

Analyzing Errors of Unsupervised Learning

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Percy Liang

Dan Klein



Unsupervised grammar induction

Goal: induce hidden syntax

The man ate a tasty sandwich

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DT — NN — VBD — DT — JJ — NN
| | | | | |
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POS tagging

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DT NN VBD

DT

JJ NN

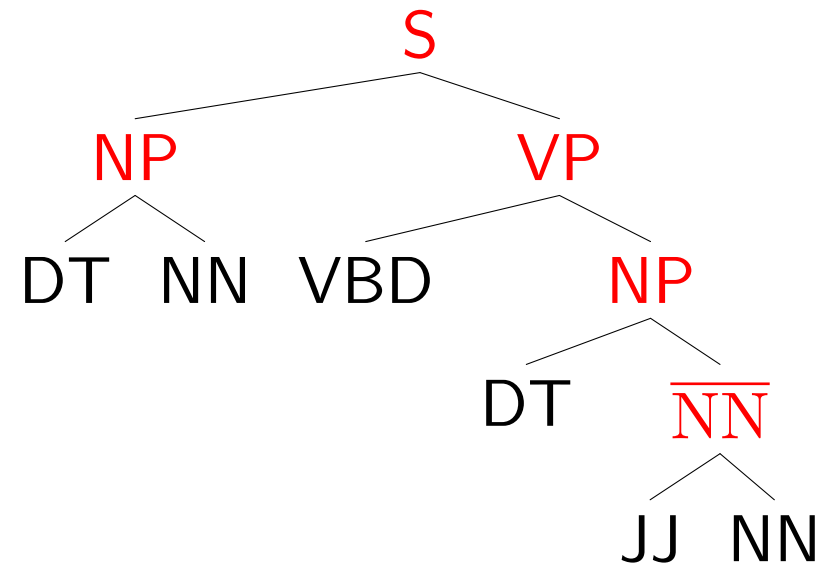
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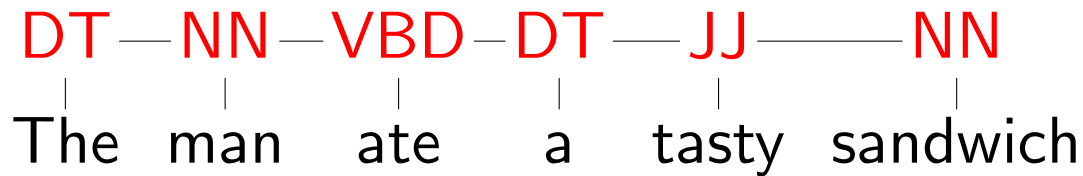
POS tagging



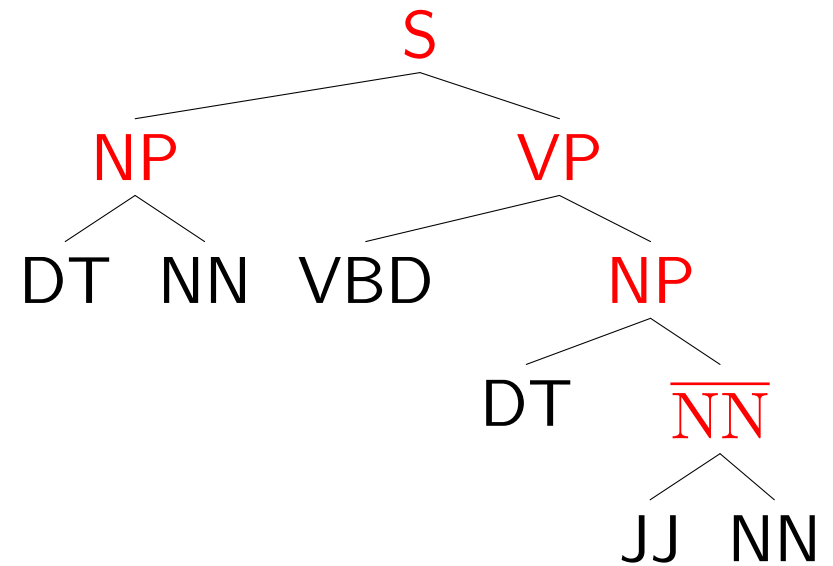
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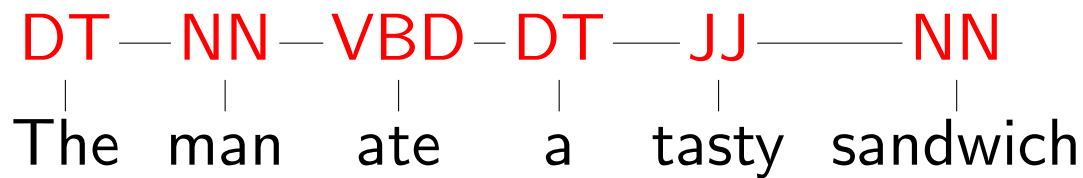
Constituency parsing

For example, on POS tagging using HMMs:

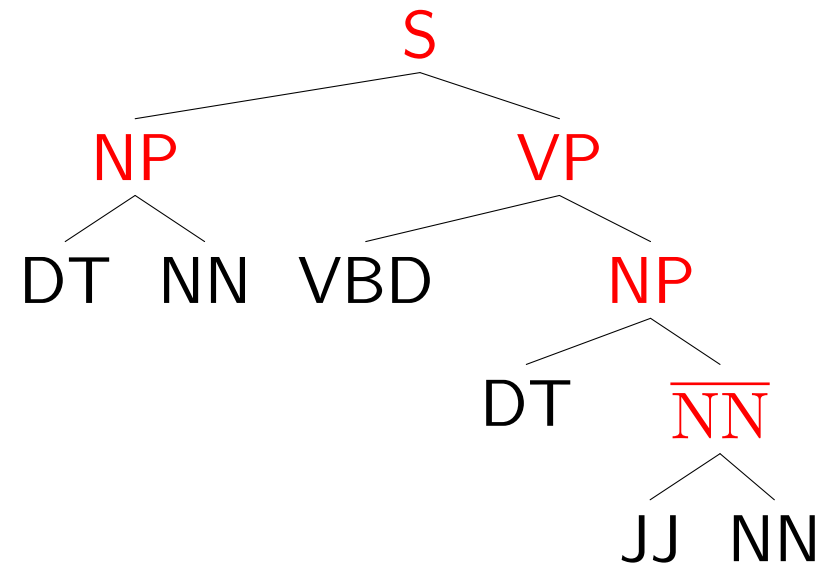
Unsupervised using EM $\approx 60\%$

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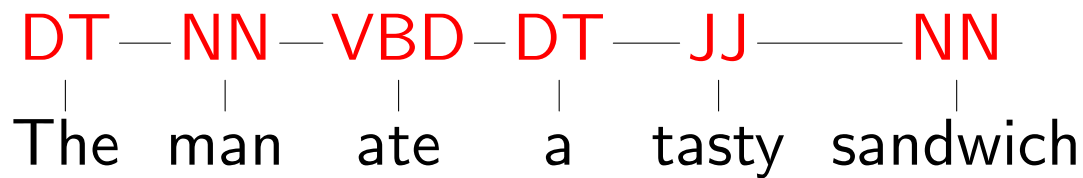
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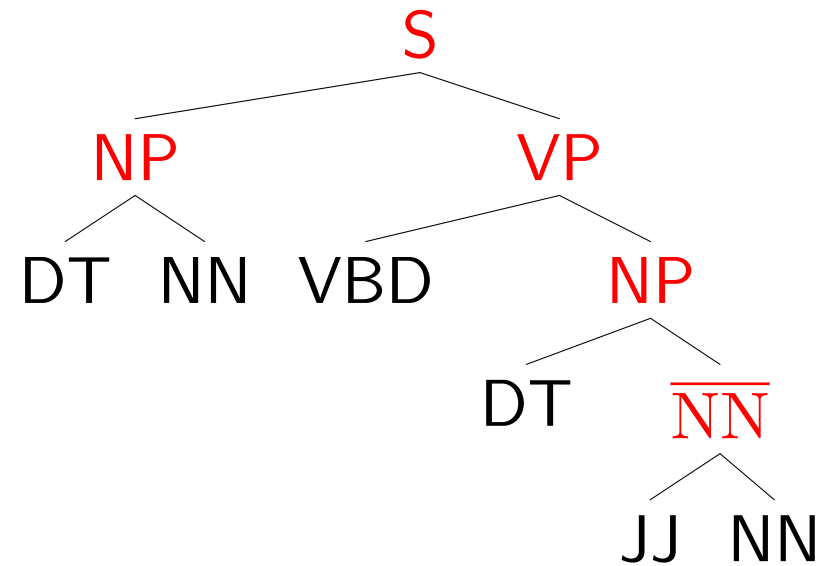
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Unsupervised grammar induction

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POS tagging



Constituency parsing

For example, on POS tagging using HMMs:

Unsupervised using EM $\approx 60\%$

Supervised $\geq 90\%$

Why does EM fail?

Four types of errors:

Four types of errors:

Optimization error

Local optima

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Local optima

Estimation error

Limited data

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Approximation error

Likelihood objective \nrightarrow accuracy

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Likelihood objective $\not\Rightarrow$ accuracy

Identifiability error

Different parameter settings \rightarrow same objective

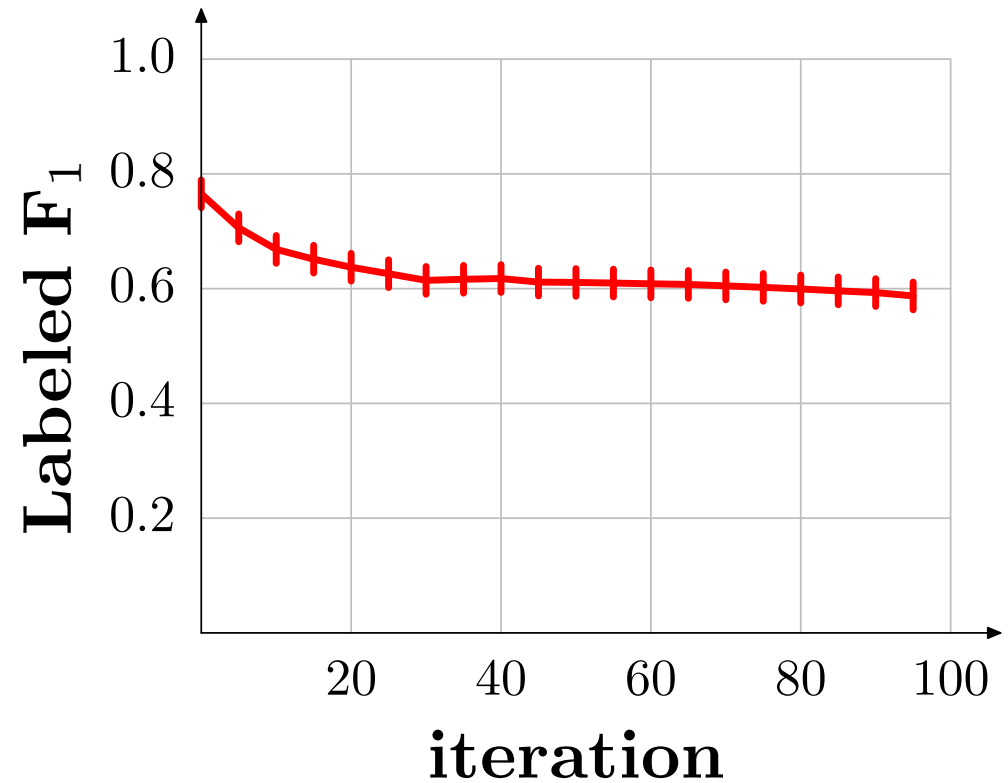
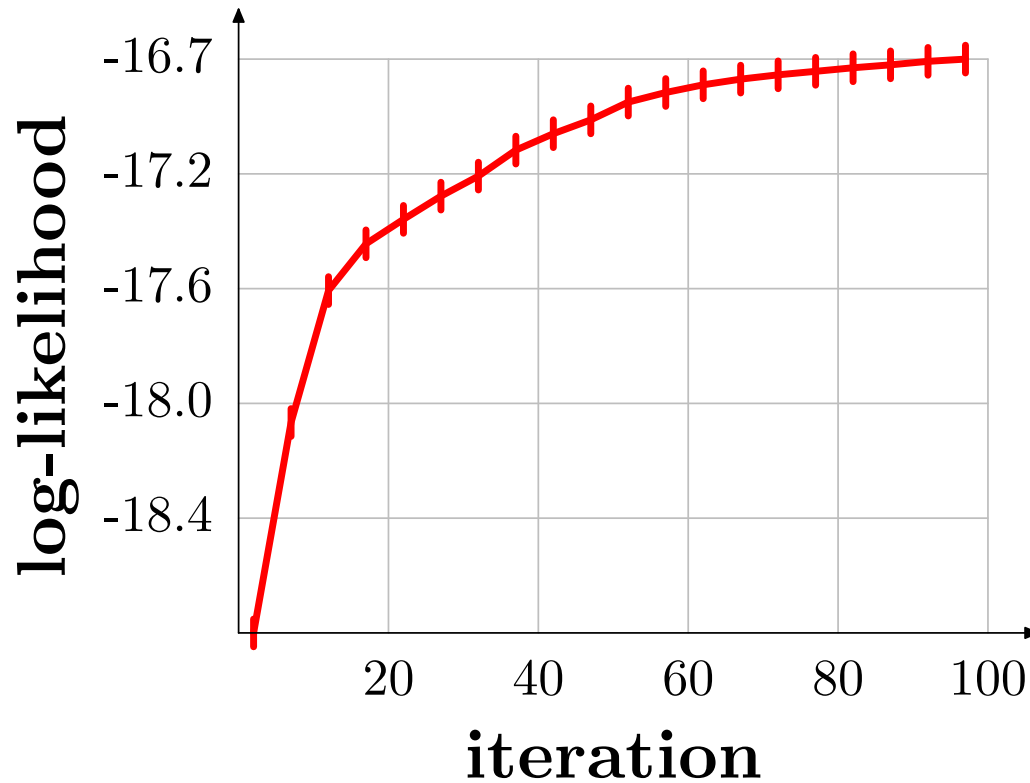
Approximation error

Problem: model likelihood $\not\Rightarrow$ prediction accuracy

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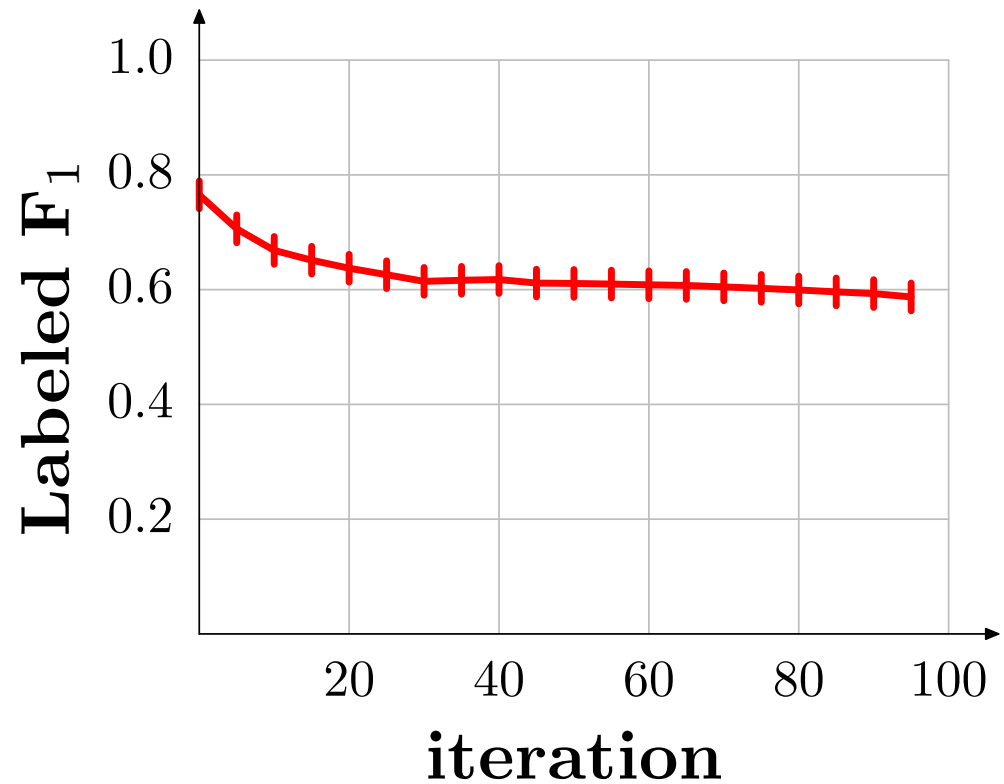
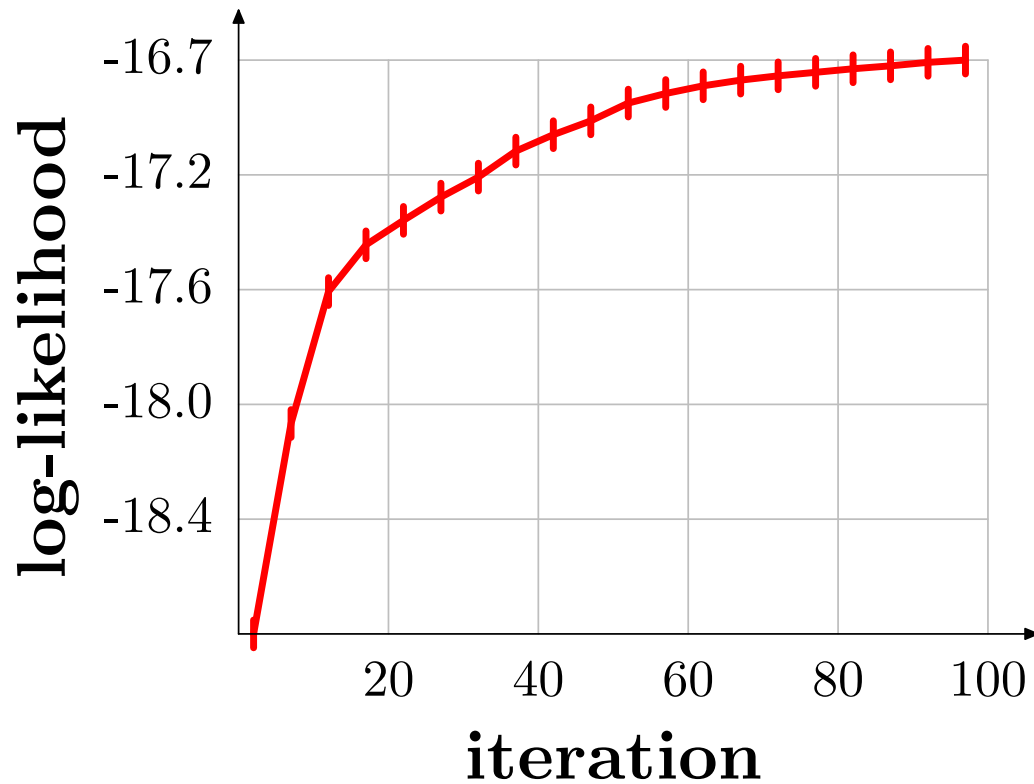
PCFG (EM starting from supervised parameter estimate):



Approximation error

Problem: model likelihood \nrightarrow prediction accuracy

PCFG (EM starting from supervised parameter estimate):



What qualitative changes is EM making?

Migrations

For the HMM:

Truth	DT	NN	NN	RB	VBD	NNS
	The chief executive allegedly made contributions					
				↓		
Iteration 1	DT	JJ	NN	RB	VBN	NNS
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JJ → NN

VBD → made



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NN → NN



JJ → NN

VBD → made



VBN → made

What are the prominent migrations over the entire corpora?

Top HMM migrations

Iteration 1

START → ~~NN~~
NNP

Sentence-initial nouns are often proper
START Revenue/~~NN~~/~~NNP~~ rose

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~~NN~~
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Noun adjuncts → adjectives (inconsistent gold tags)
*chief/~~NN~~/**JJ** executive/**NN** officer*

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Iteration 2

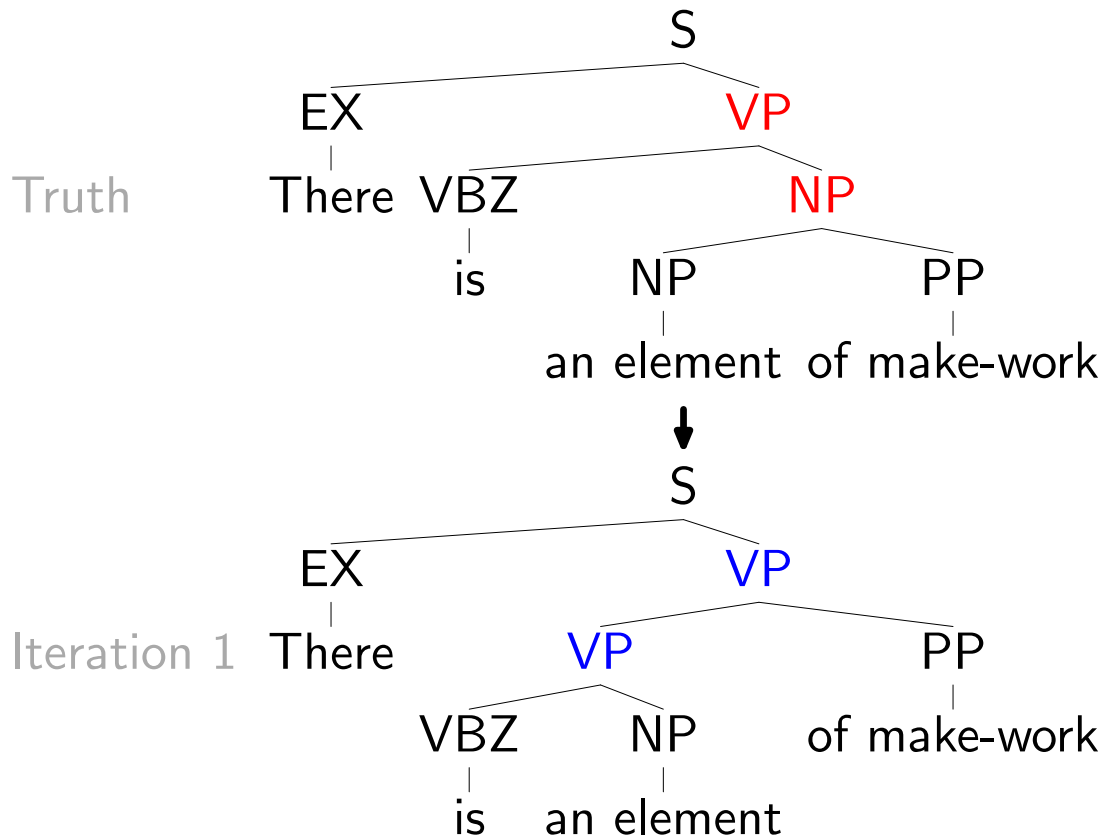
~~NN~~
JJ → NN (same as above)

START → ~~NN~~
NNP (same as above)

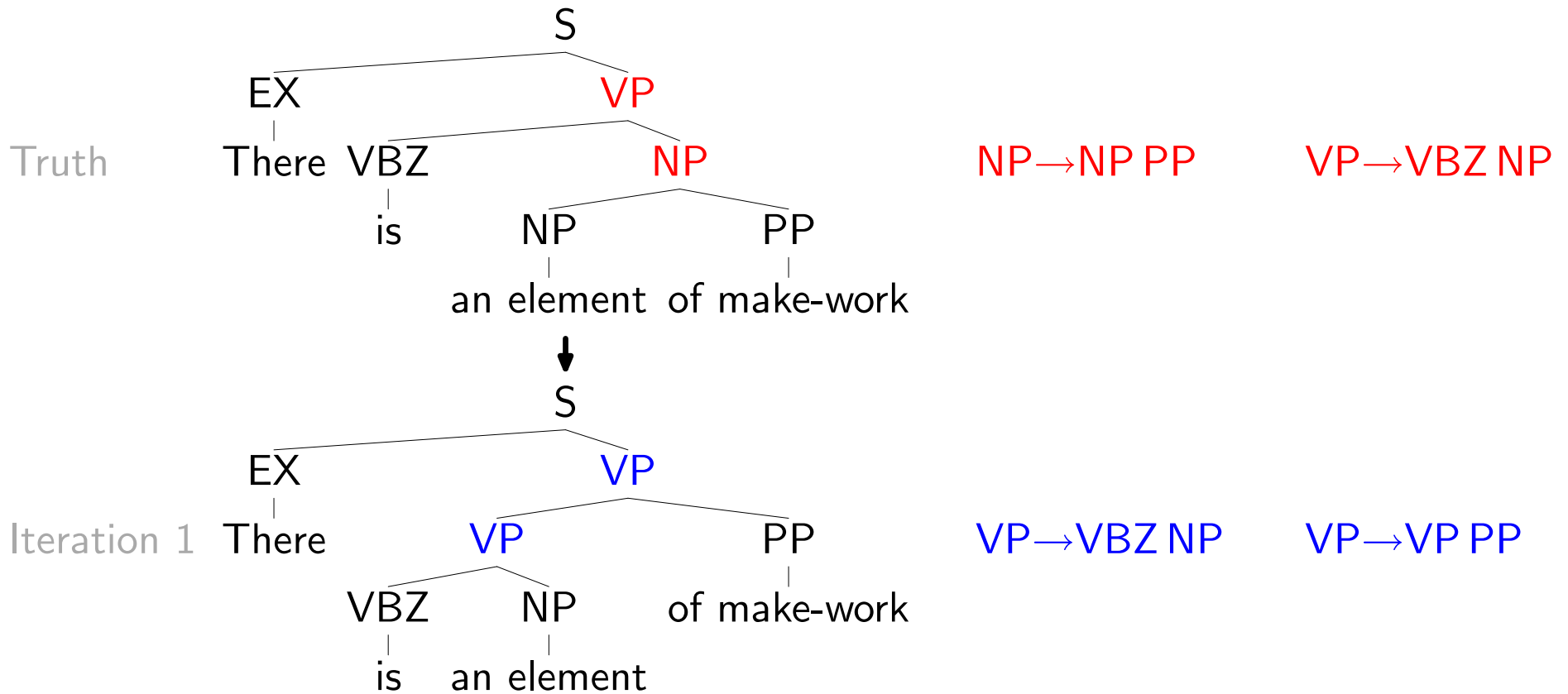
~~JJ~~
RB → TO

Inconsistent gold tags
contribute much/~~JJ~~/~~RB~~ to

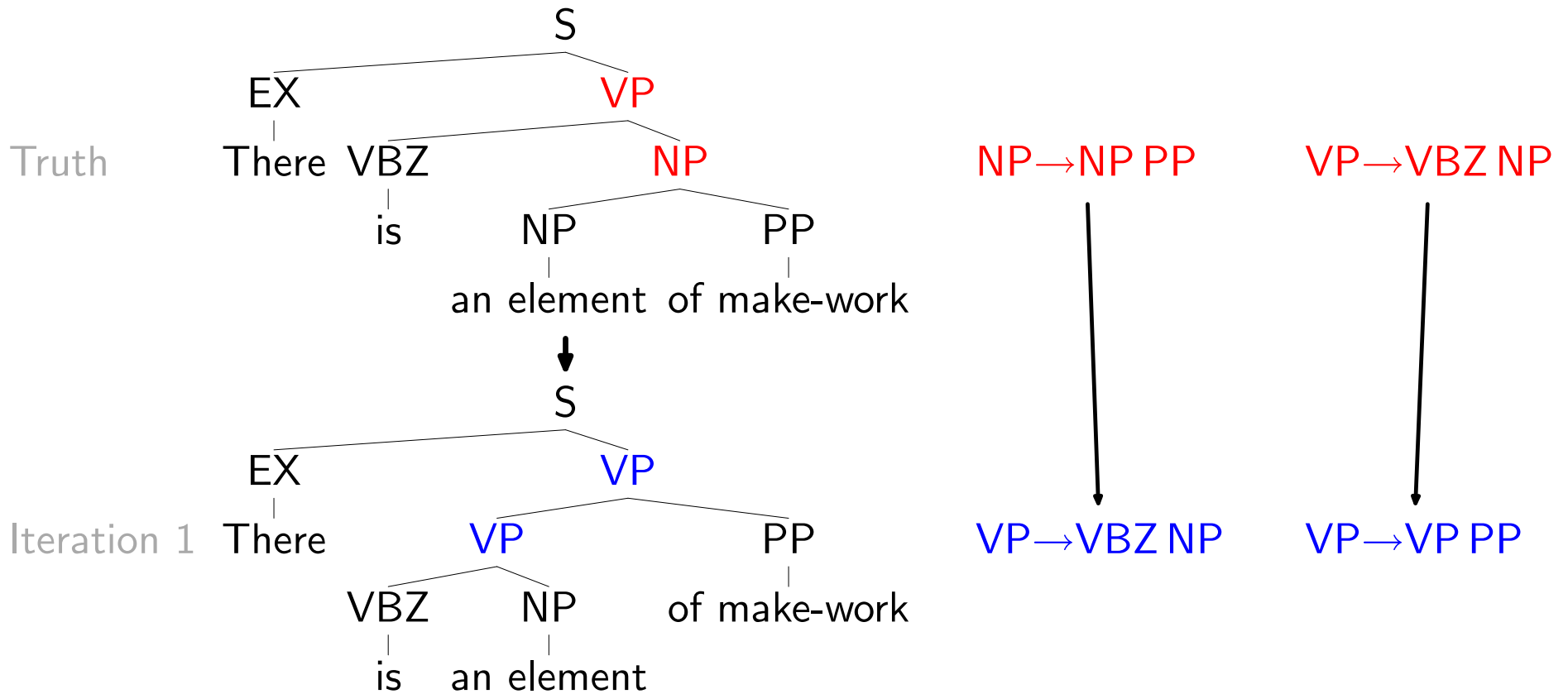
Meta-modeling for PCFGs



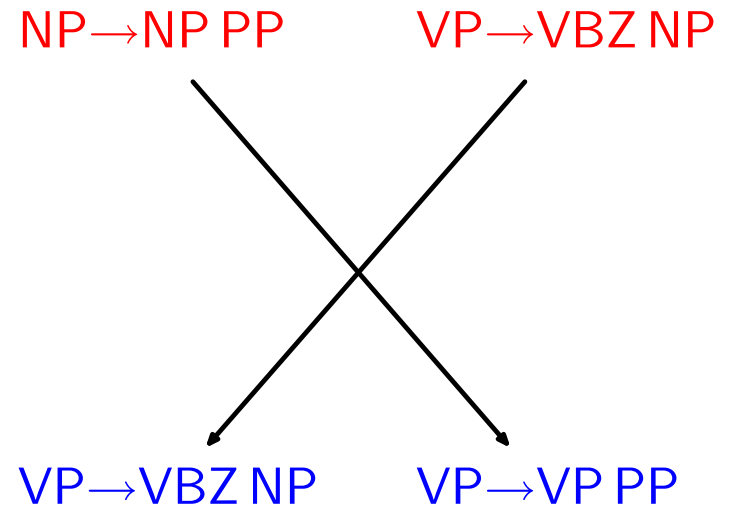
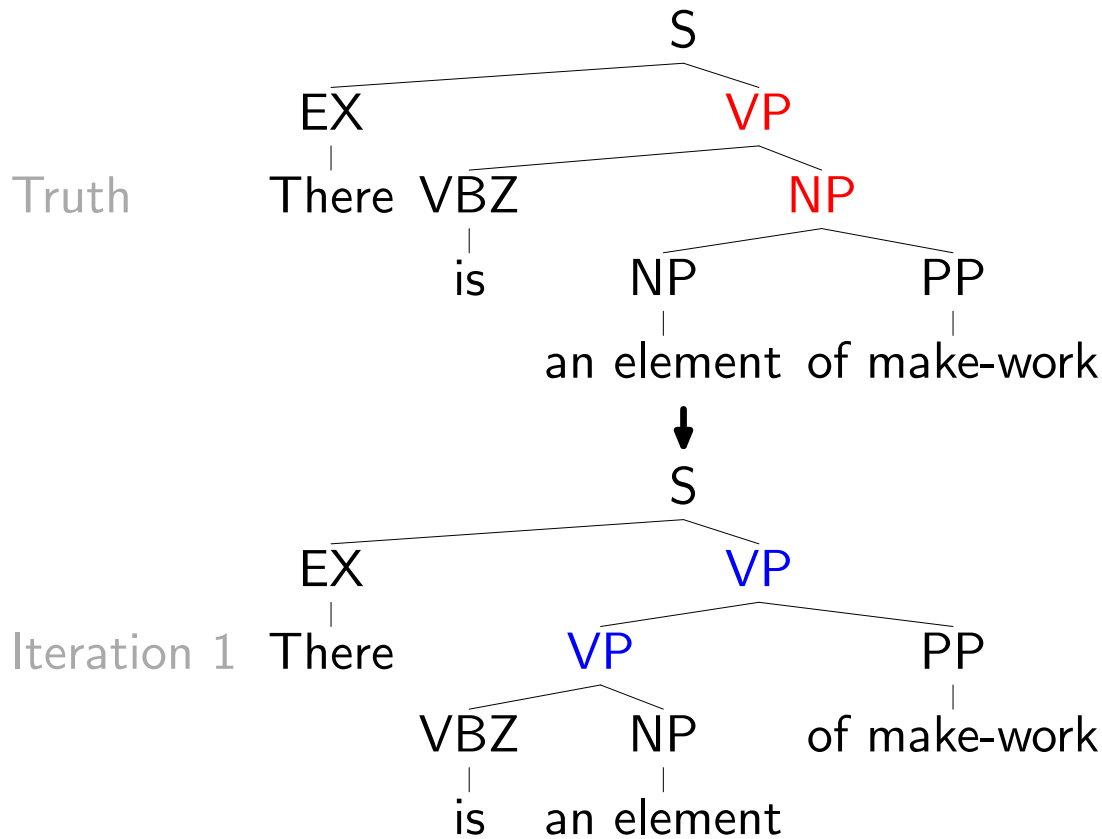
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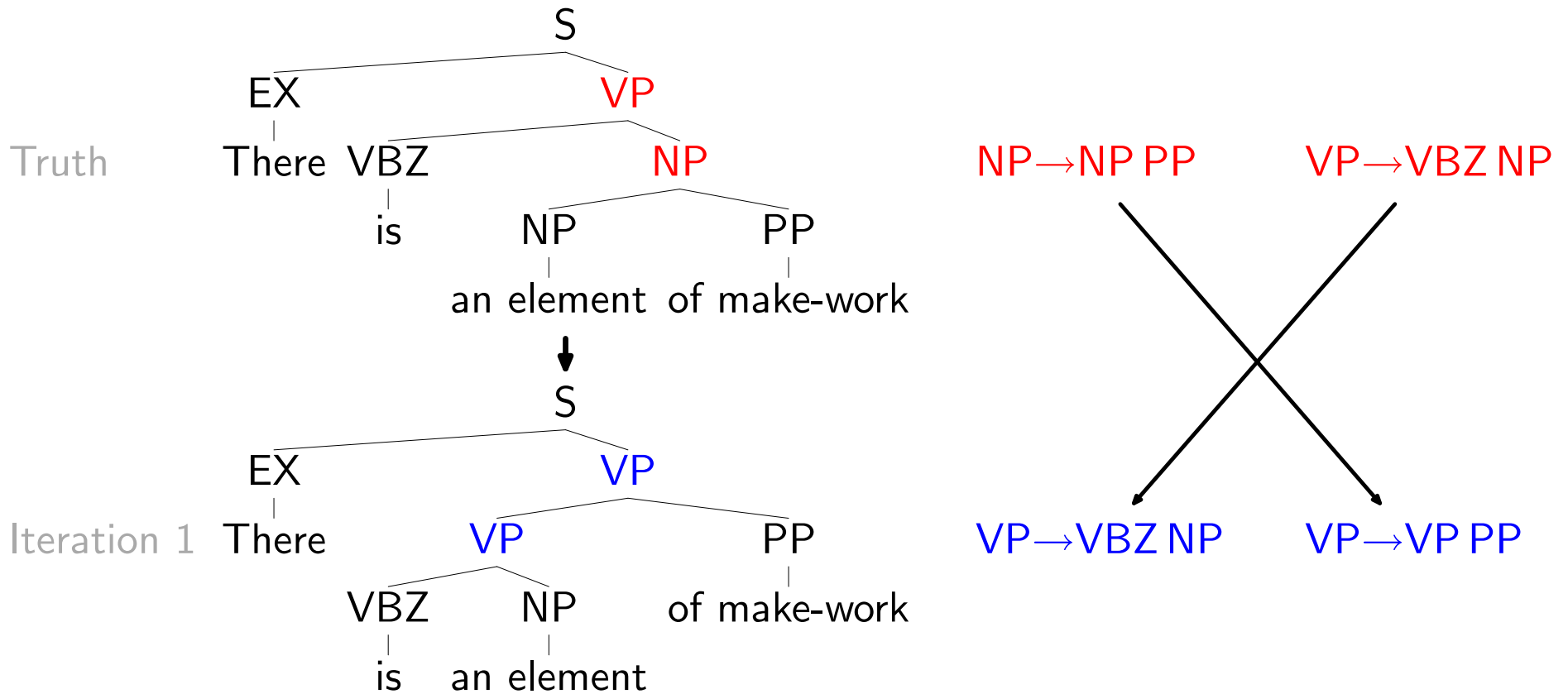


Meta-modeling for PCFGs



Migrations less clear due to uncertainty in tree structure...

Meta-modeling for PCFGs

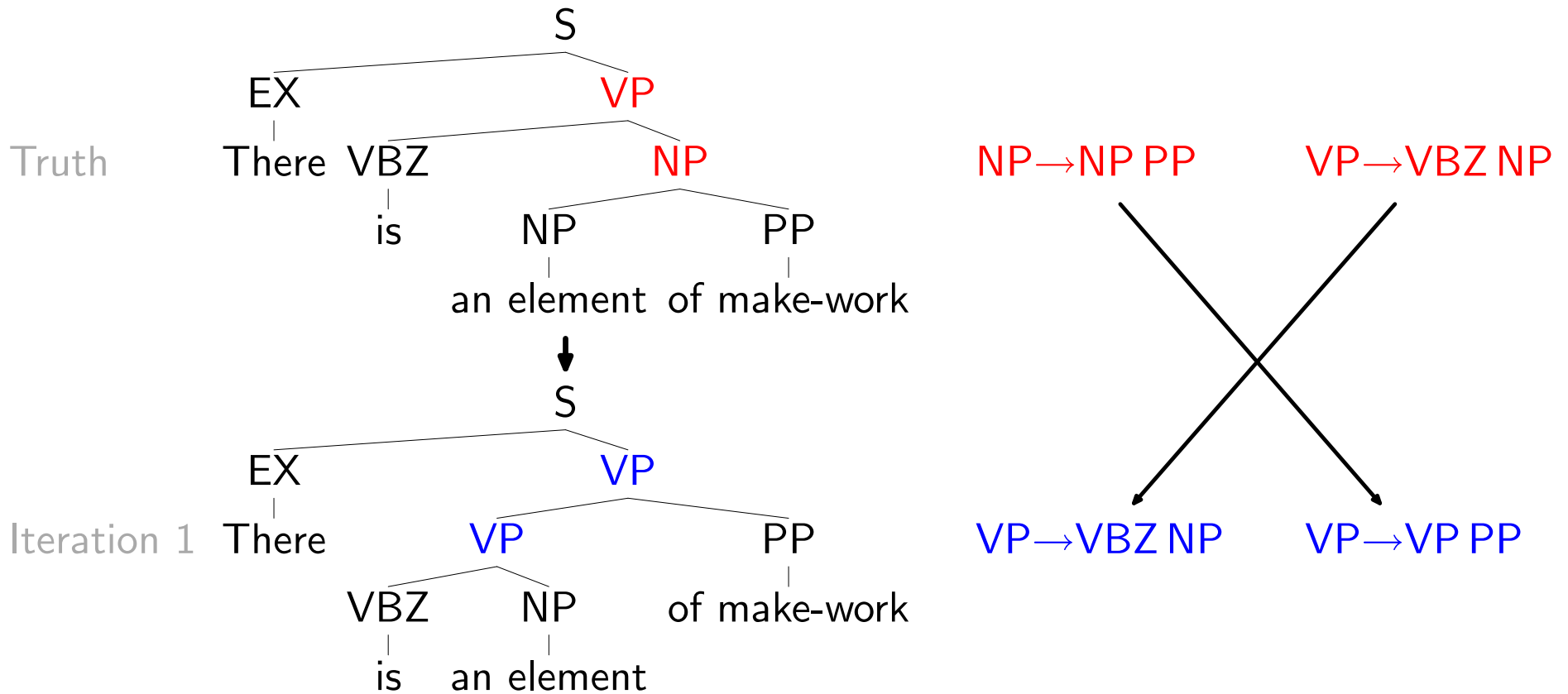


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Our approach: use a meta-model

- Migrations are hidden alignments to be learned
- Fit using EM

Meta-modeling for PCFGs



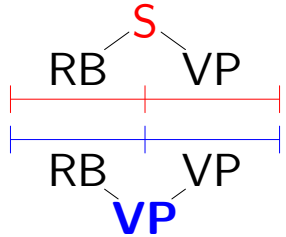
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Top PCFG migrations learned by meta-model

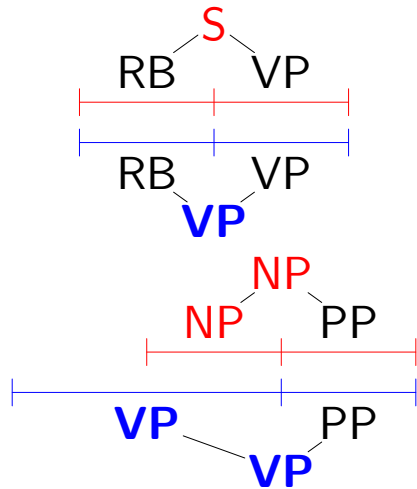
Iteration 1



Sentential adverbs \rightarrow VP adverbs

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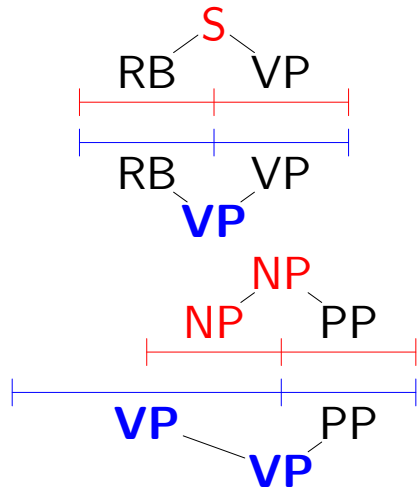


Sentential adverbs → VP adverbs

PPs raised from NPs to verbal level

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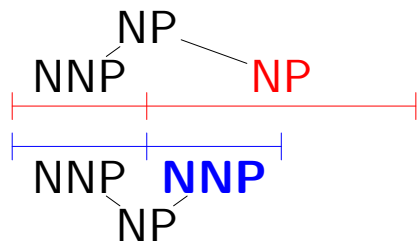
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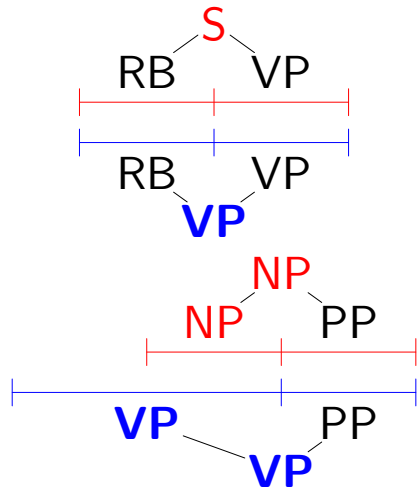
Iteration 2



Right-branching → left-branching structures

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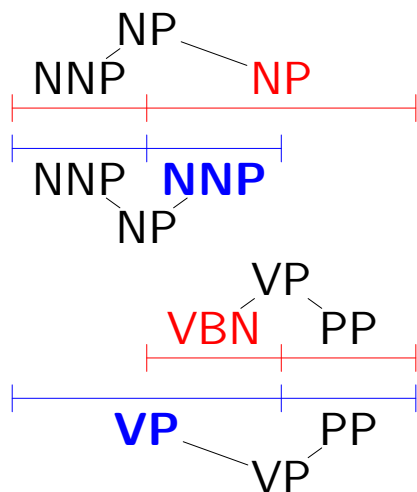
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Sentential adverbs → VP adverbs

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Iteration 2



Right-branching → left-branching structures

PP raised to higher VP

Meta-modeling summary

- Meta-model: a diagnostic tool to analyze errors systematically

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Identifiability error
Estimation error
Optimization error

Identifiability error

\mathbf{x} : input sentence

\mathbf{y} : hidden output

$p_{\theta}(\mathbf{x}, \mathbf{y})$: joint distribution with parameters θ

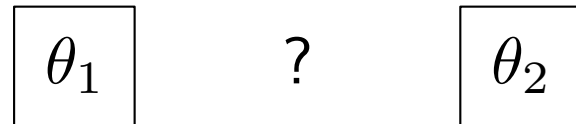
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Learning is indifferent...

$$p_{\theta_1}(\mathbf{x}) = p_{\theta_2}(\mathbf{x})$$
$$\boxed{\theta_1} \quad ? \quad \boxed{\theta_2}$$

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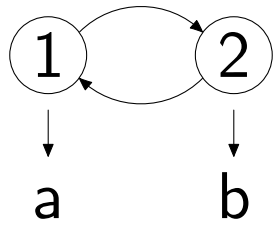
$$\boxed{\theta_1} \quad ? \quad \boxed{\theta_2}$$

but matters to prediction (bad!)

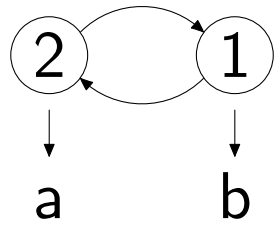
$$p_{\theta_1}(\mathbf{y} | \mathbf{x}) \neq p_{\theta_2}(\mathbf{y} | \mathbf{x})$$

Examples of non-identifiability

- Label symmetries



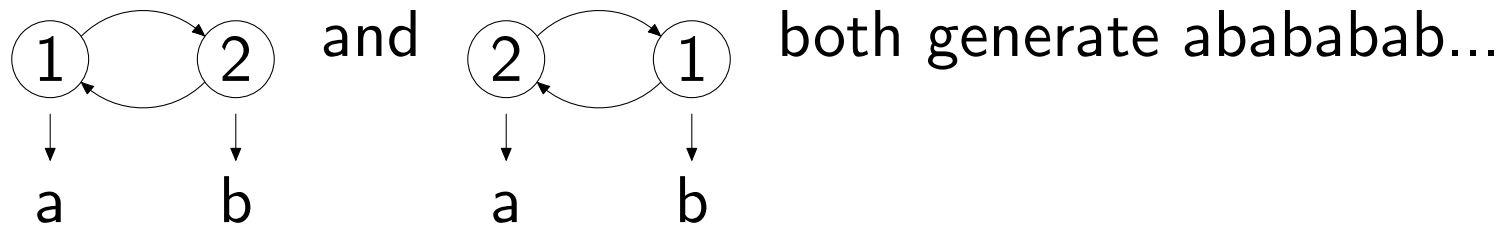
and



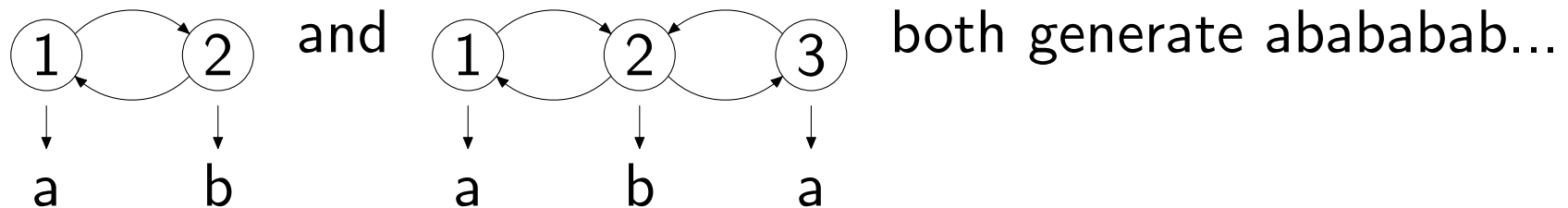
both generate abababab...

Examples of non-identifiability

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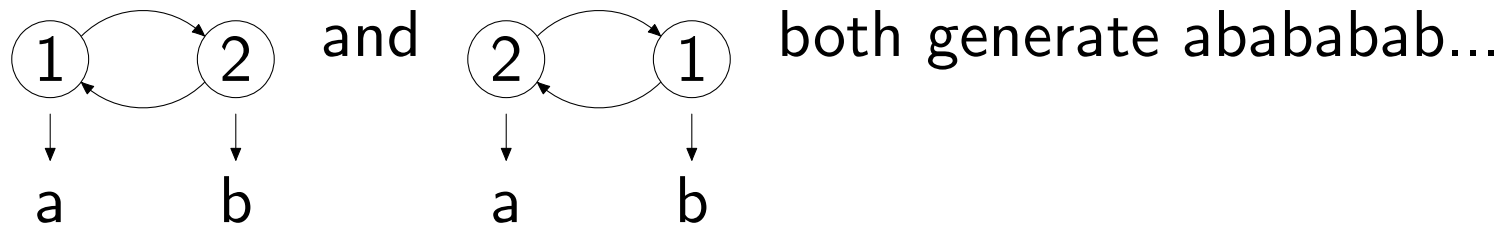


- K -state HMM (if true distribution is $< K$ -state HMM)

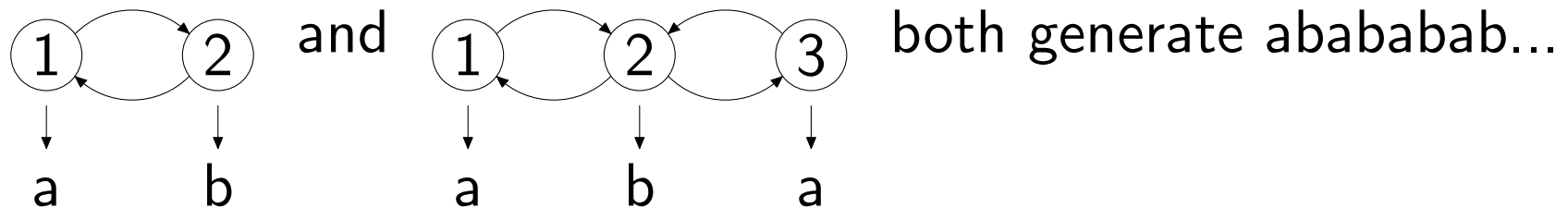


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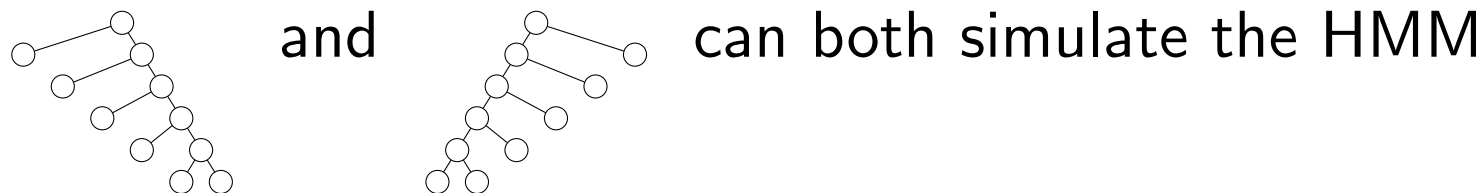
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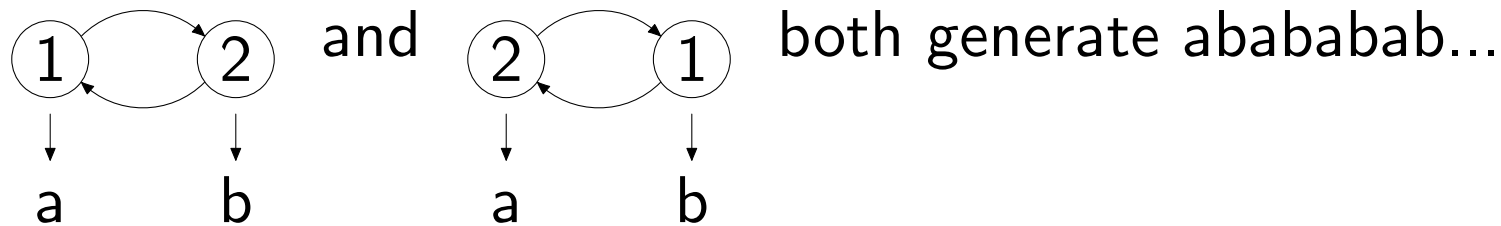


- PCFG (if true distribution is HMM)

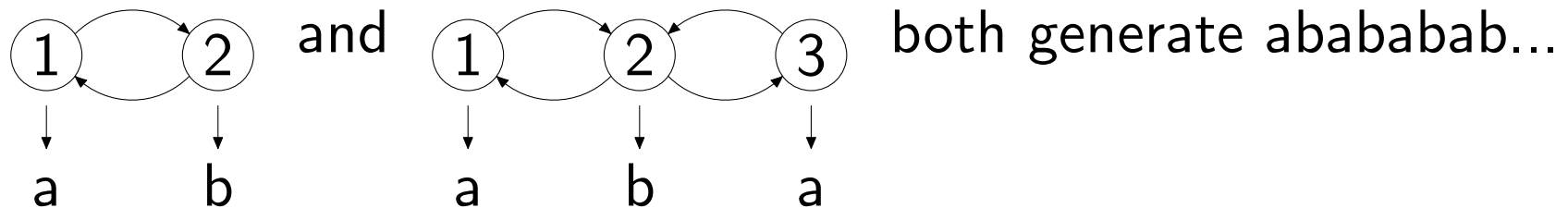


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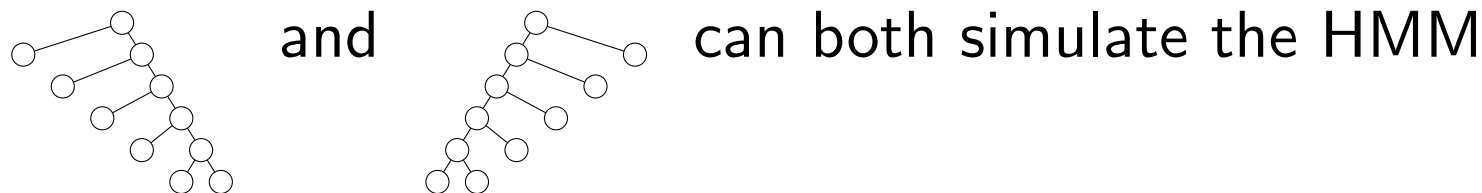
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- PCFG (if true distribution is HMM)



Real data is complex, so last two are not an issue

Identifiability and distance

Given θ_1 and θ_2 , how to measure distance between them?

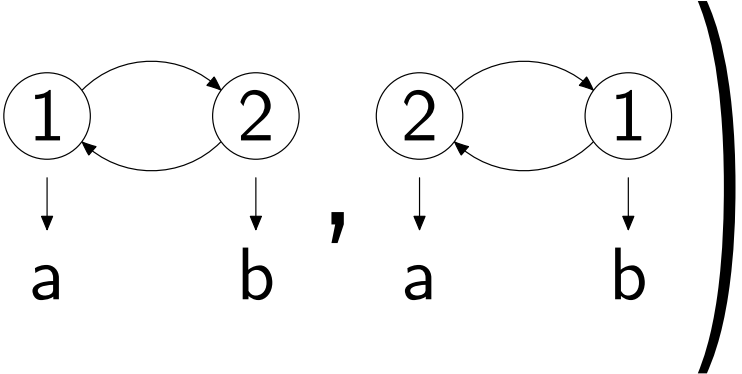
Want distance $\left(\begin{array}{c} \text{Diagram 1} \\ \text{Diagram 2} \end{array} \right) = 0$

The diagram shows two directed graphs side-by-side, enclosed in large parentheses. The left graph has two nodes, 1 and 2, in circles. Node 1 has a downward arrow to the label 'a'. Node 2 has a downward arrow to the label 'b'. There are directed edges from 1 to 2 and from 2 to 1. The right graph has two nodes, 2 and 1, in circles. Node 2 has a downward arrow to the label 'a'. Node 1 has a downward arrow to the label 'b'. There are directed edges from 2 to 1 and from 1 to 2. A comma is placed between the two graphs. To the right of the closing parenthesis is an equals sign followed by a zero.

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- We use bipartite matching to find lower and upper bounds

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- ✓ Identifiability error
- Estimation error
- Optimization error

Estimation and optimization errors

Experiment setup:

- Take some parameters θ^* (say, supervised estimate on real data)
- Use θ^* to generate synthetic data
- Can we recover θ^* using EM?

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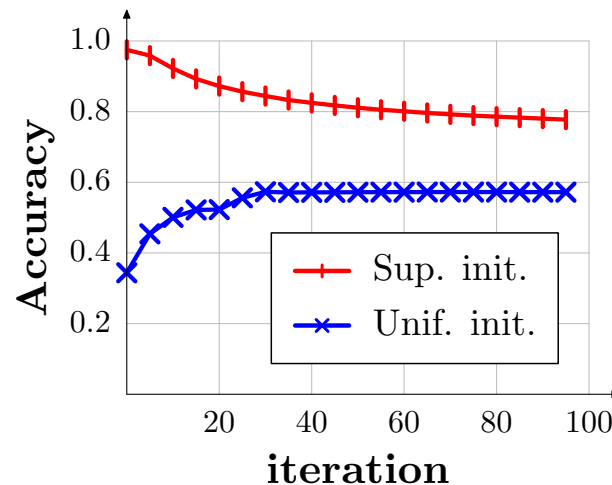
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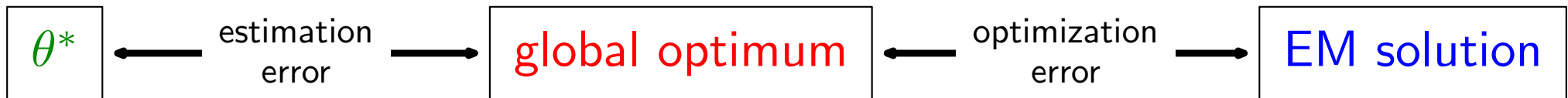
HMM on 5K examples:



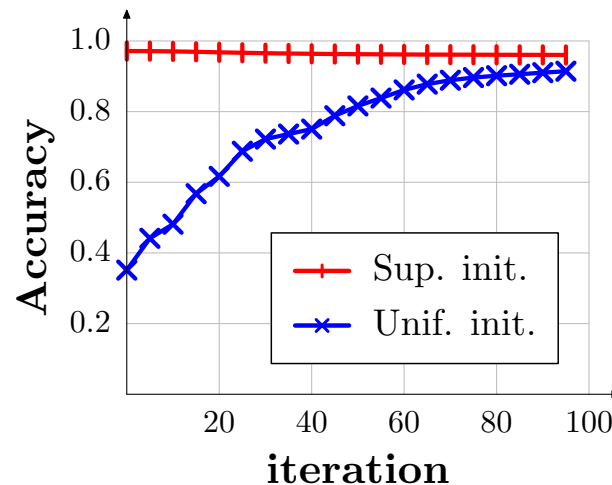
Estimation and optimization errors

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HMM on 500K examples:

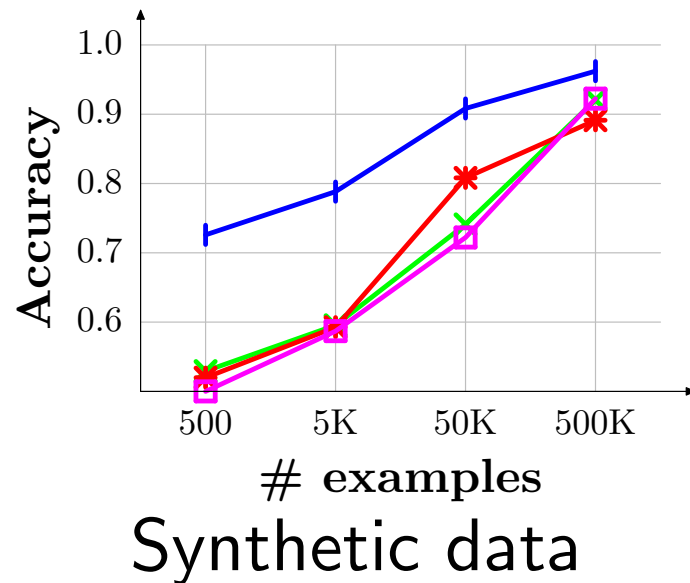


Optimization error decreases with more data

On HMM model (similar for PCFG and a dependency model):

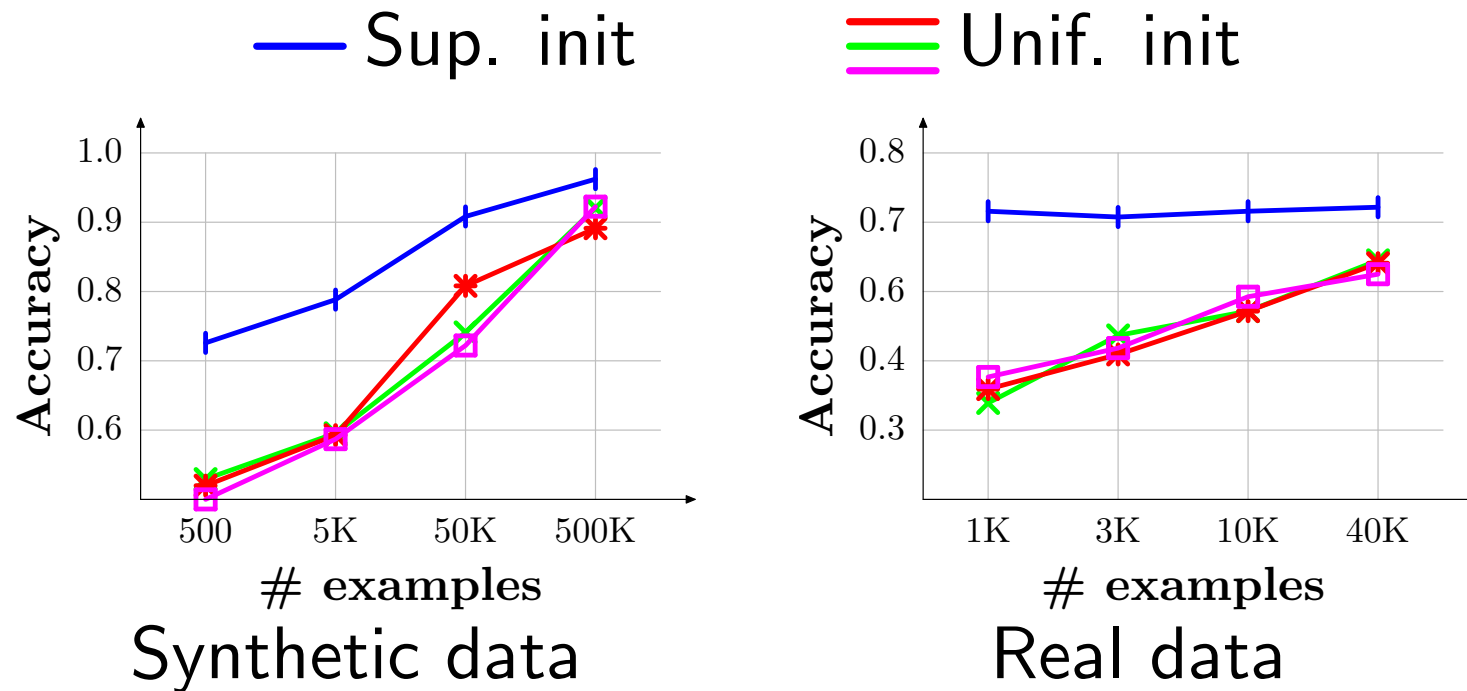
— Sup. init

— Unif. init



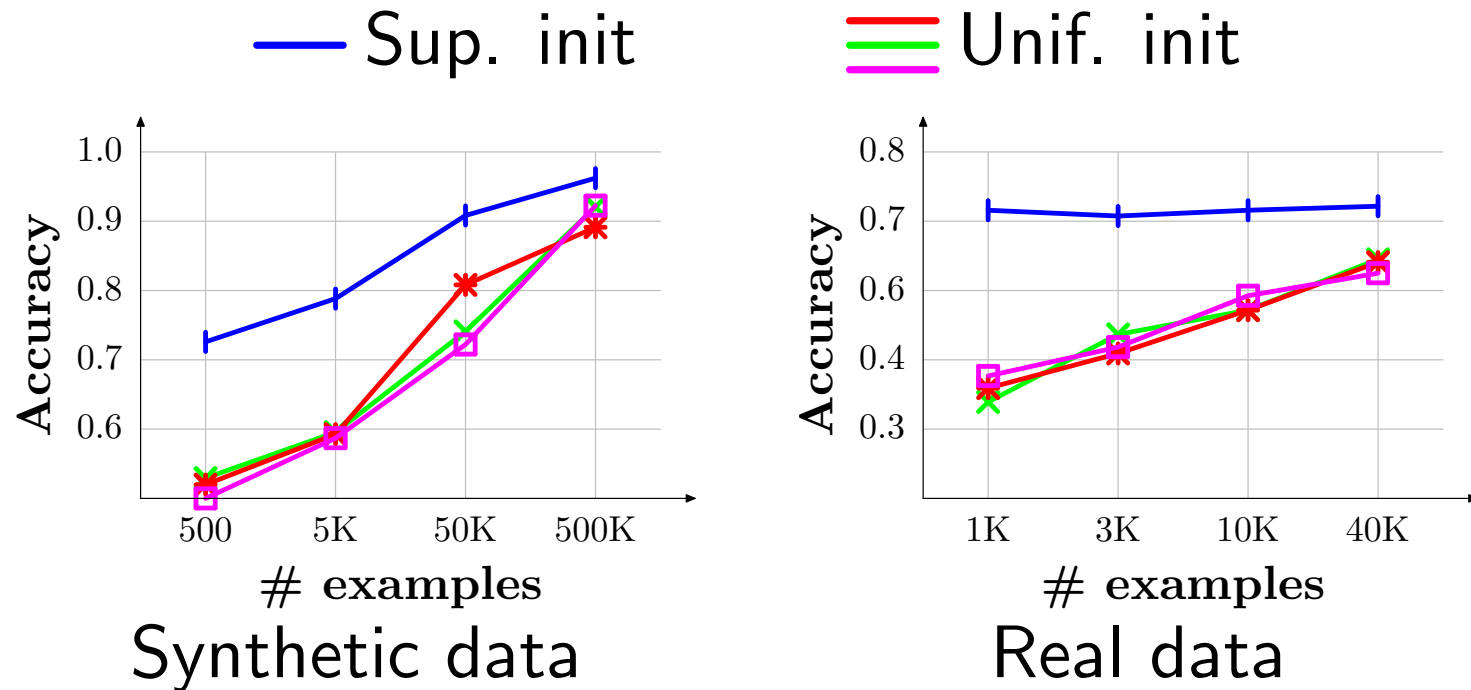
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Why does this phenomenon happen?

- **Intuition:** with more data, EM can pick up the salient patterns more easily
- Was also shown for mixture of Gaussians [Srebro, 2006]

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