

```
In [9]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from scipy import stats
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LassoCV
from sklearn.model_selection import RepeatedKFold
from sklearn.metrics import mean_squared_error
import warnings
```

```
In [2]: warnings.filterwarnings(action='ignore')
pd.options.mode.chained_assignment = None # default='warn' ---- ignores false warning
```

```
In [3]: df = pd.read_csv('medical_clean.csv')

df = df[['City', 'State', 'County', 'Zip', 'Lat', 'Lng', 'Population',
```

```
In [4]: # check for duplicates and null values
print(df.loc[df.duplicated()])
print(df.isnull().sum())
```

Empty DataFrame

Columns: [City, State, County, Zip, Lat, Lng, Population, Area, TimeZone, Job, Children, Age, Income, Marital, Gender, ReAdmis, VitD\_levels, Doc\_visits, Full\_meals\_eaten, vitD\_supp, Soft\_drink, Initial\_admin, HighBlood, Stroke, Complication\_risk, Overweight, Arthritis, Diabetes, Hyperlipidemia, BackPain, Anxiety, Allergic\_rhinitis, Reflux\_esophagitis, Asthma, Services, Initial\_days, TotalCharge, Additional\_charges]

Index: []

[0 rows x 38 columns]

City	0
State	0
County	0
Zip	0
Lat	0
Lng	0
Population	0
Area	0
TimeZone	0
Job	0
Children	0
Age	0
Income	0
Marital	0
Gender	0
ReAdmis	0
VitD_levels	0
Doc_visits	0
Full_meals_eaten	0
vitD_supp	0
Soft_drink	0
Initial_admin	0
HighBlood	0
Stroke	0
Complication_risk	0
Overweight	0

```

Arthritis      0
Diabetes       0
Hyperlipidemia 0
BackPain       0
Anxiety        0
Allergic_rhinitis 0
Reflux_esophagitis 0
Asthma         0
Services       0
Initial_days    0
TotalCharge    0
Additional_charges 0
dtype: int64

```

In [5]:

```

# check for outliers and remove (appears VitD_Levels contained the outliers)
print(df.shape)
df = df[(np.abs(stats.zscore(df.select_dtypes(include=np.number))) < 3).all(axis=1)]
print(df.shape)
print(df.head())

```

(10000, 38)

(9198, 38)

	City	State	County	Zip	Lat	Lng	Population	\
0	Eva	AL	Morgan	35621	34.34960	-86.72508	2951	
1	Marianna	FL	Jackson	32446	30.84513	-85.22907	11303	
2	Sioux Falls	SD	Minnehaha	57110	43.54321	-96.63772	17125	
3	New Richland	MN	Waseca	56072	43.89744	-93.51479	2162	
4	West Point	VA	King William	23181	37.59894	-76.88958	5287	

	Area	TimeZone	Job	...	\
0	Suburban	America/Chicago	Psychologist, sport and exercise	...	
1	Urban	America/Chicago	Community development worker	...	
2	Suburban	America/Chicago	Chief Executive Officer	...	
3	Suburban	America/Chicago	Early years teacher	...	
4	Rural	America/New_York	Health promotion specialist	...	

	Hyperlipidemia	BackPain	Anxiety	Allergic_rhinitis	Reflux_esophagitis	\
0	No	Yes	Yes	Yes	No	
1	No	No	No	No	Yes	
2	No	No	No	No	No	
3	No	No	No	No	Yes	
4	Yes	No	No	Yes	No	

	Asthma	Services	Initial_days	TotalCharge	Additional_charges
0	Yes	Blood Work	10.585770	3726.702860	17939.403420
1	No	Intravenous	15.129562	4193.190458	17612.998120
2	No	Blood Work	4.772177	2434.234222	17505.192460
3	Yes	Blood Work	1.714879	2127.830423	12993.437350
4	No	CT Scan	1.254807	2113.073274	3716.525786

[5 rows x 38 columns]

In [6]:

```

di = {'Yes': 1, 'No': 0}
di2 = {'Rural': 1, 'Suburban': 2, 'Urban': 3}
di3 = {'Divorced': 1, 'Married': 2, 'Widowed': 3, 'Never Married': 4, 'Separated': 5}
di4 = {'Male': 1, 'Female': 2, 'Nonbinary': 3}
di5 = {'Low': 1, 'Medium': 2, 'High': 3}
di6 = {'Blood Work': 1, 'Intravenous': 2, 'CT Scan': 3}
df = df.replace({'Area': di2, 'ReAdmis': di, 'Soft_drink': di, 'HighBlood': di, 'Stroke': di})
print(df.head())
df.to_csv('initial_clean.csv')

```

	City	State	County	Zip	Lat	Lng	Population	\
0	Eva	AL	Morgan	35621	34.34960	-86.72508	2951	
1	Marianna	FL	Jackson	32446	30.84513	-85.22907	11303	
2	Sioux Falls	SD	Minnehaha	57110	43.54321	-96.63772	17125	
3	New Richland	MN	Waseca	56072	43.89744	-93.51479	2162	
4	West Point	VA	King William	23181	37.59894	-76.88958	5287	

	Area	TimeZone	Job	...	\
0	2	America/Chicago	Psychologist, sport and exercise	...	
1	3	America/Chicago	Community development worker	...	
2	2	America/Chicago	Chief Executive Officer	...	
3	2	America/Chicago	Early years teacher	...	
4	1	America/New_York	Health promotion specialist	...	

	Hyperlipidemia	BackPain	Anxiety	Allergic_rhinitis	Reflux_esophagitis	\
0	0	1	1	1	0	
1	0	0	0	0	1	
2	0	0	0	0	0	
3	0	0	0	0	1	
4	1	0	0	1	0	

	Asthma	Services	Initial_days	TotalCharge	Additional_charges
0	1	1	10.585770	3726.702860	17939.403420
1	0	2	15.129562	4193.190458	17612.998120
2	0	1	4.772177	2434.234222	17505.192460
3	1	1	1.714879	2127.830423	12993.437350
4	0	3	1.254807	2113.073274	3716.525786

[5 rows x 38 columns]

```
In [7]: df.hist(figsize = (16,16))
plt.savefig('hospital_pyplot.jpg')
plt.tight_layout()
plt.close()
print('Histogram done')
```

Histogram done

```
In [8]: # bivariate analysis heatmap
ax = plt.subplots(figsize=(18,18))
ax = sns.heatmap(df.corr(), annot=True)
plt.savefig('heatmap_initial.jpg')
plt.close()
print('Initial heatmap done')
```

Initial heatmap done

```
In [10]: dfReduced = df[['Zip', 'Lat', 'Lng', 'Population', 'Age', 'ReAdmis', 'HighBl

print(dfReduced.head())
dfReduced.to_csv('reduced_clean.csv')
```

	Zip	Lat	Lng	Population	Age	ReAdmis	HighBlood	\
0	35621	34.34960	-86.72508	2951	53	0	1	
1	32446	30.84513	-85.22907	11303	51	0	1	
2	57110	43.54321	-96.63772	17125	53	0	1	
3	56072	43.89744	-93.51479	2162	78	0	0	
4	23181	37.59894	-76.88958	5287	22	0	0	

  

	Initial_days	TotalCharge	Additional_charges
0	10.585770	3726.702860	17939.403420

1	15.129562	4193.190458	17612.998120
2	4.772177	2434.234222	17505.192460
3	1.714879	2127.830423	12993.437350
4	1.254807	2113.073274	3716.525786

```
In [11]: X = np.array(dfReduced[['Zip', 'Lat', 'Lng', 'Population', 'ReAdmis', 'HighBlood'],
y = np.array(dfReduced[['Age']])
```

```
In [12]: X_train, X_test, y_train, y_test = train_test_split(X, y, train_size=0.7, random_state=
np.savetxt('training.csv', y_train)
np.savetxt('testing.csv', y_test)
```

```
In [13]: cv = RepeatedKfold(n_splits=10, n_repeats=3, random_state=1)
model = LassoCV(alphas=np.arange(0, 1, 0.01), cv=cv, n_jobs=-1)
c = model.fit(X_train, y_train)
print(model.alpha_)
d = model.score(X_test, y_test)
print(d)
e = model.predict(X_test)
np.savetxt('predictAges.csv', e)

# mean squared error
print(mean_squared_error(y_test, e))
```

```
0.02
0.889454080053387
46.98629349317084
```

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In [ ]:
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