```
# Read data. This data represents the cumulative known cases to date (https://covidtracking.com/about-data/fag)
unl = https://magithubusercontent.com/COVIDISTracking/covid-tracking-data/master/data/states_daily_4pm_et.csv
df = pd.read_csv(unl,index_col=0,parse_dates=(0))
             state positive negative pending hospitalizedCurrently hospitalizedCumulative inIcuCurrently inIcuCumulative onVentilatorCurrently onVentilatorCurrently recovered
                                                                                                                                                                                                    262.0 d15bb8087806f8dded75fb6165d7bed1bcface44 2020-05-
03T20:00:00Z 9.0
               AK 368.0 21210.0 NaN
                                                                       12.0
                                                                                                  NaN
                                                                                                                    NaN
                                                                                                                                       NaN
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                                                                                                                                                                                                                                                                                                                    21578.0 2
                                                                                                                                                                                                       NaN aa4e1102894dd9528461e4d910fab397cc40d31b 2020-05-
03T20:00:00Z 290.0
                                                                                                                                                                                                    1999.0 6b15aab296af49161db3f9fc69eef7c6ca244994 2020-05-03T20:00:00Z 76.0
                                                                       100.0
                                                                                                 427 0
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05-03
              AS
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03T20:00:00Z 0.0
                                                                                                                                                                                                                                                                                                                       57.0
                                                                                                                                                                                                    1597.0 c687661dcf3f892bcc23c38ddcb46fe809b006f1 2020-05-
03T20:00:00Z 362.0
              AZ 8640.0 72479.0 NaN
                                                                       732 N
                                                                                                1348 0
                                                                                                                   282.0
                                                                                                                                                               192 0
                                                                                                                                                                                                                                                                                        1348 0 81119 0
                                                                                                                                                                                                                                                                                                                     81110 0 1
Double-click (or enter) to edit
['iniculumently', 'iniculumulative', 'onventilatorCumulative', 'hash', 'dateChecked', 'hospitalized', 'total', 'posNeg', 'fips', 'deathIncrease', 'hospitalizedincrease', 'negativeIncrease', 'positiveIncrease', 'totalTestResultsIncrease'])
 г
                 state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults
      2020-05-03
       2020-05-03
                             7725.0 84775.0
                                                                                              427.0 1999.0 76.0
      2020-05-03 AR 3431.0 49459.0 NaN
                                                                          100.0
                                                                                                                                            E2900 0
       2020-05-03
                               0.0
                                        57.0 NoN
                                                                            NaN
                                                                                                       NoN
                                                                                                                   NoN 0.0
                                                                                                                                              57.0
      2020-05-03 AZ 8640.0 72479.0
                                                  NaN
                                                                          732.0
                                                                                                    1348.0 1597.0 362.0
                                                                                                                                            81119.0
# Create new features
# Divide positive by totalTestResults to get positive_percent
df_drop["percent_positive"] = ""
df_drop["percent_positive"] = 100*df_drop["positive"]/df_drop["totalTestResults"]
df_drop.head()
                  state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent positive
      2020-05-03
                             368.0 21210.0
                                                                                                                  262.0 9.0
                                                                             12.0
                                                                                                                                            21578.0
      2020-05-03
                    AL 7725.0 84775.0
                                                                            NaN
                                                                                                                  NaN 290.0
                                                                                                                                            92500.0
                                                                                                                                                               8.351351
                   AR 3431.0 49459.0 NaN
                                                                           100.0
                                                                                                     427.0 1999.0 76.0
                                                                                                                                                               6.487049
                                                                           732.0
                                                                                                                  1597.0 362.0
# Divide hospitalized by positive to get hospitalized_percent
import numpy as np

df_drop("hospitalized_percent") = ""

df_drop("hospitalized_percent") = np.nammax(df_drop(['hospitalizedCurrently','hospitalizedCumulative']], axis=1)

df_drop("hospitalized_percent"] = 100*df_drop("hospitalized_percent")|df_drop("positive")|

df_drop_head()
 /usr/local/lib/python3.6/dist-packages/ipykernel_launcher.py:3: RuntimeWarning: All-NaN axis encountered
This is separate from the ipykernel package so we can avoid doing imports until
                   state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent_positive hospitalized_percent
      2020-05-03 AK
                             368.0 21210.0
                                                                            12.0
                                                                                                                  262.0 9.0
                                                                                                                                            21578.0
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      2020-05-03 AI 7725.0 84775.0 NaN
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      2020-05-03 AR 3431.0 49459.0 NaN
                                                                           100.0
                                                                                                                                            52890.0
                                                                                                                                                              6.487049
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      2020-05-03 AZ 8640.0 72479.0 NaN
                                                                                                    1348.0 1597.0 362.0
                                                                                                                                            81119.0
                                                                                                                                                              10.651019
# Divide recovered by positive to get recovered_percent
df_drop["recovered_percent"] = ""
df_drop["recovered_percent"] = 190*df_drop["recovered"]/df_drop["positive"]
df_drop_head()
                  state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent positive hospitalized percent recovered percent
      2020-05-03 AK
                             368.0 21210.0
                                                                             12.0
                                                                                                       NaN
                                                                                                                  262.0 9.0
                                                                                                                                            21578.0
                                                                                                                                                               1.705441
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      2020-05-03
                   AL 7725.0 84775.0 NaN
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                                                                                                                  1597.0 362.0
                                                                                                                                             81119.0
                                                                                                                                                               10.651019
                                                                                                                                                                                      15.601852
                                                                                                                                                                                                           18.483796
# Divide death by positive to get death_percent
# Divide death by positive to get death_percent

df_drop["death_percent"] = ""

df_drop["death_percent"] = 100*df_drop["death"]/df_drop["positive"]

df_drop.head()
```

```
# Fetch the latest state population data (nst-est2019-01.csv)
from google.colab import files
uploaded = files.upload()

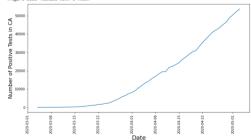
D: Choose Files | rat-est2019-01 cav | - nst-est2019-01 cav | - nst-est2019-01.csv (application/wnd.ms-excel) - 676 bytes, last modified: 4/13/2020 - 100% done Saving nst-est2819-81.csv to nst-est2819-81.csv
# Load latest state population data
         State Population
          AK 731545.0
      1 AL 4903185.0
      2 AR 3017804.0
# Add column of state populations (population) to df_drop_total_posNeg # Need to sort rows by state using index numbering from state_list
df_drop["population"] = ""
for i in range(len(df_drop)):
    for index in range(len(df_state_pop)):
        if df_drop.iloc[i, 0] = off_state_pop.iloc[index, 0]:
        df_drop.iloc[i, 13] = df_state_pop.iloc[index, 1]
df_drop.head()
             state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent positive hospitalized percent recovered percent death percent population
      2020-
                                                                                                             262.0
                       7725.0 84775.0
                                               NaN
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      2020-
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                                                                                                                                         57.0
                                                                                                                                                          0.000000
                                                                                                                                                                                                           NaN
                                                                                                                                                                                                                           NaN
# Normalize positive to state population
df_drop["positive_norm"] = ""
df_drop["positive_norm"] = df_drop["positive"]/df_drop["population"]
df_drop.nead["]
 г
              state positive negative pending hospitalizedCurrently hospitalizedCurrently hospitalizedCurrently hospitalized percent positive norm
      2020-
05-03
                        368.0 21210.0
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                                                                                                                                                                                 13.398058
                        3431.0 49459.0
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                                                                                                  427.0
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                                                                                                                                                                                                       18 483796
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                                                                                                                                                                                                                                                     0.001187
   Normalize hospitalized to state population
             state positive negative pending hospitalizedCurrently hospitalizedCurrently hospitalizedCurrently hospitalizedCurrently hospitalized morm hospitalized norm
       date
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                                                                                                             262.0 9.0
                        368.0 21210.0
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                        7725 0 84775 0
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05-03
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                                                                                                                                         57.0
                                                                                                                                                          0.000000
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               AZ 8640.0 72479.0
# Normalize recovered to state population
df_drop["recovered_norm"] = ""
df_drop["recovered_norm"] = df_drop["recovered"]/df_drop["population"]
df_drop.head()
              state positive negative pending hospitalizedCurrently hospitalizedCurrently hospitalizedCurrently hospitalized_more recovered death totalTestResults percent_positive hospitalized_percent recovered_percent death_percent population positive_norm hospitalized_norm recovered_norm
                                                                        NaN
                                                                                                                                                          8.351351
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                        7725.0 84775.0
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                                                                                                                                                                                                       18.483796
- ....-ourse ocatn to state population

df_drop["death_norm"] = ""

df_drop["death_norm"] = df_drop["death"]/df_drop["population"]

df_drop.head()
D+
              state positive negative pending hospitalizedCurrently hospitalizedCurrently hospitalized_morm recovered death totalTestResults percent positive hospitalized_percent recovered percent death_percent positive_norm hospitalized_norm recovered_norm death_norm
                         368.0 21210.0
                                                                                                   NaN
                                                                                                             262.0 9.0
                                                                                                                                       21578.0
                                                                                                                                                          1.705441
                                                                                                                                                                                  3.260870
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      2020-
05-03
                AZ 8640.0 72479.0 NaN
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                                                                                                             1597.0 362.0
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                                                                                                                                                                                 15.601852
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```

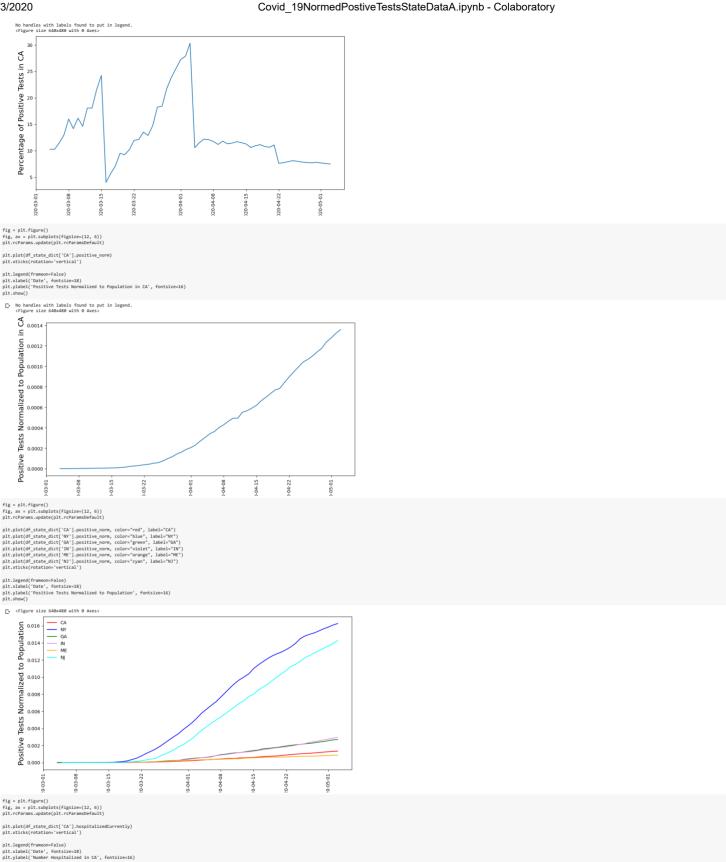
```
Covid 19NormedPostiveTestsStateDataA.ipynb - Colaboratory
5/3/2020
        [] <class 'pandas.core.frame.DataFrame'>
DatetimeIndex: 3321 entries, 2020-05-03 to 2020-01-22
Data columns (total 18 columns):
# Column Mon-Null Count Dtype
                # Get the unique values of 'state' column
state_list = df.state.unique()
state_list
        D. array(['Ax', 'Ax', 'Ax', 'Ax', 'Ax', 'CA', 'CO', 'CT', 'DC', 'DE', 'FL', 'GA', 'GO', 'H1', 'L1A', 'L1D', 'L1D', 'L1N', 'L1S', 'NC', 'L4N', 'NA', 'N
        df_state_dict['AK'].head()
                                state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent positive hospitalized percent recovered percent death_percent population positive_norm hospitalized_norm recovered_norm death_norm
                    2020-
05-03
                                                    368.0 21210.0
                                                                                                                                                                                             NaN
                                                                                                                                                                                                                 262.0 9.0
                                                                                                                                                                                                                                                                 21578.0
                                                                                                                                                                                                                                                                                                   1.705441
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                                                    364.0 19961.0
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                                  AK 355.0 18764.0
                                                                                                                                                                                                                 252.0
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                                    AK 355.0 18764.0
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                                                                                                                                                                                                                                                                 19119 0
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                                                                                                                                                                                                                                                                                                                                                                                    67 605634
                                                                                                                                                                                                                                                                                                                                                                                                                    2 535211
                                                                                                                                                                                                                                                                                                                                                                                                                                           731545 0
        df_state_dict['CA'].head()
                                state positive negative pending hospitalizedCurrently hospitalizedCumulative recovered death totalTestResults percent positive hospitalized percent recovered death percent population positive norm hospitalized norm recovered norm death norm
                                    CA 53616.0 662135.0
                                                                                                                                                                                                                                                                                                                                                                                                                      4.131229 39512223.0
                                    CA 52197.0 634606.0
                                                                                                                                        4722.0
                                                                                                                                                                                             NaN
                                                                                                                                                                                                                    NaN 2171.0
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                                  CA 48917.0 576420.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.001238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.000126
        from matplotlib import pyplot as plt
        plt.plot(df_state_dict['CA'].positive)
plt.xticks(rotation='vertical')
                   B
```

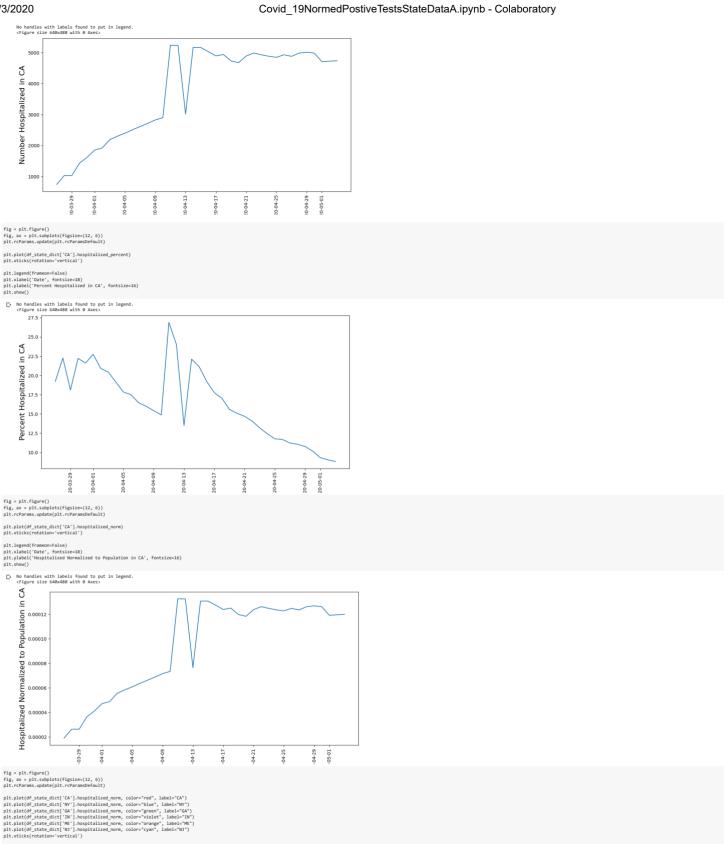


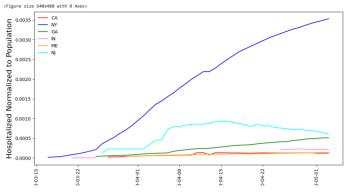
```
plt.plot(df_state_dict['CA'].percent_positive)
plt.xticks(rotation='vertical')
plt.xlabel('Date', fontsize=18)
plt.ylabel('Percentage of Positive Tests in CA', fontsize=16)
plt.show()
```

D-

plt.show()







In several states, population normalized hospitalizations plateau, although population normalized death rate continues to grow

fig = plt.figure()
fig, ar = plt.visuplots(figsize=(12, 6))
plt.rcParams.update(plt.rcParamsDefault)
plt.plot(of_state_dist('CA'].death)
plt.xticks(rotation='vertical')
plt.legend(frameon=false)
plt.label('Date', fontsize=18)
plt.vlabel('Date', fontsize=16)
plt.show()

No handles with labels found to put in legend. <Figure size 640x480 with 0 Axes>

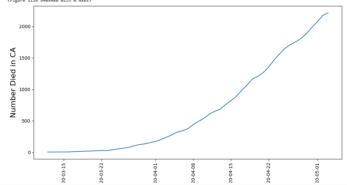
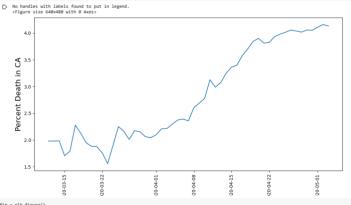


fig = plt.figure()
fig, ax = plt.subplots(figsize-(12, 6))
plt.rcParams.update(plt.rcParamsDefault)
plt.plt(fdf_state_dict['Ca'].death_percent)
plt.xticks(rotation='vertical')
plt.legend(frameon=False)
olt.ylabe('jots'_fortstats)

plt.show()



fig, as = plt.subplots(figsize=(12, 6))
plt.rcParams.update(plt.rcParamsDefault)
plt.plot(ff.state_dict['Ca'].death_norm)
plt.plt.cff.state_dict['Ca'].death_norm)
plt.legend(frameon=False)
plt.legend(frameon=False)
plt.legend(frameon=False)
plt.legend(frameon=False)

plt.legend(frameon=False)
plt.xlabel('Date', fontsize=18)
plt.ylabel('Death Normalized to Population in CA', fontsize=16)
plt.show()

C+

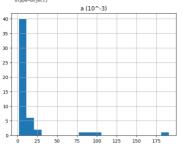
```
No handles with labels found to put in legend. 
<Figure size 640x480 with 0 Axes>
                  1e-5
        Death Normalized to Population in CA
plt.plot(df_state_dict['CA'].death_norm, color="red", label="CA")
plt.plot(df_state_dict['M'].death_norm, color="blue", label="M')
plt.plot(df_state_dict['M'].death_norm, color="plen", label="GA')
plt.plot(df_state_dict['M'].death_norm, color="red", label="M')
plt.plot(df_state_dict['M'].death_norm, color="volot", label="M')
plt.plot(df_state_dict['M'].death_norm, color="cyan", label="M')
plt.plot(df_state_dict['M'].death_norm, color="cyan", label="M')
plt.state(cycloriem="vertical")
plt.legend(frameon=False)
plt.xlabel('Date', fontsize=18)
plt.ylabel('Death Normalized to Population', fontsize=16)
plt.show()

← Figure size 640x480 with 0 Axes

           0.0010 - CA - NY - GA - IN - ME - NJ
        Normalized to Population
        Death I
               0.0002
Note how the population normalized death curves relate closely to population normalized postive test curves
# Curve fitting done at: \underline{\text{http://www.xuru.org/rt/NLR.asp\#CopyPaste}}
# Fetch the parameters for each state (AexpBx*-1.csv) that fit to positive_norm = a^*exp(b/x) # where x is the number of days from March 4, 2020 from goagle.colab import files uploaded = files.uploadef
 # Load the parameters for each state (AexpBx^-1.csv) that fit to positive_norm = a^*exp(b/x) import io
amport to
df_state_params = pd.read_csv(io.StringIO(uploaded['AexpBx^-1.csv'].decode('utf-8')))
df_state_params.head()
            State a (10^-3)
       0 AK 2.593040 -75.366476 1.0 0.996906
1 AL 12.121593 -111.222242 2.0 0.997430
       2 AR 2.941186 -75.356785 4.0 0.997586
3 AS NaN NaN NaN NaN NaN
               AZ 4.984063 -90.295019
                                                             1.0 0.998613
df_state_params.describe()
 C+
                                            b fit rank
        count 52.000000 52.000000 52.000000 52.000000
         mean 16.215254 -100.951881 1.769231 0.995682
                 31.801661 25.545128 1.095720 0.004749
                    1.952592 -185.986576 1.000000 0.972728
         25% 5.041013 -116.155268 1.000000 0.995399
                   7.113788 -99.476492 1.000000 0.997030
          max 190.553218 -49.104858 5.000000 0.999660
E+
```

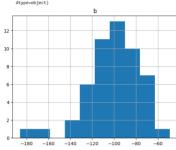
array([[<matplotlib.axes._subplots.AxesSubplot object at 0x7f093e578940>]],

df_state_params.hist(column='a (10^-3)', bins=20)



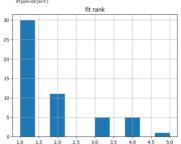
High value outliers here are NJ (fit rank 1), NY, (fit rank 1), RI (fit rank 5), and SD (fit rank 4)

df_state_params.hist(column='b', bins=10)



Low value outliers here are RI (fit rank 5) and SD (fit rank 4).

df_state_params.hist(column='fit rank')



The A*exp(B/x) functional form works extremely well for thirty of the 52 states (57.7%).

Fetch static data for each state (CovidCompleteStateData.csv)
from google.colab import files
uploaded = files.upload()

[2] Choose Files CovidCompl...teData.csv

• CovidCompleteStateData.csv(application\vnd.ms-excel) - 60510 bytes, last m Saving CovidCompleteStateData.csv to CovidCompleteStateData.csv

Load static data for each state (CovidCurrentStateData.csv)
import io
df_state_data = pd.read_csv(io.StringIO(uploaded['CovidCompleteStateData.csv'].decode('utf-8')))
df_state_data.head()

D.		ate I	Sum of NUM_Medicare_BEN		Sum of NUM_BEN_Age_65_to_74	Sum of NUM_BEN_Age_75_to_84	Sum of NUM_BEN_Age_Greater_84	Sum of NUM_Female_BEN	Sum of NUM_Male_BEN	Sum of NUM_Black_or_African_American_BEN	Sum of NUM_Asian_Pacific_Islander_BEN	Sum of NUM_Hispanic_BEN	Sum of NUM_American_IndianAlaska_Native_BEN NUM_BEN_With_
	0	AK	1820384.0	270970.0	809516.0	468255.0	175296.0	1034762.0	760009.0	62311.0	76773.0	46525.0	147917.0
	1	AL	10804823.0	2065353.0	4386595.0	2980828.0	1190504.0	6237445.0	4514041.0	1549811.0	30624.0	65500.0	5556.0
	2	AR	15892716.0	2818665.0	6370265.0	4555468.0	1848506.0	9275039.0	6507151.0	1334245.0	19642.0	108428.0	62782.0
	3	AS	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN
	4	AZ	10786064.0	886596.0	4861035.0	3377040.0	1294375.0	5944519.0	4747801.0	221183.0	61840.0	689880.0	179818.0
	5 rows	× 116	columns										

Elevation Ratio = Highest Elevation/Mean Elevation
of state_data['Elevation Ratio'] = of state_data['Highest Elevation']/of_state_data['Mean Elevation']
of state_data['Elevation Ratio'] = of state_data['Highest Elevation'],divide(of_state_data['Mean Elevation'], fill_w.

Capital Area Ratio = Capital Land Area/Capital Water Area
of state_data['Capital Area Ratio'] = of_state_data['Capital Land Area']/df_state_data['Capital Water Area']

of_state_data['Capital Land Area'] = of_state_data['Capital Land Area'].astype(float)

of_state_data['Capital Area Ratio'] = of_state_data['Capital Land Area'].divide(of_state_data['Capital Water Area'],

Boundaries = Number of boarding states + On Coast + Borders Another Country
df_state_data['Boundaries'] = df_state_data['Number of bordering states'] + df_state_data['On Coast'] + df_state_data

Latitude Difference to State Capital = Latitude - Capital Latitude

df_state_data['Latitude Difference to State Capital'] = df_state_data['Latitude'] - df_state_data['Capital Latitude']

Longitude Difference to State Capital = Capital Longitude - Longitude

df_state_data['Longitude Difference to State Capital'] = df_state_data['Capital Longitude'] - df_state_data['Longitude']

Latitude Difference to DC = Latitude - DC Latitude
df_state_data['Latitude Difference to DC'] = df_state_data['Latitude'] - 38.984722

Longitude Difference to DC = DC Longitude - Longitude
df_state_data['Longitude Difference to DC'] = -77.016389 - df_state_data['Longitude'] # Latitude Difference to US Center = Latitude - Center Latitude df_state_data['Latitude Difference to Center'] = df_state_data['Latitude'] - 39.833333

Longitude Different to US Center = Center Longitude - Longitude

df_state_data['Longitude Difference to Center'] = -98.585522 - df_state_data['Longitude'] df state data.head() State Sum of Sum 1820384.0 270970.0 809516.0 468255.0 175296.0 1034762.0 760009.0 62311.0 46525.0 809516.0 488255.0 175296.0 1034782.0 760009.0 4386595.0 2980828.0 1190504.0 6237445.0 4514041.0 6370265.0 4555468.0 1848506.0 9275039.0 6507151.0 NaN NaN NaN NaN NaN NaN 4861035.0 3377040.0 1294375.0 5944519.0 4747801.0 76773.0 147917.0 1820384.0 270970.0 10804823.0 2065353.0 15892716.0 2818665.0 NaN NaN 10786064.0 886596.0 1549811.0 30624.0 65500.0 5556.0 221183.0 61840.0 689880.0 179818.0 5 rows × 126 columns df state data.shape

Define variables for regression
df_templ = df_state_data.drop(df_state_data.index[[3, 12, 27, 42, 50]])
x = df_templ.drop('state', asts = 1)
df_templ = df_state_parass.drop(df_state_data.index[[3, 12, 27, 42, 50]])
y = df_templ[7, (30-3)']

Look at correlation coefficients
pd.set_option('display.max_columns', None)
pd.set_option('display.max_rows', 1000)
X.corr()

	Sum of NUM_Medicare_BEN	Sum of NUM_BEN_Age_Less_65	Sum of NUM_BEN_Age_65_to_74	Sum of NUM_BEN_Age_75_to_84	Sum of NUM_BEN_Age_Greater_84	Sum of NUM_Female_BEN	Sum of NUM_Male_BEN	Sum of NUM_Black_or_African_American_BEN	Sum of NUM_Asian_Pacific_Islander_BEN	Sum of NUM_Hispanic_BEN NUM_Am
Corr of NUM Medicary DEN	4.000000	0.981404	0.998624	0.998100	0.989961	0.999917	0.999897	0.896692	0.525530	0.893302
Sum of NUM_Medicare_BEN Sum of NUM_BEN_Age_Less_65	1.000000 0.981404	1.000000	0.998624	0.998100	0.989961		0.999897	0.896692	0.525530	0.893302
Sum of NUM_BEN_Age_65_to_74	0.998624	0.978099	1.000000	0.996374			0.998636	0.895722		
Sum of NUM_BEN_Age_75_to_84	0.998100		0.996374				0.998296	0.884218		
Sum of NUM_BEN_Age_Greater_84	0.989961	0.960650	0.982712	0.992601	1.000000	0.989606	0.990404	0.864777	0.561253	0.879739
Sum of NUM_Female_BEN	0.999917	0.982576	0.998372	0.997916	0.989606	1.000000	0.999658	0.899227	0.523618	0.889962
Sum of NUM_Male_BEN	0.999897	0.979741	0.998636	0.998296	0.990404	0.999658	1.000000	0.893490	0.527005	0.895526
Sum of NUM_Black_or_African_American_BEN	0.896692	0.926091	0.895722	0.884218	0.864777	0.899227	0.893490	1.000000	0.302985	0.726543
Sum of NUM_Asian_Pacific_Islander_BEN	0.525530	0.475021	0.517514	0.530001	0.561253		0.527005	0.302985	1.000000	
Sum of NUM_Hispanic_BEN	0.893302	0.827878	0.902298	0.899556	0.879739		0.895526	0.726543	0.633875	
Sum of NUM_American_IndianAlaska_Native_BEN	0.082561	0.059858	0.091513		0.065730		0.083230	-0.035649		
Sum of NUM_BEN_With_Race_Not_Elsewhere_Classified	0.823477 0.996838	0.774080 0.978894	0.803783 0.994391	0.832225 0.996119			0.824850 0.996745	0.638847 0.888851	0.739901 0.485602	0.734084
Sum of NUM_Non-Hispanic_White_BEN Sum of NUM_Minorities	0.996838	0.978894	0.994391	0.996119	0.988883 0.945115		0.958590	0.888851	0.485602	0.865686
Sum of Average Age of BEN	0.682483		0.686432	0.663590	0.637504		0.678577	0.692434		
Sum of NUM_BEN_Atrial_Fibrillation	0.990425	0.969550	0.985604	0.991418			0.990816	0.889881		
Sum of NUM_BEN_Asthma	0.995532	0.979588	0.991583	0.992903	0.991762		0.995600	0.893349	0.526980	0.880784
Sum of NUM_BEN_Cancer	0.994765	0.972149	0.992903	0.994874	0.987401	0.994443	0.994921	0.900538	0.464158	0.883930
Sum of NUM_BEN_Heart_Failure	0.997133	0.985150	0.995371	0.993915	0.984952	0.997171	0.996846	0.913138	0.484598	0.884681
Sum of NUM_BEN_Chronic_Kidney_Disease	0.997501	0.980301	0.997095	0.995430	0.984274	0.997279	0.997616	0.907084	0.485414	0.893142
Sum of NUM_BEN_Chronic_Obstructive_Pulmonary_Disease	0.986234	0.980624	0.981625	0.983999	0.978052	0.986989	0.985885	0.906585	0.429822	0.833395
Sum of NUM_BEN_Hyperlipidemia	0.996237	0.974348	0.994742	0.996423	0.987588	0.996101	0.996491	0.903165	0.477347	0.884215
Sum of NUM_BEN_Diabetes	0.997754	0.981227	0.996544	0.995687	0.985896		0.997458	0.912776		0.892805
Sum of NUM_BEN_Hypertension	0.998856		0.998079		0.986018		0.998633	0.908147		
Sum of NUM_BEN_Ischemic_Heart_Disease	0.994006	0.975145	0.991547	0.994105	0.985840	0.994115	0.993975	0.906294	0.457797	0.876921
Sum of NUM_BEN_Stroke	0.990547	0.972081	0.988818	0.990024	0.980879	0.990462	0.990642	0.919103	0.447938	0.879610
Sum of PCT_MEDICARE	0.713702	0.762102	0.716971	0.696228	0.671536	0.717742	0.709263	0.752793	0.141713	0.484719
% Urban Pop	0.246412		0.240984	0.259055	0.285836		0.250798	0.181632	0.311802	
Density (P/mi2)	-0.095479		-0.096280	-0.092015	-0.087655		-0.095250	-0.017993	-0.029294	-0.041712
Children 0-18	0.886252	0.846604	0.876226	0.888481	0.912717		0.887362	0.723346		
Adults 19-25	0.865749	0.826231 0.804492	0.852680 0.835397	0.868860 0.852982	0.900273 0.886196		0.866800 0.850081	0.698451		0.808969
Adults 26-34 Adults 35-54	0.848661	0.804492	0.835397	0.865769			0.862945	0.667717 0.695994		
Adults 55-64	0.840536	0.802214	0.822003	0.845654	0.889137		0.841899	0.678395		
65+	0.842520	0.796154	0.822919	0.852028	0.896626		0.844588	0.672577	0.691686	
Latitude	-0.400391	-0.397373	-0.403138	-0.407192	-0.381151	-0.404042	-0.398907	-0.449230	-0.184344	-0.284522
Longitude	0.046601	0.092974	0.034115	0.040031	0.057825	0.049875	0.043076	0.189118	-0.272166	-0.097547
Land Area	0.229013	0.193883	0.242084	0.230058	0.205886	0.226251	0.230971	0.128422	0.201398	0.342072
Water Area	0.042895	0.056385	0.036723	0.038782	0.050598	0.042948	0.042530	0.080106	0.048930	0.044540
Mean Elevation	-0.163276	-0.224740	-0.147730	-0.155029	-0.169641	-0.168622	-0.157738	-0.314621	0.096312	0.038452
Highest Elevation	-0.059881	-0.137582	-0.040603	-0.049835	-0.071530		-0.054634	-0.233143		
Lowest elevation	-0.354394	-0.352655	-0.344053	-0.355481	-0.373528		-0.352697	-0.312823		
Number of bordering states	0.077790		0.075964	0.059448	0.056612		0.074603	0.044175		
On Coast	0.471024	0.505115	0.442960	0.461862	0.518306 0.359742		0.470022	0.512059	0.172256	0.274148
Borders Another Country Capital Latitude	-0.388663	0.310618 -0.393979	-0.394070	0.356815 -0.392266	-0.359742		-0.386636	0.188403 -0.463654	0.423258 -0.136997	-0.269834
Capital Longitude	0.027375		0.015075	0.019608	0.037346		0.023488	0.181245		
Captial Land Area	0.008902	-0.002410	0.018688	0.009403			0.008880	-0.020032		
Capital Water Area	-0.087670		-0.083610				-0.087364	-0.097128		
Capital Mean Elevation	-0.194009	-0.217624	-0.182931	-0.190725	-0.206936	-0.197080	-0.190497	-0.249034	-0.121862	-0.049687
Capital is the Largest City	-0.171080	-0.147972	-0.165860	-0.173283	-0.194417	-0.169162	-0.172323	-0.133938	-0.129514	-0.188909
Largest City Latitude	-0.421170	-0.421496	-0.425109	-0.424938	-0.398170	-0.424688	-0.419431	-0.467601	-0.234938	-0.317480
Largest City Longitude	0.057094	0.102104	0.044423	0.050338	0.069739		0.053604	0.201824	-0.262204	-0.081857
Number of Counties	0.663716	0.710105	0.670375	0.645677	0.611985	0.666592	0.659547	0.684930	0.100824	0.503011
Became a State	-0.140415	-0.200801	-0.128869	-0.126557	-0.143131	-0.144064	-0.136075	-0.308218	0.076276	
Days Since Stayat Home Order	-0.020651	-0.019693	-0.030343	-0.027347		-0.023631	-0.018913	-0.045827		0.052471
DaysSinceFirstPositive DaysSinceTestStart	0.368252 0.290649		0.366229 0.289428	0.374653 0.297948			0.370903 0.292640	0.287102 0.233275		0.302869
15-49yearsAllcauses	0.888203		0.874919				0.889058	0.739853		
15-49yearsAsthma	0.824682		0.807656				0.825395	0.667614		
15-49yearsChronickidneydisease	0.918864	0.893772	0.909568	0.918825	0.935419	0.918655	0.918932	0.805947	0.715267	0.828201
15-49yearsChronicobstructivepulmonarydisease	0.896769	0.878089	0.880516	0.897303	0.928648	0.896643	0.896964	0.771591	0.635771	0.749901
15-49yearsDiabetesmellitus	0.912330	0.881654	0.900896	0.914260	0.936863	0.911726	0.912809	0.781996	0.692768	0.812172
15- 49yearsInterstitiallungdiseaseandpulmonarysarcoidosis	0.881251	0.864222	0.866766	0.880121	0.909006	0.881289	0.881043	0.782863	0.644183	0.738019
15-49yearsIschemicheartdisease	0.928387	0.927789	0.916634	0.923405	0.939571	0.929317	0.927199	0.849388	0.595981	0.764879
15-49yearsNeoplasms	0.887461	0.860138	0.873067	0.888685	0.919369	0.886922	0.887968	0.748507	0.729958	0.785252
15-49yearsOtherchronicrespiratorydiseases	0.906636	0.885246	0.892415	0.906637	0.934764	0.906287	0.906852	0.785146	0.652797	0.775052
15-49yearsRheumaticheartdisease	0.903473	0.893269	0.893364	0.898792	0.916822	0.903373	0.903469	0.792144	0.690715	0.785070
15-49yearsStroke	0.919789	0.898558	0.910295	0.919449		0.919891	0.919744	0.808343		0.815118
50-69yearsAllcauses	0.880146		0.863069				0.880993	0.745024		
50-69yearsAsthma	0.801803	0.765502	0.781306	0.805925			0.802845	0.641040	0.741394	0.706390
50-69yearsChronickidneydisease	0.917312	0.898401 0.872843	0.905572 0.860771	0.916416			0.917589	0.809479	0.675662 0.542692	
50-69yearsChronicobstructivepulmonarydisease 50-69yearsDiabetesmellitus	0.879259	0.872843	0.865414	0.878641 0.884673			0.879770	0.765135 0.753933		
50-69yearsDiabetesmeilitus 50-										
69yearsInterstitiallungdiseaseandpulmonarysarcoidosis	0.863191	0.840683	0.846169	0.863950	0.901043	0.862755	0.863919	0.739130	0.673635	0.725850
50-69yearsIschemicheartdisease	0.905979		0.890019				0.905613			
50-69yearsNeoplasms	0.872500		0.854035				0.873220	0.746019		
50-69yearsOtherchronicrespiratorydiseases	0.885021	0.875159	0.867604	0.883457			0.885343	0.780290		
50-69yearsRheumaticheartdisease	0.892519		0.880528 0.895108	0.886667 0.907388	0.908250 0.930491		0.892705 0.908324	0.793842 0.800720		
50-69yearsStroke 70+vearsAllcauses	0.907993	0.892290 0.819400	0.895108 0.828488	0.907388 0.854118			0.908324	0.800720 0.700955	0.656820 0.653903	
70+yearsAsthma	0.791486	0.748032	0.769602	0.799338			0.793100	0.612235		0.697428
70+yearsChronickidneydisease	0.791486	0.858325	0.769602	0.877628			0.793100			
70+yearsChronicobstructivepulmonarydisease	0.866728		0.846829				0.868173	0.721373		
70+yearsDiabetesmellitus	0.845276		0.823824	0.850785			0.846627	0.702717		
70+yearsInterstitiallungdiseaseandpulmonarysarcoidosis	0.833832	0.799993	0.814083	0.839080	0.884501	0.833012	0.835543	0.675650	0.689192	0.705929
70+yearsIschemicheartdisease	0.841243	0.819787	0.818295	0.844148	0.893641	0.840980	0.841899	0.706345	0.634076	0.681143
70+yearsNeoplasms	0.837485		0.816021	0.842466	0.889888		0.838922	0.689286		
70+yearsOtherchronicrespiratorydiseases	0.875916		0.858192	0.875620			0.876936	0.752057	0.586677	0.704251
70+yearsRheumaticheartdisease	0.844465	0.839621	0.826738	0.839480			0.845500	0.720717		0.679448
70+yearsStroke	0.871562	0.849583	0.854254	0.873252	0.910139	0.871322	0.872668	0.732812	0.659665	0.720810

AllAgesAllcauses	0.880003	0.851293	0.863399	0.882592	0.918634	0.879356	0.880977	0.736553	0.695184	0.757560
AllAgesAsthma	0.833253	0.794917	0.815812	0.836910	0.879310	0.831811	0.834191	0.674054	0.748697	0.754660
AllAgesChronickidneydisease	0.905462	0.885503	0.891504	0.905307	0.933093	0.905311	0.905943	0.788948	0.671941	0.774120
AllAgesChronicobstructivepulmonarydisease	0.877214	0.860833	0.858127	0.879265	0.916074	0.877240	0.878179	0.745966	0.580626	0.696350
AllAgesDiabetesmellitus	0.879728	0.852121	0.862209	0.882908	0.919854	0.879232	0.880670	0.745944	0.658313	0.743480
AllAgesInterstitiallungdiseaseandpulmonarysarcoidosis	0.853912	0.826093	0.835749	0.856759	0.897256	0.853326	0.855050	0.714595	0.681169	0.720490
AllAgesIschemicheartdisease	0.883535	0.870954	0.864460	0.883069	0.920941	0.883720	0.883584	0.768383	0.628518	0.717687
AllAgesNeoplasms	0.865325	0.841450	0.846325	0.867726	0.908429	0.864932	0.866273	0.728920	0.662908	0.719608
AllAgesOtherchronicrespiratorydiseases	0.903592	0.885967	0.888445	0.902970	0.932994	0.903300	0.904011	0.785664	0.622884	0.752887
AllAgesRheumaticheartdisease	0.880357	0.875286	0.866144	0.875089	0.904080	0.880168	0.880883	0.767398	0.647527	0.727462
AllAgesStroke	0.895398	0.875735	0.880674	0.895978	0.925120	0.895381	0.896074	0.771183	0.666743	0.752509
AllAgesTotal	0.880507	0.853923	0.863463	0.882813	0.919294	0.879978	0.881411	0.740657	0.683664	0.750268
Airpollution	0.889229	0.888442	0.875092	0.882964	0.910652	0.889624	0.888991	0.780168	0.654775	0.729374
Highbody-massindex	0.893797	0.872739	0.877133	0.894624	0.928715	0.893524	0.894332	0.771258	0.660813	0.753636
Highfastingplasmaglucose	0.886795	0.870124	0.868909	0.887417	0.922985	0.886779	0.887281	0.773198	0.617042	0.724166
HighLDLcholesterol	0.893215	0.882483	0.875398	0.892023	0.926304	0.893527	0.893124	0.783354	0.627247	0.727214
Highsystolicbloodpressure	0.897453	0.882631	0.880346	0.897131	0.930464	0.897547	0.897670	0.788283	0.638519	0.741135
Impairedkidneyfunction	0.889934	0.872693	0.873173	0.890034	0.923835	0.889974	0.890321	0.773115	0.657798	0.740044
Noaccesstohandwashingfacility	0.877603	0.857781	0.862453	0.876519	0.909499	0.877112	0.877940	0.756686	0.667841	0.744063
Smoking	0.881579	0.866726	0.862831	0.882612	0.919020	0.881762	0.882099	0.760942	0.604036	0.705752
Log10Pop	0.728494	0.737902	0.714057	0.722320	0.747129	0.729527	0.727871	0.666572	0.421521	0.509917
DaysSinceInfection	0.422525	0.373010	0.419727	0.431233	0.443999	0.420775	0.424402	0.365316	0.263022	0.351935
Children0-18	0.167133	0.180823	0.181296	0.159580	0.119098	0.169559	0.164119	0.169612	0.049811	0.148864
Allriskfactors	0.882815	0.860944	0.865530	0.884217	0.920134	0.882487	0.883653	0.750594	0.660552	0.737247
State Area Ratio	-0.141342	-0.180449	-0.126323	-0.134563	-0.158168	-0.143787	-0.137979	-0.260308	-0.103905	-0.027420
Elevation Ratio	0.020332	0.007311	0.029598	0.023691	-0.000223	0.021715	0.018482	0.129890	0.052935	0.012281
Capital Area Ratio	-0.119284	-0.151665	-0.109968	-0.112407	-0.124967	-0.120614	-0.117747	-0.171038	-0.076449	-0.052022
Boundaries	0.499356	0.556393	0.479330	0.477960	0.517125	0.499487	0.497522	0.455272	0.125465	0.277110
Latitude Difference to State Capital	-0.268652	-0.211068	-0.252026	-0.293417	-0.327548	-0.268822	-0.271605	-0.115573	-0.406313	-0.233715
Longitude Difference to State Capital	-0.143646	-0.133106	-0.139285	-0.150250	-0.155036	-0.143567	-0.145263	-0.100819	-0.106975	-0.108075
Latitude Difference to DC	-0.400391	-0.397373	-0.403138	-0.407192	-0.381151	-0.404042	-0.398907	-0.449230	-0.184344	-0.284522
Longitude Difference to DC	-0.046601	-0.092974	-0.034115	-0.040031	-0.057825	-0.049875	-0.043076	-0.189118	0.272166	0.097547
Latitude Difference to Center	-0.400391	-0.397373	-0.403138	-0.407192	-0.381151	-0.404042	-0.398907	-0.449230	-0.184344	-0.284522
Longitude Difference to Center	-0.046601	-0.092974	-0.034115	-0.040031	-0.057825	-0.049875	-0.043076	-0.189118	0.272166	0.097547

Note that there are many highly correlated features which need to be dropped # Create absolute value correlation matrix corr_matrix = X.corr().abs()

Select upper triangle of correlation matrix upper = corr_matrix.where(np.triu(np.ones(corr_matrix.shape), k=1).astype(np.bool))

Find index of feature columns with correlation greater than 0.95 to_drop = [column for column in upper.columns if any(upper[column] > 0.95)]

D	Sum of NUM_Medicare_BEN	Sum of NUM_Black_or_African_American_BEN	Sum of NUM_Asian_Pacific_Islander_BEN	Sum of NUM_Hispanic_BEN	Sum of NUM_American_IndianAlaska_Native_BEN	Sum of NUM_BEN_With_Race_Not_Elsewhere_Classified	Sum of Average_Age_of_BEN	Sum of PCT_MEDICARE	% Urban Pop	Density (P/mi2)	Children 0-18	Latitude	Longitude	Land Area
0	1820384.0	62311.0	76773.0	46525.0	147917.0	23372.0	996.298679	10.069041	66.0	1.2863	181405.17	61.370716	-152.404419	570665.0 §
1	10804823.0	1549811.0	30624.0	65500.0	5556.0	58660.0	3967.220634	51.254704	59.0	96.9221	1105570.08	32.806671	-86.791130	50644.0
2	15892716.0	1334245.0	19642.0	108428.0	62782.0	61250.0	3928.834167	94.570949	56.2	58.4030	686482.50	34.969704	-92.373123	52030.0
4	10786064.0	221183.0	61840.0	689880.0	179818.0	114903.0	1009.367955	14.075942	89.8	64.9550	1744612.56	33.729759	-111.431221	113595.0
5	42579588.0	2072012.0	3276415.0	5674776.0	113871.0	562214.0	4001.853612	63.398334	95.0	256.3727	9481941.36	36.116203	-119.681564	155766.0

X.info()

C+ <class 'pandas.core.frame.DataFrame'>
Int64Index: 51 entries, 0 to 55

Data	columns (total 38 columns):		
#	Column	Non-Null Count	Dtype
0	Sum of NUM Medicare BEN	51 non-null	float64
1	Sum of NUM Black or African American BEN	51 non-null	float64
2	Sum of NUM Asian Pacific Islander BEN	51 non-null	float64
3	Sum of NUM Hispanic BEN	51 non-null	float64
4	Sum of NUM American IndianAlaska Native BEN	51 non-null	float64
5	Sum of NUM BEN With Race Not Elsewhere Classified		float64
6	Sum of Average Age of BEN	51 non-null	float64
7	Sum of PCT MEDICARE	51 non-null	float64
8	% Urban Pop	51 non-null	float64
9	Density (P/mi2)	51 non-null	float64
10	Children 0-18	51 non-null	float64
11	Latitude	51 non-null	float64
12	Longitude	51 non-null	float64
13	Land Area	51 non-null	float64
14	Water Area	51 non-null	float64
15	Mean Flevation	51 non-null	float64
16	Highest Elevation	51 non-null	float64
17	Lowest elevation	51 non-null	float64
18	Number of bordering states	51 non-null	float64
19	On Coast	51 non-null	float64
20	Borders Another Country	51 non-null	float64
21	Captial Land Area	51 non-null	float64
22	Capital Water Area	51 non-null	float64
23	Capital Mean Elevation	51 non-null	float64
24	Capital is the Largest City	51 non-null	float64
25	Became a State	51 non-null	float64
26	DaysSinceStayatHomeOrder	51 non-null	float64
27	DaysSinceFirstPositive	51 non-null	float64
28	DaysSinceTestStart	51 non-null	float64
29	Log10Pop	51 non-null	float64
30	DaysSinceInfection	51 non-null	float64
31	Children0-18	51 non-null	float64
32	State Area Ratio	51 non-null	float64
33	Elevation Ratio	51 non-null	float64
34	Capital Area Ratio	51 non-null	float64
35	Boundaries	51 non-null	float64
36	Latitude Difference to State Capital	51 non-null	float64
37	Longitude Difference to State Capital	51 non-null	float64
dtyp	es: float64(38)		
memo	ry usage: 15.5 KB		

C+	Sum of NUM_Medicare_BEN	Sum of NUM_Black_or_African_American_BEN	Sum of NUM_Asian_Pacific_Islander_BEN	Sum of NUM_Hispanic_BEN	Sum of NUM_American_IndianAlaska_Native_BEN	Sum of NUM_BEN_With_Race_Not_Elsewhere_Classified	Sum of Average_Age_of_BEN	Sum of PCT_MEDICARE	% Urban Pop	Density (P/mi2)	Children 0- 18	Latitude	Longitu
count	5.100000e+01	5.100000e+01	5.100000e+01	5.100000e+01	51.000000	51.000000	51.000000	51.000000	51.000000	51.000000	5.100000e+01	51.000000	51.0000
mean	1.038431e+07	9.464777e+05	1.411691e+05	5.310095e+05	38355.372549	86379.019608	3535.191553	52.223011	74.107843	431.560508	1.486858e+06	39.464823	-93.3390
std	1.311026e+07	1.274593e+06	4.722330e+05	1.629961e+06	87337.002647	114765.665446	2518.178494	46.520870	14.885481	1647.225920	1.764935e+06	6.069546	19.2882
min	1.655870e+05	2.960000e+02	1.660000e+02	4.130000e+02	0.000000	1693.000000	70.002893	0.972106	38.700000	1.286300	1.160123e+05	21.094318	-157.4983
25%	2.252305e+06	5.366600e+04	6.445500e+03	3.101950e+04	2980.500000	17674.500000	1542.140834	13.385073	65.400000	50.604850	3.984744e+05	35.688955	-102.547!
50%	6.272609e+06	3.156040e+05	2.579200e+04	1.042170e+05	7061.000000	58660.000000	3578.360041	51.254704	74.200000	108.049700	1.013513e+06	39.849426	-89.616
75%	1.471830e+07	1.547566e+06	7.063400e+04	2.005865e+05	28506.500000	100449.000000	5214.099778	78.017975	87.550000	223.983100	1.719581e+06	43.041292	-78.988°
max	7.644909e+07	7.011107e+06	3.276415e+06	1.007620e+07	560433.000000	562214.000000	13644.965980	219.756971	100.000000	11814.541000	9.481941e+06	61.370716	-69.3819

```
Double-click (or enter) to edit
        # Train/validate split: random 75/25% train/validate split.
from sklearn.model_selection import train_test_split
       X_train, X_val, y_train, y_val = train_test_split(X, y, test_size = 0.25, random_state = 42)
       X_train.shape, y_train.shape, X_val.shape, y_val.shape
           Γ» ((38, 38), (38,), (13, 38), (13,))
          Б
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Density Children 0-
(P/mi2) 18 Latitude Longitu
                                                   Sum of Su
                                                                                                                                                                     3.800000e+01
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                                                                                                                                                                        1.001560e+06
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                                                          9.963253e+06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2056.853166 40.781616 14.338667 1899.378670 1.707713e+06 6.596259 20.5344
                                                                                                                 1.001560e+06 5.33705e+05
2.689000e+03 4.580000e+02
4.893450e+04 1.427175e+04
5.120990e+05 3.068000e+04
1.560497e+06 9.455175e+04
3.265865e+06 3.276415e+06
                                                                                                                                                                                                                                                                                                                                            2.622000e+03
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                             25%
                                                          2.518838e+06
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                                                                                                                                                                                                                                                                                                                                              1.983508e+05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5194.290279 79.721735 87.975000 249.285600 1.728603e+06 43.183955 -80.2768
                                                           4.257959e+07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        562214.000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7981.626181 185.232318 100.000000 11814.541000 9.481941e+06 61.370716 -69.3819
       # Optimizing Hyperparameters for Random Forest Regressor
from sklearn.model_selection import GridSearchCV
from sklearn.ensemble import RandomForestRegressor
        # Parameters to fit
       max depth = [2, 3, 4]
n_estimators = [35, 36, 37]
min_samples_split= [1.5, 2, 2.5]
min_samples_split= [1.5, 2, 2.5]
min_samples_split= [3.5, 4, 4.5]
max_leaf_nodes = [None]
max_features = ['auto']
ccp_ajha= [6, 0, 0.00625, 0.0125]
min_weight_fraction_leaf = [0.0, 0.00625, 0.0125]
       hyperF = dict(n_estimators = n_estimators, max_depth = max_depth,
min_samples_split = min_samples_split,
min_samples_leaf = min_samples_leaf,
max_leaf_nodes = max_leaf_nodes,
max_features = max_features,
ccp_alpha=ccp_alpha,
min_weight_fraction_leaf=min_weight_fraction_leaf)
       # Output best accuracy and best parameters
print('The score achieved with the best parameters = ', gridF.best_score_, '\n')
print('The parameters are:', gridF.best_params_)
                     Fatting 3 folds for each of 729 candidates, totalling 2187 fits

[Farallel(n_jobs-1)]: Using backend oilyaSackend with 2 concurrent workers.

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.4s

[Farallel(n_jobs-1)]: Done 4 tasks | elapsed: 1.4s

[Farallel(n_jobs-1)]: Done 6 tasks | elapsed: 1.6s

[Farallel(n_jobs-1)]: Done 9 tasks | elapsed: 1.6s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.6s

[Farallel(n_jobs-1)]: Batch computation to fast (0.1853.) Setting batch_size-2.

[Farallel(n_jobs-1)]: Batch computation to fast (0.1858.) Setting batch_size-3.

[Farallel(n_jobs-1)]: Batch computation to fast (0.1858.) Setting batch_size-3.

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.8s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 3.5s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.6s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.6s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.2s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.8s

[Farallel(n_jobs-1)]: Done 1 tasks | elapsed: 1.8
                          The parameters are: {'ccp_alpha': 0.0, 'max_depth': 3, 'max_features': 'auto', 'max_leaf_nodes': None, 'min_samples_leaf': 4, 'min_samples_split': 2, 'min_weight_fraction_leaf': 0.0, 'n_estimators': 36} (Parallel(n_jobs-1)): Done 2187 out of 2187 | elapsed: 43.35 finished
        !pip install category_encoders==2.0.0
          Collecting category encoders=2.0.8
Downloading https://files.opthombosteck.org/mackages/fielal/f792f144f33be78afeb86fa78478e8284a64263a1c89blef54e678841e/category_encoders=2.0.8-my2_my1_none-any_whl. (87k8)

**Requirement already satisfied: pathsys=0.4.1 in /usr/local/lib/python3.6/dist-packages (from category_encoders=2.0.0) (0.5.1)

*Requirement already satisfied: pandsys=0.2.1 in /usr/local/lib/python3.6/dist-packages (from category_encoders=2.0.0) (0.2.2.2.pst1)

*Requirement already satisfied: statismodels-yo-0.6.1 in /usr/local/lib/python3.6/dist-packages (from category_encoders=2.0.0) (0.10.2.2.2.pst1)

*Requirement already satisfied: statismodels-yo-0.6.1 in /usr/local/lib/python3.6/dist-packages (from category_encoders=2.0.0) (0.10.2.2.2.pst1)

*Requirement already satisfied: statismodels-yo-0.6.1 in /usr/local/lib/python3.6/dist-packages (from category_encoders=2.0.0) (0.11.2)

*Requirement already satisfied: impury=1.113 in /usr/local/lib/python3.6/dist-packages (from packages) (1.10.2.2)

*Requirement already satisfied: pyth=20217.0 in /usr/local/lib/python3.6/dist-packages (from pandsos=0.2.1.>category_encoders=2.0.0) (2.10.1)

*Requirement already satisfied: pyth=20217.0 in /usr/local/lib/python3.6/dist-packages (from pandsos=0.2.1.>category_encoders=2.0.0) (2.10.1)

*Requirement already satisfied: pyth=20217.0 in/usr/local/lib/python3.6/dist-packages (from pandsos=0.2.1.>category_encoders=2.0.0) (2.10.1)

*Requirement already satisfied: pyth=20217.0 in/usr/local/lib/python3.6/dist-packages (from scikit-learn=0.20.0->category_encoders=2.0.0) (0.11.1)

*Requirement already satisfied: pyth=20217.0 in/usr/local/lib/python3.6/dist-packages (from scikit-learn=0.20.0->category_encoders=2.0.0) (0.18.1)

*Requirement already satisfied: pyth=20217.0 in/usr/local/lib/python3.6/dist-packages (from scikit-learn=0.20.0->category_encoders=2.0.0) (0.18.1)
        from sklearn.ensemble import RandomForestRegressor
from sklearn.pipeline import make_pipeline
import category_encoders as ce
from sklearn.impute import SimpleImputer
Import citegory_
from sklearn.impute import s...,
from sklearn.impute import s...,
pipeline! = make_pipeline(
ce.oneMotEncoder(use_cat_names=True),
SimpleImputer(strategy='mean'),
RandomForestRegressor(bootstrap=True, ccp_alpha=0.0, criterion='mse',
mak_depth=3, max_features='auto', max_leaf_nodes=Nor
max_samples=None, min_impurity_decrease=0.0,
min_impurity_pititsome, min_sample_leaf=4,
min_impurity_pititsome, min_sample_leaf=4,
min_stimators=36, n_inbs=None, onb_score=False,
random_state=0, verbose=0, warm_start=False))

---
          # Get the model's training accuracy
print("Training Accurary: R^2 = ", pipeline1.score(X_train,y_train))
          # Get the model's validation accuracy
print('Validation Accuracy: R^2 = ', pipelinel.score(X_val, y_val))
        print(pipeline1.steps[2][1].feature_importances_)
          D
```

```
Feature Importances = [1.5806229e-03 5.3056745e-02 4.17410911e-03 1.59284318e-01 [1.5806229e-03 5.3056745e-02 4.17410911e-03 1.59284318e-01 2.6084125e-02 2.17672192e-02 7.81822081e-03 2.49720475e-02 2.90624300e-03 2.50540641e-02 4.59509372e-04 2.49941071e-03 1.1768719e-01 1.52212119-04 4.59509372e-04 2.49941071e-03 6.00000000e-01 1.52212109-01 4.50509372e-04 0.49941071e-03 6.000000000e-02 7.1827258e-0 5.38442596e-03 6.00000000e-00 6.00000000e-00 6.20312568e-0 5.38442596e-03 6.00000000e-00 6.00000000e-00 6.00000000e-00 7.38272568e-03 3.5662558e-03 3.56252545e-02 3.6475050e-03 6.12666322e-03]
         # Plot of feature importances from pure Random Forest Regressor

Zmatplotlib inline

import matplotlib.pyplot as plt

# Get feature importances

encoder = pipelinel.named_steps['onehotencoder']

encoded = encoder.transform(X_train)

rf = pipelinel.named_steps['randomforestregressor']

importances = pd.Series(rf.feature_importances_, encoded.columns)

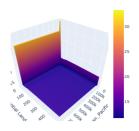
# Plot feature importances
            n = 12
plt.figure(figsize=(10,n/2))
plt.title(f'Top {n} features pipeline1')
importances1.sort_values()[-n:].plot.barh(color='grey');
                                                                                            Sum of NUM Hispanic BEN
                                               Sum of NUM_Black_or_African_American_BEN
                                                                                                                                      Boundaries -
                                                                                         Number of bordering states -
                                     Sum of NUM_American_IndianAlaska_Native_BEN
                                                                                                                                          Log10Pop
                                                                                                                                                                                                                                                                        0.075
                                                                                                                                                                                                                                                                                                         0.100
                                                                                                                                                                                                                                                                                                                                           0.125
                                                                                                                                                                                                                                                                                                                                                                               0.150
                                                                                                                                                                                                                                                                                                                                                                                                              0.175
         # Generate validation curves
Tamatpollib inline
Import numpy a mapping to pay point
Import numpy and point
Import numpy and import validation_curve
pipeline2 = make_pipeline(
ce.ordinaincoder(),
SimpleImport(),
Annotion restRegressor()
           depth = range(1, 10, 2)
train_scores, val_scores = validation_curve(
pipeline2, X_train, y_train,
param_name='randomforestregressor_max_depth',
param_rangedepth,
cv=3,
n_jobs=1
         plt.figure(dpi=150)
plt.plot(depth, np.mean(train_scores, axis=1), color='blue', label='training error')
plt.plot(depth, np.mean(val_scores, axis=1), color='red', label='validation error')
plt.title('validation (turve')
plt.vlabel('model complexty: RandomForestClassifier max_depth')
plt.vlabel('model complexty: RandomForestClassifier max_depth')
plt.vlabel('model score: Accuracy')
plt.vlabel('model score: Accuracy')
                                                                                                                                                                                                                           Validation Curve
                                                                       0
                                        score: Accuracy
                                                                 -2
                                                                 -4
                                                               -6
                                        mode
                                                             -8
                                                                                                                        training error
                                                                                                                        - validation error
                                                        -10
                                                                                                                         model complexity: RandomForestClassifier max_depth
# Get drop-column importance
column = 'Captial Land Area'
            # Fit without column
pipeline3.fit(X_train.drop(columns=column), y_train)
score_without = pipeline3.score(X_val.drop(columns=column), y_val)
print(f'Validation Accuracy without {column}: {score_without}')
            # Compare the error with & without column
print(f'Drop-Column Importance for {column}: {score_with - score_without}')
              Validation Accuracy without Captial Land Area: 0.10771758680677601
Validation Accuracy with Captial Land Area: 0.060388620867604
Drop-Column Importance for Captial Land Area: -0.03832896593917201
           # Using Eli5 library which does not work with pipelines
transformers = make_pipeline(
   ce.OneNotEncoder(use_cat_names=True),
   SimpleImputer(strategy='most_frequent')
                                                       RandomForestRegressor(bootstrap=True, ccp_alpha=0, criterion='mse', max_depth=3, max_featrues='auto', max_leaf_node=None, max_samples=None, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.0, min_lupurity_decrease=0.00_cccrease=0.0, min_lupurity_decrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_ccccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0.00_cccrease=0
```

```
model1.fit(X train transformed, y train)
                                                                               ((hootstrap=True, ccp_alpha=0, criterion='msc', max_depth=3, max_features='auto', max_leaf_nodes=Mone, max_sample=shome, min_impurity_decrease=0.0, min_impurity_split=Mone, min_samples_leaf=4, min_samples_leaf=2, min_samples_leaf=3, m_jobs=Mone, ool_score=false, n_estimators=36, m_jobs=Mone, ool_score=false, random_state=0, verbose=0, warm_start=False)
    # Get permutation importances
! pip install elis
from elis.sklearn import PermutationImportance
import elis
                model1,
scoring='r2',
n_iter=2,
random_state=42
     permuter.fit(X_val_transformed, y_val)
feature_names = X_val.columns.tolist()
     eli5.show_weights(
                b.snow_weignts(
permuter,
top=None, # show permutation importances for all features
feature_names=feature_names
Interior of the control of the contr
     from sklearn.metrics import mean_squared_error, r2_score
      # Coefficient of determination r2 for the training set
       pipeline_score = permuter.score(X_train_transformed,y_train)
print("Coefficient of determination r2 for the training set.: ", pipeline_score)
     # Coefficient of determination r2 for the validation set
pipeline_score = permuter.score(X_val_transformed,y_val)
print("Coefficient of determination r2 for the validation set.: ", pipeline_score)
      # The mean squared error
y_pred = permuter.predict(X_val_transformed)
print("Mean squared error: %.2f"% mean_squared_error(y_val, y_pred))
       Coefficient of determination r2 for the training set.: 0.3013954634238487
Coefficient of determination r2 for the validation set.: 0.06938862086760
Mean squared error: 530.47
    # Captial Land Area continues to be of importance
# Use importances for feature selection
print('Shape before removing features:', X_train.shape)
      □ Shape before removing features: (38, 38)
    from sklearn.metrics import mean squared error, r2 score
    # Coefficient of determination r2 for the training set
pipeline_score = pipeline4.score(X_train,y_train)
print("Coefficient of determination r2 for the training set.: ", pipeline_score)
     # Coefficient of determination r2 for the validation set pipeline_score = pipeline4.score(X_val,y_val) print("Coefficient of determination r2 for the validation set.: ", pipeline_score)
    # The mean squared error
y_pred = pipelines.predict(X_val)
print("Mean squared error: %.2f"% mean_squared_error(y_val, y_pred))
     Coefficient of determination r2 for the training set.: 0.26984125703310236
Coefficient of determination r2 for the validation set.: 0.11859396791756194
Nean squared error: 502.42
```

```
# Plot of features
%matplotlib inline
import matplotlib.pyplot as plt
# Get feature importances
encoder = pipeline4.named_steps['onehotencoder']
encoded = encodec.transform(X_val)
rf = pipeline4.named_steps['randomforestregressor']
importance2 = pd.Series(rf.feature_importances__, encoded.columns)
n = 12
plt.figure(figsize=(10,n/2))
plt.title(f'Top (n) features pipeline4')
importances2.sort_values()[-n:].plot.barh(color='grey');
                             Capital Area Ratio
                                  Boundaries -
                     Sum of Average_Age_of_BEN
                             Captial Land Area
                               Density (P/mi2)
                      Sum of NUM_Medicare_BEN
                                    On Coast
                                                    0.025
                                                                         0.075
                                                                                    0.100 0.125
                                                               0.050
                                                                                                        0.150
                                                                                                                   0.175
!pip install pdpbox
     Collecting pdpbox
   model2.fit(X train, y train)
    RandomForestRegressor(bootstrap=True, ccp_alpha=0, criterion='mse', max_depth=3, max_features'=uto', max_leaf_nodes=None, amx_amplex=None, min_impurity_decrease=0.0, min_impurity_split=None, min_samples_leaf=4, min_samples_pilit=2, min_ueight_fraction_leaf=0, nestimators=36, n_jobs=None, odb_score=False, random_state=0, verbose=0, warm_start=False)
pdp_interact_plot(interaction, plot_type='grid', feature_names=features2);
                 PDP interact for "Sum of NUM Asian Pacific Islander BEN" and "Captial Land Area"
           55.23
            36.4
                  166 1092.334500.33 6285.0 7368.013614.025792.049567.0
Sum of NUM_Asian_Pacific_Islander_BEN
# A two feature partical dependence plot in 30 pdp = interaction.pdp.pivot_table(

values='preds',
columns=features2[0],
index=features2[1]
)[::-1] # Slice notation to reverse index order so y axis is ascending
import plotly.graph_objs as go
target = 'Value of a parameter'
yaxis=dict(title=features2[1]),
zaxis=dict(title=target)
fig = go.Figure(surface, layout)
fig.show()
```

C+



```
Collecting shaps=0.23.0

Downloading https://files.pythochosted.org/packages/56/86/88087621f7338edy2892a992a91c255ff955466663772edleae891c6/shap-0.21.0.tar.gg (182k8)

Downloading https://files.pythochosted.org/packages/56/86/88087621f7338edy2892a992a91c255ff955466663772edleae891c6/shap-0.21.0.tar.gg (182k8)

Requirement already satisfied: cutpy in /usr/local/lib/python3.6/dist-packages (from shaps=0.21.0) (1.13.3)

Requirement already satisfied: matplotlib in /usr/local/lib/python3.6/dist-packages (from shaps=0.23.0) (0.2.1)

Requirement already satisfied: matplotlib in /usr/local/lib/python3.6/dist-packages (from shaps=0.23.0) (0.3.1)

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 ! pip install shap==0.23.0
! pip install -I shap
                                          10.098 1748/s

tcting tdom4.25.0

mloading https://files.opthombated.org/markages/d9/49/958812e8ba18035f89c901fdfc2d42f8c95947df2d5eb3c7b258019fda8/tgde-4.66.d-py2_py3-none-any_whi (63k8)

"JNB 8.3980" j7kB 8.3980" j7kB 8.3980" j7kB 8.3980" j7kB 8.3980" j7kB 8.3980" j7kB 9.3980" j7kB 9.3980 j7kB
                   # Local Interpretation using SHAP (for prediction at State # = 4, row 32)
    explainer = shap.TreeExplainer(model2)
shap_values = explainer.shap_values(X_train)
i = 32
      shap.force_plot(explainer.expected_value, shap_values[i], features=X_train.loc[i], feature_names=X_train.columns)
                   Find Shapley Forces across the training sample i (i = 0 - 37)
 X_train_processed = processor.fit_transform(X_train)
column_names = X_train.columns
shap_values_array = pd.DataFrame(columns = column_names)
for i in range(len(y_train)):
    row = X_train.iloc[[i]]
    explainer = shap.TreeExplainer(model2)
    row_processed = processor.transform(row)
    shap.values_input = explainer.shap_values(row_processed)
    shap_values_array = np.concatenate((shap_values_array, shap_values_input), axis=0)
# Create a 3D plot of force as a function of state curve displacement from mean curve and features for validation sai 
# A two feature partical dependence plot in 3D 
import plotly graph_objs as go 
surface = go.Surface(x-column_names, 
yvg_train, 
z-shap_values_array)
```

```
yaxis=dict(title= 'Value of a for state'),
zaxis=dict(title= 'Shapley Force')
                fig = go.Figure(surface, layout)
            fig.show()
                D+
            # Recursive Feature Elimination
from sklearn.feature_selection import RFE, f_regression
from sklearn.model_selection import StratifiedKFold
            rfr = RandomForestRegressor(bootstrap=True, ccp_alpha=0, max_dest_modes=None, max_dest_modes=
              #Selecting 7 features turns out to give maximum validation accuracy number.selected_features = 7 rfe = REF(frf, n_features_to_select=number_selected_features, verbose =3) rfe.fit(X_train_y_train)
                              Fitting estimator with 14 features
Fitting estimator with 13 features
Fitting estimator with 17 features
Fitting estimator with 11 features
Fitting estimator with 10 features
Fitting estimator with 9 features.
Fitting estimator with 8 features.
FRE(estimator=RandomForestRegress)
                                               string estimator with 8 features.

E(estimator=RandomForestRegressor(bootstrag=True, ccp_alpha=0, criterion=mse', max_depth=1, max_dept
              rfe_support = rfe.get_support()
rfe_feature = X_train.loc[:,rfe_support].columns.tolist()
print(str(len(rfe_feature)), 'selected features')
              from sklearn.metrics import mean squared error, r2 score
            # Coefficient of determination r2 for the training set
pipeline_score = rfe.score(X_train,y_train)
print("Coefficient of determination r2 for the training set.: ", pipeline_score)
              # Coefficient of determination r2 for the validation set
pipeline_score = rfe.score(X_val,y_val)
print("Coefficient of determination r2 for the validation set.: ", pipeline_score)
            B The mean squared error
y_pred = rfc.predict(X_val)
print("Mean squared error: %.2f*% mean_squared_error(y_val, y_pred))
              Coefficient of determination r2 for the training set.: 0.27619181813448996
Coefficient of determination r2 for the validation set.: 0.14111881952867834
Mean squared error: 489.58
              # Retain only features with highest importance from RFE
X_train_rfe_select = X_train[rfe_feature]
X_val_rfe_select = X_val[rfe_feature]
print('Shape after removing features'', X_train_rfe_select.shape, X_val_rfe_select.shape)
                □ Shape after removing features: (38, 7) (13, 7)
# Coefficient of determination r2 for the training set
pipeline_score = pipelineS.score(X_train_rfe_select,y_train)
print("Coefficient of determination r2 for the training set.: ", pipeline_score)
            # Coefficient of determination r2 for the validation set
plpeline_score = pipelineS.score(X_val_rfe_select,y_val)
print("Coefficient of determination r2 for the validation set.: ", pipeline_score)
              # The mean squared error
y_pred = pipelines.predict(X_val_rfe_select)
print("Mean squared error: %.2f"% mean_squared_error(y_val, y_pred))
              Coefficient of determination r2 for the training set.: 0.27619181813440996
Coefficient of determination r2 for the validation set.: 0.14111881952867034
Mean squared error: 489.58
            pipeline5.fit(X_val_rfe_select, y_val)
# Plot of features
%matplotlib inline
import matplotlib.pyplot as plt
            # Get feature importances
encoder = pipelines.named_steps['onehotencoder']
encoded = encoder.transform(X_val_rfe_select)
rf = pipelines.named_steps['randomforestregressor']
importances3 = pd.Series(rf.feature_importances_, encoded.columns)
          # Plot feature importances
n = number_selected_features
plt.figure(figsize=(18,n/2))
plt.title("Top (n) features pipelines")
importances3.sort_values()[-n:].plot.barh(color='grey');
```

