

Regularities in Sequences

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Chennai, December 1999

Regularities in Sequences

or a longer title...

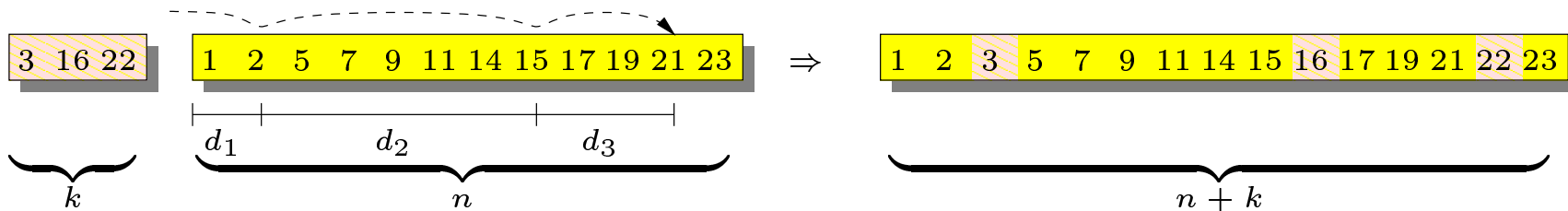
**On Using Efficient Merging in
 $\mathcal{O}(n \log n)$ Time Algorithms for
Finding Regularities in Strings**

Merging $\rightarrow \mathcal{O}(n \log n)$ Algorithms \rightarrow Strings \rightarrow Regularities

Comparison Based Merging

Problem

Merge two sorted sequences of length k and n , $k \leq n$



Solutions

- (1) Sequential merging $\Rightarrow \mathcal{O}(n + k)$ comparisons
- (2) Merging using exponential searching \Rightarrow

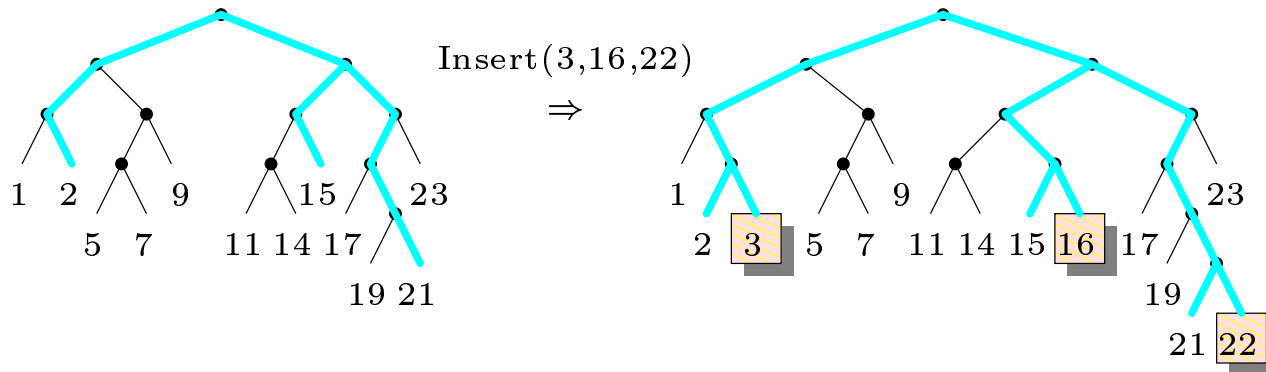
$$\Theta \left(\sum_{i=1}^k \log d_i \right) = \Theta \left(\log \binom{n+k}{k} \right) \text{ comparisons}$$

Hwang & Lin 1972

Time ?

Merging Search Trees

Brown & Tarjan 1979



Merging two balanced search trees of size k and n , $k \leq n$

- Size k tree \Rightarrow list of k elements $\mathcal{O}(k)$
 - Search for the k elements in **one traversal** of the size n tree $\mathcal{O}\left(\left|\bigcup_{i=1}^k \text{search-path}_i\right|\right)$
 - Insert the k elements $\mathcal{O}(k)$
 - Rebalance typically amortized $\mathcal{O}(k)$
- U paths ?

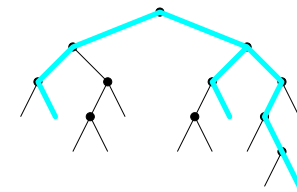
Path Theorem

Brown & Tarjan 1979

Theorem

In a **balanced** search tree with n leaves the #edges in the union of k distinct root-to-leaf paths is

$$\left| \bigcup_{i=1}^k \text{path}_i \right| = \mathcal{O} \left(\log \binom{n+k}{k} \right)$$



[e.g. AVL-trees, red-black trees, 2-3-trees, 2-4-trees, ...]

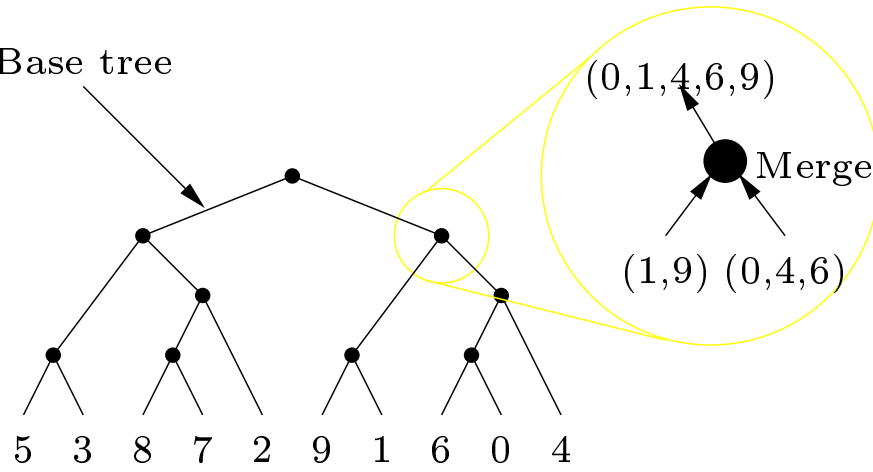
Corollary

Merging two trees of size k and n , $k \leq n$, takes time $\mathcal{O} \left(\log \binom{n+k}{k} \right)$ which is optimal

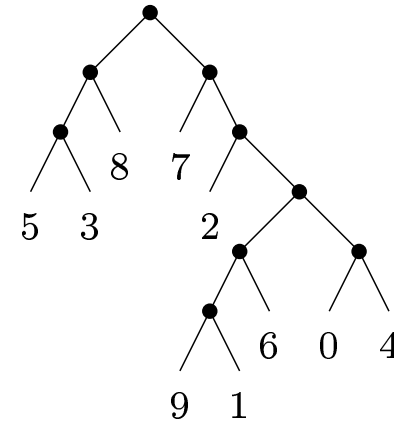
Merge Sort ?

Merge Sort — Revisited

Base tree



Balanced base tree



Unbalanced base tree

Balanced base tree	+	Sequential merging	\Rightarrow	$\mathcal{O}(n \log n)$
Unbalanced base tree	+	Sequential merging	\Rightarrow	$\mathcal{O}(n^2)$
Unbalanced base tree	+	Merging by insertion*	\Rightarrow	$\mathcal{O}(n \log^2 n)$
Unbalanced base tree	+	Optimal merging	\Rightarrow	$\mathcal{O}(n \log n)$

$n \log n ?$

Merge Sort — Analysis

Theorem

If we spend time $\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$ at each node with two children spanning respectively n_1 and n_2 leaves, $n_1 \leq n_2$, then we in a subtree spanning n leaves spend total time $\mathcal{O}(\log n!)$

Proof

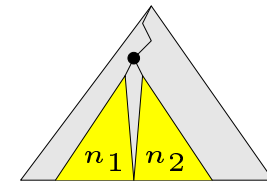
Induction, using

$$\log n_1! + \log n_2! + \log \binom{n_1 + n_2}{n_1}$$

$$= \log n_1! + \log n_2! + \log \frac{(n_1 + n_2)!}{n_1! \cdot n_2!}$$

$$= \log n_1! + \log n_2! + \log(n_1 + n_2)! - \log n_1! - \log n_2!$$

$$= \log(n_1 + n_2)! \quad \square$$



Corollary

Merging n singleton lists using any base tree takes time $\mathcal{O}(n \log n)$

Applications of the Path Lemma

Searching

Searching for the predecessors of $e_1 \leq \dots \leq e_k$ in a balanced search tree of size $n \geq k$ takes time $\mathcal{O}\left(\log \binom{n+k}{k}\right)$

Δ -Searching

The δ -predecessor of an x in a sorted list $L = (x_1, \dots, x_n)$ is defined by

$$\text{max-gap}(L) = \max\{0, x_2 - x_1, \dots, x_n - x_{n-1}\}$$

$$\Delta\text{-pred}(L, \delta, x) = \min\{y \in L \mid \text{max-gap}(L \cap [y, x]) \leq \delta\}$$

Ex. $\Delta\text{-pred}((1, 2, \underline{5}, 7, 9, 11, 14, 15, 17, 19, 21, 23), 2, 13) = 5$

- $\text{Multi-}\Delta\text{-Pred}(T, \delta, e_1, \dots, e_k)$ for each e_i finds $\Delta\text{-pred}(T, \delta, e_i)$

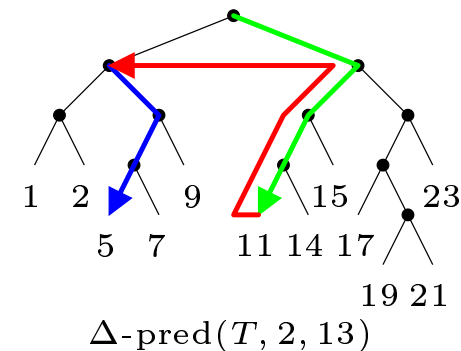
Δ -Pred ?

Δ -Searching

For each node v store $\min(T_v)$, $\max(T_v)$, and $\max\text{-gap}(T_v)$

The computation of $\Delta\text{-pred}(T, \delta, e)$ proceeds in three phases

- compute $\text{pred}(T, e)$
- search for $\Delta\text{-pred}(T, \delta, \min(T_v))$
- search for $\Delta\text{-pred}(T_v, \delta, \max(T_v))$

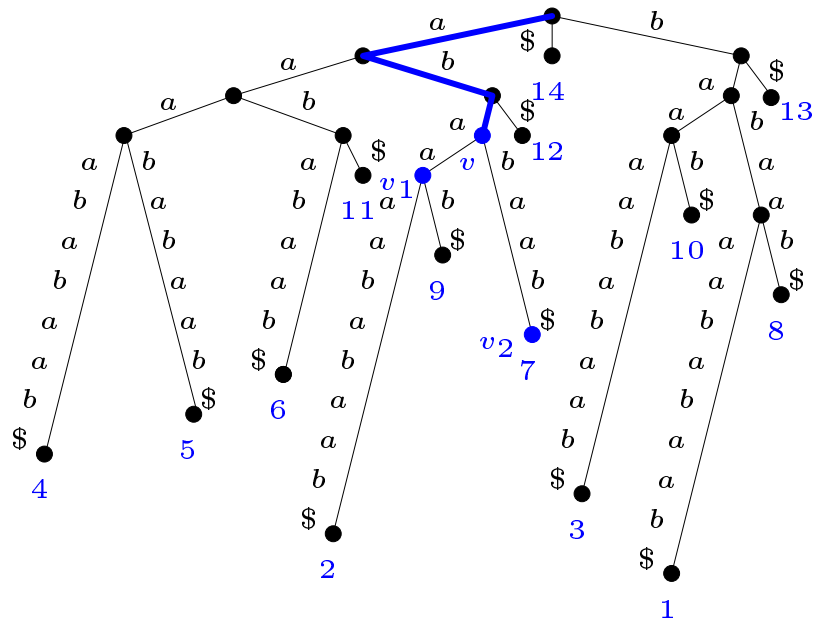


Multi- Δ -Pred($T, \delta, e_1, \dots, e_k$) can be performed in time $\mathcal{O}\left(\log \binom{n+k}{k}\right)$ when $e_1 \leq \dots \leq e_k$ and $k \leq n$, e.g. using dynamic programming to only compute $\Delta\text{-pred}(T, \delta, \min(T_v))$ and $\Delta\text{-pred}(T, \delta, \max(T_v))$ once per node

String Notation

Notation		Example	
Alphabet	Σ	$\Sigma = \{a, b\}$	
String	$S \in \Sigma^*$	$S = b a b a a a a b a b a a b$	
i th character	$S[i]$	$S[3] = b$	
Substring	$S[i..j]$	$S[3..8] = b a a a a b$	
Occurrence	(α, i)	$(b a b, 8) = b a b a a a a \underline{b a b} a a b$	
Tandem repeat	$\alpha\alpha$	$\alpha\alpha = \underline{b a} \underline{b a}$	
Pair	(α, i, j)	$(b a b a, 1, 8) = \underline{b a b a} a a a \underline{b a b a} a b$	

Path Labels and Leaf Lists



$S = b \underline{a} b a a a \underline{a} b \underline{a} b a a b$
 $L(v) = a b a$
 $LL(v_1) = \{2, 9\}$
 $LL(v_2) = \{7\}$
 $LL(v) = LL(v_1) \cup LL(v_2) = \{2, 7, 9\}$

Let v be a node in the suffix tree of S

Path-label $L(v)$ = string spelled from the root to v

Leaf-list $LL(v)$ = set of leaf labels in the subtree rooted at v

$$LL(v) = \text{all occurrences of } L(v) = \bigcup_{c \in \text{children}(v)} LL(c)$$

Regularities

We will develop $\mathcal{O}(n \log n + |\text{output}|)$ time algorithms for

- Finding all **tandem repeats** in S

$b a b a a a a \underline{b a} \underline{b a} a b$

- Finding all **maximal pairs** in S with bounded gap

$\underline{b a b a a} \underbrace{a a}_{\text{gap}} \underline{b a b a a} b$

- Finding all **maximal quasi-periodic substrings** in S

$a a a b a a b b b a b a a b a a b a a b b a a a b a$

$\overline{\hspace{15em}}$
 $\underline{\hspace{15em}}$

Tandem Repeats

Stoye & Gusfield 1998

Problem

Given S , find all substrings of S which are tandem repeats

For each tandem repeat occurrence $(\alpha\alpha, i)$ output $(|\alpha|, i)$

Ex. $S = \underline{b a b a a a} \underline{a b a b} \underline{a a b}$

Output = $(2, 1), (1, 4), (1, 5), (1, 6), (2, 7), (2, 8), (1, 11)$

Tandem repeat occ. $(|\alpha|, i)$ is **branching** iff $S[i + |\alpha|] \neq S[i + 2|\alpha|]$

Ex. $(2, 7)$ is non-branching and $(2, 8)$ is branching

$(|\alpha|, i)$ non-branching $\Rightarrow (|\alpha|, i + 1)$ tandem repeat occ.

Given all branching tandem repeat occ. we can compute all tandem repeat occ. in time $\mathcal{O}(|\text{output}|)$

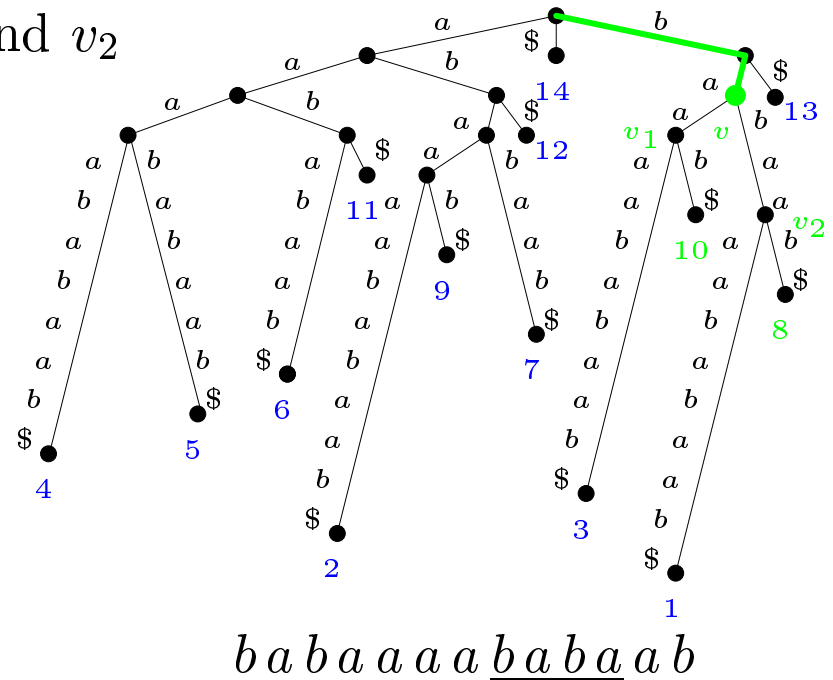
Branching ?

Branching Tandem Repeat Occurrences

Lemma

$(\alpha\alpha, i)$ is a branching tandem repeat occurrence in S iff

- \exists node v in the suffix tree of S with $L(v) = \alpha$
- v has two distinct children v_1 and v_2
- $i \in LL(v_1)$ and $i + |\alpha| \in LL(v_2)$



Branching Tandem Repeat Occurrences

Algorithm

1. Compute suffix tree (w.l.o.g. all nodes \leq two children) $\mathcal{O}(n)$
2. Bottom-up do for each node v with children v_1 and v_2
 $n_1 = |LL(v_1)| \leq |LL(v_2)| = n_2$
 - a) For each $i \in LL(v_1)$ search for $i \pm |L(v)|$ in $LL(v_2)$
 using optimal multi-search $\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$
 If $i + |L(v)| \in LL(v_2)$ then output $(|L(v)|, i)$
 If $i - |L(v)| \in LL(v_2)$ then output $(|L(v)|, i - |L(v)|)$
 - b) $LL(v) = LL(v_1) \cup LL(v_2)$
 using optimal tree merging $\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$

Total time $\mathcal{O}(n \log n)$, i.e. $\mathcal{O}(n \log n)$ branching tandem repeat occ.

Maximal Pairs with Bounded Gap

Brodal et al. 1999

A pair (α, i, j) is

right-maximal iff $S[i + |\alpha|] \neq S[j + |\alpha|]$

left-maximal iff $S[i - 1] \neq S[j - 1]$

maximal iff both right- and left-maximal

The **gap** of $(\alpha, i, j) = \max(i, j) - \min(i, j) - |\alpha|$

Ex. $b \underline{a b} a a a \underline{a b} a b a a b$

$(a b, 2, 7)$ is left-maximal but not right-maximal

$(a b a, 2, 7)$ is maximal and $\text{gap} = 2$

Problem

Given S , find all maximal pairs in S with gap in an interval $[\Delta_L, \Delta_H]$

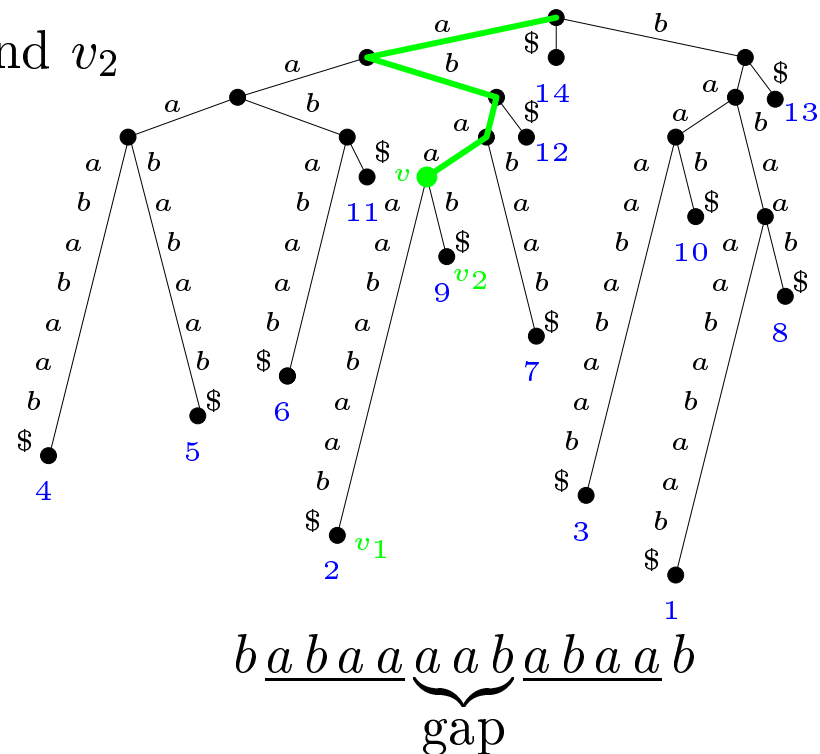
For each such pair (α, i, j) output $(|\alpha|, i, j)$

Right-Maximal Pairs

Lemma

(α, i, j) is a right-maximal pair in S iff

- \exists node v in the suffix tree of S with $L(v) = \alpha$
- v has two distinct children v_1 and v_2
- $i \in LL(v_1)$ and $j \in LL(v_2)$



Right-Maximal Pairs with Bounded Gap

Algorithm

1. Compute suffix tree (w.l.o.g. all nodes \leq two children) $\mathcal{O}(n)$

2. Bottom-up do for each node v with children v_1 and v_2

$$n_1 = |LL(v_1)| \leq |LL(v_2)| = n_2$$

a) For each $i \in LL(v_1)$ search for $i \pm (|L(v)| + \Delta_H)$ in $LL(v_2)$

$$\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$$

$j \in [i+|L(v)|+\Delta_L .. i+|L(v)|+\Delta_H] \cap LL(v_2)$ output $(|L(v)|, i, j)$

$j \in [i-|L(v)|-\Delta_H .. i-|L(v)|-\Delta_L] \cap LL(v_2)$ output $(|L(v)|, j, i)$

$$\mathcal{O}(|\text{output}|)$$

b) $LL(v) = LL(v_1) \cup LL(v_2)$

$$\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$$

Total time $\mathcal{O}(n \log n + |\text{output}|)$

Maximal ?

Maximal Pairs with Bounded Gap

Simple approach

- Compute all right-maximal pairs $(|\alpha|, i, j)$ with bounded gap
 - Output all $(|\alpha|, i, j)$ where $S[i - 1] \neq S[j - 1]$
- ... but requires too much time

Efficient approach

Divide the leaf-lists into blocks of contiguous indices with same left-character

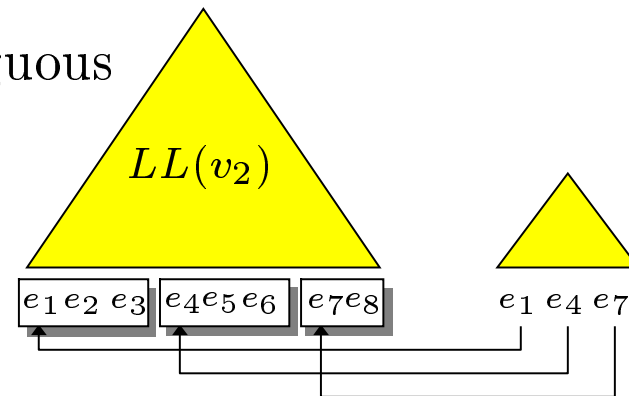
For each $LL(v)$ keep a tree with the starting points of each block

In the computation of

$$\{j \in [i + |L(v)| + \Delta_L .. i + |L(v)| + \Delta_H] \cap LL(v_2) : S[j - 1] \neq S[i - 1]\}$$

a block of j 's where $S[j - 1] = S[i - 1]$ can be skipped in time $\mathcal{O}(1)$

Total time $\mathcal{O}(n \log n + |\text{output}|)$



Finding Maximal Quasi-Periodic Substrings

Algorithm

1. Mark all v where $L(v)$ not quasi-periodic $\mathcal{O}(n \log n)$
The most complicated step — applies multi-searches and merging
2. Bottom-up do for each node v with children v_1 and v_2
 $n_1 = |LL(v_1)| \leq |LL(v_2)| = n_2$
 - a) Compute $LL(v) = LL(v_1) \cup LL(v_2)$ $\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$
 - b) For each coalescing run in $LL(v)$ identify at least one
 $i \in LL(v_1)$ in the run $\mathcal{O}(n_1)$
 - c) Apply Multi- Δ -Pred($LL(v), |L(v)|, \dots$) to the result of b) to
find the start and end of each coalescing run $\mathcal{O}\left(\log \binom{n_1+n_2}{n_1}\right)$
 - d) If v is marked then report all the coalescing runs found in c)
as maximal quasi-periodic substrings

Total time $\mathcal{O}(n \log n)$, i.e. $\mathcal{O}(n \log n)$ maximal quasi-periodic substrings

Theorems Applied

Theorem

In a **balanced** search tree with n leaves the #edges in the union of k distinct root-to-leaf paths is

$$\left| \bigcup_{i=1}^k \text{path}_i \right| = \mathcal{O} \left(\log \binom{n+k}{k} \right)$$

Theorem

If we spend time $\mathcal{O} \left(\log \binom{n_1+n_2}{n_1} \right)$ at each node with two children spanning respectively n_1 and n_2 leaves, $n_1 \leq n_2$, then we in a subtree spanning n leaves spend total time $\mathcal{O}(\log n!)$

Conclusion

- We have presented $\mathcal{O}(n \log n + |\text{output}|)$ time algorithms for finding all tandem repeats, maximal pairs with bounded gap, and maximal quasi-periodic substrings of a string
- Based on efficient merging and two related theorems
- The data structuring approach allowed us to concentrate on a few crucial underlying string properties

Open problem

- Can the time be improved to $o(n \log n)$ for finding all pairs with bounded gap or all maximal quasi-periodic substrings ?

Without an upper bound on the gaps all pairs can be found in time $\mathcal{O}(n + |\text{output}|)$

Brodal et al. 1999

A compact representation of all tandem repeats can be found in time $\mathcal{O}(n)$

Gusfield & Stoye 1998