

2021101113-anova-activity

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1 Anova-Activity

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```
[ ]: import pandas as pd
import numpy as np
from scipy.stats import shapiro, levene, f_oneway, chi2
from statsmodels.stats.multicomp import pairwise_tukeyhsd
import matplotlib.pyplot as plt
import seaborn as sns
from tabulate import tabulate
from scipy.stats import f
from scipy.stats import ttest_ind
from scipy.stats import kstest
import pingouin as pg
from statsmodels.stats.multitest import multipletests
```

2 Exam Performance

```
[2]: data = pd.read_csv('exam_scores.csv')
```

Null Hypothesis (H0): Exam performance is not affected by type of schooling

Alternative Hypothesis (H1): Type of schooling affects exam performance

```
[3]: group_stats = {
    'Groups': ['Home', 'Boarding', 'Regular'],
    'Count': [len(data['Home']), len(data['Boarding']), len(data['Regular'])],
    'Sum': [data['Home'].sum(), data['Boarding'].sum(), data['Regular'].sum()],
    'Average': [data['Home'].mean(), data['Boarding'].mean(), data['Regular'].
    ↪mean()],
    'Variance': [data['Home'].var(), data['Boarding'].var(), data['Regular'].
    ↪var()]
}
group_stats_df = pd.DataFrame(group_stats)
print(group_stats_df)
```

	Groups	Count	Sum	Average	Variance
0	Home	15	1182	78.800000	141.171429
1	Boarding	15	1078	71.866667	73.980952
2	Regular	15	1263	84.200000	50.457143

2.1 Check for Normality

```
[4]: shapiro_normality_tests = {}
for col in data.columns:
    stat, p = shapiro(data[col])
    shapiro_normality_tests[col] = {'Shapiro-Wilk Statistic': stat, 'p-value': p,
    ↪ 'Normality': p > 0.05}

lilliefors_normality_tests = {}
for col in data.keys():
    n = len(data[col])
    d, p = kstest(data[col], 'norm', args=(np.mean(data[col]), np.
    ↪ std(data[col], ddof=1)))
    lilliefors_normality_tests[col] = {'Kolmogorov-Smirnov Statistic': d,
    ↪ 'p-value': p, 'Normality': p > 0.05}

print("Shapiro Normality Tests:")
print(pd.DataFrame(shapiro_normality_tests))
print("\n")
print("Kolmogorov-Smirnov Tests with Lilliefors Significance Correction:")
print(pd.DataFrame(lilliefors_normality_tests))
```

Shapiro Normality Tests:

	Home	Boarding	Regular
Shapiro-Wilk Statistic	0.905427	0.969238	0.97489
p-value	0.115225	0.846616	0.922709
Normality	True	True	True

Kolmogorov-Smirnov Tests with Lilliefors Significance Correction:

	Home	Boarding	Regular
Kolmogorov-Smirnov Statistic	0.154949	0.114252	0.099591
p-value	0.812204	0.976888	0.994779
Normality	True	True	True

2.1.1 Check homogeneity of variances

```
[5]: levene_mean = levene(data['Home'], data['Boarding'], data['Regular'],
    ↪ center='mean')
levene_median = levene(data['Home'], data['Boarding'], data['Regular'],
    ↪ center='median')
```

```

levене_trimmed_mean = levene(data['Home'], data['Boarding'], data['Regular'],
    ↪center='trimmed',proportiontocut=0.1)
levене_adjusted_df = levene(data['Home'], data['Boarding'], data['Regular'],
    ↪center='trimmed', proportiontocut=0.05)
print("\nHomogeneity of Variances Test:")
levене_test = pd.DataFrame({
    'Center': ['Mean', 'Median', 'Trimmed Mean','Adjusted df'],
    'Test-Statistic': [levене_mean.statistic, levene_median.statistic,
    ↪levене_trimmed_mean.statistic,levене_adjusted_df.statistic],
    'p-value': [levене_mean.pvalue, levene_median.pvalue, levene_trimmed_mean.
    ↪pvalue,levене_adjusted_df.pvalue]
})

print(levене_test)
if levene_mean.pvalue > 0.05:
    print("Variance is homogenous based on mean")
elif levene_median.pvalue > 0.05:
    print("Variance is homogenous based on median")
elif levene_trimmed_mean.pvalue > 0.05:
    print("Variance is homogenous based on trimmed mean")
elif levene_adjusted_df.pvalue > 0.05:
    print("Variance is homogenous based on adjusted df")
else:
    print("Variance is not homogenous")

if levene_mean.pvalue > 0.05:
    print("\nOne-Way ANOVA Test is chosen")
else:
    print("\nRobust Welch's ANOVA Test is chosen")

```

Homogeneity of Variances Test:

	Center	Test-Statistic	p-value
0	Mean	1.674937	0.199589
1	Median	1.647691	0.204693
2	Trimmed Mean	1.948518	0.157222
3	Adjusted df	1.674937	0.199589

Variance is homogenous based on mean

One-Way ANOVA Test is chosen

2.1.2 Check for sphericity of variances

```
[6]: data = pd.DataFrame(
    {
        'Home': data['Home'],
        'Boarding': data['Boarding'],
        'Regular': data['Regular']
    }
)
mauchly_test = pg.sphericity(data)
print("\nMauchly's Test of Sphericity:")
print(mauchly_test)
statistic_value = mauchly_test[1]
p_value = mauchly_test[4]

if p_value > 0.05:
    print("Sphericity is assumed")
else:
    print("Sphericity is not assumed")
```

Mauchly's Test of Sphericity:

SpherResults(spher=True, W=0.9227681119421941, chi2=1.044905017549741, dof=2, pval=0.5930642676095743)

Sphericity is assumed

3 One-way ANOVA

```
[7]: Home = data['Home']
Boarding = data['Boarding']
Regular = data['Regular']

k = 3
N = len(Home) + len(Boarding) + len(Regular)
group_means = [np.mean(Home), np.mean(Boarding), np.mean(Regular)]
grand_mean = np.mean([np.mean(Home), np.mean(Boarding), np.mean(Regular)])
SSb = sum([len(Home) * (group_means[0] - grand_mean) ** 2,
          len(Boarding) * (group_means[1] - grand_mean) ** 2,
          len(Regular) * (group_means[2] - grand_mean) ** 2])
dfb = k-1
MSb = SSb / dfb
SSw = sum([(x - group_means[i]) ** 2 for i, data in enumerate([Home, Boarding, Regular]) for x in data])
dfw = N-k
MSw = SSw / dfw
F_value = MSb / MSw
alpha = 0.05
```

```

F_crit = f.ppf(1 - alpha, dfb, dfw)
p_value = 1-f.cdf(F_value, dfb, dfw)

anova_table = [
    ["Between Groups", f"{SSb:.6f}", dfb, f"{MSb:.6f}"],
    ["Within Groups", f"{SSw:.6f}", dfw, f"{MSw:.6f}"],
    ["Total", f"{SSb+SSw:.6f}", dfb+dfw]
]

print("ANOVA Table")
print(tabulate(anova_table, headers=["Source of Variation", "SS", "df", "MS"],
    ↪tablefmt="pretty"))

anova_table = [
    ["Between Groups", f"{F_value:.6f}",f"{p_value:.6f}" ,f"{F_crit:.6f}"],
    ["Within Groups"],
    ["Total"]
]
print("\nANOVA Table")
print(tabulate(anova_table, headers=["Source of Variation", "F", "p-value","F_
    ↪crit"], tablefmt="pretty"))

```

ANOVA Table

Source of Variation	SS	df	MS
Between Groups	1146.711111	2	573.355556
Within Groups	3718.533333	42	88.536508
Total	4865.244444	44	

ANOVA Table

Source of Variation	F	p-value	F crit
Between Groups	6.475922	0.003537	3.219942
Within Groups			
Total			

```

[8]: anova_result = f_oneway(data['Home'], data['Boarding'], data['Regular'])
print("\nOne-way ANOVA Test:")
print(f"F-statistic: {anova_result.statistic}")
print(f"p-value: {anova_result.pvalue}")

if anova_result.pvalue < 0.05:

```

```

    print("\nSince p-value < 0.05, there are significant differences between_
    ↪groups. Using a one way ANOVA we observed that the schooling method has a_
    ↪significant effect on exam performance")
    print("Main effect(F) is significant")
else:
    print("\nNo significant differences between groups.")

```

One-way ANOVA Test:

F-statistic: 6.475922406683641

p-value: 0.003536773789503349

Since p-value < 0.05, there are significant differences between groups. Using a one way ANOVA we observed that the schooling method has a significant effect on exam performance

Main effect(F) is significant

4 Effect size calculation

```

[9]: Effect_size = SSb / (SSb + SSw)
    print(f"Effect Size: {Effect_size:.6f}")
    print(f"Type of schooling explains {Effect_size*100:.2f}% of the variance in_
    ↪exam performance")

```

Effect Size: 0.235694

Type of schooling explains 23.57% of the variance in exam performance

We know there is difference between the groups, but which groups perform better or worse?

- Planned comparison (contrast) – prior to experiment (based on the literature)
- Regular schooling > (boarding or home school)
- Regular schooling – Control condition
- Boarding school – Experimental condition 1
- Home school – Experimental condition 2
- But as the no. of planned comparisons increase (>2 comparisons), the alpha level has to be adjusted, again to avoid Type I error. This is done by dividing the alpha level by the no. of comparisons. This is called Bonferroni correction.

4.0.1 Post-hoc Bonferroni for group comparisons

```

[10]: t_statistic_home_boarding, p_value_home_boarding = ttest_ind(Home, Boarding)
    t_statistic_boarding_regular, p_value_boarding_regular = ttest_ind(Boarding,
    ↪Regular)
    t_statistic_regular_home, p_value_regular_home = ttest_ind(Regular, Home)

    alpha = 0.05

```

```

alpha_corrected = alpha / 3

print("alpha corrected: ", alpha_corrected)

p_value_home_boarding_corrected = p_value_home_boarding * 3
p_value_boarding_regular_corrected = p_value_boarding_regular * 3
p_value_regular_home_corrected = p_value_regular_home * 3

table_data = [
    ['Groupwise comparisons', 'T-test p-value', 'Bonferroni-corrected p-value'],
    ['Home vs Boarding', p_value_home_boarding,
    ↪p_value_home_boarding_corrected],
    ['Boarding vs Regular', p_value_boarding_regular,
    ↪p_value_boarding_regular_corrected],
    ['Regular vs Home', p_value_regular_home, p_value_regular_home_corrected]
]
print(tabulate(table_data, headers="firstrow", tablefmt="grid"))

```

alpha corrected: 0.016666666666666666

Groupwise comparisons	T-test p-value	Bonferroni-corrected p-value
Home vs Boarding	0.07781	0.23343
Boarding vs Regular	0.00019644	0.000589321
Regular vs Home	0.142042	0.426125

4.1 Holm method for multiple comparisons

```

[11]: datasets = [('Home', data['Home']), ('Boarding', data['Boarding']), ('Regular',
    ↪data['Regular'])]
alpha = 0.05
p_values = []
for i in range(len(datasets)):
    for j in range(i + 1, len(datasets)):
        group1_name, group1_data = datasets[i]
        group2_name, group2_data = datasets[j]
        t_stat, p_value = ttest_ind(group1_data, group2_data)
        p_values.append(p_value)

table = []
table.append(['Group 1', 'Group 2', 'Significant Difference', 'p-corrected'])
reject, p_values_corrected, _, _ = multipletests(p_values, alpha=alpha,
    ↪method='holm')
index = 0

```

```

for i in range(len(datasets)):
    for j in range(i + 1, len(datasets)):
        group1_name, _ = datasets[i]
        group2_name, _ = datasets[j]
        if reject[index]:
            print(f"There is a significant difference between {group1_name} and_
↪{group2_name} (p-corrected = {p_values_corrected[index]})\n")
            table.append([group1_name, group2_name, "Yes",_
↪f"{p_values_corrected[index]}"])
        else:
            print(f"No significant difference between {group1_name} and_
↪{group2_name} (p-corrected = {p_values_corrected[index]})\n")
            table.append([group1_name, group2_name, "No",_
↪f"{p_values_corrected[index]}"])
            index += 1
print(tabulate(table, headers='firstrow', tablefmt='grid'))

```

No significant difference between Home and Boarding (p-corrected = 0.1556199753902239)

No significant difference between Home and Regular (p-corrected = 0.1556199753902239)

There is a significant difference between Boarding and Regular (p-corrected = 0.0005893211647290649)

Group 1	Group 2	Significant Difference	p-corrected
Home	Boarding	No	0.15562
Home	Regular	No	0.15562
Boarding	Regular	Yes	0.000589321

4.1.1 Tukey's HSD post-hoc test

```

[12]: if anova_result.pvalue < 0.05:
        data_melted = pd.melt(data)
        posthoc = pairwise_tukeyhsd(data_melted['value'], data_melted['variable'],_
↪alpha=0.05)
        print(posthoc)
        print(posthoc.q_crit)
        HSD = posthoc.q_crit*np.sqrt(MSw / len(data))
        print(f"HSD: {HSD:.6f}")

```



```
print(f"The mean difference between any two samples must be more than {HSD:.
↪6f} at alpha = 0.05 for the difference to be statistically significant")
```

Multiple Comparison of Means - Tukey HSD, FWER=0.05

```
=====
group1  group2 meandiff p-adj  lower  upper  reject
-----
Boarding   Home    6.9333 0.1204  -1.414 15.2806  False
Boarding  Regular  12.3333 0.0024   3.986 20.6806   True
   Home Regular    5.4  0.269 -2.9473 13.7473  False
-----
```

3.4358230206770175

HSD: 8.347306

The mean difference between any two samples must be more than 8.347306 at alpha = 0.05 for the difference to be statistically significant

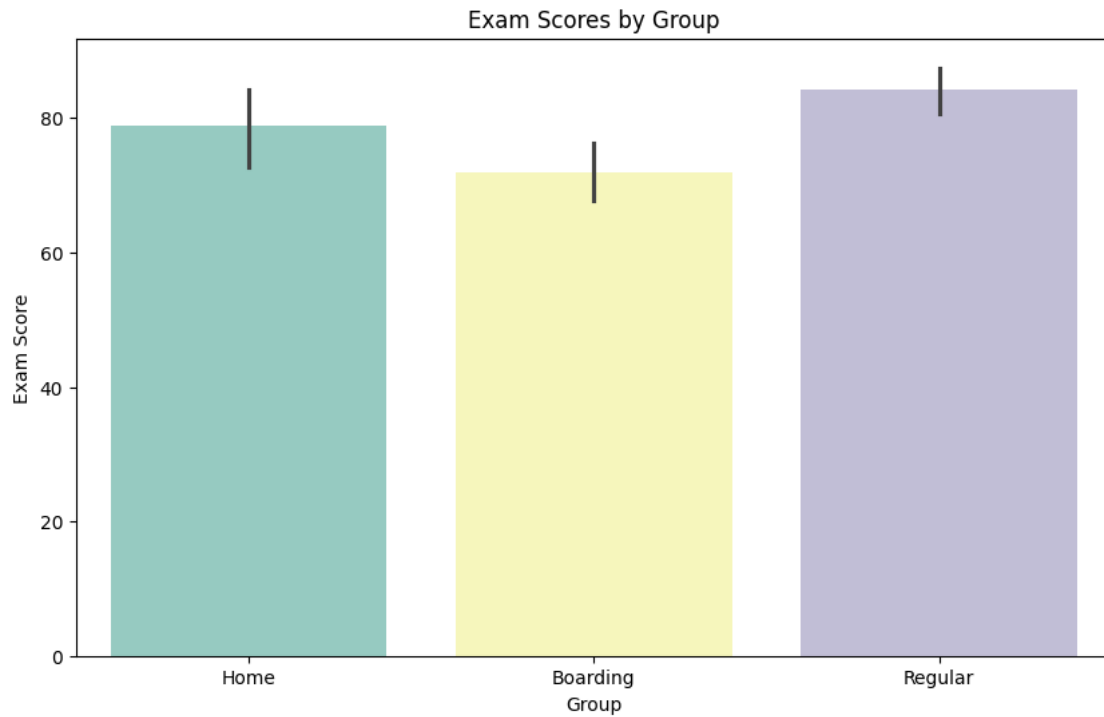
Using Bonferroni post-hoc test, we found that regular school resulted in better exam performance than boarding school ($p < .001$). There was no significant difference between the other groups

Using a one way ANOVA we observed that the schooling method has a significant effect on exam performance

Using Bonferroni post-hoc test, we found that regular school resulted in better exam performance than boarding school ($p < .001$). There was no significant difference between the other groups.

5 Plot Analyzed Data

```
[13]: data_melted = pd.melt(data)
plt.figure(figsize=(10, 6))
sns.barplot(x='variable', y='value', data=data_melted, errorbar=('ci', 95),
↪hue='variable', palette="Set3")
plt.title('Exam Scores by Group')
plt.xlabel('Group')
plt.ylabel('Exam Score')
plt.show()
print("Error bars denote confidence intervals (CI) of 95%")
```



Error bars denote confidence intervals (CI) of 95%

```
[14]: plt.figure(figsize=(10, 6))
sns.boxplot(x='variable', y='value', hue='variable', data=data.melt(),
            palette="Set3", legend=False)
plt.title('Exam Scores by Group')
plt.xlabel('Group')
plt.ylabel('Exam Score')
plt.show()
```

