DS5110 HW 3 - Due Feb. 12

Kylie Ariel Bemis 2/1/2019

Instructions

Create a directory with the following structure:

- hw3-your-name/hw3-your-name.Rmd
- hw3-your-name/hw3-your-name.pdf

where hw3-your-name.Rmd is an R Markdown file that compiles to create hw3-your-name.pdf.

Do not include data in the directory. Compress the directory as .zip.

Your solution should include all of the code necessary to answer the problems. All of your code should run (assuming the data is available). All plots should be generated using ggplot2. Missing values and overplotting should be handled appropriately. Axes should be labeled clearly and accurately.

To submit your solution, create a new private post of type "Note" on Piazza, select "Individual Student(s) / Instructor(s)" and type "Instructors", select the folder "hw3", go to Insert->Insert file in the Rich Text Editor, upload your .zip homework solution. Title your note "[hw3 solutions] your name" and post the private note to Piazza. Be sure to post it only to instructors

Part A

Problem 1

Find a dataset that is personally interesting to you. It may be a publicly-available dataset, or a dataset for which you have permission to use and share results. There are many places on to find publicly-available dataset, and simply searching Google for your preferred topic plus "public dataset" may provide many hits. Here some additional resources to get you started:

- US Government datasets (https://catalog.data.gov/dataset)
- Center for Disease Control (CDC) data (https://data.cdc.gov)
- Bureau of Labor Statistics (https://www.bls.gov/data/)
- NASA datasets (https://nssdc.gsfc.nasa.gov)
- World Bank Open Data (https://data.worldbank.org)
- Kaggle Datasets (https://www.kaggle.com/datasets)

This does not have to be the same dataset you will use for your group project.

Import the dataset into R, put it into a tidy format, and print the first ten observations of the dataset.

Problem 2

Step 1: Perform exploratory data analysis on the dataset, using the techniques learned in class. Calculate summary statistics that are of interest to you and create plots using ggplot2 that show your findings.

Step 2: Create an attractive PowerPoint or Keynote slide including your name, a description of your dataset, and your key findings, incorporating any plots and/or tables that are most relevant and interesting. Make sure you cite the source of the data!

Step 3: Export this slide to PDF, and upload it to Piazza as a public Note titled "[mini-poster] your name - dataset name" in the "miniposter" folder, along with a brief description of the dataset by the homework due date.

Part B

Problems 3–5 uses a subset of the DBLP database of bibliographic information on major computer science journals and proceedings, available from https://data.mendeley.com/datasets/3p9w84t5mr. The dataset has been processed to include predictions of the author's genders using the open-source Genderize API. The processed data has been made available in the form of SQL scripts that import the data into a MySQL database. We are primarily interested in the "general" and "authors" tables created by the "main.sql" and "authors.sql" scripts, respectively.

You have three options to load the dataset into R: (1) import the data into a MySQL database, accessed via dbplyr, (2) edit/convert the scripts and import the data into another RDBMS such as SQLite, which is then accessed via dbplyr, or (3) parse the text data in the SQL scripts into R (this is possible but difficult).

If you choose to use MySQL, the README file describes the steps to import the tables into a database, and then use dbplyr with the RMySQL package to work with the data in R.

If you choose to use another RDBMS such as SQLite (which is easier to install, and many *nix operating systems come with it installed already), you will likely need to edit or convert the scripts to be compatible.

One available conversion tool is the mysql2sqlite script available from https://github.com/dumblob/mysql2sqlite. This will convert a MySQL script to a SQLite script.

If you are using a POSIX compliant operating system, you could import the relevant tables into a SQLite database named dblp.db using the following commands in a compatible shell:

```
./mysql2sqlite main.sql | sqlite3 dblp.db
./mysql2sqlite authors.sql | sqlite3 dblp.db
```

and use dbplyr with the RSQLite package to work with the data in R.

Problem 3

Filter the data to include only the authors for whom a gender was predicted as 'male' or 'female' with a probability of 0.95 or greater, and then create a bar plot showing the total number of *distinct* male and female authors published each year. Comment on the visualization.

Problem 4

Still including only the authors for whom a gender was predicted with a probability of 0.95 or greater, create a stacked bar plot showing the *proportions* of distinct male authors vs. distinct female authors published each year. (The stacked bars for each year will sum to one.) Comment on the visualization.

Problem 5

Still including only the authors for whom a gender was predicted with a probability of 0.95 or greater, create a bar plot showing the count of papers published with (1) male first authors and (2) female first authors.

Then create a bar plot showing the count of papers published with (1) no female authors and (2) at least 1 female author. Comment on any similarities and differences between the two bar plots.