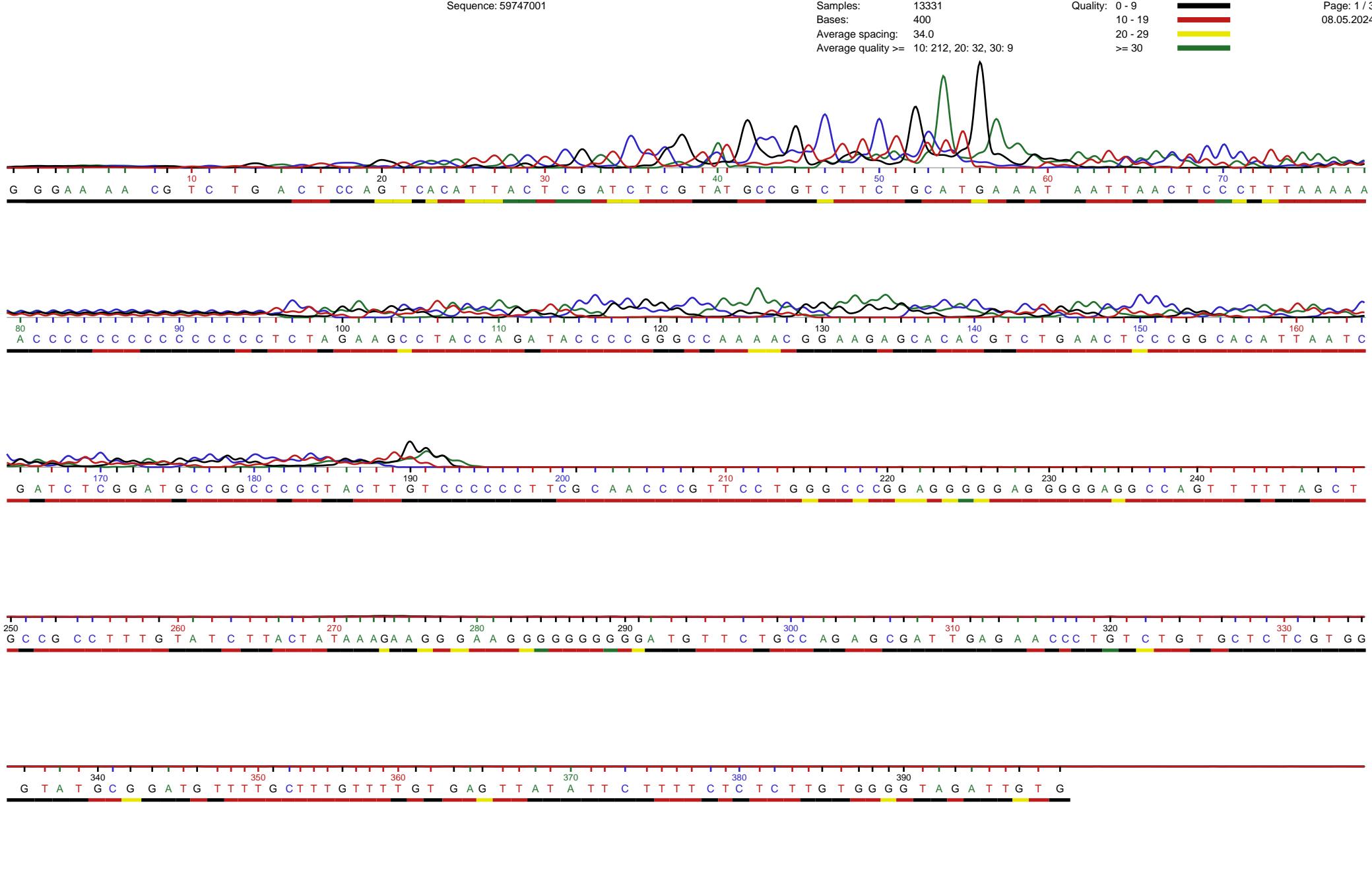Sequence: 59747001

Samples: 13331
Bases: 400
Average spacing: 34.0
Average quality >= 10: 212, 20: 32, 30: 9

Quality: 0 - 9
10 - 19
20 - 29
>= 30

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08.05.2024

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G G G A A  A A  C G  T C T G  A C T C C A G  T C A C A T  T A C T  C G A T C T C G  T A T  G C C  G T C T T C T  G C A T G A A A T  A A T T A A C T C C  C T T T A A A A A

A C C C C C C C C C  C C C C C C T C T A  G A A G C C T A C C A  G A T A C C C C  G G G  C C A A A A C  G G A A G A G C A C A C  G T C T G A A C T C C C G G  C A C A T T A A T C

G A T C T C G G A T G C C G G C C C C C T A C T T  G T  C C C C C C T T C G  C A A C C C G  T T C C T G G G  C C C G G A G G G G G A G  G G G G A G G C C A G T  T T T T A G C T

G C C G  C C T T T G  T A T C T T A C T A T A A A G A  A G G  G A A G G G G G G G G G G G A  T G T T C T G C C  A G A G C G A T  T G A G A A C C C  T G T C T G T  G C T C T C G T G G

G T A T G C G G A T G  T T T T G C T T T G T T T T  G T  G A G  T T A T A T T C  T T T T C T C T C T T G T G G G G  T A G A T T G T  G
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