



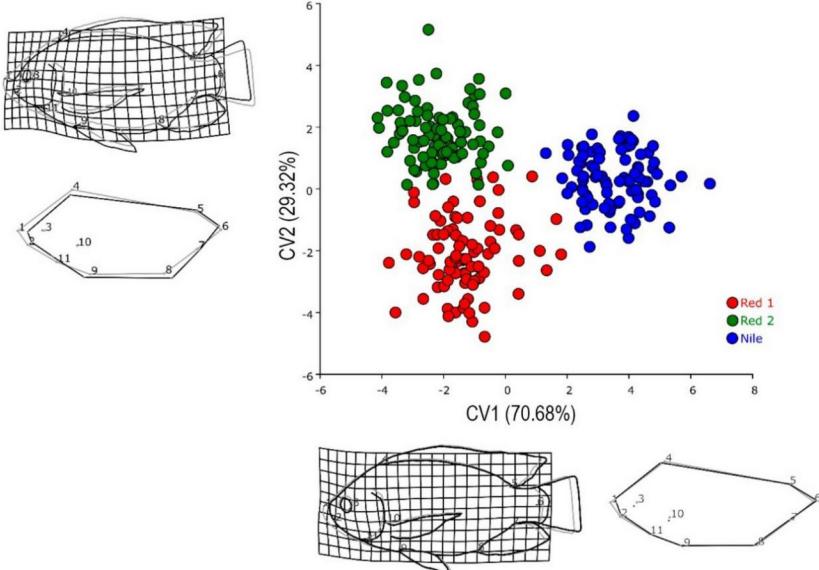
# ALPACA

*Automated Landmarking through Point-cloud Alignment and Correspondence Analysis*

# Morphometrics – EcoEvo

Morphometric analysis is widely used in ecology and evolutionary biology

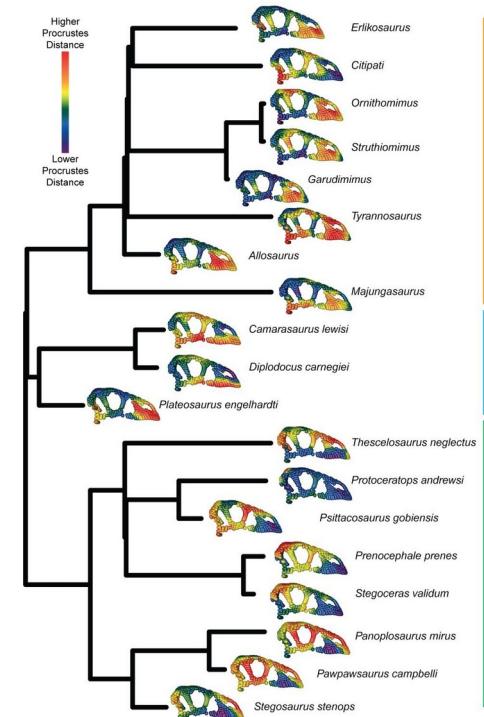
## Plasticity



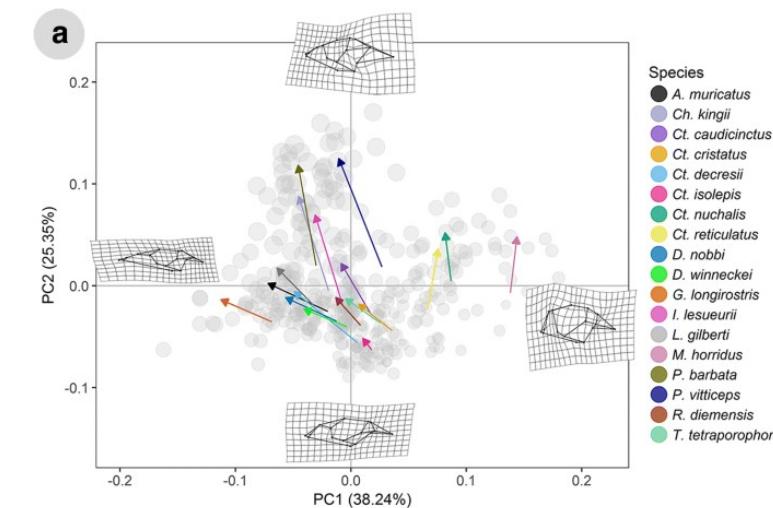
Montoya-Lopez et. al., 2019

## Evolutionary change

Felice et. al., 2020

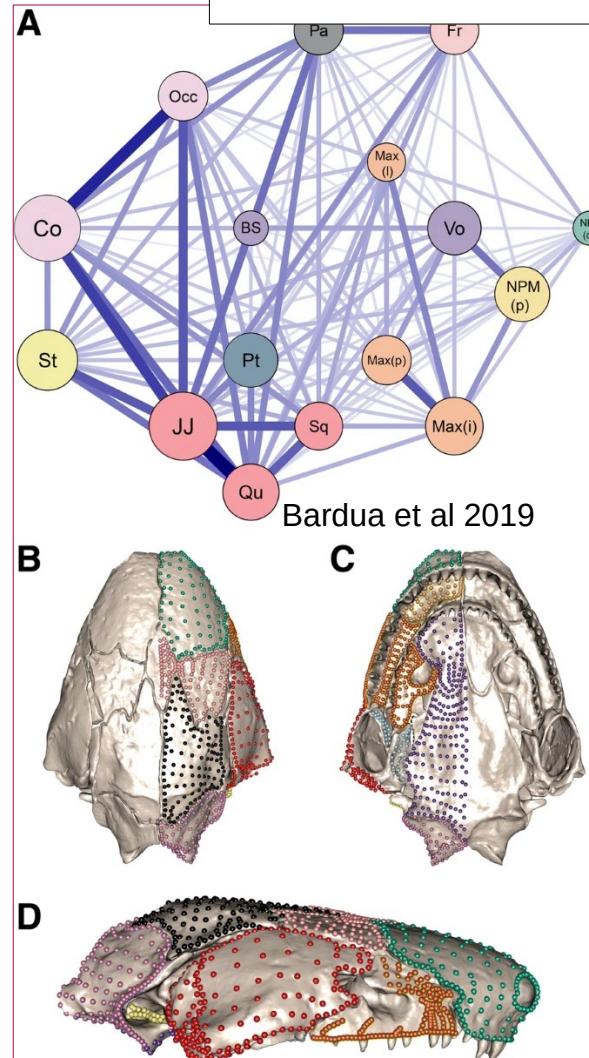


## Ontogeny

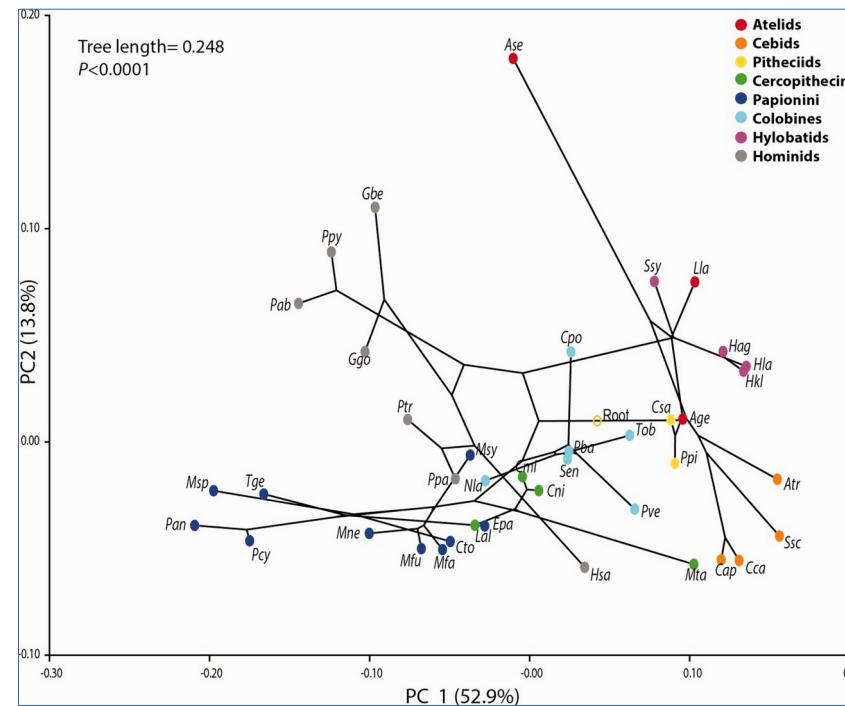


Gray et. al., 2019

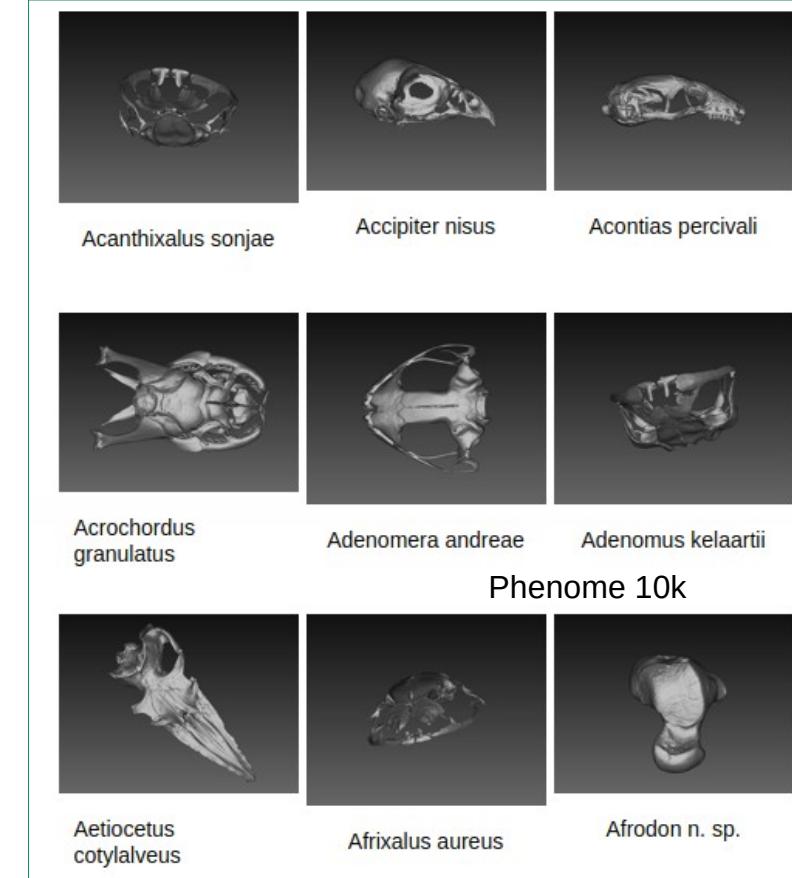
# Advances – Analysis, Data and Imaging



Modularity and  
Integration



Phylogenetics  
And  
Evolutionary  
Models

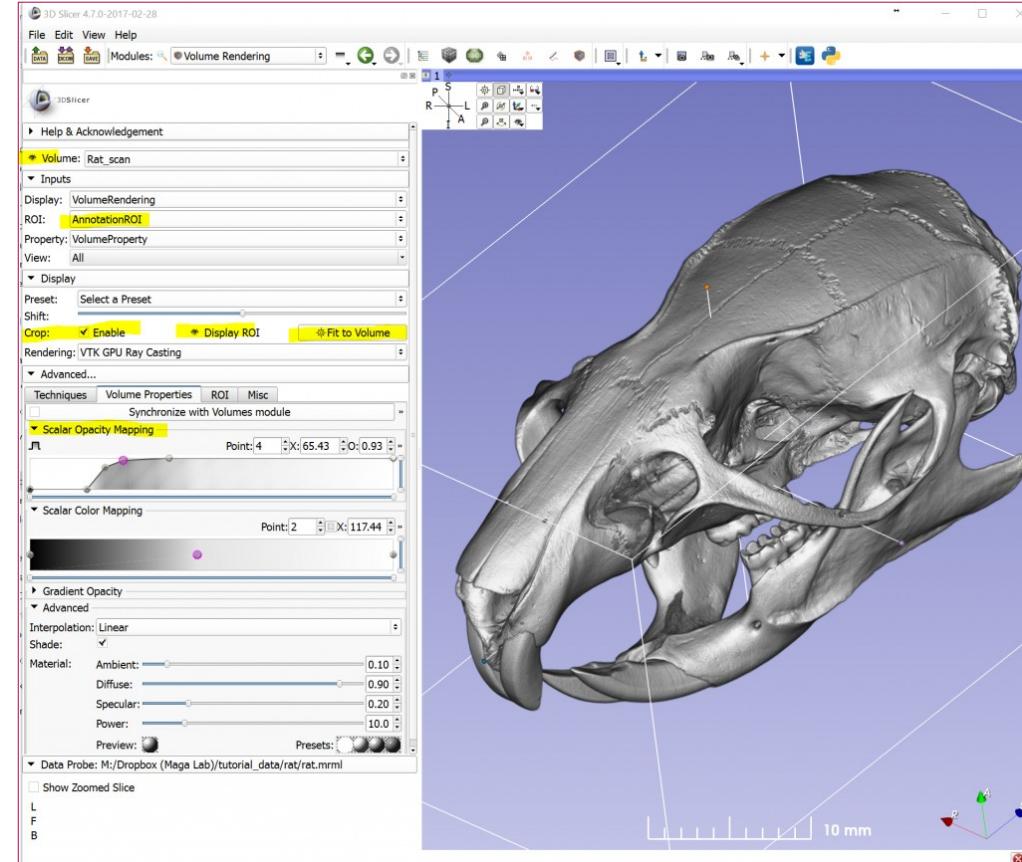


Public Databases  
and High res. images

# Data collection – Still largely manual

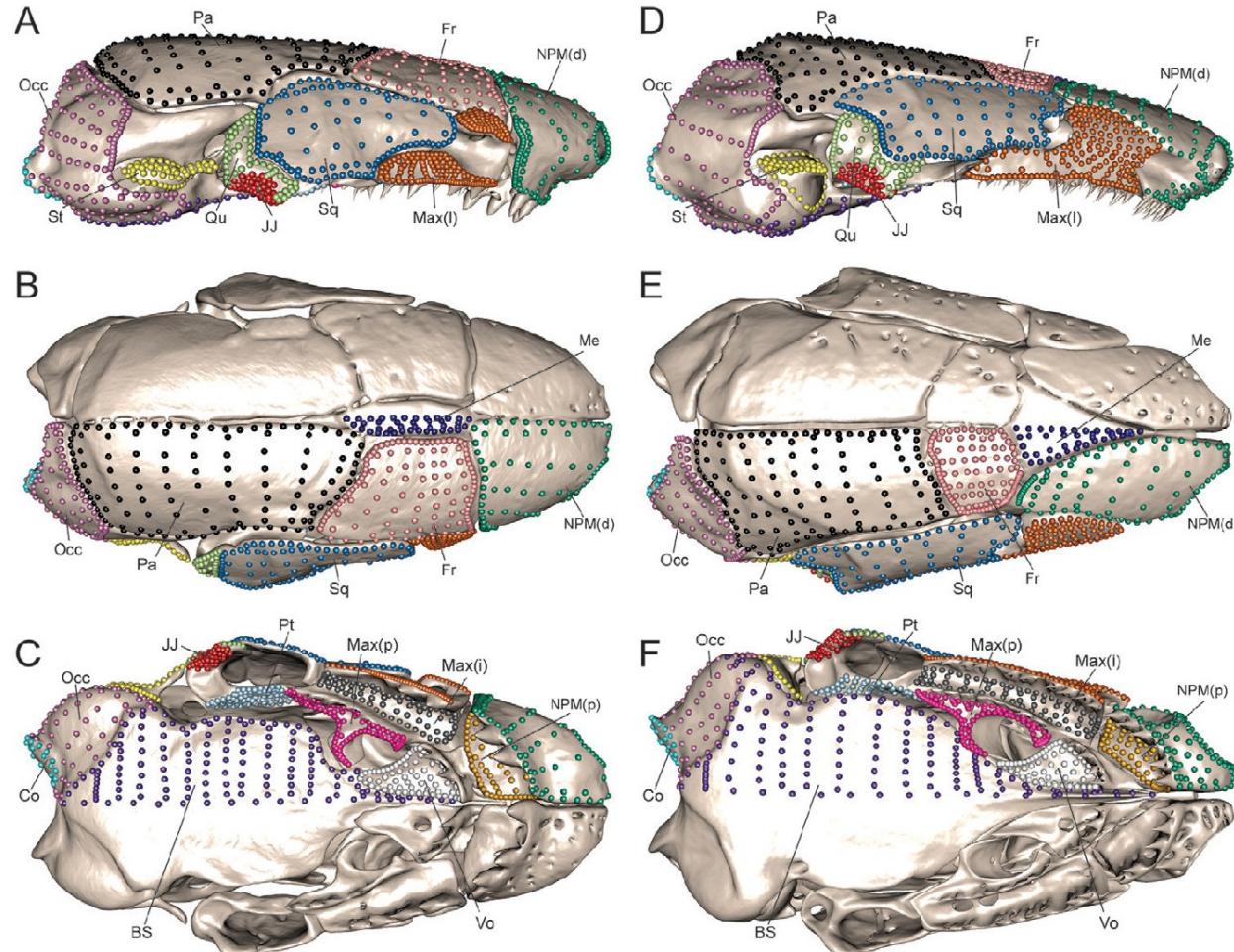


e.g., digitizers



e.g., software

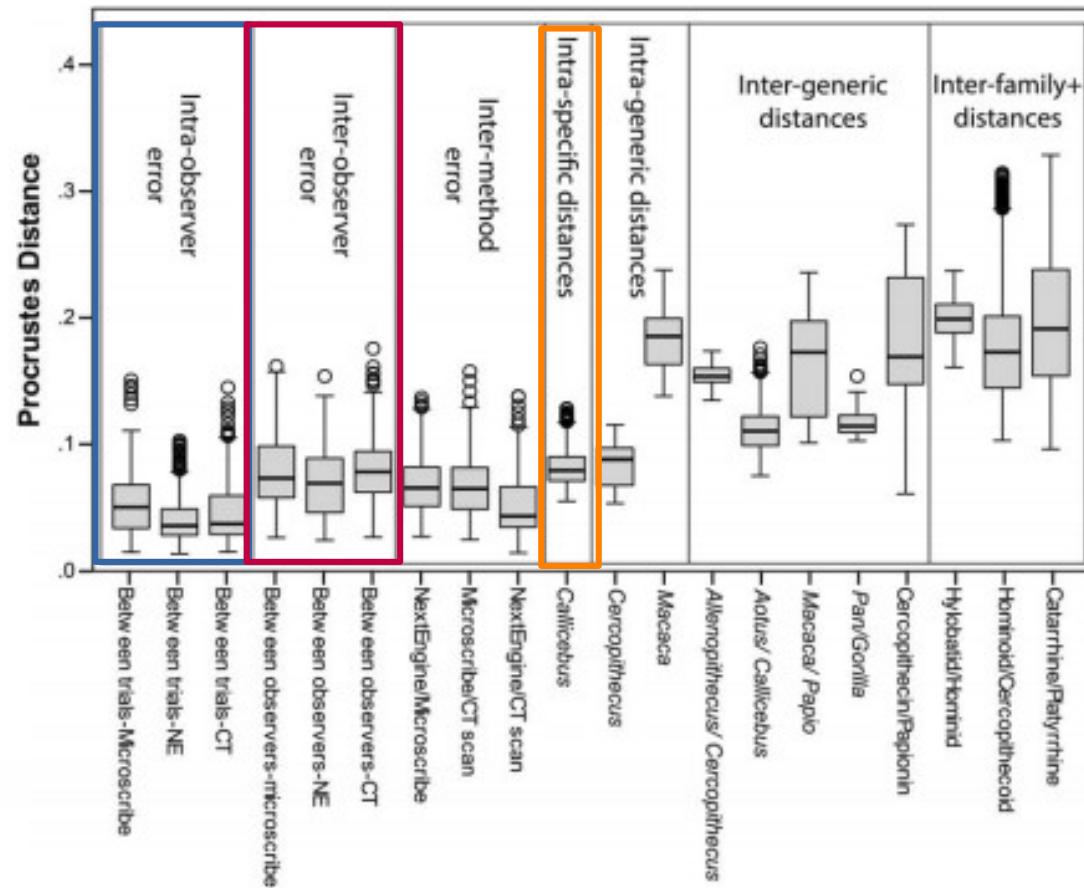
# Time consuming



66 (type 1)  
+  
336 (sliding)  
+  
736 (semi)

Marshall et. al., 2019

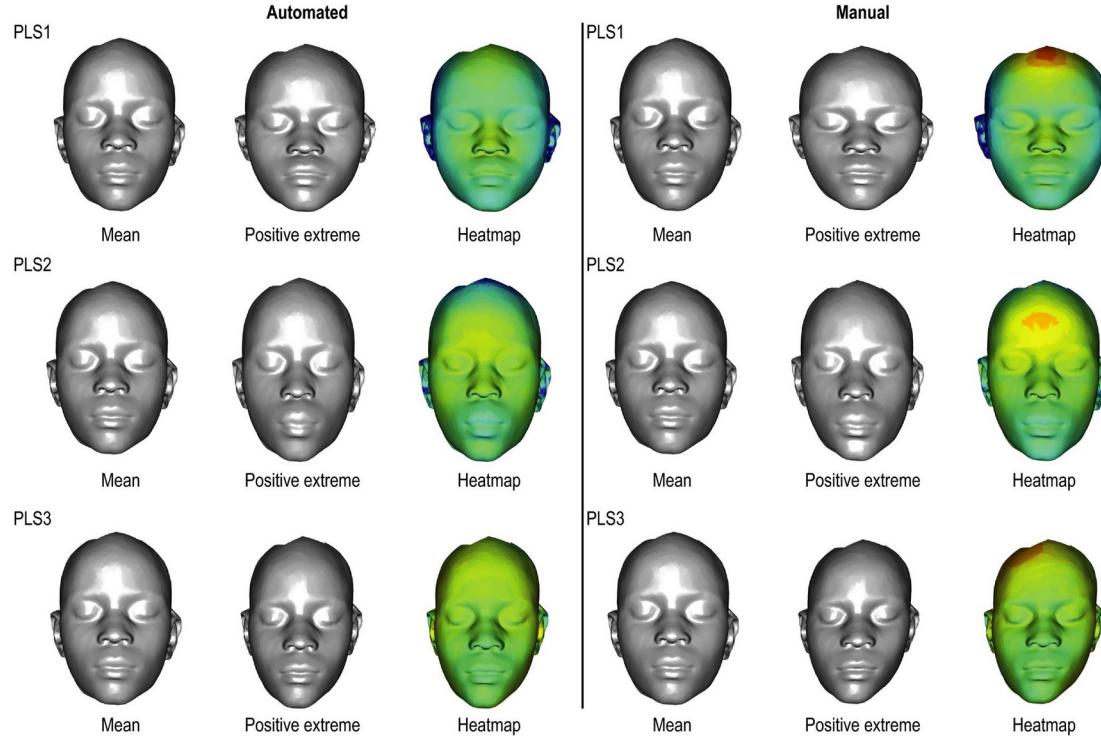
# Intra- and inter-observer error



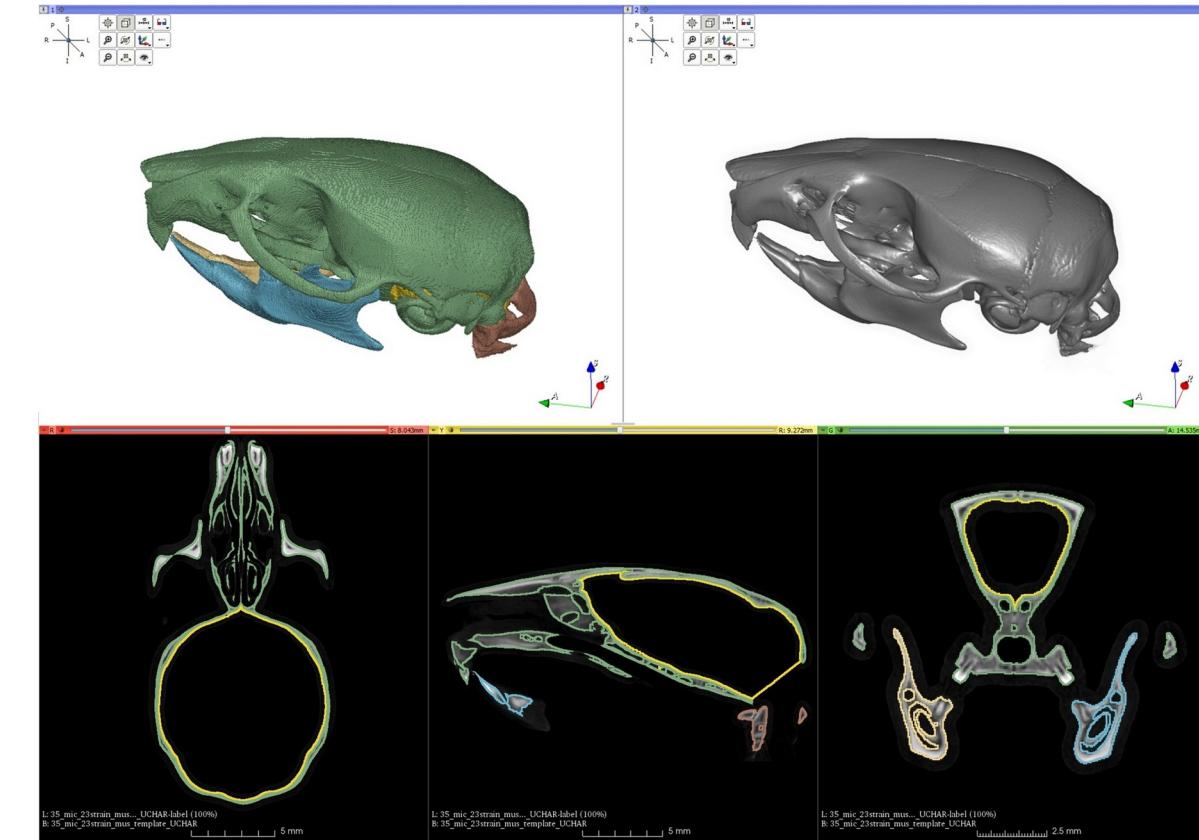
**FIGURE 4** Box plot of Procrustes distances within and among observers, methods, and taxa. Darkened bars represent the median value for each group, boxes show the interquartile range (25<sup>th</sup> to 75<sup>th</sup> percentile), and the whiskers extend to 1.5 times the interquartile range. Outliers are designated by circles

# In contrast with biomedical research

c 3D Morphs and heatmaps corresponding to PLS Scores 1-3



Li et al. "Rapid automated landmarking for morphometric analysis of three-dimensional facial scans." Journal of Anatomy (2017).



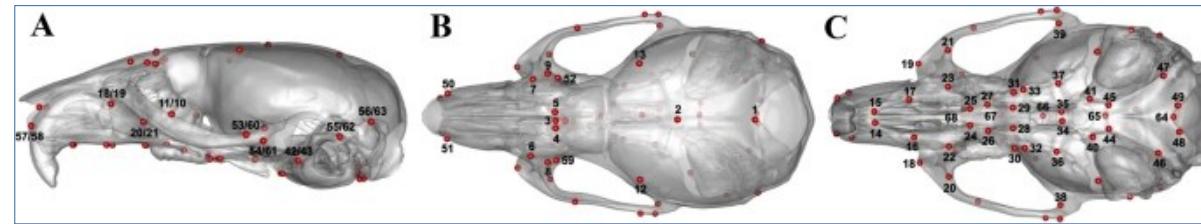
Maga et al. A population level atlas of *Mus musculus* craniofacial skeleton and automated image-based shape analysis. Journal of Anatomy (2017)

# Ecology and Evolution - Challenges

\* Lack of comparable resources

- Personnel: imaging scientist

- Hardware: automation = as much as 10 CPU/hours per specimen



e.g. Devine et al 2020

\* System-specificity

- Algorithms: species-specific and hard to generalize

# Our attempt



ALPACA

Intuitive

Users with no prior experience in image analysis could use it

Lightweight

Run in minutes  
(much less time than manual digitization)

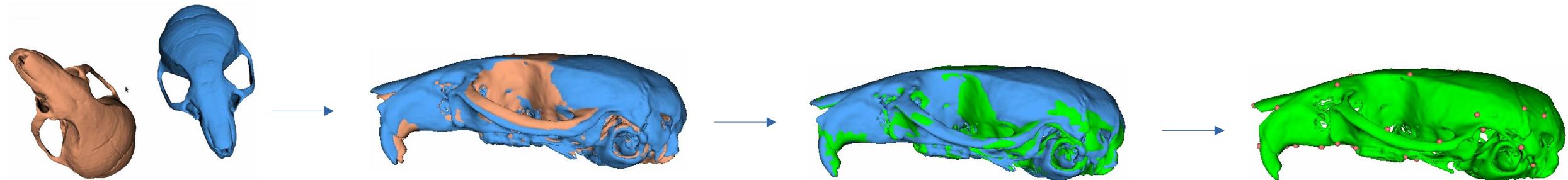
General

Can be applied to any structure of interest, given some limitations in the degree of shape variation

In many ways, it is similar to traditional methods

## Registration

is the process of finding the optimal transformation to maximize correspondence across specimens



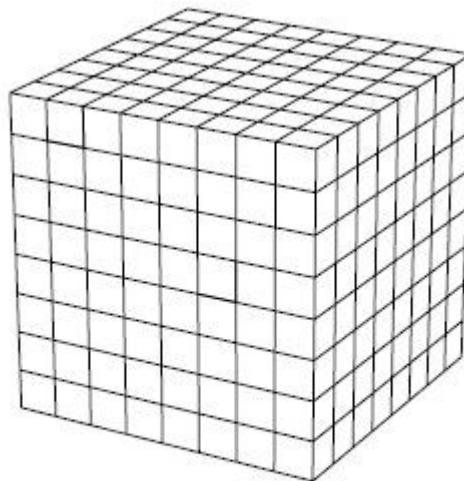
# But it is also different

Key aspect

Sparse representation can be used to find for alignment

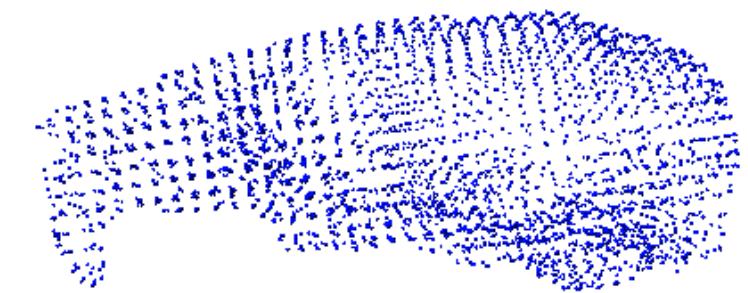


Mesh



Voxel Grid

User-specified

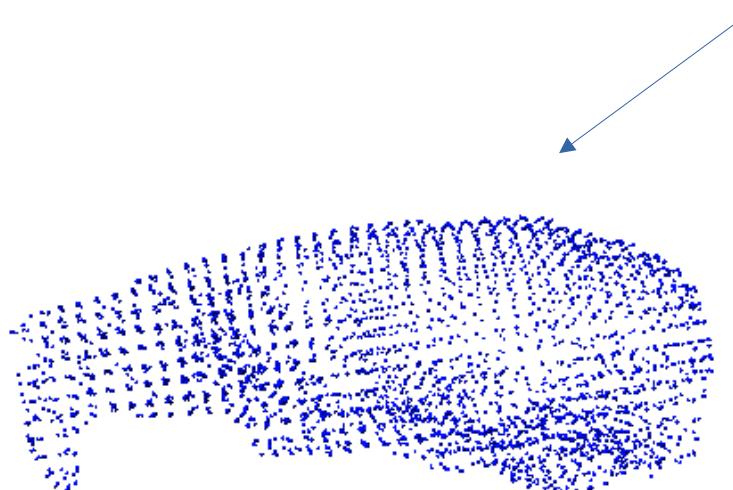


Voxel size = 0.8 mm

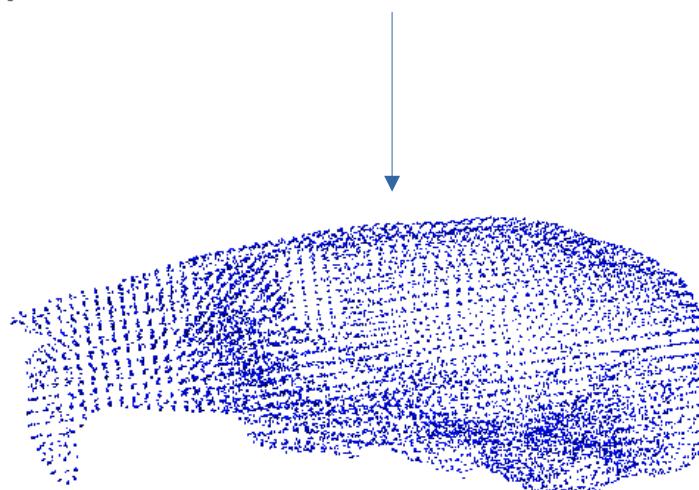
Point-cloud

# But it is also different

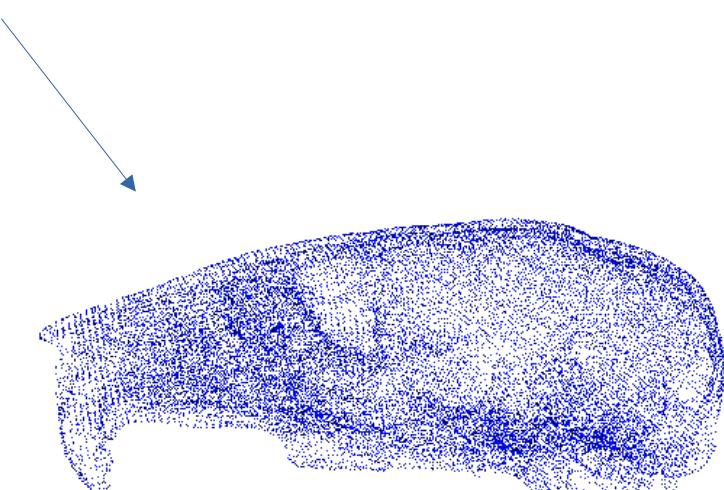
How sparse? User has control over it



Voxel size = 0.8 mm

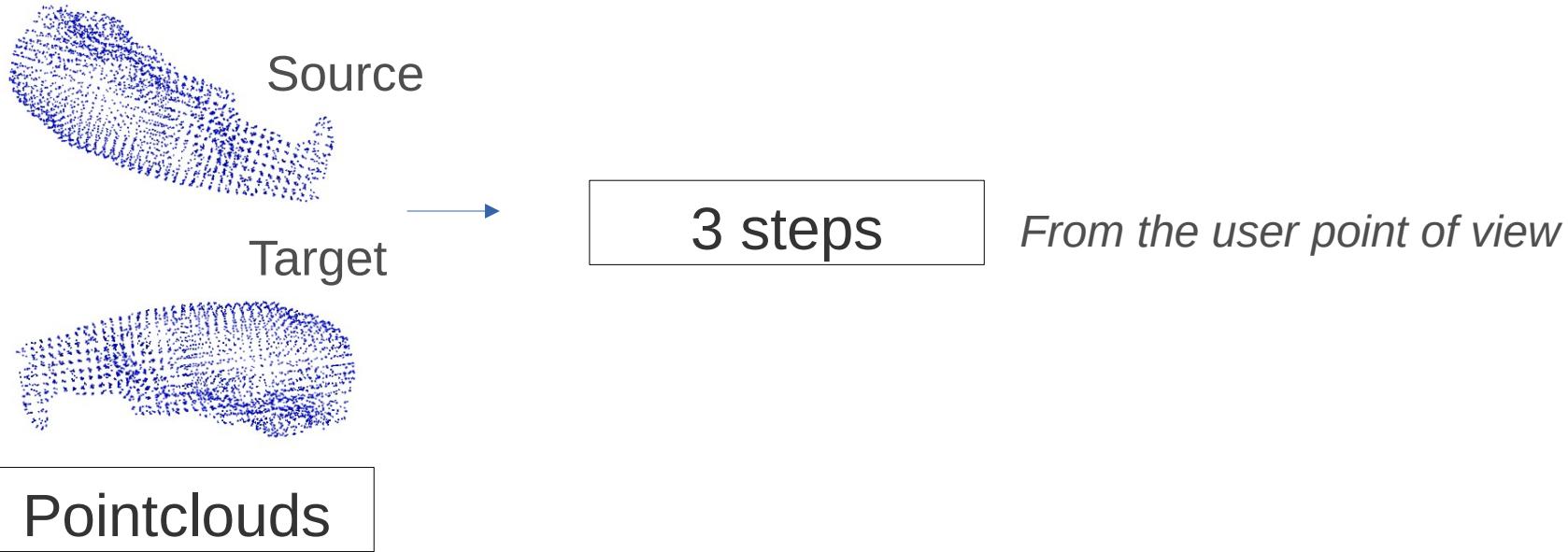


Voxel size = 0.6 mm

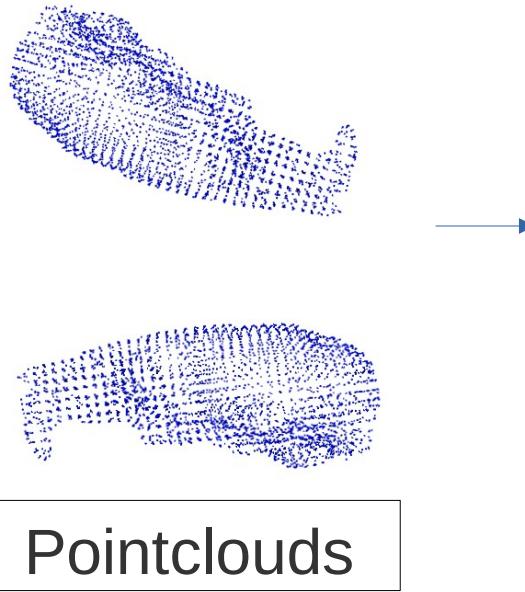


Voxel size = 0.2 mm

# Once pointclouds are obtained



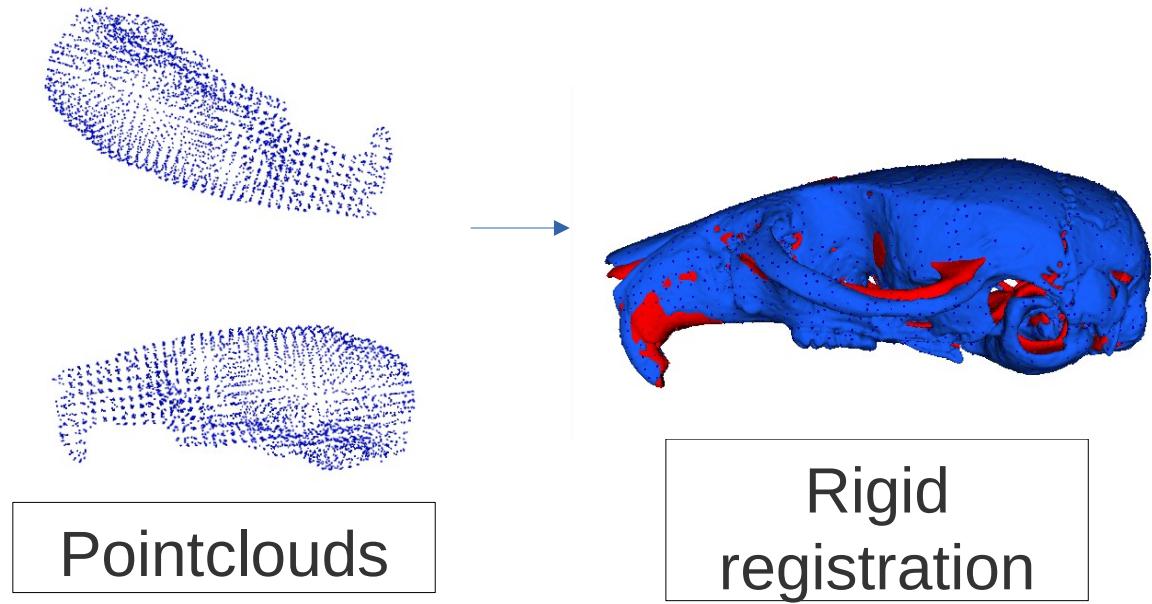
# Rigid



Pointclouds

Rigid  
registration

# Rigid



Pointclouds

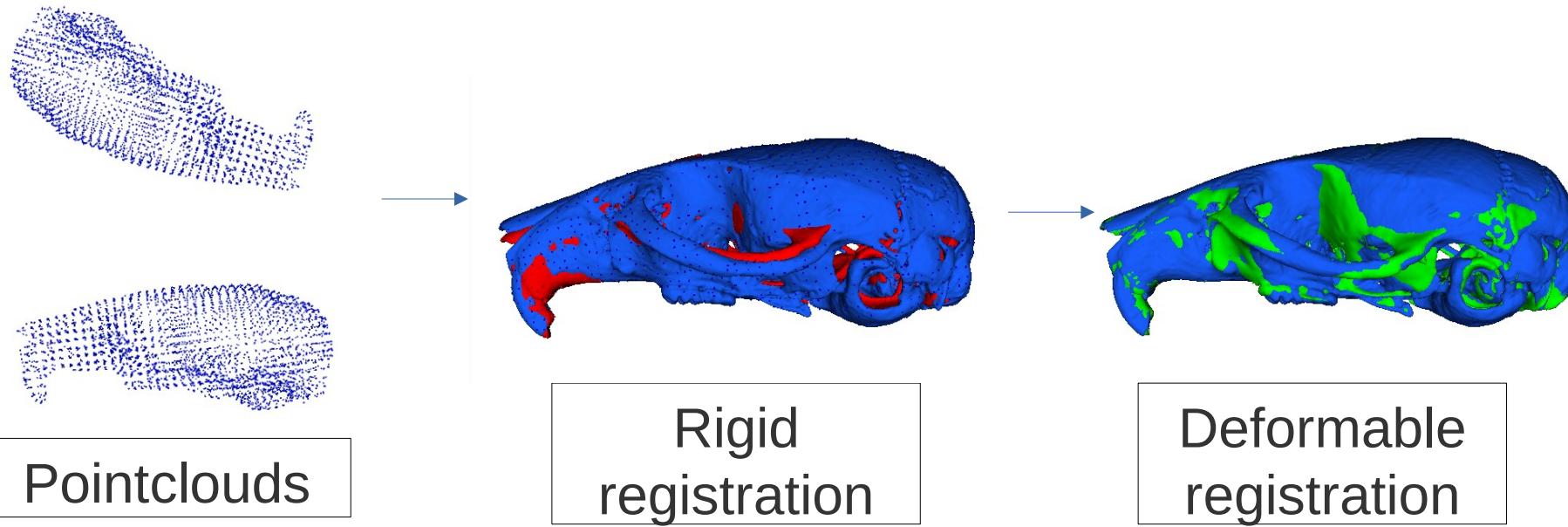
Rigid  
registration

Global registration = Finds the initial (rough) alignment (FPFH-based RANSAC)

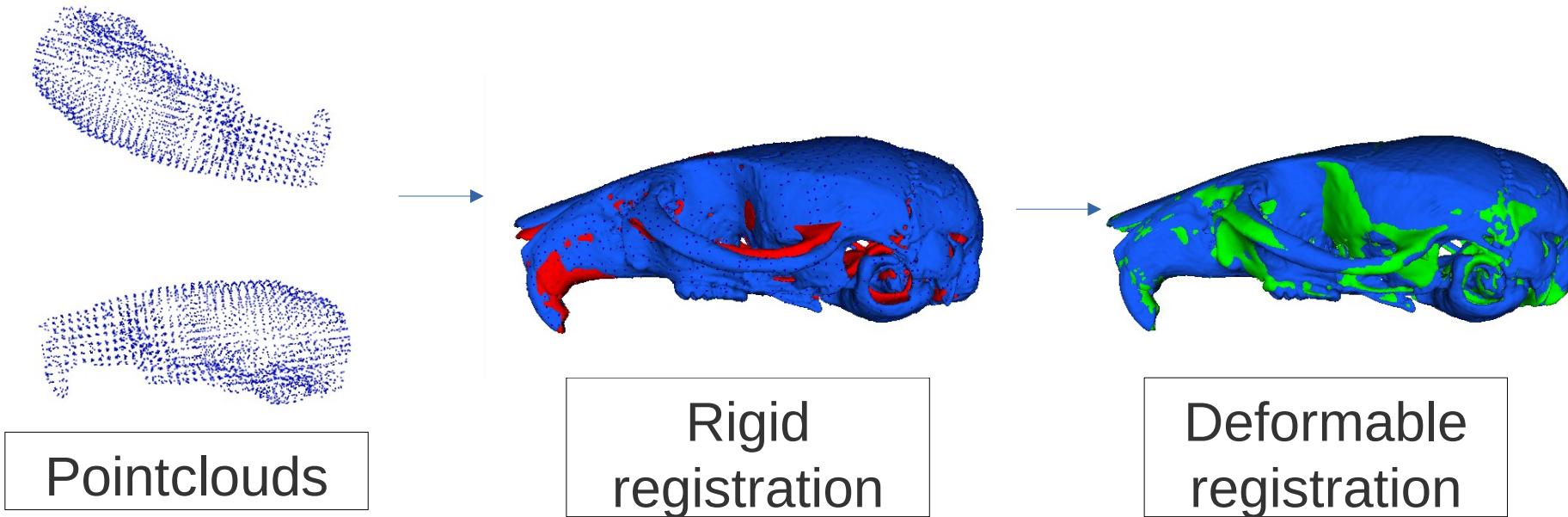
+

Local registration = Refines the rough alignment (point-to-plane ICP)

# Deformable

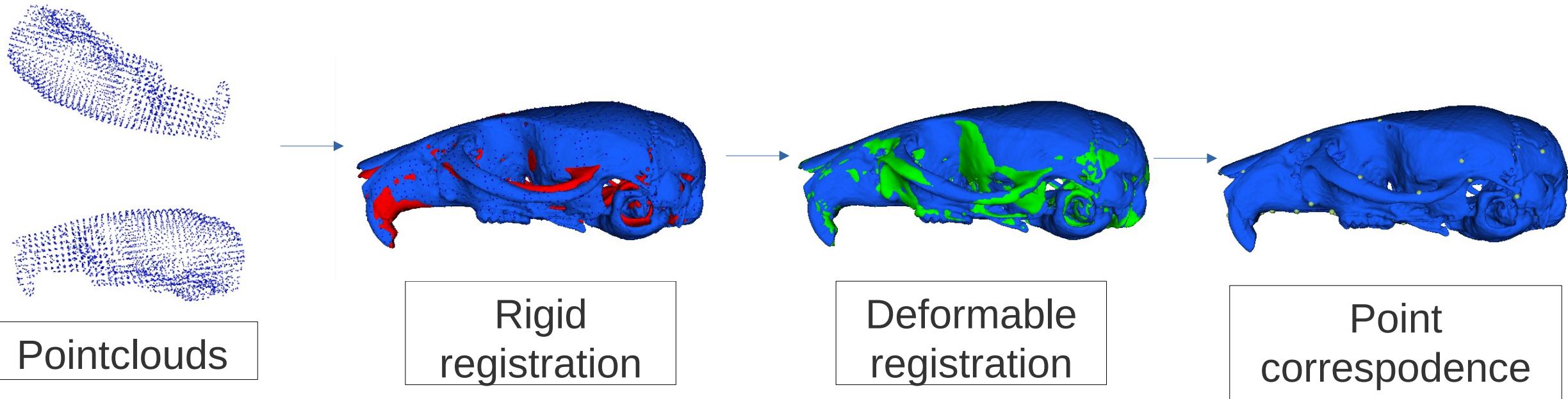


# Deformable

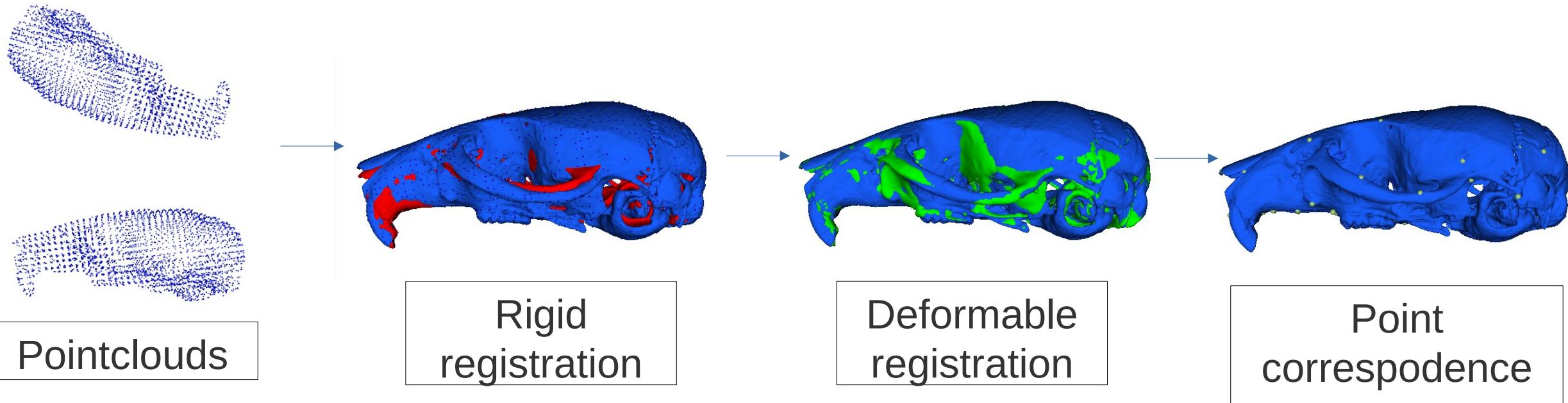


Finds the deformation necessary to closely align one pointcloud to another, given some constraint (Coherent Point Drift)

# Deformable



# Deformable



Transfer the landmark positions from the source to the target mesh

# So far - Four datasets

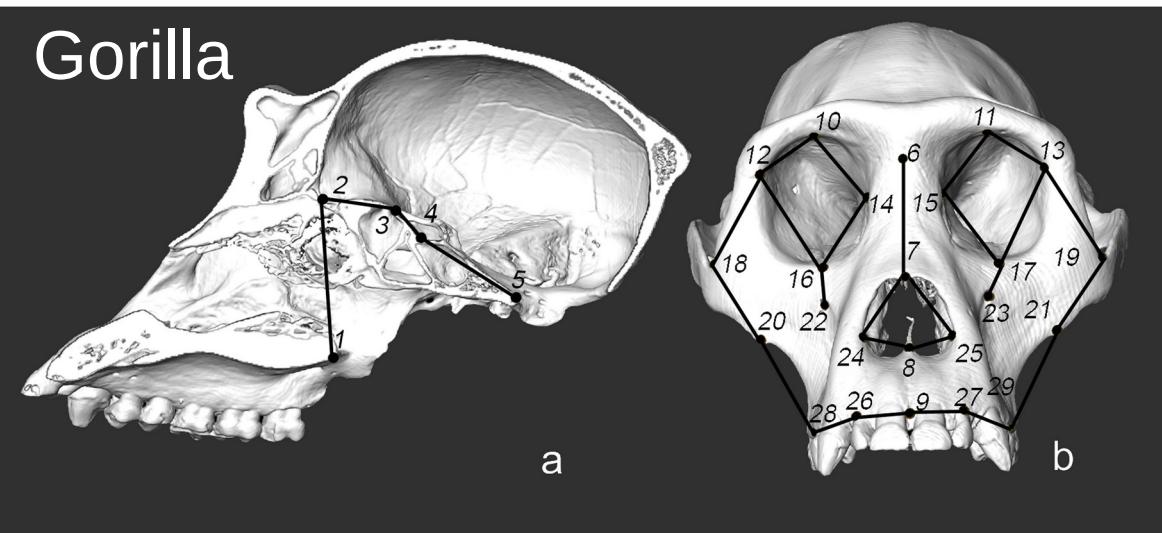
Mouse



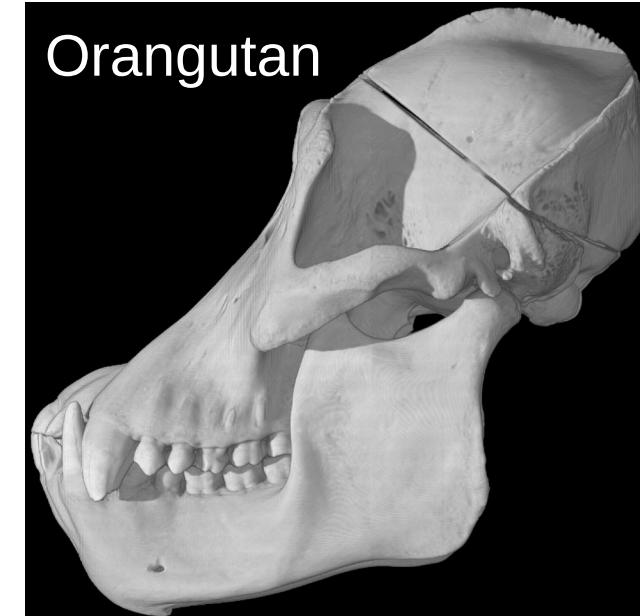
Chimps



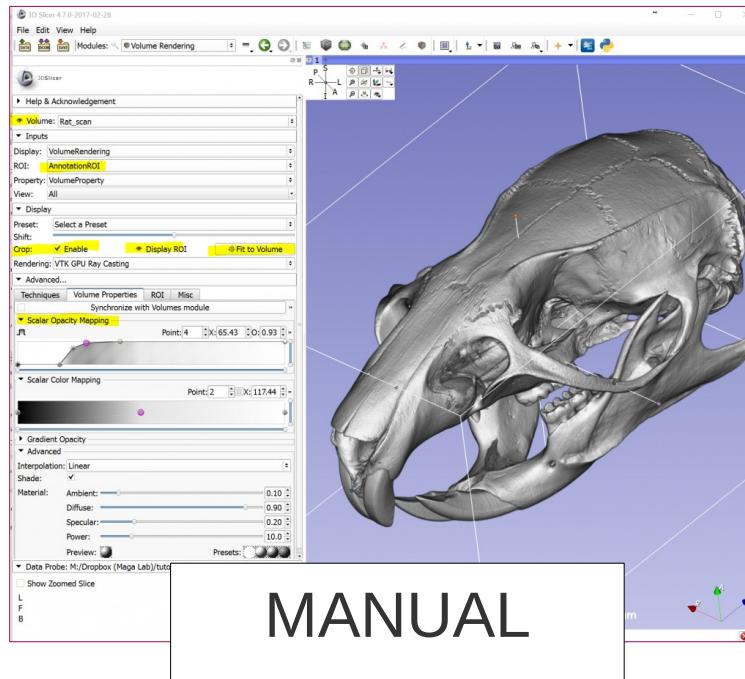
Gorilla



Orangutan



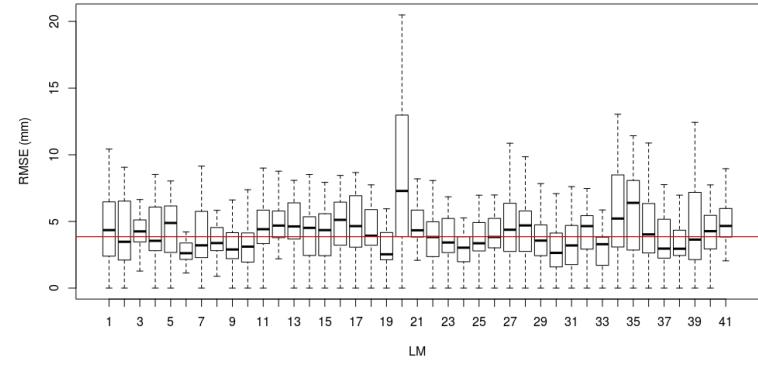
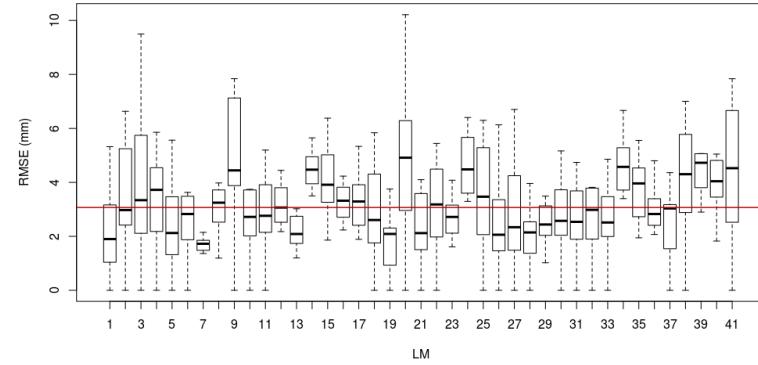
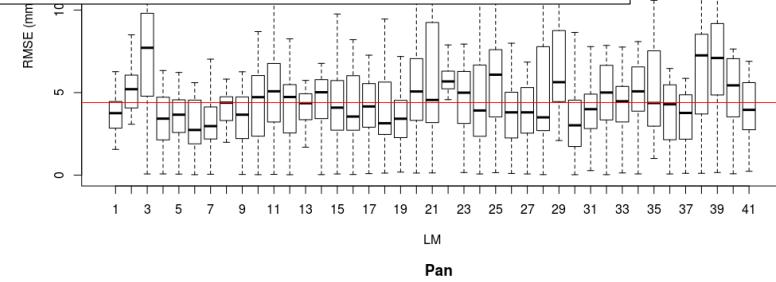
# Method comparison



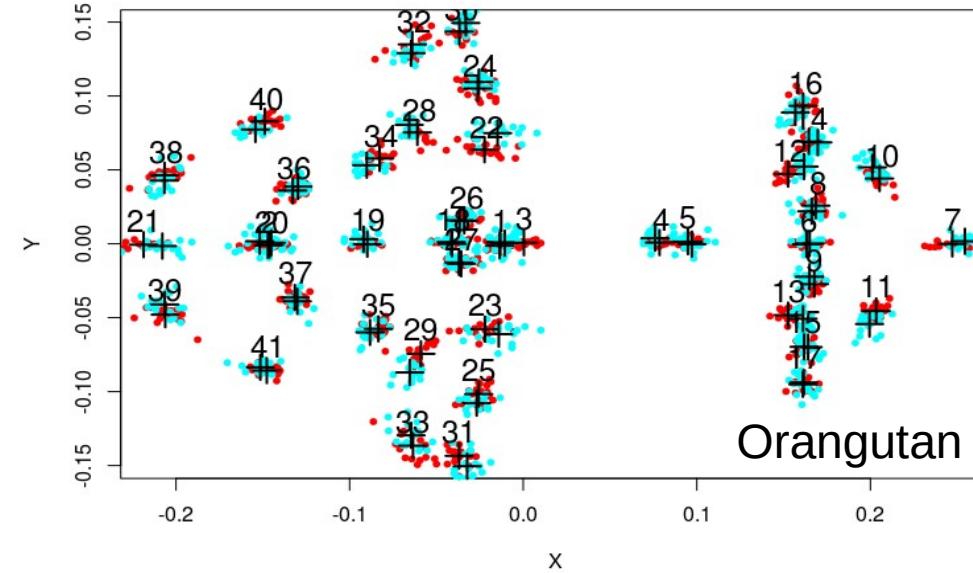
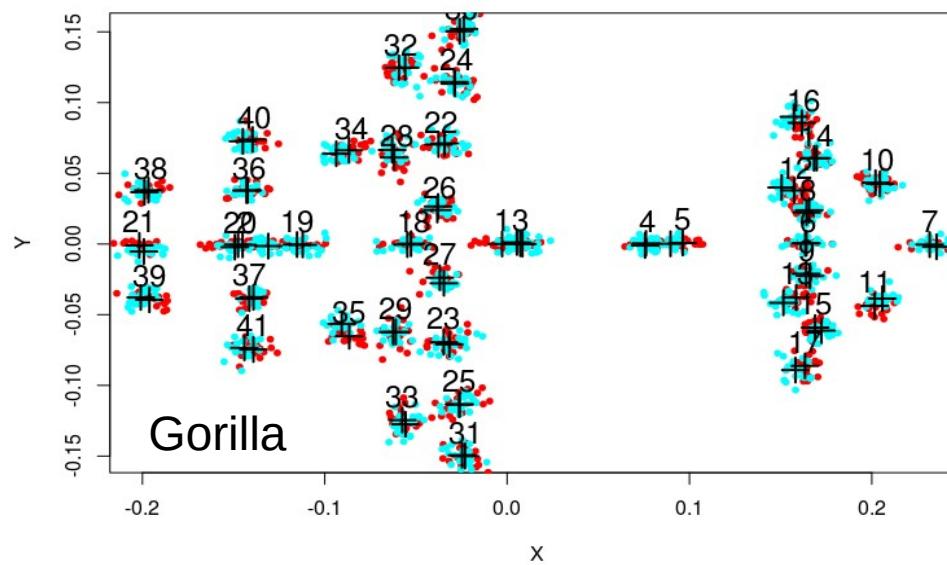
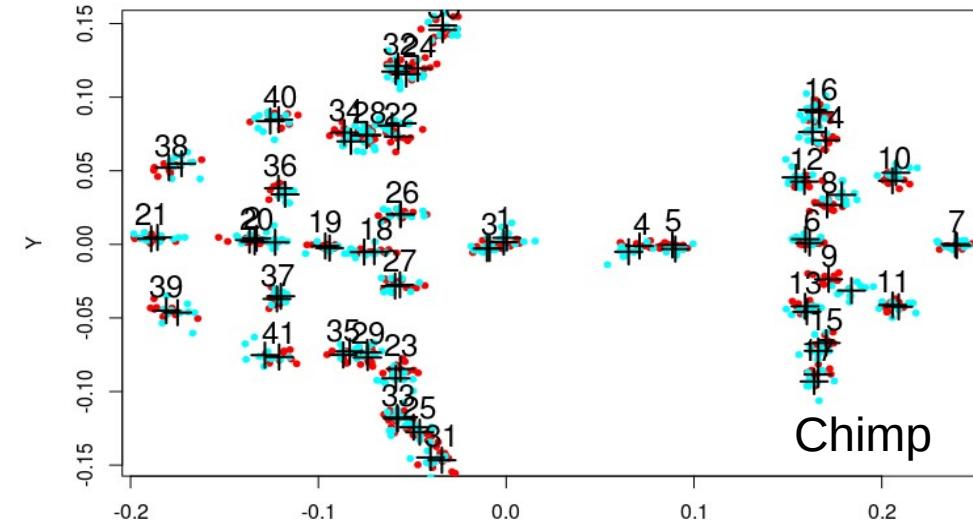
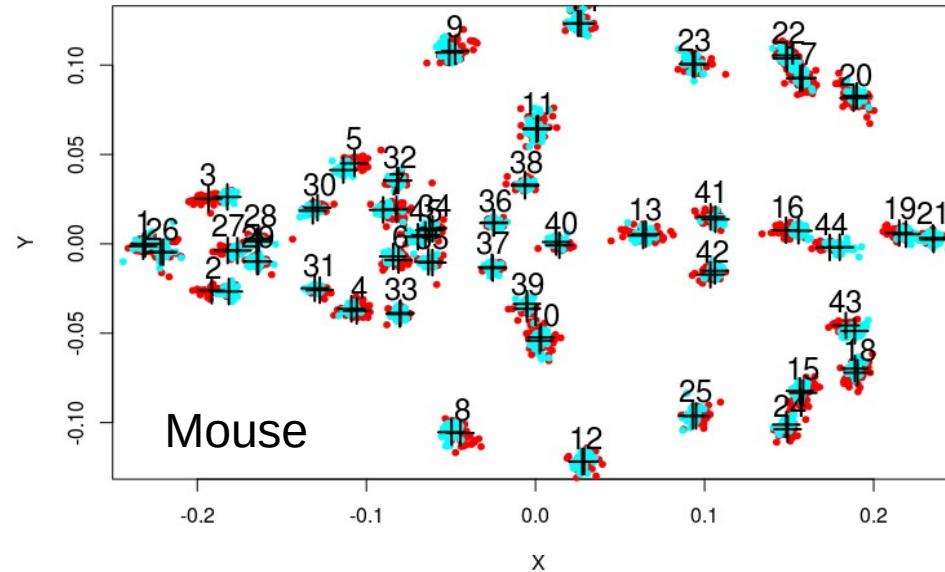
VS



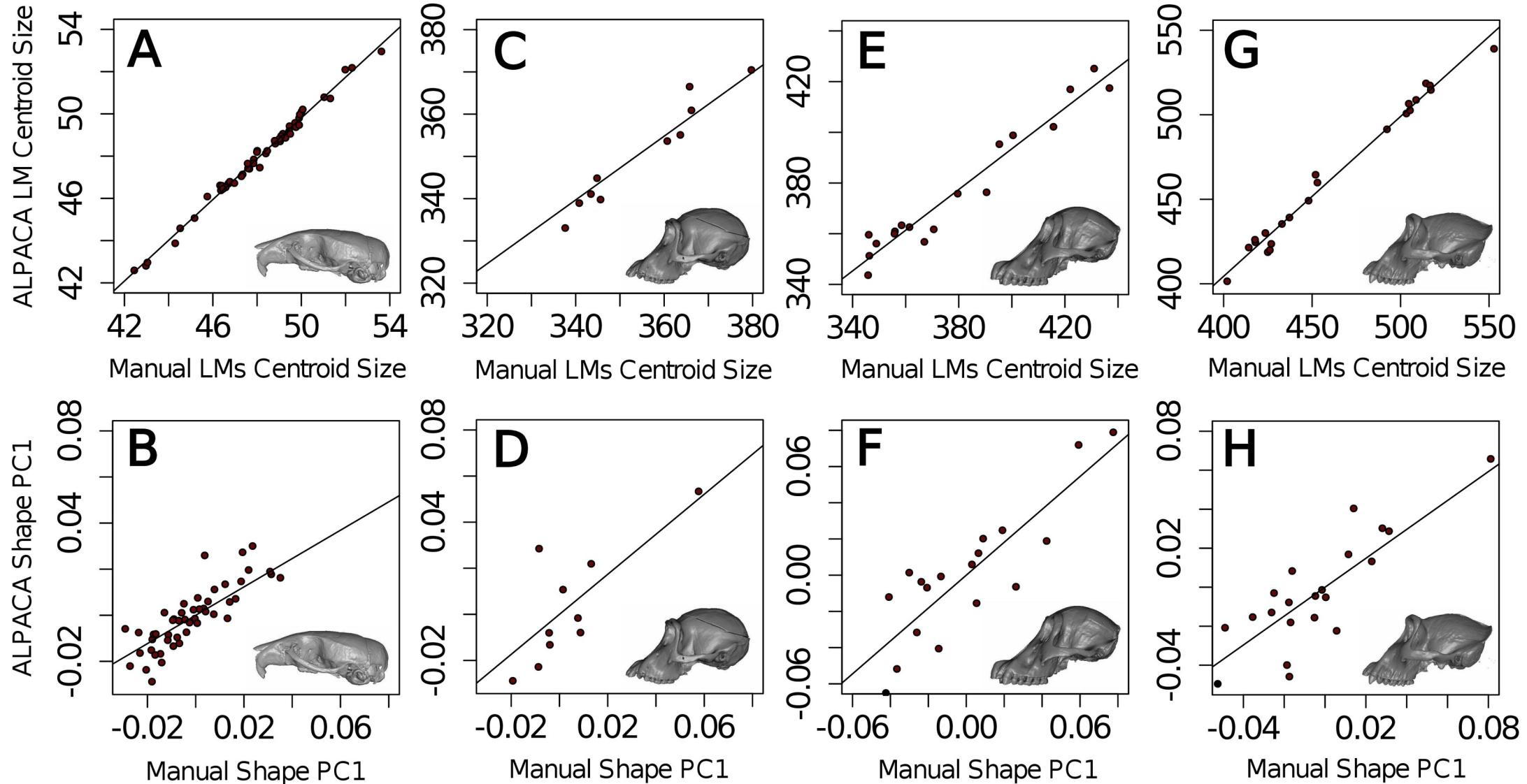
# Method comparison – Euclidean space



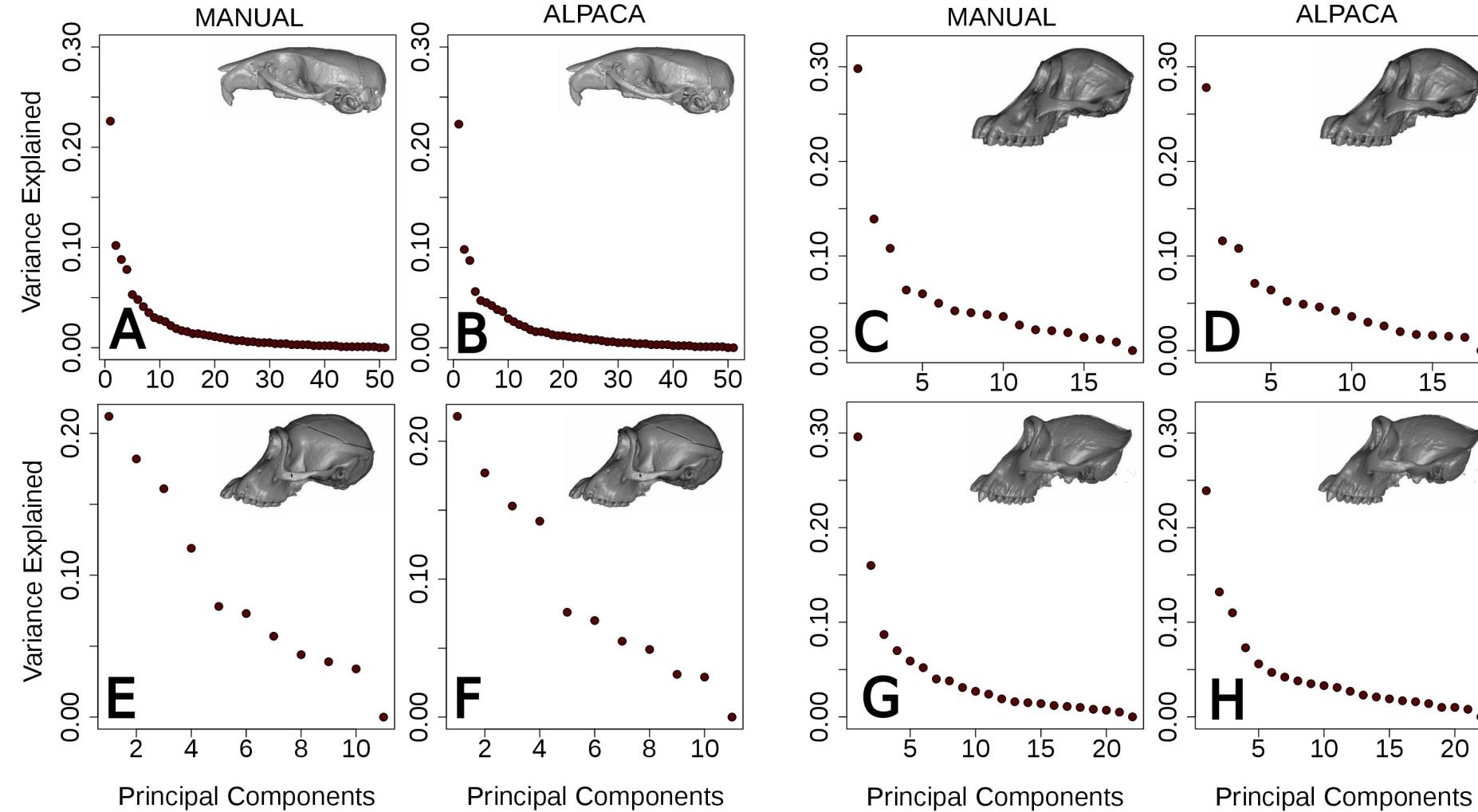
# Method comparison – LM configurations



# Method comparison - Centroid Size and PC1



# Method comparison – Scree plot



# But we are counting on you to really test it ..

**\*\* Please try with your own datasets and let us know how it goes \*\***

If you have trouble, we are here to help.

The greater the use, the better the tool will become

# But we are counting on you to really test it ..

**\*\* Please try with your own datasets and let us know how it goes \*\***

If you have trouble, we are here to help.

The greater the use, the better the tool will become

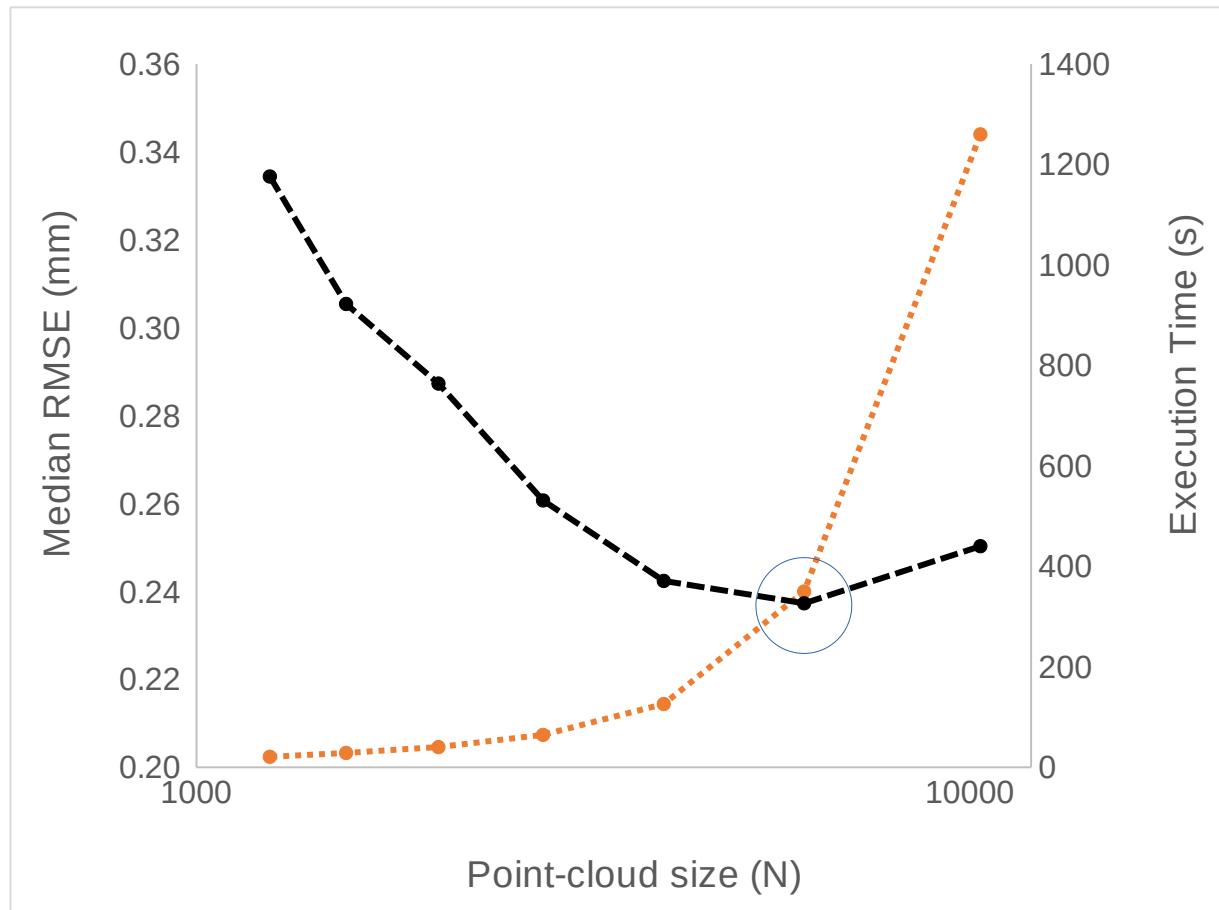
## Short term improvements – Next 4 weeks

- 1) Add scaling option
  - \* User will have the option to scale the source mesh to match the target one
- 2) Voxel size will be inferred directly from the mesh
  - \* No need to choose subsampling voxel size

# A few practical considerations

# Accuracy x Resources

\* First tip : Aim for 5,000 to 6,000 points

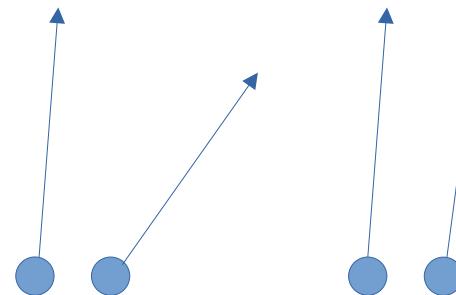


$N=62$  specimens

# Deformable – Structure specific

The deformable step has two regularization parameters ( $\alpha$  and  $\beta$ ) . These parameters will typically have to be modified to better serve your own needs.

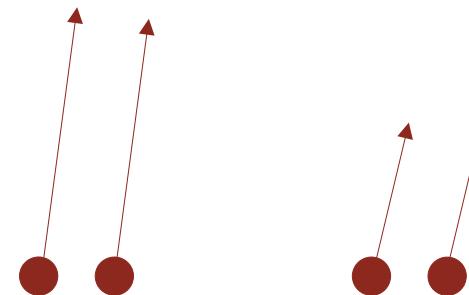
Motion coherence



Small  $\beta$

Large  $\beta$

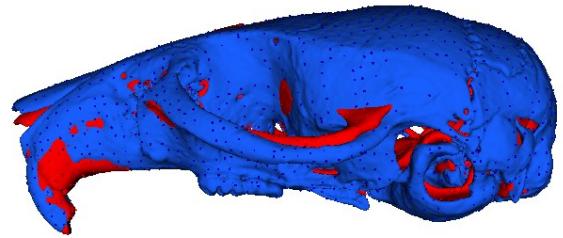
Displacement



Small  $\alpha$

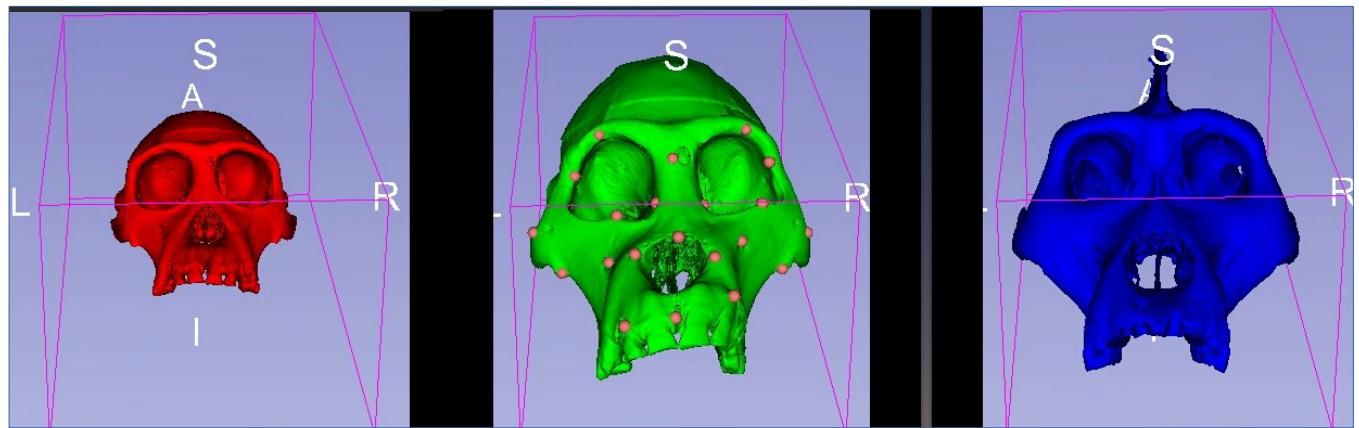
Large  $\alpha$

# Deformable – Structure specific



Small  $\beta = 2$

Large  $\alpha = 2$



Large  $\beta = 14$

Small  $\alpha = 0.05$