



# Chapter 3 Dynamic Programming

Algorithm Theory WS 2018/19

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# Dynamic Programming (DP)



#### $DP \approx Recursion + Memoization$

**Recursion:** Express problem *recursively* in terms of (a 'small' number of) *subproblems* (of the same kind)

Memoize: Store solutions for subproblems reuse the stored solutions if the same subproblems has to be solved again

Weighted interval scheduling: subproblems W(1), W(2), W(3), ...

runtime = #subproblems · time per subproblem

## **Dynamic Programming**



"Memoization" for increasing the efficiency of a recursive solution:

 Only the *first time* a sub-problem is encountered, its solution is computed and then stored in a table. Each subsequent time that the subproblem is encountered, the value stored in the table is simply looked up and returned

(without repeated computation!).

 Computing the solution: For each sub-problem, store how the value is obtained (according to which recursive rule).

## **Dynamic Programming**



Dynamic programming / memoization can be applied if

- Optimal solution contains optimal solutions to sub-problems (recursive structure)
- Number of sub-problems that need to be considered is small

# Knapsack



- <u>n items 1, ..., n</u>, each item has weight  $w_i$  and value  $v_i$
- Knapsack (bag) of capacity W
- Goal: pack items into knapsack such that total weight is at most W and total value is maximized:

$$\max \sum_{i \in S} v_i$$
s.t.  $S \subseteq \{1, ..., n\}$  and 
$$\sum_{i \in S} w_i \leq W$$

• E.g.: jobs of length  $w_i$  and value  $v_i$ , server available for W time units, try to execute a set of jobs that maximizes the total value

#### **Recursive Structure?**

#### OPT(K):



- Optimal solution: <u>O</u>
- If  $n \notin \mathcal{O}$ : OPT(n) = OPT(n-1)
- What if  $n \in \mathcal{O}$ ?
  - Taking n gives value  $v_n$
  - But, n also occupies space  $w_n$  in the bag (knapsack)
  - There is space for  $W w_n$  total weight left!

$$\underbrace{\mathsf{OPT}(n)}_{} = v_n + \text{optimal solution with first } n-1 \text{ items}$$

$$\underbrace{\mathsf{and knapsack of capacity}}_{} W - w_n$$

# A More Complicated Recursion



**OPT**(k, x): value of optimal solution with items 1, ..., k

and knapsack of capacity x

#### **Recursion:**

and knapsack of capacity 
$$x$$

wain soal! OPT( $u, W$ )

Recursion:

OPT( $k, x$ ) = wax  $\left\{ \begin{array}{c} OPT(k-1, x) \\ opt. sol. when \\ uot using item k \end{array} \right\}$ 

The analysis sense

that it a literation  $\frac{1}{2} \left\{ \begin{array}{c} ap \\ bp \end{array} \right\}$ 

The subproblems?

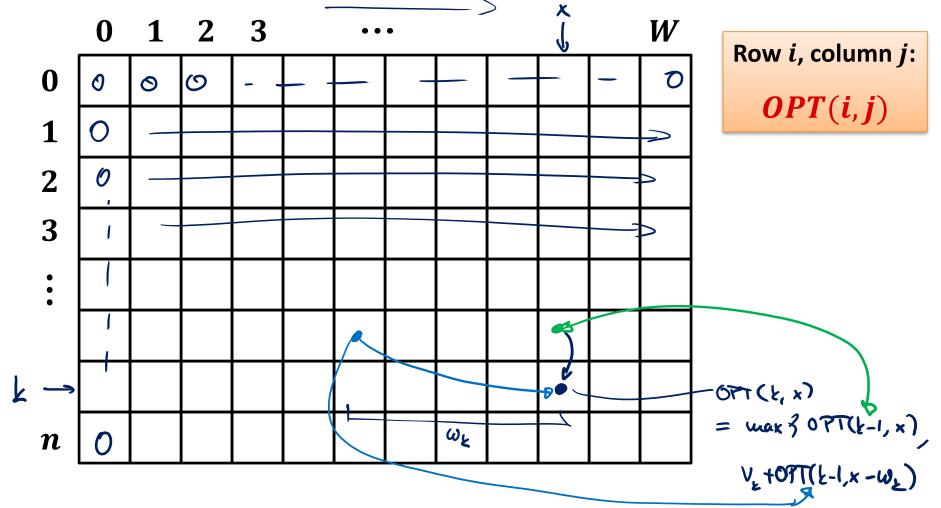
$$OPT(0, x) = 0$$

## Dynamic Programming Algorithm



Set up table for all possible OPT(k, x)-values

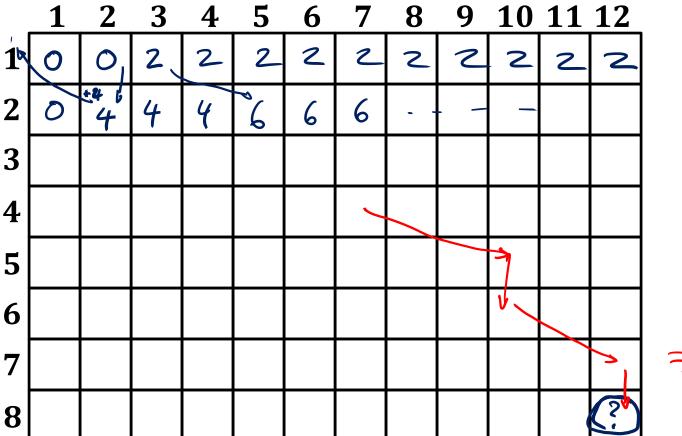
• Assume that all weights  $w_i$  are integers!



## Example



- 8 items: (3,2), (2,4), (4,1), (5,6), (3,3), (4,3), (5,4), (6,6) Knapsack capacity: 12 weight value
- $OPT(k, x) = \max\{OPT(k-1, x), OPT(k-1, x-w_k) + v_k\}$



7,5

# Running Time of Knapsack Algorithm



- Size of table:  $O(n \cdot W)$
- Time per table entry:  $O(1) \rightarrow$  overall time: O(nW)
- Computing solution (set of items to pick): Follow  $\leq n$  arrows  $\rightarrow O(n)$  time (after filling table)
- Note: Time depends on  $W \rightarrow$  can be exponential in n...
- And it is problematic if weights are not integers.

## String Matching Problems



#### **Edit distance:**

- For two given strings A and B, efficiently compute the edit distance D(A, B) (# edit operations to transform A into B) as well as a minimum sequence of edit operations that transform A into B.
- Example: mathematician → multiplication:

#### **Edit Distance**



Given: Two strings  $A=a_1a_2\dots a_m$  and  $B=b_1b_2\dots b_n$ 

**Goal:** Determine the minimum number D(A, B) of edit operations required to transform A into B

#### **Edit operations:**

- a) Replace a character from string A by a character from B
- **b) Delete** a character from string A
- c) Insert a character from string B into A

#### Edit Distance – Cost Model



- Cost for **replacing** character a by  $b: \underline{c(a, b)} \ge 0$
- Capture insert, delete by allowing  $a = \underline{\varepsilon}$  or  $b = \varepsilon$ :
  - Cost for **deleting** character  $a: c(\underline{a}, \underline{\varepsilon})$
  - Cost for **inserting** character  $b: c(\varepsilon, b)$
- Triangle inequality:

$$\underline{c(a,c)} \le c(a,b) + c(b,c)$$

→ each character is changed at most once!

• Unit cost model: 
$$c(a,b) = \begin{cases} 1, & \text{if } a \neq b \\ 0, & \text{if } a = b \end{cases}$$

#### **Recursive Structure**



Optimal "alignment" of strings (unit cost model)

bbcadfagikccm **and** abbagflrgikacc:

Consists of optimal "alignments" of sub-strings, e.g.:

-bbcagfa and 
$$-gik-ccm$$
 abb-adfl rgikacc-

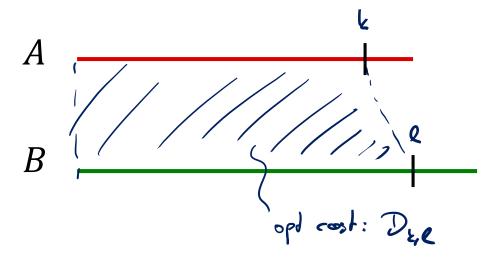
• Edit distance between  $A_{1,m}=a_1\dots a_m$  and  $B_{1,n}=b_1\dots b_n$ :

$$D(A,B) = \min_{k,\ell} \{ D(A_{1,k}, B_{1,\ell}) + D(A_{k+1,m}, B_{\ell+1,n}) \}$$

## Computation of the Edit Distance



Let 
$$A_k \coloneqq \underline{a_1 \dots a_k}$$
 ,  $B_\ell \coloneqq \underline{b_1 \dots b_\ell}$  , and 
$$\underline{D_{k,\ell}} \coloneqq \underline{D(A_k,B_\ell)}$$



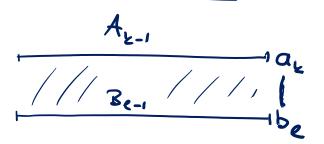
## Computation of the Edit Distance



Three ways of ending an "alignment" between  $A_k$  and  $B_\ell$ :

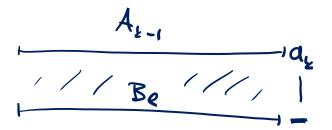
1.  $a_k$  is replaced by  $b_\ell$ :

$$D_{k,\ell} = D_{k-1,\ell-1} + c(a_k, b_\ell)$$



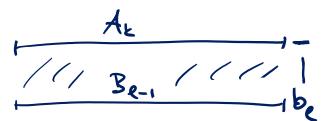
2.  $a_k$  is deleted:

$$D_{k,\ell} = D_{k-1,\ell} + \underline{c(a_k, \varepsilon)}$$



3.  $b_{\ell}$  is inserted:

$$D_{k,\ell} = D_{k,\ell-1} + c(\varepsilon, b_{\ell})$$



## Computing the Edit Distance

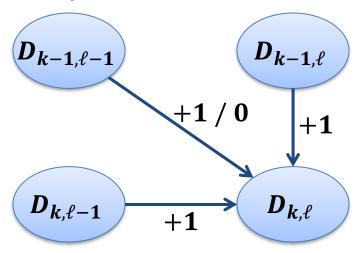


• Recurrence relation (for  $k, \ell \geq 1$ )

$$D_{k,\ell} = \min \begin{cases} D_{k-1,\ell-1} + c(a_k, b_\ell) \\ D_{k-1,\ell} + c(a_k, \varepsilon) \\ D_{k,\ell-1} + c(\varepsilon, b_\ell) \end{cases} = \min \begin{cases} D_{k-1,\ell-1} + 1 / 0 \\ D_{k-1,\ell} + 1 \\ D_{k,\ell-1} + 1 \end{cases}$$

unit cost model

• Need to compute  $D_{i,j}$  for all  $0 \le i \le k$ ,  $0 \le j \le \ell$ :



#### Recurrence Relation for the Edit Distance



#### **Base cases:**

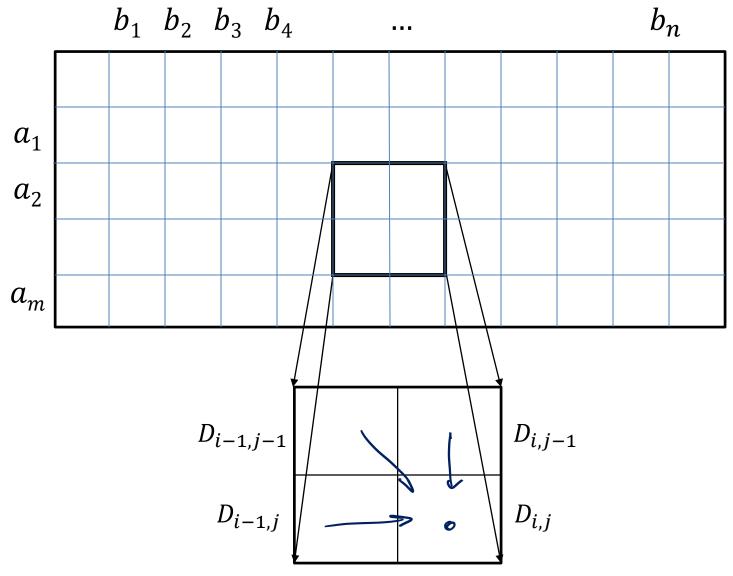
$$\begin{aligned}
D_{0,0} &= D(\varepsilon, \varepsilon) = \underline{0} \\
\overline{D_{0,j}} &= D(\varepsilon, B_j) = D_{0,j-1} + c(\varepsilon, b_j) \\
\overline{D_{i,0}} &= D(A_i, \varepsilon) = D_{i-1,0} + c(a_i, \varepsilon)
\end{aligned}$$

#### **Recurrence relation:**

$$egin{aligned} egin{aligned} oldsymbol{D_{k-1,\ell-1}} + oldsymbol{c}(oldsymbol{a_k,\ell-1}) + oldsymbol{c}(oldsymbol{a_k,\ell-1}) + oldsymbol{c}(oldsymbol{a_k,\ell-1}) + oldsymbol{c}(oldsymbol{e_k,\ell-1}) \end{aligned}$$

# Order of solving the subproblems





# Algorithm for Computing the Edit Distance



#### **Algorithm** *Edit-Distance*

**Input:** 2 strings 
$$A = a_1 \dots a_m$$
 and  $B = b_1 \dots b_n$ 

**Output:** matrix 
$$D = (D_{ij})$$

$$1 D[0,0] := 0;$$

2 for 
$$i := 1$$
 to  $m$  do  $D[i, 0] := i$ ;

3 for 
$$j := 1$$
 to  $n$  do  $D[0, j] := j$ ;

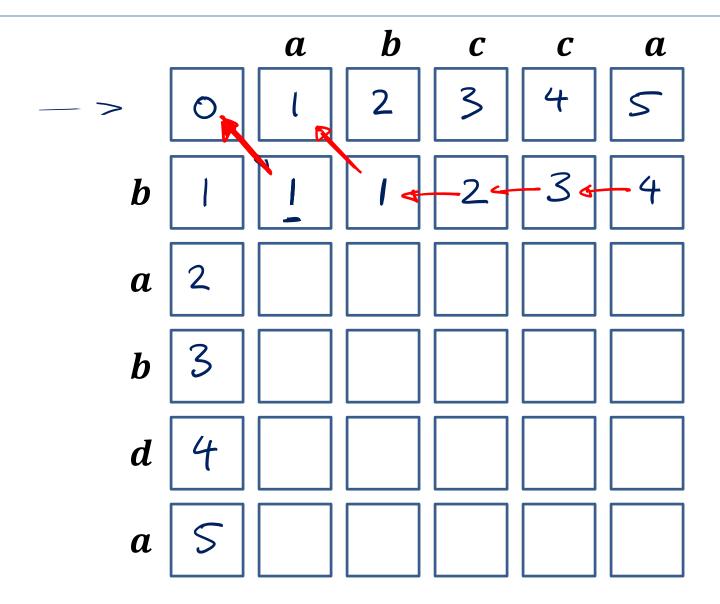
4 for 
$$i := 1$$
 to  $m$  do

5 for 
$$j := 1$$
 to  $n$  do

6 
$$D[i,j] := \min \begin{cases} D[i-1,j] + 1 \\ D[i,j-1] + 1 \\ D[i-1,j-1] + c(a_i,b_i) \end{cases}$$
;

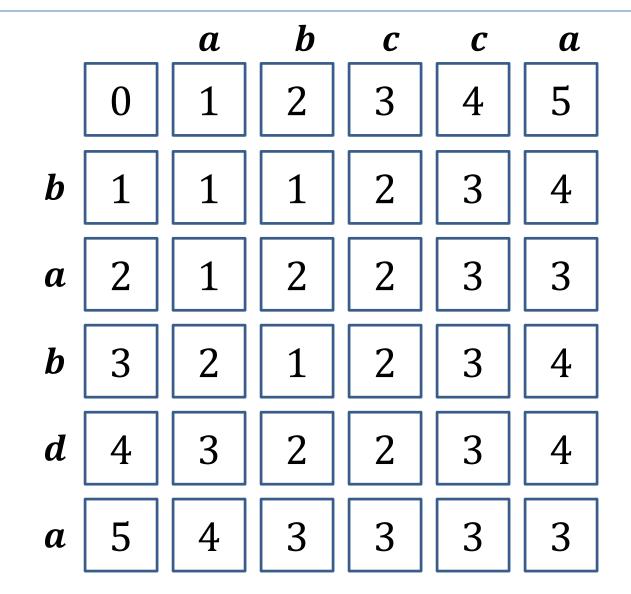
# Example





## **Edit Operations**





## Computing the Edit Operations

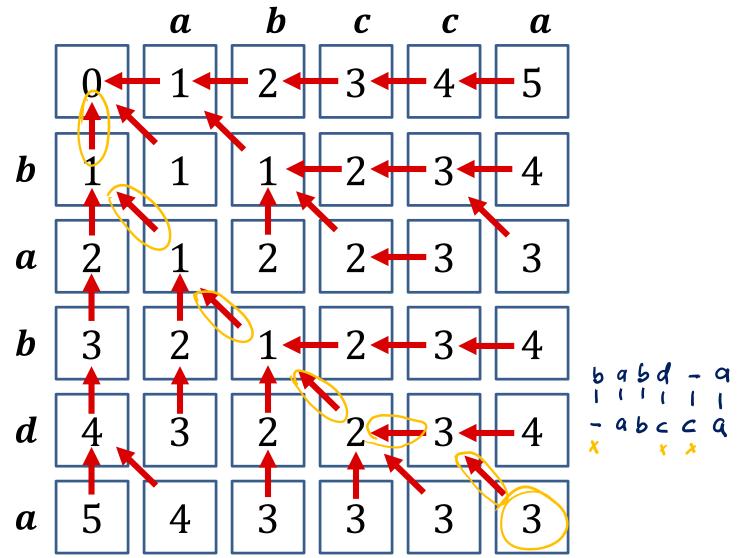


```
Algorithm Edit-Operations(i, j)
Input: matrix D (already computed)
Output: list of edit operations
1 if i = 0 and j = 0 then return empty list
2 if i \neq 0 and D[i,j] = D[i-1,j] + 1 then
     return Edit-Operations(i-1,j) \circ "delete a_i"
4 else if j \neq 0 and D[i,j] = D[i,j-1] + 1 then
     return Edit-Operations(i, j - 1) \circ ,, insert b_i"
5
  else //D[i,j] = D[i-1,j-1] + c(a_i,b_i)
     if a_i = b_i then return Edit-Operations (i-1, j-1)
     else return Edit-Operations(i-1, j-1) \circ "replace a_i by b_i"
8
```

**Initial call:** *Edit-Operations*(*m*,*n*)

# **Edit Operations**





#### Edit Distance: Summary



• Edit distance between two strings of length m and n can be computed in O(mn) time.

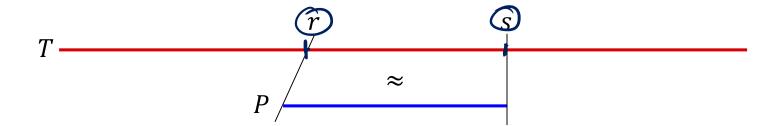
- Obtain the edit operations:
  - for each cell, store which rule(s) apply to fill the cell
  - track path backwards from cell (m, n)
  - can also be used to get all optimal "alignments"
- Unit cost model:
  - interesting special case
  - each edit operation costs 1



**Given:** strings  $T = t_1 t_2 \dots t_n$  (text) and  $P = p_1 p_2 \dots p_m$  (pattern).

**Goal:** Find an interval [r, s],  $1 \le r \le s \le n$  such that the sub-string  $T_{r,s} := t_r \dots t_s$  is the one with highest similarity to the pattern P:

$$\underset{1 \le r \le s \le n}{\operatorname{arg min}} D(T_{r,s}, P)$$





#### **Naive Solution:**

for all 
$$1 \le r \le s \le n$$
 do

compute  $D(T_{r,s}, P)$  cost:  $O((s-r) \cdot m) = O(n \cdot m)$ 

choose the minimum

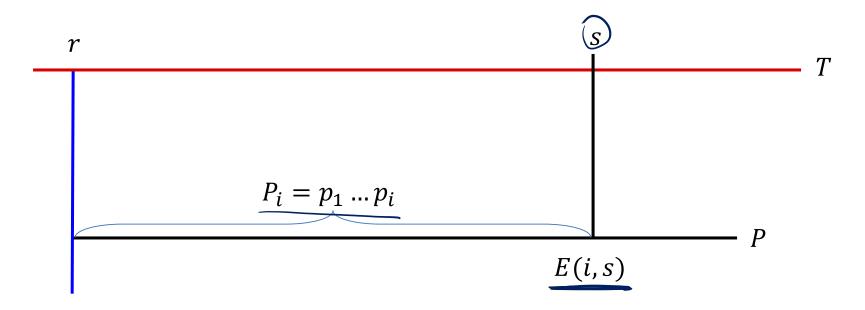
 $\longrightarrow \text{overall}: O(n^3 \cdot m)$ 

unit cost: Can be improved to  $O(n \cdot m^3)$ 



#### A related problem:

• For each position s in the text and each position i in the pattern compute the minimum edit distance E(i,s) between  $P_i = p_1 \dots p_i$  and any substring  $T_{r,s}$  of T that ends at position s.

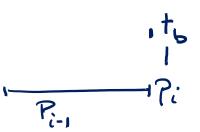




Three ways of ending optimal alignment between  $T_b$  and  $P_i$ :

1.  $t_b$  is replaced by  $p_i$ :

$$E_{b,i} = E_{b-1,i-1} + c(t_b, p_i)$$



2.  $t_b$  is deleted:

$$E_{b,i} = E_{b-1,i} + c(t_b, \varepsilon)$$

3.  $p_i$  is inserted:

$$E_{b,i} = E_{b,i-1} + c(\varepsilon, p_i)$$



#### Recurrence relation (unit cost model):

$$E_{b,i} = \min egin{cases} E_{b-1,i-1} + 1 / 0 \ E_{b-1,i} + 1 \ E_{b,i-1} + 1 \end{pmatrix}$$

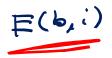
#### **Base cases:**

$$E_{0,0} = 0$$

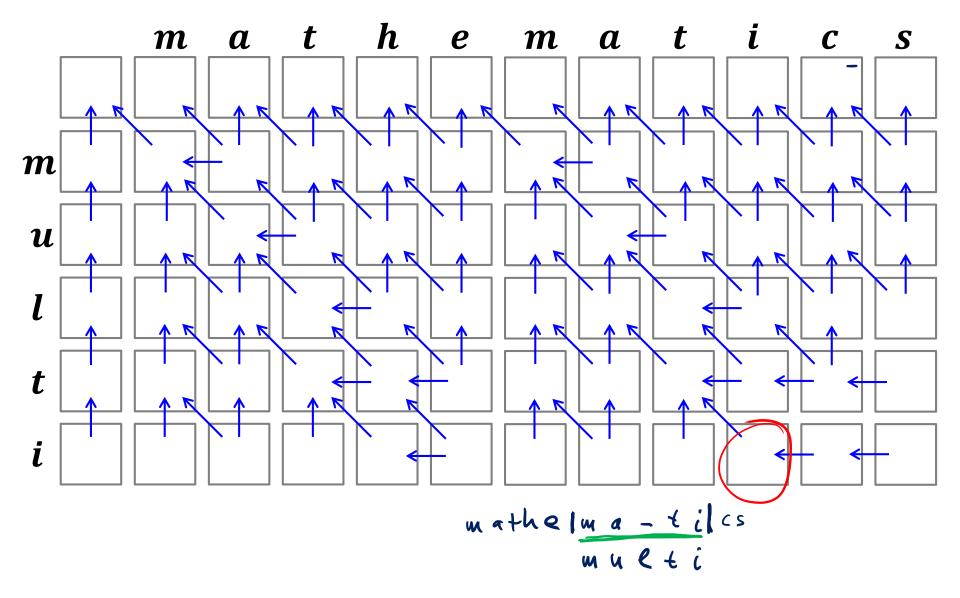
$$E_{0,i} = i$$

$$E_{i,0} = 0$$

## Example









- Optimal matching consists of optimal sub-matchings
- Optimal matching can be computed in O(mn) time
- Get matching(s):
  - Start from minimum entry/entries in bottom row
  - Follow path(s) to top row
- Algorithm to compute E(b,i) identical to edit distance algorithm, except for the initialization of E(b,0)

#### Related Problems in Bioinformatics



#### **Sequence Alignment:**

Find optimal alignment of two given DNA, RNA, or amino acid sequences.

#### **Global vs. Local Alignment:**

- Global alignment: find optimal alignment of 2 sequences
- Local alignment: find optimal alignment of sequence 1
   (patter) with sub-sequence of sequence 2 (text)