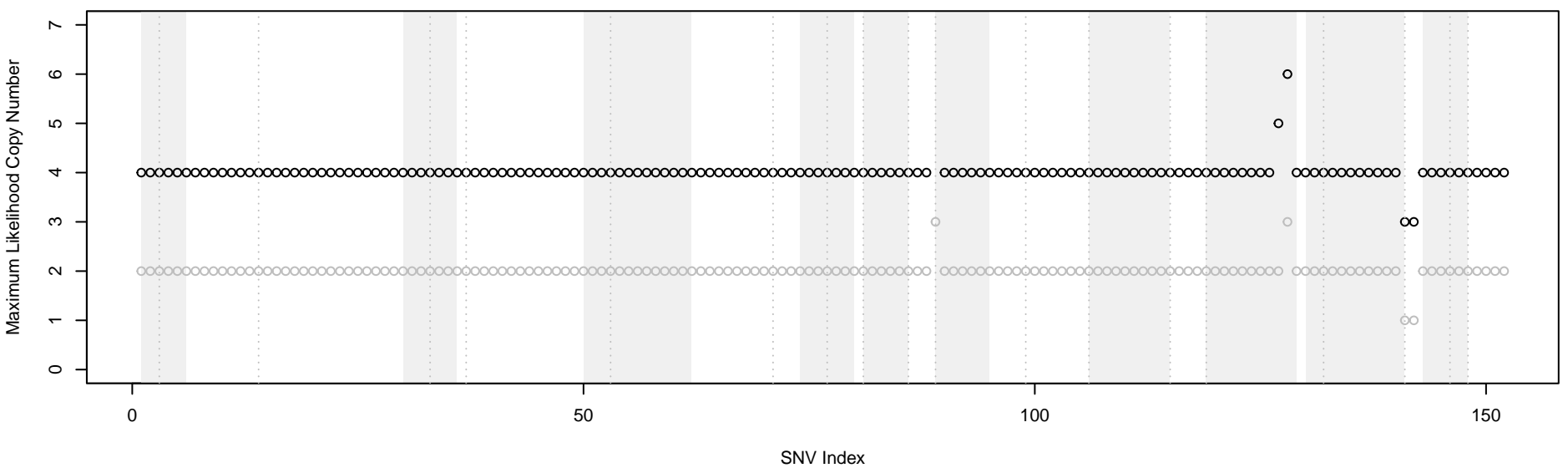
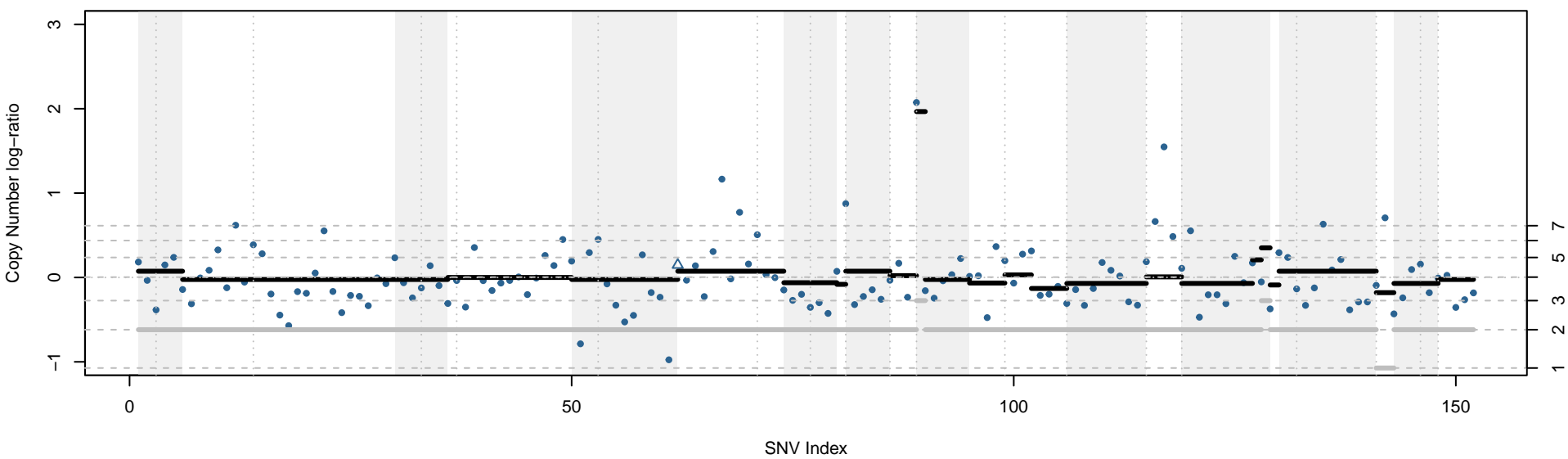
Purity: 0.54 Tumor ploidy: 3.985

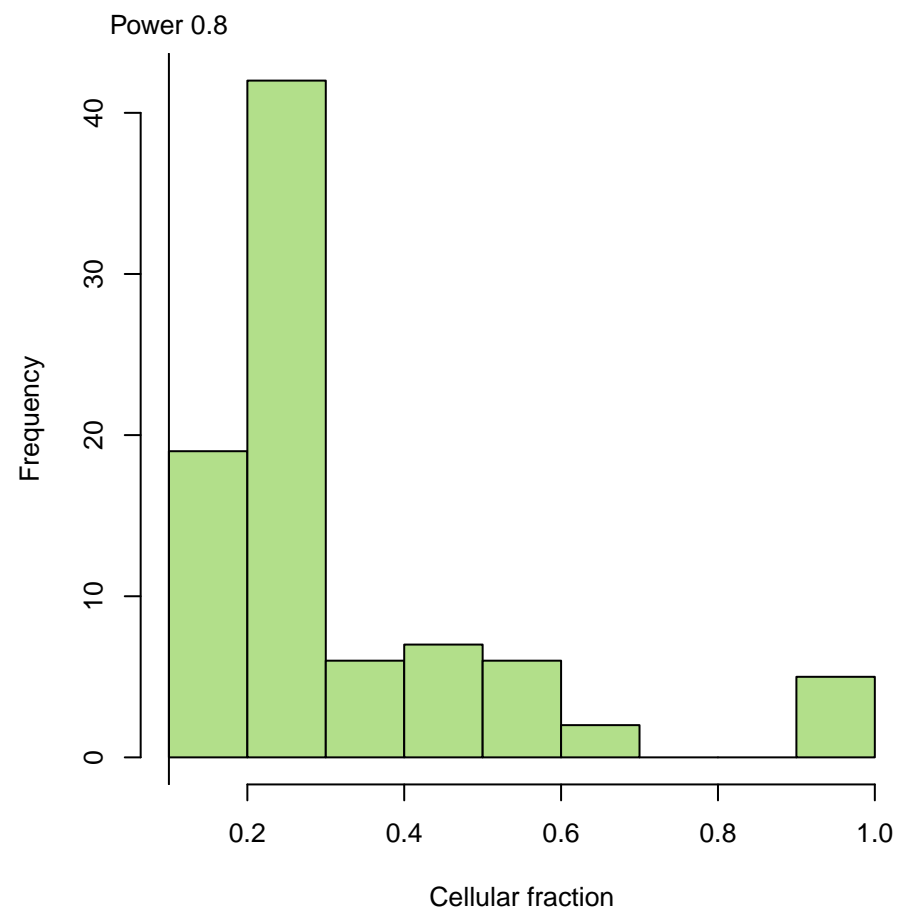
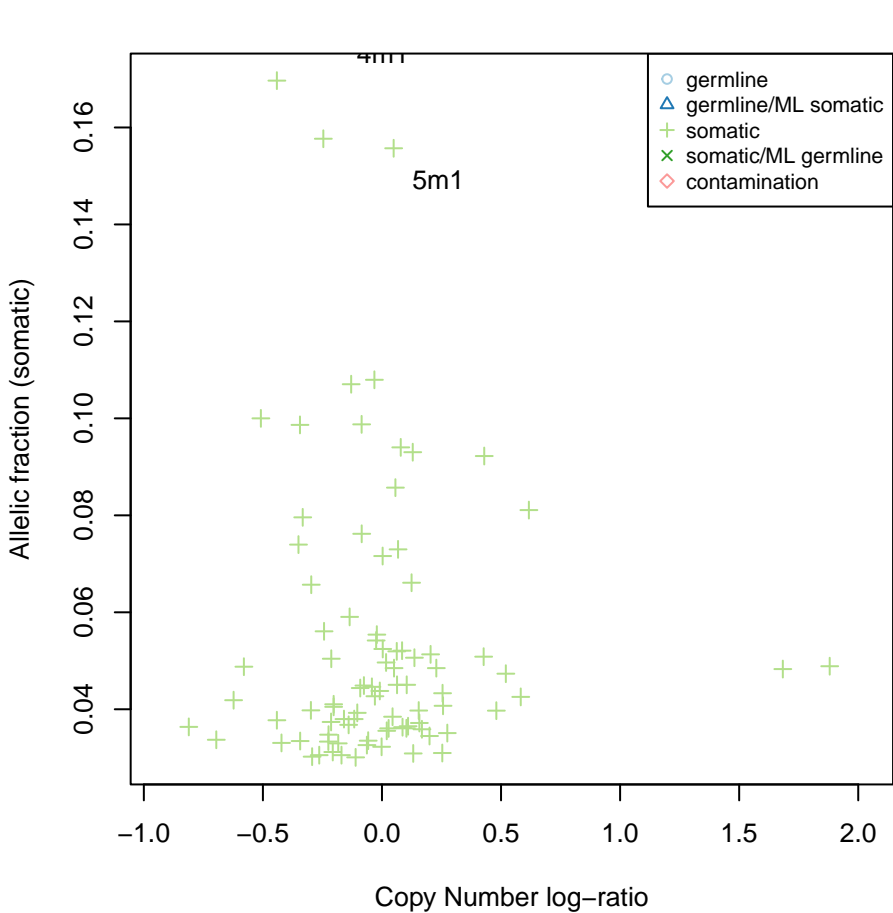
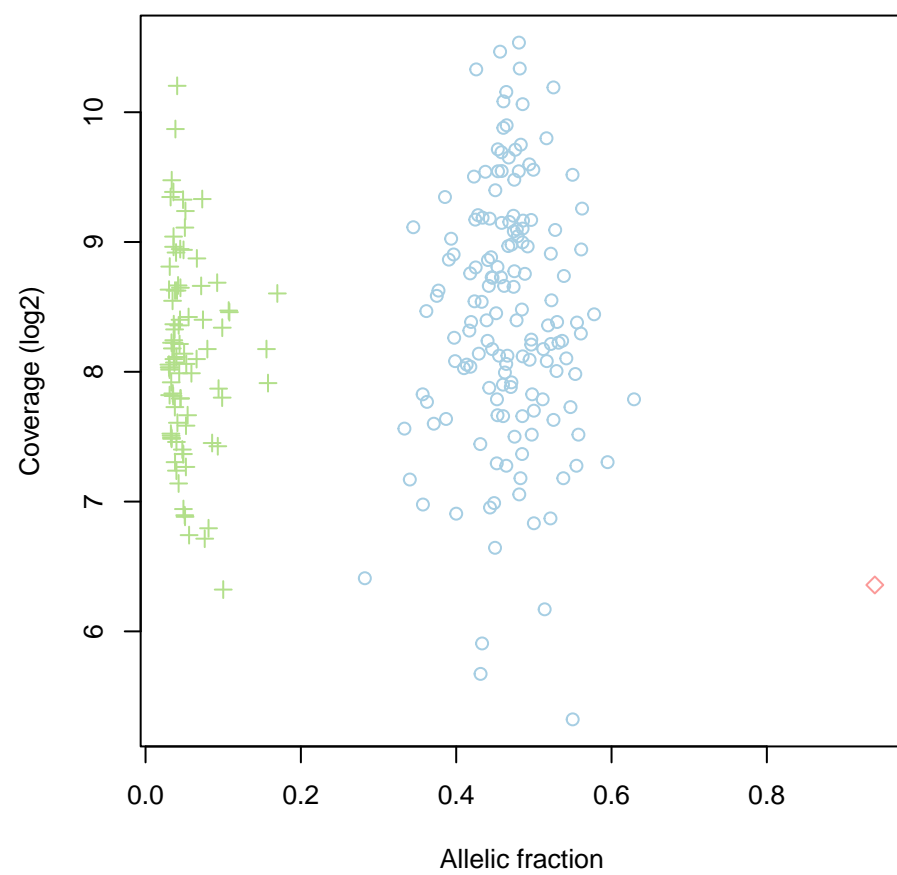
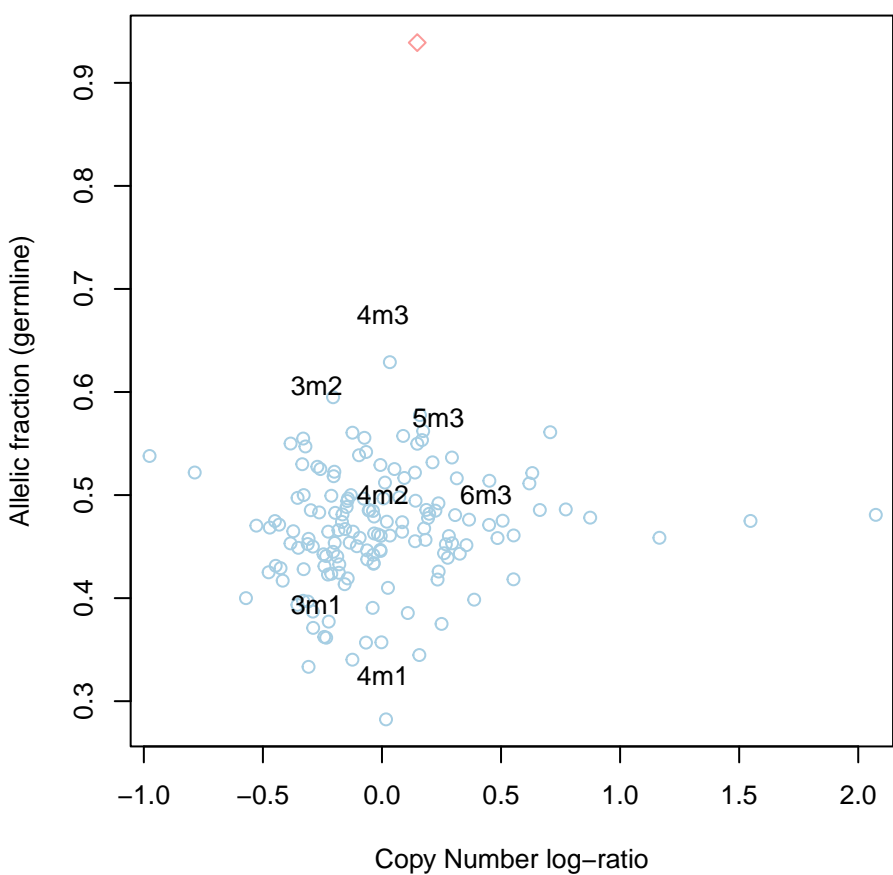


Purity: 0.54 Tumor ploidy: 3.985 SNV log-likelihood: -180.15 GoF: 81.3% Mean coverage: 108;387

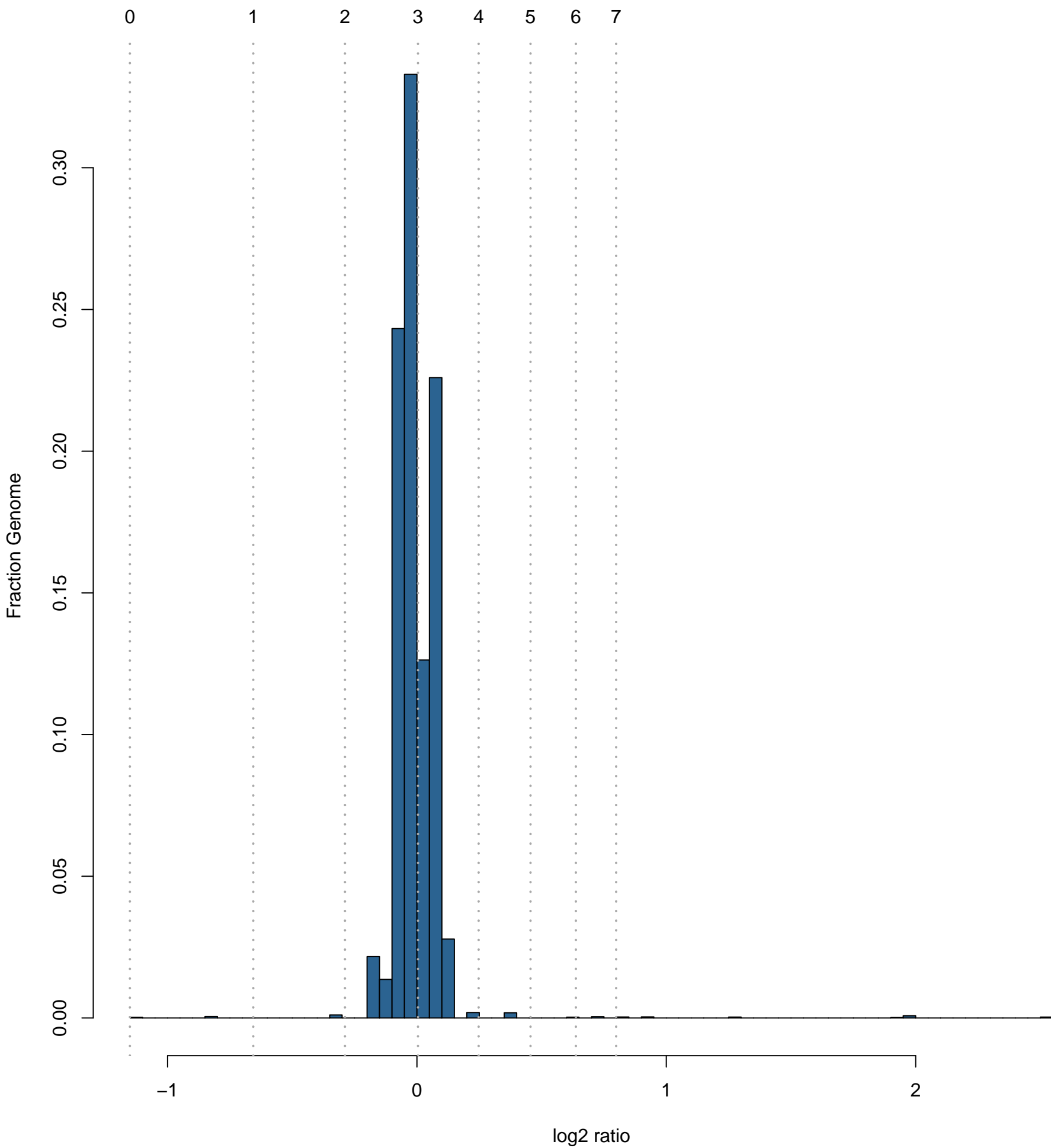


SCNA-fit log-likelihood: -15802.59

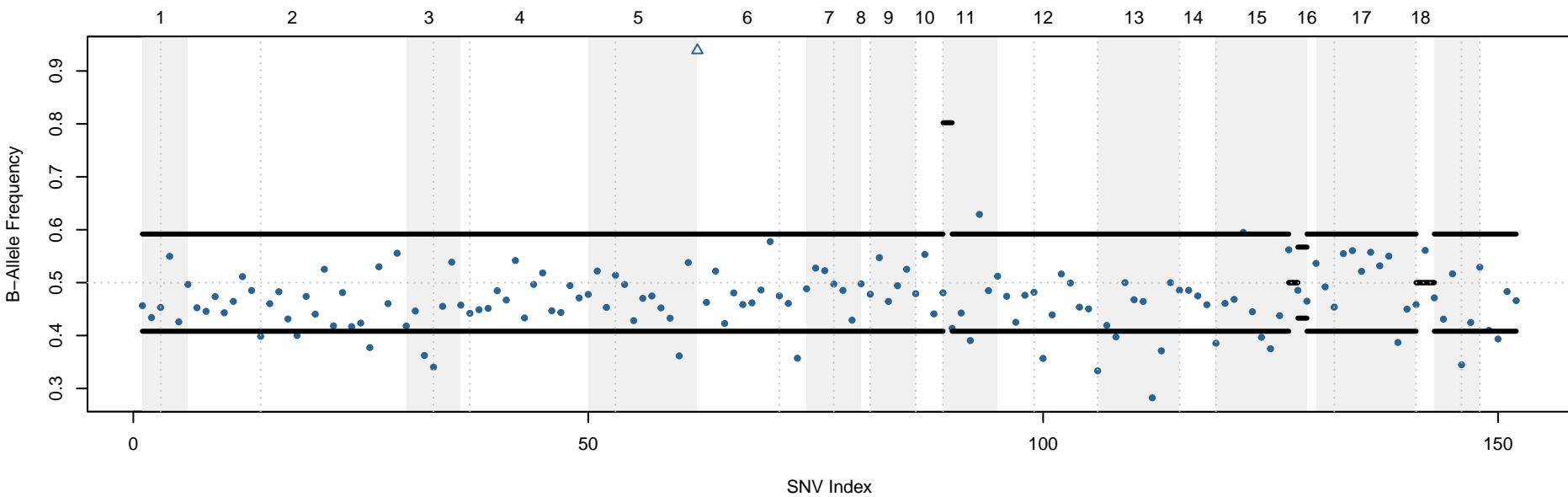




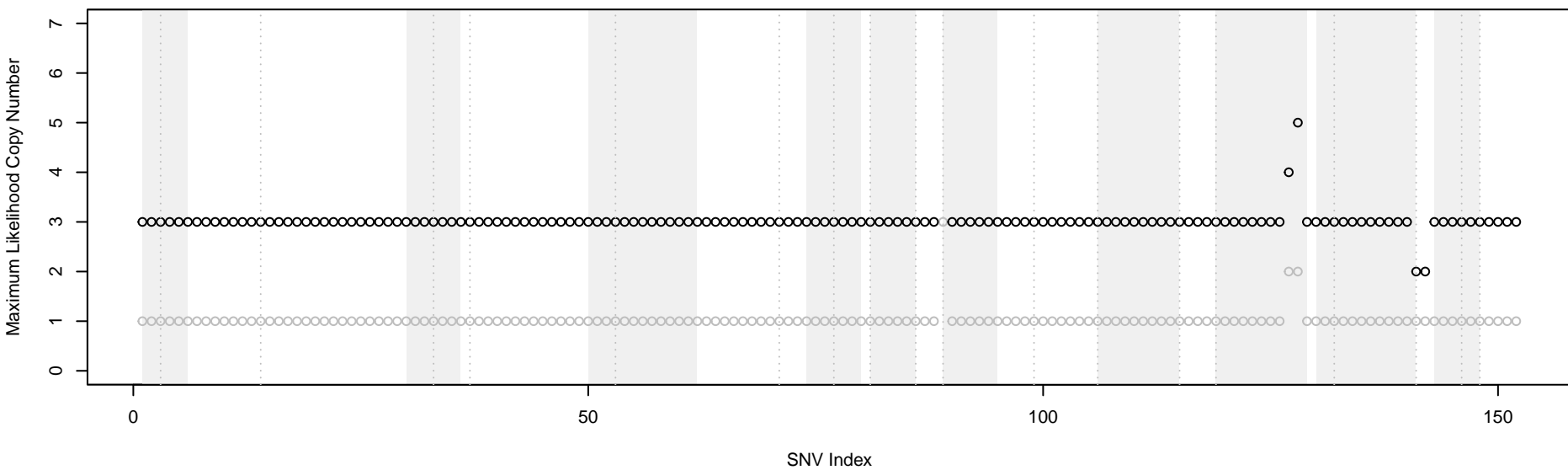
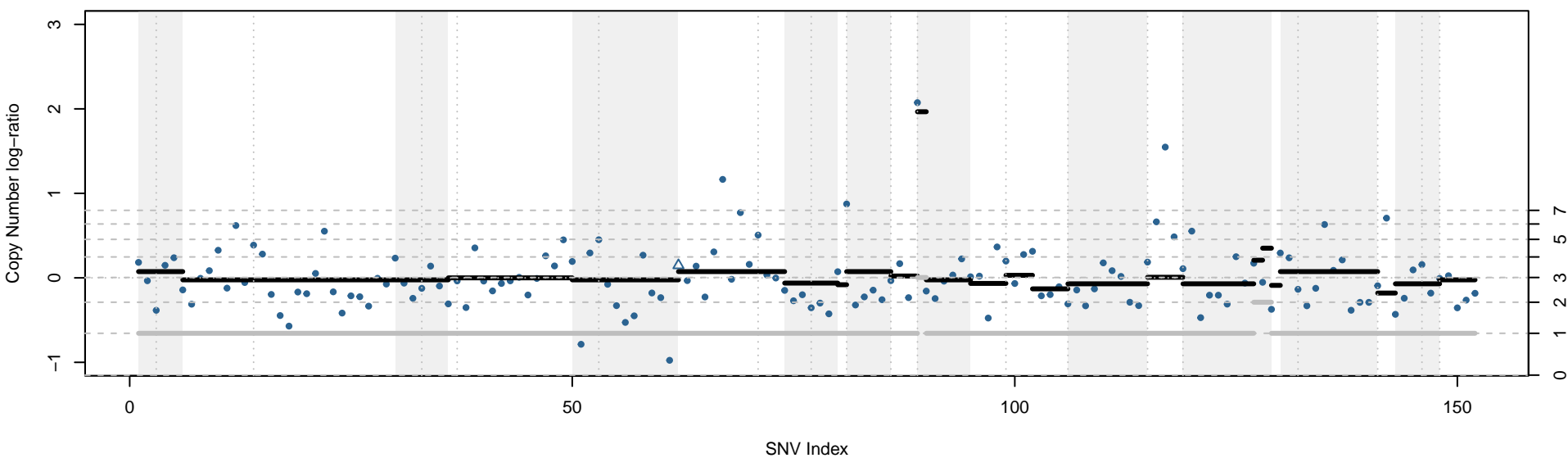
Purity: 0.45 Tumor ploidy: 2.984

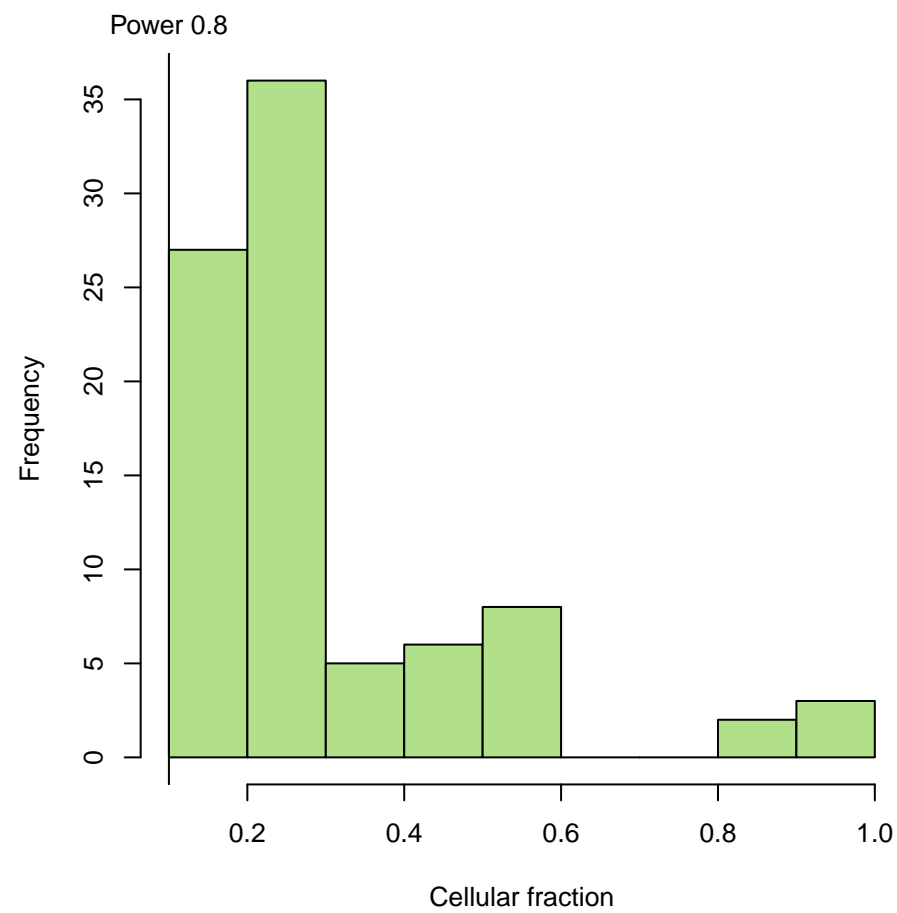
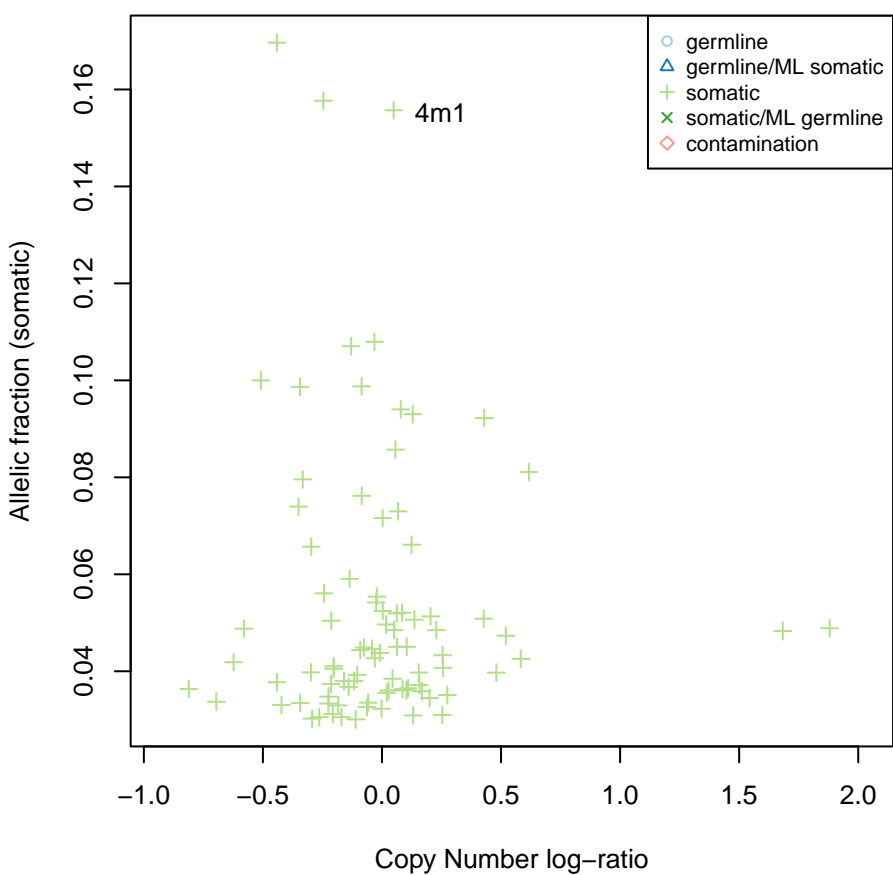
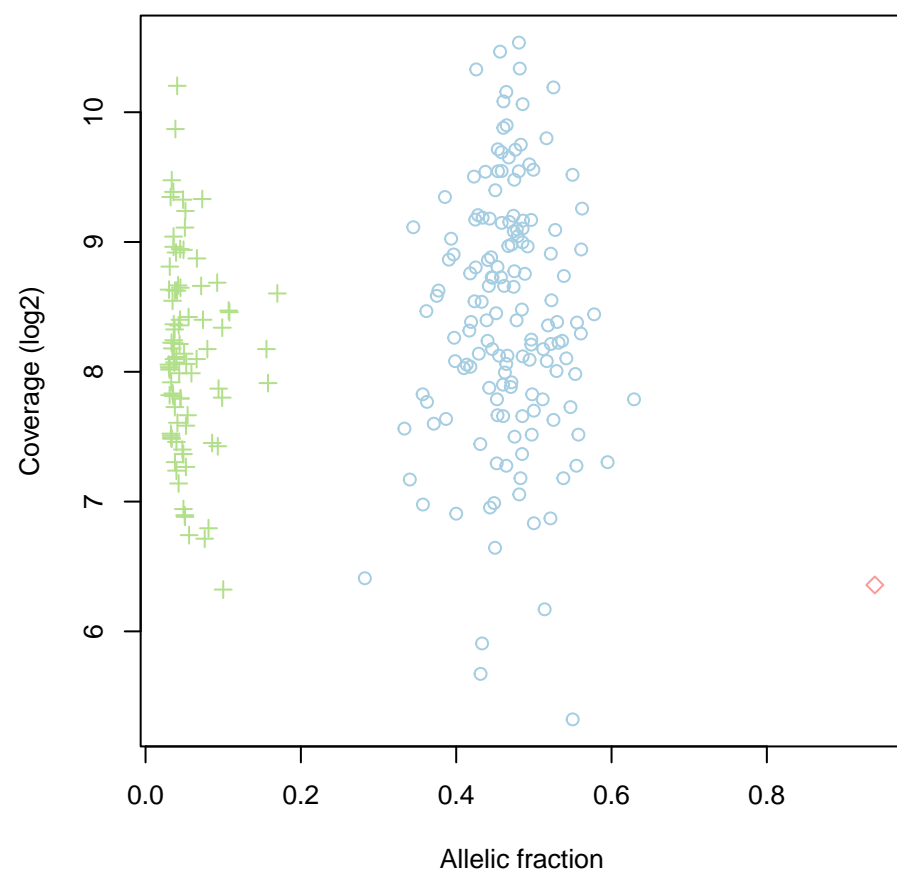
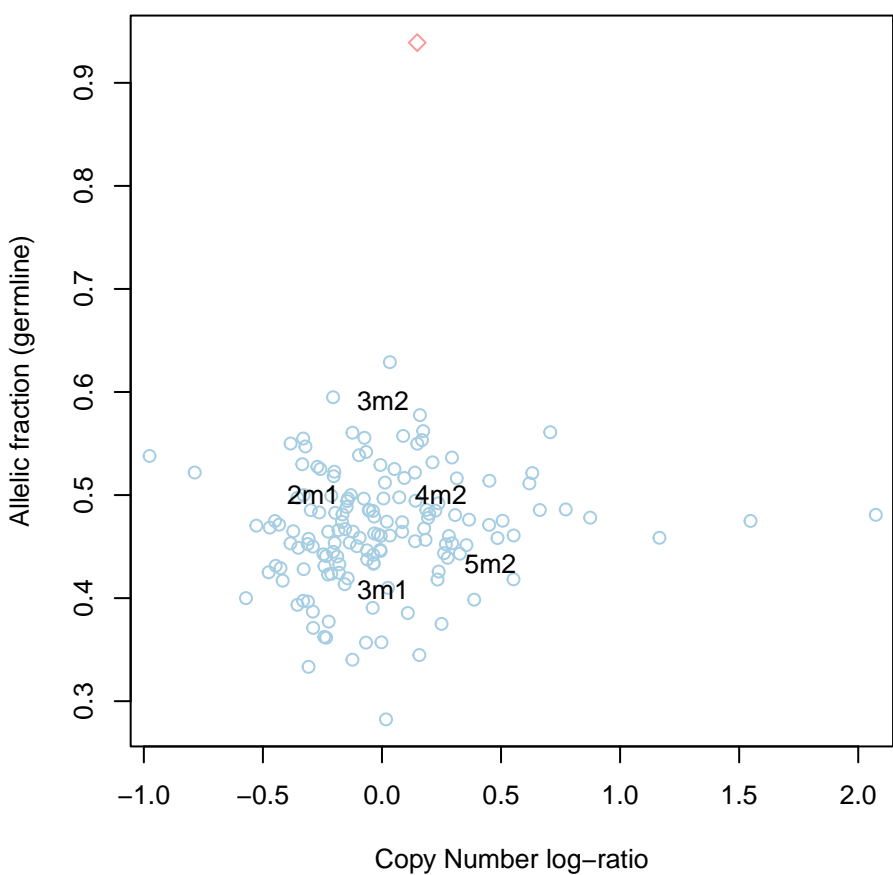


Purity: 0.45 Tumor ploidy: 2.984 SNV log-likelihood: -254.36 GoF: 79% Mean coverage: 108;387

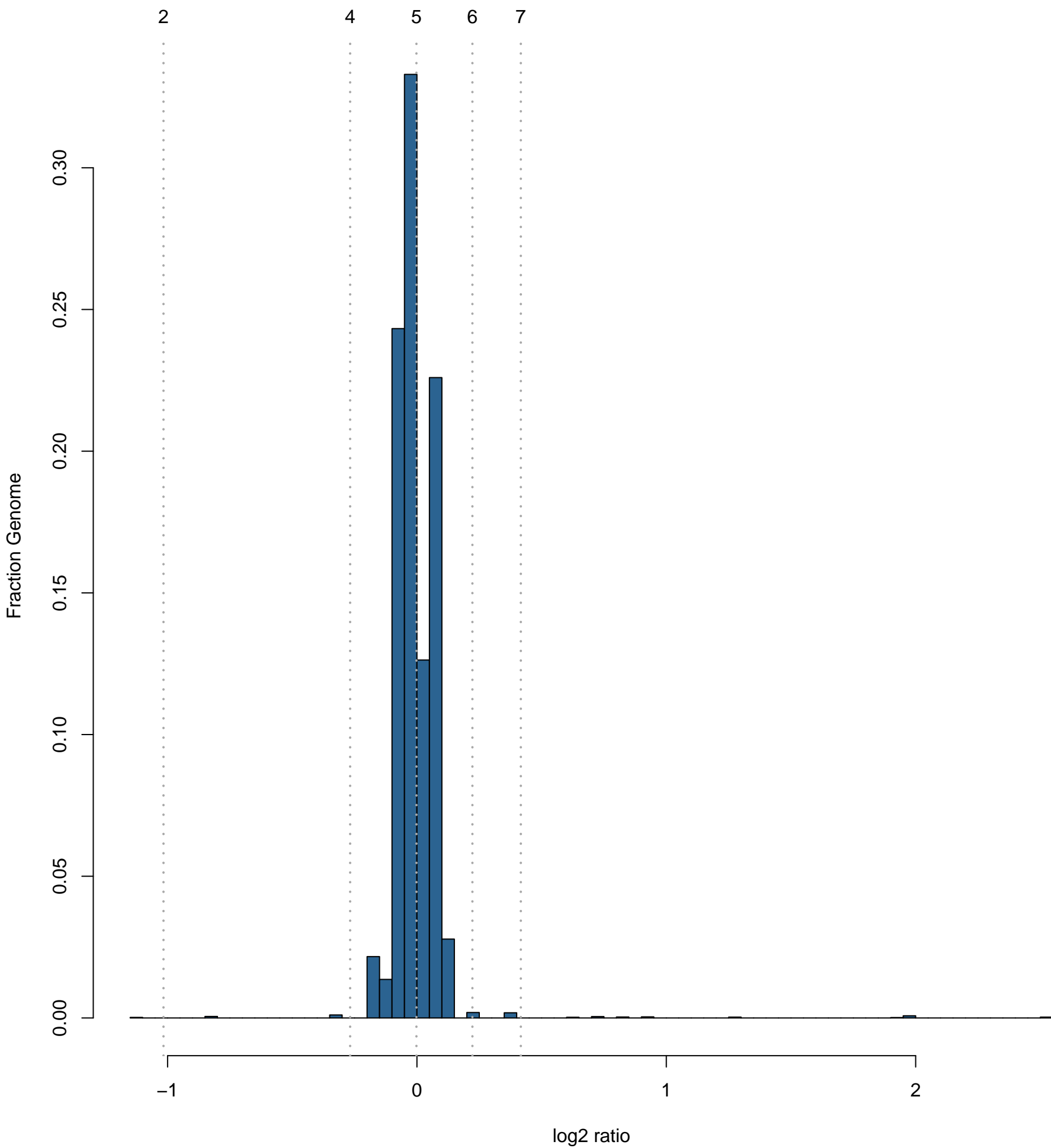


SCNA-fit log-likelihood: -15798.24

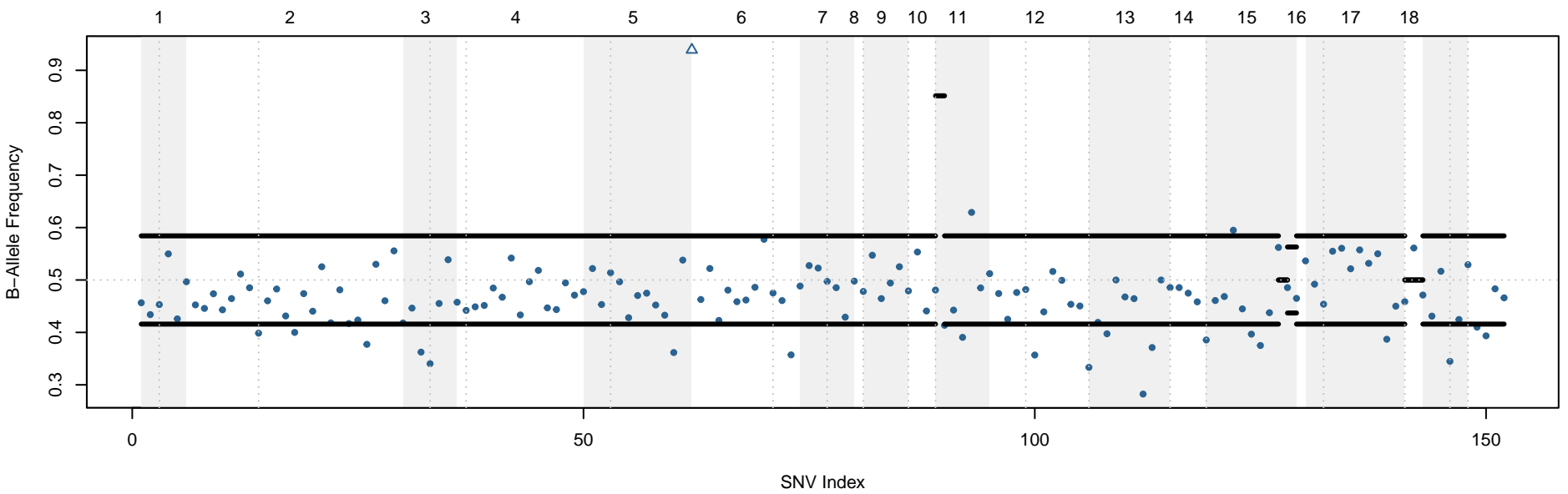




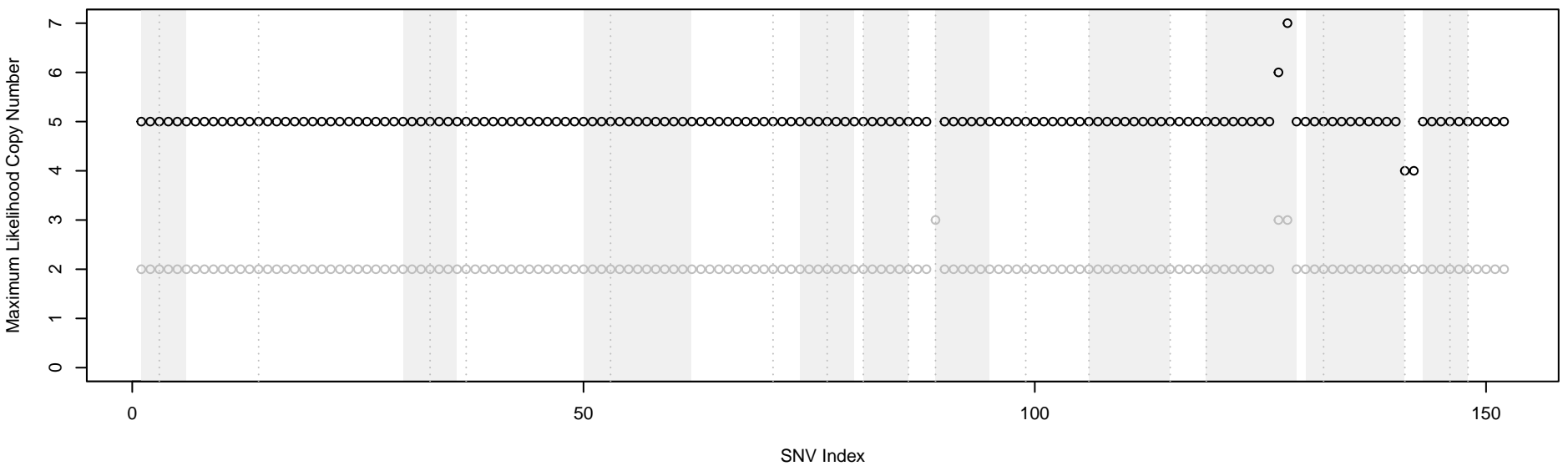
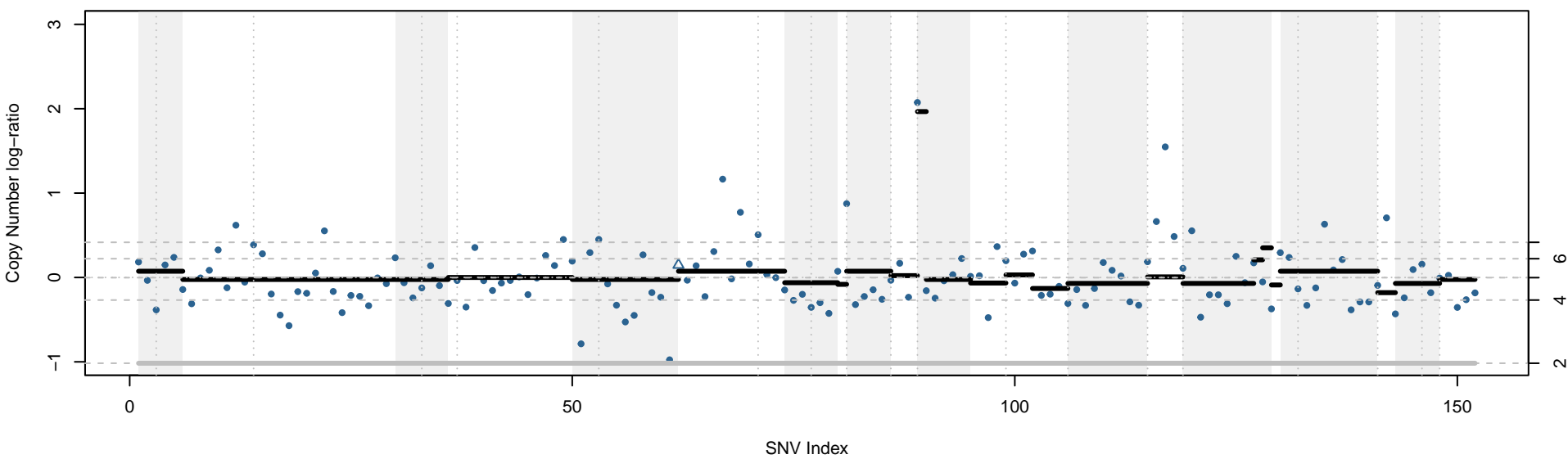
Purity: 0.68 Tumor ploidy: 5.008

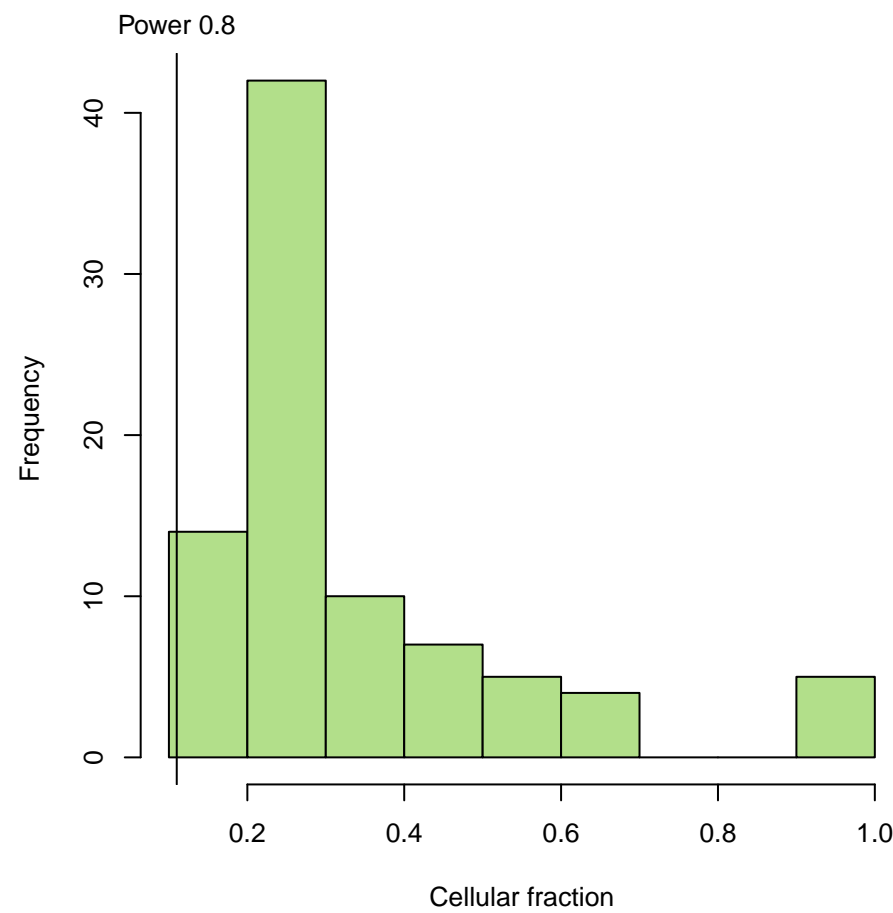
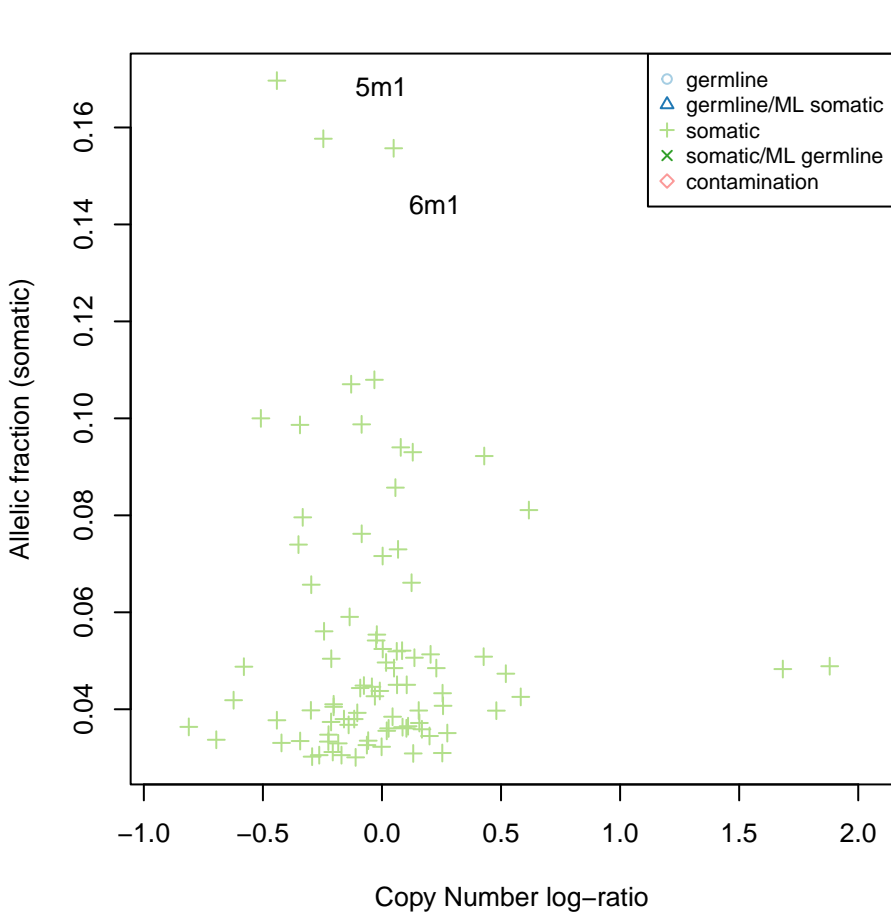
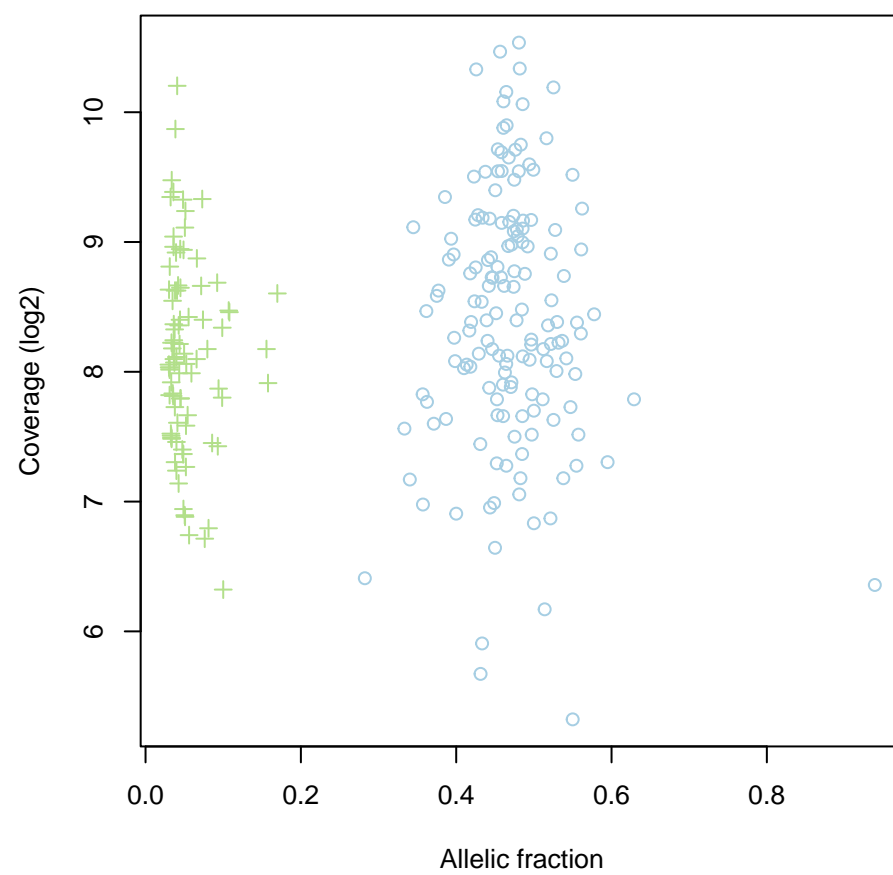
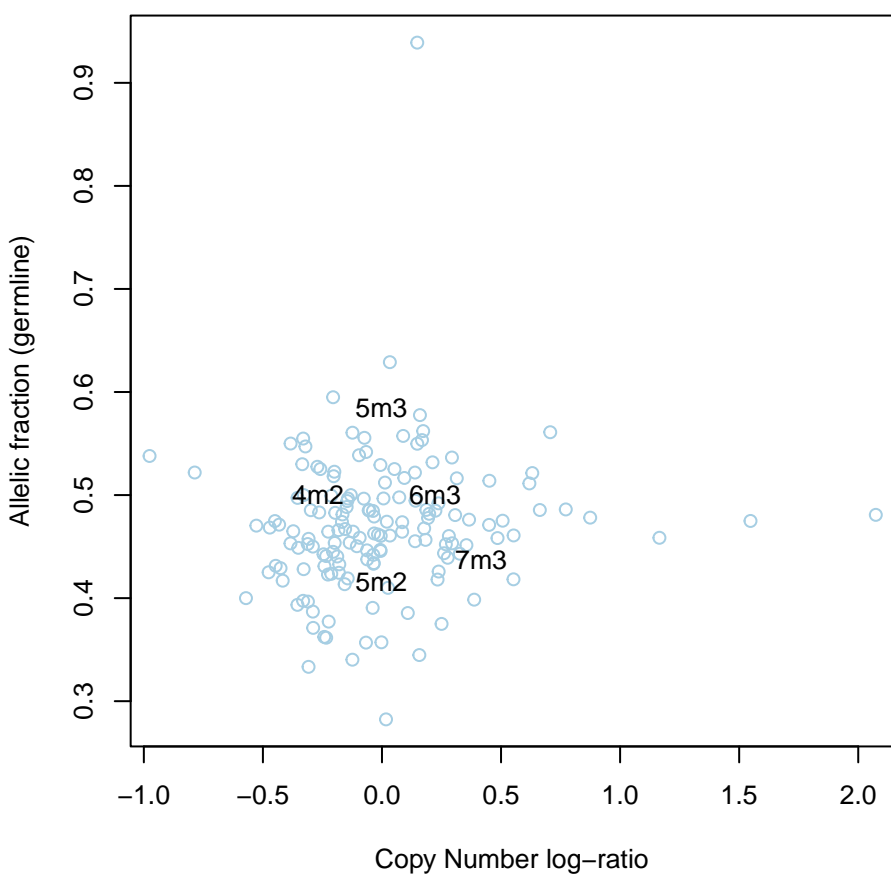


Purity: 0.68 Tumor ploidy: 5.008 SNV log-likelihood: -307.77 GoF: 83.2% Mean coverage: 108,387

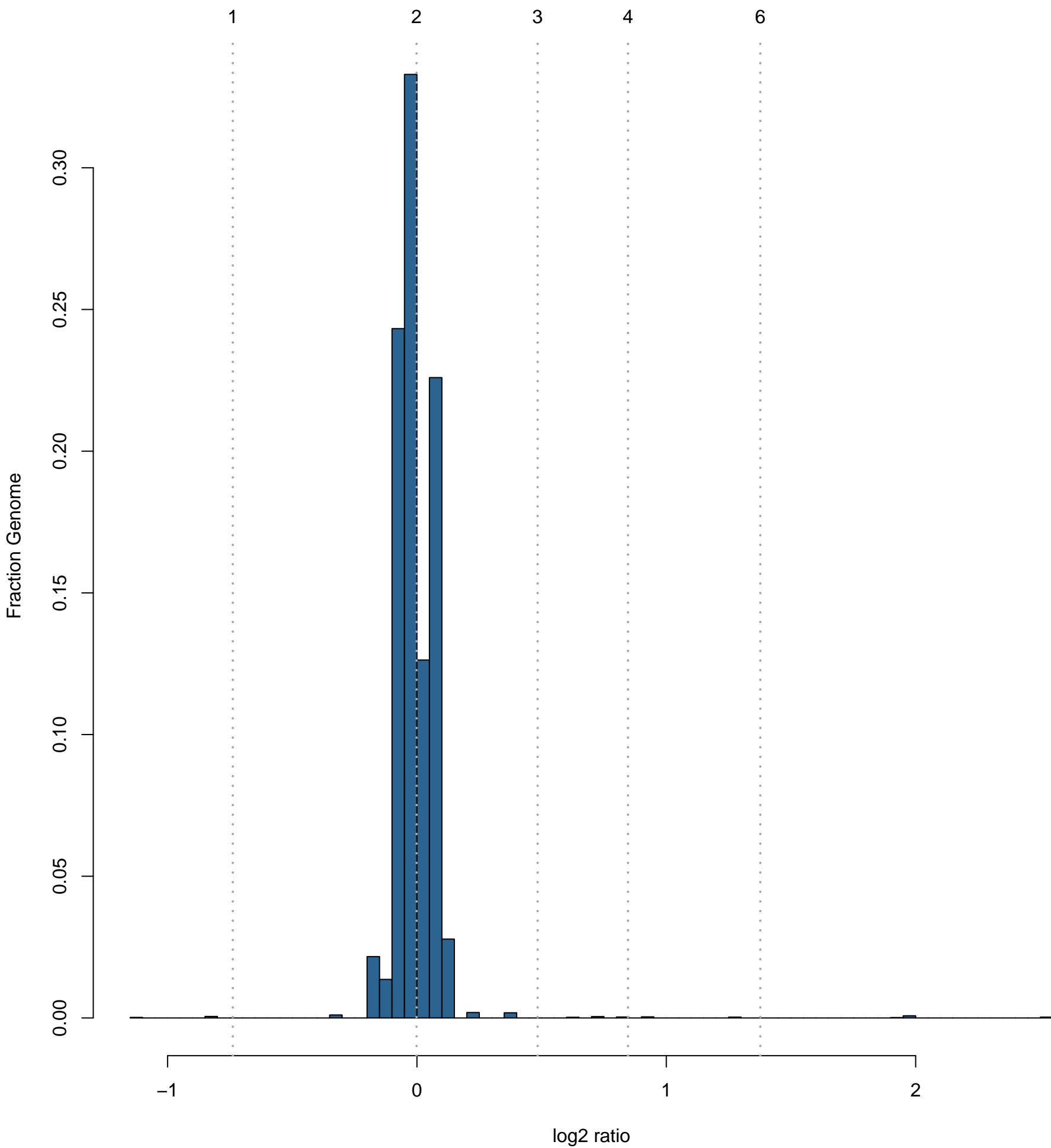


SCNA-fit log-likelihood: -15808

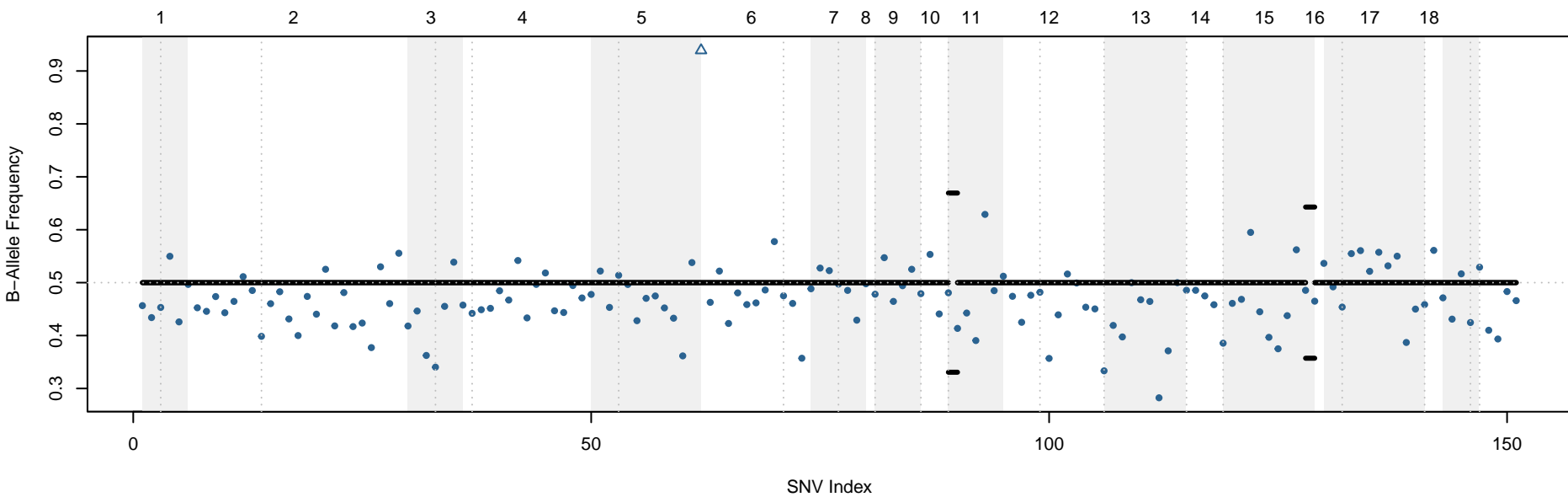




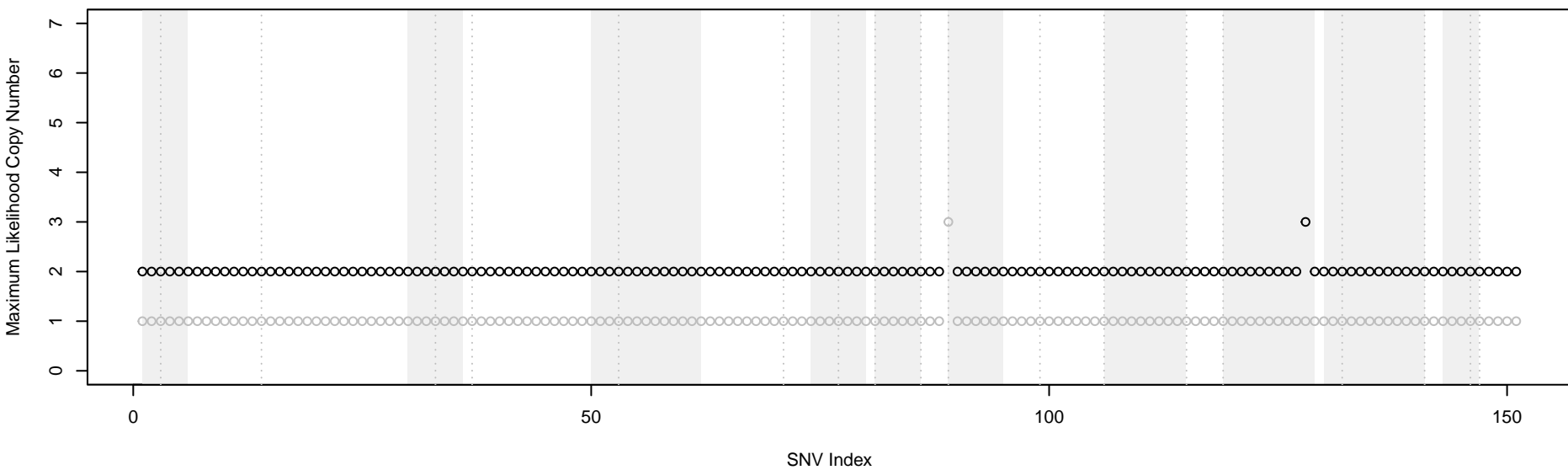
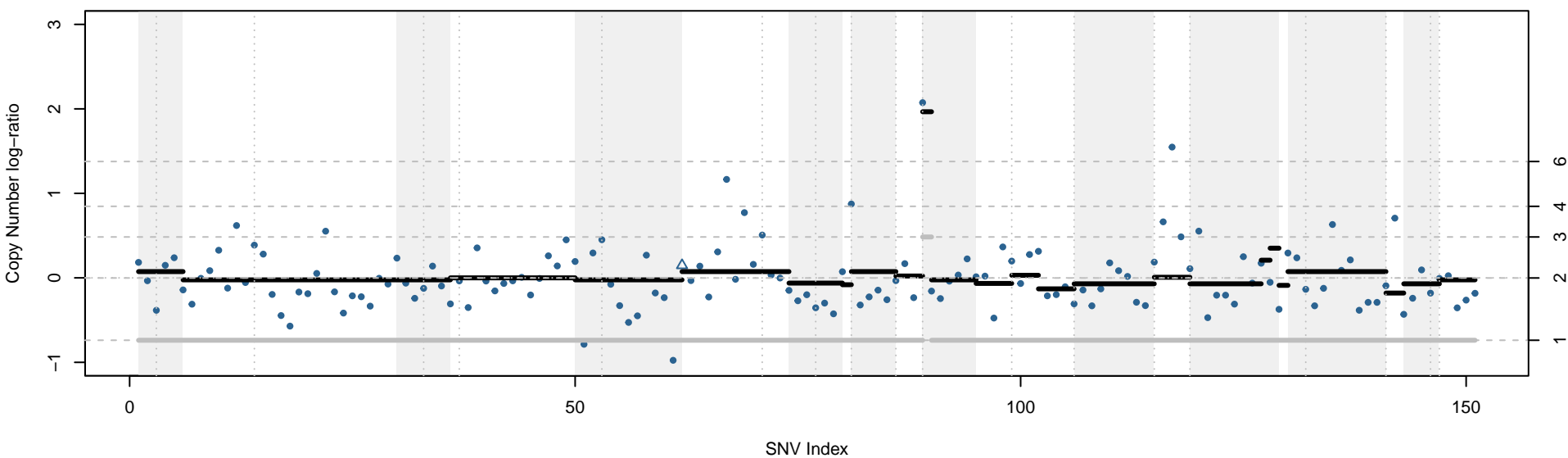
Purity: 0.8 Tumor ploidy: 2.003

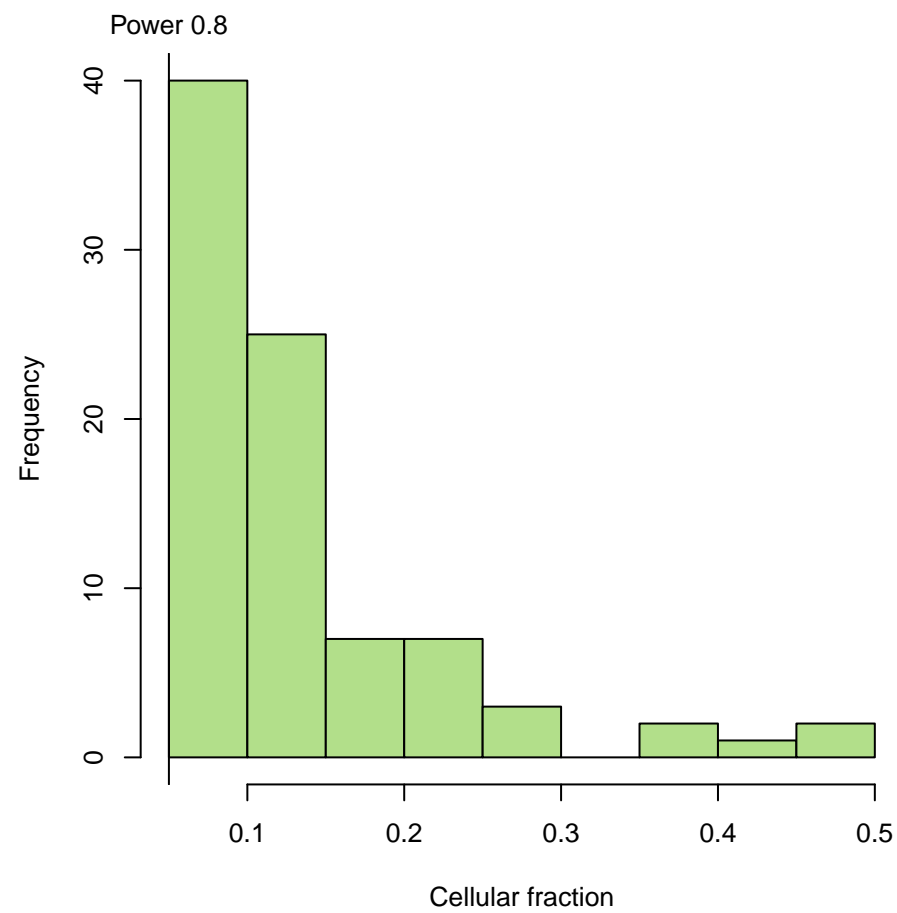
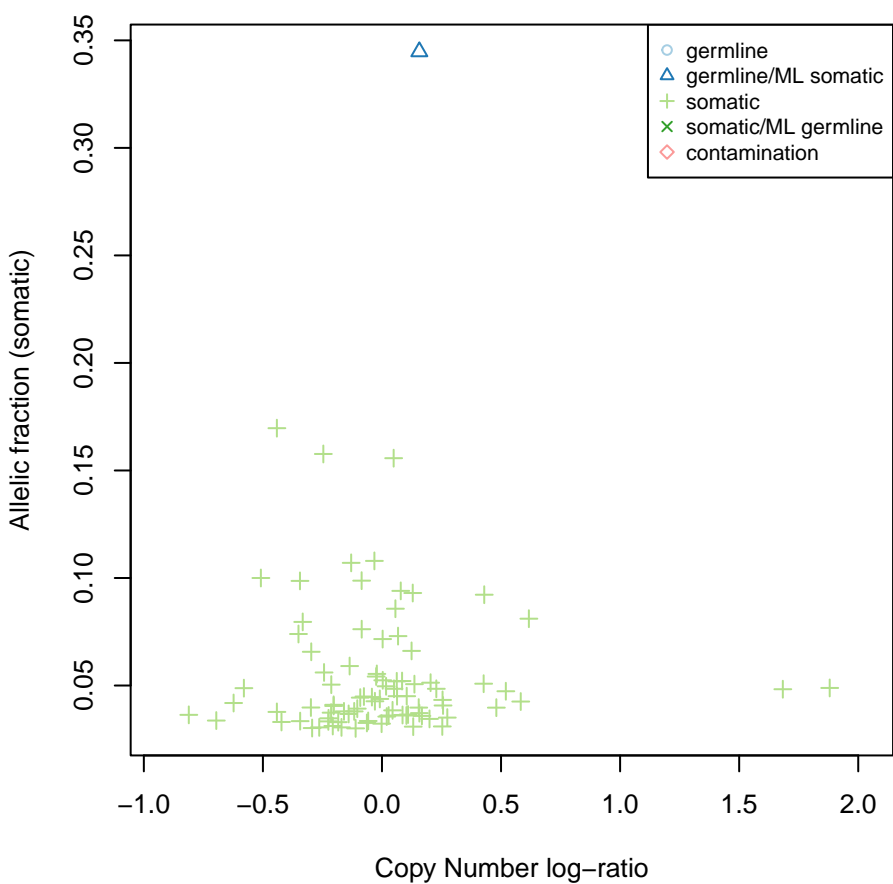
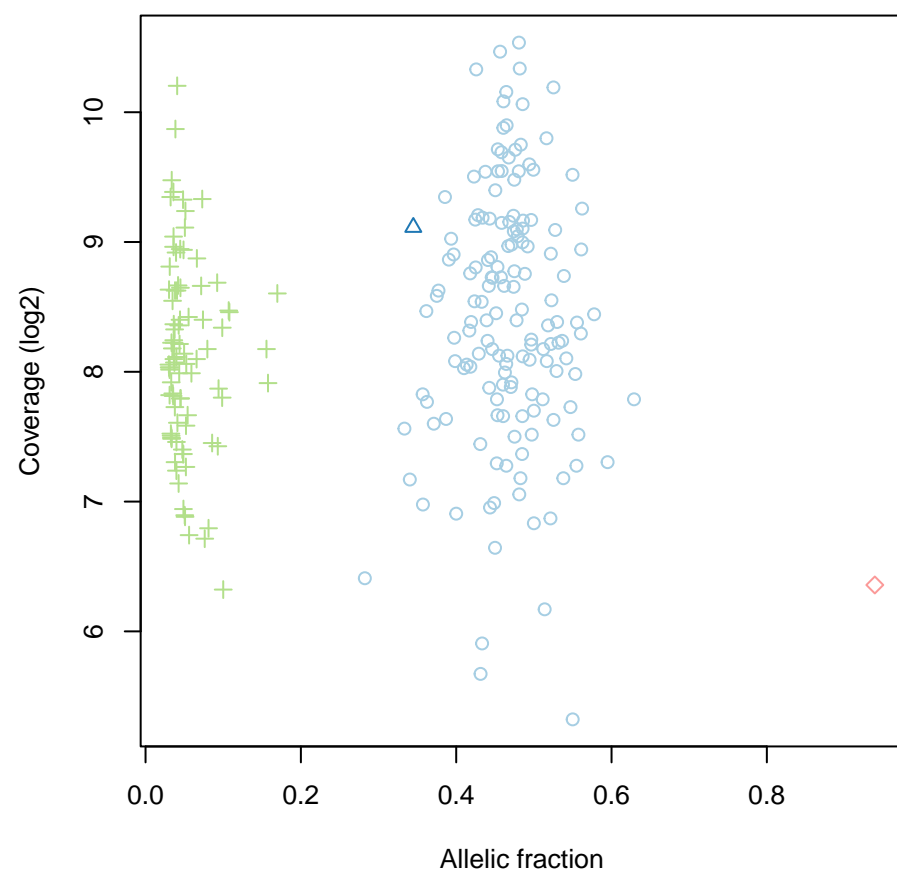
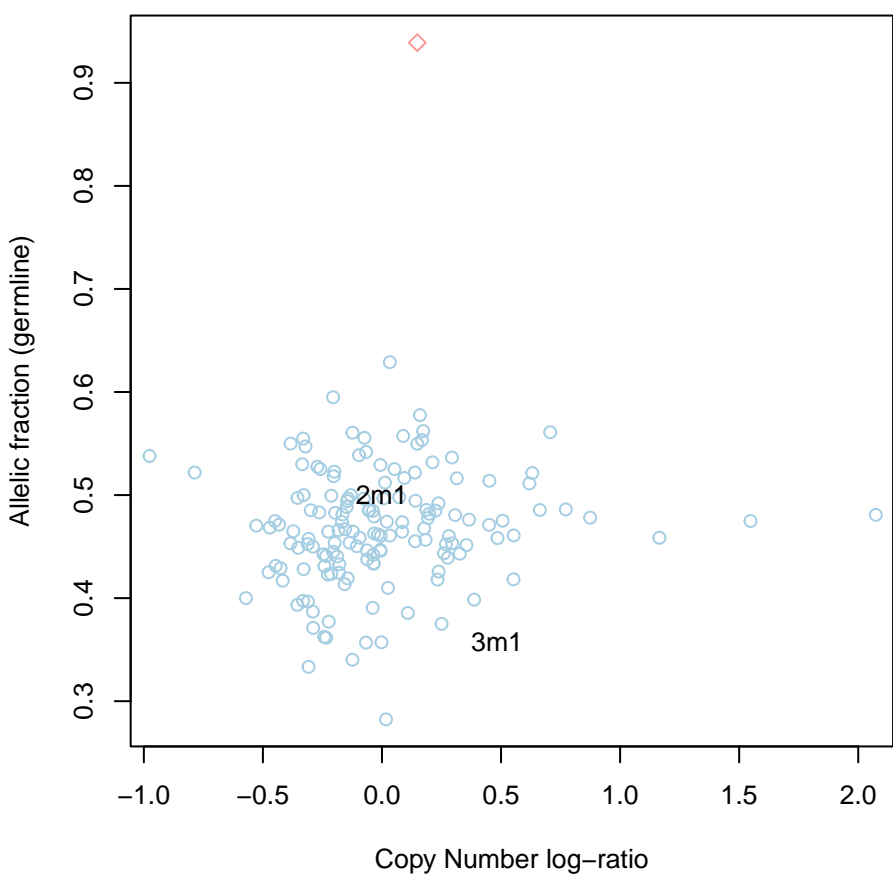


Purity: 0.8 Tumor ploidy: 2.003 SNV log-likelihood: -696.23 GoF: 0% Mean coverage: 108;387

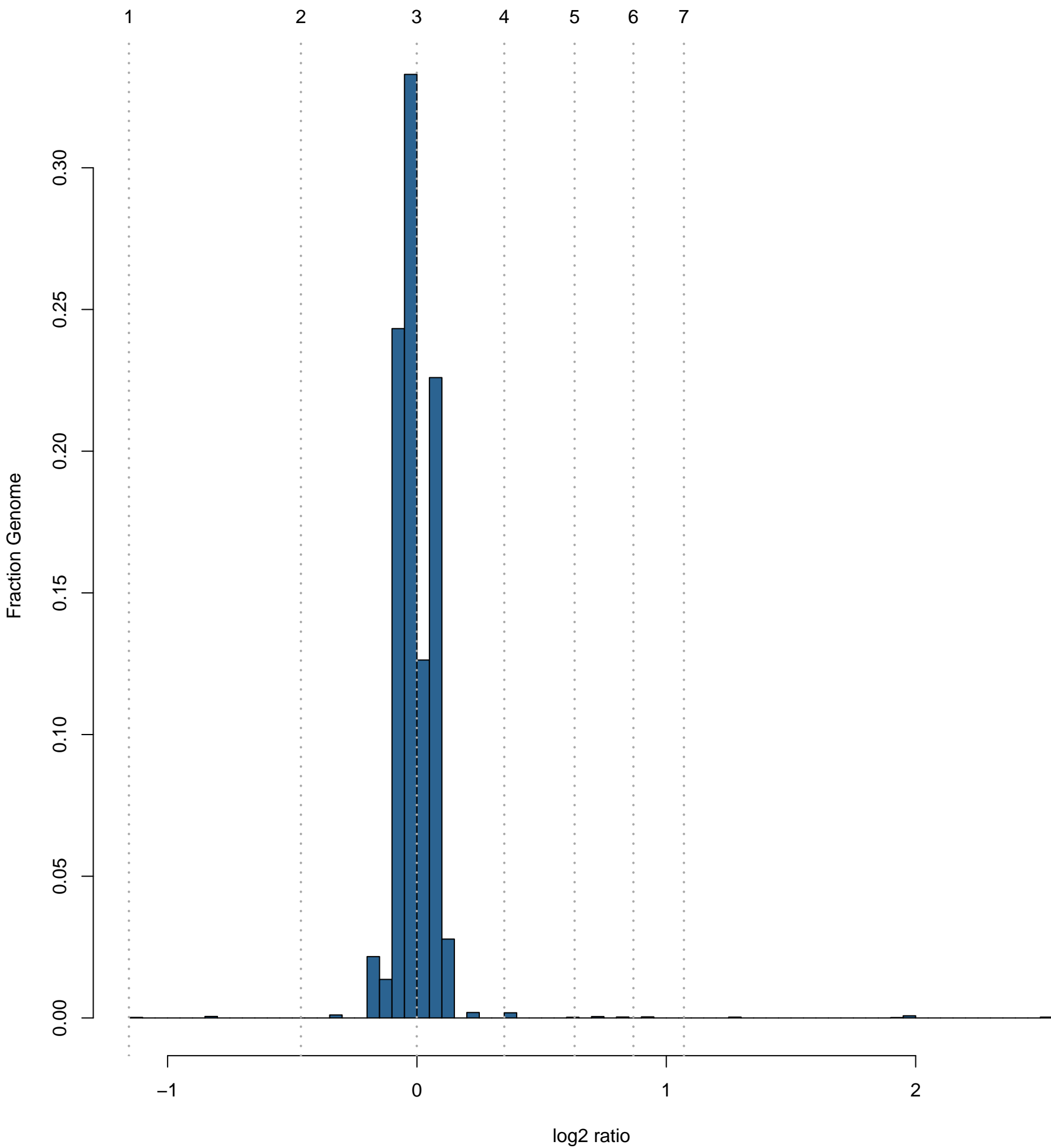


SCNA-fit log-likelihood: -15819.15

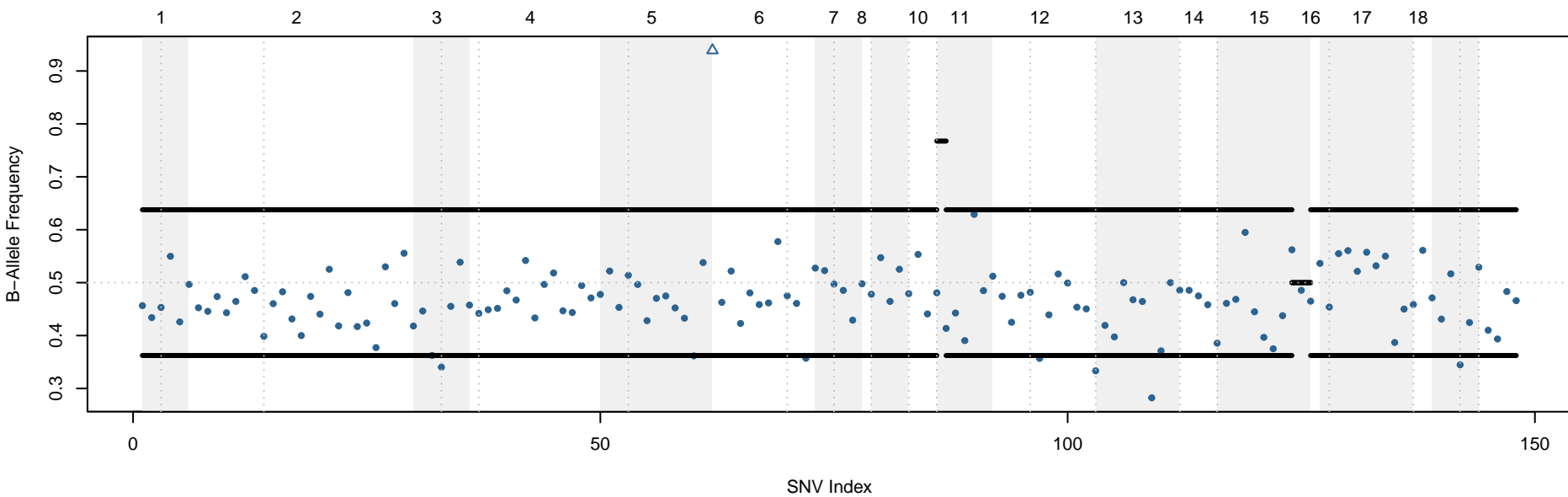




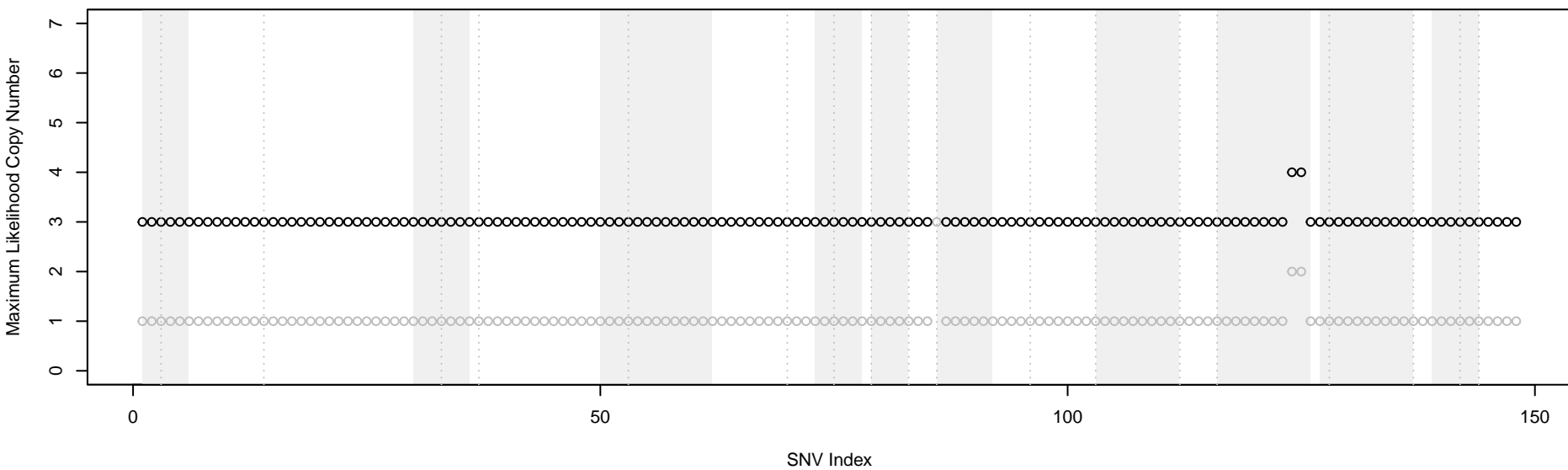
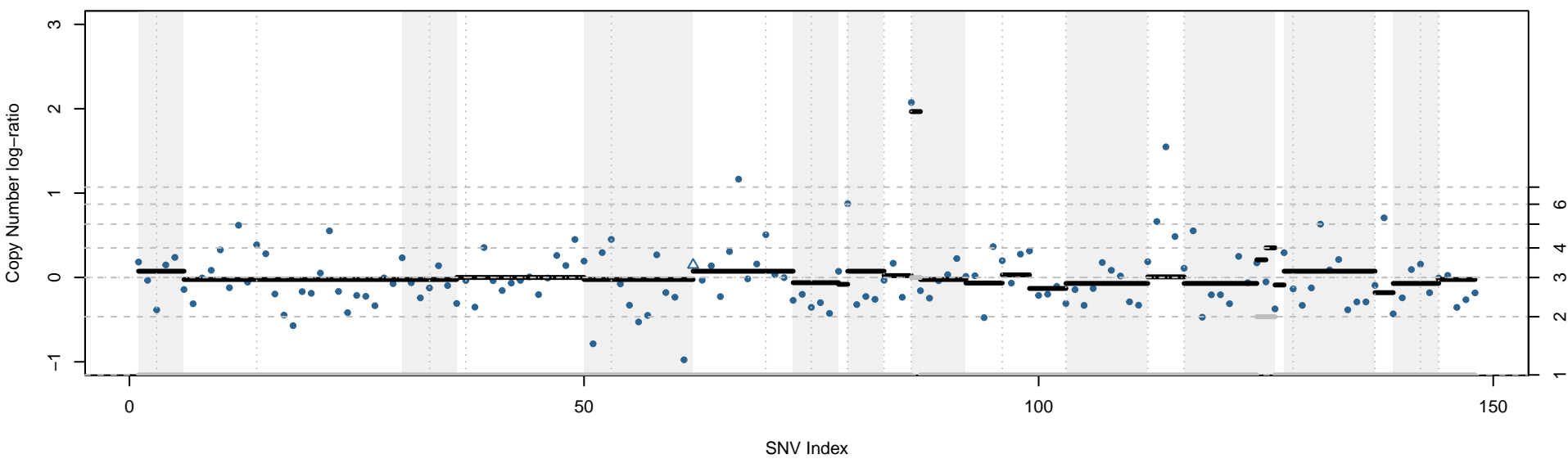
Purity: 0.76 Tumor ploidy: 3.002

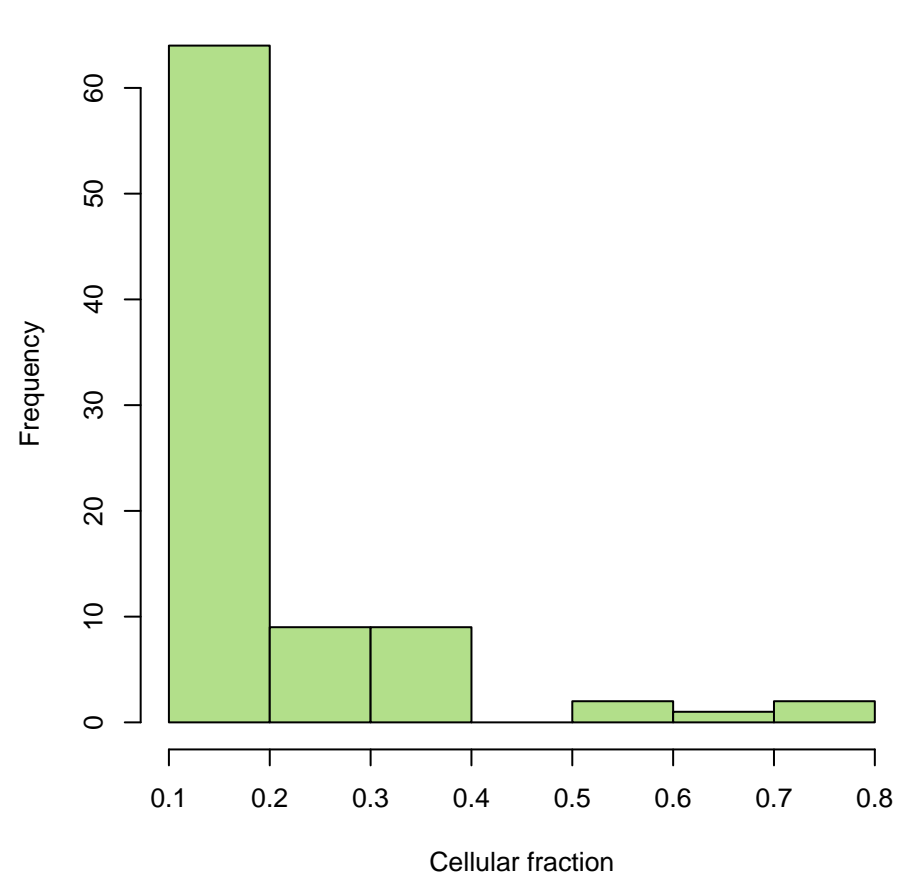
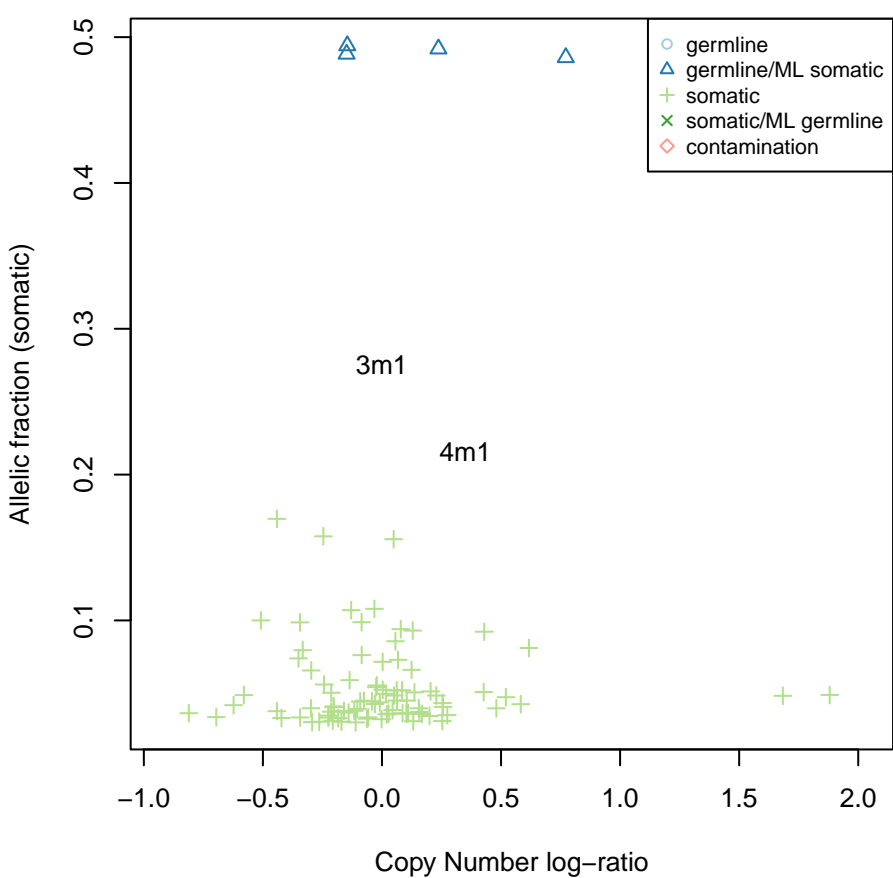
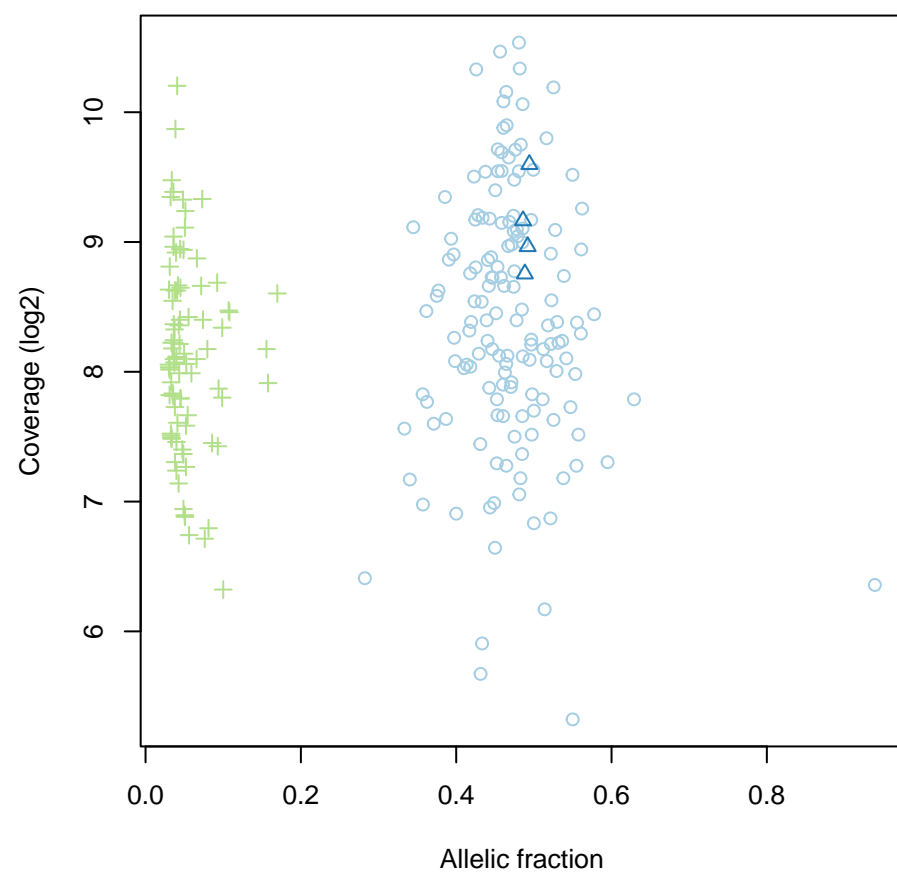
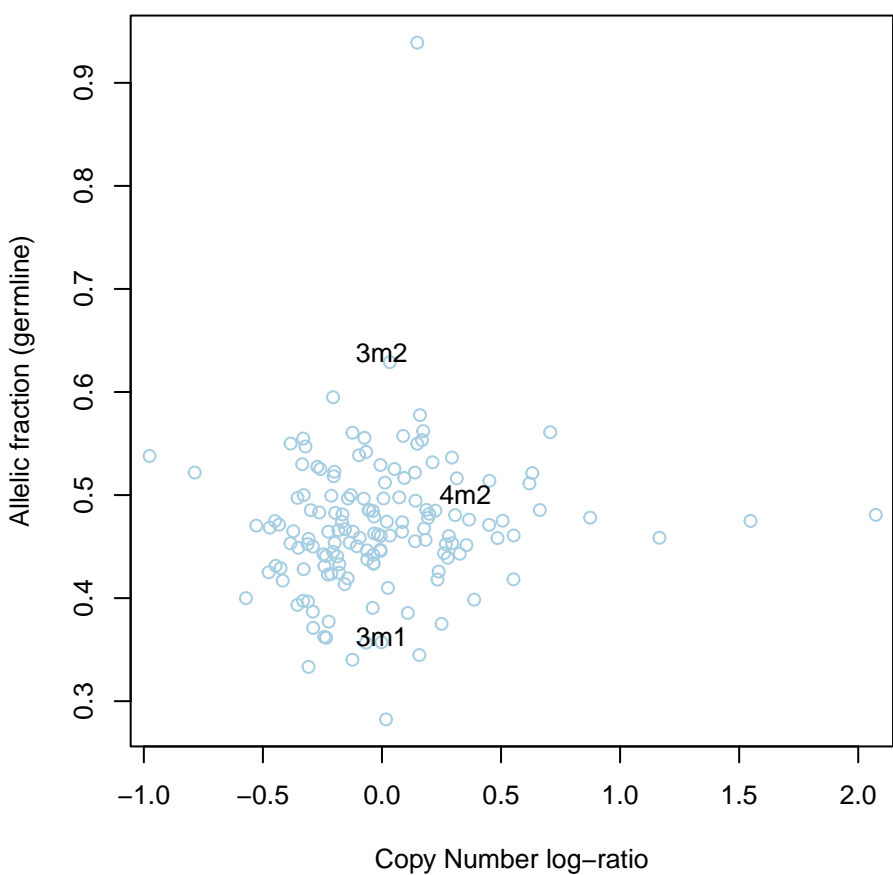


Purity: 0.76 Tumor ploidy: 3.002 SNV log-likelihood: -974.74 GoF: 41.9% Mean coverage: 108;387

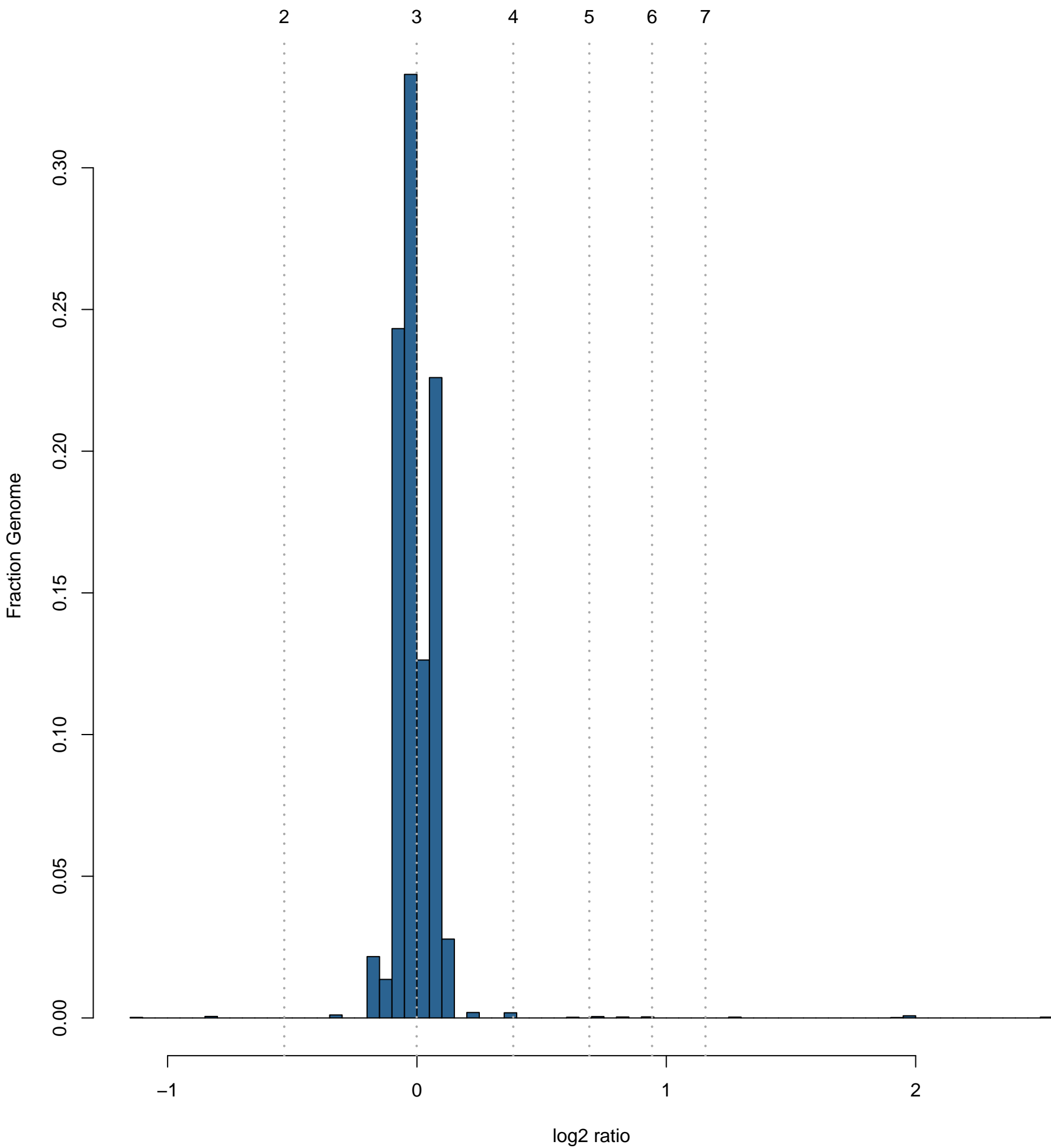


SCNA-fit log-likelihood: -15812.68

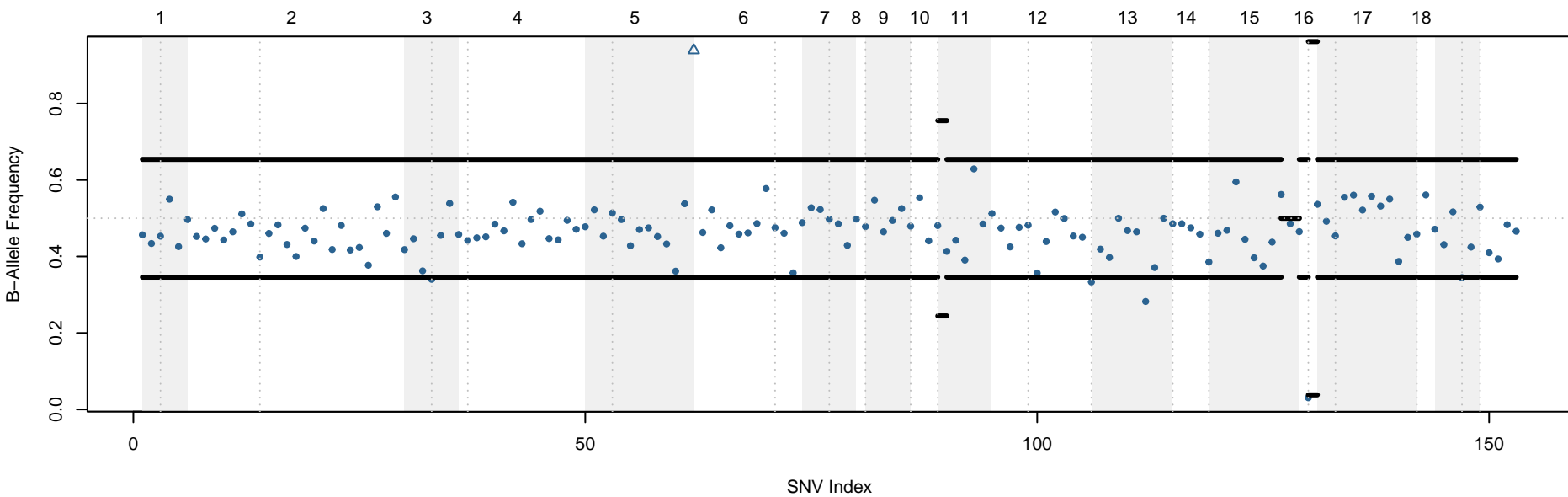




Purity: 0.89 Tumor ploidy: 3.002



Purity: 0.89 Tumor ploidy: 3.002 SNV log-likelihood: -1342.46 GoF: 23.4% Mean coverage: 108;387



SCNA-fit log-likelihood: -15814.87

