



# Non-Rigid Surface Correspondence

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# Goal

## Find maps between surfaces

- Non-rigid
- Bijective
- Smooth
- Shape preserving
- Automatic
- Efficient computation
- Provide metric
- Semantic alignment

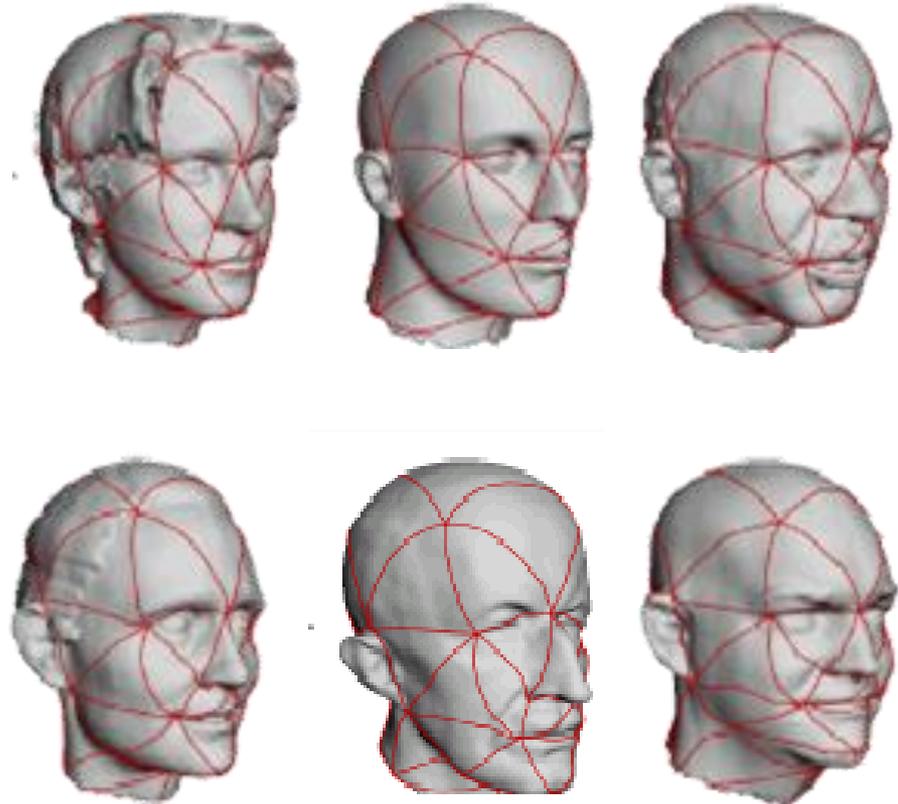




# Motivating Applications

Finding corresponding points on surfaces enables ...

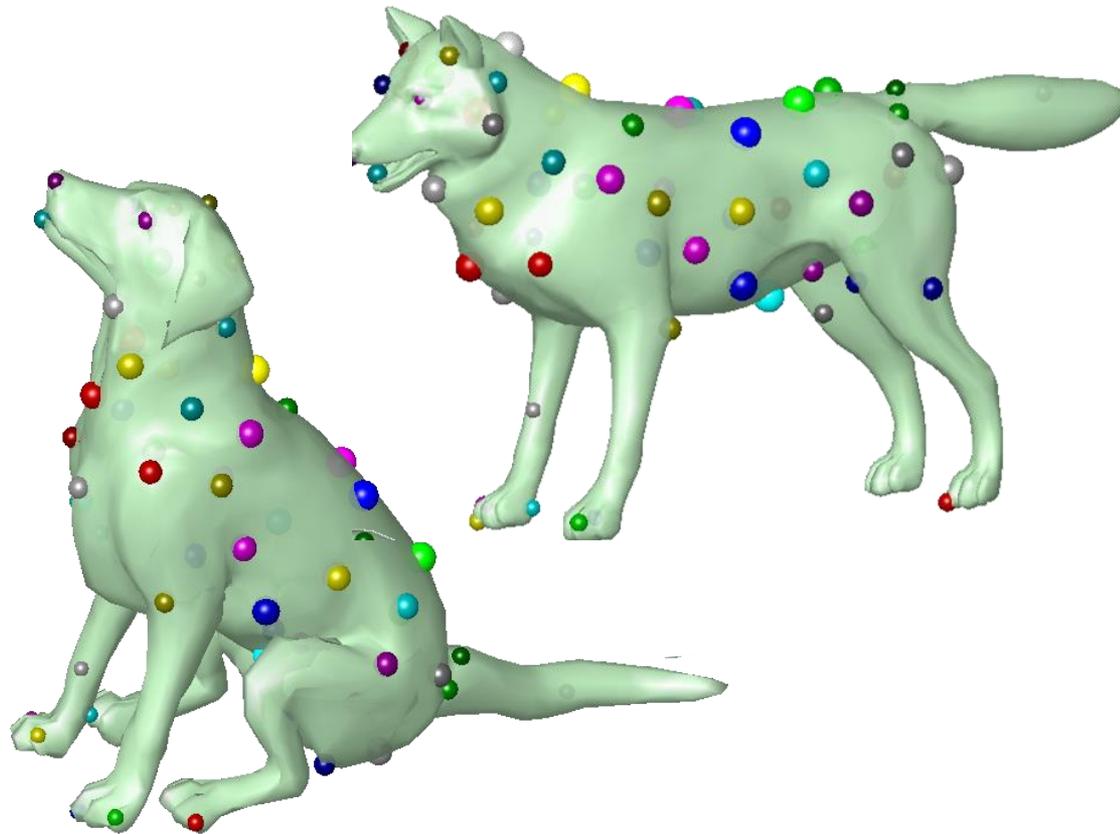
- Surface comparison
- Collection analysis
- Property transfer
- Morphing
- etc.





# Problem 1

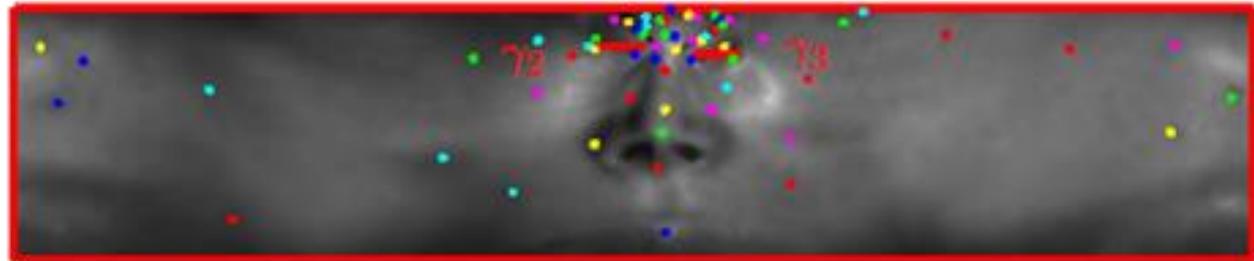
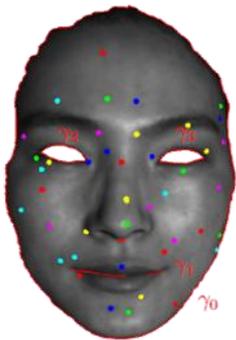
Find a sparse set of feature correspondences



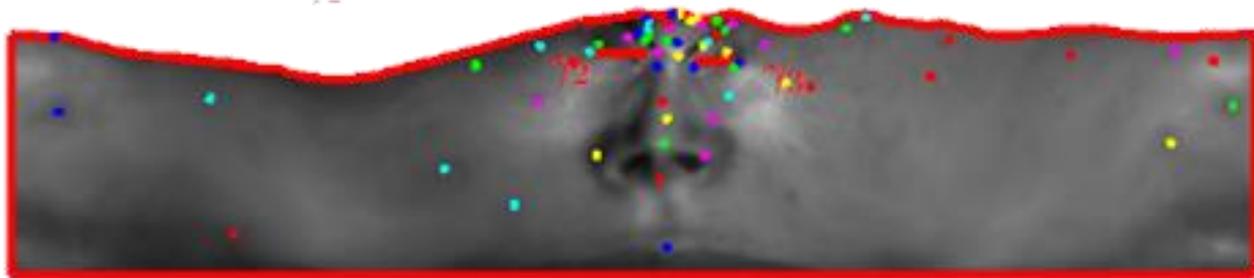
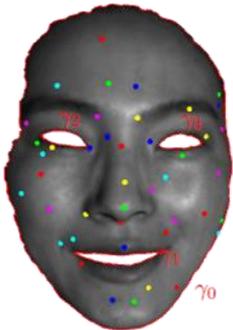


# Problem 2

Compute a dense map from  
a sparse set of feature correspondences



Zeng et al., 2008]



Least Squares Conformal Map  
(preserve angles as best as possible)



# Outline

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Introduction

Some surface mapping algorithms

- Feature correspondence search
- High-dimensional embedding
- Möbius transformations
- Blended maps

Example Application

Conclusion

Future work



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Some surface mapping algorithms

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Example Application

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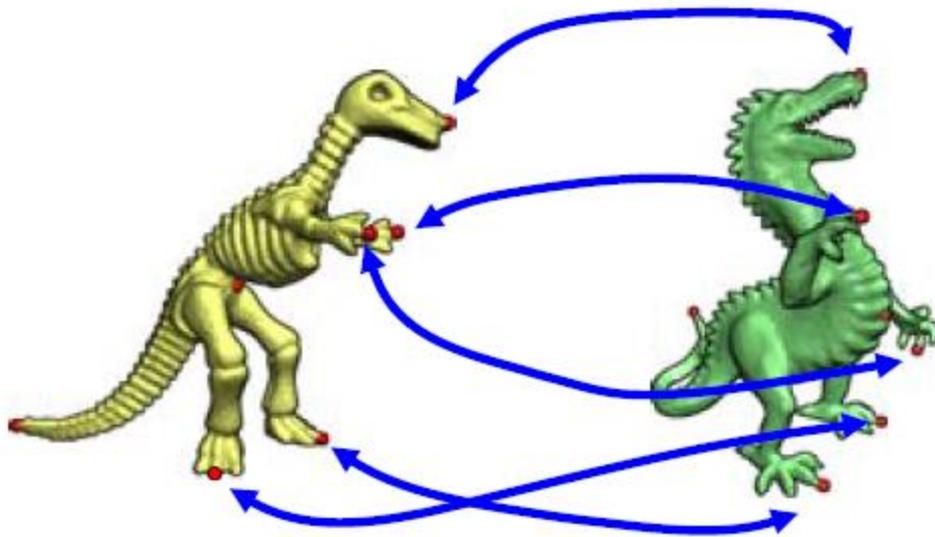
Future work

# Feature Correspondence Search



For each coarse set of feature correspondences:

- Measure the deformation required to align them
  - ... maybe by solving problem 2
- Remember the one with least deformation

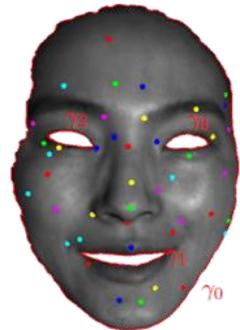
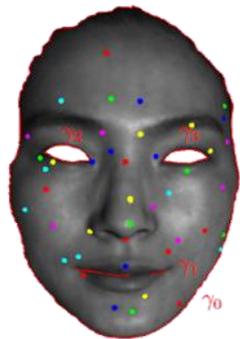




# Feature Correspondence Search

Measures of distortion:

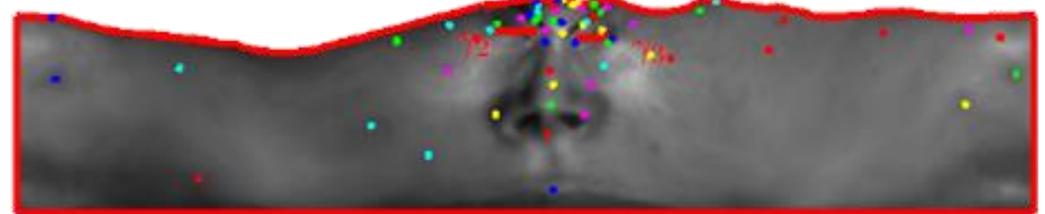
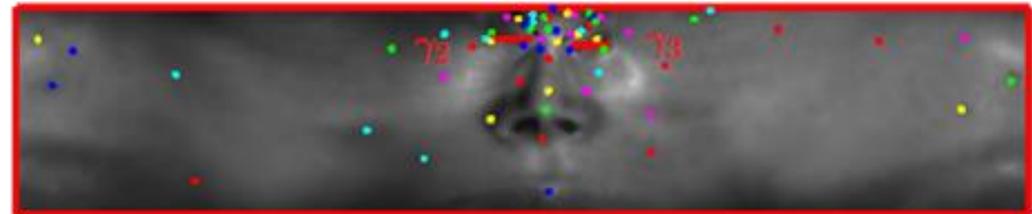
- Differences in geodesic distances
- Differences in conformal factors (angles)
- etc.



Feature points



Branch and bound  
Priority-driven search  
etc.



Least squares conformal map  
aligning corresponding feature points



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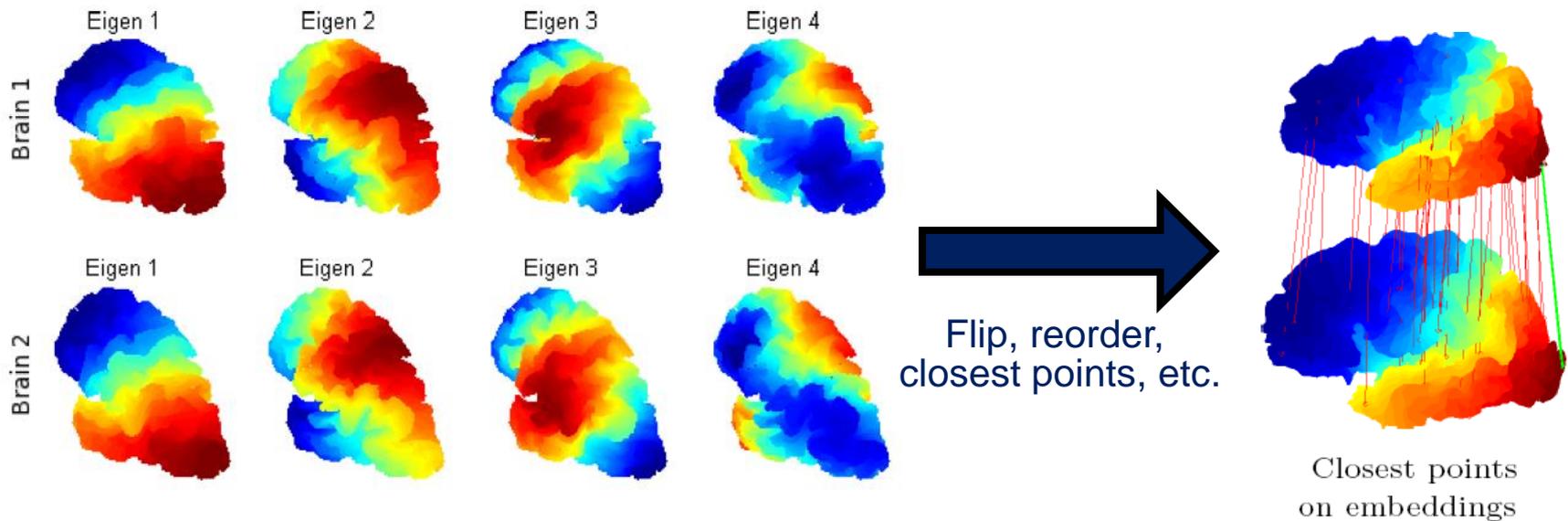
Conclusion

Future work



# High-Dimensional Embedding

Find nearest neighbors after spectral embedding



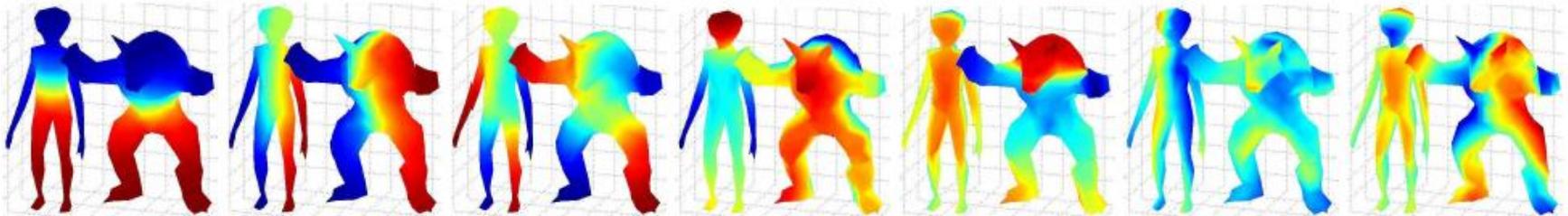
Eigenfunctions of the Laplacian

[Lombaert et al. 2011]



# High-Dimensional Embedding

Find nearest neighbors after spectral embedding



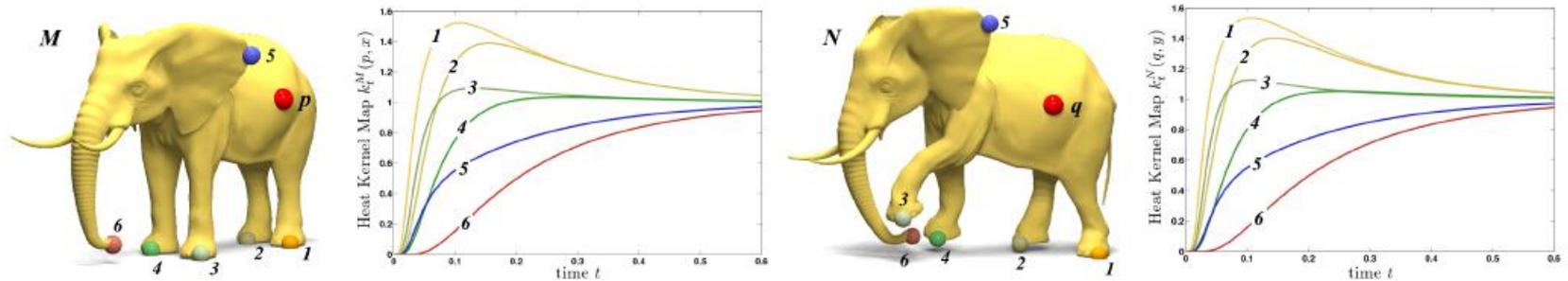
Eigenfunctions of the Laplacian

[Lombaert et al. 2011]



# High-Dimensional Embedding

Find nearest neighbors after heat kernel embedding implied by a single point correspondence



Heat Kernel Map  
[Ovsjanikov et al. 2010]



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- Blended maps

Example Application

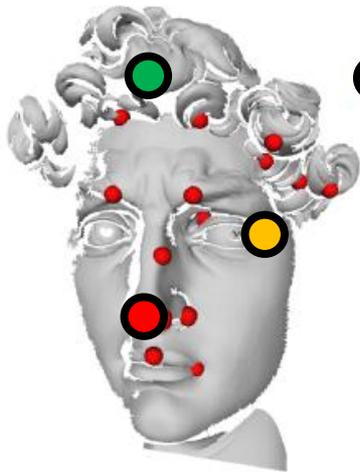
Conclusion

Future work

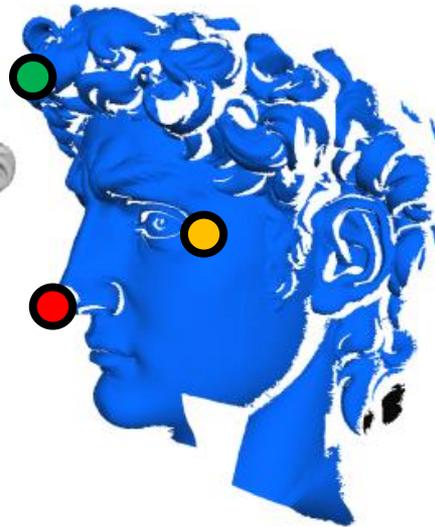


# Möbius Transformations

It would be nice to search a low-dimensional space of transformations to align non-rigid surfaces ...



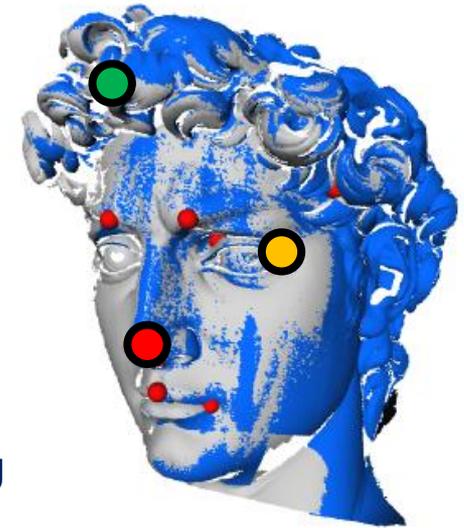
Scan A



Scan B



RANSAC  
Hough transform  
Geometric hashing  
etc.

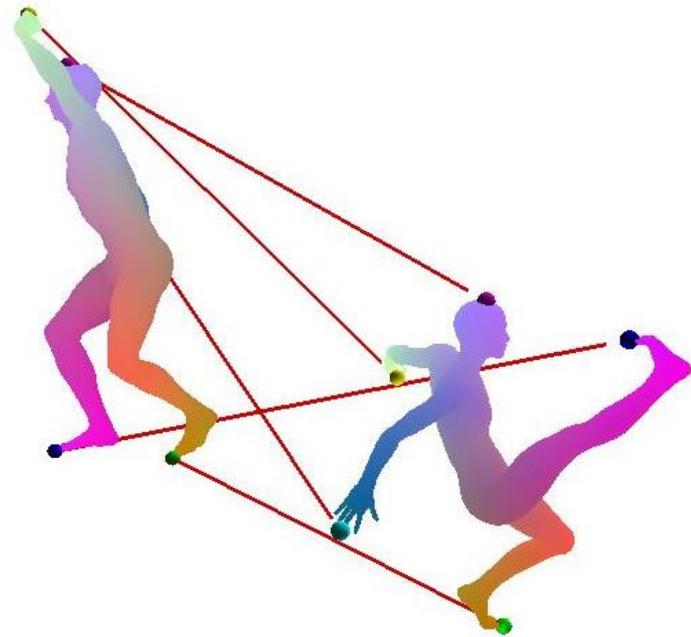
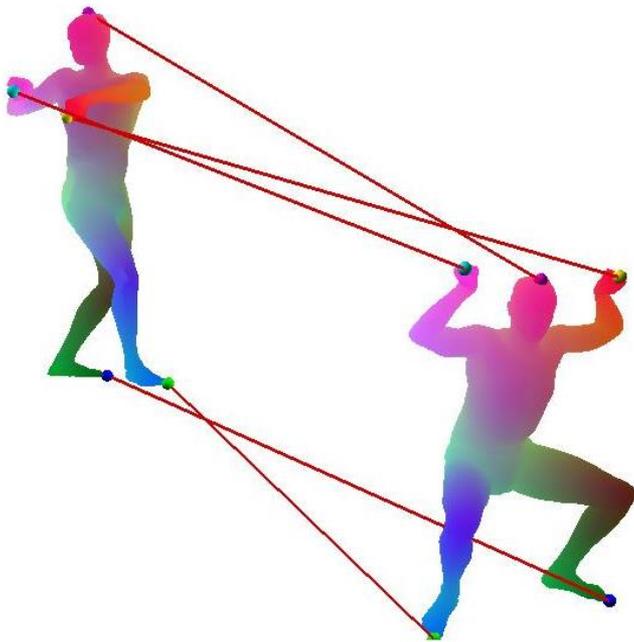


Best  
Alignment



# Key Observation

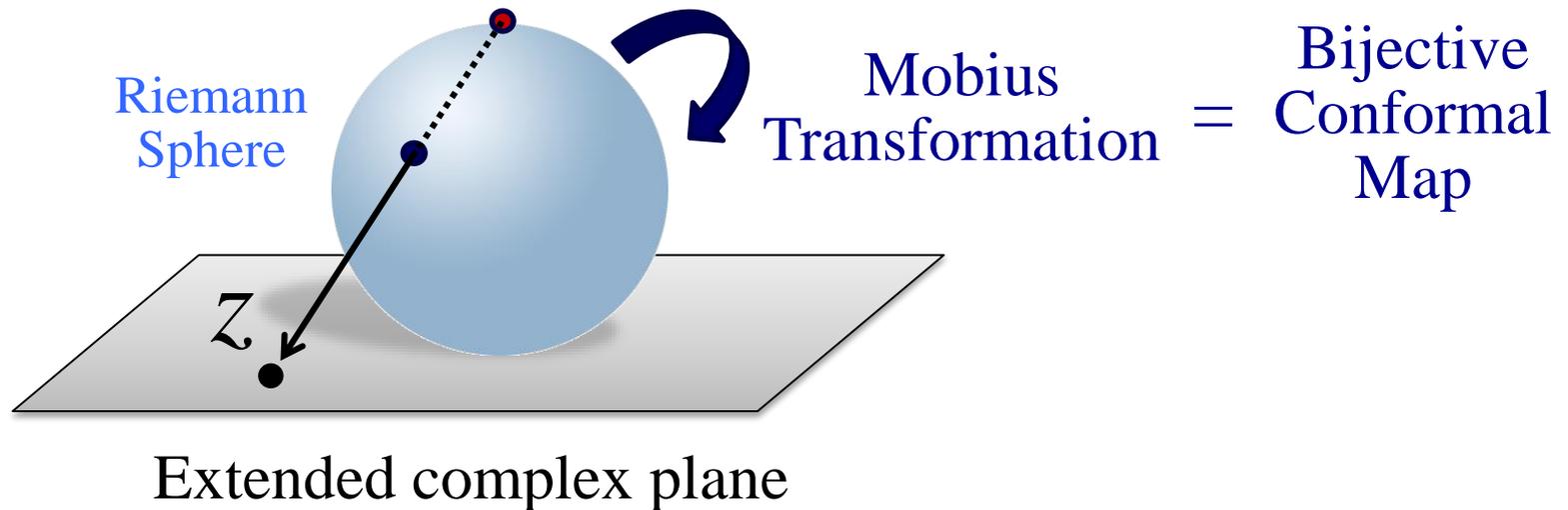
The Möbius group provides a low-dimensional space to search efficiently for the “best” conformal map between genus zero surfaces





# Möbius Transformations I

Möbius transformations are a group of functions on the extended complex plane that represent bijective, conformal maps





# Möbius Transformations II

Möbius transformations are simple rational functions:

$$f(z) = \frac{az + b}{cz + d}, \quad ad - bc \neq 0, \quad a, b, c, d \in \mathbb{C}$$

They have only six degrees of freedom  
(they can be computed analytically  
from three point correspondences)

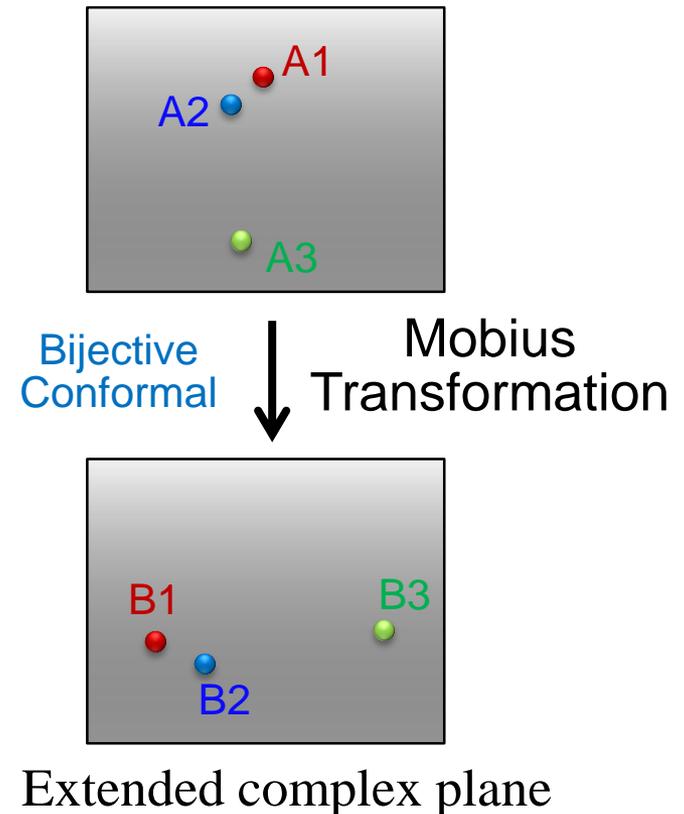


$$f(z_i) = y_i, i = 1, 2, 3$$



# Möbius Transformations III

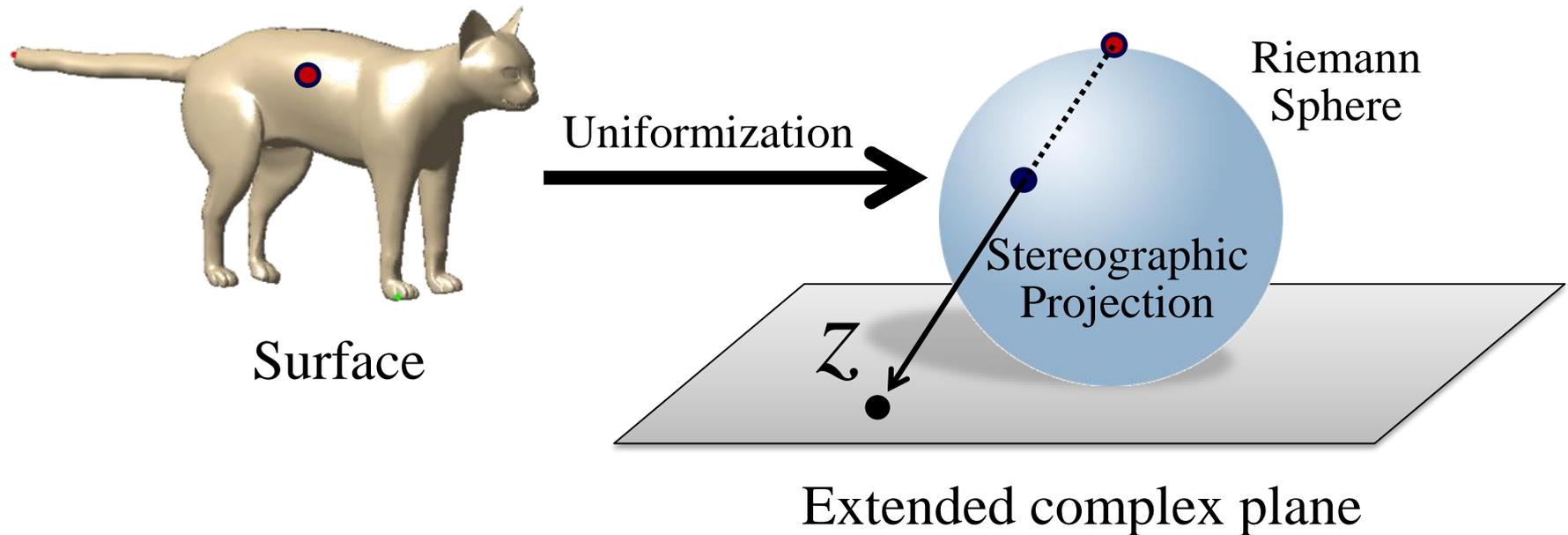
Therefore, any three point correspondences define a bijective, conformal map from the extended complex plane onto itself





# Möbius Transformations IV

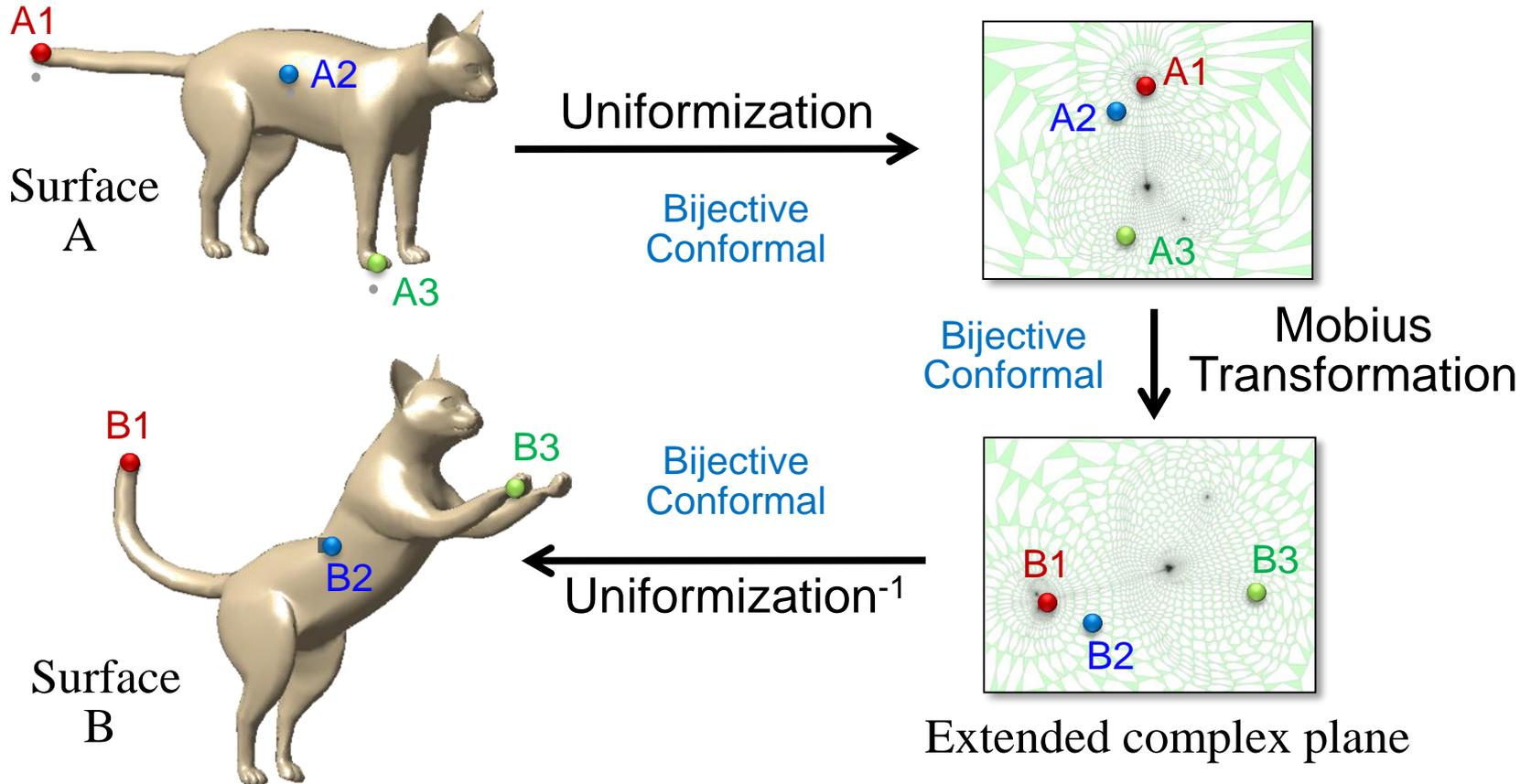
Since every genus zero surface can be mapped conformally onto the extended complex plane (Riemann sphere), ...





# Möbius Transformations V

Any three point correspondences define a bijective, conformal map between genus zero surfaces





# Möbius Transformations VI

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We can search for the “lowest distortion” bijective, conformal map between genus zero surfaces using algorithms that sample triplets of correspondences (e.g., RANSAC, Hough transform, etc.)

Polynomial-time algorithm  
for non-rigid surface mapping



# Surface Mapping Algorithm

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Example: RANSAC algorithm

For  $i = 1$  to  $\sim N^3$

Sample three points  $(A_1, A_2, A_3)$  on surface A

Sample three points  $(B_1, B_2, B_3)$  on surface B

Compute conformal map  $M: (A_1, A_2, A_3) \rightarrow (B_1, B_2, B_3)$

Remember M if distortion is smallest



# Surface Mapping Algorithm

Example: RANSAC algorithm

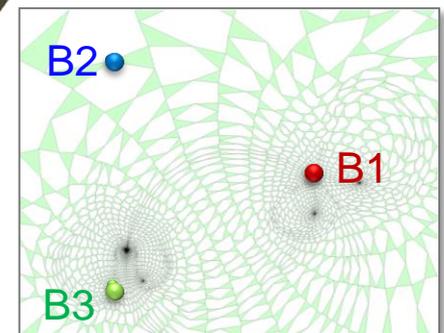
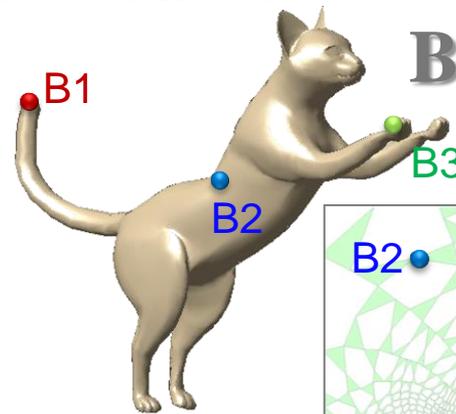
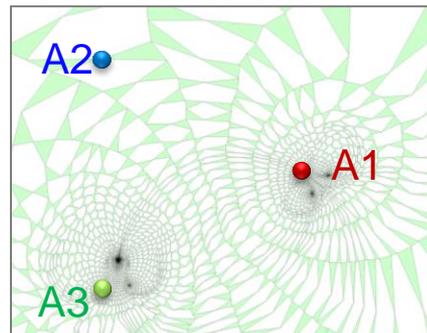
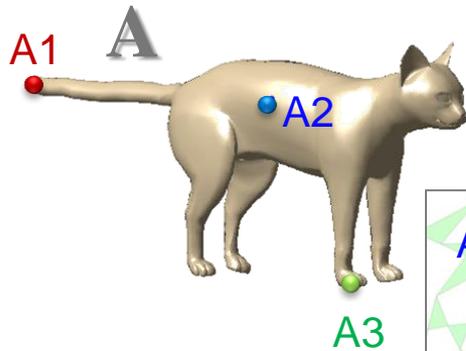
For  $i = 1$  to  $\sim N^3$

Sample three points (A1,A2,A3) on surface A

Sample three points (B1,B2,B3) on surface B

Compute conformal map  $M: (A1,A2,A3) \rightarrow (B1,B2,B3)$

Remember M if distortion is smallest



Measure distortion by relative change of area  
(deviation from isometry)



# Surface Mapping Algorithm

Example: RANSAC algorithm

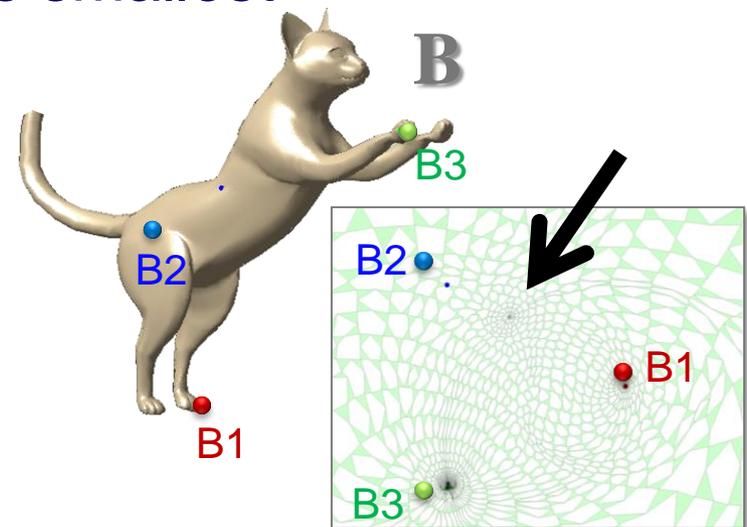
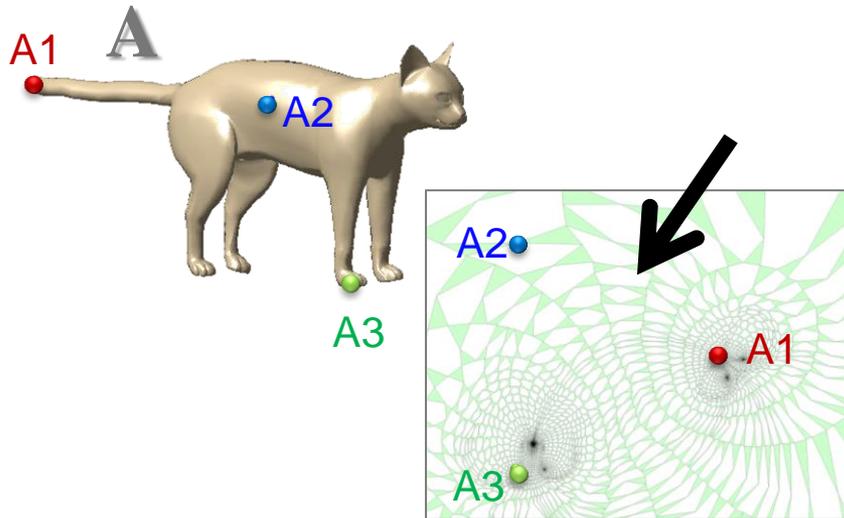
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# Surface Mapping Algorithm

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RANSAC algorithm properties:

- Non-rigid
- Bijective
- Smooth
- Shape preserving
- Automatic
- Efficient computation
- Provides metric
- Semantic alignment?



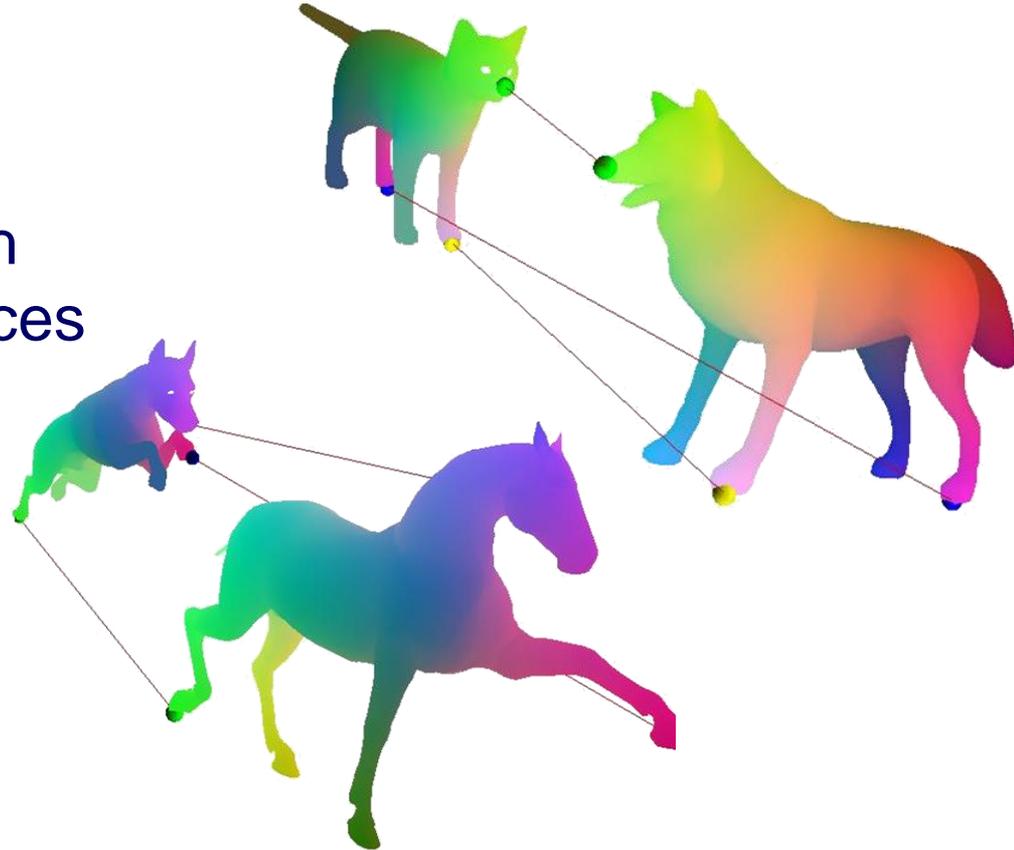
# Experimental Results

## Data:

- 51 pairs of meshes representing animals from TOSCA and SHREC Watertight data sets

## Methodology:

- Predict surface maps
- Compare to ground truth semantic correspondences

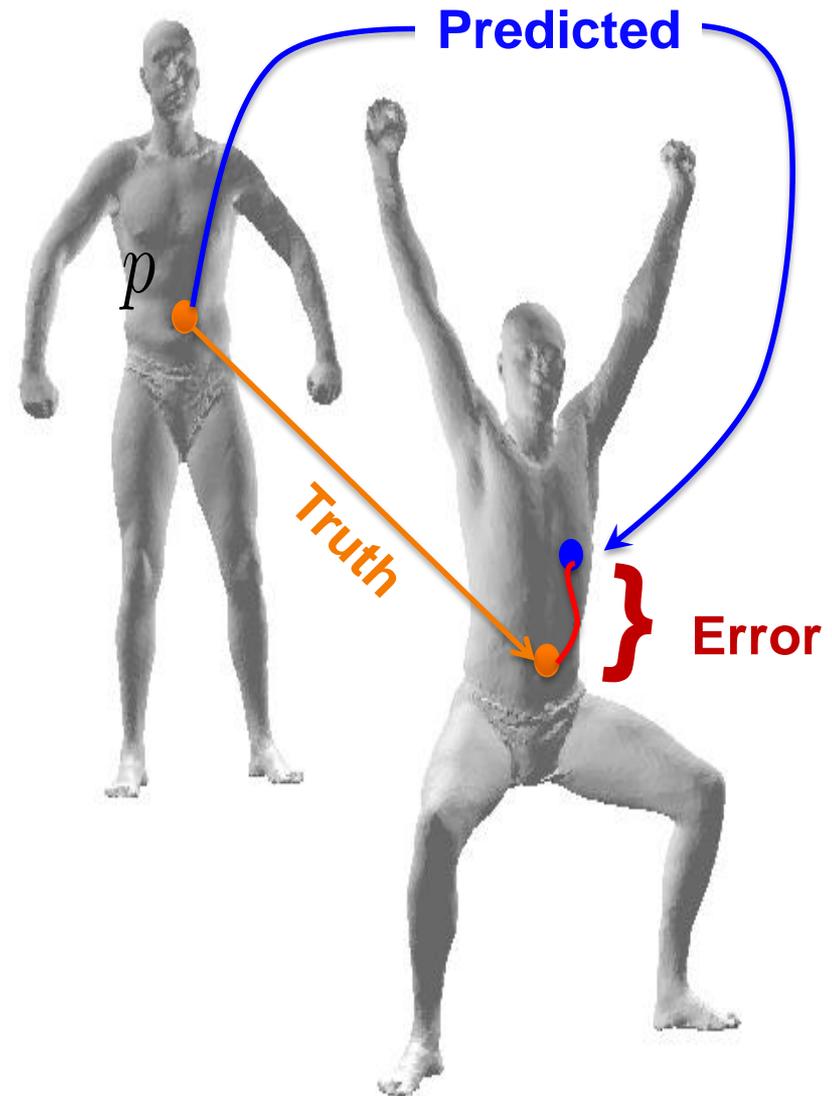




# Experimental Results

## Evaluation:

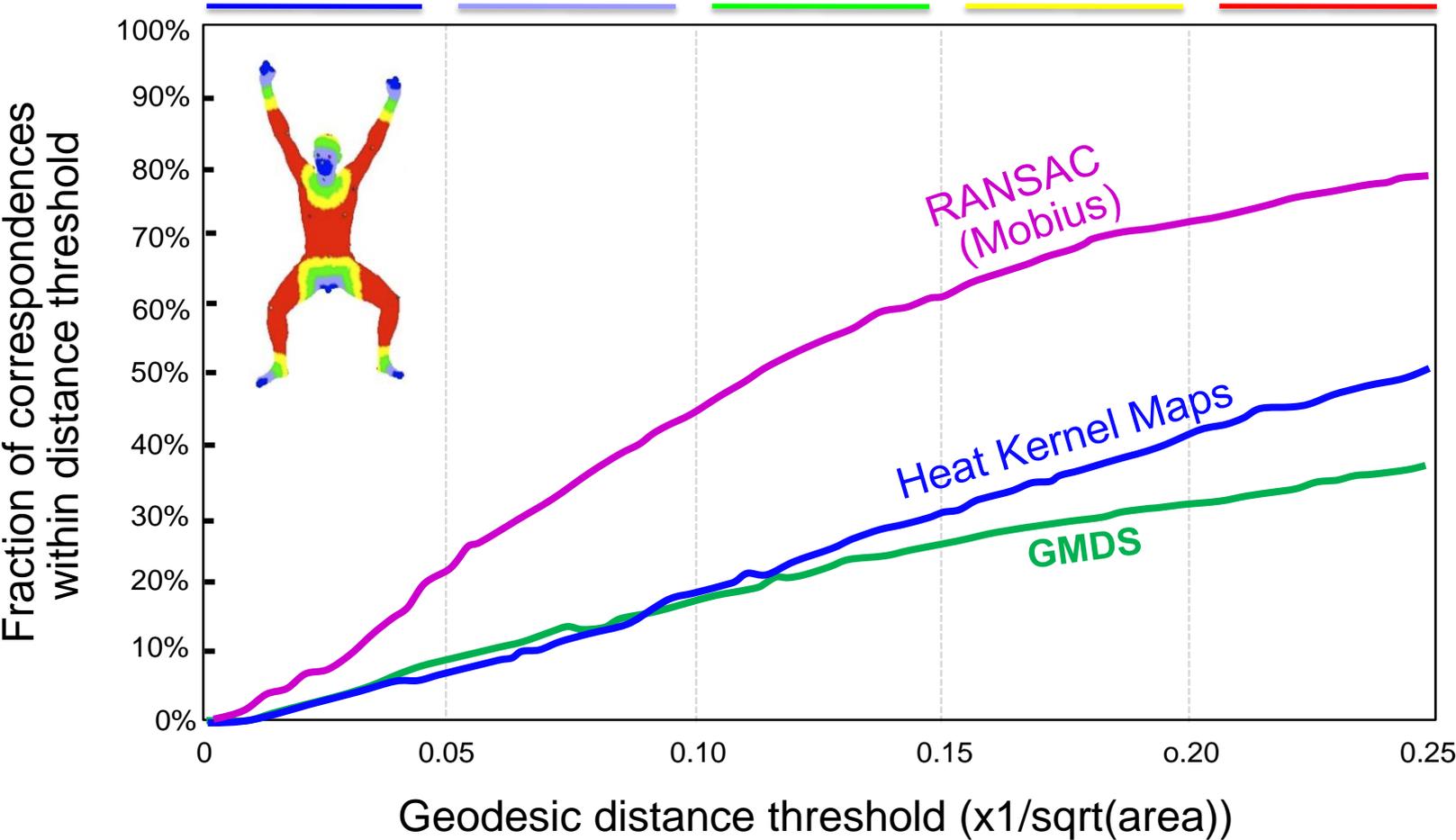
1. For every point with a ground truth correspondence, measure geodesic distance between predicted correspondence and ground truth correspondence
2. Plot fraction of points within geodesic error threshold





# Experimental Results

Results:





# Outline

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Some surface mapping algorithms

- Feature correspondence search
- High-dimensional embedding
- Möbius transformations
- **Blended maps**

Example Application

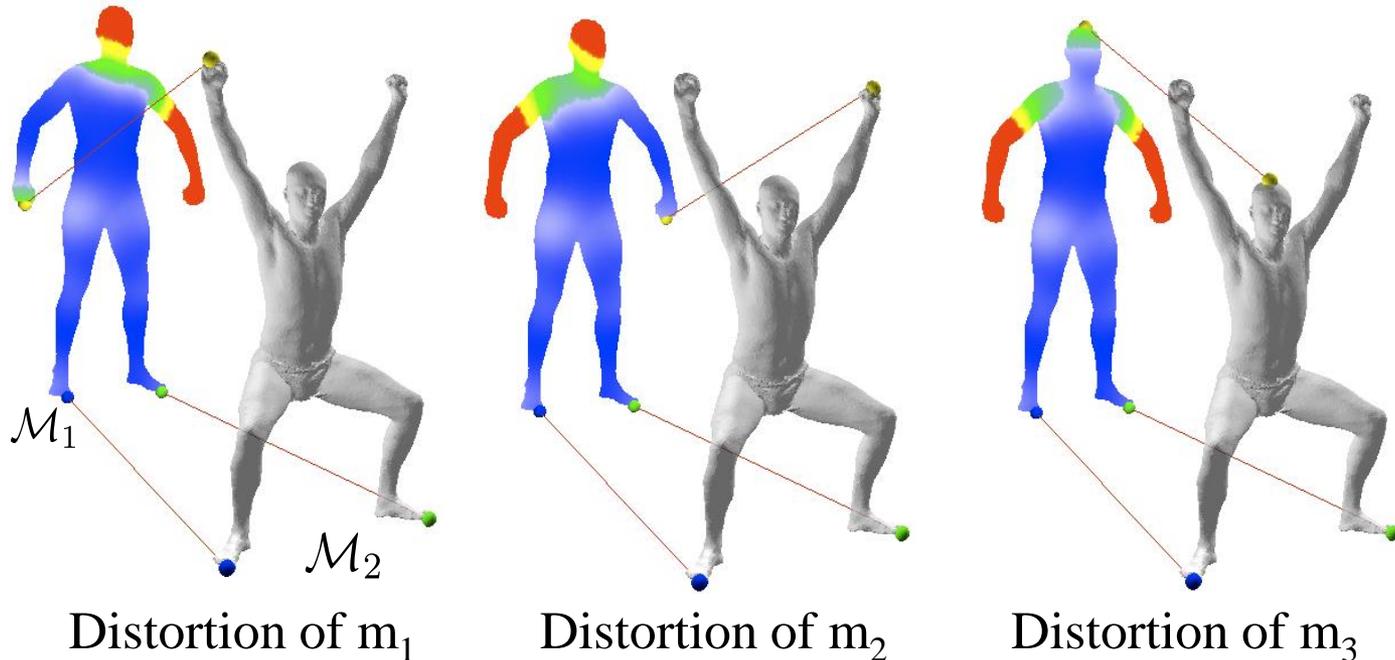
Conclusion

Future work



# Blended Maps

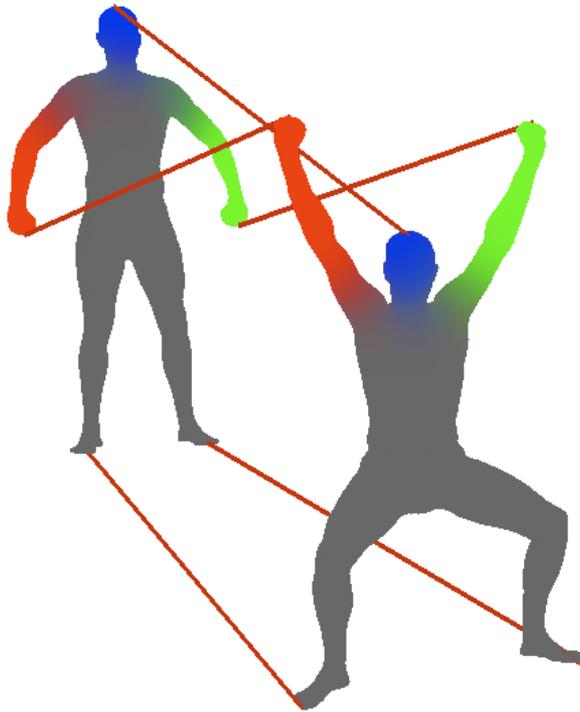
For significantly different surfaces, no single conformal map provides low distortion everywhere



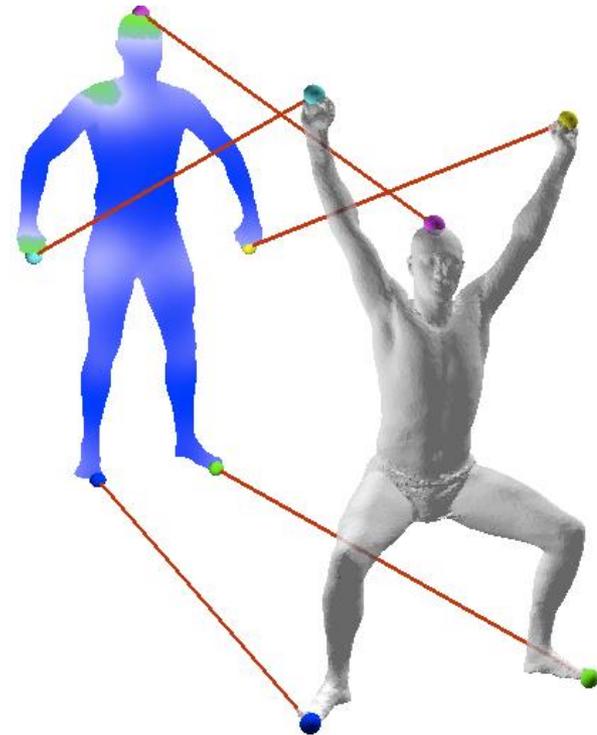


# Blended Maps

Idea: blend conformal maps with smooth weights



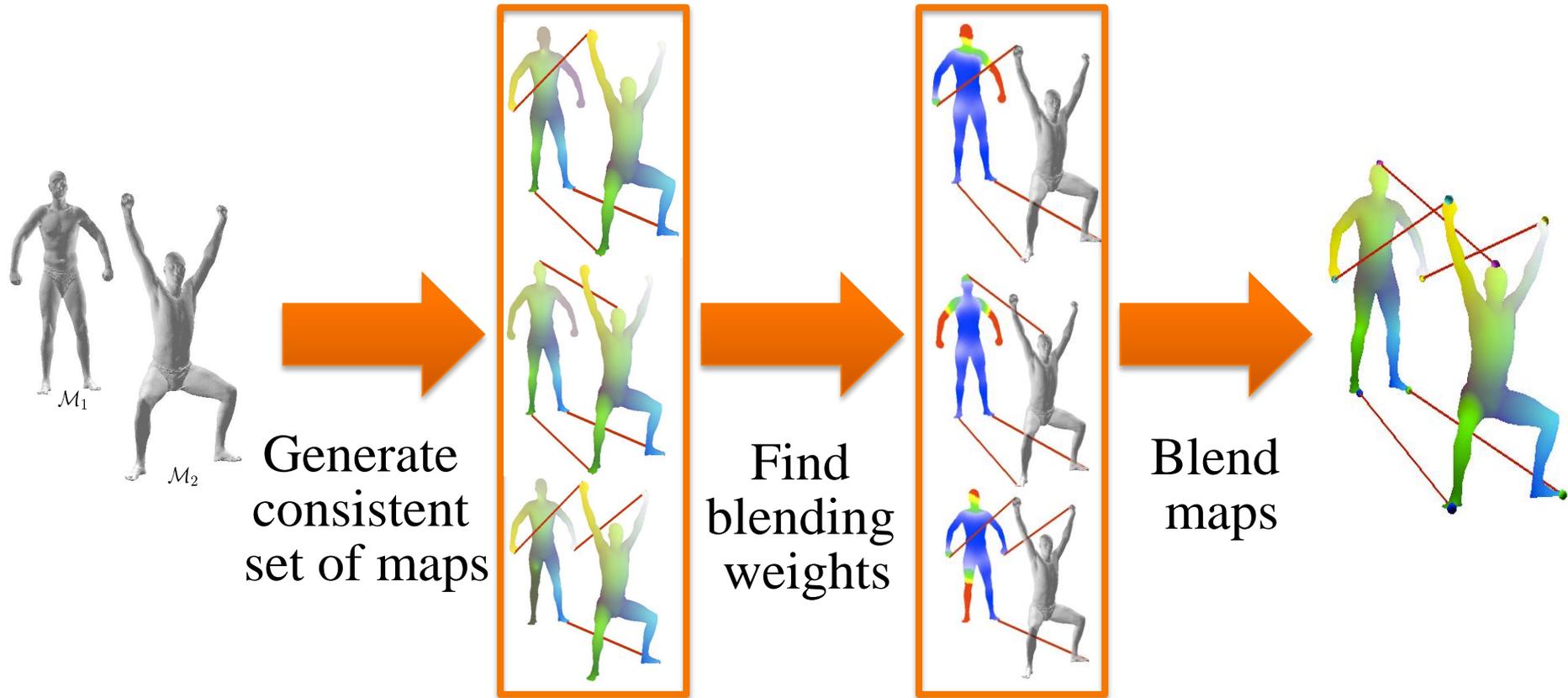
Blending Weights  
for  $m_1$ ,  $m_2$ , and  $m_3$



Distortion of the  
Blended Map



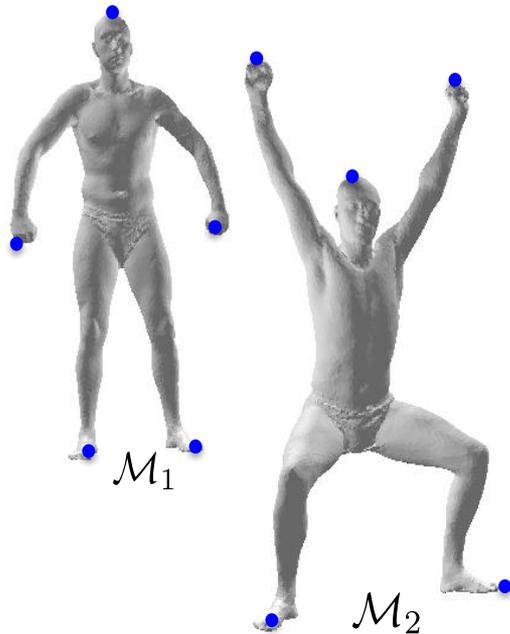
# Computing Blended Maps



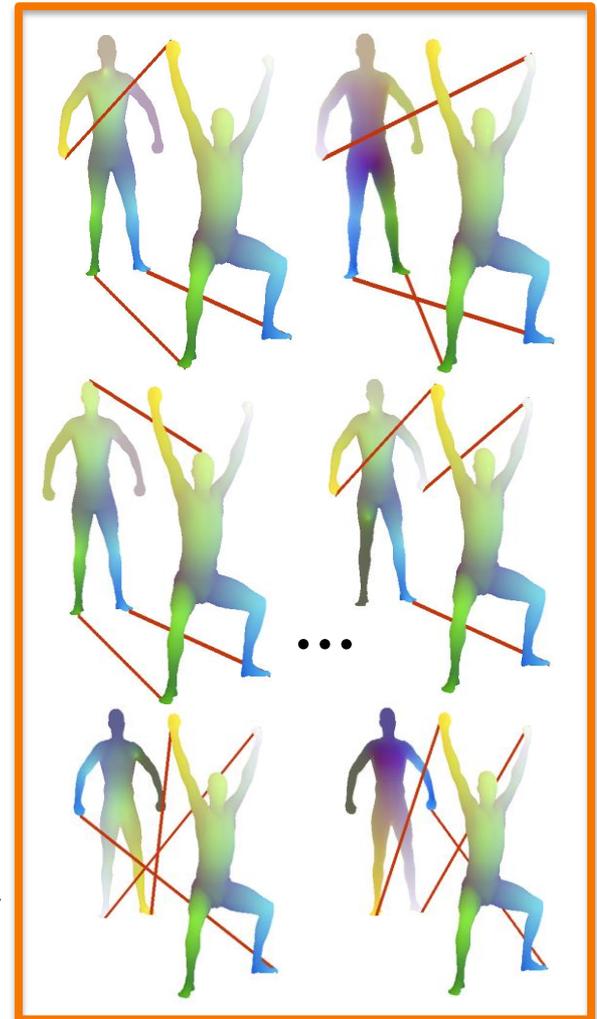


# Computing Blended Maps

1. Generate candidate maps by enumerating triplets of feature correspondences



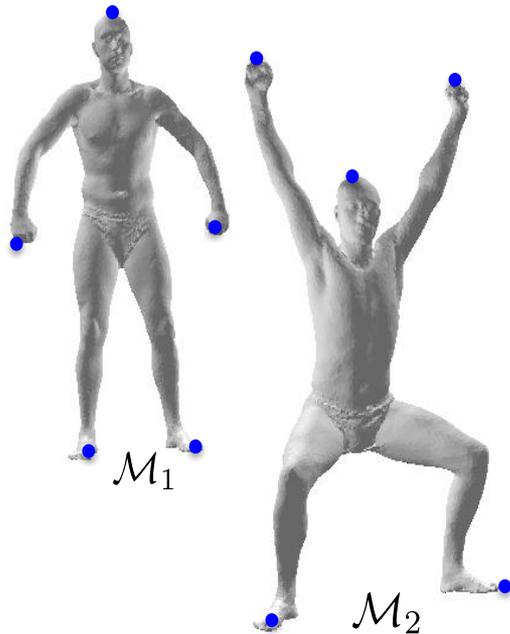
Set of candidate maps



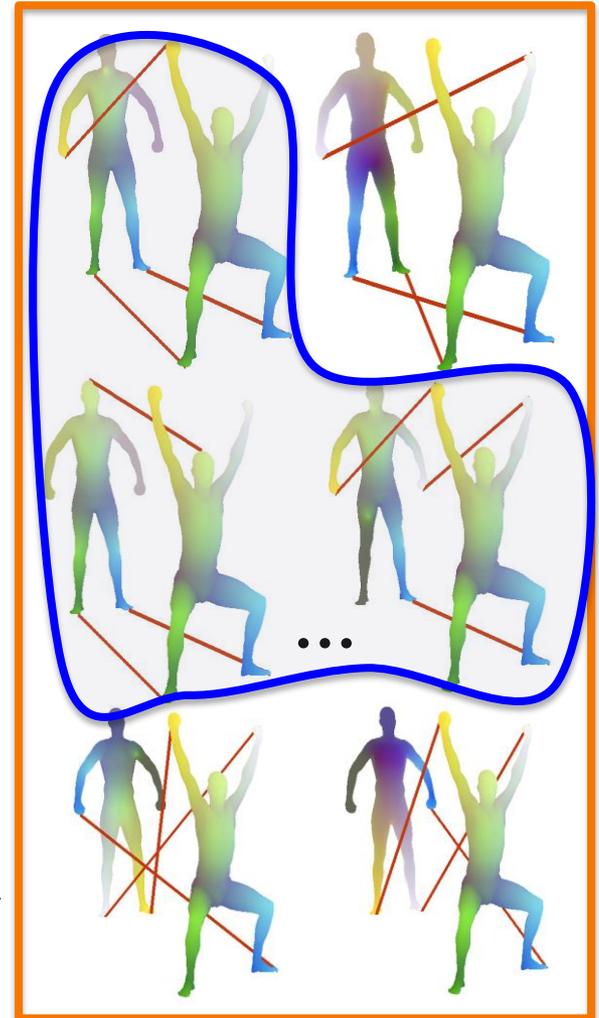


# Computing Blended Maps

2. Select consistent set of low-distortion candidate maps



Set of candidate maps

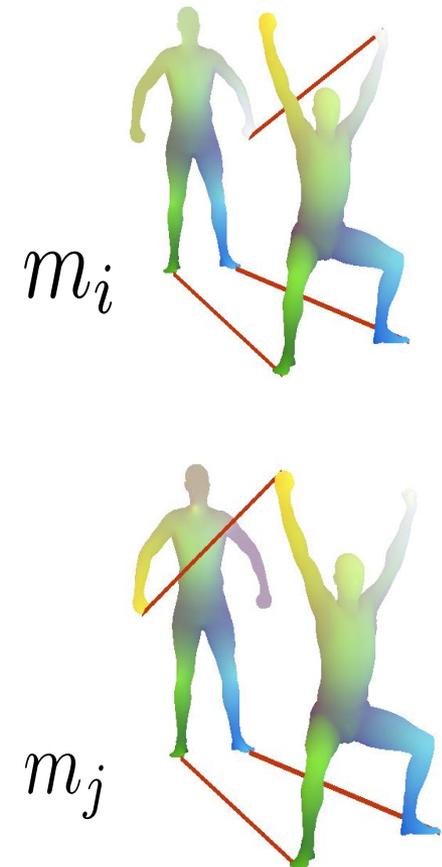
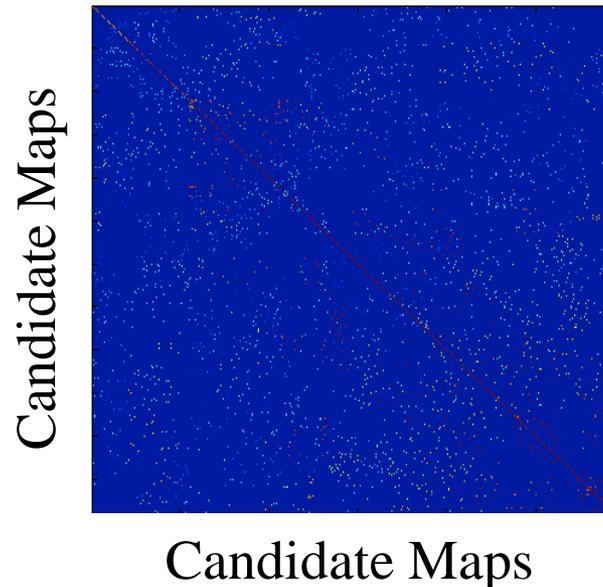




# Computing Blended Maps

2a. Define a matrix  $\mathbf{B}$  where every entry  $(i,j)$  indicates the distortion of  $m_i$  and  $m_j$  and their pairwise similarity  $S_{i,j}$

$$\mathbf{B}_{i,j} = \int_{M_1} c_i(p)c_j(p)S_{i,j}(p)dA(p)$$

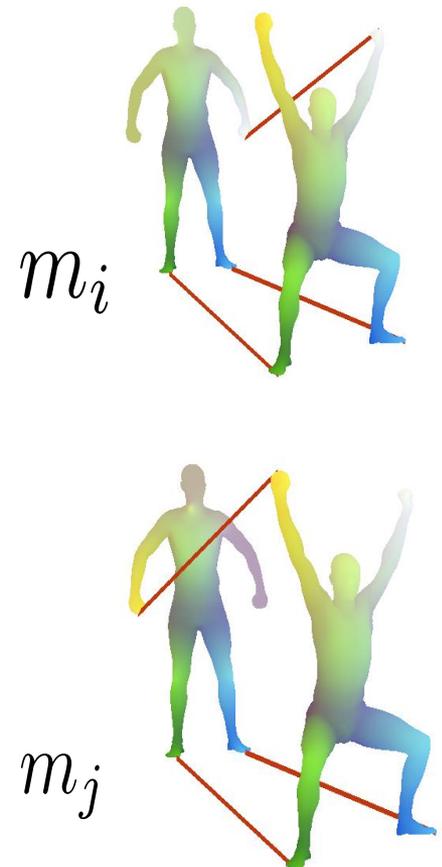
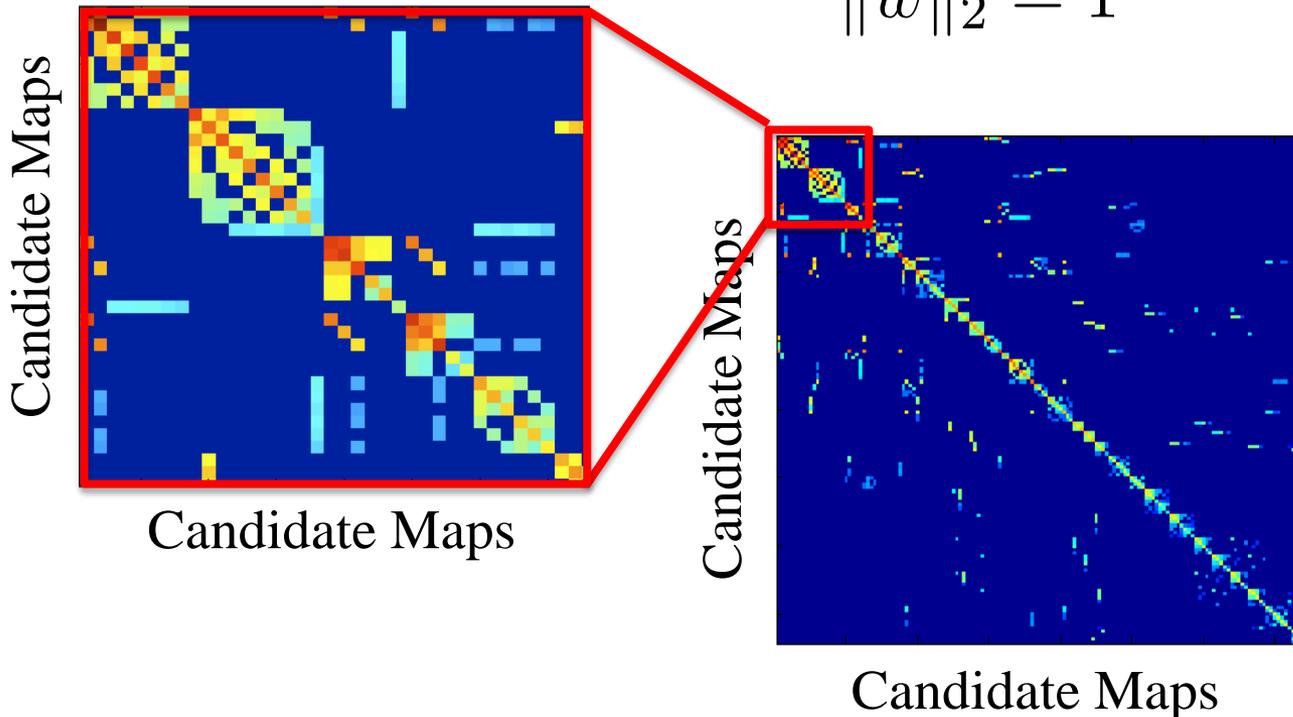




# Computing Blended Maps

2b. Find block of consistent, low-distortion maps using top eigenvector(s) of  $\mathbf{B}$

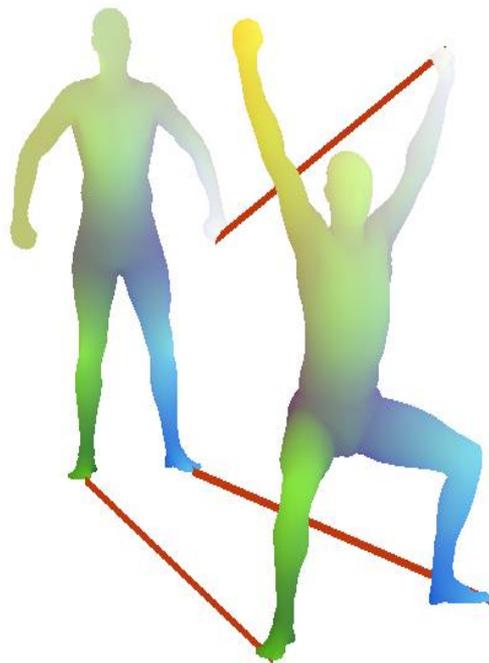
$$E_{\mathcal{M}}(\vec{w}) = \vec{w}^T \mathbf{B} \vec{w}$$
$$\|\vec{w}\|_2 = 1$$



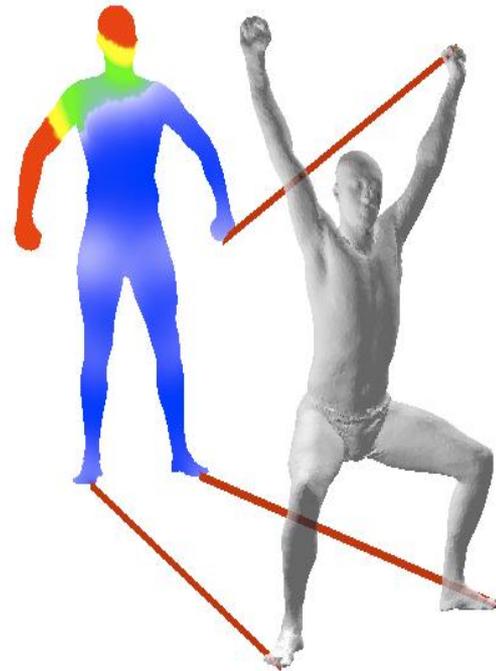


# Computing Blended Maps

3. Compute blending weight  $c_i(p)$  for every map  $m_i$  at every point  $p$  based on distortion of  $m_i$  at  $p$



Candidate Map



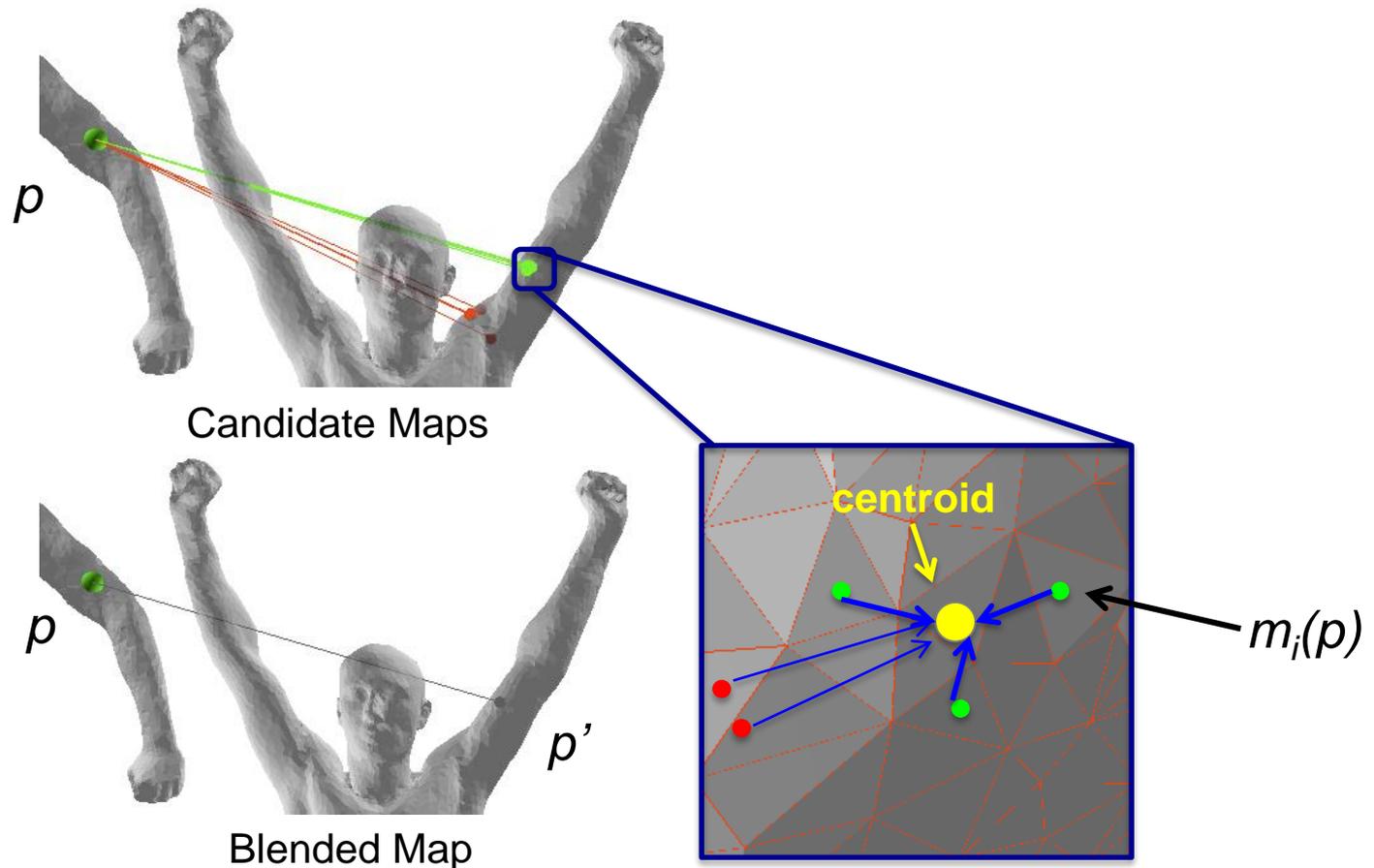
Blending Weight

$$c_i(p)$$

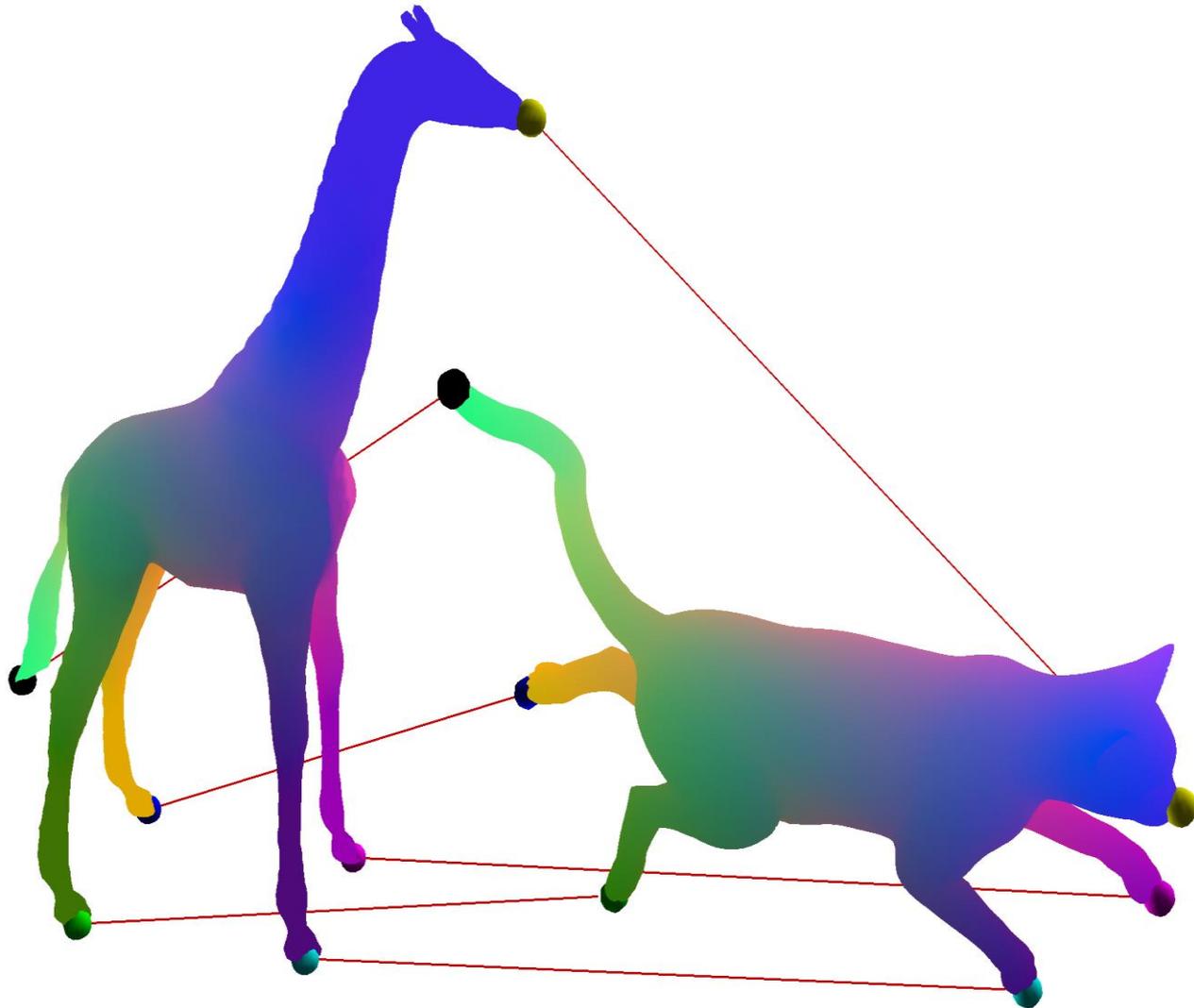


# Computing Blended Maps

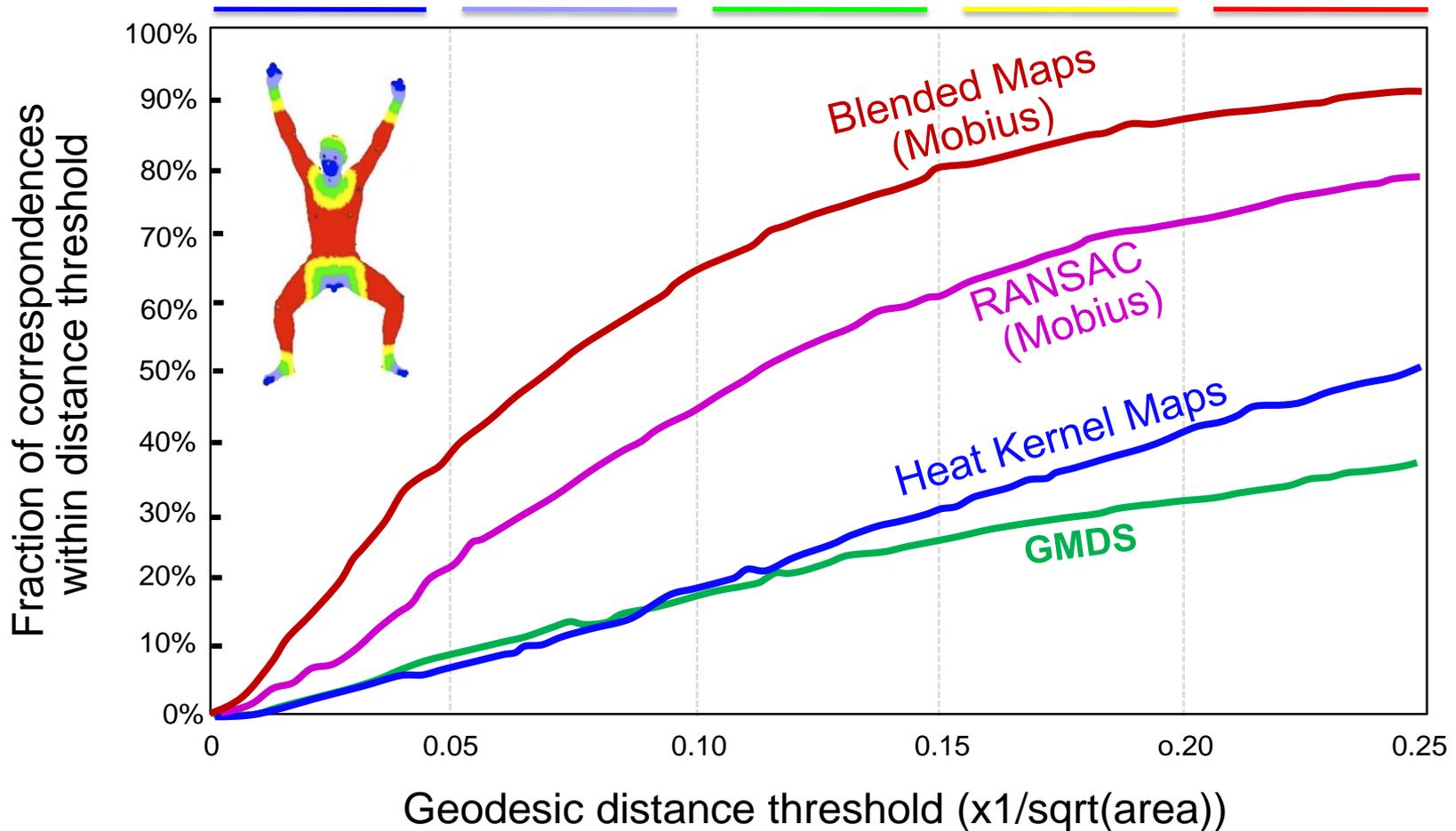
4. Define image  $p'$  of every point  $p$  as the weighted geodesic centroid of  $m_i(p)$



# Computing Blended Maps



# Experimental Results





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Example Application

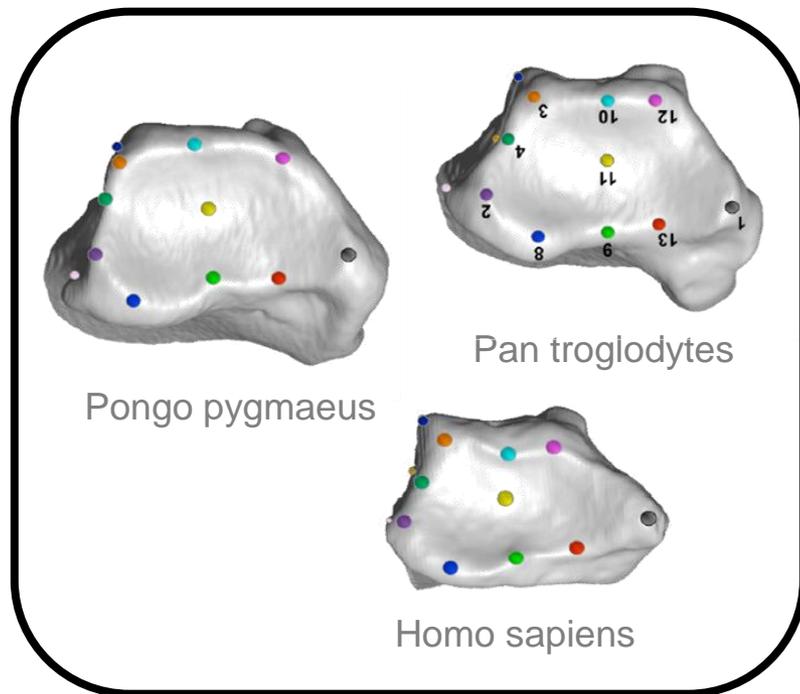
Conclusion

Future work

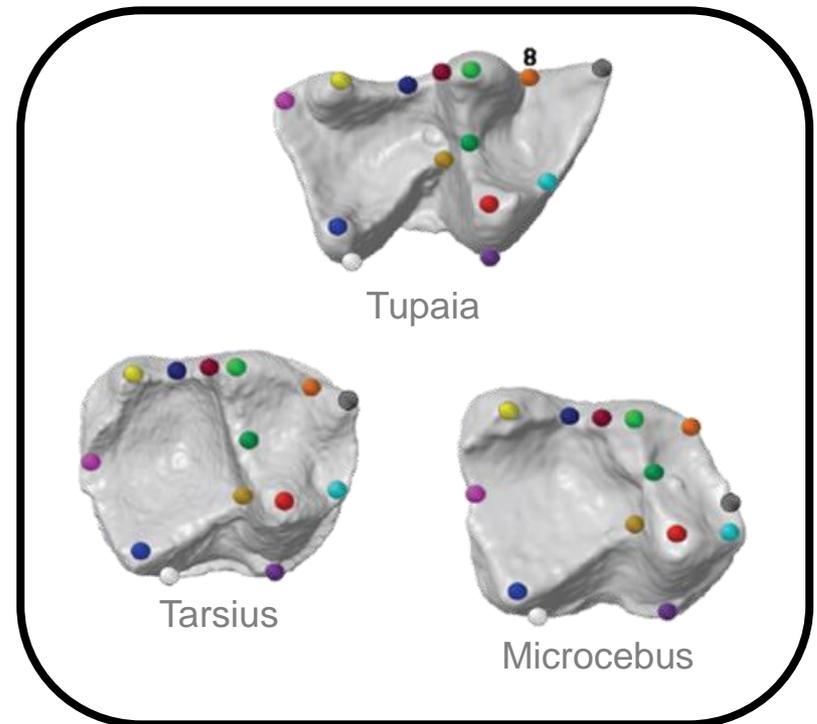


# Application

Automatically quantify the geometric similarity of anatomical surfaces



Distal Radius



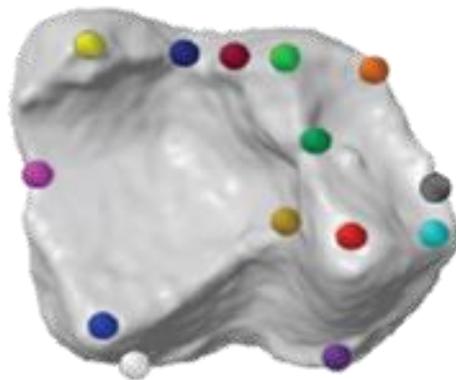
Mandibular Molar



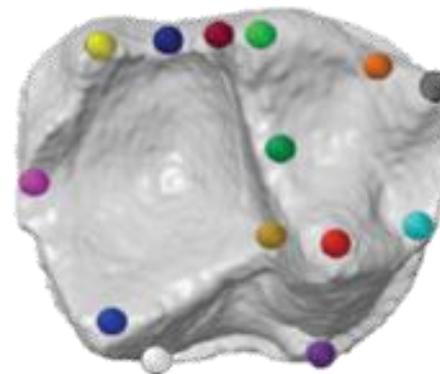
# Application

Traditional Procrustes distance:

$$d(X, Y) = \min_R \left[ \left( \sum_{i=1}^N \|R(X_i) - Y_i\|^2 \right)^{1/2} \right]$$



$X = \{ X_i \}$



$Y = \{ Y_i \}$

Human  
Specified  
Landmarks

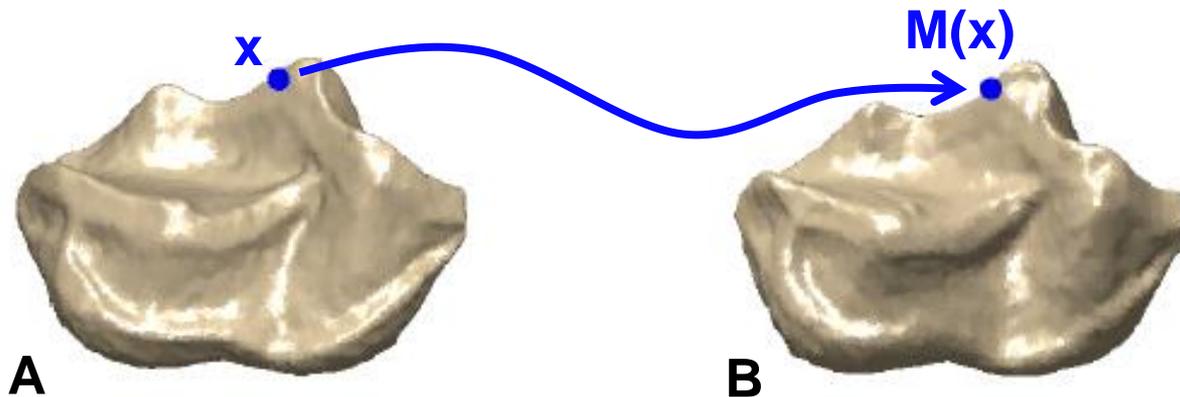




# Application

New continuous Procrustes distance:

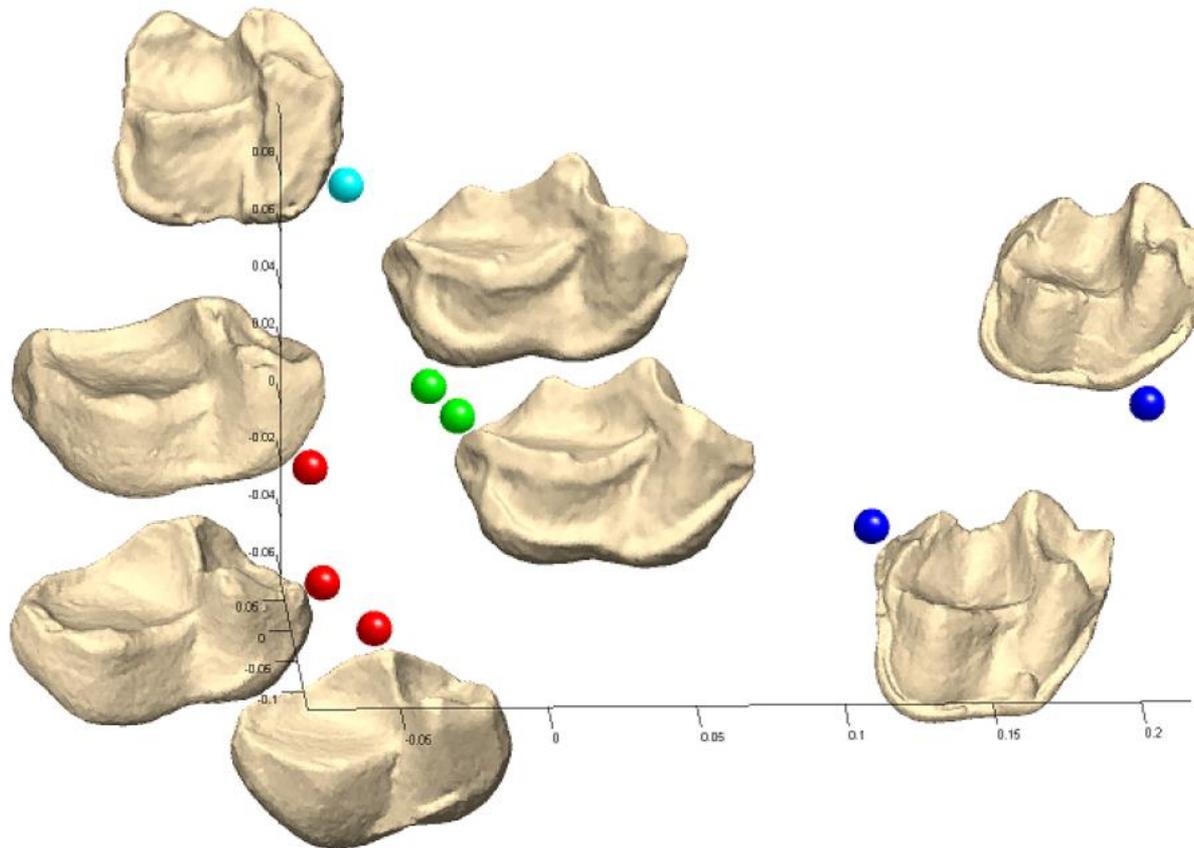
$$d(A, B) = \min_{R, M} \left[ \left( \int_A \|R(x) - M(x)\|^2 dx \right)^{1/2} \right]$$





# Application

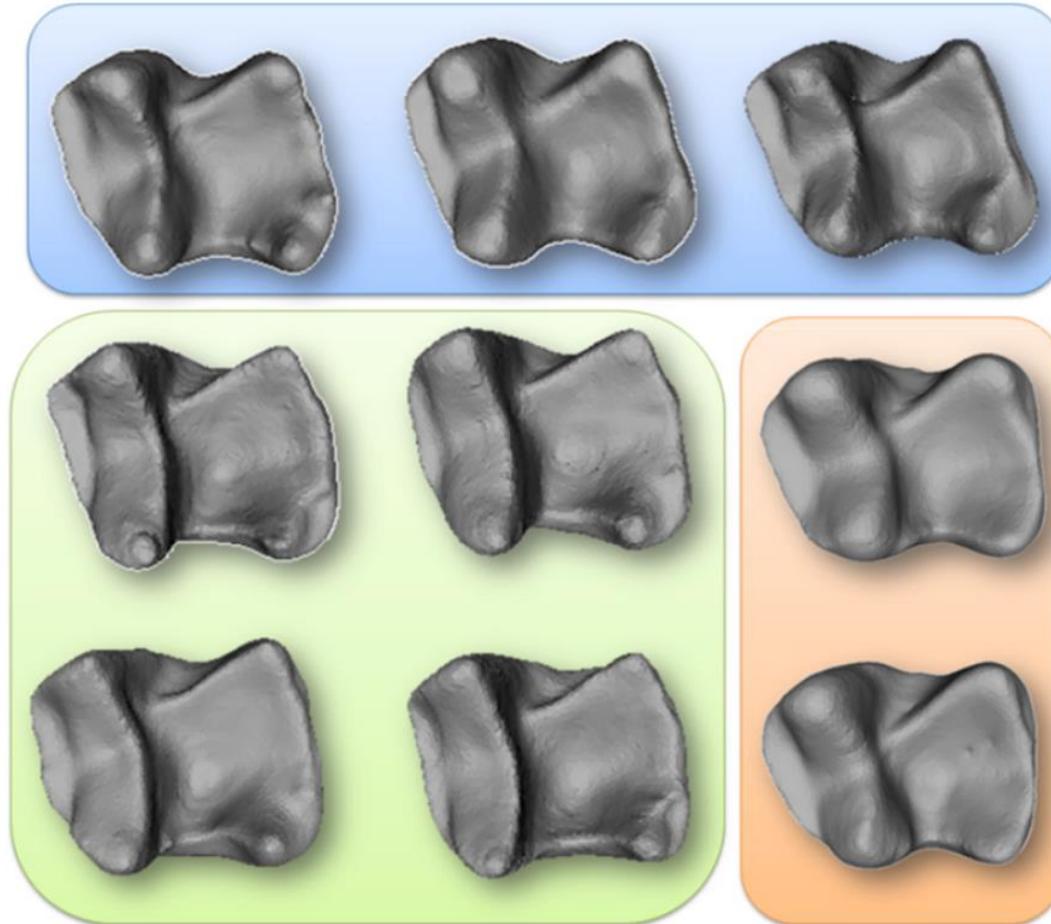
Embedding based on new distance





# Application

Clustering based on new distance



Species Groups of Galaga Genus

# Application



## Classification based on nearest-neighbors

<b>Mandibular Molar</b>	# Groups	# Objects	New Distance	Human Landmarks
Genus	24	99	90.9%	91.9%
Family	17	106	92.5%	94.3%
Order	5	116	94.8%	95.7%

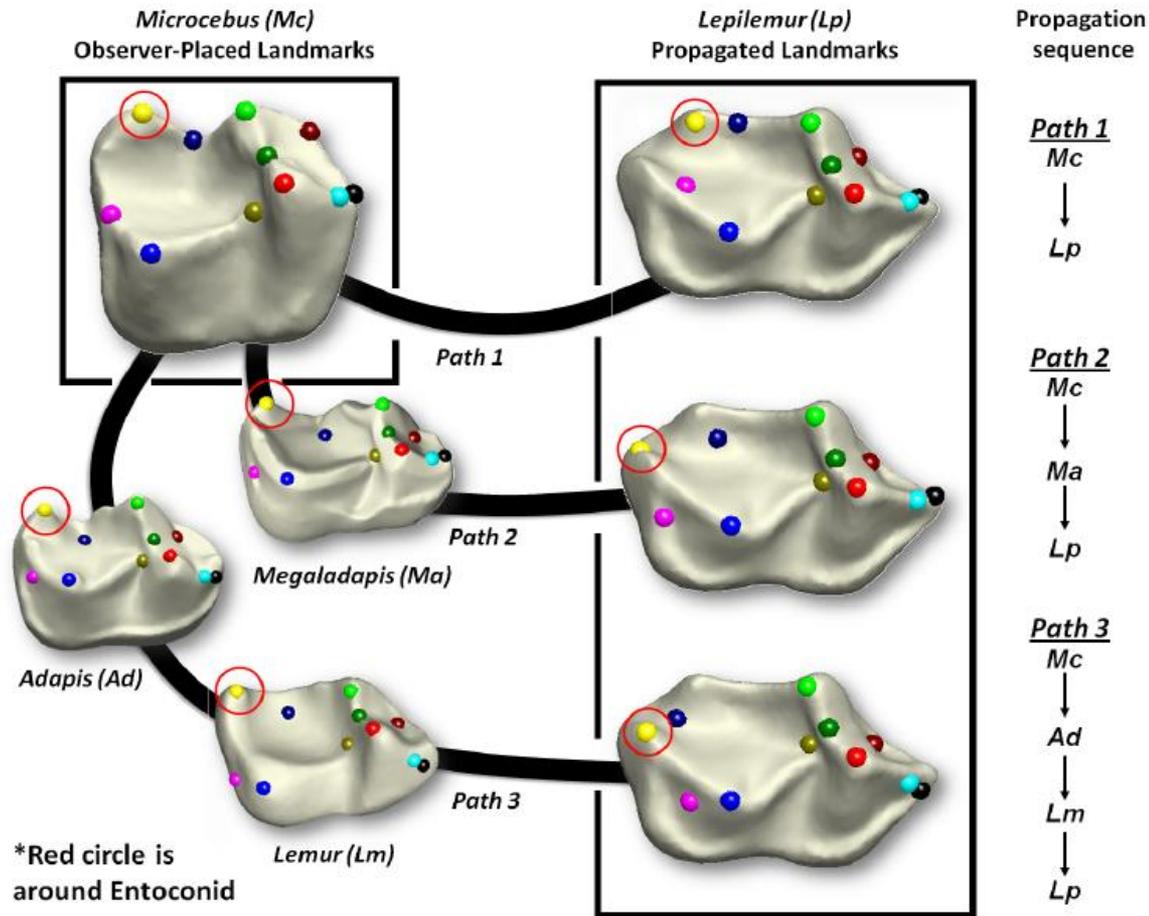
<b>First Metatarsal</b>	# Groups	# Objects	New Distance	Human1 Landmarks	Human2 Landmarks
Genus	13	59	79.9%	76.3%	88.1%
Family	9	61	91.8%	83.6%	93.4%
Superfamily	2	61	100%	100%	100%

<b>Distal Radius</b>	# Groups	# Objects	New Distance	Human Landmarks
Genus	4	45	84.4%	77.7%



# Application

## Propagating correspondences





# Acknowledgments

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## Test data

- Giorgi et al. (SHREC Watertight), Anguelov et al. (SCAPE), Bronstein et al. (TOSCA)

## Test code:

- Ovsjanikov et al. (HKM), Bronstein et al. (GMDS)

## Application

- Boyer, St. Clair, Patel, Jernvall, Puente, Daubechies

## Funding:

- NSERC, NSF, AFOSR, Intel, Adobe, Google

**Thank You!**