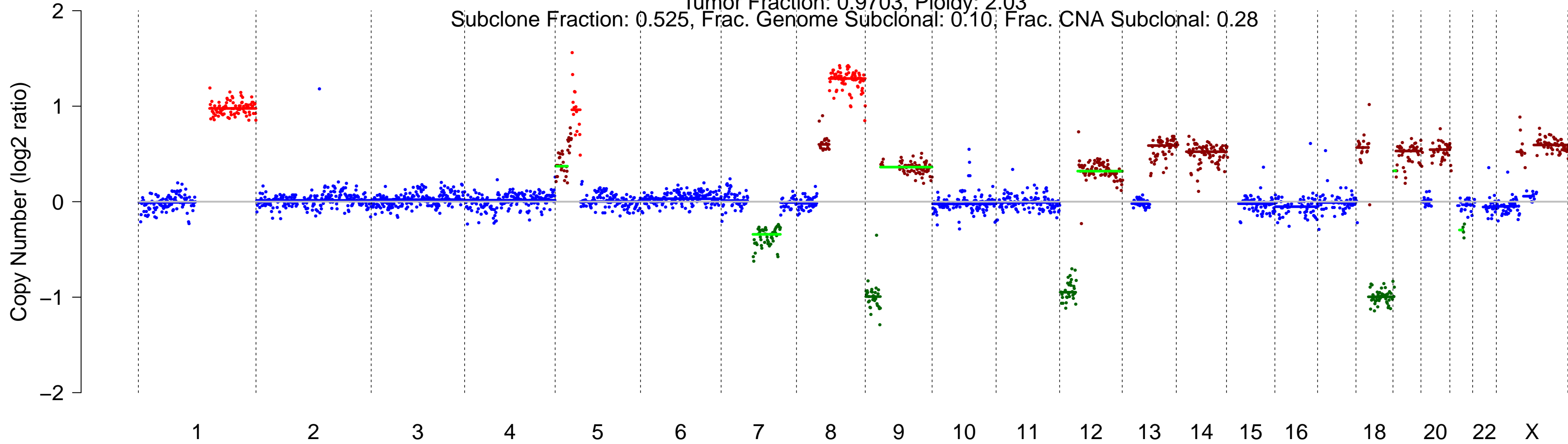


NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.5, p: 2, log likelihood: 3352

Tumor Fraction: 0.9703, Ploidy: 2.03

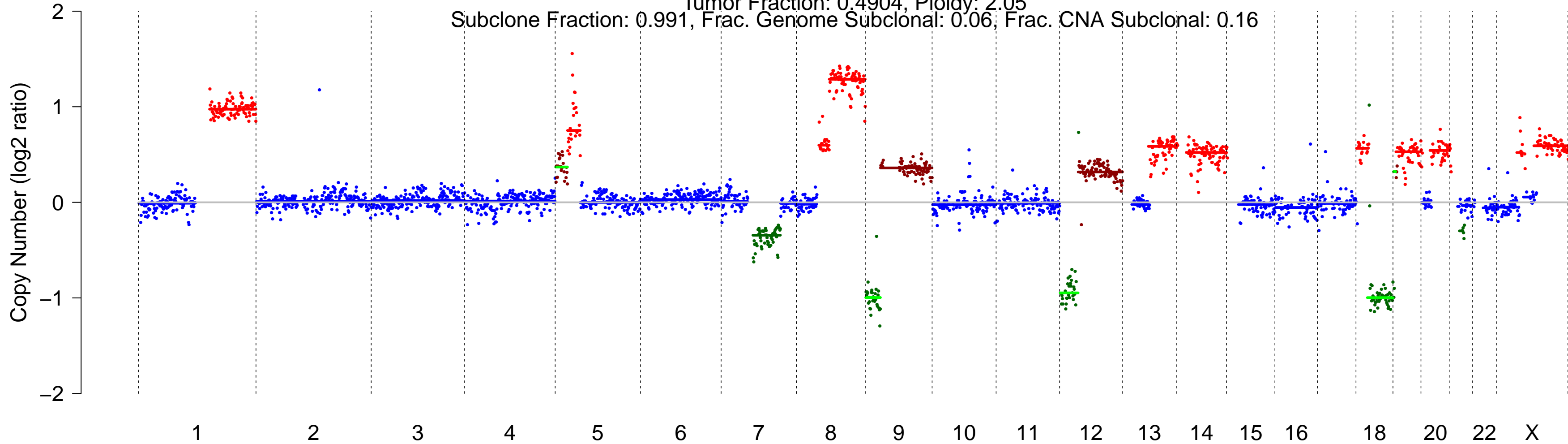
Subclone Fraction: 0.525, Frac. Genome Subclonal: 0.10; Frac. CNA Subclonal: 0.28



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.7, p: 2, log likelihood: 2898

Tumor Fraction: 0.4904, Ploidy: 2.05

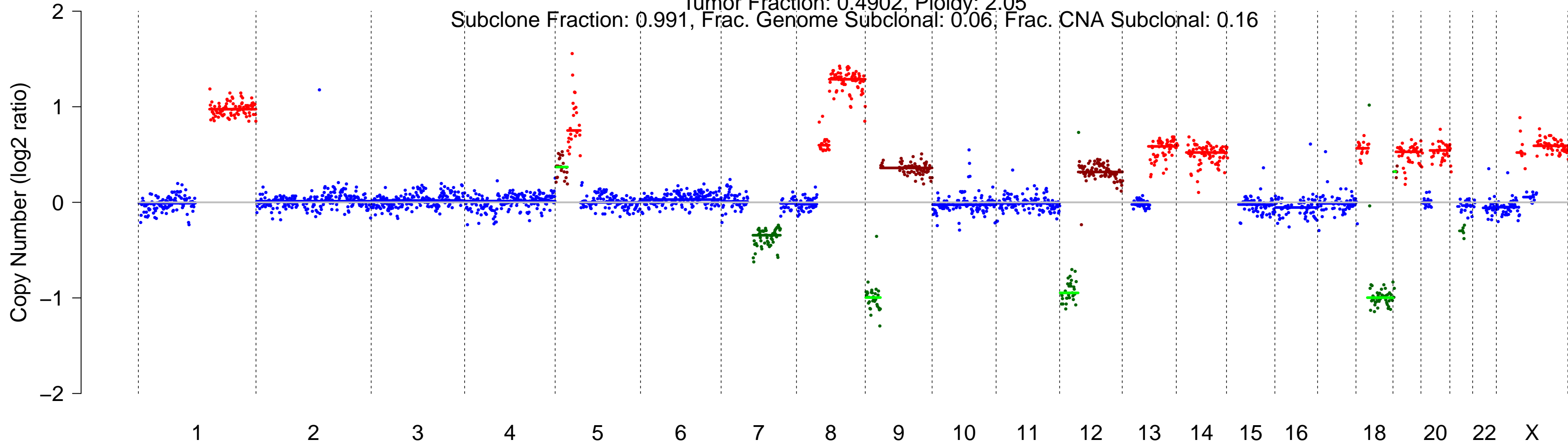
Subclone Fraction: 0.991, Frac. Genome Subclonal: 0.06; Frac. CNA Subclonal: 0.16



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.8, p: 2, log likelihood: 2898

Tumor Fraction: 0.4902, Ploidy: 2.05

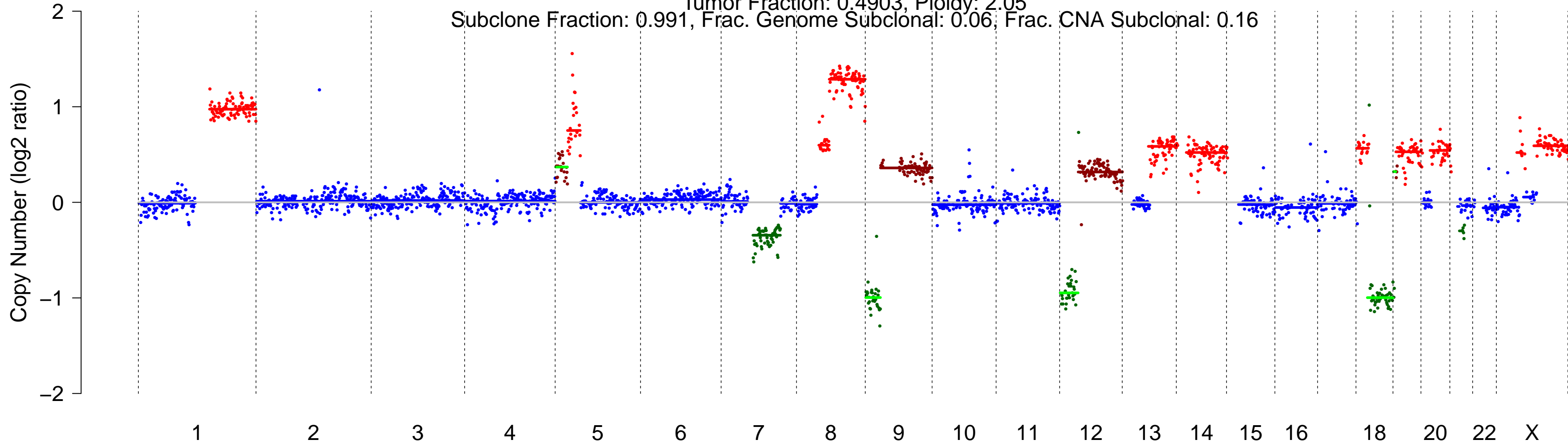
Subclone Fraction: 0.991, Frac. Genome Subclonal: 0.06; Frac. CNA Subclonal: 0.16



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.9, p: 2, log likelihood: 2898

Tumor Fraction: 0.4903, Ploidy: 2.05

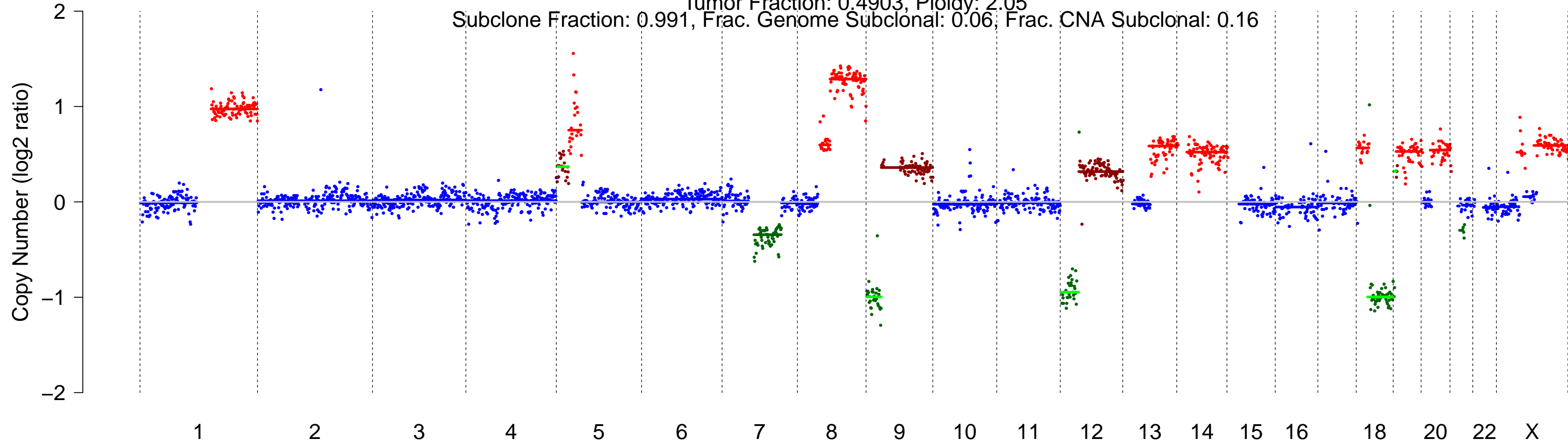
Subclone Fraction: 0.991, Frac. Genome Subclonal: 0.06; Frac. CNA Subclonal: 0.16



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.9, p: 3, log likelihood: 2898

Tumor Fraction: 0.4903, Ploidy: 2.05

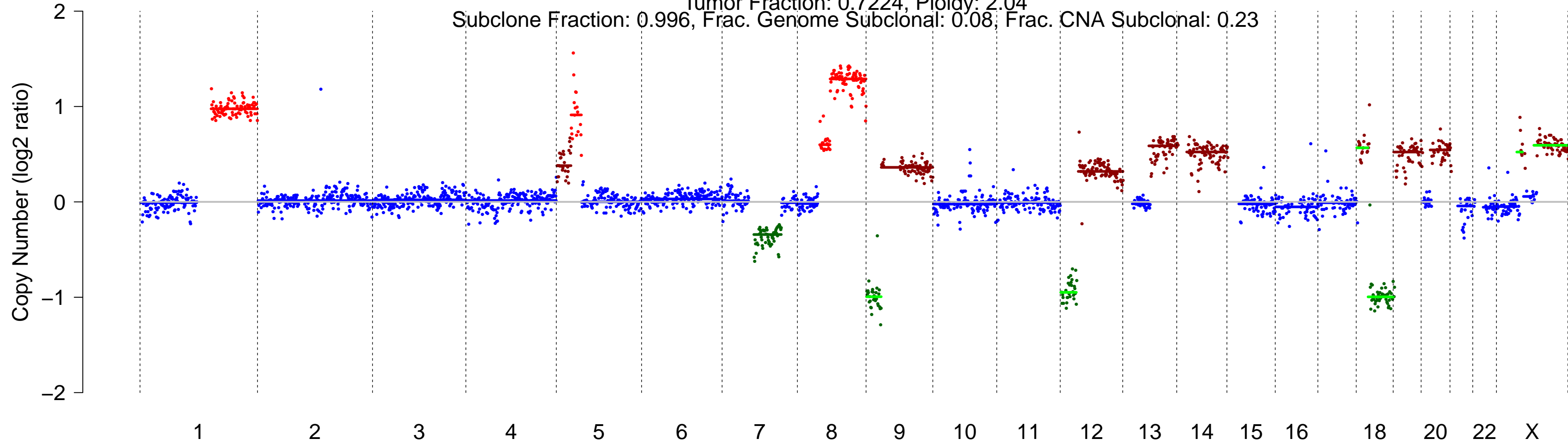
Subclone Fraction: 0.991, Frac. Genome Subclonal: 0.06; Frac. CNA Subclonal: 0.16



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.6, p: 2, log likelihood: 2805

Tumor Fraction: 0.7224, Ploidy: 2.04

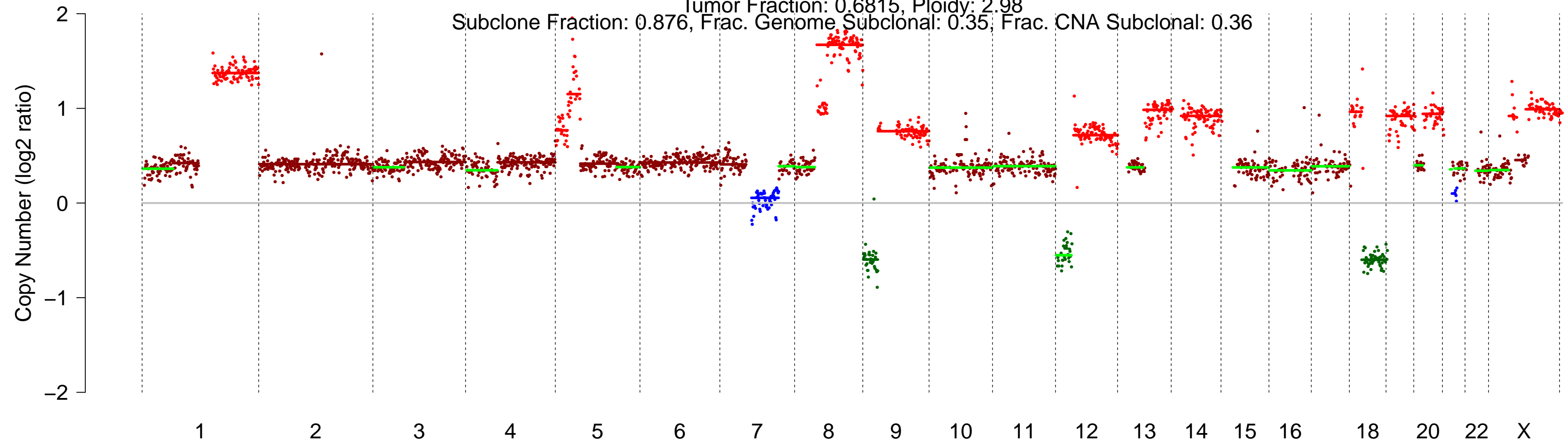
Subclone Fraction: 0.996, Frac. Genome Subclonal: 0.08; Frac. CNA Subclonal: 0.23



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.6, p: 3, log likelihood: 2615

Tumor Fraction: 0.6815, Ploidy: 2.98

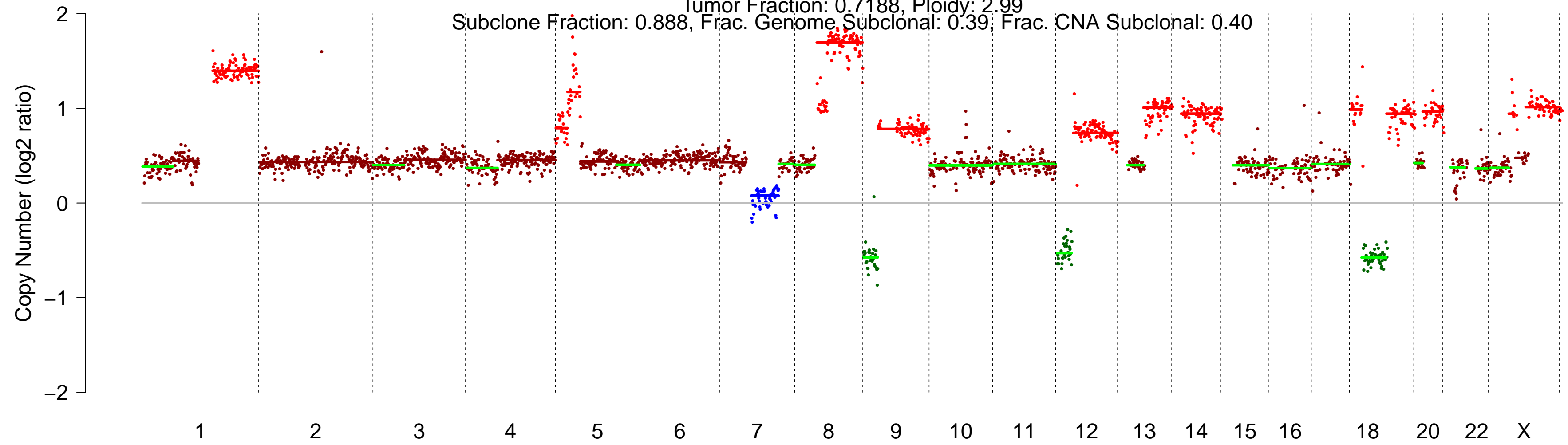
Subclone Fraction: 0.876, Frac. Genome Subclonal: 0.35; Frac. CNA Subclonal: 0.36



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.7, p: 3, log likelihood: 2601

Tumor Fraction: 0.7188, Ploidy: 2.99

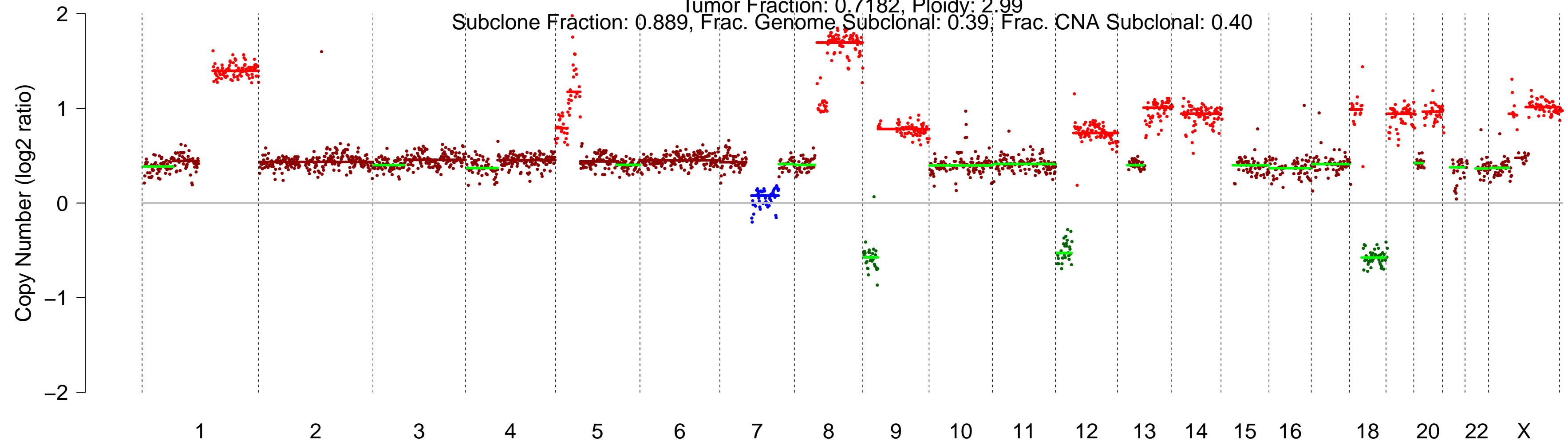
Subclone Fraction: 0.888, Frac. Genome Subclonal: 0.39, Frac. CNA Subclonal: 0.40



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.8, p: 3, log likelihood: 2601

Tumor Fraction: 0.7182, Ploidy: 2.99

Subclone Fraction: 0.889, Frac. Genome Subclonal: 0.39; Frac. CNA Subclonal: 0.40



NaomicfDNA2_S3_R1.dedup.filt.bam.1Mb, n: 0.5, p: 3, log likelihood: 2338

Tumor Fraction: 0.6696, Ploidy: 3.04

Subclone Fraction: 0.469, Frac. Genome Subclonal: 0.00, Frac. CNA Subclonal: 0.00

