Rapid Training of Information Extraction with Local and Global Data Views

by

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This dissertation focuses on fast system development for Information Extraction (IE). State-of-the-art systems heavily rely on extensively annotated corpora which are slow to build for a new domain or task. Moreover, previous systems are mostly built with local evidence such as words in a short context window or features that are extracted at the sentence level. They usually generalize poorly on new domains. This dissertation presents novel approaches for rapidly training an IE system for a new domain or task based on both local and global evidence. Specifically, we present three systems: a relation type extension system based on active learning a relation type extension system based on semi-supervised learning, and a crossdomain bootstrapping system for domain adaptive named entity extraction. The active learning procedure adopts features extracted at the sentence level as the local view and distributional similarities between relational phrases as the global view. It builds two classifiers based on these two views to find the most informative contention data points to request human labels so as to reduce annotation cost. The semi-supervised system aims to learn a large set of accurate patterns for extracting relations between names from only a few seed patterns. It estimates the confidence of a name pair both locally and globally: locally by looking at the patterns that connect the pair in isolation; globally by incorporating the evidence from the clusters of patterns that connect the pair. The use of pattern clusters can prevent semantic drift and contribute to a natural stopping criterion for semisupervised relation pattern discovery. For adapting a named entity recognition system to a new domain, we propose a cross-domain bootstrapping algorithm, which iteratively learns a model for the new domain with labeled data from the original domain and unlabeled data from the new domain. We first use word clusters as global evidence to generalize features that are extracted from a local context window. We then select self-learned instances as additional training examples using multiple criteria, including some based on global evidence.
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Dedicated to my wife and our son
Acknowledgements

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Abstract

This dissertation focuses on fast system development for Information Extraction (IE). State-of-the-art systems heavily rely on extensively annotated corpora, which are slow to build for a new domain or task. Moreover, previous systems are mostly built with local evidence such as words in a short context window or features that are extracted at the sentence level. They usually generalize poorly on new domains.

This dissertation presents novel approaches for rapidly training an IE system for a new domain or task based on both local and global evidence. Specifically, we present three systems: a relation type extension system based on active learning, a relation type extension system based on semi-supervised learning, and a cross-domain bootstrapping system for domain adaptive named entity extraction.

The active learning procedure adopts features extracted at the sentence level as the local view and distributional similarities between relational phrases as the global view. It builds two classifiers based on these two views to find the most informative contention data points to request human labels so as to reduce annotation cost.

The semi-supervised system aims to learn a large set of accurate patterns for extracting relations between names from only a few seed patterns. It estimates the confidence of a name pair both locally and globally: locally by looking at the patterns that connect the pair in isolation; globally by incorporating the evidence from the clusters of patterns that connect the pair. The use of pattern clusters can prevent semantic drift and contribute to a natural stopping criterion for semi-supervised relation pattern discovery.

For adapting a named entity recognition system to a new domain, we propose a cross-domain bootstrapping algorithm, which iteratively learns a model for the
new domain with labeled data from the original domain and unlabeled data from the new domain. We first use word clusters as global evidence to generalize features that are extracted from a local context window. We then select self-learned instances as additional training examples using multiple criteria, including some based on global evidence.
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Chapter 1

Introduction

Information Extraction (IE) is the task of extracting from texts instances of predefined types of entities and relations involving these entities. This dissertation studies the problem of rapid training of IE systems. We focus on two well-defined IE tasks: Named Entity Recognition (NER) and Relation Extraction (RE). Previous state-of-the-art systems for these two tasks are typically trained with extensively annotated corpora, making them hard to adapt to a new domain or extend to a new task. These systems are categorized as supervised systems.

In contrast to the supervised approach, both semi-supervised and active learning approaches aim to reduce the amount of annotated data so as to speed up the training of IE systems for new domains and new tasks. However, this brings many new challenges. For example, how to prevent the well-known semantic drift problem for the iterative process of semi-supervised learning? Can we find a good and natural stopping criterion? What would be the most effective sample selection method for active learning?

In the next two sections, we describe the two IE tasks, review the most relevant
previous work with emphasis on analyzing their shortcomings when applied to new domains and tasks. And then we briefly introduce our solutions for domain adaptation for NER and type extension for RE. The last section presents the outline of this dissertation.

1.1 Named Entity Recognition

1.1.1 Task

NER, as an evaluation task, was first introduced in the Sixth Message Understanding Conference (MUC-6)[34]. Given a raw English text, the goal is to identify names in it and classify them into one of the three categories: PERSON, ORGANIZATION and LOCATION. For the example sentence in Figure1.1, an NER system needs to extract \textit{Bill Gates} as a PERSON, \textit{Seattle} as a LOCATION, and \textit{Microsoft} as an ORGANIZATION. Other evaluations such as CoNLL[20], followed MUC and extended it to many other languages other than English.

\begin{verbatim}
<PERSON>Bill Gates</PERSON>, born October 28, 1955 in <LOCATION>Seattle</LOCATION>, is the former chief executive officer (CEO) and current chairman of <ORGANIZATION>Microsoft</ORGANIZATION>.
\end{verbatim}

Figure 1.1: NER Example in SGML format.

Evaluation of NER performance is done by comparing system output against answer keys prepared by a human annotator. We count as: \textit{correct} if the NE tags agree in both extent and type; \textit{spurious} if a system tag does not match the tag of the answer key; \textit{missing} if a tag in the answer key has no matching tag in the system output. We then compute \textit{recall}, \textit{precision} and \textit{F1} scores as follows:
\[
\text{recall} = \frac{\text{correct}}{\text{correct} + \text{missing}} 
\]  
\[\text{(1.1)}\]

\[
\text{precision} = \frac{\text{correct}}{\text{correct} + \text{spurious}} 
\]  
\[\text{(1.2)}\]

\[
F1 = \frac{2 \times \text{recall} \times \text{precision}}{\text{recall} + \text{precision}} 
\]  
\[\text{(1.3)}\]

Note that the MUC evaluation gives partial credit to cases where a system NE tag matched the answer key in extent but not in type, while the CoNLL scores give no such partial credit. Following most previous work, we will use the CoNLL evaluation metric instead of the MUC metric.

1.1.2 Prior Work

A comprehensive survey of NER is given by [61]. Here we only review the classical NER models and other most relevant work of this dissertation.

1.1.2.1 Supervised NER

NER is well recognized as a sequence labeling task. Given a sequence of tokens/words of a sentence, \( T = (t_1...t_n) \), the goal is to assign it the most likely sequence of name classes \( c_1...c_n \). As a name may contain multiple tokens, it is necessary and convenient to break a name class into a few subclasses. Table 1.1 shows an example using the BIO scheme, which splits a name type into \( B \) (beginning of the name type) and \( I \) (continuation of the name type). A \( O \) tag is used to represent a non-name token.
Table 1.1: NER Examples with BIO decoding scheme

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<tbody>
<tr>
<td>B-person</td>
<td>I-person</td>
<td>O</td>
<td>O</td>
<td>O</td>
<td>O</td>
<td>O</td>
<td>O</td>
<td>B-location</td>
</tr>
</tbody>
</table>

Supervised NER models are mostly Markov Models, which represent a name class as a hidden state in a Markov chain. So the task transforms to finding the most likely state sequence $s_1...s_n$ given the token sequence $t_1...t_n$.

$$S = \arg \max P(s_1...s_n|t_1...t_n)$$  \hspace{1cm} (1.4)

Applying Bayes’ rule and assuming a first order Markov process, the probability is factored into Markov transition probabilities, where the transition to the current state $s_i$ depends only on the previous state $s_{i-1}$ and the observation at the current state $t_i$.

$$S = \arg \max P(s_1...s_n|t_1...t_n) = \arg \max P(t_1...t_n|s_1...s_n)P(s_1...s_n) = \arg \max \Pi_{i=1}^n P(t_i|s_i)P(s_i|s_{i-1})$$  \hspace{1cm} (1.5)

This is a Hidden Markov Model (HMM). $P(t_i|s_i)$ refers to the emission probability, the probability of the token $t_i$ given the state $s_i$, and $P(s_i|s_{i-1})$ refers to the transition probability, the probability of the state $s_i$ given the previous state $s_{i-1}$.

In Maximum Entropy Markov Model (MEMM)\cite{56}, instead of computing the two conditional probabilities: emission and transition, the transition to the current state is conditioned on both the previous state and the entire token sequence, i.e. $P(s_i|s_{i-1}, t_1...t_n)$. Although in practice, only a short context window is used, for example, $t_{i-2}t_{i-1}t_it_{i+1}t_{i+2}$.
\[ S = \arg \max P(s_1...s_n|t_1...t_n) \]

\[ = \arg \max \prod_{i=1}^{n} P(s_i|s_{i-1}, t_1...t_n) \]

\[ \approx \arg \max \prod_{i=1}^{n} P(s_i|s_{i-1}, t_{i-2}t_{i-1}t_it_{i+1}t_{i+2}) \]  

HMM and MEMM models can be quite effective when they are tested on the same domain of texts as the training domain. However, they usually perform poorly on a domain that is different or slightly different from the training domain. For example, [21] reported that a system trained on the CoNLL 2003 Reuters dataset achieved an F-measure of 0.908 when it was tested on a similar Reuters corpus but only 0.643 on a Wall Street Journal dataset.

Looking closer at the domain effect problem, we observe that the models’ parameters (the transition probabilities of both the HMM and MEMM models and the emission probability of the HMM model) are trained to optimize the performance on domains that are similar to the training domain. A new domain may contain many names and contexts that have not been observed by the models. An unobserved token \( t_i \) has poor parameter estimation, for example, the probability that a state emits that token in an HMM model, \( P(t_i|s) \). The transition from one state to another will be poorly estimated as well, which is true for both HMM and MEMM models. For example, using the BIO scheme, if the training domain contains typical English person names (1 to 3 tokens) and the testing domain contains many transliterated foreign names (more than 3 tokens), then the transition probability from the state I-person to itself would be underestimated for the testing domain, given that this type of transition is more frequent in the testing than in the training domain.

To remedy this, supervised models would have to design sophisticated back-
off strategies[5]. Alternatively, one can annotate texts of the testing domain and re-train the models. This however, is a time consuming process and may not be reusable for other domains.

1.1.2.2 Semi-supervised NER

The portability of supervised NER models is limited by the availability of annotated data of a new domain. Semi-supervised learning aims to build a model for a new domain with only small amounts of labeled data and large amounts of unlabeled data.

Semi-supervised NER is feasible because one can usually identify and classify a name by either the name string itself (e.g., New York City), or by the context in which the name appears (e.g., He arrived in <>). A bootstrapping procedure based on the co-training [7] idea utilizes the name string itself and the context as two views of a data point in NER. Starting with a few seed names of a particular category, it first identifies and evaluates the contexts in which these names occur. It then uses selected predictive and confident contexts to match new names which are further used to learn new contexts.

Bootstrapping systems mostly focus on semantic lexicon acquisition [67][57][86], building a dictionary of a specific semantic class from a few seed examples. The evaluation is typically done by manually judging the correctness of the extracted terms, usually only for the top ranked extractions. This accuracy-based evaluation is quite different from the MUC and CoNLL evaluations. Only a few bootstrapping systems used the MUC and CoNLL evaluation metrics[86]. Customizing these systems for the MUC and CoNLL style within document NER is worth further exploration.
A second problem with semi-supervised NER is *semantic drift*. While a name typically belongs to just one class, there are names that belong to multiple classes when they are separated from contexts (e.g., *Washington can be a person or a location*). So bootstrapping for one class of names may extract names of other classes. To alleviate this problem, seeds of multiple categories are introduced to serve as *negative* categories to provide *competition* among categories[86]. However, these *negative* categories are usually identified by hand, which undermines the intention of semi-supervised learning which is to reduce human supervision.

More recently, [75] and [57] proposed to use unsupervised learning to discover these *negative* classes. They cluster either words or contexts based on *distributional similarity* and use identified clusters as *negative* classes so as to avoid the manual construction of these classes.

A third problem with bootstrapping-based systems is the lack of a natural stopping criterion. Most systems either use a fixed number of iterations or use a labeled development set to detect the right stopping point. We propose to stop bootstrapping by detecting *semantic drift* in Chapter 3. It is straightforward to detect *semantic drift* if the bootstrapping process tends to accept more members of the *negative* classes instead of the *positive* class.

### 1.1.2.3 Active Learning for NER

Active learning reduces annotation cost by selecting the most *informative* examples for requesting human labels. The most *informative* example can be the most *uncertain* example tagged by the underlying NER model. However, selecting uncertain examples may include a lot of outliers which are rare examples and may not necessarily improve NER performance. [71] proposed a multiple-criteria-based
selection method, including informativeness, representativeness, density, and diversity, so as to maximize the performance gain one can achieve in a single round of active learning by providing a good balance between selecting common instances and rare instances.

Committee-based selection has also been applied to active learning for NER[47][4]. [4] builds two Markov models based on different feature sets, uses KL-divergence to quantify the disagreement between the two models, and select the most disagreed examples to request human labels.

1.1.3 Cross-domain Bootstrapping for Named Entity Recognition

Millions of dollars have already been spent on annotating news domain data and state-of-the-art NER systems are typically trained with news data. However, these supervised models perform poorly on non-news domains. Moreover, when building a NER model for a new target domain, both semi-supervised and active learning tend to work on the target domain directly and ignore the potential benefits one can get from existing annotated news data.

We propose a cross-domain bootstrapping (CDB) algorithm to adapt a NER model trained on a news domain to a terrorism report domain without annotating examples on the latter. CDB first builds a MEMM model on the news domain, iteratively tags a large unlabeled terrorism report corpus to select self-learned labeled instances, and finally upgrades the model with these new instances. There are two major components of CDB: feature generalization and instance selection.

Feature generalization. The news-trained model is based on English language names, such as “John Smith” but is much less confident in extracting names from
other languages, such as “Abdul al-Fatah al-Jayusi”, which are more common in US terrorism reports. To use the words from the names as features, our CDB algorithm moves one level up to build word clusters and extracts clusters as features. Specifically, words are clustered in a hierarchical way based on distributional similarity. Words in terrorism reports may share the same cluster membership with those in news articles. So even if the news-trained model does not include a specific word of the terrorism report domain, the cluster level features may still fire for the terrorism report domain.

*Instance selection.* Armed with generalized features, CDB now has a better starting point to select self-learned examples. We have adopted multiple criteria for instance selection, two of which, *density* and *diversity*, again explore the cluster property of unlabeled data. The idea is to select centroid instances instead of selecting outliers. Promoting instances like “President A.P.J. Abdul Kalam says” will bring in more salient contexts than others. As models are upgraded iteratively, CDB can be slow. The *diversity* criterion aims to maximize performance gain by selecting a set of instances that are not very similar to each other during a single iteration. The difference of densities is used to prevent promoting two very similar instances.

With *feature generalization* and *instance selection*, CDB improved the news domain NER system’s performance on the terrorism report domain by 7 points of F-measure (from 66% to 73%). It also significantly outperformed traditional bootstrapping by about 3 points of F-measure. These improvements were largely due to the exploration of the clusters of unlabeled data.
1.2 Relation Extraction

1.2.1 Task

Names and entities are isolated information. Connecting related entities and labeling them with the right semantic relation type is the task of relation extraction, which would provide a more useful resource for IE-based applications. There are two types of relation extraction tasks that have been extensively studied.

**Relation mention extraction.** The US government sponsored Automatic Content Extraction (ACE) program introduced relation extraction in 2002 which was continued until 2008. Table 2.1 shows the relation types and examples defined in ACE 2004. ACE defines a relation over a pair of entities. And a relation mention is defined over a pair of entity mentions in the same sentence. Assuming that the sentence is “Adam, a data analyst for ABC Inc.”, there are two entities {Adam, a data analyst} and {ABC Inc.} in it. Adam and a data analyst are two mentions of the entity {Adam, a data analyst}. Relation mention is defined over the two closest entity mentions in a sentence. So according to the ACE definition, an EMPLOYMENT relation mention should be established between a data analyst and ABC Inc. One needs to rely on coreference information to determine the relation between Adam and ABC Inc.

**Relation extraction between names.** A large body of relation extraction work concerns extracting relations between a pair of names, again in the same sentence. So for the previous example, one needs to extract an EMPLOYMENT relation between the two names Adam and ABC Inc. More recently, the Knowledge Base Population (KBP) evaluation[42] introduces the task of slot filling, finding attributes for PERSON and ORGANIZATION in about 1 million documents.
KBP does not constrain a relation to hold in a single sentence. In fact, for some hard cases, one would have to do cross-sentence relation extraction to find certain attributes.

**Evaluation.** As a relation extraction system relies on the output of an entity extraction model, evaluating the performance of relation extraction is more complicated than that of entity extraction. To make performance comparable among different systems developed by different sites, researchers usually separate the evaluation of entity extraction from relation extraction. For example, instead of relying on system output of entity mentions, most reported ACE systems use the hand annotated entity mentions as the input of a relation extraction system. As for NER, relation extraction are usually evaluated based on *precision*, *recall*, and *f-measure*.

### 1.2.2 Prior Work

#### 1.2.2.1 Supervised relation extraction

A supervised approach casts relation extraction as a classification task. Given a collection of documents that have been annotated with entities and relations, one will build a positive example for a pair of entity mentions if the pair is annotated with a type of relation, and build a negative example if the pair is not labeled with any predefined relation types.

There are two standard learning strategies for relation extraction, *flat* and *hierarchical*. The *flat* strategy simply trains a $n+1$-way classifier for $n$ classes of relations and the non-relation class (no predefined relation holds for a pair of entity mentions). The *hierarchical* strategy separates *relation detection* from
relation classification. It first trains a binary classifier which detects whether a pair of entity mentions has a relation or not. Then a \( n \)-way classifier is trained to distinguish among the \( n \) relation classes. This hierarchical strategy was proposed based on the observation from the ACE corpora that the number of non-relation instances is usually 8 to 10 times larger than that of relation instances. The intention was to group all relation instances into one single class so as to alleviate the effect of unbalanced class distribution between the relation and non-relation classes and improve recall for relation instances.

Both feature-based and kernel-based classification approaches have been applied to relation extraction.

A feature-based classifier starts with multiple level analyses, including tokenization, syntactic, and dependency parsing of the sentence that contains a pair of entity mentions[48][93]. It then extracts a feature vector for the pair which contains various entity, sequence, syntactic, and semantic features. Table 2.2 shows an example feature set.

Pairs of entities that have the same relation type are usually connected by similar token sequences, the shortest paths in dependency trees, or have similar subtree structures in syntactic parsing trees. These structures can be modeled as features in a feature-based system but would be much more powerful if a kernel function can be defined over them. In fact, kernel functions at the token sequence[14], dependency path[13] and syntactic parsing tree[95] levels have all been proposed to extract relations, with tree kernels working as effectively as or even better than a feature-based system.

Both feature-based and kernel-based supervised relation extraction systems can give state-of-the-art performance. However, extending such systems to a new type
of relation would require annotating data for this new type of relation from scratch. This greatly impedes the application of such systems to extract a new type of relation.

1.2.2.2 Semi-supervised relation extraction

Language is redundant. A pair of names may be connected by multiple patterns (e.g., a sequence of tokens) that actually express the same relation. Table 1.2 shows sample patterns that are all indicating that there is an EMPLOYMENT relation between Bill Gates and Microsoft. Similarly, a single pattern may connect many pairs of names. See the examples in Table 1.3.

<table>
<thead>
<tr>
<th>Named Entity 1</th>
<th>Pattern</th>
<th>Named Entity 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bill Gates</td>
<td>. the head of</td>
<td>Microsoft</td>
</tr>
<tr>
<td></td>
<td>. the chairman of</td>
<td></td>
</tr>
<tr>
<td></td>
<td>. chief executive of</td>
<td></td>
</tr>
<tr>
<td></td>
<td>. chief executive officer of</td>
<td></td>
</tr>
<tr>
<td></td>
<td>. chairman of software giant</td>
<td></td>
</tr>
<tr>
<td></td>
<td>. Chairman and CEO of</td>
<td></td>
</tr>
<tr>
<td></td>
<td>. founder and chairman of</td>
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</table>

Table 1.2: Sample patterns that connect Bill Gates and Microsoft

Using the pair of names as one view and the patterns connecting the pair as another view, semi-supervised relation extraction usually adopts a co-training style bootstrapping procedure[2]. It starts with a few seed patterns that indicate the target relation to match name pairs and evaluate the confidence of these extracted name pairs. Then in the next step it uses these name pairs to search for additional patterns that are connecting these pairs. These newly discovered patterns are
Table 1.3: Sample name pairs that are connected by “, the chairman of”

<table>
<thead>
<tr>
<th>Named Entity 1</th>
<th>Pattern</th>
<th>Named Entity 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Steve Jobs</td>
<td></td>
<td>Apple</td>
</tr>
<tr>
<td>Eric Schmidt</td>
<td></td>
<td>Google</td>
</tr>
<tr>
<td>Roy J. Bostock</td>
<td>, the chairman of</td>
<td>Yahoo</td>
</tr>
<tr>
<td>Robin Li</td>
<td></td>
<td>Baidu</td>
</tr>
<tr>
<td>... ……</td>
<td></td>
<td>... ...</td>
</tr>
</tbody>
</table>

evaluated and the most confident ones are added to the seed pattern set to repeat the bootstrapping process.

Bootstrapping only requires a few seed examples to be annotated and hence can be rapidly customized for a new relation type. However, there are at least three limitations of the bootstrapping based relation extraction.

First, it is limited to extracting relations between names. As observed in the ACE corpus, relations are more frequently expressed by pronouns and nouns than by names. If the goal is to extract relation mentions as defined in the ACE evaluation, it is usually not feasible to separate the context from the pair of entity mentions to establish the two views of a relation instance. For one thing, there are many relation mentions that are just single noun phrases and there are no tokens between the two involved entity mentions. For example, a SOCIAL-FAMILY relation exists between the mention *His* and the mention *His father* as in the sentence “His father said that ...”. The relation is expressed mostly by the pair of entity mentions. The right context “said that ...” is not sufficient to indicate that there is a SOCIAL-FAMILY between the pair of mentions. For another, a relation mention that is not expressed between two names tends to be quite ambiguous. For example, a pair of pronoun and name does not determine a unique relation. Re-
placing the wild cards in “He * New York” with the pattern “lives in” will produce a *residence* relation while with “drove to the city of” will give a *locatedIn* relation.

Secondly, even for extracting relations between names, bootstrapping also faces the *semantic drift* problem. Many pairs of names may have multiple relations. For example, as shown in Table 1.4, depending on the contexts, the pair $<Bill Clinton, Arkansas>$ may be interpreted as a *birthPlace*, *governorOf*, or *locatedIn* relation. A bootstrapper for the *birthPlace* relation may accept patterns that actually indicate the other two relations, resulting in pairs of names extracted that are straying away from the semantics of the *target* relation. As for bootstrapping based NER, multi-category bootstrapping was proposed to alleviate the *semantic drift* problem[84]. However, as mentioned earlier, it is not appealing and difficult to find useful *negative* categories by hand. We propose to use pattern clusters to uncover the possible types of relations of a pair of names, leading to a principled way to introduce *negative* categories[75]. We will show an application of acquiring relation patterns in Chapter 3.

<table>
<thead>
<tr>
<th>Named Entity 1</th>
<th>Pattern</th>
<th>Named Entity 2</th>
<th>Relation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bill Clinton</td>
<td>was born in</td>
<td>Arkansas</td>
<td>birthPlace</td>
</tr>
<tr>
<td></td>
<td>was governor of</td>
<td></td>
<td>governorOf</td>
</tr>
<tr>
<td></td>
<td>visited</td>
<td></td>
<td>locatedIn</td>
</tr>
<tr>
<td>... ...</td>
<td>... ...</td>
<td></td>
<td>...</td>
</tr>
</tbody>
</table>

Table 1.4: Example of a pair of names that have multiple relation.

Moreover, similar to bootstrapped NER systems, current bootstrapped relation extraction systems lack a natural stopping criterion. As mentioned earlier, we will try to solve both *semantic drift* and *finding a natural stopping criterion* problems
at the same time.

1.2.2.3 Un-supervised relation extraction

Both supervised and semi-supervised approaches extract instances of relations of predefined types. Relations can be also discovered in an un-supervised way. The main idea is to cluster pairs of names based on the similarity of context words that are intervening between the names. Figure 1.2 shows one example from [37]. Like semi-supervised learning, it also utilizes the linguistic intuition that name pairs that have the same relation usually share similar contexts.

![Diagram showing un-supervised relation discovery]

Figure 1.2: Example of un-supervised relation discovery.

The relation discovery procedure begins with named entity tagging. Then the contexts of a pair of names are gathered together to generate a feature vector. The similarity between pairs of names is calculated using the cosine measure, which is used as the distance measure of Complete Linkage, a type of Hierarchical Agglomerative Clustering, to cluster pairs of names.

Un-supervised relation discovery can discover meaningful relations with zero annotation cost. However, it faces a few challenges to be more beneficial to poten-
tial applications. First of all, as mentioned earlier in this chapter, a pair of names may actually exhibit multiple relation types. This might affect the consistency of the labels of some of the generated relation clusters. Secondly, further improvements are needed for better alignment between generated clusters and a specific application. For example, one application might need the governor relation while another might need all relations that are related to government officials. Generating clusters at the right level of granularity with respect to the specific underlying application need further exploration.

1.2.3 Active and Semi-supervised Learning for Relation Type Extension

This dissertation studies how to rapidly and accurately extend a system for a new type of relation. To reduce annotation cost, we apply two learning algorithms for fast training of relation extraction systems: active learning and semi-supervised learning. In both approaches, we show that one can benefit from using both the local view and the global view of a relation instance to reduce annotation cost without degrading the performance of relation extraction. We briefly introduce our approaches below and will present their details in Chapter 2 and 3.

1.2.3.1 Active learning for relation type extension

We apply active learning to extract relation mentions, as defined in the ACE program. The local view involves the features extracted at the sentence level (from a sentence that contains a pair of entity mentions). As it is not reasonable to separate the pair of entity mentions and their context to establish two data views, we represent each relation instance as a relational phrase. Roughly speaking, we
define a relational phrase as the following: if there are no context tokens between
the two mentions, we treat the two mentions together as a relational phrase. If
there are tokens in between, we then use the middle token sequence as the rela-
tional phrase. We will give a formal definition of relational phrase in Chapter 2.
The global view involves the distributional similarity between relational phrases
computed from a 2-billion-token text corpus.

We build a feature-based relation classifier based on the local view. We also
build a k-nearest neighbor classifier based on the global view, classifying an unla-
beled instance based on its closest labeled examples. We then measure the degree
of disagreement between the two classifiers using KL-divergence. The instances
with the largest degree of deviation between the two classifiers are treated as the
most informative examples to request human labels.

1.2.3.2 Semi-supervised learning for relation type extension

We apply semi-supervised learning to extract relations between names. Specif-
ically, we develop a bootstrapping procedure for acquiring semantic patterns for
extracting a target relation.

The procedure begins with a small set of patterns of the target relation to match
name pairs. These name pairs should be evaluated based on their confidence of
indicating the target relation. Traditional bootstrapping evaluates the confidence
of a newly matched name pair by looking at the confidence of each individual
pattern that connects the name pair in isolation. We call this type of confidence
the local view confidence. This dissertation moves one level up to build pattern
clusters and estimates the confidence of a name pair based on the clusters of
patterns that connect the name pair as well. We call it the global view confidence.
Figure 1.3 and Figure 1.4 depict the *local* and *global* views.

<table>
<thead>
<tr>
<th>Named Entity 1</th>
<th>Pattern</th>
<th>Named Entity 2</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bill Clinton</td>
<td><em>visit</em></td>
<td>Arkansas</td>
</tr>
<tr>
<td></td>
<td><em>born in</em></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>fly to</em></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>governor of</em></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>arrive in</em></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>campaign in</em></td>
<td></td>
</tr>
</tbody>
</table>

Figure 1.3: Example of the local view of a name pair.

Figure 1.4: Example of the global view of a name pair. $C_i$ means the pattern cluster $i$.

The procedure then uses the matched name pairs to search for additional patterns and evaluate these patterns. Top confident patterns are added to the seed pattern set. And the whole process repeats until it meets a stopping criterion.

Introducing pattern clusters not only helps to more reliably estimate the confidence of matched name pairs so as to prevent *semantic drift*, but also contributes to a natural stopping criterion of the bootstrapping process. If the process tends to promote patterns that do not share the same cluster membership with the seed
patterns, then it is likely that patterns being accepted are actually indicating other
types of relations other than the target relation. Hence semantic drift occurs and
the process should be stopped.

1.3 Outline of Thesis

The rest of the dissertation is organized as follows: Chapter 2 presents in
detail the proposed active learning based relation type extension with local and
global data views. Chapter 3 focuses on the bootstrapping based relation type
extension with local and global data views[75]. We then present the cross-domain
bootstrapping algorithm for domain adaptive NER in Chapter 4[77]. We conclude
and point to future work in Chapter 5.
Chapter 2

Relation Type Extension: Active Learning with Local and Global Data Views

2.1 Introduction

Relation extraction aims to connect related entities and label them with the right semantic relationship. For example, a relation extraction system needs to detect an Employment relation between the entities He and Arkansas in the sentence He was the governor of Arkansas. The task of relation type extension is to extend an existing relation extraction system to be able to extract a new type of relation, often called the target relation, preferably in a fast, cheap and accurate way.

A supervised approach can tackle this problem in an accurate way. But it is slow and expensive as it relies on human annotation of a large quantity of
examples. Semi-supervised learning, in contrast, does not require much human effort by automatically bootstrapping a system for the target relation from only a few labeled examples and a large unlabeled corpus. However, a large gap still exists between the performance of semi-supervised and supervised systems. Moreover, their performance largely depends on the choice of seeds[81][50].

Another attractive alternative is active learning, which reduces annotation cost by requesting labels of only the most informative examples while maintaining high learning performance. It is also shown to be robust to the choice of the seeds[47]. Specifically, we focus on relation type extension with co-testing[60], an active learning approach in the co-training[7] setting. It minimizes human annotation effort by building two classifiers based on two different views of the data and asking for human labels only for contention data points, points on which the two classifiers disagree about their labels. The key to the success of co-testing is to find a natural way of splitting a data point into two views that are uncorrelated and compatible (each view is sufficient in labeling the data point).

To date, there is limited work on applying co-testing to relation type extension. The main difficulty, as we believe, is the lack of a natural way of partitioning the data into two uncorrelated and compatible views. Unlike named entity classification where one can rely on either the name string itself (Arkansas) or the context (was the governor of <>) to determine the type of the named entity[23][47], the type of a relation is mostly determined by the context in which the two entities appear. For example, it is not possible to decide the type of relation between He and Arkansas without the local context was the governor of. If the context was traveled to, then the relation of the two entities would change entirely. Thus it is not desirable to separate the entities from their context to establish two views. Instead, we treat
them together as a single view, the \textit{local view}.

Motivated by the idea of distributional similarity, we move beyond the \textit{local view} to interpret the relation between two entities. Specifically, we compute from a 2 billion token corpus the distributional similarities between relational phrases. We take these similarities as the \textit{global view} upon which we build a classifier which classifies new examples based on the $k$--\textit{nearestneighbor} algorithm. For example, if the phrase \textit{arrived in} is more similar to \textit{traveled to} than to \textit{was the governor of}, the \textit{global view} classifier classifies entities connected by \textit{arrived in} as the same relation as those connected by \textit{traveled to}. Armed with this \textit{global view} classifier and a classifier trained with features extracted from the \textit{local view}, applying co-testing to relation type extension becomes feasible.

The main contributions of active learning with \textit{local} and \textit{global} views are: it indeed introduces two \textit{uncorrelated} and \textit{compatible} views for relation extraction. It provides substantial reduction of annotation effort as compared to various baselines based on an extensive experimental study on the ACE 2004 corpus. Furthermore, it leads to faster convergence of learning.

The next section introduces our task. Section 2.3 and 2.4 describes the \textit{local} and \textit{global} view classifiers in detail. We present LGCo-Testing and baseline systems in Section 2.5 and 2.6, and evaluate them in Section 2.7. We discuss related work in Section 2.8 and conclude this chapter in Section 2.9.


2.2 Task Definition

We choose to work on the well defined relation extraction task of the ACE\textsuperscript{1} program in 2004, mostly driven by the purpose of system comparison as many published results on this data set are available. A relation is defined over a pair of entities within a single sentence. ACE 2004 defined 7 major relation types. Some examples from the annotation guideline\textsuperscript{2} are shown in Table 2.1. Following previous work, we only deal with relation mentions.

<table>
<thead>
<tr>
<th>Type</th>
<th>Example</th>
</tr>
</thead>
<tbody>
<tr>
<td>EMP-ORG</td>
<td>the CEO of Microsoft</td>
</tr>
<tr>
<td>PHYS</td>
<td>a military base in Germany</td>
</tr>
<tr>
<td>GPE-AFF</td>
<td>U.S. businessman</td>
</tr>
<tr>
<td>PER-SOC</td>
<td>his ailing father</td>
</tr>
<tr>
<td>DISC</td>
<td>each of whom</td>
</tr>
<tr>
<td>ART</td>
<td>US helicopters</td>
</tr>
<tr>
<td>OTHER-AFF</td>
<td>Cuban-American people</td>
</tr>
</tbody>
</table>

Table 2.1: ACE relation examples from the annotation guideline. Heads of entity mentions are marked.

We consider two experimental settings to simulate real world settings when we build a system for a target relation.

1. **Binary setting** where we treat one of the ACE relation types as the target relation. We use as labeled data a few labeled examples of the target relation (possibly by random selection). And all other examples in the ACE corpus are treated as unlabeled data.

2. **Multi-class setting** where we treat one of the ACE relations as the target rela-

\textsuperscript{1}Task definition: http://www.itl.nist.gov/iad/894.01/tests/ace/ ACE guidelines: http://projects.ldc.upenn.edu/ace/

\textsuperscript{2}http://projects.ldc.upenn.edu/ace/docs/EnglishRDCV4-3-2.PDF
tion and all others as *auxiliary* relations. We use as labeled data a few labeled examples of the *target* relation and all labeled *auxiliary* relation examples. All other examples in the ACE corpus are treated as unlabeled data. This multi-class setting simulates a common training scenario where one wants to extend a system trained with the extensive annotation of the ACE relation types to additional types of relations.

### 2.3 The Local View Classifier

There are two common learning approaches for building a classifier based on the local view: feature based[48][93][45] and kernel based[88][13][14][92][95]. As we want to compare LGCo-Testing with co-testing based on a feature split at the local level, we choose to build a feature based *local* classifier.

Given a relation instance $x = (s, e_i, e_j)$, where $e_i$ and $e_j$ are a pair of entities and $s$ is the sentence containing the pair, the local classifier starts with multiple level analyses of the sentence such as tokenization, syntactic parsing, and dependency parsing. It then extracts a feature vector $v$ which contains a variety of lexical, syntactic and semantic features for each relation instance. Our features are cherrypicked from previous feature based systems. Table 2.2 shows the feature set with examples.

After feature engineering, the local classifier applies machine learning algorithms to learn a function which can estimate the conditional probability $p(c | v)$, the probability of the type $c$ given the feature vector $v$ of the instance $x$. We used maximum entropy (MaxEnt) to build a binary classifier (for the binary setting) and a multi-class classifier (for the multi-class setting) because the training is fast,
<table>
<thead>
<tr>
<th>Level</th>
<th>Type</th>
<th>Description</th>
<th>Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Entity</td>
<td>ET</td>
<td>Entity types</td>
<td>ET1=PERSON; ET2=LOCATION</td>
</tr>
<tr>
<td></td>
<td>ET12</td>
<td>Combination of ET1 and ET2</td>
<td>ET12=PERSON–LOCATION</td>
</tr>
<tr>
<td></td>
<td>ML</td>
<td>Mention levels</td>
<td>ML1=NAME; ML2= NOMINAL</td>
</tr>
<tr>
<td></td>
<td>ML12</td>
<td>Combination of ML1 and ML2</td>
<td>ML12=NAME–NOMINAL</td>
</tr>
<tr>
<td></td>
<td>HE</td>
<td>Heads of entities</td>
<td>HE1=Clinton; HE2= border</td>
</tr>
<tr>
<td></td>
<td>HE12</td>
<td>Combination of HE1 and HE2</td>
<td>Clinton–border</td>
</tr>
<tr>
<td></td>
<td>BagWE</td>
<td>Bag of words of entities</td>
<td>{President, Clinton} {the, Irish, border}</td>
</tr>
<tr>
<td>Sequence</td>
<td>WBE1</td>
<td>Words before entity 1</td>
<td>[NIL]</td>
</tr>
<tr>
<td></td>
<td>WB</td>
<td>Words between</td>
<td>[travel, to]</td>
</tr>
<tr>
<td></td>
<td>WAE2</td>
<td>Words after entity 2</td>
<td>[for, an]</td>
</tr>
<tr>
<td></td>
<td>NUMWB</td>
<td># words between</td>
<td>2</td>
</tr>
<tr>
<td></td>
<td>TPatternET</td>
<td>Token pattern coupled with entity types</td>
<td>PERSON_traveled_to_LOCATION</td>
</tr>
<tr>
<td>Syntactic Parsing</td>
<td>PTP</td>
<td>Path of phrase labels connecting E1 and E2 in the parsing tree</td>
<td>NF–VF–PP</td>
</tr>
<tr>
<td></td>
<td>PTPH</td>
<td>PTP augmented with the head word of the top phrase in the path</td>
<td>NF–S–traveled–VF–PP</td>
</tr>
<tr>
<td></td>
<td>CPHRE1</td>
<td>Chunk heads before E1</td>
<td>[NIL]</td>
</tr>
<tr>
<td></td>
<td>CPHB</td>
<td>Chunk heads between</td>
<td>[travel, to]</td>
</tr>
<tr>
<td></td>
<td>CPHAE2</td>
<td>Chunk heads after E2</td>
<td>[for, ceremony]</td>
</tr>
<tr>
<td></td>
<td>CPP</td>
<td>Path of phrase labels connecting the two entities in the chunking</td>
<td>NF–VP/S–PF–NP</td>
</tr>
<tr>
<td></td>
<td>CPPH</td>
<td>CPP augmented with head words</td>
<td>NF–Clinton–VF/S–traveled–PF–te–NF–border</td>
</tr>
<tr>
<td></td>
<td>CPatternET</td>
<td>Chunk head pattern with entity types</td>
<td>PERSON_traveled_to_LOCATION</td>
</tr>
<tr>
<td>Dependency Parsing</td>
<td>DPathET</td>
<td>Shortest dependency path connecting the two entities coupled with entity types</td>
<td>PREM_nsubj_traveled_prep_to_LOC</td>
</tr>
<tr>
<td></td>
<td>ET1DW1</td>
<td>Entity type and the dependent word for E1</td>
<td>PERSON–traveled</td>
</tr>
<tr>
<td></td>
<td>ET2DW2</td>
<td>Entity type and the dependent word for E2</td>
<td>LOCATION–to</td>
</tr>
<tr>
<td></td>
<td>H1DW1</td>
<td>The head and the dependent word for E2</td>
<td>Clinton–traveled</td>
</tr>
<tr>
<td></td>
<td>H2DW2</td>
<td>Head word and the dependent word for E2</td>
<td>border–to</td>
</tr>
<tr>
<td></td>
<td>ET12Same</td>
<td>Whether E1 and E2 included in the same NP/PP/VP</td>
<td>false/false/true</td>
</tr>
<tr>
<td></td>
<td>NP/PP/VP</td>
<td>Whether E1 and E2 included in the same NP/PP/VP</td>
<td></td>
</tr>
</tbody>
</table>

Table 2.2: Sample local features for “<e_i>President Clinton</e_i> traveled to <e_j>the Irish border</e_j> for an ... ”
which is crucial for active learning as it is not desirable to keep the annotator waiting because of slow training.

### 2.4 The Global View Classifier

Building a classifier based on the *global* view involves three stages of process: extracting relational phrases, computing distributional similarities, and building the classifier based on similarities. We describe these stages in detail below.

**Extracting relational phrases.** Given a relation instance \( x = (s, e_i, e_j) \) and assuming that \( e_i \) appears before \( e_j \), we represent it as a relational phrase \( p_x \), which is defined as the n-gram that spans the head\(^3\) of \( e_i \) and that of \( e_j \). Formally, \( p_x = [\text{head}_{e_i}, \text{head}_{e_j}] \). For example, we extract *Clinton traveled to the Irish border* as the phrase for the example in Table 2.2. As our goal is to collect the tokens before and after a phrase as features to capture the similarity between phrases and long phrases are too sparse to be useful, we instead use the definition \( p_x = (e_i, e_j) \) (tokens between \( e_i \) and \( e_j \)) when the phrase contains more than 5 tokens. Thus for the example in Table 2.2, because the previously extracted phrase contains 6 tokens, we will instead use the phrase “traveled to” to represent that instance.

**Computing distributional similarities.** We first compile our 2 billion token text corpus to a database of 7-grams\(^70\) and then form 7-gram queries to extract features for a phrase. Example queries for the phrase “traveled to” are shown in Table 2.3.

We then collect the tokens that could instantiate the wild cards in the queries as features. Note that tokens are coupled with their positions. For example, if the

\(^3\)The last token of a noun group.
matched 7-gram is President Clinton traveled to the Irish border, we will extract from it the following five features: President\textunderscore -2, Clinton\textunderscore -1, the\textunderscore +1, Irish\textunderscore +2 and border\textunderscore +3.

Each phrase $P$ is represented as a feature vector of contextual tokens. To weight the importance of each feature $f$, we first collect its counts, and then compute an analogue of $tf-idf$: $tf$ as the number of corpus instances of $P$ having feature $f$ divided by the number of instances of $P$; $idf$ as the total number of phrases in the corpus divided by the number of phrases with at least one instance with feature $f$. Now the token feature vector is transformed into a $tf-idf$ feature vector. We compute the similarity between two vectors using Cosine similarity. The most similar phrases of traveled to and his family are shown in Table 2.4.

<table>
<thead>
<tr>
<th></th>
<th>traveled to</th>
<th>his family</th>
</tr>
</thead>
<tbody>
<tr>
<td>Phrase</td>
<td>Sim.</td>
<td>Phrase</td>
</tr>
<tr>
<td>visited</td>
<td>0.779</td>
<td>his staff</td>
</tr>
<tr>
<td>arrived in</td>
<td>0.763</td>
<td>his brother</td>
</tr>
<tr>
<td>worked in</td>
<td>0.751</td>
<td>his friends</td>
</tr>
<tr>
<td>lived in</td>
<td>0.719</td>
<td>his children</td>
</tr>
<tr>
<td>served in</td>
<td>0.686</td>
<td>their families</td>
</tr>
<tr>
<td>consulted with</td>
<td>0.672</td>
<td>his teammates</td>
</tr>
<tr>
<td>played for</td>
<td>0.670</td>
<td>his wife</td>
</tr>
</tbody>
</table>

Table 2.4: Sample of similar relational phrases.

**Building the classifier.** We build the global view classifier based on the $k$-
nearest neighbor idea, classifying an unlabeled example based on closest labeled examples. The similarity between an unlabeled instance \( u \) and a labeled instance \( l \) is measured by the similarity between their phrases, \( p_u \) and \( p_l \). Note that we also incorporate the entity type constraints into the similarity computation. The similarity is defined to be zero if the entity types of \( u \) and \( l \) do not match. The similarity between \( u \) and a relation type \( c \), \( \text{sim}(u, c) \), is estimated by the similarity between \( u \) and its \( k \) closest instances in the labeled instance set of \( c \) (we take the averaged similarity if \( k > 1 \); we will report results with \( k = 3 \) as it works slightly better than 1, 2, 4 and 5). Let \( h(u) \) be the classification function, we define it as follows:

\[
h(u) = \arg \max_c \text{sim}(u, c) \tag{2.1}
\]

2.5 LGCo-Testing

We first introduce a general co-testing procedure, then describe the details of the proposed LGCo-Testing.

Let \( D_U \) denote unlabeled data, and \( D_L \) denote labeled data, the co-testing procedure repeats the following steps until it converges:

1. Train two classifiers \( h_1 \) and \( h_2 \) based on two data views with \( D_L \)
2. Label \( D_U \) with \( h_1 \) and \( h_2 \) and build a contention set \( S \)
3. Select \( \bar{S} \subseteq S \) based on informativeness and request human labels
4. Update: \( D_L = D_L \cup \bar{S} \) and \( D_U = D_U \setminus \bar{S} \)

Initialization. Initialization first concerns the choice of the seeds. For the multi-class setting, it also needs to effectively introduce the instances of auxiliary
relations.

For the choice of the seeds, as we are doing simulated experiments on the ACE corpus, we take a random selection strategy and hope multiple runs of our experiments can approximate what will actually happen in real world active learning. Moreover, it was empirically found that active learning is able to rectify itself from bad seeds[47]. In all experiments for both the binary and the multi-class settings, we use as seeds 5 randomly selected target relation instances and 5 randomly selected non-relation instances (entity pairs in a sentence not connected by an ACE relation).

For the multi-class setting, we use a stratified strategy to introduce the auxiliary relation instances: the number of selected instances of a type is proportional to that of the total number of instances in the labeled data. We also make the assumption that our target relation is as important as the most frequent auxiliary relation and select these two types equally. For example, assuming that we only have two auxiliary types with 100 and 20 labeled instances respectively, we will randomly select 5 instances for the first type and 1 instance for the second type, given that we initialized our active learning with 5 target relation seeds. We also experimented with several other ways in introducing the auxiliary relation instances and none of them were as effective as the stratified strategy. For one example, using all the auxiliary instances to train the initial classifiers unfortunately generates an extremely unbalanced class distribution and tends to be biased towards the auxiliary relations. For another, selecting the same number of instances for the target relation type and all the auxiliary types does not take full advantage of the class distribution of the auxiliary types, which can be estimated with the labeled data pool.
**Informativeness measurement.** It is straightforward to get the hard labels of an instance from both the local and global view classifiers. As the local classifier uses MaxEnt which is essentially logistic regression, we take the class with the highest probability as the hard label. The hard label of the global classifier is the relation type to which the instance is most similar. As long as the two classifiers disagree about an instance’s hard label and one of the labels is our target relation, we add it to the contention set.

Quantifying the disagreement between the two classifiers is not as straightforward as getting the hard labels because the local classifier produces a probability distribution over the relation types while the global classifier produces a similarity distribution. So we first use the following formula to transform similarities to probabilities.

\[
p(c|u) = \frac{\exp(sim(u, c))}{\sum_i \exp(sim(u, c_i))} \tag{2.2}
\]

Here \(u\) is an instance that needs to be labeled, \(c\) is a specific relation type, and \(sim(u, c_i)\) is the similarity between \(u\) and one of the relation types \(c_i\).

We then use **KL-divergence** to quantify the degree of deviation between the two probability distributions. **KL-divergence** measures the divergence between two probability distributions \(p\) and \(q\) over the same event space \(\chi\):

\[
D(p||q) = \sum_{x \in \chi} p(x) \log \frac{p(x)}{q(x)} \tag{2.3}
\]

It is non-negative. It is zero for identical distributions and achieves its maximum value when distributions are peaked and prefer different labels. We rank the contention instances by descending order of **KL-divergence** and pick the top 5
instances to request human labels during a single iteration.

It is worth mentioning that, for each iteration in the multi-class setting, auxiliary instances are introduced using the stratified strategy as in the initialization step.

**Convergence detection.** We stop LGCo-Testing when we could not find contention instances.

### 2.6 Baselines

We compare our approach to a variety of baselines, including six active learning baselines, one supervised system and one semi-supervised system. We present the details of active learning baselines below, and refer the reader to the experiment section to learn more about other baselines.

**SPCo-Testing.** One of the many competitive active learning approaches is to build two classifiers based on a feature split at the local level. As reported by[45], either the sequence features or the parsing features are generally sufficient to achieve state-of-the-art performance for relation extraction. So we build one classifier based on the *sequence* view and the other based on the *parsing* view. More precisely, one classifier is built with the feature set based on \{entity, sequence\} and the other based on \{entity, syntactic parsing, dependency parsing\}. We build these two classifiers with MaxEnt. The initialization is the same as in LGCo-Testing. *KL-divergence* is used to quantify the disagreement between the two probability distributions returned by the two MaxEnt classifiers. Contention points are ranked in descending order of *KL-divergence* and the top 5 ones are used to query the annotator in one iteration. Like LGCo-Testing, SPCo-Testing stops when the
contention set is empty.

**Uncertainty AL.** This is an uncertainty-based active learning baseline. We build a single MaxEnt classifier based on the full feature set in Table 2.2 at the local level. It uses the same initialization as in LGCo-Testing. Informativeness measurement is based on *uncertainty*, which is approximated by the entropy $h(p)$ of the probability distribution of the MaxEnt classifier over all the relation types $c_i$.

$$h(p) = - \sum_i p(c_i) \log p(c_i) \quad (2.4)$$

It is also non-negative. It is zero when one relation type is predicted with a probability of 1. It attains its maximum value when the distribution is a uniform one. So the higher the entropy, the more uncertain the classifier is. So we rank instances in descending order of entropy and pick the top 5 ones to request human labels. Stopping Uncertainty AL cannot be naturally done as with co-testing. A less appealing solution is to set a threshold based on the *uncertainty* measure.

**Random AL.** This is a random selection based active leaning baseline. 5 instances are selected randomly during a single iteration. There is no obvious way to stop Random AL although one can use a fixed number of instances as a threshold, a number that might be related to the budget of a project.

The next three baselines aim to investigate the benefits of incorporating features from the *global* view into the local classifier. They are inspired by recent advances in using cluster level features to compensate for the sparseness of lexical features[59][76]. Specifically, we use the distributional similarity as a distance measure to build a phrase hierarchy using Complete Linkage. The threshold for cutting the hierarchy into clusters is determined by its ability to place the initial
seeds into a single cluster. We will revisit how the threshold is selected in Chapter 3. For extracting cluster level features, take traveled to as an example, if its cluster membership is $c$, we will extract a cluster feature $\text{phraseCluster} = c$.

**UncertaintyAL+**. The only difference between UncertaintyAL+ and UncertaintyAL is its incorporation of cluster features in building its classifier. This is essentially the active learning approach presented by [59].

**SPCo-Testing+**. It differs from SPCo-Testing only in its sequence view classifier, which is trained with additional phrase cluster features.

**LGCo-Testing+**. It differs from LGCo-Testing only in its local view classifier, which is trained with additional phrase cluster features.

### 2.7 Experiments

#### 2.7.1 Experimental Setting

We experiment with the nwire and bnews genres of the ACE 2004 data set, which are benchmark evaluation data for relation extraction. There are 4374 relation instances and about 45K non-relation instances. Documents are preprocessed with the Stanford parser\(^4\) and chunklink\(^5\) to facilitate feature extraction. Note that following most previous work, we use the hand labeled entities in all our experiments.

We do 5-fold cross-validation as most previous supervised systems do. Each round of cross-validation is repeated 10 times with randomly selected seeds. So, a total of 50 runs are performed (5 subsets times 10 experiments). We report

---

\(^4\)http://nlp.stanford.edu/software/lex-parser.shtml

\(^5\)http://ilk.uvt.nl/team/sabine/chunklink/README.html
average results of these 50 runs. Note that we do not experiment with the DISC (discourse) relation type which is syntactically different from other relations and was removed from ACE after 2004.

The size of unlabeled data is approximately 36K instances ($45K \div 5 \times 4$). Each iteration selects 5 instances to request human labels and 200 iterations are performed. So a total of 1,000 instances are presented to our annotator. This setting simulates satisfying a customer’s demand for an adaptive relation extractor in a few hours. Assuming two entities and their contexts (the annotation unit) are highlighted and an annotator only needs to mark it as a target relation or not, 4 instances per minute should be a reasonable or underestimated annotation speed. And assuming that our annotator takes a 10-minute break in every hour, he or she can annotate 200 instances per hour. We are now ready to test the feasibility and quality of relation type extension in a few hours.

2.7.2 Results

We evaluate active learning on the target relation. Penalty will be given to cases where we predict target as auxiliary or non-relation, and vice versa. To measure the reduction of annotation cost, we compare active learning with the results of [76], which is a state-of-the-art feature-based supervised system. We use the number of labeled instances to approximate the cost of active learning. So we list in Table 2.5 the F1 difference between an active learning system with different number of labeled instances and the supervised system trained on the entire corpus.

The results of LGCo-Testing are simply based on the local classifier’s predictions on the test set. For the SPCo-Testing system, a third classifier is trained with
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Table 2.5: F1 difference (in percentage) = F1 of active learning minus F1 of supervised learning. Bold numbers indicate the best performance. UN := UncertaintyAL. SP := SPCo-Testing. LG := LGCo-Testing.
the full feature set to get test results. As there is a large gap between RandomAL and the supervised system (40% F1 difference regardless of the number of labeled instances), it is excluded from Table 2.5. The three baselines with cluster level features perform similarly as their corresponding baselines without cluster phrases, e.g. UncertaintyAL+ and UncertaintyAL perform similarly. So we exclude their results too.

2.7.3 Analyses

Comparing active learning with supervised learning: LGCo-Testing trained with 1,000 labeled examples achieves results comparable to supervised learning trained with more than 35K labeled instances in both the binary and the multi-class settings. This is true even for the two most frequent relations in ACE 2004, EMP-ORG and PHYS (about 1.6K instances for EMP-ORG and 1.2K for PHYS). This represents a substantial reduction in instances annotated of 97%. So assuming our annotation speed is 200 instances per hour, we can build in five hours a competitive system for EMP-ORG and a slightly weak system for PHYS. Moreover, we can build comparable systems for the other four relations in less than 5 hours. Much of the contribution, as depicted in Figure 2.1, can be attributed to the sharp increase in precision during early stages and the steady improvement of recall in later stages of learning.

Comparing active learning systems: the clear trend is that LGCo-Testing outperforms UncertaintyAL and SPCo-Testing by a large margin in most cases for both experimental settings. This indicates its superiority in selective sampling for fast system development and adaptation. SPCo-Testing, which is based on the feature split at the local level, does not consistently beat the uncertainty based
systems. Part of the reason, as we believe, is that the sequence and parsing views are highly correlated. For example, the token sequence feature “traveled to” and the dependency path feature “nsubj’ traveled prep to” are hardly conditionally independent.

Comparing LGCo-Testing in the multi-class setting with that in the binary setting, we observe that the reduction of annotation cost by incorporating auxiliary types is more pronounced in early learning stages (#labels < 200) than in later ones, which is true for most relations. Figure 2.2 depicts this by plotting the F1 difference (between active learning and supervised learning) of LGCo-Testing in the two experimental settings against the number of labels. Besides the two relations GPE-AFF and OTHER-AFF shown in Figure 2.2, taking ART as a third example relation type, with 50 labels, the F1 difference of the multi-class LGCo-Testing is -29.8 while the binary one is -48.4, which represents a F1 improvement of 19.6 when using auxiliary types. As the number of labels increases, the multi-class setting incorporates more and more auxiliary instances, which might decrease the priors
Figure 2.2: F1 difference of LGCo-Testing for GPE-AFF and OTHER-AFF for the target relations. Hence the improvement for the target relation degrades in later learning stages.

The three baselines with cluster level features perform similarly as and do not show obvious advantage over their corresponding baselines without cluster level features, which further indicate that stirring global and local views together would not fully utilize the strength of the global view as co-testing, at least for the task of relation extraction.

To compare LGCo-Testing with semi-supervised learning, we simply take the best results in Chapter 3 where we used pattern clusters based on distributional similarities in bootstrapping, though we evaluated our system only on relations between names and only reported results for EMP-ORG, PHYS and PER-SOC. Their best F1 scores are 60 (EMP-ORG), 37 (PER-SOC), and 28 (PHYS), which are much lower than LGCo-Testing with 1000 labels. However, semi-supervised learning systems do not require human labeled examples except for the seeds. It is
impressive that with just a few seeds, semi-supervised learning can achieve F1 of 60 for the EMP-ORG relation. Combing it with active learning to further reduce the annotation cost is definitely a promising research avenue of our future work.

2.8 Related Work

To the best of our knowledge, this is the first piece of work in using active learning for relation type extension and the literature on this topic is rather limited. Our work is first motivated by co-training[7] and co-testing[60] which provide us with a solid theoretical foundation.

Our global view is mostly triggered by recent advances in using cluster level features for generalized discriminative models, including using word clusters[59] and phrase clusters[54] for name tagging and using word clusters for relation extraction[16][76].

Our multi-class setting is similar to the transfer learning setting of Jiang (2009), namely building up a system for a target relation with a few target instances and all auxiliary instances. They removed auxiliary instances from their evaluation data while we preserved auxiliary instances in our evaluation data, which unfortunately hinders a direct and fair comparison between their system and ours.

Perhaps the most relevant work is the our semi-supervised learning system in Chapter 3 which uses pattern clusters as an additional view for an enhanced confidence measure of learned name pairs. The two works differ in specific learning approaches and how the global view is used.
2.9 Conclusion

We have presented LGCo-Testing, a multi-view active learning approach for relation type extension based on \textit{local} and \textit{global} views. Evaluation results showed that LGCo-Testing can reduce annotation cost by 97\% while maintaining the performance level of supervised learning. It has prepared us well to apply active learning to real world relation type extension tasks. Combining it with semi-supervised learning to further reduce annotation cost is another promising research avenue.
Chapter 3

Relation Type Extension: Bootstrapping with Local and Global Data Views

3.1 Introduction

The Natural Language Processing (NLP) community faces new tasks and new domains all the time. Without enough labeled data of a new task or a new domain to conduct supervised learning, semi-supervised learning is particularly attractive to NLP researchers since it only requires a handful of labeled examples, known as seeds. Semi-supervised learning starts with these seeds to train an initial model; it then applies this model to a large volume of unlabeled data to get more labeled examples and adds the most confident ones as new seeds to re-train the model. This iterative procedure has been successfully applied to a variety of NLP tasks, such as hypernym/hyponym extraction[38], word sense disambiguation[87],
While semi-supervised learning can give good performance for many tasks, it is a procedure born with two defects. One is semantic drift. When semi-supervised learning is under-constrained, the semantics of newly promoted examples might stray away from the original meaning of seed examples as discussed in [11][24][15]. For example, a semi-supervised learning procedure to learn semantic patterns for the *Located-in* relation (PERSON in LOCATION/GPE) might accept patterns for the *Employment* relation (employee of GPE/ORGANIZATION) because many unlabeled pairs of names are connected by patterns belonging to multiple relations. Patterns connecting <Bill Clinton, Arkansas> include *Located-in* patterns such as “visit”, “arrive in” and “fly to”, but also patterns indicating other relations such as “governor of”, ”born in”, and “campaign in”. Similar analyses can be applied to many other examples such as <Bush, Texas> and <Schwarzenegger, California>. Without careful design, semi-supervised learning procedures usually accept bogus examples during certain iterations and hence the learning quality degrades.

The other shortcoming of semi-supervised learning is its lack of natural stopping criteria. Most semi-supervised learning algorithms either run a fixed number of iterations[2] or run against a separate labeled test set to find the best stopping criterion[1]. The former solution needs a human to keep eyeballing the learning quality of different iterations and set ad-hoc thresholds accordingly. The latter requires a separate labeled test set for each new task or domain. They make semi-supervised learning less appealing than it could be since the intention of using semi-supervised learning is to minimize supervision.

In this chapter, we propose a novel learning framework which can automatically monitor the semantic drift and find a natural stopping criterion for semi-supervised
Central to our idea is that instead of using unlabeled data directly in semi-supervised learning, we first cluster the seeds and unlabeled data in an unsupervised way before conducting semi-supervised learning. The semantics of unsupervised clusters are usually unknown. However, the cluster to which the seeds belong can serve as the target cluster. Then we guide the semi-supervised learning procedure using the target cluster. Under such learning settings, semantic drift can be automatically detected and a stopping criterion can be found: stopping the semi-supervised learning procedure when it tends to accept examples belonging to clusters other than the target cluster.

We demonstrate in this chapter the above general idea by considering a bootstrapping procedure to discover semantic patterns for extracting relations between named entities based on both local and global data views.

Traditional bootstrapping usually starts with some high-precision and high frequency seed patterns for a specific relation to match named entities. It evaluates the confidence of a newly matched name pair by looking at the confidence of each individual pattern that connects the name pair in isolation. We call this type of confidence the local view confidence. This dissertation presents a novel bootstrapping that moves one level up to build pattern clusters and estimates the confidence of a name pair based on the clusters of patterns as well. We call it the global view confidence. Using pattern clusters leads to a more reliable confidence estimation of a pair of named entities. Moreover, it uncovers that a pair of named entities might be connected by patterns indicating multiple relations by assuming that different clusters of patterns indicating different types of relations. We should give these pairs of names lower confidence. We then use newly promoted named entities to search for additional confident patterns connecting them. When patterns that
are being promoted have different cluster memberships from the seed patterns, semantic drift occurs and we should stop the bootstrapping process.

The next section describes our unsupervised pattern clusters. Section 3.3 presents the details of our novel bootstrapping procedure based on local and global data views. We evaluate our algorithms in Section 3.4 and present related work in Section 3.5. We draw conclusions and point to future work in Section 3.6.

3.2 Pattern Clusters

3.2.1 Distributional Hypothesis

The Distributional Hypothesis[36] states that words that tend to occur in similar contexts tend to have similar meanings. [53] extended this hypothesis to cover patterns (dependency paths in their case). The idea of the extension is that if two patterns tend to occur in similar contexts then the meanings of the patterns tend to be similar. For example, in “X solves Y” and “X finds a solution to Y”, “solves” and “finds a solution to” share many common Xs and Ys and hence are similar to each other. This extended distributional hypothesis serves as the basis on which we compute similarities for each pair of patterns.

3.2.2 Pattern Representation - Shortest Dependency Path

We adopt a shortest dependency path (SDP) representation of relation patterns. SDP has demonstrated its power in kernel methods for relation extraction[13]. Its capability in capturing most of the information of interest is also evidenced by a systematic comparison of effectiveness of different information extraction patterns
in[73]. For example, “nsubj <- met ->prep_in” is able to represent Located-in between “Gates” and “Seattle” while a token-based pattern would be much less general because it would have to specify all the intervening tokens.

Figure 3.1: Stanford dependency tree for sentence “Gates, Microsoft’s chairman, met with President Clinton in Seattle”.

3.2.3 Pre-processing

We tag and parse each sentence in our corpus with the NYU named entity tagger\(^2\) and the Stanford dependency parser\(^3\). Then for each pair of names in the dependency tree, we extract the SDP connecting them. Names in the path are replaced by their types. We require SDP to contain at least one verb or noun. We use the base form of words in SDP. We also require the length of the path (defined as the number of dependency relations and words in it) to be between 3 and 7. Short paths are more likely to be generic patterns such as “of” and can be handled

\(^1\)SDP is equivalent to the linked chains described in [73] when the dependency of a sentence is represented as a tree not a graph.
\(^2\)http://cs.nyu.edu/grishman/jet/license.html
\(^3\)http://nlp.stanford.edu/software/lex-parser.shtml
separately as in[62]. Very long paths are more likely to be non-relation patterns and too sparse to be useful even if they are relation patterns.

### 3.2.4 Clustering Algorithm

The basic idea of our clustering algorithm is to group all the paths (including the seed paths used later for semi-supervised learning) in our corpus into different clusters based on distributional similarities. We first extract a variety of features from the named entities $X$ and $Y$ connected by a path $P$ as shown in Table 3.1. We then compute an analogue of $tf-idf$ for each feature $f$ of $P$ as follows: $tf$ as the number of corpus instances of $P$ having feature $f$ divided by the number of instances of $P$; $idf$ as the total number of paths in the corpus divided by the number of paths with at least one instance with feature $f$. Then we adopt a vector space model, i.e., we construct a $tf-idf$ feature vector for each $P$. Now we compute the similarity between two vectors/paths using Cosine similarity and cluster all the paths using Complete Linkage.

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</tr>
<tr>
<td>Name Type of $Y$</td>
<td>RIGHT_GPE</td>
</tr>
<tr>
<td>Combination of Types of $X$ and $Y$</td>
<td>PERSON_GPE</td>
</tr>
<tr>
<td>Conjunction of String and Type of $X$</td>
<td>LEFT_Jordan_PERSON</td>
</tr>
<tr>
<td>Conjunction of String and Type of $Y$</td>
<td>RIGHT_China_GPE</td>
</tr>
<tr>
<td>Conjunction of Strings and Types of $X$ and $Y$</td>
<td>Jordan_PERSON_China_GPE</td>
</tr>
</tbody>
</table>

Table 3.1: Sample features for “$X$ visited $Y$” as in “Jordan visited China”

Some technical details deserve more attention here.

**Feature extraction:** We extract more types of features than the DIRT paraphrase discovery procedure used in [53]. [53] considered $X$ and $Y$ separately while
we also use the conjunction of X and Y. We also extract named entity types as features since we are interested in discovering relations among different types of names. Some names are ambiguous such as Jordan. We hope coupling the type with the string of the name may alleviate the ambiguity.

**Similarity measure and clustering method:** There are many ways to compute the similarity/distance between two feature vectors, such as Cosine, Euclidean, Hamming, and Jaccard coefficient. There are also many standard clustering algorithms. Our choice of the Cosine similarity measure and the Complete Linkage clustering algorithm are mostly driven by that they are the most popular similarity measure and clustering algorithm in NLP applications.

### 3.3 Semi-supervised Relation Pattern Discovery

We first present a standard bootstrapping algorithm which evaluates the confidence of a name pair based on local evidence only. And we call it *unguided bootstrapping*. Then we describe our new bootstrapping procedure whose confidence measure for name pairs is based on both local and global evidence. We call this *guided bootstrapping* because of the use of pattern clusters for better evaluating the confidence of name pairs and for detecting the stopping criterion.

#### 3.3.1 Unguided Bootstrapping

The procedure associates a precision between 0 and 1 with each pattern, and a confidence between 0 and 1 with each name pair. Initially the seed patterns for a specific relation $R$ have precision 1 and all other patterns 0. It consists of the following steps:
Step1: Use seed patterns to match new NE pairs and evaluate NE pairs.

Intuitively, for a newly matched NE pair, if many of the $k$ patterns connecting the two names are high-precision patterns then the name pair has a high confidence. The confidence is computed by the following formula.

$$Conf(N_i) = 1 - \prod_{j=1}^{k}(1 - Prec(p_j))$$  \hspace{1cm} (3.1)

Problem: While the intuition is correct, in practice this will over-rank NE pairs which are not only matched by patterns belonging to the target relation $R$ but are also connected by patterns of many other relations. This is because of the initial settings used in many semi-supervised learning systems: seeds are assigned high confidence. Thus all NE pairs matched by initial seed patterns will have very high confidence.

Suppose the target relation is Located-in, and “visited” is a seed pattern; then the <Clinton, Arkansas> example will be over-rated because we cannot take into account that it would also match patterns of other relations such as governorOf and birthPlace in a real corpus. This will cause a vicious circle, i.e., bogus NE pairs extract more bogus patterns which further extract more bogus NE pairs. We believe this flaw of the initial settings partially results in the semantic drift problem.

The problem happens because we are looking at the patterns that connect a name pair in isolation. If we insist to use only the local view to evaluate a name pair, i.e., by looking at the patterns in isolation, one can imagine that the problem can not be solved by using a different formula to replace the one presented here. A
possible solution is to study the structure of unlabeled data (NE pairs in our case) and integrate this structure information into the initial settings. Indeed, this is where pattern clusters come into play. We will demonstrate this in Section 3.3.2.

**Step 2: Use NE pairs to search for new patterns and rank patterns.**

Similar to the intuition in Step 1, for a pattern \( p \), if many of the NE pairs it matches are very confident then \( p \) has many supporters and should have a high ranking. We can use formula 3.2 to estimate the confidence of patterns and rank them.

\[
Conf(p) = \frac{Sup(p)}{|H|} \log Sup(p) \tag{3.2}
\]

Here \( |H| \) is the number of unique NE pairs matched by \( p \) and \( Sup(p) \) is the sum of the support it can get from the \( |H| \) pairs:

\[
Sup(p) = \sum_{j=1}^{|H|} Conf(N_j) \tag{3.3}
\]

The precision of \( p \) is given by the average confidence of the NE pairs matched by \( p \). Formula 3.4 normalizes the precision to range from 0 to 1. As a result the confidence of each NE pair is also normalized to between 0 and 1.

\[
Prec(p) = \frac{Sup(p)}{|H|} \tag{3.4}
\]

**Step 3: Accept patterns**

Most systems accept the \( K \) top ranked patterns in Step 2 as new seeds, subject to some restrictions such as requiring the differences of confidence of the \( K \) patterns to be within a small range.

**Step 4: Loop or stop**
The procedure now decides whether to repeat from Step 1 or to terminate. Most systems simply do not know when to stop. They either run a fixed number of iterations or use some held-out data to find one criterion that works the best for the held-out data.

### 3.3.2 Guided Bootstrapping

Recall that our clustering algorithm in Section 3.2 provides us with $K$ clusters, each of which contains $n$ ($n$ differs in different clusters) patterns. Every pattern in our corpus now has a cluster membership (the seed patterns have the same membership).

The most important benefit from our pattern clusters is that now we can measure how strongly a NE pair $N_i$ is associated with our target cluster $C_t$ (the one to which the seed patterns belong).

$$Prob(N_i \in C_t) = \frac{\sum_{p \in C_t} freq(N_i, p)}{m} \tag{3.5}$$

Here $freq(N_i, p)$ is the number of times $p$ matches $N_i$ and $m$ is the total number of pattern instances matching $N_i$.

We integrate this prior cluster distribution of each NE pair into the initial settings of our new bootstrapping procedure.

**Step1:** Use seed patterns to match new NE pairs and evaluate NE pairs.

Assumption: A good NE pair must be strongly associated with the target cluster and can be matched by multiple high-precision patterns. In other words, it has to have strong support from both the global and local evidence.
So we evaluate a NE pair by the harmonic mean of two confidence scores, namely the *global confidence* as its association with the *target* cluster and the *local confidence* given by the patterns matching it.

\[
Conf(N_i) = \frac{2 \times Local\_Conf(N_i) \times Global\_Conf(N_i)}{Local\_Conf(N_i) + Global\_Conf(N_i)} \tag{3.6}
\]

\[
Local\_Conf(N_i) = 1 - \prod_{j=1}^{k}(1 - Prec(p_j)) \tag{3.7}
\]

\[
Global\_Conf(N_i) = \text{Prob}(N_i \in C_t) \tag{3.8}
\]

Under such settings, \(<\text{Clinton, Arkansas}>\) will be assigned a lower confidence score for the *Located-in* relation than it is in the unguided bootstrapping. Even if we assign high precision to our seed patterns such as “visited” and consequently the \(Local\_Conf\) is very high, it can still be discounted by the \(Global\_Conf\). The \(Global\_Conf\) of \(<\text{Clinton, Arkansas}>\) related to the *Located-in* relation is indeed very low (less than 0.1) in our experiments.

**Step 2: Use NE pairs to search for new patterns and rank patterns.**

All the measurement functions are the same as those used in the unguided bootstrapping. However, with better ranking of NE pairs in Step 1, the patterns are also ranked better than they are in the unguided bootstrapping.

**Step 3: Accept patterns**

We also accept the \(K\) top ranked patterns.

**Step 4: Loop or stop**

Since each pattern in our corpus has a cluster membership, we can monitor the *semantic drift* easily and naturally stop: it drifts when the procedure tries to
accept patterns which do not belong to the target cluster; we can stop when the procedure tends to accept more patterns outside of the target cluster.

If our clustering algorithm can give us perfect pattern clusters, we can stop bootstrapping immediately after it accepts the first pattern not belonging to the target cluster. Then the bootstrapping becomes redundant since all it does is to consume the patterns of the target cluster.

Facing the reality of the behavior of many clustering algorithms, we allow the procedure to occasionally accept patterns outside of the target cluster but we are not tolerant when it tries to accept more patterns outside of the target cluster than patterns in it. Note that when such patterns are accepted they will be moved to the target cluster and invoke the re-computation of Global_Conf of NE pairs connected by these patterns. The ranking functions in step 1 and 2 insure that the procedure will only accept patterns which can gain strong support from NE pairs that are strongly associated with the target cluster and are connected by many confident patterns.

3.4 Experiments

3.4.1 Corpus

Our corpora contain 37 years of news articles: TDT5, NYT(94-00), APW(98-00), XINHUA(96-00), WSJ(94-96), LATWP(94-97), REUFF(94-96), REUTE(94-96), and WSJSF(87-94). It contains roughly 65 million sentences and 1.3 billion tokens.
3.4.2 Seeds

Seeds of the 3 relations we are going to test are given in table 3.2. Located-in detects relation between PERSON and LOCATION/GPE; Social (SOC) detects social relations (either business or family) between PERSON and PERSON; Employment (EMP) detects employment relations between PERSON and ORGANIZATION.

<table>
<thead>
<tr>
<th>Relation</th>
<th>Seeds</th>
</tr>
</thead>
<tbody>
<tr>
<td>locatedIn</td>
<td>nsubj’ visit dobj</td>
</tr>
<tr>
<td></td>
<td>nsubj’ travel prep_to</td>
</tr>
<tr>
<td></td>
<td>poss’ trip prep_to</td>
</tr>
<tr>
<td>SOC</td>
<td>appos friend/lawyer poss</td>
</tr>
<tr>
<td></td>
<td>appos son/spokesman prep_of/prep_for</td>
</tr>
<tr>
<td></td>
<td>nsubj’ fire dobj</td>
</tr>
<tr>
<td></td>
<td>nsubjpass’ fire agent</td>
</tr>
<tr>
<td>EMP</td>
<td>appos chairman/executive/founder prep_of</td>
</tr>
<tr>
<td></td>
<td>appos editor prep_of</td>
</tr>
<tr>
<td></td>
<td>appos director/head/officer/analyst prep_at</td>
</tr>
<tr>
<td></td>
<td>appos manager prep_with</td>
</tr>
</tbody>
</table>

Table 3.2: Seed patterns.

(nsubj, dobj, prep, appos, poss, nsubjpass, agent stand for subject, direct object, preposition, apposition, possessive, passive nominal subject and complement of passive verb). The quote marks in Table 3.2 and Table 3.3 denote inverse dependencies in the dependency path. We provide more seeds (executives and staff) for EMP because it has been pointed out in [74] that EMP contains a lot of job titles.

We work on these three relations mainly because of the availability of benchmark evaluation data. These are the most frequent relations in our evaluation
3.4.3 Unsupervised Experiments

We run the clustering algorithm described in Section 3.2 using all the 37 years’
data. We require that a pattern match at least 7 distinct NE pairs and that an
NE pair must be connected by at least 7 unique patterns. As a result, there are
635,128 patterns (22,225 unique ones) used in experiments. We use 0.005 as the
cutoff threshold of Complete Linkage. The threshold is decided by trying a series
of thresholds and searching for the maximal\(^4\) one that is capable of placing the seed
patterns for each relation into a single cluster. Table 3.3 shows the top 15 patterns
(ranked by their corpus frequency) of the cluster into which our Located-in seeds
fall.

3.4.4 Semi-supervised Experiments

To provide strong statistical evidence, we divide our data into 10 folds (com-
binations of news articles from different years and different news resources). We
then run both the unguided and guided bootstrapping on the 10 folds. For both
procedures, we accept \(n\) patterns in a single iteration (\(n\) is initialized to 2 and set
to \(n + 1\) after each iteration). We run 50 iterations in the unguided bootstrapping
and 1,325 patterns are accepted for each fold and each relation. Our guided boot-
strapping procedure stops when there are two consecutive iterations in which more
than half of the newly accepted patterns do not belong to the target cluster. Thus
the number of patterns accepted for each fold and each relation differs as the last

\(^4\)We choose the maximal value because many clusters will be merged to a single one when
the threshold is close to 0, making the clusters too general to be useful.
Table 3.3: Top 15 patterns in the Located-in Cluster.

### 3.4.5 Evaluation

The output of our bootstrapping procedures is 60 sets of patterns (20 sets per relation). We need a data set and evaluation method which can compare their effectiveness equally and consistently.

Evaluation data: ACE 2004 training data. ACE does not provide relation annotation between each pair of names. For example, in “US President Clinton said that the United States ...” ACE annotates an EMP relation between the name “US” and nominal “President”. There is no annotation between “US” and “Clinton”. However, it provides entity co-reference information which connects “President” to “Clinton”. So we take advantage of this entity co-reference information to automatically re-annotate the relations where possible to link a pair of names
within a single sentence. The re-annotation yields an EMP relation between “US” and “Clinton”. The re-annotation is reviewed by hand to avoid adding a relation linking “Clinton” and the more distant co-referent “United States”, even though “US” and “the United States” refer to the same entity. This data set provides us with 412/3492 positive/negative relation instances between names. Among the 412 positive instances, there are 188/117/35 instances for EMP/Located-in/SOC relations.

Evaluation method: We adopt a direct evaluation method, i.e., use our sets of patterns to extract relations between names on ACE data. Applying patterns to a benchmark data set can provide us with better precision/recall analyses. We use a strict pattern match strategy. We can certainly take advantage of loose match or add patterns as additional features to feature-based relation extraction systems to boost our performance but we do not want these to complicate the comparison of the guided and unguided bootstrapping procedures.

3.4.6 Results and Analyses

We average our results on the 10 folds. We plot precision against recall and semantic drift rate against iterations (Drift). We compute the semantic drift rate as the percentage of false positive instances belonging to ACE relations other than the target relation. Take EMP for example, we compute how many of the false positive instances belonging to other relations such as Located-in, SOC and other ACE relations. In all plots, red solid lines represent guided bootstrapping and blue dotted lines unguided bootstrapping.

There are a number of conclusions that can be drawn from these results. We are particularly interested in the following two questions: To what extent did we
Figure 3.2: Precision-recall curve and Drift Rate for EMP/Located-in/SOC

prevent semantic drift by the guidance of pattern clusters? Did we stop at the right point, i.e., can we keep high precision while maintaining near maximal recall?
1. It is obvious from the drift curves that our guided bootstrapping effectively prevents semantic drift. Indeed, there is no drift at all when Located-in and SOC learners terminate. Although drift indeed occurs in the EMP relation, its curve is much lower than that of the unguided bootstrapping.

2. Our guided bootstrapping terminates when the precision is still high while maintaining a reasonable recall. Our bootstrapping for EMP/SOC/Located-in terminates at F-measures of 60/37/28 (in percentage). We conducted the Wilcoxon Matched-Pairs Signed-Ranks Test on the 10 folds, comparing the F-measures of the last iteration of our bootstrapping guided by clusters and the iteration which provides the best average F-measure over the 3 relations of the unguided bootstrapping. The results show that the improvement of using clusters to guide bootstrapping is significant at a 97% confidence level.

We hypothesize that when working on dozens or hundreds of relations the gain of our procedure will be even bigger since we can effectively prevent inter-class errors.

3.5 Related Work

Recent research starts exploring unlabeled data for discriminative learning. [59] augmented name tagging training data with hierarchical word clusters and encoded cluster membership in features for improving name tagging. [54] further explored a two-stage cluster-based approach: first clustering phrases and then relying on a supervised learner to identify useful clusters and assign proper weights to cluster features. Other similar work includes [82] for name tagging, and [49] for dependency parsing.
While similar in spirit, our supervision is minimal, i.e., we only use a few seeds while the above approaches rely on a large amount of labeled data. To the best of our knowledge, the theme explored in this chapter is the first study of using pattern clusters for preventing semantic drift in semi-supervised pattern discovery.

Recent research also explored the idea of driving semi-supervised learning with explicit constraints constructed by hand such as identifying mutual exclusion of different categories (i.e., people and sport are mutually exclusive). This is termed constraint-driven learning in [17], coupled learning in [15] and counter-training in [84]. The learning quality largely depends on the completeness of explicit constraints. While we share the same goal, i.e., to prevent semantic drift, we rely on unsupervised clusters to discover implicit constraints for us instead of generating constraints by hand.

Our research is also close to semi-supervised IE pattern learners including [67], [2], [85], and many others. While they conduct bootstrapping on unlabeled data directly, we first cluster unlabeled data and then bootstrap with help from clusters.

There are also clear connections to work on unsupervised relation discovery [37][90][69]. They group pairs of names into relation clusters based on the contexts between names while we group the contexts/patterns into clusters based on features extracted from names.

3.6 Conclusions and Future Work

We presented a simple bootstrapping procedure which utilized both local and global evidence for an enhanced confidence measure of name pairs. This novel procedure can achieve the best F-1 score while maintaining a good trade-off between
precision and recall. We also demonstrated that it can effectively prevent *semantic drift* and naturally terminate.

We plan to extend this idea to improve relation extraction performance with a richer model as used in [89] and [94] than a simple pattern learner. The feature space will be much larger than the one adopted in this chapter. We will investigate how to overcome the memory bottleneck when we apply rich models to millions of instances.
Chapter 4

Cross-Domain Bootstrapping for Named Entity Recognition

4.1 Introduction

Named Entity Recognition (NER) is a fundamental information extraction task with the objective of identifying and classifying proper names into certain predefined categories such as persons, organizations and locations. Supervised NER systems perform well when they are trained and tested on data from the same domain. However, when testing on a new domain which is different or even slightly different from the domain they were trained on, their performance usually degrades dramatically. For example, [21] reported that a system trained on the CoNLL 2003 Reuters dataset achieved an F-measure of 0.908 when it was tested on a similar Reuters corpus but only 0.643 on a Wall Street Journal dataset.

The performance degradation phenomenon occurs when one has access to labeled data in one domain (the source domain) but has no labeled data in another
domain (the target domain). This is a typical situation as one might be able to expend the limited effort required to annotate a few target examples as a test bed but cannot afford to annotate additional examples for training purpose. However, it is usually the case that we have access to abundant unlabeled data in the target domain.

This chapter works on this common scenario where we have access to labeled data in the source domain and only unlabeled data in the target domain. We propose a cross-domain bootstrapping (CDB) algorithm to iteratively adapt the source domain model to the target domain. Specifically, we first train an MEMM (maximum entropy Markov model[56]) source/seed model using the labeled data in the source domain and then apply it to the unlabeled data pool of the target domain. We then select good instances based on multiple criteria and use them to re-train and upgrade the seed model.

CDB differs from previous bootstrapping algorithms in several aspects. First, the seed model is generalized with word clusters. A model trained on the source domain may perform poorly on the target domain partly because it relies on local evidence such as features involving the word identities and there are many target domain specific words (for both names and context words) that have not been observed in the source domain. This motivates our work to use word clusters as a type of global evidence to extract additional features to generalize the seed model. The assumption is that even if we have not observed a target word $W_t$ in the source domain, another word $W_s$ in the source domain might share the same cluster membership with the word $W_t$. The cluster level feature still fires even if the lexical feature is absent from the source domain. More specifically, we mix the labeled source domain corpus with the unlabeled target domain corpus
and generate the Brown word clusters[12] from this joint corpus. We then extract cluster memberships as features to augment the feature based NER system trained on the source domain.

CDB is novel in its multi-criteria-based instance selection method. Standard bootstrapping usually adopts a single criterion which is based on the confidence measure only, promoting those instances that are most confidently labeled from the unlabeled data. This might not be a problem when the data used for training the seed model and the unlabeled data are drawn from the same domain. However, in our cross domain setting, the most confidently labeled examples are those that have been observed in or are most similar to the source domain. CDB uses multiple criteria to select instances that are novel, confident, representative and diverse. It first uses novelty as a filter, maintaining only these instances that are specific to the target domain. It then ranks these novel instances based on a confidence measure at both local and global levels. Top ranked instances contribute to a candidate set. Finally, it applies representativeness and diversity measures, which again are based on global evidence, to all the candidates and selects a subset of them for promotion.

The rest of this chapter is organized as follows: The next section positions us with respect to related work. Section 4.3 briefly introduces our NER task and source and target domains. Section 4.4 describes the CDB algorithm in detail. We present an experimental study in Section 4.5 and conclude in Section 4.6.
4.2 Related Work

There is a large body of domain adaptation research on different NLP tasks. Here we only discuss work related to NER.

Supervised domain adaptation for NER works on the scenario where one has labeled data from both the source and the target domains\cite{25}\cite{28}. \cite{25} has shown that a better model can be learned from the labeled data by making three copies of the features: general, source-dependent and target-dependent. Without labeled data from the target domain, it is impossible to distinguish and jointly learn the three types of features. Our work also generalizes and augments features but is obviously different from the above approaches in that the word cluster features are extracted from an unlabeled corpus.

Semi-supervised domain adaptation for NER deals with the situation such as ours where one only has labeled data from the source domain but not the target domain\cite{46}\cite{83}. (We can also refer to this branch of research as unsupervised learning because there is no supervision from the target domain.) \cite{46} studied domain adaptation from an instance weighting perspective and proposed a balanced bootstrapping algorithm in which the small number of instances promoted from the target domain was re-weighted to have an equal weight to the large number of source instances. Their instance selection was based on a confidence measure. \cite{83} described a domain adaptive bootstrapping framework where the instances were selected based on informativeness. Neither of the two approaches generalized their seed models as we have done and both of them used a single instance selection criterion instead of the multiple criteria we have used.

Standard bootstrapping for domain-specific NER or semantic lexicon acquisition works on the target domain directly (both the seed examples and the unlabeled
data are from the target domain) and typically adopts a confidence measure for selecting new instances[39][57][67][86]. It has been shown that seed selection is very important for standard bootstrapping[81]. The way we generalize our seed model is similar, but not identical to seed selection in a sense that both of the approaches try to provide a better starting point for bootstrapping.

4.3 Task and Domains

Our NER task is similar to those defined in some benchmark evaluations such as MUC-6[34], CoNLL-2003[79] and ACE-05. Given a raw sentence, the goal is to identify name expressions and classify them into one of the following three types: PER (person), ORG (organization) and GPE (Geo-Political entity). We choose to work with these three types as they are the most frequent ones in our target domain. Figure 4.1 illustrates examples from both domains.

![Source Example](image)

Target Example1: The ruler of Saudi Arabia is Fahad bin Abdul Aziz bin Abdul Rahman Al-Sa’ud.

Target Example2: where Sheikh Abdul Sattar al-Rishawi and the Anbar Salvation Front became a force for stability.

Figure 4.1: Examples of NER task and domains

Our target domain documents are from publicly available reports (in English) on terrorism, such as those from the Combating Terrorism Center at West Point\(^2\). There are many domain-specific characteristics of our target domain. Just to mention a few: it is noisy as we automatically convert PDF documents to text files (footnotes might be inserted in the middle of natural language sentences and punctuation might not be converted correctly such as the example “Al-Sa?ud”); it is Arabic name (transliterated) rich and the naming convention is different from English names; name variation is another noticeable problem.

We choose the ACE-05 annotated data as the source domain because the degree of overlap between the ACE-05 data and the target domain data is higher than the MUC-6 and the CoNLL-2003 datasets, which are from the 90s.

4.4 Cross-Domain Bootstrapping

We first present an overview of the CDB algorithm, and then we describe in detail the generalization of a seed model with word clusters and the multi-criteria-based instance selection method.

4.4.1 Overview of the CDB Algorithm

The input to the algorithm is a labeled dataset from the source domain and an unlabeled dataset from the target domain, denoted by $D^L_S$ and $D^U_T$ respectively. Let $G$ denote the growing set which contains selected instances during each round $t$ and is initialized to be an empty set at round 0; the CDB algorithm repeats the following steps until it meets a stopping criterion.

\(^2\)http://www.ctc.usma.edu/publications/sentinel
1. Train an NER seed model $M_t$ with $D^l_S \cup G_t$, generalize it with word clusters
2. Label $D^U_T$ using $M_t$
3. Select $D^L_T \subseteq D^U_T$ based on multiple criteria
4. Update: $G_{t+1} = G_t \cup D^L_T$ and $D^U_T = D^U_T \setminus D^L_T$

The output of the CDB algorithm is an NER model which will be used to identify and classify named entities in the target domain. It is important to mention that the seed model $M$ is generalized with word clusters at each round, not just at the beginning of the algorithm.

4.4.2 Seed Model Generalization

Ungeneralized seed model: NER is typically viewed as a sequential prediction problem. Given a sentence containing a sequence of tokens, the goal is to assign a name class to each one of the tokens. Formally, let $S = (t_1...t_N)$ be an input sequence of tokens and $C = (c_1...c_N)$ be the output sequence of name classes, the prediction problem is to estimate the probability $P(C|S)$.

To facilitate the learning procedure, we use the standard BIO decoding scheme. Each name type $c$, other than the type $O$ (not a name), is split into subtypes $B$-$c$ (beginning of $c$) and $I$-$c$ (continuation of $c$). Although the less used BILOU (beginning, inside, last, outside and unit-length) scheme was claimed to outperform the BIO scheme in [65], we did not observe the same behavior in our target domain (see Section 4.5.2). The BIO representation gives us 7 classes (3 name types $\times$ 2 subtypes $+$ 1 not a name class).

We build an MEMM[56] model with the following customary features:

1. current token $t_i$
2. lower case tokens in a 5-token-window \( t_{i-2}t_{i-1}t_it_{i+1}t_{i+2} \)

3. a word type feature (all-capitalized, initial-capitalized, all-digits, etc.) for each token in the context window

4. previous prediction \( c_{i-1} \)

5. conjunction of \( c_{i-1} , 1, 2 \) and 3

6. gazetteer membership of context tokens in a dictionary of country and US state names.

Note that we do not extract POS tags as features since a good target domain POS tagger is not available to us.

As the model makes a local decision, that is, it predicts the name class for each individual token based on its feature vector, it might produce illegal transitions between name classes (e.g., B-PER followed by I-ORG). So we run the Viterbi algorithm to select the sequence of name classes with the highest probability.

**Generalizing seed model with word clusters:** The sparsity of lexical features is a notorious problem in many supervised NLP systems. Recent advances in generating word classes from unlabeled corpora and adding them as features has proven to be an effective way of generalizing the local lexical features to alleviate sparsity\[59\][65][80]. The unavailability of a cross-domain unlabeled corpus hinders the direct adaptation of this technique to our cross-domain setting. Ideally, we would prefer an unlabeled corpus containing words of both domains so that the word classes can generalize for both domains. So we propose to generate a joint corpus by mixing the labeled source data with the unlabeled target data.

We then follow previous research and use the Brown algorithm\[12\] to generate word clusters from the joint corpus. The Brown algorithm is a hierarchical clustering algorithm which initially assigns each word to its own cluster and then
repeatedly merges the two clusters which cause the least loss in average mutual information between adjacent clusters based on bigram statistics. By tracing the pairwise merging steps, one can obtain a word hierarchy which can be represented as a binary tree. A word can be compactly represented as a bit string by following the path from the root to itself in the tree, assigning a 0 for each left branch, and a 1 for each right branch. Table 4.1 shows some words and their bit string representations obtained from the joint corpus.

By using prefixes of different lengths one can produce word clusters of various granularities so as to avoid the commitment to a single cluster. We used clusters with lengths 4, 6, 10 and 20 and augmented the previous, current and next token features with word clusters[59][65][80]. For example, when we extract features for the current token “John”, we will add a cluster feature $curPrefix_6=110100$ when we use length 6. (Note that the cluster feature is a nominal feature, not to be confused with an integer feature.) Now, even if we have not observed “Abdul” in our source domain, its cluster level feature still fires given that the $curPrefix_6$ feature is the same for both “John” and “Abdul”.

<table>
<thead>
<tr>
<th>Bit string</th>
<th>Examples</th>
</tr>
</thead>
<tbody>
<tr>
<td>110100011</td>
<td>John, James, Mike, Steven, Dan, ...</td>
</tr>
<tr>
<td>11010011101</td>
<td>Abdul, Mustafa, Abi, Abdei, ...</td>
</tr>
<tr>
<td>11010011111</td>
<td>Shaikh, Shaykh, Sheikh, Sheik, ...</td>
</tr>
<tr>
<td>1101000011</td>
<td>President, Pope, Vice, ...</td>
</tr>
<tr>
<td>111111110</td>
<td>Qaeda, Qaida, qaeda, QAEDA, ...</td>
</tr>
<tr>
<td>00011110000</td>
<td>FDA, NYPD, FBI, ...</td>
</tr>
<tr>
<td>000111100100</td>
<td>Taliban, ...</td>
</tr>
</tbody>
</table>

Table 4.1: An example of words and their bit string representations. Bold ones are transliterated Arabic words.

It is worth mentioning an additional benefit of using word clusters: different
Arabic name variants are grouped together such as variants of “Shaikh” and variants of “Qaeda” in Table 4.1. Without analyzing and comparing the internal structure of the words, such as computing the edit distance between different words, the clustering algorithm itself is able to capture this domain-specific knowledge.

4.4.3 Multi-Criteria-based Instance Selection

Most standard bootstrapping algorithms use a confidence measure as the single selection criterion. In practice, this works well under the single domain setting. In a cross-domain setting like ours, the most confidently labeled instances are highly correlated with the source domain and hence contain little information about the target domain. In contrast, the CDB algorithm adopts an instance selection method based on multiple criteria.

**Instance:** We define an instance $I = \langle v, c \rangle$ as the feature vector $v$ and the name class $c$ of the current token $t_i$ under consideration. Although sentence seems to be a more natural unit than token for a bootstrapped NER system[41], our sentences contain many target domain specific names and context words which undermines the reliability of any confidence measure defined over a sentence. However, when broken down to the token level, it is easy to design a relatively reliable confidence measure, as the feature vector $v$ is essentially extracted from a short context window $t_{i-2}t_{i-1}t_it_{i+1}t_{i+2}$. Also, the feature vector does contain the transition from the previous name class to the current class as we include the prediction of the previous token $t_{i-1}$ as a feature (The class of the previous token is known after we run Viterbi over the whole sentence). Moreover, the NER model outputs normalized probabilities predicting the name classes based on the vector $v$ and it is convenient to add the vector to the feature file for re-training the NER model.
**Novelty:** Novelty prefers an instance $I$ that contains target-domain-specific tokens in its context window and can be confidently labeled by the seed model. If all the context tokens have been observed in the source domain then the instance contains less target domain information than others. However, if all the 5 tokens are target-domain-dependent then the seed model’s prediction of the instance might not be reliable. So we tried different values (1, 2 and 3) for the number of target-domain-specific tokens in the context window and different positions (token index in the range $[i-2, i+2]$) and found that the following simple measure worked the best: if the current token $t_i$ is the only target-domain-specific token then the instance is considered to be novel.

**Confidence:** A reliable confidence measure is crucial to the success of a bootstrapping algorithm. If one bogus instance is selected, it will lead to the selection of many other bogus instances. CDB’s confidence measure not only considers how confident an instance is labeled locally but also globally.

The local confidence of an instance $I$ is defined as the posterior entropy of the 7 name classes $C$ given the instance’s feature vector $v$.

$$LocalConf(I) = - \sum_{c_i} p(c_i|v) \log p(c_i|v) \quad (4.1)$$

It is non-negative. It achieves its minimum value 0 when the MEMM model predicts a class $c_i$ with probability 1 (more precisely when $p(c_i|v) = 1$). It achieves its maximum value when the predictions are evenly distributed over the 7 classes. So the lower the value, the more confident the instance is.

The global confidence concerns how other occurrences of the current token $t_i$ in the instance $I$ are labeled in the whole corpus. The linguistic intuition here
is that one name usually belongs to one class in a corpus\cite{29}\cite{51}\cite{82}. The CDB algorithm would prefer to select name tokens that can be consistently labeled in the whole corpus. So we gather all other occurrences of \( t_i \) in the corpus, generate their feature vectors and take as the name class of each occurrence the one with the highest probability returned by the MEMM model. The BI tags of the class are then deleted, for example, B-PER would become PER. This is because a name token can be at different positions (e.g., Smith can be either B-PER or I-PER). So global confidence uses 4 name classes instead of 7. We then compute the global confidence as below:

\[
GlobalConf(I) = -\sum_{c_i} p(c_i) \log p(c_i)
\]  

(4.2)

where \( p(c_i) \) is the corpus level probability of \( t_i \) belonging to the class \( c_i \). It is defined as the number of times \( t_i \) is predicted with the class \( c_i \) divided by the total number of occurrences of \( t_i \) in the corpus. For example, if “Abdul” appears 10 times in the corpus and is predicted 8 times as PER, then the probability of “Abdul” belonging to the class PER is 0.8. Similar to the local confidence measure, the lower the value of the global confidence, the more confident the instance is.

We then propose a final measure to combine the two confidence measures. We simply take the product of the two measures.

\[
ComConf(I) = LocalConf(I) \times GlobalConf(I)
\]  

(4.3)

**Density:** In addition to the most confident instances, CDB also aims to select the most representative instances. We use a density measure to evaluate the representativeness of an instance. The density of an instance \( i \) is defined as the
average similarity between \( i \) and all other instances \( j \) in the corpus. The most *representative* instance is the one with the largest *density* value.

\[
Density(i) = \frac{\sum_{j=1\land j\neq i}^{N} Sim(i, j)}{N - 1} \quad (4.4)
\]

where \( N \) is the total number of instances in the corpus and \( Sim(i, j) \) is the similarity between the two instances, which is defined as the standard Jaccard Similarity between the feature vectors \( u \) and \( v \) of the two instances. The Jaccard Similarity between \( u \) and \( v \) is defined as the number of matched features of \( u \) and \( v \) divided by the number of unique features in the union of \( u \) and \( v \). The match function for a feature \( f \) returns 1 if the values of \( f \) in \( u \) and \( v \) are the same and 0 otherwise.

Alternatively, we could find the angle between the two feature vectors and compute the Cosine Similarity between them. However, as all the features for NER take discrete values the simpler Jaccard Similarity suffices to capture the similarity between feature vectors.

**Diversity:** CDB also aims to select instances as diverse as possible. Intuitively, if we have observed an instance and its similar instances a sufficient number of times then we cannot learn more new information from them. Take the instance “, said * in his” for example, where * is the current token, which we restrict to be a target-domain-specific token (*novelty*) and is highly likely to be a person; it is confident at both *local* and *global* levels given that the context is salient and it is probably very dense too. However, repeatedly selecting such instances is a waste of time because no additional benefit can be gained for CDB.

So *globally*, once an instance has been selected, it is removed from the unlabeled target corpus. The CDB algorithm will never select it again in the following rounds. *Locally*, in a single round, when we evaluate an instance \( i \), we will compare the
difference between $i$ and all the instances $j$ that have already been selected. If the difference is large enough, we accept $i$; otherwise we reject it. One possibility of measuring the difference is to directly use the similarity measure $Sim(i, j)$. But this tends to reduce the chance of selecting dense instances given that a dense instance has many similar instances and tends to occur more frequently than others. For example, if we already selected the instance “, said Abdul in his”, the chance of selecting other similar instances “, said * in his” is low. We then turn to a compromise measure to compute the difference between instances $i$ and $j$ which is defined as the difference of their density values. By setting a small threshold for $diff(i, j)$, dense instances still have a higher chance to be selected while a certain degree of diversity is achieved at the same time.

$$diff(i, j) = Density(i) - Dentisty(j)$$  (4.5)

**Order of applying different criteria:** CDB first applies the novelty measure to all the instances in the corpus to filter out non-novel instances, and then it computes the confidence score for each novel instance. Instances are then ranked in increasing order of confidence score (lower value means higher confidence) and the top ranked $M$ instances will be used to generate a candidate set. CDB now applies the density measure to all the members in the candidate set and ranks the instances in descending order of density (larger value means higher density). Finally, CDB accepts the first instance (with the highest density) in the candidate set and selects other candidates based on the $diff$ measure.
4.5 Experiments

4.5.1 Data, Evaluation and Parameters

Source domain data: Table 4.2 summarizes the source domain data (ACE 2005) used in this paper. The 2005 dataset contains 6 genres: Broadcast Conversations (bc), Broadcast News (bn), Conversational Telephone Speech (cts), Newswire (nw), Usenet (un) and Weblog (wl). We randomly selected 10 documents from each genre for testing purposes.

<table>
<thead>
<tr>
<th>Genre</th>
<th>Training (#doc)</th>
<th>Test (#doc)</th>
</tr>
</thead>
<tbody>
<tr>
<td>bc</td>
<td>50</td>
<td>10</td>
</tr>
<tr>
<td>bn</td>
<td>216</td>
<td>10</td>
</tr>
<tr>
<td>cts</td>
<td>29</td>
<td>10</td>
</tr>
<tr>
<td>nw</td>
<td>97</td>
<td>10</td>
</tr>
<tr>
<td>un</td>
<td>39</td>
<td>10</td>
</tr>
<tr>
<td>wl</td>
<td>109</td>
<td>10</td>
</tr>
<tr>
<td>Total</td>
<td>540 (285K words)</td>
<td>60 (31K words)</td>
</tr>
</tbody>
</table>

Table 4.2: source domain data.

Target domain data: Table 4.3 lists the sizes of the unlabeled and the labeled corpus as well as the number of instances of the 3 name types in the labeled corpus. The labeled data (for testing purpose) is annotated according to the ACE 2005 guideline\(^3\).

Corpora for generating word clusters: To study the impact of unlabeled corpora on cross-domain NER, we downloaded the word clusters generated by [65]\(^4\) and [80]\(^5\). Following them, we used Liang’s implementation of the Brown algorithm

\(^3\)http://projects.ldc.upenn.edu/ace/docs/English-Entities-Guidelines_v5.6.1.pdf
\(^4\)http://cogcomp.cs.illinois.edu/page/software_view/4
\(^5\)http://metaoptimize.com/projects/wordreprs/
<table>
<thead>
<tr>
<th>Data</th>
<th>Size</th>
</tr>
</thead>
<tbody>
<tr>
<td>Unlabeled/Labeled</td>
<td>10M/23K words</td>
</tr>
<tr>
<td>PERSON</td>
<td>771 instances</td>
</tr>
<tr>
<td>ORGANIZATION</td>
<td>585 instances</td>
</tr>
<tr>
<td>GPE</td>
<td>559 instances</td>
</tr>
</tbody>
</table>

Table 4.3: *target* domain data.

and generated 1,000 word clusters for both the TDT5 (the English portion) and the joint corpus[52]. The TDT5 is selected because it contains news from the year 2003 and some ACE 2005 training documents are also from 2003.

<table>
<thead>
<tr>
<th>Data</th>
<th>Size(#words)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reuters 1996 (from [64])</td>
<td>43M</td>
</tr>
<tr>
<td>Cleaned RCV1 (from [78])</td>
<td>37M</td>
</tr>
<tr>
<td>TDT5 (LDC2006T18)</td>
<td>83M</td>
</tr>
<tr>
<td>Joint Corpus (ACE training + Unlabeled <em>target</em> data)</td>
<td>10M</td>
</tr>
</tbody>
</table>

Table 4.4: Corpora for generating word clusters.

**Evaluation:** Evaluation is done at the named entity level, not the BIO tags level. Boundary errors are penalized. We use the CoNLL scoring metric and report precision, recall and F1 scores.

**Parameters:** As the CDB algorithm uses several parameters, we summarize them in Table 4.5 for easy reference. Because CDB runs the Viterbi algorithm on each sentence, it is time consuming to run it on the whole unlabeled data. So we divided them sequentially into 6 batch sets and picked a random set for bootstrapping in each iteration. The candidate set contains more instances than the CDB algorithm will actually select because of the density and diversity measures. As the majority of tokens belong to the not-a-name class, we select the same amount of name/not-a-name instances in order to provide a balanced distribution between
the name and the not-a-name classes. We tried many different parameter values and those in Table 4.5 were selected by eyeballing the quality of selected instances.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Size or Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Batch Set</td>
<td>60K sentences (roughly 1.7M tokens)</td>
</tr>
<tr>
<td>Candidate Set</td>
<td>2000/2000 name/not-a-name instances</td>
</tr>
<tr>
<td>$D_f^t$</td>
<td>300/300 name/not-a-name instances</td>
</tr>
<tr>
<td>Iterations</td>
<td>30</td>
</tr>
<tr>
<td>$diff(i, j)$</td>
<td>0.001</td>
</tr>
</tbody>
</table>

Table 4.5: Parameters for CDB.

4.5.2 Performance of the source Domain Model

We build two source models using the ACE 2005 training data in Table 4.2. The first model is an HMM model with the BILOU states encoding. The second model is the MEMM model with the BIO encoding using the conventional features as described in Section 4.4.2 (no word cluster features). Their performances on both the source and the target domains are summarized in Table 4.6.

<table>
<thead>
<tr>
<th>Model</th>
<th>P</th>
<th>R</th>
<th>F1</th>
<th>Domain</th>
</tr>
</thead>
<tbody>
<tr>
<td>HMM(BILOU)</td>
<td>82.49</td>
<td>81.16</td>
<td>81.82</td>
<td>source</td>
</tr>
<tr>
<td>HMM(BILOU)</td>
<td>53.29</td>
<td>57.52</td>
<td>55.33</td>
<td>target</td>
</tr>
<tr>
<td>MEMM(BIO)</td>
<td>84.68</td>
<td>81.54</td>
<td>83.08</td>
<td>source</td>
</tr>
<tr>
<td>MEMM(BIO)</td>
<td>70.02</td>
<td>61.86</td>
<td>65.69</td>
<td>target</td>
</tr>
</tbody>
</table>

Table 4.6: Performance of source models over the 3 name types.

Table 4.6 shows that although both models achieve F1 of 80s on the source domain, they generalize poorly on the target domain. Comparing the HMM model with the MEMM model, it seems that the feature based model generalizes better to the target domain than the generative HMM model even though we did use...
the word type features as a back-off model similar to [5]. We did not observe the advantage of using the BILOU scheme as reported in [65] for both the source and the target domains. Although we could not determine the exact reason why this happens for the source domain, for the target domain, it contains long transliterated Arabic names implying that the state transition from “I” to “I” is more common in the target than the source domain. This type of transition might not be observed enough and estimated sufficiently by the fine grained BILOU scheme.

4.5.3 Performance of the Generalized Model

We augmented the source model with word clusters (as described in Section 4.4.2) from the four unlabeled corpora in Table 4.4. Their performance on the target domain is shown in Table 4.7.

<table>
<thead>
<tr>
<th>Word clusters</th>
<th>P</th>
<th>R</th>
<th>F1</th>
</tr>
</thead>
<tbody>
<tr>
<td>No Cluster</td>
<td>70.02</td>
<td>61.86</td>
<td>65.69</td>
</tr>
<tr>
<td>Reuters 1996</td>
<td>69.26</td>
<td>64.26</td>
<td>66.67</td>
</tr>
<tr>
<td>RCV1</td>
<td>66.33</td>
<td>64.42</td>
<td>65.36</td>
</tr>
<tr>
<td>TDT5</td>
<td>70.76</td>
<td>66.51</td>
<td>68.57</td>
</tr>
<tr>
<td>Joint Corpus</td>
<td>72.82</td>
<td>66.61</td>
<td>69.58</td>
</tr>
</tbody>
</table>

Table 4.7: Performance of the generalized source model.

Table 4.7 shows the superiority of using a joint corpus to generate word clusters: the 10M words joint corpus outperformed the other 3 larger corpora. The TDT5 corpus is more than 8 times larger than the joint corpus, but is still 1 point behind. Using word clusters from the Reuters corpora (Reuters 1996 and RCV1) have shown to improve NER systems’ performance on the CoNLL 2003 NER task[65][80]. But they provided limited performance gain for our model when testing on the target domain. The results shown here indicate the necessity
of using a joint corpus or ideally a general purpose corpus for generalizing the source domain model for cross-domain NER.

### 4.5.4 Performance of CDB

We start with the generalized source/seed model and run the CDB algorithm on the unlabeled target domain corpus using the parameters specified in Table 4.5. As mentioned earlier, the seed model is generalized sequentially, that is, word cluster features are used during each round. We plot $F_1$ score against iteration in Figure 4.2. The results are obtained by testing the updated model during each round on the labeled target domain data. The results are averaged on 10 runs.

There are several clear trends in Figure 4.2. First, without using the novelty measure (the line at the bottom), CDB performs worse than generalized seed model (GSM). Although the seed model is already generalized with word clusters, the most confidently labeled instances might still be more similar to the source than the target domain. This indicates that novelty is a necessary measure for cross-domain NER. Comparing the two confidence measures: $ComConf$ and $LocalConf$, in general, $ComConf$ outperforms $LocalConf$. After using the novelty measure, all instances are new to our seed model. So there is some degree of uncertainty when the model tries to make a local prediction. Not only considering the local prediction, but also considering how the same token is labeled globally, the $ComConf$ measure seems to be a better choice in a cross-domain setting.

Regarding the density and the diversity measures, both of them further improve the performance. $Density$, however, does not perform well in the first 6 iterations. We checked the instances that had been selected during these iterations and found that many of them appear with very strong context words such as $Mr.$, $President$, $80$
Figure 4.2: Performance of CDB. GSM stands for generalized seed model at iteration 0; + means with and - means without.

*General and said. They are considered representative instances according to our density measure. They can be regarded as cross-domain contexts which might have been learned by the generalized and un-generalized source domain models. In contrast, the diversity measure not only considers how representative an instance is but also prefers a certain degree of difference among the selected instances. Hence, the diversity measure has achieved the best result CDB could get so far.

The best $F$ score with the diversity measure is 72.85, a 7.16 improvement compared to the source model. The $F$ score at the last iteration is 72.45, a 6.76 improvement compared to the source model.
We also run a standard bootstrapping procedure with the un-generalized seed model and with the same parameters used for the CDB procedure. The performance trends of using different instance selection criteria are similar to those of the CDB algorithm. The best F score, 70.12, is also obtained with the diversity measure. This further confirms that the multiple criteria proposed in this paper are better than a single criterion. CDB with generalized seed model outperformed the standard bootstrapping by more than 2 points which further indicates the usefulness of the combination of feature generalization and multi-criteria-based instance selection methods proposed in this paper.

4.6 Conclusion

We have described a general cross-domain bootstrapping algorithm for adapting a model trained only on a source domain to a target domain. We have improved the source model’s F score by around 7 points. This is achieved without using any annotated data from the target domain and without explicitly encoding any target-domain-specific knowledge into our system. The improvement is largely due to the incorporation of the global evidence in feature generalization and the multi-criteria-based instance selection method.

Our immediate future work is to find a natural stopping criterion for the bootstrapping procedure, perhaps through the detection of semantic drift.[57][75]. Gazetteer resources have proven to be a powerful knowledge base for improving NER performance[22]. The only gazetteer in CDB now is a country and US state list. So another promising research avenue is to study how to automatically learn or mine a target domain named entity dictionary to further improve our system’s
performance.
Chapter 5

Conclusion and Future Work

5.1 Conclusion

Recent advances in information extraction move beyond narrowly restricted domains and shift the focus to a variety of new emerging domains. These include terrorism reports, scientific publications, legal and medical documents, microblog texts such as tweets and Foursquare tips, collaboratively generated web documents such as Wikipedia, and many more. This however, brings a great many new tasks and challenges, and it is not desirable to hire human experts to annotate data for every new task of every new domain.

This dissertation moves a step further to solve these emerging problems with low cost and in a fast and accurate way. Specifically, we have presented three systems: the relation type extension system based on active learning (Chapter 2), the relation type extension system based on semi-supervised learning (Chapter 3), and the cross-domain bootstrapping system for domain adaptive named entity extraction (Chapter 4).
The main contribution of this dissertation is its exploration of using both local and global evidence for fast system development. The benefits of using these two types of evidence are substantial: the co-testing procedure based on the local and global views of relation instances reduced annotation cost by 97% while maintaining the performance level of supervised learning. The use of pattern clusters as the global view in semi-supervised relation pattern acquisition not only greatly improved the quality of learned patterns but also contributed to a natural stopping criterion. The generalization of local lexical features using word clusters and the instance selection based on global evidence in the cross-domain bootstrapping procedure were able to improve the source model’s performance on the target domain by 7% F1 without annotating any target domain data.

5.2 Future Work

Active Learning for Relation Type Extension. For investigating the feasibility of the proposed active learning approach, this dissertation performed simulated active learning experiments with the ACE 2004 annotated corpora. This simplifies the evaluation of active learning from many factors that might arise in real world active learning settings: the effect of downstream component errors such as name and entity extraction, the effect of annotation consistency by one or more annotators, the time of model re-training in each iteration, etc. We are excited to explore these factors in our future real world active learning experiments.

Another direction of this branch of research, as mentioned in the last section of Chapter 2, is to combine semi-supervised learning with active learning to further reduce annotation cost.
Semi-supervised Learning for Relation Type Extension. Currently the seed patterns used are defined manually, which is far from being a desirable and principled way. A promising research avenue is to study the automatic discovery and selection of seed patterns. More specifically, we will focus on how to conduct seed discovery and selection based on the *global* evidence, i.e., pattern clusters. A two-stage procedure might move this idea forward: manually define some *initial* seed patterns and then choose the *good* ones from the *target* cluster as the *actual* seeds.

Cross-domain Bootstrapping for Named Entity Recognition. In domain adaptation for named entity recognition, one may have a dictionary of names for the *target* domain but no labeled *target* domain data. This represents a more and more realistic scenario because of the emergence of collaboratively generated knowledge bases such as Wikipedia. There are two future directions for our cross-domain bootstrapping to accommodate this scenario: 1) extracting dictionary-based features to further generalize features; 2) combining with distantly annotated data (by aligning a dictionary with texts to generate training data) to further improve performance.

Other Future Research Avenues. As the ultimate goal of information extraction is to power other applications, I see the opportunity to expand the scope of my research to contribute to other fields of study. For example, the information retrieval community shows great interest in entity-oriented search such as people search, job search and product search. Extracting the entities and the attributes of them (similar to relation extraction) with high performance is a first and crucial step for finding relevant documents for the entities. It would be exciting to bridge the gap between disciplines.
Bibliography


