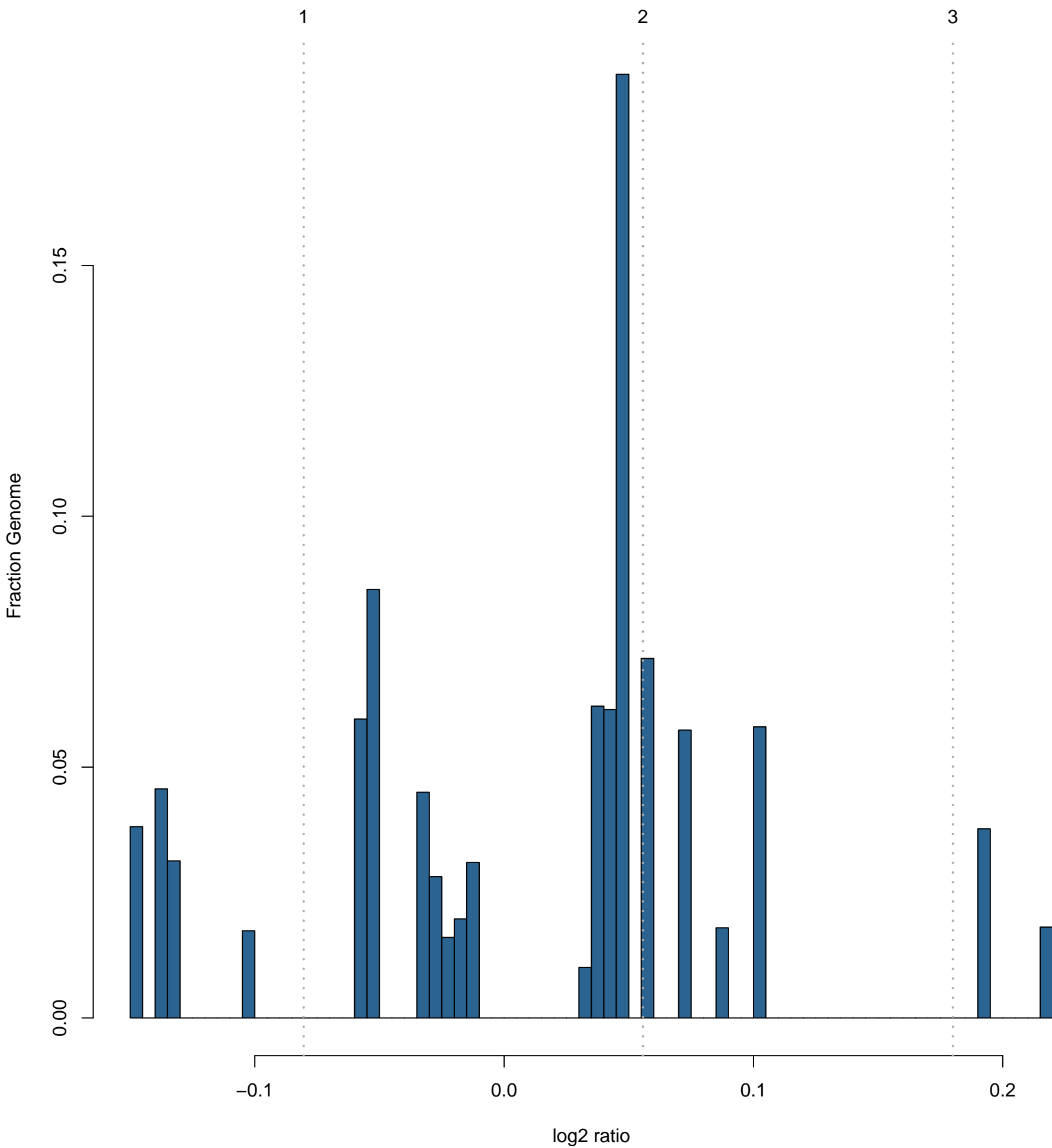
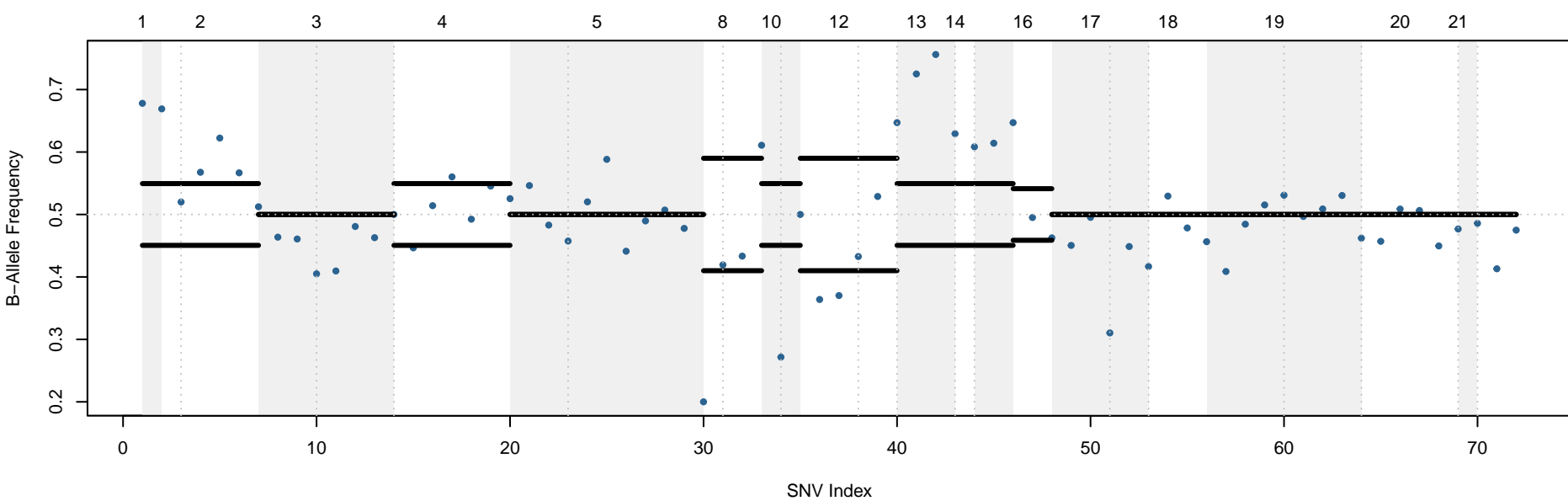


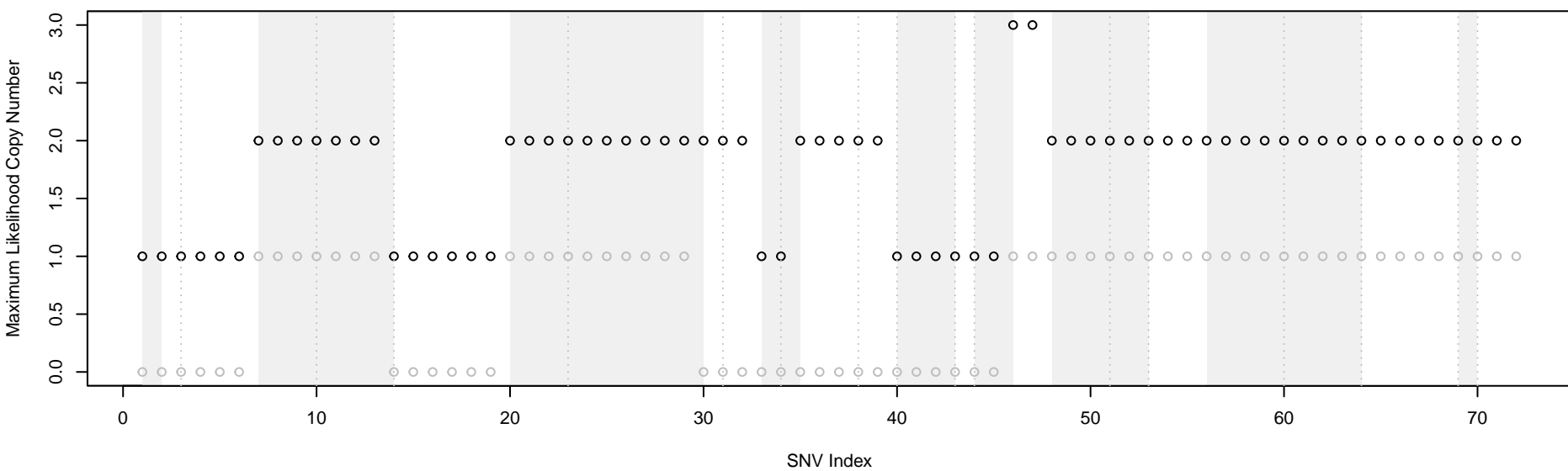
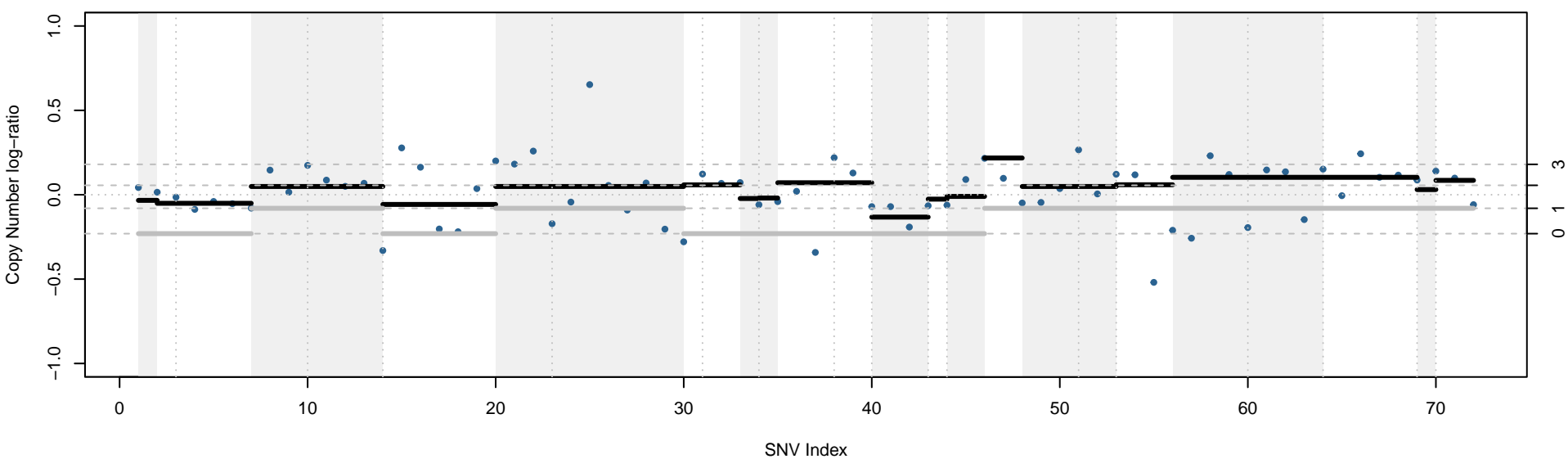
Purity: 0.18 Tumor ploidy: 1.579

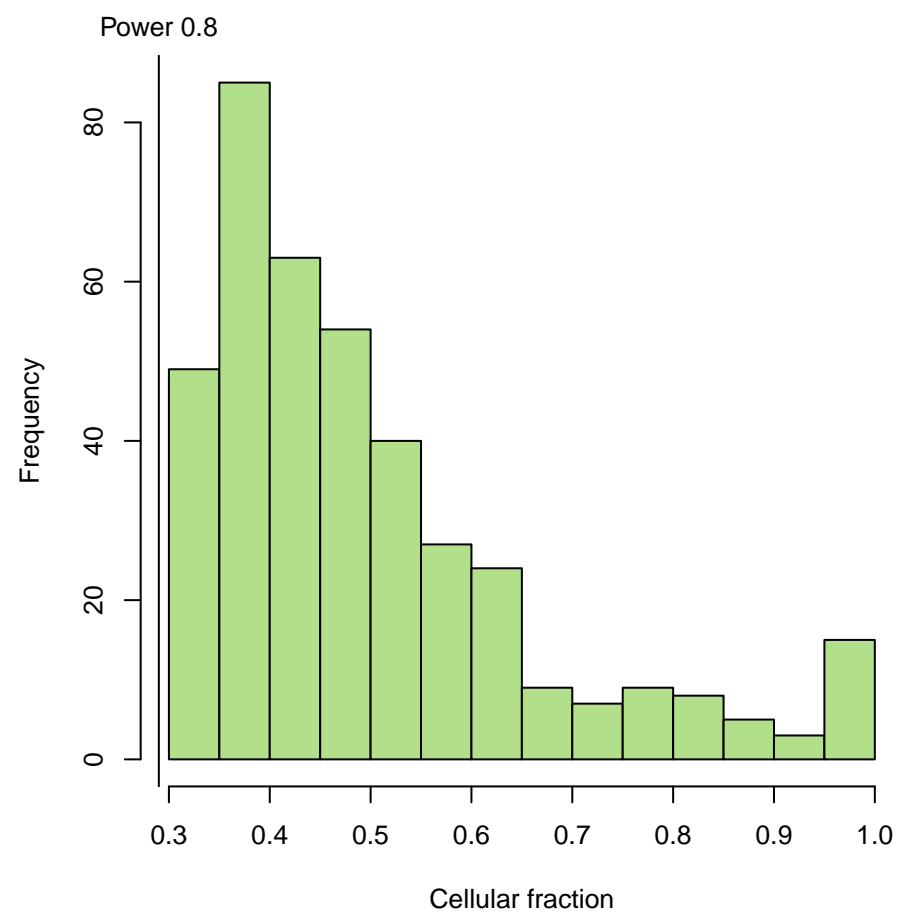
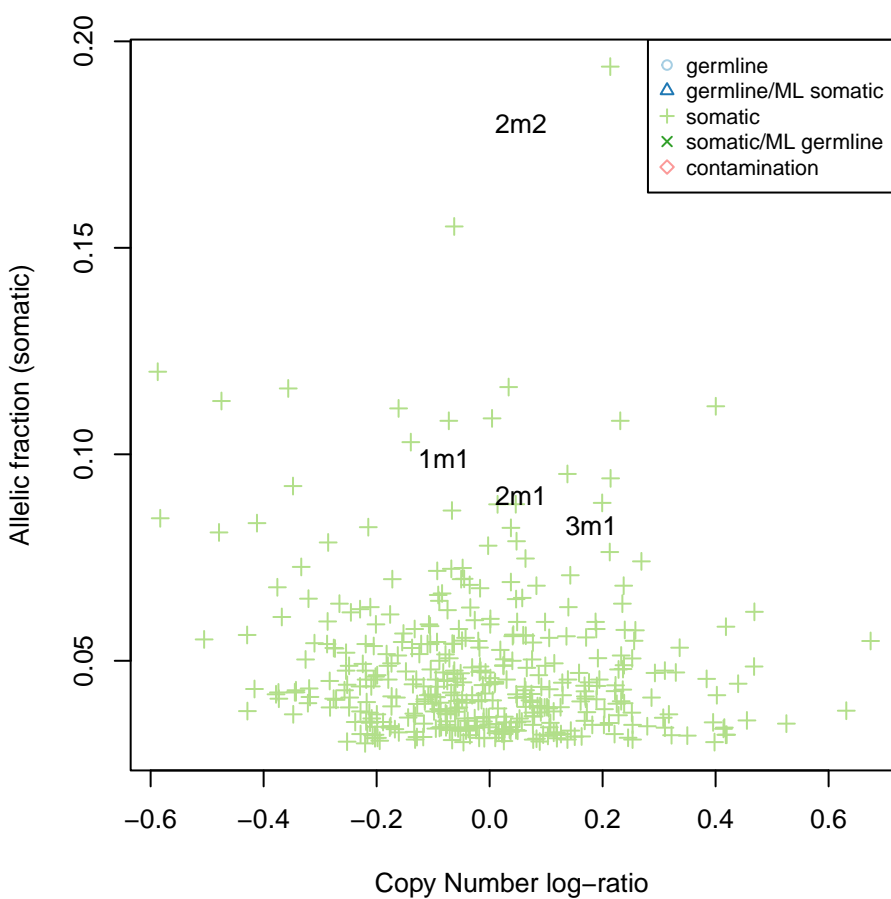
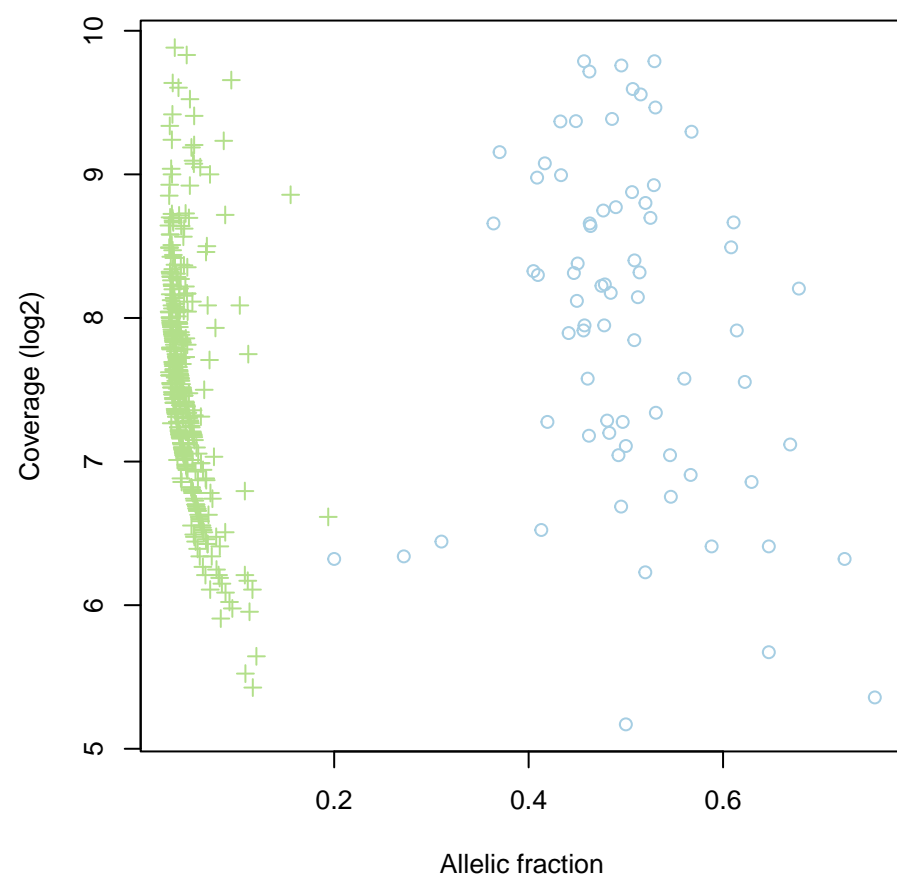
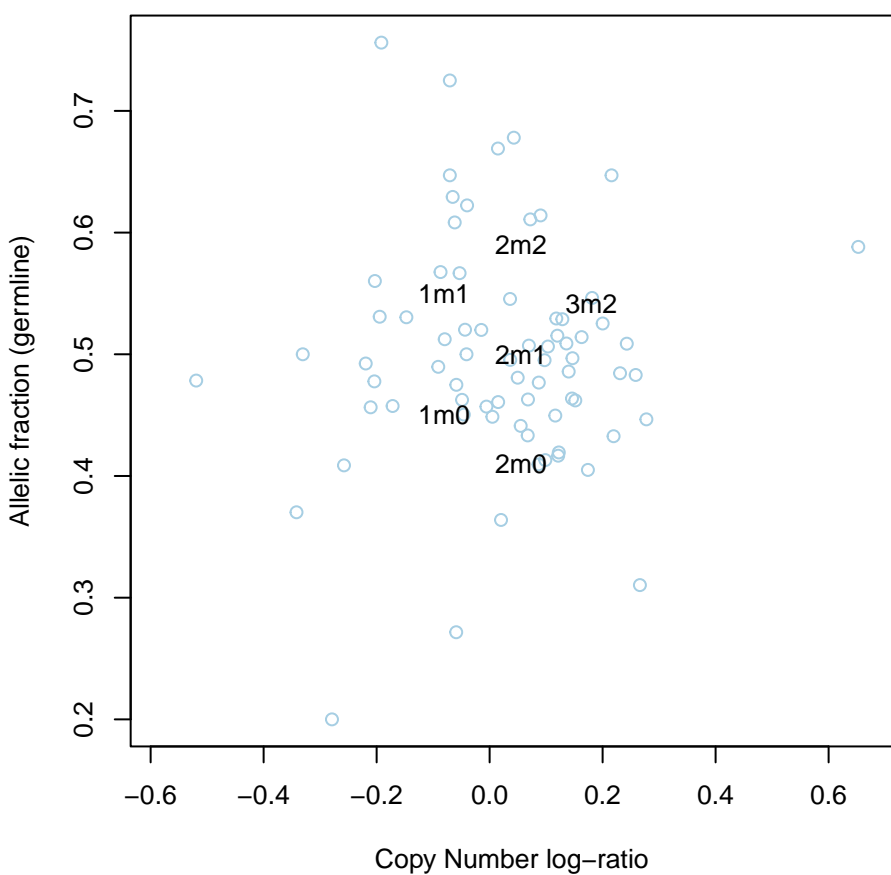


Purity: 0.18 Tumor ploidy: 1.579 SNV log-likelihood: 401.64 GoF: 92.9% Mean coverage: 233;281

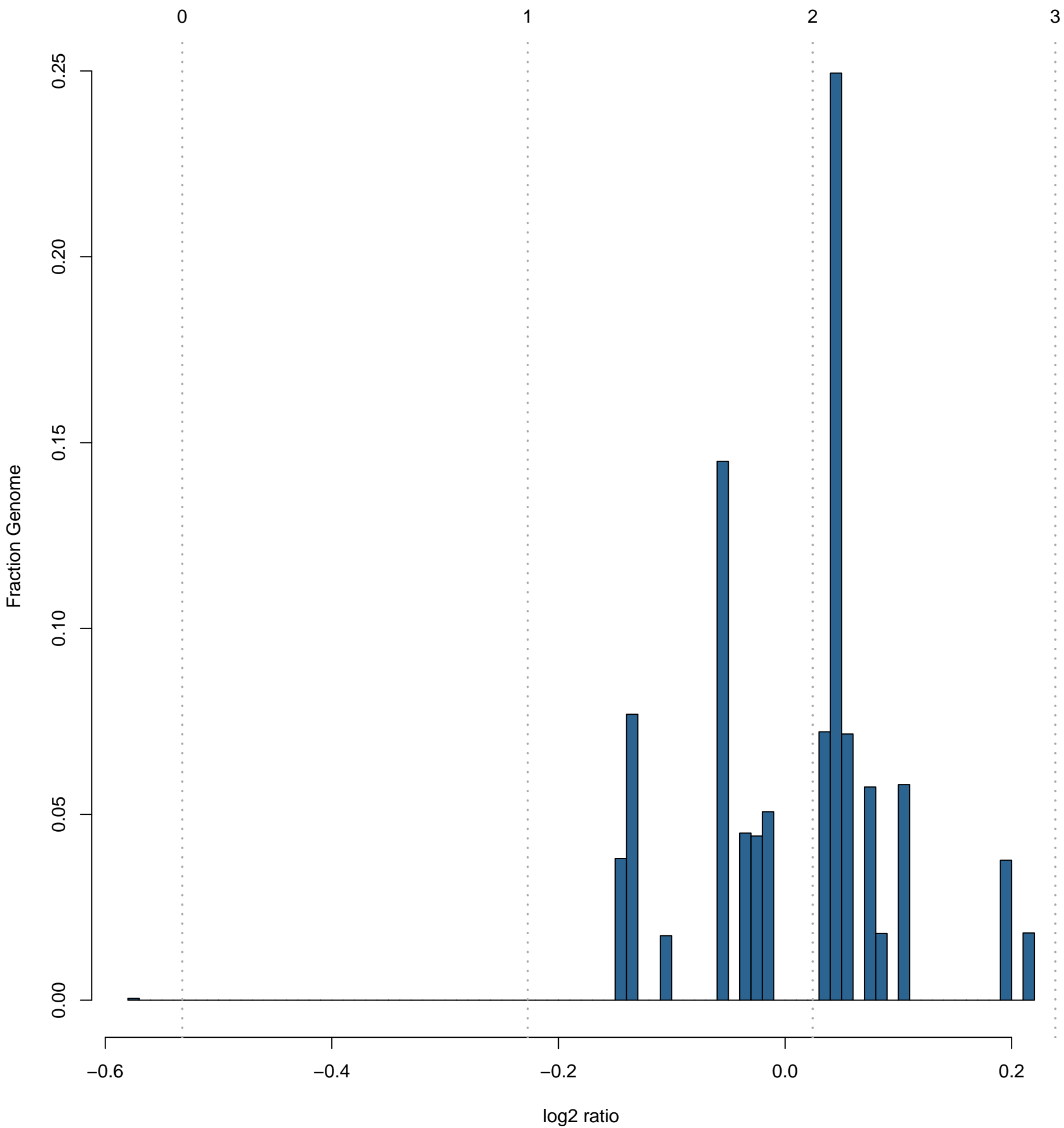


SCNA-fit log-likelihood: -5119.5

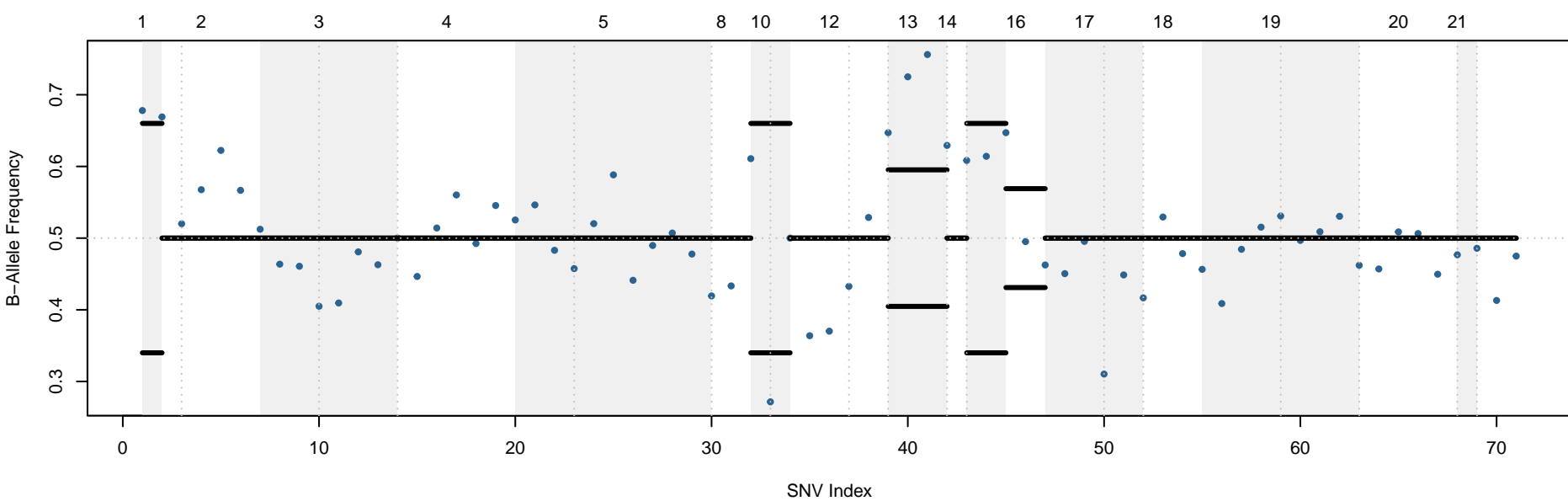




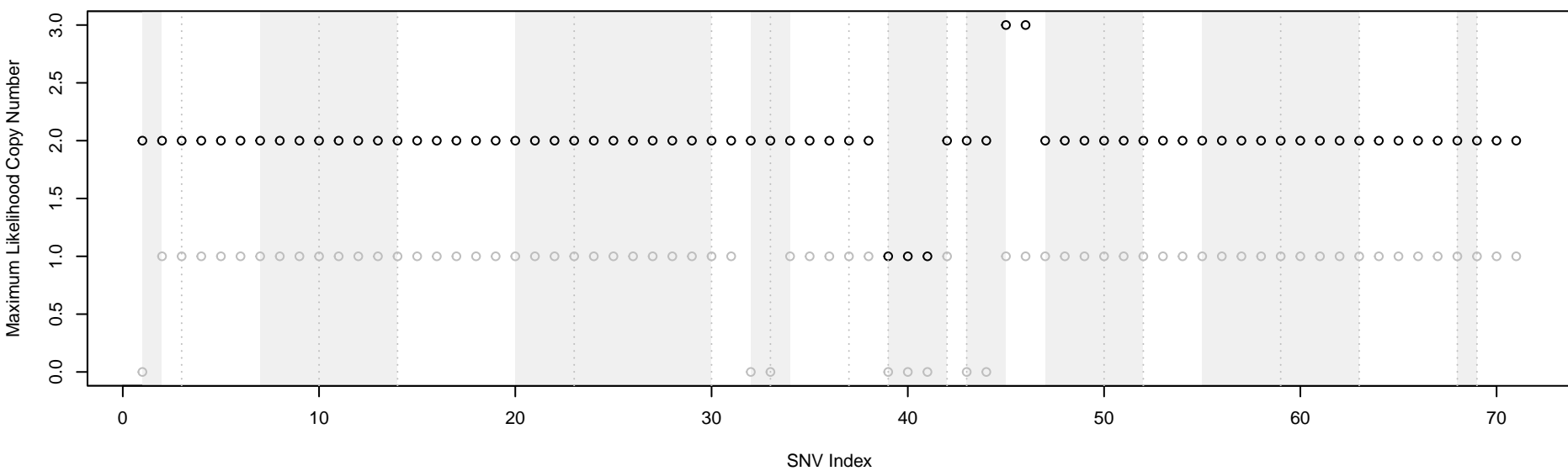
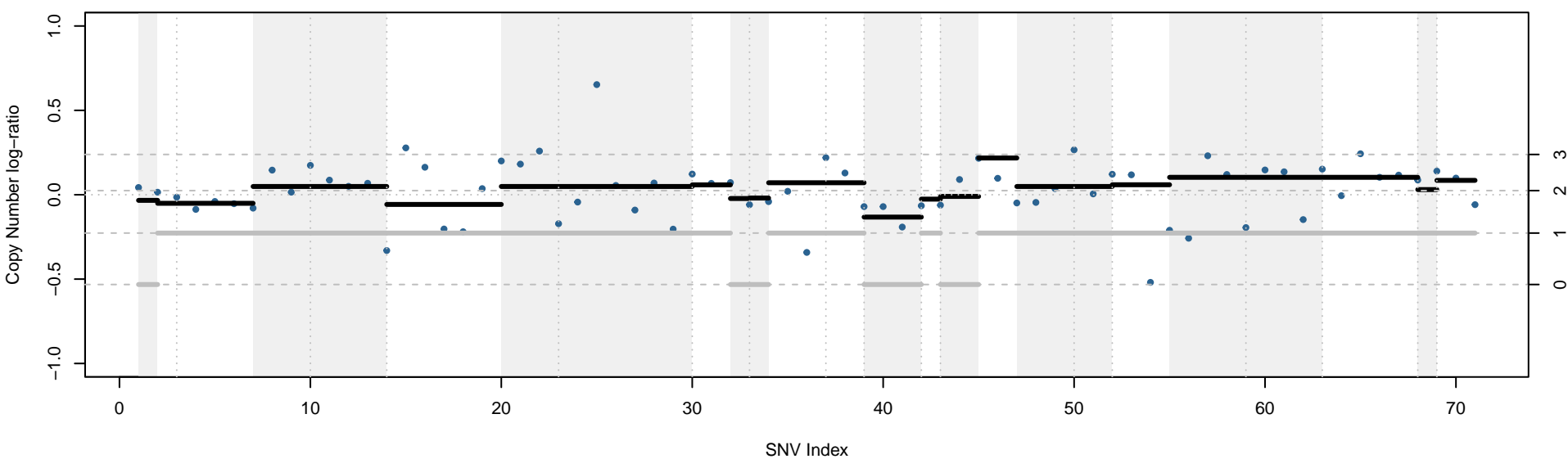
Purity: 0.32 Tumor ploidy: 1.895

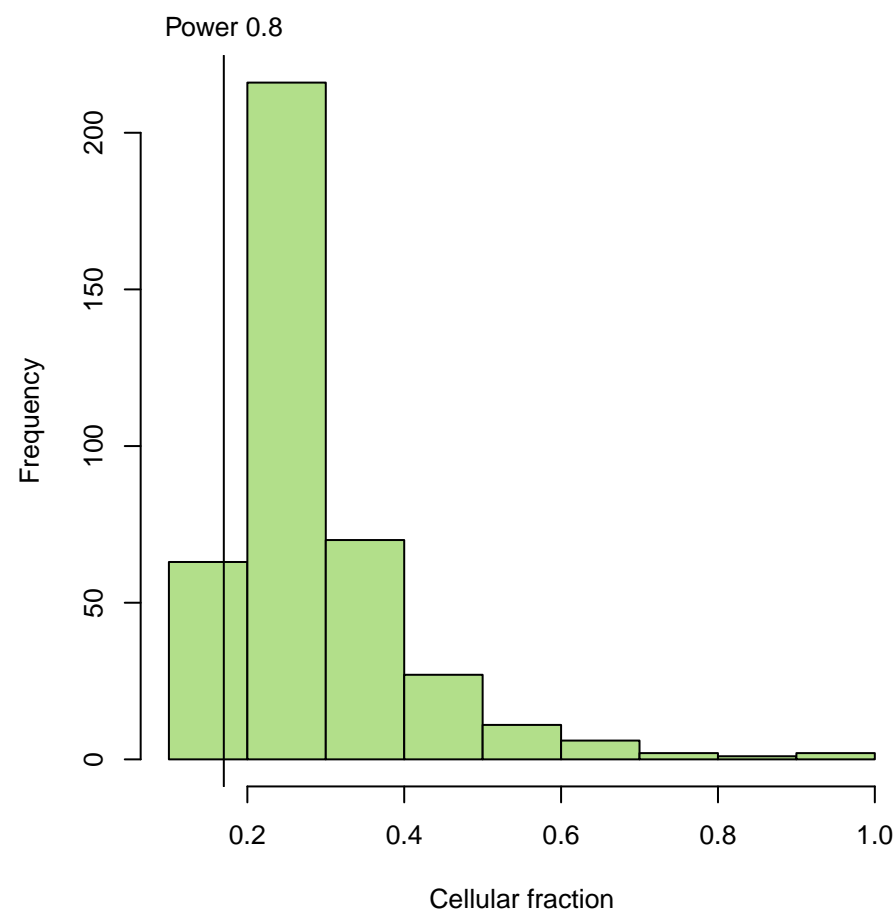
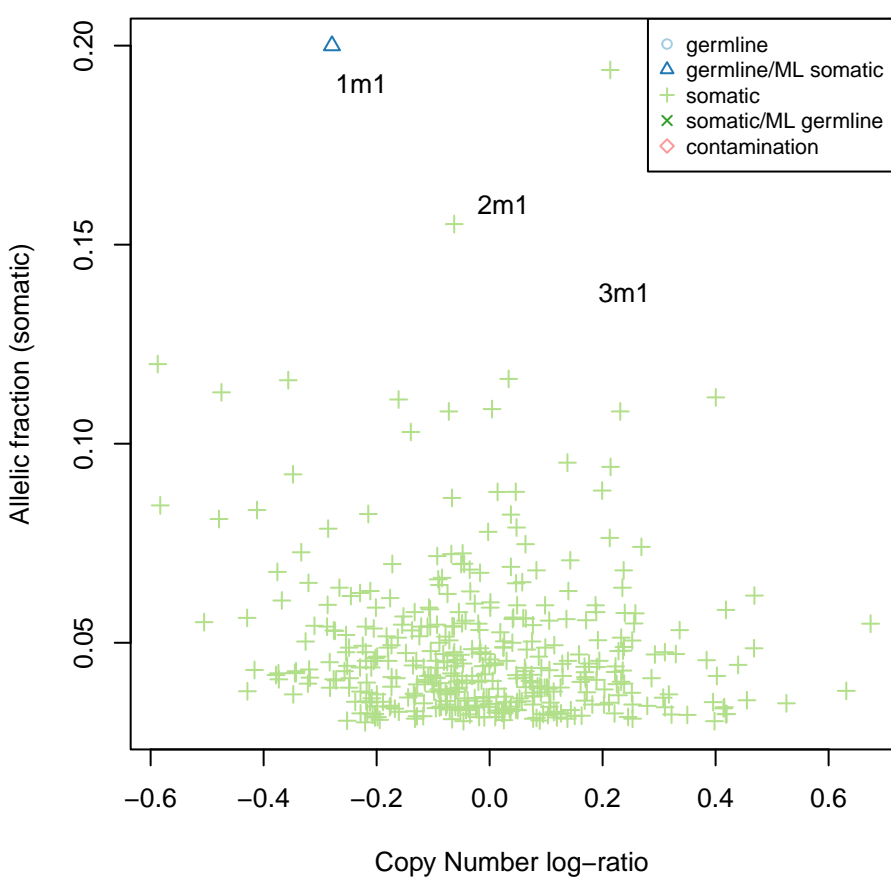
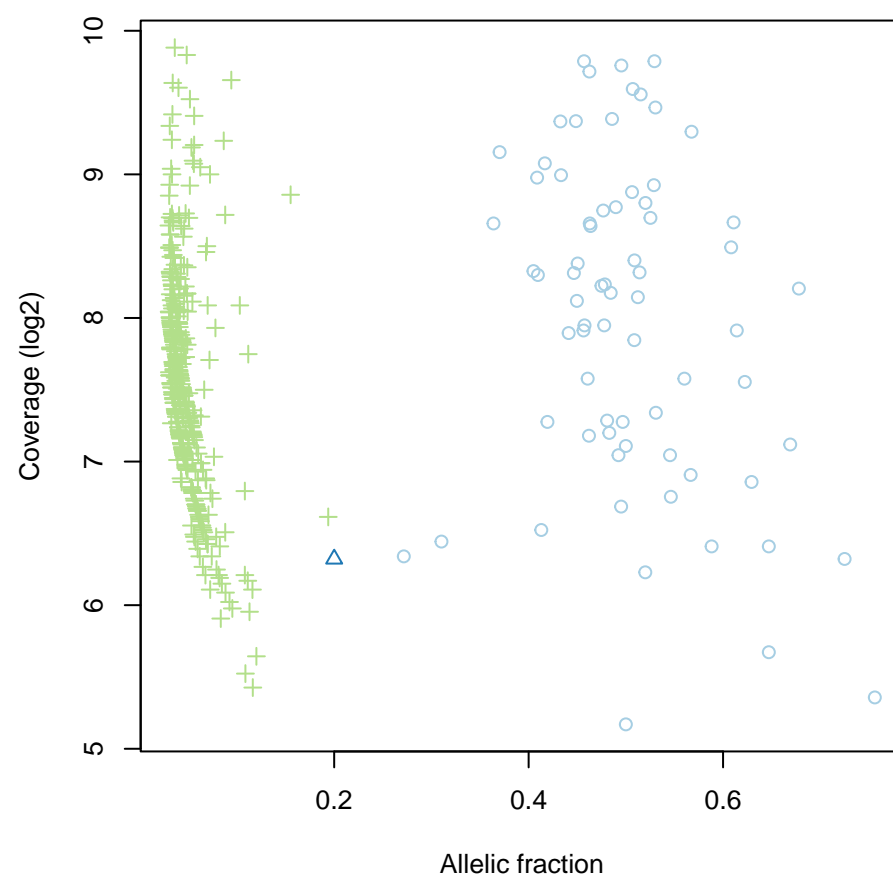
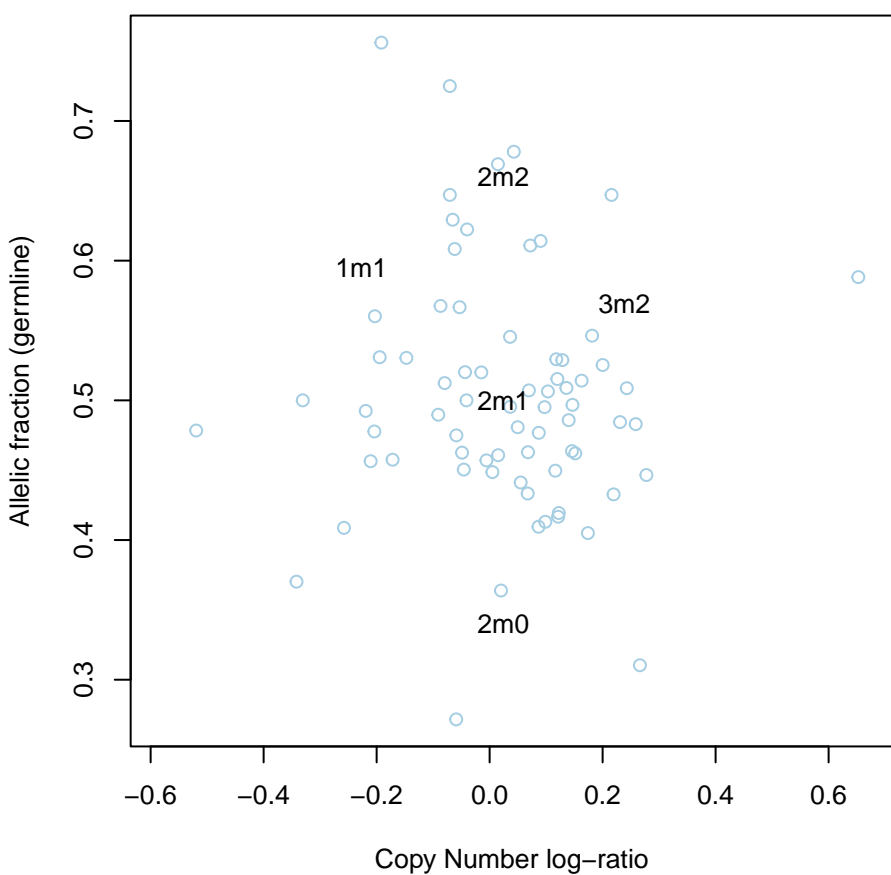


Purity: 0.32 Tumor ploidy: 1.895 SNV log-likelihood: 304.78 GoF: 68.8% Mean coverage: 233;281

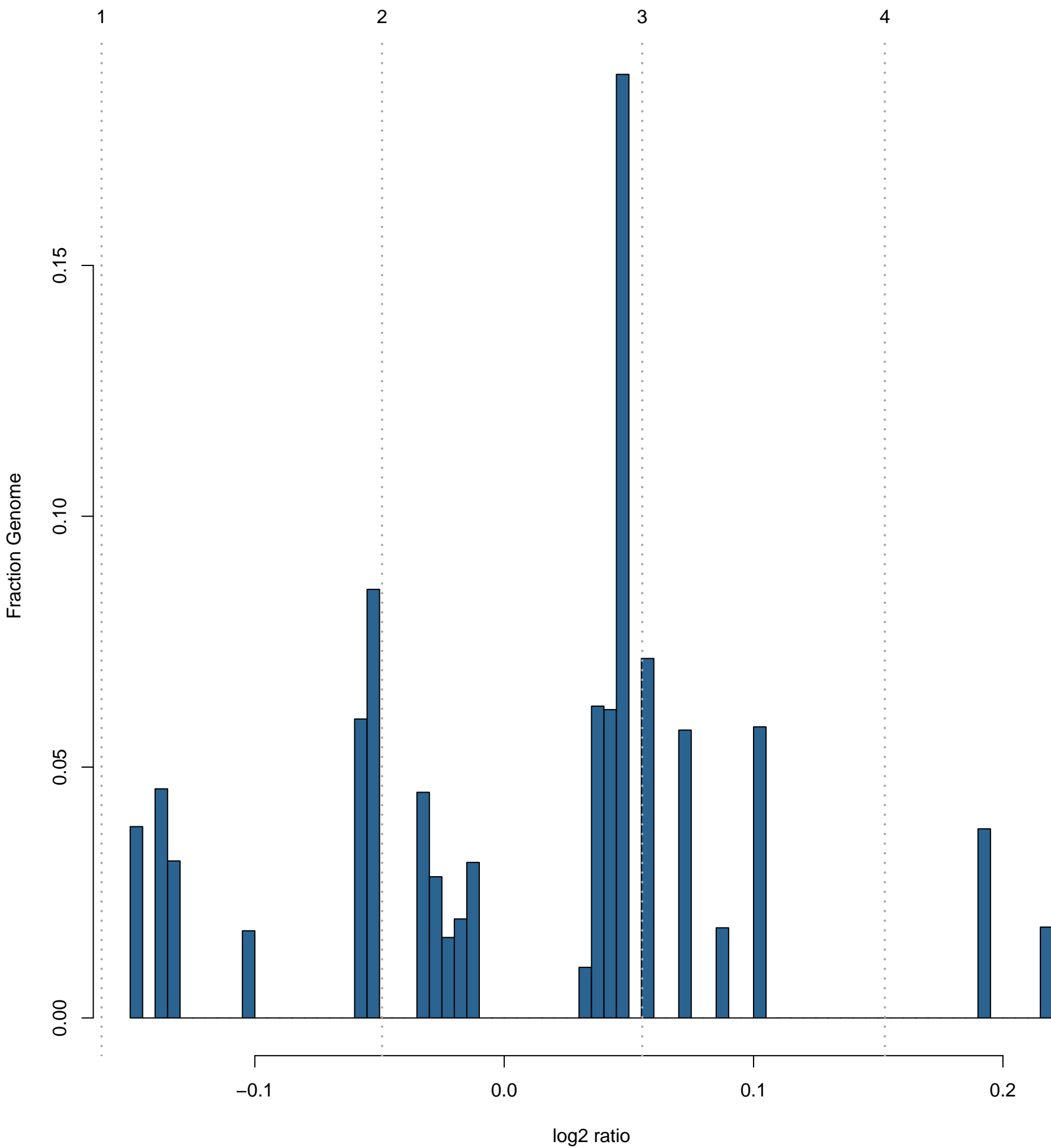


SCNA-fit log-likelihood: -5311.88

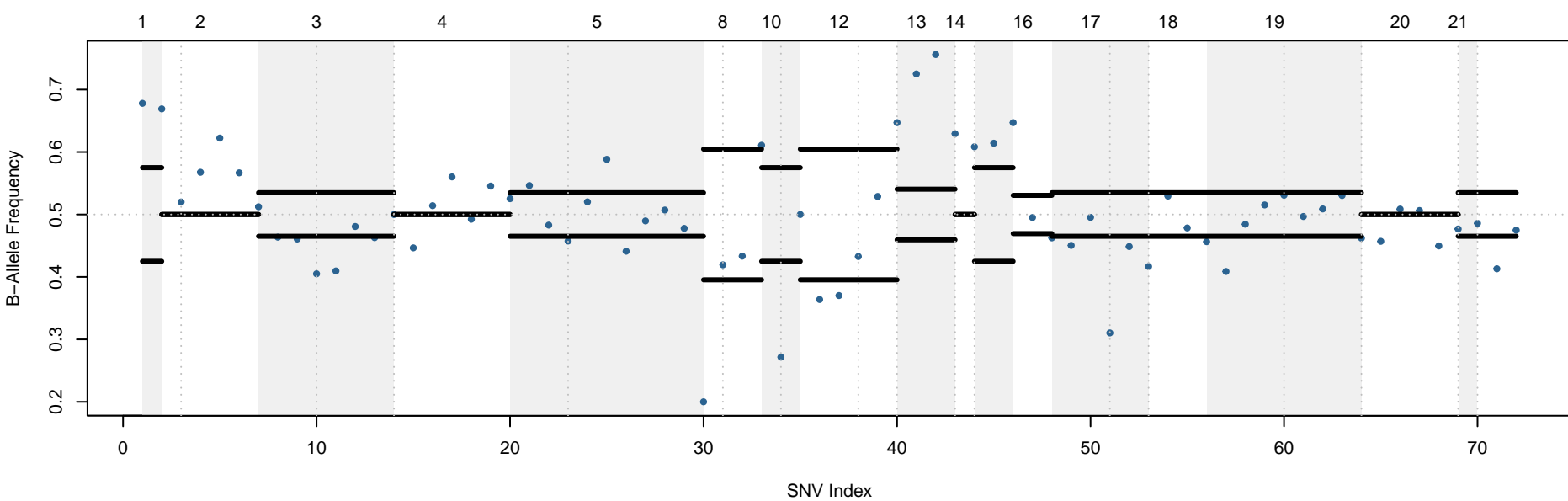




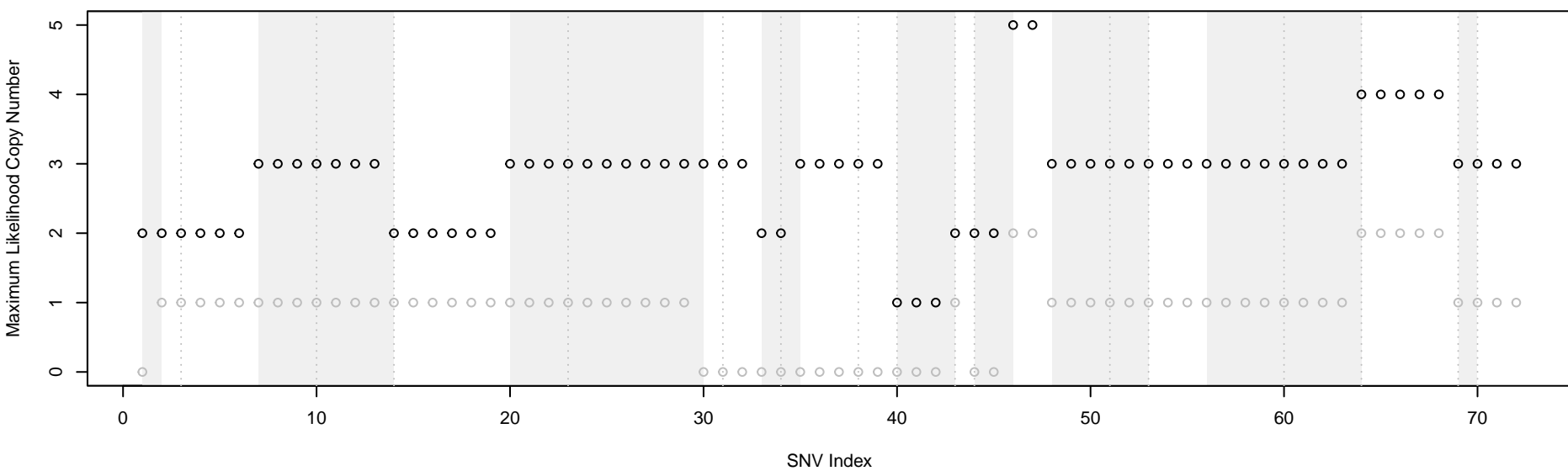
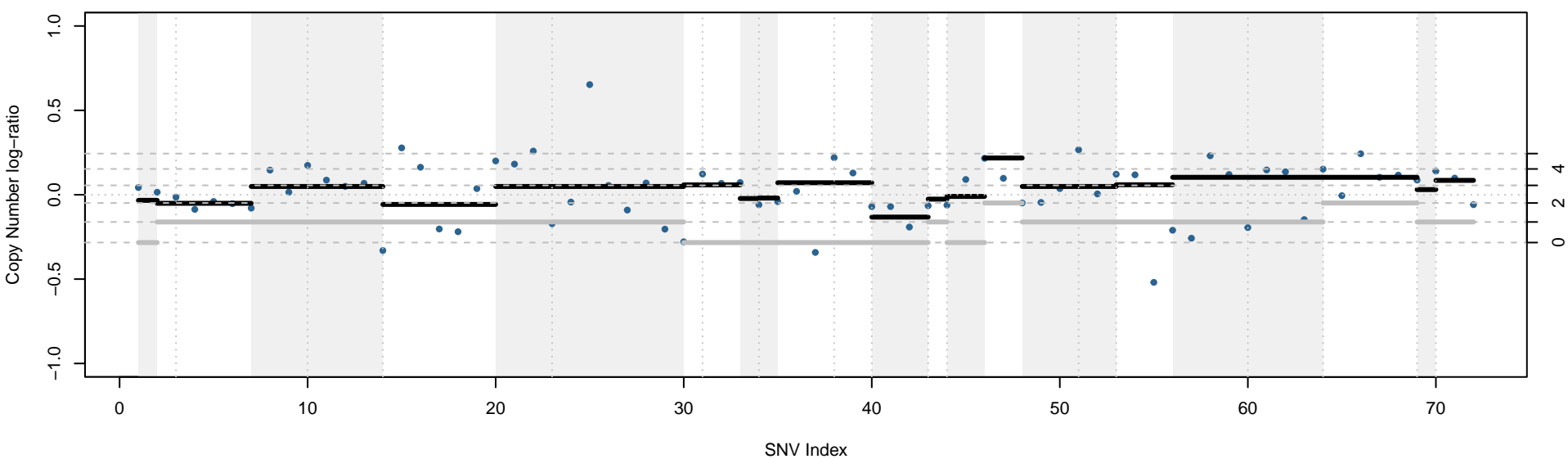
Purity: 0.15 Tumor ploidy: 2.461



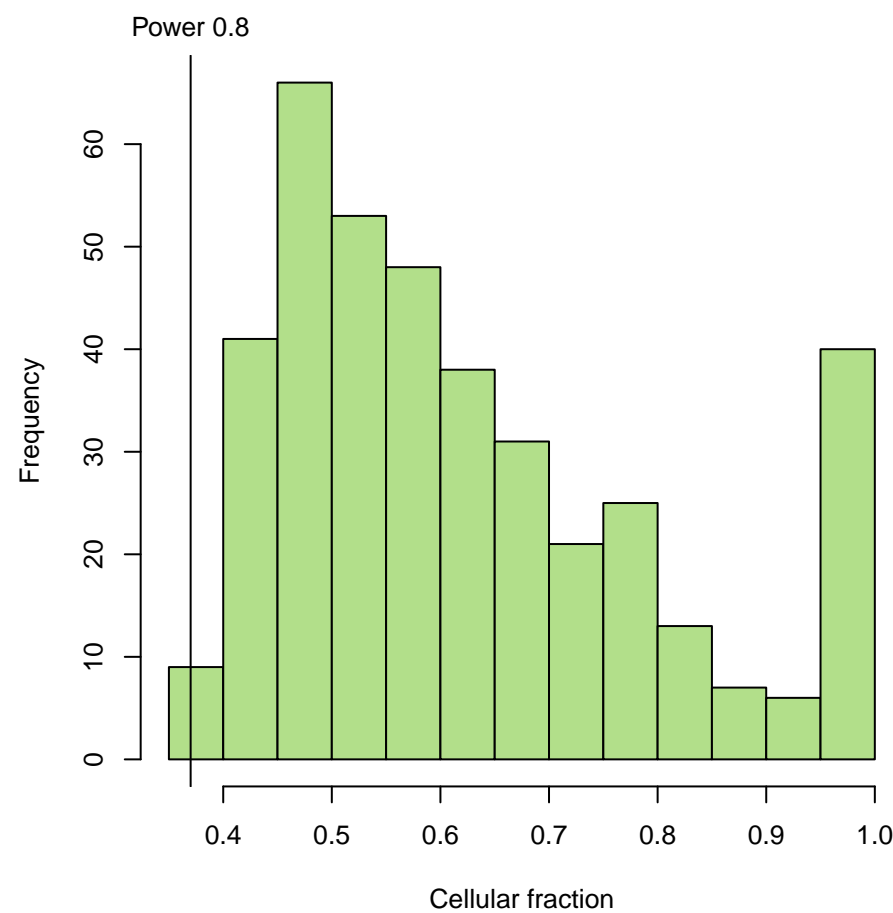
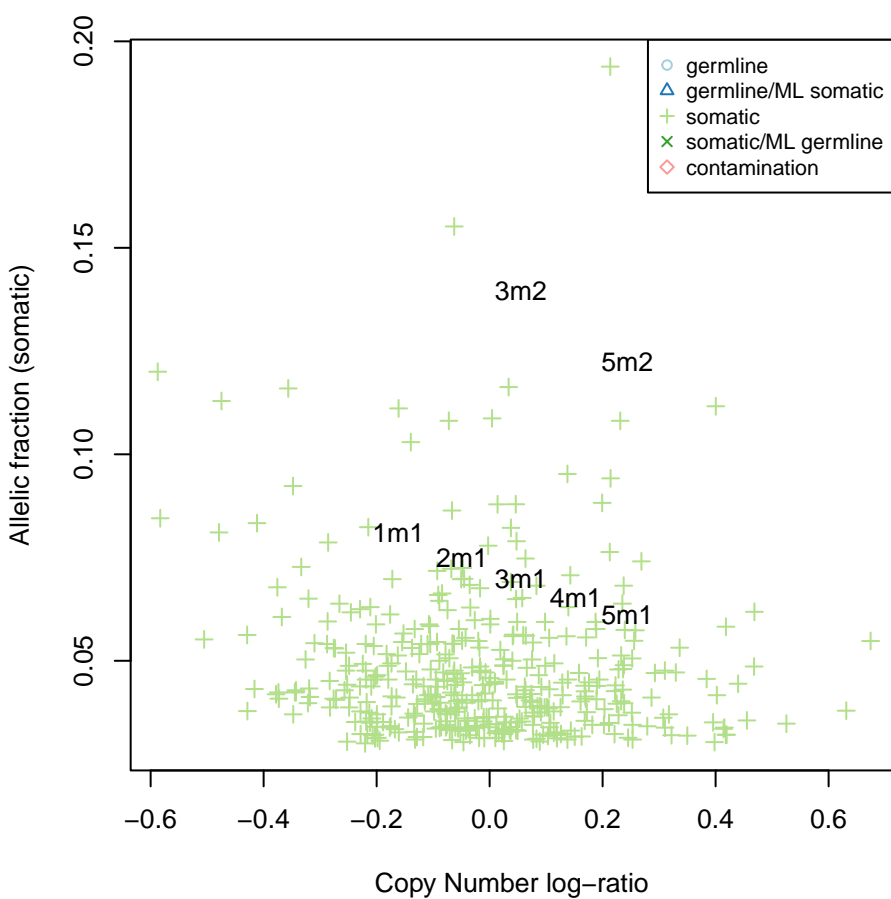
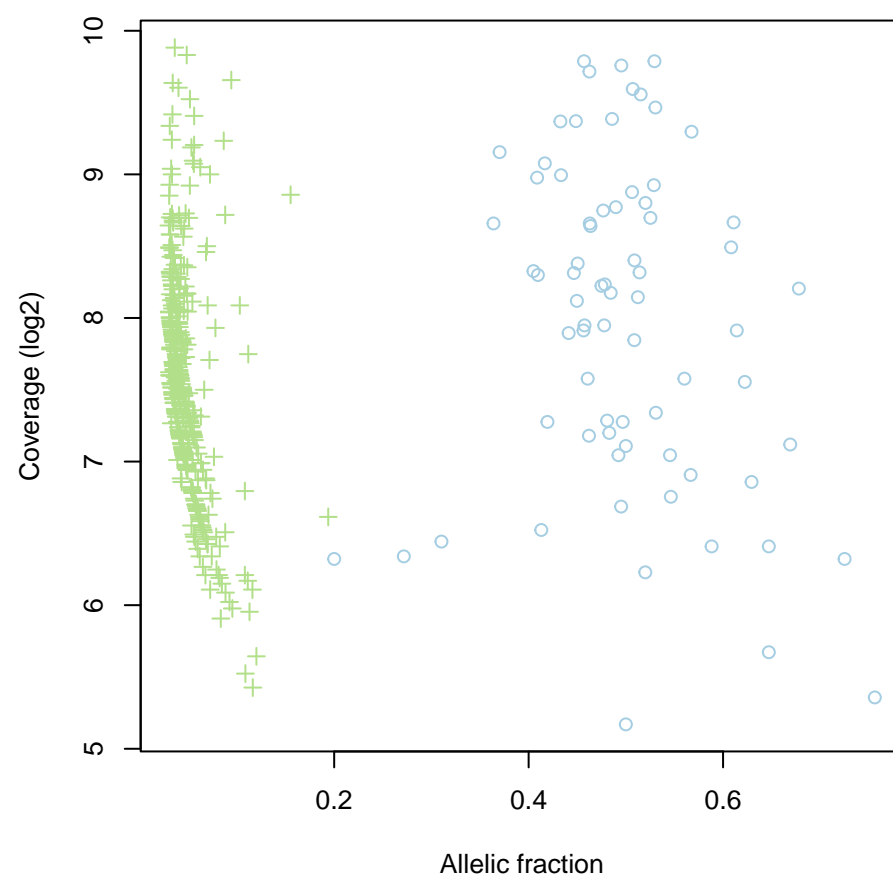
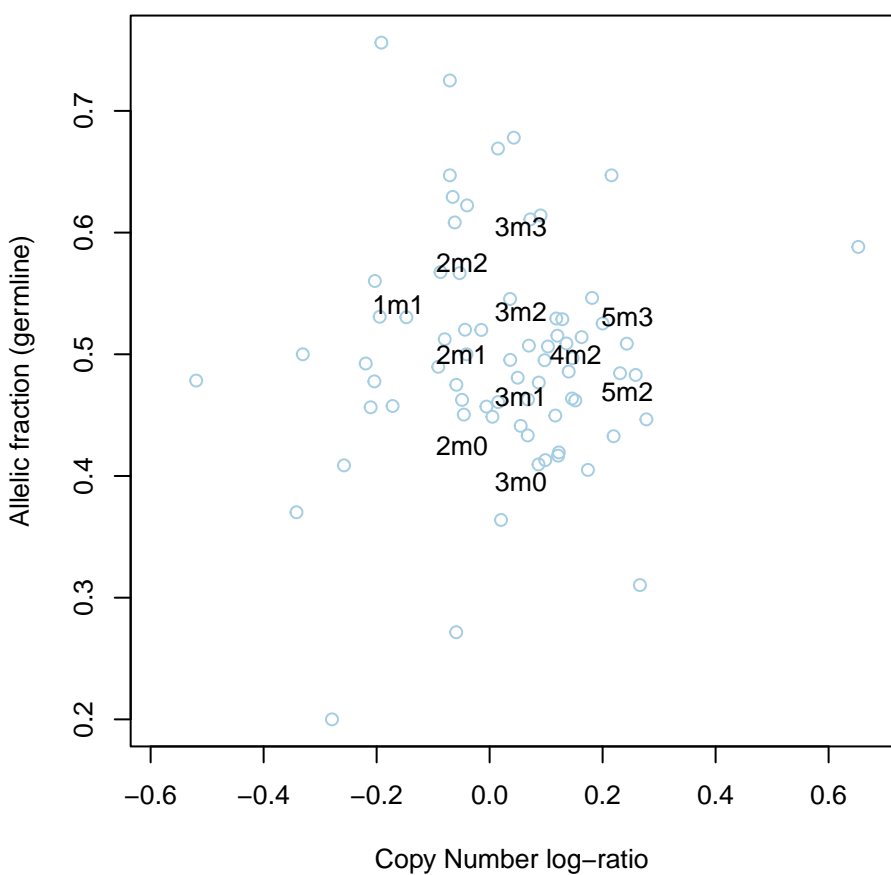
Purity: 0.15 Tumor ploidy: 2.461 SNV log-likelihood: 133.92 GoF: 96.2% Mean coverage: 233;281



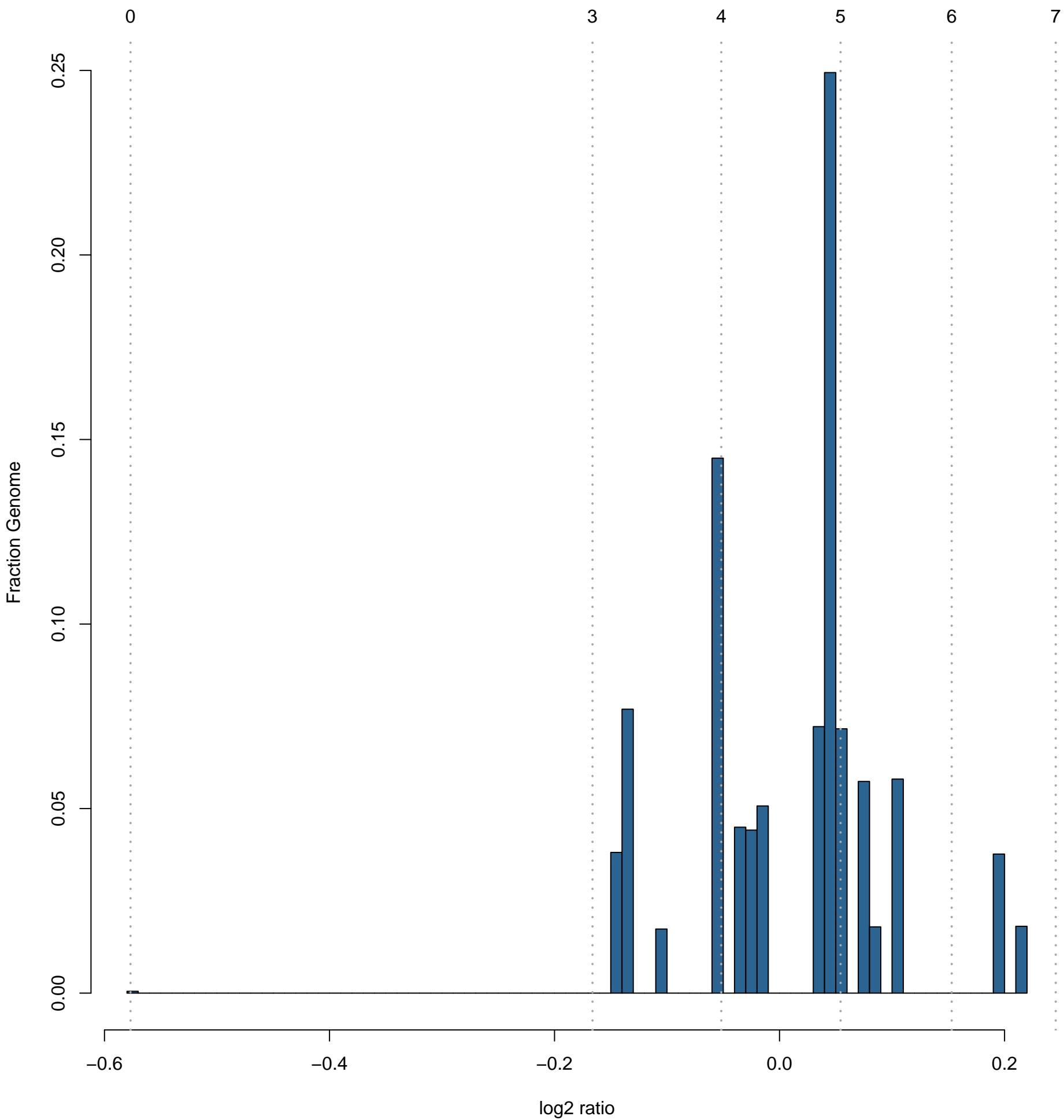
SCNA-fit log-likelihood: -5040.87



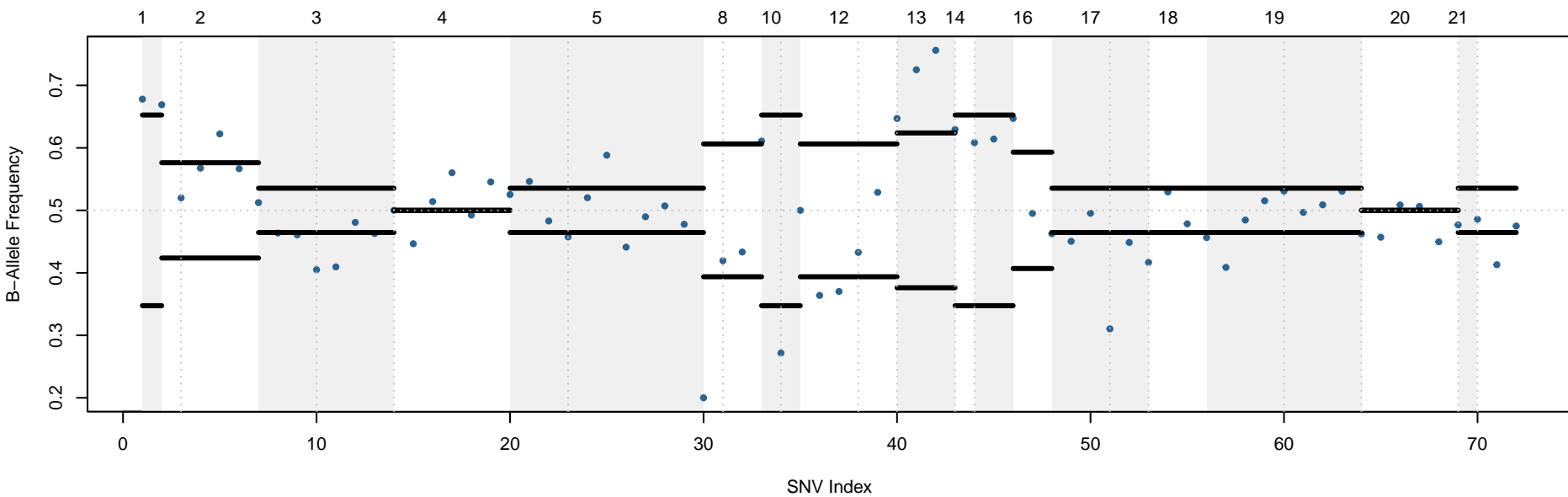




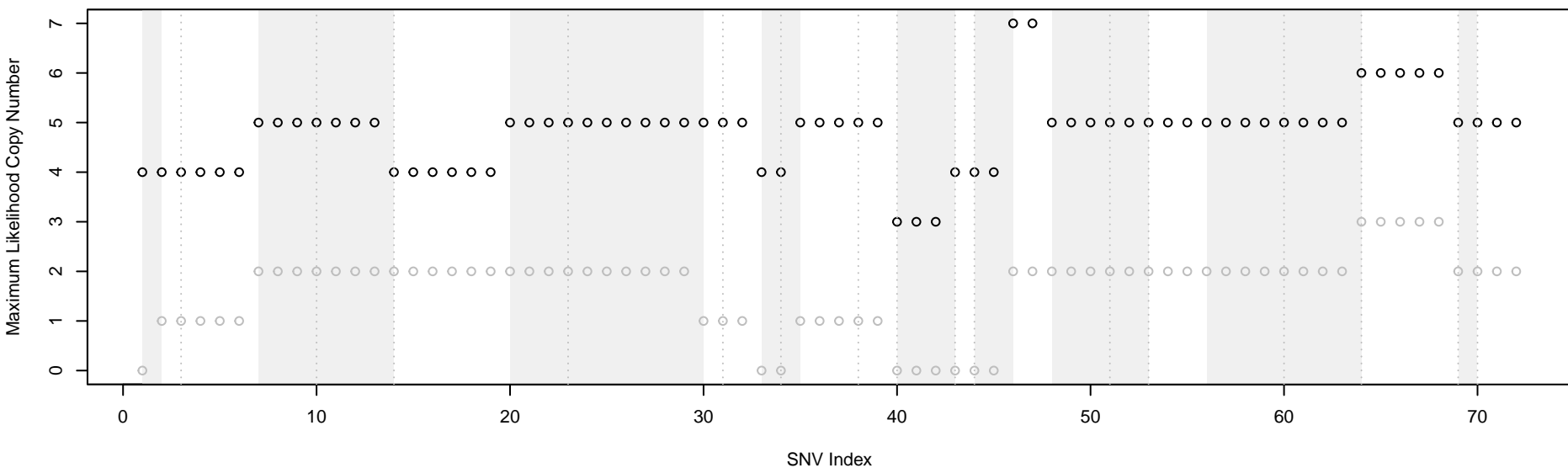
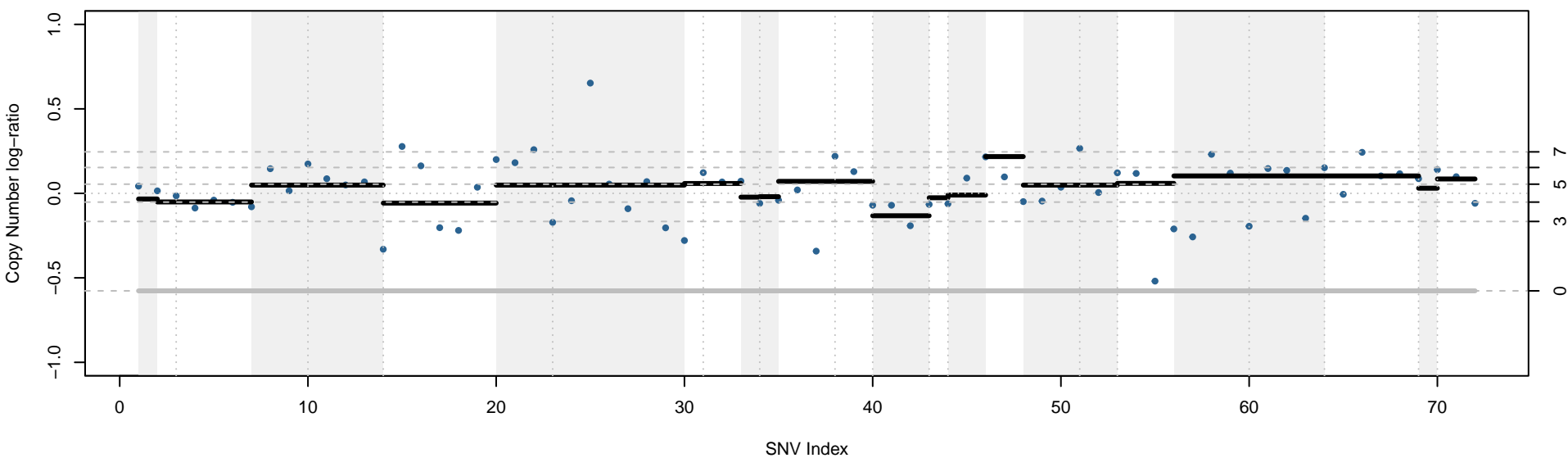
Purity: 0.18 Tumor ploidy: 4.479

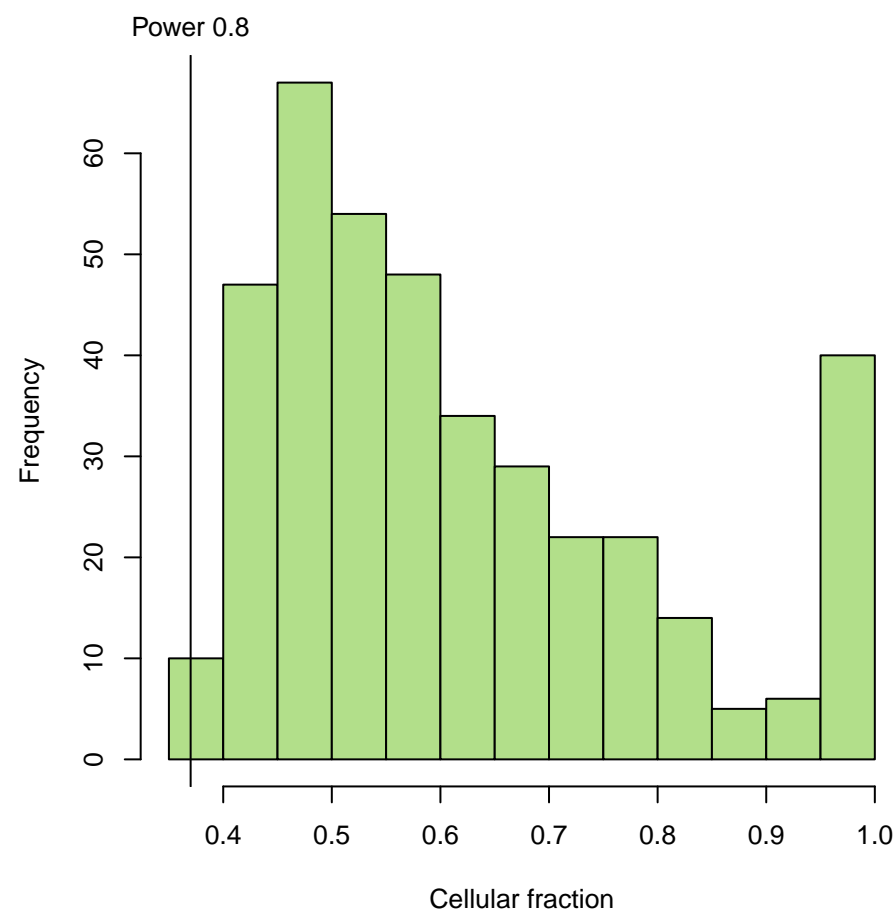
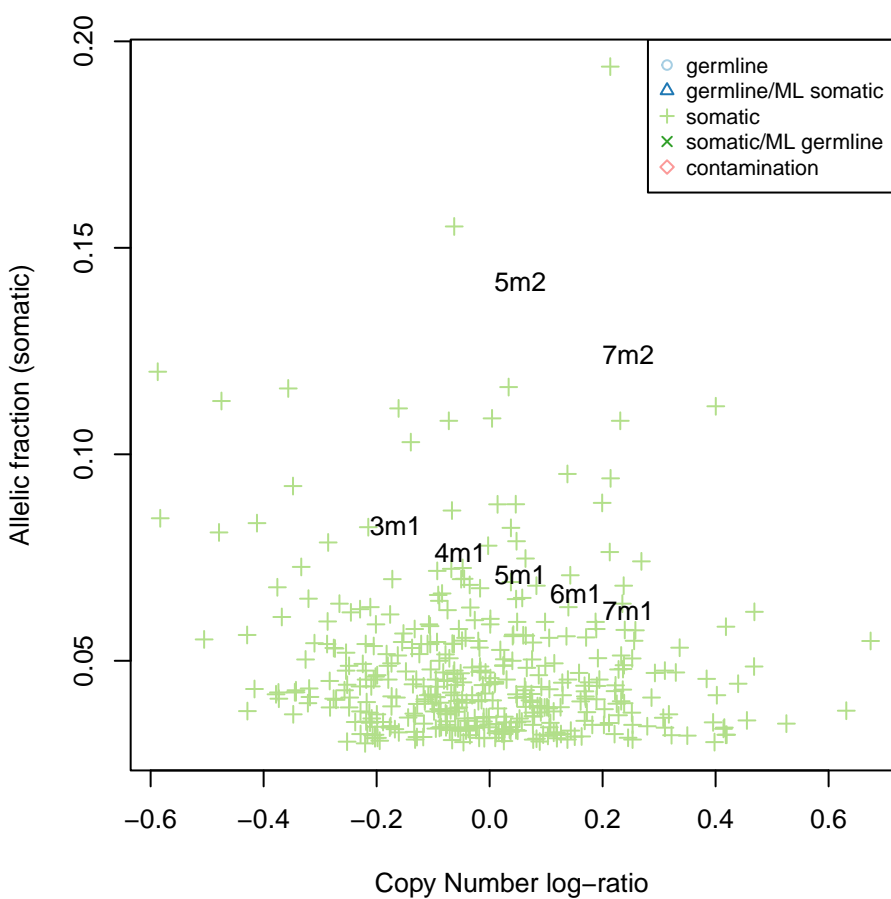
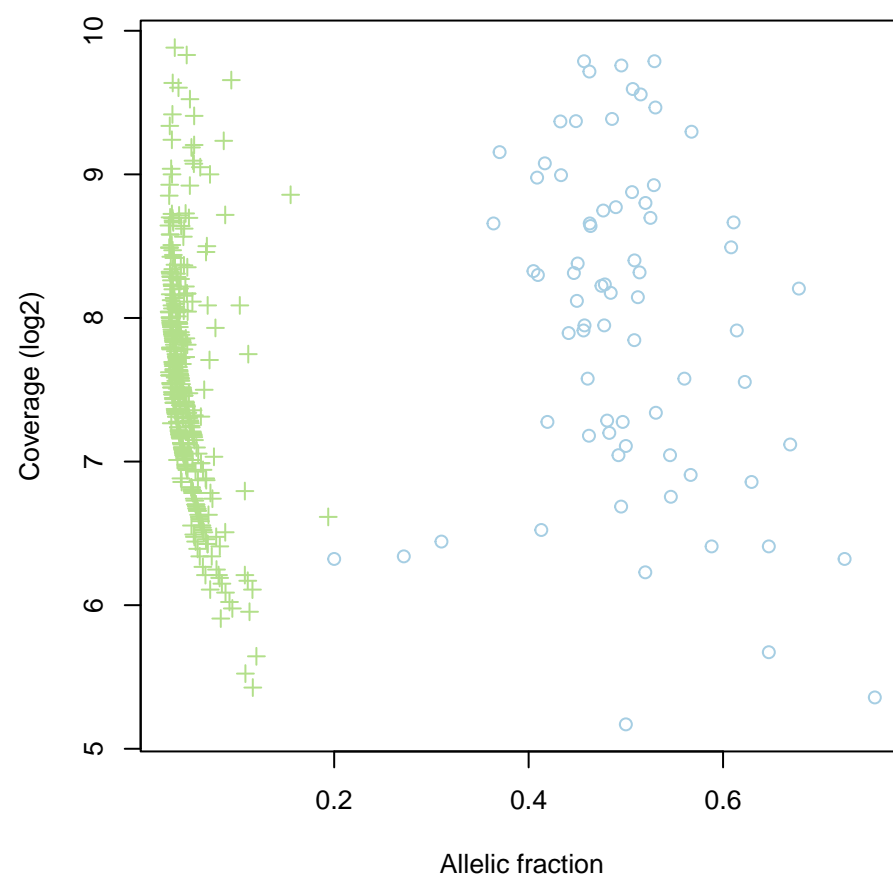
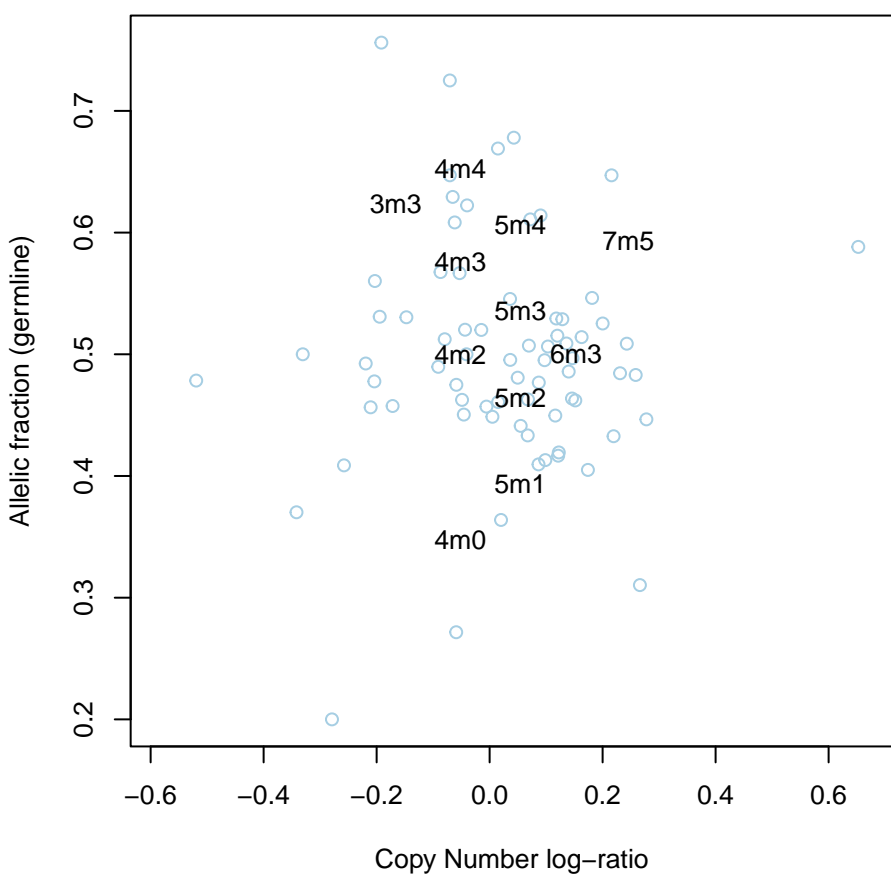


Purity: 0.18 Tumor ploidy: 4.479 SNV log-likelihood: -140.18 GoF: 96.9% Mean coverage: 233;281

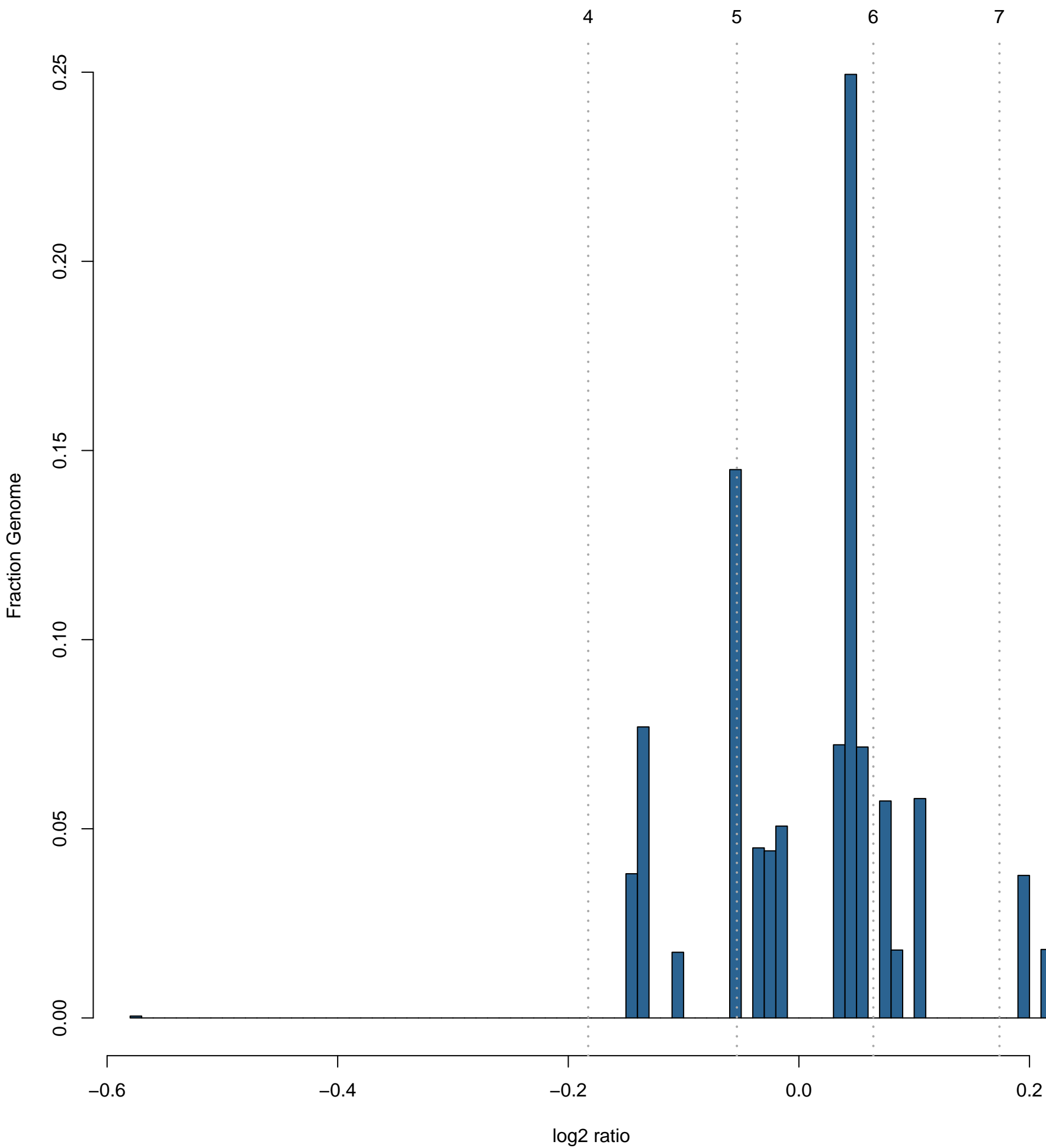


SCNA-fit log-likelihood: -5038.49

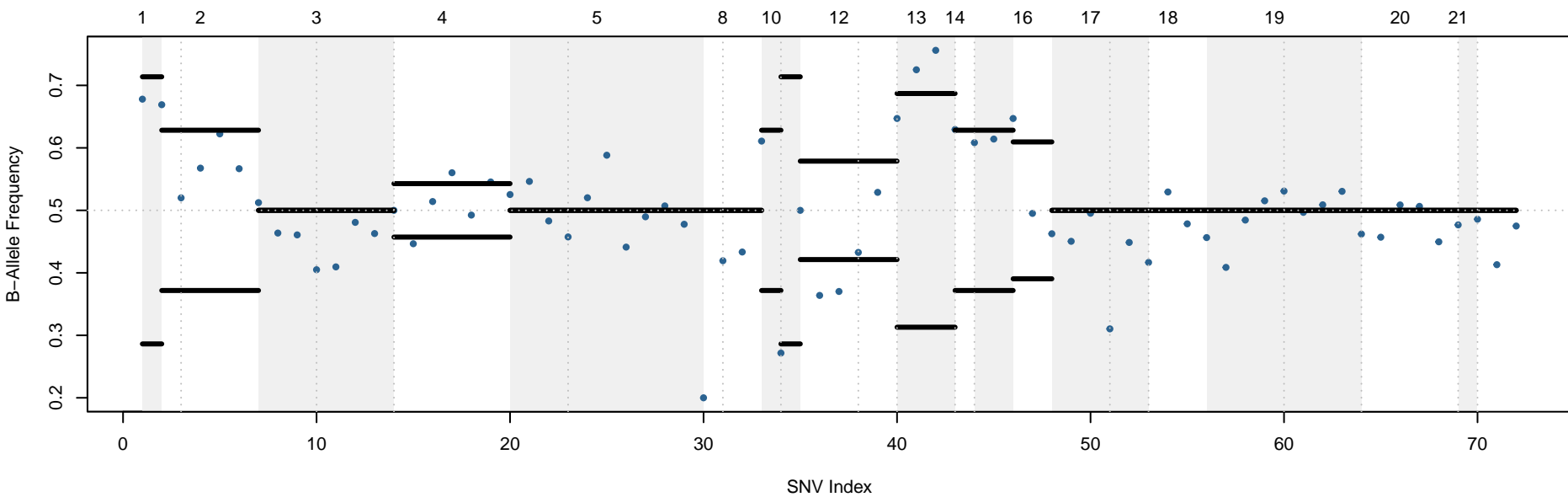




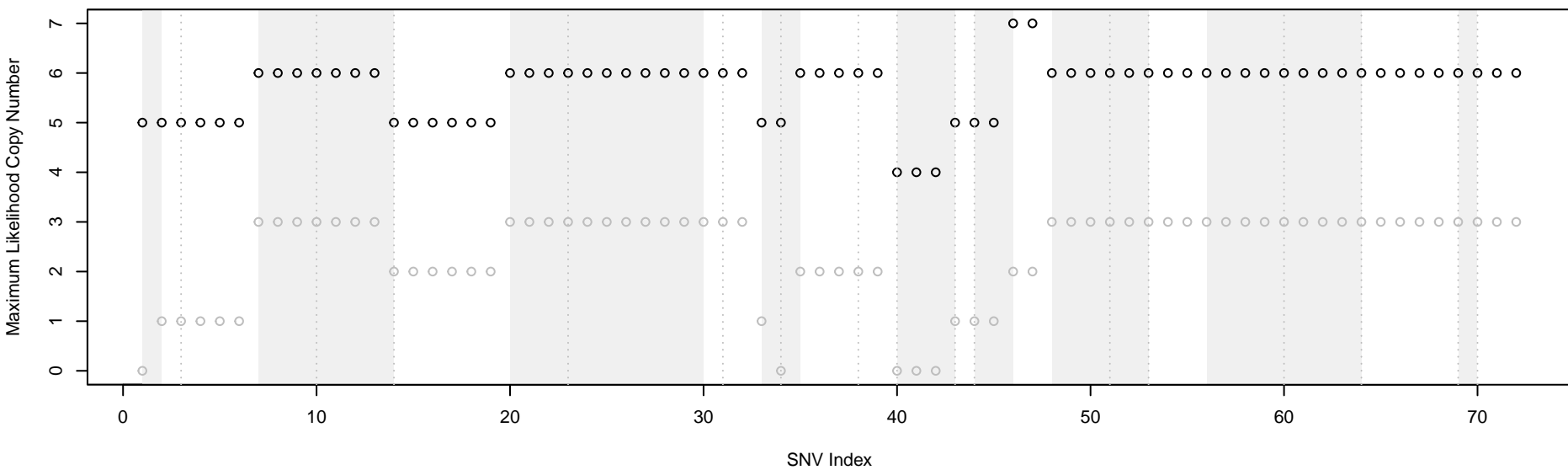
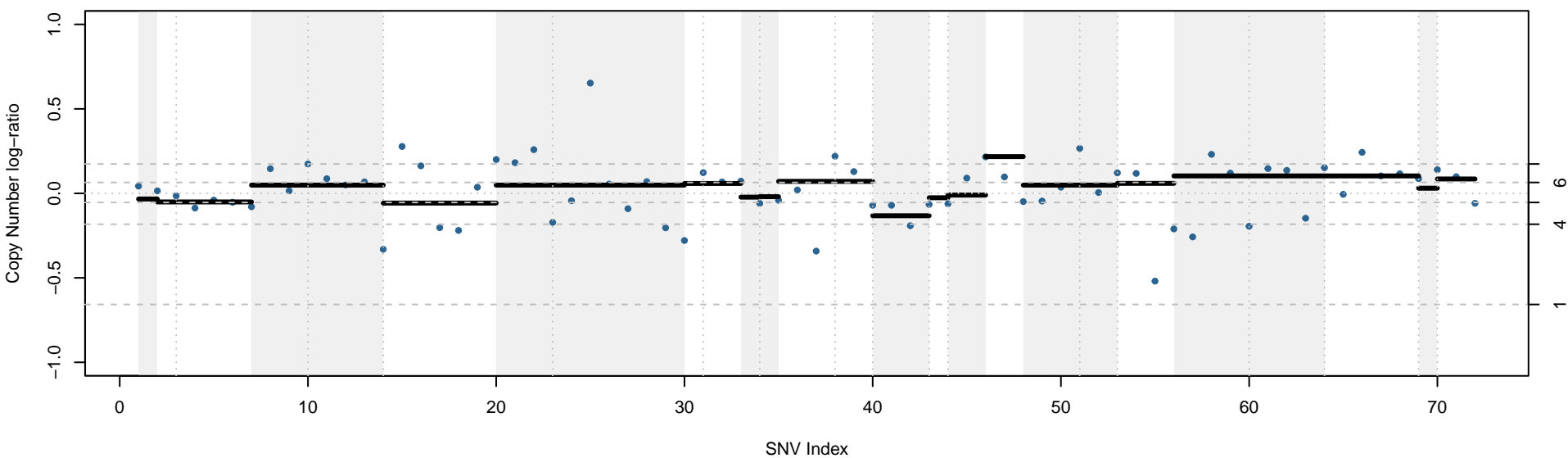
Purity: 0.23 Tumor ploidy: 5.444

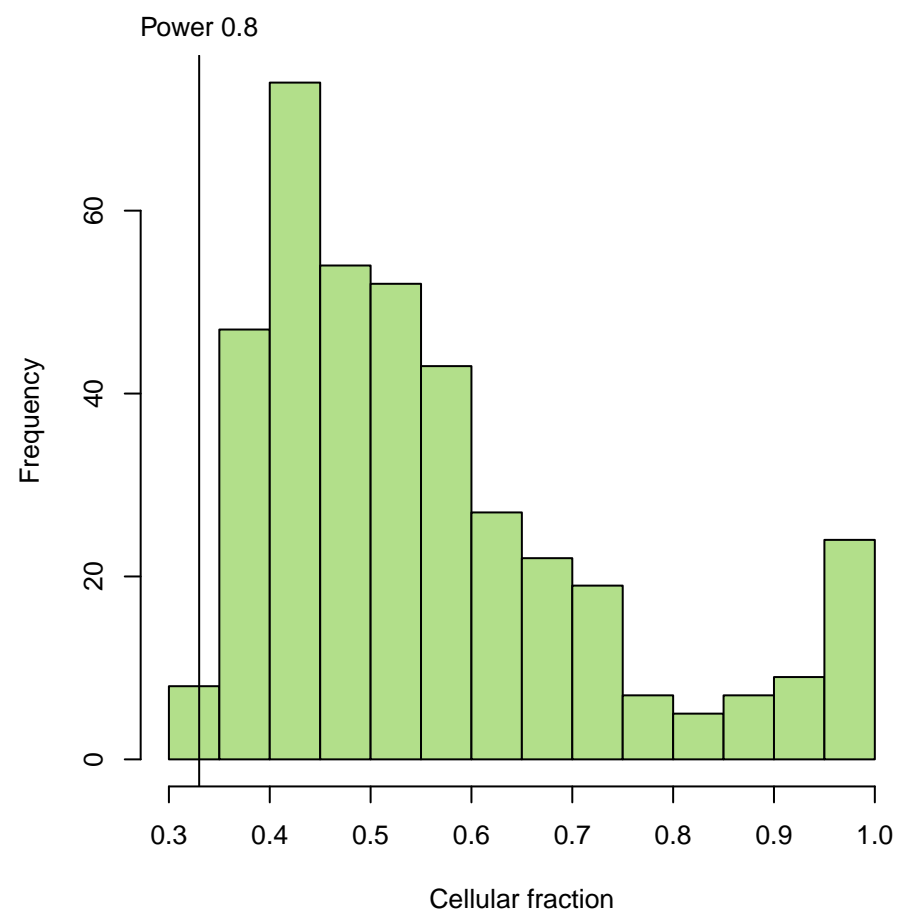
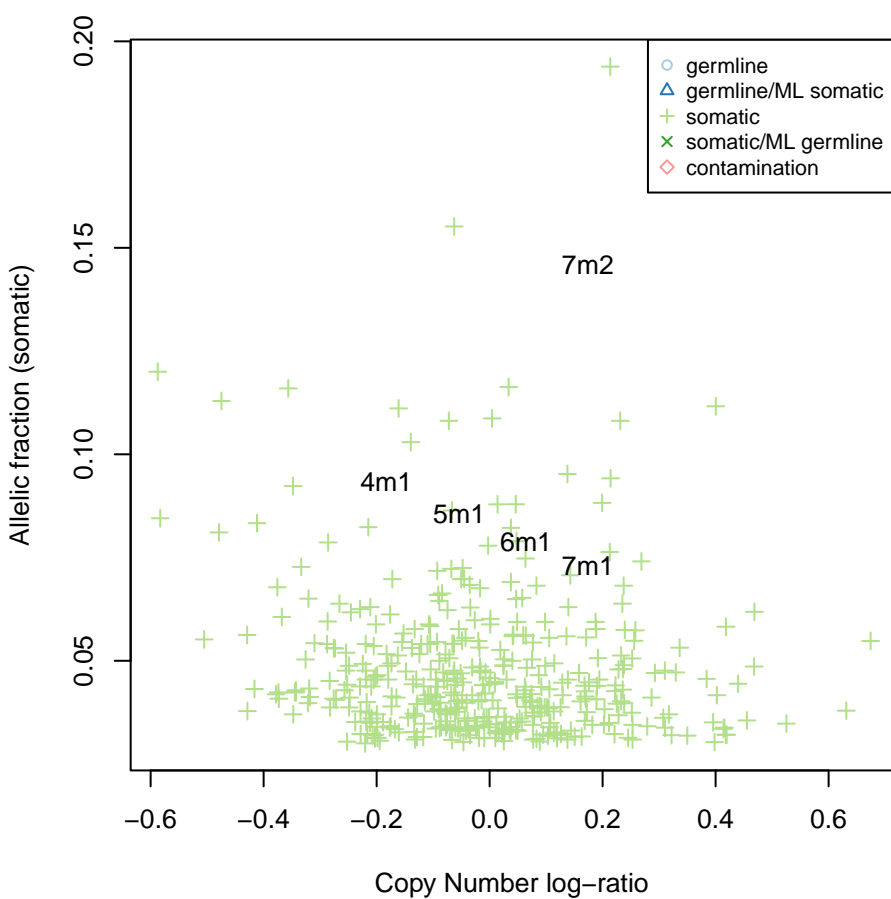
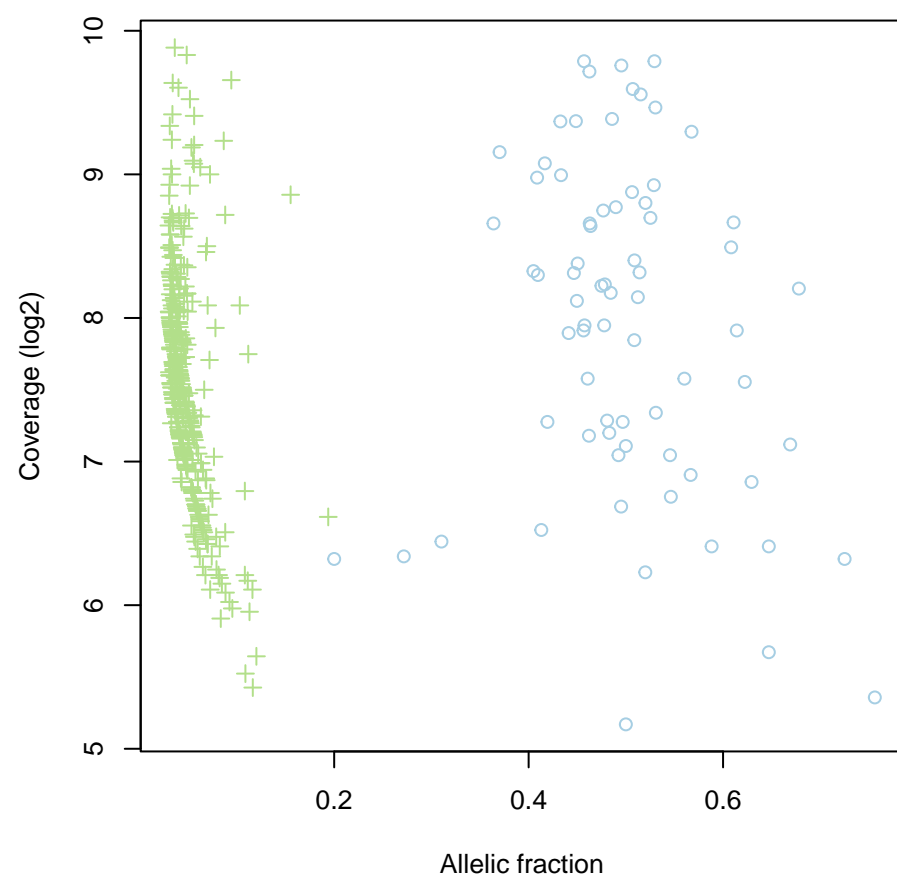
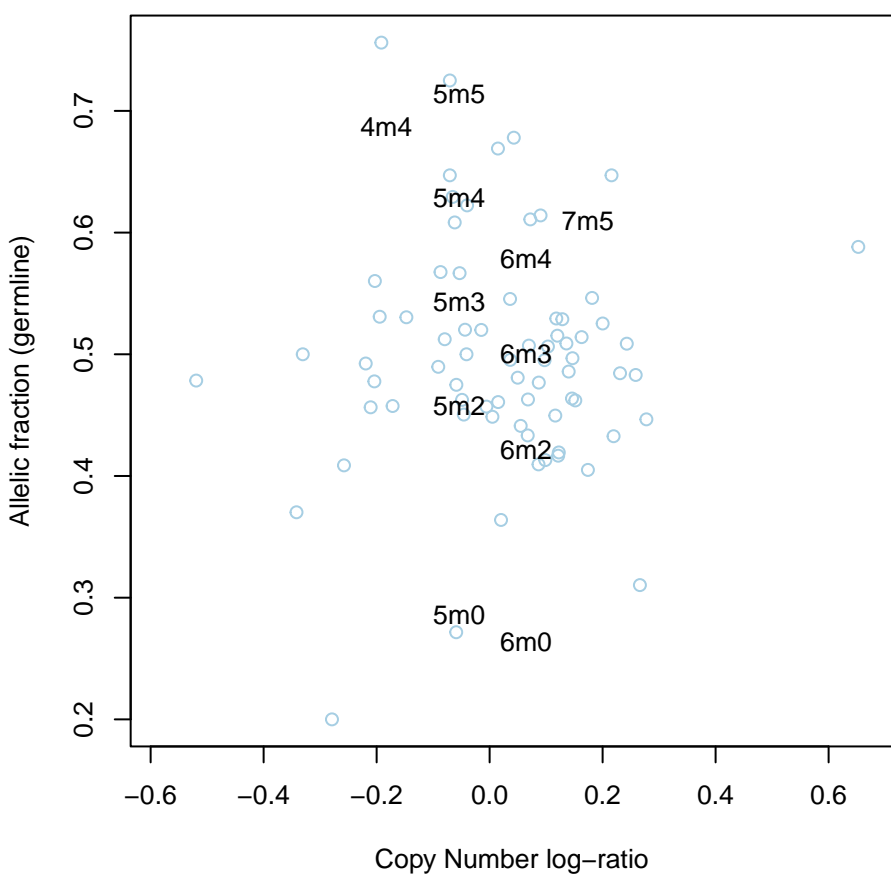


Purity: 0.23 Tumor ploidy: 5.444 SNV log-likelihood: -174.24 GoF: 95.7% Mean coverage: 233;281

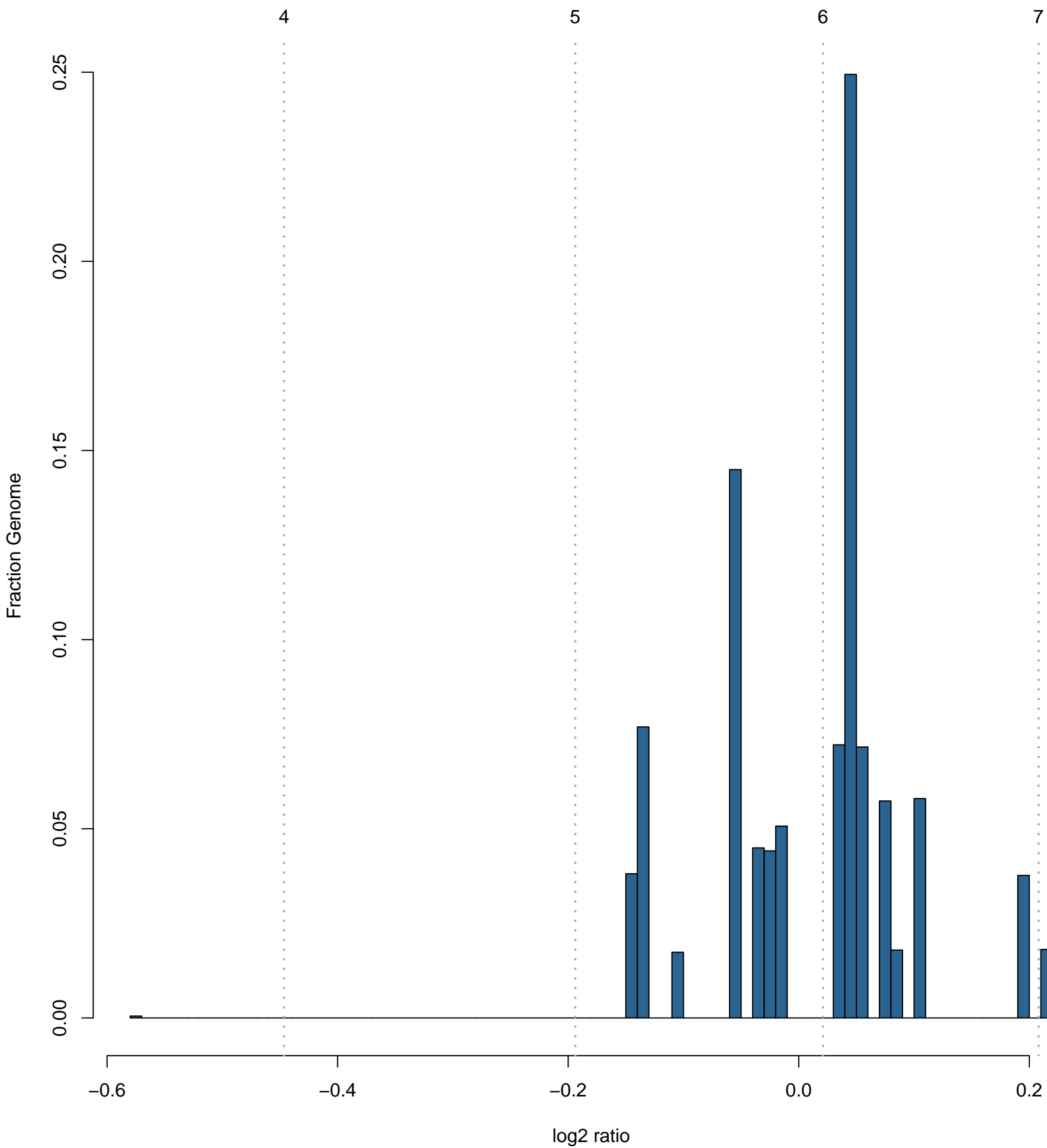


SCNA-fit log-likelihood: -5057.42



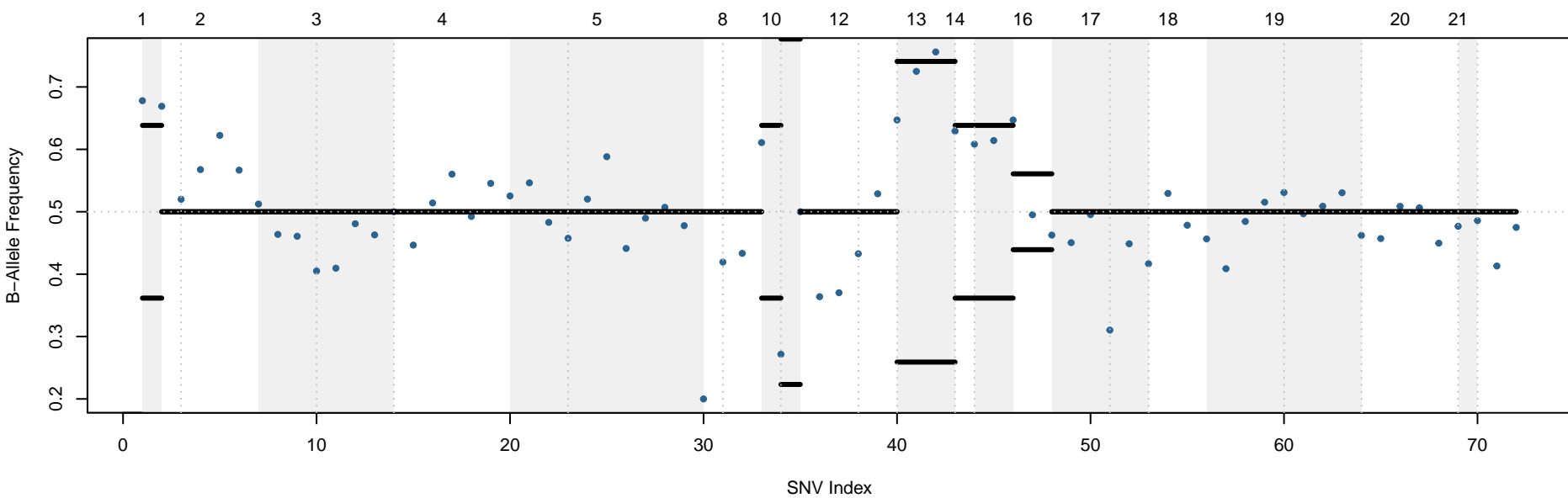


Purity: 0.62 Tumor ploidy: 5.895

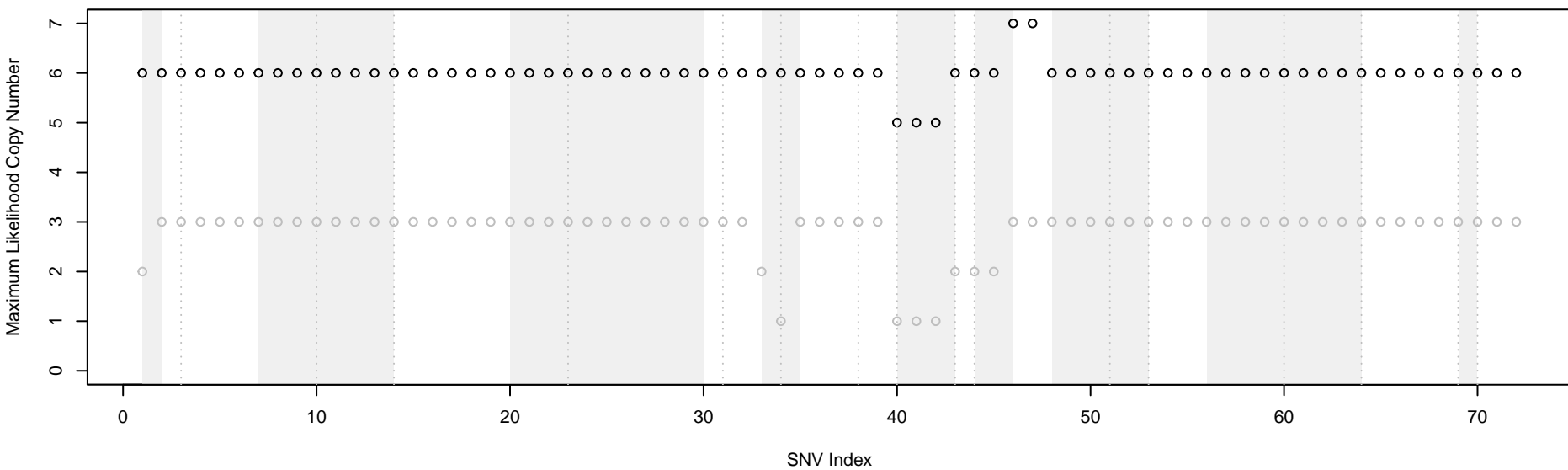
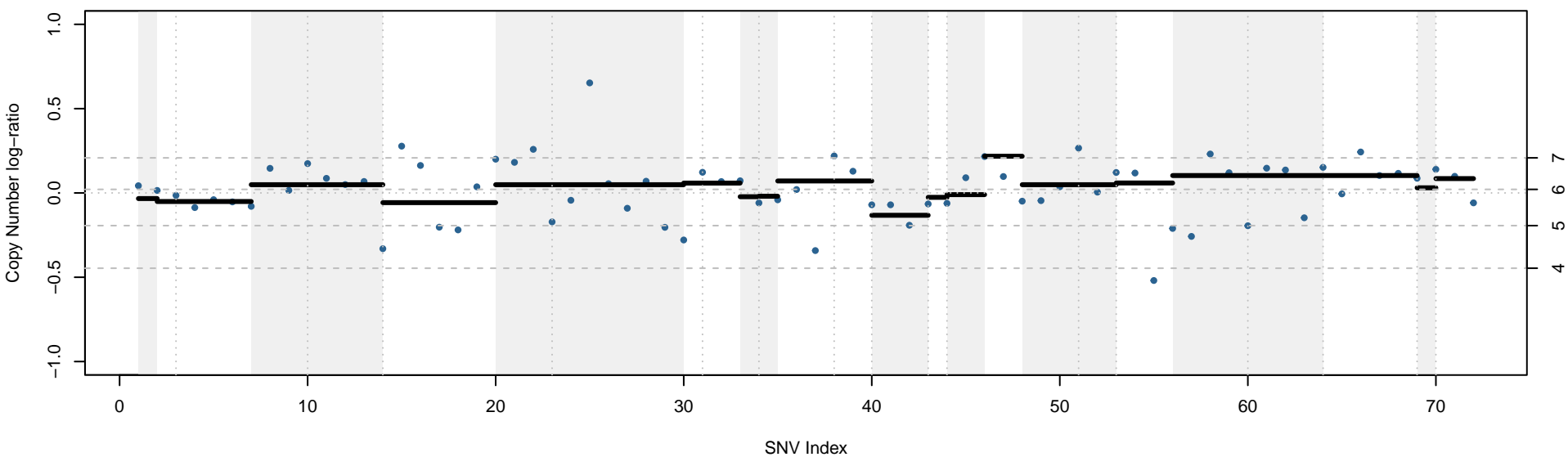


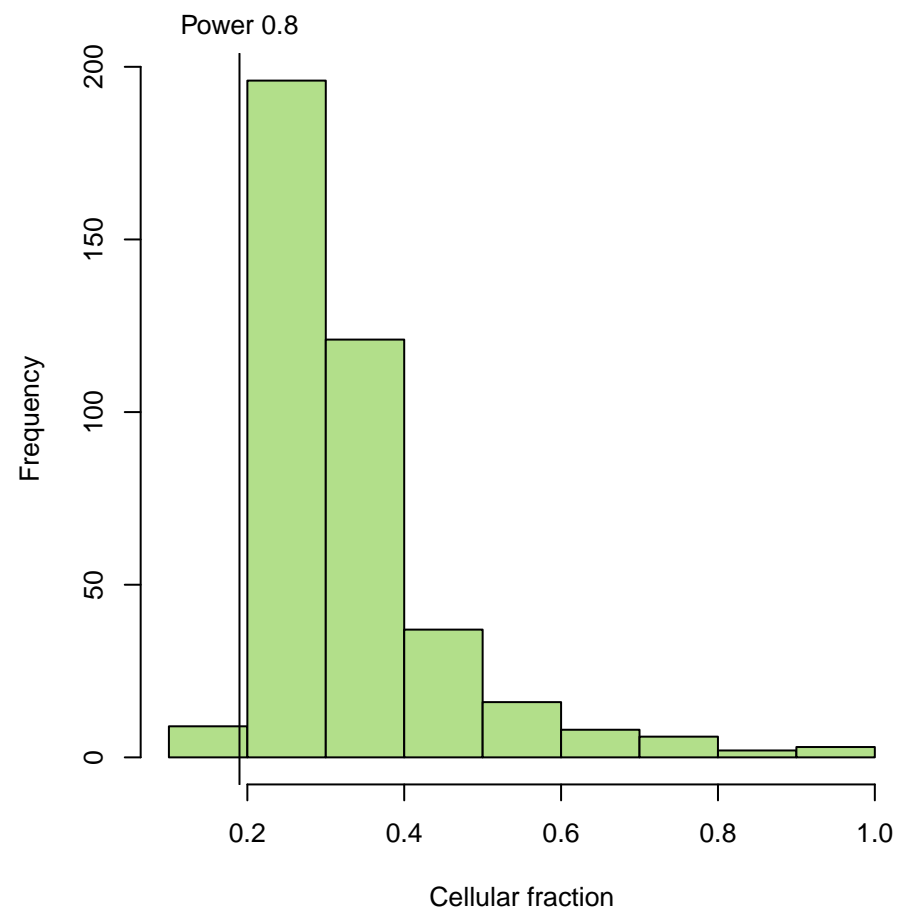
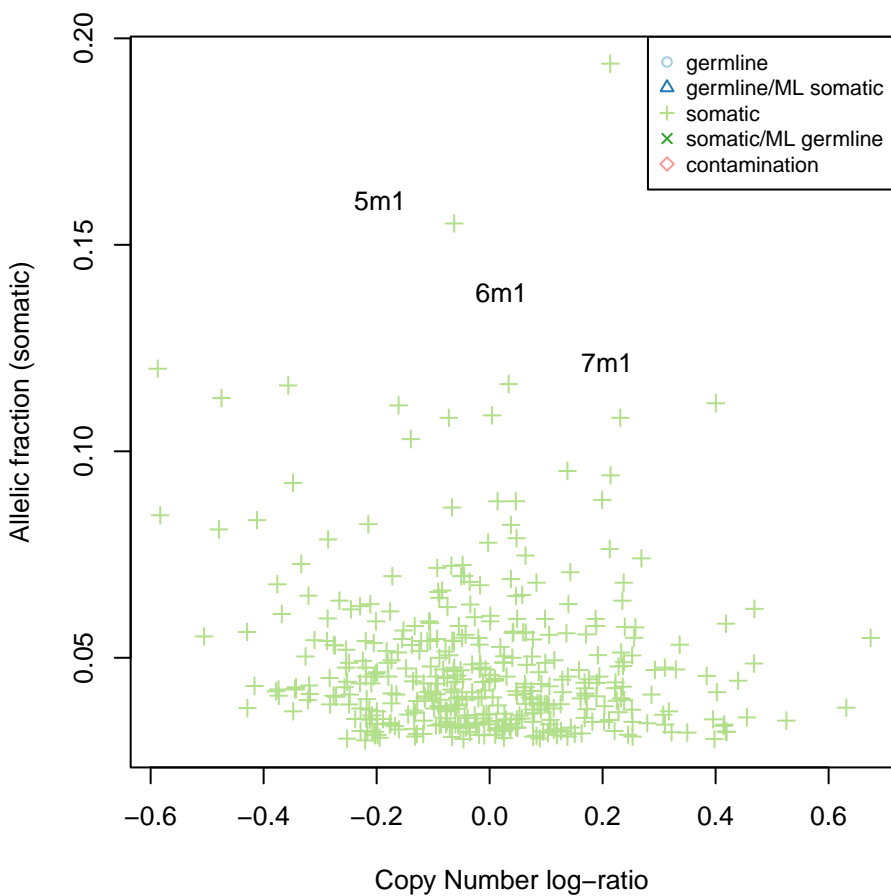
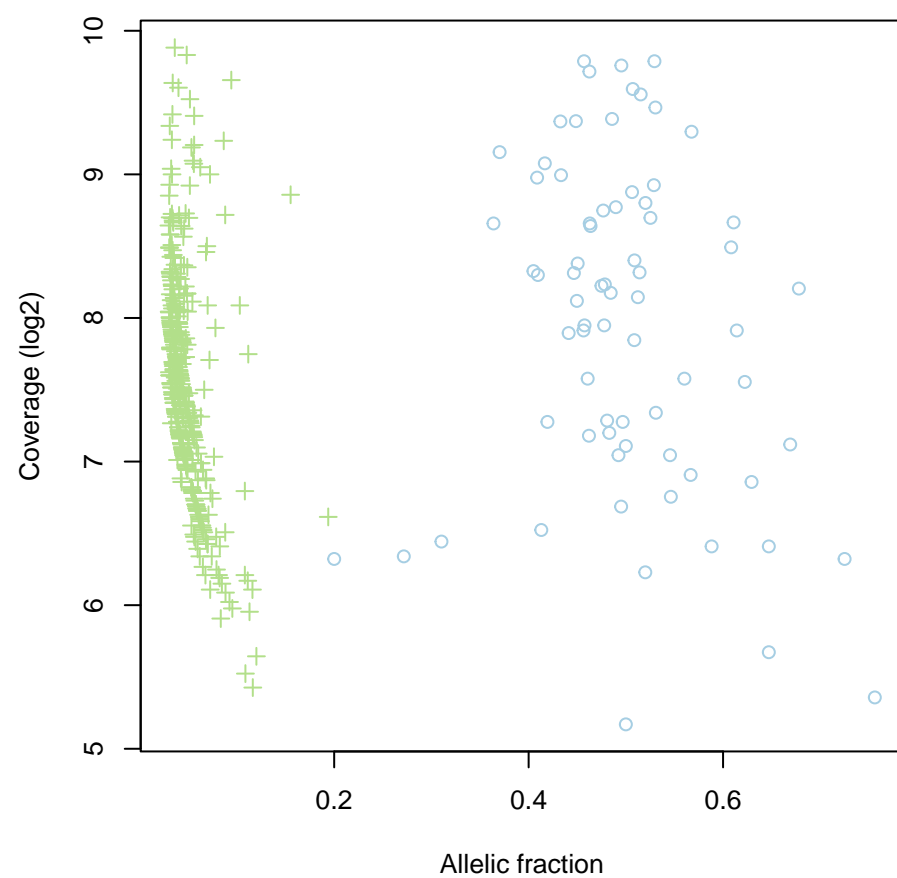
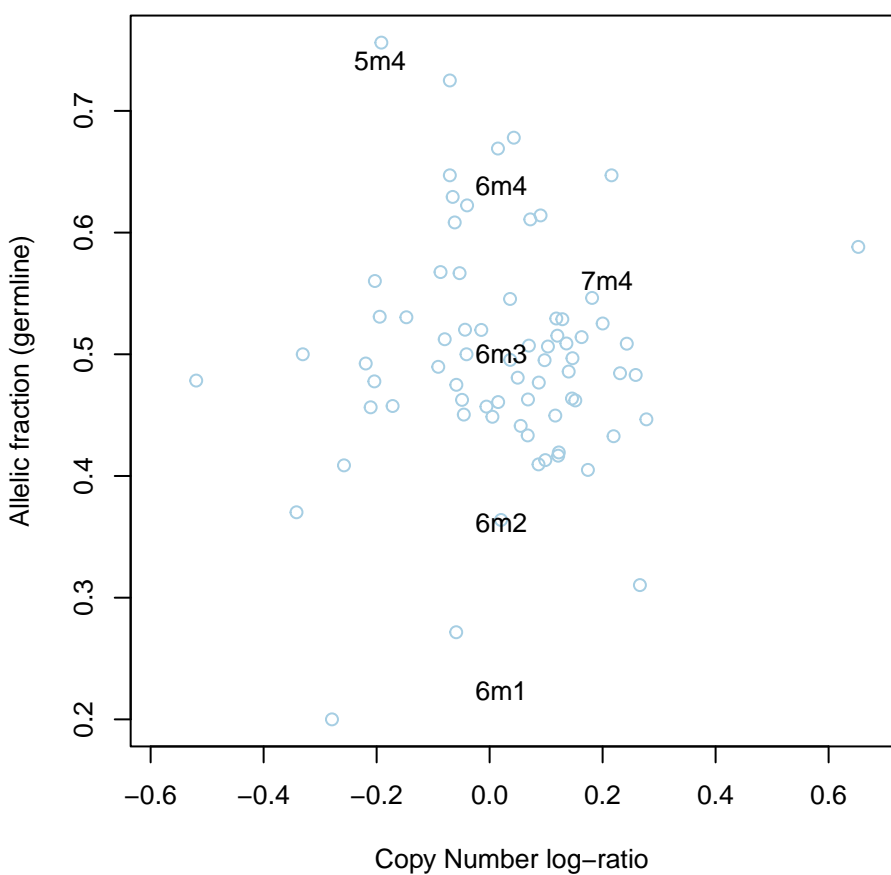


Purity: 0.62 Tumor ploidy: 5.895 SNV log-likelihood: -145.44 GoF: 79.3% Mean coverage: 233;281

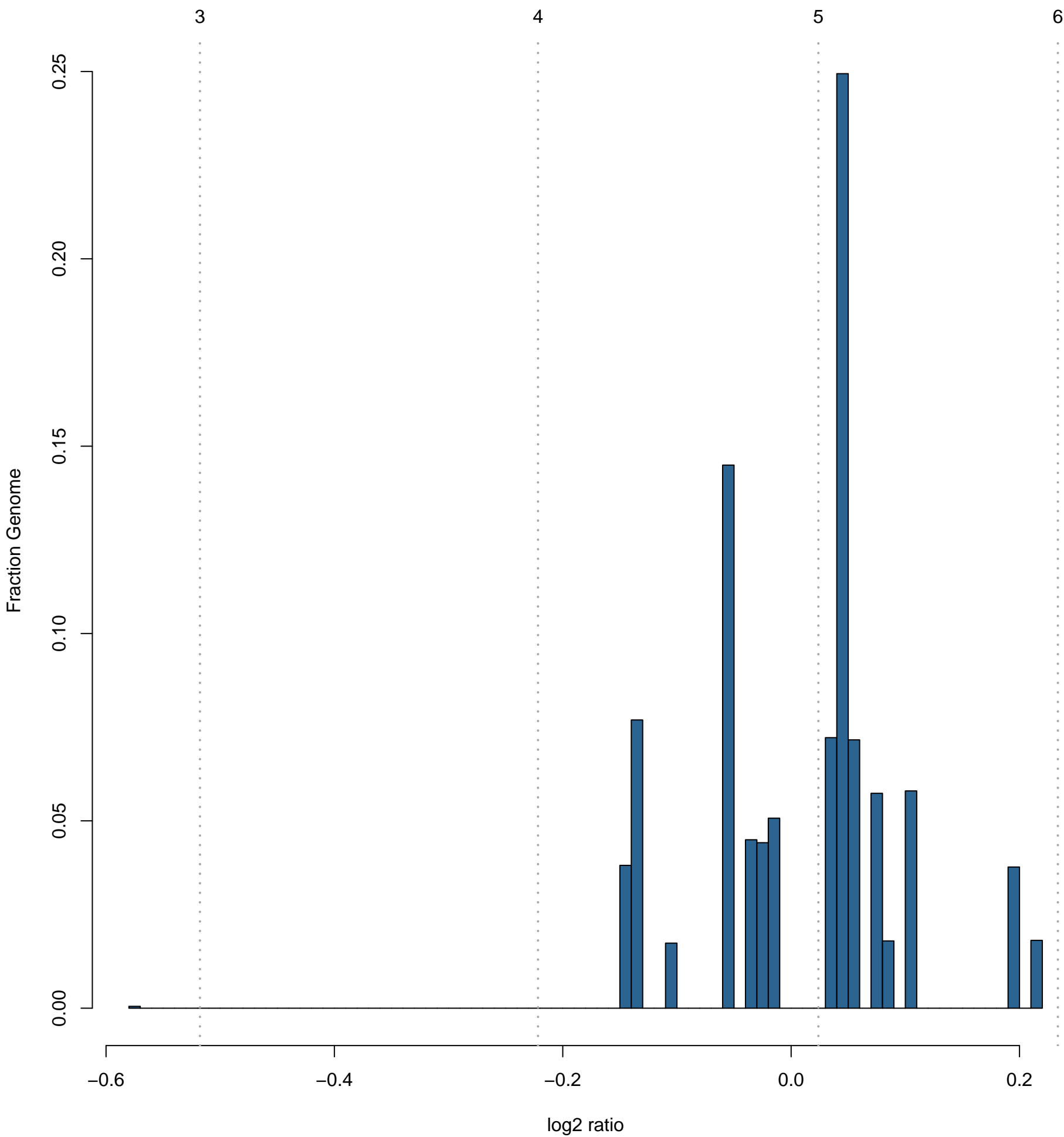


SCNA-fit log-likelihood: -5238.2

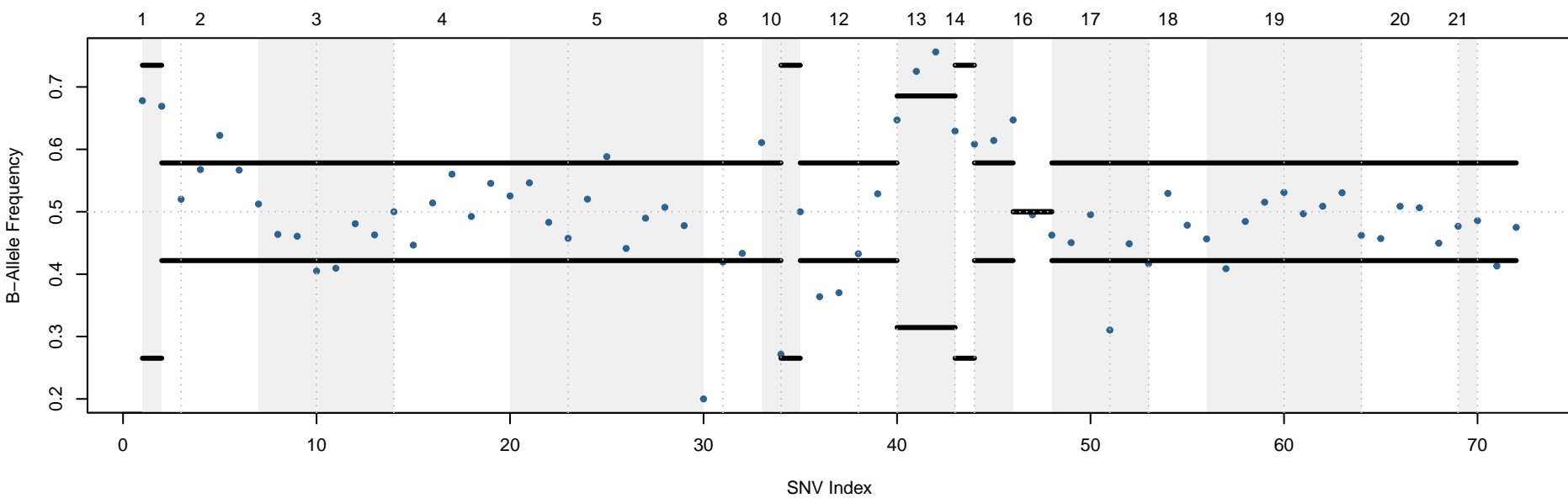




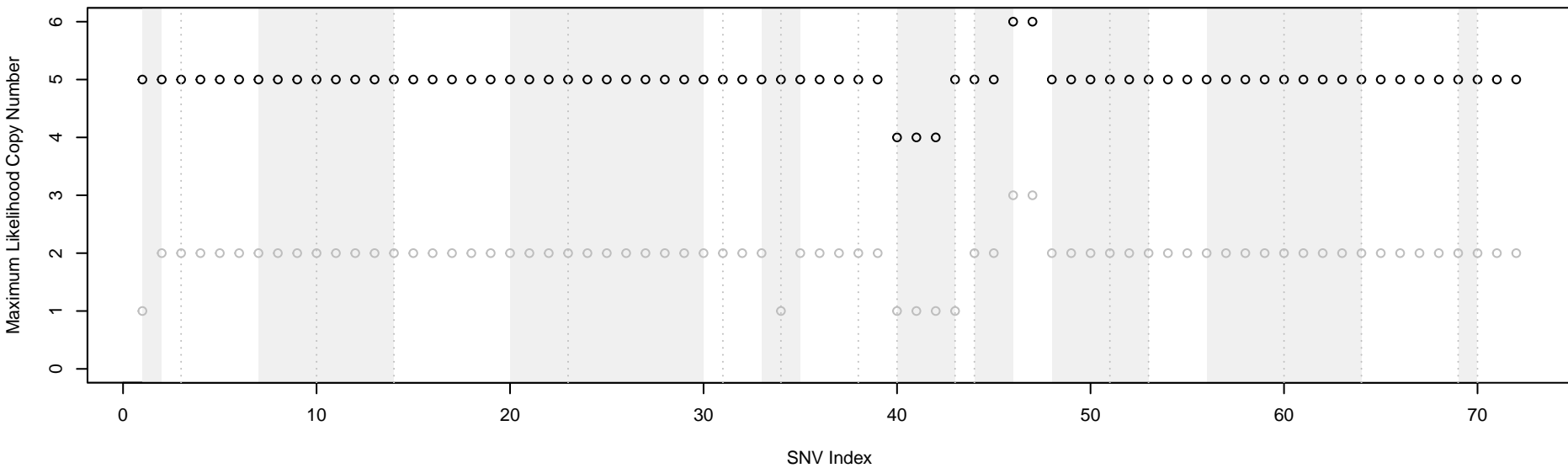
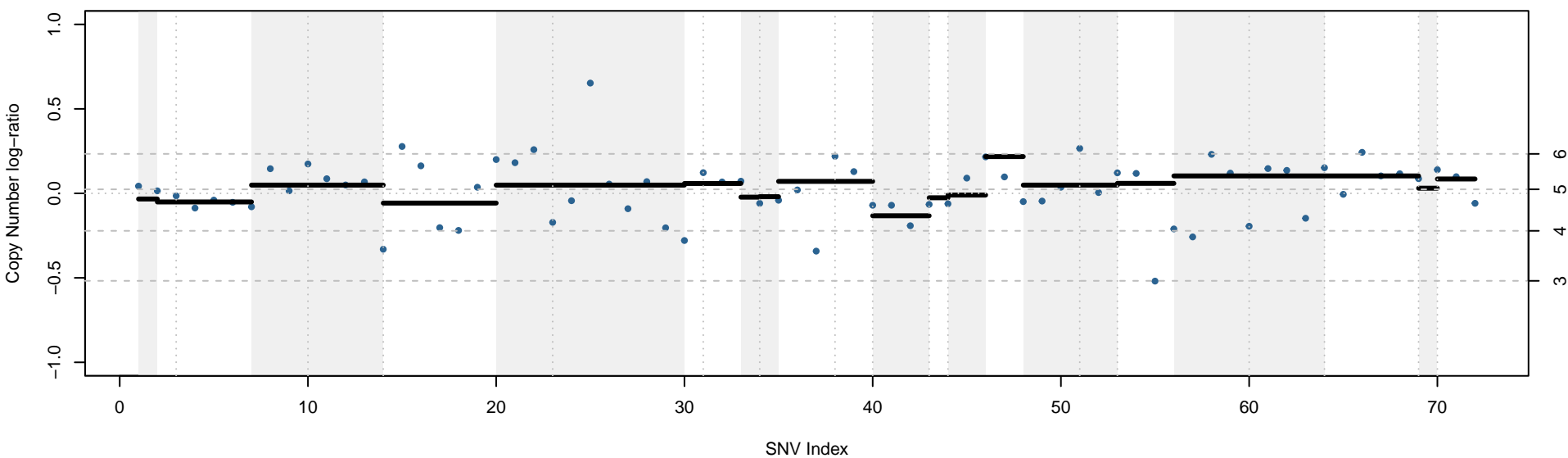
Purity: 0.59 Tumor ploidy: 4.895

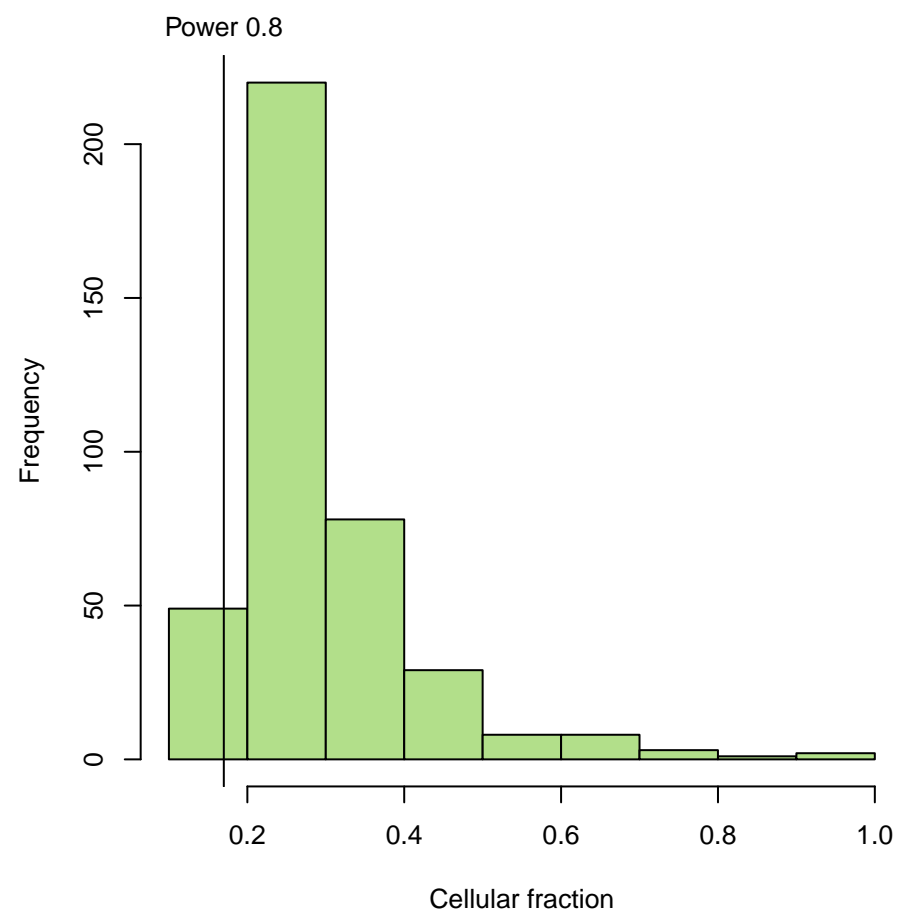
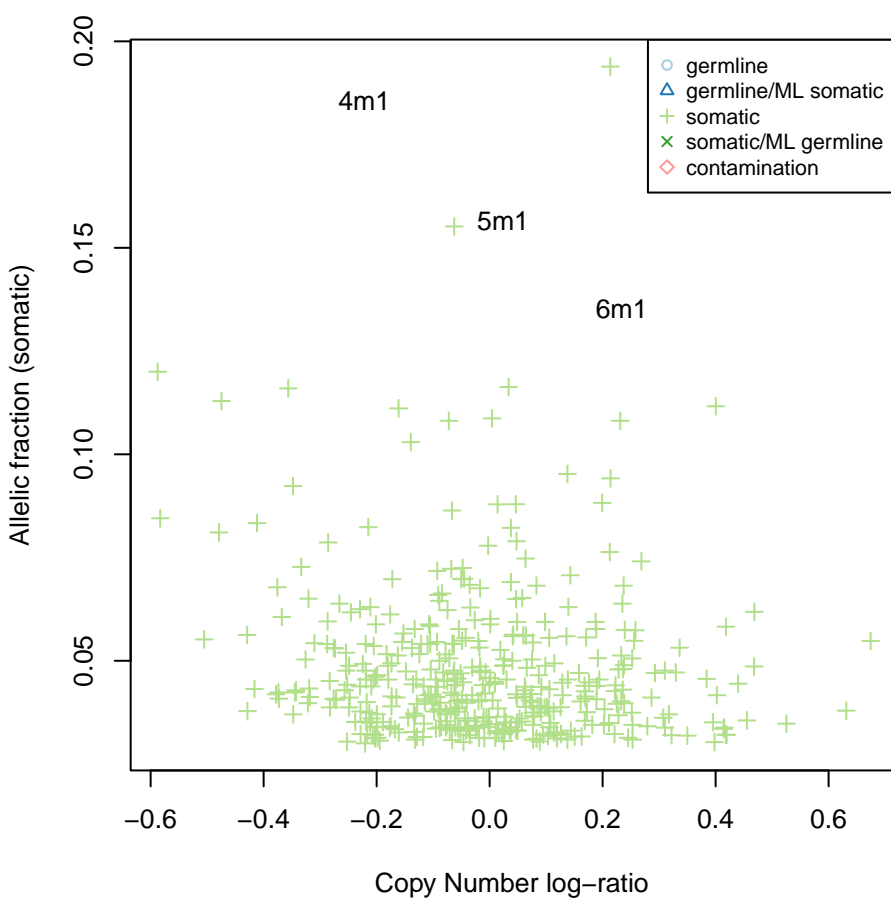
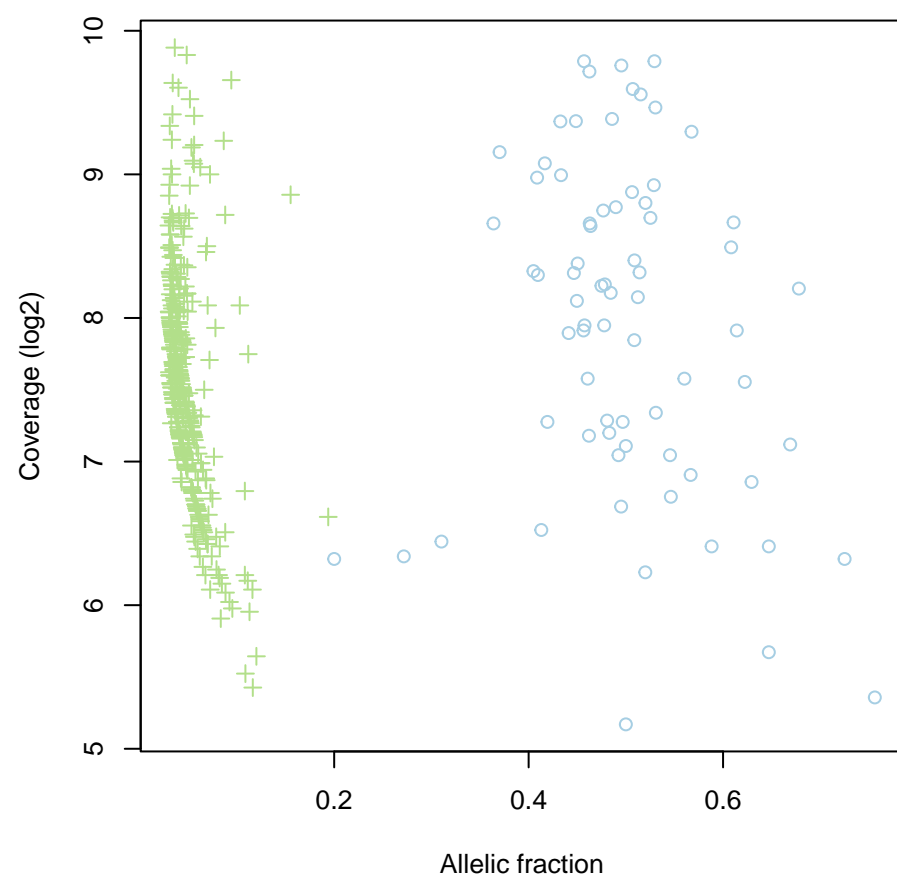
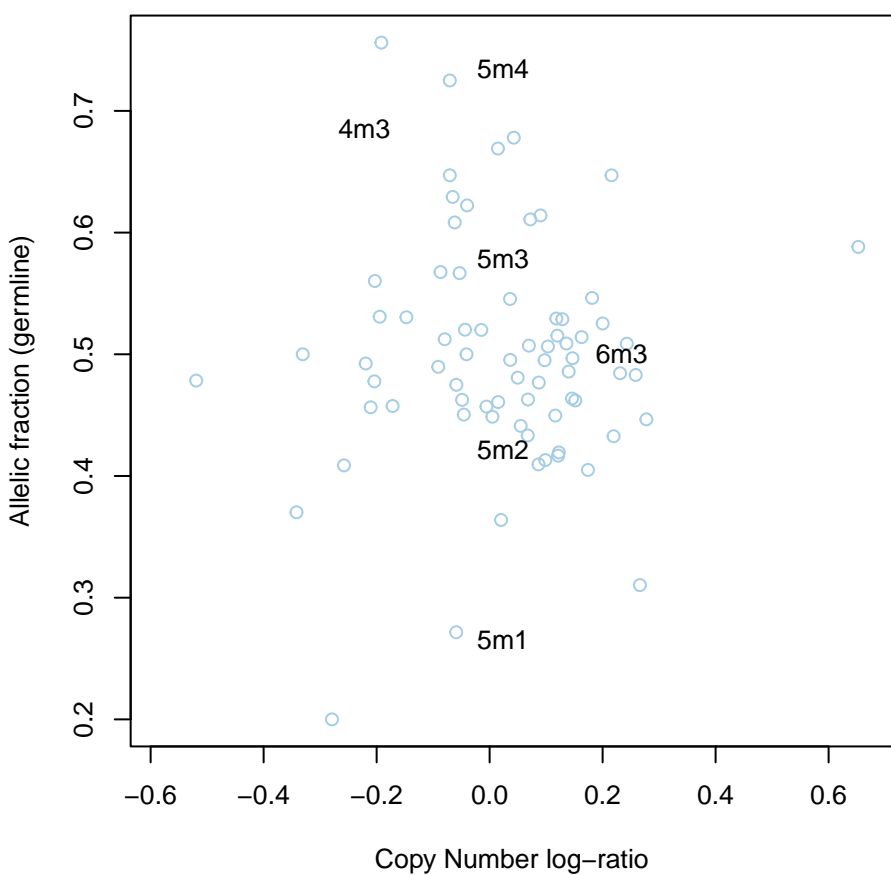


Purity: 0.59 Tumor ploidy: 4.895 SNV log-likelihood: -236.56 GoF: 70.8% Mean coverage: 233;281

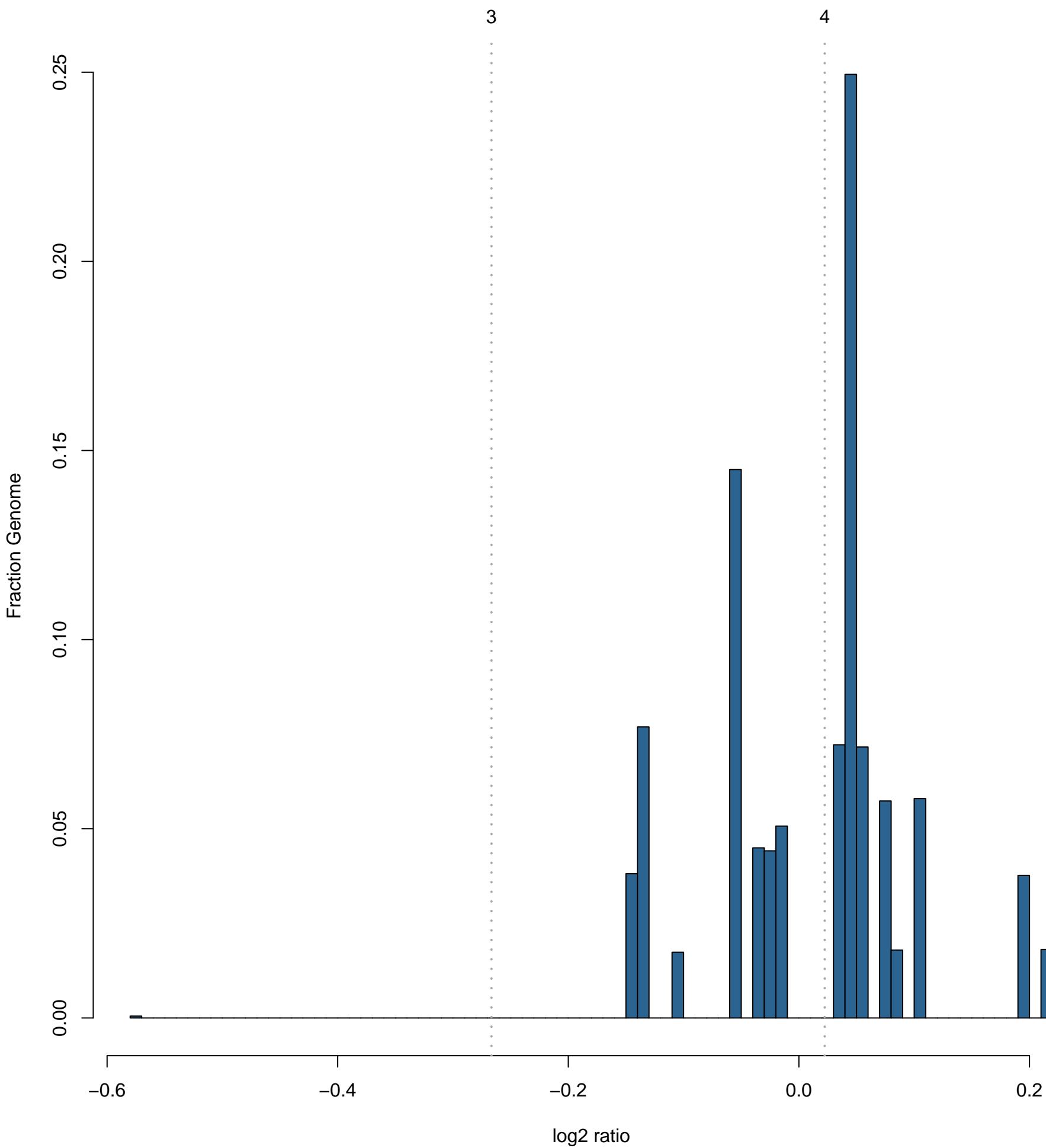


SCNA-fit log-likelihood: -5297.13

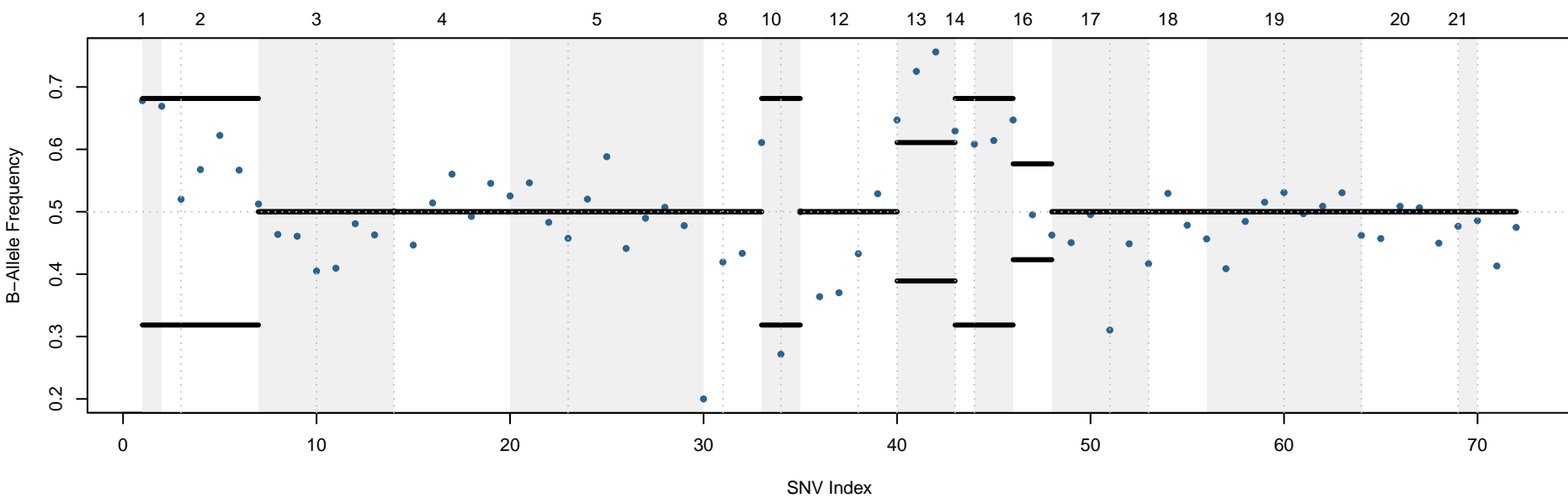




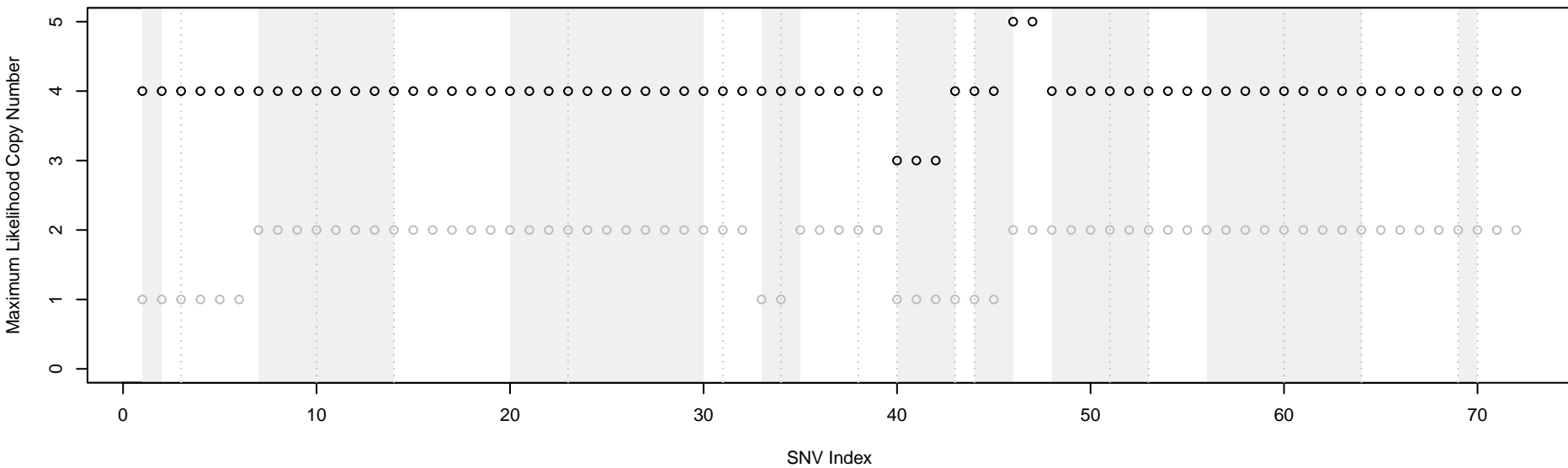
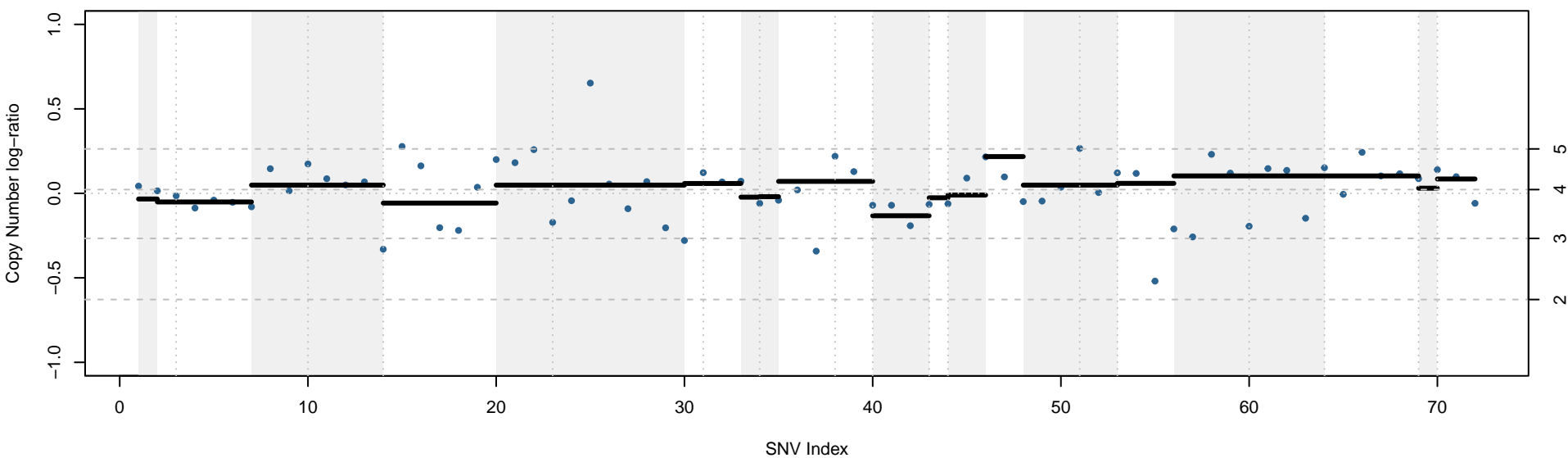
Purity: 0.57 Tumor ploidy: 3.915

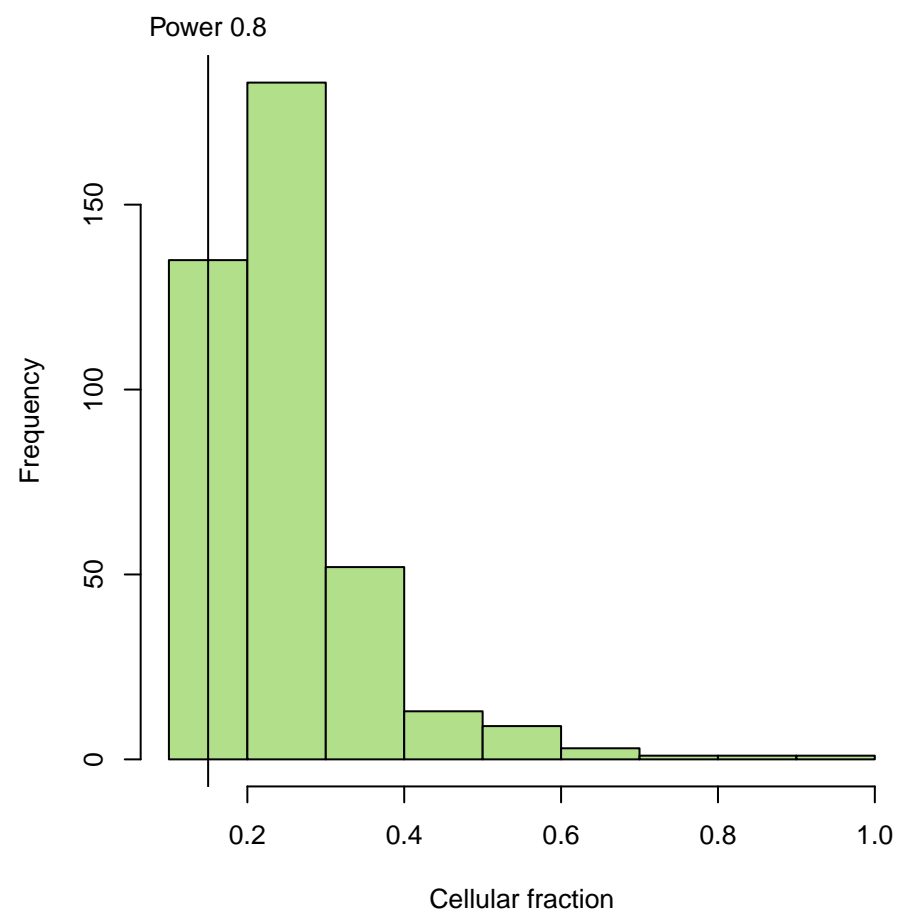
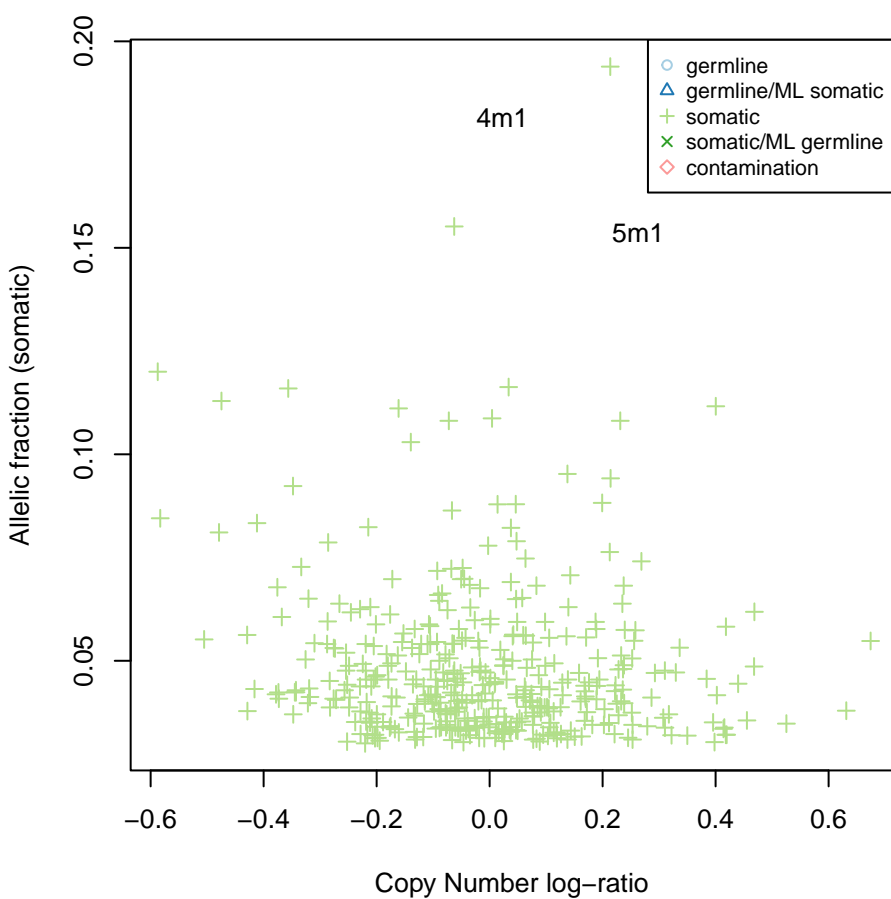
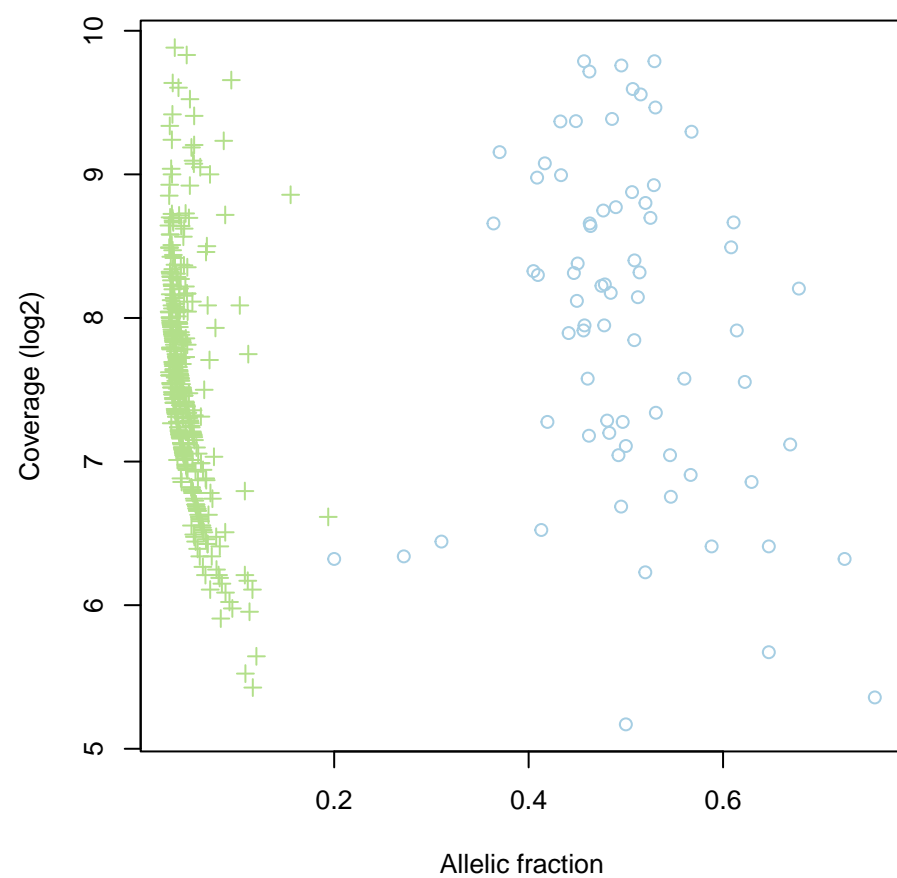
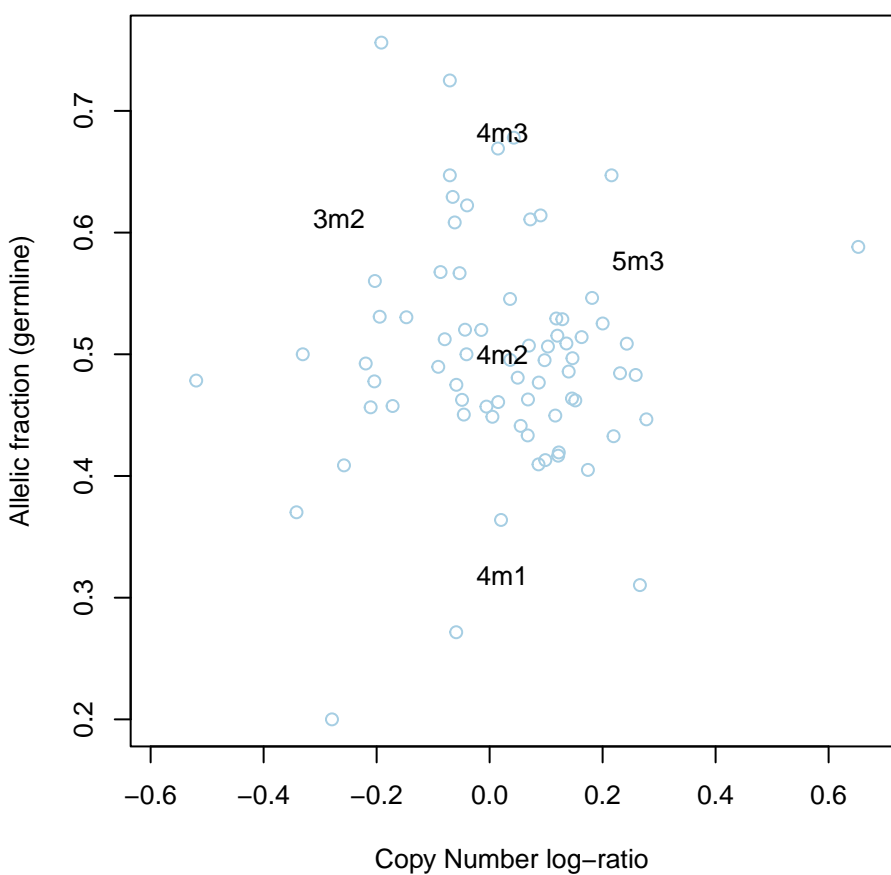


Purity: 0.57 Tumor ploidy: 3.915 SNV log-likelihood: -237.21 GoF: 56.4% Mean coverage: 233;281



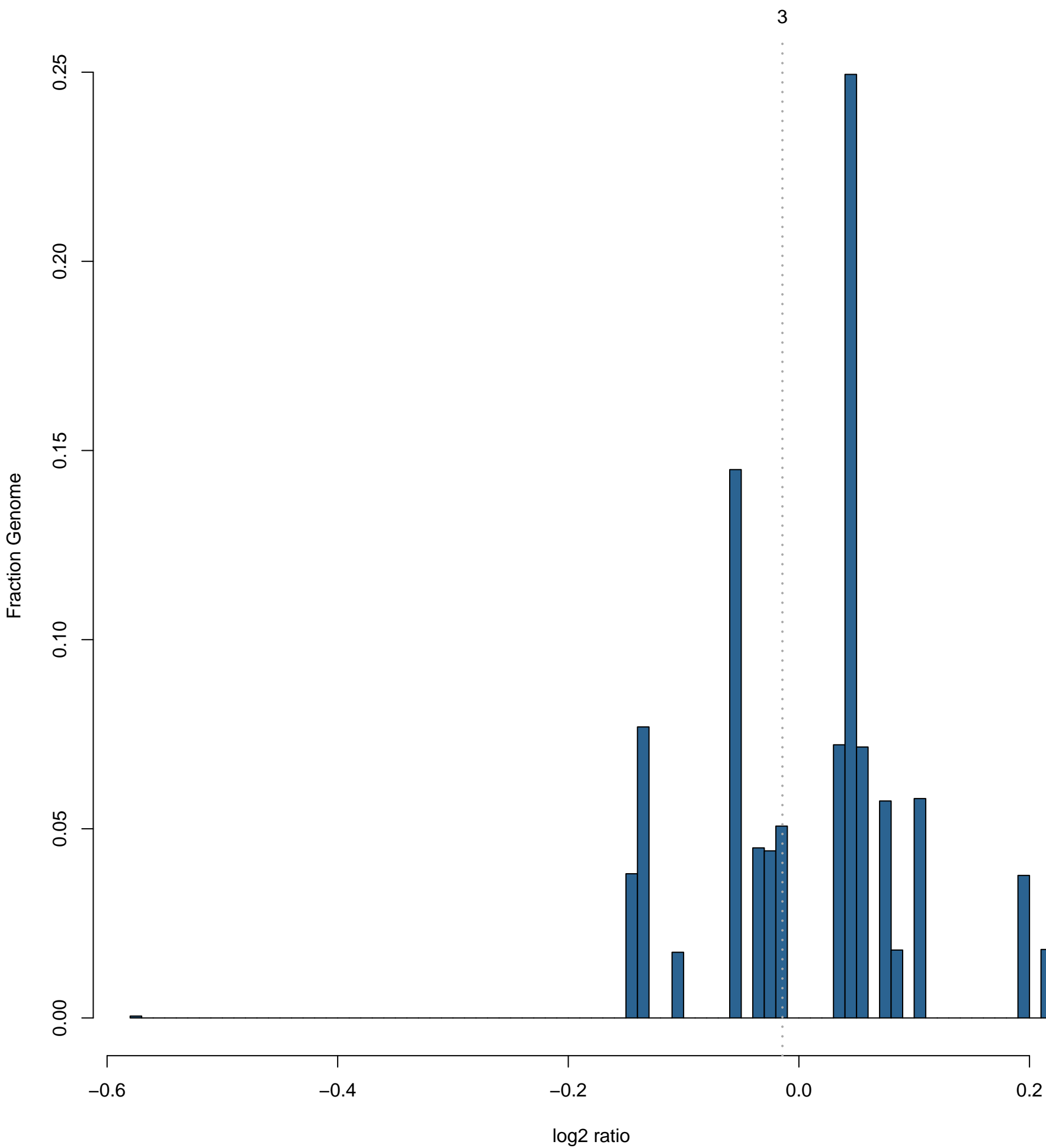
SCNA-fit log-likelihood: -5421.08



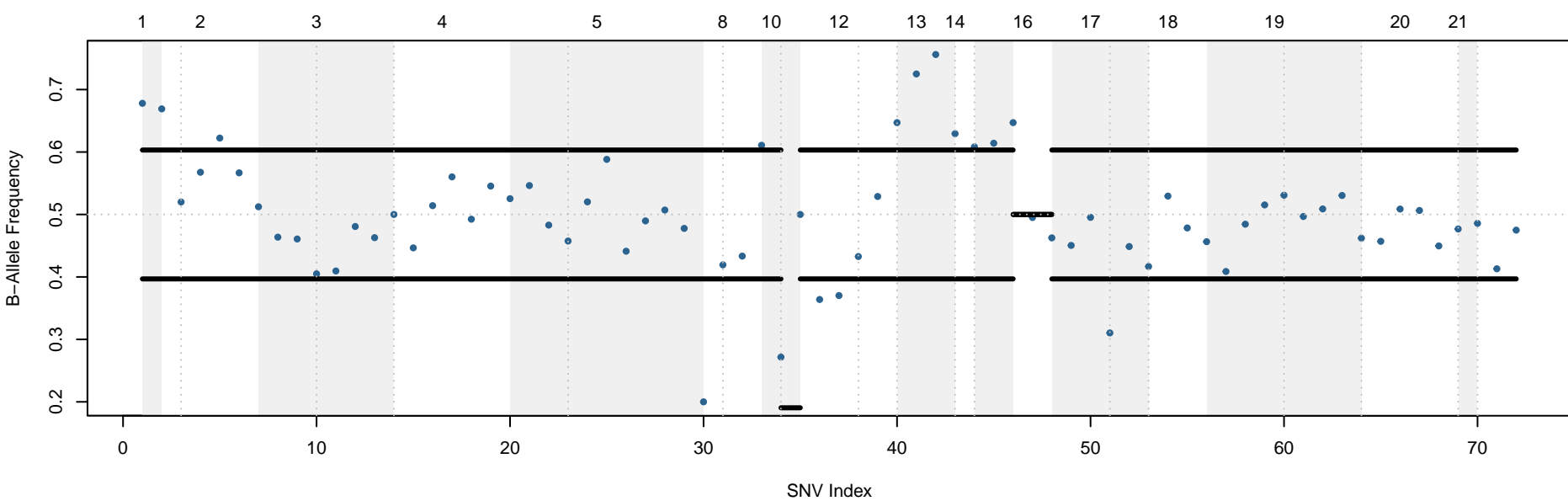




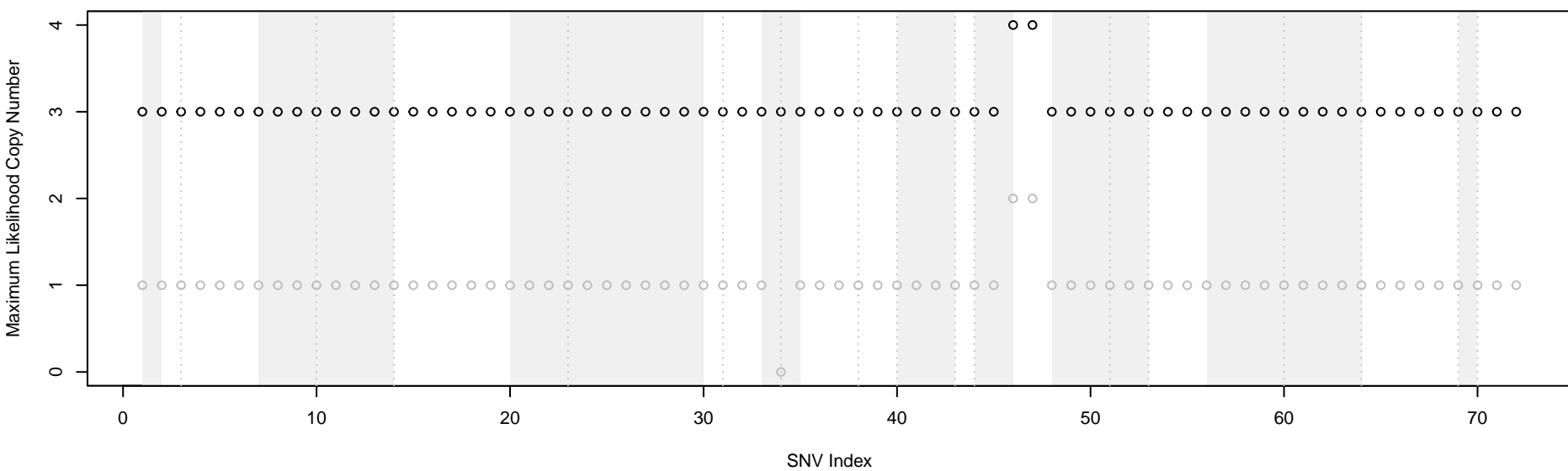
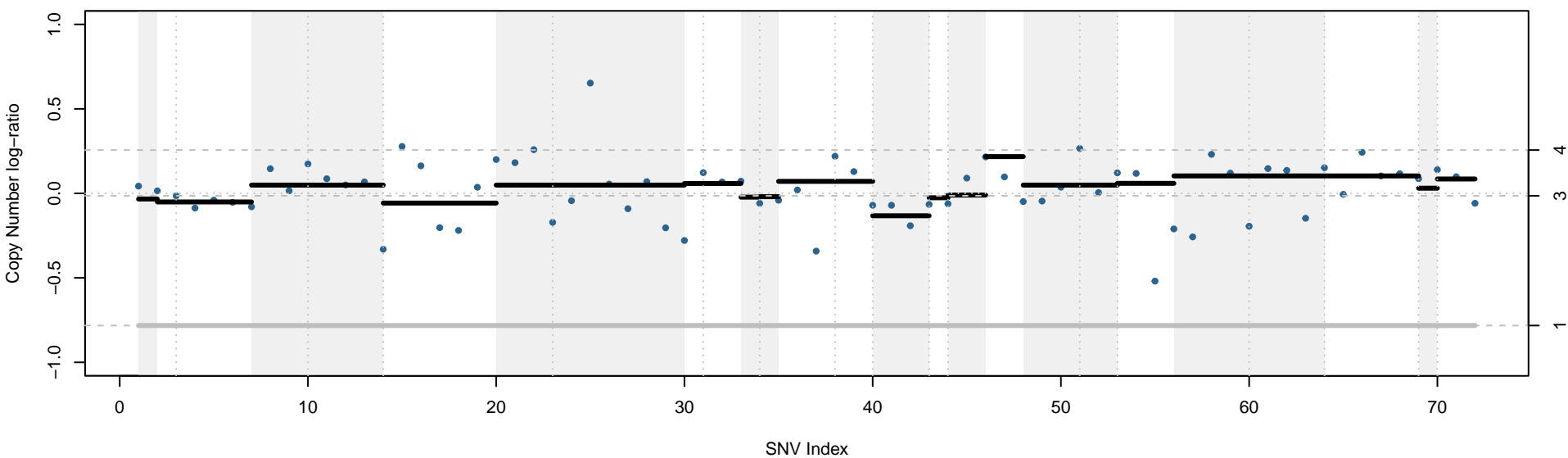
Purity: 0.52 Tumor ploidy: 3.048

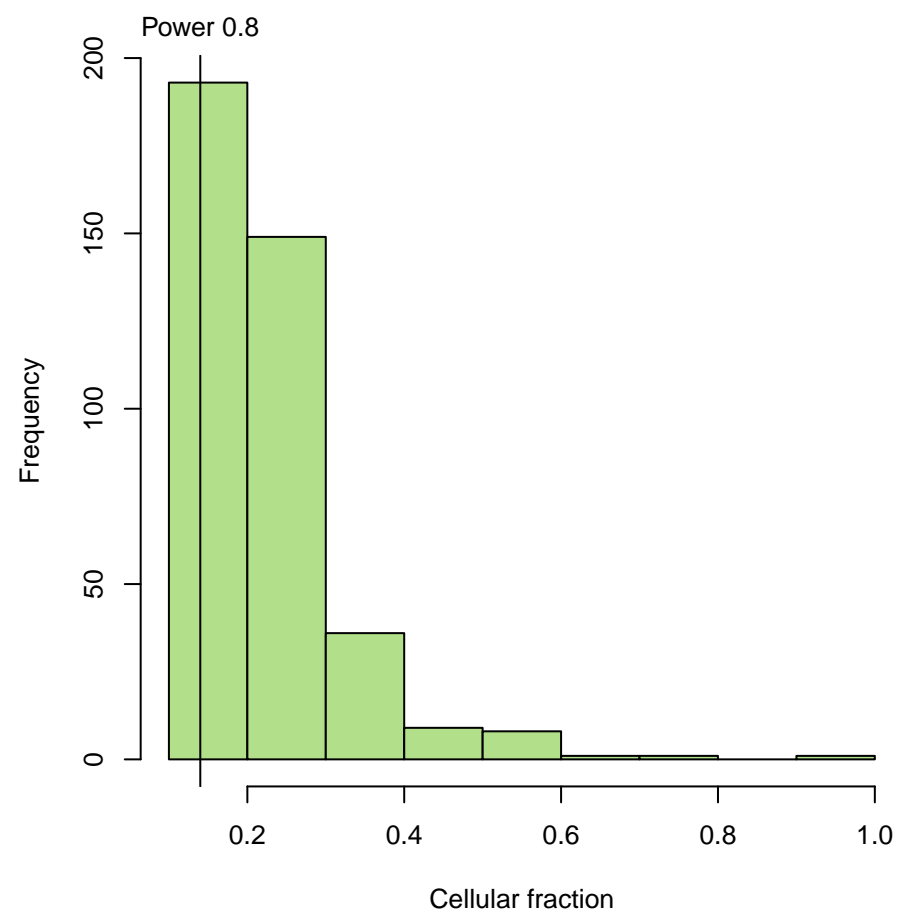
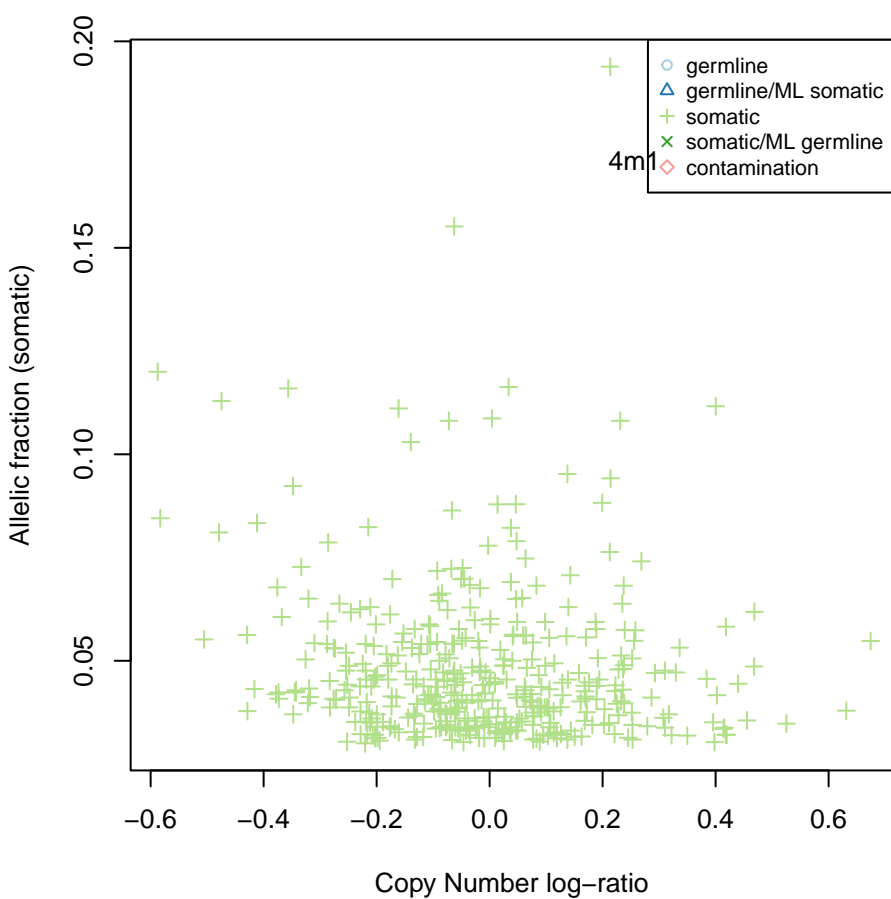
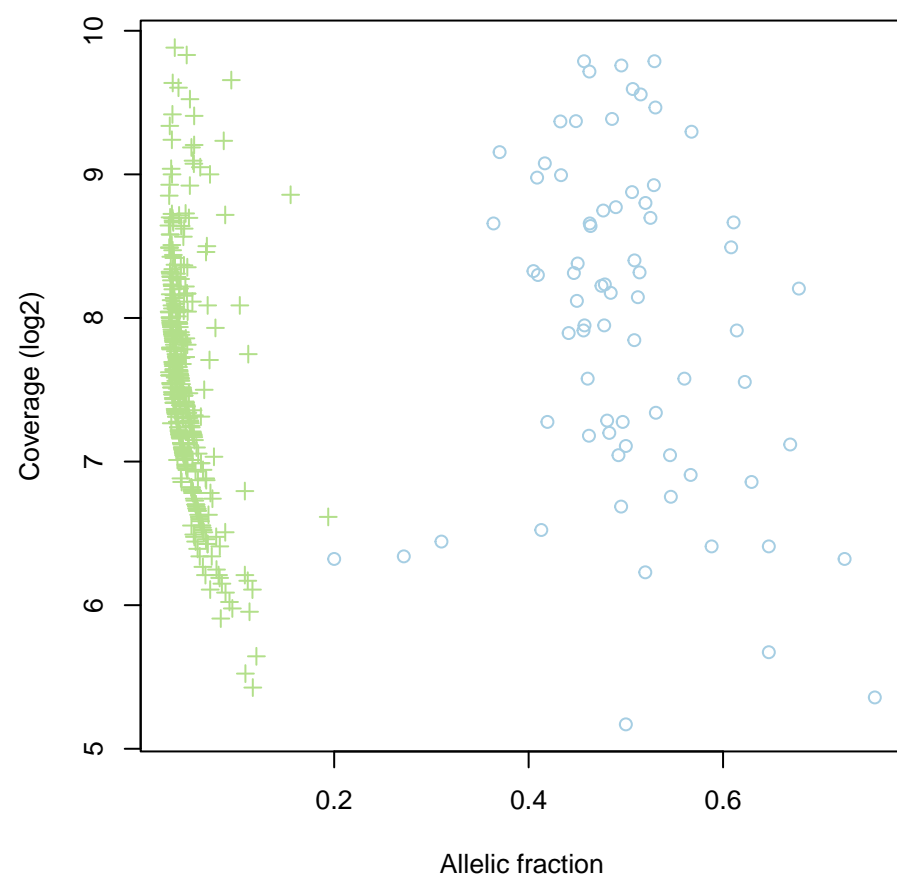
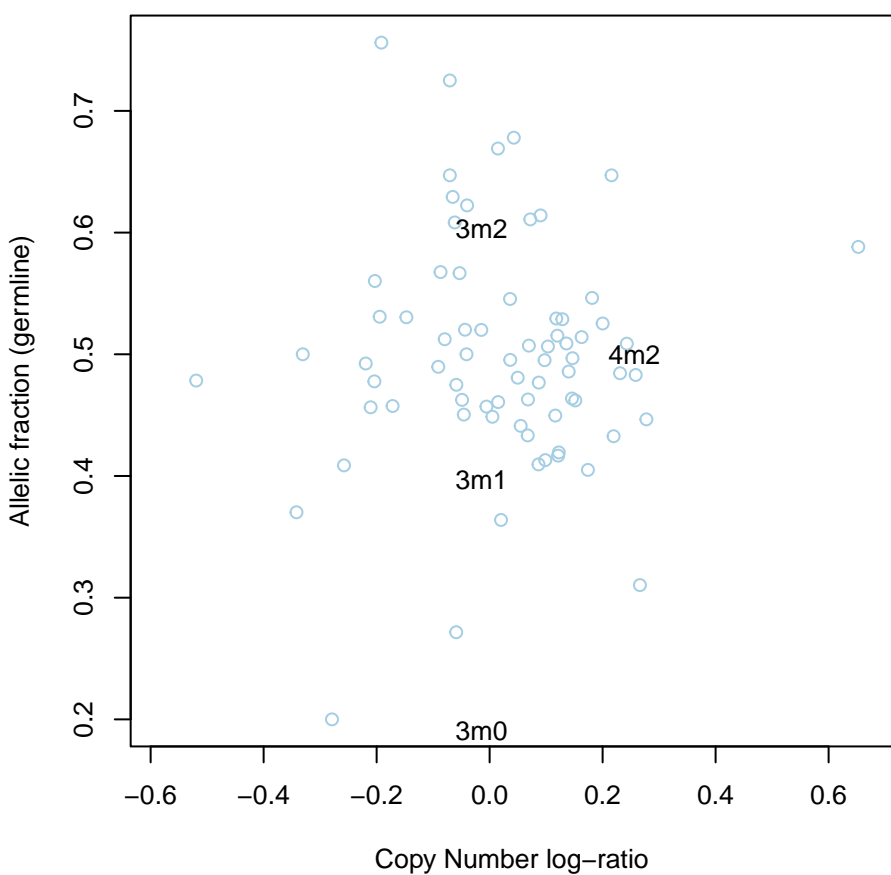


Purity: 0.52 Tumor ploidy: 3.048 SNV log-likelihood: -394.9 GoF: 44.8% Mean coverage: 233;281

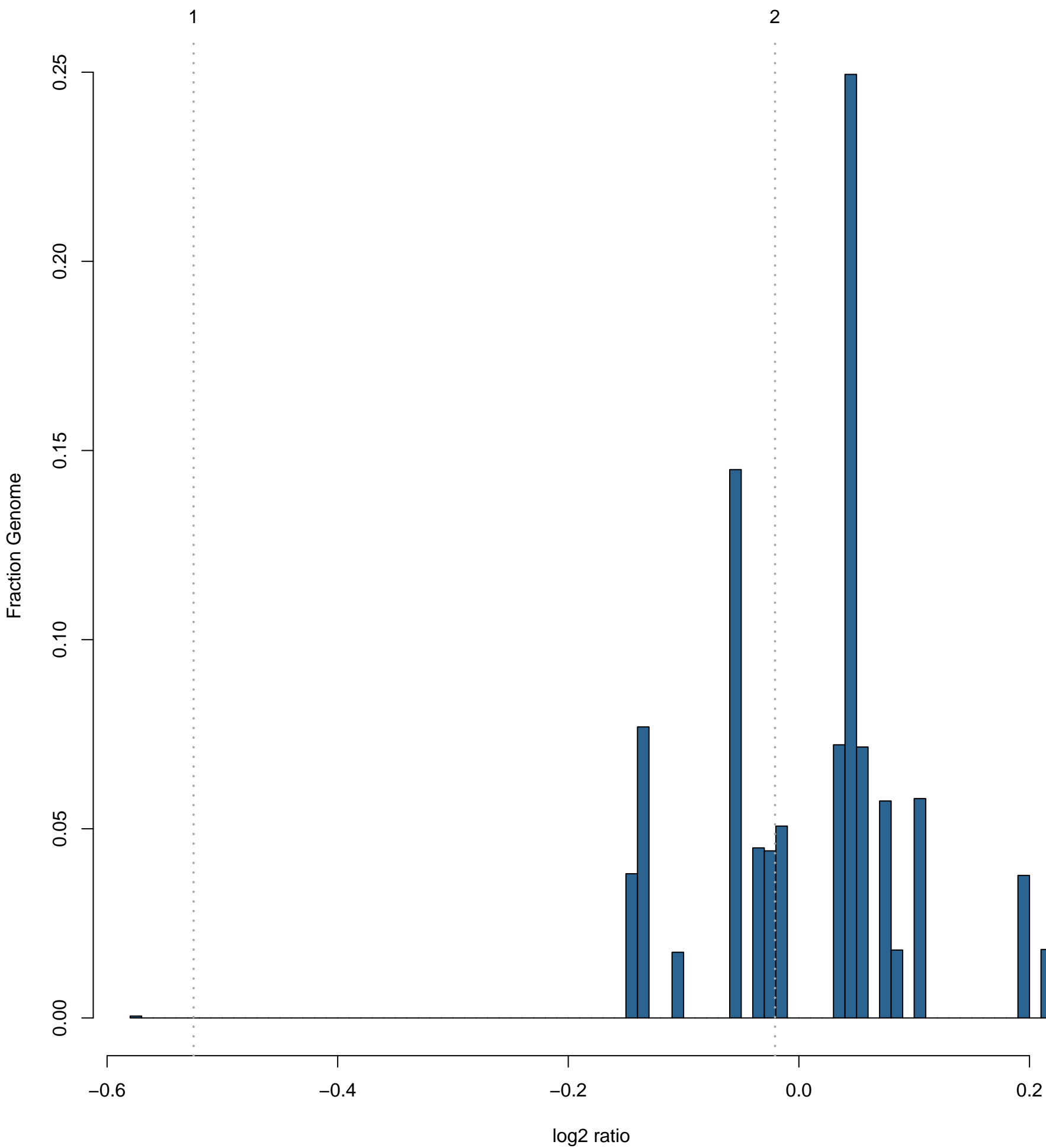


SCNA-fit log-likelihood: -5518.4

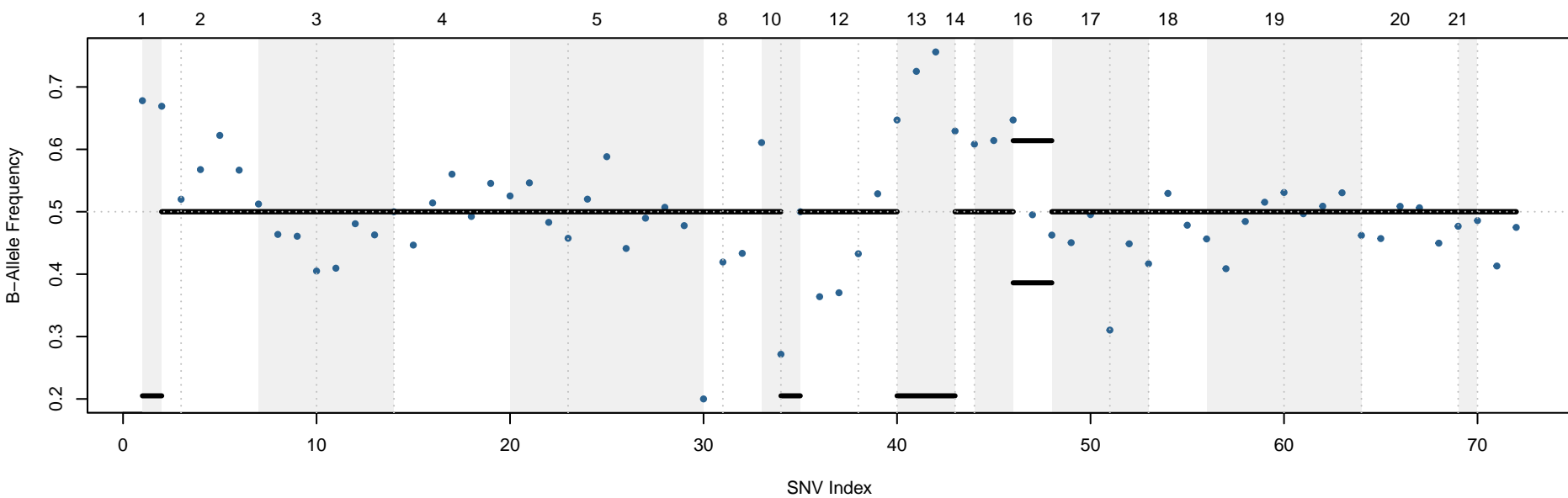




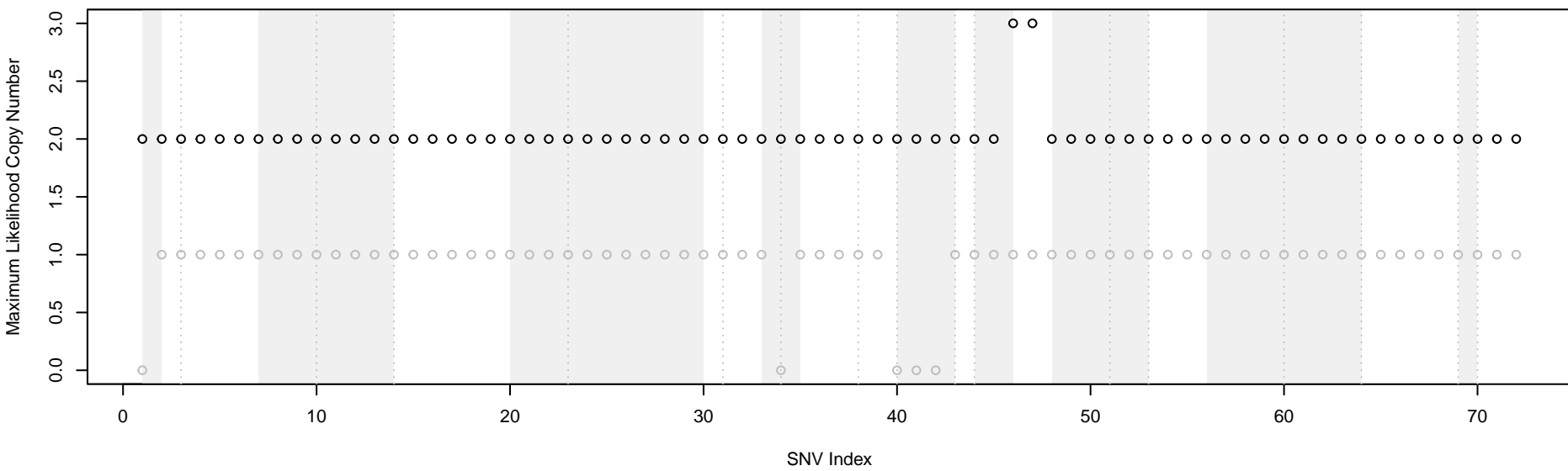
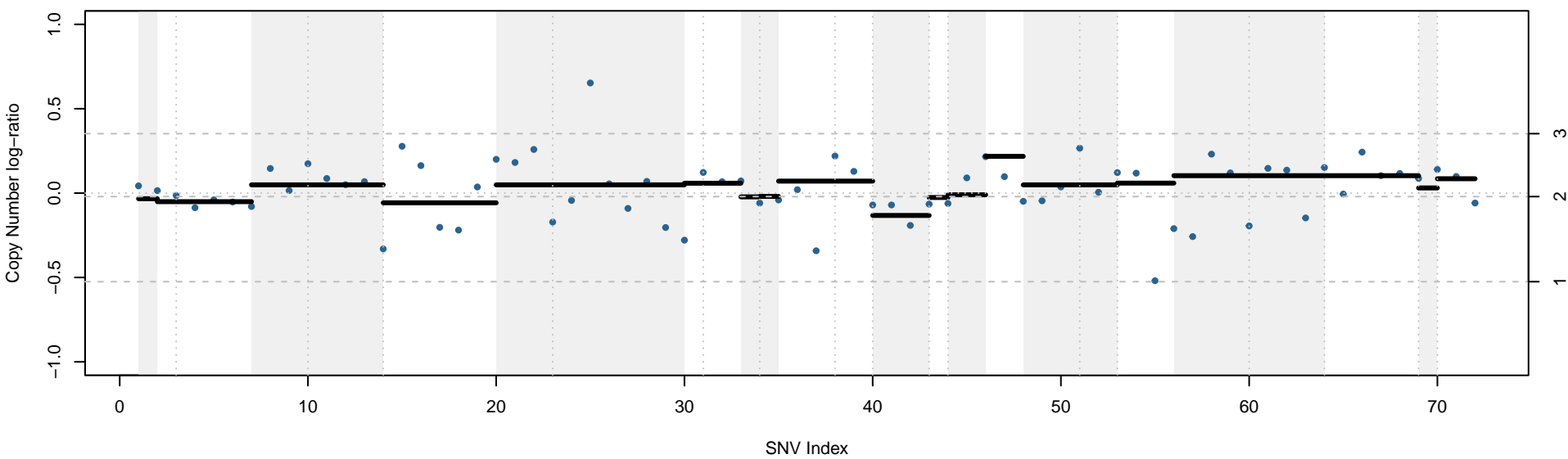
Purity: 0.59 Tumor ploidy: 2.049

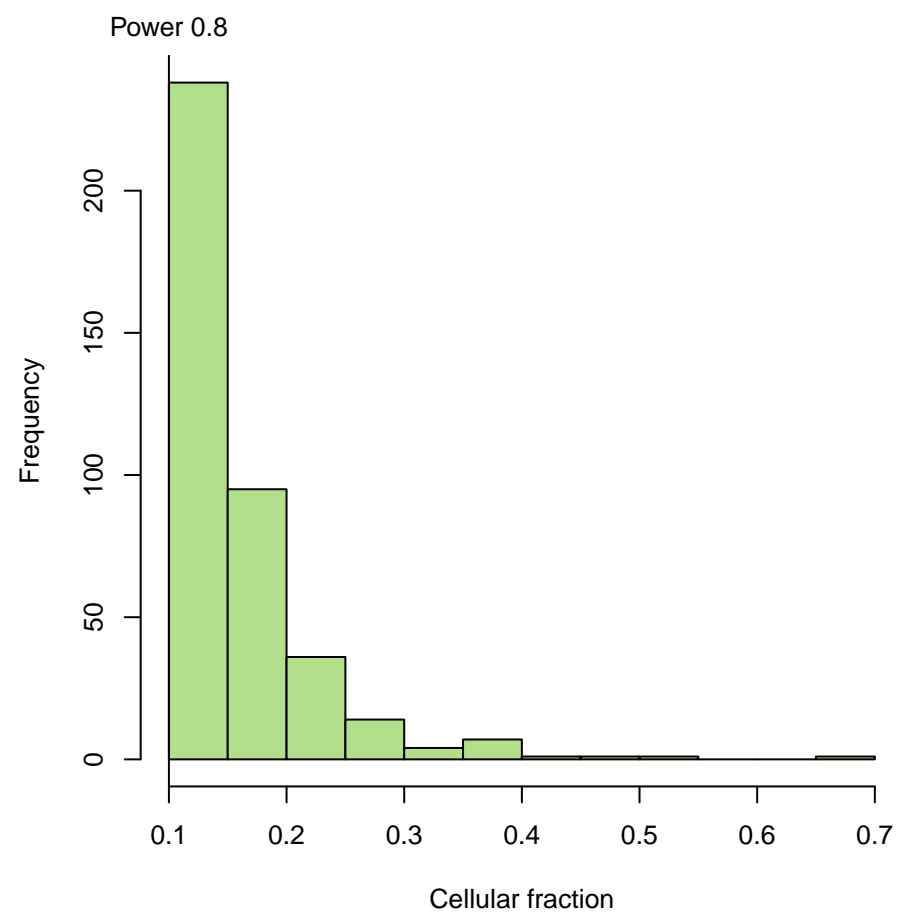
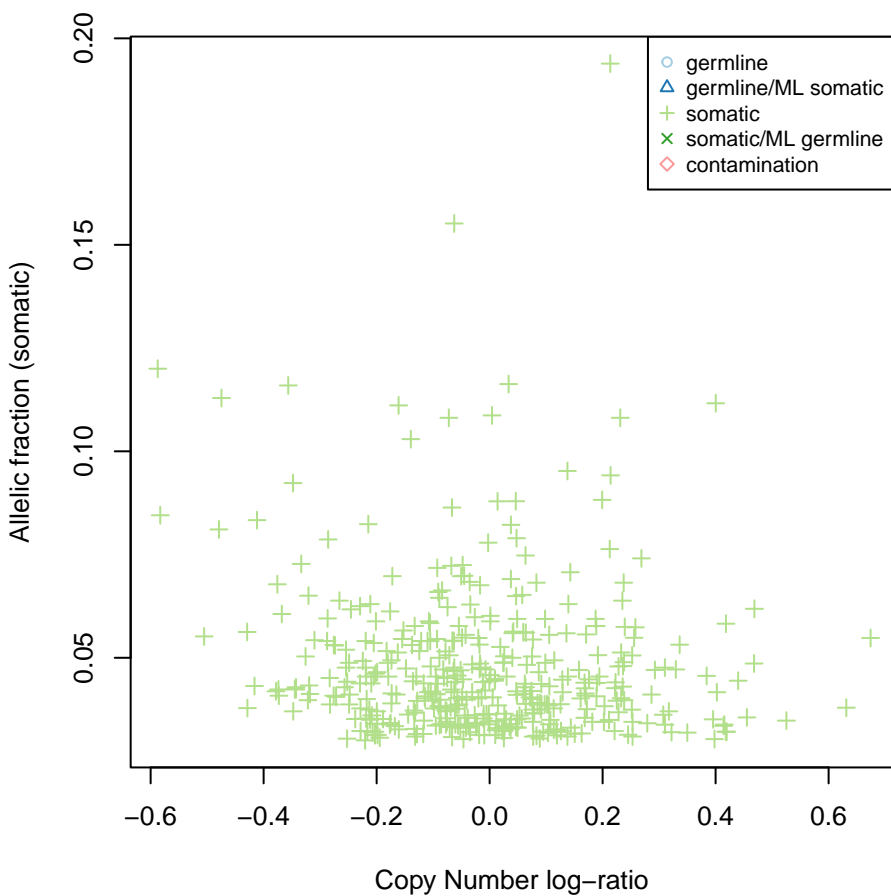
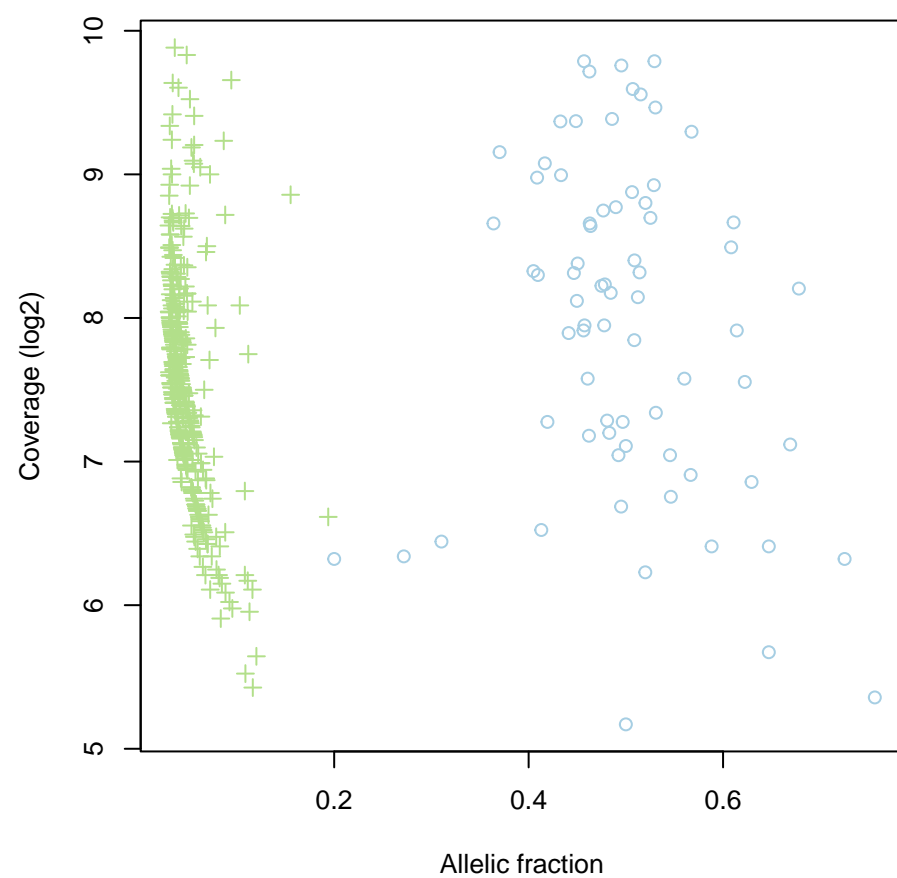
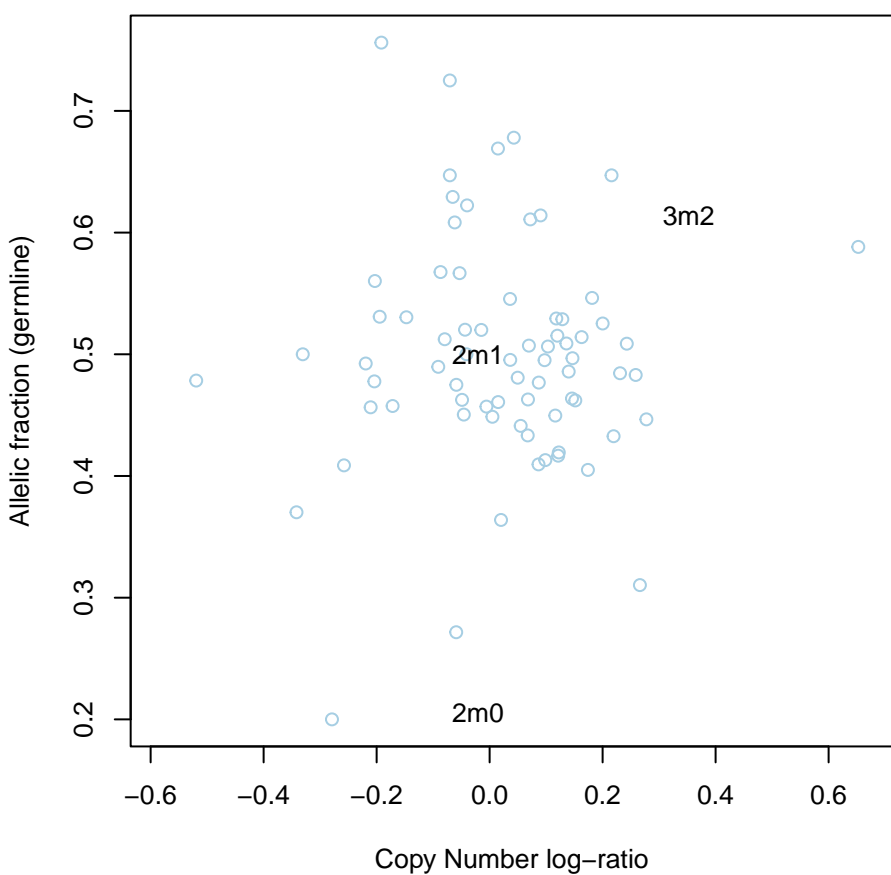


Purity: 0.59 Tumor ploidy: 2.049 SNV log-likelihood: -1158.87 GoF: 0% Mean coverage: 233;281

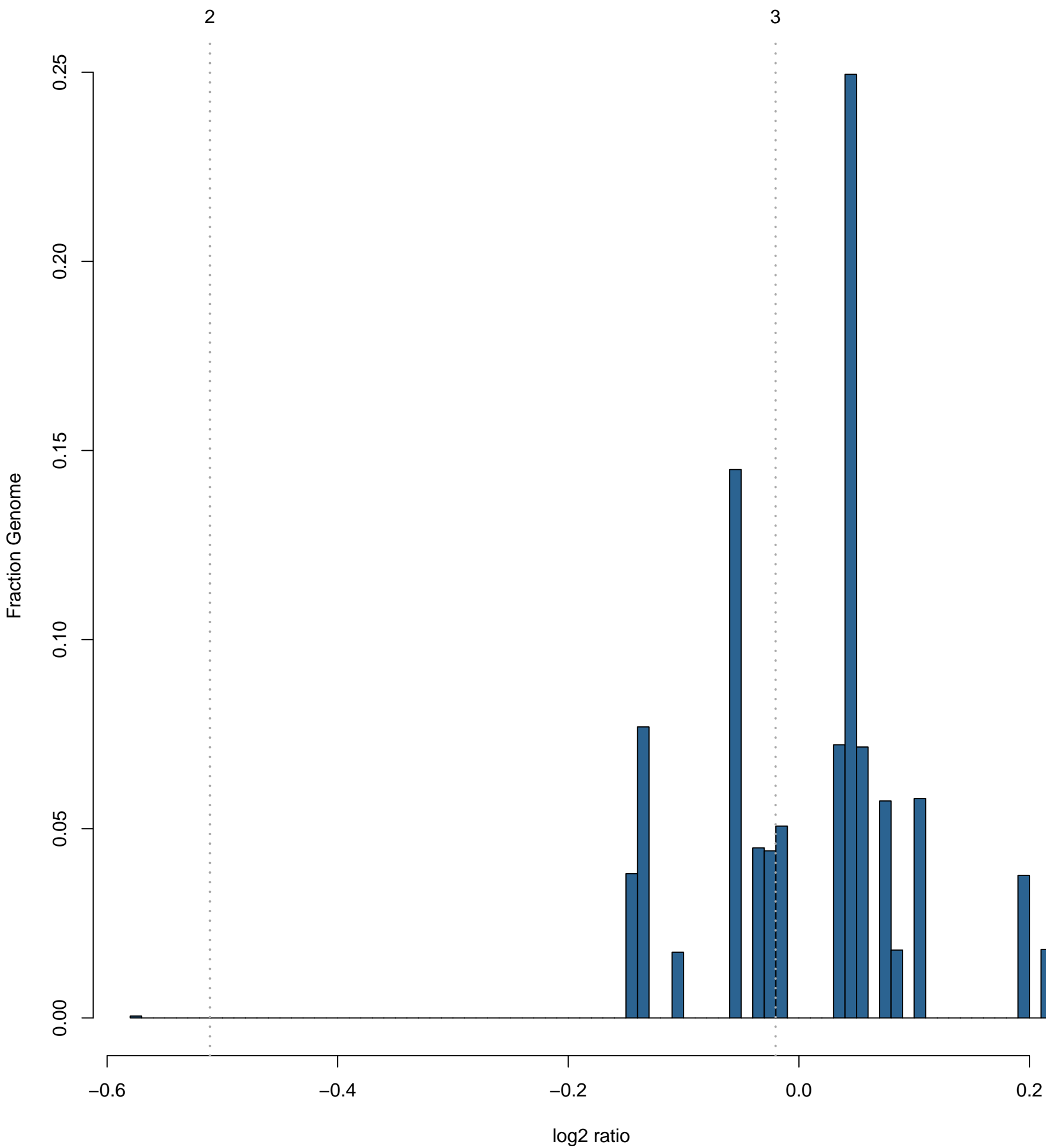


SCNA-fit log-likelihood: -5649.4

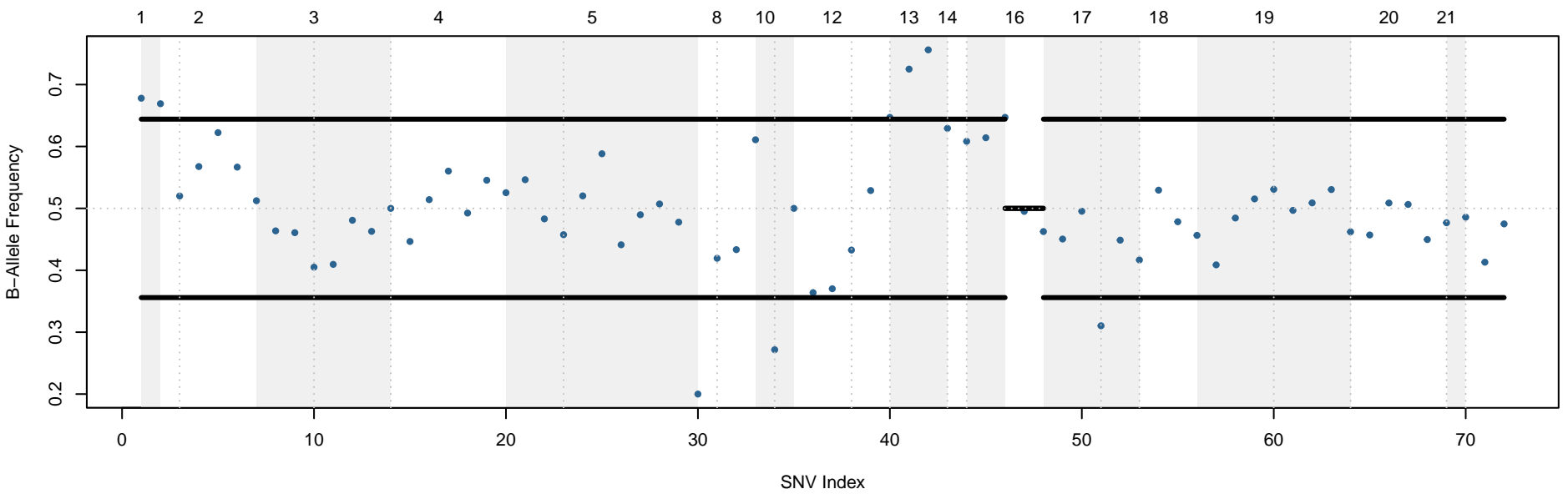




Purity: 0.81 Tumor ploidy: 3.049



Purity: 0.81 Tumor ploidy: 3.049 SNV log-likelihood: -1580.69 GoF: 0% Mean coverage: 233;281



SCNA-fit log-likelihood: -5636.16

