



NSF Engineering Research Center for  
Computer Integrated Surgical Systems and  
Technology



LABORATORY FOR  
**Computational  
Sensing + Robotics**  
THE JOHNS HOPKINS UNIVERSITY

## Registration – Part 3

600.455/655 Computer Integrated Surgery



100 YEARS  
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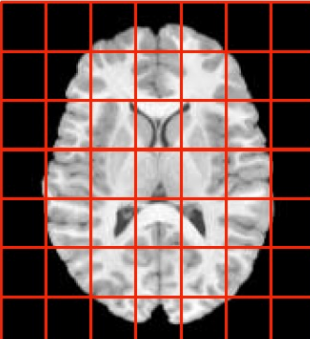
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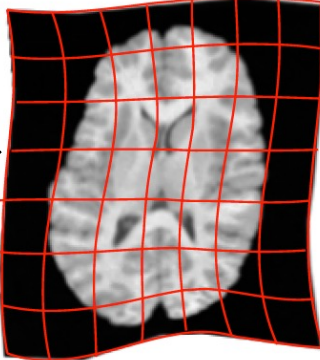
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### Deformable Registration




$\text{Im}(\vec{x})$

→

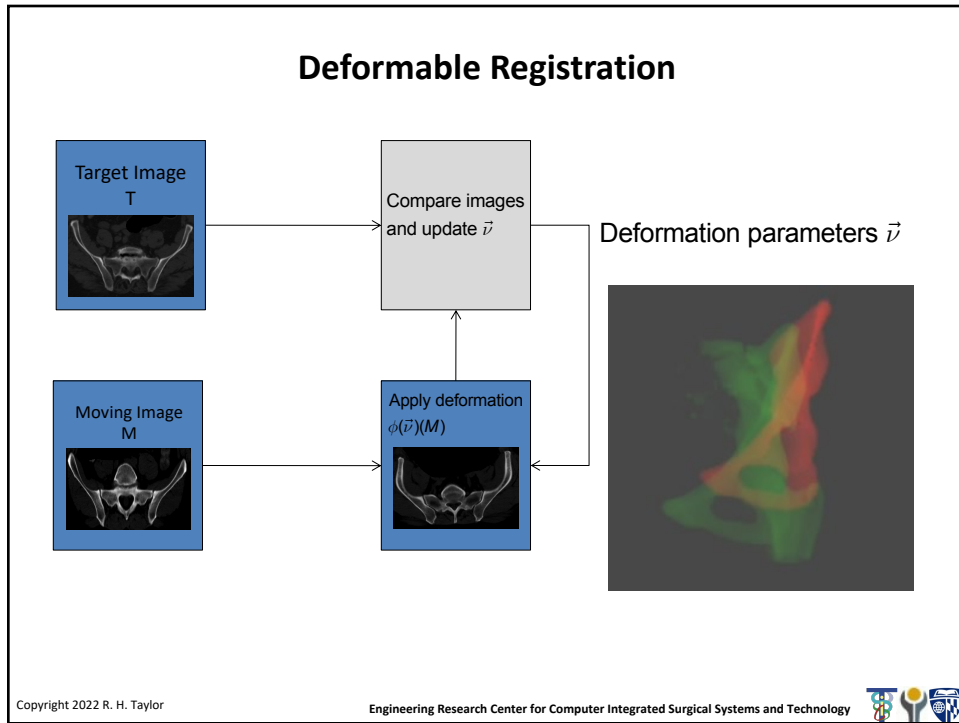


$\text{Im}(\Phi(\vec{\rho}, \vec{x}))$

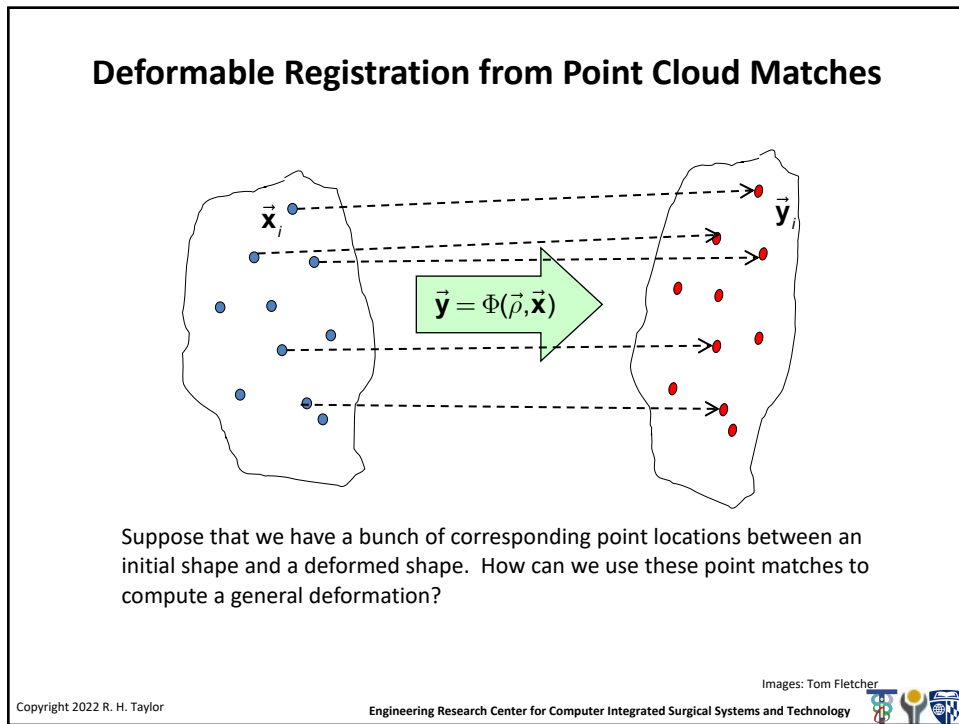
- Many different ways to parameterize the deformation function
  - Typically some version of a spline or radial basis function
- One desirable (though not universal) property: diffeomorphism
  - A function  $\Phi$  is diffeomorphic if  $\Phi$  is bijective and both  $\Phi$  and  $\Phi^{-1}$  are smooth

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## Deformable warping from point cloud matches

- One answer would be the deformable Coherent Point Drift algorithm (Myronenko & Song, *IEEE PAMI*, 2010)
- Another answer might make use of what we learned in programming assignments
  - E.g., fit Bernstein or B-spline polynomials to determine distortion.

$$\vec{u} = \text{TrimToBox}(\vec{x})$$

$$\vec{y} = \sum_{i,j,k} \vec{c}_{i,j,k} B_i(u_x) B_j(u_y) B_k(u_z)$$

or

$$\vec{y} = \sum_{i,j,k} \vec{c}_{i,j,k} N_i(u_x) N_j(u_y) N_k(u_z)$$

- Note: In this case, the coefficients will also parameterize the “shape”

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## Radial Basis Functions

Given a scalar function  $\phi(\bullet)$  and a set of sample points  $\vec{p}_k$  with associated deformations  $\vec{d}_k$ , one can represent the deformation  $\Phi$  at a point  $\vec{x}$  by

$$\Phi(\vec{x}) = \sum_k \vec{d}_k \phi_k(\|\vec{x} - \vec{p}_k\|)$$

- Many possible functions to use for  $\phi$ 
  - Common choices include Gaussians and “thin plate splines”, which have non-compact support (i.e.,  $\Phi(y) > 0$  for arbitrarily large  $y$ )
  - Others have compact support (i.e.,  $\Phi(y) = 0$  for  $|y| >$  some value)\*

\* See: M. Fornefett, K. Rohr, and H. S. Stiehl, "Radial basis functions with compact support for elastic registration of medical images", *Image and Vision Computing*, vol. 19- 1, 2001, pp. 87-96, 2001.  
<http://www.sciencedirect.com/science/article/pii/S0262885600000573>  
[http://dx.doi.org/10.1016/S0262-8856\(00\)00057-3](http://dx.doi.org/10.1016/S0262-8856(00)00057-3)

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## Thin Plate Splines

- Minimum energy spline deformations

$$TPS(\vec{v}; \vec{a}, \mathbf{B}, \mathbf{C}, \mathbf{P}) = \vec{a} + \mathbf{B} \cdot \vec{v} + \sum_i \vec{c}_i U(\|\vec{v} - \vec{p}_i\|)$$

where  $U(r) = r^2 \log(r)$  for 2D images

- Global support
- Popularized by Fred Bookstein for analysis of anatomic variation
  - F. L. Bookstein, *Morphometric tools for landmark data*, Geometry and biology: Cambridge University Press, 1991.

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## Thin Plate Splines Digression

- Some citations (from G. Donato and S. Belongie, “Approximation Methods for Thin Plate Spline Mappings and Principal Warps”, 2002; [http://www.cs.ucsd.edu/Dienst/UI/2.0/Describe/ncstrl.ucsd\\_cse/CS2003-0764](http://www.cs.ucsd.edu/Dienst/UI/2.0/Describe/ncstrl.ucsd_cse/CS2003-0764) )

- [1] C. T. H. Baker. *The numerical treatment of integral equations*. Oxford: Clarendon Press, 1977.
- [2] S. Belongie, J. Malik, and J. Puzicha. Matching shapes. In *Proc. 5th Int'l. Conf. Computer Vision*, volume 1, pages 454–461, July 2001.
- [3] F. L. Bookstein. Principal warps: thin-plate splines and decomposition of deformations. *IEEE Trans. Pattern Analysis and Machine Intelligence*, 11(6):567–585, June 1989.
- [4] H. Chui and A. Rangarajan. A new algorithm for non-rigid point matching. In *Proc. IEEE Conf. Comput. Vision and Pattern Recognition*, pages 44–51, June 2000.
- [5] M.H. Davis, A. Khotanzad, D. Flammig, and S. Harms. A physics-based coordinate transformation for 3-d image matching. *IEEE Trans. Medical Imaging*, 16(3):317–328, June 1997.
- [6] F. Girosi, M. Jones, and T. Poggio. Regularization theory and neural networks architectures. *Neural Computation*, 7(2):219–269, 1995.
- [7] M. J. D. Powell. A thin plate spline method for mapping curves into curves in two dimensions. In *Computational Techniques and Applications (CTACS)*, Melbourne, Australia, 1995.
- [8] A.J. Smola and B. Schölkopf. Sparse greedy matrix approximation for machine learning. In *ICML*, 2000.
- [9] G. Wahba. *Spline Models for Observational Data*. SIAM, 1990.
- [10] Y. Weiss. Smoothness in layers: Motion segmentation using nonparametric mixture estimation. In *Proc. IEEE Conf. Comput. Vision and Pattern Recognition*, pages 520–526, 1997.
- [11] C. Williams and M. Seeger. Using the Nyström method to speed up kernel machines. In T. K. Leen, T. G. Dietterich, and V. Tresp, editors. *Advances in Neural Information Processing Systems 13: Proceedings of the 2000 Conference*, pages 682–688, 2001.

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## M-dimensional Thin Plate Spline Summary

Given

$$TPS(\vec{v}; \vec{a}, \mathbf{B}, \mathbf{C}, \mathbf{P}) = \vec{a} + \mathbf{B} \bullet \vec{v} + \sum_i \vec{c}_i U(\|\vec{v} - \vec{p}_i\|)$$

where

$$U(r) = r^2 \log(r) \quad \text{for 2D}$$

$$= r^2 \log(r^2) \quad \text{for 3D}$$

$$\vec{v} = [v_1, \dots, v_M]^T$$

$$\vec{p}_i = [p_{1,i}, \dots, p_{M,i}]^T$$

$$\mathbf{P} = [\vec{p}_1, \dots, \vec{p}_N]^T$$

$$\mathbf{C} = [\vec{c}_1, \dots, \vec{c}_N]$$

$$\mathbf{B} = [\vec{b}_1, \dots, \vec{b}_M]$$

Note: Some sources give

$$U(r) = \begin{cases} r^{4-m} \ln(r) & \text{for } m=2 \text{ or } 4 \\ r^{4-m} & \text{otherwise} \end{cases}$$

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## M-dimensional Thin Plate Spline Fitting

Given

$$\mathbf{V} = [\vec{v}_1, \dots, \vec{v}_N] \quad \mathbf{F} = [\vec{f}_1, \dots, \vec{f}_N]$$

find  $\vec{a}$ ,  $\mathbf{B}$ ,  $\mathbf{C}$  such that

$$\vec{f}_i = TPS(\vec{v}_i; \vec{a}, \mathbf{B}, \mathbf{C}, \mathbf{V})$$

To do this, solve the linear system

$$\begin{bmatrix} \mathbf{K}_{[N \times N]} & \vec{1}_{[N \times 1]} & \mathbf{V} \\ \vec{1}_{[1 \times N]} & 0 & 0 \\ \mathbf{V}^T & 0 & \mathbf{0}_{[M \times M]} \end{bmatrix} \begin{bmatrix} \mathbf{C}^T \\ \vec{a}^T \\ \mathbf{B}^T \end{bmatrix} = \begin{bmatrix} \mathbf{F}^T \\ 0 \\ \mathbf{0}_{[M \times 1]} \end{bmatrix}$$

where

$$\mathbf{K}_{i,j} = \mathbf{K}_{j,i} = U(\|\vec{v}_i - \vec{v}_j\|) \quad \text{with } U(r) = r^2 \log r \text{ or } U(r) = r^2 \log r^2$$

$$\mathbf{K}_{i,j} = (\vec{v}_i - \vec{v}_j) \bullet (\vec{v}_i - \vec{v}_j) \log \left( \sqrt{(\vec{v}_i - \vec{v}_j) \bullet (\vec{v}_i - \vec{v}_j)} \right)$$

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### TPS 2D case

Given a set of points  $\vec{p}_i = [x_i, y_i]$  and corresponding points  $\vec{p}_i^* = [x_i^*, y_i^*]$ , we want to find TPS parameters such that  $\vec{p}_i^* = TPS(\vec{p}_i; \vec{a}, \mathbf{B}, \mathbf{C}, \mathbf{P})$

To do this, we solve the least squares problem

$$\begin{bmatrix} 0 & \cdots & U_{1,k} & \cdots & U_{1,N} & 1 & x_1 & y_1 \\ \vdots & \ddots & & U_{ij} & & \vdots & \vdots & \vdots \\ U_{k,1} & \cdots & 0 & \cdots & U_{k,N} & 1 & x_k & y_k \\ \vdots & U_{ij} & & \ddots & \vdots & \vdots & \vdots & \vdots \\ U_{N,1} & \cdots & U_{N,k} & \cdots & 0 & 1 & x_N & y_N \\ 1 & \cdots & 1 & \cdots & 1 & 0 & 0 & 0 \\ x_1 & \cdots & x_k & \cdots & x_N & 0 & 0 & 0 \\ y_1 & \cdots & y_k & \cdots & y_N & 0 & 0 & 0 \end{bmatrix} \bullet \begin{bmatrix} \vec{c}_1 \\ \vdots \\ \vdots \\ \vec{c}_N \\ \vec{a} \\ \vec{b}_x \\ \vec{b}_y \end{bmatrix} = \begin{bmatrix} \vec{p}_1^* \\ \vdots \\ \vec{p}_k^* \\ \vdots \\ \vec{p}_N^* \\ \vec{0} \\ \vec{0} \\ \vec{0} \end{bmatrix}$$

where  $U_{i,j} = U_{j,i} = U(\|\vec{p}_i - \vec{p}_j\|)$

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### M-dimensional Thin Plate Spline Fitting

Define

$$\mathbf{L}_{[M+N+1 \times M+N+1]} = \begin{bmatrix} \mathbf{K}_{[N \times N]} & \vec{\mathbf{1}}_{[N \times 1]} & \mathbf{V} \\ \vec{\mathbf{1}}_{[1 \times N]} & \mathbf{0} & \mathbf{0} \\ \mathbf{V}^T & \mathbf{0} & \mathbf{0}_{[M \times M]} \end{bmatrix}$$

If there are many points, this matrix may be expensive to invert or even pseudo-invert. There are various methods to deal with this problem. These include

- Use a random sample of the  $\vec{v}_i$  to approximate the solution
- Use a random sample of the basis functions & all data to solve problem in least squares sense
- Use matrix approximation methods

See

[http://www.cs.ucsd.edu/Dienst/UI/2.0/Describe/ncstrl.ucsd\\_cse/CS2003-0764](http://www.cs.ucsd.edu/Dienst/UI/2.0/Describe/ncstrl.ucsd_cse/CS2003-0764)

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## Other Radial Basis Functions

Note that the function  $U(r)$  in the previous discussion is an example of a more general class of "radial basis functions". These functions can be used in deformable registration in much the same way as the TPS function used above. Other commonly used radial basis functions include

$$U(r) = (r^2 + c^2)^\mu \text{ for } \mu \in \mathbb{R}_+$$

$$U(r) = (r^2 + c^2)^{-\mu} \text{ for } \mu \in \mathbb{R}_+$$

$$U(r) = e^{-r^2/2\sigma^2}$$

The last one is probably the most popular for global support. There are also radial basis functions with "compact" support. For example\*

$$\Psi(r, \sigma) = \begin{cases} \left(1 - \frac{r}{\sigma}\right)^{k+1+|d|/2} & \text{if } 0 \leq r \leq \sigma \\ 0 & \text{otherwise} \end{cases}$$

\* See: M. Fornefett, K. Rohr, and H. S. Stiehl, "Radial basis functions with compact support for elastic registration of medical images", *Image and Vision Computing*, vol. 19-1, A12, pp. 87-96, 2001.  
<http://www.sciencedirect.com/science/article/pii/S0262885600000573>  
[http://dx.doi.org/10.1016/S0262-8856\(00\)00057-3](http://dx.doi.org/10.1016/S0262-8856(00)00057-3)

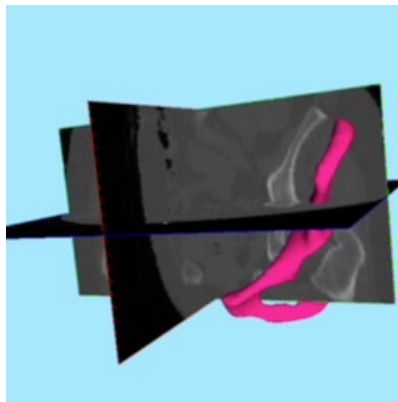
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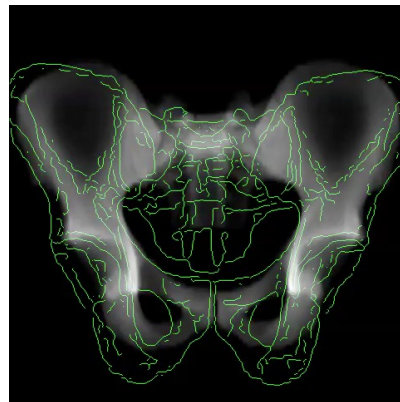


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## Deformable Registration to Statistical "Atlases"



Deformable 3D/3D  
Jianhua Yao



Deformable 2D/3D  
Ofri Sadovskiy

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## Deformable Atlas-based Registration

- Much of the material that follows is derived from the Ph.D. thesis work of J. Yao, Ofri Sadowsky, and Gouthami Chintalapani:
  - J. Yao, “Statistical bone density atlases and deformable medical image registrations”, Ph. D. Thesis, Computer Science, The Johns Hopkins University, Baltimore, 2001.
  - O. Sadowsky, "Image Registration and Hybrid Volume Reconstruction of Bone Anatomy Using a Statistical Shape Atlas," Ph.D. Thesis, Computer Science, The Johns Hopkins University, Baltimore, 2008
  - G. Chintalapani, Statistical Atlases of Bone Anatomy and Their Applications, Ph.D. thesis in Computer Science, The Johns Hopkins University, Baltimore, Maryland, 2010.
- A number of other authors, including
  - Cootes et al. 1999 – “Active Appearance Models”
  - Feldmar and Ayache 1994
  - Ferrant et al. 1999
  - Fleute and Lavallee 1999
  - Lowe 1991
  - Maurer et al. 1996
  - Shen and Davatzikos 2000

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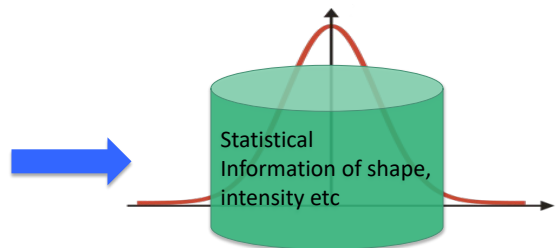
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## What is a “Statistical Atlas” ?

- An atlas that incorporates statistics of anatomical shape and intensity variations of a given population



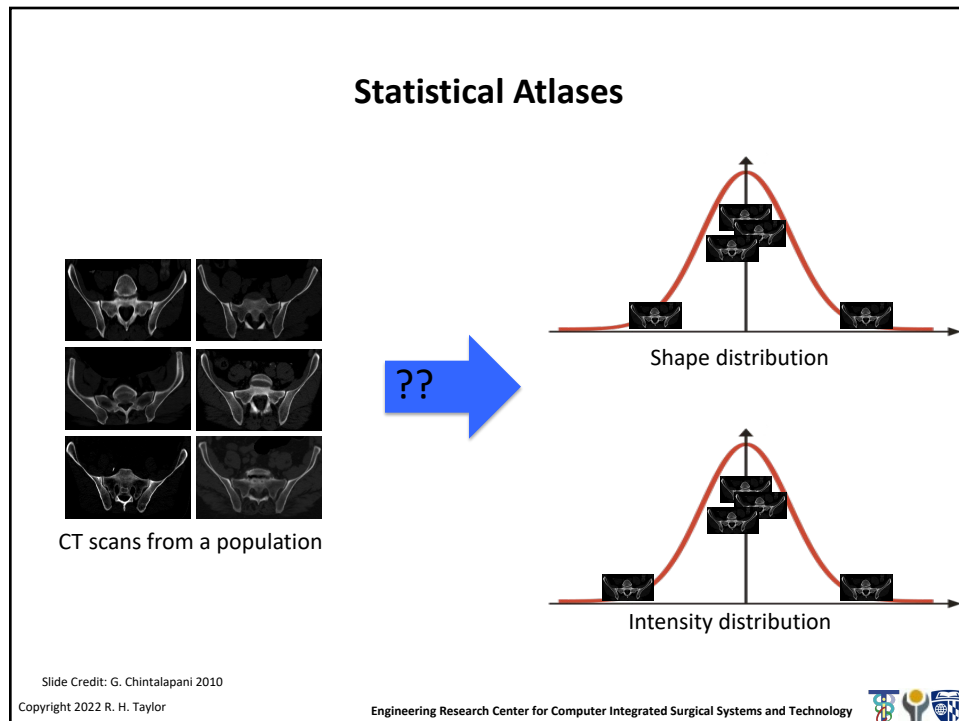
Credit: G. Chintalapani 2010

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## Statistical models

- The next few slides will review the use of the Singular Value Decomposition (SVD) in constructing statistical shape models.
- There is a close relationship between this material and the “principal components analysis” (PCA) methods you may have encountered in a statistics class.

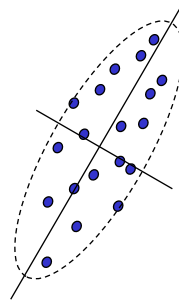
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## Principal Components Analysis (PCA)

Suppose that you have a set of  $N$  vectors  $\vec{a}_i$  in an  $M$  dimensional space?  
Is there a natural "coordinate system" for these vectors?



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## Principal Components Analysis (PCA)

We proceed as follows

$$\vec{a}^{(avg)} = \frac{\sum_i \vec{a}_i}{N}; \quad \vec{b}_i = \vec{a}_i - \vec{a}^{(avg)}; \quad \mathbf{B} = [\vec{b}_1, \dots, \vec{b}_N];$$

Then form the singular value decomposition

$$\mathbf{B} = \mathbf{U} \Sigma \mathbf{V}^T = \mathbf{U} \begin{bmatrix} \Sigma^{(N)} \\ \mathbf{0} \end{bmatrix} \mathbf{V}^T \quad \text{where } \Sigma^{(N)} = \text{diag}(\sigma_1, \dots, \sigma_N)$$

Then we note that  $\mathbf{B}\mathbf{B}^T = \mathbf{U}\Sigma^2\mathbf{U}^T$ . Of course  $\mathbf{U}$  is huge, but we have the following useful fact. We note that

$$\mathbf{B} = [\vec{u}_1, \dots, \vec{u}_N, \vec{u}_{N+1}, \dots, \vec{u}_M] \begin{bmatrix} \sigma_1 & & & \\ & \ddots & & \\ & & \sigma_N & \\ & & & 0 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \end{bmatrix} \mathbf{V}^T = [\vec{u}_1, \dots, \vec{u}_N] \Sigma^{(N)} \mathbf{V}^T = \mathbf{U}^{(N)} \Sigma^{(N)} \mathbf{V}^T$$

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## Principal Components Analysis (PCA)

This means that any column  $\vec{b}_k$  of  $\mathbf{B}$  may be expressed as a linear combination of the first  $N$  columns of  $\mathbf{U}$

$$\mathbf{B} = [\vec{u}_1, \dots, \vec{u}_N] \Sigma^{(N)} \mathbf{V}^T = \mathbf{U}^{(N)} \Sigma^{(N)} \mathbf{V}^T$$

$$\vec{b}_k = \lambda_1^{(k)} \vec{u}_1 + \dots + \lambda_N^{(k)} \vec{u}_N = \mathbf{U}^{(N)} \Lambda^{(k)}$$

where

$$\Lambda^{(k)} = \text{transpose}(\mathbf{U}^{(N)}) \vec{b}_k$$

So

$$\vec{a}_k = \vec{a}^{(avg)} + \vec{b}_k = \vec{a}^{(avg)} + \lambda_1^{(k)} \vec{u}_1 + \dots + \lambda_N^{(k)} \vec{u}_N$$

But often the last few values of the  $\lambda_k$  are small. If we ignore all but the first  $D$  values, we have

$$\vec{a}_k \approx \vec{a}^{(avg)} + \lambda_1^{(k)} \vec{u}_1 + \dots + \lambda_D^{(k)} \vec{u}_D$$

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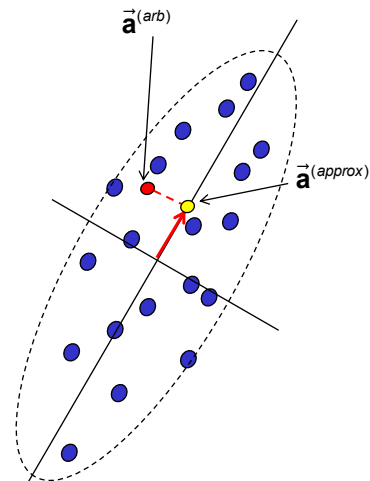
## Principal Components Analysis (PCA)

Suppose now that we have an arbitrary  $\vec{a}^{(arb)}$ . We can approximate  $\vec{a}^{(arb)}$  as follows:

$$\vec{b}^{(arb)} = \vec{a}^{(arb)} - \vec{a}^{(avg)}$$

$$\Lambda^{(arb)} = \text{transpose}(\mathbf{U}^{(D)}) \vec{b}^{(arb)}$$

$$\vec{a}^{(arb)} \approx \vec{a}^{(avg)} + \lambda_1^{(arb)} \vec{u}_1 + \dots + \lambda_D^{(arb)} \vec{u}_D$$



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## Statistical Atlases & PCA

Given a set of  $N$  models  $\vec{\mathbf{X}}^{(j)} = [\vec{\mathbf{x}}_k^{(j)}]^T = [\dots x_k^{(j)}, y_k^{(j)}, z_k^{(j)}, \dots]$ , compute

$$\vec{\mathbf{X}}^{(avg)} = \begin{bmatrix} \vdots \\ \vec{\mathbf{x}}_k^{(avg)} \\ \vdots \end{bmatrix} \text{ where } \vec{\mathbf{x}}_k^{(avg)} = \frac{1}{N} \sum_j \vec{\mathbf{x}}_k^{(j)} \text{ and the differences}$$

$$\vec{\mathbf{D}}^{(j)} = \vec{\mathbf{X}}^{(j)} - \vec{\mathbf{X}}^{(avg)} = \begin{bmatrix} \vdots \\ \vec{\mathbf{d}}_k^{(j)} \\ \vdots \end{bmatrix} \text{ where } \vec{\mathbf{d}}_k^{(j)} = \vec{\mathbf{x}}_k^{(j)} - \vec{\mathbf{x}}_k^{(avg)}. \text{ Create the matrix}$$

$$\mathbf{D} = \begin{bmatrix} \dots & \vec{\mathbf{D}}^{(j)} & \dots \end{bmatrix}_{[3Nvertices \times N]} = \begin{bmatrix} \vec{\mathbf{d}}_1^{(1)} & \dots & \vec{\mathbf{d}}_k^{(1)} & \dots & \vec{\mathbf{d}}_1^{(1)} \\ \vdots & & \vdots & & \vdots \\ \vec{\mathbf{d}}_k^{(1)} & \dots & \vec{\mathbf{d}}_k^{(j)} & \dots & \vec{\mathbf{d}}_k^{(N)} \\ \vdots & & \vdots & & \vdots \\ \vec{\mathbf{d}}_{Nvertices}^{(1)} & \dots & \vec{\mathbf{d}}_{Nvertices}^{(j)} & \dots & \vec{\mathbf{d}}_{Nvertices}^{(N)} \end{bmatrix}$$

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## Statistical Atlases & PCA

Compute the singular value decomposition of  $\mathbf{D}$

$$\mathbf{D} = \mathbf{U}\Sigma\mathbf{V}^T \quad \text{where } \Sigma = \begin{bmatrix} \text{diag}(\vec{\sigma}) \\ \mathbf{0} \end{bmatrix}$$

$$\mathbf{D} = \mathbf{U} \begin{bmatrix} \text{diag}(\vec{\sigma})\mathbf{V}^T \\ \mathbf{0} \end{bmatrix}$$

Note that

$$\frac{1}{N-1} \mathbf{D}^T \mathbf{D} = \frac{1}{N-1} \mathbf{V} \Sigma \mathbf{U}^T \mathbf{U} \Sigma \mathbf{V}^T = \frac{1}{N-1} \mathbf{V} \Sigma^2 \mathbf{V}^T$$

$$\frac{1}{N-1} \mathbf{D} \mathbf{D}^T = \frac{1}{N-1} \mathbf{U} \Sigma \mathbf{V}^T \mathbf{V} \Sigma \mathbf{U}^T = \frac{1}{N-1} \mathbf{U} \Sigma^2 \mathbf{U}^T$$

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## Statistical Atlases & PCA

Any individual model  $\mathbf{D}^{(j)}$  can be written as a linear combination of the columns of  $\mathbf{U}$ . Treating  $\vec{\mathbf{D}}^{(j)}$  as a column vector, we can write this as

$$\vec{\mathbf{D}}^{(j)} = \mathbf{U} \bullet \begin{bmatrix} \lambda_1^{(j)} \\ \vdots \\ \lambda_N^{(j)} \\ \vec{\mathbf{0}} \end{bmatrix} \quad \text{where} \quad \begin{bmatrix} \lambda_1^{(j)} \\ \vdots \\ \lambda_N^{(j)} \\ \vec{\mathbf{0}} \end{bmatrix} \text{ is the } j^{\text{th}} \text{ column of } \begin{bmatrix} \text{diag}(\vec{\sigma})\mathbf{V}^T \\ \mathbf{0} \end{bmatrix}$$

If we define

$$\mathbf{M} = \begin{bmatrix} \mathbf{U}^{(1)} & \dots & \mathbf{U}^{(N)} \end{bmatrix} \quad (\text{i.e., the first } N \text{ columns of } \mathbf{U})$$

we get the expression

$$\vec{\mathbf{D}}^{(j)} = \mathbf{M}\vec{\lambda} \quad \text{where } \vec{\lambda} \text{ is the } j^{\text{th}} \text{ column of } (\text{diag}(\vec{\sigma})\mathbf{V}^T).$$

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## Statistical Atlases & PCA

Note that while  $\mathbf{U}$  is  $3N_{\text{vertices}} \times 3N_{\text{vertices}}$  (i.e., huge),  $\mathbf{M}$  has only the first  $N$  columns, since there are at most  $N$  non-zero singular values

In fact, we usually also truncate even more, only saving columns corresponding to relatively large singular values  $\sigma_i$ . Since the standard algorithms for SVD produce positive singular values  $\sigma_i$  sorted in descending order, this is easy to do.

Note also, that since the columns of  $\mathbf{M}$  are also columns of  $\mathbf{U}$ , they are orthogonal. Hence  $\mathbf{M}^T\mathbf{M} = \mathbf{I}_{N \times N}$ . But  $\mathbf{M}\mathbf{M}^T = \mathbf{C}$  will be an  $3N_{\text{vertices}} \times 3N_{\text{vertices}}$  matrix that will not in general be diagonal.

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## Statistical Atlases & PCA

As a practical matter, it is not a good idea to ask your SVD program to produce the full matrix  $\mathbf{U}$  for an  $3N_{\text{vertices}} \times N$  matrix  $\mathbf{D}$ . Many SVD packages give you the option to compute only the singular values  $\vec{\sigma}$  and the right hand side matrix  $\mathbf{V}$  or its transpose. Then,  $\mathbf{M}$  can be computed from

$$\mathbf{M} \text{diag}(\vec{\sigma}) \mathbf{V}^T = \mathbf{D}$$

$$\mathbf{M} \text{diag}(\vec{\sigma}) = \mathbf{D} \mathbf{V}$$

$$\mathbf{M} = \mathbf{D} \mathbf{V} \text{diag}(\vec{\sigma})^{-1}$$

$$= \mathbf{D} \mathbf{V} \begin{bmatrix} 1/\sigma_1 & 0 & \dots & \dots & 0 \\ 0 & \ddots & & & \vdots \\ \vdots & & 1/\sigma_k & & \vdots \\ \vdots & & & \ddots & 0 \\ 0 & \dots & \dots & 0 & 1/\sigma_N \end{bmatrix}$$

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## Statistical Atlases & PCA

Similarly, given a vector  $\vec{\mathbf{D}}^{(inst)}$  we can find a corresponding vector  $\vec{\lambda}^{(inst)}$  from the following

$$\vec{\mathbf{D}}^{(inst)} = \mathbf{M} \vec{\lambda}^{(inst)}$$

$$\begin{aligned} \mathbf{M}^T \vec{\mathbf{D}}^{(inst)} &= \mathbf{M}^T \mathbf{M} \vec{\lambda}^{(inst)} \\ &= \vec{\lambda}^{(inst)} \end{aligned}$$

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## Statistical Atlases & PCA

Suppose that we select  $\vec{\lambda} = [\lambda_1, \dots, \lambda_N]^T$  as a random variable with some distribution having expected value  $E(\vec{\lambda}) = \vec{\mathbf{0}}$  and covariance

$$\text{cov}(\vec{\lambda}) = E(\vec{\lambda} \bullet \vec{\lambda}^T) = \begin{bmatrix} E(\lambda_1^2) & \dots & E(\lambda_1 \lambda_N) \\ \vdots & \ddots & \vdots \\ E(\lambda_N \lambda_1) & \dots & E(\lambda_N^2) \end{bmatrix} = \Sigma^2$$

and compute a corresponding random model  $\vec{\mathbf{X}}(\vec{\lambda})$

$$\vec{\mathbf{X}}(\vec{\lambda}) = \vec{\mathbf{X}}^{(avg)} + \mathbf{M} \bullet \vec{\lambda}$$

What can we say about the expected value and covariance of  $\vec{\mathbf{X}}(\vec{\lambda})$ ?

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## Statistical Atlases & PCA

For the expected value, we have

$$\begin{aligned} E(\vec{\mathbf{X}}(\vec{\lambda})) &= E(\vec{\mathbf{X}}^{(avg)} + \mathbf{M} \bullet \vec{\lambda}) \\ &= \vec{\mathbf{X}}^{(avg)} + \mathbf{M} \bullet E(\vec{\lambda}) = \vec{\mathbf{X}}^{(avg)} + \mathbf{M} \bullet \vec{\mathbf{0}} \\ &= \vec{\mathbf{X}}^{(avg)} \end{aligned}$$

Then

$$\begin{aligned} \text{cov}(\vec{\mathbf{X}}(\vec{\lambda})) &= E(\vec{\mathbf{D}}(\vec{\lambda}) \bullet \vec{\mathbf{D}}(\vec{\lambda})^T) \quad \text{where } \vec{\mathbf{D}}(\vec{\lambda}) = \vec{\mathbf{X}}(\vec{\lambda}) - \vec{\mathbf{X}}^{(avg)} \\ &= E(\mathbf{M} \bullet \vec{\lambda} \bullet \vec{\lambda}^T \bullet \mathbf{M}) \\ &= \mathbf{M} \bullet E(\vec{\lambda} \bullet \vec{\lambda}^T) \bullet \mathbf{M}^T \\ &= \mathbf{M} \bullet \Sigma^2 \bullet \mathbf{M}^T \end{aligned}$$

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### Statistical Atlases & PCA

Thus, if we assemble a representative sample set of models  $\vec{\mathbf{X}}^{(l)}$ , and compute the average model  $\vec{\mathbf{X}}^{(avg)}$  and the SVD of the corresponding matrix  $\mathbf{D} = \left[ \dots \left( \vec{\mathbf{X}}^{(l)} - \vec{\mathbf{X}}^{(avg)} \right) \right]$ , then we have a way of generating an arbitrary number of models

$$\vec{\mathbf{X}}^{(inst)} = \vec{\mathbf{X}}^{(avg)} + \mathbf{M} \vec{\lambda}^{(inst)} = \vec{\mathbf{X}}^{(avg)} + \sum_k \vec{\mathbf{M}}^{(k)} \lambda_k^{(inst)}$$

with the same mean and covariance. I.e., we know how the individual features  $\vec{\mathbf{x}}_k^{(inst)}$  co-vary.

Further, given a representative model instance  $\vec{\mathbf{X}}^{(inst)}$  we can compute a corresponding set of mode weights  $\vec{\lambda}^{(inst)}$  from

$$\vec{\lambda}^{(inst)} = \mathbf{M}^T \left( \vec{\mathbf{X}}^{(inst)} - \vec{\mathbf{X}}^{(avg)} \right)$$

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### Statistical Atlas

Thus, one representation of a statistical "atlas" of models consists of

- An average model  $\vec{\mathbf{X}}^{(avg)}$
- An eigen matrix  $\mathbf{M}$  of variation modes
- A diagonal covariance matrix  $\Sigma^2$  for the modes

This information may be used in many ways, including

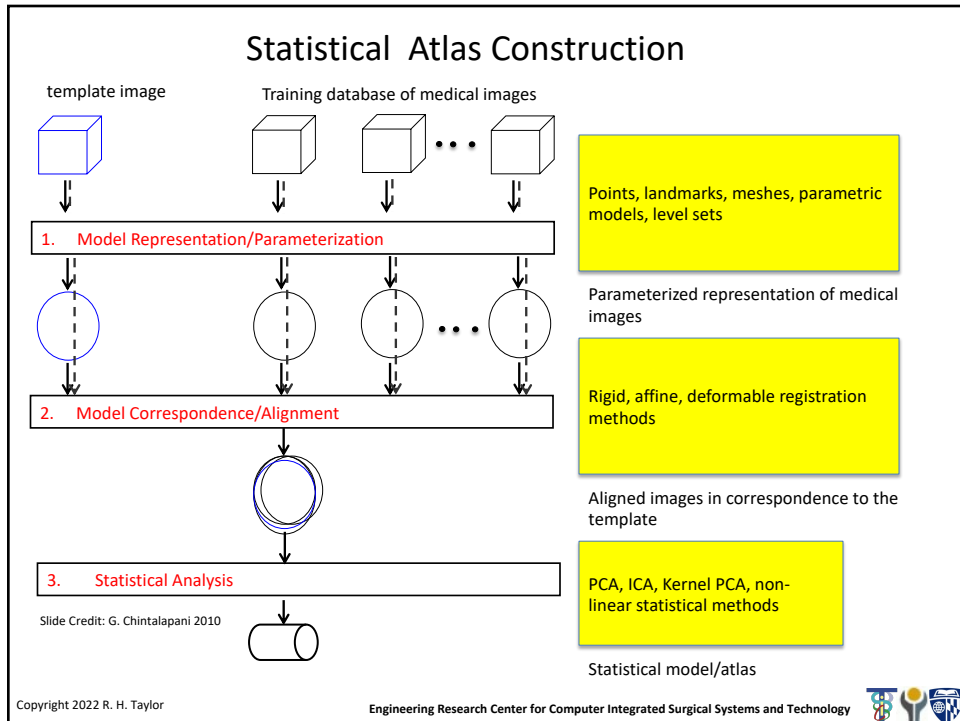
- Atlas-based deformable segmentation/registration
- Statistical analysis of anatomic variation
- etc.

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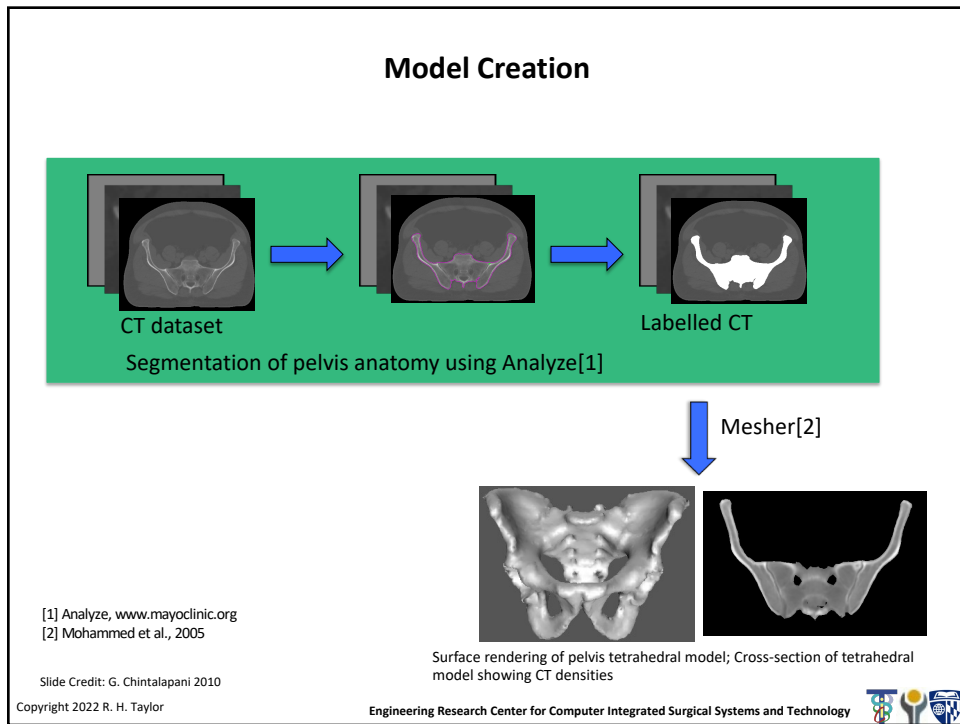
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### Model Representation

- Tetrahedral mesh represents shape
- Bernstein polynomials approximate CT density within each tetrahedron[1,2]

where

$$P^d(\mathbf{u}) = \sum_{|\mathbf{k}|=d} C_{\mathbf{k}} B_{\mathbf{k}}^d(\mathbf{u})$$

$$\mathbf{k} = (k_0, k_1, k_2, k_3) \quad \mathbf{u} = (u_0, u_1, u_2, u_3)$$

$$|\mathbf{k}| = k_0 + k_1 + k_2 + k_3 \quad |\mathbf{u}| = 1$$

$$B_{\mathbf{k}}^d(\mathbf{u}) = \frac{d!}{k_0!k_1!k_2!k_3!} u_0^{k_0} u_1^{k_1} u_2^{k_2} u_3^{k_3}$$

- Alternative is to use voxels directly after deformation to mean shape

[1] Yao, PhD Thesis, 2002; [2] Sadowsky, PhD Thesis, 2008

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Credit: G. Chintalapani 2010

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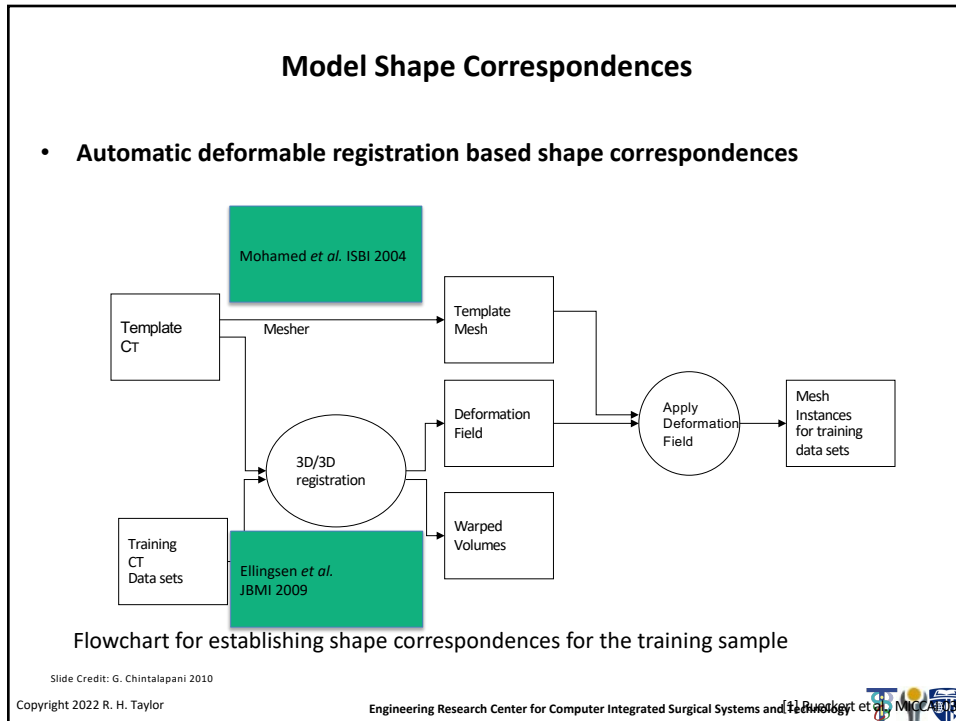
### Model Correspondence

- Need to establish a common coordinate frame for the training database

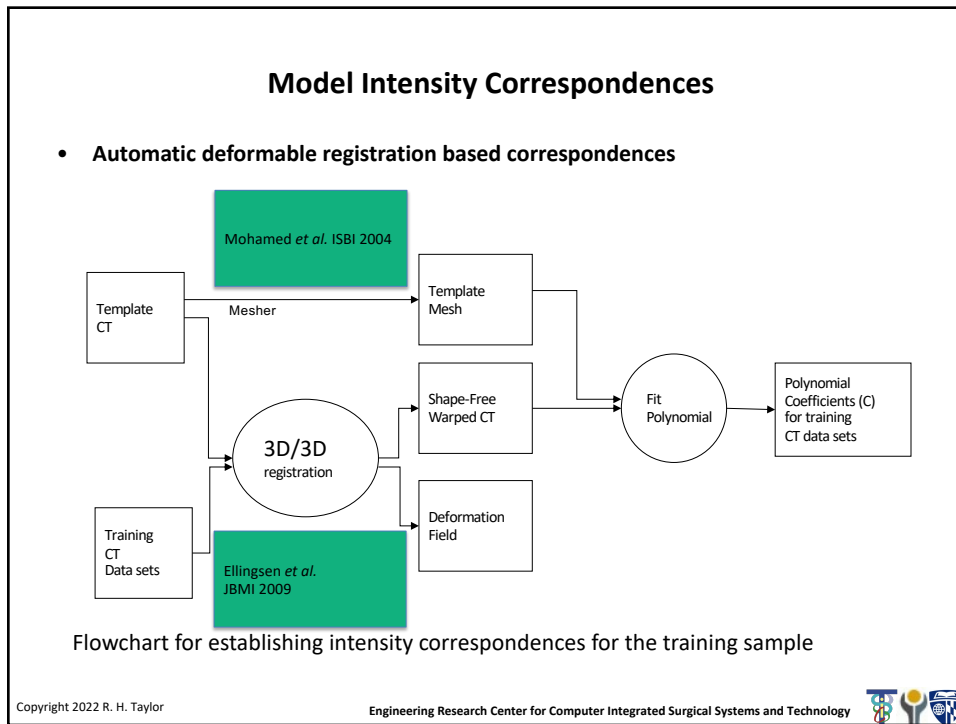
- Need to establish point correspondence between the training datasets

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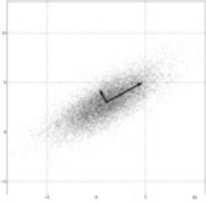
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### Shape Statistics: Principal Component Analysis

- Given N mesh instances of training sample, create matrix of the vertices

$$S = \begin{bmatrix} \hat{x}_1 & \hat{x}_2 & \dots & \hat{x}_N \\ \hat{y}_1 & \hat{y}_2 & \dots & \hat{y}_N \\ \hat{z}_1 & \hat{z}_2 & \dots & \hat{z}_N \end{bmatrix}_{3n \times N} = \begin{bmatrix} x_{11} & x_{12} & \dots & x_{1N} \\ y_{11} & y_{12} & \dots & y_{1N} \\ z_{11} & z_{12} & \dots & z_{1N} \\ \vdots & \vdots & \ddots & \vdots \\ y_{n1} & y_{n2} & \dots & y_{nN} \\ z_{n1} & z_{n2} & \dots & z_{nN} \end{bmatrix}$$


- Compute mean and subtract the mean from the sample
- Compute  $\mathfrak{S} = S - \bar{s} = S - \frac{1}{N} \sum_{i=1}^N \hat{s}_i$


$$SVD(\mathfrak{S}) = UDV^T$$

With principal components in U and eigen values

$$\lambda = \frac{1}{N-1} DD^T$$

- Alternative:** compute SVD of deformation field

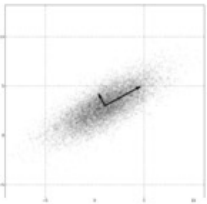
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### Principal Component Analysis

- Given the PCA model, any data instance can be expressed as a linear combination of the principal components

$$\bar{s} + \sum_{k=1}^{N-1} U_k \lambda_k$$


- Compact model → fewer components
- Select first 'd' components represented by the 'd' eigen values

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## Statistical Shape and Intensity Models

- Shape statistical model: Mesh vertices become data matrix

$$\bar{s} + \sum_{k=1}^d U_k \lambda_k = \bar{s} + U^T \lambda$$

- Intensity statistical model: Polynomial coefficients become data matrix

$$\bar{c} + \sum_{k=1}^p Y_k \mu_k = \bar{c} + Y^T \mu$$

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## Deformable Registration Between Shape/Density Atlas and Patient CT

- Goal: Register and Deform the statistical density atlas to match patient anatomy
- Significance:
  - Building patient specific model with same topology (mesh structure) as the atlas
  - Automatic segmentation
  - Accumulatively building models for training set
  - Pathological diagnosis

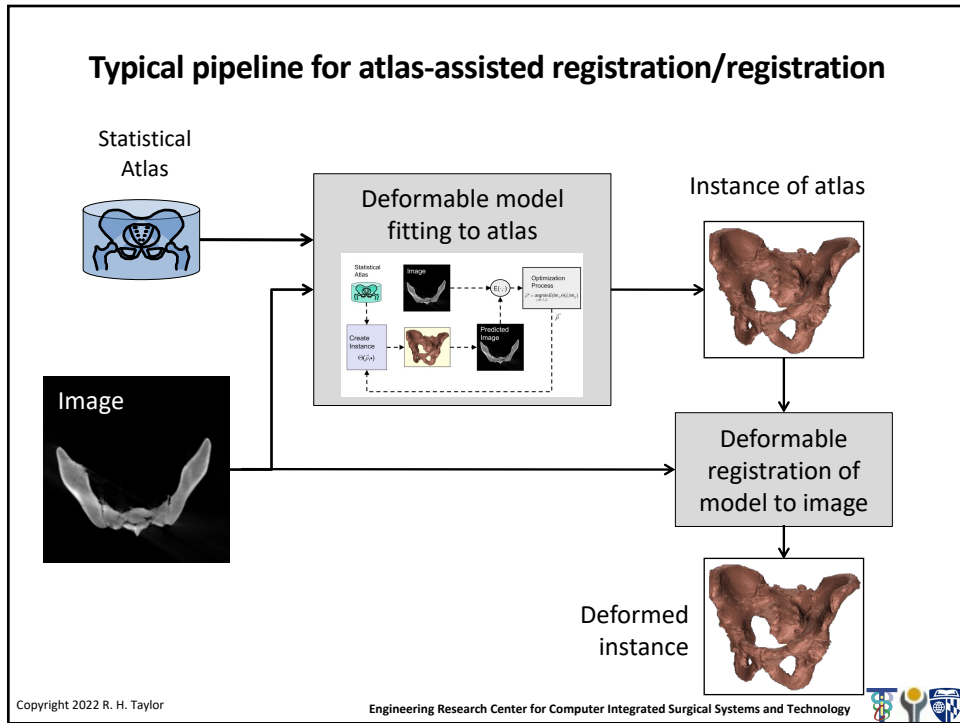
Jianhua Yao

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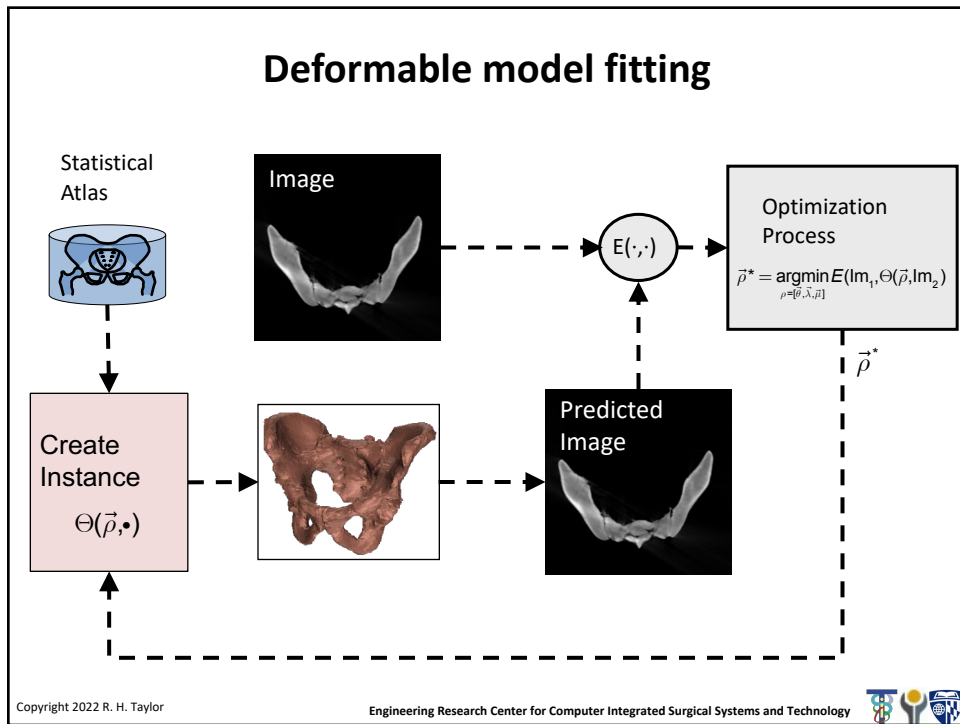
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## Deformable Registration Scheme

- Affine Transformation
  - Translation  $T=(t_x, t_y, t_z)$
  - Rotation  $R=(r_x, r_y, r_z)$
  - Scale  $S=(s_x, s_y, s_z)$  [Similarity if  $s_x=s_y=s_z$ ]
- Global Deformation
  - Statistical deformation mode ( $M_i$ )
- Local Deformation
  - Adjustment of every vertex



## Optimization Algorithm

- Direction Set (Powell's) method in multi-dimensions
  - Search the parameter space to minimize the cost functions
  - Advantage
    - Don't need to compute derivative of cost functions
    - Much fewer evaluations than downhill simplex methods
- Alternatives
  - Downhill Simplex (similar advantages)
  - Covariance Matrix Adaptation Evolution Strategy (CMA-ES) method (similar advantages)
  - Levenberg-Marquardt (requires computing gradients)
  - Many others



## Local Deformation

- Motivation: Statistical deformation can't capture all the variability due to the limited number of models in the training set
- Locally adjust the location of vertices to match the boundary of the bone and the interior density properties
- Use multiple-layer flexible mesh template matching to find the correspondence between model vertices and image voxels
- Apply radial basis function (or other scheme) based on vertex-to-voxel location matches

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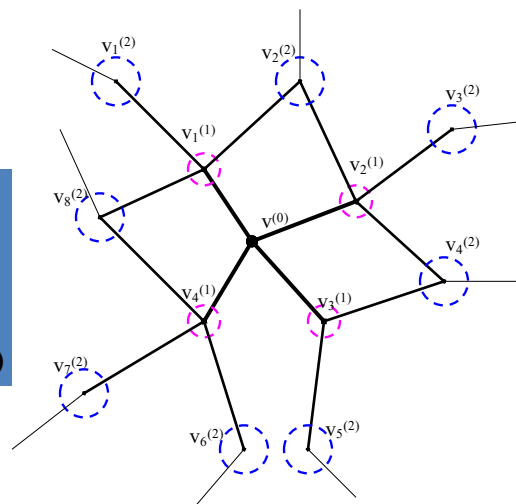


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## Multiple-layer Flexible Mesh Template

- Each vertex on the model defines a mesh template
- Template is in the form

$$(0, \text{Sphere}(v_1^{(1)} - v^{(0)}, r_1), \\ \text{Sphere}(v_2^{(1)} - v^{(0)}, r_1), \dots, \\ \text{Sphere}(v_1^{(2)} - v^{(0)}, r_2), \\ \text{Sphere}(v_1^{(2)} - v^{(0)}, r_2), \dots)$$



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## Template matching

For each pixel location  $\vec{x}_0$  :

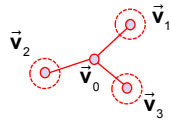
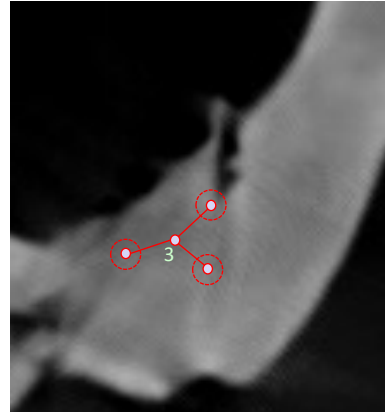
Place  $\vec{v}_0$  at  $\vec{x}_0$

For each neighbor  $\vec{v}_k$

Find the  $\vec{x}_k$  near  $\vec{v}_k$  that minimizes  $E(\vec{x}_k, \vec{v}_k)$

Score  $(\vec{x}_0) = E(\vec{x}_0, \vec{v}_0) + \sum_k w_k E(\vec{x}_k, \vec{v}_k)$

Pick the  $\vec{x}_0$  with the best score



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## Template matching

For each pixel location  $\vec{x}_0$  :

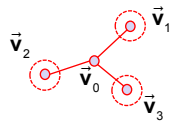
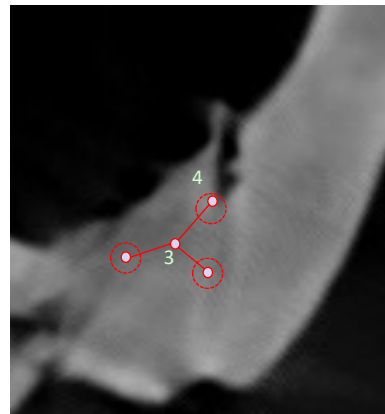
Place  $\vec{v}_0$  at  $\vec{x}_0$

For each neighbor  $\vec{v}_k$

Find the  $\vec{x}_k$  near  $\vec{v}_k$  that minimizes  $E(\vec{x}_k, \vec{v}_k)$

Score  $(\vec{x}_0) = E(\vec{x}_0, \vec{v}_0) + \sum_k w_k E(\vec{x}_k, \vec{v}_k)$

Pick the  $\vec{x}_0$  with the best score



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## Template matching

For each pixel location  $\vec{x}_0$  :

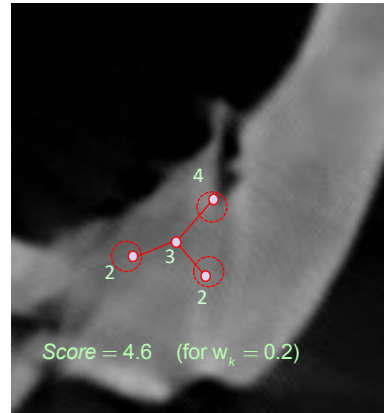
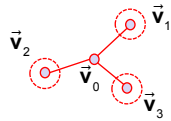
Place  $\vec{v}_0$  at  $\vec{x}_0$

For each neighbor  $\vec{v}_k$

Find the  $\vec{x}_k$  near  $\vec{v}_k$  that minimizes  $E(\vec{x}_k, \vec{v}_k)$

Score ( $\vec{x}_0$ ) =  $E(\vec{x}_0, \vec{v}_0) + \sum_k w_k E(\vec{x}_k, \vec{v}_k)$

Pick the  $\vec{x}_0$  with the best score



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## Template matching

For each pixel location  $\vec{x}_0$  :

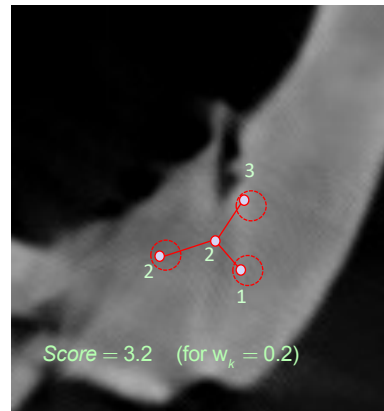
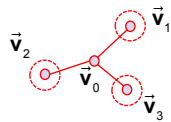
Place  $\vec{v}_0$  at  $\vec{x}_0$

For each neighbor  $\vec{v}_k$

Find the  $\vec{x}_k$  near  $\vec{v}_k$  that minimizes  $E(\vec{x}_k, \vec{v}_k)$

Score ( $\vec{x}_0$ ) =  $E(\vec{x}_0, \vec{v}_0) + \sum_k w_k E(\vec{x}_k, \vec{v}_k)$

Pick the  $\vec{x}_0$  with the best score

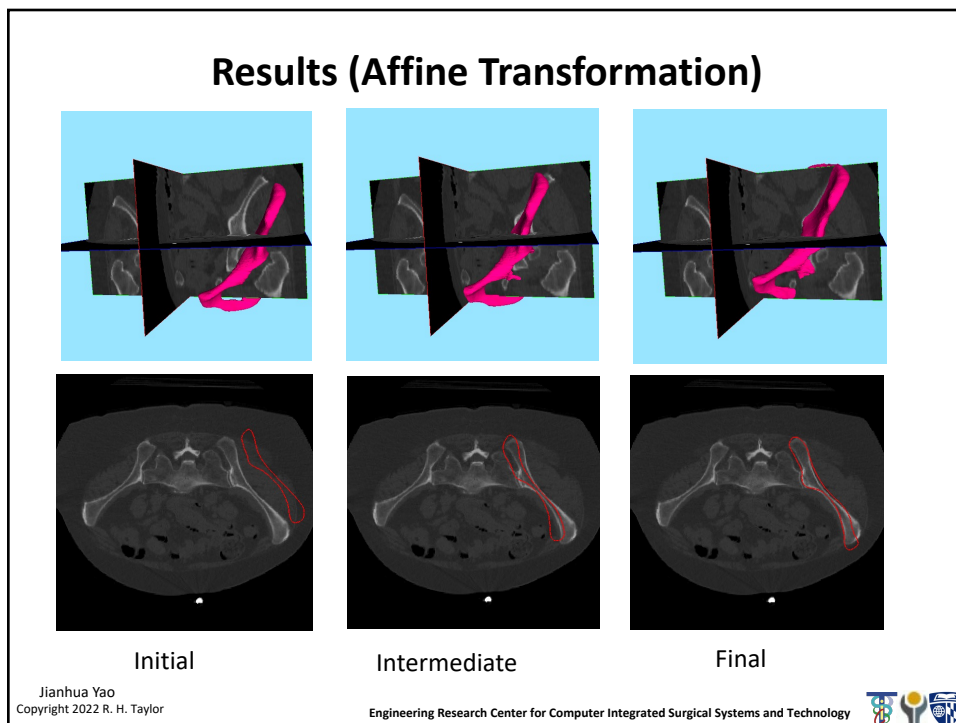


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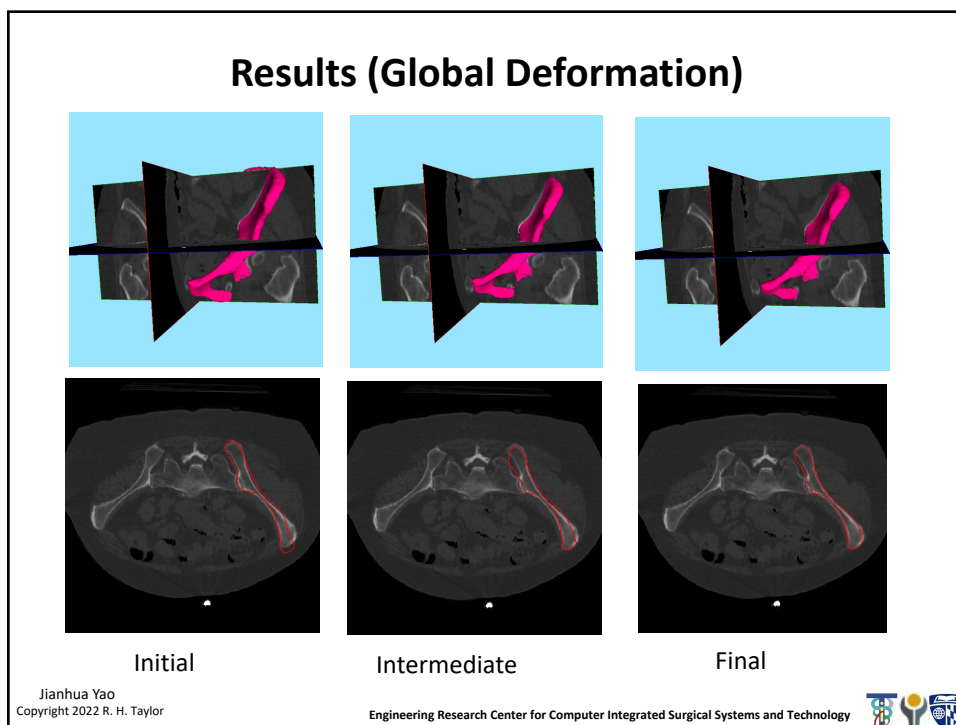
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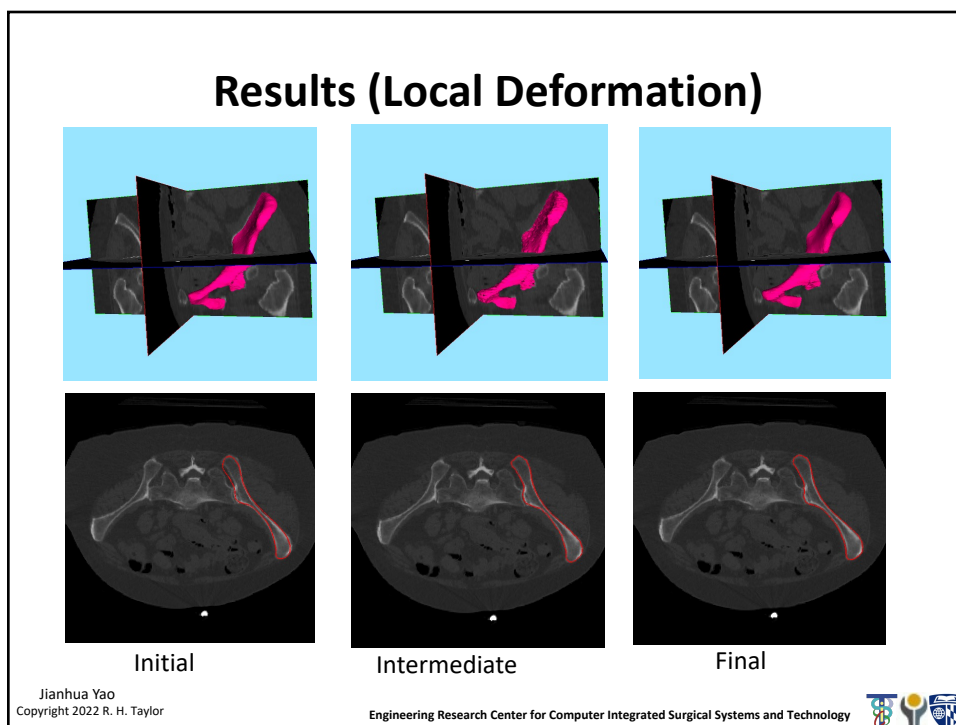
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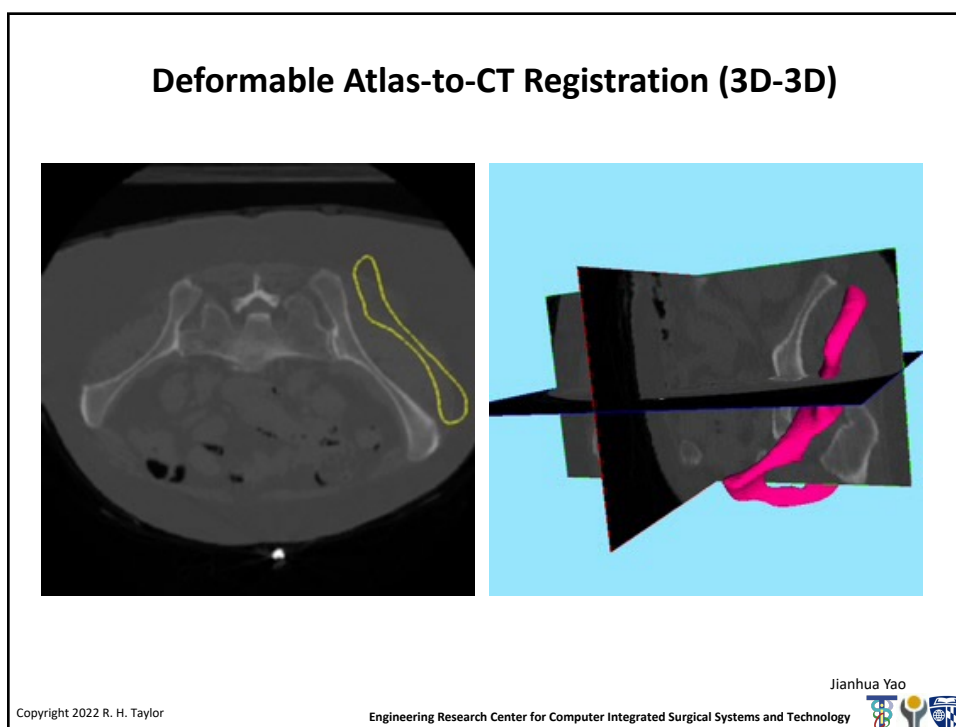
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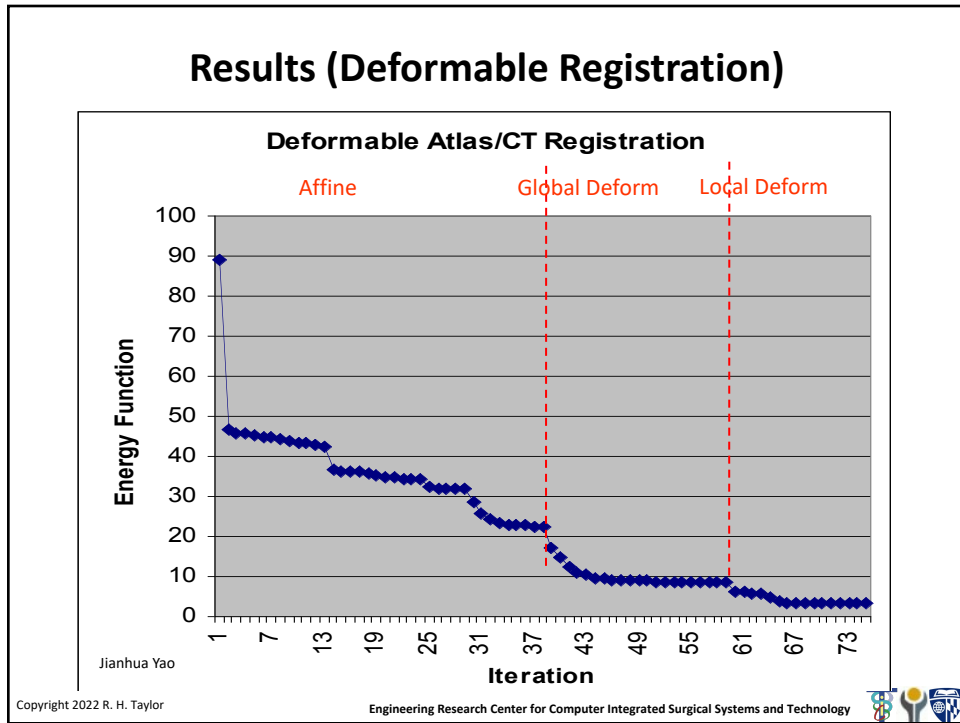


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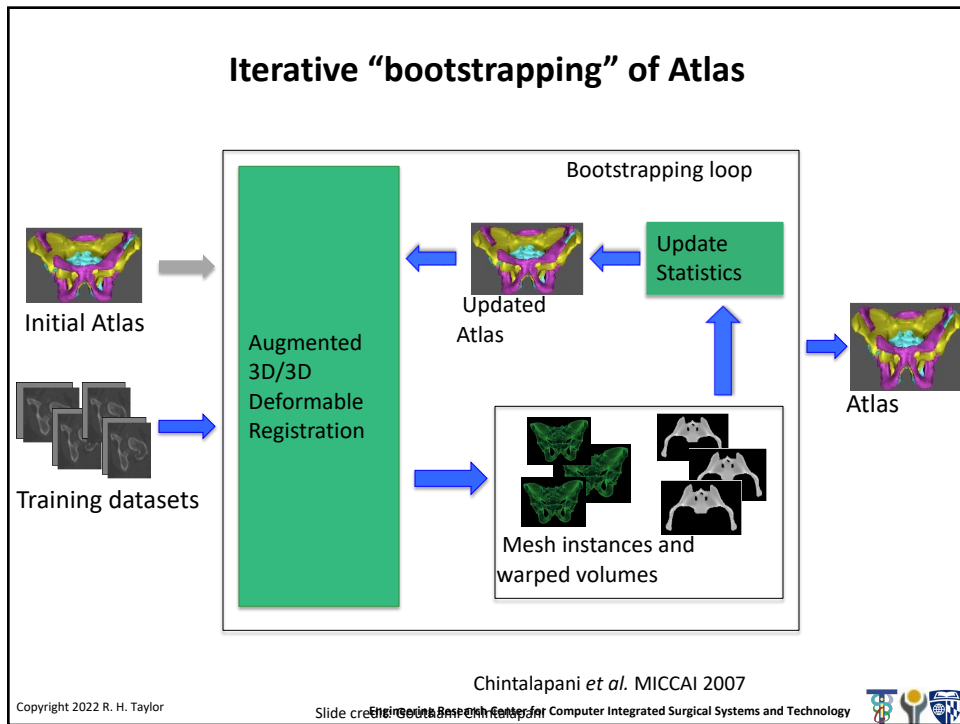


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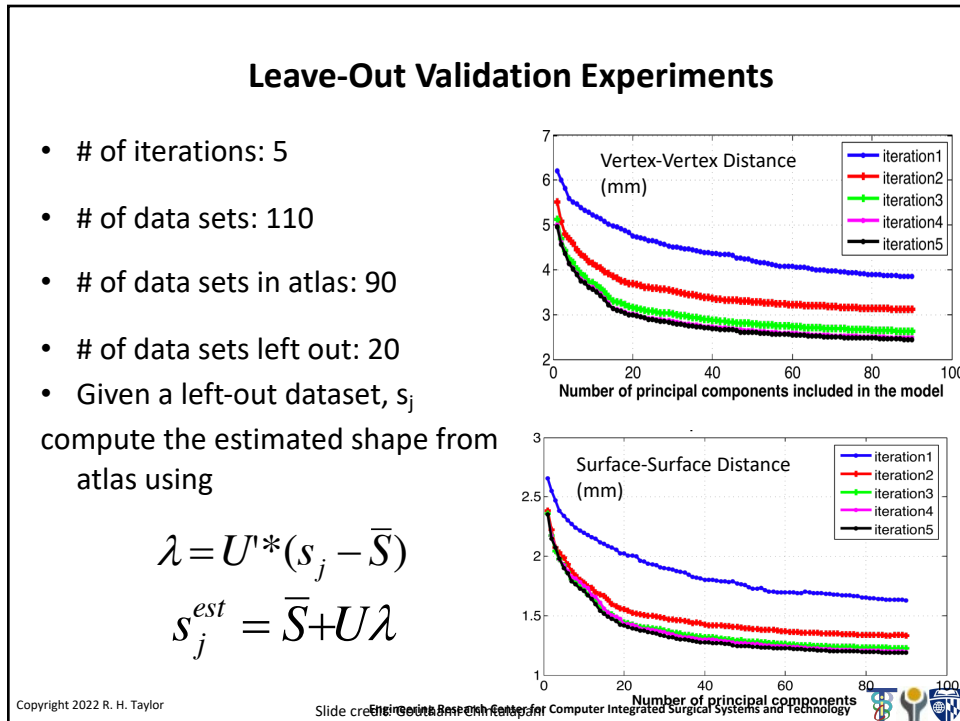




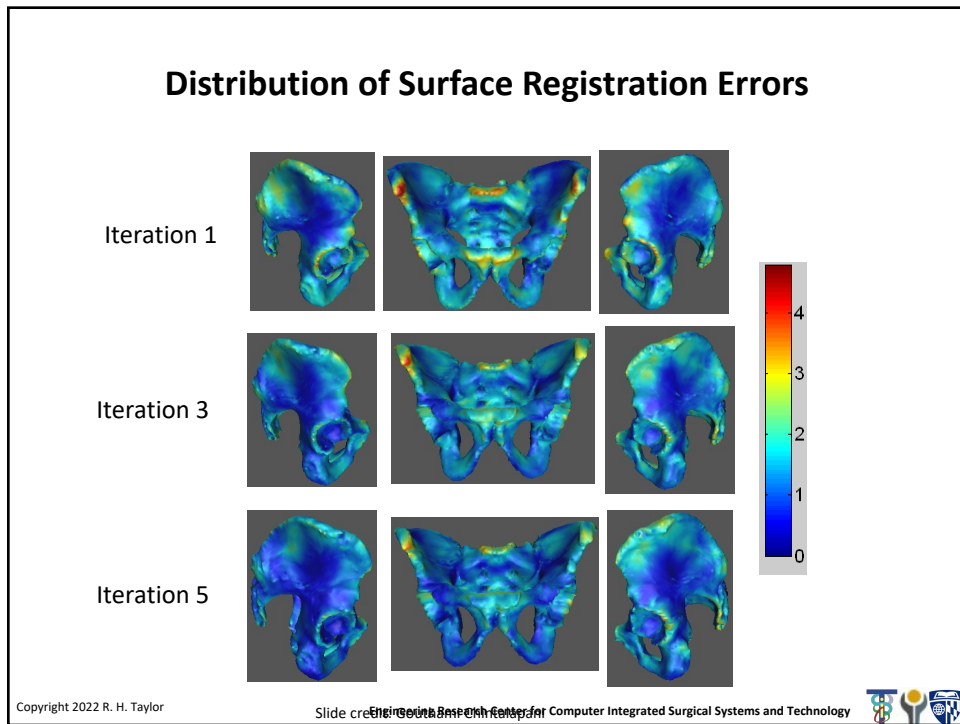
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### Choice of Initial Template

- Claim:
  - iterative method does not depend on the choice of template
- Criteria:
  - Mean shape converges
  - Modes exhibit similar deformation patterns
- Experimental setup:
  - Three random templates
  - Atlases with and without bootstrapping compared
- Result
  - All three atlases exhibit similar deformation patterns after bootstrapping

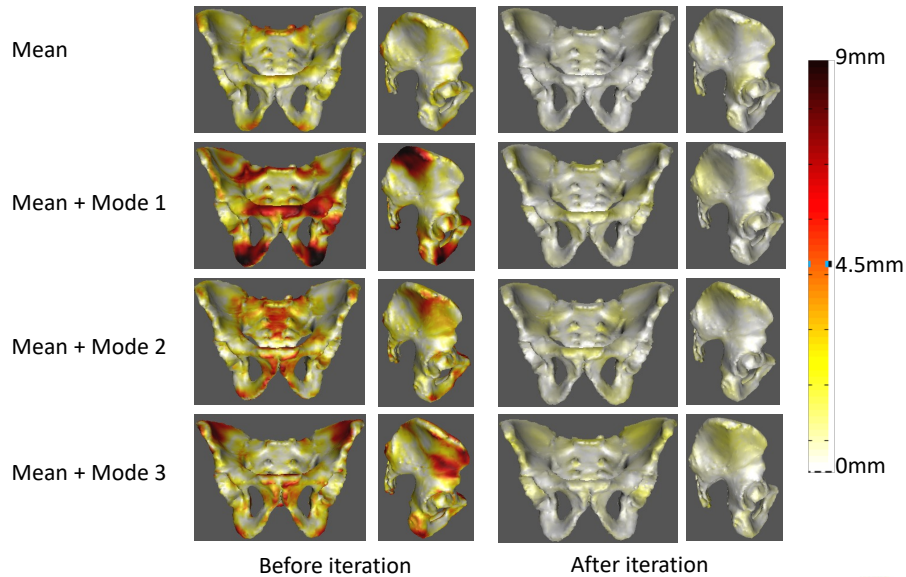
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### Average Difference between Atlases 1,2 and 3



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## Training Sample Size

- Goal:
  - To determine the size of the training sample to build a stable statistical atlas
- Criteria:
  - Atlas is stable
  - No significant improvement in residual error
- Experimental setup:
  - Varying sample size 20, 40, 60, 80
  - Leave-20-out validation test
- Result:
  - Minimum of 50 data sets are required for pelvis atlas

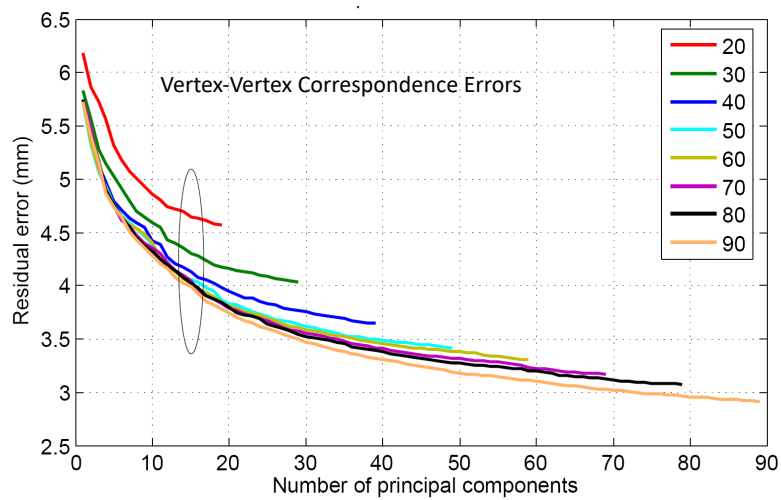
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## Training Sample Size

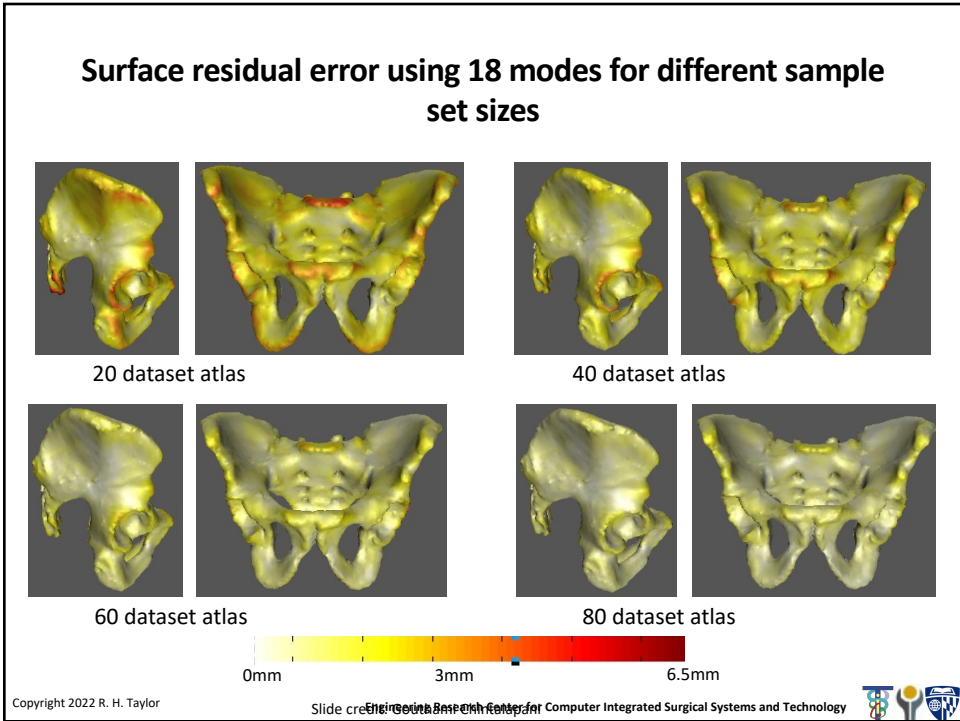


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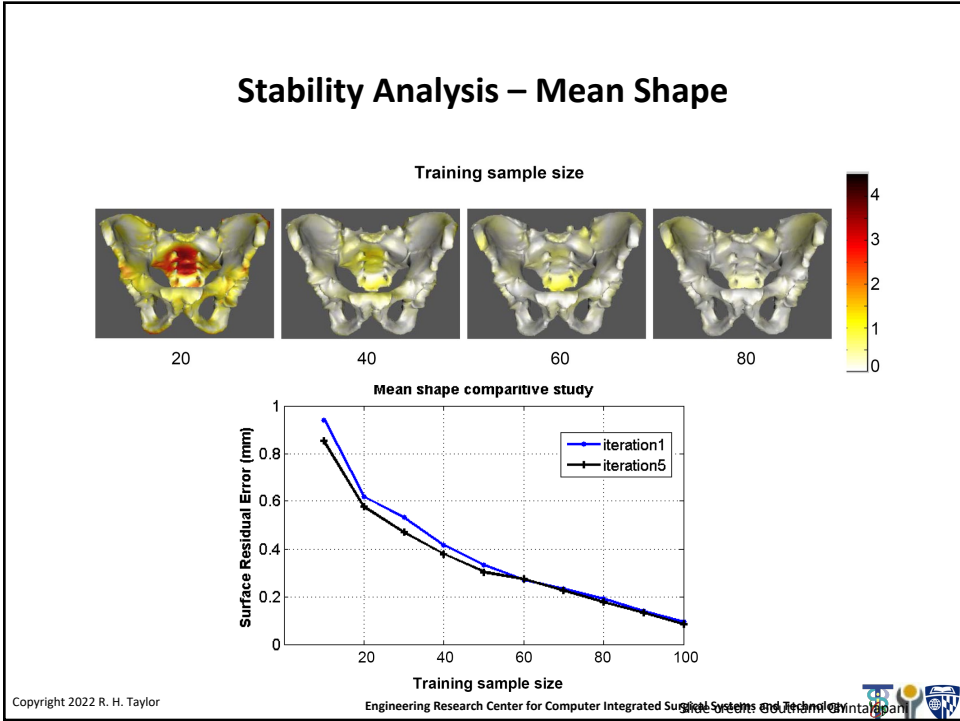
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## Shape Atlas Mesh Refinement

- Note that the methods described so far all assume that the vertices of the mesh after deformable registration all correspond to each other
  - This is often not the case
  - Also, some image segmentation methods we would like to use do not always produce the same surface mesh
  - Is there anything we can do???
- Yes: The basic idea is to do deformable registration of statistical model vertices to the surface(s) to find corresponding points, and then iterate.

Mesh Vertex Improvement  
[\(click here\)](#)

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## Deformable registration between density atlas and a set of 2D X-Rays

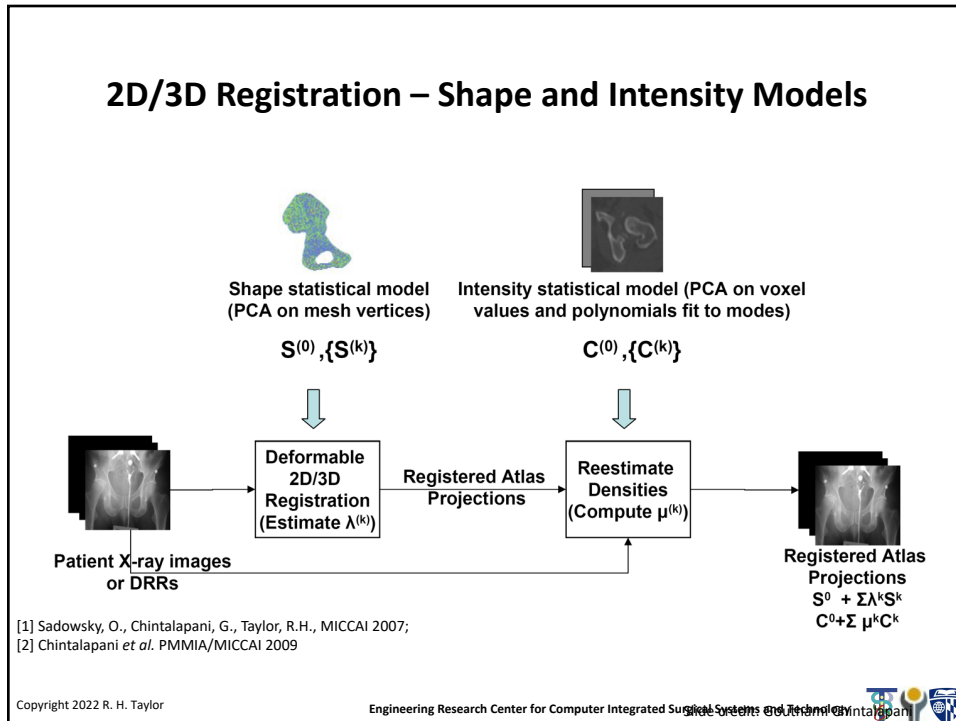
- Goal: Register and Deform the statistical density atlas to match intraoperative x-rays
- Significance:
  - Build virtual patient specific CT without real patient CT
  - Register pre-operative models and intra-operative images
  - Map predefined surgical procedure and anatomical landmarks into intra-operative images

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### 2D/3D Registration – Shape and Intensity

(1)	(2)	(3)	(4)	(5)
#	$S^{true} - S^{est}$ (mm)	RMS ( $V^{true}, V^{est}_{mean}$ ) (HU)	RMS( $V^{true}, V^{est}_{modes}$ ) (HU)	$\frac{\Delta}{((3)-(4))}$ %
1	1.94	109.92	58.88	<b>46.43</b>
2	1.62	128.32	96.0	<b>25.19</b>
3	1.90	98.4	77.12	<b>21.63</b>
4	2.60	51.68	41.6	<b>19.50</b>
5	2.48	109.44	84.8	<b>22.51</b>
6	1.95	73.44	50.56	<b>31.15</b>
7	2.30	72.96	47.52	<b>34.84</b>
8	2.93	101.28	85.76	<b>15.32</b>
avg	<b>2.21</b>	<b>93.18</b>	<b>67.78</b>	<b>27.07</b>

Avg surface registration accuracy: 2.21mm  
 Avg. reduction in RMS errors intensity: 27%

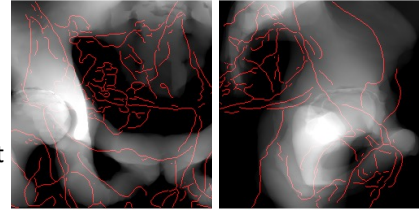
Table 1: Residual errors from leave-out-validation tests of the augmented registration algorithm. Column 2 shows the surface distance after 2D/3D shape registration. Columns 3 shows residual errors when using mean density only and column 4 shows residual errors with mean density and density modes. The % reduction in RMS error between columns 3 and 4 is given in Column 5

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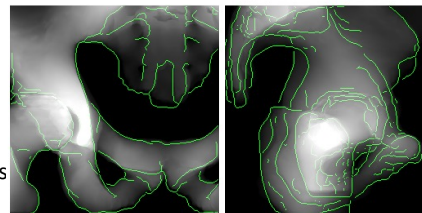
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## 2D/3D Registration – Hip Model

- **Problem:** To create patient specific models using atlas
  - single organ atlases are insufficient
- **Our approach:** Develop a multi-component atlas
  - Use hip atlas instead of a pelvis or femur atlas
  - Extend atlas building framework to incorporate hip joint
  - Extend the registration framework to incorporate articulated hip joint
- **Results**
  - Multi-component atlas registration is accurate compared to individual organ atlas



Pelvis atlas registered to hip projection images



Hip atlas registered to hip projection images

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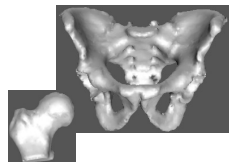
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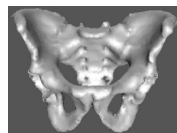
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## Multi-Component Atlas

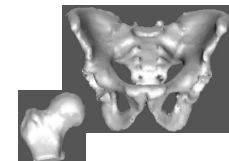
1. Two components – pelvis and femur
2. Create mesh instances of pelvis and femur separately
3. Align pelvis and femur meshes together
4. Align pelvis meshes
5. Align femur meshes
6. Concatenate pelvis and femur meshes
7. PCA on the concatenated mesh



Combined Rigid+Scale



Separate Rigid



Combined Statistical Analysis

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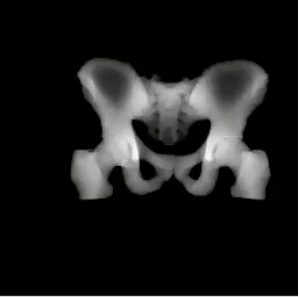
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### Multi-Component Hip Atlas

PC1
PC2
PC3

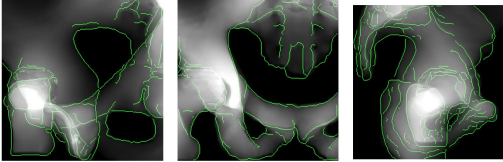
[1] Chintalapani *et al.* CAOS 2009 Slide credit: Gouthami Chintalapani

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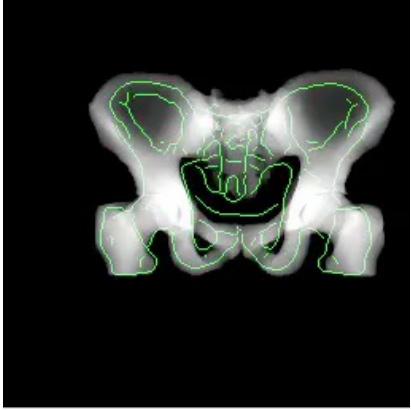
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### 2D/3D Registration – Hip Model

- Registration with truncated images
  - FOV: 160mm
  - Three views
- Avg surface registration accuracy: 2.15 mm



Atlas projections overlaid on DRR images after registration



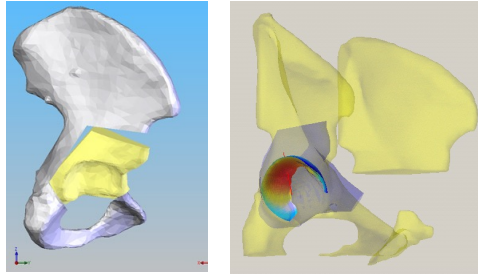
2D/3D deformable registration

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## Applications – Hip Osteotomy



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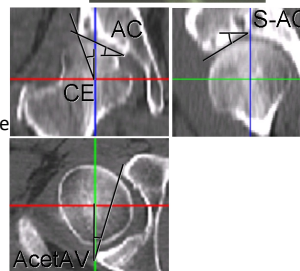
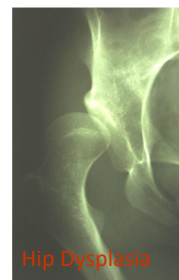
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## Background

- **Hip dysplasia:**
  - Malformation of the hip (normally a ball and socket joint)
  - Significant cause of osteoarthritis, especially in young adults
- **Surgery goals:**
  - Reduce pain symptoms
  - Realign joint to contain the femoral head
  - Diminish risk for degenerative joint changes
  - Improve contact pressure distribution
- **Periacetabular Osteotomy (PAO):**
  - Maintains pelvic structural stability
  - Preserves viable vascular supply
  - Technically challenging tool placement and realignment procedure
- **Limitations of current navigation systems:**
  - Lack the ability to track bone fragment alignment
  - Do not provide anatomical measurements
  - Omit biomechanical-based planning and guidance
  - Ignore the risk of reducing joint range-of-motion



Anatomical measurements used to diagnose hip dysplasia

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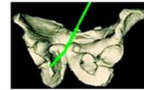
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## Biomechanical Guidance System (BGS)


- **BGS Preoperatively:**
  - Plans surgical cuts
  - Optimizes contact pressures and joint realignment
  - Calculates anatomical-based angles that are meaningful to the surgical team
- **BGS Intraoperatively:**
  - Tracks surgical tools and bone fragment alignment
  - Computes resulting contact pressures
  - Calculates hip range-of-motion
  - Visualizes the surgical cuts
  - Displays radiation-free Digitally Reconstructed Radiographs (DRR)




BGS



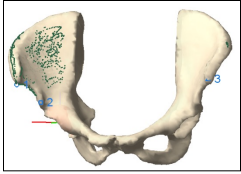
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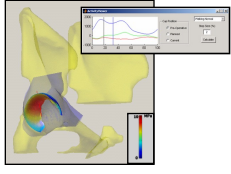
Contact Pressures



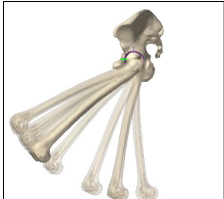
Patient



Model to Patient  
Registration



Joint contact-pressure after PAO



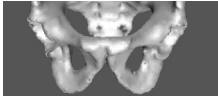
Hip-range-of-motion

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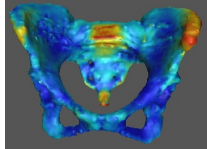
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## Atlas Based Extrapolation of CT

- **Problem:** Partial CT scans of patients
  - Dose minimization for young female patients
  - But the BGS needs full pelvis CT for planning
- **My approach:** Use atlas to predict the missing data
  - Robust probabilistic atlases
  - Improve prediction using pre-op and intra-op x-ray images
- **Preliminary Results**
  - Comparable to the registration errors from full CT scans



Typical pre-operative CT scan of a dysplastic patient undergoing osteotomy



Distribution of surface registration errors of a patient pelvis model estimated from partial CT scan

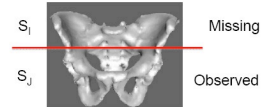
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## Atlas Adaptation to Partial Data

Given a statistical shape model with mean  $\bar{S}$  and modes  $\mathbf{U} = \{U^{(1)} \dots U^{(M)}\}$   
Rearrange vertex indices and partition model into components corresponding to known and unknown parts

$$\bar{S} = \begin{bmatrix} \bar{S}_I \\ \bar{S}_J \end{bmatrix} \quad \mathbf{U} = \begin{bmatrix} \mathbf{U}_I \\ \mathbf{U}_J \end{bmatrix}$$



Find a set of registration parameters  $(s, \mathbf{R}, \vec{p}, \vec{\lambda})$

$$(s, \mathbf{R}, \vec{p}, \vec{\lambda}) = \operatorname{argmin} \left\| S_J^{(obs)} - \left( s\mathbf{R}(\bar{S}_J + \mathbf{U}_J \vec{\lambda}) + \vec{p} \right) \right\|$$

Estimate the total shape as

$$S^{(est)} = \begin{bmatrix} \left( s\mathbf{R}(\bar{S}_I + \mathbf{U}_I \vec{\lambda}) + \vec{p} \right) \\ S_J^{(obs)} \end{bmatrix}$$

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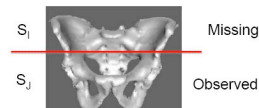
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## Atlas Adaptation to Partial Data with Xray Images



➤ 2D/3D registration[2] of inferred data with X-ray images

$$(s, \mathbf{R}, \vec{p}, \vec{\lambda}) = \operatorname{argmax} \sum_k MI(I_k, DRR(DensityAtlas, s\mathbf{R}(\bar{S}_J + \mathbf{U}_J \vec{\lambda}) + \vec{p}))$$

➤ Final atlas extrapolated model is given as

$$S^{(est)} = \begin{bmatrix} \left( s\mathbf{R}(\bar{S}_I + \mathbf{U}_I \vec{\lambda}) + \vec{p} \right) \\ S_J^{(obs)} \end{bmatrix}$$

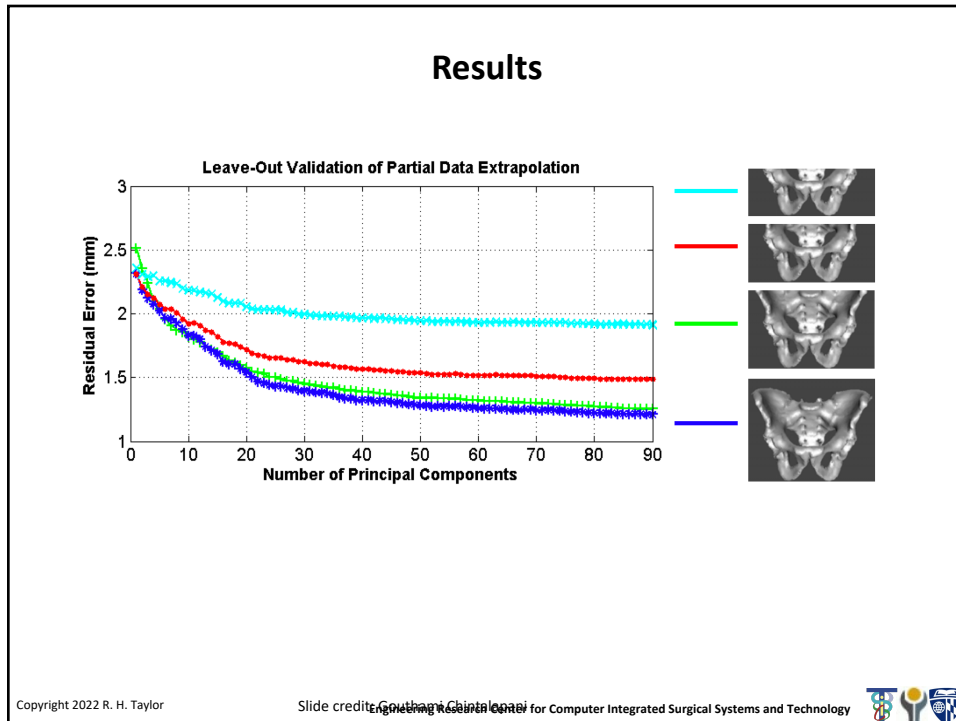
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### Results – Atlas Experiments

#	Full CT				Partial CT				Partial CT + X-ray			
	mean	max	std	95%	mean	max	std	95%	mean	max	std	95%
1	1.41	8.20	1.06	3.45	1.97	14.06	1.69	5.17	1.37	10.94	1.13	3.54
2	1.88	7.25	1.42	4.71	2.15	12.25	1.73	5.28	1.73	14.78	1.71	4.51
3	1.55	7.72	1.20	3.77	2.45	11.33	2.08	6.89	1.41	6.81	1.10	3.54
4	1.32	5.77	1.01	3.27	1.69	9.06	1.43	4.58	1.21	6.80	1.03	3.27
5	1.72	8.29	1.17	3.79	1.62	6.87	1.24	3.93	1.36	8.17	1.13	3.61
6	1.69	10.58	1.55	4.78	2.64	14.87	2.27	7.18	1.71	11.33	1.54	5.06
avg	1.59	7.96	1.23	3.96	2.08	11.40	1.74	5.50	1.46	9.80	1.27	3.92

Atlas inferred CT using full CT scan

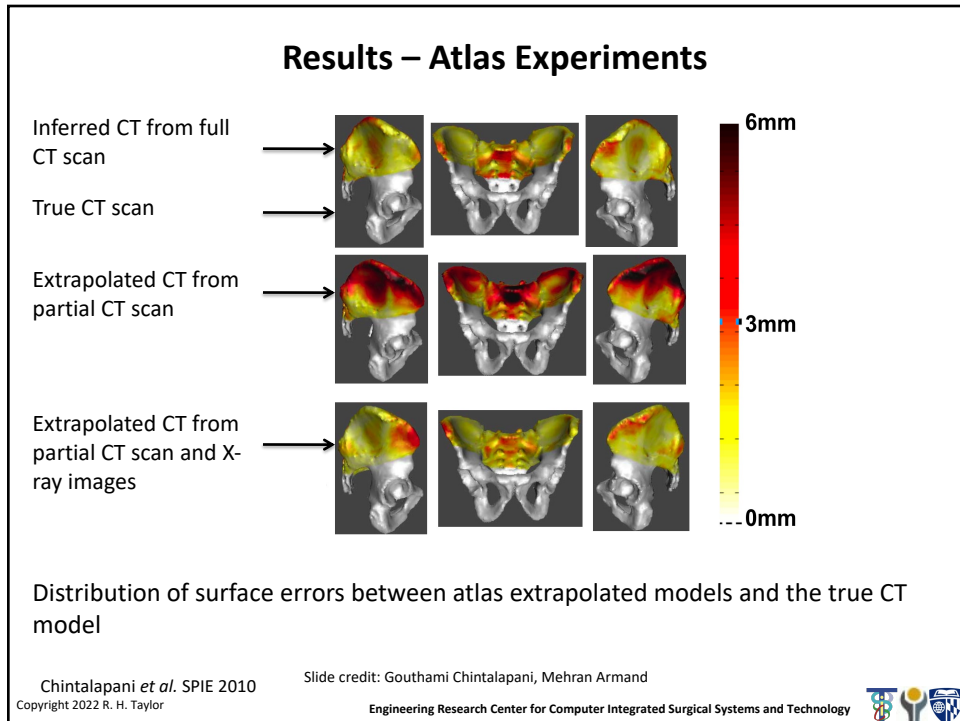
Atlas extrapolated CT using partial CT scan

Atlas extrapolated CT using partial CT scan and X-ray images

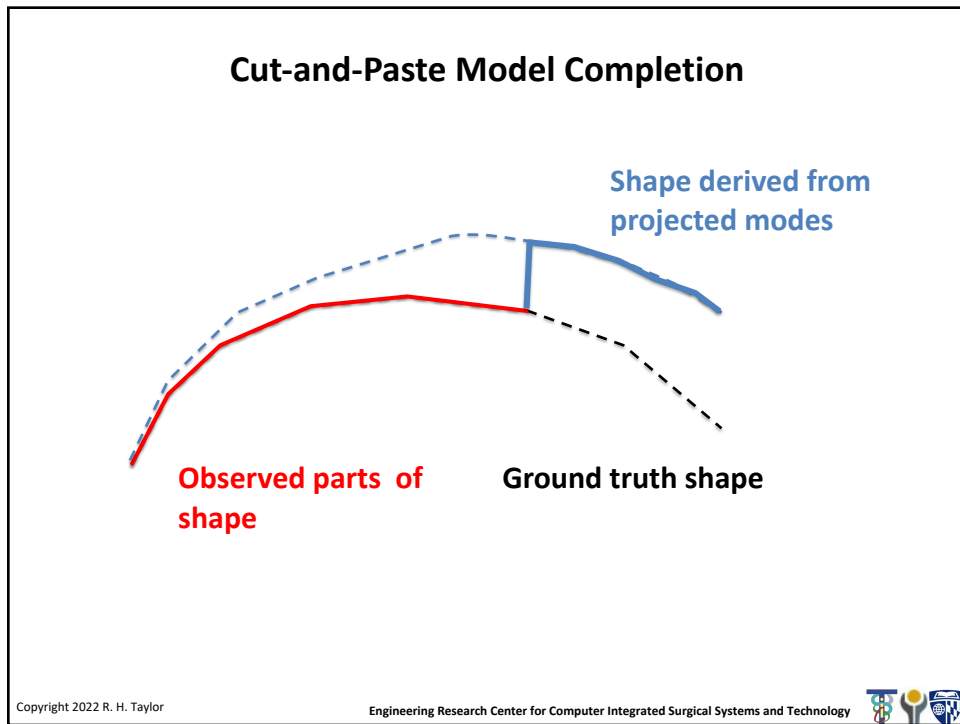
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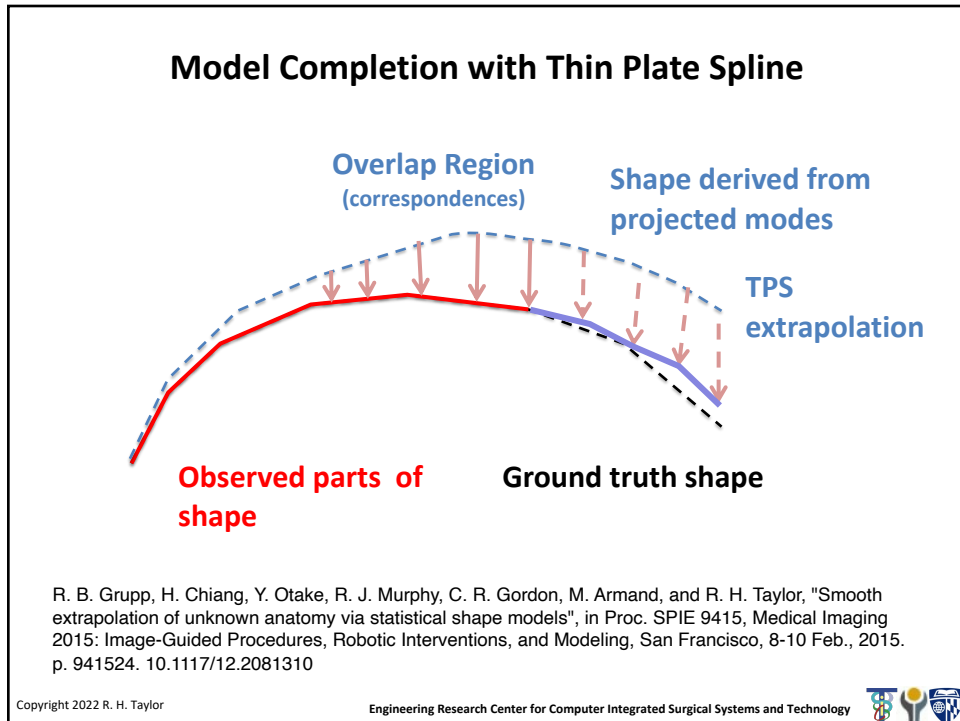
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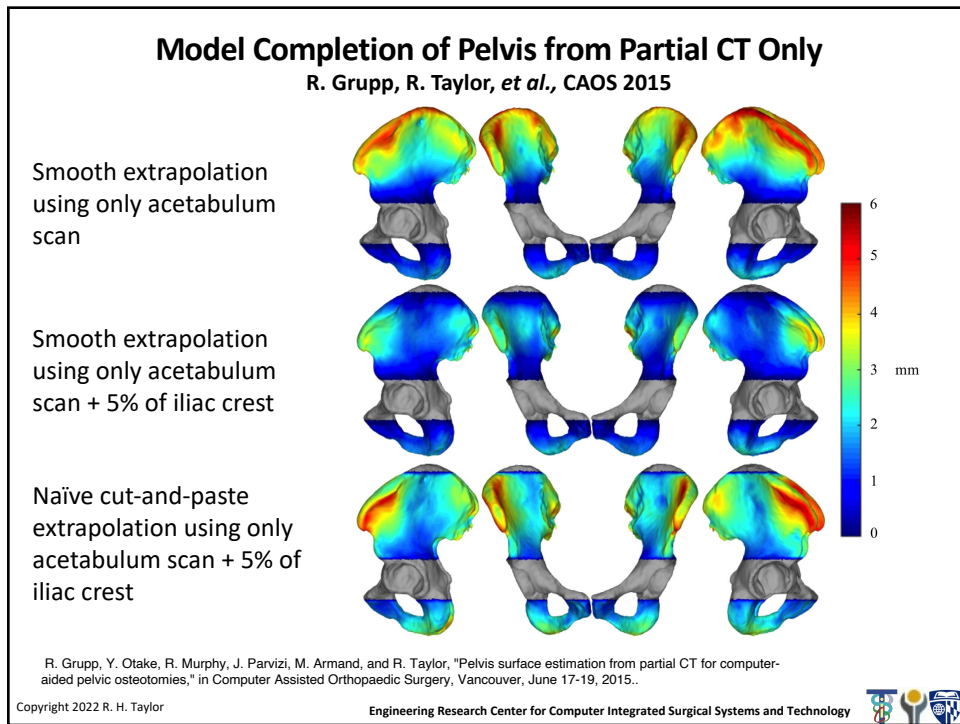
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## Osteotomy Simulations

➤ Atlas extrapolated model is used primarily for two reasons:

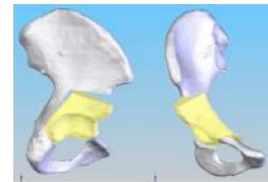
1. Model to patient registration

- simulation experiments
- six leave out experiments
- FRE error metric



2. Fragment tracking

- Simulated osteotomy cuts
- Applied known transformation to the Fragment
- Computed the fragment transformation
- Compared it to the known transformation

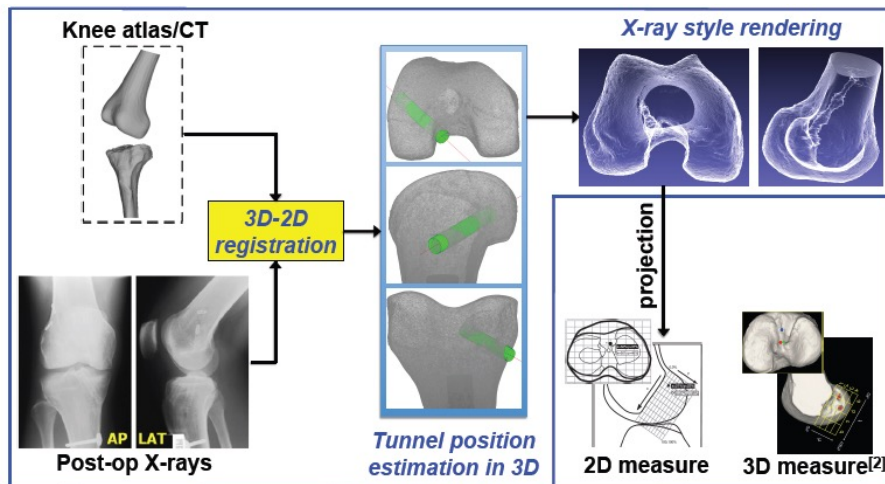


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## Statistical Assessment of ACL Tunnel Positions

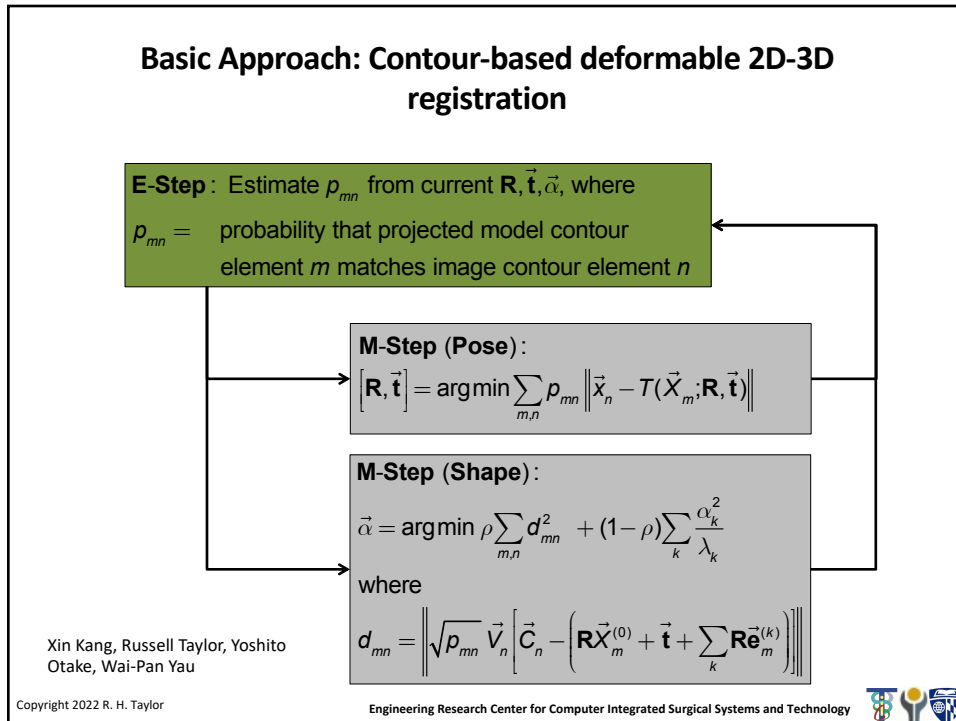
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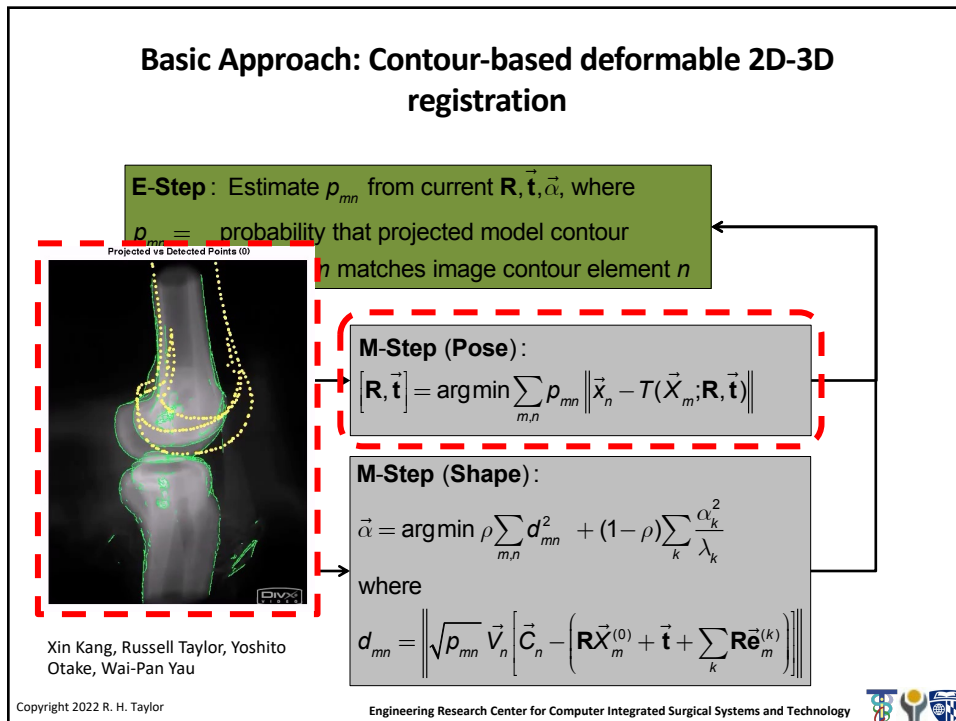
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### Basic Approach: Contour-based deformable 2D-3D registration

# of iteration	Max err (mm)
#1	10.32
#2	7.50
#3	6.63
#4	6.17
#5	5.33
#6	4.98
#7	4.64
#8	3.95
#9	3.81
#10	3.29
Final	3.02

Current  $\mathbf{R}, \vec{\mathbf{t}}, \vec{\alpha}$ , where  
 deformed model contour  
 image contour element  $n$

(Pose):  

$$\arg \min \sum_{m,n} \rho_{mn} \|\vec{\mathbf{x}}_n - T(\vec{\mathbf{X}}_m; \mathbf{R}, \vec{\mathbf{t}})\|$$

M-Step (Shape):  

$$\vec{\alpha} = \arg \min \rho \sum_{m,n} d_{mn}^2 + (1-\rho) \sum_k \frac{\alpha_k^2}{\lambda_k}$$
 where  

$$d_{mn} = \left\| \sqrt{\rho_{mn}} \vec{\mathbf{V}}_n \left[ \vec{\mathbf{C}}_n - \left( \mathbf{R} \vec{\mathbf{X}}_m^{(0)} + \vec{\mathbf{t}} + \sum_k \mathbf{R} \vec{\mathbf{e}}_m^{(k)} \right) \right] \right\|$$

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### C-arm Distortion

➤ What is distortion ?

- Avg distortion: **2.14 mm/pixel**
- max distortion: **4.60 mm/pixel**

➤ How to rectify images ?

- Phantom based correction
- Polynomial functions to model distortion

Example C-arm images showing distortion, straight metal wires appear curved due to distortion

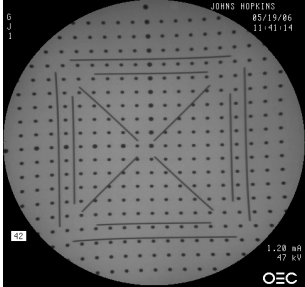
Typical bi-planar phantom used for C-arm calibration

$$(u_d, v_d) = \sum_{i=0}^n \sum_{j=0}^n C_{ij} B_{ij}(u_0, v_0)$$

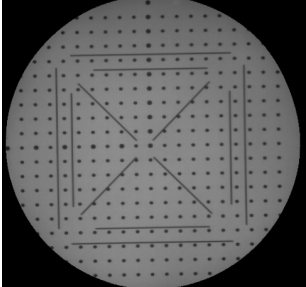
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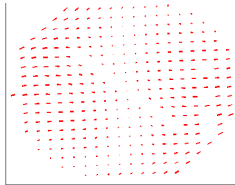
## C-Arm Distortion Correction



Warped X-ray image of the phantom



Dewarped X-ray image



Distortion vector map

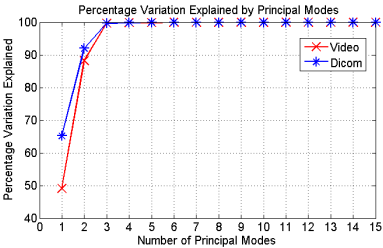
$$\vec{\Delta d} = (\Delta u, \Delta v) = (u_d, v_d) - (u_0, v_0)$$

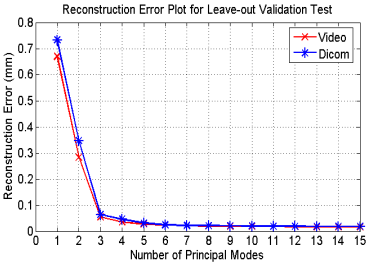
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## Statistical Characterization of C-Arm Distortion correction using PCA

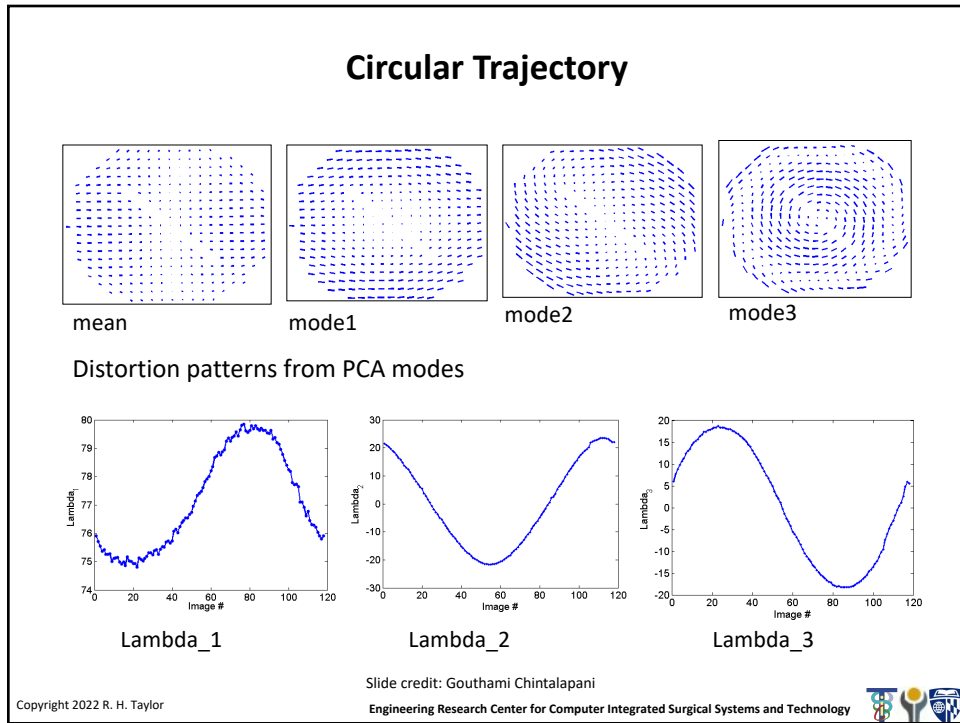
- Principal component analysis on distortion maps
  - 120 images, one every 3 degrees approx., along propeller axis (similar to the full sweep data used for 3D reconstruction)
  - 200 images to span the sphere defined by the "C" of the c-arm



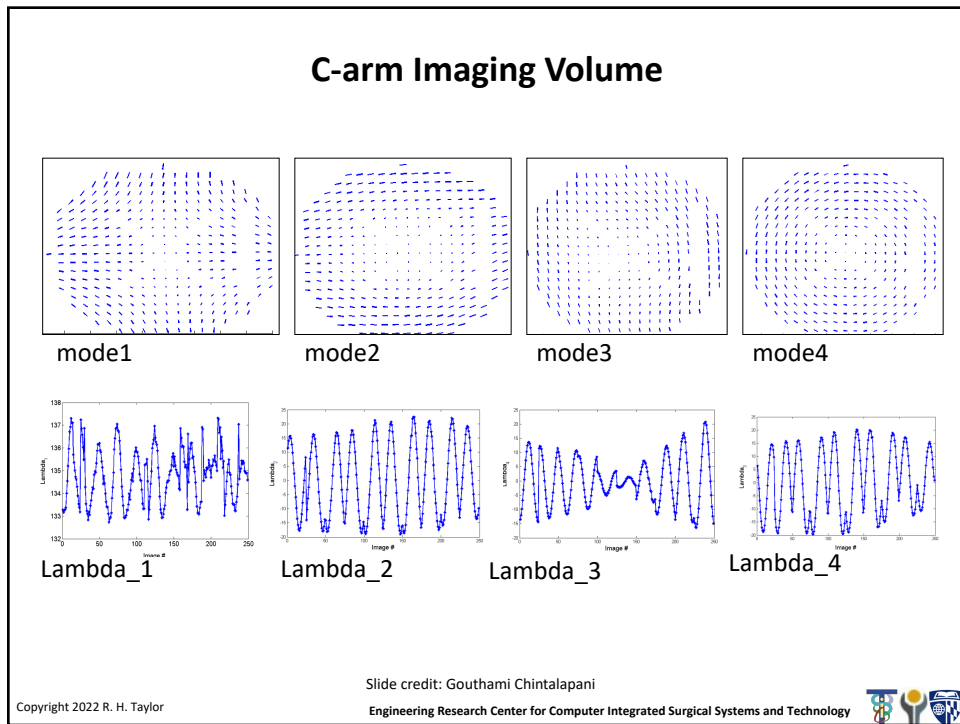


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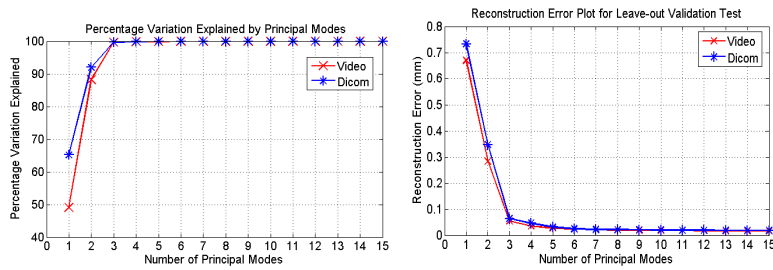
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### Eigen Analysis of Distortion Maps

- First three modes are significant and explain 99% of the variation
- Leave-out validation tests indicate that the distortion parameters can be recovered with an accuracy of less than 0.1 mm/pixel.



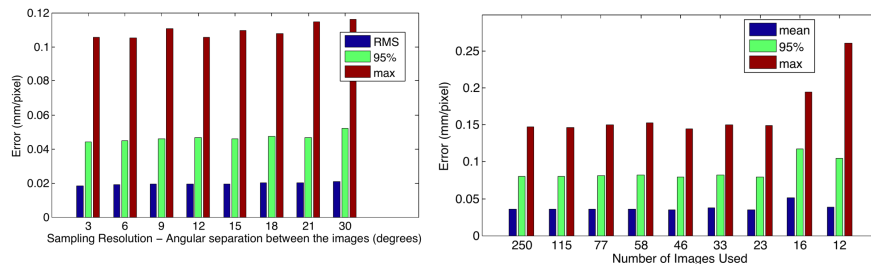
$$\epsilon = \left\| \Delta \vec{d} - \left( M_0 + \sum_{i=1}^n \lambda_i D_i \right) \right\|^2$$

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### Sampling Resolution

- How many images are required to statistically characterize the distortion patterns ?



Circular Trajectory

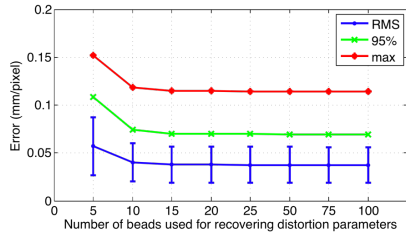
C-arm Imaging Volume

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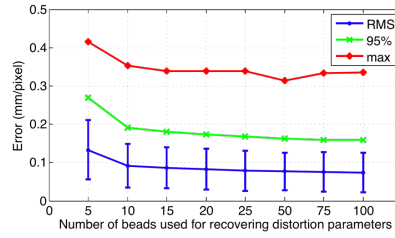
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### Recovering Distortion Parameters

- Use as few beads as possible to recover the distortion mode parameters



Circular Arc



C-arm Imaging Volume

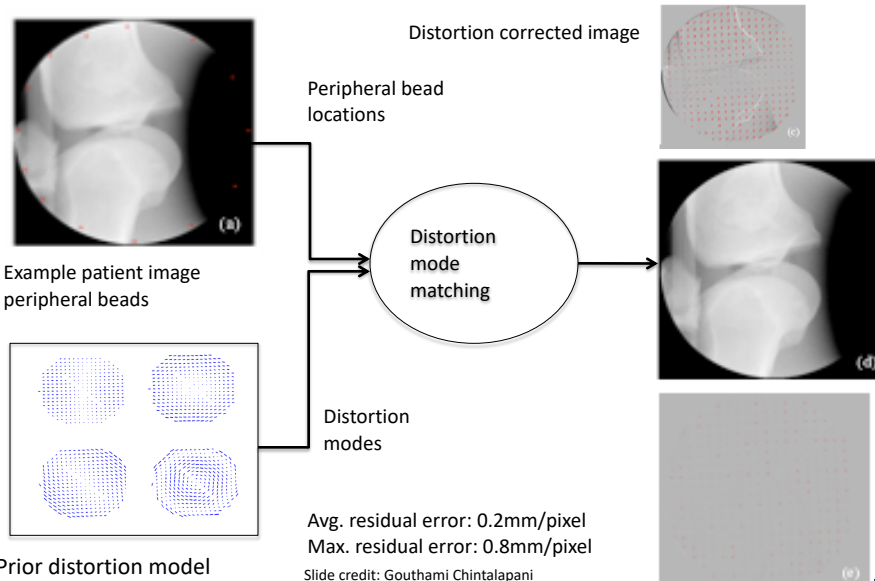
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### Small Phantom based Distortion Correction



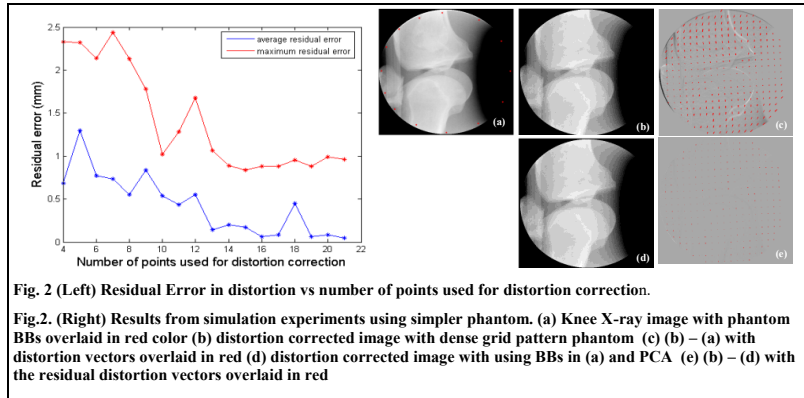
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## Small Phantom based Distortion Correction



Statistical Characterization of C-arm Distortion with Intra-operative Application

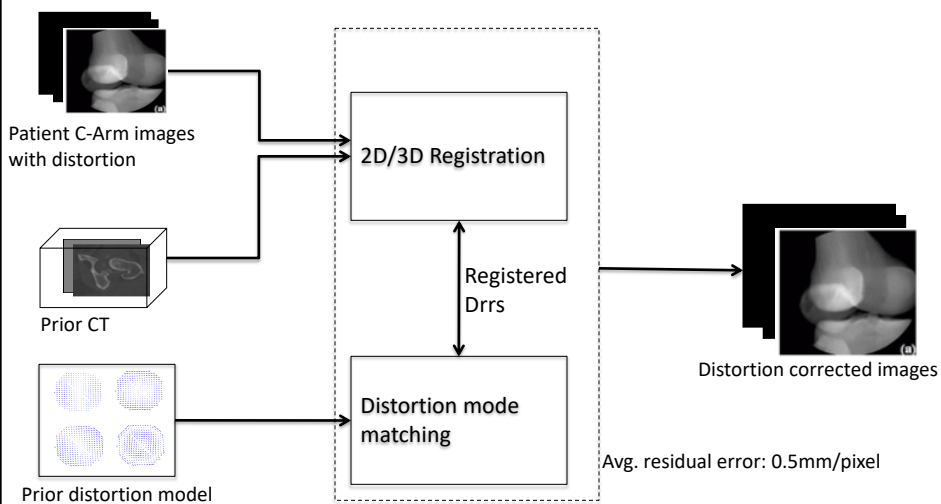
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## Using Patient CT as Fiducial



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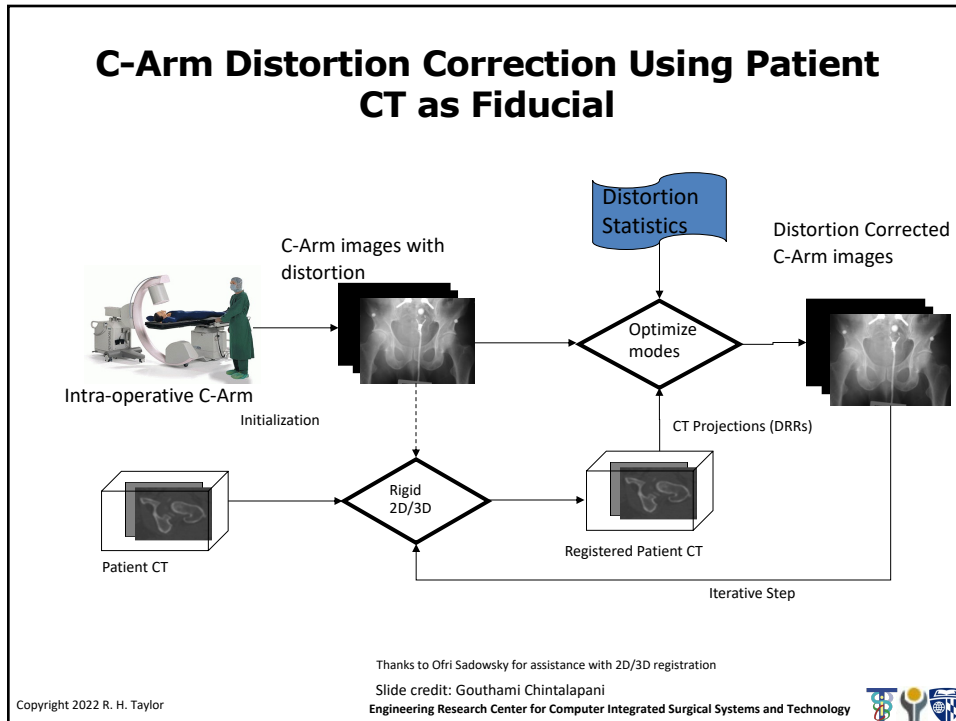
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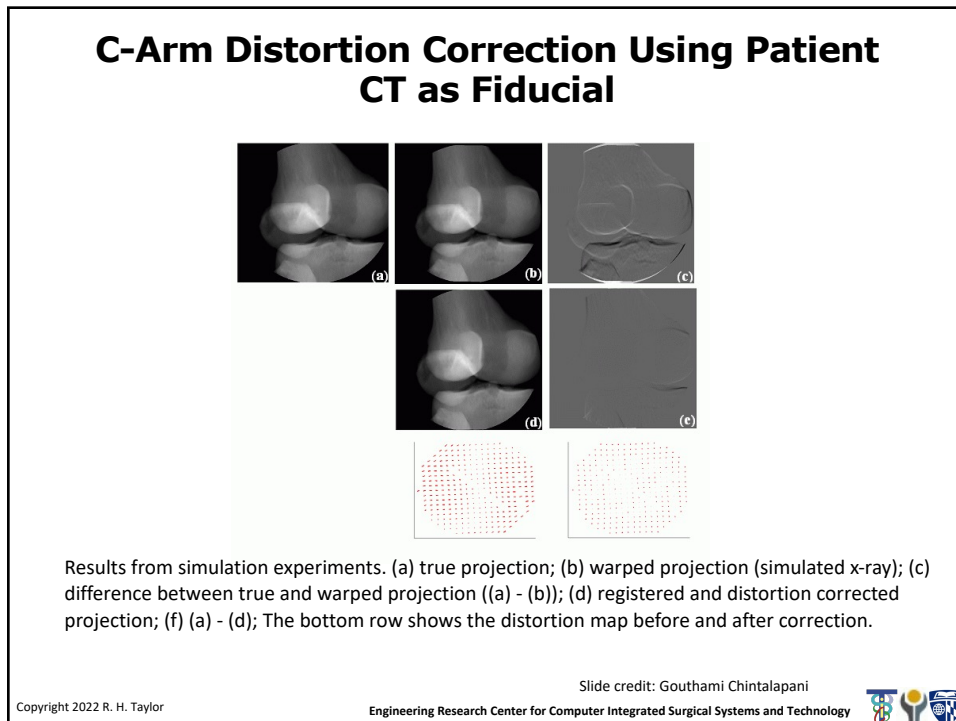
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