TotalDeaths_2020

July 18, 2021

1 Total Deaths 2020

1.0.1 In this analysis, we will analyze the following:

- How different the number of deaths were in the United States for the years 2015 through 2020 and compare them.
- The total number of deaths per jurisdiction, per cause, per year.

1.1 Import libraries

We first need to import Python libraries that will be used with our data analyses and load in the deaths by jurisdiction and cause csv file directly from the source.

[1]:
```
import seaborn as sns
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import re
from IPython.display import display
import statsmodels.api as sm
import statsmodels.formula.api as smf

%matplotlib inline
```

2 Read in .csv file

[2]:
```
data = pd.read_csv("https://data.cdc.gov/api/views/u6jv-9ijr/rows.csv?
↪accessType=DOWNLOAD&bom=true&format=true")
```

[3]:
```
data.head()
```

[3]:

	Jurisdiction	Week Ending Date	State Abbreviation	Year	Week	\
0	Alabama	2015-01-10	AL	2015	1	
1	Alabama	2015-01-10	AL	2015	1	
2	Alabama	2016-01-09	AL	2016	1	
3	Alabama	2016-01-09	AL	2016	1	
4	Alabama	2017-01-07	AL	2017	1	

```
                       Cause Group  Number of Deaths  \
0  Alzheimer disease and dementia             120.0
1  Alzheimer disease and dementia             120.0
2  Alzheimer disease and dementia              76.0
3  Alzheimer disease and dementia              76.0
4  Alzheimer disease and dementia              96.0

                    Cause Subgroup Time Period Suppress Note  \
0  Alzheimer disease and dementia   2015-2019      NaN  NaN
1  Alzheimer disease and dementia   2015-2019      NaN  NaN
2  Alzheimer disease and dementia   2015-2019      NaN  NaN
3  Alzheimer disease and dementia   2015-2019      NaN  NaN
4  Alzheimer disease and dementia   2015-2019      NaN  NaN

   Average Number of Deaths in Time Period  Difference from 2015-2019 to 2020  \
0                                      103                                NaN
1                                      103                                NaN
2                                      103                                NaN
3                                      103                                NaN
4                                      103                                NaN

   Percent Difference from 2015-2019 to 2020                  Type
0                                        NaN  Predicted (weighted)
1                                        NaN            Unweighted
2                                        NaN  Predicted (weighted)
3                                        NaN            Unweighted
4                                        NaN  Predicted (weighted)
```

2.1 Data cleanup

2.1.1 Drop uneeded columns

Lets drop all of the columns that will not be needed in our data analyses. We are interested in Jurisdiction, Year, and Cause Group so we will drop every other column.

[4]:
```
data.drop('Week Ending Date', axis = 1, inplace = True)
data.drop('State Abbreviation', axis = 1, inplace = True)
data.drop('Week', axis = 1, inplace = True)
data.drop('Cause Subgroup', axis = 1, inplace = True)
data.drop('Time Period', axis = 1, inplace = True)
data.drop('Suppress', axis = 1, inplace = True)
data.drop('Note', axis = 1, inplace = True)
data.drop('Average Number of Deaths in Time Period', axis = 1, inplace = True)
data.drop('Difference from 2015-2019 to 2020', axis = 1, inplace = True)
data.drop('Percent Difference from 2015-2019 to 2020', axis = 1, inplace = True)
data.drop('Type', axis = 1, inplace = True)
```

We will check for missing values and handle them.

[5]:
```
data.isnull().sum()
```

[5]:
```
Jurisdiction          0
Year                  0
Cause Group           0
Number of Deaths    134
dtype: int64
```

Lets look at these null values.

[6]:
```
data[data.isna().any(axis=1)]["Jurisdiction"].unique()
```

[6]:
```
array(['Connecticut', 'North Carolina', 'Delaware'], dtype=object)
```

Lets replace these null values with the median from every year for every cause of death and check for null values again.

[7]:
```
data["Number of Deaths"] = data.groupby(["Jurisdiction", "Year", "Cause␣
↪Group"]).transform(lambda x: x.fillna(int(x.median())))
```

[8]:
```
data.isna().sum()
```

[8]:
```
Jurisdiction        0
Year                0
Cause Group         0
Number of Deaths    0
dtype: int64
```

Lets check the data types.

[9]:
```
data.dtypes
```

[9]:
```
Jurisdiction         object
Year                  int64
Cause Group          object
Number of Deaths    float64
dtype: object
```

'Number of Deaths' is of type float and should be of type int so lets change it and check the data types again.

[10]:
```
data = data.astype({'Number of Deaths':'int'})
data.dtypes
```

[10]:
```
Jurisdiction        object
Year                 int64
Cause Group         object
Number of Deaths     int64
dtype: object
```

Lets check for strange values.

[11]:
```
data.describe()
```

[11]:
```
                Year  Number of Deaths
count  356518.000000     356518.000000
mean     2017.770143        157.986413
std         1.877661        683.299929
min      2015.000000         11.000000
25%      2016.000000         23.000000
50%      2018.000000         44.000000
75%      2019.000000         95.000000
max      2021.000000      12433.000000
```

Lets see how many rows have a jurisdiction that is 'United States'.

[12]:
```
display(data[data['Jurisdiction'] == 'United States'].head())
display(data[data['Jurisdiction'] == 'United States'].count())
```

```
       Jurisdiction  Year                     Cause Group  Number of Deaths
29278  United States  2015  Alzheimer disease and dementia              6187
29279  United States  2015  Alzheimer disease and dementia              6187
29280  United States  2016  Alzheimer disease and dementia              5155
29281  United States  2016  Alzheimer disease and dementia              5155
29282  United States  2017  Alzheimer disease and dementia              5844

Jurisdiction        8788
Year                8788
Cause Group         8788
Number of Deaths    8788
dtype: int64
```

We need specific jurisdictions so lets drop all of the rows that contain 'United States' in the 'Jurisdiction' column and check if they are gone.

[13]:
```
index = data[(data['Jurisdiction'] == 'United States')].index
data.drop(index, inplace = True)
display(data[data['Jurisdiction'] == 'United States'].count())
```

```
Jurisdiction        0
Year                0
Cause Group         0
Number of Deaths    0
dtype: int64
```

Before we analyze this data, Python likes it better when variables do not have spaces, so lets get rid of the spaces.

[14]:
```
data.columns = ['Jurisdiction', 'Year', 'CauseGroup', 'NumberOfDeaths']
data.head()
```

[14]:

	Jurisdiction	Year	CauseGroup	NumberOfDeaths
0	Alabama	2015	Alzheimer disease and dementia	120
1	Alabama	2015	Alzheimer disease and dementia	120
2	Alabama	2016	Alzheimer disease and dementia	76
3	Alabama	2016	Alzheimer disease and dementia	76
4	Alabama	2017	Alzheimer disease and dementia	96

Since the data looks good, we can now start analyzing the dataset.

2.2 Dataset Analyses

Dataset: Weekly_counts_of_death_by_jurisdiction_and_cause_of_death analyses

2.2.1 Seperate data by year

The data needs to be seperated by year so we can analyze deaths per year. We will create a new dataframe for every year in the data called 'df_year' replacing year with the actual year.

[15]:
```
for year in data['Year'].unique():
    globals()['df_%s' % year] = data[data.Year == year].reset_index(drop = True)
```

Lets sum the total number of deaths for all years and total number of deaths per year. We will store the values for total deaths in a variable called 'totalDeaths_all' and store each years total deaths in a seperate variable called 'sum_year' replacing year with the actual year.

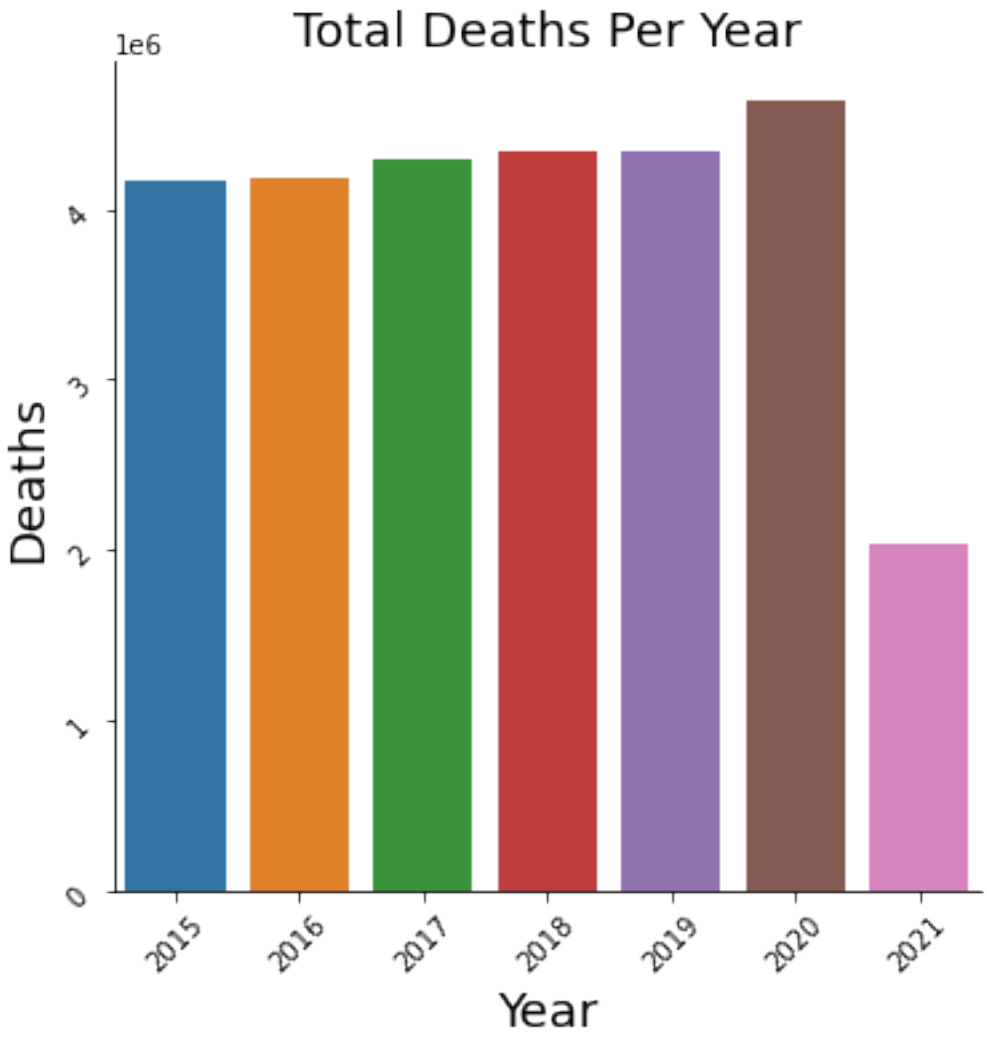
[16]:
```
sum_years = data.groupby([data.Year]).sum()

totalDeaths_all = int(sum(sum_years['NumberOfDeaths']))

i = 0
for year in sum_years.index:
    globals()['sum_%s' % year] = int(sum_years.iloc[i])
    i+=1
```

We will now create a dataframe of years and deaths to analyze and compare and visualize it with a bar plot.

[17]:
```
years = data['Year'].unique()

years_total = []
for year in years:
    years_total.append(str(year))
years_total.append('Total')

deaths = [item for item in sum_years['NumberOfDeaths']]
deaths.append(totalDeaths_all)

rate = []
```

```
for i in range(len(deaths)-1):
    rate.append(round(deaths[i]/totalDeaths_all, 5))

rateSum = sum(rate)
rate.append(rateSum)

df_deaths = np.array([years_total, deaths, rate]).T
df_deaths = pd.DataFrame(df_deaths)
df_deaths.columns = ['Year', 'Deaths', 'Rate']

df_ndeaths = df_deaths

df_ndeaths.drop(df_ndeaths.tail(1).index,inplace=True) # drop first n rows

df_ndeaths = df_ndeaths.astype({'Year':'int', 'Deaths':'int', 'Rate':'float'})

display(df_ndeaths)

sns.catplot(x = 'Year', y = 'Deaths', kind = 'bar', data = df_ndeaths)
plt.title("Total Deaths Per Year", size=18)
plt.xlabel("Year", size=18)
plt.ylabel("Deaths", size=18)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()
```

	Year	Deaths	Rate
0	2015	4169618	0.14884
1	2016	4185822	0.14942
2	2017	4301084	0.15353
3	2018	4343758	0.15506
4	2019	4341440	0.15497
5	2020	4637121	0.16553
6	2021	2035063	0.07264



2.2.2 Deaths by cause and year

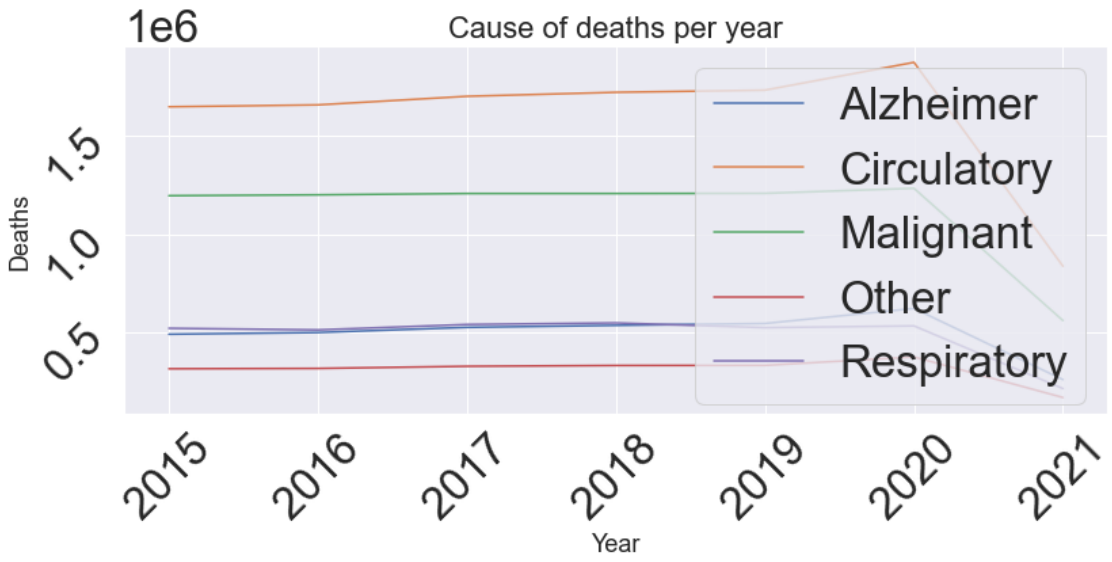
Lets analyze deaths by cause and year. We will seperate each year by cause, create a dataframe and plot the results.

[29]:
```
# Get Alzheimer deaths per year and store in a variable called␣
↪'deathsAlzheimer_year' where year is the actual year
for year in data['Year'].unique():
    globals()['deathsAlzheimer_%s' % year] = int(data.query(f"CauseGroup ==␣
↪'Alzheimer disease and dementia' and Year == {year}")['NumberOfDeaths'].
↪sum())

# Get Malignant deaths per year and store in a variable called␣
↪'deathsMalignant_year' where year is the actual year                    ␣
↪
```

```
for year in data['Year'].unique():
    globals()['deathsMalignant_%s' % year] = int(data.query(f"CauseGroup ==␣
 ↪'Malignant neoplasms' and Year == {year}")['NumberOfDeaths'].sum())

# Get Respiratory deaths per year and store in a variable called␣
 ↪'deathsRespiratory_year' where year is the actual year                    ␣
 ↪
for year in data['Year'].unique():
    globals()['deathsRespiratory_%s' % year] = int(data.query(f"CauseGroup ==␣
 ↪'Respiratory diseases' and Year == {year}")['NumberOfDeaths'].sum())

# Get Circulatory deaths per year and store in a variable called␣
 ↪'deathsCirculatory_year' where year is the actual year                    ␣
 ↪
for year in data['Year'].unique():
    globals()['deathsCirculatory_%s' % year] = int(data.query(f"CauseGroup ==␣
 ↪'Circulatory diseases' and Year == {year}")['NumberOfDeaths'].sum())

# Get Other deaths per year and store in a variable called 'deathsOther_year'␣
 ↪where year is the actual year                                             ␣
 ↪
for year in data['Year'].unique():
    globals()['deathsOther_%s' % year] = int(data.query(f"CauseGroup == 'Other␣
 ↪select causes' and Year == {year}")['NumberOfDeaths'].sum())

yearCauseDeaths = data.groupby([data.Year, data.CauseGroup]).sum().reset_index()

myDict = {}
for i in range(len(yearCauseDeaths)):
    if myDict == {}:
        myDict = {yearCauseDeaths.iloc[i,0]:[yearCauseDeaths.iloc[i,-1]]}
    else:
        if yearCauseDeaths.iloc[i,0] in myDict.keys():
            myDict[yearCauseDeaths.iloc[i,0]].append(yearCauseDeaths.iloc[i,-1])
        else:
            myDict.update({yearCauseDeaths.iloc[i,0]:[yearCauseDeaths.
 ↪iloc[i,-1]]})

colNames = yearCauseDeaths["CauseGroup"].unique()
col_names = []

for item in colNames:
    col_names.append(item.split(" ")[0])

df_deathsCausePerYear = pd.DataFrame(myDict.values(), columns = col_names)
df_deathsCausePerYear
```

```
years = myDict.keys()

df_deathsCausePerYear.insert(0, "Year", years)

display(df_deathsCausePerYear)

fig = plt.gcf()
fig.set_size_inches(13, 5)
plt.plot(df_deathsCausePerYear.Year, "Alzheimer", data = df_deathsCausePerYear)
plt.plot(df_deathsCausePerYear.Year, "Circulatory", data =␣
 ↪df_deathsCausePerYear)
plt.plot(df_deathsCausePerYear.Year, "Malignant", data = df_deathsCausePerYear)
plt.plot(df_deathsCausePerYear.Year, "Other", data = df_deathsCausePerYear)
plt.plot(df_deathsCausePerYear.Year, "Respiratory", data =␣
 ↪df_deathsCausePerYear)
plt.title("Cause of deaths per year", size = 22)
plt.xlabel("Year", size = 18)
plt.ylabel("Deaths", size = 18)
plt.xticks(rotation = 45)
plt.yticks(rotation = 45)
plt.legend()
plt.show()
```

	Year	Alzheimer	Circulatory	Malignant	Other	Respiratory
0	2015	490130	1649144	1196044	314026	520274
1	2016	498712	1658822	1200410	315756	512122
2	2017	525012	1702588	1207174	326626	539684
3	2018	535206	1722912	1207218	330710	547712
4	2019	544754	1733832	1208100	331254	523500
5	2020	621421	1875620	1233553	374082	532445
6	2021	258620	836478	559593	167456	212916



From the results of Deaths Per Cause Per Year, we can see that the number of deaths increase every year in every cause except respiratory. Respiratory deaths are the only cause that the number of deaths decrease and increase but it stays consistent. We can visualize that circulatory deaths are the number one cause of deaths every year followed by malignant and then alzheimer and respiratory are about the same.

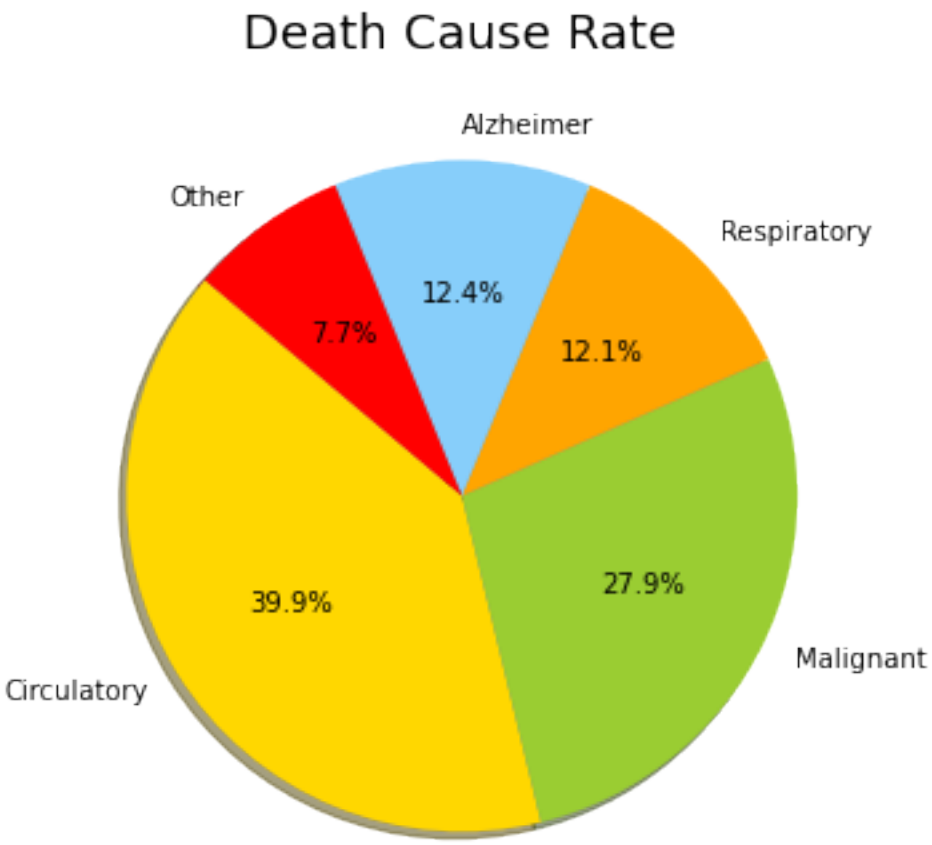
2.2.3 Total deaths by cause

Now, lets analyze deaths by cause, create a dataframe and plot the results in a pie plot.

[19]:
```
deaths_alzheimer = int(data.loc[data['CauseGroup'] == 'Alzheimer disease and␣
↪dementia', 'NumberOfDeaths'].sum())
deaths_malignant = int(data.loc[data['CauseGroup'] == 'Malignant neoplasms',␣
↪'NumberOfDeaths'].sum())
deaths_respiratory = int(data.loc[data['CauseGroup'] == 'Respiratory diseases',␣
↪'NumberOfDeaths'].sum())
deaths_circulatory = int(data.loc[data['CauseGroup'] == 'Circulatory diseases',␣
↪'NumberOfDeaths'].sum())
deaths_other = int(data.loc[data['CauseGroup'] == 'Other select causes',␣
↪'NumberOfDeaths'].sum())

array_circulatory = ['Circulatory', deaths_circulatory,␣
↪round(deaths_circulatory/totalDeaths_all, 4)]
array_malignant = ['Malignant', deaths_malignant, round(deaths_malignant/
↪totalDeaths_all, 4)]
array_respiratory = ['Respiratory', deaths_respiratory,␣
↪round(deaths_respiratory/totalDeaths_all, 4)]
```

```
array_alzheimer = ['Alzheimer', deaths_alzheimer, round(deaths_alzheimer/
 ↪totalDeaths_all, 4)]
array_other = ['Other', deaths_other, round(deaths_other/totalDeaths_all, 4)]

df_cause = pd.DataFrame([array_circulatory, array_malignant, array_respiratory,␣
 ↪array_alzheimer, array_other])

df_cause.columns = ['Cause', 'Deaths', 'Rate']

display(df_cause)

fig, ax = plt.subplots(figsize = (5, 6), subplot_kw = dict(aspect = "equal"))

labels = df_cause['Cause']
sizes = df_cause['Rate']
colors = ['gold', 'yellowgreen', 'orange', 'lightskyblue', 'red']

# Plot
plt.pie(sizes, labels = labels, colors = colors,
autopct='%1.1f%%', shadow = True, startangle=140)

plt.axis('equal')
plt.title('Death Cause Rate', size=18)
plt.show()
```

	Cause	Deaths	Rate
0	Circulatory	11179396	0.3991
1	Malignant	7812092	0.2789
2	Respiratory	3388653	0.1210
3	Alzheimer	3473855	0.1240
4	Other	2159910	0.0771



The pie plot above shows the total percentage of deaths per cause for years 2015-2020. The visualization of the pie plot also shows circulatory deaths as the number one cause of deaths at about 39% and malignant deaths at 28% which with both of these together adds up to almost 70%.

2.2.4 Death rate by cause and year

[20]:
```
rateCause_2015 = ['2015', round(deathsAlzheimer_2015/sum_2015, 4),␣
↪round(deathsMalignant_2015/sum_2015, 4),
                  round(deathsRespiratory_2015/sum_2015, 4),␣
↪round(deathsCirculatory_2015/sum_2015, 4),
                  round(deathsOther_2015/sum_2015, 4)]
rateCause_2016 = ['2016', round(deathsAlzheimer_2016/sum_2016, 4),␣
↪round(deathsMalignant_2016/sum_2016, 4),
                  round(deathsRespiratory_2016/sum_2016, 4),␣
↪round(deathsCirculatory_2016/sum_2016, 4),
                  round(deathsOther_2016/sum_2016, 4)]
```

```
rateCause_2017 = ['2017', round(deathsAlzheimer_2017/sum_2017, 4),␣
 ↪round(deathsMalignant_2017/sum_2017, 4),
                   round(deathsRespiratory_2017/sum_2017, 4),␣
 ↪round(deathsCirculatory_2017/sum_2017, 4),
                   round(deathsOther_2017/sum_2017, 4)]
rateCause_2018 = ['2018', round(deathsAlzheimer_2018/sum_2018, 4),␣
 ↪round(deathsMalignant_2018/sum_2018, 4),
                   round(deathsRespiratory_2018/sum_2018, 4),␣
 ↪round(deathsCirculatory_2018/sum_2018, 4),
                   round(deathsOther_2018/sum_2018, 4)]
rateCause_2019 = ['2019', round(deathsAlzheimer_2019/sum_2019, 4),␣
 ↪round(deathsMalignant_2019/sum_2019, 4),
                   round(deathsRespiratory_2019/sum_2019, 4),␣
 ↪round(deathsCirculatory_2019/sum_2019, 4),
                   round(deathsOther_2019/sum_2019, 4)]
rateCause_2020 = ['2020', round(deathsAlzheimer_2020/sum_2020, 4),␣
 ↪round(deathsMalignant_2020/sum_2020, 4),
                   round(deathsRespiratory_2020/sum_2020, 4),␣
 ↪round(deathsCirculatory_2020/sum_2020, 4),
                   round(deathsOther_2020/sum_2020, 4)]

df_rateCause = pd.DataFrame([rateCause_2015, rateCause_2016, rateCause_2017,␣
 ↪rateCause_2018, rateCause_2019, rateCause_2020])
df_rateCause.columns = ['Year', 'Alzheimer', 'Malignant', 'Respiratory',␣
 ↪'Circulatory', 'Other']

display(df_rateCause)
```

	Year	Alzheimer	Malignant	Respiratory	Circulatory	Other
0	2015	0.1175	0.2868	0.1248	0.3955	0.0753
1	2016	0.1191	0.2868	0.1223	0.3963	0.0754
2	2017	0.1221	0.2807	0.1255	0.3959	0.0759
3	2018	0.1232	0.2779	0.1261	0.3966	0.0761
4	2019	0.1255	0.2783	0.1206	0.3994	0.0763
5	2020	0.1340	0.2660	0.1148	0.4045	0.0807

2.2.5 Death trends per cause

Lets analyze the death trends per cause and plot them seperately.

[21]:
```
df_deathsCausePerYear
```

[21]:

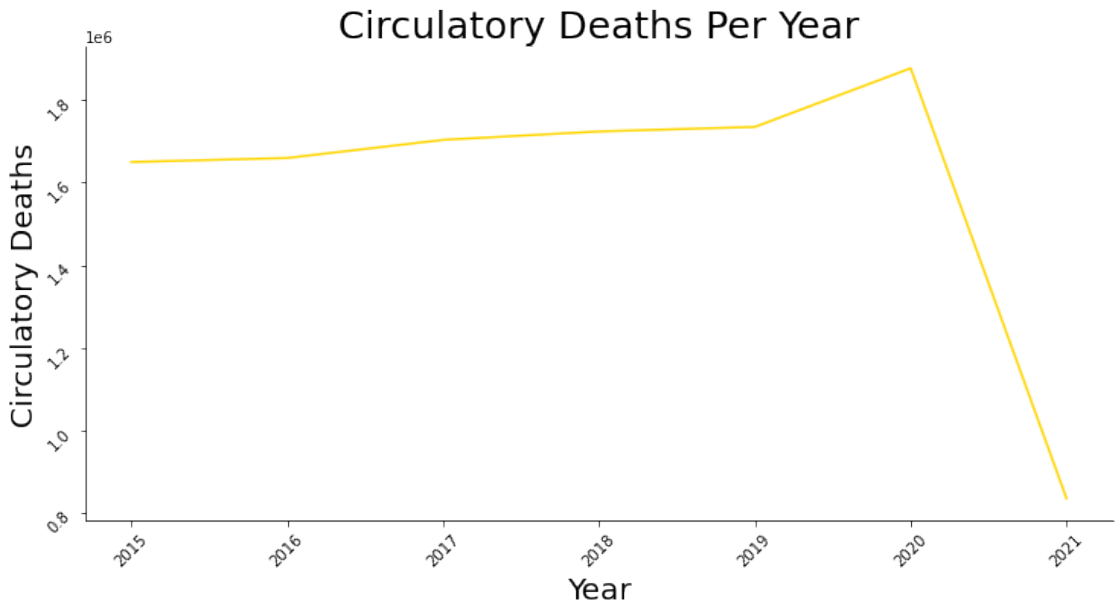
	Year	Alzheimer	Circulatory	Malignant	Other	Respiratory
0	2015	490130	1649144	1196044	314026	520274
1	2016	498712	1658822	1200410	315756	512122
2	2017	525012	1702588	1207174	326626	539684
3	2018	535206	1722912	1207218	330710	547712

```
4  2019      544754       1733832     1208100  331254       523500
5  2020      621421       1875620     1233553  374082       532445
6  2021      258620        836478      559593  167456       212916
```

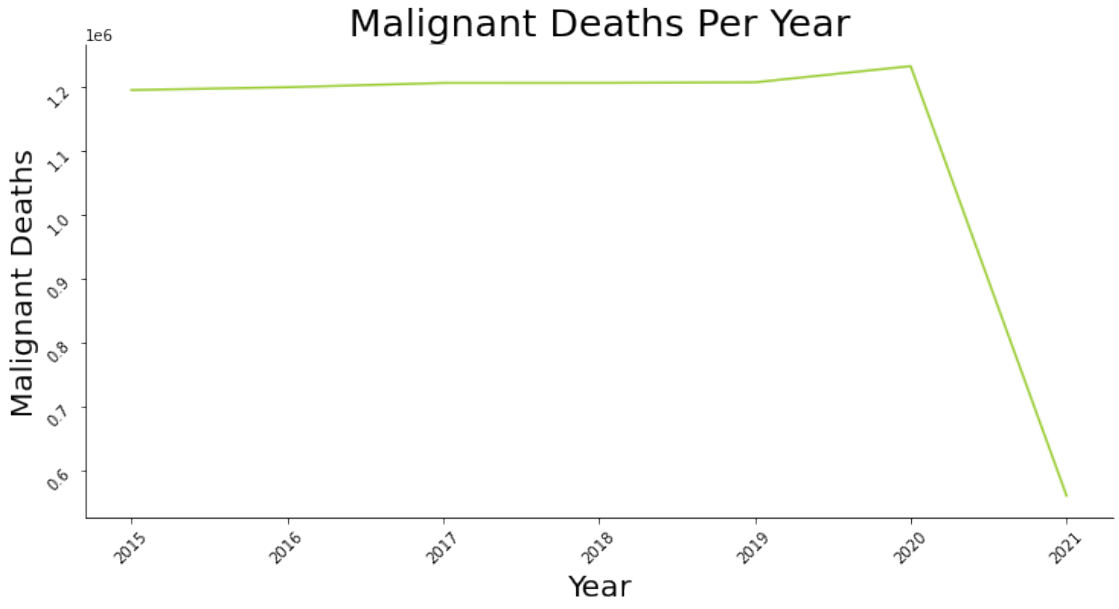
[22]:
```
data_circulatory = pd.DataFrame([df_deathsCausePerYear['Year'],␣
 ↪df_deathsCausePerYear['Circulatory']])
data_circulatory = data_circulatory.transpose().set_index('Year')
display(data_circulatory)
# Draw Plot
sns.relplot(x=data_circulatory.index, y='Circulatory', kind='line', height=5,␣
 ↪aspect=2, color='gold', data=data_circulatory)
plt.title("Circulatory Deaths Per Year", size=25)
plt.xlabel("Year", size=20)
plt.ylabel("Circulatory Deaths", size=20)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()

data_malignant = pd.DataFrame([df_deathsCausePerYear['Year'],␣
 ↪df_deathsCausePerYear['Malignant']])
data_malignant = data_malignant.transpose().set_index('Year')
display(data_malignant)
# Draw Plot
sns.relplot(x=data_malignant.index, y='Malignant', kind = 'line', height=5,␣
 ↪aspect=2, color='yellowgreen', data=data_malignant)
plt.title("Malignant Deaths Per Year", size=25)
plt.xlabel("Year", size=20)
plt.ylabel("Malignant Deaths", size=20)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()

data_respiratory = pd.DataFrame([df_deathsCausePerYear['Year'],␣
 ↪df_deathsCausePerYear['Respiratory']])
data_respiratory = data_respiratory.transpose().set_index('Year')
display(data_respiratory)
# Draw Plot
sns.relplot(x=data_respiratory.index, y='Respiratory', kind = 'line', height=5,␣
 ↪aspect=2, color='orange', data=data_respiratory)
plt.title("Respiratory Deaths Per Year", size=25)
plt.xlabel("Year", size=20)
plt.ylabel("Respiratory Deaths", size=20)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()
```

```
data_alzheimer = pd.DataFrame([df_deathsCausePerYear['Year'],␣
 ↪df_deathsCausePerYear['Alzheimer']])
data_alzheimer = data_alzheimer.transpose().set_index('Year')
display(data_alzheimer)
# Draw Plot
sns.relplot(x=data_alzheimer.index, y='Alzheimer', kind = 'line', height=5,␣
 ↪aspect=2, color='lightskyblue', data=data_alzheimer)
plt.title("Alzheimer Deaths Per Year", size=25)
plt.xlabel("Year", size=20)
plt.ylabel("Alzheimer Deaths", size=20)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()

data_other = pd.DataFrame([df_deathsCausePerYear['Year'],␣
 ↪df_deathsCausePerYear['Other']])
data_other = data_other.transpose().set_index('Year')
display(data_other)
# Draw Plot
sns.relplot(x=data_other.index, y='Other', kind = 'line', height=5, aspect=2,␣
 ↪color='red', data=data_other)
plt.title("Other Deaths Per Year", size=25)
plt.xlabel("Year", size=20)
plt.ylabel("Other Deaths", size=20)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()
```

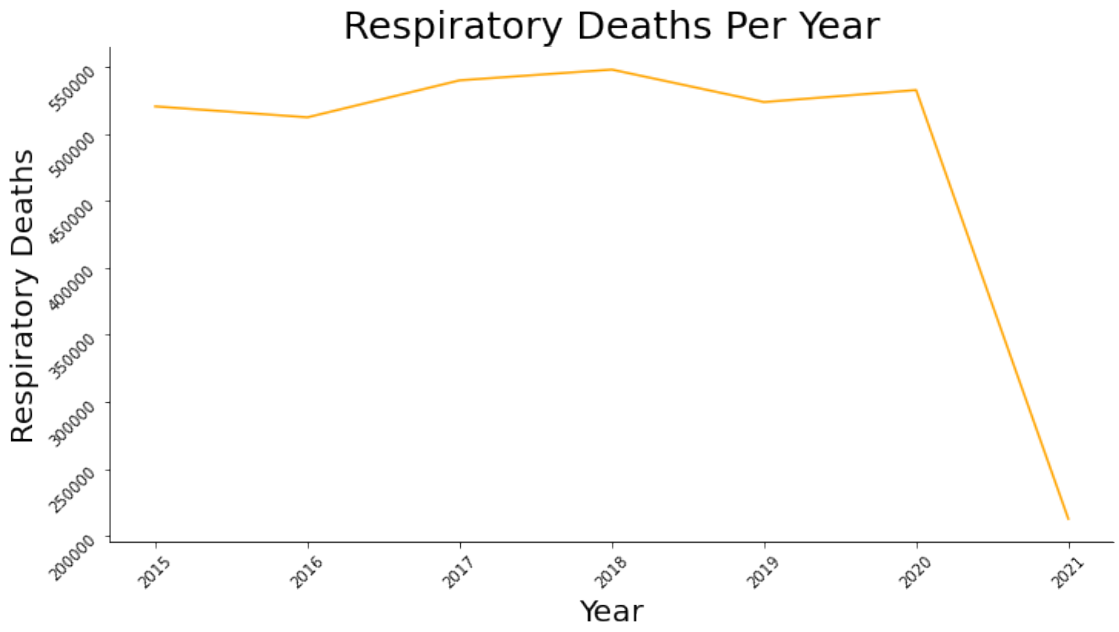
Year	Circulatory
2015	1649144
2016	1658822
2017	1702588
2018	1722912
2019	1733832
2020	1875620
2021	836478



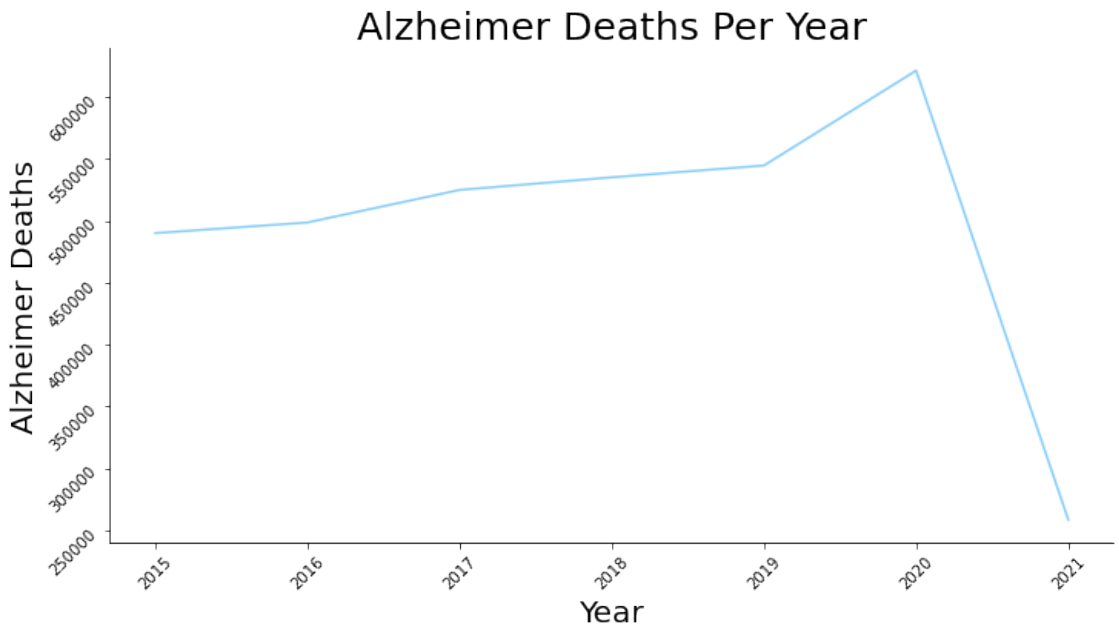
Year	Malignant
2015	1196044
2016	1200410
2017	1207174
2018	1207218
2019	1208100
2020	1233553
2021	559593



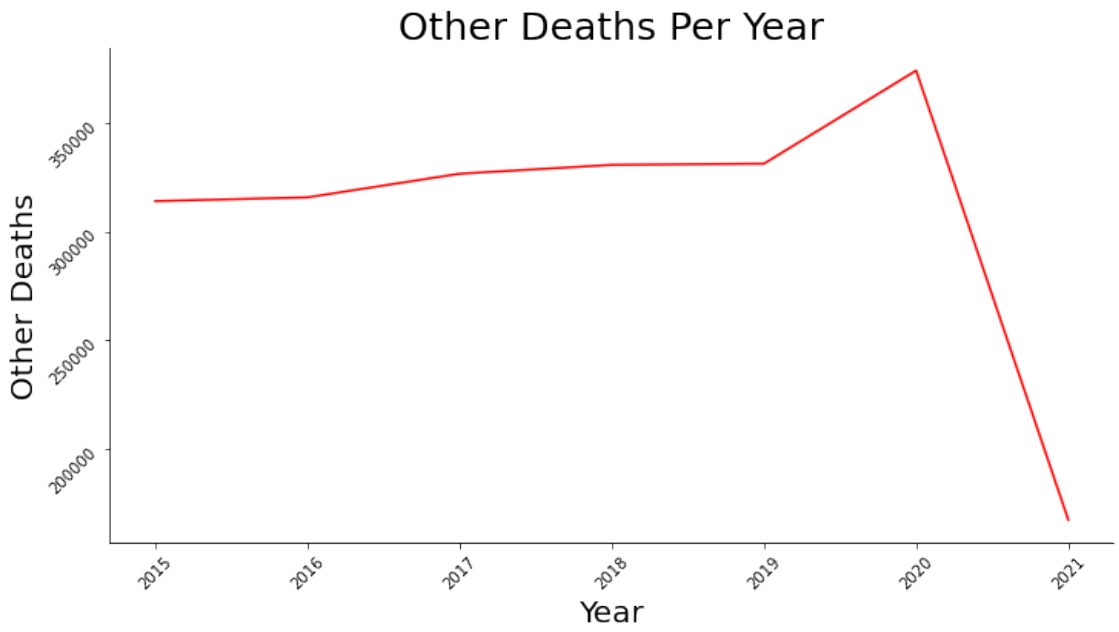
Year	Respiratory
2015	520274
2016	512122
2017	539684
2018	547712
2019	523500
2020	532445
2021	212916



Year	Alzheimer
2015	490130
2016	498712
2017	525012
2018	535206
2019	544754
2020	621421
2021	258620



Year	Other
2015	314026
2016	315756
2017	326626
2018	330710
2019	331254
2020	374082
2021	167456



According to the visualizations above, you can see the death trends of every category increase every year except respiratory. This is interesting because in the year 2020, most of the country had lockdown restrictions for the COVID-19 pandemic.

2.2.6 Total deaths by year and jurisdiction

Lets use Pandas groupby function to get the total deaths by year and jurisdiction.

[23]:
```
yearJurisdiction_df = data.groupby(['Year', 'Jurisdiction']).sum()
yearJurisdiction_df
```

[23]:

Year	Jurisdiction	NumberOfDeaths
2015	Alabama	77032
	Alaska	2254
	Arizona	79724
	Arkansas	47900
	California	411494
…		…
2021	Virginia	53252
	Washington	41044
	West Virginia	9254
	Wisconsin	36857
	Wyoming	1471

[371 rows x 1 columns]

2.2.7 Total deaths by jurisdiction

Now, lets get the total deaths by jurisdiction for all 6 years without the year and sort them. We will drop year index.

[24]:
```
jurisdiction_deaths = data.groupby('Jurisdiction', as_index=False).sum()
jurisdiction_deaths.drop('Year', axis = 1, inplace = True)
display(jurisdiction_deaths.head())

jurisdiction_deaths_sorted = jurisdiction_deaths.sort_values('NumberOfDeaths',␣
↪ascending=False).reset_index(drop=True)
display(jurisdiction_deaths_sorted.head())
```

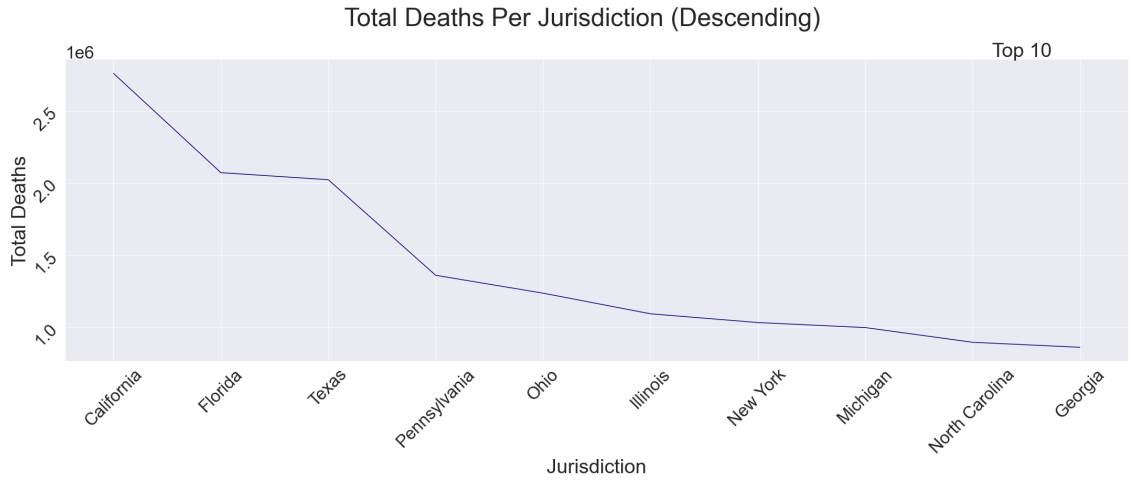
	Jurisdiction	NumberOfDeaths
0	Alabama	535321
1	Alaska	14676
2	Arizona	565455
3	Arkansas	321691
4	California	2763202

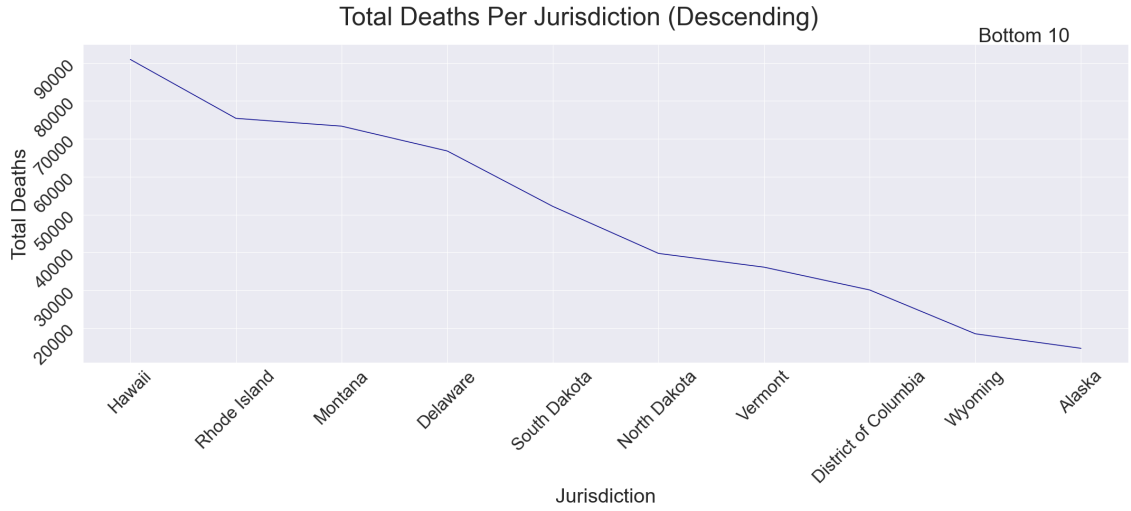
	Jurisdiction	NumberOfDeaths
0	California	2763202
1	Florida	2072697
2	Texas	2023671
3	Pennsylvania	1358740
4	Ohio	1234320

Lets plot the results in a line plot. We will plot the top 10 and then the bottom 10.

[25]:
```
sns.set(font_scale = 3)
sns.relplot(x='Jurisdiction', y='NumberOfDeaths', kind = 'line', height=10,␣
↪aspect=3, color='darkblue', data=jurisdiction_deaths_sorted.head(10))
plt.suptitle("Total Deaths Per Jurisdiction (Descending)", size=48, y=1.05)
plt.title("Top 10", size=38, x=0.9)
plt.xlabel("Jurisdiction", size=38)
plt.ylabel("Total Deaths", size=38)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()

sns.set(font_scale = 3)
sns.relplot(x='Jurisdiction', y='NumberOfDeaths', kind = 'line', height=10,␣
↪aspect=3, color='darkblue', data=jurisdiction_deaths_sorted.tail(10))
plt.suptitle("Total Deaths Per Jurisdiction (Descending)", size=48, y=1.05)
plt.title("Bottom 10", size=38, x=0.9)
plt.xlabel("Jurisdiction", size=38)
plt.ylabel("Total Deaths", size=38)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
```

```
plt.show()
```





2.2.8 2020 deaths by jurisdiction sorted from most to least

Now, lets sort the deaths from most to least for the year 2020.

[26]:
```
jurisdiction_deaths2020_sorted = yearJurisdiction_df.copy()
jurisdiction_deaths2020_sorted = jurisdiction_deaths2020_sorted.loc[(2020)].
↪sort_values('NumberOfDeaths', ascending=False).reset_index()

# Display the head
display(jurisdiction_deaths2020_sorted.head())

# Display the tail.
```

```
display(jurisdiction_deaths2020_sorted.tail())
```

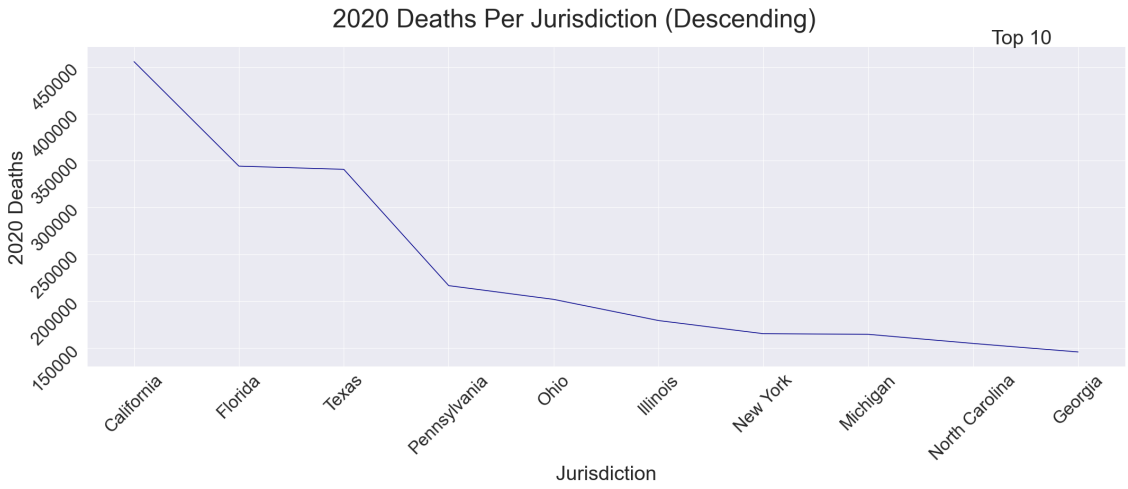
```
   Jurisdiction  NumberOfDeaths
0    California          455788
1       Florida          344224
2         Texas          340776
3  Pennsylvania          216650
4          Ohio          201957
```

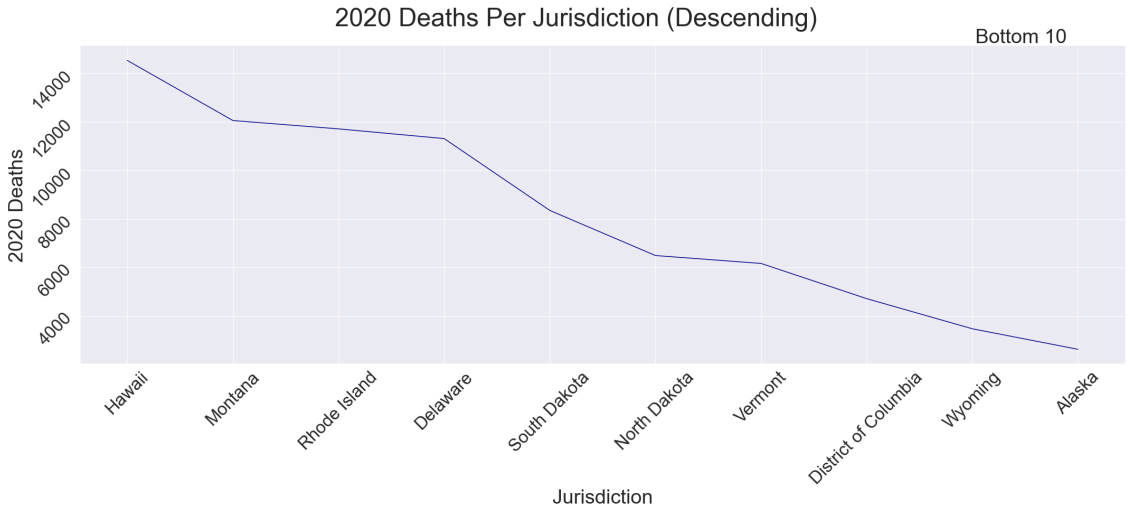
```
            Jurisdiction  NumberOfDeaths
48          North Dakota            6492
49               Vermont            6166
50  District of Columbia            4714
51               Wyoming            3472
52                Alaska            2626
```

2.3 Plot the Results

Lets plot the results for the year 2020 from most deaths to least deaths in 2 seperate plots. First, we will plot the top 10 and then we will plot the bottom 10.

[27]:
```
sns.set(font_scale = 3)
sns.relplot(x='Jurisdiction', y='NumberOfDeaths', kind = 'line', height=10,␣
 ↪aspect=3, color='darkblue', data=jurisdiction_deaths2020_sorted.head(10))
plt.suptitle("2020 Deaths Per Jurisdiction (Descending)", size=48, y=1.05)
plt.title("Top 10", size=38, x=0.9)
plt.xlabel("Jurisdiction", size=38)
plt.ylabel("2020 Deaths", size=38)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()

sns.set(font_scale = 3)
sns.relplot(x='Jurisdiction', y='NumberOfDeaths', kind = 'line', height=10,␣
 ↪aspect=3, color='darkblue', data=jurisdiction_deaths2020_sorted.tail(10))
plt.suptitle("2020 Deaths Per Jurisdiction (Descending)", size=48, y=1.05)
plt.title("Bottom 10", size=38, x=0.9)
plt.xlabel("Jurisdiction", size=38)
plt.ylabel("2020 Deaths", size=38)
plt.xticks(rotation=45)
plt.yticks(rotation=45)
plt.show()
```


2020 Deaths Per Jurisdiction (Descending)
Top 10
2020 Deaths
450000
400000
350000
300000
250000
200000
150000
California
Florida
Texas
Pennsylvania
Ohio
Illinois
New York
Michigan
North Carolina
Georgia
Jurisdiction


2020 Deaths Per Jurisdiction (Descending)
Bottom 10
2020 Deaths
14000
12000
10000
8000
6000
4000
Hawaii
Montana
Rhode Island
Delaware
South Dakota
North Dakota
Vermont
District of Columbia
Wyoming
Alaska
Jurisdiction

[ ]: