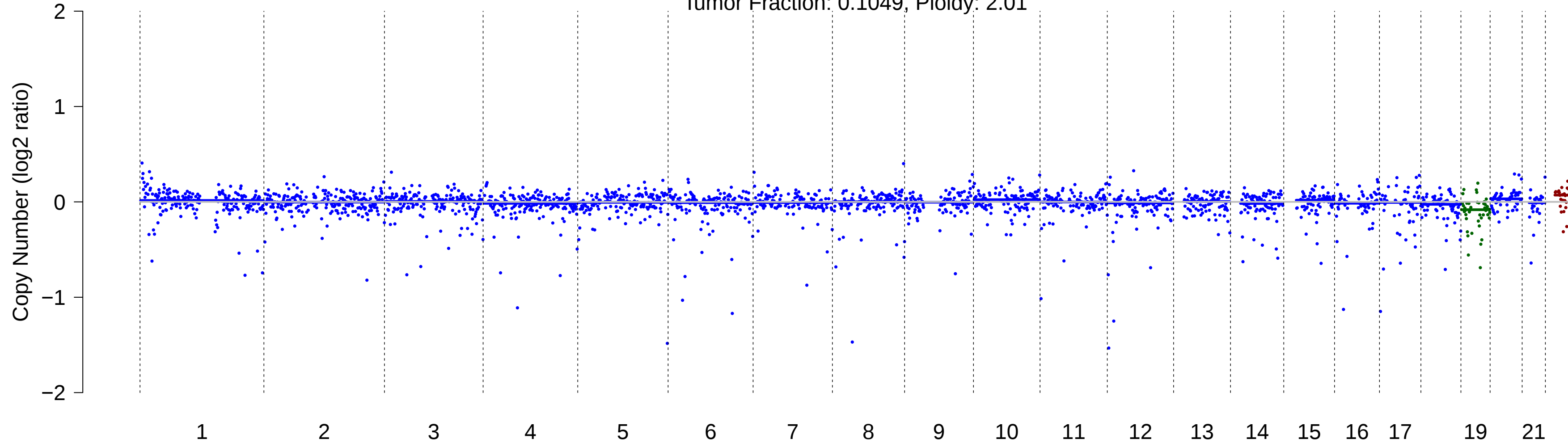


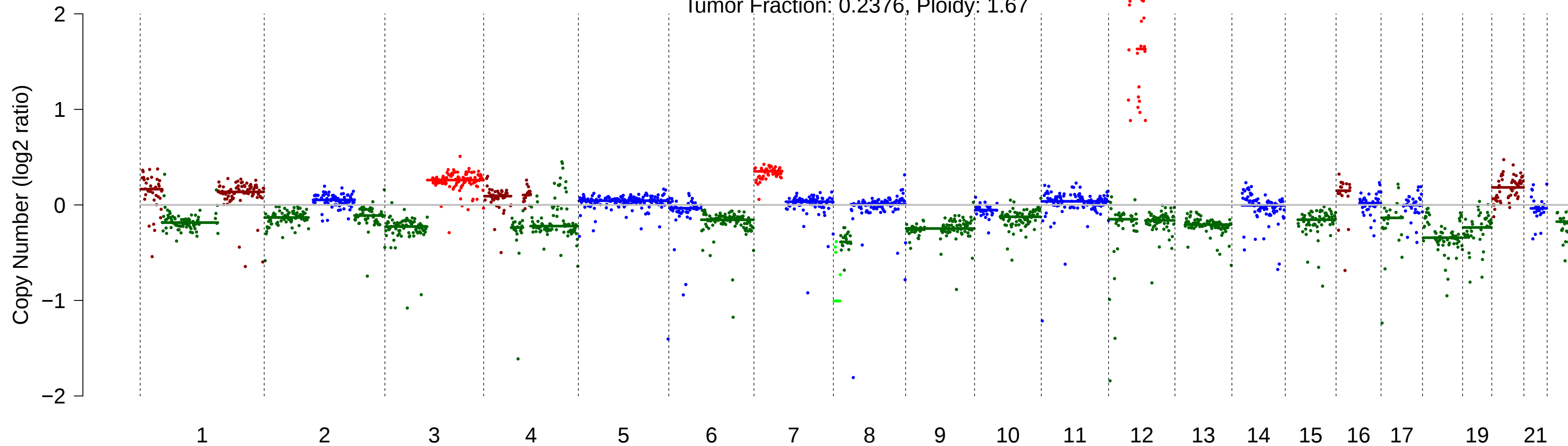
HU002_11.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2893

Tumor Fraction: 0.1049, Ploidy: 2.01



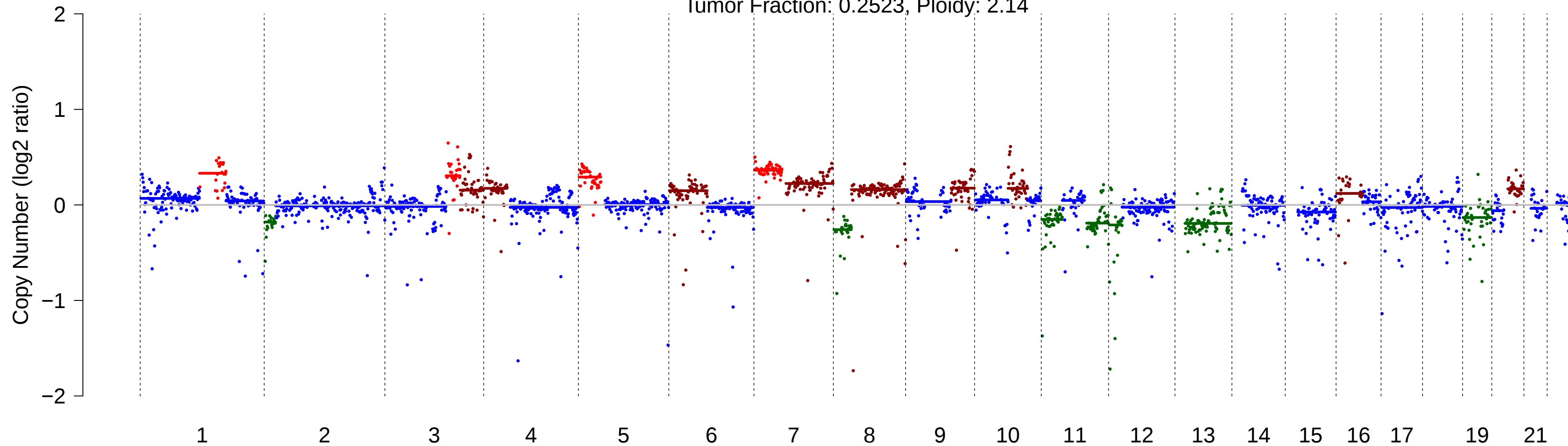
19_326.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2436

Tumor Fraction: 0.2376, Ploidy: 1.67



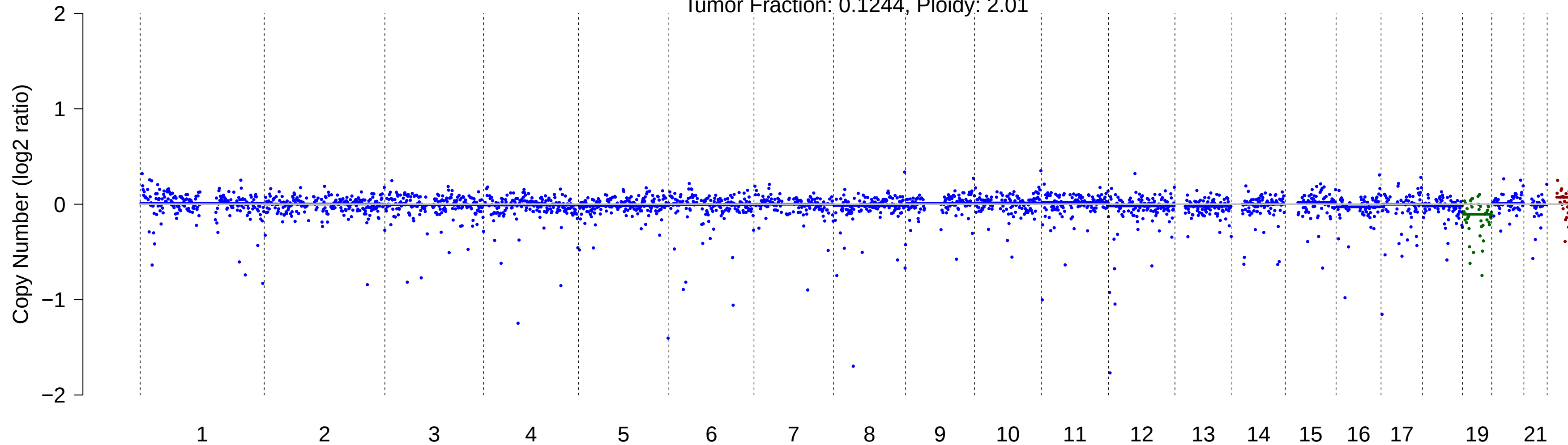
BC01.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2587

Tumor Fraction: 0.2523, Ploidy: 2.14



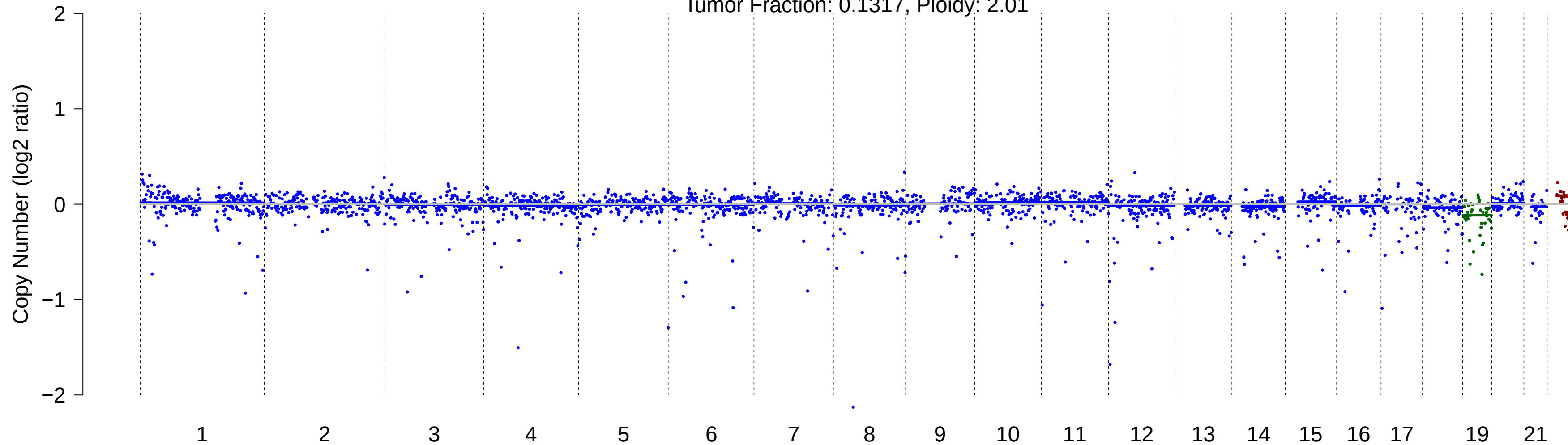
BC02.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 3016

Tumor Fraction: 0.1244, Ploidy: 2.01



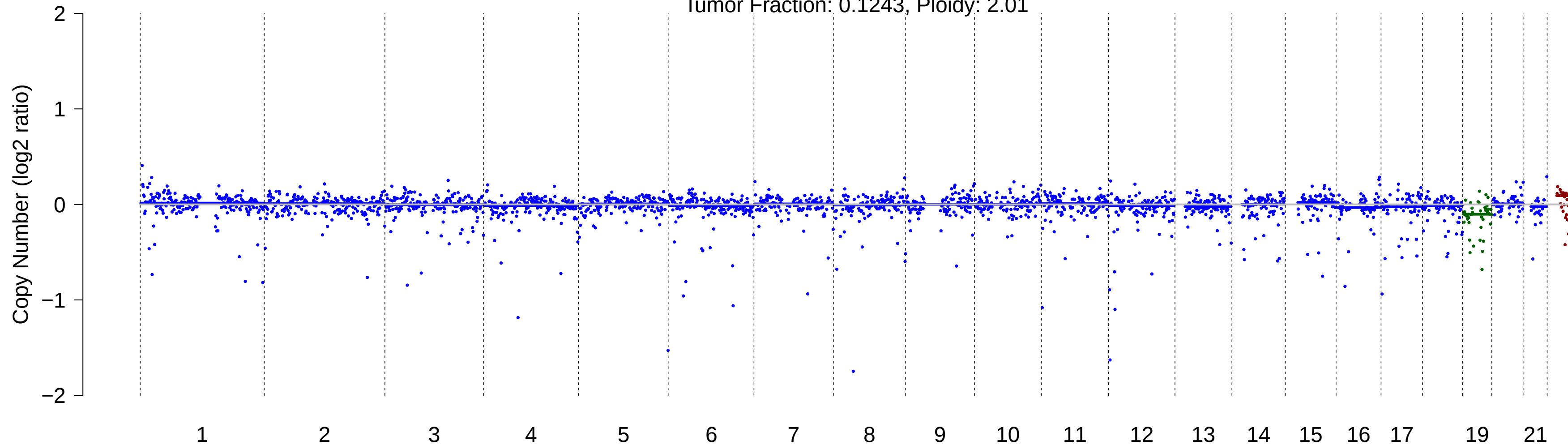
BC03.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2997

Tumor Fraction: 0.1317, Ploidy: 2.01



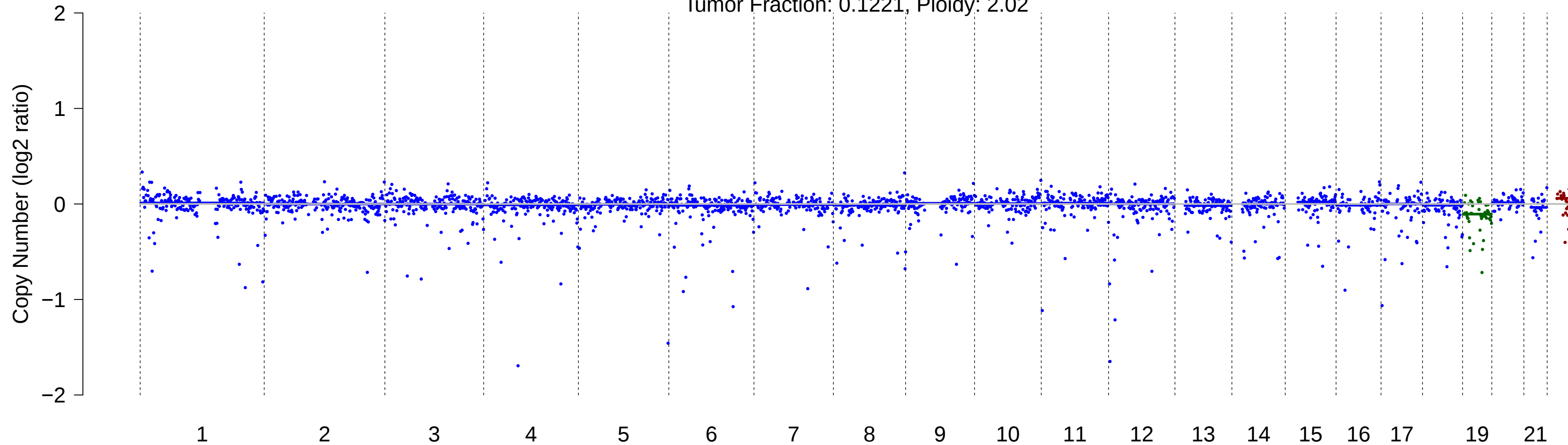
BC04.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 3109

Tumor Fraction: 0.1243, Ploidy: 2.01



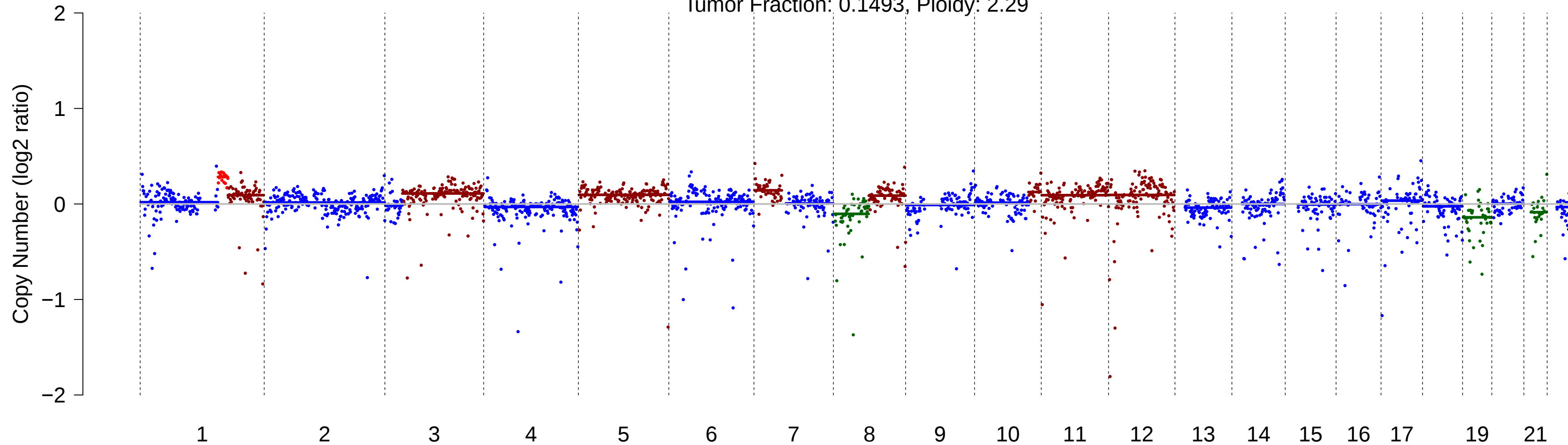
BC05.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 3324

Tumor Fraction: 0.1221, Ploidy: 2.02



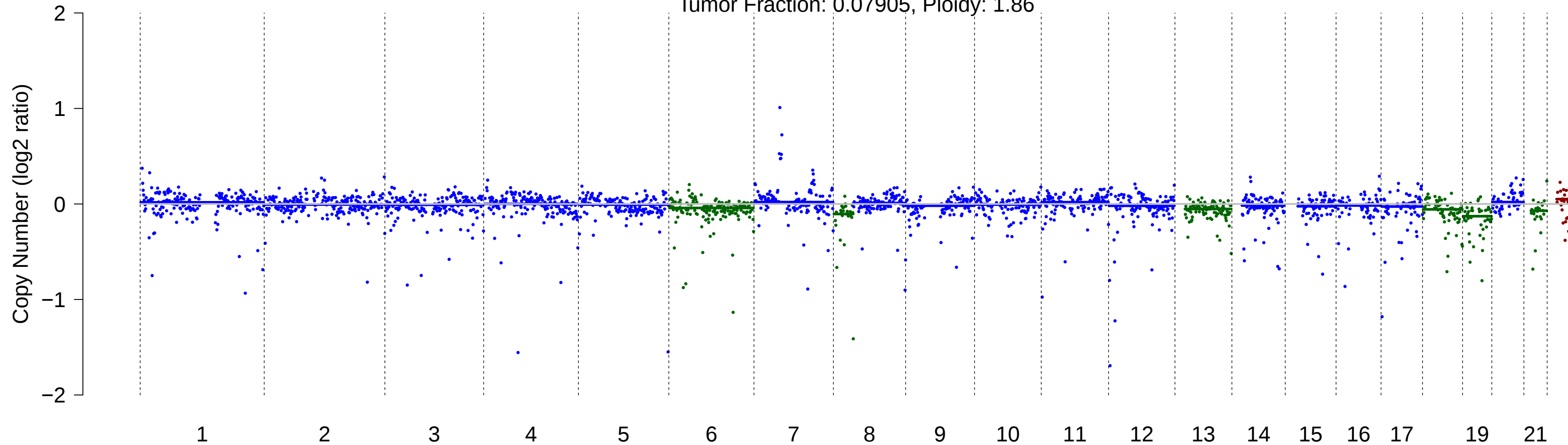
BC08.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2759

Tumor Fraction: 0.1493, Ploidy: 2.29



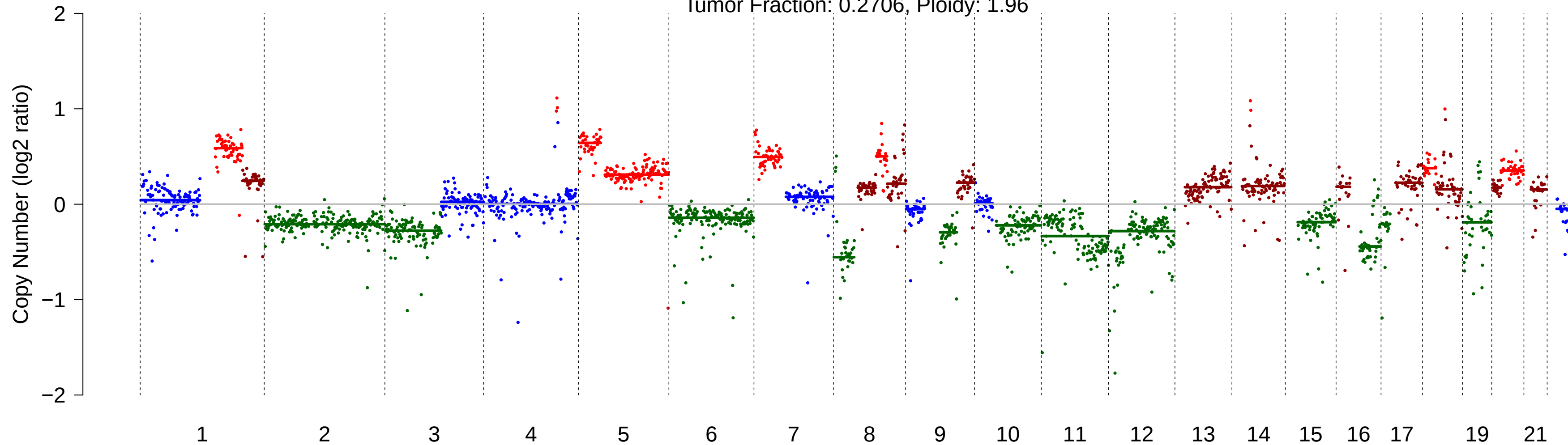
BC09.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2999

Tumor Fraction: 0.07905, Ploidy: 1.86



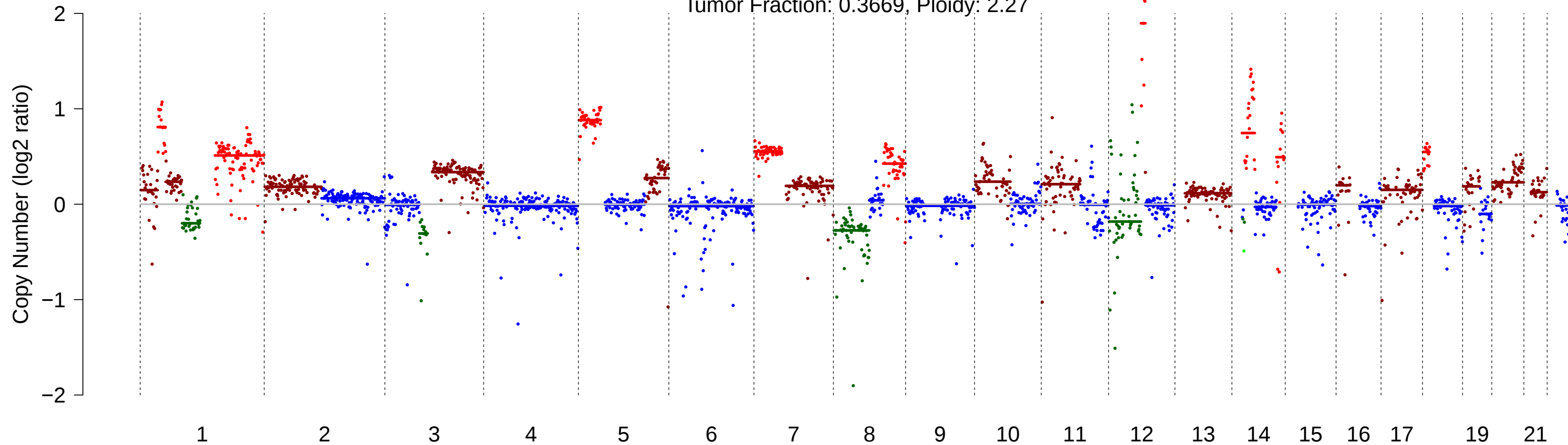
BC10.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2010

Tumor Fraction: 0.2706, Ploidy: 1.96



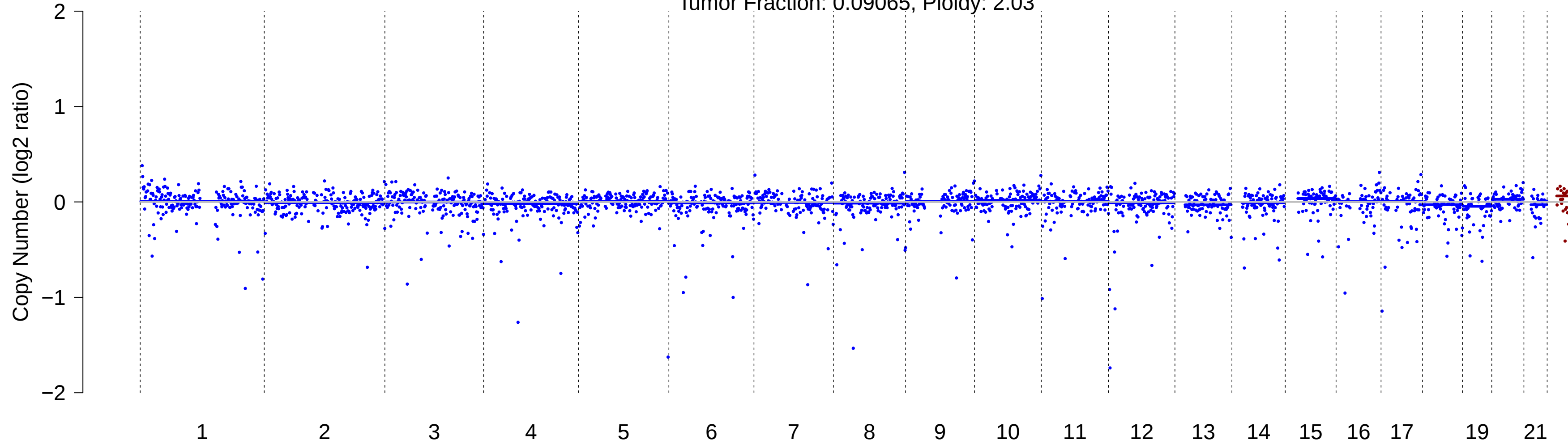
BC11.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2081

Tumor Fraction: 0.3669, Ploidy: 2.27



HU002_10.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2867

Tumor Fraction: 0.09065, Ploidy: 2.03



HU002_12.HAC.rehead.sort, n: 0.5, p: 2, log likelihood: 2627

Tumor Fraction: 0.08011, Ploidy: 2

