LGSA-001-195344_S3, n: 0.2, p: 2, log likelihood: 3580 Tumor Fraction: 0.9946, Ploidy: 1.84 Subclone Fraction: 0.635, Frac. Genome Subclonal: 0.34, Frac. CNA Subclonal: 1.00 Copy Number (log2 ratio)