

Approximate Cartesian Tree Matching: an Approach Using Swaps

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Foreword

- It's about pattern matching in time series.
- Motivations:
share prices in stock markets, temperatures, notes in music, GST data in bioinformatics...
- In recent years, new pattern matching problems such as Order-Preserving Matching and Cartesian Tree Matching have been introduced.
- To the best of our knowledge, no approximate pattern matching problem existed in the Cartesian tree framework.

Outline

- 1 Preliminaries
- 2 Characterization
- 3 Swap Graph
- 4 Conclusion

Preliminaries

Prerequisites

- Sequences of integers
- A total order $<$
- All elements of x are distinct and numbered from 1 to n (the length of x)

Cartesian tree matching (1)

Cartesian tree [Vuillemin, 1980]

A sequence x of length n can be associated to its Cartesian tree $C(x)$ according to the following rules:

- if x is empty, then $C(x)$ is the empty tree;
- if $x[1 \dots n]$ is not empty and $x[i]$ is the smallest value of x , $C(x)$ is the Cartesian tree with:
 - i as its root,
 - $C(x[1 \dots i - 1])$ as the left subtree,
 - $C(x[i + 1 \dots n])$ as the right subtree.

Cartesian tree matching (1)

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 - i as its root,
 - $C(x[1 \dots i - 1])$ as the left subtree,
 - $C(x[i + 1 \dots n])$ as the right subtree.

NB: In our examples, we will label the nodes with the values instead of the indices

Cartesian tree matching (1)

x 4 5 6 2 1 7 8 3 9

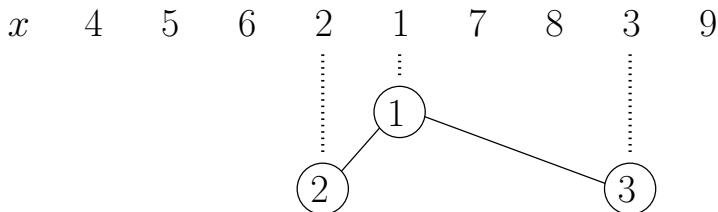
Cartesian tree matching (1)

x 4 5 6 2 1 7 8 3 9

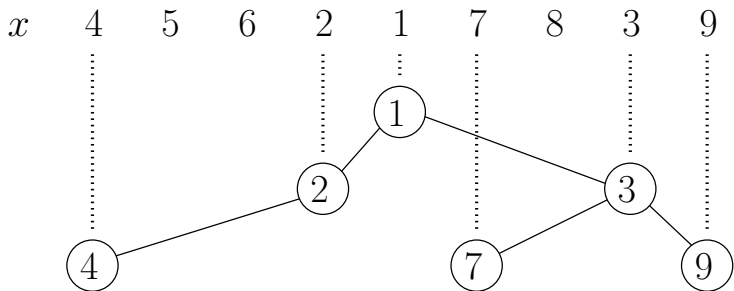
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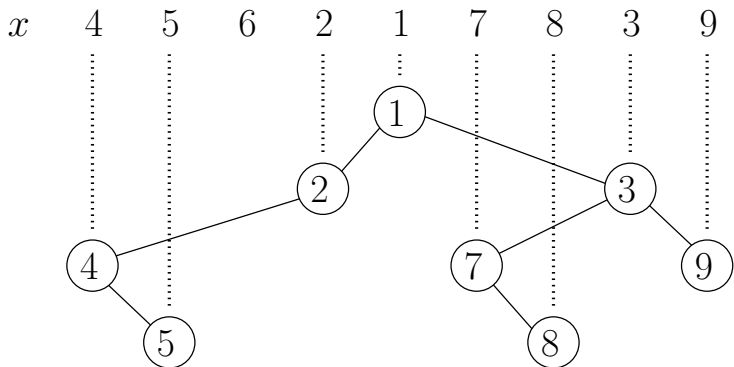
Cartesian tree matching (1)



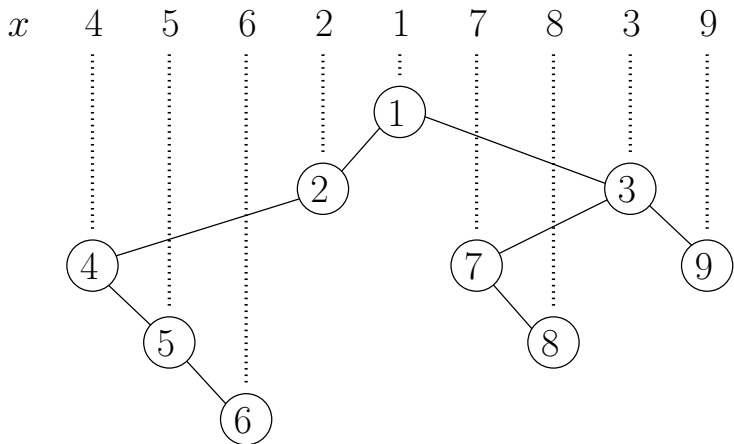
Cartesian tree matching (1)



Cartesian tree matching (1)



Cartesian tree matching (1)



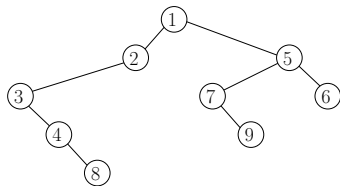
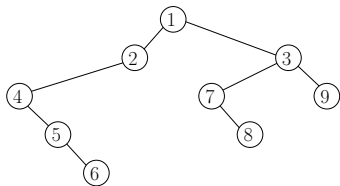
Cartesian tree matching (2)

Similarity

Two sequences x and y are similar if they share the same Cartesian tree, and we note $x \approx_{CT} y$.

Cartesian tree matching (2)

4 5 6 2 1 7 8 3 9 \approx_{CT} 3 4 8 2 1 7 9 5 6



Cartesian tree matching (2)

Cartesian tree matching [Park, Amir, Landau and Park, 2019]

The Cartesian tree matching (CTM) problem is the following:
Given a pattern p and a text t , find every factor f of t such that $f \approx_{CT} p$.

Cartesian tree matching (3)

Parent-distance [PALP19]

Given a sequence $x[1 \dots n]$, the parent-distance representation of x is an integer sequence $\overrightarrow{PD}_x[1 \dots n]$, which is defined as follows:

$$\overrightarrow{PD}_x[i] = \begin{cases} i - \max_{1 \leq j < i} \{j \mid x[j] < x[i]\} & \text{if such } j \text{ exists} \\ 0 & \text{otherwise} \end{cases}$$

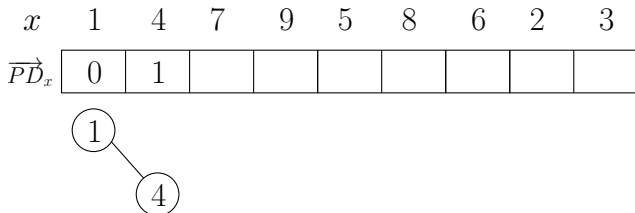
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x									

Cartesian tree matching (3)

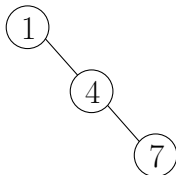
x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0								
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Cartesian tree matching (3)



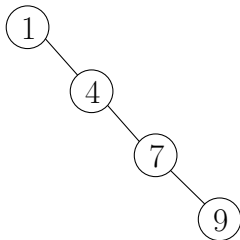
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1						



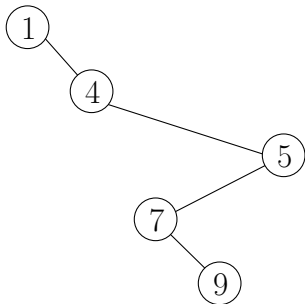
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1					



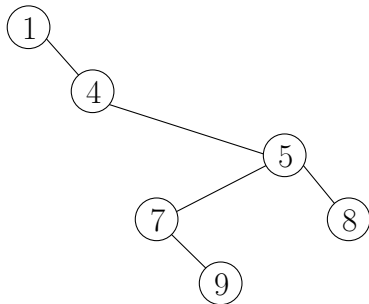
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1	3				



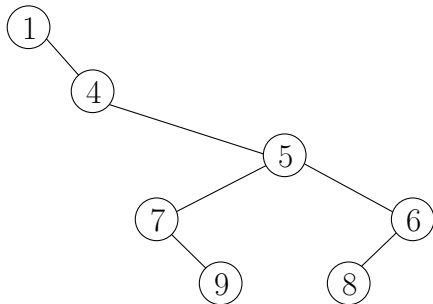
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1	3	1			



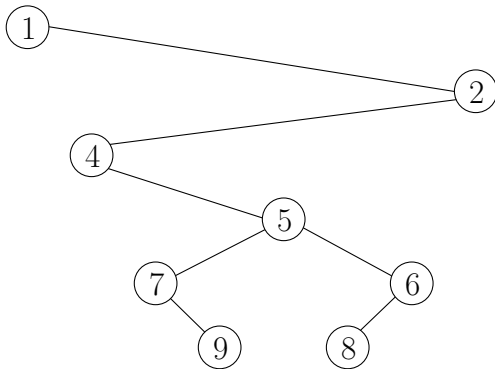
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1	3	1	2		



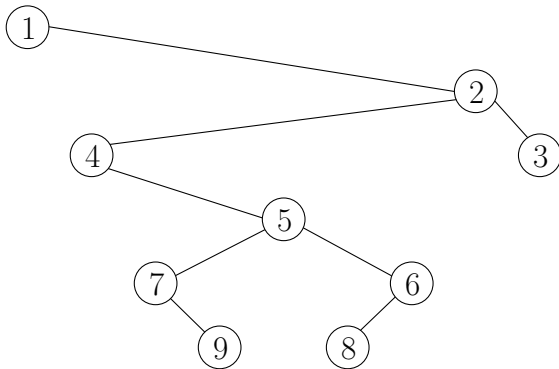
Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1	3	1	2	7	



Cartesian tree matching (3)

x	1	4	7	9	5	8	6	2	3
\overrightarrow{PD}_x	0	1	1	1	3	1	2	7	1



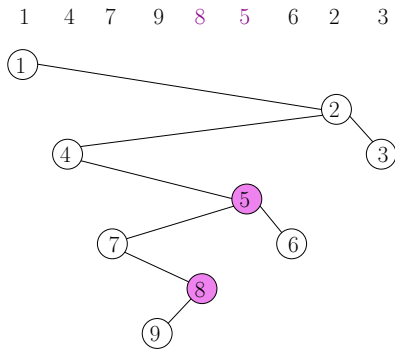
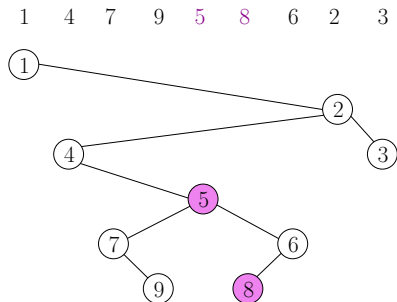
Approximate CTM (1)

Swap

Let x and y be two sequences of length n , and $i \in \{1, \dots, n-1\}$, we denote $y = \tau(x, i)$ to describe a swap, that is:

$$y = \tau(x, i) \text{ if } \begin{cases} x[j] = y[j], \forall j \notin \{i, i+1\} \\ x[i] = y[i+1] \\ x[i+1] = y[i] \end{cases}$$

Approximate CTM (1)



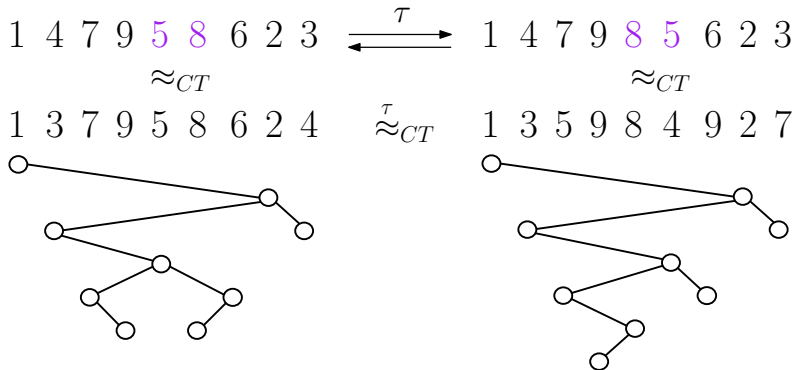
Approximate CTM (2)

Approximate CTM

Let x and y be two sequences of length n , we have $x \stackrel{\tau}{\approx}_{CT} y$ if:

$$\left\{ \begin{array}{l} x \approx_{CT} y \\ \text{or} \\ \exists x', y', \exists i \in \{1, \dots, n-1\}, x' \approx_{CT} x, y' \approx_{CT} y, \\ x' = \tau(y', i) \text{ and } y' = \tau(x', i) \end{array} \right.$$

Approximate CTM (2)



Approximate CTM (3)

Reverse parent-distance

Given a sequence $x[1 \dots n]$, the reverse parent-distance of x is an integer sequence $\overleftarrow{PD}_x[1 \dots n]$, which is defined as follows:

$$\overleftarrow{PD}_x[i] = \begin{cases} \min_{i > j \geq n} \{j \mid x[i] > x[j]\} - i & \text{if such } j \text{ exists} \\ 0 & \text{otherwise} \end{cases}$$

Approximate CTM (3)

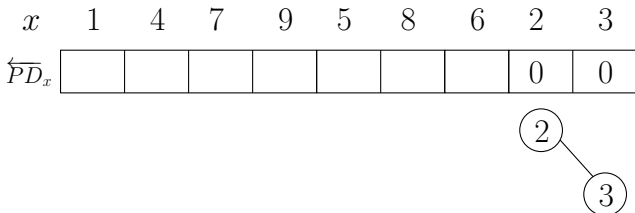
x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>	<input type="text"/>

Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x									0

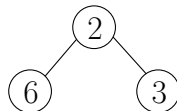
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Approximate CTM (3)



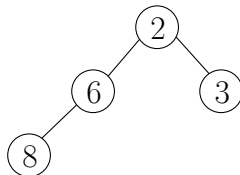
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x							1	0	0



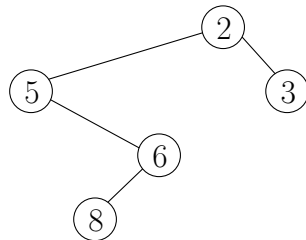
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x						1	1	0	0



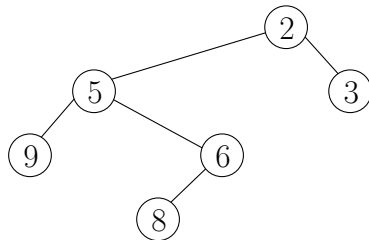
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x					3	1	1	0	0



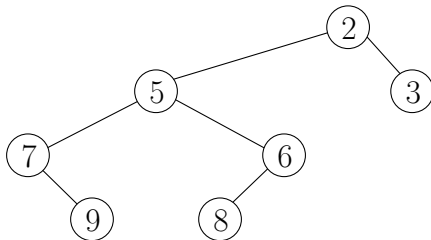
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x				1	3	1	1	0	0



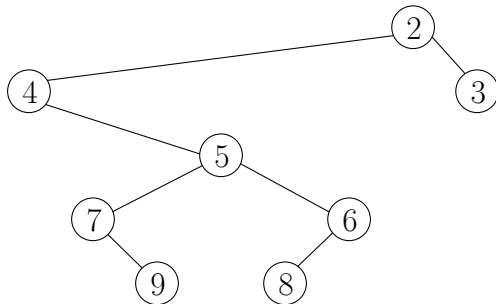
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x			2	1	3	1	1	0	0



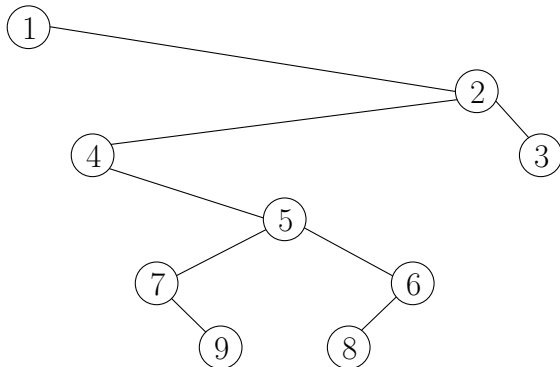
Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x		6	2	1	3	1	1	0	0

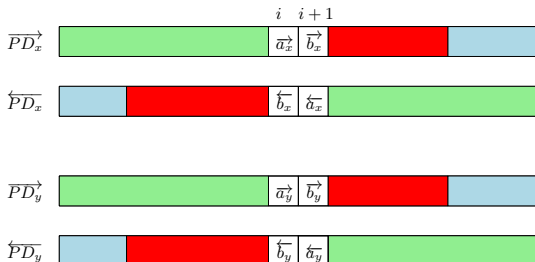


Approximate CTM (3)

x	1	4	7	9	5	8	6	2	3
\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0

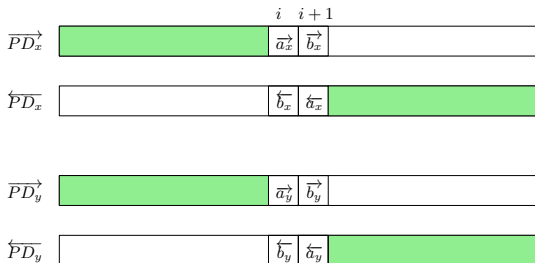


Overview



In the following, let us consider a simple example where $y = \tau(x, i)$ and $x[i] < x[i + 1]$.

Green zones



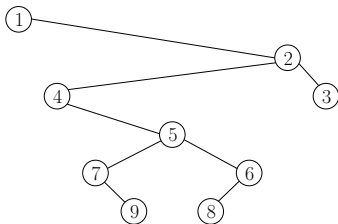
Green zones lemma

The green zones of \overrightarrow{PD}_x and \overrightarrow{PD}_y (resp. \overleftarrow{PD}_x and \overleftarrow{PD}_y) are equal.

Green zones

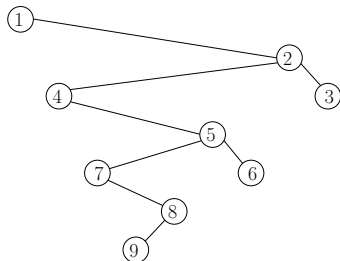
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\vec{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

\vec{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

\overleftarrow{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\vec{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

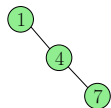
\vec{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

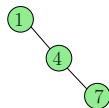
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\vec{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

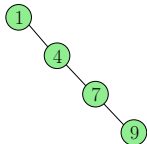
\vec{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

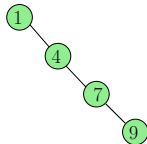
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\vec{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

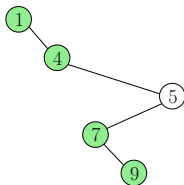
\vec{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

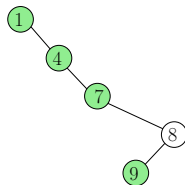
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\vec{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

\vec{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0
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3

y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

\overleftarrow{PD}_y	0	6	3	1	1	2	1	0	0
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3

Green zones

x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

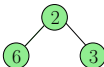
\overleftarrow{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

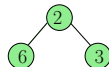
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

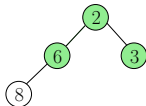
\overleftarrow{PD}_y	0	6	3	1	1	2	1	0	0
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Green zones

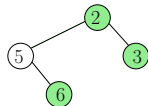
x	1	4	7	9	5	8	6	2	3
\vec{PD}_x	0	1	1	1	3	1	2	7	1

\overleftarrow{PD}_x	0	6	2	1	3	1	1	0	0
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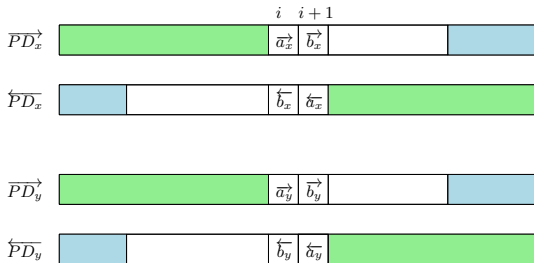


y	1	4	7	9	8	5	6	2	3
\vec{PD}_y	0	1	1	1	2	4	1	7	1

\overleftarrow{PD}_y	0	6	3	1	1	2	1	0	0
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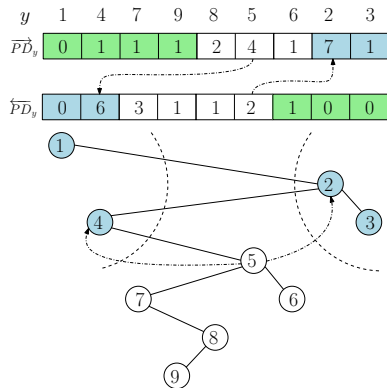
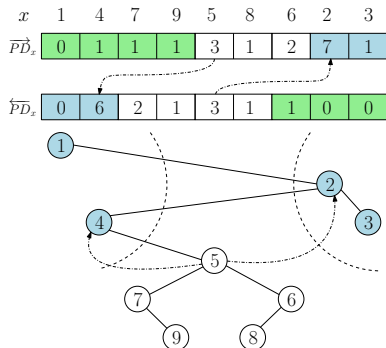
Blue zones



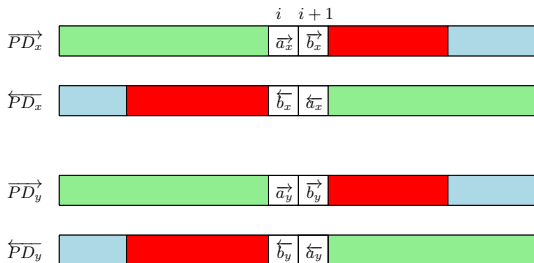
Blue zones lemma

The blue zones of \overrightarrow{PD}_x and \overrightarrow{PD}_y (resp. \overleftarrow{PD}_x and \overleftarrow{PD}_y) are equal.

Blue zones



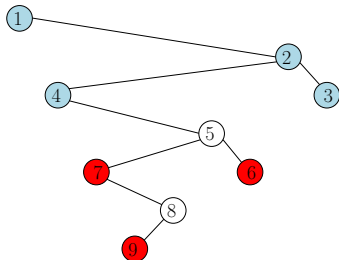
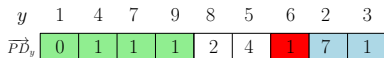
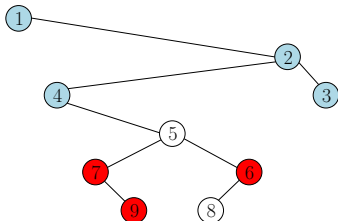
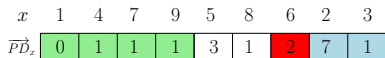
Red zones



Red zones lemma

The red zones of \overrightarrow{PD}_x and \overleftarrow{PD}_x (resp. \overrightarrow{PD}_y and \overleftarrow{PD}_y) differ by at most one.

Red zones



Local

x	1	4	7	9	5	8	6	2	3
$\vec{P}\vec{D}_x$	0	1	1	1	3	1	2	7	1

$\vec{P}\vec{D}_x$	0	6	2	1	3	1	1	0	0
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y	1	4	7	9	8	5	6	2	3
$\vec{P}\vec{D}_y$	0	1	1	1	2	4	1	7	1

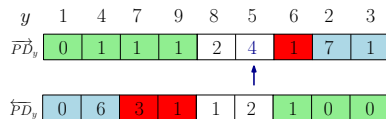
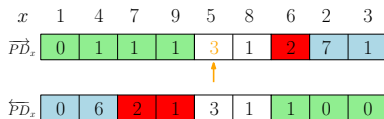
$\vec{P}\vec{D}_y$	0	6	3	1	1	2	1	0	0
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Local lemma

$$\textcircled{1} \overleftarrow{b}_y = 1$$

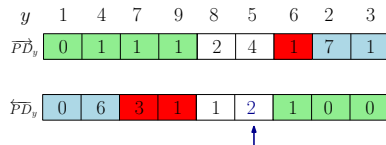
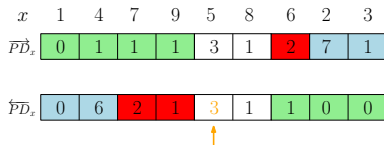
Local



Local lemma

$$2 \quad \vec{b}_y = \begin{cases} 0 & \text{if } \vec{a}_x = 0 \\ \vec{a}_x + 1 & \text{otherwise} \end{cases}$$

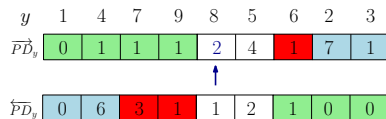
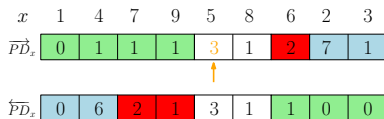
Local



Local lemma

$$\textcircled{3} \quad \overleftarrow{a}_y = \begin{cases} 0 & \text{if } \overleftarrow{b}_x = 0 \\ \overleftarrow{b}_x - 1 & \text{otherwise} \end{cases}$$

Local



Local lemma

$$\textcircled{4} \quad \vec{a}_y \leq \begin{cases} i - 1 & \text{if } \vec{a}_x = 0 \\ \vec{a}_x & \text{otherwise} \end{cases}$$

A parent-distance based algorithm

Algorithm 1: *DoubleParentDistanceMethod*(p, t)

Input : A pattern p and a text t

Output: The occurrences that $\overset{\tau}{\approx}_{CT} p$ in t

```

1  $(\overrightarrow{PD}_p, \overleftarrow{PD}_p) \leftarrow$  Compute the parent-distance tables of  $p$ ;
2 for  $j \in \{1, \dots, |t| - |p| + 1\}$  do
3    $(\overrightarrow{PD}_x, \overleftarrow{PD}_x) \leftarrow$  Compute the parent-distance tables of
    $x = t[j \dots j + p - 1]$ ;
4   if  $\overrightarrow{PD}_p = \overrightarrow{PD}_x$  then
5     | An occurrence has been found;
6   else
7     | foreach Eligible position for a swap do
8       | | if Lemmas Blue, Red and Local hold then
9       | | | An occurrence has been found;
```

A parent-distance based algorithm

Complexity

The parent-distance based algorithm has a worst-case time complexity of $\Theta(mn)$ and a $\Theta(m)$ space complexity (where m is the length of the pattern and n the length of the text).

Swap graph

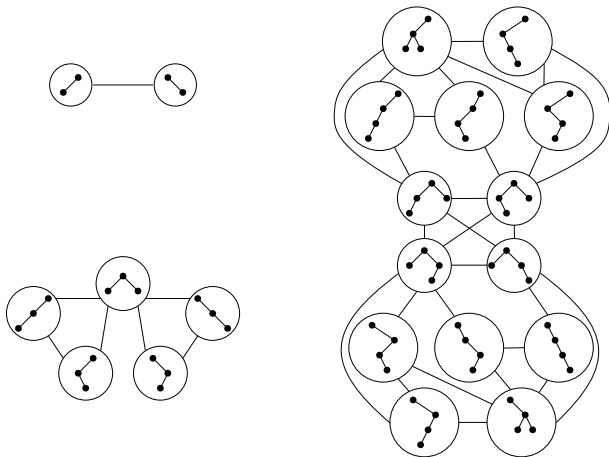
Definition

The swap graph of Cartesian trees for a given n is a graph where:

- The vertices are the Cartesian trees of size n
- There is an edge between two vertices T and T' if there exist 2 sequences x and y such that:

$$C(x) = T, C(y) = T' \text{ and } x \stackrel{\tau}{\approx}_{CT} y$$

Swap graph



Lower bound

Number of Cartesian trees

The number of Cartesian tree T with n nodes is the n -th Catalan number:

$$C_n = \frac{1}{n+1} \binom{2n}{n} = O\left(\frac{4^n}{n^{3/2}}\right)$$

Neighbours lemma

The number of neighbours $|ng(T)|$ of a given Cartesian tree T is bounded, and we have:

$$n - 1 \leq |ng(T)| \leq \lceil 3(n - 1) - 2(\log_2(n + 1) - 1) \rceil$$

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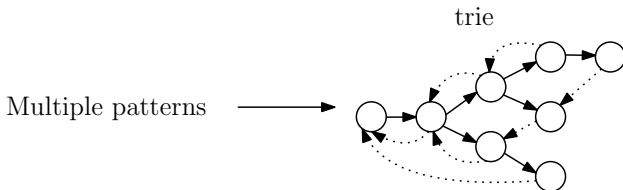
$$n - 1 \leq |ng(T)| \leq \lceil 3(n - 1) - 2(\log_2(n + 1) - 1) \rceil$$

A lower bound for the graph diameter

The diameter of the swap graph is $\Omega\left(\frac{n}{\ln n}\right)$.

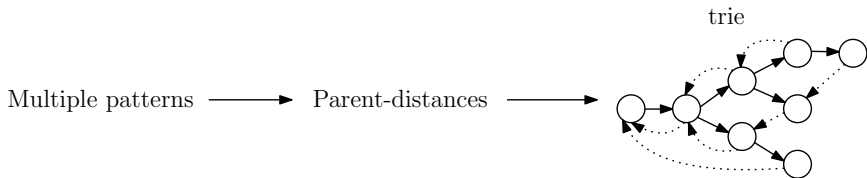
An Aho-Corasick based algorithm

[Alfred V. Aho, Margaret J. Corasick 1975]

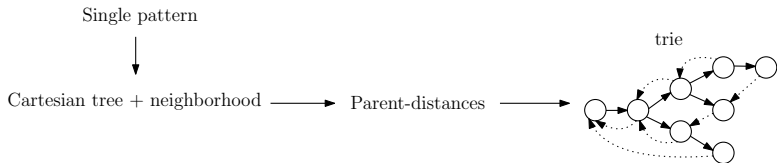


An Aho-Corasick based algorithm

[S. G. Park, A. Amir, G. M. Landau, K. Park 2019]



An Aho-Corasick based algorithm



An Aho-Corasick based algorithm

Complexity

The Aho-Corasick based algorithm has an $O((m^2 + n) \log m)$ worst-case time complexity and an $O(m^2)$ space complexity (where m is the length of the pattern and n the length of the text).

Closing words

Perspectives

- Generalize our results
- Use another representation of CT
- Introduce new metrics for approximate CTM
- Filtration
- Average complexity

Thank you for your attention!