Applied Algorithms CSCI-B505 / INFO-I500

Lecture 10.

Dynamic Programming - I

- Dynamic Programming
 - Fibonacci
 - Binomial Coefficients
 - Edit distance
 - Some Variants of Edit Distance
 - Longest Common Subsequence

Dynamic Programming

- Find the minimum or maximum of a combinatorial challenge (combinatorial opt.)
- Exhaustive search guarantees the optimum, but very expensive
- Greedy approach (!) is more reasonable, but no guarantees (in general)
- Dynamic programming aims to compute the optimum with a good complexity by storing the results of some prior computations for the sake of some others later.
- DP is particularly useful when there is a reductive solution but with significant overlaps between the recursive steps.

Bad Recursions

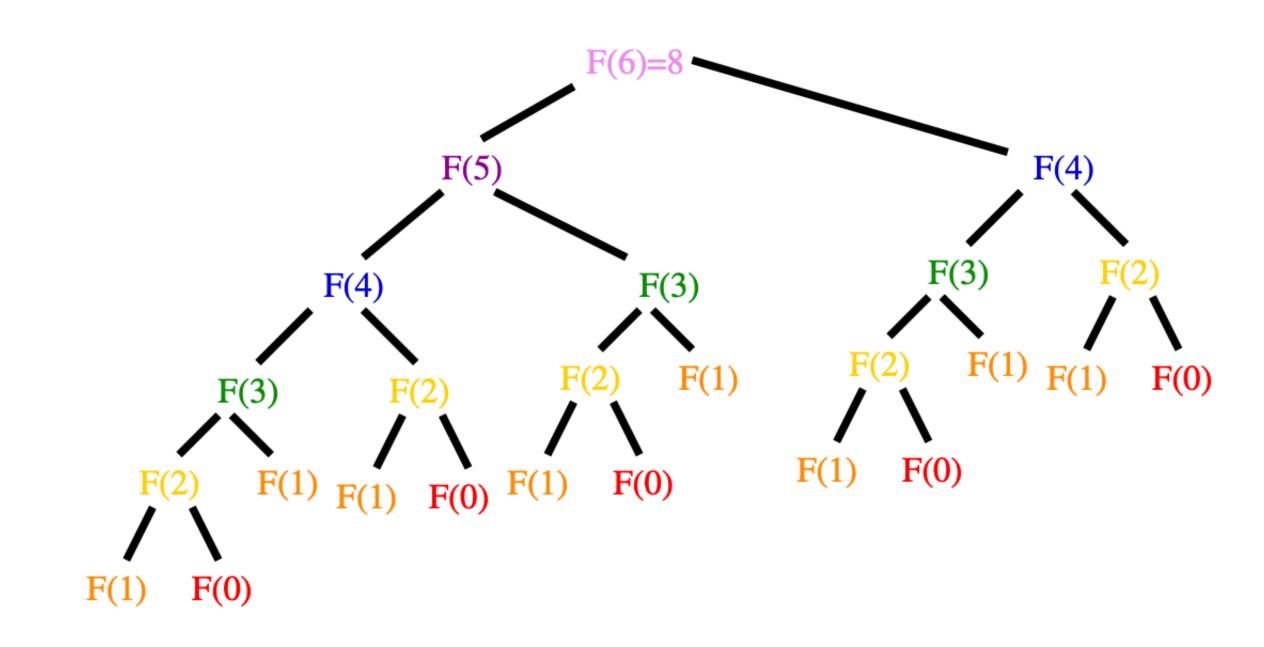
Same computation is repeated many times in the recursion tree!

Example: Fibonacci with recursion ...

```
long fib_r(int n) {
    if (n == 0) {
        return(0);
    }

if (n == 1) {
        return(1);
    }

return(fib_r(n-1) + fib_r(n-2));
}
```



- Time complexity is **exponential** $\approx 1.6^n$. Why?
- $F(n) > 1.6^n$, for large n, and since we will sum up to this by reaching that much of leaves.

Bad Recursions

A better recursion with caching!

```
F(4)
                F(5)
                               F(3)
       F(4)
F(3)
            F(2)
   F(1)
```

```
long fib_c(int n) {
   if (f[n] == UNKNOWN) {
       f[n] = fib_c(n-1) + fib_c(n-2);
   return(f[n]);
long fib_c_driver(int n) {
   int i; /* counter */
   f[0] = 0;
   f[1] = 1;
   for (i = 2; i <= n; i++) {
       f[i] = UNKNOWN;
   return(fib_c(n));
```

• Allocating O(n) space returns the result in O(n) time.

Simple DP for Fibonacci

• Allocating O(1) space returns the result in O(n) time.

Binomial Coefficients

$$\binom{n}{k} = \frac{n!}{(n-k)! \cdot k!}$$

- It can be directly calculated. However, overflows are possible even with small n, k values.
- Recursive calculation avoids such overflows

```
long binomial_coefficient(int n, int k) {
   int i, j;
             /* counters */
   long bc[MAXN+1][MAXN+1]; /* binomial coefficient table */
   for (i = 0; i <= n; i++) {
       bc[i][0] = 1;
   for (j = 0; j <= n; j++) {
      bc[j][j] = 1;
   for (i = 2; i <= n; i++) {
      for (j = 1; j < i; j++) {
          bc[i][j] = bc[i-1][j-1] + bc[i-1][j];
   return(bc[n][k]);
```

$$\binom{n}{k} = \binom{n-1}{k-1} + \binom{n-1}{k}$$

Approximate String Matching

Given two strings s1 and s2, in how many steps can we alter s1 to become s2?

The alterations we are allowed to do are:

- Substitution: Change a specific symbol of s1 to match a symbol of s2,
 - e.g., s1: shot s2:spot
- Insertion: Insert a new symbol into s1 to match a corresponding symbol on s2
 - e.g., s1: ago s2:agog
- Deletion: Delete symbol from s1
 - e.g., s1: hour s2:our

We will assume each operation has the equal cost of 1.

Approximate String Matching by Recursion

Let edit distance between $P_1P_2...P_i$ and $T_1T_2...T_j$ be D[i,j].

There are three possibilities:

1.
$$D[i,j] = D[i-1,j-1] + (1|0)$$

- 2. D[i,j] = D[i,j-1] + 1 Extra symbol on T, $indel(T_j)$, either insertion on P or a deletion from T, thus, named **indel**
- 3. D[i,j] = D[i-1,j] + 1 Extra symbol on P, $indel(P_i)$ either insertion on T or a deletion from P, $indel(P_i)$

Approximate String Matching by Recursion

```
int string_compare_r(char *s, char *t, int i, int j) {
   int k;
           /* counter */
   int opt[3];  /* cost of the three options */
   int lowest_cost; /* lowest cost */
   if (i == 0) { /* indel is the cost of an insertion or deletion */
       return(j * indel(' '));
   if (j == 0) {
       return(i * indel(' '));
                                                                        D[i,j] = D[i-1,j-1] + (1|0)
                    /* match is the cost of a match/substitution */
               = string_compare_r(s,t,i-1,j-1) + match(s[i],t[j]);
   opt [MATCH]
                                                                          \rightarrow D[i, j] = D[i, j-1] + 1
   opt[INSERT] = string_compare_r(s,t,i,j-1) + indel(t[j]); ______
   opt[DELETE] = string_compare_r(s,t,i-1,j) + indel(s[i]);_
   lowest_cost = opt[MATCH];
                                                                           D[i, j] = D[i - 1, j] + 1
   for (k = INSERT; k <= DELETE; k++) {
       if (opt[k] < lowest_cost) {</pre>
           lowest_cost = opt[k];
```

return(lowest_cost);

Recursive Program with exponential time ($\approx 3^n$) since at each recursion, three children are born.

D[5][6]

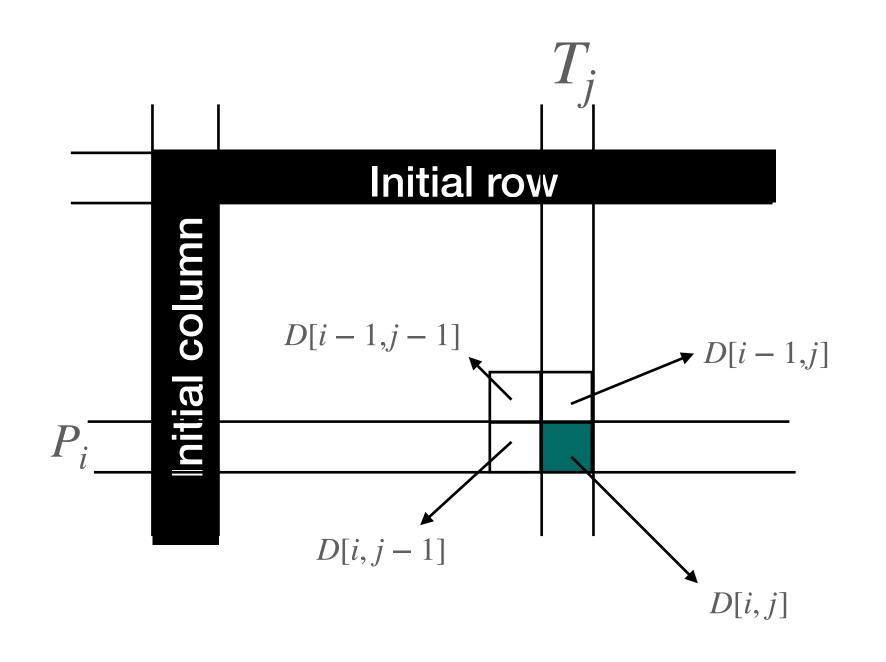
D[4][5] D[5][5] D[4][6]

D[3][4] D[4][4] D[3][5] D[4][4] D[5][4] D[4][5] D[3][5] D[4][5] D[3][6]

BAD RECURSION!?

Approximate String Matching by Dynamic Programming

	T		у	O	u	_	S	h	O	u	1	d
P	pos	0	1	2	3	4	5	6	7	8	9	10
:		<u>0</u>	1	2	3	4	5	6	7	8	9	10
t:	1	<u>1</u>	1	2	3	4	5	6	7	8	9	10
h:	2	2	<u>2</u>	2	3	4	5	5	6	7	8	9
o:	3	3	3	<u>2</u>	3	4	5	6	5	6	7	8
u:	4	4	4	3	<u>2</u>	3	4	5	6	5	6	7
-:	5	5	5	4	3	<u>2</u>	3	4	5	6	6	7
s:	6	6	6	5	4	3	<u>2</u>	3	4	5	6	7
h:	7	7	7	6	5	4	3	<u>2</u>	<u>3</u>	4	5	6
a:	8	8	8	7	6	5	4	3	3	<u>4</u>	5	6
l:	9	9	9	8	7	6	5	4	4	4	<u>4</u>	5
t:	10	10	10	9	8	7	6	5	5	5	5	<u>5</u>
		1 1										



- Initial row and column values are fixed.
- We just fill the matrix with a row-major or column major traversal.

Approximate String Matching by Dynamic Programming

```
int string_compare(char *s, char *t, cell m[MAXLEN+1][MAXLEN+1]) {
    int i, j, k;
                 /* counters */
    int opt[3];  /* cost of the three options */
    for (i = 0; i <= MAXLEN; i++) {
                                                             #define MATCH
                                                                                       /* enumerated type symbol for match */
        row_init(i, m);
                                                                                        /* enumerated type symbol for insert */
                                                             #define INSERT
        column_init(i, m);
                                                                                        /* enumerated type symbol for delete */
                                                             #define DELETE
   for (i = 1; i < strlen(s); i++) {
       for (j = 1; j < strlen(t); j++) {
           opt[MATCH] = m[i-1][j-1].cost + match(s[i], t[j]);
           opt[INSERT] = m[i][j-1].cost + indel(t[j]);
           opt[DELETE] = m[i-1][j].cost + indel(s[i]);
           m[i][j].cost = opt[MATCH];
           m[i][j].parent = MATCH;
           for (k = INSERT; k <= DELETE; k++) {
               if (opt[k] < m[i][j].cost) {
                   m[i][j].cost = opt[k];
                   m[i][j].parent = k;
                                                                                                  The edit distance between P and T
```

DP solution of edit distance works in $O(n \cdot m)$ -time and -space.

Approximate String Matching by Dynamic Programming

```
void reconstruct_path(char *s, char *t, int i, int j,
                               cell m[MAXLEN+1][MAXLEN+1]) {
    if (m[i][j].parent == -1) {
        return;
    if (m[i][j].parent == MATCH) {
        reconstruct_path(s, t, i-1, j-1, m);
        match_out(s, t, i, j);
        return;
    if (m[i][j].parent == INSERT) {
        reconstruct_path(s, t, i, j-1, m);
        insert_out(t, j);
        return;
    if (m[i][j].parent == DELETE) {
        reconstruct_path(s, t, i-1, j, m);
        delete_out(s, i);
        return;
```

• It is also possible to gather it from the cost matrix as well.

The parent information can be

separately maintained to

reconstruct the path.

```
the hount of the land of the l
```

Some Variants of the Edit Distance

- Edit distance has many different variants to solve different problems.
- All fill the DP matrix with some slight differences that make a significant effect.
 - Different cost functions
 - The position of the final result on the matrix
 - Different initialization of the matrix
 - Different traceback actions

Some Variants of the Edit Distance

Substring Matching: On a long text T we aim to spot P, wherever it matches best.

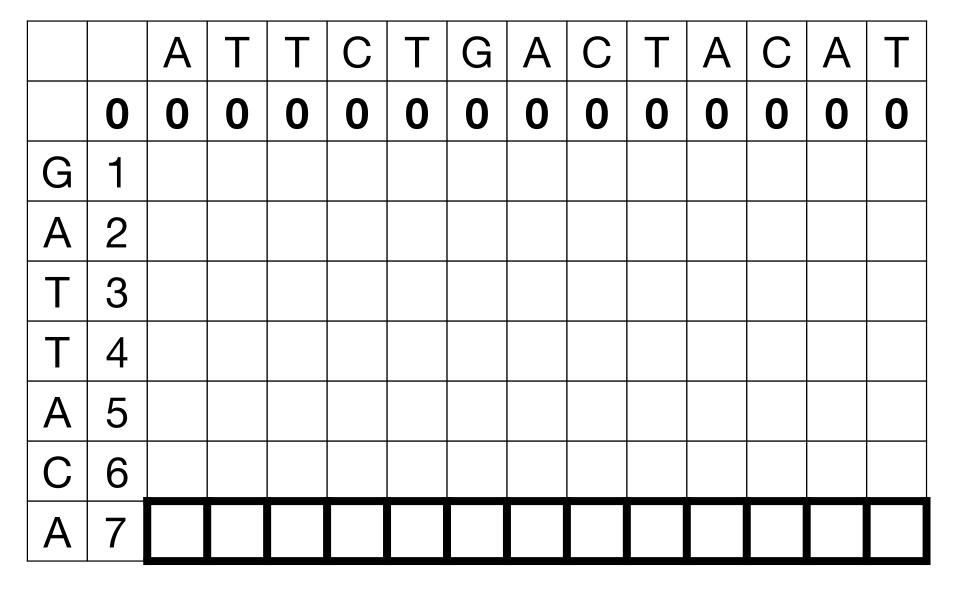
Example: Searching for the best alignment of a relatively short DNA sequence on a

long DNA sequence of a human around 3 gigabases long.

Edit Distance

		Α	Т	T	С	T	G	Α	С	Т	Α	С	Α	T
	0	1	2	3	4	5	6	7	8	9	10	11	12	13
G	1													
Α	2													
Т	3													
Т	4													
Α	5													
С	6													
A	7													

Substring Match



Find minimum cost on the last row

Some Variants of the Edit Distance

Longest Common Subsequence: Between P

and T we investigate the longest match of possibly scattered sequence of symbols

Another way is to use normal edit distance, but preventing substitutions! How?

		d	е	m	0	С	r	а	t	S
	0	0	0	0	0	0	0	0	0	0
r	0	0	0	0	0	0	1	1	1	1
е	0	0	1	1	1	1	1	1	1	1
р	0	0	1	1	1	1	1	1	1	1
u	0	0	1	1	1	1	1	1	1	1
b	0	0	1	1	1	1	1	1	1	1
I	0	0	1	1	1	1	1	1	1	1
i	0	0	1	1	1	1	1	1	1	1
С	0	0	1	1	1	2	2	2	2	2
а	0	0	1	1	1	2	3	3	3	3
n	0	0	1	1	1	2	3	3	3	3
S	0	0	1	1	1	2	3	3	3	4

P: democrats

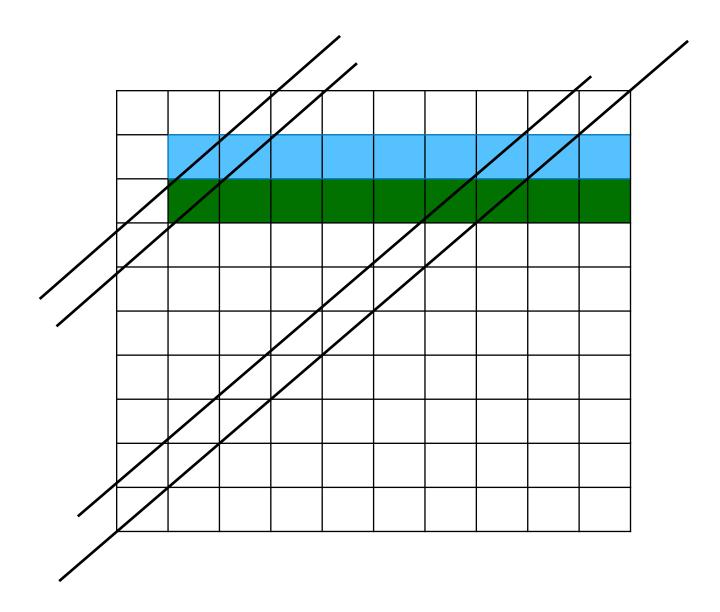
T: republicans

LCS(P,T):ecas

Edit Distance Computation Challenges

The matrix needs $O(n \cdot m)$, quadratic, space. Is there a way to reduce it?

The computation takes $O(n \cdot m)$ time, would it be possible to speed it up via parallelization?



Reading assignment

 Read the Dynamic Programming chapters from the text books, particularly from Cormen and Skiena.