## Module 5: Probabilistic Blocking, Part II

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## Agenda

- ► Data Cleaning Pipeline
- Blocking
- ▶ Probabilistic Blocking
- Locality Sensitive Hashing (LSH)
- Jaccard Similarity
- Shingling
- Putting it together
- Limitations

## Load R packages

```
library(RecordLinkage)
library(blink)
library(knitr)
```

## Data Cleaning Pipeline

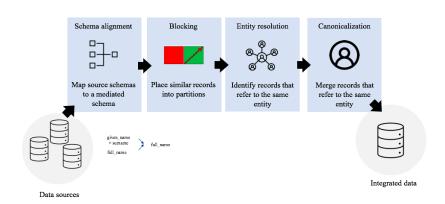


Figure 1: Data cleaning pipeline.

## Blocking

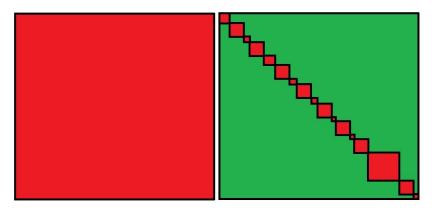


Figure 2: Left: All to all record comparison. Right: Example of resulting blocking partitions.

#### LSH

Locality sensitive hashing (LSH) is a fast method of blocking for record linkage that orginates from the computer science literature.

### Finding similar records

Our goal is to find *similar* records, where the records are assumed to be strings

How do we define similar?

## Jaccard similarity

We will work with the *Jaccard similarity*:

$$Jac(S,T) = \frac{\mid S \cap T \mid}{\mid S \cup T \mid}.$$

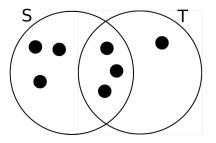


Figure 3: Two sets S and T with Jaccard similarity 3/7. The two sets share 3 elements in common, and there are 7 elements in total.

### How to represent data as sets?

We want to talk about the similarity of our data (records) $\Rightarrow$  we need to compare sets of records!

We can construct a set of short strings from the data

► This is useful because similar datasets will have many common elements (common short strings)

We can do construct these short strings using shingling

# *k*-shingling (how-to)

1. Think of the data set as a string of characters

2. A k-shingle (k-gram) is any sub-string (word) of length k found within the a record of the data set

3. Associate with each data set the set of k-shingles that appear one or more times

### Let's try

Suppose our data set is the string "Hello world", then

▶ the set of 2-shingles is {he, el, II, lo, ow, wo, or, rl, ld}

▶ the set of 3-shingles is {hel, ell, llo, low, owo, wor, orl, rld}

#### Your turn

We have the following two records:

```
# load RL data
data("RLdata500")

# select only 2 records
records <- RLdata500[129:130, c(1,3)]
names(records) <- c("First name", "Last name")

# inspect records
kable(records)</pre>
```

	First name	Last name
129	MICHAEL	
130	MICHAEL	MEYER

# Your turn (continued)

1. Compute the 2-shingles for each record

2. Using Jaccard similarity, how similar are they?

3. What do you learn from this exercise?

#### Your turn solution

- The 2-shingles for the first record are {mi, ic, ch, ha, ae, el, lv, vo, og, ge, el} and for the second are {mi, ic, ch, ha, ae, el, lm, me, ey, ye, er}
- 2. There are 6 items in common {mi, ic, ch, ha, ae, el} and 15 items total {mi, ic, ch, ha, ae, el, lv, vo, og, ge, lm, me, ey, ye, er}, so the Jaccard similarity is  $\frac{6}{15} = \frac{2}{5} = 0.4$
- 3. You should have learned that this is very tedious to do by hand!

### Useful packages/functions in R

We don't want to do this by hand!

Here are some useful packages in R that can help us!

```
library(textreuse) # text reuse/document similarity
library(tokenizers) # shingles
```

# Shingling

We can use the following functions to create k-shingles and calculate Jaccard similarity for our data

```
# get k-shingles
tokenize_character_shingles(x, n)

# calculate jaccard similarity for two sets
jaccard_similarity(a, b)
```

#### Citation Data Set

Research paper headers and citations, with information on authors, title, institutions, venue, date, page numbers and several other fields

```
library(devtools)
install_github("resteorts/cora")
library(cora)
library(ggplot2)
```

# Citation Data Set

data(cora) # load the cora data set
str(cora) # structure of cora

```
##
   'data.frame': 1879 obs. of 16 variables:
                : int 1 2 3 4 5 6 7 8 9 10 ...
##
   $ id
##
   $ title
                : 'noquote' chr "Inganas and M.R" NA NA 1
   $ book title : 'noquote' chr
                                NA NA NA NA ...
##
##
   $ authors
                : 'noquote' chr "M. Ahlskog, J. Paloheime
##
   $ address
                : 'noquote' chr
                                NA NA NA NA ...
                                "1994" "1994" "1994" "19
##
   $ date
                : 'noquote' chr
                : 'noquote' chr
##
   $ year
                                NA NA NA NA ...
```

## \$ editor : 'noquote' chr NA NA NA NA ... "Andersson, J Appl. Phys ## \$ journal : 'noquote' chr ## \$ volume : 'noquote' chr "76" "76" "76" "76" ... ## \$ pages : 'noquote' chr "893" "893" "893" "893"

18 / 24

## \$ publisher : 'noquote' chr NA NA NA NA ...
## \$ institution: 'noquote' chr NA NA NA NA ...
## \$ type : 'noquote' chr NA NA NA NA ...

#### Your turn

Using the title, authors, and journal fields in the cora dataset,

 Get the 3-shingles for each record (hint: use tokenize\_character\_shingles)

 Obtain the Jaccard similarity between each pair of records (hint: use jaccard\_similarity)

## Your turn (solution)

```
# get only the columns we want
# number of records
n <- nrow(cora)
# create id column
dat <- data.frame(id = seq len(n))</pre>
# get columns we want
dat <- cbind(dat, cora[, c("title", "authors", "journal")])</pre>
# 1. paste the columns together and tokenize for each record
shingles <- apply(dat, 1, function(x) {
  # tokenize strings
  tokenize_character_shingles(paste(x[-1], collapse=" "), n = 3)
})
```

## Your turn (solution)

```
# 2. Jaccard similarity between pairs
# empty holder for similarities
jaccard <- expand.grid(record1 = seq_len(n),</pre>
                        record2 = seq len(n))
# don't need to compare the same things twice
jaccard <- jaccard[jaccard$record1 < jaccard$record2,]</pre>
time <- Sys.time() # for timing comparison</pre>
jaccard$similarity <- apply(jaccard, 1, function(pair) {</pre>
  # get jaccard for each pair
  jaccard_similarity(shingles[[pair[1]]], shingles[[pair[2]]])
})
# timing
time <- difftime(Sys.time(), time, units = "secs")</pre>
```

This took took 108.7 seconds  $\approx 1.81$  minutes

### 



# Your turn (solution, cont'd)

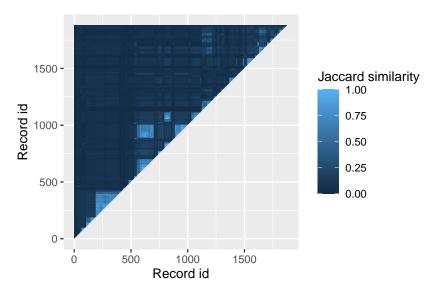


Figure 5: Jaccard similarity for each pair of records. Light blue indicates the two records are more similar and dark blue indicates less similar.

### Summary

For a data set of size n, the number of comparisons we must compute is

$$\frac{n(n-1)}{2}$$
.

For our set of records, we needed to compute 1,764,381 comparisons

For very large data sets, we need something faster (where we filter out records that are not similar).

A better approach for data sets of any realistic size is to use *hashing*, which we will look at next time.