

Module 0: Introduction Biological Image Analysis with the Bioimage Analysis “Toolbelt”

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Objective of Tutorial

- Define emerging techniques (some more established than others) for analyzing biological images
- Emphasis on **Segmentation** and **Tracking** of cells
- Provide Matlab code examples that
 - Demonstrate concepts
 - Allow you (or your student or your colleague) to immediately apply fairly complex image analysis methods to your data



Outline

- Module 1:
 - Active contours for segmentation
 - Active contours for tracking
- Module 2:
 - Scale space for segmentation
- Module 3:
 - Graph theoretic segmentation of filamentous objects
 - Graph cuts for segmentation
- Module 4:
 - Particle filters for tracking



Bioimage Analysis Toolbelt

- Written in Matlab with some routine in C/C++
- Runs on PC and Macintosh platforms
- <http://viva.ee.virginia.edu/toolbelt.html>



Basic Directions

- Each zipped file is a directory with code and a sample data set
 1. Unzip the directory
 2. Open Matlab and change to that directory
 3. If a file named "compile.m" exists, type:

compile (at the Matlab prompt)

*and press enter... I don't need to say that, do I?!

(This file will automatically compile any C/C++ code – installation of compilers may be necessary for your machine. You may have to run "mex -setup" if you've never used mex before. You may need help getting this started if you're not the programming geek type!)



Basic Directions

4. If you are using a PC, type

example

To run the example code.

On a Mac, type

examplemac



Tools on the Toolbelt

- **Active contour segmentation** through the Active Model Toolbox (**AMT**) – parametric active contour segmentation for 2D and 3D data; plus initialization by **PIG!**
- **Active contour tracking**: using snakes to track cells in 2D videos
- **Diffusion**: anisotropic diffusion for enhancing and segmenting biological images; **SRAD** for signal-dependent noise



Tools on the Toolbelt

- **Neuron Segmentation**: a graph theoretic method for segmenting filamentous objects in 3D
- **Particle filter tracking**: tracking cells with a Monte Carlo approach
- **Kalman filter tracking**: using the KF to track cells in a constant velocity model



Caveats and Additions

- Many of the tools in the toolbelt need
 - Matlab's image processing toolbox
- The neuron segmentation tool needs
 - Matlab's bioinformatics toolbox
 - A 3D Viewer – We recommend Hanchuan Peng's V3D: www.vaa3d.org
- The GVF implementation in the active contour segmentation tool is due to [Dr. Jerry Prince's](#) lab, used with their permission



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