

Lecture 13

Parsing

Part 1: Definition and Representation

Machine Learning for Structured Data
Vlad Niculae · LTL, UvA · <https://vene.ro/mlsd>

Parsing

1 Definition and Representation

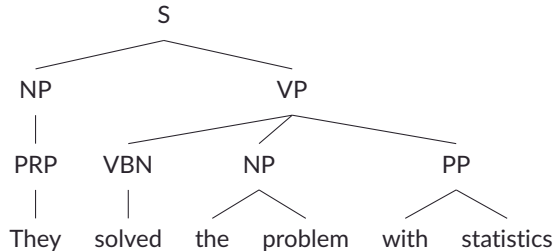
2 Bracketing: Algorithm

3 Extensions and Evaluation

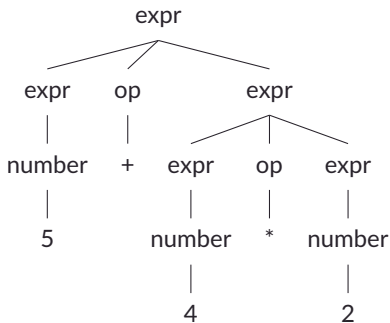
Syntactic Analysis

Syntax is an underlying structure of languages, often analyzed using parse trees.

human language (English):

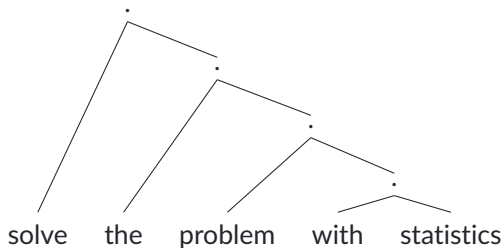
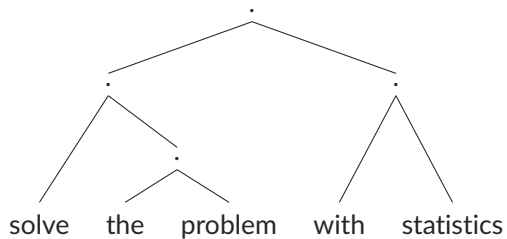


programming language (Python):



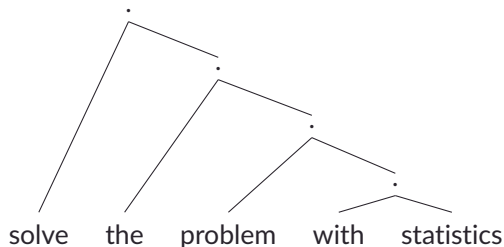
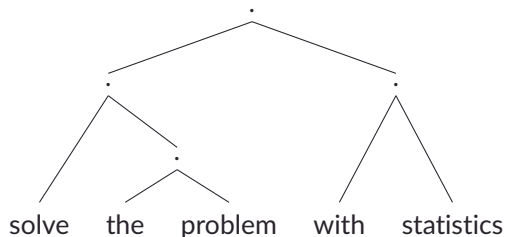
Binary Parsing

There are many kinds of complicated syntactic analysis formalisms. For simplicity, we focus on: binary trees. Let's start without labels too.



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A binary parse tree with no labelling is the same thing as a bracketing:

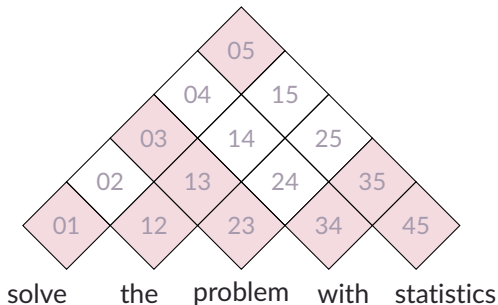
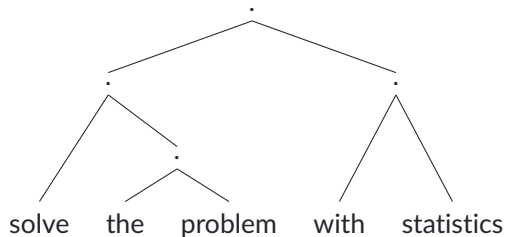
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Bracketing: Representation

Assign a score a_{ij} to the span from i to j (fencepost).

The score of a parse tree is the sum of all scores of its (nested!) spans.

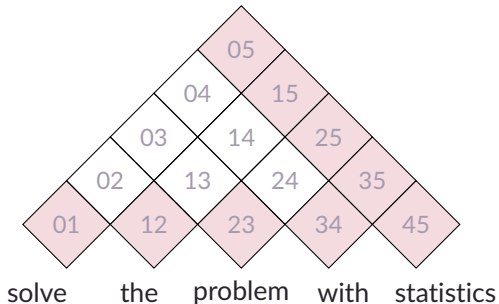
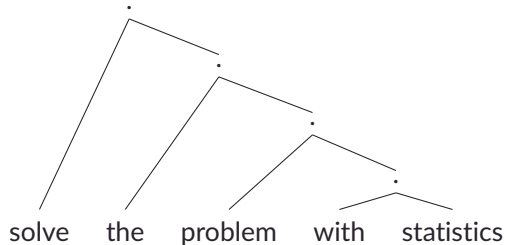


$$\text{score}(y) = a_{01} + a_{12} + a_{23} + a_{34} + a_{45} + a_{13} + a_{35} + a_{03} + a_{05}$$

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Part 2: Bracketing: Algorithm

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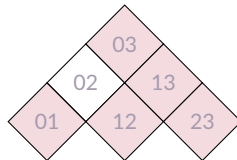
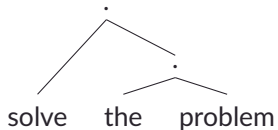
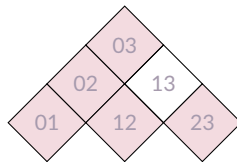
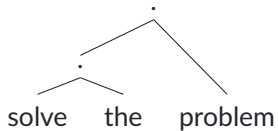
① Definition and Representation

② **Bracketing: Algorithm**

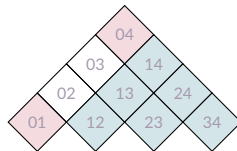
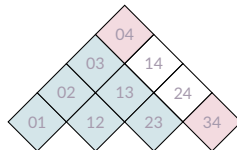
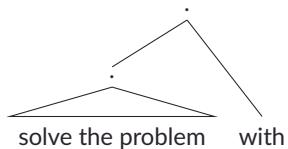
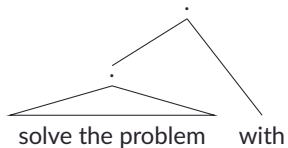
③ Extensions and Evaluation

Algorithm

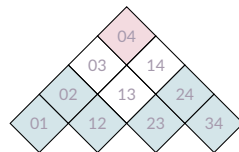
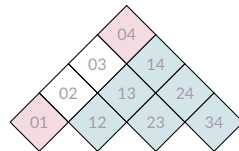
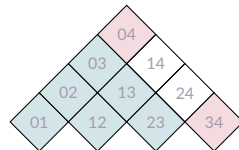
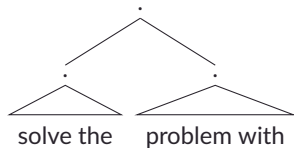
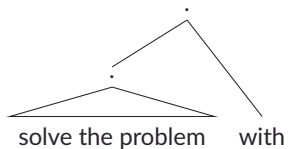
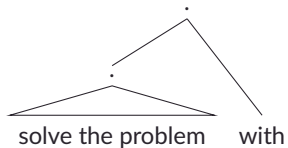
Possible parses of the subsequence (0, 3):



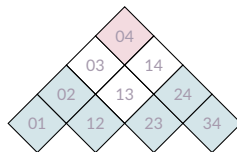
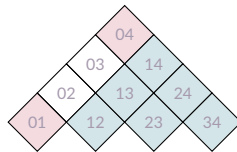
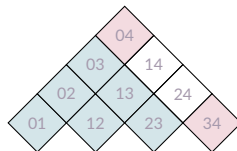
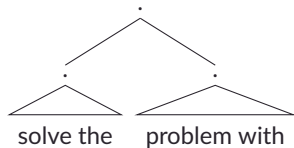
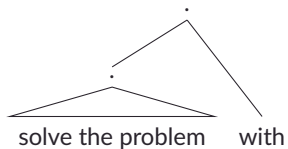
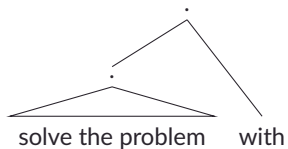
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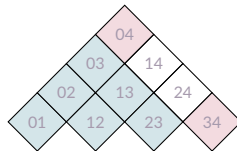
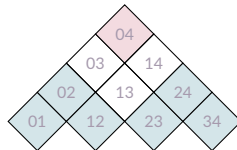
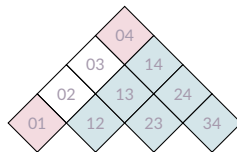


Possible parses of the subsequence (0, 4): *see the pattern?*



Dynamic Programming for Bracketing

In general: a partial parse that covers subsequence (i, j) must consist of two partial parses: one covering (i, k) and one covering (k, j) for some $i < k < j$.



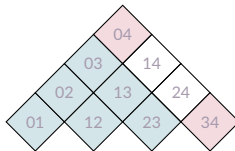
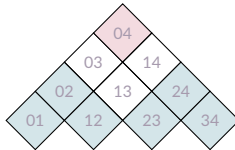
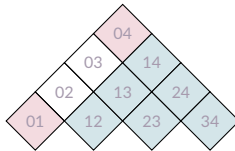
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Define M_{ij} as the maximum-scoring parse of subtree from i to j . Then:

$$M_{i,i+1} = a_{i,i+1}$$

$$M_{i,j} = \max_{i < k < j} a_{i,j} + M_{i,k} + M_{k,j}$$



Dynamic Programming for Bracketing

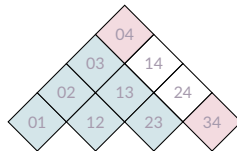
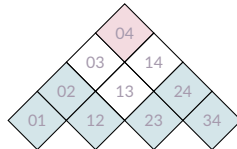
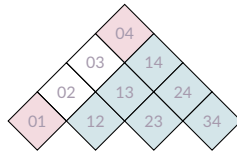
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Fill in the table bottom-up:
dynamic programming.



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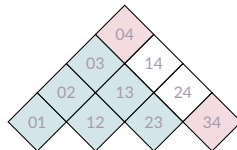
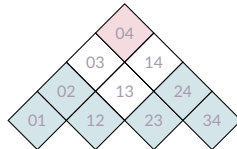
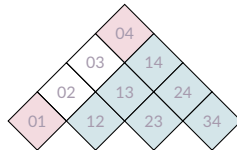
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Fill in the table bottom-up:
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CYK algorithm: Cocke, Younger, Kasami
independently discovered it in the 1960s.



The CYK Algorithm

input: Scores $a_{i,j}$ for $0 \leq i < j \leq n$
 $M_{i,j} = 0, \quad \pi_{i,j} = -1$, for $0 \leq i < j \leq n$.
 $M_{i,i+1} = a_{i,i+1}$ for $0 \leq i < n$.

Forward: compute max. scores for each span recursively

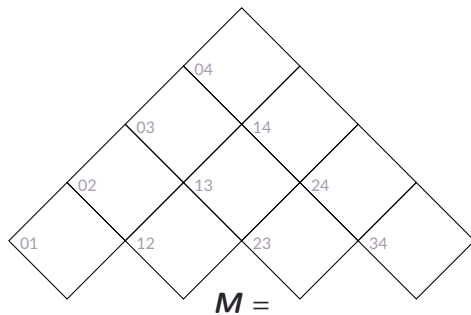
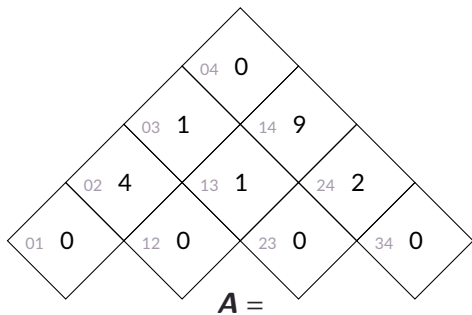
```
for  $s = 2$  to  $n$  do  
  for  $i = 0$  to  $n - s$  do  
     $j = i + s$   
     $M_{i,j} = \max_{i < k < j} a_{i,j} + M_{i,k} + M_{k,j}$   
     $\pi_{i,j} = \arg \max_{i < k < j} a_{i,j} + M_{i,k} + M_{k,j}$ 
```

Backward: follow backpointers

```
 $y^* = ()$ ,  $Q = \{(0, n)\}$ .  
while  $Q$  not empty do  
   $\text{pop } (i, j)$  from  $Q$   
   $y^* = y^* + (i, j)$   
   $k = \pi_{i,j}$   
  push  $(i, k)$  and  $(k, j)$  to  $Q$  if  $k > 0$ .
```

output: The highest-scoring bracketing y^* , and its total score f^* .

The CYK Algorithm: Example



The *Inside* Algorithm for $\log Z$

input: Scores $a_{i,j}$ for $0 \leq i < j \leq n$

$Q_{i,j} = 0$, for $0 \leq i < j \leq n$.

$Q_{i,i+1} = a_{i,i+1}$ for $0 \leq i < n$.

Forward: compute logsumexp for each span recursively

for $s = 2$ to n **do**

for $i = 0$ to $n - s$ **do**

$j = i + s$

$Q_{i,j} = \log \sum_{i < k < j} \exp a_{i,j} + Q_{i,k} + Q_{k,j}$

CYK vs Segmentation

- The two algorithms have the same inputs: a table of scores for every possible segment.
- The segmentation problem seeks the best low-level chunking.
- CYK seeks an entire tree of chunk “splits”.
- Segmentation is the simplest possible DAG. CYK cannot be represented as a DAG at all!

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Part 3: Extensions and Evaluation

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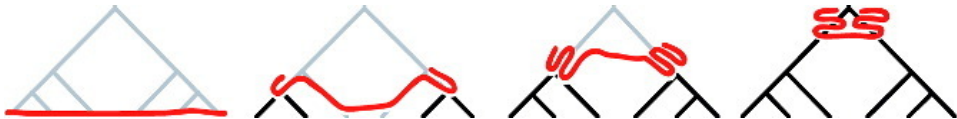
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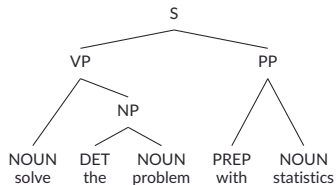
③ Extensions and Evaluation

Protein Folding as Binary Parsing



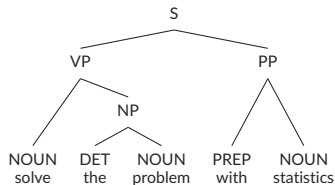
Julia Hockenmaier, Aravind K. Joshi, Ken A. Dill,
Routes are trees: The parsing perspective on protein folding. *Proteins*, 66-1, 2007.

Labelled Parsing



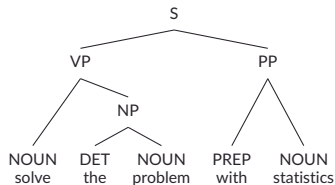
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Labelled Parsing



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- In this case, like for segmentation, we can pick the best label for each segment before starting Viterbi, and ignore the rest.

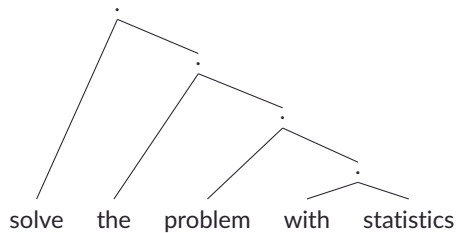
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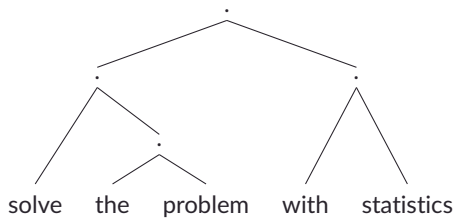
- Simple case: replace all segments with labeled segments (i, j, c) .
- In this case, like for segmentation, we can pick the best label for each segment before starting Viterbi, and ignore the rest.
- We may want “transition scores”
e.g., prefer S out of NP VP, dislike S out of VP PP.
 - related to *probabilistic context-free grammars*
 - handled by a similar DP algorithm,
:wejk higher complexity
(loop also over all combinations of labels).

Evaluation

y_{pred}

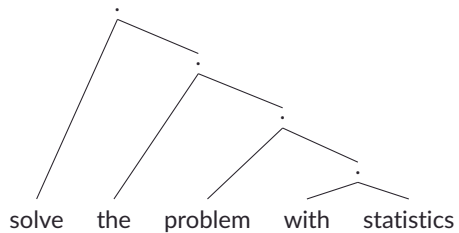


y_{true}



Evaluation

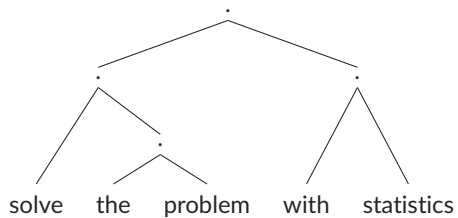
y_{pred}



Predicted spans:

(0, 1), (0, 5), (1, 2), (1, 5), (2, 3), (2, 5), (3, 4), (3, 5), (4, 5)

y_{true}

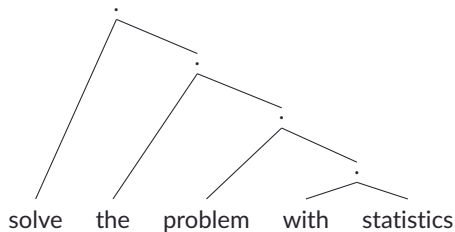


True spans:

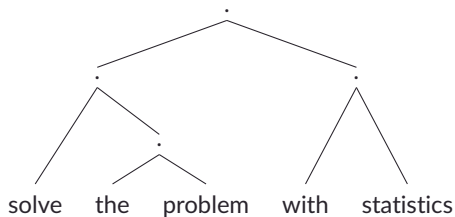
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Evaluation

y_{pred}



y_{true}



Predicted spans:

(0, 1), (0, 5), (1, 2), (1, 5), (2, 3), (2, 5), (3, 4), (3, 5), (4, 5)

$$P = \frac{\text{n. correct}}{\text{n. predicted}}$$

$$R = \frac{\text{n. correct}}{\text{n. true}}$$

True spans:

(0, 1), (0, 3), (0, 5), (1, 2), (1, 3), (2, 3), (3, 4), (3, 5), (4, 5)

$$F_1 = \frac{2PR}{P+R}$$

Note: in the unlabelled case, $P=R$, since the number of segments in a bracketing is always the same.

In the labelled case: usually common to compute per-label $P/R/F$, averaged over the entire dataset.

In linguistic applications, "real" parsing evaluation is more complicated, since trees are not binary.

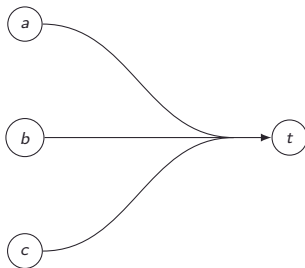


Hyperedges and Hypergraphs

There is a formalism that generalizes DAGs and can express the CYK parsing problem, but its details are too complicated for our scope. Nevertheless, here is a glimpse.

Given nodes $V = \{1, 2, \dots, n\}$

- instead of **edges**: $(s, t) : s \in V, t \in V$.
- define **hyperedges**: $((s_1, \dots, s_k), t) : s_i \in V, t \in V$.





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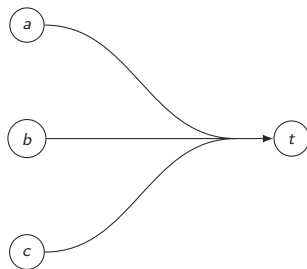
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- instead of **edges**: $(s, t) : s \in V, t \in V$.
- define **hyperedges**: $((s_1, \dots, s_k), t) : s_i \in V, t \in V$.


Any directed graph can be represented as a directed hypergraph: if (s, t) is an edge in G , then make $((s), t)$ a hyperedge in HG .

Generalizations of DAG and topological sort exist; and Viterbi & Forward algorithms work.

Read more: Liang Huang, [Advanced Dynamic Programming in Semiring and Hypergraph Frameworks](#), COLING 2008 tutorial.



Summary

- Binary parsing / bracketing can be solved with dynamic programming (even if it can't be represented as a DAG)
- Applications in computational linguistics:
related to *grammars*.
-  Can generalize the algorithms seen to compute logsumexp and sampling with DP, using a *hypergraphs* formalism.