

6. Strings

[String](#). Ordered list of characters.

[Ex.](#) Natural languages, Java programs, genomic sequences,

"The digital information that underlies biochemistry, cell biology, and development can be represented by a simple string G's, A's, T's and C's. This string is the root data structure of an organism's biology." -M. V. Olson

Using Strings in Java

[String concatenation](#). Append one string to end of another string.

[Substring](#). Extract a contiguous list of characters from a string.



```
String s = "strings";           // s = "STRINGS"
char c = s.charAt(2);          // c = 'R'
String t = s.substring(2, 7);   // t = "RINGS"
String u = s + t;              // u = "STRINGSRINGS"
```

String Implementation in Java

[Memory](#). 28 + 2N bytes for virgin string!

could use byte array instead of String to save space

```
public final class String implements Comparable<String> {
    private char[] value; // characters
    private int offset; // index of first char into array
    private int count; // length of string
    private int hash; // cache of hashCode

    private String(int offset, int count, char[] value) {
        this.offset = offset;
        this.count = count;
        this.value = value;
    }
    public String substring(int from, int to) {
        return new String(offset + from, to - from, value);
    }
    ...
}
```

String vs. StringBuilder

String. [immutable] Fast substring, slow concatenation.
StringBuilder. [mutable] Slow substring, fast append.

Radix Sorting

```
public static String reverse(String s) {
    String r = "";
    for (int i = s.length() - 1; i >= 0; i--)
        r += s.charAt(i);
    return r;
}
```

quadratic time

```
public static String reverse(String s) {
    StringBuilder r = new StringBuilder("");
    for (int i = s.length() - 1; i >= 0; i--)
        r.append(s.charAt(i));
    return r.toString();
}
```

linear time

Reference: Chapter 13, Algorithms in Java, 3rd Edition, Robert Sedgewick.

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Robert Sedgewick and Kevin Wayne • Copyright © 2005 • <http://www.Princeton.EDU/~cos226>

Radix Sorting

Radix sorting.

- Specialized sorting solution for strings.
- Same ideas for bits, digits, etc.

Applications.

- Sorting strings.
- Full text indexing.
- Plagiarism detection.
- Burrows-Wheeler transform. [see data compression]
- Computational molecular biology.

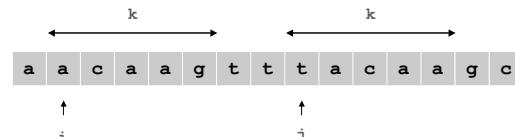
An Application: Redundancy Detector

Longest repeated substring.

- Given a string of N characters, find the longest repeated substring.
- Ex: a a c a a g t t t a c a a g c
- Application: computational molecular biology.

Dumb brute force.

- Try all indices i and j , and all match lengths k , and check.
- $O(W N^3)$ time, where W is length of longest match.



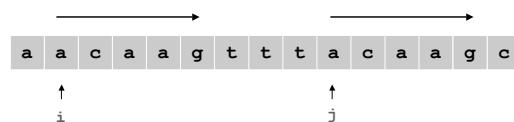
An Application: Redundancy Detector

Longest repeated substring.

- Given a string of N characters, find the longest repeated substring.
- Ex: **a a c a a g t t t a c a a g c**
- Application: computational molecular biology.

Brute force.

- Try all indices i and j for start of possible match, and check.
- $O(WN^2)$ time, where W is length of longest match.

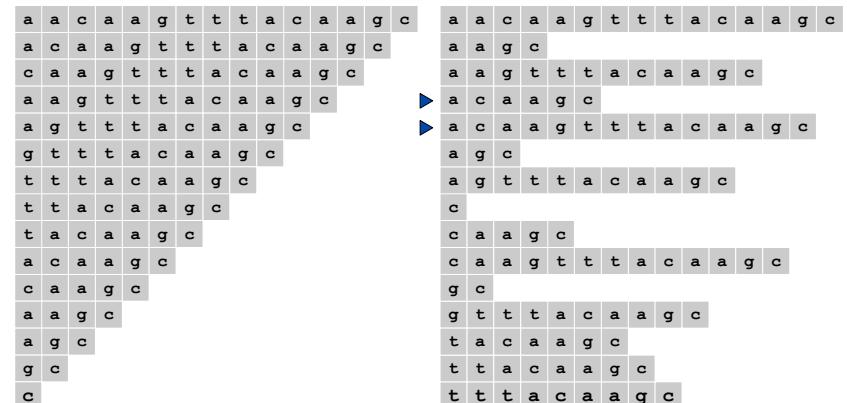


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A Sorting Solution

Suffix sort.

- Form N suffixes of original string.
- Sort to bring longest repeated substrings together.



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String Sorting

Notation.

- String = variable length sequence of characters.
- W = max # characters per string.
- N = # input strings.
- R = radix.

256 for extended ASCII, 65,536 for original UNICODE

Java syntax.

- Array of strings: `String[] a`
- Number of strings: `N = a.length`
- The i^{th} string: `a[i]`
- The d^{th} character of the i^{th} string: `a[i].charAt(d)`
- Strings to be sorted: `a[0], \dots, a[N-1]`

Suffix Sorting: Java Implementation

Java implementation.

```
public class LRS {
    public static void main(String[] args) {
        String s = StdIn.readAll();           read input
        int N = s.length();

        String[] suffixes = new String[N];
        for (int i = 0; i < N; i++)
            suffixes[i] = s.substring(i, N);

        Arrays.sort(suffixes);
        System.out.println(lcp(suffixes));
    }
}
```

create suffixes (linear time)

sort and find longest match (bottleneck)

longest common prefix of adjacent strings

```
% java LRS < moby dick.txt
,- Such a funny, sporty, gamy, jesty, jokey, hoky-poky lad, is the Ocean, oh! Th
```

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String Sorting Performance

	String Sort	Suffix (sec)
	Worst Case	Moby Dick
Brute	$W N^2$	36,000 \ddagger
Quicksort	$W N \log N \dagger$	9.5

W = max length of string.
 N = number of strings.

1.2 million for Moby Dick

\ddagger estimate
 \dagger probabilistic guarantee

Key Indexed Counting

Key indexed counting.

- Count frequencies of each letter. [0th character]

```
int N = a.length;
int[] count = new int[256+1];
for (int i = 0; i < N; i++) {
    char c = a[i].charAt(d);
    count[c+1]++;
}
```

frequencies

a	count
0	d a b
1	a d d
2	c a b
3	f a d
4	f e e
5	b a d
6	d a d
7	b e e
8	f e d
9	b e d
10	e b b
11	a c e

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Key Indexed Counting

Key indexed counting.

- Count frequencies of each letter. [0th character]
- Compute cumulative frequencies.

Key Indexed Counting

Key indexed counting.

- Count frequencies of each letter. [0th character]
- Compute cumulative frequencies.
- Use cumulative frequencies to rearrange strings.

```
for (int i = 0; i < N; i++) {
    char c = a[i].charAt(d);
    temp[count[c]++] = a[i];
}
```

rearrange

a	count	temp
0	d a b	a 0
1	a d d	b 2
2	c a b	c 5
3	f a d	d 6
4	f e e	e 8
5	b a d	f 9
6	d a d	g 12
7	b e e	
8	f e d	
9	b e d	
10	e b b	
11	a c e	

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```
for (int i = 1; i < 256; i++)
    count[i] += count[i-1];
```

cumulative counts

a	count
0	d a b
1	a d d
2	c a b
3	f a d
4	f e e
5	b a d
6	d a d
7	b e e
8	f e d
9	b e d
10	e b b
11	a c e

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Key Indexed Counting

Key indexed counting.

- Count frequencies of each letter. [0th character]
- Compute cumulative frequencies.
- Use cumulative frequencies to rearrange strings.

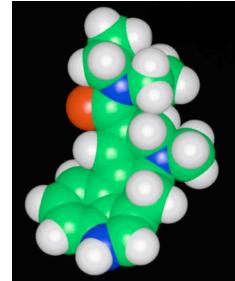
```
for (int i = 0; i < N; i++)
    a[i] = temp[i];
```

copy back

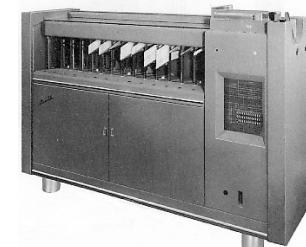
a	count	temp
0 a d d	a 2	0 a d d
1 a c e	b 5	1 a c e
2 b a d	c 6	2 b a d
3 b e e	d 8	3 b e e
4 b e d	e 9	4 b e d
5 c a b	f 12	5 c a b
6 d a b	g 12	6 d a b
7 d a d		7 d a d
8 e b b		8 e b b
9 f a d		9 f a d
10 f e e		10 f e e
11 f e d		11 f e d

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Least significant digit radix sort. Ancient method used for card-sorting.



Lysergic Acid Diethylamide, Circa 1960



Card Sorter, Circa 1960

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LSD Radix Sort

Least significant digit radix sort.

- Consider digits from right to left:
use key-indexed counting to **stable** sort by character

0 d a b	0 d a b	0 a c e
1 a d d	1 c a b	1 a d d
2 c a b	2 e b b	2 b a d
3 f a d	3 a d d	3 b e d
4 f e e	4 f a d	4 b e e
5 b a d	5 b a d	5 c a b
6 d a d	6 d a d	6 d a b
7 b e e	7 f e d	7 a d d
8 f e d	8 b e d	8 f e d
9 b e d	9 f e e	9 b e d
10 e b b	10 b e e	10 f e e
11 a c e	11 a c e	11 f e d

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LSD Radix Sort

Least significant digit radix sort.

- Consider digits from right to left:
use key-indexed counting to **stable** sort by character

```
public static void lsd(String[] a) {
    int W = a[0].length();
    for (int d = W-1; d >= 0; d--) {
        // do key-indexed counting sort on digit d
        ...
    }
}
```

Assumes fixed length strings (length = W)

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LSD Radix Sort: Correctness

Pf 1. [left-to-right]

- If two strings differ on first character, key-indexed sort puts them in proper relative order.
- If two strings agree on first character, stability keeps them in proper relative order.

Pf 2. [right-to-left]

- If the characters not yet examined differ, it doesn't matter what we do now.
- If the characters not yet examined agree, later pass won't affect order.

now	s	o	b	a	c
for	n	d	w	a	g
tip	a	g	o	a	o
ilk	l	w	j	a	g
dim	d	i	m	d	a
tag	t	a	c	b	c
jot	j	o	w	a	a
sob	s	u	e	c	e
nob	n	u	g	a	g
sky	e	g	g	g	g
hut	f	o	w	f	e
ace	f	e	e	e	ee
bet	f	e	r	f	or
men	g	i	g	g	ig
egg	h	u	t	h	ut
few	i	l	k	i	lk
jay	j	a	y	j	am
owl	j	a	m	j	ay
joy	j	o	t	j	ot
rap	j	o	y	j	oy
gig	m	e	n	m	en
wee	n	o	w	n	o
was	n	o	b	o	o
cab	o	w	l	o	wl
wad	r	a	p	r	ap
caw	s	o	sk	y	sk
cue	s	k	s	b	sob
fee	t	i	t	g	tag
tap	t	a	p	t	ap
ago	t	a	r	t	ar
tar	t	a	ti	p	ti
jam	w	e	w	d	wa
dug	w	as	w	s	as
and	w	ad	we	e	wee

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LSD Radix Sort Correctness

Running time. $\Theta(W(N + R))$.

why doesn't it violate $N \log N$ lower bound?

Advantage. Fastest sorting method for random fixed length strings.

Disadvantages.

- Accesses memory "randomly."
- Inner loop has a lot of instructions.
- Wastes time on low-order characters.
- Doesn't work for variable-length strings.
- Not much semblance of order until very last pass.

Goal. Find fast algorithm for variable length strings.

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MSD Radix Sort

Most significant digit radix sort.

- Partition file into 256 pieces according to first character.
- Recursively sort all strings that start with the same character, etc.

Q. How to sort on d^{th} character?

A. Use key-indexed counting.

now	a	c	e	a	c	e	a	c
for	a	g	o	a	g	o	a	g
tip	a	n	d	a	n	d	a	n
ilk	b	e	t	b	e	t	b	e
dim	c	a	b	c	a	b	c	a
tag	c	a	w	c	a	w	c	a
jot	c	u	e	c	u	e	c	u
sob	d	i	m	d	i	m	d	i
nob	d	u	g	d	u	g	d	u
sky	e	g	g	e	g	g	e	g
hut	f	o	w	f	o	w	f	o
ace	f	e	e	f	e	e	f	e
bet	f	e	r	f	e	r	f	e
men	g	i	g	g	i	g	g	i
egg	h	u	t	h	u	t	h	u
few	i	l	k	i	l	k	i	l
jay	j	a	y	j	a	y	j	a
owl	j	a	m	j	a	m	j	a
joy	j	o	t	j	o	t	j	o
rap	j	o	y	j	o	y	j	o
gig	m	e	n	m	e	n	m	e
wee	n	o	w	n	o	w	n	o
was	n	o	b	n	o	b	n	o
cab	o	w	l	o	w	l	o	w
wad	r	a	p	r	a	p	r	a
caw	s	o	sk	y	sk	y	o	sk
cue	s	k	s	b	s	b	k	s
fee	t	i	t	g	tag		i	t
tap	t	a	p	t	ap		a	p
ago	t	a	r	t	ar		a	r
tar	t	a	ti	p	ti		a	ti
jam	w	e	w	d	wa		e	w
dug	w	as	w	s	as		as	w
and	w	ad	we	e	wee		ad	we

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MSD Radix Sort Implementation

```
public static void msd(String[] a) {
    int N = a.length;
    msd(a, 0, N-1, 0);
}

private static void msd(String[] a, int l, int r, int d) {
    if (r <= l) return;

    // key-indexed counting sort on digit d of a[l] to a[r]
    int[] count = new int[256+1];
    ...

    // recursively sort 255 subfiles - assumes '\0' terminated
    for (int i = 0; i < 255; i++)
        msd(a, l + count[i], l + count[i+1] - 1, d+1);
}
```

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String Sorting Performance

	String Sort	Suffix (sec)
	Worst Case	Moby Dick
Brute	$W N^2$	36,000 \ddagger
Quicksort	$W N \log N \dagger$	9.5
LSD *	$W(N + R)$	-
MSD	$W(N + R)$	395

R = radix.

W = max length of string.

N = number of strings.

\ddagger estimate

* fixed length strings only

\dagger probabilistic guarantee

↑
1.2 million for Moby Dick

MSD Radix Sort: Small Files

Disadvantages.

- Too slow for small files.
 - ASCII: 100x slower than insertion sort for N = 2
 - UNICODE: 30,000x slower for N = 2
- Huge number of recursive calls on small files.

Solution. Cutoff to insertion sort for small N.

Consequence. Competitive with quicksort for string keys.

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String Sorting Performance

	String Sort	Suffix (sec)
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Quicksort	$W N \log N \dagger$	9.5
LSD *	$W(N + R)$	-
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MSD with cutoff	$W(N + R)$	6.8

R = radix.

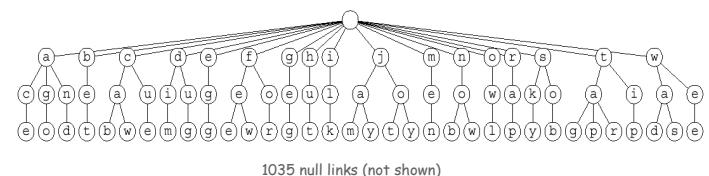
W = max length of string.

N = number of strings.

↑
1.2 million for Moby Dick

Recursive Structure of MSD Radix Sort

Trie structure. Describe recursive calls in MSD radix sort.



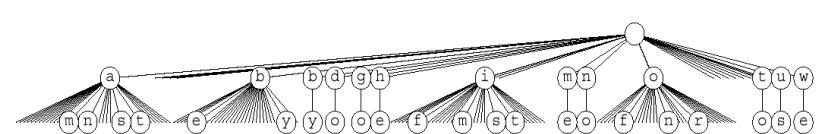
1035 null links (not shown)

Problem. Algorithm touches lots of empty nodes ala R-way tries.

- Tree can be as much as 256 times bigger than it appears!

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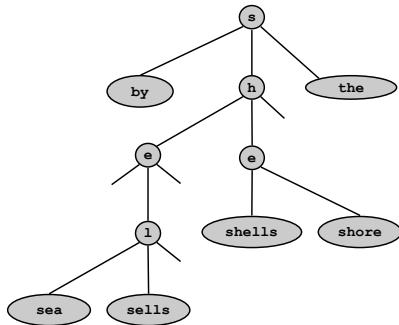
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Correspondence With Sorting Algorithms

Correspondence between trees and sorting algorithms.

- BSTs correspond to quicksort recursive partitioning structure.
- R-way tries corresponds to MSD radix sort.
- What corresponds to ternary search tries?



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3-Way Radix Quicksort

Idea 1. Use d^{th} character to "sort" into 3 pieces instead of 256, and sort each piece recursively.

Idea 2. Keep all duplicates together in partitioning step.

actinian	conobite	actinian
jeffrey	conelrad	bracteal
coenobite	actinian	conobite
conelrad	bracteal	conelrad
secureness	secureness	cumin
cumin	dilatedly	chariness
chariness	inkblot	centesimal
bracteal	jeffrey	cankerous
displease	displease	circumflex
millwright	millwright	millwright
repertoire	repertoire	repertoire
dourness	dourness	dourness
centesimal	southeast	southeast
fondler	fondler	fondler
interval	interval	interval
reversionary	reversionary	reversionary
dilatedly	cumin	secureness
inkblot	chariness	dilatedly
southeast	centesimal	inkblot
cankerous	cankerous	jeffrey
circumflex	circumflex	displease

3-way partition

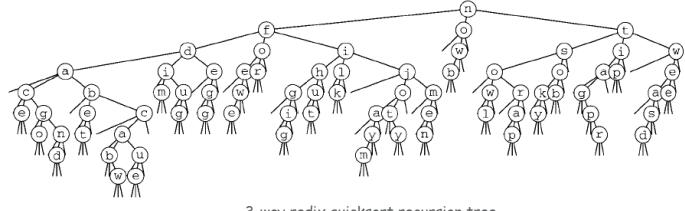
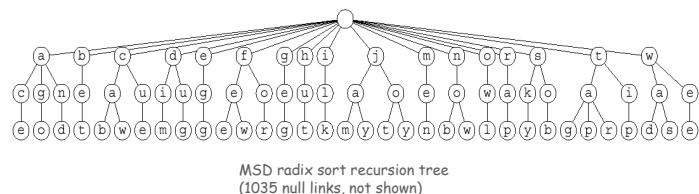
now	gig	ace	ago	ago
for	for	bet	bet	ace
tip	dug	dug	and	and
ilk	ilk	cab	ace	b et
dim	dim	dim	c ab	
tag	ago	ago	c ue	
jot	and	and	c ue	
sob	fee	egg	egg	
nob	cue	cue	dug	
sky	raw	dim		
hut	hut	fee		
ace	ace	for		
bet	bet	is ew		
man	man	lk		
egg	egg	gig		
few	few	hut		
jay	jay	jam		
owl	owl	jay		
joy	joy	joy		
rap	rap	jam	jo t	
gig	owl	owl	m en	
wad	wad	r ap		
cas	sky	sky	sky	
cue	now	we p	sob	
fee	sob	sob	tip	ta r
tap	tap	tap	t ap	ta p
ago	tag	tag	tag	tag
tar	tar	tar	tar	tar
dug	tip	tip	w as	
and	now	wee	wee	
jam	rap	wad	wad	

3-way radix quicksort

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Recursive Structure of MSD Radix Sort vs. 3-Way Quicksort

3-way radix quicksort collapses empty links in MSD tree.



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3-Way Radix Quicksort

```

private static void quicksortX(String a[], int lo, int hi, int d) {
    if (hi - lo <= 0) return;
    int i = lo-1, j = hi;
    int p = lo-1, q = hi;
    char v = a[hi].charAt(d);

    while (i < j) {
        repeat until pointers cross
        while (a[i+1].charAt(d) < v) if (i == hi) break; find i on left and
        while (v < a[--j].charAt(d)) if (j == lo) break; j on right to swap
        if (i > j) break;
        exch(a, i, j);
        if (a[i].charAt(d) == v) exch(a, ++p, i);
        if (a[j].charAt(d) == v) exch(a, j, --q);
    }
    if (p == q) {
        if (v != '\0') quicksortX(a, lo, hi, d+1);
        return;
    }
    if (a[i].charAt(d) < v) i++;
    for (int k = lo; k <= p; k++) exch(a, k, j--);
    for (int k = hi; k >= q; k--) exch(a, k, i++);
    quicksortX(a, lo, j, d);
    if ((i == hi) && (a[i].charAt(d) == v)) i++;
    if (v != '\0') quicksortX(a, j+1, i-1, d+1);
    quicksortX(a, i, hi, d);
}
  
```

sort 3 pieces recursively

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Quicksort vs. 3-Way Radix Quicksort

Quicksort.

- $2N \ln N$ string comparisons on average.
- Long keys are costly to compare if they differ only at the end, and this is common case!
- absolutism, absolut, absolutely, absolute.

3-way radix quicksort.

- Avoids re-comparing initial parts of the string.
- Uses just "enough" characters to resolve order.
- $2N \ln N$ character comparisons on average for random strings.
- Sub-linear sort for large W since input is of size NW .

Theorem. Quicksort with 3-way partitioning is OPTIMAL.

Pf. Ties cost to entropy. Beyond scope of 226.

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	String Sort	Suffix (sec)
	Worst Case	Moby Dick
Brute	$W N^2$	36,000 §
Quicksort	$W N \log N$ †	9.5
LSD *	$W(N + R)$	-
MSD	$W(N + R)$	395
MSD with cutoff	$W(N + R)$	6.8
3-way radix quicksort	$W N \log N$ †	2.8

R = radix.

W = max length of string.

N = number of strings.

§ estimate

* fixed length strings only

† probabilistic guarantee

1.2 million for Moby Dick

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Suffix Sorting: Worst Case Input

Length of longest match small.

- Hard to beat 3-way radix quicksort.

Length of longest match very long.

- 3-way radix quicksort is quadratic.
- Ex: two copies of Moby Dick.

Can we do better? $\Theta(N \log N)$? $\Theta(N)$?

Observation. Must find longest repeated substring while suffix sorting to beat N^2 .

```

abcdefghi
abcdeghiaabcdefghi
bcdefghi
bcdefghiaabcdefghi
cdefghi
cdefghiaabcdefgh
defghi
efghiabcdefghi
efghi
fghiabcdefghi
fghi
ghiabcdefghi
fhi
hiabcdefghi
hi
iabcdefghi
i

```

Input: "abcdefghiabcdefghi"

Suffix Sorting in Linearithmic Time: Key Idea

0 babaaaabcbabaaaaaa0	17 0babaaaabcbabaaaaaa
1 abaaaabcbabaaaaaa0b	16 a0babaaaabcbabaaaa
2 baaaabcbabaaaaaa0ba	15 aa0babaaaabcbabaaa
3 aaaabcbabaaaaaa0bab	14 aaa0babaaaabcbabaa
4 aaabcbabaaaaaa0babaa	3 aaaabcbabaaaaaa0bab
5 abcbbabaaaaaa0babaa	12 aaaa0babaaaabcbab
6 abcbabaaaaaa0babaaa	13 aaaa0babaaaabcbaba
7 bcbabaaaaaa0babaaaa	4 aaabcbabaaaaaa0babaa
8 cbabaaaaaa0babaaaab	5 aabcbabaaaaaa0babaa
9 babaaaaaa0babaaaabc	1 abaaaabcbabaaaaaa0b
10 abaaaaaa0babaaaabcb	10 abaaa0babaaaabcb
11 baaaaaa0babaaaabcba	6 abcbabaaaaaa0babaaa
12 aaaaa0babaaaabcbab	2 baaaabcbabaaaaaa0ba
13 aaaa0babaaaabcbaba	11 baaa0babaaaabcbab
14 aaa0babaaaabcbabaa	0 babaaaabcbabaaaaaa0
15 aa0babaaaabcbabaaa	9 babaaaa0babaaaabc
16 a0babaaaabcbabaaaa	7 bcbabaaaaaa0babaaaa
17 0babaaaabcbabaaaaaa	8 cbabaaaaaa0babaaaab

0 + 4 = 4

9 + 4 = 13

Input: "babaaaabcbabaaaaaa"

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Manber's MSD algorithm.

- Phase 0: sort on first character using key-indexed sorting.
- Phase i: given list of suffixes sorted on first 2^{i-1} characters, create list of suffixes sorted on first 2^i characters
- Finishes after $\lg N$ phases.

Manber's LSD algorithm.

- Same idea but go from right to left.
- $O(N \log N)$ guaranteed running time.
- $O(N)$ extra space (but need several auxiliary arrays).

Best in theory. $O(N)$ but more complicated to implement.

	String Sort	Suffix Sort (seconds)	
	Worst Case	Moby Dick	AesopAesop
Brute	$W N^2$	36,000 §	3,990 §
Quicksort	$W N \log N$ †	9.5	167
LSD *	$W(N + R)$	-	-
MSD	$W(N + R)$	395	memory
MSD with cutoff	$W(N + R)$	6.8	162
3-way radix quicksort	$W N \log N$ †	2.8	400
Manber ‡	$N \log N$	17	8.5

R = radix.

W = max length of string.

N = number of strings.

§ estimate

* fixed length strings only

† probabilistic guarantee

‡ suffix sorting only

1.2 million for Moby Dick
191 thousand for Aesop's Fables