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import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from scipy import stats
from sklearn.datasets import load_iris

iris = load_iris()
df = pd.DataFrame(data=iris.data, columns=iris.feature_names)
df['species'] = pd.Categorical.from_codes(iris.target,
iris.target_names)

print("CORRELATION MATRIX ANALYSIS")
print("-----")

correlation_matrix = df.iloc[:, 0:4].corr()
print("Correlation Matrix:")
print(correlation_matrix)
print("\n")

print("COVARIANCE ANALYSIS")
print("-----")

covariance_matrix = df.iloc[:, 0:4].cov()
print("Covariance Matrix:")
print(covariance_matrix)
print("\n")

print("ANOVA ANALYSIS")
print("-----")
features = iris.feature_names
print("One-way ANOVA results for each feature across different
species:")
for feature in features:
    groups = [df[df['species'] == species][feature] for species in
iris.target_names]
    f_val, p_val = stats.f_oneway(*groups)

    print(f"\nFeature: {feature}")
    print(f"F-value: {f_val:.4f}")
    print(f"p-value: {p_val:.8f}")

    if p_val < 0.05:
        print("Result: There are significant differences between species for
this feature")
    else:
        print("Result: No significant differences between species for this
feature")

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plt.figure(figsize=(10, 8))
sns.heatmap(correlation_matrix, annot=True, cmap='coolwarm', vmin=-1,
vmax=1)
plt.title('Correlation Matrix Heatmap of Iris Features')
plt.tight_layout()
plt.savefig('correlation_heatmap.png')

plt.figure(figsize=(10, 8))
sns.heatmap(covariance_matrix, annot=True, cmap='viridis')
plt.title('Covariance Matrix Heatmap of Iris Features')
plt.tight_layout()
plt.savefig('covariance_heatmap.png')

plt.figure(figsize=(12, 10))
for i, feature in enumerate(features):
plt.subplot(2, 2, i+1)
sns.boxplot(x='species', y=feature, data=df)
plt.title(f'Distribution of {feature} by Species')
plt.tight_layout()
plt.savefig('anova_boxplots.png')

plt.figure(figsize=(12, 10))
sns.pairplot(df, hue='species')
plt.tight_layout()
plt.savefig('feature_relationships.png')

print("\nAnalysis complete. All visualizations have been saved.")
```

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ishadpande@Argos:~/Documents/dev/practicals/PS$ /usr/bin/python /home/ishadpande/Documents/dev/practicals/PS/4.py
CORRELATION MATRIX ANALYSIS
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Correlation Matrix:
      sepal length (cm)  sepal width (cm)  petal length (cm)  petal width (cm)
sepal length (cm)      1.000000      -0.117570      0.871754      0.817941
sepal width (cm)      -0.117570      1.000000      -0.428440      -0.366126
petal length (cm)      0.871754      -0.428440      1.000000      0.962865
petal width (cm)      0.817941      -0.366126      0.962865      1.000000

COVARIANCE ANALYSIS
-----
Covariance Matrix:
      sepal length (cm)  sepal width (cm)  petal length (cm)  petal width (cm)
sepal length (cm)      0.685694      -0.042434      1.274315      0.516271
sepal width (cm)      -0.042434      0.189979      -0.329656      -0.121639
petal length (cm)      1.274315      -0.329656      3.116278      1.295609
petal width (cm)      0.516271      -0.121639      1.295609      0.581006

```

ANOVA ANALYSIS

One-way ANOVA results for each feature across different species:

Feature: sepal length (cm)

F-value: 119.2645

p-value: 0.00000000

Result: There are significant differences between species for this feature

Feature: sepal width (cm)

F-value: 49.1600

p-value: 0.00000000

Result: There are significant differences between species for this feature

Feature: petal length (cm)

F-value: 1180.1612

p-value: 0.00000000

Result: There are significant differences between species for this feature

Feature: petal width (cm)

F-value: 960.0071

p-value: 0.00000000

Result: There are significant differences between species for this feature

Analysis complete. All visualizations have been saved.





