zstept-assignment-7

April 20, 2021

1 Assignment 7

In this assignment we will be using an extracted subset of a CDC dataset covering individual US Covid Cases

 $\label{link:https://data.cdc.gov/Case-Surveillance/COVID-19-Case-Surveillance-Public-Use-Data-with-Ge/n8mc-b4w4$

2 Steps to Complete

To complete this assignment, complete each step below. Please turn in both your .ipynb file, and a PDF your notebook

Review the link above describing the full dataset, due to its size I have extracted a subset covering virginia only

- Step 0 Copy and save this notebook. Name it using your GMU netid, netid-assignment-7
 For example (for me) this would be jboone-assignment-7.ipynb
- Step 1 Connect to the dataset provided
 - The data is provided in CSV file located in the ./data subdirectory (see va-cases-0.csv)
- Step 2 Summarize the dataset as prompted
- Step 3 Clean the dataset as prompted
- Step 4 Add the new table to our existing SQLite database (created in the Pandas Part 1 notebook)
- Step 5 Build an index on the new table
- Step 6 Query the database using pandas

```
[1]: # basics
import os
import sys
import numpy as np
import matplotlib.pyplot as plt
import matplotlib.dates as mdates
from mpl_toolkits.axes_grid1 import make_axes_locatable

# needed for direct http requests for COVID data
import io
import requests
```

```
# time
import time

# datetimes
import datetime as dt

# pandas - depending on your environment
# you may need to add these modules to your environment via

# pip install pandas
# # OR
#
# conda install pandas
#
import pandas as pd

# sqlite
import sqlite3 as sql
```

2.1 Step 1 - Connect to the Data Source

```
[2]: filepath = 'assignment-7/data/va-cases-0.csv'
db_dir = '/Users/zachstept/OneDrive - George Mason University/GMU/Spring 2021/
→CDS 302/' # note this should point to a location on your drive
→ of the database resulting from running the pandas-0.ipynb notebook (see week
→11)

# Your Code Here:
va_cases = pd.read_csv(db_dir+filepath)
```

2.2 Step 2 - Summarize

Your Code Here

```
[3]: # Display the number of rows and columns in the dataset
    # Your Code Here
    rows = len(va_cases.axes[0])
    cols = len(va_cases.axes[1])
    print("Number of Rows: " + str(rows))
    print("Number of Columns: " + str(cols))

Number of Rows: 533688
    Number of Columns: 20
[4]: # Display the columns in the dataset
```

```
for col in va_cases.columns:
    print(col)

Unnamed: 0
```

case_month res_state state_fips_code res_county county_fips_code age_group sex race ethnicity case_positive_specimen_interval case_onset_interval process exposure_yn current_status symptom_status hosp_yn icu_yn death_yn underlying_conditions_yn

[5]: # Display the data types for each column
Your Code Here
datatypes = va_cases.dtypes
datatypes

[5]: Unnamed: 0 int64 case month object res_state object state_fips_code float64 res_county object county_fips_code float64 age_group object sex object object race ethnicity object case_positive_specimen_interval float64 case_onset_interval float64 process object exposure_yn object current_status object symptom_status object object hosp_yn icu_yn object

death_yn object underlying_conditions_yn object

dtype: object

```
[6]: # dump the summary contents of the data frame, or print the first few rows # Your Code Here
va_cases.head(10)
```

```
[6]:
        Unnamed: 0 case_month res_state
                                            state_fips_code res_county
               5196
                        2020-02
                                        VA
                                                         51.0
                                                                      NaN
               5197
                                        VA
                                                         51.0
     1
                        2020-02
                                                                      NaN
     2
               5198
                        2020-03
                                        VA
                                                         51.0
                                                                      NaN
     3
               5199
                        2020-03
                                        VA
                                                         51.0
                                                                      NaN
     4
               5200
                        2020-03
                                        VA
                                                         51.0
                                                                      NaN
     5
               5201
                        2020-03
                                        VA
                                                         51.0
                                                                      NaN
     6
               5202
                        2020-03
                                        VΑ
                                                         51.0
                                                                      NaN
     7
                                                                      NaN
               5203
                        2020-03
                                        VA
                                                         51.0
     8
               5204
                        2020-03
                                        VA
                                                         51.0
                                                                      NaN
     9
               5205
                                        VA
                                                         51.0
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                        2020-03
                                       sex race ethnicity \
        county_fips_code age_group
     0
                       NaN
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        case_positive_specimen_interval
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     8
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                                       NaN
                                                                   Missing
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     9
                                       NaN
                                                              0.0 Missing
                                                                                 Missing
                    current_status symptom_status hosp_yn
                                                                icu_yn death_yn
                                                               Missing
        Laboratory-confirmed case
                                        Symptomatic
                                                         Yes
                                                                             NaN
        Laboratory-confirmed case
                                        Symptomatic
                                                         Yes
                                                               Missing
                                                                             NaN
```

```
2 Laboratory-confirmed case
                                Symptomatic
                                                Yes Missing
                                                                    No
3 Laboratory-confirmed case
                                Symptomatic
                                                 No Missing
                                                                    No
4 Laboratory-confirmed case
                                Symptomatic
                                                 No
                                                     Missing
                                                                    No
5 Laboratory-confirmed case
                                Symptomatic
                                                 No Missing
                                                                    No
6 Laboratory-confirmed case
                                Symptomatic
                                                 No Missing
                                                                    No
7 Laboratory-confirmed case
                                Symptomatic
                                                 No Missing
                                                                    No
8 Laboratory-confirmed case
                                Symptomatic
                                                     Missing
                                                 No
                                                                    No
9 Laboratory-confirmed case
                                Symptomatic
                                                 No Missing
                                                                    No
  underlying_conditions_yn
0
                       NaN
1
                       NaN
2
                       NaN
3
                       NaN
4
                       NaN
5
                       NaN
6
                       NaN
7
                       NaN
```

2.3 Step 3 - Perform minor clean-up on the dataset

NaN

NaN

8

9

```
[7]: # display the unique values present for the counties present (use: res_county)
# Your Code Here
va_cases.res_county.unique()
```

```
[7]: array([nan, 'ACCOMACK', 'ALBEMARLE', 'AMHERST', 'ARLINGTON', 'AUGUSTA',
            'BEDFORD', 'BOTETOURT', 'CAROLINE', 'CHESAPEAKE', 'CHESTERFIELD',
            'CULPEPER', 'DANVILLE', 'DINWIDDIE', 'FAIRFAX', 'FLUVANNA',
            'FREDERICK', 'FREDERICKSBURG', 'GLOUCESTER', 'GOOCHLAND',
            'HALIFAX', 'HAMPTON', 'HENRICO', 'HOPEWELL', 'ISLE OF WIGHT',
            'KING GEORGE', 'LEE', 'LOUISA', 'LYNCHBURG', 'MANASSAS',
            'MECKLENBURG', 'MONTGOMERY', 'NEW KENT', 'NEWPORT NEWS', 'NORFOLK',
            'ORANGE', 'PITTSYLVANIA', 'PORTSMOUTH', 'POWHATAN', 'PULASKI',
            'RICHMOND CITY', 'ROANOKE', 'ROANOKE CITY', 'ROCKBRIDGE',
            'ROCKINGHAM', 'SALEM', 'STAFFORD', 'TAZEWELL', 'VIRGINIA BEACH',
            'WARREN', 'WINCHESTER', 'WISE', 'WYTHE', 'YORK', 'ALEXANDRIA',
            'BUCHANAN', 'CAMPBELL', 'FAUQUIER', 'FRANKLIN', 'HENRY',
            'PETERSBURG', 'PRINCE WILLIAM', 'RUSSELL', 'SCOTT', 'SHENANDOAH',
            'WAYNESBORO', 'CARROLL', 'CHARLOTTESVILLE', 'HARRISONBURG', 'PAGE',
            'PRINCE EDWARD', 'SMYTH', 'SPOTSYLVANIA', 'WASHINGTON', 'HANOVER',
            'JAMES CITY', 'SUFFOLK', 'LOUDOUN', 'PRINCE GEORGE', 'STAUNTON'],
           dtype=object)
```

```
[8]: # Note that there are null values for res_county (nan)
      # Drop the rows from the dataframe if the res county field is null
      # Hint: you can also think of this as only keeping the rows where the
      →res_county column is not null
      # Your Code
      va_cases = va_cases[va_cases['res_county'].notna()]
 [9]: # display the unique values present for the counties present (again)
      # to ensure that the nan values have been removed
      # Your Code Here
      va_cases.res_county.unique()
 [9]: array(['ACCOMACK', 'ALBEMARLE', 'AMHERST', 'ARLINGTON', 'AUGUSTA',
             'BEDFORD', 'BOTETOURT', 'CAROLINE', 'CHESAPEAKE', 'CHESTERFIELD',
             'CULPEPER', 'DANVILLE', 'DINWIDDIE', 'FAIRFAX', 'FLUVANNA',
             'FREDERICK', 'FREDERICKSBURG', 'GLOUCESTER', 'GOOCHLAND',
             'HALIFAX', 'HAMPTON', 'HENRICO', 'HOPEWELL', 'ISLE OF WIGHT',
             'KING GEORGE', 'LEE', 'LOUISA', 'LYNCHBURG', 'MANASSAS',
             'MECKLENBURG', 'MONTGOMERY', 'NEW KENT', 'NEWPORT NEWS', 'NORFOLK',
             'ORANGE', 'PITTSYLVANIA', 'PORTSMOUTH', 'POWHATAN', 'PULASKI',
             'RICHMOND CITY', 'ROANOKE', 'ROANOKE CITY', 'ROCKBRIDGE',
             'ROCKINGHAM', 'SALEM', 'STAFFORD', 'TAZEWELL', 'VIRGINIA BEACH',
             'WARREN', 'WINCHESTER', 'WISE', 'WYTHE', 'YORK', 'ALEXANDRIA',
             'BUCHANAN', 'CAMPBELL', 'FAUQUIER', 'FRANKLIN', 'HENRY',
             'PETERSBURG', 'PRINCE WILLIAM', 'RUSSELL', 'SCOTT', 'SHENANDOAH',
             'WAYNESBORO', 'CARROLL', 'CHARLOTTESVILLE', 'HARRISONBURG', 'PAGE',
             'PRINCE EDWARD', 'SMYTH', 'SPOTSYLVANIA', 'WASHINGTON', 'HANOVER',
             'JAMES CITY', 'SUFFOLK', 'LOUDOUN', 'PRINCE GEORGE', 'STAUNTON'],
            dtype=object)
[10]: # Note the column 'Unnamed: O' - this is likely a row identifier from the
      \hookrightarrow original
      # data set - drop this column
      # Your Code Here
      va_cases = va_cases.drop(['Unnamed: 0'], axis = 1)
[11]: | # Drop any row where age_group and sex are are both null
      # Your Code Here
      va_cases = va_cases[va_cases['age_group'].notna() & va_cases['sex'].notna()]
[12]: | # add two new columns that split case_month into year, month
      # Some hints:
      # You may add a new column to a dataframe in more than one way
      # in the lectures, I use the insert() function, but you can also
      # do this directly...like this:
```

```
#
# df['new_column'] = (some function of another column)
#
# Further hints - see the .str and .slice functions in pandas
#
# Your Code Here
va_cases['year'] = pd.DatetimeIndex(va_cases['case_month']).year
va_cases['month'] = pd.DatetimeIndex(va_cases['case_month']).month
```

2.4 Step 4 - Add the new table to the database

```
[14]: # Save the new table to the existing database
# Your Code Here
conn = sql.connect(db_dir+'covid-19.db')
va_cases.to_sql('va_cases', conn, if_exists='replace', index=False)
conn.close()
```

2.5 Step 5 - Create an Index on the New Table

```
[15]: # Create the SQL commands to create the index as strings - index on UID
# Your Code Here
ind0 = '''
drop index if exists va_cases_uid_idx;
'''
ind1 = '''
create index va_cases_uid_idx on va_cases(uid);
''''
```

```
[16]: # Connect to the database and create the new index on the UID attribute by \rightarrow executing the
```

```
# SQL you defined above
# Your Code Here
conn = sql.connect(db_dir+'covid-19.db')
cursor = conn.cursor()
cursor.execute(ind0)
cursor.execute(ind1)
conn.close()
```

2.6 Step 6 - Query the New Database

```
[29]: | # Query - this query will be simple and only involve the new table
      # define two new python variables selected_year, and selected_month (integers)
      # Your Code Here
      selected_year = 2020
      selected_month = 11
      # Now define a python string holding the query
      # The query should select the number of rows in the
      # case details table for each age group during the month, year selected
      # Hint: this is a simple query, one table with a group by on age_group
      # Your Code Here
      query = '''
          SELECT count(UID) as number_of_case, age_group
          FROM va cases
          WHERE month == {0} AND year == {1}
          GROUP BY age_group;
      '''.format(selected_month, selected_year)
      # Now execute and time the query
      # Your Code Here
      conn = sql.connect(db_dir+'covid-19.db')
      t0 = time.time()
      va_cases_by_age_group = pd.read_sql(query, conn)
      t = time.time() - t0
      print('Query executed in {:.3f} seconds'.format(t))
      conn.close()
      # show the result dataframe
      # Your Code Here
      va_cases_by_age_group
```

Query executed in 0.100 seconds

```
[29]: number_of_case age_group
0 5794 0 - 17 years
1 31150 18 to 49 years
2 11447 50 to 64 years
```

3 7280 65+ years 4 28 Missing