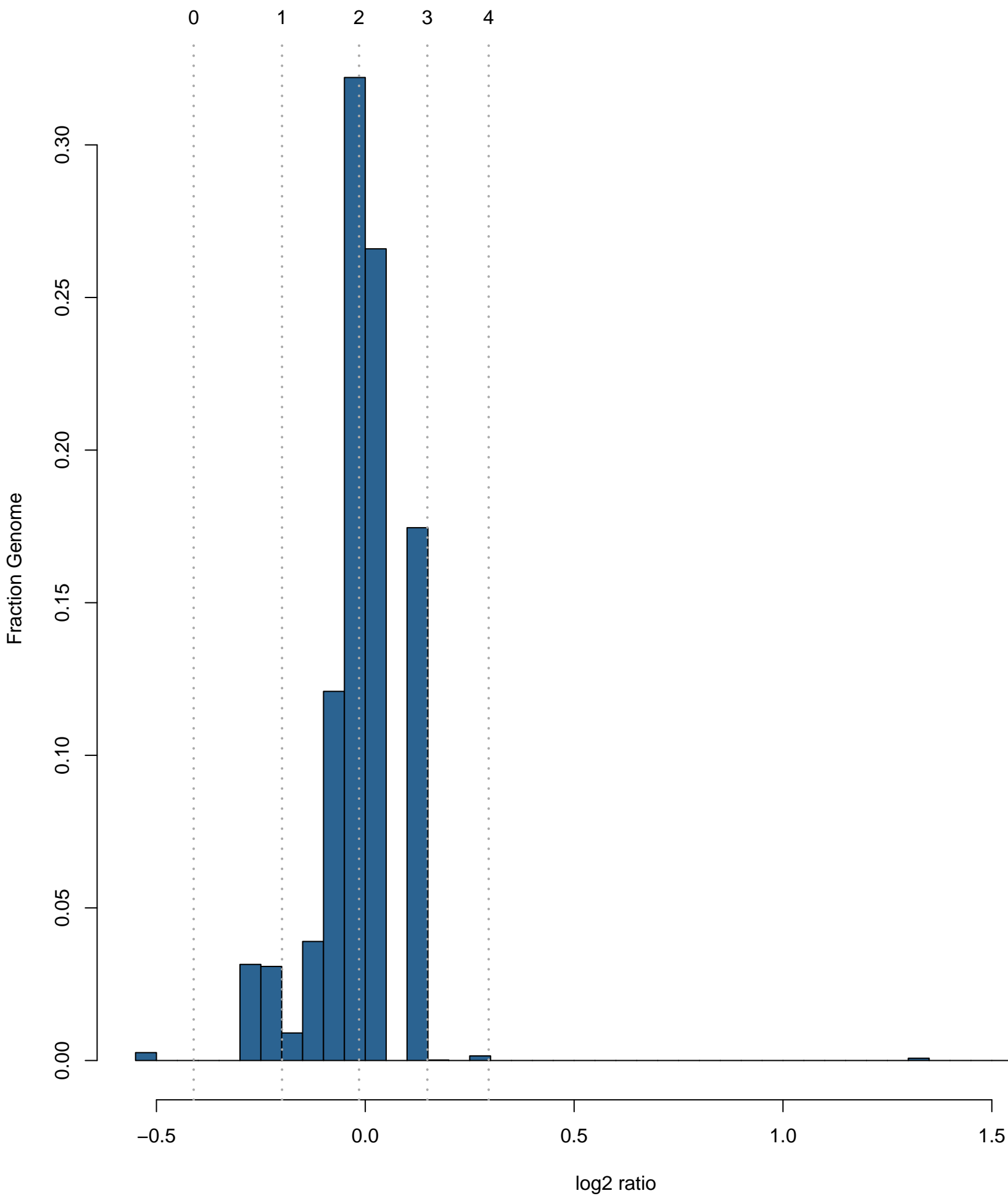
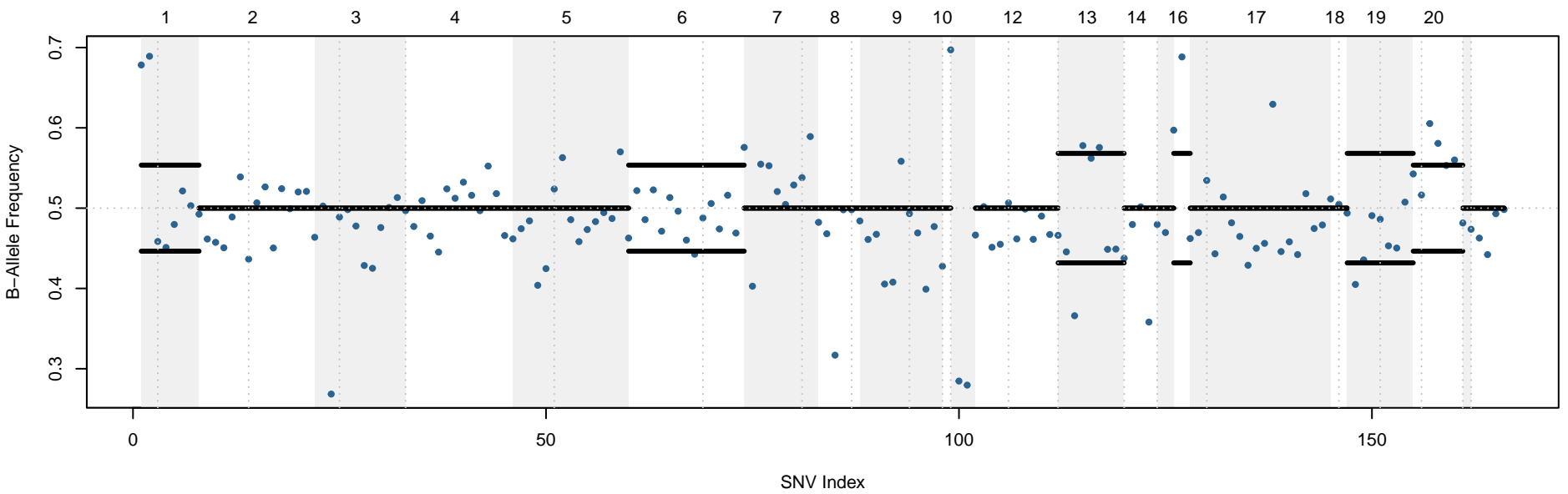


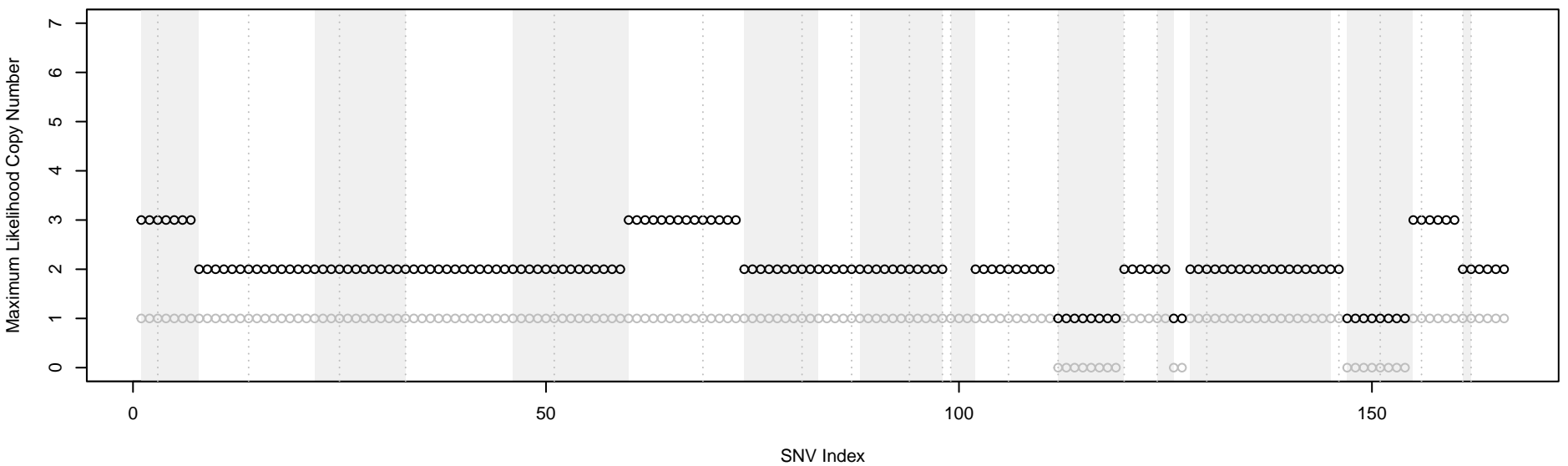
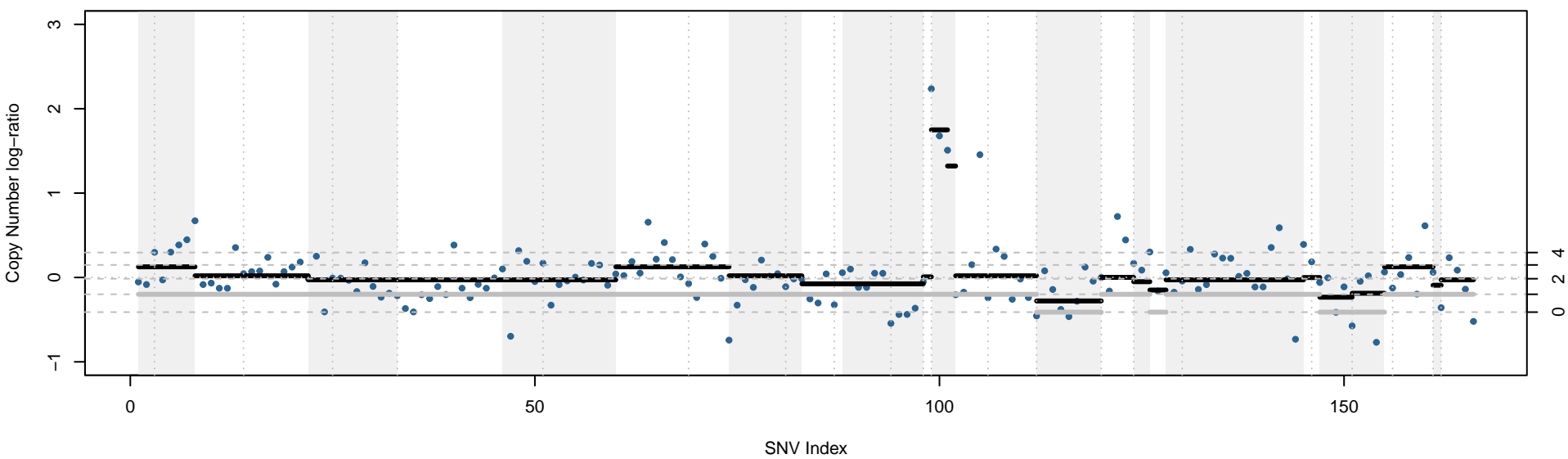
Purity: 0.24 Tumor ploidy: 2.087

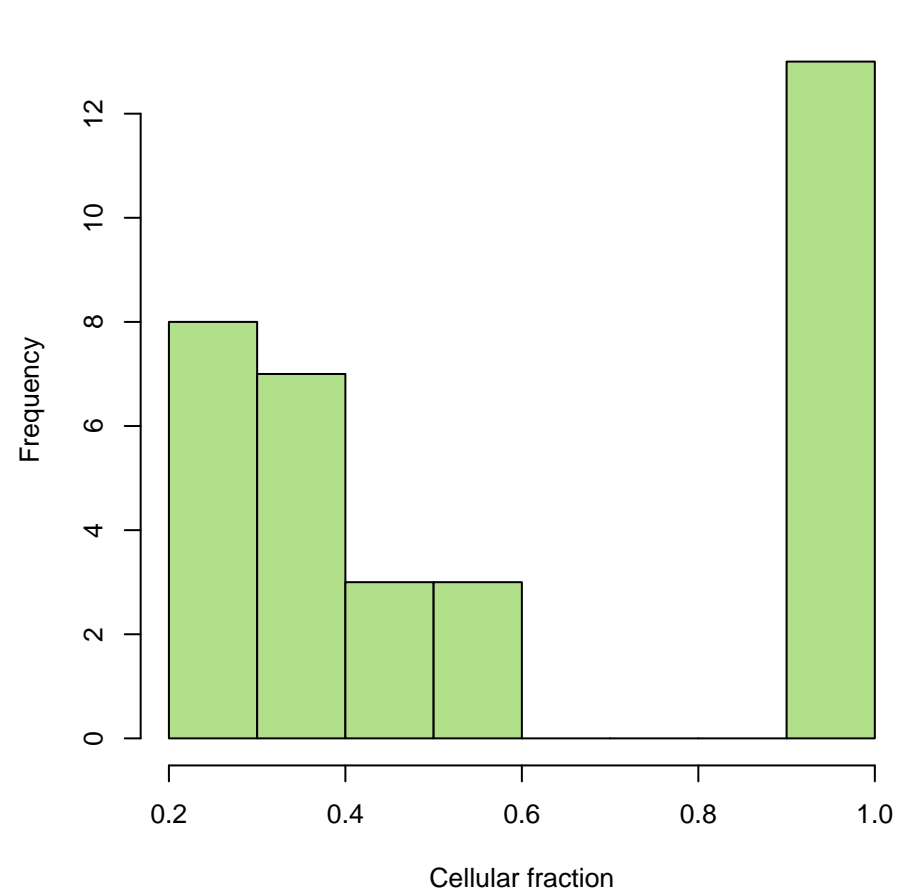
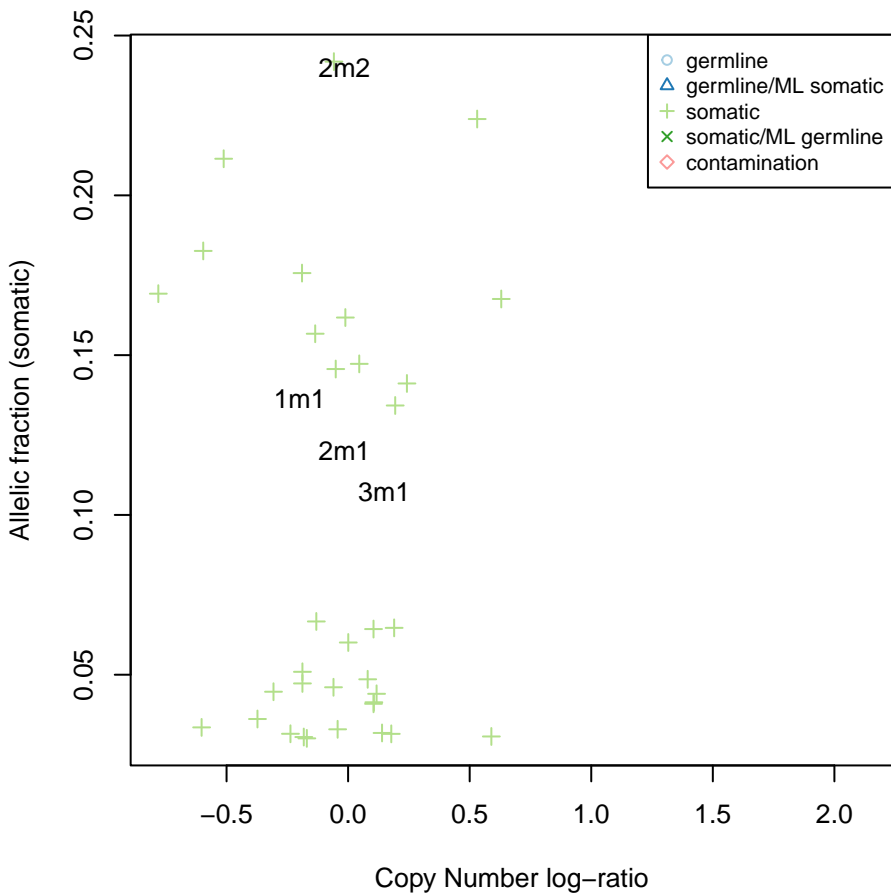
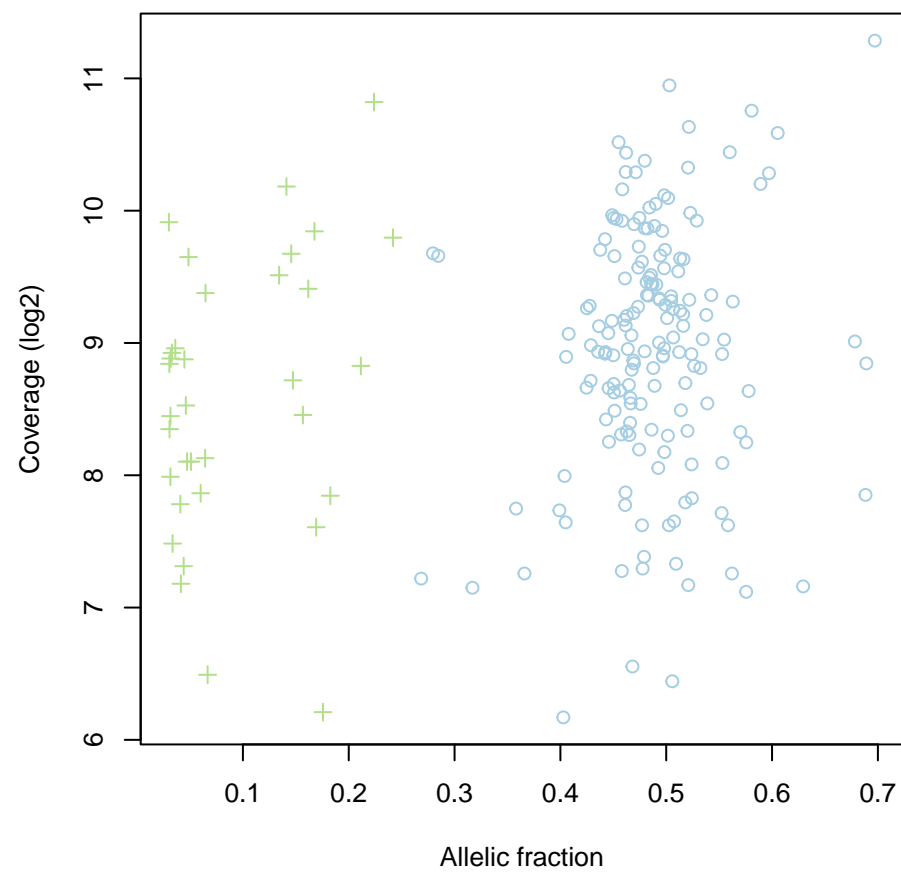
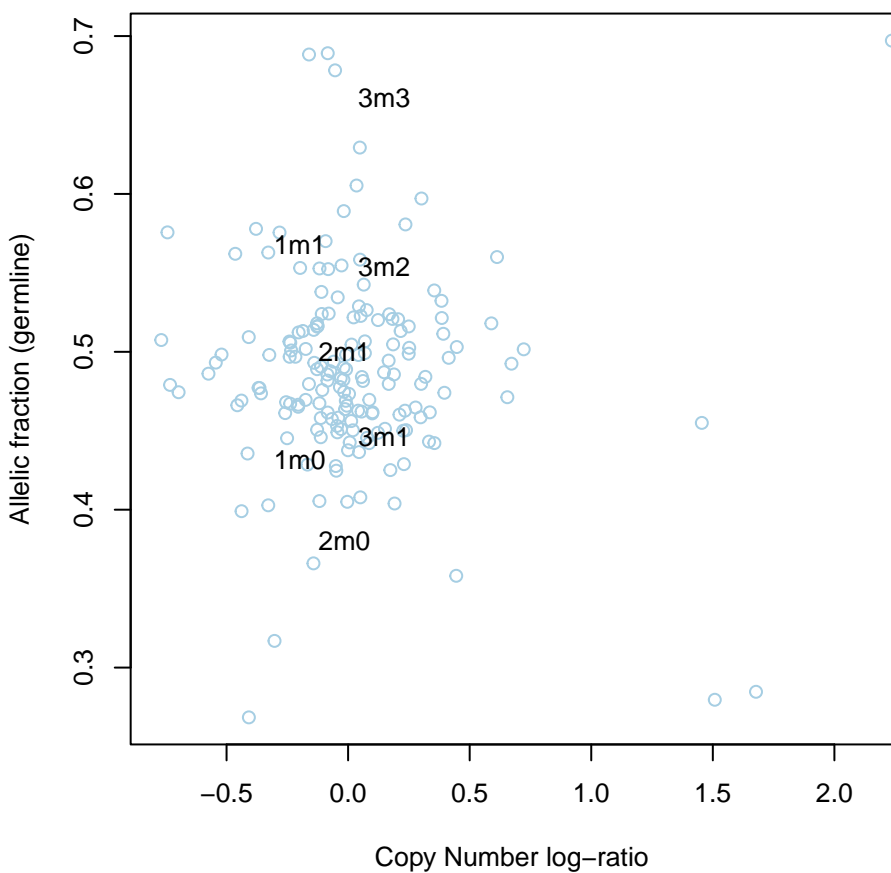


Purity: 0.24 Tumor ploidy: 2.087 SNV log-likelihood: -13.96 GoF: 92.2% Mean coverage: 584;392

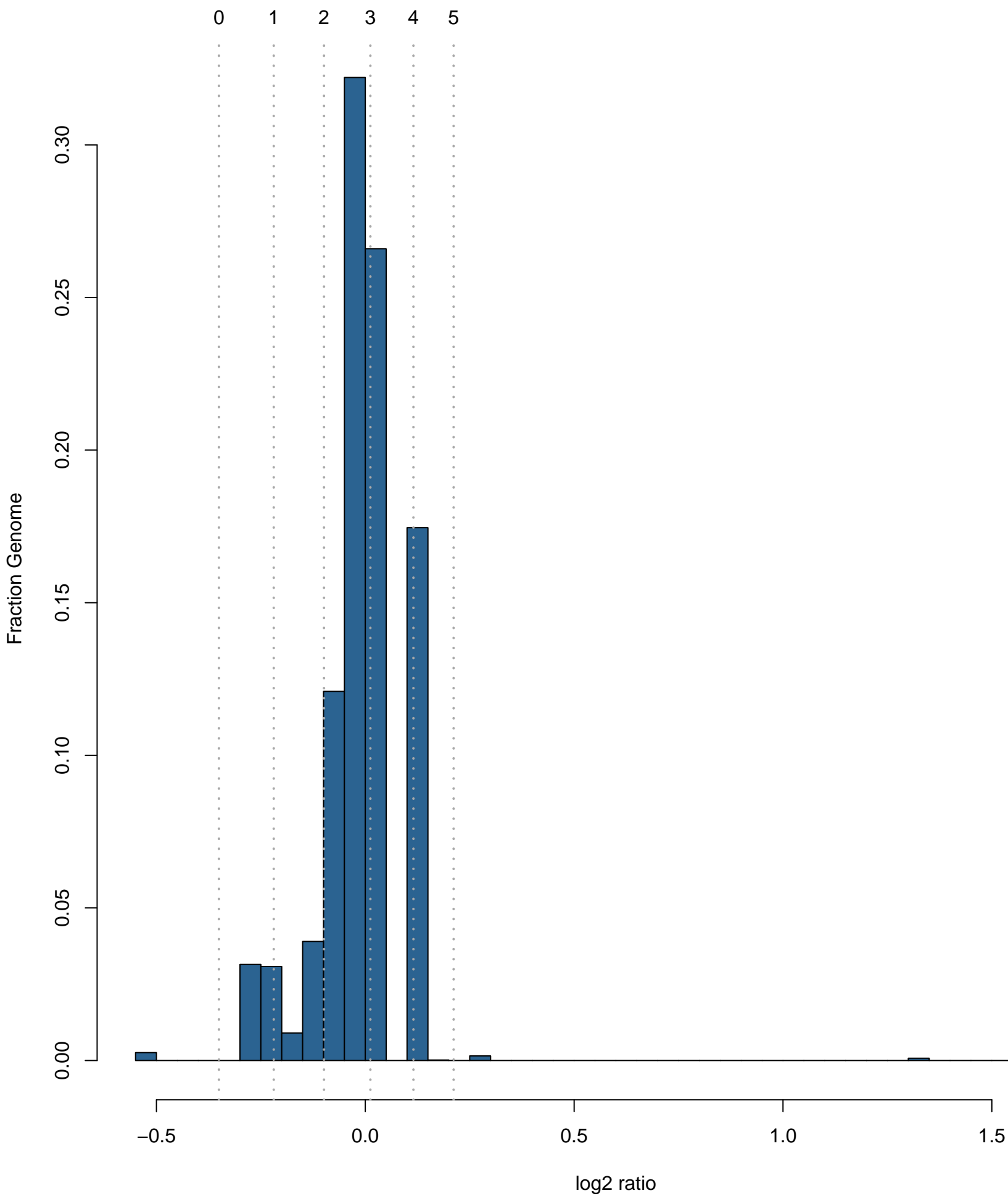


SCNA-fit log-likelihood: -8710.56

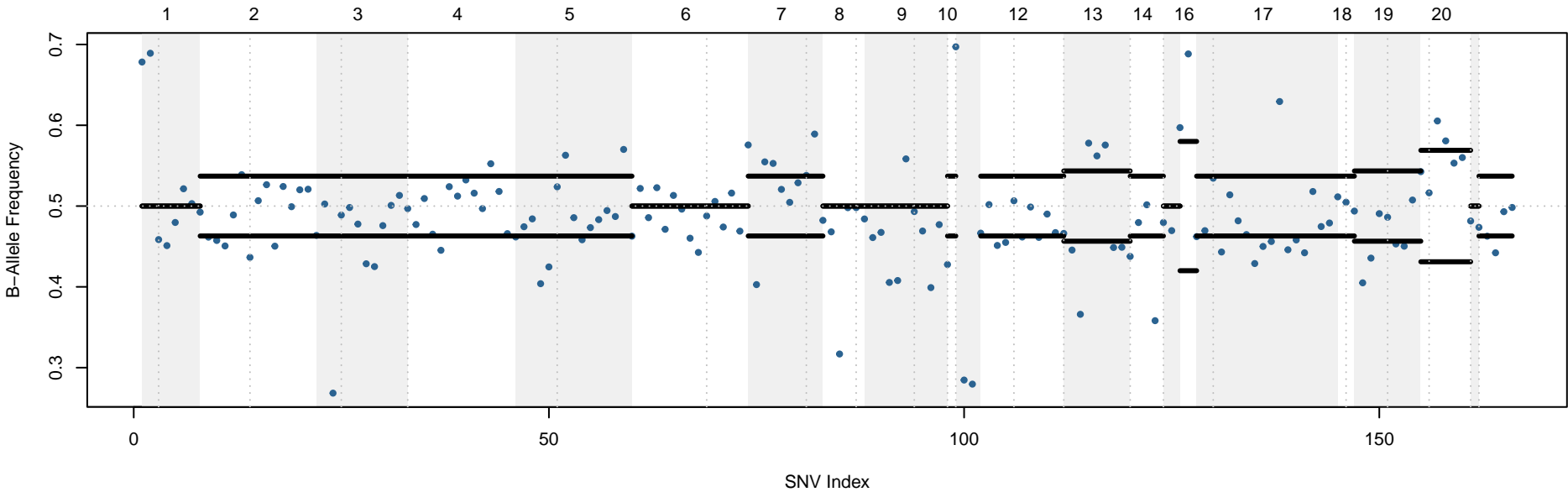




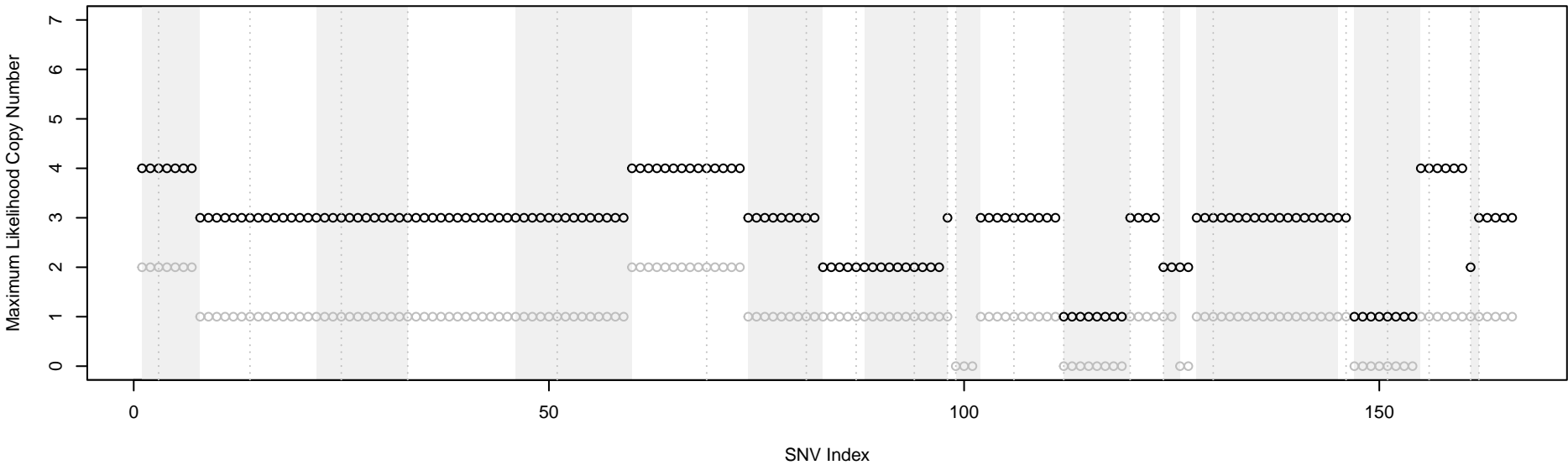
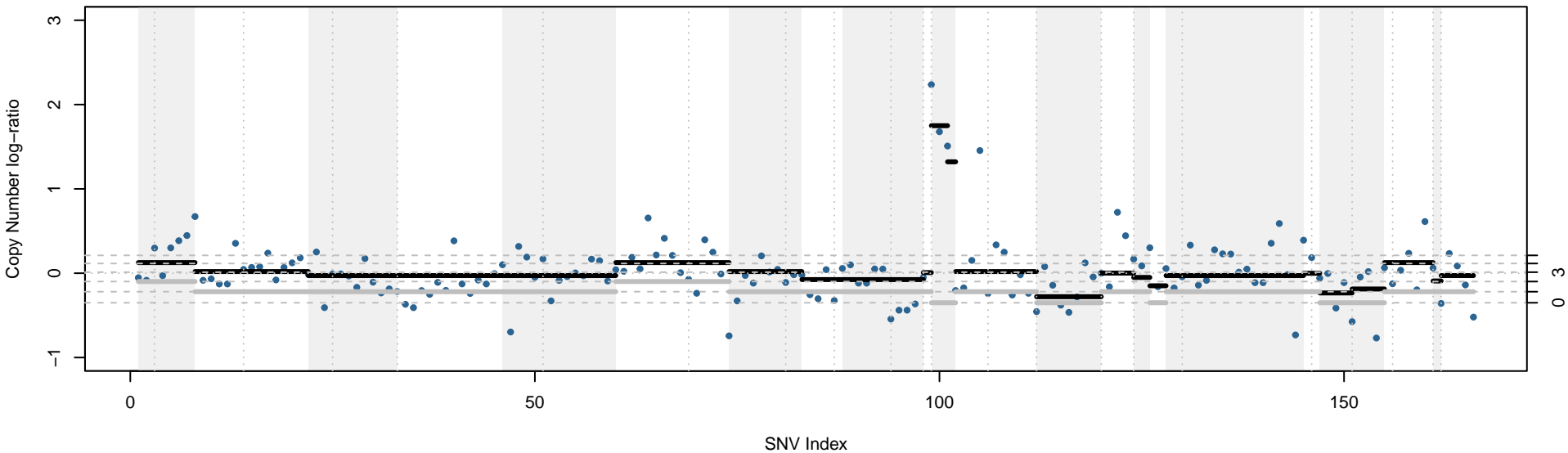
Purity: 0.16 Tumor ploidy: 2.886

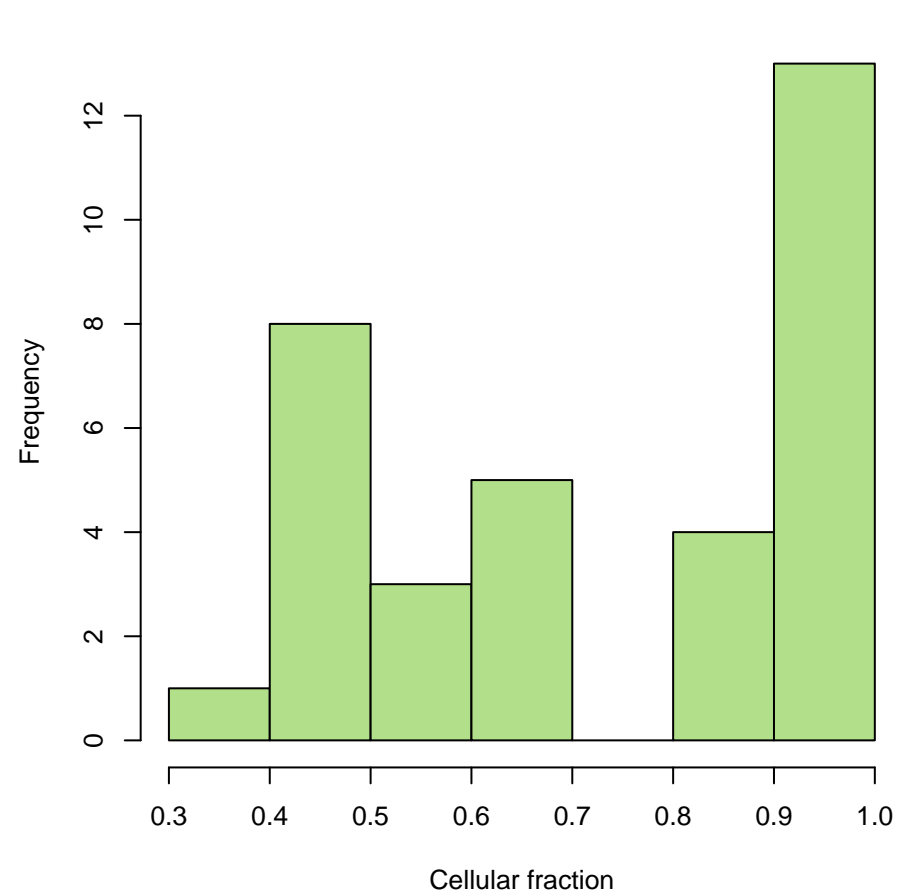
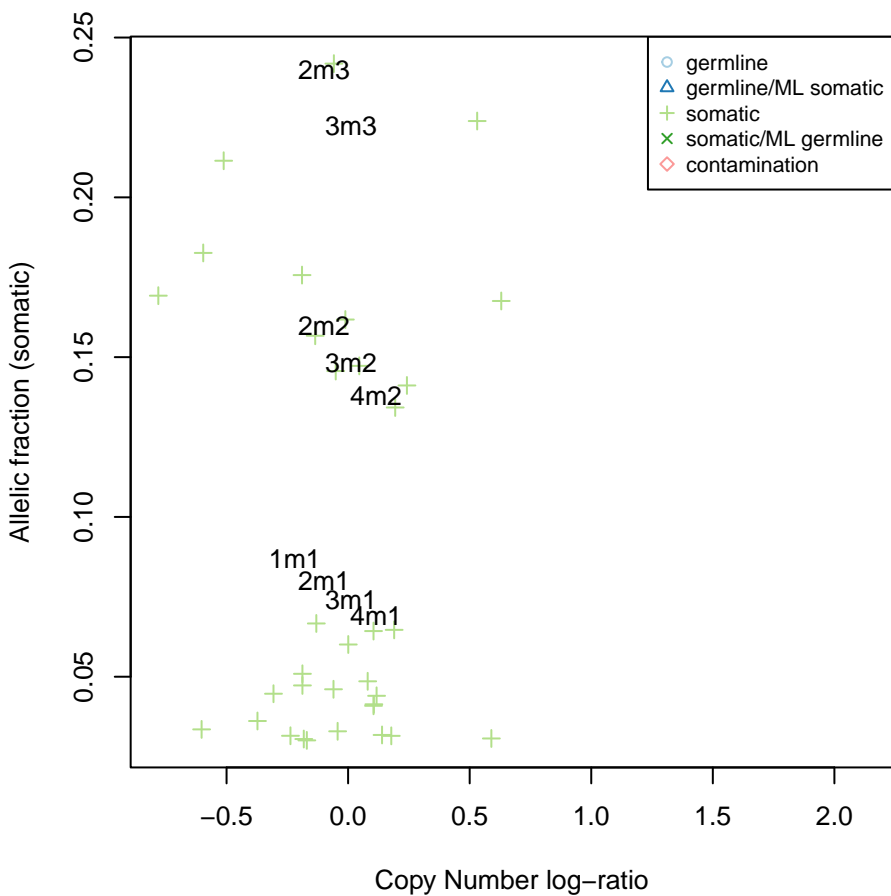
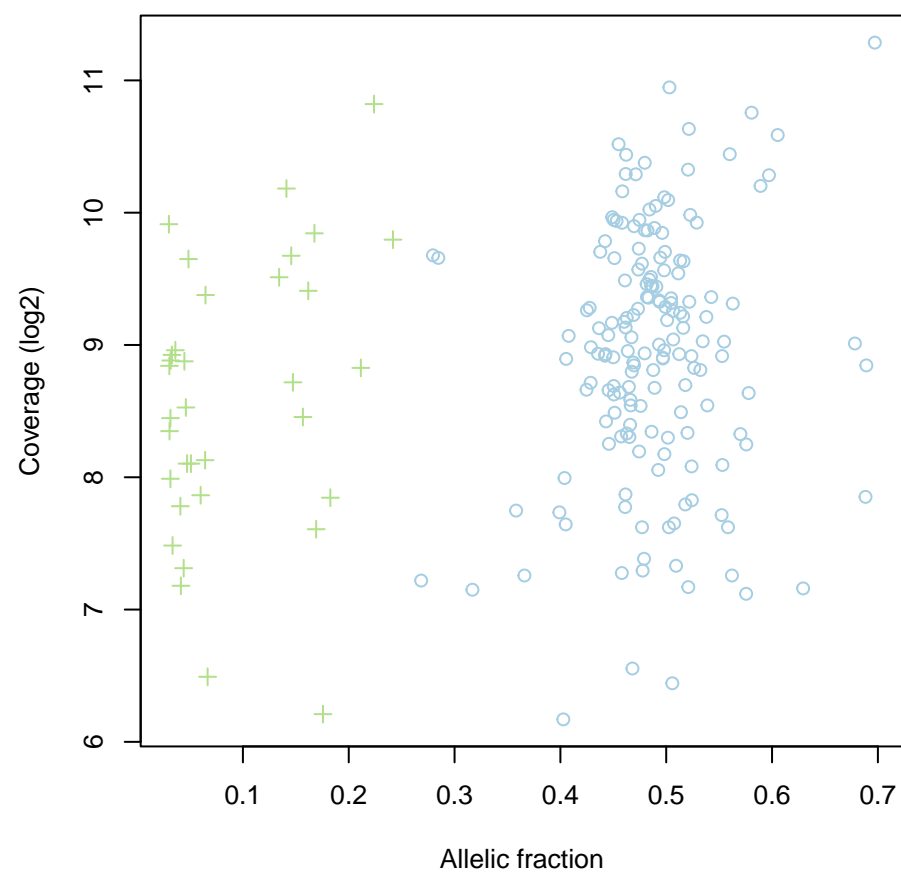
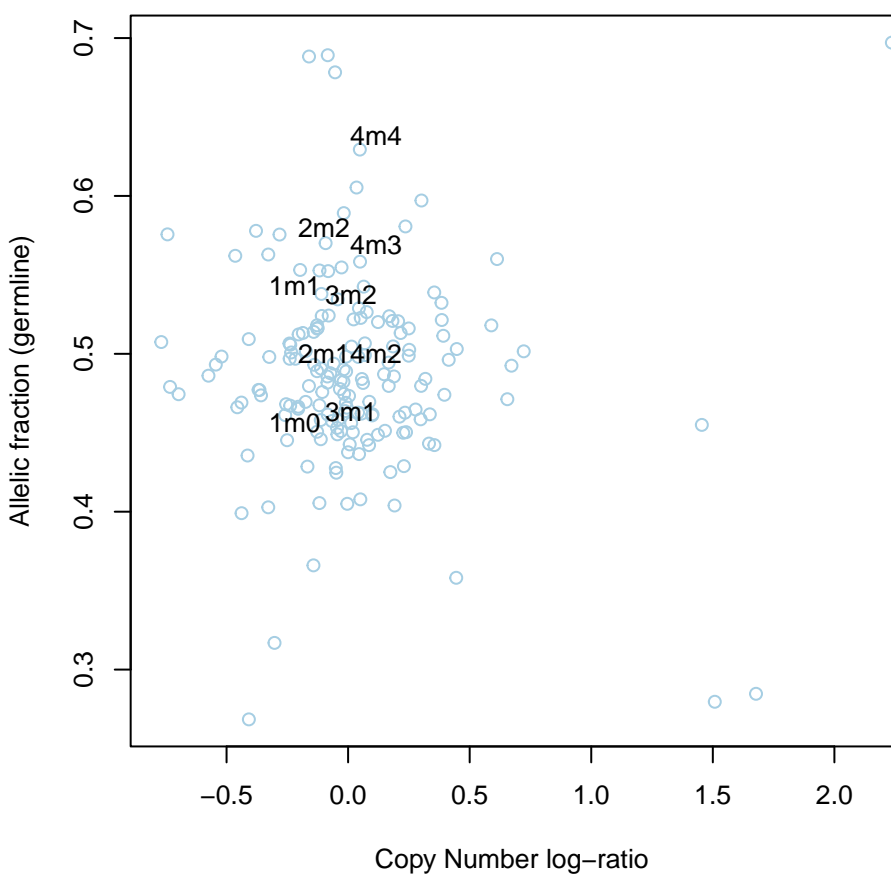


Purity: 0.16 Tumor ploidy: 2.886 SNV log-likelihood: -54.36 GoF: 94% Mean coverage: 584;392

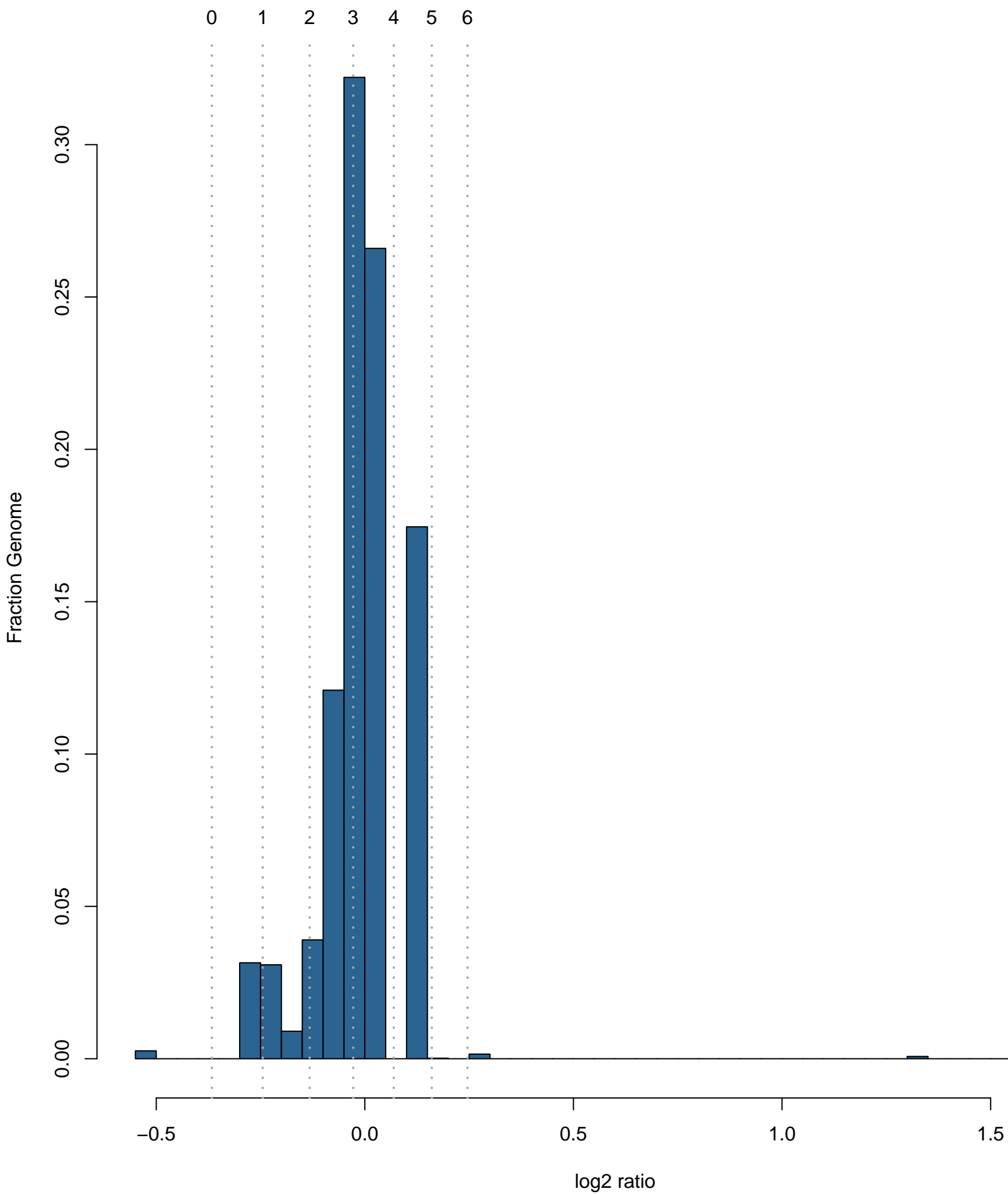


SCNA-fit log-likelihood: -8705.39

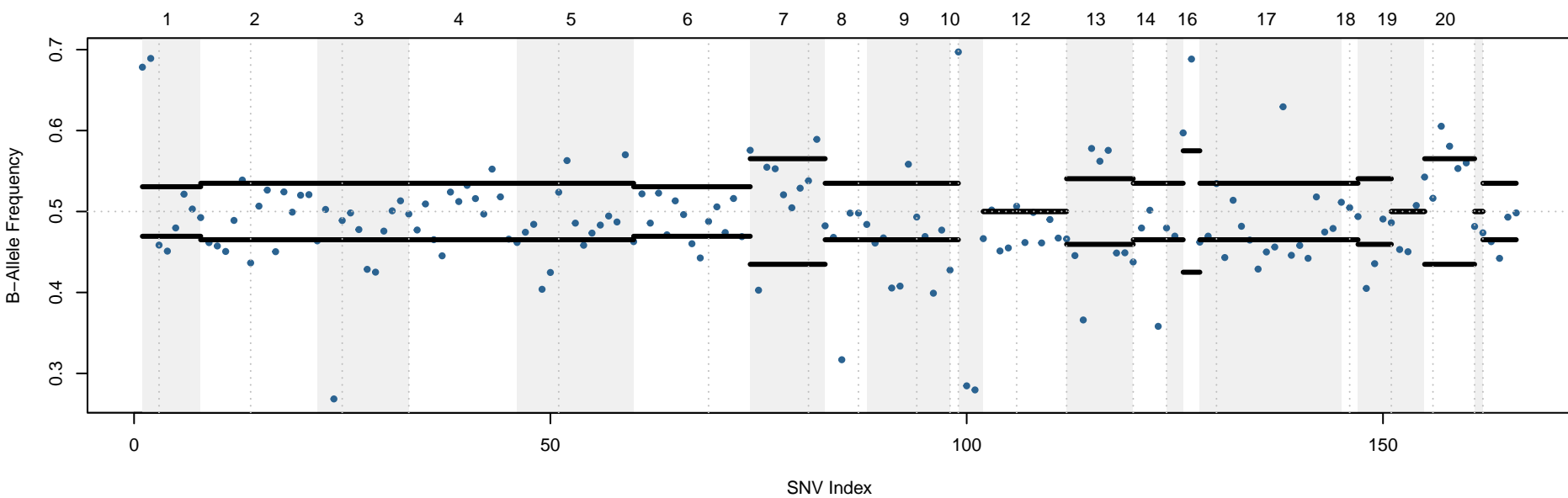




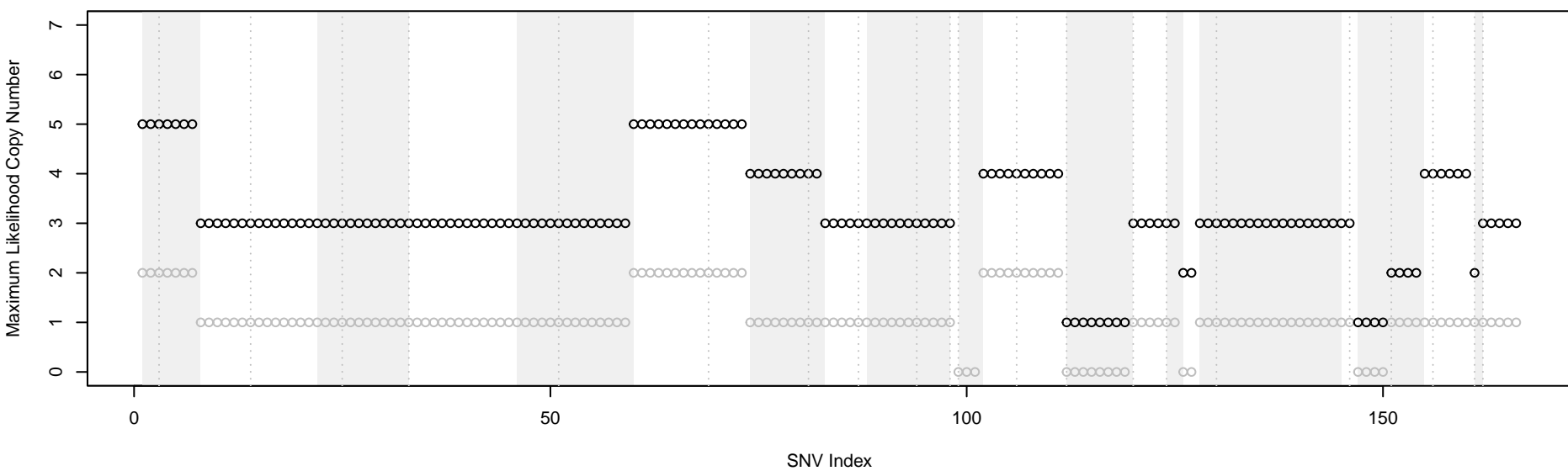
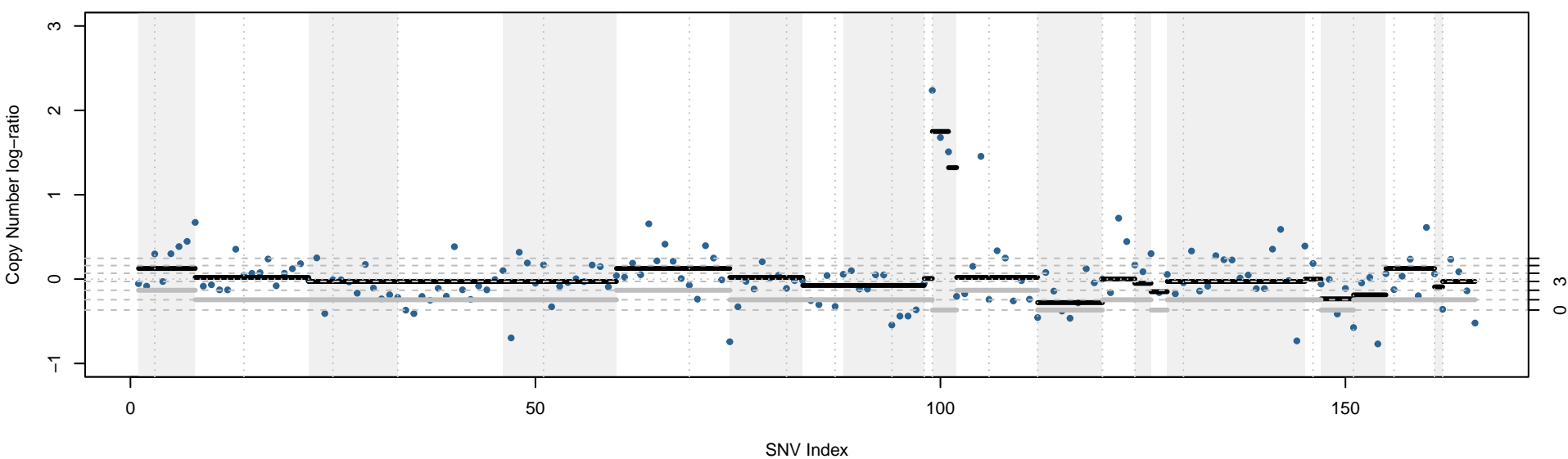
Purity: 0.15 Tumor ploidy: 3.28



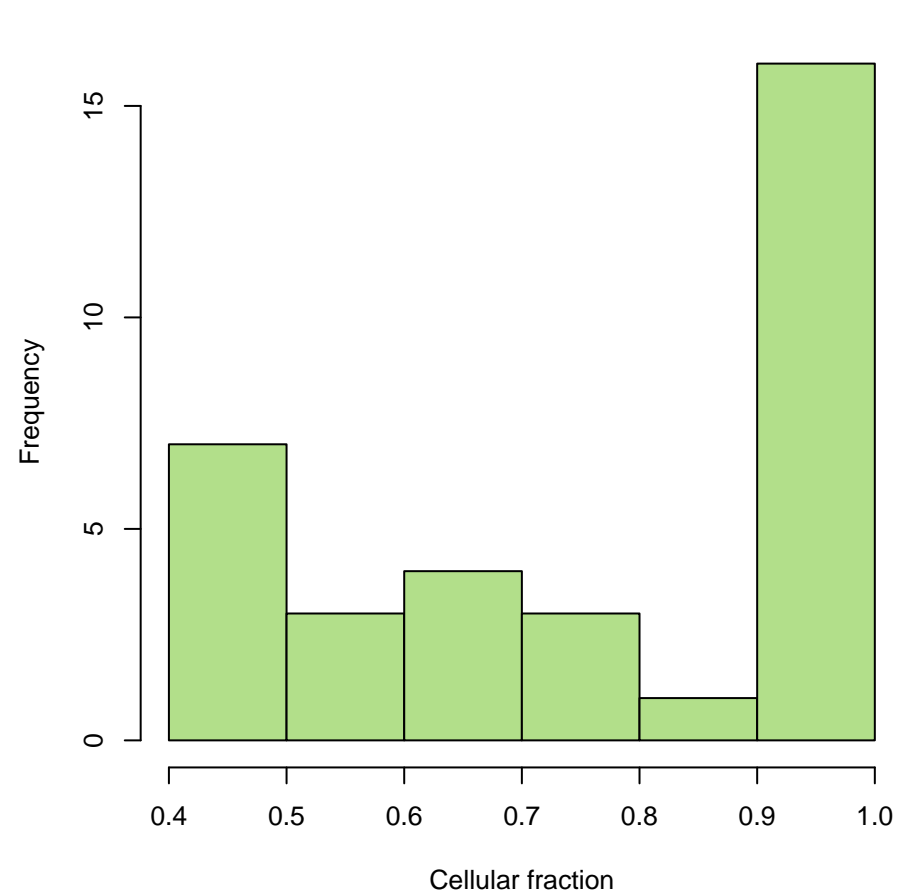
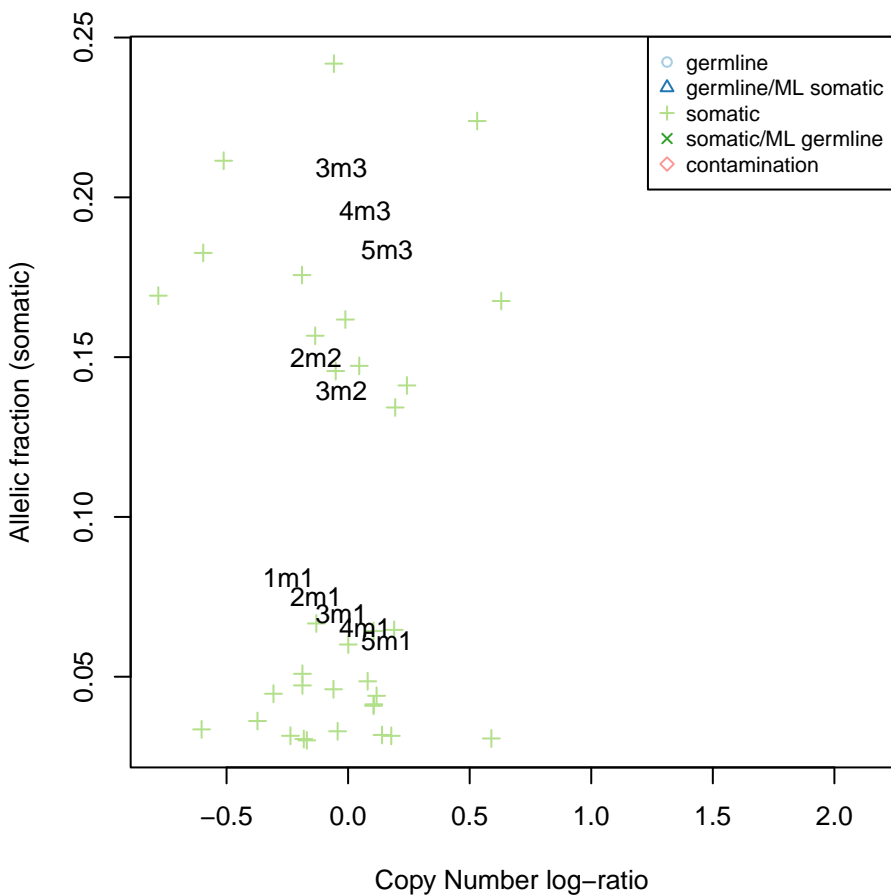
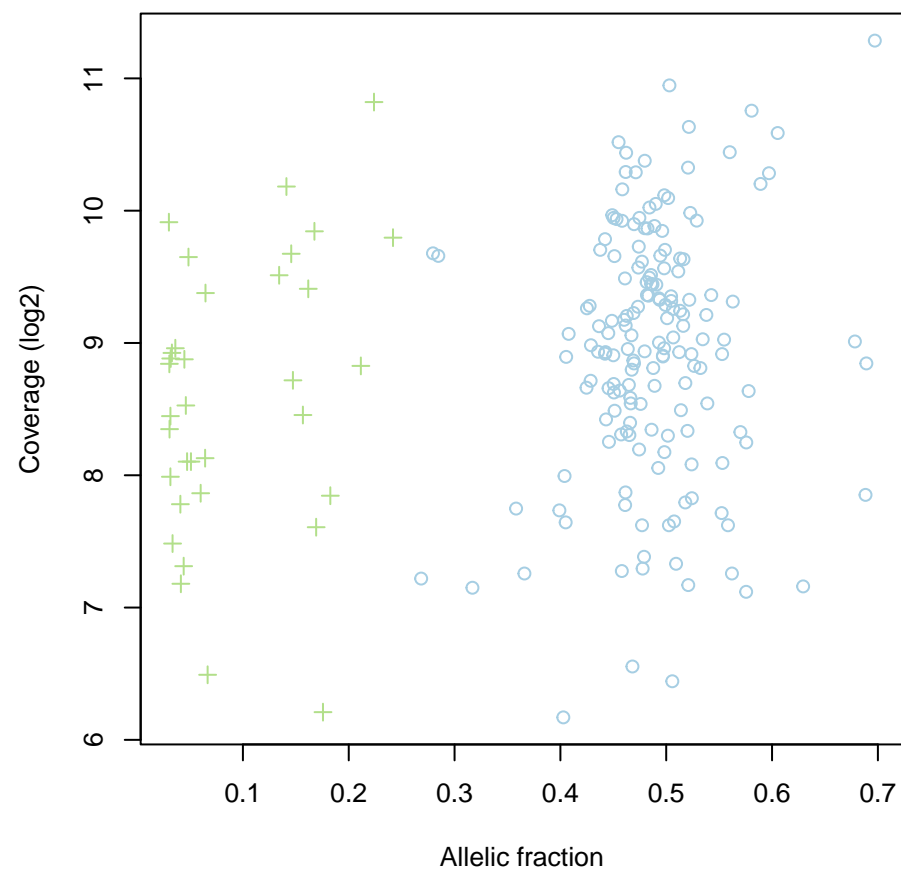
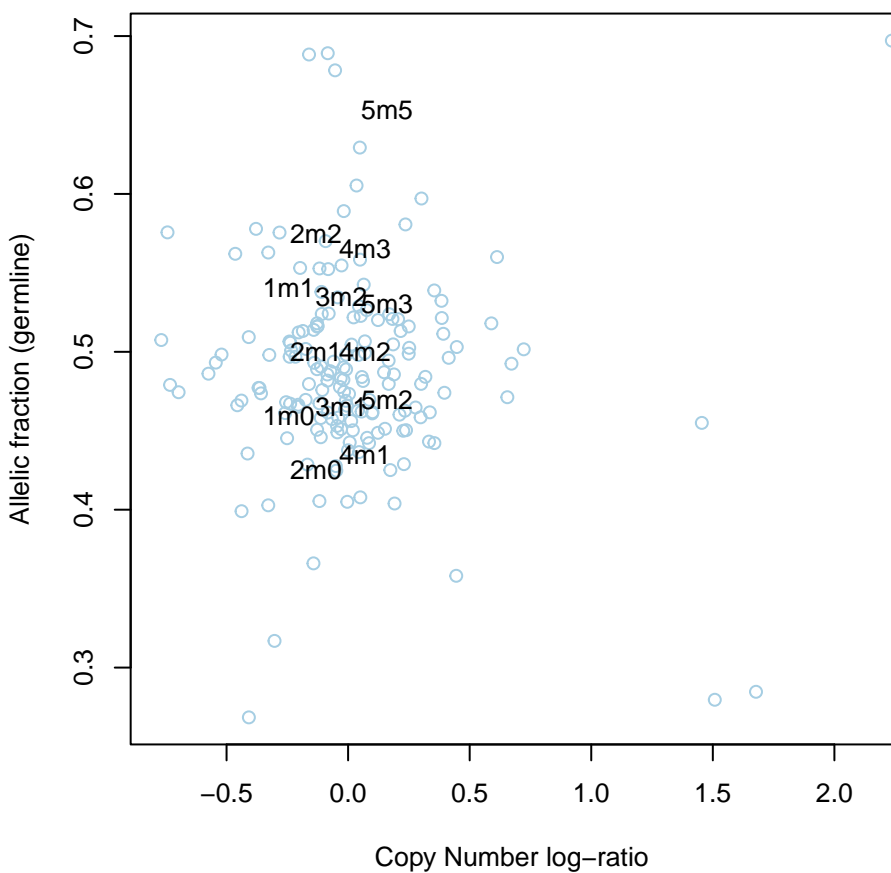
Purity: 0.15 Tumor ploidy: 3.28 SNV log-likelihood: -75.66 GoF: 94.4% Mean coverage: 584,392



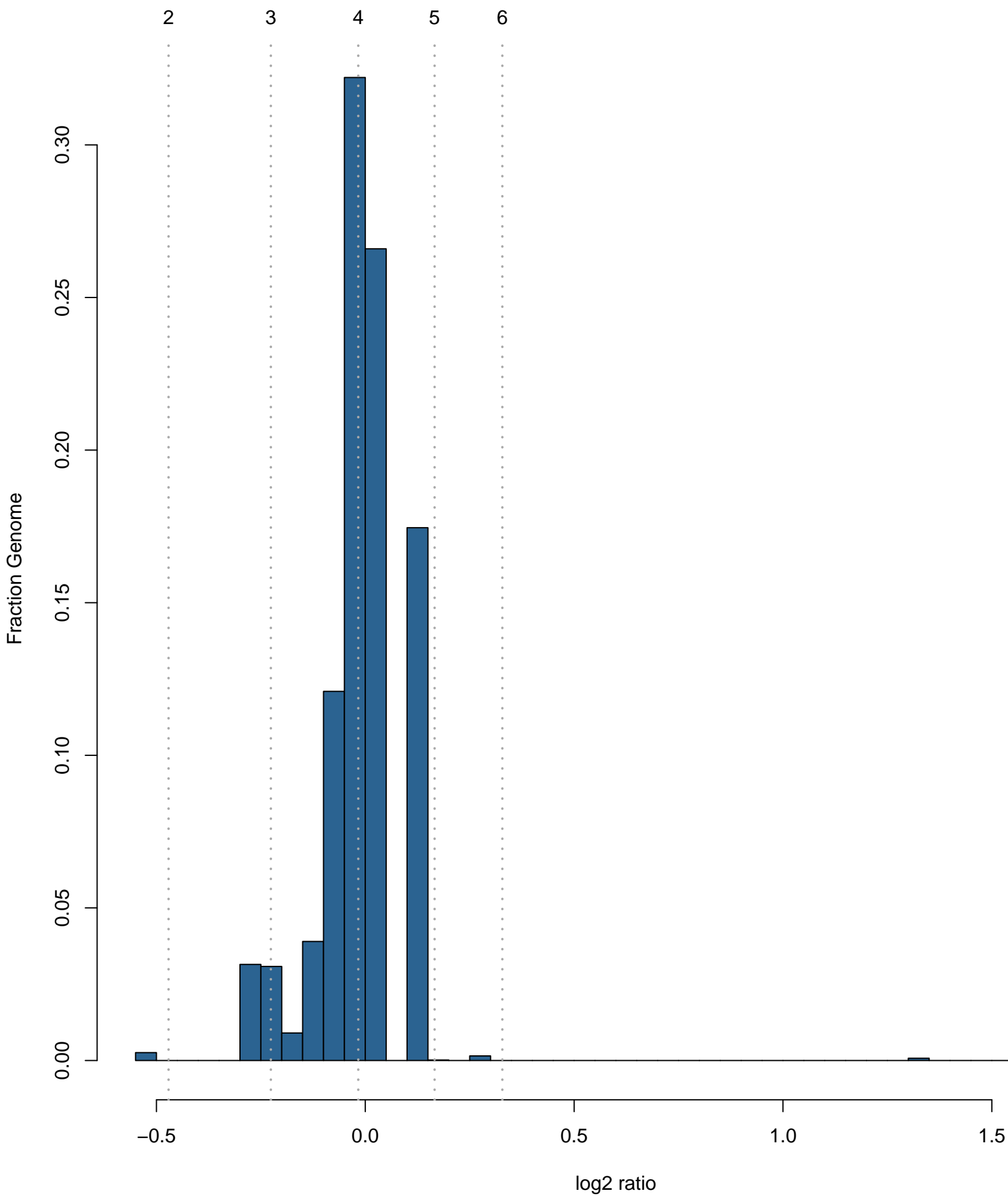
SCNA-fit log-likelihood: -8697.27



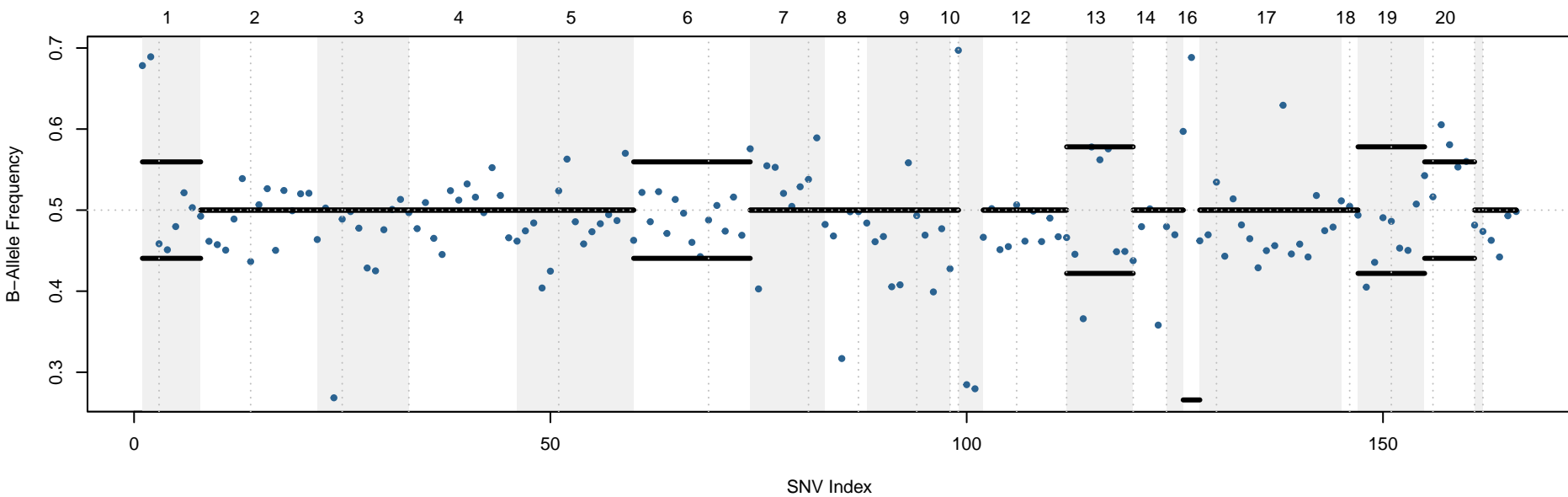




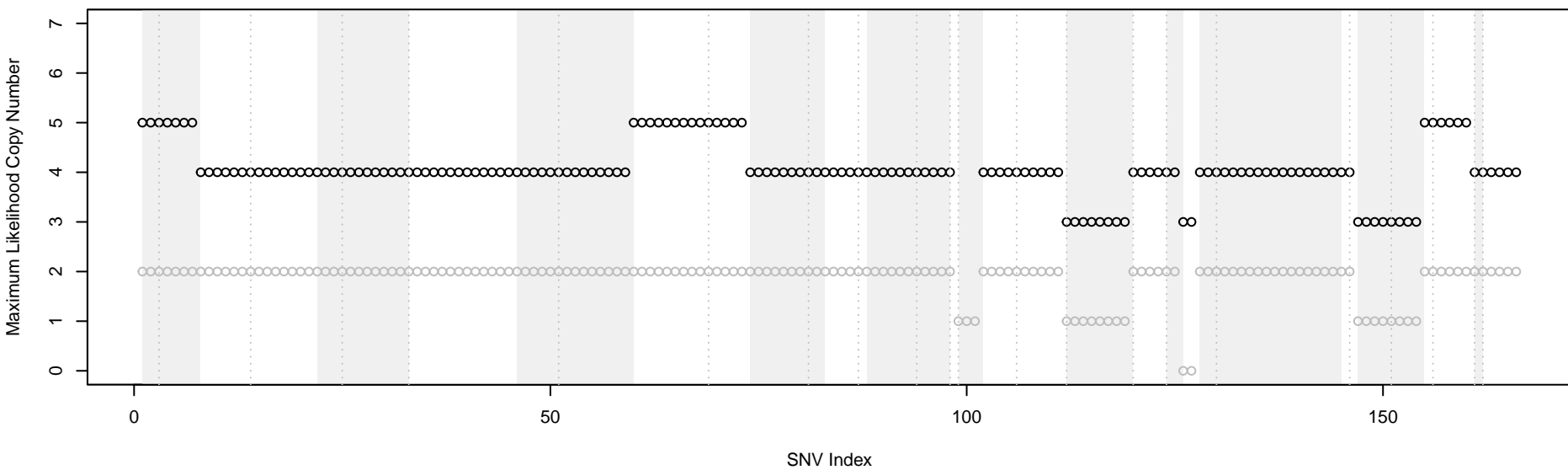
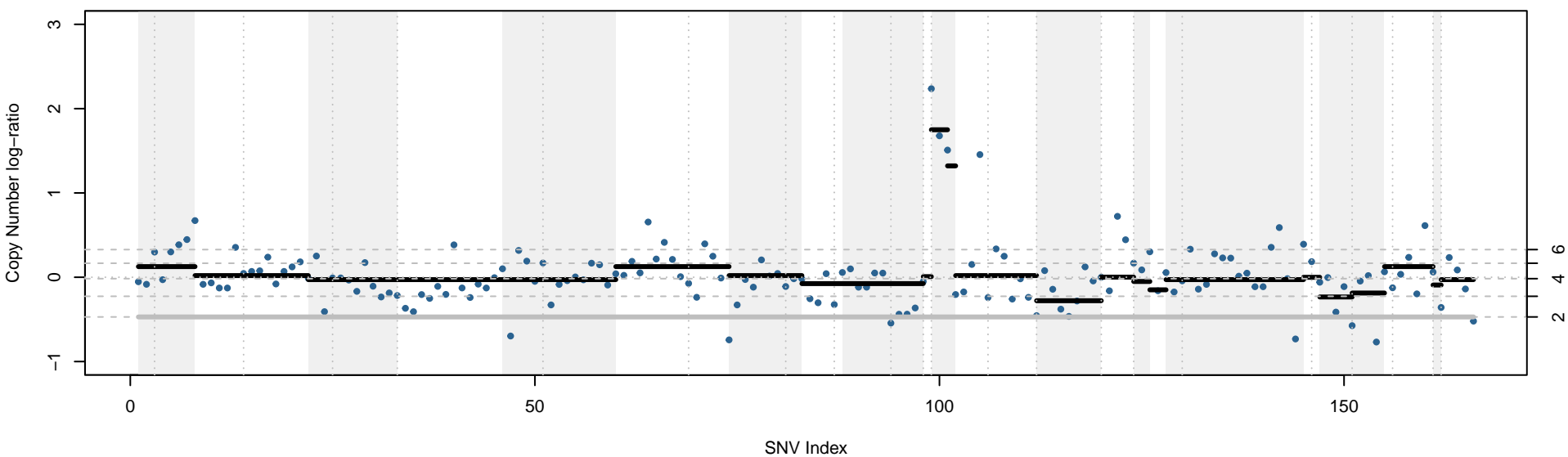
Purity: 0.37 Tumor ploidy: 4.087

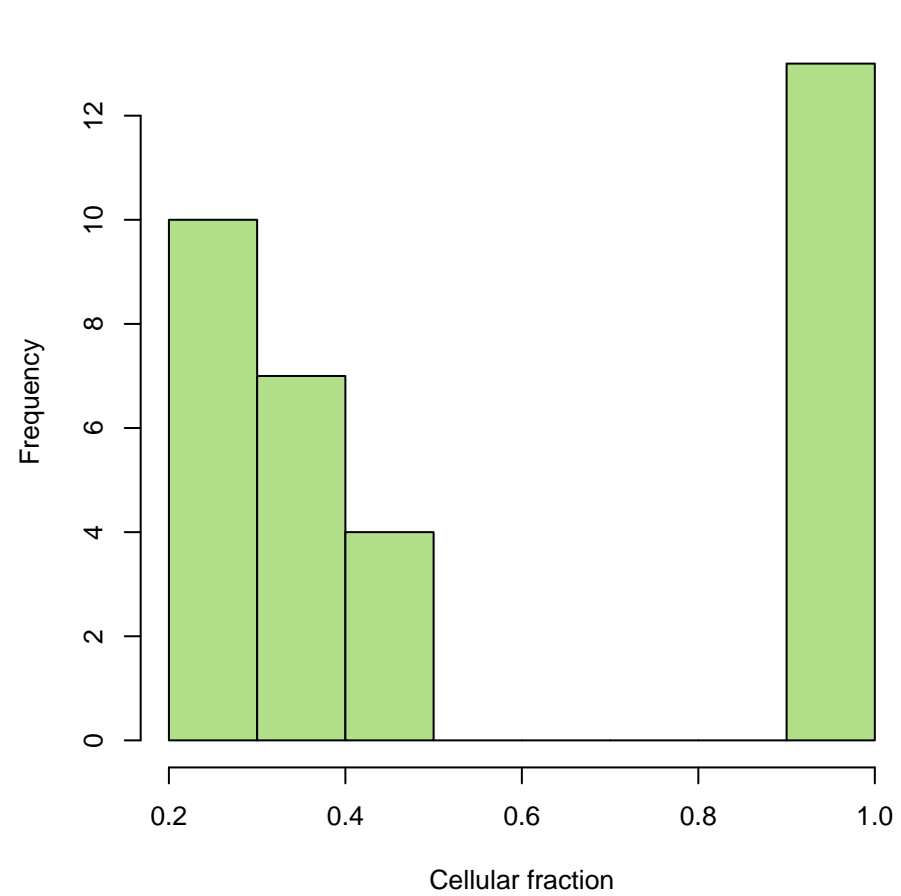
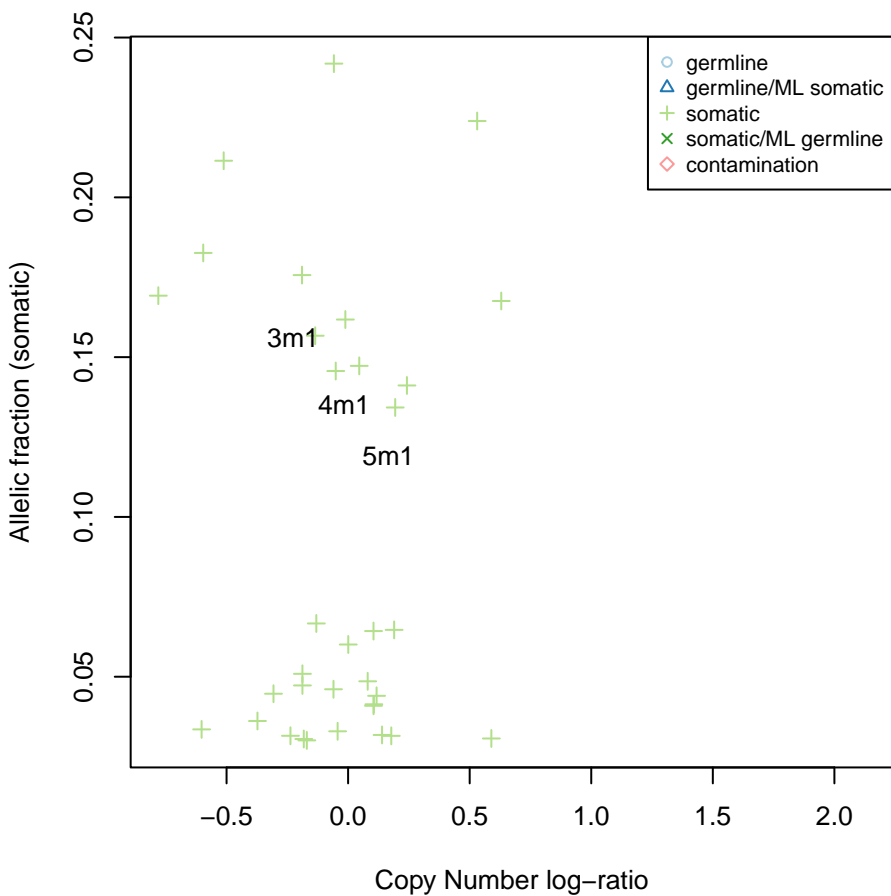
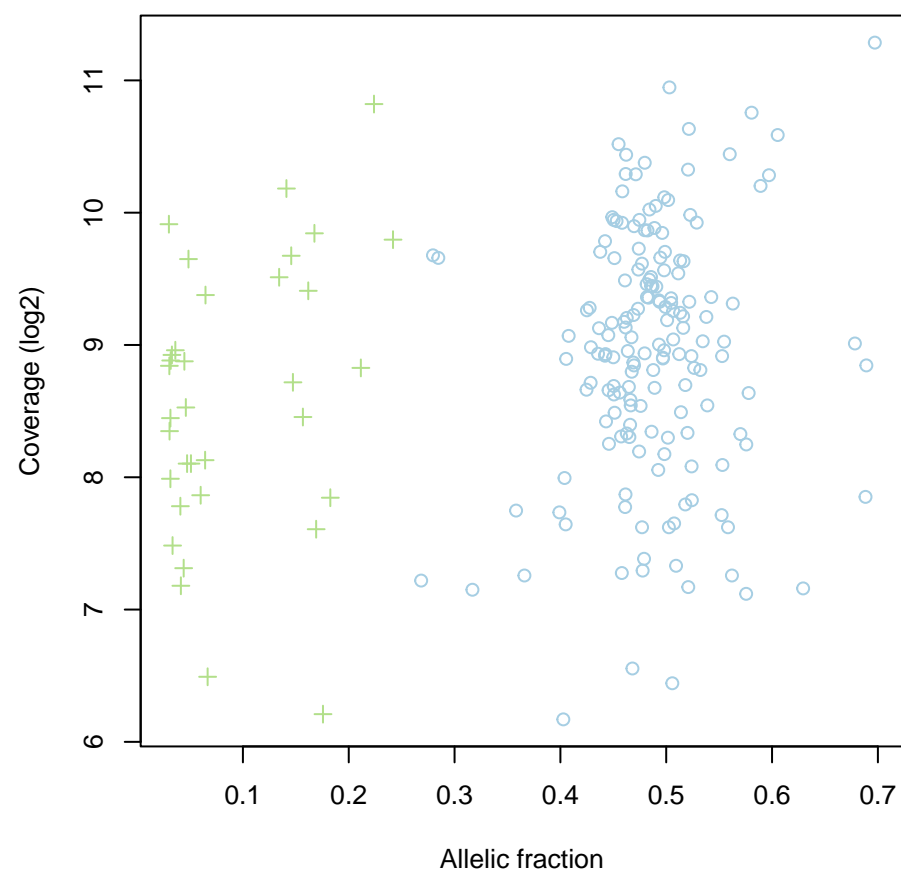
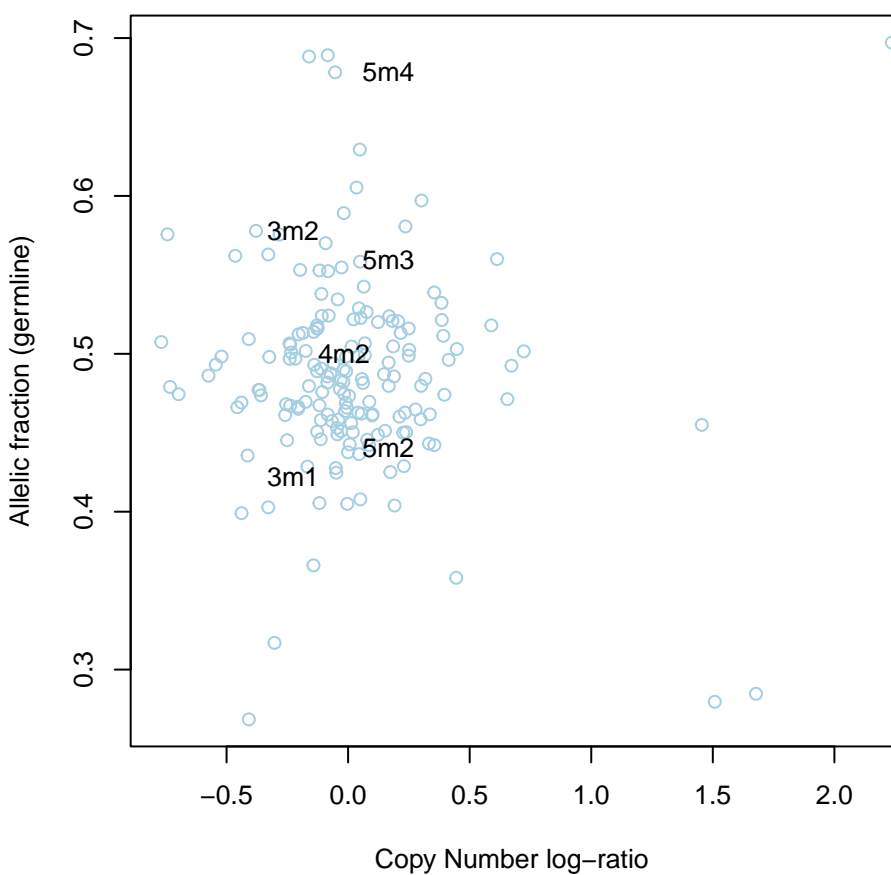


Purity: 0.37 Tumor ploidy: 4.087 SNV log-likelihood: -111.23 GoF: 91.5% Mean coverage: 584,392

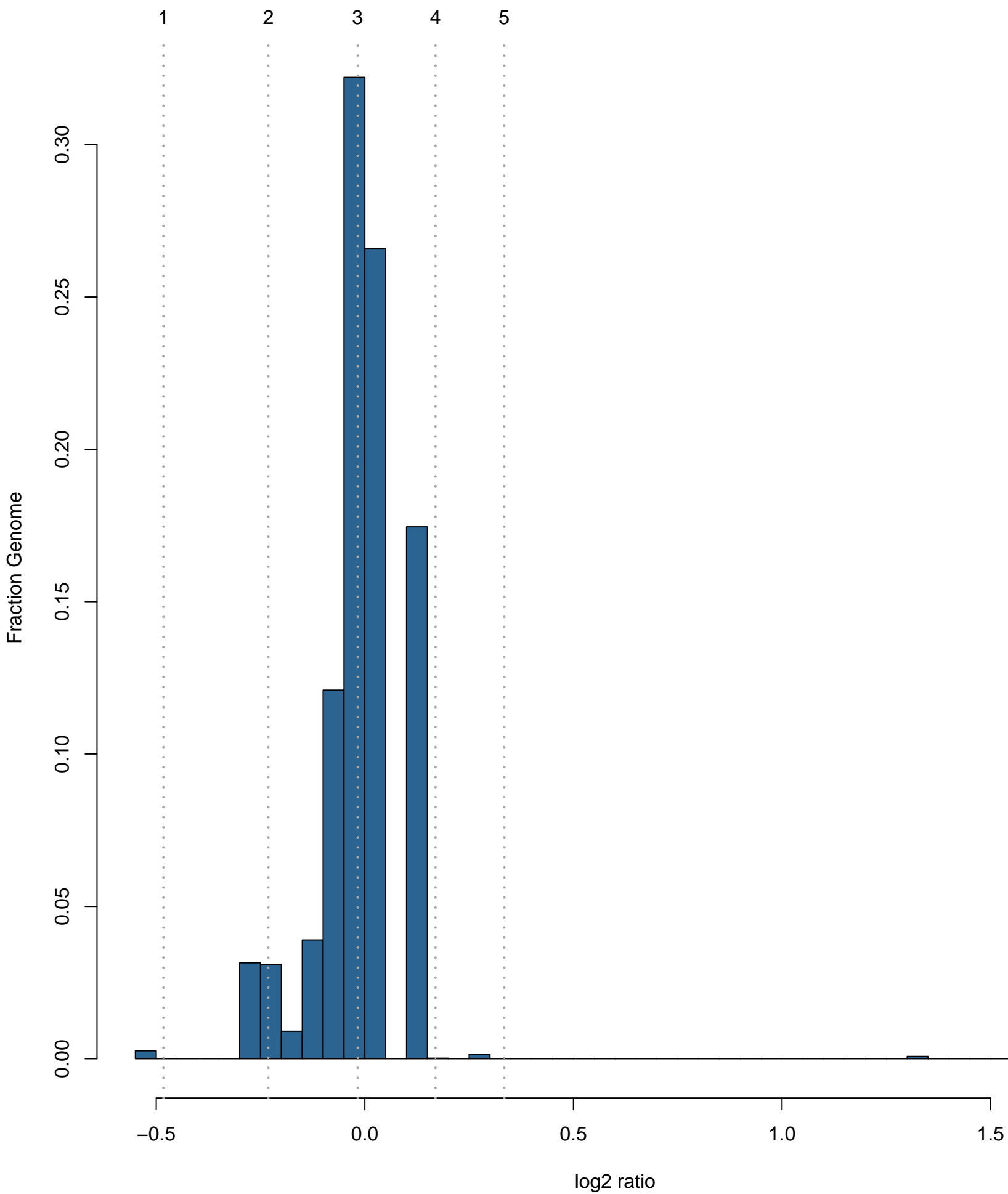


SCNA-fit log-likelihood: -8714.57

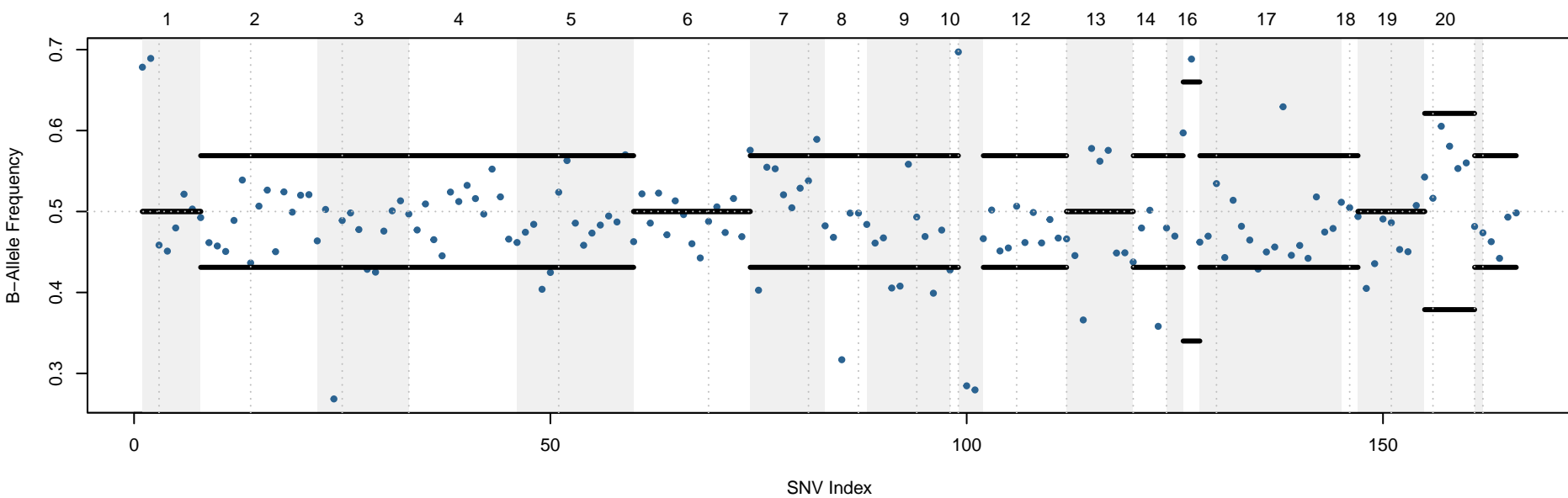




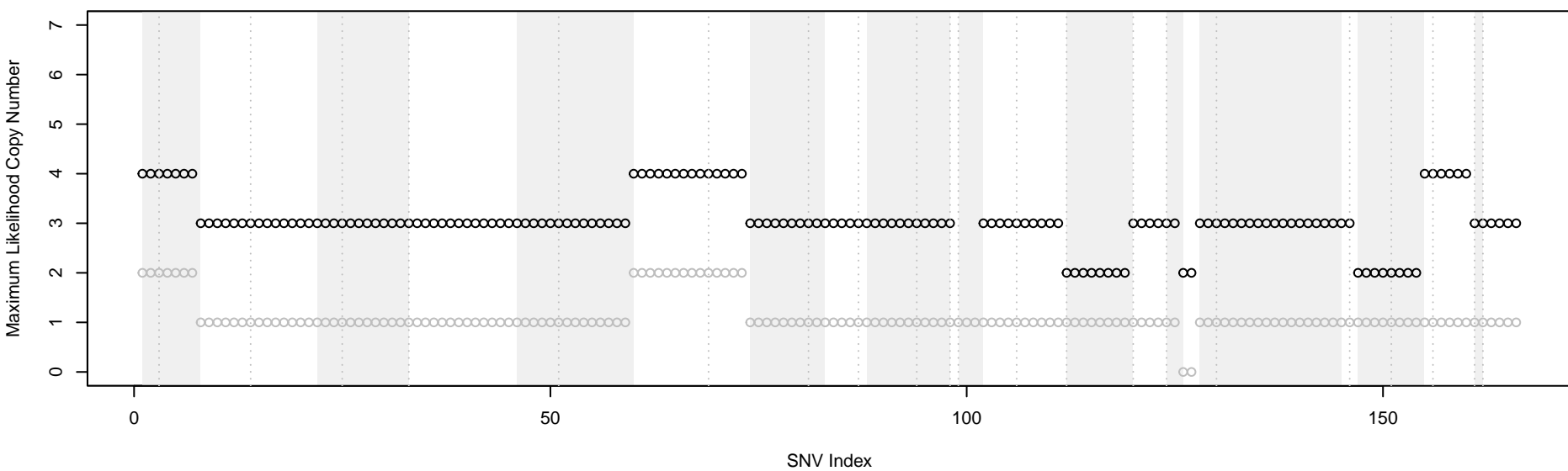
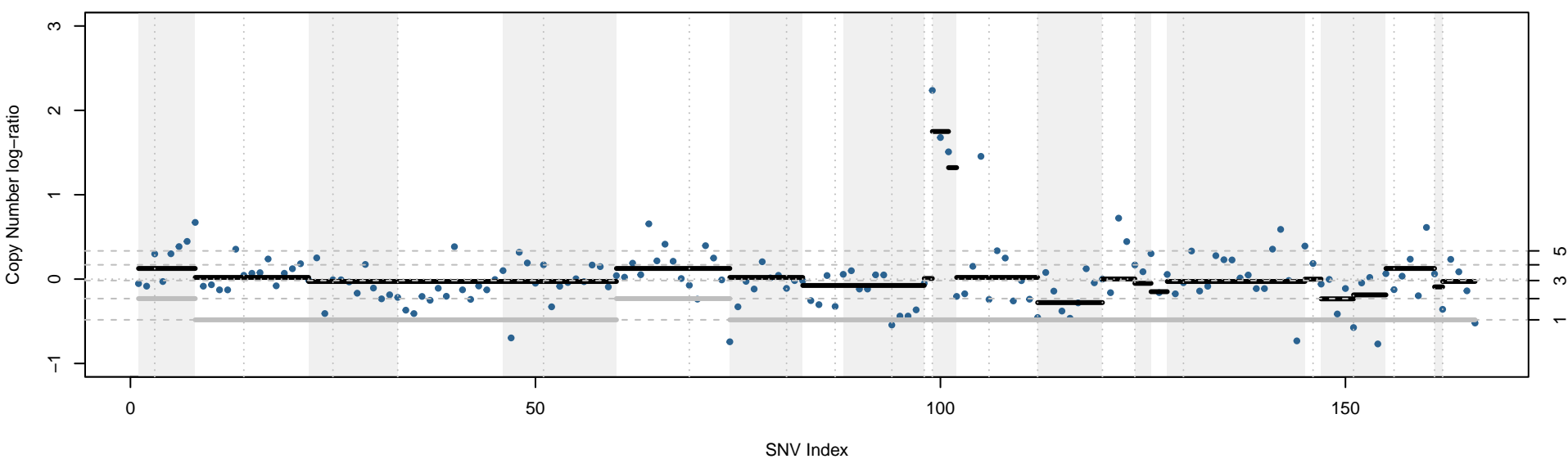
Purity: 0.32 Tumor ploidy: 3.087

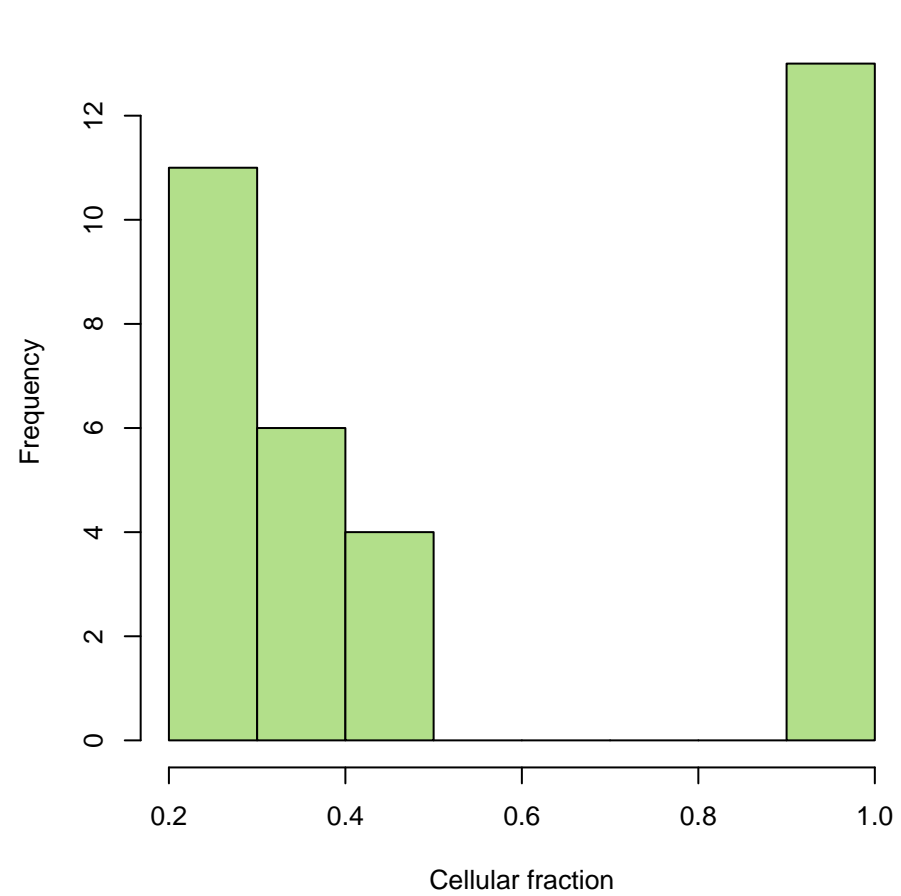
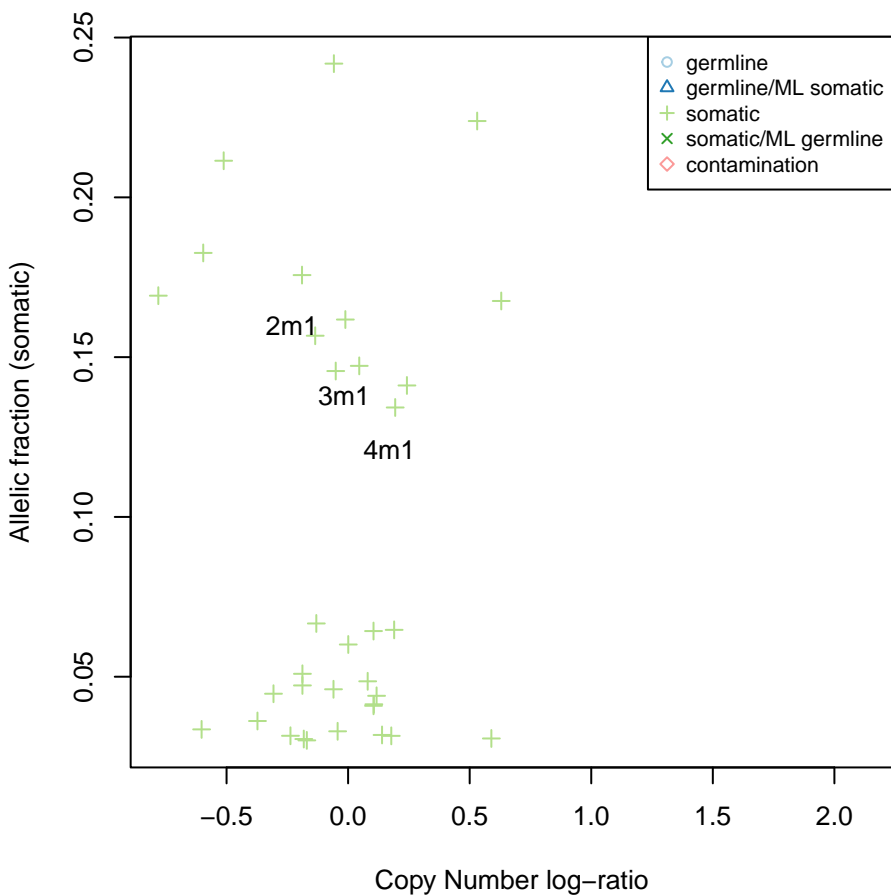
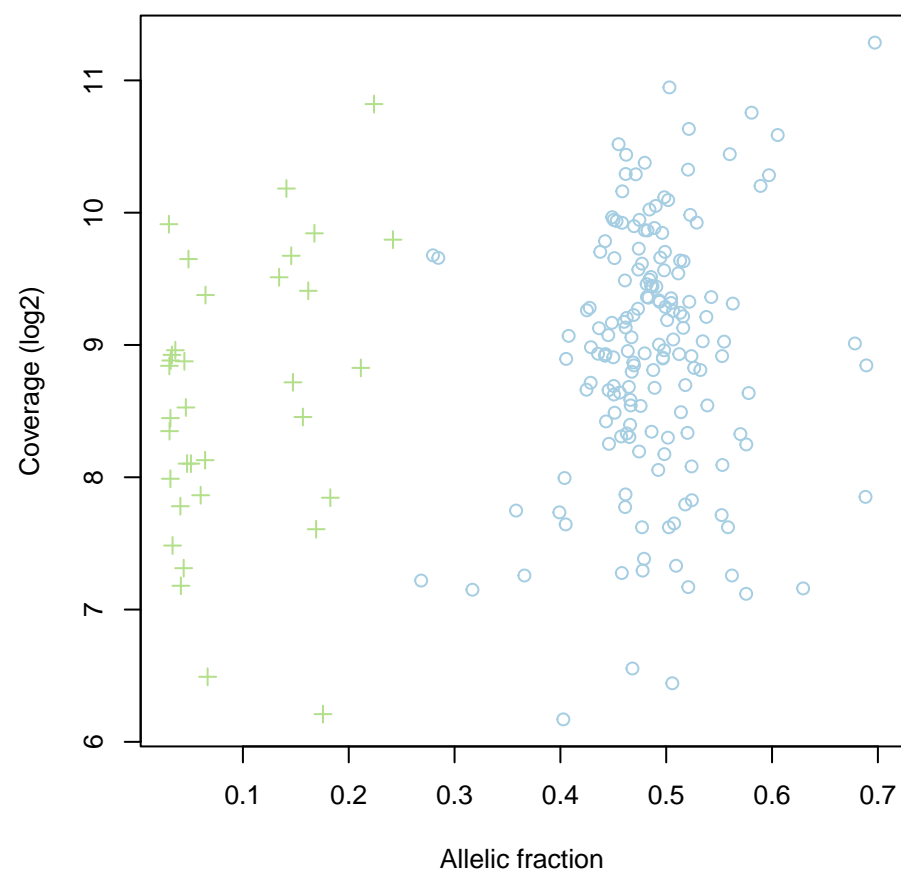
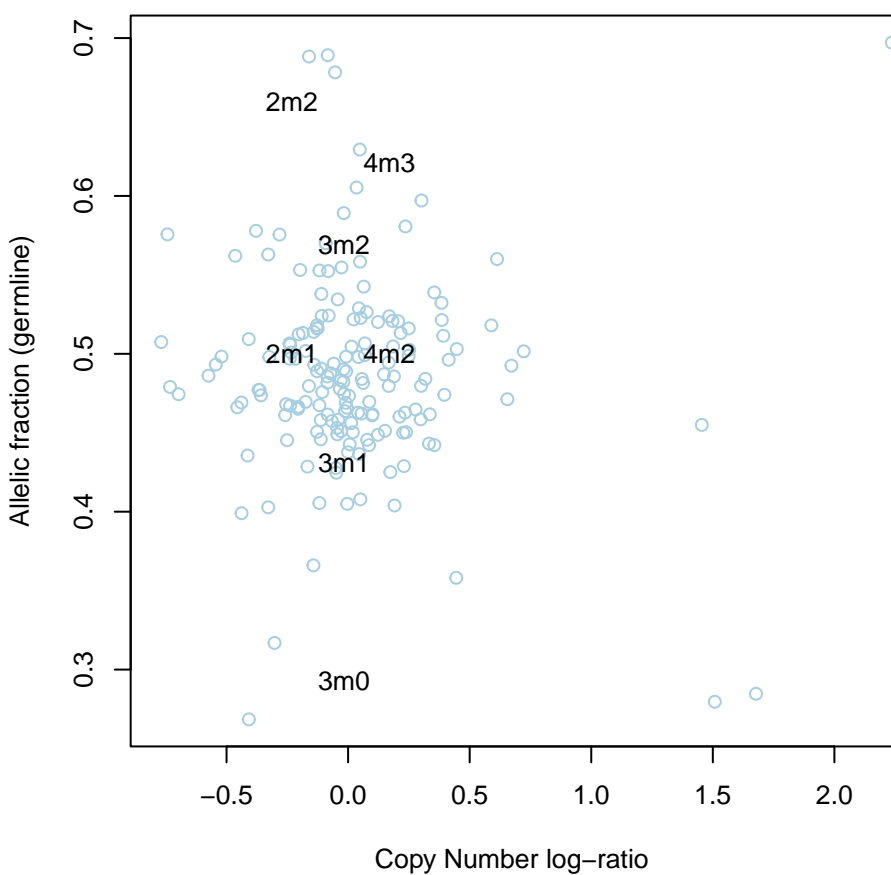


Purity: 0.32 Tumor ploidy: 3.087 SNV log-likelihood: -123.6 GoF: 91.1% Mean coverage: 584;392

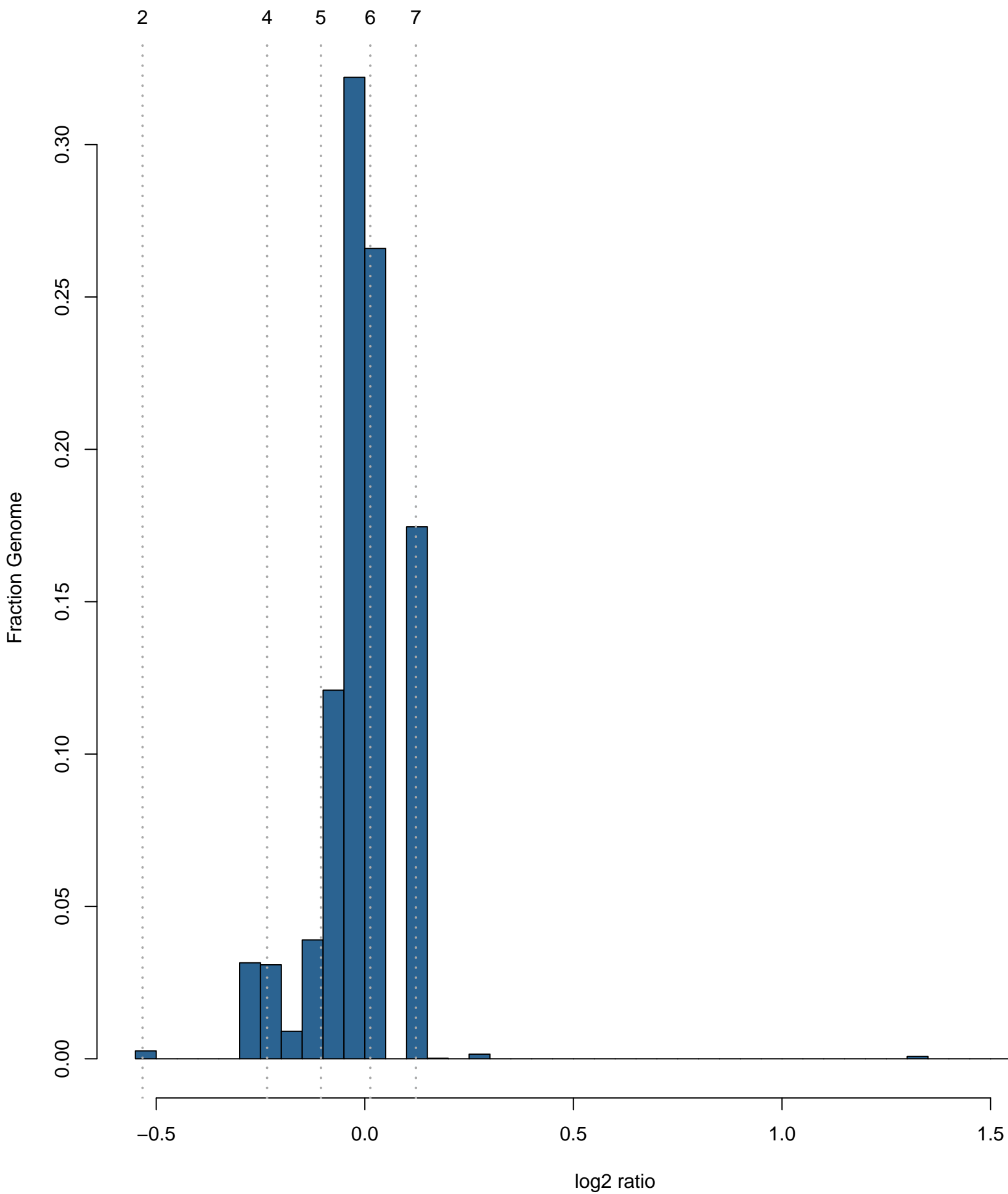


SCNA-fit log-likelihood: -8716.22



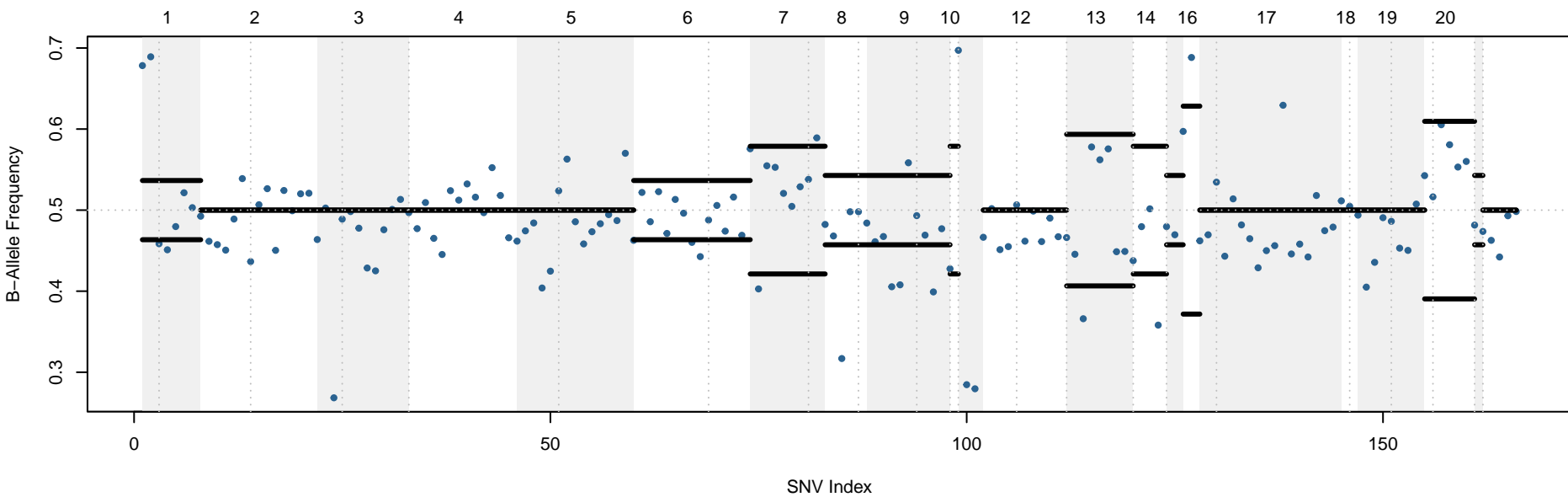


Purity: 0.23 Tumor ploidy: 5.885

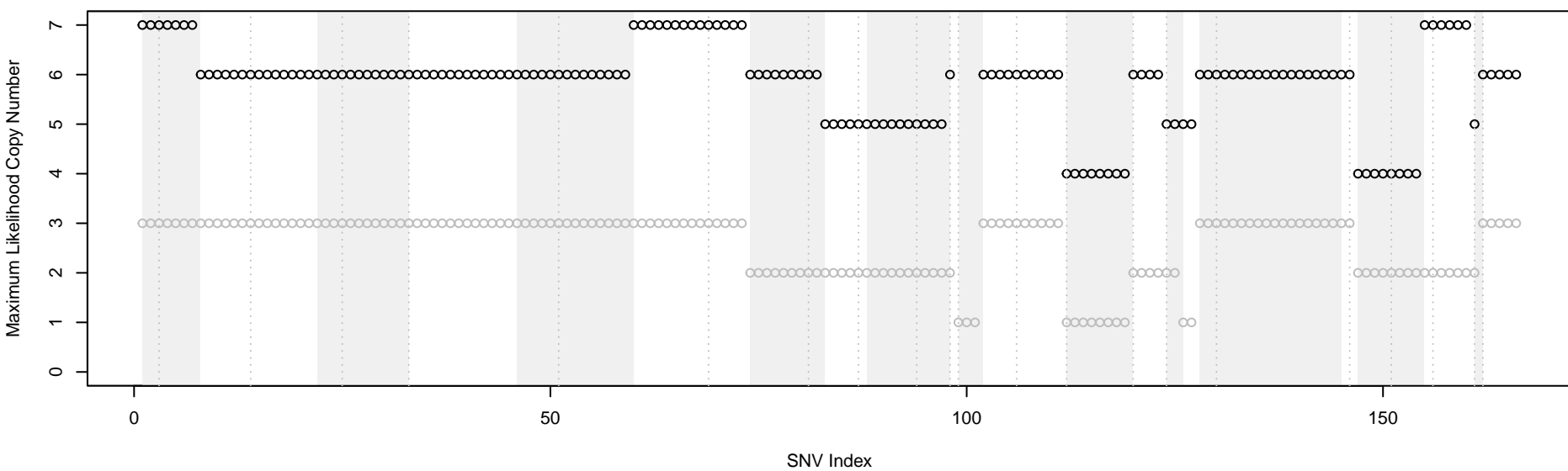
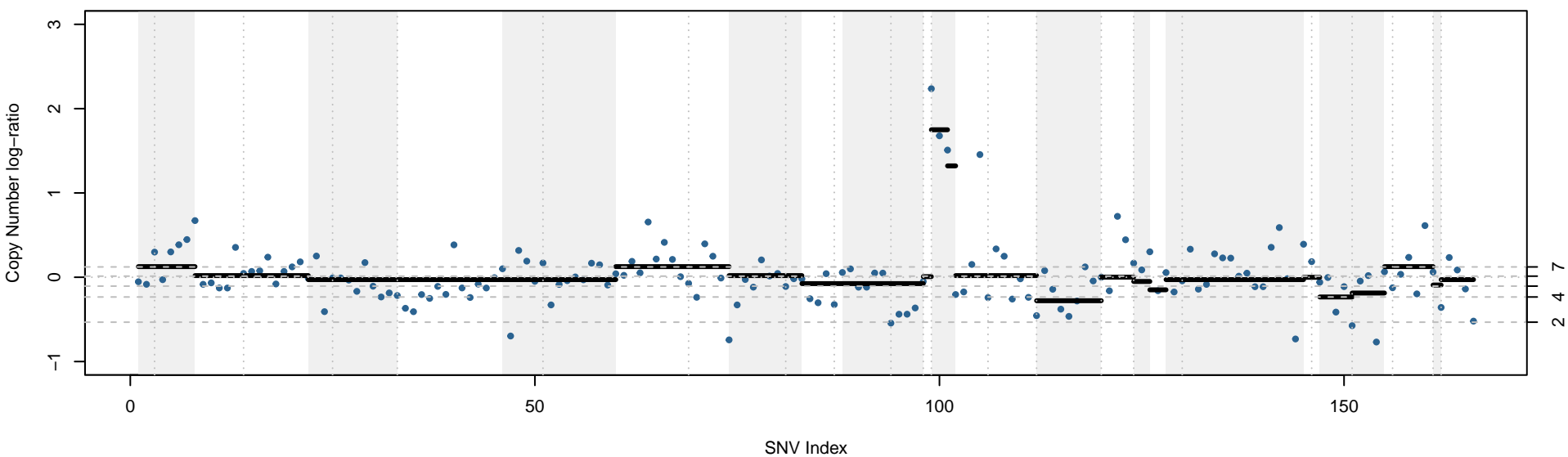


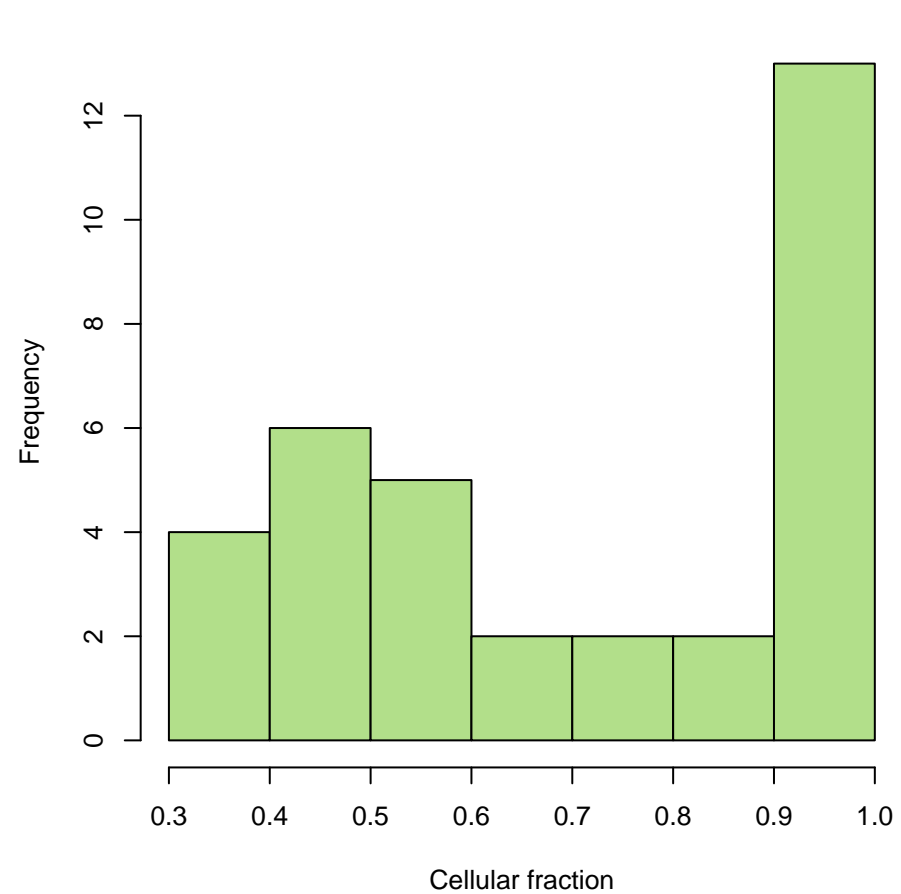
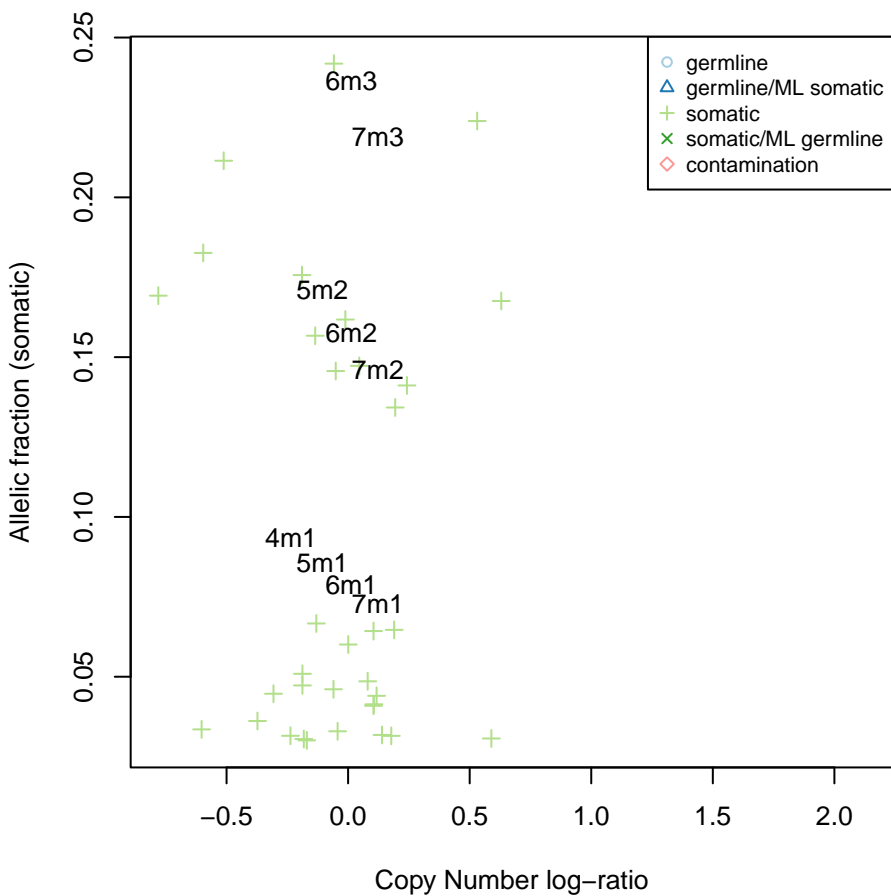
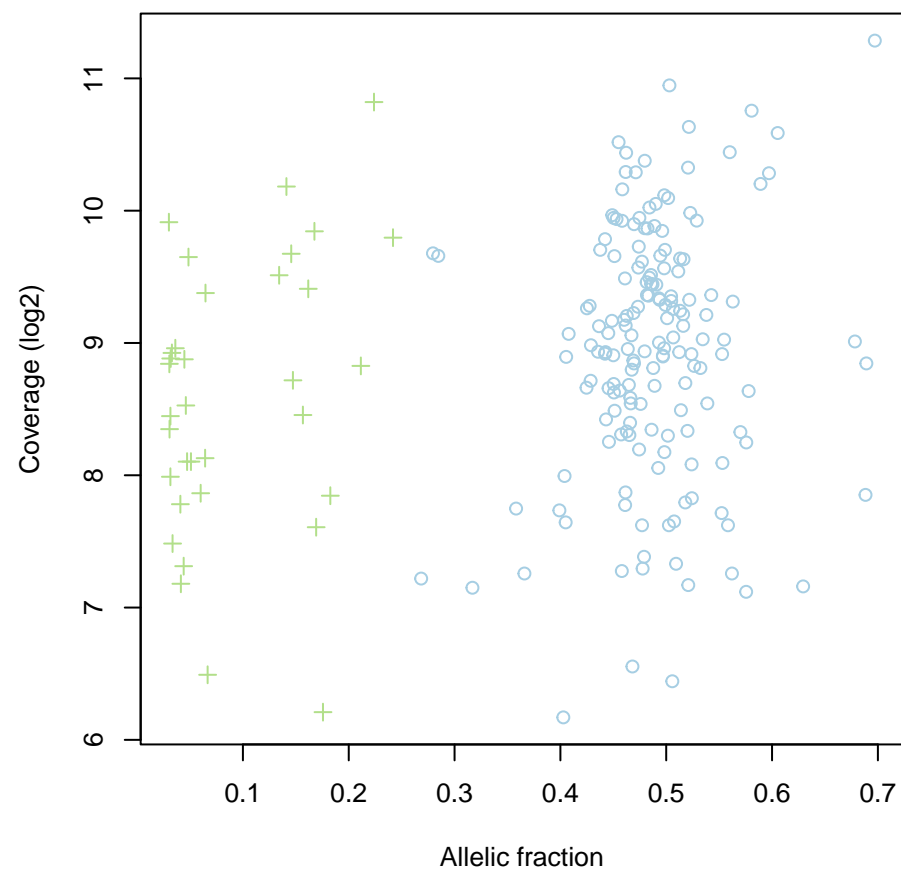
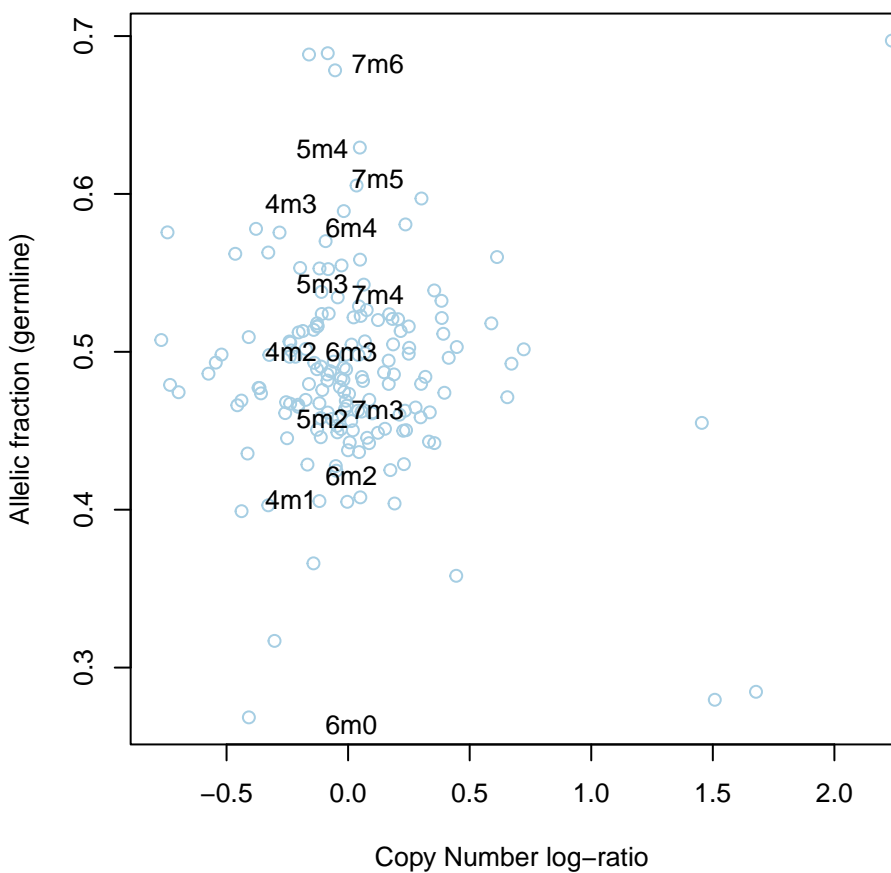


Purity: 0.23 Tumor ploidy: 5.885 SNV log-likelihood: -184.93 GoF: 93.3% Mean coverage: 584,392

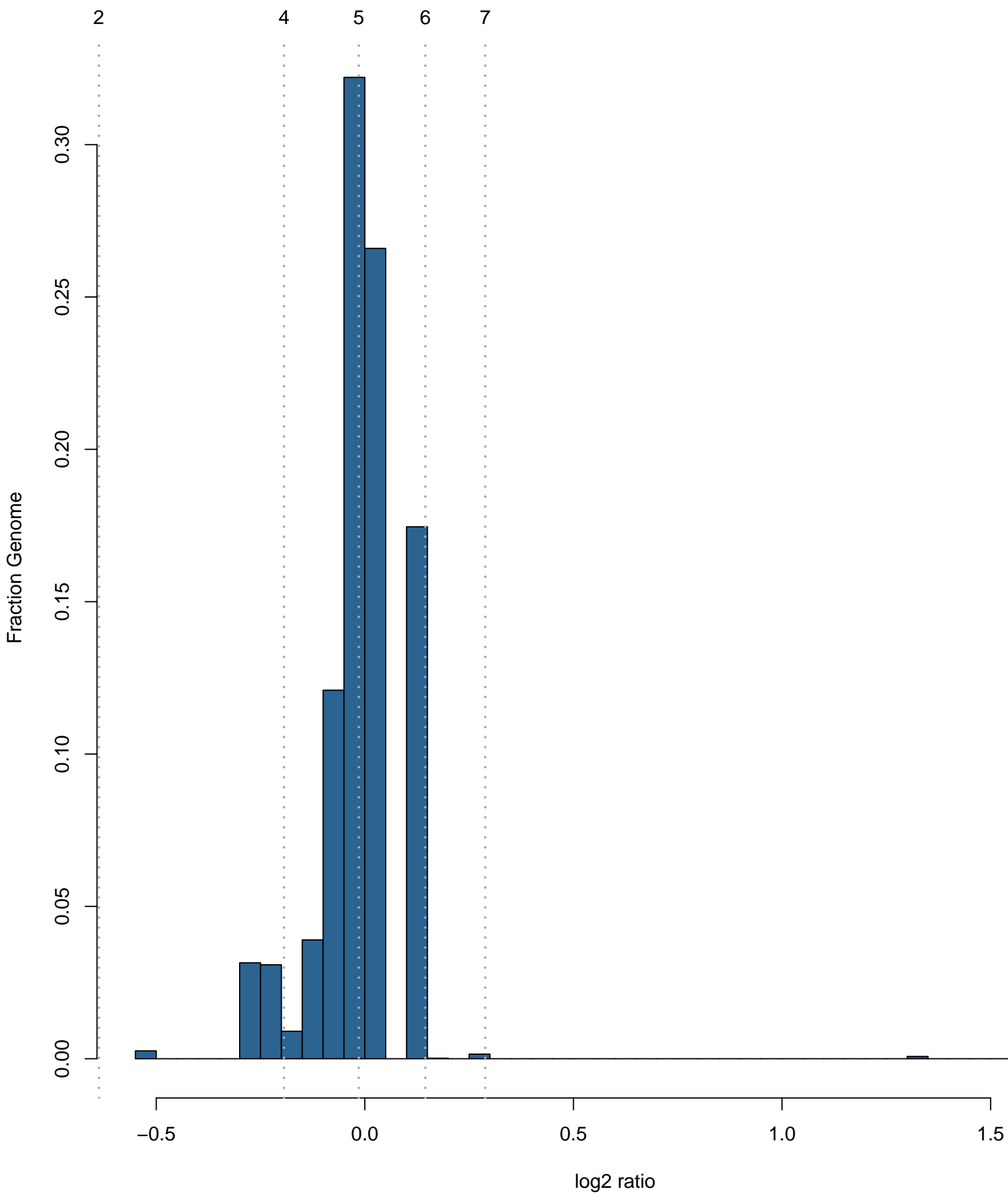


SCNA-fit log-likelihood: -8703.12

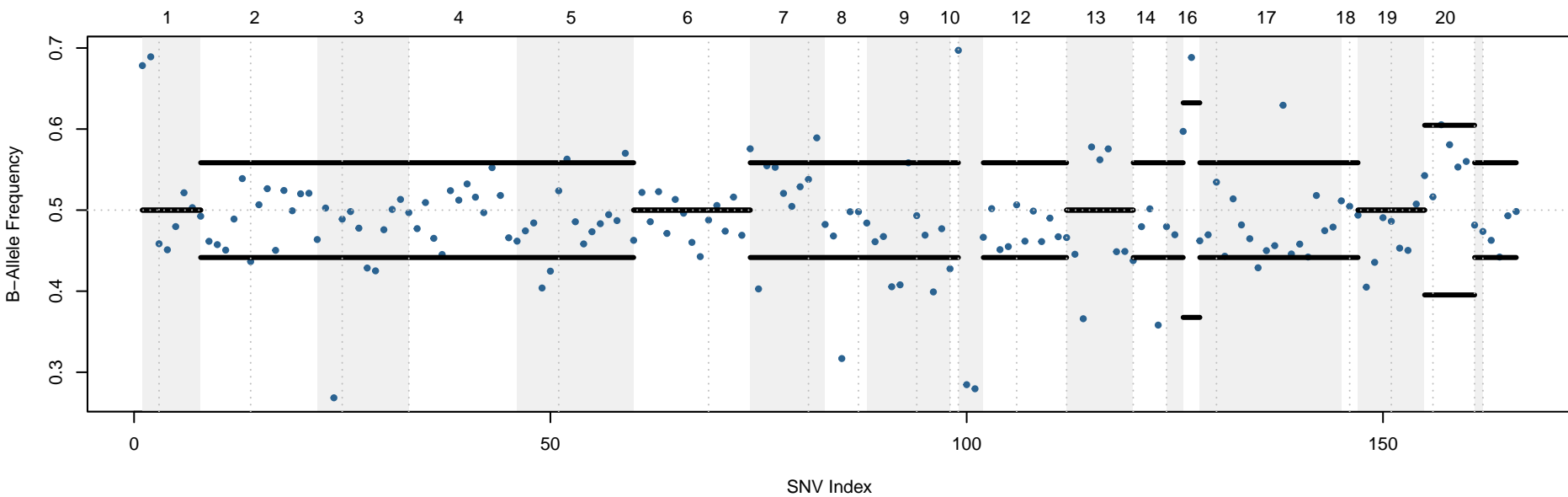




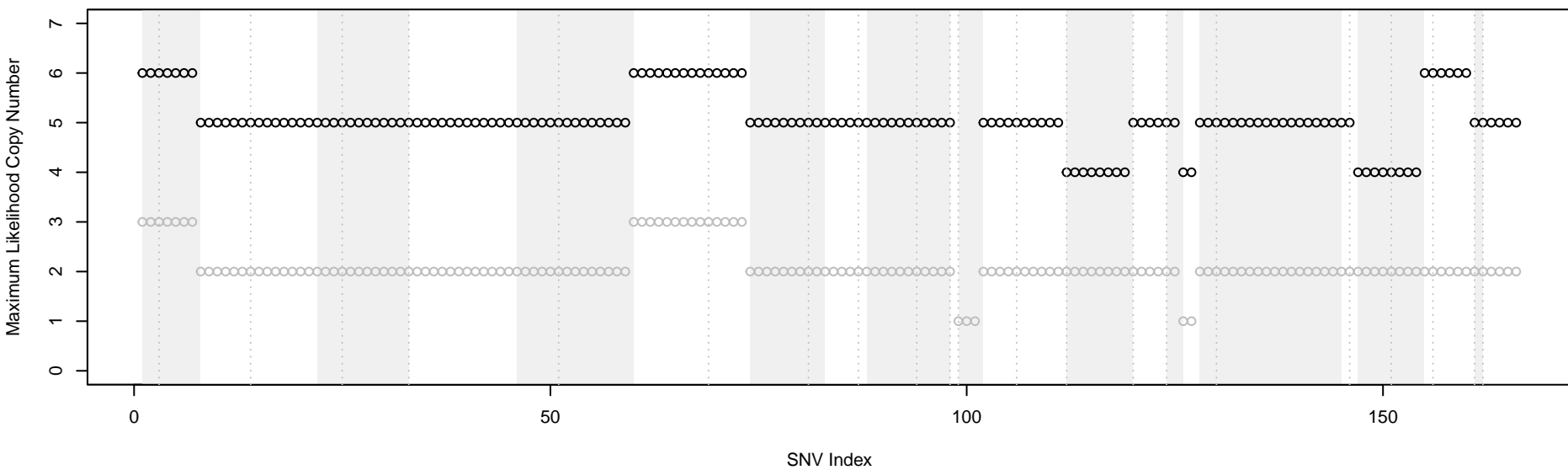
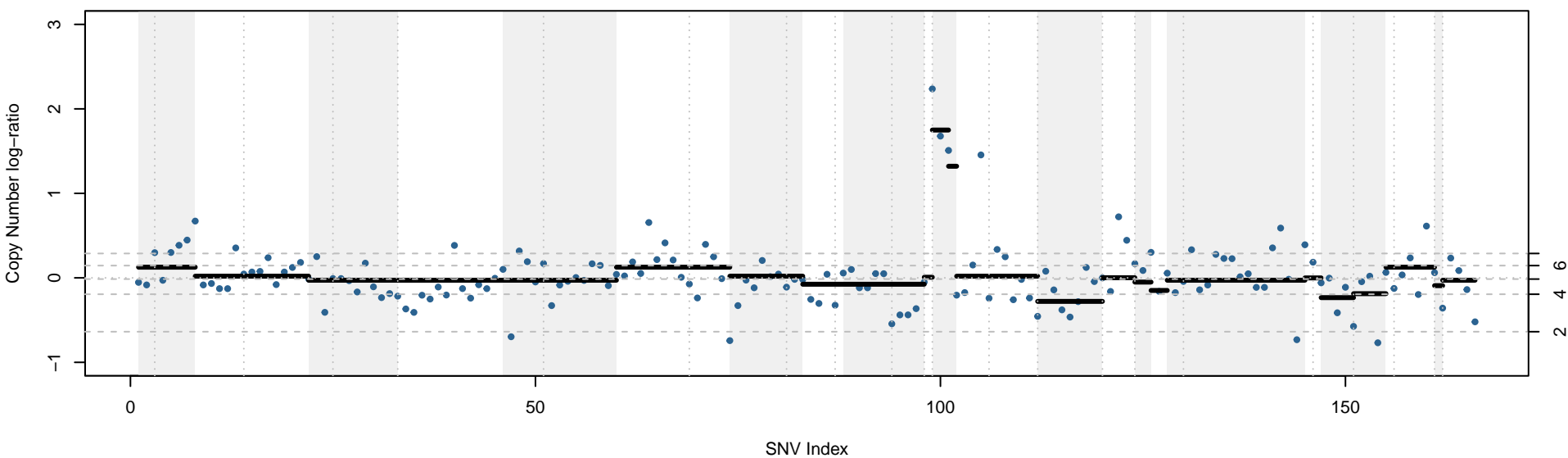
Purity: 0.36 Tumor ploidy: 5.086

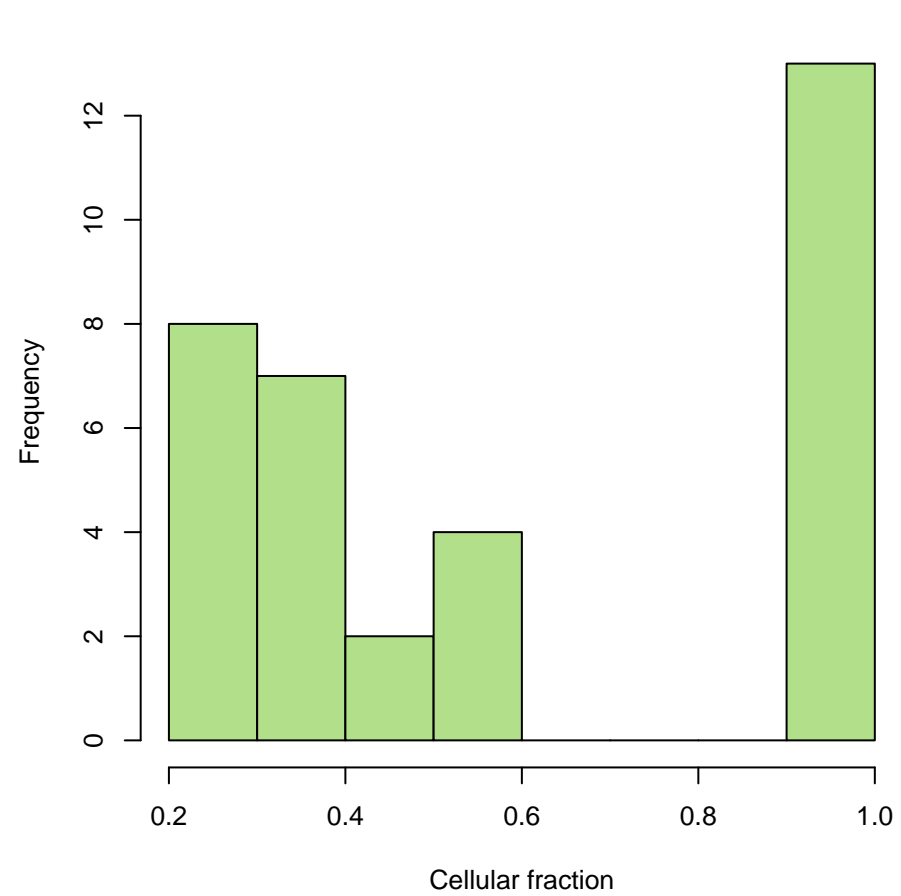
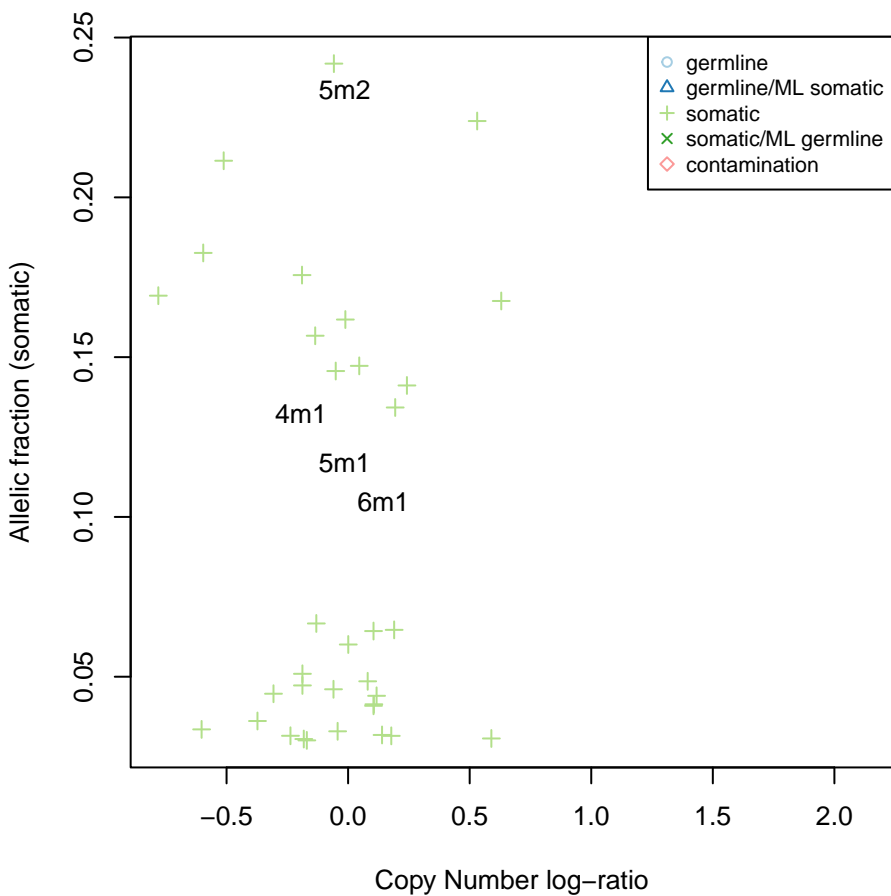
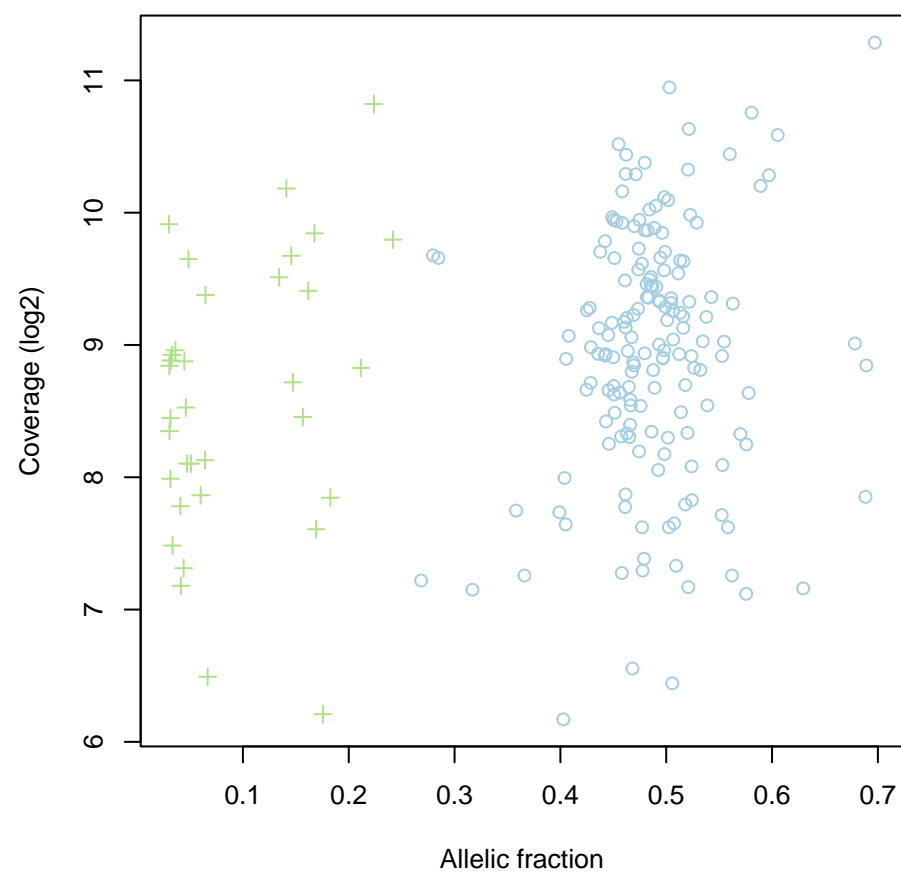
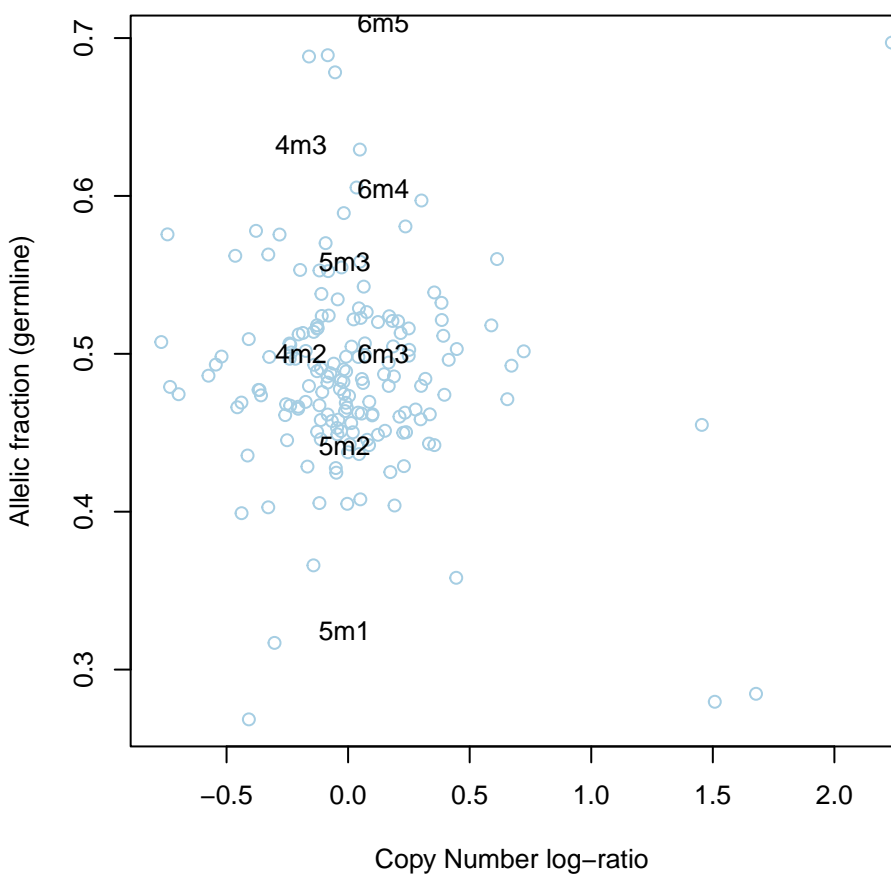


Purity: 0.36 Tumor ploidy: 5.086 SNV log-likelihood: -198.38 GoF: 92.4% Mean coverage: 584,392

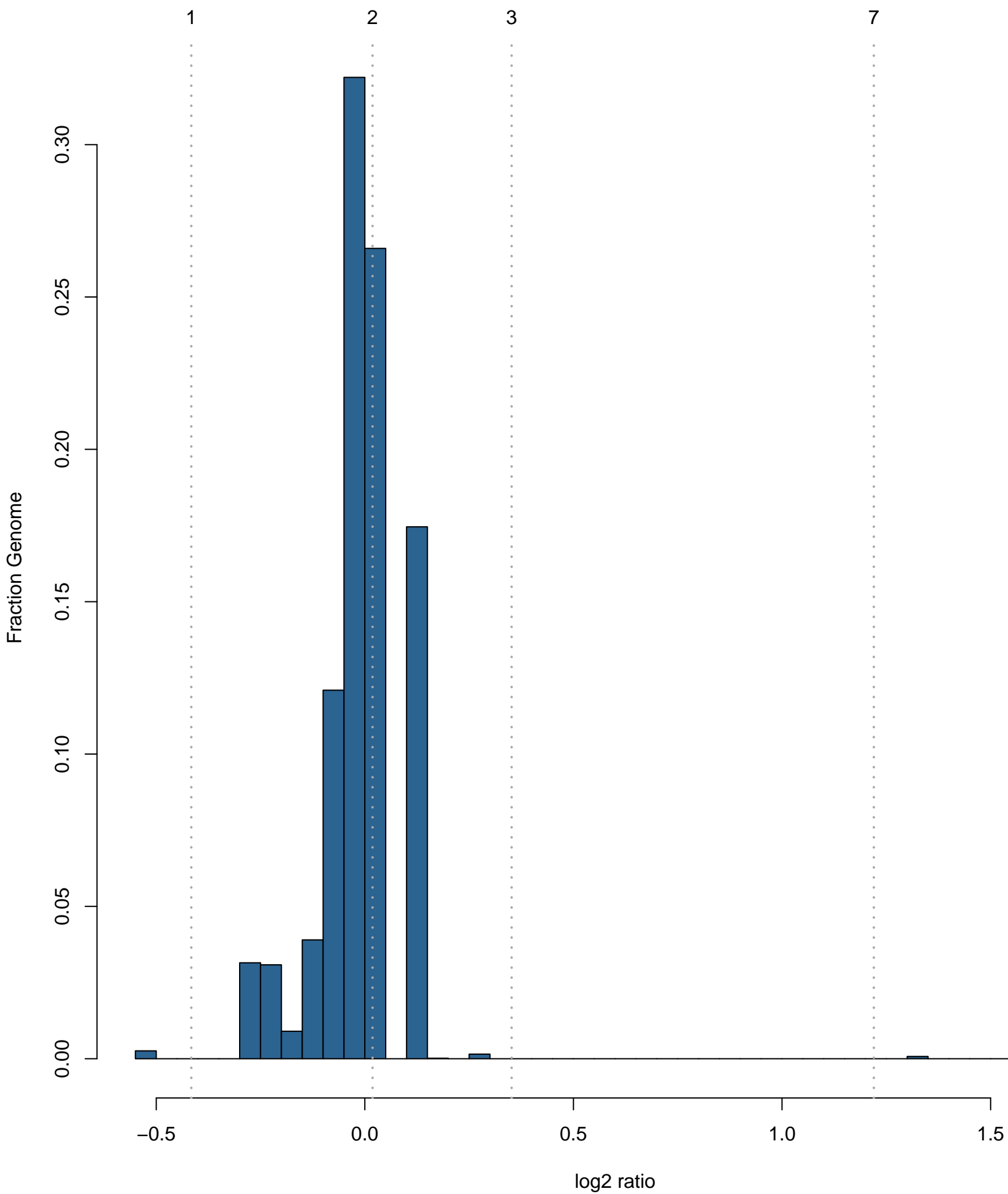


SCNA-fit log-likelihood: -8709.88

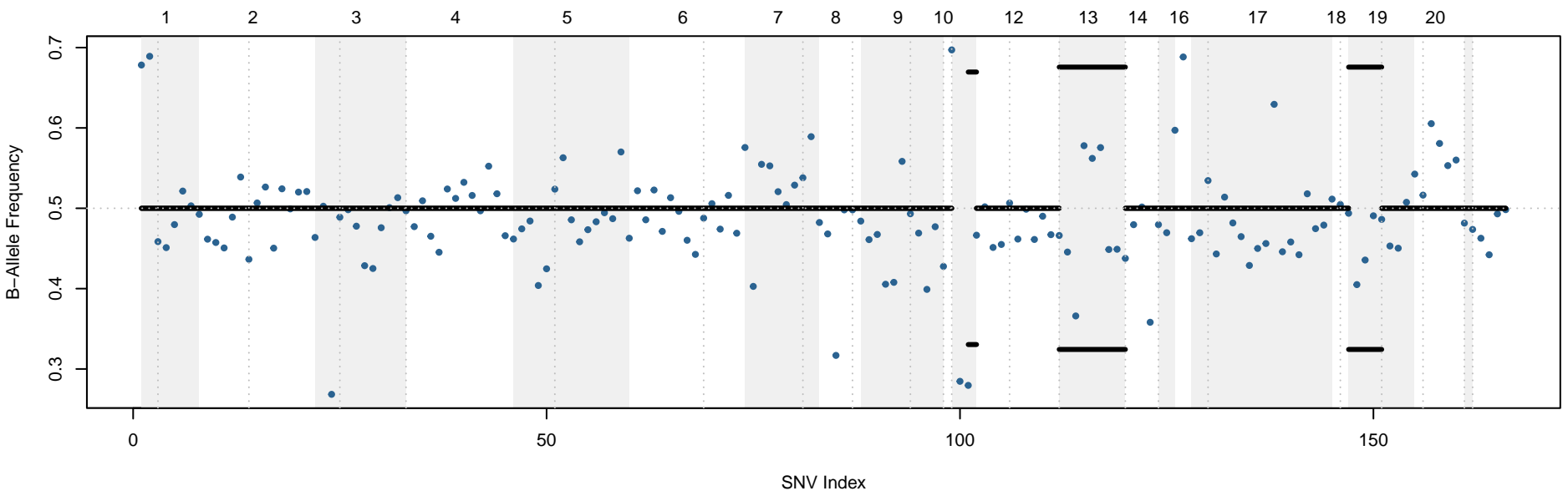




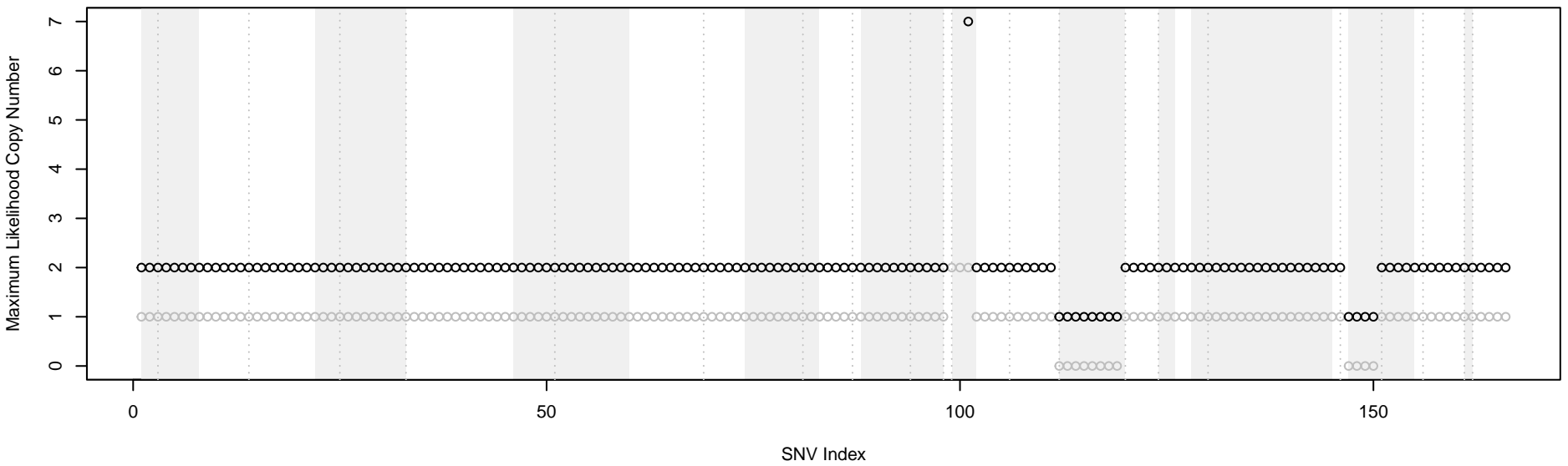
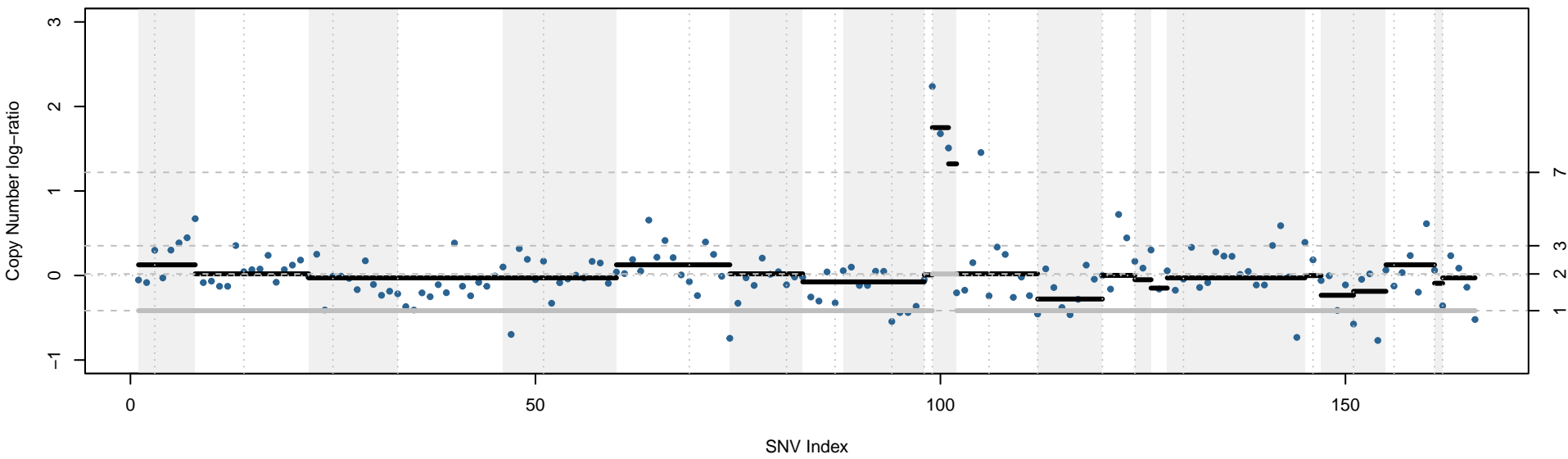
Purity: 0.52 Tumor ploidy: 1.951

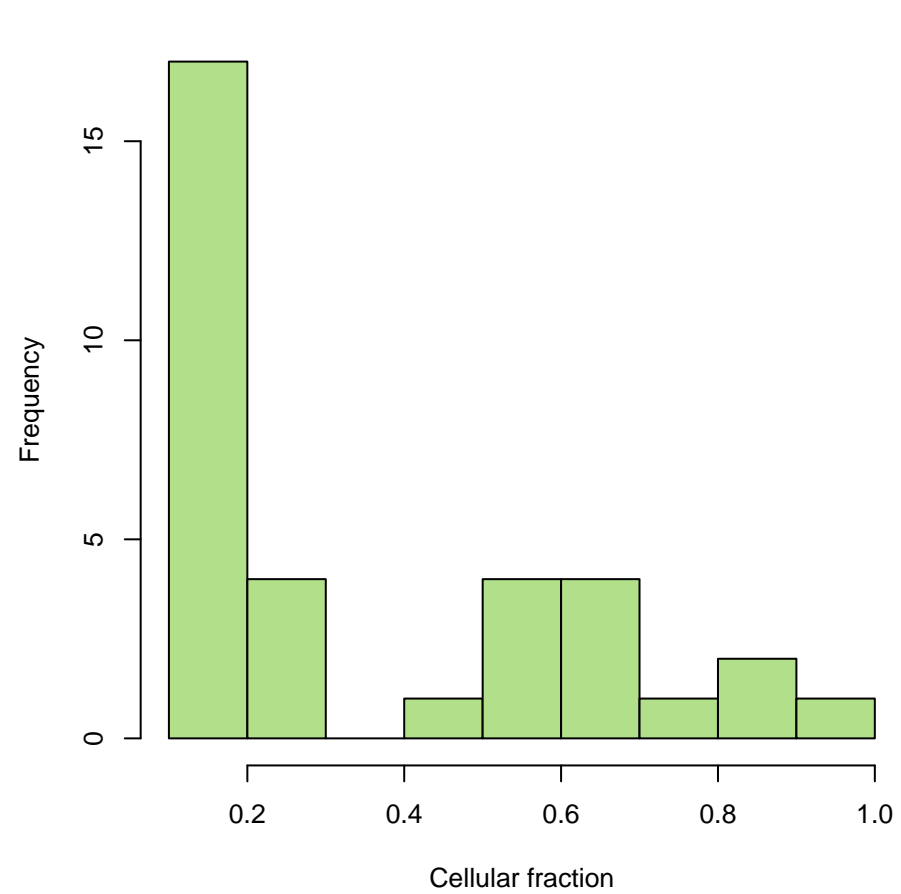
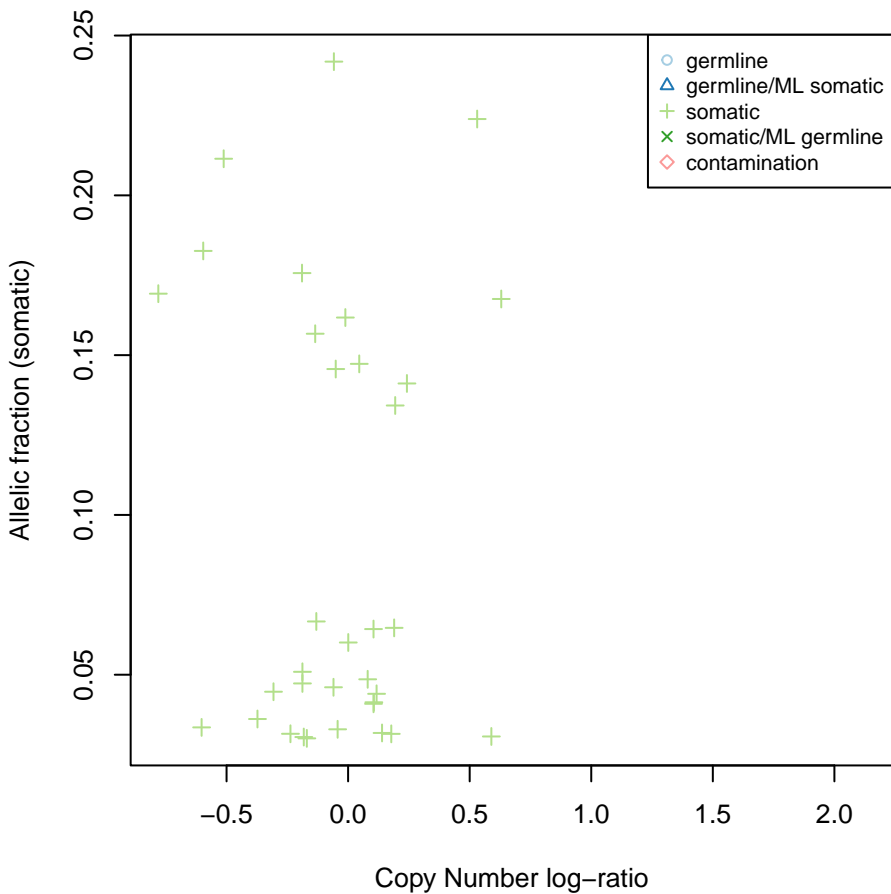
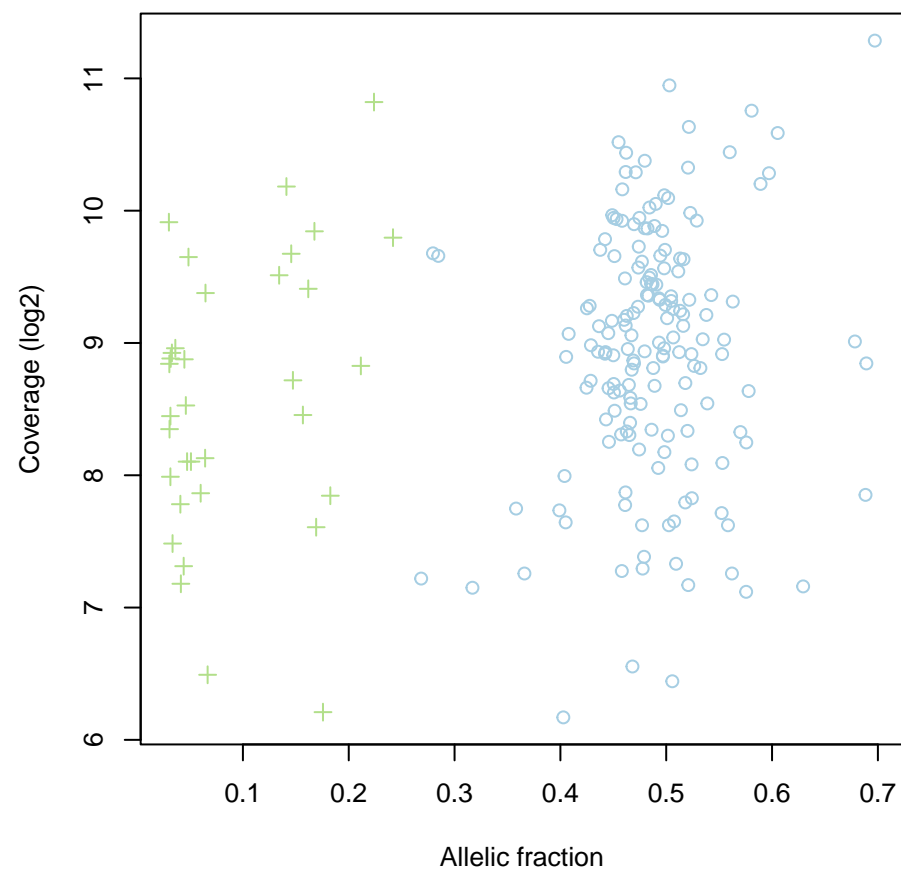
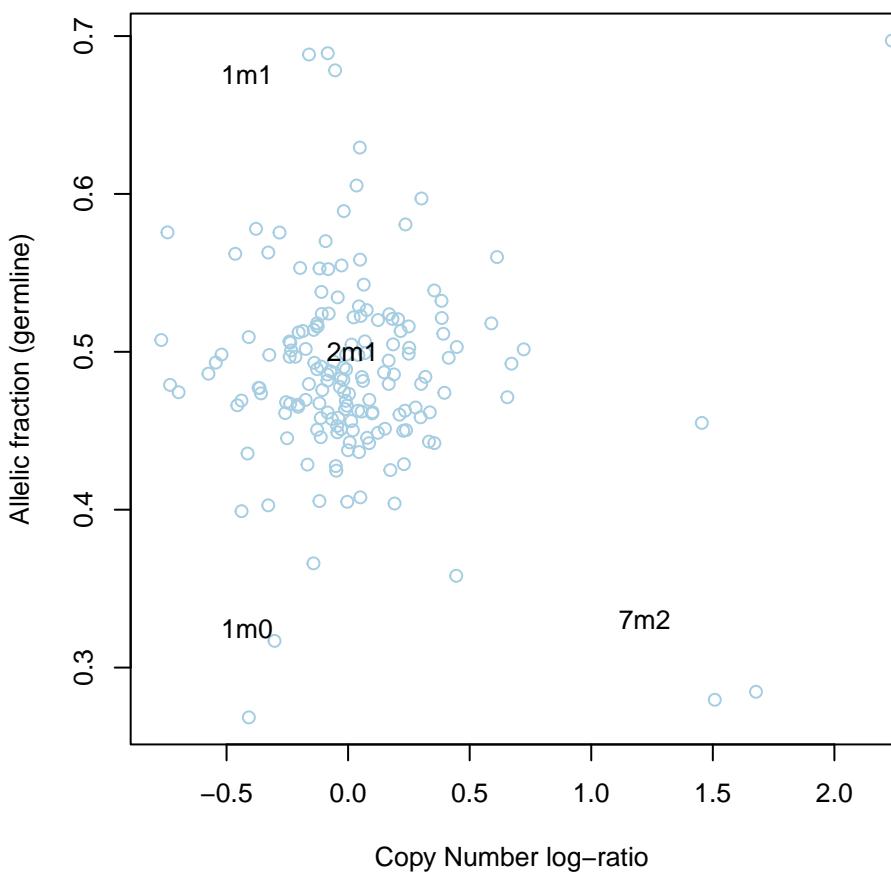


Purity: 0.52 Tumor ploidy: 1.951 SNV log-likelihood: -170.82 GoF: 78.3% Mean coverage: 584,392



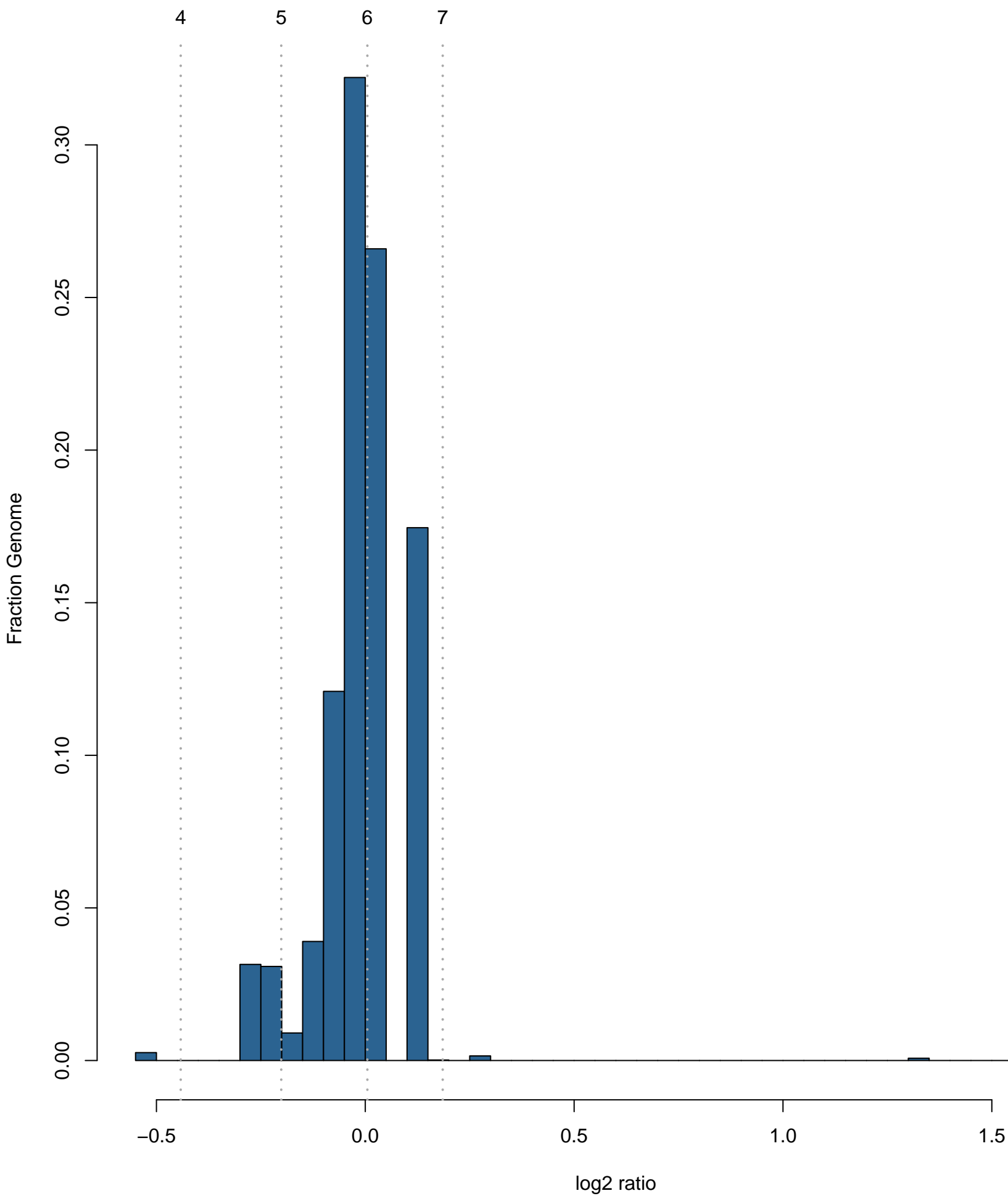
SCNA-fit log-likelihood: -8842.12



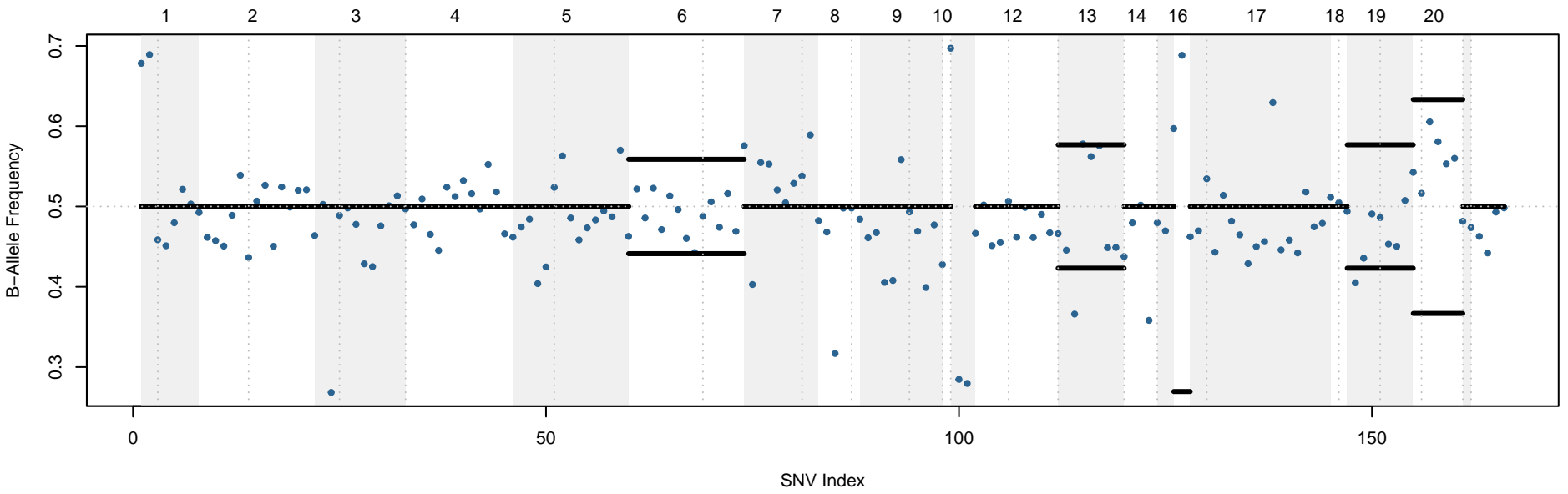




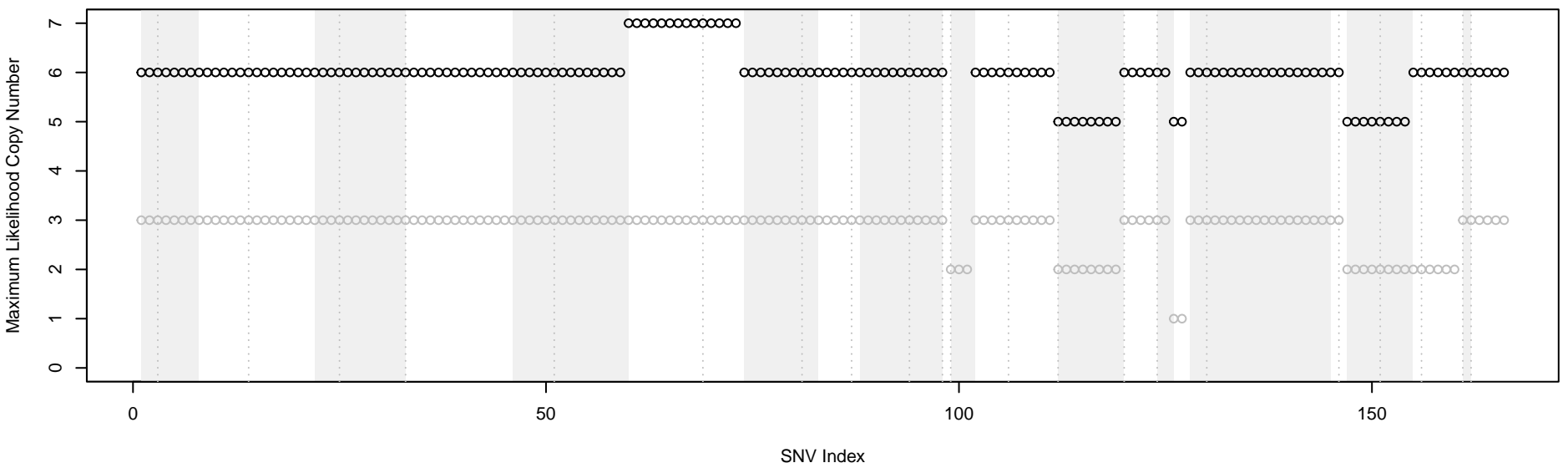
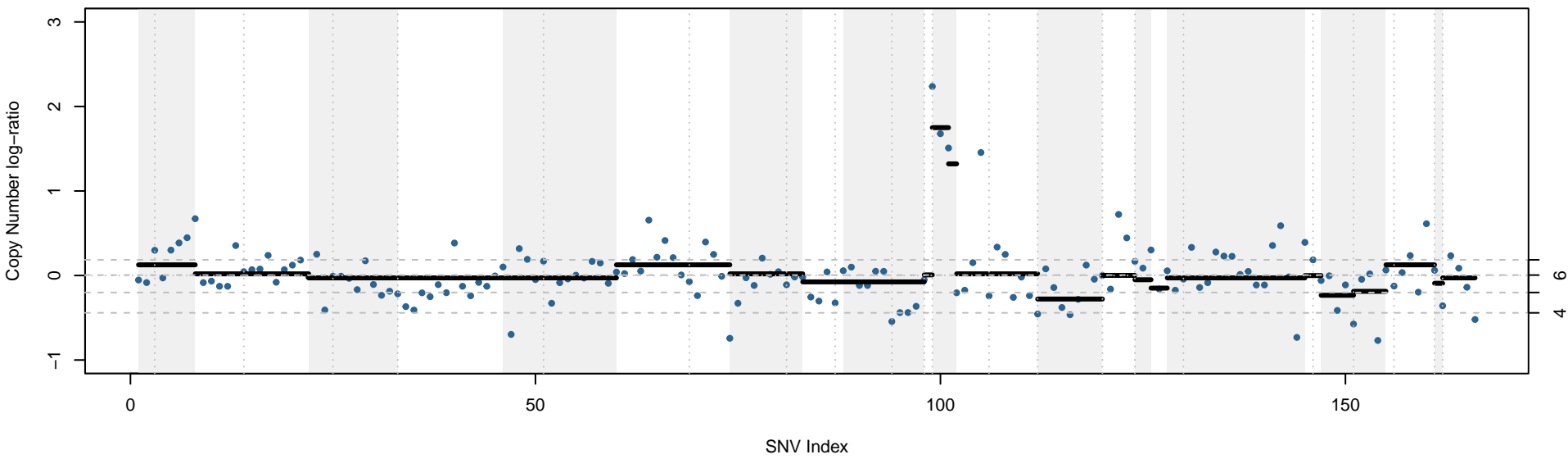
Purity: 0.57 Tumor ploidy: 5.974

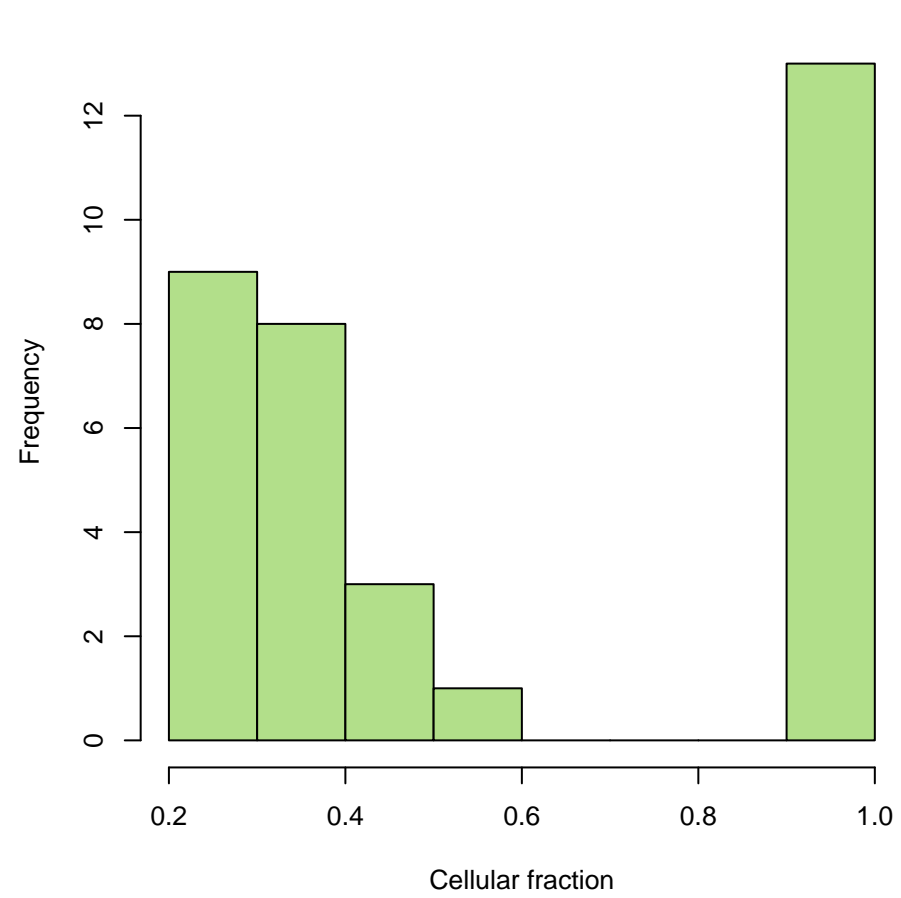
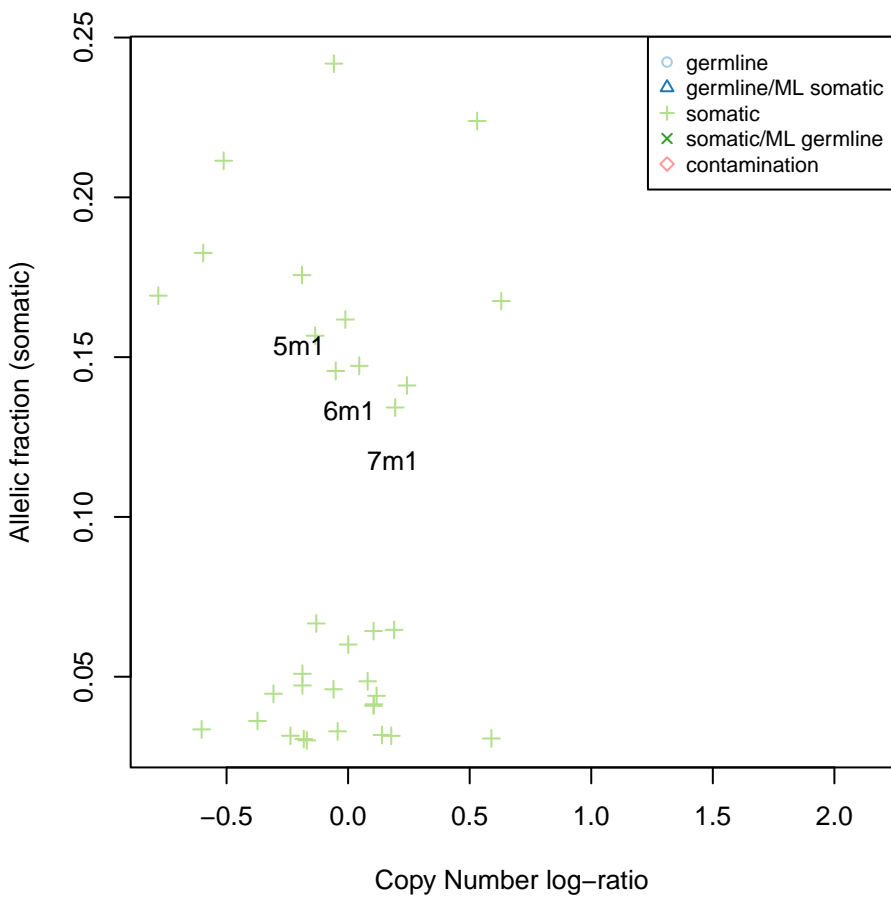
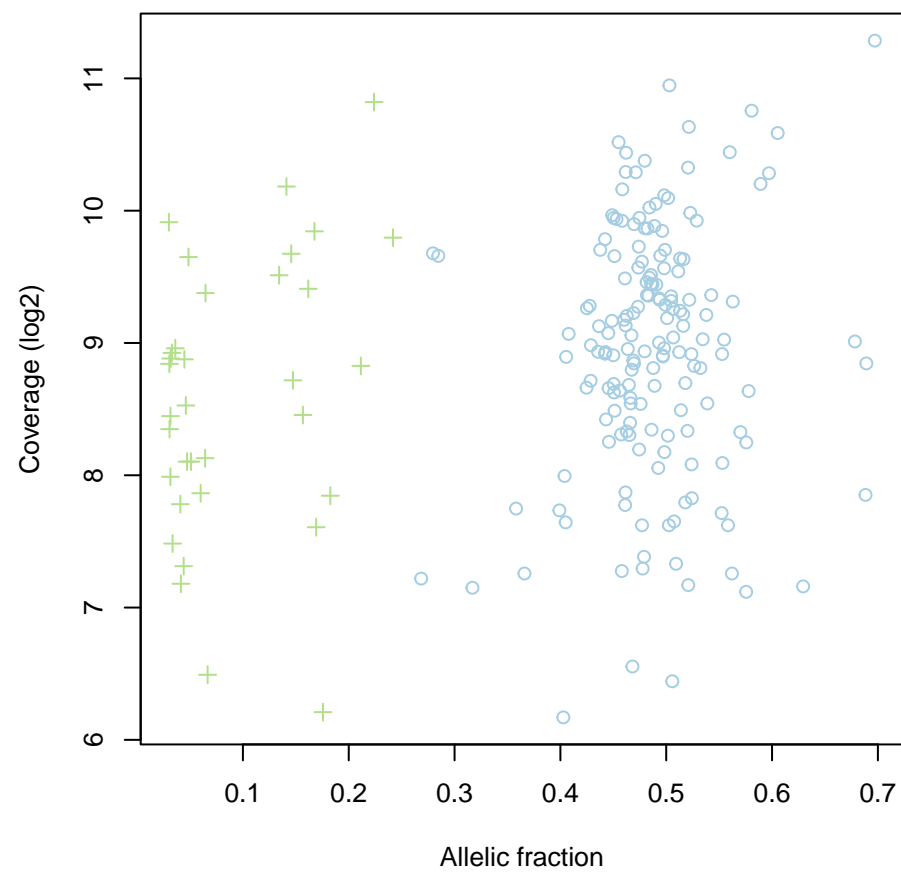
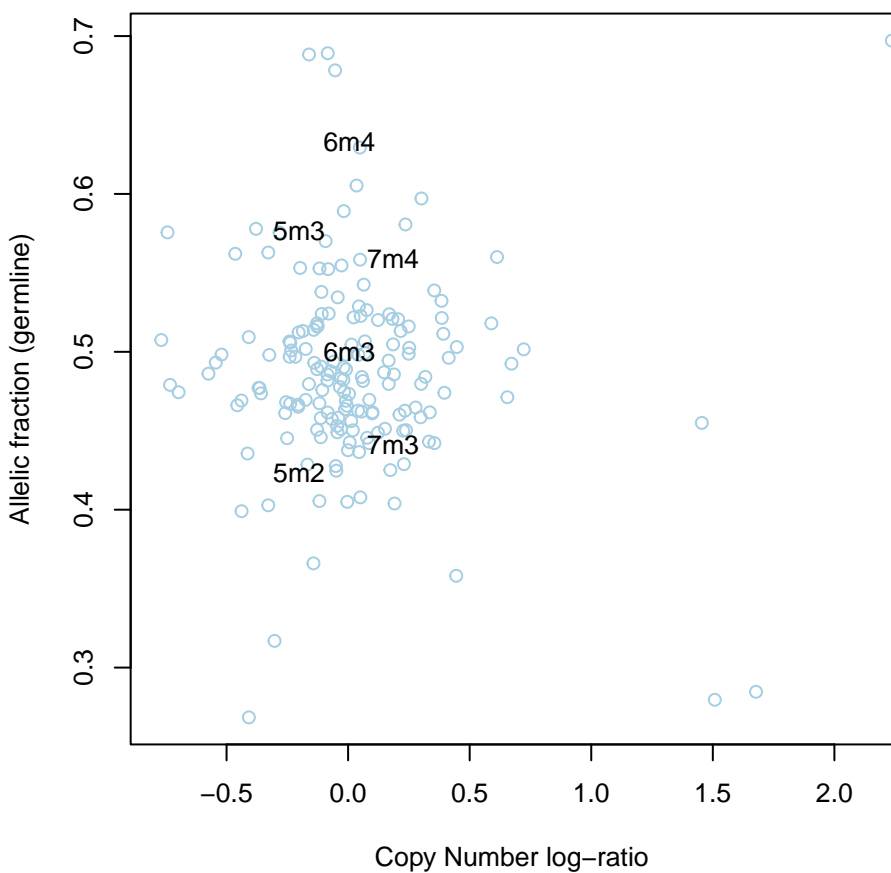


Purity: 0.57 Tumor ploidy: 5.974 SNV log-likelihood: -220.08 GoF: 90.4% Mean coverage: 584,392

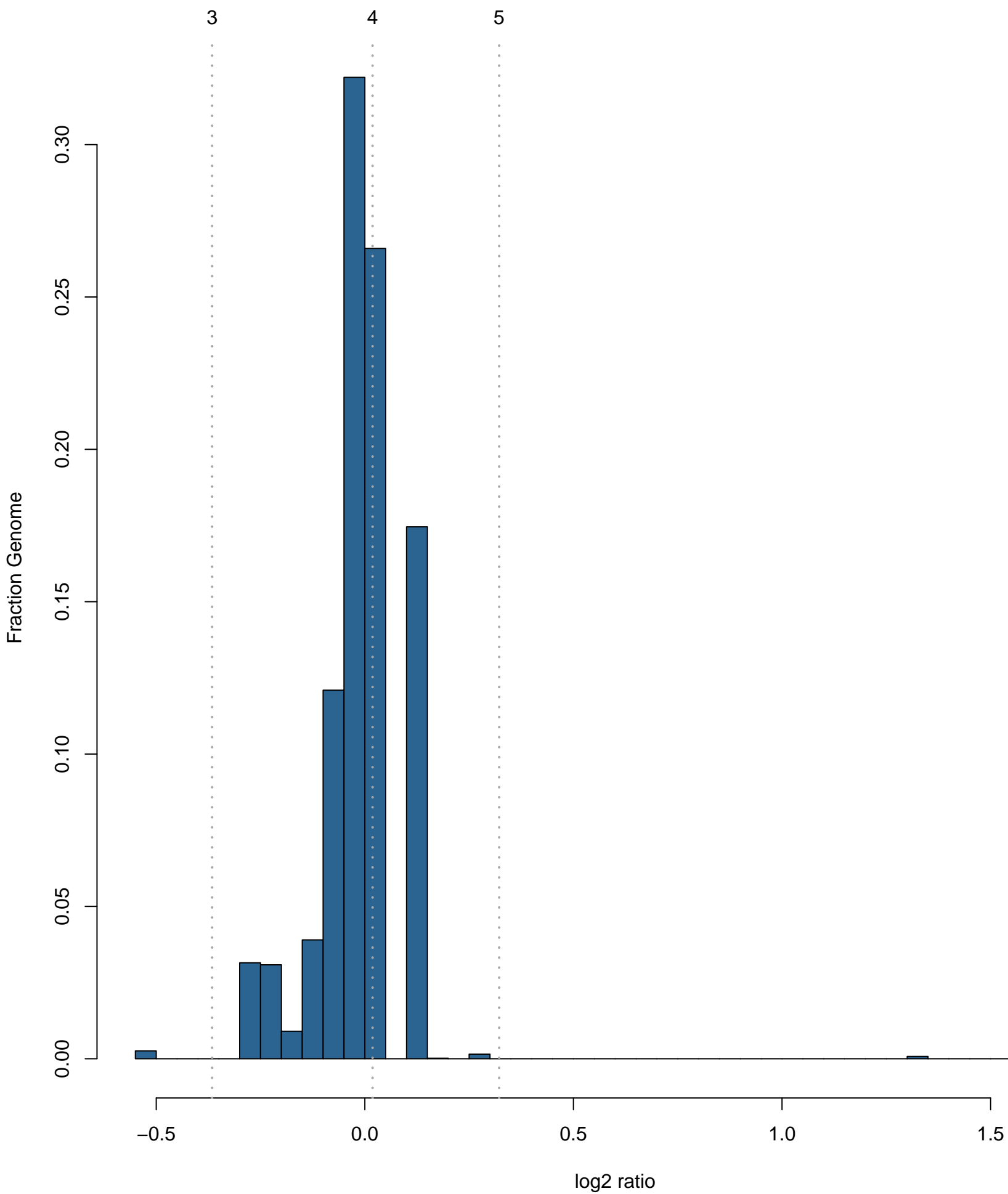


SCNA-fit log-likelihood: -8755.05

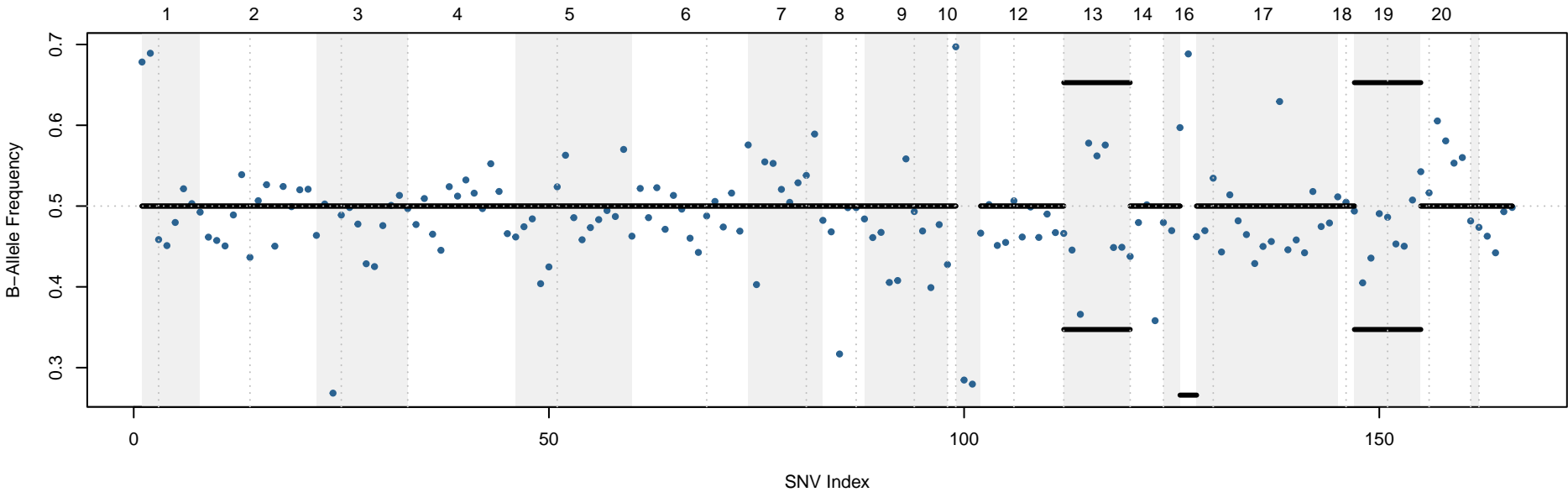




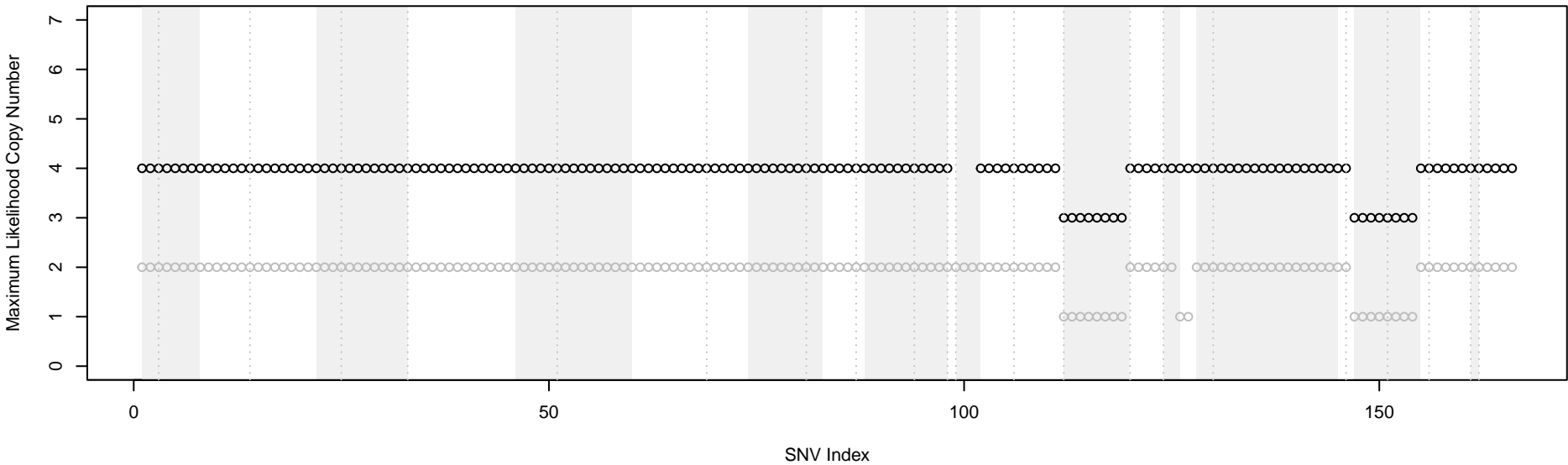
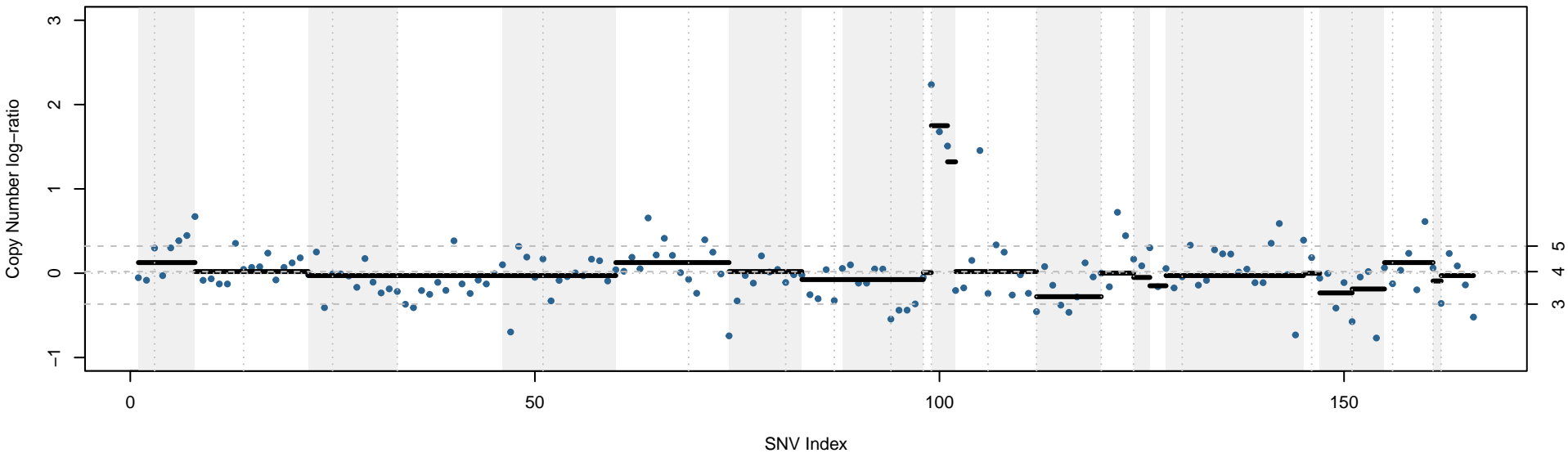
Purity: 0.88 Tumor ploidy: 3.945

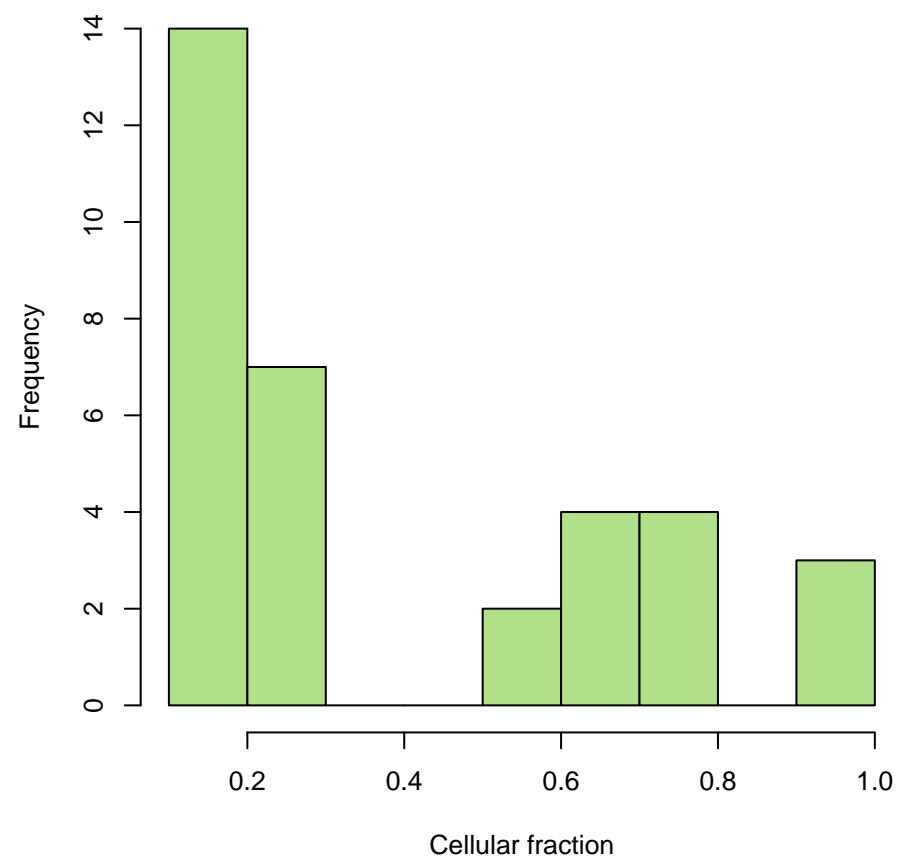
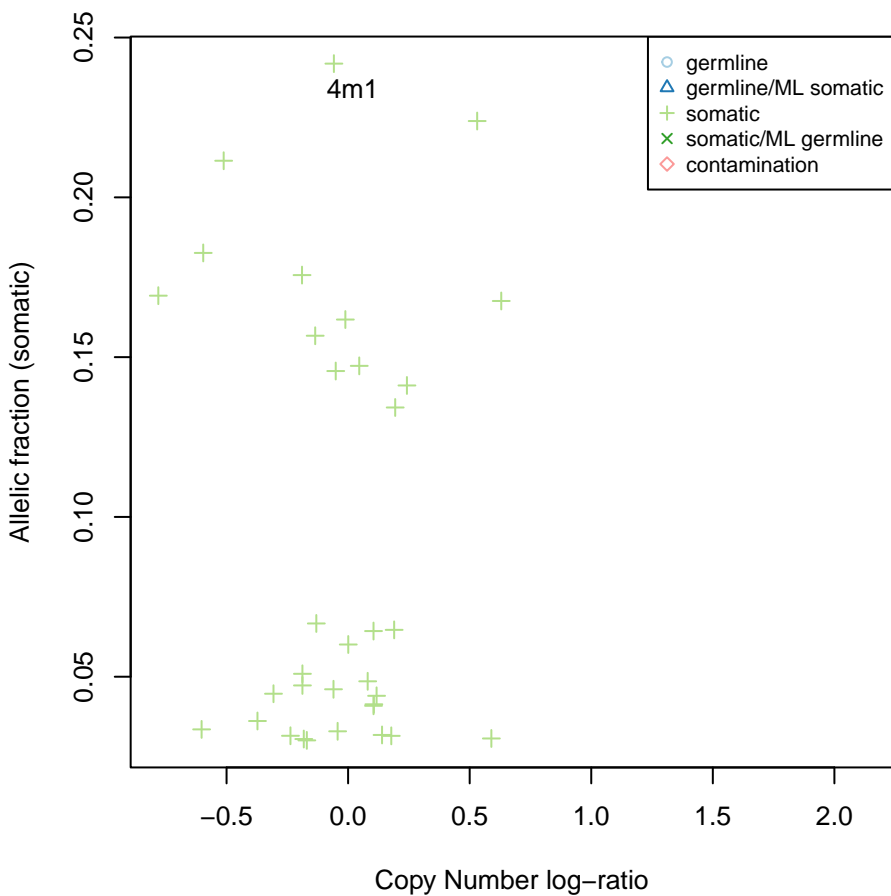
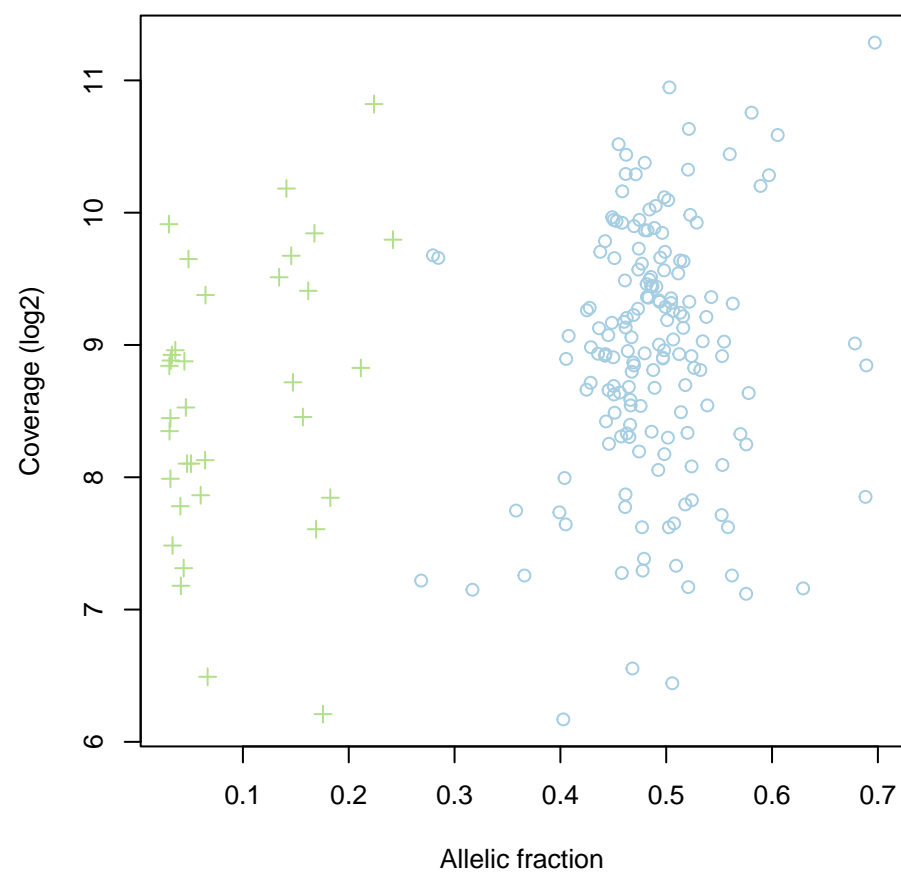
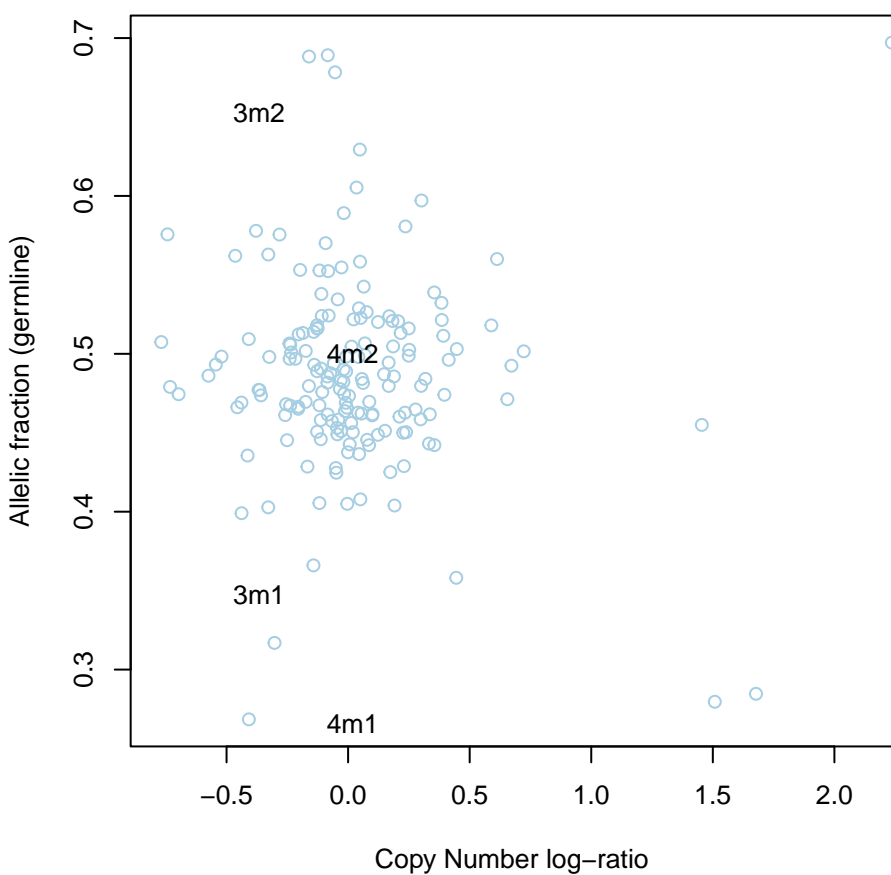


Purity: 0.88 Tumor ploidy: 3.945 SNV log-likelihood: -230 GoF: 80.7% Mean coverage: 584;392

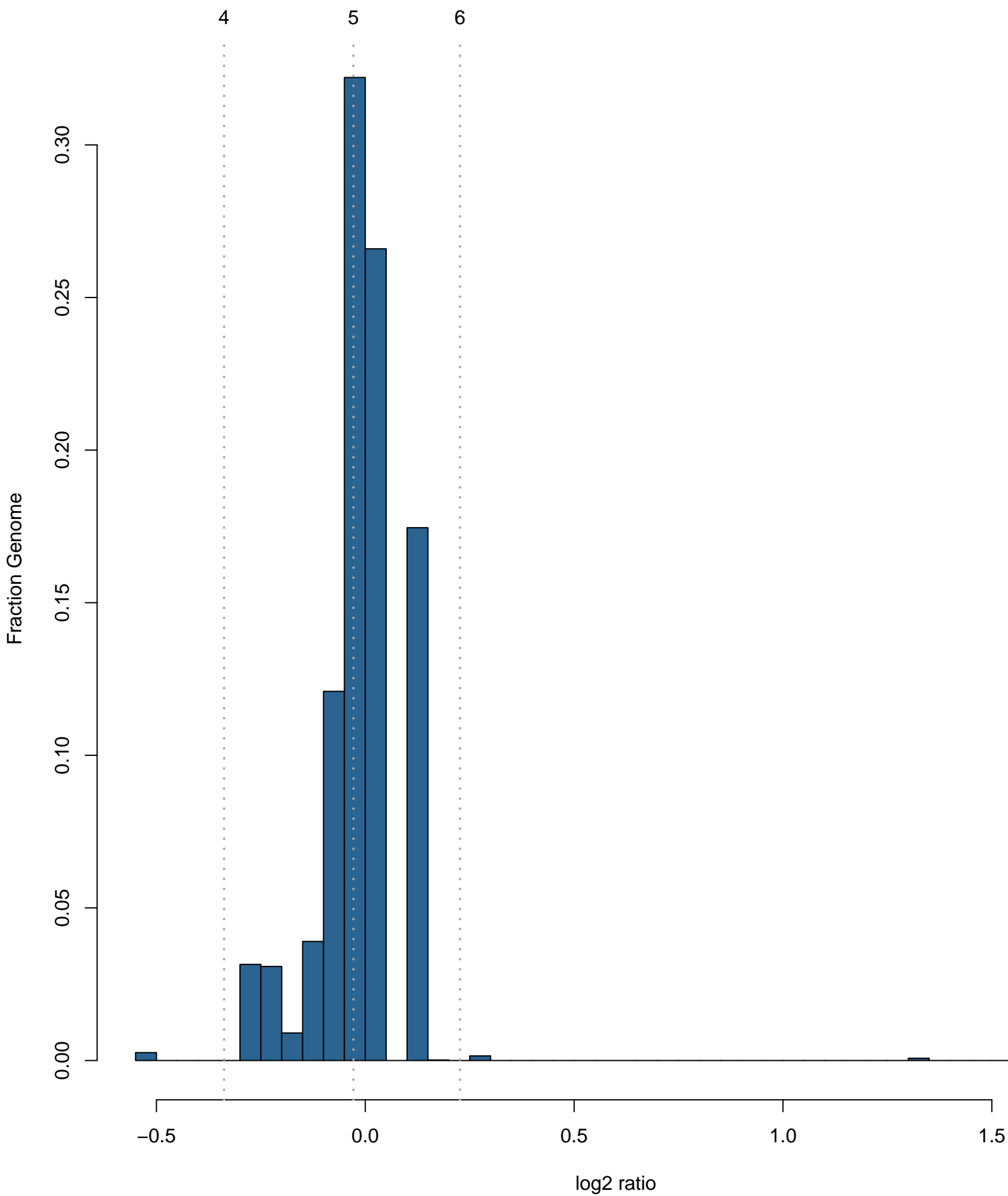


SCNA-fit log-likelihood: -8830.36

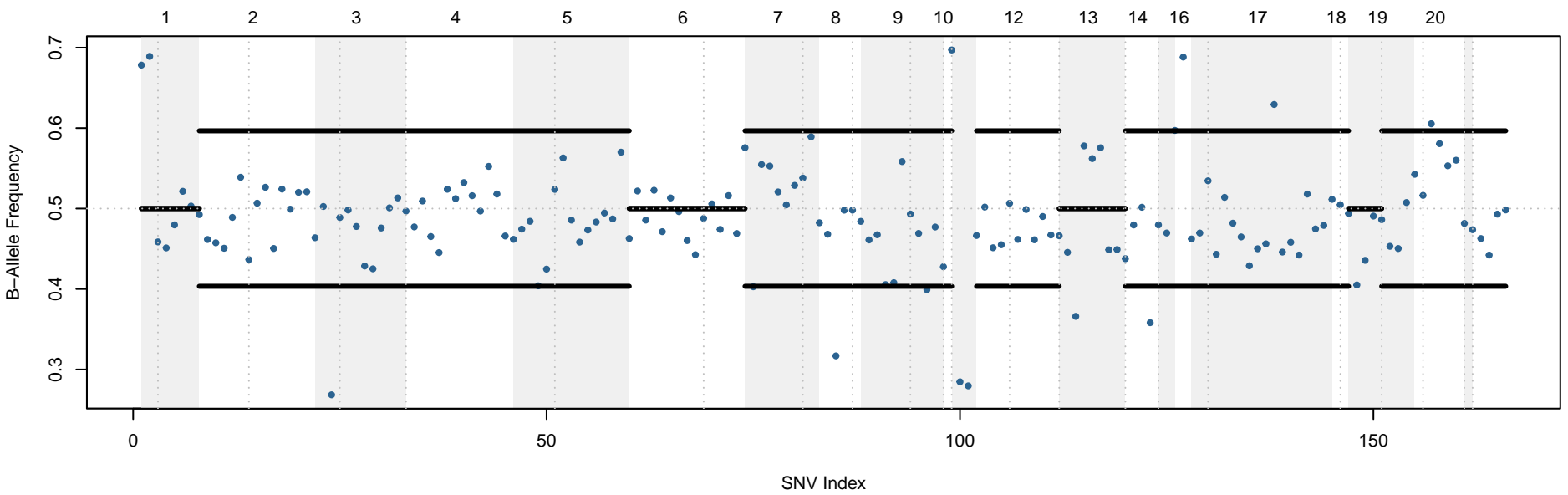




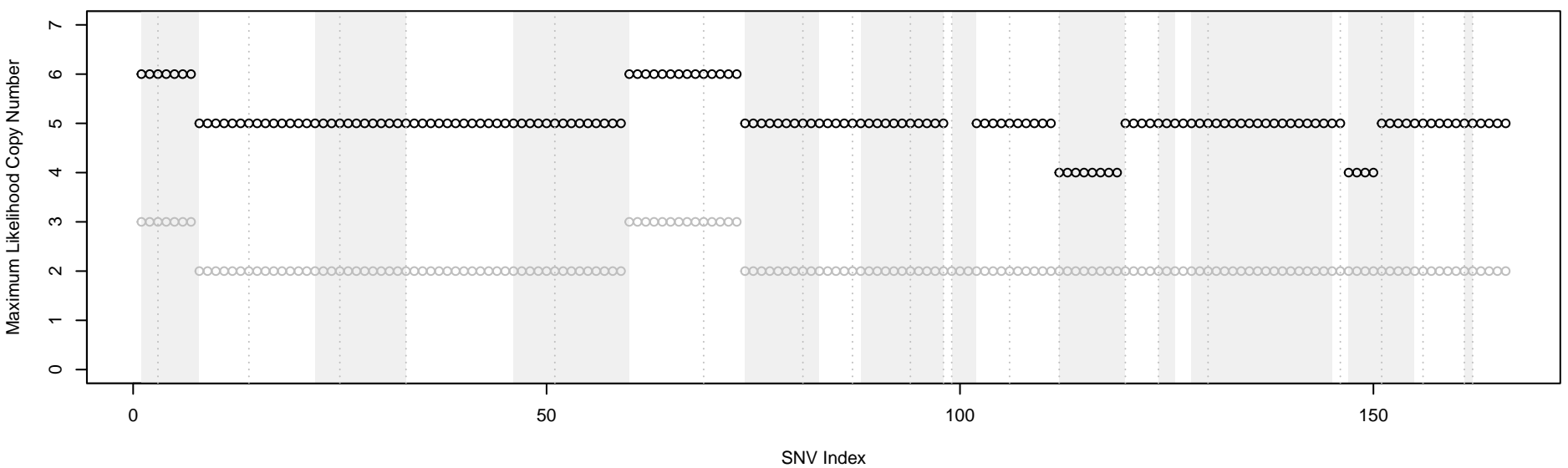
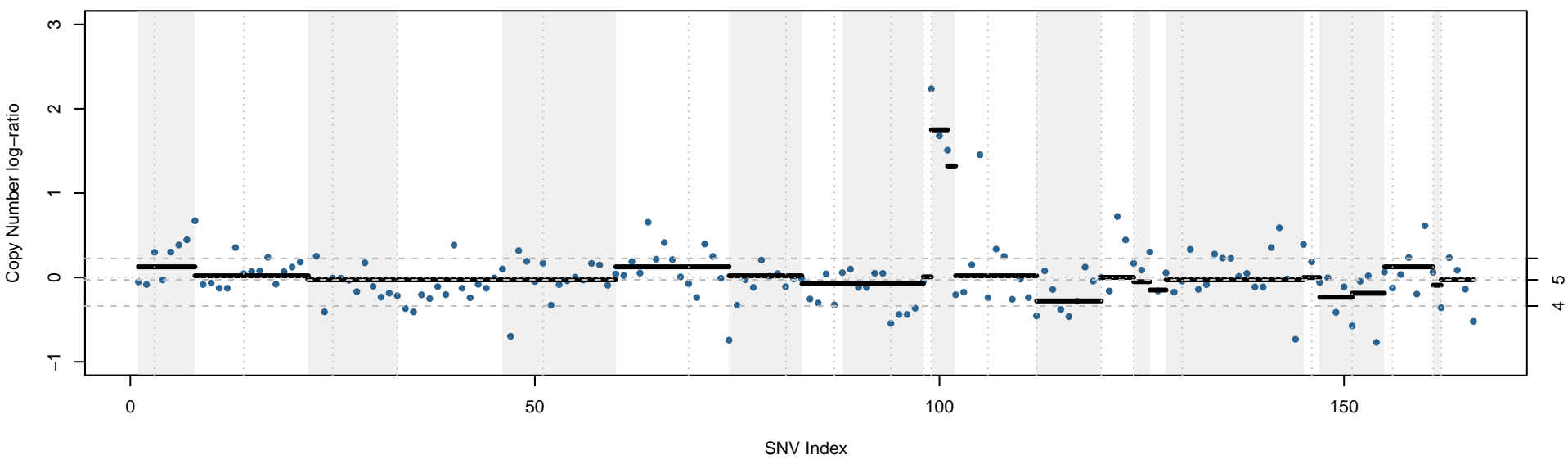
Purity: 0.92 Tumor ploidy: 5.102



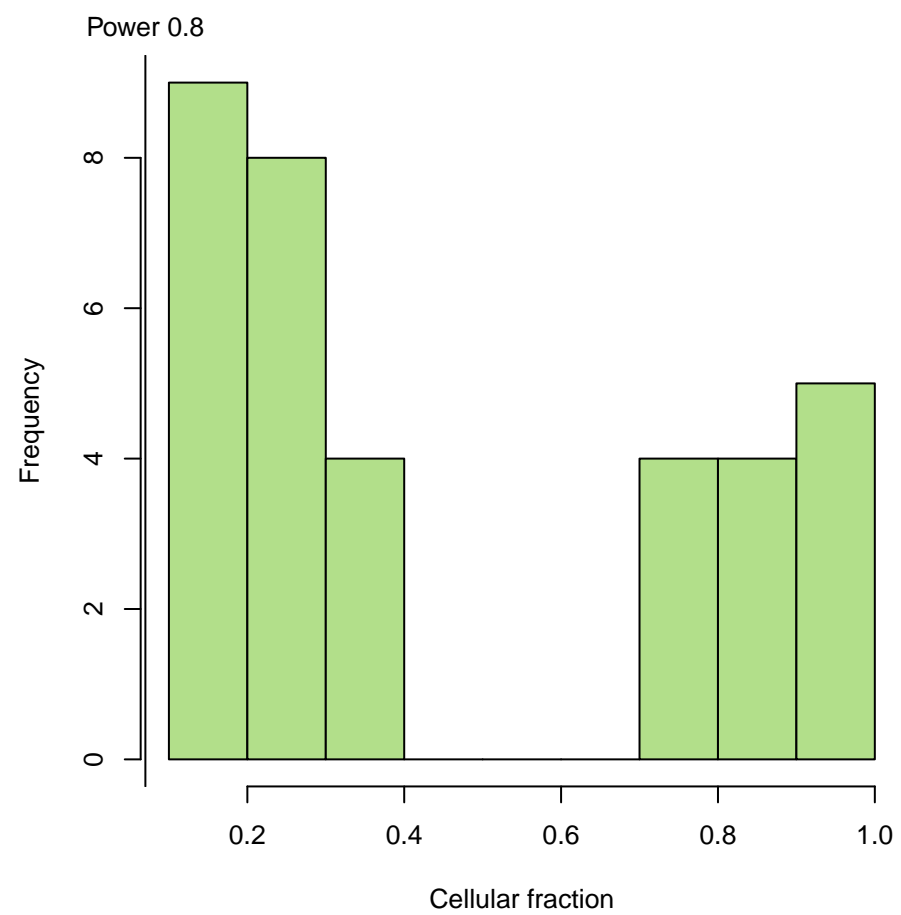
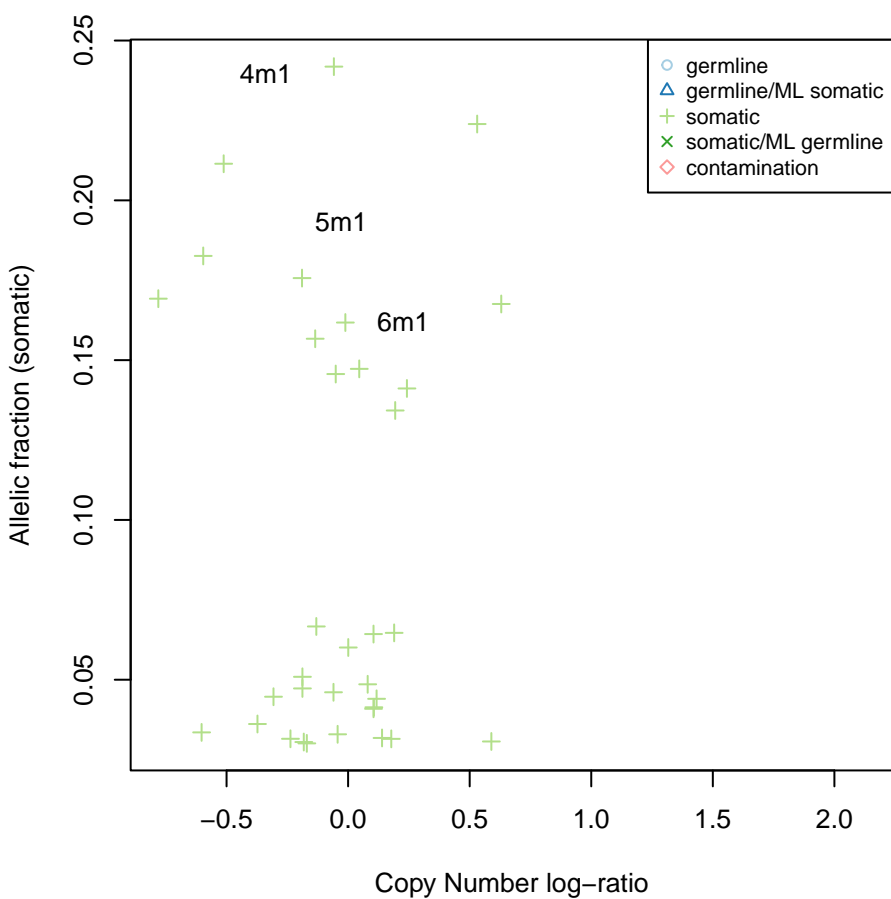
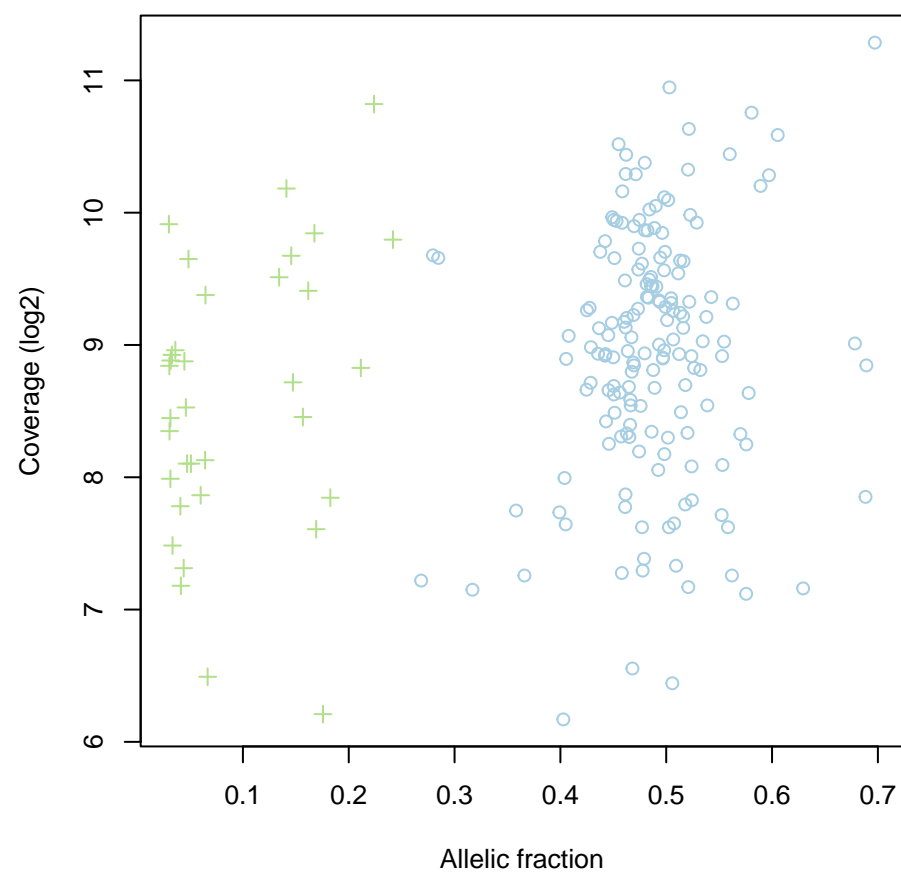
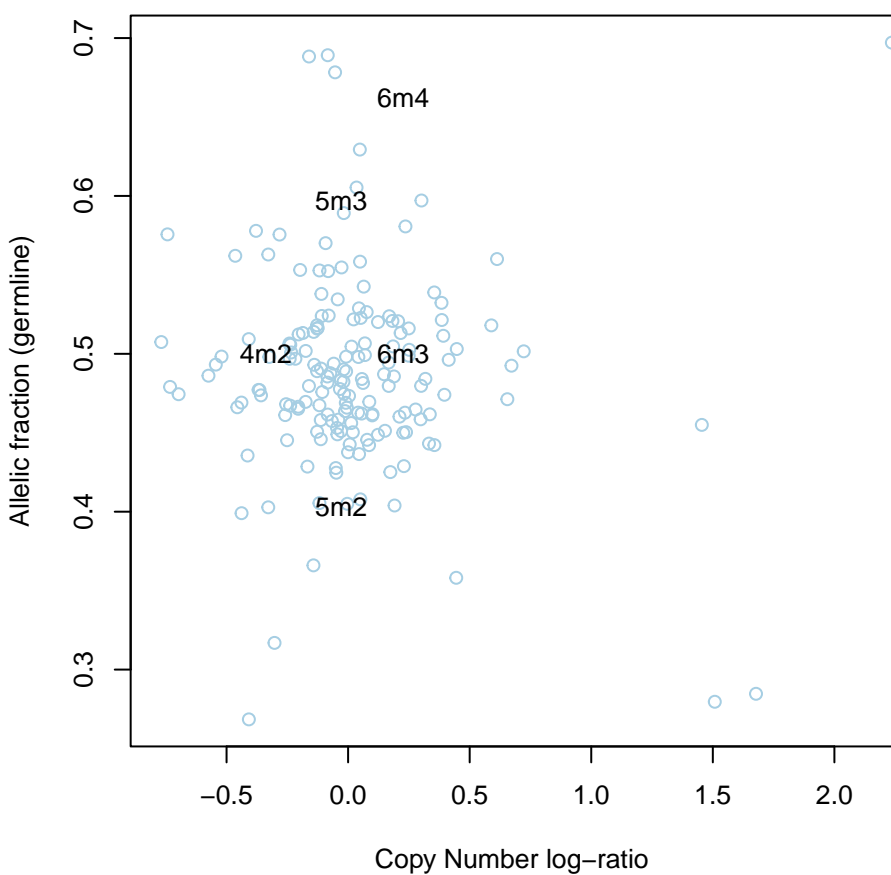
Purity: 0.92 Tumor ploidy: 5.102 SNV log-likelihood: -434.05 GoF: 82.6% Mean coverage: 584,392



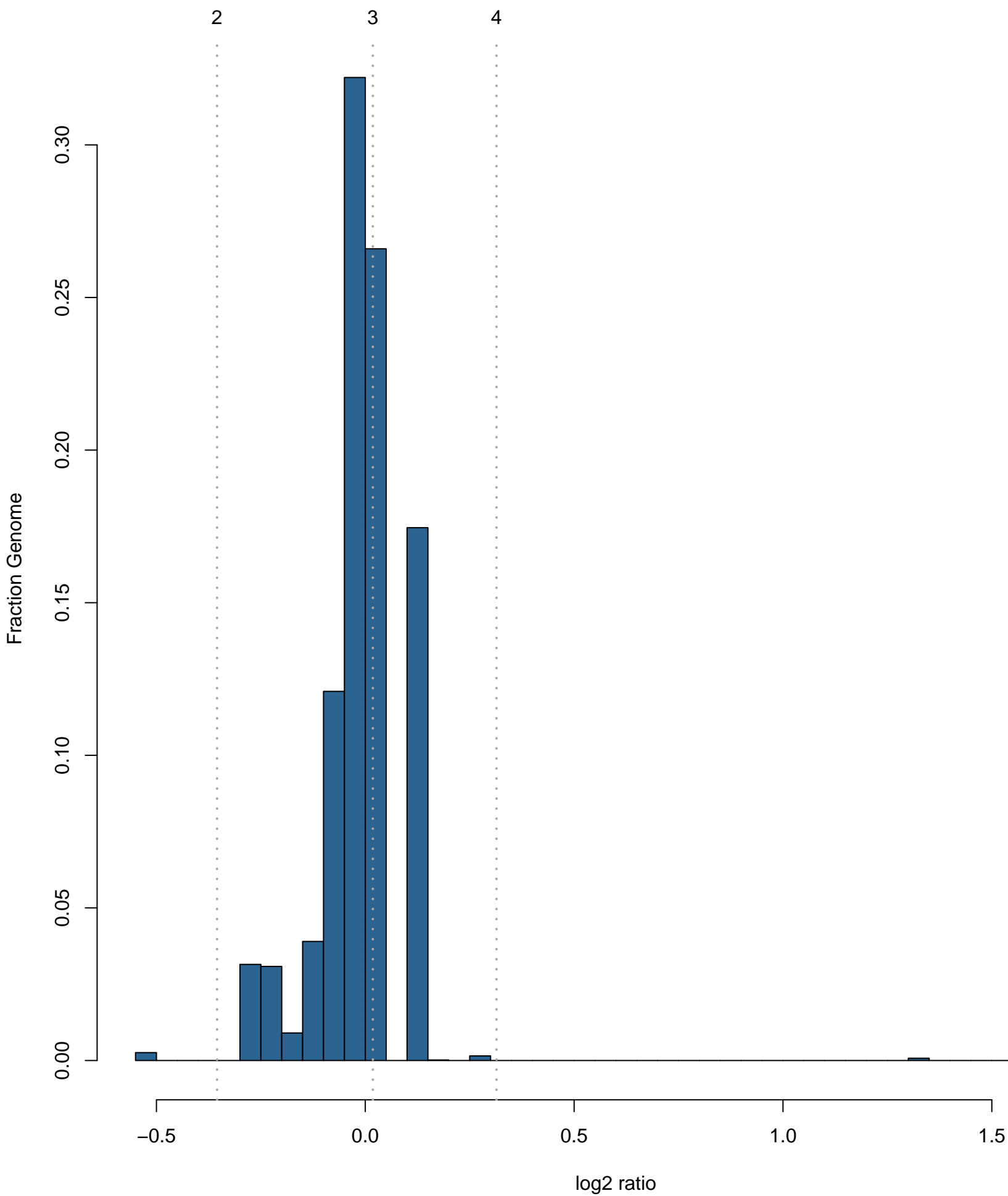
SCNA-fit log-likelihood: -8773.02



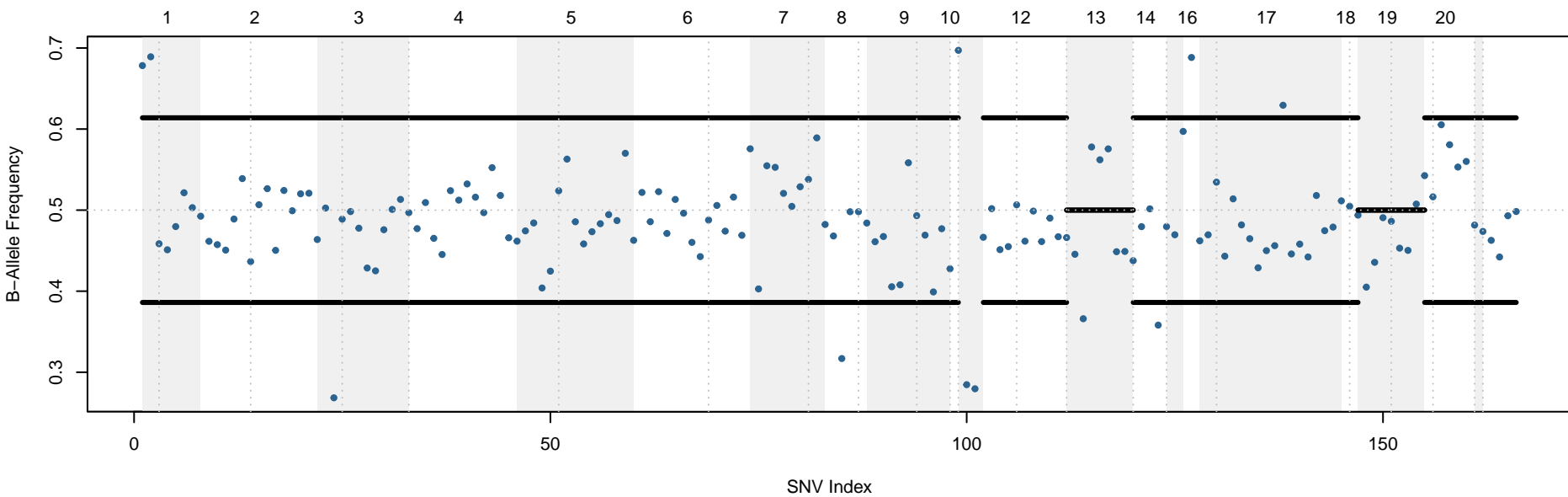




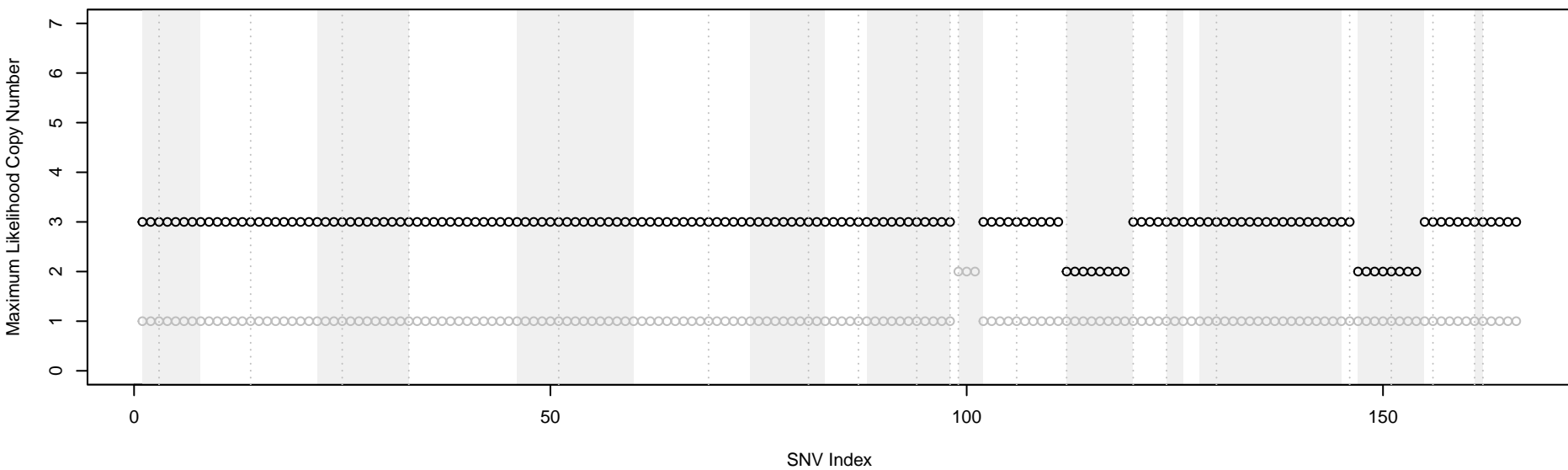
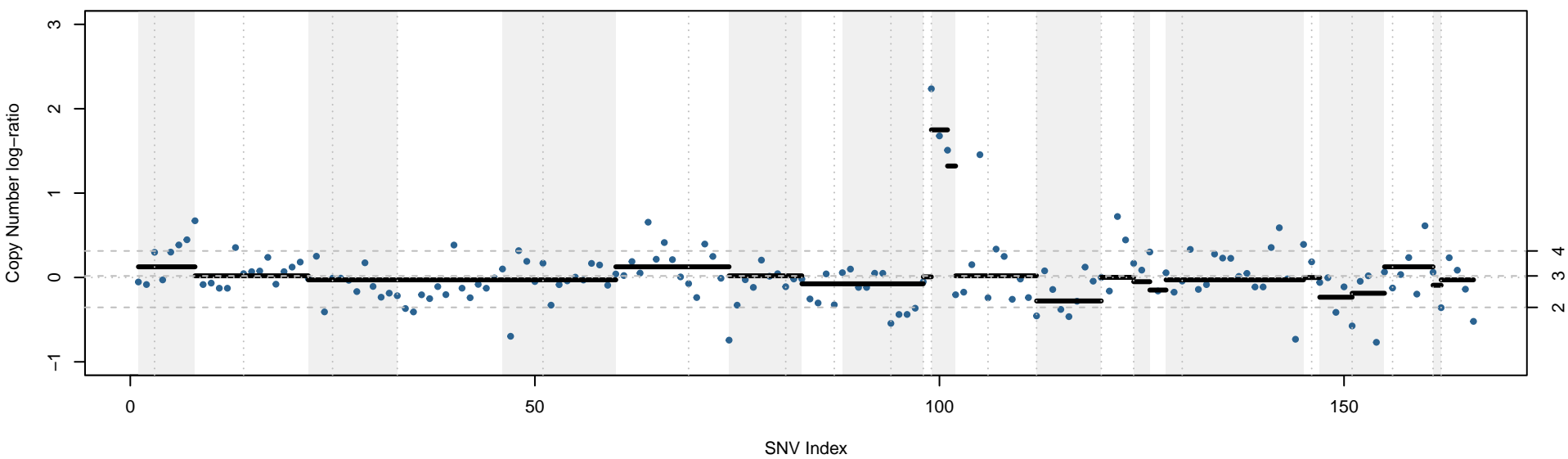
Purity: 0.59 Tumor ploidy: 2.945

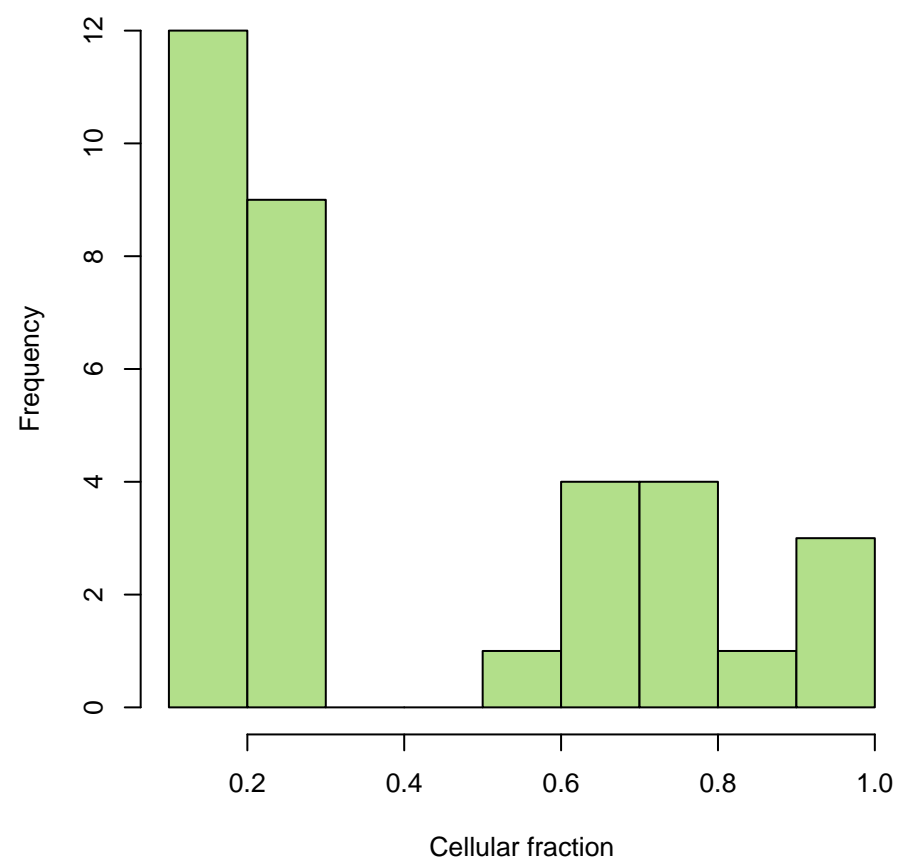
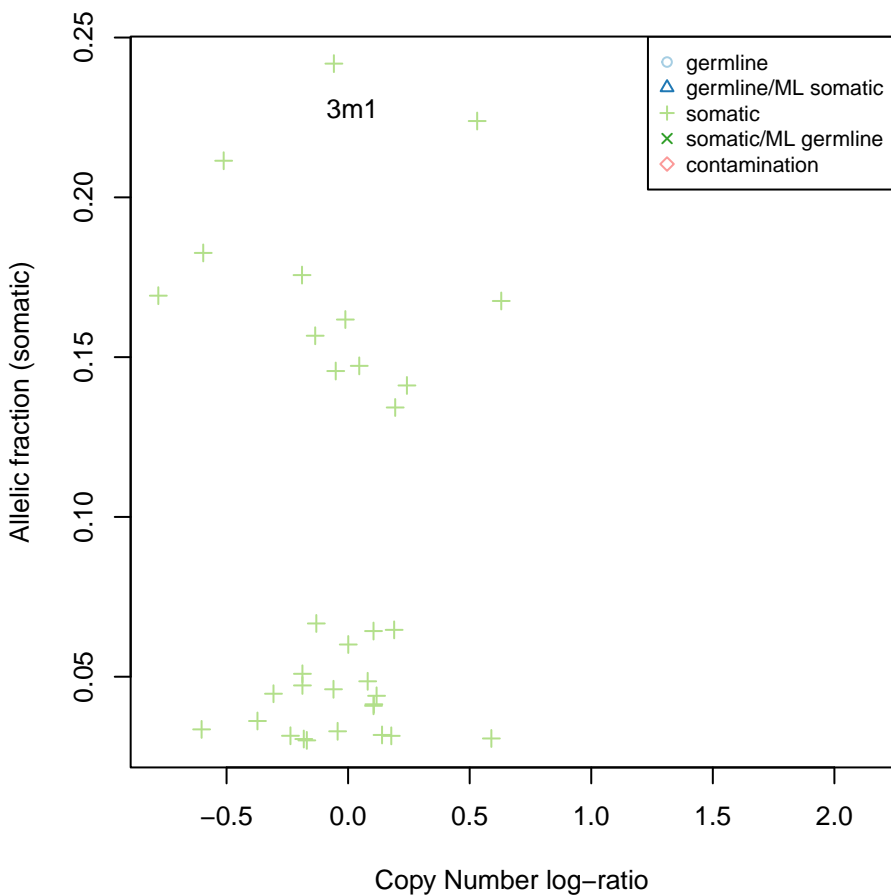
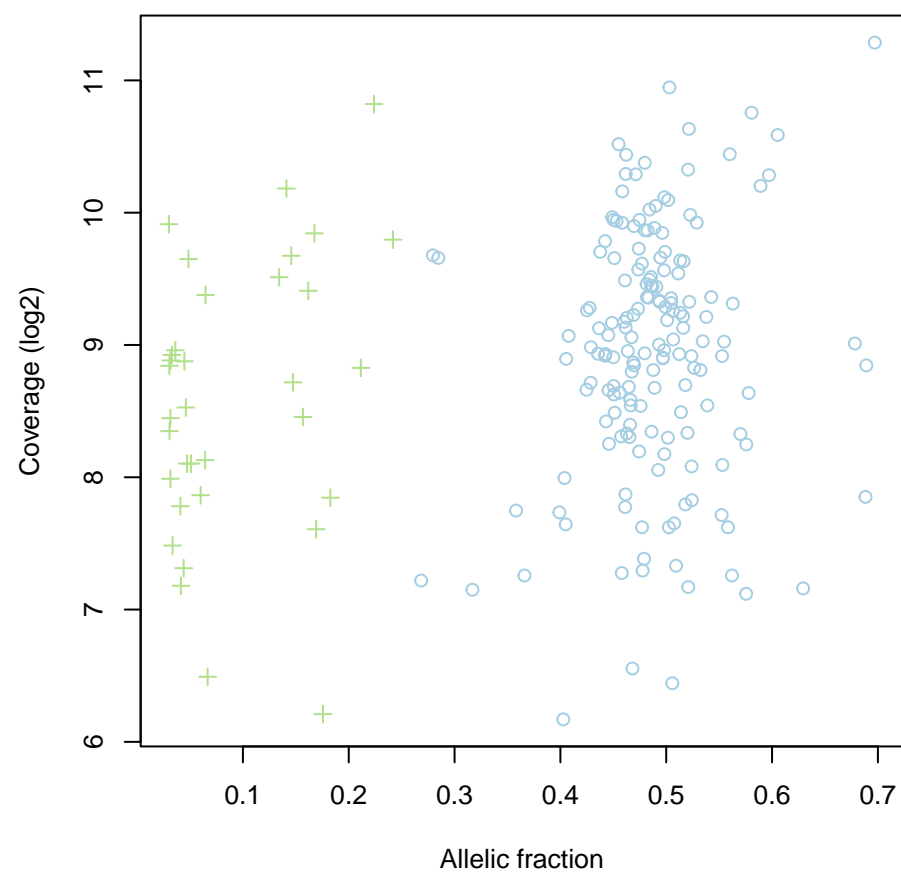
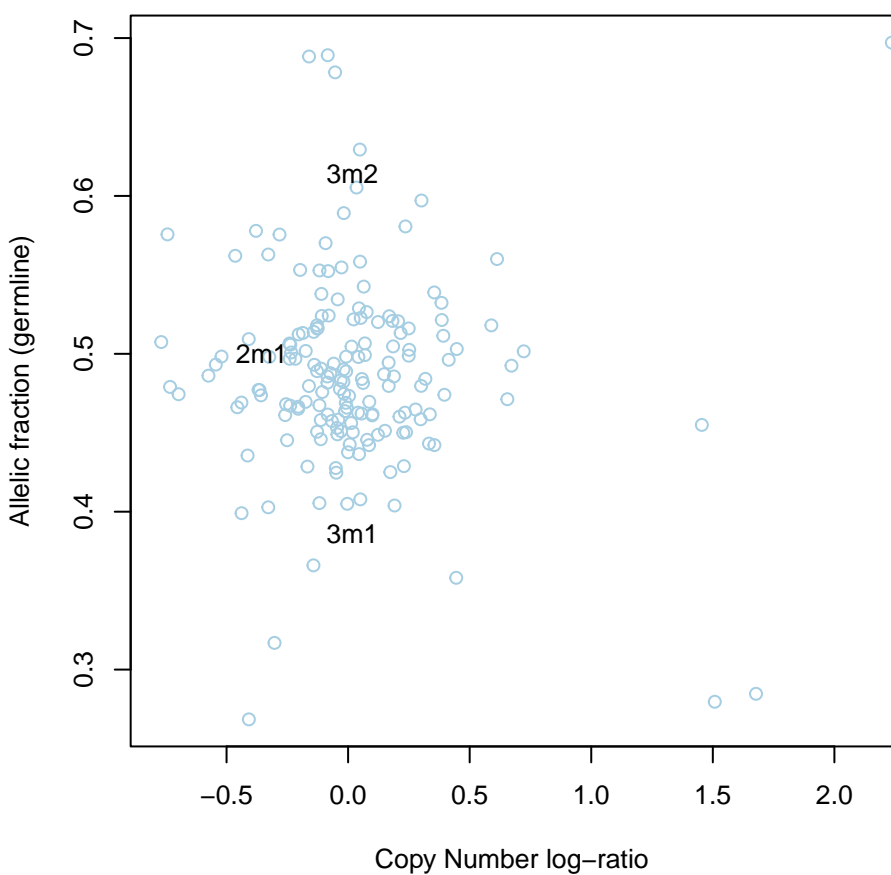


Purity: 0.59 Tumor ploidy: 2.945 SNV log-likelihood: -623.16 GoF: 74.5% Mean coverage: 584,392

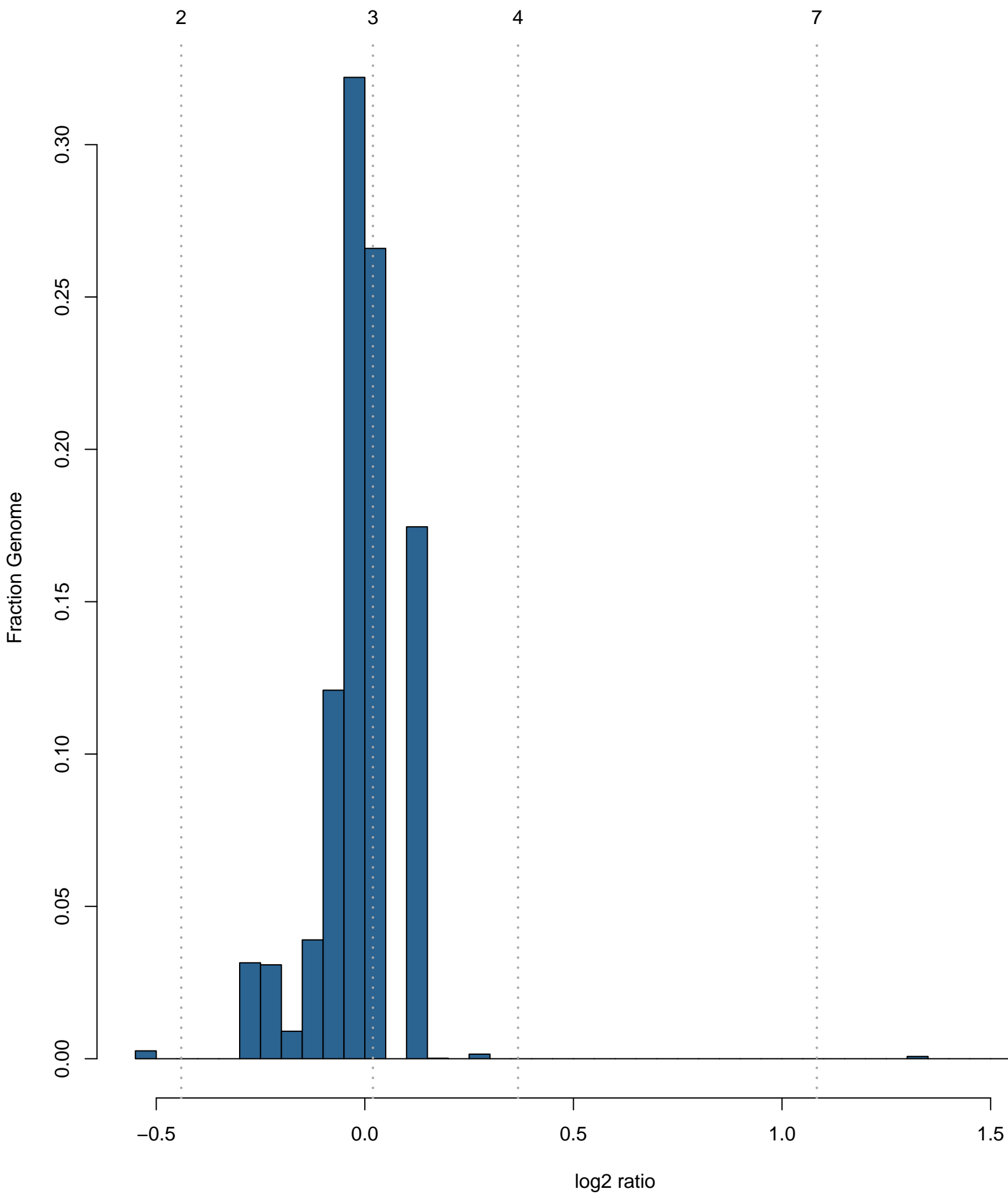


SCNA-fit log-likelihood: -8825.62

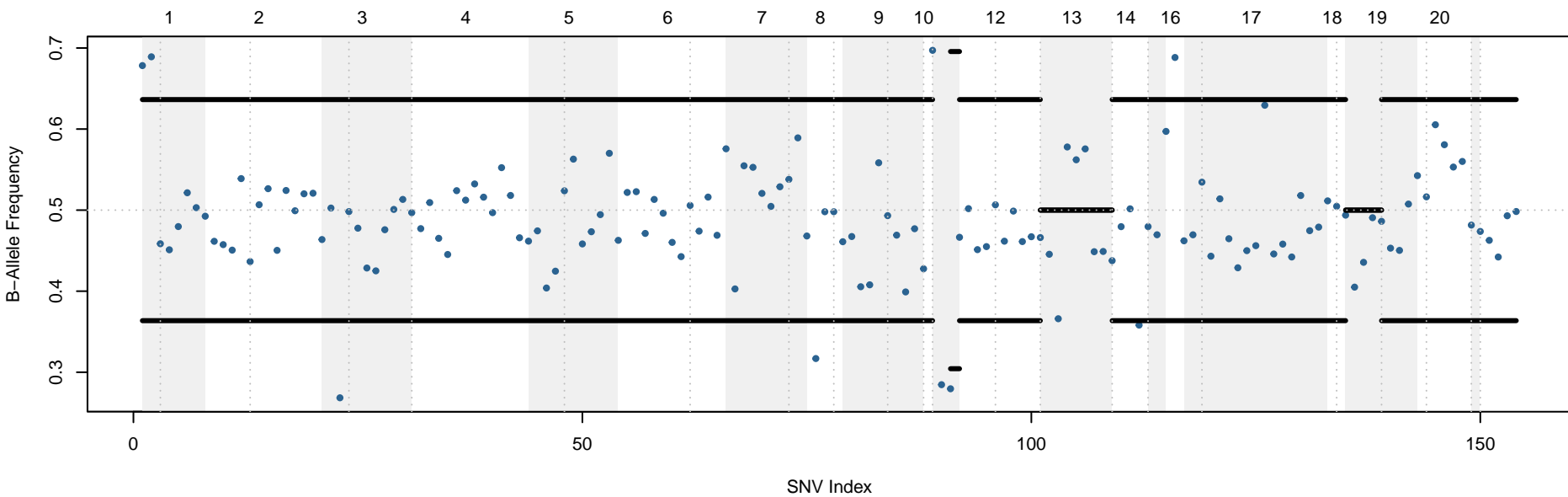




Purity: 0.75 Tumor ploidy: 2.951



Purity: 0.75 Tumor ploidy: 2.951 SNV log-likelihood: -1031.94 GoF: 63% Mean coverage: 584;392



SCNA-fit log-likelihood: -8855.14

