

String Sorting in Python - Comparison of Several Algorithms

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TESTING DATA

| Dataset | Number | alphabet | Sum of | |
|----------------|------------|----------|-----------|--|
| Dalasei | of strings | size | LCP array | |
| dna.100MB | 618 | 15 | 4501 | |
| dna.200MB | 1114 | 15 | 8948 | |
| proteins.100MB | 359505 | 24 | 18853436 | |
| proteins.200MB | 709116 | 24 | 50076184 | |
| urls.100MB | 3284368 | 114 | 94113004 | |
| urls.200MB | 6576059 | 114 | 191545831 | |
| words.100MB | 18502734 | 211 | 83643408 | |
| words.200MB | 37003241 | 220 | 168115390 | |

Table 1: Data set used for comparing the algorithms

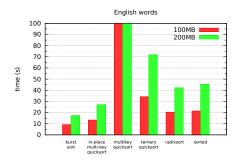
PERFORMANCE GRAPHS

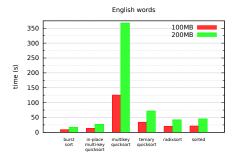
The graphs below show the time and space requirements of several algorithms on two texts. The algorithms are divided into three groups:

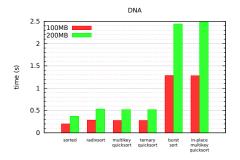
New algorithms based on reference point ranks, repetition shortcuts and wavelet trees

Improved implementations of wavelet trees and algorithms from [?]

Prior algorithms from [?, ?]







TEST RESULTS

| | sorted | MSD | Multikey | Ternary | Burst | In-place |
|----------------|------------------|------------|-----------|-----------|--------|--------------------|
| | (Python builtin) | Radix sort | quicksort | quicksort | sort | mulitkey quicksort |
| dna.100MB | 0.2 | 0.284 | 0.276 | 0.276 | 1.284 | 1.28 |
| dna.200MB | 0.372 | 0.532 | 0.52 | 0.52 | 2.44 | 2.484 |
| proteins.100MB | 0.768 | 7.024 | 7.2 | 1.908 | 8.705 | 4.252 |
| proteins.200MB | 1.532 | 20.921 | 23.301 | 3.272 | 24.67 | 10.793 |
| urls.100MB | 5.072 | 10.893 | 25.062 | 8.585 | 8.185 | 5.348 |
| urls.200MB | 10.601 | 21.641 | 64.836 | 16.921 | 16.245 | 11.697 |
| words.100MB | 21.449 | 20.357 | 125.384 | 34.182 | 9.193 | 13.313 |
| words.200MB | 45.311 | 42.147 | 367.687 | 71.788 | 17.361 | 27.09 |

Table 2: Running times for each algorithm with different data sources