Module X: Blocking

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Reading

- ▶ Binette and Steorts (2020)
- ► Steorts, Ventura, Sadinle, Fienberg (2014)
- ► Murray (2016)

Agenda

- ► Data Cleaning Pipeline
- Blocking
- ► Traditional Blocking
- Probabilistic Blocking

Load R packages

```
knitr::opts_chunk$set(echo = TRUE, fig.width=4, fig.height=
library(RecordLinkage)
library(blink)
```

Data Cleaning Pipeline

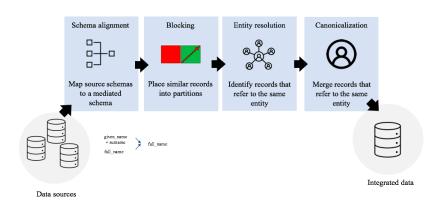


Figure 1: Data cleaning pipeline.

Blocking

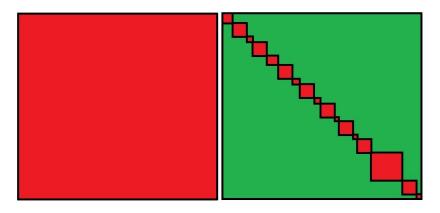


Figure 2: Left: All to all record comparison. Right: Example of resulting blocking partitions.

Blocking

- ▶ Blocking partitions similar records into partitions/blocks.
- ► ER is only performed within each blocks.

Traditional Blocking

- A deterministic (fixed) partition is formed based upon the data.
- ▶ A partition is created by treating certain fields that are thought to be nearly error-free as fixed.
- ▶ Benefits: simple, easy to understand, and fast to implement.
- Downsides: the blocks are treated as error free, which is not usually accurate and can lead to errors in the ER task that cannot be accounted for.

Example: Blocking on date of birth year.

Probabilistic Blocking

► A probability model is used to cluster the data into blocks/partitions.

Example: Fellegi-Sunter (1969), or Locality Sensitive Hashing

Under both blocking approaches, record pairs that do not meet the blocking criteria are automatically classified as non-matches.

Example: Traditional blocking

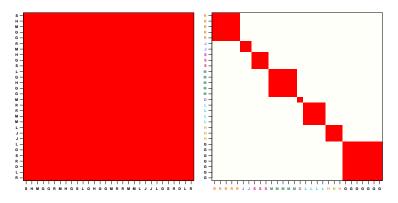


Figure 3: All-to-all record comparisons (left) versus partitioning records into blocks by lastname initial and comparing records only within each partition (right).

Example: RLdata500

```
library(RecordLinkage)
data(RLdata500)
head(RLdata500)
```

```
##
    fname c1 fname c2 lname c1 lname c2 by bm bd
     CARSTEN
## 1
                 <NA>
                         MF.TF.R.
                                   <NA> 1949
                                             7 22
        GF.R.D
                         BAUER.
                                   <NA> 1968 7 27
## 2
                 <NA>
## 3
      ROBERT
                 <NA> HARTMANN
                                   <NA> 1930 4 30
## 4
      STEFAN
                 <NA>
                         WOI.FF
                                   <NA> 1957 9 2
        RALF
                 <NA> KRUEGER
## 5
                                   <NA> 1966 1 13
                 <NA> FRANKE
## 6
     JUERGEN
                                   <NA> 1929 7 4
```

```
# Total number of all to all record comparisons choose(500,2)
```

[1] 124750

[1] 20

```
# Block by last name initial
last_init <- substr(RLdata500[,"lname_c1"], 1, 1)
head(last_init)

## [1] "M" "B" "H" "W" "K" "F"

# Total number of blocks
length(unique(last_init))</pre>
```

5 56 2 6 38 12

##

```
# Total number of records per block
recordsPerBlock <- table(last_init)
head(recordsPerBlock)

## last_init
## A B D E F G</pre>
```

```
# Block sizes can vary
plot(recordsPerBlock,
     cex.axis=0.6, xlab="", ylab="")
  20
```

```
# Total number of records pairs per block
choose(recordsPerBlock, 2)
## last init
        B D E F G
                             H J
##
    Α
                                     K I.
## 10 1540 1 15 703 66 496 28 1035 78 2850
## T
         V
             W
              Z
     1 21 1326 10
##
# Reduction on comparison space
sum(choose(recordsPerBlock, 2))
```

What is the overall dimension reduction form the original space to the reduced space induced by blocking?

Recall the original space of comparisons was

```
choose(500, 2)
```

[1] 124750

We have reduced the number of comparisons to

```
sum(choose(recordsPerBlock, 2))
```

[1] 14805

How do we calculate the reducation ratio?

The reduction ratio is

RR = % comparisons eliminated by blocking.

```
(choose(500, 2) - sum(choose(recordsPerBlock, 2))) /
choose(500, 2)
```

```
## [1] 0.8813226
```

How do we calculate the reducation ratio (via a function)?

```
reduction.ratio <- function(block.labels) {
   n_all_comp = choose(length(block.labels), 2)
   n_block_comp = sum(choose(table(block.labels), 2))

   (n_all_comp - n_block_comp) / n_all_comp
}
reduction.ratio(last_init)</pre>
```

[1] 0.8813226

Pairwise Precision

```
labels = unique(last init)
# Number of matching pairs among blocks
n matches = sapply(labels, function(label) {
  # Records in a given blocks
  records = which(last_init == label)
  # Number of matches in that block
  sum(duplicated(identity.RLdata500[records]))
})
# Total number of pairs
n_pairs = sum(choose(table(last_init), 2))
sum(n matches) / n pairs
## [1] 0.003377237
```

Pairwise Precision

```
precision <- function(block.labels, IDs) {</pre>
  labels = unique(block.labels)
  # Number of matching pairs among blocks
  n_matches = sapply(labels, function(label) {
    records = which(block.labels == label)
    sum(duplicated(IDs[records]))
  })
  # Total number of pairs
  n_pairs = sum(choose(table(block.labels), 2))
  sum(n_matches) / n_pairs
precision(last_init, identity.RLdata500)
```

Pairwise Recall

```
recall <- function(block.labels, IDs) {
  precision(IDs, block.labels)
}</pre>
```

Italian Survey on Household and Wealth (SHIW)

We will now explore a case study to the SHIW install.packages("italy", repos='http://cran.us.r-project.org/li> ## Installing package into '/Users/rebeccasteorts/Library/ ## (as 'lib' is unspecified) ## ## The downloaded binary packages are in /var/folders/bv/xhclmwh90zg08bvwnjvxtrz80000gn/T//Rtmpm #library(devtools) #devtools::install_github("cleanzr/italy") library(italy)

SHIW

- ► The SHIW is a sample survey 383 households conducted by the Bank of Italy every two years.
- The data set is anonymized to remove first and last name (and other sensitive information).
- The following attribute information is available:
 - PARENT (parental status)
 - GENDER
 - ANASC (year of birth)
 - NASCREG (working status)
 - CIT (employment status)
 - ACOM4C (branch of activity)
 - STUDIO (town size)
 - Q (quality of life status)
 - QUAL (whether or not Italian national)
 - SETT (highest educational level obtained)
 - ireq (region of italy)

Explore Data

##

head(italy08) # first year of SHIW

##		Id	PARENI	PEV	ANASC	NASCREG	CII	ACUM4C	SIODIO	Ų
##	1	1040021	1	2	1948	16	1	0	5	1
##	2	1040022	10	2	1952	16	1	0	7	1
##	3	1110521	1	1	1972	20	1	2	5	1
##	4	1110522	3	1	1935	20	1	2	2	3
##	5	1110523	3	2	1941	20	1	2	3	3

## 6	119401	1	1	1941	7	1	0	4	3
head(italy10)	# seco	ond y	year oj	f SHIW				
	. ,	DADENIE	anv.	ANTAGG	MAGGDEG	атш	4 00 14 0	OTTID TO	0

##		id	PARENT	SEX	ANASC	NASCREG	CIT	ACOM4C	STUDIO	Q
##	1	1040021	1	2	1948	16	1	0	5	3
##	2	1040022	11	2	1952	16	1	0	7	1

##	1	1040021	1	2	1948	16	1	0	5 3
##	2	1040022	11	2	1952	16	1	0	7 1
##	3	1110521	1	2	1941	20	1	2	3 3

Reformat Data

```
id08 <- italy08$id
id10 <- italy10$id
id <- c(italy08$id, italy10$id) # combine the id
italy08 <- italy08[-c(1)] # remove the id
italy10 <- italy10[-c(1)] # remove the id
italy <- rbind(italy08, italy10)</pre>
```

Your turn

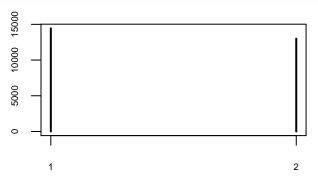
- Construct a blocking criterion for the SHIW data set.
- Provide code to construct the blocks
- Are your blocks well balanced?
- What is the reduction ratio?
- What is the pairwise recall and precision?
- Would you recommend your blocking criterion for an ER task? Why or why not.

```
I will block on gender. Why?
# block by gender
blockByGender <- italy$SEX
recordsPerBlock <- table(blockByGender)
head(recordsPerBlock)

## blockByGender
## 1 2
## 14442 12993</pre>
```

The block sizes are similar. But note, they are still quite large.

```
# Checking block sizes
plot(recordsPerBlock,
cex.axis=0.6, xlab="", ylab="")
```



```
print(rr <- reduction.ratio(blockByGender))</pre>
```

[1] 0.4986234

We have reduced the overall space by rougly 50 percent.

```
precision(blockByGender, id)

## [1] 3.599727e-05

recall(blockByGender, id)
```

```
## [1] 0.9113109
```

This is not an optimal blocking criterion as ideally, we would want both the precision and recall to be close to 1.

```
blockRule <- italy$SEX && italy$ANASC
precision(blockRule, id)

## [1] NaN
recall(blockRule, id)

## [1] 0.9998658</pre>
```