Structure Prediction

Part 0:

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

Structure Prediction

- 1 Overview
- 2 Structured Inputs
 - Recap: Encoding Sequences. RNN, CNN, Transformers
 - **Encoding Graph**
 - RNN vs GNN
 - Permutation equivariance
 - **GNN Variants**
- Structured Output
 - Probabilistic Models of Structures
 - Directed Acyclic Graphs
 - Algorithms for paths in DAGs: Maximization, Probabilities, Sampling
 - Application: Sequence Tagging

Machine Learning



Understanding, choosing, designing:

- models
- learning algorithms
- evaluation metrics
- experiment methodology

to learn and evaluate mappings from inputs x to outputs y.

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Understanding, choosing, designing:

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... for Structures



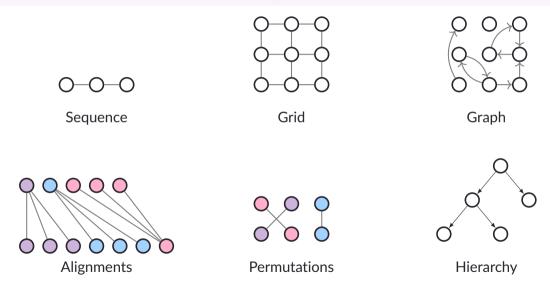
structure, noun: the way in which a complex object's parts are organized in relationship to one another.

Many objects we want to do ML on have interesting structure:

language, images, shapes, networks...

This lecture: how to work with <u>structure</u> in the <u>input</u> and the <u>output</u>.

A few examples of structure



Structures in NLP

- **Sequence** of (sub)words/characters: the usual way we encode linguistic data.
- **Segmentation** into entities / events / sections / speakers / ...
- Inter-word dependencies: syntactic or semantic analysis (graphs, trees)
- Alignment: between multi-lingual documents / speech to phonemes / ...

Structure is at the heart of all models and algorithms designed for NLP.

Recap: ML classifiers

Learn to map from inputs $x \in \mathcal{X}$ to corresponding outputs $y \in \mathcal{Y}$ given a set of training pairs (x, y).

Classification:
$$\mathcal{Y} = \{1, 2, \dots, K\}$$
.

Feature encoder $\phi : \mathcal{X} \to \mathbb{R}^d$. (could be hand-crafted or a neural net)

To make predictions:

$$\hat{y}(x) = \underset{y \in \mathcal{Y}}{\operatorname{argmax}} \, \boldsymbol{w}_y \cdot \boldsymbol{\phi}(x)$$

Another way to think of this:

weight matrix W with rows w_1, \ldots, w_k :

 $\mathbf{a}(x) = \mathbf{W} \boldsymbol{\phi}(x) \in \mathbb{R}^K$ is a vector of scores for each of the k classes

$$score(y; x) = [\mathbf{a}(x)]_y$$

The highest-scoring class wins:

$$\hat{y}(x) = \underset{y \in \mathcal{Y}}{\operatorname{argmax}} \operatorname{score}(y; x)$$

Recap: Probabilistic Classifiers, Logistic Regression

We can give a probabilistic interpretation to the ML classifier by interpreting scores as probabilities by applying <u>softmax</u>:

$$\Pr(y \mid x) = \frac{\exp(\mathsf{score}(y; x))}{Z}, \quad \text{where} \quad Z = \sum_{y \in \mathcal{Y}} \exp(\mathsf{score}(y; x)).$$

$$y \quad 1 \quad 2 \quad 3 \quad 4$$

$$\mathsf{score}(y; x) \quad -1.5 \quad 0.2 \quad 0.9 \quad -1.1$$

$$\Pr(y \mid x) \quad 0.05 \quad 0.29 \quad 0.58 \quad 0.08$$

This motivates logistic regression as a training objective (loss): train params to maximize $\sum_{(x,y)\in\mathcal{D}}\log \Pr(y\mid x)$.

Why is softmax the way it is:

exp ensures all probabilities are non-negative.

Z is the normalizing constant to ensure probabilities sum to 1.

Handling structures

We made no assumptions about the form of $x \in \mathcal{X}$:

this is abstracted into the feature encoder $\phi(x)$.

In the next part (30min), we cover feature encoders for structured objects.

Afterward, we will look at structured outputs \mathcal{Y} .

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Encoding Sequences

You've already seen several methods for encoding $\underline{\text{text}}$ as a sequence of discrete tokens. We recap.

Sequence input: Bag-of-words

Simple but powerful idea: for each vocabulary item, a feature that counts it:

 $\phi_i(x)$ = number of occurrences of word v_i in x.

This leads to:

		\.		Pook	Fairly	good	15	long	nt	the	this
	text	ϕ_1	ϕ_1	ϕ_2	ϕ_3	ϕ_4	ϕ_5	ϕ_6	ϕ_7	ϕ_8	ϕ_9
x_1	"this book is good!"	1	0	1	0	1	1	0	0	0	1
<i>x</i> ₂	"fairly long book"	0	0	1	1	0	0	1	0	0	0
<i>X</i> 3	"the book isn't good."	0	1	1	0	1	1	0	1	1	0

Variants: zero-one, normalized frequencies.

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Variants: zero-one, normalized frequencies.

Order is lost: ϕ ("doesn't word order matter") = ϕ ("word order doesn't matter")

Sequence inputs: Getting some structure back

Sequential order = a fundamental structure of language.

n-grams: treat *n* consecutive tokens as a single one.

Bigram tokenization:

"the book isn't good." \rightarrow [the_book, book_is, is_n't, n't_good, good_.]

This captures some local order.

Can even combine: 1-gram \cup 2-gram $\cup ... \cup n$ -gram: ¹

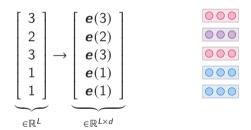
But, it comes at a cost: how many features are needed?

¹Ensure combination is reversible or else we won't be able to distinguish features. For instance, here, _ must not appear in any unigram.

Embeddings of Discrete Tokens

Neural networks perform continuous operations.

For sequential **discrete** data, (language, DNA, etc), we must first represent each token as a continuous "embedding" vector.



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$$\begin{bmatrix}
3 \\
2 \\
3 \\
1 \\
1
\end{bmatrix}
\rightarrow
\begin{bmatrix}
e(3) \\
e(2) \\
e(3) \\
e(1) \\
e(1)
\end{bmatrix}$$

$$\in \mathbb{R}^{L} \times d$$

The function e(i) retrieves the *i*th row from an *embedding matrix* $\mathbf{E} \in \mathbb{R}^{|V| \times d}$.

The embeddings could be fixed or learned as model parameters.

Continuous Bag Of Words

Different-length sequences can be encoded by pooling their embeddings.

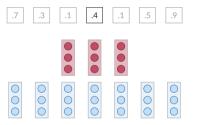


- average pooling: $z = \frac{1}{L}(z_1 + ... + z_L)$
- max pooling: $[z]_j = \max([z_1]_j, ..., [z_L]_j$ (coordinate-wise)

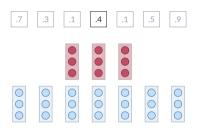
Just like in the standard bag of words, word order doesn't matter.

aka 1-d convolution with d channels

Denote L=sequence length,
 d=embedding size, k=window size.

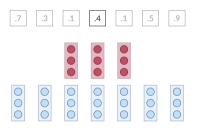


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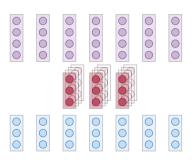
- Denote L=sequence length,
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- In the single-channel case, a filter was a dim-k vector. Now, a filter is a $d \times k$ matrix.

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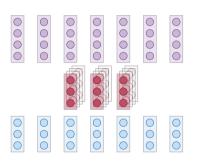
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- Apply m filters in parallel: output is a dim-m vector per window:
 a "layer" maps (L, d) → (L, m), for any L.

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- Apply m filters in parallel: output is a dim-m vector per window:
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- Kind of like "continuous" n-grams!

Recurrent Neural Networks (RNN)

Recurrently encoding a sequence of input vectors $(x_1, \ldots, x_n) \to (z_1, \ldots, z_n)$:

$$\mathbf{z}_{t} = \boldsymbol{\phi}(\mathbf{x}_{t}, \mathbf{z}_{t-1})$$

$$\mathbf{z}_{0} \quad \mathbf{z}_{1} \quad \mathbf{z}_{2} \quad \mathbf{z}_{3} \quad \dots$$

$$(hidden states)$$

$$\mathbf{x}_{1} \quad \mathbf{x}_{2} \quad \mathbf{x}_{3} \quad \dots$$

The simplest RNN is the Elman RNN:

$$z_t = \sigma \left(\underbrace{wx_t}_{\text{lin. func. of inputs}} + \underbrace{vz_{t-1}}_{\text{lin. func. of prev. state}} + b \right)$$

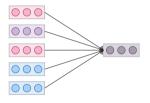
Each hidden state depends on the previous ones. Therefore, cannot parallelize, must compute in order $z_1, z_2, ...$

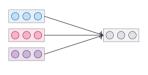
The initial state z_0 is a fixed parameter.

The final state z_n has seen the entire sequence.

Let's talk about pooling.

$$z = \mathsf{AveragePool}(z_1, \dots, z_n) := \frac{1}{n} \sum_{j=1}^n z_j$$



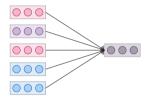


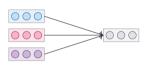
Used to get one representation of a variable-size set or sequence.

Combine n input vectors into one single output vector, with equal contribution.

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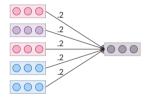
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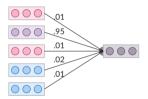
Combine n input vectors into one single output vector, with equal contribution.

But what if some of the inputs should contribute more than others?

Weighted Average Pooling

$$z = \sum_{i} \alpha_{i} z_{i}$$
, where $\alpha_{i} \geq 0$, $\sum_{i} \alpha_{i} = 1$

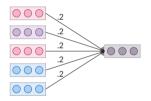


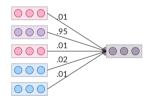


The weights α control the relative importance of the inputs.

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The weights α control the relative importance of the inputs.

But how to come up with these weights? How to decide what's important in a given context?

Attention

Key idea: have a representation of the "context" as a vector $\mathbf{q} \in \mathbb{R}^d$.

Then, say the importance of z_i is proportional to its alignment (\sim angle) to q:

$$\alpha_{i} = \underbrace{\frac{\exp(\boldsymbol{q} \cdot \boldsymbol{z}_{i})}{\sum_{j} \exp(\boldsymbol{q} \cdot \boldsymbol{z}_{j})}}_{[\text{softmax}([\boldsymbol{q} \cdot \boldsymbol{z}_{1}, ..., \boldsymbol{q} \cdot \boldsymbol{z}_{n}])]_{i}}; \qquad \text{Attn}(\boldsymbol{q}; \boldsymbol{z}_{1}, ..., \boldsymbol{z}_{n}) := \sum_{i} \alpha_{i} \boldsymbol{z}_{i}.$$

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This is the basic attention mechanism:

Pool a bunch of vectors, with varying weights, depending on how aligned they are with a context.

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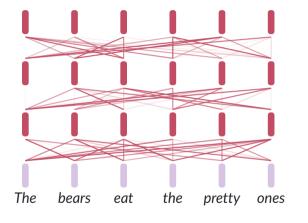
Pool a bunch of vectors, with varying weights, depending on how aligned they are with a context.

What could be the context?

- Could be just a static learned parameter.
- If training on multiple tasks or domains, q can be an embedding of the domain.
- In machine translation (say EN→NL), z_i are the EN words,
 q can be an embedding of the last NL word predicted (one by one).

Transformer

Stacked multi-head attention (+ some annoying details like LayerNorm)



- Combines some of the strengths of CNN and RNN:
- Global even without much depth: every output depends on every input.
- Parallelizable: each position and each head can be computed separately. (still one layer at a time)
- Sequence-aware thanks to positional embeddings.

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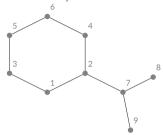
Application: Sequence Segmentation

Encoding general graphs

Graph-structured data: proteins, molecules, social networks, etc.

A graph G = (V, E):

- $V = \{1, ..., n\}$ is the set of nodes.
- $E \subseteq V \times V$ are the edges, e.g., $(u, v) \in E$ means an edge from u to v
- Directed vs undirected graphs: in a nutshell, undirected means
 (u, v) ∈ E ← (v, u) ∈ E.
- the adjacenty matrix $\mathbf{A} \in \{0, 1\}^{n \times n}$ encodes the set of edges E: $a_{uv} = 1 \iff (u, v) \in E$.



Each node can have a *type* (e.g., carbon, hydrogen, ...).

For simplicity, we assume all edges are of the same type.

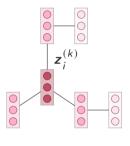
Graph datasets

Two main scenarios, but the tools we use are the same

- **1.** Each data point $x^{(i)}$ is a graph.
 - e.g., molecule solubility, malicious software detection, protein classification, ...
 can be given as a sequence of node labels (x₁⁽ⁱ⁾,...,x_{ni}⁽ⁱ⁾)
 - can be given as a sequence of node labels $(x_1^{(i)}, \dots, x_{n_i}^{(i)})$ and an adjacency matrix $\mathbf{A}^{(i)}$
 - this is what you have in assignment 1
- 2. Data points are parts of one big graph.
 - e.g., node classification (classifying bots on twitter), link prediction (instagram follow suggestions), community detection, ...
 - much harder to set up experiments, dev set/test set, etc.

Node representations with graph neural nets

Encoding a **graph** of input vectors $(x_1, ..., x_n) \rightarrow (z_1, ..., z_n)$:



- We apply an iterative process.
- At iteration 0, $z_i^{(0)} = x_i$ (the input embedding)
- At each iteration, a node's embedding is updated as a function of the embeddings of its neighbors, i.e., message passing along the edges:

$$\begin{aligned} & \boldsymbol{m}_i^{(k)} = \sum_{j \in N(i)} \boldsymbol{z}_j^{(k)} \\ & \boldsymbol{z}_i^{(k+1)} = \phi \left(\boldsymbol{W}_{\text{self}} \boldsymbol{z}_i^{(k)} + \boldsymbol{W}_{\text{neigh}} \boldsymbol{m}_i^{(k)} + \boldsymbol{b} \right) \end{aligned}$$

Apply this update in parallel for every node, then repeat.

The message received by each node is a sum of its neighbors' embeddings:

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Denote by $Z \in \mathbb{R}^{n \times d}$ the matrix of stacked node embeddings, (n =num. nodes, d =embedding dimension).

The *i*th column of the adjacency matrix a_i encodes the (in-)neighbors of node *i*.

$$\boldsymbol{a}_i^{\mathsf{T}}\boldsymbol{Z} =$$

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$$\boldsymbol{a}_i^{\top} \boldsymbol{Z} = \sum_i a_{ij} \boldsymbol{z}_j = \boldsymbol{m}_i$$

Compute all messages at once:

$$M = A^{\mathsf{T}}Z$$

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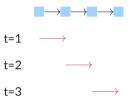
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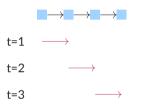
The sequence (chain) graph is also a graph, we could use a GNN.

RNN: sequential updates

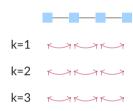


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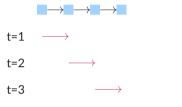


GNN: parallel local updates

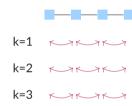


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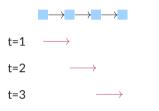
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- Propagates through entire sequence with *L* "messages".
- Embeddings only aware of nodes to the left (without bidirectional "hack")
- Defined for sequences only (some extensions possible).

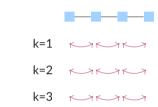
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GNN: parallel local updates



- After k iterations, every node got updates from its neighborhood up to k steps away.
- Can be used for any graph.

Pooling

As defined, a GNN gives us rich embeddings of every node.

To get a single embedding of the entire graph, we turn again to pooling.

Unlike for RNNs, there is no single node that could be taken as representative of the entire graph (especially if k is small and the graph is wide).

We turn to the kind of pooling used for CNNs:

- **1.** average pooling: $z = \frac{1}{n}(z_1 + \ldots + z_n)$
- **2.** max pooling: $[z]_j = \max([z_1]_j, ..., [z_n]_j)$

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Permutation equivariance

The structure of a graph doesn't change if we number the nodes in another order.

The output of a GNN should not change either.

Mathematically, given a graph represented as (X, A), for any permutation matrix P, a GNN satisfies

$$\mathsf{GNN}(\boldsymbol{P}\boldsymbol{X},\boldsymbol{P}\boldsymbol{A}\boldsymbol{P}^{\top}) = \boldsymbol{P}\,\mathsf{GNN}(\boldsymbol{X},\boldsymbol{A}).$$

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Encoding Graphs

RNN vs GNN

Permutation equivariance

GNN Variants

- Structured Outputs
 - Probabilistic Models of Structures
 - Directed Acyclic Graphs

Algorithms for paths in DAGs: Maximization, Probabilities, Sampling

Application: Sequence Tagging

Application: Sequence Segmentation

GNN variants

Many variations can be built on top of this idea.

- The update $z_i^{(k+1)} = \phi(\boldsymbol{W}_{\text{self}} z_i^{(k)} + \boldsymbol{W}_{\text{neigh}} \boldsymbol{m}_i^{(k)} + \boldsymbol{b})$ resembles an RNN. \rightarrow gated variants (GGNN)!
- Separate weight matrices per iteration ($m{W}_{\{ ext{self,neigh}\}}^{(k)}, m{b}^{(k)}$)
- Supporting different edge types:
 - first, notice that $W_{\text{neigh}} \sum_{i} z_{j} = \sum_{i} W_{\text{neigh}} z_{j}$.
 - then, if e(i,j) is the type of the edge from i to j, we could compute $\sum_{j} \mathbf{W}_{e(i,j)} \mathbf{z}_{j}$.
- Different normalization over neighbors (more next time).

Self-Attention for Graphs

Self-attention (and thus Transformers) are permutation equivariant.

Remember in GNN we computed the message from neighbors as a sum:

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Instead, self-attention over neighbors:

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In other words: self-attention constrained by the adjacency structure

(no attention allowed where there is no edge)

Ethylene (
$$C_2H_4$$
): ${}^{\rm H}_{\rm H}$ ${}^{\rm C} = {}^{\rm C} {}^{\rm H}_{\rm H}$



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So far, we've studied this scenario:

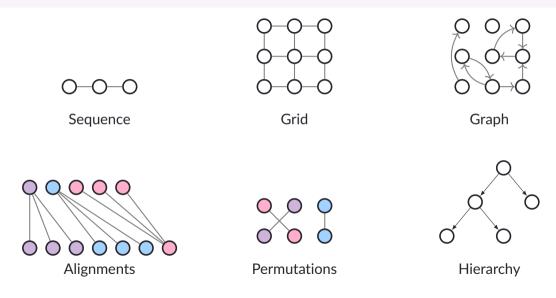
- Structured inputs
- Familiar unstructured outputs: classification / regression.

 $S_0 \longrightarrow \bullet \longrightarrow V$

In the next part of class, we study **structured outputs**.



Reminder: Kinds of Structure



- discrete objects
- made of smaller parts
- which interact with each other and/or constrain each other.

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Example: What are the possible ways to assign 4 jockeys to 4 horses?

$$\mathcal{Y} = \{(1, 2, 3, 4), (1, 2, 4, 3), (1, 3, 2, 4), \dots, (4, 3, 2, 1)\}$$

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What is $|\mathcal{Y}|$?

Recap: Logistic Regression and Perceptron Losses

The two losses we've seen for multi-class classification: (changing notation slightly)

$$L_{\mathsf{LR}}(y) = -\log \mathsf{Pr}(Y = y | x) \\ = -\operatorname{score}(y) + \log \sum_{y' \in \mathcal{Y}} \exp \left(\operatorname{score}(y') \right) \\ L_{\mathsf{Perc}}(y) \\ = -\operatorname{score}(y) + \max_{y' \in \mathcal{Y}} \operatorname{score}(y')$$

For classification:

- we had $\mathcal{Y} = \{1, 2, ..., K\}$
- the model (linear or NN) outputs a vector \mathbf{a} of scores for each class, so $score(y) = a_y$.

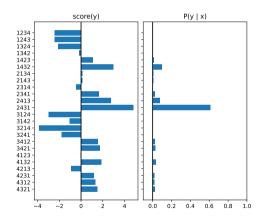
Can we generalize this to structured \mathcal{Y} ?

Probabilistic Models of Structures

Our model must be able to assign a score to every possible structure, $score(y; x, \theta)$. For brevity we just write score(y), but remember it depends on input and params.

From this, we can get a probability distribution over possible structures:

$$Pr(y \mid x) = \frac{\exp(\mathsf{score}(y))}{\sum_{y' \in \mathcal{Y}} \exp(\mathsf{score}(y'))}$$



Modelling challenges

Essential computational prerequisites:

- score(*y*)
- for prediction: $arg max_{y \in \mathcal{Y}} score(y)$
- for learning: $\log \sum_{y \in \mathcal{Y}} \exp (\operatorname{score}(y))$

The challenges: unlike multi-class classification,

- Y can vary for each data point (e.g., with n. horses)
- $|\mathcal{Y}|$ can get very large: we can't just for-loop over it.

Generally intractable!

But, for certain structures and scoring functions, efficient algorithms exist.

The Road Ahead

In the rest of the class, we shall cover a wide range of structured output tasks:

- Sequence labelling
- Sequence segmentation
- Alignments between sequences;
- Assignments and permutations
- Grid / graph labelling

While there is no general-purpose structure prediction algorithm, we shall learn a formalism that will get you far.

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Computations For Structures

Recall: Structured outputs are:

- discrete objects
- made of smaller parts
- which interact with each other and/or constrain each other,

and we must know how to compute:

- score(*y*)
- for prediction: $arg max_{y \in \mathcal{Y}} score(y)$
- for learning: $\log \sum_{y \in \mathcal{Y}} \exp(\operatorname{score}(y))$

For large problems, we can't enumerate ${\cal Y}$ (could be exponentially large).

So, we must actually make use of its structure.

Recap: Graphs

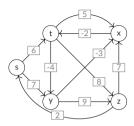
Definition 1: Weighted directed graph

A weighted directed graph is G = (V, E, w) where:

- *V* is the set of vertices (nodes) of *G*.
- $E \subset V \times V$ is the set of arcs of G: $uv \in E$ means there is an arc from node $u \in V$ to node $v \in V$ $(u \neq v)$.

Arcs are ordered pairs, so $uv \neq vu$.

• $w: E \to \mathbb{R}$ is a weight function assigning a weight to each edge.



Recap: Graphs

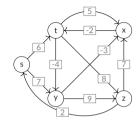
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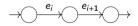
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Definition 2: Paths

A path A in G is a sequence of edges: $A = e_1 e_2 \dots e_k$, with each $e_i \in E$, two-by-two "linked", i.e., if $e_i = u_i v_i$ and $e_{i+1} = u_{i+1} v_{i+1}$ then we must have $v_i = u_{i+1}$.



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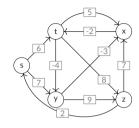
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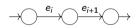
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The weight of a path is the sum of arc weights: $w(A) = \sum_{e \in P} w(e)$.

We denote path concatenation by $A_1 A_2$ (when legal).

Directed Acyclic Graphs

Definition 3: Cycle

A cycle is a path $e_1e_2\dots e_k$ wherein the last edge e_k points to the node from which the first edge e_1 departs.



Directed Acyclic Graphs

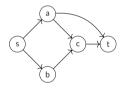
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A DAG is a directed graph that contains no cycles.



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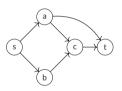
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Definition 4. Topological ordering

A topological ordering of a directed graph G = (V, E) is an ordering of its nodes v_1, v_2, \ldots, v_n such that if $v_i v_i \in E$ then i < j.

G is a DAG if and only if G admits a topological ordering.

Rough intuition: "backward" edges against the ordering \iff cycles.



TOs: s, a, b, c, t s, b, a, c, t

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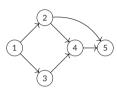
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Paths In DAGs

Label nodes in topological order $V = \{1, ..., n\}$.

Let \mathcal{Y}_i be the set of paths starting at 1 and ending at i.

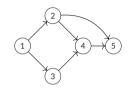


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Let's assume our space of structures is $\mathcal{Y} = \mathcal{Y}_n$.



Important things to compute:

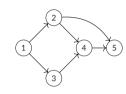
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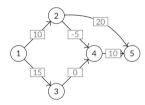
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Later, I'll show you some structured problems that can be usefully reduced to paths in a DAG, and some that cannot.

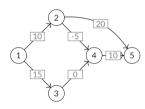
Max-Scoring Path

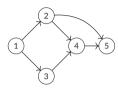
- The greedy path from 1 to 5 might not be best.
- From Data Structures and Algorithms you might recall Dijkstra's algorithm.
 - Requires no "negative cycles" always true for DAGs.
 - Complexity: $\Theta(|V| \log |V| + |E|)$ with "Fibonacci heaps"; $\Theta(|V|^2)$ with a straightforward implementation. .



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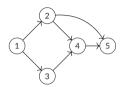
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- In the case of DAGs, we can do better.





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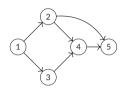
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Insight 1.

Any path from to i is an extension of some path to predecessor $j \in P_i$ by arc ji.

In other words: if $y \in \mathcal{Y}_i$ then $y = y' \hat{j}i$ for some $j \in P_i$ and some $y' \in \mathcal{Y}_j$.



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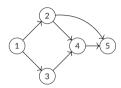
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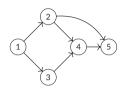
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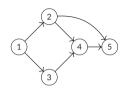
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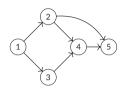
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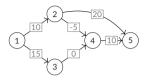
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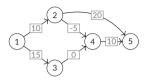
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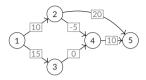


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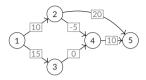
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General Viterbi algorithm for DAGs

input: Topologically-ordered DAG $G = (V, E, w), V = \{1, ..., n\}$ **output:** maximum path weights $m_1, ..., m_n$.

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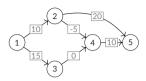
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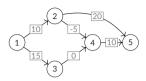
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input: Topologically-ordered DAG $G = (V, E, w), V = \{1, \dots, n\}$ **output:** maximum path weights m_1, \ldots, m_n . initialize $m_1 \leftarrow 0$ for i = 2, ..., n do $m_i \leftarrow \max_{i \in P_i} (m_j + w(ji))$ $\pi_i \leftarrow \arg\max_{i \in P_i} (m_j + w(ji))$ Reconstruct path: follow backpointers **output:** optimal path y from 1 to n (optional) $v = []: i \leftarrow n$ while i > 1 do $v \leftarrow \pi : i \cap v$ $i \leftarrow \pi_i$



 $m_i = \max_{j \in P_i} (m_j + w(ji))$ holds for any graph; but we would chase our own tail forever.

Insight 2.

In a topologically-ordered DAG, any path from 1 to i must only contain nodes i < i.

(So, we may compute m_1, \ldots, m_n in order.)

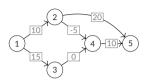
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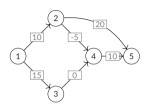
Complexity: $\Theta(|V| + |E|)$.



General Viterbi algorithm for DAGs

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initialize m_1 \leftarrow 0
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   m_i \leftarrow \max_{j \in P_i} (m_j + w(ji))
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Reconstruct path: follow backpointers
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y = []; i \leftarrow n
while i > 1 do
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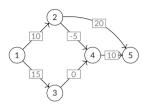


A weighted DAG induces a probability distributions over all paths from 1 to *n*:

$$Pr(y) = \frac{\exp(w(y))}{\sum_{y' \in \mathcal{Y}_n} \exp(w(y'))}$$

У	w(y)	$\exp(w(y))$	Pr(y)
$1 \rightarrow 2 \rightarrow 5$			
$1 \rightarrow 2 \rightarrow 4$	→ 5		
$1 \rightarrow 3 \rightarrow 4$	→ 5		

To assess Pr(y) even for a single path, the denominator sums over all paths.

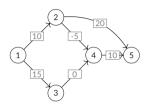


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	10 + 20 = 30 10 - 5 + 10 = 15 15 + 0 + 10 = 25		

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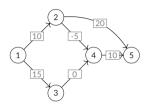


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Since $\exp w(y)$ can be huge, it's better to work with log-probabilities:

$$\log \Pr(y) = w(y) - \log \sum_{y' \in \mathcal{Y}_n} \exp w(y')$$

so we aim to compute this log-sum-exp directly.

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Proposition: DP recurrence for log-sum-exp.

$$q_i = \log \sum_{j \in P_i} \exp(q_j + w(ji))$$

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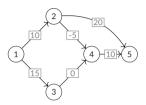
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$$= \log \sum_{j \in P_i} \exp \left(\log \sum_{y' \in \mathcal{Y}_j} \exp(w(y')) + w(ji) \right)$$
$$= \log \sum_{i \in P_i} \exp \left(q_i + w(ji) \right).$$

The Forward Algorithm



General forward algorithm for DAGs

input: Topologically-ordered DAG $G = (V, E, w), V = \{1, ..., n\}$ output: $q_n := \log \sum_{y \in \mathcal{Y}_n} \exp w(y)$.

initialize
$$q_1 \leftarrow 0$$

for $i = 2, ..., n$ do
 $q_i \leftarrow \log \sum_{j \in P_i} \exp(q_j + w(ji))$

Complexity: $\Theta(|V| + |E|)$.

Lets us calculate the log-probability of any given sequence log Pr(y).

Can use autodiff to get $\nabla_w \log \Pr(y)$.



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The pattern:

- $x \oplus y = \max(x, y)$; $x \otimes y = x + y$ form a semiring over $\mathbb{R} \cup \{-\infty\}$.
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This is a very productive generalization that leads to other algorithms too:

- the boolean semiring $x \oplus y = x \lor y$, $x \otimes y = x \land y$ over $\{0, 1\}$ yields an algorithm for path existence;
- there is a semiring that leads to top-k paths.

Sampling Paths

Goal: draw samples from the distribution over paths: $y_1, \ldots, y_k \sim \Pr(Y = y)$. Motivation:

- analyze not just the most likely path, but a set of "typical" paths
- perform inferences

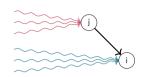
$$\mathbb{E}_{\mathsf{Pr}(Y)}[F(Y)]$$

for arbitrary functions *F*,

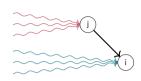
• train structured latent variable models

Probability that the last arc of a path ending in *i* is *ji*:

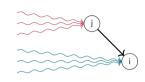
Pr(ji|y ends in i) =



$$\Pr(ji|y \text{ ends in } i) = \frac{\sum_{[y';ji] \in \mathcal{Y}_i} \exp(w(y') + w(ji))}{\sum_{y \in \mathcal{Y}_i} \exp(w(y))}$$



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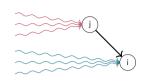


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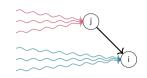
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All paths end in n, so draw the final arc jn first.

Repeat same reasoning on the subgraph with nodes $1, \ldots, j$, i.e., replace n with j and repeat until we hit 1.

Resembles the backpointers from Viterbi: think "stochastic backpointers".

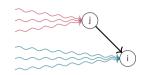
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Forward filtering, backward sampling for DAGs

input: Topologically-ordered DAG; **output:** y: a sample from Pr(y).

$$\begin{aligned} & \text{initialize } q_1 \leftarrow 0 \\ & \textbf{for } i = 2, \dots, n \, \textbf{do} \\ & q_i \leftarrow \log \sum_{j \in P_i} \exp \left(q_j + w(ji) \right) \end{aligned}$$

$$y = []; i \leftarrow n$$

while $i > 1$ do
sample $j \in P_i$ w.p. $p_j = \exp(w(ji) + q_j - q_i)$
 $y \leftarrow ji \cap y$
 $i \leftarrow j$

Conclusions

If we can cast our problem as finding paths in a DAG, then dynamic programming (DP) lets us calculate:

- $\operatorname{argmax}_{y \in \mathcal{Y}} \operatorname{score}(y)$
- $\log \sum_{y \in \mathcal{Y}} \exp \operatorname{score}(y)$ and therefore probabilities
- samples from the distribution over structures

in linear time $\Theta(|V| + |E|)$.

Next we see a bunch of structures that fit this pattern, and some that do not.



Structure Prediction

- 1 Overview
- 2 Structured Inputs

Recap: Encoding Sequences. RNN, CNN, Transformers

Encoding Graphs

RNN vs GNN

Permutation equivariance

GNN Variants

3 Structured Outputs

Probabilistic Models of Structures

Directed Acyclic Graphs

Algorithms for paths in DAGs: Maximization, Probabilities, Sampling

Application: Sequence Tagging

Application: Sequence Segmentation

Given a sequence of *n* items $x = (x_1, ..., x_n)$, assign to each of them one of *K* tags:

$$\mathbf{y} = (y_1, \dots, y_n)$$
 where each $y_i \in \{1, \dots, K\}$.

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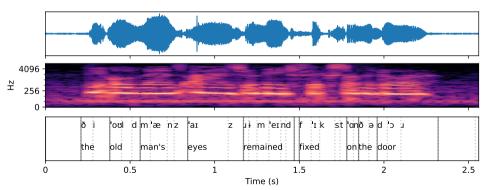
Example 1: Part-of-speech (POS) tagging in NLP

the old man the boat
$$\mathbf{y}_a$$
 det adj noun det noun \mathbf{y}_b det noun verb det noun

Given a sequence of *n* items $\mathbf{x} = (x_1, \dots, x_n)$, assign to each of them one of *K* tags:

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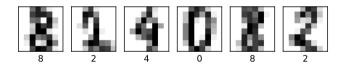
Example 2: Frame-level phoneme classification (may be part of speech recognition)



Given a sequence of *n* items $\mathbf{x} = (x_1, \dots, x_n)$, assign to each of them one of *K* tags:

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 where each $y_i \in \{1, \dots, K\}$.

Example 3: Optical character recognition



Characterizing The Output Space

Given a sequence of *n* items $\mathbf{x} = (x_1, \dots, x_n)$, assign to each of them one of *K* tags:

$$\mathbf{y} = (y_1, \dots, y_n)$$
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Input $\mathbf{x} = (x_1, \dots, x_n)$, e.g., a sequence of words.

Output $y = (y_1, ..., y_n)$, e.g., a sequence of part-of-speech tags.

For each data point (sentence), |y| = |x|; different data points have different lengths.

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For each data point (sentence), |y| = |x|; different data points have different lengths.

For fixed length n, some possible outputs:

- $(1, 1, \ldots, 1, 1) \in \mathcal{Y}$
- $(1, 1, ..., 1, 2) \in \mathcal{Y}$
- $(K, K, \ldots, K, K) \in \mathcal{Y}$

How many in terms of n?

Part-Of-Speech Tags

	Tag	Description	Example
	ADJ	Adjective: noun modifiers describing properties	red, young, awesome
Open Class	ADV	Adverb: verb modifiers of time, place, manner	very, slowly, home, yesterday
C	NOUN	words for persons, places, things, etc.	algorithm, cat, mango, beauty
Sen	VERB	words for actions and processes	draw, provide, go
O	PROPN	Proper noun: name of a person, organization, place, etc	Regina, IBM, Colorado
	INTJ	Interjection: exclamation, greeting, yes/no response, etc.	oh, um, yes, hello
	ADP	Adposition (Preposition/Postposition): marks a noun's	in, on, by, under
S		spacial, temporal, or other relation	
Closed Class Words	AUX	Auxiliary: helping verb marking tense, aspect, mood, etc.,	can, may, should, are
≥	CCONJ	Coordinating Conjunction: joins two phrases/clauses	and, or, but
ass	DET	Determiner: marks noun phrase properties	a, an, the, this
D C	NUM	Numeral	one, two, first, second
sed	PART	Particle: a preposition-like form used together with a verb	up, down, on, off, in, out, at, by
12	PRON	Pronoun: a shorthand for referring to an entity or event	she, who, I, others
	SCONJ	Subordinating Conjunction: joins a main clause with a	that, which
		subordinate clause such as a sentential complement	
er	PUNCT	Punctuation	; , ()
Other	SYM	Symbols like \$ or emoji	\$, %
	X	Other	asdf, qwfg

Figure 8.1 The 17 parts of speech in the Universal Dependencies tagset (Nivre et al., 2016a). Features can be added to make finer-grained distinctions (with properties like number, case, definiteness, and so on).

POS Tagging Evaluation

Evaluation: sequence-level accuracy

$$\frac{\sum_{i=1}^{N_{\text{valid}}} \mathbf{y}^{(i)} = \hat{\mathbf{y}}^{(i)}}{N_{\text{valid}}}$$

or micro-averaged tag accuracy (writing $n^{(i)} = |\mathbf{y}^{(i)}|$):

$$\frac{\sum_{i=1}^{N_{\text{valid}}} \sum_{j=1}^{n^{(i)}} y_j^{(i)} = \hat{y}_j^{(i)}}{\sum_{i=1}^{N_{\text{valid}}} n^{(i)}}$$

Example:

PRO **VERB** NUM NOUN **ADV** true: pred: PRO **VERB** NUM NOUN PRO words: there 70 children there are

true: INTJ pred: X words: eeeeek

Writing
$$\mathbf{y} = (y_1, \dots, y_n)$$
, take $score(\mathbf{y}) = \sum_j a_{j,y_j}$.

	det	noun	adj	verb
the	5	Q	Q	Q
⊿ QId man	Ö	3	3	1
the	5	ŏ	ŏ	Ö
boat	0	5	0	Ο

Writing
$$\mathbf{y} = (y_1, \dots, y_n)$$
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	the	old	man	the	boat
y _a	det	adj	noun	det	noun
\boldsymbol{y}_b	det	noun	verb	det	noun



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$$\mathbf{y} = (y_1, \dots, y_n)$$
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$$\mathbf{y}_a$$
 det adj noun det noun \mathbf{y}_b det noun verb det noun

$$score(\boldsymbol{y}_a) = 21$$

 $score(\boldsymbol{y}_b) =$

Writing
$$\mathbf{y} = (y_1, \dots, y_n)$$
, take $score(\mathbf{y}) = \sum_j a_{j,y_j}$.

$$score(\boldsymbol{y}_a) = 21$$

$$score(\boldsymbol{y}_b) = 17$$

A first attempt: separate classifier for each position.

1. embed and encode x, eg, with a CNN.

$$(x_1,\ldots,x_n)\to(z_1,\ldots,z_n)$$

2. For each position *j*, apply a classification head with *K* outputs. E.g.,

$$\boldsymbol{a}_j = \boldsymbol{W}^\top \boldsymbol{z}_j + \boldsymbol{b}$$

Think of **A** as a matrix with n rows and K columns, where $a_{j,c}$ is the score of assigning tag c at position j.

3. Writing $\mathbf{y} = (y_1, \dots, y_n)$, take score $(\mathbf{y}) = \sum_j a_{j,y_j}$.

```
words = [21, 79, 14] # indices
emb = Embedding(vocab_sz, dim)
clf = Linear(dim, n_tags)
# optionally add RNN, CNN, whatever
Z = emb(words) # (3 \times dim)
A = clf(Z)  # (3 × n_tags)
# computing the score of a given tag sequence:
v = \begin{bmatrix} 2 & 0 & 2 \end{bmatrix}
v_score = sum(A[i. vi]
               for y, yi in enumerate(y))
# or. if you want to be fancy/fast:
y_score = A[torch.arange(len(y)), y].sum()
```

With our score(\mathbf{y}) = $\sum_{j} a_{j,y_{j}}$, can we compute:

 $\max_{\boldsymbol{y} \in \mathcal{Y}} \operatorname{score}(\boldsymbol{y})$

	det	noun	adi	verb
the	5	0	0	0
∡ o <u>l</u> d	Q	1	3	Q
man	Ď	3	Ŏ	1
boat	Ö	5	ŏ	ŏ

With our score(y) = $\sum_i a_{j,y_i}$, can we compute:

$$\max_{\mathbf{y} \in \mathcal{Y}} \operatorname{score}(\mathbf{y})$$

$$= \max_{y_1 \in [K], \dots, y_n \in [K]} \operatorname{score}([y_1, \dots, y_n])$$

	det	noun	adj	verb
the	5	Q	0	0
Old	Ö	1 3	3	0
the	5	ŏ	ŏ	Ö
oat	0	5	0	0

With our score(y) = $\sum_i a_{j,y_i}$, can we compute:

$$\max_{\mathbf{y} \in \mathcal{Y}} \operatorname{score}(\mathbf{y})$$

$$= \max_{y_1 \in [K], \dots, y_n \in [K]} \operatorname{score}([y_1, \dots, y_n])$$

$$= \max_{y_1 \in [K], \dots, y_n \in [K]} \sum_{j} a_{j, y_j}$$

	det	noun	adj	verb
the	5	Q	0	0
old	Ŏ	1	3	Q
nan	Ď	3	Ŏ	1
tne	Š	Ď	Ŏ	Ŏ
oat	U	2	U	U

With our score(\mathbf{y}) = $\sum_{i} a_{i,y_i}$, can we compute:

$$\max_{\mathbf{y} \in \mathcal{Y}} \operatorname{score}(\mathbf{y})$$

$$= \max_{\mathbf{y_1} \in [K], \dots, y_n \in [K]} \operatorname{score}([y_1, \dots, y_n])$$

$$= \max_{\mathbf{y_1} \in [K], \dots, y_n \in [K]} \sum_{j} a_{j, y_j}$$

$$= \sum_{j} \max_{\mathbf{y_j} \in [K]} a_{j, y_j}$$

det noun adjverb the 5 0 0 0 Ald 0 1 3 0 man 0 3 0 1 the 5 0 0 0 booat 0 5 0 0	nan the	5 0 0	0	O	verb 0 0 1 0 0	
---	------------	-------------	---	---	-------------------------------	--

With our score(\mathbf{y}) = $\sum_{i} a_{j,y_i}$, can we compute:

$$\max_{\mathbf{y} \in \mathcal{Y}} \operatorname{score}(\mathbf{y})$$

$$= \max_{\mathbf{y_1} \in [K], \dots, y_n \in [K]} \operatorname{score}([y_1, \dots, y_n])$$

$$= \max_{\mathbf{y_1} \in [K], \dots, y_n \in [K]} \sum_{j} a_{j, y_j}$$

$$= \sum_{j} \max_{\mathbf{y_j} \in [K]} a_{j, y_j}$$

So, $arg max_y score(y)$ is made up of the tags selected independently at each position.

	det	noun	adj	verb	
the old	5 0	0 1	0	0	
man the	<u>0</u>	3	0	1	
oat	0	5	0	0	

With our score(\mathbf{y}) = $\sum_{j} a_{j,y_{j}}$, can we compute:

$$\log \sum_{\pmb{y} \in \mathcal{Y}} \exp\left(\mathsf{score}(\pmb{y})\right)$$

	det	noun	adj	verb
the	5	Q	Q	Q
∡ o <u>ı</u> a man	0	3	0	1
the	5	ŏ	ŏ	Ö
boat	O	5	0	O

With our score(\mathbf{y}) = $\sum_{i} a_{j,y_i}$, can we compute:

$$\log \sum_{\mathbf{y} \in \mathcal{Y}} \exp (\operatorname{score}(\mathbf{y}))$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \exp \sum_{j=1}^n a_{j,y_j}$$

	det	noun	adj	verb
the	5	0	Q	0
nan	ŏ	3	ŏ	1
oat	Ŏ	5	ŏ	ŏ

With our score(\mathbf{y}) = $\sum_{i} a_{i,y_i}$, can we compute:

$$\log \sum_{\mathbf{y} \in \mathcal{Y}} \exp(\mathsf{score}(\mathbf{y}))$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \exp \sum_{j=1}^n a_{j,y_j}$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \prod_{j=1}^n \exp a_{j,y_j}$$

	det	noun	adj	verb	
the	5	Q	Q	Q	
₽ ld man	Ö	3	3	1	
the	5	ŏ	ŏ	Ò	
boat	0	5	0	0	

With our score(\mathbf{y}) = $\sum_{i} a_{i,y_i}$, can we compute:

$$\log \sum_{\mathbf{y} \in \mathcal{Y}} \exp(\operatorname{score}(\mathbf{y}))$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \exp \sum_{j=1}^n a_{j,y_j}$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \prod_{j=1}^n \exp a_{j,y_j}$$

$$= \log \prod_{j=1}^n \sum_{y_j=1}^K \exp a_{j,y_j}$$

	det	nour	adj	verb
the	5	Q	Q	Q
<u>Aol</u> d	Ö	1	3	0
man	5	ŏ	ŏ	Q
boat	Ŏ	5	Ŏ	Ŏ

With our score(\mathbf{y}) = $\sum_{i} a_{j,y_i}$, can we compute:

$$\log \sum_{\mathbf{y} \in \mathcal{Y}} \exp (\operatorname{score}(\mathbf{y}))$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \exp \sum_{j=1}^n a_{j,y_j}$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \prod_{j=1}^n \exp a_{j,y_j}$$

$$= \log \prod_{j=1}^n \sum_{y_j=1}^K \exp a_{j,y_j}$$

$$= \sum_{j=1}^n \log \sum_{y_j=1}^K \exp a_{j,y_j}$$

	det	noun	adj	verb
the	5	0	Q	Q
man	ŏ	3	Ŏ	1
the	5	Q	0	0

Normalizing Constant (log-sum-exp)

With our score(\mathbf{y}) = $\sum_{i} a_{i,V_i}$, can we compute:

$$\log \sum_{\mathbf{y} \in \mathcal{Y}} \exp (\operatorname{score}(\mathbf{y}))$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \exp \sum_{j=1}^n a_{j,y_j}$$

$$= \log \sum_{y_1=1}^K \dots \sum_{y_n=1}^K \prod_{j=1}^n \exp a_{j,y_j}$$

$$= \log \prod_{j=1}^n \sum_{y_j=1}^K \exp a_{j,y_j}$$

$$= \sum_{j=1}^n \log \sum_{y_i=1}^K \exp a_{j,y_j}$$

	det	noun	adj	verb
the	5	Q	Q	Q
<u>⊿ol</u> d man	Ö	1	3	0
the	5	ŏ	ŏ	Ò
oat	0	5	0	0

Probabilistic interpretation: independence

$$\log \Pr(\mathbf{y}) = \operatorname{score}(\mathbf{y}) - \log \sum_{\mathbf{y}' \in \mathcal{Y}} \exp \operatorname{score}(\mathbf{y}')$$
$$= \sum_{j} \underbrace{\left(a_{j,y_{j}} - \log \sum_{k \in [K]} \exp a_{j,k}\right)}_{\log \Pr(y_{j})}$$

For sequence tagging, the separable (fully-local) score

$$\mathsf{score}(\boldsymbol{y}) = \sum_{j} a_{j,y_j}$$

amounts to applying a probabilistic classifier to each of the n positions separately! (any "magic" comes from the feature representation / neural net encoder.)

Can we design a richer score(y) taking into account the sequential structure of y?

Entirely global model: like classification, where each possible sequence is a class.

```
\begin{array}{c} \textbf{y} \ \mathsf{score}(\textbf{y}) \\ \text{det det det det det} & -1000 \\ \text{det det det det noun} & -940 \\ \text{det det det det verb} & -800 \\ \text{det noun verb det noun} & 400 \\ \text{verb verb verb verb} & -1100 \\ \end{array}
```

As expressive as possible: score is any function of the sequence.

Entirely global model: like classification, where each possible sequence is a class.

```
\begin{array}{c} \textbf{y} \ \mathsf{score}(\textbf{y}) \\ \text{det det det det det} \\ \text{det det det noun} \\ \text{det det det det verb} \\ \text{det noun verb det noun} \\ \text{verb verb verb verb} \end{array} \begin{array}{c} \textbf{y} \\ -1000 \\ -800 \\ \text{det noun} \\ \text{verb verb verb} \end{array}
```

As expressive as possible: score is any function of the sequence.

But completely intractable: $O(K^n)$ time and space.

Entirely global model: like classification, where each possible sequence is a class.

```
\begin{array}{c} \textbf{y} \ \mathsf{score}(\textbf{y}) \\ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \\ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{noun} \\ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{det} \ \mathsf{verb} \\ \mathsf{det} \ \mathsf{noun} \ \mathsf{verb} \ \mathsf{det} \ \mathsf{noun} \\ \mathsf{verb} \ \mathsf{verb} \ \mathsf{verb} \ \mathsf{verb} \ \mathsf{verb} \\ \end{array}
```

As expressive as possible: score is any function of the sequence.

But completely intractable: $O(K^n)$ time and space.

Structure output prediction is about the space in between these two extremes.

Idea: scoring transitions between adjacent tags

score(
$$\mathbf{y}$$
) = $\sum_{j=1}^{n} a_{j,y_j} + \sum_{j=2}^{n} t_{y_{j-1},y_j}$

For example, score([NOUN, DET, VERB]) = $+a_{2,DET}a_{1,NOUN} + a_{3,VERB} + t_{NOUN,DET} + t_{DET,VERB}$

Scoring Transitions Between Tags

A rich scorer that takes into account the sequential nature of y while still allowing efficient computation:

scoring transitions between adjacent tags

score(
$$\mathbf{y}$$
) = $\sum_{j=1}^{n} a_{j,y_j} + \sum_{j=2}^{n} t_{y_{j-1},y_j}$

For example, score([NOUN, DET, VERB]) = $a_{1,NOUN} + a_{2,DET} + a_{3,VERB} + t_{NOUN,DET} + t_{DET,VERB}$

Sequence Modeling With Transition Scores

score(
$$\mathbf{y}$$
) = $\sum_{j=1}^{n} a_{j,y_j} + \sum_{j=2}^{n} t_{y_{j-1},y_j}$

The tag scores $\mathbf{A} \in \mathbb{R}^{n \times K}$ can be computed as before (e.g., with a convnet.)

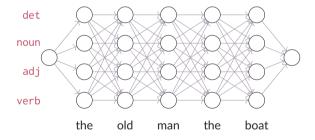
The transition scores $T \in \mathbb{R}^{K \times K}$:

- could be a learned parameter. (size does not depend on *n*)
- could be predicted by the neural net as a function of x.

Unlike in the separable case, with transition scores, we no longer get n parallel classifiers: the different tags impact one another. (This makes the model more expressive and more interesting.)

Sequence Tagging As A DAG

$$score(y) = \sum_{j=1}^{n} a_{j,y_j} + \sum_{j=2}^{n} t_{y_{j-1},y_j}$$



$$G = (V, E, w) \text{ where:}$$

$$V = \{(j, c) : j \in [n], c \in [K]\}$$

$$\cup \{s, t\}$$

$$E = \{(j - 1, c') \rightarrow (j, c) : j \in [2, n], c, c' \in [K]\}$$

$$\cup \{s \rightarrow (1, c) : c \in [K]\}$$

$$\cup \{(n, c) \rightarrow t : c \in [K]\}$$

$$w((j - 1, c') \rightarrow (j, c)) = a_{j,c} + t_{c',c}$$

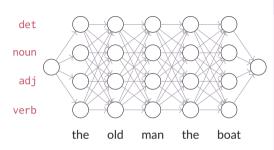
$$w(s \rightarrow (1, c)) = a_{1,c}$$

$$w((n, c) \rightarrow t) = 0$$

$$|V| \in \Theta(nK); \quad |E| \in \Theta(nK^2)$$

Topological ordering?

Viterbi For Sequence Tagging



General Viterbi (reminder sketch)

initialize
$$m_1 \leftarrow 0$$

for $i = 2, ..., n$ do
 $m_i \leftarrow \max_{j \in P_i} (m_j + w(ji))$
 $\pi_i \leftarrow \arg\max_{j \in P_i} (m_j + w(ji))$
follow backpointers to get best path

Viterbi for sequence tagging

input: Unary scores A ($n \times K$ array) Transition scores T ($K \times K$ array)

Forward: compute scores recursively $m_{1c} = a_{1c}$ for all $c \in [K]$ for j = 2 to n do for c = 1 to K do $m_{j,c} \leftarrow \max_{c' \in [K]} \left(m_{j-1,c'} + a_{j,c} + t_{c',c} \right)$ $\pi_{j,c} \leftarrow \arg\max_{c' \in [K]} \left(m_{j-1,c'} + a_{j,c} + t_{c',c} \right)$ $f^* = \max_{c' \in [K]} m_{n,c'}$

Backward: follow backpointers

$$y_n = \arg \max_{c'} m_n(c')$$

for $j = n - 1$ down to 1 do
 $y_j = \pi_{j+1, y_{j+1}}$

output: f^* and $y^* = [y_1, \dots, y_n]$

 $m_{i,c}$ is stored as a matrix M, same shape as A.

Apply $m_{1,c} = a_{1,c}$ to get the first row: (copied from **A**)

Then iteratively: $m_{j,c} = \max_{c' \in [K]} m_{j-1,c'} + a_{j,c} + t_{c',c}$

At the end, take the maximum over the last row.

det noun adj verb the សុុ៤ nan

boat

 $m_{i,c}$ is stored as a matrix M, same shape as A.

Apply $m_{1,c} = a_{1,c}$ to get the first row: (copied from **A**)

Then iteratively: $m_{j,c} = \max_{c' \in [K]} m_{j-1,c'} + a_{j,c} + t_{c',c}$

At the end, take the maximum over the last row.

```
the 5 0 0 0 0 mid the boat
```

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At the end, take the maximum over the last row.





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At the end, take the maximum over the last row.

```
the 5 0 0 0 mid 1 man the boat
```

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 $m_{i,c}$ is stored as a matrix M, same shape as A.

Apply $m_{1,c} = a_{1,c}$ to get the first row: (copied from **A**)

Then iteratively: $m_{j,c} = \max_{c' \in [K]} m_{j-1,c'} + a_{j,c} + t_{c',c}$

At the end, take the maximum over the last row.

det noun adj verb the 5 0 0 0 Mild 1 9 10 4 man 8 15 11 12 the 18 13 14 17 boat 18 26 20 17



 $m_{i,c}$ is stored as a matrix M, same shape as A.

Apply $m_{1,c} = a_{1,c}$ to get the first row: (copied from **A**)

Then iteratively: $m_{j,c} = \max_{c' \in [K]} m_{j-1,c'} + a_{j,c} + t_{c',c}$

At the end, take the maximum over the last row.

det noun adj verb the 5 0 0 0 Mild 1 9 10 4 man 8 15 11 12 the 18 13 14 17 boat 18 26 20 17

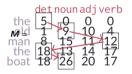
To find the best tag sequence y^* , keep track of the path.

 $m_{i,c}$ is stored as a matrix M, same shape as A.

Apply $m_{1,c} = a_{1,c}$ to get the first row: (copied from **A**)

Then iteratively: $m_{j,c} = \max_{c' \in [K]} m_{j-1,c'} + a_{j,c} + t_{c',c}$

At the end, take the maximum over the last row.



To find the best tag sequence y^* , keep track of the path.

The Two Main Recurrences Of Sequence Tagging:

(Dynamic programming applied to the sequence tagging DAG)

$$\begin{split} m_{j,c} &= \max_{c' \in [K]} & \left(m_{j-1,c'} + a_{jc} + t_{c'c} \right), \\ q_{j,c} &= \log \sum_{c' \in [K]} \exp \left(q_{j-1,c'} + a_{jc} + t_{c'c} \right). \end{split}$$

The Forward Algorithm

Forward algorithm for sequence tagging

```
input: Unary scores A (n \times K array)

Transition scores T (K \times K array)

Forward: compute scores recursively

q_{1,c} = a_{1,c} for all c \in [K]

for j = 2 to n do

for c = 1 to K do

q_{j,c} = \log \sum_{c' \in [K]} \exp \left(q_{j-1,c'} + a_{j,c} + t_{c',c}\right)

return \log Z = \log \sum_{c' \in [K]} \exp \left(q_{n,c'}\right)
```

```
the
                old
                         man
                                   the
                                           boat
       det
                adj
                         noun
                                   det
                                           noun
                                                    score(y_a) = 25
\mathbf{y}_a
       det
                         verb
                                   det
                                           noun
                                                    score(y_b) = 26
y<sub>b</sub>
                noun
                                                      score(y_c) = 1
y<sub>C</sub>
      noun
               noun
                         noun
                                  noun
                                           noun
```

```
the 5.00 0.00 0.00 0.00 φ 1.73 9.00 10.00 4.19 man 8.18 15.01 11.05 12.70 the 18.88 13.92 14.37 17.03 boat 18.08 26.88 20.90 18.38
```

	the	old	man	the	boat	
\mathbf{y}_{a}	det	adj	noun	det	noun	$score(y_a) = 25$
y _b	det	noun	verb	det	noun	$score(y_b) = 26$
\boldsymbol{y}_{c}	noun	noun	noun	noun	noun	$score(y_c) = 1$

 $\log Z \approx 26.885$

	the	old	man	the	boat	
\mathbf{y}_{a}	det	adj	noun	det	noun	$score(y_a) = 25$
y _b	det	noun	verb	det	noun	$score(y_b) = 26$
\boldsymbol{y}_{c}	noun	noun	noun	noun	noun	$score(y_c) = 1$

$$\log Z \approx 26.885$$

$$\log P(y_a) = \text{score}(y_a) - \log Z = 25 - 26.885 = -1.885$$

	the	old	man	the	boat	
\mathbf{y}_{a}	det	adj	noun	det	noun	$score(y_a) = 25$
y _b	det	noun	verb	det	noun	$score(y_b) = 26$
\boldsymbol{y}_{c}	noun	noun	noun	noun	noun	$score(y_c) = 1$

$$\log Z \approx 26.885$$

$$\log P(y_a) = \text{score}(y_a) - \log Z = 25 - 26.885 = -1.885$$

$$\log P(y_b) = \text{score}(y_b) - \log Z = 26 - 26.885 = -0.885$$

	the	old	man	the	boat	
\mathbf{y}_a	det	adj	noun	det	noun	$score(y_a) = 25$
y _b	det	noun	verb	det	noun	$score(y_b) = 26$
\mathbf{y}_c	noun	noun	noun	noun	noun	$score(y_c) = 1$

$$\log Z \approx 26.885$$

$$\log P(y_a) = \text{score}(y_a) - \log Z = 25 - 26.885 = -1.885$$

$$\log P(y_b) = \text{score}(y_b) - \log Z = 26 - 26.885 = -0.885$$

$$\log P(y_c) = \text{score}(y_c) - \log Z = 1 - 26.885 = -25.885$$

Putting It All Together

At this point, we have all the ingredients needed to train a probabilistic sequence tagger with transition scores!

- Receiving an input sequence x, the model returns unary and transition scores A and T.
- **2.** If we're at test time: run Viterbi to get predicted sequence; compute accuracies etc.
- 3. If training time: run Forward algorithm to compute the training objective $-\log P(y \mid x) = -\operatorname{score}(y) + \log \sum_{v' \in \mathcal{Y}} \exp \operatorname{score}(y')$.

This probabilistic model is often known as a Linear-Chain Conditional Random Field.

(Historically, Linear-Chain CRFs didn't use neural net scorers, but the math doesn't change. Today I prefer to teach it this way.)

Structure Prediction

- 1 Overview
- 2 Structured Inputs

Recap: Encoding Sequences. RNN, CNN, Transformers

Encoding Graphs

RNN vs GNN

Permutation equivariance

GNN Variants

3 Structured Outputs

Probabilistic Models of Structures

Directed Acyclic Graphs

Algorithms for paths in DAGs: Maximization, Probabilities, Sampling

Application: Sequence Tagging

Application: Sequence Segmentation

The rod cutting problem: We have a rod of length *n* units, and we can cut it at every marker. What cuts to make to maximize the total value of the resulting pieces?



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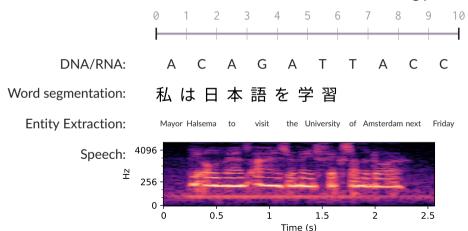
Word segmentation: 私は日本語を学習

DNA/RNA:

The rod cutting problem: We have a rod of length n units, and we can cut it at every marker. What cuts to make to maximize the total value of the resulting pieces?



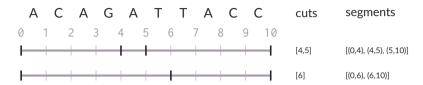
The rod cutting problem: We have a rod of length *n* units, and we can cut it at every marker. What cuts to make to maximize the total value of the resulting pieces?

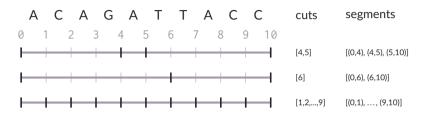


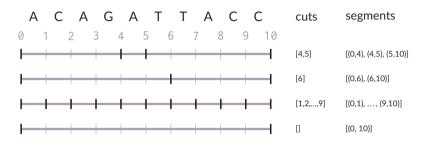
Representing and scoring segmentations

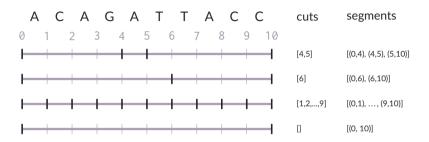


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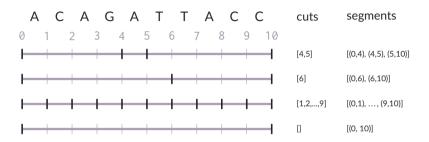




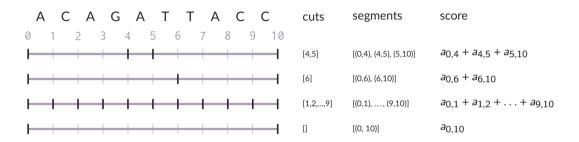




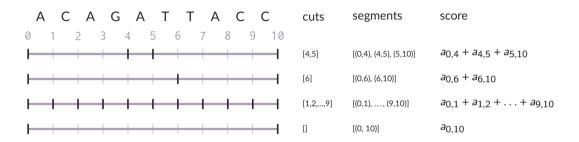
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- How many possible segments?
- How many possible segmentations?

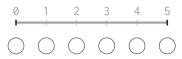


- How many possible segments?
- How many possible segmentations?
- Scoring: assign a score to every possible segment (i, j).

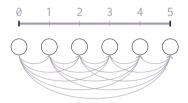


- How many possible segments?
- How many possible segmentations?
- Scoring: assign a score to every possible segment (i, j).
- You can visualize this as the "upper triangle" of a $(n+1) \times (n+1)$ matrix:





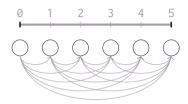
Nodes: one per fencepost. $V = \{0, 1, ..., n\}$.



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Edges: one per segment.

$$E = \{(i,j) : 0 \le i < j \le n\}.$$

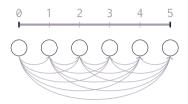


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Topologic order?

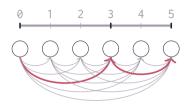


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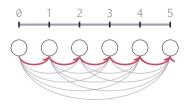


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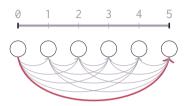


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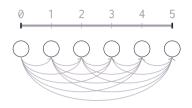


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Topologic order?



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Edges: one per segment. $E = \{(i, j) : 0 \le i < j \le n\}.$

Topologic order?

Any path from 0 to *n* corresponds to a segmentation of the sequence.

Viterbi for segmentation

input: segment scores $\mathbf{A} \in \mathbb{R}^{n \times n}$

Forward: compute recursively $m_1 = a_{01}$; $\pi_1 = 0$ for j = 2 to n do $m_j \leftarrow \max_{0 \le i < j} m_i + a_{ij}$ $\pi_j \leftarrow \arg\max_{0 \le i < j} m_i + a_{ij}$ $f^* = m_n$ Backward: follow backpointers $y^* = []; j \leftarrow n$ while j > 0 do $y^* = [(\pi_j, j)] + y^*$ $j = \pi_j$

Analogously, we can obtain a *Forward* algorithm for log *Z*: exercise for you.

Evaluation



True segments: y = [(0,3), (3,5), (5,6), (6,11)]

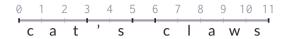
A few possible predictions:

$$\hat{\pmb{y}}_a = [(0,11)]$$

$$\hat{\boldsymbol{y}}_b = [(0,1), (1,2), \dots, (10,11)]$$

$$\hat{\boldsymbol{y}}_c = [(0,3), (3,5), (5,11)]$$

Evaluation



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The number of predicted and true segments differ.

A common way to evaluate in this scenario is:

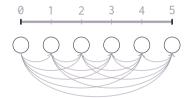
$$precision = \frac{\text{n. correctly predicted segments}}{\text{n. predicted segments}}$$

$$recall = \frac{\text{n. correctly predicted segments}}{\text{n. true segments}}$$

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

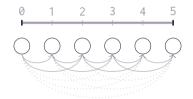
More advanced metrics can partially reward overlaps.

Extension 1: bounded segment length



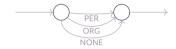
- can be much faster if we limit segment lengts to $L \ll n$.
- in terms of the DAG: discard edges ij where j i > L
- exercise: how does this impact the complexity of Viterbi?

Extension 1: bounded segment length



- can be much faster if we limit segment lengts to $L \ll n$.
- in terms of the DAG: discard edges ij where j i > L
- exercise: how does this impact the complexity of Viterbi?

Extension 2: labeled segments



- each segment also receives a label (e.g., PERSON, ORGANIZATION, NONE...)
- the labels are independent given the cuts: for any two nodes in the DAG, we only need to pick the best edge between them.

Extension 3: labeled + transitions

- drawing inspiration from sequence tagging: what if we want a reward/penalty for consecutive PERSON→ORGANIZATION segments?
- labels no longer independent given cuts.
- still solvable via DP, but must keep track of transitions.
- essentially a combination of the sequence tagging DAG and the segmentation DAG.

Summary

- Segmentations of a length-n sequence: $O(2^n)$ possible segmentations, $O(n^2)$ possible segments.
- Dynamic programming gives polynomial-time probabilistic segmentation models.
- Extensions can accommodate maximum lengths, labels, transitions.