



School on Medical Physics for
Radiation Therapy:
Dosimetry and
Treatment Planning for Basic and
Advanced Applications



25 March – 5 April 2019
Trieste, Italy

Further information:
Activity URL: <http://indico.ictp.it/event/8651/smr3278@ictp.it>

Methods for radiomics analysis

Michele Avanzo
Medical Physicist
Centro di Riferimento Oncologico
IRCSS Aviano (PN) mavanzo@cro.it

ICTP Trieste 4/2/2019

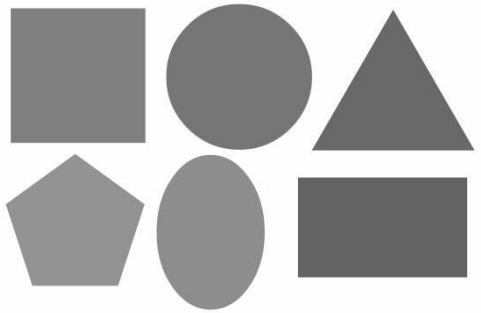
“Images are more than pictures, they are data”



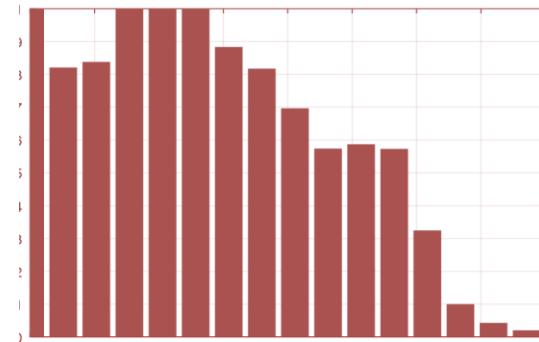
1	3	4	5	3	2	2	1
1	2	3	4	3	3	2	3
1	1	2	3	3	3	3	4
1	1	1	2	2	2	2	3
1	1	0	1	1	1	1	2
0	0	0	1	2	0	0	1
0	0	0	0	0	0	0	1
0	0	0	0	0	0	0	0

Radiomic features

Shape



Histogram (1st Order)



Textural (2nd order)



Higher
order



Radiomic features

Shape

$$compactness\ 2 = 36\pi \frac{A^2}{V^3}$$

Histogram (1st Order)

$$kurtosis = \frac{\frac{1}{N} \sum_i (X(i) - \bar{X})^4}{\left(\frac{1}{N} \sum_i (X(i) - \bar{X})^2 \right)^2}$$

$$entropy = \sum_i (P(i) \log_2 P(i))$$

Textural (2nd order)

$$autocorrelation = \sum_{i,j} i * j * P(i, j)$$

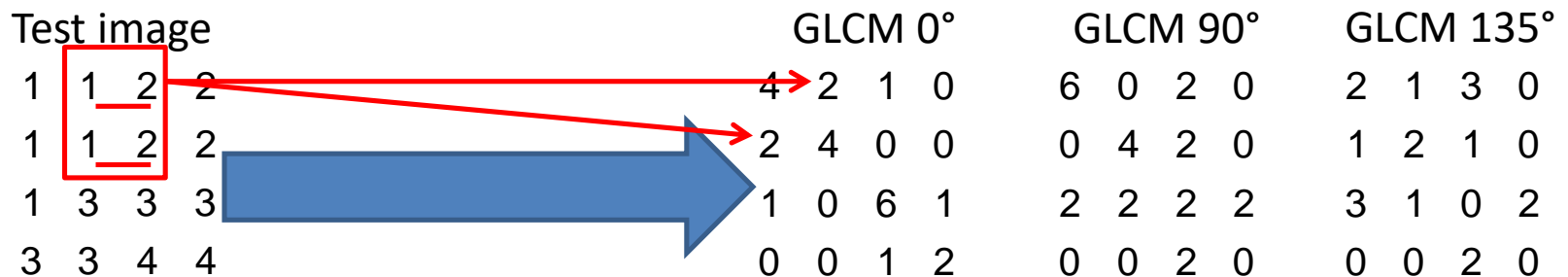
$$cluster\ shade = \sum_{i,j} (i + j - 2\mu)^3 * P(i, j)$$

Higher order

$$coarseness = \frac{1}{\varepsilon + \sum_i P(i)s(i)}$$

Textural features

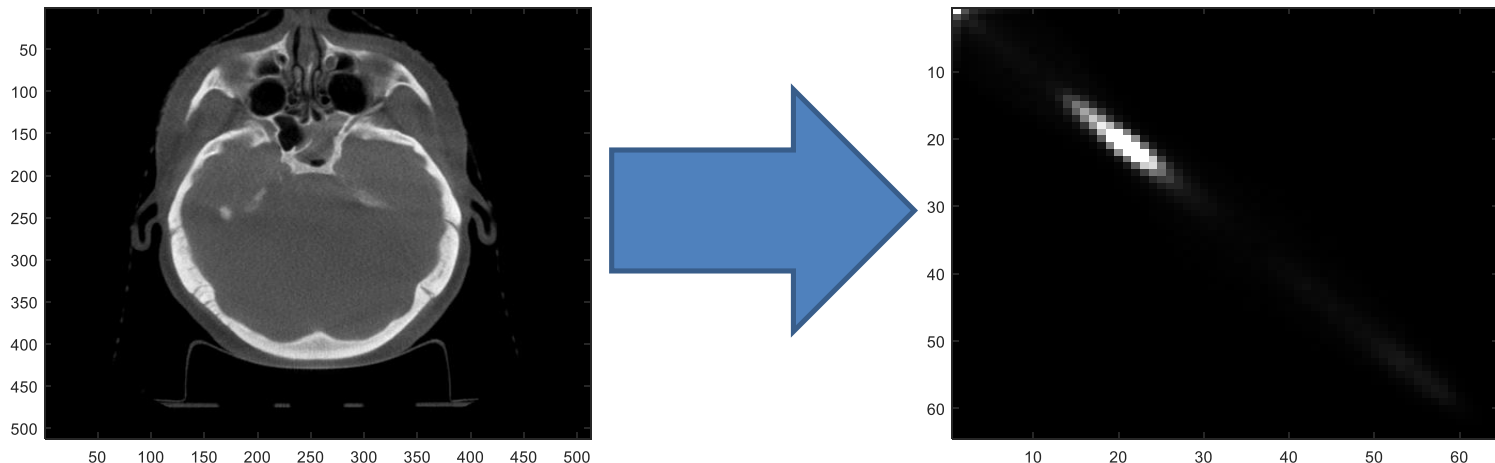
- The gray-level co-occurrence matrix (GLCM) is a matrix whose row and column numbers represent gray values, and the cells contain the number of times corresponding gray values are in a certain relationship (angle, distance).



GLCM with distance one pixel along directions 0°, 90°, 135°

Textural features

- The gray-level co-occurrence matrix (GLCM) is a matrix whose row and column numbers represent gray values, and the cells contain the number of times corresponding gray values are in a certain relationship (angle, distance).



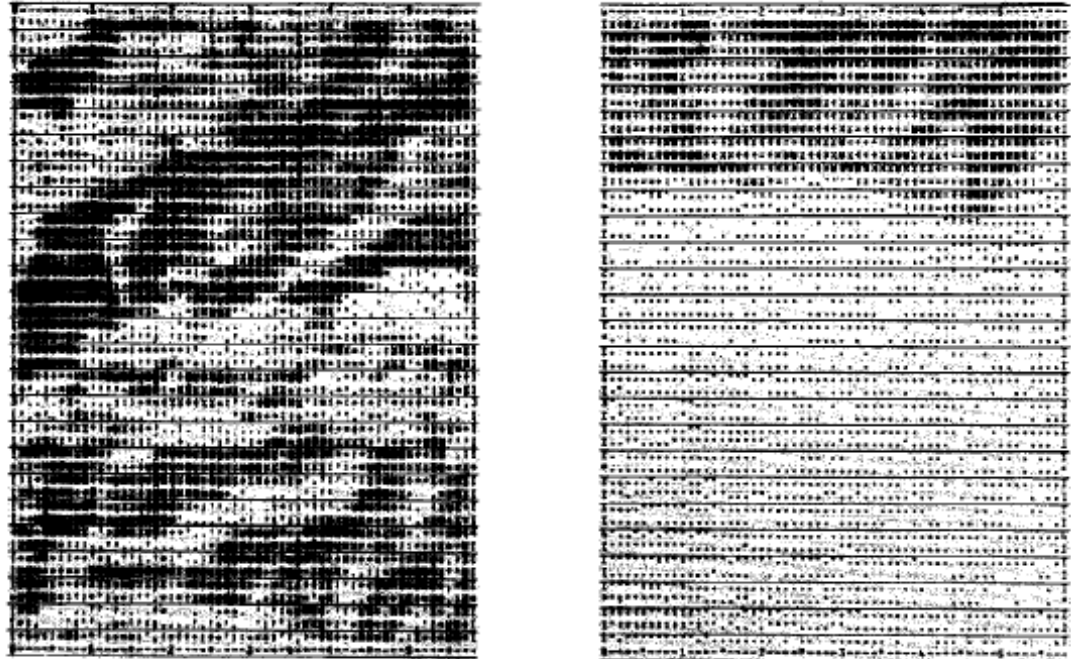
$autocorrelation = \sum_{i,j} i * j * P(i, j)$ represents the correlation of the image along the specified direction

$$cluster\ shade = \sum_{i,j} (i + j - 2\mu)^3 * P(i, j)$$

$P(i, j)$ = element of GLCM, μ = average of GLCM

When were features born?

- GLCM



Grassland				Water Body		
Angle	ASM	Contrast	Correlation	ASM	Contrast	Correlation
0°	.0128	3.048	.8075	.1016	2.153	.7254
45°	.0080	4.011	.6366	.0771	3.057	.4768
90°	.0077	4.014	.5987	.0762	3.113	.4646
135°	.0064	4.709	.4610	.0741	3.129	.4650
Avg.	.0087	3.945	.6259	.0822	2.863	.5327

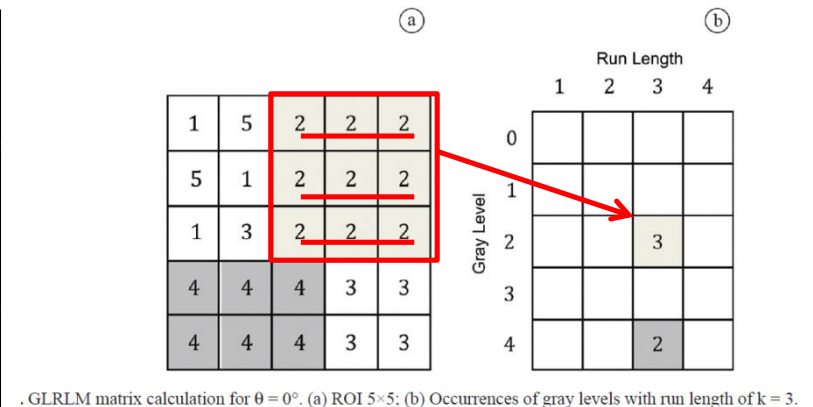
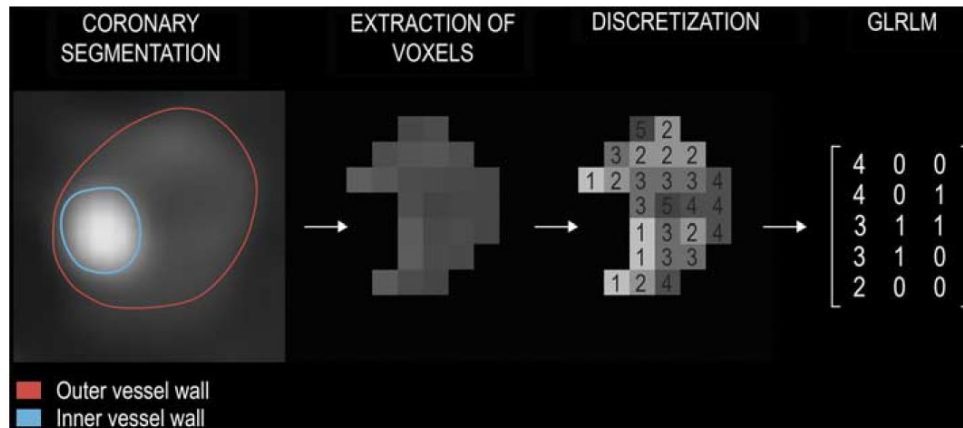
(a)

(b)

Fig. 4. Textural features for two different land-use category images.

Textural features

- Gray Level Run Length Matrix (GLRLM) is a two-dimensional matrix in which each element describes the number of times j a gray level i appears consecutively in the direction specified



Higher order variables

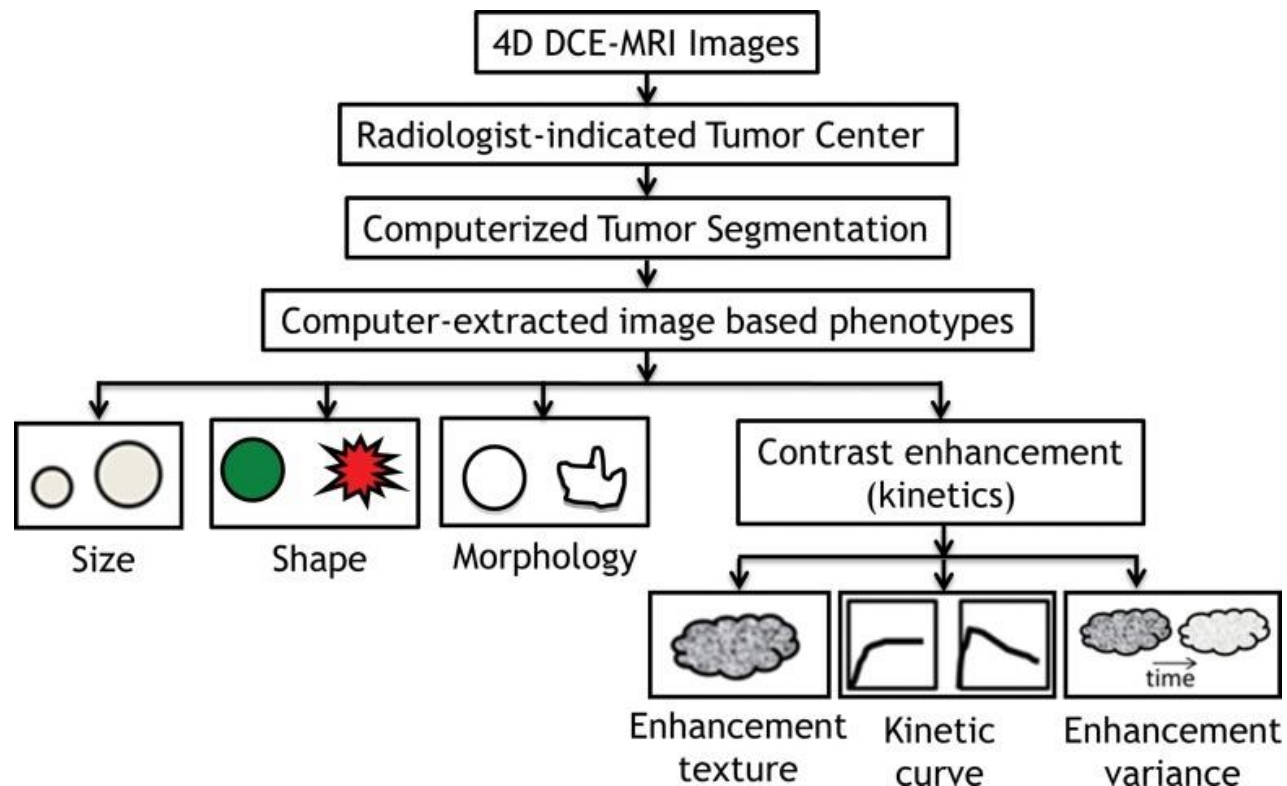
- In the neighborhood gray-tone difference matrix (NGTDM), the i th entry is a summation of the differences between all pixels with gray-tone i and the average value of their surrounding neighbors

Image				
3	2	0	1	0
1	2	1	3	0
3	1	0	2	3
1	2	3	0	3
0	0	0	0	1

NGTMD	
j	$S(j)$
0	3.25
1	1.00
2	2.00
3	4.25

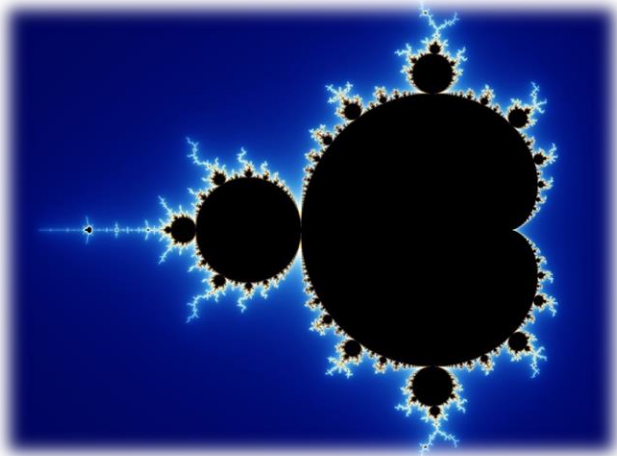
Kynetic variables

- Pharmacokinetics (uptake rate of contrast agent, washout...)
- Evolution in time of radiomic features in 4D DCE-MRI



Other features

Fractal

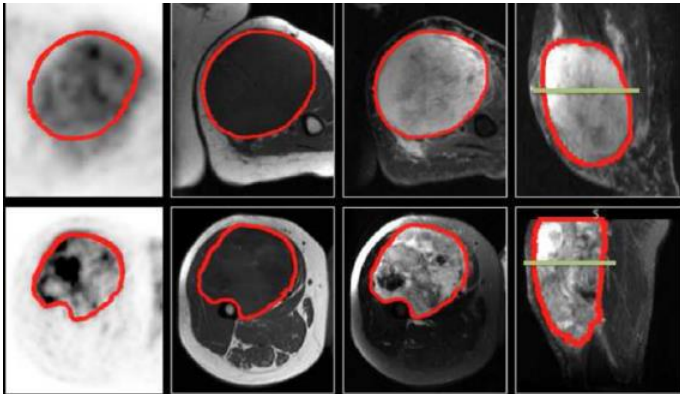


Hausdorff's fractal dimension refers to self-repeating textures of a pattern as one magnifies the feature:

$$D_0 = -\lim_{\varepsilon \rightarrow 0} (\log_{\varepsilon} N(\varepsilon)) = \lim_{\varepsilon \rightarrow 0} \frac{\log(N(\varepsilon))}{\log(\varepsilon^{-1})}$$

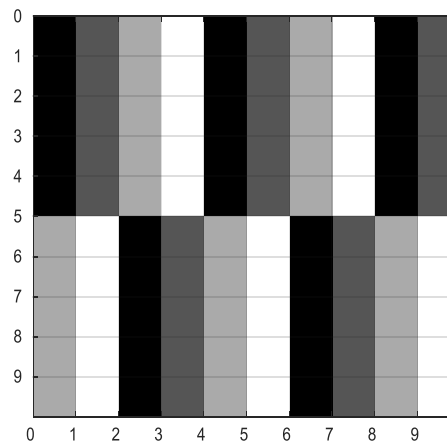
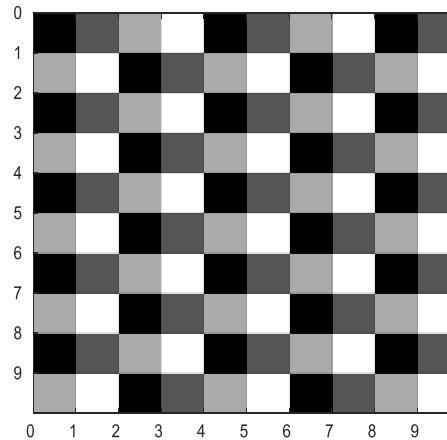
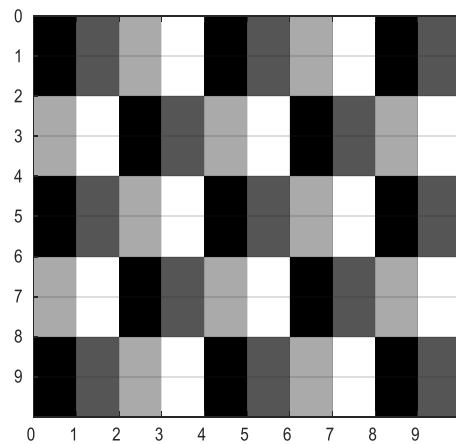
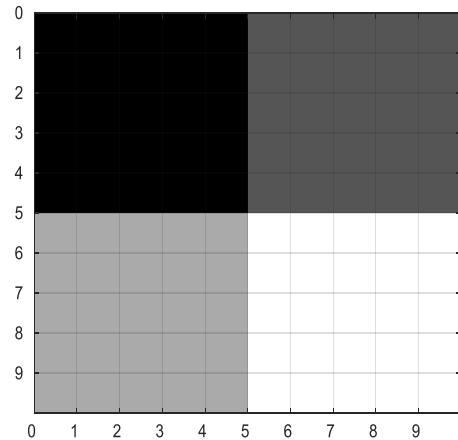
where $N(\varepsilon)$ is the number of $\varepsilon \times \varepsilon$ squares needed to cover the 2D area.

Fusion

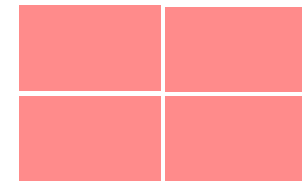


Wavelet discrete transform can be used to fuse images. The weight of wavelet bands in fusion can be used as a feature

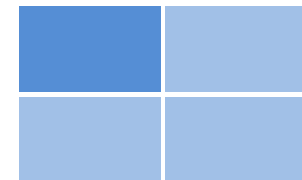
Radiomic features



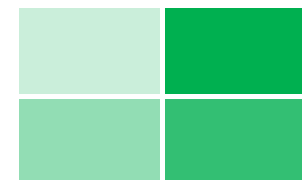
Histogram (1st Order)



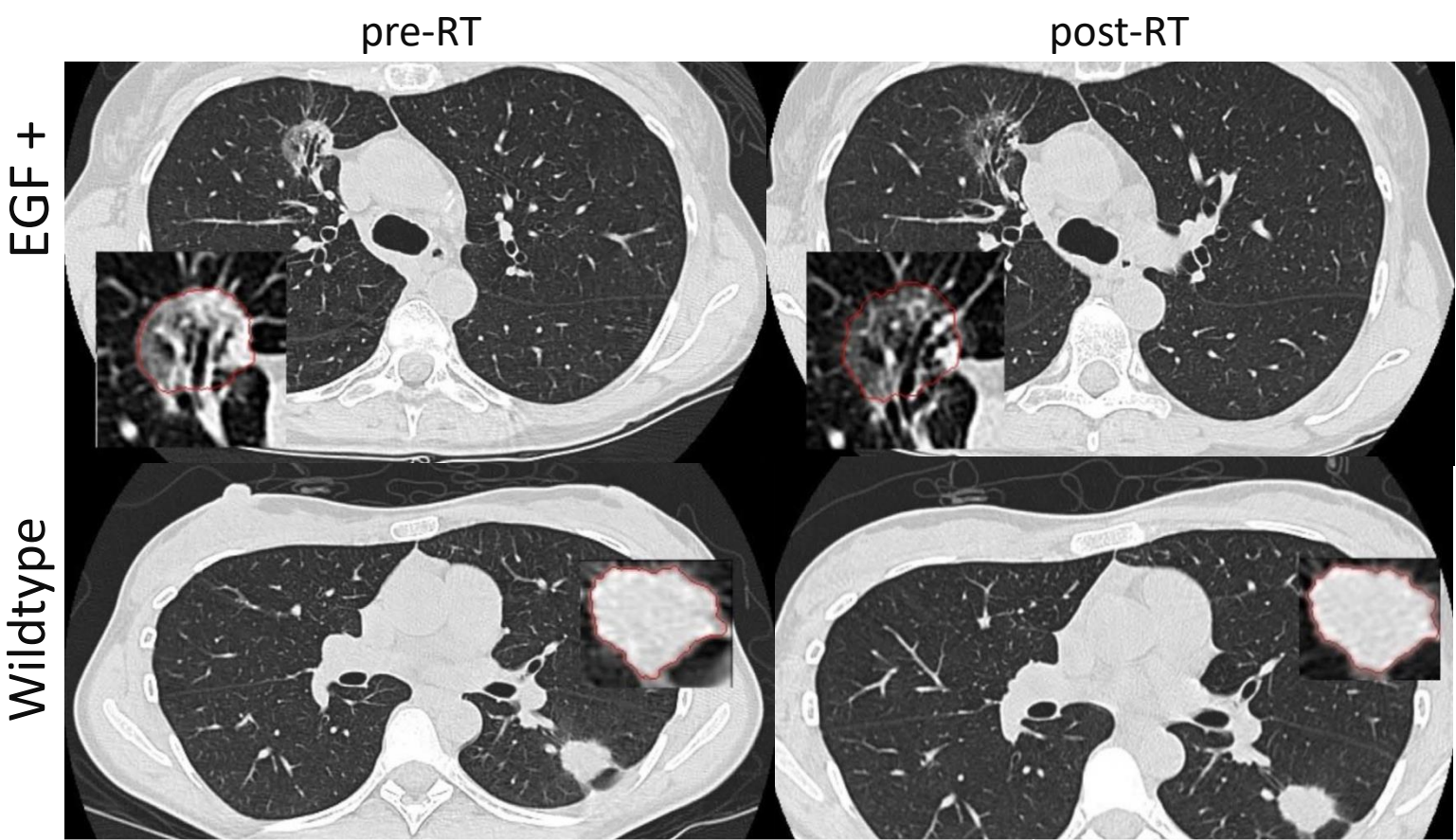
Textural (2nd order)



Higher order



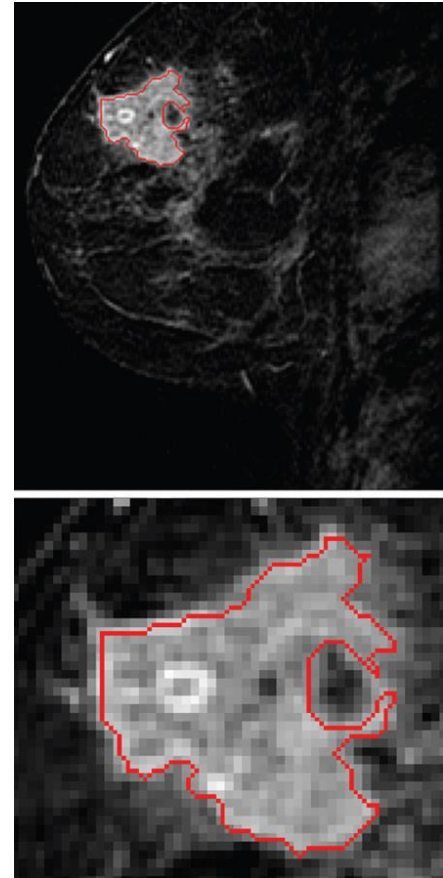
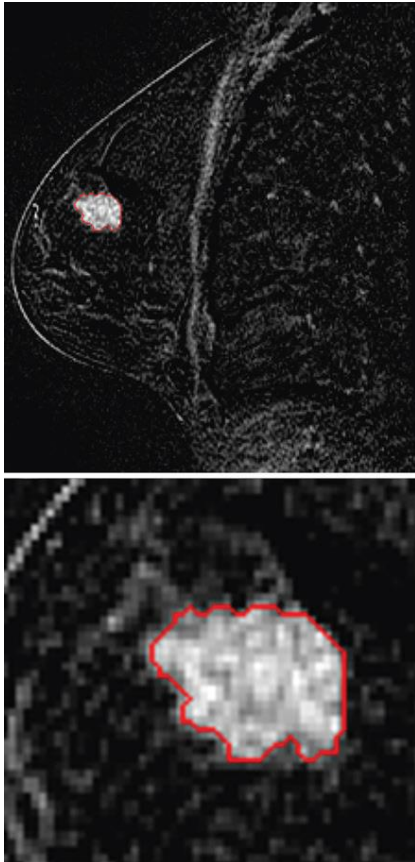
Radiomic features vs EGF mutation status



EGFR status	CT acquisition	Volume	Radius_Std	Shape_SI6	Gabor_Energy-dir135-w3	Gabor_Energy-dir45-w9	Laws_Energy-10	Laws_Energy-13
EGFR positive	Baseline (Fig 1-a)	7766.5	1.522	0.145	5337.9	419770.4	475.2	1369.6
	Followup (1-b)	7195.8	1.657	0.151	4043.5	327365.1	512.0	1352.9
	Change	-570.6	0.135	0.006	-1294.4	-92405.3	36.8	-16.6
Wild type	Baseline (Fig 1-c)	3502.4	1.422	0.173	11601.7	419578.9	367.7	353.9
	Followup (1-d)	4522.8	1.251	0.165	10605.5	361191.5	326.3	349.3
	Change	1020.4	-0.171	-0.009	-996.2	-58387.4	-41.5	-4.5

Breast Cancer

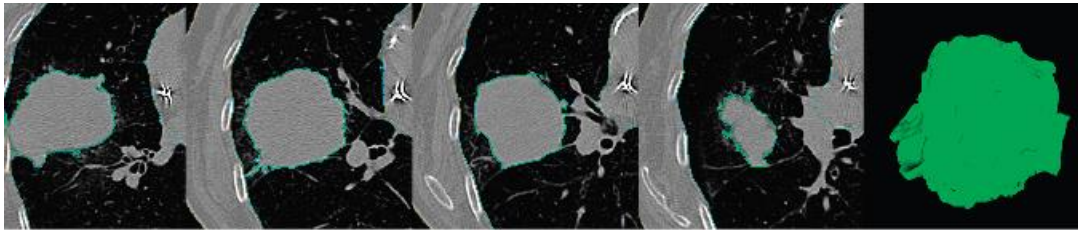
ER, PR, positive, HER2 negative, stage II
invasive breast cancer, good prognosis.



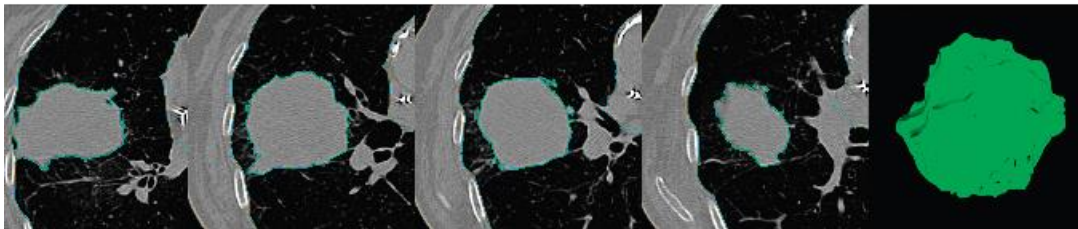
ER, PR, HER2 negative, stage II invasive
breast , poor prognosis

Reproducibility (Test-retest)

- Measured from repeated measurements on same conditions



First baseline scan



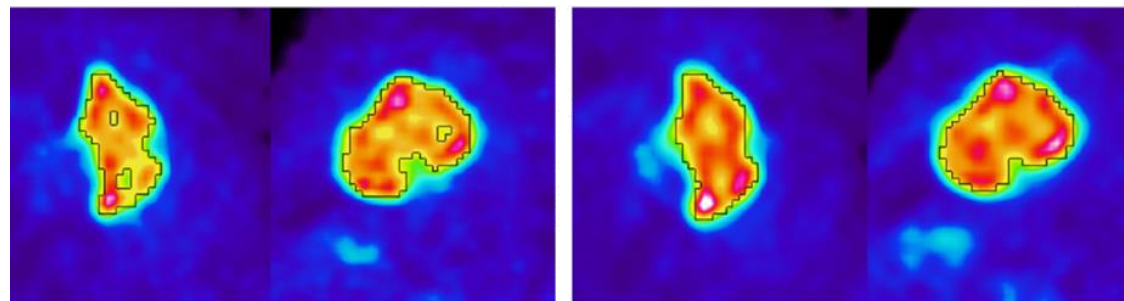
Second baseline scan



(A) 93(42.4%) over 219 features were stable (Concordance
(B) Correlation Coefficient above 0.85) respectively in the
RIDER dataset

Textural features are more reproducible with respect to maximum and mean SUV.

63% of features stable (Intraclass correlation coefficient > 0.9)

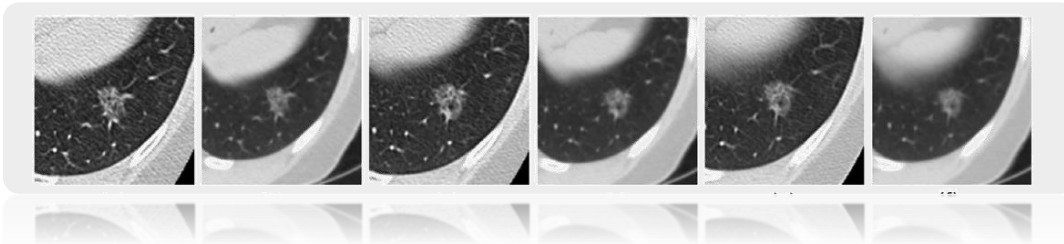
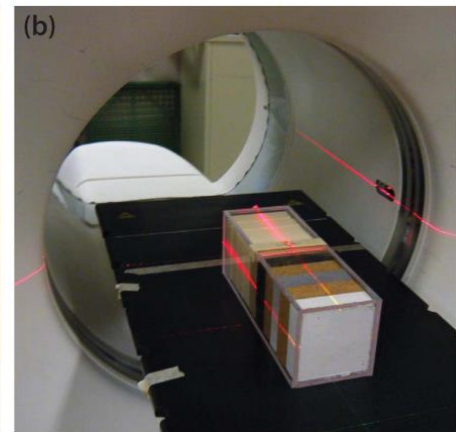
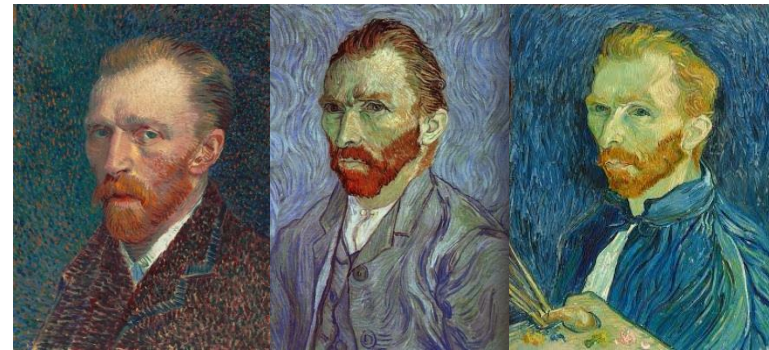


Robustness: CT

- Robustness is variability with changing conditions (e.g. reconstruction parameters, scanner, patient position)

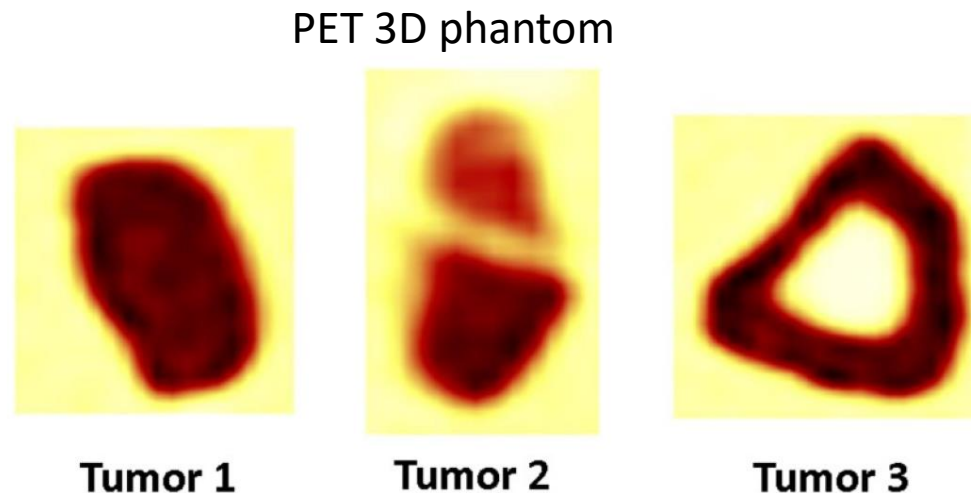
Radiomic features from CT are sensitive to:

- Scanner
- Slice thickness
- reconstruction algorithms
- Segmentation



Robustness: PET

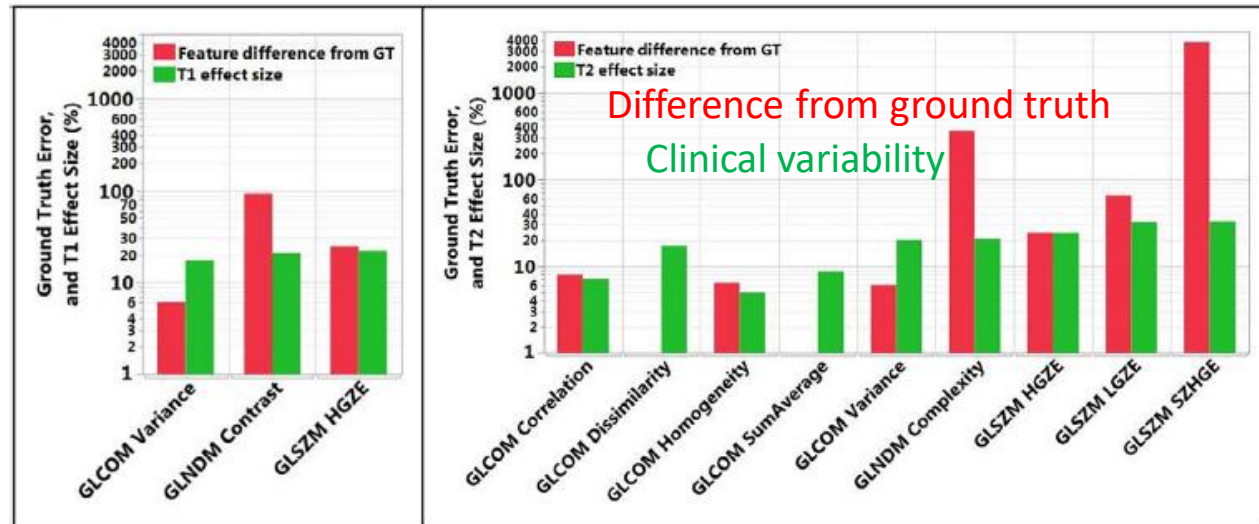
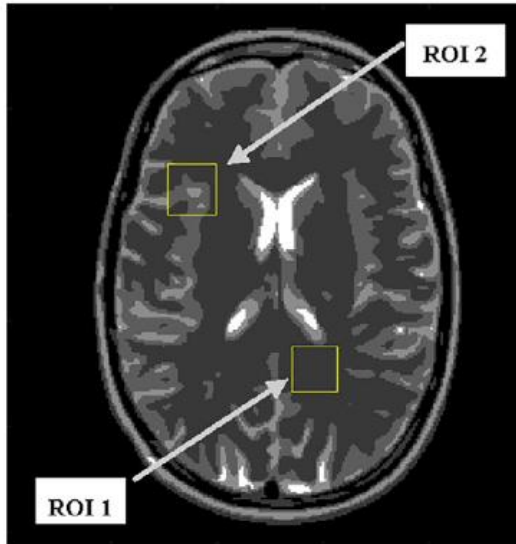
- Image reconstruction algorithm (OSEM, TOF, PSF, PSFTOF)
- The method of quantization or discretization, where voxel intensities are grouped into equally spaced bins, also affects reproducibility
- Scan duration (\approx noise)
- Segmentation



Robustness: MRI

- Radiomic features extracted from MRI scans depend on the pulse sequence, field of view, field strength, and slice thickness
- Effect of reconstruction (iterative vs non iterative) algorithm is small

a) Digital ground truth phantom used as input to a MRI simulator in Matlab.



Which are the most stable features?

	FIRST ORDER	SHAPE METRICS	TEXTURE ANALYSIS	COMMENTS
ROI SEGMENTATION				
MANUAL DELINEATION	◆	◆◆◆	◆◆◆	Mainly PET studies and one multi-center CT study. Shape metrics from PET may be less subject to inter-observer differences. Semi-automated methods generally improve reproducibility.
SEMI-AUTO / AUTO	◆	◆◆	◆◆	
IMAGE RECONSTRUCTION				
RECONSTRUCTION FILTER	◆	◆◆	◆◆◆	Consistent in a few CT and PET studies of NSCLC.
VOXEL SAMPLING	◆◆	◆◆	◆◆◆	
IMAGE ACQUISITION SETTINGS				
RESPIRATORY MOTION	◆◆	◆◆	◆◆	Consistent over single-institution PET and CBCT studies of NSCLC.
SCATTERED RADIATION	◆◆	?	◆◆	In one CBCT study of NSCLC, but did not evaluate shape metrics.
CT SCANNER	◆◆	◆◆	◆◆	In one multi-institutional CT study in NSCLC, effects were similar in magnitude to inter-patient differences.
DIGITAL IMAGE PRE-PROCESSING				
NOISE AND SMOOTHING	◆◆	?	◆◆	Single-center CBCT and planning CT study in H&N; smoothing and noise have less effect than high-pass and logarithmic filters.
INTENSITY DISCRETIZATION	◆◆	◆◆	◆◆	Consistent in H&N studies of perfusion CT and PET, bin size may have less impact in PET.
CONSENSUS ABOUT MOST STABLE OR LEAST STABLE RADIOMIC FEATURES	<p>Entropy was consistently among the most repeatable/reproducible first-order features. There were inconsistent findings for skewness and kurtosis.</p> <p>Certain shape metrics may be reproducible in PET, and slightly less reproducible in CT, though it is unclear which individual features prove to be stable.</p> <p>No emergent pattern or consensus for highly reproducible textural features. Coarseness and contrast were among the least reproducible.</p>			

◆ less likely ◆◆ probable ◆◆◆ highly likely influenced by parameters

Good repeatability is a necessary, but not sufficient condition for high predictive power of a feature,

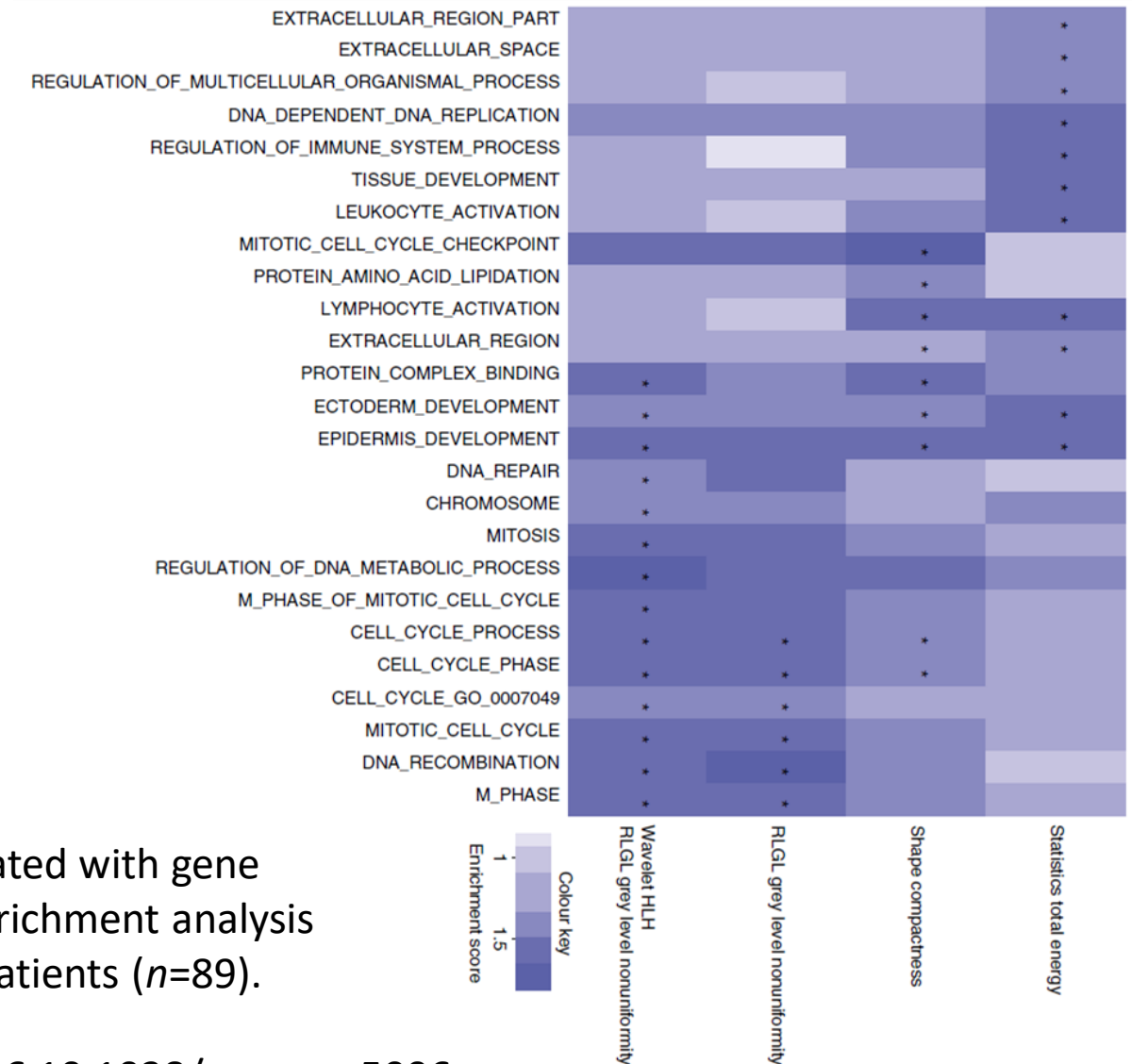
If a feature has a low repeatability, its predictive power must be low, too

If a feature has a good repeatability, we cannot conclude anything about its predictive power

Radiomics and biology

- Radiomic features provide a description of the appearance of the tumor in the medical image
- Medical images are not the tumor, but a representation, but...
- ...in biopsy-based assays, the extracted sample does not always represent the entire population of tumor cells, and...
- radiomic features assess the comprehensive three-dimensional tumor bulk by means of imaging information

Radiomics and biology



Radiomic features are associated with gene expression using gene-set enrichment analysis (GSEA) in a data set of lung patients ($n=89$).

Radiomics and biology

- Tumor histology (squamous cell carcinoma, large cell carcinoma, adenocarcinoma and “not otherwise specified”)

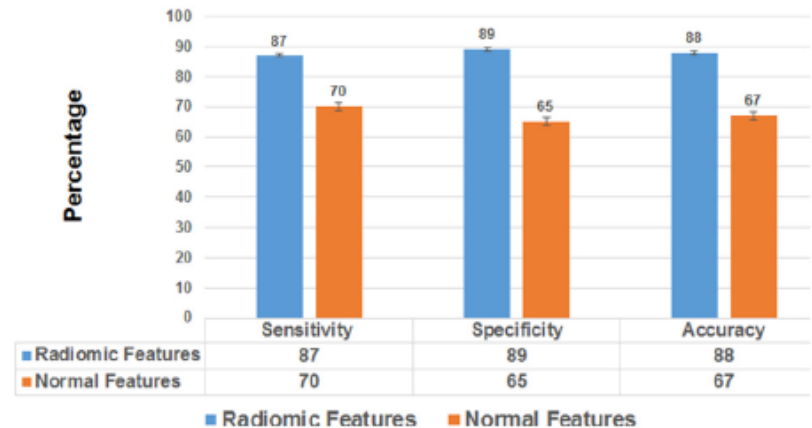


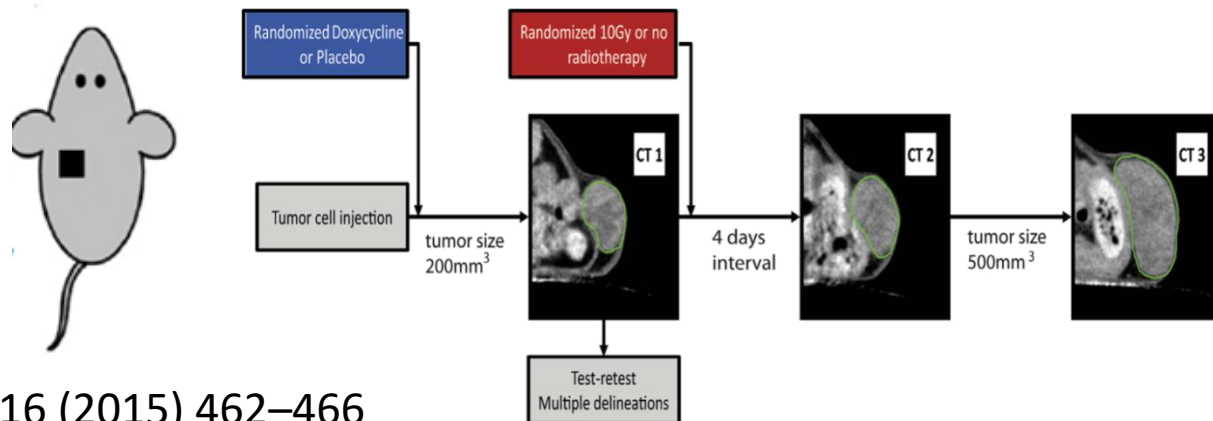
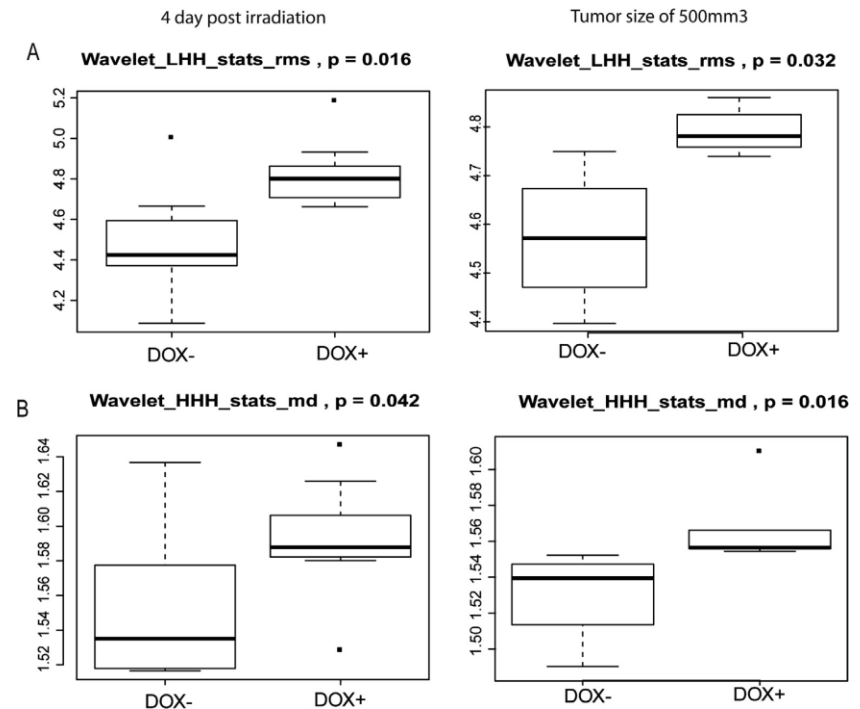
Figure 2. Classification metrics using radiomic and normal features.

- *ALK/ROS1/RET* fusion-positive tumor
 - younger age, advanced tumor stage, solid tumor on CT, SUV_{max} tumor mass, kurtosis and variance
 - sensitivity and specificity, 0.73 and 0.70, respectively.

Patil, Tomography 2 (4) DECEMBER 2016

Biology and radiomics: causal effect?

- Tumor cells of colon cancer(HCT116, GADD34 inducibili) injected in the flank of nude mice
- Some mice had placebo other received a drug which induces overexpression of gene GADD34 in the tumor
- CT scan was acquired and radiomic features extracted in both cohorts

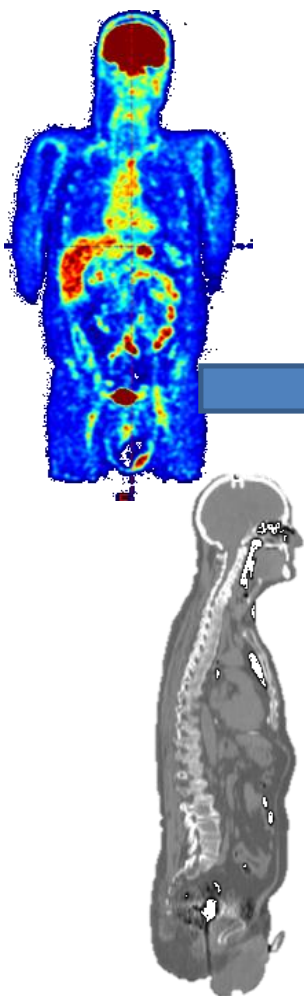


Definition of radiomics

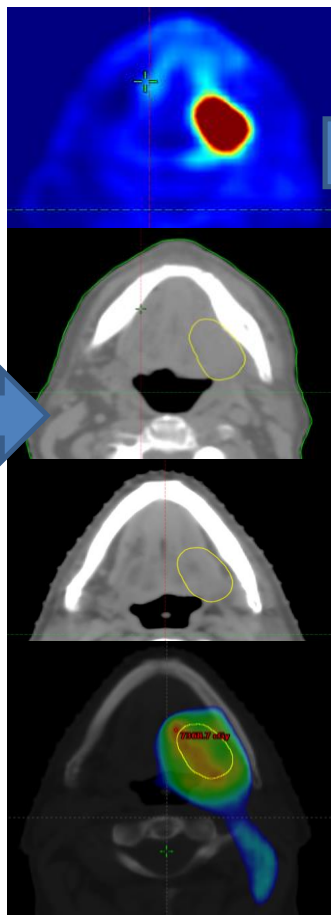
- The term radiomics originates from the words “radio” which refers to radiology, i.e. medical images in the broad sense (CT, PET, MRI, US, mammography etc.), and “omics”, first used in the term genomics to indicate the mapping of human genome, indicating large scale analysis
- The goal of radiomics is prediction of biological or clinical endpoints by:
 - quantitative analysis of tumor/organ at risk through extraction of a large amount of radiomic features
 - use of machine learning for building predicting models

Radiomics: workflow

I. Imaging



II. Contouring



III. Pre- Processing, filtering

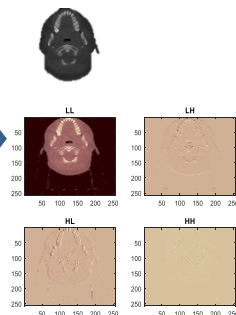
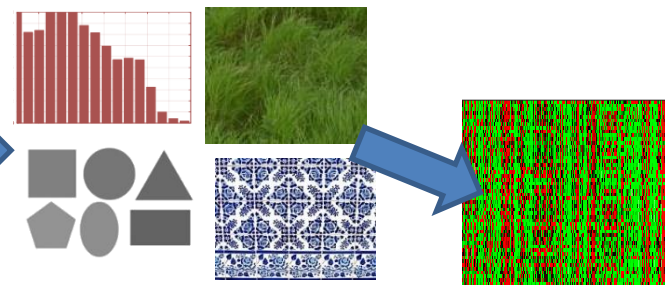
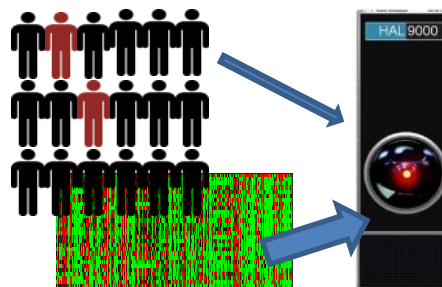


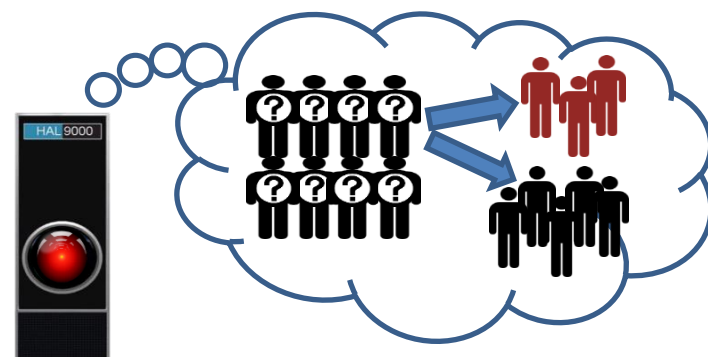
Image features



V. Machine learning



VI. Validation



Pre-processing

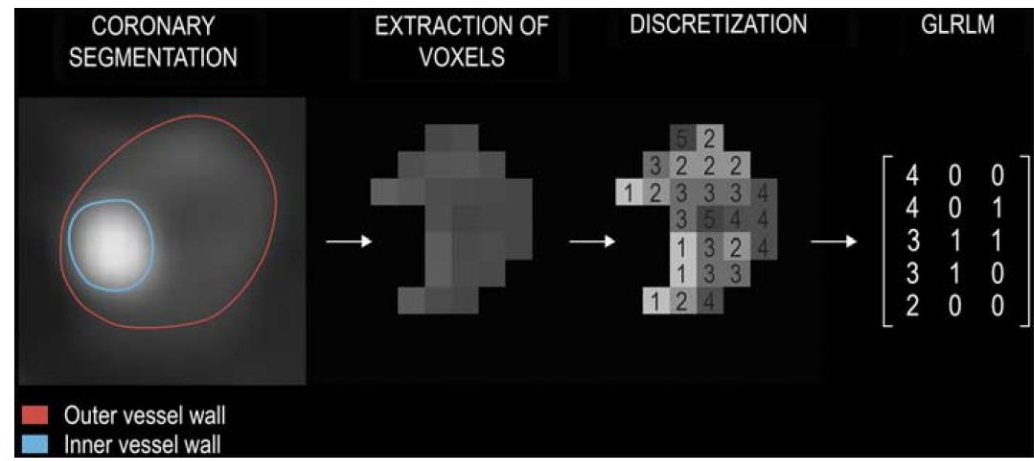
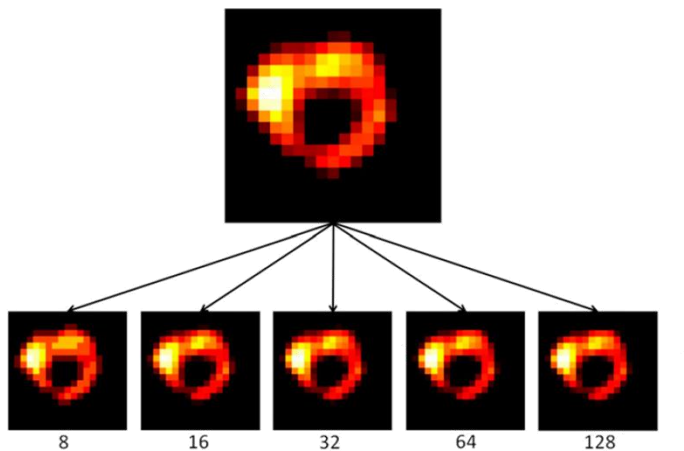
Preprocessing aims at reduce noise and calculation time and to harmonize images of different patients:

1) Discretization of the intensity levels. 2 methods are used: :

- “fixed bin size”, where intensity levels are grouped into bins of fixed size, such as 25 Hounsfield Units nella CT
- “fixed bin number”, where the number of levels are fixed, e.g. 32 or 64

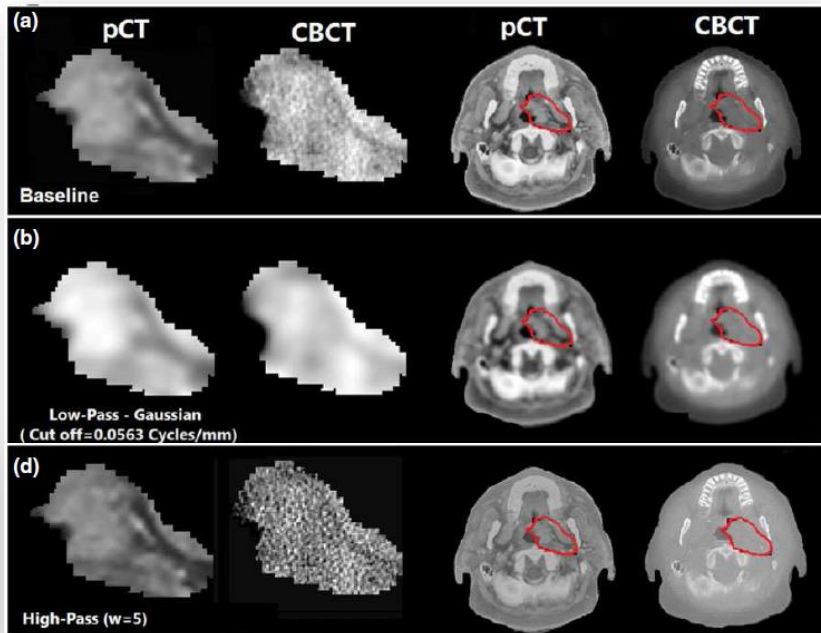
2) Resampling of image into voxels with size e.g. $3 \times 3 \times 3 \text{ mm}^3$.

Interpolation algorithms used: nearest neighbour, trilinear, tricubic convolution, tricubic spline interpolation



Filtration

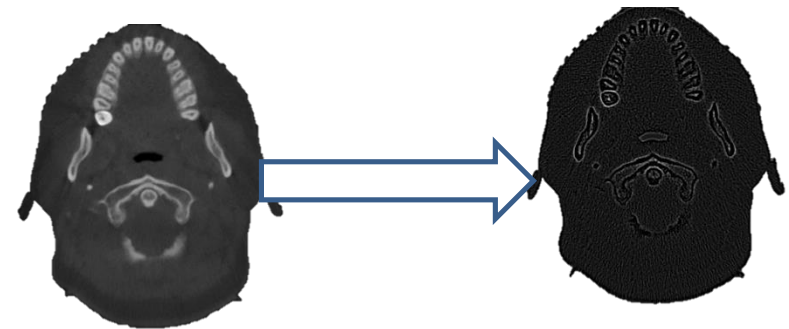
Low-pass and high pass filters:



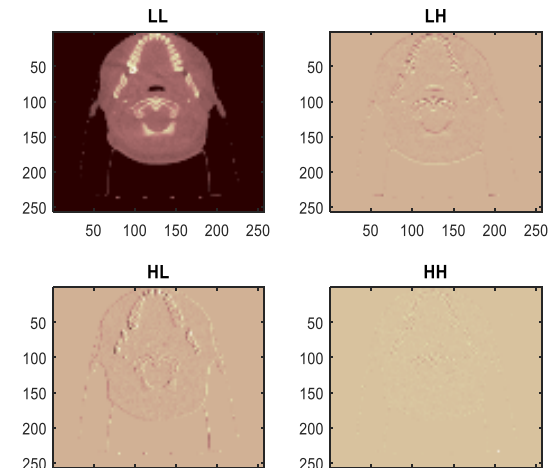
Filter Laplacian of Gaussian (LOG):

$$Log(x, y) = -\frac{1}{\pi\sigma^4} \left[1 - \frac{x^2 + y^2}{2\sigma^2} \right] e^{-\frac{x^2 + y^2}{2\sigma^2}}$$

σ = radius of gaussian



Wavelet Transform 2D:



Definitions of radiomic features

- Some papers report comprehensive formulas of radiomic features:
Kickengereder et al, Radiology 2016;;160845.
Aerts et. Al, NATURE COMMUNICATIONS | 5:4006 | DOI: 10.1038/ncomms5006
- Some inconsistencies in definitions:

Paper 1

$$\text{compactness 2} = 36\pi \frac{V^2}{A^3}$$

Paper 2

$$\text{compactness 2} = 36\pi \frac{A^2}{V^3}$$



Cornell University

[arXiv.org](#) > [cs](#) > [arXiv:1612.07003](#)

Computer Science > Computer Vision and Pattern Recognition

Image biomarker standardisation initiative

Alex Zwanenburg, Stefan Leger, Martin Vallières, Steffen Löck, for the Image Biomarker Standardisation Initiative

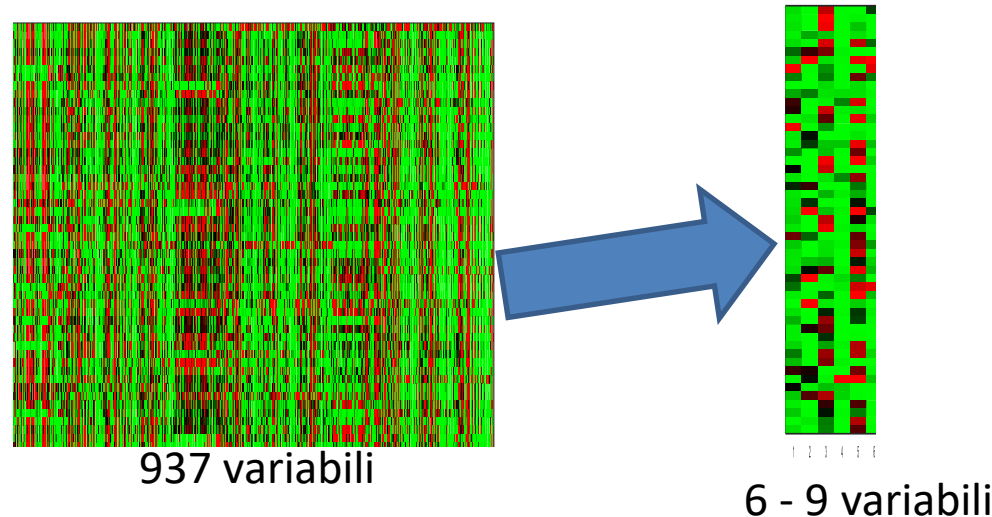
(Submitted on 21 Dec 2016 (v1), last revised 17 Sep 2018 (this version, v7))

Open-source softwares

- ePAD, Stanford University, doi.org/10.1016/B978-0-12-812133-7.00013-2
- PyRadiomics/Radiomics , Harvard Medical School *10.1158/0008-5472.CAN-17-0339*
- Texture Analysis Toolbox, Martin Vallières,
<https://github.com/mvallieres/radiomics/tree/master/TextureToolbox>
- Quantitative Image Feature Engine (QIFE) Stanford University, *10.1007/s10278-017-0019-x*
- IBEX: MD Anderson Cancer Center, doi: *10.1118/1.4908210*.
- MaZda, Technical University of Lodz, Poland, [doi:10.1016/j.nima.2012.09.006](https://doi.org/10.1016/j.nima.2012.09.006)
- LifeX , Gustave Roussy, Parigi, *10.1158/0008-5472.CAN-18-0125*

Feature selection

- The building of a radiomic models has two phases.
- In the first, feature selection, the variables are reduced by eliminating those that are:
 - Redundant, because they are inter-correlated
 - Not predictive (not associated with the outcome)



Feature selection methods

- *minimum redundancy maximum relevance (mRMR)* calculates mutual information (MI) between a set of features and the outcome. The set of features with maximum MI is selected.
- *RELIEF (RELevance In Estimating Features)*, ranks the features according to how well they separate patients with different outcomes but similar values of features:
 - Better score to features with different values in patients with different outcome
 - Penalizes features which have different values in patients with the same outcome
- *Stepwise selection* is an iterative process which adds or removes features to a model at each step. Then the variables are included in the model according to a statistical test with null hypothesis that the variable has zero coefficient in the model-

Machine learning

- Radiomic signature: combination of variables with high predictive power
- Classifier: model to classify the patient e.g. responder, non responder to therapy

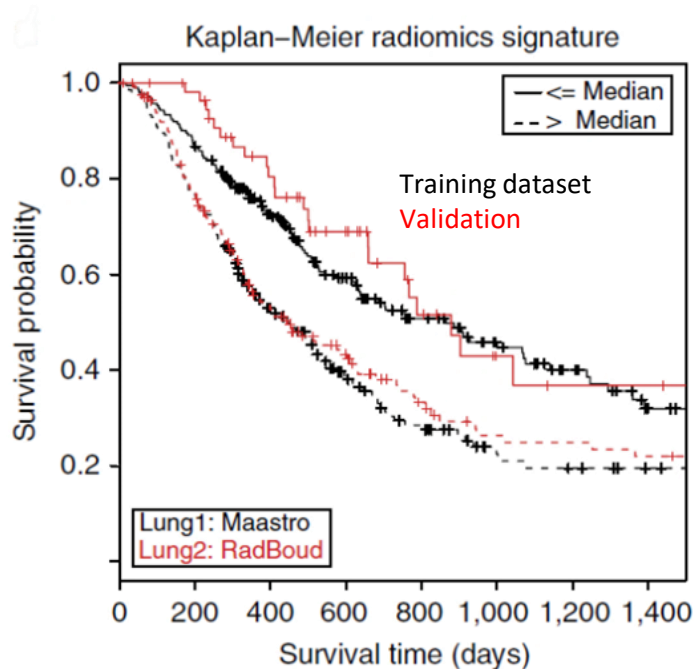


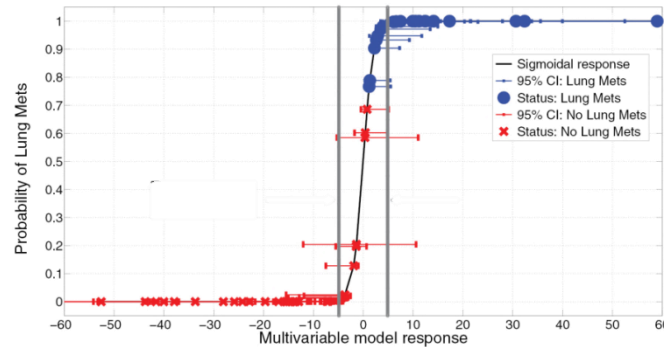
TABLE 6. Summary of Classification Results Obtained by 10-Fold CV

Parameter set	Algorithm	TP rate	FP rate	Specificity	Precision
T1	ANN	0.968	0.091	0.909	0.968
	k-NN	0.935	0.091	0.909	0.967
T2	ANN	0.968	0.273	0.727	0.909
	k-NN	0.935	0.182	0.818	0.935
DWI	ANN	0.903	0.182	0.818	0.933
	k-NN	0.968	0.182	0.818	0.938

FP, false-positive; TP, true-positive.

Machine learning

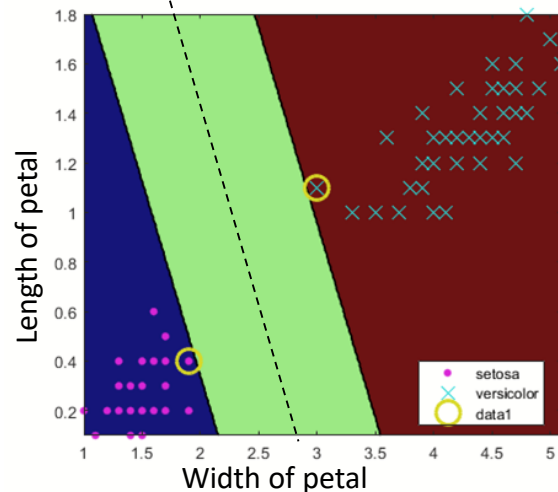
- Logistic Regression



- Support Vector Machine

<https://www.mathworks.com/help/stats/support-vector-machines-for-binary-classification.html>

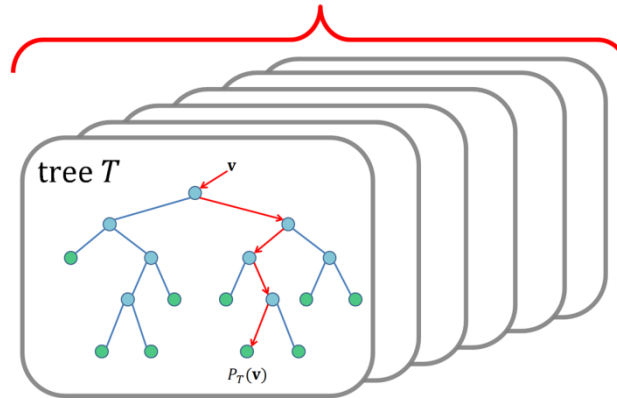
Classificazione automatica dell'iris



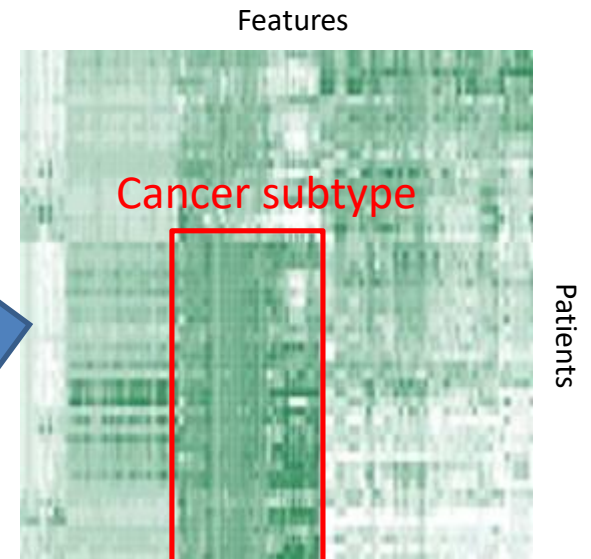
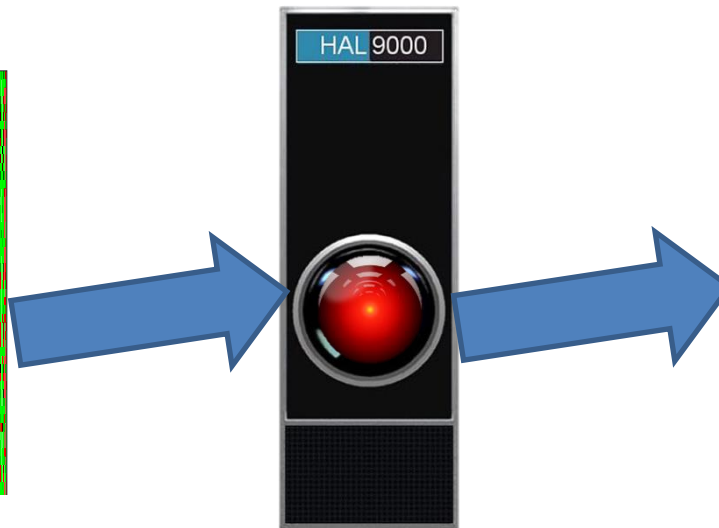
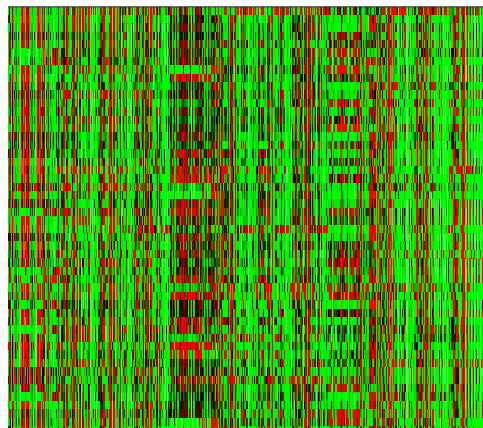
Machine learning

- Random forest

Decision Forest

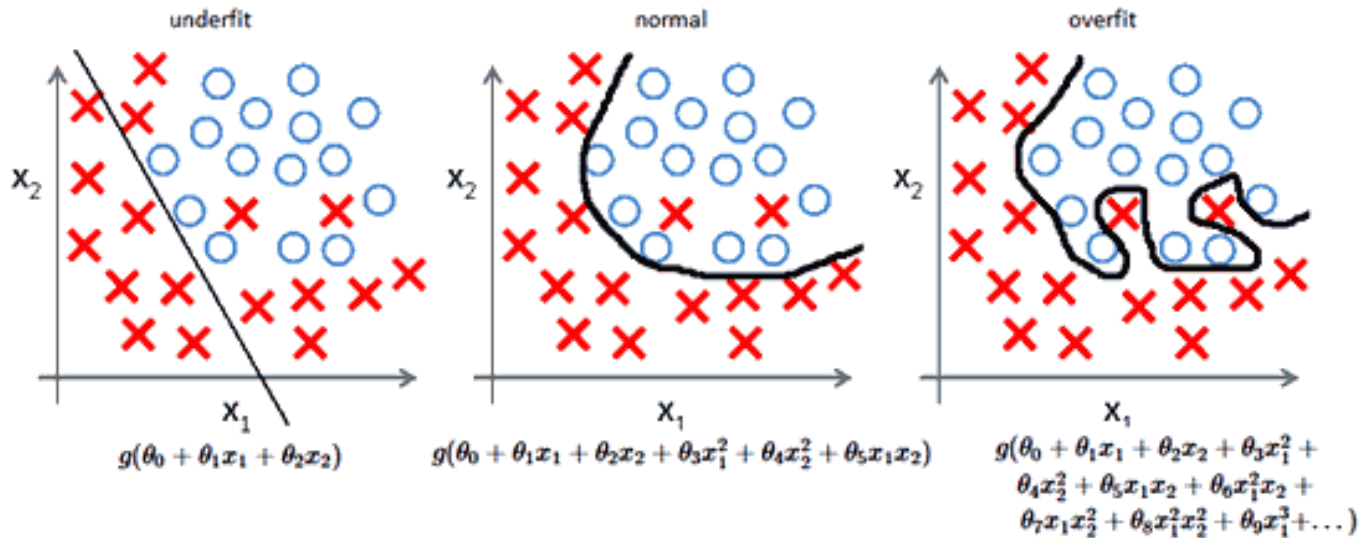


- Unsupervised methods



Overfitting

- Too many variables --> risk of overfitting



- The overfitted model fails when used on a dataset different from the training dataset (poor generalizability)
- Overfitting can be avoided with careful feature selection and validation

Validation

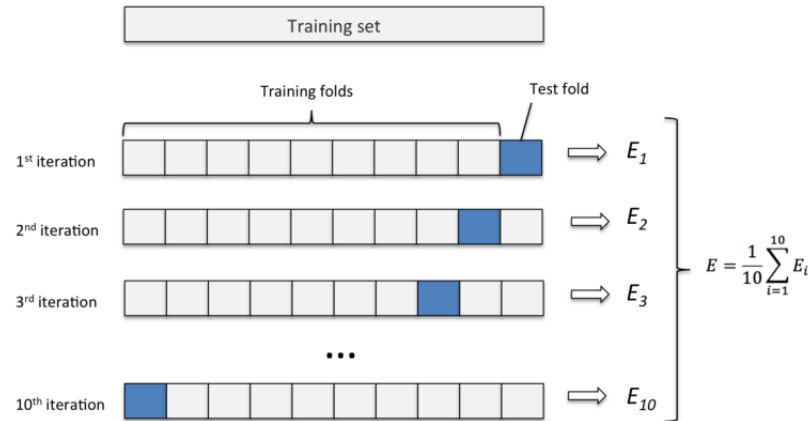
According to TRIPOD (Transparent reporting of a multivariable prediction model for individual prognosis or diagnosis) criteria, there are the following validation methods:

- 1) Developing and validating on the same data, which gives apparent performance. This evaluation is usually optimistic estimation of the true performance
- 2) Developing the models using all the data, then using resampling techniques to evaluate the performance
- 3) Randomly split the data into 2 groups for development and validation separately
- 4) Split the data non-randomly (e.g. by location or time), which is stronger than 3)
- 5) Develop the model using one data set and validate on separate data. Stronger than performing posterior splitting of data



Resampling techniques

- Cross validation



- Bootstrap

~~1~~ 2 ~~3~~ 4 ~~5~~ 6 7 8 9 10

Bootstrap sample 1

1 4 3 5 5 3 7 8 9 10

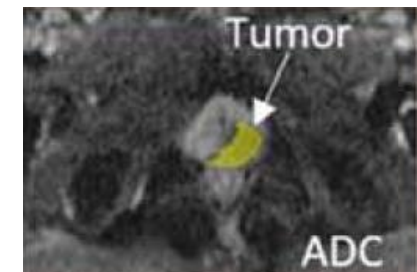
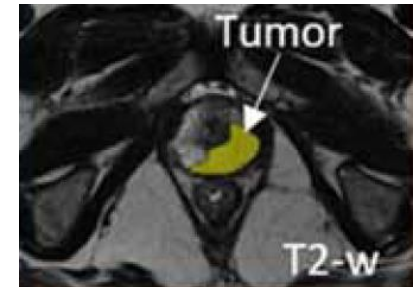
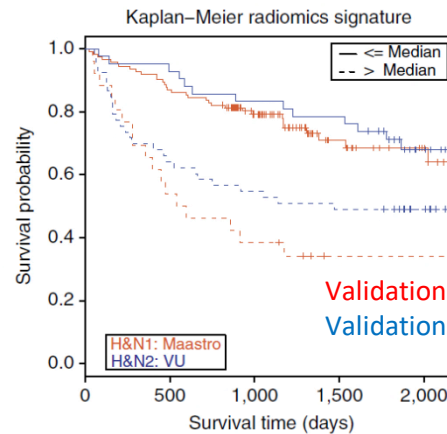
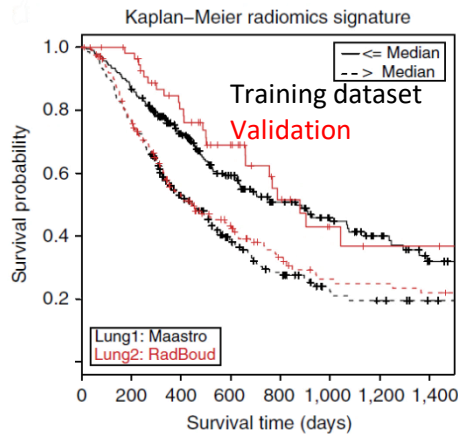
1 ~~2~~ 3 ~~4~~ ~~5~~ 6 7 ~~8~~ 9 10

Bootstrap sample 2

1 2 3 1 5 8 5 8 9 1

- Other techniques: “jackknife” or leave-one-out (LOOCV), where a patient is removed from analysis at each iteration

Examples of predictive models

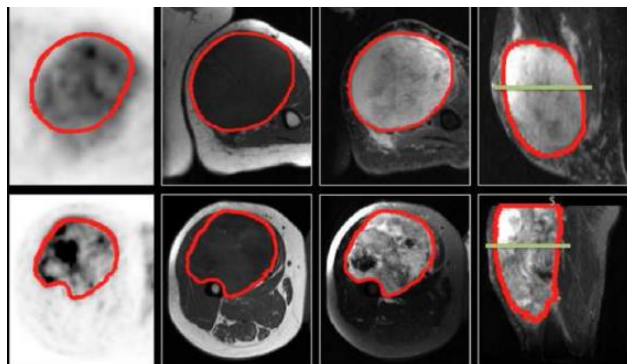


- Survival for lung and H&N squamous cell carcinoma

Aerts et. al NATURE COMMUNICATIONS | 5:4006 | DOI: 10.1038/ncomms5006

Gleason score and biovhemical relapse in prostate tumor

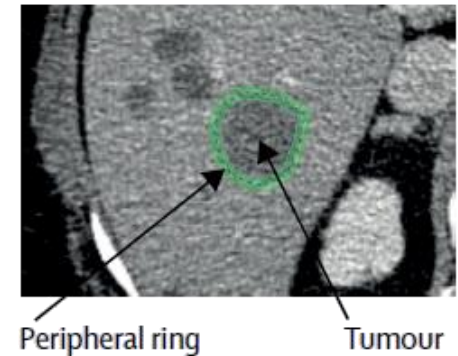
Gnep, J. MAGN. RESON. IMAGING 2016



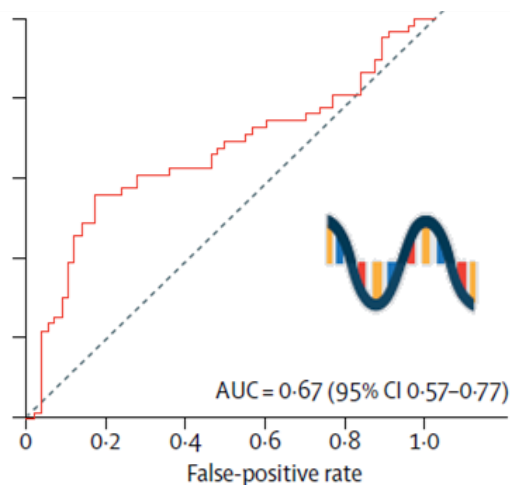
Distant metastases from sarcoma of extremities

Immunotherapy

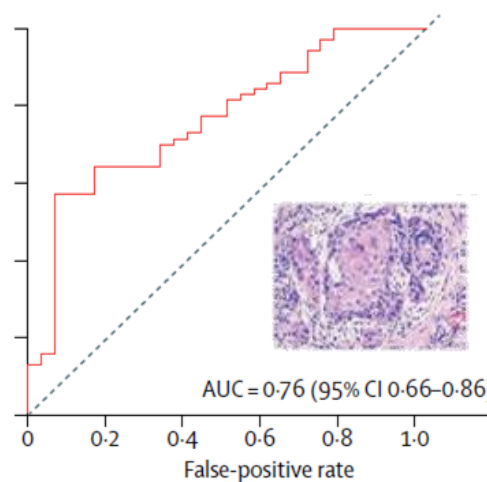
- Model for immunotherapy
- Training set of 135 patients with different tumors
- Radiomic signature for presence of CD8 antigens estimated from RNA sequencing



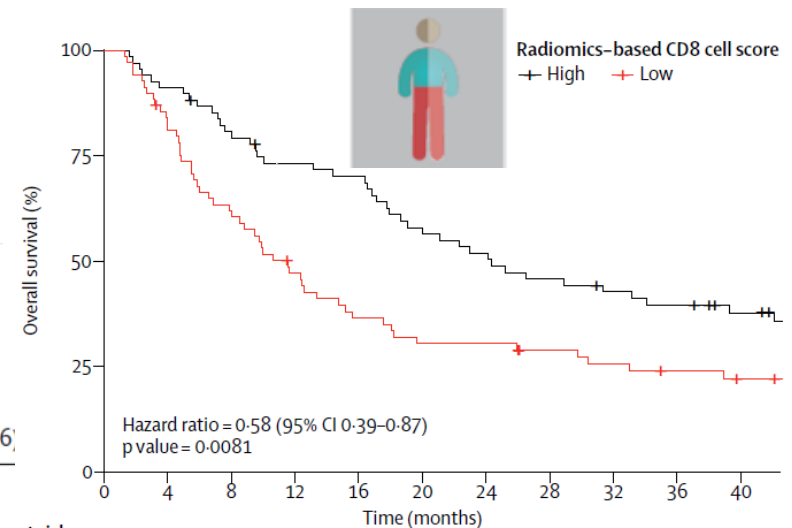
Gene expression of CD8 cells (119 pts)



Phenotype of tumor desert-immune (few CD8 cells) vs inflamed (many cells CD8), 100 pts



Survival of patients treated with immunotherapy (137 pts)

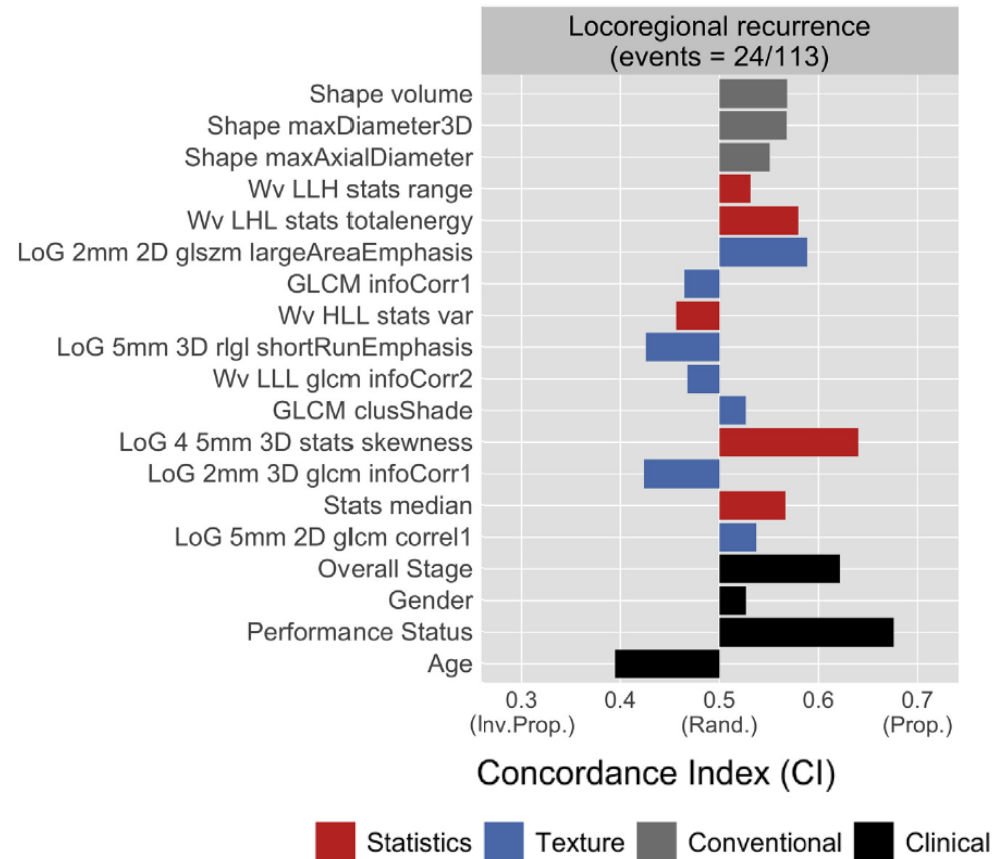


Prediction for local recurrence in SBRT

113 patients

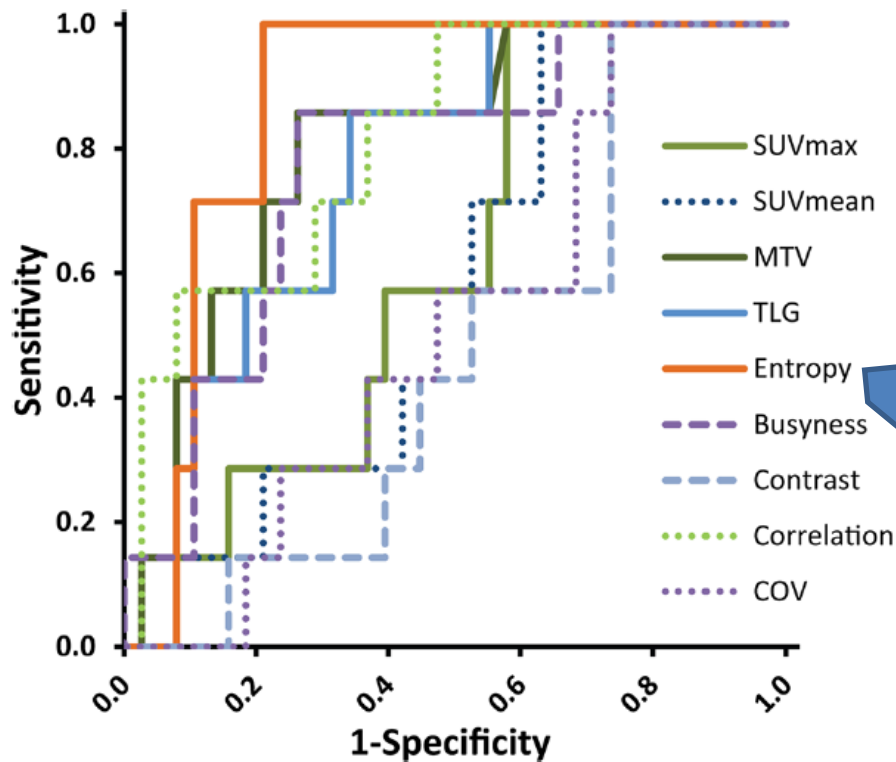
close to the chest wall:
10–12 Gy * 5 fractions,
12–14 Gy * 4 fractions
Other: 18 Gy * 3

Free breathing CT



No feature had significant correlation with recurrence!

Prediction for local recurrence in SBRT from PET



45 patients

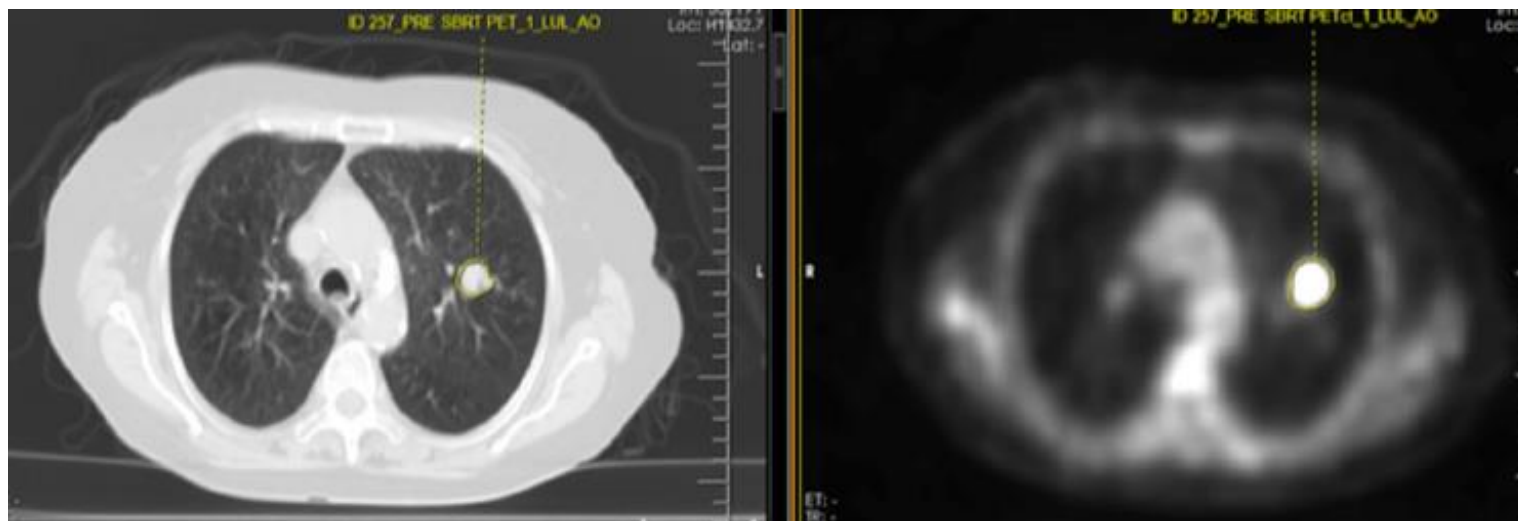
24–45 Gy delivered in 3–5 fractions.
Dose prescribed to the 60% isodose
which had to cover 100% PTV

Significant correlation of
several textural parameters
with local recurrence.
AUC value for entropy of 0.872

Figure 1 Value of textural and standard PET parameters for prediction of local recurrence. ROC curves for prediction of local recurrence through different PET parameters. Coarseness is the same curve as busyness.

Regional control after SBRT:PET/CT

- Radiomics on PET/CT for prediction of control and survival in SBRT-treated lung cancer patients.

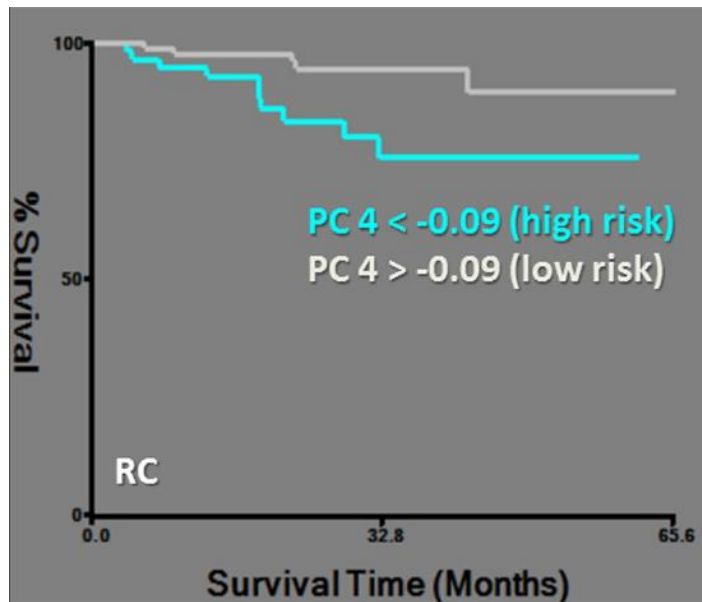


150 patients, 172 cancers

48-56 Gy SBRT Fractionation not included

Regional control after SBRT:PET/CT

- Radiomics on PET/CT for prediction of control

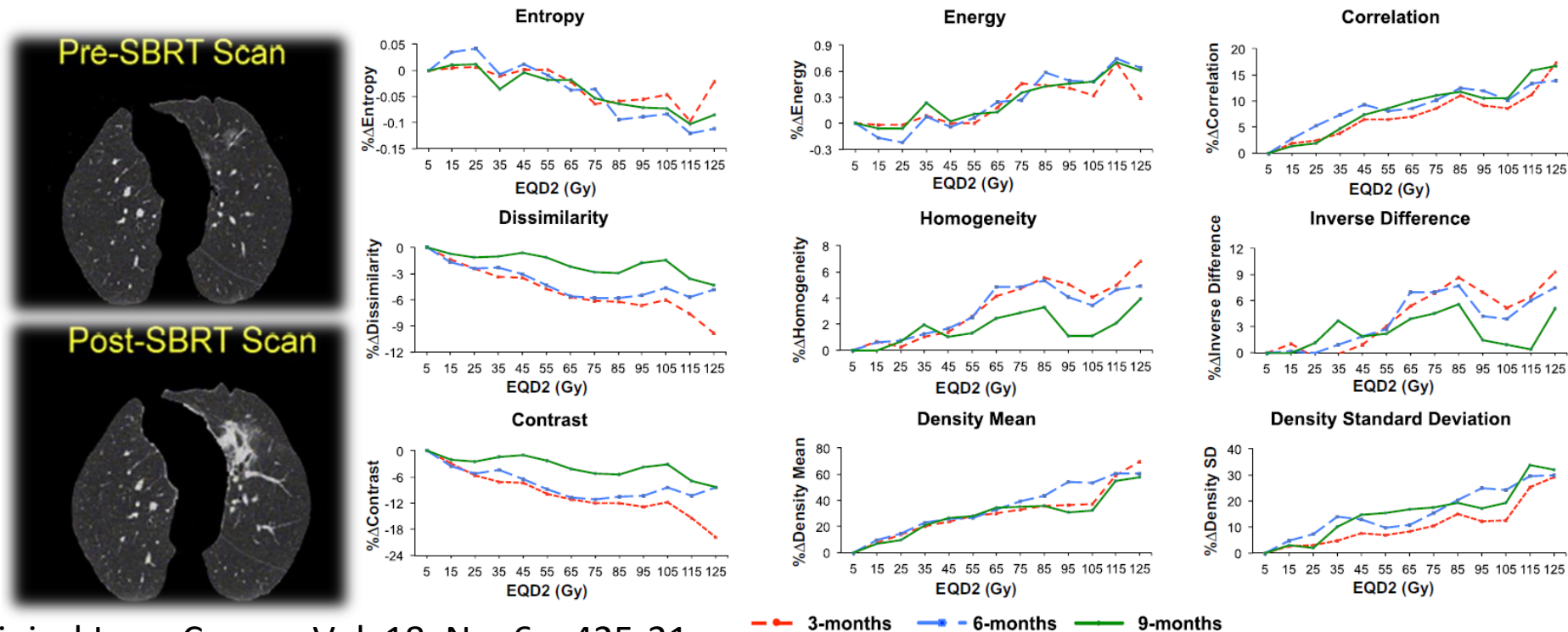


Subgroups of low and high recurrence free survival were determined by a cut-off value of 0.09 for radiomic signature PC4

Radiomic signature "PC4"	
1 st order	Kurtosis (PET)
	Skewness* (PET)
Textural	Homogeneity (PET)
	Normalized Entropy (PET)
Shape	Area regularity (PET)
	Area regularity (CT)
	Perimeter regularity (2) (PET)

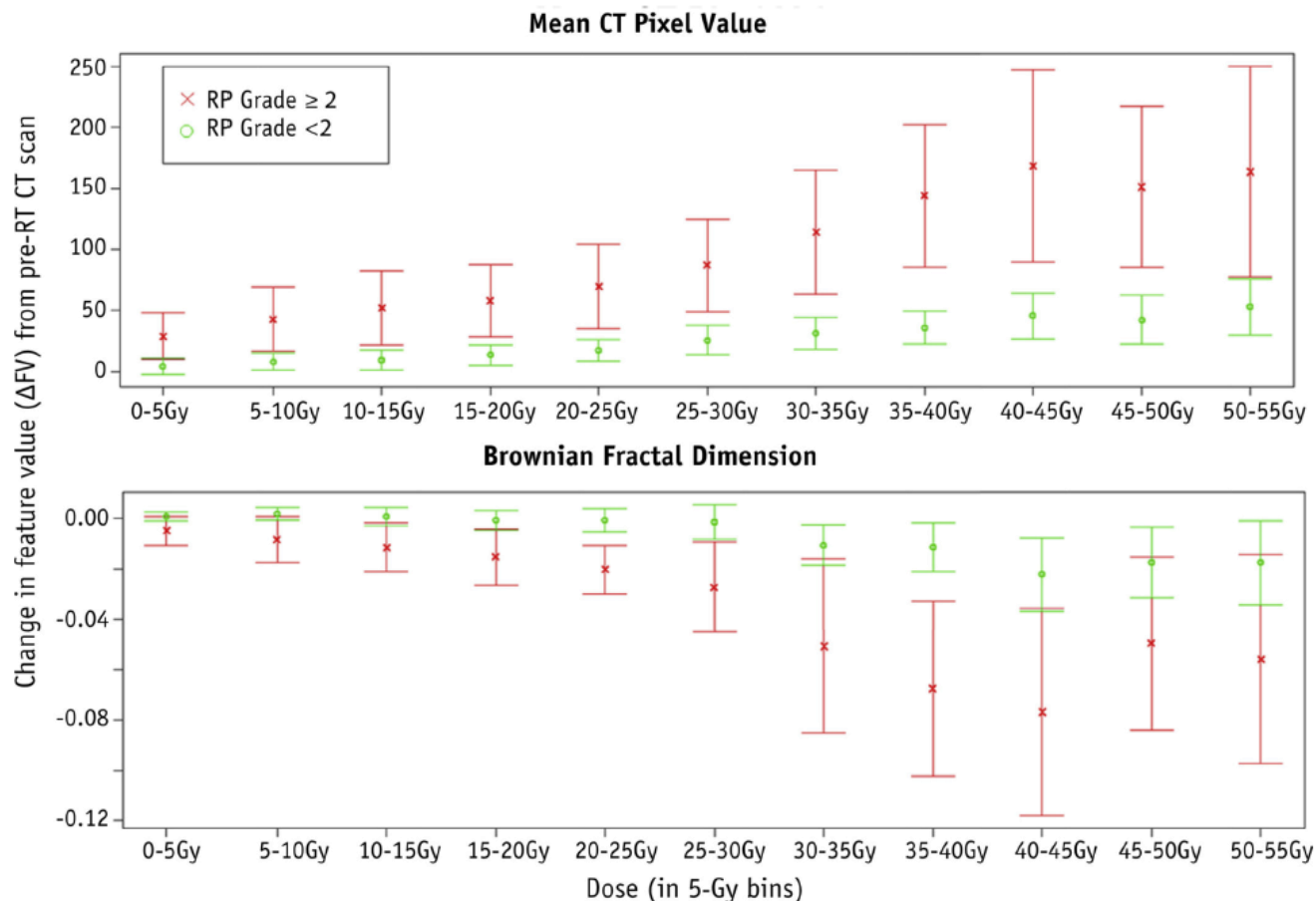
Lung injury

- Radiomic features significantly correlates with lung-injury scored by oncologist post-SBRT (18 Gy*3, 12.5 Gy*4, 12 Gy*5)
- GLCM features outperformed histogram features



Prediction of radiation pneumonitis

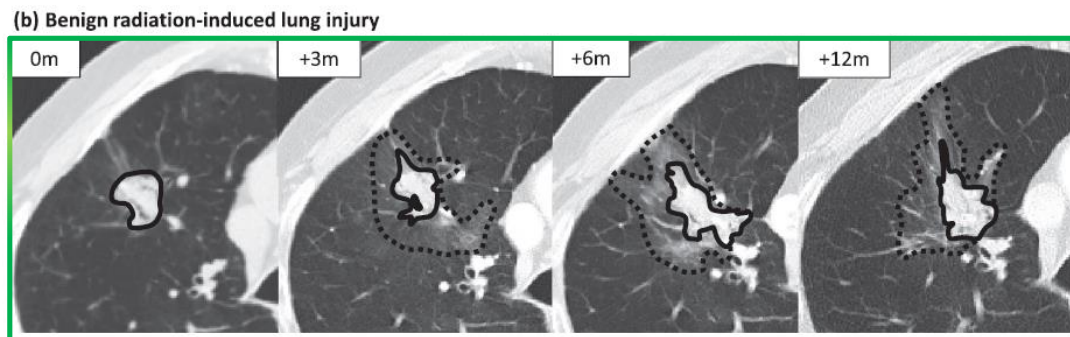
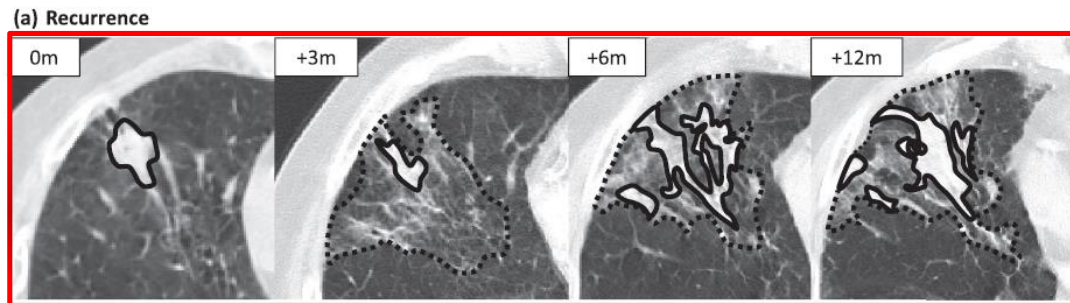
- 50.4 Gy, non-SBRT, esophageal cancer



Differentiation of recurrence

- On two-fold CV, first-order features yielded 73% accuracy, second order 76%–77%
- longest axial diameter and 3D volume, gave 60% and 57%

Recurrence



Benign
changes

FIG. 2. Manual delineations of post-SABR consolidative and ground-glass opacity findings throughout follow-up for a patient with recurrence (a) and radiation-induced lung injury (b). The zero-month (0m) time point indicates the pretreatment lesion. The solid lines enclose consolidative regions and the dashed lines enclose ground-glass opacity regions.

Radiomics of oropharyngeal tumor

- Observational, retrospective, monoinstitutional study at the CRO - Aviano
- Collaboration among Medical Physics, Radiotherapy, Nuclear Medicine, Radiology
- Has the objective of building a predictive model for:
 - HPV status, and
 - response (complete/not complete) at 3 months from the end of radiotherapy

From radiomic analysis of pretreatment images of the patient and dose distribution

Radiomics and HPV status

- The tumors in HPV-positive patients appear more homogeneous and small in CT

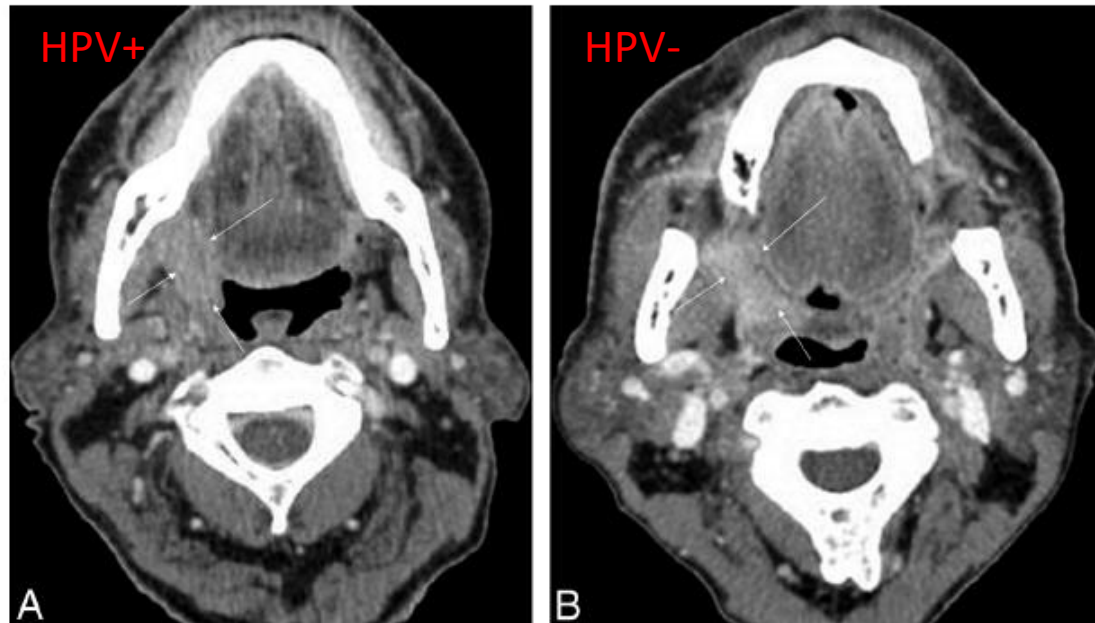
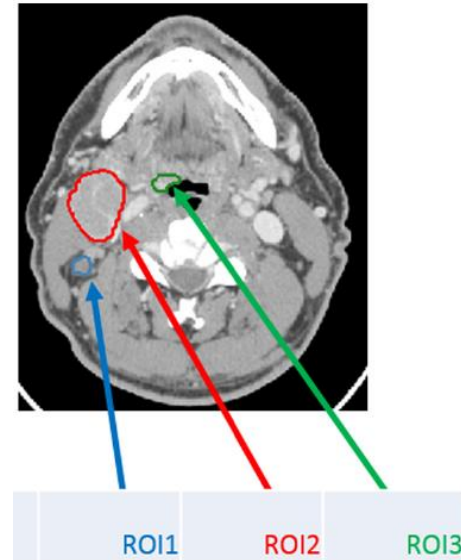


FIG 3. Representative examples of patients with HPV-positive and HPV-negative SCC. A, HPV-negative right tonsillar squamous cell carcinoma (arrows) in a 65-year-old man. B, HPV-positive right tonsillar squamous cell carcinoma (arrows) in a 65-year-old man.

Radiomics and HPV status

- Model based on contrast-enhanced CT, 315 patients oropharyngeal
- 150 patients for training, 165 validation
- Model had AUC of 0.915 in validation



Yu K, Clinical and Translational Radiation Oncology 7 (2017) 49–54

- Model for prediction of HPV determined from p16
- CT, no contrast
- Multicentric database of 778 patient, randomly split into training dataset (80%) and validation ($N = 150$).
- The model scored AUC=0.764 in validation

Oropharyngeal: local control

- 465 pazienti
- Local control proven pathologically (biopsy and/or resection) or radiologically
- Analysis on contrast enhanced CT
- Radiomic signature based on:
 - Intensity Direct Local Range Max: average of range (max-min) for every voxel with respect to surrounding region
 - Neighbor Intensity Difference Complexity: measures the perceived complexity in the image

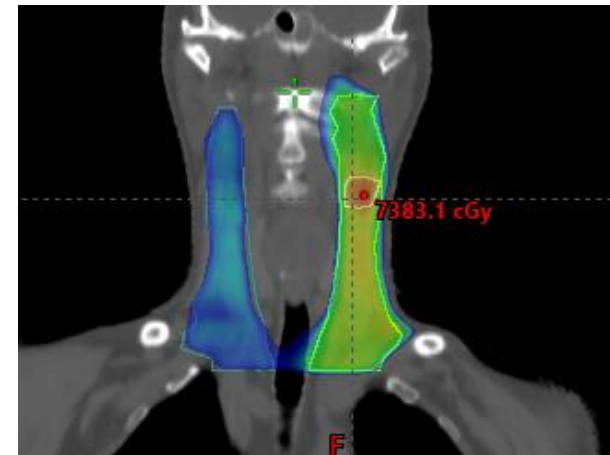
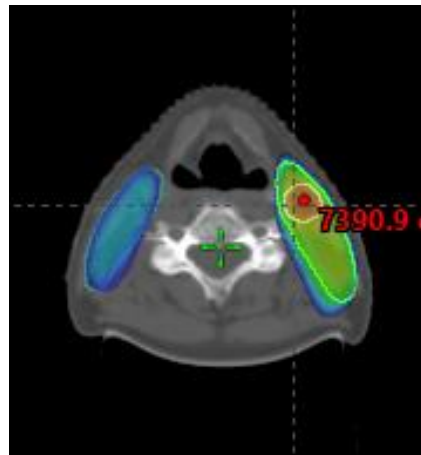
The radiomic signature had higher predictive capability than variables HPV status and administered therapy

Methods

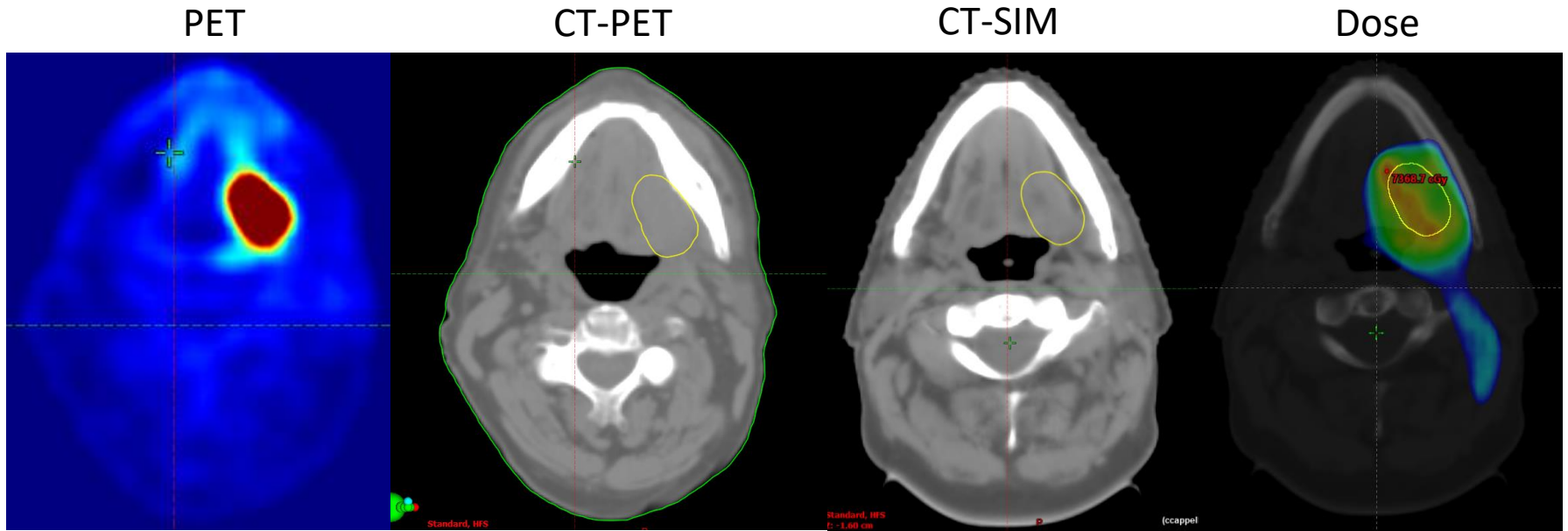
- 51 Patients treated with IMRT
- 70.95 Gy to microscopic disease
- 62,70 Gy to high risk lymph-nodes
- 59,10 Gy “ low risk “ “

Characteristics of patients

Patients	51
Male/female	41/10
Chemotherapy (no, Concomitant, neoadjuvant, neoad.+conc.)	1/12/36/2
Stage TNM 8°: 1, 2, 3, 4A, 4B	14/8/4/21/4
HPV Status (+,-)	28/23



Methods



- Tumor was contoured by one clinician using PET
- Contour reported on CT-PET and simulation CT using image registration
- Variables extracted also from dose distribution

Protocols of acquisition

PET Philips Gemini TF 16

Average injected activity of ^{18}F -FDG was 280 MBq

Algorithm of reconstruction PET “Blob-OS-TF”,

a 3D ordered subset iterative TOF reconstruction technique

Matrici 144×144 con voxel $4 \times 4 \times 4 \text{ mm}^3$

CT-PET Philips Gemini TF 16

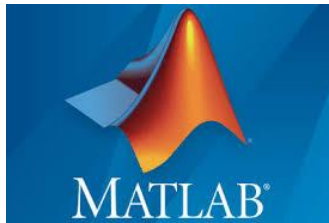
- 120 kV, 108 mA average, pitch 0.83, acquisition time 0.5 s
- Slice thickness 5 mm, kernel: ‘B’ body

CT-SIM Toshiba Aquilion/LB

- 120 kV, average tube current 300 mA , rotation time 0.75 s
- Slice thickness 2 mm, kernel: ‘C13’

Methods

- Software written “in-house” in Matlab, benchmarking with Ibex



IBEX is Intended for Research Use Only.

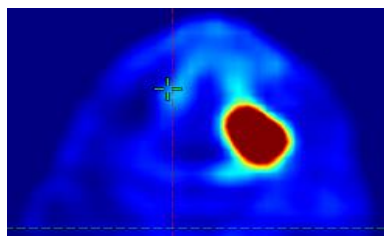
IBEX^{V1.0 β}

Imaging Biomarker Explorer Software

- 21 shape variables
- 47 textural (+ filters gaussian, LOG, median)
- 5 higher order (“ “ “ ”)
- In total: 937 features per patient
- Stepwise feature selection, support vector machine
- Cross-validation

Preliminary results (1)

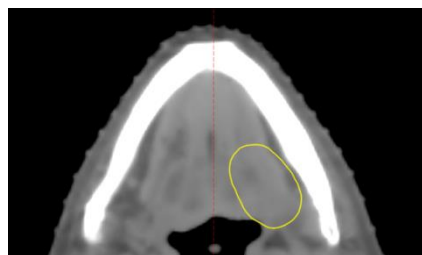
- Model for HPV status:
- 1 shape (solidity), 2 simulation CT, 1 PET, 2 dose variables were selected
- In the cross-validation:
 - Sensitivity (positive on patient with HPV+): 85,2%%
 - Specificity (negative on patient HPV-): 83,3%



Inv.Diff.Norm PET

Measures local inhomogeneity

		Real	
		HPV+	HPV-
Predicted	HPV+	23	4
	HPV-	4	20

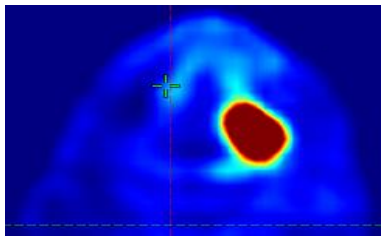


GLCM Cluster Prominence
Measures variability of values

Preliminary results (2)

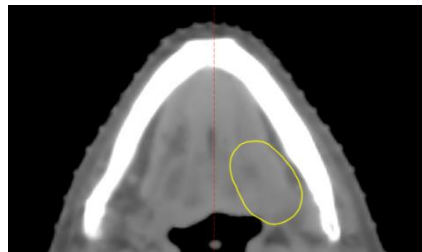
- Model for complete response 3 months from therapy:
- 1 shape (roundness), 3 simulation CT, 4 PET features were selected
- In the cross-validation:
 - Sensitivity (positive on patient with HPV+): 100,0%
 - Specificity (negative on patient HPV-): 95,1%

Matrice di confusione		Real	
		RC+	RC-
Predicted	RC+	39	2
	RC-	0	10



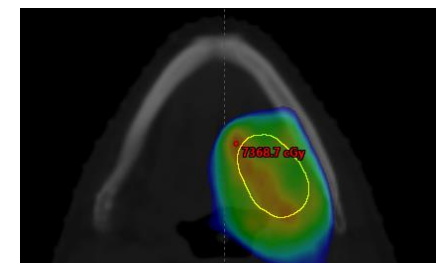
SRLGE PET

Describes presence of stripes of low value voxels



Long run emphasis CT-SIM
Presece of stripes of voxels with same value

Dose Range
Related to inhomogeneity of dose



Conclusions

- Radiomics is entering its mature phase:
 - The number of radiomic papers is increasing exponentially
 - More and more radiomic studies have solid validation
 - more attention than in the past to feature reproducibility
- If you want to approach radiomics:
 - Read some of the many excellent reviews on the subject
 - Read the Imaging Biomarker Standardisation Initiative
 - Download and use open source software

School on Medical Physics for Radiation Therapy: Dosimetry and Treatment Planning for Basic and Advanced Applications



25 March – 5 April 2019
Trieste, Italy

Further information:

Activity URL: <http://indico.ictp.it/event/8651/smr3278@ictp.it>

Thank you for your attention!

Michele Avanzo
Medical Physicist
Centro di Riferimento Oncologico
IRCSS Aviano (PN) mavano@cro.it

CRO
AVIANO