



Dipartimento di Fisica e Astronomia "Ettore Majorana"

RADIOMICS ANALYSIS OF 11C-METHIONINE PET/CT IMAGES THROUGH A MACHINE LEARNING APPROACH

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INTRODUCTION DEFINITION OF RADIOMICS:

RADIOMICS

Radiomics is a new multi-disciplinary science that aims to quantitatively analyze a diagnostic image, through the extraction of features with a statistical and informatic approach, to provide information on the pathophysiology of the underlying tumor, which cannot be qualitatively appreciated by the human eye.

WORKFLOW IN RADIOMICS



Radiomics and radiogenomics in lung cancer: A review for the clinician

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CATEGORIES OF FEATURES

SHAPE FEATURES

STATISTICS FEATURES

TEXTURAL FEATURES

AIM OF THE RESEARCH

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Is to identify a relevant prognostic model concerning a real clinical problem, in a dataset of 56 patients with brain tumors underwent 11C-methionine-PET exams, to discriminate between low- and high-grade tumors.





11C-MET 1



Data Acquisition: The images presenting the required conditions was acquired by different Tomographs:

Tomographs	Number of images	LOW GRADE "0"	HIGH GRADE "1"
ALL	56	17	39
GE DISCOVERY 690	32	9	23
Siemens BIOGRAPH HORIZON	24	8	

LifeX Software:

Local Image Features Extraction — LIFEx — C. Nioche, F. Orlhac, I. Buvat

T = 40% SUVmax

LIFEx version 6.nn, Last update of document: 2019/12/26

• Visualization of the image:

Pre-processing: 0-64 intensities / 0-20 SUV

- Segmentation of the volume of interest: Threshold method = $\begin{cases} tumor & if \quad f(x,y) \ge T \\ non \ tumor & if \quad f(x,y) < T \end{cases}$
- Extraction of features (44):
 - 4 Shape features
 - 4 Conventional features
 - 5 Histogram features
 - 31 Textural features
- The files acquired for each patient were two:
- 1. The extraction of features from the segmented volume (Threshold ROI).
- 2. A ROI of 81 voxels was designed on the SUV max, to eliminate dependence on the segmented volume (Fixed ROI).

EXPERIMENTAL METHOD



First Order	Shape SHAPE_Sphericity SHAPE_Compacity SHAPE_Volume_mL SHAPE_Volume_vx	Histogram HISTO_Skewness HISTO_Kurtosis HISTO_Entropy_log10 HISTO_Entropy_log2 HISTO_Energy		CONV_(min, mean, sd, max) CONV_peak 0.5mL CONV_peak 10.5mL CONV_peak 1mL CONV_TLG (for TP, MN)	
Second Order	GLCM vx : distance with neighbours v GLCM_homogeneity v GLCM_Energy v GLCM_Contrast v GLCM_Correlation v GLCM_Entropy v GLCM_Dissimilarity	GLRM CGLRM_GRE GLRM_LGRE GLRM_GRE GLRM_SRLGE GLRM_SRLGE GLRM_SRLGE GLRM_SRLGE GLRM_GLNU GLRM_GLNU GLRM_RP		GLDM GGLM_Coarseness IGLDM_Contrast IGLDM_Busyness	GLZLM GLZUM_LCE GLZUM_LCZE GLZUM_LCZE GLZUM_SZLGE GLZUM_SZLGE GLZUM_SZLGE GLZUM_LZLGE GLZUM_LZLGE GLZUM_ZLNU GLZUM_ZLNU GLZUM_ZLNU GLZUM_ZP
Output	ROI Output	Histogram Out. Show histogram Save histogram in fi	ie	Matrix Output- Save GLCM in file in each direction Save GLRLM in file Save NGLDM in file Save GLZLM in file	



DATA ANALYSIS



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Proposed Method of Features Selection

- For each column of matrix X, calculate the point biserial correlation (pbc) index between 1. variables x (features) and dichotomic variable y.
 - \bar{x}_{1} be the corresponding averages the values

of x which are paired with y = 1 (n₁)

 \bar{x}_{0} be the corresponding averages the values of x which

are paired with y=0 (n₀)

 \bar{x} the total average (n)

 $pbc = \frac{\sqrt{\frac{n_1 n_0}{n} (\bar{x}_1 - \bar{x}_0)}}{\left[\sum_{i=0}^{1} \sum_{i=1}^{n_i} (x_{ii} - \bar{x})^2\right]^{1/2}}$

- 2. Sort the X columns in descending order of absolute value of point biserial correlation index
- 3. Start a WHILE-DO cycle, and it is done a logistic regression analysis, through function 'FITGLM' in Matlab, where one of the outputs is the p-value:

WHILE (p-value decreases) Add regressor column and perform logistic regression DO Add last regressor column

Relevant Features						
TOMOGRAPH	FEATURES	p-VALUE	TOMOGRAPH	FEATURES	p-VALUE	
ALL	NGLDM	0.1615	ALL	SHAPE Sphericity	0.0314	
	Busyness	0.3207		SHAPE Compacity	0.0215	
	GLZLM LZLGE			HISTO Kurtosis	0.0232	
GE	GLRLM LRLGE	0.05	GE	GLRLM LRLGE	0.0481	
	GLZLM LZLGE	0.137		GLRLM LGRE	0.117	1
SIEMENIS	UISTO Skownoog	0.0126	OTEMENIC		0.00026	5
SIEWIENS		0.0130	SIEMENS	GLCM Correlation	0.00036	
	CUNV SUV	0.0136		SHAPE Compacity	0.0014	
	Fixed ROI					

DATA ANALYSIS

Machine Learning Model: Discriminant Analysis

- Classification: the process of predicting a qualitative response Y, starting from a set of predictors X
- Supervised learning: use labeled data (to map the data to the desired output classification)

Training set:

a set of examples, represented as a pair of an input vector (features) and a desired **–** output value (target or category label).

K-fold Cross Validation



K=5

Classifier: the classification algorithm analyzes the training data and infers a hypothesis (function), which can be used to predict new outputs. In matlab the function "mdl = fitcdiscr(X,Y)" returns a discriminant analysis classifier based on the input variables X and response Y.

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Validation set:

The performance of a model is estimated on a validation sample

Each time, one of the k subsets is used as the validation set, and the other k-1 subsets are put together to form a training set.

This step is useful in the case of a small mount of data.

DATA ANALYSIS

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Performance Evaluation



1-Specificità

RESULTS

Threshold ROI

Fixed ROI

TOMOGRAPH	SENSITIVITY	SPECIFICITY	ACCURACY Error	TOMOGRAPH	SENSITIVITY	SPECIFICITY	ACCURACY Error
ALL	52.44 %	76.62 %	70.31 % 29.69%	ALL	41.17 %	63.60 %	57.25 % 42.75%
GE	71.76 %	83.76 %	80.51 % 19.49%	GE	28.52 %	88.47 %	71.64 % 28.36%
SIEMENS	86.67 %	84.86 %	84.98 % 15.02%	SIEMENS	76.67 %	71.81 %	72.88 % 27.12%

- 1. Our workflow achieves, using an innovative feature selection method, good performance, in terms of accuracy, which is greater than 70.31% and specificity, which is greater than 71.81%, in all cases except when we consider both scanners using the fixed ROI.
- 2. Higher performance values are found for the data set of threshold ROI. We can conclude that there are no particular advantages in using a homogeneous region of interest (e.g. fixed ROI) rather the a variable ROI among patients (e.g. thresholding method).

RESULTS

0.9

0.8

0.7

0.0 gt

5 0.5

Lrue

0.3

0.2

0

ROC Curves for Discriminat Analysis Classification

0.4 0.6 0.8

TOMOGRAPH

SIEMENS

False positive rate

AUC

(95%CI)

78.91%

58.80/99.01%

ROC Curves for Discriminat Analysis Classification

0.2

9

HISTOskewness

- Reference Line

Predicted Probability

CONVENTIONAL_BIM_SUV_volume (# mL)1

GLCM_orrelation

ROC CURVE - FIXED ROI









CONCLUSIONS

• Low-performance values are noted for the AUC (< 78.91%): The main cause is the low number of patients. In addition, the patient dataset is unbalanced. For better performance, a machine learning algorithm should work with the same number of "objects" for both classes. The group that shows the most symmetry is the one related to Siemens, in fact, in all the results we notice higher values.

• In all cases lower values are noticed when the scanners are united in the same group "ALL". This is a particular limitation: radiomics features are affected by acquisition parameters, reconstruction algorithm, slice thickness, voxel size, thus hampering multicenter studies.

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Radiomics is a new discipline and for this reason it needs standardization and future studie