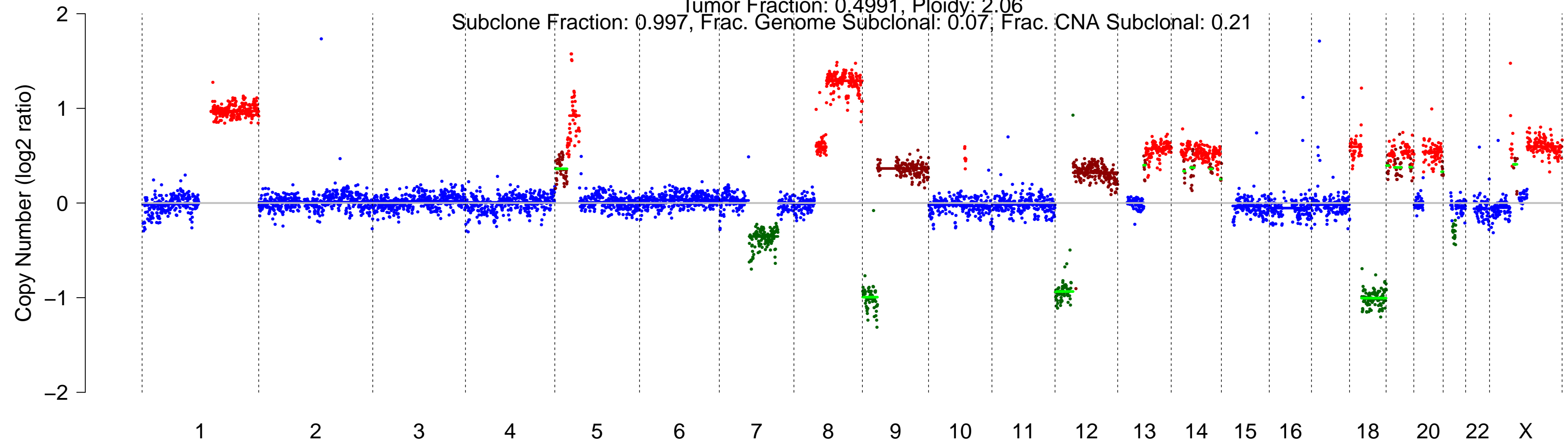


NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.7, p: 2, log likelihood: 5579

Tumor Fraction: 0.4991, Ploidy: 2.06

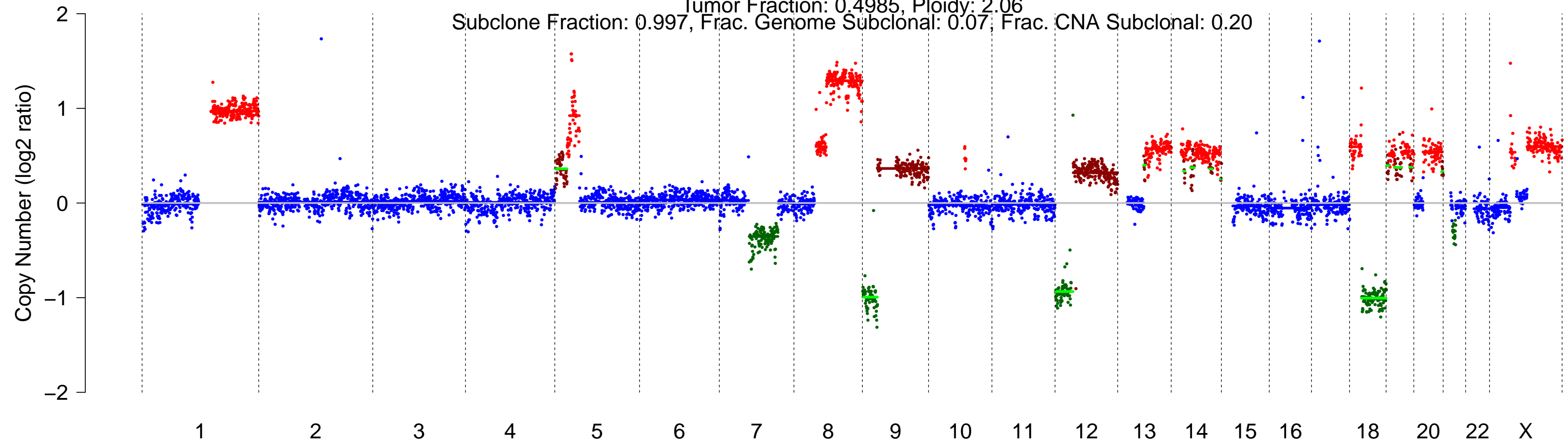
Subclone Fraction: 0.997, Frac. Genome Subclonal: 0.07, Frac. CNA Subclonal: 0.21



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.9, p: 2, log likelihood: 5579

Tumor Fraction: 0.4985, Ploidy: 2.06

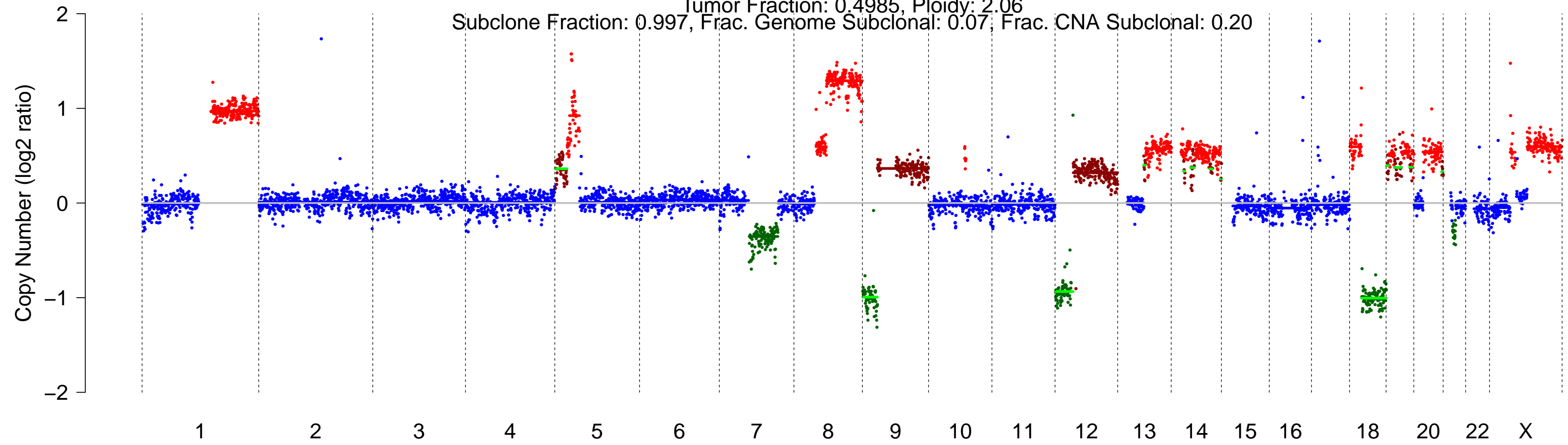
Subclone Fraction: 0.997, Frac. Genome Subclonal: 0.07, Frac. CNA Subclonal: 0.20



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.9, p: 3, log likelihood: 5578

Tumor Fraction: 0.4985, Ploidy: 2.06

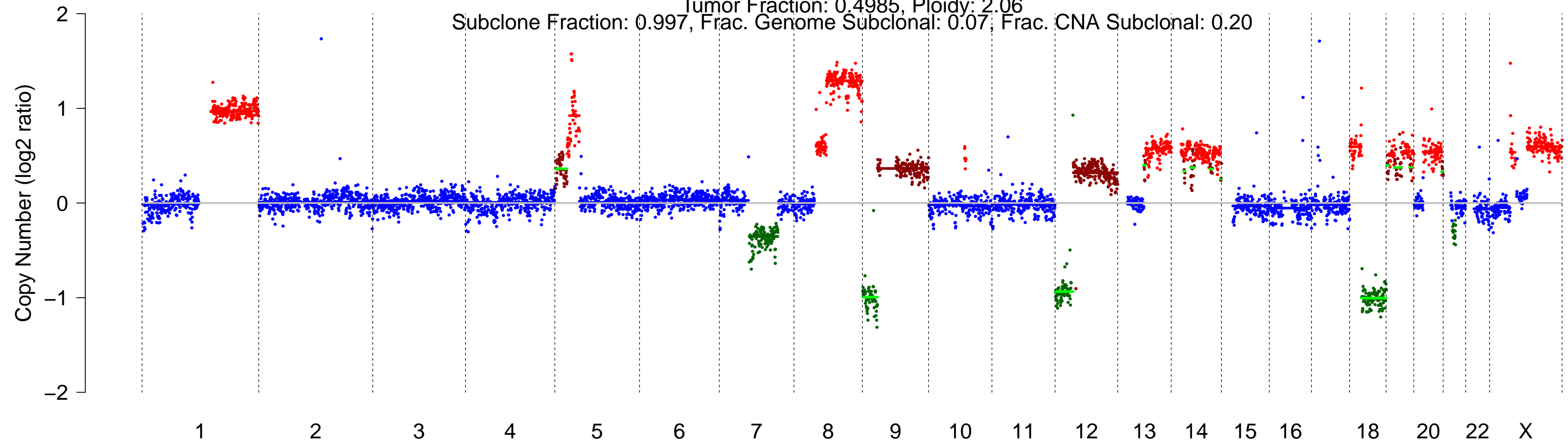
Subclone Fraction: 0.997, Frac. Genome Subclonal: 0.07, Frac. CNA Subclonal: 0.20



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.8, p: 2, log likelihood: 5577

Tumor Fraction: 0.4985, Ploidy: 2.06

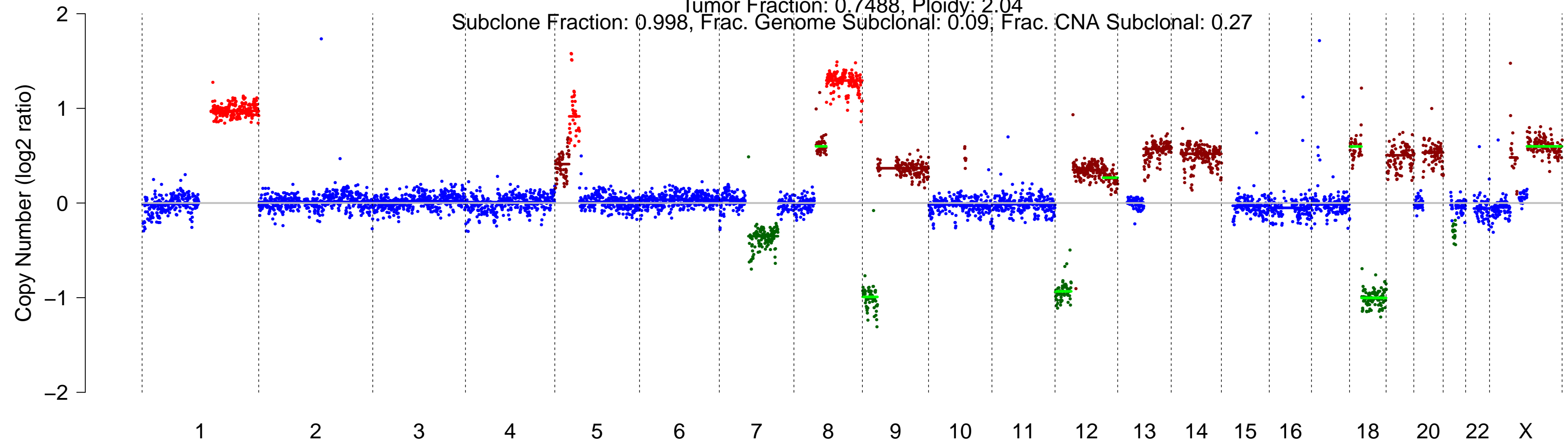
Subclone Fraction: 0.997, Frac. Genome Subclonal: 0.07, Frac. CNA Subclonal: 0.20



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.5, p: 2, log likelihood: 5397

Tumor Fraction: 0.7488, Ploidy: 2.04

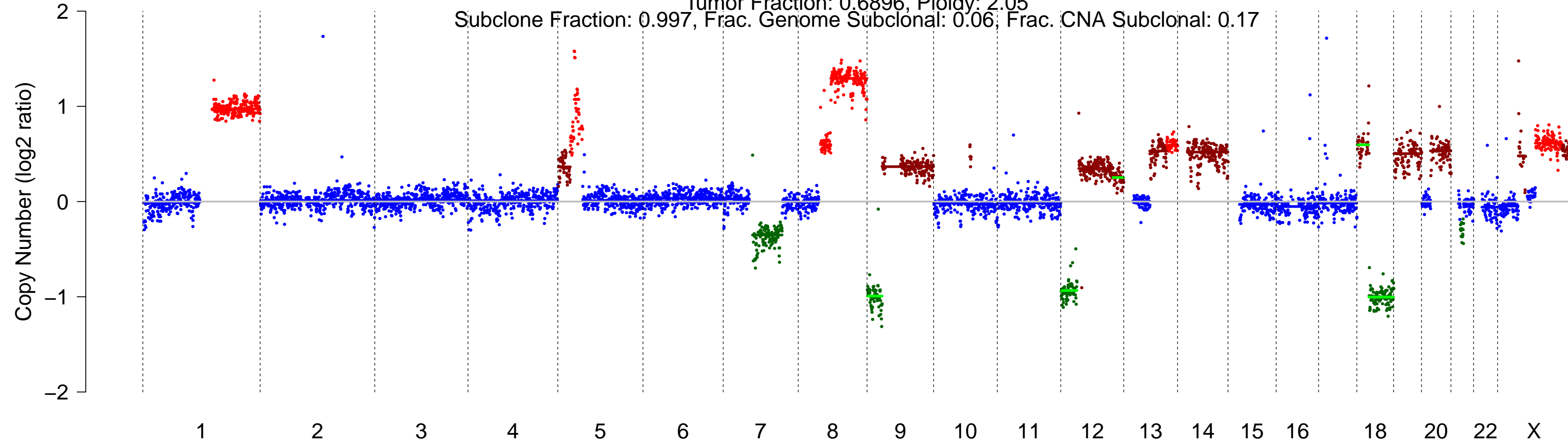
Subclone Fraction: 0.998, Frac. Genome Subclonal: 0.09; Frac. CNA Subclonal: 0.27



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.6, p: 2, log likelihood: 5381

Tumor Fraction: 0.6896, Ploidy: 2.05

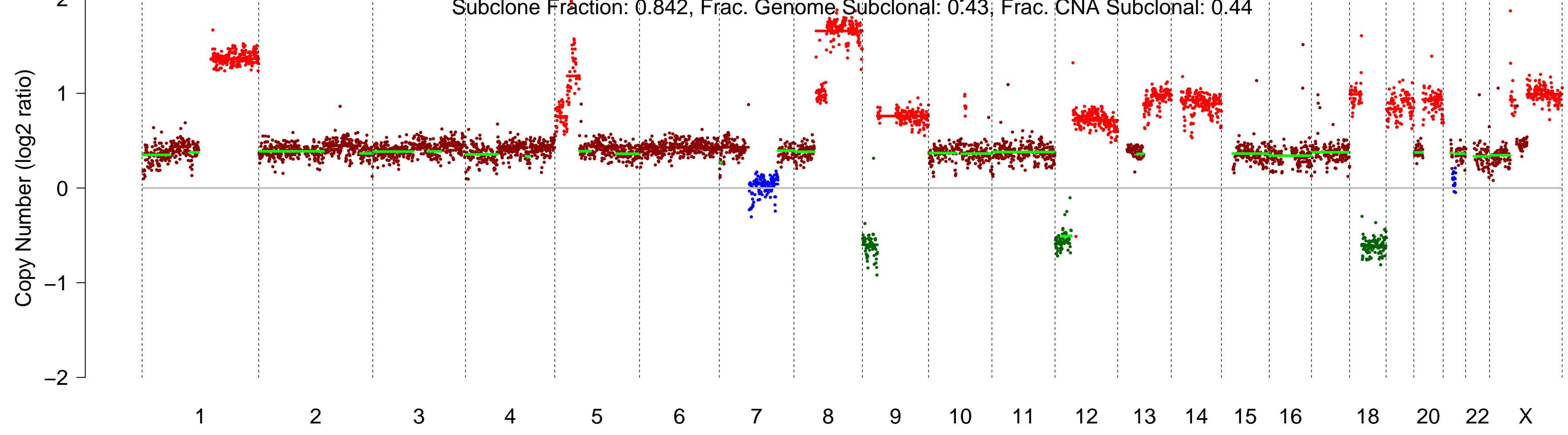
Subclone Fraction: 0.997, Frac. Genome Subclonal: 0.06; Frac. CNA Subclonal: 0.17



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.6, p: 3, log likelihood: 5117

Tumor Fraction: 0.6976, Ploidy: 2.96

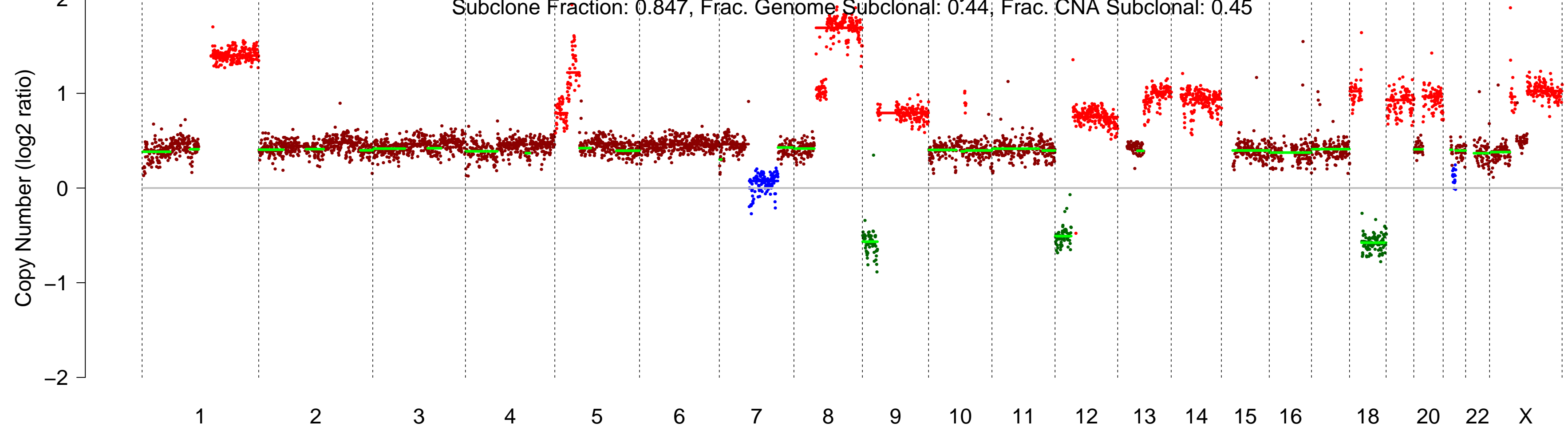
Subclone Fraction: 0.842, Frac. Genome Subclonal: 0.43, Frac. CNA Subclonal: 0.44



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.7, p: 3, log likelihood: 5113

Tumor Fraction: 0.7552, Ploidy: 2.97

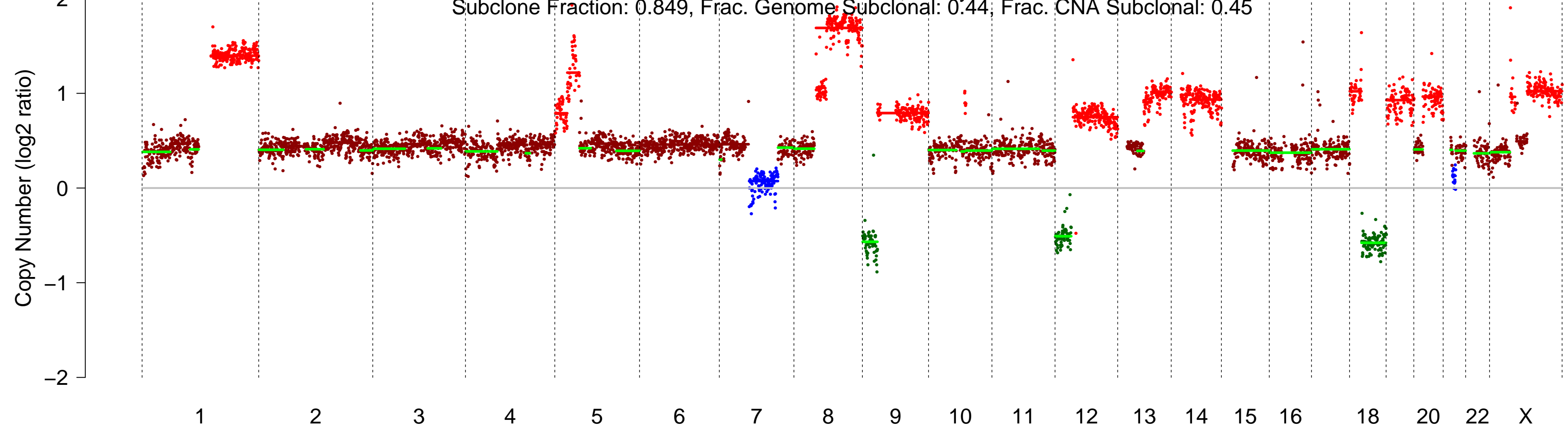
Subclone Fraction: 0.847, Frac. Genome Subclonal: 0.44, Frac. CNA Subclonal: 0.45



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.8, p: 3, log likelihood: 5112

Tumor Fraction: 0.7535, Ploidy: 2.96

Subclone Fraction: 0.849, Frac. Genome Subclonal: 0.44, Frac. CNA Subclonal: 0.45



NaomicfDNA2_S3_R1.dedup.filt.bam.500Kb, n: 0.5, p: 3, log likelihood: 5111

Tumor Fraction: 0.6857, Ploidy: 2.95

Subclone Fraction: 0.828, Frac. Genome Subclonal: 0.43, Frac. CNA Subclonal: 0.44

