



[08.10.2021]

Bulky mediastinal lymphoma classification with ML-techniques

Activity report

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Meeting ML_INFN @ Firenze [8 October 2021]

Objective

Classification of bulky mediastinal lymphoma using **Machine Learning** techniques applied to PET and CT images.

Lymphoma types:

- 1. Hodgkin (HL)
- 2. Gray Zone (GZ)
- 3. Primary Mediastinal Lymphoma (PML)

PLANNED

From low-level data

- Data: PET and CT images
- <u>Technique</u>: Image classification
- Algorithm: Convolutional Neural Nets

IN PROGRESS

From high-level data

- <u>Data:</u> features drawn by <u>LIFEx</u> from images
- Technique: Binary and multiclass classification
- Algorithm: Logistic regression + Random Forest

First attempts: high-level dataset

<u>Database of 119 rows x 101 columns</u>

- different data types → need for homogeneous dataset
 - numbers
 - dates
 - o texts
 - o intervals
- dim[instance space] ~ dim[feature space] → need for reducing the feature space

Public repository on GitHub to **prepare**, **clean** and **study** the high-level dataset through <u>Jupyter Notebooks</u>:

- data_preparation -- data format correction
- data_visualization -- correlated features removal
- binary classification -- classification in HL and non-HL
- multiclass classification -- classification in HL, GZ and PML

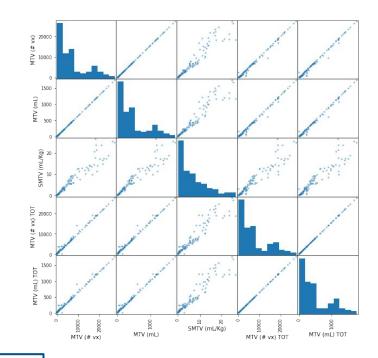


Feature list

```
ID
Lymphoma typer (HL =1, GZ = 2, PML =3)
                                                                 NGLDM Coarseness
                                                                                               CHECK Cluster(s)ToSmall
Data nascita
                                                                 NGLDM Contrast
                                                                                               GLCM Homogeneity[=InverseDifference]
dPET staging
                           SHAPE_Volume(mL)
                                                                                               GLCM_Energy[=AngularSecondMoment]
                                                                 NGLDM Busyness
SUVmin (SUV)
                           SHAPE Volume(vx)
                                                                                               GLCM Contrast[=Variance]
                                                                 GLZLM SZE
SUVmean (SUV)
                           SHAPE Sphericity[onlyFor3DROI])
                                                                 GLZLM LZE
                                                                                               GLCM Correlation
SUVstd (SUV)
                           SHAPE Surface(mm2)[onlyFor3DROI]
                                                                 GLZLM LGZE
                                                                                               GLCM Entropy log10
SUVmax (SUV)
                           SHAPE Compacity[onlyFor3DROI]
                                                                                               GLCM Entropy log2[=JointEntropy]
                                                                 GLZLM HGZE
MTV (\# vx)
                           PARAMS_DistanceOfNeighbours
                                                                 GLZLM SZLGE
                                                                                               GLCM Dissimilarity
MTV (mL)
                           PARAMS NumberOfGreyLevels
                                                                 GLZLM SZHGE
                                                                                               GLRLM SRE
SMTV (mL/Kg)
                           PARAMS BinSize
                                                                 GLZLM_LZLGE
                                                                                               GLRLM LRE
TLG (SUV*mL)
                           PARAMS_IntensityResampling
                                                                GLZLM LZI
                                                                          CONVENTIONAL SUVbwn
                                                                                              GLRLM LGRE
STLG (SUV*mL/Kg)
                           PARAMS BoundsRangeOfValueAfterDiscre
                                                                GLZLM GLI
                                                                          CONVENTIONAL SUVbwn
                                                                                              GLRLM HGRE
MTV (# vx) TOT
                           PARAMS_ZSpatialResampling
                                                                 GLZLM_ZLI
                                                                                              GLRLM_SRLGE
                                                                          CONVENTIONAL SUVbws
MTV (mL) TOT
                                                                                                                   number
                           PARAMS YSpatialResampling
                                                                GLZLM ZP
                                                                                              GLRLM SRHGE
                                                                          CONVENTIONAL SUVbwn
SMTV (mL/Kg) TOT
                           PARAMS XSpatialResampling
                                                                                                                   date
                                                                TimePosit
                                                                                              GLRLM_LRLGE
                                                                          CONVENTIONAL SUVbw
TLG (SUV*mL) TOT
                                                                 zLocation
                                                                                              GLRLM LRHGE
                                                                                                                   text
                                         e only for PEI or NM)
                                                                          CONVENTIONAL SUVbwd
STLG (SUV*mL/Kg) TOT
                                        tized volume sought
                                                                                              GLRLM GLNU
                                                                           CONVENTIONAL SUVbw
                                                                                                                   interval
   DISCRETIZED SUVbwpeakSphere1mL(value only for PET or NM)
                                                                                              GLRLM_RLNU
                                                                           CONVENTIONAL SUVbws
   DISCRETIZED TLG(mL)[onlyForPETorNM]
                                                                                              GLRLM RP
                                                                           CONVENTIONAL SUVbwl
   DISCRETIZED_HISTO_Skewness
                                                                           CONVENTIONAL SUVbw
   DISCRETIZED HISTO Kurtosis
                                                                           CONVENTIONAL SUVbwpeakSphere0,5mL:discretized volume sought
   DISCRETIZED HISTO ExcessKurtosis
                                                                           CONVENTIONAL SUVbwpeakSphere0,5mL(value only for PET or NM)
   DISCRETIZED_HISTO_Entropy_log10
                                                                           CONVENTIONAL SUVbwpeakSphere1mL:discretized volume sought
   DISCRETIZED HISTO Entropy log2
                                                                           CONVENTIONAL SUVbwpeakSphere1mL(value only for PET or NM)
   DISCRETIZED_HISTO_Energy[=Uniformity]
                                                                           CONVENTIONAL TLG(mL)[onlyForPETorNM]
```

Dataset cleaning

- Features with no clear contents dropped
 - o text data
 - o intervals
- Variables with strong correlations dropped
 - $\circ \ \rho_{X,Y} \geq |0.75| {\rightarrow} \, \mathrm{strong} \; \mathrm{correlation}$
- New feature added
 - o age, from dates subtraction (then dropped)



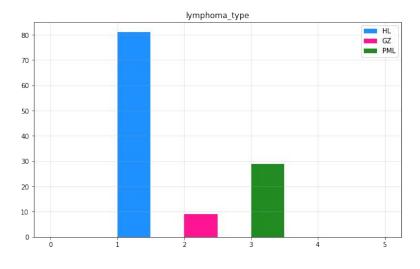
99 → **17 FEATURES**

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Bulky mediastinal lymphoma classification with ML-techniques

binary_classification

Instance space



- Missing information
 - o some features not available for all the instances
 - o instances **dropped** to keep homogeneity
 - \circ 119 \rightarrow 101 rows (instances)
- Dataset strongly unbalanced
 - HL-class is <u>over-represented</u> w.r.t. the other two ones
 - o classification suffers from unbalancing
- Binary classification
 - HL and non-HL classification
 - o dataset a bit more balanced
 - o NHI-HI ratio: 55.4%
- Train/Test split: **80%/20%**
- k-Folds Cross Validation with k=3

Performance measures: accuracy, precision & recall

The **accuracy** is generally not the preferred performance measure for classifiers, especially when one are dealing with **unbalanced datasets**. In fact, classifiers succeed in recognizing instances belonging to most frequent label, resulting in <u>high accuracy</u> even if the other instances are mis-classified.

In order to ensure a model evaluation <u>robust</u> against unbalanced datasets, one can use **precision** or **recall** scores.

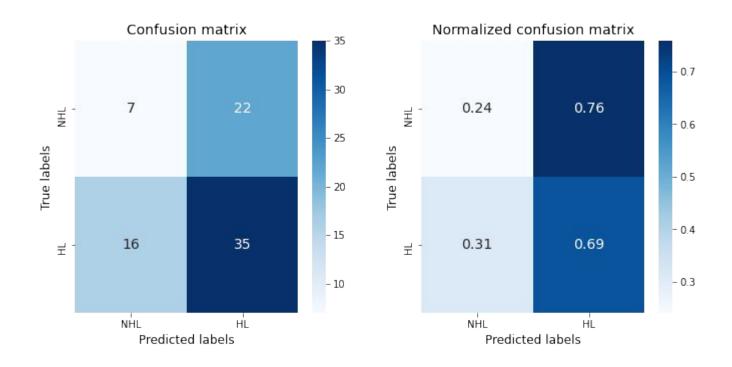
In the Jupyter Notebook we have used RECALL!

$$ccuracy = \frac{TP + TN}{TP + FP + TN + FN} \qquad precision = \frac{TP}{TP + FP}$$

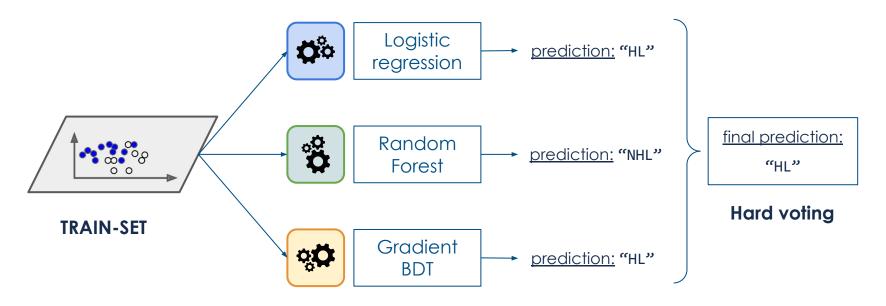
Predicted labels Negative Positive Negative True **False Negative Positive Actual labels** (FP) (TN) False True **Negative Positive** (FN) (TP)

$$recall = \frac{TP}{TP + FN}$$

Dummy Classifier: performance

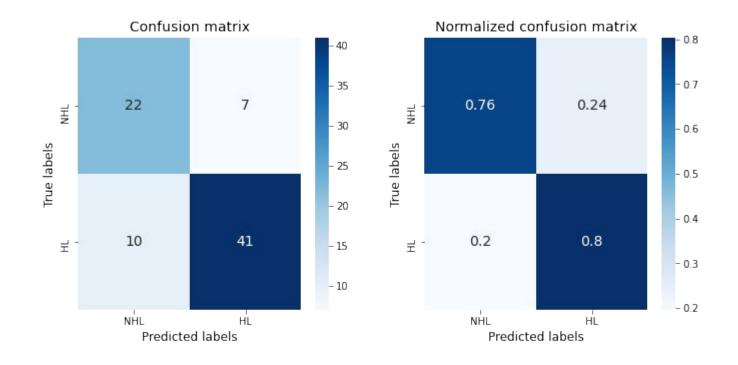


Model combination

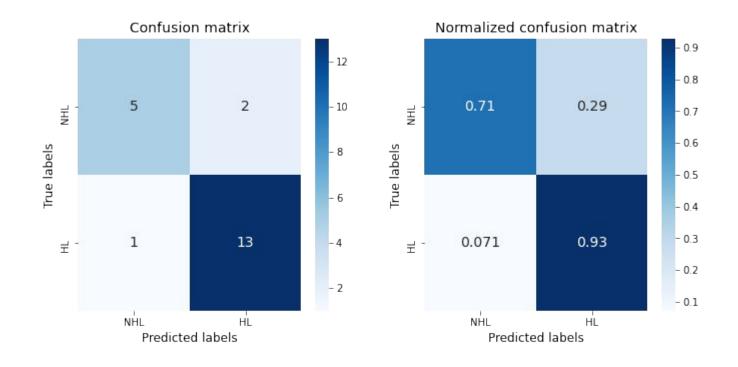


Diverse predictors

Model combination: performance



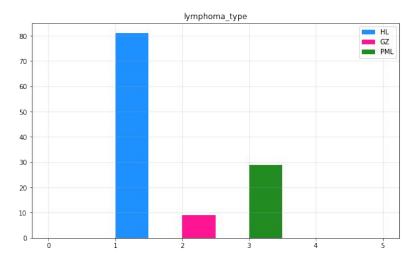
Results on the test-set



Bulky mediastinal lymphoma classification with ML-techniques

 ${\tt multiclass_classification}$

Instance space

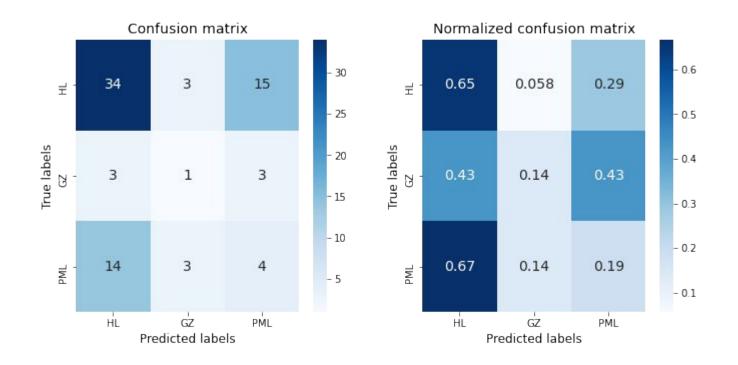


- Missing information
 - o some <u>features not available</u> for all the instances
 - instances dropped to keep homogeneity
 - \circ 119 \rightarrow 101 rows (instances)
- Dataset strongly unbalanced
 - HL-class is <u>over-represented</u> w.r.t. the other two ones
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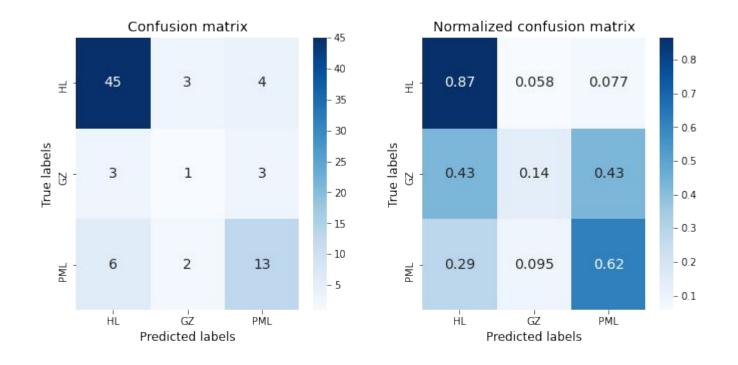
Multiclass classification

- HL, GZ and PML classification
- o GZ-HL ratio: 13.8%
- o PML-HL ratio: 41.5%
- Train/Test split: **80%/20%**
- k-Folds Cross Validation with k=3

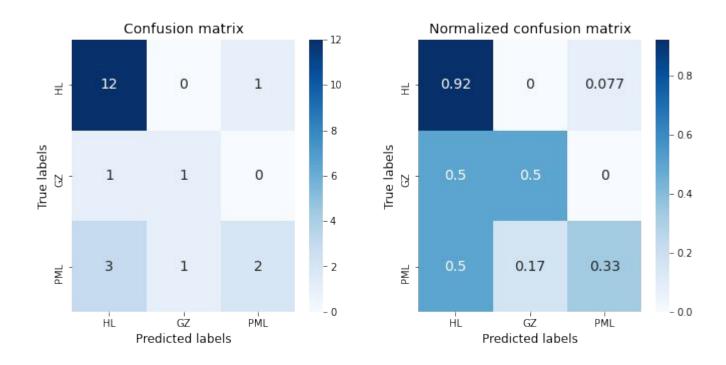
Dummy Classifier: performance



Model combination: performance



Results on the test-set



Conclusion

- Preliminary studies
- There is room for improvements
 - increasing the dataset → more data
 - balancing the dataset → expanding low-represented classes
 - o working with low-level data → image classification techniques
- Binary classification looks promising
 - GZ-class should be treated as a classification uncertainty
 - custom probability thresholds → multiclass classification

Open to any kind of suggestions!

Backup

Train/Test split

DATASET | size: 101 | ratio: 100%

TRAIN-SET | size: 80 | ratio: 80%

TEST-SET | size: 21 | ratio: 20%

To ensure that our Machine Learning model will be able to **generalize** well to new cases, namely to behave like a *true predictor*, we should avoid to use <u>all the dataset</u> to train our model. A better option is to **split the data into two sets**:

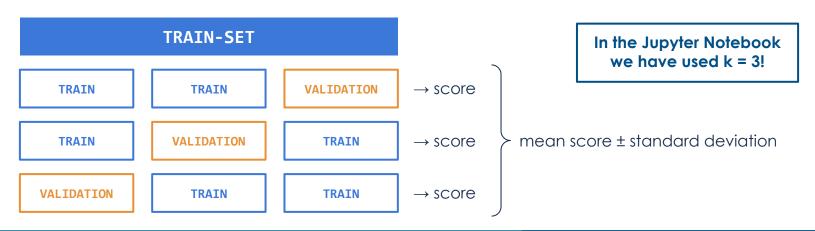
- training set to train the model
- test set to test the model

The error rate on new cases is called the **generalization error**, and we can estimate it evaluating the trained model on the test set.

Model optimization: k-Folds Cross Validation

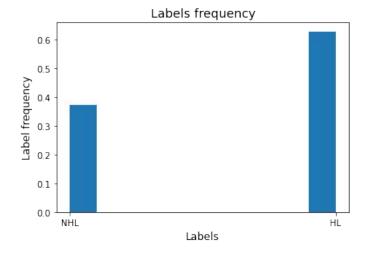
In the search of the best possible predictor, we should avoid that our model <u>learns</u> by heart the training instances, because it will result in a **greater generalization error**.

A possible solution is to use **k-Folds Cross Validation**, an algorithm based on the splitting of the training set into k different subsets. We will use k-1 subsets to train the model and leave the last subset to validate it. Then, the average performance of our model against each of the folds can be used as a robust score to build an **optimization problem**, namely to find the best possible model.



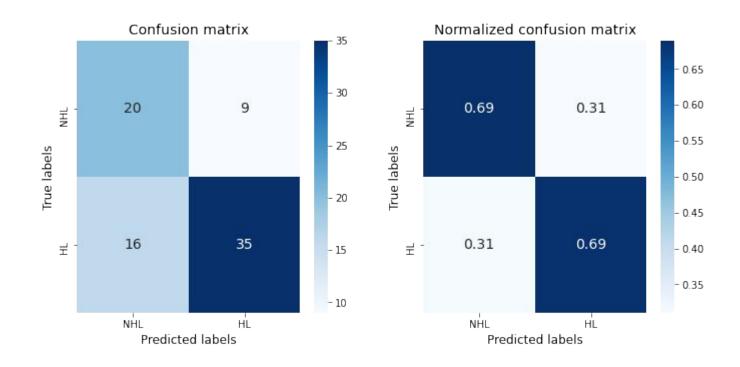
Dummy Classifier: model baseline

- The **Dummy Classifier** is a simple classifier that makes predictions using simple rules:
 - o computes the frequency of each label
 - o interprets such frequencies as <u>probabilities</u>
 - predicts new instances label on the basis of computed probabilities
- This classifier is useful as a simple baseline to compare with other (real) classifiers

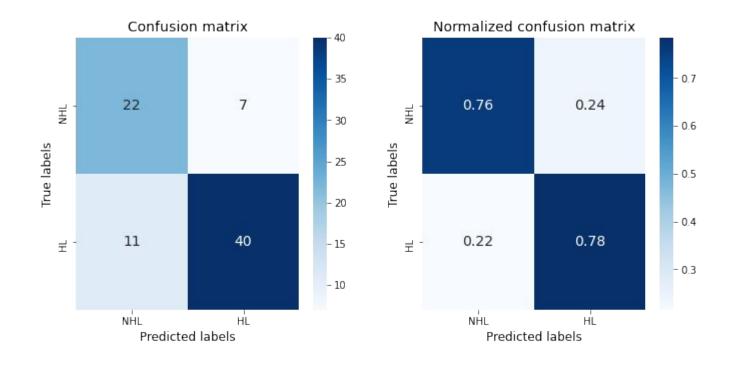


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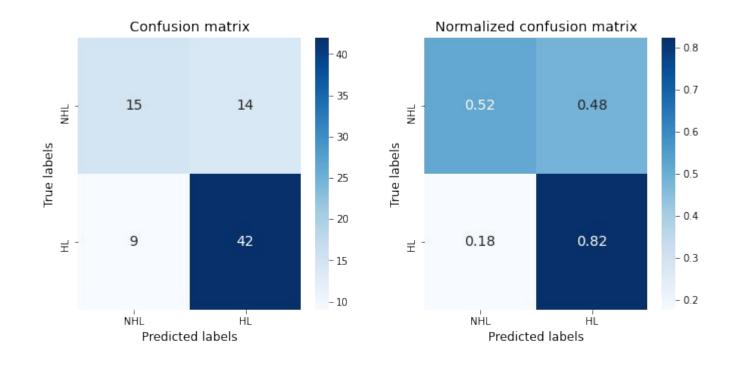
Logistic regression: performance



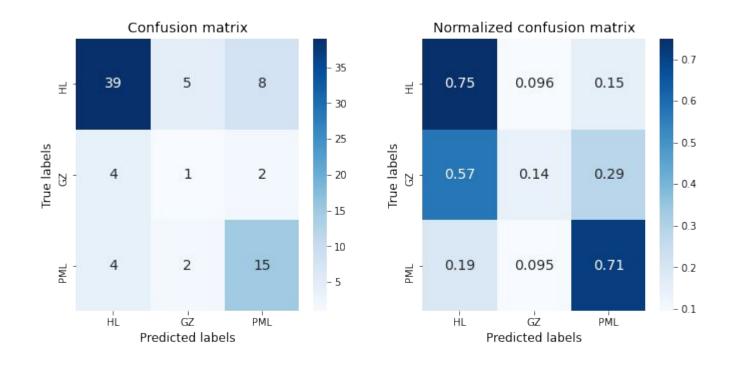
Random Forest classifier: performance



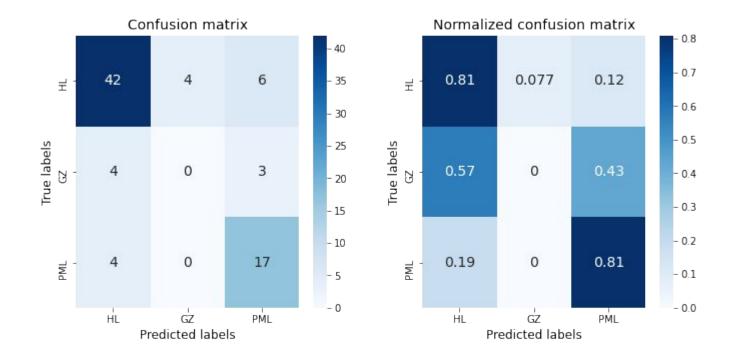
Gradient BDT classifier: performance



Logistic regression: performance



Random Forest classifier: performance



Gradient BDT classifier: performance

