Pattern matching with don't cares and few errors

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Matching with don't cares & few erros Introduction

Pattern matching

$$\begin{split} &\Sigma: \text{ alphabet } \\ &t = t_1 t_2 \dots t_n \in \Sigma^n: \text{ text } \\ &p = p_1 p_2 \dots p_m \in \Sigma^m: \text{ pattern } \end{split}$$

Exact pattern matching

Given: a pattern $p \in \Sigma^n$, a text $t \in \Sigma^m$.

Goal: find all the places that *p* matches *t*.

• O(n) time [Boyer & Moore 1977; Knuth, Morris & Pratt 1977]



Matching with don't cares & few erros Introduction

Pattern matching

Pattern matching with don't cares

Given: a pattern $p \in \Sigma^n$, a text $t \in \Sigma^m$, where p, t may contain ' ϕ 's. **Goal:** find all the places that p matches t.

• $\Theta(n \log m)$ [Cole& Hariharan 2002; Clifford $\times 2$ & 2007].



Matching with don't cares & few erros Introduction

Pattern matching (k-mismatch)

k-mismatch WITHOUT don't cares

Given: a pattern $p \in \Sigma^n$, a text $t \in \Sigma^m$ and an integer $k \ge 0$. **Goal:** find all the places that p matches t with $\le k$ mismatches.

- $\Theta(n\sqrt{m\log m})$ time [Abrahamson 1987; Kosaraju 1987].
- $\Theta(n\sqrt{k \log k})$ [Amir, Lewenstein & Porat 2004].



Pattern matching (k-mismatch)

k-mismatch with don't cares

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Goal: find all the places that p matches t with $\leq k$ mismatches.

- $O(nm^{1/3}k^{1/3}\log^{2/3}m)$ time [Clifford & Porat 2007].
 - ' ϕ 's: permitted in either the pattern or text, but not both.
- Extend Kosaraju & Abrahamson's work with little extra work: $\Theta(n\sqrt{m\log m})$ time.
- No other previous efficient algorithm for this problem.



Contribution of this paper

k-mismatch with don't cares

Given: a pattern $p \in \Sigma^n$, a text $t \in \Sigma^m$ and an integer $k \ge 0$, where p, t may contain ' ϕ 's. **Goal:** find all the places that p matches t with < k mismatches.

This paper:

- Two randomized $\tilde{\Theta}(nk)$ time algorithms.
 - A randomized $\Theta(nk \log m \log n)$ time algorithm.
 - Further improved $\rightarrow \Theta(n(k + \log m \log k) \log n)$ time.
- A deterministic $\Theta(nk^2 \log^2 m)$ time algorithm (group testing).
- $\Theta(nk \operatorname{polylog} m)$ time using k-selectors.



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k-mismatch & Hamming distance

HD(i): the Hamming distance between p and t[i,...,i+m-1].
 * φ matches any symbol in Σ.

$$HD_k(i) = \begin{cases} HD(i) & \text{if } HD(i) \le k \\ \bot & \text{otherwise.} \end{cases}$$

- HD_k(i) ≠⊥
 ⇒ There is a k-mismatch between p and t at alignment i.
- For example, $HD_2(1) = 2$, $HD_2(5) = 1$

i: 1 2 3 4 5 6 7 8 9 10 *t*: A A C φ G A φ T T G *p*: A φ G G A

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 - *i*: 1 2 3 4 5 6 7 8 9 10
 - $t: A A C \phi G A \phi T T G$
 - p: A ϕ G G A



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i: 1 2 3 4 5 6 7 8 9 10
t: A A C
$$\phi$$
 G A ϕ T T G

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Transformation from matching to coding...

• The key observation by [Clifford×2 2007]:

$$\sum_{j=1}^{m} (p_j - t_{i+j-1})^2 = \sum_{j=1}^{m} (p_j^2 - 2p_j t_{i+j-1} + t_{i+j-1}^2).$$

$$\sum_{j=1}^m p_j' t_{i+j-1}' (p_j - t_{i+j-1})^2.$$

where

$$p_j' = \left\{ egin{array}{ccc} 0 & ext{if } p_j = `\phi' & t_i' = \left\{ egin{array}{ccc} 0 & ext{if } t_i = `\phi' & 1 & 0 & ext{therwise.} \end{array}
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where

$$p'_{j} = \begin{cases} 0 & \text{if } p_{j} = `\phi' \\ 1 & \text{otherwise;} \end{cases} \quad t'_{i} = \begin{cases} 0 & \text{if } t_{i} = `\phi' \\ 1 & \text{otherwise.} \end{cases}$$

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Matching with don't cares & few erros Preliminaries

The cross-correlation

$$(t \otimes p)[i] := \sum_{j=1}^{m} p_j t_{i+j-1}, \quad 0 \le i \le n-m+1.$$

- The above cross-correlation (convolution) can be calculated in Θ(n log m) time.
 - Fast Fourier Transform (FFT).



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Starting by 1-mismatch...

The 1-mismatch problem

To determine where p and t have EXACTLY ONE mismatch.

Masking out a number of positions in p with \u03c6's at random (each of prob. (k - 1)/k).

- The resulting pattern: *subpattern*.
- * It is likely that exactly one mismatch can be found.



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 - The resulting pattern: subpattern.
 - \star It is likely that exactly one mismatch can be found.



Algorithm 1-mismatch $\Theta(n \log m)$

Input: Pattern *p*, text *t*.

Output: B[i]: mismatch location in t for each alignment where HD(i) = 1.

- Compute $A_0[i] = \sum_j (p_j t_{i+j-1})^2 p'_j t'_{i+j-1};$
- 2 Compute $A_1[i] = \sum_j (i+j-1)(p_j t_{i+j-1})^2 p'_j t'_{i+j-1};$
- (a) for each $i \in \{0, 1, \dots, n\}$ do
 - a. if $A_0[i] \neq 0$ then
 - $B[i] \leftarrow A_1[i]/A_0[i];$

b. else

• $B[i] \leftarrow No_Mismatch;$

• for each $i \in \{0, 1, \dots, n\}$ s.t. $B[i] \neq \texttt{No_Mismatch}$ do

• if
$$(p[B[i] - i + 1] - t[B[i]])^2 \neq A_0[i]$$
 then
 $B[i] \leftarrow \text{More_Than_1_Mismatch};$



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- Running 1-mismatch for $\Theta(k \log n)$ times.
- Each time we get the location of a mismatch (with the subpattern) if one occurs.
- Similar to the concept of solving the Coupon Collector's problem.
- Yet, the last coupon is always the most difficult to get!



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Correctness?
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Correctness of the first randomized algorithm

• For $HD(i) \leq k$:

- One fixed single mismatch is found in one iteration with prob. $\geq \left(\frac{k-1}{k}\right)^{k-1} \cdot \frac{1}{k} > \frac{1}{e^k}.$
 - This one is not found after Θ(k log n) iterations with prob.
 ≤ (1 − 1/ek)^{Θ(k log n)} ≤ n^{-c} for some constant c.
- Applying union bound to derive the overall error prob.
- For HD(i) > k:
 - Extra checking stage by computing

$$C[i] = \sum_{j=1}^m (i+j-1)(p_j-t_{i+j-1})^2 p_j' t_{i+j-1}', ext{ for each } i.$$

"Correct" C[i] for each distinct 1-mismatch using the found A₀[i
 Keep track of found mismatches (using a binary search tree).



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The clever ideas

Instead of finding ALL mismatches at once, find HALF of them! Efficient & with high prob. of success.

- Save the effort in dealing with previous found mismatches.
 Correct the sums in A₀ and A₁ prior to their being used.
- The left mismatches to be found → increase the sampling rate!



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The clever ideas

- Instead of finding ALL mismatches at once, find HALF of them!
 Efficient & with high prob. of success.
- Save the effort in dealing with previous found mismatches.
 - Correct the sums in A_0 and A_1 prior to their being used.
- $\bullet\,$ The left mismatches to be found \rightarrow increase the sampling rate!



A faster recursive randomized algorithm

Input: Pattern p, text t, and an integer $k \ge 0$. **Output:** Array $O[i] = HD_k(p, t[i, ..., i + m - 1]).$

- 1 Initialize *E* and set $k_0 \leftarrow k$;
- ② for $s \leftarrow 0$ to $\lfloor \log k \rfloor$ do
 - a. for times $\leftarrow 1$ to $\Theta(k_s + \log n)$ do /* Sample and Match stage */

I. Sample subpattern p^* with sample rate $1/k_s$;

II. self-correcting-1-mismatch(p^*, t, E);

- b. Update *E* according to the mismatches found in the iterations; c. $k_{s+1} \leftarrow k_s/2$;
- $L[i] \leftarrow$ total number of distinct mismatches found at alignment *i*;
- Check at each position i in t that all mismatches were found;
- **③** $O[i] \leftarrow L[i]$, if all mismatches were found, otherwise $O[i] \leftarrow \bot$.



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A faster recursive randomized algorithm $\Theta(n(k + \log m \log k) \log n)$

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Thanks for your attention.



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Matching with don't cares & few erros

Appendix



Joseph C.-C. Lin (Academia Sinica, TW) Matching with don't cares & few erros

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E

Half different mismatches can be found whp in each stage

•
$$k_s = k/2^s$$

Lemma 4.5

After $\Theta(k_s + \log n)$ iterations of the sample and match stage, the locations and values of $\leq k_s/2$ different mismatches will remain to be found whp.

- X_i: 1 if a mismatch at the *i*th stage is found, 0 otherwise.
- $E(\sum_{i=1}^{\omega} X_i) \ge \omega/e$, for $\omega = \Theta(k_s + \log n)$.
 - $\star\,$ Sum of random variables \rightarrow using Chernoff bounds.
 - $\geq \omega/2e$ mismatches can be found whp.
- Found mismatches are entirely contained in any size-(k_s/2) set with prob. ≤ 2^{ω/2e} (k_s/2).



The number of times to handling previously found mismatches

Lemma 4.7

If the sample and match stage is run $r = \Theta(k_s + \log n)$ times, # times a previously discovered mismatch is found is $O(nk(k_s + \log n)/k_s)$ whp.

• $X_{i,j}$: whether p_i was replaced with ϕ in iteration j.

•
$$\Pr(X_{i,j} = 1) = 1/k_s$$
.

- a_i : # mismatches previously found at p_i in all alignments.
- $X = \sum_{i \in [m], j \in [r]} a_i X_{i,j}$ is $O(nk(k_s + \log n)/k_s)$ whp.
 - Using Chernoff bound once again (though a different formula).
- * The overall time complexity can be derived easily.



The Chernoff-Hoeffding bounds

Theorem 4.4

Assume that X_1, \ldots, X_m are i.i.d. random variables, $X_i \in \{0, 1\}$. Let $\mu_i = E(X_i)$. Then

$$Pr\left(\sum_{i=1}^m X_i \leq (1-\delta)m\mu\right) < e^{-m\mu\delta^2/2}.$$

Theorem 4.6

Let X_1, \ldots, X_m be discrete, independent random variables s.t. $E(X_i) = 0$ and $|X_i| \le 1$ for all *i*. Let $X = \sum_{i=1}^m X_i$. Then

$$Pr(X \ge \lambda \sqrt{\operatorname{Var}(X)}) \le e^{-\lambda^2/4}.$$

In the proof of Theorem 4.9

Let us concentrate on the *i*th stage of the recursion. At this stage, we need to solve the $k/2^i$ -mismatch problem, by running the self-correcting 1-mismatch algorithm $\Theta(2^i + \log n)$ times. ...

- Shouldn't it be $\Theta(k/2^i + \log n)$ times?
 - Though it doesn't affect the result.

