

```
# IRIS FLOWER CLASSIFICATION PROJECT
## PART 3 – Exploratory Data Analysis and Hypothesis Testing
```

```
!pip install seaborn scipy
```

```
Requirement already satisfied: seaborn in /usr/local/lib/python3.12/dist-packages (0.13.2)
Requirement already satisfied: scipy in /usr/local/lib/python3.12/dist-packages (1.16.3)
Requirement already satisfied: numpy!=1.24.0,>=1.20 in /usr/local/lib/python3.12/dist-packages (from seaborn) (2.0.2)
Requirement already satisfied: pandas>=1.2 in /usr/local/lib/python3.12/dist-packages (from seaborn) (2.2.2)
Requirement already satisfied: matplotlib!=3.6.1,>=3.4 in /usr/local/lib/python3.12/dist-packages (from seaborn) (3.10.0)
Requirement already satisfied: contourpy>=1.0.1 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: fonttools>=4.22.0 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: kiwisolver>=1.3.1 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: packaging>=20.0 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: pillow>=8 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn) (11.1.0)
Requirement already satisfied: pyparsing>=2.3.1 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: python-dateutil>=2.7 in /usr/local/lib/python3.12/dist-packages (from matplotlib!=3.6.1,>=3.4->seaborn)
Requirement already satisfied: pytz>=2020.1 in /usr/local/lib/python3.12/dist-packages (from pandas>=1.2->seaborn) (2025.2)
Requirement already satisfied: tzdata>=2022.7 in /usr/local/lib/python3.12/dist-packages (from pandas>=1.2->seaborn) (2025.3)
Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.12/dist-packages (from python-dateutil>=2.7->matplotlib!=3.6.1)
```

```
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt

from sklearn.datasets import load_iris
```

```
iris = load_iris()

df = pd.DataFrame(
    iris.data,
    columns=iris.feature_names
)

df["species"] = iris.target

df["species"] = df["species"].map({
    0: "setosa",
    1: "versicolor",
    2: "virginica"
})

df.head()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	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	

Next steps: [Generate code with df](#) [New interactive sheet](#)

The Iris dataset is loaded using the machine learning library :contentReference[oaicite:0]{index=0}. The dataset contains four measurement features and three flower species categories.

```
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
 #   Column           Non-Null Count  Dtype  
--- 
 0   sepal length (cm)  150 non-null   float64
 1   sepal width (cm)  150 non-null   float64
 2   petal length (cm) 150 non-null   float64
 3   petal width (cm)  150 non-null   float64
```

```
4 species      150 non-null    object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

```
df.describe()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	grid icon
count	150.000000	150.000000	150.000000	150.000000	grid icon
mean	5.843333	3.057333	3.758000	1.199333	grid icon
std	0.828066	0.435866	1.765298	0.762238	grid icon
min	4.300000	2.000000	1.000000	0.100000	grid icon
25%	5.100000	2.800000	1.600000	0.300000	grid icon
50%	5.800000	3.000000	4.350000	1.300000	grid icon
75%	6.400000	3.300000	5.100000	1.800000	grid icon
max	7.900000	4.400000	6.900000	2.500000	grid icon

```
df.isnull().sum()
```

	0
sepal length (cm)	0
sepal width (cm)	0
petal length (cm)	0
petal width (cm)	0
species	0

```
dtype: int64
```

The dataset contains 150 records with no missing values. All features are numerical measurements suitable for analysis.

```
df.groupby("species").mean()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	grid icon
species					grid icon
setosa	5.006	3.428	1.462	0.246	grid icon
versicolor	5.936	2.770	4.260	1.326	grid icon
virginica	6.588	2.974	5.552	2.026	grid icon

```
df.groupby("species").median()
```

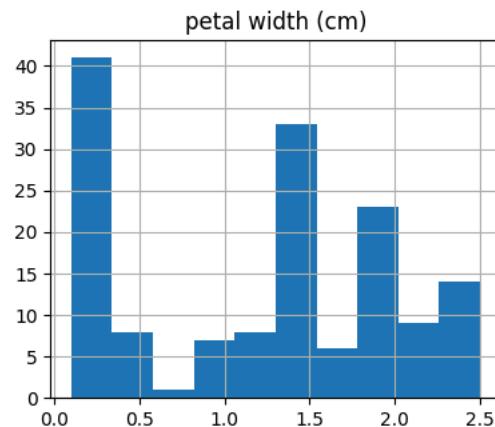
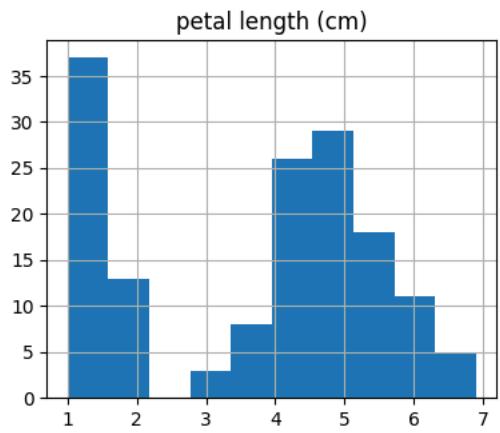
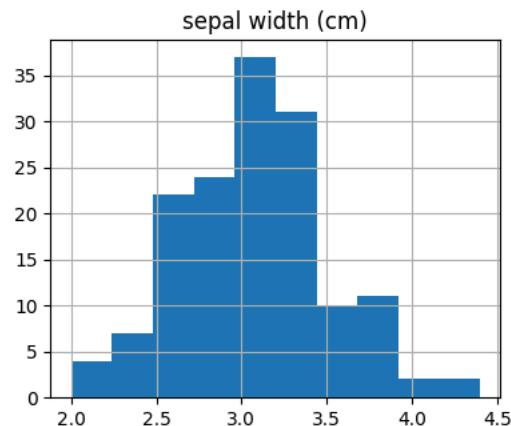
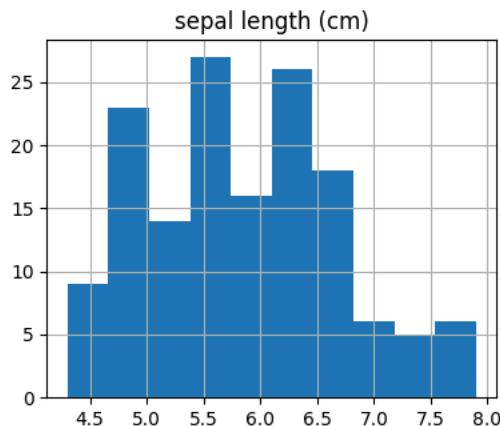
	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	grid icon
species					grid icon
setosa	5.0	3.4	1.50	0.2	grid icon
versicolor	5.9	2.8	4.35	1.3	grid icon
virginica	6.5	3.0	5.55	2.0	grid icon

```
df.groupby("species").std()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	grid icon
species					grid icon
setosa	0.352490	0.379064	0.173664	0.105386	grid icon
versicolor	0.516171	0.313798	0.469911	0.197753	grid icon
virginica	0.635880	0.322497	0.551895	0.274650	grid icon

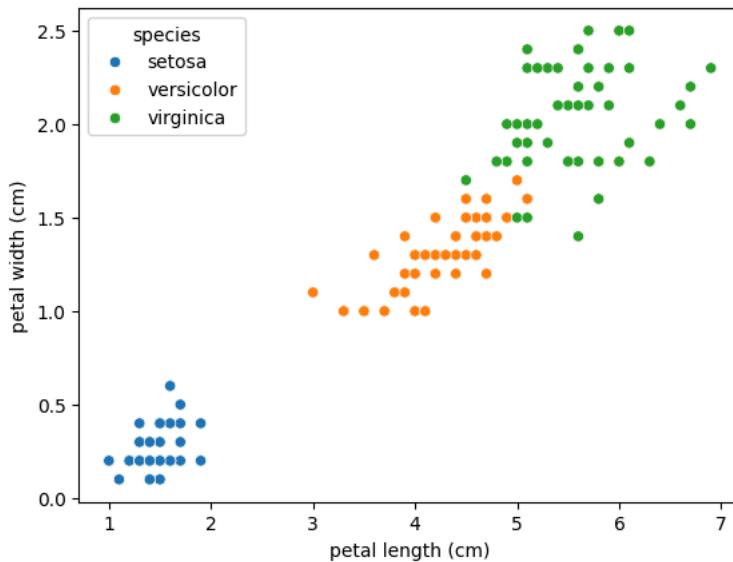
Petal measurements show major differences among species compared to sepal measurements.

```
df.hist(figsize=(10,8))
plt.show()
```



Histograms show distribution of flower measurements. Setosa species has smaller petal sizes compared to others.

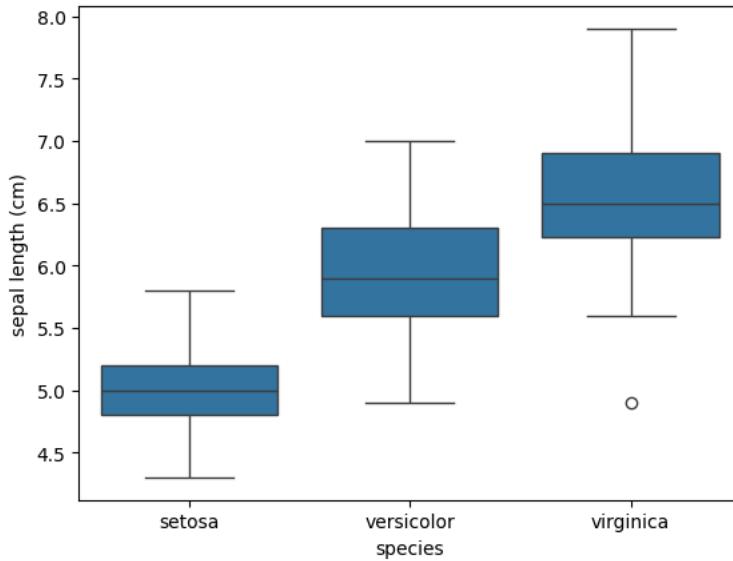
```
sns.scatterplot(
    data=df,
    x="petal length (cm)",
    y="petal width (cm)",
    hue="species"
)
plt.show()
```



Petal length and petal width provide clear separation among Iris species.

```
sns.boxplot(
x="species",
y="sepal length (cm)",
data=df
)

plt.show()
```



Box plots show variation and median differences among species.

```
from scipy.stats import ttest_ind

setosa=df[df["species"]=="setosa"]

versicolor=df[df["species"]=="versicolor"]

ttest_ind(
setosa["sepal length (cm)"],
versicolor["sepal length (cm)"]
)

TtestResult(statistic=np.float64(-10.52098626754911), pvalue=np.float64(8.985235037487079e-18), df=np.float64(98.0))
```

Since the p-value is less than 0.05, the null hypothesis is rejected. Sepal length differs significantly between Setosa and Versicolor species.

```
from scipy.stats import f_oneway

virginica=df[df["species"]=="virginica"]

f_oneway(
    setosa["petal width (cm)"],
    versicolor["petal width (cm)"],
    virginica["petal width (cm)"]
)

F_onewayResult(statistic=np.float64(960.0071468018067), pvalue=np.float64(4.169445839443833e-85))
```

The ANOVA test shows significant differences in petal width among the three species.

Key Insights:

1. Petal measurements strongly distinguish Iris species.
2. Setosa species forms a separate cluster.
3. Versicolor and Virginica partially overlap.
4. Statistical tests confirm significant measurement differences.