LGSA-001-195344_S3, n: 0.5, p: 3, log likelihood: 3375 Tumor Fraction: 0.9199, Ploidy: 2.74
Subclone Fraction: 0.07, Frac. Genome Subclonal: 0.08, Frac. CNA Subclonal: 0.11 Copy Number (log2 ratio)