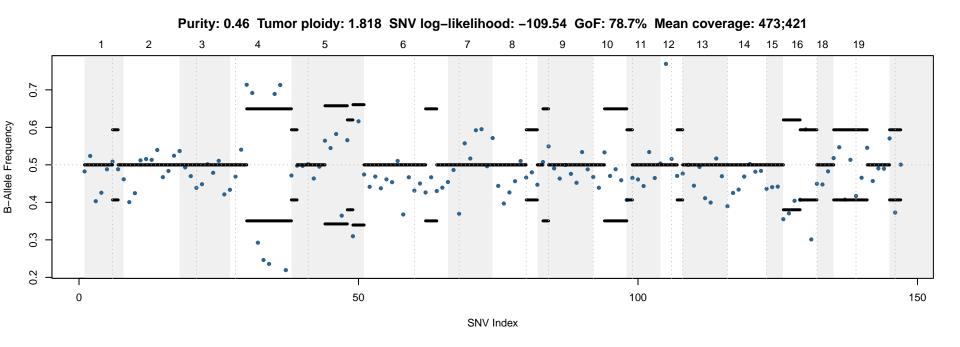
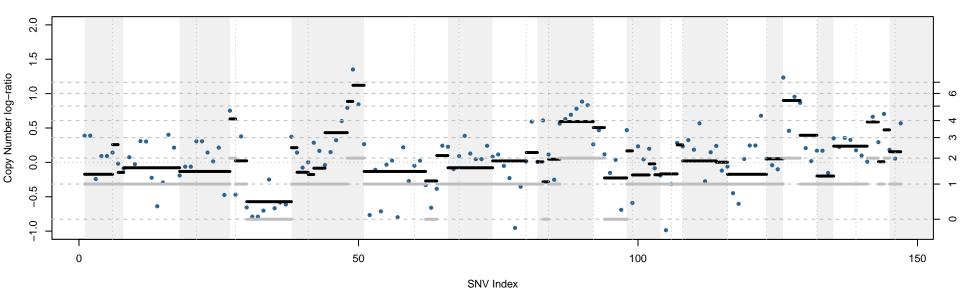
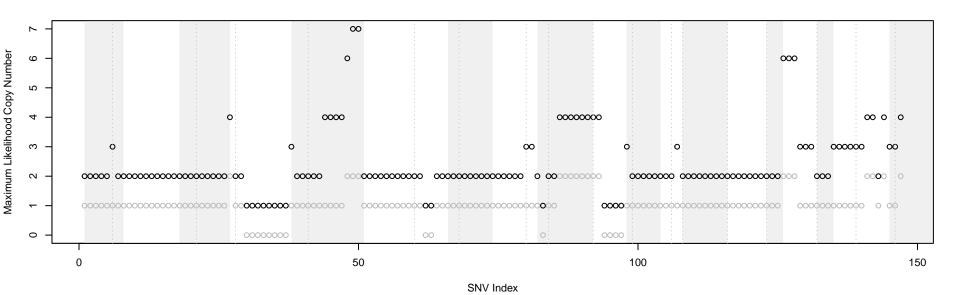
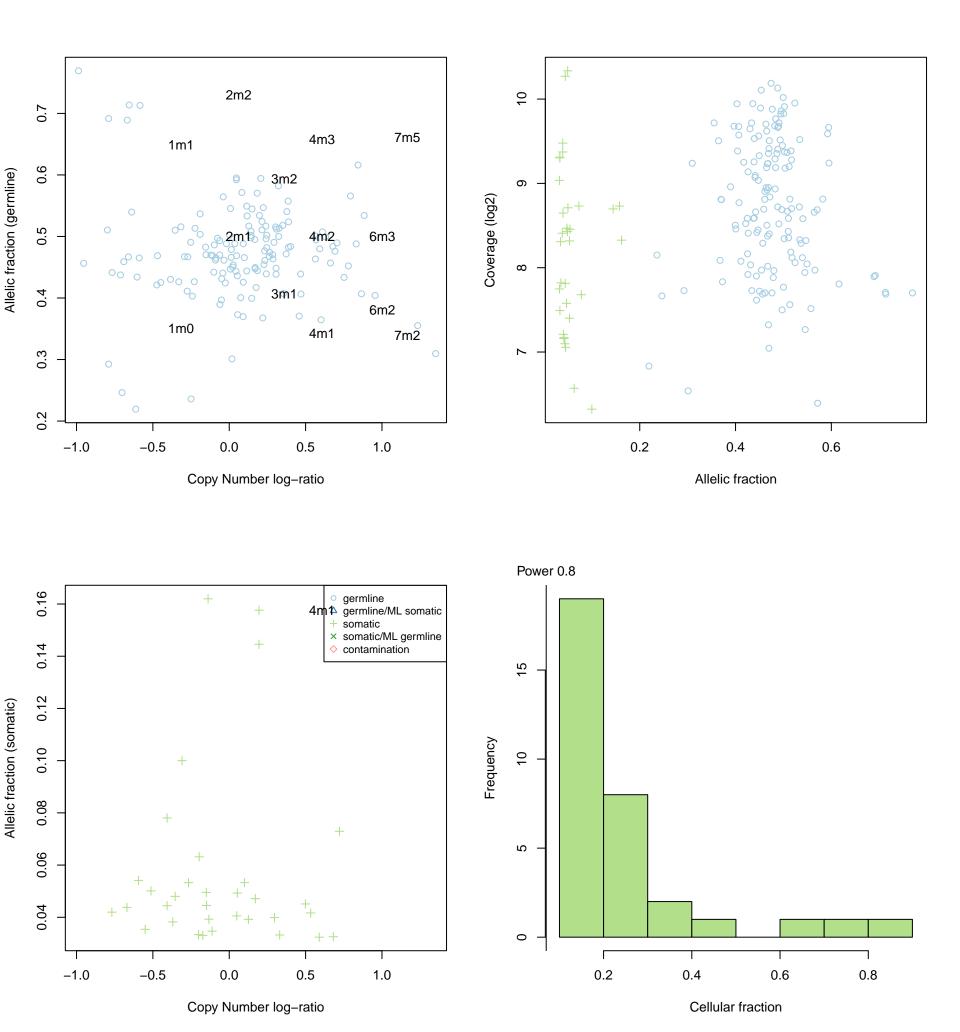
Purity: 0.46 Tumor ploidy: 1.818 2 7 0 5 6 3 Fraction Genome 0.05 0.00 -1.5 -1.0 -0.5 0.0 0.5 1.0 log2 ratio



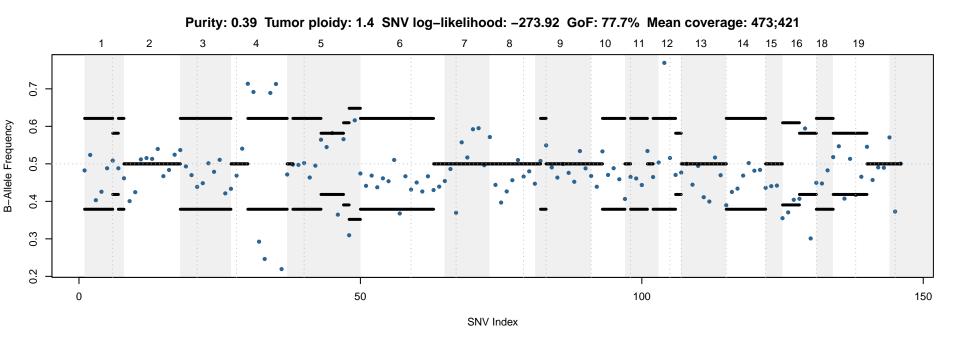
SCNA-fit log-likelihood: -3891.52







Purity: 0.39 Tumor ploidy: 1.4 2 7 3 4 6 0 5 Fraction Genome 0.05 0.00 -1.0 -0.5 0.0 0.5 1.0 log2 ratio



SCNA-fit log-likelihood: -4098.14

