LESSON 14 LAB

Basic datasets management, Coding in Colab and Matlab

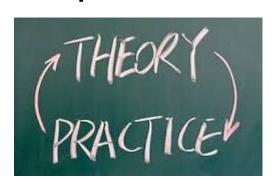


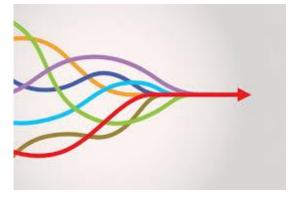
Laboratory outline

- LABORATORY (in a separate file): Loading and basic dataset management
 - Colab example
 - Matalab example
- The laboratory is based on the IRIS dataset
- Please refer to the Lesson #14 for the description of the IRIS dataset

What in the exam?

- In the exam there will be no (exact) coding
- Nevertheless, is it important to study the application of the concept of the theory in practical cases and examples in order to better understand the notions





Knowledge

You will be asked to comment some example of code

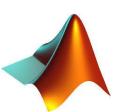
LAB CLASS

COLAB



- Environment preparation
- Data loading

MATLAB



- Load the dataset
- Create the data structures (X and Y)

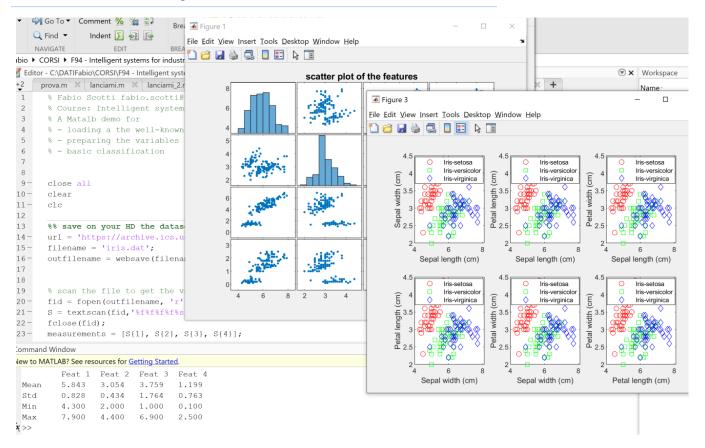
- Preliminary data analysis
- Plot the data
- Normalization
- Classification example

Best thing to do... (Matlab)

Download the example and let them run!



laboratory_matlab_IRIS_Dataset.m



The example is commented... explore the code and try to modify it and/or add new plots

Best thing to do... (Colab)

- Download the code
- laboratory_COLAB_IRIS_Dataset.ipynb
- → upload it to your Gdrive

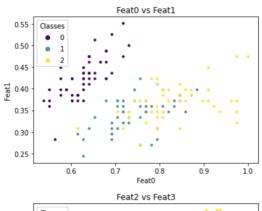
https://colab.research.google.com/

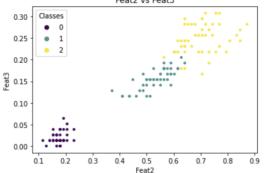
> open it with Colab and let it run

(use Google Chrome – with login)

Plot features and classes

```
[41] import matplotlib.pyplot as plt
     # select 2 features to plot
     feat0 = xNumpy_Norm[:,0]
     feat1 = xNumpy_Norm[:,1]
     # params
     area = np.pi*3
     fig1, ax1 = plt.subplots()
     scatter1 = ax1.scatter(feat0, feat1, s=area, c=yNumpy)
     plt.title('Feat0 vs Feat1')
     plt.xlabel('Feat0')
     plt.ylabel('Feat1')
     legend1 = ax1.legend(*scatter1.legend_elements(),
                         loc="upper left", title="Classes")
     ax1.add_artist(legend1)
     plt.show()
     # select 2 features to plot
     feat2 = xNumpy_Norm[:,2]
     feat3 = xNumpy Norm[:,3]
     # params
     area = np.pi*3
     # Plot
     fig2, ax2 = plt.subplots()
     scatton? - av2 scatton/foat2 foat2 s-anoa s-vNumny)
```



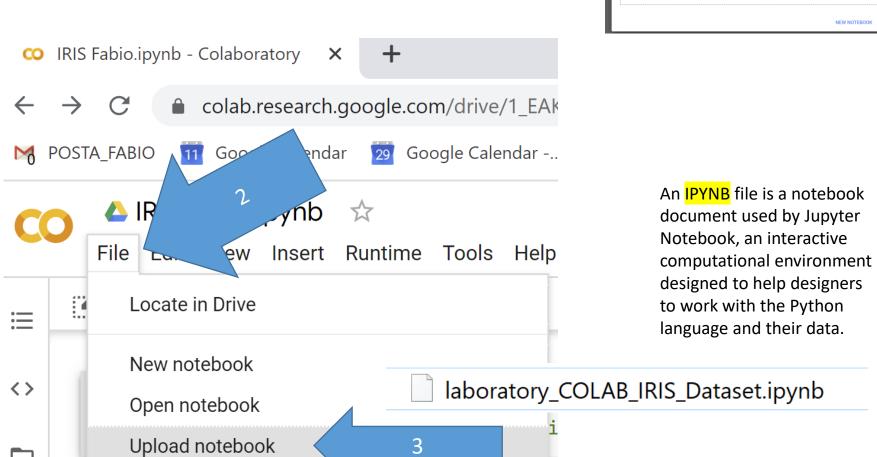


The examples are commented

Explore them and try to modify them or to add new plots

Upload your *.ipynb

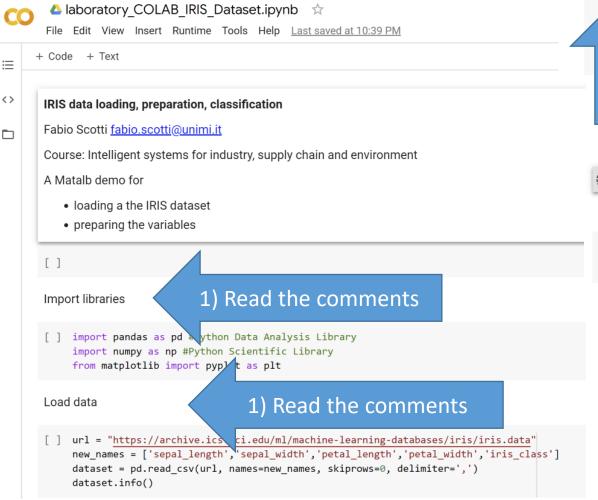
https://colab.research.google.com/





Import libraries

Let's it run



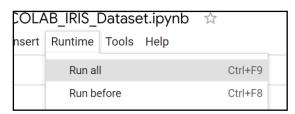
[] import pandas import numpy a from matplotli



Import libraries

[] import pandas as pd #Python Data Analysis Library import numpy as np #Python Scientific Library from matplotlib import pyplot as plt

Clik on the [] to run the cell



Or «Run all»



Toolboxes Loading, EDA/plotting in Python/Colab

And let's have a look about Matplotlib







Environment preparation

Panda: libraries for dataset preprocessing

Numpy: libraries for array manipulation

Import libraries

[74] import pandas as pd #Python Data Analysis Library
import numpy as np #Python Scientific Library

NumPy (pronounced /ˈnʌmpaɪ/ (NUM-py) or sometimes /ˈnʌmpi/ (NUM-pee)) is a library for the Python programming language, adding support for large, multi-dimensional arrays and matrices, along with a large collection of high-level mathematical functions to operate on these arrays.

Data loading

Iris dataset

Use the panda library to read the CSV

Load data

```
[75] url = "https://archive.ics.uci.edu/ml/machine-learning-databases/iris/iris.data"
   new_names = ['sepal_length','sepal_width','petal_length','petal_width','iris_class']
   dataset = pd.read_csv(url, names=new_names, skiprows=0, delimiter=',')
   dataset.info()
```

Data loading

dataset.info() gives us organized information

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 150 entries, 0 to 149
Data columns (total 5 columns):
    Column Non-Null Count
                               Dtype
0 sepal_length 150 non-null float64
1 sepal_width 150 non-null float64
2 petal_length 150 non-null float64
3 petal_width 150 non-null float64
    iris_class 150 non-null
                               object
dtypes: float64(4), object(1)
memory usage: 6.0+ KB
```

Data loading

dataset.head(n) to view the first n samples

		sepal_length	sepal_width	petal_length	petal_width	iris_class
	0	5.1	3.5	1.4	0.2	Iris-setosa
	1	4.9	3.0	1.4	0.2	Iris-setosa
	2	4.7	3.2	1.3	0.2	Iris-setosa
	3	4.6	3.1	1.5	0.2	Iris-setosa
	4	5.0	3.6	1.4	0.2	Iris-setosa
	5	5.4	3.9	1.7	0.4	Iris-setosa
	6	4.6	3.4	1.4	0.3	Iris-setosa
	7	5.0	3.4	1.5	0.2	Iris-setosa
	8	4.4	2.9	1.4	0.2	Iris-setosa
	9	4.9	3.1	1.5	0.1	Iris-setosa

First, we need to separate the features from the classes

• Different preprocessing for features and classes (Necessary? YES! To avoid Data Leakage!!!)

```
[77] y = dataset['iris_class']
x = dataset.drop(['iris_class'], axis=1)

print ("dataset : ",dataset.shape)
print ("x : ",x.shape)

print ("y : ",y.shape)
```

First, we need to separate the features from the classes

- Feature matrix has 4 columns (the features)
- Target vector has 1 column (the class)

```
C→ dataset : (150, 5)
x : (150, 4)
y : (150,)
```

Use the numpy library to convert data into matrix format

Ease of access to portions of the matrices

```
[78] xNumpy = x.to_numpy()
yNumpy = y.to_numpy()
```

We can access a single feature or a single samples

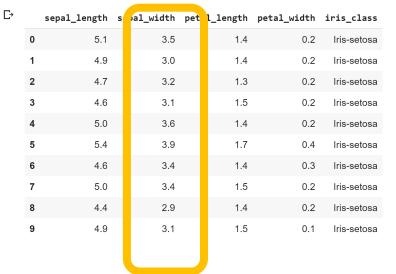
Read a specific feature: matrix slicing by column

```
[79] feature_to_extract = 1
  feature = xNumpy[:,feature_to_extract]
  print(feature)
```

We can access a single feature or a single samples

Read a specific feature: matrix slicing by column

```
[79] feature_to_extract = 1
    feature = xNumpy[:,feature_to_extract]
    print(feature)
```



We can access a single feature or a single samples

Read a specific feature: matrix slicing by column

```
feature to extract = 1
 feature = xNumpy[:,feature to extract]
                                                                       Debugging!!
 print(feature)
                                      3.8 3.2 3.7 3.3 3.2 3.2 3.1 2.3
2.8 2.8 3.3 2.4 2.9 2.7 2.
                              2.2 2.9 2.9 3.1 3.
                          3.
               2.8 3.
                      2.9 2.6 2.4 2.4 2.7 2.7 3.
                                                  3.4 3.1 2.3 3.
                      2.9 2.9 2.5 2.8 3.3 2.7 3.
                  2.5 2.8 3.2 3.
                                 3.8 2.6 2.2 3.2 2.8 2.8 2.7 3.3 3.2
              2.8 3.8 2.8 2.8 2.6 3. 3.4 3.1 3. 3.1 3.1 3.1 2.7 3.2
               3.4 3. ]
       2.5 3.
```



·						
•		sepal_length	sepal_width	petal_length	petal_width	iris_class
	0	5.1	3.5	1.4	0.2	Iris-setosa
	1	4.9	3.0	1.4	0.2	Iris-setosa
	2	4.7	3.2	1.3	0.2	Iris-setosa
	3	4.6	3.1	1.5	0.2	Iris-setosa
	4	5.0	3.6	1.4	0.2	Iris-setosa
	5	5.4	3.9	1.7	0.4	Iris-setosa
	6	4.6	3.4	1.4	0.3	Iris-setosa
	7	5.0	3.4	1.5	0.2	Iris-setosa
	8	4.4	2.9	1.4	0.2	Iris-setosa
	9	4.9	3.1	1.5	0.1	Iris-setosa

We can access a single feature or a single samples

Read a specific sample: matrix slicing by row

```
[80] sample_to_extract = 1
sample = xNumpy[sample_to_extract,:]
print(sample)
```

We can access a single feature or a single samples

Read a specific sample: matrix slicing by row

```
[80] sample_to_extract = 1
    sample = xNumpy[sample_to_extract,:]
    print(sample)
```

Debugging!!



₽		sepal_length	sepal_width	petal_length	petal_width	iris_class
	0	5.1	3.5	1.4	0.2	Iris-setosa
	1	4.9	3.0	1.4	0.2	Iris-setosa
	2	4.7	3.2	1.3	0.2	Iris-setosa
	3	4.6	3.1	1.5	0.2	Iris-setosa
	4	5.0	3.6	1.4	0.2	Iris-setosa
	5	5.4	3.9	1.7	0.4	Iris-setosa
	6	4.6	3.4	1.4	0.3	Iris-setosa
	7	5.0	3.4	1.5	0.2	Iris-setosa
	8	4.4	2.9	1.4	0.2	Iris-setosa
	9	4.9	3.1	1.5	0.1	Iris-setosa

 sepal_length
 sepal_width
 petal_length
 petal_width
 iris_class

 0
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 3.5
 1.4
 0.2
 Iris-setosa

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 4.9
 3.0
 1.4
 0.2
 Iris-setosa

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We can access a single feature or a single samples

Read a specific sample: matrix slicing by row

С→

```
[80] sample_to_extract = 1
    sample = xNumpy[sample_to_extract,:]
    print(sample)
```

$$\Box$$
 [4.9 3. 1.4 0.2]

Data normalization: ensure that the data is in the range [0-1]

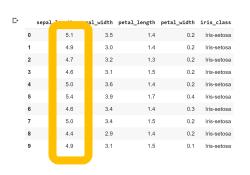
 Min-max normalization: (x-min(x)) / (max(x)-min(x))

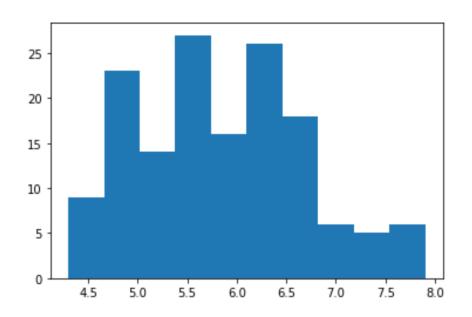
```
[81] xNumpy_Norm = (xNumpy - np.min(xNumpy)) / (np.max(xNumpy) - np.min(xNumpy))
```

GEN TO SERVICE OF THE SERVICE OF THE

Data normalization: ensure that the data is in the range [0-1]

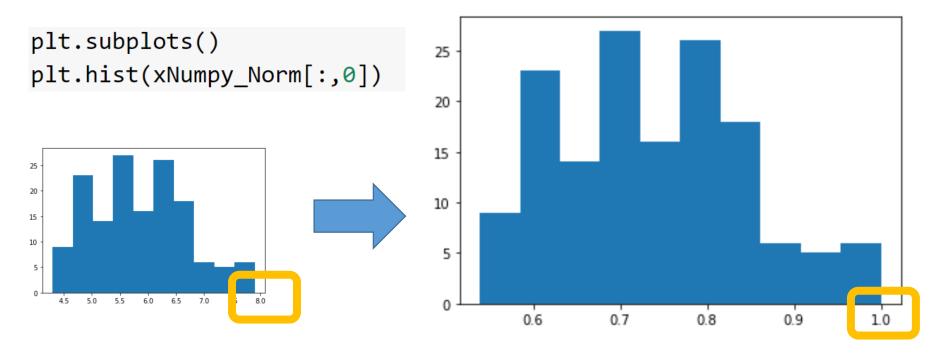
Non-normalized data





Data normalization: ensure that the data is in the range [0-1]

Normalized data



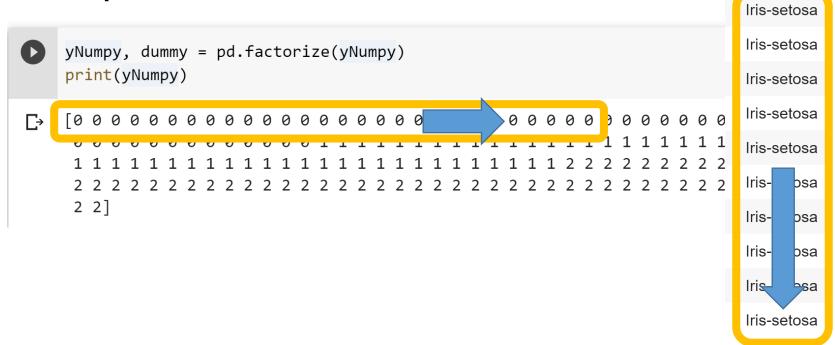
Data conversion: from categorical to decimal

Class data is in categorical format



Data conversion: from categorical to decimal

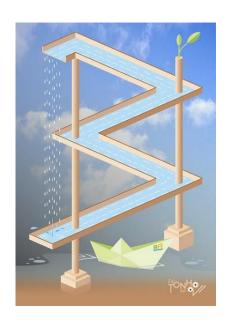
Conversion to decimal for easier manipulation



iris_class

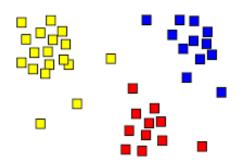
Visualizing the features and the classes allows us to have a first visual check on the discriminatory capability

 Drawback: visualization is limited to 2 or 3 dimensions



Visualizing the features and the classes allows us to have a first visual check on the discriminatory capability

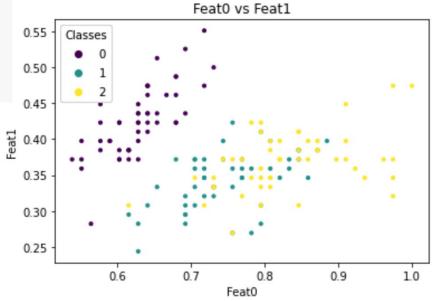
- 1. Select features to plot
- 2. Assign color based on the class



```
import matplotlib.pyplot as plt
# select 2 features to plot
feat0 = xNumpy_Norm[:,0]
feat1 = xNumpy_Norm[:,1]
# params
area = np.pi*3
# Plot
fig1, ax1 = plt.subplots()
scatter1 = ax1.scatter(feat0, feat1, s=area, c=yNumpy)
plt.title('Feat0 vs Feat1')
plt.xlabel('Feat0')
plt.ylabel('Feat1')
legend1 = ax1.legend(*scatter1.legend elements(),
                    loc="upper left", title="Classes")
ax1.add artist(legend1)
plt.show()
```

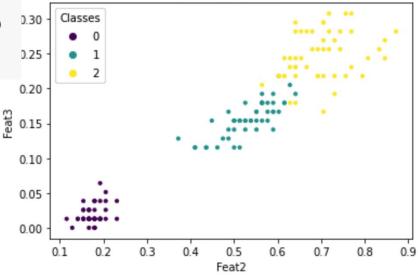
Feat0 + Feat1:

- Good discrimination for class 0
- Bad discrimination for class 1 and 2 (classes are mixed)



Feat2 + Feat3:

Better discrimination for all classes



Feat2 vs Feat3



Toolboxes Loading, EDA/plotting in Matlab



Load the dataset (1/2)

% Load the dataset and cast data

url = 'https://archive.ics.uci.edu/ml/machine-learningdatabases/iris/iris.data';

S = textscan(fid, '%f%f%f%f%s', 'delimiter', ',');

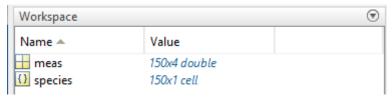
fclose(fid);

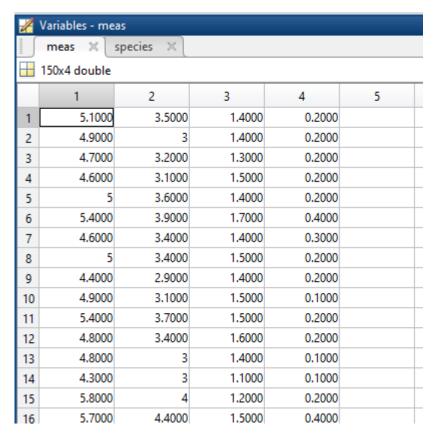
meas = $[S{1}, S{2}, S{3}, S{4}];$

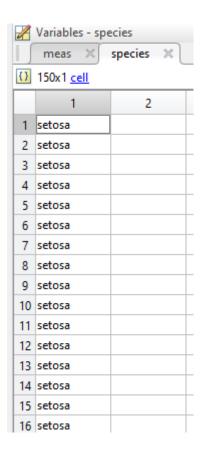
species = $S{5}$;

% Equivalent to "load fisheriris"

Load the dataset (2/2)

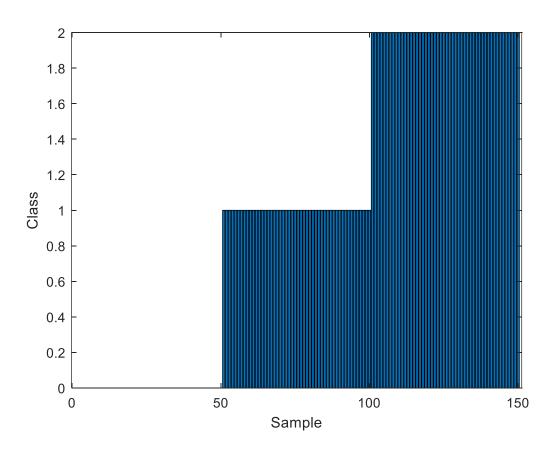






Create the data structures

```
% Problem
P = meas;
% Targets
T = zeros(size(species));
for i = 1: size(T,1)
  switch species{i}
     case 'setosa'
       T(i) = 0;
     case 'versicolor'
       T(i) = 1;
     case 'virginica'
       T(i) = 2;
  end
end
figure
bar(T)
xlabel('Sample')
ylabel('Class')
```



Plotmatrix() \rightarrow Feat_i VS Feat_j

%% creating the feature matrix Problem (X in the course)

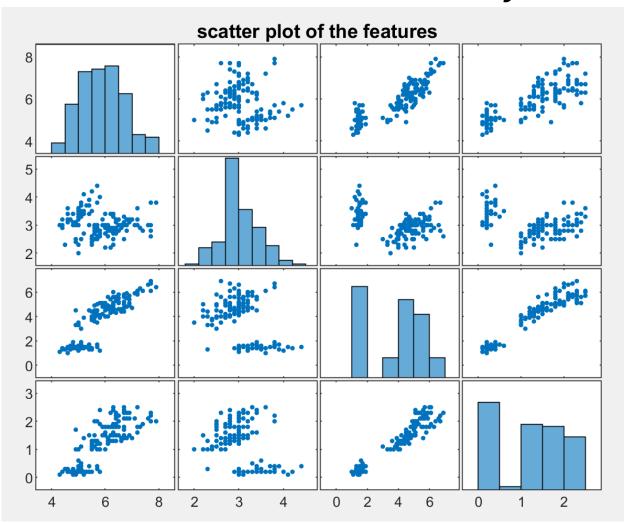
P = meas;

% plot the feature matrix

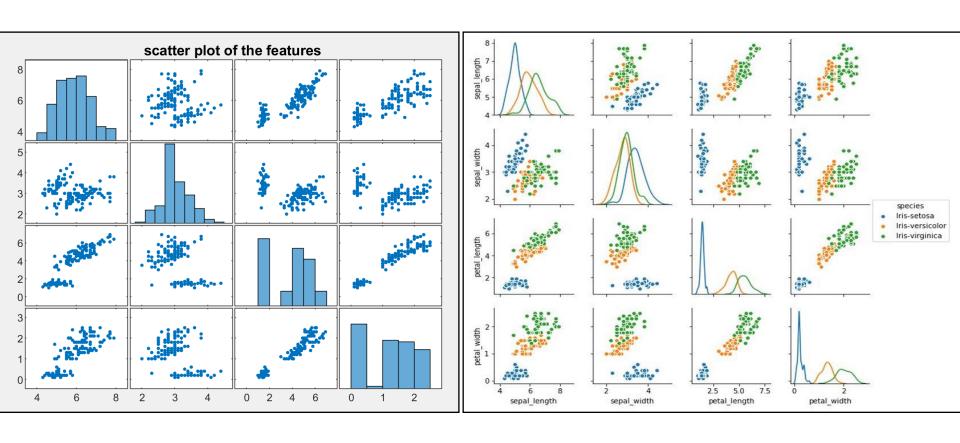
figure

plotmatrix(P)

title('scatter plot of the features')

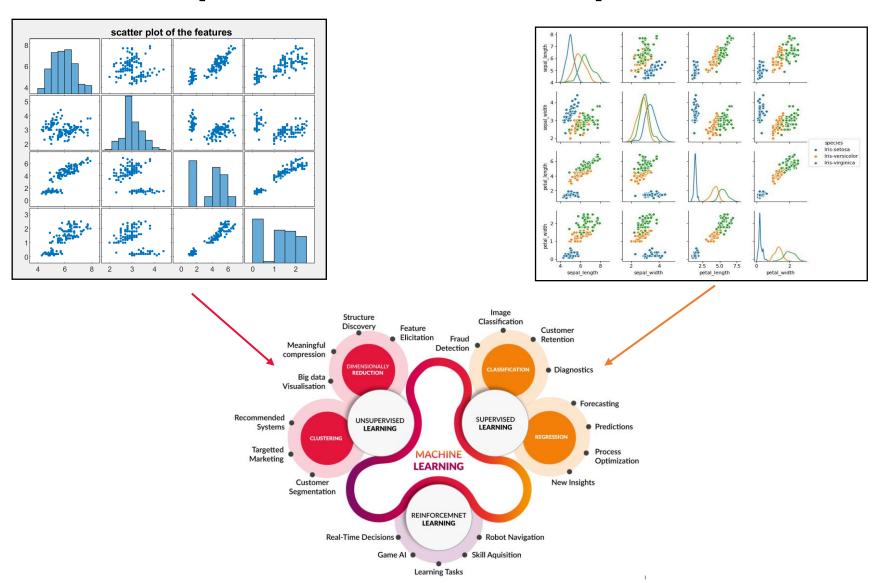


plotmatrix() VS pairplot ()

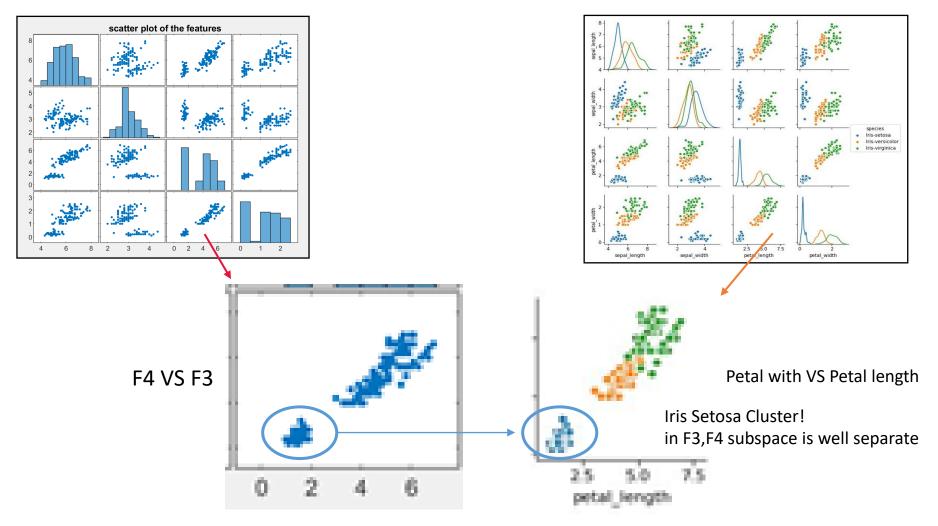


Unsupervised VS Supervised

Unsupervised VS Supervised



Clusters in the unsupervised analysis



Preliminary data analysis

```
statisticsP = zeros(4, size(P,2));
statisticNames{1} = 'Mean';
statisticNames{2} = 'Std ';
statisticNames{3} = 'Min ';
statisticNames{4} = 'Max ';
statisticsP(1,:) = mean(P);
statisticsP(2,:) = std(P);
statisticsP(3,:) = min(P);
statisticsP(4,:) = max(P);
fprintf('\t\tFeat 1\tFeat 2\tFeat 3\tFeat 4\n');
for i = 1:4
  fprintf('%s\t', statisticNames{i});
  for j = 1: size(P,2)
     fprintf('%03.03f\t', statisticsP(i,j));
  end
```

fprintf('\n')

end

0.2000 4.7000 3.2000 1.3000 1.5000 4.6000 3.1000 0.2000 1.4000 3.6000 0.2000 1.7000 5.4000 3.9000 0.4000 4.6000 3.4000 1.4000 0.3000 Matlab is natively 3.4000 1.5000 0.2000 working on 4.4000 2.9000 1.4000 0.2000 1.5000 0.1000 4.9000 3.1000 columns 1.5000 5.4000 3.7000 0.2000 4.8000 3.4000 1.6000 0.2000 4.8000 1.4000 0.1000 4.3000 1.1000 0.1000 mean(P) → [5.843 3.057 3.758 1.199]

5.1000

4.9000

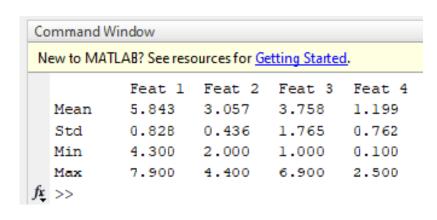
3.5000

0.2000

0.2000

1.4000

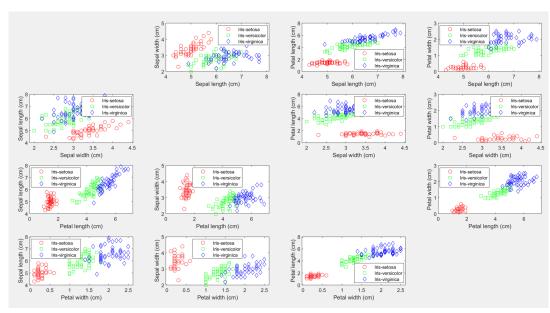
1.4000



Plot the data (1/2)

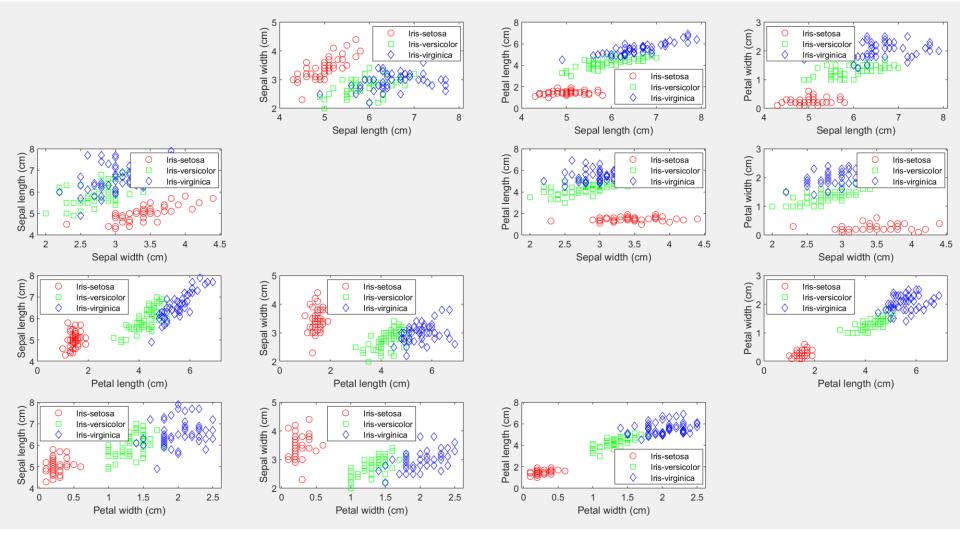
```
% Plot the dataset
variableNames{1} = 'Sepal length (cm)';
variableNames{2} = 'Sepal width (cm)';
variableNames{3} = 'Petal length (cm)';
variableNames{4} = 'Petal width (cm)';
iCount = 1;
figure
for i = 1:4
  for j = 1 : 4
     if (i\sim=j)
       subplot(4, 4, iCount)
       gscatter(P(:,i), P(:,j), species, 'rgb', 'osd');
       xlabel(variableNames{i})
       ylabel(variableNames{j})
     end
     iCount = iCount + 1;
  end
```

end



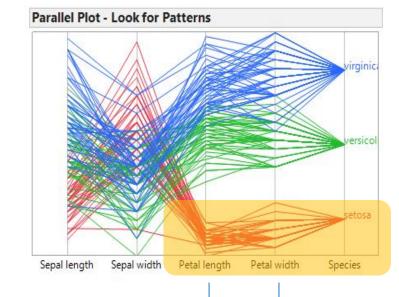
You can create your own visualization plots and dashboard! This is a valuable part of the job...

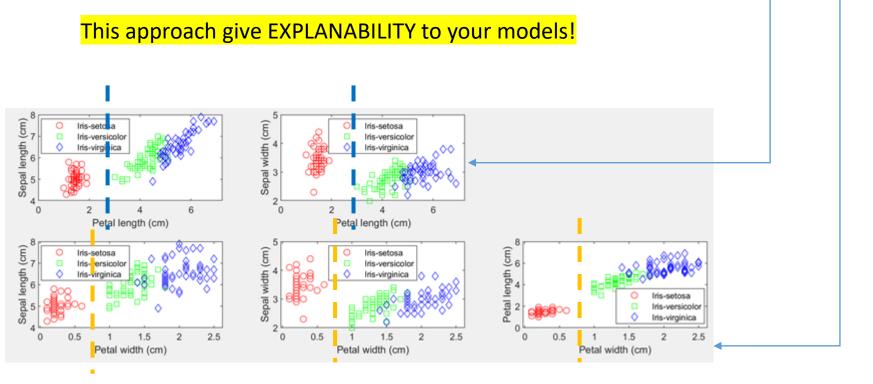
Plot the data (2/2)



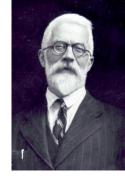
EDA: application

From the Parallel Plot we already verified the separation at least of the Setosa Class
Just using the Petal length or Petal width







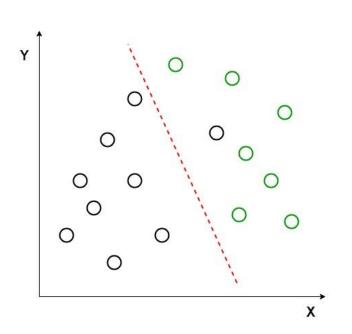


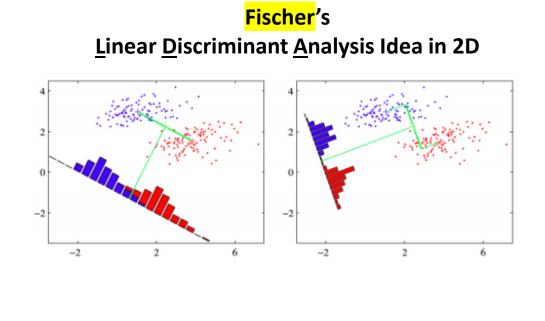
% Linear Discriminant% Analysis

Ida = fitcdiscr(P,T);

IdaClass = resubPredict(Ida);

Just an example: we will present this topic later in the course





Linear Classification

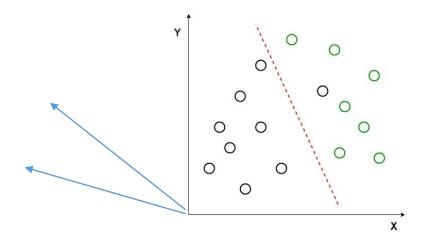
% Linear Discriminant Analysis

lda = fitcdiscr(P,T);

% predict values

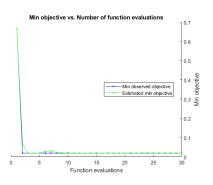
% using the predictor data stored in Ida.X.

ldaClass = resubPredict(lda);



0 0.2000 0 0.2000 0 0.2000 0 0.2000	000	3.5000	5.1000	1
0 0.2000 0 0.2000 0 0.2000	000	_		
0 0.2000 0 0.2000 0 0.2000		3	4.9000	2
0.2000	000	3.2000	4.7000	3
	000	3.1000	4.6000	4
0.4000	000	3.6000	5	5
	000	3.9000	5.4000	6
0.3000	000	3.4000	4.6000	7
0.2000	000	3.4000	5	8
0.2000	000	2.9000	4.4000	9
0.1000	000	3.1000	4.9000	10
0.2000	000	3.7000	5.4000	11
0.2000	000	3.4000	4.8000	12
0.1000	000	3	4.8000	13
0.1000	000	3	4.3000	14
0.2000	000	4	5.8000	15
0.4000	000	4.4000	5.7000	16

Fischer's Linear Discriminant Analysis in 4 dimension

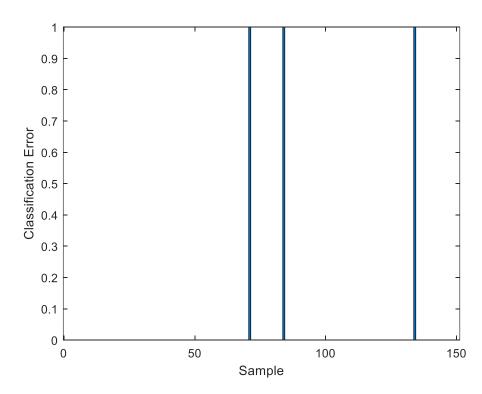


Classification error

% Classification error

```
errorVector = IdaClass ~= T;
```

figure
bar(errorVector);
xlabel('Sample')
ylabel('Classification Error')



Figures of merit (1/2)

```
Command Window
New to MATLAB? See resources for Getting Started.
  Total classification error = 3
  Std of the classification error = 0.140
  Total classification error = 2.000 %
  Std of the classification error = 14.047 %
  confusionMatrix =
       50
              48 1
                     49
f_{\underline{x}} >>
```

Just an example: we will present this topic later in the course

Figures of merit

```
Std of the classification error = 0.140
% Figures of merit
                                              Total classification error = 2.000 %
                                              Std of the classification error = 14.047 %
totalError = sum(errorVector);
stdError = std(errorVector);
totalErrorPerc = mean(errorVector)*100;
stdErrorPerc = stdError * 100;
fprintf('Total classification error = %d \n', totalError);
fprintf('Std of the classification error = %03.03f \n', stdError);
fprintf('Total classification error = %03.03f %% \n', totalErrorPerc);
fprintf('Std of the classification error = %03.03f %% \n', stdErrorPerc);
%Confusion Matrix
                                                         confusionMatrix
confusionMatrix = confusionmat(IdaClass,T)
                                                              50
            Just an example:
we will present this topic later in the course
```

Total classification error = 3