

# Convolution kernels for natural language (Collins and Duffy, 2001)

LING 572

Advanced Statistical Methods for NLP

February 20, 2020

# Highlights

- Introduce a tree kernel
- Show how it is used for reranking

# Reranking

# Reranking

- Training data:

$\{(x_i, y_i)\}$  and for each  $x_i$ , a set of candidates  $\{y_{ij}\}$ .  
and one of  $y_{ij}$  is the same as  $y_i$ .

- Goal: create a module that reranks candidates

- The reranker is used as a post-processor.

- In this paper, build a reranker for parsing

$x_i$  is a sentence,  $y_{ij}$  is a parse tree.

Notation:  $\{(s_i, t_i)\}$ ,  $C(s_i) = \{x_{ij}\}$

# Formulating the problem

$$\{(s_i, t_i)\}, C(s_i) = \{x_{ij}\}$$

$h(x_{ij})$  is the feature vector of candidate  $x_{ij}$ .

Let  $x_{i1}$  be the correct parse for  $s_i$ .

Training: calculate  $\vec{w}$

$$\text{Decoding: } x^* = \operatorname{argmax}_{x \in C(s)} \vec{w} \cdot h(x)$$

# Reranking: Training

Minimize  $\|\vec{w}\|^2$  subject to the constraints

$$\vec{w} \cdot h(x_{i1}) \geq \vec{w} \cdot h(x_{ij}), \forall i, \forall j \geq 2$$



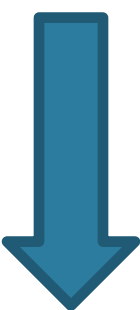
$$\vec{w} \cdot (h(x_{i1}) - h(x_{ij})) \geq 1, \forall i, \forall j \geq 2$$



Recall that in SVM  $\vec{w} = \sum_i \alpha_i y_i \vec{x}_i$

$$\vec{w} = \sum_{(i,j)} \alpha_{ij} (h(x_{i1}) - h(x_{ij}))$$

$$f(x) = \vec{w} \cdot x = \sum_{ij} \alpha_{ij} (h(x_{i1}) \cdot h(x) - h(x_{ij}) \cdot h(x))$$



With the kernel trick

$$f(x) = \sum_{ij} \alpha_{ij} (K(x_{i1}, x) - K(x_{ij}, x))$$

# Perceptron training

$$f(x) = \vec{w} \cdot x = \sum_{ij} \alpha_{ij} (h(x_{i1}) \cdot h(x) - h(x_{ij}) \cdot h(x))$$

$$\alpha_{i,j} = 0;$$

for each sentence  $i$

for each  $j > 1$

if  $f(x_{i1}) < f(x_{ij})$  then  $\alpha_{ij}++$ ;

# Tree kernel

$$f(x) = \sum_{ij} \alpha_{ij} (K(x_{i1}, x) - K(x_{ij}, x))$$

$$K: X \times X \rightarrow R$$

Each member of  $X$  is a parse tree.

What is a good tree kernel?



# A tree kernel

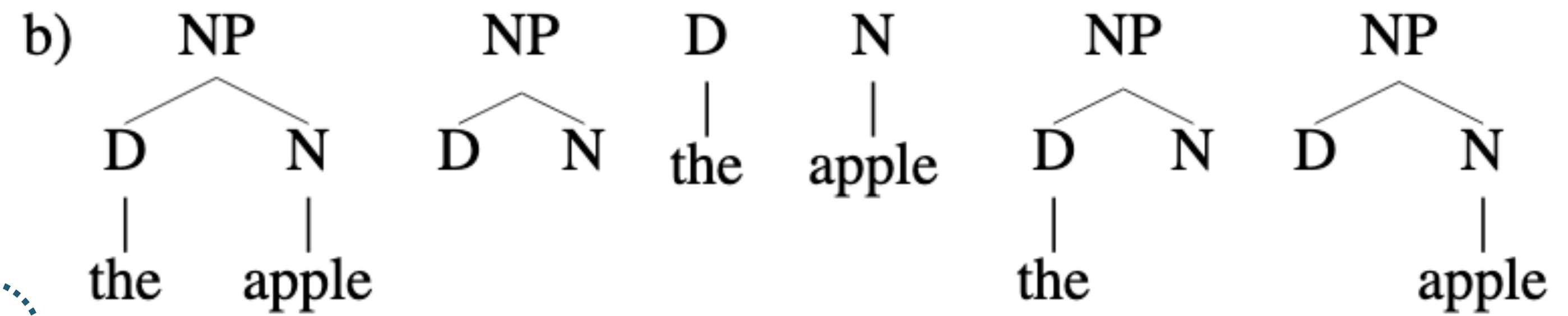
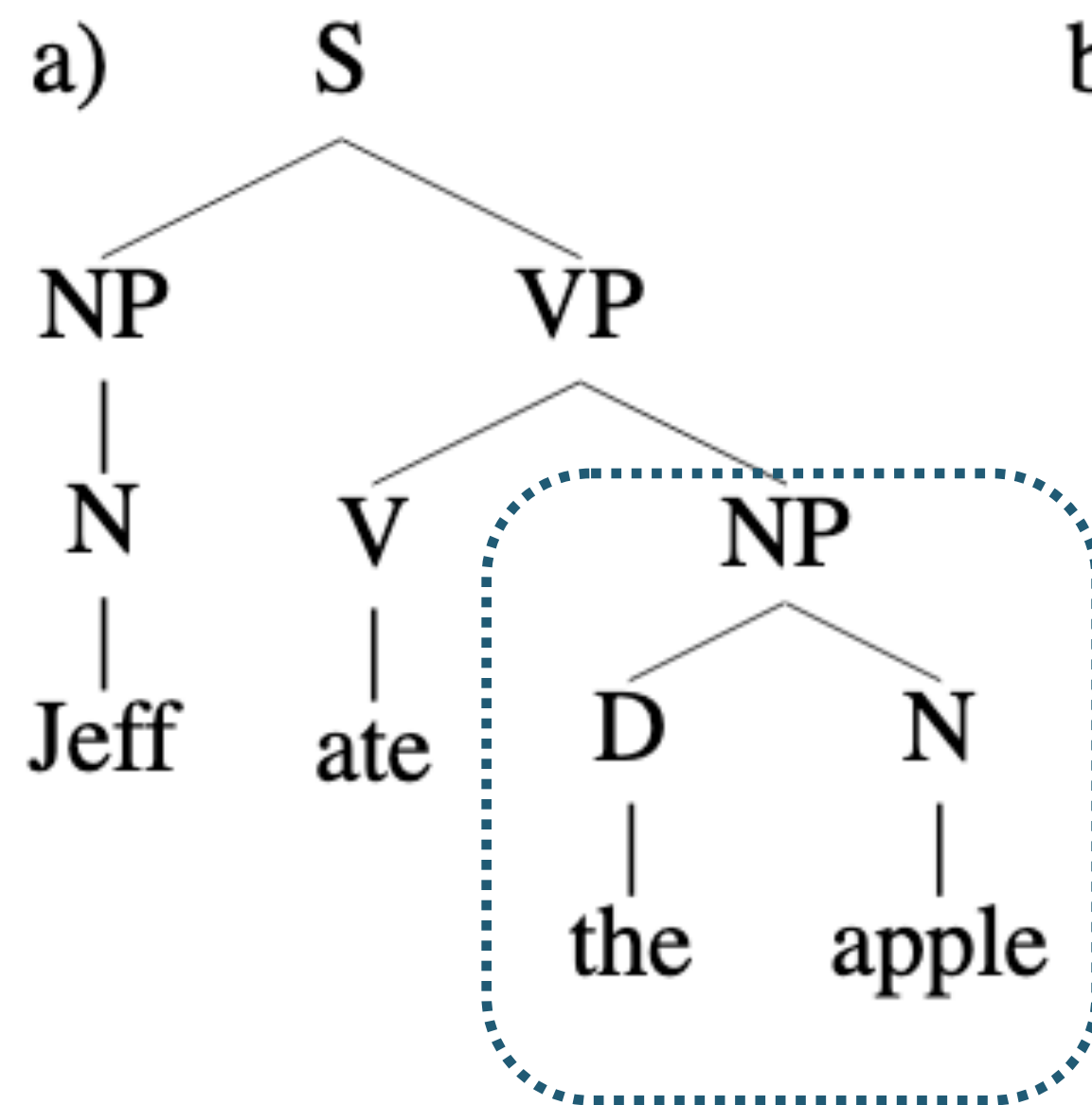
# Intuition

- Given two trees  $T1$  and  $T2$ , the more subtrees  $T1$  and  $T2$  share, the more similar they are.
- Method:
  - For each tree, enumerate all the subtrees
  - Count how many are in common
- Do it in an efficient way

# Definition of subtree

- A subtree is a subgraph which has **more than** one node, with the restriction that entire (not partial) rule productions must be included.
- “A subtree rooted at node  $n$ ” means “a subtree whose root is  $n$ ”.

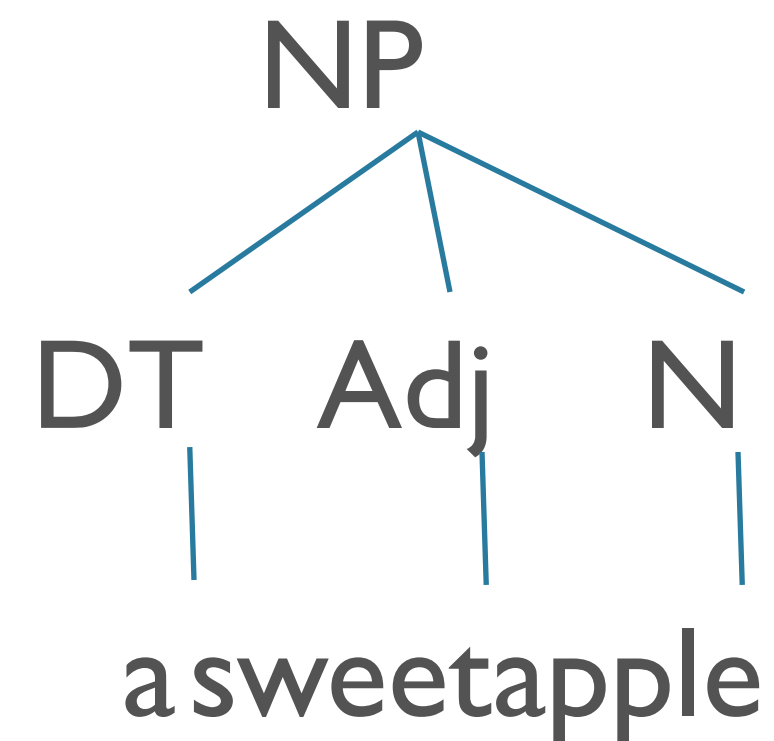
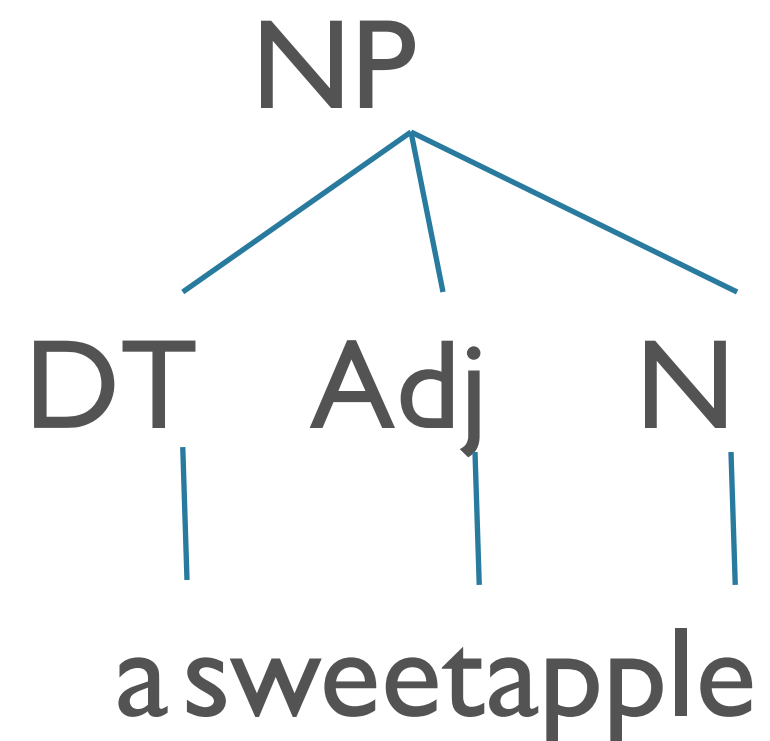
# An example



# $C(n1, n2)$

$C(n1, n2)$  counts the number of common subtrees rooted at  $n1$  and  $n2$ .

$C(n1, n2) = ??$



# Calculating $C(n1, n2)$

If the productions at  $n1$  and  $n2$  are different

then  $C(n1, n2) = 0$

else if  $n1$  and  $n2$  are pre-terminals

then  $C(n1, n2) = 1$

else  $C(n_1, n_2) = \prod_{j=1}^{nc(n1)} (1 + C(ch(n1, j), ch(n2, j)))$

# Representing a tree as a feature vector

Let  $ST$  be the set of sub-trees in **any** tree

$$ST = \{s_1, s_2, \dots, s_n, \dots\}$$

Let  $h_i(T)$  be the num of occurrences of  $s_i$  in  $T$

$$h(T) = (h_1(T), h_2(T), \dots, h_n(T), \dots)$$

$$I_i(n) = \begin{cases} 1 & \text{if } s_i \text{ is a subtree rooted at } n. \\ 0 & \text{otherwise} \end{cases}$$

$$h_i(T_1) = \sum_{n_1 \in N_1} I_i(n_1), \text{ where } N_1 \text{ is the set of nodes in } T_1$$

$$h_i(T_2) = \sum_{n_2 \in N_2} I_i(n_2)$$

# A tree kernel

$$\begin{aligned} h(T_1) \cdot h(T_2) &= \sum_i h_i(T_1) h_i(T_2) \\ &= \sum_i \left( \sum_{n_1 \in N_1} I_i(n_1) \right) * \left( \sum_{n_2 \in N_2} I_i(n_2) \right) \\ &= \sum_i \sum_{n_1 \in N_1} \sum_{n_2 \in N_2} I_i(n_1) I_i(n_2) \\ &= \sum_{n_1 \in N_1} \sum_{n_2 \in N_2} \left( \sum_i I_i(n_1) I_i(n_2) \right) \\ &= \sum_{n_1 \in N_1} \sum_{n_2 \in N_2} C(n_1, n_2) \end{aligned}$$

$K(T_1, T_2) = h(T_1) \cdot h(T_2)$  can be calculated in  $O(|N_1||N_2|)$



# Properties of this kernel

- The value of  $K(T_1, T_2)$  depends greatly on the size of the trees  $T_1$  and  $T_2$ .

$$K'(T_1, T_2) = \frac{K(T_1, T_2)}{\sqrt{K(T_1, T_1)K(T_2, T_2)}}$$

- $K(T, T)$  could be huge. The output would be dominated by the most similar tree.

=> The model would behave like a nearest neighbor rule

# Down-weighting the contribution of large subtrees when calculating $C(n1, n2)$

If the productions at  $n1$  and  $n2$  are different

then  $C(n1, n2) = 0$

else if  $n1$  and  $n2$  are pre-terminals

then  $C(n_1, n_2) = \lambda$

else  $C(n_1, n_2) = \lambda \prod_{j=1}^{nc(n1)} (1 + C(ch(n1, j), ch(n2, j)))$

# Experimental results

# Experiment setting

- Data:
  - Training data: 800 sentences,
  - Dev set: 200 sentences
  - Test set: 336 sentences
  - For each sentence, 100 candidate parse trees
- Learner: voted perceptron
- Evaluation measure: 10 runs and report the average parse score
- Baseline (with PCFG): 74% (labeled f-score)

# Results

Depth	1	2	3	4	5	6
Score	$73 \pm 1$	$79 \pm 1$	$80 \pm 1$	$79 \pm 1$	$79 \pm 1$	$78 \pm 0.01$
Improvement	$-1 \pm 4$	$20 \pm 6$	$23 \pm 3$	$21 \pm 4$	$19 \pm 4$	$18 \pm 3$

With different max subtree size

Scale	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9
Score	$77 \pm 1$	$78 \pm 1$	$79 \pm 1$	$79 \pm 1$	$79 \pm 1$	$79 \pm 1$	$79 \pm 1$	$79 \pm 1$	$78 \pm 1$
Imp.	$11 \pm 6$	$17 \pm 5$	$20 \pm 4$	$21 \pm 3$	$21 \pm 4$	$22 \pm 4$	$21 \pm 4$	$19 \pm 4$	$17 \pm 5$

# Summary

- Show how to use a SVM or a perceptron learner for the reranking task.
- Define a tree kernel that can be calculated in polynomial time.
  - Note: the number of features is infinite.
- The reranker improves parse score from 74% to 80%.