Introduction to Probabilistic Record Linkage

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What is Record Linkage?

- Record linkage is the task of identifying duplicate records over noisy datasets.
- Easy with unique identifiers, difficult when faced with errors
- **Bipartite matching** is the specific goal of matching one record in one dataset to most one match in another dataset

Record Linkage in Practice

Duke TODAY



MAKING SENSE OF SYRIA'S MURKY DEATH TOLL



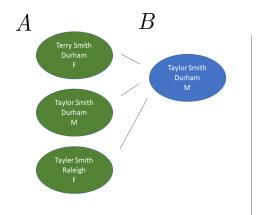
Linkage for Downstream Analysis

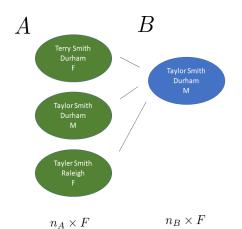
Response Variable	Personal Identification Information		

Personal Identification Information		Covariates	

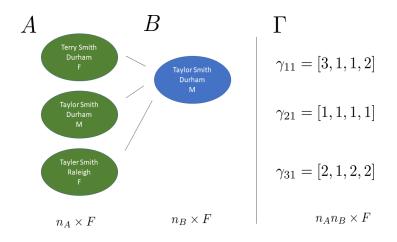
Linkage for Downstream Analysis

Response Variable	Personal Identification Information	2	Personal Identification Information	Covariates
		•		



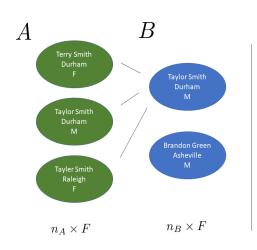


- n_A, n_B records in A, B
- \bullet F=4 features for comparison
 - First name
 - Last name
 - City
 - Gender
- $L = \{3, 3, 2, 2\}$ levels of comparison



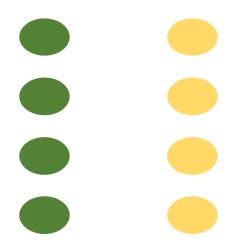
Represent linkage structure through vector $\mathbf{Z} = \{Z_1, \dots, Z_{n_B}\}$, where

$$Z_j = egin{cases} i, & ext{if records } i \in A ext{ and } j \in B ext{ match;} \\ n_A + 1, & ext{if record } j \in B ext{ has no match in } A; \end{cases}$$

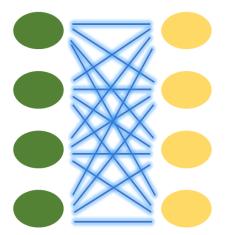


$$Z_1 = 2$$

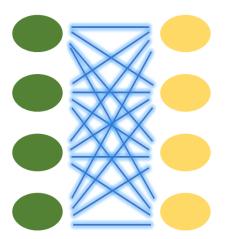
$$Z_2 = n_A + 1$$



 $n_A n_B$ independent decisions

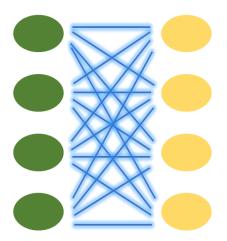


 $n_A n_B$ independent decisions



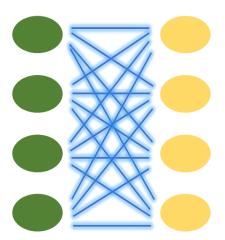
 scalable to large datasets (fastlink, Enamorado et al 2019)

 $n_A n_B$ independent decisions



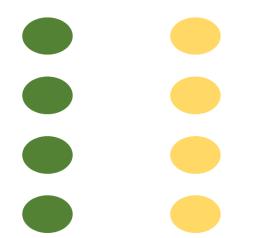
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- not bipartite, requires post-processing

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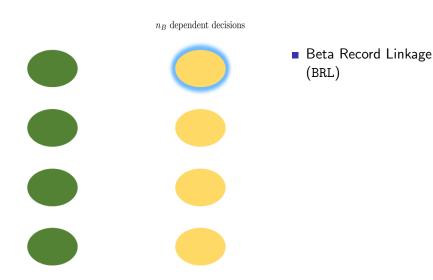


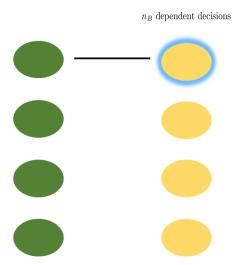
- scalable to large
 datasets (fastlink,
 Enamorado et al
 2019)
- not bipartite, requires post-processing
- overmatches, leading to inaccurate parameter estimation

- The Fellegi and Sunter model is essentially a mixture model
- We posit that comparison vectors come from two different distributions, one for matching, and another for nonmatching record pairs
- The model learns the parameters for each distribution and independently classifies each record pair as matching or nonmatching.

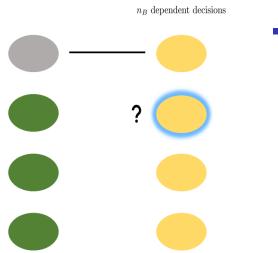


Beta Record Linkage (BRL)

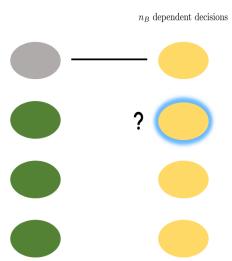




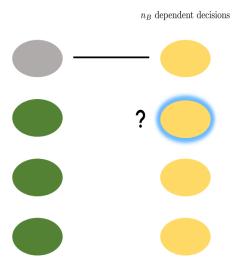
Beta Record Linkage (BRL)



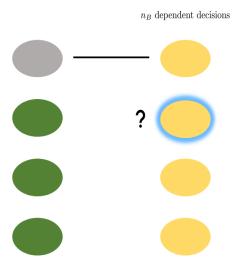
Beta Record Linkage (BRL)



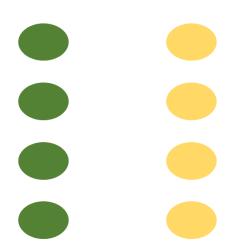
- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing

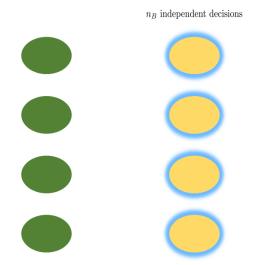


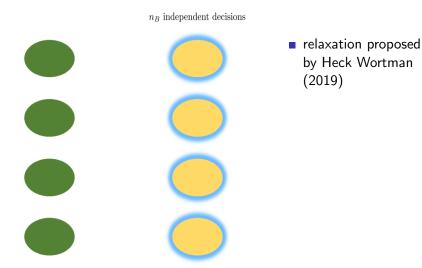
- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing
- high accuracy for linkage and other parameters

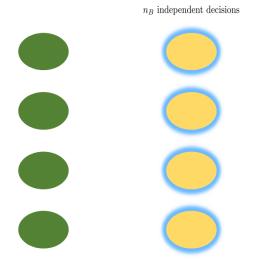


- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing
- high accuracy for linkage and other parameters
- inherently serial, not scalable to large linkage tasks



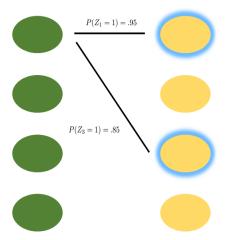






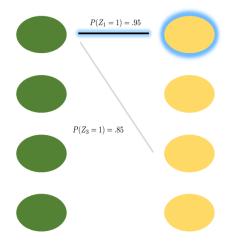
- relaxation proposed by Heck Wortman (2019)
- minimal loss of accuracy, large computational gains





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- relaxation proposed by Heck Wortman (2019)
- minimal loss of accuracy, large computational gains
- allows for "one to many" matchings
- simple
 postprocessing to
 obtain bipartite
 matching

Model Parameters

- Both models utilize **m** and **u** parameters to model the data.
- Define $m_{f\ell} = P(\gamma_{ij}^f = \ell | \text{Records Match})$ and $u_{f\ell} = P(\gamma_{ij}^f = \ell | \text{Records Do Not Match})$
- We can think of **m** as reliability parameters, that show how likely it is for a feature to remain unchanged across files
- We can think of u as discernment parameters, that show how unique the values of a feature can be

Fellegi Sunter Weights

- We multiply these parameters for all f and the relevant ℓ to get $\mathbf{m}_{ij} = P(\gamma_{ij}|\text{Records Match})$ and $\mathbf{u}_{ij} = P(\gamma_{ij}|\text{Records Do Not Match})$
- These probabilities form the likelihood ratio

$$w_{ij} = \frac{P(\gamma_{ij}|\text{Records Match})}{P(\gamma_{ij}|\text{Records Do Not Match})}$$

■ This ratio is large when \mathbf{m}_{ij} is large and \mathbf{u}_{ij} is small

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Constructing Comparison Vectors

- We measure the distance between two fields using whatever metric we like, and then threshold this distance into discrete values
- We can use binary agreement, or use multiple agreement levels to capture partial agreement
- Existing packages perform binary agreement, numerical distance, and string distance

String Distance

- Many packages compare strings through the normalized Levenstein distance. This is the minimum number of edits required to transform one string to the other, divided by the length of the larger string.
- Additional metrics could be language specific Soundex codes
- Care should be taken when transforming metric into discreet agreement level. For example, should a Soundex match be regarded as a full match or a partial match?
- Code from existing packages can be edited to accommodate additional string distances

Conditionally Independent Features

- The record linkage model assumes that agreement levels are conditionally independent, given the matching status of the records. This is satisfied for most fields
- For example, knowing that two records agree on birth month does not tell you anything about whether they agree on birth year. These fields are independent
- In contrast, if two records disagree on department, we know they probably disagree on city. These fields are not independent
- Using dependent features is essentially using the same information multiple times, and can harm results

Conditionally Independent Features

- One solution to dependent features is to create a more nuanced comparison incorporating multiple features
- For example, you can define full agreement for "location" to be matching on city AND department, partial agreement to be matching on only department, and no agreement not be matching on neither.
- Current packages do not execute this type of comparison, but this can be done with some work
- Depending on the type of dependent fields, I may recommend keeping both or just omitting one.

Conditionally Independent Features

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Random Sampling and Generalization

- To avoid calculating $n_A n_B$ comparison vectors, we can use blocking
- We only create comparison vectors when records agree on a particular field
- It may be tempting to block on a unique identifier, but this can create blocks that are too small for record linkage to run effectively

Thoughts for DANE

- With fastLink, you may be able to do the entire linkage task $(60,000 \times 1,000,000)$ records at one time. If not blocking by gender or by department should make it feasible
- You should not need to use the generalization technique
- fastLink currently does not do Soundex comparisons, but using allowing for partial agreement on strings should be sufficient. fastLink code can be modified to incorporate Soundex comparison, but may take some work

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Performance Metrics with Known Labels

- When true matching status is known, we use two primary metrics:
- To balance these two metrics, we sometimes also consider:
 - F-Measure = Recall × Precision Recall + Precision

Performance Metrics without Known Labels

- Without known labels, evaluating model performance is more difficult, but there are general checks you can make
 - The histogram of match probabilities should be clearly separated, with most record pairs having probabilities near 0 or near 1, and minimal number of pairs in between
 - Fellegi Sunter weights for impactful matching variables should be high. Weights for partial agreement should be lower than those for full agreement
 - Visually inspect a subset of matches and nonmatches. In particular, examine matches near the decision threshold.
- These approaches are demonstrated in the accompanying Rmarkdown file

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Software

- fastLink can be found on GitHub
- BRL can also be found on GitHub
- fabl is not yet publicly available
- Due to the scale of DANE's record linkage task, I recommend you to use fastLink

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References

- Fellegi I. and Sunter A., A theory for record linkage, J. Am. Stat. Assoc. 64 (1969), pp. 1183–1210
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- Sadinle M., Bayesian estimation of bipartite matchings for record linkage, J. Am. Stat. Assoc. 112 (2017), pp. 600–612