▼ EXPLORATORY DATA ANALYSIS

#IMPORT ALL NECCESSARY MODULES

import pandas as pd

import numpy as np

import matplotlib.pyplot as plt

import seaborn as sns

#import data set

df=sns.load_dataset("iris")

df

₽		sepal_length	sepal_width	petal_length	petal_width	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
	145	6.7	3.0	5.2	2.3	virginica
	146	6.3	2.5	5.0	1.9	virginica
	147	6.5	3.0	5.2	2.0	virginica
	148	6.2	3.4	5.4	2.3	virginica
	149	5.9	3.0	5.1	1.8	virginica

→ SUMMARY OF DATA

150 rows × 5 columns

df.info()

<class 'pandas.core.frame.DataFrame'>

→ statistical summary

df.describe()

	sepal_length	sepal_width	petal_length	petal_width
count	150.000000	150.000000	150.000000	150.000000
mean	5.843333	3.057333	3.758000	1.199333
std	0.828066	0.435866	1.765298	0.762238
min	4.300000	2.000000	1.000000	0.100000
25%	5.100000	2.800000	1.600000	0.300000
50%	5.800000	3.000000	4.350000	1.300000
75%	6.400000	3.300000	5.100000	1.800000
max	7.900000	4.400000	6.900000	2.500000

check for null values

df.isnull().sum()

sepal_length 0
sepal_width 0
petal_length 0
petal_width 0
species 0
dtype: int64

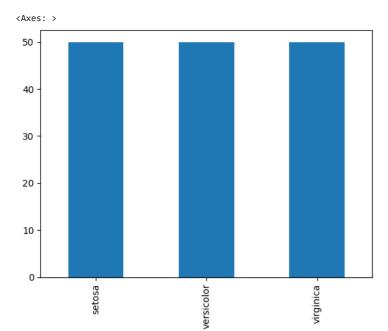
#covariance of data
df.cov()

	sepal_length	sepal_width	petal_length	petal_width
sepal_length	0.685694	-0.042434	1.274315	0.516271
sepal_width	-0.042434	0.189979	-0.329656	-0.121639
petal_length	1.274315	-0.329656	3.116278	1.295609
petal_width	0.516271	-0.121639	1.295609	0.581006

#visualisation
plt.figure(figsize=(10,6))
sns.heatmap(df.cov(),annot=True)



df.species.value_counts().plot(kind="bar")



#check for duplicates df.duplicated()

> False False False 0 1 2 3 4 False ... False 145 146 False 147 148 False False 149 False Length: 150, dtype: bool

df.shape

(150, 5)

df[df.duplicated()]

	sepal_length	sepal_width	petal_length	petal_width	species
142	5.8	2.7	5.1	1.9	virginica

#drop the duplicates df.drop_duplicates(inplace=True)

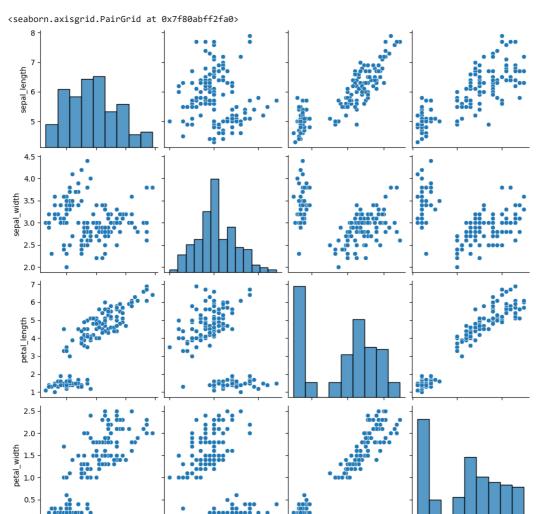
df

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149 rows × 5 columns

0.0

sepal_length



petal_length

petal_width

sns.scatterplot(x=df.petal_length,y=df.petal_width,hue=df["species"])

