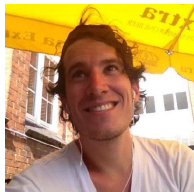


Splitting Compounds By Semantic Analogy



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Introduction

Compound words...

- ▶ ... make life hard for standard NLP applications, incl. MT
- ▶ ... are often modeled with shallow information (e.g. Moses frequency-based splitter)

Question: Can we use distributional semantics to do deeper processing of compounds in a simple way?



Splitting compounds for SMT

- ▶ Koehn and Knight (2003) showed PBMT systems can better deal with compounds if they are split into their meaningful parts
- ▶ Difficulty: many possible splits, we need to choose the correct ones

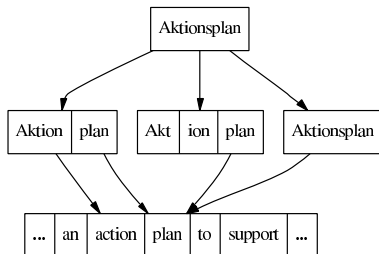


Figure: Compound splitting example from Koehn and Knight (2003).



Compounds and the semantic vector space

Semantic vector space

- ▶ Word embeddings saw surge of successful applications recently
- ▶ Basic idea: "You shall know a word by the company it keeps"
 - Words are mapped to vectors of real numbers in low dimensional space
 - These vectors are estimated on large amounts of text data using a neural network



Compounds and the semantic vector space

Semantic vector space

- ▶ Mikolov et al. (2013) showed that word embeddings capture some linguistic phenomena:
 - *king is to man what queen is to woman*
 $v(\text{king}) - v(\text{man}) + v(\text{woman}) \approx v(\text{queen})$
 - *cars is to car what dogs is to dog*
 $v(\text{cars}) - v(\text{car}) + v(\text{dog}) \approx v(\text{dogs})$



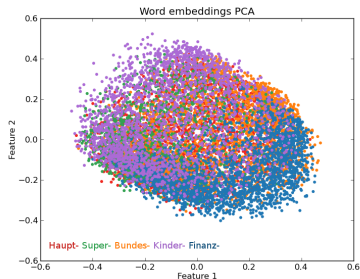
Compounds and the semantic vector space

Morphology induction from word embeddings

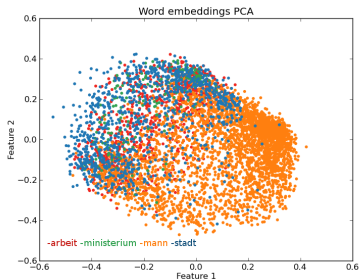
- ▶ Soricut and Och (2015) exploit these regularities to induce morphology from word embeddings
- ▶ Method:
 - Extract prefix and suffix replacement rules from the vocabulary
 - Keep 1000 examples of each rule
 - Judge how well each pair explains the other pairs:
cars is to *car* what *dogs* is to *dog*?
 - Find most representative examples for each rule



Compounds and the semantic vector space



(a) Compounds with same modifier.



(b) Compounds with the same head.



The analogy test

- ▶ We model compounds based on their modifiers
- ▶ Potential compound splits are judged by how similar they are to a set of prototypical compounds for each modifier

Analogy test: *Mauszeiger* is to *Zeiger* what *Mausklick* is to *Klick*?

(mouse pointer)

(pointer)

(mouse click)

(click)



Extracting potential compound splits

For all words in the vocabulary:

- ▶ Extract all possible string prefixes ≥ 4 :
Bundespräsident \rightarrow *Bund, Bunde, Bundes, ...*
- ▶ Judge each Modifier+Compound pair by how well it explains others



Judging potential compound splits

All potential compounds with prefix *Maus*

Maus|kostüm
Maus|zeiger
Maus|stämme
Maus|klick
Maus|hirn
Maus|tasten
Maus|ersatz
Maus|mutanten
Maus|knopf
Maus|steuerung
Maus|bewegung
Maus|gene
Maus|clicks
Maus|hirs
Maus|zeiger
Maus|hirnen
Maus|bedienung
...
(up to 500)



Judging potential compound splits

All potential compounds \times All potential compounds

Maus|kostüm
 Maus|zeiger
 Maus|stämme
 Maus|klick
 Maus|hirn
 Maus|tasten
 Maus|ersatz
 Maus|mutanten
 Maus|knopf
 Maus|steuerung
 Maus|bewegung
 Maus|gene
 Maus|klicks
 Maus|hirs
 Maus|zeiger
 Maus|hirnen
 Maus|bedienung

...
 (up to 500)

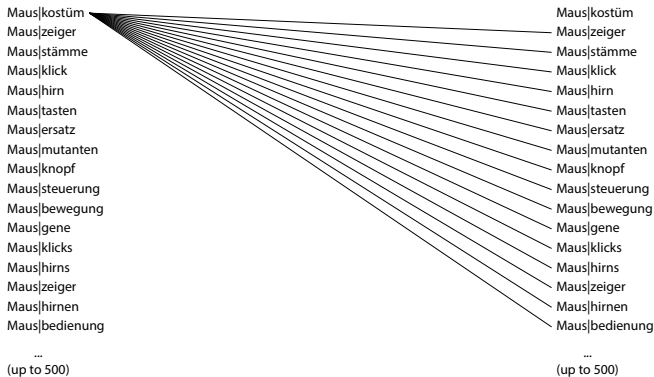
Maus|kostüm
 Maus|zeiger
 Maus|stämme
 Maus|klick
 Maus|hirn
 Maus|tasten
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 Maus|mutanten
 Maus|knopf
 Maus|steuerung
 Maus|bewegung
 Maus|gene
 Maus|klicks
 Maus|hirs
 Maus|zeiger
 Maus|hirnen
 Maus|bedienung

...
 (up to 500)



Judging potential compound splits

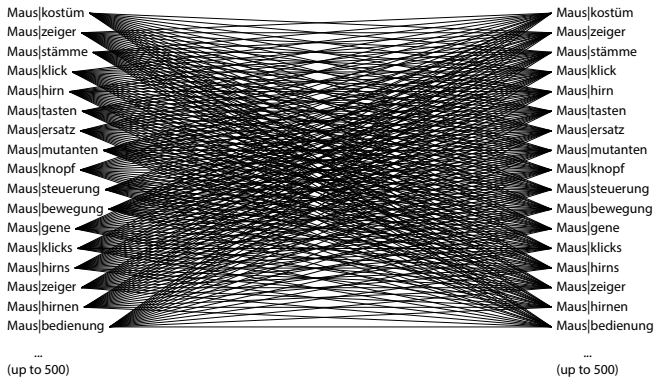
All potential compounds \times All potential compounds





Judging potential compound splits

All potential compounds \times All potential compounds





Judging potential compound splits

Maus|zeiger
Maus|stämme
Maus|klick
Maus|hirn

Maus|zeiger
Maus|stämme
Maus|klick
Maus|hirn

Perform analogy test: *Mauszeiger* is to *Zeiger* what *Mausklick* is to *Klick*?

(mouse pointer)

(pointer)

(mouse click)

(click)



Computational considerations

- ▶ **Analogy test is expensive!**
- ▶ True and predicted vectors:
 - $V_{\text{Mausklick}}$
 - $\hat{V}_{\text{Mausklick}} = \text{Mauszeiger} - \text{Zeiger} + \text{Klick}$
- ▶ Two evaluation functions: RANK and COSINE



Computational considerations

- ▶ Exact but slow implementation:

$$\text{RANK}(v_{\text{compd}}, \hat{v}_{\text{compd}}) = \text{RANK OF } v_{\text{compd}} \text{ IN } \underset{w \in V}{\text{arg sort}} \left[\text{COSINE}(v_w, \hat{v}_{\text{compd}}) \right]$$

- ▶ Approximate but fast implementation:
 - Approximate k-nearest neighbor search
 - We use the Spotify Annoy library (C++) to perform the search
- ▶ *Maus|zeiger* explains *Maus|klick* **IFF**

$$\text{RANK}(v_{\text{compd}}, \hat{v}_{\text{compd}}) < 100 \quad \mathbf{AND} \quad \text{COSINE}(v_{\text{compd}}, \hat{v}_{\text{compd}}) > 0.5$$



Prototypes

Compounds that are good examples of a compound modifier.

- ▶ These are best at explaining other similar modifier+compound pairs
- ▶ We call this set the modifier's *prototypes*



Extracting prototypes

Maus|kostüm
Maus|zeiger
Maus|stämme
Maus|klick
Maus|hirn
Maus|tasten
Maus|ersatz
Maus|mutanten
Maus|knopf
Maus|steuerung
Maus|bewegung
Maus|gene
Maus|klicks
Maus|hirs
Maus|zeiger
Maus|hirnen
Maus|bedienung

...
(up to 500)

Maus|kostüm
Maus|zeiger
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Maus|gene
Maus|klicks
Maus|hirs
Maus|zeiger
Maus|hirnen
Maus|bedienung

...
(up to 500)



Extracting prototypes

Maus|kostüm
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Maus|hirs
Maus|zeiger
Maus|hirnen
Maus|bedienung

...
(up to 500)

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Maus|hirnen
Maus|bedienung

...
(up to 500)



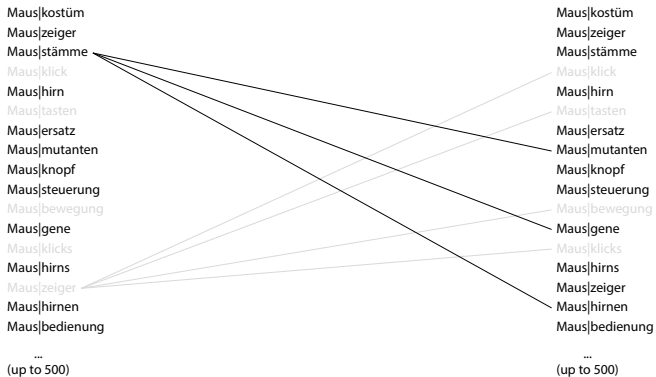
Extracting prototypes

Maus|kostüm
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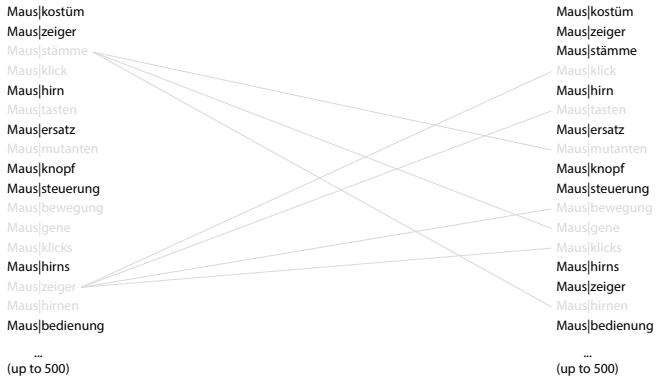


Extracting prototypes





Extracting prototypes





Extracted prototypes for *Maus-*

Prototype	Evidence words
V-Zeiger	-Bewegung -Klicks -Klick -Tasten -Zeiger
V-Stämme	-Mutanten -Gene -Hirnen -Stämme
V-Kostüm	-Knopf -Hirn -Hirns -Kostüm
V-Steuerung	-Ersatz -Bedienung -Steuerung



Compound splitting: *Mausmutation*

Mausmutation

- ▶ We start from the left...





Compound splitting: *Mausmutation*

Mausmutation
→

- ▶ Do I know the modifier *Mau*? No!



Compound splitting: *Mausmutation*

Mausmutation
→

- ▶ Do I know the modifier *Maus*? Yes!



Compound splitting: *Mausmutation*

Mausmutation



- ▶ Do I know the modifier *Maus*? Yes!

Prototypes:

- -Zeiger
- -Stämme
- -Kostüm
- -Steuerung



Compound splitting: *Mausmutation*

Mausmutation



- ▶ Do I know the modifier *Maus*? Yes!

Prototypes:

- Zeiger
- Stämme ✓
- Kostüm
- Steuerung

→ *Mausmutation* is to *Mutation* what *Mausstämme* is to *Stämme*.



Compound splitting: *Mausmutation*

Mausmutation



- ▶ Do I know the modifier *Mausm*? No!



Compound splitting: *Mausmutation*

Mausmutation

- ▶ And so on...



Compound splitting: *Mausmutation*

Maus|mutation

- ▶ The prototype with the highest score will be our split!
- ▶ Recurse...



Compound splitting: *Plantage*

Plantage

- ▶ Let's try another example...





Compound splitting: *Plantage*

Plantage
→

- ▶ Do I know the modifier *Plan*? Yes!



Compound splitting: *Plantage*

Plantage
→

- ▶ Do I know the modifier *Plan*? Yes! Prototypes:
 - -Feststellung
 - -Wert
 - -Fertiger
 - ...



Compound splitting: *Plantage*

Plantage
→

- ▶ Do I know the modifier *Plan*? Yes! Prototypes:
 - -Feststellung
 - -Wert
 - -Fertiger
 - ...



Compound splitting: *Plantage*

Plantage

- ▶ No compound split!

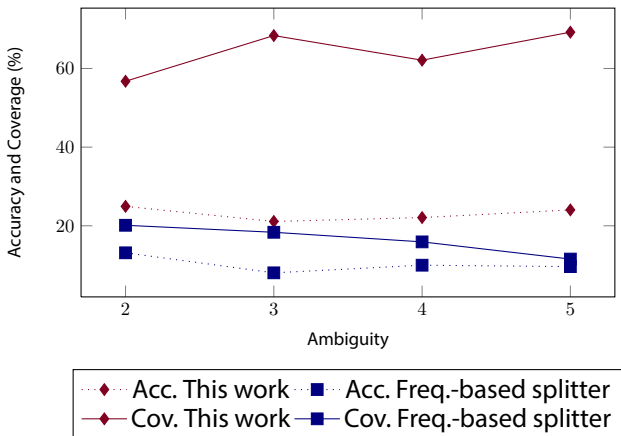


Intrinsic evaluation

- ▶ Evaluation on human-annotated dataset (Henrich and Hinrichs, 2011)
 - ~50k compounds
 - only binary splits
- ▶ Baseline: Frequency-based Moses compound splitter (Koehn and Knight, 2003)
- ▶ We evaluate:
 - Accuracy: $\frac{|\text{correct splits}|}{|\text{compounds}|}$
 - Coverage: $\frac{|\text{compounds split}|}{|\text{compounds}|}$



Intrinsic evaluation





Machine translation experiments (German to English)

	(a) No comp. splitting			(b) Rare: $c(w) < 20$			(c) All words		
	Splits	BLEU	MTR	Splits	BLEU	MTR	Splits	BLEU	MTR
Moses splitter	0	17.6	25.5	231	17.6	25.7	244	17.9	25.8 ^A
This work				744	18.2 ^{ABC}	26.1 ^{ABC}	1616	17.7	26.3 ^A

^A Stat. sign. against (a) at $p < 0.05$ ^B Stat. sign. against Moses splitter at same $c(w)$ at $p < 0.05$

^C Stat. sign. against best Moses splitter (c) at $p < 0.05$



Conclusion

- ▶ Regularities in semantic vector space can be used to model composition of compounds
- ▶ We can extract modifiers and prototypes (Soricut and Och, 2015)
- ▶ Compound splitting algorithm:
 - Good intrinsic performance on gold standard
 - Improved translation quality (standard PBMT setup)
 - Especially adept at splitting highly ambiguous compounds



Thank You!

Any questions?



References

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