An Introduction to Entity Resolution

Rebecca C. Steorts

Department of Statistical Science, affiliated faculty in Computer Science, Biostatistics and Bioinformatics, the information initiative at Duke (iiD) and the Social Science Research Institute (SSRI)

Duke University and U.S. Census Bureau

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Entity resolution (record linkage or de-duplication) joins multiple data sets removes duplicate entities often in the absence of a unique identifier.

Motivations

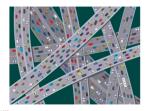
In Syria, we have duplicated information regarding individuals who have died in the conflict.

In the census, we have duplicated information of individuals that fill out census forms every 10 years.

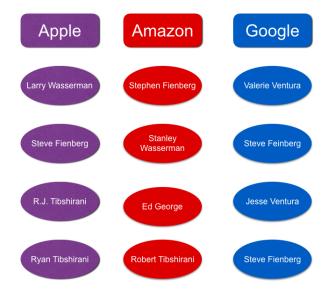
Goal: Estimation of the sample size and standard errors. Additional Goal: Posterior densities, uncertainty quantification, etc.



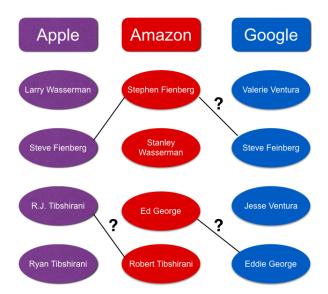
BAD NEWS FOR EVERYONE! THE 2020 CENSUS IS ALREADY IN TROUBLE



A graph with no edges



The entity resolution graph



The node of Larry Wasserman

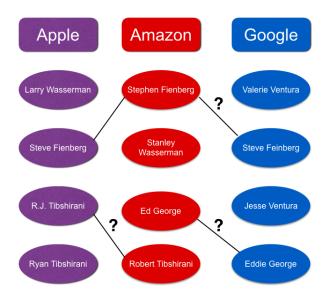


The node of Larry Wasserman



1014 Murray Hill Avenue Pittsburgh, PA 15217 412-361-3146

The entity resolution graph



Who's the real Steve Fienberg?



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Who's the real Steve Fienberg?

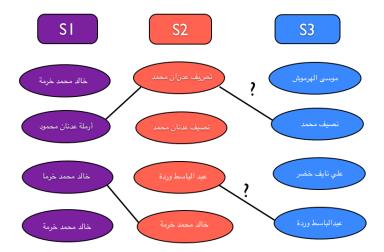


Stephen Fienberg

240 Collins Drive Pittsburgh, PA 50-54 412-793-3313 537 N Neville Street Pittsburgh, PA 15213 65+ 412-683-5599

These are clearly not the same Steve Fienberg!

Syrian Civil War



Entity Resolution

Why is entity resolution difficult?

Goals of Entity Resolution

Suppose that we have a total of M records in D data sets.

- 1 We seek models that are much less than $O(M^2)$ (quadratic).
- We seek models that are reliable, accurate, fit the data well, and account for the uncertainty of the model.

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Terminology

- De-duplication
- 2 Record linkage
- 8 Entity resolution
- 4 Blocking

De-duplication



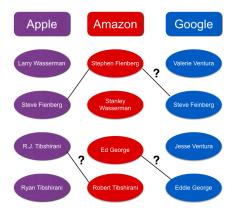
De-duplication

Much of the literature can be grouped into the case of de-duplication.

Common examples from both academia and industry are the following:

logistic regression, random forests, support vector machines, Bayesian adaptive regression trees, and locality sensitive hashing.

The entity resolution graph



De-duplication refers to duplication within just one database. **Record linkage** refers to duplication across multiple databases. **Entity resolution** refers to both duplication across and within databases (the entire graph).

Blocking

Often one performs blocking due to the fact that record linkage problems require a quadratic number of comparisons.

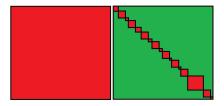


Figure: All-to-all record comparisons (left) versus partitioning records into blocks and comparing records only within each partition (right).

We will assume some method of blocking is embedded within a record linkage procedure.

Blocking

The most common method used for blocking is typically

- 1 deterministic blocking method
- 2 probabilistic blocking method

Examples include blocking on features (deterministic) or probabilistic types such as locality sensitive hashing.

See Christen (2012); Steorts, Ventura, Sadinle, Fienberg (2014); Chen, Shrivastava, Steorts (2018).

Common Methods for Entity Resolution

- Match on a unique identifier if it exists.
- Perform exact matching.
- Perform a likelihood ratio or hypothesis test.

[Newcombe (1959), Fellegi and Sunter (1969)].

Unique Identifier

Suppose that each feature has a unique identifier that we are sure is accurate, like social security number.

Then we can unique match records based on the unique identifier.

Problems occur this unique identifier is missing or has noise in it, etc.

Exact Matching

In exact matching, we compare all features. We decide if the record is a match if they agree on all features. Otherwise, we decide the record is a non-match.



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Why would this method be bad for evaluation purposes?

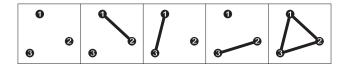
Fellegi and Sunter Method

- Newcombe (1959), Fellegi and Sunter (1969): two databases, all-to-all comparison of records.
- Neyman Pearson hypothesis test with a threshold t.
- If record *i* and *j* are above *t*, then we have a match.
- Otherwise, a non-match.

Fellegi and Sunter Method

- Computationally intractable.
- Transitivity not preserved.
- If 1 matches 2, and 2 matches 3, then 1 does NOT necessarily match 3.

Major limitations and major flaws of this method!



Evaluation Metrics

How do we evaluate performance of a particular record linkage method?

Evaluation Metrics

- Recall
- 2 Precision
- 3 Reduction Ratio
- 4 Estimated Sample Size
- 5 Standard Error of Estimated Sample Size
- 6 Run Time
- Robustness to Tuning Parameters
- 8 Do Supervised Methods Overfit the Data

Evaluation Metrics

- 1 Pairs of data can be linked in both the handmatched training data (which we refer to as "truth") and under the estimated linked data. We refer to this situation as true positives (TP).
- 2 Pairs of data can be linked under the truth but not linked under the estimate, which are called false negatives (FN).
- 3 Pairs of data can be not linked under the truth but linked under the estimate, which are called false positives (FP).
- Pairs of data can be not linked under the truth and also not linked under the estimate, which we refer to as true negatives (TN).

Recall, Precision, Reduction Ratio

$$Recall = \frac{TP}{TP + FN} = 1 - FNR.$$

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Recall, Precision, Reduction Ratio

$$\begin{aligned} \text{Recall} &= \frac{TP}{TP + FN} = 1 - FNR. \\ \text{Precision} &= \frac{TP}{TP + FP} = 1 - FPR. \end{aligned}$$

Reduction ratio (RR) measures the relative reduction of the comparison space from the de-duplication or hashing technique.

See Christen (2012), Steorts, Ventura, Sadinle, Fienberg (2014) for a formal definition.

Other Evaluation Metrics

- 1 Estimated Sample Size
- Standard Error of Estimated Sample Size
- 3 Run Time
- 4 Robustness to Tuning Parameters
- 5 Do Supervised Methods Overfit the Data
- 1 The estimated sample size and standard error must be defined for each method, but this is not difficult to do in practice.
- 2 Any method can be evaluated also for the run time, so one can gauge computationally costs.
- 3 Robustness of tuning parameters should be explored from a Bayesian and a frequentist perspective.
- 4 It's also essential to make sure that supervised methods do not overfit the data (see Steorts (2015)).

RLdata500 dataset

Let's look at an example on data that is available from the Record Linkage package in R, where we compare many different methods according to the evaluation metrics that we have laid out.

We will first describe the data set and then compare the following methods in R:

- blink
- 2 logistic regression
- 3 random forests
- 4 Bayesian adaptive regression trees

RLdata500 dataset

```
fname_c1 fname_c2 lname_c1 lname_c2 by bm bd
  CARSTEN
             <NA>
                    METER.
                             <NA> 1949 7 22
     GERD
             <NA> BAUER
                             <NA> 1968
                                      7 27
3
   ROBERT
             <NA> HARTMANN
                             <NA> 1930
                                      4 30
4
   STEFAN
             <NA>
                    WOLFF <NA> 1957
5
     RALF
           <NA> KRUEGER
                           <NA> 1966 1 13
6
  JUERGEN
             <NA>
                   FRANKE
                             <NA> 1929
                                       7 4
```

The RLdata500 data set consists of 500 records with 10 percent duplication.

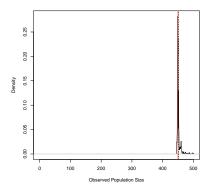


Figure: Shared MPMMS graphical representation of simulation study. Only makes

one false positive set.

Figure: Posterior density for N in simulation study. The FNR and FPR: 0.04 and 0.02.

Procedure	FNR	FDR
blink (Steorts (2015))	0.02	0.08
Exact Matching	1	0
Near-Twin	1	0
BART (10% training)	0.10	0.16
BART (20% training)	0.07	0.11
BART (50% training)	0.03	0.04
BART (full data)	0.02	0
Random Forests (10% training)	0.05	0.15
Random Forests (20% training)	0.04	0.09
Random Forests (50% training)	0.02	0.06
Random Forests (full data)	0.04	0.06
Logistic Regression (10% training)	0.09	0.16
Logistic Regression (20% training)	0.06	0.07
Logistic Regression (50% training)	0.02	0.01
Logistic Regression (full data)	0.02	0

Table: False negative rate (FNR) and false discovery rate (FDR) for the proposed EB methodology and five other record linkage methods. For the supervised methods, we run 100 iterations of each one and average these ◆□▶ ◆□▶ ◆■▶ ◆■▶ ■ 夕♀◎ such that overfitting is not occurring.

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Robustness

How do we make sure a method is robust?

For a semi-supervised method, we want to make sure that it's robust to different choices of the training/test data and any tuning paramater(s).

For probabilistic and Bayesian methods, we want to make sure these methods are robust to choices of hyper-parameters and/or tuning parameters.

Robustness, computational time complexity, and sensitivity analysis can be further explored in Steorts (2015) and Steorts, Hall, Fienberg (2016).

Some Key Papers

- Christen (2012); Steorts, Ventura, Sadinle, Fienberg (2016) [Fair Comparisons]
- Sadinle (2014), Steorts, Hall, Fienberg (2016) [Fair Comparisons]
- Vatsalan, Christen, O'Keefe, Verykios (2014) [Evaluations of Record Linkage and Privacy]
- Victoria Stodden and David Dunson [Reproducibility and Fairness]
- 6 Chen, Shrivastava, Steorts (2018) [Reproducibility, Fairness, and Evaluations]
- Oavid Banks, Megan Price, Patrick Ball [Ethical Use of Data with respect to Human Rights]

This Short Course

- An Introduction to Entity Resolution (Just Completed)
- 2 An Introduction to Blocking (Demos)
- 3 Bayesian Approaches to Graphical Entity Resolution (Demos)
- 4 Distributed Bayesian Entity Resolution (Demos)

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There are other following materials that may be of interest after completion of this short course:

- An Introduction to Locality Sensitive Hashing (Demos)
- Unique Entity Estimation Applied to the Syrian Conflict (Demos

https://resteorts.github.io/record-linkage-tutorial/

Thank you! Questions?

Contact: beka@stat.duke.edu