

## HIGHLIGHTS

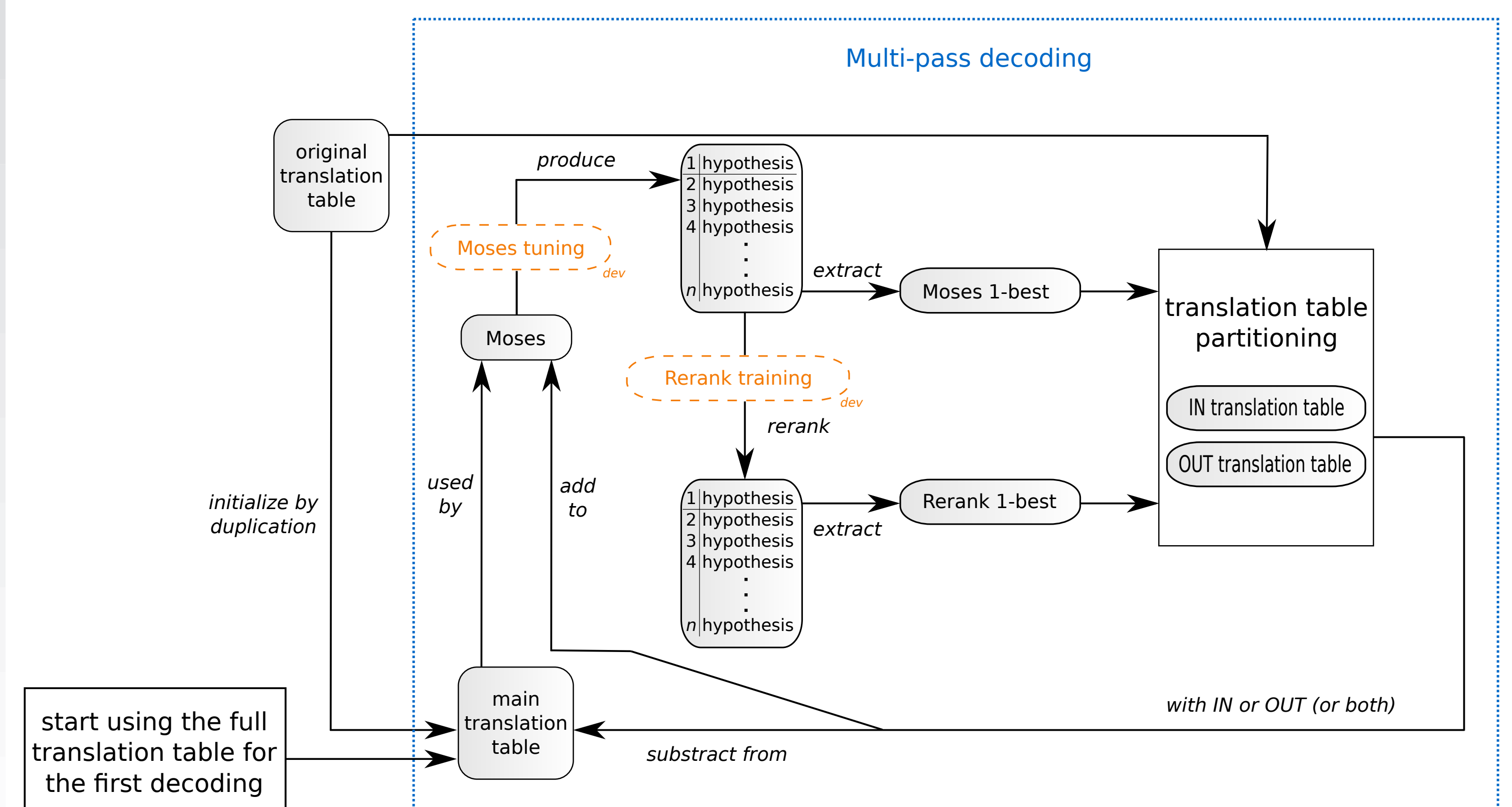
### Using complex features in SMT

- some complex features are not easily integrable during decoding:
  - need of a complete hypothesis (e.g syntactic features)
  - computational cost (e.g neural network models)
  - not available during a first decoding (e.g word posteriors)
- such features are usually used through single-pass reranking

### Main ideas

- exploit a reranking pass result during decoding
- isolate in separate translation tables the possibly misused bi-phrases to better optimize their feature weights

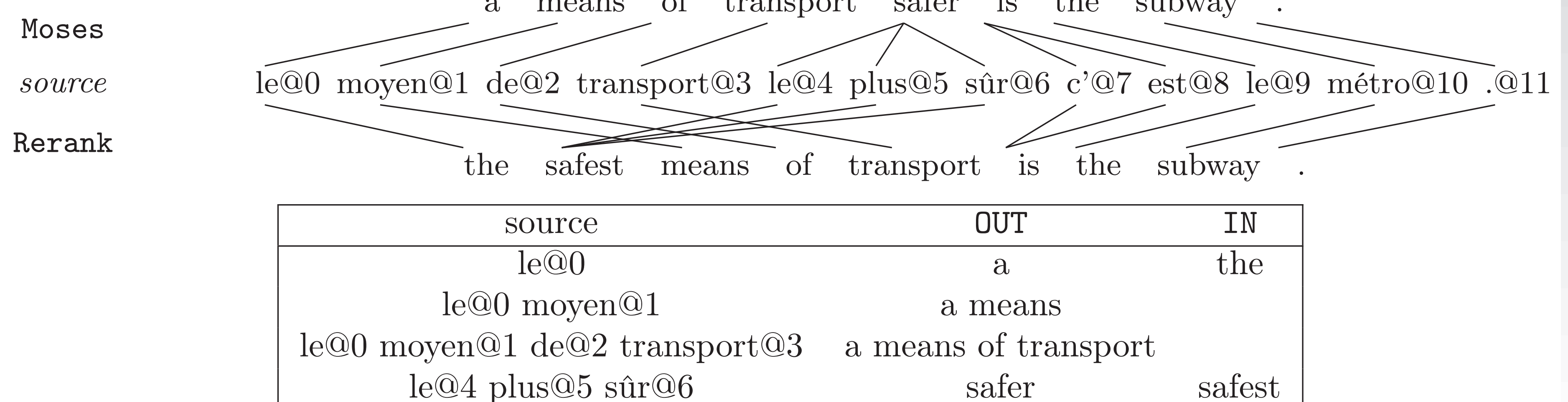
## THE MULTI-PASS DECODING PROCEDURE



## IN AND OUT TRANSLATION TABLES

- **IN**: contains bi-phrases of the reranking 1-best not in the decoder 1-best
- **OUT**: contains bi-phrases of the decoder 1-best not in the reranking 1-best
- extracted bi-phrases are removed from the original translation table
- IN or OUT (or both) are added to the next decoding pass

## TRANSLATION TABLE PARTITIONING WITH LOCATED TOKENS



## EXPERIMENTAL SETTINGS

- Moses tuned with kb-mira
- reranking system (Rerank) trained with kb-mira using the Moses 1,000-best
- complex features used during reranking:

| Features      | medical |       | news  |       |
|---------------|---------|-------|-------|-------|
|               | En→Fr   | Fr→En | En→Fr | Fr→En |
| Moses         | 38.8    | 37.1  | 31.1  | 28.6  |
| + MosesNorm   | 38.9    | 37.2  | 31.1  | 28.7  |
| + NeuralNet   | 41.9    | 38.9  | 32.5  | 29.8  |
| + POSLM       | 39.2    | 37.7  | 31.1  | 28.9  |
| + WPP         | 39.1    | 37.1  | 31.2  | 28.6  |
| + POSTagRatio | 38.9    | 37.3  | 31.1  | 28.8  |
| + Syntax      | 38.8    | 37.2  | 31.2  | 28.9  |
| + IBM1        | 39.1    | 37.2  | 30.9  | 28.8  |
| Rerank        | 42.8    | 40.1  | 32.5  | 29.9  |

## MULTI-PASS DECODING RESULTS FOR ALL CONFIGURATIONS

| Configuration |        | medical En→Fr |         | medical Fr→En |         | news En→Fr  |         | news Fr→En  |         |
|---------------|--------|---------------|---------|---------------|---------|-------------|---------|-------------|---------|
|               |        | test          | # iter. | test          | # iter. | test        | # iter. | test        | # iter. |
| baseline      | Moses  | 38.8          | -       | 37.1          | -       | 31.1        | -       | 28.6        | -       |
|               | Rerank | 42.8          | -       | 40.1          | -       | 32.5        | -       | 29.9        | -       |
| OUT           | Moses  | 41.8          | 4       | 38.7          | 3       | 31.8        | 1       | 29.2        | 1       |
|               | Rerank | 43.8          | 4       | 40.5          | 3       | 32.9        | 1       | 30.3        | 1       |
| IN            | Moses  | 43.2          | 4       | 39.9          | 3       | 32.4        | 2       | 29.3        | 2       |
|               | Rerank | <b>44.2</b>   | 4       | <b>41.0</b>   | 3       | <b>33.2</b> | 2       | <b>30.4</b> | 2       |
| IN and OUT    | Moses  | 42.4          | 4       | 38.7          | 3       | 32.1        | 2       | 29.2        | 2       |
|               | Rerank | 43.5          | 4       | 40.6          | 3       | 32.9        | 2       | <b>30.4</b> | 2       |

- IN alone gives the best results
- "IN and OUT" performs between the configurations using IN and OUT alone
- most configurations converge quickly (between 1 and 4 iterations)

## CONCLUSION & FUTURE WORK

### Conclusion

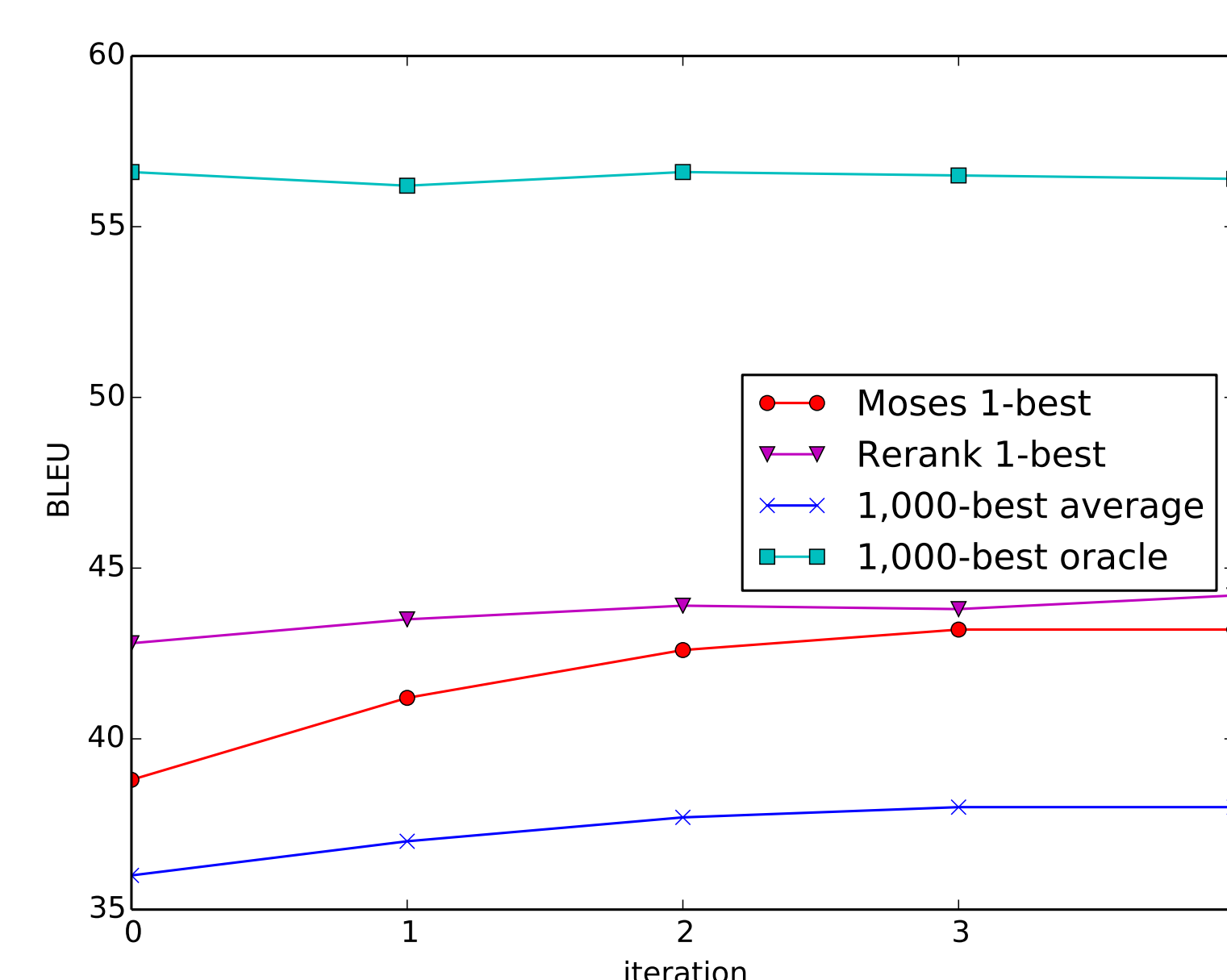
- strong and consistent improvements for all configurations
- simple criterion to iteratively partition the translation table
- makes a better use of complex features than single-pass reranking

### Future work

- add features to the new translation tables to use more informations about the reranking result during decoding
- use the result of a rewriting system (Marie and Max, 2014) to guide the decoder
- enhance the diversity in the  $n$ -best list (Chatterjee and Cancedda, 2010; Gimpel et al., 2013) to train a better reranking system
- add more complex features to the reranking pass

## INCREASING OF THE $n$ -BEST QUALITY

- analysis for the IN configuration (medical En→Fr):



- quickly reduces the gap between Moses and Rerank BLEU scores
- 1,000-best average BLEU increases over the iterations