LGSA-001-195344_S3, n: 0.3, p: 3, log likelihood: 3352 Tumor Fraction: 0.9222, Ploidy: 2.74 Subclone Fraction: 0.172, Frac. Genome Subclonal: 0.02, Frac. CNA Subclonal: 0.03 Copy Number (log2 ratio)