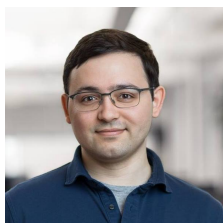


.\|

Analyzing Uncertainty in Neural Machine Translation



Myle Ott

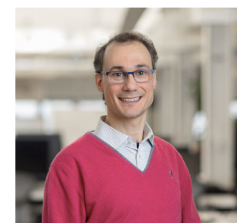
myleott@fb.com



Michael Auli



David Grangier



Marc'Aurelio Ranzato

Facebook AI Research

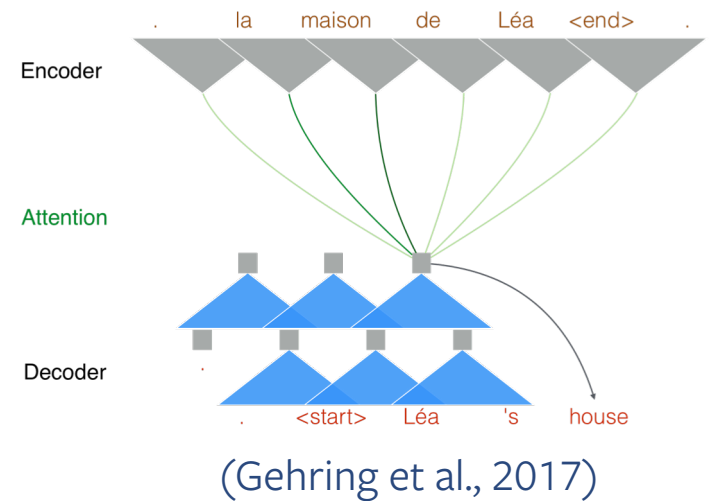
. \ | Background

Neural Machine Translation

Input: source sentence $X = \{x_1, \dots, x_N\}$

Output: target translation $Y = \{y_1, \dots, y_T\}$

$$p(Y|X; \theta) = \prod_{t=1}^T p(y_t | y_{1:t-1}, X; \theta)$$



.\ | Background

Training: maximum likelihood (autoregressive) with cross entropy loss

$$\mathcal{L}_{\text{ML}} = \sum_{t=1}^T \log p(y_t | y_{1:t-1}, X; \theta)$$

Inference: sampling or MAP

$$\hat{y}_{\text{MAP}} = \arg \max_{w_{1:T}} \sum_t \log p(w_t | w_{1:t-1}, X; \theta)$$

. \ | Background

Training: maximum likelihood (autoregressive) with cross entropy loss

$$\mathcal{L}_{\text{ML}} = \sum_{t=1}^T \log p(y_t | y_{1:t-1}, X; \theta)$$

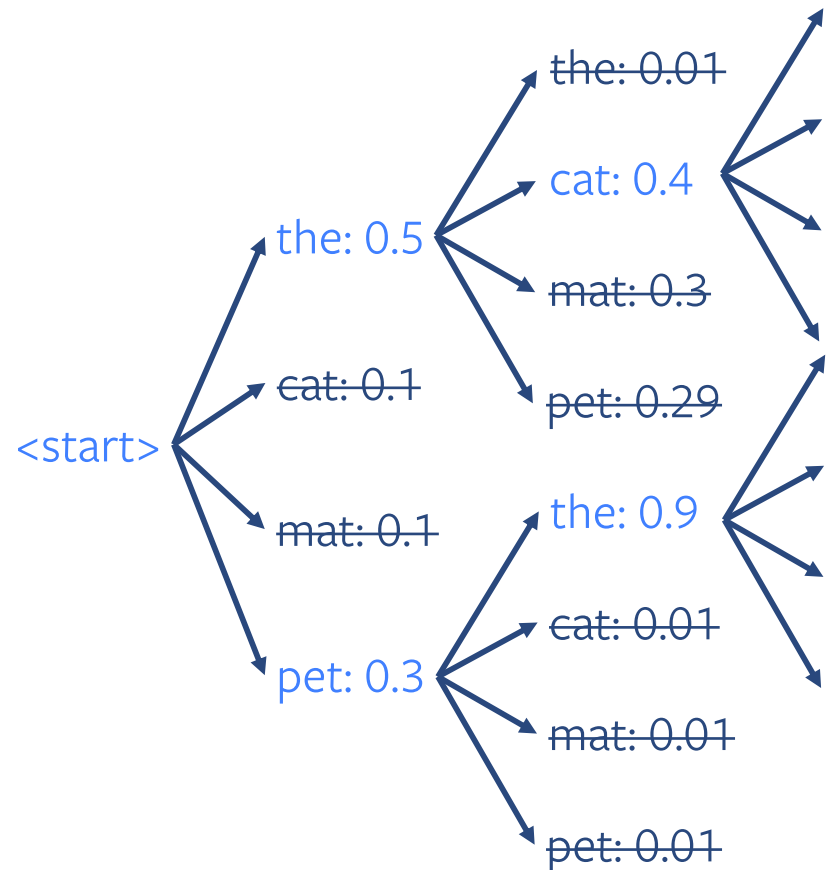
Inference: sampling or MAP Intractable to enumerate

$$\hat{y}_{\text{MAP}} = \boxed{\arg \max_{w_{1:T}}} \sum_t \log p(w_t | w_{1:t-1}, X; \theta)$$

.\\ | Background

Approximate inference with
beam search

- Decode sequence left-to-right and keep K best hypotheses at each step
- Equiv. to greedy search when the beam width (K) = 1



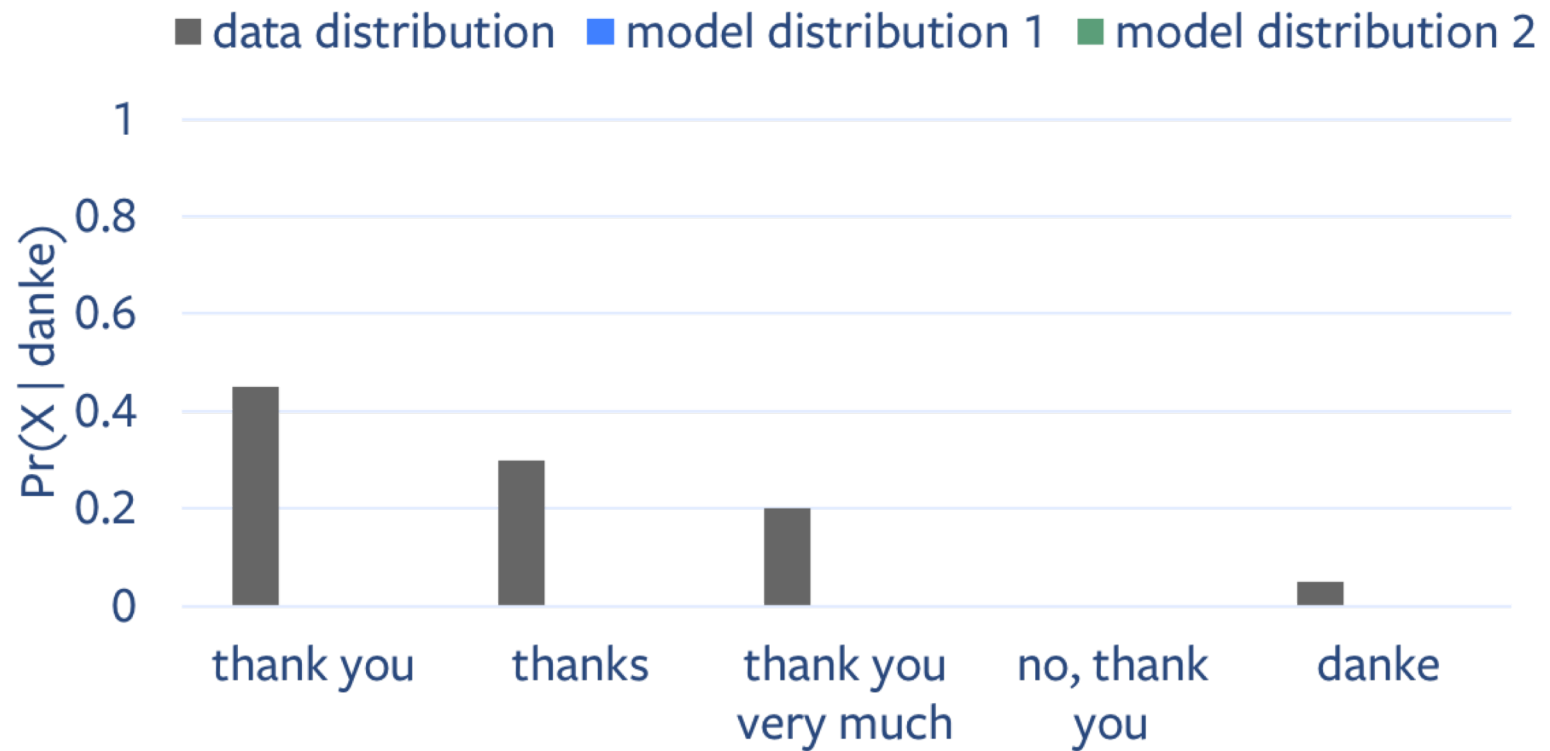
.\| This work

Goal: Investigate the effects of **uncertainty** in NMT model fitting and search

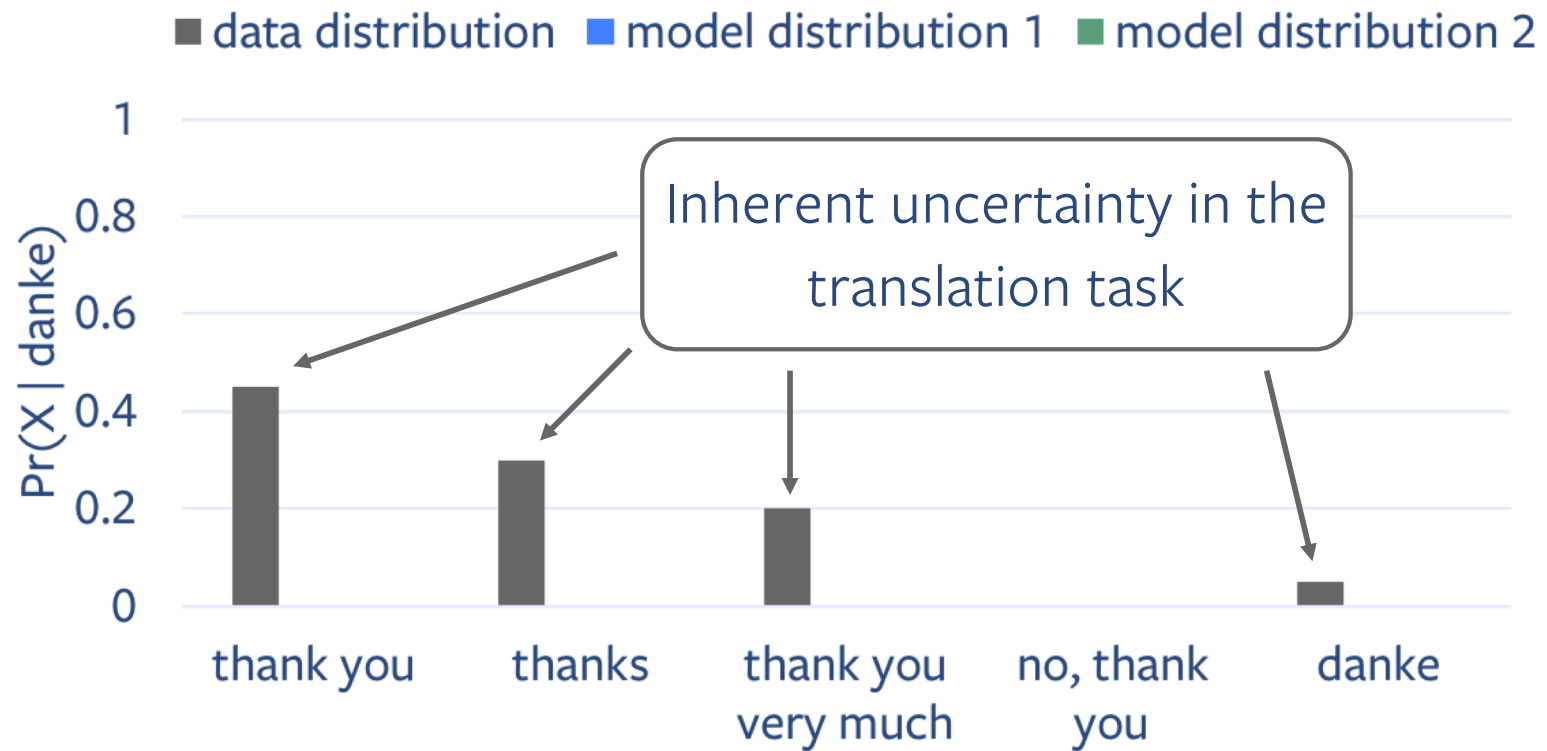
.\| This work



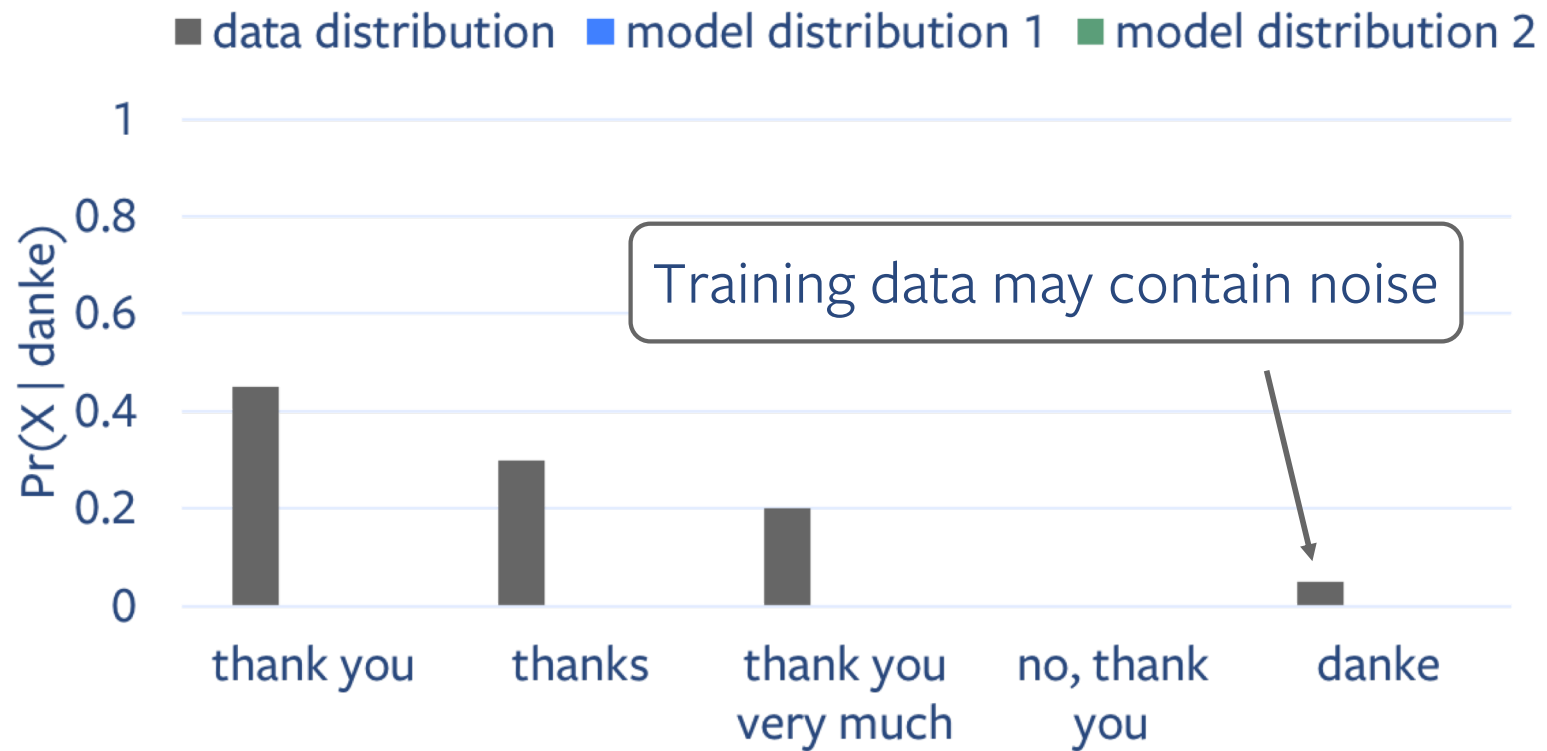
.\| This work



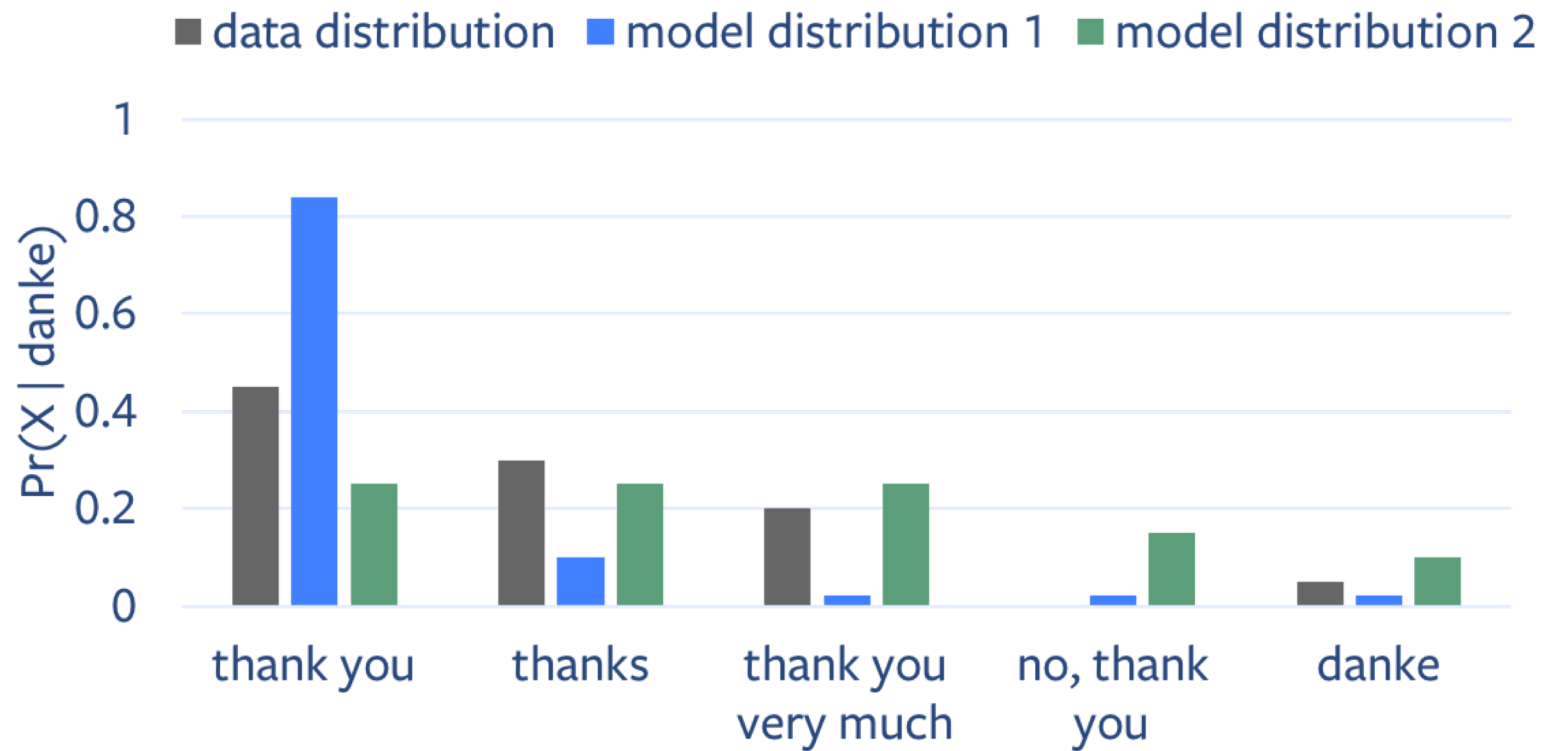
.\\ This work



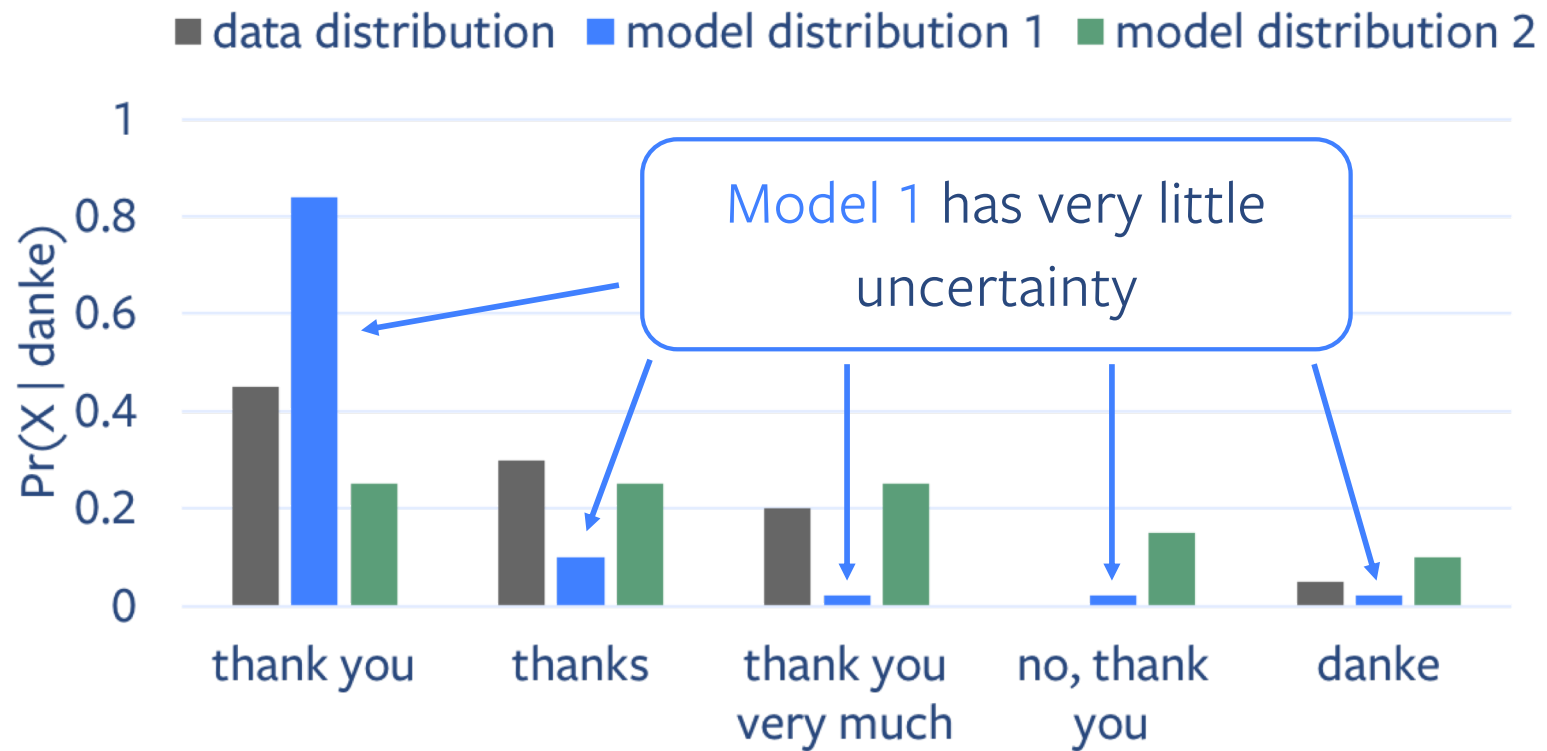
.\\ This work



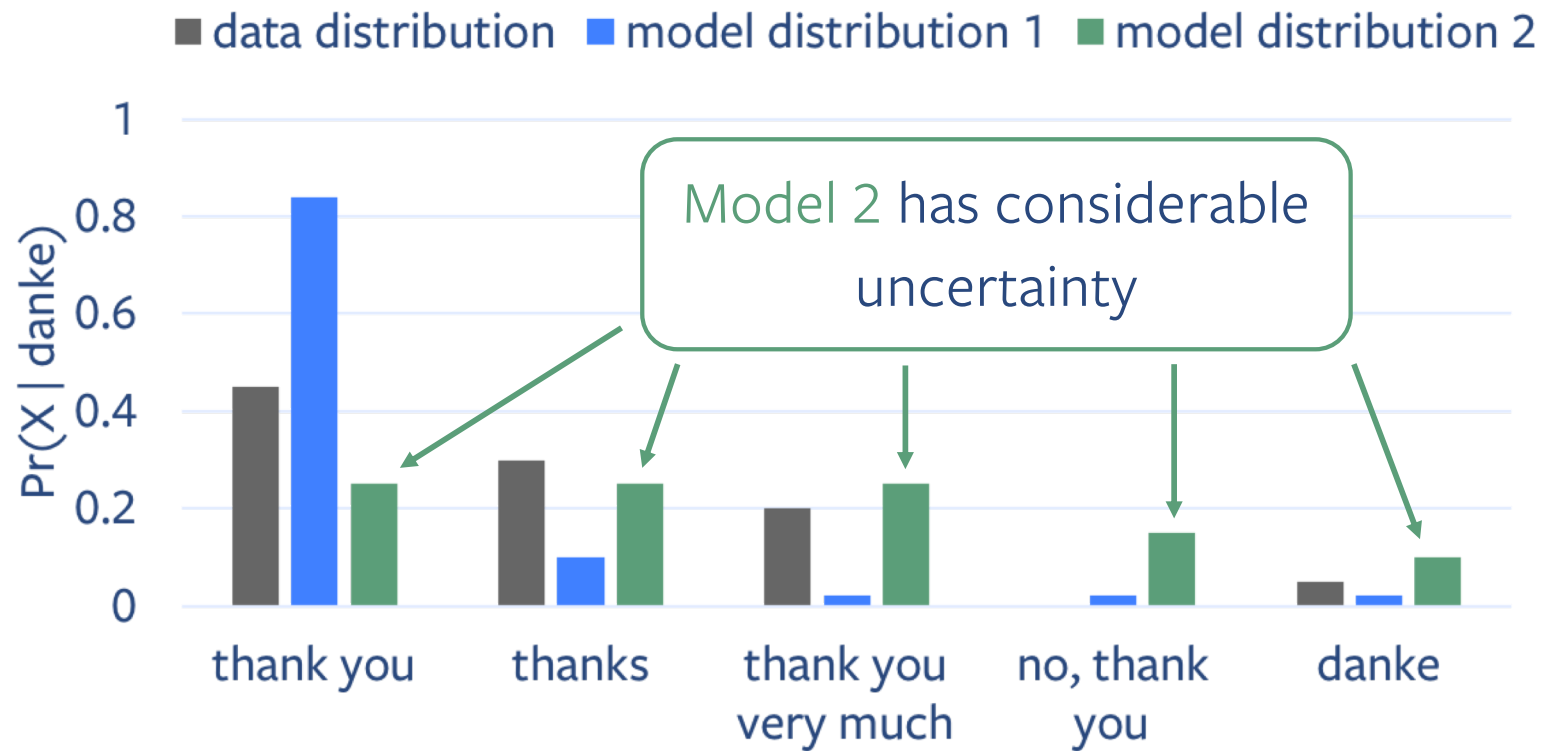
.\| This work



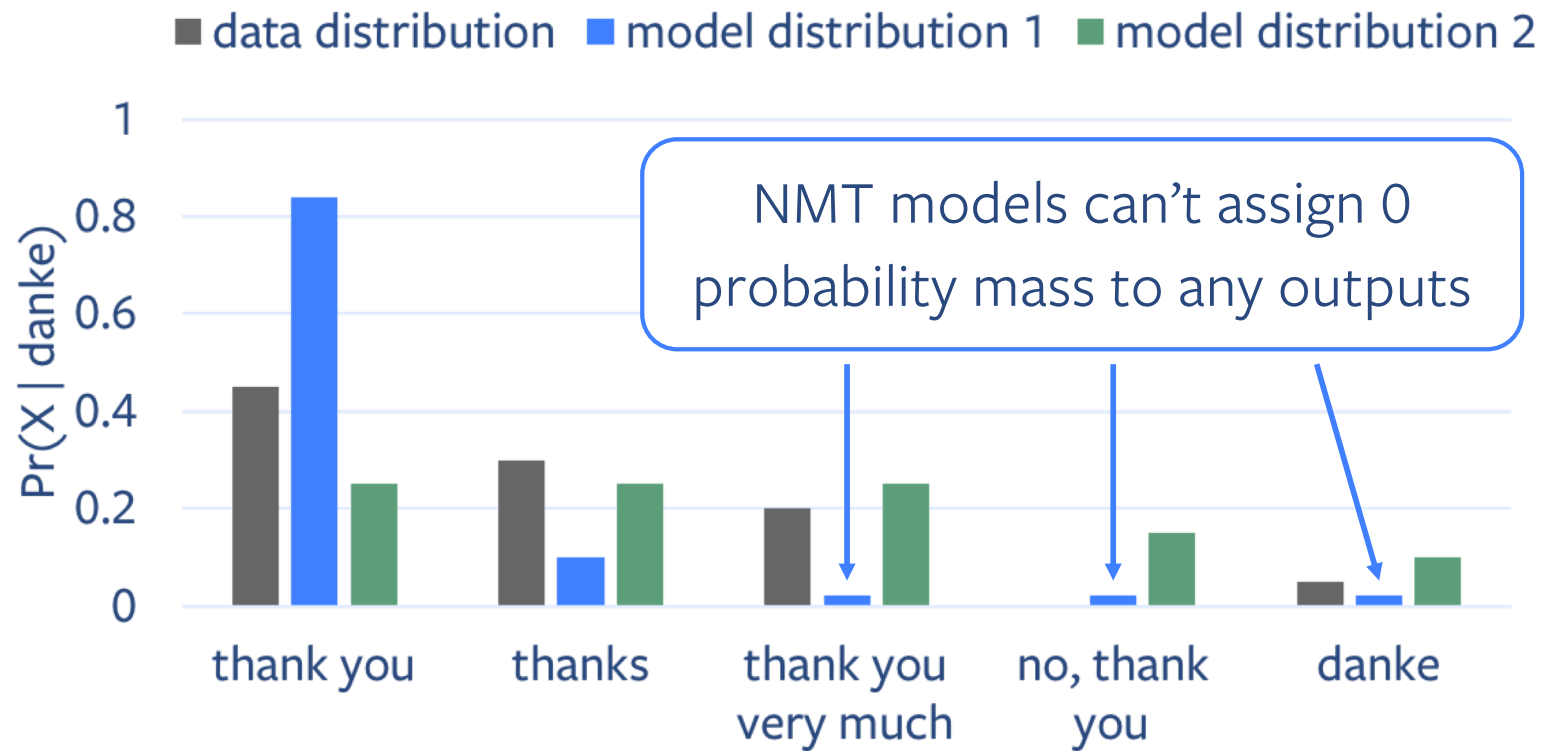
.\\ This work



.\\ This work



.\\ This work



. \ | This work

Goal: Investigate the effects of **uncertainty** in NMT model fitting and search

- Do NMT models capture uncertainty, and how is this uncertainty represented in the model's output distribution?
- How does uncertainty affect search?
- How closely does the model distribution match the data distribution?
- How do we answer these questions with (typically) only a single reference translation per source sentence?

. \ | Experimental setup

Convolutional sequence-to-sequence models* (Gehring et al., 2017)

Evaluation: compare translations with **BLEU** (Papineni et al., 2002)

- Modified n-gram precision metric, values from 0 (worst) to 100 (best)

Datasets: WMT14 English-French and English-German

- Mixture of news, parliamentary and web crawl data

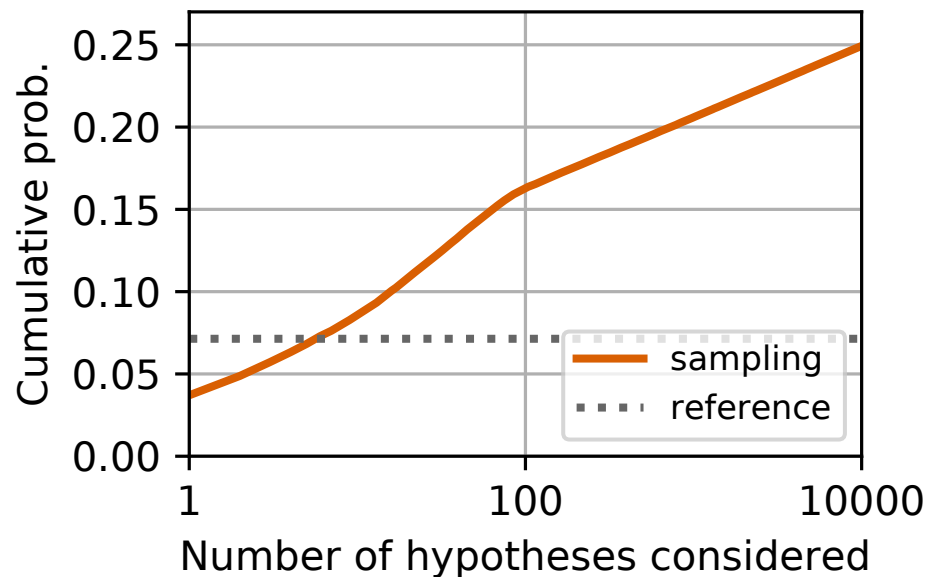
* Results hold for other tested architectures too, e.g., LSTM

. \ | Do NMT models capture uncertainty?

Question: How much uncertainty is there in the model's output distribution?

Experiment: How many independent samples does it take to cover most of the sequence-level probability mass?

. \ | Do NMT models capture uncertainty?



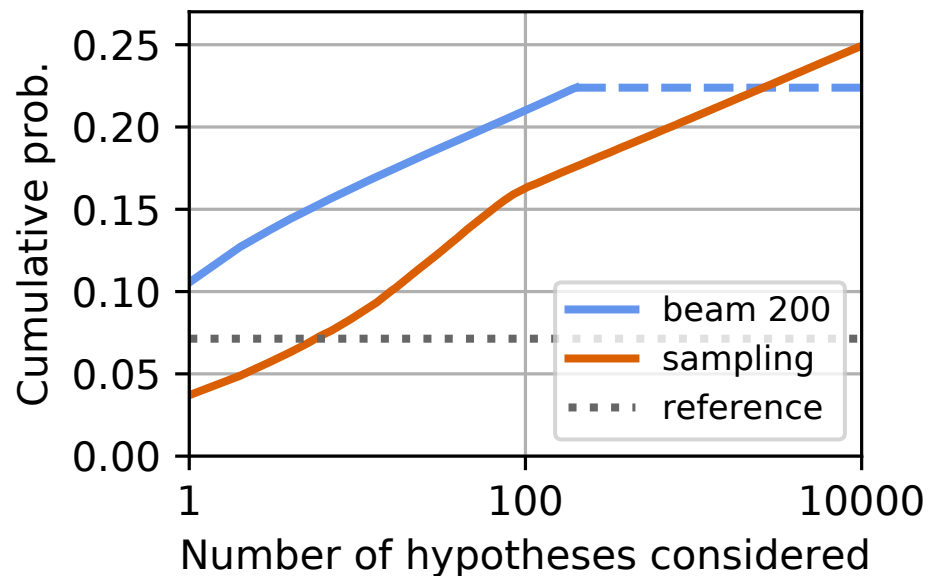
(WMT14 En-Fr)

Model's output distribution is **highly uncertain!**

- Even after 10K samples we cover only 25% of sequence-level probability mass

What about **beam search**?

.\\ | Do NMT models capture uncertainty?



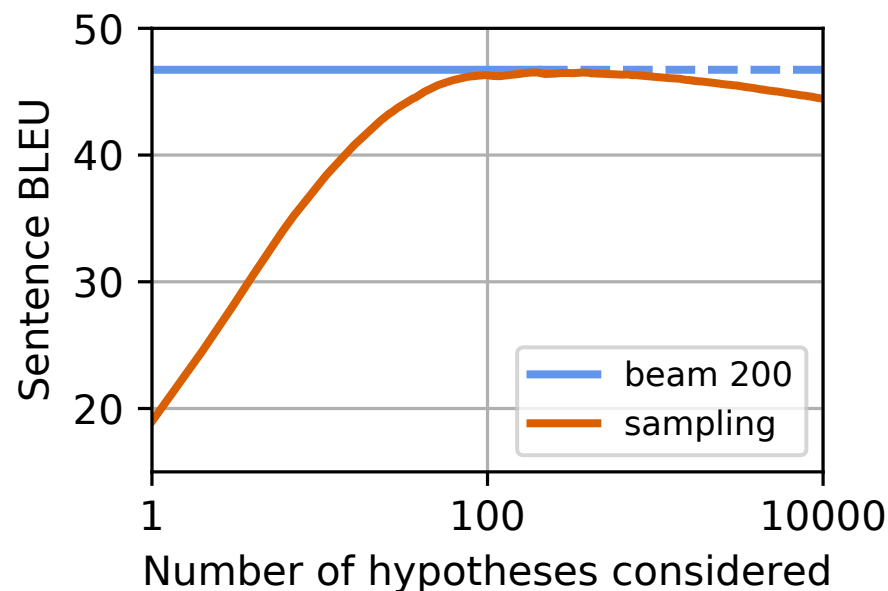
(WMT14 En-Fr)

Beam search is very efficient!

The reference score (....) is lower than beam hypotheses

What is the quality (BLEU) of these translations?

. \ | Uncertainty and Search



(WMT14 En-Fr)

Beam search is efficient and produces accurate translations

Sampling produces increasingly likely hypotheses, but these get worse BLEU after ~200

.\| Uncertainty and Search

Source: The first nine episodes of Sheriff Callie 's Wild West will be available (...)

Reference: Les neuf premiers épisodes de shérif Callie' s Wild West seront disponibles (...)

Hypothesis: The first nine episodes of Sheriff Callie 's Wild West will be available (...)

.\\| Uncertainty and Search

Source: The first nine episodes of Sheriff Callie 's Wild West will be available (...)

log probs:	-4.53	-0.02	-0.28	-0.11	-0.01	-0.001	-0.004	-0.002
------------	-------	-------	-------	-------	-------	--------	--------	--------

Hypothesis: The first nine episodes of Sheriff Callie 's Wild West will be available (...)

.\ | Uncertainty and Search

Copies* make up 2.0% of the WMT14 En-Fr training set,
but are over-represented in the output of beam search

- Among beam hypotheses, copies account for:

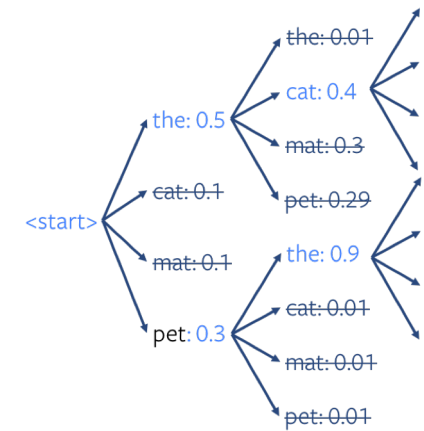
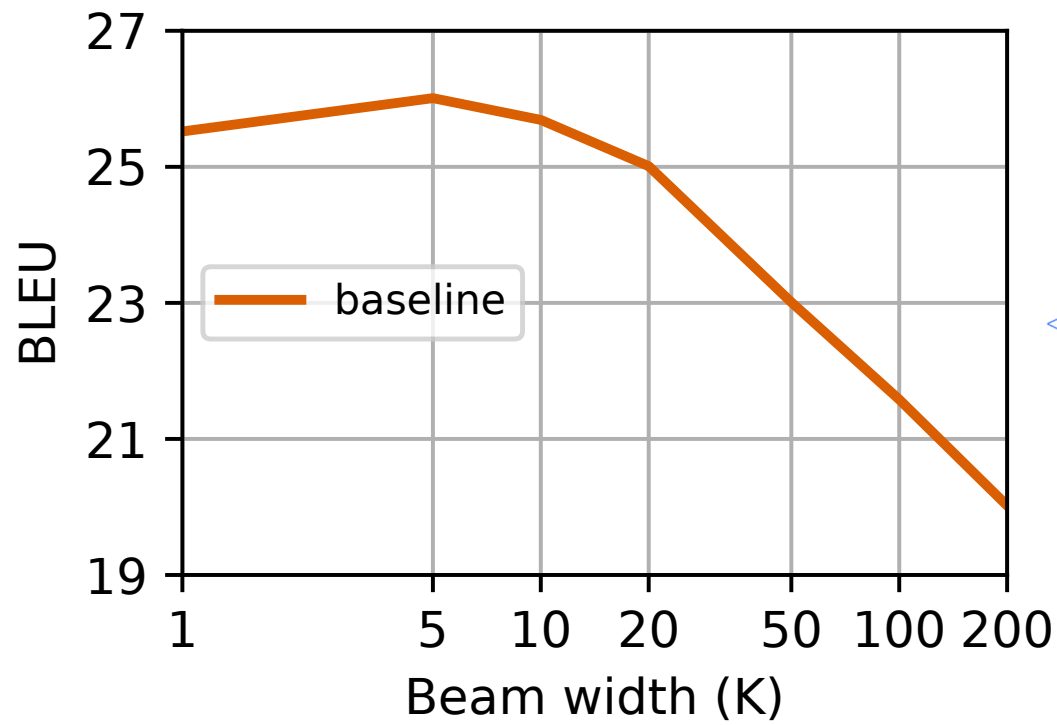
Beam=1: 2.6%

Beam=5: 2.9%

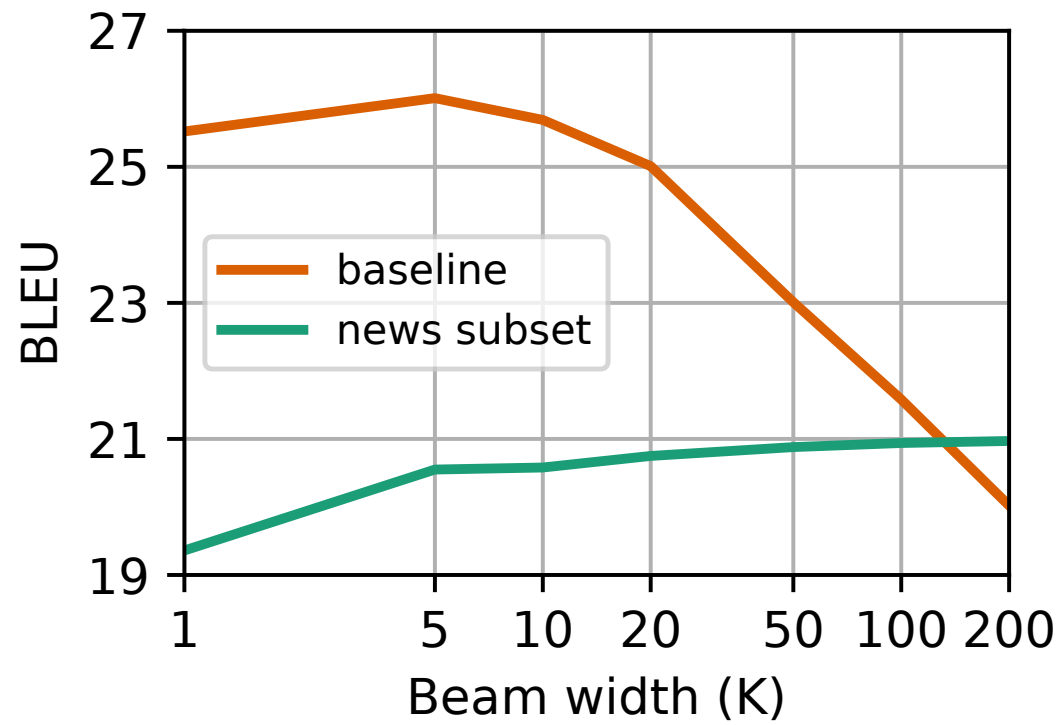
Beam=20: 3.5%

* a copy is a translation that shares
>= 50% of its unigrams with the source

.\\| Uncertainty and Search



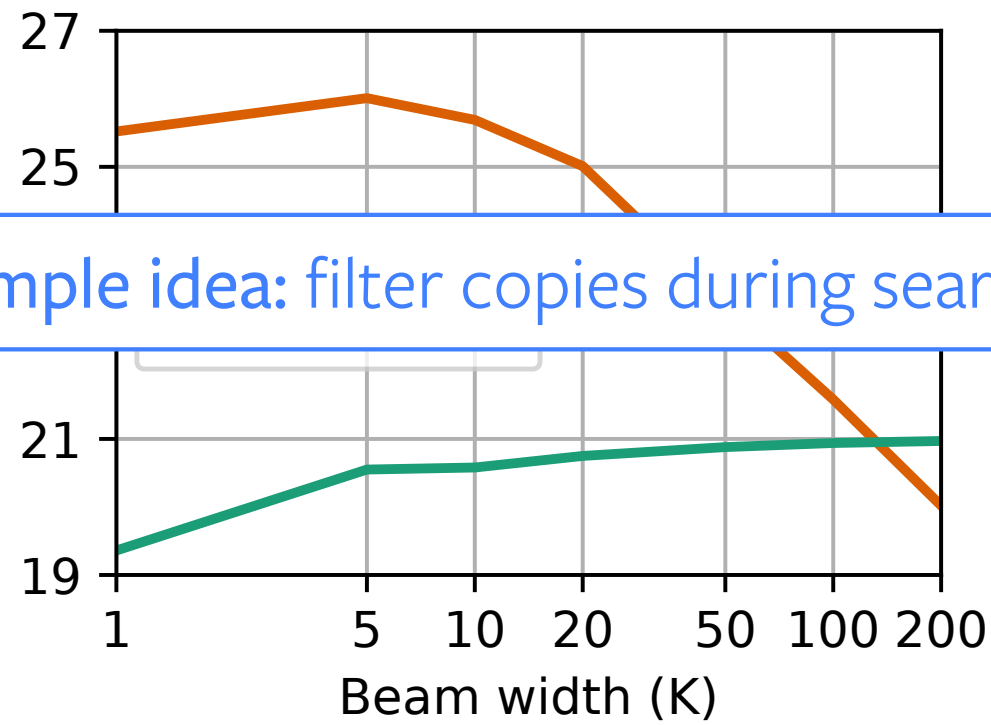
.\| Uncertainty and Search



(WMT17 En-De)

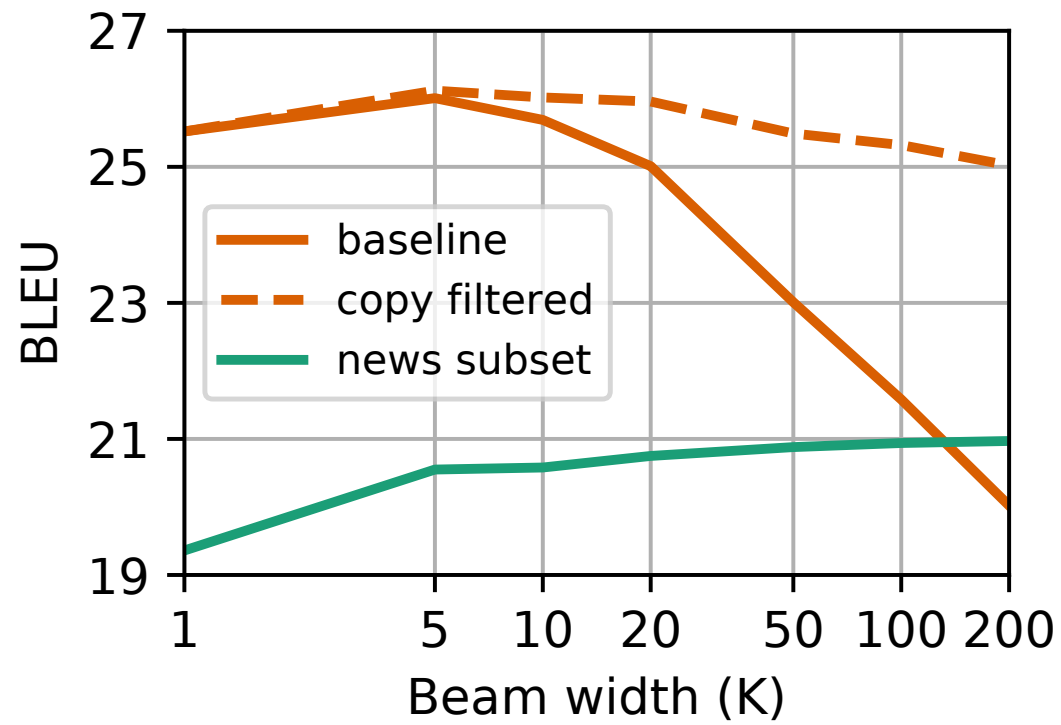
.\\| Uncertainty and Search

A simple idea: filter copies during search



(WMT17 En-De)

.\| Uncertainty and Search



(WMT17 En-De)

.\ | Do NMT models capture uncertainty?

Yes, with interesting effects on search!

Follow-up: How is it represented? Does it match the data distribution?

Challenging because:

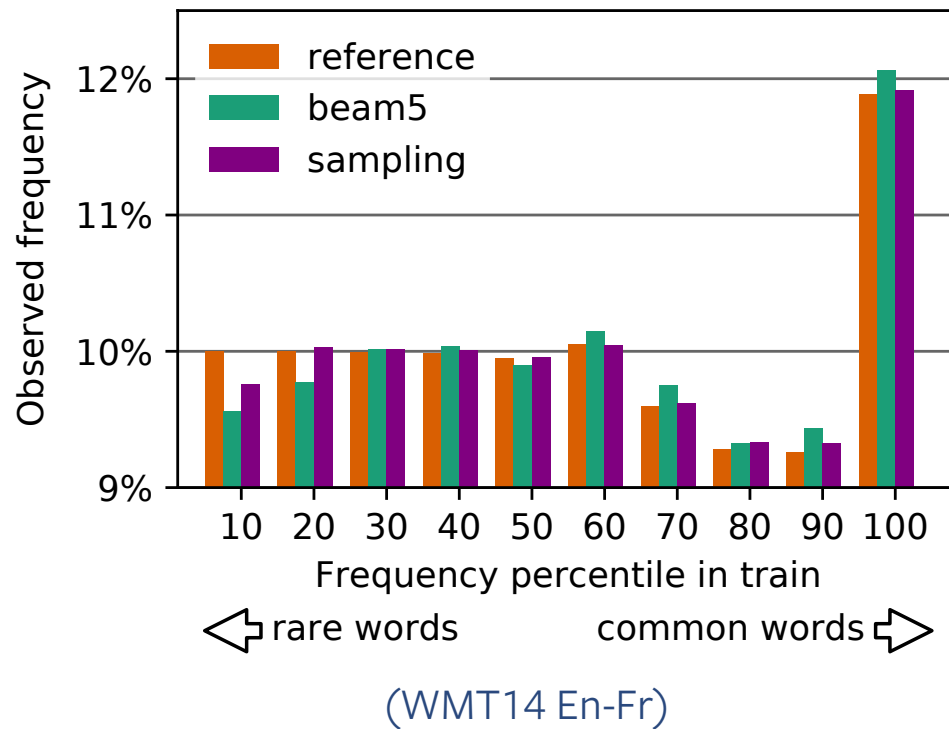
- We typically observe only a single sample from the data distribution for each source sentence (i.e., one reference translation)
- The model distribution is intractable to enumerate

. \ | Necessary matching conditions

What are the necessary conditions for the model distribution to match the data distribution:

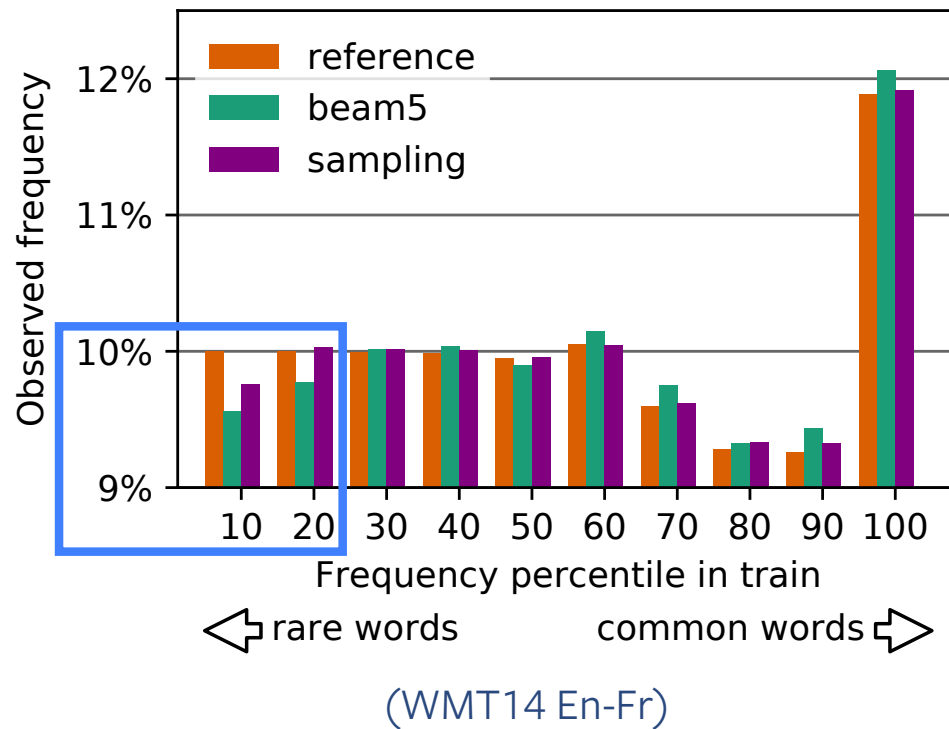
- ...at the **token level**?
- ...at the **sequence level**?
- ...when considering **multiple reference translations**?

.\\ Necessary matching conditions—Token Level



Histogram of unigram frequencies

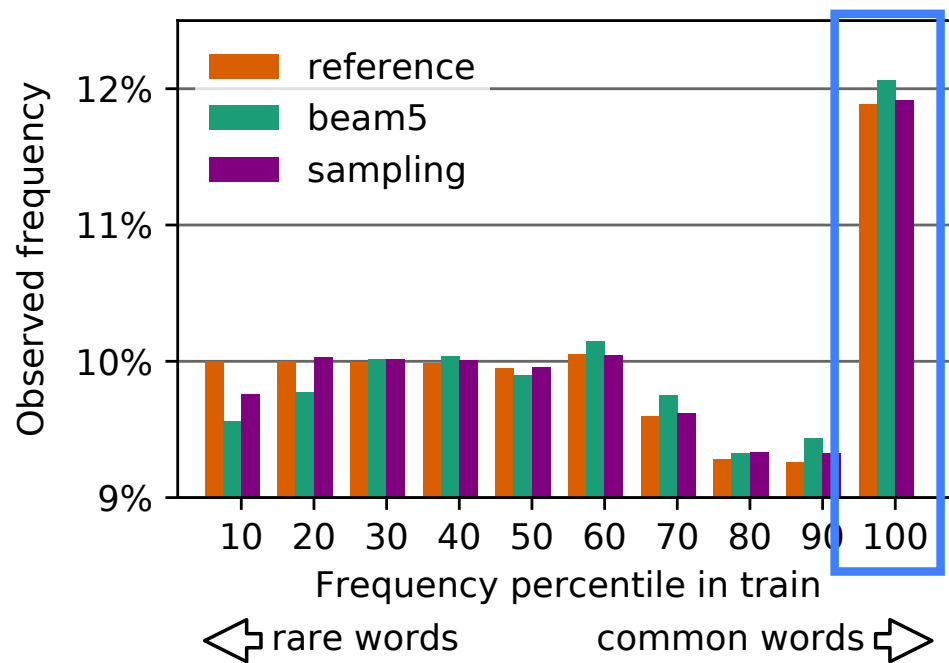
.\\ Necessary matching conditions—Token Level



Histogram of unigram frequencies

Beam under-estimates the rarest words

.\\ Necessary matching conditions—Token Level



Histogram of unigram frequencies

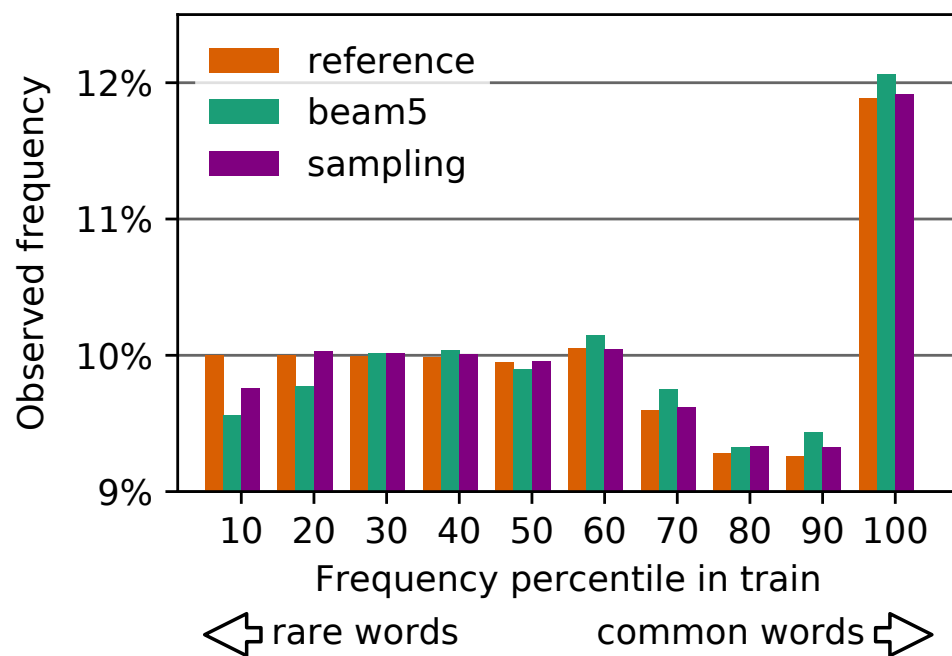
Beam under-estimates the rarest words

Beam over-estimates frequent words.

We should expect this!

(WMT14 En-Fr)

.\\ Necessary matching conditions—Token Level



(WMT14 En-Fr)

Histogram of unigram frequencies

Beam under-estimates the rarest words

Beam over-estimates frequent words.

We should expect this!

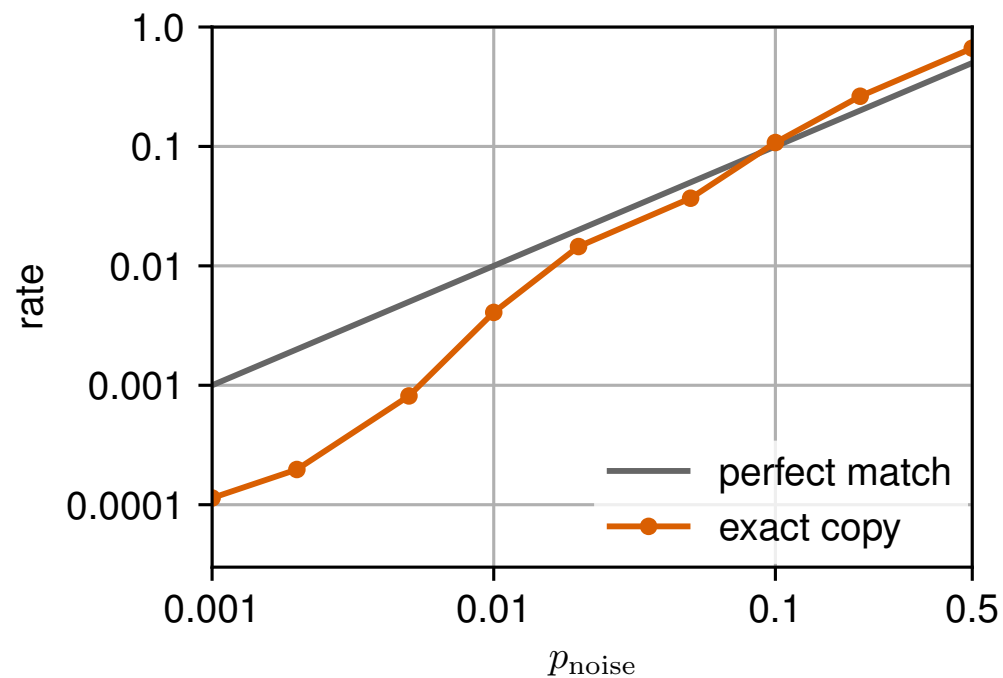
Sampling mostly matches the reference data distribution

. \ | Necessary matching conditions—Sequence Level

Synthetic experiment:

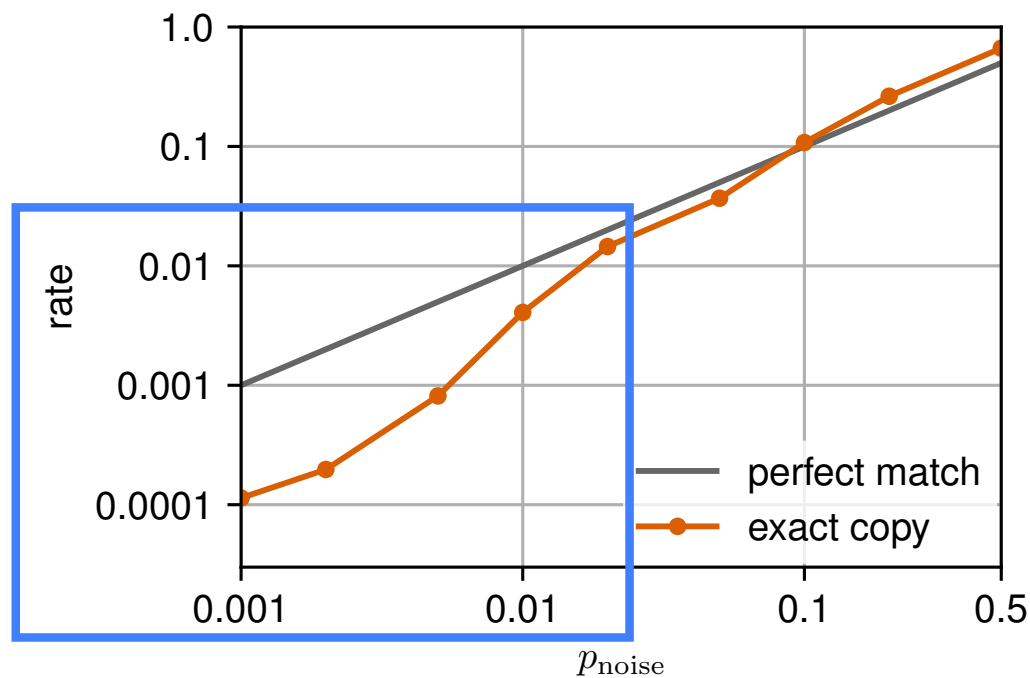
- Retrain model on news subset of WMT, which does not contain copies
- Artificially introduce copies in the training data with probability p_{noise}
- Measure rate of copies among sampled hypotheses

.\\ Necessary matching conditions—Sequence Level



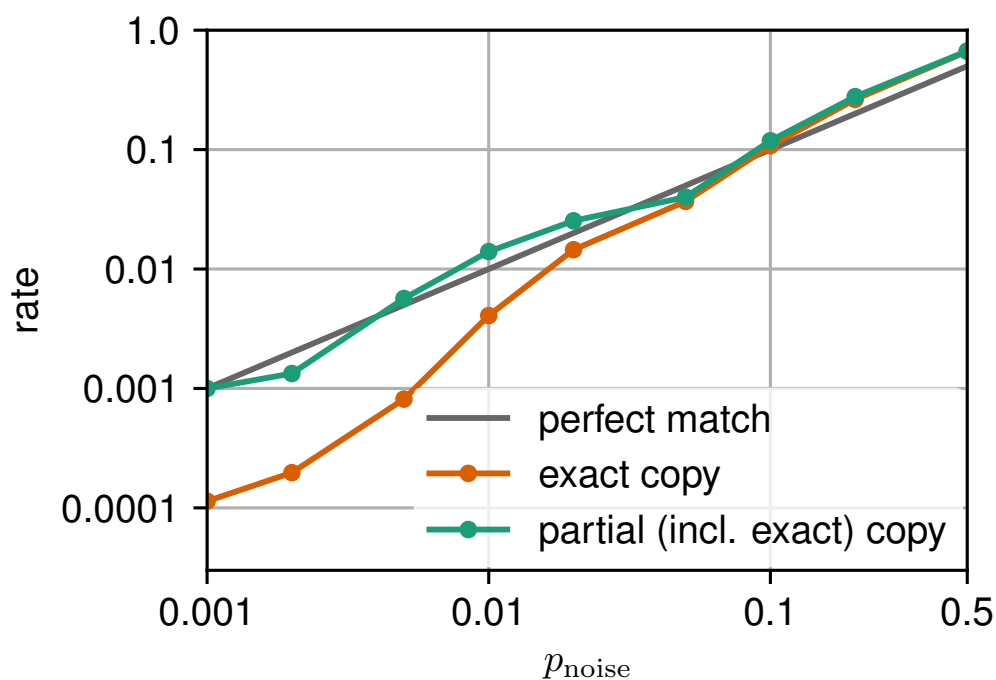
(WMT17 En-De)

.\\ Necessary matching conditions—Sequence Level



Model under-estimates
copies at a sequence level

.\\ Necessary matching conditions—Sequence Level

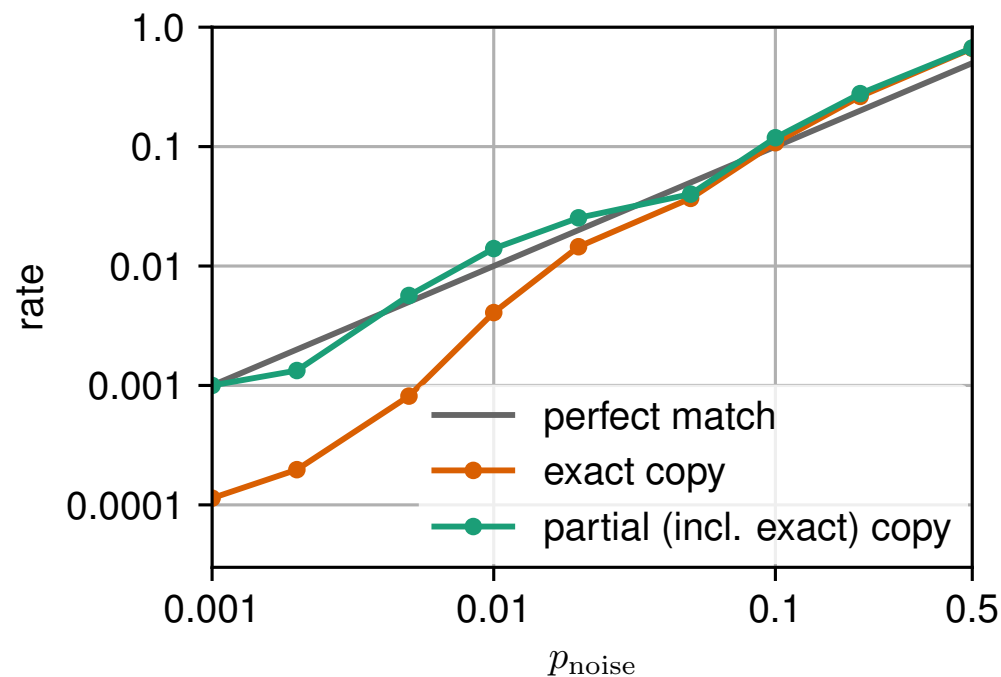


p_{noise} controls rate of exact copies

Partial copies* do not appear in training, yet...

* A partial copy has a unigram overlap of $\geq 50\%$ with the source

.\\ Necessary matching conditions—Sequence Level



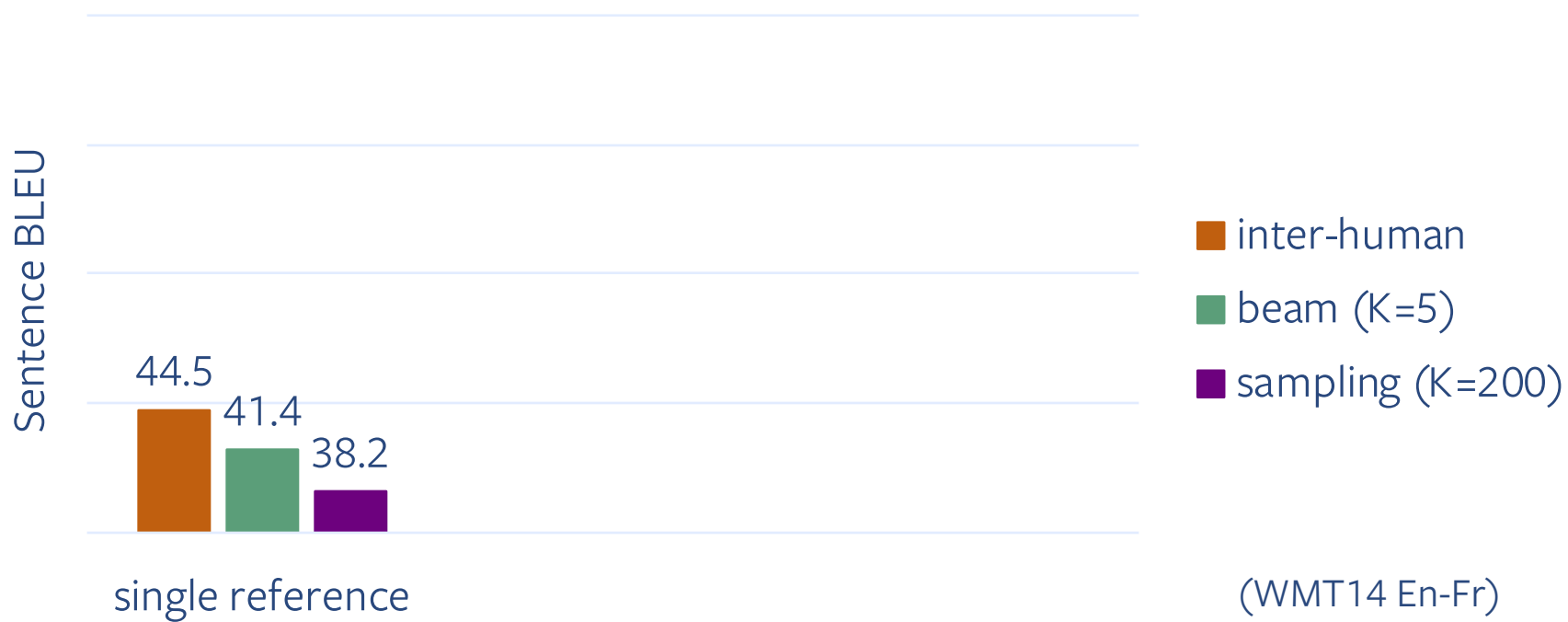
The model smears
probability mass in
hypothesis space!

. \ | Necessary matching conditions—with Mult. References

Question: Can we use BLEU to assess how well the model distribution matches the data distribution?

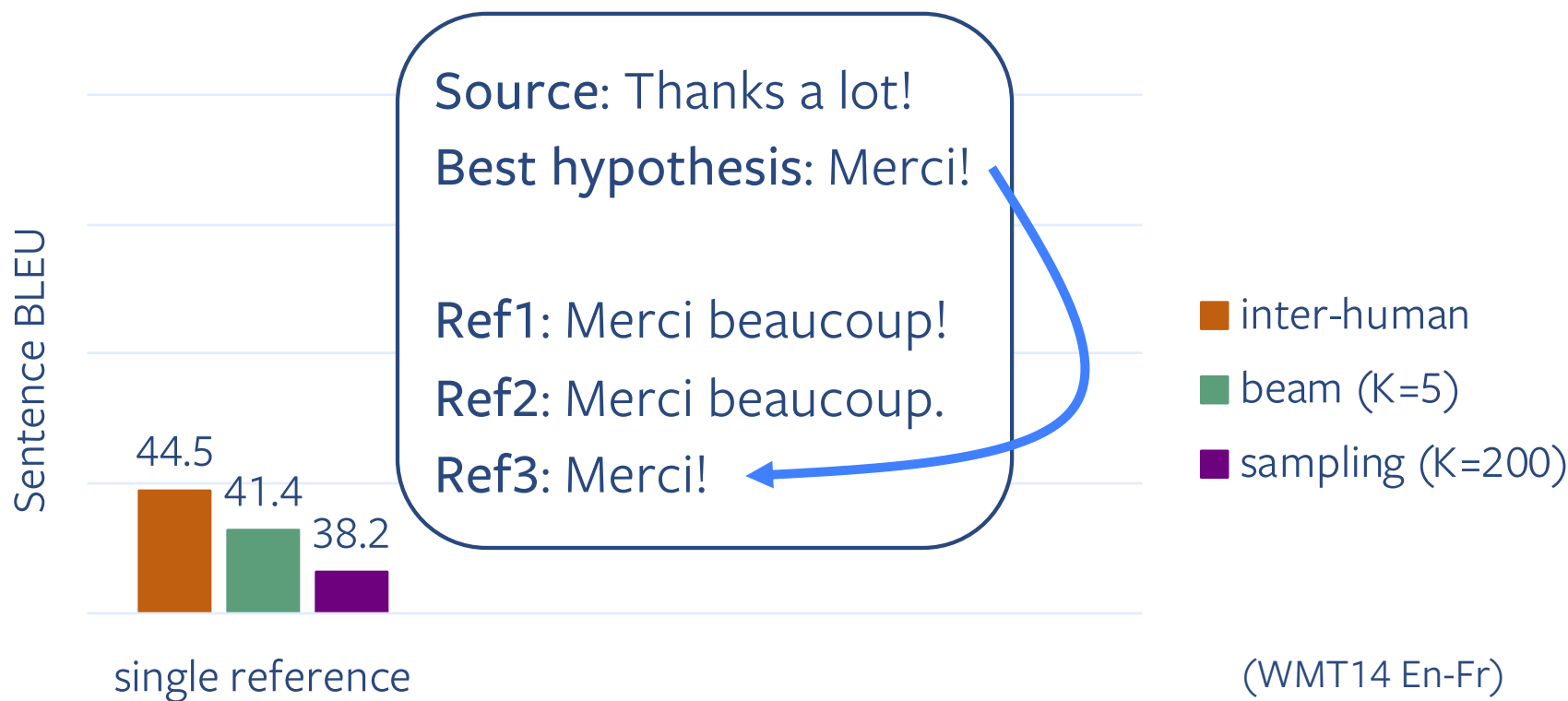
- Collect 10 additional reference translations from distinct human translators
- 500 sentences (En-Fr) and 500 sentences (En-De)
- 10K sentences total
- Available at: github.com/facebookresearch/analyzing-uncertainty-nmt

.\|



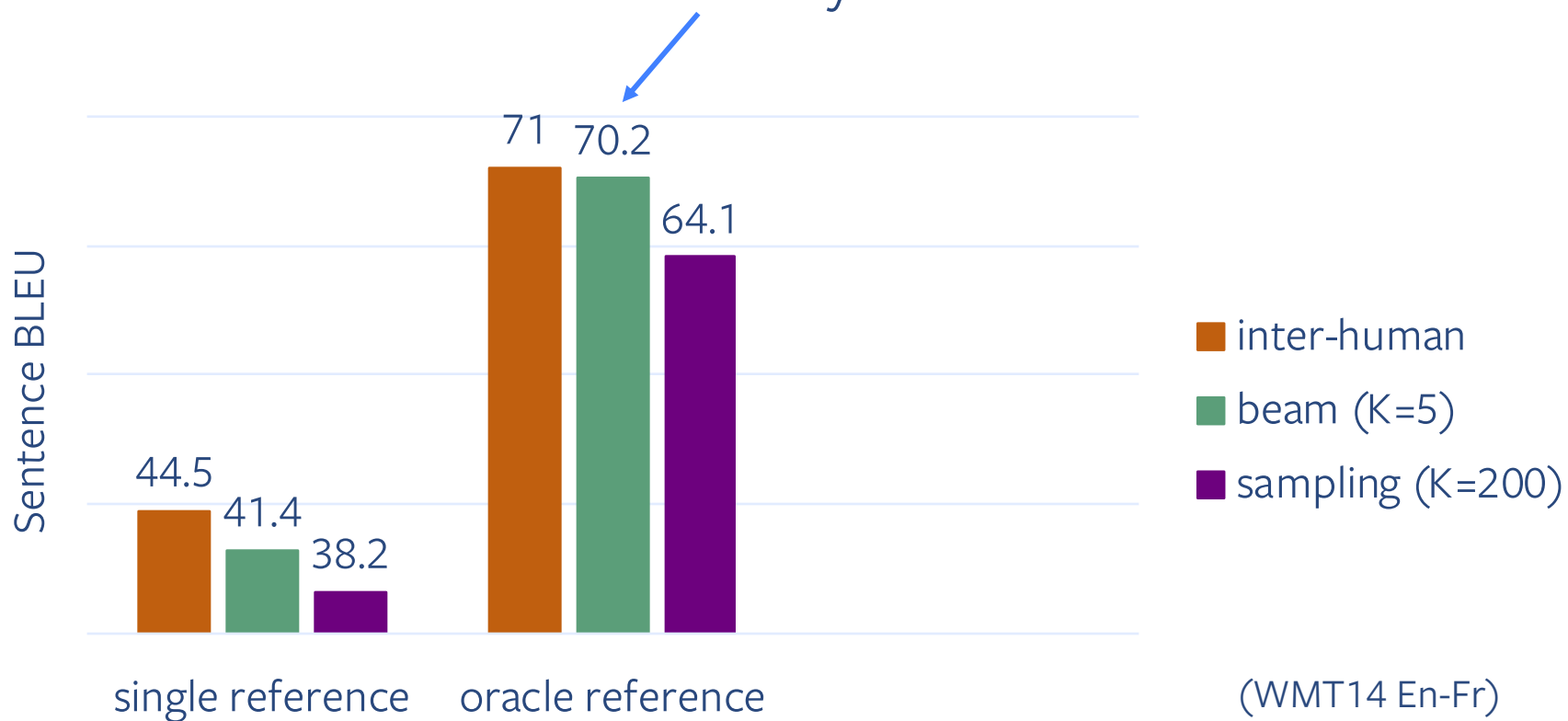
. \ |

oracle reference: BLEU w.r.t. best matching reference



.\|

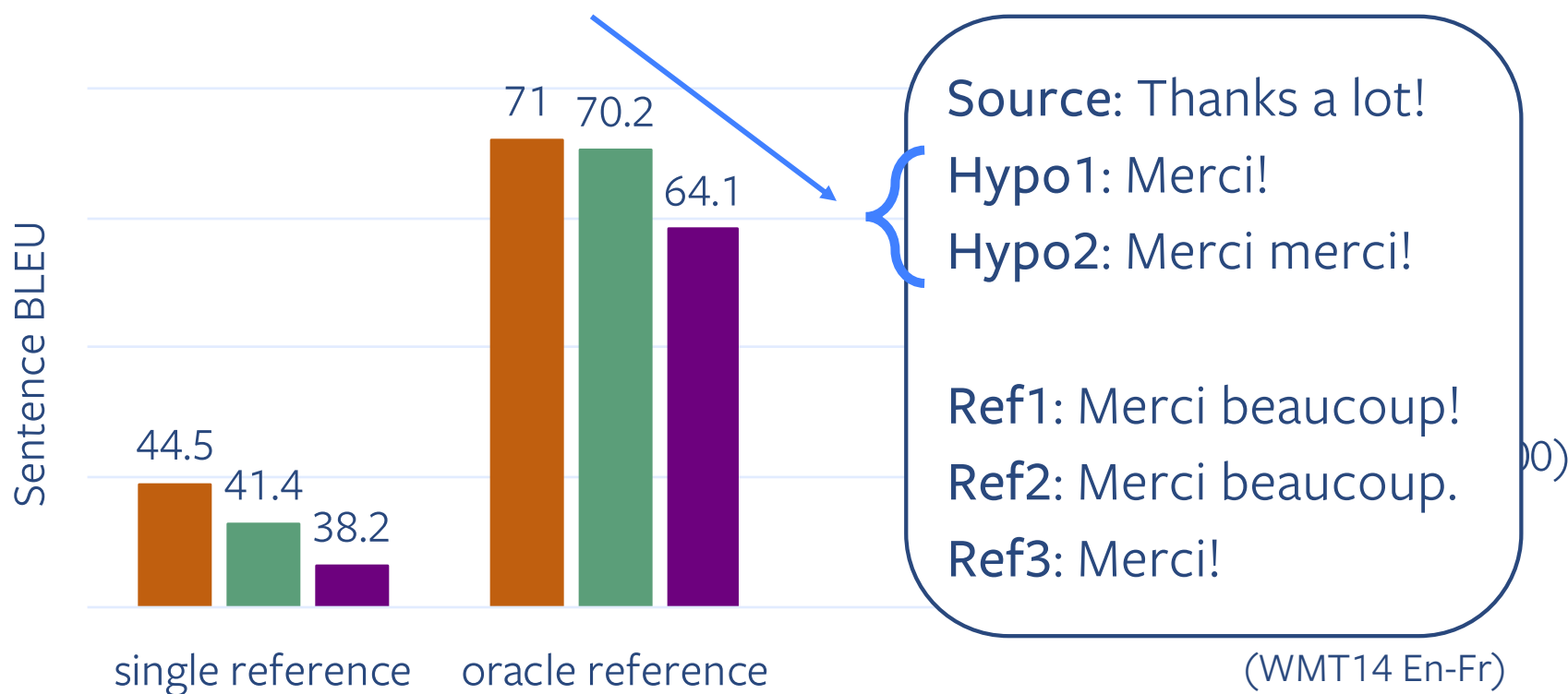
The best **beam** hypothesis
is very close to a reference



. \ |

average oracle:

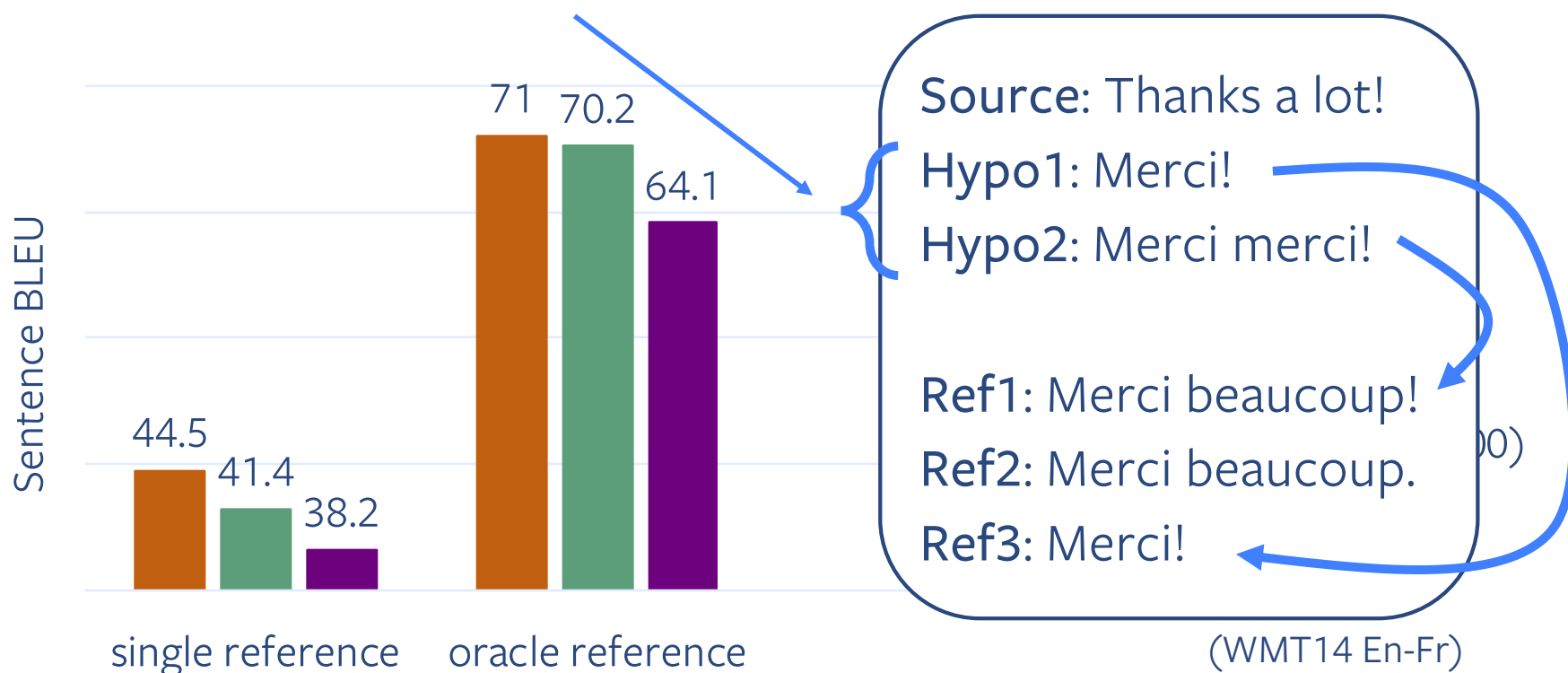
average oracle reference BLEU over top-K hypotheses



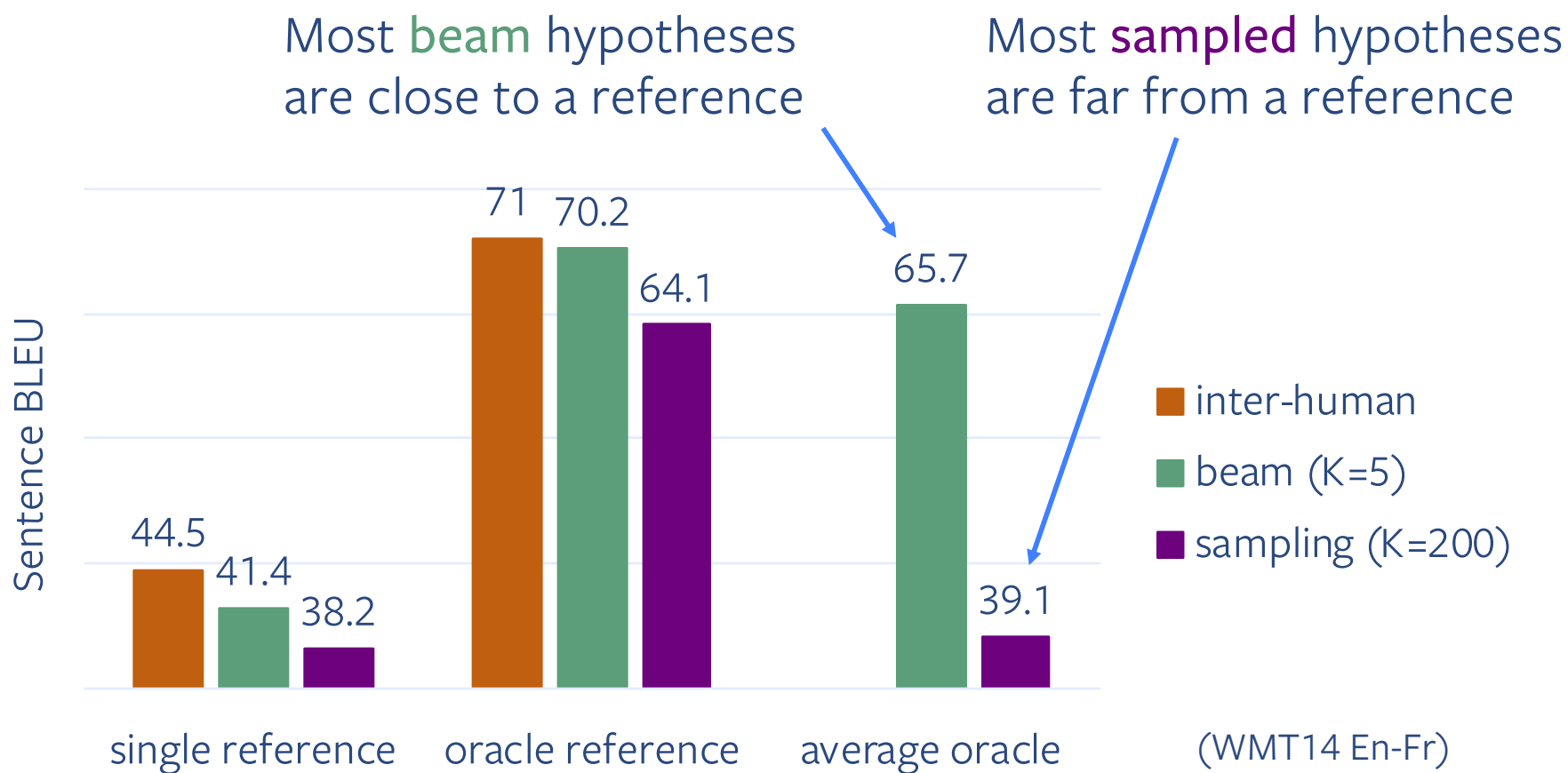
. \ |

average oracle:

average oracle reference BLEU over top-K hypotheses

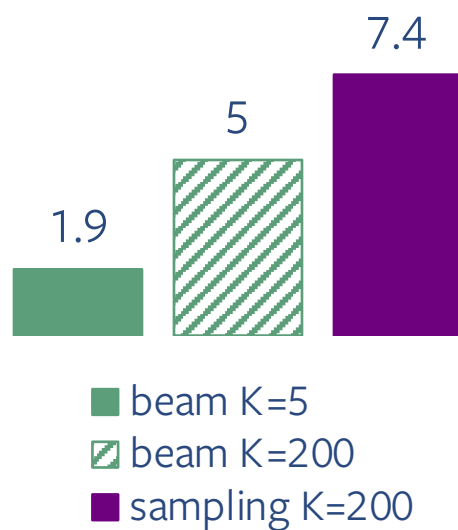


. \ |



.\|

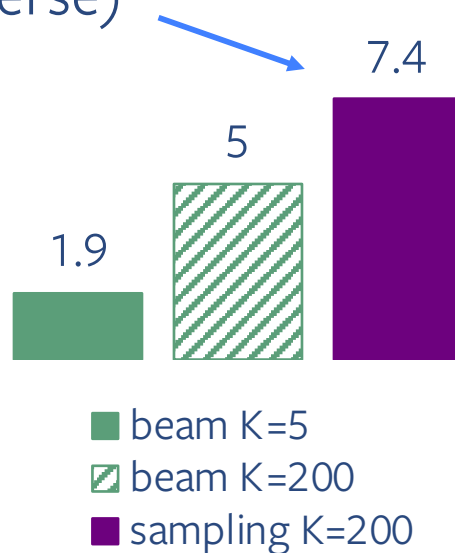
refs covered: number of distinct references
(out of 10) matched to at least one hypothesis



. \ |

refs covered: number of distinct references
(out of 10) matched to at least one hypothesis

Sampling covers more
hypotheses (is more diverse)
than **beam search**



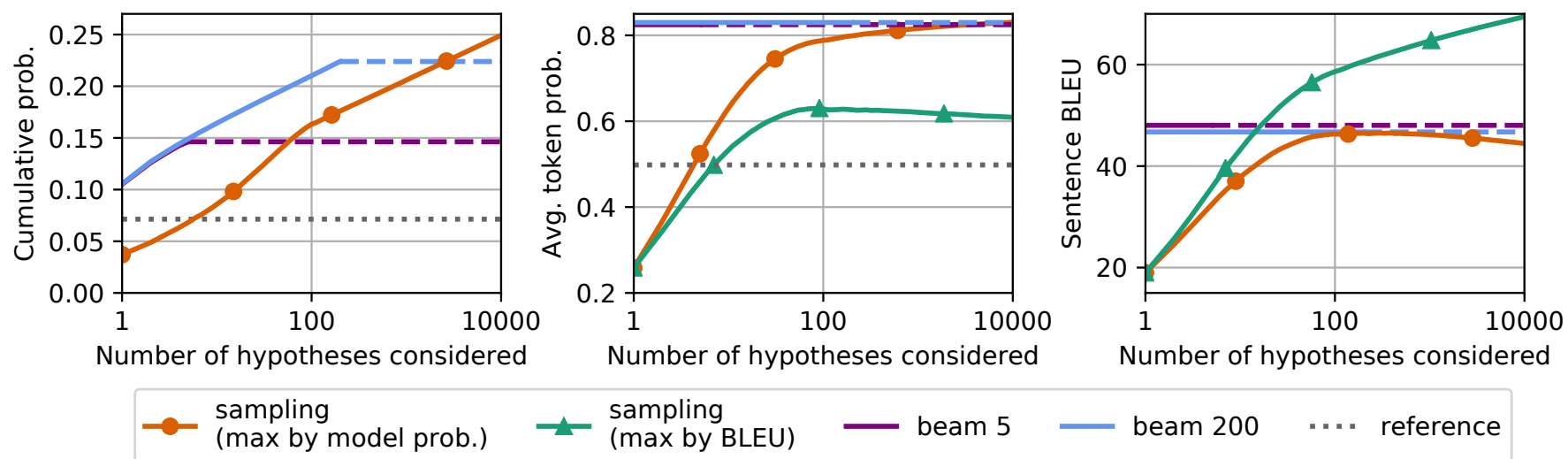
.\ | Conclusion

Poster #163

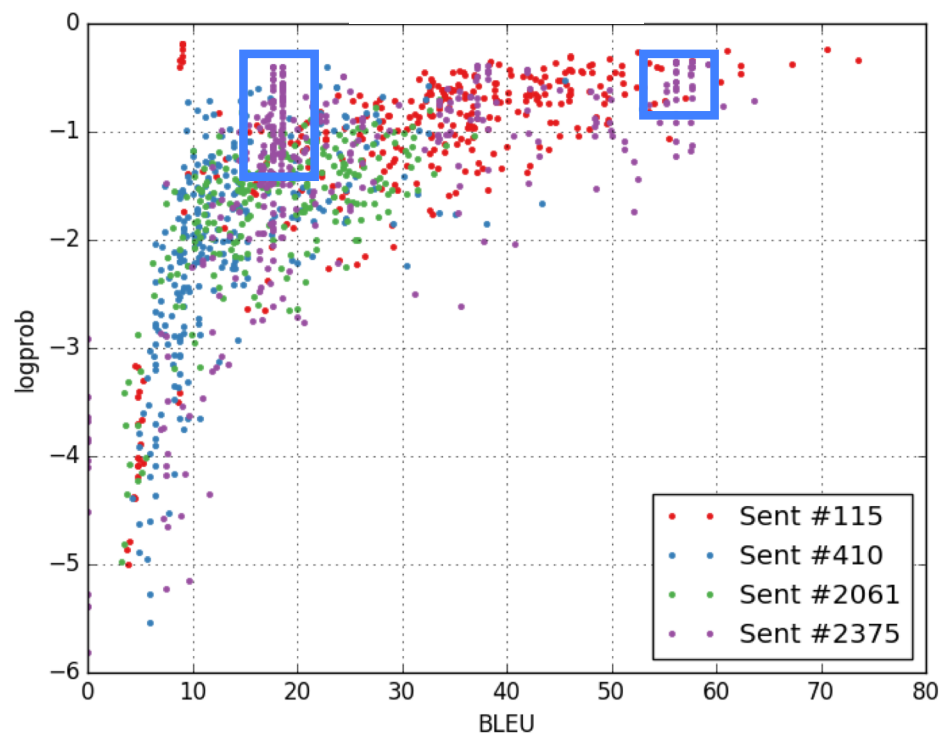
- NMT models capture uncertainty in their output distributions
- Beam search is **efficient** and **effective**, but prefers frequent words
- Degradation with large beams is mostly due to **copying**, but this can be mitigated by **filtering**
- Models are well calibrated at the token level, but **smear probability mass** at the sequence level
- Smearing may be responsible for **lack of diversity** in beam search outputs

Dataset link: github.com/facebookresearch/analyzing-uncertainty-nmt

. \ |



.\|



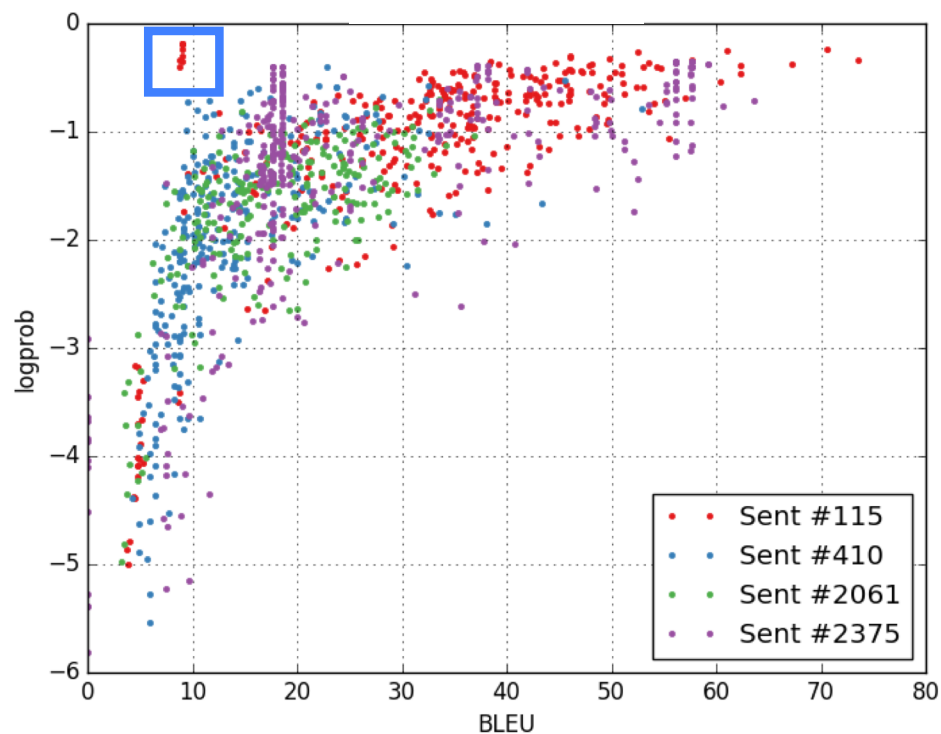
Source: Should this election be decided two months after we stopped voting?

Ref: Cette élection devrait-elle être décidée deux mois après que le vote est terminé?

Low BLEU: Ce choix devrait-il être décidé deux mois après la fin du vote?

High BLEU: Cette élection devrait-elle être décidée deux mois après l'arrêt du scrutin?

. \ |



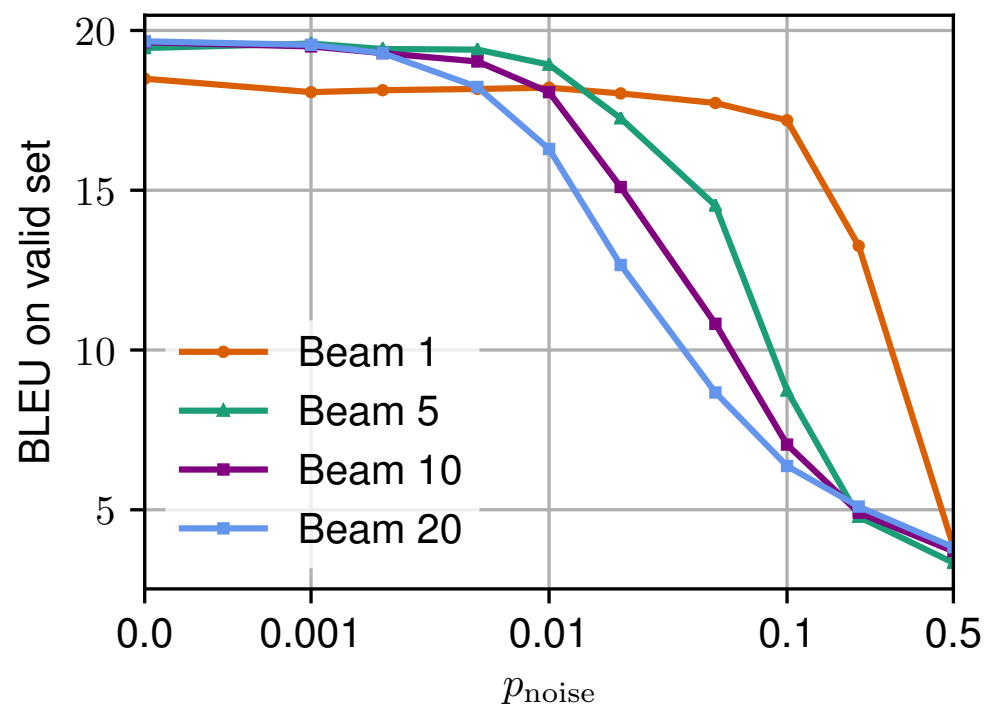
Source: The first nine episodes of Sheriff
<unk> 's Wild West will be available (...)

Ref: Les neuf premiers épisodes de <unk>
<unk> s Wild West seront disponibles (...)

Low BLEU: The first <unk> <unk> of <unk>
<unk> s Wild West will be available (...)

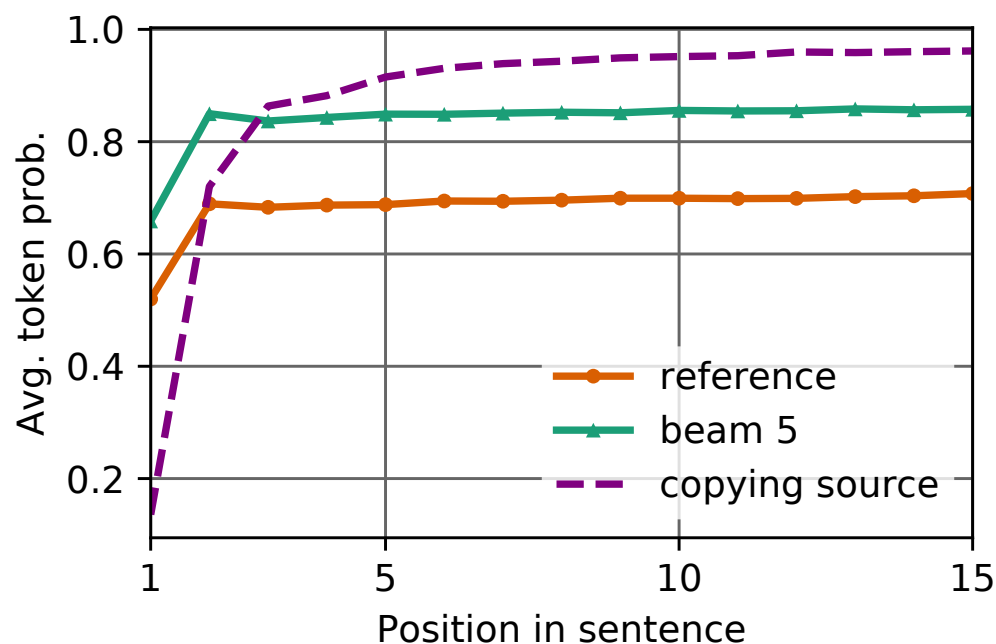
Output is a “copy” in the source language!

. \ |



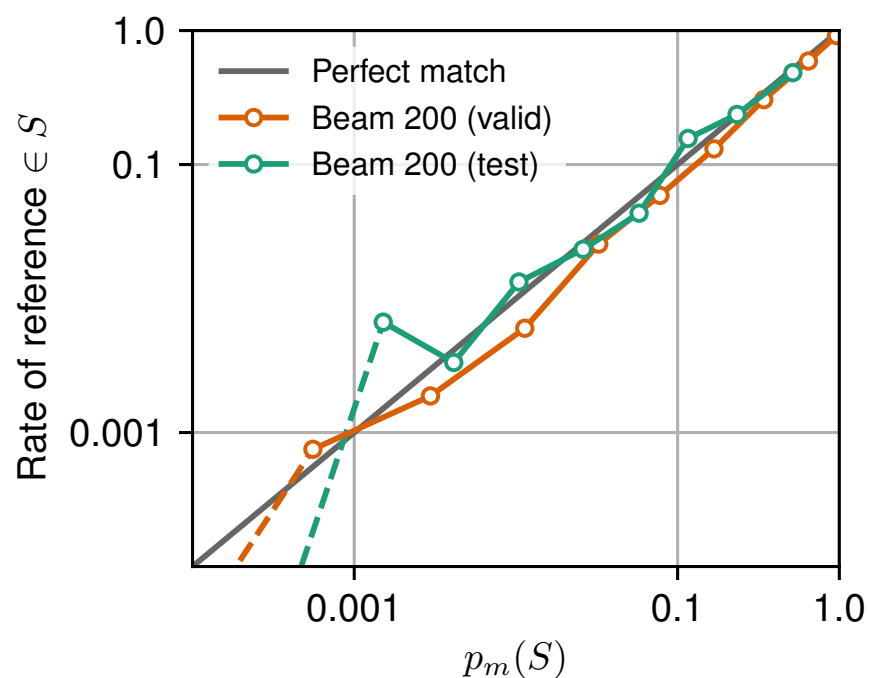
- Train model on news subset of WMT, which does not contain copies
- Artificially introduce copies in training data with p_{noise}
- Small amounts of copy noise lead to a large drop in BLEU for beam $k=20$

. \ |



- During decoding we pay a large penalty for the first copied word
- Subsequently, there is little uncertainty—just continue copying
- Large beams increase chance of reaching the “copy” mode

. \ | Necessary matching conditions—Sequence Level



Set-level calibration

(Guo et al., 2017; Kuleshov & Liang, 2015)

$$\mathbb{E}_{x \sim p_d} [\mathbb{I}\{x \in S\}] = p_m(S)$$

- x-axis: model score of 200 beam hypos
- y-axis: rate at which reference translation is among beam hypos

.\ | Necessary matching conditions—with Mult. References

	beam		sampling
	$k = 5$	$k = 200$	$k = 200$
Prob. covered	4.7%	11.1%	6.7%
Sentence BLEU			
single reference	41.4	36.2	38.2
oracle reference	70.2	61.0	64.1
average oracle	65.7	56.4	39.1
- # refs covered	1.9	5.0	7.4
Corpus BLEU (multi-bleu.pl)			
single reference	41.6	33.5	36.9
10 references	81.5	65.8	72.8