ASSESSING THE PROGNOSTIC IMPACT OF 3D CT IMAGE **TUMOUR RIND TEXTURE** FEATURES ON LUNG CANCER SURVIVAL MODELLING

Alanna Vial, David Stirling, Matthew Field, Montserrat Ros, Christian Ritz, Martin Carolan, Lois Holloway, Alexis A. Miller





Health Illawarra Shoalhaven Local Health District





Introduction

- Develop a technique for developing prognostic image characteristics, termed radiomics, for non-small cell lung cancer based on a tumour edge region-based analysis.
- Texture features were extracted from the rind of the tumour in a publicly available 3D CT data set to predict two-year survival.
- Medical experts have expressed a need for this data to be analysed
 o improve treatment planning and improve overall patient survivability.
- Impact of providing personalised automated radiotherapy treatment.

What is Radiomics?

• Radiomics: is the application of textual image processing and data mining in the field of radiology. This is a novel approach to improve treatment outcomes, by finding attributes and features in CT imaging data.



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Why Personalised Radiotherapy?

Current Issues

Patients vary significantly, however clinical trials can't match every variation exactly, hence treatment plans can be incorrect.

Data Mining

- Terabytes of cancer data available to be mined.
- It is proposed that through data mining all data better treatment options can be applied.

Radiomics

 Radiomics is the proposed method for solving this problem.



Key Challenges

Preprocessing the data

- Remove duplicated
 data
- Normalise data to ensure it is comparing like to like
- Transform data from DICOM images to MATLAB files for simpler analysis.

 Discover Radiomic signatures and other textual features to find relationship with 2-year survivability

Finding Features

Modelling the data

Use data mining and image processing techniques such as Logistic Regression or Support Vector Machines to model data.

Extracting the Rind

Region exploration around the GTV



Deriving the Boundary Masks

To derive the mask, we first define the set of points representing the inside and outside rind, let $\mathbf{x} \in \mathbb{R}^3$ and $\mathbf{z} \in \mathbb{R}^3$ define positions vectors in the 3-Dimensional image set frame. Further, let $\mathcal{V} \subset \mathbb{R}^3$ be the set of all points in the gross tumour volume (GTV) as given by the expert or any other segmented region of interest (ROI) and $\hat{\mathcal{V}} \subset \mathcal{V}$ is defined as all points on the boundary of this region.

Suppose there is a distance function that can be constructed for each ROI, $d(\hat{\mathcal{V}}) : \mathbb{R}^3 \to \mathbb{R}$, which will be defined by the following

$$d(\mathbf{x}, \hat{\mathcal{V}}) = \begin{cases} \min \|\mathbf{x} - \mathbf{z}\|, \ \forall \mathbf{z} \in \hat{\mathcal{V}}, \ \forall \mathbf{x} \notin \mathcal{V} \\ -\min \|\mathbf{x} - \mathbf{z}\|, \ \forall \mathbf{z} \in \hat{\mathcal{V}}, \ \forall \mathbf{x} \in \mathcal{V} \end{cases}$$

Now the outer boundary region can be expressed by the set

$$S_o = \{ \mathbf{x} \mid 0 < d(\mathbf{x}, \hat{\mathcal{V}}) < L \},\tag{1}$$

where L is the width of a region that contains all line segments perpendicular to the surface. So is therefore a region external to the ROI. Conversely, we have

$$S_i = \{ \mathbf{x} \mid -L < d(\mathbf{x}, \hat{\mathcal{V}}) < 0 \},$$
(2)

defining an inner region such that $S_i \subseteq \mathcal{V}$. Colloquially, we refer to the region, $S = \{S_i \cup S_o\}$, as the rind of a given ROI, obtained by expanding or contracting the ROI uniformly from the surface by L. 8

Rind Comparison















Note: the dark blue background is not included in this analysis.

Rind analysis

• Logistic Regression



Whole volume - Training data Whole Volume - Cross Validation, AUC is 0.6032 Survival Probability True positive rate <=Median >Median 0.5 0.5 0 0 500 1000 1500 2000 0.2 0.4 0.6 0.8 Survival Time (days) False positive rate Inside Rind - Cross Validation, AUC is 0.56626 Inside Rind - Training data Survival Probability <=Median >Median 0.5 0 1500 0.2 0.6 0.8 0 500 1000 2000 0 0.4 Survival Time (days) False positive rate Outside Rind - Cross Validation, AUC is 0.61429 **Outside Rind - Training data** Survival Probability <=Median >Median 0.5 0 0 500 1000 1500 2000 0 0.2 0.4 0.6 0.8 1 Survival Time (days) False positive rate

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Kaplan Meier Survival Curves – Three Features



Kaplan Meier Survival Curves – Texture Features

AUC Results from 20 iterations of logistic regression using ten fold cross validation

Volume	Three Features	Texture Features
Analysed	$AUC\pm\sigma$	$AUC \pm \sigma$
Whole vol.	$0.589 {\pm} 0.015$	0.661 ± 0.016
Inner rind	$0.558 {\pm} 0.011$	0.679 ± 0.024
Outer rind	0.598±0.011	0.689 ± 0.015
Outer rind with vol.	$0.583{\pm}0.013$	0.699 ± 0.011
Vol. excl. inner rind	$0.584{\pm}0.013$	0.624 ± 0.015

Conclusion

- Explored the textual radiomic features in a whole 3D tumour volume, compared to the inside and outside rind of the tumour only, for CT images of NSCLC.
- The derived models were compared against the previous methods of training radiomic signatures that are descriptive of the whole tumour volume.
- Radiomic features derived solely from regions external, but neighbouring, the tumour were shown to also have prognostic value.
- Textual features found within the tumour rinds were very similar to the textures in the entire tumour volume, and this analysis predicted two-year survival with an improved accuracy of 3% for survival classification using textures from the outside rind compared to the whole volume.

Clinical Implications

- It is important to note, that there is significant clinical uncertainty in defining the tumour boundaries of the GTV and that this should be considered when viewing these results.
- These results indicate that while the centre of the tumour is currently the main clinical target for radiotherapy treatment, the tissue immediately around the tumour is also clinically important.

Future Work

- One limitation of this work is in the arbitrary selection of the radiomic features.
- In future work we will use a machine learning algorithm such as convolutional neural networks to determine which radiomic features have the highest prognostic significance for determining two-year survival.
- Furthermore the radiomic features themselves are determined based on general mathematical formulas associated with analysing an image, in a future study we plan to use modern image processing techniques to determine alternative radiomic features.



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