Approximate String Matching Techniques

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Keywords: String Matching, Data Quality, Record Matching, Record Linkage, Data Warehousing.

Abstract: Data quality is a key to success for all kinds of businesses that have information applications involved, such as data integration for data warehouses, text and web mining, information retrieval, search engine for web applications, etc. In such applications, matching strings is one of the popular tasks. There are a number of approximate string matching techniques available. However, there is still a problem that remains unanswered: for a given dataset, how to select an appropriate technique and a threshold value required by this technique for the purpose of string matching. To challenge this problem, this paper analyses and evaluates a set of popular token-based string matching techniques on several carefully designed different datasets. A thorough experimental comparison confirms the statement that there is no clear overall best technique. However, some techniques do perform significantly better in some cases. Some suggestions have been presented, which can be used as guidance for researchers and practitioners to select an appropriate string matching technique and a corresponding threshold value for a given dataset.

1 INTRODUCTION

Data quality is a key to success for all kinds of businesses that have information applications involved, such as data integration for data warehouse applications; text and web mining, information retrieval and search engines for web applications. When data need to be integrated from multiple sources, it always has a problem: how to identify data records that refer to equivalent entities, which is called a duplicate detection problem. The “approximate string matching problem” is generally used as a black-box function with some threshold parameter in the context of duplicate detection problem. In general, string matching deals with the problem of whether two strings refer to the same entity.

String matching, also referred as record matching or record linkage in statistics community has been a persistent and well-known problem for decades (Elmagarmid et al., 2007). For a brief review of the topic of record linkage, see the work by Herzog et al., (2010). To deal with string matching, there are mainly two types of matching techniques: character-based and token-based techniques. Character-based similarity techniques are designed to handle well typographical errors. However, it is often the case that typographical conventions lead to rearrangement of words e.g., “John Smith” vs. “Smith, John”. In such cases, character-based techniques fail to capture the similarity of the entities. Token-based techniques are designed to compensate for this problem (Elmagarmid et al., 2007). Therefore, generally speaking, character-based similarity techniques are good for the single word problem, such as name matching while token-based for the matching with more than one word. For instance, for e-commerce datasets used in (Köpcke et al., 2010), token-based techniques could be more appropriate, given the length of the fields. This paper will focus on popular token-based techniques. Although these techniques are designed for dealing with this kind of problems, since there is no clear overall best string matching technique for all kinds of datasets (Christen, 2006), a question still remains unanswered for researchers and practitioners: for a given dataset, how to select an appropriate technique and a threshold value required by this technique for the purpose of string matching. In past decade, several researchers have challenged this problem (Bilenko et al, 2003; Christen, 2006; Cohen, Ravikumar and Fienberg, 2003; Hassanzadeh et al., 2007; Peng et al., 2012). However, none of them have done such a
comprehensive analysis and comparison work that is done for token-based techniques in this paper. The contributions of this paper are to overview 12 popular token-based string matching techniques, evaluate whether the following factors will have effect on the performance: the error rate in a dataset, the threshold value, the selected type of strings in a dataset, and the size of a dataset, by using sixty three carefully designed datasets.

The rest of this paper is structured as follows. Related works are described in next section. Section 3 introduces techniques that will be examined. The main contribution of this paper is presented in section 4 that describes the preparation of the datasets, the experiments, the analysis and comparisons. Finally, this paper is concluded and future work pointed out in section 5.

2 RELATED WORK

Bilenko et al. (2003) and Cohen et al. (2003) evaluated and compared a set of existing string matching techniques, which include popular character-based techniques, token-based techniques and hybrid techniques. They claimed that the Monge-Elkan performed best on average and SoftTFIDF performed best overall. However, their works did not consider the effect of the error rate, the type of errors in a string, the token length in a string and the size of a dataset on the performance. Besides, regarding the threshold value used for matching, their work only mentioned that a suitable threshold value was chosen, but not mentioned how and whether or not this value was universal for all considered techniques.

Peter Christen (2006) thoroughly discussed the characteristics of personal names and the potential sources of variations and errors in them, and also evaluated a number of commonly used name matching techniques, considering given names, surnames and full names separately, and proposed nine useful recommendations for technique selection when dealing with name matching problems. Particularly, the author pointed out the importance of choosing a suitable threshold value. It was argued that it was a difficult task to select a proper threshold value and even small changes of the threshold could result in dramatic drops in matching quality. However, his focus was on personal name matching techniques. Also, similar to Cohen et al’s work (2003), the author did not consider any effect of the error rate, the token length in a string and the size of a dataset on the performance.

Hassanzadeh et al., (2007) presented an overview of several string matching techniques and thoroughly evaluated their accuracy on several datasets with different characteristics and common quality problems. Similar to this paper, the work was focused on token-based string matching techniques. The effect of types of errors and the amount of errors were both considered. Types of errors considered include edit errors, token swap and abbreviation replacement. Their experiment results showed that types of errors and the amount of errors both had significant effect on the performance. It was claimed that the threshold value used for the matching task would influence the individual performance of matching techniques. However the token length in a string and the size of datasets were not considered.

Recently, Peng et al., (2012) presented an evaluation work on techniques for name matching. The work considered a variety of factors, such as the error rate, the size of a dataset, which might have effect on the performance of such techniques. Their preliminary experimental results confirmed that there is no overall clear best technique. The work claimed that the error rate in the dataset has effect on threshold values. However, they only considered character-based matching techniques.

3 STRING MATCHING TECHNIQUES

String matching that allows errors is also called approximate string matching. The problem, in general, is to “find a text where a text given pattern occurs, allowing a limited number of “errors” in the matches.” (Navarro, 2001) Each technique uses a different error model, which defines how different two strings are.

In token-based similarities, two strings, s and t can be converted into two token sets, where each token is a word. A similarity function, Sim() is used to define whether strings s and t are similar or not, based on a given threshold value. In this paper, 12 token-based similarity techniques are considered. In the rest of this section, if there is no reference given, the description of algorithm’s formula follows the work in (Hassanzadeh et al., 2007).

3.1 Notations

Given two relations: $R = \{r_i : 1 \leq i \leq N_1\}$ and $S = \{s_j : 1 \leq j \leq N_2\}$, where $|R| = N_1$, $|S| = N_2$, for a similarity
function \( \text{sim} \), a pair of records, \((r_x, s_y) \in R \times S\) is considered to be similar if \( \text{sim}(r_x, s_y) \geq \theta \), where \( \theta \) is a given threshold value.

### 3.2 Edit Similarity or Levenshtein

The Levenshtein distance (Navarro, 2001) is defined to be the minimum number of edit operations required to transform string \( s_1 \) into \( s_2 \). Edit operations are delete, insert, substitute and copy. It can be calculated by:

\[
\text{Sim}_{\text{Leven}}(s_1, s_2) = 1.0 - \frac{\text{dist}(s_1, s_2)}{\max(|s_1|, |s_2|)}
\]

where \( \text{dist}(s_1, s_2) \) refers to the actual Levenshtein distance function which returns a value of 0 if the strings are the same or a positive number of edits if they are different. The value of such a measure is between 0.0 and 1.0 where the bigger the value, the more similar between the two strings.

### 3.3 GES

The Gap Edit Similarity (GES) measure uses gap penalties. (Chaudhuri et al., 2006). GES defines the similarity between two strings as a minimum cost required to transform string \( s_1 \) into \( s_2 \). The similarity measure for two strings, \( s_1, s_2 \) can be calculated by:

\[
\text{Sim}_{\text{GES}}(s_1, s_2) = 1 - \min(\frac{tc(s_1, s_2)}{wt(s_1)}, 1.0)
\]

where \( wt(s_1) \) is the sum of weights of all tokens in \( s_1 \), and \( tc(s_1, s_2) \) is a sequence of the following transformation operations:

- **Token insertion**: inserting a token \( t \) in \( s_1 \) with cost \( w(t) \), \( c_m \) where \( c_m \) is the insertion factor constant and is in the range between 0 and 1. In our experiments, \( c_m = 1 \);
- **Token deletion**: deleting a token \( t \) from \( s_1 \) with cost \( w(t) \);
- **Token replacement**: replacing a token \( t_1 \) by \( t_2 \) in \( s_1 \) with cost \( (1 - \text{sim}_{\text{Leven}}(t_1, t_2)) \cdot w(t) \).

### 3.4 Jaccard/Weighted Jaccard

Jaccard similarity is the fraction of tokens in \( s_1 \) and \( s_2 \) that are present in both of the strings. Weighted Jaccard is the weighted version of Jaccard similarity. The similarity measure for two strings, \( s_1, s_2 \) can be calculated by:

\[
\text{Sim}_{\text{wJaccard}}(s_1, s_2) = \frac{\sum_{t \in s_1 \cap s_2} w_R(t)}{\sum_{t \in s_1 \cup s_2} w_R(t)}
\]

where \( w_R(t) \) is a function of weight that reflects the frequency of token \( t \) in relation \( R \):

\[
w_R(t) = \log\left(\frac{n - n_t + 0.5}{n_t + 0.5}\right)
\]

where \( n \) is the number of tuples in the relation \( R \) and \( n_t \) is the number of tuples in \( R \) containing the token \( t \).

### 3.5 TF-IDF

TF-IDF, term frequency–inverse document frequency, is a similarity measure that is widely used in information retrieval community (Cohen, Ravikumar and Fienberg, 2003). The similarity measure for two strings, \( s_1, s_2 \) can be calculated by:

\[
\text{Sim}_{\text{TFIDF}}(s_1, s_2) = \sum_{w \in s_1 \cap s_2} V(w, s_1) \cdot V(w, s_2)
\]

\[
V'(w, s) = \text{log}(\text{TF}_{w, s} + 1) \cdot \text{log}(\text{IDF}_w)
\]

\[
V(w, s) = \frac{V'(w, s)}{\sum_{w'} V'(w, s')^2}
\]

where \( \text{TF}_{w, s} \) is the frequency of token \( w \) in \( s \), \( \text{IDF}_w \) is the inverse of the fraction of names in the corpus that contains \( w \).

### 3.6 SoftTFIDF

SoftTFIDF is a hybrid similarity measure by Cohen et al., (2003), in which similar tokens are considered as well as tokens in \( s_1 \) \( \cap \) \( s_2 \). The similarity measure for two strings, \( s_1, s_2 \) can be calculated by:

\[
\text{Sim}_{\text{SoftTFIDF}}(s_1, s_2) = \sum_{w \in \text{CLOSE}(\theta, s_1, s_2)} V(w, s_1) \cdot V(w, s_2) \cdot D(w, s_2)
\]

where \( \text{CLOSE}(\theta, s_1, s_2) \) is the set of tokens \( t \) such that for some \( t_i \in \text{CLOSE}(\theta, s_1, s_2) \), \( \text{sim}(t_i, s_2) > \theta \), where \( \text{sim}(\cdot) \) is a secondary similarity function, and \( D(w, s_2) = \max_{t \in \text{CLOSE}(\theta, s_1, s_2)} \text{sim}(w, s_2) \), for \( w \in \text{CLOSE}(\theta, s_1, s_2) \). In the experiments, Jaro-Winkler similarity function is used as the secondary similarity function and \( \theta = 0.9 \).

### 3.7 Cosine TF-IDF

Cosine TF-IDF is a well-established technique used in Information Retrieval and uses vectors to analyse the strings and to calculate the similarity between the pattern and the string. The similarity is determined by the cosine of the angle between these vectors. The formula is shown below for the
similarity, as well as the formula for how the vectors are assigned:

\[ \text{Sim}_{\text{cosine}}(s_1, s_2) = \sum_{t \in s_1 \cap s_2} w_s(t) \cdot w_{s_2}(t) \]

where \( w_s(t) \) and \( w_{s_2}(t) \) are normalized tf-idf weights for each common token in \( s_1 \) and \( s_2 \) respectively. The normalized tf-idf weight of token \( t \) in a given string \( s \) is defined as follows:

\[ w_s(t) = \frac{w'_s(t)}{\sqrt{\sum_{t' \in s} w'_s(t')^2}} \]

where \( w'_s(t) \) is the term frequency of token \( t \) within string \( s \) and \( \text{idf}(t) \) is the inverse document frequency with respect to the entire relation \( R \).

### 3.8 BM25

BM25 is an efficient string searching technique that is used as the foundation for many others. The algorithm looks for instances of P (pattern) in S (text) by performing comparisons of characters at a range of different alignments. The similarity measure for two strings, \( s_1, s_2 \), can be calculated by:

\[ \text{Sim}_{\text{BM25}}(s_1, s_2) = \sum_{t \in s_1 \cap s_2} w'_{s_1}(t) \cdot w_{s_2}(t) \]

where

\[ w'_{s_1}(t) = \frac{(k_2 + 1) \cdot tf_{s_1}(t)}{k_2 + tf_{s_1}(t)} \]

\[ w_{s_2}(t) = w_R^{(1)}(t) \cdot \frac{(k_1 + 1) \cdot tf_{s_2}(t)}{k_2 + tf_{s_2}(t)} \]

and

\[ w_R^{(1)}(t) = \log \left( \frac{n - n_t + 0.5}{n_t + 0.5} \right) \]

\[ K(s) = k_1 (1 - b) + b \cdot \frac{|s|}{\text{avg}_s} \]

where \( tf_s(t) \) is the frequency of the token \( t \) in string \( s \), \( |s| \) is the number of tokens in \( s \), \( \text{avg}_s \) is the average number of tokens per record in relation \( R \), \( n_t \) is the number of records containing the token \( t \) and \( k_t \), \( k_s \) and \( b \) are a set of independent parameters, where \( k_t \in [1,2] \), \( k_s = 8 \) and \( b \in [0.6, 0.75] \).

### 3.9 Hidden Markov

Hidden Markov Model (HMM) has a formula wherein a language model is given to each individual document. HMM is a model that is assigned to actual strings whereby the probability of returning a particular string is measured against another string. The following formula shows the score for HMM used in this experiment:

\[ \text{Sim}_{\text{HMM}}(s_1, s_2) = \prod_{t \in s_1} (a_0 P(t|GE) + a_1 P(t|s_2)) \]

where \( a_0 \) and \( a_1 = 1 - a_0 \) are the transition states probabilities of the Markov model and \( P(t|GE) \) and \( P(t|s_2) \) are determined by:

\[ P(t|GE) = \frac{\sum_{s \in R} \text{number of times } t \text{ appears in } s}{\sum_{s \in R} |s|} \]

\[ P(t|s_2) = \frac{\text{number of times } t \text{ appears in } s}{|s_2|} \]

### 3.10 WHIRL

The similarity of two document vectors \( \vec{v} \) and \( \vec{w} \) is usually interpreted as the cosine of the angle between \( \vec{v} \) and \( \vec{w} \). The value of \( \text{Sim}_{\text{WHIRL}}(\vec{v}, \vec{w}) \) is always between one and zero and is given by the following formula (Cohen, 2000):

\[ \text{Sim}_{\text{WHIRL}}(\vec{v}, \vec{w}) = \frac{\vec{v} \cdot \vec{w}}{||\vec{v}|| \cdot ||\vec{w}||} \]

where \( \vec{v} \) is related to the “importance” of the term \( t \) in the document represented by \( \vec{v} \), and \( \vec{w} \) is related to the relevance of term \( t \) in the document represented by \( \vec{w} \). \( ||\vec{v}|| \) is the length of the vector \( \vec{v} \) and \( ||\vec{w}|| \) is the length of the vector \( \vec{w} \). Two documents are similar when they share many “important” terms (Cohen, 2000).

### 3.11 Affine Gap

Some sequences are much more likely to have a big gap, rather than many small gaps. For example, a biological sequence is much more likely to have one big gap of length 10, due to a single insertion or deletion event, than it is to have 10 small gaps of length 1. Affine gap penalties use a gap opening penalty, \( o \), and a gap extension penalty, \( e \). A gap of length \( l \) is then given a penalty \( o + (l - 1)e \). So that gaps are discouraged, \( o \) is almost always negative. Because a few large gaps are better than many small gaps, \( e \), though negative, is almost always less negative than \( o \), so as to encourage gap extension, rather than gap introduction.

In the Affine Gap Penalty model, a gap is given a weight \( W_g \) to “open gap” and another weight \( W_s \) to “extend the gap”. The formula for the model is:

\[ W_{\text{total}} = W_g + q W_s \]
where \( q \) is the length of the gap, \( W_g \) is the weight to “open the gap” and \( W_s \) is the weight to “extend the gap” with one more space (Vingron and Waterman, 1994).

### 3.12 Fellegi-Sunter

Fellegi-Sunter makes comparison between recorded characteristics and values in two records and makes a decision based on whether or not the comparison pair make up the same object or event (Fellegi and Sunter, 1969).

Decisions are made according to three types which will be referred to as \( A_1 \) (a link), \( A_2 \) (a possible link) and \( A_3 \) (a non-link). When members of the comparison pair are unmatched then Fellegi-sunter equation calculates the score by determining the probability of the error occurring between the matched pair defined as:

\[
\mu = \sum_{\gamma \in \Omega} u(\gamma) P(A_1|\gamma)
\]

\[
\lambda = \sum_{\gamma \in \Omega} m(\gamma) P(A_3|\gamma)
\]

where \( u(\gamma) \) and \( m(\gamma) \) are probabilities of realizing and discovering a comparison vector. The summation is the total space \( P \) of all possible realizations.

### 4 EXPERIMENTS AND EVALUATION

In our experiments, we focus on the performance of the above 12 popular token-based string matching techniques. Datasets designed cover a wide range of characteristics, such as the level of “dirtiness”, the token length in a string, the size of datasets and the type of errors. It is expected that the experiment results should show that all these characteristics have significant effect on the performance and optimal threshold values.

#### 4.1 Datasets Preparation

In the absence of common datasets for data cleaning, we prepare our data for experiments as follows.

The datasets that are used are based on real Electoral Roll data. First, a one million record dataset was extracted, from which an address list was created, which includes House number, Street, City and County, Postcode. This list contains 10000 clean, non-duplicate addresses, with an ID associated to each of the records.

Erroneous records were introduced by doing the following five operations manually to the address fields of records: inserting, deleting, substituting, replacing characters and swapping tokens. The replacing type of errors includes abbreviation errors, such as replacing St. with Street or vice versa. The level of the dirtiness of a dataset is divided into three levels: Low, Medium and High. Static percentages are used as a guideline for such a classification: for low error datasets, the percentage was set to 20%; for medium, 50% and for high, 80%.

The dataset sizes that were considered consisted of six sizes with 500, 1000, 2000, 4000, 8000 and 10000 records respectively. The average record length was approximately 8.2 words per record. The errors in the datasets were distributed in a uniform way. Experiments have also been run on datasets generated using different parameters. For each size, there were three datasets generated with mixed types of errors, having a different error rate associated. This contributes eighteen datasets with mixed type of errors introduced.

In addition, datasets with single type of errors were also considered in our experiment. The insertion and deletion errors were incorporated solely into each dataset and the algorithms were measured against each other to see how they performed against each technique. With regard to how these error types were distributed amongst the data, the distribution went as follows: Insertion errors were performed on a single character per record that would correspond to a given percentage distribution rate dependent on the error rate being analyzed. For example, for the low error dataset for 1000 records, there were 20% of those records that had a single “insertion” error implemented. So 200 of those 1000 records had a character “added” that would simulate an error. There were 18 datasets incorporated with only insertion errors.

Deletion followed a similar approach for each error rate and took a single character away from the number of records corresponding to the respective percentage error rate. There were 18 such datasets as well.

Token lengths were measured for the 8000 size dataset with the “low”, “medium” and “high” error rates taken into consideration. Three different token lengths were used, “Short”, “Medium” and “Long”. For the short tokens, the only fields considered for each record were for Street and City. Medium length considered County and the Long length size added PostCode which was the original size of the record.
The average length used for low was 2.1 tokens per record, medium was 5.6 and long was 8.2. Therefore in total there are sixty three datasets for the experiment.

4.2 Measures

A target string is a positive if it is returned by a technique; otherwise it is a negative. A positive is a true positive if the match does in fact denote the same entity; otherwise it is a false positive. A negative is a false negative if the un-match does in fact denote the same entity; otherwise it is a true negative.

We evaluate the matching quality using the F-measure ($F$) that is based on precision and recall:

$$F1 = \frac{2 \times P + R}{P + R}$$

with $P$ (precision) and $R$ (recall) defined as:

$$P = \frac{|true positives|}{(|true positives| + |false positives|)}$$

$$R = \frac{|true positives|}{(|true positives| + |false negatives|)}$$

Clearly, a trade-off between recall and precision exists, if most targets are matched, recall will be high but precision will be low. Conversely if precision is high, recall will be low. F1-measure is a way of combining the recall and precision into a single measure of overall performance (Rijsbergen, 1979). In our experiments, precision, recall and F1-measure are measured against different value of similarity thresholds, $\theta$. For the comparison of different techniques, the maximum F1-measure score across different thresholds is used.

4.3 Results

In this section, testing results on the sixty three carefully designed datasets are analysed and evaluated. Results show that in general, the size of a dataset is not significantly sensitive to the accuracy (the best F1-score) relative to the threshold values.

4.3.1 Effect of Error Rate on Threshold Values

Figure 1: Optimum threshold value for different techniques on datasets with three different error rates.

Figure 2: Maximum F1-score for different techniques on datasets of 8000 records with three different error rates.

Figure 3: Maximum F1-score for different techniques on datasets of 8000 records with three different token length associated with medium error rates.

Figure 4: Maximum F1-score for different techniques on datasets of 10000 records only having insertion errors with three different error rates.
rates. The results confirm that the level of “dirtiness” in a dataset has significant effect on threshold selection. From the figure, it says the higher the error rate in the dataset, the lower the threshold value is required in order to achieve the maximum F1-score, except the case for BM25. For example W/Jaccard achieves the maximum F1-score on datasets with low error rate at the threshold value of 0.85, with medium and high error rate at threshold values of 0.57 and 0.47 respectively. However, BM25 achieves the maximum F1-score on datasets with high error rate at the threshold value of 0.60, but with medium error rate at the threshold value of 0.59.

4.3.2 Effect of Error Rate on Performance

Experiment results show that the “dirtiness” of a dataset has great effect on the overall performance. As an example, Figure 2 shows the maximum F1-scores for the 12 different techniques on datasets of size 8000 with three different error rate associated. It shows that for datasets with low error rate associated, HMM, BM25 and Cosine TF-IDF are the best three performers among the 12 algorithms, while Affine Gap, WHIRL and SoftTFIDF are the worst performers. For datasets with medium and high error rate associated, Affine Gap joined HMM and BM25 as the top three performers, while Cosine TF-IDF drops out of the best 5. This shows that Affine Gap is suitable for datasets with higher level of “dirtiness”, and Cosine TF-IDF is suitable for datasets with lower level of “dirtiness”. In general, the performance decreases along with the increase of the “dirtiness” in a dataset, except technique, Affine Gap, which has higher maximum F1-score on datasets with medium error rate associated than on datasets with low error rate associated.

4.3.3 Effect of Size of Datasets on Performance

Our experiment results show that the size of datasets does not have significant effect on maximum F1-score values, given optimum threshold values.

4.3.4 Effect of Length of Strings on Threshold Values

Figure 3 shows the maximum F1-score for different techniques on datasets of 8000 records with three different token lengths associated with medium error rates. It says in general, performance decreases along with the increase of lengths, except Cosine TF-IDF, where the performance on long length datasets is slightly better than that on medium length. From results for different techniques on datasets with medium length and long length tokens respectively, associated with three different error rate, it can be seen that BM25, HMM, cosine TF-IDF and W/Jaccard performed better than others, and Affine Gap, WHIRL and SoftTFIDF performed badly when the token length is long. However, the performance of Affine Gap, WHIRL and SoftTFIDF increased significantly when the token length is medium.

4.3.5 Effect of Type of Errors on Performance

Figure 4 shows the maximum F-score for different techniques on datasets of 10000 records only having insertion errors with three different error rates. For datasets with low error rate associated, W/Jaccard, GES and Edit Similarity are the best three performers, while Fellegi-Sunter, Cosine TF-IDF and BM25 perform worse than the rest. The performance of BM25 increases along with the increase of the level of “dirtiness” in datasets, while the performance of GES and Edit Similarity decreases along with the increase of the level of “dirtiness” in datasets. It is noted that BM25 is among the best three performers in most of cases when the error types in datasets are mixed. See section 4.3.2.

4.3.6 Effect of Timing

Results also show, when the error rate is low, Affine, GES, Felligi-Sunter and Edit Similarity cost less time among the twelve algorithms whereas SoftTFIDF costs the most time. The pattern is similar when the error rate is medium. However, SoftTFIDF costs the least time when the error rate is high. HMM, TF-IDF and WHIRL also cost less time than the rest when the error rate is high. Our experiment results agree that smaller datasets cost less time, and the time used increases when the error rate increases. In particular, time used on datasets with high error rate associated is significantly more than that on datasets with low or medium error rate associated.

4 CONCLUSIONS AND FUTURE WORK

This paper has analysed and evaluated twelve popular token-based name string matching
techniques. A comprehensive comparison of the twelve techniques has been done based on a series of experiments on 63 carefully designed datasets with different characteristics, such as the rate of errors, the type of error, the number, the length of tokens in a string, and the size of a dataset. The comparison results confirmed the statement that there is no clear best technique. The characteristics considered all have significant effect on performance of these techniques, except the size of a dataset. In general, HMM and BM25 perform better than others, especially on smaller sized datasets, but consume much more time. Cosine TF-IDF and TF-IDF are better on larger datasets with a higher error rate associated. Results also show that techniques that perform well on datasets incorporated with mixed type of errors do not secure a similar performance on datasets incorporated with a single type of errors. For example, BM25 didn’t perform well on datasets with low error rate, incorporated only with insertion errors. Similarly, HMM didn’t perform well on datasets with low error rate, incorporated only with deletion errors. The token length also has an effect on the performance. For example, some techniques, such as Affine Gap, WHIRL and SoftTFIDF performed much better when the token length is medium than that of the token length when it is long.

Regarding the threshold value, the results show that the level of “dirtiness” in a dataset has significant effect on threshold selection. In general, the higher the error rate in the dataset, the lower the threshold value is required in order to achieve the maximum F1-score.

The work introduces a number of further investigations, including: 1) to do more experiments on datasets with more characteristics, such as the number of tokens in strings etc.; 2) to do further analysis in order to evaluate whether there is a method to select a threshold value for any of the matching techniques on a given dataset.

REFERENCES


