Spring 2019

CSCI 621: Digital Geometry Processing

# 5.2 Surface Registration



## Acknowledgement

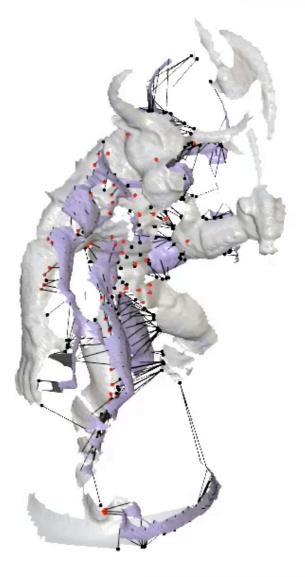
#### Images and Slides are courtesy of

- Prof. Szymon Rusinkiewicz, Princeton University
- ICCV Course 2005: http://gfx.cs.princeton.edu/proj/ iccv05\_course/



## **Surface Registration**

# Align two partially-overlapping meshes given initial guess for relative transform

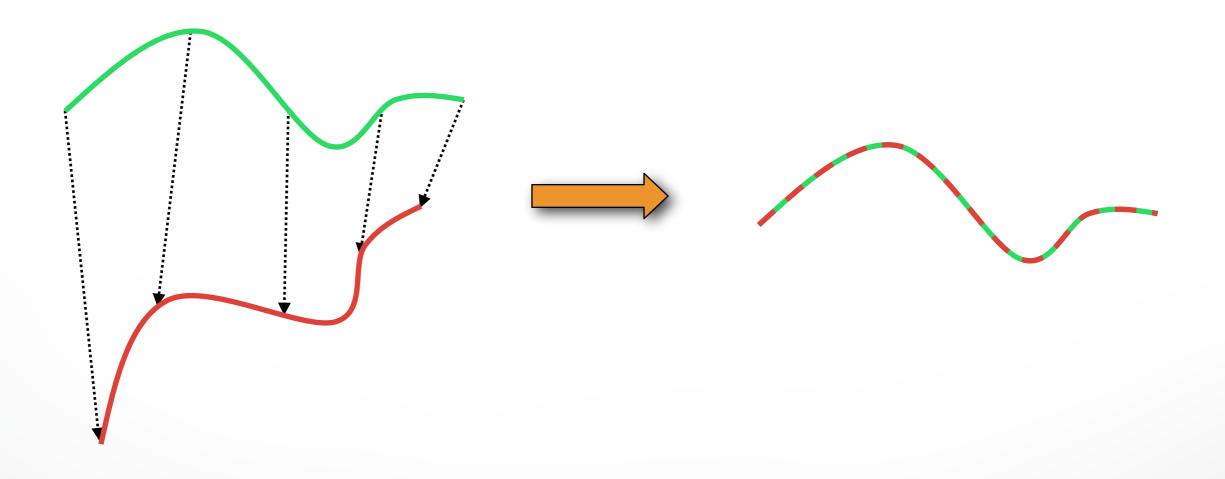


## Outline

- ICP: Iterative Closest Points
- Classification of ICP variants
  - Faster alignment
  - Better robustness
- ICP as function minimization

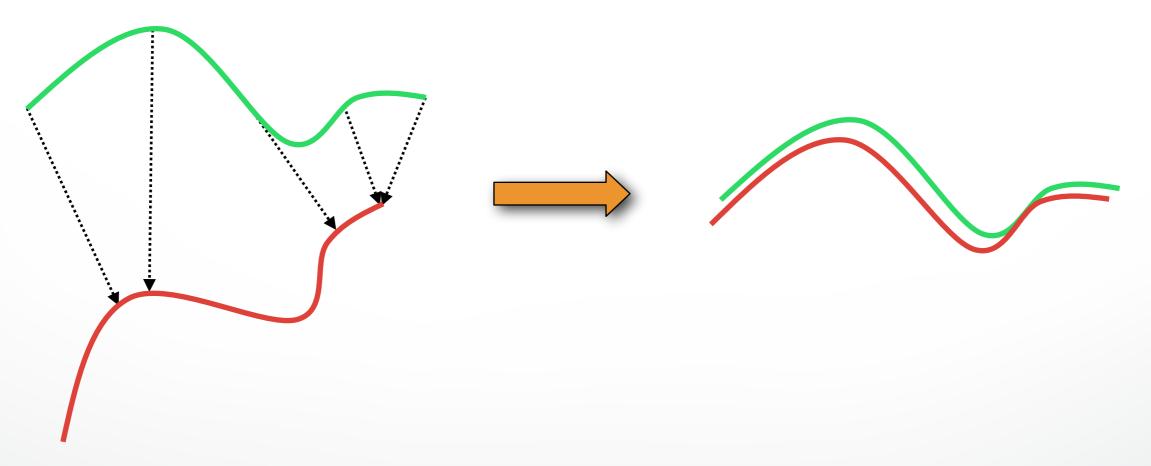
## **Aligning 3D Data**

If correct correpondences are known, can find correct relative rotation/translation



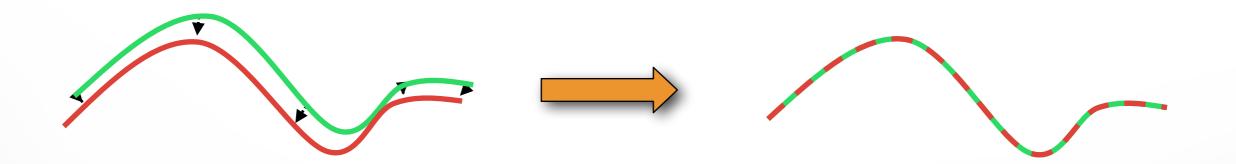
# **Aligning 3D Data**

- How to find correspondences: User input? Feature detection? Signatures?
- 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, using data structure such as k-d tree
- **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])

## **Shape Matching: Translation**

Define bary-centered point sets

$$\bar{\mathbf{p}} := \frac{1}{m} \sum_{i=1}^{m} \mathbf{p}_i \qquad \bar{\mathbf{q}} := \frac{1}{m} \sum_{i=1}^{m} \mathbf{q}_i$$

$$\hat{\mathbf{p}}_i := \mathbf{p}_i - \bar{\mathbf{p}} \qquad \hat{\mathbf{q}}_i := \mathbf{q}_i - \bar{\mathbf{q}}$$

Optimal translation vector t maps barycenters onto each other

$$t = \bar{p} - R\bar{q}$$

## **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

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

# 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 (outliers) 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

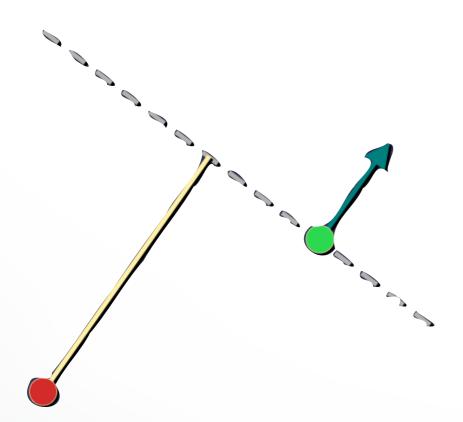
#### **Comparisons of many variants in**

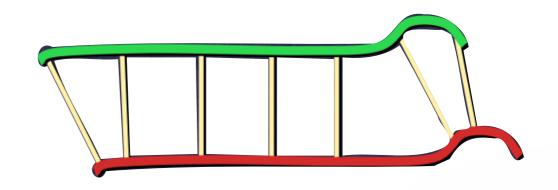
• [Rusinkiewicz & Levoy, 3DIM 2001]

- 1. Selecting source points (from one or both meshes)
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## **Point-to-Plane Error Metric**

Using point-to-plane distance instead of point-to-point lets 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**

Overconstrained linear system

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

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

1

• Solve using least squares

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

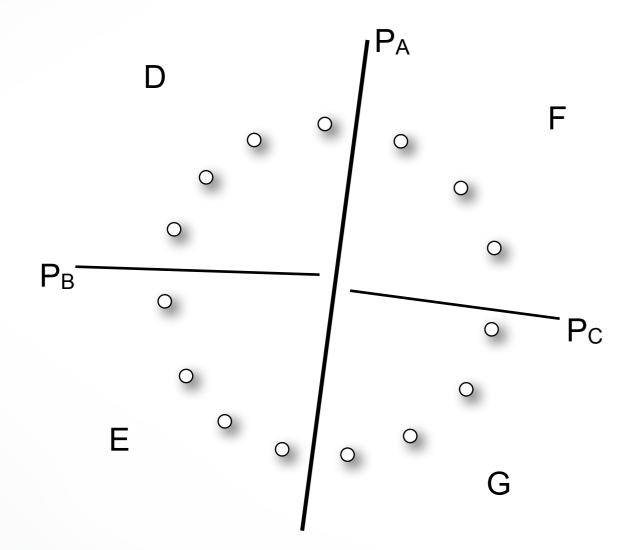
## Improving ICP Stabilitiy

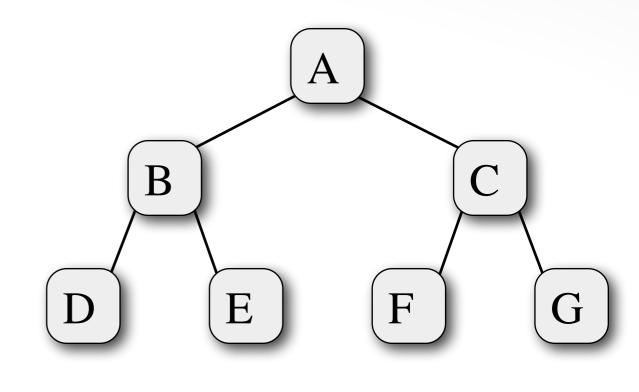
- Closest compatible point
- Stable sampling

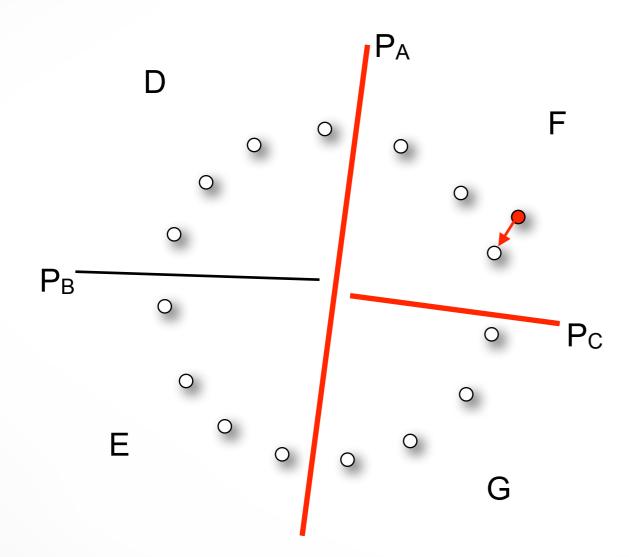
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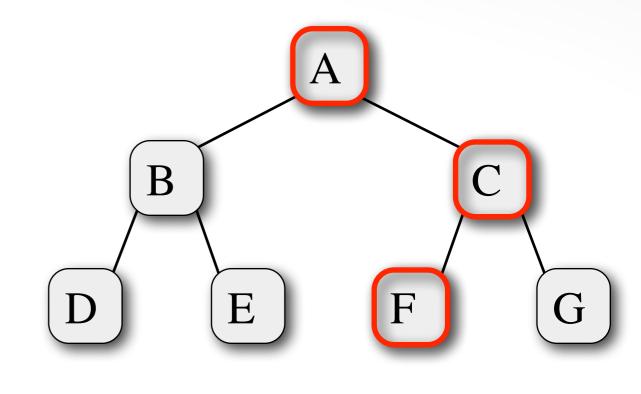
- Find closest point of a query point
  - Brute force: O(n) complexity

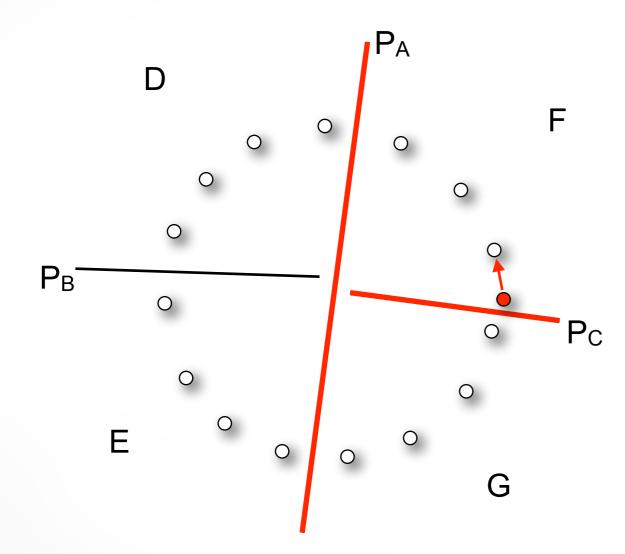
- Use Hierarchical BSP tree
  - Binary space partitioning tree (general kD-tree)
  - Recursively partition 3D space by planes
  - Tree should be balanced, put plane at median
  - log(n) tree levels, complexity O(nlog n)

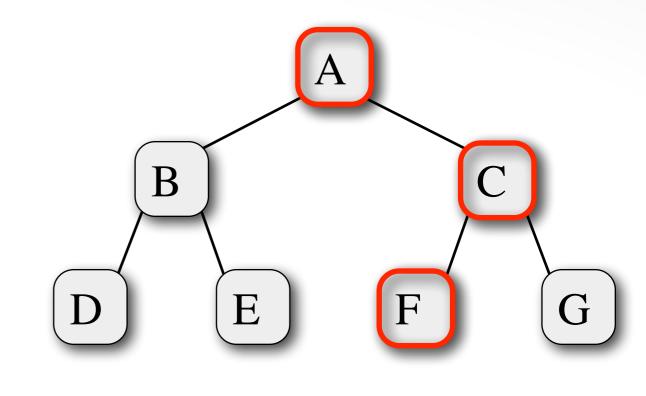


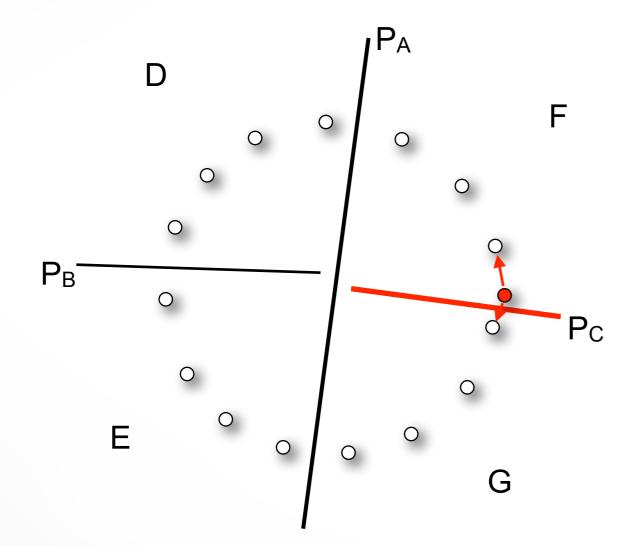


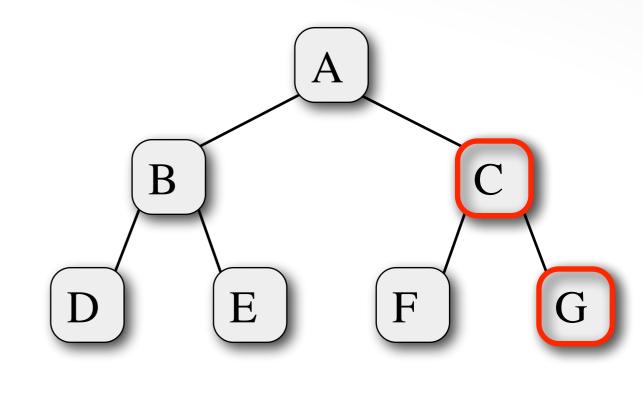


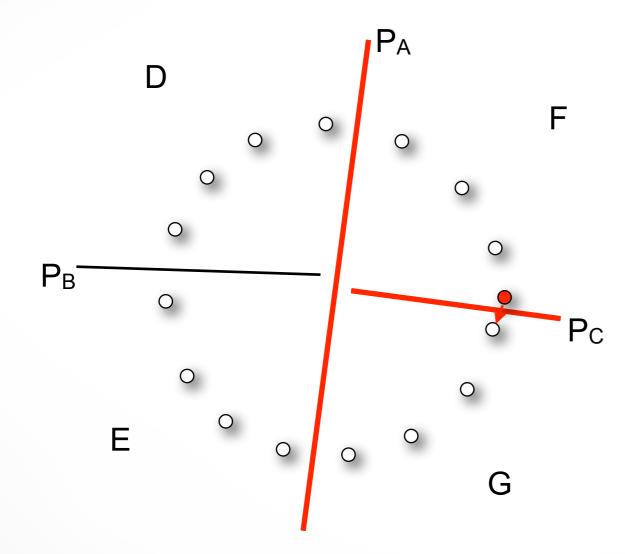


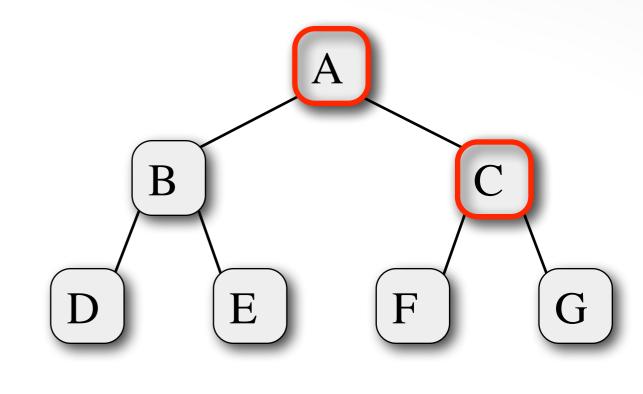












```
BSPNode::dist(Point x, Scalar& dmin)
{
  if (leaf node())
    for each sample point p[i]
      dmin = min(dmin, dist(x, p[i]));
  else
  {
    d = dist to plane(x);
    if (d < 0)
      left child->dist(x, dmin);
      if (|d| < dmin) right child->dist(x, dmin);
    }
    else
      right child->dist(x,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

## **Closest 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)

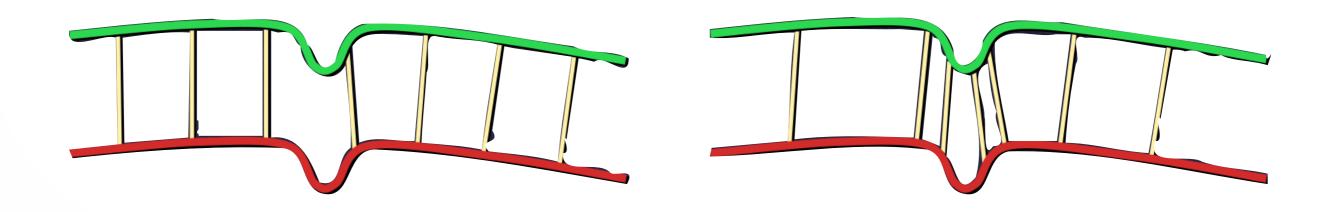
#### 1. Selecting source points (from one or both meshes)

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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

## **Stable Sampling**



Uniform Sampling

Stable Sampling

## **Covariance Matrix**

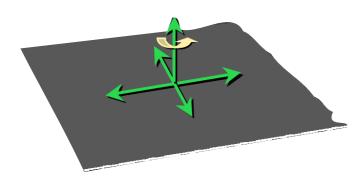
• Aligning transform is given by  $\mathbf{A}^{\top}\mathbf{A}\mathbf{x} = \mathbf{A}^{\top}\mathbf{b}$ , where

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

• Covariance matrix  $\mathbf{C} = \mathbf{A}^{\top} \mathbf{A}$  determines the change in error when surfaces are moved from optimal alignment

# **Sliding Directions**

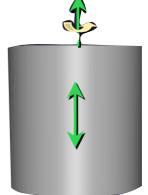
• Eigenvectors of  $\mathbf{C}$  with small eigenvalues correspond to sliding transformations



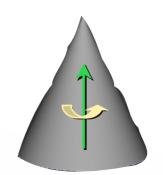
**3 small eigenvalues**2 translation1 rotation



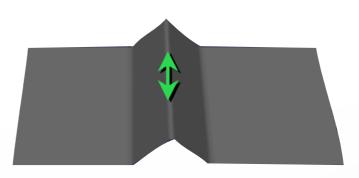
**3 small eigenvalues** 3 rotation



2 small eigenvalues1 translation1 rotation



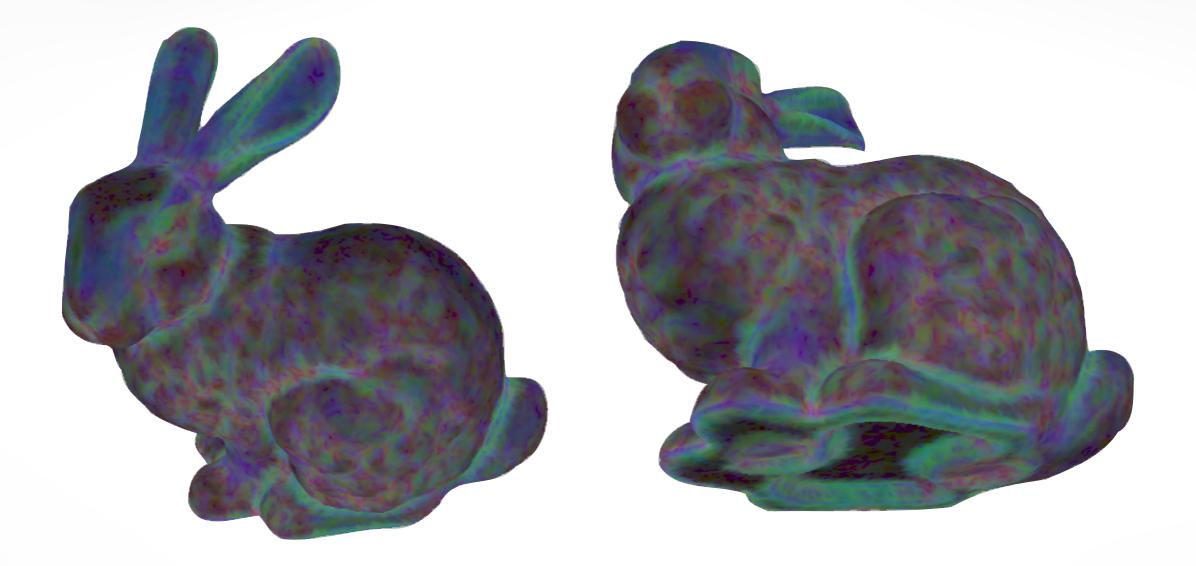
small eigenvalue
 rotation

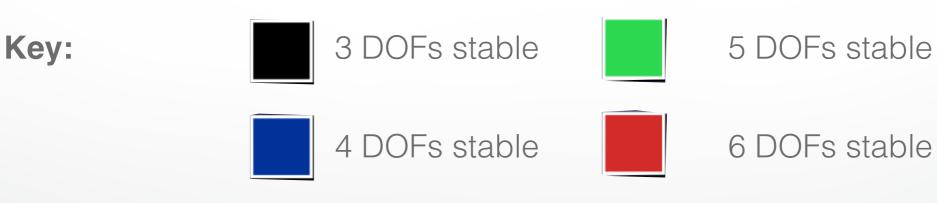


**1 small eigenvalue** 1 translation

[Gelfand]

## **Stability Analysis**





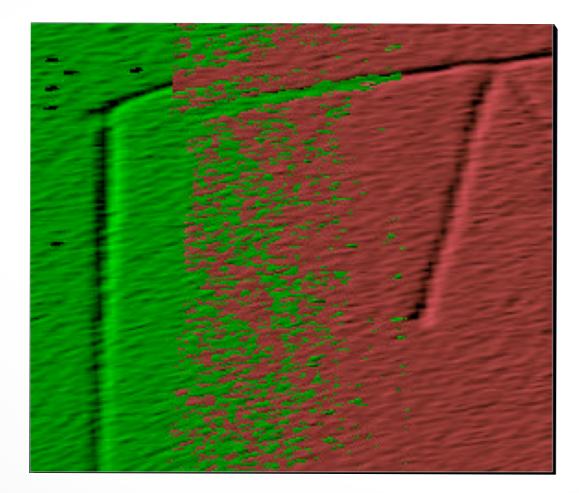
## **Sample Selection**

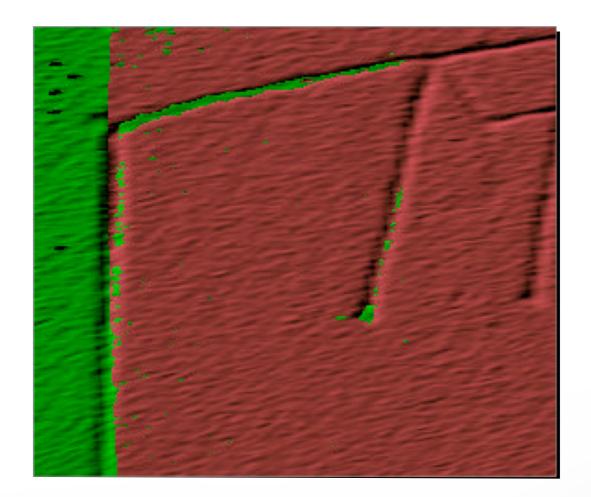
- Select points to prevent small eigenvalues
  - Based on C obtained from sparse sampling

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

## Result

Stability-based or normal-space sampling important for smooth areas with small features





#### Random Sampling

#### 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

# Improving ICP Speed

Projection-based matching

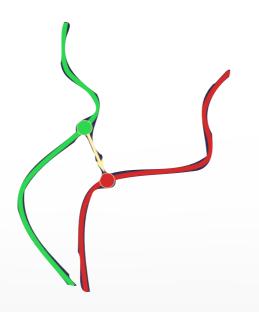
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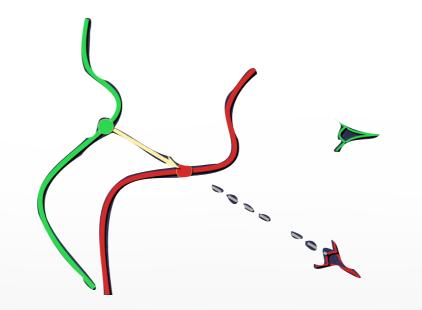
# **Finding Corresponding Points**

- Finding Closest point is most expensive stage of the ICP algorithm
  - Brute force search O(n)
  - Spatial data structure (e.g., k-d tree) O(log n)



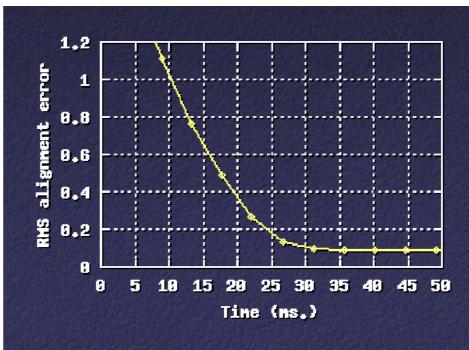
## **Projection to Find Correspondence**

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



#### **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, vs. a few seconds



# Application

- 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

#### Examples



[Rusinkiewicz et al. '02]

Artec Group

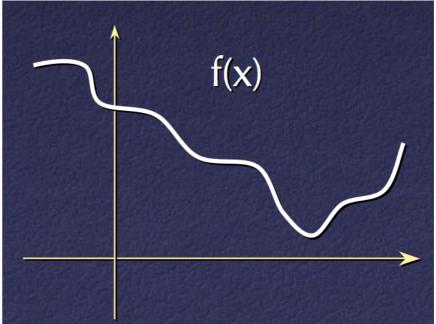
[Newcombe et al. '11] KinectFusion

#### **Theoretical Analysis of ICP Variants**

- One way of studying performance is via empirical tests on various scenes
- How to analyze performance analytically?
- For example, when does point-to-plane help? Under what conditions does projection-based matching work?

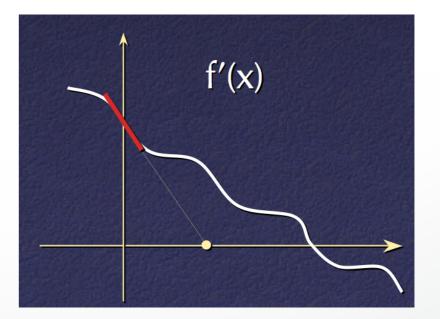
## What does ICP do?

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



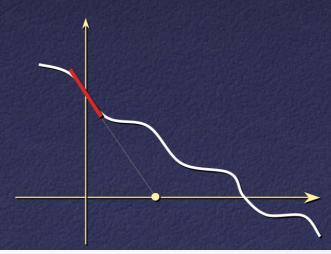
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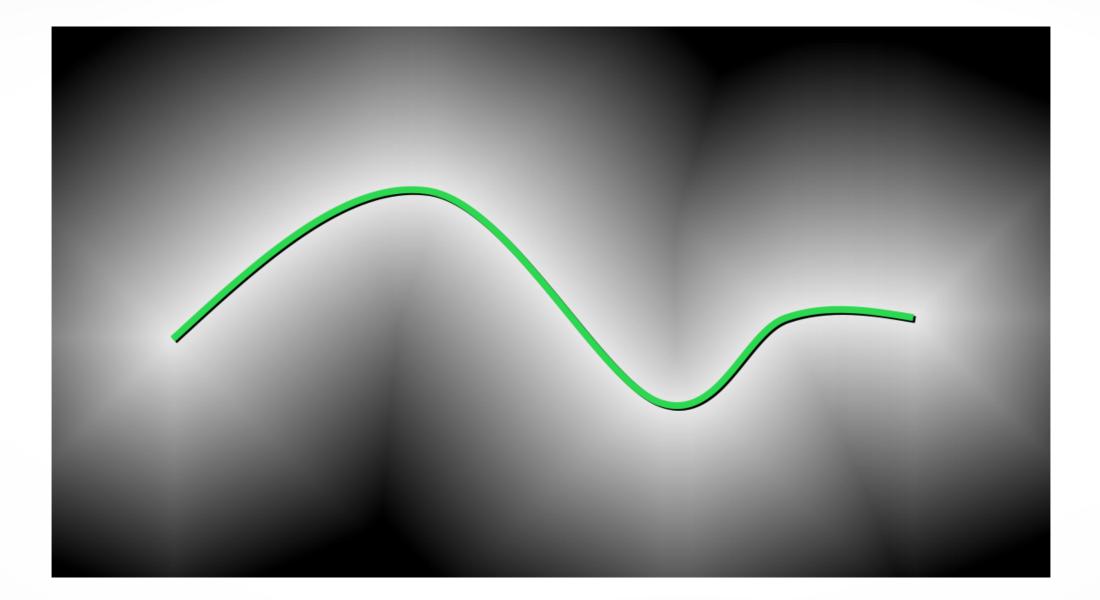


## What does ICP do?

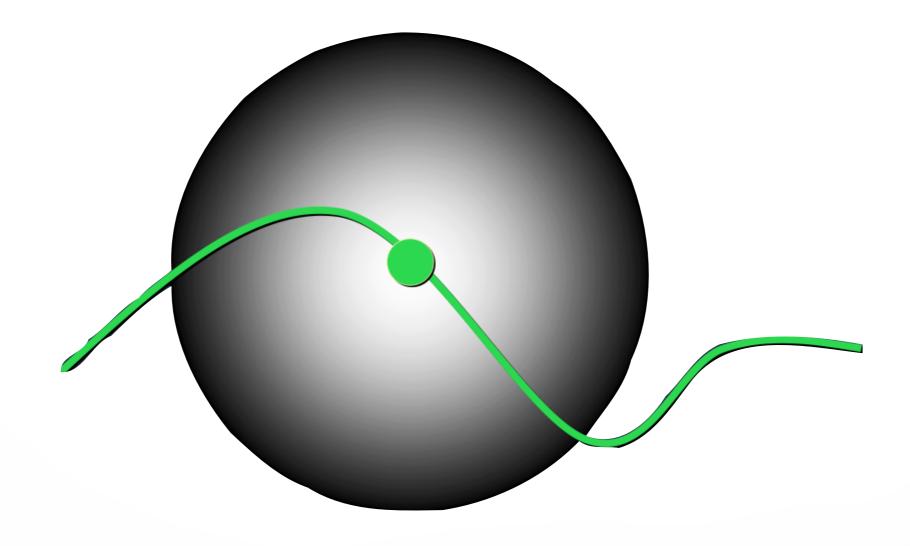
- Two ways of thinking about ICP:
  - Solving correspondence problem
  - Minimizing point-to-surface squared distance
- ICP is like Newton's method on an approximation of the distance function
  - ICP variants affect shape of the global error function or local approximation



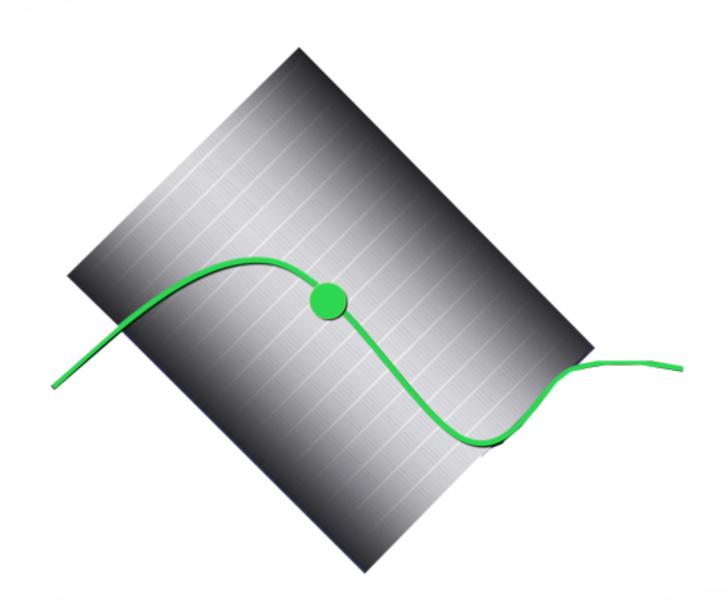
#### **Point-to-Surface Distance**



#### **Point-to-Point Distance**

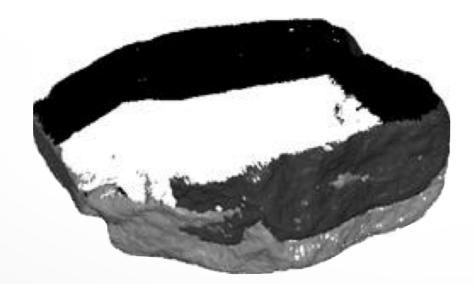


#### **Point-to-Plane Distance**



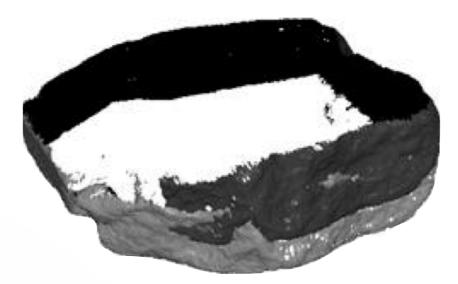
#### **Global Registration Goal**

- Given: n scans around an object
- Goal: align them all
- First attempt: apply ICP to each scan to one other



#### **Global Registration Goal**

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 all previous scans [Masuda '96]
- Disadvantages:
  - Order dependent
  - Doesn't spread out error

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

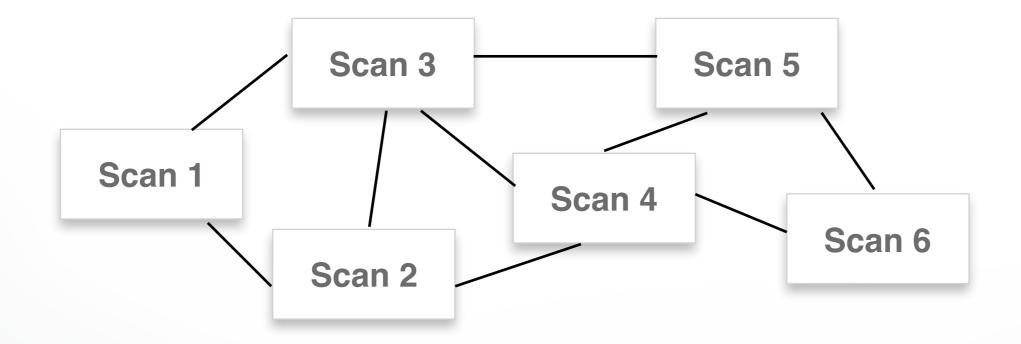
- While not converged:
  - For each scan:
    - For each point:
      - For every other scan
        - Find closest point
  - Minimize error w.r.t. transforms of **all** scans
- Disadvantage:
  - Solve (6n)x(6n) matrix equation, where n is number of scans

## **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]

#### **Graph Methods**

 Many global registration algorithms create a graph of pairwise alignments between scans



# Sharp et al. 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 global reg phase use precomputed corresponding points.

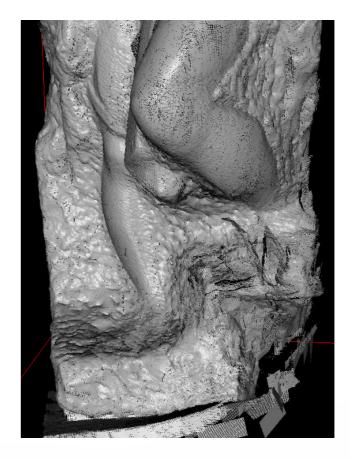
## Lu and Milios Algorithm

- Perform pairwise ICPs, record optimal rotation/translation and covariance for each
- Least squares simultaneous minimization of all errors (covariance-weighted)
- Requires linearization of rotations
  - Worse than the ICP case, since don't converge to (incremental rotation) = 0

#### **Bad ICP in Global Registration**

#### One bad ICP can throw off the entire model!





Correct Global Registration Global Registration Including Bad ICP

#### Literature

- Rusinkiewicz & Levoy, Efficient Variants of the ICP Algorithm, 3DIM 2001
- Chen & Medioni, "Object modeling by registration of multiple range images", ICRA1991
- Besl & McKay: A method for registration of 3D shapes, PAMI 1992
- Horn: Closed-form solution of absolute orientation using unit quaternions, Journal Opt. Soc. Amer. 4(4), 1987
- Gelfand et al: Geometrically Stable Sampling for the ICP Algorithm, 3DIM, 2001.
- Pulli, Multiview Registration for Large Data Sets, 3DIM 1999

#### http://cs621.hao-li.com

# Thanks!

