

Used Linux Commands for file size

4. The suffix array and BWT are significantly more memory- and time-efficient than the uncompressed suffix trie, which takes ~60s and 528MB due to its $O(n^2)$ structure. In contrast, ****suffix arrays (~30MB)**** and ****BWT (~8KB)**** offer drastic performance gains, making them ideal for large-scale genomic analysis.

[illegible]

[illegible]

[illegible]

T1G1A1C2A2C1T1A1C1T1G1T2G1C1G1A1G1C1T1A1C1A1T2C1G1C1A1G1T4C1A2G2A2T
1C1G1T2C1G1A2T1G1A1T2A1C1G3C2A2T1A3C1G1c1g1t1c1t1a1t1c1a2c2t1g1t2a1t1a1g
1t1c2t3a1g1c1a2c1t1g1c1a2c1t2g2a1c1t1g3t1a1c3t2c1t1g1t2a2c1a3t1a1t2c4t2a1c1a2t
2T1G2C1T2A3T1A1C2T1A1T1G1T4A2T1C4T2A1C1T1A1G1A2C1G2C1A1C1T4C3t1c1t3a1
c5a1t1a1t4a1g4a1t1c1t1a1g4c2t1c3t1g2G1A1C1T1G1A1G1A1C2A4T1G1A2G2T2A1C1g1
c1a3g1t1g2c1t1g1t2a1t2c1t2a2g1G1A2G

- b) rle: 8868 sequence: 7920
- c) Original size: 7920 characters
 - a. Compressed size: 11714 characters
 - b. Compression ratio: 1.48
 - c. ✗ RLE compression was not effective.

Part 3:

1. Linear Search in 1.049896 seconds
2. BWT Search in 0.008646 seconds
3. BWT search as it is clearly faster.