Combining Unsupervised and Supervised Alignments for MT: An Empirical Study

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Abstract

Word alignment plays a central role in statistical MT (SMT) since almost all SMT systems extract translation rules from word aligned parallel training data. While most SMT systems use unsupervised algorithms (e.g. GIZA++) for training word alignment, supervised methods, which exploit a small amount of human-aligned data, have become increasingly popular recently. This work empirically studies the performance of these two classes of alignment algorithms and explores strategies to combine them to improve overall system performance. We used two unsupervised aligners, GIZA++ and HMM, and one supervised aligner, ITG, in this study. To avoid language and genre specific conclusions, we ran experiments on test sets consisting of two language pairs (Chinese-to-English and Arabic-to-English) and two genres (newswire and weblog). Results show that the two classes of algorithms achieve the same level of MT performance. Modest improvements were achieved by taking the union of the translation grammars extracted from different alignments. Significant improvements (around 1.0 in BLEU) were achieved by combining outputs of different systems trained with different alignments. The improvements are consistent across languages and genres.

1 Introduction

Word alignment plays a central role in training statistical machine translation (SMT) systems since almost all SMT systems extract translation rules from word aligned parallel training data. Until recently, most SMT systems used GIZA++ (Och and Ney, 2003), an unsupervised algorithm, for aligning parallel training data. In recent years, with the availability of human aligned training data, supervised methods (e.g. the ITG aligner (Haghighi et al., 2009)) have become increasingly popular.

The main objective of this work is to show the two classes (unsupervised and supervised) of algorithms are complementary and combining them will improve overall system performance. The use of human aligned training data allows supervised methods such as ITG to more accurately align frequent words, such as the alignments of Chinese particles (e.g. “bei”, “de”, etc) to their English equivalents (e.g. “is/are/was/..”, “of”, etc). On the other hand, supervised methods can be affected by sub-optimal alignments in hand-aligned data. For example, the hand-aligned data used in our experiments contain some coarse-grained alignments (e.g. “lian-he guo” to “United Nations”) although fine-grained alignments (“lian-he” to “United” and “guo” to “Nations”) are usually more appropriate for SMT. Unsupervised methods are less likely to be affected by this problem. We used two well studied unsupervised aligners, GIZA++ (Och and Ney, 2003) and HMM (Liang et al., 2006) and one supervised aligner, ITG (Haghighi et al., 2009) as representatives in this work.

We explored two techniques to combine different alignment algorithms. One is to take the union of the translation rules extracted from alignments produced by different aligners. This is motivated by studies that showed that the coverage of translation rules is critical to SMT (DeNeefe et al., 2007). The
other method is to combine the outputs of different MT systems trained using different aligners. Assuming different systems make independent errors, system combination can generate a better translation than those of individual systems through voting (Rosti et al., 2007).

Our work differs from previous work in two ways. Past studies of combining alternative alignments focused on minimizing alignment errors, usually by merging alternative alignments for a sentence pair into a single alignment with the fewest number of incorrect alignment links (Ayan and Dorr, 2006). In contrast, our work is based on the assumption that perfect word alignment is impossible due to the intrinsic difficulty of the problem, and it is more effective to resolve translation ambiguities at later stages of the MT pipeline. A main focus of much previous work on word alignments is on theoretical aspects of the proposed algorithms. In contrast, the nature of this work is purely empirical. Our system was trained on a large amount of training data and evaluated on multiple languages (Chinese-to-English and Arabic-to-English) and multiple genres (newswire and weblog). Furthermore, we used a state of the art string-to-tree decoder (Shen et al., 2008) to establish the strongest possible baseline. In comparison, experiments in previous studies typically used one language pair and one genre (usually newswire), a reduced amount of training data and a phrase based decoder.

This paper is organized as follows. Section 2 describes the three alignment algorithms. Section 3 describes the two methods used to combine these aligners to improve MT. The experimental setup used to compare these methods is presented in Section 4. Section 5 shows the results including a discussion. Section 6 discusses related work. Section 7 concludes the paper.

2 Alignment Algorithms

We used three aligners in this work: GIZA++ (Och and Ney, 2003), jointly trained HMM (Liang et al., 2006), and ITG (Haghighi et al., 2009). GIZA++ is an unsupervised method based on models 1-5 of Brown et al. (1993). Given a sentence pair e → f, it seeks the alignment a that maximizes the probability $P(f, a|e)$. As in most previous studies using GIZA++, we ran GIZA++ in both directions, from e to f and from f to e, and symmetrized the bidirectional alignments into one, using a method similar to the grow-diagonal-final method described in Och and Ney (2003). We ran GIZA++ up to model 4.

The jointly trained HMM aligner, or HMM for short, is also unsupervised but it uses a small amount of hand-aligned data to tweak a few high level parameters. Low level parameters are estimated in an unsupervised manner like GIZA++.

The ITG aligner is a supervised method whose parameters are tuned to optimize alignment accuracy on hand-aligned data. It uses the inversion transduction grammar (ITG) (Wu, 1997) to narrow the space of possible alignments. Since the ITG aligner uses features extracted from HMM alignments, HMM was run as a prepossessing step in our experiments. Both the HMM and ITG aligners are publicly available.

3 Methods of Combining Alternative Alignments for MT

We explored two methods of combining alternative alignments for MT. One is to extract translation rules from the three alternative alignments and take the union of the three sets of rules as the single translation grammar. Procedurally, this is done by concatenating the alignment files before extracting translation rules. We call this method unioned grammar. This method greatly increases the coverage of the rules, as the unioned translation grammar has about 80% more rules than the ones extracted from the individual alignment in our experiments. As such, decoding is also slower.

The other is to use system combination to combine outputs of systems trained using different aligners. Due to differences in the alignment algorithms, these systems would produce different hypotheses with independent errors. Combining a diverse set of hypotheses could improve overall system performance. While system combination is a well-known technique, to our knowledge this work is the first to apply it to explicitly exploit complementary alignment algorithms on a large scale.

Since system combination is an established technique, here we only briefly discuss our system com-

\[http://code.google.com/p/berkeleyaligner/\]
bination setup. The basic algorithm was described in Rosti et al. (2007). In this work, we use incremental hypothesis alignment with flexible matching (Rosti et al., 2009) to produce the confusion networks. 10-best lists from all systems are collected first. All 1-best hypotheses for each segment are used as confusion network skeletons, the remaining hypotheses are aligned to the confusion networks, and the resulting networks are connected in parallel into a joint lattice with skeleton specific prior probabilities estimated from the alignment statistics on the initial arcs. This lattice is expanded with an unpruned bi-gram language model and the system combination weights are tuned directly to maximize the BLEU score of the 1-best decoding outputs. Given the tuned system combination weights, a 300-best list is extracted from the lattice, the hypotheses are re-scored using an unpruned 5-gram language model, and a second set of system combination weights is tuned to maximize the BLEU score of the 1-best hypothesis of the re-scored 300-best list. The same re-scoring step is also applied to the outputs of individual systems.

4 Experiment Setup

To establish strong baselines, we used a string-to-tree SMT system (Shen et al., 2008), one of the top performing systems in the NIST 2009 MT evaluation, and trained it with very large amounts of parallel and language model data. The system used large sets of discriminatively tuned features (up to 55,000 on Arabic) inspired by the work of Chiang et al. (2009). To avoid drawing language, genre, and metric specific conclusions, we experimented with two language pairs, Arabic-English and Chinese-English, and two genres, newswire and weblog, and report both BLEU (Papineni et al., 2002) and TER (Snover et al., 2006) scores. Systems were tuned to maximize BLEU on the tuning set using a procedure described in Devlin (2009).

The sizes of the parallel training corpora are 238M words (target side) for Arabic-English MT and 265M words for Chinese-English. While the majority of the data is publicly available from the Linguistic Data Consortium (LDC), some of the data is available under the DARPA GALE program. Due to the mixing of different data sources, some test sentences have four reference translations while the rest have only one. The average number of references per test sentence varies across test sets. For this reason, MT scores are not comparable across test sets. Table 1 shows the size and the average number of references per sentence of the test sets.

Two hand-aligned corpora were used to train the ITG aligner: LDC2009E82 (Arabic-English) and LDC2009E83 (Chinese-English). We re-tokenized the corpora using our tokenizers and projected the LDC alignments to our tokenization heuristically. The projection was not perfect and sometimes created very coarse-grained alignments. We used a set of filters to remove such problematic data. We ended up with 3,667 Arabic-English and 879 Chinese-English hand-aligned sentence pairs with sufficient quality for training automatic aligners.
5 Results

Three baseline systems were trained using the three different aligners. Case insensitive BLEU and TER scores for Arabic newswire, Arabic weblog, Chinese newswire, and Chinese weblog are shown in Tables 2, 3, 4, and 5, respectively\(^2\). The BLEU scores on the Test set are fairly similar but the ordering between different alignment algorithms is mixed between different languages and genres. To compare the two alignment combination strategies, we trained a system using the union of the rules extracted from the alternative alignments (union in the tables) and a combination of the three baseline system outputs (3 syscomb in the tables). The system with the unioned grammar was also added as an additional system in the combination marked by 4 syscomb.

As seen in the tables, unioned grammar and system combination improve MT on both languages (Arabic and Chinese) and both genres (newswire and weblog). While there are improvements on both SysCombTune and Test, the results on SysCombTune are not totally fair since it was used for tuning system combination weights and as validation for optimizing weights of the MT systems. Therefore our discussion will focus on results on Test. (We did not show scores on Tune because systems were directly tuned on it.) Statistical significance is determined at 95% confidence level using the bootstrap method described in Koehn (2004), and is only applied on results obtained on the blind Test set.

For unioned grammar, the overall improvement in BLEU is modest, ranging from 0.1 to 0.6 point compared with the best baseline system, with little change in TER score. The improvements in BLEU score are statistically significant for Arabic (both genres), but not for Chinese. The improvements in TER are not significant for either language.

System combination produces bigger improvements in performance. Compared with the best baseline system, the improvement in BLEU ranges from 0.8 to 1.6 point. There are also noticeable improvements in TER, around 1.0 point. The TER improvements are mostly explained by the hypothesis alignment algorithm which is closely related to TER scoring (Rosti et al., 2009). The results are interesting because all three baseline systems (GIZA++, HMM and ITG) are identical except for the word alignments used in rule extraction. The results confirm that the aligners are indeed complementary, as we conjectured earlier. Also, the four-system combination yields consistent gains over the three-system combination, suggesting that the system using the unioned grammar is somewhat complementary to the three baseline systems. The statistical test indicates that both the three and four system combinations are significantly better than the single best alignment system for all languages and genres in BLEU and TER. In most cases, they are also significantly better than unioned grammar.

Somewhat surprisingly, the GIZA++ trained system is slightly better than the ITG trained system on all genres but Chinese weblog. However, we should point out that such a comparison is not entirely fair. First, we only ran ITG on short sentences. (For long sentences, we had to settle for HMM alignments for computing reasons.) Second, the hand-aligned data used for ITG training are not very clean, as we said before. The ITG results could be improved if these problems were not present.

\(^2\)Dagger (†) indicates statistically better results than the best individual alignment system. Double dagger (‡) indicates statistically better results than both best individual alignment and unioned grammar. Bold indicates best Test set performance among individual alignment systems.
### 5.1 Discussion

Inter-aligner agreements provide additional evidence about the differences between the aligners. Suppose on a common data set, the sets of alignment links produced by two aligners are $A$ and $B$, we compute their agreement as \( \frac{|A \cap B|}{|A| + |A \cap B|/|B|} \). (This is the average of recall and precision of one set by treating the other set as reference.) The agreement between GIZA++ and ITG is around 78% on a subset of the Arabic-English parallel data. The agreements between GIZA++ and HMM, and between HMM and ITG are slightly higher, around 83%. Since ITG could not align long sentences, we only used short sentences (at most 60 words in length) in our calculation.

Due to the large differences between the aligners, significantly more rules were extracted with the unioned grammar method in our experiments. On average, the size of the grammar (number of rules) was increased by about 80% compared with the baseline systems. The larger grammar results in more combinations of partial theories in decoding. However, for computing reasons, we kept the beam size of the decoder constant despite the increase in grammar size, potentially pruning out good theories. Performance could be improved further if larger beam sizes were used. We will leave this to future work.

### 6 Related Work

Ayan and Dorr (2006) described a method to minimize alignment errors by combining alternative alignments into a single alignment for each sentence pair. Deng and Zhou (2009) used the number of extractable translation pairs as the objective function for alignment combination. Och and Ney (2003) and Koehn et al. (2003) used heuristics to merge the bidirectional GIZA++ alignments into a single alignment. Despite differences in algorithms and objective functions in these studies, they all attempted to produce a single final alignment for each sentence pair. In comparison, all alternative alignments are directly used by the translation system in this work.

The unioned grammar method in this work is very similar to Giménez and Márquez (2005), which combined phrase pairs extracted from different alignments into a single phrase table. The difference
from that work is that our focus is to leverage complementary alignment algorithms, while theirs was to leverage alignments of different lexical units produced by the same aligner.

Some studies leveraged other types of differences between systems to improve MT. For example, de Gispert et al. (2009) combined systems trained with different tokenizations.

The theory behind the GIZA++ aligner was due to Brown et al. (1993). The theory of Inversion Transduction Grammars (ITG) was due to Wu (1997). The ITG aligner (Haghighi et al., 2009) used in this work extended the original ITG to handle blocks of words in addition to single words. The use of HMM for word alignment can be traced as far back as to Vogel et al. (1996). The HMM aligner used in this work was due to Liang et al. (2006). It refined the original HMM alignment algorithm by jointly training two HMMs, one in each direction. Furthermore, it used a small amount of supervised data to tweak some high level parameters, although it did not directly use the supervised data in training.

7 Conclusions

We explored two methods to exploit complementary alignment algorithms. One is to extract translation rules from all alternative alignments. The other is to combine outputs of different MT systems trained using different aligners. Experiments on two language pairs and two genres show consistent improvements over the baseline systems.

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