

Technique for Reliable Use of Images in Medical Physics TRU-IMP A Dynamic GUI

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Overview

TRU-IMP

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Context & Goal

Where to find it

What it does

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Where to find it

What it does

- ▶ Work in Progress!
- ▶ More features to be added with users' inputs and comments.

Context: Nuclear Medicine

- ▶ Nuclear medicine uses radiopharmaceuticals to detect functional activity in an organism;
- ▶ It is a type of functional imaging, i.e. not anatomical;
- ▶ An anatomical image (eg. CT, MRI, US) can be added to have both anatomical and functional features;
- ▶ Many physical factors affect the quality of the produced image.

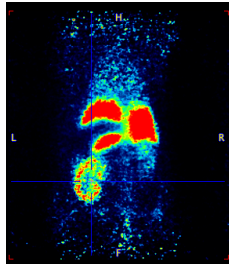


Figure: Nuclear medicine acquisition of a rat injected with a radiopharmaceutical gathering in the kidneys.

Context: Dynamic Imaging

- ▶ In a dynamic context, many timeframes are made to create an evolving acquisition;
- ▶ This allows to track the movements of the radiopharmaceuticals through the organism;
- ▶ This can be done, for instance, by using a Time-Activity Curve (TAC);
- ▶ New factors are present that reduce the quality of the image, among which movements between each timeframe.

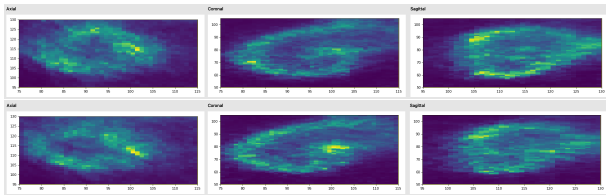


Figure: Two subsequent timeframes of a dynamic acquisition on a rat centered around the left kidney

- ▶ Segmentations is the act of selecting a Region of Interest (ROI);
- ▶ This can be done manually or automatically;
- ▶ With dynamic images, the notion of a good segmentation is **problematic**.

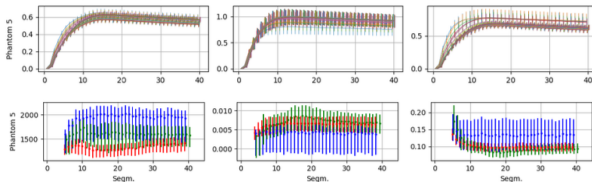


Figure: TACs and pharmacokinetics parameters for a simple dynamic phantom using 3 different segmentation schemes. What is seen is that the TACs agree, but not the parameters.

- ▶ Develop a tool to analyze segmentations in a dynamic context;
- ▶ Determine the adequacy of current methods;
- ▶ Techniques for Reliable Use of Images in Medical Physics (TRU-IMP)

Where to Find It

- ▶ Currently hosted on GitHub



Figure: Scan Me (QR Code)

What is Needed

- ▶ If using the GUI packaged: **nothing**
- ▶ If using the source code: Python 3.X with some packages:
 - ▶ Numpy;
 - ▶ Scipy;
 - ▶ SkImage;
 - ▶ Matplotlib;
 - ▶ Time, OS, Numba*;
 - ▶ Dynesty;
 - ▶ PyDicom;
 - ▶ Pickle.

TRU-IMP: The Basics

- ▶ A free tool to work on 4D Dicom Images (.dcm) and soon NIfTI (.nii) ;
- ▶ Main uses:
 - ▶ Segmentations;
 - ▶ Comparison of segmentations;
 - ▶ Uncertainty on TACs;
 - ▶ Extraction of pharmacokinetic parameters.

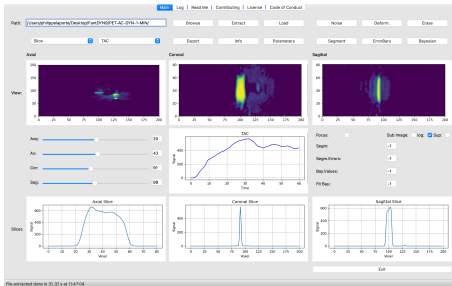


Figure: Basic view of the GUI, with an acquisition open

- ▶ Many segmentation schemes implemented:
 - ▶ Geometric;
 - ▶ Gradient-based;
 - ▶ Filling-based;
 - ▶ Statistics-based:
 - ▶ ICM;
 - ▶ Fuzzy C-Means (FCM).
- ▶ The user can select most of the hyperparameters manually;
- ▶ All segmentation schemes are based on a reference timeframe (static segmentation).

TRU-IMP: Comparisons

- ▶ In a dynamic setting, multiple timeframes can be used as a reference;
- ▶ Many segmentation schemes are available;
- ▶ To compare segmentations, many tools are available directly and graphically:
 - ▶ Dice & Jaccard coefficients;
 - ▶ Overlapping of segmentation and image;
 - ▶ The loss functions and convergence parameters (when relevant)

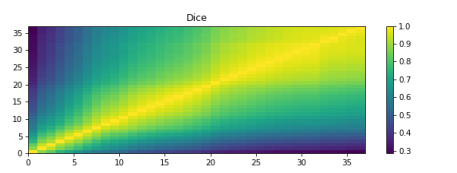


Figure: The Sørensen-Dice coefficients for different segmentation on a dynamic acquisition of a simple dynamic phantom

TRU-IMP: Uncertainties

- ▶ Comparing Time-Activity Curves (TACs) requires error bars;
- ▶ Segmentations normally don't produce them directly;
- ▶ The introduction of uncertainties is done by deformation of the Region of Interest (ROI):
 - ▶ Linear shifts;
 - ▶ Expansions/contractions;
 - ▶ Rotations;
- ▶ Another method is to sample the fuzzy segmentation (when relevant).

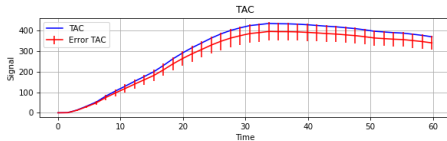


Figure: TAC and an associated uncertainty induced by linear shifts

TRU-IMP: Pharmacokinetic Parameters

- ▶ A model of displacement of the radiopharmaceutical can be used;
- ▶ Parameters dictate the flow between sections of the image;
- ▶ These parameters can be determined by fitting the TAC.

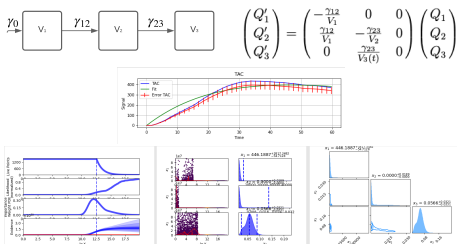


Figure: Pharmacokinetic model of the simple phantom with the fitted curve as per the extracted parameters.

The bottom image represents the convergence analysis of the *Dynesty* module

- ▶ Compatibility with .nii files;
- ▶ Export of the segmentations directly;
- ▶ Work with multiple acquisitions at the same time;
- ▶ Segmentations with errors from a propagation of uncertainties;
- ▶ Exploration of parameter space.

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Where to Find It



Figure: Scan Me (QR Code)