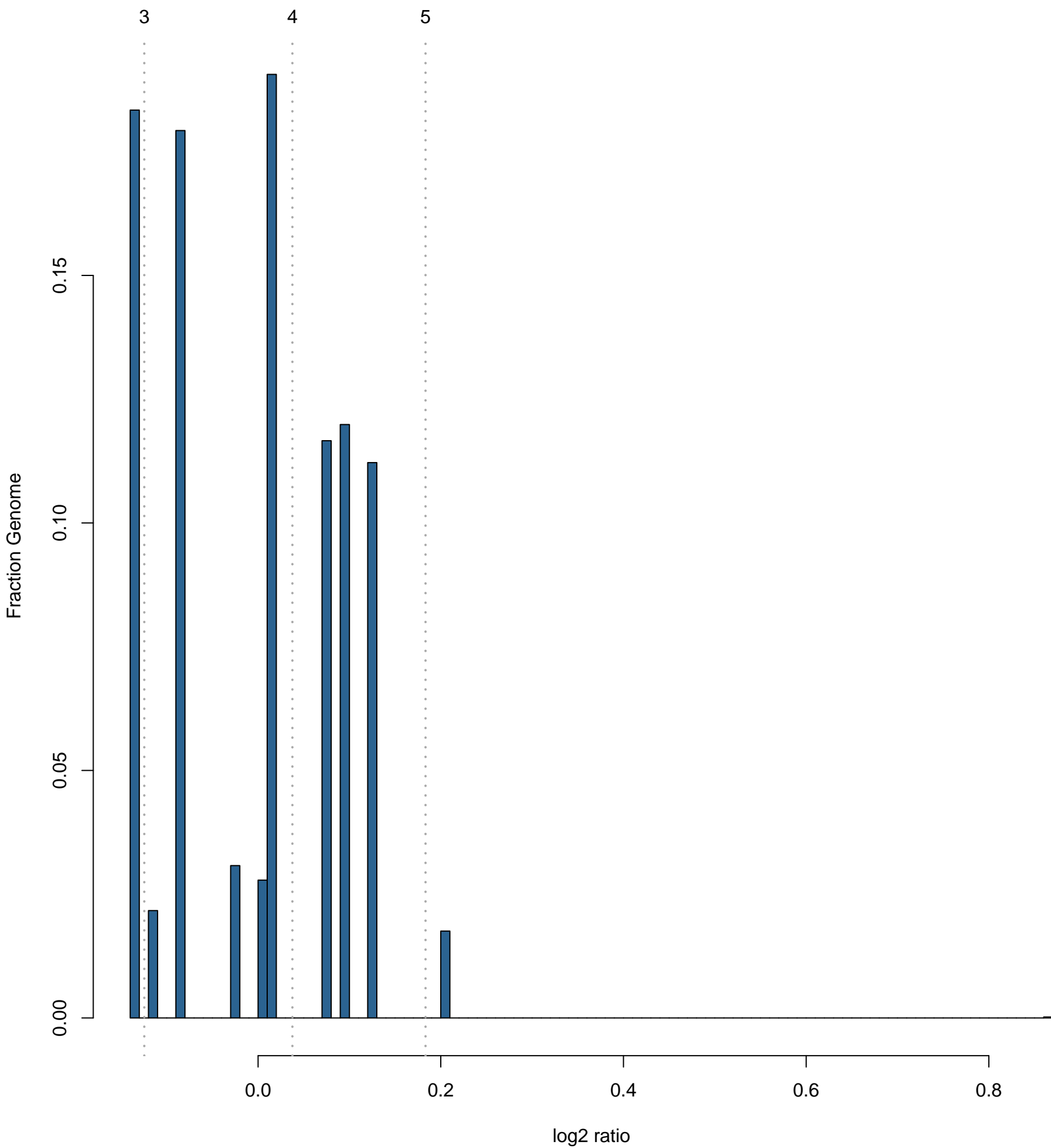
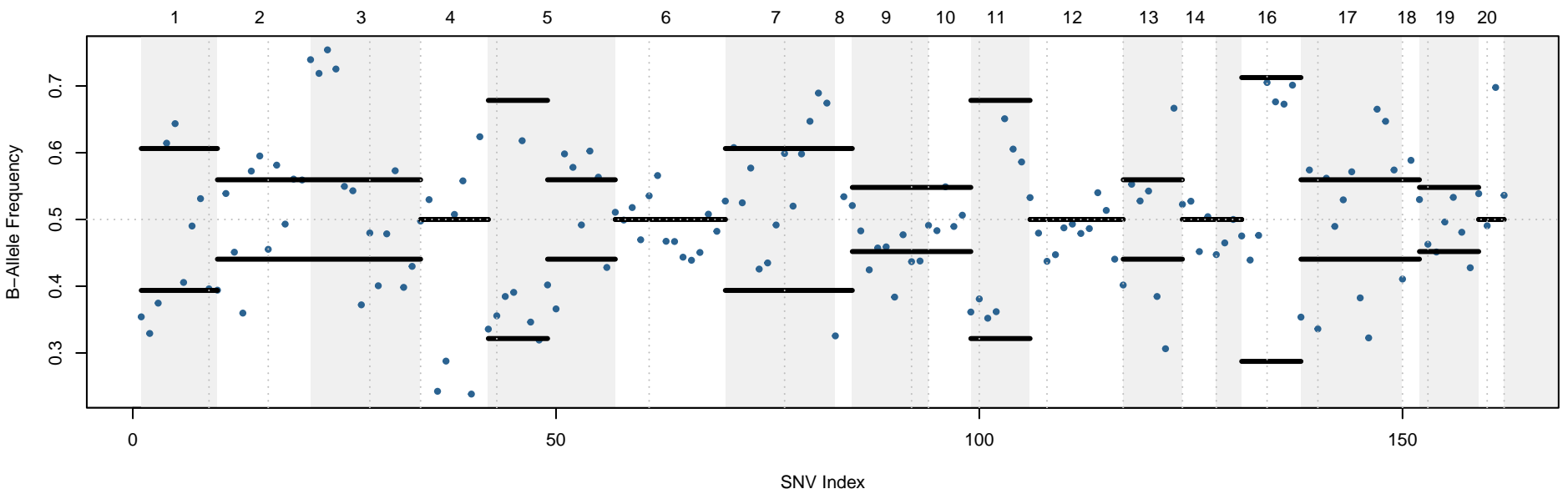


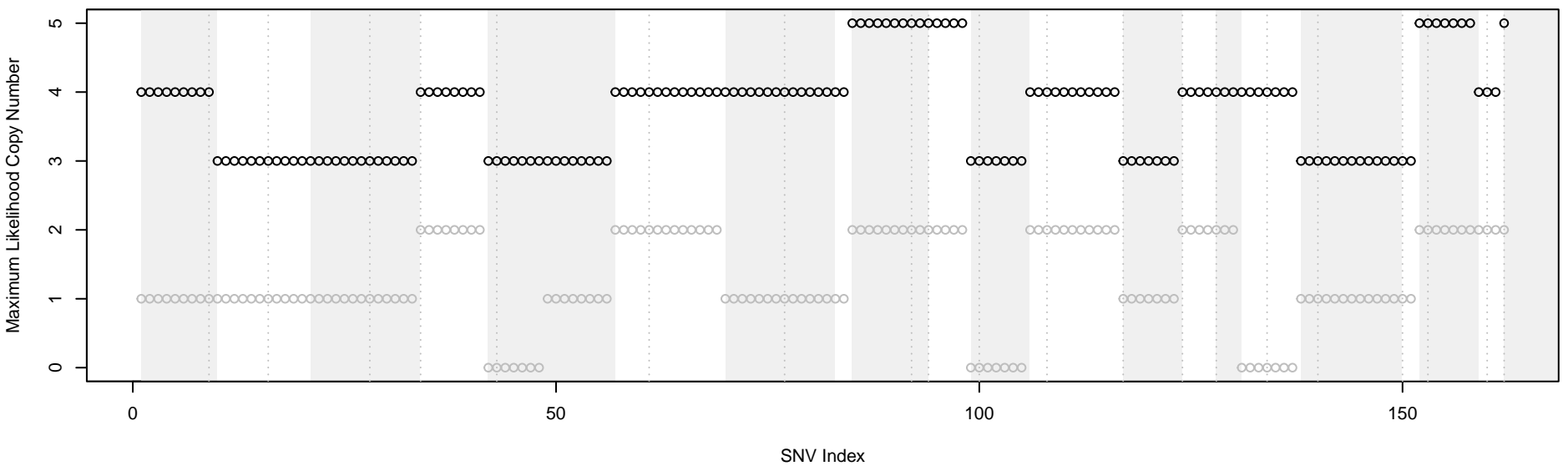
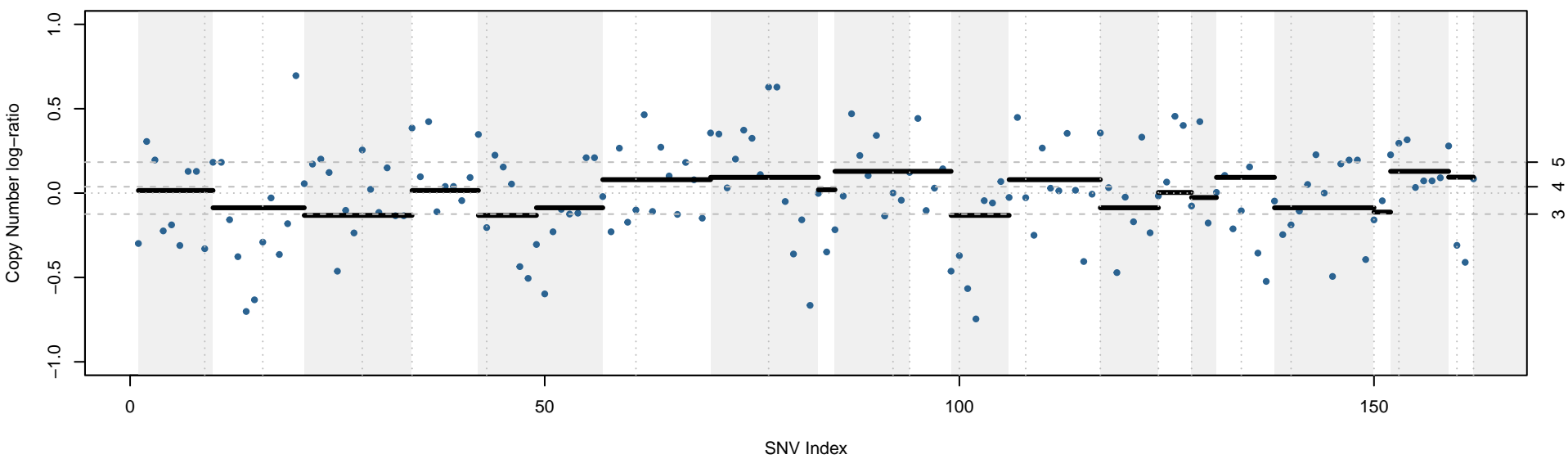
Purity: 0.27 Tumor ploidy: 3.758

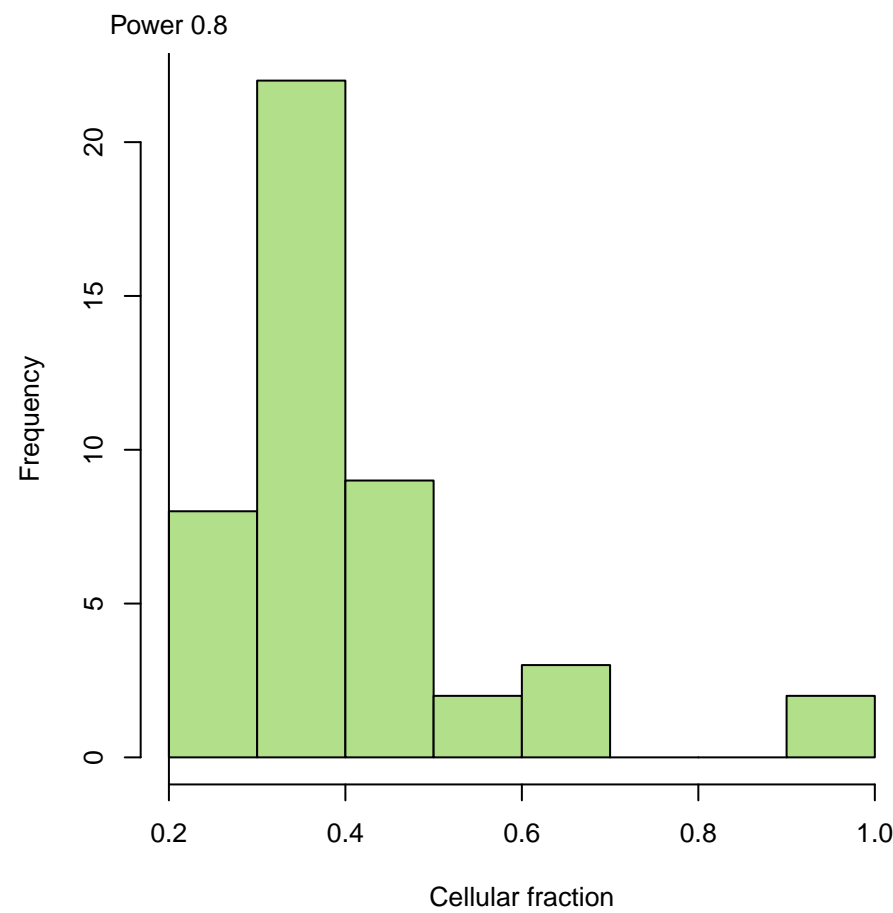
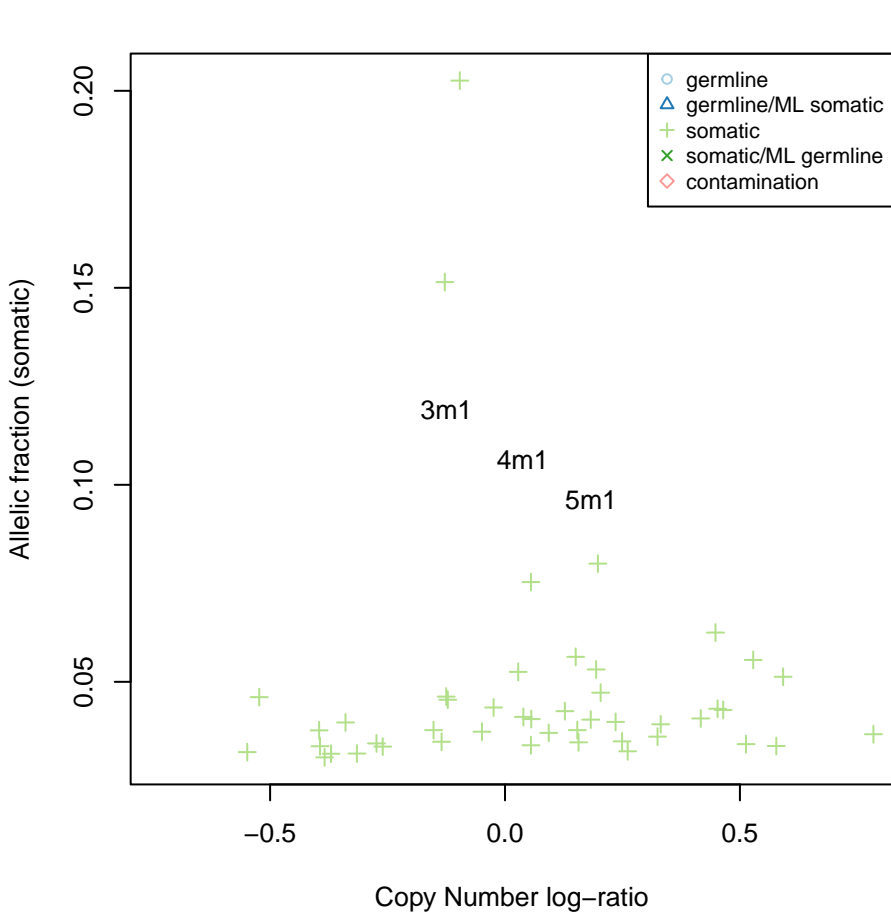
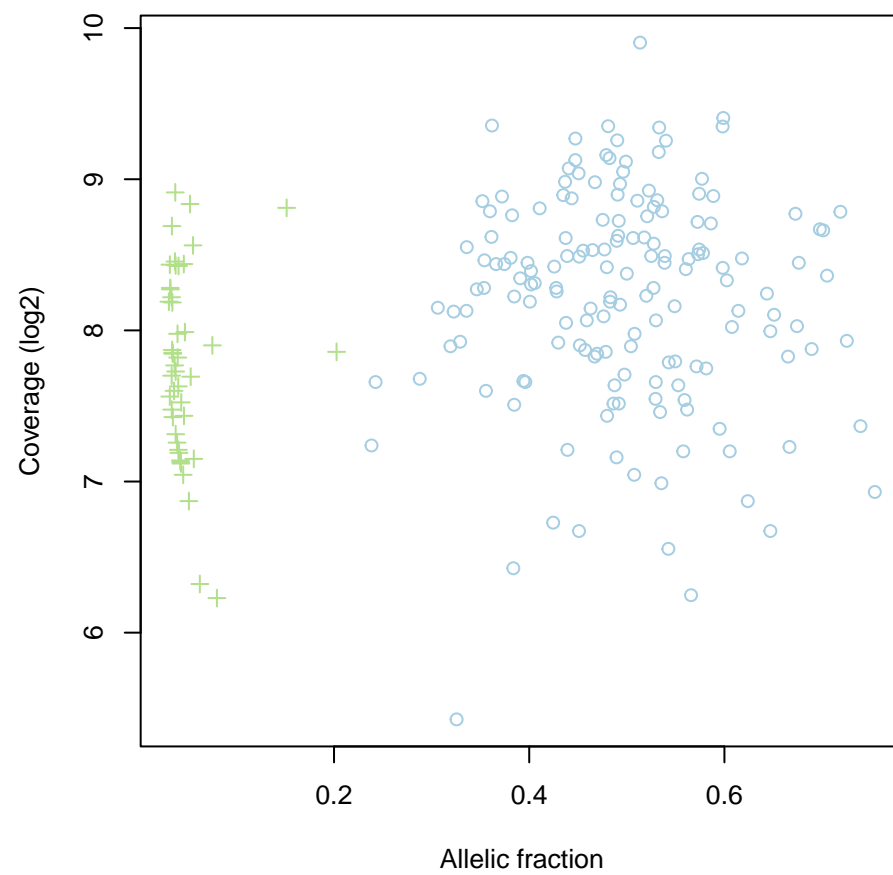
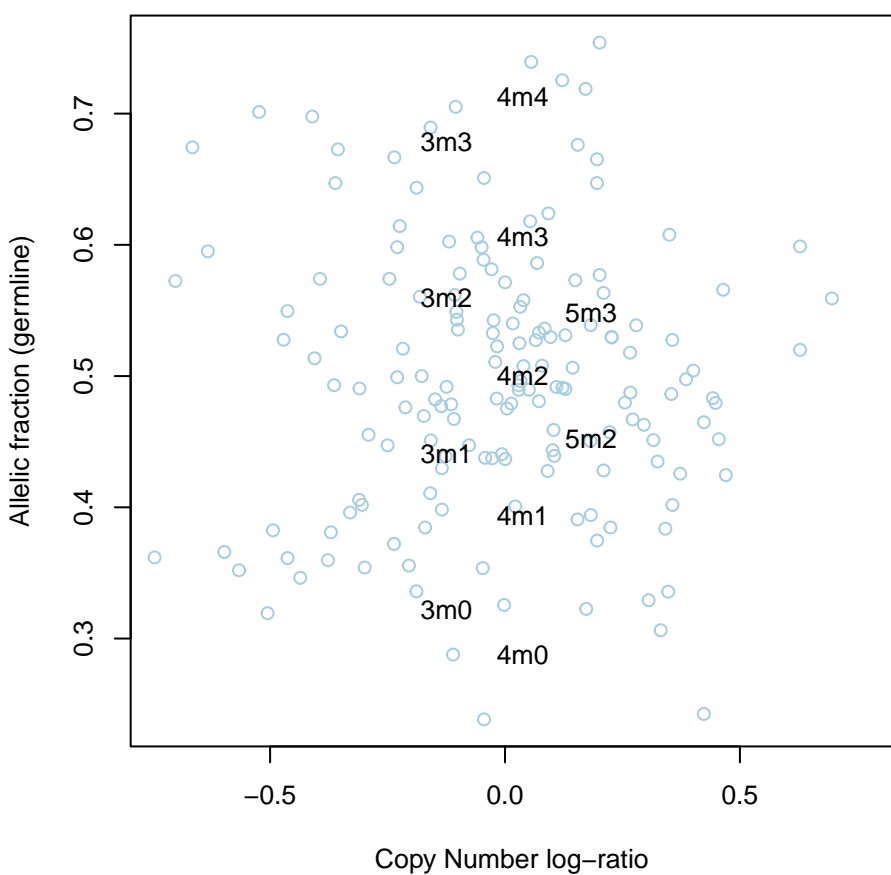


Purity: 0.27 Tumor ploidy: 3.758 SNV log-likelihood: -176.96 GoF: 93.5% Mean coverage: 312;424

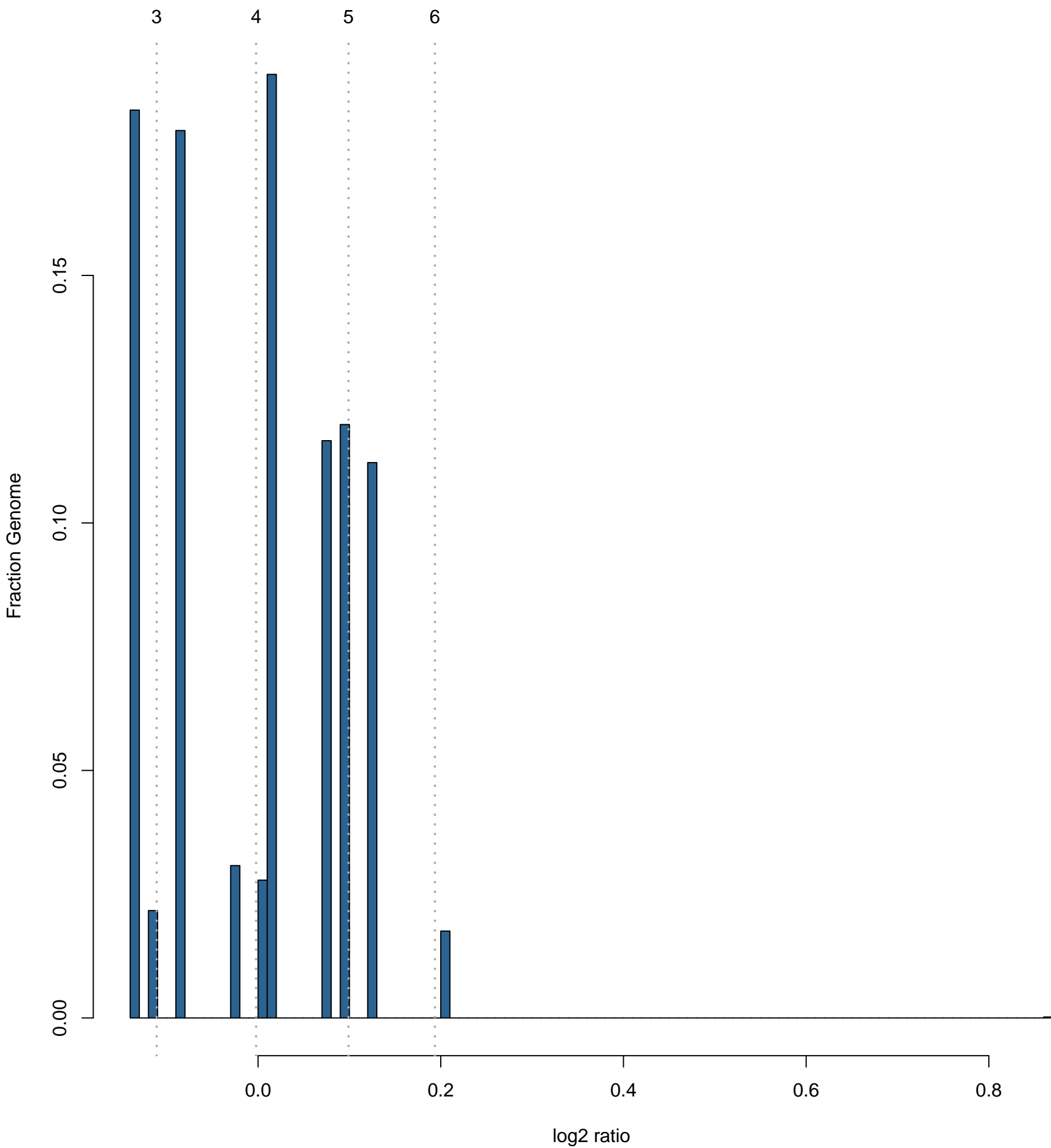


SCNA-fit log-likelihood: -19401.5

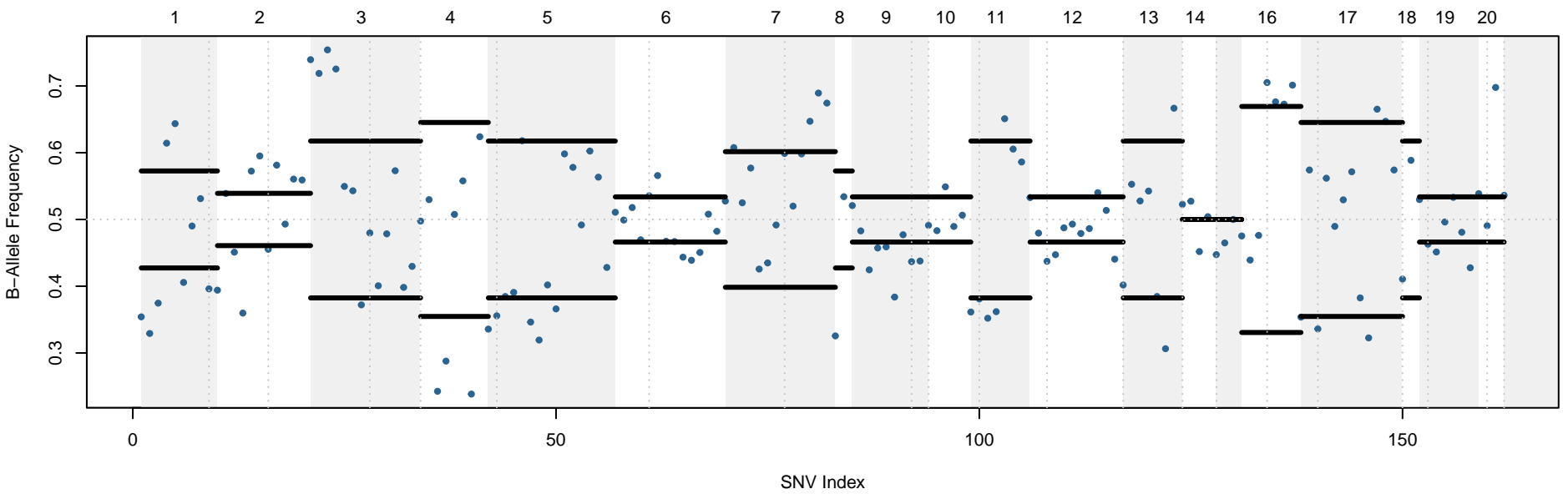




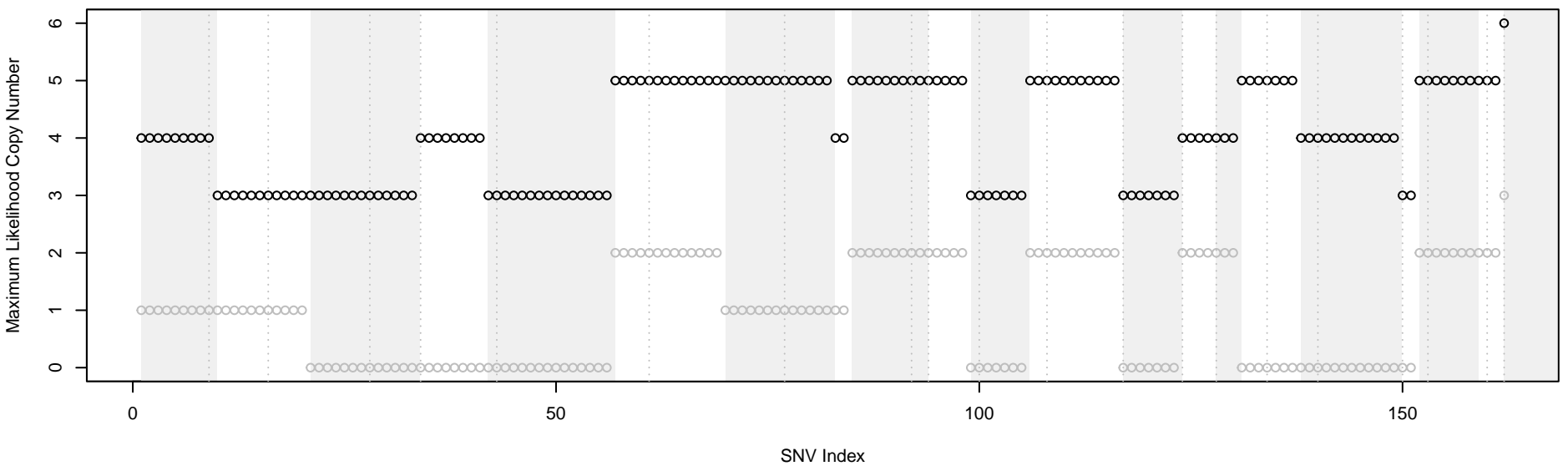
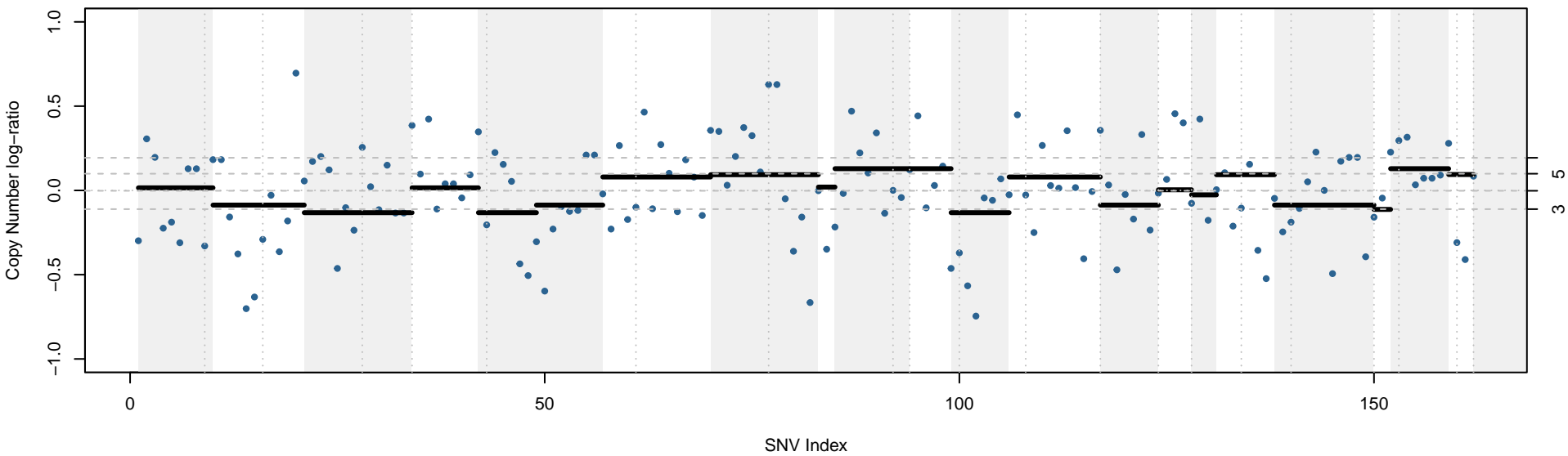
Purity: 0.17 Tumor ploidy: 4.021

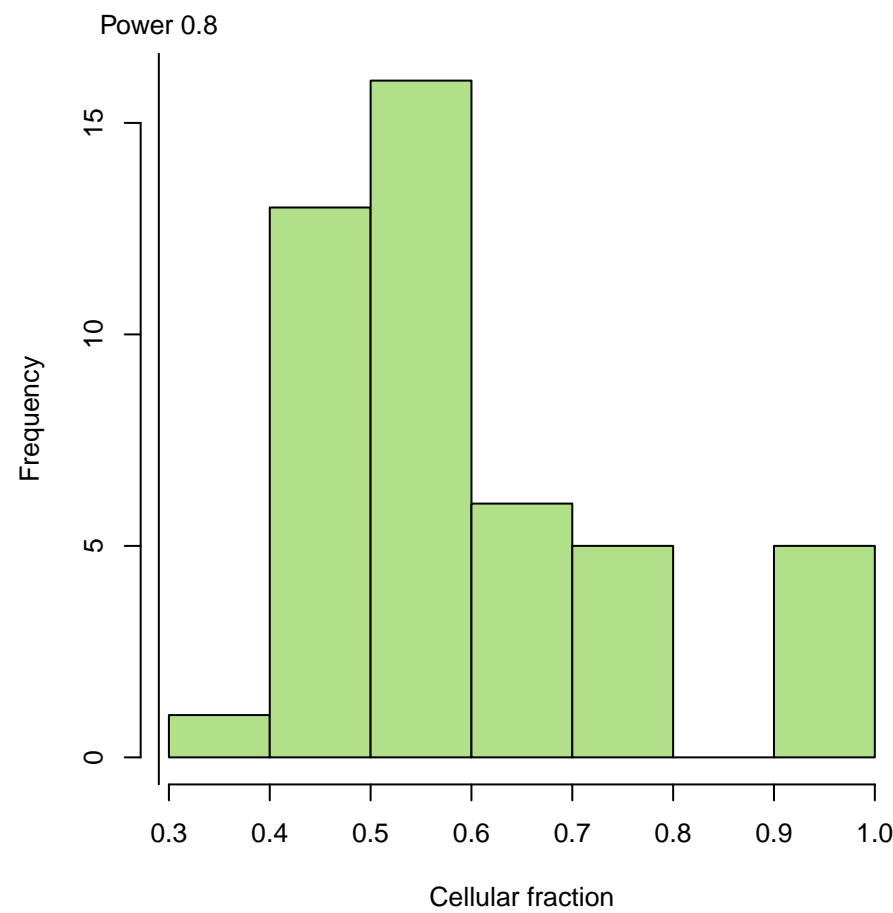
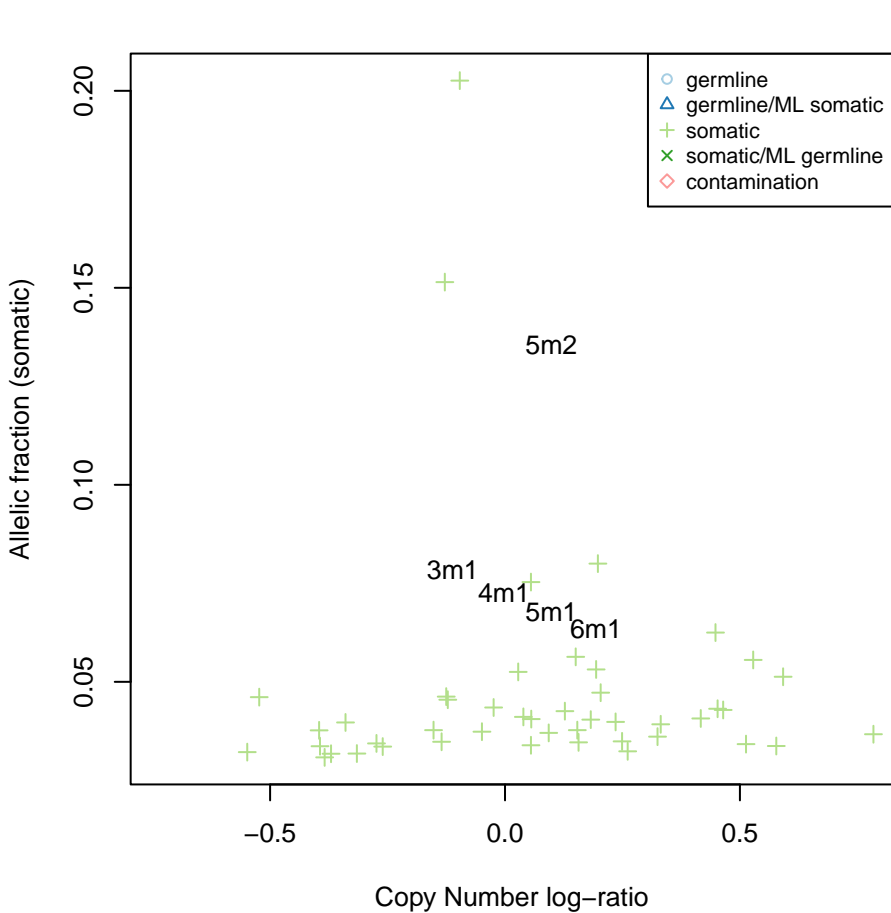
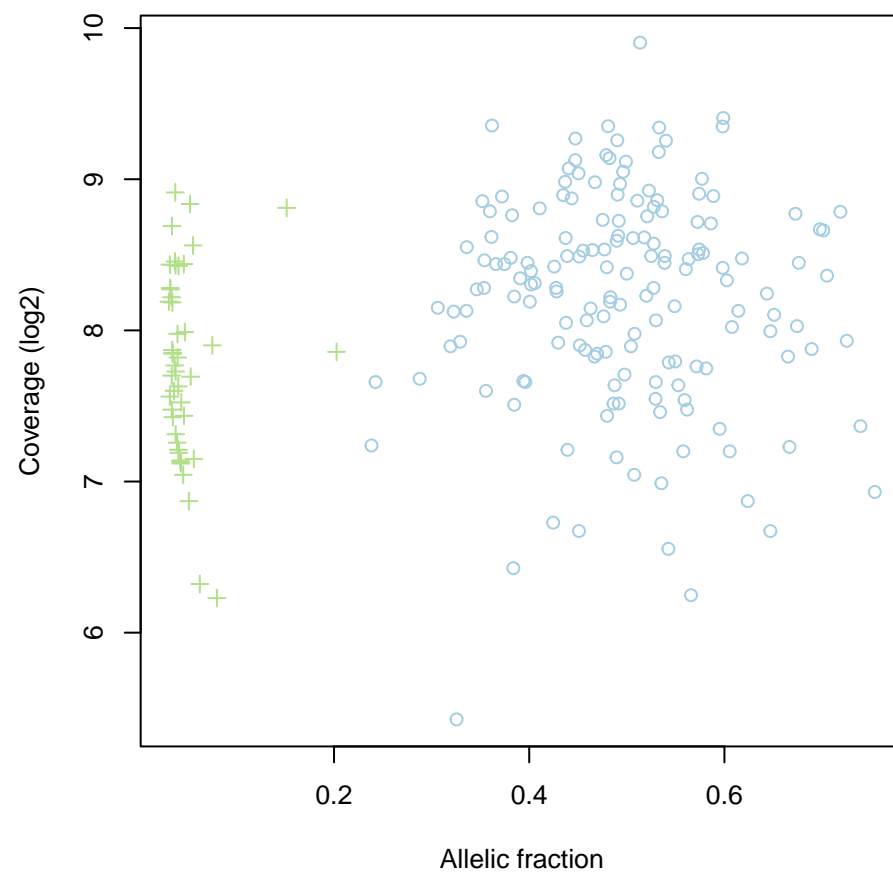
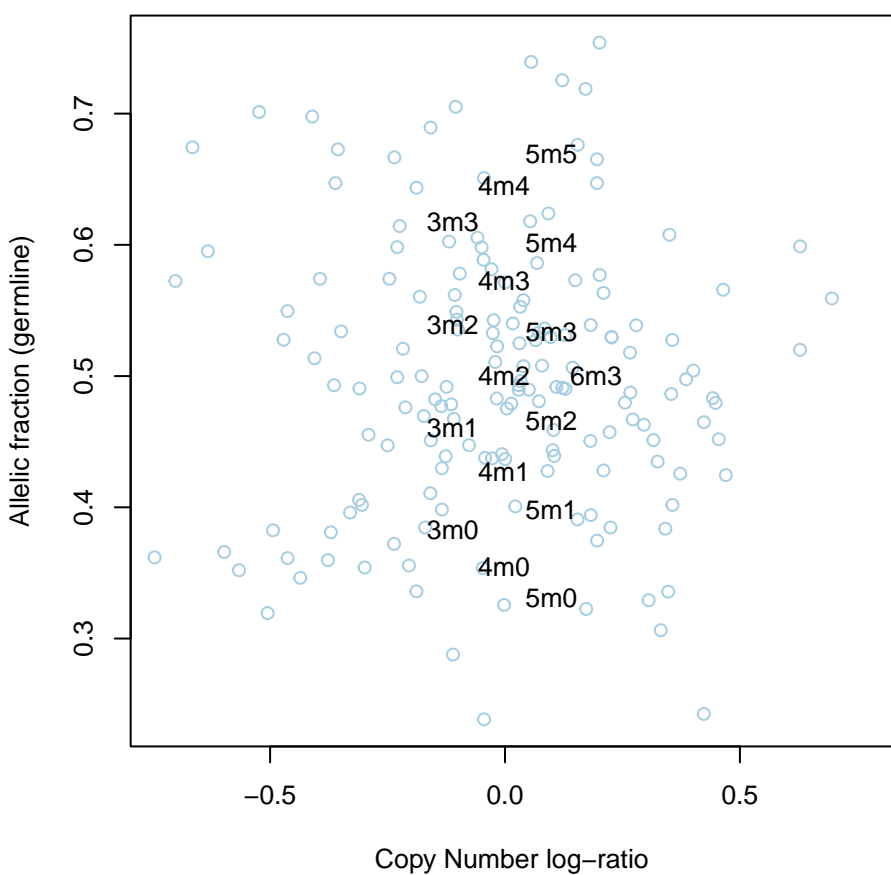


Purity: 0.17 Tumor ploidy: 4.021 SNV log-likelihood: -185.05 GoF: 94.7% Mean coverage: 312;424

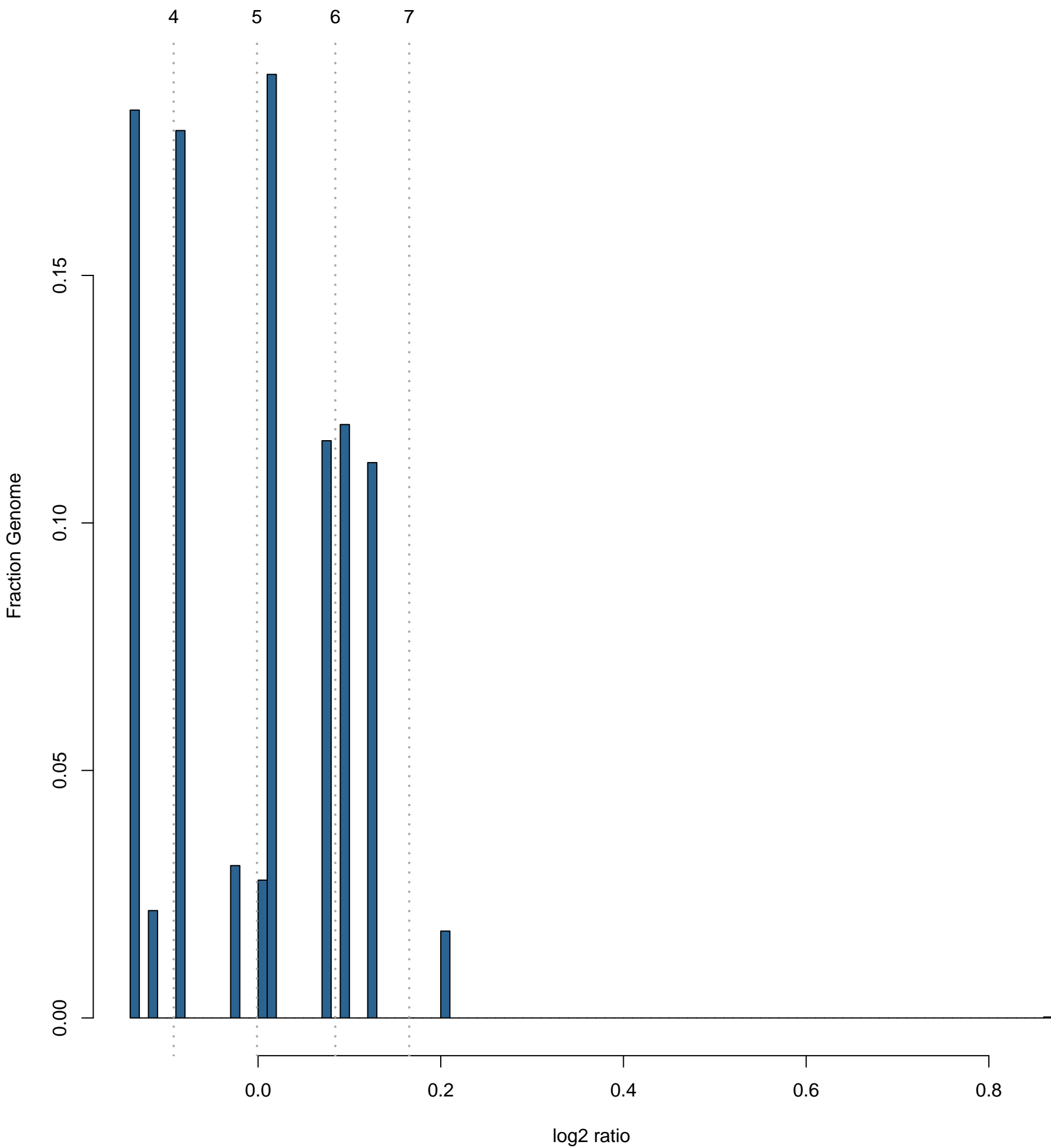


SCNA-fit log-likelihood: -19375.18

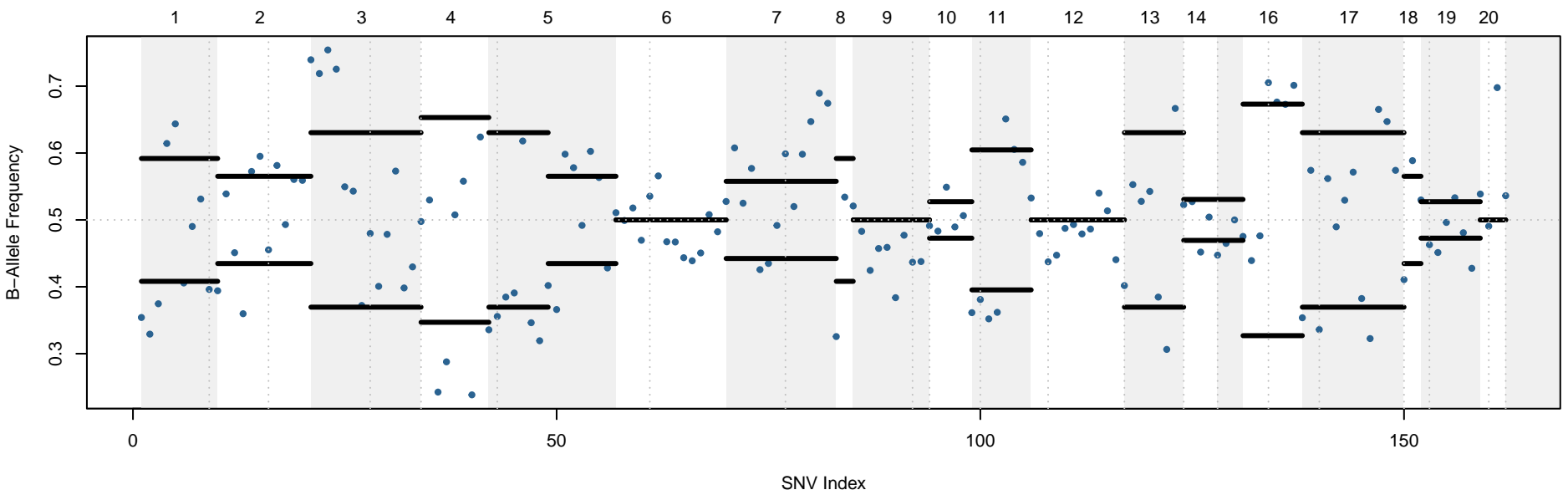




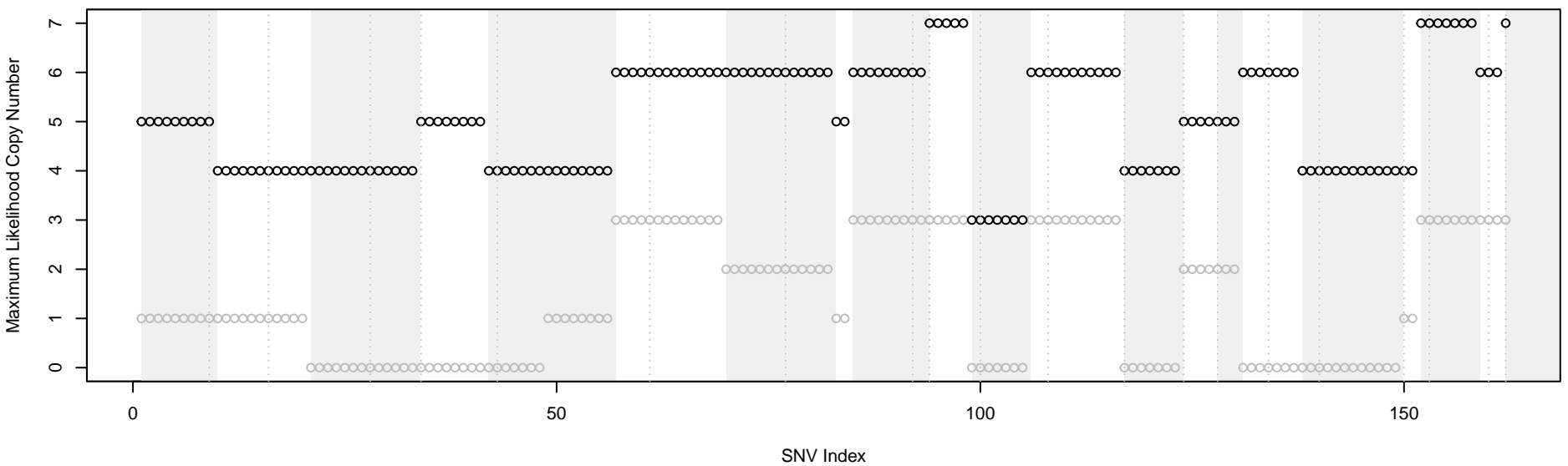
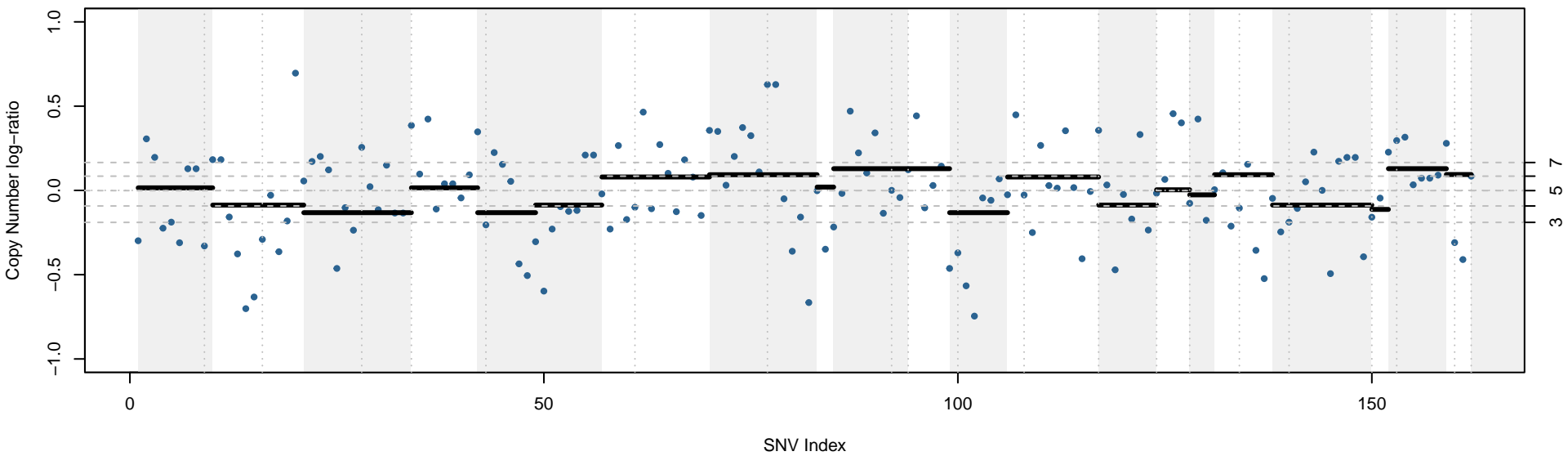
Purity: 0.15 Tumor ploidy: 5.014



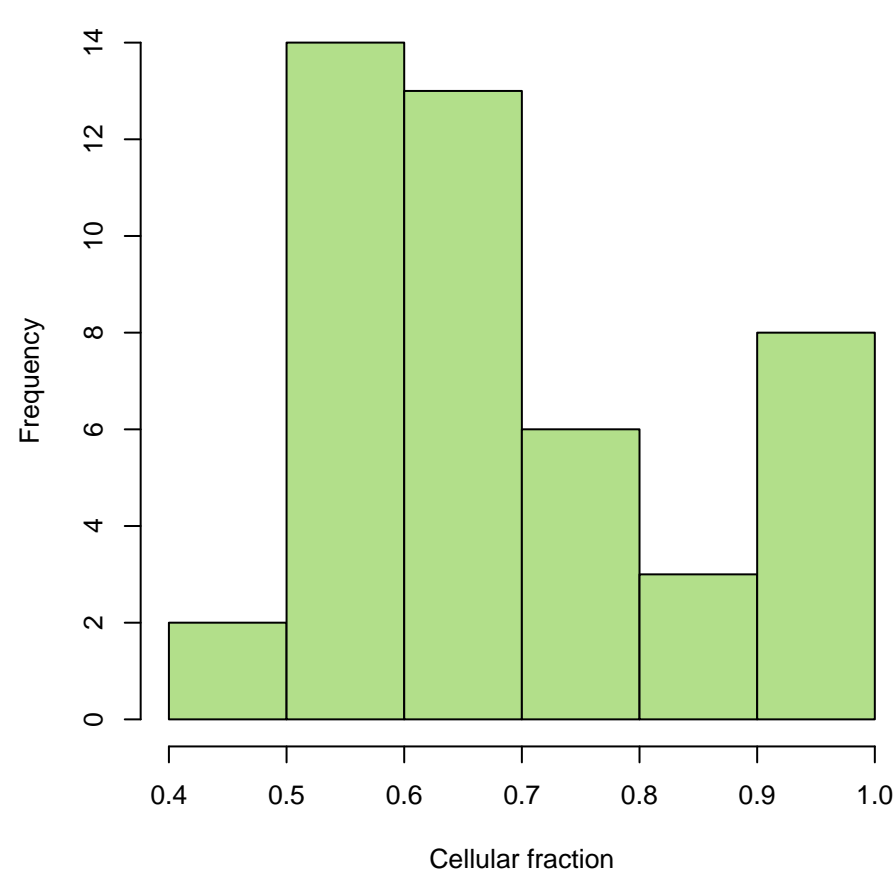
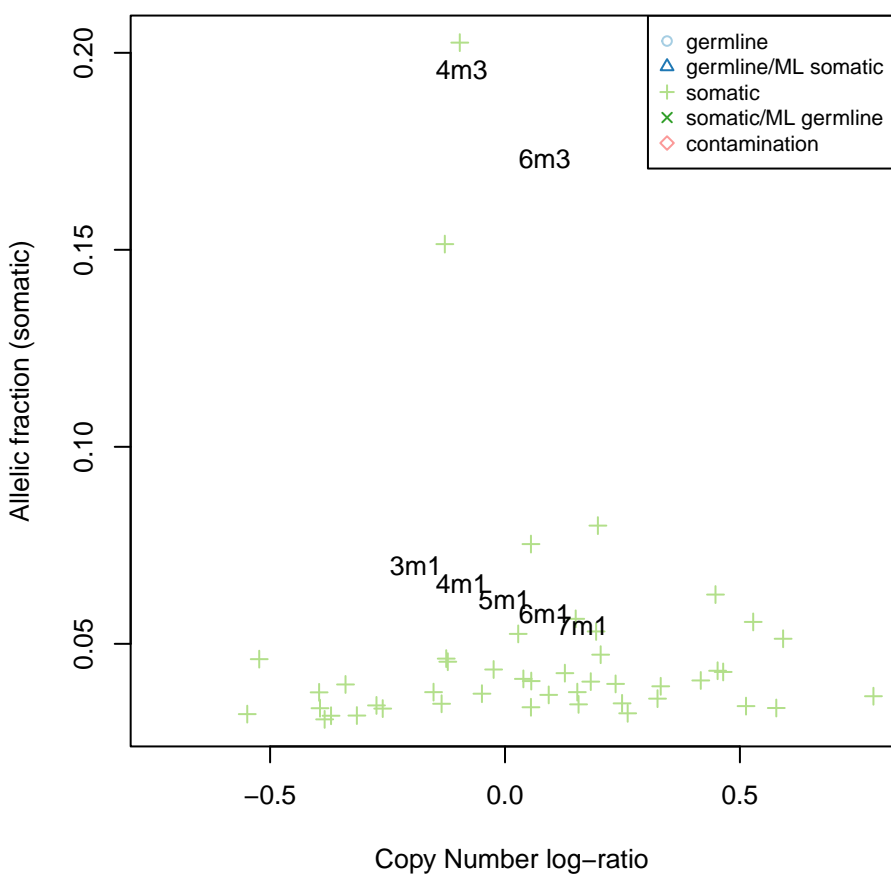
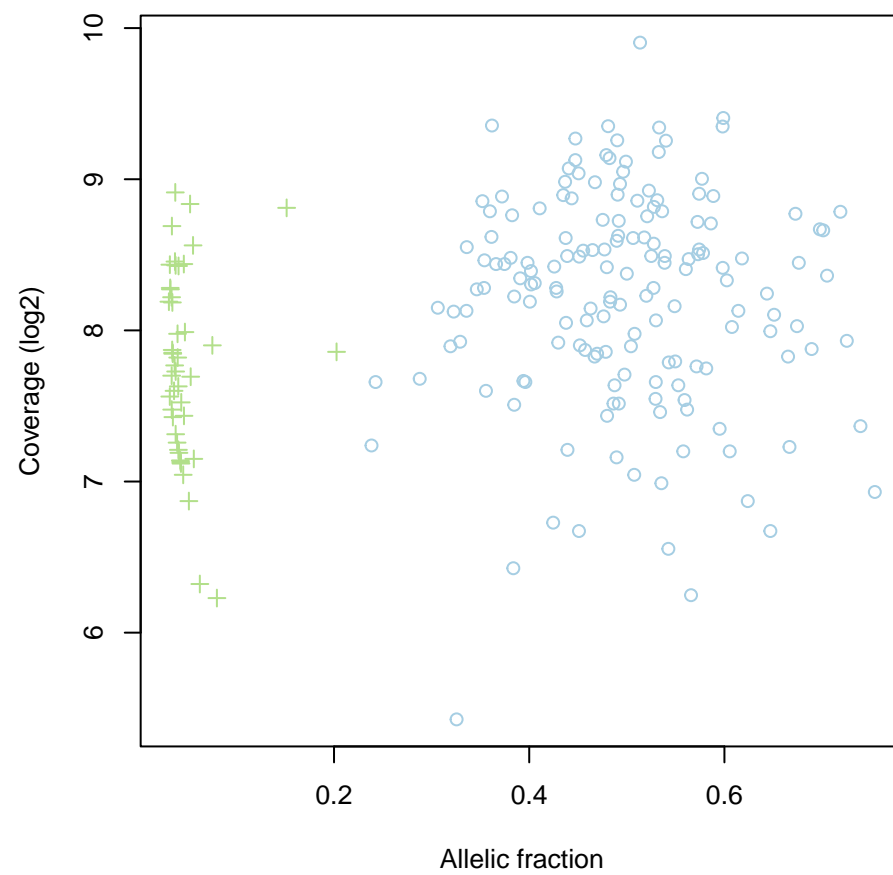
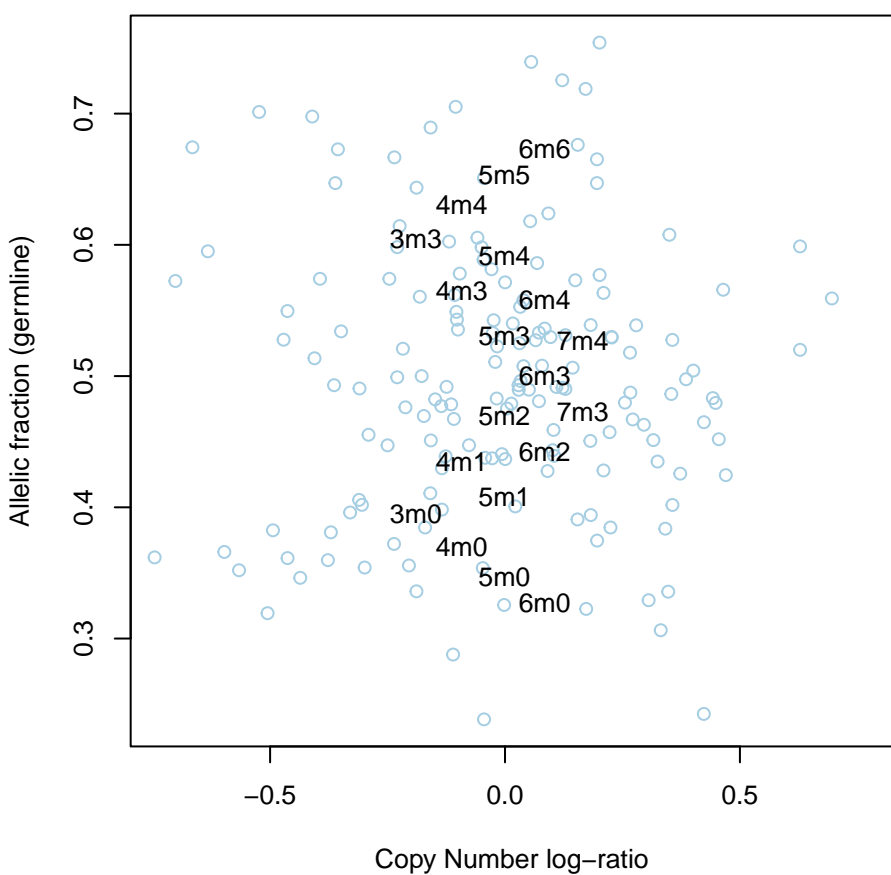
Purity: 0.15 Tumor ploidy: 5.014 SNV log-likelihood: -203.27 GoF: 94.8% Mean coverage: 312;424



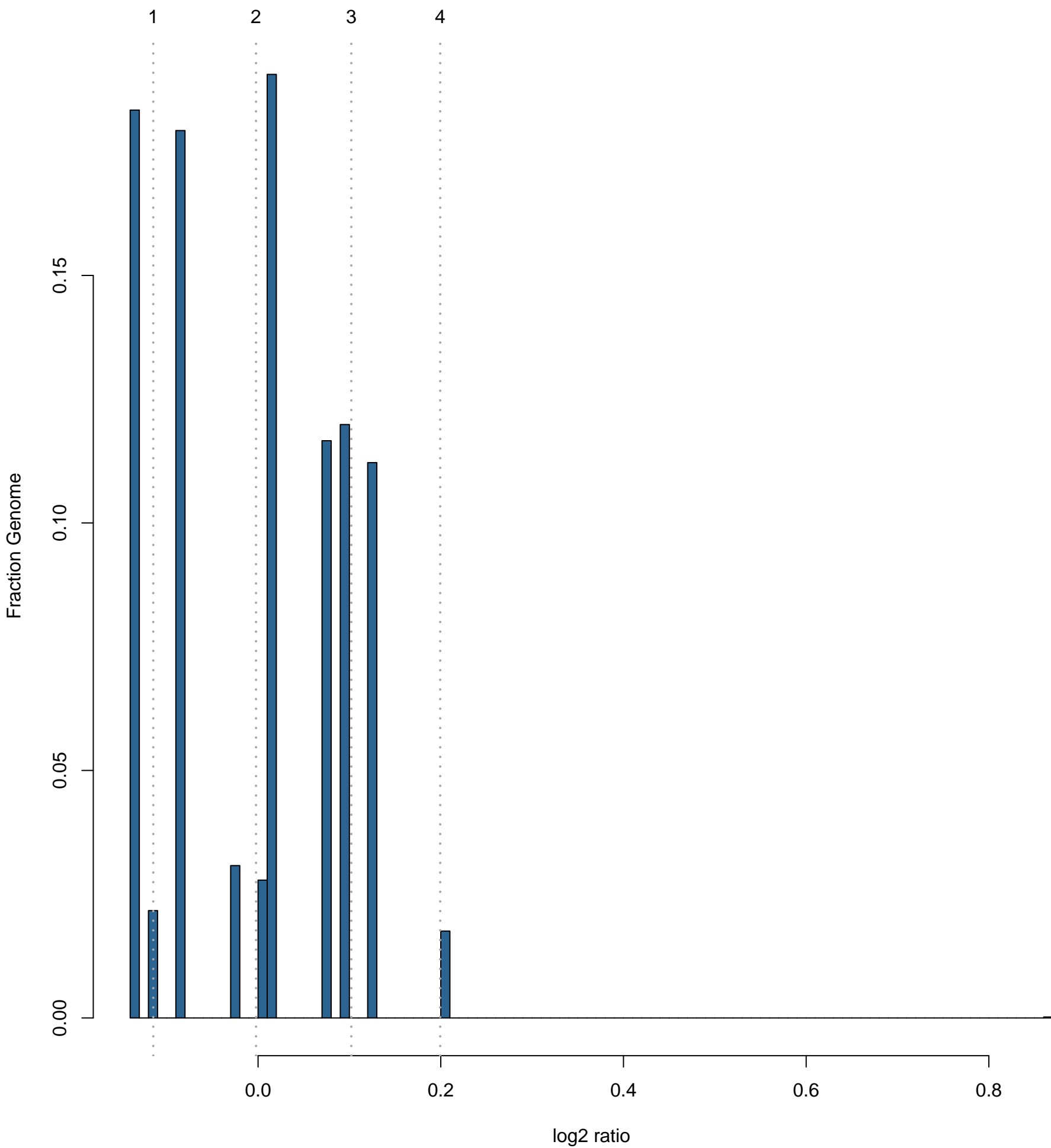
SCNA-fit log-likelihood: -19375.76



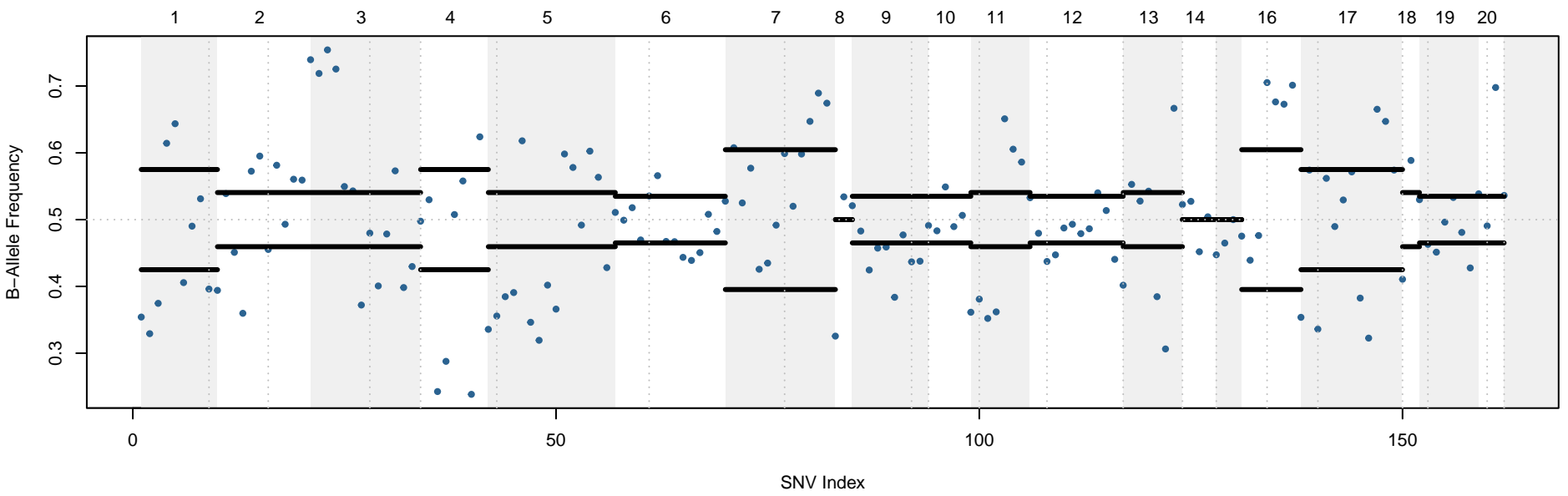




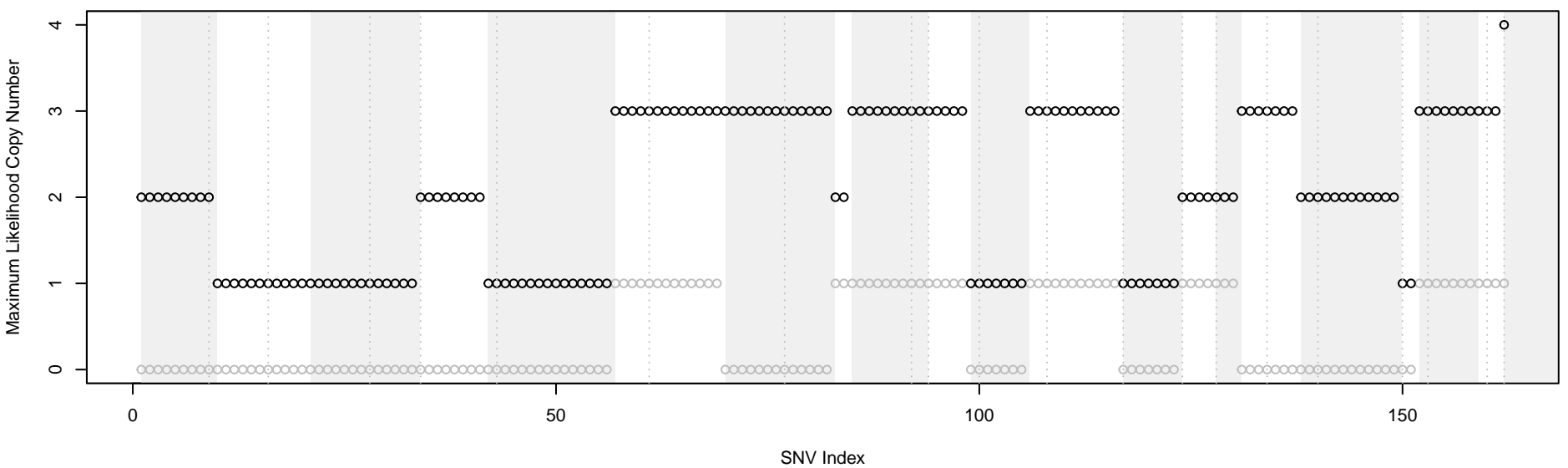
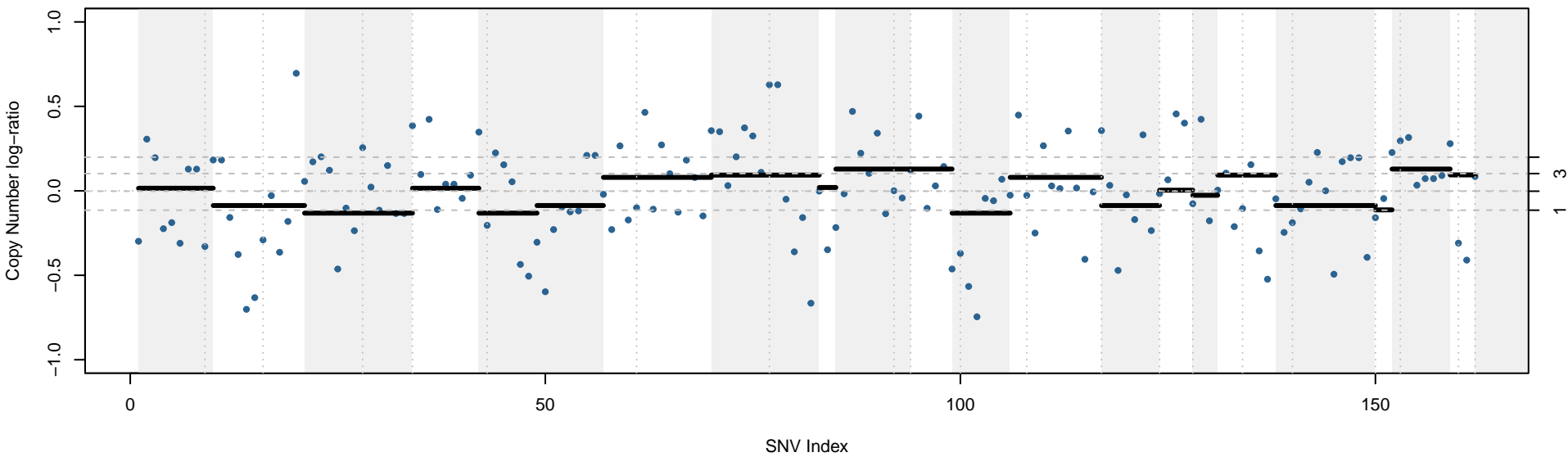
Purity: 0.15 Tumor ploidy: 2.021

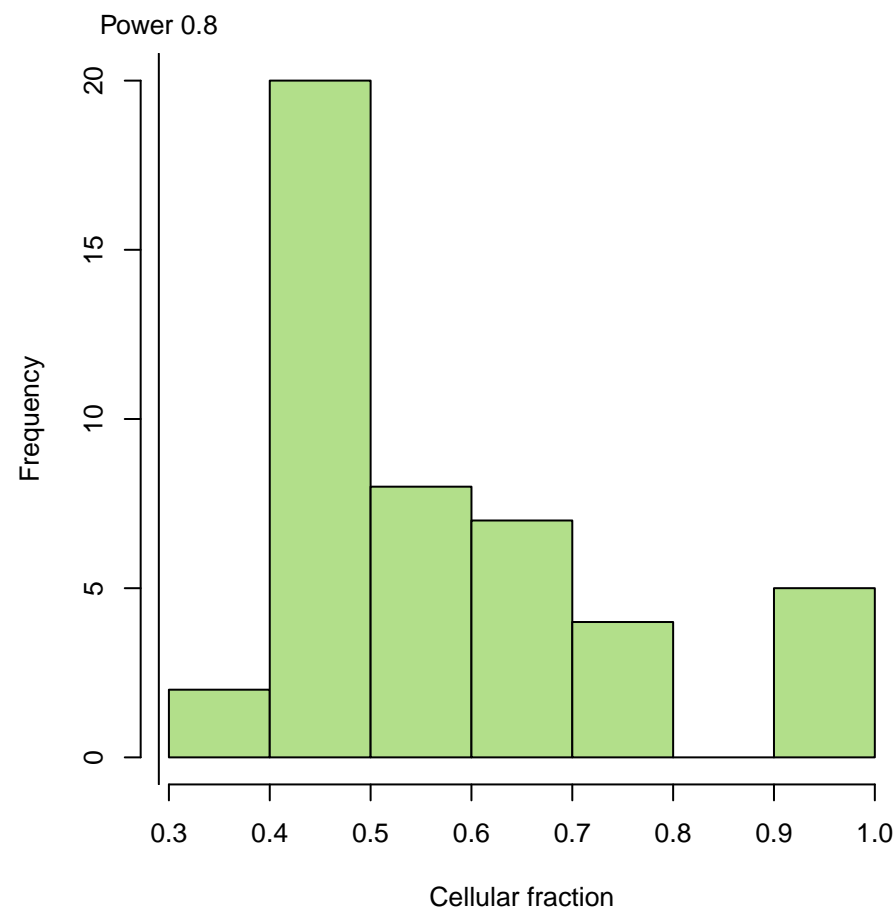
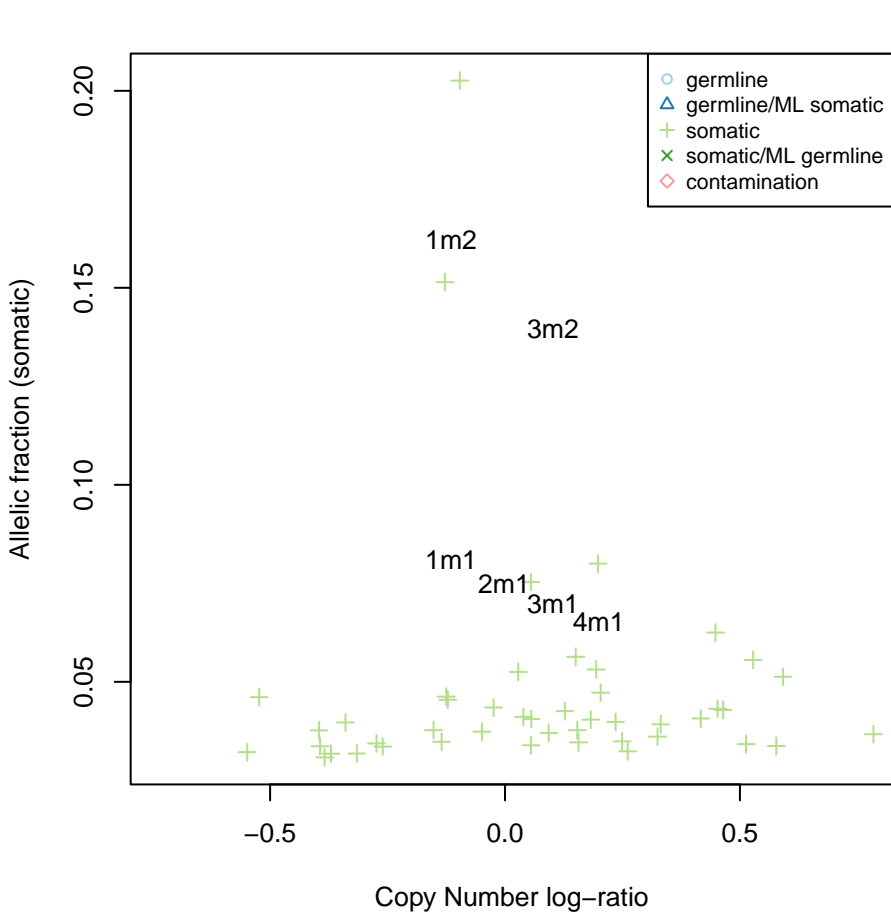
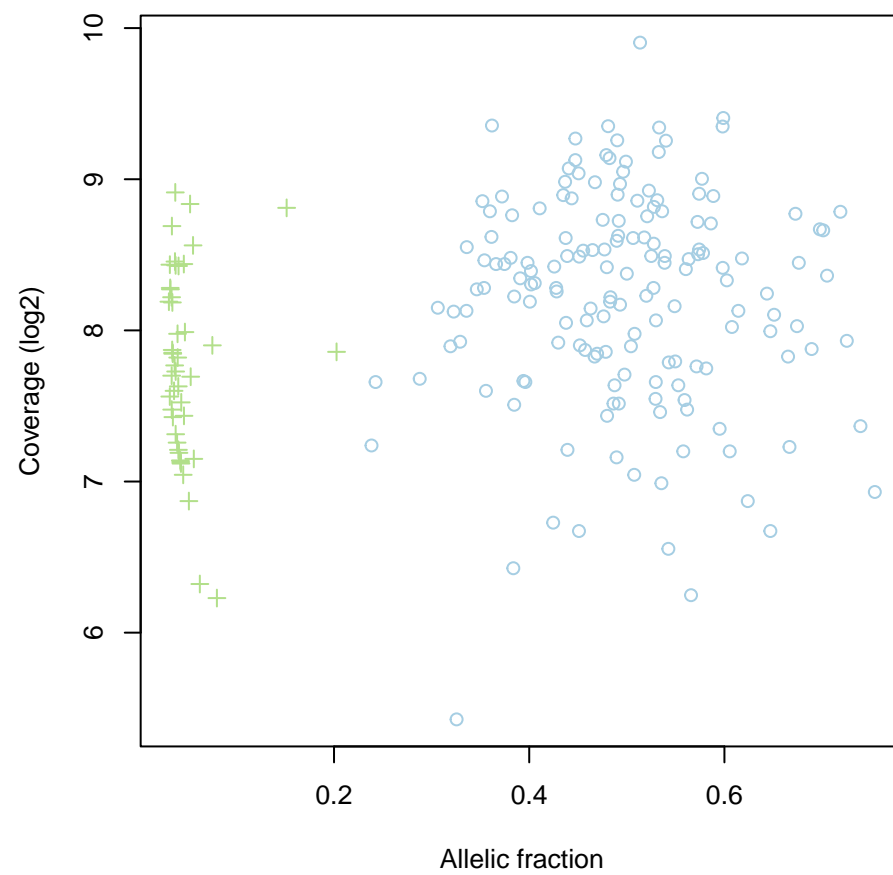
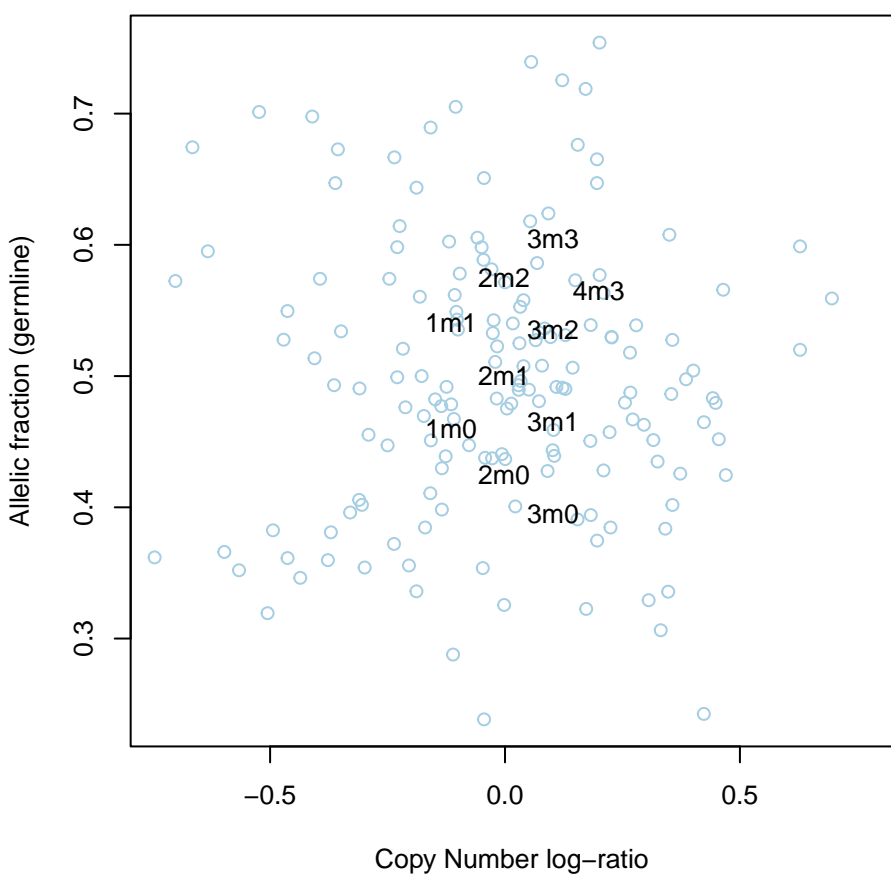


Purity: 0.15 Tumor ploidy: 2.021 SNV log-likelihood: -228.3 GoF: 90.1% Mean coverage: 312;424

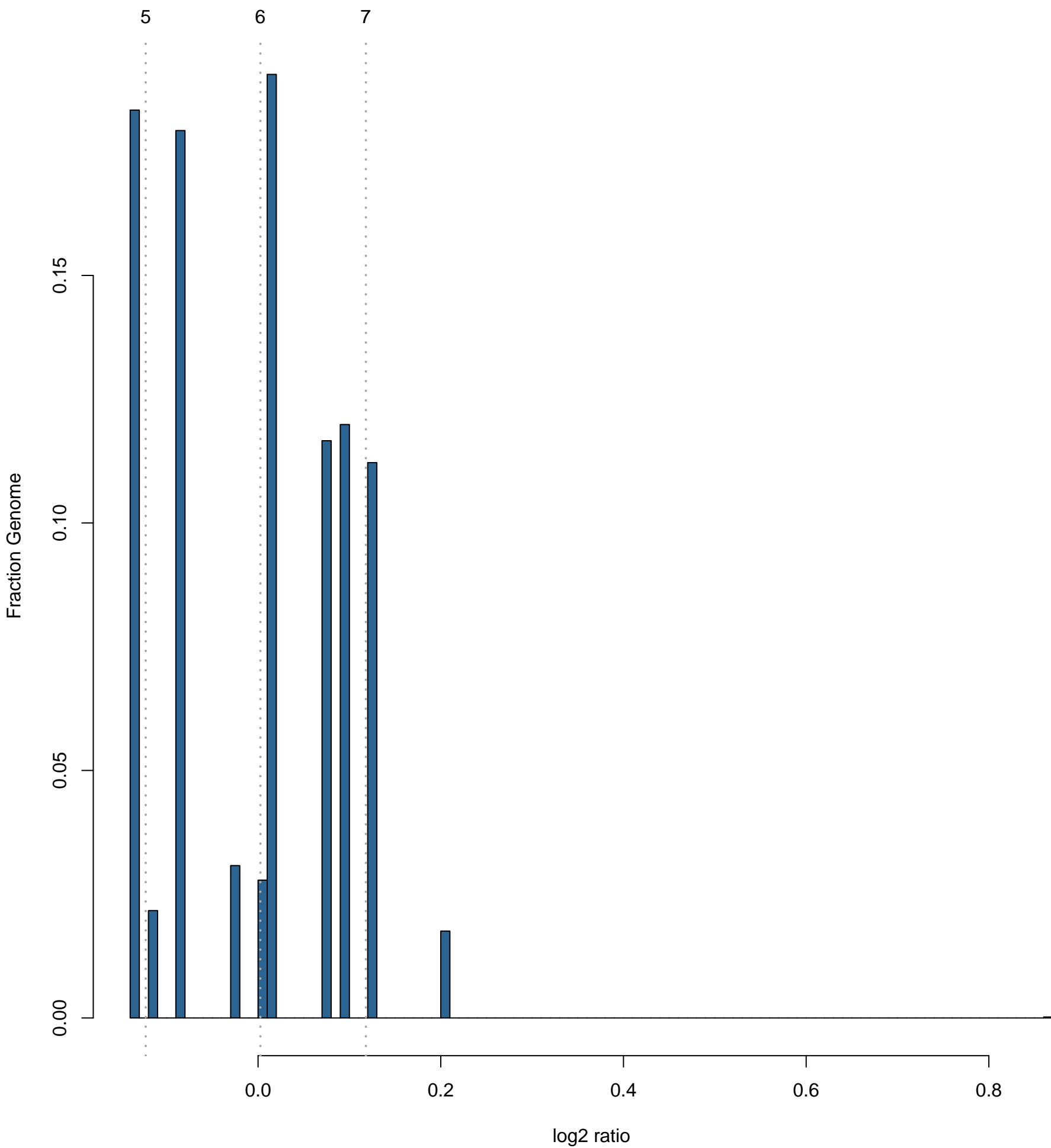


SCNA-fit log-likelihood: -19374.99

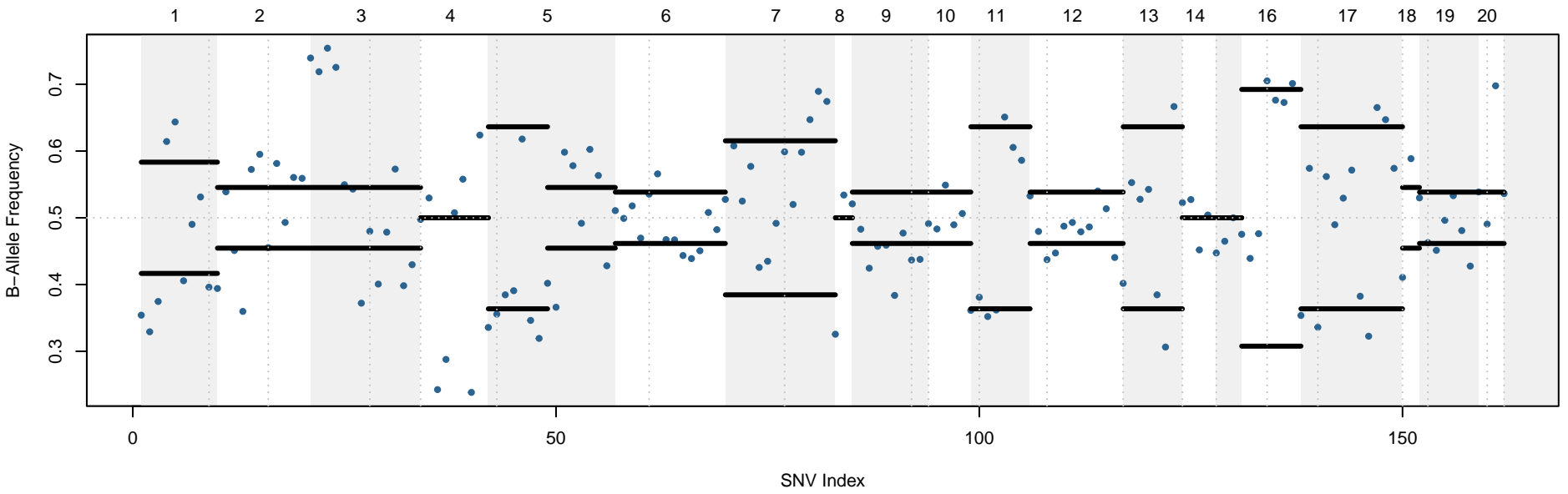




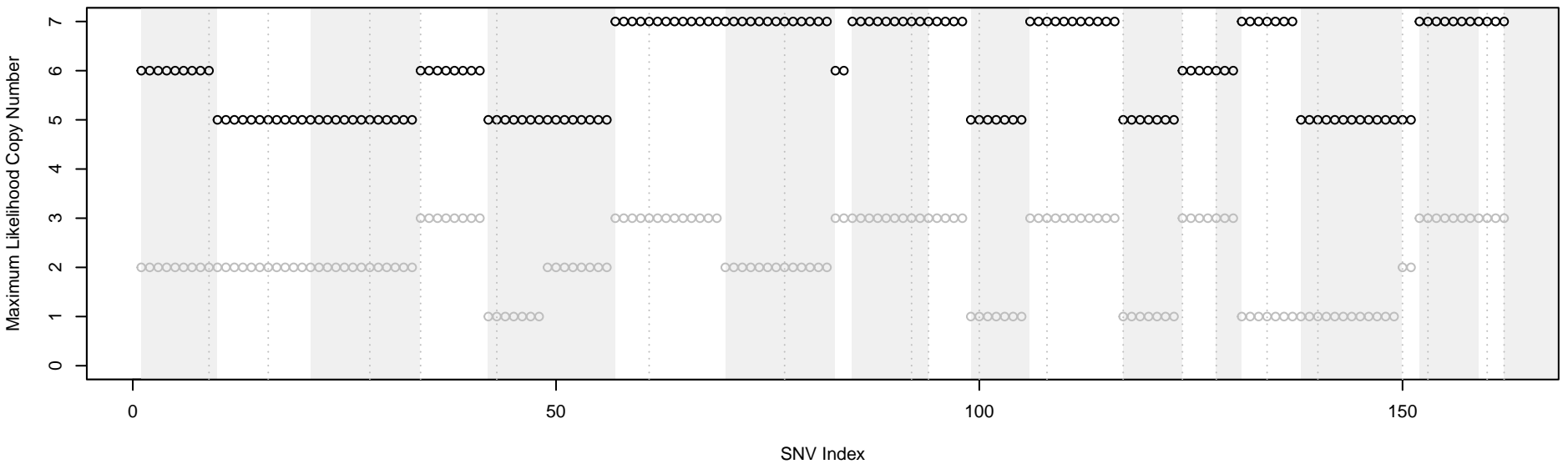
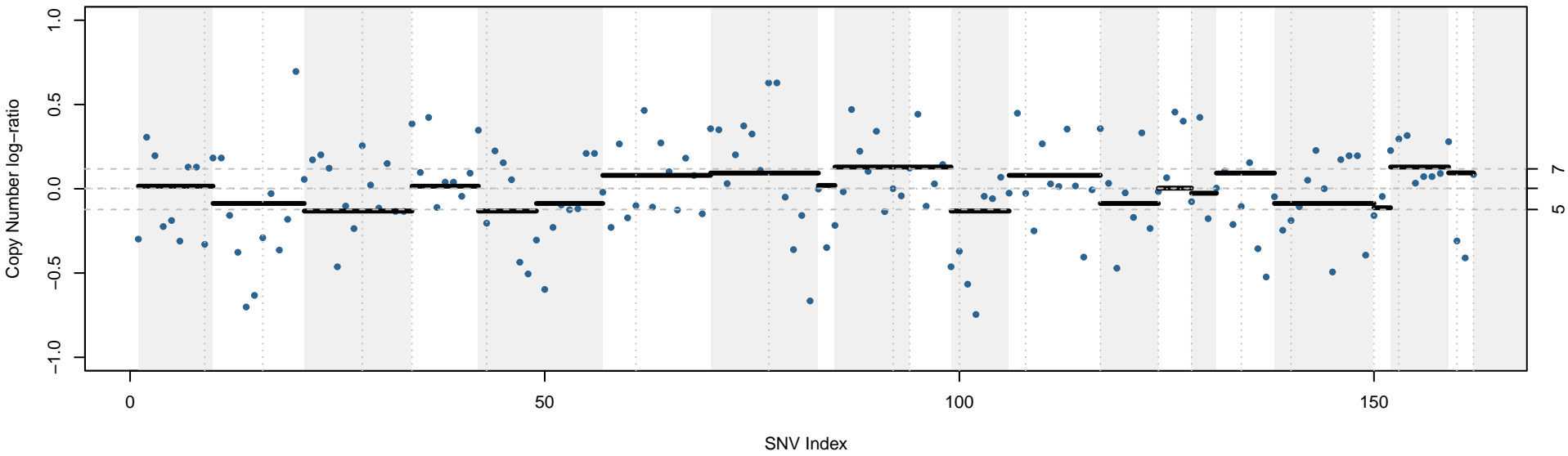
Purity: 0.25 Tumor ploidy: 5.979

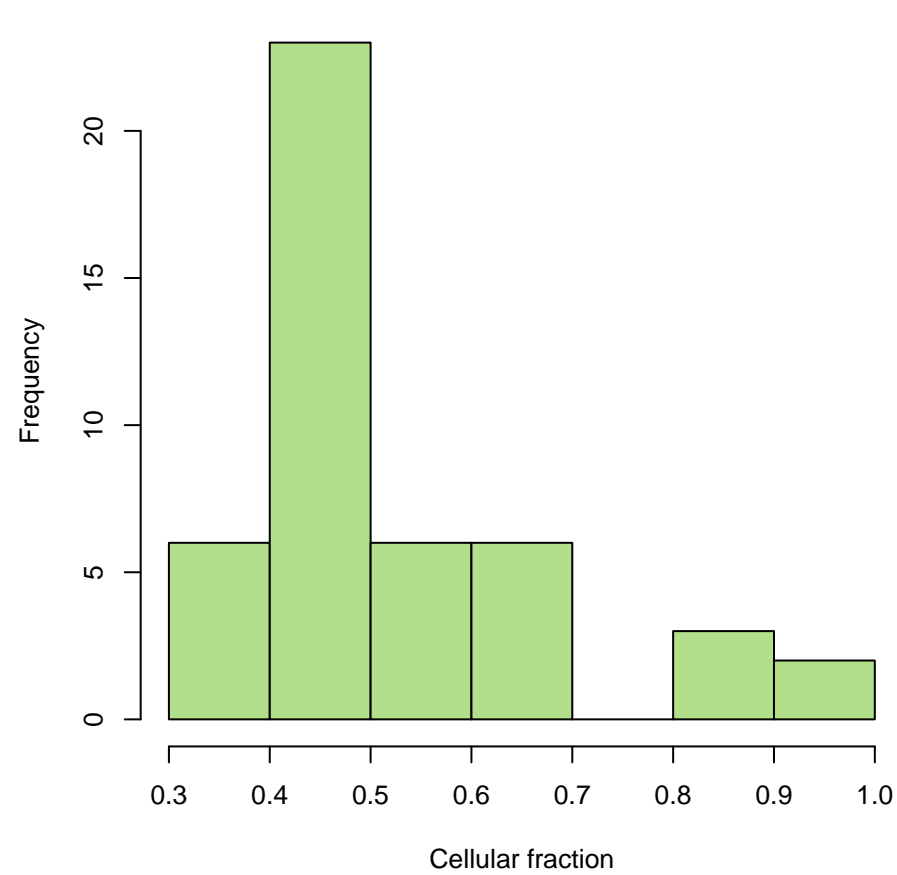
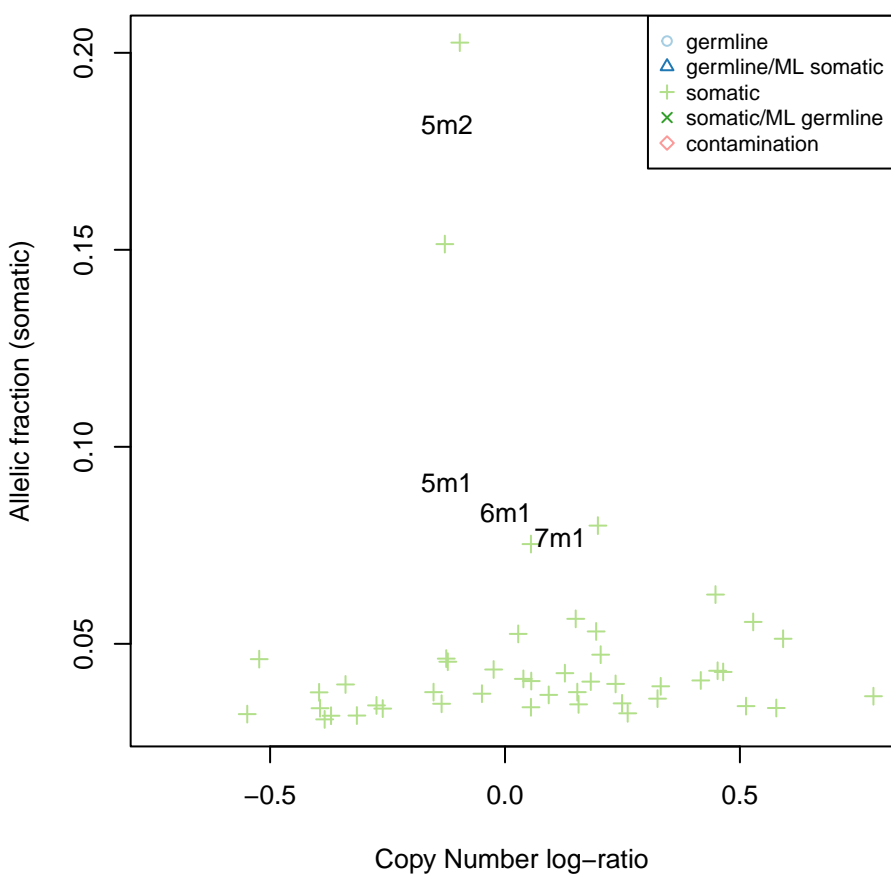
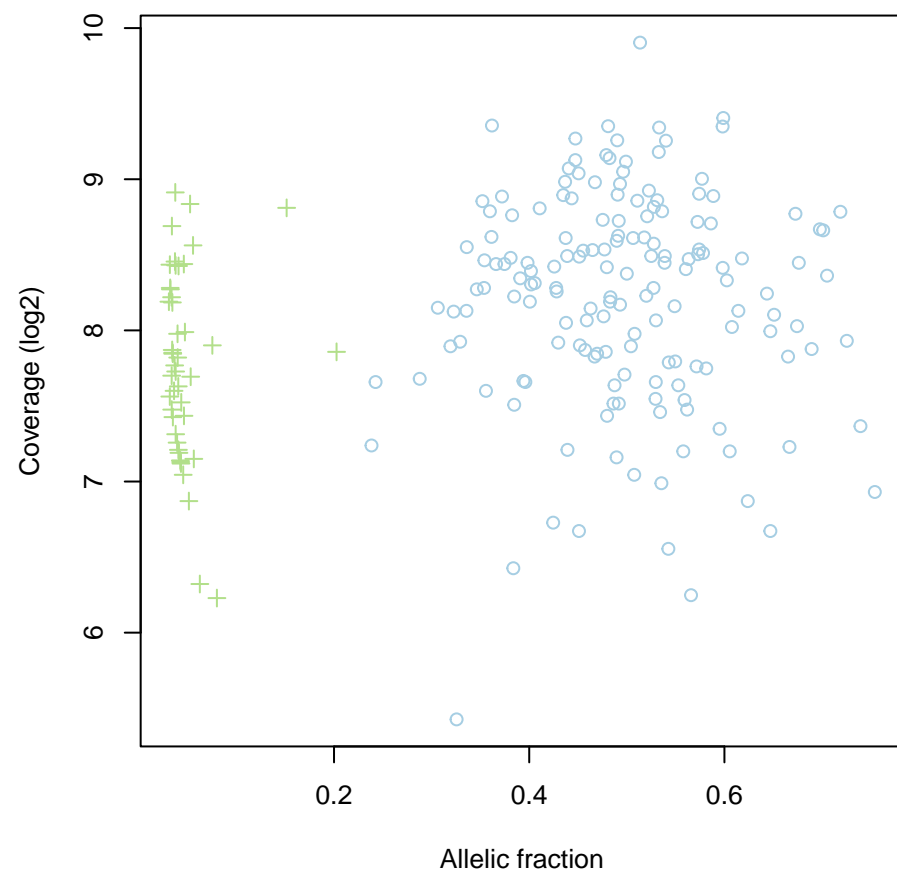
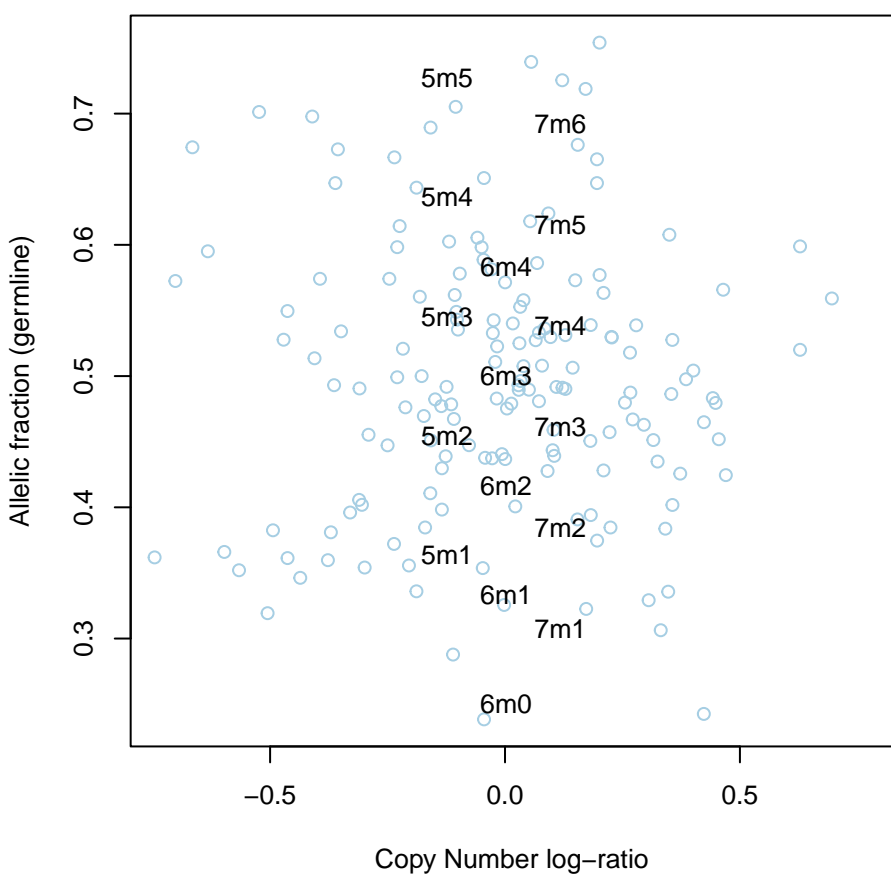


Purity: 0.25 Tumor ploidy: 5.979 SNV log-likelihood: -266.85 GoF: 95.1% Mean coverage: 312;424

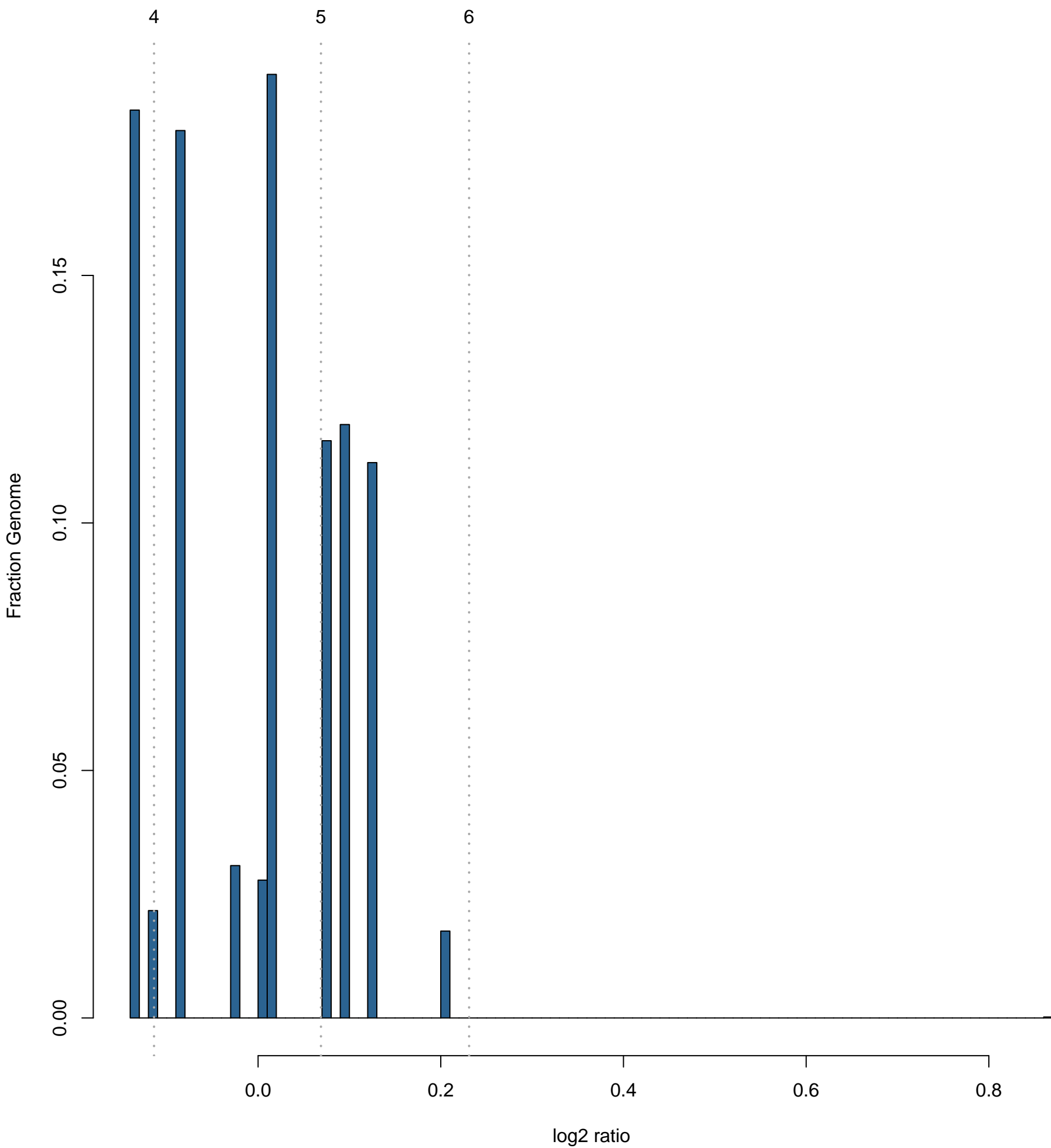


SCNA-fit log-likelihood: -19383.38



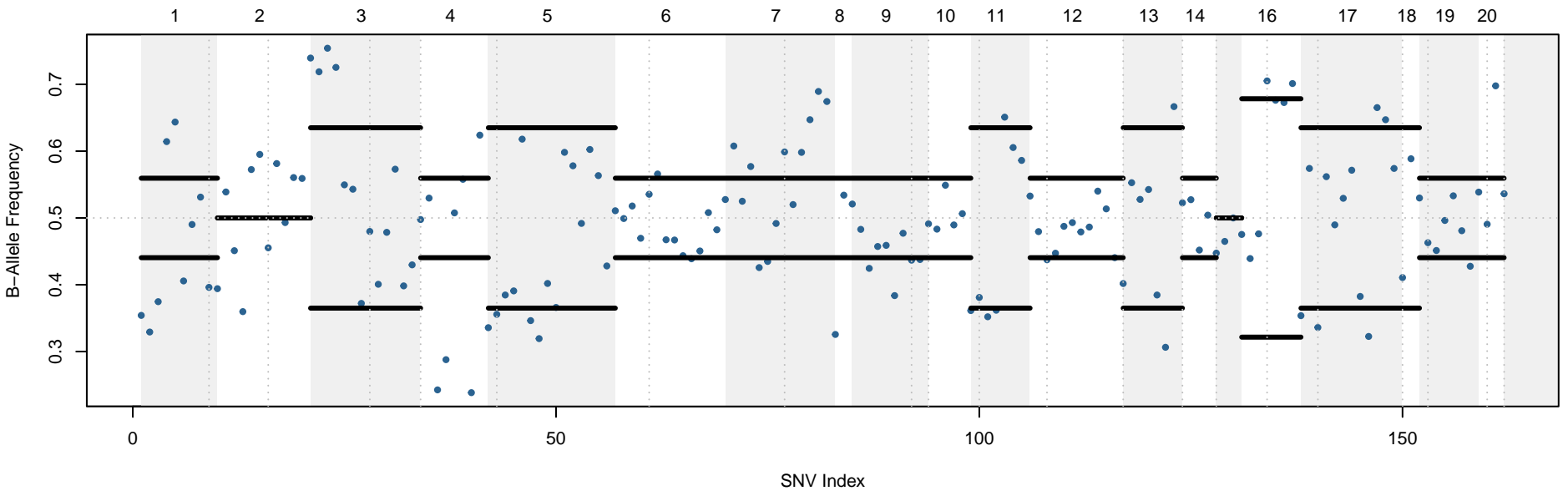


Purity: 0.37 Tumor ploidy: 4.608

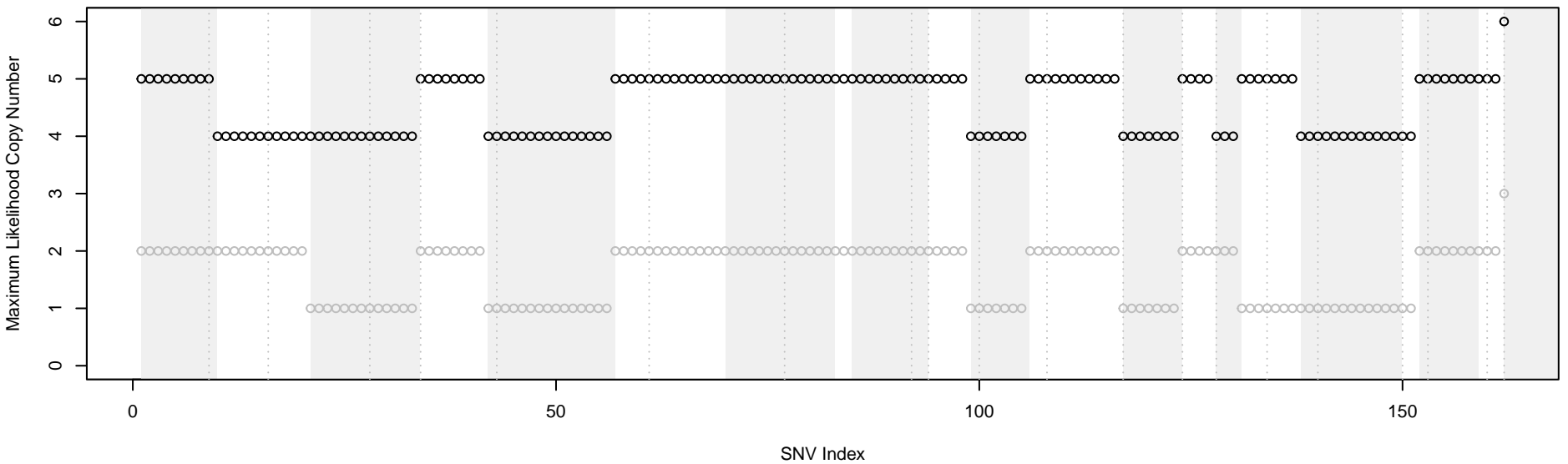
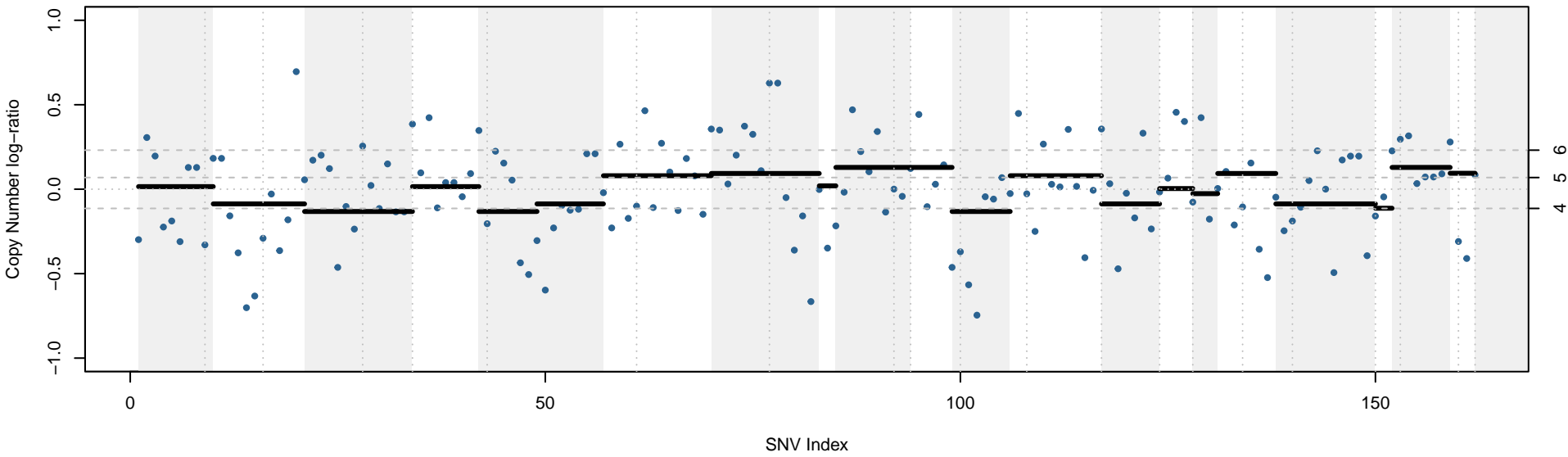


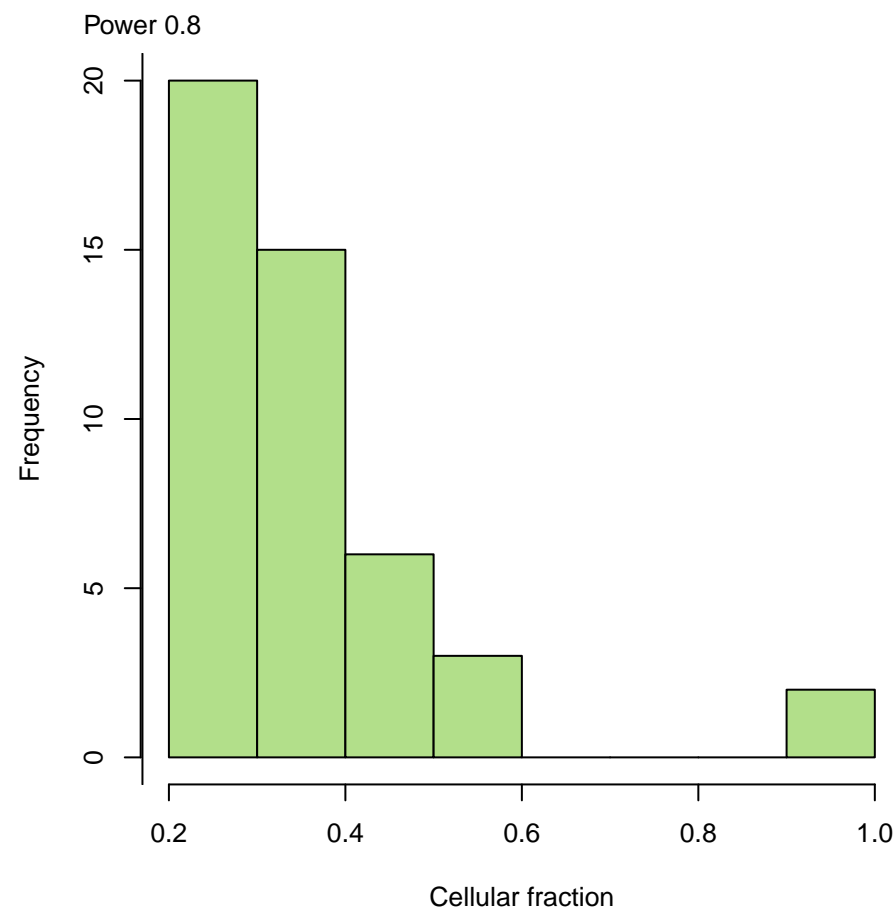
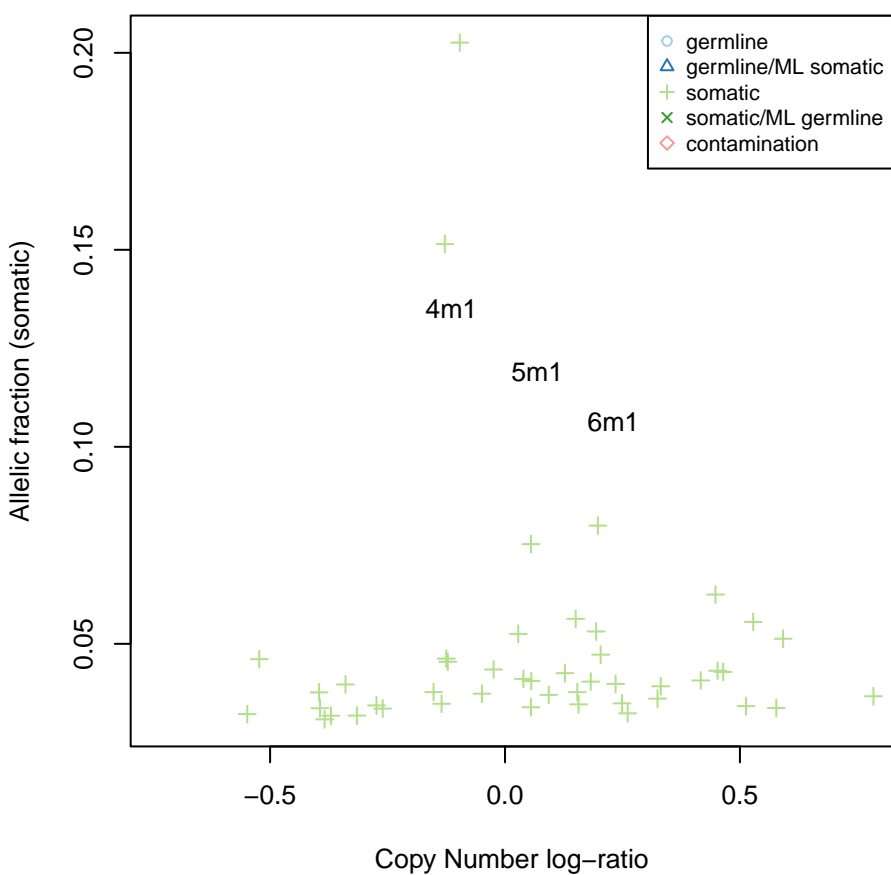
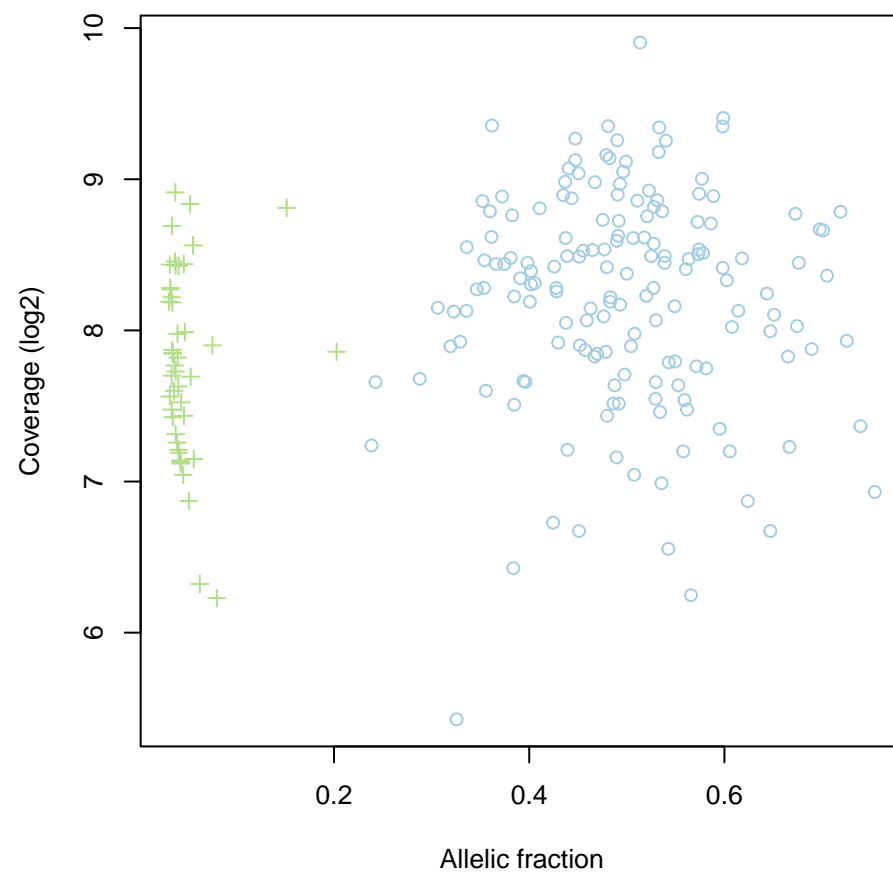
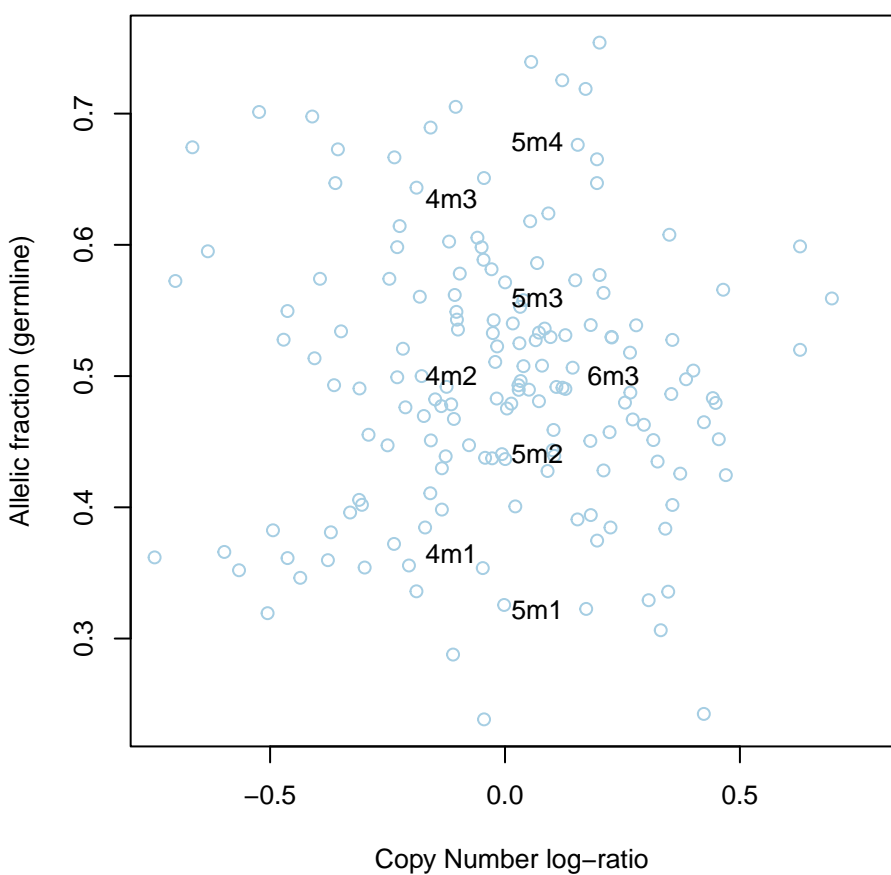


Purity: 0.37 Tumor ploidy: 4.608 SNV log-likelihood: -259.4 GoF: 91.1% Mean coverage: 312;424

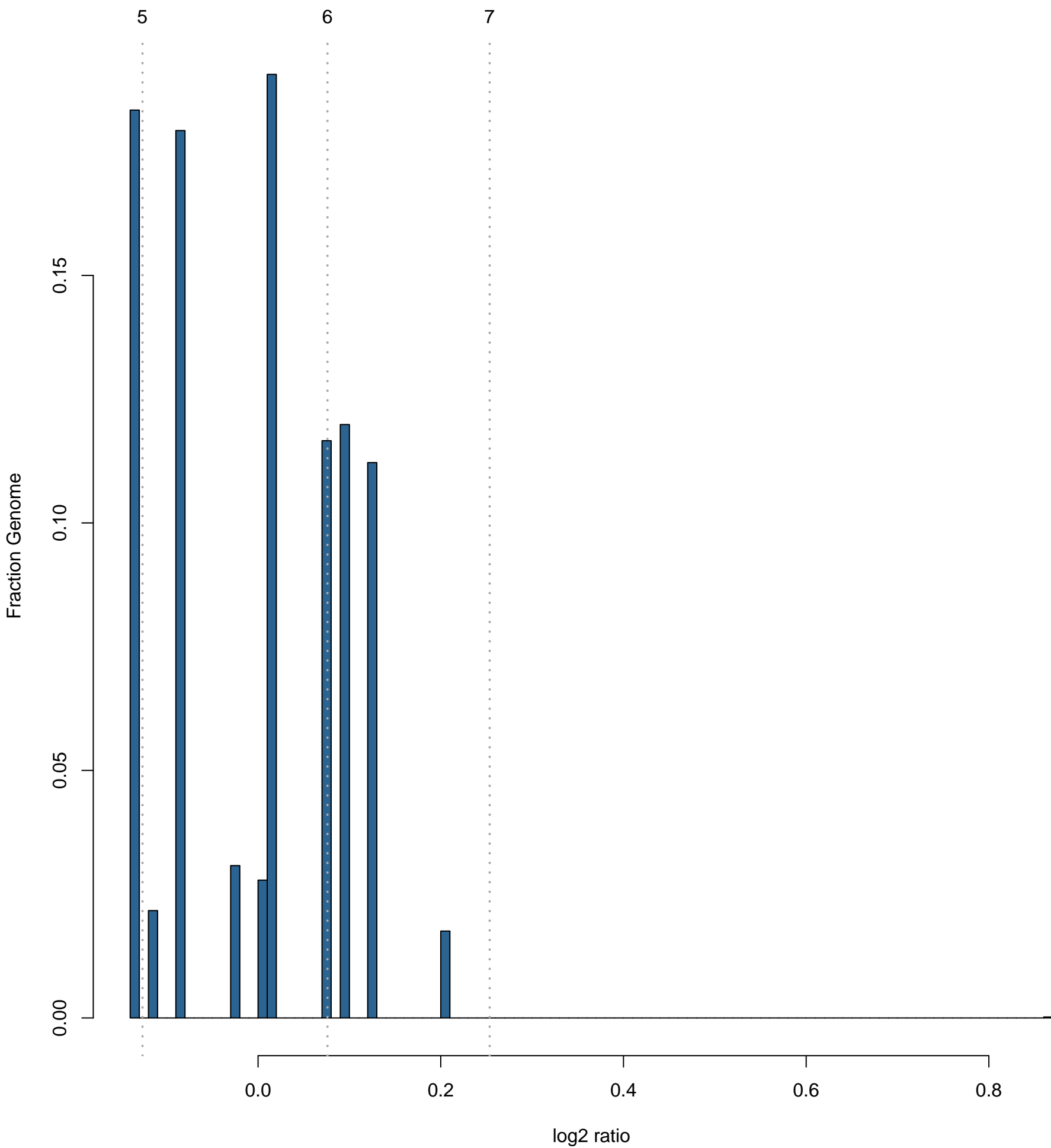


SCNA-fit log-likelihood: -19406.76

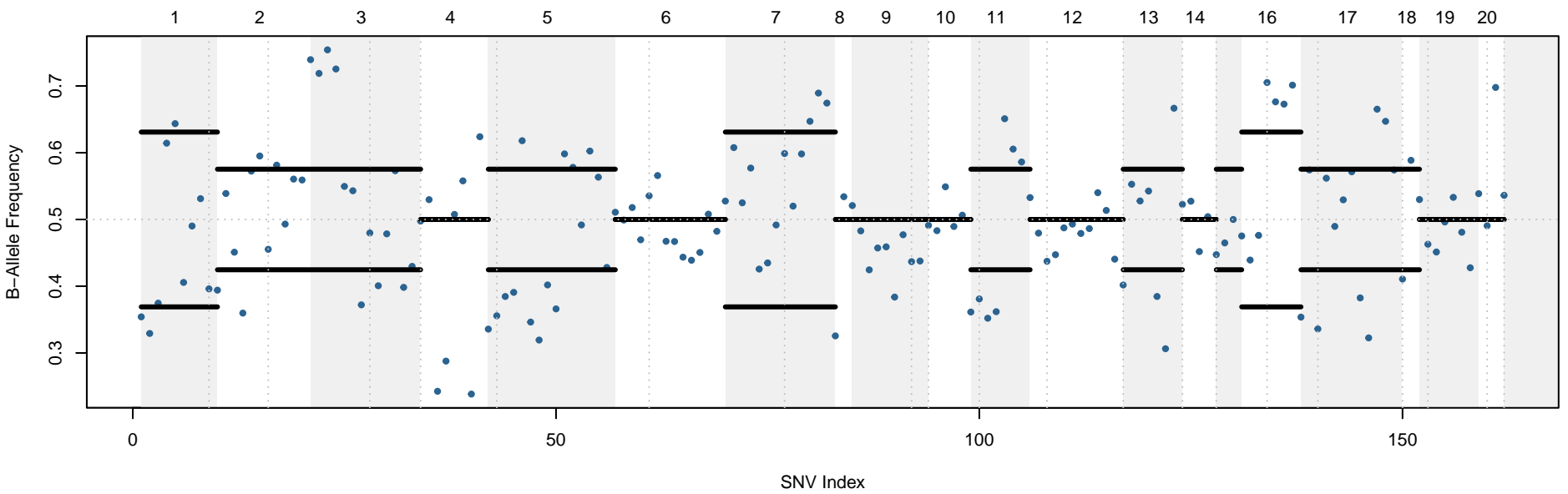




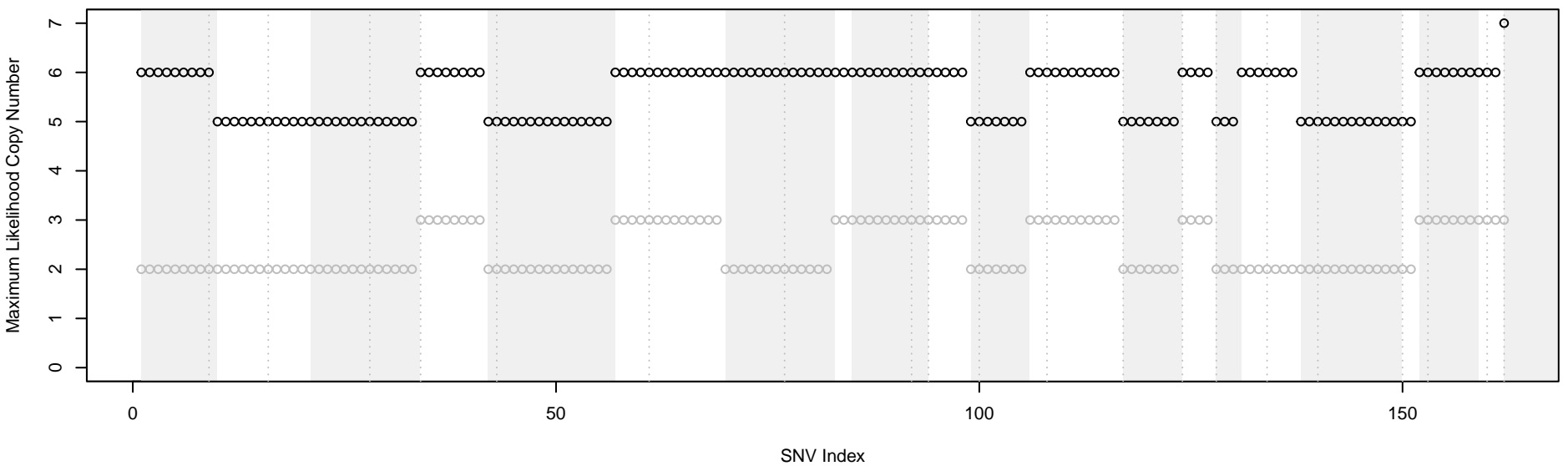
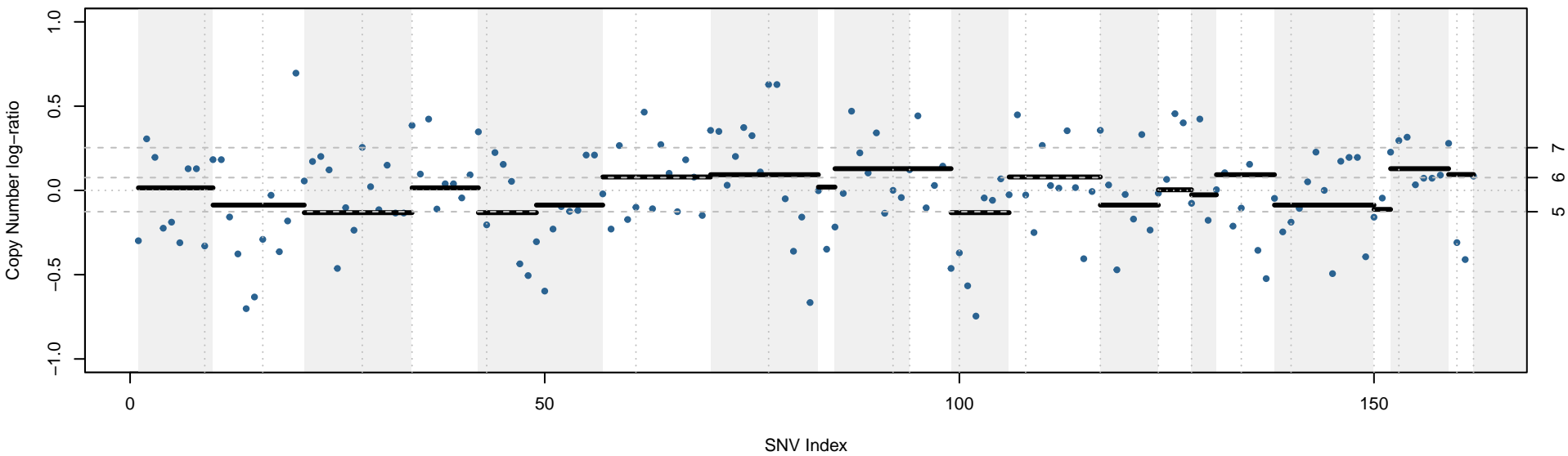
Purity: 0.55 Tumor ploidy: 5.608

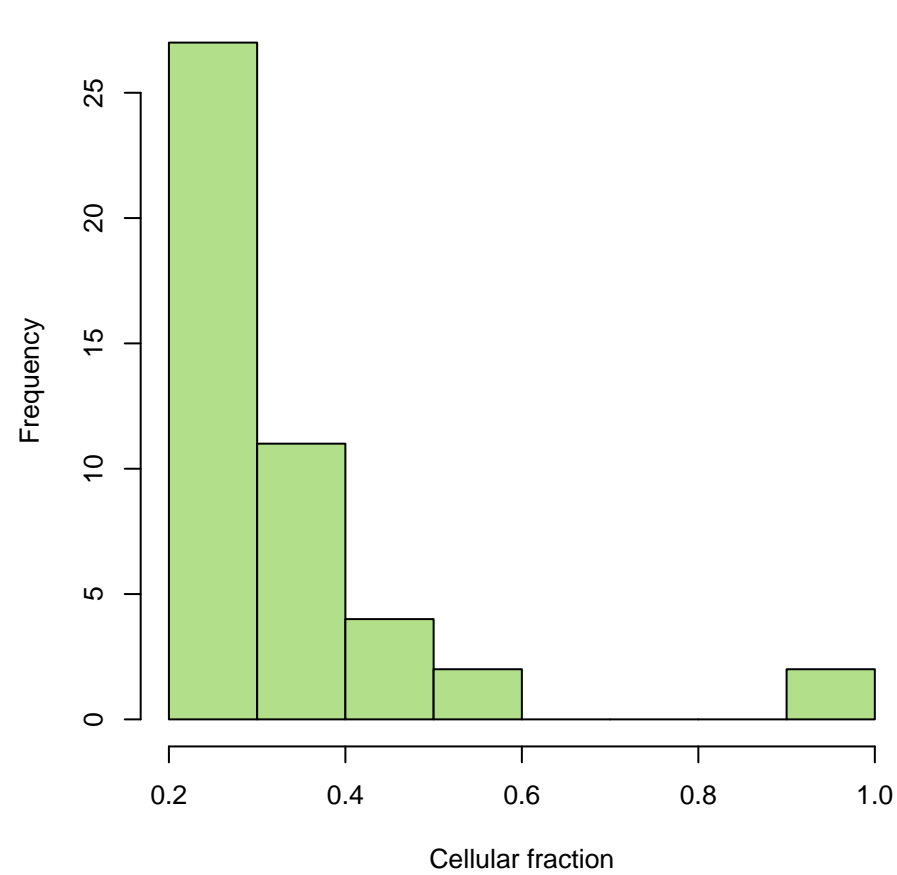
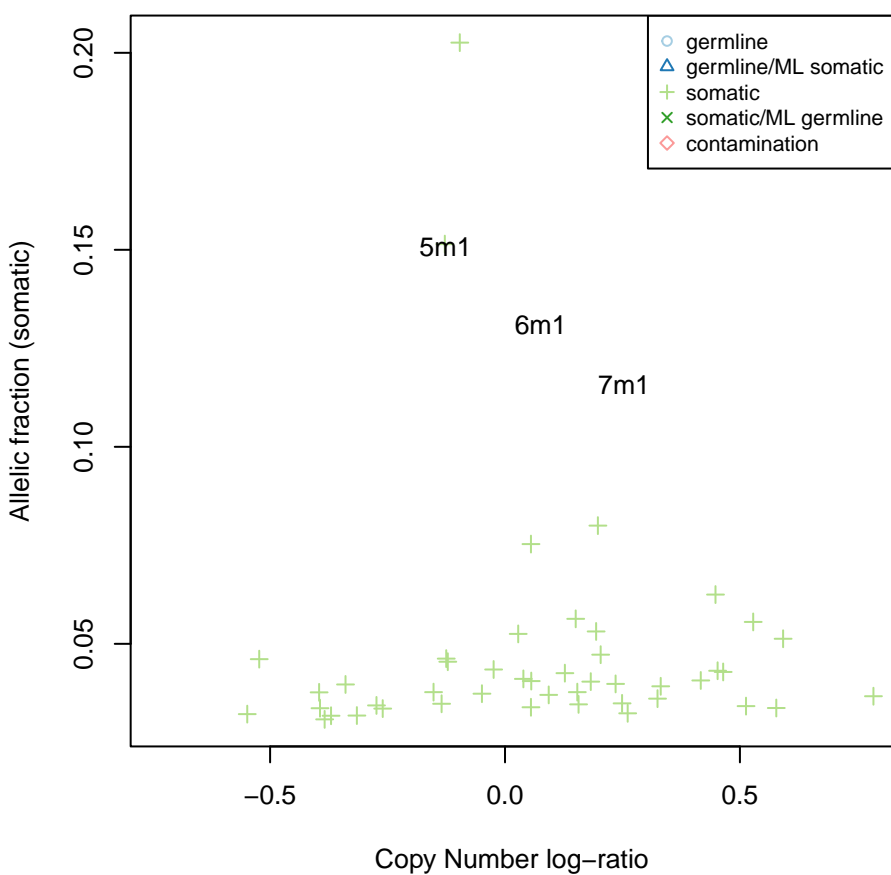
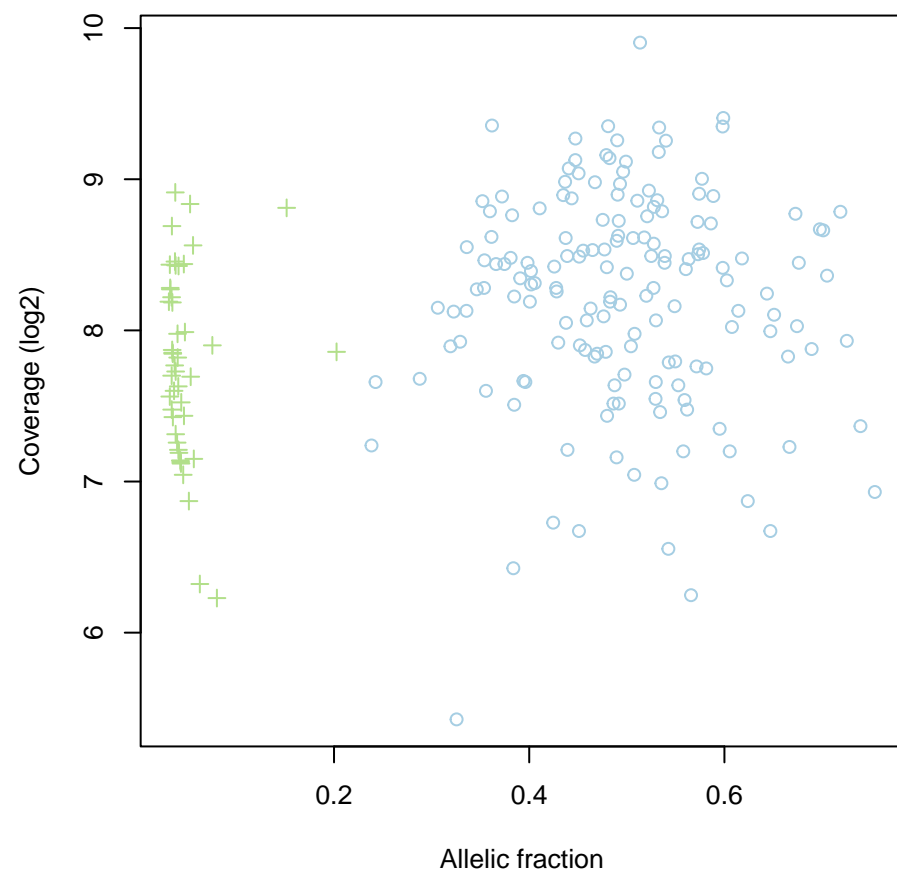
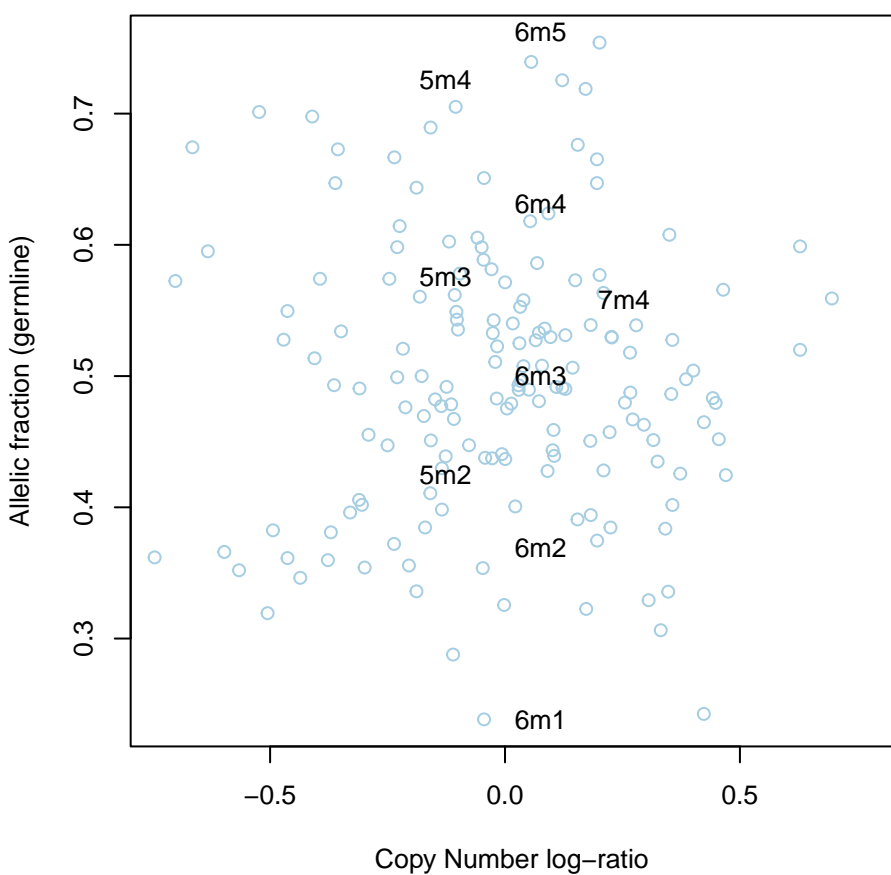


Purity: 0.55 Tumor ploidy: 5.608 SNV log-likelihood: -261.06 GoF: 90.1% Mean coverage: 312;424

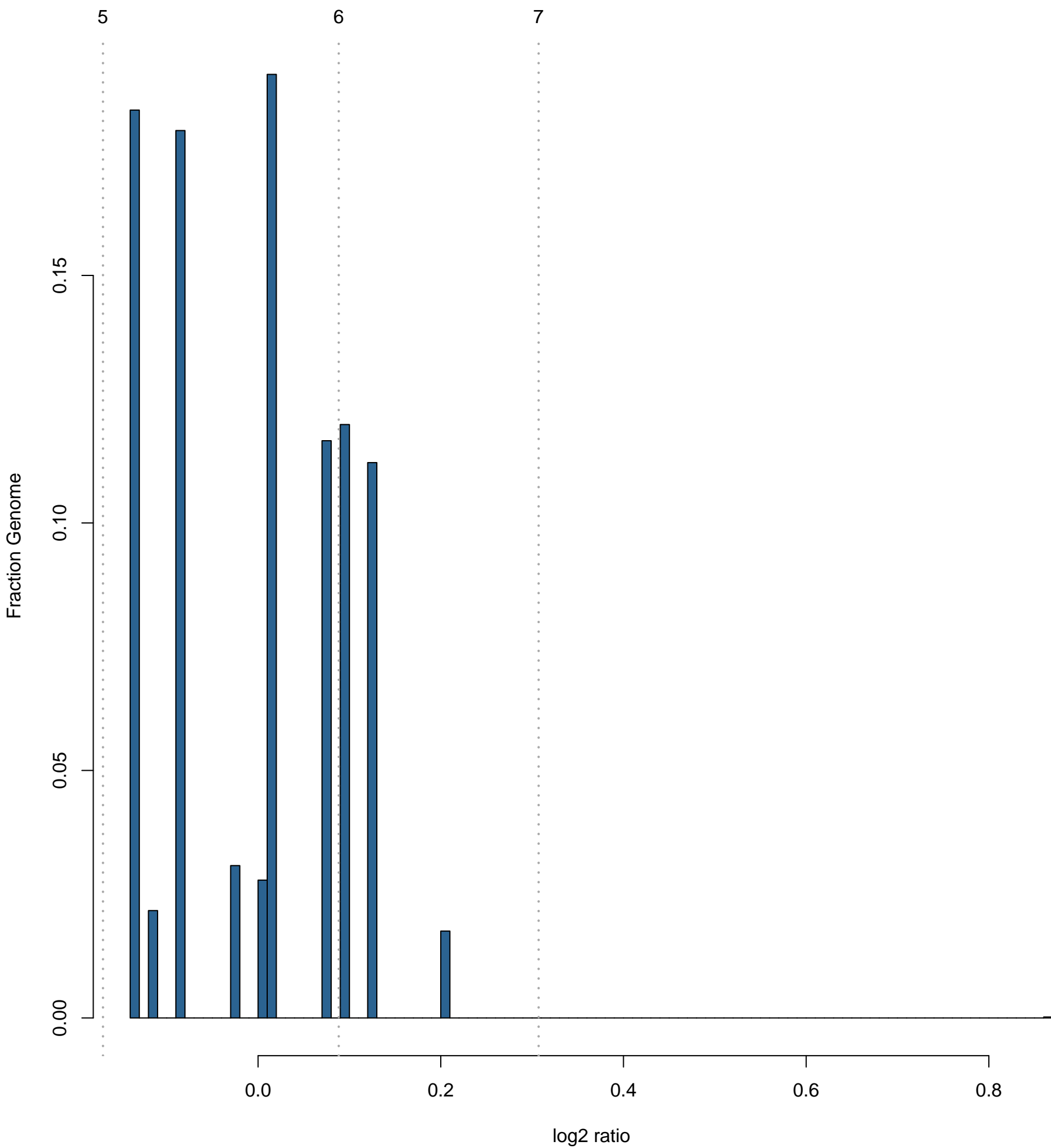


SCNA-fit log-likelihood: -19413.61

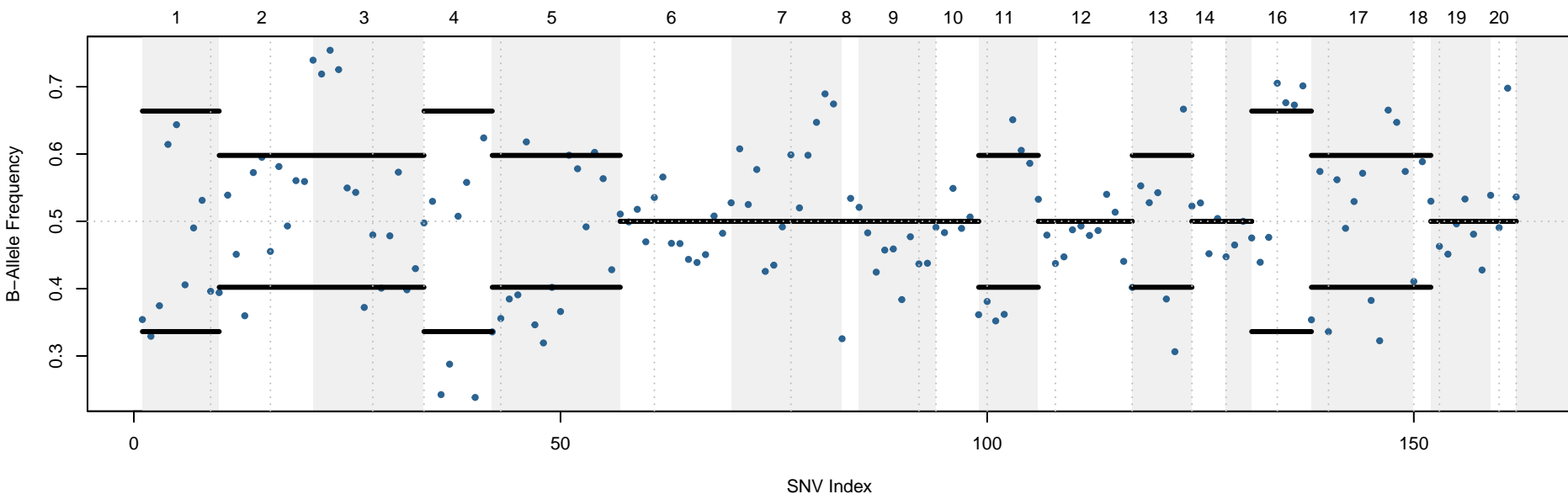




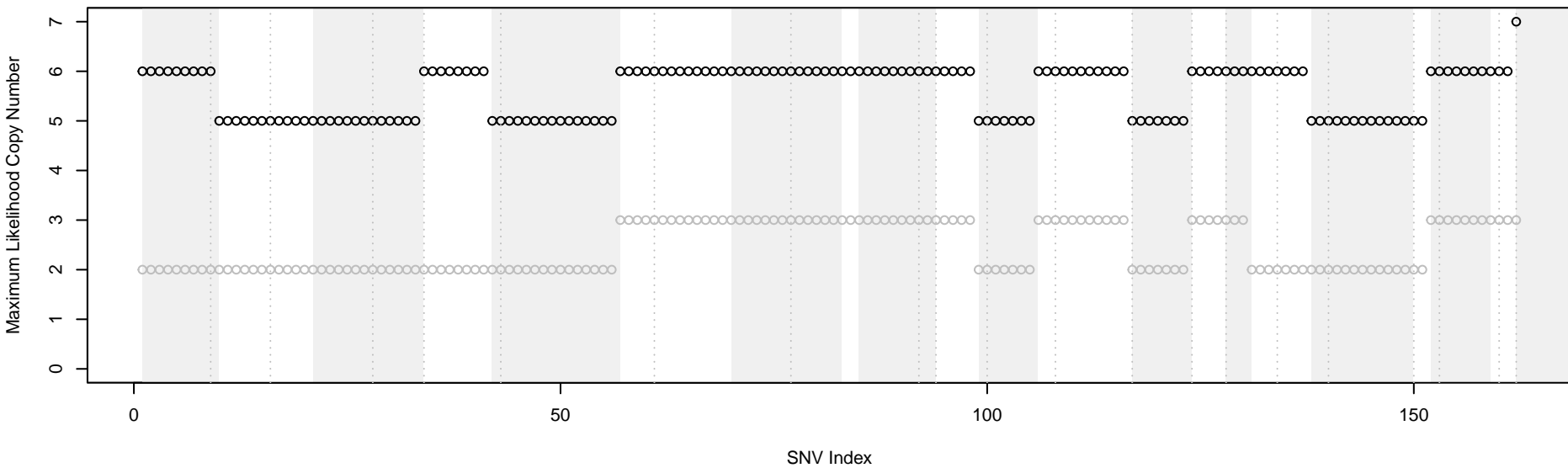
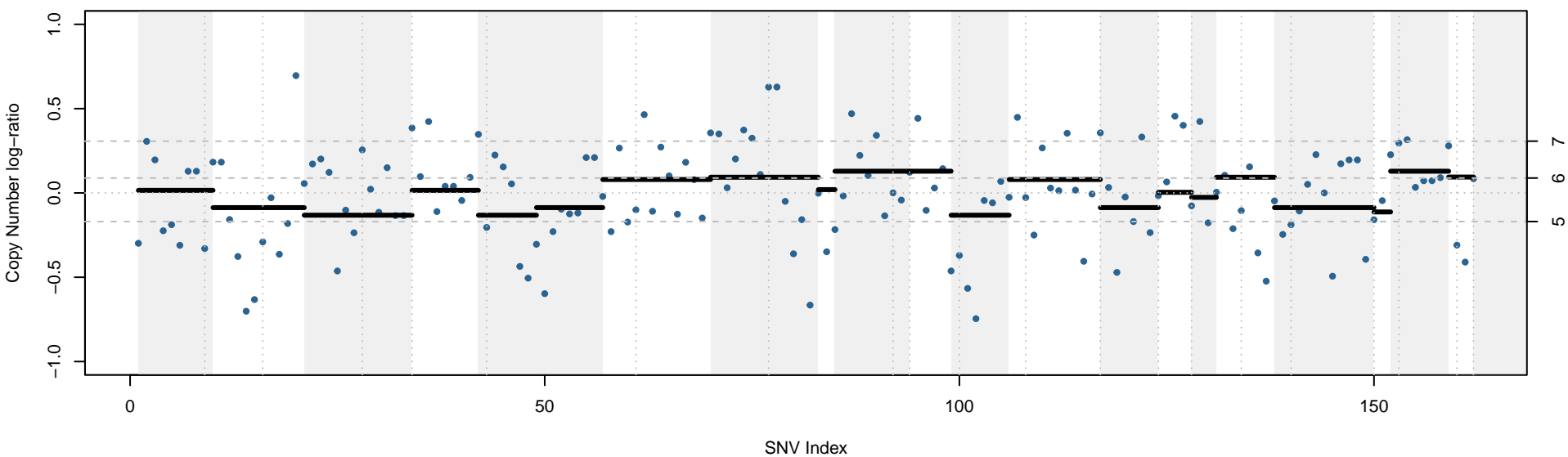
Purity: 0.95 Tumor ploidy: 5.638

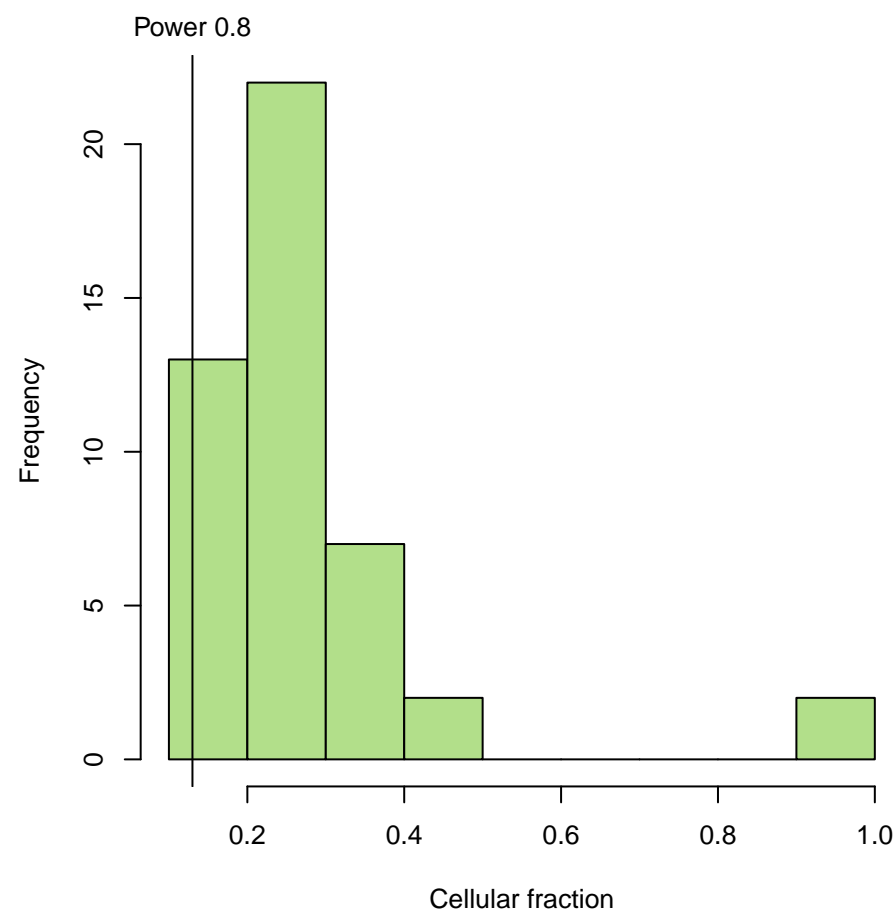
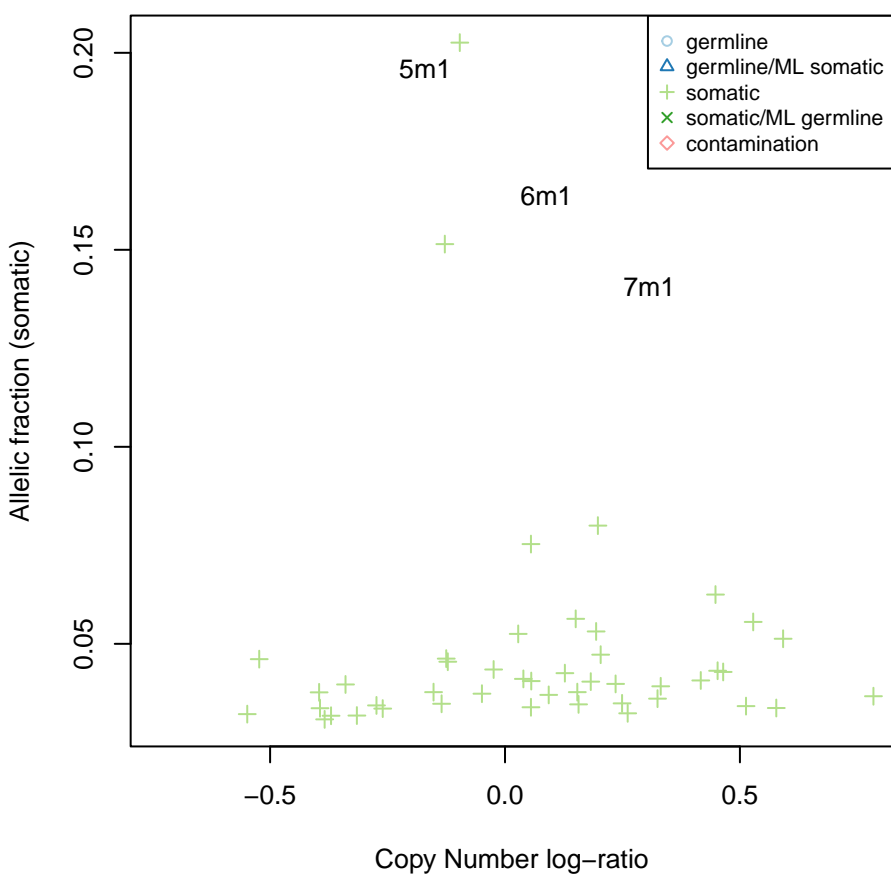
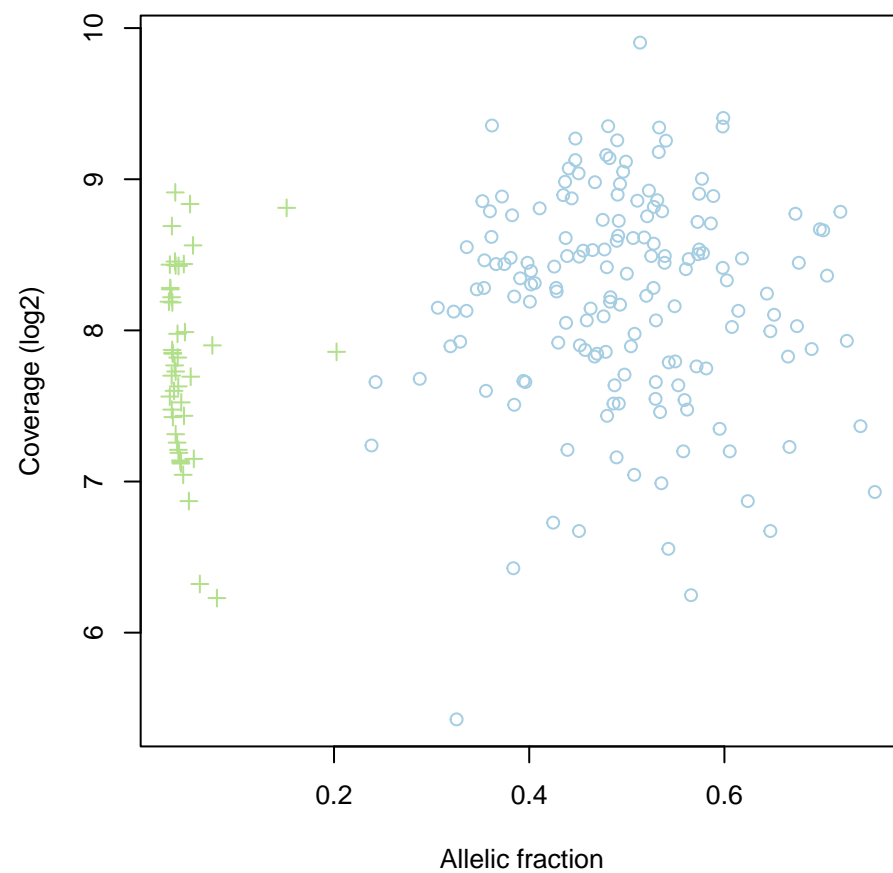
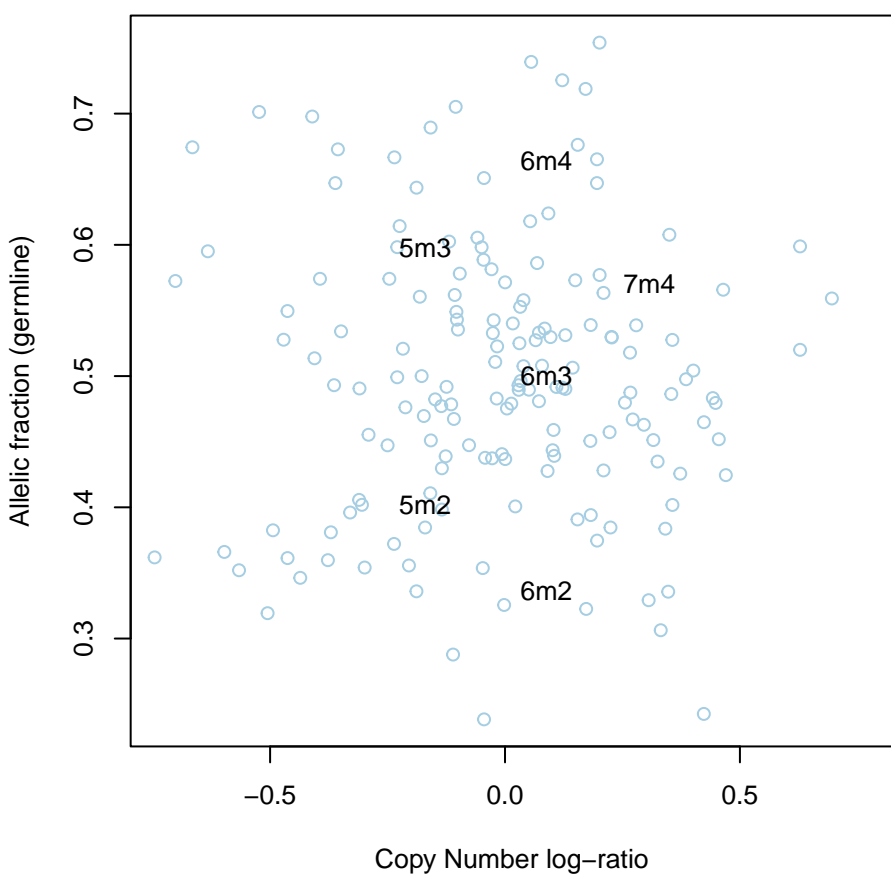


Purity: 0.95 Tumor ploidy: 5.638 SNV log-likelihood: -284.02 GoF: 85.8% Mean coverage: 312;424



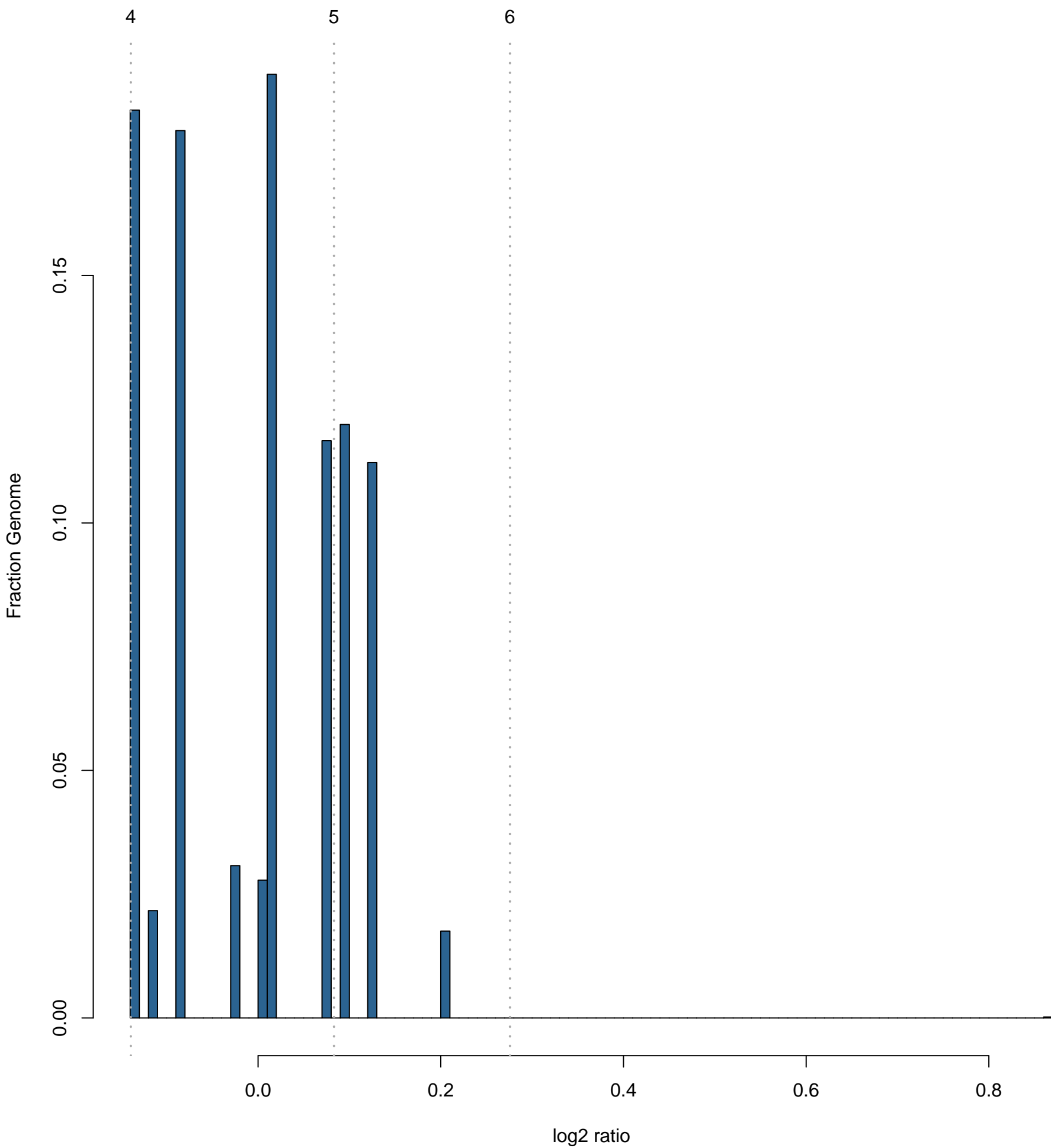
SCNA-fit log-likelihood: -19460.99



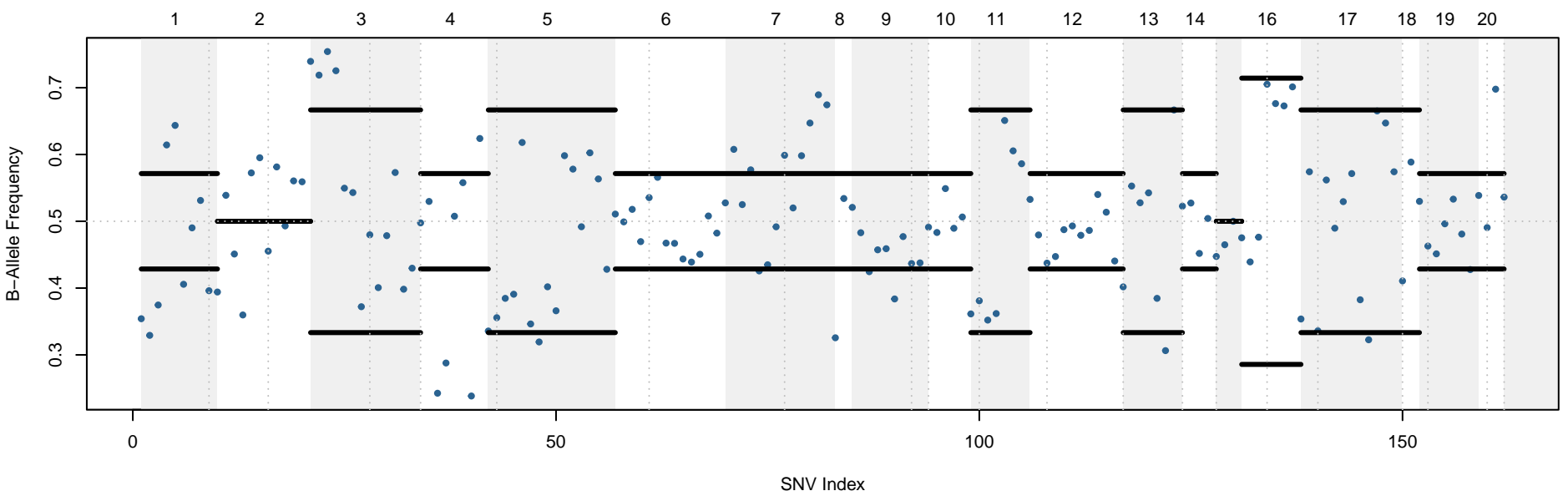




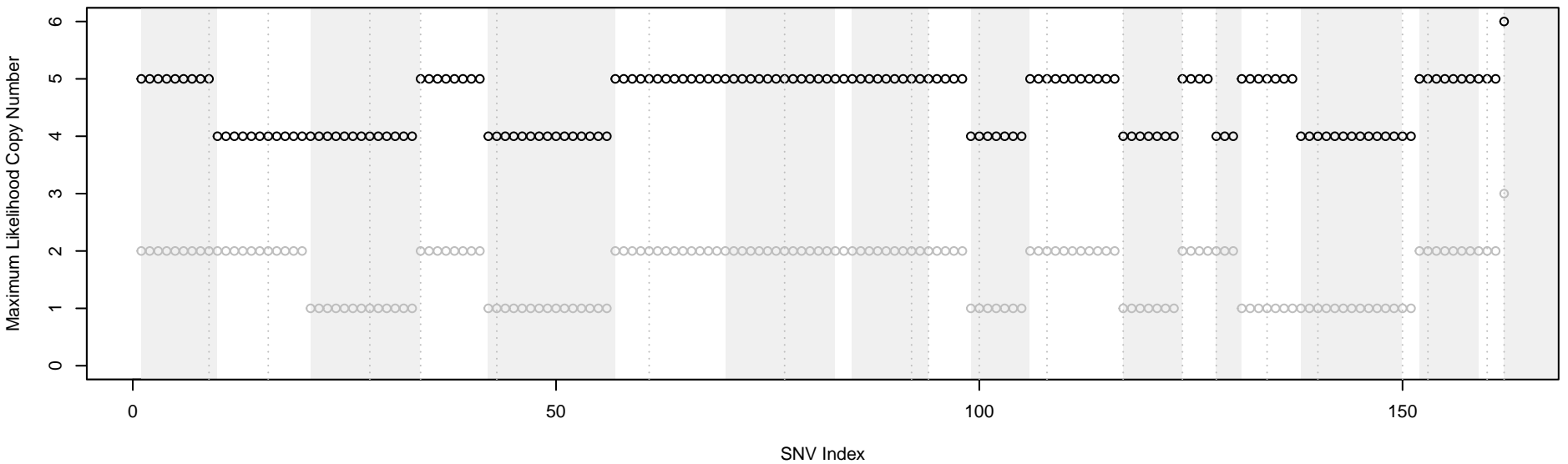
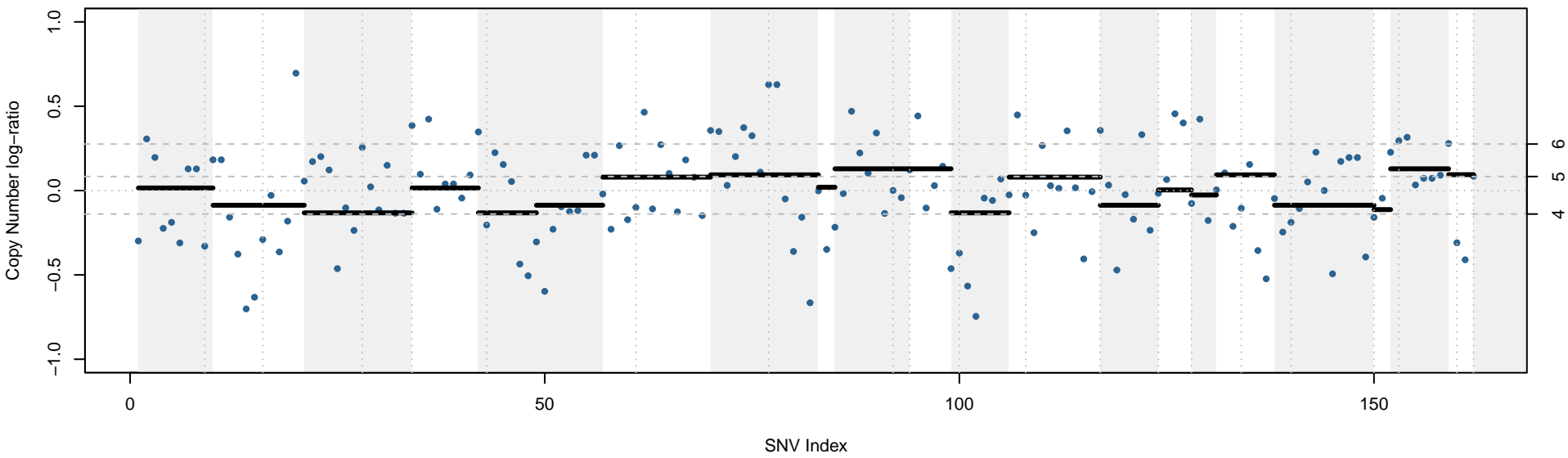
Purity: 0.5 Tumor ploidy: 4.608

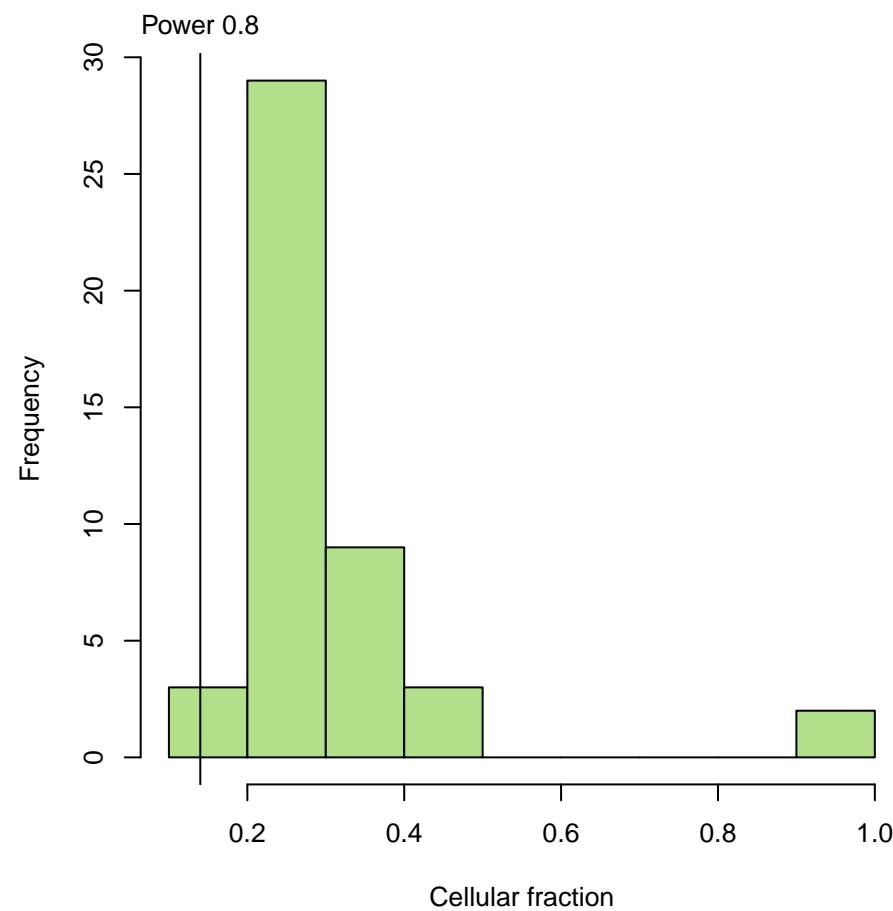
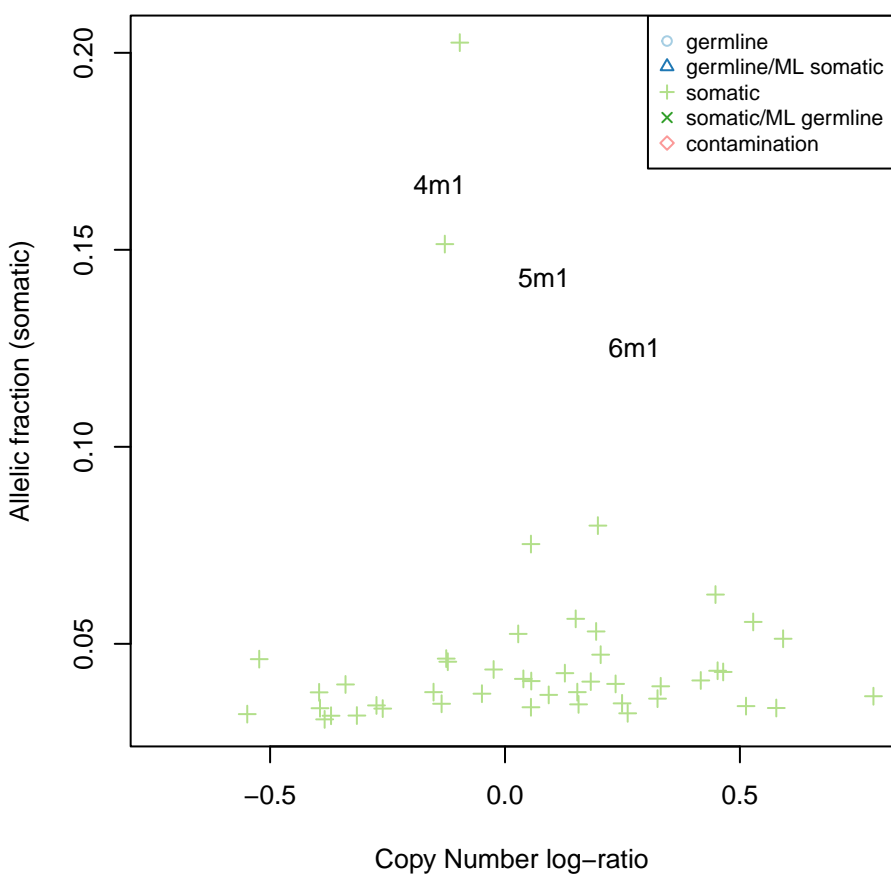
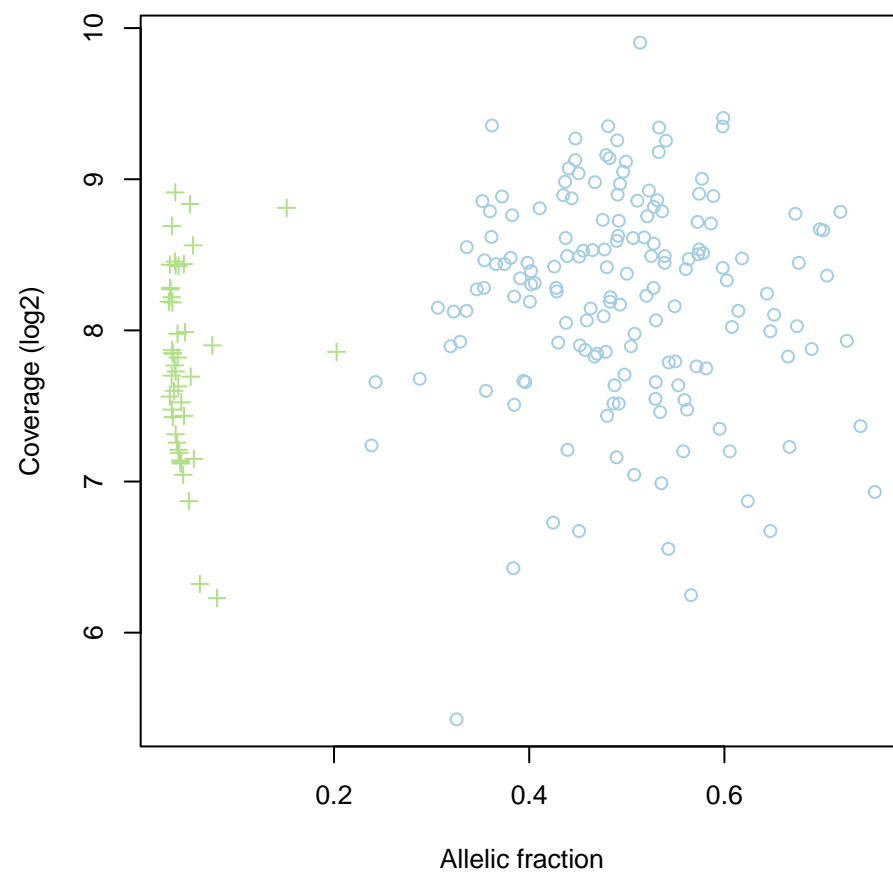
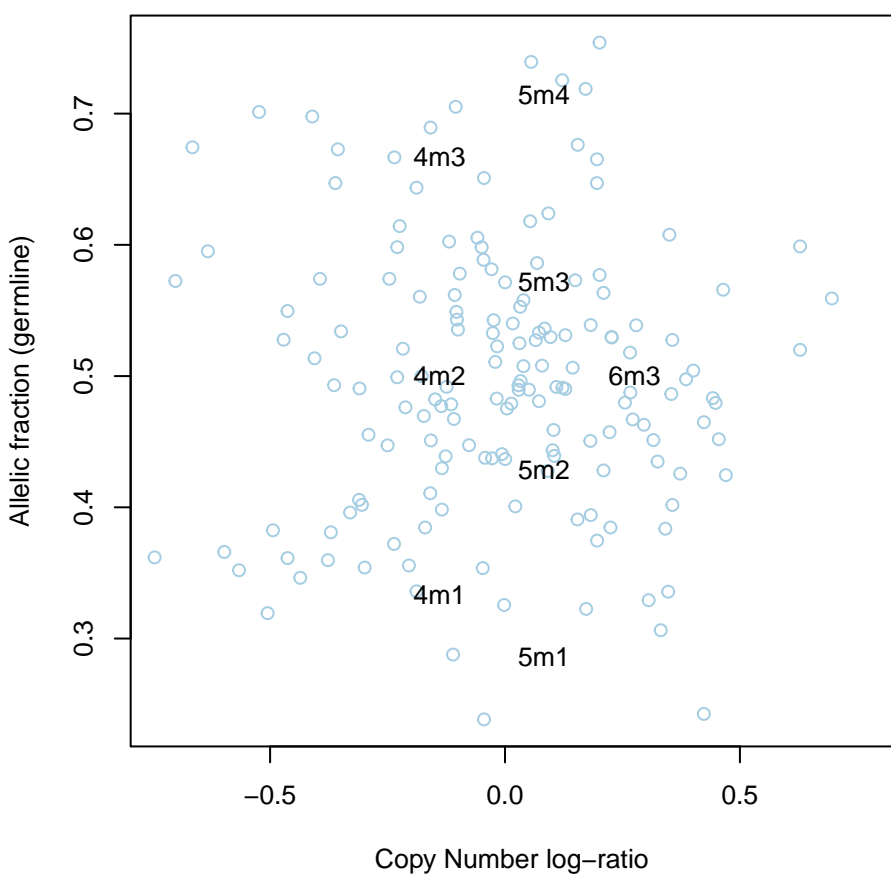


Purity: 0.5 Tumor ploidy: 4.608 SNV log-likelihood: -344.79 GoF: 87.7% Mean coverage: 312;424

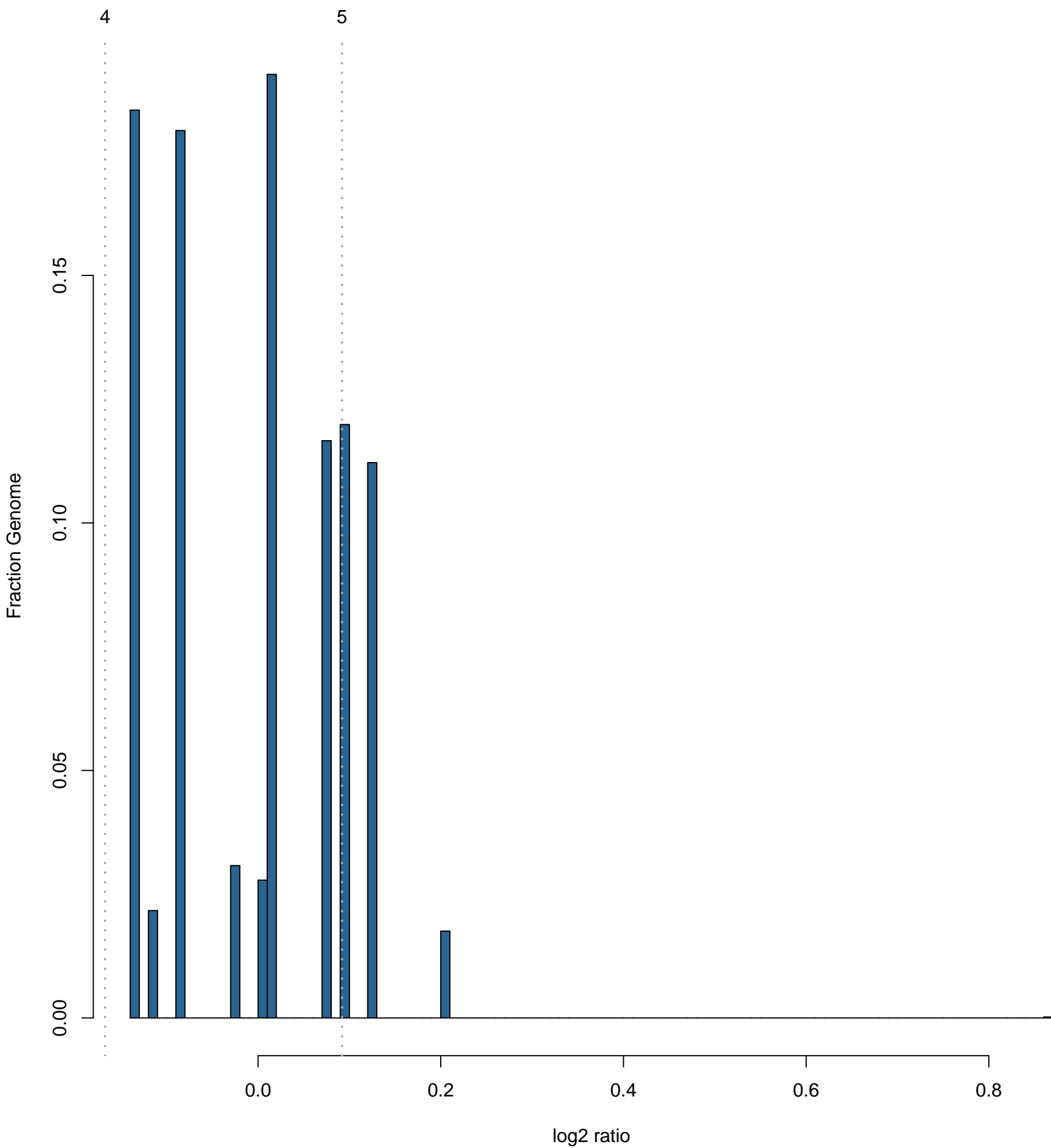


SCNA-fit log-likelihood: -19426.02

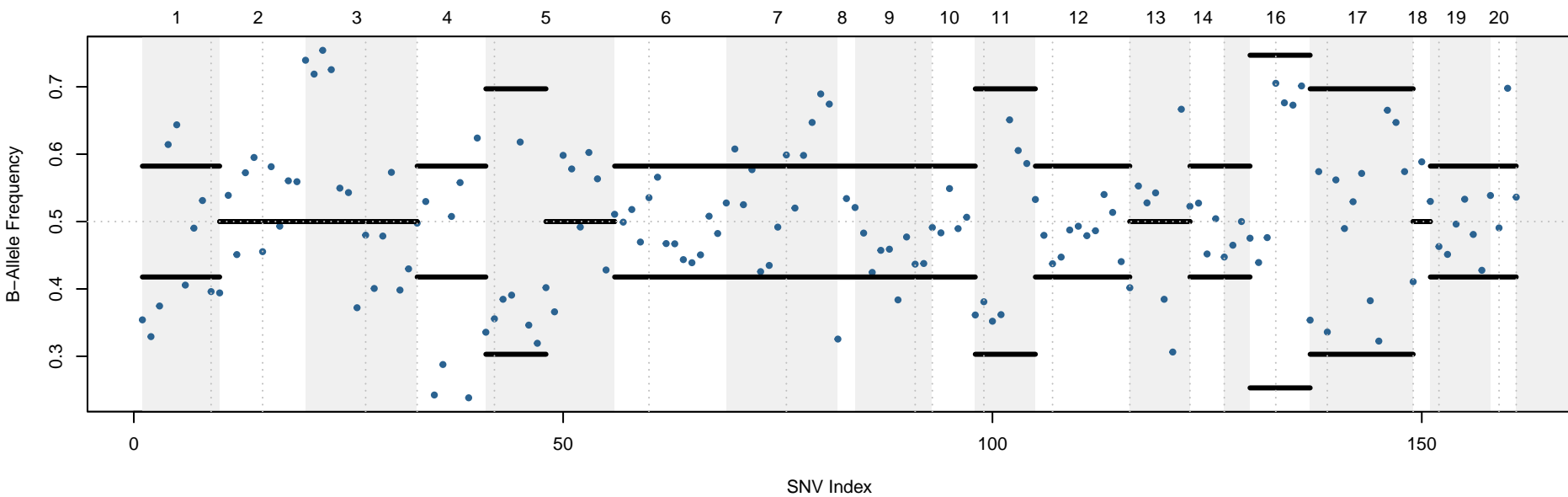




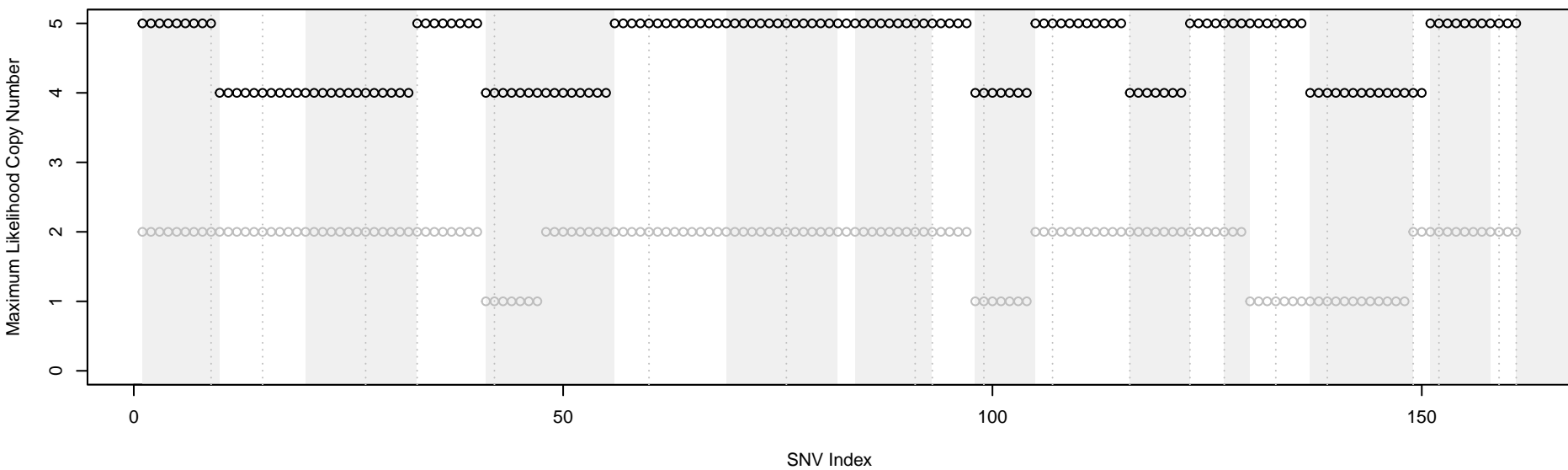
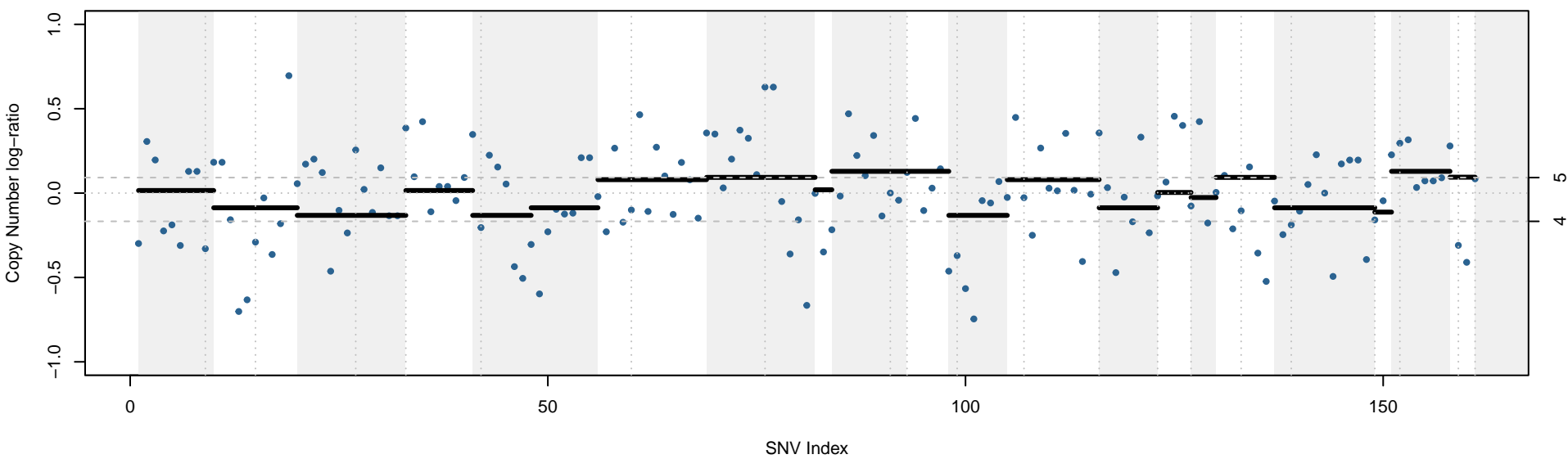
Purity: 0.65 Tumor ploidy: 4.625

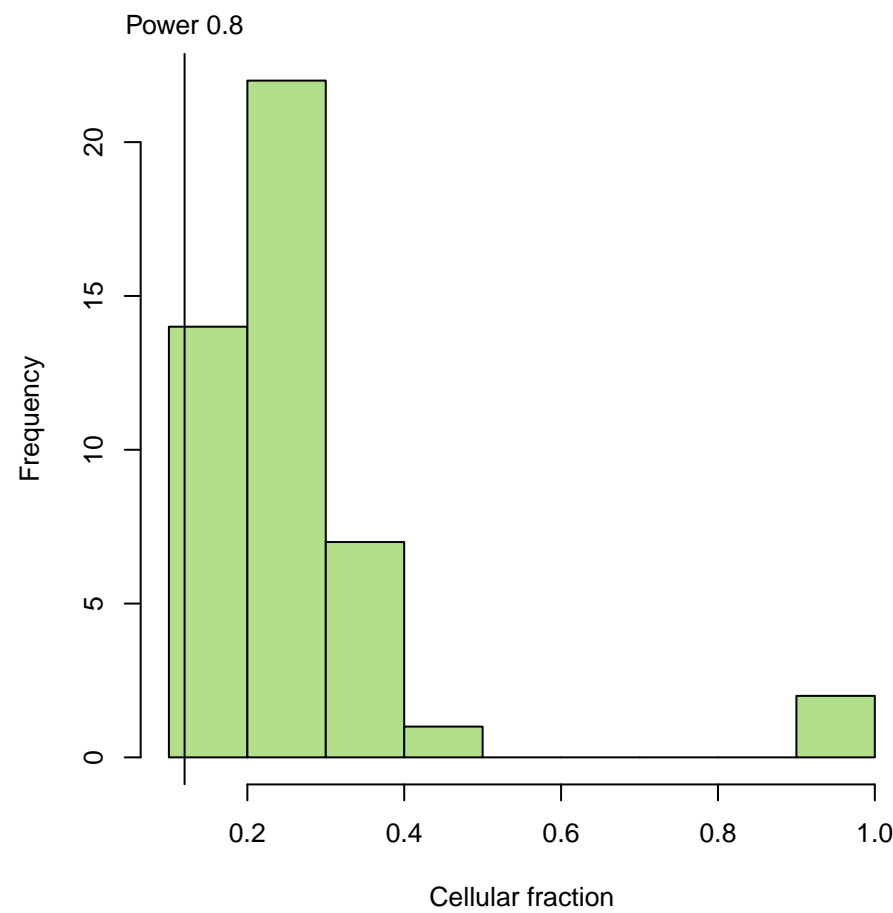
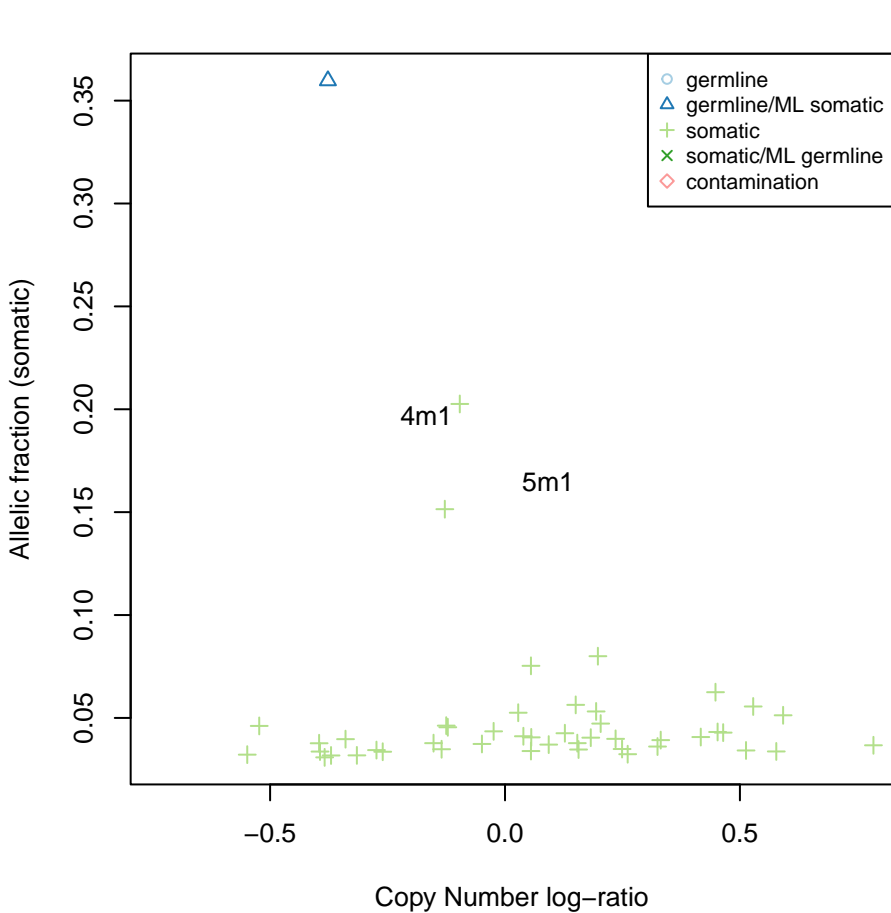
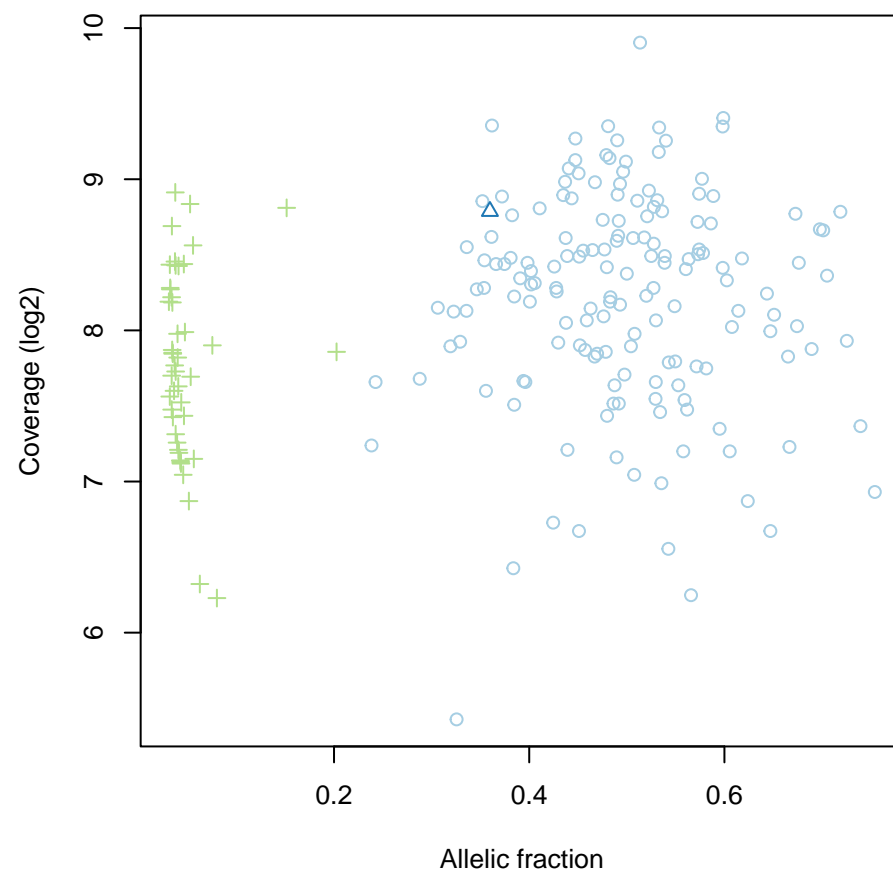
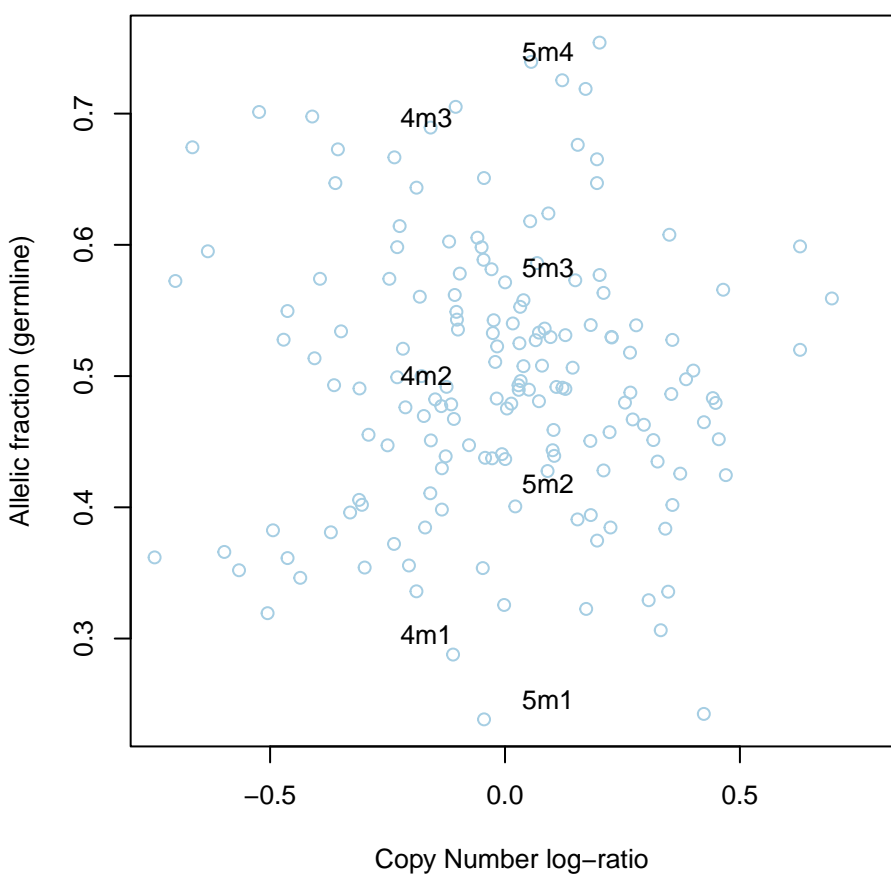


Purity: 0.65 Tumor ploidy: 4.625 SNV log-likelihood: -452.07 GoF: 82.6% Mean coverage: 312;424

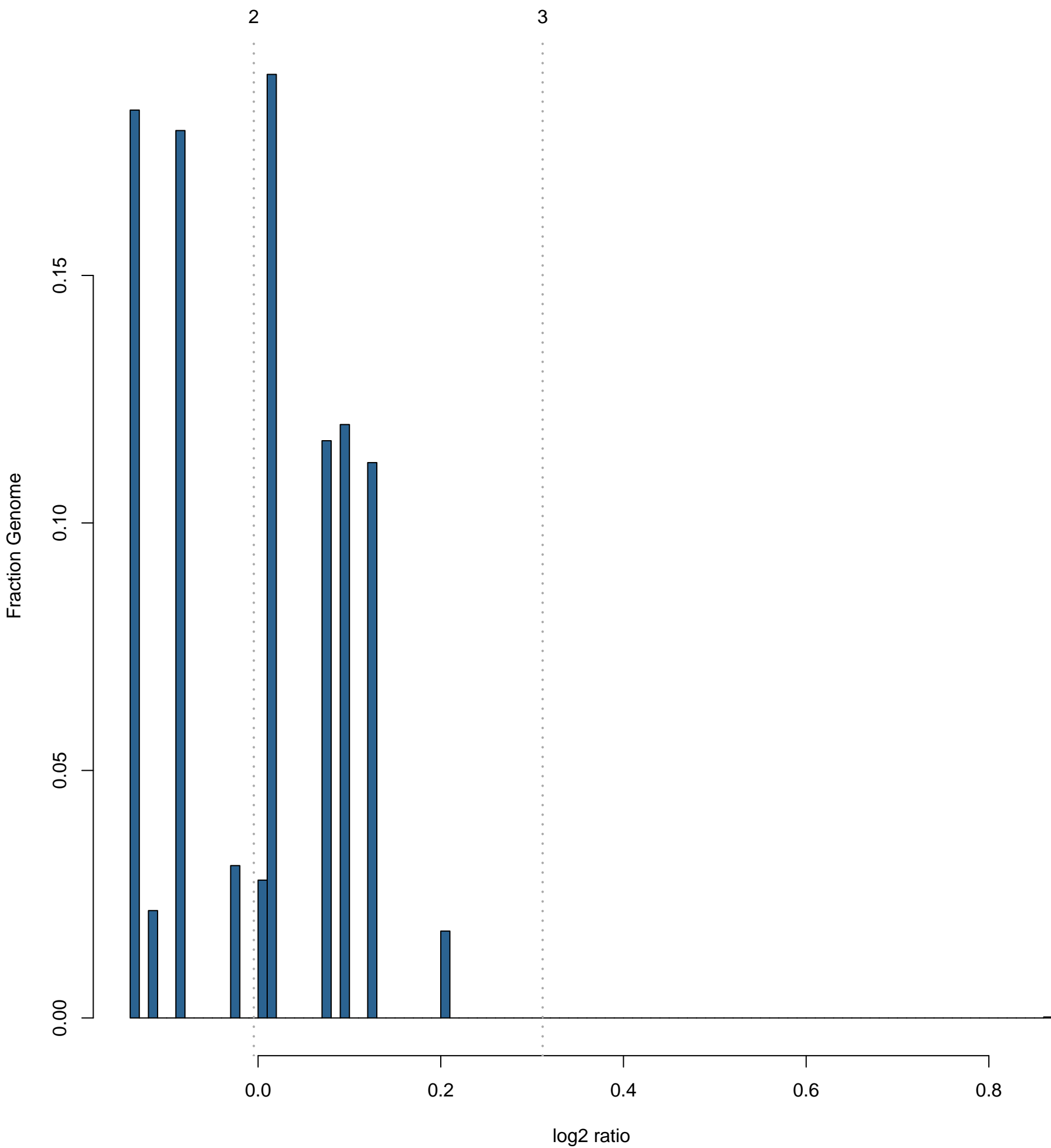


SCNA-fit log-likelihood: -19461.55

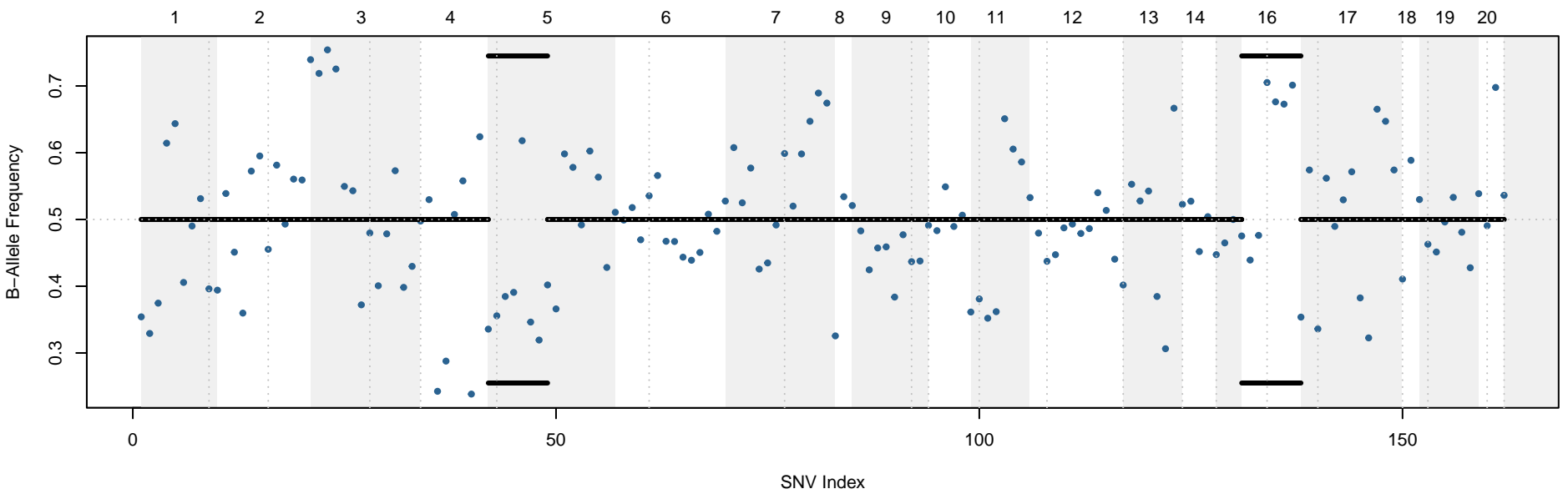




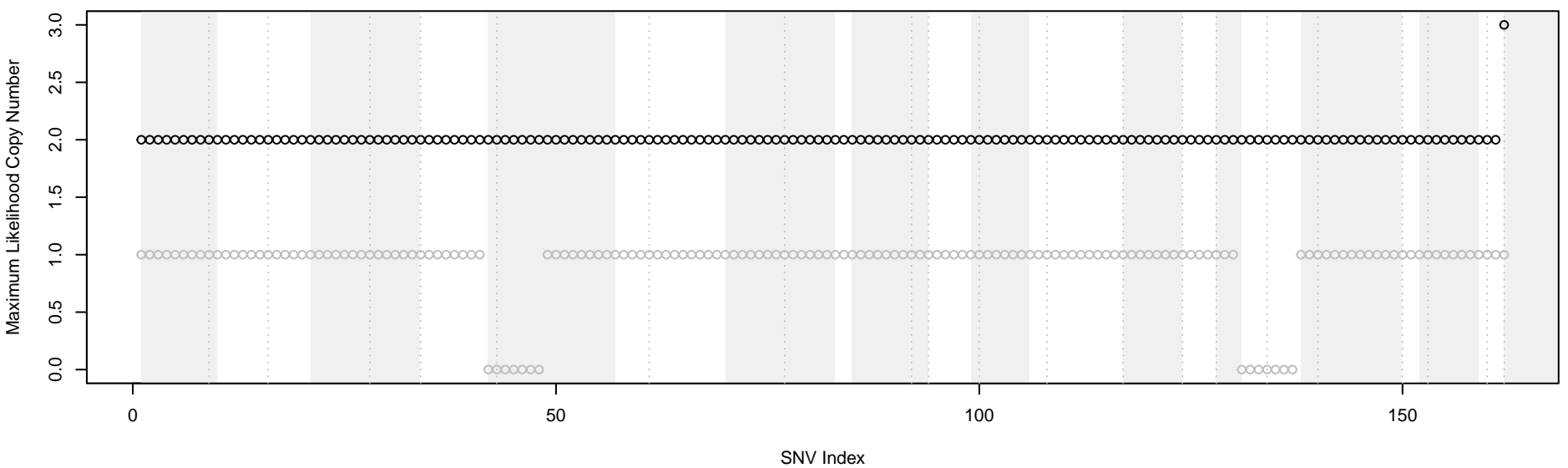
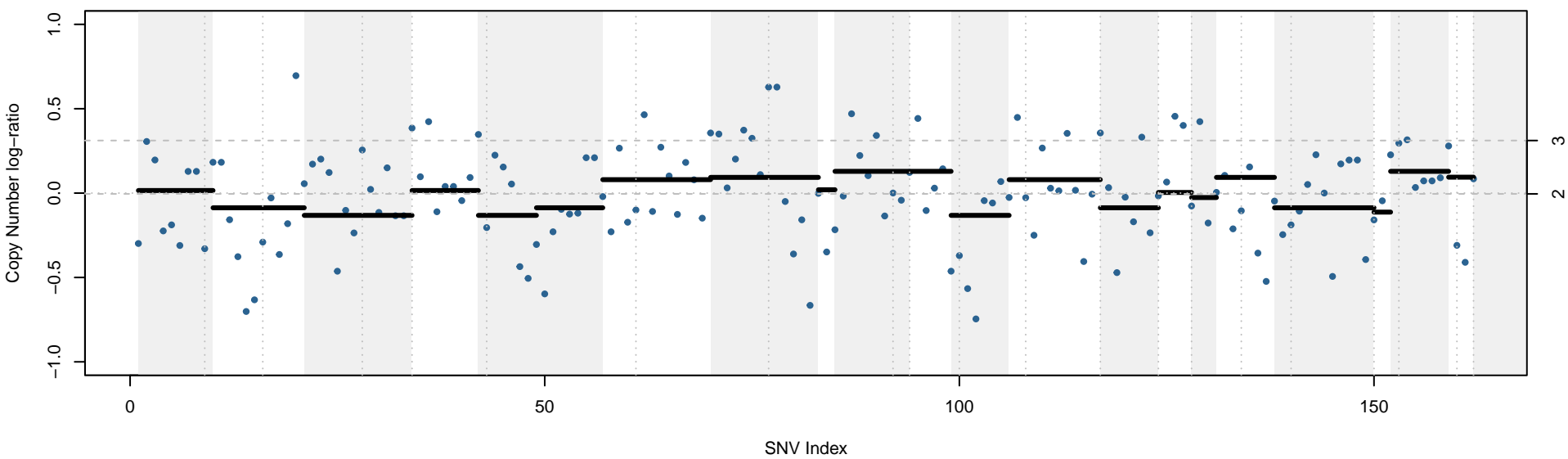
Purity: 0.49 Tumor ploidy: 2.013



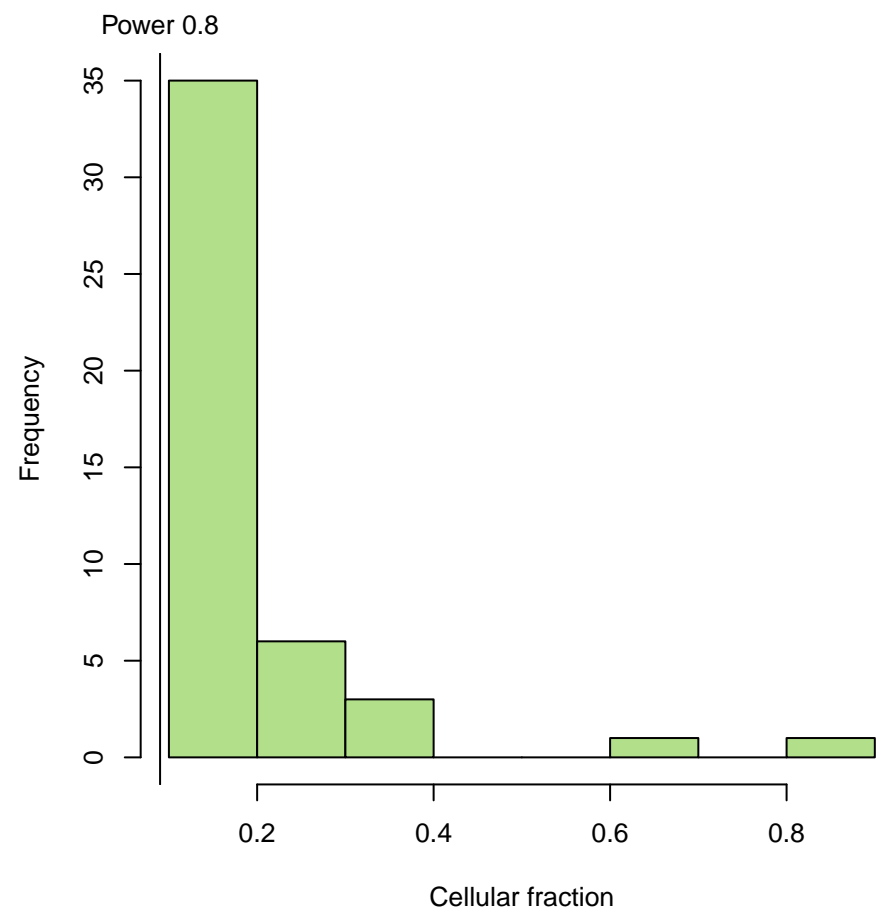
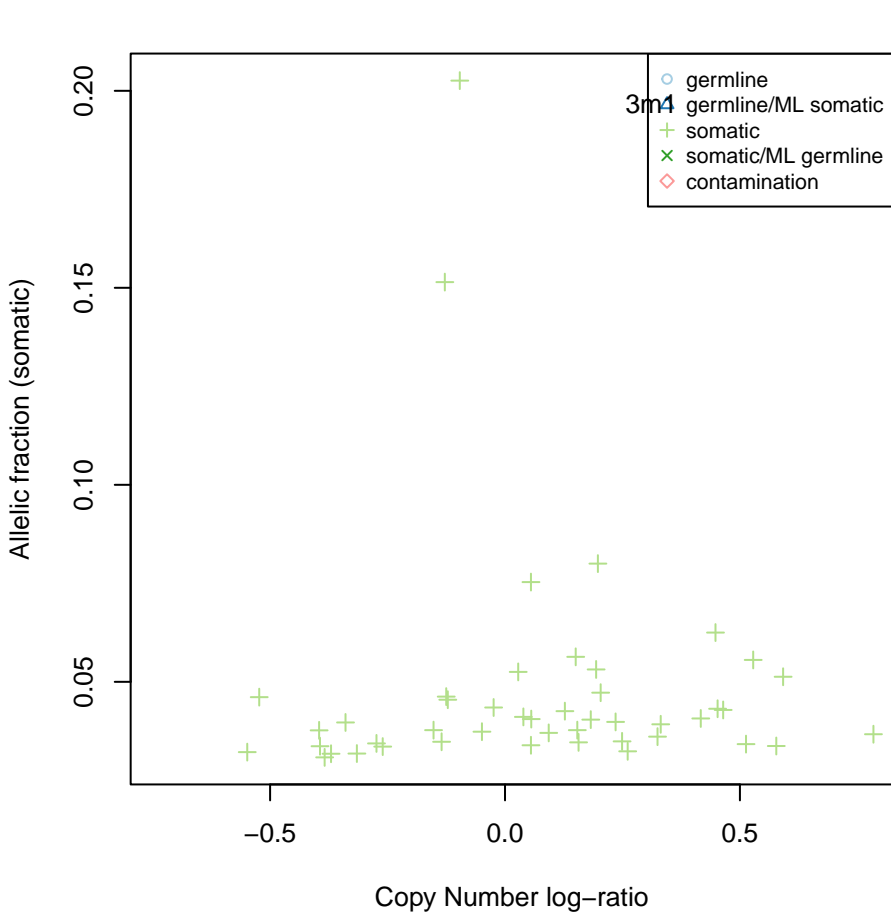
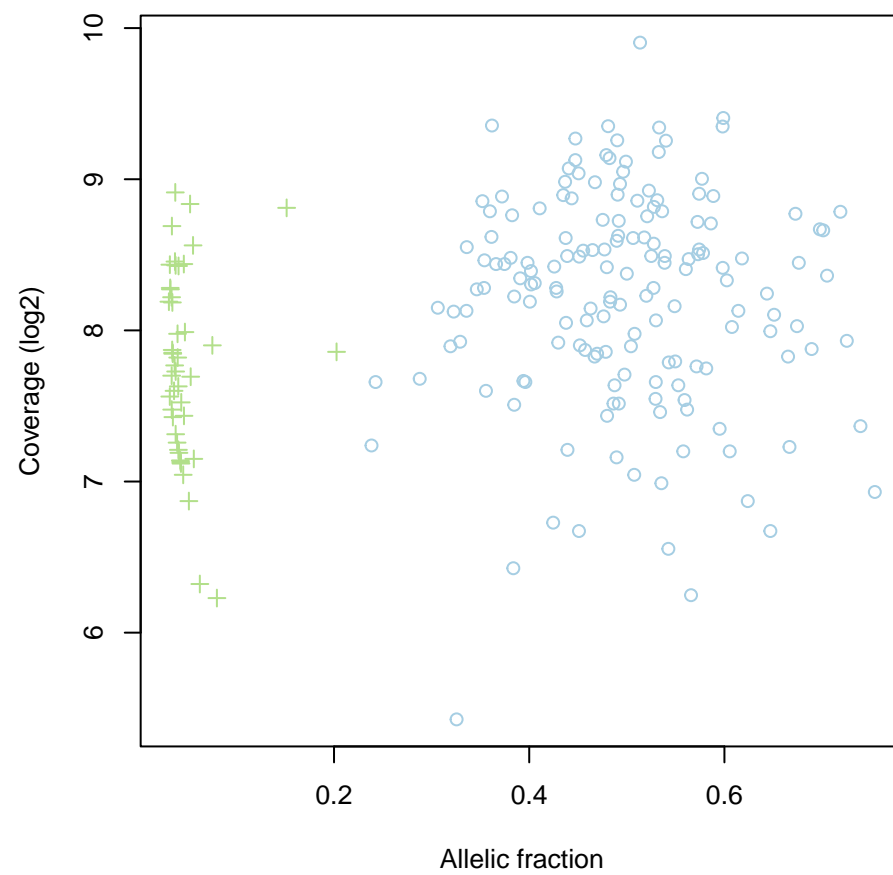
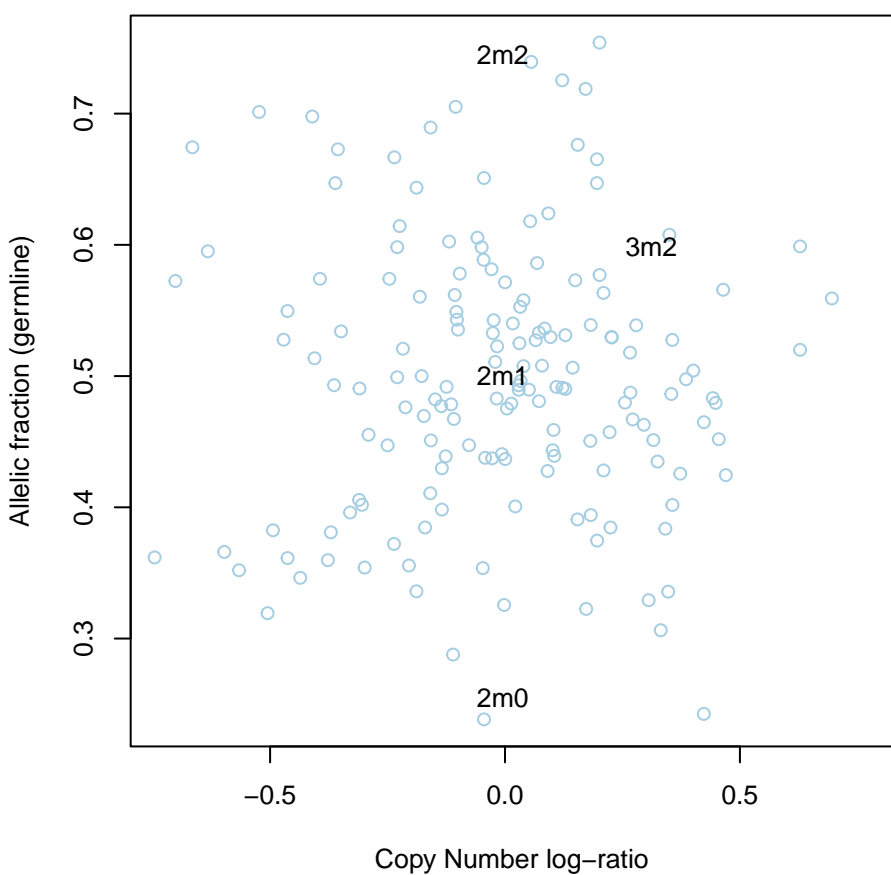
Purity: 0.49 Tumor ploidy: 2.013 SNV log-likelihood: -446.25 GoF: 68.8% Mean coverage: 312;424



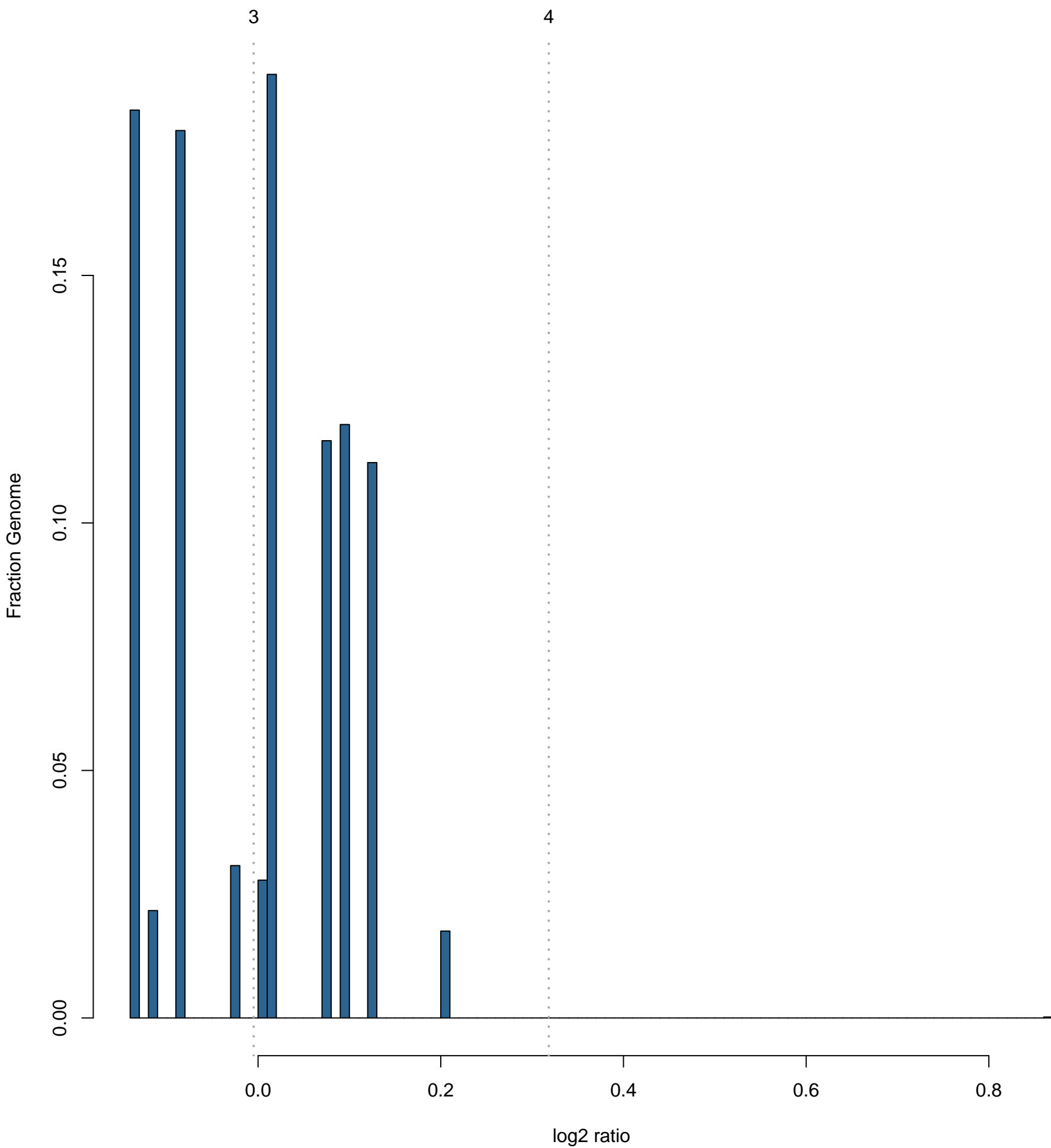
SCNA-fit log-likelihood: -19596.57



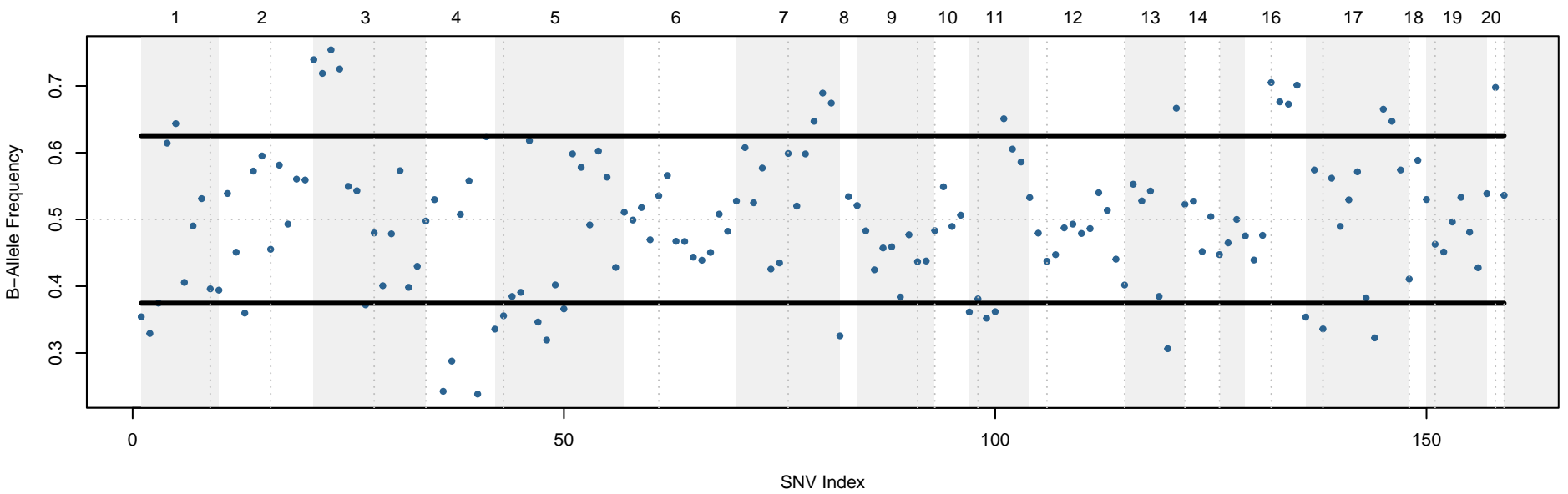




Purity: 0.67 Tumor ploidy: 3.013



Purity: 0.67 Tumor ploidy: 3.013 SNV log-likelihood: -559.15 GoF: 65.9% Mean coverage: 312;424



SCNA-fit log-likelihood: -19597.52

