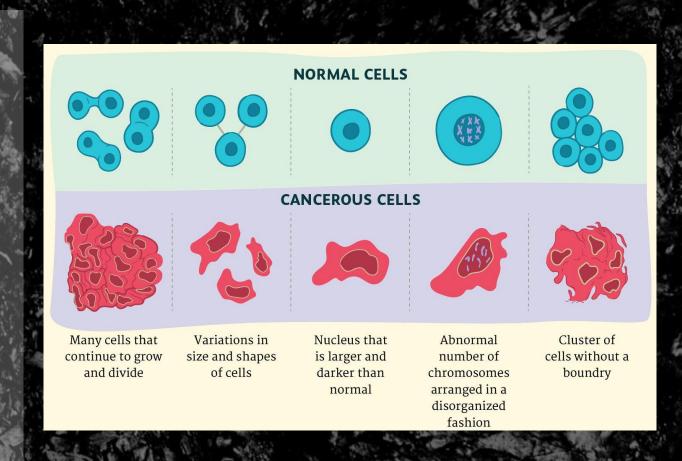
HADACA project

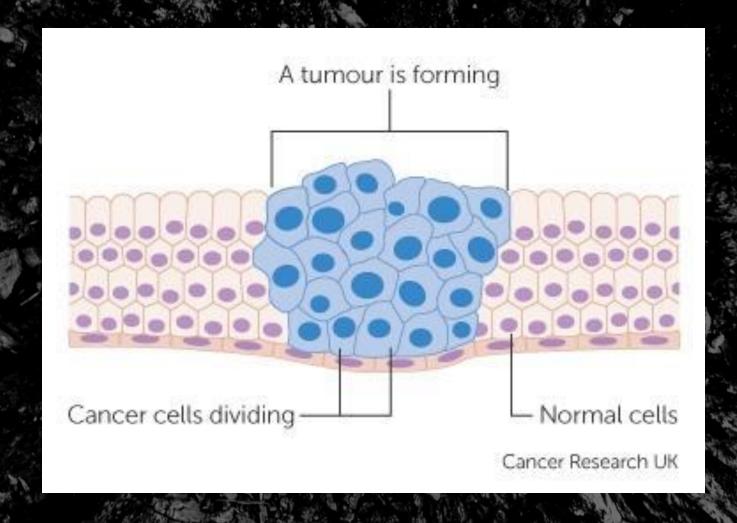
Group FACTOR

What is this project about?

- About 90.5 million people have cancer.
- Only in 2017, 400.000 new types of cancer were discovered.
- Unfortunately, even in age of technologies, there is still no absolute cure for it.

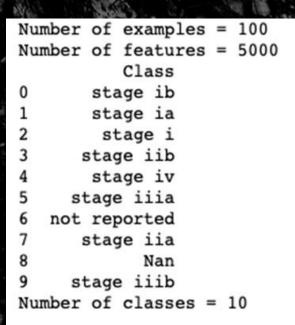


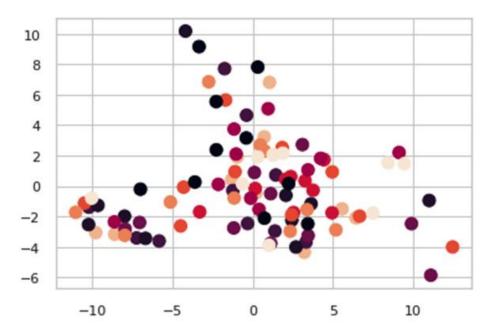
This project is a way to help the research advance and fight the scourge of 21st century.



We will use Machine Learning to automate the process of diagnosing the tumors on different stages and make it easier to define the exact stage of cancer.
We have data based on dataset of 5000 real patients with 10

different types of cancer.

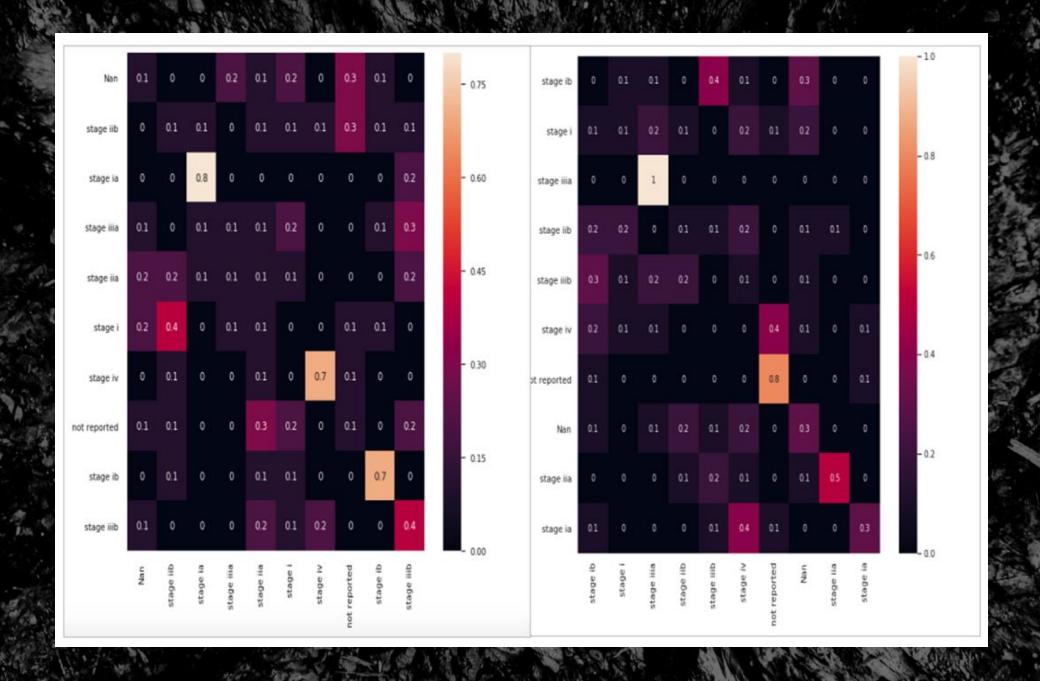


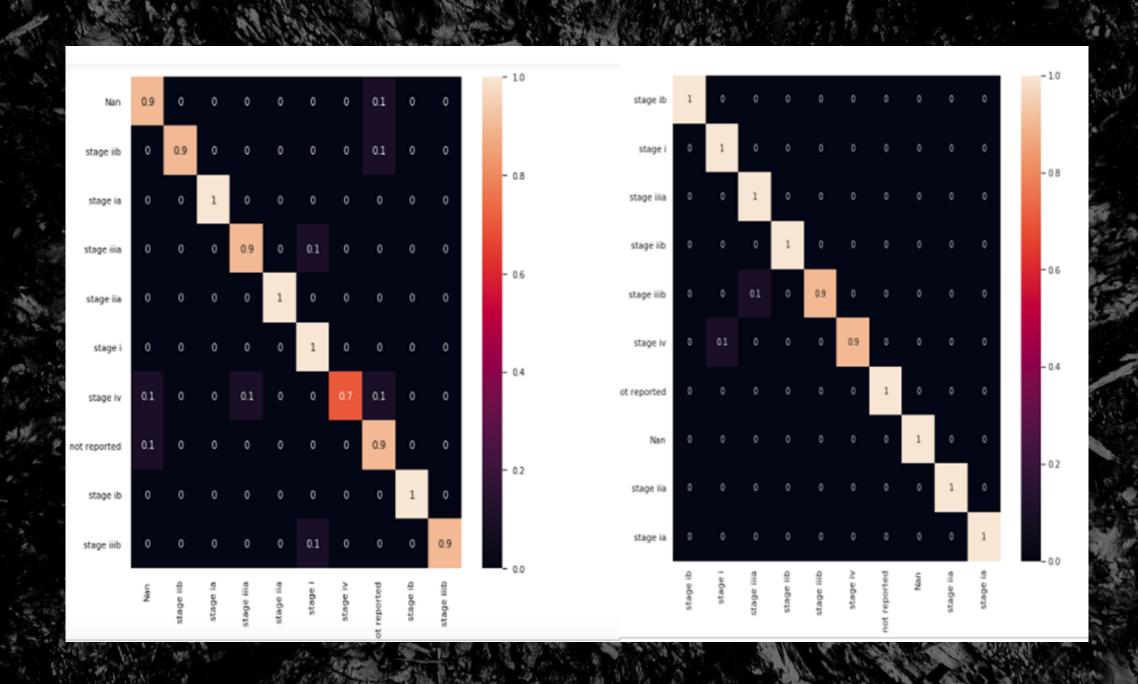




Confusion matrix

• The confusion matrix is, in the terminology of supervised learning, a tool for measuring the quality of a classification system. One of the interests of the confusion matrix is that it quickly indicates whether a classification system is able to classify correctly.







• Preprocessing is all about sorting and selecting accurate data to optimize the process of classification and to make it as easy and fast as possible.

```
import pandas as pd
                                                      // Importing the librairies we need
From data.io import read as df
Data = read as df('data dir + '/' + 'data name')
                                                      // Importing the dataset
X = dataset.iloc[:.:-1].values
                                                      // Create a matrix of features in our dataset
From sklearn.preprocessing import Imputer
                                                      // Taking care of missing data
Imputer = Imputer(missing values = "NaN",
                                                      // With the class called imputer in sklearn.preprocessing
          strategy = "mean", axis = 0)
                                                           we will search for missing data
Imputer = imputer.fit(X[:,1:3])
X[:, 1:3] = imputer.transform(X[:, 1:3])
                                                      // Now we will just replace the missing values with
                                                         the mean of the column by the method transform.
from sklearn.model selection import train test split // We separe the data into three datasets, for training,
X train, X test, Y train, Y test =
                                                         testing and validation.
           train test split(X,Y, test size=0.2)
from sklearn.preprocessing import StandardScaler
sc X = StandardScaler()
X train = sc X.fit transform(X train)
                                                      // We applied a feature scaling to the data :
X \text{ test} = \text{sc } X.\text{transform}(X \text{ test})
                                                          It is a method used to standardize the range of
                                                          independent variables or features of data.
```

Classification

• There are many ways to carry on the classification of data. One of them is to use the perceptron. We used the multiclass perceptron.

```
Algorithm 5 PERCEPTRONTRAIN(D, MaxIter)
 w_d \leftarrow o, for all d = 1 \dots D
                                                                          // initialize weights
 b \leftarrow 0
                                                                              // initialize bias
 3: for iter = 1 ... MaxIter do
       for all (x,y) \in D do
          a \leftarrow \sum_{d=\tau}^{D} w_d x_d + b
                                                     // compute activation for this example
          if ya \leq o then
             w_d \leftarrow w_d + yx_d, for all d = 1 \dots D
                                                                           // update weights
             b \leftarrow b + y
                                                                                // update bias
          end if
       end for
 ... end for
 12: return w_0, w_1, ..., w_D, b
Algorithm 6 PerceptronTest(w_0, w_1, ..., w_D, b, \hat{x})
 a \leftarrow \sum_{d=1}^{D} w_d \hat{x}_d + b
                                                 // compute activation for the test example
 2: return SIGN(a)
```

Conclusion: (Results)

$\underline{\textbf{Table 2}}: \textbf{Preliminary Results}$

Dataset	Base Estimator (starting kit)	Naive Bayes Classifier	SVM	Decision Tree	Random Forest	Perceptron
Training	0.8991	0.3860	0.8994	0.8996	0.9980	0.2100
Cross- Validation	0.6872	0.8233	0.834	0.8677	0.8666	0.6581
Validation	0.41	0.49	0.42	0.43	0.39	0.67

RESULTS									
#	User	Entries	Date of Last Entry	Prediction score	Duration 🛦	Detailed Results			
1	luc.gibaud	4	01/24/19	0.9843 (1)	0.00 (1)	View			
2	Malikkazi	3	01/24/19	0.9804 (2)	0.00 (1)	View			
3	takfarinas.nait-larbi	5	03/22/19	0.9790 (3)	0.00 (1)	View			
4	doctor	25	03/22/19	0.9790 (3)	0.00 (1)	View			
5	martin.bauw	32	01/24/19	0.9721 (4)	0.00 (1)	View			
6	Cancer	23	03/24/19	0.9506 (5)	0.00 (1)	View			
7	HEALTH	11	03/24/19	0.8776 (6)	0.00 (1)	View			
8	Cure	19	03/24/19	0.5932 (7)	0.00 (1)	View			
9	Zhengying	4	02/08/19	0.3589 (8)	0.00 (1)	View			
10	FACTOR	8	03/26/19	0.2399 (9)	0.00 (1)	View			
11	OmarAbdoulayeBADIANE	2	03/26/19	0.2399 (9)	0.00 (1)	View			