

Maths DataSet Analysis-3

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1.Runs Test:

```
import csv
import random
import statistics
import math
import scipy.stats as stats
import matplotlib.pyplot as plt
def runs_test(data):
    n = len(data)
    median = sorted(data)[n // 2]
    runs_above_median = 0
    runs_below_median = 0
    current_run = 1
    print("median: ",median)
    for i in range(1, n):
        if data[i] > median:
            if data[i - 1] <= median:
                runs_above_median += 1
            else:
                current_run += 1
        elif data[i] < median:
            if data[i - 1] >= median:
                runs_below_median += 1
            else:
                current_run += 1
    total_runs = runs_above_median + runs_below_median
    print("total_runs : ",total_runs)
    n1=0 #no.of A's
    n2=0 #no.of B's
    for i in range(n):
        if(data[i]>median):
            n1+=1
        elif(data[i]<median):
            n2+=1
```

```

    #use the standard table to find the values for obtained n1,n2--r1,r2 at
    alpha=0.05
    print("no.of A's and B's respectively are: ",n1,n2)

csv_file_path = '../diabetesdataset.csv'
with open(csv_file_path, 'r', newline='') as csvfile:
    csv_reader = csv.reader(csvfile)
    header = next(csv_reader)
    data = list(csv_reader)
    data1 = [float(row[0]) for row in data] # Assuming the values are in the
    first column

sample_size = 35
if sample_size > len(data):
    sample_size= len(data)

random_samples = random.sample(data, sample_size)

```

```

PS C:\Users\nvsku\OneDrive\Desktop\maths-analysis-1\analysis-3> python -u "c:\Users\nvsku\OneDrive\Desktop\maths-analysis-1\analysis-3\RunsTest.py"
median: 5.0
total_runs : 20
no.of A's and B's respectively are: 15 17
sampledata1=[float(row[0])for row in random_samples]

runs_test(sampledata1)

```

output:

Use critical values of r table to find the values r1 and r2.

Here n1=15 and n2=17

From the table;

r1=11

r2=23

And obtained r=total_runs=20

As $r_1 < r < r_2$; accept H_0 : Data is random

2. Sign test:

```
import csv
import random
import scipy.stats as stats
import matplotlib.pyplot as plt
import math
# H0: Median =5.0
# Ha: Median!=5.0

def sign_test(data,sample_size):

    median = 5.0 #assuming median be 5 in H0
    positives = 0
    negatives = 0
    zeroes=0
    for i in range(0, sample_size):
        if data[i] > median:
            positives+=1
        elif data[i] < median:
            negatives+=1
        else:
            zeroes+=1

    n=(sample_size-zeroes)
    print("no.of zeroes : ",zeroes)
    print("no.of positive signs : ",positives)
    print("no.of negative signs : ",negatives)
    print("critical n value(sample_size-no.of zeroes) : ",n)

    z_cal=abs((min(positives,negatives))-(n/2))/math.sqrt(sample_size/2)
    print("z_calculated : ",z_cal)

csv_file_path = '../diabetesdataset.csv'
with open(csv_file_path, 'r', newline='') as csvfile:
    csv_reader = csv.reader(csvfile)
    header = next(csv_reader)
    data = list(csv_reader)
    data1 = [float(row[0]) for row in data] # Assuming the values are in the
first column
```

```

sample_size = 35
if sample_size > len(data):
    sample_size = len(data)

random_samples = random.sample(data, sample_size)
sampledata1=[float(row[0])for row in random_samples]

sign_test(sampledata1,sample_size)

```

output:

```

PS C:\Users\nvsku\OneDrive\Desktop\maths-analysis-1\analysis-3> python -u "c:\Users\nvsku\OneDrive\Desktop\
no.of zeroes : 4
no.of positive signs : 12
no.of negative signs : 19
critical n value(sample_size-no.of zeroes) : 31
z_calculated : 0.836600265340756

```

Conclusion:

- Find z_{tab} value from the standard table and compare with z_{cal} .
- If sampleSize > 26:
 - If $z_{\text{cal}} \leq z_{\text{tab}}$: Accept H_0
 - Else: reject H_0
- If sampleSize ≤ 26:
 - If $z_{\text{cal}} \leq z_{\text{tab}}$: Reject H_0
 - Else: accept H_0

From the table:

$Z_{\text{tab}} = 1.9600$

$Z_{\text{cal}} = 0.8366$

$z_{\text{cal}} < z_{\text{tab}}$ and sample size > 26:

Accept H_0

3. Wilcoxon Rank Sum Test:

```
import pandas as pd
import scipy.stats as stats
import random

print("\nWilcoxon Rank Sum Test\n")
df = pd.read_csv('../diabetesdataset.csv')

# Separate your data into two groups (e.g., group1 and group2)
group1 = df['Glucose(before Lunch)']
group2 = df['Glucose(after Lunch)']

print("Group 1 (Glucose(before Lunch)):")
print(f"H0: Null hypothesis - No significant difference between the groups.")
print(f"Ha: Alternative hypothesis - There is a significant difference between the groups.")

sample_size_group1 = 10 # Choose the sample size for group 1
sample_size_group2 = 12 # Choose the sample size for group 2

print("\tSample size Group 1: " + str(sample_size_group1))
sample_group1 = random.sample(group1.tolist(), sample_size_group1)
print("\tSample size Group 2: " + str(sample_size_group2))
sample_group2 = random.sample(group2.tolist(), sample_size_group2)

# Perform the Wilcoxon Rank Sum Test
statistic, p_value = stats.ranksums(sample_group1, sample_group2)

# Output the results
print("\tWilcoxon Rank Sum Test statistic:", statistic)
print("\tP-value:", p_value)

# Interpret the results
alpha = 0.05 # significance level
if p_value < alpha:
    conclusion = (
        f'''\tSince p-value < alpha\n'''
        f"Reject the null hypothesis: There is a significant difference between the groups.")
else:
    conclusion = (
        f'''\tSince p-value >= alpha\n'''
        f"Fail to reject the null hypothesis: There is no significant difference between the groups.")

print(conclusion)
```

Output:

```
Wilcoxon Rank Sum Test

Group 1 (Glucose(before Lunch)):
H0: Null hypothesis - No significant difference between the groups.
Ha: Alternative hypothesis - There is a significant difference between the groups.
Sample size Group 1: 10
Sample size Group 2: 12
Wilcoxon Rank Sum Test statistic: -0.9890707100936805
P-value: 0.3226285469329776
Since p-value >= alpha
Fail to reject the null hypothesis: There is no significant difference between the groups.
```

4. Wilcoxon Signed Rank Test(sample size > 30):

```
import pandas as pd
import scipy.stats as stats
import random

print("\nWilcoxon Signed-Rank Test\n")

df = pd.read_csv('../diabetesdataset.csv')

# Assuming you have a paired dataset with 'Glucose(before Lunch)' and
# 'Glucose(after Lunch)' columns

print("Paired Data (Glucose(before Lunch) and Glucose(after Lunch)):")
print(f"H0: Null hypothesis - No significant difference between the two paired samples.")
print(f"Ha: Alternative hypothesis - There is a significant difference between the two paired samples.")

sample_size = 34 # Choose the sample size

print("\tSample size: " + str(sample_size))
sample_before = random.sample(df['Glucose(before Lunch)'].tolist(), sample_size)
sample_after = random.sample(df['Glucose(after Lunch)'].tolist(), sample_size)

# Perform the Wilcoxon Signed-Rank Test
statistic, p_value = stats.wilcoxon(sample_before, sample_after)

# Output the results
print("\tWilcoxon Signed-Rank Test statistic:", statistic)
print("\tP-value:", p_value)

# Interpret the results
```

```

alpha = 0.05 # significance level
if p_value < alpha:
    conclusion = (
        f'''\tSince p-value < alpha\n'''
        f"Reject the null hypothesis: There is a significant difference
between the paired samples.")
else:
    conclusion = (
        f'''\tSince p-value >= alpha\n'''
        f"Fail to reject the null hypothesis: There is no significant
difference between the paired samples.")

print(conclusion)

```

output:

```

Wilcoxon Signed-Rank Test

Paired Data (Glucose(before Lunch) and Glucose(after Lunch)):
H0: Null hypothesis - No significant difference between the two paired samples.
Ha: Alternative hypothesis - There is a significant difference between the two paired samples.
Sample size: 34
Wilcoxon Signed-Rank Test statistic: 201.0
P-value: 0.10122521140147
Since p-value >= alpha
Fail to reject the null hypothesis: There is no significant difference between the paired samples.

```

b. Sample size ≤ 30 :

```

import pandas as pd
import scipy.stats as stats
import random

print("\nWilcoxon Signed-Rank Test\n")

df = pd.read_csv('../diabetesdataset.csv')

# Assuming you have a paired dataset with 'Glucose(before Lunch)' and
'Glucose(after Lunch)'

print("Paired Data (Glucose(before Lunch) and Glucose(after Lunch)):")
print(f"H0: Null hypothesis - No significant difference between the two paired
samples.")
print(f"Ha: Alternative hypothesis - There is a significant difference between
the two paired samples.")

sample_size = 25 # Choose the sample size (n < 30)

```

```

print("\tSample size: " + str(sample_size))

# Sample the data for 'Glucose(before Lunch)' and 'Glucose(after Lunch)'
columns
sample_before = random.sample(df['Glucose(before Lunch)'].tolist(),
sample_size)
sample_after = random.sample(df['Glucose(after Lunch)'].tolist(), sample_size)

# Perform the Wilcoxon Signed-Rank Test
statistic, p_value = stats.wilcoxon(sample_before, sample_after)

# Output the results
print("\tWilcoxon Signed-Rank Test statistic:", statistic)
print("\tP-value:", p_value)

# Interpret the results
alpha = 0.05 # significance level
if p_value < alpha:
    conclusion = (
        f'''\tSince p-value < alpha\n'''
        f"Reject the null hypothesis: There is a significant difference
between the paired samples.")
else:
    conclusion = (
        f'''\tSince p-value >= alpha\n'''
        f"Fail to reject the null hypothesis: There is no significant
difference between the paired samples.")
print(conclusion)

```

output:

```

Wilcoxon Signed-Rank Test

Paired Data (Glucose(before Lunch) and Glucose(after Lunch)):
H0: Null hypothesis - No significant difference between the two paired samples.
Ha: Alternative hypothesis - There is a significant difference between the two paired samples.
Sample size: 25
Wilcoxon Signed-Rank Test statistic: 150.5
P-value: 0.7509929537773132
Since p-value >= alpha
Fail to reject the null hypothesis: There is no significant difference between the paired samples.

```