

# Introduction to Bayesian Entity Resolution

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# What is “Bayesian”?

- 1 Setting up a *full probability model* – a joint probability distribution for all observable and unobservable quantities

$p(\mathbf{x}|\boldsymbol{\theta})$  – likelihood

$p(\boldsymbol{\theta})$  – prior

- 2 Conditioning on observed data – calculating and interpreting the appropriate *posterior distribution*

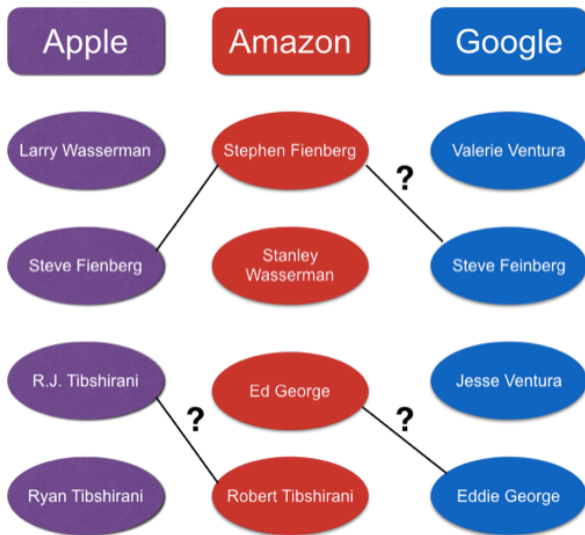
$$p(\boldsymbol{\theta}|\mathbf{x}) = \frac{p(\mathbf{x}, \boldsymbol{\theta})}{p(\mathbf{x})} = \frac{p(\mathbf{x}|\boldsymbol{\theta})p(\boldsymbol{\theta})}{p(\mathbf{x})} \propto p(\mathbf{x}|\boldsymbol{\theta})p(\boldsymbol{\theta})$$

# Why Bayesian Entity Resolution

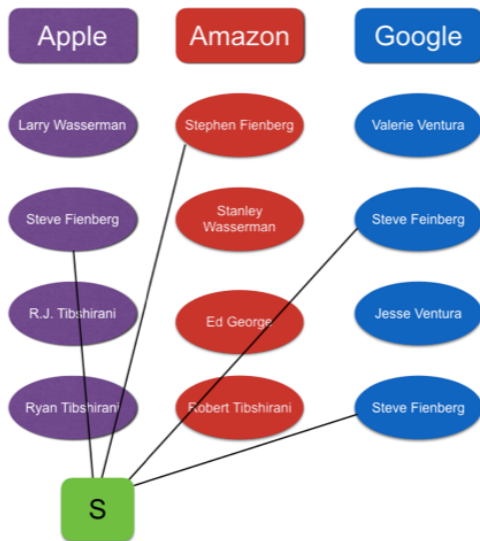
- 1 Entity resolution can be treated as a clustering problem.
- 2 Records are clustering to a latent entity.
- 3 This results in the model becoming a bipartite graph, which allows one to estimate latent individuals across multiple high dimensional databases.
- 4 The Bayesian paradigm naturally allows uncertainty quantification of the entity resolution process, a full posterior distribution, credible intervals, etc.
- 5 Theoretical properties have recently been explored for latent variable models, supporting the above approach.

[Copas and Hilton (1990), Tancredi and Liseo (2011), Steorts, Barnes, Neiswanger (2017), Zanella et al. (2016)]

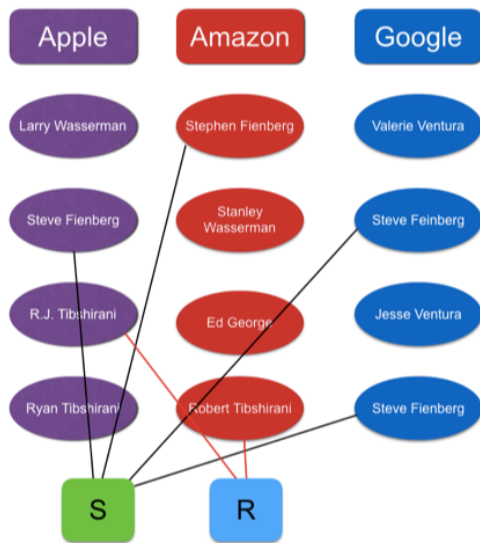
# The entity resolution graph



# The latent variable approach



# The latent variable approach

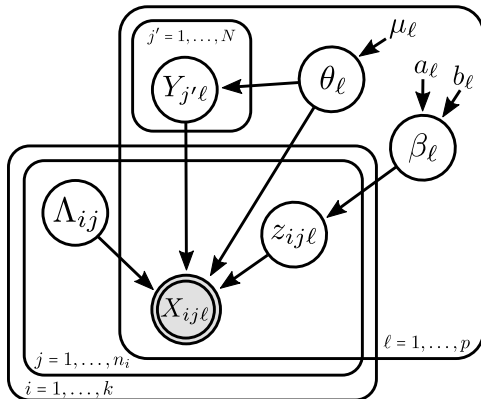


# Notation

- $X_{ij\ell}$ : observed value of the  $\ell$ th field for the  $j$ th record in the  $i$ th data set,  $1 \leq i \leq k$  and  $1 \leq j \leq n_i$ .
- $Y_{j'\ell}$ : true value of the  $\ell$ th field for the  $j'$ th latent individual.
- $\lambda_{ij}$ : latent individual to which the  $j$ th record in the  $i$ th list corresponds.  $\mathbf{\Lambda}$  is the collection of these values..
  - e.g. Five records in one list  $\mathbf{\Lambda} = \{1, 1, 2, 3, 3\} \rightarrow 3$  latent entities or clusters.
- $z_{ij\ell}$ : indicator of whether a distortion has occurred for record field value  $X_{ij\ell}$

# Graphical Record Linkage

Graphical model representation of [Steorts et al. \(2016\)](#):



- $\Lambda_{ij}$  represents the linkage structure  $\rightarrow$  **uniform prior**.
- Requires information about the number of latent entities a priori and it is very informative.



# Bayesian Entity Resolution

Previous literature: not balanced regarding modeling, handling high-dimensional data, and uncertainty of multiple databases.

- Bayesian model: simultaneously links and de-duplicates.
- Assume records are noisy, distorted.
- We have a novel representation ( $\Lambda$ ): linkage structure.
- The **strength** of a Bayesian approach is that transitivity of the linked records is nearly automatic.
- Our data structure provides uncertainty estimates of linked records that can be propagated into later analyses.

[[Steorts](#), Hall, and Fienberg (2014), [Steorts](#), Hall, and Fienberg (2016)]

# Empirically Motivated Priors

- The major weakness of [Steorts, Hall, and Fienberg \(2016\)](#) is the fact that it did not handle text (string) data.
- Steorts (2015) overcomes this issue by taking an empirical Bayesian approach, and making extensive comparisons to supervised methods.

# Model Specification: String model

- The distortion of string-valued variables is modeled using a probabilistic mechanism based on some measure of distance between the true and distorted strings.

$$P(X_{ij\ell} = w | \lambda_{ij}, Y_{\lambda_{ij\ell}}, z_{ij\ell}) = \frac{\alpha_{\ell} \exp[-cd(w, Y_{\lambda_{ij\ell}})]}{\sum_{w \in S_I} \alpha_{\ell} \exp[-cd(w, Y_{\lambda_{ij\ell}})]}$$

where  $c$  is a parameter that needs to be specified and  $d$  represents a string metric distance e.g. [Levenshtein](#) or [Jaro-Winkler](#).

# Model Specification: Likelihood Function

$$X_{ij\ell} = w | \lambda_{ij}, Y_{\lambda_{ij\ell}}, z_{ij\ell} \stackrel{iid}{\sim} \begin{cases} \delta(Y_{\lambda_{ij\ell}}), & \text{if } z_{ij\ell} = 0 \\ F_{\ell}(Y_{\lambda_{ij\ell}}), & \text{if } z_{ij\ell} = 1 \text{ and } \ell \leq p_s \\ G_{\ell}, & \text{if } z_{ij\ell} = 1 \text{ and } \ell > p_s \end{cases}$$

- $z_{ij\ell} = 0$ , then  $X_{ij\ell} = Y_{\lambda_{ij\ell}}$
- $F_{\ell}$  is the string model in the last slide.
- $G_{\ell}$  is the empirical distribution function of the categorical data.

# Model Specification: Hierarchical Model

$$Y_{\lambda_{ij}\ell} \stackrel{iid}{\sim} G_{\ell}$$

$$z_{ij\ell} | \beta_{i\ell} \stackrel{iid}{\sim} \text{Bernoulli}(\beta_{i\ell})$$

$$\beta_{i\ell} \stackrel{iid}{\sim} \text{Beta}(a, b)$$

$$\lambda_{ij} \stackrel{iid}{\sim} \text{DiscreteUniform}(1, \dots, N)$$

where  $a, b, N$  are unknown parameters that must be estimated or fixed.

- $\beta_{i\ell}$  represent the distortion probabilities of the fields.
- The parameters  $a$  and  $b$  for the Beta prior need to be specified.
- The number of latent entities or clusters needs to be specified in advance.

# blink package

R package that removes duplicate entries from multiple databses using the empirical Bayes graphical method:

```
install.packages("blink")
```

- Formatting data for use with blink
- Tuning parameters
- Running the Gibbs sampler (estimate model parameters)
- Output

## RLdata500 data

We will continue with the RLdata500 dataset in the RecordLinkage package consisting of 500 records with 10% duplication.

```
library(blink) # load blink library
library(RecordLinkage) # load data library
data("RLdata500") # load data
head(RLdata500) # take a look
```

##	fname_c1	fname_c2	lname_c1	lname_c2	by	bm	bd
## 1	CARSTEN	<NA>	MEIER	<NA>	1949	7	22
## 2	GERD	<NA>	BAUER	<NA>	1968	7	27
## 3	ROBERT	<NA>	HARTMANN	<NA>	1930	4	30
## 4	STEFAN	<NA>	WOLFF	<NA>	1957	9	2
## 5	RALF	<NA>	KRUEGER	<NA>	1966	1	13
## 6	JUERGEN	<NA>	FRANKE	<NA>	1929	7	4

# Formatting the data

```
# categorical variables
```

```
X.c <- as.matrix(RLdata500[, c("by", "bm", "bd")])
```

```
# string variables
```

```
X.s <- as.matrix(RLdata500[, c("fname_c1", "lname_c1")])
```

X.c and X.s include all files stacked on top of each other, for categorical and string variables respectively

```
# keep track of which rows of are in which files
```

```
file.num <- rep(c(1, 2, 3), c(200, 150, 150))
```



# Tuning parameters

## Hyperparameters

```
# Subjective choices for distortion probability prior  
# parameters of a Beta(a,b)  
a <- 1  
b <- 999
```

## Distortion

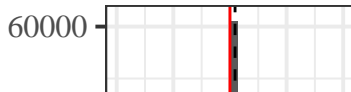
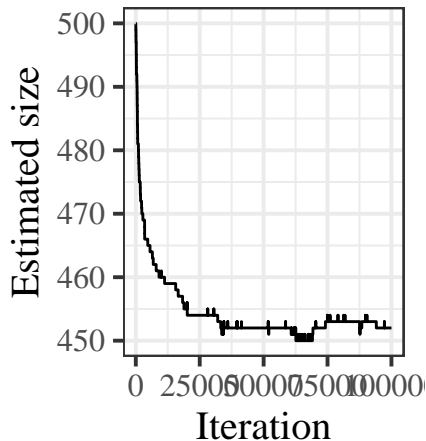
```
# string distance function example  
d <- function(s1, s2) {  
  adist(s1, s2) # approximate string distance  
}  
  
# steepness parameter  
c <- 1
```

# Running the Gibbs sampler

```
lam.gs <- rl.gibbs(file.num = file.num, # file  
                  X.s = X.s, X.c = X.c, # data  
                  num.gs = 100000, # iterations  
                  a = a, b = b, # prior params  
                  c = c, d = d, # distortion  
                  M = 500) # max # latents
```

# Output

```
# count how many unique latent individuals  
size_est <- apply(lam.gs, 1, function(x) {  
  length(unique(x))  
})
```



# Evaluation

```
# estimated pairwise links
est_links_pair <- pairwise(links(lam.gs[-(1:25000), ]))

# true pairwise links
true_links_pair <- pairwise(links(matrix(identity.RLdata500, nrow = 1)))

#comparison
comparison <- links.compare(est_links_pair, true_links_pair, counts.only = TRUE)

# precision
precision <- comparison$correct/(comparison$incorrect + comparison$correct)

# recall
recall <- comparison$correct/(comparison$correct + comparison$missing)

# results
c(precision, recall)
```

```
## [1] 1.00 0.96
```

## Your turn

Using the title, authors, year, and journal columns in the cora dataset from the RLdata package,

- 1 Let's only use data with **complete cases** (for simplicity):
- 2 Format the data to use with `blink`
  - Which columns are string vs. categorical?
  - Of the remaining data, assume that the rows 1-200 are from database 1, 201-400 are from database 2, and 401-560 are from database 3
- 3 Create tuning parameters for your model
  - Think about prior hyperparameters as well as the string distortion function
- 4 Run the Gibbs sampler 50 times to update the linkage structure
- 5 **Extra:** Evaluate your estimated linkage structure using precision and recall

# Your turn (solution)

```
# 1. complete cases of data onl
# load data
library(RLdata)
data("cora")
data("cora_gold")

not_missing <- complete.cases(cora[, c("year", "journal", "title", "authors")])

# 2. formatting data
# categorical variables
X.c <- as.matrix(cora[not_missing, c("year", "journal")])

# string variables
X.s <- as.matrix(cora[not_missing, c("title", "authors")])

# keep track of which rows of are in which files
file.num <- rep(c(1, 2, 3), c(200, 200, 160))
```

# Your turn (solution, cont'd)

```
# 3. hyperparameters

# Subjective choices for distortion probability prior
# parameters of a Beta(a,b)
a <- 1
b <- 999

# string distance function example
d <- function(s1, s2) {
  adist(s1, s2) # approximate string distance
}

# steepness parameter
c <- 1

# 4. run the gibbs sampler
lam.gs <- rl.gibbs(file.num = file.num, # file
                  X.s = X.s, X.c = X.c, # data
                  num.gs = 10, # iterations
                  a = a, b = b, # prior params
                  c = c, d = d, # distortion
                  M = nrow(X.s)) # max # latents
```

# Your turn (extra solution)

```
# 5. (extra) evaluation

# estimated pairwise links
est_links_pair <- pairwise(links(lam.gs))

# true pairwise links, only for those included
# get true number
included_idx <- seq_len(sum(not_missing))
names(included_idx) <- cora[not_missing, "id"]
included_pairs <- subset(cora_gold, id1 %in% names(included_idx) & id2 %in% names(included_idx))

# get index for graphmaking
included_pairs <- apply(included_pairs, 2, function(col) {
  names <- as.character(col)
  included_idx[names]
})

# need list of pairs to compare
true_links_pair <- split(included_pairs, seq(nrow(included_pairs)))

# comparison
comparison <- links.compare(est_links_pair, true_links_pair, counts.only = TRUE)

# precision
precision <- comparison$correct/(comparison$incorrect + comparison$correct)

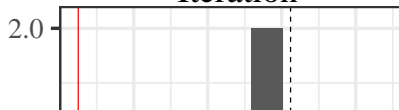
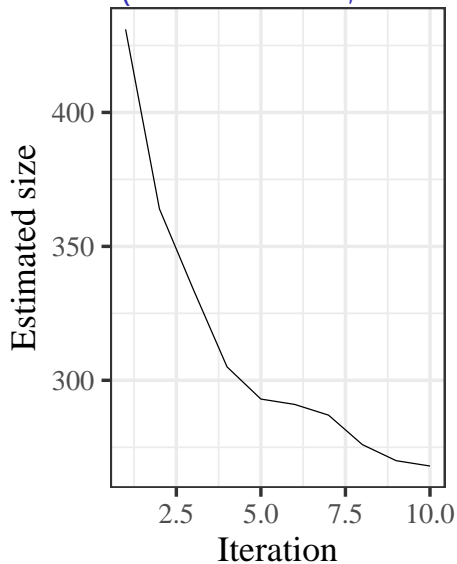
# recall
recall <- comparison$correct/(comparison$correct + comparison$missing)

# results
c(precision, recall)
```

```
## [1] 1.000000000 0.002649919
```



## Your turn (extra solution, cont'd)



# References

- Steorts, R. (2015), Entity Resolution with Empirically Motivated Priors, Bayesian Analysis 10(4), pp. 849–875.
- Steorts et al. (2016). A Bayesian Approach to Graphical Record Linkage and De-duplication, Journal of the American Statistical Association, 111:516, pp.1660-1672.
- Sadinle, M. (2014). Detecting duplicates in a homicide registry using a bayesian partitioning approach. The Annals of Applied Statistics 8(4), pp. 2404–2434
- Zanella et al. (2016). Flexible Models for Microclustering with Applications to Entity Resolution, Advances in Neural Information Processing Systems (NIPS) 29, pp. 1417-1425.
- Miller et al. (2015). The Microclustering Problem: When the Cluster Sizes Don't Grow with the Number of Data Points. NIPS Bayesian Nonparametrics: The Next Generation Workshop Series.