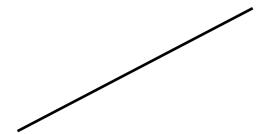
# Homework 1

- Leaderboard
- Read through, submit the default output
- Time for questions on Tuesday

# Agenda

- Focus on Homework 1
  - Review IBM Models 1 & 2
  - Inference (compute best alignment from a corpus given model parameters)
  - Parameter estimation (find model parameters)
- Discuss HMM model and IBM models 3–5

$$p(\text{target} \mid \text{source}) = \sum_{\text{alignments}} p(\text{alignment}) \prod_{\text{target word}} p(\text{target word} \mid \text{source word})$$



Model 1: uniform

Model 2: absolute positioning

Word translation table

#### Review: IBM Models 1 & 2

- A generative model is a data creation story
- IBM models: p(f | e) how f is generated from e
  - Assuming the data was generated by the model, which way did it most likely happen?
  - Always ask, what are the parameters of the model?
  - Each step of the story needs parameters

- Given: an English sentence **e**, parameters **q** and **t**
- Choose a French length m
- For each French word position *i* ∈ 1...*m*
  - Choose a source word position  $\mathbf{a_i} = \mathbf{q}(\mathbf{j} \mid \mathbf{i}, \mathbf{l}, \mathbf{m})$
  - Choose a translation probability t(f<sub>i</sub> | e<sub>ai</sub>)

# Model Parameters

**t**(**f** | **e**)

**q**(**j** | **i**, **l**, **m**)

Models 1 & 2

f	е	p(f   e)
le	the	0.42
la	the	0.4
programme	the	0.001
a	has	0.78

Model 1

$$\frac{1}{l+1}$$

Model 2

### Task 1: Inference

- Input: a sentence pair (e,f) and a model (t,q)
- Models 1 & 2: each link is generated independently
  - For each target word, compute most likely alignment link

$$p(a_i = j \mid e, f) = q(j \mid i, l, m)t(f_i \mid e_{a_i})$$

Choose the one that maximizes this probability

- Input: a sentence pair (e,f) and a model (t,q)
- Models 1 & 2: each link is generated independently

$$p(a_i = j \mid e, f) = q(j \mid i, l, m)t(f_i \mid e_{a_i})$$

NULL And the program has been implemented

Le programme a ete mis en application

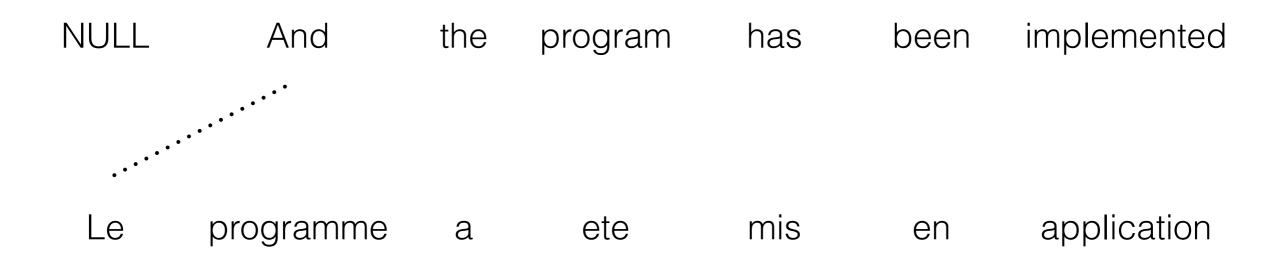
- Input: a sentence pair (e,f) and a model (t,q)
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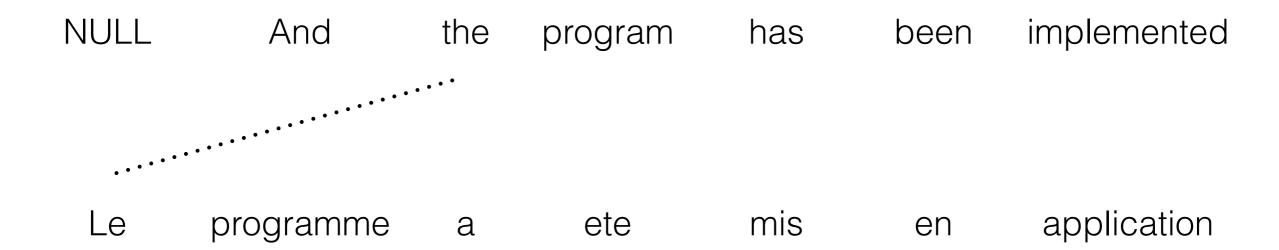
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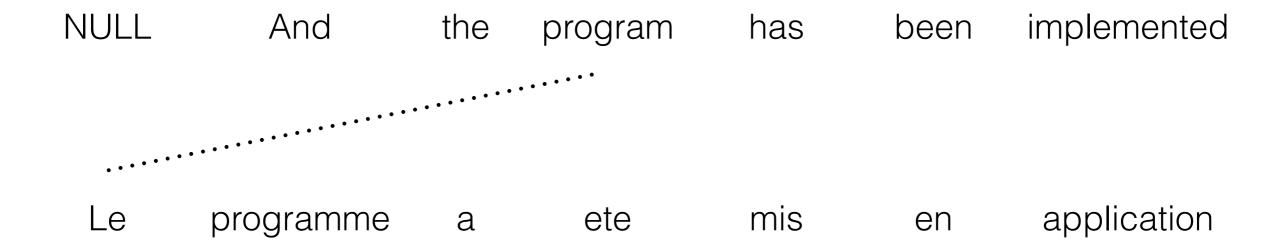
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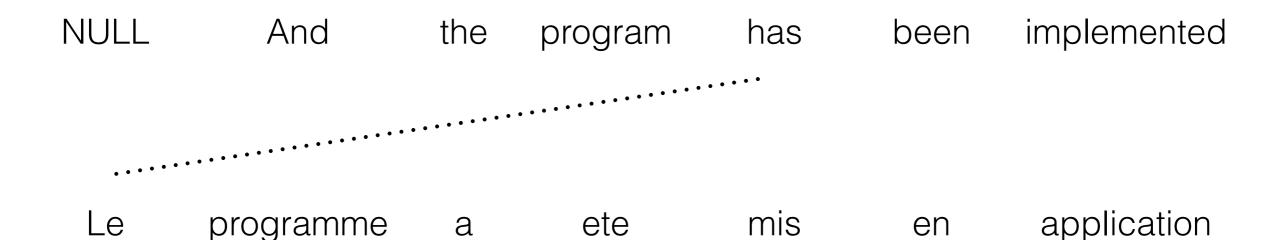
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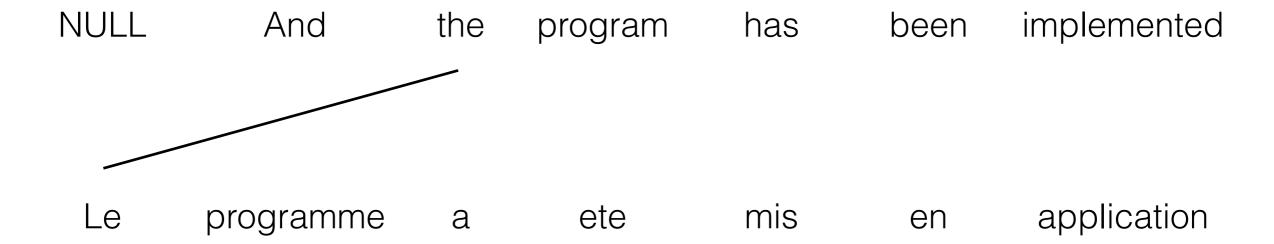
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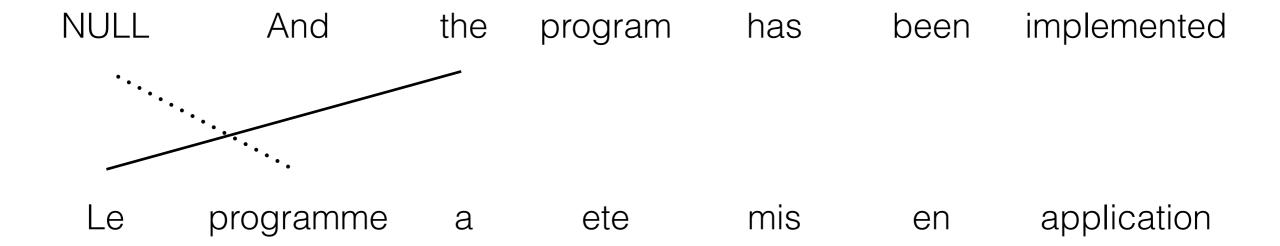
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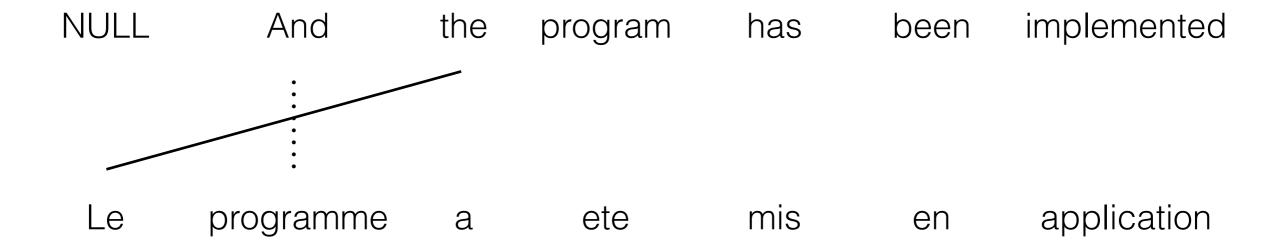
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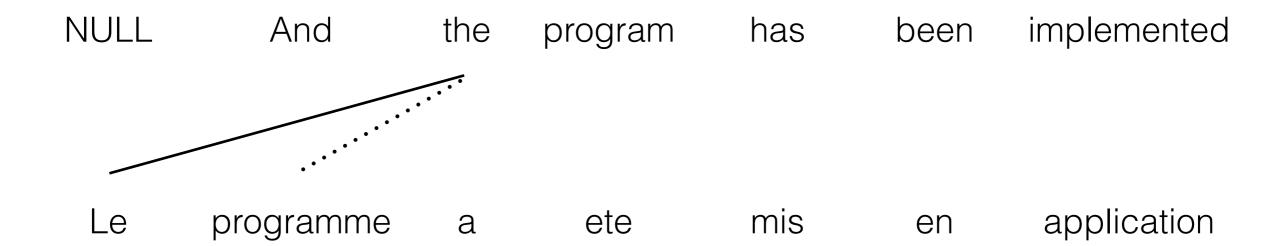
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#### Task 2: Parameter Estimation

 Guess parameters, compute expectations, adjust, repeat

```
initialize parameters t and q to something repeat until convergence for every sentence for every target position j for every source position i count(f_j, e_i) += P(a_i = j | e_i, f_j) count(e_i) += P(e_i = e_i, e_i) count(e_i) += P(e_i = e_i, e_i) count(e_i, e_i, e_i) += P(e_i = e_i, e_i) t(e_i) count(e_i) += P(e_i = e_i, e_i) t(e_i) += Count(e_i) count(e_i) count(e_i) count(e_i) count(e_i) q(e_i) | count(e_i) count(e_i) count(e_i) q(e_i) | count(e_i) count(e_i) | count(
```

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  - Each word and alignment link are generated separately; there are no dependencies between alignment links at all

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  - Each word and alignment link are generated separately; there are no dependencies between alignment links at all
  - ★ The cost of easy inference here is an overly simplistic model

#### Pros and cons

- Some drawbacks of word based alignments
  - All reorderings have the same probability
  - Alignments are independent
  - No notion of multiword alignments
  - Alignments are asymmetric
  - No morphology
  - No syntax

#### Pros and cons

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MODEL 2

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# Building intuitions

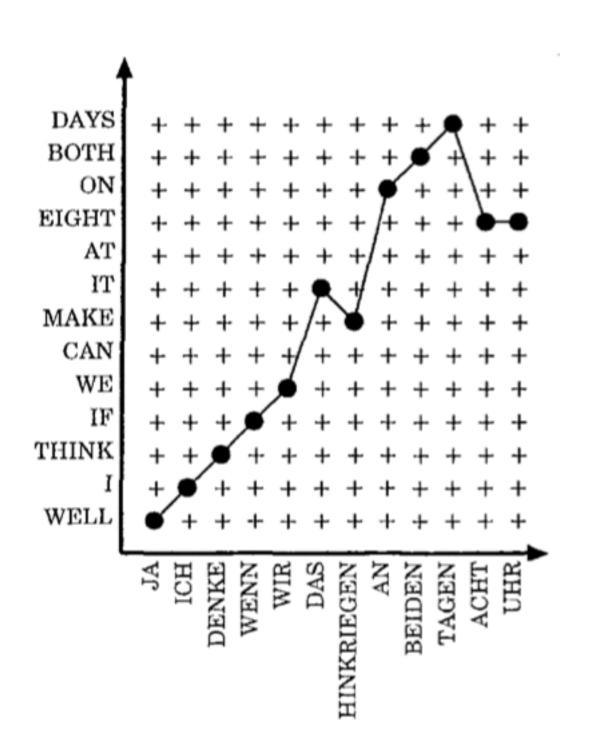
- Model 2 still generates all alignments independently
- Let's try to think of something we might change
  - Using an important (and oft-avoided) tool in the scientist's toolkit: looking at the data
  - We'll use Picaro, a tool for alignment visualization github.com/joshua-decoder/picaro

#### Discuss

5 minutes, with a neighbor or two

- What patterns did you see in the alignments?
   (order them from simplest to most complex)
- Pick one pattern: how might you model it? What parameters would you need?

# Vogel, Ney, & Tillmann ('96)



We now propose an HMM-based alignment model. The motivation is that typically we have a strong localization effect in aligning the words in parallel texts (for language pairs from Indoeuropean languages): the words are not distributed arbitrarily over the sentence positions, but tend to form clusters. Fig. 1 illustrates this effect for the language pair German–English.

Each word of the German sentence is assigned to a word of the English sentence. The alignments have a strong tendency to preserve the local neighborhood when going from the one language to the other language. In many cases, although not always, there is an even stronger restriction: the difference in the position index is smaller than 3.

(Hidden Markov Model)

$$p(a \mid e, m) = \prod_{i=1}^{m} q(a_i = j \mid i, l, m)$$

$$p(a \mid e, m) = \prod_{i=1}^{m} p(a_i \mid a_{i-1})$$

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Model 2 used the absolute positions of words

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 A better idea: relative positioning using position differences

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A "jump" probability

 What are the parameters of this alignment model?

- What are the parameters of this alignment model?
  - A simple table

jump distance	prob
-3	0.03
-2	0.05
-1	0.12
0	0.2
1	0.3
2	0.09
3	0.08

#### HMM Model

- What are the parameters of this alignment model?
  - A simple table
  - Other ways?

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#### HMM Model

- What are the parameters of this alignment model?
  - A simple table
  - Other ways?
- What else might we like to condition on?

jump distance	prob
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### HMM Model

- What's different about inference with this model?
  - Alignment links are no longer (conditionally) independent!
  - Inference (and EM) now require something more complicated (dynamic programming)

- Some drawbacks of word based alignments
  - All reorderings have the same probability



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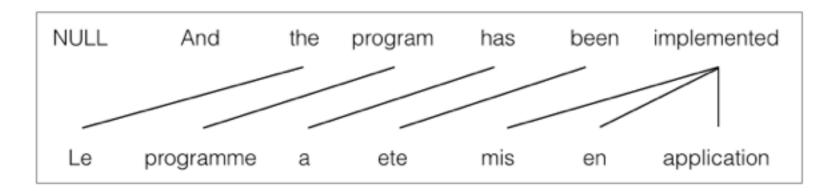
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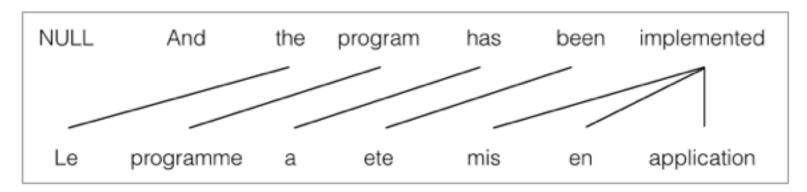
MODEL 2

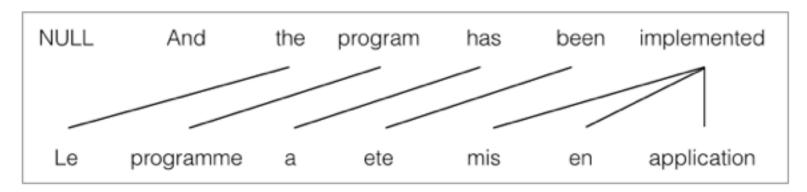
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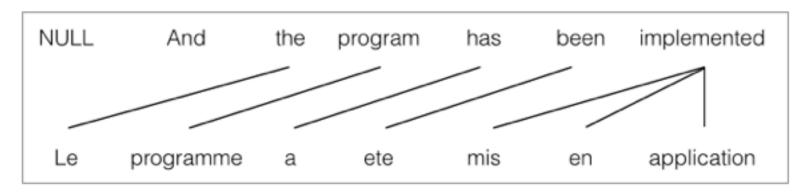
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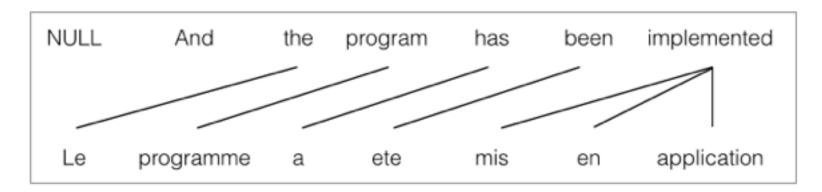




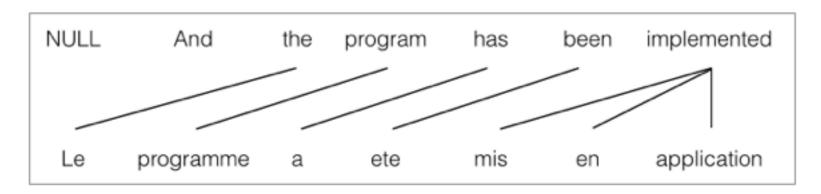




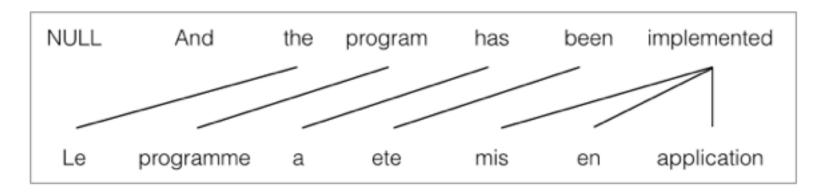
• Fertility: some words produce more translations



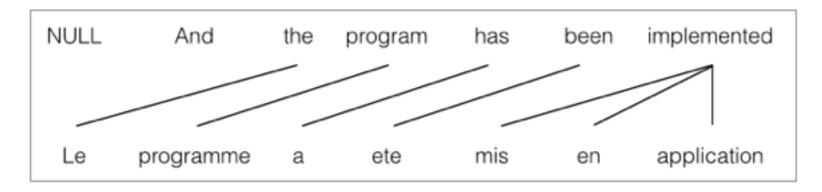
• Allowed in previous models, but not permitted / discouraged



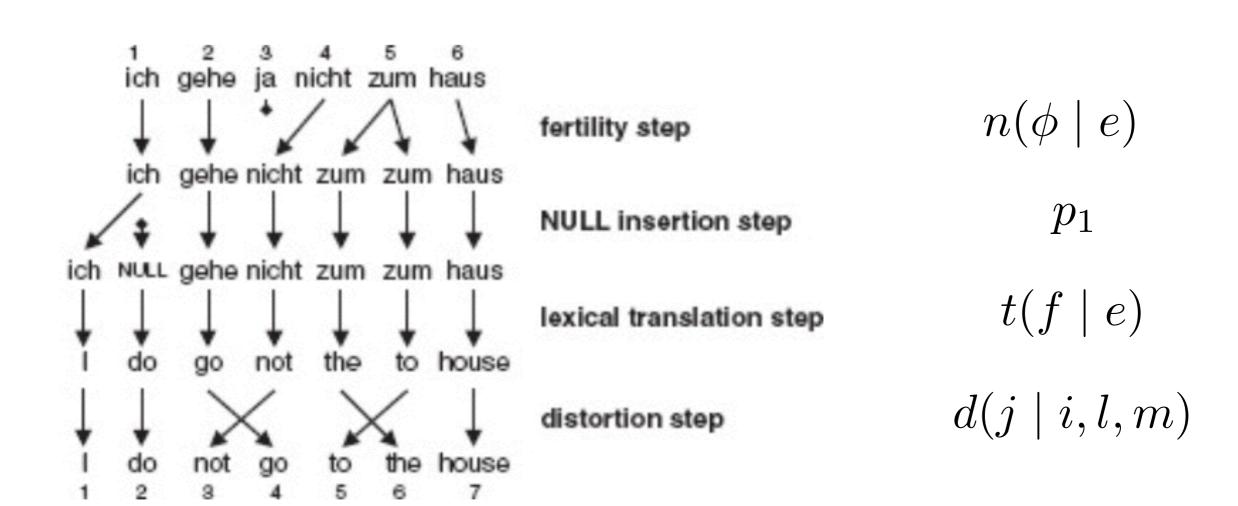
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- NULL fertility?



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- NULL fertility?
  - No, more related to sentence length



- Allowed in previous models, but not permitted / discouraged
- NULL fertility?
  - No, more related to sentence length
  - Instead, randomly insert NULL after each word with probability p



# Fertility

- The complete alignment p(a | f, m) no longer factorizes to independent alignment decisions
- Now have to resort to sampling
- Basic idea
  - Seed Model 3 parameters with best Model 2 alignment
  - Randomly make small changes, collect Model 3 counts every once in a while

- Some drawbacks of word based alignments
  - All reorderings have the same probability

MODEL 2

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- \_\_\_\_

MODEL 2

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MODEL 3

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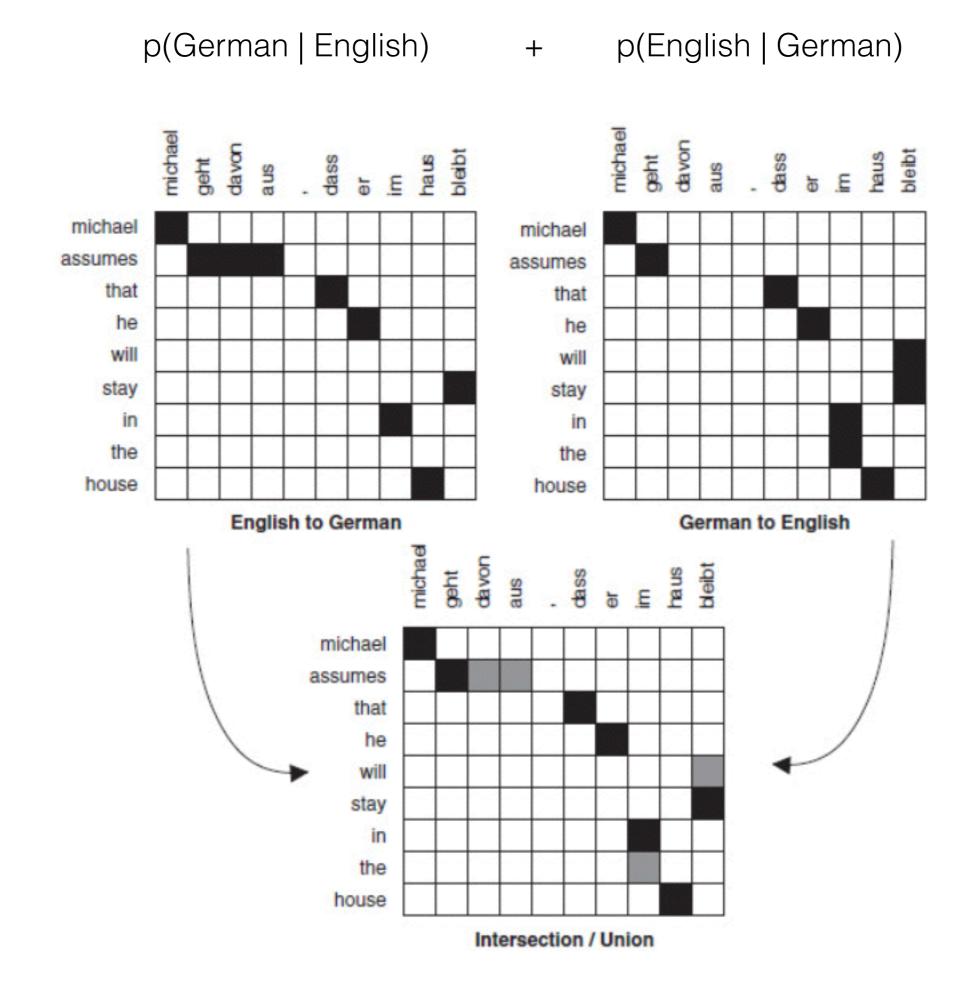
# Higher IBM Models

- Increasingly model new phenomena at the cost of model complexity
  - Model 4: cepts and relative distortion
  - Model 5: solves deficiency of Model 4
- Inference is now accomplished with sampling

Alignments are still asymmetric (why?)

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  - All models explain each target word f with a link to a single source word e

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- Alignments are still asymmetric (why?)
  - All models explain each target word f with a link to a single source word e
- Solution: build two models and combine them
- Used for phrase-based translation (next week)

# Summary

- Lexical alignment: IBM Models 1–5
  - Model 1: word-based translation
  - Model 2: +non-uniform alignments
  - Model 3: +fertility
  - Model 4: +cepts and distortion
  - Model 5: –deficiency
- HMM alignment: relative positioning

# Key points

- General tradeoff between complexity of model and ease of inference
- Modeling ideas come from general knowledge and looking at the data
- Keep things concrete with a generative story and being explicit about how parameters are represented
- Simple models are useful for initializing more complicated ones

# Big Picture

