

# Hierarchical Feature Matching for Shape Analysis

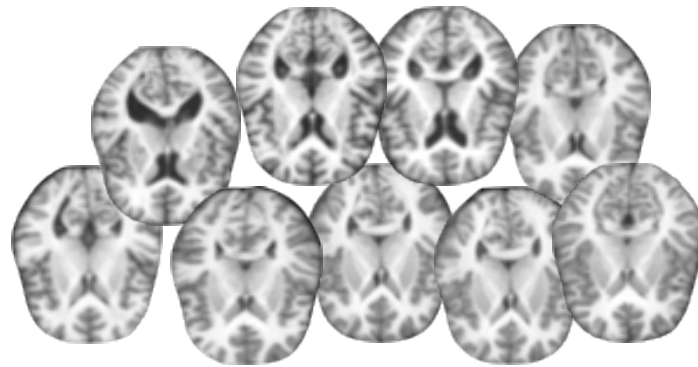
Ross Whitaker

BIRS - Aug 2011

# Some Recent Work

- Suppose we have lots of examples (images/shapes)

- 100s - 1000s



- Opportunities

- Possibly learn important shape characteristics
  - Chance of finding similar shapes as needed

- Challenges

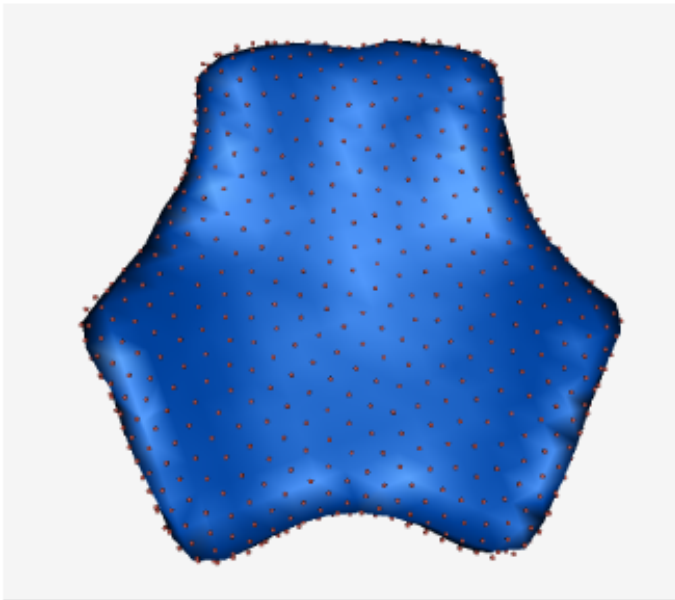
- Making systematic use of so much data
  - Quickly finding what you want

# The Talk I Had Planned

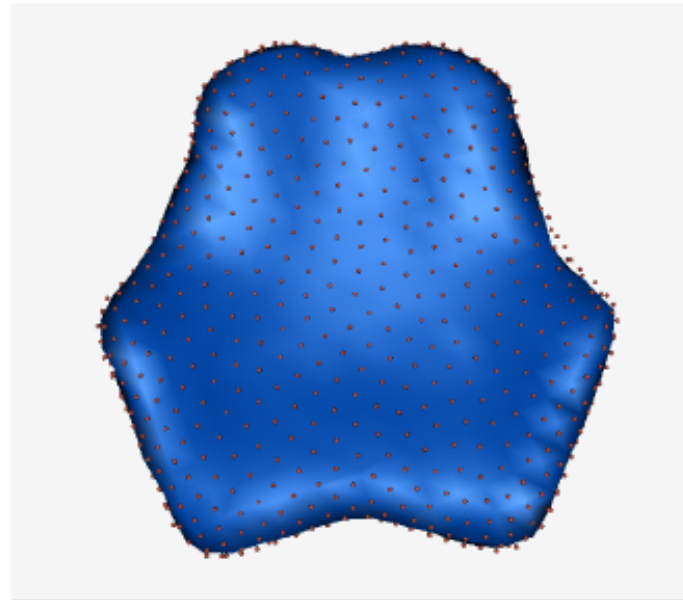
1. Statistics of ensembles for shape correspondence (Cates, Datar)
2. Learning manifolds of large collections of brain images (Gerber)
3. Hierarchical feature-based shape matching for fast neighbor lookup (Zhu)

**This Talk**

# Ensemble-Based Shape Correspondence

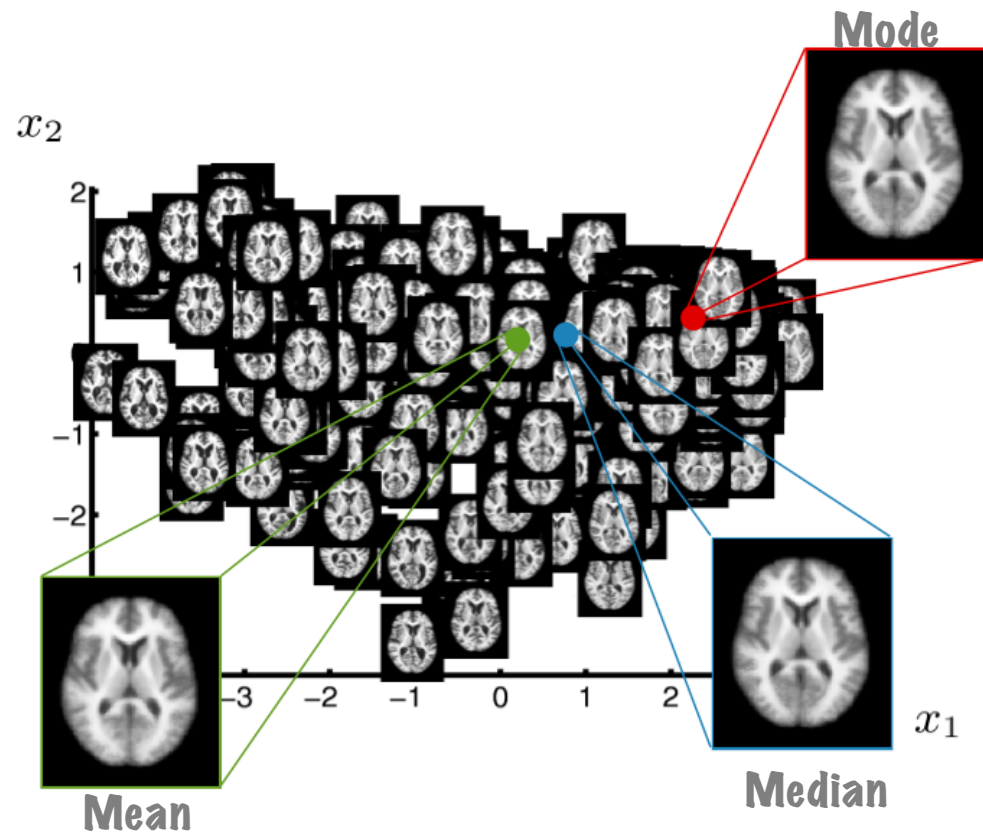
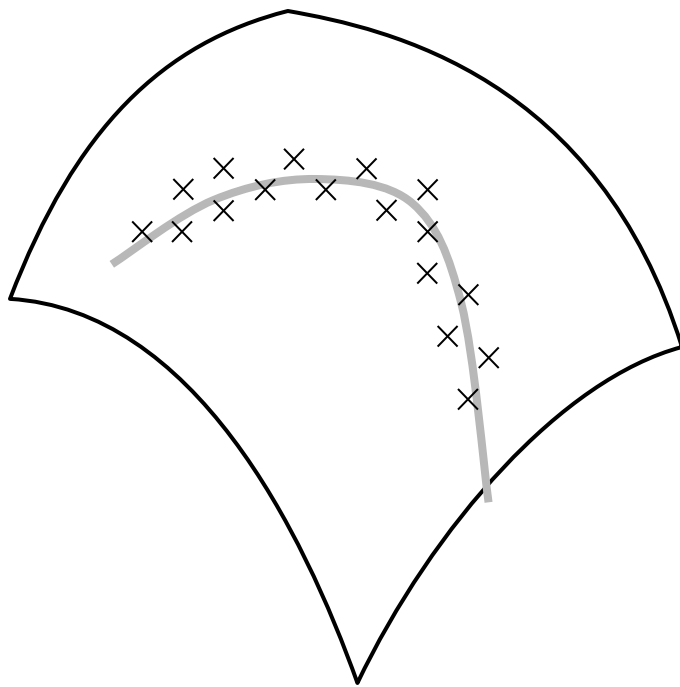


*Mean shape: LacZ -/-*



*Wild Type*

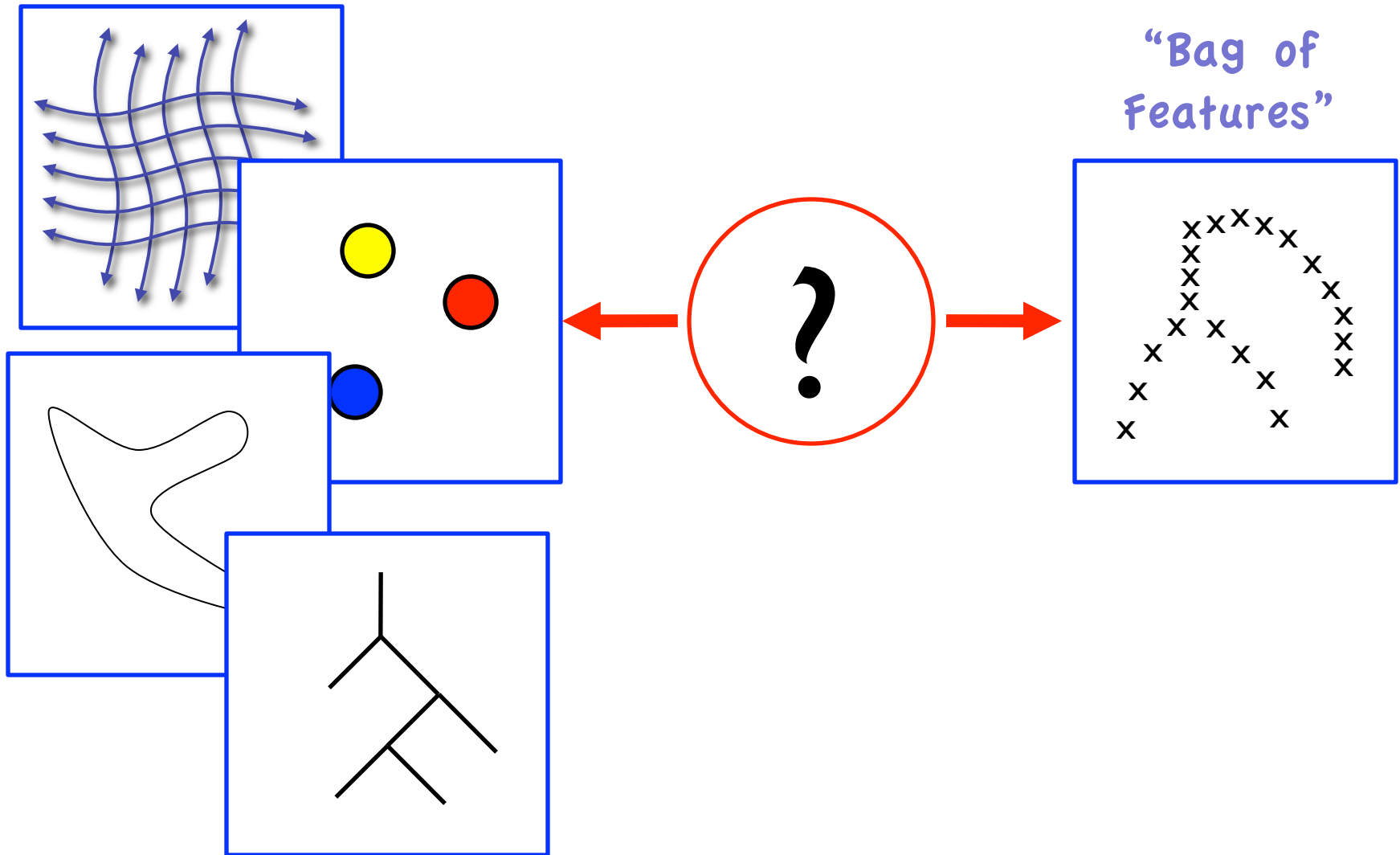
# Learned Manifolds of Brain Images



# Some Observations

- Shape analysis hinges on “correspondence”
- Shapes similar => correspondence easy
  - Shapes very different => hard (optimization)
- Ensembles help with correspondence
  - Statistical models regularize/constrain problem
  - Rely on nearest neighbors in shape space
- Roadblocks to analysis
  - Getting data into the correct “framework”
  - Optimizations and lack of generality

# Shape Representations

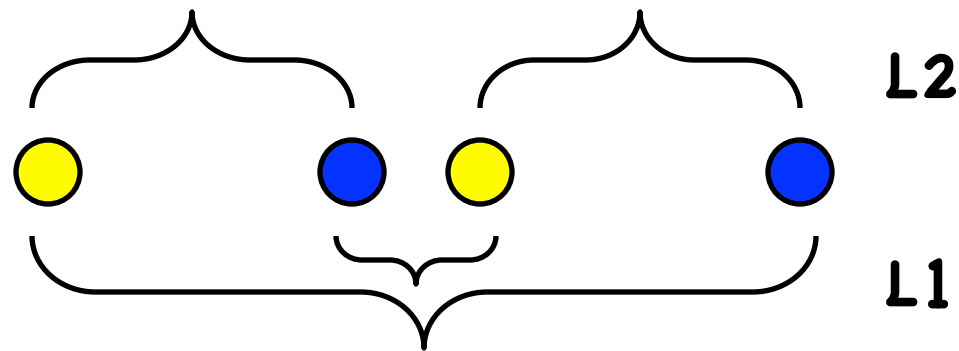


# Bipartite Matching

- Find the matches that minimize L1 norm

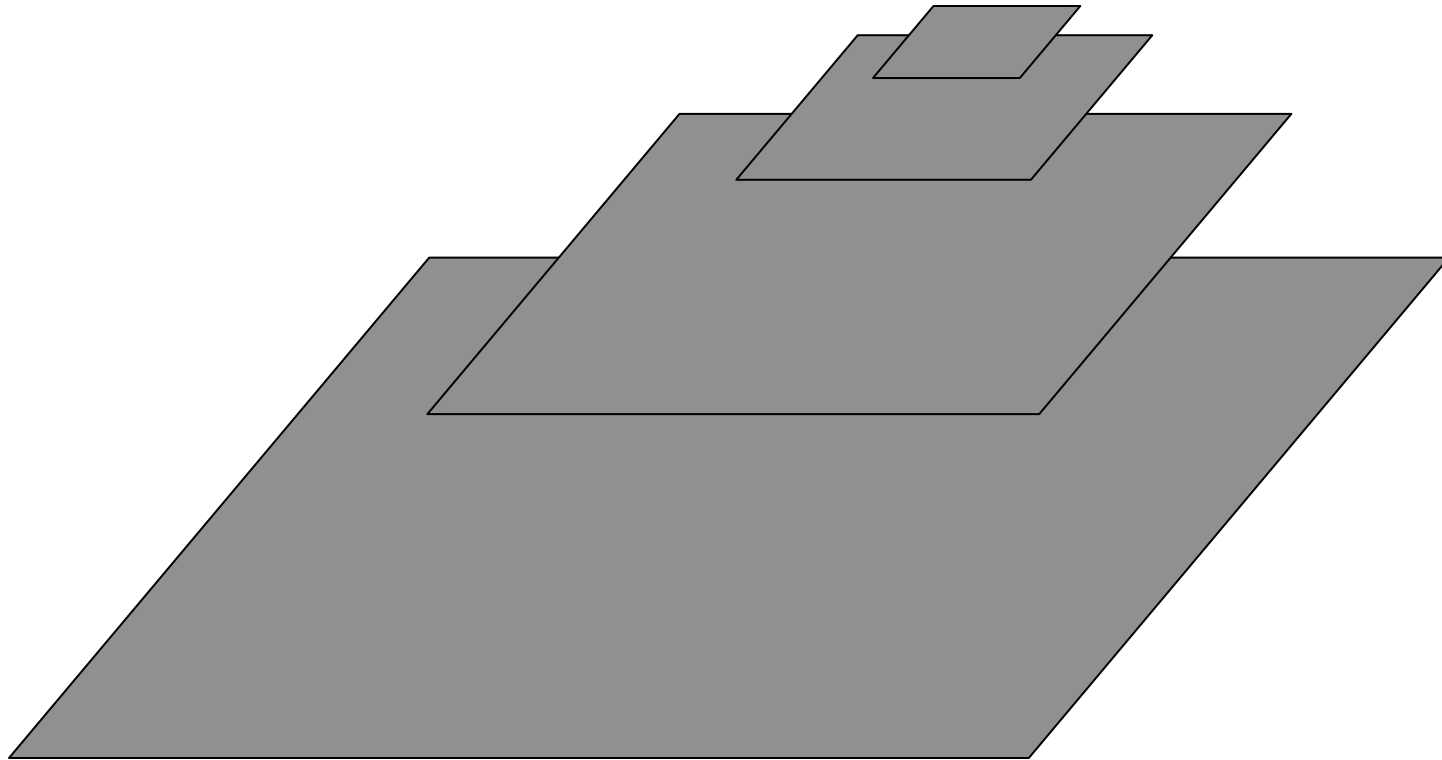
$$C(X, Y) = \min_{\pi} (|X - \pi Y|)$$

- L1 is agnostic about certain correspondences





# Pyramid Matching



# Pyramid Matching

- Form a multiresolution set of histograms (of features)

$$\Psi(\mathbf{X}) = [H_0(\mathbf{X}), \dots, H_{L-1}(\mathbf{X})]$$

- Pyramid match distance/similarity

$$\mathcal{P}_\Delta(\Psi(\mathbf{Y}), \Psi(\mathbf{Z})) = \sum_{i=0}^{L-1} w_i N_i$$

- $N_i$  is number of matches at each level

$$\mathcal{I}(\mathbf{A}, \mathbf{B}) = \sum_{j=1}^r \min(\mathbf{A}^{(j)}, \mathbf{B}^{(j)}) \quad N_i = \mathcal{I}(H_i(\mathbf{Y}), H_i(\mathbf{Z})) - \mathcal{I}(H_{i-1}(\mathbf{Y}), H_{i-1}(\mathbf{Z}))$$

- Cost/distance vs similarity

$$w_i = d2^i \quad w_i = \frac{1}{d2^i}$$

# Pyramid Matching

- Form a multiresolution set of histograms (of feature sets A and B)

$$H(A) = [h_1(A), h_2(A), \dots, h_L(A)]$$

$$H(B) = [h_1(B), h_2(B), \dots, h_L(B)]$$

$$\text{Bin size } \frac{1}{d^{2^l}}$$

- Pyramid match distance/similarity

$$\kappa(A, B) = \sum_{l=1}^L w_l N_l$$

- $N_i$  is number of matches at each level

$$N_l(A, B) = I_l(A, B) - I_{l-1}(A, B) \quad I_l(A, B) = \sum_{\text{bins } i} \min \left( h_l^{(i)}(A), h_l^{(i)}(B) \right)$$

- Cost/distance vs similarity

$$w_i = d^{2^i} \quad w_i = \frac{1}{d^{2^i}}$$

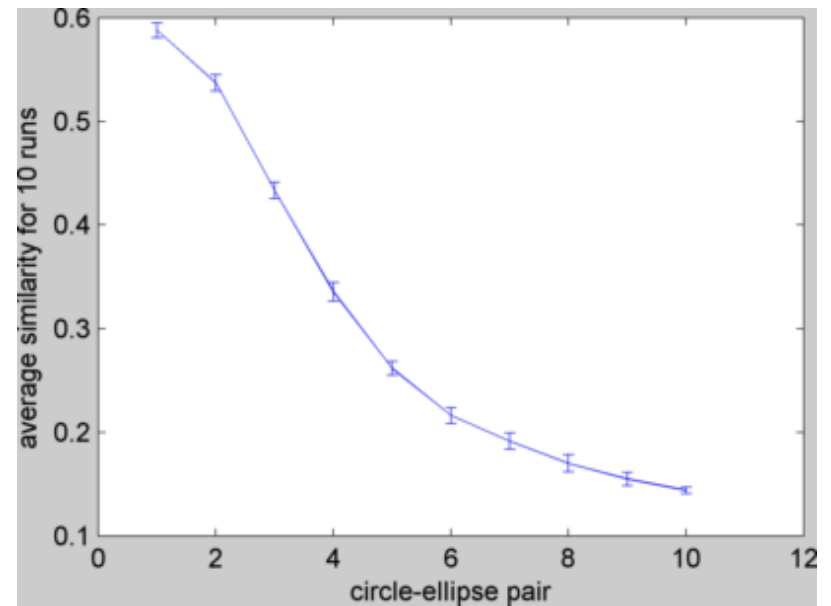
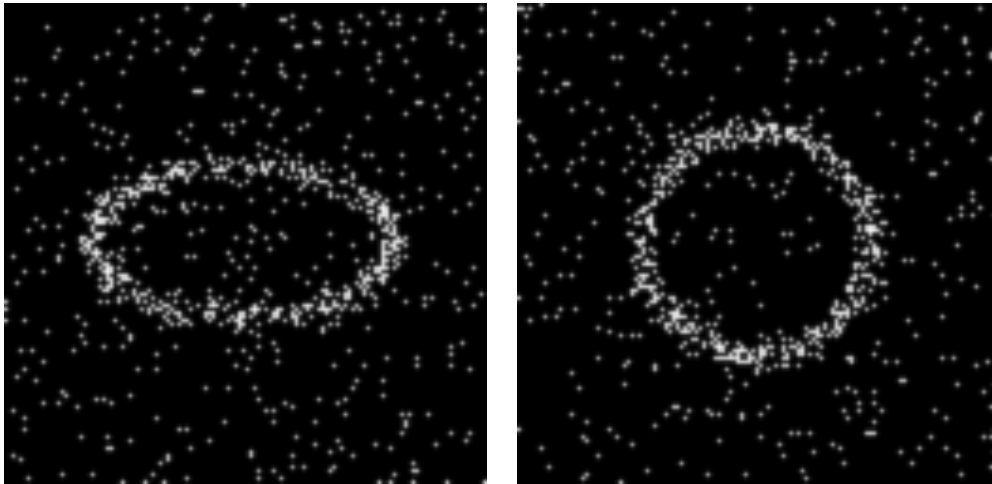
# Pyramid Match Kernels

## Grauman 2006

- The distance case approximates L1 (expected)
- Distance case -> metric
- Similarity case -> Mercer kernel
  - Also robust to outliers/mismatches
- Technicalities
  - How to deal with different numbers of features
  - Should be “normalized”
- Use x-y coordinates and coded features
  - SPM, Lazebnik 2006

# Some Experiments

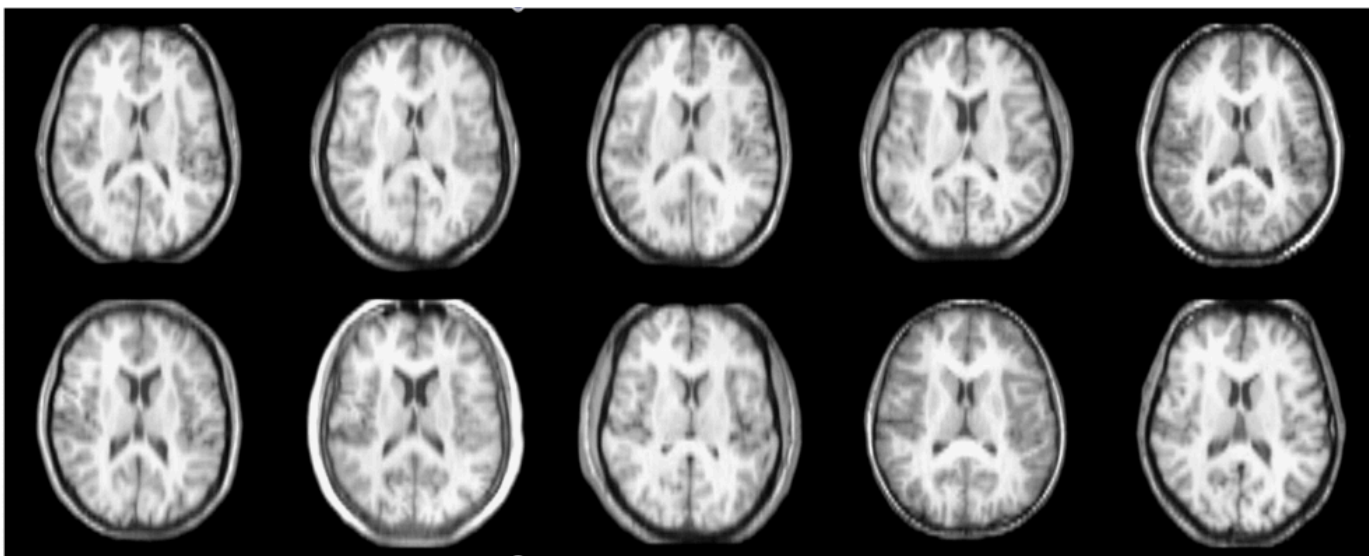
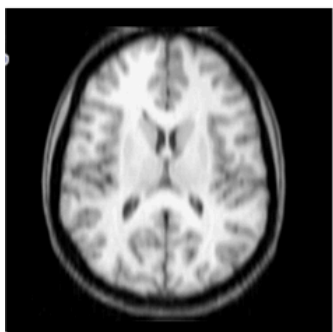
- Random points from circles/ellipses
- Up to 15% random points (mismatches)
- Kernels moderately robust (distinguish shapes)
- Distance less so...



# Image Segmentation

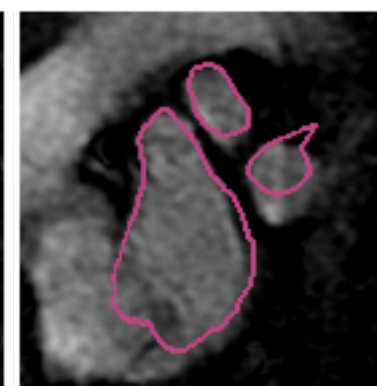
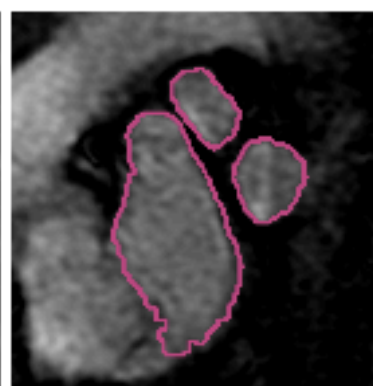
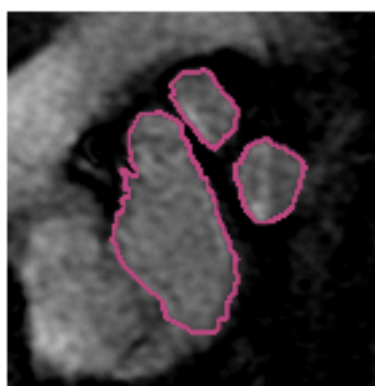
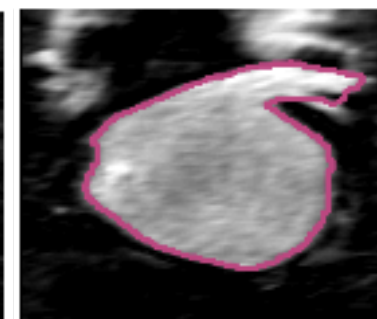
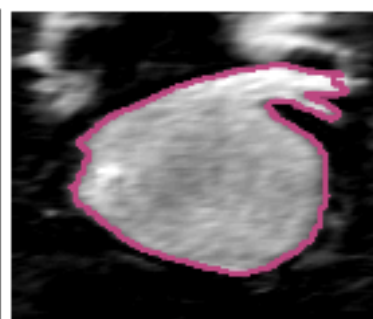
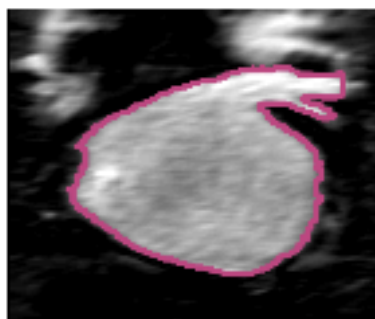
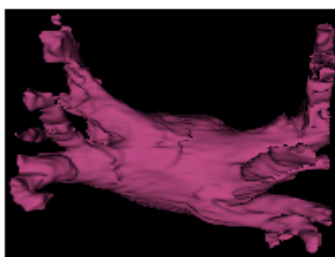
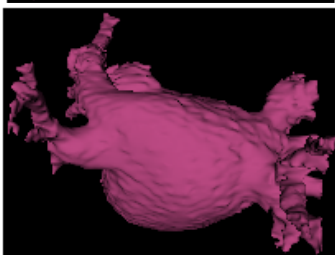
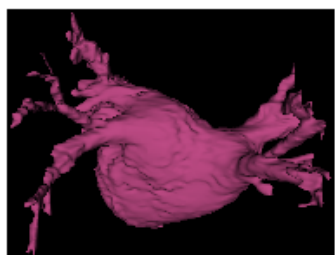
## How to Choose a Template?

- Single individual? Not general
- Average of whole population? Too general
- Choose a set of segmented images that



# Choosing Similar Templates

- Nonparametric estimators on the space of objects [Depa, et al. 2010]



(a) expert

(b) WV

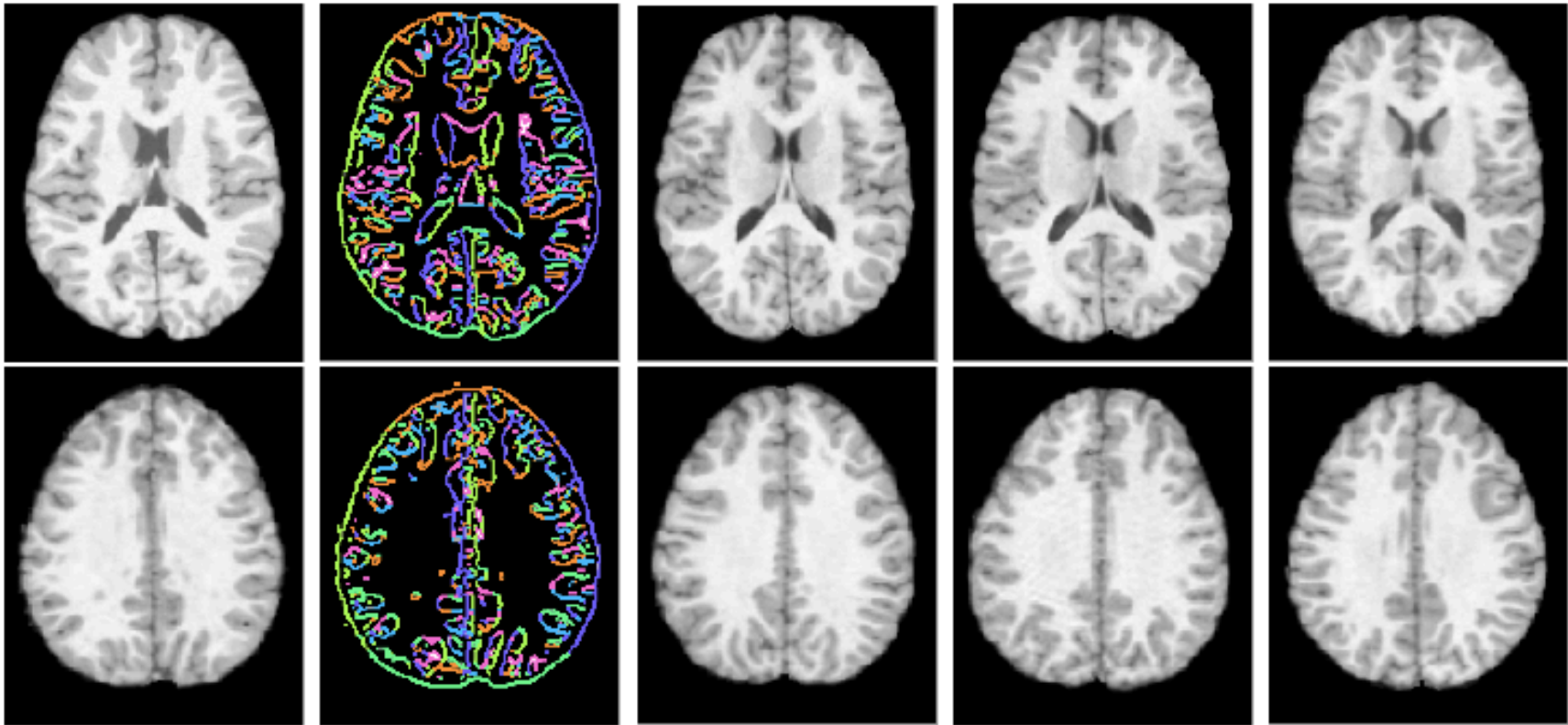
(d) AT

# Challenge

- Given (potentially) thousands or millions of examples, how do we find the most similar images (shape)?
- Zhu, et al. , MICCAI 2011
  - Use spm as an approximate, fast shape “lookup” for very large sets of examples
  - Strategy: (i) Features (ii) Codebooks (iii) SPM for shape similarity



# Nearest Neighbor Lookup (Brains)



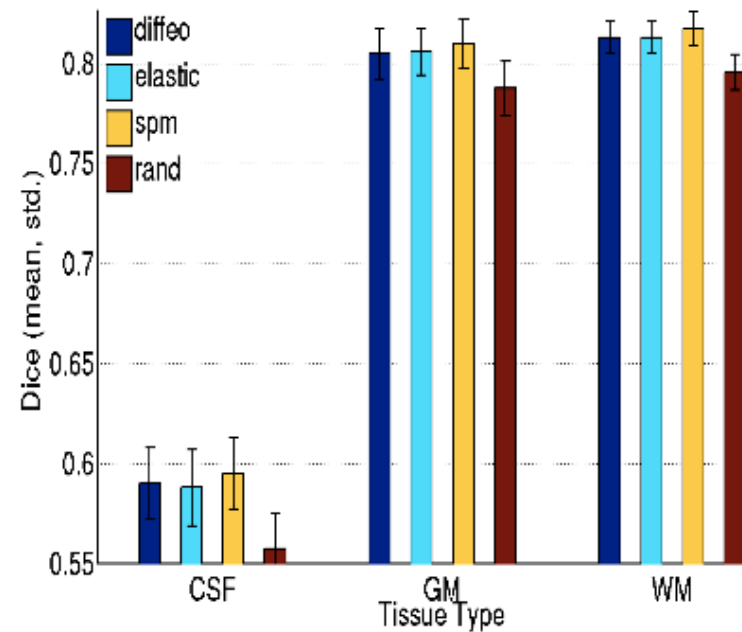
# Nearest Neighbor Lookup Segmentation Performance

<i>k</i> -NN Accuracy							
	Diff <sub>1</sub>	Diff <sub>2</sub>	Elas <sub>1</sub>	Elas <sub>2</sub>	SPM <sub>1</sub>	SPM <sub>6</sub>	SPM <sub>18</sub>
Diff <sub>1</sub>	1	0.39	0.22	0.35	0.25	0.32	0.32
Diff <sub>2</sub>		1	0.51	0.69	0.45	0.53	0.53
Elas <sub>1</sub>			1	0.45	0.36	0.36	0.36
Elas <sub>2</sub>				1	0.42	0.52	0.53
SPM <sub>1</sub>					1	0.56	0.52
SPM <sub>6</sub>						1	0.86
SPM <sub>18</sub>							1

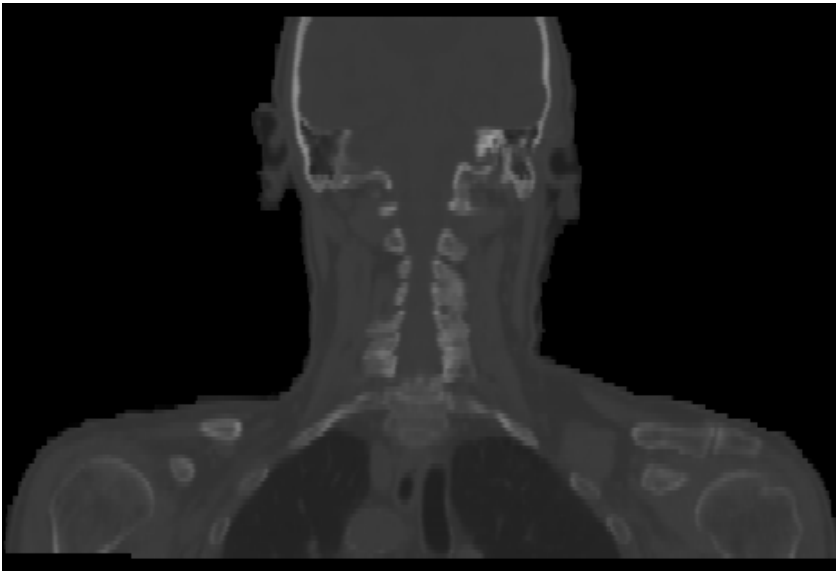
Average $\epsilon$ -Ball Radius Ratio							
	Diff <sub>1</sub>	Diff <sub>2</sub>	Elas <sub>1</sub>	Elas <sub>2</sub>	SPM <sub>1</sub>	SPM <sub>6</sub>	SPM <sub>18</sub>
Diff <sub>1</sub>	1	1.24	1.30	1.25	1.38	1.33	1.32
Diff <sub>2</sub>	1.26	1	1.20	1.16	1.33	1.29	1.26
Elas <sub>1</sub>	1.29	1.19	1	1.23	1.29	1.27	1.27
Elas <sub>2</sub>	1.13	1.07	1.10	1	1.12	1.09	1.09

Tissue classification

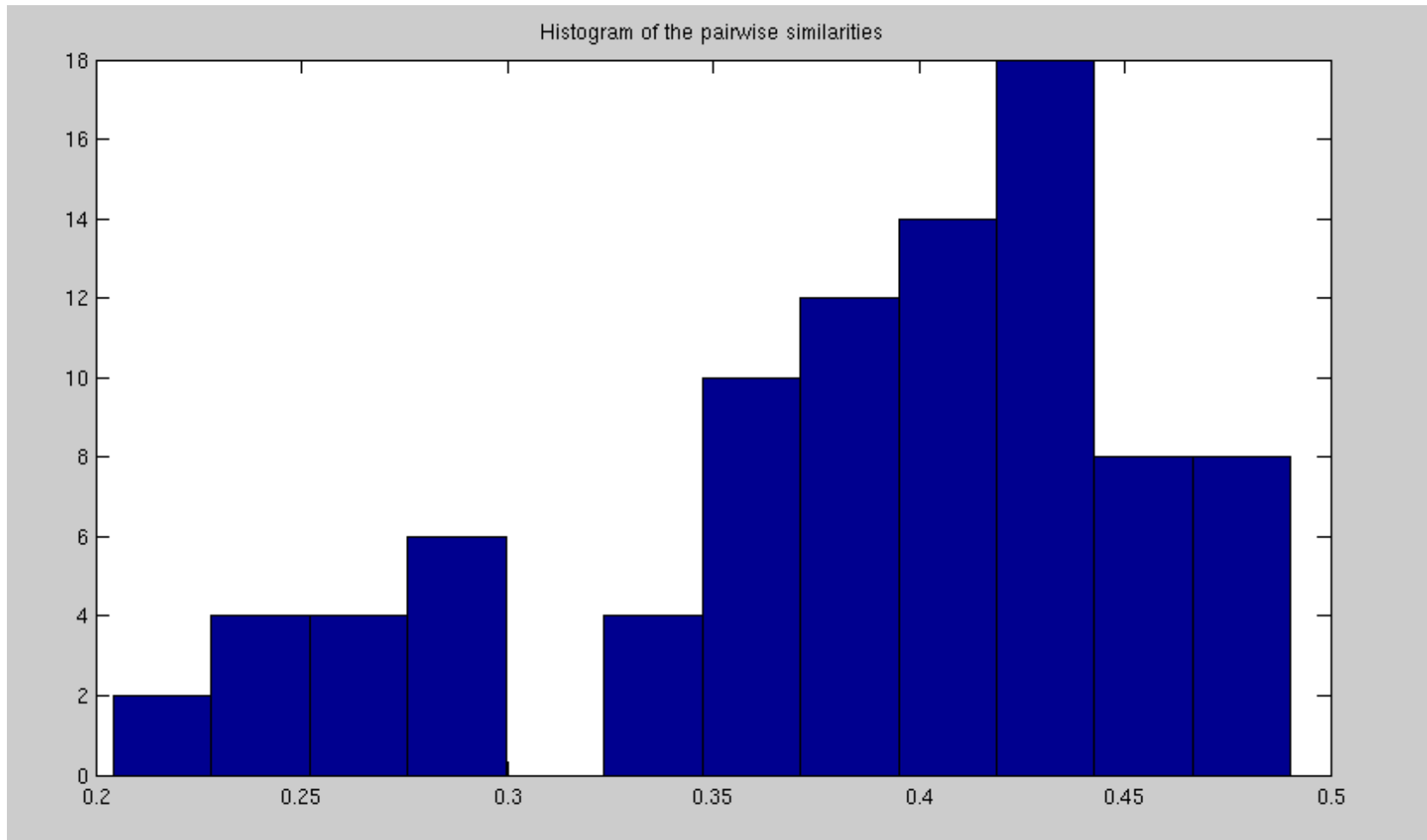


# Another Example

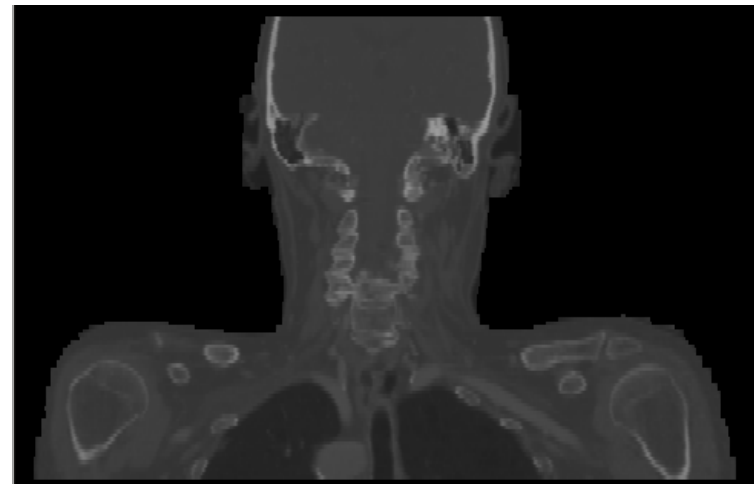
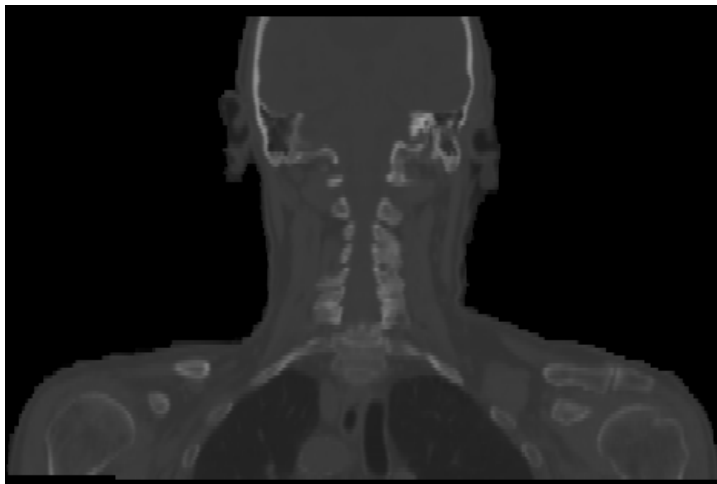
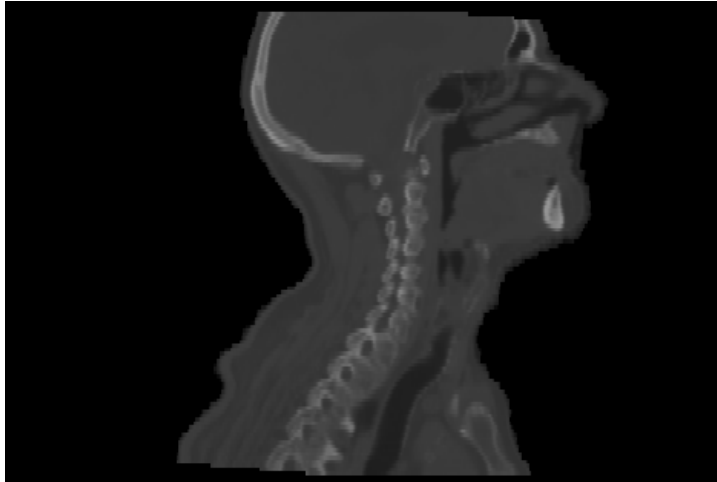
- Head/neck CT for radiation treatment
- Can we reuse old segmentations?
  - E.g. from large database



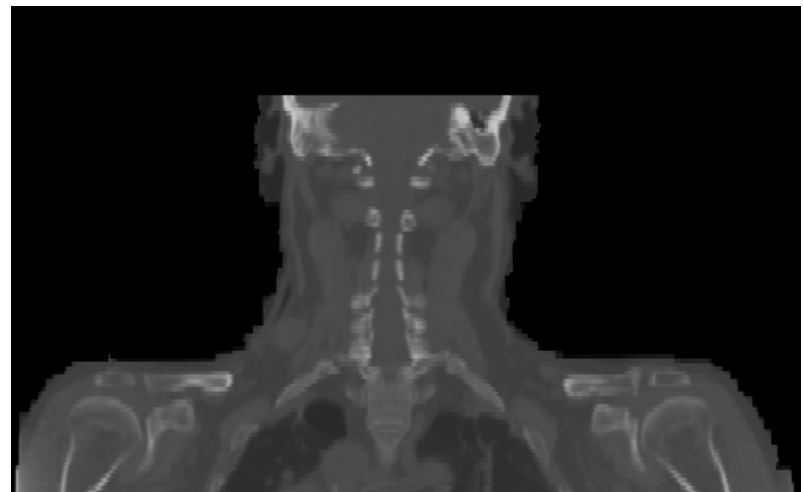
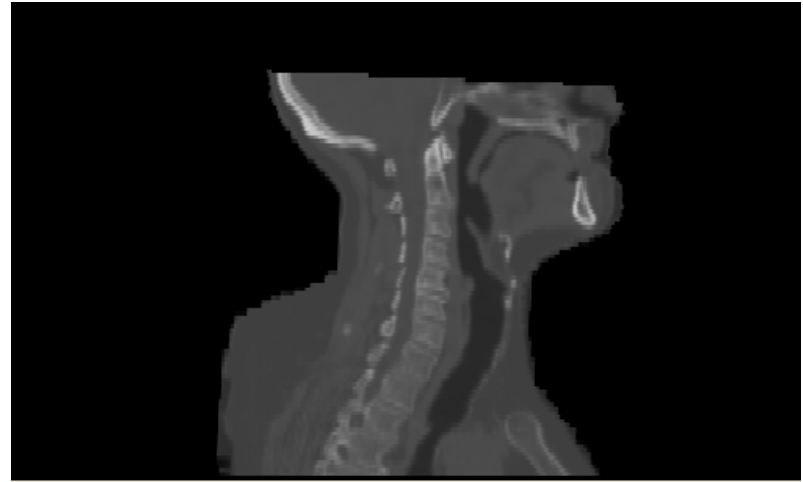
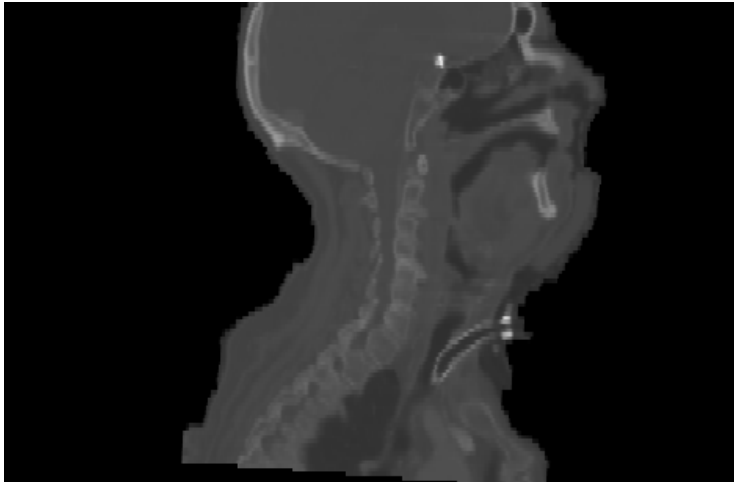
# Experiment - 10 Scans



# Good Match



# Bad Match



# Speed Is Important

- Floating point operations (est. for vol)
  - LDDM -  $10^{13}$
  - Elastic registration -  $10^{11}$
  - SPM -  $10^8$
- Can we do more with this representation?
  - E.g. statistics