### Introduction

Since I <u>already programmed</u> the IBM Model 1 for a course on Statistical Machine Translation, the submitted code is just a polished and commented version of it. I am also attaching a draft of the paper *Leveraging Neural Machine Translation for Word Alignment*, which you may find interesting. Please do not share it outside of your team as it is currently in a review process. It covers three topics: (1) using an NMT to get word alignments, (2) various hard alignment extractor methods and (3) combination of all in a small neural network. The extractor methods are in brackets. Hard alignment extractors are reused from this projects and this document describes them very superficially and without mathematical notation. The original version was based on Kohen's pseudocode, which he recently <u>revised</u>. For the graphics I also used <u>Slow Align</u>, a small tool for displaying word alignment (needs major reworking).

The submitted files:

```
img/
 align {a3,a4,int}.svg # illustrations for this document
src/
 aligner.py
                         # main aligner code
 run_eval.py
                         # runs a configuration and evaluates it
                           runs and evaluates fast align configurations
  fastalign.sh
                         # hard alignment extractors, re-used from LeverageAlign
 extractor.py
                         # used for intersection alignments extrensically
 intersector.pv
                         # computes forward and backward alignments and intersects them
 intersect_reverse.py
Leverage_Align.pdf
                         # draft of Levaraging Neural Machine Translation for Word Alignment paper
                         # rust implementation of IBM1
rust/main.rs
```

### **Baseline**

Baseline observed.

Precision = 0.243110 Recall = 0.544379 AER = 0.681684

#### **Custom Models**

## **IBM1 (A0)**

Even though the code uses numpy whenever possible, it still takes considerable amount of time to run on all 100k examples (~10 minutes for 5 steps). This version simply takes the maximum aligned token. This makes a strong assumption that every token is aligned to exactly one other token.

Results for IBM1  $\,/\,$  A0 with 5 EM steps.

### Threshold (A2)

Results for A2(0.35) with 7 EM steps.

```
python3 ./src/aligner.py -n 100000 -e A2 -s 7 | python3 ./jhu-mt-hw/hw2/score-alignments --data jhu-mt-hw/
Precision = 0.685567
Recall = 0.778107
AER = 0.280435
```

## Proportion of best (A3)

This extraction method examines all target tokens and selects the best one. All other alignments which have scores of at least alpha times the maximum one are taken. Here alpha is a parameter. In the following example consider alpha=0.4, then *menu* would align to *Popupmenü* and *Dialogfeld* (mistakenly) and not to others.

Choose Export from the pop-up menu at the bottom of the dialog box .

Wählen Sie im Popupmenü unten im Dialogfeld die Option " Exportieren . "

0.4

Results for A3(0.98) with 7 EM steps.

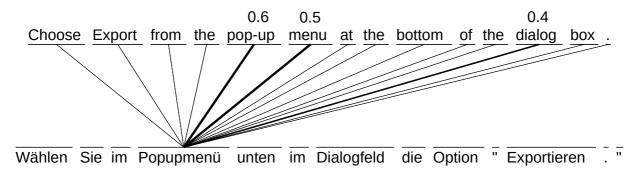
0.2

0.6

python3 ./src/aligner.py -n 100000 -e A3 -s 7 | python3 ./jhu-mt-hw/hw2/score-alignments --data jhu-mt-hw/
Precision = 0.537281
Recall = 0.878698
AER = 0.370400

#### Proportion of reverse best (A4)

The same approach can be done from the target token perspective. Surprisingly, this works a bit better than A3. In the example for alpha=0.5, *Popupmenü* would align to *pop-up*, *menu* and *dialog* (mistakenly).

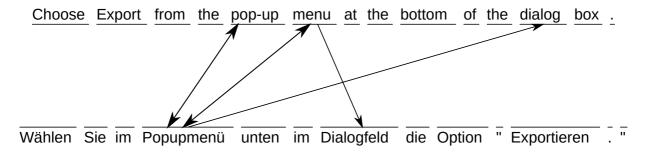


Results for A4(0.98) with 7 EM steps.

python3 ./src/aligner.py -n 100000 -e A4 -s 7 | python3 ./jhu-mt-hw/hw2/score-alignments --data jhu-mt-hw/
Precision = 0.590789
Recall = 0.846154
AER = 0.330601

#### Intersection (A2\*A3\*A4)

As described by Koehn, intersection in alignment directions increases precision. The intersection of different alignment methods also improves the performance. Intersecting the previous two examples leads to *Popupmenü* being aligned to *pop-up* and *menu*.



The main disadvantage of both A3 and A4 is that even with alpha=1, they insist that every token is aligned. Thresholding the alignments improves the performance. Results for A2(0.25)  $\$  A3(0.98)  $\$  A4(0.98) with 7 EM steps.

```
python3 ./src/aligner.py -n 100000 -e A2*A3*A4 -s 7 | python3 ./jhu-mt-hw/hw2/score-alignments --data jhu-
Precision = 0.758483
Recall = 0.784024
AER = 0.231228
```

#### Intersection And Reverse (A2'\*A3'\*A4' \* A2\*A3\*A4)

The final step combines the previous approach by computing the alignment from the other direction and intersecting it as well. In this version only alignments which are within 7 places of each other are considered. This promotes diagonal alignment. A more elegant solution would be to not create this hard boundary, but only increase scores to alignments on the diagonal. The decision process would then be delegated to the extractors.

Results for A2(0.25) \cap A3(0.9) \cap A4(0.9) \cap A2'(0.25) \cap A3'(0.9) \cap A4'(0.9) with 5 EM steps.

```
src/intersect_reverse.sh
Precision = 0.879452
Recall = 0.718935
AER = 0.197724
```

The third sentence of the dataset:

```
Alignment 2 KEY: ( ) = guessed, * = sure, ? = possible
   (*)
(*)
(*)
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```

## fast\_align

The script src/fastalign.sh runs fast\_align with best practice arguments (-v Dirichlet prior and -do close to diagonal). EM steps are default 5.

```
Precision = 0.788122
Recall = 0.831361
AER = 0.196670
```

Alignment of the third sentence in the data by fast\_align. Neither the current implementation nor fast\_align guessed the *ont été fixées* phrase correctly, but fast\_align performed slightly better. This is true even for overall performance, where fast\_align is better by 0.001. I am however sure, that if one would choose better parameters, then both fast\_align and the proposed aligner would produce even better results.

```
Alignment 2 KEY: ( ) = guessed, * = sure, ? = possible
  (*)
(*)
(*)
                                                                               ils
                  ( )
                                                                               sont
                                                                               restreints
                     ( )
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```

Without -d the results are:

Precision = 0.645161 Recall = 0.766272 AER = 0.313448

# **Performance**

I was quite concerned with the implementation speed. So I wrote it in Rust (rust/). The time went from ~10min to ~15s. This is comparable to fast\_align (~14s). The comparison is however not fair from both sides, since the Rust implentation uses only IBM1 EM computation, but on the other fast\_align is multi-threaded.

Only the A0 extraction method was implemented. The results are almost the same as for the IBM1 / A0 model. The small differences are caused by unstable argmax.

Precision = 0.618585 Recall = 0.798817 AER = 0.323890