

Approximate Cartesian Tree Matching with Substitutions

Panagiotis Charalampopoulos, *Jonas Ellert*, Manal Mohamed

Cartesian tree matching with substitutions

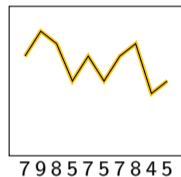
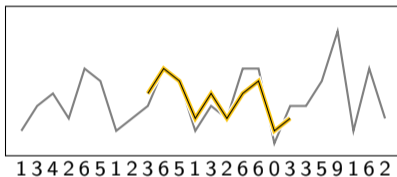
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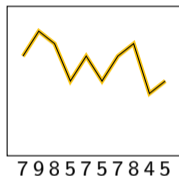
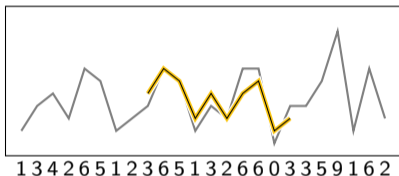


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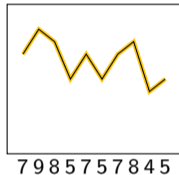
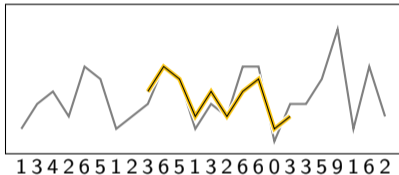


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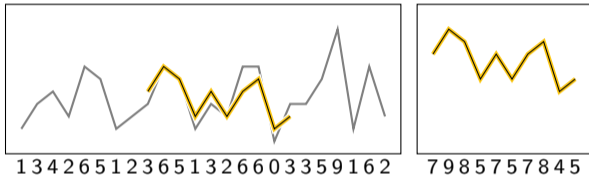
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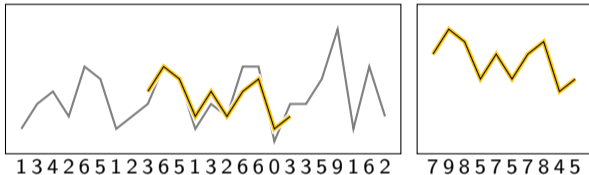
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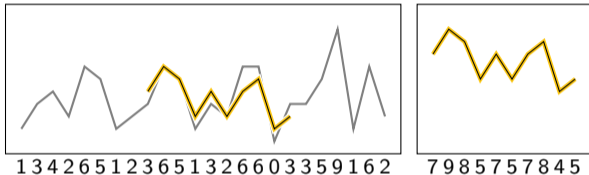
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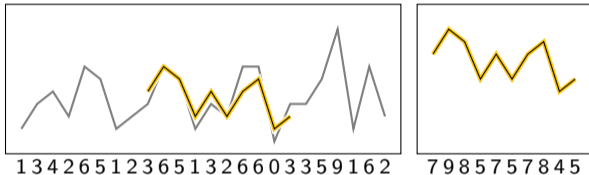
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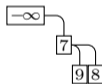
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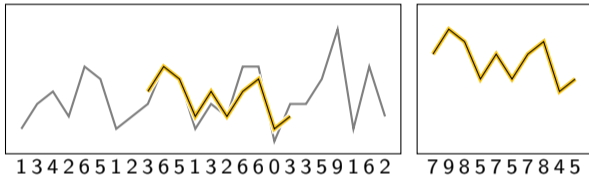
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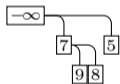
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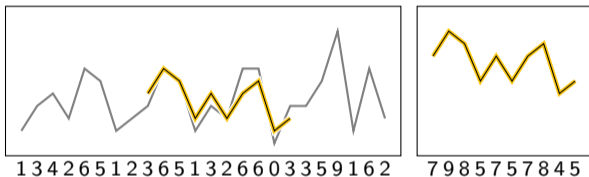
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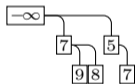
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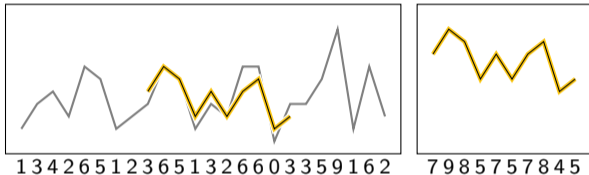
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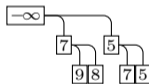
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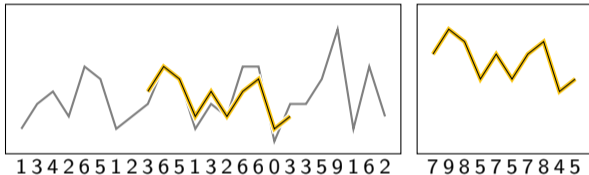
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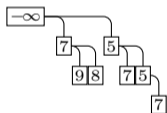
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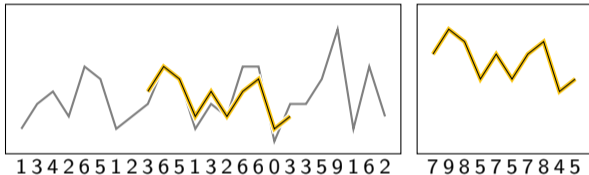
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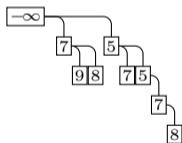
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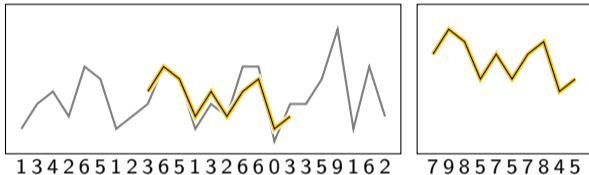


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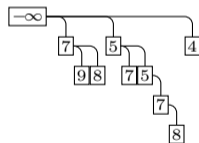
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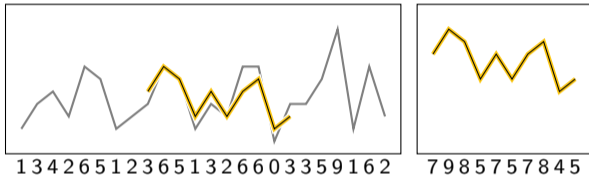


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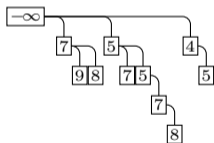
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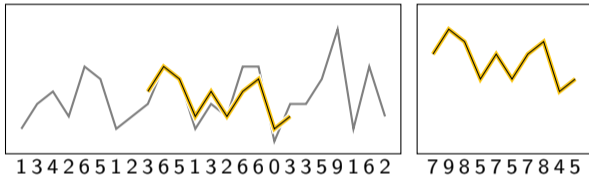


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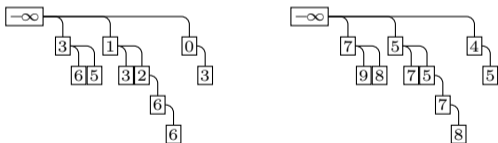
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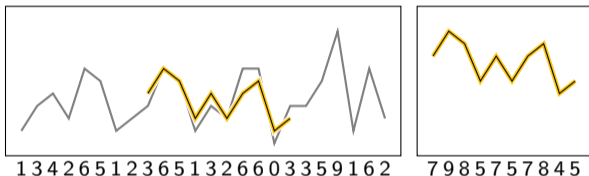
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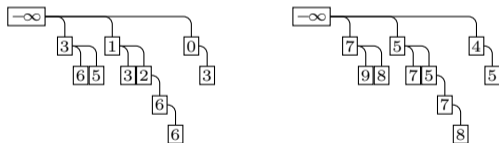
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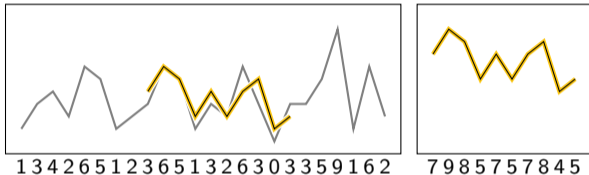
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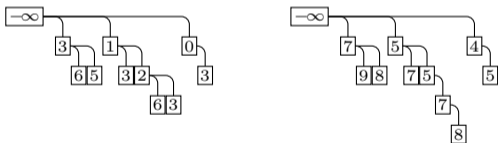
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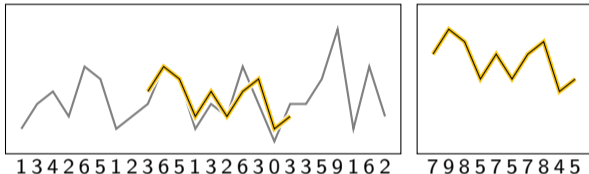
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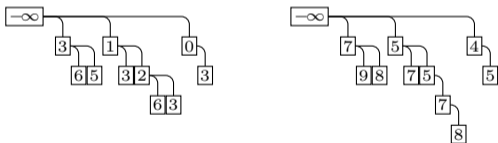
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solution: allow k substitutions in text
before computing tree

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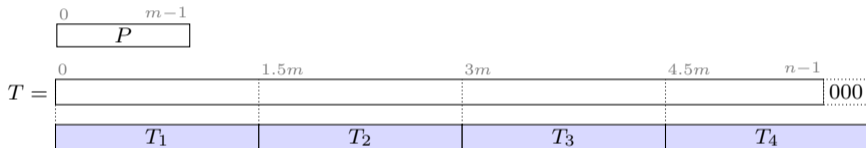
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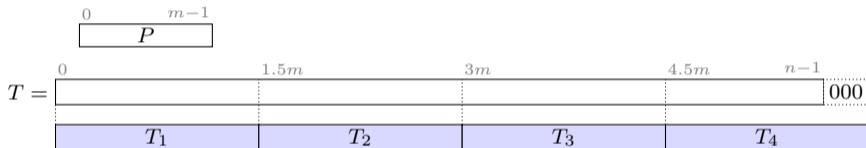


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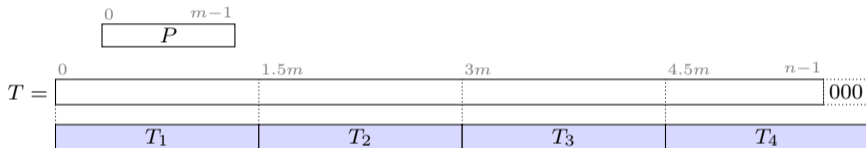


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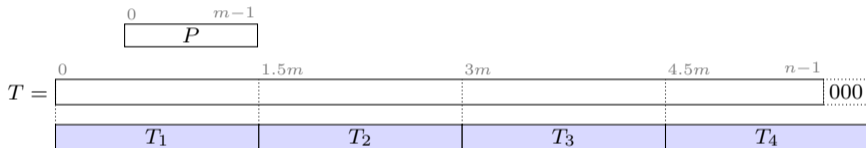


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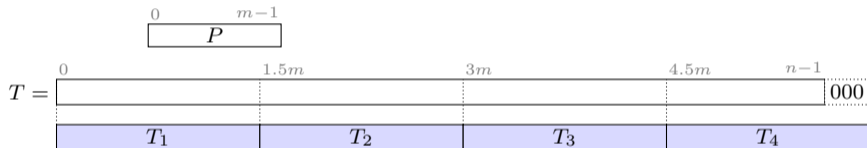


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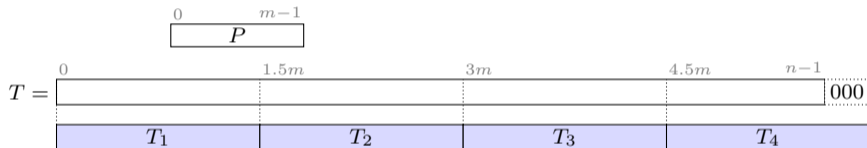


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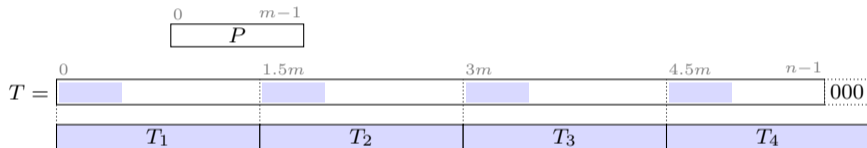


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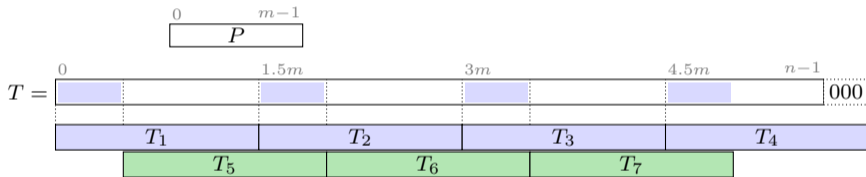


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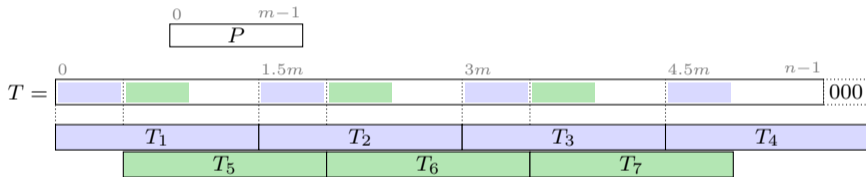


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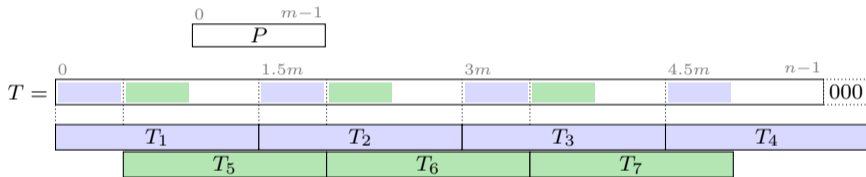


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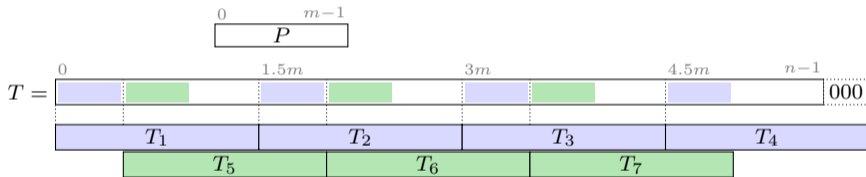


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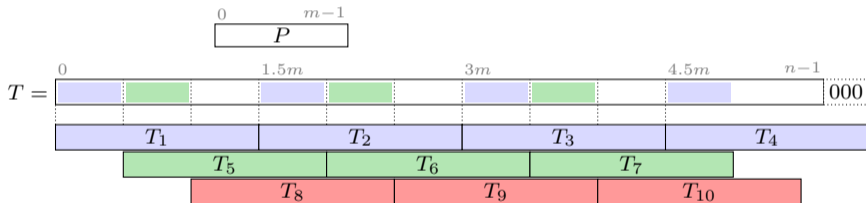


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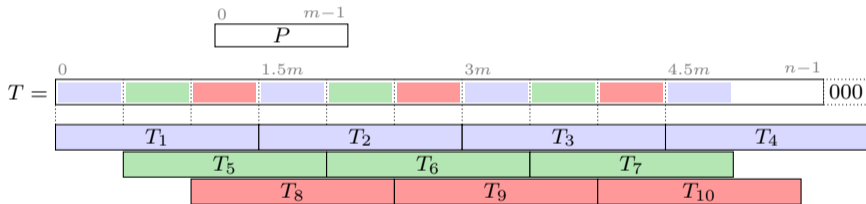


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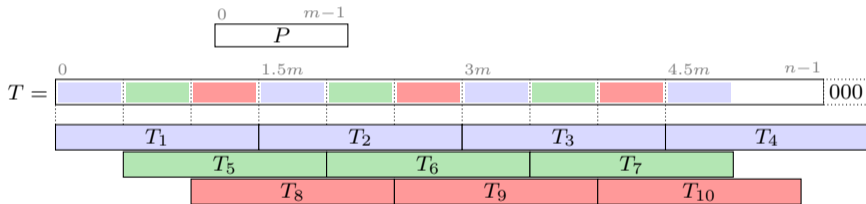


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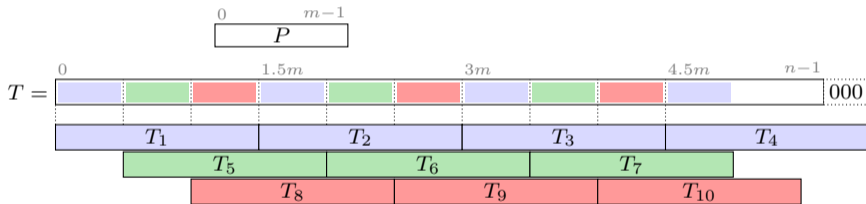


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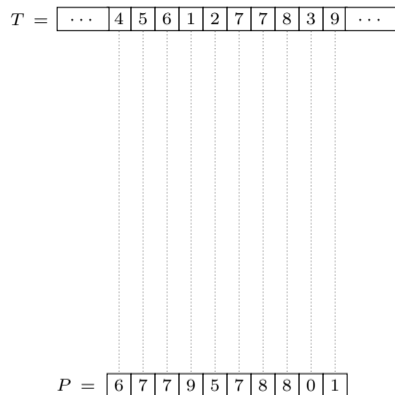
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- $\mathcal{O}(n/m)$ blocks, as promised!

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- text $T[1..1.5m]$, pattern $P[1..m]$, threshold k
- output all fragments $T[i..i+m)$ at Hamming distance $\leq k$ to a string that has same CT as P

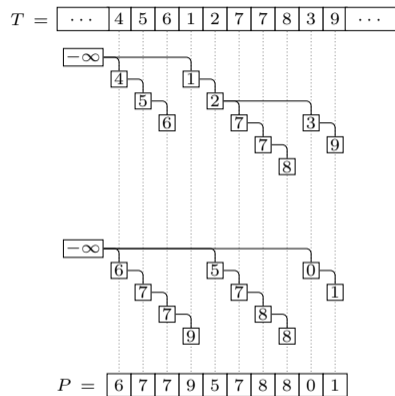
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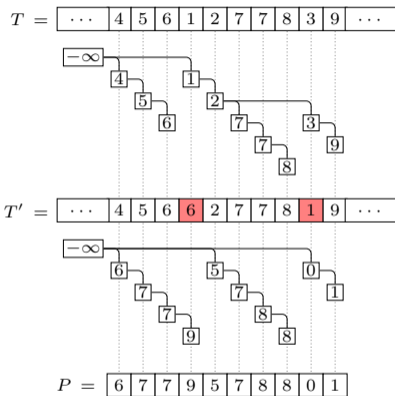
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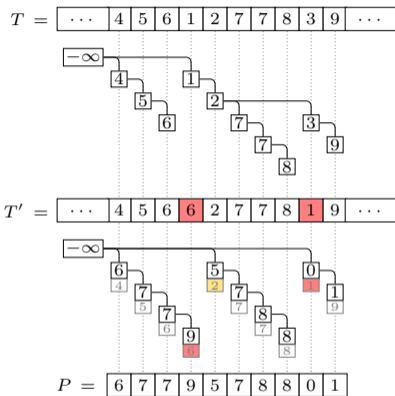
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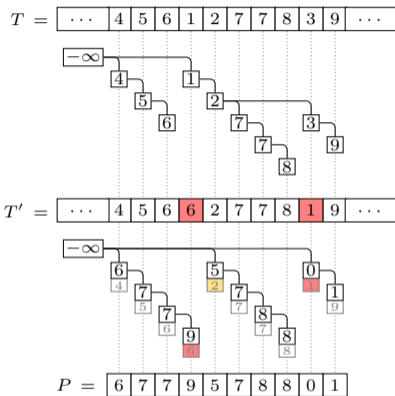


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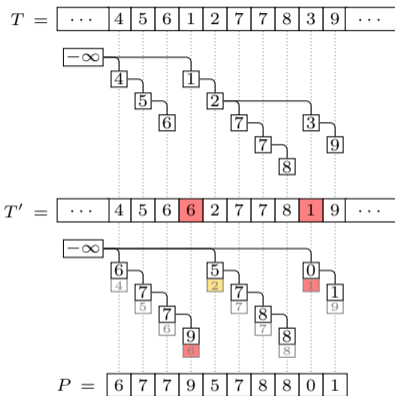
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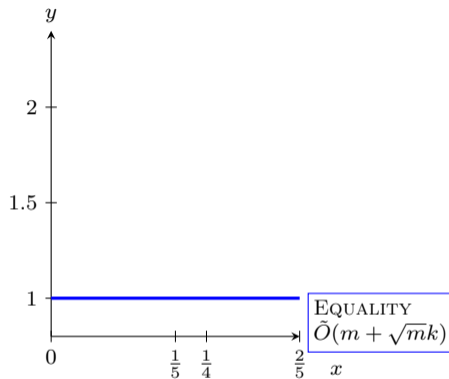
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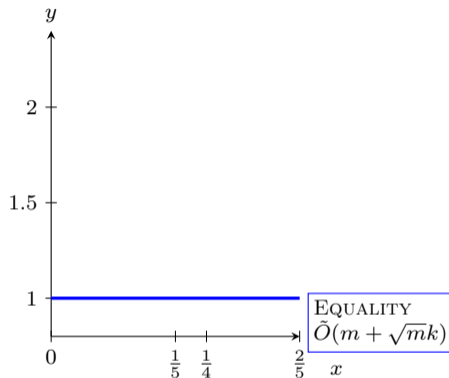
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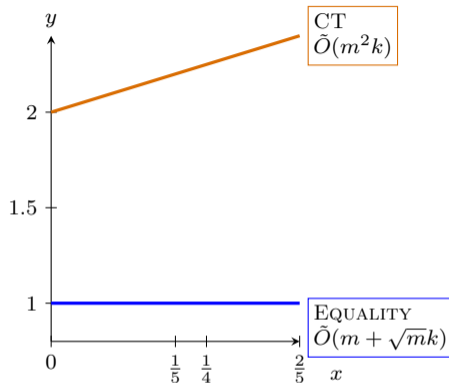
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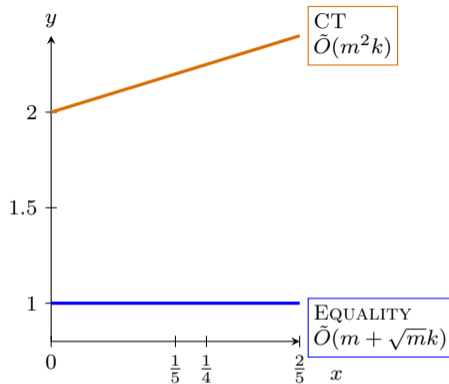
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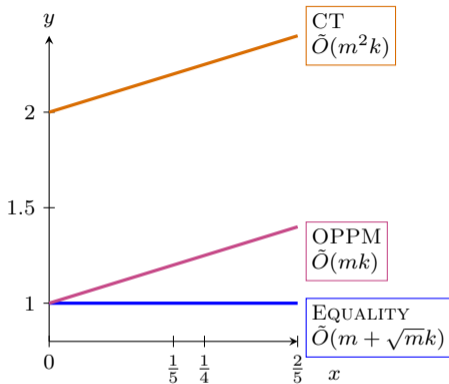
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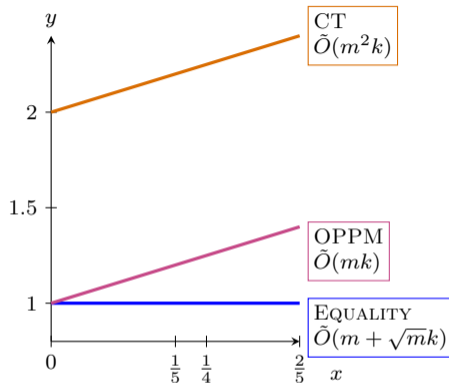
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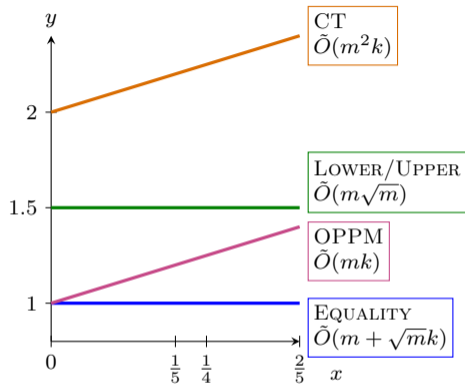
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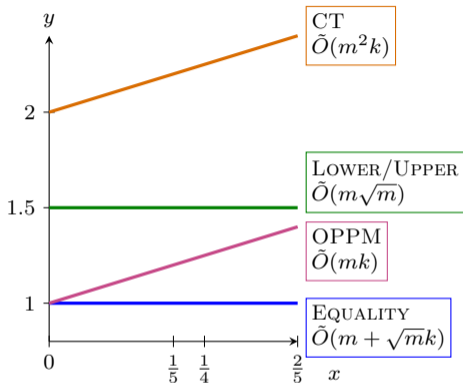
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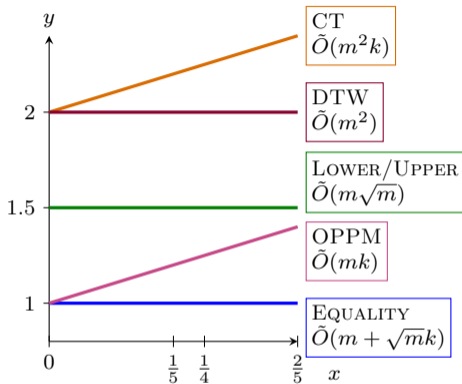
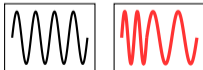
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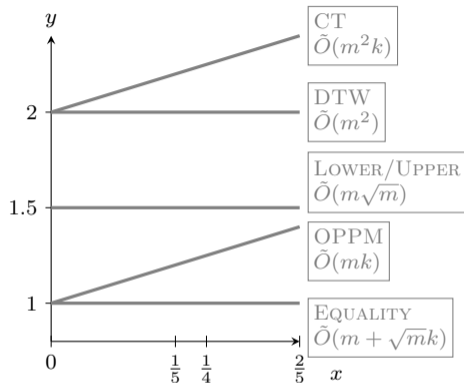
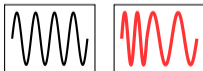
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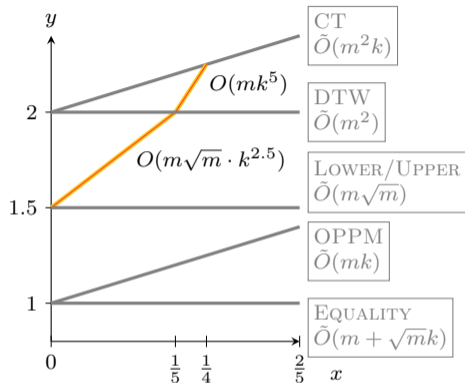
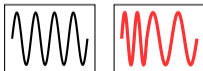
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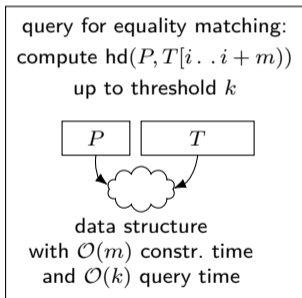
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query for equality matching:
compute $\text{hd}(P, T[i..i+m])$
up to threshold k

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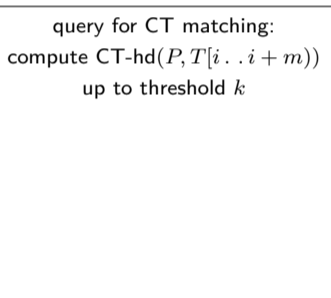
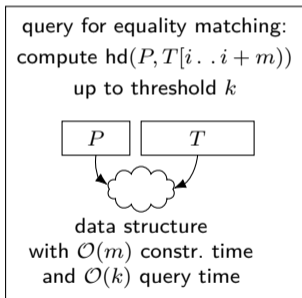
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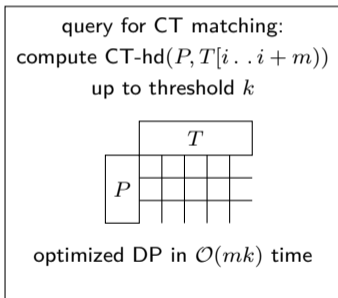
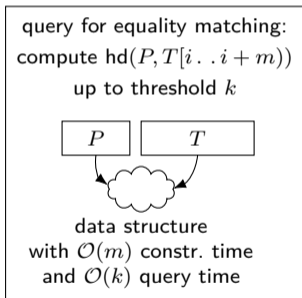
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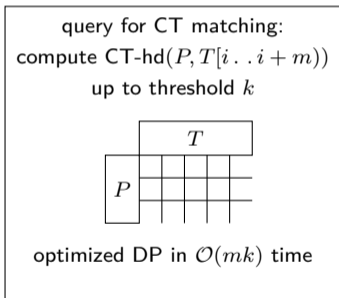
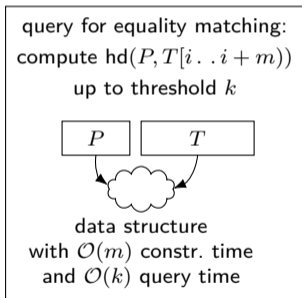
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- algorithms for equality matching heavily exploit periodicity

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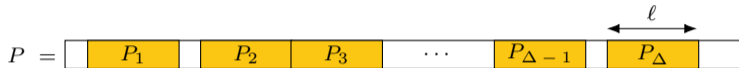
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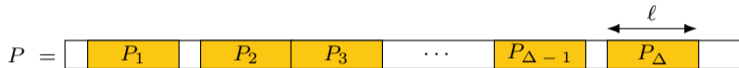
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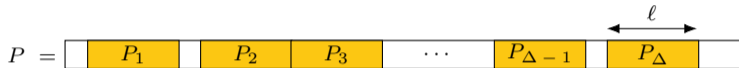
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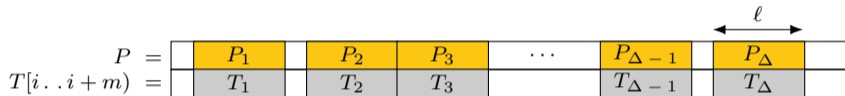
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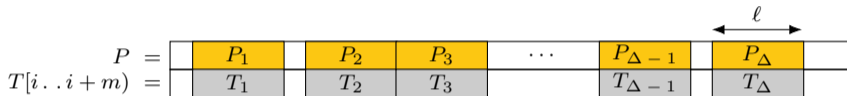
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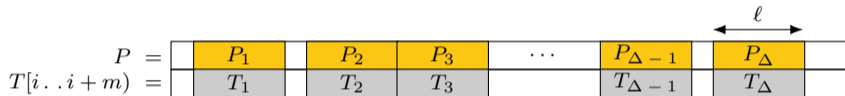


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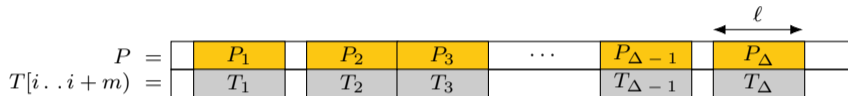


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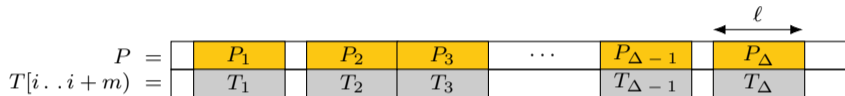


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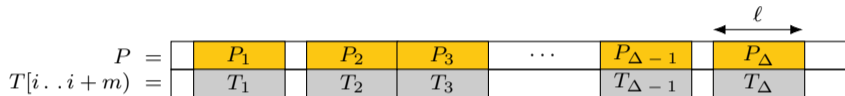


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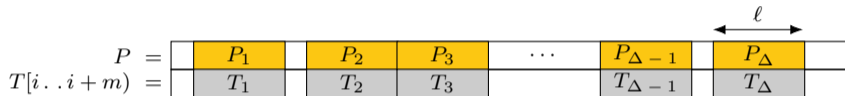


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- if $T[i..i+m]$ and P have same CT after k substitutions in T then P_i and T_i have same CT for at least $\Delta - k \geq \Delta/2$ choices of i
- use exact CT matching to find all occurrences of all P_i in T in $\tilde{O}(m\Delta)$ time
- at most $\mathcal{O}(m)$ occurrences in total, at most $\mathcal{O}(m/\Delta)$ text fragments have enough matches
- verify each in $\mathcal{O}(mk)$ time
- overall time $\tilde{O}(m^2k/\Delta + m\Delta) = \tilde{O}(m\sqrt{m} \cdot \text{poly}(k))$ using $\Delta = \Theta(\sqrt{m}/\text{poly}(k))$

What if we can't find enough anchors?

- fix some parameter $\Delta \in [2k, \sqrt{m}/2]$ to define some anchor substrings of P
 - Δ non-overlapping fragments of P
 - each of length $\ell \approx m/(2\Delta)$
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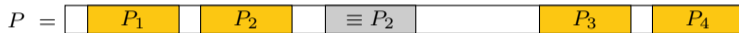
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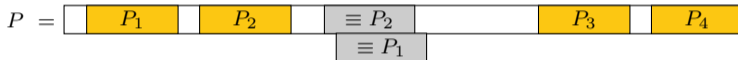
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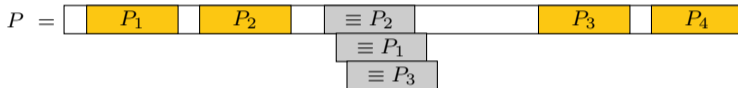
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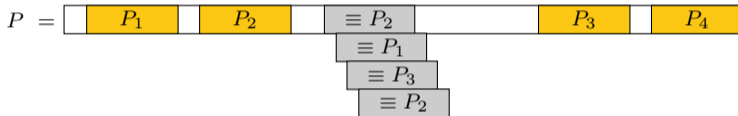
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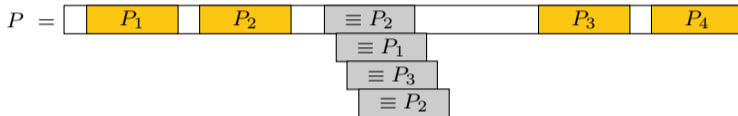
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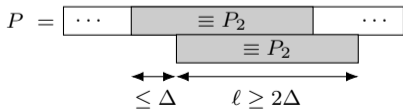
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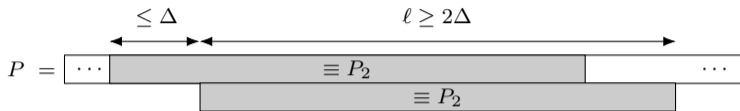


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- zoom in...



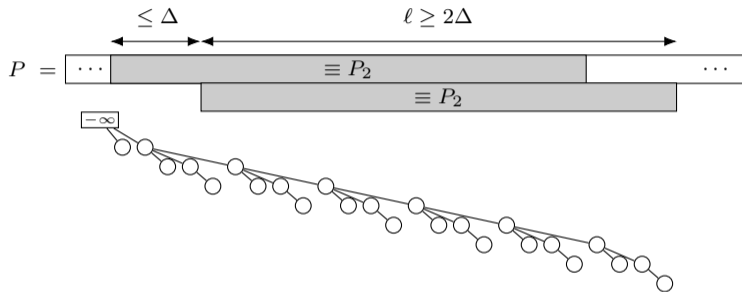
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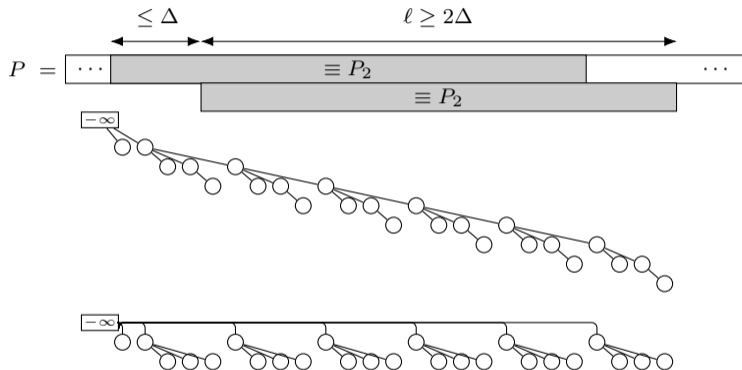
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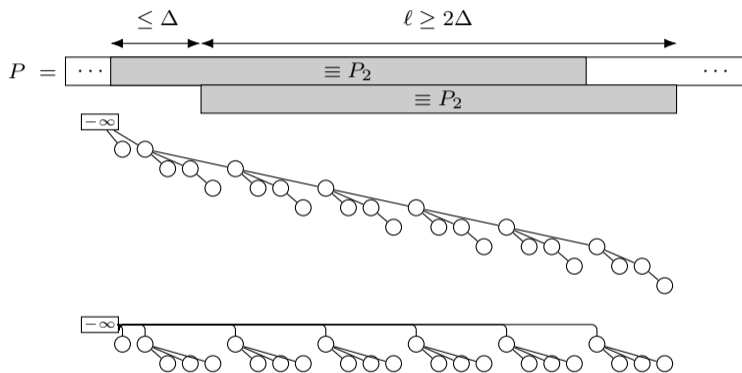
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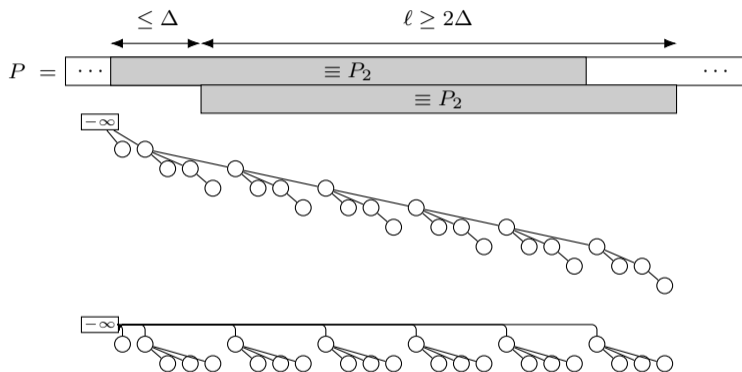
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- highly periodic with respect to structure of CT

What if we can't find enough anchors?

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- highly periodic with respect to structure of CT
- adapt all the tricks for periodic fragments from equality pattern matching

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Open questions:

- What is the correct complexity?
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- CT/OPPM/DTW... how do they actually work on real data?

References

-  Boneh, Itai et al. (2024). “Optimal Dynamic Time Warping on Run-Length Encoded Strings”. In: *51st International Colloquium on Automata, Languages, and Programming, ICALP 2024, Tallinn, Estonia, July 8-12, 2024*. Vol. 297. LIPIcs. Schloss Dagstuhl - Leibniz-Zentrum für Informatik, 30:1–30:17. DOI: 10.4230/LIPICs.ICALP.2024.30.
-  Charalampopoulos, Panagiotis, Jonas Ellert, and Manal Mohamed (2026). “Approximate Cartesian Tree Matching with Substitutions”. In: *43rd International Symposium on Theoretical Aspects of Computer Science, STACS 2026, Grenoble, France, March 9-13, 2026*. Vol. 364. LIPIcs. Schloss Dagstuhl - Leibniz-Zentrum für Informatik, 26:1–26:21. DOI: 10.4230/LIPICs.STACS.2026.26.
-  Gawrychowski, Pawel and Przemyslaw Uznanski (2016). “Order-preserving pattern matching with k mismatches”. In: *Theor. Comput. Sci.* 638, pp. 136–144. DOI: 10.1016/J.TCS.2015.08.022.
-  — (2018). “Towards Unified Approximate Pattern Matching for Hamming and L_1 Distance”. In: *45th International Colloquium on Automata, Languages, and Programming, ICALP 2018, Prague, Czech Republic, July 9-13, 2018*. Vol. 107. LIPIcs. Schloss Dagstuhl - Leibniz-Zentrum für Informatik, 62:1–62:13. DOI: 10.4230/LIPICs.ICALP.2018.62.
-  Kim, Sungmin and Yo-Sub Han (2025). “Approximate Cartesian tree pattern matching”. In: *Theor. Comput. Sci.* 1056, Article 115506. DOI: 10.1016/J.TCS.2025.115506.
-  Labib, Karim, Przemyslaw Uznanski, and Daniel Wolleb-Graf (2019). “Hamming Distance Completeness”. In: *30th Annual Symposium on Combinatorial Pattern Matching, CPM 2019, Pisa, Italy, June 18-20, 2019*. Vol. 128. LIPIcs. Schloss Dagstuhl - Leibniz-Zentrum für Informatik, 14:1–14:17. DOI: 10.4230/LIPICs.CPM.2019.14.