HCS-1070, n: 0.5, p: 2, log likelihood: 1381 Tumor Fraction: 0.5969, Ploidy: 1.78
Subclone Fraction: 0.392, Frac. Genome Subclonal: 0.13, Frac. CNA Subclonal: 0.33 Copy Number (log2 ratio)

HCS-1070, n: 0.6, p: 2, log likelihood: 1381 Tumor Fraction: 0.5968, Ploidy: 1.78
Subclone Fraction: 0.393, Frac. Genome Subclonal: 0.13, Frac. CNA Subclonal: 0.33 Copy Number (log2 ratio)

HCS-1070, n: 0.4, p: 2, log likelihood: 1380 Tumor Fraction: 0.5969, Ploidy: 1.78
Subclone Fraction: 0.392, Frac. Genome Subclonal: 0.13, Frac. CNA Subclonal: 0.33 Copy Number (log2 ratio)

HCS-1070, n: 0.7, p: 2, log likelihood: 1380 Tumor Fraction: 0.5969, Ploidy: 1.78
Subclone Fraction: 0.393, Frac. Genome Subclonal: 0.13, Frac. CNA Subclonal: 0.33 Copy Number (log2 ratio)

HCS-1070, n: 0.3, p: 2, log likelihood: 1379 Tumor Fraction: 0.597, Ploidy: 1.78
Subclone Fraction: 0.392, Frac. Genome Subclonal: 0.13, Frac. CNA Subclonal: 0.33 Copy Number (log2 ratio)

HCS-1070, n: 0.3, p: 3, log likelihood: 1104 Tumor Fraction: 0.8805, Ploidy: 2.75
Subclone Fraction: 0.795, Frac. Genome Subclonal: 0.12, Frac. CNA Subclonal: 0.16 Copy Number (log2 ratio)