Detect and Remove Outliers using Pandas

February 14, 2025

1 Detect and Remove Outliers using Python (Pandas)

https://www.geeksforgeeks.org/detect-and-remove-the-outliers-using-python/ An Outlier is a data item/object that deviates significantly from the rest of the (so-called normal) objects. Identifying outliers is important in statistics and data analysis because they can have a significant impact on the results of statistical analyses. The analysis for outlier detection is referred to as outlier mining.

Outliers can skew the mean (average) and affect measures of central tendency, as well as influence the results of tests of statistical significance.

1.1 Using barplot graph

1.1.1 Outlier detection

```
[1]: import sklearn
    from sklearn.datasets import load_diabetes
    import pandas as pd
    import matplotlib.pyplot as plt

diabetes = load_diabetes()

column_name = diabetes.feature_names
    df_diabetes = pd.DataFrame(diabetes.data)
    df_diabetes.columns = column_name
    print(df_diabetes.head())
    print(df_diabetes.info())
```

```
bmi
                                          s1
                                                   s2
                                                            s3
                                                               \
                                  bp
       age
               sex
 1 - 0.001882 - 0.044642 - 0.051474 - 0.026328 - 0.008449 - 0.019163 0.074412
2 0.085299 0.050680 0.044451 -0.005670 -0.045599 -0.034194 -0.032356
3 -0.089063 -0.044642 -0.011595 -0.036656 0.012191
                                             0.024991 -0.036038
 0.005383 -0.044642 -0.036385
                            0.021872 0.003935
                                             0.015596 0.008142
       s4
                s5
0 -0.002592 0.019907 -0.017646
1 -0.039493 -0.068332 -0.092204
2 -0.002592 0.002861 -0.025930
3 0.034309 0.022688 -0.009362
```

4 -0.002592 -0.031988 -0.046641 <class 'pandas.core.frame.DataFrame'> RangeIndex: 442 entries, 0 to 441 Data columns (total 10 columns):

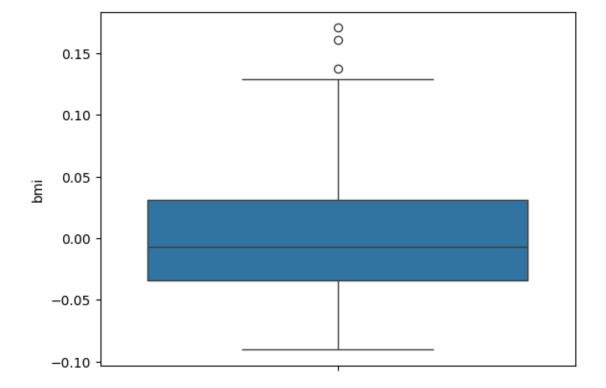
#	Column	Non-Null Count	Dtype
0	age	442 non-null	float64
1	sex	442 non-null	float64
2	bmi	442 non-null	float64
3	bp	442 non-null	float64
4	s1	442 non-null	float64
5	s2	442 non-null	float64
6	s3	442 non-null	float64
7	s4	442 non-null	float64
8	s5	442 non-null	float64
9	s6	442 non-null	float64
		/ >	

dtypes: float64(10)
memory usage: 34.7 KB

None

```
[2]: import seaborn as sns
sns.boxplot(df_diabetes['bmi'])
```

[2]: <Axes: ylabel='bmi'>



1.1.2 Outlier removal

```
import seaborn as sns
import matplotlib.pyplot as plt

def removal_box_plot(df, column, threshold):
    sns.boxplot(df[column])
    plt.title(f'Original Box Plot of {column}')
    plt.show()

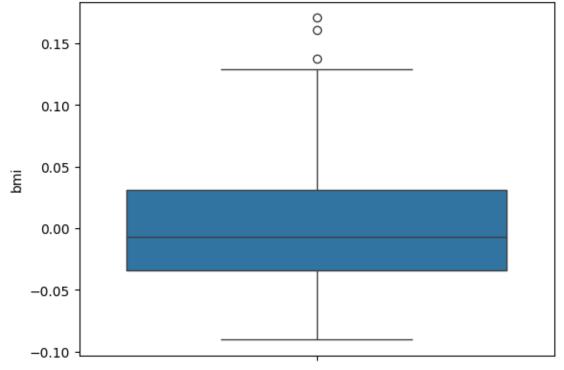
    removed_outliers = df[df[column] <= threshold]

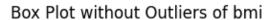
    sns.boxplot(removed_outliers[column])
    plt.title(f'Box Plot without Outliers of {column}')
    plt.show()
    return removed_outliers

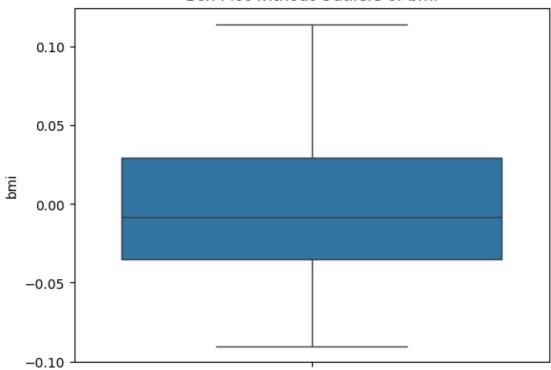
threshold_value = 0.12

no_outliers = removal_box_plot(df_diabetes,'bmi',threshold_value)</pre>
```

Original Box Plot of bmi



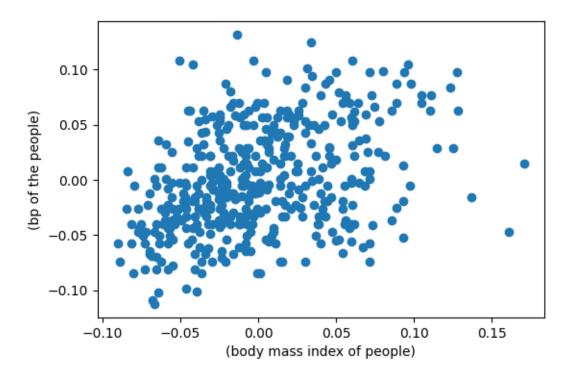




1.2 Using scatterplot graph

1.2.1 Outlier Detection

```
[4]: fig, ax = plt.subplots(figsize=(6,4))
ax.scatter(df_diabetes['bmi'], df_diabetes['bp'])
ax.set_xlabel('(body mass index of people)')
ax.set_ylabel('(bp of the people)')
plt.show()
```



1.2.2 Outlier Removal

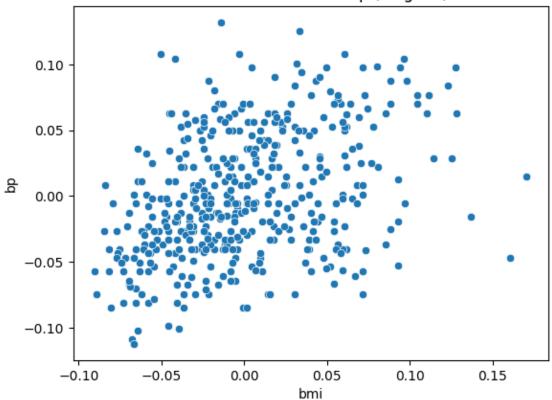
```
[10]: import numpy as np
      import seaborn as sns
      import matplotlib.pyplot as plt
      def removal_scatter_plot(df, column1, column2):
          # Scatter plot before removing outliers
          sns.scatterplot(x=df[column1], y=df[column2])
          plt.title(f'Scatter Plot of {column1} vs {column2} (Original)')
          plt.xlabel(column1)
          plt.ylabel(column2)
          plt.show()
          # Removing outliers using the specified condition
          outlier_indices = np.where((df[column1] > 0.12) & (df[column2] < 0.8))</pre>
          removed_outliers = df.drop(outlier_indices[0])
          # Check if there are any data points left after outlier removal
          if removed_outliers.empty:
              print(f"No data points remain after removing outliers with the given
       ⇔condition.")
          else:
              # Scatter plot after removing outliers
```

```
sns.scatterplot(x=removed_outliers[column1],__
y=removed_outliers[column2])
    plt.title(f'Scatter Plot of {column1} vs {column2} (Without Outliers)')
    plt.xlabel(column1)
    plt.ylabel(column2)
    plt.show()

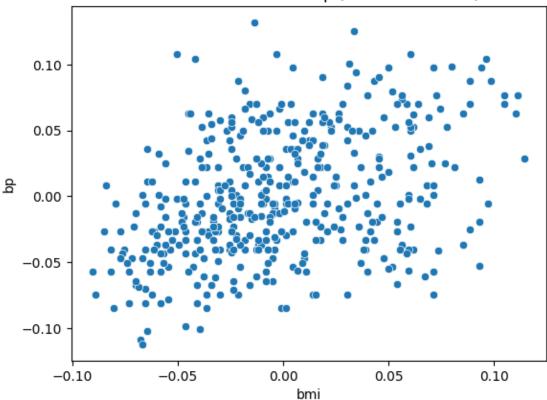
return removed_outliers

# Apply the function
no_outliers = removal_scatter_plot(df_diabetes, 'bmi', 'bp')
```

Scatter Plot of bmi vs bp (Original)



Scatter Plot of bmi vs bp (Without Outliers)



1.3 Using Z-score

Zscore = (data_point - mean) / std. deviation

1.3.1 Outlier detection

```
[29]: from scipy import stats import numpy as np

z = np.abs(stats.zscore(df_diabetes['age']))
print(z)
#print(df_diabetes['age'].info())

[0.80050009 0.03956713 1.79330681 1.87244107 0.11317236 1.94881082 0.9560041 1.33508832 0.87686984 1.49059233 2.02518057 0.57139085 0.34228161 0.11317236 0.95323959 1.1087436 0.11593688 1.48782782 0.80326461 0.57415536 1.03237385 1.79607132 1.79607132 0.95323959 1.33785284 1.41422259 2.25428981 0.49778562 1.10597908 1.41145807 1.26148309 0.49778562 0.72413034 0.6477606 0.34228161 1.02960933 0.26591186 0.19230663 0.03956713 0.03956713 0.11317236 2.10155031 1.26148309 0.41865135 0.95323959 0.57139085 1.18511334 1.64333183
```

```
1.41145807 0.87963435 0.72413034 1.25871858 1.1087436 0.19230663
1.03237385 0.87963435 0.87963435 0.57415536 0.87686984 1.33508832
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1.33508832 0.26591186 0.26591186 0.19230663 0.65052511 2.02518057
0.11317236 2.17792006 1.48782782 0.26591186 0.34504612 0.80326461
0.03680262 0.95323959 1.49059233 0.95323959 1.1087436 0.9560041
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1.18234883 0.57415536 0.03680262 0.03956713 0.34228161 0.34228161
1.94881082 1.25871858 0.57415536 0.4950211 2.02518057 0.57139085
0.41865135 0.80050009 0.87686984 0.41865135 1.79607132 0.41865135
1.03237385 1.33508832 1.02960933 0.11317236 0.11593688 0.11593688
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0.49778562 2.10155031 0.57415536 0.6477606 2.17792006 1.41145807
1.1087436  0.57415536  0.80326461  0.18954211  0.26591186  1.41145807
0.95323959 1.41145807 0.57139085 1.18234883 0.72413034 0.4950211
1.02960933 0.6477606 2.17792006 0.34228161 1.26148309 0.57415536
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0.34228161 0.03956713 0.26591186 1.56419757 0.87686984 0.19230663
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```

```
0.42141587 0.41865135 1.33785284 0.57139085 0.34504612 0.6477606
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1.79607132 1.10597908 0.26591186 1.25871858 0.49778562 0.34228161
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0.11593688 0.87686984 0.9560041 0.9560041 ]
```

1.3.2 Identifying Threshold value using z-score

```
[32]: from scipy import stats
import numpy as np

# Calculate z-scores for the 'age' column
z = np.abs(stats.zscore(df_diabetes['age']))

# Define the threshold using the 95th percentile
threshold = np.percentile(z, 95)

# Print the threshold value
print("Threshold value:", threshold)

# Identify outliers
outliers = np.where(z > threshold)

# Print the results
print("Z-scores:", z)
print("Outliers:", outliers)
```

```
Threshold value: 1.8724410718097853

Z-scores: [0.80050009 0.03956713 1.79330681 1.87244107 0.11317236 1.94881082 0.9560041 1.33508832 0.87686984 1.49059233 2.02518057 0.57139085 0.34228161 0.11317236 0.95323959 1.1087436 0.11593688 1.48782782 0.80326461 0.57415536 1.03237385 1.79607132 1.79607132 0.95323959 1.33785284 1.41422259 2.25428981 0.49778562 1.10597908 1.41145807 1.26148309 0.49778562 0.72413034 0.6477606 0.34228161 1.02960933 0.26591186 0.19230663 0.03956713 0.03956713 0.11317236 2.10155031
```

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1.26148309 0.41865135 0.95323959 0.57139085 1.18511334 1.64333183
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0.6477606  0.03956713  0.95323959  1.56419757  0.80326461  0.26867637
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0.18954211 1.71693706 0.6477606 0.57139085 1.26148309 0.11317236
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0.49778562 1.56696208 0.11593688 1.26148309 0.42141587 0.80050009
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0.9560041 0.19230663 0.34504612 0.19230663 0.41865135 1.10597908
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0.34228161 0.03956713 0.34228161 1.49059233 1.02960933 0.11317236
0.72413034 0.4950211 0.41865135 0.9560041 1.10597908 0.11593688
0.18954211 0.49778562 0.87963435 1.56696208 0.72413034 1.26148309
1.79607132 1.10597908 0.26591186 1.25871858 0.49778562 0.34228161
2.32789504 0.42141587 0.34504612 1.02960933 1.18511334 0.57139085
1.33508832 1.1087436 0.19230663 0.11317236 1.56419757 1.1087436
1.71693706 0.11593688 0.57415536 1.1087436 0.18954211 0.42141587
0.18954211 0.57415536 0.34228161 0.26867637 1.18511334 0.87686984
0.11593688 0.87686984 0.9560041 0.9560041 ]
Outliers: (array([ 5, 10, 26, 41, 77, 79, 102, 106, 131, 136, 204, 211,
223,
     226, 242, 281, 311, 321, 344, 374, 402]),)
```

1.3.3 Outlier Removal

```
[28]: import numpy as np
      def removal_outliers_zscore(df, column, z_column, threshold):
          # Calculate outlier indices based on z-score threshold for the specified,
       ⇔column
          outlier_indices = np.where(z_column > threshold)[0]
          removed_outliers = df.drop(outlier_indices)
          # Print DataFrame shapes
          print(f"Original DataFrame Shape: {df.shape}")
          print(f"DataFrame Shape after Removing Outliers: {removed_outliers.shape}")
          return removed_outliers
      # Define the z-score threshold
      threshold z = 2
      # Calculate z-scores for the 'age' column
      z = np.abs((df_diabetes['age'] - df_diabetes['age'].mean()) /__

¬df_diabetes['age'].std())
      # Apply the function
```

```
no_outliers = removal_outliers_zscore(df_diabetes, 'age', z, threshold_z)
```

```
Original DataFrame Shape: (442, 10)
DataFrame Shape after Removing Outliers: (426, 10)
```

1.4 Using IQR (Inter Quartile Range)

```
IQR = Quartile3 - Quartile1
```

Syntax: numpy.percentile(arr, n, axis=None, out=None) Parameters: arr :input array. n : percentile value.

```
[34]: Q1 = np.percentile(df_diabetes['bmi'], 25, method='midpoint')
Q3 = np.percentile(df_diabetes['bmi'], 75, method='midpoint')
IQR = Q3 - Q1
print(IQR)
```

0.06520763046978838

To define the outlier base value is defined above and below dataset's normal range namely Upper and Lower bounds, define the upper and the lower bound (1.5*IQR value is considered):

```
upper = Q3 + 1.5IQR lower = Q1 - 1.5IQR
```

In the above formula as according to statistics, the 0.5 scale-up of IQR (new_IQR = IQR + 0.5*IQR) is taken, to consider all the data between 2.7 standard deviations in the Gaussian Distribution.

```
[37]: upper = Q3 + 1.5 * IQR
    upper_array = np.array(df_diabetes['bmi'] >= upper)
    print("Upper Bound:", upper)
    print(upper_array.sum())

lower = Q1 - 1.5 * IQR
    lower_array = np.array(df_diabetes['bmi'] <= lower)
    print("Lower Bound:", lower)
    print(lower_array.sum())</pre>
```

```
Upper Bound: 0.12879000811776306
3
Lower Bound: -0.13204051376139045
```

1.5 Outlier removal in Dataset using IQR

```
[5]: import sklearn
from sklearn.datasets import load_diabetes
import pandas as pd
import numpy as np
```

```
diabetes = load_diabetes()
column_name = diabetes.feature_names
df_diabetes = pd.DataFrame(diabetes.data)
df_diabetes.columns = column_name
df_diabetes.head()
print("Old Shape:", df_diabetes.shape)
Q1 = df_diabetes['bmi'].quantile(0.25)
Q3 = df_diabetes['bmi'].quantile(0.75)
IQR = Q3 - Q1
lower = Q1 - 1.5 * IQR
upper = Q3 + 1.5 * IQR
upper_array = np.where(df_diabetes['bmi'] >= upper)[0]
lower_array = np.where(df_diabetes['bmi'] <= lower)[0]</pre>
df_diabetes.drop(index=upper_array, inplace=True)
df_diabetes.drop(index=lower_array, inplace=True)
print("New Shape:", df_diabetes.shape)
Old Shape: (442, 10)
New Shape: (439, 10)
```

[]: