LGSA-001-195344\_S3, n: 0.6, p: 2, log likelihood: 3601 Tumor Fraction: 0.6334, Ploidy: 1.74 Subclone Fraction: 0.151, Frac. Genome Subclonal: 0.01, Frac. CNA Subclonal: 0.03 Copy Number (log2 ratio)