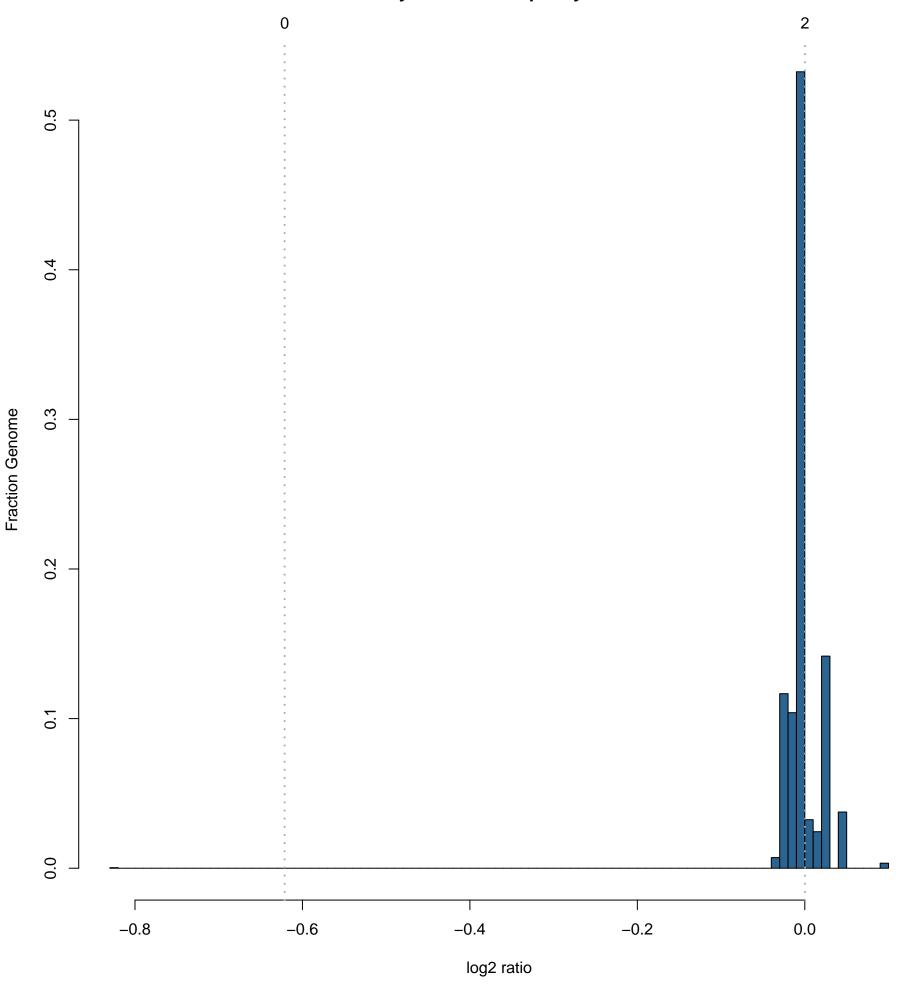
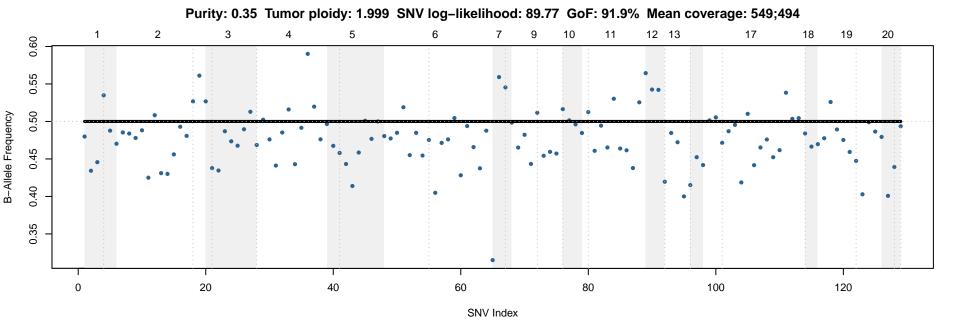
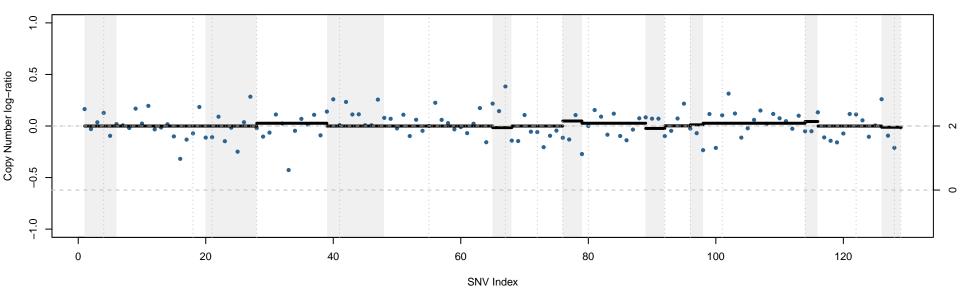
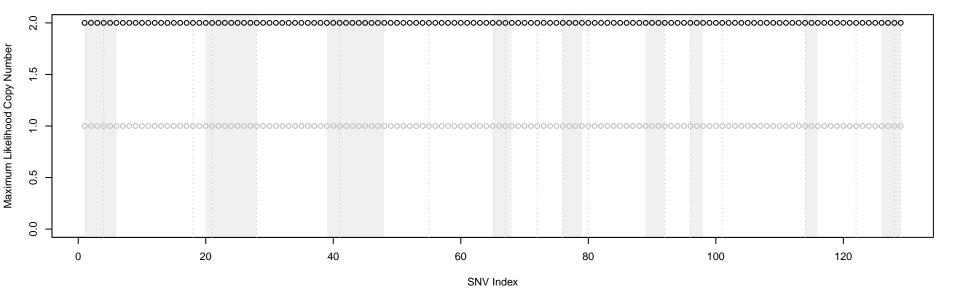
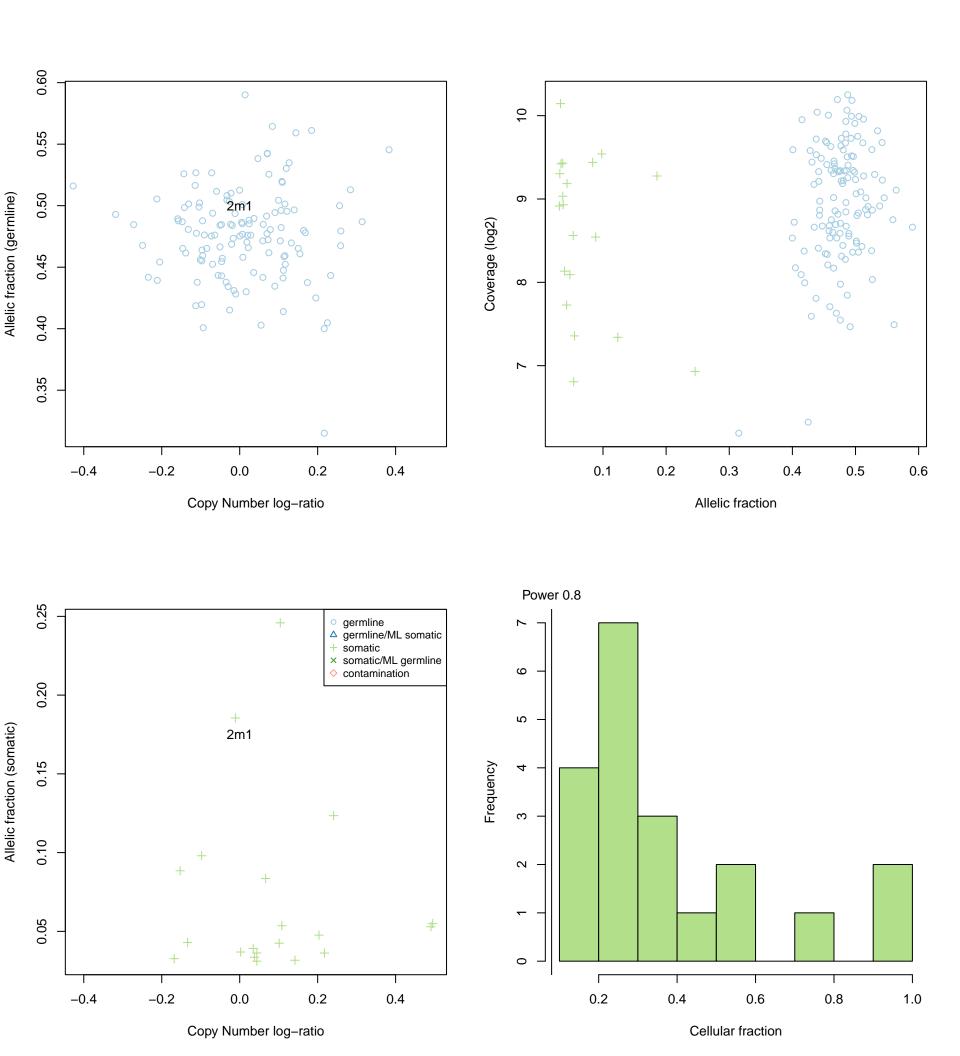
Purity: 0.35 Tumor ploidy: 1.999



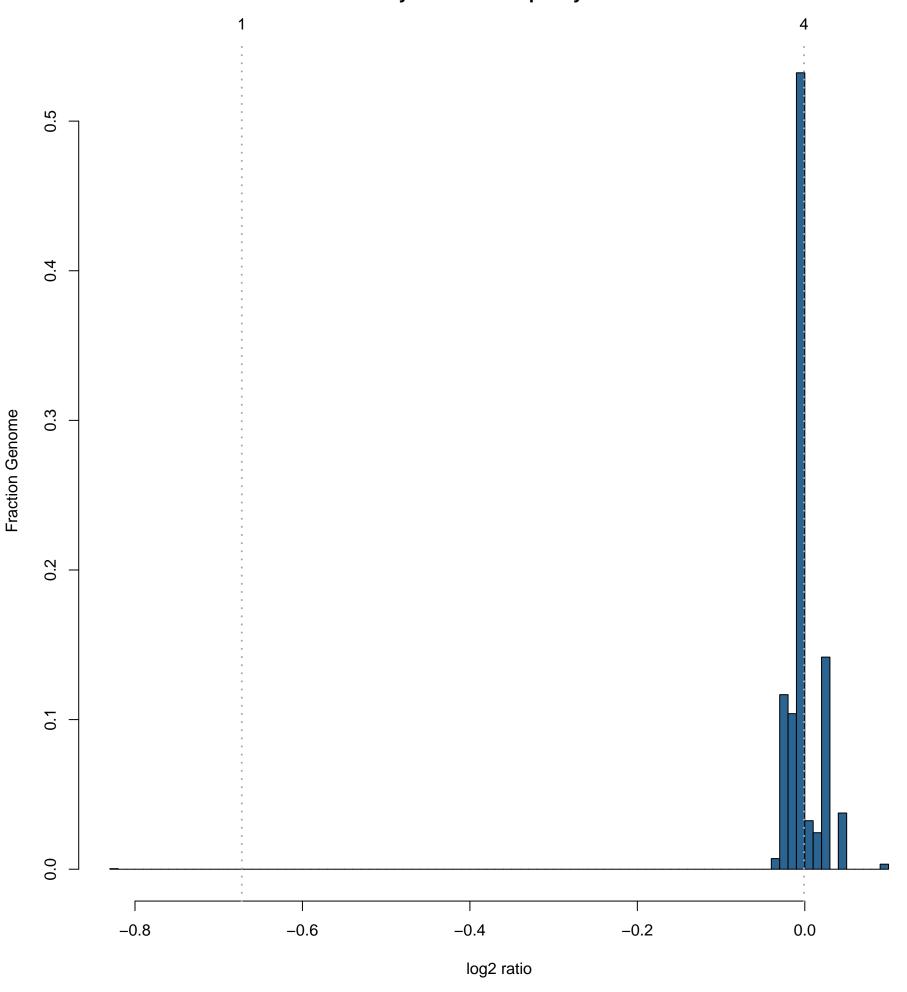


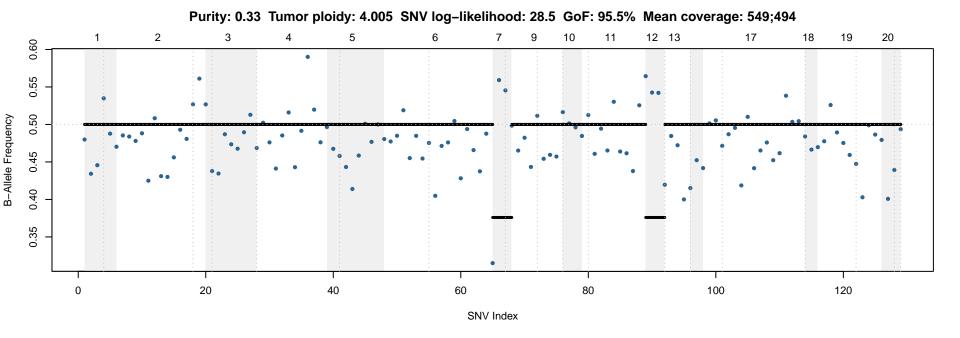


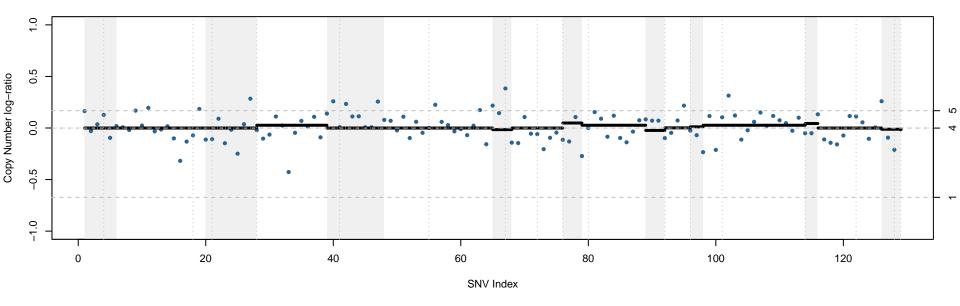


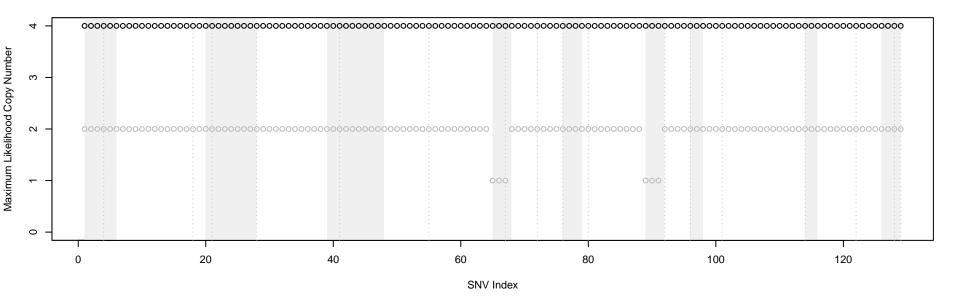


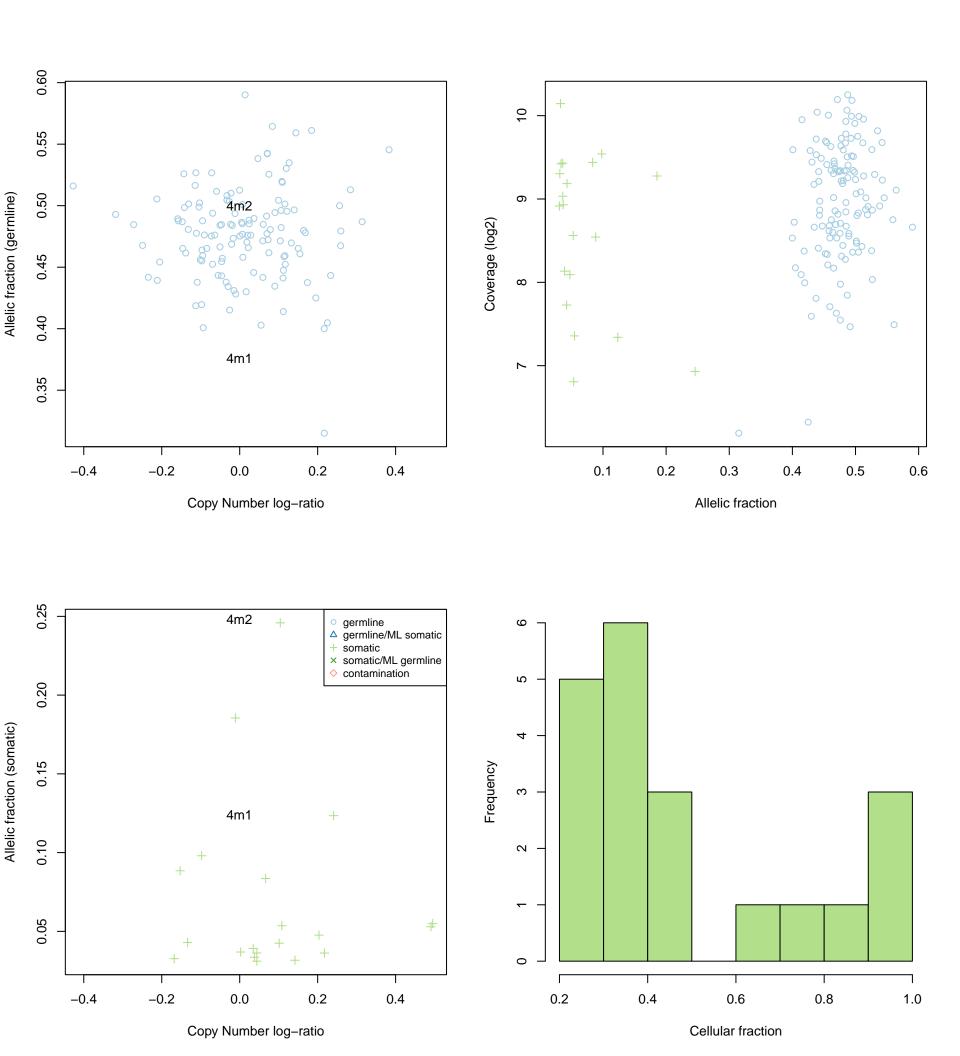
Purity: 0.33 Tumor ploidy: 4.005



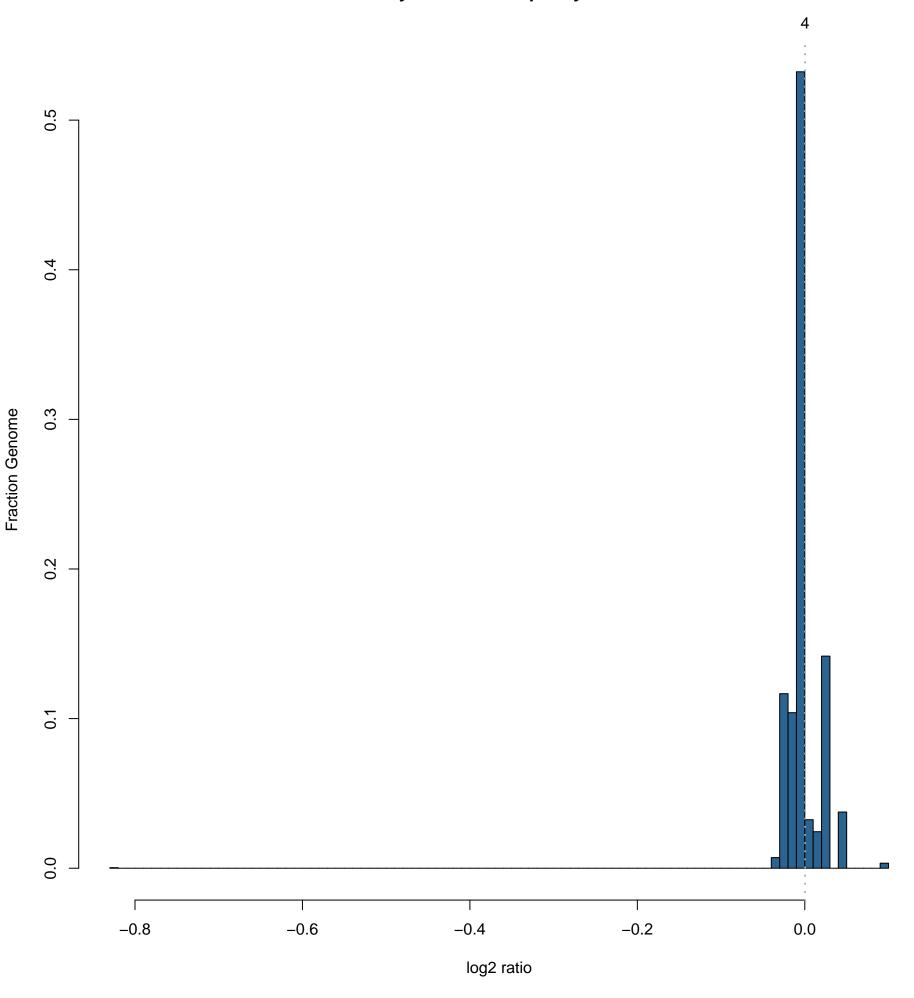


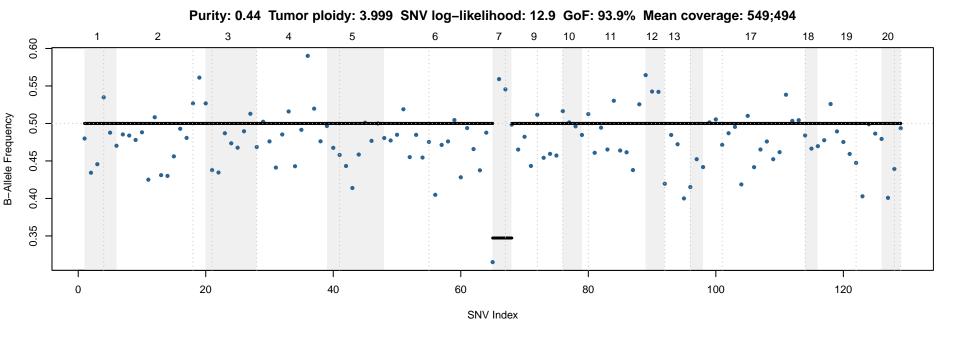


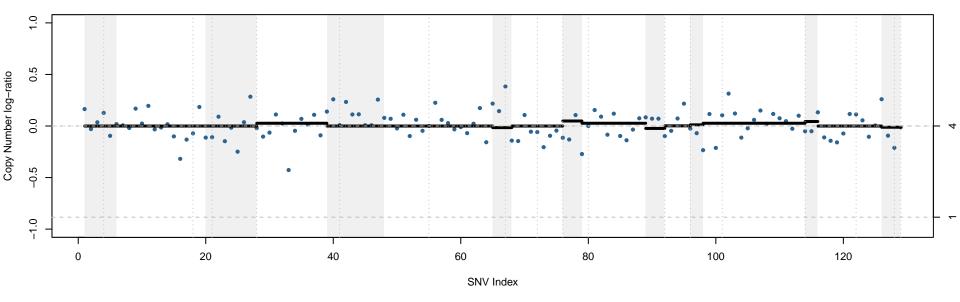


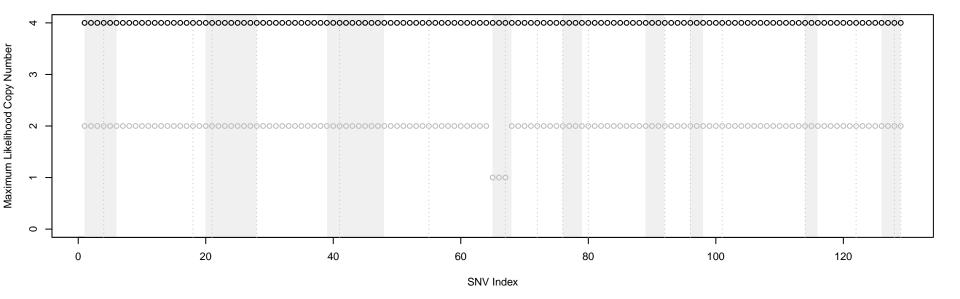


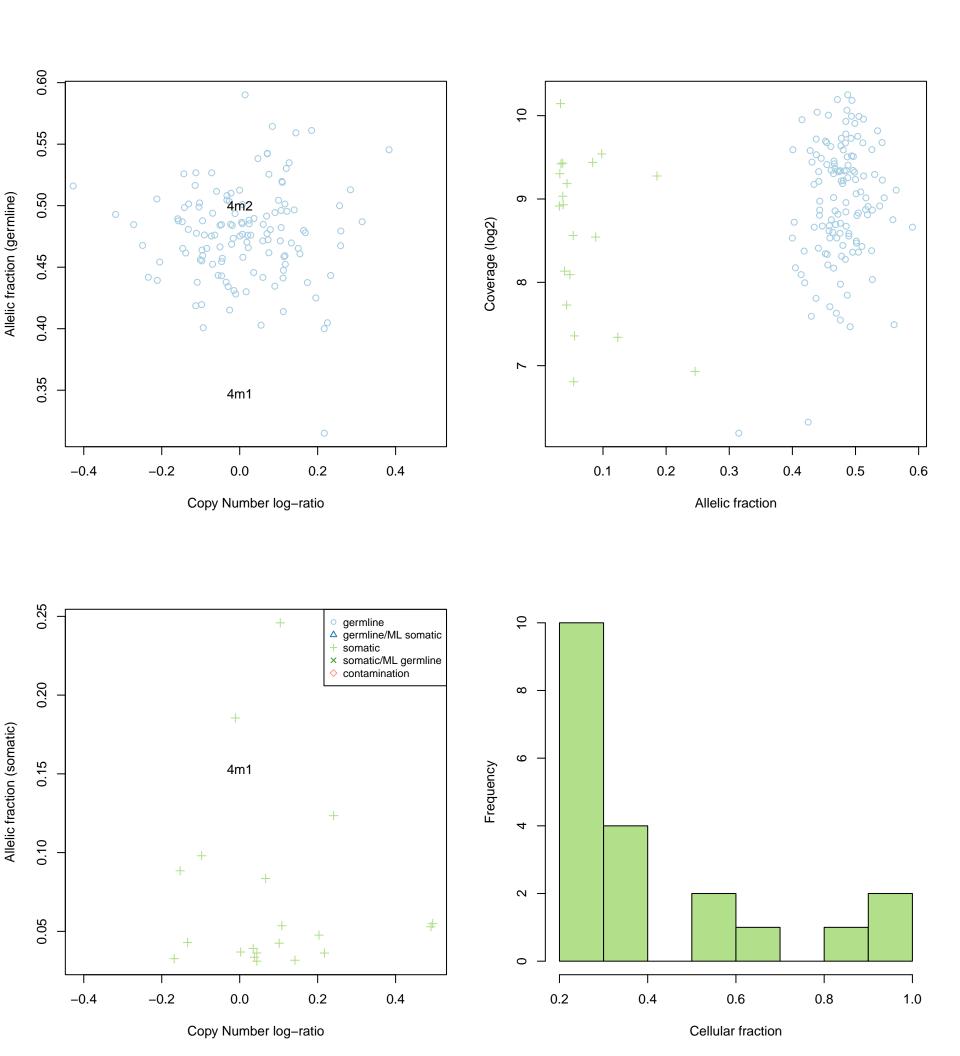
Purity: 0.44 Tumor ploidy: 3.999



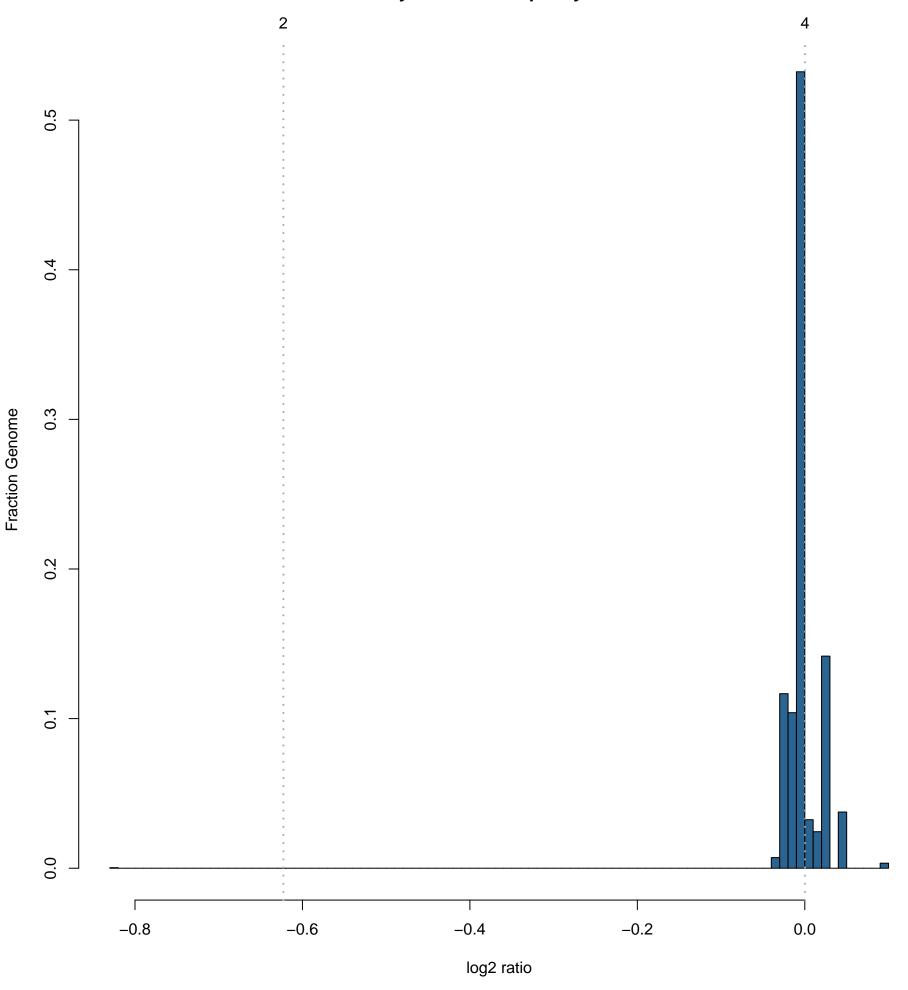


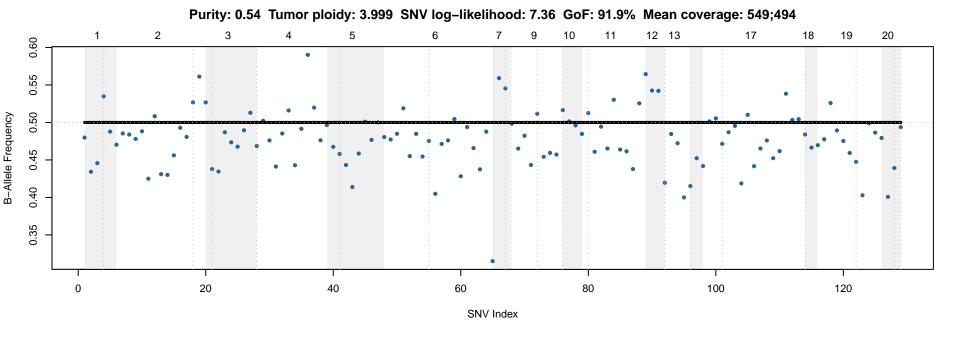


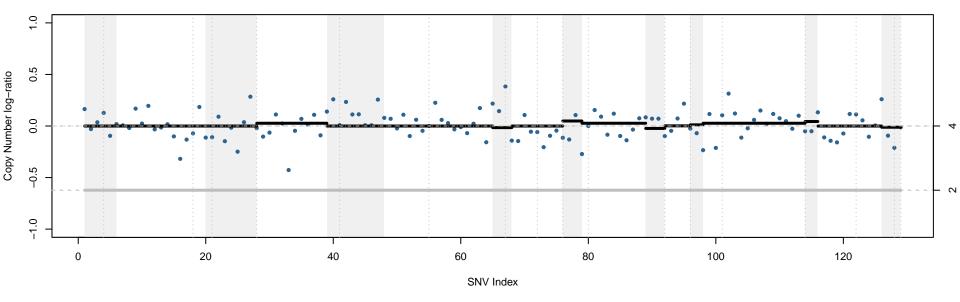


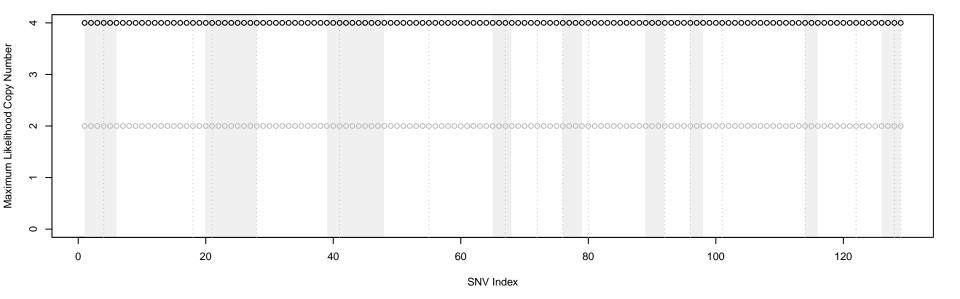


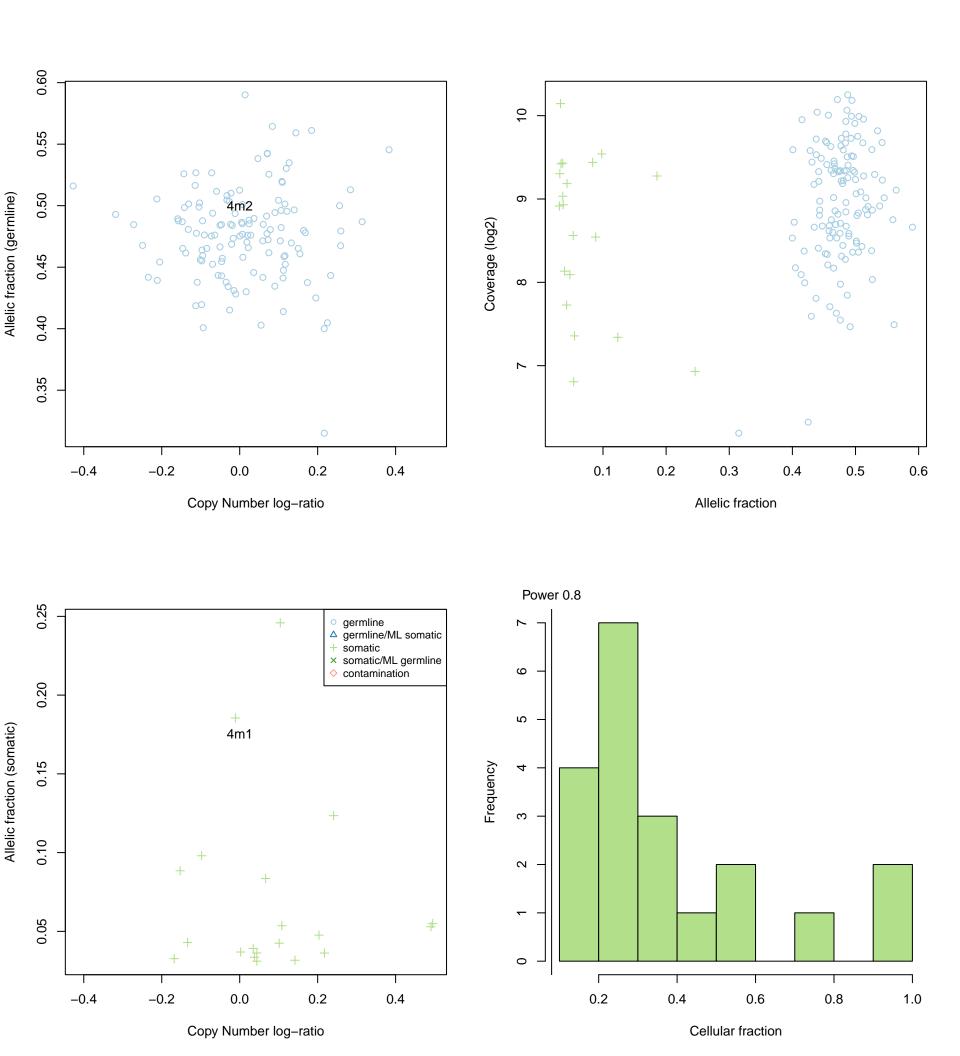
Purity: 0.54 Tumor ploidy: 3.999



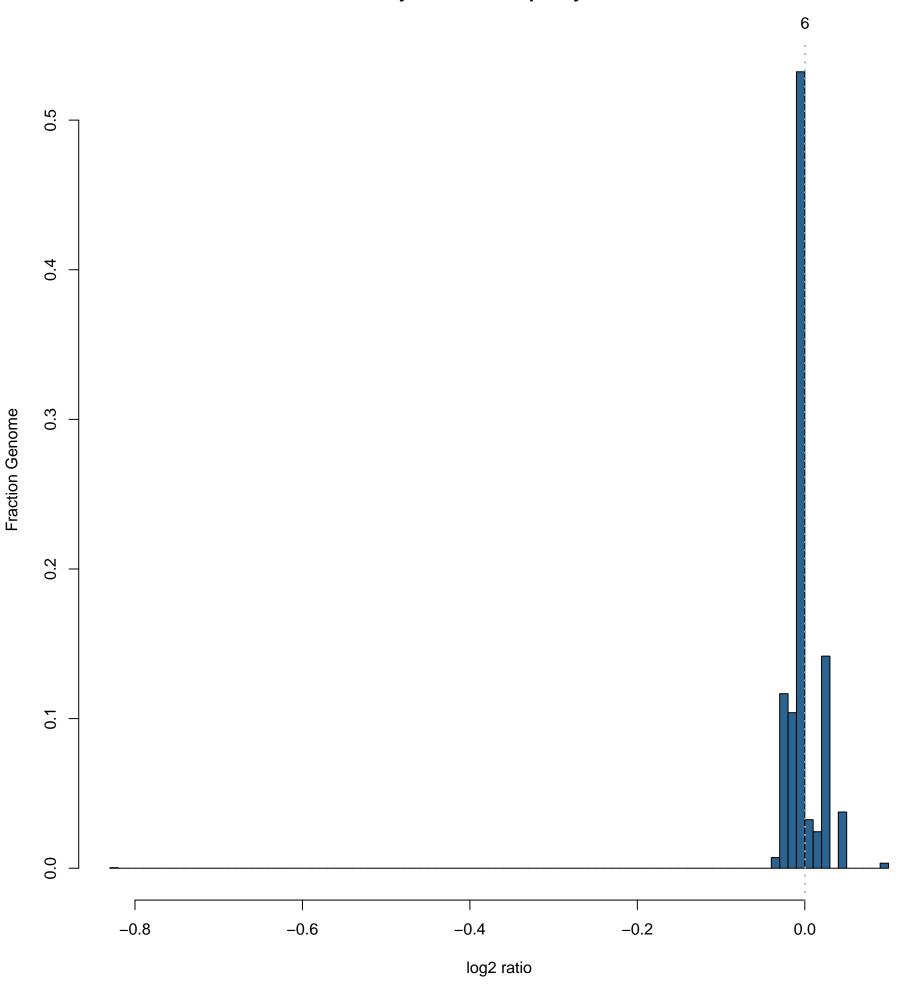


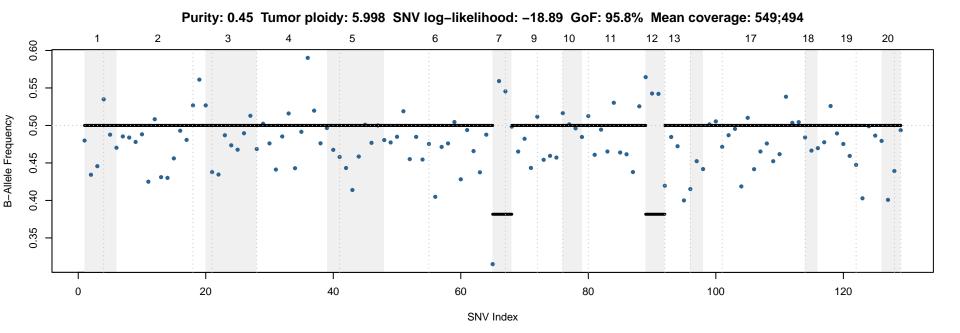




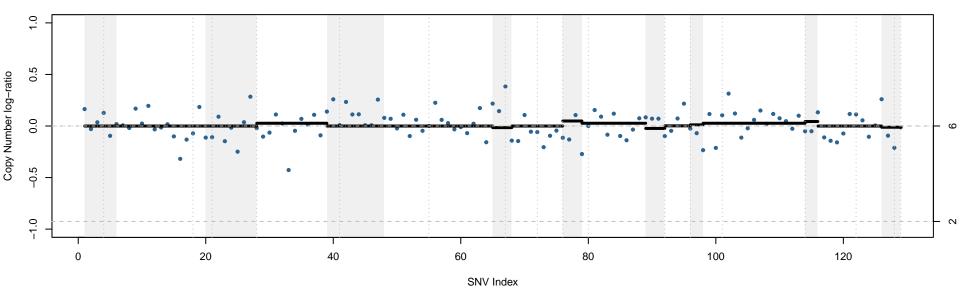


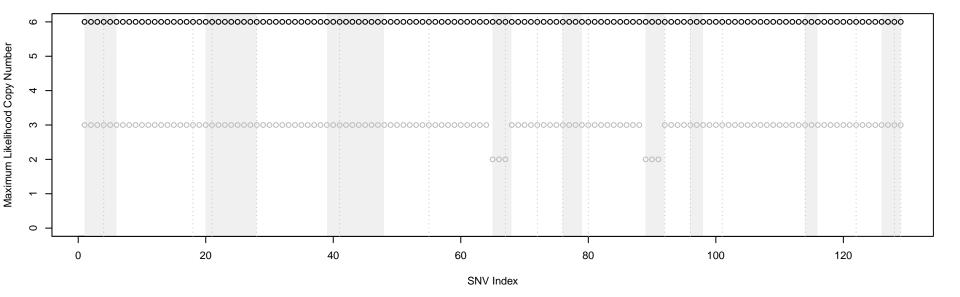
Purity: 0.45 Tumor ploidy: 5.998

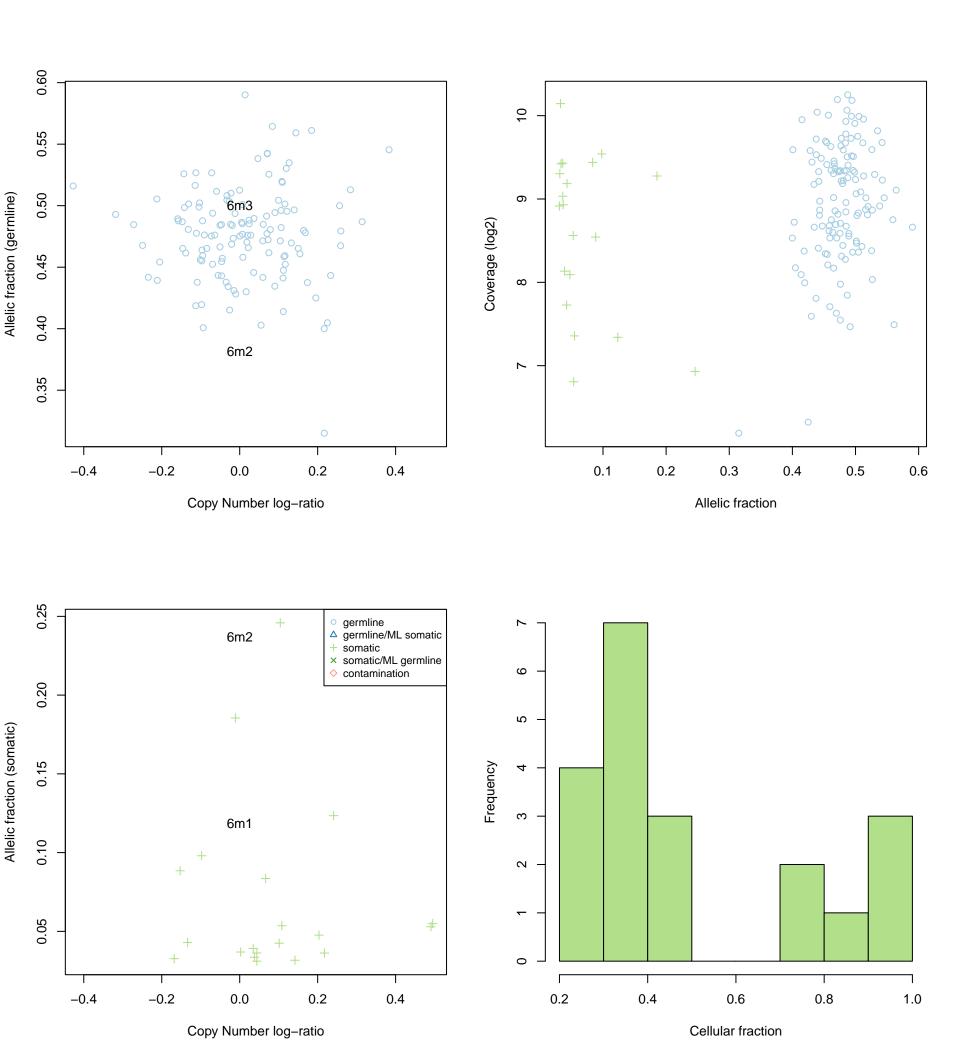




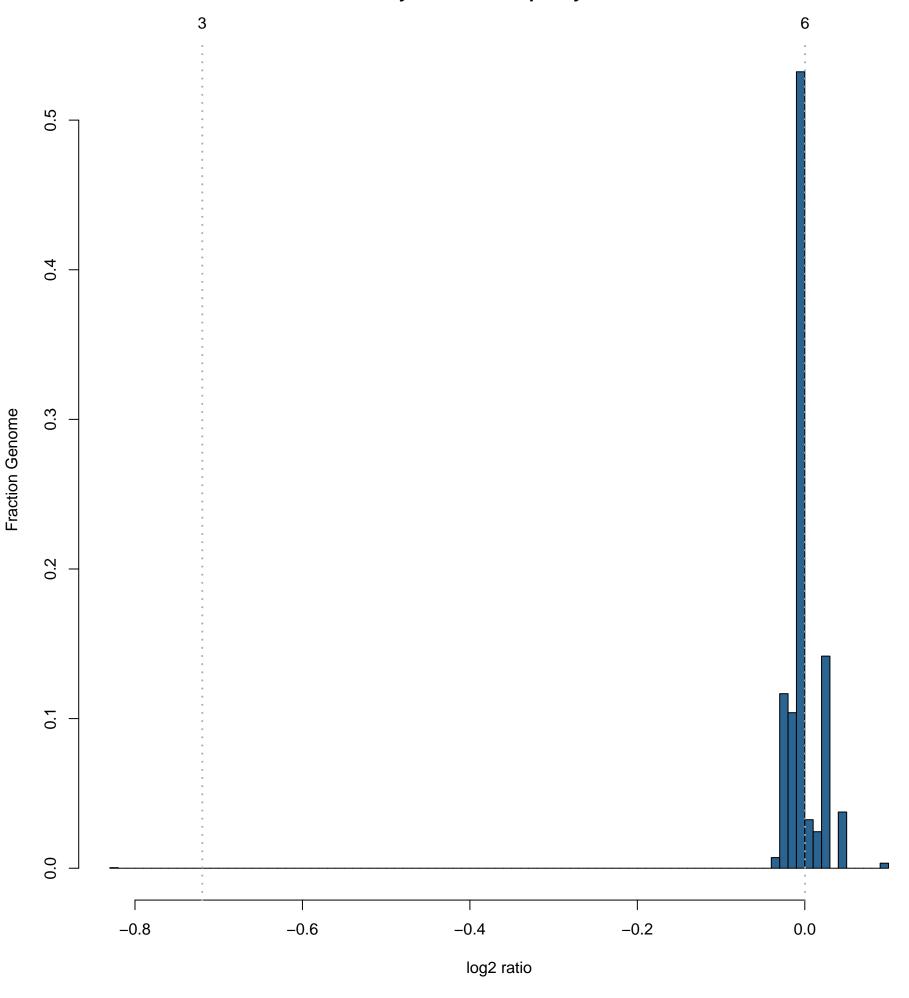
SCNA-fit log-likelihood: -4217.24

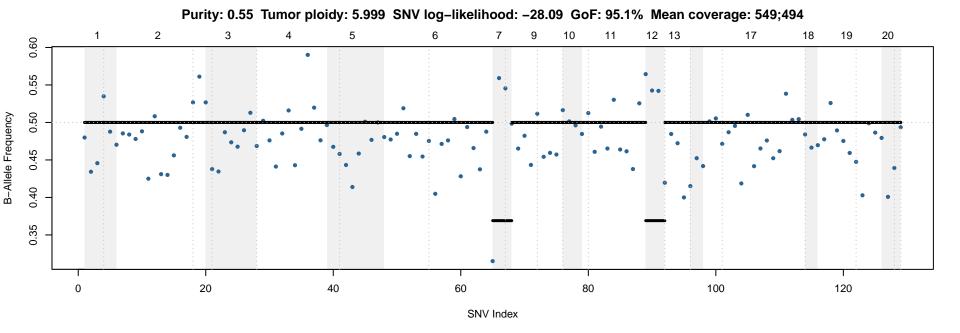


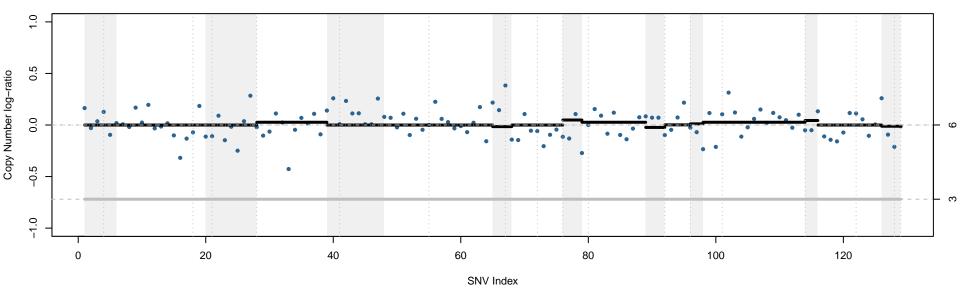


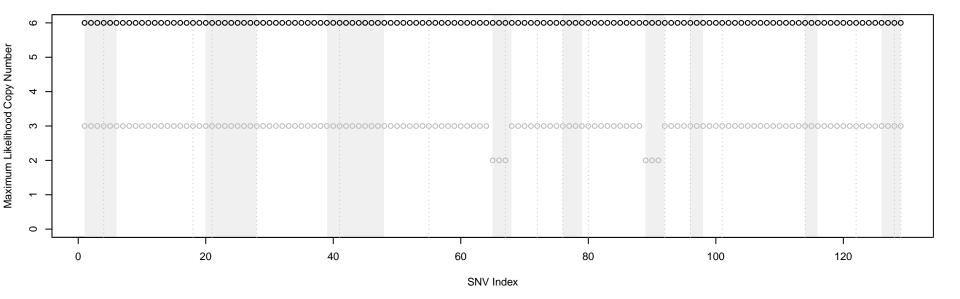


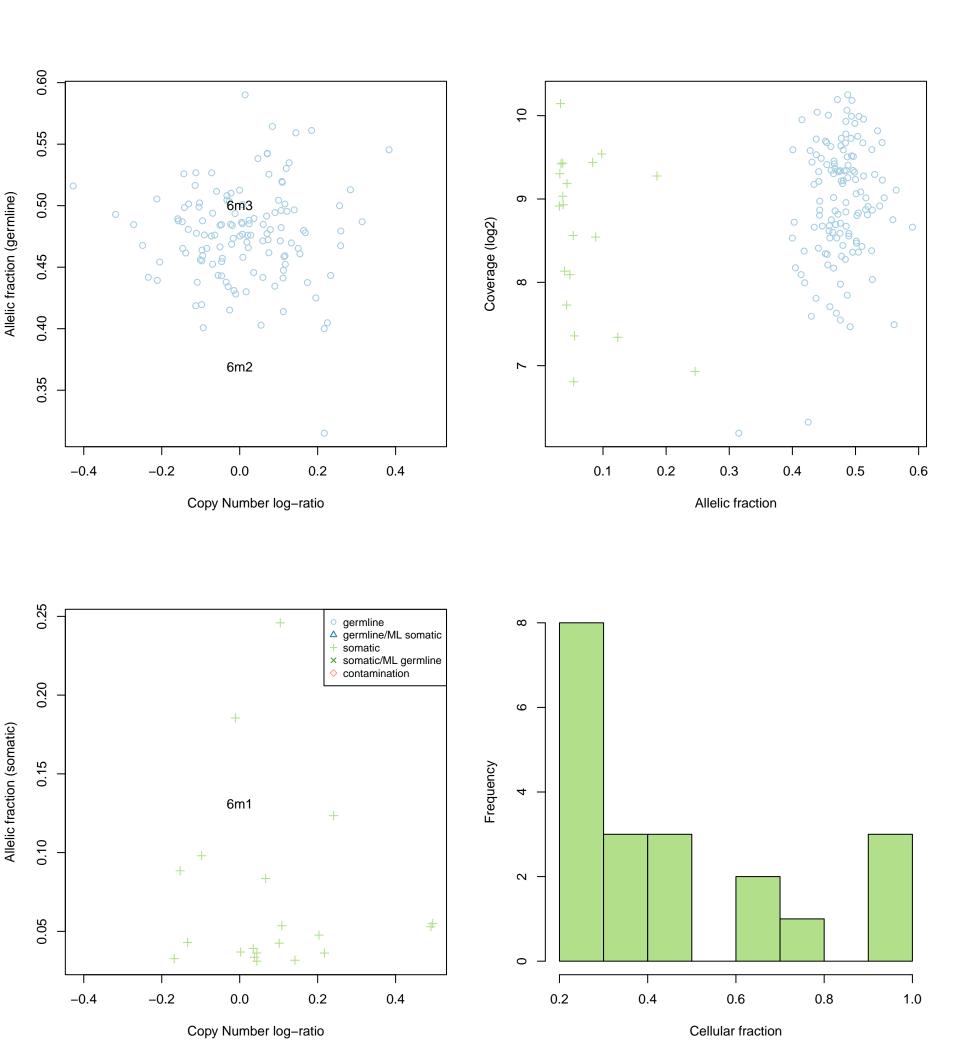
Purity: 0.55 Tumor ploidy: 5.999



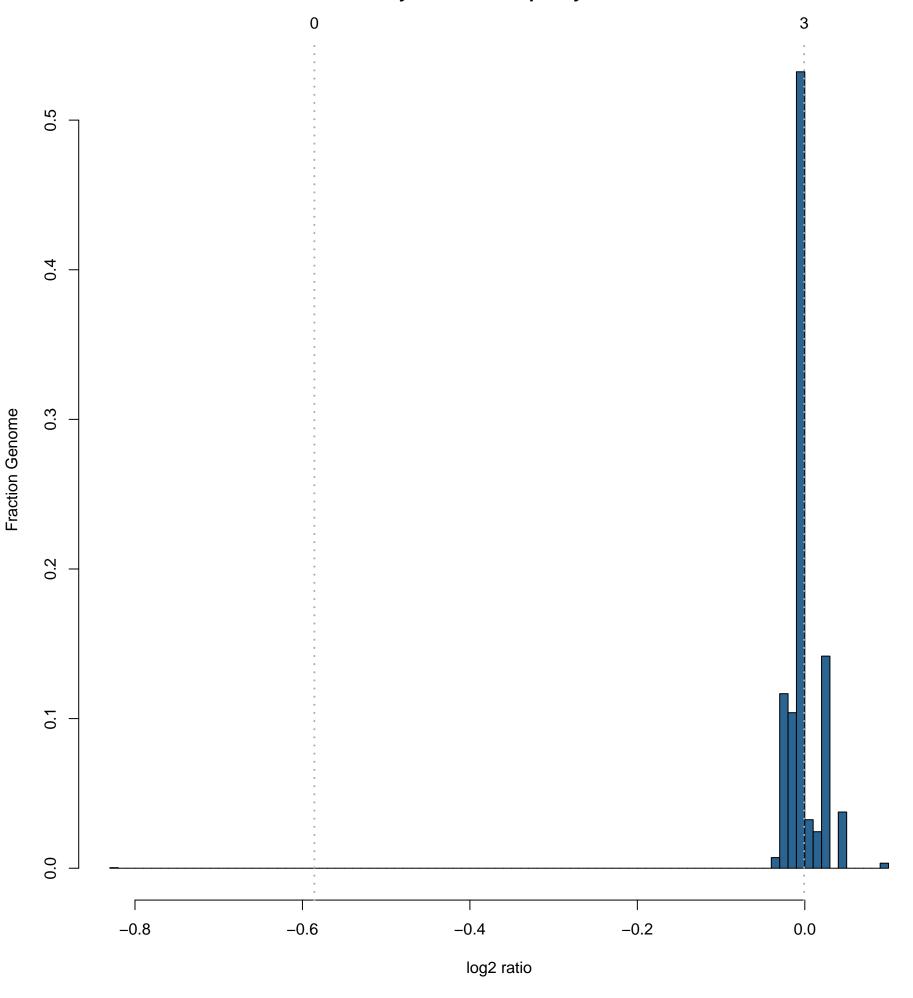


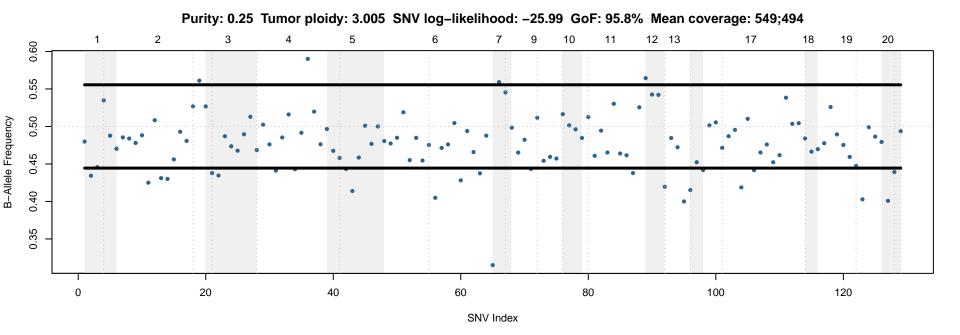


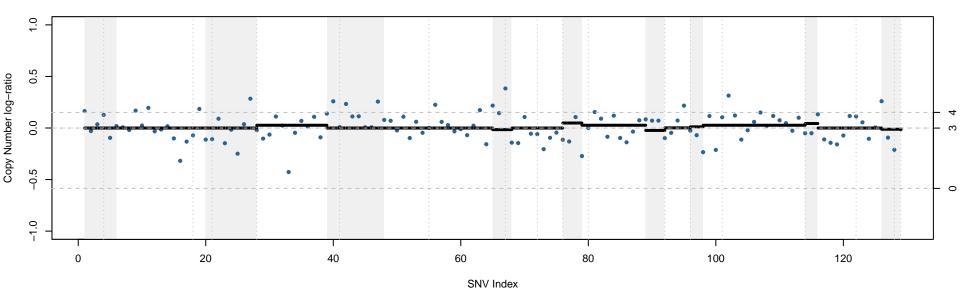


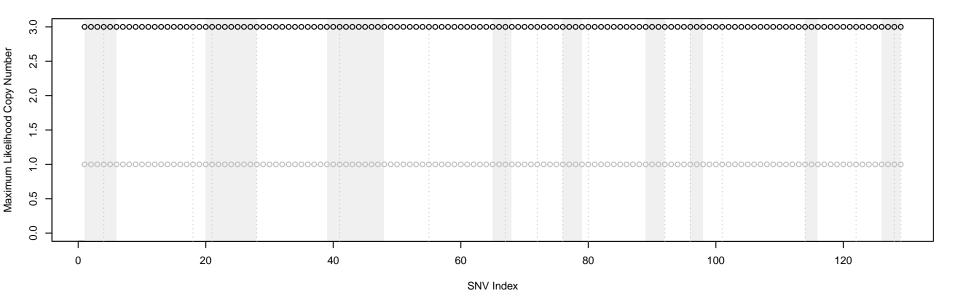


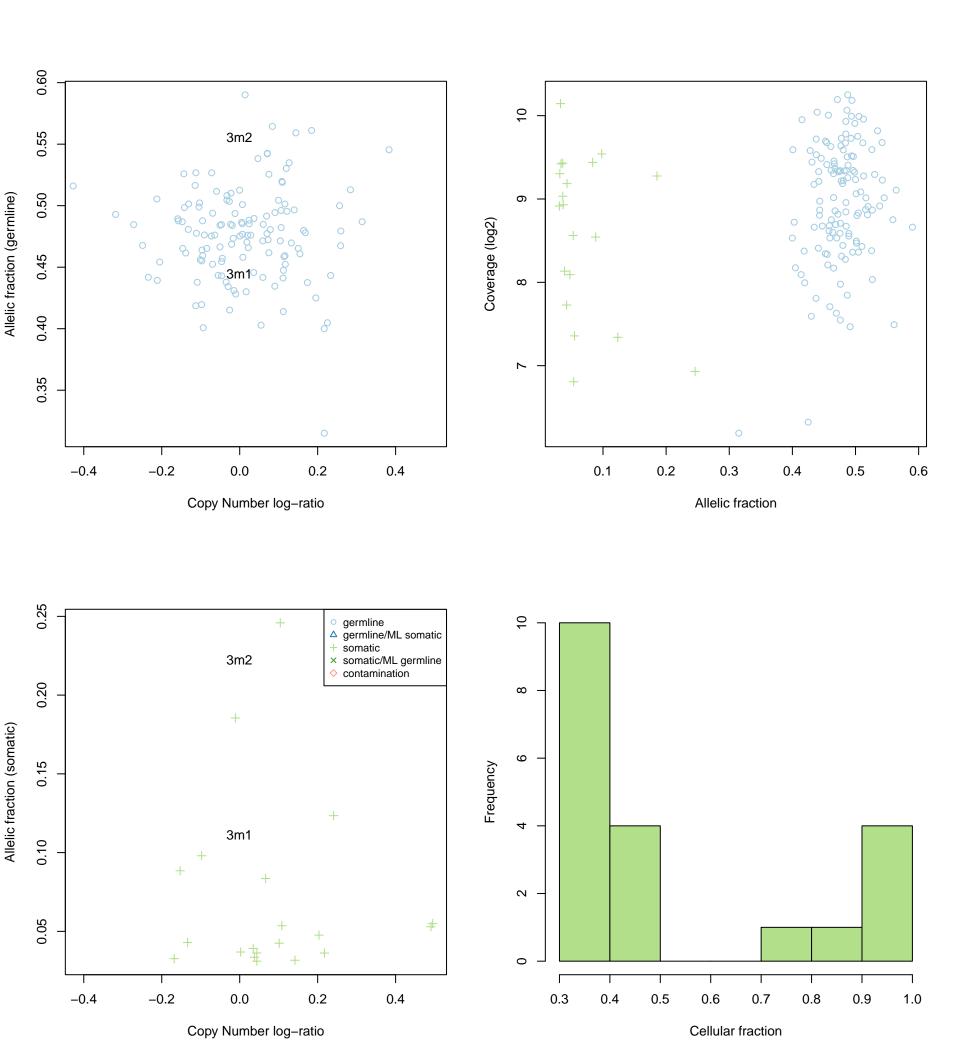
Purity: 0.25 Tumor ploidy: 3.005



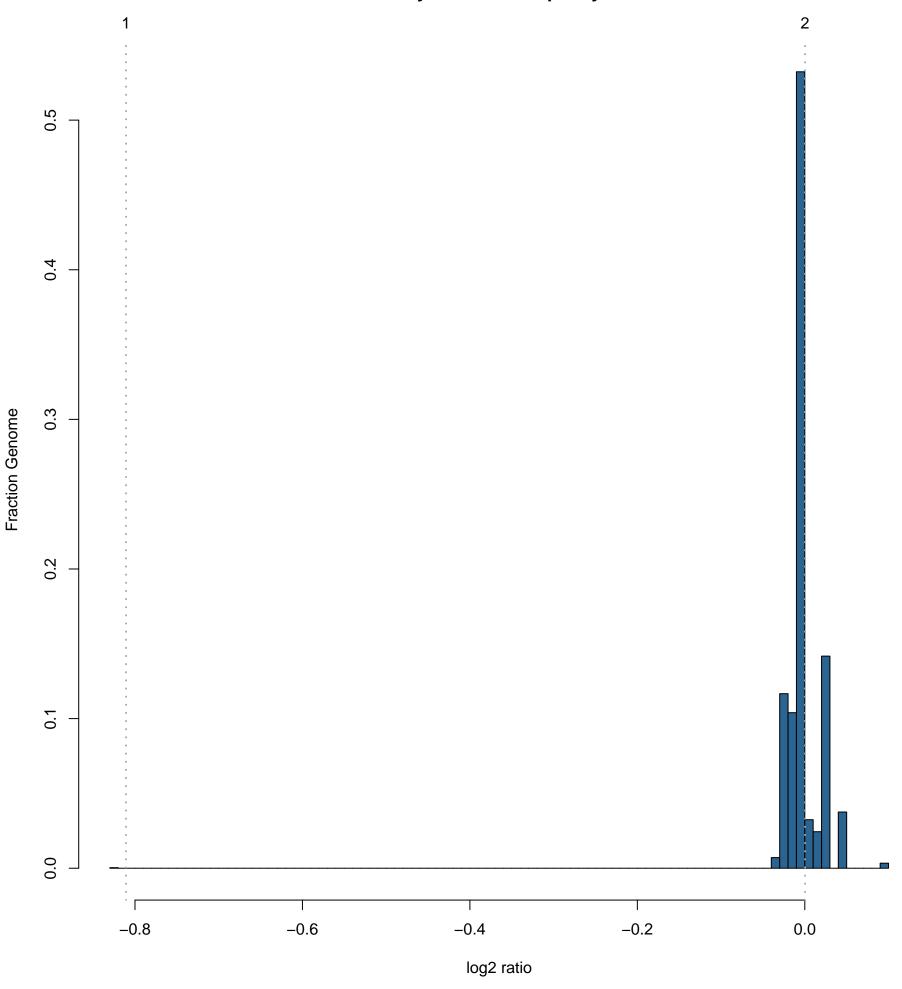


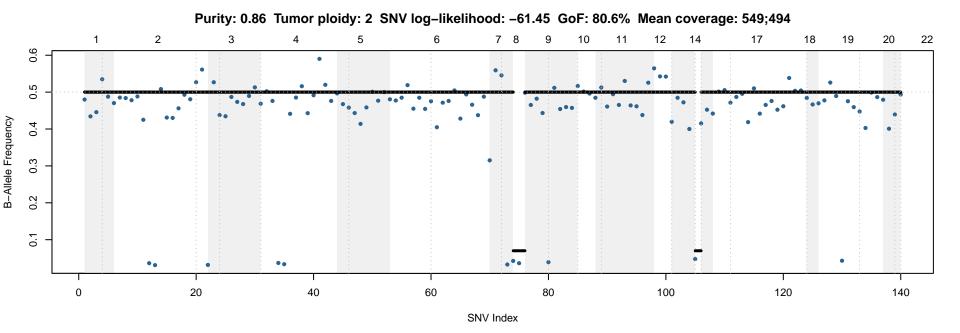


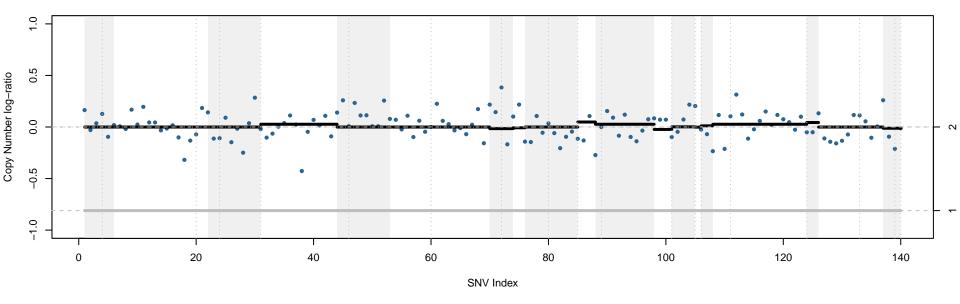


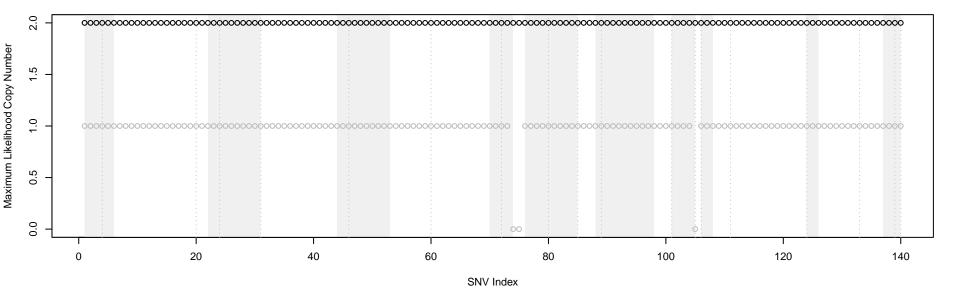


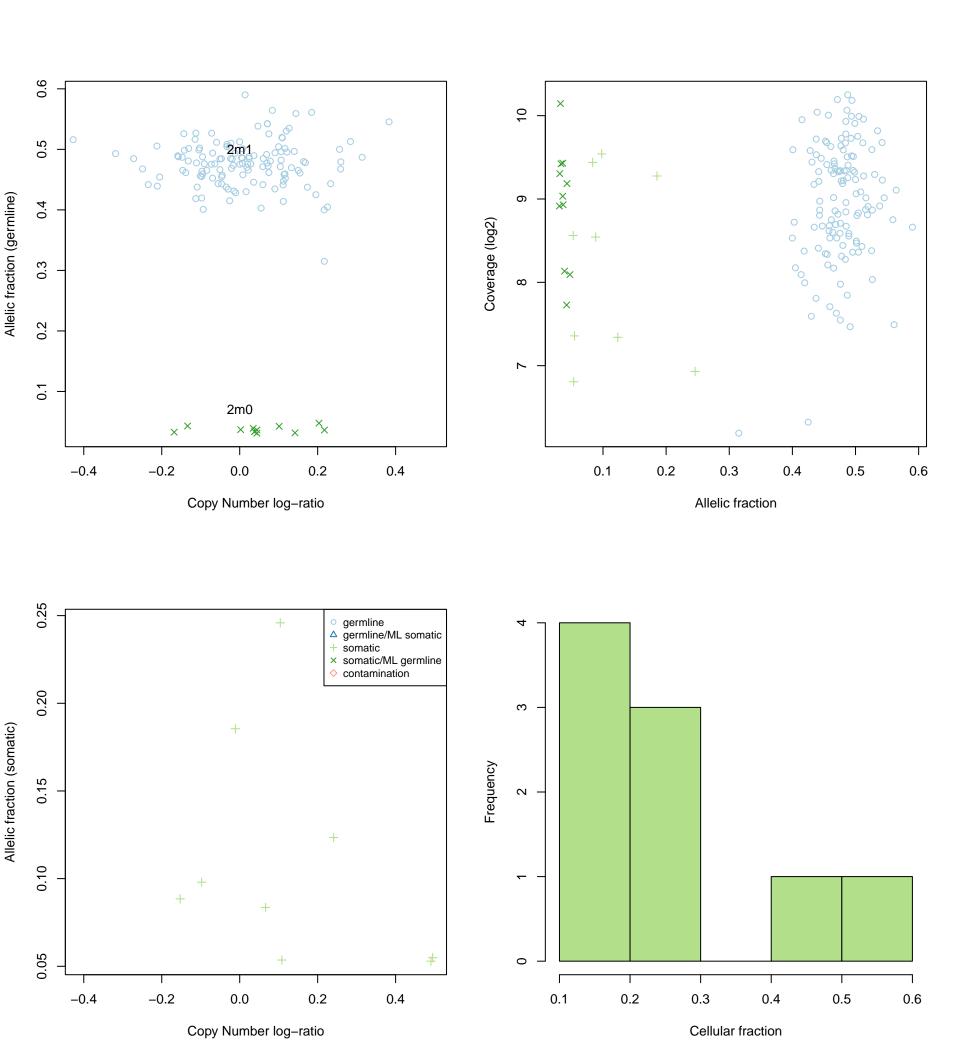
Purity: 0.86 Tumor ploidy: 2



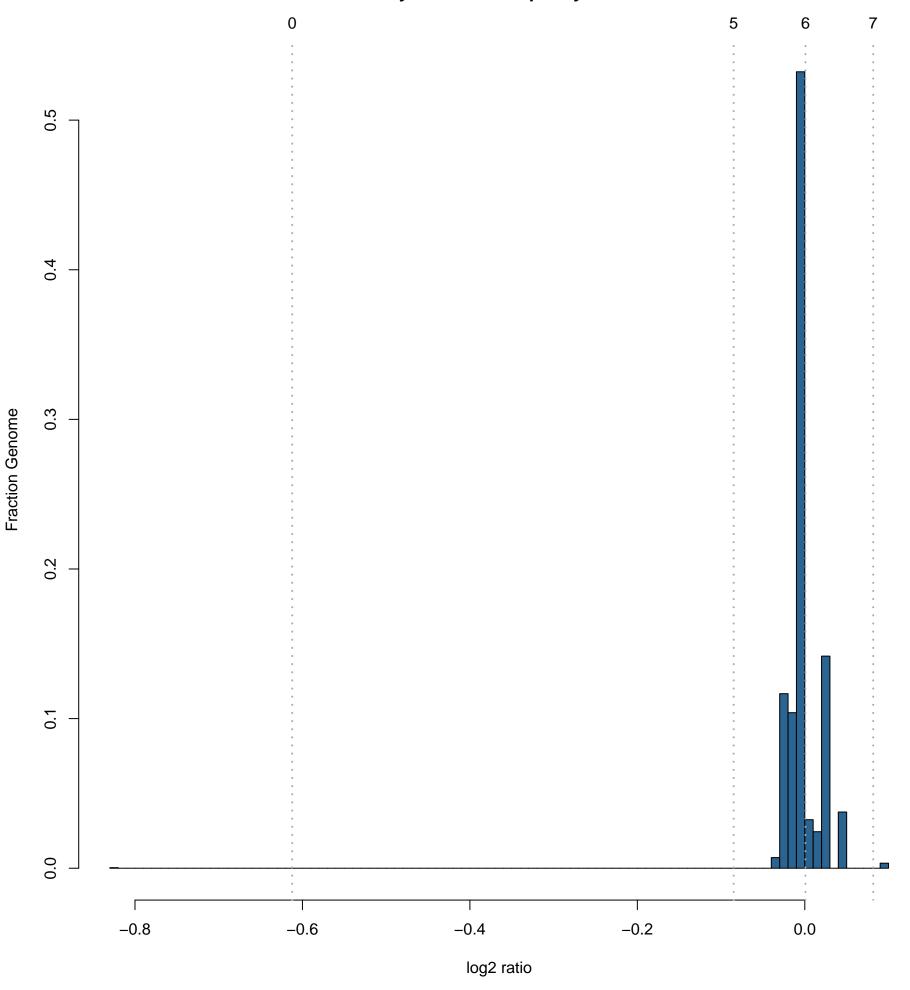


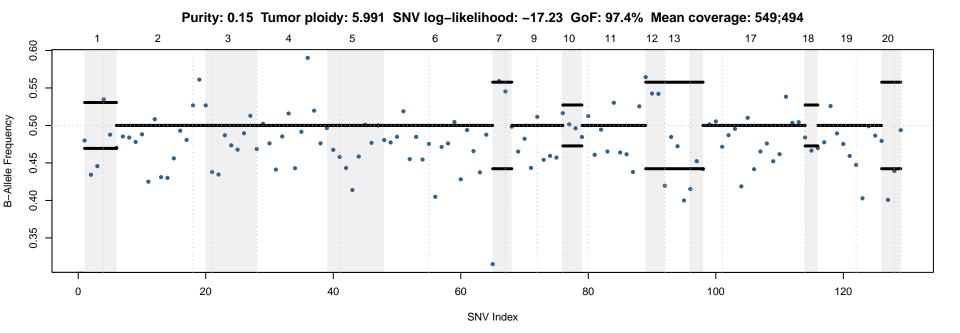


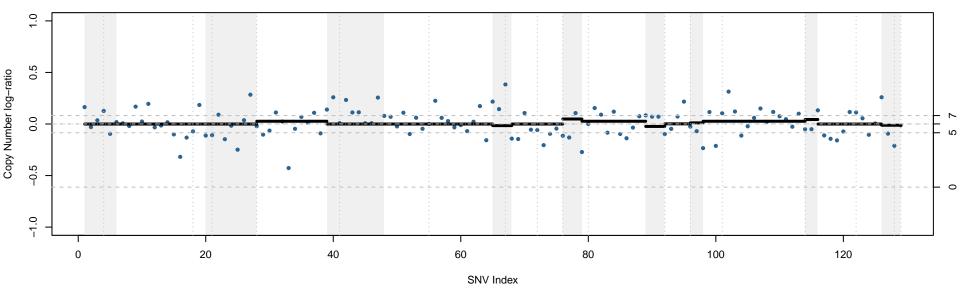


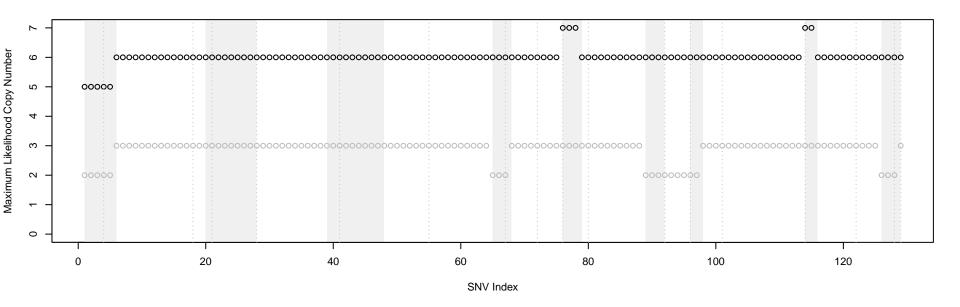


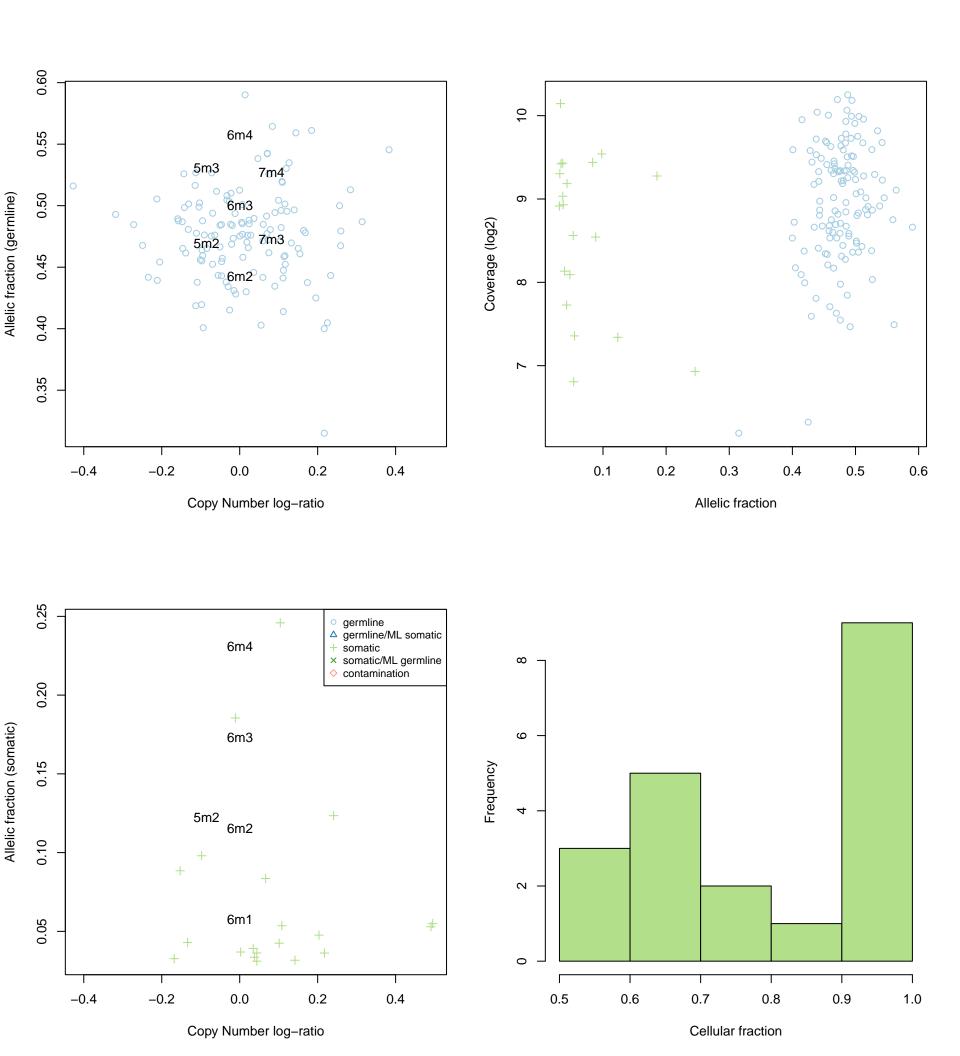
Purity: 0.15 Tumor ploidy: 5.991



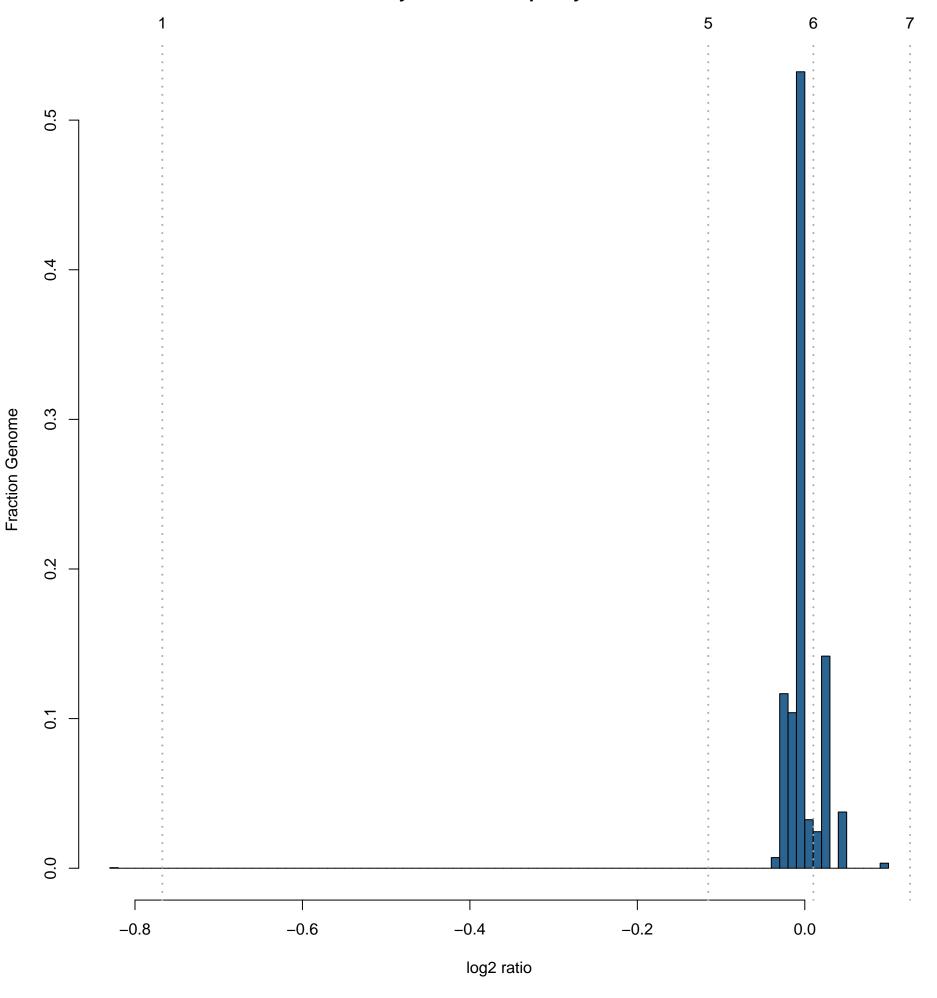


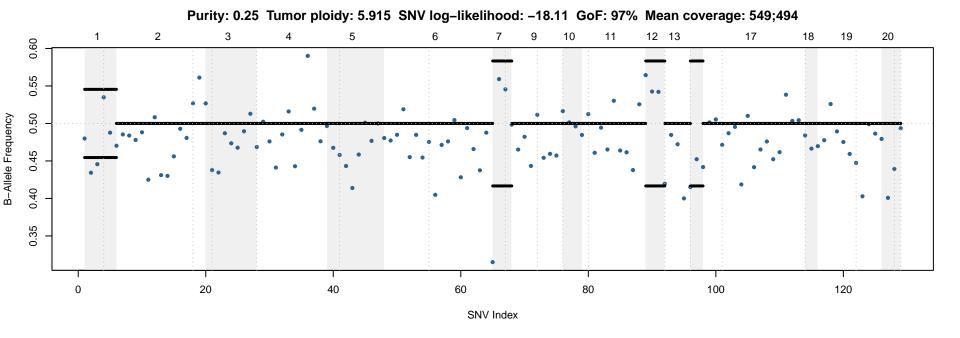


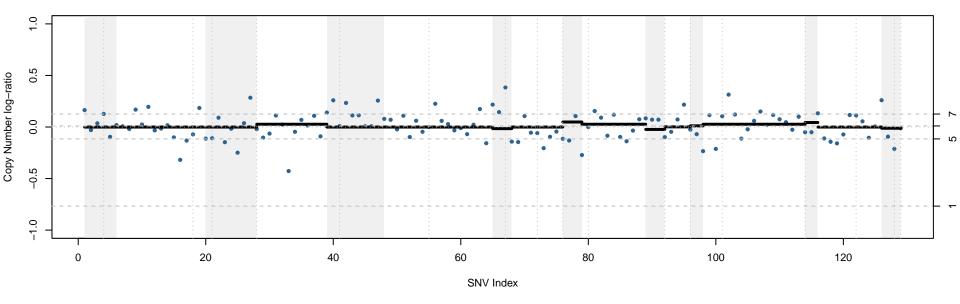


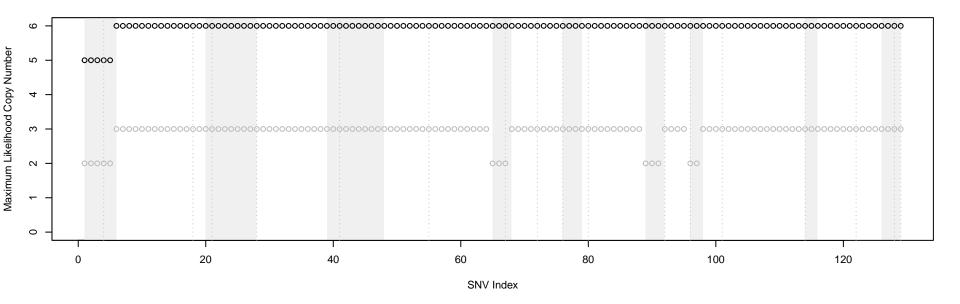


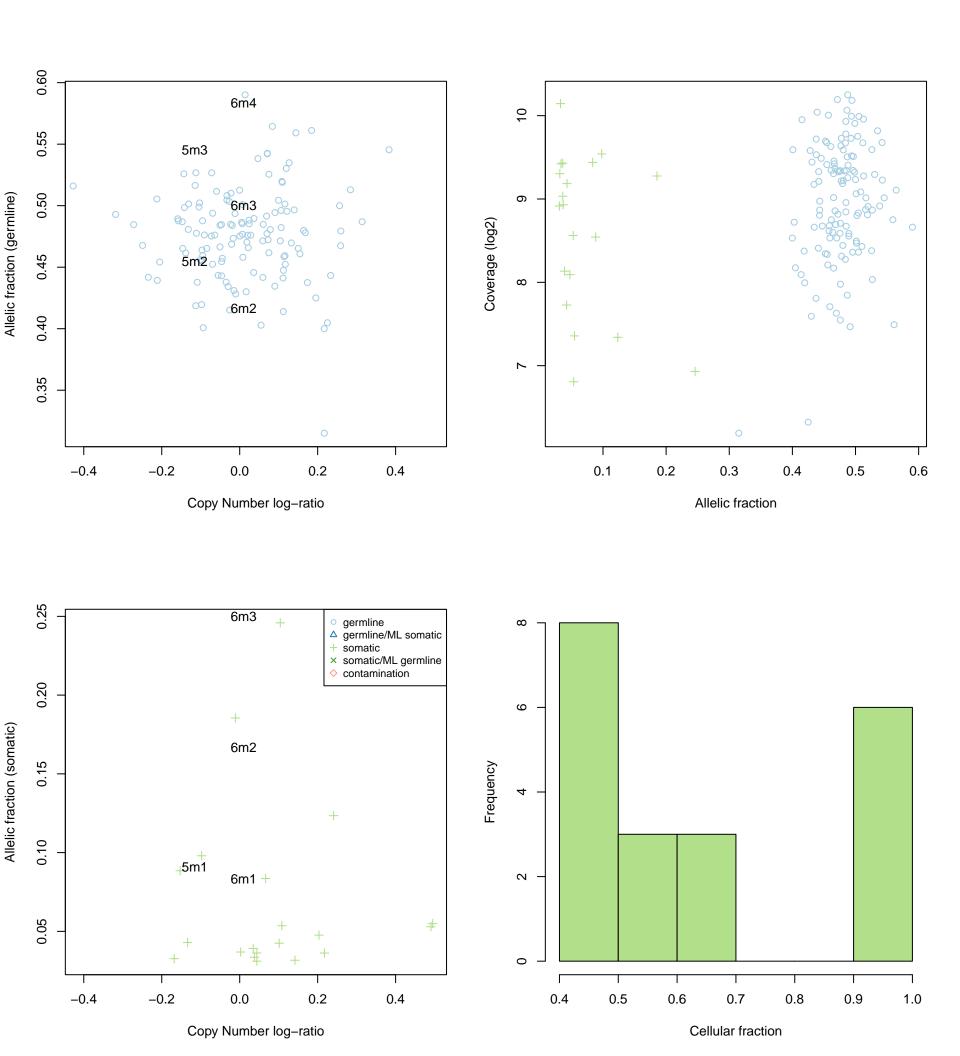
Purity: 0.25 Tumor ploidy: 5.915



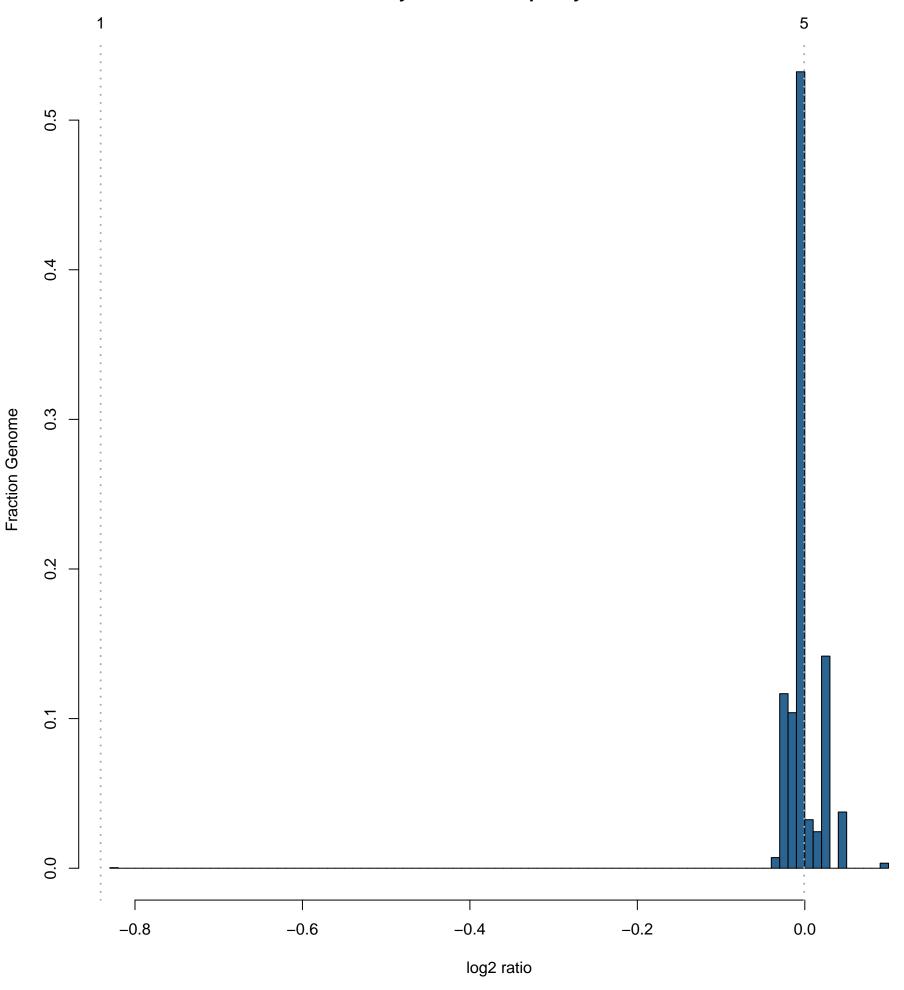


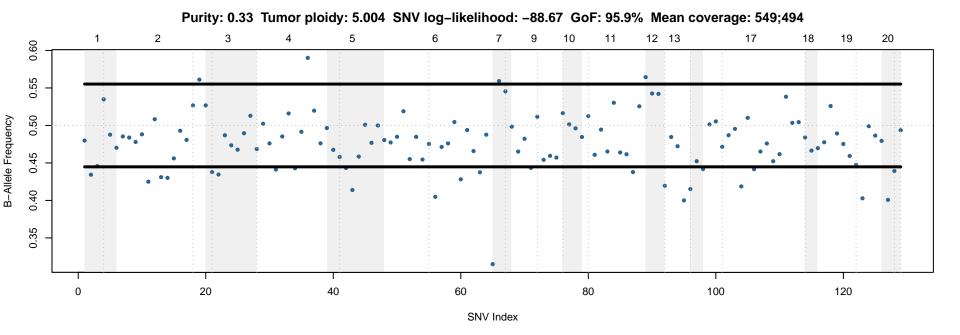


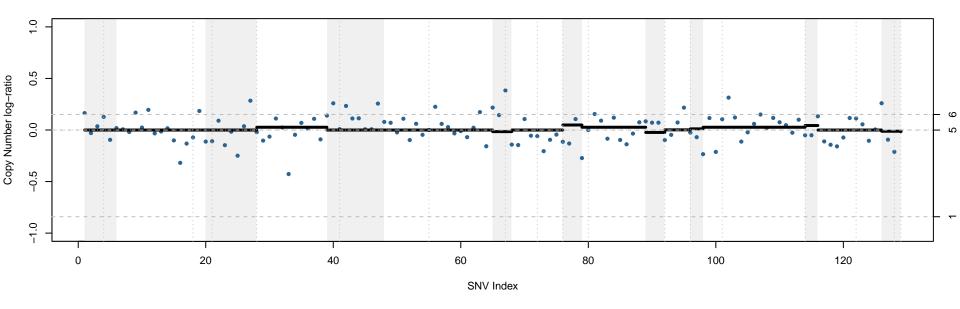


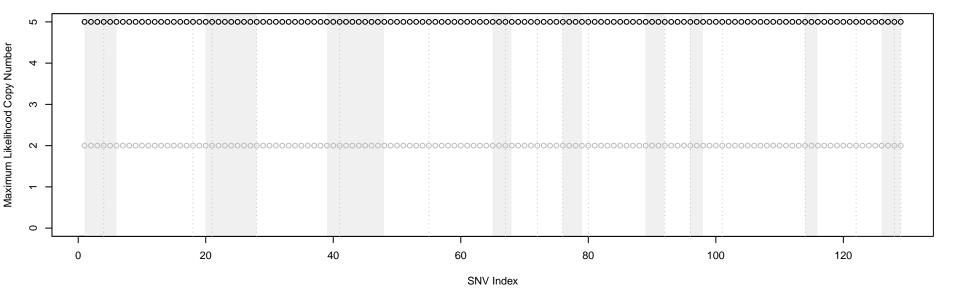


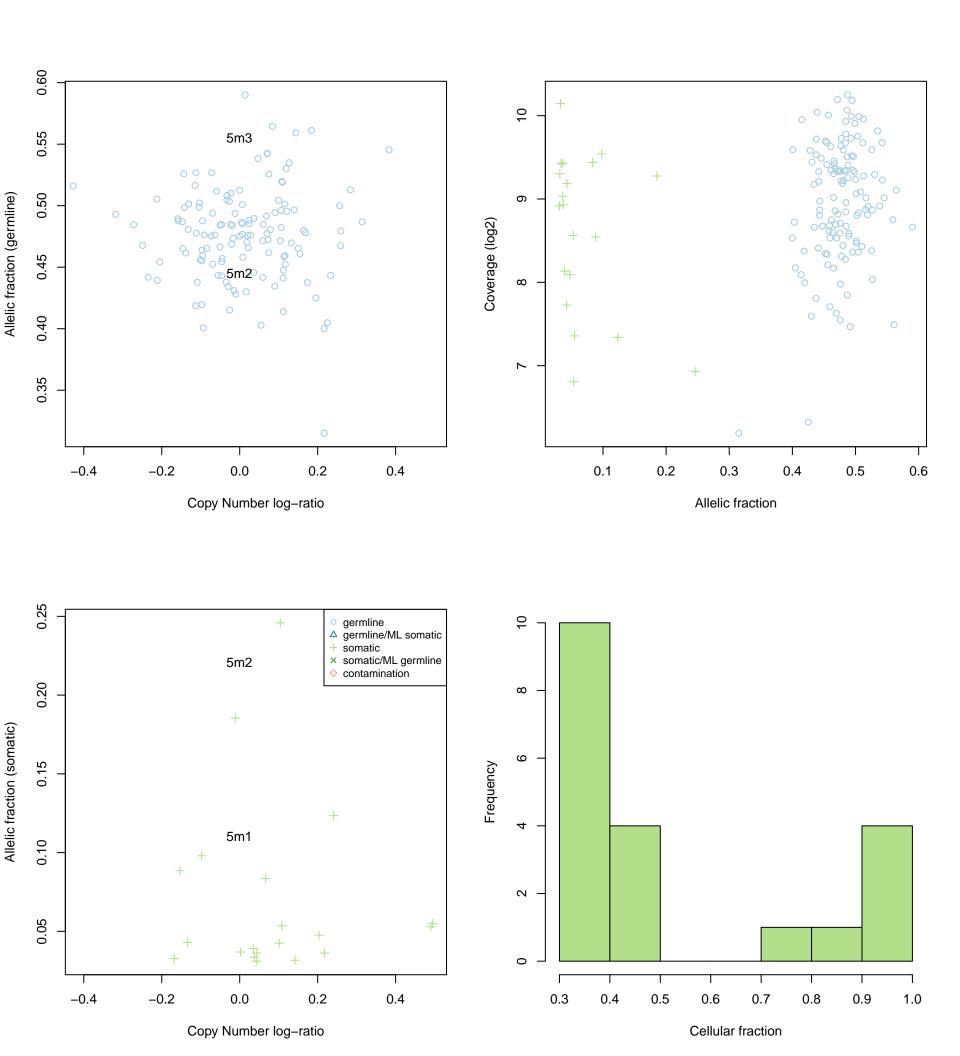
Purity: 0.33 Tumor ploidy: 5.004



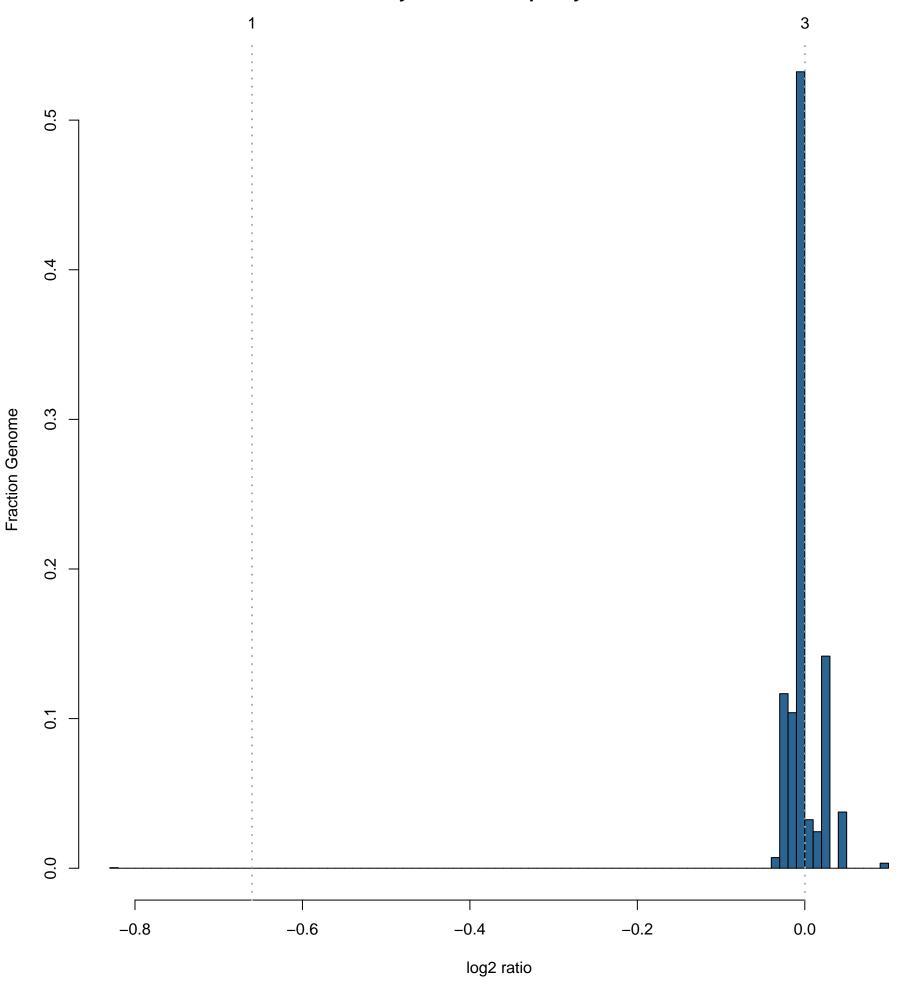


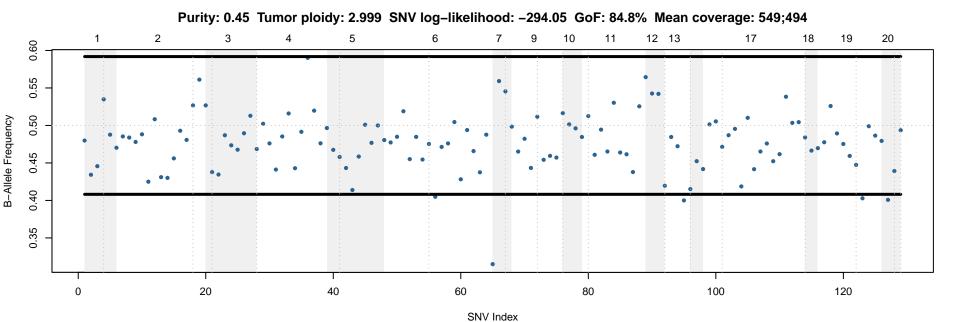




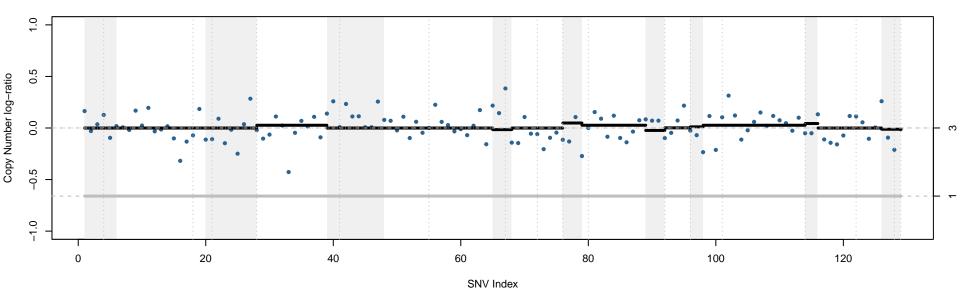


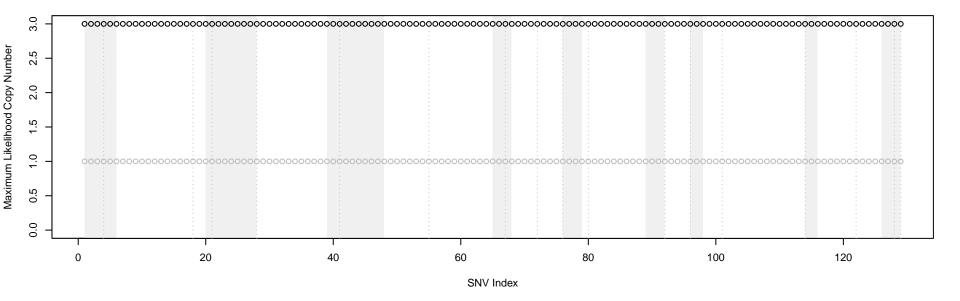
Purity: 0.45 Tumor ploidy: 2.999

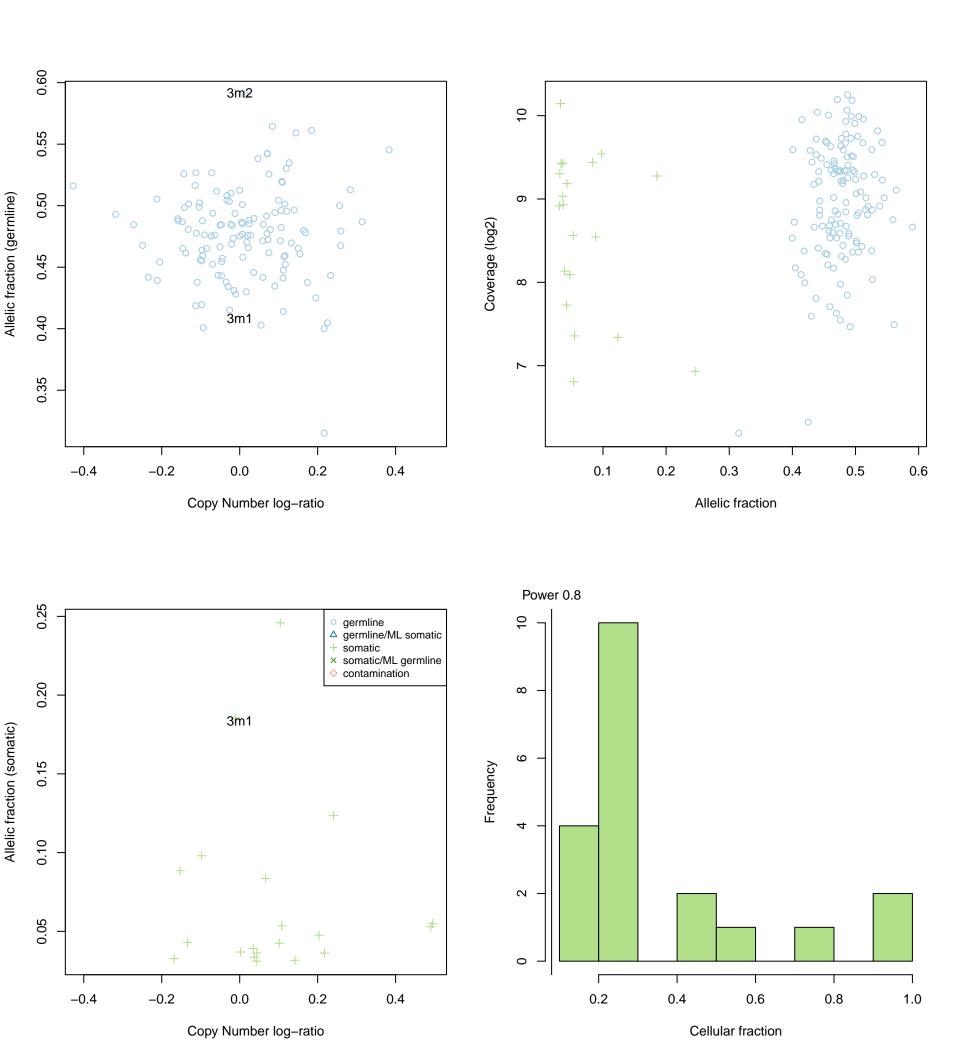




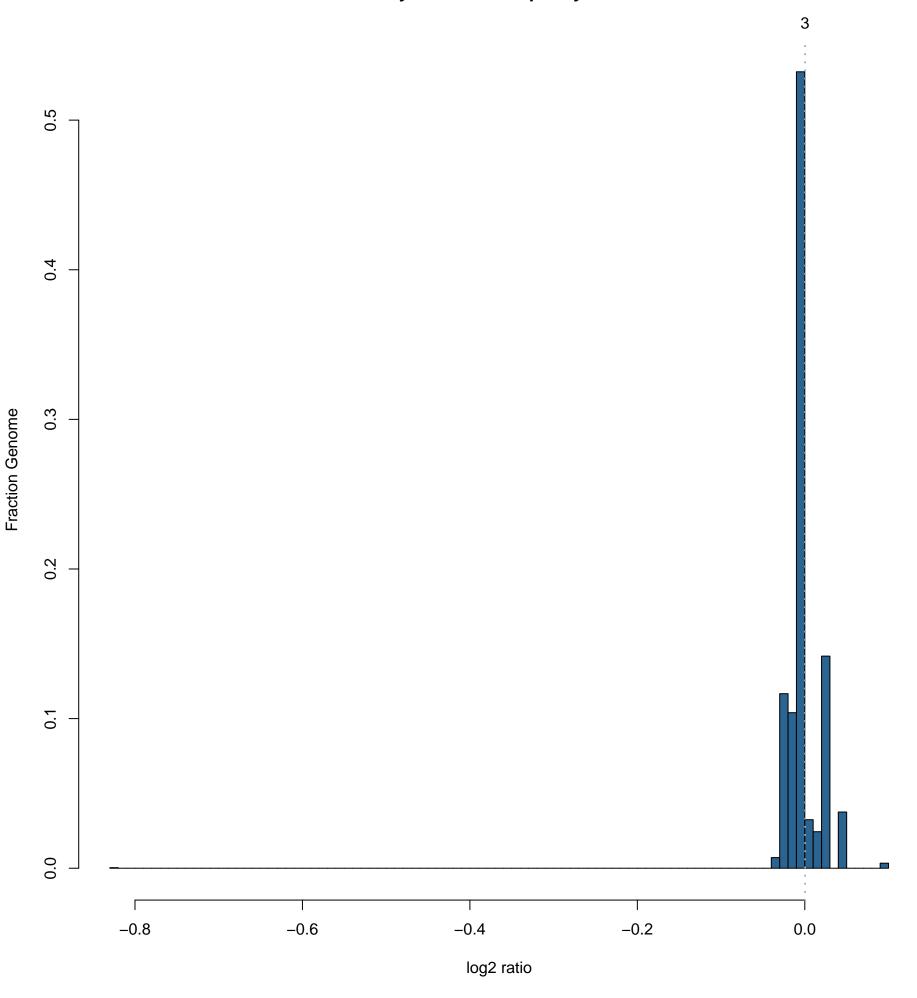
SCNA-fit log-likelihood: -4217.57

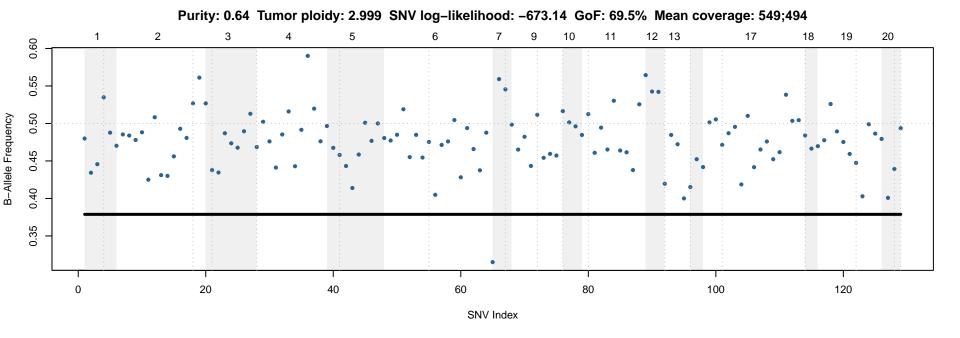


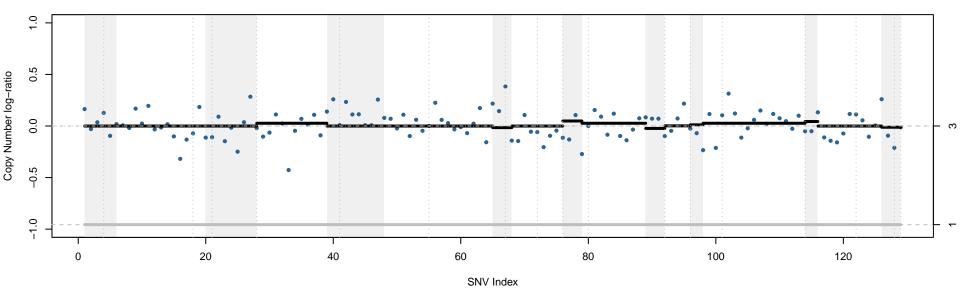


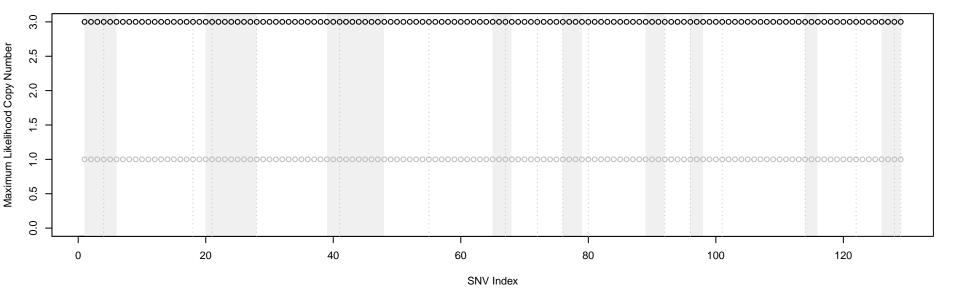


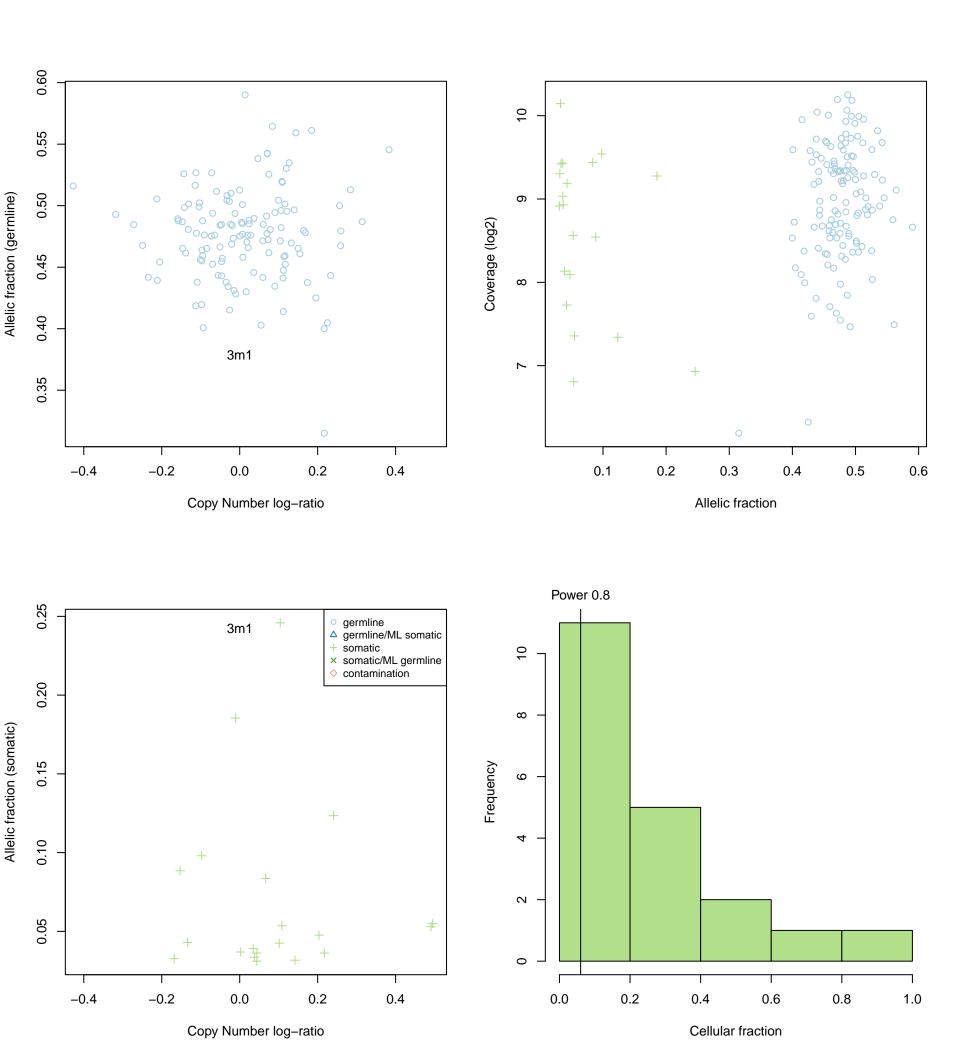
Purity: 0.64 Tumor ploidy: 2.999











Purity: 0.84 Tumor ploidy: 3 2 3 Fraction Genome 0.3 0.2 0.1 0.0 -0.8 -0.6 -0.4 -0.2 0.0

log2 ratio

