Fully Automatic Compilation of Portuguese-English and Portuguese-Spanish Parallel Corpora

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Abstract. This paper reports the fully automatic compilation of parallel corpora for Brazilian Portuguese. Scientific news texts available in Brazilian Portuguese, English and Spanish are automatically crawled from a multilingual Brazilian magazine. The texts are then automatically aligned at document- and sentence-level. The resulting corpora contain about 2,700 parallel documents totaling over 150,000 aligned sentences each. The quality of the corpora and their usefulness are tested in an experiment with machine translation.

1. Introduction

Parallel corpora are collections of texts that are mutual translations. Machine Translation (MT) systems have recently become very popular due to the latest advances in the field of Statistical Machine Translation (SMT). The most advanced SMT approaches, such as phrase-based [Koehn et al. 2003] and hierarchical SMT [Chiang 2005], learn how to perform translation from parallel corpora.

A popular way of building a parallel corpus is collecting large amounts of data from the web. The alignment between documents, sentences and words allows the development of resources such as dictionaries and SMT systems.

[Esplà-Gomis and Forcada 2010] present Bitextor, a language-independent tool to harvest parallel documents from multilingual web sites. It uses features from the HTML structure of the documents and their URLs to perform document alignment. [Aziz et al. 2008] describe the creation of a Brazilian Portuguese-Spanish (pt-es) parallel corpus of scientific texts. The corpus is manually crawled from the Brazilian magazine Pesquisa FAPESP Online¹ and the document alignment is manually performed, followed by automatic sentence alignment using the Translation Corpus Aligner (TCA) method from [Hofland 1996]. The same magazine and methodology were used to create a Brazilian Portuguese-English (pt-en) corpus. Both pt-es and pt-en corpora are freely available² and will be referred to as Fapesp-v1. Despite small (~18K sentences, the Fapesp-v1 corpora have been used in interesting applications: rule-based MT [Caseli et al. 2006], SMT [Aziz et al. 2009] and coreference resolution [Souza and Orasan 2011].

In this paper we present a sentence-aligned parallel corpora (pt-en and pt-es) using all the currently available issues of the magazine Pesquisa FAPESP Online. Unlike in [Aziz et al. 2008], our compilation is fully automatic from the data crawling to the sentence alignment. Unlike Bitextor, data crawling is fit to our type of text, resulting in very good recall and precision at document-level alignment with surface-based features.

¹http://revistapesquisa.fapesp.br/
²www.nilc.icmc.usp.br/lacioweb/english/corpora.htm
2. Methodology

The compilation of the corpora is divided in three steps: data crawling, document alignment and sentence alignment, as described in the following sections.

2.1. Data Crawling

The source for our corpora is the Brazilian scientific magazine *Pesquisa FAPESP Online*, which contains parallel electronic documents in three languages: Brazilian Portuguese (original release), English and Spanish (human translations). The magazine currently contains approximately 2,700 parallel articles and it is monthly updated.

We use GNU wget\(^3\) and a URL template to download HTML pages from the magazine’s website. For a given issue we first download its index, a single page containing links to articles, and then parse those links and download the actual articles. Issues are identified by a sequential number that is consistent across the different versions of the website (original, English and Spanish). Therefore the alignment of issues is straightforward. However, there is no obvious correspondence between articles’ identifiers, and therefore content-based document alignment techniques need to be applied. The crawling took about 1 hour and resulted in 3,658 original articles (233,490 sentences), 2,740 English translations (184,587 sentences) and 2,750 Spanish translations (180,525 sentences). Hereafter we will refer to this compilation as Fapesp-v2.

2.2. Document Alignment

Assuming the existence of a bilingual dictionary (take pt–en as example), we use a very simple alignment technique:

1. For every issue we gather its pt and en articles, lowercase and tokenize them;
2. For every pt document we replace the original words by their n-best translations according to our bilingual dictionary;
3. We represent documents as distributional profiles (DP), that is, a vector that contains a word \(w\) and its frequency \(f\) in the document: profile : \(w \rightarrow f\);
4. For every possible document pair we compute the cosine similarity of their DPs.
5. We align each en document to its best pt candidate if the similarity score is above a given threshold.

We compiled bilingual dictionaries using the small corpus Fapesp-v1 running 5 iterations of IBM Model 1 [Brown et al. 1993]. IBM Model 1 estimates a word-based translation probability distribution from a sentence aligned corpus. We set the number of best translations from the dictionary to 3 and the similarity metric threshold to 0.15 after a short round of evaluation described in Section 3. The same procedure was used to align pt–es articles. We obtained 2,675 pt–en and 2,668 pt–es parallel texts.

2.3. Sentence Alignment

We use an implementation of TCA specially written for aligning Brazilian Portuguese to a foreign text [Caseli and Nunes 2003]. While the most desired alignment type is the substitution (1-1), i.e., one source sentence aligned to one target sentence, other alignment types occur as consequence of different translation decisions. 85% of the alignments produced

\(^3\)www.gnu.org/software/wget/
are substitutions. For the remaining cases, the following heuristics were designed (based on manual inspection) to convert as many as possible cases into valid 1-1 substitutions. Figure 1 presents examples of the application of these heuristics.

**H1:** A deletion followed by an insertion, or vice-versa, many times result from a wrong decision of the automatic aligner, in reality the two alignment sets should be a single 1-1 alignment.

**H2:** Alignments of the type 1-2 or 2-1 usually result from incorrect sentence splitting or different translation decisions. For instance, a text separated by a colon in the source text is separated by a period in the target text. If two chunks of one sentence, separated by a punctuation mark, independently align to each of the two sentences in the other language, the original alignment is replaced by two 1-1 alignments. We align the chunks using the same metric used to align documents.

| H1 | Uma hiptese na contram | [A hypothesis on the wrong side of the road] |
|----|------------------------|-------------------------------------------|
| H2 | [So peixes que, como os linguados, dos quais so aparentados, tm os dois olhos e o colorido do corpo de um lado s - e assim conseguem se camuflar]; [“Vivem enterrados na area e os predadores no os enxergam”, relata Menezes]; | [They are fish that, like flounders, to which they are related, have their two eyes and the body colored on one side only – and thus succeed in camouflaging themselves]; [“They live buried in the sand, and their predators do not see them”, Menezes reports]; |

**Figure 1. Examples of the application of our alignment heuristics**

In our experiments with pt-en, for example, ~10K sentence pairs were generated using the conversion heuristics. After the application of the heuristics, all the remaining non-substitution alignments sets (3% of the total) were removed from the corpus, since these are likely to contain inconsistencies. The resulting Fapesp-v2 corpora contain 156,712 pt-en and 152,238 pt-es sentence pairs.

### 3. Experiments

To assess the quality of the corpus, we performed intrinsic (pt-en) and extrinsic evaluations (pt-en and pt-es). Starting with the document alignment strategy, we manually checked 310 pairs of pt-en texts. A cosine threshold of 0.15 guaranteed that 97.66% of the en documents were aligned to pt ones with 99.35% accuracy.

For the pt-en sentence alignment, we assessed 900 substitution (1-1) alignments produced by the TCA aligner and 300 resulting from the conversion heuristics. Results showed that 98% of the original substitution alignments were correct and that the heuristics H1 and H2 achieved an accuracy of 95.91% and 82%, respectively.

For both document and sentence alignment, we expect the results for the pt-es corpus to be similar or even more positive, given the proximity of the language pair.

#### 3.1. Phrase-based SMT

To assess the utility of the corpora in an external task, we used them to build Phrase-based SMT (PBSMT) systems and evaluated their quality using automatic metrics. For each language, we split the corpus in one training, two developments and two test sets as shown in Table 1 (the last column points to the HTML documents in the website).
We trained a PBSMT model using the Moses toolkit [Koehn et al. 2007] following its “baseline” settings and truecased data [Koehn et al. 2008]. For comparison purposes, we trained a model using the old version of the training set (18,232 sentence pairs): pt-en Fapesp-v1 (UoW-fapesp-v1), and another using the new version: pt-en Fapesp-v2 (UoW-fapesp-v2). In both experiments we used the Fapesp-v2 dev for tuning the features of the PBSMT model. The systems were tested using 3 test sets: i) Fapesp-v1 test (667 sentences), ii) Fapesp-v2 test-a, and iii) Fapesp-v2 test-b. We assessed the system’s performances in terms of BLEU [Papineni et al. 2002] and its case-sensitive version BLEU-c. We also compared the translations against those produced by Google Translate, an off-the-shelf, out-of-domain PB-SMT system. Table 2 shows that the model trained using our automatically compiled corpus significantly outperforms both Google Translate and a model trained on the previous, smaller version of the corpus, UoW-fapesp-v1, in all test sets.

### Table 2. Performance on the test sets

| PBSMT system | UoW-fapesp-v1 | UoW-fapesp-v2 | Google |
|--------------|---------------|---------------|--------|
|              | BLEU | BLEU-c | BLEU | BLEU-c | BLEU | BLEU-c |
| fapesp-v1    | 39.99 | 37.94 | **57.09** | **54.25** | 37.62 | 36.97 |
| fapesp-v2-a  | 43.24 | 40.98 | **57.69** | **54.87** | 40.4 | 39.86 |
| fapesp-v2-b  | 30.5 | 28.8 | **44.69** | **42.15** | 38.35 | 37.71 |

For pt-es, a PBSMT system trained using Fapesp-v2 achieved BLEU scores of 70.42 (Fapesp-v1 test), 70.26 and 71.49 (Fapesp-v2 test-a and test-b).

### 4. Conclusions

We have presented an approach to automatically build parallel corpora involving Brazilian Portuguese. The corpora are aligned at the sentence level and contain over 150K pt-en and pt-es sentence pairs. This is comparable in size to the News Commentaries parallel corpora used for standard MT evaluations by the Workshop on Machine Translation. We showed that the proposed strategies for document and sentence alignment achieve good performance and demonstrated the usefulness of our corpora for building SMT systems.

Both pt-en and pt-es corpora are available, along with the tools for the complete pipeline of corpus creation. We also intend to keep updating the corpora as the magazine releases additional issues. We expect these resources will be useful for people working with multilingual applications that require parallel corpora involving Brazilian Portuguese.

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4. [http://www.statmt.org/wmt11/](http://www.statmt.org/wmt11/)
5. [http://pers-www.wlv.ac.uk/~in1676/resources](http://pers-www.wlv.ac.uk/~in1676/resources)
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