From Strings to Seaweeds Modern Tools for Classical Problems

Philip Wellnitz National Institute of Informatics Tokyo

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(National Institute of Informatics). (Universitã© Paris Citã©, CNRS, IRIF, F-75013, Paris, France); (Universitat PolitÃ" cnica de Catalunya) •

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An Example

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•••• An Example

Task: Find Saarbrücken in a text.

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•••• An Example

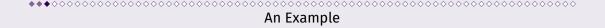
Task: Find Saarbrücken in a text.

Or Saarbruecken.

•••• An Example

Task: Find Saarbrücken in a text.

Or Saarbruecken. Or Sarrebruck.



Task: Find Saarbrücken in a text.

Or Saarbruecken. Or Sarrebruck. Or Saarbrucken, Saarbr|cken, Saarbrücken,

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The Approximate String Matching Problem

Approximate String Matching

Given a text *T*, a pattern string *P*, and an integer *k*, identify the (starting positions of) substrings of *T* that are at **edit distance** of at most *k* to *P*.



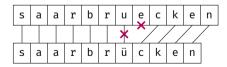
early 1980's

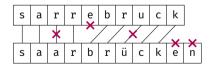
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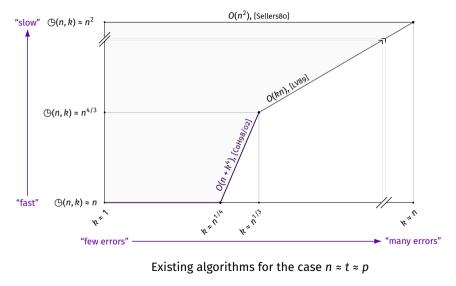
Edit distance: minimum number of insertions, deletions, or substitutions of single characters to transform one string into another string

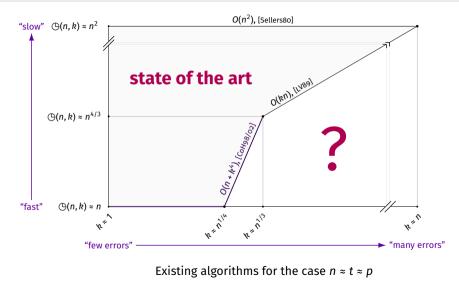


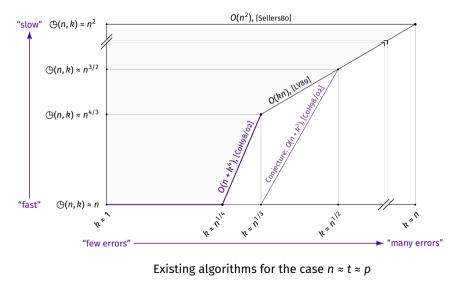


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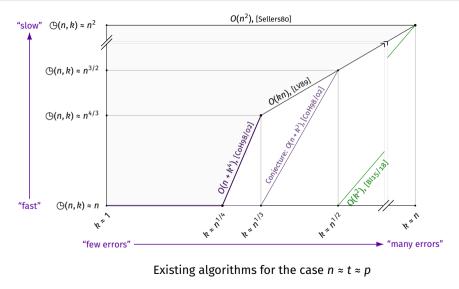
••••• State-of-the-Art Algorithms



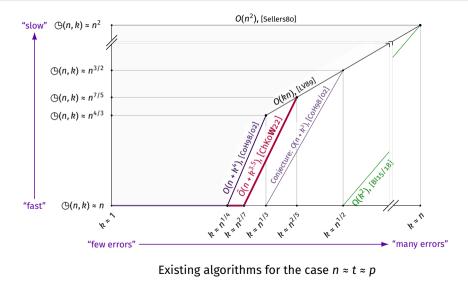




•••••• State-of-the-Art Algorithms



••••• State-of-the-Art Algorithms



••••••••Outline

Intro

Some Background and Basic Definitions (Some) Classical Algorithms for Approximate String Matching

Structural Insights into the Solution Structure The Marking Trick and How to Handle Easy Patterns Dynamic Puzzle Matching and How to Handle Not So Easy Patterns

On Puzzles and Seaweeds

Some Technical Definitions

Detour: On the Name of Seaweeds

Alignment Graphs and Permutation Matrices of Strings

Solving Dynamic Puzzle Matching via the Seaweed Method

Outlook: Pattern Matching with Weighted Edits and Open Problems

Approximate String Matching

For a text T, a pattern P, and an integer k,

Identify the (starting positions of) substrings of T that are at edit distance of at most k to P.

• Naive idea: Compute edit distance to pattern starting from any position in the text

→ Need to compute edit distance between strings

Approximate String Matching

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◆ Naive idea: Compute edit distance to pattern starting from any position in the text
 → Need to compute edit distance between strings

Edit Distance (ED)

For two strings T and P, compute their edit distance

(minimum number of insertions, deletions, and substitutions to transform T into P).

 Textbook dynamic programming algorithm from around 1970, runs in time O(tp) (writing t = |T|, p = |P|)

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From Edit Distance to Approximate String Matching

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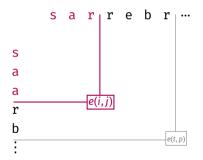
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- Textbook dynamic programming algorithm from around 1970, runs in time O(tp) (writing t = |T|, p = |P|)
- ♦ Write e(i, j) for the ED of T[0..i) and P[0..j)
 → e(t, p) is the ED of T and P.
- Clearly, e(i, 0) = i (delete T[0..i)) and e(0, j) = j (insert P[0..j))
- Observe

e(i,j) = min

$$\begin{cases}
e(i - 1,j) + 1 & (delete from T) \\
e(i,j - 1) + 1 & (insert in T) \\
e(i - 1,j - 1) \\
+(T[i] \neq P[j]) & (match/subst)
\end{cases}$$



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- Observe

$$f(i,j) = \min \begin{cases} e(i-1,j) + 1 & (\text{delete from } T) \\ e(i,j-1) + 1 & (\text{insert in } T) \\ e(i-1,j-1) \\ +(T[i] \neq P[j]) & (\text{match/subst}) \end{cases}$$

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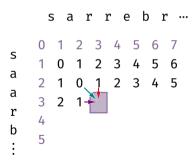
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	5	5 6	a 1	r 1	c e	e k	נכ		••
c	0	1	2	3	4	5	6	7	
S	1	0	1	2	3	4	5	6	
a	2	1	0	1	2	3	4	5	
a	3	2	1	1					
r	4								
b	5								
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	9	5 6	a 1	C 1	6	e k) 1		
-	0	1	2	3	4	5	6	7	
s	1	0	1	2	3	4	5	6	
a a	2	1	0	1	2	3	4	5	
a r	3	2	1	1	2	3	4	5	
b	4	3	2	1	1	2	3	4	
:	5	4	3	2	2	2	2	3	

Approximate String Matching

For a text T, a pattern P, and an integer k,

Identify the (starting positions of) substrings of T that are at edit distance of at most k to P.

- Textbook dynamic programming algorithm from ~1970 1980, runs in time O(tp) (writing t = |T|, p = |P|), [Sellers 1980, rediscovered several times]
- Write e(i, j) for the ED of T[0..i) and P[0..j)

after deleting an arbitrary prefix of T

- Clearly, e(i, 0) = 0 (delete T[0..i)) and e(0, j) = j (insert P[0..j))
- Observe

$$e(i,j) = \min \begin{cases} e(i-1,j)+1 & (\text{delete from } T) \\ e(i,j-1)+1 & (\text{insert in } T) \\ e(i-1,j-1) \\ +(T[i] \neq P[j]) & (\text{match/subst}) \end{cases}$$

							-	-	
	0	0	0	0	0	0	0	0	
S	1	0	1	1	1	1	1	1	
a	2	1	0	1	2	2	2	2	
а	3	2	1	1	2	3	3	3	
r	4	3	2	1	1	2	3	3	
b	5	4	3	2	2	2	2	3	
:									

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	s	a	r	r	е	b	r	
_	0	0	0	0	0	0	0	0
s	1	0	1	1	1	1	1	1
a a	2	1	0	1	2	2	2	2
a n	3	2	1	1	2	3	3	3
r b	4	3	2	1	1	2	3	3
D	5	4	3	2	2	2	2	3
nding	pos	sitio	ns o	f sa	arb	wit	h k	= 2.

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- Textbook dynamic programming algorithm from ~1970 1980, runs in time O(tp) (writing t = |T|, p = |P|), [Sellers 1980, rediscovered several times]
- Write e(i,j) for the ED of $T^{R}[0..i)$ and $P^{R}[0..j)$

after deleting an arbitrary prefix of T^R

- Clearly, e(i, 0) = 0 (delete T[0..i)) and e(0, j) = j (insert P[0..j))
- Observe

$$e(i,j) = \min \begin{cases} e(i-1,j) + 1 & (\text{delete from } T) \\ e(i,j-1) + 1 & (\text{insert in } T) \\ e(i-1,j-1) \\ +(T[i] \neq P[j]) & (\text{match/subst}) \end{cases}$$

	1	r ł) (9 1	r 1	r a	a :	s
b	0	0	0	0	0	0	0	0
	1	1	0	1	1	1	1	1
r	2	1	1	1	1	1	2	2
a a	3	2	2	2	2	2	1	2
a s	4	3	3	3	3	3	2	2
5	5	4	4	4	4	4	3	2
artir	og n	ociti	ione	of		ah u	ith.	h = 2

Starting positions of saarb with k = 2.

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•••••••••• Detour: More on (Historic) Applications

Approximate String Matching

For a text *T*, a pattern *P*, and an integer *k*,

Identify the (starting positions of) substrings of T that are at edit distance of at most k to P.

(Variations of) Sellers' O(tp) Algorithm were (are?) popular for

- (Have seen) Text Retrieval: Find occurrences of patterns or phrases in a text, accounting for misspellings or conversion errors
- Signal Processing: Find patterns in signals, accounting for transmission errors
- Computational Biology: Find specific patterns in DNA sequences, accounting for mutations or evolutionary alterations

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There must be faster algorithms out there!

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There must be faster algorithms out there!



Approximate String Matching

For a text T, a pattern P, and an integer k,

Identify the (starting positions of) substrings of T that are at **edit distance** of at most k to P.

So far, did not use k

→ Do not need values *e*(*i*,*j*) > *k*

How do we compute just values ≤ k?
 → Compute values diagonally.

	1	C 1	: a	a s	5 1	c a	a a	a :	5 1	r ł	n a	a s	5 1	C 1	c a	a s	5
	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
r	1	0	0	1	1	0	1	1	1	0	1	1	1	0	0	1	1
a	2	1	1	0	1	1	0	1	2	1	1	1	2	1	1	0	1
a s	3	2	2	1	1	2	1	0	1	2	2	1	2	2	2	1	1
s r	4	3	3	2	1	2	2	1	0	1	2	2	1	2	3	2	1
-	5	4	3	3	2	1	2	2	1	0	1	2	2	1	2	3	2
a a	6	5	4	3	3	2	1	2	2	1	1	1	2	2	2	2	3
a s	7	6	5	4	4	3	2	1	2	2	2	1	2	3	3	2	3
S	8	7	6	5	4	4	3	2	1	2	3	2	1	2	3	3	2

Finding saarsaar with $k \le 2$ edits.

Approximate String Matching

For a text T, a pattern P, and an integer k,

Identify the (starting positions of) substrings of T that are at **edit distance** of at most k to P.

◆ So far, did not use k → Do not need values e(i, j) > k

♦ How do we compute just values ≤ k?
 → Compute values diagonally.

	r	r	а	S	1	r a	a	a	S	r	h	a	S	r	ri	a s	5
0	0	()	0	0	0	0	0	0	0	0	0	0	0	0	0	0
1	0	()	1	1	0	1	1	1	0	1	1	1	0	0	1	1
2	1	1	I	0	1	1	0	1	2	1	1	1	2	1	1	0	1
3	2	2	2	1	1	2	1	0	1	2	2	1	2	2	2	1	1
4	3	1	3	2	1	2	2	1	0	1	2	2	1	2	3	2	1
5	4	1.1	3	3	2	1	2	2	1	0	1	2	2	1	2	3	2
6	5	L	÷	3	3	2	1	2	2	1	1	1	2	2	2	2	3
7	6		5	4	4	3	2	1	2	2	2	1	2	3	3	2	3
8	7	6	5	5	4	4	3	2	1	2	3	2	1	2	3	3	2

Finding saarsaar with $k \le 2$ edits.

Approximate String Matching

For a text T, a pattern P, and an integer k,

Identify the (starting positions of) substrings of T that are at **edit distance** of at most k to P.

- ◆ So far, did not use k
 → Do not need values e(i, j) > k
- How do we compute just values ≤ k?
 → Compute values diagonally.

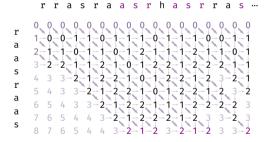
1	. 1	r a	a s	5 1	r a	a a	a	S 1	r I	h i	a :	5 1	r 1	c a	a s	5
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
1	0	0	1	1	0	1	1	1	0	1	1	1	0	0	1	1
2	1	1	0	1	1	0	1	2	1	1	1	2	1	1	0	1
3	2	2	1	1	2	1	0	1	2	2	1	2	2	2	1	1
4	3	3	2	1	2	2	1	0	1	2	2	1	2	3	2	1
5	4	3	3	2	1	2	2	1	0	1	2	2	1	2	3	2
6	5	4	3	3	2	1	2	2	1	1	1	2	2	2	2	3
7	6	5	4	4	3	2	1	2	2	2	1	2	3	3	2	3
8	7	6	5	4	4	3	2	1	2	3	2	1	2	3	3	2

Finding saarsaar with $k \le 2$ edits.

Approximate String Matching

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Finding saarsaar with $k \le 2$ edits.

Approximate String Matching

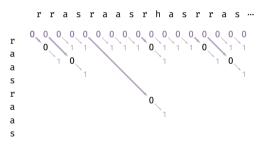
For a text T, a pattern P, and an integer k,

- ◆ So far, did not use k
 → Do not need values e(i, j) > k
- How do we compute just values ≤ k?

 → Compute furthest position on diag d reachable with ℓ = 0, ..., k edits.

 → Jump over equal substrings in O(1) time (Kangaroo Jumps).

 → In total: O((t + p)k) time time [Landau, Vishkin'89].



Finding saarsaar with $k \leq 0$ edits.

Approximate String Matching

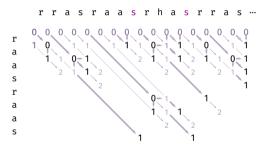
For a text T, a pattern P, and an integer k,

- ◆ So far, did not use k
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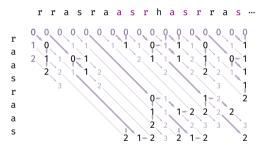


Finding saarsaar with $k \le 1$ edits.

Approximate String Matching

For a text T, a pattern P, and an integer k,

- ◆ So far, did not use k
 → Do not need values e(i, j) > k
- How do we compute just values ≤ k?
 → Compute furthest position on diag d reachable with l = 0, ..., k edits.
 → Jump over equal substrings in O(1) time (Kangaroo Jumps).
 → In total: O((t + p)k) time time [Landau Vishkin'80]



Finding saarsaar with $k \le 2$ edits.

Approximate String Matching

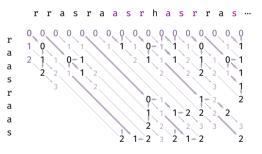
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Finding saarsaar with $k \le 2$ edits.

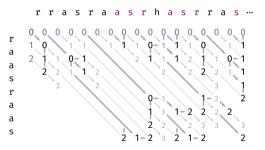
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Finding saarsaar with $k \le 2$ edits.

There must be faster algorithms out there!

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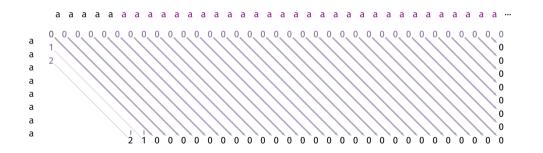
...



Philip Wellnitz 12-2

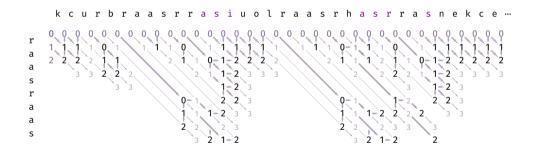
Idea: Can we filter out diagonals that will never lead to an occurrence?

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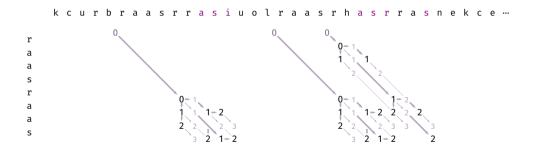


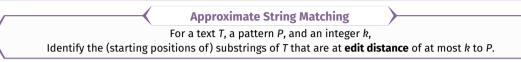
Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 12-4

Idea: Can we filter out diagonals that will never lead to an occurrence? Sometimes we can; and if we cannot, there might be structure to exploit!



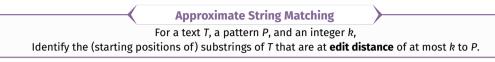
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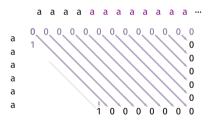
Idea: Find parts of P that are rare in T → Can filter out candidates for occurrences

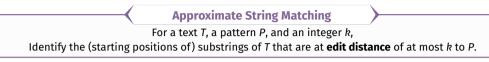
- Seen: Highly repetitive parts R are bad...
 ~→ R may appear ≈ t times in T
- Non-repetitive parts B are good!
 → B may appear only ≈ t/b times in T
 → Problem: B may appear approximately in T



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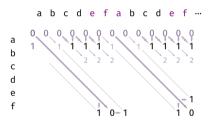


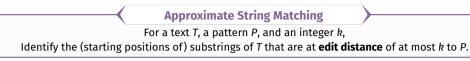


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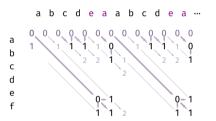
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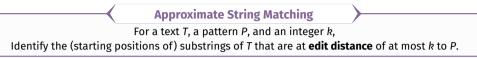




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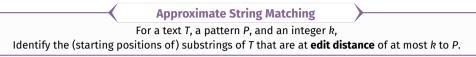
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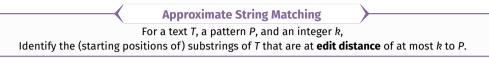
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- Leads to O(t + k⁴ · t/p) algorithm (more details soon) [Cole,Hariharan'02] (announced '98)





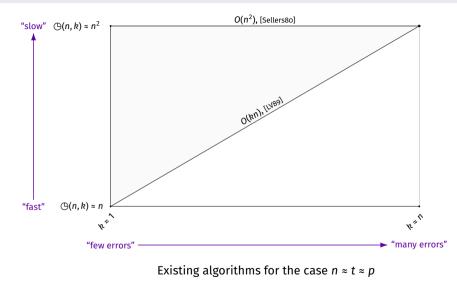
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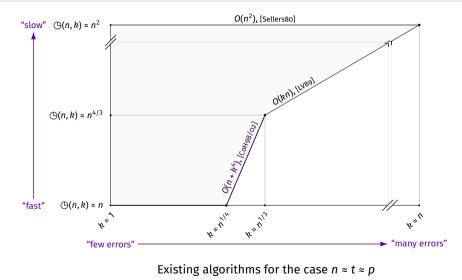


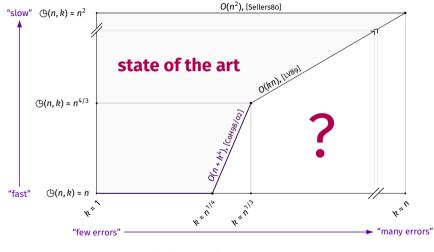


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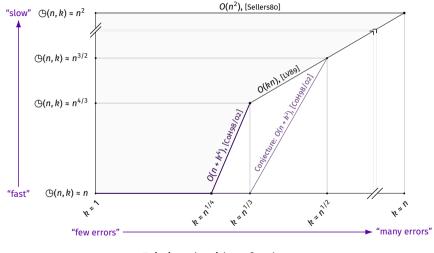




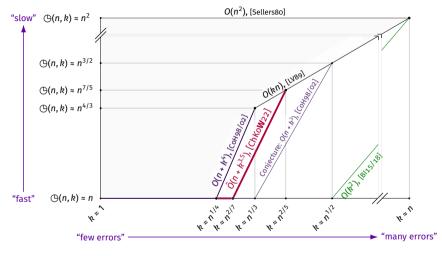




Existing algorithms for the case $n \approx t \approx p$



Existing algorithms for the case $n \approx t \approx p$



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Why did a new improvement take 24 years?

• Simpler problems were not fully understood!

• Example 1: The complexity of Edit Distance was settled only in 2014

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 \rightsquigarrow Yields lower bound of $O(t + k^2 \cdot t/p)$ for Approximate String Matching.

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- Example 2: String Matching with Mismatches (no insertions or deletions of chars)
 - → O(tk)_algorithm [Landau, Vishkin'**86**]
 - $\rightarrow \tilde{O}(t\sqrt{k})$ algorithm [Amir, Lewenstein, Porat'**04**]
 - $\rightarrow \tilde{O}(t + t/p \cdot k^2)$ algorithm [CFPSS'**16**]

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Optimal String Matching with Mismatches

There is a $\tilde{O}(t + kt/\sqrt{p})$ -time algorithm for String Matching with Mismatches and no significantly faster algorithm exits.

(Unless there is a major breakthrough for combinatorial Boolean Matrix Multiplication.)

[GU18] (ann. 2017)

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[GU18] (ann. 2017)

Beating Cole and Hariharan's Algorithm

How do we obtain faster algorithms?

New insights into the solution structure of Approximate String Matching Beating Cole and Hariharan's Algorithm

How do we obtain faster algorithms?

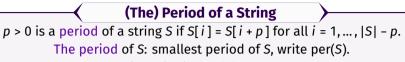
New insights into the solution structure of Approximate String Matching



Structural Results for Approximate String Matching

Step o:

What is the solution structure of Exact String Matching?



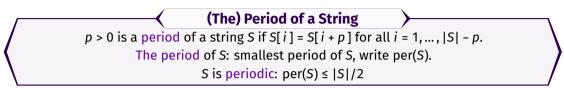
S is periodic: $per(S) \le |S|/2$

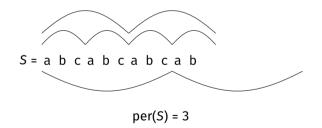


S = a b c a b c a b c a b

per(S) = 3

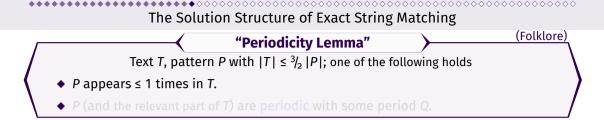
Philip Wellnitz 21-1

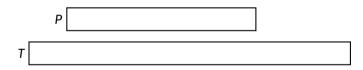


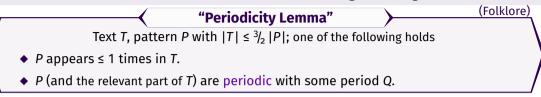


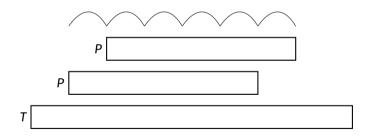
6 and 9 also periods of S

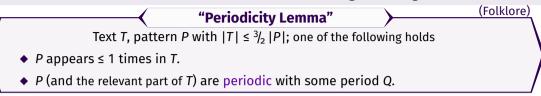
Philip Wellnitz 21-2

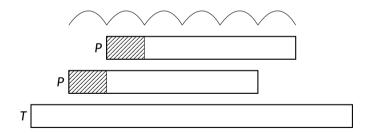


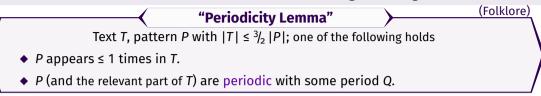


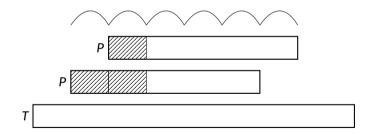


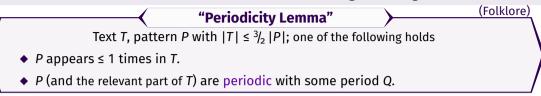


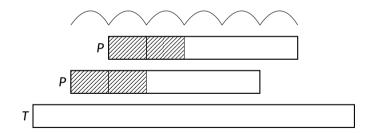


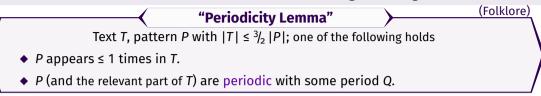


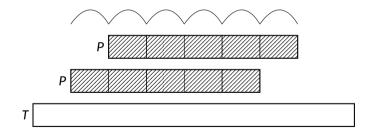


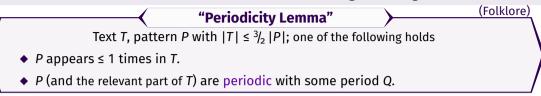


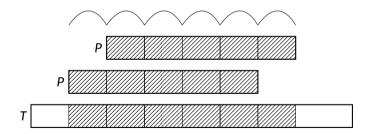


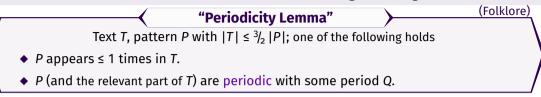


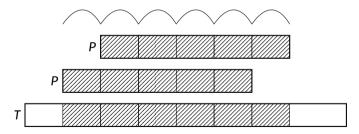










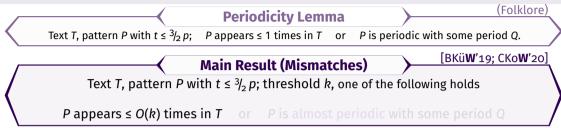


The Standard Trick: For $|T| \gg |P|$ consider separately O(n/m) fragments of T of length $\leq \frac{3}{2}m$ that overlap by m - 1 positions.

Step 1:

What is the solution structure of String Matching with Mismatches?

Structural Results for Approximate String Matching

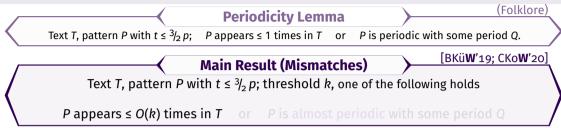


Ta ···· a c a a a ··· a c a a ··· a c a

a…ac…c appears 2k times

aa…aca has approximate period a (is at HD ≤ 2k from Q[∞])

Structural Results for Approximate String Matching

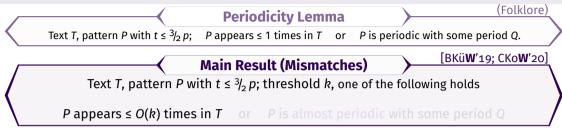


a…ac…c appears 2k times

ramacaamac paamaca

aa…aca has approximate period a (is at HD ≤ 2*k* from Q[∞])

Structural Results for Approximate String Matching



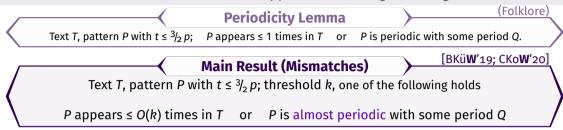
P a … a c … c

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Structural Results for Approximate String Matching



P a … a c … c

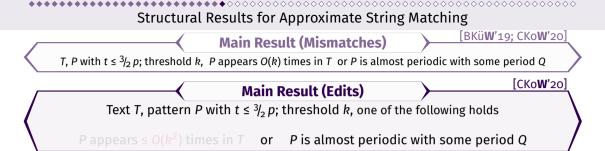
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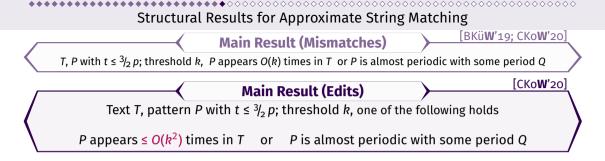
T a ··· a c a a a ··· a c a a ··· a c a a ··· a c a

aa…aca has approximate period a (is at HD $\leq 2k$ from Q^{∞})

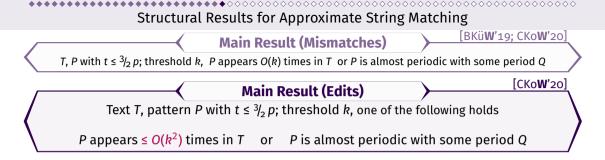
Step 1.5:

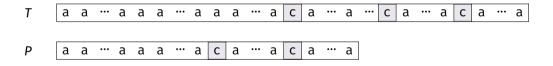
The solution structure of Approximate String Matching.



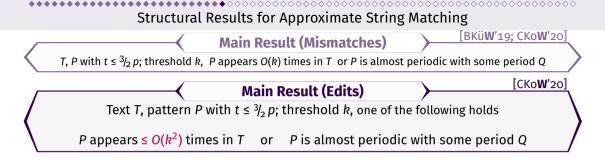


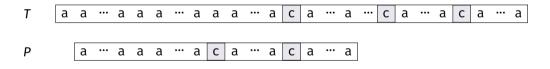
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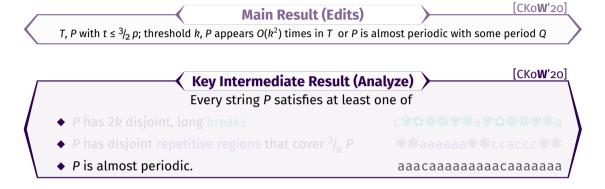


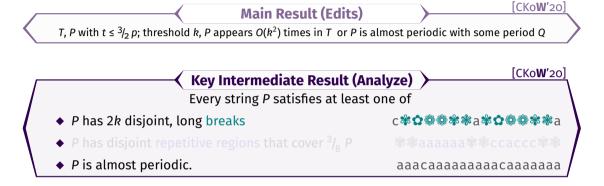
aa···aaa···aca···a appears $\approx k^2$ times

Main Result (Edits)

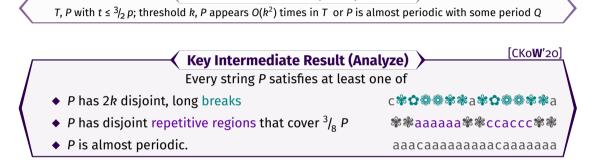
T, P with $t \leq 3/2 p$; threshold k, P appears $O(k^2)$ times in T or P is almost periodic with some period Q

[CKo**W**'20]

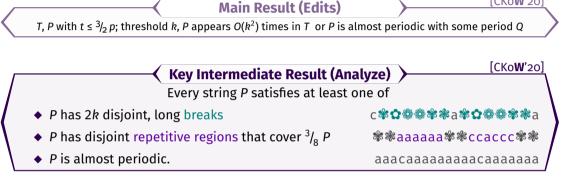




Main Result (Edits)

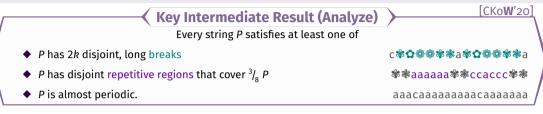


[CKo**W**'20

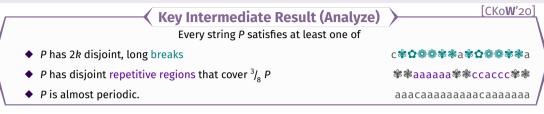


Analyze implies Main Result.

[CKo**W**'20

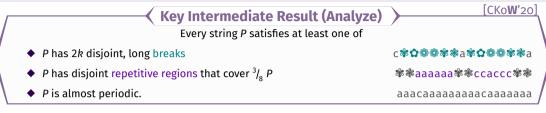






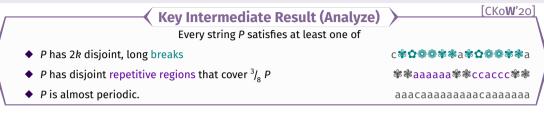


Process P from left to right, p/8k new characters at a time.



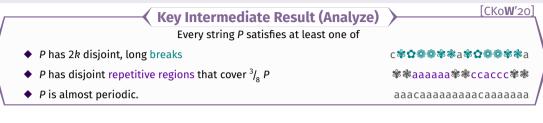


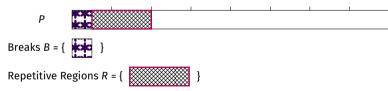
• If a fragment is a break, add it to the found breaks.



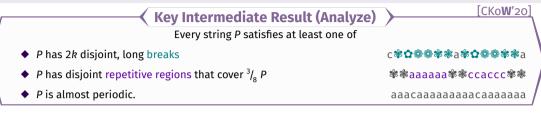


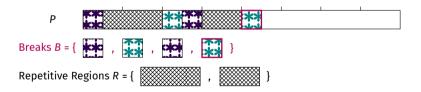
• Otherwise, find the shortest prefix (longer than *p*/8*k*) that is a repetitive region.



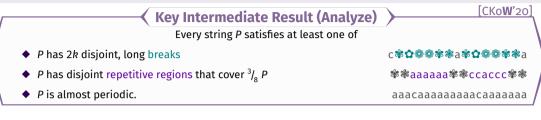


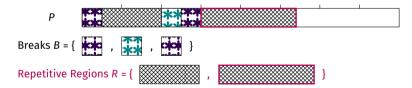
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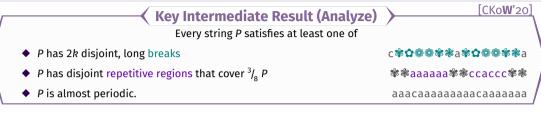


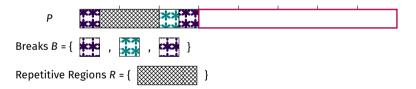
• If we found 2k breaks, return the breaks.



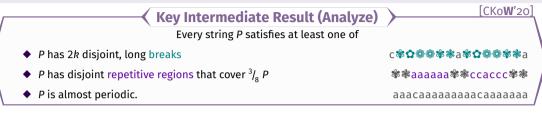


• If the total length of the repetitive regions is > $3/8 \cdot p$, return the repetitive regions.





• If we reach the end of *P*, try to find a single repetitive region starting from the end.

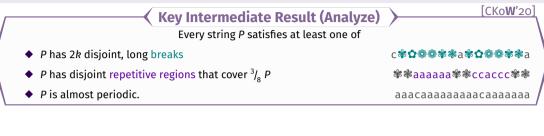


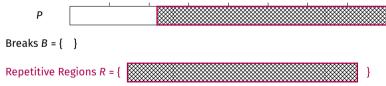


Breaks B = { }

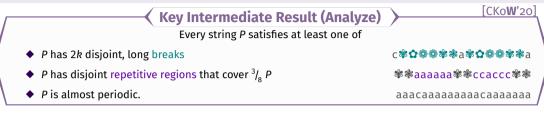
Repetitive Regions R = { }

• If we reach the end of P, try to find a single repetitive region starting from the end.





• If we found a repetitive region, return it.

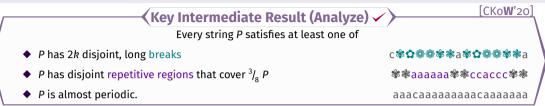




Repetitive Regions R = { }

• If we again don't obtain a repetitive region, *P* is almost periodic.

Structural Results for Approximate String Matching



How do we turn our insights into faster algorithms?

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Need to tackle three cases.

- P contains 2k disjoint breaks;
- P contains disjoint repetitive regions R_i;
- *P* is almost periodic

How do we turn our insights into faster algorithms?

Need to tackle three two cases.

- *P* contains 2*k* disjoint breaks;
- ◆ P contains disjoint repetitive regions R;; →→ Follows from the other two cases.
- *P* is almost periodic

Have: 2k disjoint breaks B_1, \dots, B_{2k} in P such that

• $|B_i| = \Theta(|P|/k)$ and • $|B_i|$ is not periodic

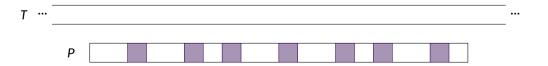
• In any *k*-edit occ, at least 1 break is matched exactly.



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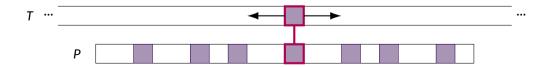
• Algorithm idea: For each B_i, find exact matches of B_i in T



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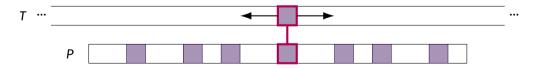
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- Have at most |T|/(|B_i|/2) = Θ(k|T|/|P|) exact occ's of B_i
 → O(k²|T|/|P|) calls to [LV'89]; O(k⁴|T|/|P|) time in total

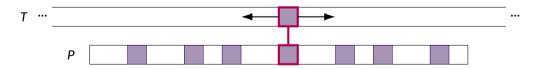


Philip Wellnitz 31-5

Have: 2k disjoint breaks B_1, \dots, B_{2k} in P such that

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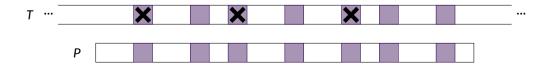
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 Run [LV'89] only for positions in T where at least k breaks match exactly ~> How?



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Philip Wellnitz 31-9

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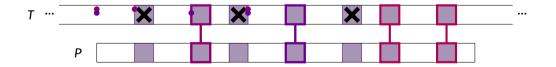
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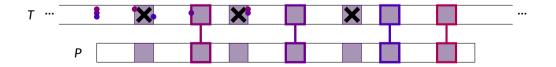
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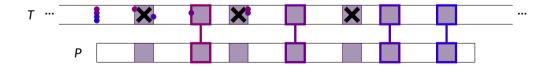
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- ◆ If $B_i = P[\ell_i...) = T[a...)$, add a mark to $T[\lfloor (a \ell_i)/k \rfloor]$; run [LV'89] for pos w/ ≥ k marks $\rightarrow O(k^2|T|/|P|)$ marks in total; $\rightarrow O(k|T|/|P|)$ calls to [LV'89]; $\rightarrow O(k^3|T|/|P|)$ time



How do we turn our insights into faster algorithms?

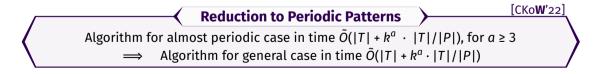
Need to tackle three one case.

- P contains 2k disjoint breaks; \rightsquigarrow Adaption of [Cole,Hariharan'98] yields $O(|T| + |T|/|P| \cdot k^3)$.
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Reduction to Periodic Patterns

Algorithm for almost periodic case in time $\tilde{O}(|T| + k^a \cdot |T|/|P|)$, for $a \ge 3$ \implies Algorithm for general case in time $\tilde{O}(|T| + k^a \cdot |T|/|P|)$

Need to tackle: P is almost periodic

- In [CKoW'20], use elaborate marking scheme to obtain O(|T| + |T| / |P| · k⁴) algorithm
 → Not faster than [Cole, Hariharan'98]
- In [CKoW'22], trade-off between
 - Refinement of algorithm from [CKoW'20]
 - New algorithm based on "Seaweed Technology" of [Tiskin'10,'15]
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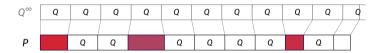
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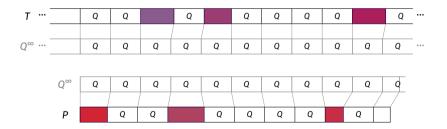
Have: P is at ED $\leq 2k$ to a string with period Q of len O(|P|/k)

If P is close to Q[∞], then so is T → have alignments A_p : P → Q[∞] and A_T : T → Q[∞]
 A_T and A_p induce tile partition of P and T; all but O(k) tiles are Q



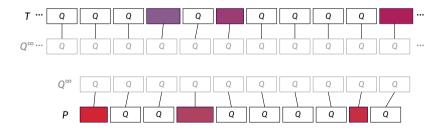
Have: P is at ED $\leq 2k$ to a string with period Q of len O(|P|/k)

• If P is close to Q^{∞} , then so is $T \rightsquigarrow$ have alignments $A_p : P \rightsquigarrow Q^{\infty}$ and $A_T : T \rightsquigarrow Q^{\infty}$ A_T and A_n induce tile partition of P and T: all but O(k) tiles are O



Have: P is at ED $\leq 2k$ to a string with period Q of len O(|P|/k)

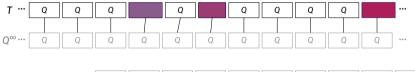
• If P is close to Q^{∞} , then so is $T \rightsquigarrow$ have alignments $A_p : P \rightsquigarrow Q^{\infty}$ and $A_T : T \rightsquigarrow Q^{\infty}$ A_T and A_p induce tile partition of P and T; all but O(k) tiles are Q

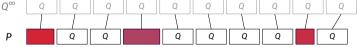


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- If P is close to Q^{∞} , then so is $T \rightsquigarrow$ have alignments $A_p : P \rightsquigarrow Q^{\infty}$ and $A_T : T \rightsquigarrow Q^{\infty}$ A_T and A_p induce tile partition of P and T; all but O(k) tiles are Q
- ◆ Imagine shifting P along T, one tile/Q at a time
 → For each shift, want to compute occs

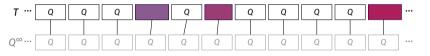
→ Between shifts, O(k) tiles get aligned to a new/different tile

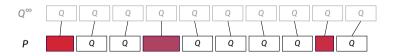




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- Imagine shifting *P* along *T*, one tile/*Q* at a time
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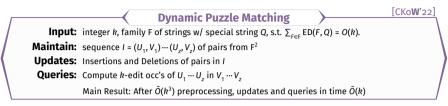
Dynamic Puzzle Matching[CKoW'22]Input: integer k, family F of strings w/ special string Q, s.t. $\Sigma_{F \in F}$ ED(F, Q) = O(k).Maintain: sequence $I = (U_1, V_1) \cdots (U_z, V_z)$ of pairs from F²Updates: Insertions and Deletions of pairs in IQueries: Compute k-edit occ's of $U_1 \cdots U_z$ in $V_1 \cdots V_z$

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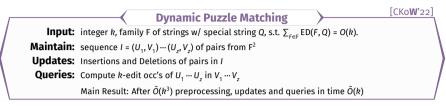
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After $\tilde{O}(k^3)$ preprocessing, updates and queries in time $\tilde{O}(k)$



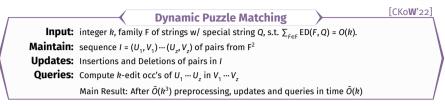
• For simplicity: assume $|Q| \approx \sqrt{|P|}$; ignore handling of initial and final pairs





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- > k copies of Q in $P \implies \ge 1$ copy of Q matched exactly

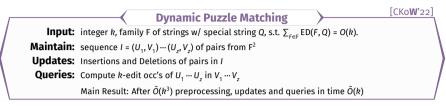




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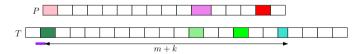
→ Starting pos of *k*-edit occ's in *T* within *O*(*k*) from endpoints of tiles

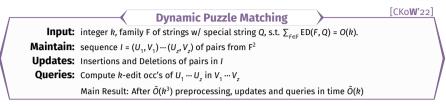




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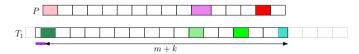
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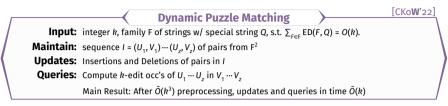




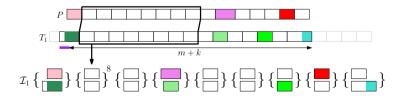
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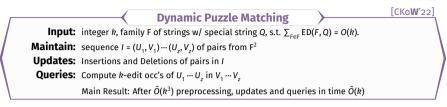




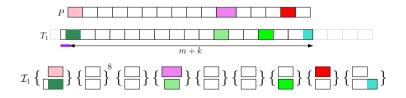
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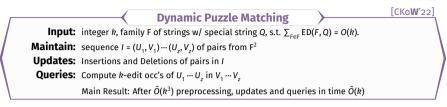
Philip Wellnitz 35-From Strings to Seaweeds: Modern Tools for Classical Problems



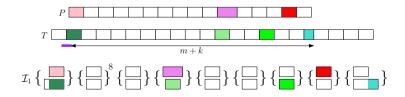
- For simplicity: assume $|Q| \approx \sqrt{|P|}$; ignore handling of initial and final pairs
- Goal: Iterate over all I_i's with one DPM instance

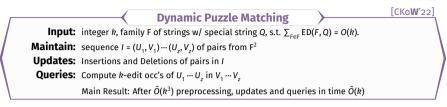


Philip Wellnitz 35-From Strings to Seaweeds: Modern Tools for Classical Problems

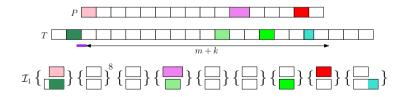


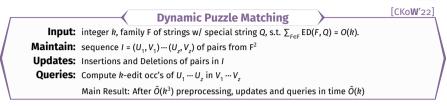
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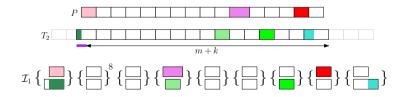


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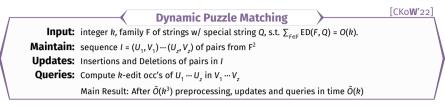




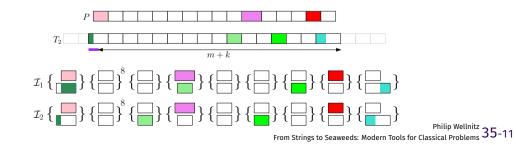
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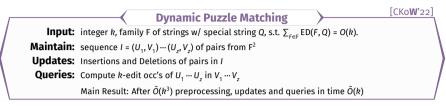


Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 35-10



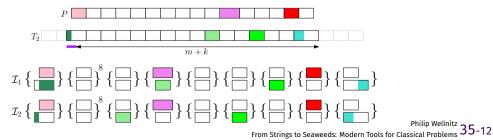
- For simplicity: assume $|Q| \approx \sqrt{|P|}$; ignore handling of initial and final pairs
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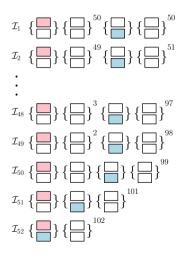




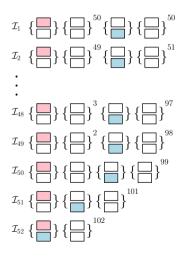
- For simplicity: assume $|Q| \approx \sqrt{|P|}$; ignore handling of initial and final pairs
- Goal: Iterate over all I_i 's with one DPM instance

 \rightsquigarrow Over $\Theta(\sqrt{|P|})$ shifts of P, need $O(\sqrt{|P|}k)$ DPM-updates $\rightsquigarrow \tilde{O}(k^3 + \sqrt{|P|}k^2)$ time



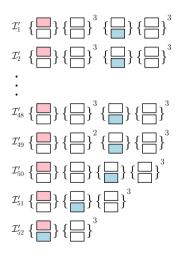


Example k = 2.

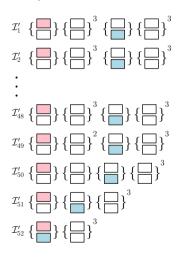


For plain run (Q, Q)^y, at least y - k copies of Q matched exactly in k-edit occ

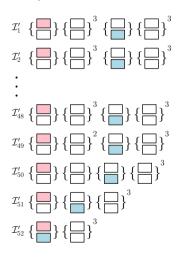
Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems **36-2**



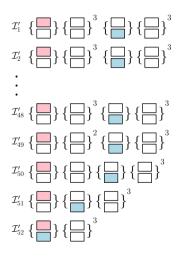
- For plain run (Q, Q)^y, at least y k copies of Q matched exactly in k-edit occ
- Cap exponents of plain runs at k + 1



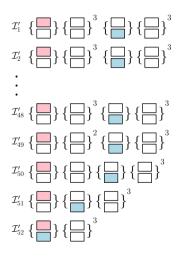
- For plain run (Q, Q)^y, at least y k copies of Q matched exactly in k-edit occ
- Cap exponents of plain runs at k + 1
 - $\rightsquigarrow O(k)$ DPM-updates per pair of special tiles $\rightsquigarrow O(k^2)$ pairs of special tiles $\rightsquigarrow \tilde{O}(k^4)$ time



- For plain run (Q, Q)^y, at least y k copies of Q matched exactly in k-edit occ
- Cap exponents of plain runs at \sqrt{k}
 - $\rightsquigarrow O(\sqrt{k})$ DPM-updates per pair of special tiles $\rightsquigarrow O(k^2)$ pairs of special tiles $\rightsquigarrow \tilde{O}(k^{3.5})$ time

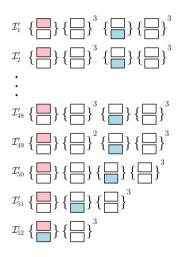


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 → Filter out false positives using another marking scheme → Õ(k^{3.5}) time in total (boring)

Example k = 2.

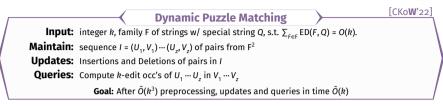


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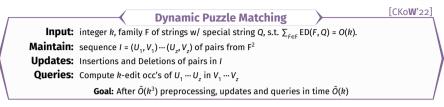
Main Result

Pattern P, text T, threshold k; can compute starting pos of all k-edit occ's in time $\tilde{O}(|T| + k^{3.5} |T|/|P|)$.

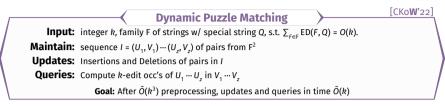
[CKo**W**'22]



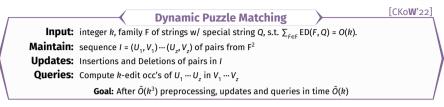
- Store edit distance information for each pair $(U_i, V_i) \rightarrow$ suitable permutation matrices allow this in O(k) space
- Show how to compose the information of different pairs in a suitable way ~---"seaweed product"
- Show how to compute said product efficiently



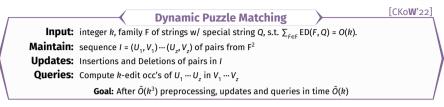
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General Idea

- Store edit distance information for each pair $(U_i, V_i) \rightsquigarrow$ suitable permutation matrices allow this in O(k) space
- Show how to compose the information of different pairs in a suitable way ~---"seaweed product"
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Content Warning: Some technical computations ahead!

A ∈ {0,1}^{n×n} is permutation matrix: every row and every column has exactly one 1
 → 1-to-1 corresponds to permutation in S_n

• For $n \times n$ matrix A, distribution matrix A^{Σ} is $(n + 1) \times (n + 1)$ matrix with

$$A^{\Sigma}[i,j] := \sum_{i' \ge i} \sum_{j' < j} A[i',j']$$

• For $(n + 1) \times (n + 1)$ matrix A, density matrix A^{\Box} is $n \times n$ matrix with

$$A^{\Box}[i,j] := A[i+1,j] + A[i,j+1] - A[i+1,j+1] - A[i,j]$$

$$\begin{pmatrix} (0,0) & (n-1,0) \\ 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix}$$

$$(0,n-1) & (n-1,n-1)$$

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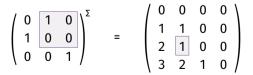
/ 0	1	0 \		• • •
1	0	0) 0 1 /	$ \left(\begin{array}{rrrr} 1 & 2 & 3 \\ 2 & 1 & 3 \end{array}\right) $	\times i
\ 0	0	1 /	(2 1 3)	

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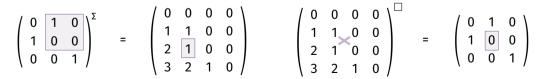


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Lemma (Relation Density-Distribution Matrix)

For any A, have $(A^{\Sigma})^{\Box} = A$; for simple A, have $(A^{\Box})^{\Sigma} = A$ (first row and last col of A are o)

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Lemma (Properties of Permutation Matrices)

Perm. matrix A; $A^{\Sigma}[i, 0] = A^{\Sigma}[n-1, j] = 0$ and $A^{\Sigma}[0, j] = j$; $A^{\Sigma}[i, 0] = n-1 - i$ and $A^{\Sigma}[i, j] \ge j - i$

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(Working with Permutation Matrices) [Tiso7, CP10] $For <math>n \times n$ permutation matrix A can store in O(n) space and can access $A^{\Sigma}[i, j]$ in $O(\log n / \log \log n)$

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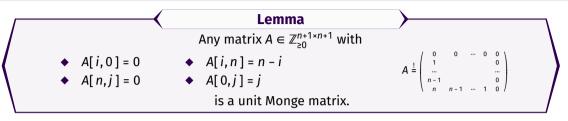
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Detour: Characterizing Unit-Monge Matrices

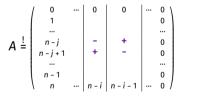


• Show: each row/column of A^{\Box} sums to 1

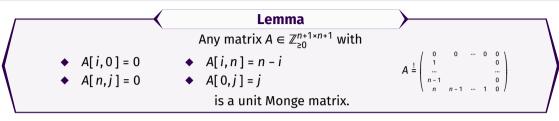
Consider

$$\sum_{j} A^{\Box}[i,j] = \sum_{j} A[i+1,j] + A[i,j+1] - A[i+1,j+1] - A[i,j]$$

= $A[i+1,0] + A[i,n] - A[i+1,n] - A[i,0]$
= $0 + (n-i) - (n-i-1) - 0$
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Detour: Characterizing Unit-Monge Matrices

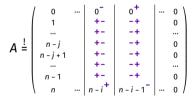


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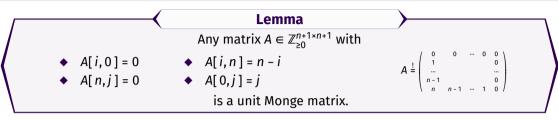
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Detour: Characterizing Unit-Monge Matrices

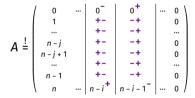


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Monge Matrix × (min,+)-Product

- A is simple: first row and last col are o; for simple A have $A^{\Sigma \Box} = A^{\Box \Sigma} = A$
- ♦ A is Monge if $A^{\Box} \ge 0$; A is unit-Monge if A^{\Box} is a permutation matrix
- ◆ For matrices A, B, the (min-+)-product A ⊙ B is

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(A \circ B)[i,k] := \min_{j} (A[i,j] + B[j,k])
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Recall: For adjacency matrix A of (weighted) graph G, A^{ol} stores l-hop dist in G

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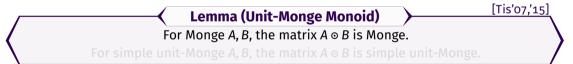
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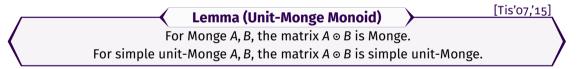
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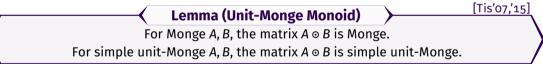
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Recall: For adjacency matrix A of (weighted) graph G, $A^{\circ \ell}$ stores ℓ -hop dist in G



Proof: Board or exercise.

- A is simple: first row and last col are o; for simple A have $A^{\Sigma \Box} = A^{\Box \Sigma} = A$
- ◆ A is Monge if $A^{\Box} \ge 0$; A is unit-Monge if A^{\Box} is a permutation matrix
- ◆ For matrices A, B, the (min-+)-product A ⊙ B is

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```

Lemma (Unit-Monge Monoid)

For Monge A, B, the matrix $A \circ B$ is Monge. For simple unit-Monge A, B, the matrix $A \circ B$ is simple unit-Monge.

• For perm matrices A, B, define seaweed product $A \square B := (A^{\Sigma} \odot B^{\Sigma})^{\square}$

```
• A \square B is different from normal perm concat: I^R = \begin{pmatrix} 0 & \cdots & 0 & 1 \\ 1 & 0 & 1 & 0 \\ 0 & 1 & 0 & 1 \\ 1 & 0 & \cdots & 0 \end{pmatrix} is a zero
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Philip Wellnitz 41-

[Tis'07,'15]

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Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 41-8

[Tis'07,'15]

$$\begin{pmatrix} 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad \bullet \quad \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0 & 1 \\ 0 & 1 & 0 \end{pmatrix} =$$

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$$\begin{pmatrix} 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix}
\models \begin{pmatrix} 1 & 0 & 0 \\ 0 & 0 & 1 \\ 0 & 1 & 0 \end{pmatrix} = \begin{pmatrix} 0 & 0 & 1 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \end{pmatrix} \\ \stackrel{\checkmark}{\downarrow} (\cdot)^{\Sigma} \qquad \stackrel{\checkmark}{\downarrow} (\cdot)^{\Sigma} \qquad \stackrel{\checkmark}{\downarrow} (\cdot)^{\Box} \\ \begin{pmatrix} 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 \\ 2 & 1 & 0 & 0 \\ 3 & 2 & 1 & 0 \end{pmatrix}
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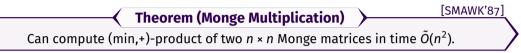
$$\begin{pmatrix} 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix} \quad \bullet \quad \begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix} = \\ \begin{matrix} \zeta (\cdot)^{\Sigma} & & \zeta (\cdot)^{\Sigma} \\ \begin{pmatrix} 0 & 0 & 0 & 0 \\ 1 & 1 & 0 & 0 \\ 2 & 1 & 0 & 0 \\ 3 & 2 & 1 & 0 \end{pmatrix} \\ \bullet \begin{pmatrix} 0 & 0 & 0 & 0 \\ 1 & 1 & 1 & 0 \\ 2 & 2 & 1 & 0 \\ 3 & 2 & 1 & 0 \end{pmatrix}$$

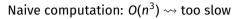
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Naive computation: $O(n^3) \rightsquigarrow$ too slow

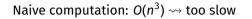
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(Proof skipped.)



Theorem (Monge Multiplication)

Can compute (min,+)-product of two $n \times n$ Monge matrices in time $\tilde{O}(n^2)$.

(Proof skipped.)

Theorem (Unit Monge Multiplication)

Can compute (min,+)-product of two $n \times n$ simple unit-Monge matrices in time $O(n \log n)$.

[SMAWK'87]

[Tis'07,'15]

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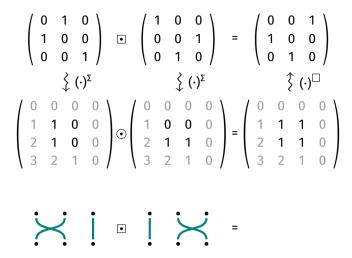
Proof idea (details messy, but not too complicated).

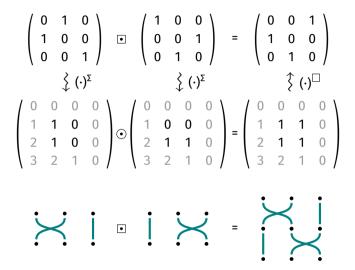
- Divide and Conquer: Split each matrix into two; reduce to just two subproblems
- For conquer step, use elaborate, but easy to compute function to decide which solution to propagate from subproblem

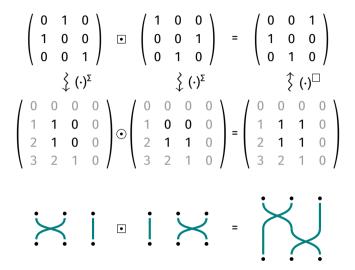
[SMAWK'87]

[Tis'07,'15]

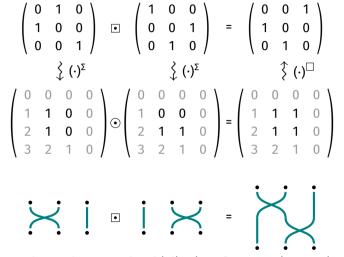
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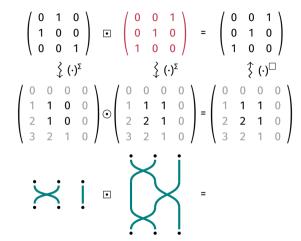
Philip Wellnitz 45-4

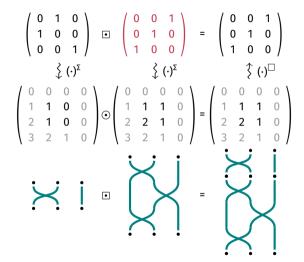


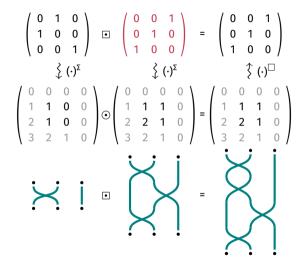
Example matches normal multiplication of permutation matrices.

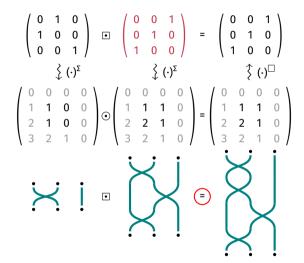
Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 45-5

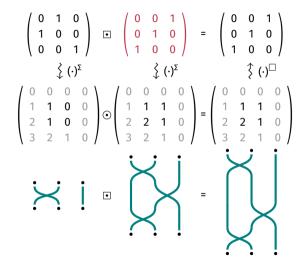
$$\begin{pmatrix} 0 & 1 & 0 \\ 1 & 0 & 0 \\ 0 & 0 & 1 \end{pmatrix} \square \begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix} = \begin{pmatrix} 0 & 0 & 1 \\ 0 & 1 & 0 \\ 1 & 0 & 0 \end{pmatrix}$$
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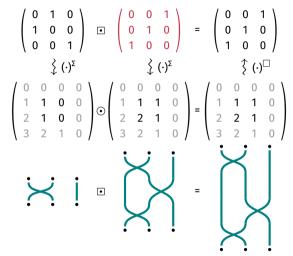




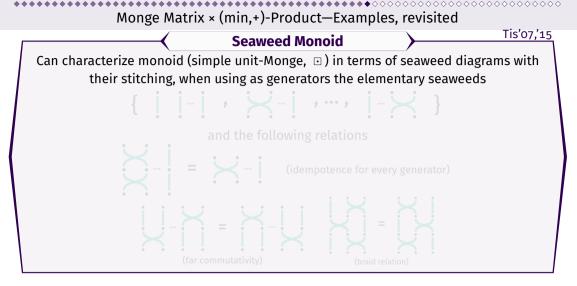






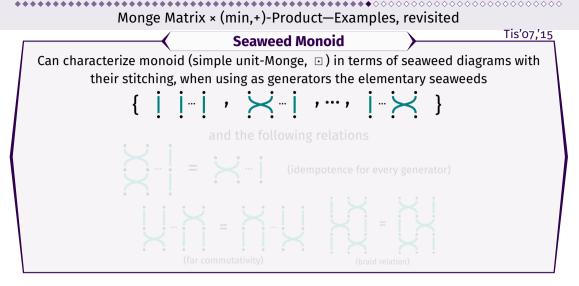


Can view seaweed product in terms of these diagrams, but need some extra simplification rules.



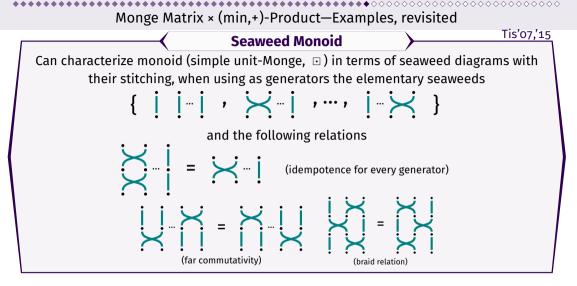
- Elementary seaweeds are just identity matrix and transposition matrices.
- Idempotence is crucial difference from standard perm conca

From Strings to Seaweeds: Modern Tools for Classical Problems 47-1



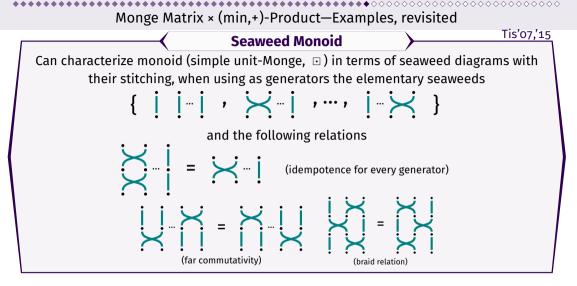
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Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 47-2



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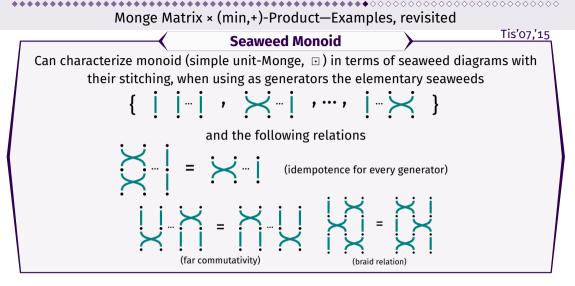
From Strings to Seaweeds: Modern Tools for Classical Problems 47-3



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Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 47



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Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 47-5

A Final Example: Stitching with *I*

• Consider following task:

Have permutations $(1, ..., n) \mapsto (\sigma(1), ..., \sigma(n))$ and $(1, ..., n) \mapsto (\varrho(1), ..., \varrho(n))$ Want permutation $(1, ..., n, n + 1, ..., 2n) \mapsto (\sigma(1), ..., \sigma(n), \varrho(1), ..., \varrho(n))$

→ Prime application: string concatenation

♦ Claim:

$$P_{\sigma\varrho} = \begin{pmatrix} P_{\sigma} & 0\\ 0 & P_{\varrho} \end{pmatrix} = \begin{pmatrix} P_{\sigma} & 0\\ 0 & I_{|\varrho|} \end{pmatrix} = \begin{pmatrix} I_{|\sigma|} & 0\\ 0 & P_{\varrho} \end{pmatrix}$$

• Easy proof via seaweeds:



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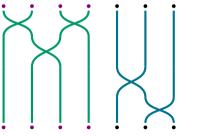
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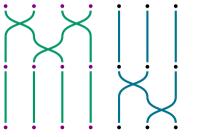
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Seaweeds in a Nutshell

- Somewhat weird matrix product (min-plus product of prefix-sum matrices of permutation matrices)
- Seaweed product can be computed efficiently

~> Important special case: stitching with I)

• Next, and most importantly: can reinterpret the original DP matrix for ED as prefix-sum matrix of a permutation matrix (after some massaging)

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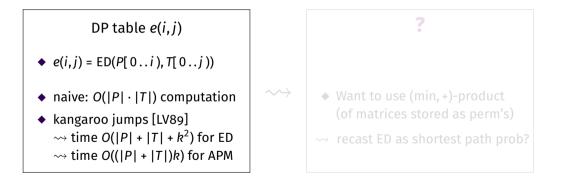
Using Seaweeds for Faster ED Computations

Given *P*, *T*, need to compute ED between arbitrary substrings ~> **Goal**: Use seaweeds for speed-ups over standard DP



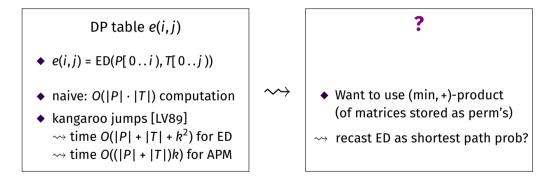
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Recall classical DP (again):

- ♦ Write e(i, j) for the ED of T[0..i) and P[0..j)
 → e(t, p) is the ED of T and P.
- Clearly, e(i, 0) = i (delete T[0..i)) and e(0, j) = j (insert P[0..j))

Observe

	e(i,j) = min {				e(i – 1,j) + 1 e(i,j – 1) + 1 e(i – 1,j – 1)			(delete from T) (insert in T) (match/subst)										
						l		+(T	[i]≠	P[j])		(m	ato	:h/	su	bs	t)	
	S	a	r	r	e	b	r					s				-		r
S										s	0	1 0 1	2	3	4	5	6	7
a										a	1	0	1	2	3	4	5	6
a			_							а	2	1	0		2	3	4	5
r			-e(i,j)						r	3	2	1-					
b							_	L		b :	4							
:							-e(t, p)		-	5							

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$e(i,j) = \min \left\{ \right.$		(delete from <i>T</i>) (insert in <i>T</i>)						
l		(match/subst)						
$\mathbf{s} \mathbf{a} \mathbf{r} \mathbf{r} \mathbf{e} \mathbf{b}$	r a a r	s a r r e b r 0 1 2 3 4 5 6 7 1 0 1 2 3 4 5 6 2 1 0 1 2 3 4 5 6 3 2 1						

5

Consider different,

but related alignment graph:

- Vertex (i, j) for every $i \in [0..|T|], j \in [0..|P|]$
- Horizontal edge (i, j) → (i + 1, j) of weight 1 (deletion from T)
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- Diagonal edge (i, j) → (i + 1, j + 1) of weight (T[i] ≠ P[j]) (match/subst)

→ shortest (0, 0) to (|T|, |P|) path is ED of T and P
 → recover DP by running Dijkstra from (0, 0)





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]	e(i – 1, j) + 1	(delete from T)						
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sar reb		sarrebr						
	C.	0 1 2 3 4 5 6 7						
	5	0 1 2 3 4 5 6 7 1 0 1 2 3 4 5 6						
	a	2 1 0 1 2 2 4 5						

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s a r r e b	r a a r b :	s a r r e b r … 0 1 2 3 4 5 6 7 1 0 1 2 3 4 5 6 2 1 0 1 2 3 4 5 3 2 1 1 4 5

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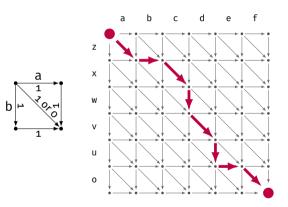


Alignment Graph of P and T

- Vertex (i,j) for every $i \in [0..|T|], j \in [0..|P|]$
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 \rightarrow shortest (0, 0) to (|T|, |P|) path is ED of T and P

- shortest (*i*, *j*) to (*i*', *j*') path is ED of T[*i*..*i*') and P[*j*..*j*'))
- ↔ For stitching: Need (just)
 boundary-to-boundary dist's
- Define D_{P,T}[a, b] := dist(in_a, out_b)



dist((0,0),(|T|,|P|)) = ED(T,P)

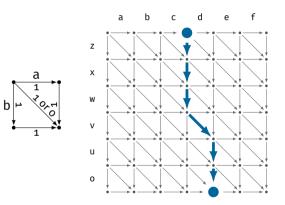
Alignment Graph of P and T

- Vertex (i,j) for every $i \in [0..|T|], j \in [0..|P|]$
- Horizontal edge (i, j) → (i + 1, j) of weight 1 (deletion from T)
- Vertical edge (i, j) → (i, j + 1) of weight 1 (insertion in T)
- Diagonal edge (i,j) → (i + 1, j + 1) of weight (T[i] ≠ P[j]) (match/subst)

 \rightarrow shortest (0, 0) to (|T|, |P|) path is ED of T and P

shortest (*i*, *j*) to (*i*', *j*') path is ED of T[*i*..*i*') and P[*j*..*j*'))

- ↔ For stitching: Need (just)
 boundary-to-boundary dist's
- Define D_{P,T}[a, b] := dist(in_a, out_b)



dist((i, 0), (i', |P|)) = ED(T[i ... i'), P)

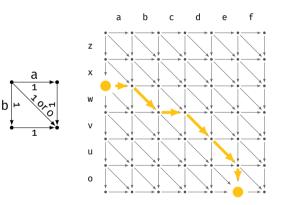
Alignment Graph of P and T

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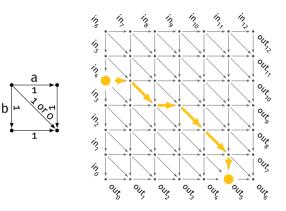
dist((i, j), (i', j')) = ED(T[i ... i'), P[j ... j'))

Alignment Graph of P and T

- Vertex (i,j) for every $i \in [0..|T|], j \in [0..|P|]$
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- shortest (i, j) to (i', j') path is ED of T[i..i') and P[j..j'))
- → For stitching: Need (just)
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 - Define D_{P,T}[a, b] := dist(in_a, out_b)



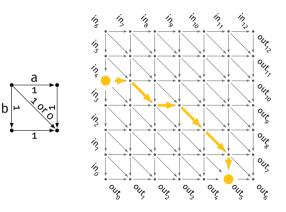
dist $(in_{a}, out_{b}) = ED(T[max(a - |P|, 0)..min(b, |T|)),$ P[max(|P| - a, 0)..min(|P| + |T| - b, |P|)))

Alignment Graph of P and T

- Vertex (i,j) for every $i \in [0..|T|], j \in [0..|P|]$
- Horizontal edge (i, j) → (i + 1, j) of weight 1 (deletion from T)
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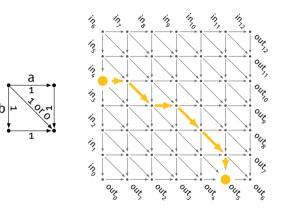


 $\begin{aligned} \mathsf{dist}(\mathsf{in}_a,\mathsf{out}_b) &= \mathsf{ED}(T[\max(a - |P|, 0) \dots \min(b, |T|)), \\ P[\max(|P| - a, 0) \dots \min(|P| + |T| - b, |P|))) \end{aligned}$

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- ◆ Problem: Some D_{p,T}[a, b] are ∞
 → use undirected edges
- Suffices to show (recall earlier slide):

Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$



dist (in_a , out_b) = ED(T[max(a - |P|, 0)..min(b, |T|)), P[max(|P| - a, 0)..min(|P| + |T| - b, |P|)))

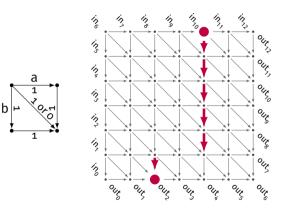
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Philip Wellnitz

Consider Alignment Graph of P and T

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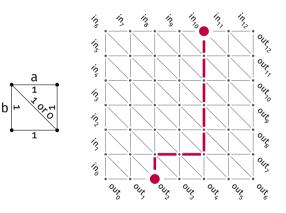


dist(in_a , out_b) = ED(T[max(a - |P|, 0)..min(b, |T|)), P[max(|P| - a, 0)..min(|P| + |T| - b, |P|)))

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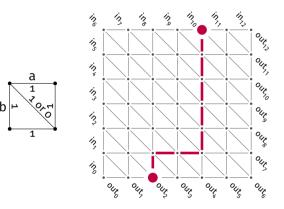
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dist (in_a , out_b) * ED(T[max(a - |P|, 0) ... min(b, |T|)), P[max(|P| - a, 0) ... min(|P| + |T| - b, |P|)))

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
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- Suffices to show (recall earlier slide):
 - $D_{P,T}$ is non-negative and integer • Border satisfies (n := |P| + |T|) $D_{P,T} \doteq \begin{pmatrix} 0 & 0 & 0 & 0 \\ 0 & - & 0 & 0 \\$

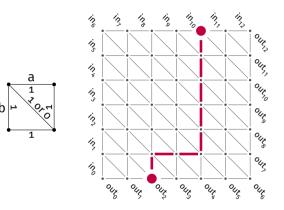


dist (in_a, out_b) $\stackrel{*}{=}$ ED(T[max(a - |P|, 0).. min(b, |T|)), P[max(|P| - a, 0).. min(|P| + |T| - b, |P|)))

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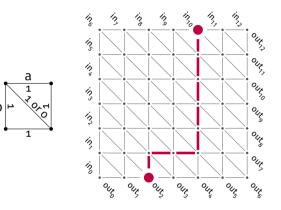


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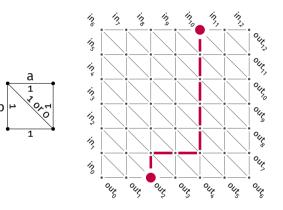


dist (in_a, out_b) $\stackrel{*}{=}$ ED(T[max(a - |P|, 0).. min(b, |T|)), P[max(|P| - a, 0).. min(|P| + |T| - b, |P|)))

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- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):
 - *D_{P,T}* is non-negative and integer ✓
 Border satisfies (*n* := |*P*| + |*T*|)
 D_{P,T} = $\begin{pmatrix}
 0 & 0 & \cdots & 0 & 0 \\
 1 & \cdots & 0 & 0 \\
 n & -1 & \cdots & 1 & 0 \\
 n & -1 & \cdots & 1 & 0
 \end{aligned}$ *D_{P,T}*[*a*, 0] = 0*D_{P,T}*[*a*, 0] =*b*

Idea: Force boundary values to be correct $\rightarrow D'_{P,T}[a, b] := (D_{P,T}[a, b] - a + b)/2$

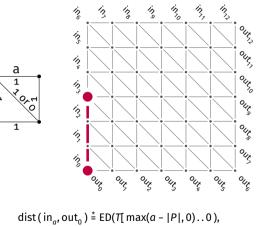


dist (in_a , out_b) * ED(T[max(a - |P|, 0)..min(b, |T|)), P[max(|P| - a, 0)..min(|P| + |T| - b, |P|)))

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
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 Border satisfies (*n* := |*P*| + |*T*|)
 D_{P,T} = $\begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & \cdots & 0 & 0 \\ \cdots & \cdots & 0 & 0 \\ n-1 & \cdots & 1 & 0 & 0 \end{pmatrix}$ *D_{P,T}*[*a*, 0] = 0
 D_{P,T}[*a*, n] = *n a D_{P,T}*[*n*, *b*] = 0
 D_{P,T}[0, *b*] = *b*

Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$

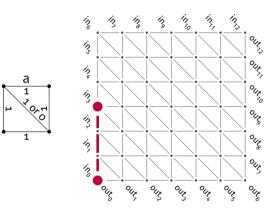


P[max(|P| - a, 0) . . |P|))

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
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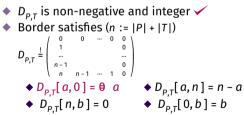
Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$



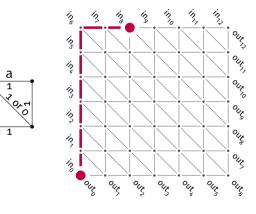
dist (in_a, out₀) = ED(ε , P[max(|P| - a, 0).. |P|)) = a

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
 → use undirected edges
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):



Idea: Force boundary values to be correct $\rightarrow D'_{p,\tau}[a, b] := (D_{p,\tau}[a, b] - a + b)/2$



 $dist(in_a, out_0) = a$

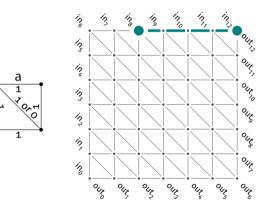
h

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
 → use undirected edges
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):

•
$$D_{P,T}$$
 is non-negative and integer
• Border satisfies $(n := |P| + |T|)$
 $D_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & \cdots & 0 & 0 \\ n & 1 & \cdots & 0 \\ n & n & 1 & \cdots & 1 \end{pmatrix}$
• $D_{P,T}[a, 0] = \Theta \quad a$
• $D_{P,T}[a, n] = n - a$
• $D_{P,T}[n, b] = 0$
• $D_{P,T}[0, b] = b$

Idea: Force boundary values to be correct $\rightarrow D'_{p,\tau}[a, b] := (D_{p,\tau}[a, b] - a + b)/2$

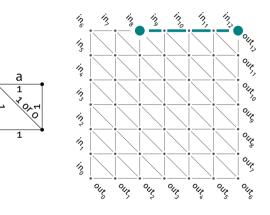


 $dist(in_a, out_n) = n - a$

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
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Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$

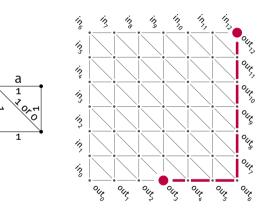


dist(in_a, out_n) = n - a

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
 → use undirected edges
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):
 - $D_{P,T}$ is non-negative and integer • Border satisfies (n := |P| + |T|) $D_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & & & 0 & 0 \\ n & -1 & & & 0 & 0 \\ n & n & -1 & & 0 & 0 \end{pmatrix}$ • $D_{P,T}[a, 0] = \Theta \ a$ • $D_{P,T}[a, n] = n - a$ • $D_{P,T}[n, b] = \Theta \ n - b$ • $D_{P,T}[0, b] = b$

Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$

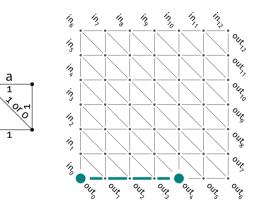


 $dist(in_n, out_b) = n - b$

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
 → use undirected edges
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):

Idea: Force boundary values to be correct $\rightarrow D'_{p,T}[a, b] := (D_{p,T}[a, b] - a + b)/2$

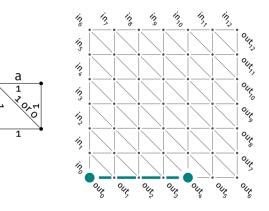


 $dist(in_0, out_b) = b$

Consider Alignment Graph of P and T

- Define D_{P,T}[a, b] := dist(in_a, out_b)
- ◆ Problem: Some D_{P,T}[a, b] are ∞
 → use undirected edges
- \rightsquigarrow Hope: $D_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):
 - $D_{P,T}$ is non-negative and integer • Border satisfies (n := |P| + |T|) $D_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & \cdots & 0 & 0 \\ n-1 & \cdots & 0$

Idea: Force boundary values to be correct $\rightsquigarrow D'_{P,T}[a, b] := (D_{P,T}[a, b] - a + b)/2$



dist (in_a , out_b) * ED(T[max(a - |P|, 0)..min(b, |T|)), P[max(|P| - a, 0)..min(|P| + |T| - b, |P|)))

Consider Alignment Graph of P and T

• Define $D_{P,T}[a, b] := dist(in_a, out_b)$ and $D'_{P,T}[a, b] := (D_{P,T}[a, b] - a + b)/2$

\rightsquigarrow Hope: $D'_{P,T}^{\Box}$ is permutation matrix

- Suffices to show (recall earlier slide):
 - D'_{P,T} is non-negative and integer
 - Border satisfies (n := |P| + |T|

$$D'_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & & & 0 \\ \cdots & & & \cdots \\ n-1 & & 0 \\ n & n-1 & \cdots & 1 & 0 \end{pmatrix}$$

Have:

◆ $D'_{p,T}[a, n] = ((n - a) - a + n)/2 = n - a$ ◆ $D'_{p,T}[0, b] = (b - 0 + b)/2 = b$

Consider Alignment Graph of P and T

• Define $D_{PT}[a, b] := dist(in_a, out_b)$ an

d
$$D'_{P,T}[a, b] := (D_{P,T}[a, b] - a + b)/2$$

- \rightsquigarrow Hope: D'_{PT}^{\Box} is permutation matrix
- Suffices to show (recall earlier slide):
 - D'_{P τ} is non-negative and integer
 - Border satisfies (n := |P| + |T|)

$$D'_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & & 0 \\ \cdots & & \cdots \\ n & n - 1 & \cdots & 1 & 0 \end{pmatrix}$$

Have:

♦
$$D'_{P,T}[a, 0] = (a - a + 0)/2 = 0$$

♦ $D'_{P,T}[n, b] = ((n - b) - n + b)/2 = 0$

◆ $D'_{P,T}[a, n] = ((n - a) - a + n)/2 = n - a$ ◆ $D'_{P,T}[0, b] = (b - 0 + b)/2 = b$

Consider Alignment Graph of P and T

• Define $D_{PT}[a, b] := dist(in_a, out_b)$ and

d
$$D'_{P,T}[a, b] := (D_{P,T}[a, b] - a + b)/2$$

- \rightarrow Hope: D'_{PT}^{\Box} is permutation matrix
- Suffices to show (recall earlier slide):
 - D'_{P τ} is non-negative and integer
 - Border satisfies (n := |P| + |T|)

$$D'_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & & 0 & \\ \cdots & & \cdots & \cdots & \\ n & n & -1 & \cdots & 1 & 0 \end{pmatrix} \checkmark$$

Have:

♦
$$D'_{P,T}[a, 0] = (a - a + 0)/2 = 0$$

♦ $D'_{P,T}[n, b] = ((n - b) - n + b)/2 = 0$

• $D'_{P,T}[a, n] = ((n - a) - a + n)/2 = n - a$ • $D'_{PT}[0,b] = (b-0+b)/2 = b$

Consider Alignment Graph of P and T

• Define $D_{p_T}[a, b] := \text{dist}(in_a, \text{out}_b)$ and $D'_{p_T}[a, b] := (D_{p_T}[a, b] - a + b)/2$

- \rightarrow Hope: D'_{PT}^{\Box} is permutation matrix
- Suffices to show (recall earlier slide):
 - D'_{PT} is non-negative and integer
 - Border satisfies (n := |P| + |T|)

$$D'_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & & 0 & \\ \cdots & & & \cdots & \\ n-1 & & & 0 \\ n & n-1 & \cdots & 1 & 0 \end{pmatrix} \checkmark$$

Have:

♦
$$D'_{P,T}[a, 0] = (a - a + 0)/2 = 0$$

♦ $D'_{P,T}[n, b] = ((n - b) - n + b)/2 = 0$

• $D'_{P,T}[a,n] = ((n-a) - a + n)/2 = n - a$ • $D'_{PT}[0,b] = (b-0+b)/2 = b$

Consider Alignment Graph of P and T

- Define $D_{P,T}[a, b] := \text{dist}(\text{in}_a, \text{out}_b)$ and $D'_{P,T}[a, b] := (D_{P,T}[a, b] a + b)/2$
- \rightsquigarrow Hope: $D'_{P,T}^{\Box}$ is permutation matrix
- Suffices to show (recall earlier slide):
 - ◆ D'_{P,T} is non-negative and integer → Problem!
 - Border satisfies (n := |P| + |T|)

$$D'_{P,T} \stackrel{!}{=} \begin{pmatrix} 0 & 0 & \cdots & 0 & 0 \\ 1 & \cdots & & 0 \\ \cdots & \cdots & \cdots & \cdots \\ n-1 & \cdots & 1 & 0 \end{pmatrix} \checkmark$$

Have:

♦
$$D'_{P,T}[a, 0] = (a - a + 0)/2 = 0$$

♦ $D'_{P,T}[n, b] = ((n - b) - n + b)/2 = 0$

◆ $D'_{P,T}[a, n] = ((n - a) - a + n)/2 = n - a$ ◆ $D'_{P,T}[0, b] = (b - 0 + b)/2 = b$

Substitutions Are Too Cheap!

• Define $D_{P,T}[a,b] := \text{dist}(\text{in}_a, \text{out}_b)$ and $D'_{P,T}[a,b] := (D_{P,T}[a,b] - a + b)/2$ \rightarrow Hope: $D'_{P,T}^{\Box}$ is permutation matrix \rightarrow Problem: $D'_{P,T}^{\Box}[a,b]$ not integer

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• Every character subst. flips parity of dist $(in_a, out_b) \rightsquigarrow$ "forbid subst's" / increase cost to 2



dist (in_a , out_b) = |b - a|good (|b - a| - a + b always divisible by 2)



dist (in_a, out_b) = max($|x_a - x_b|$, $|y_a - y_b|$) – LCS($P_{a,b}, T_{a,b}$) red: cost 1 (bad), green: cost 0 (saving of 2, good)

Philip Wellnitz 55-1

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Philip Wellnitz 55-2

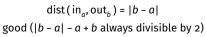
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Philip Wellnitz 55-3

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Deletion Distance (DD)

Min number of insertions or deletions of characters to transform T into P.

Embedding DD into ED

Write $S^{\ddagger} := S[0] \ddagger S[1] \ddagger \dots S[|S|] \ddagger$. Then $DD(P^{\ddagger}, T^{\ddagger}) = 2ED(P, T)$.

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 In alignment graph: corresponds to removing all diag edges of weight 1 (and adding vertices/edges corresponding to \$)

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From Strings to Seaweeds: Modern Tools for Classical Problems 50-4

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Main ResultConsider (deletion distance) alignment graph of $P^{\$}$ and $T^{\$}$;define $D_{P^{\$,T}\$}[a,b] := dist(in_a, out_b)$ and $D'_{P^{\$,T}\$}[a,b] := (D_{P^{\$,T}\$}[a,b] - a + b)/2$.
Then, $D'_{P^{\$,T}\$}^{\Box}$ is a permutation matrix.

- Can (easily) show: (min, +)-product of D'_{p\$,7}\$ matrices ("stitching") yields deletion distances for concatenated strings
 → Can use seaweed product for fast computation
- Some technicalities (skipped):
 - stitching requires some overlap in at least one string
 - ♦ want to compute only ≈ k diagonals; requires a "restriction" operation
 - formally defining the object we use to represent pairs of strings in DPM
 - **•** ...

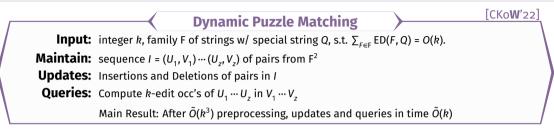
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 - formally defining the object we use to represent pairs of strings in DPM
 - **•** ...

Using Dynamic Puzzle Matching



General Idea

- Store edit distance information for each pair (U_i, V_i) \sim Seaweed/permutation matrix of (U_i, V_i) allow this in O(k) space
 - Set we can be it the matrix of $(0_j, v_j)$ allow this in O(x) space
- $oldsymbol{\diamond}$ In preprocessing, build matrix for every possible pair of strings from family F
- Use seaweed product of [Tis'07,'15] to compose the information of different pairs Õ(k) per stitch
- Use BST on top to support update queries

Adding Weights

What about the weighted setting?

Adding Weights

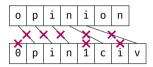
What about the weighted setting?

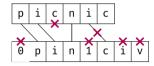
10. The weighted case. In the weighted case, deletions of different characters and the various substitutions may have differing costs, but, by way of normalization, all will be required to have cost at least 1.

The approximate matches with $\cot \leq k$ can be found using essentially the same algorithm; the only change needed is to the Landau–Vishkin algorithm to take into account the differing costs. The details are left to the reader.

—Cole, Hariharan, 2002

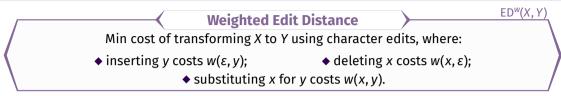




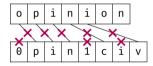


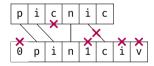
ED(0PIN1CIV, OPINION) = 5

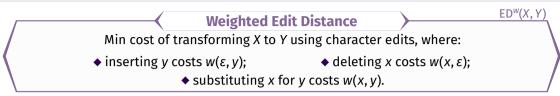
ED(0PIN1CIV, PICNIC) = 5



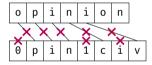
w(0,0) := 1 w(1,1) := 1 w(C,0) := 1 w(*,*) := 2 $w(*,\varepsilon) := 1$ $w(\varepsilon,*) := 10$

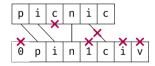






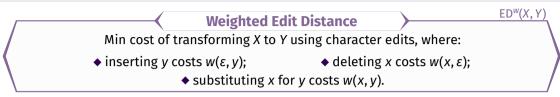
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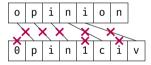


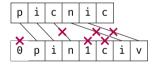
ED^w(0PIN1CIV, OPINION) = 6

ED^w(0PIN1CIV, PICNIC) ≤ 14



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ED^w(0PIN1CIV, OPINION) = 6

ED^w(0PIN1CIV, PICNIC) = 8

Computing (Weighted) Edit Distance

How fast can we compute the (weighted) edit distance of two strings? $\rightsquigarrow O(n^2)$ (recalled this earlier!)

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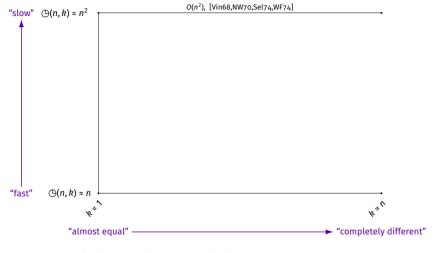
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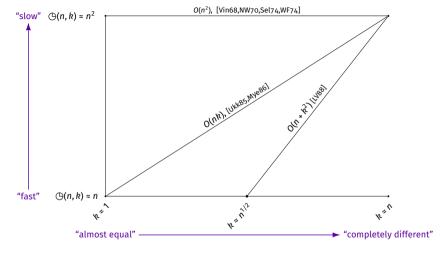
For Pattern Matching, enough to compute (weighted) edit distance if small (at most *k*) → Bounded (Weighted) Edit Distance

Algorithms for (Bounded) Edit Distance



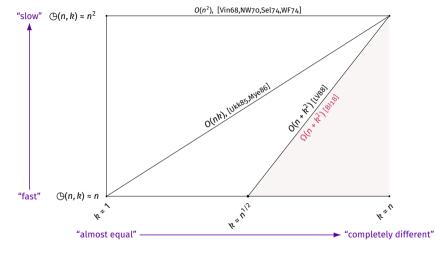
Existing algorithms for Edit Distance ED(X, Y), where $|X|, |Y| \le n$

Algorithms for (Bounded) Edit Distance



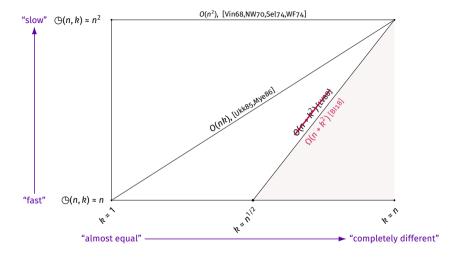
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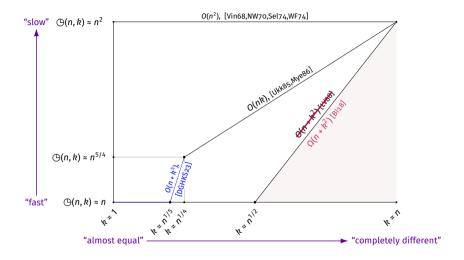


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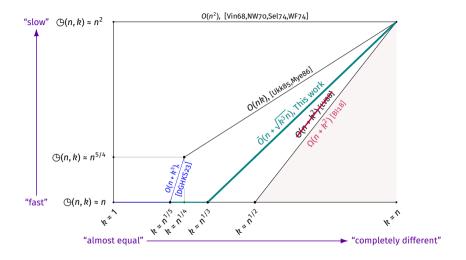
What about Weighted Edit Distance?



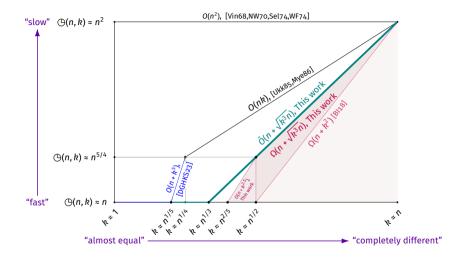
Existing algorithms for Weighted Edit Distance $ED^{w}(X, Y)$, where $|X|, |Y| \le n$



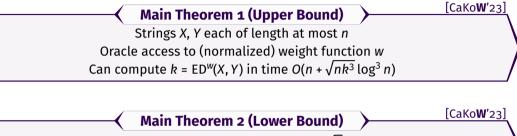
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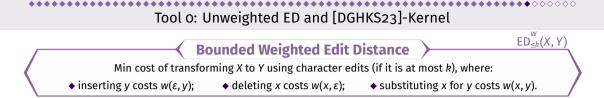


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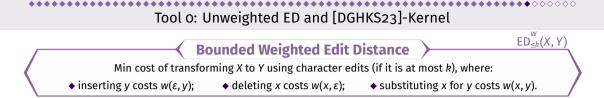


Assuming the APSP Hypothesis and for $\sqrt{n} \le k \le n$, Main Theorem 1 is tight (up to $n^{o(1)}$ -factors)

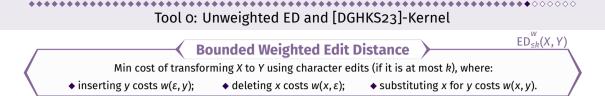
Tool O: Unweighted ED and [DGHKS23]-Kernel Bounded Weighted Edit Distance Min cost of transforming X to Y using character edits (if it is at most k), where: \Rightarrow inserting y costs $w(\varepsilon, y)$; \Rightarrow deleting x costs $w(x, \varepsilon)$; \Rightarrow substituting x for y costs w(x, y).



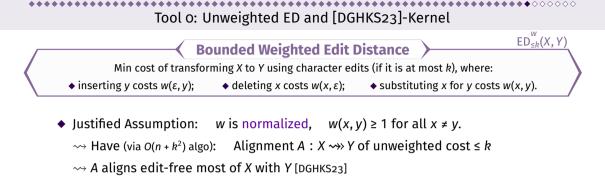
• Justified Assumption: w is normalized, $w(x, y) \ge 1$ for all $x \ne y$.



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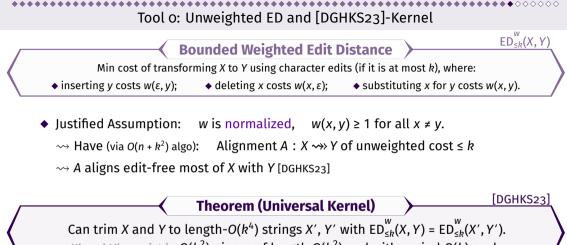


Justified Assumption: w is normalized, w(x, y) ≥ 1 for all x ≠ y.
 → Have (via O(n + k²) algo): Alignment A : X → Y of unweighted cost ≤ k
 → A aligns edit-free most of X with Y [DGHKS23]





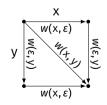
Philip Wellnitz From Strings to Seaweeds: Modern Tools for Classical Problems 66-5

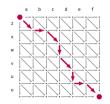


X' and Y' consist in $O(k^2)$ pieces of length $O(k^2)$ and with period O(k) each

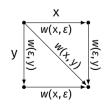
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- Idea: Use (directed) alignment graph AG of X and Y (seaweeds don't work here)
- Idea²: Trim AG to O(k) diagonals $\rightarrow ED_{\leq k}^{W}(X, Y)$ is distance $(0, 0) \rightarrow (|X|, |Y|)$
- Idea³: Split AG according to structure of X and Y
 Compute all b-to-b dist [Kleino5] + stitch together results ((min, +)-product [SMAWK87])

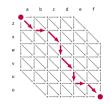
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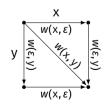


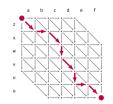
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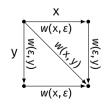


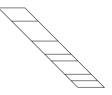
- X and Y consist in $O(k^2)$ pieces of length $O(k^2)$ and with period O(k) each
- Idea: Use (directed) alignment graph AG of X and Y (seaweeds don't work here)
 → ED^W(X, Y) is distance (0, 0) → (|X|, |Y|)
- Idea²: Trim AG to O(k) diagonals → ED^W_{≤k}(X, Y) is distance (0, 0) → (|X|, |Y|)
 → AG has O(k⁵) vertices → Dijkstra yields Õ(k⁵) algo
- Idea³: Split AG according to structure of X and Y
 Compute all b-to-b dist [Kleino5] + stitch together results ((min, +)-product [SMAWK87])





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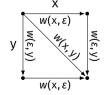


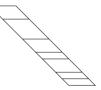
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- Idea³: Split AG according to structure of X and Y
 Compute all b-to-b dist [Kleino5] + stitch together results ((min, +)-product [SMAWK87])
 for periodic pieces: fast exponentiation;

$$\rightsquigarrow k^2 \cdot \tilde{O}(k^2)$$
 for periodic pieces + $k^2 \cdot \tilde{O}(k^2)$ for stitching





Tool 2: Divide and Conquer

- X and Y consist in $O(k^2)$ periodic pieces of length $O(k^2)$ and with period O(k) each
- Idea: Use AG, trimmed to O(k) diags $\rightsquigarrow ED_{\leq k}^{W}(X, Y)$ is distance $(0, 0) \rightsquigarrow (|X|, |Y|)$
- Idea³: Compute all b-to-b dist for periodic pieces [Kleino5] + fast exponentiation; stitch together results using min-plus product [SMAWK87]
- Idea⁴: Use Divide-and-Conquer to reduce number of periodic pieces to O(k) $\rightarrow k \cdot \tilde{O}(k^2)$ for periodic pieces (and padding) + $k \cdot \tilde{O}(k^2)$ for stitching

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Tool 3: Compressibility instead of Periodicity

- X and Y consist in O(k) pieces of length $O(k^2)$ each
- Idea: Use AG, trimmed to O(k) diags $\rightsquigarrow ED_{\leq k}^{W}(X, Y)$ is distance $(0, 0) \rightsquigarrow (|X|, |Y|)$
- Idea³: Compute all b-to-b dist for periodic pieces [Kleino5] + fast exponentiation; stitch together results using min-plus product [SMAWK87]
- Idea⁴: Use Divide-and-Conquer to reduce number of periodic pieces to O(k)
- Idea⁵: Use tailor-made compressibility measure instead of periodicity + (w)ED algorithms for compressed strings $\sim \tilde{O}(n + \sqrt{k^3 n})$ time in total

◆◇◇ Main Results

Main Theorem 1 (Upper Bound) 🗸

Strings X, Y each of length at most n Oracle access to (normalized) weight function w Can compute $k = ED^{w}(X, Y)$ in time $O(n + \sqrt{nk^3} \log^3 n)$

Main Theorem 2 (Lower Bound)

Assuming the APSP Hypothesis and for $\sqrt{n} \le k \le n$, Main Theorem 1 is tight (up to $n^{o(1)}$ -factors)

•• Pattern Matching with Weighted Edits—Next Steps

- Have good algorithm for Bounded Weighted Edit Distance
- ◆ (ongoing work) Use structural result and directed alignment graph for Õ(|T| + k⁴|T|/|P|) for PM w/ WE
- Seaweeds don't help, but fast multiplication of Monge Matrices still does (at a cost of an extra O(k))
- Still need to solve a lot of extra technical challenges (actually obtaining an analogue of [LV89] for verifying starting positions; need to generalize DPM; ...)

• Summary and Future Directions

Main Results of today

- Structural characterization of strings and PM w/ Edits
- Main tools required for $\tilde{O}(|T| + k^{3.5}|T|/|P|)$ algo for PM w/ E
- Some ideas of how to generalize to the weighted setting

Big Open Questions

- PM w/ Edits in $o(|T| + k^3 \cdot |T|/|P|)$
- PM w/ Weighted Edits in $o(|T| + k^4 \cdot |T|/|P|)$
- Better algorithms known for small integer weights [GK24⁺], might give better PM algos in that setting



