

Manual of Disambiguation

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1 Overview

The source code is an implementation of authorship disambiguation algorithm described by Vetle Torvik et al 2009. While it is originally designed for the disambiguation of inventors in the United States Patent and Trademark Office (USPTO) database, minor modifications in the source file can provide solutions for problems of the similar kind.

The algorithm uses Bayesian theorem to calculate the probability of match between two records based on the comparison between the two records in significant attributes such as names, geographical location and companies, etc. The algorithm includes three major steps: 1. Blocking, 2. Training, and 3. Disambiguation. The details of each step are described in the following sections.

The source code is written in C++ and uses many features of C++, such as polymorphism, template, STL library, etc. Therefore, a fairly good understanding of C++ is expected to fully understand the code. The code also outsources IBM CPLEX for quadratic programming. Import and export of the disambiguation result may need some APIs of SQL/SQLite. The code is compiled in Linux by GNU gcc/g++/make.

2 Work Flow

2.1 Schematic

This is a schematic of the whole disambiguation process. Each array represents one step, with the square box of identical color representing the source code files that implements the step, and the round box of identical color representing the necessary definitions of relevant classes and/or functions.

The disambiguation is an iterative process: the result of previous disambiguation serves as the input of the next round disambiguation. For each round of disambiguation, there are three major steps: blocking, training and disambiguation.

2.2 Blocking

Blocking of records is the first step of disambiguation. The whole point of blocking is to avoid most unnecessary comparisons without much accuracy loss. Records of similar attributes are blocked together since they are likely to be a match. Because the aim of disambiguation is individual inventors, the most significant attributes names are usually adopted as the blocking rules. It is critical to remember that ONLY records within the same blocks are compared and thus disambiguated.

Iterative blocking in a permissive way is applied in the disambiguation, because it is assumed that after each round of disambiguation, blocks are well disambiguated and thus similar and/or more stringent blockings are meaningless. For example, the first round of blocking is exact first names + exact last names, the second can be first 5 characters of first names + first 8 characters of last names, and the third round can be first 3 characters of first names + first 5 characters of last names.

2.3 Training

The point of training is to obtain the ratios database for similarity profiles based on training sets.

2.3.1 Similarity Profiles

Similarity profiles are defined as multi-dimensional variables that describe the similarity of two records. Each record has many attributes, such as first name, last name, assignee, street, city, country, etc. Similarity profiles are selected attributes that are believed to be influential in pairwise comparisons; thus, they are decided by users and can be different in each round of disambiguation. For example, similarity profiles in the first round of disambiguation can be first name + last name + assignee + city, while those in the second round can be last name + assignee + street. Each attribute in a similarity profile is given a score to evaluate the similarity of the attribute from the comparison of the two records. The score is discrete and is usually non-negative integers. Scoring of each attribute is totally user-defined, and, conceptually, has to be fully understood by users. Similarity profiles refer to the multi-dimensional integers obtained from the comparison.

2.3.2 Training sets

Training sets contain pairs of records that are believed with high confidence to be either match or non-match. In the USPTO patent inventor disambiguation, there are four training sets:

1. match pairs to train personal information based on patent information: this set is created by selecting record pairs belonging to the same blocks and having at least two common coauthors.
2. non-match pairs to train personal information based on patent information: this set is created by selecting record pairs that share the same patent id but different author sequence of that patent.
3. match pairs to train patent information based on personal information: this set is created by selecting record pairs that share exact rare names.
4. non-match pairs to train patent information based on personal information: this set is created by selecting record pairs whose names are both rare and different.

2.3.3 Ratios database

The ratio value of a given similarity profile, or R-value, is defined as the frequency of the similarity profile in the match set divided by the frequency of the same similarity profile in the non-match set. Ratios database, therefore, is the mapping of all possible similarity profiles to their corresponding r-values. In the patent disambiguation, personal information is assumed to be independent from patent information, so r-value of a similarity profile is the multiple of the r-value of the personal information part of the similarity profile and that of the patent information part of it. By using the four training sets described above, the ratios database can be obtained. Usually the ratios database is implemented as a binary search tree. Since it is likely that some similarity profiles are not observed in match or non-match sets, interpolation and extrapolation of those profiles become necessary. Moreover, the similarity profiles are assumed to be monotonic. Quadratic programming is adopted to accomplish the monotonicity enforcement, the interpolation and the extrapolation.

2.4 Disambiguation

2.4.1 Clusters

The smallest unit of disambiguation is a cluster, which contains one or more records. A cluster is interpreted as a unique inventor. The ultimate goal of disambiguation is to finalize clusters and their components.

2.4.2 Probability of Match

When two records are compared to find out the probability of match, their similarity profiles are calculated first by comparing relevant attributes. Then the corresponding r-value is looked up from the ratios database. Finally, the Bayesian theorem is applied to get the final probability of match, with the help of a priori probability which is block-dependent.

2.4.3 Cluster-based Disambiguation

During disambiguation, clusters belonging to the same blocks are compared. No inter-block cluster comparisons would happen. When two clusters are compared, their component records are compared exhaustively to find the interactive force between the two clusters. The sum of the interactive probabilities is compared to internal probabilities of both clusters. If the comparison passes a predetermined threshold, the two clusters are determined as of a same inventor and are merged together. The process continues independently within each block until no merge can happen. Once all blocks are disambiguated, the disambiguation process of the current round ends.

3 Configuration

3.1 Prerequisites

- A high-end workstation (usually 8 CPUs and 24G RAM). 24G RAM is probably the minimum memory requirement. Since multi-threading is supported, more CPUs generally reduce time cost.
- Linux (Ubuntu, Fedora or other compatibles)
- GCC/G++ (4.0 or higher)
- IBM CPLEX
- SQLite3 (3.6 or higher) (This is not mandatory but necessary if one needs to out put the result to a sqlite3 database).

4 Configuration and Compilation of Executables

4.1 Configure IBM CPLEX

IBM CPLEX package for academia is usually free for download and use. Trial versions will not work as limitations apply. Follow the instructions to install. Pay attention to the architecture (32/64 bit).

4.1.1 Configure SQLite3

SQLite3 is a public domain, light weight SQL database engine. Download from website and install.

4.1.2 Configure Makefile

Edit the file `makefile` in the source code package. Things include

- The path of IBM CPLEX library and header files (pay attention to architecture.)
- The path of SQLite3 library and header files
- System libraries and header files (math library, pthread library, etc).
- Optimization flags (optional).

4.1.3 Compile

In the command line, in the directory where the file makefile is, type make. Source codes will compile automatically. A successful compilation will generate several executables, namely exedisambig and txt2sqlite3. The former is the main disambiguation program, and the latter is the executable that dumps the disambiguation result into sqlite3 databases. Errors in compilation are usually caused by incorrect configurations in the makefile file. If necessary, type make clean to remove all the newly built object files, and make to rebuild.

4.2 Configuration of Disambiguation

The disambiguation is configured by two parts: the disambiguation engine and the blocking rules.

4.2.1 Engine Configuration

The configuration of the disambiguation engine includes the following items:

- Working directory: the directory where all the intermediary and final files will be saved.
- Original CSV File: the source database of disambiguation in text format, which is usually exported from a SQL database. The first line of the file consists of names of each attribute/column, and the rest lines are data.
- Number of threads: the number of maximum threads to allow multi-threading of the program. Usually it is set to be the number of CPUs in the computer.
- Generate stable training sets: match and non-match training sets from rare names are usually very stable. Therefore, if those training sets already exist, this switch can be set to false. However, since the creation of the stable training sets generally does not take too much time, it is usually recommended to be true.
- Use available ratios database: this switch controls whether or not training should take place. Since training is usually somewhat time-costly (4 hours), this switch can be set to true if training is unnecessary, such as when debugging. But generally this option should be set to false.
- Thresholds: the thresholds series to determine whether or not two records are a match should be set in a permissive way, such as 0.99, 0.98, 0.95. The reason to do this is to allow most similar records conglomerate first, which can help improve the accuracy.
- Necessary attributes: This option selects those necessary attributes to load into the memory from the original CSV file, because not all the data in the original CSV file need to be loaded. These necessary attributes should be exactly the same as those appearing in the first line of the CSV file.
- Adjust prior by frequency: This option controls the evaluation of the priori probability for each block. It is an option that should always set to be true unless modification of the code occurs after thorough understanding of the code.
- Debug mode: this switch indicates whether the program is for debugging or not. Set to **false** if a normal disambiguation is desired. In debug mode, the program will look for the debug configuration file debug_block_x.txt in the working directory, where x is the current round of disambiguation. The debug configuration file contains all the specified blocks that are of interest, and the program only disambiguates those blocks.
- Number of training pairs: the maximum of pairs of records obtained for training. Usually set to 10 million.

- Starting round: this number specifies the round of disambiguation to start with.
- Starting file: this string specifies the file of previous disambiguation result from which the disambiguation starts. If the starting round is 1, the starting file just needs to be a valid location in the file system, because it will be automatically created or overwritten.
- Postprocess after each round: the switch determines whether or not post-processing should be applied after each round of disambiguation. It can be either true or false, although true is recommended.

4.2.2 Blocking Configuration

The blocking configuration file determines the blocking rules and the similarity profile components for each round of disambiguation. It is in the format of:

```
[ Round X ]
```

```
Attribute Name 1: String Manipulation parameters
```

```
Attribute Name M: String Manipulation parameters
```

```
Active Similarity Attributes: attribute names of similarity profile components.
```

```
[ Round Y ]
```

```
Attribute Name 1: String Manipulation parameters
```

```
Attribute Name N: String Manipulation parameters
```

```
Active Similarity Attributes: attribute names of similarity profile components.
```

In the current engine, a typical example is:

```
[ Round 5 ]
```

```
Firstname: 1 : 0 , 3 , true
```

```
Middlename: 1 : 0 , 0 , false
```

```
Lastname: 0 : 0, 5, true
```

```
ACTIVE SIMILARITY ATTRIBUTES: Firstname, Middlename, Lastname, Latitude, Coauthor,  
Assignee
```

This means that the blocking identification for round 5 is created from three attributes:

- First name: the raw data is read from the index 1 position of the first name attribute object (see more details in source codes), and is truncated from the index 0 position of the string for a maximum length of 3 characters in a forward direction (left to right).
- Middle name: the raw data is read from the index 1 position of the middle name attribute object (see more details in source codes), and is truncated from the index 0 position of the string for a maximum length of 0 characters in a backward direction (right to left).
- Last name: the raw data is read from the index 0 position of the last name attribute object (see more details in source codes), and is truncated from the index 0 position of the string for a maximum length of 5 characters in a forward direction (left to right). And the components of similarity profiles are: first name, middle name, last name, latitude (geographical location), coauthor, assignee (company).



Figure 1: An Algorithm.

4.3 Export Disambiguation Results

The results of disambiguation can be exported to a SQLite database by running the `txt2sqlite3` executable file.

5 Implementation Details

5.1 Data Structure

The disambiguation engine is designed in an object-oriented way. The key concepts in the whole engine include:

- **Attributes:** conceptualized as the information in each column in the database. The attribute class has a complicated hierarchy and thus requires much more understanding. As for the implementation, each CONCRETE attribute class has multiple pointers to data strings. See more details in the source code.
- **Records:** a vector of attributes representing a full record in the database. More strictly, each record contains a vector of pointers to attributes.
- **Clusters:** conceptualized as a unique inventor having multiple records. Each cluster contains a list of pointers to records.

The following picture shows the data structure of the disambiguation.

5.2 Customization of Attributes

The attribute class has several layers of inheritance, each of which implements some more functionality and thus becomes more concrete. At the very low level of the hierarchy, there are several modes to choose.

Ideally one only needs to pick up a mode to inherit if a concrete class is supposed to be defined. It is also the class creator's responsibility to know and to define (or override) the following data members and member functions for a given concrete class:

- Class name: the name of the concrete class.
- Class name specifier string: the name specifier of the concrete class, which can be used to retrieve data from the original data file.
- Class group: the training group of the class, such as personal side, patent side, or none.
- Name specifiers of interactive classes: the specifiers of the classes that is necessary for the current class in order for a viable comparison definition.
- Number of interactive classes: the number of interactive classes.
- Maximum value: the maximum score of the comparison function of the class.
- Compare: the comparison function of the class.
- Other functions if necessary.

Here is a typical example of the customization of the Firstname attribute and the Latitude attribute. For Firstname:

- Class name: `cFirstname`. This class has no interaction with other classes and thus simply inherits from the mode `cAttribute_Single_Mode<cFirstname>`.
- Class name specifier string: `Firstname`. This is exactly the same as the one in the original database.
- Class group: `Personal`. First name is in the personal information side.
- Name specifiers of interactive classes: (empty).
- Number of Interactive classes: 0
- Maximum value: 4. The rating is user-defined and has to be consistent with the comparison function.
- Compare: Override the parent class definition.
- Other functions:
- `split_string`: defines how the data is extracted from the original database and stored in the class object.
- `exact_compare`: defines how the exact comparison function works.

For Latitude:

- Class name: `cLatitude`. This class has interaction with several other attributes, so it inherits from `cAttribute_Interactive_Mode<cLatitude, cLatitude_Data>`, where `cLatitude_Data` contains the real data and inherits from `cAttribute_Singe_Mode<cLatitude_Data>`.
- Class name specifier string: `Latitude`.
- Class group: `Patent`
- Name specifiers of interactive classes: `Longitude, City, Country`
- Number of interactive classes: 3
- Maximum value: 5.
- Compare: Overridden.

5.3 Customization of Cluster Comparisons

Comparison between two clusters, a critical part of disambiguation, includes several configurable steps that can be very influential to the results. When two clusters are compared:

- The two representatives of the clusters are compared to get a post-probability. If the probability does not pass a minimum screening threshold, the clusters are judged to be of different inventors and should not merge. The comparison of the clusters stops here.
- If the probability passes the minimum threshold, then:
 - Another threshold, dependent on the features of the clusters, is calculated and will be used as a caliber in the further disambiguation of the two clusters. The threshold depends on the cohesions of the two clusters, the career gap between the clusters, the number of common locations between the clusters, etc.
 - A priority queue of a given size is created. The size of the queue depends on the size of both clusters. The larger the clusters, the larger the queue size.
 - Exhaustive comparisons between members of the two clusters are performed, and the results are fed into the priority queue in the following manner.
- If the priority queue is not full, the new result fills into the queue.
- If the queue is full, the new result is compared with the lowest value in the queue.
- If the new result is greater than the lowest value, and the lowest value is less than the caliber threshold, then the new result simply replaces the lowest value in the priority queue.
- If the new results is less than the lowest value but greater than the caliber threshold, the new result is added to the priority queue.
- If the new result is less than the lowest value and also less than the caliber threshold, discard the new result.
 - The average value of the priority queue is then calculated after the exhaustive comparisons finish. If the average is greater than the caliber threshold, the two clusters are labeled as of a same inventor and will be followed by a merge of the clusters; otherwise, the comparison of the clusters stops here.
- If the clusters are labeled as a merge, the two clusters will merge into a new bigger cluster, with the previously calculated average value of the priority queue as its new cohesion value (or with-in-cluster density). A new representative of the new cluster will be decided later.

Therefore, the customization of the cluster comparisons can be in:

- The screening threshold
- The caliber threshold
- The size of the priority queue
- The way to calculate the new cohesion value (the with-in-cluster density)