

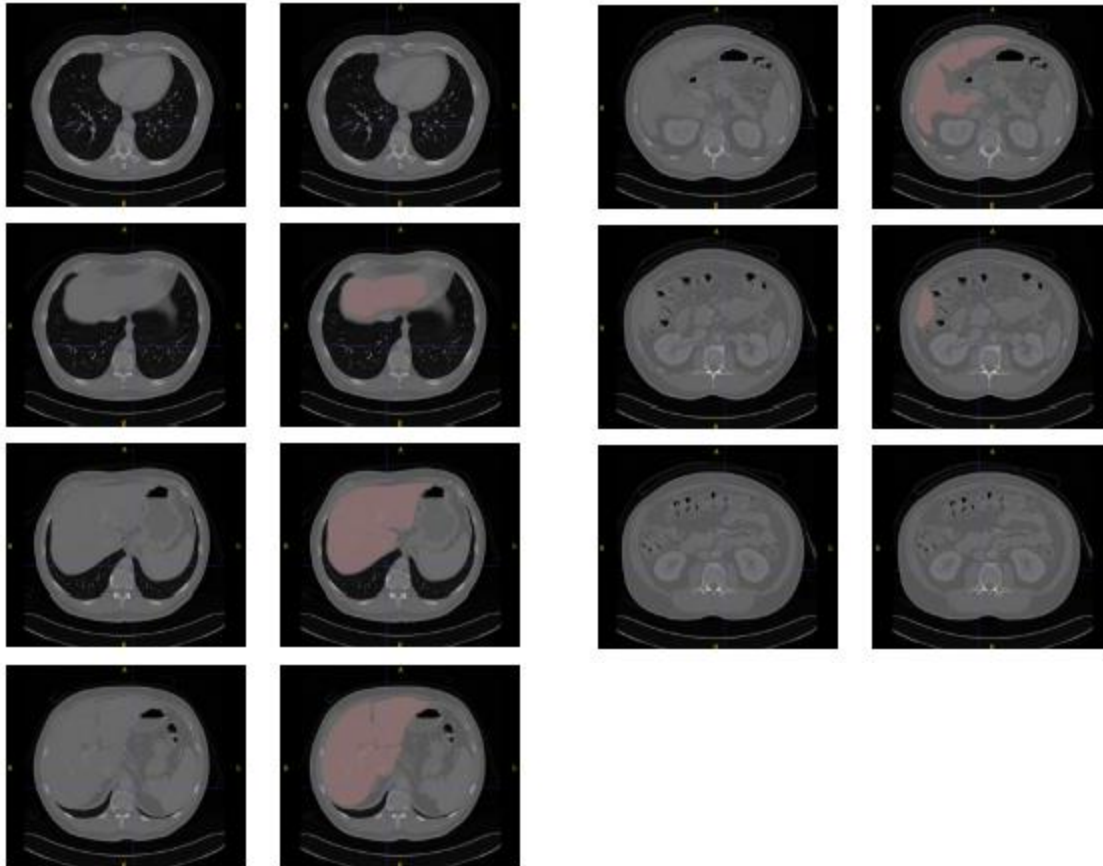
Model-Based Organ Segmentation: Recent Methods

Jiun-Hung Chen
General Exam Paper
2009

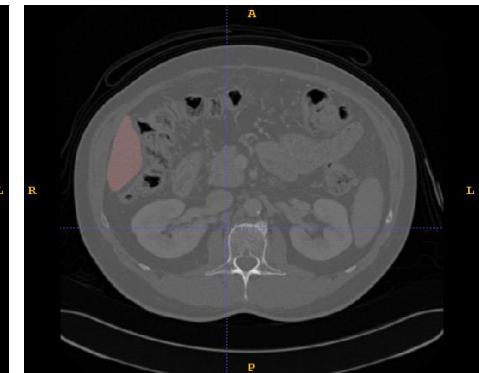
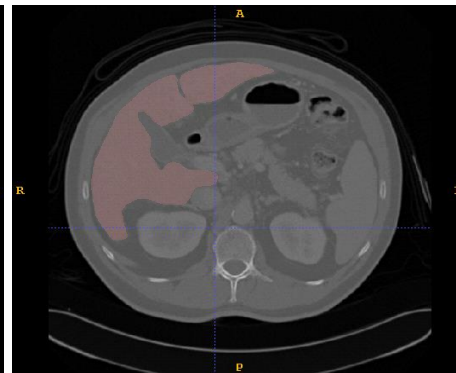
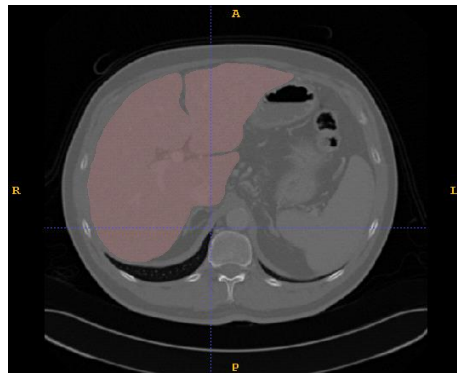
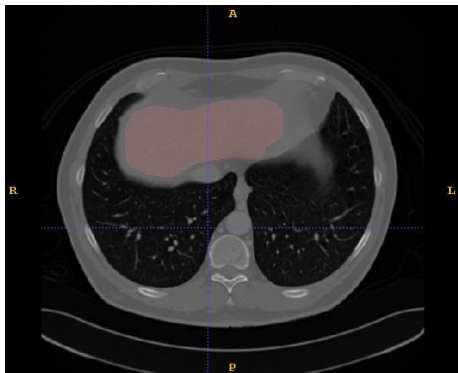
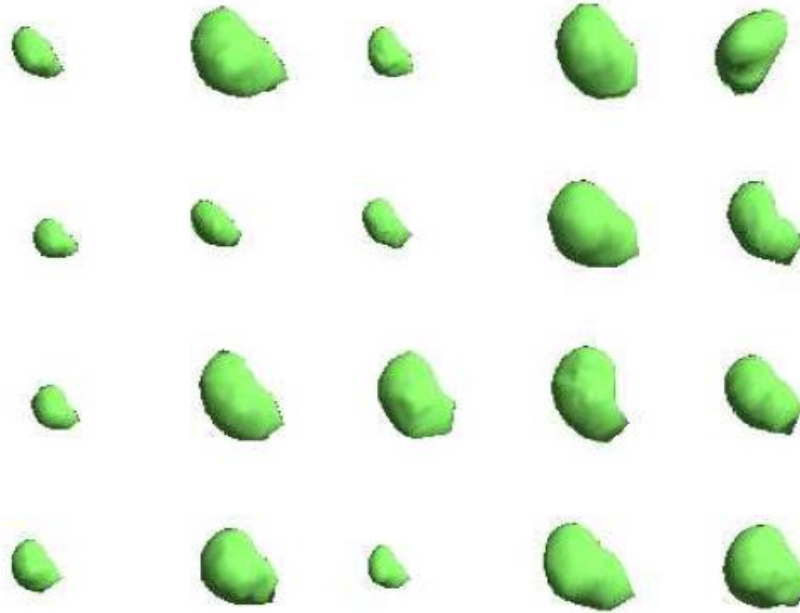
Problem Statement

- Learn how to segment new, unseen CT images from a set of training CT images with ground truth organs marked.
- Goal: Minimize the training errors while generalizing to the new CT images

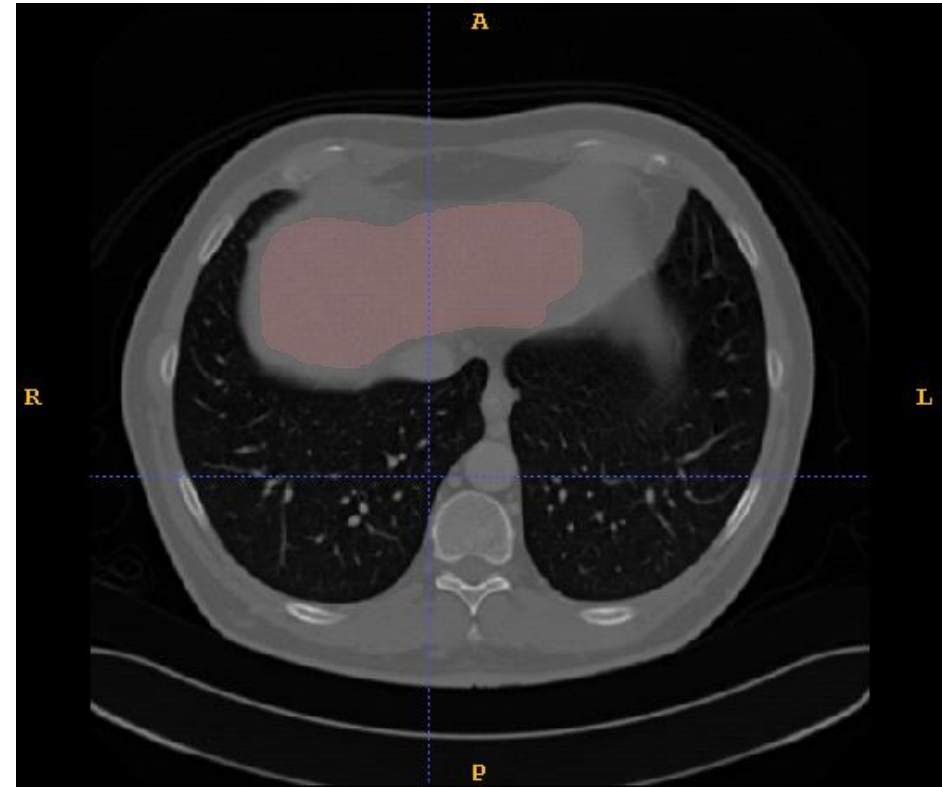
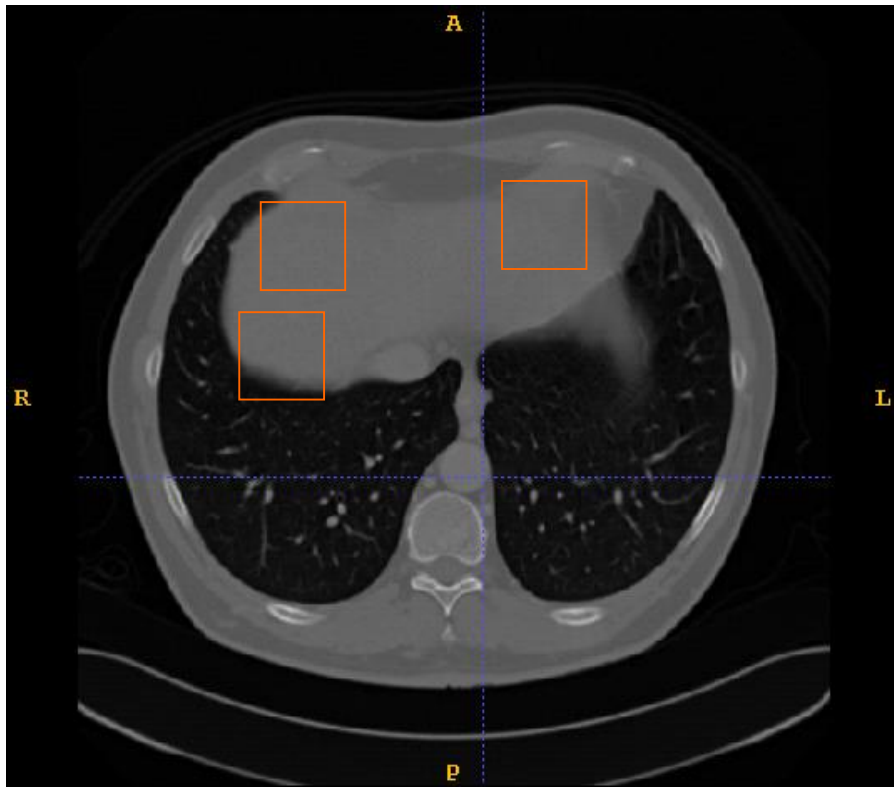
Problem Organ: The Liver



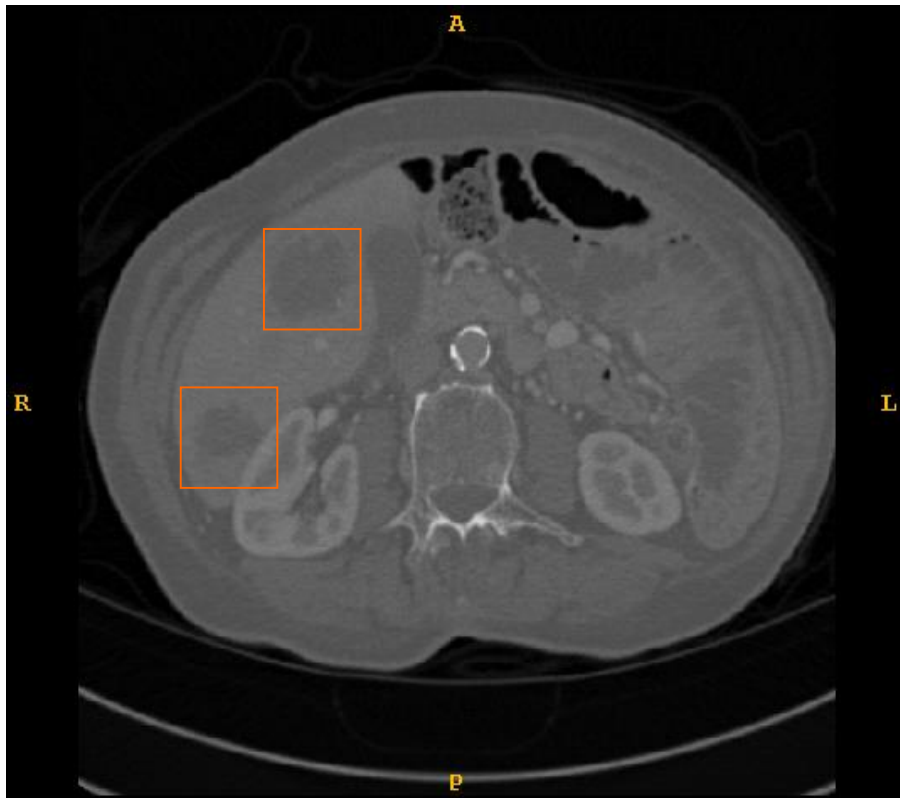
Why Difficult? (Shape Variations)



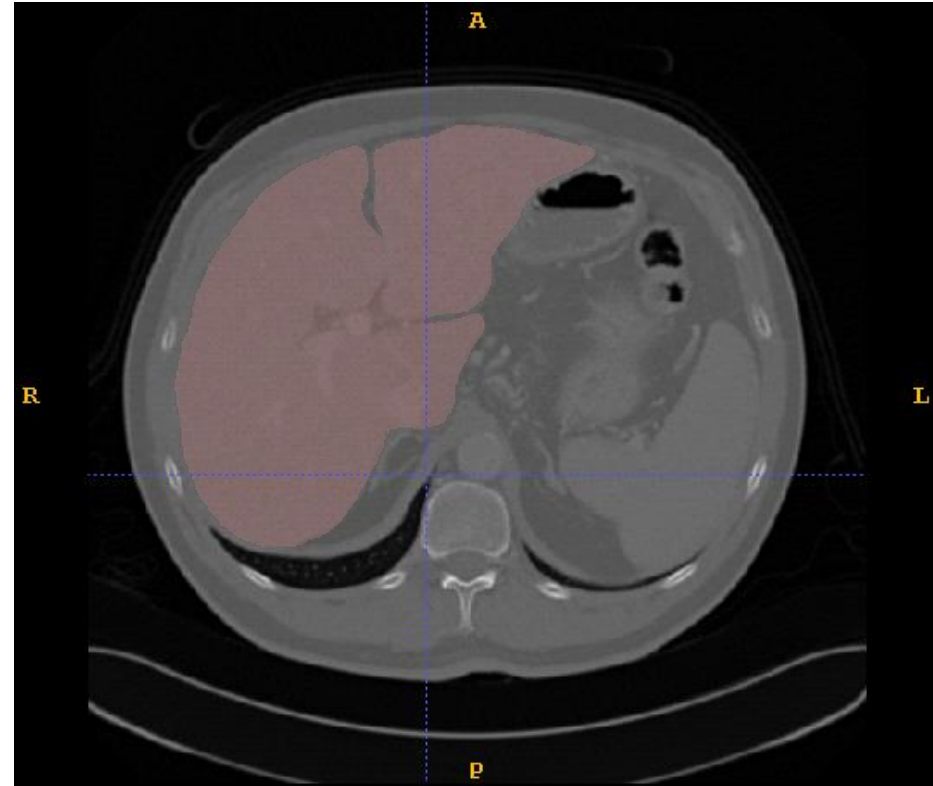
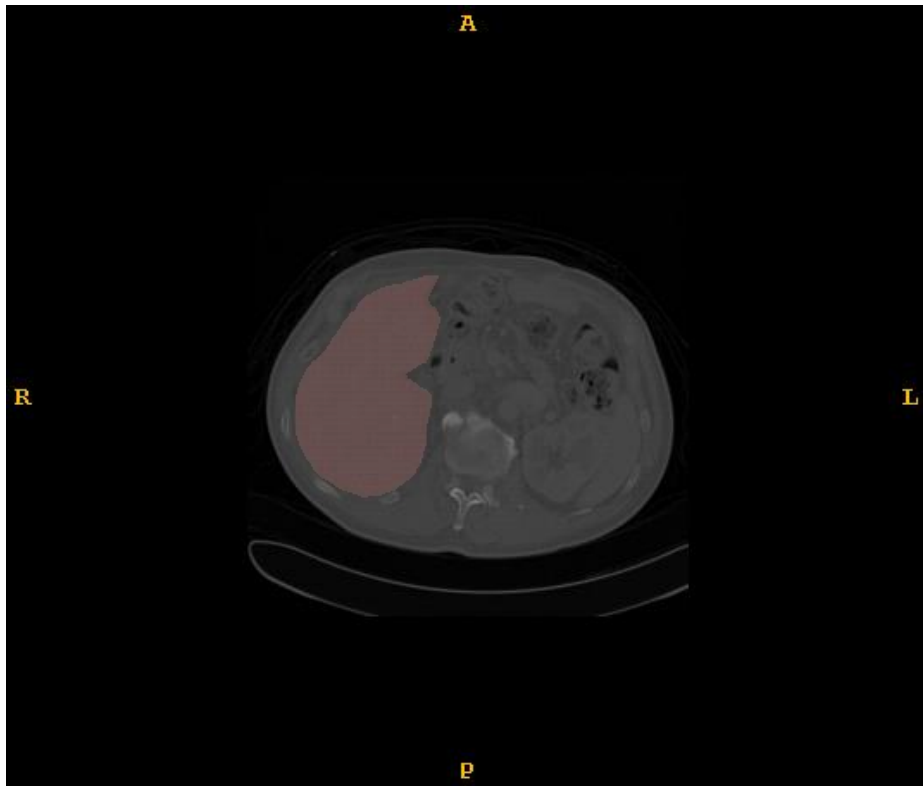
Why Difficult? (Similar Appearances)



Why Difficult? (Appearance Changes)

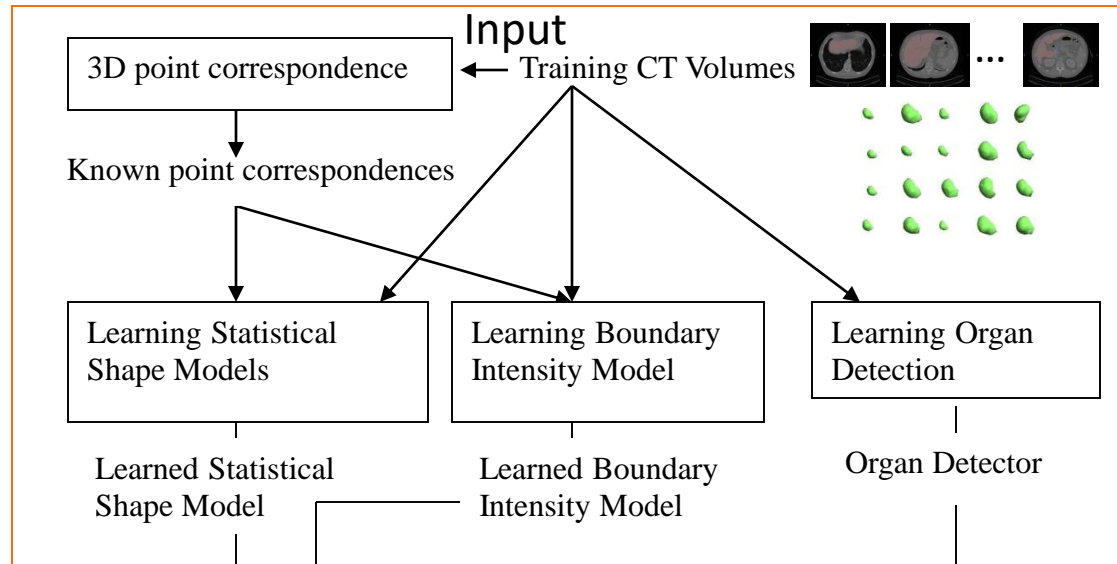


Why Difficult? (Position Changes)

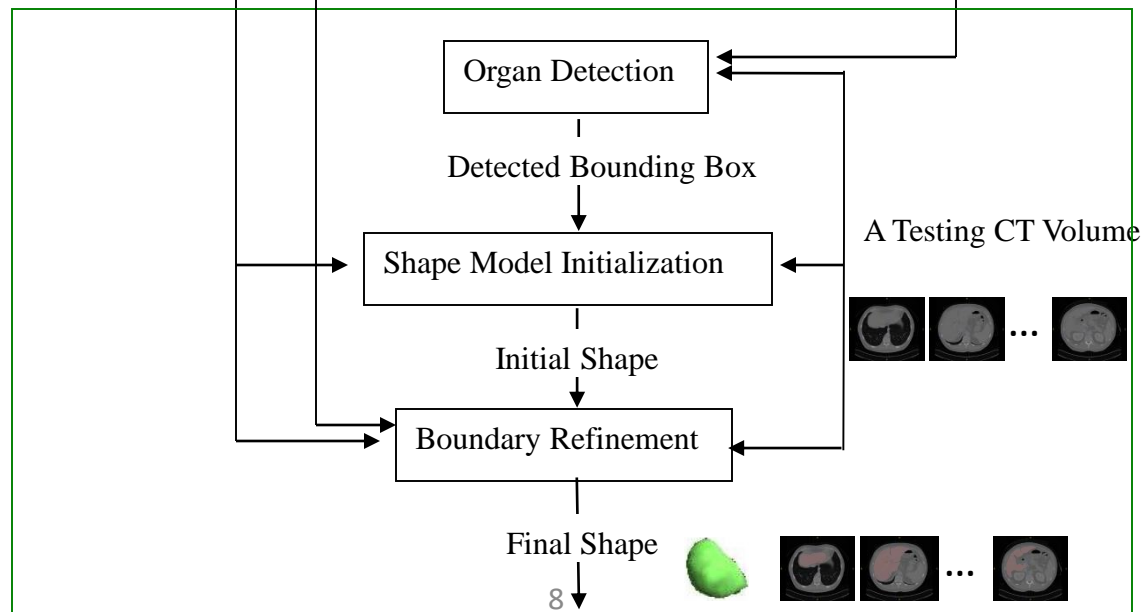


Active Shape Model Based Framework

Training phase



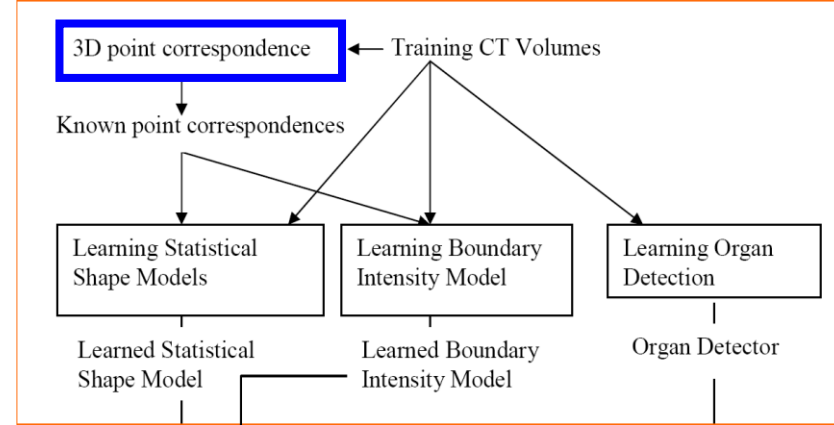
Testing phase



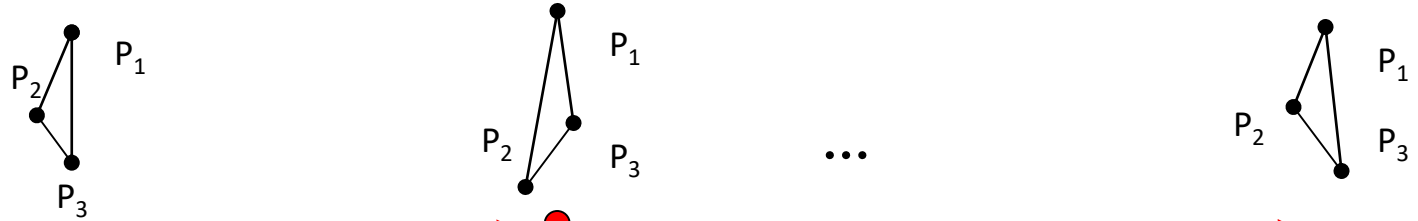
Active Shape Models: Training

- Shapes are modeled in a training phase using a set of CT volumes whose ground truth segmentations are given.
- There are 4 steps to the training phase.
 1. Find **3D point correspondences** on training meshes.
 2. Learn a **statistical 3D shape model** of the shapes.
 3. Learn a **boundary intensity model** for each vertex.
 4. Learn an **organ detector** that finds bounding boxes.

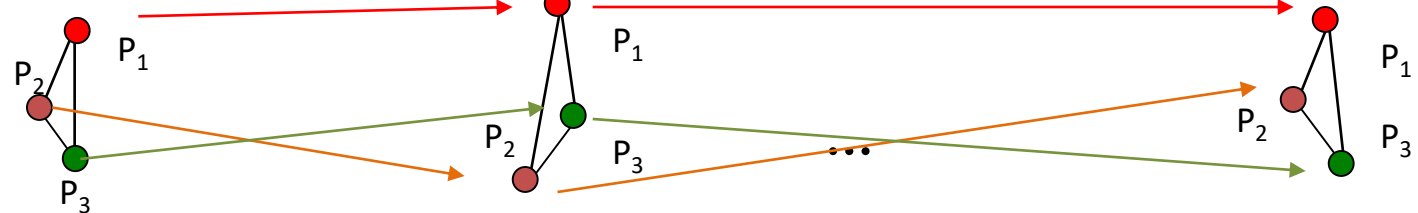
First find 3D point correspondences
Among all the training models of
The organ being modeled.



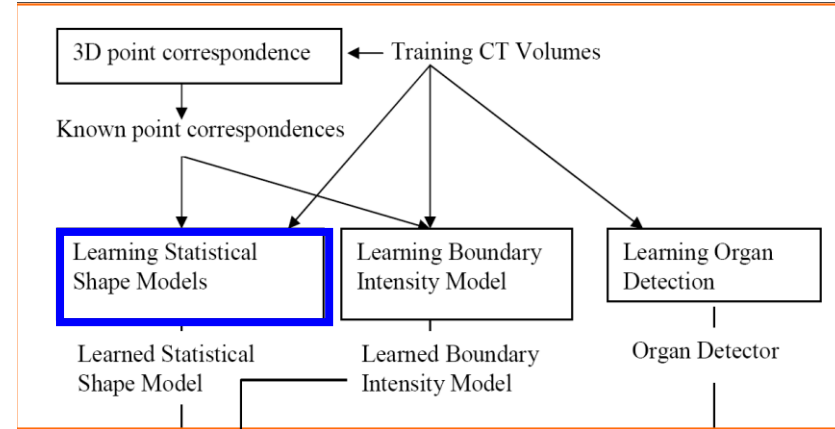
Input



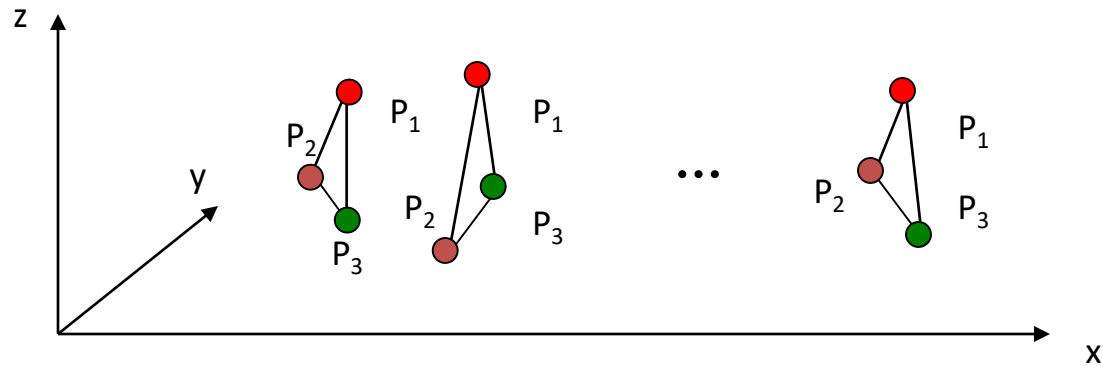
Output



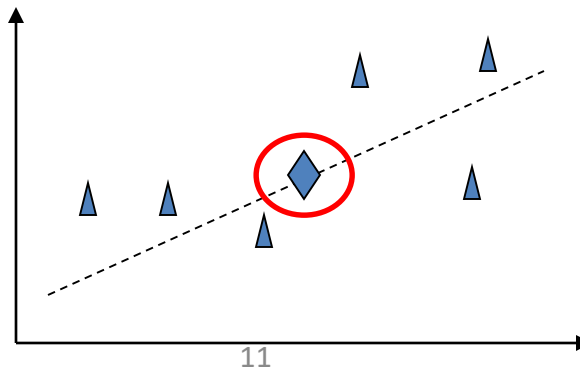
Next learn a statistical shape model of that organ.



Input

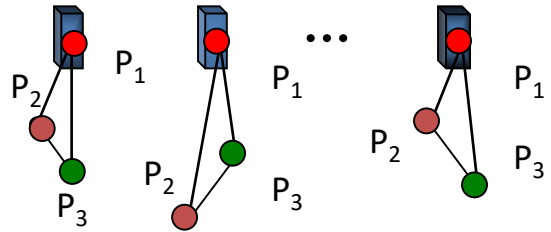


Output



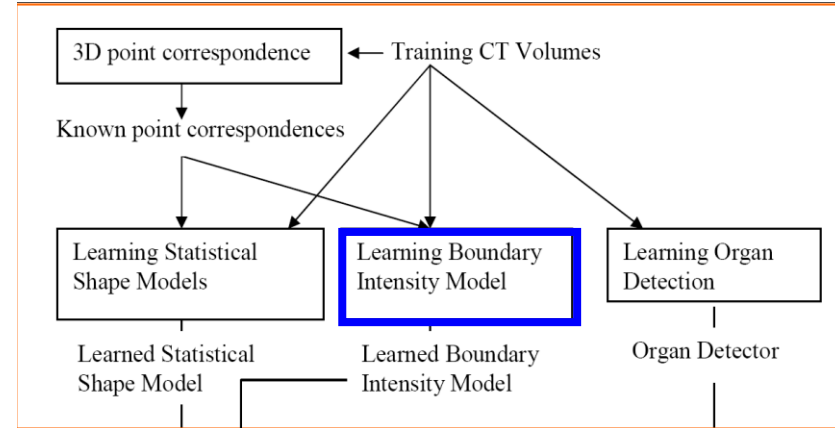
simplified
linear model

Next learn a model of the boundary intensities.

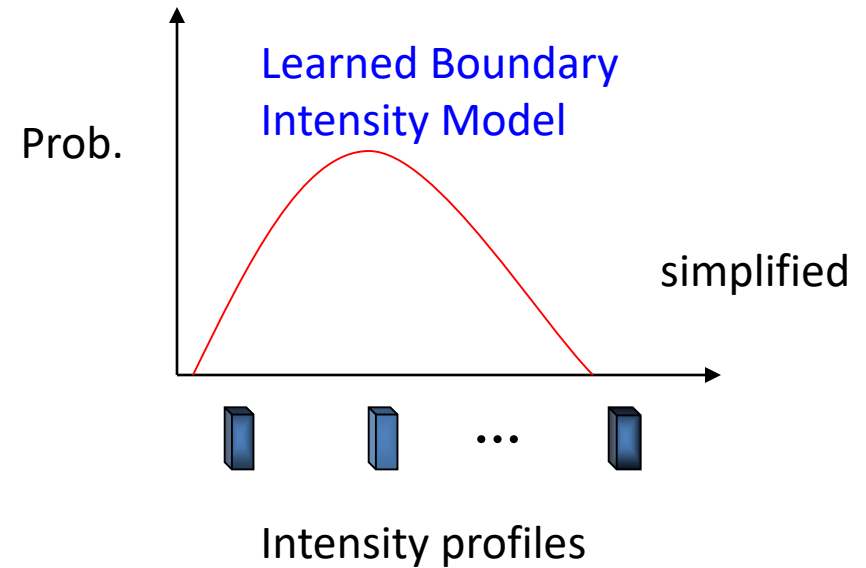
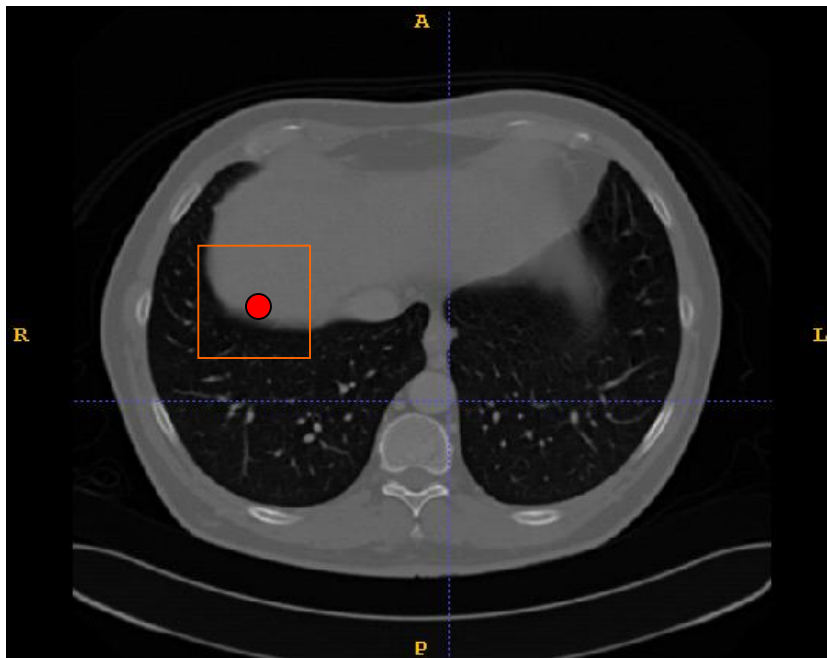


Intensity profiles

Input ●

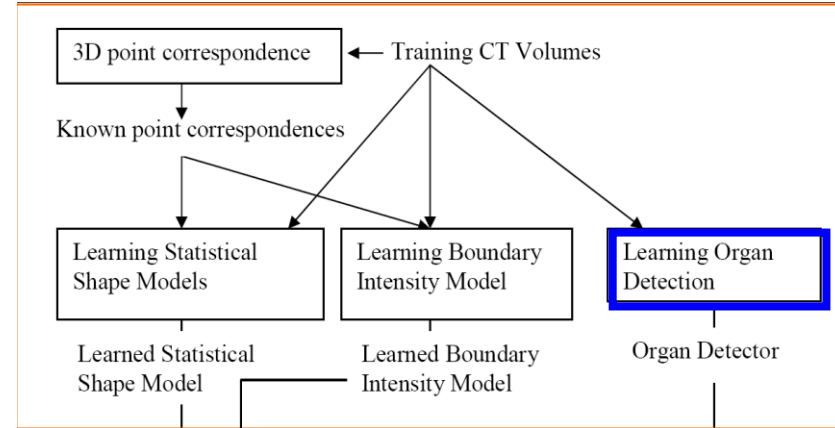


Output ●

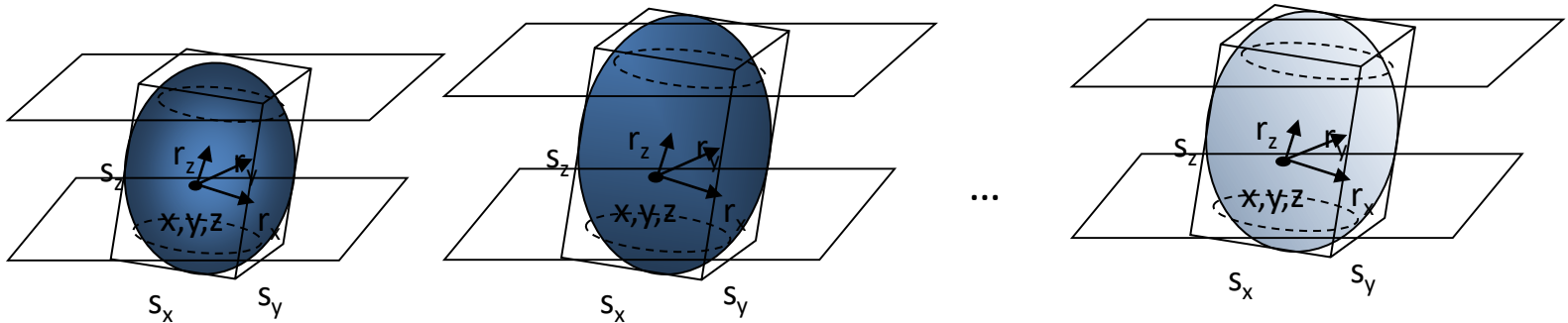


Finally train a classifier to find the organ inside a bounding box.

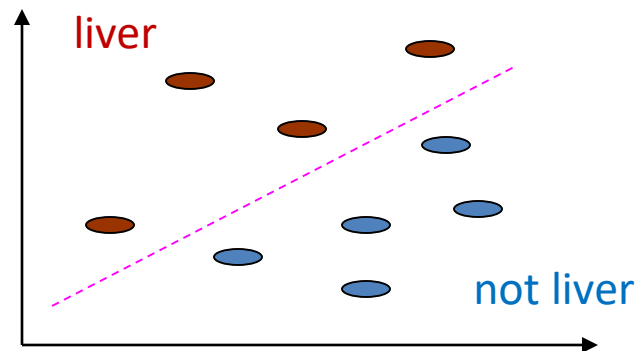
- Given a bounding box and the CT slices inside it, a classifier learns to decide if *everything* inside the box is liver or not.



Input



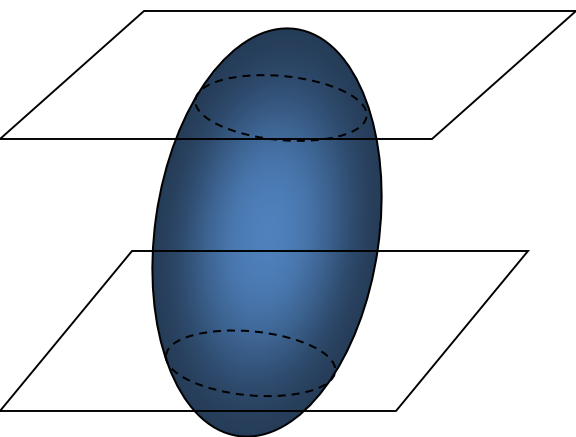
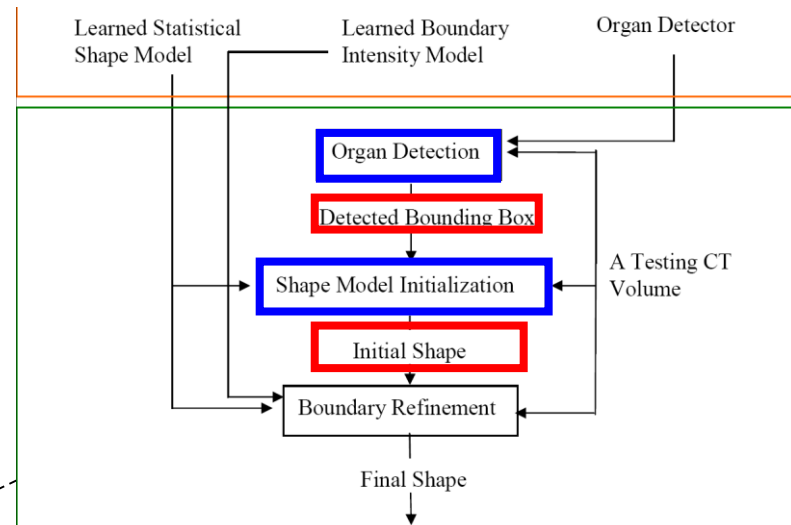
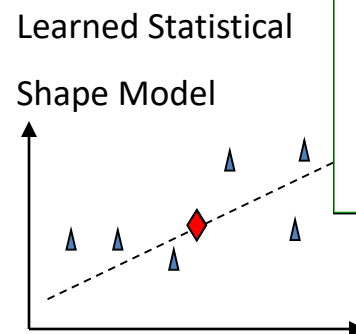
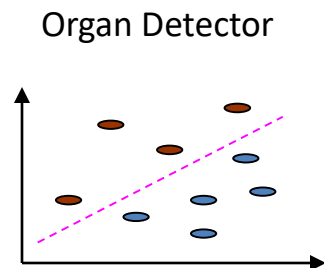
Output



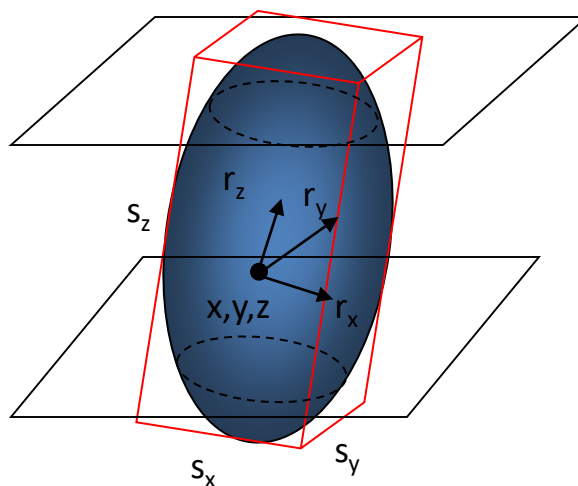
Where do you get the bounding box?

Active Shape Models: Testing

- There are 3 steps to the testing phase
 1. **organ detection:** use the learned organ detector to detect the organ in the testing volume and return a bounding box
 2. **shape model initialization:** initialize the learned statistical model based on the detected bounding box
 3. **boundary refinement:** use the learned boundary intensity model to estimate the refinement to the model for this shape

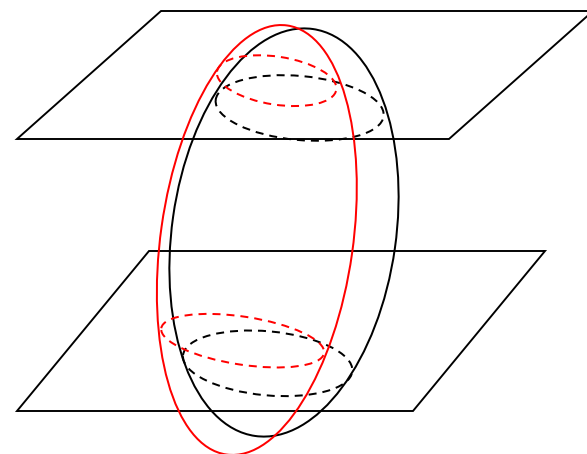


Test Organ



Find Bounding Box

Initialize Shape Model



Boundary Refinement

Methods for Point Correspondences

1. Principal Component Analysis (PCA)

PCA takes in the points of each shape in the training set. It produces a set of basis vectors (the components).

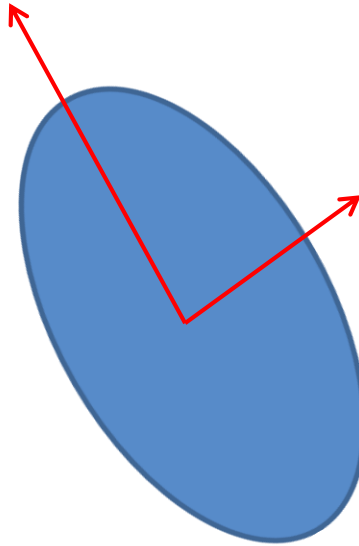
Each shape can then be represented as a linear combination of these components.

$$\tilde{\mathbf{x}} = \bar{\mathbf{x}} + \sum_{k=1}^K c_k \mathbf{b}_k \text{ where } \bar{\mathbf{x}} \text{ is the mean shape}$$

The optimal K projection axes \mathbf{b}_k , $k = 1$ to K are the **eigenvectors of the covariance matrix** of the training set of points corresponding to the **K largest eigenvalues**.

Intuitive Meaning of Principal Components

eigenvector corresponding to highest
eigenvalue



eigenvector corresponding to second
eigenvalue

Eigenimages for Face Recognition

training
images

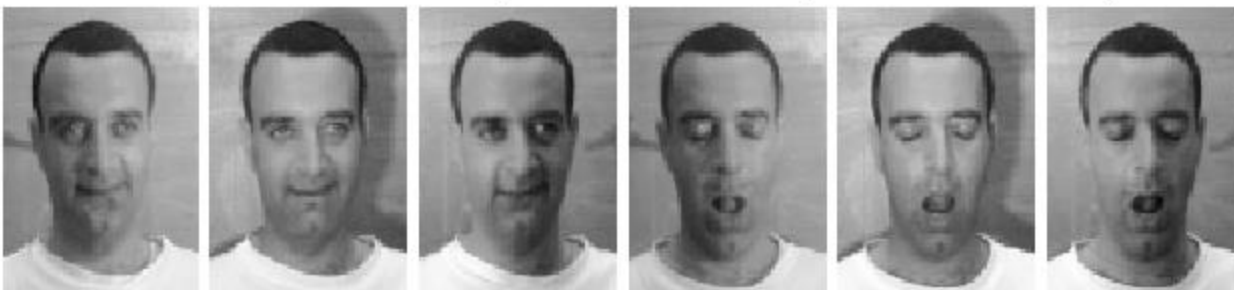


mean
image



3 eigen-
images

linear
approx-
imations

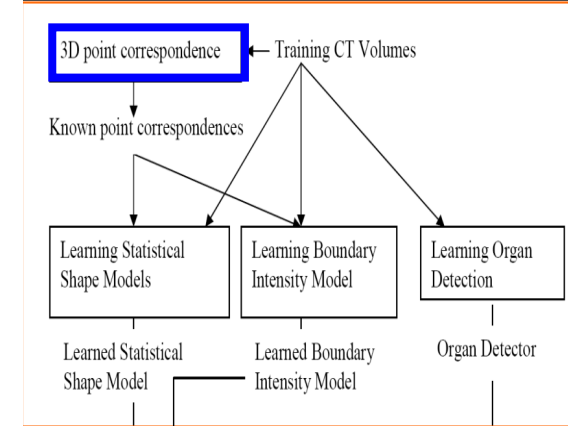


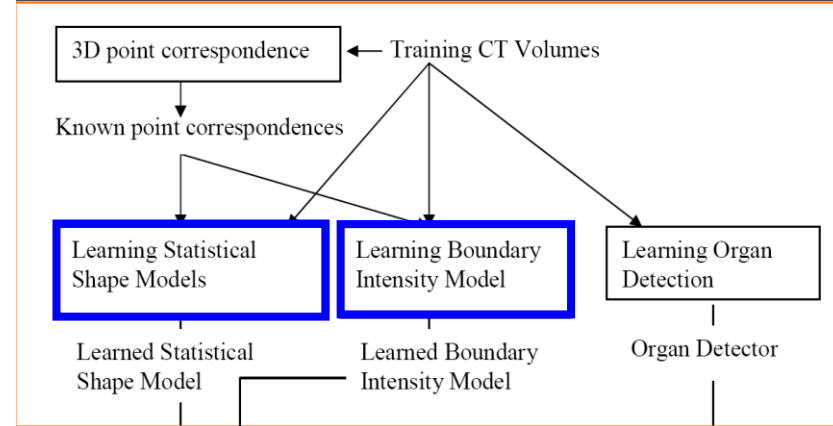
3D Point Correspondence (MDL)

- Goal: Find 3D Point Correspondence
- Idea: **Minimize MDL-based objective function**
 - Evaluate the quality of the correspondence

$$F = \sum_{k=1}^N L_k \text{ with } L_k = \begin{cases} 1 + \log(\lambda_k / \lambda_{cut}), & \text{if } \lambda_k \geq \lambda_{cut} \\ \lambda_k / \lambda_{cut}, & \text{otherwise} \end{cases}$$

- The λ_k s are the **eigenvalues** from PCA.
- How: Gradient descent
 - Manipulate correspondences by parameterization and re-parameterization.





- Statistical Shape Models



- Principal Component Analysis (PCA)
- Kernel PCA

In either case the shape model consists of the PCA mean and basis. Any shape can be represented.

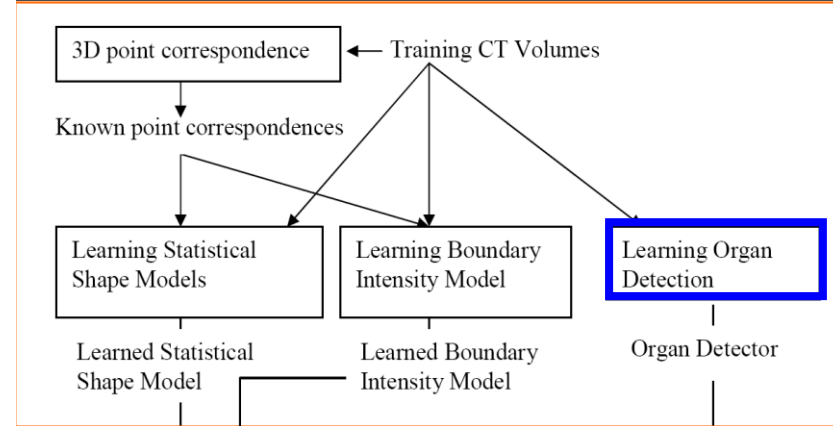
- Boundary Intensity Models

- ☐ Gaussian distribution
- ☐ AdaBoosted histogram classifiers
- ☐ Heuristics

Cootes et al. [IEEE PAMI 01], Twining et al. [BMVC'01]

Cootes et al. [IEEE PAMI 01], Li [ICCV'05], Kainmuller et al. [MICCAI'07]

Organ Detection (MSL)



- Goal: Find the bounding box
 - The parameter space is 9D.
 - 3D positions, 3D scales and 3D orientations.
- Idea
 - Uniform and exhaustive search is unnecessary
- How: decompose the problem into three steps
 - position estimation, position-scale estimation and finally position-scale-orientation estimation.

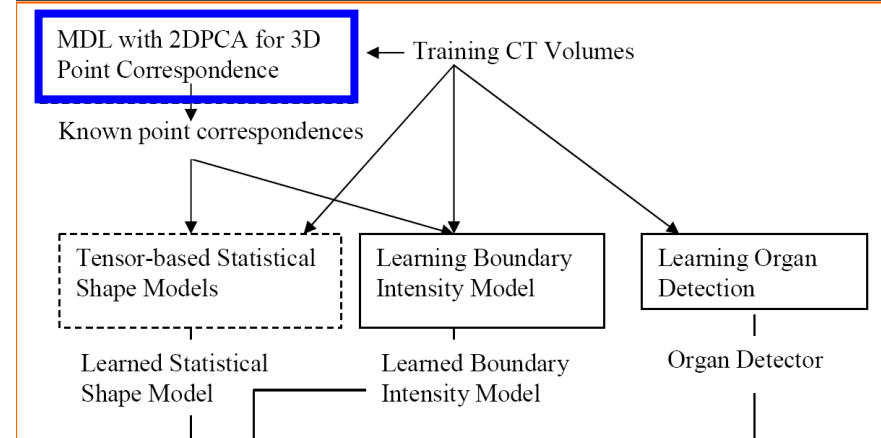
Two ASM-based Systems

Kainmuller et al. [MICCAI'07]	Ling et al. [CVPR'08]
Statistical Shape Models	
<ul style="list-style-type: none"> – PCA – 43 CT volumes 	<ul style="list-style-type: none"> – PCA, hierarchical shape pyramids – 75 volumes
Boundary Intensity Model	
<ul style="list-style-type: none"> – Heuristics 	<ul style="list-style-type: none"> – A boundary classifier
Liver Detection	
<ul style="list-style-type: none"> – Lungs detection and DICOM info 	<ul style="list-style-type: none"> – MSL (marginal space learning)
Performance	
<ul style="list-style-type: none"> – Ranked first in a recent liver segmentation competition. – 10 testing volumes – 1.1mm (the average symmetric surface distance) – 15 minutes. 	<ul style="list-style-type: none"> – 5 fold cross validation – 1.59 mm (the average symmetric surface distance) – 1.38 mm (the median) – 12 seconds.

Experimental Setting

- Datasets:
 - 4 types of organs (livers, left kidneys, right kidneys, spleens)
 - 15-20 subjects
- Leave-one-out cross validation
- Measure the reconstruction error
- Metrics: Euclidean and Hausdorff distance

MDL-2DPCA



- MDL-based objective function

$$F = \sum_{k=1}^N L_k \text{ with } L_k = \begin{cases} 1 + \log(\lambda_k / \lambda_{cut}), & \text{if } \lambda_k \geq \lambda_{cut} \\ \lambda_k / \lambda_{cut}, & \text{otherwise} \end{cases}$$

- Idea: Generalize the objective function to 2DPCA space
 - Replace eigenvalues from PCA with those from 2DPCA
 - How: Gradient descent
- Comparisons: original MDL vs. MDL-2DPCA

2DPCA

- When we use normal PCA, our representation of a shape is a vector

$$(x_1, y_1, z_1, x_2, y_2, z_2, \dots, x_K, y_K, z_K)^T$$

- When we use 2DPCA, our representation of a shape is a 2D matrix (or tensor)

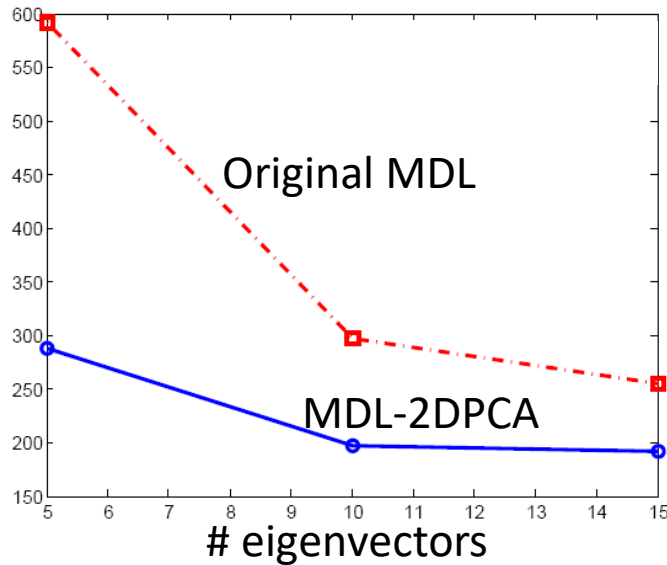
$$\begin{bmatrix} x_1 & y_1 & z_1 \\ x_2 & y_2 & z_2 \\ x_K & y_K & z_K \end{bmatrix}$$

- And so are the components.

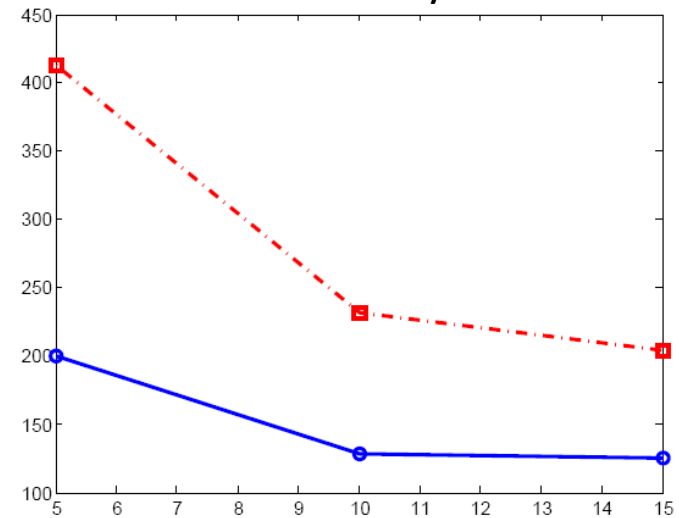
Results (3D Point Correspondences)

Reconstruction error

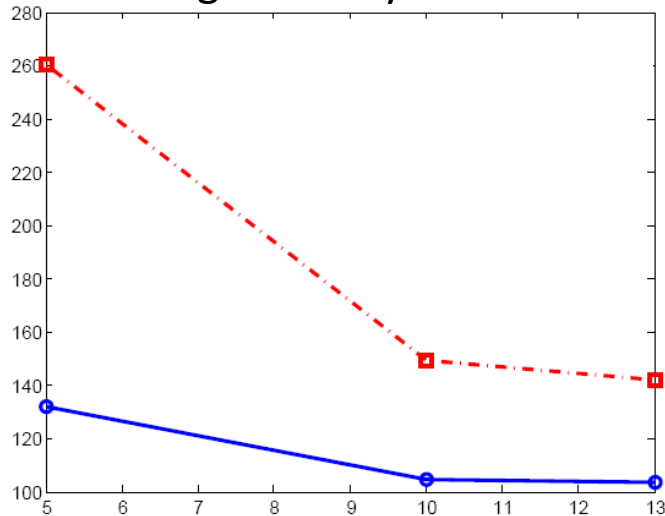
Livers



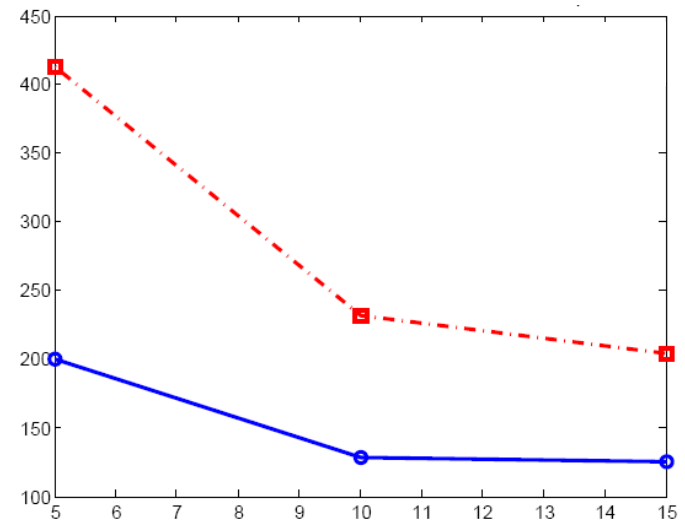
Left Kidneys



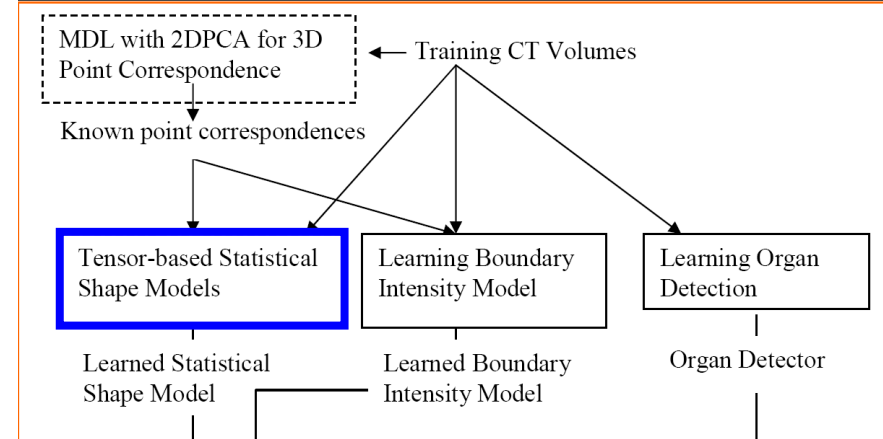
Right Kidneys



Spleens



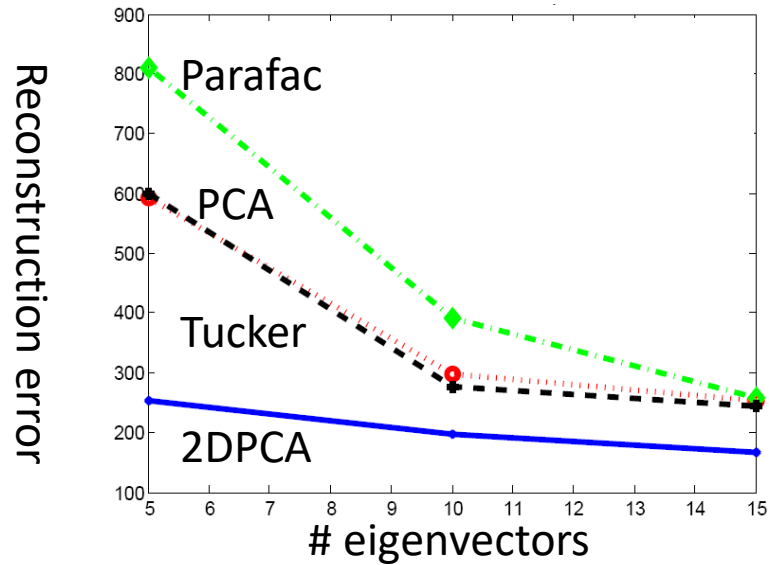
Tensor-based SSM



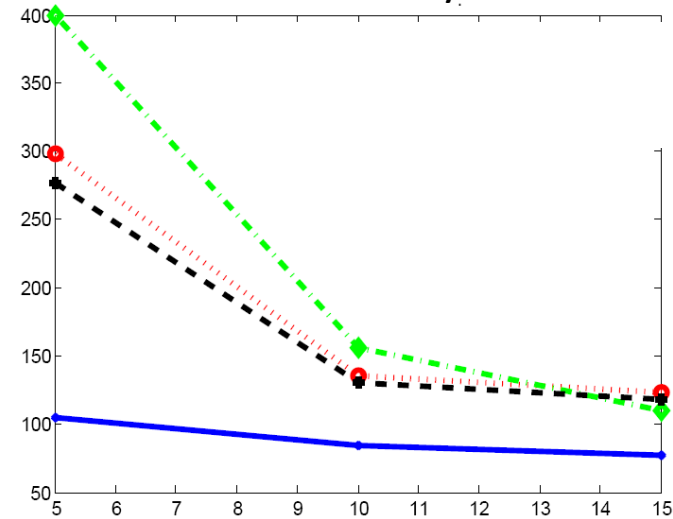
- Idea: Tensor-based dimension reduction methods
 - 2DPCA
 - Parafac model
 - Tucker decomposition
- Comparisons: PCA vs. Tensor-based dimension reduction

Results (Statistical Shape Models)

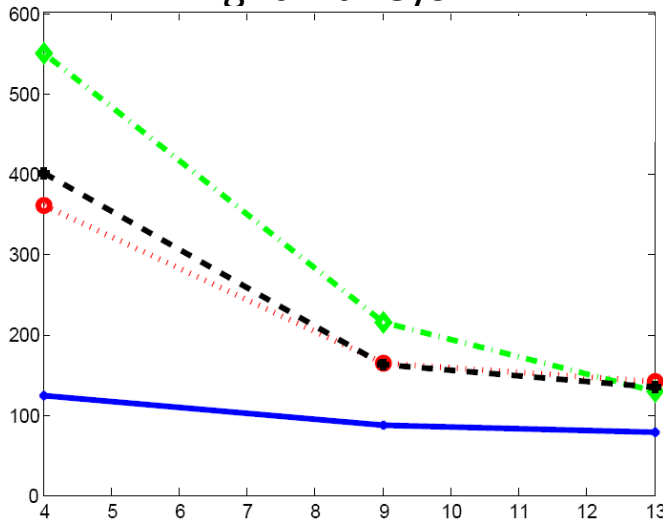
Livers



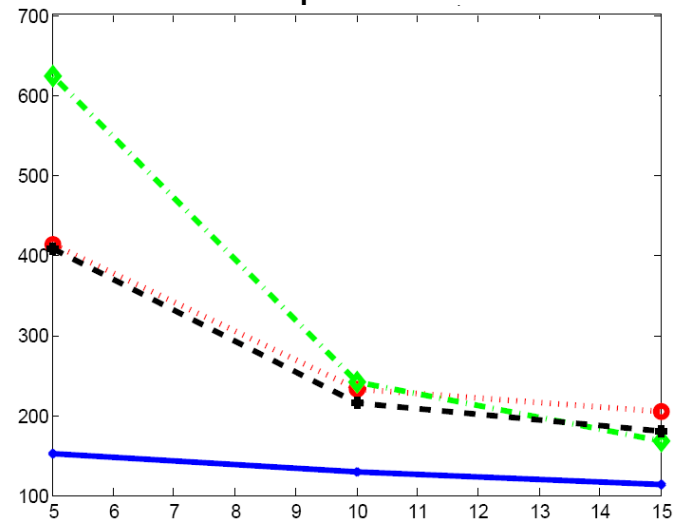
Left Kidneys



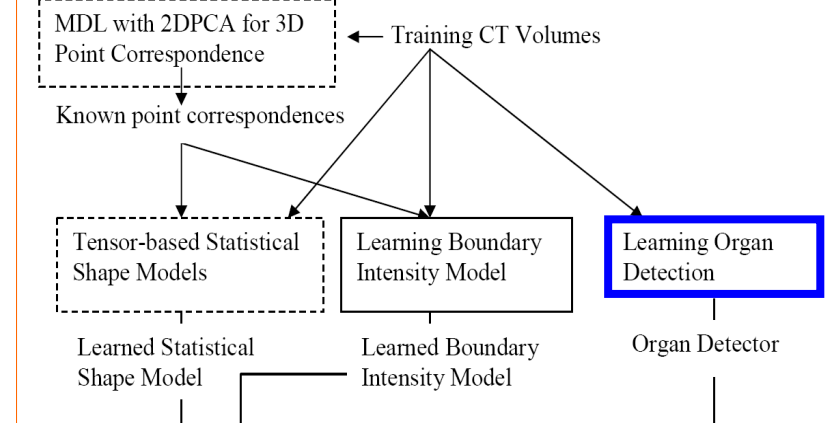
Right Kidneys



Spleens



Organ Detection (Boosting Approach)



- Idea: Classify whether an image block contains an organ of interest
- How:
 - Partition slices into non-overlapping 32x32 blocks
 - Global features: gray-tone histogram of the image slice and its slice index
 - Local features: the position of a block, the mean and variance of its intensity values, and its intensity histogram.
 - 20,000 SVM linear classifiers + Adaboosting
- Comparisons: Manual vs. Adaboosting

Results (Organ Detection)

	Positive (predicted)	Negative (predicted)
Positive (actual)	96.23%	3.77%
Negative (actual)	4.57%	95.43%

Livers (Training)

	Positive (predicted)	Negative (predicted)
Positive (actual)	91.23%	8.77%
Negative (actual)	6.57%	93.43%

Livers (Testing)

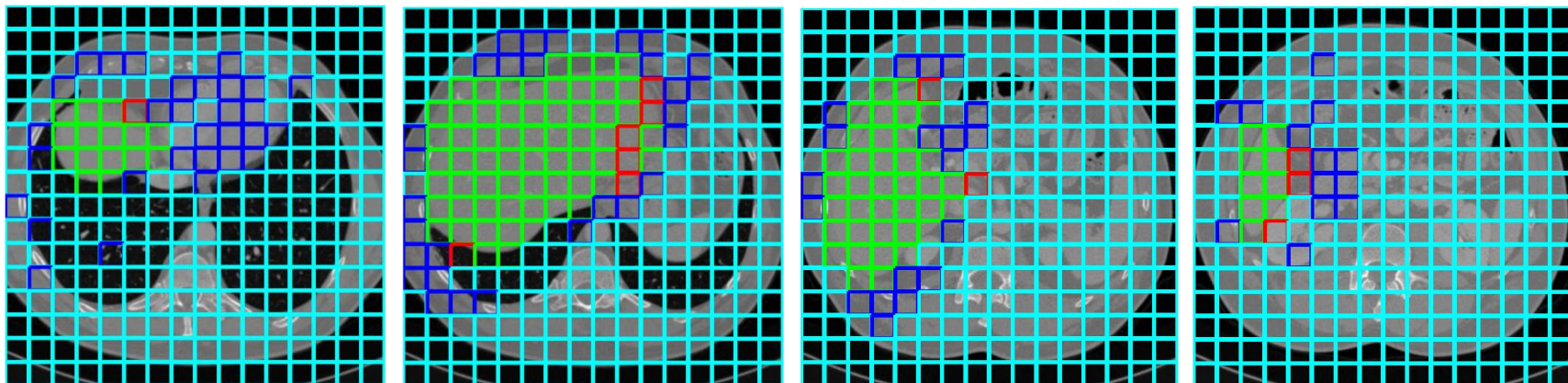
True Positive : Green,

False Positive: Blue

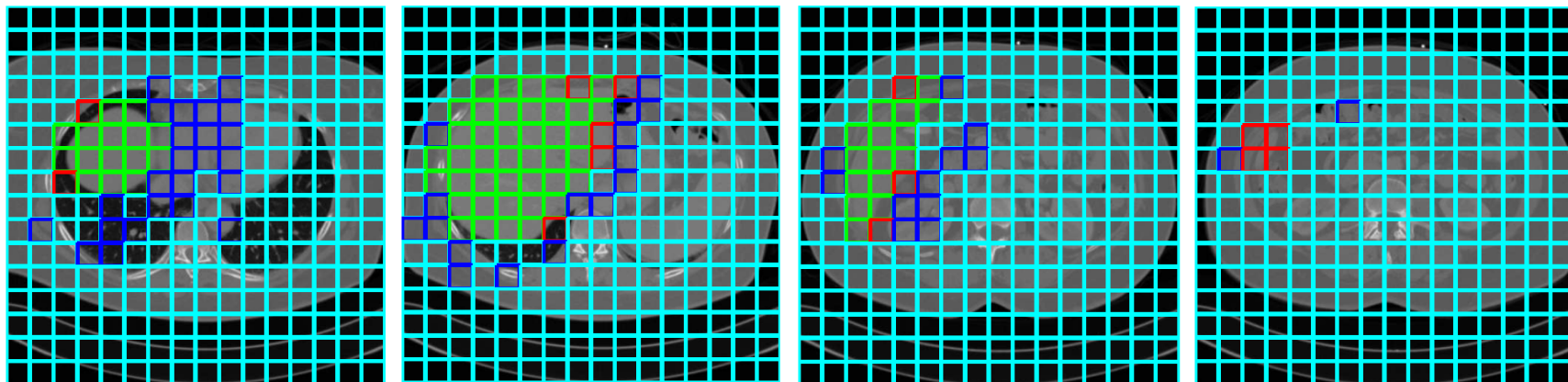
False Negative: Red,

True Negative: Cyan

Sub. 1



Sub. 2



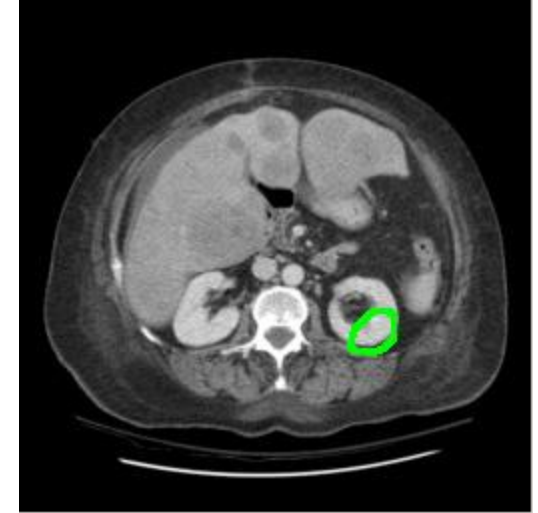
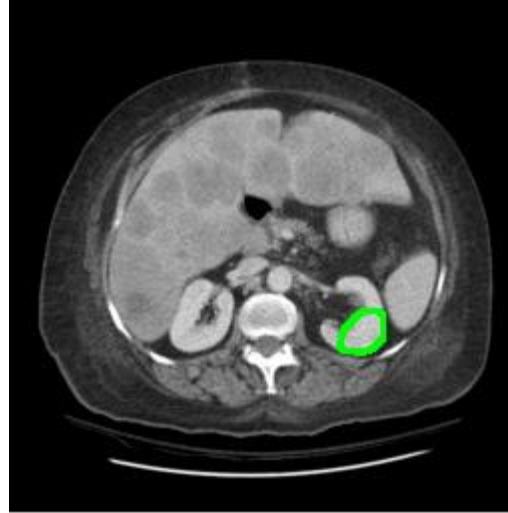
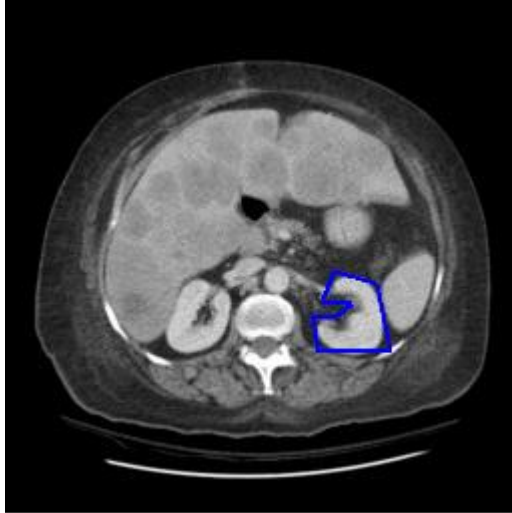
Results (Boundary Refinement)

Initial Contour

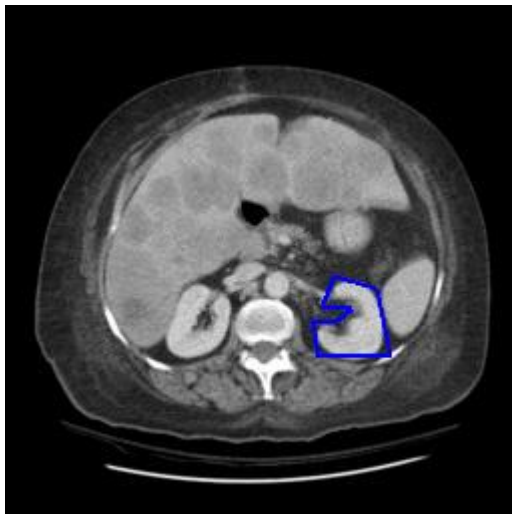
Slice 1

Slice 2

without



with



What have we got?

- A great design.
- Several conference papers.
- No complete system, just pieces, some pieces never really done.
- Would be a space for a course or longer project.
- Livers came from here: <http://sliver07.org/>
- Here is a newer one:
<https://www.virtualskeleton.ch/ShapeChallenge/Start2014>
- LOTS of data challenges here: https://grand-challenge.org/all_challenges/