# Digital Geometry - Surface Registration

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http://jjcao.github.io/DigitalGeometry/

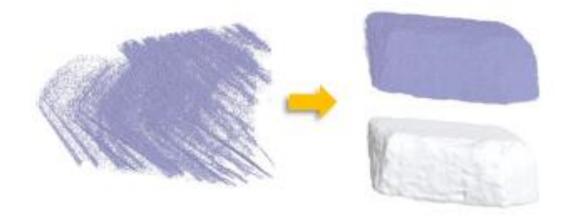
#### **Definition**

- Surface registration is the process of identifying and matching corresponding regions
- across multiple scans given in arbitrary initial positions,
- and estimating the corresponding rigid transforms that best align the scans to each other.



2018/2/28

## **Applications**



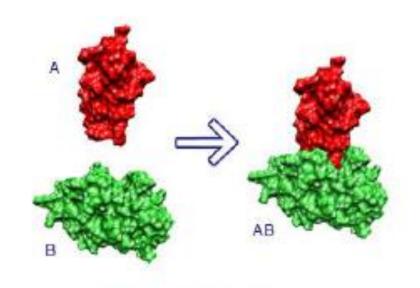
Surface reconstruction



Object completion



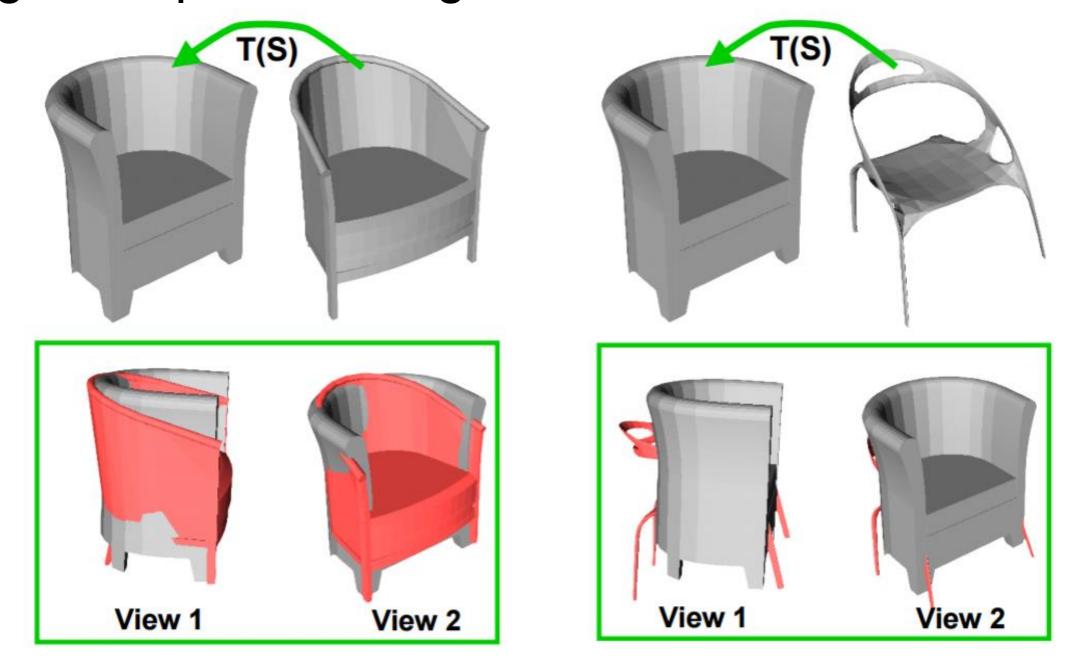
Fragment assembly



Protein docking

ZU 10/Z/Z0

## Rigid Shape Matching – Search a transformation



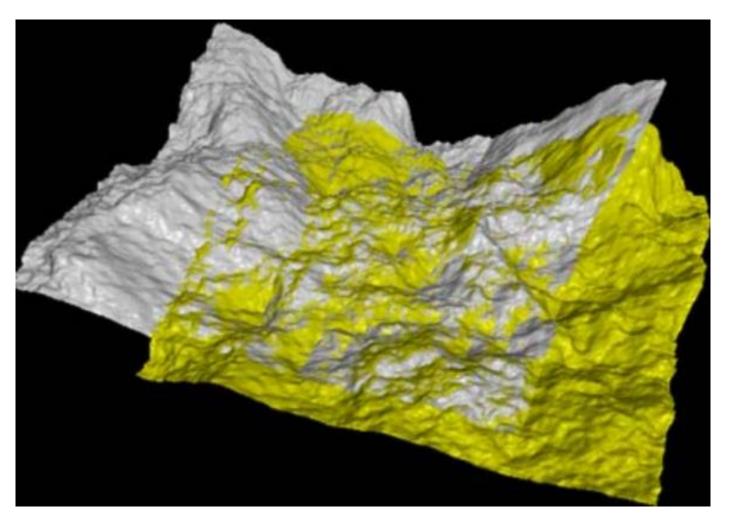
#### Correspondence Problem Classification

- How many meshes?
  - Two: Pairwise registration
  - More than two: multi-view registration
- Initial registration available?
  - Yes: Local optimization methods
  - No: Global methods
- Class of transformations?
  - Rotation and translation: Rigid-body
  - Non-rigid deformations

## Pairwise Rigid Registration Goal

Align two partially overlapping meshes given initial guess for relative

transform



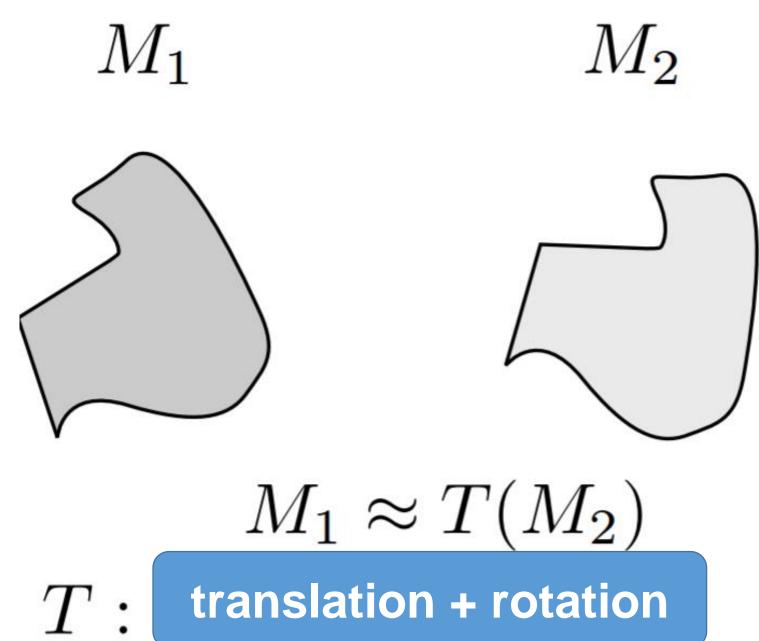
#### Outline

Basic ICP: Iterative Closest Points

- Classification of ICP variants
  - Faster alignment
  - Better robustness
- Global Registration

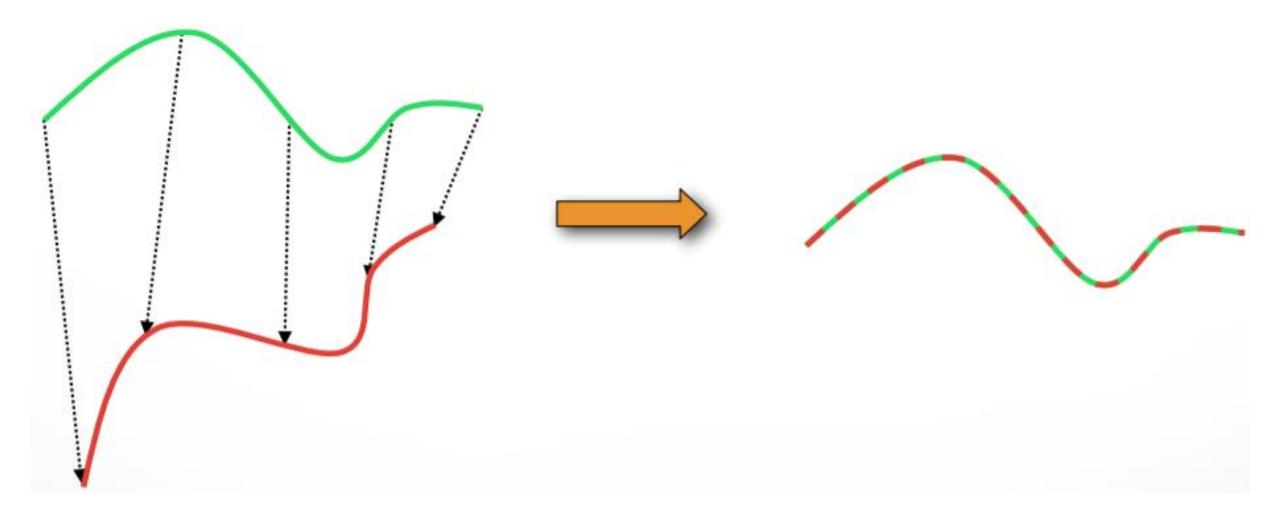
## ICP: Iterative Closest Points

## Objective



## Aligning 3D Data

• If correct **correspondences** are **known**, can find correct relative rotation/translation

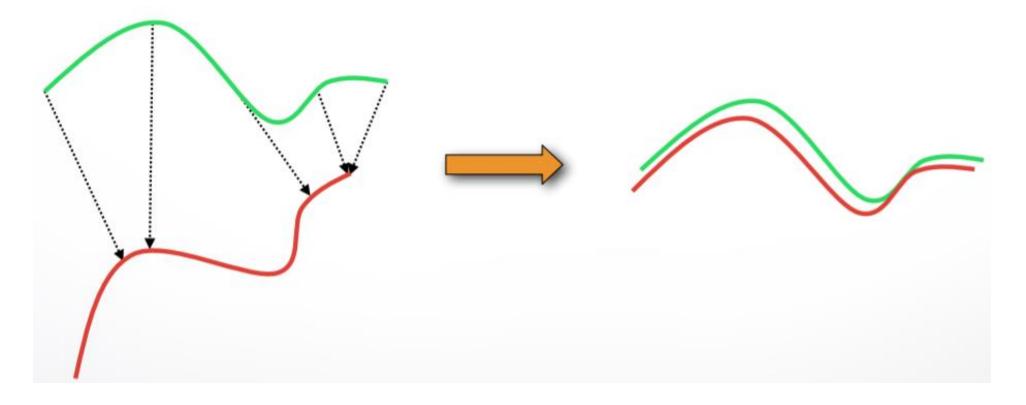


## Aligning 3D Data

How to find correspondences: User input? Feature detection?
 Signatures?

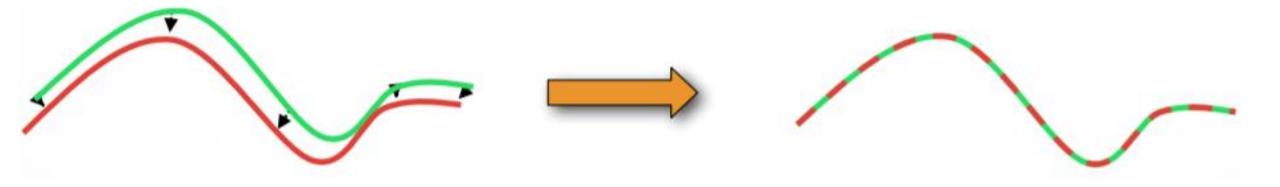
a seeminglyradical guess

Alternatives: assume closest points correspond



## Aligning 3D Data

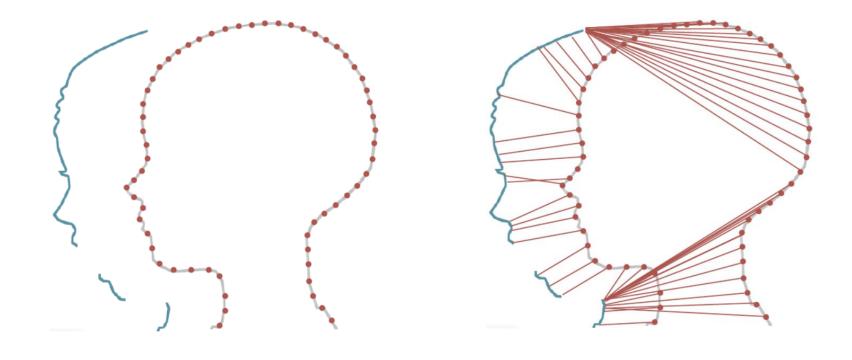
- ... and iterate to find alignment
  - Iterative Closest Points (ICP) [Besl & Mckay]
- Converges if starting position "close enough"



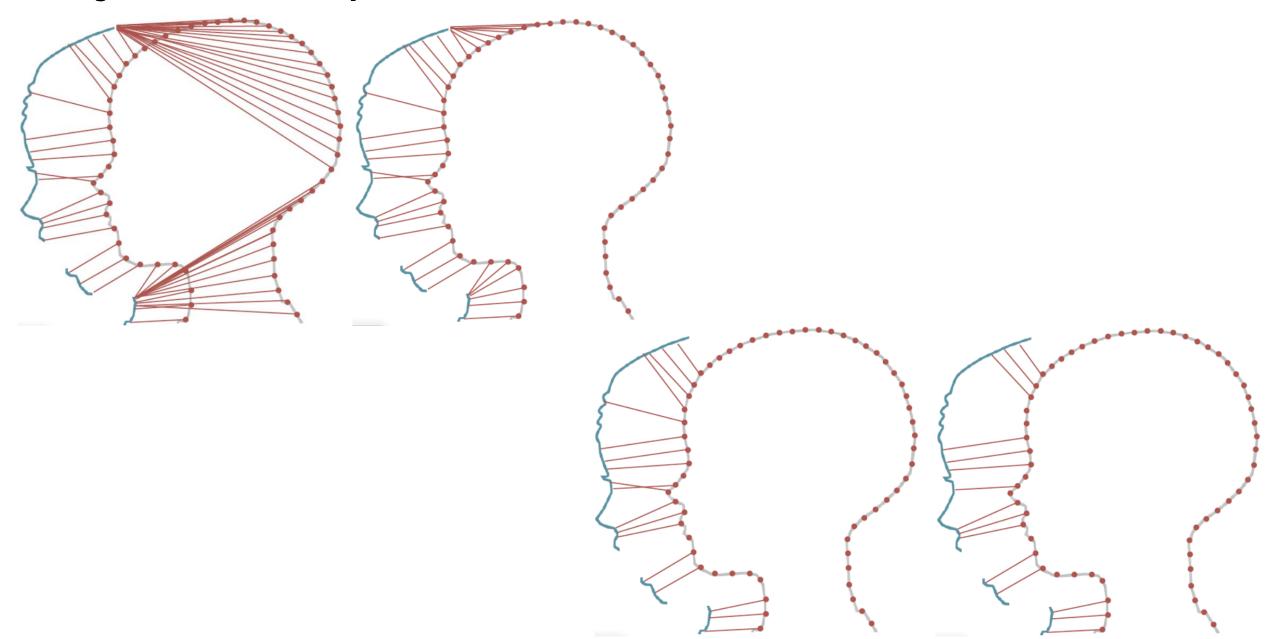
#### **Basic ICP**

• Select e.g., 1000 random points

• Match each to closest point on other scan



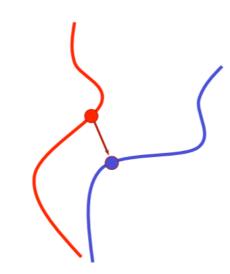
## Reject some pairs



#### **Basic ICP**

• Select e.g., 1000 random points

- Match each to closest point on other scan
- Reject pairs with distance > k times median



Construct error function:

$$E = \sum \|\mathbf{R}\mathbf{p}_i + \mathbf{t} - \mathbf{q}_i\|^2$$

 Minimize (closed form solution in [Horn 87]) (Also this note: Least-Squares Rigid Motion Using SVD by Olga Sorkine)

## Shape Matching: Translation first. why?

Assume R is fixed and denote  $F(\mathbf{t}) = \sum_{i=1}^{n} w_i ||(R\mathbf{p}_i + \mathbf{t}) - \mathbf{q}_i||^2$ . We can find the optimal translation by taking the derivative of F w.r.t.  $\mathbf{t}$  and searching for its roots:

$$0 = \frac{\partial F}{\partial \mathbf{t}} = \sum_{i=1}^{n} 2w_i \left( R\mathbf{p}_i + \mathbf{t} - \mathbf{q}_i \right) =$$

$$= 2\mathbf{t} \left( \sum_{i=1}^{n} w_i \right) + 2R \left( \sum_{i=1}^{n} w_i \mathbf{p}_i \right) - 2 \sum_{i=1}^{n} w_i \mathbf{q}_i.$$
(2)

Denote

$$\bar{\mathbf{p}} = \frac{\sum_{i=1}^{n} w_i \mathbf{p}_i}{\sum_{i=1}^{n} w_i}, \quad \bar{\mathbf{q}} = \frac{\sum_{i=1}^{n} w_i \mathbf{q}_i}{\sum_{i=1}^{n} w_i}.$$
 (3)

By rearranging the terms of (2) we get

$$\mathbf{t} = \bar{\mathbf{q}} - R\bar{\mathbf{p}}.\tag{4}$$

In other words, the optimal translation  $\mathbf{t}$  maps the transformed weighted centroid of P to the weighted centroid of Q. Let us plug the optimal  $\mathbf{t}$  into our objective function:

$$\mathbf{t} = \bar{\mathbf{q}} - R\bar{\mathbf{p}}.\tag{4}$$

In other words, the optimal translation  $\mathbf{t}$  maps the transformed weighted centroid of P to the weighted centroid of Q. Let us plug the optimal  $\mathbf{t}$  into our objective function:

$$\sum_{i=1}^{n} w_i \| (R\mathbf{p}_i + \mathbf{t}) - \mathbf{q}_i \|^2 = \sum_{i=1}^{n} w_i \| R\mathbf{p}_i + \bar{\mathbf{q}} - R\bar{\mathbf{p}} - \mathbf{q}_i \|^2 =$$
 (5)

$$= \sum_{i=1}^{n} w_i \|R(\mathbf{p}_i - \bar{\mathbf{p}}) - (\mathbf{q}_i - \bar{\mathbf{q}})\|^2.$$
 (6)

We can thus concentrate on computing the rotation R by restating the problem such that the translation would be zero:

$$\mathbf{x}_i := \mathbf{p}_i - \bar{\mathbf{p}}, \quad \mathbf{y}_i := \mathbf{q}_i - \bar{\mathbf{q}}. \tag{7}$$

So we look for the optimal rotation R such that

$$R = \underset{R}{\operatorname{argmin}} \sum_{i=1}^{n} w_i \| R \mathbf{x}_i - \mathbf{y}_i \|^2.$$
 (8)

#### Shape Matching: Rotation

Approximate nonlinear rotation by general matrix

$$\min_{\mathbf{R}} \sum_{i} \|\hat{\mathbf{p}}_{i} - \mathbf{R}\hat{\mathbf{q}}_{i}\|^{2} \rightarrow \min_{\mathbf{A}} \sum_{i} \|\hat{\mathbf{p}}_{i} - \mathbf{A}\hat{\mathbf{q}}_{i}\|^{2}$$

The least squares linear transformation is

$$\mathbf{A} = \left(\sum_{i=1}^{m} \hat{\mathbf{p}}_{i} \hat{\mathbf{q}}_{i}^{T}\right) \cdot \left(\sum_{i=1}^{m} \hat{\mathbf{q}}_{i} \hat{\mathbf{q}}_{i}^{T}\right)^{-1} \in \mathbb{R}^{3 \times 3}$$

SVD & Polar decomposition extracts rotation from A

$$\mathbf{A} = \mathbf{U} \mathbf{\Sigma} \mathbf{V}^T \quad \rightarrow \quad \mathbf{R} = \mathbf{U} \mathbf{V}^T$$

#### Iterative Closest Points (ICP)-The classic version

- Given two point sets P and Q, establish correspondence between each point at P and its closest point at Q.
- Iterate between two steps:
  - 1. Use the estimated correspondence to estimate the best rigid **transformation** and align the shapes.
  - 2. Derive new **correspondence** from the new alignment.
- Stop when there is no significant change.
- The initial alignment is critical!

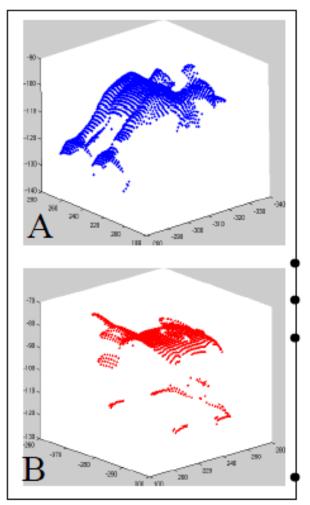
#### Suggestions:

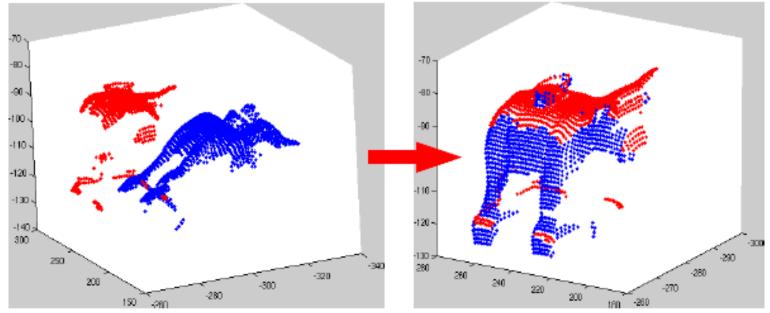
- 1. Use a set of set of feature points/ user markers.
- 2. Use PCA to align the shapes.
- 3. Use the symmetry axes to align the shapes.

## Registration – ICP algorithm

Example : registration of 3D model parts (toy cow)







Input: point clouds acquired from non-aligned viewpoints (from 3D range scanner) & initial estimation of registration

**Output:** Transformation

Variants on the following stages of ICP have been proposed:

- 1. Selecting source points (from one or both meshes)
- 2. Matching to points in the other mesh
- 3. Weighting the correspondences
- 4. Rejecting certain (outlier) point pairs
- 5. Assigning an error metric to the current transform
- 6. Minimizing the error metric w.r.t. transformation

- Can analyze various aspects of performance:
  - Speed
  - Stability
  - Tolerance of noise and/or outliers
  - Maximum initial misalignment
- Comparisons of many variants in
  - [Rusinkiewicz & Levoy, 3DIM 2001]: Efficient Variants of the ICP Algorithm

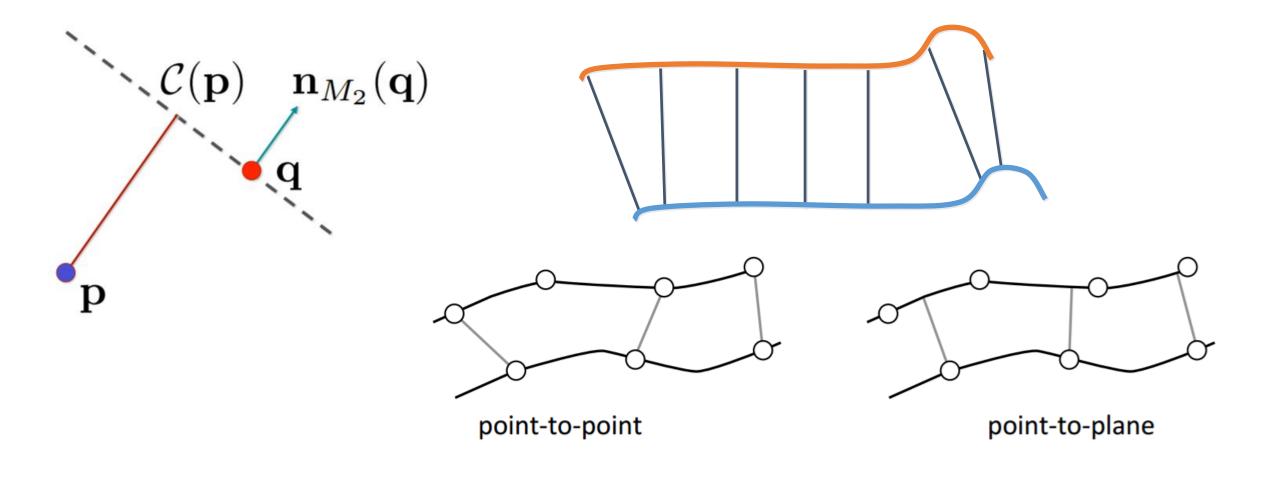
- 1. Selecting source points (from one or both meshes)
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a huge (an order of magnitude or more) difference in convergence

- 5. Assigning an error metric to the current transform
- 6. Minimizing the error metric w.r.t. transformation

#### Point-to-Plane Error Metric

- Using point-to-plane distance instead of point-to-point
- allows flat regions slide along each other [Chen & Medioni 91]



#### Point-to-Plane Error Metric

Error function:

$$E = \sum \left( (\mathbf{R}\mathbf{p}_i + \mathbf{t} - \mathbf{q}_i)^{\top} \mathbf{n_i} \right)^2$$

where  ${f R}$  is a rotation matrix,  ${f t}$  is a translation vector

• Linearize (i.e. assume that  $\sin \theta \approx \theta$ ,  $\cos \theta \approx 1$ ):

$$E \approx \sum \left( (\mathbf{p}_i - \mathbf{q}_i)^{\top} \mathbf{n}_i \right) + \mathbf{r}^{\top} (\mathbf{p}_i \times \mathbf{n}_i) + \mathbf{t}^{\top} \mathbf{n}_i)^2 \qquad \mathbf{r} = \begin{bmatrix} r_x \\ r_y \\ r_z \end{bmatrix}$$

Result: overconstrained linear system

#### Point-to-Plane Error Metric

$$E \approx \sum ((\mathbf{p}_i - \mathbf{q}_i)^{\top} \mathbf{n}_i) + \mathbf{r}^{\top} (\mathbf{p}_i \times \mathbf{n}_i) + \mathbf{t}^{\top} \mathbf{n}_i)^2$$

Overconstrained linear system

$$\mathbf{A}\mathbf{x} = \mathbf{b}$$

$$\mathbf{A} = \begin{bmatrix} \leftarrow & \mathbf{p}_1 \times \mathbf{n}_1 & \rightarrow & \leftarrow & \mathbf{n}_1 & \rightarrow \\ \leftarrow & \mathbf{p}_2 \times \mathbf{n}_1 & \rightarrow & \leftarrow & \mathbf{n}_2 & \rightarrow \\ \vdots & & & \vdots & & \end{bmatrix} \quad \mathbf{x} = \begin{bmatrix} r_x \\ r_y \\ r_z \\ t_x \\ t_y \\ t_z \end{bmatrix} \quad \mathbf{b} = \begin{bmatrix} -(\mathbf{p}_1 - \mathbf{q}_1)^{\top} \mathbf{n}_1 \\ -(\mathbf{p}_2 - \mathbf{q}_2)^{\top} \mathbf{n}_2 \\ \vdots & & \vdots \end{bmatrix}$$

Solve using least squares

$$\mathbf{A}^{\top} \mathbf{A} \mathbf{x} = \mathbf{A}^{\top} \mathbf{b}$$
$$\mathbf{x} = (\mathbf{A}^{\top} \mathbf{A})^{-1} \mathbf{A}^{\top} \mathbf{b}$$

#### Rotation matrix from axis and angle

For some applications, it is helpful to be able to make a rotation with a given axis. Given a unit vector  $\mathbf{u} = (u_X, u_Y, u_Z)$ , where  $u_X^2 + u_Y^2 + u_Z^2 = 1$ , the matrix for a rotation by an angle of  $\theta$  about an axis in the direction of  $\mathbf{u}$  is [3]

$$R = egin{bmatrix} \cos heta + u_x^2 \left( 1 - \cos heta 
ight) & u_x u_y \left( 1 - \cos heta 
ight) - u_z \sin heta & u_x u_z \left( 1 - \cos heta 
ight) + u_y \sin heta \ u_y u_x \left( 1 - \cos heta 
ight) + u_z \sin heta & \cos heta + u_y^2 \left( 1 - \cos heta 
ight) & u_y u_z \left( 1 - \cos heta 
ight) - u_x \sin heta \ u_z u_x \left( 1 - \cos heta 
ight) - u_y \sin heta & u_z u_y \left( 1 - \cos heta 
ight) + u_x \sin heta & \cos heta + u_z^2 \left( 1 - \cos heta 
ight) \ \end{bmatrix}$$

A derivation of this matrix from first principles can be found in section 9.2. here.<sup>[4]</sup>

This can be written more concisely as

$$R = \cos heta \mathbf{I} + \sin heta [\mathbf{u}]_ imes + (1 - \cos heta) \mathbf{u} \otimes \mathbf{u}$$

where  $[\mathbf{u}]_{\times}$  is the cross product matrix of  $\mathbf{u}$ ,  $\otimes$  is the tensor product and /is the Identity matrix, or

alternatively as 
$$R_{jk} = egin{cases} \cos^2rac{ heta}{2} + \sin^2rac{ heta}{2}\left(2u_j^2 - 1
ight), & ext{if } j = k \ 2u_ju_k\sin^2rac{ heta}{2} + \epsilon_{jkl}u_l\sin heta, & ext{if } j 
eq k \end{cases}$$

where  $\epsilon_{jkl}$  is the Levi-Civita symbol with  $\epsilon_{123}=1$ . This is a matrix form of Rodrigues' rotation formula, (or the equivalent, differently parameterized Euler–Rodrigues formula) with<sup>[5]</sup>

$$\mathbf{u}\otimes\mathbf{u}=\mathbf{u}\mathbf{u}^T=egin{bmatrix} u_x^2 & u_xu_y & u_xu_z\ u_xu_y & u_y^2 & u_yu_z\ u_xu_z & u_yu_z & u_z^2 \end{bmatrix}, \qquad [\mathbf{u}]_ imes =egin{bmatrix} 0 & -u_z & u_y\ u_z & 0 & -u_x\ -u_y & u_x & 0 \end{bmatrix}.$$

$$E = \sum ((\mathbf{R}\mathbf{p}_i + \mathbf{t} - \mathbf{q}_i)^{\top} \mathbf{n}_i)^2 \quad \Longrightarrow \quad E \approx \sum ((\mathbf{p}_i - \mathbf{q}_i)^{\top} \mathbf{n}_i) + \mathbf{r}^{\top} (\mathbf{p}_i \times \mathbf{n}_i) + \mathbf{t}^{\top} \mathbf{n}_i)^2$$

$$R = \cos\theta I + \sin\theta [r]_{\times} + (1 - \cos\theta I) r(r)^{T}$$

• Key: 
$$(Rp_i)^T n_i = ((I + [r]_{\times})p_i)^T n_i$$

 $([r]_{\times}p_i)^T n_i ?= (r)^T (p_i \times n_i)$ 

$$\mathbf{a} imes\mathbf{b}=[\mathbf{a}]_{ imes}\mathbf{b}=egin{bmatrix}0&-a_3&a_2\a_3&0&-a_1\-a_2&a_1&0\end{bmatrix}egin{bmatrix}b_1\b_2\b_3\end{bmatrix}$$

- $(r \times p_i)^T n_i$  ?=  $(r)^T (p_i \times n_i)$
- $(r \times p_i)^T n_i = n_i \cdot (r \times p_i)$ ;  $(r)^T (p_i \times n_i) = r \cdot (p_i \times n_i)$
- $n_i \cdot (r \times p_i)$ ? =  $r \cdot (p_i \times n_i)$

$$\mathbf{a} \cdot (\mathbf{b} \times \mathbf{c}) = \mathbf{b} \cdot (\mathbf{c} \times \mathbf{a}) = \mathbf{c} \cdot (\mathbf{a} \times \mathbf{b})$$

## Improving ICP Stability

- Closest compatible point
- Stable sampling

- 1. Selecting source points (from one or both meshes)
- 2. Matching to points in the other mesh
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#### Closest Compatible Point

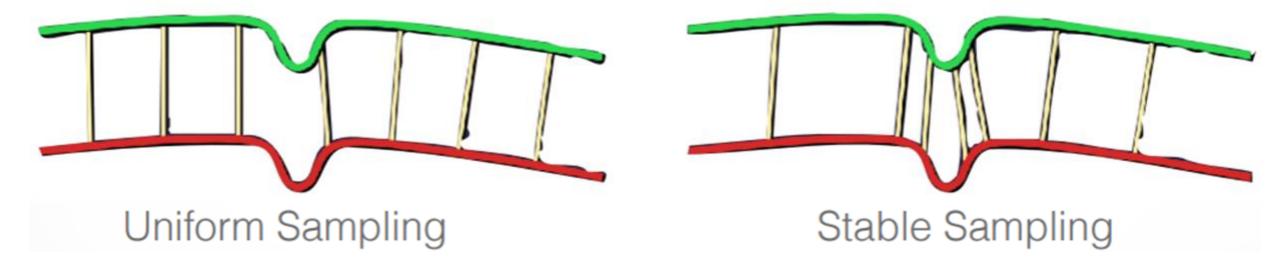
- Closest point often a bad approximation to corresponding point
- Can improve matching effectiveness by restricting match to compatible points
  - Compatibility of colors [Godin et al. '94]
  - Compatibility of normals [Pulli '99]
  - Other possibilities: curvature, higher-order derivatives, and other local features (remember: data is noisy)
  - New challenge: how to assign proper weights to color, normal or ...

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## Selecting Source Points

- Use all points
- Uniform subsampling
- Random sampling
- Stable sampling [Gelfand et al. 2003]
  - Select samples that constrain all degrees of freedom of the rigid-body transformation

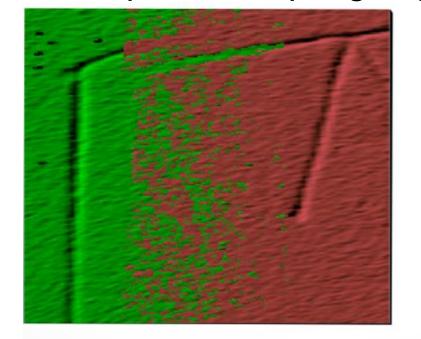


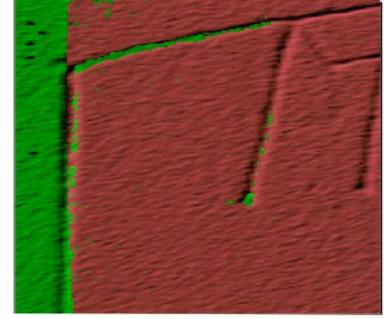
## Sample Selection

- Simpler variant: normal-space sampling
  - select points with uniform distribution of normals
  - Pro: faster, does not require eigenanalysis
  - Con: only constrains translation

Stability-based or normal-space sampling important for smooth areas

with small features





Normal-space Sampling

## Selection vs. Weighting

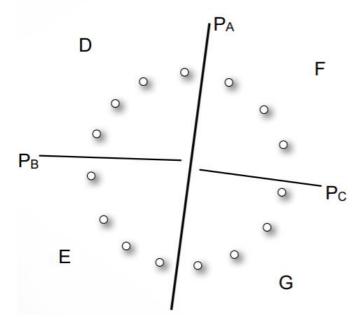
- Could achieve same effect with weighting
- Hard to ensure enough samples in features except at high sampling rates
- However, have to build special data structure
- Preprocessing / run-time cost tradeoff

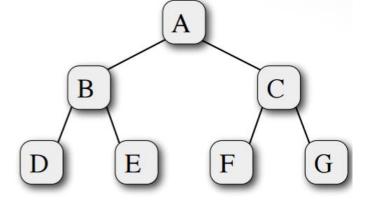
#### **ICP** Variants

- Can analyze various aspects of performance:
  - Speed
  - Stability
  - Tolerance of noise and/or outliers
  - Maximum initial misalignment
- Comparisons of many variants in
  - [Rusinkiewicz & Levoy, 3DIM 2001]

#### Closest Point Search

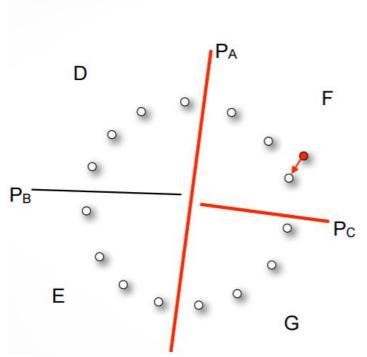
- most expensive stage of the ICP algorithm
  - Brute force search O(n)

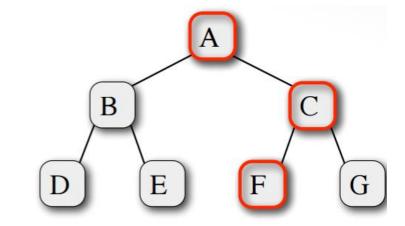




- Use Hierarchical BSP tree
  - Binary space partitioning tree (general kD-tree)
  - Recursively partition 3D space by planes
  - Tree should be balanced, put plane at median
  - log(n) tree levels, complexity O(nlog n)

#### **BSP Closest Point Search**

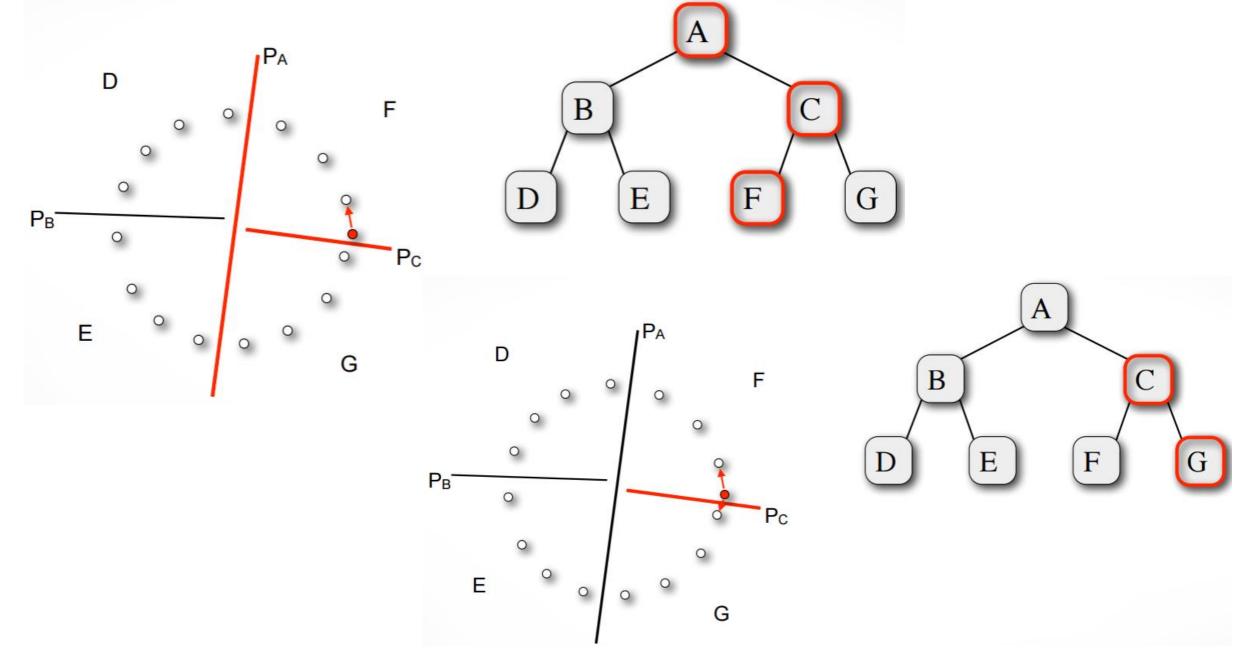




#### Implement BSPNode::dist() with the following info:

```
BSPNode left_child, right_child;
vector<Point> p; // p[i] is the ith point
...
bool leaf_node();
void dist(Point x, Scalar& dmin): x: the query point, dmin: min distance
between x and its closest point in the Tree.
};
float dist = dist_to_plane(x); // a signed distance for assigning x to left
or right of the plane & ...
```

#### How to handle this?

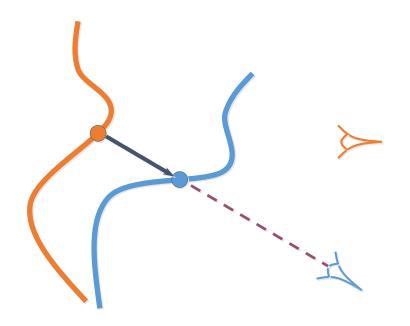


#### **BSP Closest Point Search**

```
BSPNode::dist(Point x, Scalar& dmin)
  if (leaf node())
    for each sample point p[i]
      dmin = min(dmin, dist(x, p[i]));
  else
    d = dist_to_plane(x);
    if (d < 0)
      left child->dist(x, dmin);
      if (|d| < dmin) right_child->dist(x, dmin);
    else
      right child->dist(x, dmin);
      if (|d| < dmin) left_child->dist(x, dmin);
```

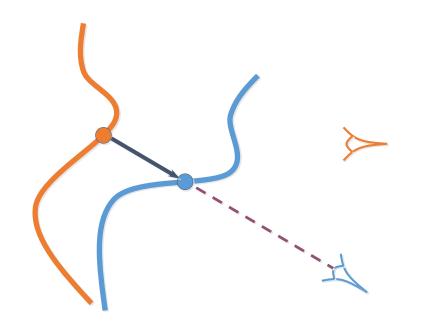
#### Closest Point Search

- most expensive stage of the ICP algorithm
  - Brute force search O(n)
  - Binary space partitioning tree (general kD-tree) O(nlog n)



#### Projection to Find Correspondence

- Idea: use a simpler algorithm to find correspondences
- For range images, can simply project point by "reverse calibration" [Blais 95]
  - Constant-time
  - Does not require precomputing a spatial data structure



$$oldsymbol{ ilde{x}}_{\scriptscriptstyle S} = oldsymbol{K} \left[ egin{array}{c} oldsymbol{R} & oldsymbol{t} \end{array} 
ight] oldsymbol{p}_w = oldsymbol{P} oldsymbol{p}_w$$
Calibration Matrix

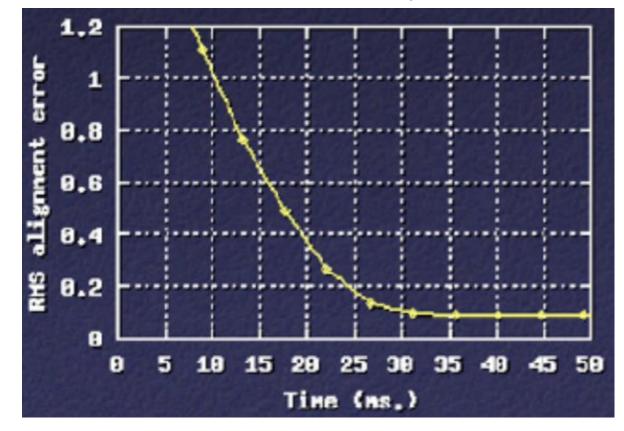
Camera Matrix

## Projection-Based Matching

- Slightly worse performance per iteration
- Each iteration is one to two orders of magnitude faster than closest point

Result: can align two range images in a few milliseconds (0.001)

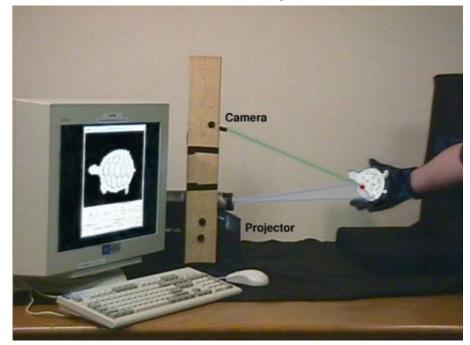
seconds), vs. a few seconds



#### **Applications**

- Given:
  - A scanner that returns range images in real time
  - Fast ICP
  - Real-time merging and rendering
- Result: 3D model acquisition
  - Tight feedback loop with user
  - Can see and fill holes while scanning

## Scanner Layout







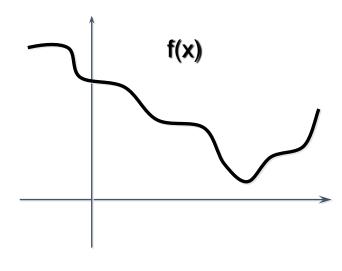


Artec Group

[Newcombe et al. '11] KinectFusion

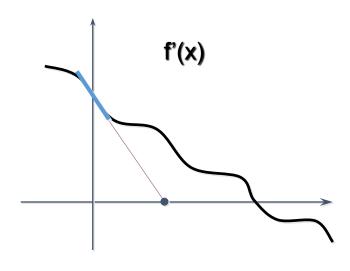
#### What Does ICP Do?

- Two ways of thinking about ICP:
  - Solving the correspondence problem
  - Minimizing point-to-surface squared distance
- ICP is like Newton's method on an approximation of the distance function



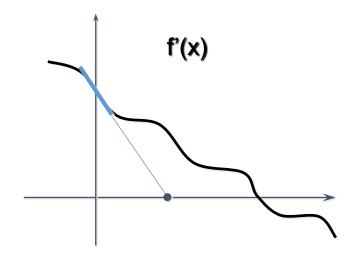
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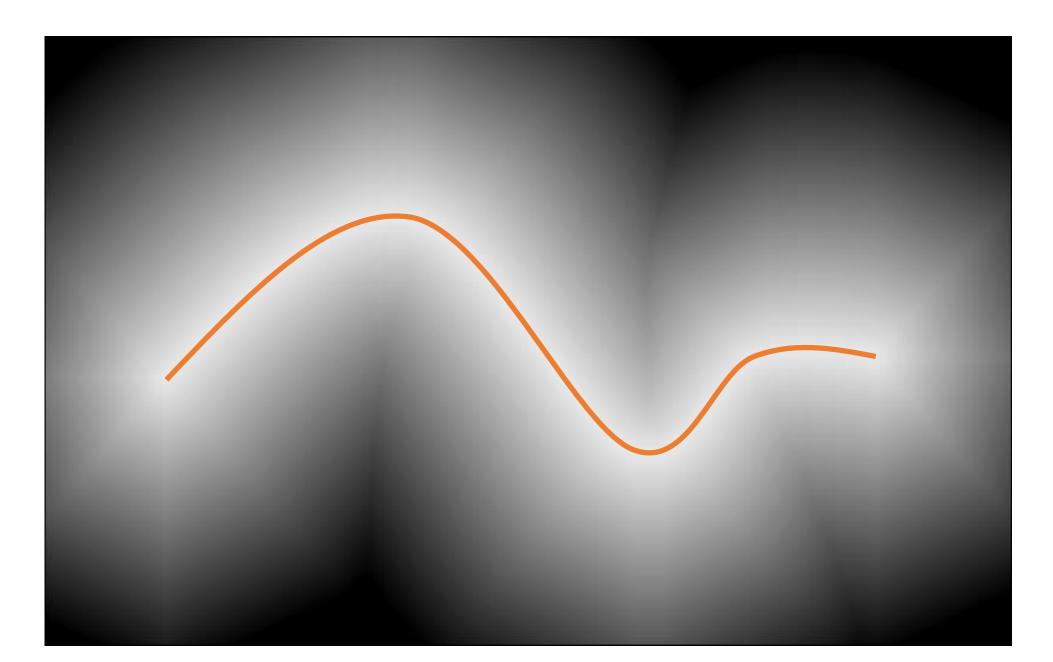


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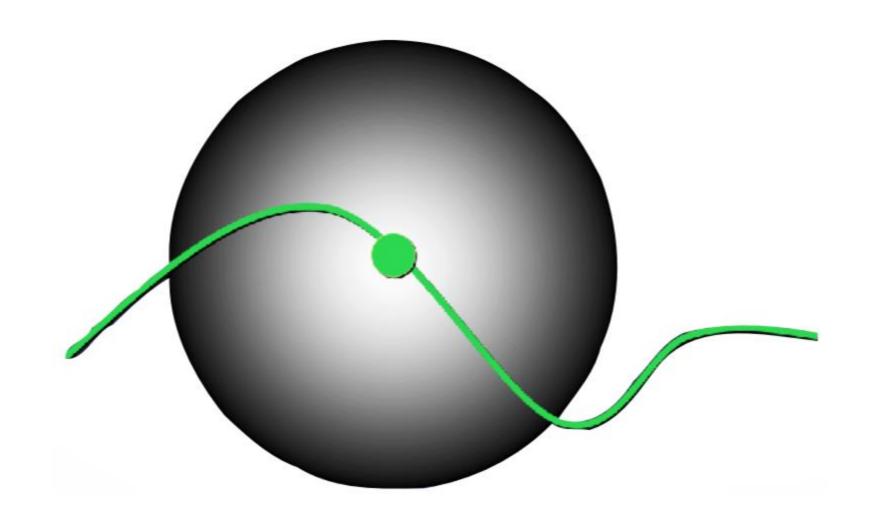
- Two ways of thinking about ICP:
  - Solving the correspondence problem
  - Minimizing point-to-surface squared distance
- ICP is like Newton's method on an approximation of the distance function
  - ICP variants affect shape of global error function or local approximation



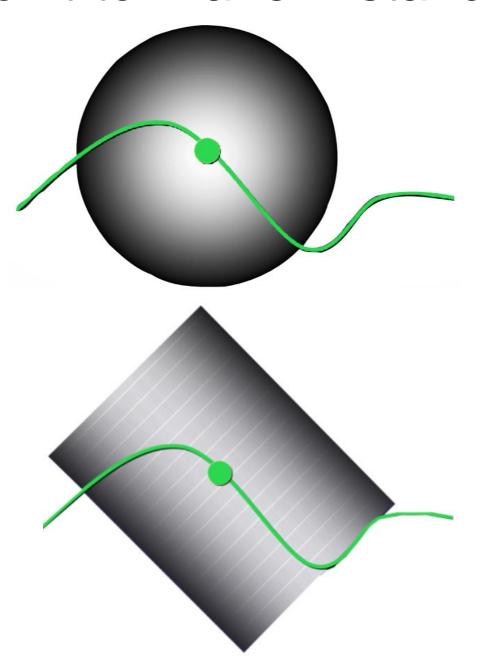
#### Point-to-Surface Distance

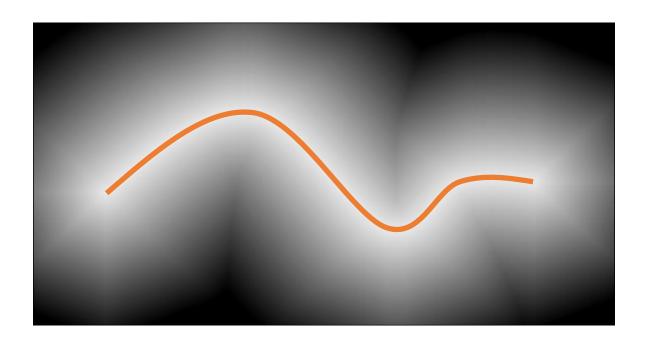


## Point-to-Point Distance



#### Point-to-Plane Distance



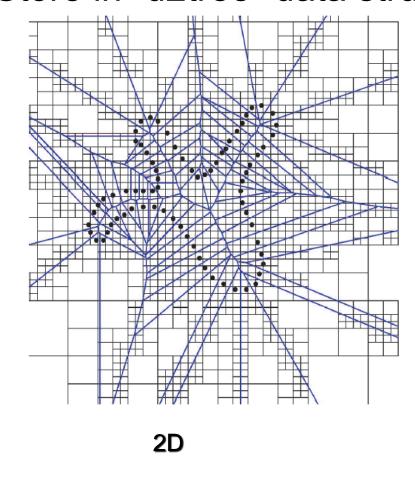


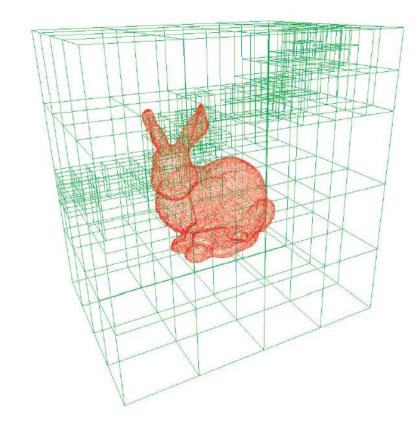
#### Soft Matching and Distance Functions

- Soft matching equivalent to standard ICP on (some) filtered surface
- Produces filtered version of distance function
  - ⇒ fewer local minima
- Multiresolution minimization [Turk & Levoy 94]
   or soft assign with simulated annealing (good description in [Chui 03])

#### Mitra et al.'s Optimization

- Precompute piecewise-quadratic approximation to distance field throughout space
- Store in "d2tree" data structure





3D

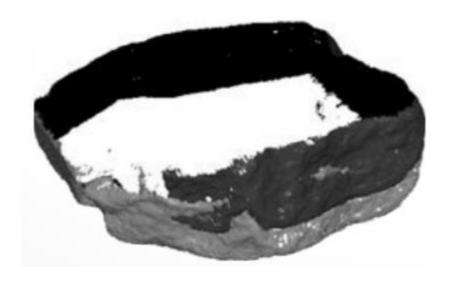
#### Mitra et al.'s Optimization

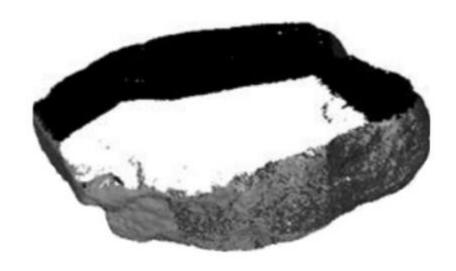
- Precompute piecewise-quadratic approximation to distance field throughout space
- Store in "d2tree" data structure

- At run time, look up quadratic approximants and optimize using Newton's method
  - More robust, wider basin of convergence
  - Often fewer iterations, but more precomputation

## Global Registration Goal

- Given: n scans around an object
- Goal: align them all
- First attempt: apply ICP to each scan to one other
- Want method for distributing accumulated error among all scans





#### Approach #1: Avoid the Problem

 In some cases, have 1 (possibly low-resolution) scan that covers most surface

Align all other scans to this "anchor" [Turk 94]

Disadvantage: not always practical to obtain anchor scan

## Approach #2: The Greedy Solution

- Align each new scan to the union of all previous scans [Masuda '96]
- Disadvantages:
  - Order dependent
  - Doesn't spread out error
    - This sometimes avoids catastrophic accumulation of error, but really isn't guaranteed to do anything.

#### Approach #3: The Brute-Force Solution

- While not converged:
  - For each scan:
    - For each point:
      - For every other scan
        - Find closest point
  - Minimize error w.r.t. transforms of all scans

- Disadvantage:
  - Solve (6n)x(6n) matrix equation, where n is number of scans

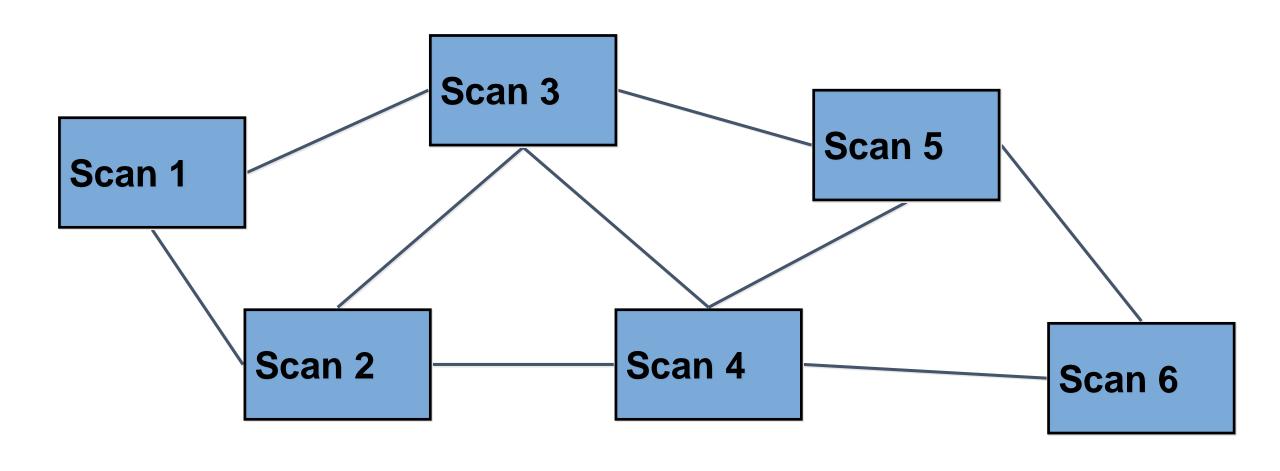
## Approach #3a: Slightly Less Brute-Force Solution

- While not converged:
  - For each scan:
    - For each point:
      - For every other scan
        - Find closest point
  - Minimize error w.r.t. transforms of this scans

- Faster than previous method (matrices are 6x6) [Bergevin '96, Benjemaa '97]
- Previous: Solve (6n)x(6n) matrix equation, where n is number of scans

#### Graph Methods

 Many globalreg algorithms create a graph of pairwise alignments between scans



## Pulli's Algorithm

 Perform pairwise ICPs, record sample (e.g. 200) of corresponding points

- For each scan, starting w. most connected
  - Align scan to existing set
  - While (change in error) > threshold
    - Align each scan to others
- All alignments during globalreg phase use precomputed corresponding points

## Sharp et al. Algorithm

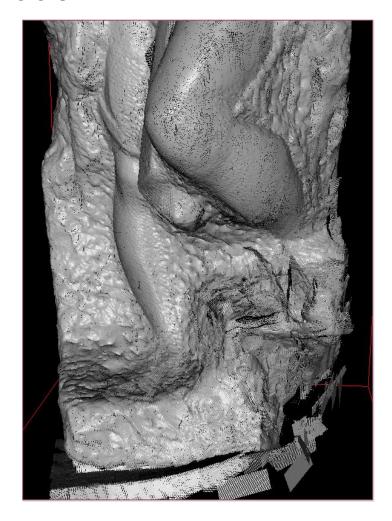
- Perform pairwise ICPs, record only optimal rotation/translation for each
- Decompose alignment graph into cycles
- While (change in error) > tolerance
  - For each cycle:
    - Spread out error equally among all scans in the cycle
  - For each scan belonging to more than 1 cycle:
    - Assign average transform to scan

## Bad ICP in Globalreg

One bad ICP can throw off the entire model!



**Correct Globalreg** 



**Globalreg Including Bad ICP** 

## Acknowledgement

- Data-Driven Shape Anlaysis CS468 @Stanford.edu, Spring 2014
   CSCI 621: Digital Geometry Processing SS 2017 @ USC
- 3D Scan Matching and Registration, Part II, ICCV 2005 Short Course

#### Literature

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2018/2/28

# Thanks