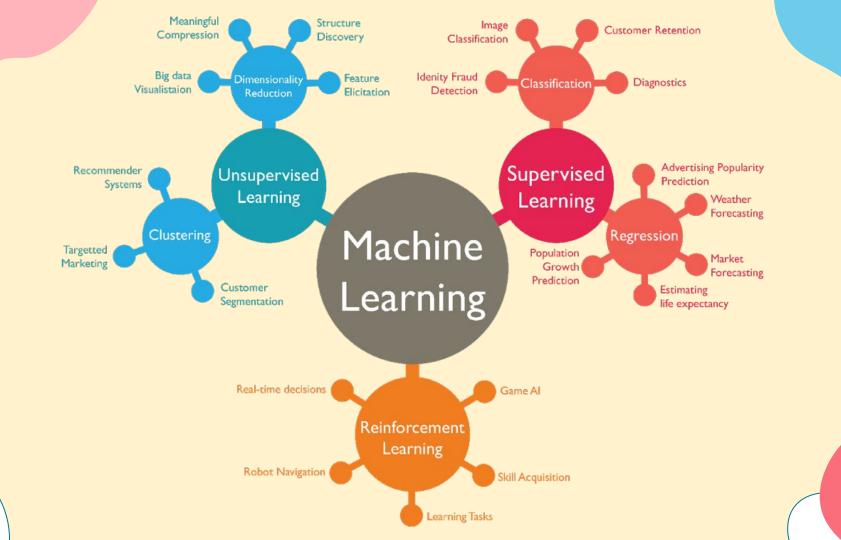
# Capítol 1.2: Machine Learning

Aina Palacios

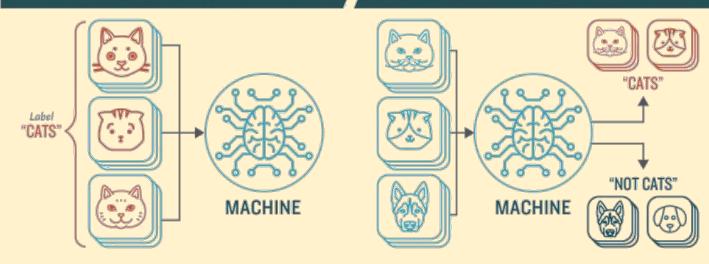


#### How **Supervised** Machine Learning Works

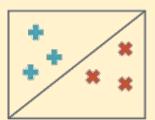
STEPI

Provide the machine learning algorithm categorized or "labeled" input and output data from to learn STEP 2

Feed the machine new, unlabeled information to see if it tags new data appropriately. If not, continue refining the algorithm

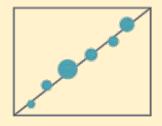


#### TYPES OF PROBLEMS TO WHICH IT'S SUITED



#### **CLASSIFICATION**

Sorting items into categories



#### REGRESSION

Identifying real values (dollars, weight, etc.)

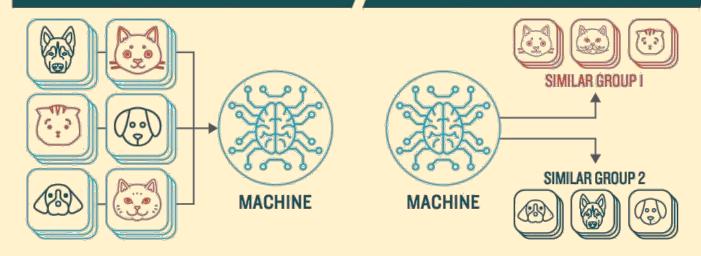
#### **How Unsupervised Machine Learning Works**

STEPI

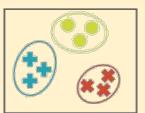
Provide the machine learning algorithm uncategorized, unlabeled input data to see what patterns it finds

STEP 2

Observe and learn from the patterns the machine identifies



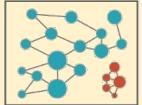
#### TYPES OF PROBLEMS TO WHICH IT'S SUITED



#### **CLUSTERING**

#### Identifying similarities in groups

For Example: Are there patterns in the data to indicate certain patients will respond better to this treatment than others?



#### **ANOMALY DETECTION**

#### Identifying abnormalities in data

For Example: Is a hacker intruding in our network?

## Entorn de treball

#### **Pandas**

Per poder treballar la base de dades



#### Llibreria Sklearn

Per poder aplicar Machine Learning així com preprocessats



- 1. Determinar objectiu i estudi de les dades
- 2. Preprocessat
- 3. Procés enginyeria
- 4. Determinar train i test
- 5. Escollir i entrenar el model
- 6. Mètriques i validació creuada
- 7. Modificar paràmetres

- 1. Determinar objectiu i estudi de les dades
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# 1. Determinar Objectiu i estudi de les dades

Primerament, hem de determinar quin és l'objectiu que estem perseguint. A vegades hem de fixar-nos que les mostres recollides contenen informació a priori i posterior de l'esdeveniment:

- Per exemple, si busquem si un paquet arribarà tard al seu destí i tenim una base de dades amb tot el recorregut del paquet, la informació a posteriori de l'entrega, així com els motius de retard, són dades que no podem utilitzar, ja que no són indicadors prioris del retard de l'entrega.

El nostre objectiu pot ser classificar, predir una regressió o aplicar una tècnica de clústering.

Per poder entendre millor el que estem fent, és important fer un estudi de totes les característiques de la meva mostra. D'aquesta manera, podem aplicar **preprocessats** de les mostres per ajudar als algoritmes a treballar millor.

#### Anàlisi descriptiu i gràfic

- 1. Determinar objectiu i estudi de les dades
- 2. Preprocessat
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## 2. Preprocessat

És important netejar les dades abans d'aplicar cap algoritme sobre elles

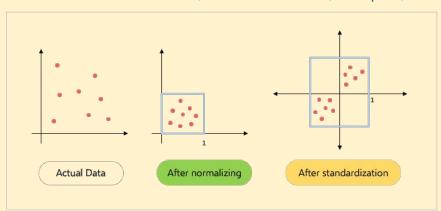
- 1. **Dades no existents**: Nans, Unknown, Null, ext.
  - a. Si són numèriques podem posar a 0, fer interpolació, mitjana,...
  - b. Si són categòriques podem posar un identificador
  - c. Si no es poden substituir, hauriem d'eliminar les dades
- 2. **Dades anòmales**: Hi ha dades que poden no tenir sentit, aplicar el coneixement previ. Un outlier no és una dada anòmala.

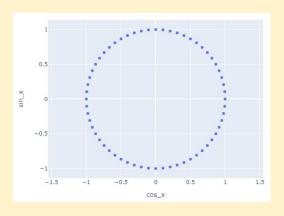
- 1. Determinar objectiu i estudi de les dades
- 2. Preprocessat
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## 3. Procés enginyeria

No aplicar procés sobre el target o invertir el procés

- 3. **Procés d'enginyeria**: Existeixen diferents transformacions que ajuden al model a predir millor.
  - a. Si són gaussianes -> Estandardització
  - b. Si la seva distribució no és normal -> Normalització
  - c. Si conté outliers -> RobustScaler o altres
  - d. Si són categòriques -> Dummys o enumeració
  - e. Altres: Polimorfisme, transformacions, cícliques,...

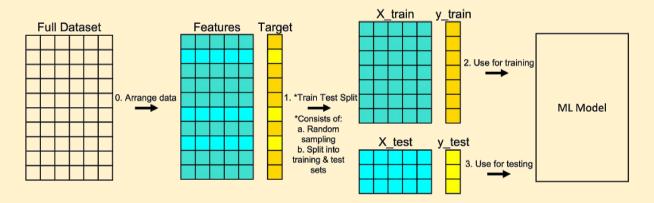




- 1. Determinar objectiu i estudi de les dades
- 2. Preprocessat
- 3. Procés enginyeria
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## 4. Determinar train i test

Si treballem amb Supervised Machine Learning, necessitem dividir les dades entre train i test per tal de poder determinar el millor **model** per aconseguir el nostre objectiu.



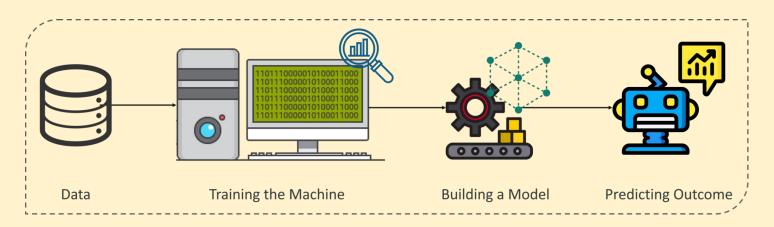
El test l'utilitzarem per veure com de bé el model funciona. Normalment representa un 30% o 20% de les dades.

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2)
```

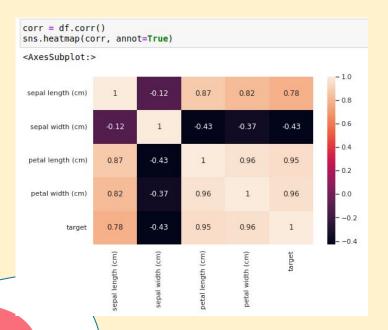
- 1. Determinar objectiu i estudi de les dades
- 2. Preprocessat
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## **Machine Learning**

- → Enllaç a curs de <u>Machine Learning</u>
- → Enllaç Machine Learning amb Sklearn
- → Cheeting sheet Machine Learning amb Sklearn



## Classification



```
from sklearn import neighbors, datasets, preprocessing
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
iris = datasets.load_iris()
```

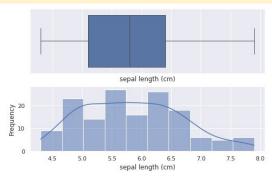
import pandas as panda

import numpy as np

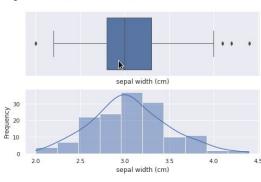
sep	al length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
0	5.1	3.5	1.4	0.2	0.0
1	4.9	3.0	1.4	0.2	0.0
2	4.7	3.2	1.3	0.2	0.0
3	4.6	3.1	1.5	0.2	0.0
4	5.0	3.6	1.4	0.2	0.0

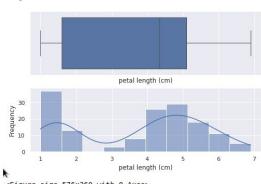
```
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings("ignore")
for i in df.columns:
   plt.figure()
    plt.tight layout()
    sns.set(rc={"figure.figsize":(8, 5)})
    f, (ax box, ax hist) = plt.subplots(2, sharex=True)
    plt.gca().set(xlabel= i,ylabel='Frequency')
    sns.boxplot(df[i], ax=ax box , linewidth= 1.0)
    sns.histplot(df[i], ax=ax hist , bins = 10,kde=True)
```





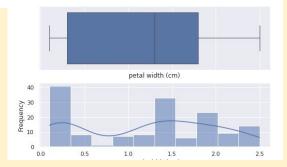
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<Figure size 576x360 with 0 Axes>

<Figure size 576x360 with 0 Axes>

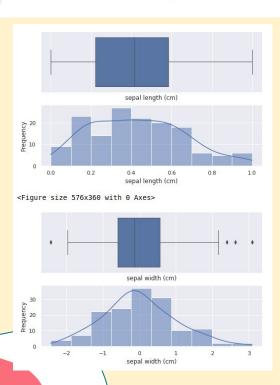


#### 1.2 Preprocessat

: df.isnull().values.any()

: False

No tenim valors nuls i considerem que no existeixen dades anomeles



- La distribució de sepal width és normal, per tant aplicarem estandarització
- · Les altres variables no contenen outliers, per tant utilitzarem normalització

```
from sklearn.preprocessing import StandardScaler
from sklearn.preprocessing import MinMaxScaler
standColumns = ['sepal width (cm)']
scalerStand = preprocessing.StandardScaler().fit(df[standColumns])
df[standColumns] = scalerStand.transform(df[standColumns])
normColumns = ['sepal length (cm)', 'petal length (cm)','petal width (cm)']
scalerNorm = preprocessing.MinMaxScaler().fit(df[normColumns])
df[normColumns] = scalerNorm.transform(df[normColumns])
import matplotlib.pyplot as plt
import seaborn as sns
import warnings
warnings.filterwarnings("ignore")
for i in df.columns:
    plt.figure()
   plt.tight layout()
    sns.set(rc={"figure.figsize":(8, 5)})
    f, (ax box, ax hist) = plt.subplots(2, sharex=True)
    plt.gca().set(xlabel= i,ylabel='Frequency')
    sns.boxplot(df[i], ax=ax box , linewidth= 1.0)
    sns.histplot(df[i], ax=ax hist , bins = 10,kde=True)
```

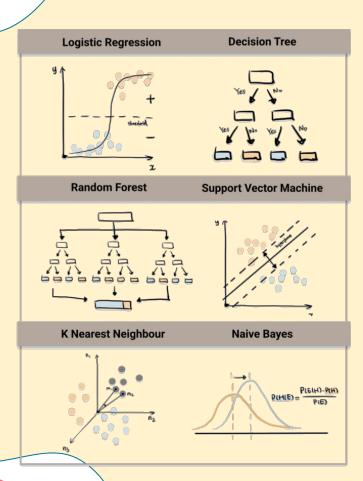
#### 1.3 Test/train

```
from sklearn.model_selection import train_test_split

X = df.drop(['target'],axis=1)
y = df['target']

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.2)
```

Hem utilitzat un test del 20% perquè tenim poques mostres



#### 1.4 Models

```
from sklearn.linear_model import LogisticRegression
lr = LogisticRegression(random_state=42)
```

```
from sklearn.svm import SVC
svc = SVC(kernel='linear')
```

```
from sklearn.ensemble import RandomForestClassifier
rf = RandomForestClassifier(max depth=2, random state=42)
```

```
Training the models
```

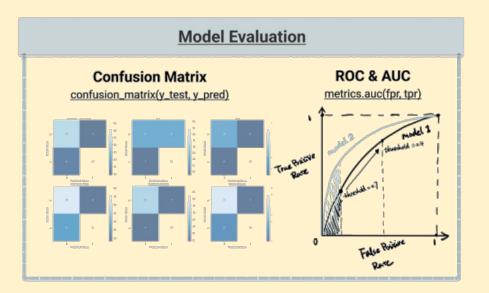
```
lr.fit(X_train, y_train)
svc.fit(X_train, y_train)
rf.fit(X_train, y_train)
```

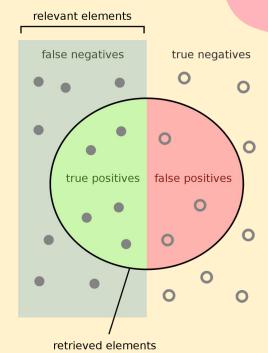
```
y_pred_lr = lr.predict(X_test)
y_pred_svc = svc.predict(X_test)
y_pred_rf = rf.predict(X_test)
```

#### y\_pred\_rf

array([2, 1, 2, 0, 2, 1, 2, 0, 0, 1, 1, 1, 1, 0, 0, 2, 0, 0, 0, 2, 2, 1, 1, 2, 2, 2, 1, 2, 1, 0])

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How many retrieved items are relevant?

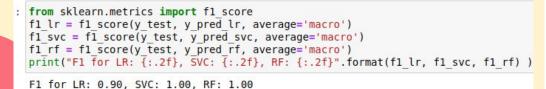
Precision =

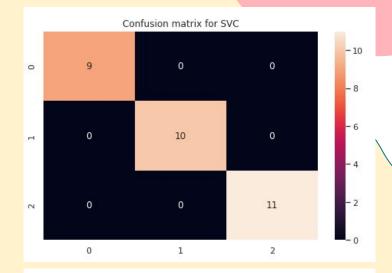
How many relevant items are retrieved?

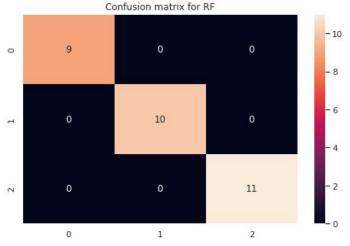


#### 5 Evaluació

```
: from sklearn.metrics import confusion matrix
  cf matrix lr = confusion matrix(y test, y pred lr)
  cf matrix svc = confusion matrix(y test, y pred svc)
  cf matrix rf = confusion matrix(y test, y pred rf)
: sns.heatmap(cf matrix lr, annot=True).set(title='Confusion matrix for LR')
: [Text(0.5, 1.0, 'Confusion matrix for LR')]
                  Confusion matrix for LR
                                                     - 10
  0
                                        11
  N
            0
                          1
```

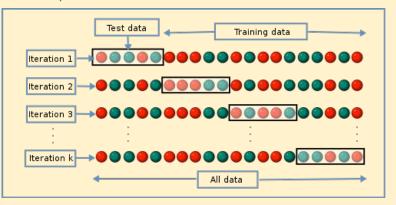






## Validació creuada

Quan creem el test/train pot donar la casualitat que el nostre set afavoreix el nostre model. Per tal de trobar el valor real dels nostres paràmetres, utilitzem CrossValidation.



#### **Cross Validation**

```
from sklearn.model_selection import cross_val_score
print(cross_val_score(lr, X, y, cv=5, scoring='fl_macro'))
```

## Validació creuada

```
cv_lr = cross_val_score(lr, X, y, cv=5, scoring='f1_macro')
print("F1 for LR mean: {:.2f}, std: {:.2f}".format(cv_lr.mean(), cv_lr.std()) )

cv_svc = cross_val_score(svc, X, y, cv=5, scoring='f1_macro')
print("F1 for SVC mean: {:.2f}, std: {:.2f}".format(cv_svc.mean(), cv_svc.std()) )

cv_rf = cross_val_score(rf, X, y, cv=5, scoring='f1_macro')
print("F1 for RF mean: {:.2f}, std: {:.2f}".format(cv_rf.mean(), cv_rf.std()) )

F1 for LR mean: 0.92, std: 0.05
F1 for SVC mean: 0.96, std: 0.03
F1 for RF mean: 0.95, std: 0.03
```

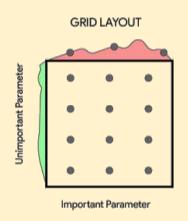
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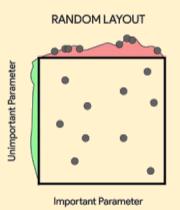
## 7. Modificar paràmetres

El model conté diferents **paràmetres** que modifiquen el seu comportament.

És important intentar trobar aquells paràmetres que donen els millors resultats.

Podem intentar trobar-los directament o utilitzant el **GridSearch** o un **Randomized Parameter Optimization**, que ens provarà totes les combinacions possibles que li diguem i trobarà el millor resultat.

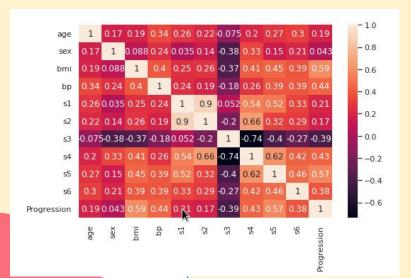




#### 6. Millors paràmetres

```
from sklearn.model selection import RandomizedSearchCV
params = {"solver": ['newton-cg', 'lbfgs', 'liblinear', 'sag', 'saga'], "penalty": ["l1", "l2", "elasticnet"]}
rsearch = RandomizedSearchCV(estimator=lr,
                             param distributions=params,cv=5,
                             n iter=14, random state=42, scoring='f1 macro')
rsearch.fit(X, y)
print(rsearch.best score )
print(rsearch.best params )
0.9530138477506899
{'solver': 'saga', 'penalty': 'l1'}
y pred RS ls = LogisticRegression(random state=42, solver = 'saga', penalty = 'l1')\
    .fit(X train,y train).predict(X test)
cf matrix RS lr = confusion matrix(y test, y pred RS ls)
sns.heatmap(cf matrix RS lr. annot=True).set(title='Confusion matrix for LR')
[Text(0.5, 1.0, 'Confusion matrix for LR')]
                Confusion matrix for LR
                                                  - 10
         11
                       11
                        1
                                       2
```

# Regression



#### **Machine Learning**

#### Regression

Utilitzarem la base de dades Iris, que ens proporciona un Data Set sobre tipus de plantes i els seus pètals

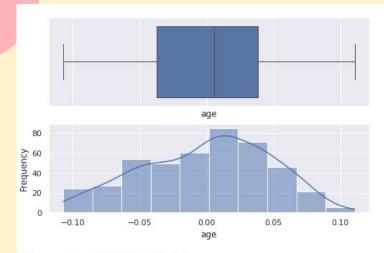
```
from sklearn import neighbors, datasets, preprocessing
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score
diabetes = datasets.load diabetes()
```

import pandas as panda
import numpy as np

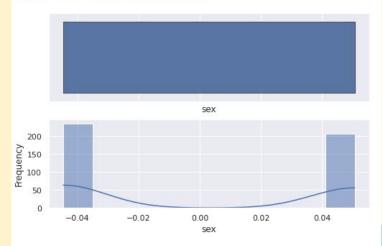
df = panda.DataFrame(diabetes.data,columns=diabetes.feature\_names)
df['Progression'] = diabetes.target

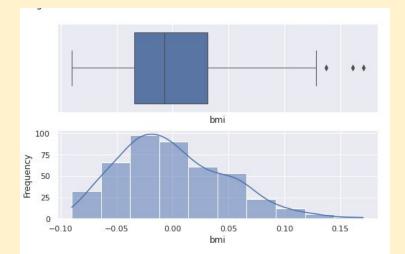
df.head()

	age	sex	bmi	bp	s1	s2	s3	s4	<b>s</b> 5	s6	Progression
0	0.038076	0.050680	0.061696	0.021872	-0.044223	-0.034821	-0.043401	-0.002592	0.019907	-0.017646	151.0
1	-0.001882	-0.044642	-0.051474	-0.026328	-0.008449	-0.019163	0.074412	-0.039493	-0.068332	-0.092204	75.0
2	0.085299	0.050680	0.044451	-0.005670	-0.045599	-0.034194	-0.032356	-0.002592	0.002861	-0.025930	141.0
3	-0.089063	-0.044642	-0.011595	-0.036656	0.012191	0.024991	-0.036038	0.034309	0.022688	-0.009362	206.0
4	0.005383	-0.044642	-0.036385	0.021872	0.003935	0.015596	0.008142	-0.002592	-0.031988	-0.046641	135.0

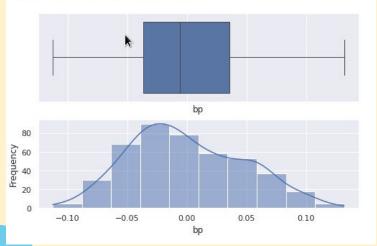


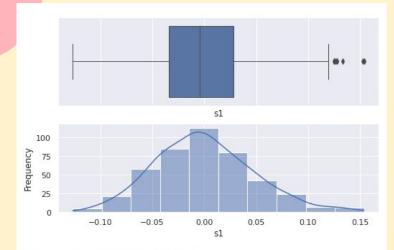
<Figure size 576x360 with 0 Axes>



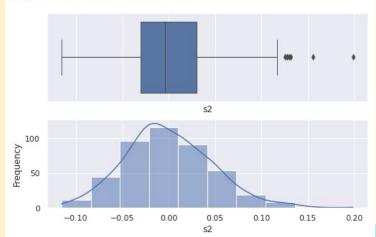


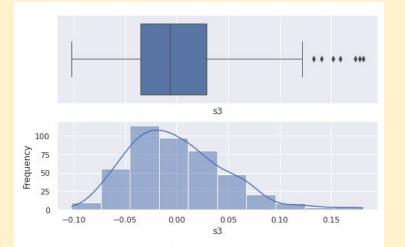
<Figure size 576x360 with 0 Axes>



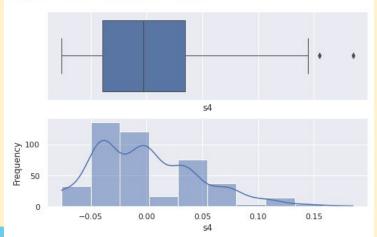


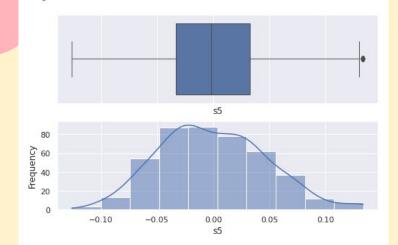
<Figure size 576x360 with 0 Axes>



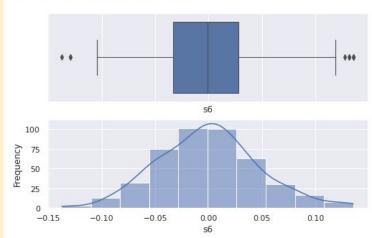


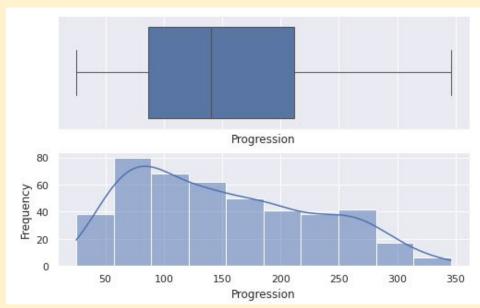
<Figure size 576x360 with 0 Axes>





<Figure size 576x360 with 0 Axes>



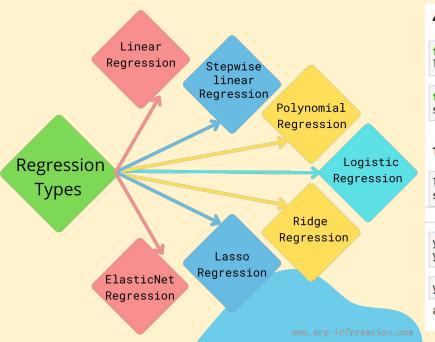


#### 3 Test/train

```
from sklearn.model_selection import train_test_split

X = df.drop(['Progression'],axis=1)
y = df['Progression']

X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 0.3)
```



#### 4 Models

from sklearn.linear\_model import LinearRegression
lr = LinearRegression()

from sklearn.svm import SVR
svr = SVR()

#### Training the models

lr.fit(X\_train, y\_train)
svr.fit(X train, y train)

# - R - R -

y\_pred\_lr = lr.predict(X\_test)
y\_pred\_svr = svr.predict(X\_test)

y\_pred\_lr[1:5]

array([135.69448628, 196.53507727, 74.45375974, 87.28375355])

#### 5 Evaluació

	Model	R2	MSE
0	Linear Regression	0.502632	3289.974895
1	Support Vector Machines	0.157471	5573 138008

#### **Cross Validation**

```
from sklearn.model_selection import cross_val_score

lr = LinearRegression()
cv_lr = cross_val_score(lr, X, y, cv=5, scoring='r2')
cv_svr = cross_val_score(svr, X, y, cv=5, scoring='r2')
```

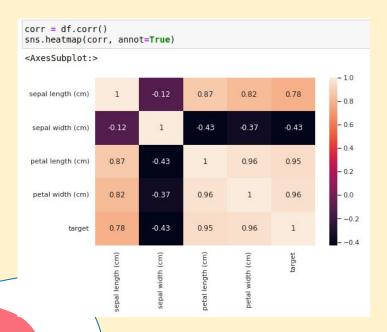
```
cv_lr_mean = [cv_lr.mean(), cv_svr.mean()]
metrics['R2 with CV'] = cv_lr_mean
metrics.head()
```

	Model	R2	MSE	R2 with CV	
0	Linear Regression	0.255382	3761.752362	0.482316	
1	Support Vector Machines	0.126417	4413.269770	0.161335	

#### 6. Millors paràmetres

```
from sklearn.model selection import RandomizedSearchCV
param = {'kernel' : ('linear', 'poly', 'rbf', 'sigmoid'),'C' : [1,5,10],'degree' : [3,8],
         'coef0' : [0.01,10,0.5], 'gamma' : ('auto', 'scale')}
lr = LinearRegression()
rsearch = RandomizedSearchCV(estimator=svr,
                              param distributions=param, cv=5,
                              n iter=10, random state=42, scoring='r2')
rsearch.fit(X, y)
print(rsearch.best score )
print(rsearch.best params )
0.47480829897832244
{'kernel': 'linear', 'gamma': 'scale', 'degree': 3, 'coef0': 0.5, 'C': 10}
svr bp = SVR(kernel='linear', qamma='scale', degree= 3,coef0= 0.5,C= 10).fit(X train, y train)
y pred svr bp = svr bp.predict(X test)
metrics bp = panda.DataFrame( data = [['SVR best params', r2 score(y test, y pred svr bp),
                          mean squared error(y test, y pred svr bp),
                          cross val score(svr bp, X, y, cv=5, scoring='r2').mean()]],
                                   columns = ['Model', 'R2', 'MSE', 'R2 with CV'])
metrics = metrics.append(metrics bp, ignore index=True)
metrics.head()
                         R2
               Model
                                  MSE R2 with CV
       Linear Regression 0.255382 3761.752362
                                        0.482316
1 Support Vector Machines 0.126417 4413.269770
                                        0.161335
        SVR best params 0.273890 3668.252470
                                        0.474808
```

# Unsupervised



```
from sklearn import neighbors, datasets, preprocessing
from sklearn.model selection import train test split
from sklearn.metrics import accuracy score
iris = datasets.load iris()
```

```
import numpy as np
df= pd.DataFrame(data= np.c [iris['data'], iris['target']],
```

import pandas as pd

columns= iris['feature names'] + ['target']) df.head()

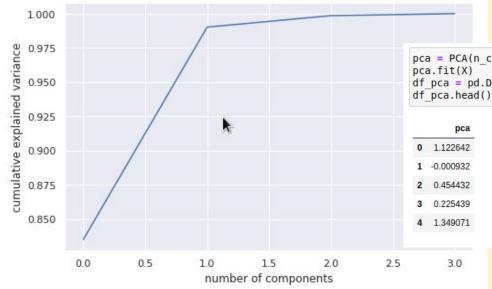
se	epal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target
0	5.1	3.5	1.4	0.2	0.0
1	4.9	3.0	1.4	0.2	0.0
2	4.7	3.2	1.3	0.2	0.0
3	4.6	3.1	1.5	0.2	0.0
4	5.0	3.6	1.4	0.2	0.0

#### 3 PCA (Principal Component Analisis)

```
from sklearn.decomposition import PCA

X = df.drop(['target'],axis=1)
y = df['target']

pca = PCA().fit(X)
plt.plot(np.cumsum(pca.explained_variance_ratio_))
plt.xlabel('number of components')
plt.ylabel('cumulative explained variance');
```

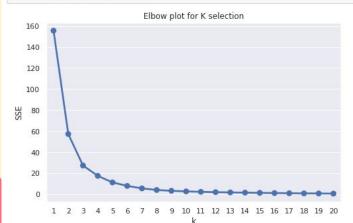


pca = PCA(n\_components=1)
pca.fit(X)
df\_pca = pd.DataFrame(pca.transform(X), columns=['pca'], index=df.index)
df pca.head()

#### 4 Models

```
from sklearn.cluster import KMeans
from kneed import KneeLocator
def elbow plot(df):
    """Create elbow plot from normalized data"""
    sse = {}
    sse r = []
    for k in range(1, 21):
        kmeans = KMeans(n clusters=k, random state=1)
        kmeans.fit(df)
        sse[k] = kmeans.inertia
        sse r.append(kmeans.inertia )
    plt.title('Elbow plot for K selection')
    plt.xlabel('k')
    plt.ylabel('SSE')
    sns.pointplot(x=list(sse.keys()),
                 y=list(sse.values()))
    plt.show()
    return sse r
```

```
sse = elbow_plot(df_pca)
```



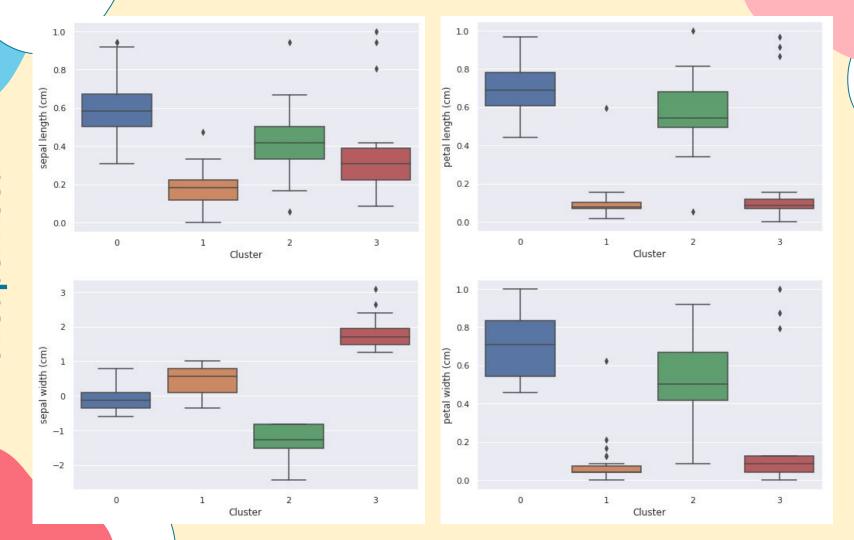
```
kl = KneeLocator(range(1, 21), sse, curve="convex", direction="decreasing")
kl.elbow
```

4

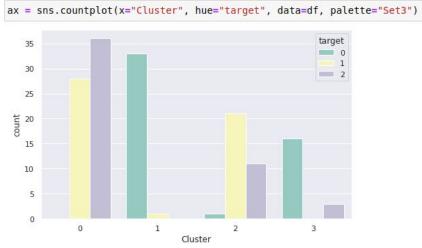
```
k_means = KMeans(n_clusters=4, random_state=42)
k_means.fit(X)
```

```
df['Cluster'] = y_pred
df.head()
```

	sepal length (cm)	sepal width (cm)	petal length (cm)	petal width (cm)	target	Cluster
0	0.222222	1.019004	0.067797	0.041667	0	1
1	0.166667	-0.131979	0.067797	0.041667	0	1
2	0.111111	0.328414	0.050847	0.041667	0	1
3	0.083333	0.098217	0.084746	0.041667	0	1
4	0.194444	1.249201	0.067797	0.041667	0	3







# Jahem acabat la part 2! Gràcies a tots!

