A Local ROI-specific Atlas-based Segmentation of Prostate Gland and Transitional Zone in Diffusion MRI Junjie Zhang¹, Sameer Baig¹, Alexander Wong², Masoom A. Haider¹, Farzad Khalvati¹ Image: Construction of Prostate Gland



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Introduction

Segmentation of prostate and related anatomic structure, such as transitional zone, in medical images facilitates prostate cancer detection, as well as a number of other clinical practices.

Usually the first step in computer-aided detection of prostate cancer.

The current practice of manually contouring the prostate gland in MR images is a tedious task.

Computer-assisted segmentation methods could reduce the time spent manually contouring the prostate gland and potentially reduce the inter-user variability of diagnosis.

Related Researches



A popular class of algorithms in literature for prostate segmentation is *atlas-based segmentation* (ABS) algorithms [1].

In ABS, registration between atlas images and the target image, in particular for MR images, can be difficult:

- Large variability of the MR images in terms of image intensity characteristics (e.g. scanner variability)
- Structure (e.g., different field of views (FOVs) and different imaging center), and anatomical variabilities of scanned regions.

Method

We propose a semi-automatic *local ROI-specific atlasbased segmentation* (LABS) method to segment prostate gland and transitional zone in diffusion MR images (Figure 1), inspired by ABS and a *sequential registration-based segmentation* (SRS) method [2]:

- Minimize user interaction;
- Focus on the vicinity of prostate for atlas matching and atlas-to-target registration;

Step I: User-specified Bounding Box

The user specifies the bounding boxes (BBs) of the prostate gland on key slices (e.g., the base, middle, and the apex);



Figure 1: The proposed local ROI-specific atlas-based segmentation (LABS) pipeline. Each step is highlighted in blocks with different colors.

Results and Conclusion

The algorithm was validated on diffusion MRIs of 100 patients using leave-one-out method*.



Quantitative Analysis

Tables 1, 2 and Figure 2 present results with both 3 and 5 user-specified bounding boxes, and the segmentation results with (w.) and without (w/o. or original) post-

Enlarged BBs are interpolated across slices to produce the prostate volume of interest (VOI).

Step II: Atlas Selection

- Corresponding prostate VOIs are extracted from atlas database w.r.t. the prostate VOI of target patient;
- Best matched VOI is then selected from atlas based on two criteria: the similarity measurement and volume ratio.

Step III: ROI-based Registration

Similar to ABS, best matched VOI is registered to target VOI using an affine registration method.

Step IV: Transformation and Post-processing

Mask VOI in atlas is then transformed and scaled to the size of interpolated BBs to produce segmentation for both prostate and transitional zone.

* The pipeline is implemented in ProCanVAS.

processing, by Dice Similarity Coefficient (DSC).

Table 1. DSC (%) of prostate gland segmentation.

	3 BBs	5 BBs	$\overline{\Delta DSC}$
w/o. post-processing	76.8 ±8.2	78.3 ±6.5	+1.5
w. post-processing	80.2 ±4.7	85.4 ±3.2	+5.2
$\overline{\Delta DSC}$	+3.4	+7.1	

Table 2. DSC (%) of transitional zone segmentation.

	3 BBs	5 BBs	$\overline{\Delta DSC}$
w/o. post-processing	69.1 ±9.6	70.6 ±8.4	+1.5
w. post-processing	73.7 ±6.8	77.3 ±5.9	+3.6
$\overline{\Delta DSC}$	+4.6	+6.8	



Figure 2: Results (DSC) for prostate gland (left) and transitional zone (right) segmentation.



