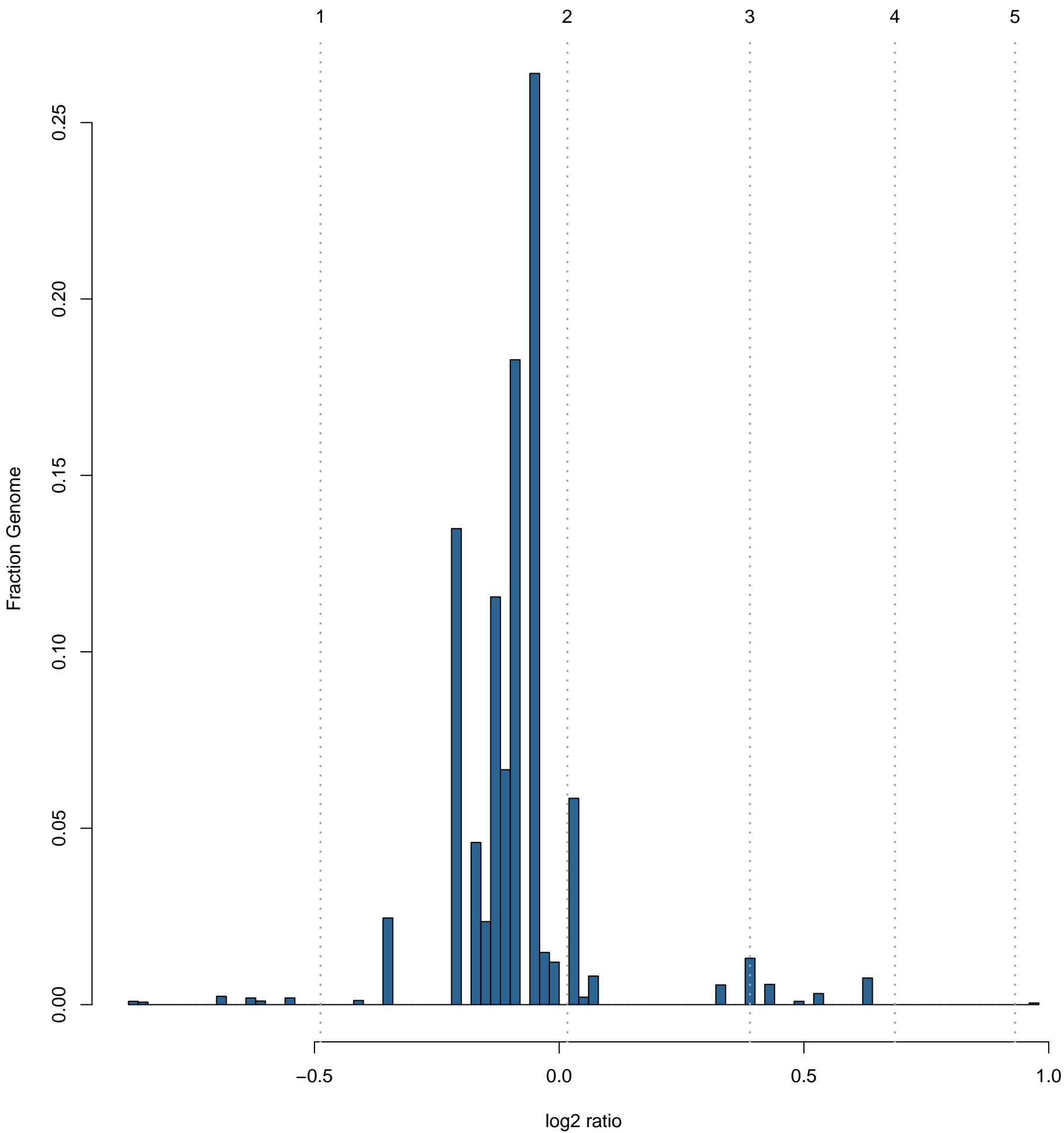
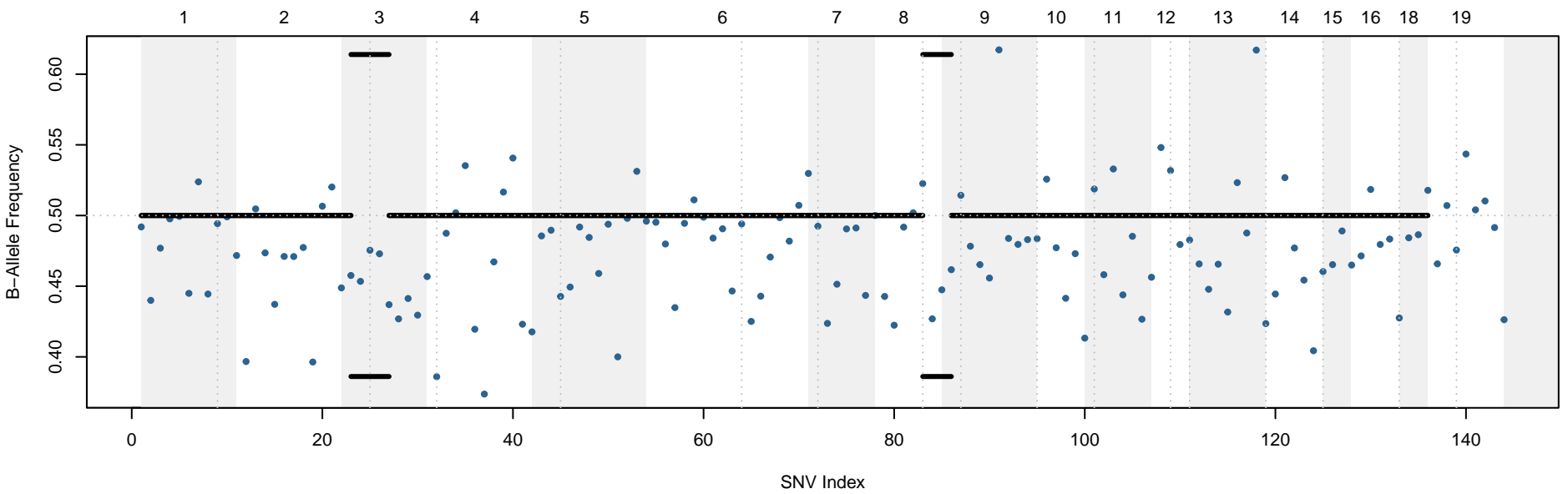


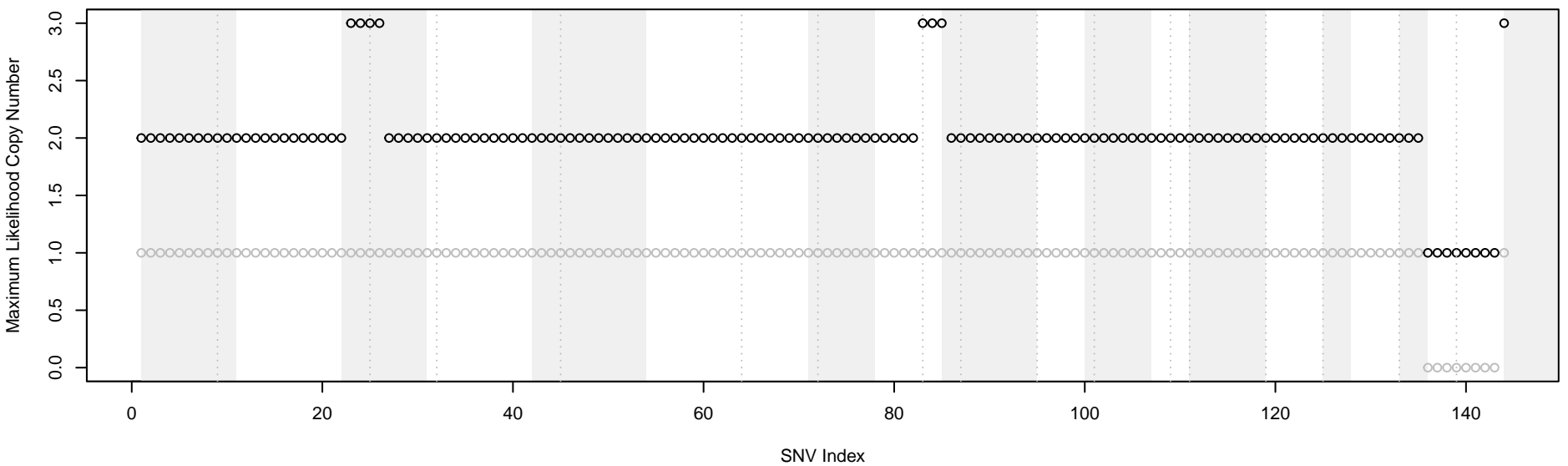
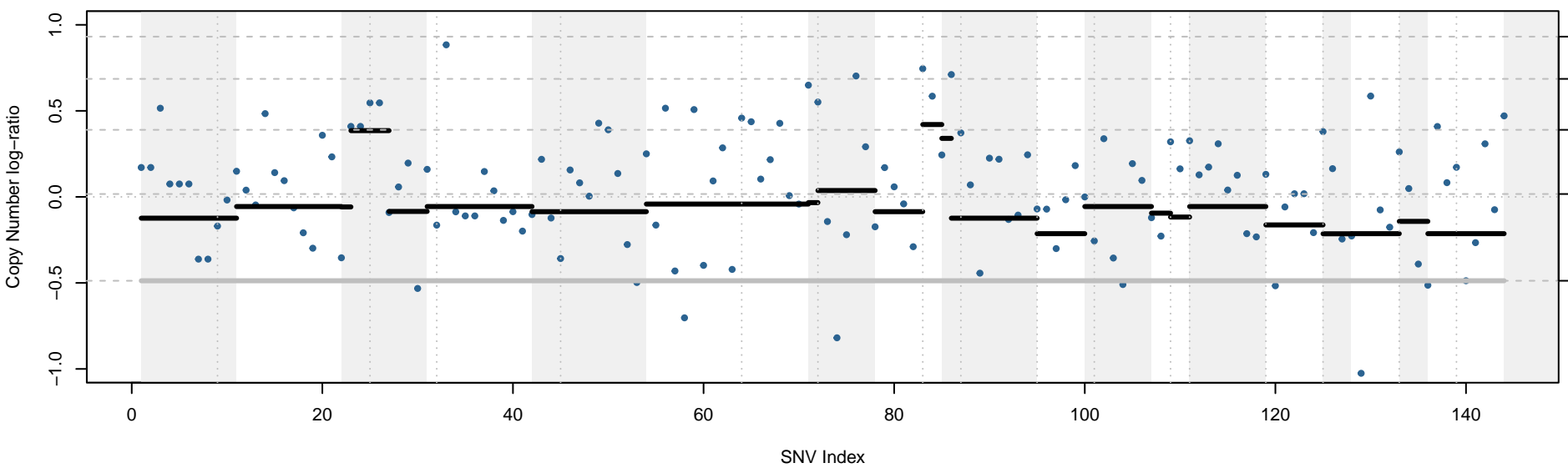
Purity: 0.59 Tumor ploidy: 1.961

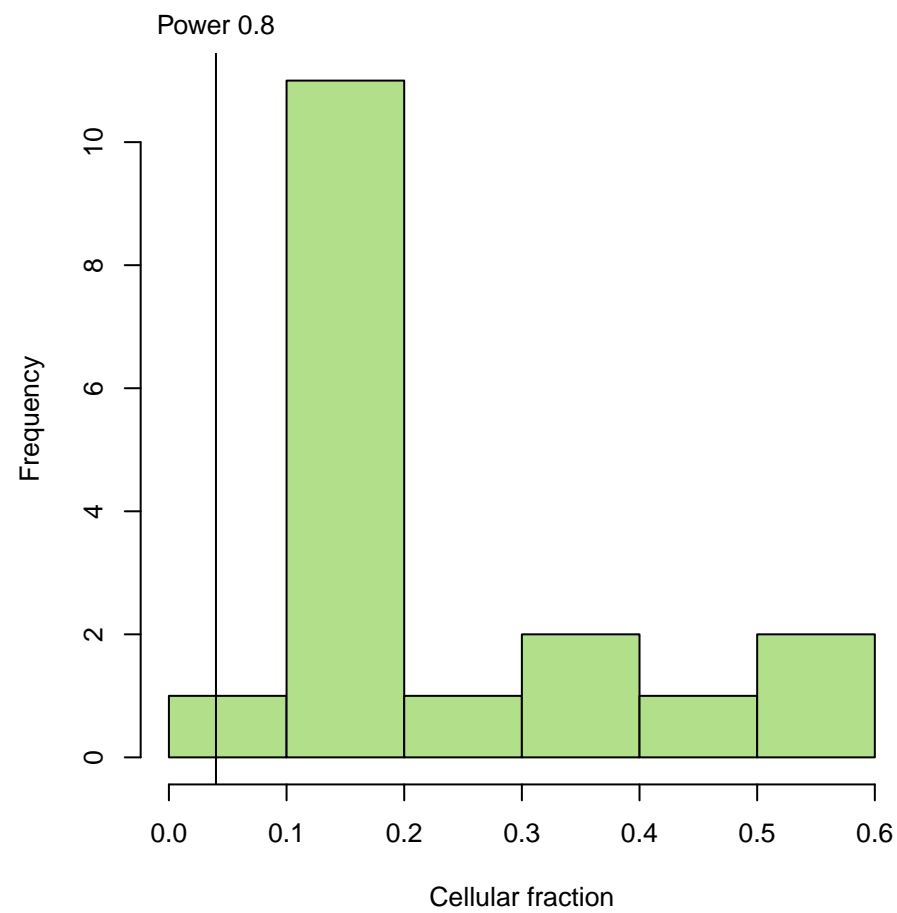
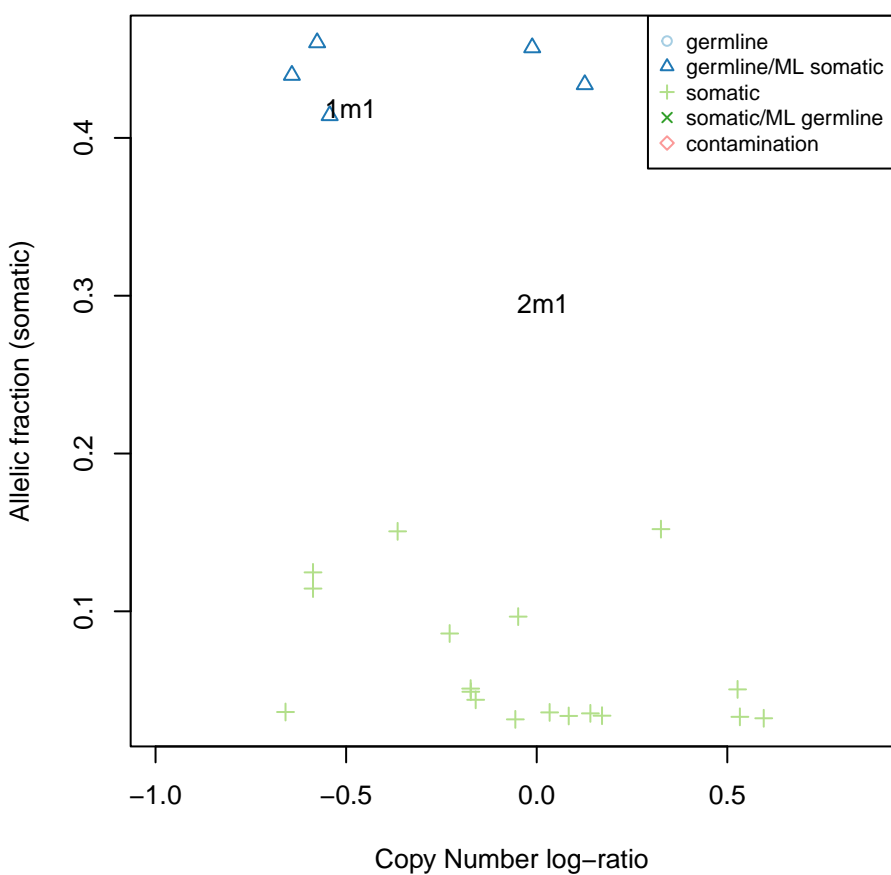
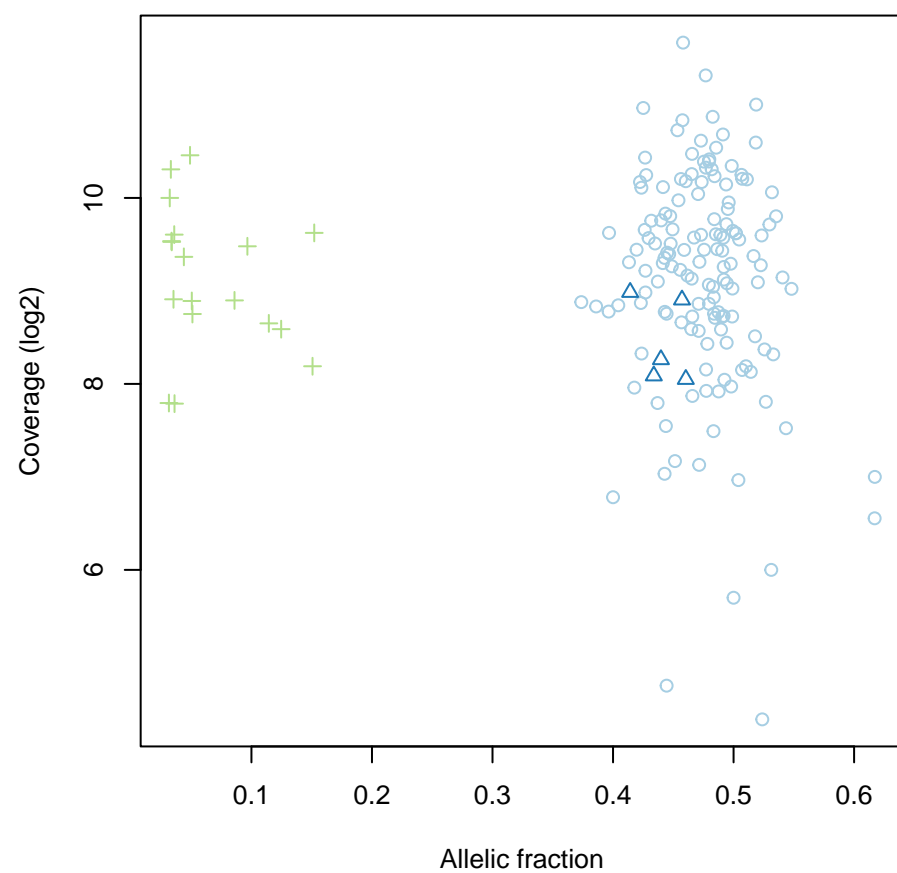
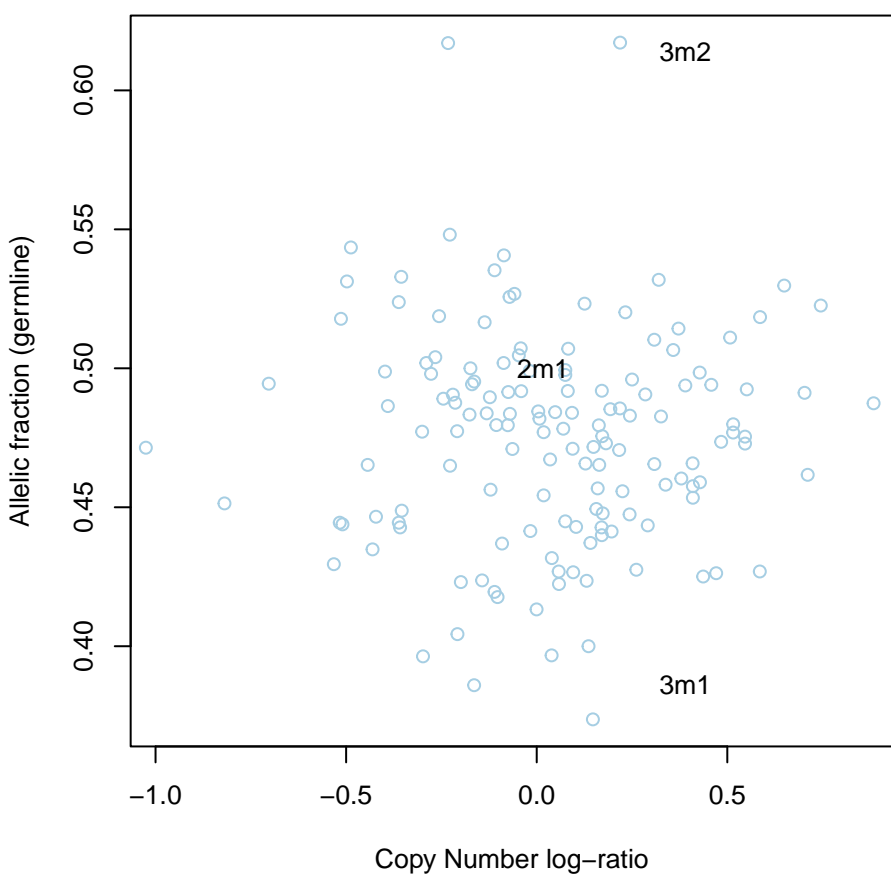


Purity: 0.59 Tumor ploidy: 1.961 SNV log-likelihood: -154.68 GoF: 74.6% Mean coverage: 714,414

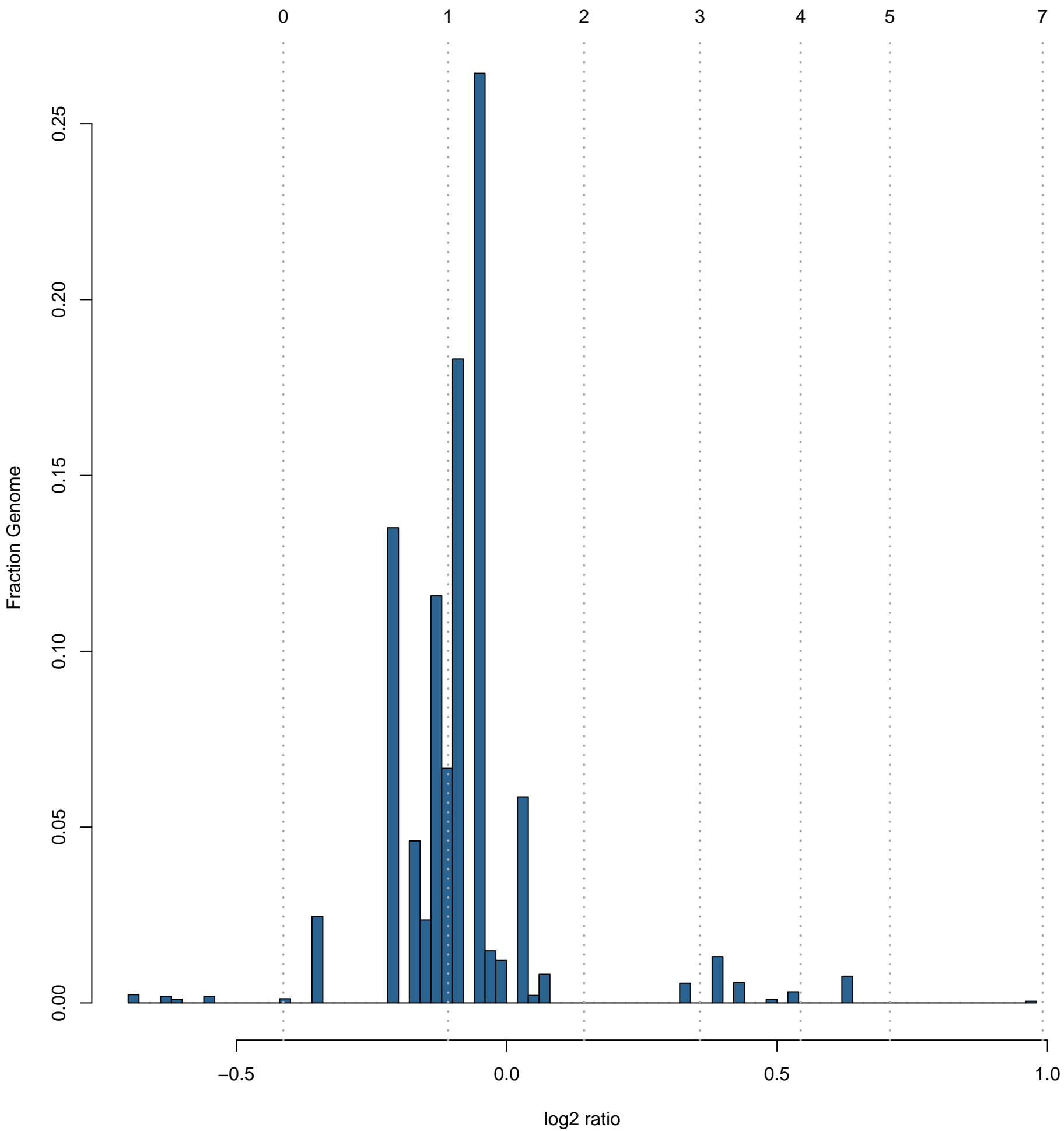


SCNA-fit log-likelihood: -9465.69

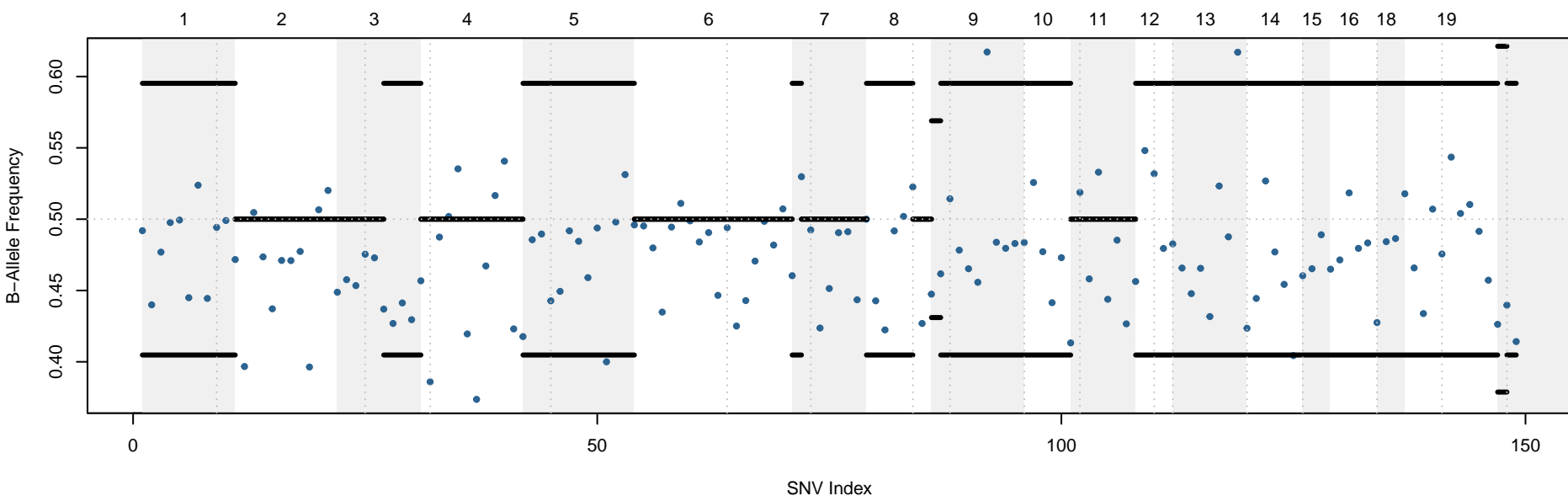




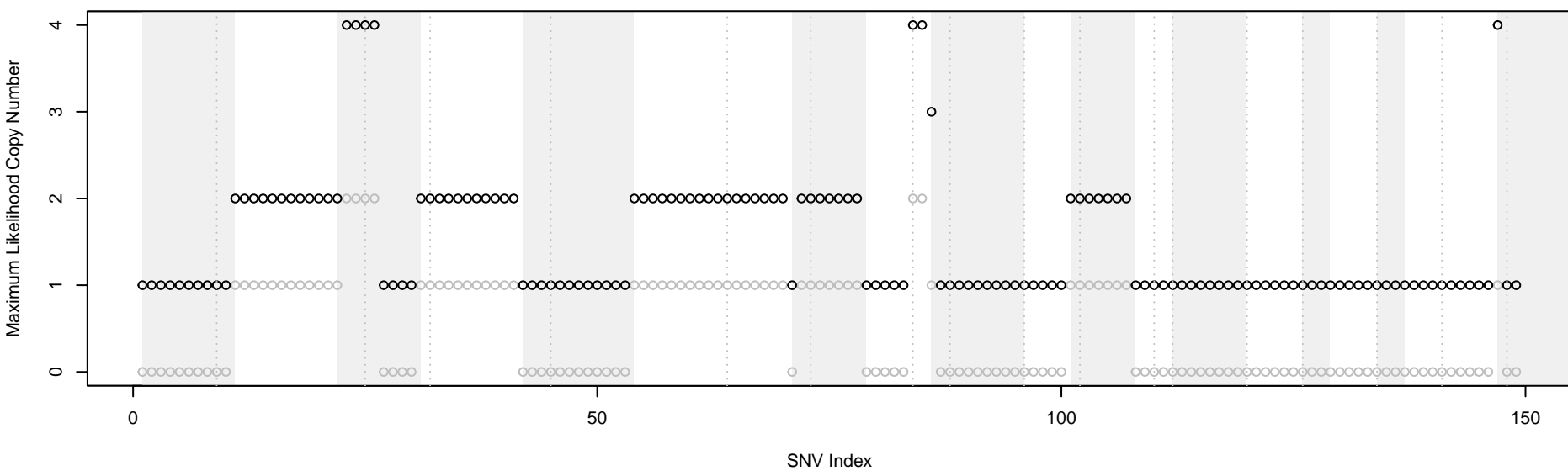
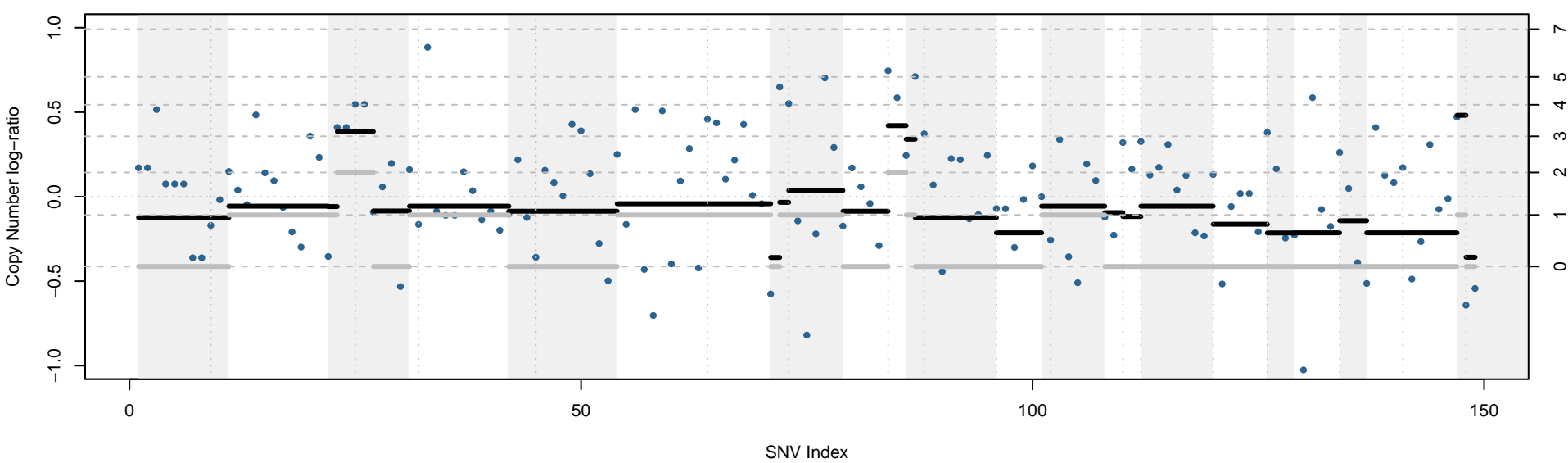
Purity: 0.32 Tumor ploidy: 1.409

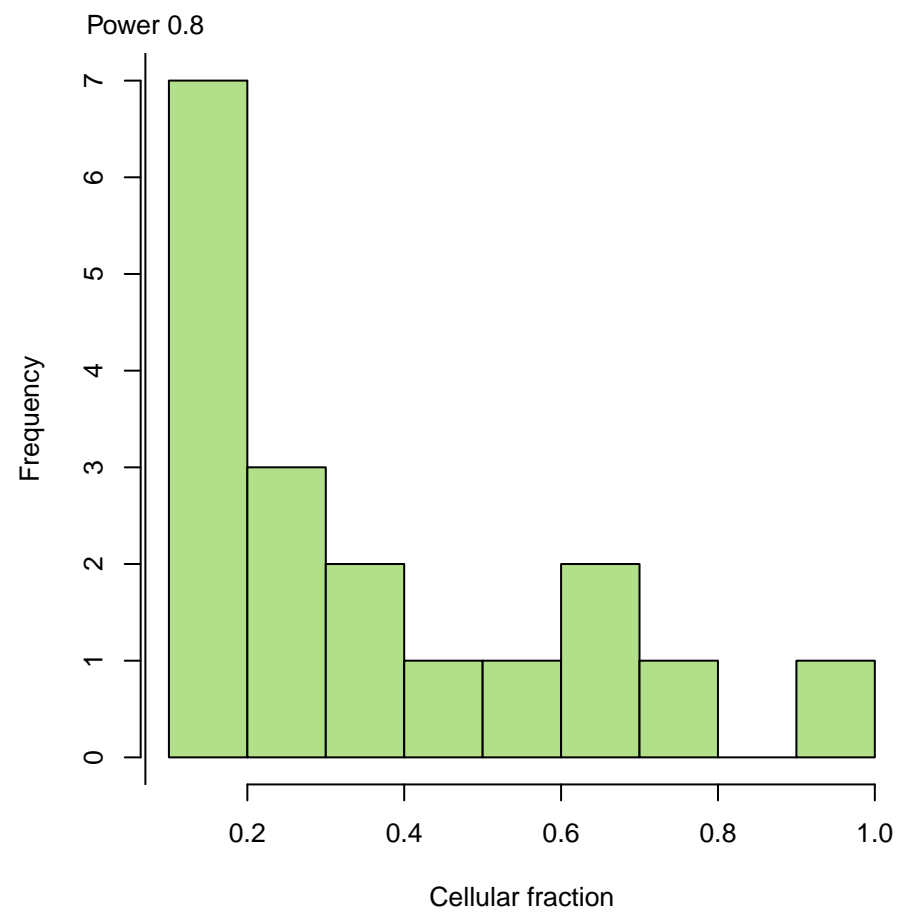
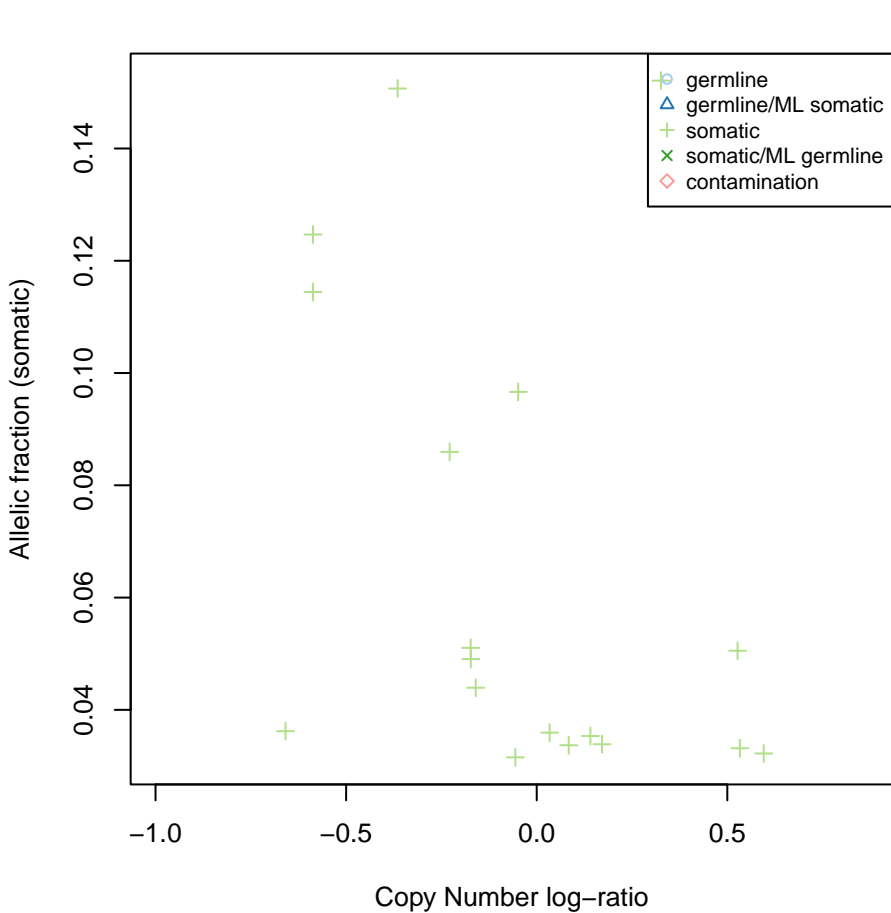
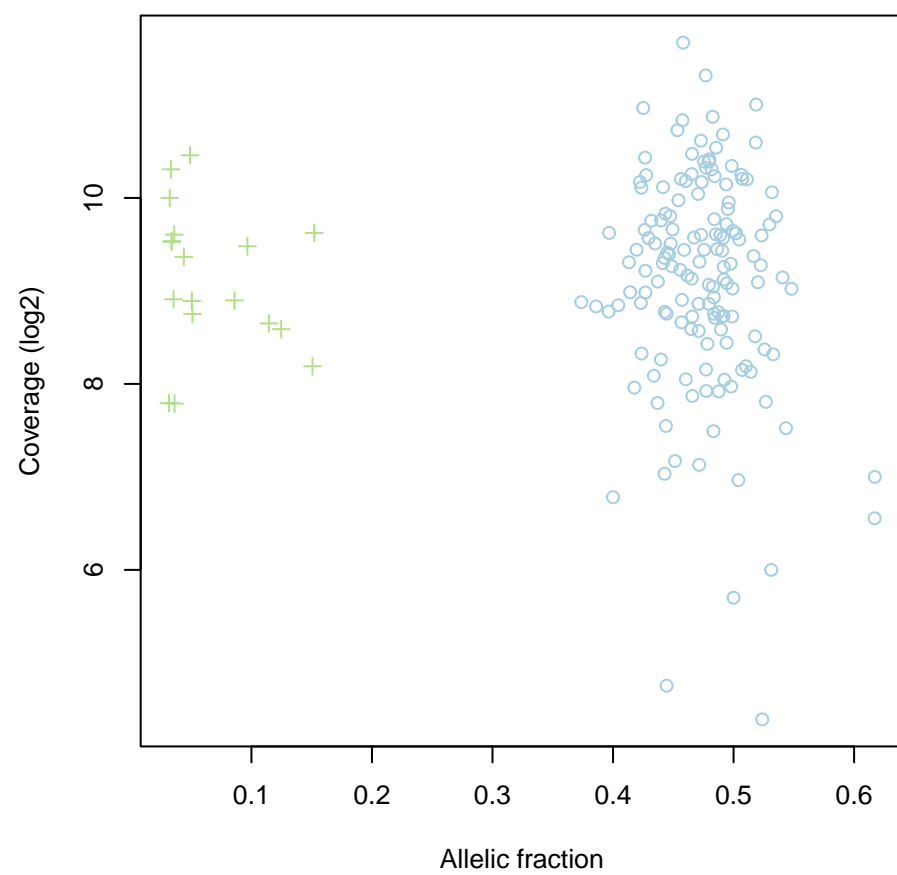
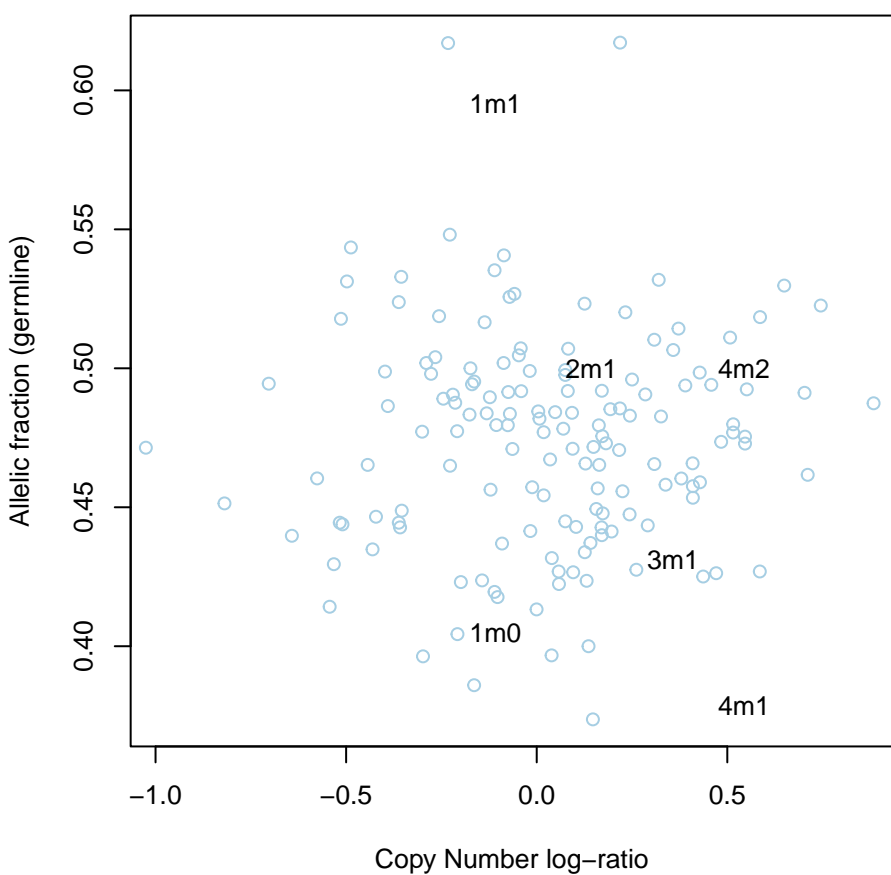


Purity: 0.32 Tumor ploidy: 1.409 SNV log-likelihood: -210.08 GoF: 88.5% Mean coverage: 714;414

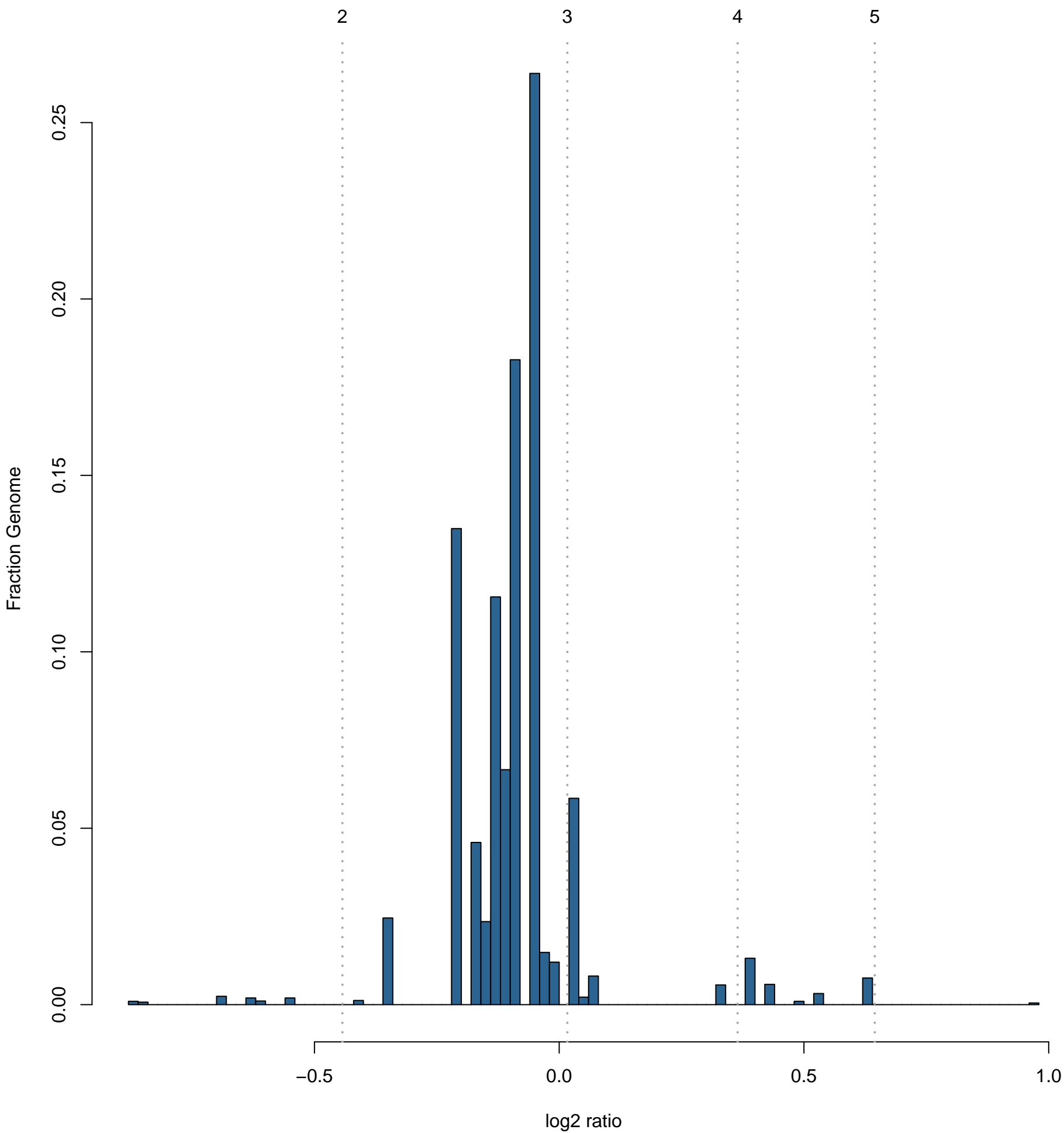


SCNA-fit log-likelihood: -9385.13

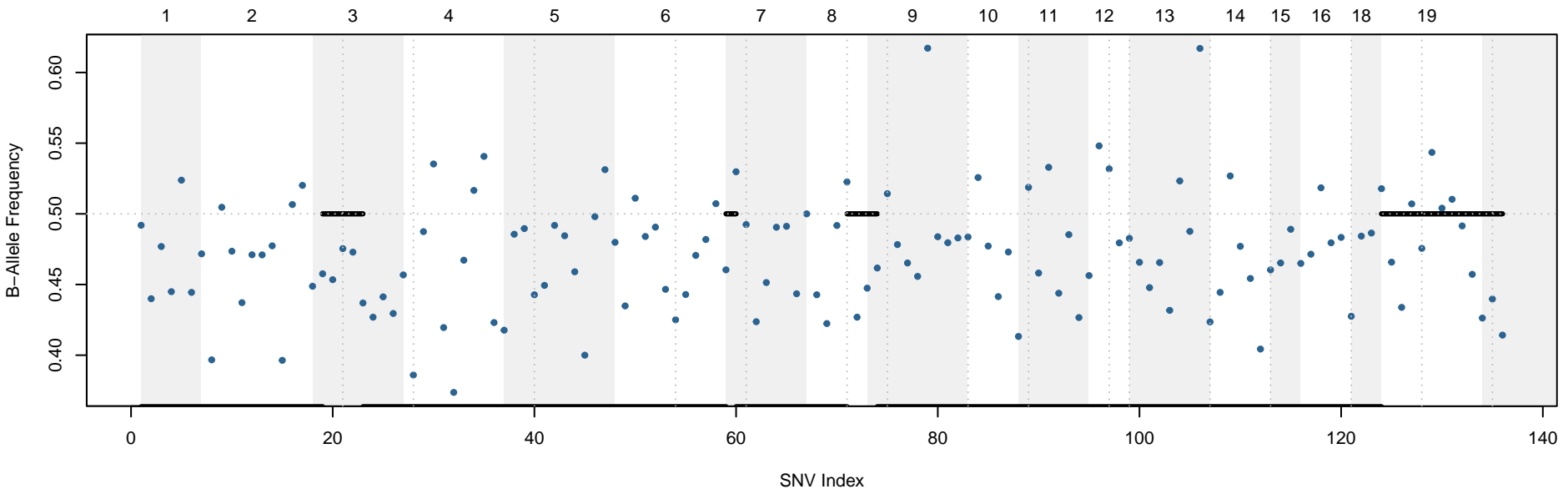




Purity: 0.75 Tumor ploidy: 2.958



Purity: 0.75 Tumor ploidy: 2.958 SNV log-likelihood: -836.56 GoF: 68.3% Mean coverage: 714,414



SCNA-fit log-likelihood: -9428.83

