

CPE311 Computational Thinking with Python**Name:** Ballesteros, Angelo**Section:** CPE22S2**Performed on:** 6/20/2024**Submitted on:** 6/20/2024**Submitted to:** Engr. Roman M. Richard**6.1 Intended Learning Outcomes**

1. Use pandas and numpy data analysis tools.
2. Demonstrate how to analyze data using numpy and pandas

6.2 Resources:

- Personal Computer
- Jupyter Notebook
- Internet Connection

✓ **6.3 Supplementary Activities:****Exercise 1**

Run the given code below for exercises 1 and 2, perform the given tasks without using any Python modules.

```
: import random
random.seed(0)
salaries = [round(random.random()*1000000, -3) for _ in range(100)]
```

Using the data generated above, calculate the following statistics without importing anything from the statistics module in the standard library (<https://docs.python.org/3/library/statistics.html>) and then confirm your results match up to those that are obtained when using the statistics module (where possible):

- Mean
- Median
- Mode (hint: check out the Counter in the collections module of the standard library at <https://docs.python.org/3/library/collections.html#collections.Counter>)
- Sample variance
- Sample standard deviation

```
import random # data generated
random.seed(0)
salaries = [round(random.random()*1000000, -3) for _ in range(100)]
```

```
from statistics import median
from math import isnan
from itertools import filterfalse
```

```
def cal_mean(data): # mean function
    return sum(data) / len(data)
```

```
mean = cal_mean(salaries)
```

```

def cal_median(data): # median function
    sorted_data = sorted(data)
    n = len(sorted_data)
    midpoint = n // 2

    if n % 2 == 1:
        return sorted_data[midpoint]
    else:
        return (sorted_data[midpoint - 1] + sorted_data[midpoint]) / 2

median = cal_median(salaries)

from collections import Counter

def cal_mode(data): # mode function
    frequency = Counter(data)
    mode_data = frequency.most_common(1)
    return mode_data[0][0] if mode_data else None

mode = cal_mode(salaries)

def cal_sampleVar(data): # sample variance function
    mean = cal_mean(data)
    squared_diff = [(x - mean) ** 2 for x in data]
    return sum(squared_diff) / (len(data) - 1)

sampleVariance = cal_sampleVar(salaries)

def cal_standardDev(data): # sample standard deviation function
    variance = cal_sampleVar(data)
    return variance ** 0.5

sample_standardDev = cal_standardDev(salaries)

# output
print(f"Mean:", mean)
print(f"Median:", median)
print(f"Mode:", mode)
print(f"Sample Variance:", sampleVariance)
print(f"Sample Standard Deviation:", sample_standardDev)

```

```

→ Mean: 585690.0
   Median: 589000.0
   Mode: 477000.0
   Sample Variance: 70664054444.44444
   Sample Standard Deviation: 265827.11382484

```

```
# reference of output
```

```
import pandas as pd
```

```
df = pd.DataFrame(salaries)
df.describe()
```

```

→

```

	0
count	100.000000
mean	585690.000000
std	265827.113825
min	1000.000000
25%	403500.000000
50%	589000.000000
75%	816750.000000
max	996000.000000

Exercise 2

Using the same data, calculate the following statistics using the functions in the statistics module where appropriate:

- Range
- Coefficient of variation Interquartile range
- Quartile coefficient of dispersion

```
range = max(salaries) - min(salaries) # range function

mean = cal_mean(salaries) # COV function
standardDev = cal_standardDev(salaries)
COV = (standardDev / mean) * 100

def cal_iqr(data): # interquartile range function
    sorted_data = sorted(data)
    q1 = cal_median(sorted_data[:len(sorted_data) // 2])
    q3 = cal_median(sorted_data[(len(sorted_data) + 1) // 2:])
    return q3 - q1


iqr = cal_iqr(salaries)

def cal_qd(data): # quartile dispersion function
    sorted_data = sorted(data)
    q1 = cal_median(sorted_data[:len(sorted_data) // 2])
    q3 = cal_median(sorted_data[(len(sorted_data) + 1) // 2:])
    return (q3 - q1) / (q3 + q1)

qd = cal_qd(salaries)

# output

print(f"Range:", range)
print(f"Coefficient of Variation:%", COV)
print(f"Interquartile Range:", iqr)
print(f"Quartile Coefficient of Dispersion:", qd)
```

 Range: 995000.0
Coefficient of Variation:% 45.38699889443903
Interquartile Range: 417500.0
Quartile Coefficient of Dispersion: 0.3417928776094965

Exercise 3: Pandas for Data Analysis

Load the diabetes.csv file. Convert the diabetes.csv into dataframe

Perform the following tasks in the diabetes dataframe:

1. Identify the column names
2. Identify the data types of the data
3. Display the total number of records
4. Display the first 20 records
5. Display the last 20 records
6. Change the Outcome column to Diagnosis

-
7. Create a new column Classification that display "Diabetes" if the value of outcome is 1 , otherwise "No Diabetes"
 8. Create a new dataframe "withDiabetes" that gathers data with diabetes
 9. Create a new dataframe "noDiabetes" thats gathers data with no diabetes
 10. Create a new dataframe "Pedia" that gathers data with age 0 to 19
 11. Create a new dataframe "Adult" that gathers data with age greater than 19
 12. Use numpy to get the average age and glucose value.
 13. Use numpy to get the median age and glucose value.
 14. Use numpy to get the middle values of glucose and age.
 15. Use numpy to get the standard deviation of the skinthickness.

```
# uploading csv and convert it into dataframe
```


```
import pandas as pd
from google.colab import files
```

```
uploaded = files.upload()
```

```
diabetes = list(uploaded.keys())[0]
```

```
df = pd.read_csv(diabetes)
```

```
print(df)
```

 Choose Files diabetes.csv

- **diabetes.csv**(text/csv) - 23873 bytes, last modified: 6/20/2024 - 100% done

Saving diabetes.csv to diabetes (2).csv


	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	\
0	6	148	72	35	0	33.6	
1	1	85	66	29	0	26.6	
2	8	183	64	0	0	23.3	
3	1	89	66	23	94	28.1	
4	0	137	40	35	168	43.1	
..	
763	10	101	76	48	180	32.9	
764	2	122	70	27	0	36.8	
765	5	121	72	23	112	26.2	
766	1	126	60	0	0	30.1	
767	1	93	70	31	0	30.4	

	DiabetesPedigreeFunction	Age	Outcome
0	0.627	50	1
1	0.351	31	0
2	0.672	32	1
3	0.167	21	0
4	2.288	33	1
..
763	0.171	63	0
764	0.340	27	0
765	0.245	30	0
766	0.349	47	1
767	0.315	23	0

```
[768 rows x 9 columns]
```


```
column_names = list(df.columns) # identify column names
```

```
print(column_names)
```

 ['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome']

```
df_types = df.dtypes # identify the data types
```


```
print(df_types)
```

 Pregnancies int64
 Glucose int64
 BloodPressure int64
 SkinThickness int64
 Insulin int64
 BMI float64
 DiabetesPedigreeFunction float64
 Age int64
 Outcome int64
 dtype: object

```
print(f"Number of records: {df.shape[0]}") # total number of records
```

 Number of records: 768

```
print(df.head(20)) # first 20 records
```




	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	\
0	6	148	72	35	0	33.6	
1	1	85	66	29	0	26.6	
2	8	183	64	0	0	23.3	
3	1	89	66	23	94	28.1	
4	0	137	40	35	168	43.1	
5	5	116	74	0	0	25.6	

6	3	78	50	32	88	31.0
7	10	115	0	0	0	35.3
8	2	197	70	45	543	30.5
9	8	125	96	0	0	0.0
10	4	110	92	0	0	37.6
11	10	168	74	0	0	38.0
12	10	139	80	0	0	27.1
13	1	189	60	23	846	30.1
14	5	166	72	19	175	25.8
15	7	100	0	0	0	30.0
16	0	118	84	47	230	45.8
17	7	107	74	0	0	29.6
18	1	103	30	38	83	43.3
19	1	115	70	30	96	34.6

	DiabetesPedigreeFunction	Age	Outcome
0	0.627	50	1
1	0.351	31	0
2	0.672	32	1
3	0.167	21	0
4	2.288	33	1
5	0.201	30	0
6	0.248	26	1
7	0.134	29	0
8	0.158	53	1
9	0.232	54	1
10	0.191	30	0
11	0.537	34	1
12	1.441	57	0
13	0.398	59	1
14	0.587	51	1
15	0.484	32	1
16	0.551	31	1
17	0.254	31	1
18	0.183	33	0
19	0.529	32	1

```
df.tail(20) # last 20 records
```



	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigree
748	3	187	70	22	200	36.4	
749	6	162	62	0	0	24.3	
750	4	136	70	0	0	31.2	
751	1	121	78	39	74	39.0	
752	3	108	62	24	0	26.0	
753	0	181	88	44	510	43.3	
754	8	154	78	32	0	32.4	
755	1	128	88	39	110	36.5	
756	7	137	90	41	0	32.0	
757	0	123	72	0	0	36.3	
758	1	106	76	0	0	37.5	
759	6	190	92	0	0	35.5	
760	2	88	58	26	16	28.4	
761	9	170	74	31	0	44.0	
762	9	89	62	0	0	22.5	
763	10	101	76	48	180	32.9	
764	2	122	70	27	0	36.8	
765	5	121	72	23	112	26.2	
766	1	126	60	0	0	30.1	
767	1	93	70	31	0	30.4	

```
df = df.rename(columns={'Outcome': 'Diagnosis'}) # change outcome to diagnosis
```

```
# create new column Classification that display Diabetes if outcome is 1, otherwise No Diabetes
```

```
df['Classification'] = df['Diagnosis'].apply(lambda x: 'Diabetes' if x == 1 else 'No Diabetes')
print(df.head(20))
```

```

Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin  BMI  \
0            6      148             72             35      0  33.6
1            1       85             66             29      0  26.6
2            8     183             64              0      0  23.3
3            1       89             66             23     94  28.1
4            0     137             40             35    168  43.1
5            5     116             74              0      0  25.6
6            3       78             50             32     88  31.0
7           10     115              0              0      0  35.3
8            2     197             70             45    543  30.5
9            8     125             96              0      0   0.0
10           4     110             92              0      0  37.6
11          10     168             74              0      0  38.0
12          10     139             80              0      0  27.1
13           1     189             60             23    846  30.1
14           5     166             72             19    175  25.8
15           7     100              0              0      0  30.0
16           0     118             84             47    230  45.8
17           7     107             74              0      0  29.6
18           1     103             30             38     83  43.3
19           1     115             70             30     96  34.6

```

```

DiabetesPedigreeFunction  Age  Diagnosis  Classification
0            0.627    50           1      Diabetes
1            0.351    31           0    No Diabetes
2            0.672    32           1      Diabetes
3            0.167    21           0    No Diabetes
4            2.288    33           1      Diabetes
5            0.201    30           0    No Diabetes
6            0.248    26           1      Diabetes
7            0.134    29           0    No Diabetes
8            0.158    53           1      Diabetes
9            0.232    54           1      Diabetes
10           0.191    30           0    No Diabetes
11           0.537    34           1      Diabetes
12           1.441    57           0    No Diabetes
13           0.398    59           1      Diabetes
14           0.587    51           1      Diabetes
15           0.484    32           1      Diabetes
16           0.551    31           1      Diabetes
17           0.254    31           1      Diabetes
18           0.183    33           0    No Diabetes
19           0.529    32           1      Diabetes

```

```
withDiabetes = df.loc[df['Diagnosis'] == 1] # dataframe withDiabetes
```

```
noDiabetes = df.loc[df['Diagnosis'] == 0] # dataframe noDiabetes
```

```
pedia = df.loc[(df['Age'] >= 0) & (df['Age'] <= 19)] # dataframe Pedia
```

```
adult = df.loc[(df['Age'] > 19)] # dataframe Adult
```

```
# average age and glucose value
```

```
import numpy as np
```

```
average_age = np.mean(df['Age'])
```

```
average_glucose = np.mean(df['Glucose'])
```

```
print(f"Average age:", average_age)
```

```
print(f"Average glucose value:", average_glucose)
```

```

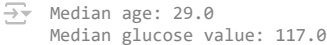
Average age: 33.240885416666664
Average glucose value: 120.89453125

```

```
# median age and glucose value

median_age = np.median(df['Age'])
median_glucose = np.median(df['Glucose'])

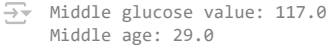
print(f"Median age:", median_age)
print(f"Median glucose value:", median_glucose)
```



```
# middle values of glucose and age

mid_glucose = np.median(df['Glucose'])
mid_age = np.median(df['Age'])

print(f"Middle glucose value:", mid_glucose)
print(f"Middle age:", mid_age)
```




```
# standard deviation of skin thickness

import numpy as np

skinthickness_std = np.std(df['SkinThickness'])

print(f"Standard deviation of skinthickness:", skinthickness_std)
```



6.4 Conclusion

In conclusion, I have applied the intended learning outcomes for the activity. I have use the pandas and numpys for importing and analyzing the given data which is the diabetes file. I got a little problem in exercise 1 and 2 because the statistics value does not match in the statistics table of the random generated data. Furthermore, this activity does connect in my previous course which is MATH 019A - Engineering Data Analysis so it is easy for me in understanding the measures of central tendency but in terms of coding it in Python, it was hard so I did get some guides. Lastly, I thought it was not possible in converting the csv file into dataframe since Google Colab does not have some uploading button, turns out there was a code to upload files and it displays the dataframe of the uploaded file.

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