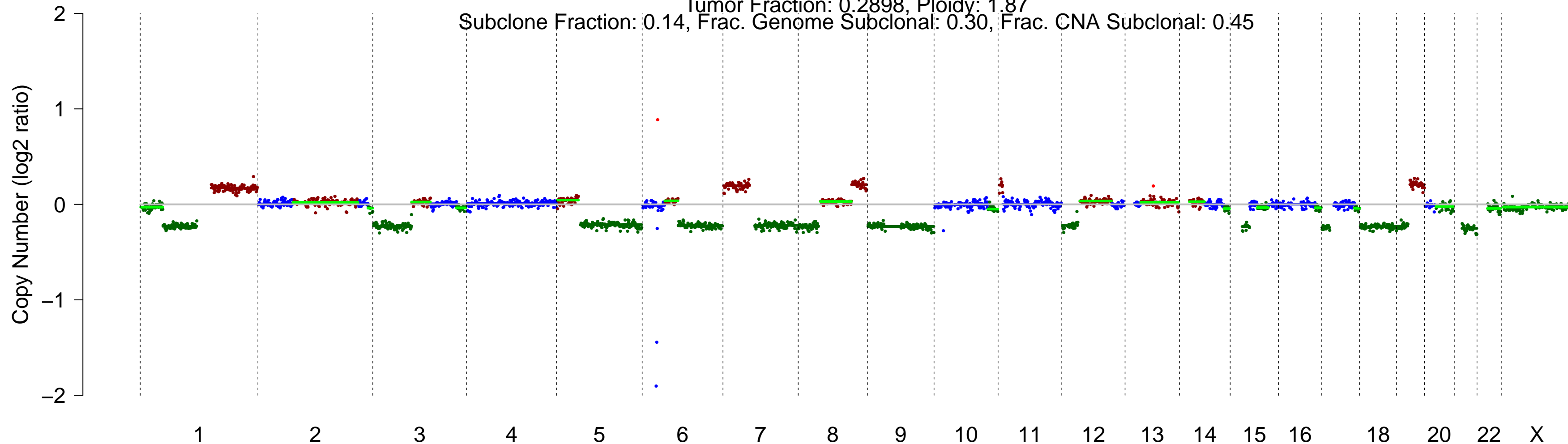


sample20200117-1_ZXY-WGS_I7, n: 0.7, p: 2, log likelihood: 5746

Tumor Fraction: 0.2898, Ploidy: 1.87

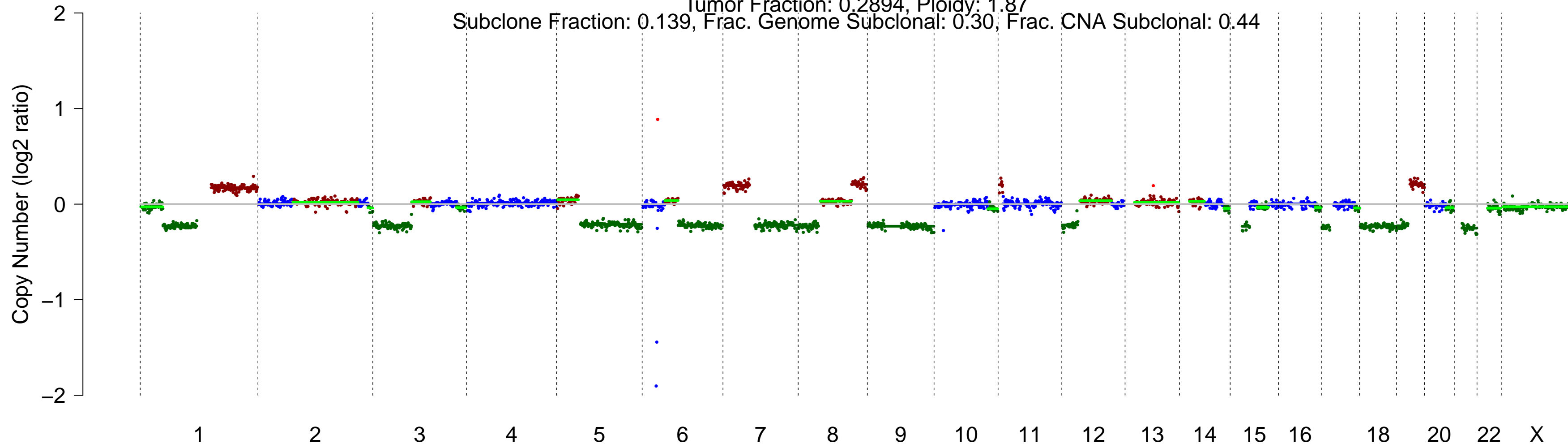
Subclone Fraction: 0.14, Frac. Genome Subclonal: 0.30, Frac. CNA Subclonal: 0.45



sample20200117-1_ZXY-WGS_I7, n: 0.8, p: 2, log likelihood: 5744

Tumor Fraction: 0.2894, Ploidy: 1.87

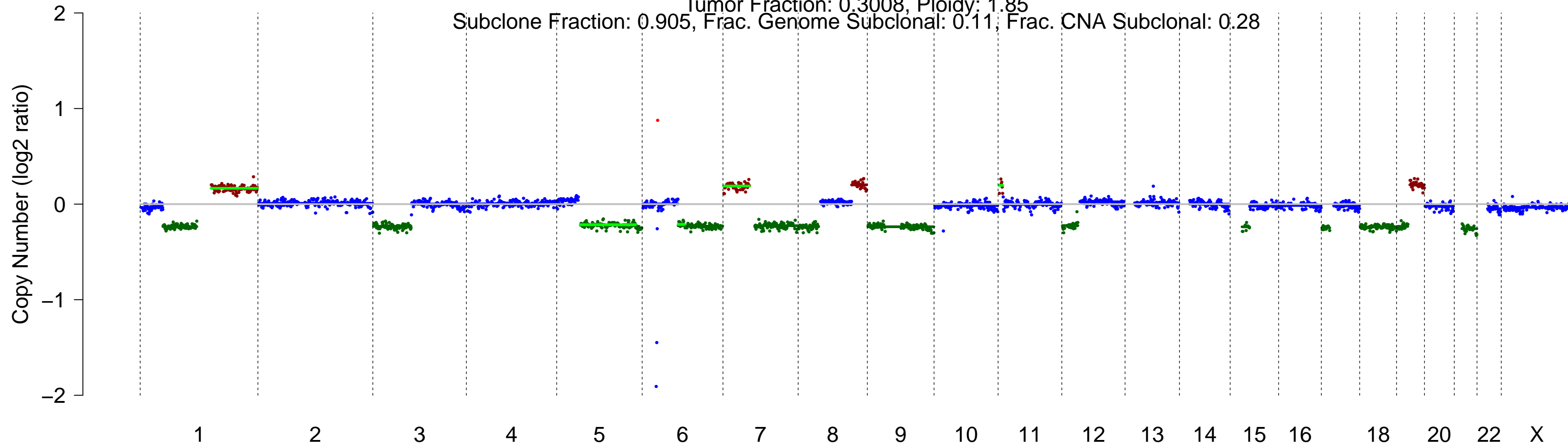
Subclone Fraction: 0.139, Frac. Genome Subclonal: 0.30, Frac. CNA Subclonal: 0.44



sample20200117-1_ZXY-WGS_I7, n: 0.6, p: 2, log likelihood: 5590

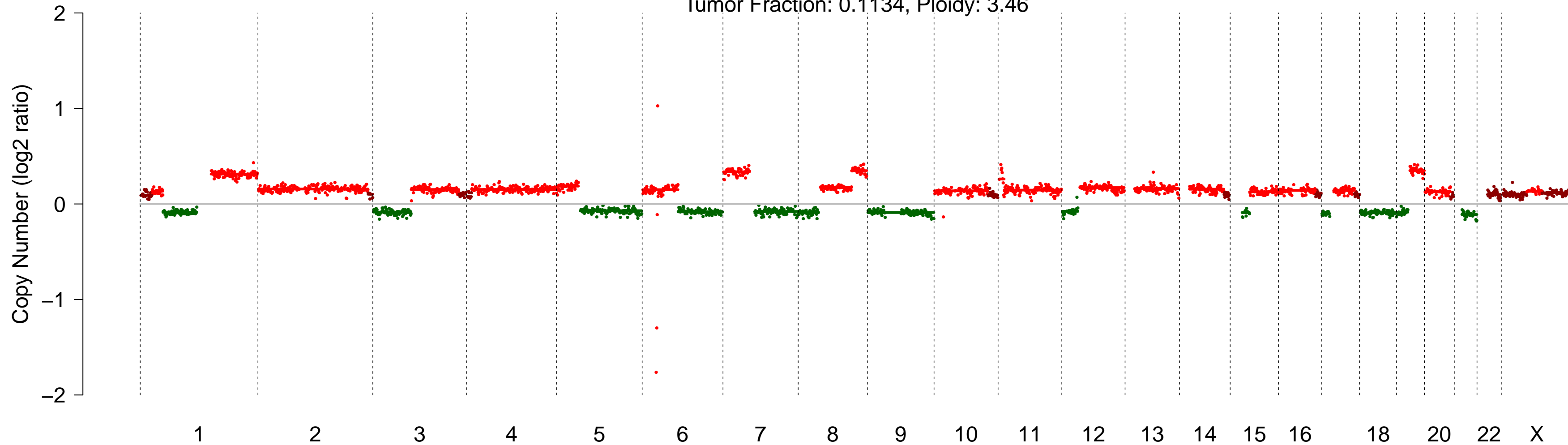
Tumor Fraction: 0.3008, Ploidy: 1.85

Subclone Fraction: 0.905, Frac. Genome Subclonal: 0.11, Frac. CNA Subclonal: 0.28



sample20200117-1_ZXY-WGS_I7, n: 0.9, p: 5, log likelihood: 4975

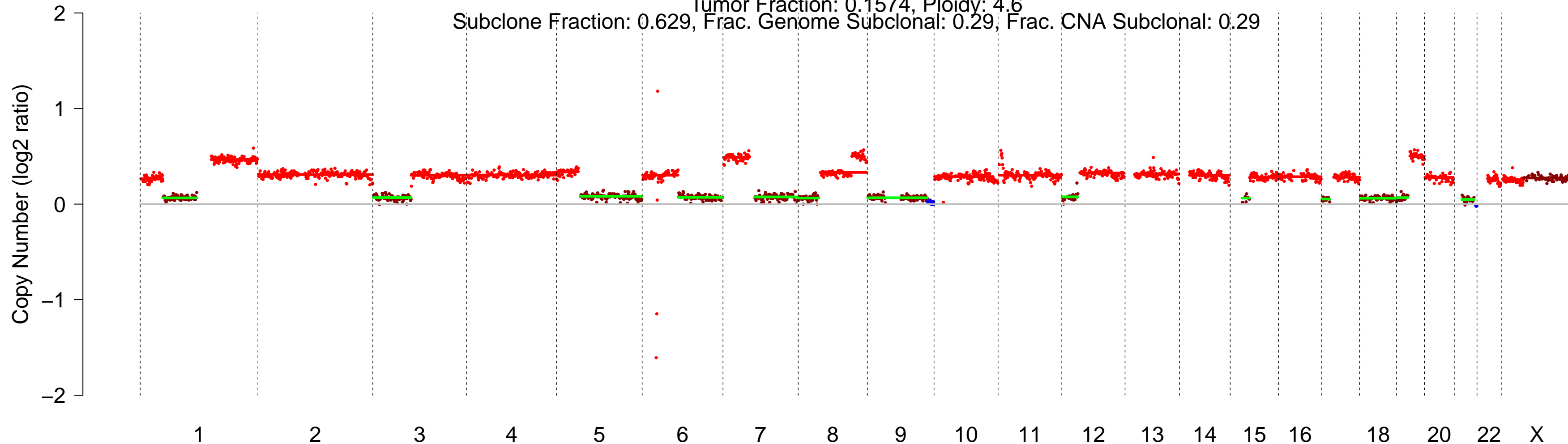
Tumor Fraction: 0.1134, Ploidy: 3.46



sample20200117-1_ZXY-WGS_I7, n: 0.8, p: 5, log likelihood: 4249

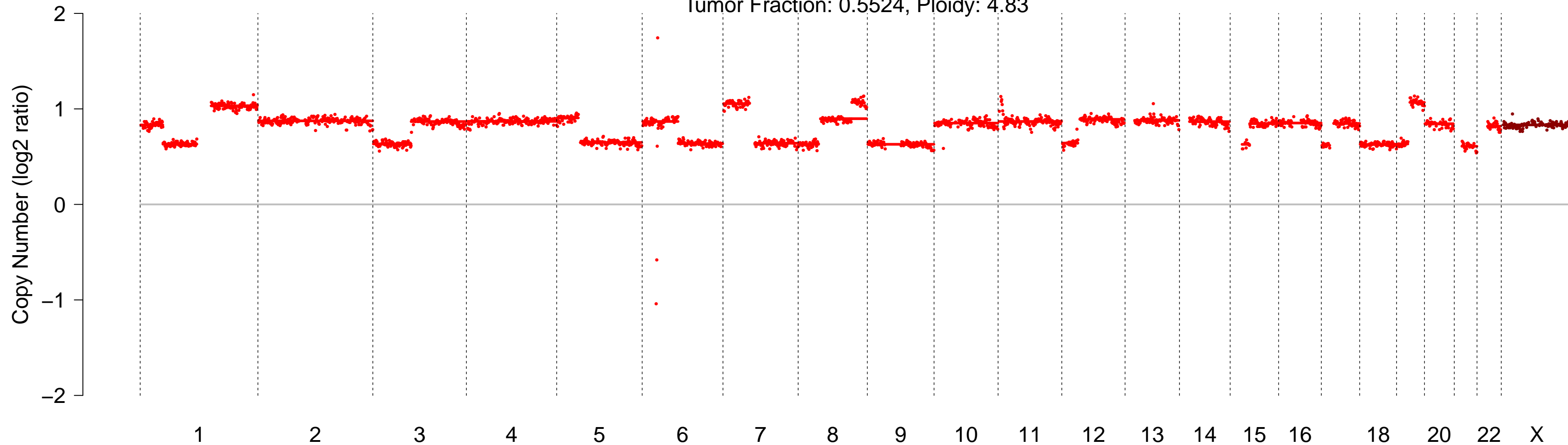
Tumor Fraction: 0.1574, Ploidy: 4.6

Subclone Fraction: 0.629, Frac. Genome Subclonal: 0.29, Frac. CNA Subclonal: 0.29



sample20200117-1_ZXY-WGS_I7, n: 0.7, p: 5, log likelihood: 4194

Tumor Fraction: 0.5524, Ploidy: 4.83



sample20200117-1_ZXY-WGS_I7, n: 0.5, p: 5, log likelihood: 4193

Tumor Fraction: 0.5495, Ploidy: 4.83



sample20200117-1_ZXY-WGS_I7, n: 0.6, p: 5, log likelihood: 4193

Tumor Fraction: 0.5506, Ploidy: 4.83

