Efficient and Scalable Bipartite Matching through Fast Beta Linkage (fabl)

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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
- Far ranging applications in business, public health, and human rights
- **Bipartite matching** is the specific goal of matching one record in one dataset to most one match in another dataset

Record Linkage in Practice

Monkey Cage - Analysis

Georgia's 'exact match' law could potentially harm many eligible voters.



Georgia gubernatorial candidates Stacey Abrams, left, and Brian Kemp on May 20 in Atlanta, (John Amis/AP)

By Ted Enamorado October 20, 2018



Record Linkage in Practice

Monkey Care - Anah

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Georgia gubernatorial candidates Stacey Abrams, left, and Brian Kemp on May 20 in Atlanta, (John Amis/AP)

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DNC Announces New National Record Linkage System

APRIL 24, 2020

f y

Algorithm developed by DNC expert in the field of record linkage will increase organizing efficiency by 9 percent and provide campaigns with more comprehensive view of the overall electorate

Linkage for Downstream Analysis

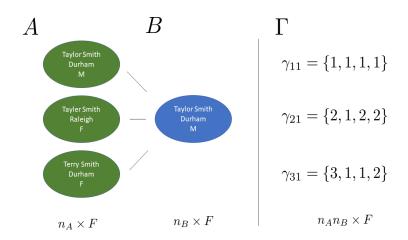
Response Variable	Personal Identification Information				

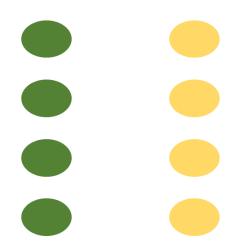
Personal Identification Information		Covariates			

Linkage for Downstream Analysis

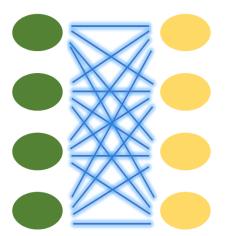
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		•		

Linkage through Comparison Vectors

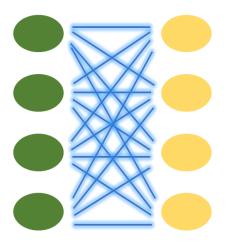




 $n_A n_B$ independent decisions

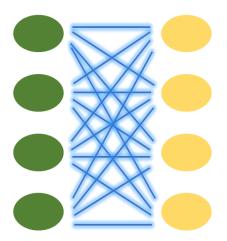


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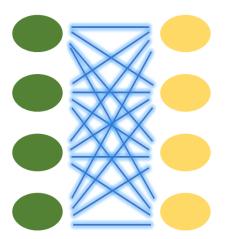
scalable to large
datasets (fastlink,
Enamorado et al
2017)

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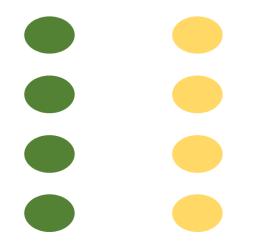


- scalable to large datasets (fastlink, Enamorado et al 2017)
- no transitive closure, requires post-processing

 $n_A n_B$ independent decisions

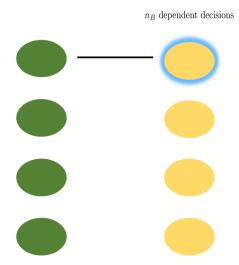


- scalable to large datasets (fastlink, Enamorado et al 2017)
- no transitive closure, requires post-processing
- overmatches, leading to inaccurate parameter estimation

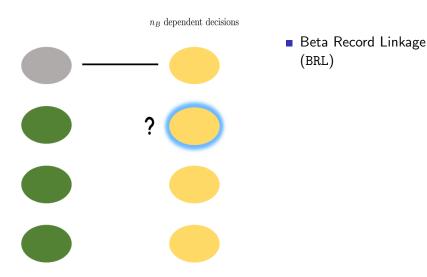


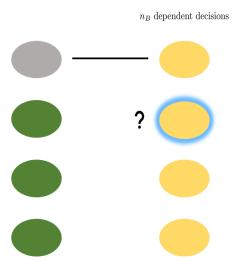
Beta Record Linkage (BRL)



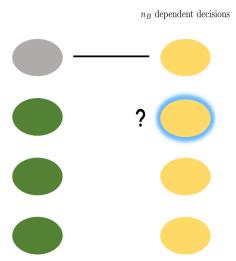


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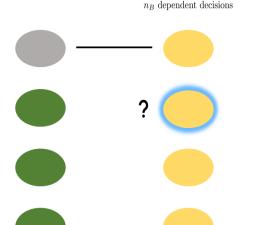




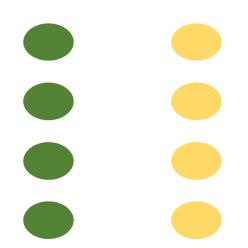
- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing

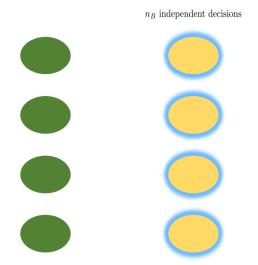


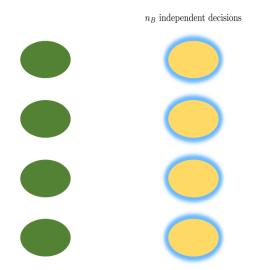
- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing
- high accuracy for linkage and other parameters



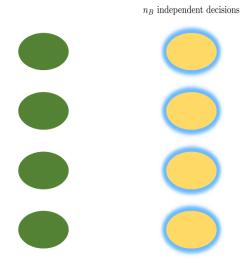
- Beta Record Linkage (BRL)
- strictly enforces one-to-one matching, no post-processing
- high accuracy for linkage and other parameters
- inherently serial, not scalable to large linkage tasks







 simple mathematical change, large computational gains



- simple mathematical change, large computational gains
- minimal loss of accuracy for linkage and other parameters, minimal post-processing

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Notation

- File A with records indexed $i \in \{1, ..., n_A\}$ and file B with records $j \in \{1, ..., n_B\}$, with $n_A \ge n_B$. We use F features for linkage, with L_f possible levels of agreement on feature f.
- $oldsymbol{\Gamma} \in \mathbb{R}^{n_A n_B imes F}$ matrix of comparison vectors where $\gamma_{ij}^f \in \{1,\dots,L_f\}$
- $m_{fl} = P\left(\gamma_{ij}^f = I|Z_j = i\right)$
- $u_{fl} = P\left(\gamma_{ij}^f = I|Z_j \neq i\right)$

Fast Beta Linkage (fabl)

$$P(\Gamma | \mathbf{Z}, \mathbf{m}, \mathbf{u}) = \prod_{j=1}^{n_B} \prod_{i=1}^{n_A} \left[\prod_{f=1}^F \prod_{l=1}^{L_f} m_{fl}^{I(Z_j = i)} u_{fl}^{I(Z_j \neq i)} \right]^{I(\gamma_{ij}^f = l)}$$

$$\mathbf{m_f} \sim \text{Dirichlet}(\alpha_{f1}, \dots, \alpha_{fL_f})$$

$$\mathbf{u_f} \sim \text{Dirichlet}(\beta_{f1}, \dots, \beta_{fL_f})$$

$$Z_j | \lambda = \begin{cases} \frac{1}{n_A} \lambda & z_j \leq n_A; \\ 1 - \lambda & z_j = n_A + 1 \end{cases}$$

$$\lambda \sim \text{Beta}(\alpha_{\lambda}, \beta_{\lambda})$$

Model specification allows for parallel/distributed computing, hashing of comparison vectors, and storage efficient indexing (SEI)

Hashing

- Recognize there are at most $P = \prod_{f=1}^{F} L_f$ unique agreement patterns, regardless of number of records. When (i,j) pair exhibits agreement pattern p, say h(i,j) = p.
- Reduce data to sufficient statistics
 - $r_{p_j} = \{i \mid (i,j) \in h_p\}$
 - $\blacksquare H_{p_j} = ||r_{p_j}||$
 - $\blacksquare H_p = \sum_j H_{p_j}$
- Run Gibbs sampler at level of agreement patterns, not record pairs
 - Sample the agreement of pattern $h(z_j, j)$, instead of record label z_j .
 - $lue{}$ Use number of matches for each pattern to update m and u
 - lacksquare Back fill record labels at the end through r_{p_j}
- Reduces computational complexity from $O(n_A \times n_B \times F)$ to $O(P \times n_B \times F)$



Managing Large Data

- **Distributed Computing** Partition data in to chunks $\{A_I\}$ and $\{B_J\}$. Compare records, hash results, compute summary statistics in parallel, and synthesize results.
- Storage Efficient Indexing (SEI) Store at most small number R many record labels in each r_{p_j} , remove highly unlikely record labels from memory. Proper weights for calculations maintained through summary statistics $\{H_p\}$ and $\{H_{p_i}\}$.
- Hashing plus SEI can reduce memory requirements by > 99.
 - Simulation of 20,000 × 20,000 linkage task with 4 fields. Naive approach requires 6.4GB of storage for all-to-all comparisons, hashing and SEI requires 90MB.

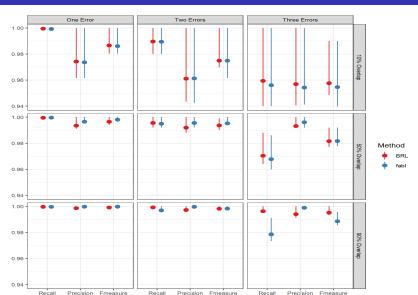
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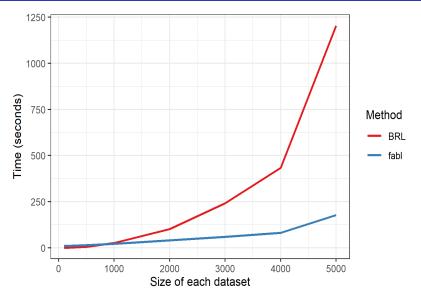
Three Simulation Studies

- We compare fabl against BRL in three simulation studies
 - Measure precision and recall on 100 simulated datasets and varying levels of error and duplication across files
 - Measure speed when both n_A and n_B are increasing
 - Measure speed when n_A is increasing and $n_B = 500$ is fixed.

Accuracy Simulation



Speed Simulation 1



Speed Simulation 2

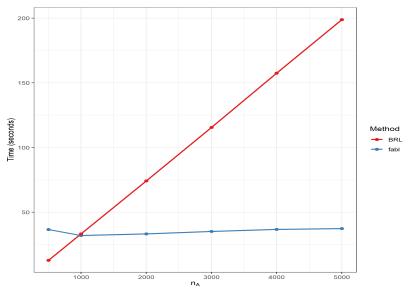


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El Salvador Case Study

- El Salvadoran Civil War (1980-1991)
- Lists of casualties collected by multiple organizations
 - Salvadoran Human Rights Commissions (CDHES), $n_A = 4420$
 - El Rescate Tutela Regal (ERTL), $n_B = 1323$
 - Features include first name, last name, date and place of death
- We aim to find duplicate records across files
 - Particularly difficult because families are often killed together, and some children share names with parents

Run Time

Linkage Results with P = 2048 patterns

	DID	time (sec)
fabl	5565	296.79
BRL	5562	269.45

Run Time

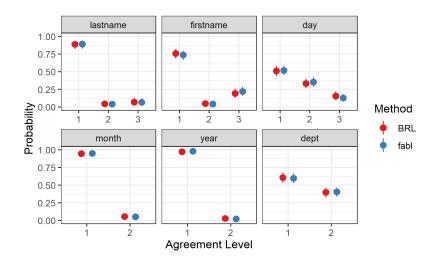
Linkage Results with P = 2048 patterns

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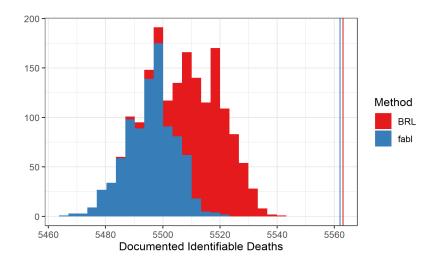
Linkage Results with P = 216 patterns

	DID	time (sec)
fabl	5564	60.42
BRL	5562	253.21

Posterior Inference



Posterior Inference



Violations of One-to-One Matching

	lastname	firstname	dataset	day	month	year	dept	muni
825	PINEDA	ROSA	CDHES	6	4	1984	NA	NA
826	PINEDA	ROSA MARIA	CDHES	6	4	1984	NA	NA
2776	PINEDA	ROSA MARIA	ER-TL	4	4	1984	CUSCATLAN	NA

Violations of One-to-One Matching

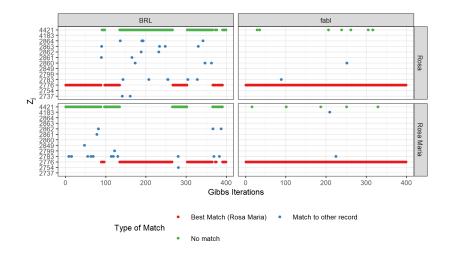


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Extensions and Directions

- Linkage in the face of duplicates within datasets
- Models that allow reliability of information to differ by subgroup in the data
- Linkage over blocked data (allows for much larger linkage tasks)