

Lung_Cancer_EDA_DB_notebook

February 12, 2023

```
[1]: # Christine Orosco
# Store dataset to a local SQLite DB and conduct EDA
```

```
[2]: # import libraries
import pandas as pd
import numpy as np
import sqlite3
import json
from sqlite3 import Error
import matplotlib.pyplot as plt
import seaborn as sns
```

```
[3]: # Set context to `paper`
sns.set(rc={"font.size":15,"axes.labelsize":10})
#fig, ax = plt.subplots(figsize=(10,10))
sns.set(color_codes=True)
```

0.1 Read the flat file

```
[4]: # Read csv file into df
clean_csv_df = pd.read_csv('~/.cancer_csv.csv')
clean_csv_df.head(5)
```

```
[4]:
```

	Year	State	State_Code	HHS_Region	Age_Range	Benchmark	Locality	\
0	2010	Alabama	AL	4	0-84	2010 Fixed	All	
1	2010	Alaska	AK	10	0-84	2010 Fixed	All	
2	2010	Arizona	AZ	9	0-84	2010 Fixed	All	
3	2010	Arkansas	AR	6	0-84	2010 Fixed	All	
4	2010	California	CA	9	0-84	2010 Fixed	All	

	Observed_Deaths	Population	Expected_Deaths	Potentially_Excess_Deaths	\
0	8879.0	4704052.0	6279.0		2600.0
1	798.0	705520.0	664.0		134.0
2	8917.0	6288617.0	8177.0		740.0
3	5547.0	2864516.0	3893.0		1654.0
4	46065.0	36652988.0	41572.0		4493.0

	Percent_Potentially_Excess_Deaths
0	29.3
1	16.8
2	8.3
3	29.8
4	9.8

0.2 Connect to SQLite DB

```
[5]: # Connect to DB and load table from pandas into sqlite
# Csv file
conn = sqlite3.connect('~/.cancer_data.db')
c1 = conn.cursor()
```

```
[6]: clean_csv_df.to_sql("clean_csv", conn, if_exists="replace")
```

```
[7]: # Query table for data
results = c1.execute('Select * from clean_csv limit 10')
for row in results:
    print(row)
```

```
(0, 2010, 'Alabama', 'AL', 4, '0-84', '2010 Fixed', 'All', 8879.0, 4704052.0,
6279.0, 2600.0, 29.3)
(1, 2010, 'Alaska', 'AK', 10, '0-84', '2010 Fixed', 'All', 798.0, 705520.0,
664.0, 134.0, 16.8)
(2, 2010, 'Arizona', 'AZ', 9, '0-84', '2010 Fixed', 'All', 8917.0, 6288617.0,
8177.0, 740.0, 8.3)
(3, 2010, 'Arkansas', 'AR', 6, '0-84', '2010 Fixed', 'All', 5547.0, 2864516.0,
3893.0, 1654.0, 29.8)
(4, 2010, 'California', 'CA', 9, '0-84', '2010 Fixed', 'All', 46065.0,
36652988.0, 41572.0, 4493.0, 9.8)
(5, 2010, 'Colorado', 'CO', 8, '0-84', '2010 Fixed', 'All', 5879.0, 4959583.0,
5623.0, 263.0, 4.5)
(6, 2010, 'Connecticut', 'CT', 1, '0-84', '2010 Fixed', 'All', 5454.0,
3489199.0, 4723.0, 731.0, 13.4)
(7, 2010, 'Delaware', 'DE', 3, '0-84', '2010 Fixed', 'All', 1631.0, 882190.0,
1212.0, 419.0, 25.7)
(8, 2010, 'District of\nColumbia', 'DC', 3, '0-84', '2010 Fixed', 'All', 855.0,
591408.0, 657.0, 199.0, 23.3)
(9, 2010, 'Florida', 'FL', 4, '0-84', '2010 Fixed', 'All', 34229.0, 18367185.0,
28894.0, 5335.0, 15.6)
```

```
[8]: # Load Website data file into pandas
clean_html_df = pd.read_csv('~/.website_deaths.csv')
clean_html_df.head(5)
```

```
[8]:
```

	State	Avg_Ann_Count	Recent_Trend	AA_Death_Rate_100K(95%CI)	\
0	Kentucky	3307	falling	60.4	
1	West Virginia	1416	falling	53.6	
2	Arkansas	2002	falling	52.8	
3	Mississippi	1880	falling	52.8	
4	Tennessee	4212	falling	51.3	

	AA_CI_lower	AA_CI_upper	Recent_5yr_Death_Rate(95%CI)	\
0	59.5	61.3	-5.6	
1	52.3	54.8	-1.9	
2	51.8	53.9	-3.1	
3	51.7	53.9	-2.6	
4	50.6	52.0	-5.4	

	5yr_Death_Rate_CI_lower	5yr_Death_Rate_CI_upper
0	-7.7	-3.4
1	-2.2	-1.5
2	-3.9	-2.3
3	-3.0	-2.1
4	-7.3	-3.5

```
[9]: # Connect to DB and load table from pandas into sqlite
# website file
c1 = conn.cursor()
clean_html_df.to_sql('clean_html', conn, if_exists='replace')
conn.commit()
```

```
[10]: # Query table for data
results = c1.execute('Select * from clean_html limit 10')
for row in results:
    print(row)
```

```
(0, 'Kentucky', 3307, 'falling', 60.4, 59.5, 61.3, -5.6, -7.7, -3.4)
(1, 'West Virginia', 1416, 'falling', 53.6, 52.3, 54.8, -1.9, -2.2, -1.5)
(2, 'Arkansas', 2002, 'falling', 52.8, 51.8, 53.9, -3.1, -3.9, -2.3)
(3, 'Mississippi', 1880, 'falling', 52.8, 51.7, 53.9, -2.6, -3.0, -2.1)
(4, 'Tennessee', 4212, 'falling', 51.3, 50.6, 52.0, -5.4, -7.3, -3.5)
(5, 'Oklahoma', 2323, 'falling', 50.0, 49.1, 50.9, -2.5, -3.1, -2.0)
(6, 'Indiana', 3843, 'falling', 48.7, 48.0, 49.5, -4.5, -6.3, -2.6)
(7, 'Missouri', 3762, 'falling', 48.7, 48.0, 49.4, -4.6, -6.0, -3.2)
(8, 'Alabama', 2998, 'falling', 48.5, 47.8, 49.3, -3.3, -3.9, -2.7)
(9, 'Louisiana', 2569, 'falling', 47.5, 46.7, 48.4, -4.7, -6.6, -2.7)
```

```
[11]: # # Load API data file into pandas
clean_api_df = pd.read_csv('~/.api_deaths.csv')
conn.commit()
clean_api_df.head(5)
```

```
[11]: State_Code  yearstart  yearend    Strat Value_Type    Value
      0         AK        2010      2014 Overall    AA_Nbr    255.0
      1         AL        2010      2014 Overall    AA_Nbr   3149.0
      2         AR        2010      2014 Overall    AA_Nbr   2118.0
      3         AZ        2010      2014 Overall    AA_Nbr   2736.0
      4         CA        2010      2014 Overall    AA_Nbr  12590.0
```

```
[12]: # Connect to DB and load table from pandas into sqlite
      # website file
      c1 = conn.cursor()
      clean_api_df.to_sql('clean_api', conn, if_exists='replace')
      conn.commit()
```

```
[13]: # Query table for data
      results = c1.execute('Select * from clean_api limit 10')
      for row in results:
          print(row)
```

```
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0)
(1, 'AL', 2010, 2014, 'Overall', 'AA_Nbr', 3149.0)
(2, 'AR', 2010, 2014, 'Overall', 'AA_Nbr', 2118.0)
(3, 'AZ', 2010, 2014, 'Overall', 'AA_Nbr', 2736.0)
(4, 'CA', 2010, 2014, 'Overall', 'AA_Nbr', 12590.0)
(5, 'CO', 2010, 2014, 'Overall', 'AA_Nbr', 1591.0)
(6, 'CT', 2010, 2014, 'Overall', 'AA_Nbr', 1712.0)
(7, 'DC', 2010, 2014, 'Overall', 'AA_Nbr', 241.0)
(8, 'DE', 2010, 2014, 'Overall', 'AA_Nbr', 568.0)
(9, 'FL', 2010, 2014, 'Overall', 'AA_Nbr', 11896.0)
```

0.3 Merge 3 tables into 1

```
[14]: # Merge the 3 tables into 1 large table within Sqlite using SQL JOIN
      c1.execute('create table big_tab as \
      select a.*, b.* , c.* from clean_api a \
      inner join clean_csv b on b.State_Code = a.State_Code \
      inner join clean_html c on c.State = b.State')
```

```
[14]: <sqlite3.Cursor at 0x7fe9ca7cb2d0>
```

```
[15]: conn.commit()
```

```
[44]: # select the data from the table that contains all three tables joined together
      results = c1.execute('select * from big_tab limit 10')
      for row in results:
          print(row)
```

```
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 1, 2010, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 798.0, 705520.0, 664.0, 134.0, 16.8, 33, 'Alaska',
```

```

228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 53, 2011, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 836.0, 717624.0, 692.0, 145.0, 17.3, 33, 'Alaska',
228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 103, 2012, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 830.0, 726076.0, 719.0, 111.0, 13.4, 33, 'Alaska',
228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 154, 2013, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 913.0, 729443.0, 734.0, 179.0, 19.6, 33, 'Alaska',
228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 205, 2014, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 882.0, 730801.0, 749.0, 138.0, 15.6, 33, 'Alaska',
228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(0, 'AK', 2010, 2014, 'Overall', 'AA_Nbr', 255.0, 255, 2015, 'Alaska', 'AK', 10,
'0-84', '2010 Fixed', 'All', 884.0, 732295.0, 769.0, 117.0, 13.2, 33, 'Alaska',
228, 'falling', 36.6, 34.3, 38.9, -7.5, -10.3, -4.6)
(1, 'AL', 2010, 2014, 'Overall', 'AA_Nbr', 3149.0, 0, 2010, 'Alabama', 'AL', 4,
'0-84', '2010 Fixed', 'All', 8879.0, 4704052.0, 6279.0, 2600.0, 29.3, 8,
'Alabama', 2998, 'falling', 48.5, 47.8, 49.3, -3.3, -3.9, -2.7)
(1, 'AL', 2010, 2014, 'Overall', 'AA_Nbr', 3149.0, 52, 2011, 'Alabama', 'AL', 4,
'0-84', '2010 Fixed', 'All', 8992.0, 4724997.0, 6403.0, 2591.0, 28.8, 8,
'Alabama', 2998, 'falling', 48.5, 47.8, 49.3, -3.3, -3.9, -2.7)
(1, 'AL', 2010, 2014, 'Overall', 'AA_Nbr', 3149.0, 102, 2012, 'Alabama', 'AL',
4, '0-84', '2010 Fixed', 'All', 8963.0, 4741919.0, 6545.0, 2418.0, 27.0, 8,
'Alabama', 2998, 'falling', 48.5, 47.8, 49.3, -3.3, -3.9, -2.7)
(1, 'AL', 2010, 2014, 'Overall', 'AA_Nbr', 3149.0, 153, 2013, 'Alabama', 'AL',
4, '0-84', '2010 Fixed', 'All', 9011.0, 4751746.0, 6678.0, 2333.0, 25.9, 8,
'Alabama', 2998, 'falling', 48.5, 47.8, 49.3, -3.3, -3.9, -2.7)

```

```

[17]: # Combine the data from the tables by importing each table into a DF then
      ↪ combine all DFs into 1 large DF using pandas
      # PD.JOIN

```

```

[18]: # Read the three tables from sqlite into seperate DFs
clean_csv_df = pd.read_sql_query("SELECT * from clean_csv", conn,
      ↪ index_col=['State_Code', 'State'])
clean_csv_df.head(5)

```

```

[18]:
      index  Year  HHS_Region  Age_Range  Benchmark  Locality \
State_Code State
AL      Alabama      0   2010           4      0-84  2010 Fixed  All
AK      Alaska      1   2010          10      0-84  2010 Fixed  All
AZ      Arizona      2   2010           9      0-84  2010 Fixed  All
AR      Arkansas      3   2010           6      0-84  2010 Fixed  All
CA      California    4   2010           9      0-84  2010 Fixed  All

      Observed_Deaths  Population  Expected_Deaths \

```

State_Code	State			
AL	Alabama	8879.0	4704052.0	6279.0
AK	Alaska	798.0	705520.0	664.0
AZ	Arizona	8917.0	6288617.0	8177.0
AR	Arkansas	5547.0	2864516.0	3893.0
CA	California	46065.0	36652988.0	41572.0

		Potentially_Excess_Deaths	\
State_Code	State		
AL	Alabama	2600.0	
AK	Alaska	134.0	
AZ	Arizona	740.0	
AR	Arkansas	1654.0	
CA	California	4493.0	

		Percent_Potentially_Excess_Deaths
State_Code	State	
AL	Alabama	29.3
AK	Alaska	16.8
AZ	Arizona	8.3
AR	Arkansas	29.8
CA	California	9.8

```
[19]: clean_html_df = pd.read_sql_query("SELECT * from clean_html", conn,
    ↪index_col='State')
clean_html_df.head(5)
```

	index	Avg_Ann_Count	Recent_Trend	AA_Death_Rate_100K(95%CI)	\
State					
Kentucky	0	3307	falling	60.4	
West Virginia	1	1416	falling	53.6	
Arkansas	2	2002	falling	52.8	
Mississippi	3	1880	falling	52.8	
Tennessee	4	4212	falling	51.3	

	AA_CI_lower	AA_CI_upper	Recent_5yr_Death_Rate(95%CI)	\
State				
Kentucky	59.5	61.3	-5.6	
West Virginia	52.3	54.8	-1.9	
Arkansas	51.8	53.9	-3.1	
Mississippi	51.7	53.9	-2.6	
Tennessee	50.6	52.0	-5.4	

	5yr_Death_Rate_CI_lower	5yr_Death_Rate_CI_upper
State		
Kentucky	-7.7	-3.4
West Virginia	-2.2	-1.5

Arkansas	-3.9	-2.3
Mississippi	-3.0	-2.1
Tennessee	-7.3	-3.5

```
[20]: clean_api_df = pd.read_sql_query("SELECT * from clean_api", conn,
↳ index_col='State_Code')
```

```
[21]: clean_api_df.head(5)
```

```
[21]:
```

	index	yearstart	yearend	Strat	Value_Type	Value
State_Code						
AK	0	2010	2014	Overall	AA_Nbr	255.0
AL	1	2010	2014	Overall	AA_Nbr	3149.0
AR	2	2010	2014	Overall	AA_Nbr	2118.0
AZ	3	2010	2014	Overall	AA_Nbr	2736.0
CA	4	2010	2014	Overall	AA_Nbr	12590.0

0.4 Combine 3 Dataframes into 1

```
[22]: # Combine all three DFs into 1 DF using PD.JOIN
```

```
big_df1 = clean_csv_df.join(clean_api_df, on='State_Code',
↳ lsuffix="State_Code_x", rsuffix="State_Code_y")\
    .join(clean_html_df, on='State', lsuffix="State_x", rsuffix="State_y" )
big_df1.columns
```

```
[22]: Index(['indexState_Code_x', 'Year', 'HHS_Region', 'Age_Range', 'Benchmark',
'Locality', 'Observed_Deaths', 'Population', 'Expected_Deaths',
'Potentially_Excess_Deaths', 'Percent_Potentially_Excess_Deaths',
'indexState_Code_y', 'yearstart', 'yearend', 'Strat', 'Value_Type',
'Value', 'index', 'Avg_Ann_Count', 'Recent_Trend',
'AA_Death_Rate_100K(95%CI)', 'AA_CI_lower', 'AA_CI_upper',
'Recent_5yr_Death_Rate(95%CI)', '5yr_Death_Rate_CI_lower',
'5yr_Death_Rate_CI_upper'],
dtype='object')
```

```
[23]: # drop columns
```

```
big_df1.drop(columns = ['indexState_Code_x', 'HHS_Region', 'index',
↳ 'indexState_Code_y', 'Locality'], axis=1, inplace=True)
```

```
[24]: big_df1.dtypes
```

```
[24]: Year                int64
Age_Range              object
Benchmark              object
Observed_Deaths        float64
Population              float64
```

Expected_Deaths	float64
Potentially_Excess_Deaths	float64
Percent_Potentially_Excess_Deaths	float64
yearstart	int64
yearend	int64
Strat	object
Value_Type	object
Value	float64
Avg_Ann_Count	float64
Recent_Trend	object
AA_Death_Rate_100K(95%CI)	float64
AA_CI_lower	float64
AA_CI_upper	float64
Recent_5yr_Death_Rate(95%CI)	float64
5yr_Death_Rate_CI_lower	float64
5yr_Death_Rate_CI_upper	float64
dtype:	object

```
[25]: # Reset the index to State_Code and Year
big_df1.reset_index(inplace=True)
```

0.5 Create Visualization Functions

```
[ ]:
```

```
[27]: # Create Histogram Plot Function
def Hist_Plot(subtitle, xlab, ser1, xval):
    """Plot histogram"""

    # Main Title
    fig = plt.figure(figsize=(15,10))
    title = fig.suptitle(subtitle, fontsize=14, fontweight="bold")

    fig.subplots_adjust(top=0.88, wspace=0.3)

    # Histogram
    ax1 = fig.add_subplot(1,1,1)
    ax1.set_xlabel(xlab)
    ax1.set_ylabel(ylab)
    sns.histplot(data=ser1, x=xval, bins=20, color='darksalmon')
    plt.show()

    return

# Create Scatter Plot function
def sns_Scatter(subtitle, xlab, ylab, xval, yval, data):
```



```

"""Scatter plot"""
fig = plt.figure(figsize=(15,10))
title = fig.suptitle(subtitle, \
                    fontsize=14, fontweight="bold")
fig.subplots_adjust(top=0.88, wspace=0.3)
# Scatter Plots
ax1 = fig.add_subplot(1,1,1)
ax1.set_xlabel(xlab)
ax1.set_ylabel(ylab)
sns.scatterplot(x = xval, y = yval, data=ser1)
plt.show()
return

# Create a barplot function

def sns_bar(subtitle, ser1, xlab, ylab, xval, yval):
    """Plot a bar plot"""

    fig = plt.figure(figsize=(15,10))
    title = fig.suptitle(subtitle, \
                        fontsize=14, fontweight="bold")
    fig.subplots_adjust(top=0.88, wspace=0.3)
    ax1 = fig.add_subplot(1,1,1)
    ax1.set_xlabel(xlab)
    ax1.set_ylabel(ylab)
    sns.barplot(data=ser1, x=xval, y=yval, label=xlab, ax=ax1)
    plt.show()
    return

# Create a line chart function

def sns_line(subtitle, ser1, xlab, ylab, xval, yval, t_hue):
    """Plot a line plot"""

    fig = plt.figure(figsize=(15,10))
    title = fig.suptitle(subtitle, \
                        fontsize=14, fontweight="bold")
    fig.subplots_adjust(top=0.88, wspace=0.3)
    ax1 = fig.add_subplot(1,1,1)
    ax1.set_xlabel(xlab)
    ax1.set_ylabel(ylab)

    sns.lineplot(data=ser1, x = xval, y = yval, hue=t_hue)
    plt.show()
    return

# Create a regression chart function

```

```
def sns_reg(subtitle, ser1, xlab, ylab, xval, yval):
    """Plot a linear regression"""

    fig = plt.figure(figsize=(15,10))
    title = fig.suptitle(subtitle, \
                        fontsize=14, fontweight="bold")
    fig.subplots_adjust(top=0.88, wspace=0.3)
    ax1 = fig.add_subplot(1,1,1)
    ax1.set_xlabel(xlab)
    ax1.set_ylabel(ylab)

    sns.regplot(data=ser1, x = xval, y = yval)
    plt.show()
    return
```

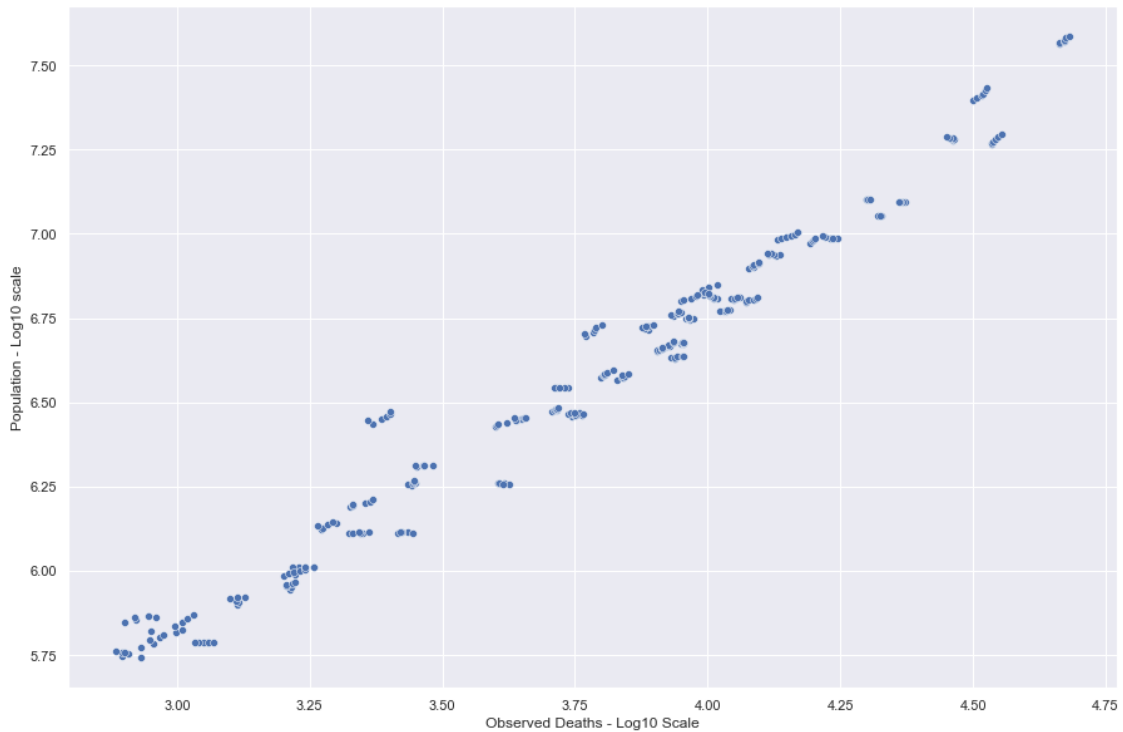
0.6 Create Scatter Plots

```
[28]: # create a DF with only the observed_deaths and Year
ser1 = big_df1[['Observed_Deaths', 'Population']].copy()
```

```
[29]: # Compute log values for better visualization
ser1['Population_log'] = np.log10(ser1[['Population']])
ser1['Observed_Deaths_log'] = np.log10(ser1[['Observed_Deaths']])
```

```
[30]: # Create a Scatter plot for Population and observed number of deaths
subtitle = 'Observed Lung Cancer Deaths per Population growth for each State'
xval = 'Observed_Deaths_log'
yval = 'Population_log'
xlab = 'Observed Deaths - Log10 Scale'
ylab = 'Population - Log10 scale'
sns_Scatter(subtitle, xlab, ylab, xval, yval, ser1)
```

Observed Lung Cancer Deaths per Population growth for each State

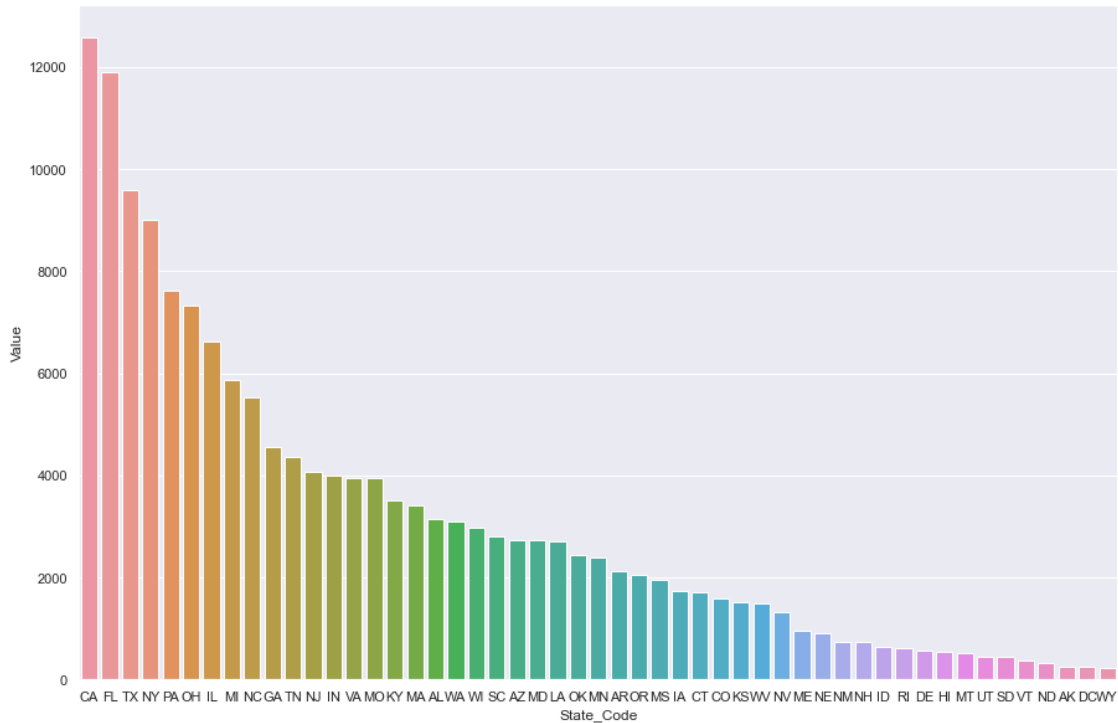


0.7 Create Bar Plots

```
[31]: # Create a bar chart of the Age_adjusted Number per 100K deaths for each state.
      # Drop duplicates because values are repeated for each year. Only need one row
      ↪per state
ser1 = big_df1[['State_Code', 'Value']].sort_values('Value', ascending = False).
      ↪copy()
ser1.drop_duplicates(keep='first', inplace=True)
```

```
[45]: # Define Plot params
ser1 = big_df1[['State_Code', 'Value']].sort_values('Value', ascending = False).
      ↪copy()
subtitle = "Age_Adjusted Lung Cancer Deaths per 100K of population for each
      ↪State"
xlab = "State"
xval = 'State_Code'
ylab = 'Age_Adjusted Annual Lung Cancer Number'
yval = 'Value'
sns_bar(subtitle, ser1, xlab, ylab, xval, yval)
```

Age_Adjusted Lung Cancer Deaths per 100K of population for each State



0.8 Create Line Charts

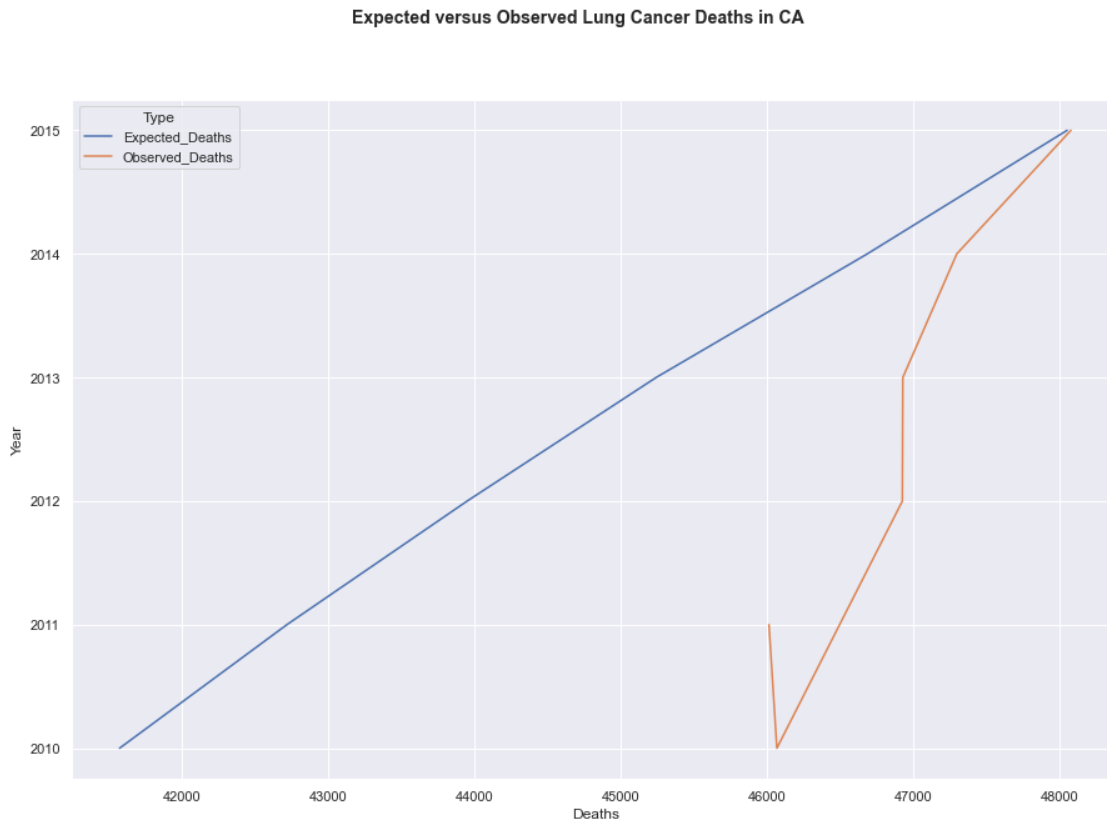
[33]: # Create a line chart showing the observed_deaths vs expected deaths for 5 years for CA.

[34]: # Reshape the data to show two lines representing Observed Deaths and Expected Deaths

```
tmp_df = big_df1[['Expected_Deaths', 'Observed_Deaths', 'State_Code', 'Year']].copy()
ser1 = tmp_df[tmp_df['State_Code'] == 'CA'].copy()
ser1.drop(columns='State_Code', inplace=True)
tmp = pd.melt(ser1, id_vars=['Year'], var_name='Type', value_name='Nbr_Of_Deaths')
tmp.head(5)
```

	Year	Type	Nbr_Of_Deaths
0	2010	Expected_Deaths	41572.0
1	2011	Expected_Deaths	42716.0
2	2012	Expected_Deaths	43948.0
3	2013	Expected_Deaths	45240.0
4	2014	Expected_Deaths	46683.0

```
[46]: # Create line plot showing the two categories of values
ser1 = tmp
subtitle = "Expected versus Observed Lung Cancer Deaths in CA"
xlab = 'Deaths'
xval = 'Nbr_Of_Deaths'
ylab = 'Year'
yval = 'Year'
t_hue = 'Type'
sns_line(subtitle, ser1, xlab, ylab, xval, yval, t_hue)
```



```
[36]: # Show the observed deaths for California because the line graph shows an
      ↪ anomaly
tmp2 = big_df1[['State_Code', 'Year', 'Observed_Deaths']].copy()
tmp2[(tmp2['State_Code'] == 'CA')]
```

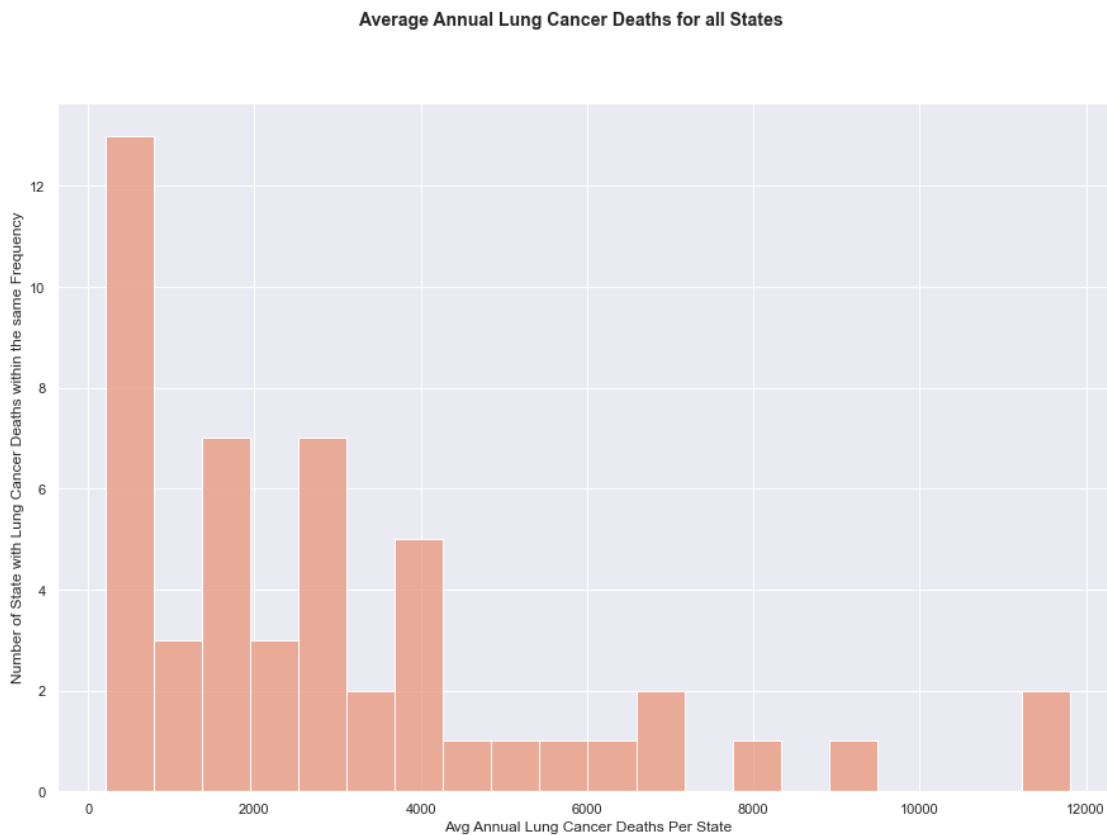
```
[36]:   State_Code  Year  Observed_Deaths
4         CA  2010         46065.0
51        CA  2011         46011.0
106       CA  2012         46923.0
157       CA  2013         46926.0
208       CA  2014         47295.0
```

0.9 Create Histogram

```
[37]: # Create a histogram of the average annaul death rates for all states.
```

```
[48]: # Select a subset for the histogram. Drop duplicate rows as all the years
      ↪reflect the same value
ser1 = big_df1[['State_Code', 'Avg_Ann_Count']].copy()
ser1.drop_duplicates(keep='first', inplace=True)
```

```
[50]: # Define Plot Params
subtitle = "Average Annual Lung Cancer Deaths for all States"
xlab = "Avg Annual Lung Cancer Deaths Per State"
xval = 'Avg_Ann_Count'
ylab = 'Number of State with Lung Cancer Deaths within the same Frequency'
yval = 'Value'
Hist_Plot(subtitle, xlab, ser1, xval)
```

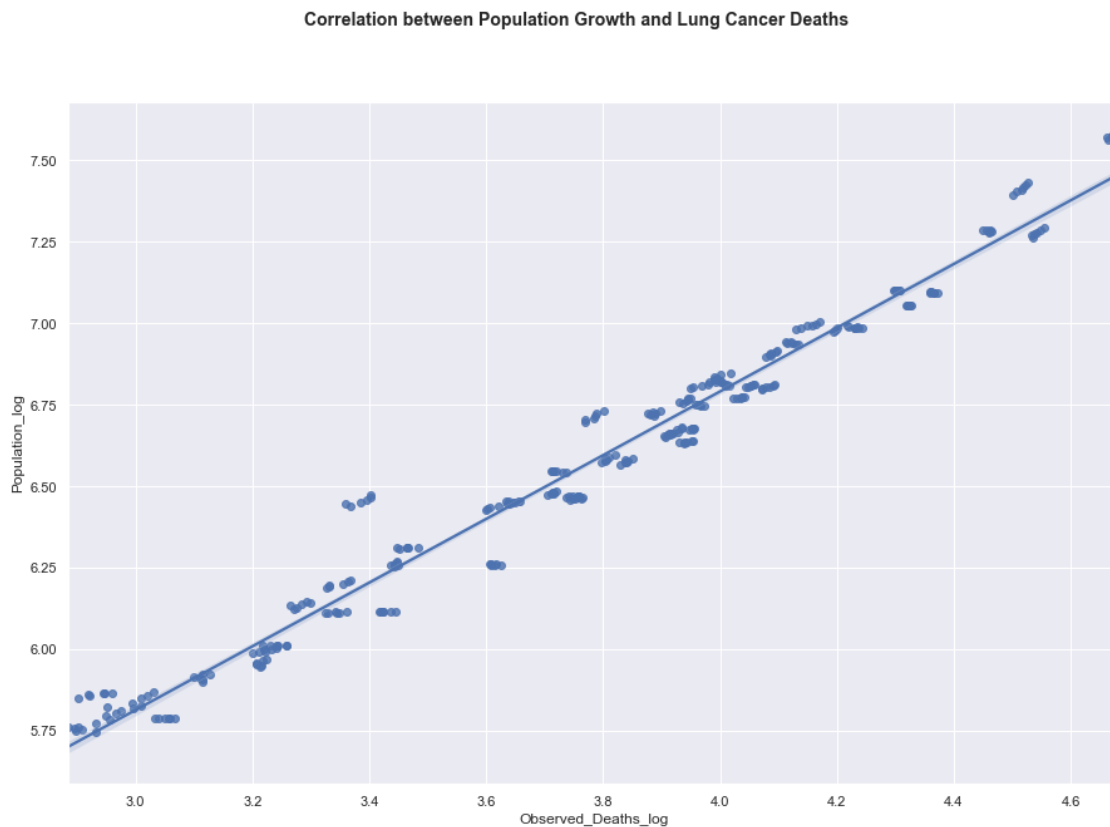


0.10 Create Linear Regression Plot

```
[40]: # Create a linear regression plot to show correlation between population growth ↵  
      ↪and cancer deaths
```

```
[41]: # Create subset for with observed deaths and population for each state  
      # Change the population to log10 for better visualization  
      ser1 = big_df1[['State_Code', 'Population', 'Year', 'Observed_Deaths']].copy()  
      ser1['Population_log'] = np.log10(ser1[['Population']])  
      ser1['Observed_Deaths_log'] = np.log10(ser1[['Observed_Deaths']])
```

```
[42]: # Create Regression Plot  
      subtitle = "Correlation between Population Growth and Lung Cancer Deaths"  
      xval = "Observed_Deaths_log"  
      xlab = 'Number of Lung Cancer Deaths'  
      ylab = 'Population'  
      yval = 'Population_log'  
      sns_reg(subtitle, ser1, xlab, ylab, xval, yval)
```



```
[ ]:
```