

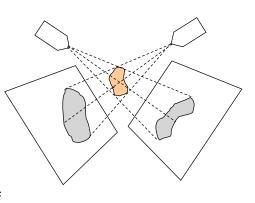
Feature-Based 2D-3D Registration

Given

- 3D surface model of an anatomic structure
- Multiple 2D x-ray projection images taken at known poses relative to some coordinate system C
- Initial estimate of the pose F of the anatomic object relative to the x-ray imaging coordinate system C

Goal

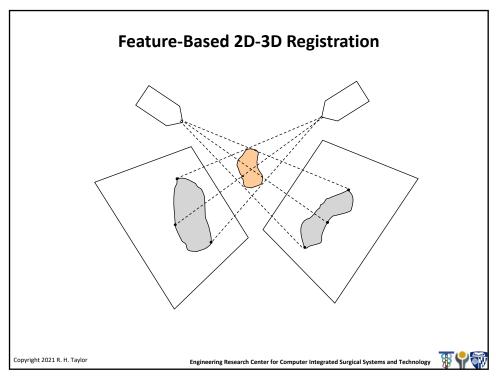
- Compute an accurate value for F

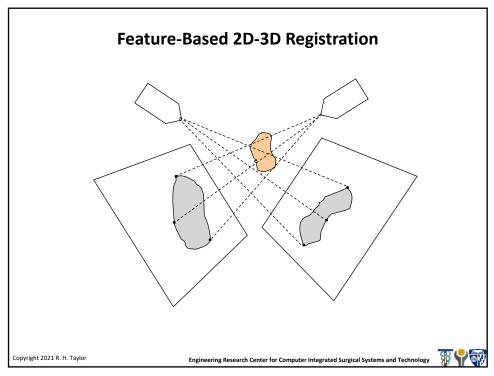


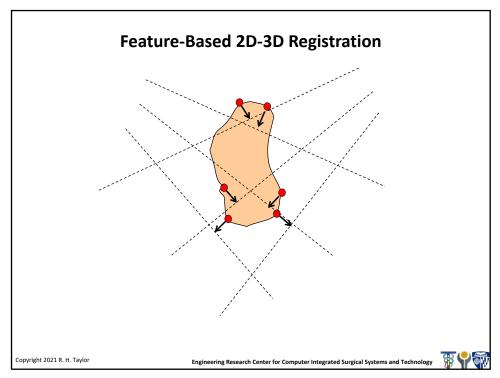
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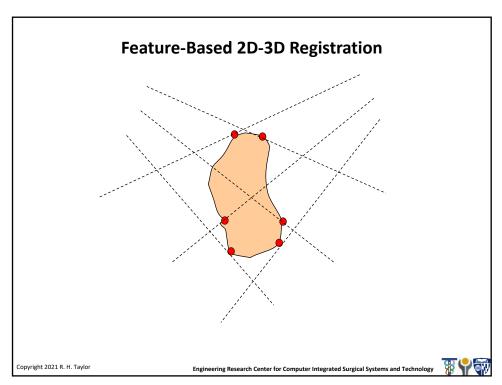
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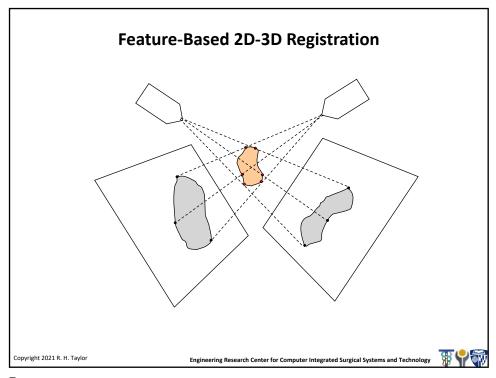


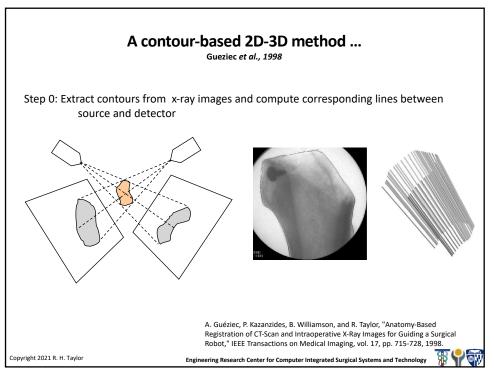










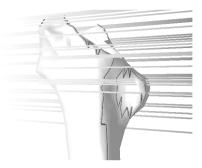


A contour-based 2D-3D method ...

Gueziec et al., 1998

Step 1: Given the current estimate for F = [R,t], compute the apparent projection contours of the model for each viewing direction.

Step 2: For each x-ray path line line L_i , identify the closest point p_i on an apparent projection contour. This will give a set of points on the body surface to be moved toward the corresponding x-ray lines



A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

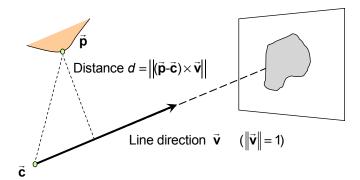
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A contour-based 2D-3D method ...

Gueziec et al., 1998



Note: It is convenient to use the x-ray source position (i.e., the center of convergence for a bundle of x-ray projection lines) as the value for $\vec{\mathbf{c}}$.

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

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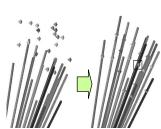
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A contour-based 2D-3D method ...

Gueziec et al., 1998

Step 3: Solve an optimization problem to compute a value of F that minimizes the distance between the p_i and the L_i.



$$\min_{\mathbf{R}, \hat{\mathbf{t}}} \sum_{i} d_{i}^{2} = \min_{\mathbf{R}, \hat{\mathbf{t}}} \sum_{i} \left\| \vec{\mathbf{v}}_{i} \times \left(\mathbf{c}_{i} - \left(\mathbf{R} \vec{\mathbf{p}}_{i} + \vec{\mathbf{t}} \right) \right) \right\|^{2}$$

$$= \min_{\mathbf{R}, \hat{\mathbf{t}}} \sum_{i} \left\| \mathbf{v}_{i} \times \left(\vec{\mathbf{c}}_{i} - \left(\mathbf{R} \vec{\mathbf{p}}_{i} + \vec{\mathbf{t}} \right) \right) \right\|^{2}$$

 $= \min_{\substack{\mathbf{R}, \mathbf{t} \\ \text{Step 4: Iterate steps 1-3 until reach convergence}}} \left\| \mathbf{skew} \left(\vec{\mathbf{v}}_i \right) \bullet \left(\mathbf{c}_i - \left(\mathbf{R} \vec{\mathbf{p}}_i + \vec{\mathbf{t}} \right) \right) \right\|^2$

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

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Computational Note

Gueziec uses the Cayley parameterization for rotations:

$$\mathbf{R}(\vec{\mathbf{u}}) = (\mathbf{I} - \mathbf{skew}(\vec{\mathbf{u}}))(\mathbf{I} + \mathbf{skew}(\vec{\mathbf{u}}))^{-1}$$

This leads to the approximation

$$\mathbf{R}(\vec{\mathbf{u}}) \approx \mathbf{I} + \mathbf{skew}(2\vec{\mathbf{u}})$$

which is similar to our familiar $\mathbf{R}(\vec{\alpha}) \approx \mathbf{I} + \mathbf{skew}(\vec{\alpha})$.

He also uses the notation \mathbf{U} =skew($\vec{\mathbf{u}}$). So $\mathbf{R}(\vec{\mathbf{u}}) = (\mathbf{I} - \mathbf{U})(\mathbf{I} + \mathbf{U})^{-1}$

Similarly, we will see $V=skew(\vec{v})$, etc.

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A countour-based 2D-3D method ...

Gueziec et al., 1998

Gueziec compared three different methods for performing the minimization in Step 3:

- Levenberg Marquardt (LM) nonlinear minimization.
- Linearization and constrained minimization
- Use of a Robust M-Estimator

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

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Levenberg-Marquardt ...

(Following development in Gueziec et al., 1998)

Define
$$f_i(\vec{x}) = \|\mathbf{V}_i(\vec{\mathbf{c}}_i - \mathbf{R}(\vec{\mathbf{u}})\vec{\mathbf{p}}_i - \vec{\mathbf{t}})\|$$
 where $\vec{x}^t = [\vec{\mathbf{u}}^t, \vec{\mathbf{t}}^t], \mathbf{V}_i = skew(\vec{\mathbf{v}}_i)$

Our goal is to minimize

$$\varepsilon(\vec{x}) = \sum_{i} f_{i}(\vec{x})^{2} = \sum_{i} \left\| \mathbf{V}_{i} \left(\vec{\mathbf{c}}_{i} - \mathbf{R}(\vec{\mathbf{u}}) \vec{\mathbf{p}}_{i} - \vec{\mathbf{t}} \right) \right\|^{2}$$

We note that $\varepsilon(\vec{x})$ is nonlinear. Levenberg-Marquardt is a widely used optimization method for problems of this type. However, it requires us to evaluate the partial derivitives $\partial f_i / \partial x_j$. Gueziec worked these out symbolically for his problem

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Levenberg-Marquardt ...

(Following development in Gueziec et al., 1998)

Define
$$f_i(\vec{x}) = \left\| \mathbf{V}_i \left(\vec{\mathbf{c}}_i - \mathbf{R}(\vec{\mathbf{u}}) \vec{\mathbf{p}}_i - \vec{\mathbf{t}} \right) \right\|$$
 where $\vec{x}^t = [\vec{\mathbf{u}}^t, \vec{\mathbf{t}}^t], \mathbf{V}_i = skew(\vec{\mathbf{v}}_i)$

$$\mathbf{J} = \begin{bmatrix} \cdots & \frac{\partial f_i}{\partial \vec{\mathbf{x}}} & \cdots \end{bmatrix} = \begin{bmatrix} \cdots & \frac{\partial f_i}{\partial \vec{\mathbf{u}}} & \cdots \\ & \frac{\partial f_i}{\partial \vec{\mathbf{t}}} & \cdots \end{bmatrix}$$

$$\frac{\partial f_i}{\partial \vec{\mathbf{t}}} = \frac{\mathbf{V}_i^t \mathbf{V}_i (\mathbf{R} \vec{\mathbf{p}}_i - \mathbf{c} + \vec{\mathbf{t}})}{f_i}$$

$$\frac{\partial f_i}{\partial \vec{\mathbf{u}}} = \left(\frac{\partial \mathbf{R} \vec{\mathbf{p}}_i}{\partial \vec{\mathbf{u}}}\right)^t \frac{\mathbf{V}_i^t \mathbf{V}_i (\mathbf{R} \vec{\mathbf{p}}_i - \mathbf{c} + \vec{\mathbf{t}})}{f_i}$$



Details on this may be found in reference [45] of Gueziec's paper

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

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Levenberg-Marquardt ...

(Following development in Gueziec et al., 1998)

Step 1: Pick λ = a small number; pick initial guess for \vec{x}

Step 2: Evaluate $f_i(\vec{x})$ and **J** and solve the least squares problem

$$\begin{bmatrix} \vdots \\ (\mathbf{J}^{t}\mathbf{J} + \lambda \mathbf{I})\Delta \vec{\mathbf{x}} - \mathbf{J}^{t}f_{i} \\ \vdots \end{bmatrix} = \begin{bmatrix} \vdots \\ 0 \\ \vdots \end{bmatrix}$$

for $\Delta \vec{x}$.

Step 3: $\vec{x} \leftarrow \vec{x} + \Delta \vec{x}$; update λ .

Step 4: Evaluate termination condition. If not done, go back to to step 2

Note: Usually λ starts small and grows larger. Consult standard references (e.g., Numerical Recipes) for more information.

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Constrained Linearized Least Squares ...

(Following development in Gueziec et al., 1998)

Step 0: Make an initial guess for \mathbf{R} and \mathbf{t}

Step 1: Compute $\vec{\mathbf{p}}_i \leftarrow \mathbf{R}\vec{\mathbf{p}}_i + \vec{\mathbf{t}}$

Step 2: Define $\mathbf{P}_i = skew(\vec{\mathbf{p}}_i)$, $\mathbf{V}_i = skew(\vec{\mathbf{v}}_i)$

Step 3: Solve the least squares problem:

$$\varepsilon^{2} = \min \begin{bmatrix} \vdots & \vdots \\ 2\mathbf{V}_{i}^{\mathbf{P}_{i}} & \mathbf{V}_{i} \\ \vdots & \vdots \end{bmatrix} \begin{bmatrix} \vec{\mathbf{u}} \\ \Delta \vec{\mathbf{t}} \end{bmatrix} - \begin{bmatrix} \vdots \\ \mathbf{V}_{i}(\vec{\mathbf{c}}_{i} - \vec{\mathbf{p}}_{i}) \end{bmatrix}^{2} \text{ subject to } \|\vec{\mathbf{u}}\| \leq \rho$$

where ρ is sufficiently small so that I+2U approximates a rotation

Step 4: Compute $\Delta \mathbf{R} = (\mathbf{I} - \mathbf{U})(\mathbf{I} + \mathbf{U})^{-1}$ Update $\mathbf{p}_i \leftarrow \Delta \mathbf{R} \mathbf{p}_i + \Delta \vec{\mathbf{t}}; \ \mathbf{R} \leftarrow \Delta \mathbf{R} \mathbf{R}; \ \vec{\mathbf{t}} \leftarrow \Delta \mathbf{R} \vec{\mathbf{t}} + \Delta \vec{\mathbf{t}}$

Step 5: If ε is small enough or some othe termination condition is met, then stop. Otherwise go back to Step 2.

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

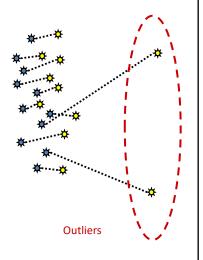
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Robust Pose Estimation ...

 Basic idea is to identify outliers and give them little or no weight.



R. Kumar and A. R. Hanson, "Robust methods for estimating pose and a sensitivity analysis," Comput. Vision, Graphics, Image Processing-IU, vol. 60, no. 3, pp. 313–342, 1994.

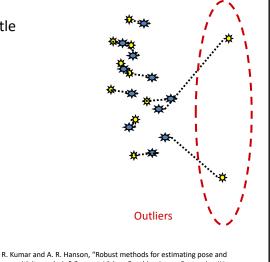
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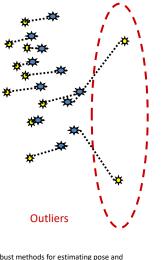
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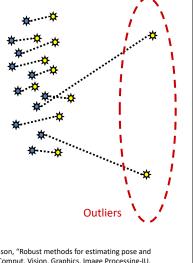
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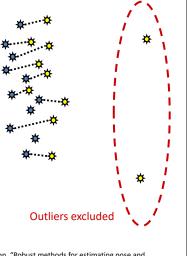
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Robust Pose Estimation ...

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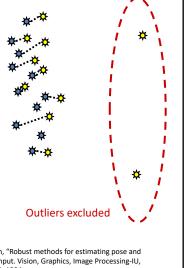
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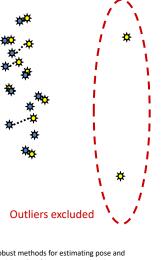
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Robust Pose Estimation ...

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Robust M-Estimator ...

(Following development in Gueziec et al., 1998)

Step 0: Make an initial guess for \mathbf{R} and \mathbf{t}

Step 1: Compute $\vec{\mathbf{p}}_i \leftarrow \mathbf{R}\vec{\mathbf{p}}_i + \vec{\mathbf{t}}$

Step 2: Define $\mathbf{P}_i = skew(\vec{\mathbf{p}}_i)$, $\mathbf{V}_i = skew(\vec{\mathbf{v}}_i)$,

Step 3: Solve a robust linearized problem

$$\varepsilon = \underset{\vec{\mathbf{u}}, \Delta \mathbf{t}}{\operatorname{argmin}} \sum_{i} \rho \left(\frac{0.6745 \ \mathbf{e}_{i}}{median(\{\mathbf{e}_{i}\})} \right) \text{ where } \mathbf{e}_{i} = \left\| \mathbf{V}_{i}(\vec{\mathbf{p}}_{i} - \mathbf{c}_{i} + 2\mathbf{P}_{i}\vec{\mathbf{u}} + \Delta \vec{\mathbf{t}}) \right\|$$

(See next slide)

Step 4: Compute $\Delta \mathbf{R} = (\mathbf{I} - \mathbf{U})(\mathbf{I} + \mathbf{U})^{-1}$

Update $\mathbf{p}_i \leftarrow \Delta \mathbf{R} \mathbf{p}_i + \Delta \vec{\mathbf{t}}$; $\mathbf{R} \leftarrow \Delta \mathbf{R} \mathbf{R}$; $\vec{\mathbf{t}} \leftarrow \Delta \mathbf{R} \vec{\mathbf{t}} + \Delta \vec{\mathbf{t}}$

Step 5: If ε is small enough or some othe termination condition is met, then stop. Otherwise go back to Step 2.

A. Guéziec, P. Kazanzides, B. Williamson, and R. Taylor, "Anatomy-Based Registration of CT-Scan and Intraoperative X-Ray Images for Guiding a Surgical Robot," IEEE Transactions on Medical Imaging, vol. 17, pp. 715-728, 1998.

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Robust M-Estimator ...

(Following development in Gueziec et al., 1998)

Step 3.0: Set $\vec{\mathbf{u}} = \vec{\mathbf{0}}$, $\Delta \mathbf{t} = \vec{\mathbf{0}}$

Step 3.1: Compute $\mathbf{e}_i = \left| \left| \mathbf{V}_i(\vec{\mathbf{p}}_i - \vec{\mathbf{c}}_i + 2P_i\vec{\mathbf{u}} + \Delta \vec{\mathbf{t}}) \right| \right|$, $s = median(\left\{ \cdots, \mathbf{e}_i, \cdots \right\}) / 0.6745$,

Step 3.2: Solve $\mathbf{C}\vec{\mathbf{x}} = \vec{\mathbf{d}}$, where $\vec{\mathbf{x}}^t = [\vec{\mathbf{u}}^t, \vec{\mathbf{t}}^t]$

$$\mathbf{C} = \sum_{i} \Psi(\frac{\mathbf{e}_{i}}{\mathbf{s}}) \begin{bmatrix} 2\mathbf{P}_{i}\mathbf{W}_{i}\mathbf{P}_{i} & \mathbf{P}_{i}\mathbf{W}_{i} \\ 2\mathbf{P}_{i}\mathbf{W}_{i} & \mathbf{W}_{i} \end{bmatrix} \text{ and } \vec{\mathbf{d}} = \sum_{i} \Psi(\frac{\mathbf{e}_{i}}{\mathbf{s}}) \begin{bmatrix} \mathbf{P}_{i}\mathbf{W}_{i}(\vec{\mathbf{c}}_{i} - \vec{\mathbf{p}}_{i}) \\ \mathbf{W}_{i}(\vec{\mathbf{c}}_{i} - \vec{\mathbf{p}}_{i}) \end{bmatrix}$$

where
$$\mathbf{W}_{i} = \mathbf{V}_{i}^{t} \mathbf{V}_{i} = \mathbf{I} - \vec{\mathbf{v}}_{i} \mathbf{v}_{i}^{t}$$
 $\Psi(\mu) = \begin{cases} \mu \left(1 - \mu^{2} / \alpha^{2}\right)^{2} & \text{if } \|\mu\| \leq \alpha \\ 0 & \text{otherwise} \end{cases}$

(**Note**: We use α =2)

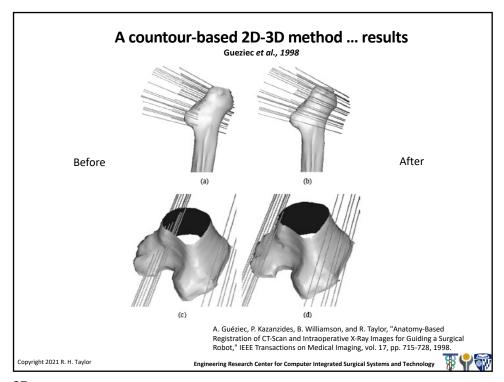
Step 3.3: Iterate steps 3.1 and 3.2 until a suitable termination condition is reached.

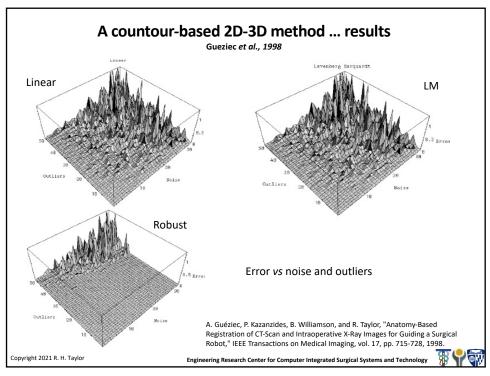
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A contour-based 2D-3D method ... times

Gueziec et al., 1998

TABLE I

AVERAGE EXECUTION TIMES IN MS FOR THE THREE REGISTRATION METHODS APPLIED TO DATA SETS THAT COMPRISE 100 POINTS (TOP) AND 20 POINTS (BOTTOM)

Number Points/Method	LM	Linear	Robust
100 points (CPU time)	790	690	28
20 points (CPU time)	200	42	9.6

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Sample Set Analysis

- **Question:** How good is a particular set of 3D sample points for the purpose of registration to a 3D surface?
- Long line of authors have looked at this question
- Next few slides are based on the work of David Simon, et al (1995)

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Sample Set Analysis: Distance Estimates

Let

$$F(\mathbf{x}) = 0$$

be the implicit equation of a surface, then one good estimate of the distance of a point \mathbf{x} to the surface is

$$D(\mathbf{x}) = \frac{F(\mathbf{x})}{\|\nabla F(\mathbf{x})\|}$$

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Sample set analysis: sensitivity

Let \mathbf{x}_s be a point on the surface, and let $T(\overline{\eta})$ represent a small perturbation with parameters $\overline{\eta}$ with respect to the surface of point \mathbf{x}_s :

$$\mathbf{x}_s' = T(\overline{\eta})\mathbf{x}_s$$

Then we define $\mathbf{V}(\mathbf{x}_s)$ to be

$$\mathbf{V}(\mathbf{x}_s) = \frac{\partial D(T(\overline{\eta})\mathbf{x}_s)}{\partial \overline{\eta}} = \begin{bmatrix} \mathbf{n}_s \\ \mathbf{x}_s \times \mathbf{n}_s \end{bmatrix}$$

where \mathbf{n}_s is the unit normal to the surface at \mathbf{x}_s . So,

$$D(\mathbf{T}(\overline{\eta})\mathbf{x}_s)) \simeq \mathbf{V}^T(\mathbf{x}_s)\overline{\eta}$$

Squaring this gives

$$D^{2}(\mathbf{T}(\overline{\eta})\mathbf{x}_{s})) \simeq \overline{\eta}_{T}\mathbf{V}(\mathbf{x}_{s})\mathbf{V}^{T}(\mathbf{x}_{s})\overline{\eta}$$
$$= \overline{\eta}^{T}\mathbf{M}(\mathbf{x}_{s})\overline{\eta}$$

Note that ${\bf M}$ is 6×6 positive, semi-definite, symmetric matrix.

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Sample set analysis: sensitivity

For a region \mathcal{R} , define

$$E_{R}(\overline{\eta}) = \overline{\eta}^{T} \left[\sum_{\mathbf{x}_{s} \in \mathcal{R}} \mathbf{M}(\mathbf{x}_{s}) \right] \overline{\eta}$$

$$= \overline{\eta}^{T} \mathbf{\Psi}_{\mathcal{R}} \overline{\eta}$$

$$= \overline{\eta}^{T} \mathbf{Q} \mathbf{\Lambda} \mathbf{Q}^{T} \overline{\eta}$$

$$= \sum_{1 \leq i \leq 6} \lambda_{i} (\overline{\eta}^{T} \cdot \mathbf{q}_{i})^{2}$$

• Note that the eigenvectors \mathbf{q}_i correspond to small differential transformations $\mathbf{T}(\mathbf{q}_i)$, and can sort eigenvalues so that

$$\lambda_1 \ge \lambda_2 \ge \ldots \ge \lambda_6$$

- Note that eigenvector \mathbf{q}_1 corresponds to direction of greatest constraint.
- Similarly, can also think of \mathbf{q}_6 as the least constrained direction.

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Sample Set Analysis: Goodness Measures

- Magnitude of smallest eigenvalue (Simon)
- (Kim and Khosla)

$$\frac{\sqrt[6]{\lambda_1 \cdot \ldots \cdot \lambda_6}}{\lambda_1 + \ldots + \lambda_6}$$

• Nahvi

$$\frac{\lambda_6^2}{\lambda_1}$$

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Sample Set Selection

- One blind search method (similar to Simon, 1995) is:
 - Randomly select sample points on surface
 - (prune for reachability)
 - evaluate goodness of sample set using some criterion
 - repeat many times and choose the best one found

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Sample Set Selection

- · Refinement of blind search (hill climbing):
 - Randomly select sample points on surface
 - (prune for reachability)
 - evaluate goodness of sample set using some criterion
 - replace a point from sample set with a randomly selected point
 - evaluate goodness
 - if better, keep it
 - else revert to original point and try again
- Variations include simulated annealing, "genetic" algorithms

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Sample Set Selection: Another Alternative

- ullet Select large number of random points ${f x}_s$
- Prune for reachability
- For each point, compute constraint direction $\mathbf{V}_s = \mathbf{V}(\mathbf{x}_s)$. To a first approximation, a measurement at \mathbf{x}_s with accuracy ϵ_s constrains $\overline{\eta}$ by

$$|\mathbf{V}_s\overline{\eta}|\leq\epsilon_s$$

• Now select subset of the \mathbf{x}_s that minimizes, e.g.,

$$\min_{\delta_s} \max \overline{\eta}^T \mathbf{S} \overline{\eta}$$

subject to

$$\begin{array}{rcl} \{\delta_s \ \in \ \{0,1\} \\ |\delta_s \mathbf{V}_s \overline{\eta}| \ \leq \ \epsilon_s \\ \sum\limits_s \delta_s \ \leq \ \text{subsetsize} \end{array}$$

There are various ways to do this.

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Sample Set Selection: Another Alternative (con'd)

• One can also minimize other forms, e.g.,

$$\min_{s} \max_{i} |\sigma_{i}\eta_{i}|$$

subject to similar constraints

• An alternative is to minimize the number of sample points required to ensure that some constraints on $\overline{\eta}$ are guaranteed to be met. E.g.,

$$\min_{\delta_s} \sum \delta_s$$

such that

$$\delta_s \in \{0, 1\}$$
$$\xi \leq \xi_{limit}$$

where

$$\xi = \max_{\overline{\eta}} \overline{\eta}^T \mathbf{S} \overline{\eta}$$

or some other form subject to

$$|\delta_s \mathbf{V}_s \overline{\eta}| \leq \epsilon_s$$

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Probabilistic Registration

- Registration methods typically use some optimization algorithm to find a "best" transformation between one data set and the other.
- It makes sense to try to find the "most likely" registration transformation.
- ICP minimizes sum-of-squares distances.
- This is equivalent to assuming that point-pair match probabilities are independent and symmetric Gaussian distributions based on distances
- But there are a number of other methods that explicitly consider probabilities ...

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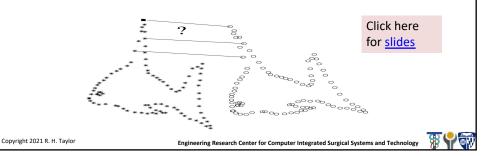
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Coherent Point Drift

- A. Myronenko and X. Song, "Point-Set Registration: Coherent Point Drift", *IEEE Trans. on Pattern Analysis and Machine Intelligence*, vol. 32- 12, pp. 2262-2275, 2010.
- · Alignment of point clouds
 - Fast method follows "EM" paradigm
 - Tolerates outliers and noise
 - Transformations: Rigid, affine, general deformable



Quiz

- The link will be in the chat
- You have 2 minutes

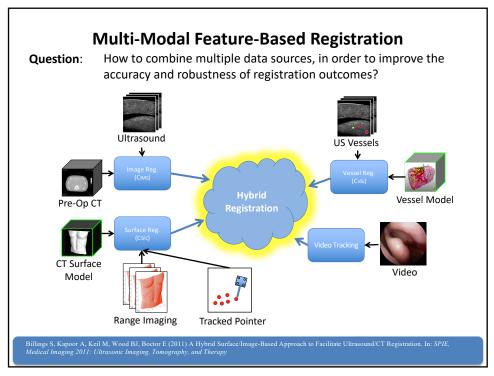


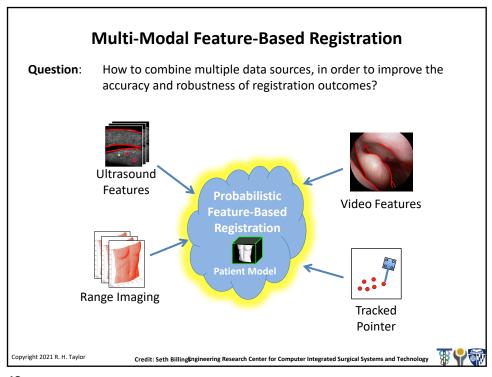
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Iterative Closest Point (ICP) Revisited



- Widely popular and useful method for point cloud to surface registration introduced by Besl & McKay in 1992
- Many variants proposed since its inception affecting all aspects of the algorithm (robustness, matching criteria, match alignment, etc.)

Matching Phase:

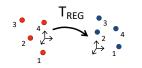
for each point in the source shape, find the closest point on the target shape

$$y_i = C_{CP}(T(x_i), \Psi) = \underset{y \in \Psi}{\operatorname{argmin}} \|y - T(x_i)\|_2$$

> Registration Phase:

compute transformation to minimize sum of square distances between matches

$$T = \underset{T}{\operatorname{argmin}} \sum_{i=1}^{n} \|\boldsymbol{y_i} - T(\boldsymbol{x_i})\|_2^2$$



S. Billings and R. H. Taylor, "Iterative Most Likely Oriented Point Registration", in Medical Image Computing and Computer-Assisted Interventions (MICCAI), Boston, October, 2014.

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Most-Likely Point Paradigm Illustrated with ICP

1. Probability Model: isotropic Gaussian

$$f_{\text{match}}(\mathbf{x} \mid \mathbf{y}, \sigma^2) = \frac{1}{(2\pi\sigma^2)^{3/2}} \cdot e^{-\frac{1}{2\sigma^2} \|\mathbf{y} - \mathbf{x}\|^2}$$

2. Match Phase:

$$\begin{aligned} \mathbf{y}_i &= \underset{\mathbf{y}_i \in \boldsymbol{\Psi}}{\operatorname{argmax}} f_{\text{match}}(\mathbf{T}(\mathbf{x}_i) \, | \, \mathbf{y}_i, \sigma^2) \\ &= \underset{\mathbf{y}_i \in \boldsymbol{\Psi}}{\operatorname{argmax}} \, \frac{1}{(2\pi\sigma^2)^{3/2}} \cdot e^{-\frac{1}{2\sigma^2} \|\mathbf{y}_i - \mathbf{T}(\mathbf{x}_i)\|^2} \\ &\to \underset{\mathbf{y}_i \in \boldsymbol{\Psi}}{\operatorname{argmin}} \, \|\mathbf{y}_i - \mathbf{T}(\mathbf{x}_i)\| \end{aligned}$$

3. Registration Phase:

$$\begin{split} \mathbf{T} &= \underset{\mathbf{T}}{\operatorname{argmax}} \prod_{i}^{n} f_{\operatorname{match}}(\mathbf{T}(\mathbf{x}_{i}) \, | \, \mathbf{y}_{i}, \sigma^{2}) \\ &= \underset{\mathbf{T}}{\operatorname{argmax}} \prod_{i}^{n} \frac{1}{(2\pi\sigma^{2})^{3/2}} \cdot e^{-\frac{1}{2\sigma^{2}} \|\mathbf{y}_{i} - \mathbf{T}(\mathbf{x}_{i})\|^{2}} \\ &\rightarrow \underset{\mathbf{T}}{\operatorname{argmax}} \left[-n \log \left((2\pi\sigma^{2})^{3/2} \right) - \frac{1}{2\sigma^{2}} \sum_{i}^{n} \|\mathbf{y}_{i} - \mathbf{T}(\mathbf{x}_{i})\|^{2} \right] \\ &\rightarrow \underset{\mathbf{T}}{\operatorname{argmin}} \sum_{i}^{n} \|\mathbf{y}_{i} - \mathbf{T}(\mathbf{x}_{i})\|^{2} \end{split}$$

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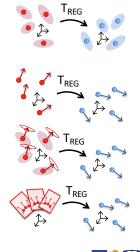
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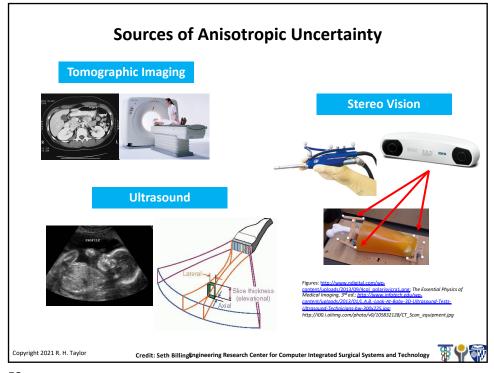
Outline of Registration Algorithms

- ICP Iterative Closest Point
 - isotropic position data
- IMLP Iterative Most Likely Point
 - anisotropic position data
 - robust to outliers
- IMLOP Iterative Most Likely Oriented Point
 - isotropic position & orientation data
- G-IMLOP Generalized IMLOP
 - anisotropic position & orientation data
- P-IMLOP Projected IMLOP
 - anisotropic position & projected orientation data



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Prior Work: Anisotropic Registration

Generalized Total Least Squares ICP (GTLS-ICP)

Estépar RSJ, Brun A, Westin C-F (2004) Robust generalized total least squares iterative closest point registration. In: MICCAI 2004

- Registration Phase
 - · anisotropic noise model
 - ad-hoc implementation less accurate / efficient; can be unstable
- Match Phase
 - isotropic (i.e. closest-point matching)
- Generalized ICP (G-ICP)
 - Registration Phase
 - anisotropic noise model limited to model locally-linear surface regions surrounding each feature point of a point cloud shape
 - · uses off-the-shelf conjugate gradient solver
 - Match Phase
 - isotropic (i.e. closest-point matching)

Segal A, Haehnel D, Thrun S (2009) Generalized-ICP. In: Robotics: Science and Systems V

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Prior Work: Anisotropic Registration

Anisotropic ICP (A-ICP)

Maier-Hein L, Franz AM, Dos Santos TR, Schmidt M, Fangerau M, et al. (2012) Convergent iterative closest-point algorithm to accomodate anisotropic and inhomogenous localization error. *IEEE Trans Pattern Anal Mach Intell* 34: 1520–1532.

- Registration Phase
 - · anisotropic noise model
 - ad-hoc implementation does not fully account for noise in both shapes (i.e., lacks ability to reorient the data-shape covariances during optimization)
- Match Phase
 - anisotropic noise model with non-optimal matching (finds minimal Mahalanobis distance match rather than most-likely match)
 - inefficient implementation; also cannot guarantee that the "best" match is found
- Initializes registration by ICP (due to inefficient match phase)

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Iterative Most Likely Point (IMLP)

Probability Model: anisotropic Gaussian

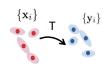
$$f_{\text{match}}(\mathbf{x} \mid \mathbf{y}, \Sigma_{\mathbf{x}}, \Sigma_{\mathbf{y}}) = \frac{1}{(2\pi)^{3/2} |\Sigma_{\mathbf{x}} + \Sigma_{\mathbf{y}}|^{1/2}} \cdot e^{-\frac{1}{2}(\mathbf{y} - \mathbf{x})^T (\Sigma_{\mathbf{x}} + \Sigma_{\mathbf{y}})^{-1} (\mathbf{y} - \mathbf{x})}$$

Match Phase:

$$egin{aligned} \left[\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}
ight] &= \operatorname*{argmin}_{\left[\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}
ight] \in \Psi} \left[\left.\log \left\langle \mathbf{R}\mathbf{\Sigma}_{\mathrm{x}i}\mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i}
ight|
ight) \\ &+ \left.\left(\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i)
ight)^{\mathrm{\scriptscriptstyle T}} (\mathbf{R}\mathbf{\Sigma}_{\mathrm{x}i}\mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i})^{-1} (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i))
ight] \end{aligned}$$

Registration Phase:

$$\mathbf{T} = \underset{\mathbf{T} = [\mathbf{R}, \mathbf{t}]}{\operatorname{argmin}} \sum_{i}^{n} (\mathbf{y}_{i} - \mathbf{T}(\mathbf{x}_{i}))^{T} (\mathbf{R} \mathbf{\Sigma}_{xi} \mathbf{R}^{T} + \mathbf{\Sigma}_{yi})^{-1} (\mathbf{y}_{i} - \mathbf{T}(\mathbf{x}_{i}))$$



Billings SD, Boctor EM, Taylor RH (2015) Iterative Most-Likely Point Registration (IMLP): A Robust Algorithm for Computing Optimal Shape Alignment. *PLoS One* 10: e0117688

IMLP: Match Phase



- Due to anisotropic distance metric, standard KD-tree search techniques do not apply.
- Approach: PD-tree search with modified node test



PD Tree Constructed by Datum Positions

Constructing the PD tree:

- 1. Add all datums to a root node
- 2. Compute covariance of datum positions within the node
- 3. Create minimally-sized bounding box aligned to the covariance eigenvectors
- 4. Partition node along the direction of greatest extent
- 5. Form left and right child nodes from the datums in each partition
- 6. Repeat from Step 2 for left and right child nodes until # datums in node < threshold or node size < threshold

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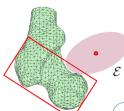
IMLP: Match Phase



Searching the PD tree:

Assume the current match candidate has a match error equal to $E_{\text{best}} \,$

Question: can any feature in this node possibly provide a match error less than E_{best}?



$$\begin{split} [\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}] &= \operatorname*{argmin}_{[\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}] \in \boldsymbol{\Psi}} \left[\left. \log(\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i} \right|) \right. \\ &+ \left. (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i))^{\mathrm{\scriptscriptstyle T}} (\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i})^{-1} (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i)) \right] \end{split}$$

True if: $(\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i))^T (\mathbf{R} \mathbf{\Sigma}_{xi} \mathbf{R}^T + \mathbf{\Sigma}_{\mathrm{node}})^{-1} (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i)) < E_{\mathrm{best}} - log_{\min}$

Node of the PD Tree

Node Test: if the ellipsoid

$$\mathcal{E} = \{ \mathbf{y} \mid (\mathbf{y} - \mathrm{T}(\mathbf{x}_i))^{\mathrm{\scriptscriptstyle T}} (\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{node}})^{-1} (\mathbf{y} - \mathrm{T}(\mathbf{x}_i)) \leq E_{\mathrm{best}} - log_{\min} \}$$

intersects the bounding box of the node, then search the node

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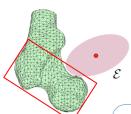
IMLP: Match Phase



Searching the PD tree:

Assume the current match candidate has a match error equal to Ebest

Question: can any feature in this node possibly provide a match error less than Ebest?



$$\begin{split} [\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}] &= \operatorname*{argmin}_{[\mathbf{y}_i, \mathbf{\Sigma}_{\mathrm{y}i}] \in \boldsymbol{\Psi}} \left[\left. \log(\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i} \right|) \right. \\ &+ \left. (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i))^{\mathrm{\scriptscriptstyle T}} (\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{y}i})^{-1} (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i)) \right] \end{split}$$

True if: $(\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i))^T (\mathbf{R} \mathbf{\Sigma}_{xi} \mathbf{R}^T + \mathbf{\Sigma}_{\mathrm{node}})^{-1} (\mathbf{y}_i - \mathrm{T}(\mathbf{x}_i)) < E_{\mathrm{best}} - log_{\min}$

Details in Billings' Thesis

Node Test: if the ellipsoid Node of the PD Tree

 $\mathcal{E} = \{\mathbf{y} \mid (\mathbf{y} - \mathrm{T}(\mathbf{x}_i))^{\mathrm{\scriptscriptstyle T}} (\mathbf{R} \mathbf{\Sigma}_{\mathrm{x}i} \mathbf{R}^{\mathrm{\scriptscriptstyle T}} + \mathbf{\Sigma}_{\mathrm{node}})^{-1} (\mathbf{y} - \mathrm{T}(\mathbf{x}_i)) \leq E_{\mathrm{best}} \vdash log_{\min} \}^{\mathrm{\scriptsize J}}$ intersects the bounding box of the node, then search the node

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IMLP: Registration Phase



1. Re-formulate the cost function from an unconstrained optimization

$$\mathbf{T} = \operatorname*{argmin}_{[\mathbf{R},\mathbf{t}]} \sum_{i=1}^{n} (\mathbf{y}_i - \mathbf{R}\mathbf{x}_i - \mathbf{t})^T (\mathbf{R}\boldsymbol{\Sigma}_{\mathrm{x}i}\mathbf{R}^T + \boldsymbol{\Sigma}_{\mathrm{y}i})^{-1} (\mathbf{y}_i - \mathbf{R}\mathbf{x}_i - \mathbf{t})$$

to a constrained optimization

constrained optimization
$$\mathbf{T} = \operatorname*{argmin}_{[\mathbf{R},\mathbf{t}]} \sum_{i=1}^n (\mathbf{x}_i - \mathbf{x}_i^*)^T \mathbf{\Sigma}_{xi}^{-1} (\mathbf{x}_i - \mathbf{x}_i^*) + \sum_{i=1}^n (\mathbf{y}_i - \mathbf{y}_i^*)^T \mathbf{\Sigma}_{yi}^{-1} (\mathbf{y}_i - \mathbf{y}_i^*)$$
 subject to:
$$\mathbf{F}_i(\mathbf{x}_i^*, \mathbf{y}_i^*, \mathbf{R}, \mathbf{t}) = \mathbf{y}_i^* - \mathbf{R} \mathbf{x}_i^* - \mathbf{t} = 0$$
 Generalized Total Least Squares (GTLS)

 x_i^* - true (unknown) data-point position yi* - true (unknown) model-point position

2. Linearize the constraints with a Taylor series centered at the measured (known) data

$$\begin{split} F_i(\mathbf{x}_i^*, \mathbf{y}_i^*, \mathbf{R}, \mathbf{t}) &\approx F_{Li}^k(\mathbf{x}_i, \mathbf{y}_i, \mathbf{d}\alpha, \mathbf{dt}) \\ &= F_i^0(\mathbf{x}_i, \mathbf{y}_i, \mathbf{R}_k, \mathbf{t}_k) - \mathbf{r}_{yi} + \mathbf{R}_k \mathbf{r}_{xi} + \mathrm{skew}(\mathbf{R}_k \mathbf{x}_i) \mathbf{d}\alpha - \mathbf{dt} = 0 \end{split}$$

Note using: $\Delta \mathbf{R} \approx \mathbf{I} + \text{skew}(\mathbf{d}\alpha)$ $\mathbf{r}_{xi} = \mathbf{x}_i - \mathbf{x}_i^*$ $\mathbf{r}_{vi} = \mathbf{y}_i - \mathbf{y}_i^*$

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IMLP: Registration Phase

- 3. Apply the method of Lagrange multipliers to solve constrained optimization.
 - 3a. Form the Lagrange function using the linearized constraints

$$\mathcal{L}(\mathbf{d}\alpha, \mathbf{dt}, \lambda) = \sum_{i=1}^{n} \mathbf{r}_{xi}^{T} \mathbf{\Sigma}_{xi}^{-1} \mathbf{r}_{xi} + \sum_{i=1}^{n} \mathbf{r}_{yi}^{T} \mathbf{\Sigma}_{yi}^{-1} \mathbf{r}_{yi} + \sum_{i=1}^{n} \lambda_{i}^{T} \mathbf{F}_{Li}^{k}(\mathbf{x}_{i}, \mathbf{y}_{i}, \mathbf{d}\alpha, \mathbf{dt})$$

3b. Solve zero gradient w.r.t. the optimization parameters and the Lagrange multipliers

$$\mathbf{dp} = \begin{bmatrix} \mathbf{d}\alpha \\ \mathbf{dt} \end{bmatrix} \quad \mathbf{f}^0 = \begin{bmatrix} \mathbf{f}_1^0 \\ \vdots \\ \mathbf{f}_n^0 \end{bmatrix} \quad \mathbf{J} = \begin{bmatrix} \mathrm{skew}(\mathbf{R}_k\mathbf{x}_1) & -\mathbf{I} \\ \vdots & \vdots \\ \mathrm{skew}(\mathbf{R}_k\mathbf{x}_n) & -\mathbf{I} \end{bmatrix} \quad \boldsymbol{\Sigma} = \begin{bmatrix} \mathbf{F}_x^0 \boldsymbol{\Sigma}_{\mathbf{x}} \mathbf{F}_x^{0T} + \boldsymbol{\Sigma}_{\mathbf{y}} \end{bmatrix}$$

$$\mathbf{F}_x^0 = \begin{bmatrix} -\mathbf{R}_k & & \\ & \ddots & \\ & & -\mathbf{R}_k \end{bmatrix} \quad \boldsymbol{\Sigma}_{\mathbf{x}} = \begin{bmatrix} \boldsymbol{\Sigma}_{\mathbf{x}1} & & \\ & \ddots & \\ & & \boldsymbol{\Sigma}_{\mathbf{x}n} \end{bmatrix} \quad \boldsymbol{\Sigma}_{\mathbf{y}} = \begin{bmatrix} \boldsymbol{\Sigma}_{\mathbf{y}1} & & \\ & \ddots & \\ & & \boldsymbol{\Sigma}_{\mathbf{y}n} \end{bmatrix}$$

4. Iteratively solve 3b by linear least squares until convergence.

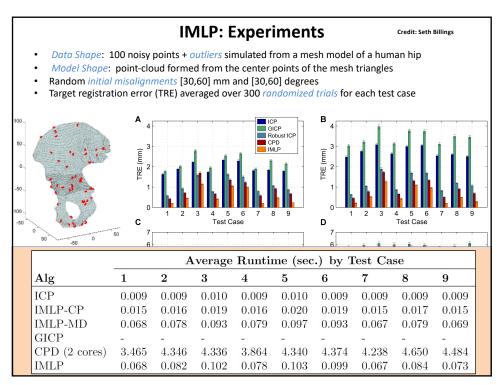
$$\mathbf{R}_{k+1} = \mathbf{R}(\mathbf{d}\alpha)\mathbf{R}_k$$
, $\mathbf{t}_{k+1} = \mathbf{t}_k + \mathbf{d}\mathbf{t}$

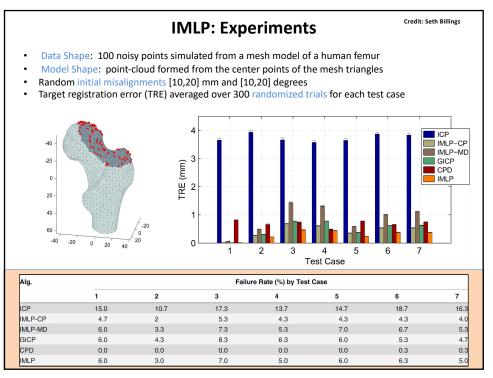
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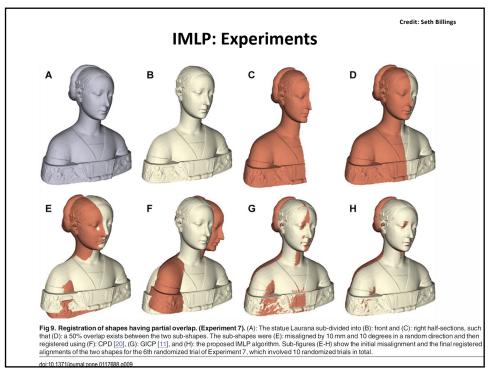
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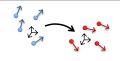
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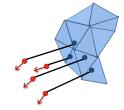
Iterative Most Likely Oriented Point (IMLOP)



> Matching Phase:

for each oriented point in the source shape, find the most likely oriented point on the target shape

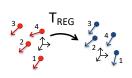
$$\boldsymbol{y_i} = \operatorname{C}_{\operatorname{MLP}}(T(\boldsymbol{x_i}), \boldsymbol{\varPsi}) = \operatorname*{argmax}_{\boldsymbol{y} \in \boldsymbol{\varPsi}} f_{\operatorname{match}}(T(\boldsymbol{x_i}), \boldsymbol{y})$$



> Registration Phase:

compute transformation to maximize the likelihood (i.e. minimize negative log-likelihood) of oriented point matches

$$T = \underset{T}{\operatorname{argmin}} \left(\frac{1}{2\sigma^2} \sum_{i=1}^{n} \| \boldsymbol{y}_{pi} - T(\boldsymbol{x}_{pi}) \|_2^2 - k \sum_{i=1}^{n} \boldsymbol{y}_{ni}^T R \boldsymbol{x}_{ni} \right)$$

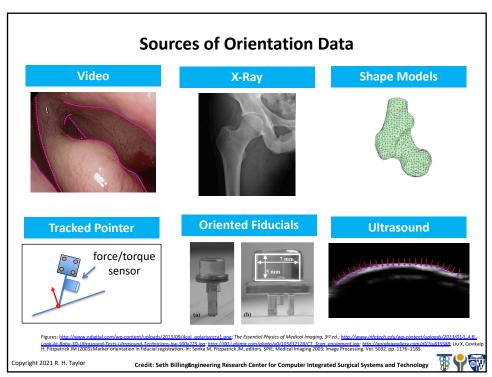


S. Billings and R. H. Taylor, "Iterative Most Likely Oriented Point Registration", in Medical Image Computing and Computer-Assisted Interventions (MICCAI), Boston, October, 2014.

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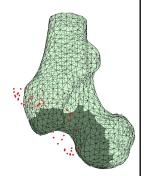




Experiments

Performance comparison of IMLOP vs. ICP was made through a simulation study using a human femur surface mesh segmented from CT imaging.

- source shape created by randomly sampling points from the mesh surface (10, 20, 35, 50, 75, and 100 points tested)
- Gaussian [wrapped Gaussian] noise added to the source points (0, 0.5, 1.0, and 2.0 mm [degrees] tested)
- Applied random misalignment of [10,20] mm / degrees
- 300 trials performed for each sample size / noise level
- Registration accuracy (TRE) evaluated using 100 validation points randomly sampled from the mesh
- Registration failures automatically detected using threshold on final residual match errors



Example source point cloud sampled from dark region of target mesh.

ICP: threshold on position residuals only

IMLOP: threshold on position & orientation residuals

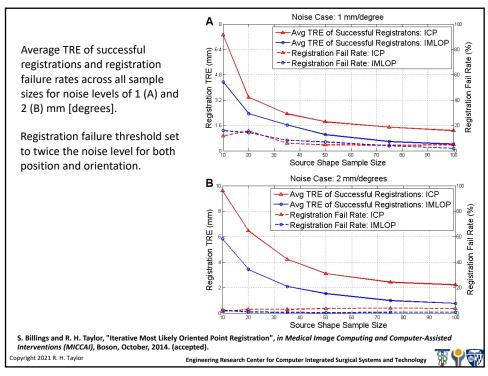
S. Billings and R. H. Taylor, "Iterative Most Likely Oriented Point Registration", in Medical Image Computing and Computer-Assisted Interventions (MICCAI), Boston, October, 2014.

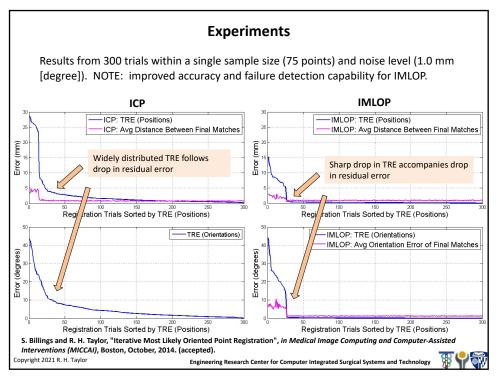
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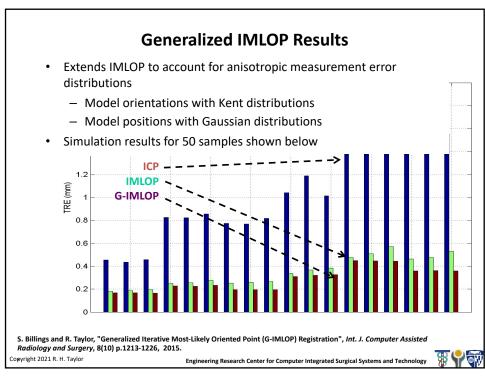
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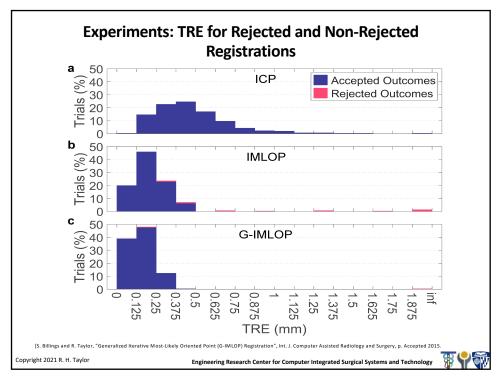
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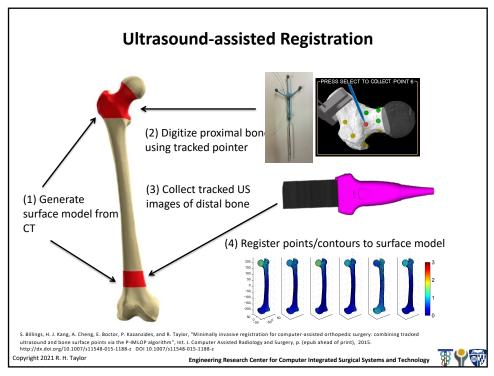
70

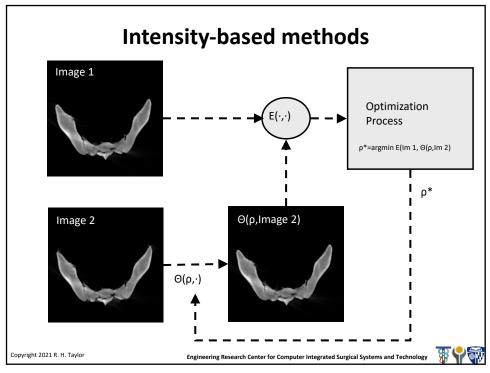












Intensity-based methods

- Typically performed between images
- The "features" in this case are the intensities associated with pixels (2D) or voxels (3D) in the images.
- · General framework:

$$\vec{\rho}^* = \min_{\vec{\rho}} E\left(Image_1, \Theta\left(\vec{\rho}, Image_2\right)\right)$$

• Methods differ mostly in choice of transformation function $\Theta(\cdot)$ and Energy function $E(\cdot,\cdot)$,

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Typical energy functions (not an exhaustive list)

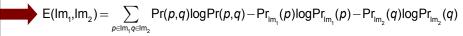
Normalized image subtraction

$$E(Im_{1},Im_{2}) = \sum_{\overline{k}} \frac{\left|Im_{1}\left[\overline{k}\right] - Im_{2}\left[\overline{k}\right]\right|}{\max_{\overline{j}} \left(\left|Im_{1}\left[\overline{j}\right] - Im_{2}\left[\overline{j}\right]\right|\right)}$$

Normalized cross correlation (NCC)

$$\mathsf{E}(\mathsf{Im}_{\scriptscriptstyle{1}},\mathsf{Im}_{\scriptscriptstyle{2}}) = \frac{\sum_{\overline{k}} \left(\mathsf{Im}_{\scriptscriptstyle{1}}\left[\overline{k}\right] - avg(\mathsf{Im}_{\scriptscriptstyle{1}})\right) \left(\mathsf{Im}_{\scriptscriptstyle{2}}\left[\overline{k}\right] - avg(\mathsf{Im}_{\scriptscriptstyle{2}})\right)}{\sqrt{\sum_{\overline{k}} \left(\mathsf{Im}_{\scriptscriptstyle{1}}\left[\overline{k}\right] - avg(\mathsf{Im}_{\scriptscriptstyle{1}})\right)^{2}}\sqrt{\sum_{\overline{k}} \left(\mathsf{Im}_{\scriptscriptstyle{2}}\left[\overline{k}\right] - avg(\mathsf{Im}_{\scriptscriptstyle{2}})\right)^{2}}}$$

Mutual information



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Mutual Information

- First proposed independently in 1995 by Collignon and Viola & Wells.
- · Very widely practiced
- Is able to co-register images with very different sensor modalities so long as there is a stable relationship between intensities in one modality with those in another
- · Many "flavors" and variations

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Mutual Information

Entropy

 $H(a) = Pr(a) \log Pr(a)$

 $H(a,b) = Pr(a,b) \log Pr(a,b)$

Mutual Information (Viola & Wells '95, Colligen '95)

Similarity (A,B) = H(A) + H(B) - H(A,B)

Normalized mutual information (Maes et al. '97)

Similarity
$$(A,B) = \frac{H(A) + H(B)}{H(A,B)}$$

Objective function

 $\mathsf{E}(\mathsf{Im}_{\scriptscriptstyle{1}},\!\mathsf{Im}_{\scriptscriptstyle{2}}) = -\mathit{Similarity}(\mathsf{Im}_{\scriptscriptstyle{1}},\!\mathsf{Im}_{\scriptscriptstyle{2}})$

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Basic Idea of Intensity-Based 2D/3D Registration

- Assumes a pre-op CT is available
- Simulate many C-Arm images and choose the most similar to the intraoperative image
- Solves the following optimization problem:

 $\underset{\theta \in SE(3)}{\operatorname{argmin}} \mathcal{S}(I_{\operatorname{Intra-Op}}, \mathcal{P}(\theta, I_{\operatorname{CT}}))$



Slide credit: Robert Grupp

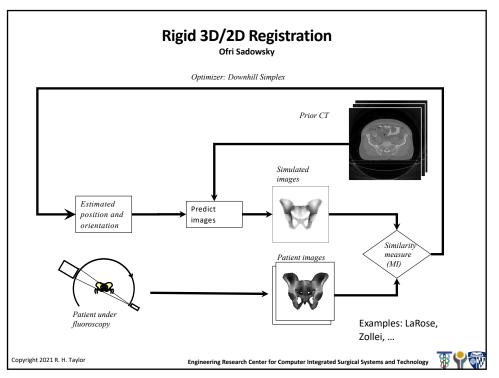
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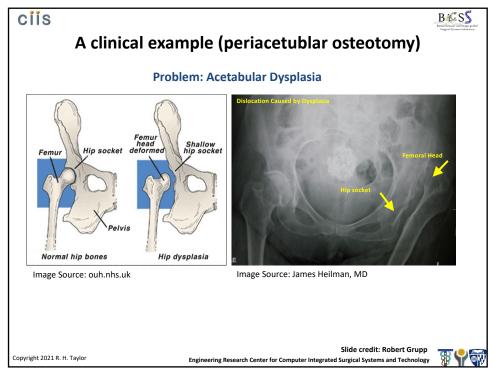


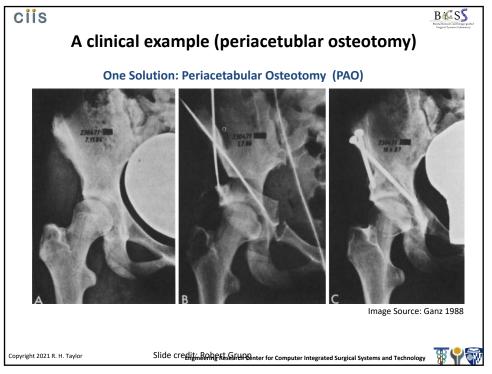


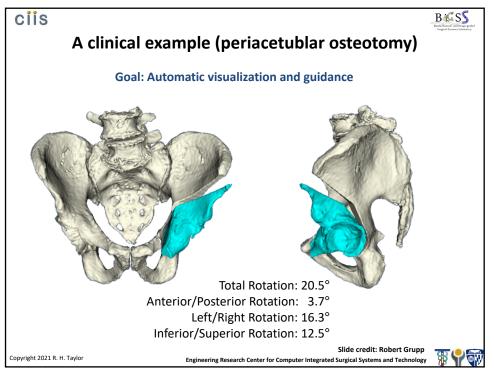
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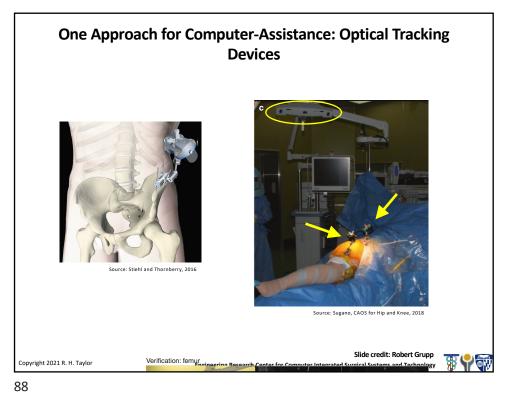




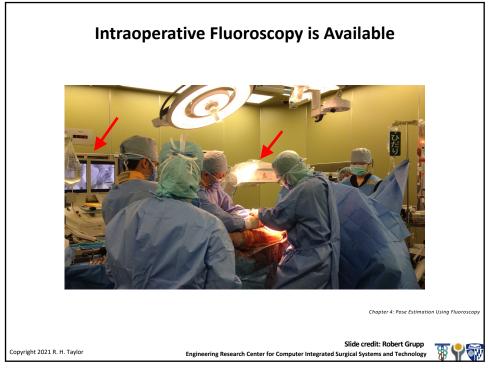


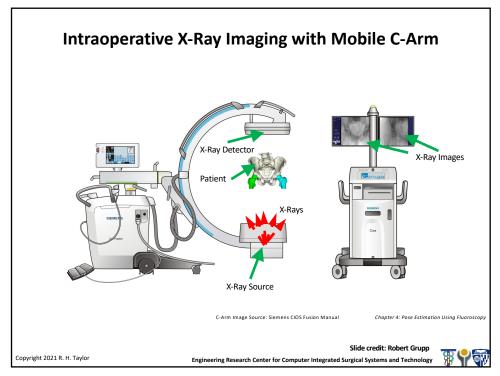


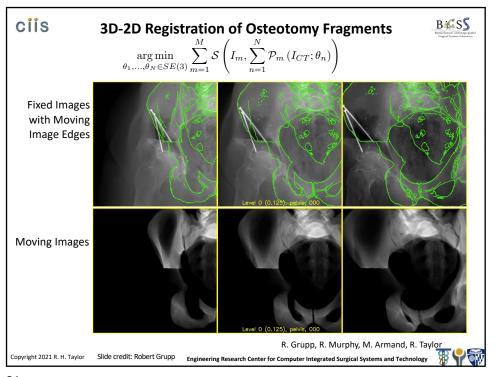




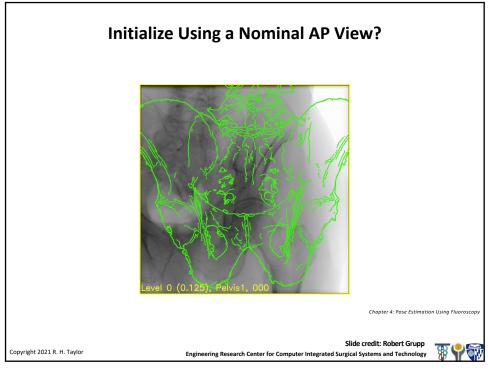






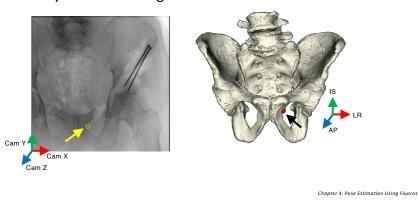


• Compute the Sobel derivatives in the X and Y directions of the two input images: $\nabla_X I_1, \ \nabla_X I_2, \ \nabla_Y I_1, \ \nabla_Y I_2$ • Compute NCC between the corresponding gradient images: $\mathcal{S}(I_1,I_2) = NCC(\nabla_X I_1, \nabla_X I_2) + NCC(\nabla_Y I_1, \nabla_Y I_2)$ $\nabla_X I_1 \qquad \nabla_X I_2 \qquad \nabla_Y I_1 \qquad \nabla_Y I_2$ R. Grupp, R. Murphy, M. Armand, R. Taylor Copyright 2021 R. H. Taylor Slide credit: Robert Grupp Engineering Research Center for Computer Integrated Surgical Systems and Technology



Use a Single Landmark to Initialize Registration

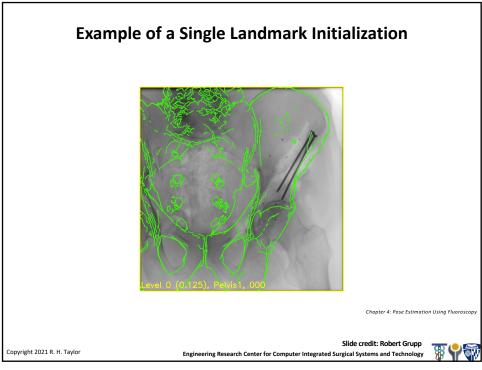
- Assume the pelvis is in an AP orientation this may be computed preoperatively
- Manually annotate a single landmark to recover translation



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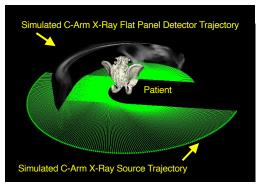




Automatically Initialize Second and Third Views

- Constrain C-arm motion to orbital rotation
- Perform an exhaustive search over $\pm 90^{\circ}$ in 1° increments



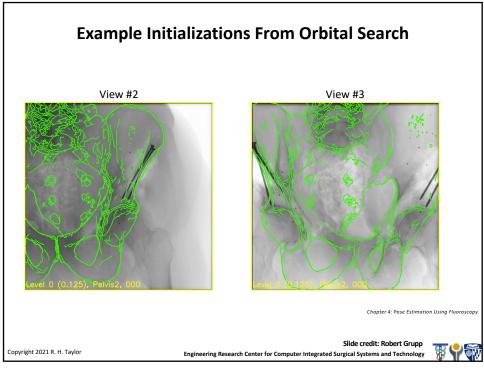


Chapter 4: Pose Estimation Using Fluoroscop

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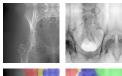
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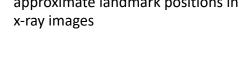


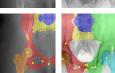


Automatic Landmark-Based Initialization

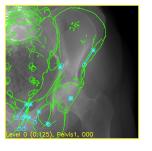
• Train a CNN to recognize approximate landmark positions in x-ray images







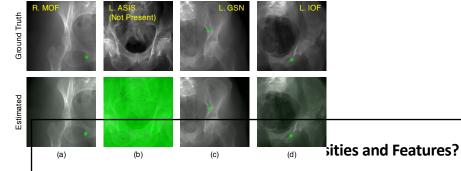
- Use landmark-based 2D-3D registration to initialize registration
- Combine landmark and intensity objective functions



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• Registration objective function:

$$\min_{\theta_{P},\theta_{LF},\theta_{RF} \in SE(3)} \underbrace{\lambda \mathcal{S}\left(\mathcal{P}\left(\theta_{P},\theta_{LF},\theta_{RF}\right),I\right) + \left(1-\lambda\right) \mathcal{R}\left(\theta_{P},\theta_{LF},\theta_{RF}\right)}_{\text{Image Similarity Term}} \\ + \left(1-\lambda\right) \mathcal{R}\left(\theta_{P},\theta_{LF},\theta_{RF}\right)$$

- Usually, regularization penalizes the amount of rotation and translation away from initialization
- Why not directly include the landmark re-projection as regularization?

$$\mathcal{R}\left(\theta_{P}\right) = \frac{1}{2\sigma_{\ell}^{2}} \sum_{l=1}^{N_{L}} \left\| \mathcal{P}\left(p_{\mathrm{3D}}^{(l)}; \theta_{P}\right) - p_{\mathrm{2D}}^{(l)} \right\|_{2}^{2}$$

 Can also think of this as running landmark registration and regularizing on image appearance

Chapter 6: Automatic and Rob

regularizing on image appearance

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