LGSA-001-195344\_S3, n: 0.9, p: 2, log likelihood: 3315 Tumor Fraction: 0.3766, Ploidy: 2.48
Subclone Fraction: 0.985, Frac. Genome Subclonal: 0.17, Frac. CNA Subclonal: 0.17 Copy Number (log2 ratio)