LGSA-001-195344_S3, n: 0.3, p: 2, log likelihood: 3576 Tumor Fraction: 0.6326, Ploidy: 1.74 Subclone Fraction: 0.872, Frac. Genome Subclonal: 0.01, Frac. CNA Subclonal: 0.04 Copy Number (log2 ratio)