LGSA-001-195314_S6, n: 0.2, p: 2, log likelihood: 3429 Tumor Fraction: 0.6985, Ploidy: 2.01 Subclone Fraction: 0.35, Frac. Genome Subclonal: 1.03, Frac. CNA Subclonal: 3.85 Copy Number (log2 ratio)