Accelerated Natural Language Processing 2018

Lecture 13: (Features), Parsing as search and as dynamic programming

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1. Features [Material in slides 1--8 are not examinable]

The name feature has been around in Linguistics for a long time

The basic idea is to capture generalisations by decomposing monolithic categories into collections of simpler features $% \left({{{\rm{c}}_{\rm{s}}}} \right)$

Originally developed for phonology, where we might have e.g.

/i/ +high, +front /e/ -high, +front /o/ -high, -front /u/ +high, -front

Where we can now 'explain' why *fi/* and */u/* behave similarly in certain cases, while *fi/* and */e/* go together in other cases.

Those are all binary features

- sometimes also used at the level of syntax
- +/- singular; +/- finite

2. Features, cont'd

But more often we find features whose values are some enumerated type

person: {1st,2nd,3rd}; number: {sg, pl}; ntype: {count, mass}

We'll follow J&M and write collections of features like this:

person 3rd number pl

It will be convenient to generalise and allow features to take feature bundles as values:

ntype	count		
agreement	person number	3rd pl	
	ntype agreement	ntype count agreement person number	

3. Features in use

We can now add feature bundles to categories in our grammars

In practice we allow some further notational conveniences:

Not all features need be specified (e.g. the number feature for 'sheep')
 In rules, we allow the values of features to be variables
 And we can add constraints in terms of those variables to rules

For example

Γ	ntype	count		
N	agreement	person number	3rd pl	→ men dogs cats
-				

NP[agreement $x] \rightarrow D[$ agreement y] N[agreement z] x = y = z

4. Features: more than notational convenience?

At one level, features are just a convenience

The allow us to write lexicon entries and rules more transparently

But they also "capture generalisations"

If we write a pair of rules using some (in principle opaque) complex category labels, they are not obviously related in any way:

$S \rightarrow NP_{sg}VP_{sg}$ $S \rightarrow NP_{pl}VP_{pl}$

It appears as if we have to justify each of these independently

or that we might have had one without the other

or had S → NP_{sg}VP_{p1} just as well

Whereas when we write

 $S \rightarrow NP VP [\langle NP agreement \rangle = \langle VP agreement \rangle]$

we are making a stronger claim, even though 'behind the scenes' this single line corresponds to a collection of simple atomic-category rules

5. Infinity again: categories

Once you move to feature bundles as the values of features

You can in principle have an infinite number of categories
 By having one or more recursive features

- And, as with infinite numbers of rules, that actually changes your position on the Chomsky hierarchy
- One strand of modern grammatical theory
- From GPSG to HPSG to so-called sign-based grammatical theories

Puts essentially all the expressive power of the grammar into feature structures

6. Unification

When we write '=' between two feature paths or feature variables, we mean more than an equality test

Consider the noun phrase "a sheep", and the following rules





 $NP[agreement x] \rightarrow D[agreement y] N[agreement z] x = y = z$

The resulting parse tree reveals that we have not only tested for compatibility between the various feature structures, we ve actually merged them: $NF|_{Byregent} \odot]$

(D) N [agreement (D) [person 3rd] number sg] sheep

where by the 0 we mean that all three agreement values are the the same feature structure

7. Unification, cont'd

The implications of unification run deep

a Nagreement © person a linumber a sheep

The three occurrences of @ don't just appear the same

- They are the same
- That is, a single structure, shared 3 times
 So any change to one in the future will be a change to all
- As would be the case with e.g. "the sheep runs" or "the sheep run"

J&M give a detailed introduction to unification, which is what this is called, in section 15.2 (J&M 2nd ed.), and a formal definition in section 15.4.

The directed acyclic graph (DAG) way of drawing feature structures used in J&M 15.4 makes clearer when necessary structure identity is the case, as opposed to contingent value equality



8. Parsers

A **parser** is an algorithm that computes a structure for an input string given a grammar. All parsers have two fundamental properties:

Directionality The sequence in which the structures are constructed Almost always top-down or bottom-u

Search strategy The order in which the search space of possible analyses is explored • Usually depth-first, breadth-first or best-first

9. Recursive Descent Parsing

A **recursive descent** parser treats a grammar as a specification of how to break down a top-level goal into subgoals This means that it works very similarly to a particular blind approach to constructing a rewriting

interpretation derivation

Directionality Top-down:

starts from the start symbol of the grammar
works down to the terminals

Search strategy Depth-first:

 expands the left-most unsatisfied non-terminal · until it gets to a terminal which either matches the next item in the input or it doesn't

10. Recursive Descent Parsing: preliminaries

- We're trying to build a parse tree, given
- a grammar
 an input, i.e. a sequence of terminal symbols

As for any other depth-first search, we may have to backtrack

- · So we must keep track of backtrack points
- And whenever we make a choice among several rules to try, we add a backtrack point consisting of a partial tree
- the remaining as-yet-unexplored rules and the as-yet-unconsumed items of input
- Note that, to make the search go depth-first
- · We'll use a stack to keep track

· That is, we'll operate last in, first out (LIFO)

Finally, we'll need a notion of where the focus of attention is in the tree we're building

• We'll call this the subgoal

11. Recursive Descent Parsing: Algorithm sketch

We start with

 a tree consisting of an 'S' node with no children This node is currently the subgoal An empty stack
An input sequence



Repeatedly

- If the subgoal is a non-terminal

 Choose a rule from the set of rules in the grammar whose left-hand sides match the subgoal
 For example, the very first time around the loop, we might choose

 S - NP VP
 - b. add children to the subgoal node corresponding to the symbols in the right-hand side of the chosen rule, in order In our example, that is two children, \mathbb{N} and \mathbb{VP}



ii. Advance the subgoal iii. Go back to (1) c. Otherwise (they don't match), Backtrack

12. Recursive Descent Parsing: Algorithm

sketch. concluded

The three imperative actions in the preceding algorithm are defined as follows:

- Choose Pick one member from the set of rules 1. If the set has only one member, you're done
 - Otherwise, push a new backtrack point onto the stack
 With the unchosen rules, the current tree and subgoal and the current (unconsumed) input sequence
- Advance Change the subgoal, as follows:
 - If the current subgoal has a sibling to its right, pick that
 - Failing which, if the current subgoal is not the root, set the subgoal to the current subgoal's parent, and go back to (1)

 - a Tailing which, if the input is empty, we win
 The current subgoal is the 'S' at the root, and it is the top node of a complete parse tree for the original input
 Otherwise, Backtrack

Backtrack

- Try to, as it were, change your mind. That is: Unless the stack is empty, **pop** the top backtrack point off the backtrack stack and
 a. Set the tree, subgoal and input from it
 b. **Choose** a rule from its set of rules c. Go back to step (1b) of the algorithm
- Otherwise (the stack is step (1) of the algorithm
 Otherwise (the stack is step)
 We lose!
 There is no parse for the input with the grammar
- We'll see the operation of this algorithm in detail in this week's lab

13. Search Strategies

Schematic view of the top-down search space:



In depth-first search the parser

- · explores one branch of the search space at a time
- For example, using a stack (last-in, first-out) of incomplete trees to try to expand





In breadth-first search the parser

 explores all possible branches in parallel For example, using a **queue** (first-in, first out) of incomplete trees to try to expand









The bottom-up search space works, as the name implies, from the leaves upwards Trying to build and combine small trees into larger ones
 The parser we look at in detail in the next lectures works that way

14. Shift-Reduce Parsing

Search strategy does not imply a particular directionality in which structures are built. Recursive descent parsing searches depth-first and builds top-down Although Shift-reduce parsing also searches depth-first, in contrast it builds structures bottom-up.

It does this by repeatedly

shifting terminal symbols from the input string onto a stack
 reducing some elements of the stack to the LHS side of a rule when they match its

As described, this is just a recogniser

· You win if you end up with a single '5' on the stack and no more input

Actual parsing requires more bookkeeping

Given certain constraints, it is possible to pre-compute auxiliary information about the grammar and exploit it during parsing so that no backtracking is required.

Modern computer languages are often parsed this way

· But grammars for natural languages don't (usually) satisfy the relevant constraints

15. Global and Local Ambiguity

A string can have more than one structural analysis (called **global ambiguity**) for one or both of two reasons:

 Grammatical rules allow for different attachment options: · Lexical rules that allow a word to be in more than one word class

Within a single analysis, some sub-strings can be analysed in more than one way

- even if not all these sub-string analyses 'survive'
 That is, if they are not compatible with any complete analysis of the entire string
 This is called **local ambiguity**
- Local ambiguity is very common in natural languages as described by formal grammars

All depth-first parsing is inherently **serial**, and serial parsers can be massively inefficient when faced with **local ambiguity**.

16. Complexity

Depth-first parsing strategies demonstrate other problems with "parsing as search":

- Structural ambiguity in the grammar and lexical ambiguity in the words (that is, words occurring under more than one part of speech) may lead the parser down a wrong path
- wrong path 2. So the same sub-tree may be built several times = whenever a path fails, the parser abandons any subtrees computed since the last backtrack point, backtracks and starts again
- The complexity of this **blind backtracking** is exponential in the worst case because of repeated **re-analysis** of the same sub-string.

· We'll experience this first-hand in this week's lab

Chart parsing is the name given to a family of solutions to this problem

17. Dynamic Programming

It seems like we should be able to avoid the kind of repeated reparsing a simple recursive descent parser must often do

A CFG parser, that is, a context-free parser, should be able to avoid re-analyzing sub-strings · because the analysis of any sub-string is independent of the rest of the parse



The parser's exploration of its search space can exploit this independence if the parser uses dynamic programming.

Dynamic programming is the basis for all chart parsing algorithms.

18. Parsing as dynamic programming

Given a problem, dynamic programming systematically fills a table of solutions to sub-pro

- A process sometimes called memoisation
- Once solutions to all sub-problems have been accumulated
- DP solves the overall problem by composing them
- For parsing, sub-problems are analyses of sub-strings
- which can be memoised

in a chart
also know as a well-formed substring table, WFST

Each entry in the chart or WFST corresponds to a complete **constituent** (sub-tree), indexed by the start and end of the sub-string that it covers

Active chart parsing goes further, and uses the chart for partial results as well

19. Depicting a WFST/Chart

A well-formed substring table (aka chart) can be depicted as either a matrix or a graph · Both contain the same information

When a WFST (aka chart) is depicted as a matrix:

- Rows and columns of the matrix correspond to the start and end positions of a span of items from the input
 That is, starting right before the first word, ending right after the final one
- A cell in the matrix corresponds to the sub-string that starts at the row index and ends at the column index A cell can contain
- can contain information about the **type** of constituent (or constituents) that span(s) the substring pointers to its sub-constituents (In the case of **active** chart parsers, **predictions** about what constituents might follow the substring)

20. Depicting a WFST as a matrix

Here's a sample matrix, part-way through a parse



0 See 1 with 2 a 3 telescope 4 in 5 hand 6

We can read this as saying:

There is a PP from 1 to 4

 Because there is a Prep from 1 to 2
 and an NP from 2 to 4

21. Depicting a WFST as a graph

A sample graph, for the same situation mid-parse

 Here, nodes (or vertices) apresent positions in the text string, starting before the first words of the final word.
 action engels
 action engels<



22. Algorithms for chart parsing



23. CKY Algorithm

CKY (Cocke, Kasami, Younger) is an algorithm for recognising constituents and recording them in the chart (WFST).

CKY was originally defined for Chomsky Normal Form



(Much more recently, this restriction has been lifted in a version by <u>Lang and Leiss</u>)
 The example below follows them in part, also allowing unary rules of the form A → B

We can enter constituent A in cell (i,j) iff either

there is a rule A → b and
 b is found in cell (i, j)

or if there is a rule λ → B ⊂ and there is at least one k between i and j such that
 a is found in cell (i, k)
 C is found in cell (k, j)

24. CKY parsing, cont'd

Proceeding systematically bottom-up, CKY guarantees that the parser only looks for rules which might yield a constituent from is to j **after** it has found all the constituents that might contribute to it, that is

- That are shorter than it is
 That end at or to the left of j
- That end at or to the left or 3
 This guarantees that every possible constituent will be found



Note that this process manifests the fundamental weakness of blind bottom-up parsing:

 Large numbers of constituents will be found which do not participate in the ultimately spanning 'correct' analyses.

25. Visualising the chart: YACFG

Grammatical rules Lexical rules

$S \rightarrow NP VP$	Det → a the (determiner)	
NP → Det Nom	N → fish frogs soup (noun)	
$NP \rightarrow Nom$	Prep → in for (preposition)	
Nom → N SRel	TV → saw ate (transitive verb)	
Nom \rightarrow N	IV → fish swim (intransitive verb)	
VP → TV NP	Relpro → that (relative pronoun)	
$VP \rightarrow IV PP$		
$VP \rightarrow IV$		
PP → Prep NP		
SRel → Relpro VP		
Nom: nominal (the part of the NP after the determiner, if any)		

SRel: subject relative clause, as in the frogs that ate fish.

Non-terminals occuring (only) on the LHS of lexical rules are sometimes called pre-terminals

In the above grammar, that's Det, N, Prep, TV, IV, Relpro

Sometimes instead of sequences of words

we just parse sequences of pre-terminals
At least during grammar development