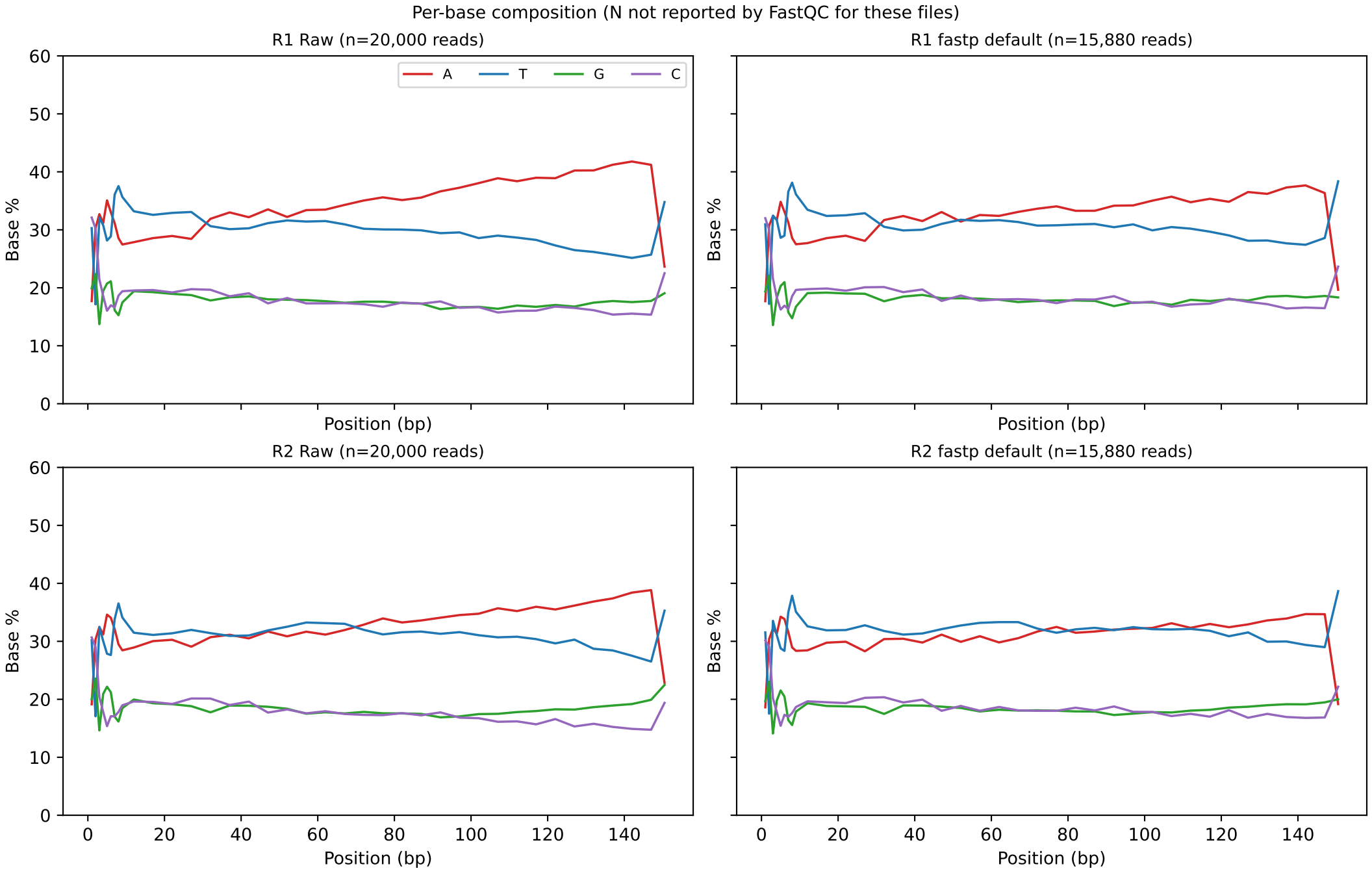
Per-base composition (N not reported by FastQC for these files)
R1 Raw (n=20,000 reads)
R1 fastp default (n=15,880 reads)
R2 Raw (n=20,000 reads)
R2 fastp default (n=15,880 reads)
A
T
G
C
Base %
Position (bp)
0
10
20
30
40
50
60
0
20
40
60
80
100
120
140