

# CSE 554

## Lecture 7: Alignment

Fall 2012

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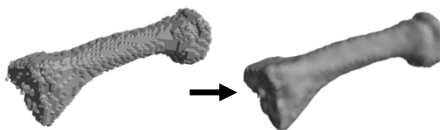
Alignment

Slide 1

## Review

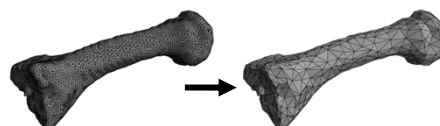
- Fairing (smoothing)

- Relocating vertices to achieve a smoother appearance
- Method: centroid averaging



- Simplification

- Reducing vertex count
- Method: edge collapsing



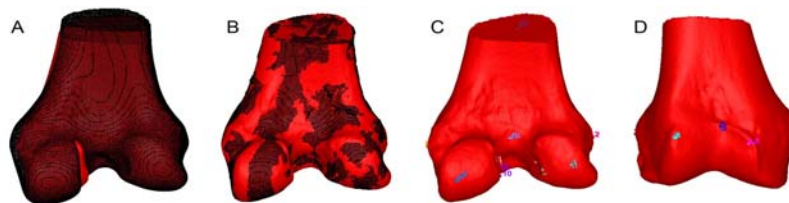
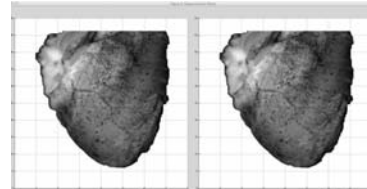
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# Registration

- Fitting one model to match the shape of another
  - Automated annotation
  - Tracking and motion analysis
  - Shape and data comparison



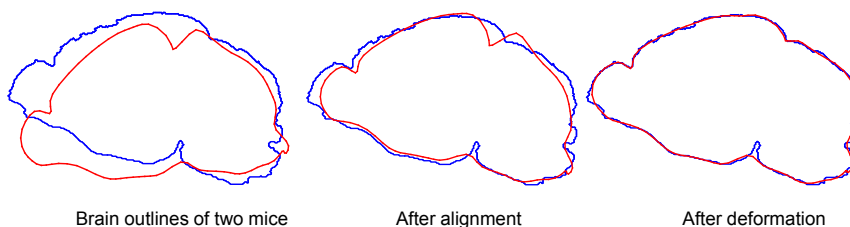
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# Registration

- Challenges: global and local shape differences
  - Imaging causes global shifts and tilts
  - • Requires **alignment**
  - The shape of the organ or tissue differs in subjects and evolve over time
    - Requires **deformation**



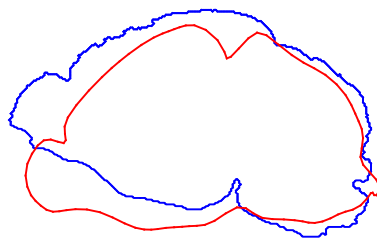
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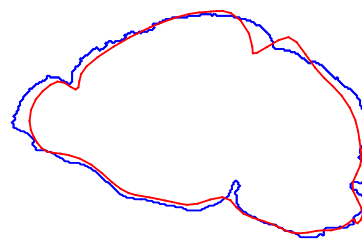
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# Alignment

- Registration by **translation** or **rotation**
  - The structure stays “rigid” under these two transformations
    - Called **rigid-body** or **isometric** (distance-preserving) transformations
  - Mathematically, they are represented as matrix/vector operations



Before alignment



After alignment

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Alignment

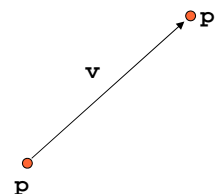
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# Transformation Math

- Translation
  - Vector addition:  $\mathbf{p}' = \mathbf{v} + \mathbf{p}$

— 2D: 
$$\begin{pmatrix} p'_x \\ p'_y \end{pmatrix} = \begin{pmatrix} v_x \\ v_y \end{pmatrix} + \begin{pmatrix} p_x \\ p_y \end{pmatrix}$$

— 3D: 
$$\begin{pmatrix} p'_x \\ p'_y \\ p'_z \end{pmatrix} = \begin{pmatrix} v_x \\ v_y \\ v_z \end{pmatrix} + \begin{pmatrix} p_x \\ p_y \\ p_z \end{pmatrix}$$



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# Transformation Math

- Rotation

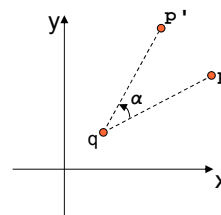
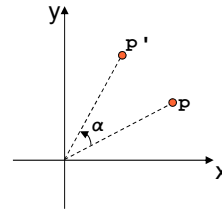
- Matrix product:  $\mathbf{p}' = \mathbf{R} \cdot \mathbf{p}$

- 2D:  $\begin{pmatrix} p'_x \\ p'_y \end{pmatrix} = \mathbf{R} \cdot \begin{pmatrix} p_x \\ p_y \end{pmatrix}$

$$\mathbf{R} = \begin{pmatrix} \cos[\alpha] & -\sin[\alpha] \\ \sin[\alpha] & \cos[\alpha] \end{pmatrix}$$

- Rotate around the origin!
  - To rotate around another point q:

$$\mathbf{p}' = \mathbf{R} \cdot (\mathbf{p} - \mathbf{q}) + \mathbf{q}$$



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# Transformation Math

- Rotation

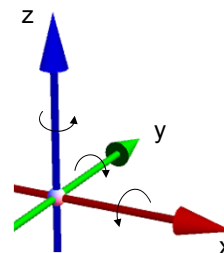
- Matrix product:  $\mathbf{p}' = \mathbf{R} \cdot \mathbf{p}$

- 3D:  $\begin{pmatrix} p'_x \\ p'_y \\ p'_z \end{pmatrix} = \mathbf{R} \cdot \begin{pmatrix} p_x \\ p_y \\ p_z \end{pmatrix}$

Around X axis:  $\mathbf{R}_x = \begin{pmatrix} 1 & 0 & 0 \\ 0 & \cos[\alpha] & -\sin[\alpha] \\ 0 & \sin[\alpha] & \cos[\alpha] \end{pmatrix}$

Around Y axis:  $\mathbf{R}_y = \begin{pmatrix} \cos[\alpha] & 0 & \sin[\alpha] \\ 0 & 1 & 0 \\ -\sin[\alpha] & 0 & \cos[\alpha] \end{pmatrix}$

Around Z axis:  $\mathbf{R}_z = \begin{pmatrix} \cos[\alpha] & -\sin[\alpha] & 0 \\ \sin[\alpha] & \cos[\alpha] & 0 \\ 0 & 0 & 1 \end{pmatrix}$



Any arbitrary 3D rotation  
can be composed from  
these three rotations

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# Transformation Math

- Properties of an arbitrary rotational matrix
  - **Orthonormal** (orthogonal and normal):  $\mathbf{R} \cdot \mathbf{R}^T = \mathbf{I}$ 
    - Examples:
 
$$\begin{pmatrix} \cos[\alpha] & -\sin[\alpha] \\ \sin[\alpha] & \cos[\alpha] \end{pmatrix} \cdot \begin{pmatrix} \cos[\alpha] & \sin[\alpha] \\ -\sin[\alpha] & \cos[\alpha] \end{pmatrix} = \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}$$

$$\begin{pmatrix} 1 & 0 & 0 \\ 0 & \cos[\alpha] & -\sin[\alpha] \\ 0 & \sin[\alpha] & \cos[\alpha] \end{pmatrix} \cdot \begin{pmatrix} 1 & 0 & 0 \\ 0 & \cos[\alpha] & \sin[\alpha] \\ 0 & -\sin[\alpha] & \cos[\alpha] \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$$
    - **Easy to invert:**  $\mathbf{R}^{-1} = \mathbf{R}^T$
  - Any orthonormal matrix represents a rotation around some axis (not limited to X,Y,Z)

# Transformation Math

- Properties of an arbitrary rotational matrix
  - Given an orthonormal matrix, the **angle** of rotation represented by the matrix can be easily calculated from the **trace** of the matrix
    - Trace: sum of diagonal entries
    - 2D: The trace equals  $2 \cos(a)$ , where  $a$  is the rotation angle
    - 3D: The trace equals  $1 + 2 \cos(a)$
  - The larger the trace, the smaller the rotation angle

# Transformation Math

- Eigenvectors and eigenvalues
  - Let  $M$  be a square matrix,  $v$  is an eigenvector and  $\lambda$  is an eigenvalue if:
$$M \cdot v = \lambda \cdot v$$
    - If  $M$  represents a rotation (i.e., orthonormal), the rotation axis is an eigenvector whose eigenvalue is 1.
  - There are at most  $m$  distinct eigenvalues for a  $m$  by  $m$  matrix
  - Any scalar multiples of an eigenvector is also an eigenvector (with the same eigenvalue).

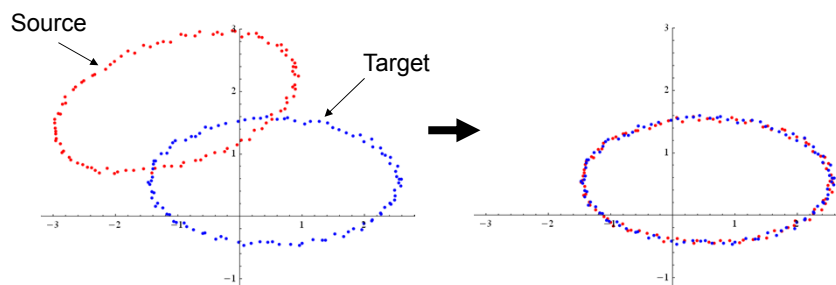
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# Alignment

- Input: two models represented as point sets
  - Source and target
- Output: locations of the translated and rotated source points



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# Alignment

- Method 1: Principal component analysis (PCA)
  - Aligning principal directions
- Method 2: Singular value decomposition (SVD)
  - Optimal alignment given prior knowledge of correspondence
- Method 3: Iterative closest point (ICP)
  - An iterative SVD algorithm that computes correspondences as it goes

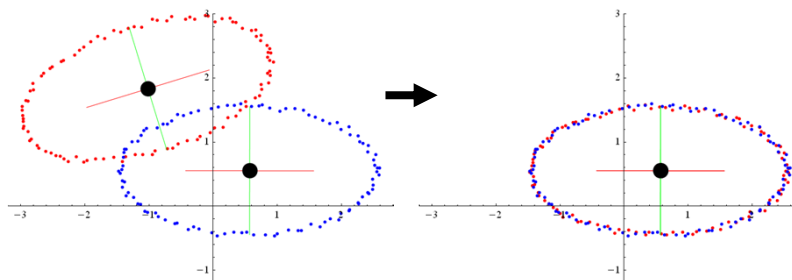
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## Method 1: PCA

- Compute a shape-aware coordinate system for each model
  - Origin: Centroid of all points
  - Axes: Directions in which the model varies most or least
- Transform the source to align its origin/axes with the target



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# Method 1: PCA

- Computing axes: Principal Component Analysis (PCA)

- Consider a set of points  $p_1, \dots, p_n$  with centroid location  $c$

- Construct matrix  $P$  whose  $i$ -th column is vector  $p_i - c$

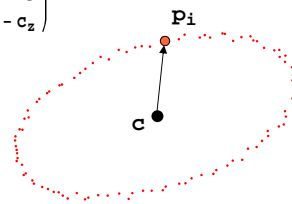
- 2D (2 by  $n$ ):  $P = \begin{pmatrix} p_{1x} - c_x & p_{2x} - c_x & \dots & p_{nx} - c_x \\ p_{1y} - c_y & p_{2y} - c_y & \dots & p_{ny} - c_y \end{pmatrix}$

- 3D (3 by  $n$ ):  $P = \begin{pmatrix} p_{1x} - c_x & p_{2x} - c_x & \dots & p_{nx} - c_x \\ p_{1y} - c_y & p_{2y} - c_y & \dots & p_{ny} - c_y \\ p_{1z} - c_z & p_{2z} - c_z & \dots & p_{nz} - c_z \end{pmatrix}$

- Build the covariance matrix:  $M = P \cdot P^T$

- 2D: a 2 by 2 matrix

- 3D: a 3 by 3 matrix



# Method 1: PCA

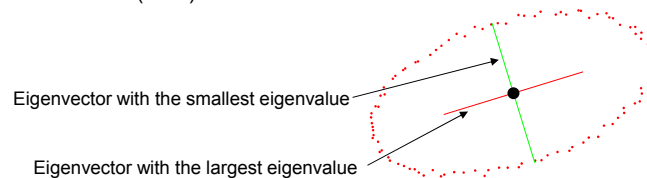
- Computing axes: Principal Component Analysis (PCA)

- Eigenvectors of the covariance matrix represent principal directions of shape variation

- The eigenvectors are un-singed and orthogonal (2 in 2D; 3 in 3D)

- Eigenvalues indicate amount of variation along each eigenvector

- Eigenvector with largest (smallest) eigenvalue is the direction where the model shape varies the most (least)





## Method 1: PCA

- PCA-based alignment

- Let  $c_S, c_T$  be centroids of source and target.
- First, translate source to align  $c_S$  with  $c_T$ :

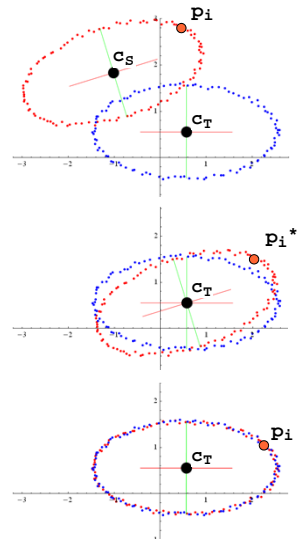
$$p_i^* = p_i + (c_T - c_S)$$

- Next, find rotation  $R$  that aligns two sets of PCA axes, and rotate source around  $c_T$ :

$$p_i' = c_T + R \cdot (p_i^* - c_T)$$

- Combined:

$$p_i' = c_T + R \cdot (p_i - c_S)$$



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## Method 1: PCA

- Finding rotation between two sets of **oriented** axes

- Let  $A, B$  be two matrices whose **columns** are the axes
  - The axes are orthogonal and normalized (i.e., both  $A$  and  $B$  are orthonormal)
- We wish to compute a rotation matrix  $R$  such that:

$$R \cdot A = B$$

- Notice that  $A$  and  $B$  are orthonormal, so we have:

$$R = B \cdot A^{-1} = B \cdot A^T$$

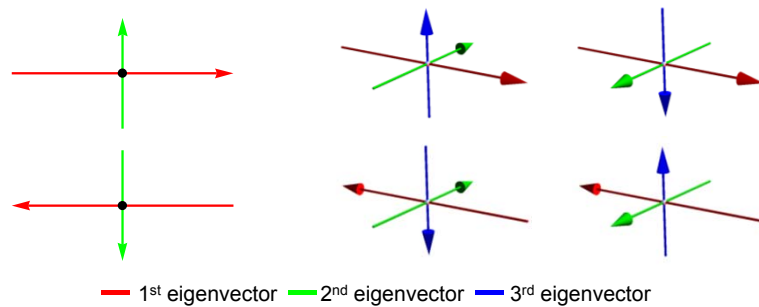
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## Method 1: PCA

- Assigning orientation to PCA axes
  - There are 2 possible orientation assignments in 2D
  - In 3D, there are 4 possibilities (observing the right-hand rule)



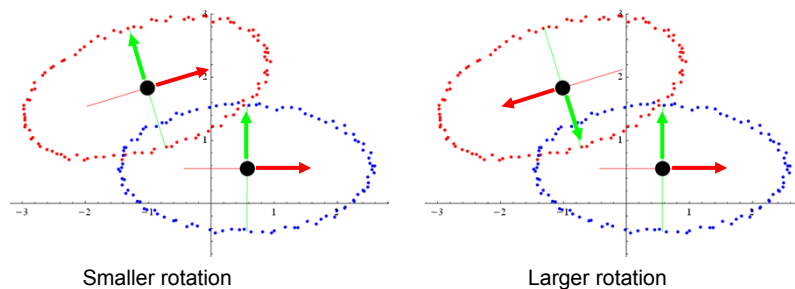
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## Method 1: PCA

- Finding rotation between two sets of **un-oriented** axes
  - Fix the orientation of the target axes.
  - For each orientation assignment of the source axes, compute  $R$
  - Pick the  $R$  with smallest rotation angle (by checking the trace of  $R$ )



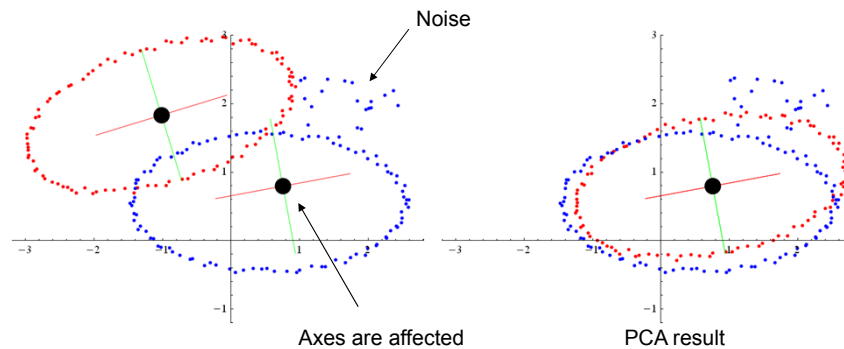
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# Method 1: PCA

- Limitations
  - Centroid and axes are affected by noise



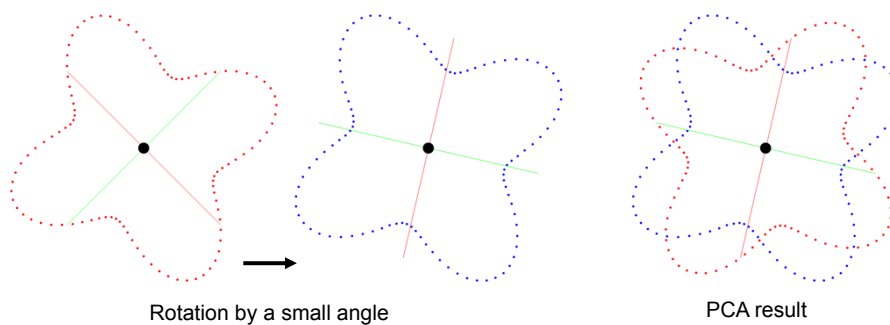
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# Method 1: PCA

- Limitations
  - Axes can be unreliable for circular objects
    - Eigenvalues become similar, and eigenvectors become unstable



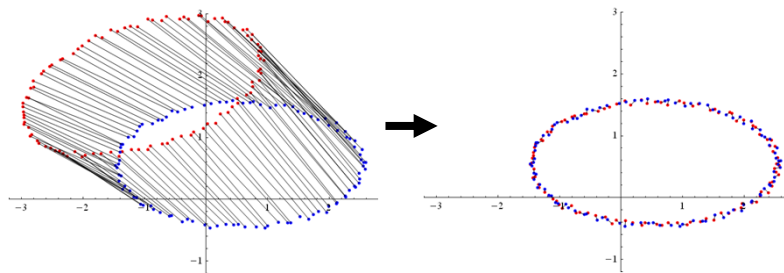
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## Method 2: SVD

- Optimal alignment between corresponding points
  - Assuming that for each source point, we know where the corresponding target point is



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## Method 2: SVD

- Formulating the problem
  - Source points  $p_1, \dots, p_n$  with centroid location  $c_s$
  - Target points  $q_1, \dots, q_n$  with centroid location  $c_T$ 
    - $q_i$  is the corresponding point of  $p_i$
  - After centroid alignment and rotation by some  $R$ , a transformed source point is located at:

$$p_i' = c_T + R \cdot (p_i - c_s)$$

- We wish to find the  $R$  that minimizes sum of pair-wise distances:

$$E = \sum_{i=1}^n \|q_i - p_i'\|^2$$

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## Method 2: SVD

- An equivalent formulation
  - Let  $P$  be a matrix whose  $i$ -th column is vector  $p_i - c_S$
  - Let  $Q$  be a matrix whose  $i$ -th column is vector  $q_i - c_T$
  - Consider the cross-covariance matrix:

$$\mathbf{M} = \mathbf{P} \cdot \mathbf{Q}^T$$

- Find the orthonormal matrix  $R$  that maximizes the trace:

$$\text{Tr} [\mathbf{R} \cdot \mathbf{M}]$$

## Method 2: SVD

- Solving the minimization problem
  - Singular value decomposition (SVD) of an  $m$  by  $m$  matrix  $M$ :

$$\mathbf{M} = \mathbf{U} \cdot \mathbf{W} \cdot \mathbf{V}^T$$

- $U, V$  are  $m$  by  $m$  orthonormal matrices (i.e., rotations)
- $W$  is a diagonal  $m$  by  $m$  matrix with non-negative entries
- The orthonormal matrix (rotation)  $\mathbf{R} = \mathbf{V} \cdot \mathbf{U}^T$  is the  $R$  that maximizes the trace  $\text{Tr} [\mathbf{R} \cdot \mathbf{M}]$
- SVD is available in *Mathematica* and many Java/C++ libraries

## Method 2: SVD

- SVD-based alignment: summary

- Forming the cross-covariance matrix

$$\mathbf{M} = \mathbf{P} \cdot \mathbf{Q}^T$$

- Computing SVD

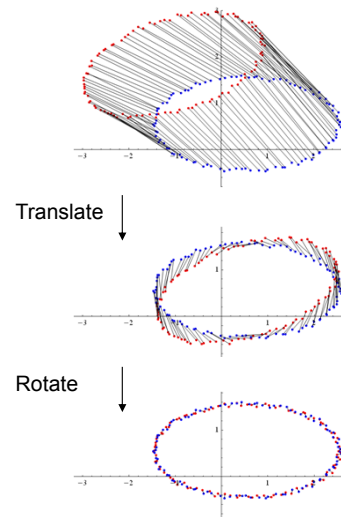
$$\mathbf{M} = \mathbf{U} \cdot \mathbf{W} \cdot \mathbf{V}^T$$

- The rotation matrix is

$$\mathbf{R} = \mathbf{V} \cdot \mathbf{U}^T$$

- Translate and rotate the source:

$$\mathbf{p}_i' = \mathbf{c}_T + \mathbf{R} \cdot (\mathbf{p}_i - \mathbf{c}_S)$$



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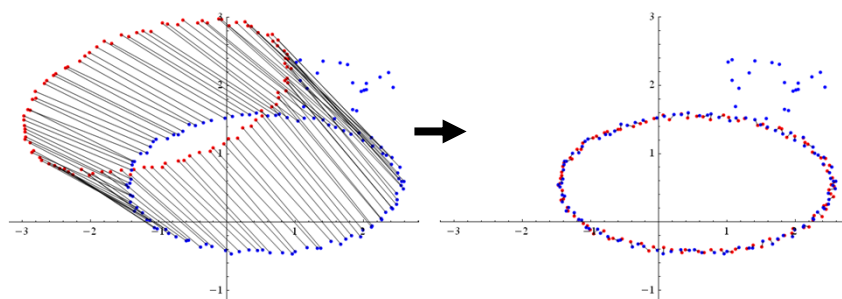
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## Method 2: SVD

- Advantage over PCA: more stable

- As long as the correspondences are correct



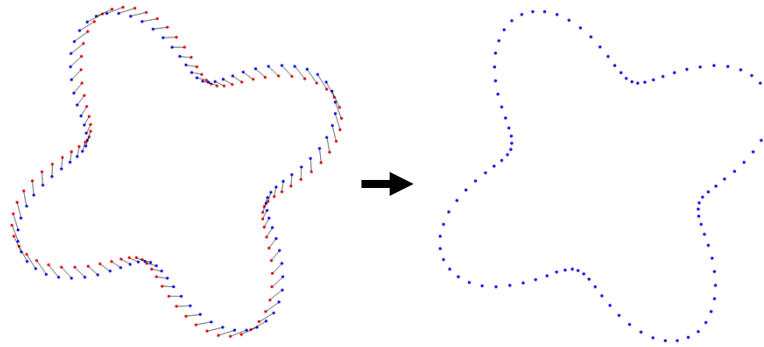
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## Method 2: SVD

- Advantage over PCA: more stable
  - As long as the correspondences are correct



## Method 2: SVD

- Limitation: requires accurate correspondences
  - Which are usually not available

## Method 3: ICP

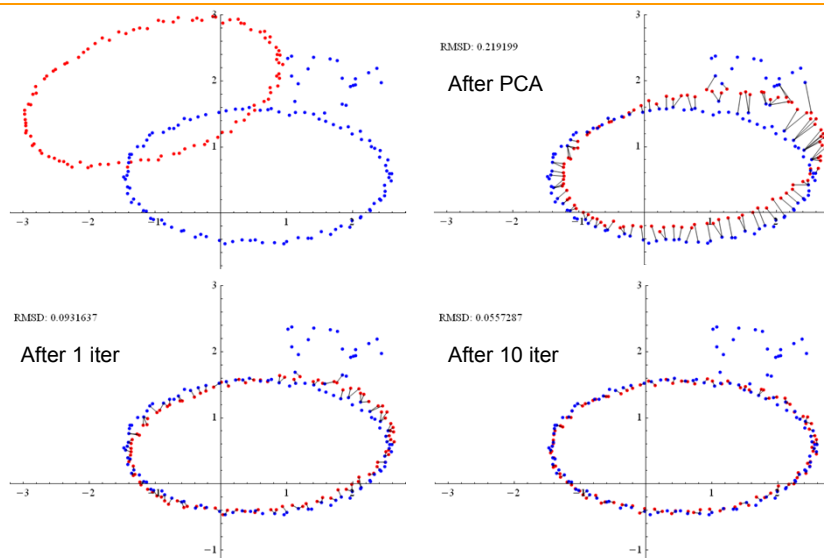
- The idea
  - Use PCA alignment to obtain initial guess of correspondences
  - Iteratively improve the correspondences after repeated SVD
- Iterative closest point (ICP)
  - 1. Transform the source by PCA-based alignment
  - 2. For each transformed source point, assign the closest target point as its corresponding point. Align source and target by SVD.
    - Not all target points need to be used
  - 3. Repeat step (2) until a termination criteria is met.

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## ICP Algorithm



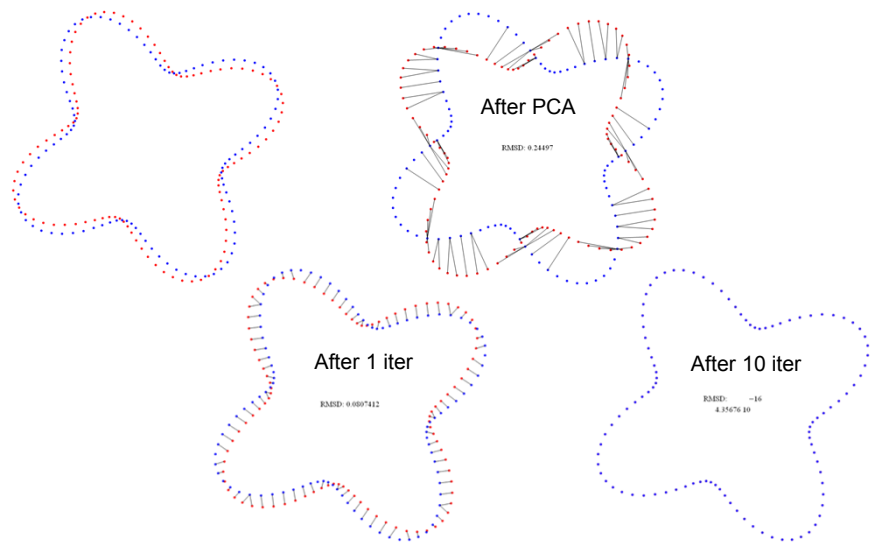
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# ICP Algorithm



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# ICP Algorithm

- Termination criteria
  - A user-given maximum iteration is reached
  - The **improvement** of fitting is small

- Root Mean Squared Distance (RMSD):

$$\sqrt{\frac{\sum_{i=1}^n \|q_i - p_i\|^2}{n}}$$

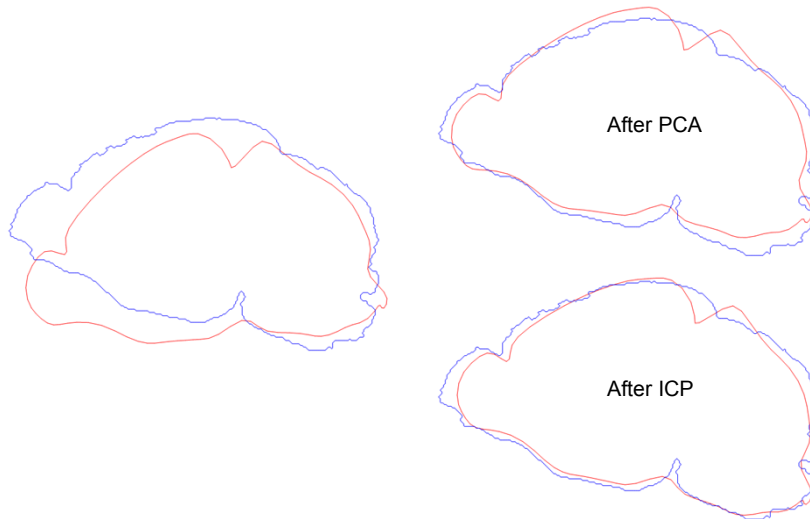
- Captures average deviation in all corresponding pairs
- Stops the iteration if the difference in RMSD before and after each iteration falls beneath a user-given threshold

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## More Examples

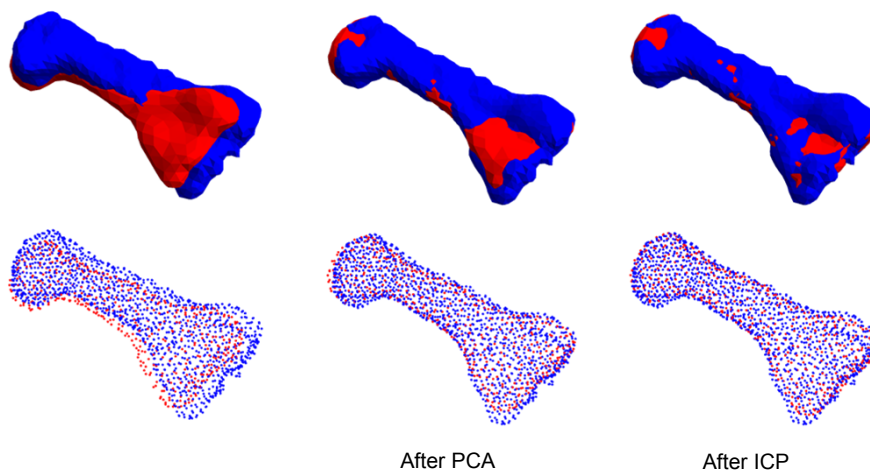


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## More Examples



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