LGSA-001-195344_S3, n: 0.9, p: 3, log likelihood: 3241 Tumor Fraction: 0.3769, Ploidy: 2.48
Subclone Fraction: 0.89, Frac. Genome Subclonal: 0.07, Frac. CNA Subclonal: 0.07 Copy Number (log2 ratio)