Medical Imaging Analysis : Lab 2. MevisLab

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1 Introduction

The aim of this lab is to learn the basic structure and use of the MevisLab framework for quick prototyping of medical imaging applications. In this lab you will learn:

- Basic MevisLab filters and structure.
- How to develop medical image analysis prototypes
- Develop a breast boundary segmentation framework.

2 MeVislab

2.1 Installation

MeVisLab is an environment developed by MeVis Medical Solutions AG and Fraunhofer MEVIS in Germany. You can download and install the software at http://www.mevislab.de/download/. Although this is beyond the interest of the lab, MevisLab allows to program your own filters and modules in different programming languages. Documentation is at http://www.mevislab.de/docs/2.5/MeVisLab/Resources/Documentation/Publish/SDK/GettingStarted/. You can check the main modules and the help and examples linked to those (see next section).

2.2 Common MeVisLab Blocks

MeVisLab is used by joining blocks, you can look for them and place them using the Search Modules on top right of the screen (Fig. 1). You can get information on the blocks use by using the **Show Example Network** and **Help** options.

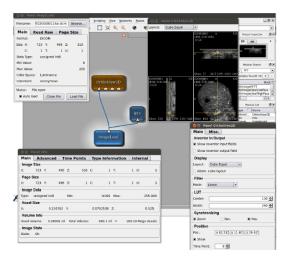


Figure 1: Main MevisLab window

The following list enumerates some basic blocks.

- ImageLoad/ImageSave used to load and save dicom images.
- itkImageFileReader/itkImageFileWriter used to load and save .nii images. If you use it to load dicom images, the pixel dimension will not be understood correctly.

- Info used to get access to the image information.
- OrthoView2D Visualize 3D images with the axial, coronal and sagittal views.
- View2D Visualize data from one view. Useful for 2D images.
- GVROrthoOverlay Permits to show on one visualizer some specified regions as overlay.
- ImageStatistics Calculate several image parameters. You can specify a mask image.
- Arithmetic Used to do any arithmetic operations on images.
- **SubImage** Extracts subimages from an input image based on either voxel start/size, voxel start/end or world start/end. Can also be used to create a region larger than the input image.
- AnonymizeMacro: allows to anonymize patient contained in the Dicom
- Threshold: Threshold an image.
- Connected Components: Obtain the connected components of an image.
- Morphology: Performs morphological operations.
- DimensionSliceClone: Clones a given slice into the given dimension a number of times.
- Orthoprojection: projects a 3D volume into a 2D image using different operations (max, min,mean,...).

3 Breast Boundary segmentation

Open MevisLab and follow the Getting Started Tutorial that introduces you to MevisLab. Given the 3D breast ultrasound volumes given, the aim is to develop a breast boundary segmentation algorithm in order to separate the breast area from the background, as shown in the following figure.

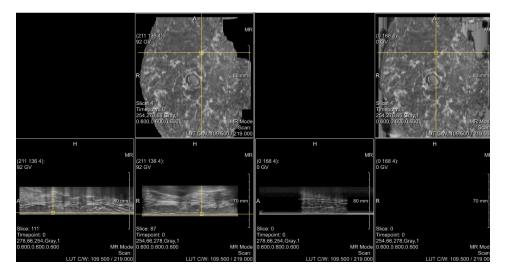


Figure 2: Example of breast segmentation: Original (right) and Masked ABUS volumes (left)

You can use any existing modules, but you should aim at a **fully automatic algorithm**: minimal (or no) user intervention, and the parameters must be the same for segmenting all images (no parameter tweaking for each image).

Questions (to be included in the report)

- \bullet The MeV islab script (file) that implements the image segmentation
- Report indicating:
 - The steps followed to segment the breast. There is no need to describe the algorithms, but you should explain why you apply them.
 - All segmentation results.
 - Limitations of your approach (if any).
 - Encountered problems found in the assignment and/or suggestions