



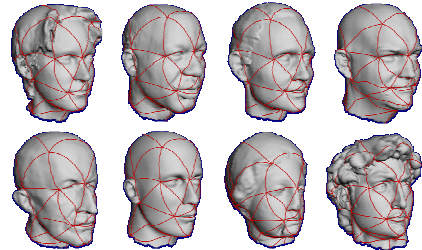
# Surface Registration

Thomas Funkhouser  
COS 526, Fall 2006



## Goal

- Establish 1 to 1 mapping between points on one 3D surface and corresponding points on a different surface



Praun



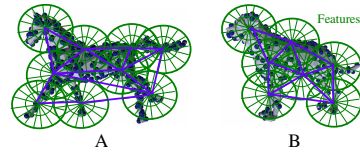
## Motivation - Matching

- Determine geometric similarity of surfaces



## Motivation - Matching

- Determine geometric similarity of surfaces



$$D(A,B) = \sum_{\text{Correspondences}} \Delta\text{FeatureShape} + \sum_{\text{Correspondence Pairs}} \Delta\text{SpatialConsistency}$$



## Motivation – Common parameterization

- Registration provides consistent parameterization
  - Allows for basic operations like matching, mean, signal processing, etc.

$$\frac{1}{n} \left( \text{Head}_1 + \text{Head}_2 + \text{Head}_3 + \dots \right) = \text{Mean Head}$$

Praun



## Motivation - Morphing

- Smoothly transition from one surface to another
  - When registered, simply use linear combinations of vertices



Allen03

## Motivation – Attribute Transfer



- Copy attributes from one surface to another
  - Texture transfer (below)
  - Deformation weight transfer
  - Segmentation transfer

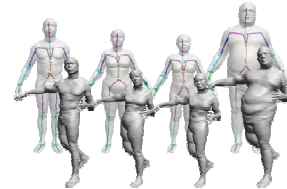


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## Motivation – Attribute Transfer



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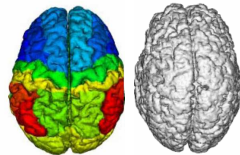


Allen03

## Motivation – Attribute Transfer



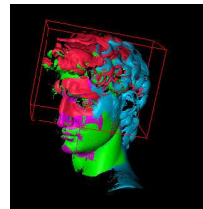
- Copy attributes from one surface to another
  - Texture transfer
  - Deformation weight transfer
  - Segmentation transfer (below)



## Motivation – Scanning



- Combine multiple scans to form complete surface
  - Must align scans from different views

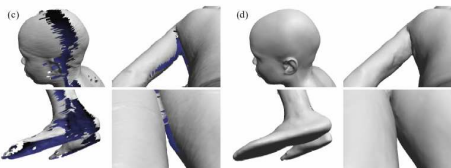


Rusinkiewicz

## Motivation – Hole Filling



- Use surface of one model to fill holes of another
  - e.g., to fix surfaces captured with range scanners



## Registration Goal



- Find minimal dissimilarity measure between surfaces over class of possible transformations



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## Registration Methods



- Underlying issues:
  - Transformation type
  - Surface representation
  - Dissimilarity measure
  - Algorithmic strategy

## Registration Methods - Part 1



- Transformation Type
  - Rigid: mutual distances of points within a model are conserved during transformation

$$x_B = R_{AB}x_A + t_{AB}$$

R is a rotation matrix and t is a translation vector

- Non-rigid: account for surface deformations in the transformation
  - e.g., Affine transformation
  - e.g., Thin plate spline

## Registration Methods - Part 2



- Surface Representation
  - Surface description
    - Points, mesh, splines, etc.
  - Surface features
    - Curvature extrema, saddle points, ridges, etc.
  - Shape descriptors
    - Harmonic shape descriptors, spin images

## Registration Methods - Part 3



- Dissimilarity measure
  - Distance
    - Distances between corresponding points after alignment
  - Deformation
    - Amount of deformation implied by alignment

## Registration Methods - Part 4



- Algorithmic strategy
  - Optimization
    - Iterative methods
    - Simulated annealing
  - Voting
    - Pose clustering
    - Geometric Hashing
    - Generalized Hough Transform

## Example – Registering Human Bodies



- Algorithm Input
  - Set of human range images
  - Set of colored feature markers
- Algorithm Goal
  - Develop correspondence from template to target
  - Compute affine transform for each vertex
  - Minimize error function

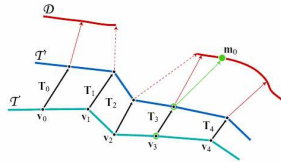
*Allen et al.  
The Space of Human Body Shapes  
Siggraph 2003*



## Optimization Variables



- Algorithm viewed as optimization problem
  - Given an initial template surface with vertices  $v_i$
  - Corresponding affine transformation matrices  $T_i$
  - Current state is  $T_i v_i$  for all  $i$  (see diagram)
  - Find values of  $T_i$  to minimize objective function
  - Attempts to find a "good fit" (blue) of template (cyan) to target (red)



## Objective Function



- Objective Function has three weighted terms
  - Data error
  - Smoothness error
  - Marker error

$$E = \alpha E_d + \beta E_s + \gamma E_m$$

- Will use different weights in each phase of process
  - Multistep / Multi-resolution fitting process

## Objective Function – Marker Error



- Measures distance between pre-labeled markers
  - Correspondences set up beforehand

$$E_m = \sum_{i=1}^m \|\mathbf{T}_{\kappa_i} \mathbf{v}_{\kappa_i} - \mathbf{m}_i\|^2$$

## Objective Function – Data Error



- Data error term requires current match to be close to target
  - Uses distance from each transformed vertex to the target surface
  - Weighted by confidence measure (from scanning)
  - Hole regions have weight = 0
  - Sums total error

$$E_d = \sum_{i=1}^n w_i \text{dist}^2(\mathbf{T}_i \mathbf{v}_i, \mathcal{D})$$

- Distance function
  - Uses transformed template vertex
  - Takes minimum distance to "compatible" vertices in target

## Objective Function – Smoothness



- Measures smoothness of deformation applied to template
  - $E_s$  measures change in  $T_i$  between adjacent vertices
  - Encourages similarly-shaped features to be mapped to each other

$$E_s = \sum_{\{i,j\} \in \text{edges}(\mathcal{T})} \|\mathbf{T}_i - \mathbf{T}_j\|_F^2$$

- Uses Frobenius norm (vector L2 norm)

## Algorithm Procedure



- Minimize error function using L-BFGS-B algorithm
  - Quasi-Newton method with limited memory usage

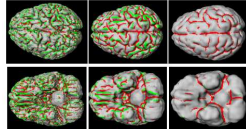
$$E = \alpha E_d + \beta E_s + \gamma E_m$$

- Make four passes over data (2 low res, 2 high res)
  - Fit markers (low res,  $\alpha = 0, \beta = 1, \gamma = 10$ )
  - Refit using data term (low res,  $\alpha = 1, \beta = 1, \gamma = 10$ )
  - Repeat in high resolution (hi res,  $\alpha = 1, \beta = 1, \gamma = 10$ )
  - Refit using predominantly data term (hi res,  $\alpha = 10, \beta = 1, \gamma = 1$ )

## Example – Aligning Brains



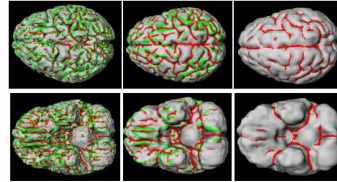
- Algorithm Input
  - Set of human brain surfaces
  - Prelabeled reference brain mesh (low resolution)
- Algorithm Goal
  - Correspondence from template to target
  - Identify particular features in the brain (gyri, sulci)
  - Minimize error function



## Transformation types



- In brains, we see homothetic deformation (local uniform stretch) when aligning features



## Objective Function



- Minimizes error between vertex and feature point

$$O_{ij} = d_{ij} \cdot n_{ij} \cdot f_{ij}$$

- Euclidean distance measure

$$d_{ij} = 1 + \sqrt{[x_i - x_j]^2 + [y_i - y_j]^2 + [z_i - z_j]^2}$$

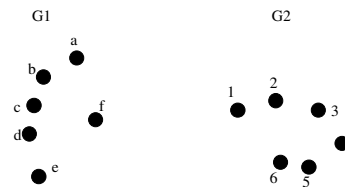
- Surface Normal Match

$$n_{ij} = 2 - \vec{n}_i \cdot \vec{n}_j$$

- Feature Match

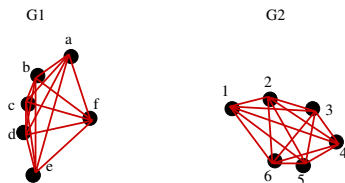
$$f_{ij} = \begin{cases} 1.0, & \text{if } (t_i, t_j) = \begin{cases} (\text{sulcus, sulcus}) \\ (\text{gyrus, gyrus}) \end{cases} \\ & \text{(no feature, no feature)} \\ 2.8, & \text{if } (t_i, t_j) = \begin{cases} (\text{sulcus, no feature}) \\ (\text{gyrus, no feature}) \end{cases} \\ 3.0, & \text{if } (t_i, t_j) = (\text{sulcus, gyrus}) \end{cases} \dots$$

## Example – Aligning Point Sets



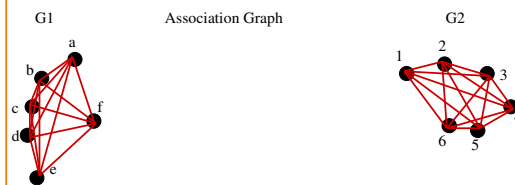
Consider rigid transformations

## Association Graph



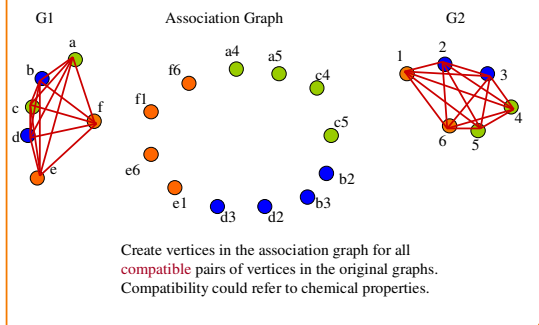
Represent both points sets as complete graphs (G1 and G2).  
(edges connect all pairs of vertices within each point set)

## Association Graph

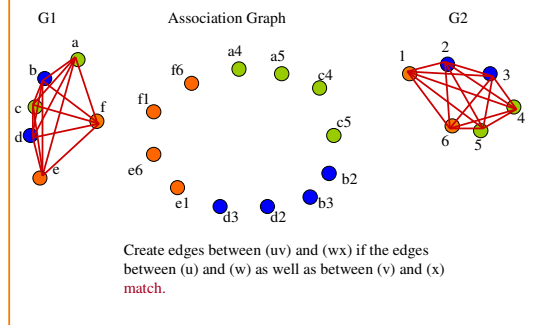


Create vertices in the association graph for all compatible pairs of vertices in the original graphs.  
This can lead to a large number of vertices.

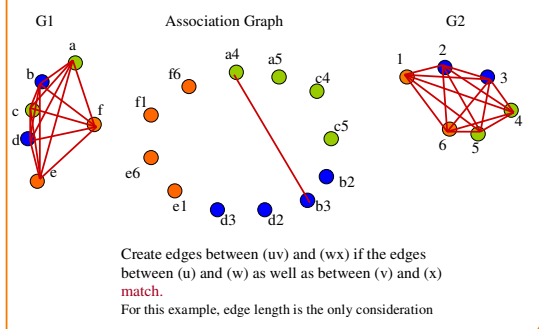
## Association Graph



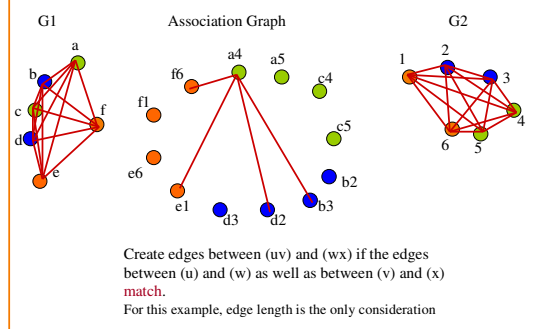
## Association Graph



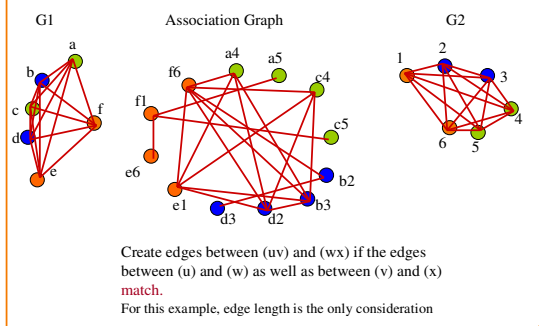
## Association Graph



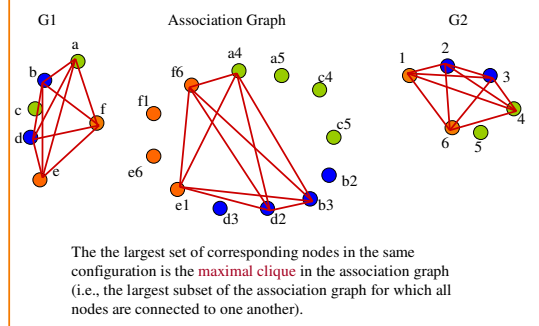
## Association Graph



## Association Graph



## Association Graph



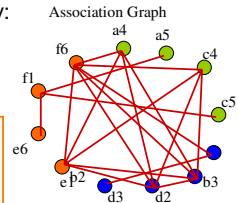
## Association Graph



- Computational complexity:
  - $O(2^n)$  for  $n$  points
  - NP-complete

```

Find the Maximal Clique{
  return Cliques(empty, all nodes)
}
Cliques(X, Y){
  if (no node in Y-X is connected to all of X){
    return X;
  }else{
    y = node in Y connected to all of X;
    return Largest(Cliques(X union y, Y),
                  Cliques(X, Y-y));
  }
}
    
```



[Schmitt02, Brown82]

## Iterative Closest Points (ICP)



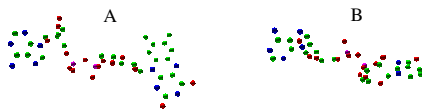
- Assume closest points correspond
  - Avoid finding one-to-one correspondences
- Rigid body transformations
- Greedy optimization procedure
  - Start with rough guess for alignment
  - Iteratively refine transform

[Besl92]

## Iterative Closest Point



- Given two point sets

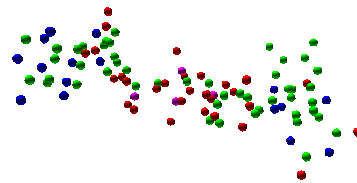


[Besl92]

## Iterative Closest Point



- Given two point sets and an initial guess for the transformation that aligns them

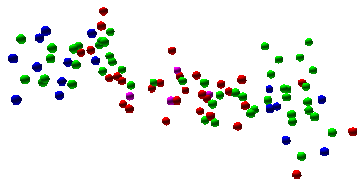


[Besl92]

## Iterative Closest Point



- Assume closest points correspond

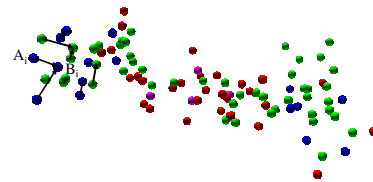


[Besl92]

## Iterative Closest Point



- Assume closest points correspond:  $A \rightarrow B$

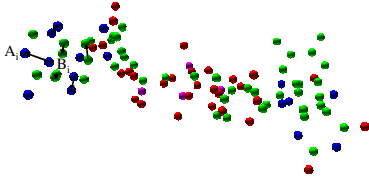


[Besl92]

## Iterative Closest Point



- Assume closest points correspond:  $A \rightarrow B$  and  $B \rightarrow A$

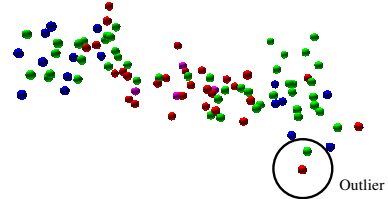


[Bes192]

## Iterative Closest Point



- Rejecting outliers

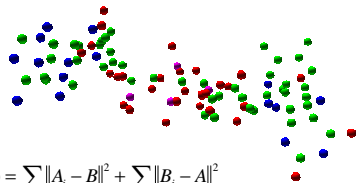


[Bes192]

## Iterative Closest Point



- Find the transformation that optimally aligns proposed correspondences (superposition)



$$d(A, B) = \sum_{A_i \in A} \|A_i - B\|^2 + \sum_{B_j \in B} \|B_j - A\|^2$$

[Bes192]

## Iterative Closest Point



- Iterate until convergence
  - Select source points (from one or both surfaces)
  - Match to points in the other molecule
  - Weight the correspondences
  - Reject outlier point pairs
  - Compute an error metric for the current transform
  - Minimize the error metric w.r.t. transformation

Computational complexity

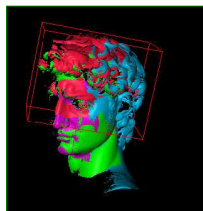
- $O(k * n \log n)$  for  $n$  points per set and  $k$  iterations  
§  $k$  iterations \*  $O(n)$  points \*  $O(\log n)$  to find closest point

Slide courtesy of Szymon Rusinkiewicz

## ICP – Aligning Surfaces (Scans)



- Start with manual initial alignment



Slide courtesy of Szymon Rusinkiewicz

## ICP - Aligning Surfaces (Scans)



- Improve alignment using ICP



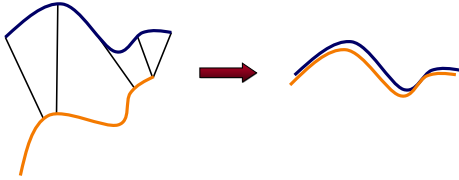
Slide courtesy of Szymon Rusinkiewicz



## ICP - Aligning Surfaces (Scans)



- Assume closest points correspond, compute the best transform...



Slide courtesy of Szymon Rusinkiewicz

## ICP - Aligning Surfaces (Scans)



- ... and iterate to find alignment
- Converges to some local minimum
- Correct if starting position "close enough"



Slide courtesy of Szymon Rusinkiewicz

## Pose Clustering



- General method
  - Enumerate possible transformations
  - Vote for best one
- Methods
  - Pose clustering
  - Geometric Hashing
  - Generalized Hough Transform

## Pose Clustering



- Discretize transformations and scoring

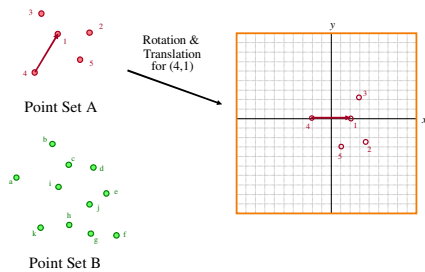


[Wolfson97]

## Pose Clustering



- Discretize transformations and scoring

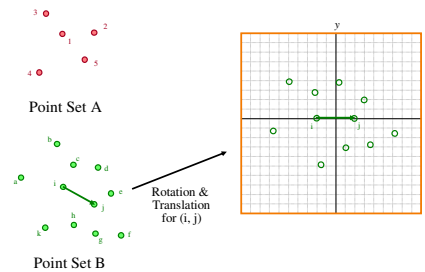


[Wolfson97]

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- Discretize transformations and scoring

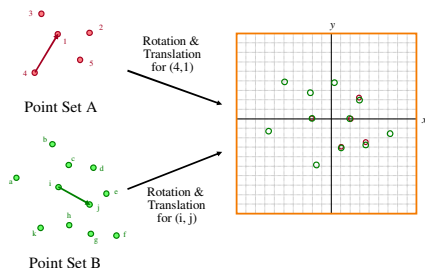


[Wolfson97]

## Pose Clustering



- Discretize transformations and scoring

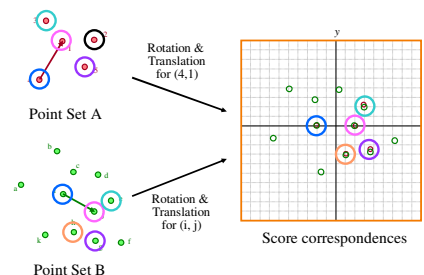


[Wolfson97]

## Pose Clustering



- Discretize transformations and scoring

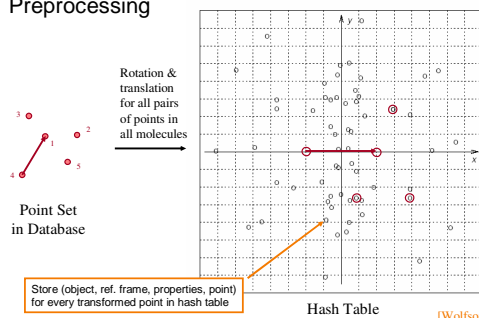


[Wolfson97]

## Geometric Hashing



- Preprocessing

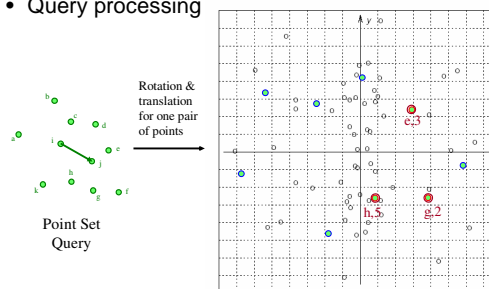


[Wolfson97]

## Geometric Hashing



- Query processing



[Wolfson97]

## Geometric Hashing



- Preprocessing
  - For each triple of points
  - Compute reference frame
  - For each point
    - Transform point into reference frame
    - Hash (molecule, ref. frame, properties, point)
- Query processing
  - Choose any triple of points
  - Compute reference frame
  - For each point
    - Transform point into reference frame
    - For each entry in hash bin for transformed point
    - Vote for (object, ref. frame)

## Geometric Hashing



- Preprocessing complexity
  - $O(n^4)$  for  $n$  points per binding site
    - $O(n^3)$  possible triples \*  $O(n)$  transformations per triple
- Query complexity
  - $O(m)$  \* binsize for  $m$  points in query binding site
    - 1 triple \*  $O(m)$  transformations per triple \* binsize hash processing per transformation

[Wolfson97]

## Summary



- Different methods for different ...
  - Transformation types
  - Surface representations
  - Dissimilarity measures