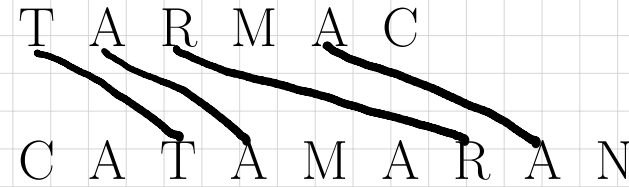


Dynamic Programming II

Recall the maximum common subsequence problem from last day:



More sophisticated: count # changes

e.g., You : Pythagorus

Google : Pythagoras ?
1 change

You : recurrence

Google : recurrence ?
2 changes

A change is:

- add a letter } gap
- delete a letter }
- replace a letter — mismatch

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The problem comes up in bioinformatics for DNA strings.

DNA is a sequence of chromosomes, i.e., a string over the alphabet A, C, T, G.

This is called edit distance.

Two string can be aligned in different ways:

e.g. A A C A T
A A A A G

3 changes
(2 gaps, 1 mismatch)

e.g. A A C A T
A A A A G

2 changes
(2 mismatches)

Problem: Given 2 strings $x_1..x_m$ and $y_1..y_n$, compute their edit distance.
I.e., find the alignment that gives the minimum number of changes.

Dynamic Programming Algorithm

Subproblem: $M(i, j)$ = minimum number of changes to match $x_1..x_{i-1}x_i$ and $y_1..y_{j-1}y_j$.

choices:- match x_i to y_i , pay replacement cost if they differ

- match x_i to blank (delete x_i)

- match y_j to blank (add y_j)

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$$M(i, j) = \min \begin{cases} M(i-1, j-1) & \text{if } x_i = y_j \\ r + M(i-1, j-1) & \text{if } x_i \neq y_j \\ d + M(i-1, j) & \text{match } x_i \text{ to blank} \\ a + M(i, j-1) & \text{match } y_j \text{ to blank} \end{cases} \quad \text{where:}$$

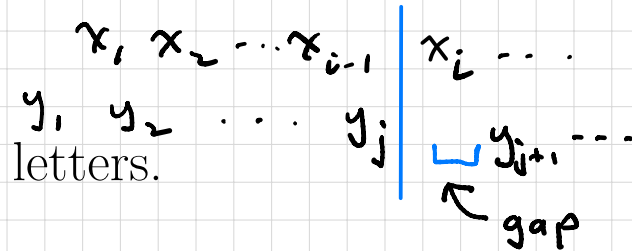
r = replacement cost
 d = delete cost
 a = add cost

So far, we used $r = d = a = 1$ (i.e., count # changes).

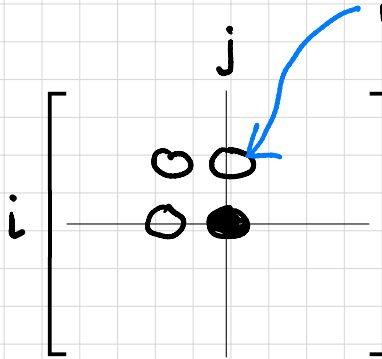
More sophisticated: $r(x_i, y_j)$ - replacement cost depends on the letters.

e.g., $r(a, s) = 1$ because these keys are close on typewriter

$r(a, c) = 2$... not too close



In what order do we solve subproblems? Same as last day.



matrix $m \times n$

$M[0..m, 0..n]$

for $i = 0..m$: $M(i, 0) = id$ - delete i letters

for $j = 0..n$: $M(0, j) = ja$ - add j letters

for $i = 1..m$

for $j = 1..n$

$M(i, j) = \dots$

}

fill matrix in order $\left[\begin{smallmatrix} \Rightarrow \\ \Rightarrow \\ \Rightarrow \end{smallmatrix} \right]$
(or could do columns first)

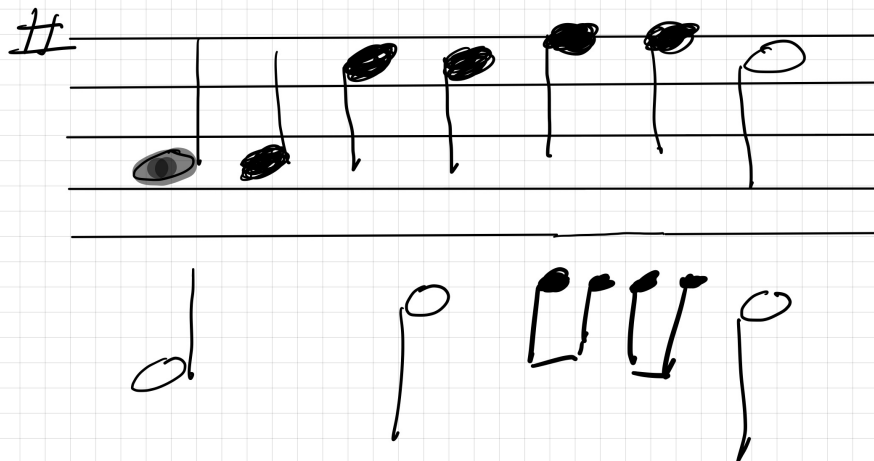
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Analysis: $O(nm)$ time and $O(nm)$ space

<https://powcoder.com> (not solving problems, constant time each)

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A different application: music pattern matching

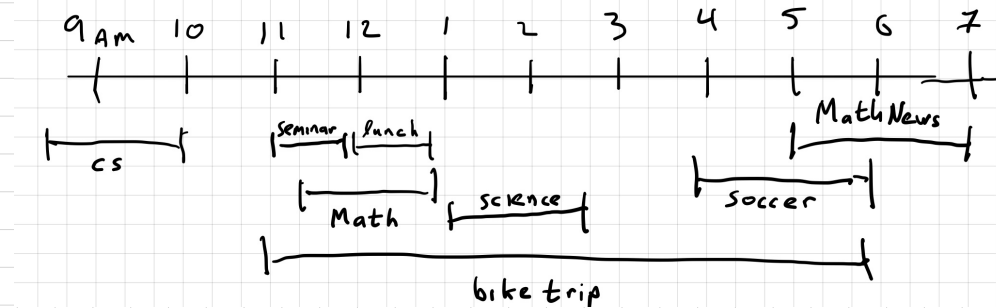


← match this

← to this

use replacement rules that allow $d \rightarrow \text{half note}$

Recall Interval Scheduling aka Activity Selection: Given a set of intervals I , find a maximum size subset of disjoint intervals:



Weighted Interval Scheduling

Weighted Interval Scheduling: Given I and weight $w(i)$ for each $i \in I$, find set $S \subseteq I$ such that no two intervals overlap and maximize $\sum_{i \in S} w(i)$.

e.g., you have preferences for certain activities

A more general problem:

- I is a set of element ("items")
- $w(i)$ = weight of item i
- some pairs (i, j) conflict

Find a maximum weight subset $S \subset I$ with no conflicting pairs.

Can be modeled as a graph: vertex = item edge = conflict

Problem is Max Weight Independent Set and we will see later that it is NP-complete.

A general approach to finding max weight independent set.
Consider one item i . Either we choose it or not.

$$\text{OPT}(I) = \max\{\text{OPT}(I - \{i\}), w(i) + \text{OPT}(I')\} \quad \text{where } I' = \text{intervals disjoint from } i$$

In general this recursive solution does not give polynomial time.

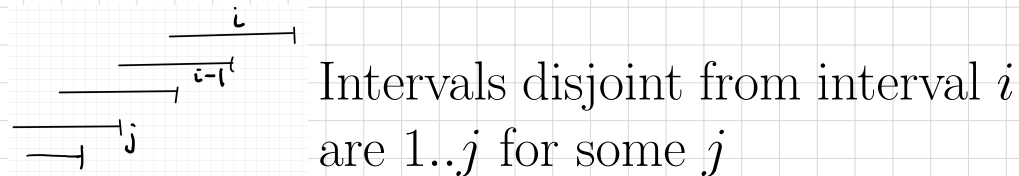
$$T(n) = 2T(n-1) + O(1) \implies T(n) \in \Theta(2^n)$$

Essentially, we may end up solving subproblems for each of the 2^n subsets of I .

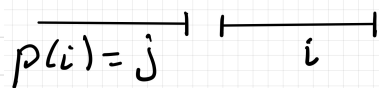
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When I = set of intervals, we can do better with dynamic programming.

Order intervals $1..n$ by right endpoint
something nice happens



For each i , let $p(i)$ = largest index $j < i$
s.t. interval j is disjoint from interval i .



Now we can solve subproblems.

Let $M(i)$ = max weight subset of
intervals $1..i$

$$M(i) = \max\{M(i-1), w(i) + M(p(i))\}$$

A Dynamic Programming algorithm – computes the actual set, not just weight

Sort intervals $1..n$ by right endpoint.

$M(0) := 0; \quad S(0) := \emptyset$ *S stores the set*

for i **from** 1 **to** n **do**

Compute $p(i)$ $O(n)$ { $p(i) := i - 1$

while $p(i) \neq 0$ **and** intervals i and $p(i)$ overlap **do**

$p(i) := p(i) - 1$

od

if $M(i-1) \geq w(i) + M(p(i))$ **then**

$M(i) := M(i-1); \quad S(i) := S(i-1)$

else

$M(i) := w(i) + M(p(i)); \quad S(i) := \{i\} \cup S(p(i))$

fi

od

Final answer: weight $M(n)$, set $S(n)$

Time: n subproblems, each $O(n)$

so total of $O(n^2) + O(n \log n)$ to sort.

Space: $O(n^2)$ - storing n sets, each $O(n)$

Next:

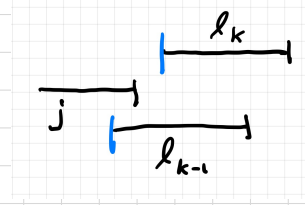
1. computing all $p(i)$ values before-hand to save time
2. computing S by backtracking to save space

How to compute $p(i)$: We use sorted order $1..n$ by right endpoint
AND sorted order $\ell_1..\ell_n$ by left endpoint

```

 $j := n$ 
for  $k$  from  $n$  downto 1 do
  while  $\ell_k$  overlaps  $j$  do
     $j := j - 1$ 
  od
   $p(\ell_k) := j$ 
od

```



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Run time $\Theta(n)$ after sorting

<https://powcoder.com>

Final algorithm:

```

Sort intervals  $1..n$  by right endpoint.
Sort intervals by left endpoint.
Compute  $p(i)$  for all  $i$ .
 $M(0) := 0$ 
for  $i$  from 1 to  $n$  do
   $M(i) := \max\{M(i-1), w(i) + M(p(i))\}$ 
od

```

Runtime: $\underbrace{O(n \log n)}_{\text{sort}} + \underbrace{O(n)}_{p(*)} + O(n \cdot c)$
 ↗ time per subproblem
 ↘ # subproblems

Backtracking to compute S : Use recursive routine to S -OPT

```

S-OPT( $i$ )
  if  $i = 0$  then
    return  $\emptyset$ 
  elif  $M(i - 1) \geq w(i) + M(p(i))$  then
    return S-OPT( $i - 1$ )
  else
    return  $\{i\} \cup \text{S-OPT}(p(i))$ 
  fi

```

The set we want is $S\text{-OPT}(n)$.

Time: $O(n)$

Space: $O(n)$

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Summary

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- A general idea to find an optimal subset is to solve subproblems where one element is in or out

Exponential in general; can sometimes be efficient

- Key ideas of dynamic programming:
 - Identify subproblems (not too many) together with
 - an order of solving them such that each subproblem can be solved by combining a few previously solved subproblems.