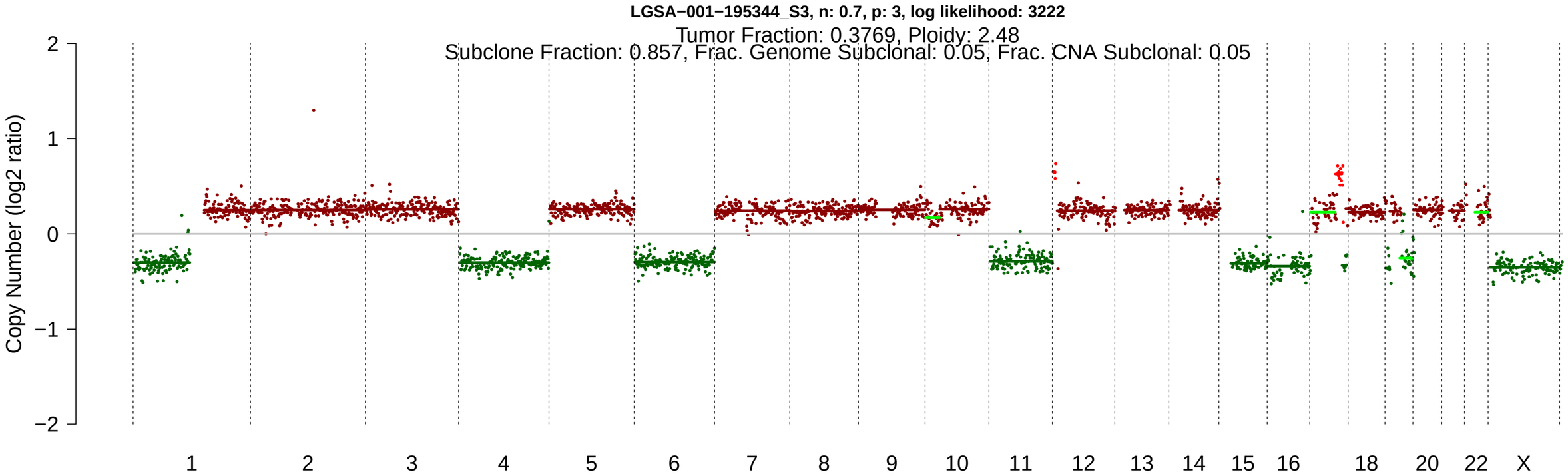
LGSA−001−195344_S3, n: 0.7, p: 3, log likelihood: 3222
Tumor Fraction: 0.3769, Ploidy: 2.48
Subclone Fraction: 0.857, Frac. Genome Subclonal: 0.05, Frac. CNA Subclonal: 0.05
Copy Number (log2 ratio)
2
1
0
−1
−2
1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
18
20
22
X