!wget https://raw.githubusercontent.com/uiuc-cse/data-fa14/gh-pages/data/iris.csv

#importing packages

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

iris = pd.read_csv('iris.csv') #reasding the csv file

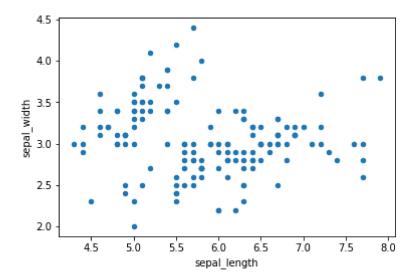
iris #printing the csv file

species	petal_width	petal_length	sepal_width	sepal_length	
setosa	0.2	1.4	3.5	5.1	0
setosa	0.2	1.4	3.0	4.9	1
setosa	0.2	1.3	3.2	4.7	2
setosa	0.2	1.5	3.1	4.6	3
setosa	0.2	1.4	3.6	5.0	4
virginica	2.3	5.2	3.0	6.7	145
virginica	1.9	5.0	2.5	6.3	146
virginica	2.0	5.2	3.0	6.5	147
virginica	2.3	5.4	3.4	6.2	148
virginica	1.8	5.1	3.0	5.9	149

150 rows × 5 columns

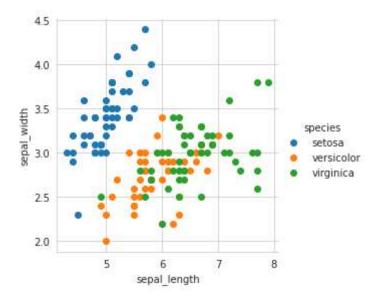
iris.plot(kind='scatter', x= 'sepal_length', y='sepal_width') #plotting
https://colab.research.google.com/drive/1UNeafpDvFNCYDhU1f6xf4LozluP5QAQD#scrollTo=bx00WHuN6Wo-&printMode=true

plt.show()



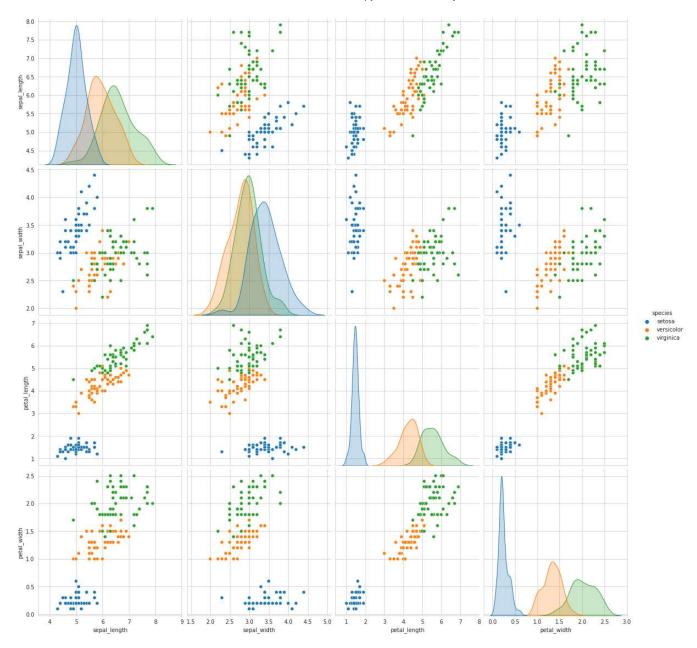
sns.set_style('whitegrid')

sns.FacetGrid(iris, hue='species',height =4).map(plt.scatter, 'sepal_length','sepal_width').a
plt.show()



sns.pairplot(iris, hue='species', height=4) #plotting
plt.show()

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