

Life Expectancy data analysis

August 16, 2024

1 Data preprocessing before building the machine learning model

1.1 STEPS of preprocessing data

#step:1] import the necessary library #step:2] Read dataset #step:3] sanity check of data #step:4] exploratory data analysis(EDA) #step:5] missing value treatment #step:6] outlier treatment #step:7] duplicate garbag value treatment #step:8] noemalization #step:9] encoding od data

```
[32]: ##Import necessary library
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np
```

```
[33]: ##Read the data
data = pd.read_csv("Life Expectancy Data.csv")
```

```
[34]: ## find the top 3 rows
data.head(3)
```

```
[34]:      Country  Year  Status  Life expectancy  Adult Mortality  \
0  Afghanistan  2015  Developing             65.0             263.0
1  Afghanistan  2014  Developing             59.9             271.0
2  Afghanistan  2013  Developing             59.9             268.0

      infant deaths  Alcohol  percentage expenditure  Hepatitis B  Measles  ...  \
0                62     0.01             71.279624             65.0     1154  ...
1                64     0.01             73.523582             62.0     492  ...
2                66     0.01             73.219243             64.0     430  ...

      Polio  Total expenditure  Diphtheria  HIV/AIDS  GDP  Population  \
0      6.0              8.16             65.0      0.1  584.259210  33736494.0
1     58.0              8.18             62.0      0.1  612.696514   327582.0
2     62.0              8.13             64.0      0.1  631.744976  31731688.0

      thinness  1-19 years  thinness 5-9 years  \
0              17.2              17.3
1              17.5              17.5
```

2	17.7	17.7
	Income composition of resources	Schooling
0	0.479	10.1
1	0.476	10.0
2	0.470	9.9

[3 rows x 22 columns]

```
[35]: ##sanity check the data
data.shape
```

```
[35]: (2938, 22)
```

```
[36]: data.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 2938 entries, 0 to 2937
Data columns (total 22 columns):
#   Column                                Non-Null Count  Dtype
---  -
0   Country                               2938 non-null   object
1   Year                                  2938 non-null   int64
2   Status                               2938 non-null   object
3   Life expectancy                       2928 non-null   float64
4   Adult Mortality                       2928 non-null   float64
5   infant deaths                         2938 non-null   int64
6   Alcohol                              2744 non-null   float64
7   percentage expenditure                2938 non-null   float64
8   Hepatitis B                           2385 non-null   float64
9   Measles                               2938 non-null   int64
10  BMI                                   2904 non-null   float64
11  under-five deaths                     2938 non-null   int64
12  Polio                                 2919 non-null   float64
13  Total expenditure                     2712 non-null   float64
14  Diphtheria                            2919 non-null   float64
15  HIV/AIDS                              2938 non-null   float64
16  GDP                                    2490 non-null   float64
17  Population                            2286 non-null   float64
18  thinness 1-19 years                   2904 non-null   float64
19  thinness 5-9 years                    2904 non-null   float64
20  Income composition of resources        2771 non-null   float64
21  Schooling                             2775 non-null   float64
dtypes: float64(16), int64(4), object(2)
memory usage: 505.1+ KB
```

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

```
[37]: Country          0
      Year             0
      Status           0
      Life expectancy  10
      Adult Mortality  10
      infant deaths    0
      Alcohol          194
      percentage expenditure  0
      Hepatitis B      553
      Measles          0
      BMI              34
      under-five deaths  0
      Polio            19
      Total expenditure 226
      Diphtheria       19
      HIV/AIDS         0
      GDP              448
      Population       652
      thinness 1-19 years  34
      thinness 5-9 years  34
      Income composition of resources 167
      Schooling        163
      dtype: int64
```

```
[38]: data.isnull().sum()/data.shape[0]*100
```

```
[38]: Country          0.000000
      Year             0.000000
      Status           0.000000
      Life expectancy  0.340368
      Adult Mortality  0.340368
      infant deaths    0.000000
      Alcohol          6.603131
      percentage expenditure  0.000000
      Hepatitis B      18.822328
      Measles          0.000000
      BMI              1.157250
      under-five deaths  0.000000
      Polio            0.646698
      Total expenditure  7.692308
      Diphtheria       0.646698
      HIV/AIDS         0.000000
      GDP              15.248468
      Population       22.191967
      thinness 1-19 years  1.157250
      thinness 5-9 years  1.157250
      Income composition of resources  5.684139
```

```
Schooling
dtype: float64
```

```
5.547992
```

```
[39]: data.duplicated().sum()
```

```
[39]: 0
```

```
[40]: ##EDA
data.describe().T
```

```
[40]:
```

	count	mean	std \
Year	2938.0	2.007519e+03	4.613841e+00
Life expectancy	2928.0	6.922493e+01	9.523867e+00
Adult Mortality	2928.0	1.647964e+02	1.242921e+02
infant deaths	2938.0	3.030395e+01	1.179265e+02
Alcohol	2744.0	4.602861e+00	4.052413e+00
percentage expenditure	2938.0	7.382513e+02	1.987915e+03
Hepatitis B	2385.0	8.094046e+01	2.507002e+01
Measles	2938.0	2.419592e+03	1.146727e+04
BMI	2904.0	3.832125e+01	2.004403e+01
under-five deaths	2938.0	4.203574e+01	1.604455e+02
Polio	2919.0	8.255019e+01	2.342805e+01
Total expenditure	2712.0	5.938190e+00	2.498320e+00
Diphtheria	2919.0	8.232408e+01	2.371691e+01
HIV/AIDS	2938.0	1.742103e+00	5.077785e+00
GDP	2490.0	7.483158e+03	1.427017e+04
Population	2286.0	1.275338e+07	6.101210e+07
thinness 1-19 years	2904.0	4.839704e+00	4.420195e+00
thinness 5-9 years	2904.0	4.870317e+00	4.508882e+00
Income composition of resources	2771.0	6.275511e-01	2.109036e-01
Schooling	2775.0	1.199279e+01	3.358920e+00

	min	25%	50% \
Year	2000.00000	2004.000000	2.008000e+03
Life expectancy	36.30000	63.100000	7.210000e+01
Adult Mortality	1.00000	74.000000	1.440000e+02
infant deaths	0.00000	0.000000	3.000000e+00
Alcohol	0.01000	0.877500	3.755000e+00
percentage expenditure	0.00000	4.685343	6.491291e+01
Hepatitis B	1.00000	77.000000	9.200000e+01
Measles	0.00000	0.000000	1.700000e+01
BMI	1.00000	19.300000	4.350000e+01
under-five deaths	0.00000	0.000000	4.000000e+00
Polio	3.00000	78.000000	9.300000e+01
Total expenditure	0.37000	4.260000	5.755000e+00
Diphtheria	2.00000	78.000000	9.300000e+01
HIV/AIDS	0.10000	0.100000	1.000000e-01

GDP	1.68135	463.935626	1.766948e+03
Population	34.00000	195793.250000	1.386542e+06
thinness 1-19 years	0.10000	1.600000	3.300000e+00
thinness 5-9 years	0.10000	1.500000	3.300000e+00
Income composition of resources	0.00000	0.493000	6.770000e-01
Schooling	0.00000	10.100000	1.230000e+01

	75%	max
Year	2.012000e+03	2.015000e+03
Life expectancy	7.570000e+01	8.900000e+01
Adult Mortality	2.280000e+02	7.230000e+02
infant deaths	2.200000e+01	1.800000e+03
Alcohol	7.702500e+00	1.787000e+01
percentage expenditure	4.415341e+02	1.947991e+04
Hepatitis B	9.700000e+01	9.900000e+01
Measles	3.602500e+02	2.121830e+05
BMI	5.620000e+01	8.730000e+01
under-five deaths	2.800000e+01	2.500000e+03
Polio	9.700000e+01	9.900000e+01
Total expenditure	7.492500e+00	1.760000e+01
Diphtheria	9.700000e+01	9.900000e+01
HIV/AIDS	8.000000e-01	5.060000e+01
GDP	5.910806e+03	1.191727e+05
Population	7.420359e+06	1.293859e+09
thinness 1-19 years	7.200000e+00	2.770000e+01
thinness 5-9 years	7.200000e+00	2.860000e+01
Income composition of resources	7.790000e-01	9.480000e-01
Schooling	1.430000e+01	2.070000e+01

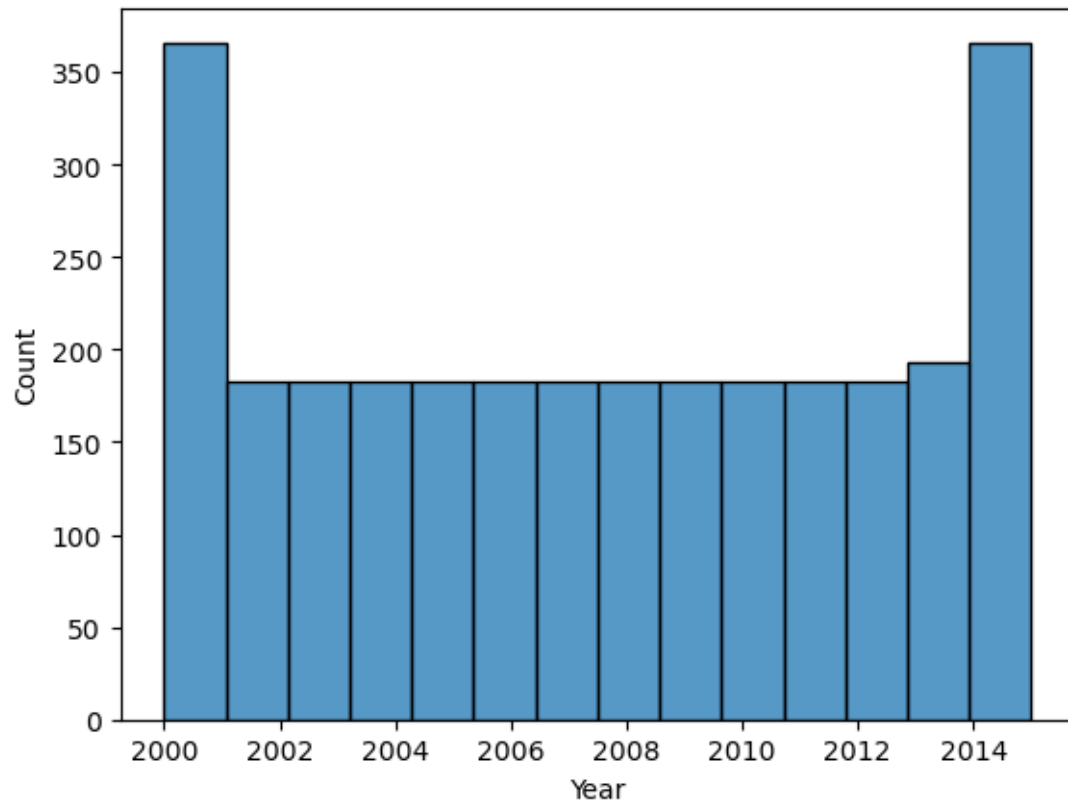
```
[41]: data.describe(include="object")
```

```
[41]:
```

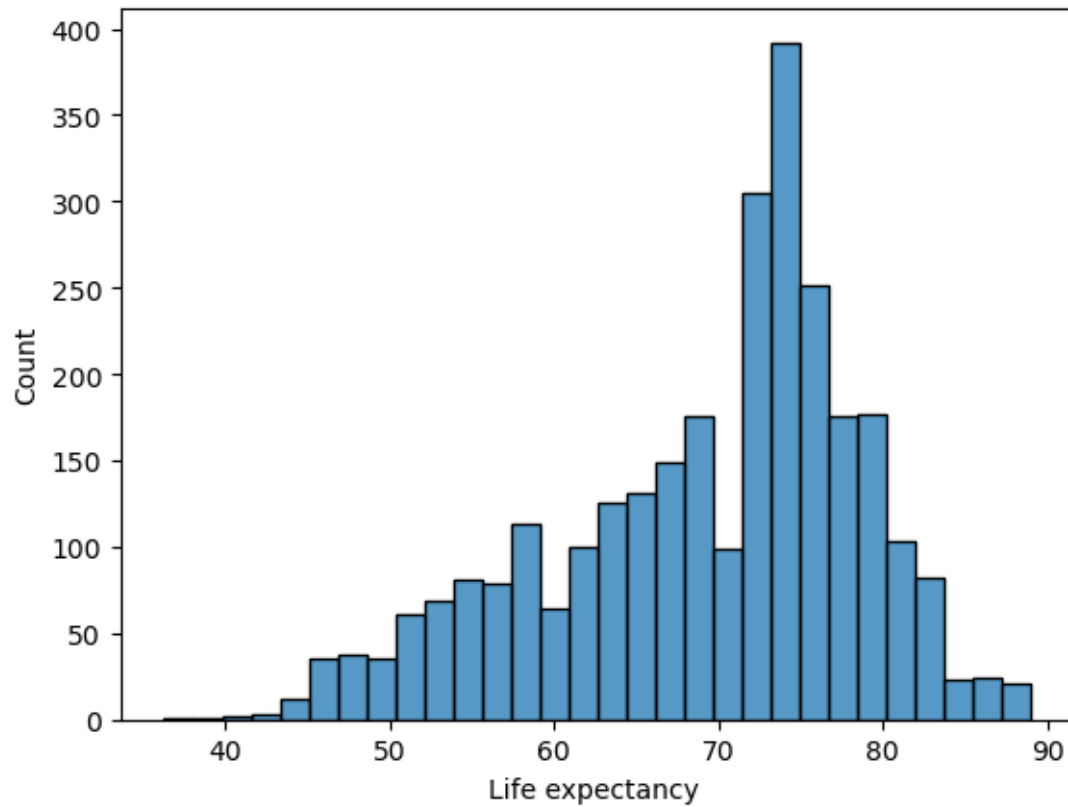
	Country	Status
count	2938	2938
unique	193	2
top	Afghanistan	Developing
freq	16	2426

```
[42]: ## histogram to understand the destrribution
for i in data.select_dtypes(include="number").columns:
    sns.histplot(data=data, x=i) # Corrected data argument
    plt.show() # Correctly close the function call
```

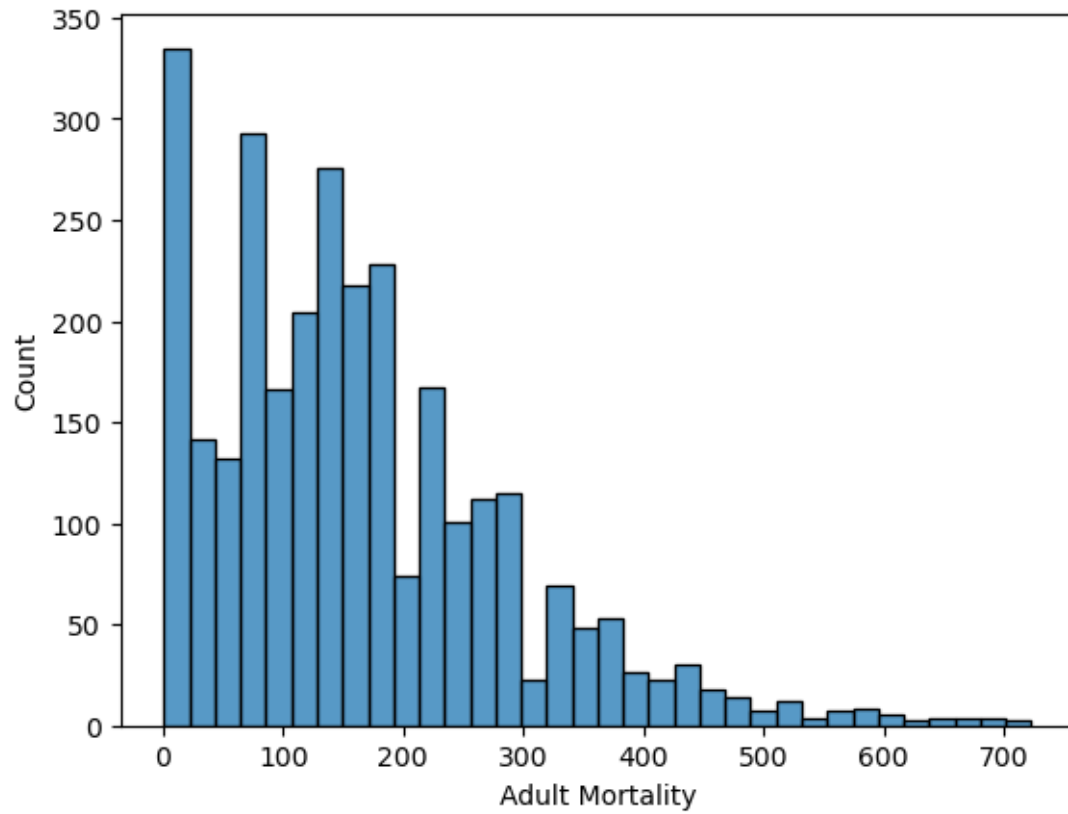
```
C:\Users\Vikas\anaconda3\lib\site-packages\seaborn\_oldcore.py:1119:
FutureWarning: use_inf_as_na option is deprecated and will be removed in a
future version. Convert inf values to NaN before operating instead.
with pd.option_context('mode.use_inf_as_na', True):
```



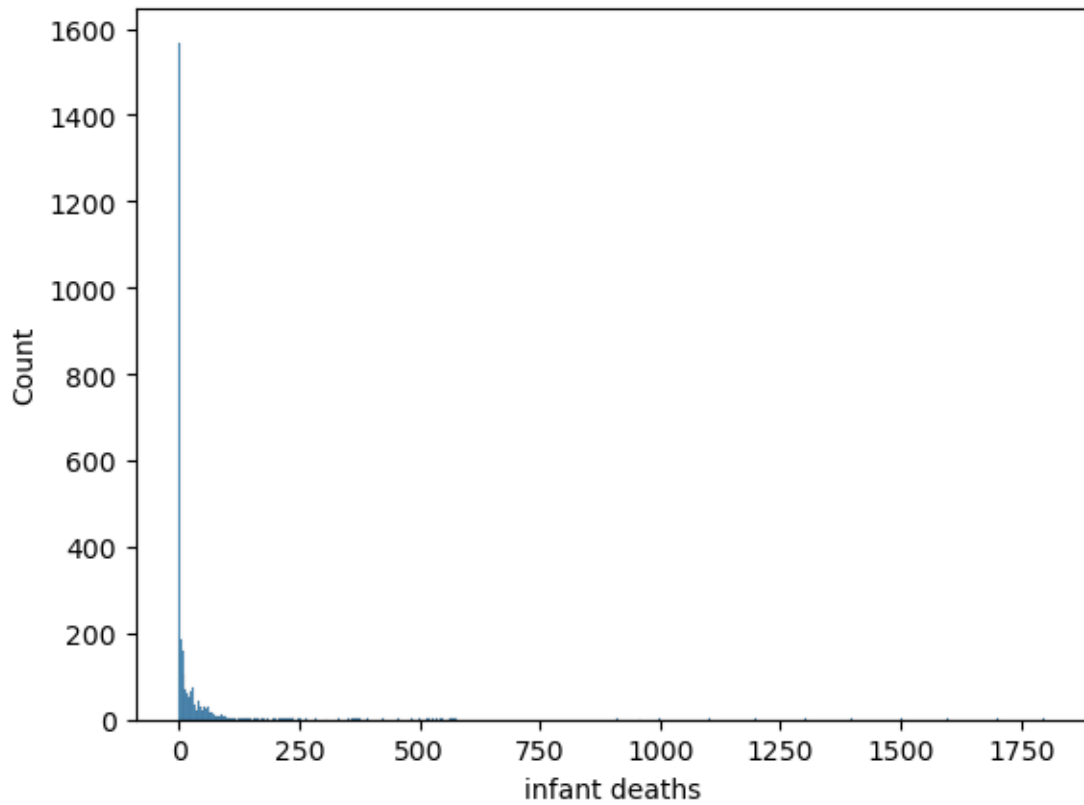
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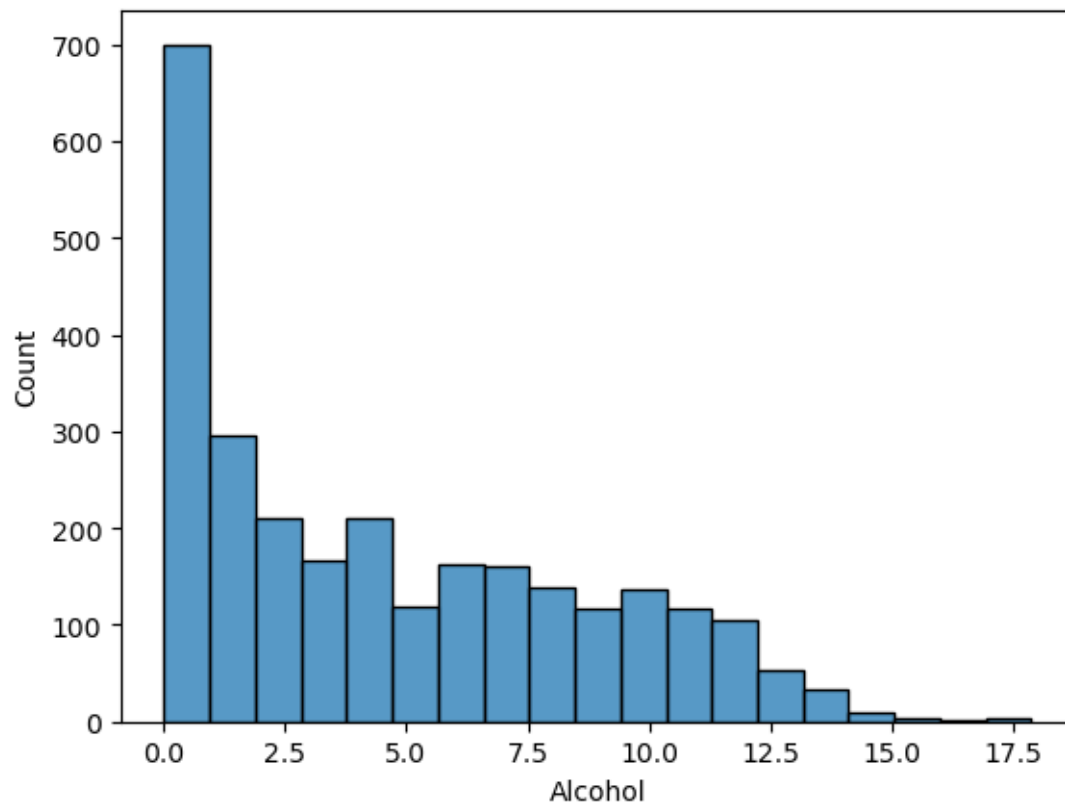
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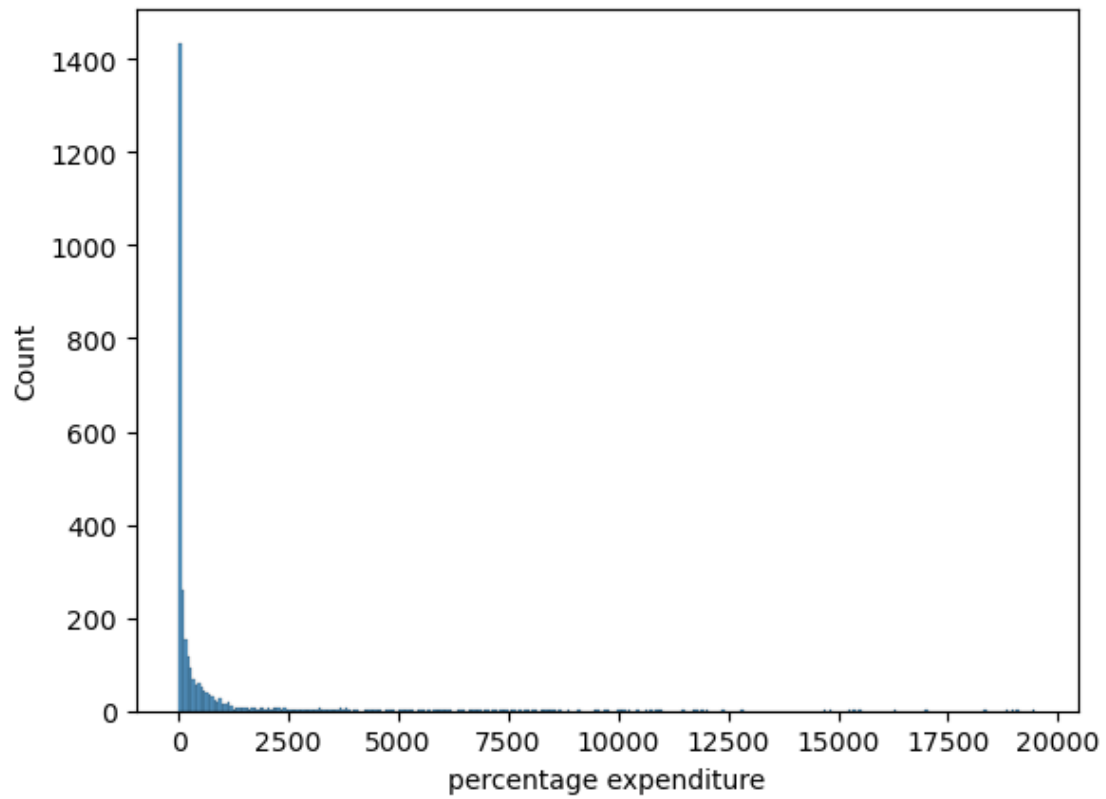
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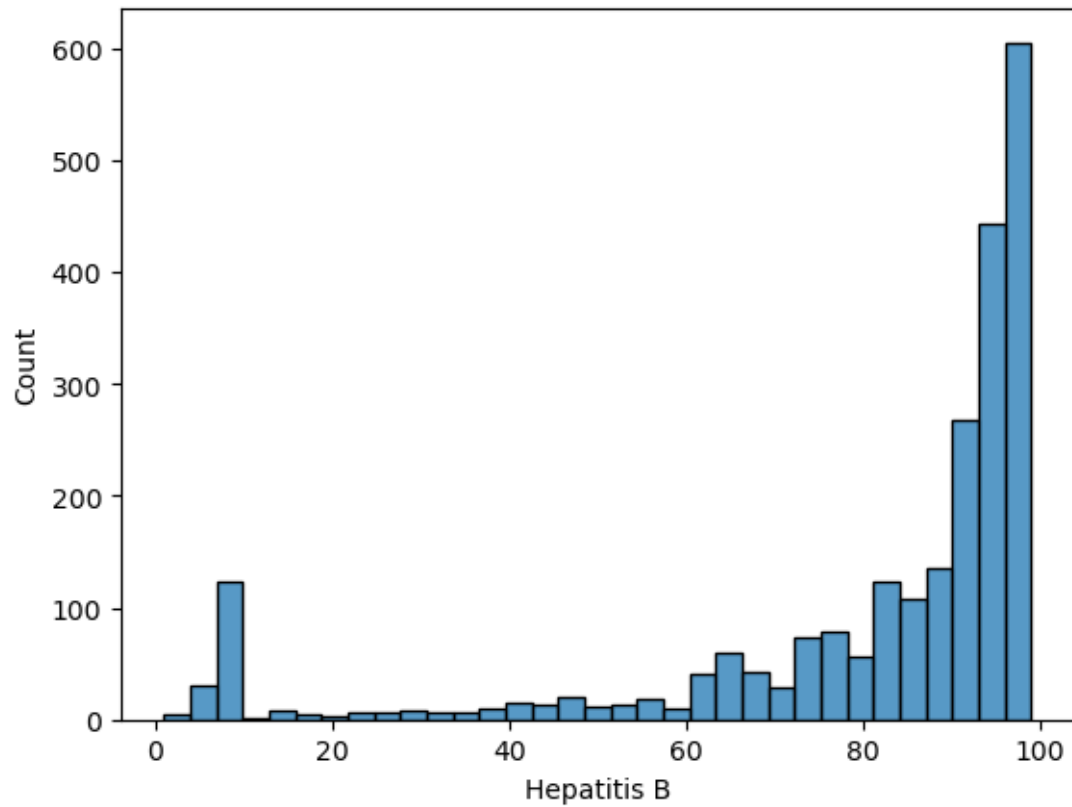
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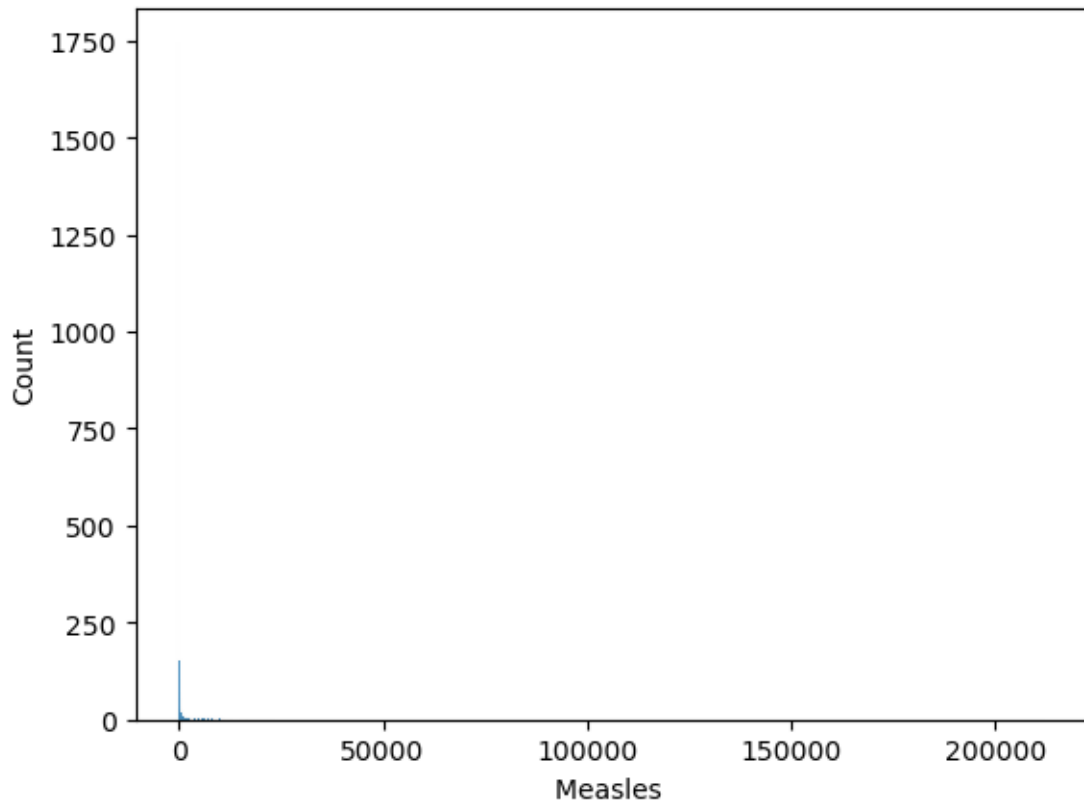
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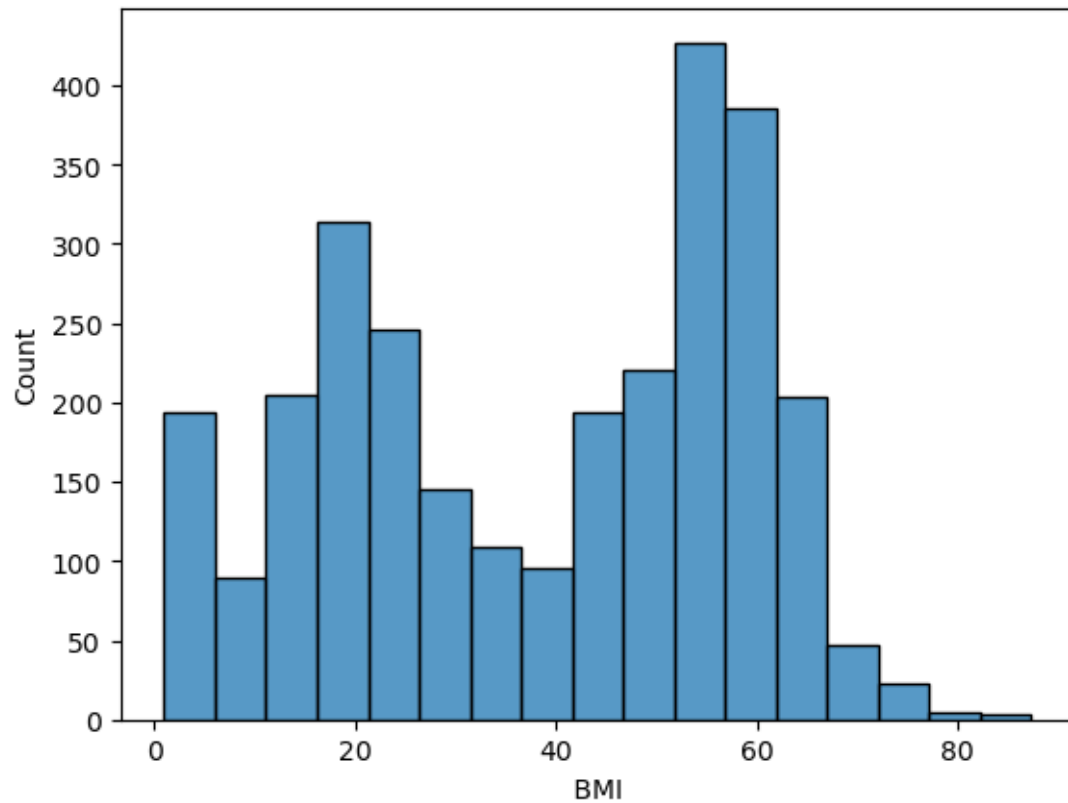
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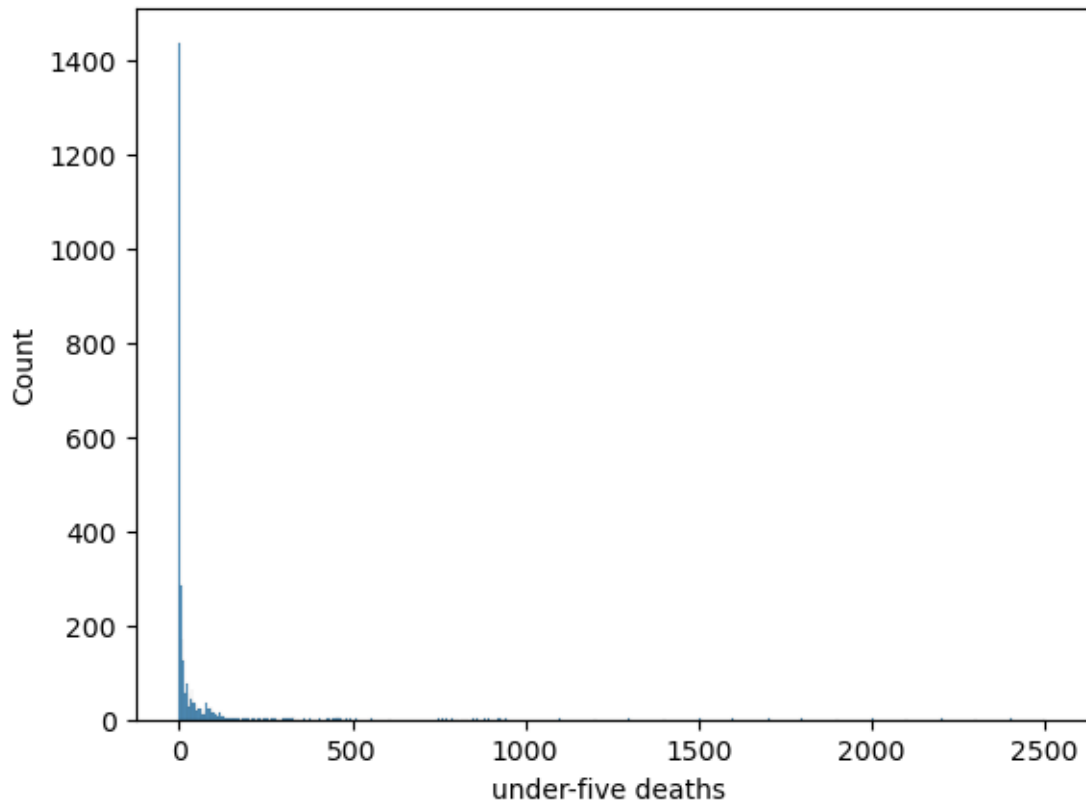
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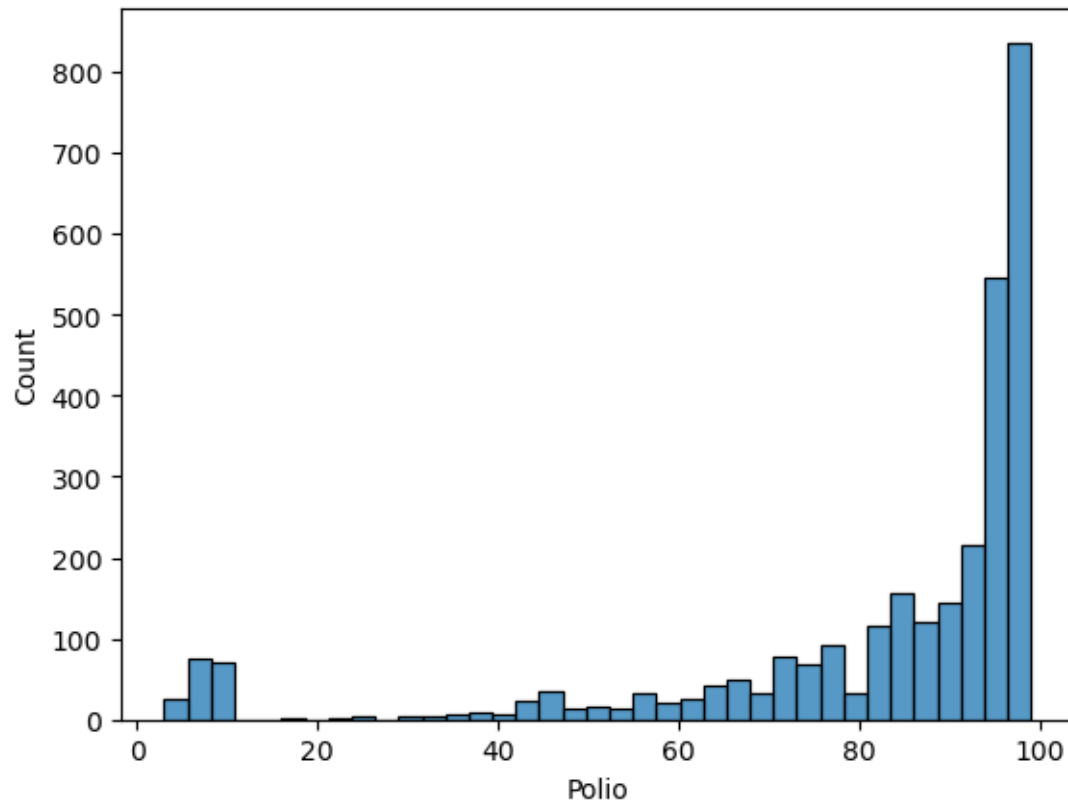
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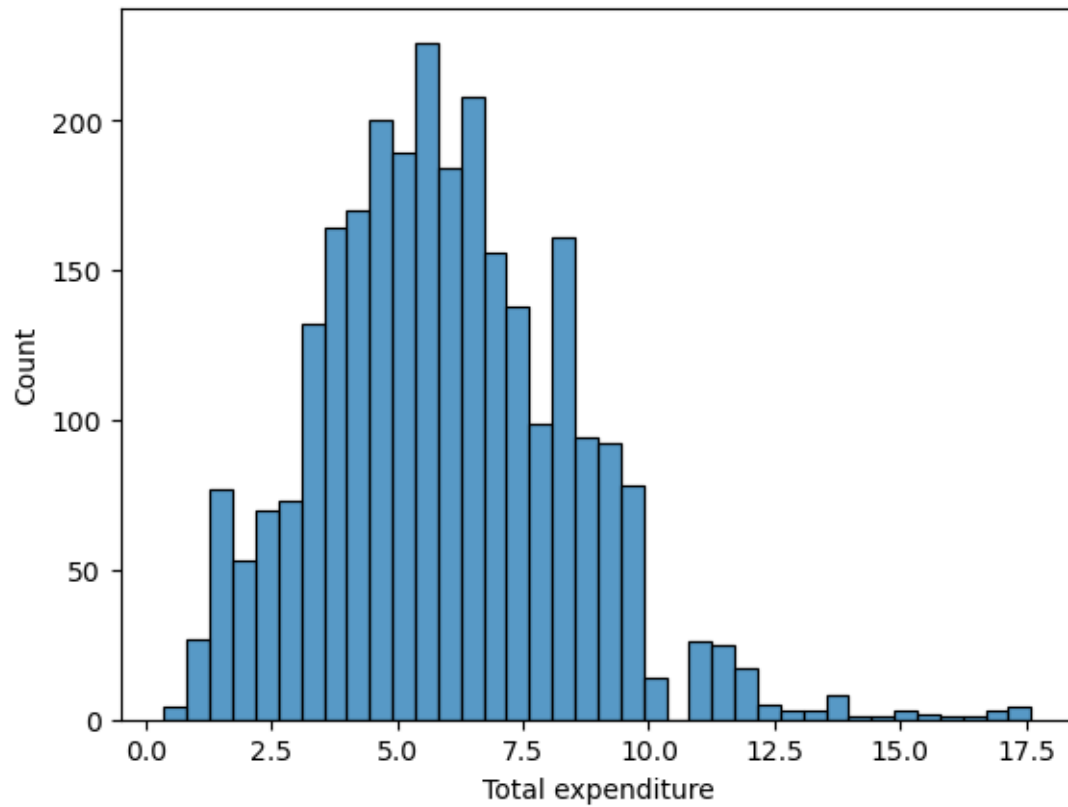
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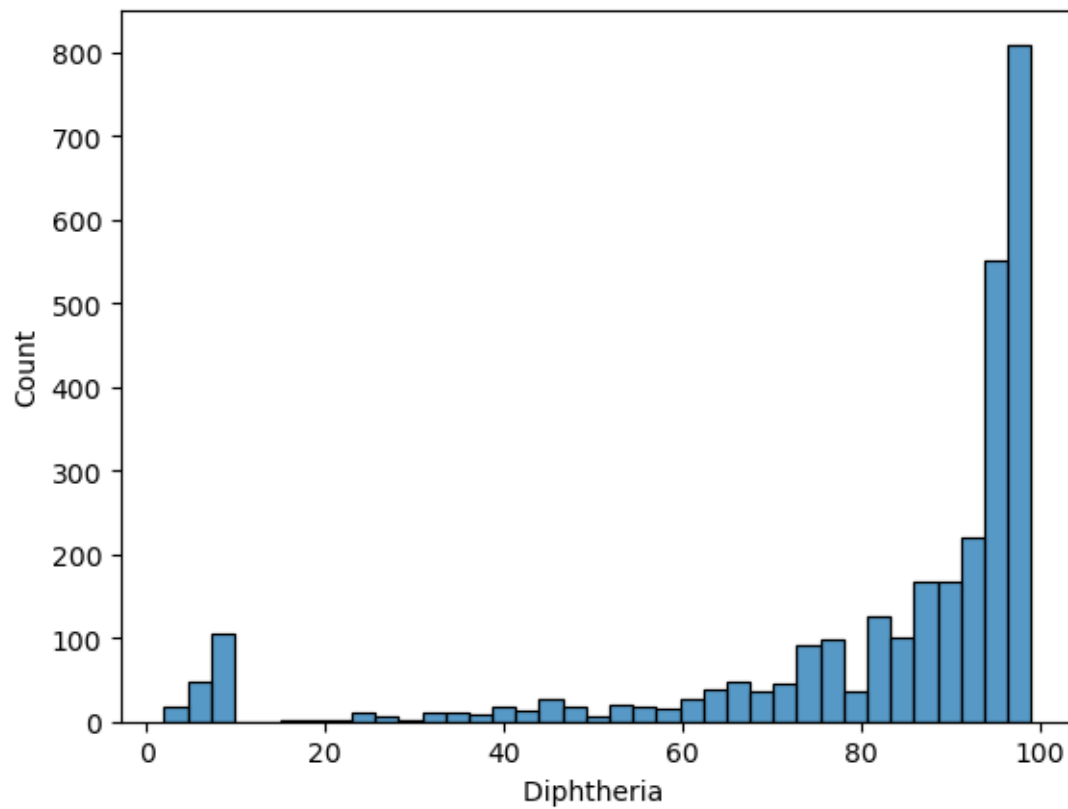
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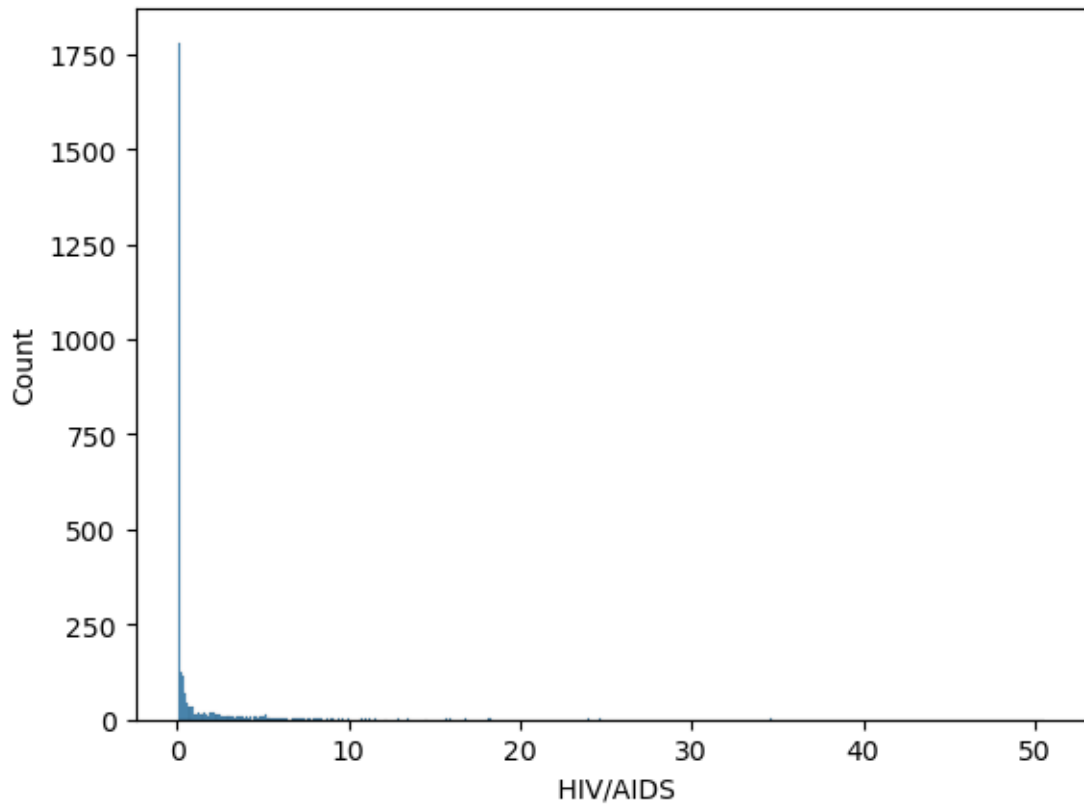
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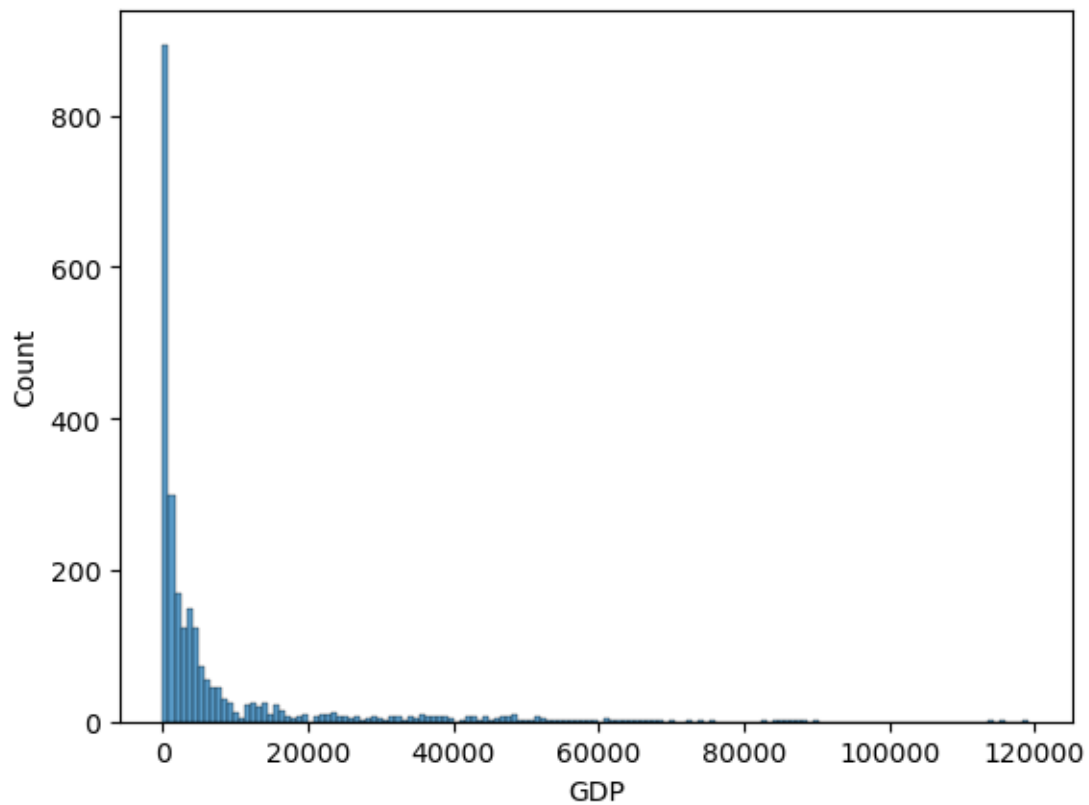
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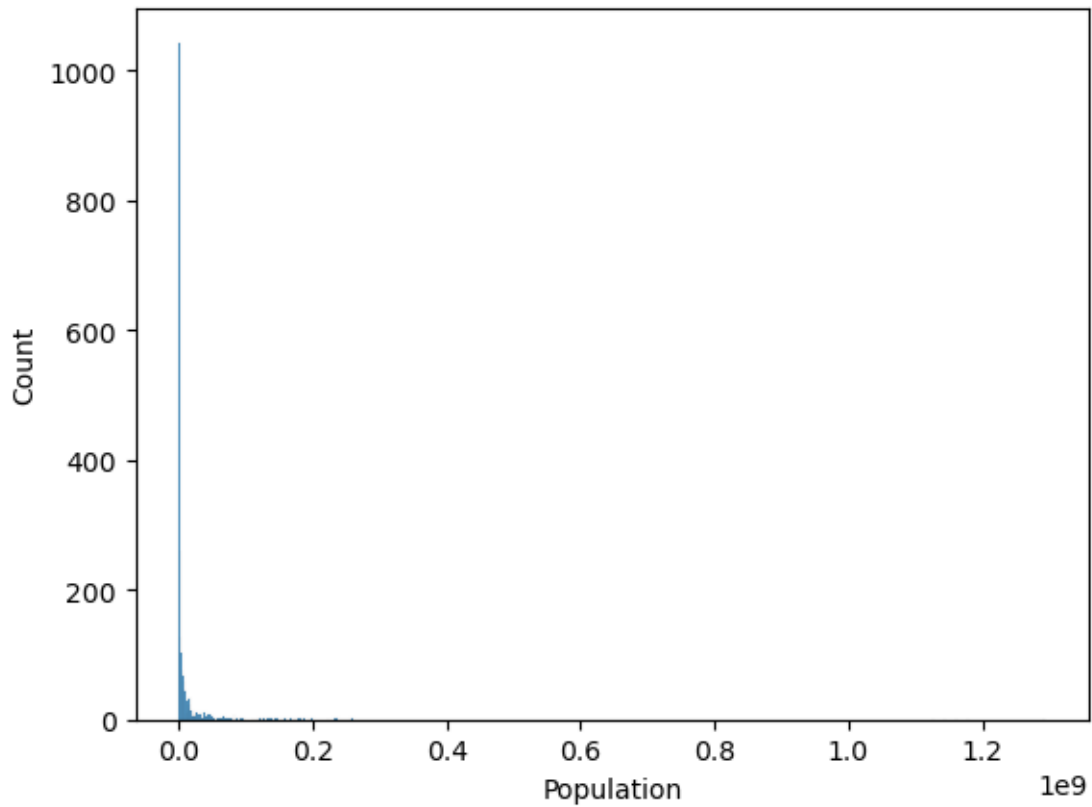
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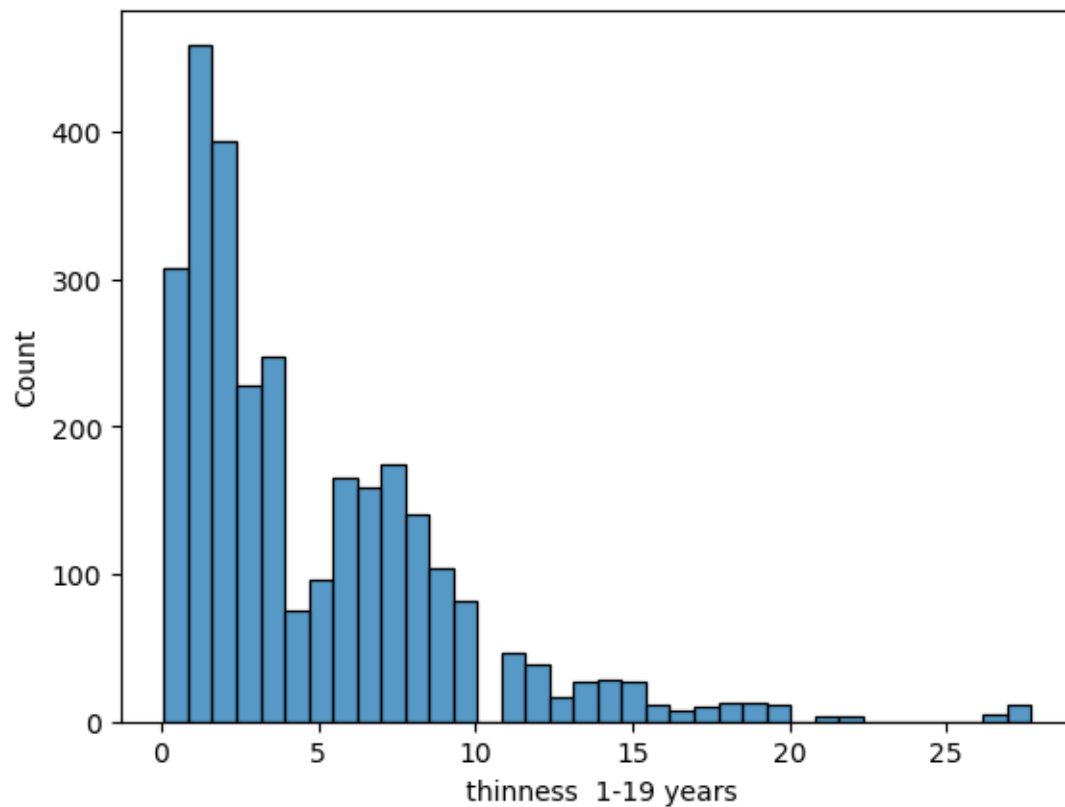
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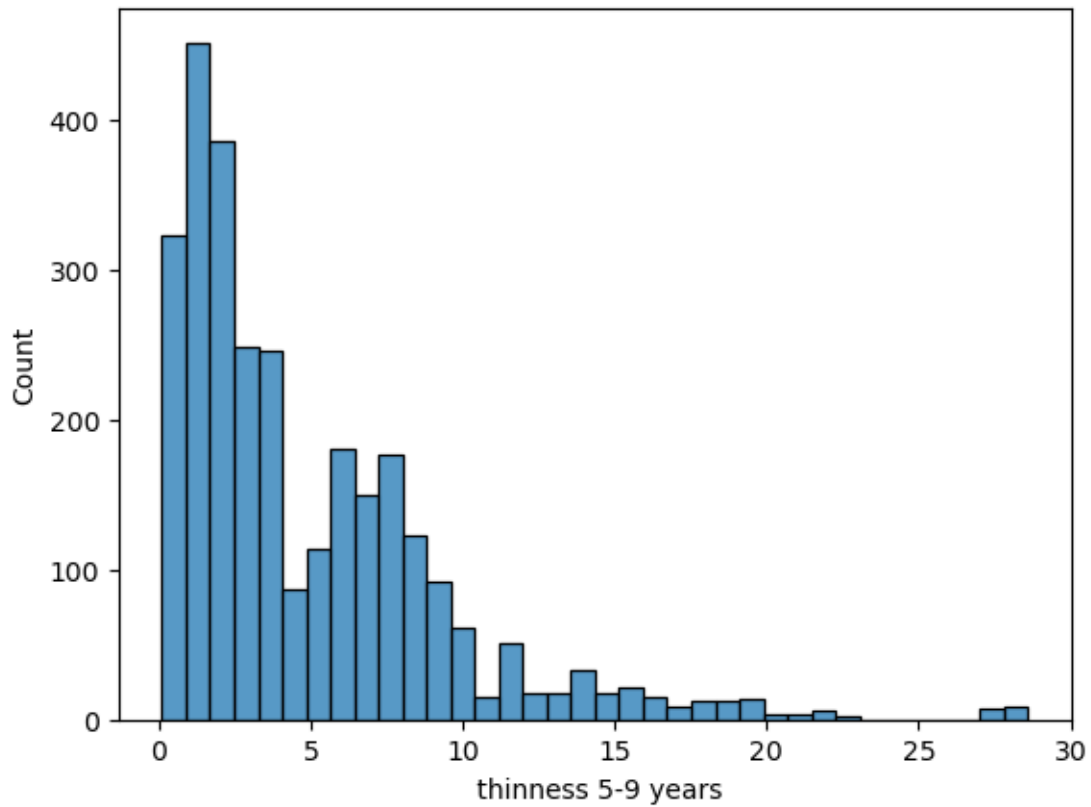
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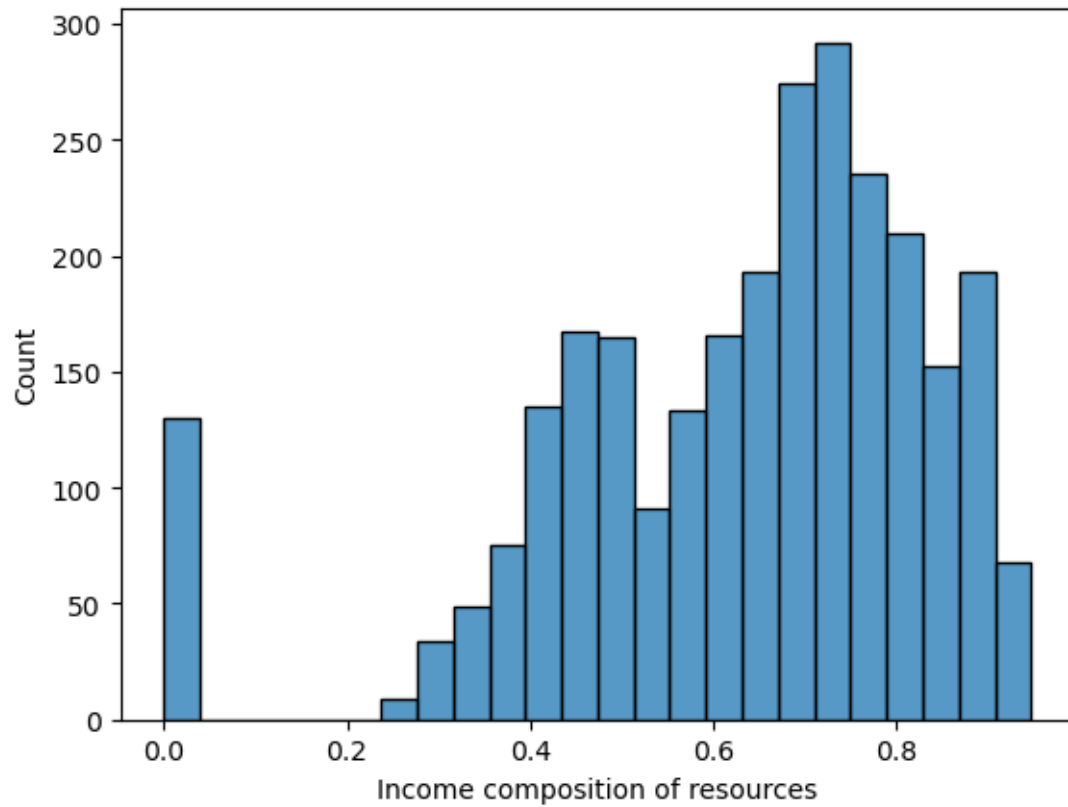
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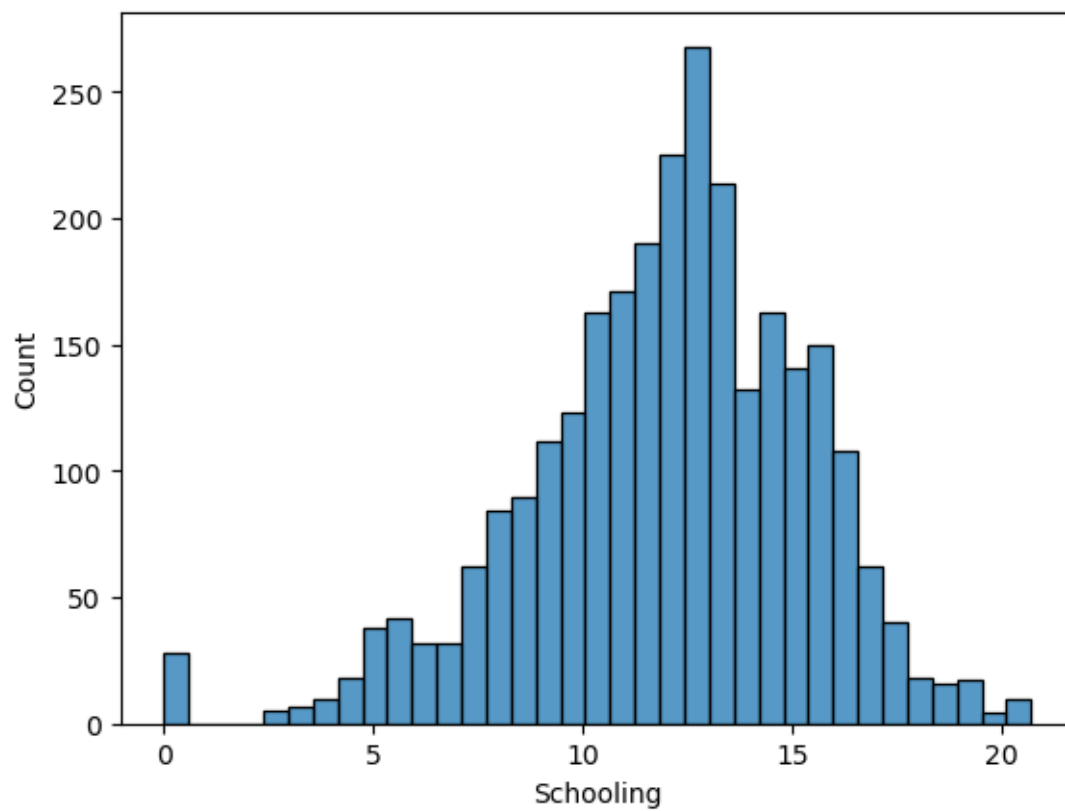
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FutureWarning: use_inf_as_na option is deprecated and will be removed in a
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```



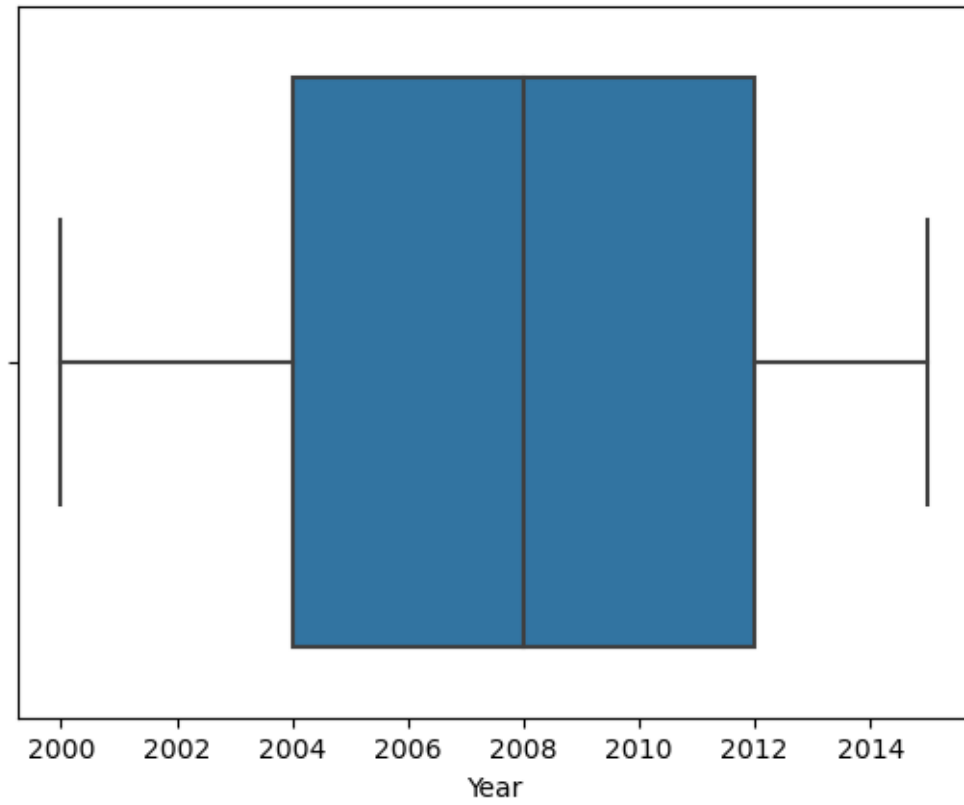
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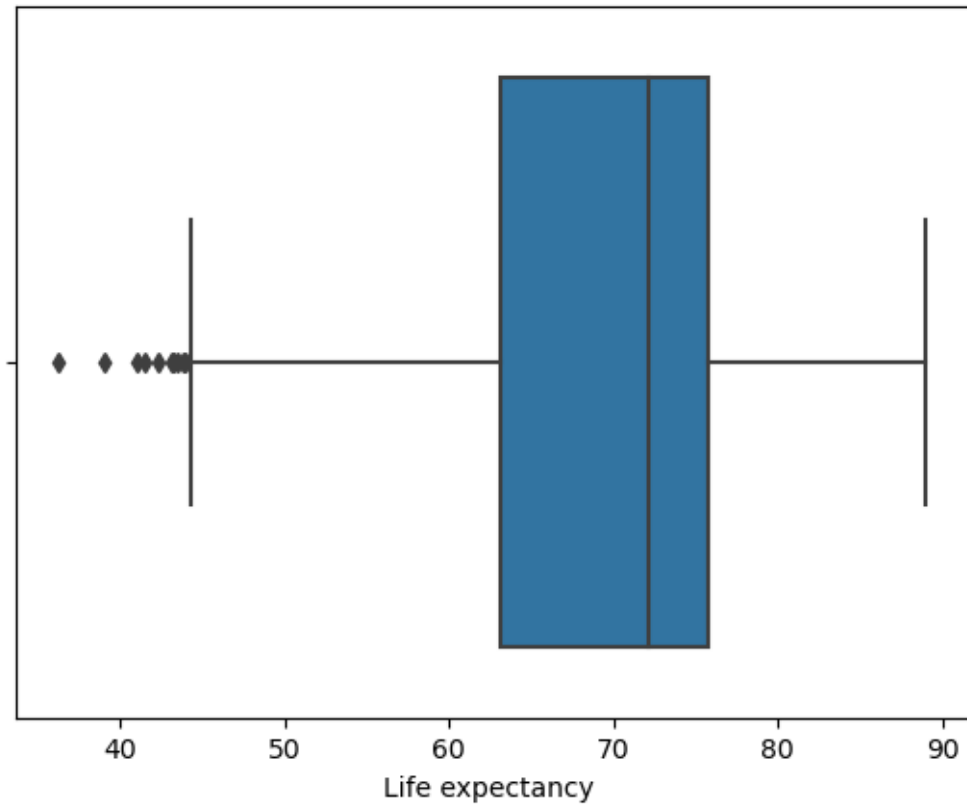


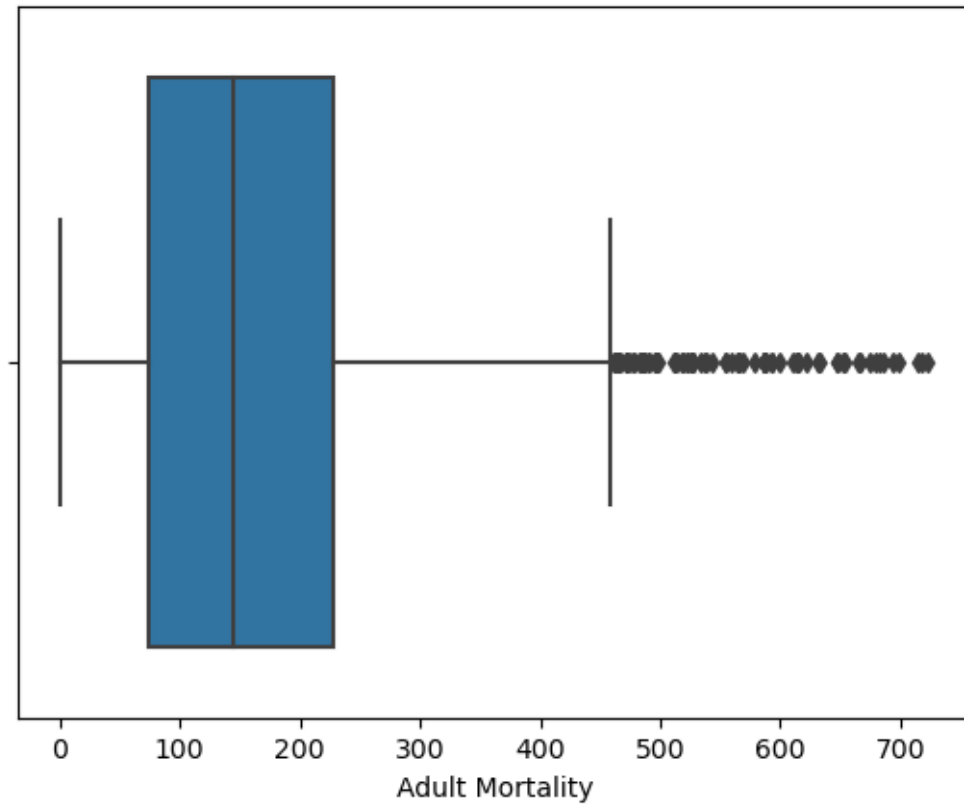
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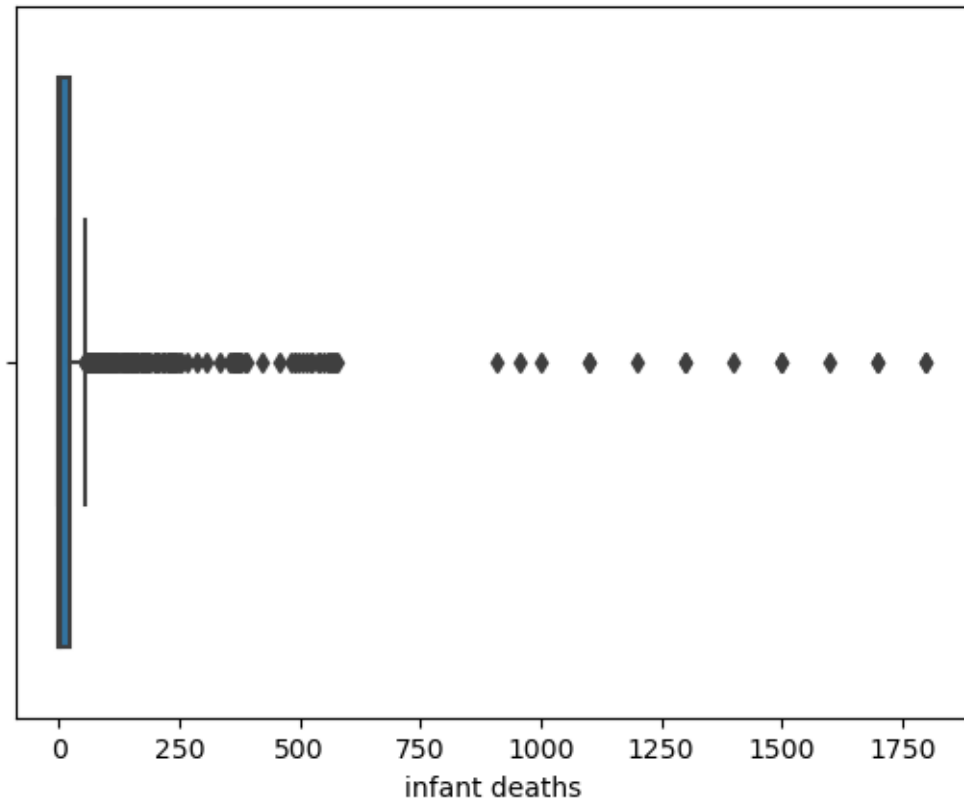



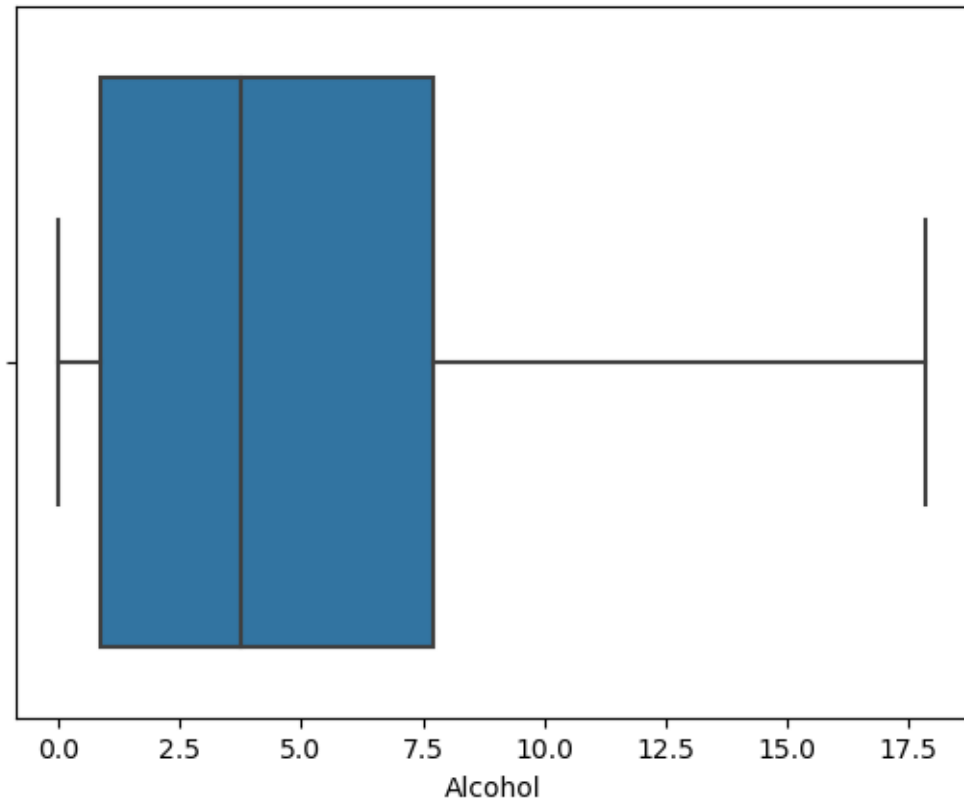
```
[43]: ## Boxplot to identify the outlier
for i in data.select_dtypes(include="number").columns:
    sns.boxplot(data=data, x=i)
    plt.show()
```

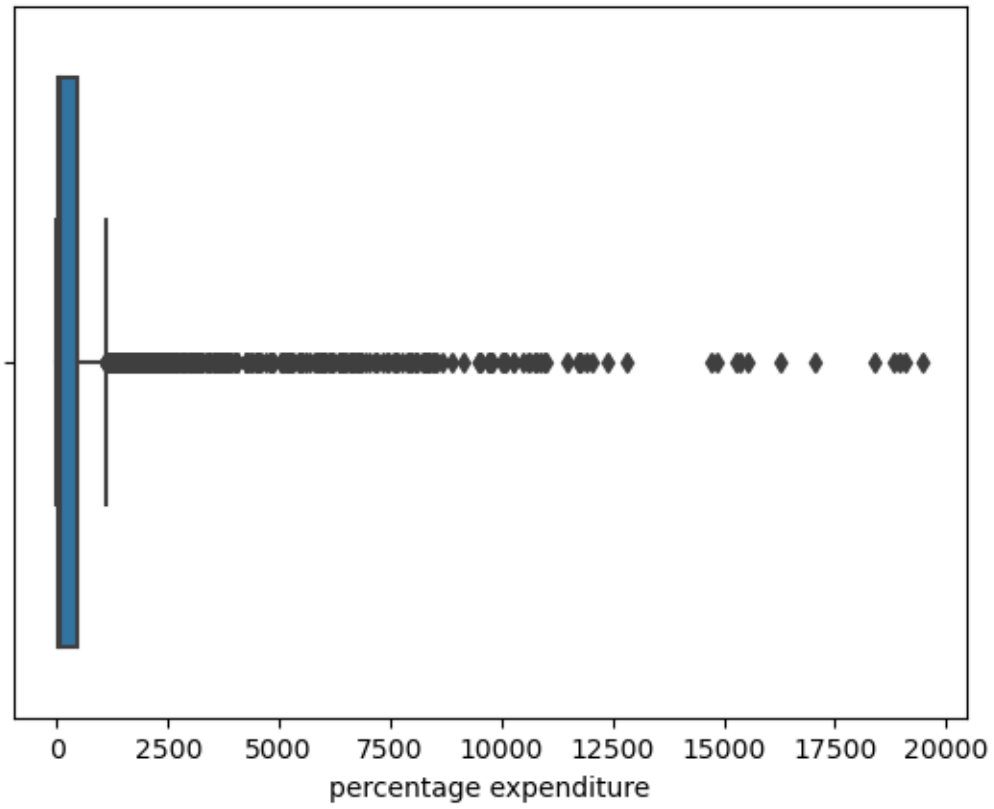


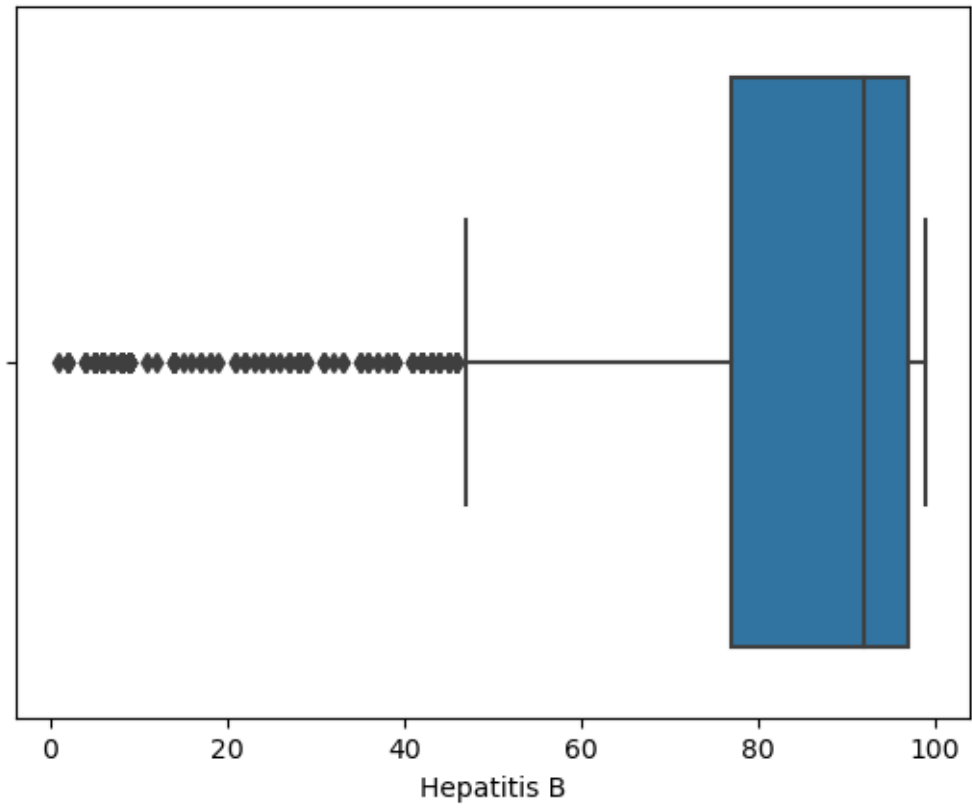


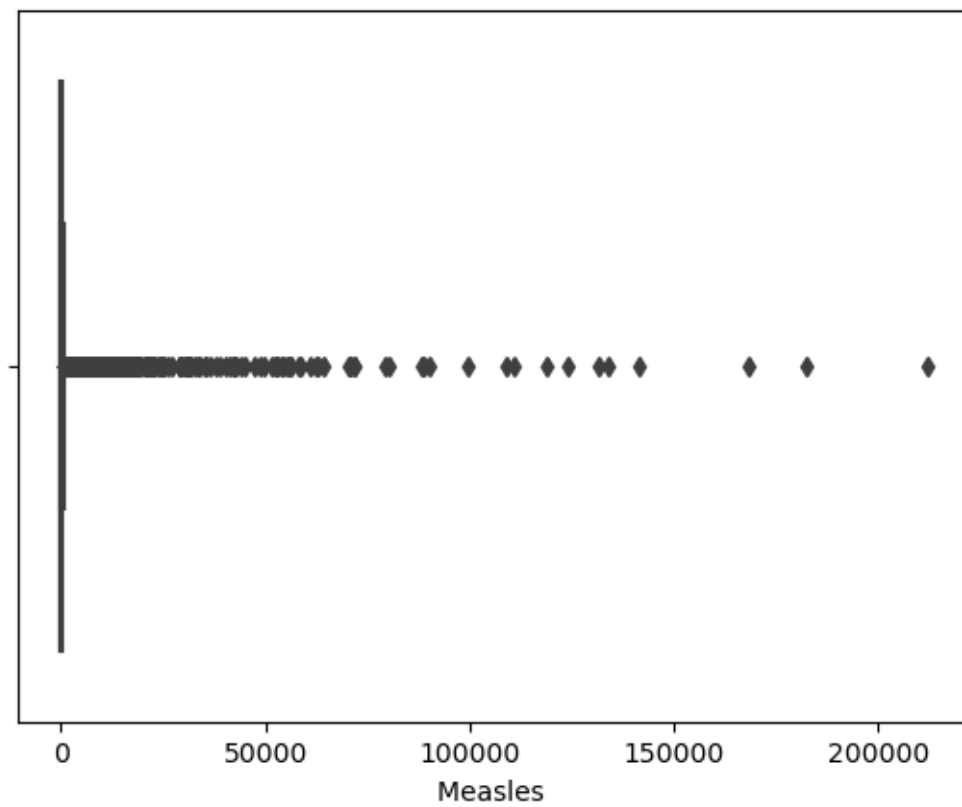


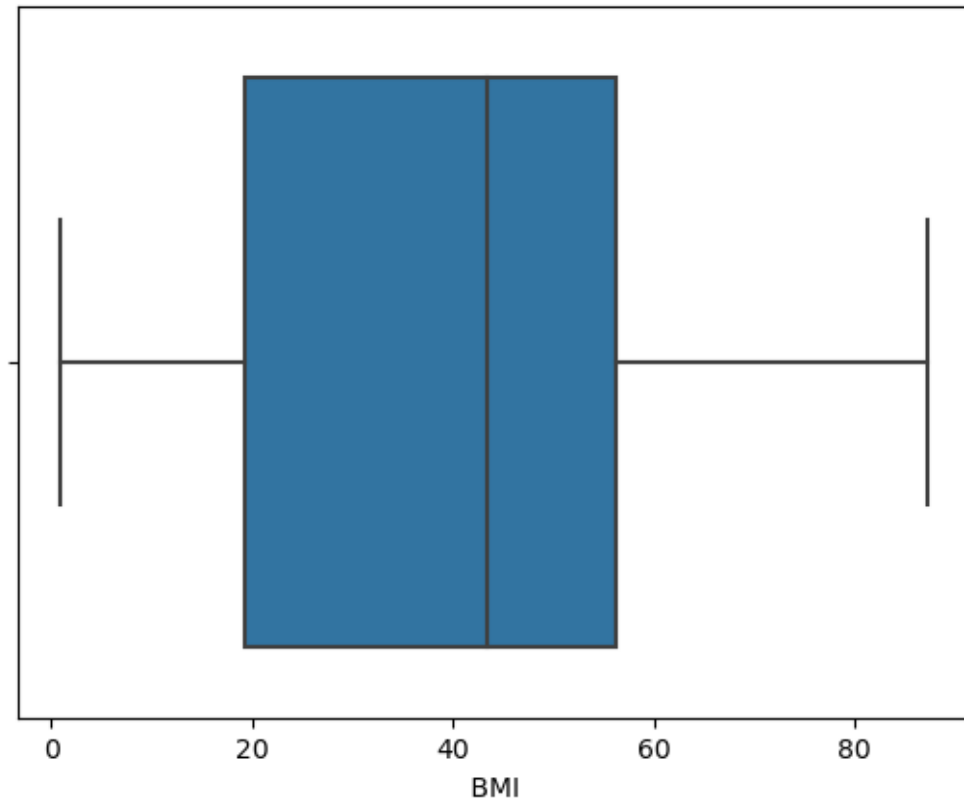


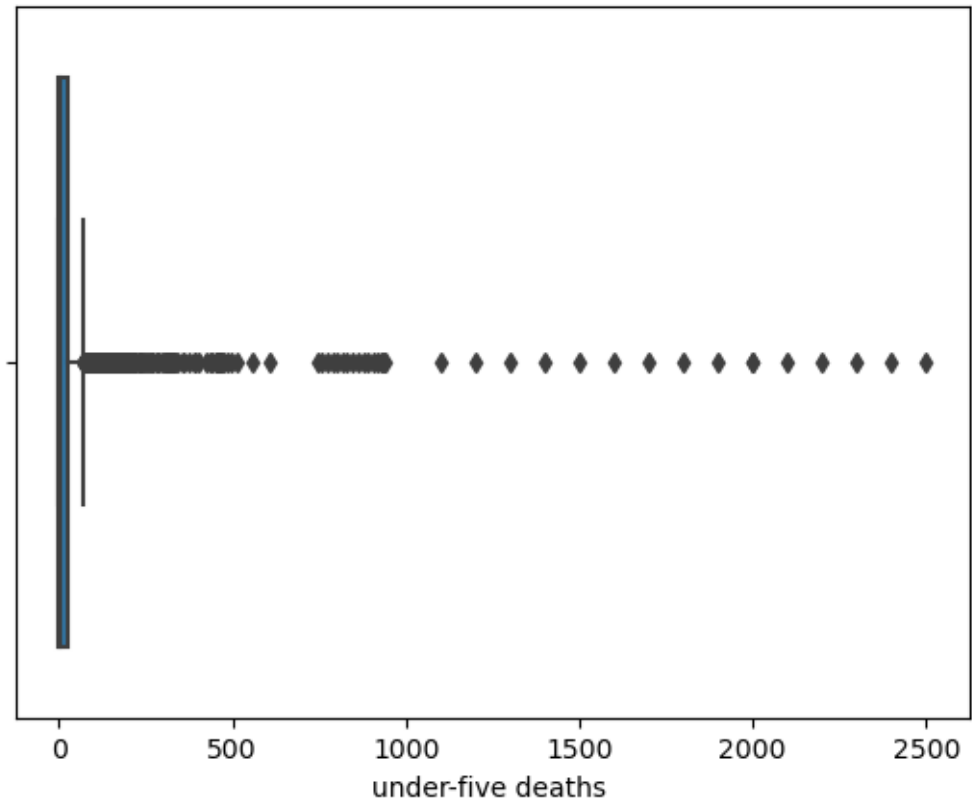


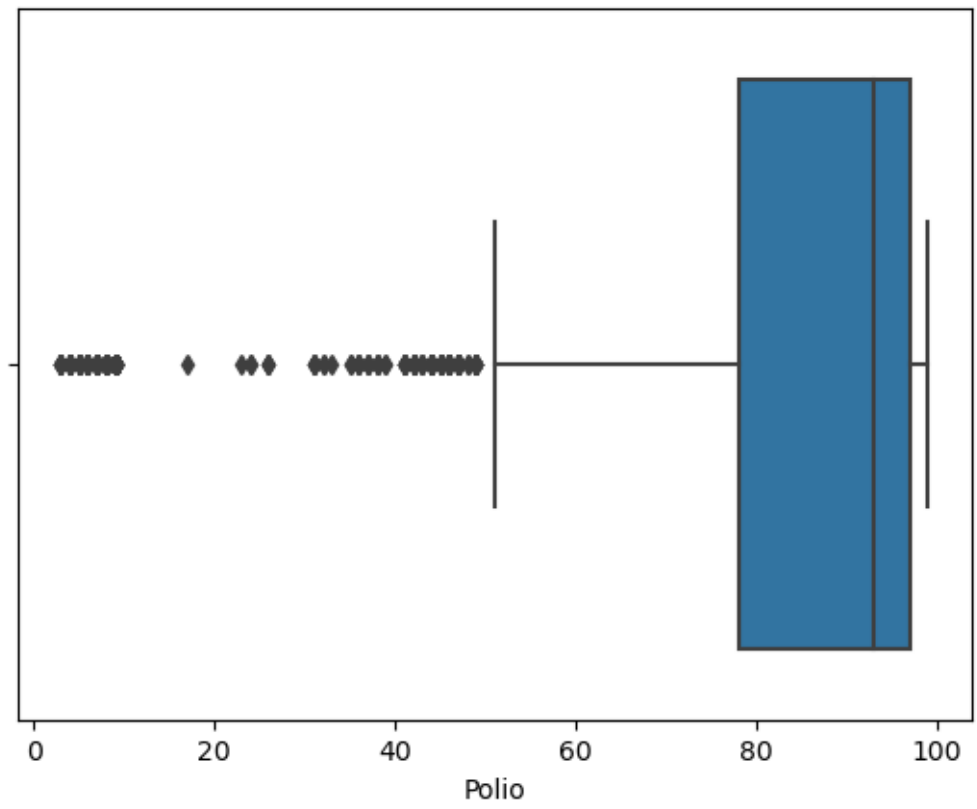


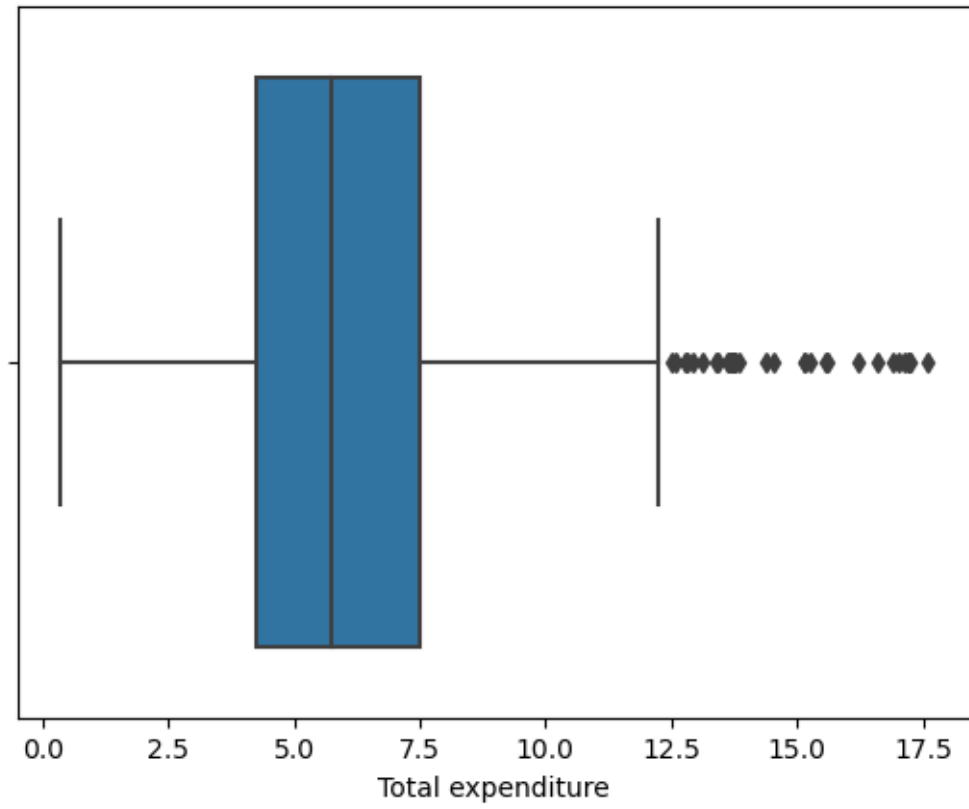


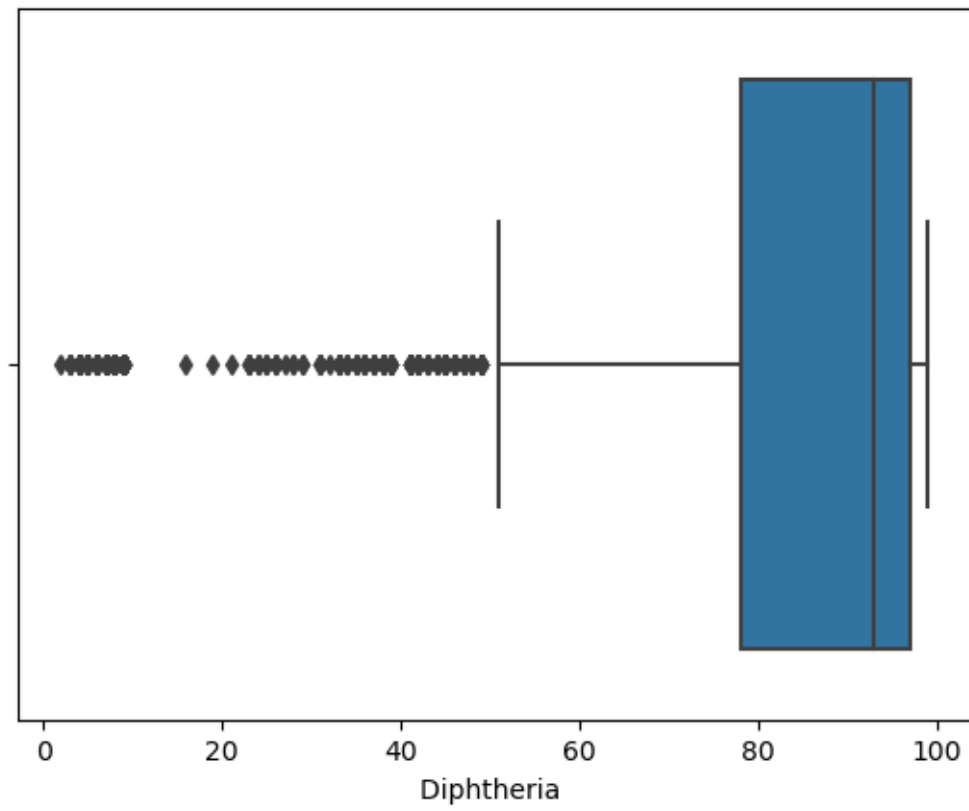


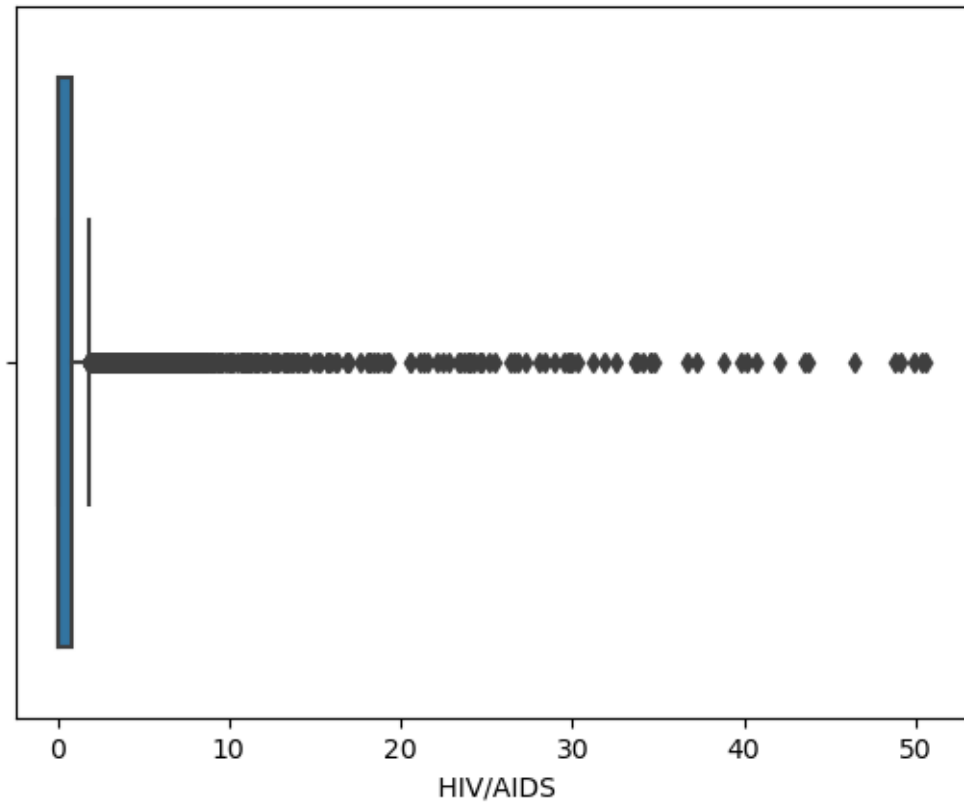


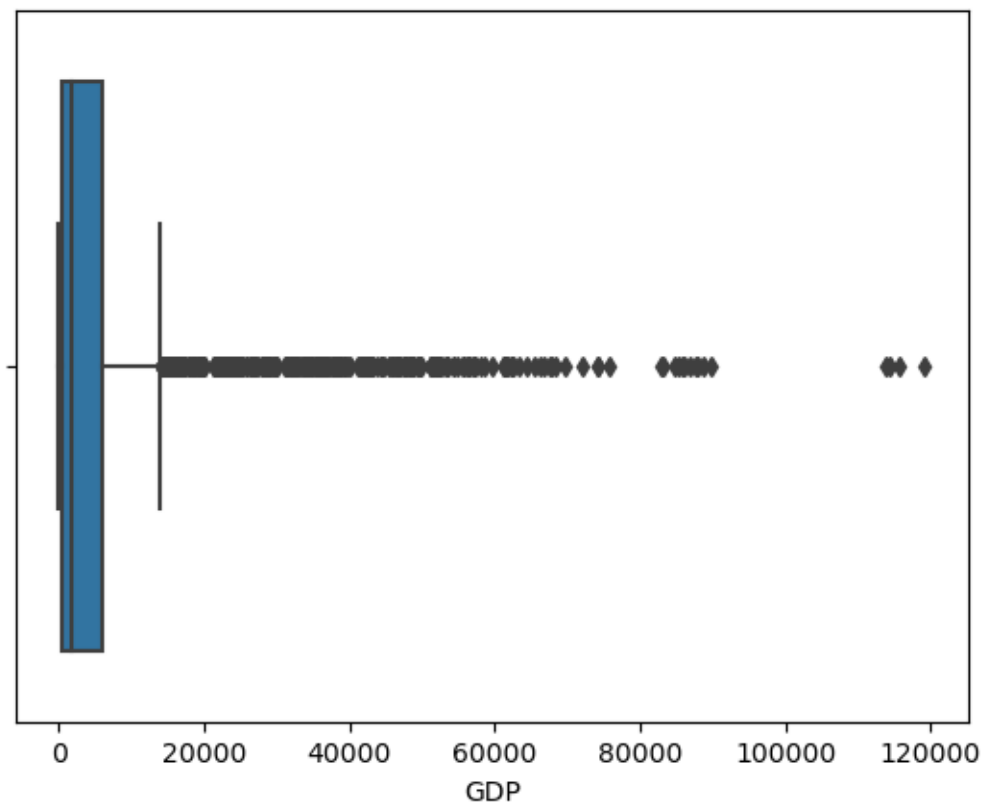


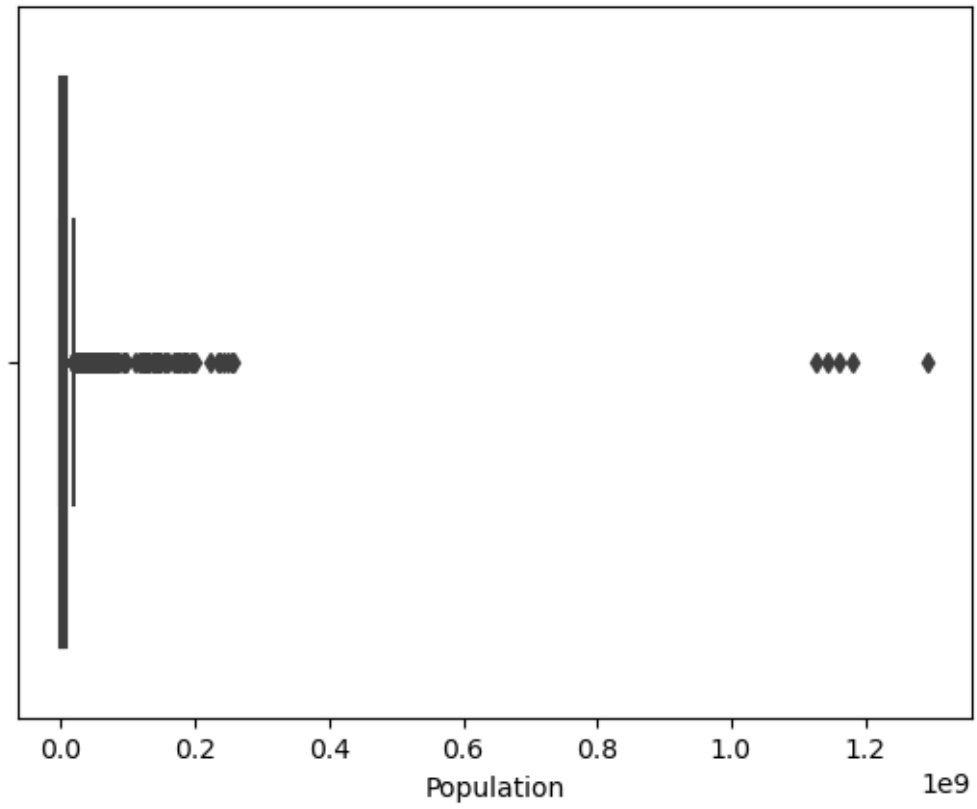


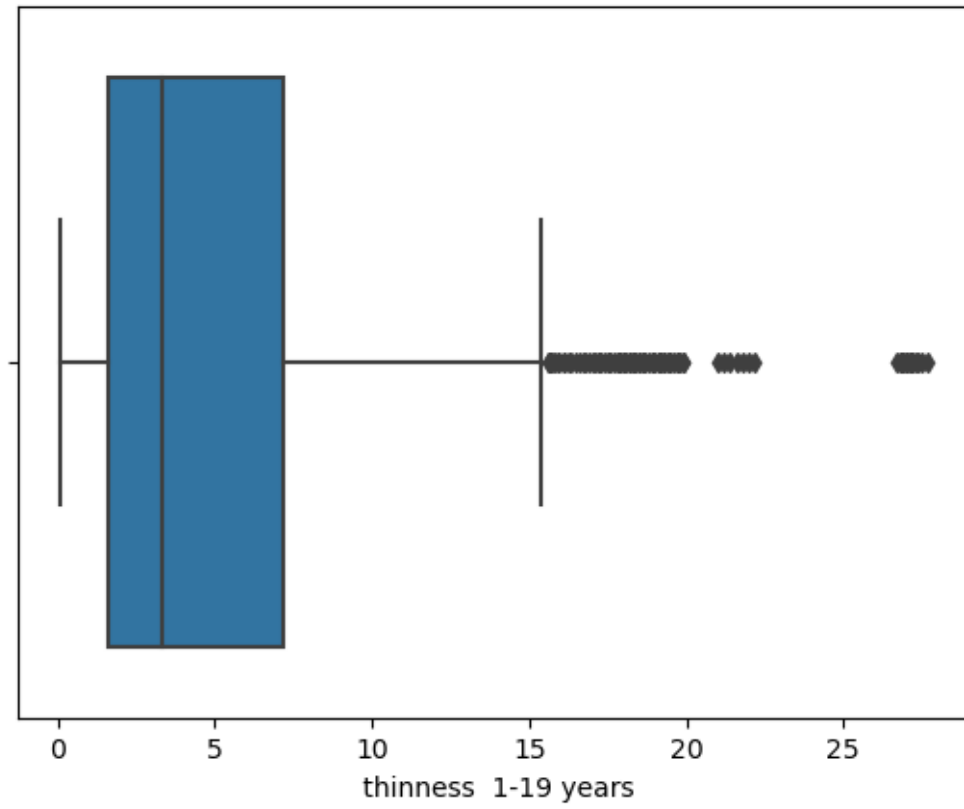


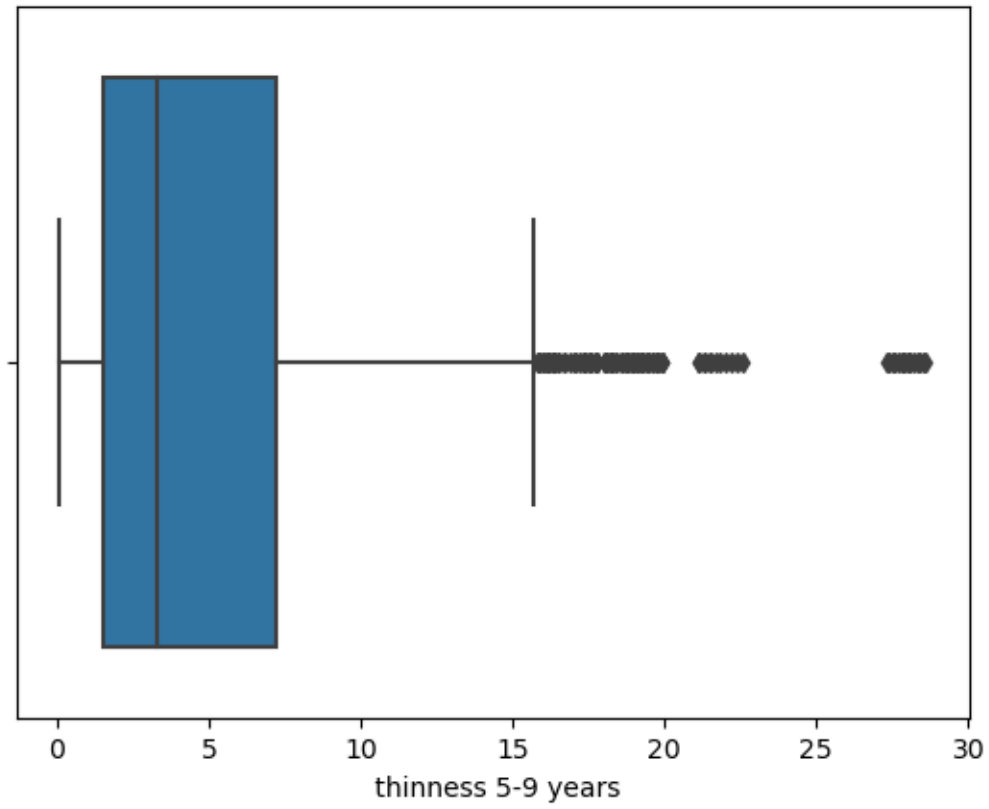


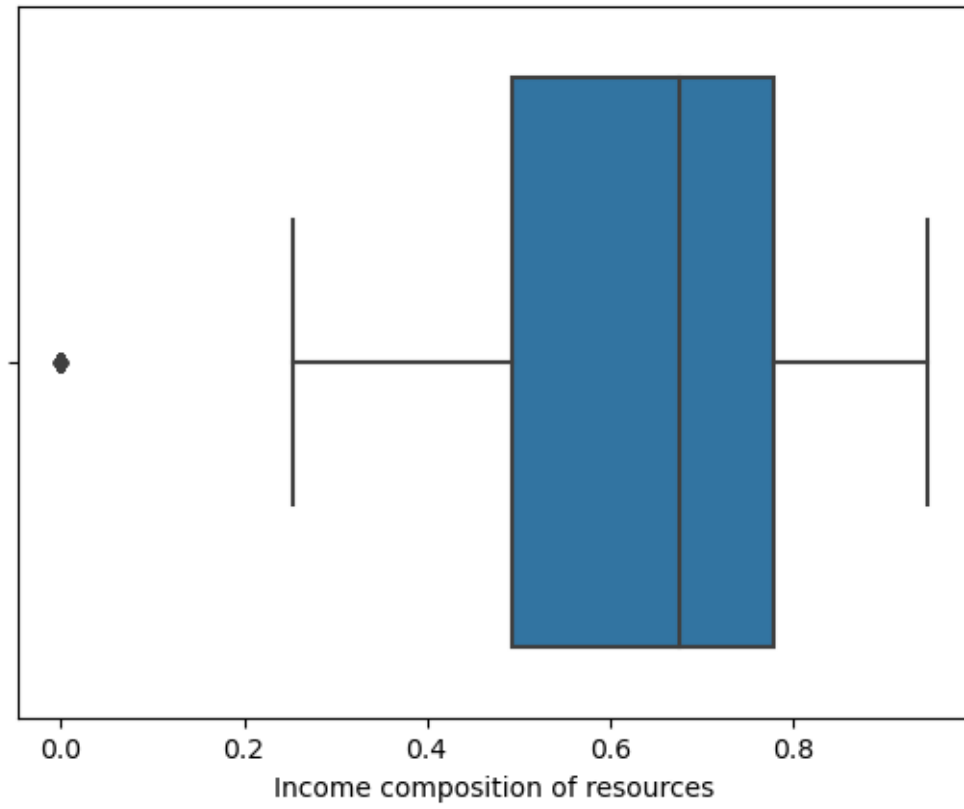


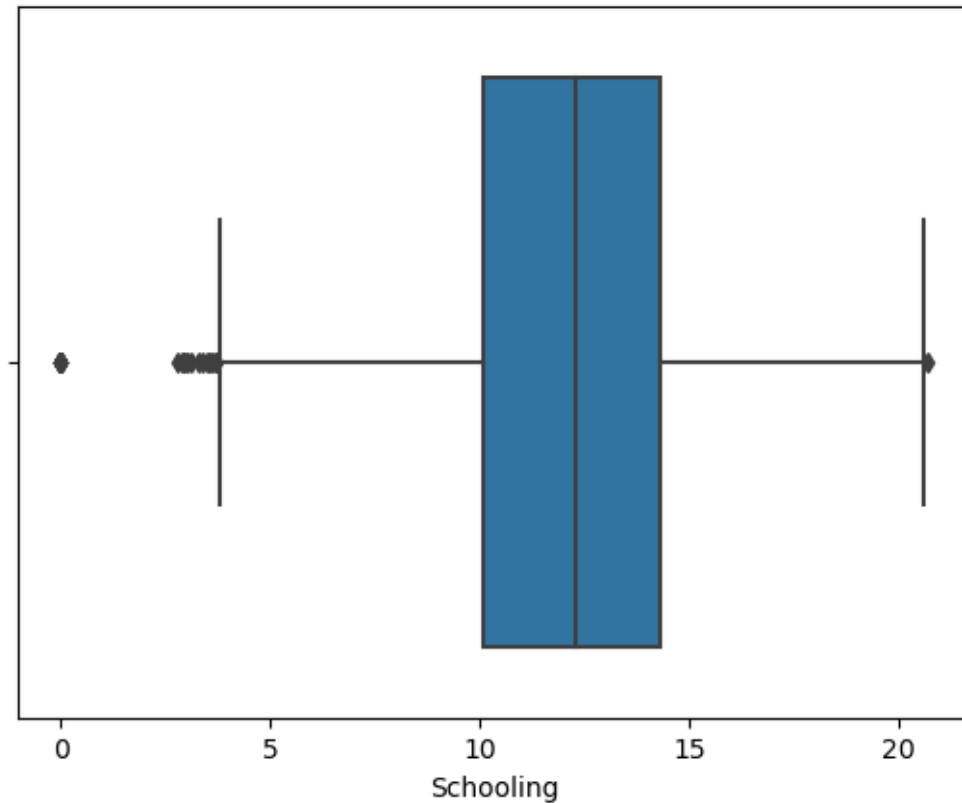












```
[47]: data.rename(columns={'Measles ': 'Measles'}, inplace=True)
```

```
[50]: print(data.columns)
data.rename(columns={' thinness 1-19 years': 'thinness_1_19_years'},
            inplace=True)
data.columns = data.columns.str.replace('\s+', ' ', regex=True).str.strip()
data.columns = data.columns.str.replace('\s+', ' ', regex=True).str.strip()
print(data.columns) # Check cleaned column names
```

```
Index(['Country', 'Year', 'Status', 'Life expectancy', 'Adult Mortality',
       'infant deaths', 'Alcohol', 'percentage expenditure', 'Hepatitis B',
       'Measles', 'BMI', 'under-five deaths', 'Polio', 'Total expenditure',
       'Diphtheria', 'HIV/AIDS', 'GDP', 'Population', 'thinness 1-19 years',
       'thinness 5-9 years', 'Income composition of resources', 'Schooling'],
      dtype='object')
```

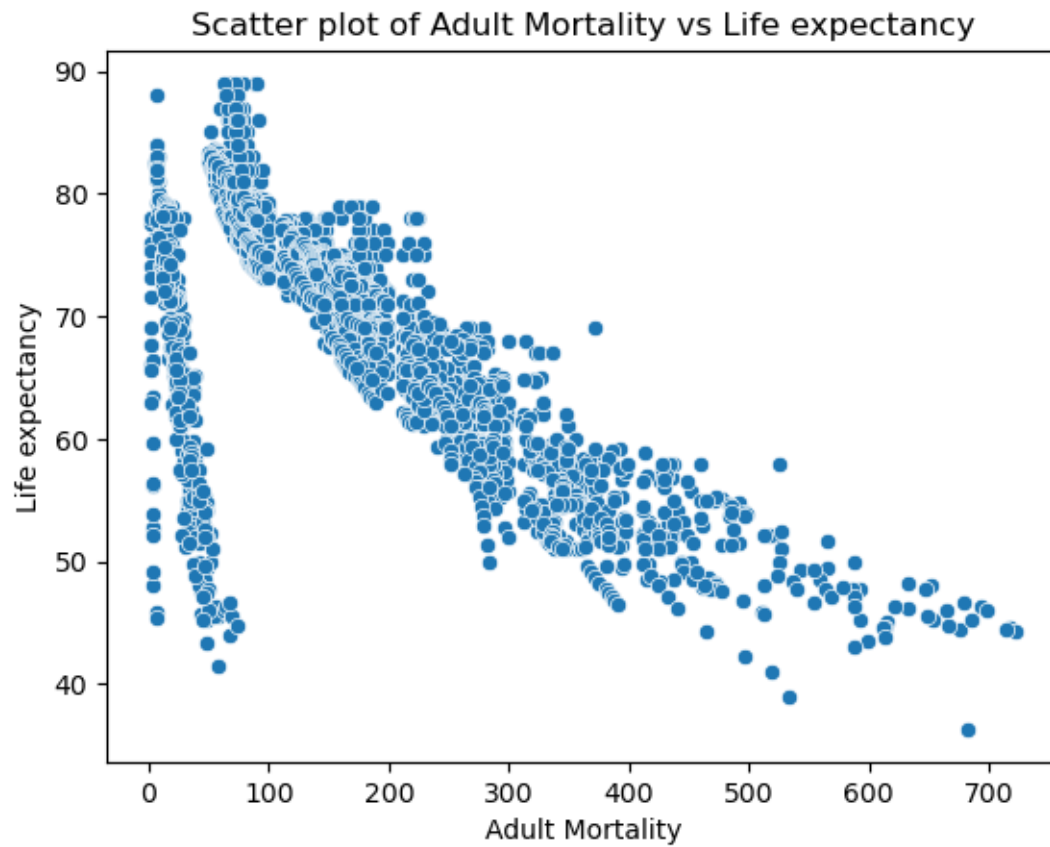
```
Index(['Country', 'Year', 'Status', 'Life expectancy', 'Adult Mortality',
       'infant deaths', 'Alcohol', 'percentage expenditure', 'Hepatitis B',
       'Measles', 'BMI', 'under-five deaths', 'Polio', 'Total expenditure',
       'Diphtheria', 'HIV/AIDS', 'GDP', 'Population', 'thinness 1-19 years',
       'thinness 5-9 years', 'Income composition of resources', 'Schooling'],
      dtype='object')
```

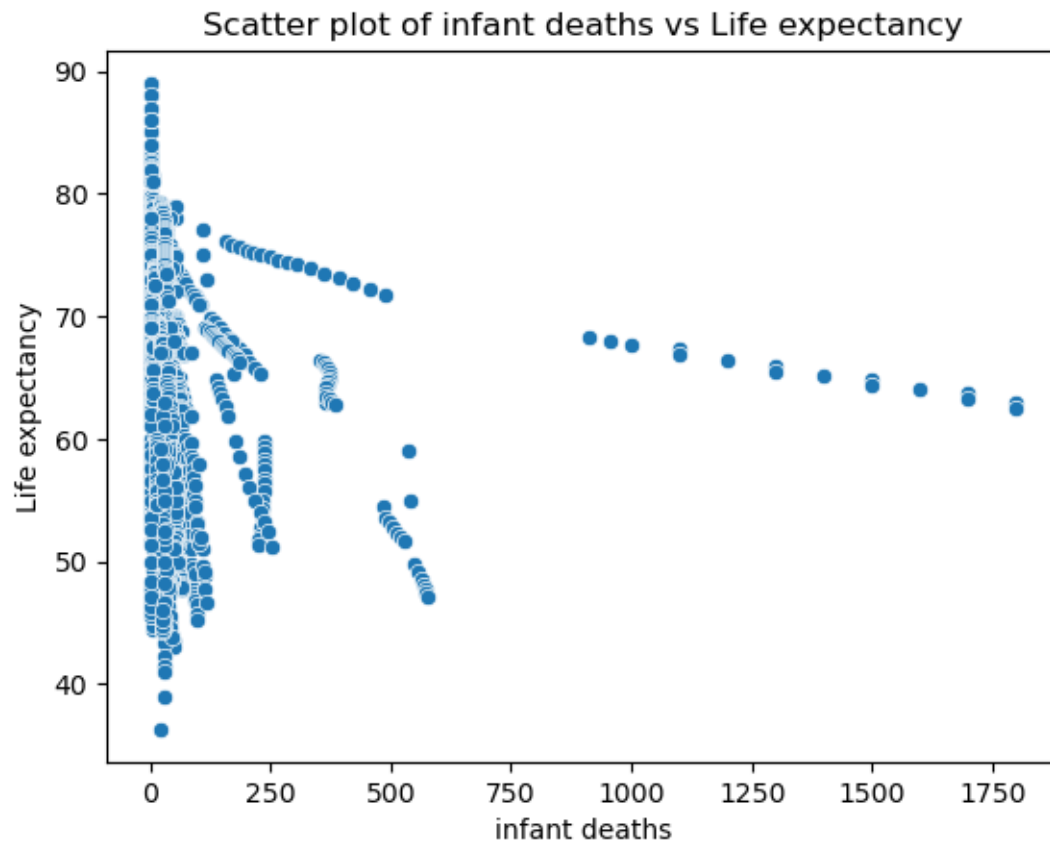
```
[51]: ## scatterplot to understand the relationship
# Strip any leading/trailing whitespace from column names
data.columns = data.columns.str.strip()

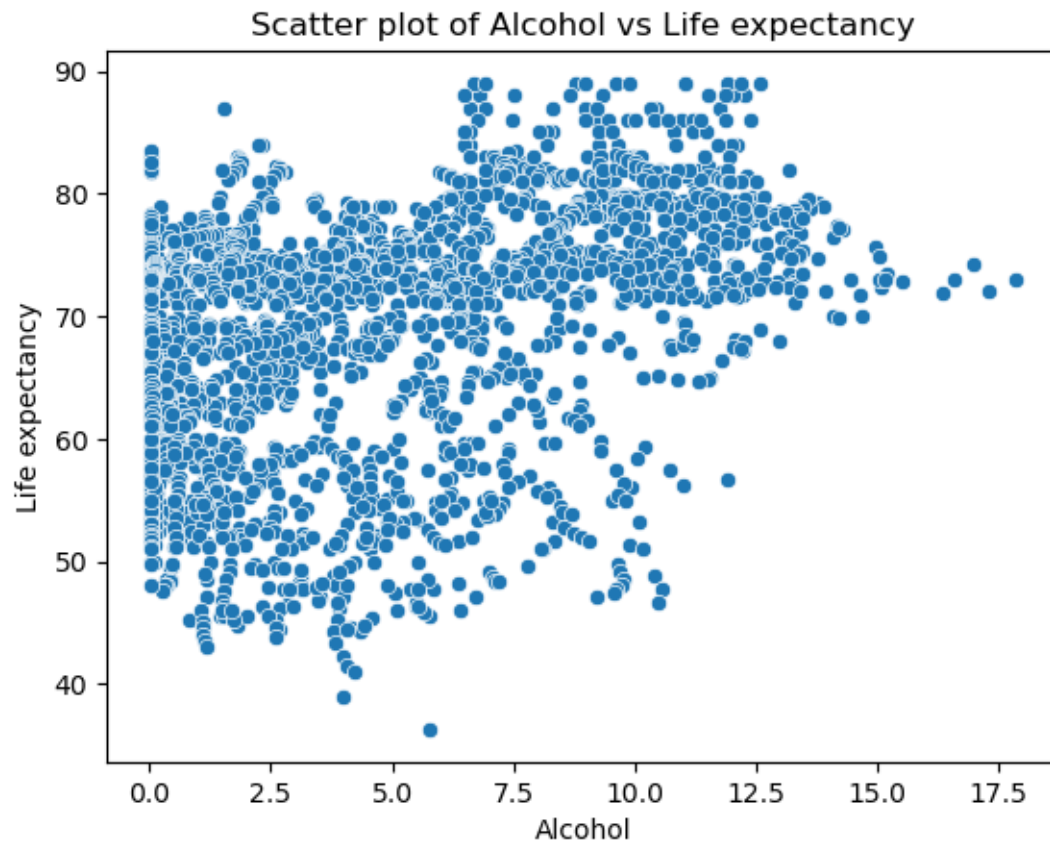
# List of columns to create scatter plots for
columns = ['Year', 'Adult Mortality', 'infant deaths', 'Alcohol', 'percentage_
↳expenditure',
           'Hepatitis B', 'Measles', 'BMI', 'under-five deaths', 'Polio',
           'Total expenditure', 'Diphtheria', 'HIV/AIDS', 'GDP', 'Population',
           'thinness 1-19 years', 'thinness 5-9 years', 'Income composition of_
↳resources',
           'Schooling']

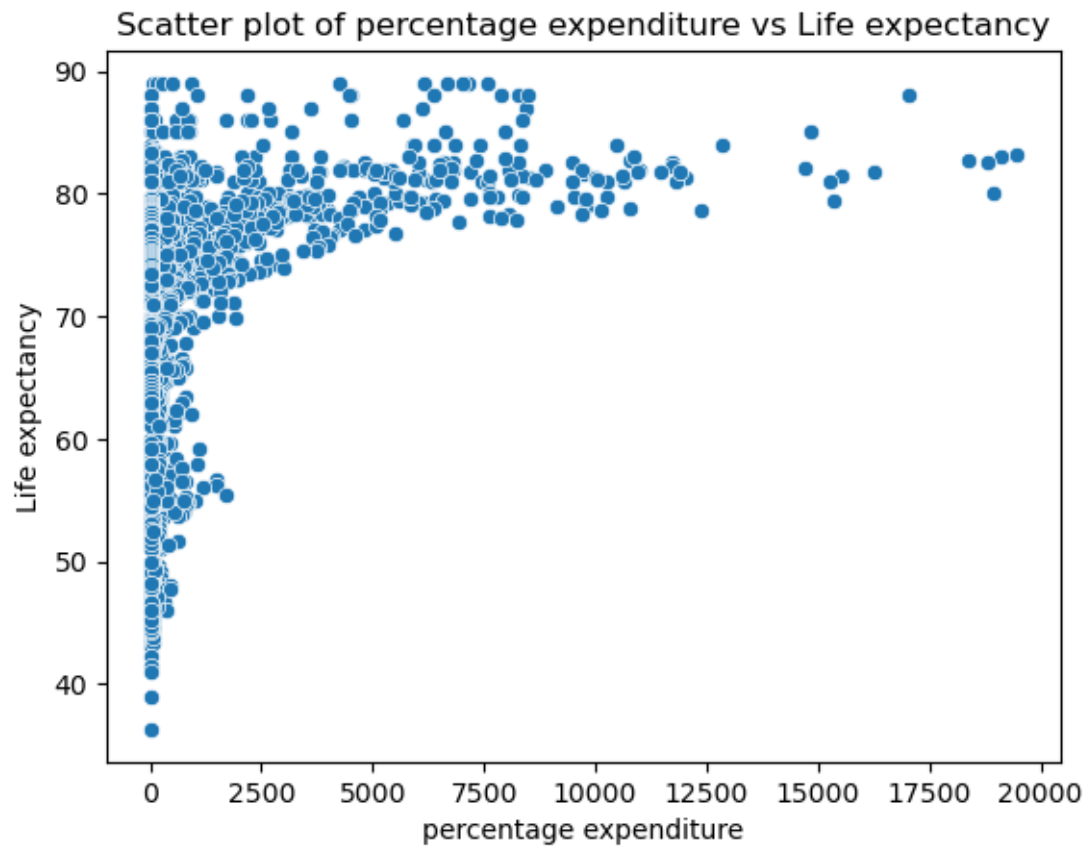
# Loop through each column and create a scatter plot against 'Life expectancy'
for i in columns:
    sns.scatterplot(data=data, x=i, y='Life expectancy') # Make sure 'Life_
↳expectancy' is correct
    plt.title(f'Scatter plot of {i} vs Life expectancy')
    plt.show()
```

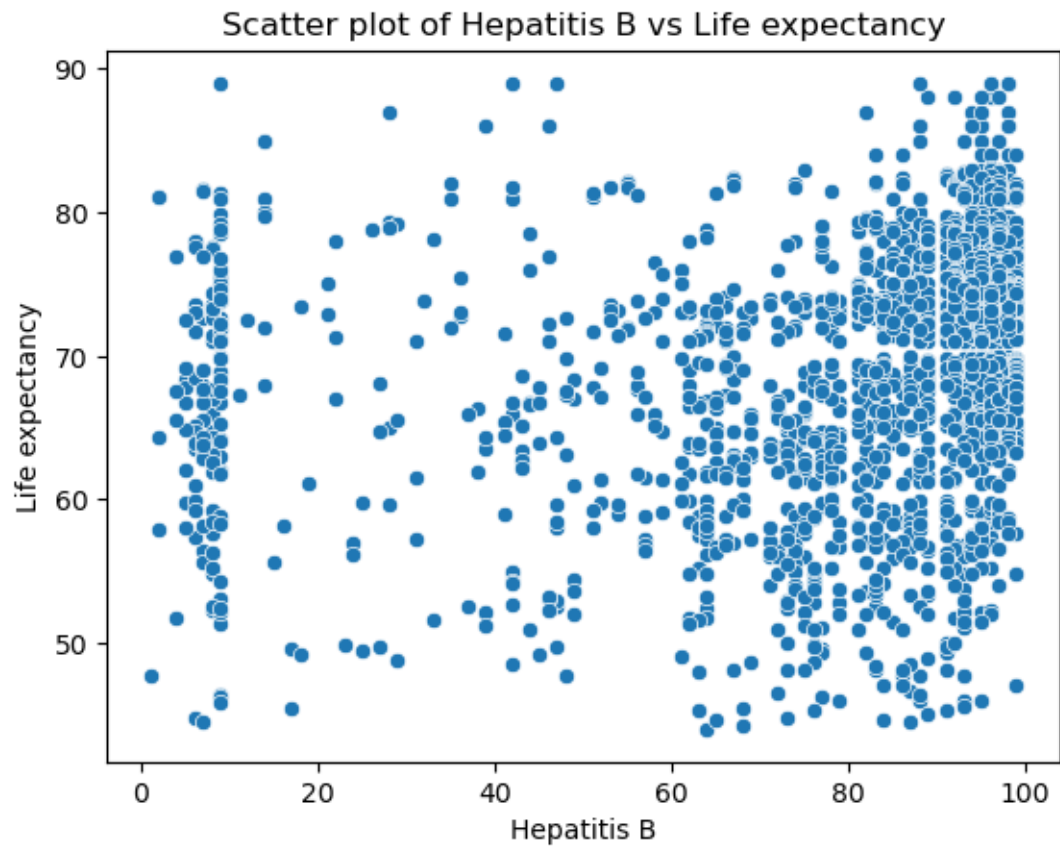




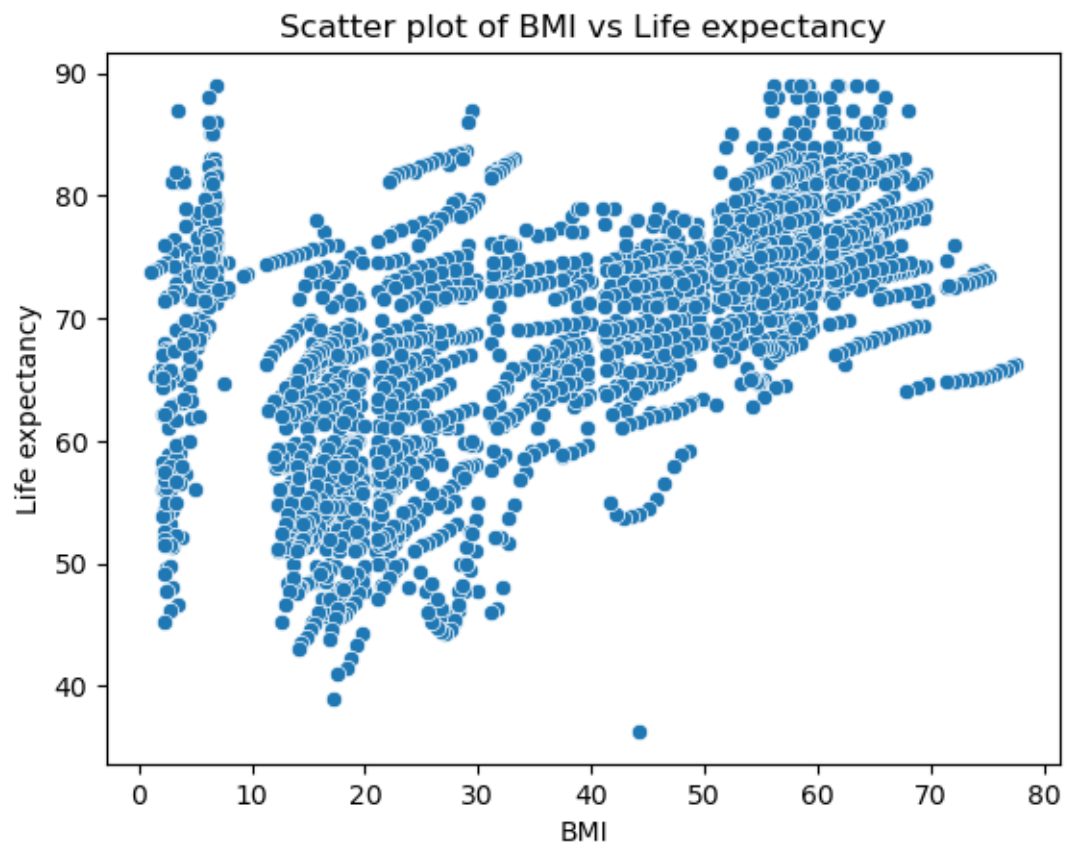




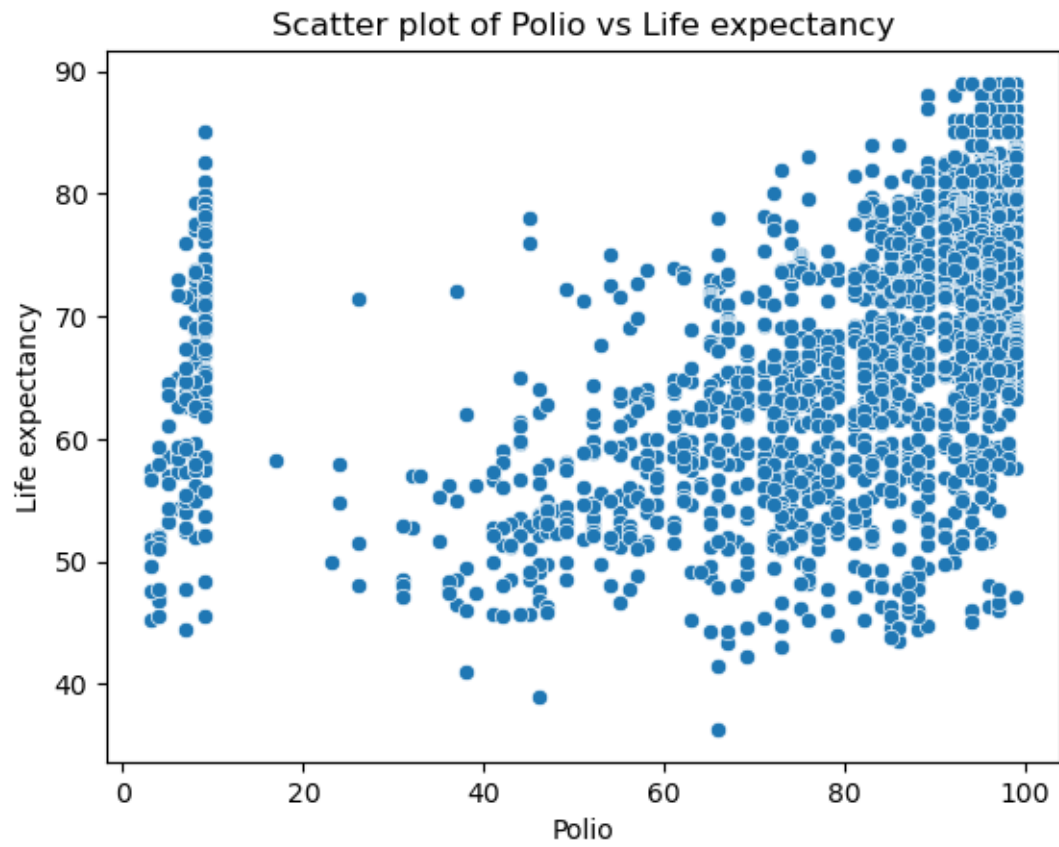


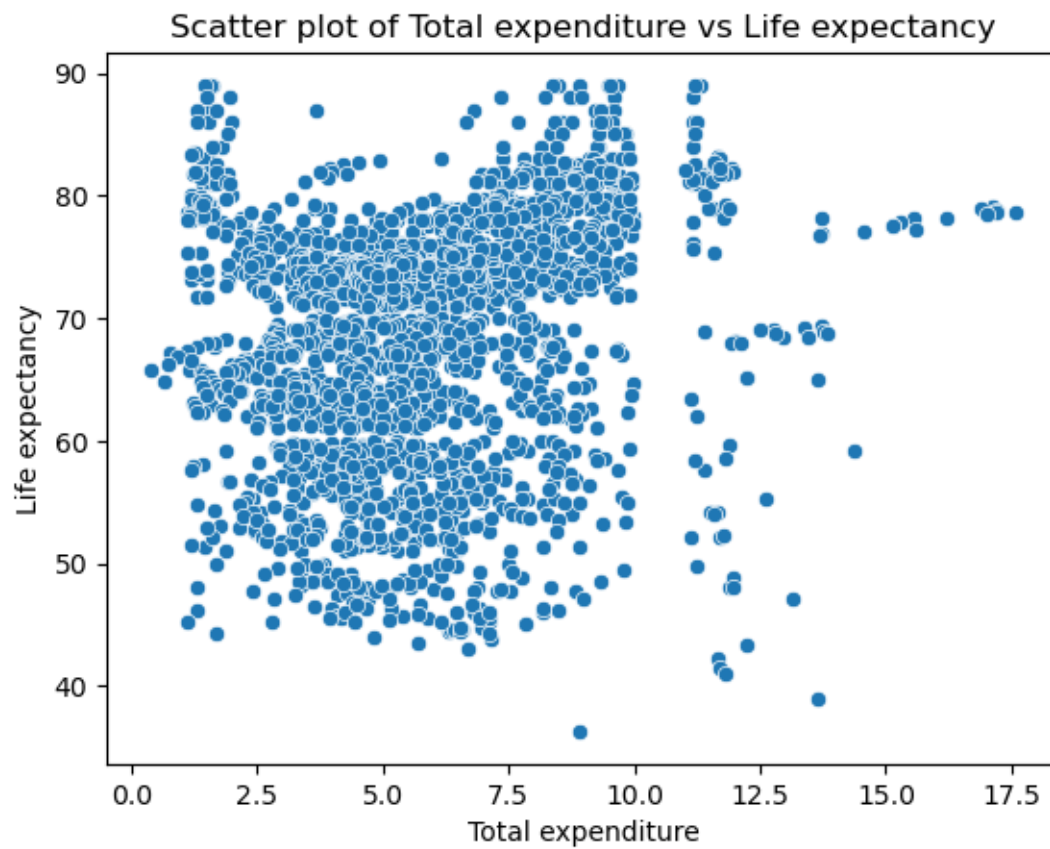


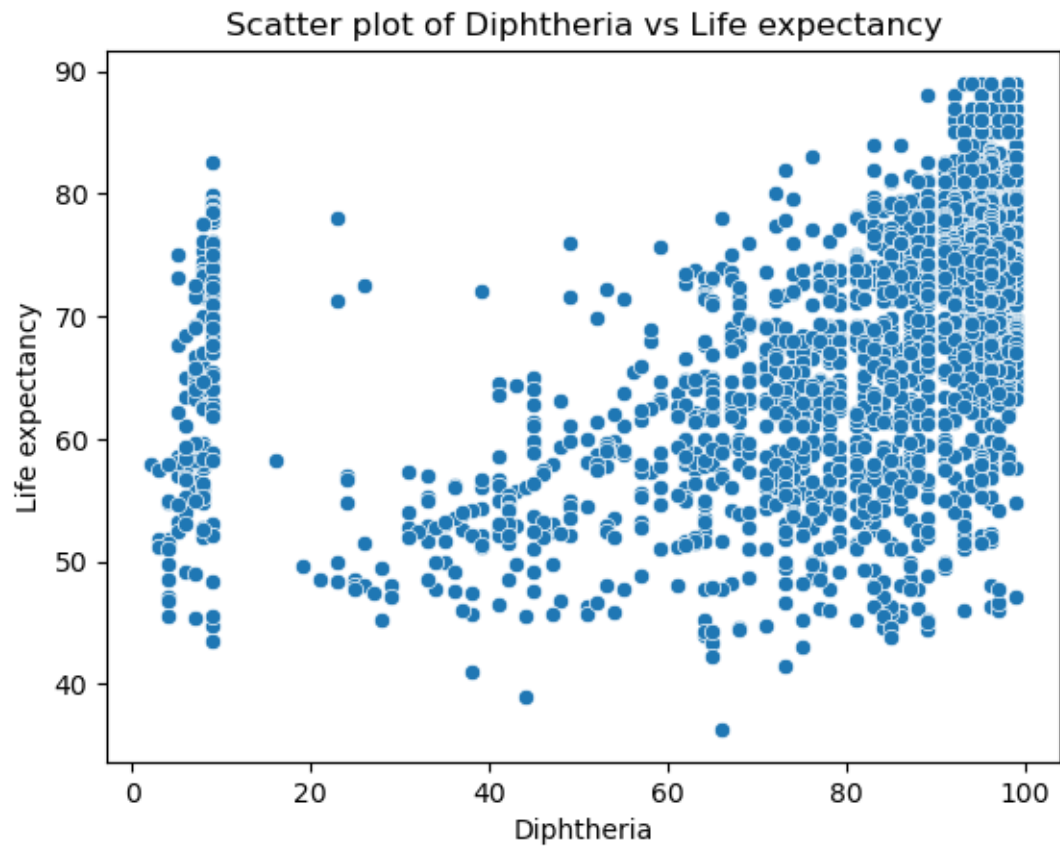


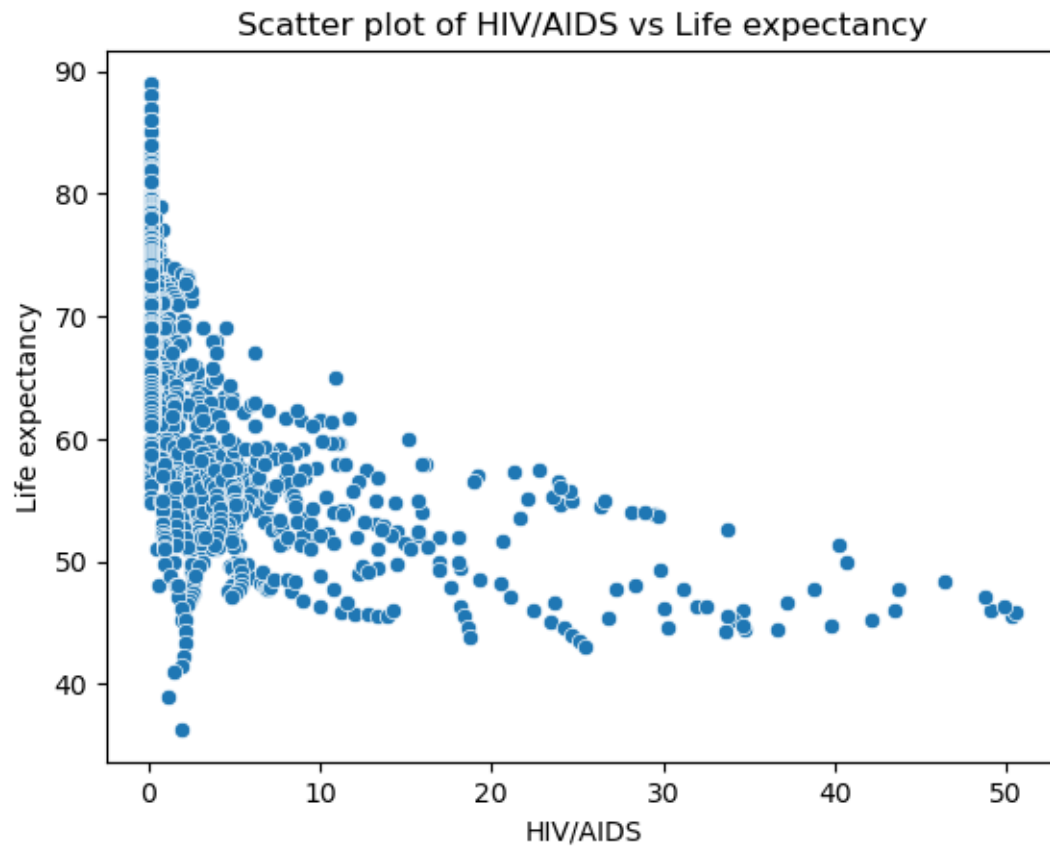


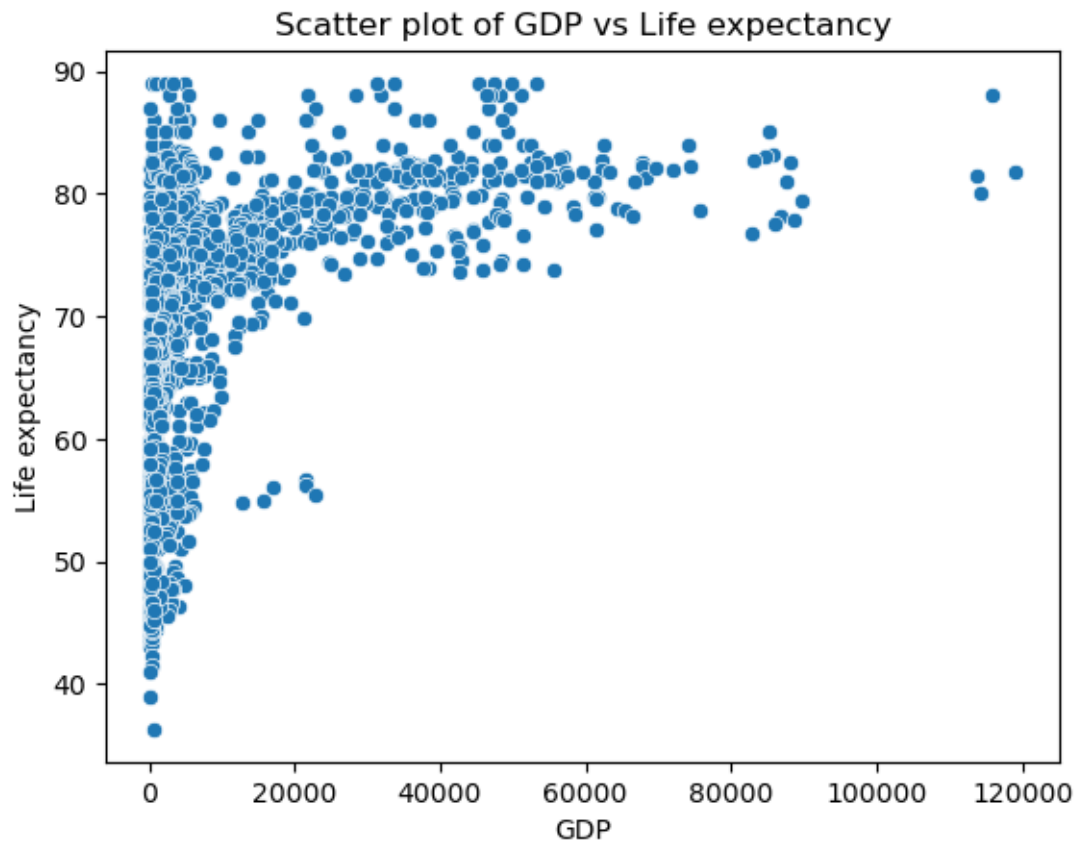


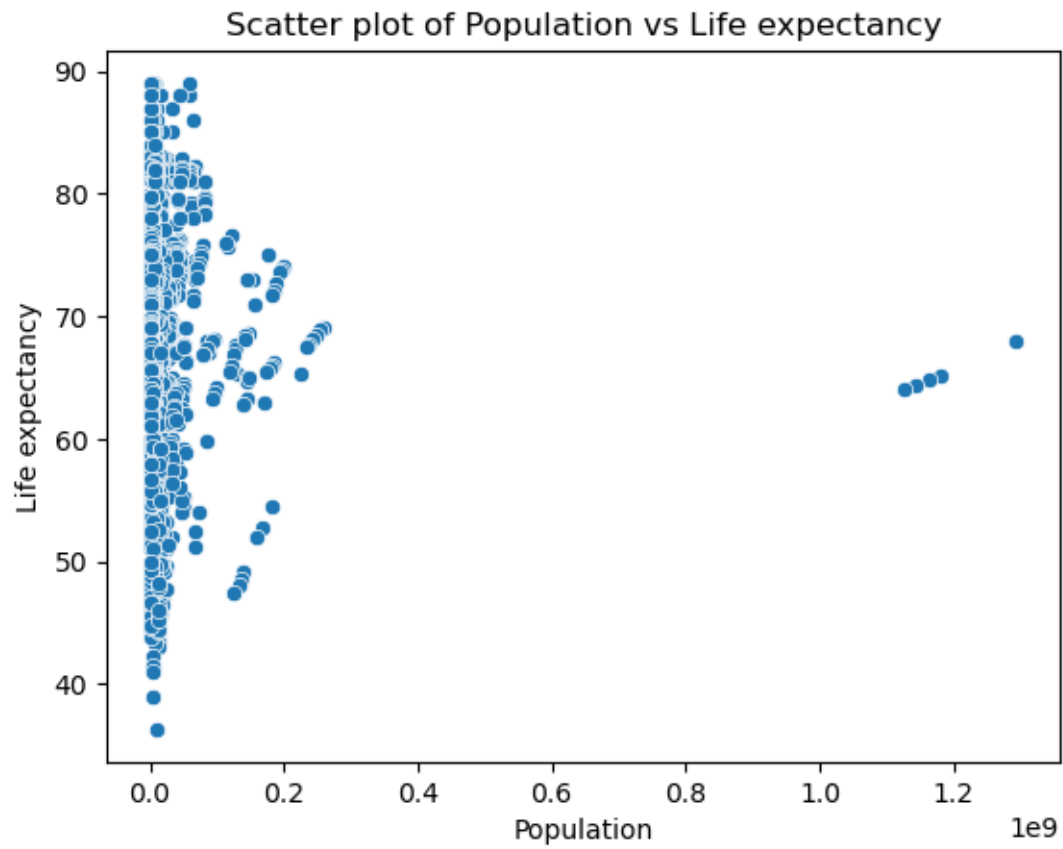


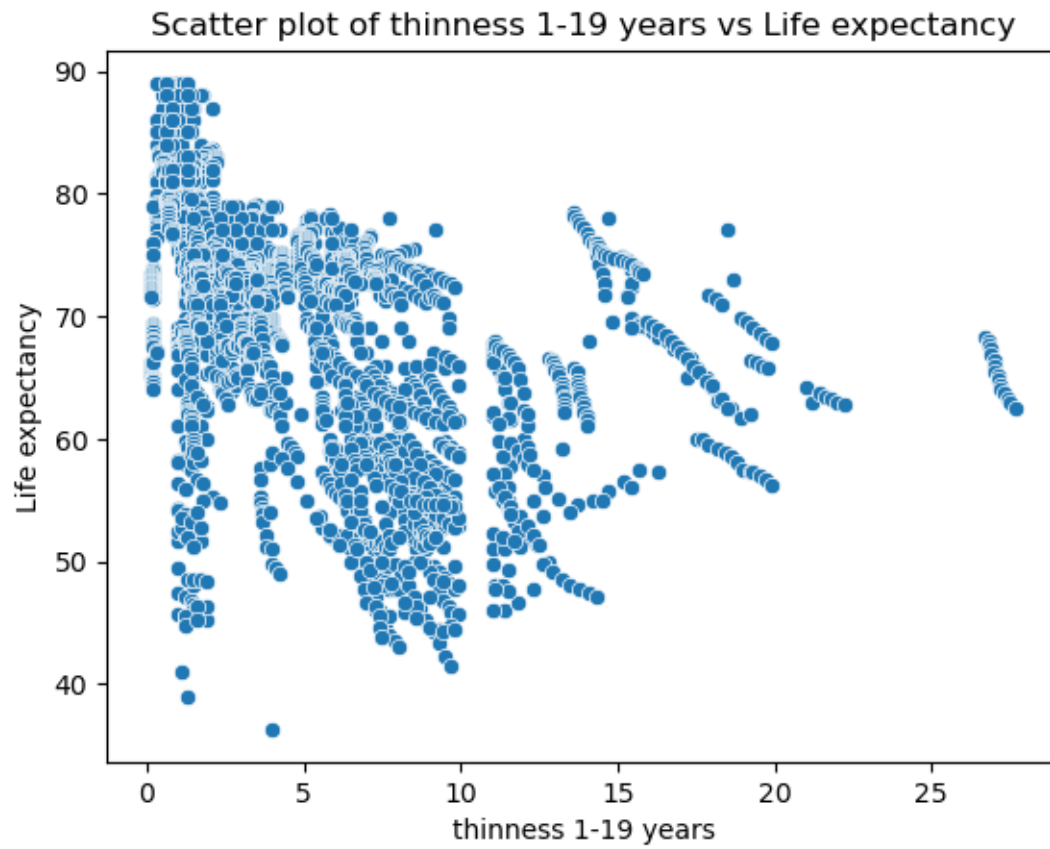


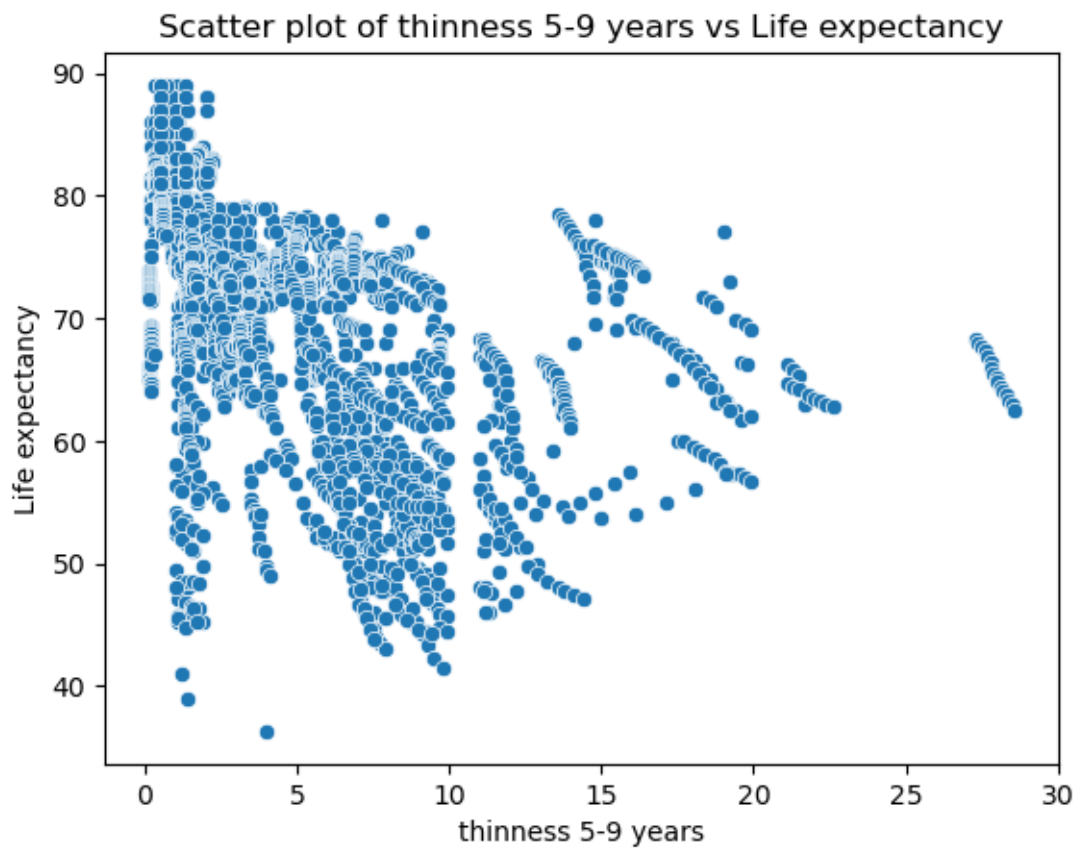


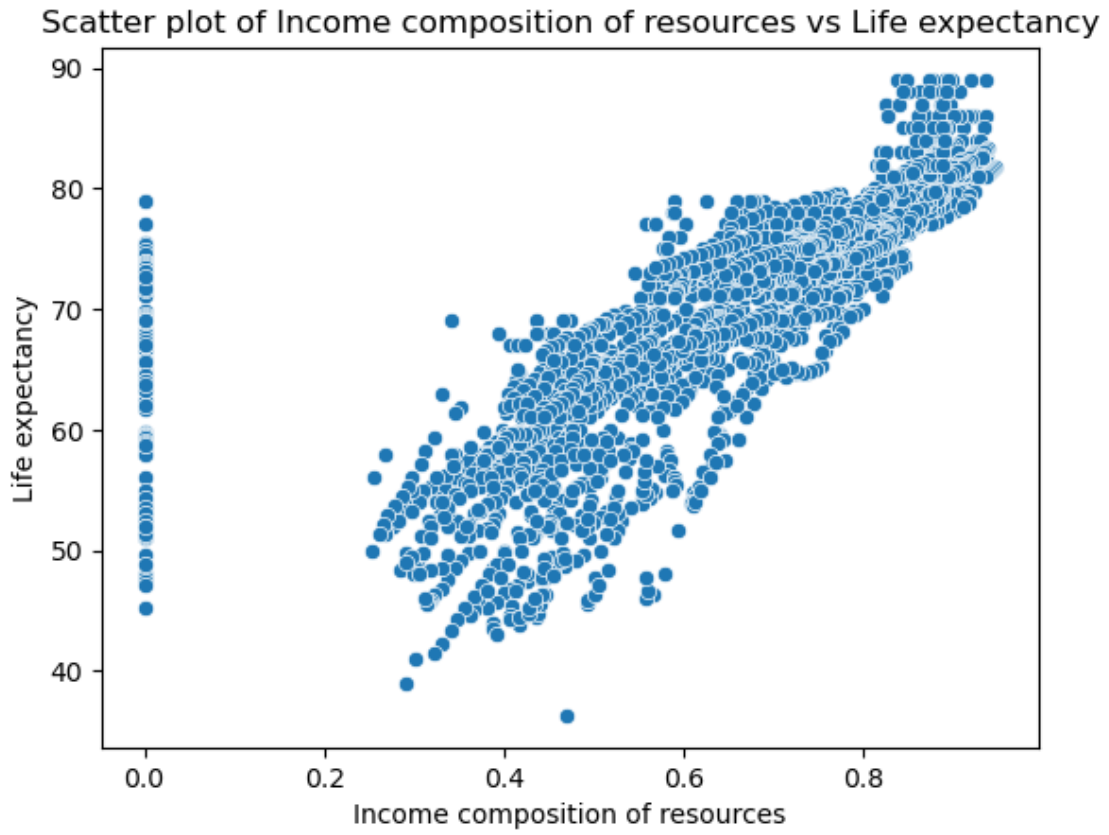


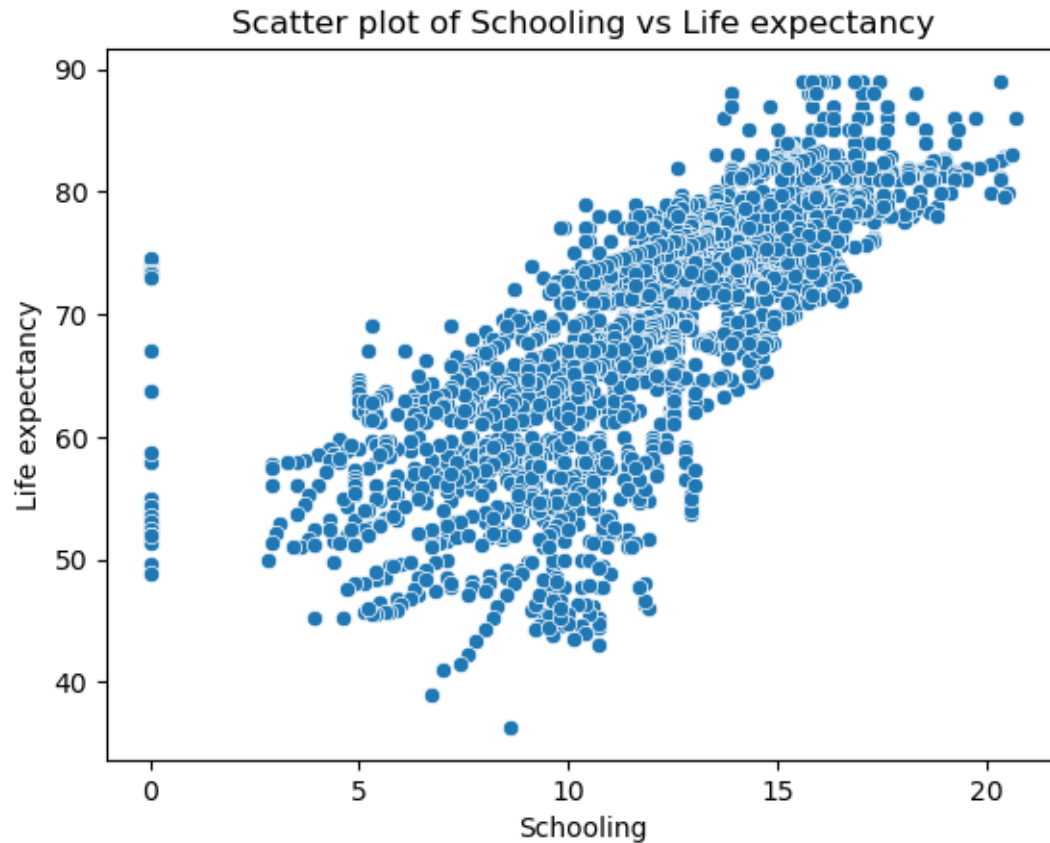












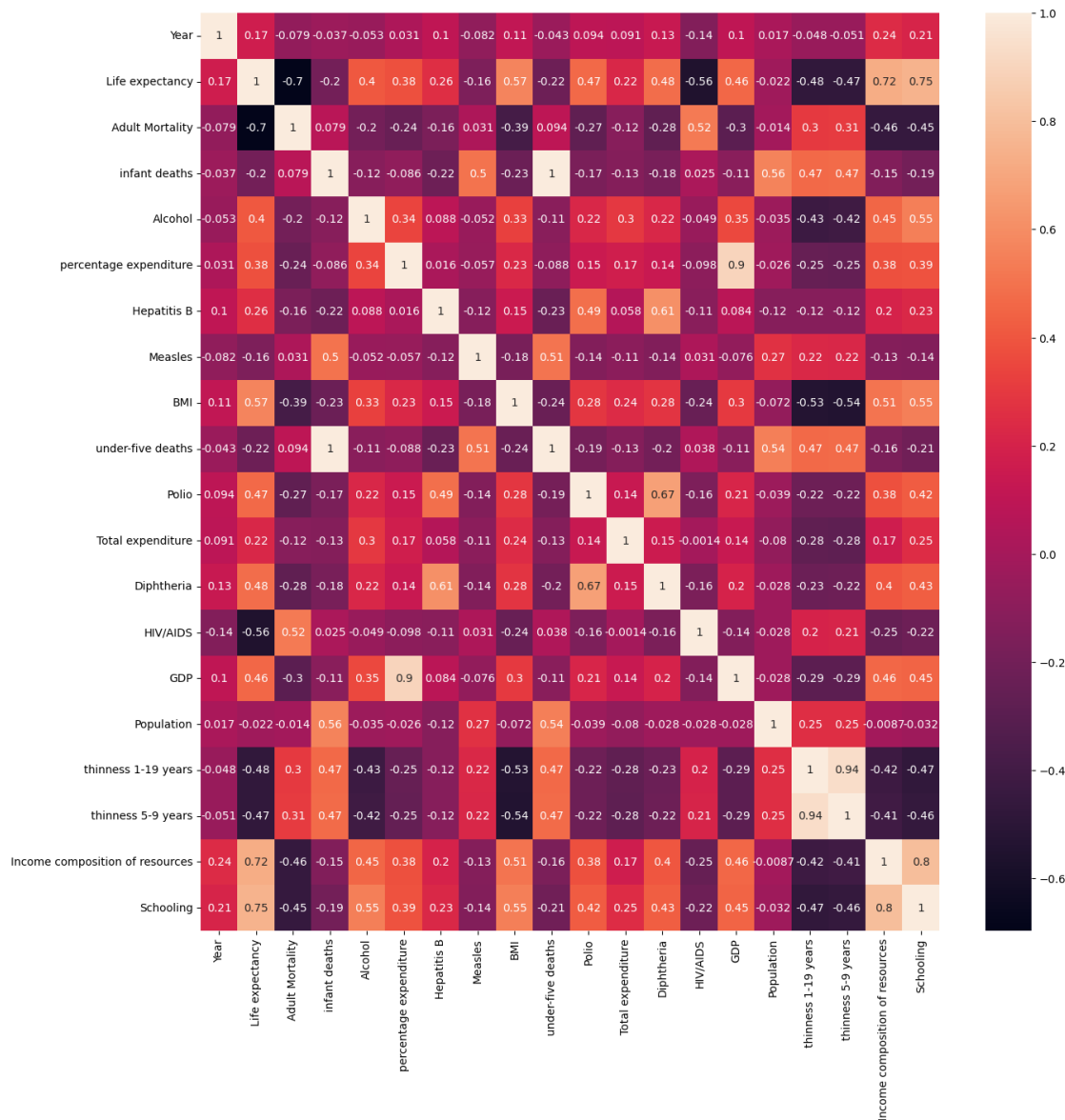
```
[23]: data.select_dtypes(include='number').columns
```

```
[23]: Index(['Year', 'Life expectancy ', 'Adult Mortality', 'infant deaths',
        'Alcohol', 'percentage expenditure', 'Hepatitis B', 'Measles ', ' BMI ',
        'under-five deaths ', 'Polio', 'Total expenditure', 'Diphtheria ',
        ' HIV/AIDS', 'GDP', 'Population', ' thinness 1-19 years',
        ' thinness 5-9 years', 'Income composition of resources', 'Schooling'],
        dtype='object')
```

```
[54]: ## correlation of heatmap
s=data.select_dtypes(include="number").corr()
```

```
[59]: plt.figure(figsize=(15, 15))
sns.heatmap(s,annot=True)
```

```
[59]: <Axes: >
```

2 Missing values treatments

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

```
[60]: Country      0
      Year         0
      Status       0
      Life expectancy 10
      Adult Mortality 10
      infant deaths  0
```

Alcohol	194
percentage expenditure	0
Hepatitis B	553
Measles	0
BMI	34
under-five deaths	0
Polio	19
Total expenditure	226
Diphtheria	19
HIV/AIDS	0
GDP	448
Population	652
thinness 1-19 years	34
thinness 5-9 years	34
Income composition of resources	167
Schooling	163
dtype:	int64

```
[61]: for i in ["BMI","Polio","Income composition of resources"]:
      data[i].fillna(data[i].median(),inplace=True)
```

C:\Users\Vikas\AppData\Local\Temp\ipykernel_23700\3528658117.py:2:
FutureWarning: A value is trying to be set on a copy of a DataFrame or Series through chained assignment using an inplace method.
The behavior will change in pandas 3.0. This inplace method will never work because the intermediate object on which we are setting values always behaves as a copy.

For example, when doing 'df[col].method(value, inplace=True)', try using 'df.method({col: value}, inplace=True)' or df[col] = df[col].method(value) instead, to perform the operation inplace on the original object.

```
data[i].fillna(data[i].median(),inplace=True)
```

C:\Users\Vikas\AppData\Local\Temp\ipykernel_23700\3528658117.py:2:
FutureWarning: A value is trying to be set on a copy of a DataFrame or Series through chained assignment using an inplace method.
The behavior will change in pandas 3.0. This inplace method will never work because the intermediate object on which we are setting values always behaves as a copy.

For example, when doing 'df[col].method(value, inplace=True)', try using 'df.method({col: value}, inplace=True)' or df[col] = df[col].method(value) instead, to perform the operation inplace on the original object.

```
data[i].fillna(data[i].median(),inplace=True)
```

C:\Users\Vikas\AppData\Local\Temp\ipykernel_23700\3528658117.py:2:

FutureWarning: A value is trying to be set on a copy of a DataFrame or Series through chained assignment using an inplace method.
The behavior will change in pandas 3.0. This inplace method will never work because the intermediate object on which we are setting values always behaves as a copy.

For example, when doing 'df[col].method(value, inplace=True)', try using 'df.method({col: value}, inplace=True)' or df[col] = df[col].method(value) instead, to perform the operation inplace on the original object.

```
data[i].fillna(data[i].median(),inplace=True)
```

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

```
[62]: Country          0
      Year            0
      Status          0
      Life expectancy  10
      Adult Mortality  10
      infant deaths    0
      Alcohol         194
      percentage expenditure  0
      Hepatitis B      553
      Measles          0
      BMI              0
      under-five deaths  0
      Polio            0
      Total expenditure  226
      Diphtheria       19
      HIV/AIDS         0
      GDP              448
      Population       652
      thinness 1-19 years  34
      thinness 5-9 years  34
      Income composition of resources  0
      Schooling        163
      dtype: int64
```

```
[64]: ## remove the missing values knnImputabser
      from sklearn.impute import KNNImputer
      impute=KNNImputer()
```

```
[65]: for i in data.select_dtypes(include="number").columns:
      data[i]=impute.fit_transform(data[[i]])
```

```
[66]: # after we check the null values
      data.isnull().sum()
```

```
[66]: Country          0
      Year            0
      Status          0
      Life expectancy  0
      Adult Mortality  0
      infant deaths    0
      Alcohol          0
      percentage expenditure 0
      Hepatitis B      0
      Measles          0
      BMI              0
      under-five deaths 0
      Polio            0
      Total expenditure 0
      Diphtheria       0
      HIV/AIDS         0
      GDP              0
      Population       0
      thinness 1-19 years 0
      thinness 5-9 years 0
      Income composition of resources 0
      Schooling        0
      dtype: int64
```

3 Outliers treatment

```
[77]: ## Decide whether to do outlier treatment or not, if do how? outlier treatment
      ↪we do the continues numerical columns
      def wisker(col):
          q1, q3 = np.percentile(col, [25, 75]) # Calculate the first and third
          ↪quartile
          iqr = q3 - q1 # Calculate the interquartile range (IQR)
          lw = q1 - 1.5 * iqr # Lower whisker
          uw = q3 + 1.5 * iqr # Upper whisker
          return lw, uw
```

```
[78]: data.columns
```

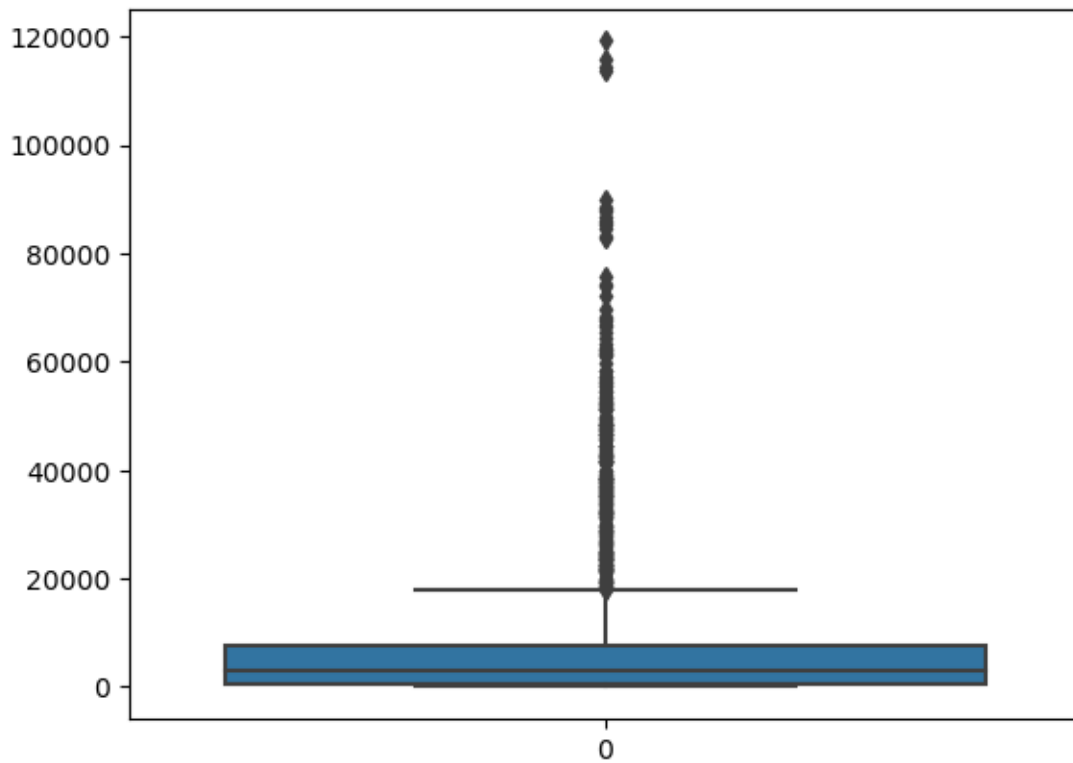
```
[78]: Index(['Country', 'Year', 'Status', 'Life expectancy', 'Adult Mortality',
          'infant deaths', 'Alcohol', 'percentage expenditure', 'Hepatitis B',
          'Measles', 'BMI', 'under-five deaths', 'Polio', 'Total expenditure',
          'Diphtheria', 'HIV/AIDS', 'GDP', 'Population', 'thinness 1-19 years',
          'thinness 5-9 years', 'Income composition of resources', 'Schooling'],
          dtype='object')
```

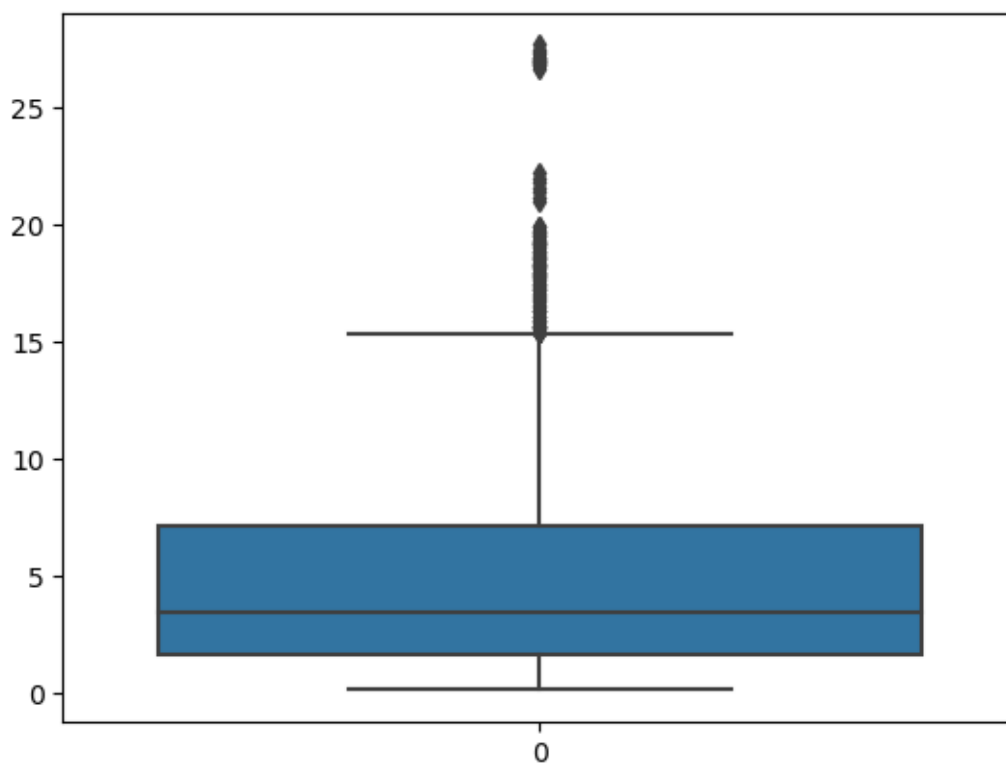
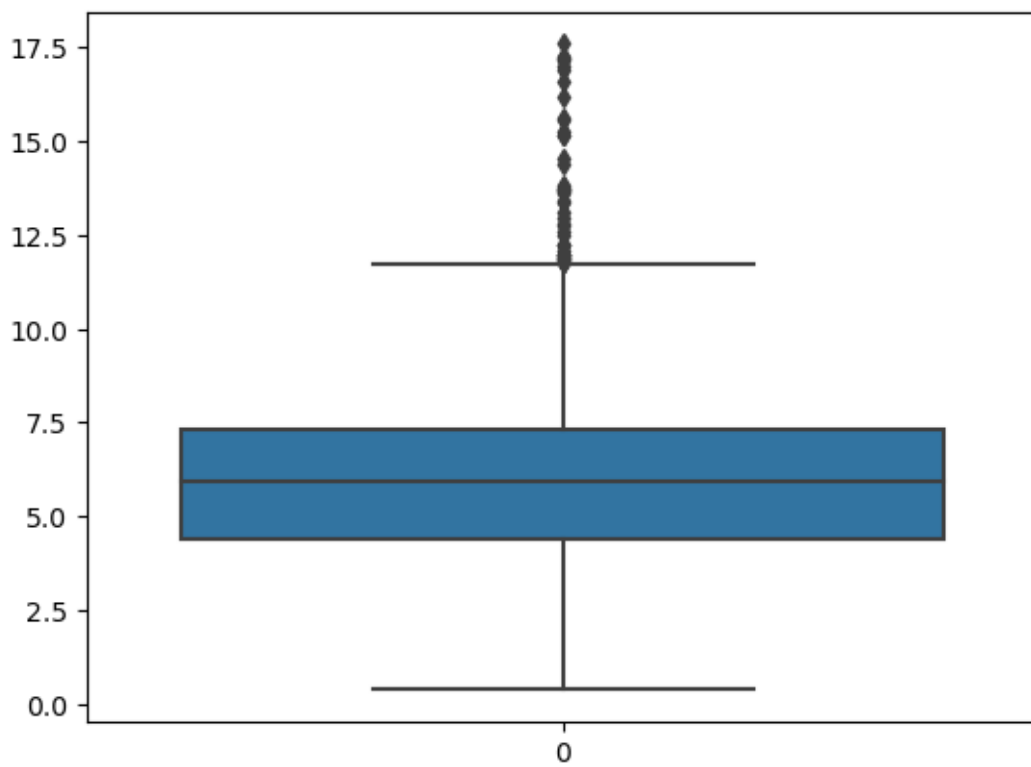
```
[80]: wisker(data['GDP'])
```

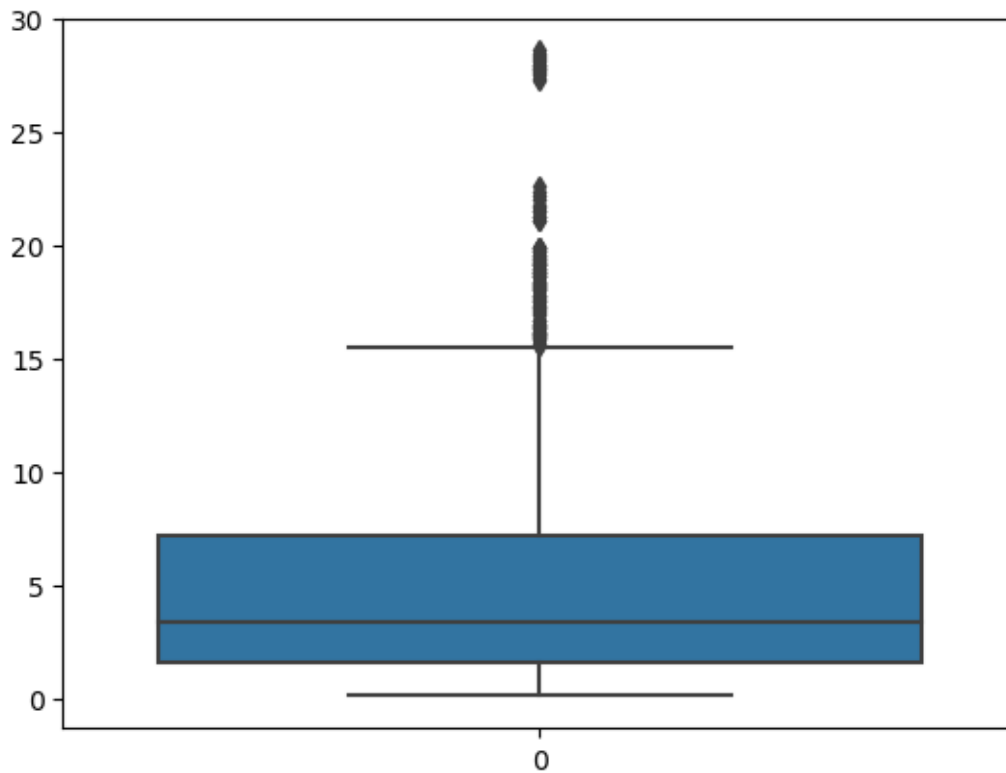
```
[80]: (-9773.52021495771, 17837.165679596183)
```

```
[ ]: for i in ['GDP', 'Total expenditure', 'thinness 1-19 years', 'thinness 5-9 years']:  
    lw, uw=wisker(data[i])  
    data[i]=np.where(data[i]<lw, lw, data[i])  
    data[i]=np.where(data[i]>uw, uw, data[i])
```

```
[81]: for i in ['GDP', 'Total expenditure', 'thinness 1-19 years', 'thinness 5-9 years']:  
    sns.boxplot(data[i])  
    plt.show()
```







```
[83]: dummy=pd.get_dummies(data=data,columns=["Country","Status"],drop_first=True)
```

```
[84]: dummy
```

```
[84]:
```

	Year	Life expectancy	Adult Mortality	infant deaths	Alcohol	\
0	2015.0	65.0	263.0	62.0	0.01	
1	2014.0	59.9	271.0	64.0	0.01	
2	2013.0	59.9	268.0	66.0	0.01	
3	2012.0	59.5	272.0	69.0	0.01	
4	2011.0	59.2	275.0	71.0	0.01	
...	
2933	2004.0	44.3	723.0	27.0	4.36	
2934	2003.0	44.5	715.0	26.0	4.06	
2935	2002.0	44.8	73.0	25.0	4.43	
2936	2001.0	45.3	686.0	25.0	1.72	
2937	2000.0	46.0	665.0	24.0	1.68	

	percentage expenditure	Hepatitis B	Measles	BMI	under-five deaths	\
0	71.279624	65.0	1154.0	19.1	83.0	

1	73.523582	62.0	492.0	18.6	86.0
2	73.219243	64.0	430.0	18.1	89.0
3	78.184215	67.0	2787.0	17.6	93.0
4	7.097109	68.0	3013.0	17.2	97.0
...
2933	0.000000	68.0	31.0	27.1	42.0
2934	0.000000	7.0	998.0	26.7	41.0
2935	0.000000	73.0	304.0	26.3	40.0
2936	0.000000	76.0	529.0	25.9	39.0
2937	0.000000	79.0	1483.0	25.5	39.0

	Country_United States of America	Country_Uruguay \
0	False	False
1	False	False
2	False	False
3	False	False
4	False	False
...
2933	False	False
2934	False	False
2935	False	False
2936	False	False
2937	False	False

	Country_Uzbekistan	Country_Vanuatu \
0	False	False
1	False	False
2	False	False
3	False	False
4	False	False
...
2933	False	False
2934	False	False
2935	False	False
2936	False	False
2937	False	False

	Country_Venezuela (Bolivarian Republic of)	Country_Viet Nam \
0	False	False
1	False	False
2	False	False
3	False	False
4	False	False
...
2933	False	False
2934	False	False
2935	False	False

2936		False	False
2937		False	False

	Country_Yemen	Country_Zambia	Country_Zimbabwe	Status_Developing
0	False	False	False	True
1	False	False	False	True
2	False	False	False	True
3	False	False	False	True
4	False	False	False	True
...
2933	False	False	True	True
2934	False	False	True	True
2935	False	False	True	True
2936	False	False	True	True
2937	False	False	True	True

[2938 rows x 213 columns]

[]: