Statistics Practical Implementations in Python

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Measure of Central Tendency

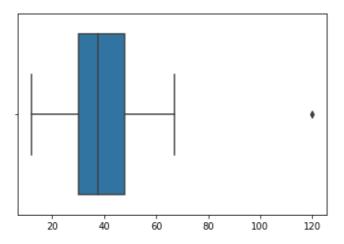
esult in an error or misinterpretation.

warnings.warn(

<AxesSubplot:>

Out[10]:

```
1. Mean
2. Median
3. Mode
In [5]:
## DataSet lists For Practices:-
ages = [23, 24, 32, 45, 12, 43, 67, 45, 32, 56, 32, 120]
In [6]:
## Finding Mean & Median via Numpy Library.
import numpy as np
print(np.mean(ages))
print(np.median(ages))
44.25
37.5
In [7]:
## We can also find Mean & Median via Statistics Library.
import statistics
print(statistics.mean(ages))
print(statistics.median(ages))
44.25
37.5
In [8]:
## Findind Mode via Statistics Library.
import statistics
statistics.mode(ages)
Out[8]:
In [10]:
## For Finding Outlier, We Basically Use Boxplox.
import seaborn as sns
sns.boxplot(ages)
C:\Users\Lenovo\anaconda3\lib\site-packages\seaborn\ decorators.py:36: FutureWarning: Pas
s the following variable as a keyword arg: x. From version 0.12, the only valid positiona
l argument will be `data`, and passing other arguments without an explicit keyword will r
```



5 Number Summary

```
In [12]:
# To Find q1 And q3 in Dataset...
## Where, q1 is 25 Percentile And q3 is 75 percentile.
import numpy as np
q1, q3 = np.percentile(ages,[25,75])
In [23]:
## Here we are finding the value of q1 and q3.
```

```
## Here we are finding the value of q1 and q3.
print( f"q1 is {q1} and q3 is {q3}")
```

q1 is 30.0 and q3 is 47.75

In [24]:

```
# To check outlier [Lower Fence - Higher Fence]
# Where IQR is Inter Quantile Range.

IQR = q3 -q1
lower_fence = q1 - 1.5*(IQR)
higher_fence = q3 + 1.5*(IQR)
print(f"Lower Fence is {lower_fence} and Higher Fence is {higher_fence}")
```

Lower Fence is 3.375 and Higher Fence is 74.375

Measure Of Dispersion

1. Variance

2. Standard Deviation

```
In [26]:
```

```
## In Statistics Library Variance are Comes via sample Variances Formuala statistics.variance(ages)
```

```
Out[26]:
```

795.2954545454545

In [27]:

```
np.var(ages, axis = 0) ### In Numpy Library Variance are comes via Population Variances Formula.
```

Out[27]:

```
729.0208333333334
```

```
In [28]:
```

```
## We are Approaching Manual Step to Find Population Variance
def variance(data):
   n = len(ages)
    ## mean of the data
   mean = sum(data)/n
    ## Variance
    deviation = [(x - mean) ** 2 for x in data]
    variance = sum(deviation)/n ## here we are using Population Variance formula
   return variance
In [39]:
variance (ages)
```

Out[39]:

729.0208333333334

In [46]:

```
## We are Approaching Manual Step to Find Sample Variance
def variance(data):
   n = len(ages)
    ## mean of the data
   mean = sum(data)/n
    ## Variance
   deviation = [(x - mean) ** 2 for x in data]
   variance = sum(deviation)/(n-1) ## here we are using Sample Variance Formula.
   return variance
```

In [47]:

```
variance (ages)
```

Out[47]:

795.2954545454545

In [50]:

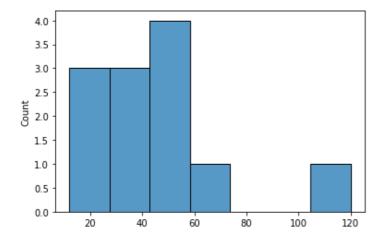
```
# Now, We are doing Manual Approaching for Finding Variances with Degree of Freedom (dof)
## Where dof = 0 means it is Population variances.
## Where dof = 1 means it is sample variances.
def variance(data, dof = 0):
   n = len(ages)
    ## mean of the data
   mean = sum(data)/n
   ## Variance
   deviation = [(x - mean) ** 2 for x in data]
   variance = sum(deviation)/(n- dof) ## here we are using Sample Variance Formula.
   return variance
```

In [53]:

```
## here we are finding Population Variance with dof =0
variance(ages, dof =0)
```

```
Out[53]:
729.0208333333334
In [54]:
## here we are finding Sample Variance with dof =1.
variance(ages, dof =1)
Out[54]:
795.2954545454545
In [57]:
## In Statistics Library , we are finding sample Variance as below :-
statistics.variance(ages)
Out[57]:
795.2954545454545
In [58]:
## In Statistics Library, we are finding Population Variance as below :-
statistics.pvariance(ages)
Out[58]:
729.0208333333334
In [59]:
## For Finding Standard Deviation in statistics we use Maath Library.
import math
math.sqrt(statistics.pvariance(ages))
Out[59]:
27.000385799712813
Histograms And PDF
In [60]:
import seaborn as sns
sns.histplot(ages)
Out[60]:
```

<AxesSubplot:ylabel='Count'>



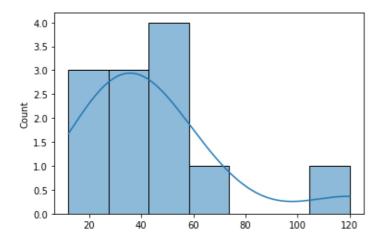
```
In [61]:
```

import seaborn as sns

```
sns.histplot(ages, kde = True) ## Where Kde = kernal density Estimators
```

Out[61]:

<AxesSubplot:ylabel='Count'>



In [62]:

```
## Now, we check practical with some dataset:-

df = sns.load_dataset('iris')
```

In [64]:

```
## head() basically used for top 5 data present in any dataset
df.head()
```

Out[64]:

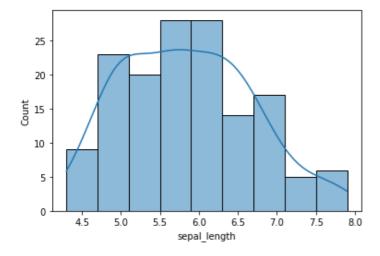
	sepal_length	sepal_width	petal_length	petal_width	species
0	5.1	3.5	1.4	0.2	setosa
1	4.9	3.0	1.4	0.2	setosa
2	4.7	3.2	1.3	0.2	setosa
3	4.6	3.1	1.5	0.2	setosa
4	5.0	3.6	1.4	0.2	setosa

In [65]:

```
sns.histplot(df['sepal_length'], kde = True)
```

Out.[65]:

<AxesSubplot:xlabel='sepal_length', ylabel='Count'>



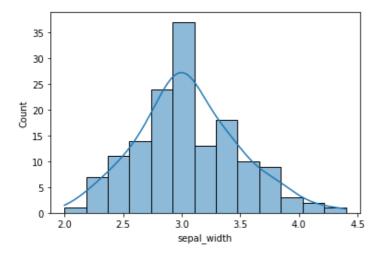
In [66]:

sns histolot (df['senal width'] kde = True)

DIDONITED CREEK CALL DONAL WINGE IN MAC ILUC!

Out[66]:

<AxesSubplot:xlabel='sepal_width', ylabel='Count'>

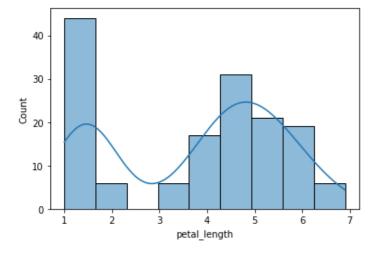


In [67]:

```
sns.histplot(df['petal length'], kde = True)
```

Out[67]:

<AxesSubplot:xlabel='petal_length', ylabel='Count'>

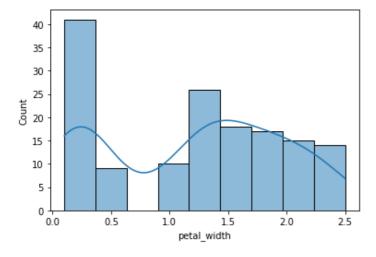


In [68]:

```
sns.histplot(df['petal_width'], kde = True)
```

Out[68]:

<AxesSubplot:xlabel='petal_width', ylabel='Count'>



In [69]:

How to Charte a Normal Distributed Date with the halo of Normal

```
s = np.random.normal(0.5,0.2,1000)
```

In [70]:

S

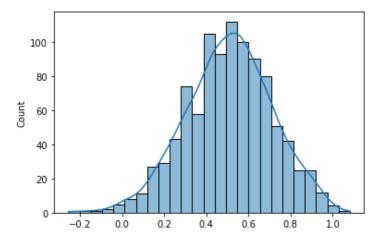
Out[70]:

```
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```

```
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                                      0.58337134, 0.44831011,
0.60837269,
0.86297816,
             0.22612841, 0.35540339, 0.65471009, 0.37348811,
0.78254754,
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0.19187389, 0.52205473, 0.6943523, 0.50771795, 0.90842507,
0.56352208, 0.80372771, 0.29839909, 0.87376001, 0.50740368,
0.64445538, 0.87769576, 0.70409311, 0.42865279, 0.38760179,
```

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0.50948434,	0.2801032 ,	0.65500205,	0.45845769,	0.4928895 ,
•		•	•	•
0.69861493,	0.39686519,	0.61688589,	0.7057442 ,	0.47549625,
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•	•			•
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0.41857073,	0.55611789,	0.57307487,	0.56814482,	0.5204143 ,
•	•	· · · '	•	•
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0.92482301,	0.61970749,	0.5288567,	0.08224764,	0.63566643,
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•		· ·		•
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0.18120982,	0.48097886,	0.60338321,	0.26968114,	0.88911387,
•	•	·	•	•
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	0.0094769 ,	•	•	•
0.81149316,	0.07477097,	0.63572231,	-0.0192333 ,	0.91736001,
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0.47108847,	0.65550511,	0.29168258,	0.77578801,	0.62602314,
0.53007176,	0.6189141 ,	0.98419809,	0.78040766,	0.45319585,
-0.08478568,	0.94572126,	0.54690887,	0.03163383,	0.57363574,
0.7510476	0.4842176 ,	0.42652436,	0.89163488,	0.3757975])
	, ,	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	,	,

<AxesSubplot:ylabel='Count'>



Other Distribution

Log Normal Distribution, Power Law distribution

```
In [78]:
```

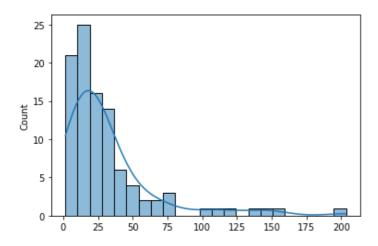
```
## To Find Log Normal Distributions:-
mu, sigma = 3., 1. ## Where mu is Means and sigma is Standard deviation.
s = np.random.lognormal(mu, sigma, 100)
```

In [79]:

```
## Histogram Plot for Log Normal Distribution dataset
sns.histplot(s, kde = True)
```

Out[79]:

<AxesSubplot:ylabel='Count'>

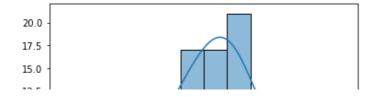


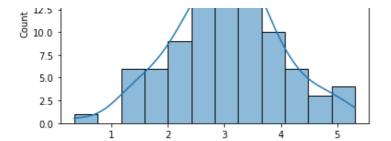
In [80]:

```
sns.histplot(np.log(s), kde = True)
```

Out[80]:

<AxesSubplot:ylabel='Count'>





Check Whether Distribution is Normal Distribution or not.

```
In [83]:
```

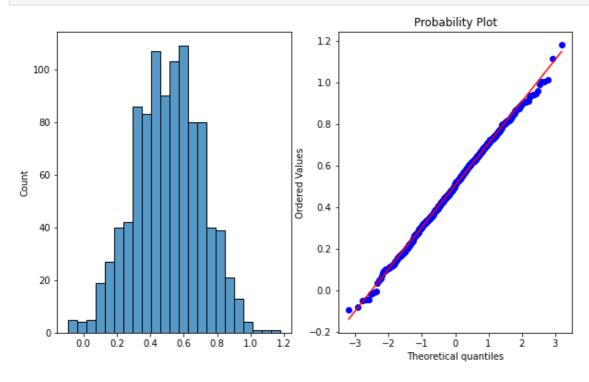
```
# If you want to check whether feature is Guassian or Normal distribution .
## Q-Q Plot

import matplotlib.pyplot as plt
import scipy.stats as stat
import pylab

def plot_data(sample):
   plt.figure(figsize = (10,6))
   plt.subplot(1,2,1)
   sns.histplot(sample)
   plt.subplot(1,2,2)
   stat.probplot(sample, dist='norm', plot = pylab)
   plt.show()
```

In [84]:

```
## Creating a Normal Distributed data:-
s = np.random.normal(0.5, 0.2, 1000)
plot_data(s)  ## Here, in below Result data comes in a straight line of Probability Plot
, so it is a Normal Distribution.
```

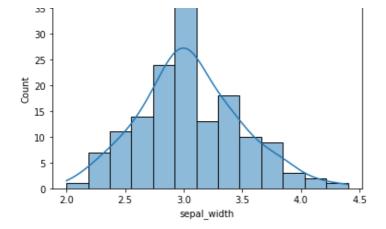


```
In [86]:
```

```
sns.histplot(df['sepal_width'], kde= True)
```

Out[86]:

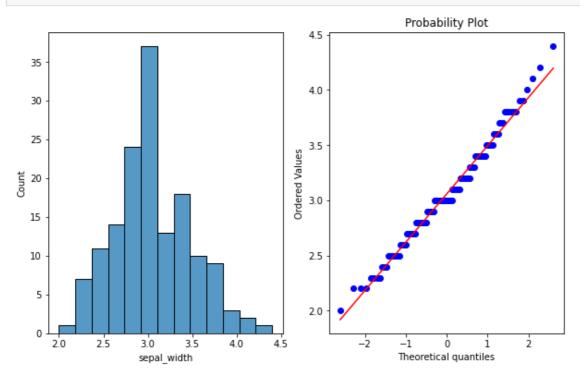
```
<AxesSubplot:xlabel='sepal_width', ylabel='Count'>
```



In [87]:

Here, in below Result data comes in a straight line of Probability Plot, so it is a No rmal Distribution.

```
plot_data(df['sepal_width'])
```



Check Whether Distribution is Log Normal Distribution or not.

```
In [92]:
```

```
## To Find Log Normal Distributions:-
mu, sigma = 3., 1. ## Where mu is Means and sigma is Standard deviation.
sample = np.random.lognormal(mu, sigma, 100)
```

In [93]:

```
## If you want to check whether feature is Log Normal distribution .

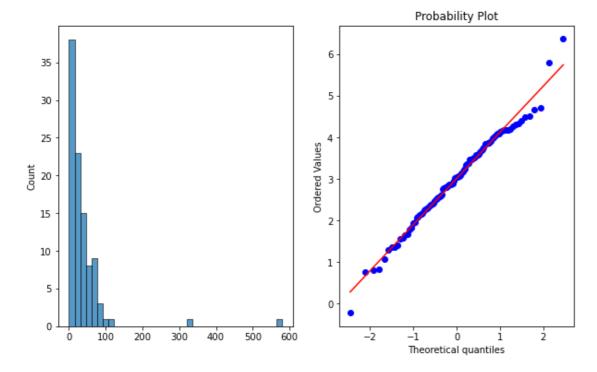
import matplotlib.pyplot as plt
import scipy.stats as stat
import pylab

def plot_data(sample):
    plt.figure(figsize = (10,6))
    plt.subplot(1,2,1)
```

```
sns.histplot(sample)
plt.subplot(1,2,2)
stat.probplot(np.log(sample), dist='norm', plot = pylab)
plt.show
```

In [94]:

```
## Here, in below Result data comes in a straight line of Probability Plot, so it is a L
og Normal Distribution.
plot_data(sample)
```



Pearson and Sperman Rank Correlation

```
In [95]:
```

```
## For Practicing we are taking Tips Dataset:-
df = sns.load_dataset('tips')
```

In [96]:

df.head()

Out[96]:

	total_bill	tip	sex	smoker	day	time	size
0	16.99	1.01	Female	No	Sun	Dinner	2
1	10.34	1.66	Male	No	Sun	Dinner	3
2	21.01	3.50	Male	No	Sun	Dinner	3
3	23.68	3.31	Male	No	Sun	Dinner	2
4	24.59	3.61	Female	No	Sun	Dinner	4

In [97]:

```
import pandas as pd
```

In [98]:

```
## In Pandas Dafault, Corr() takes Pearson Correlation:-
df.corr()
```

Out[98]:

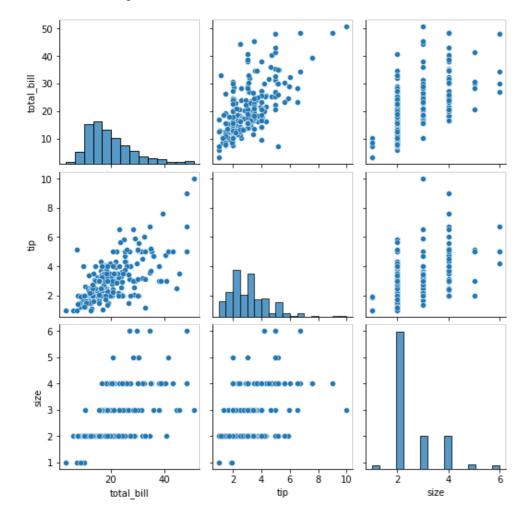
	total_bill	tip	size
total_bill	1.000000	0.675734	0.598315
tip	0.675734	1.000000	0.489299
size	0.598315	0.489299	1.000000

In [99]:

To Check Pearson Correlation in Diagramatically format, we use. sns.pairplot(df)

Out[99]:

<seaborn.axisgrid.PairGrid at 0x1edc25ea550>



Thank You