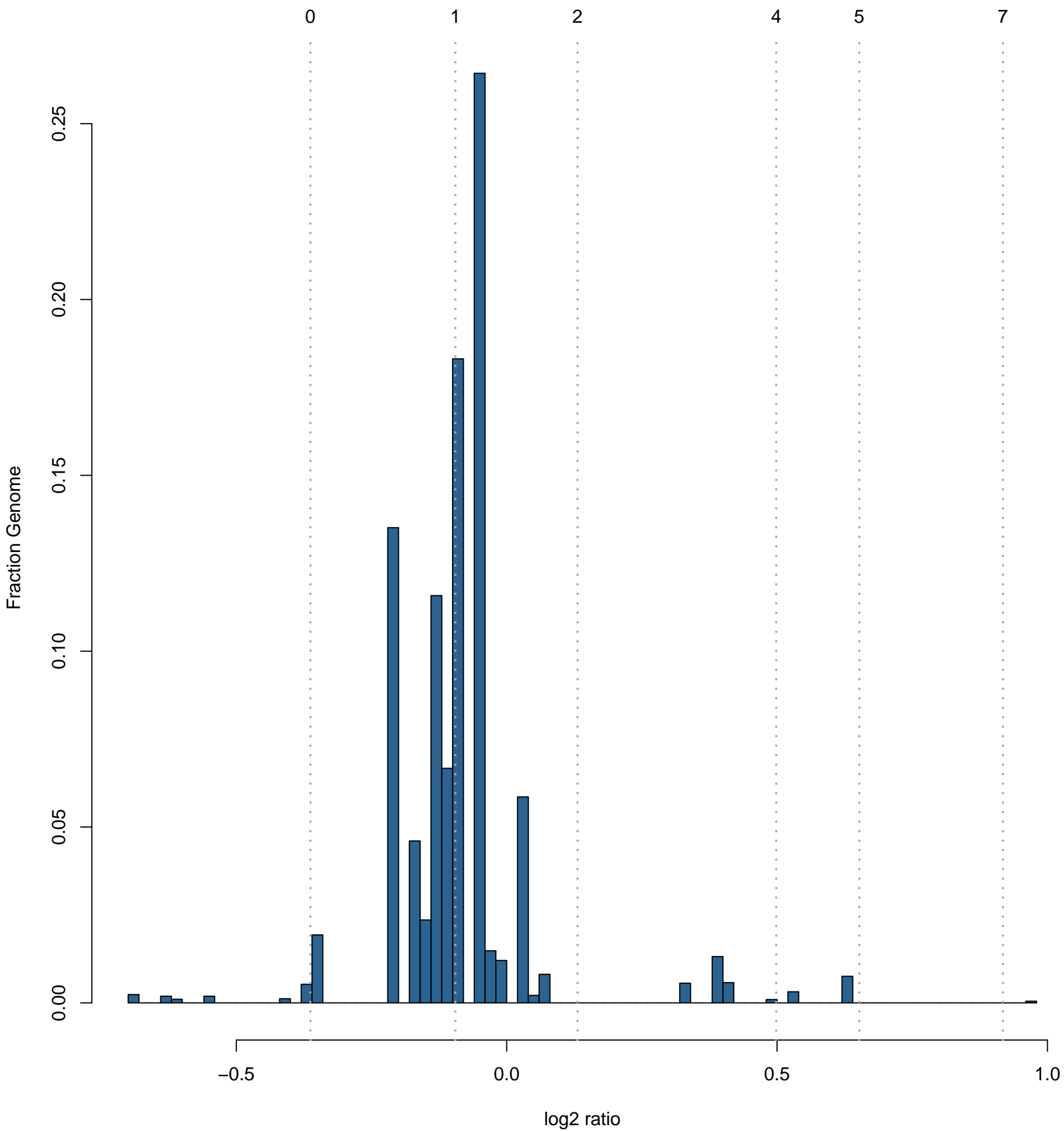
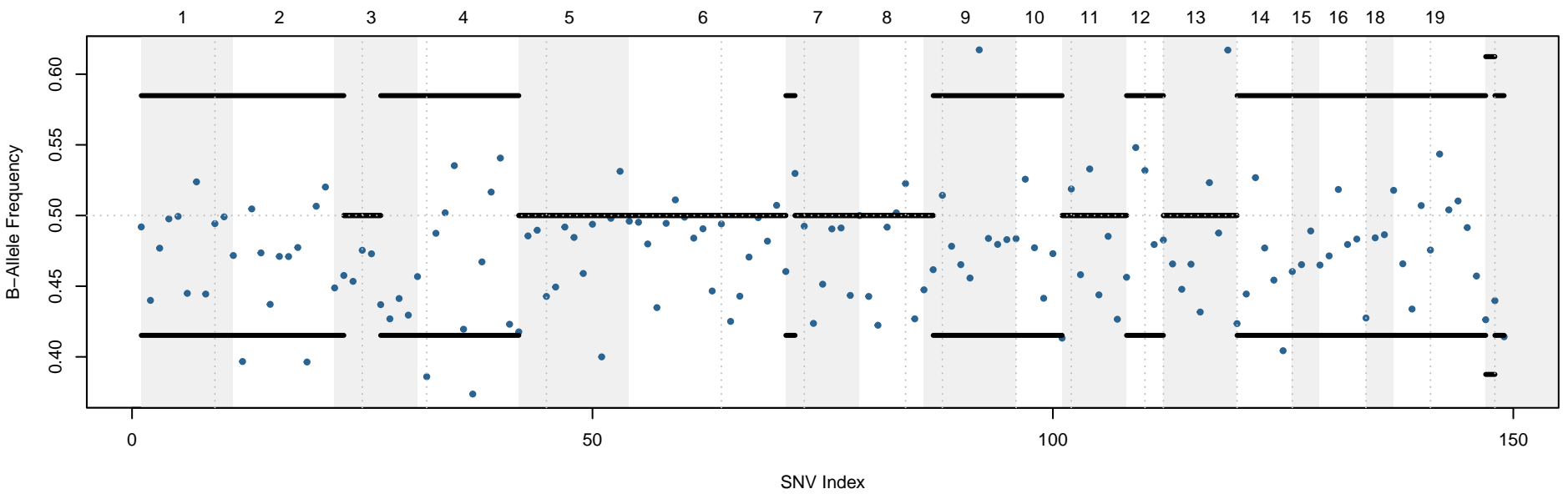


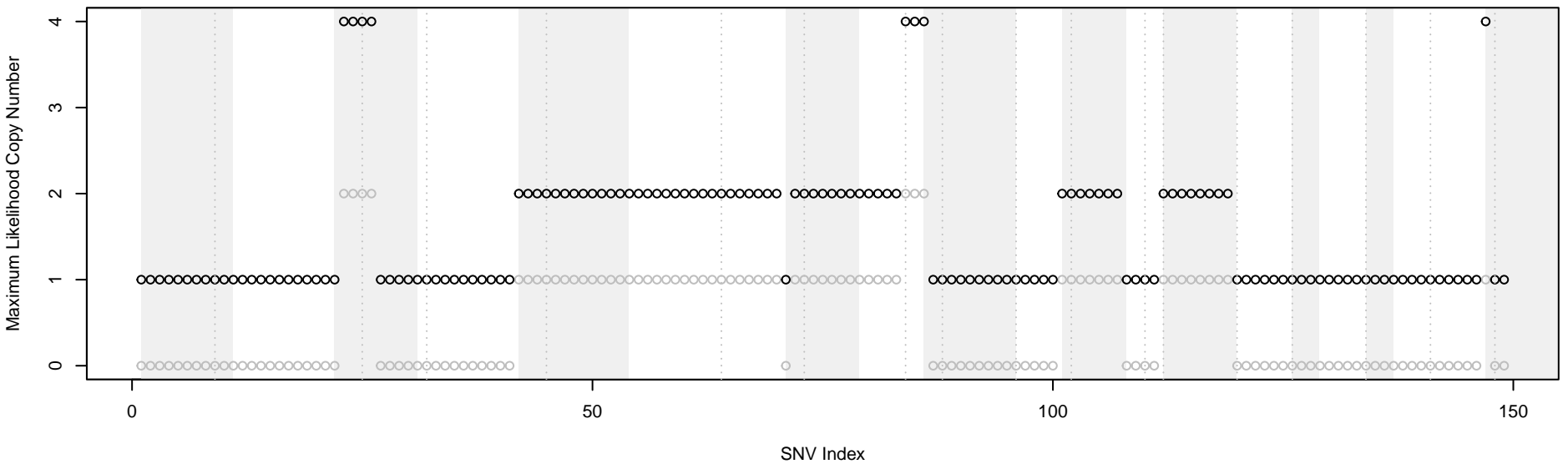
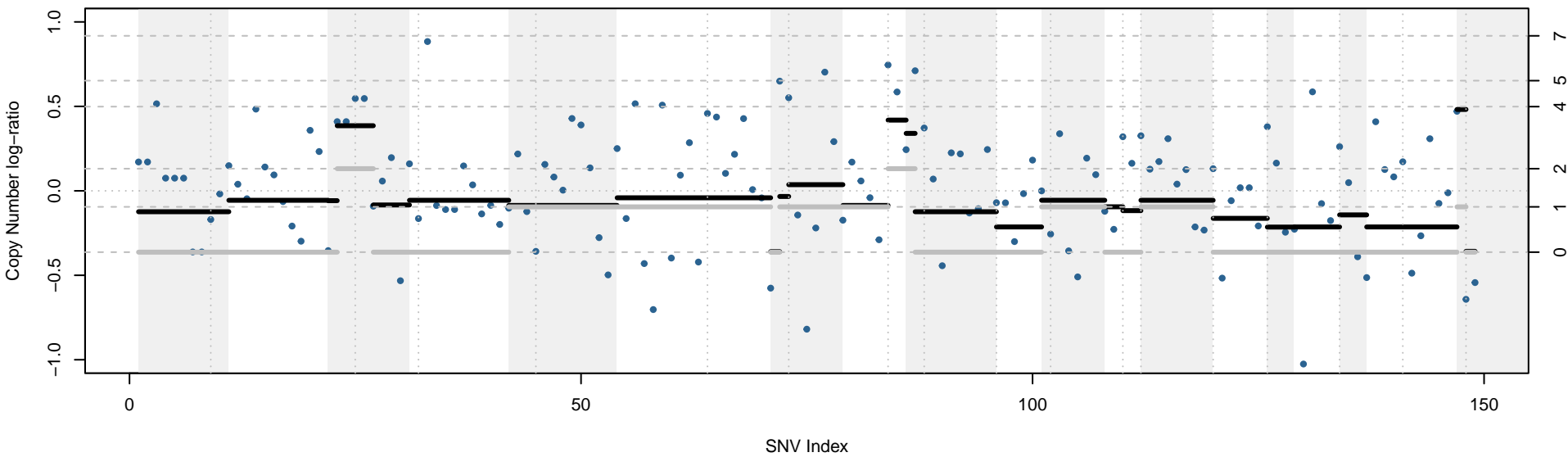
Purity: 0.29 Tumor ploidy: 1.401

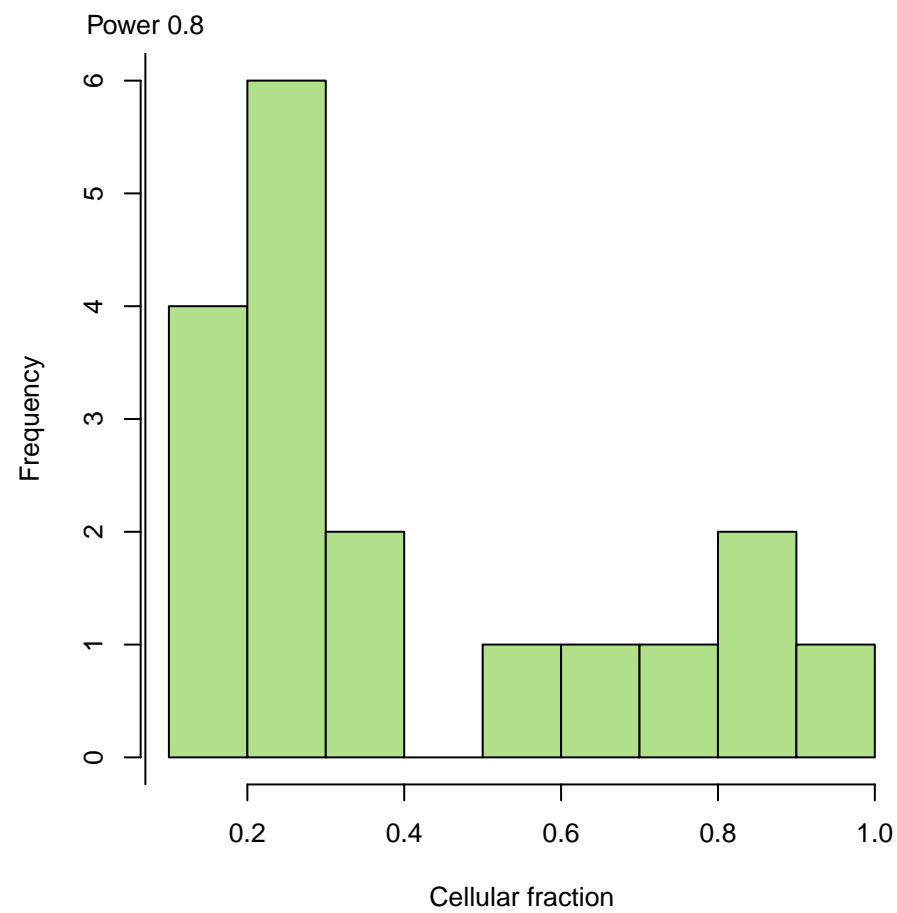
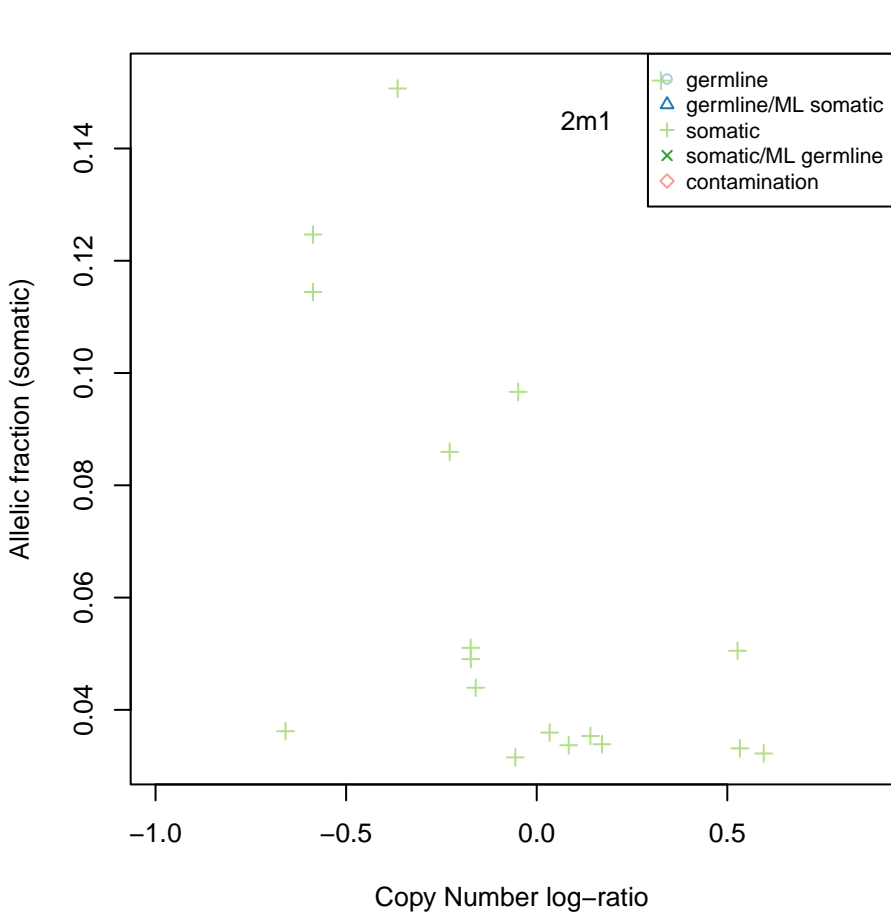
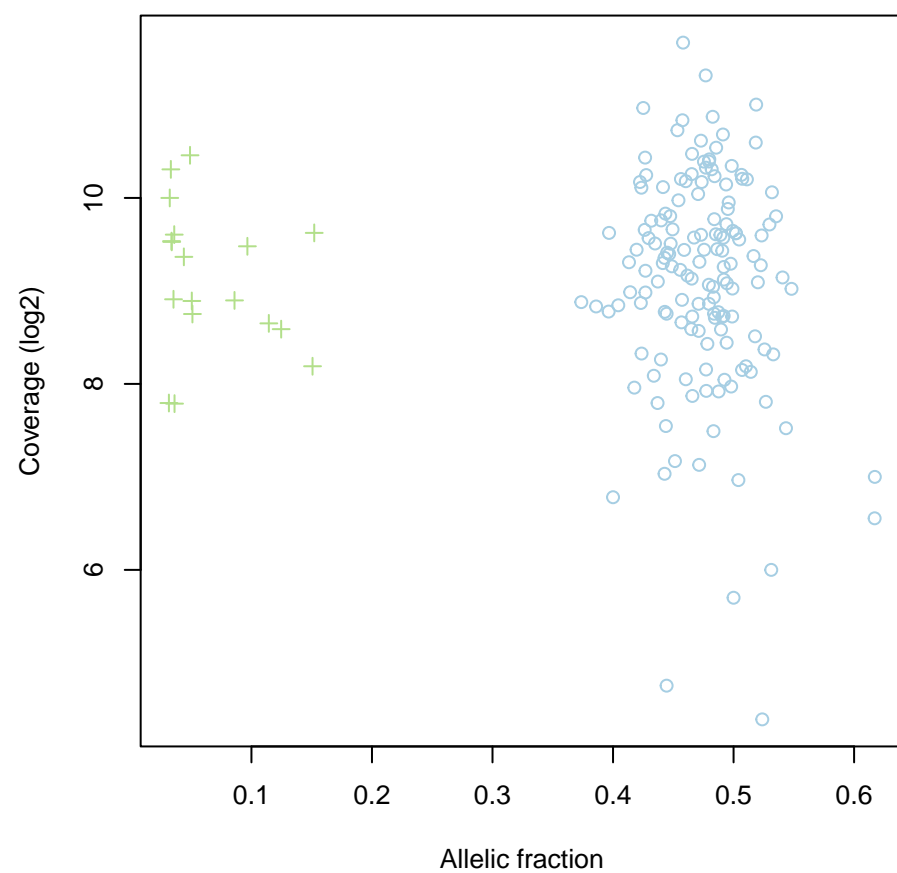
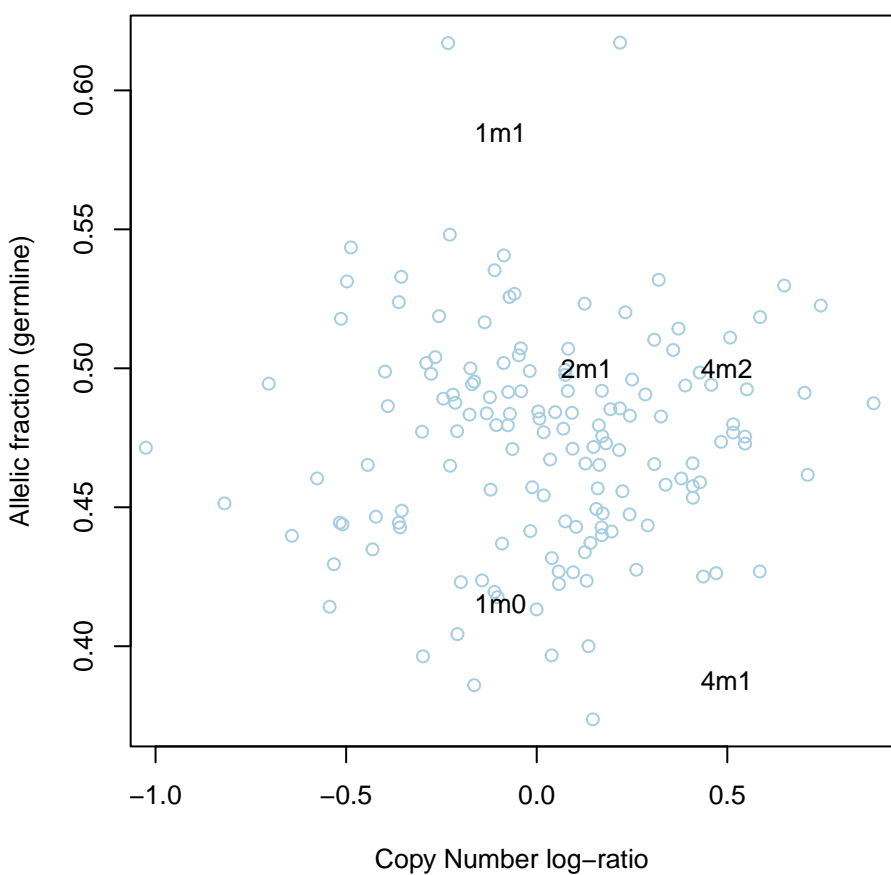


Purity: 0.29 Tumor ploidy: 1.401 SNV log-likelihood: -104.88 GoF: 91.5% Mean coverage: 714;414

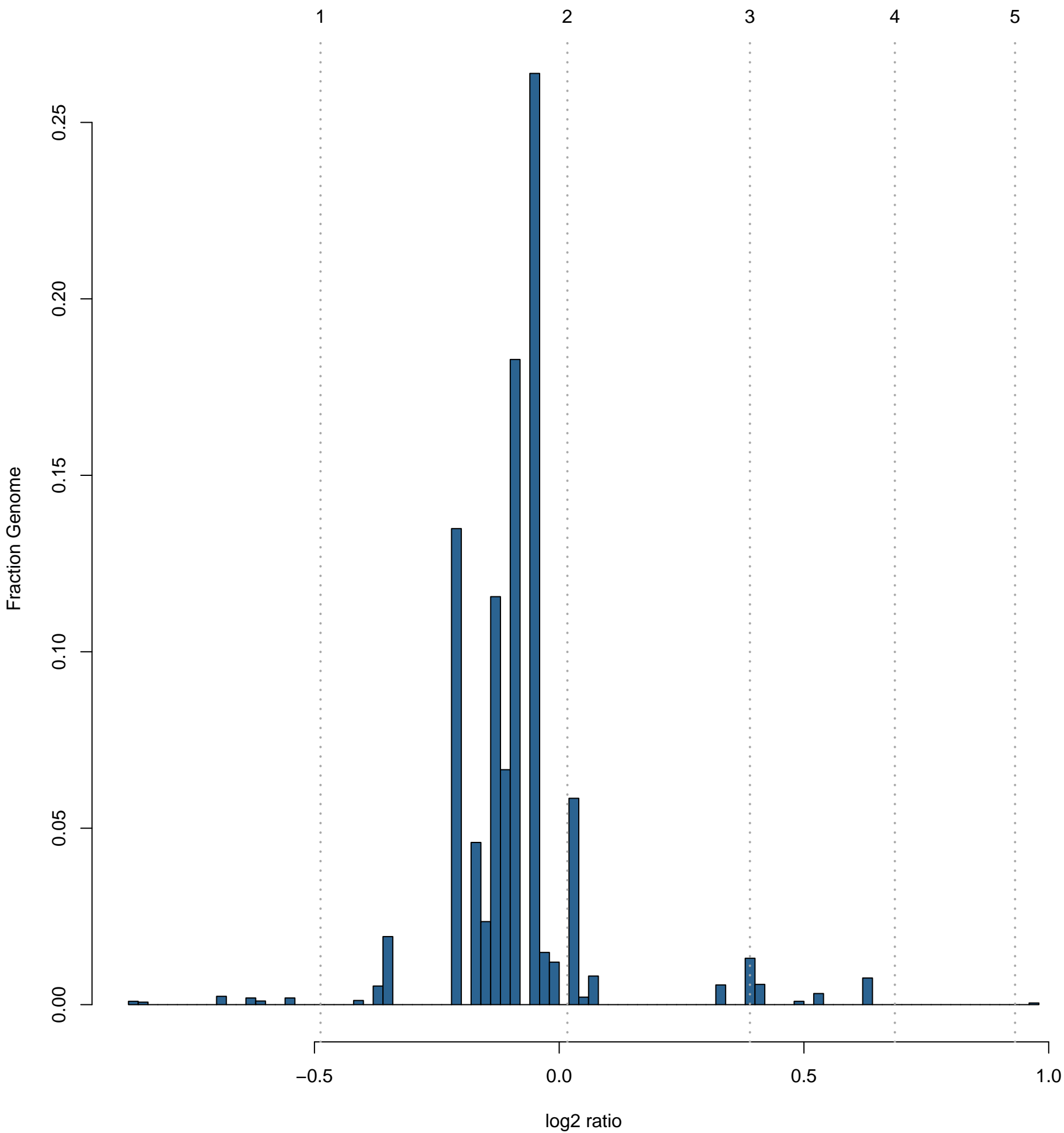


SCNA-fit log-likelihood: -9389.46

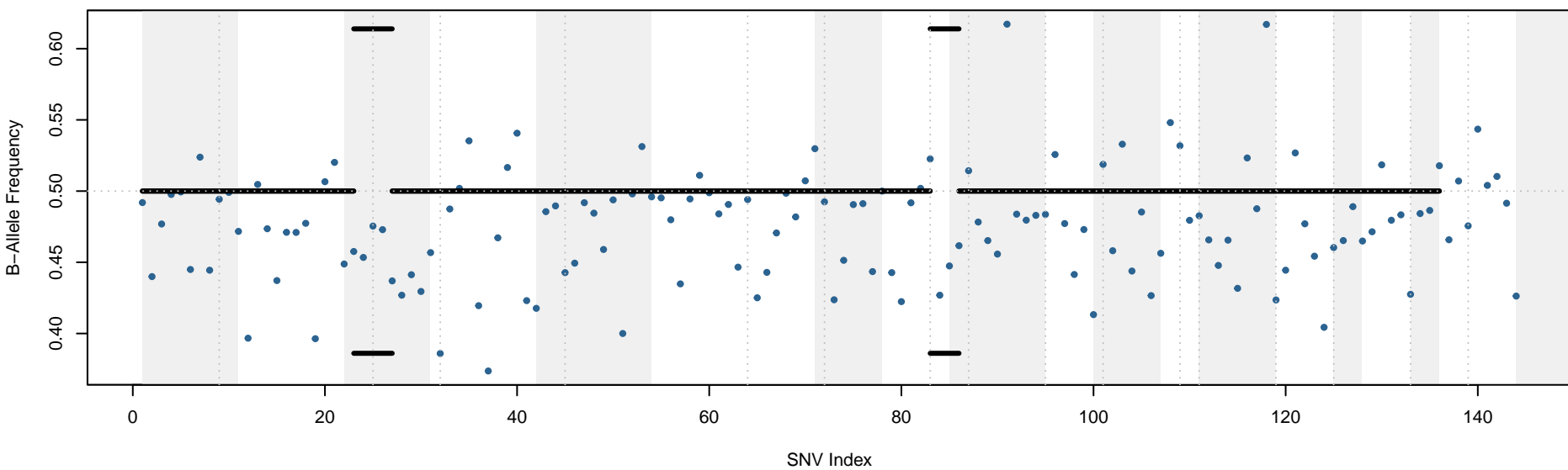




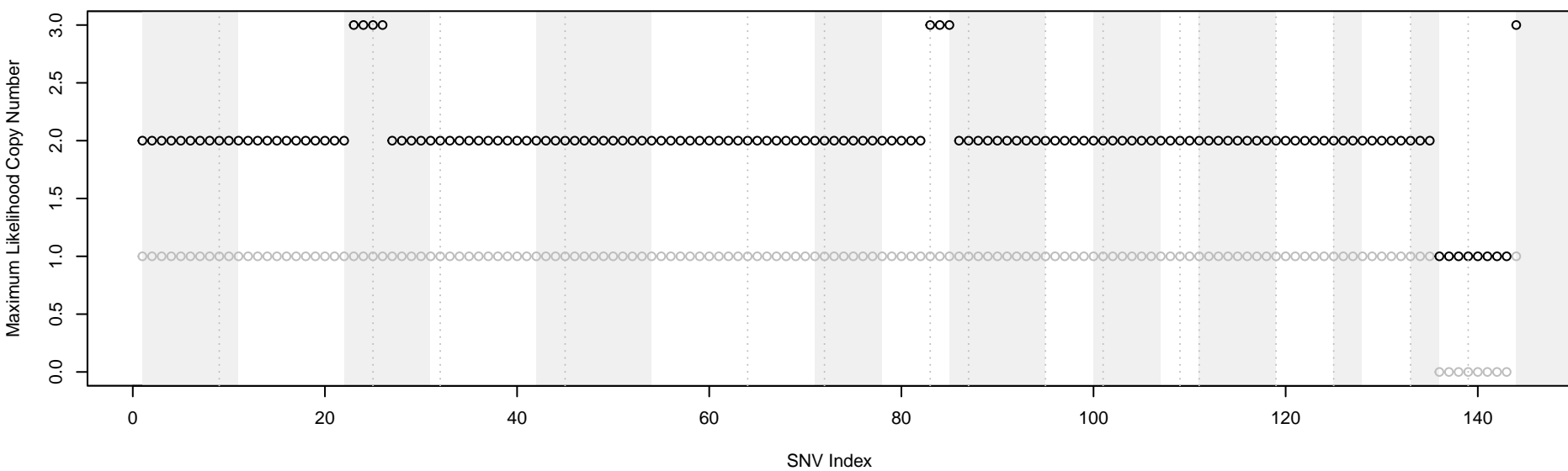
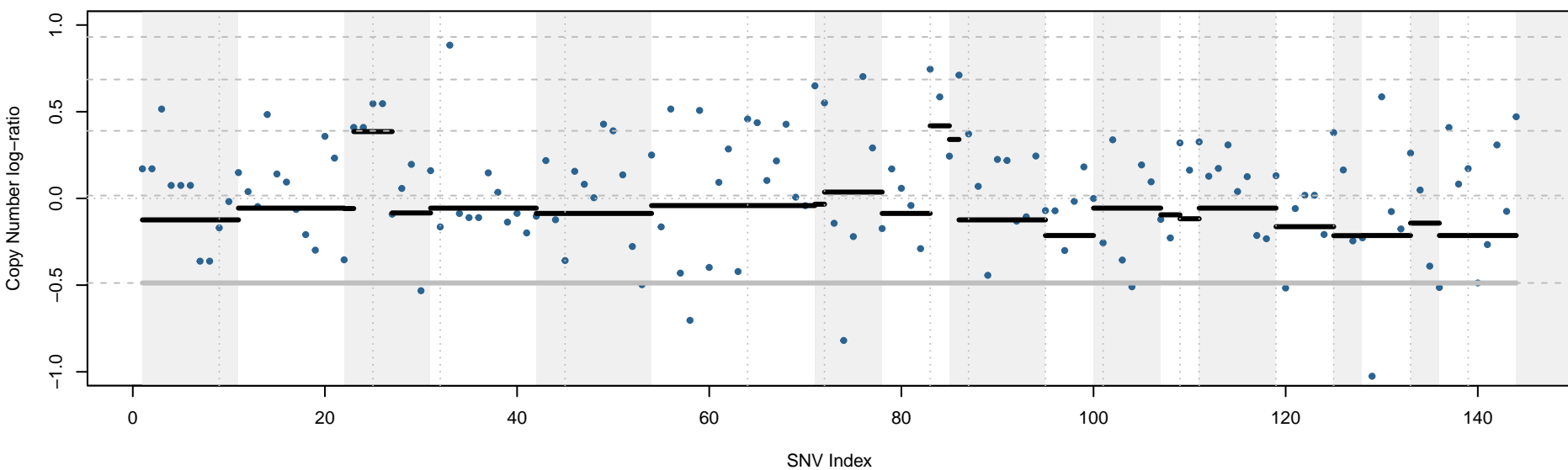
Purity: 0.59 Tumor ploidy: 1.961

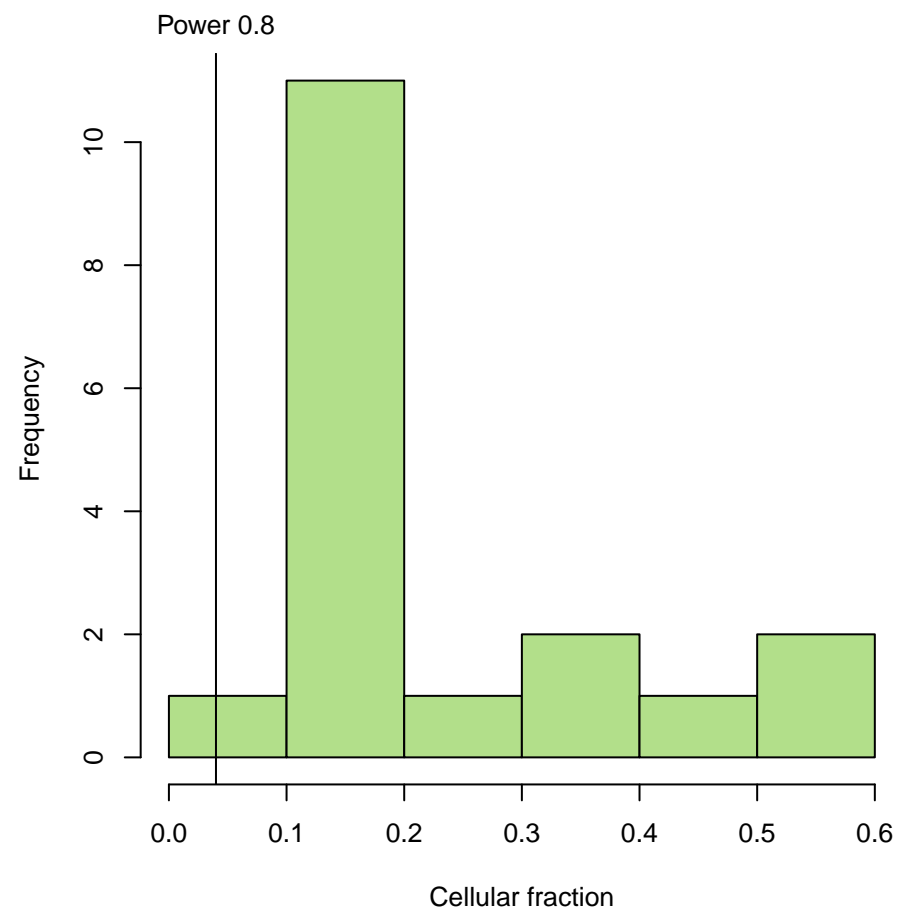
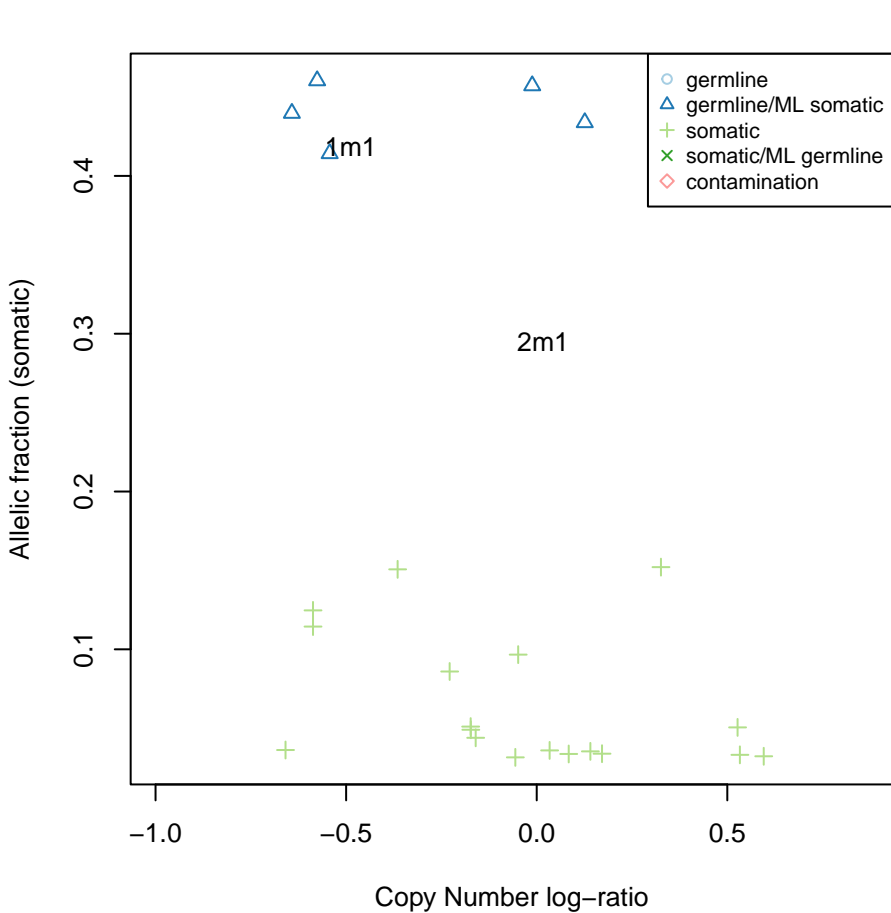
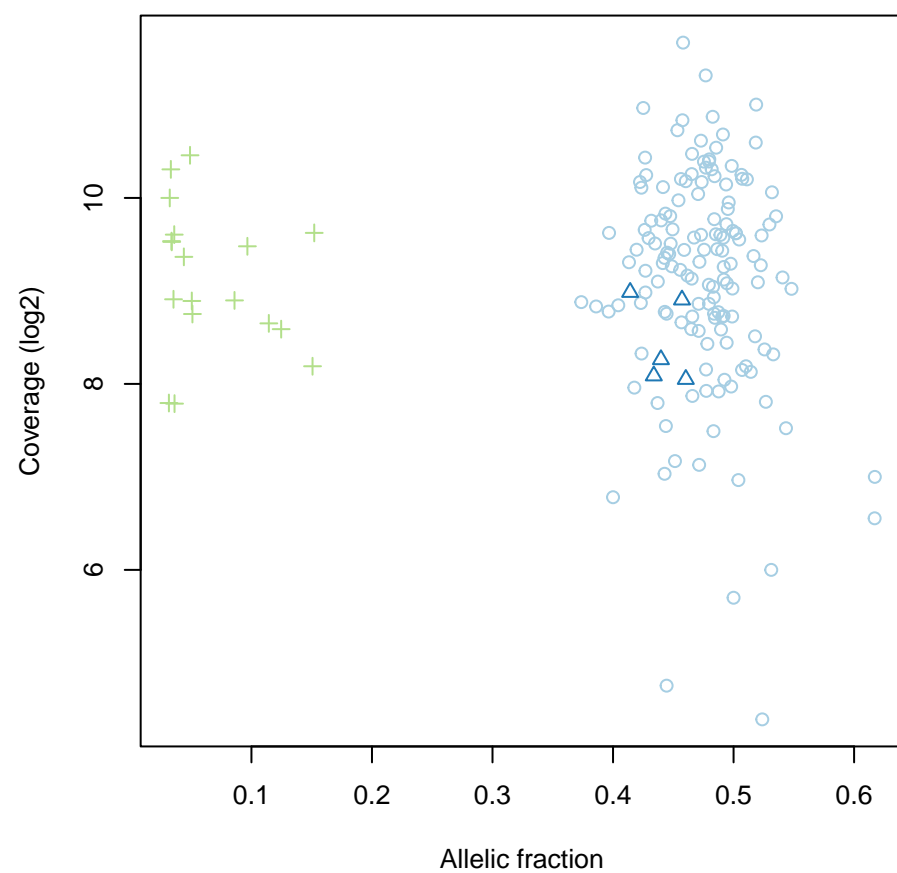
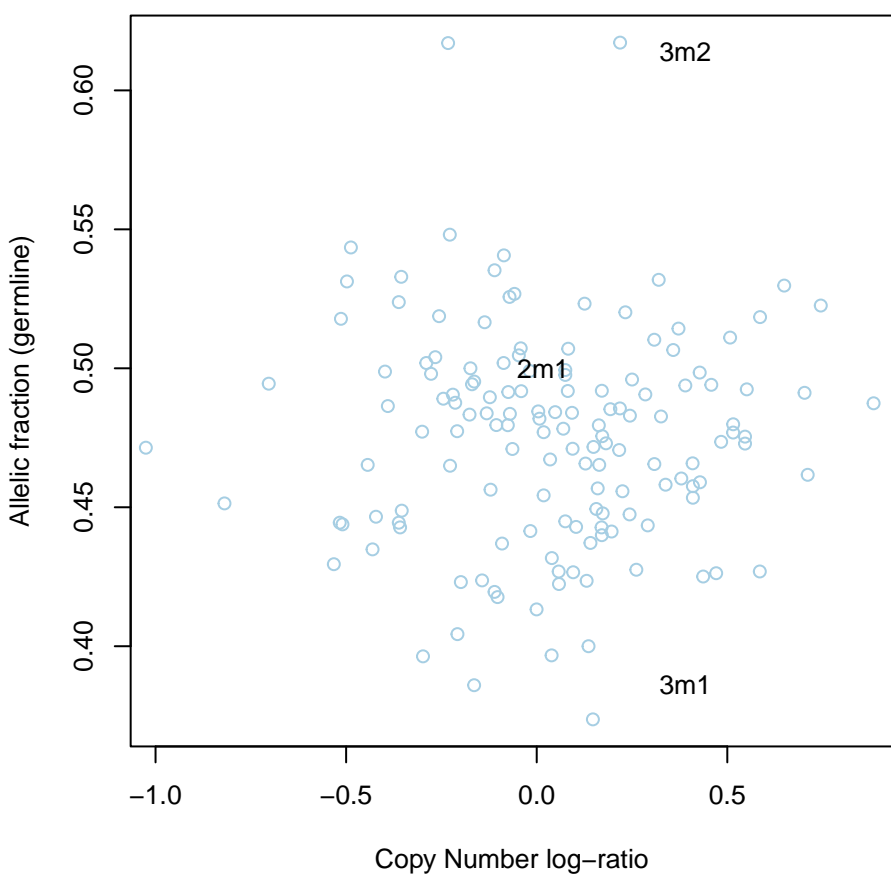


1                    2                    3                    4                    5                    6                    7                    8                    9                    10                    11                    12                    13                    14                    15                    16                    18                    19

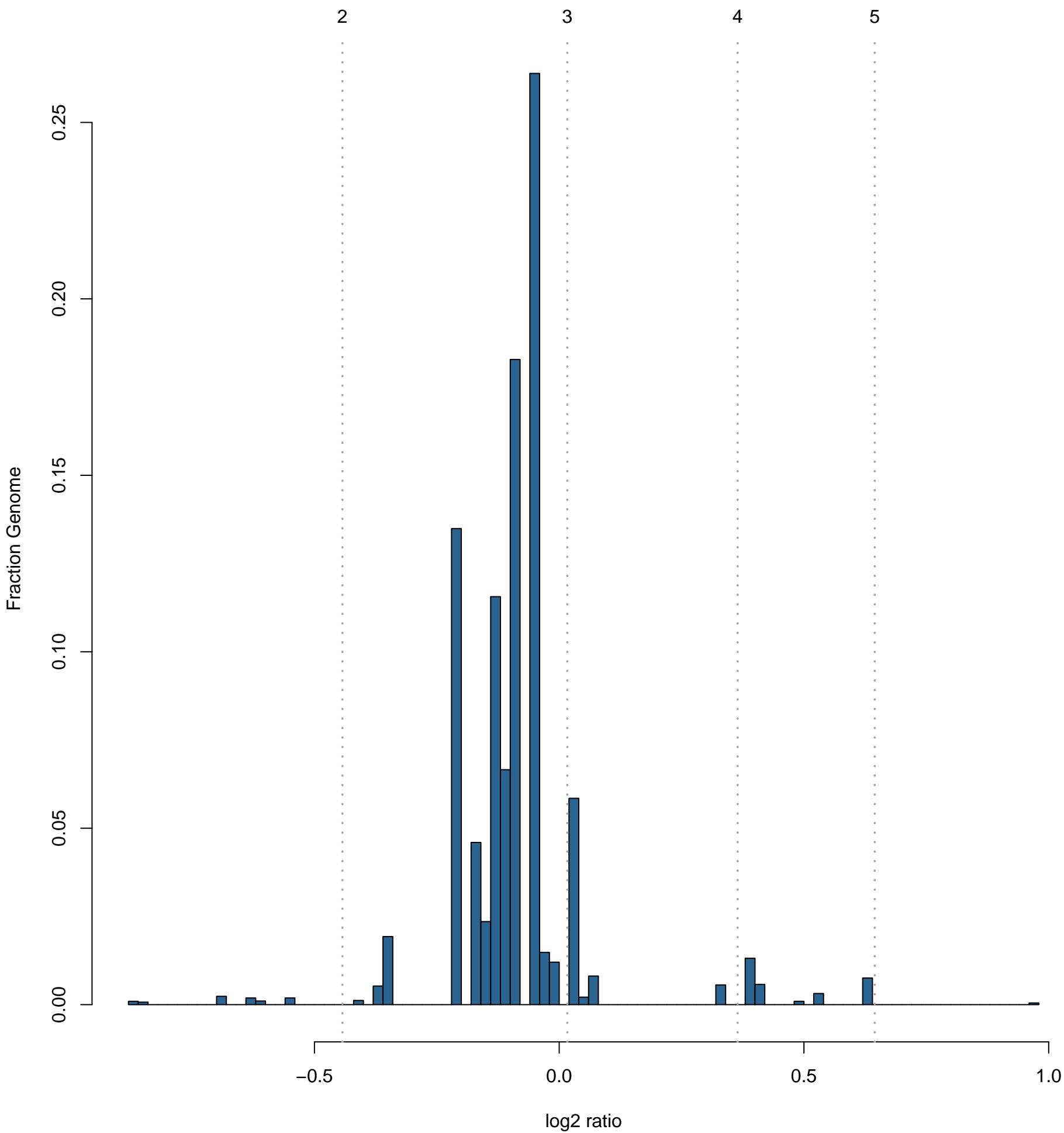


**SCNA-fit log-likelihood: -9467.28**

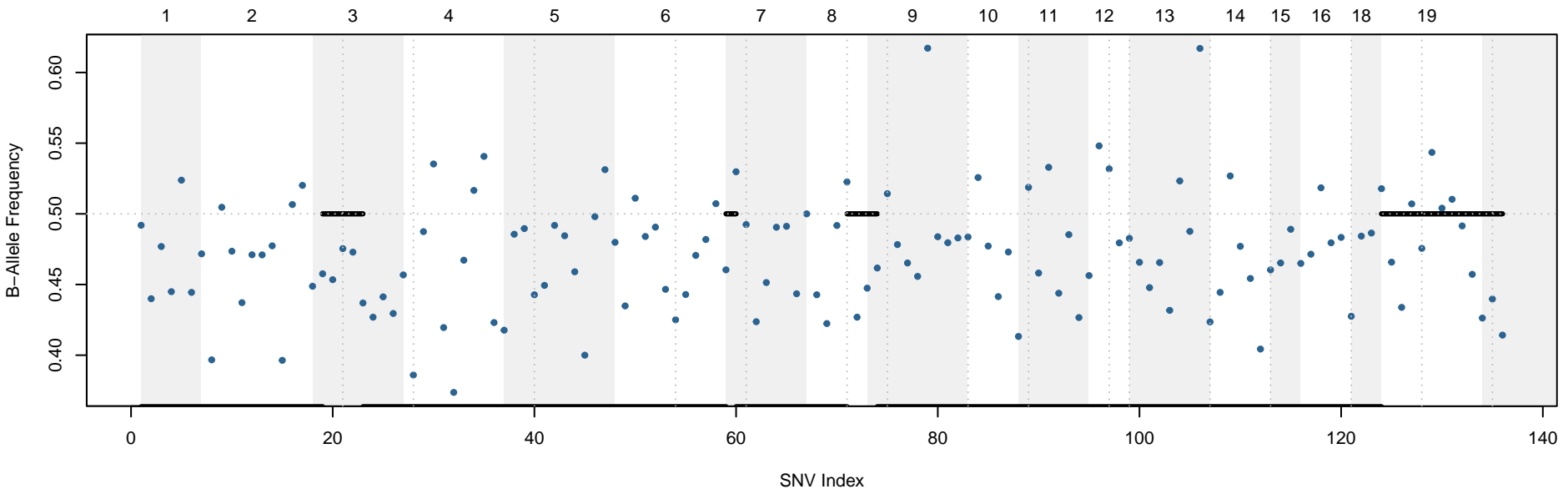




Purity: 0.75 Tumor ploidy: 2.958



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



SCNA-fit log-likelihood: -9430.41

