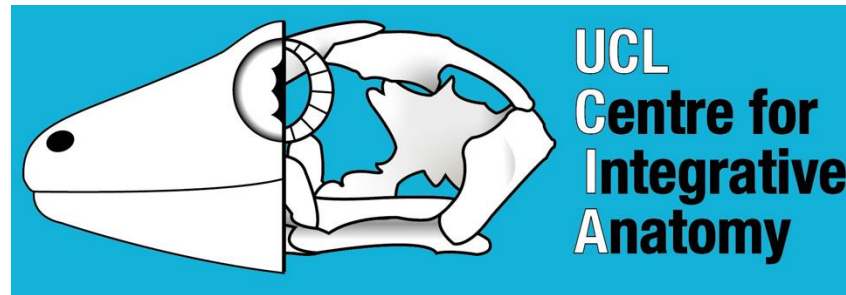
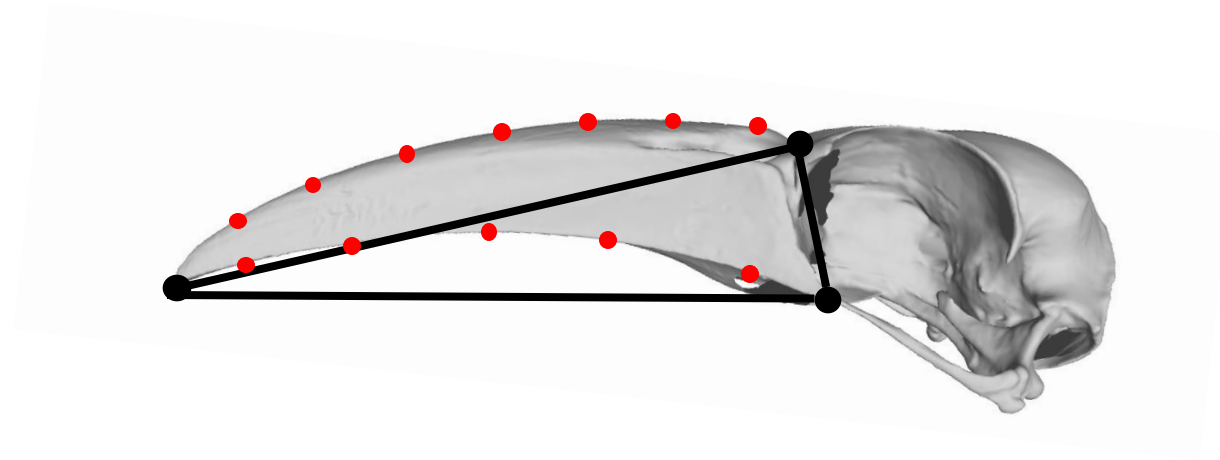
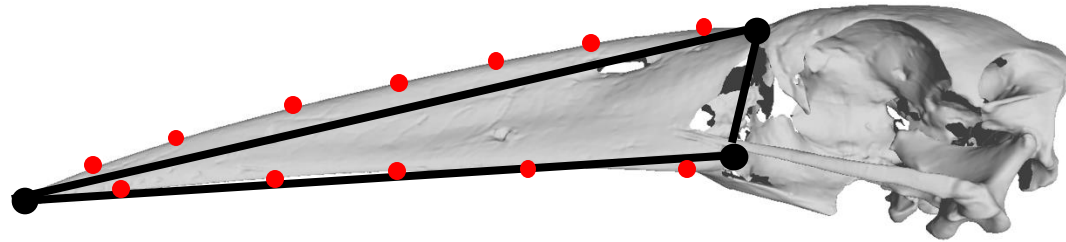


Advanced Shape Analysis

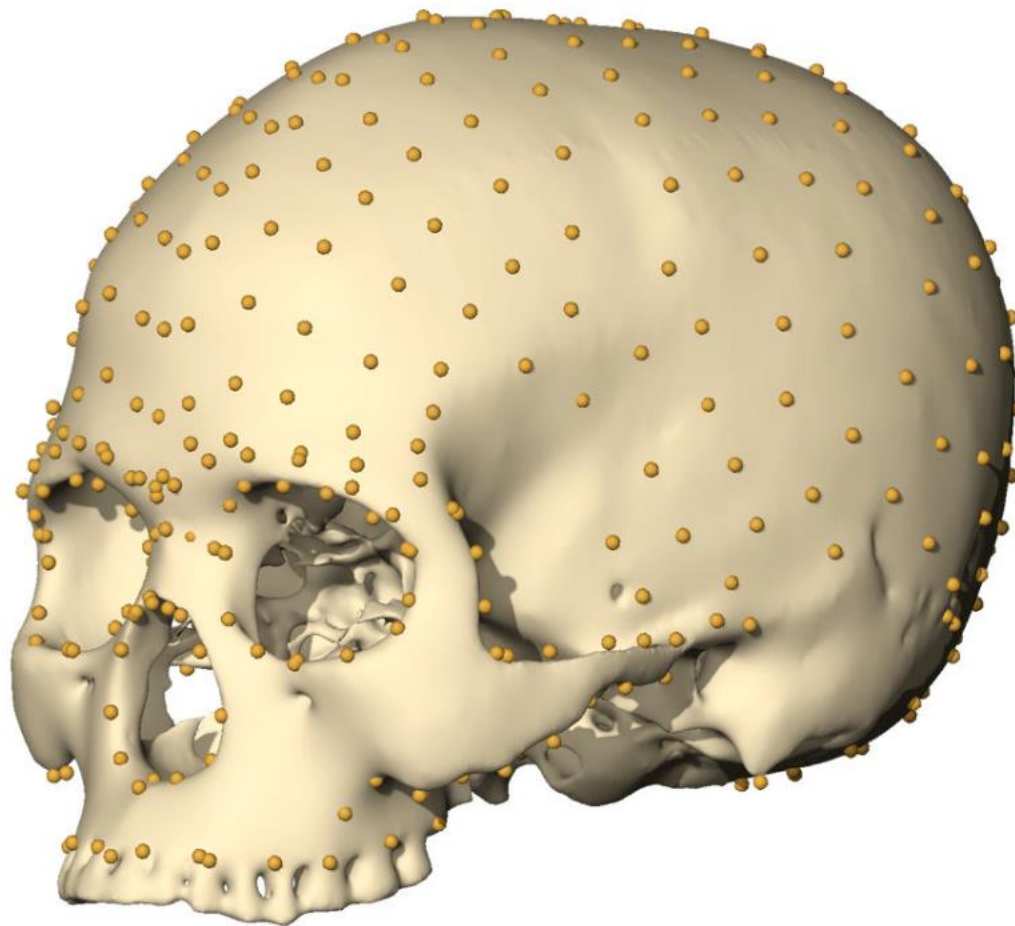
Ryan N. Felice



Sliding Semilandmarks

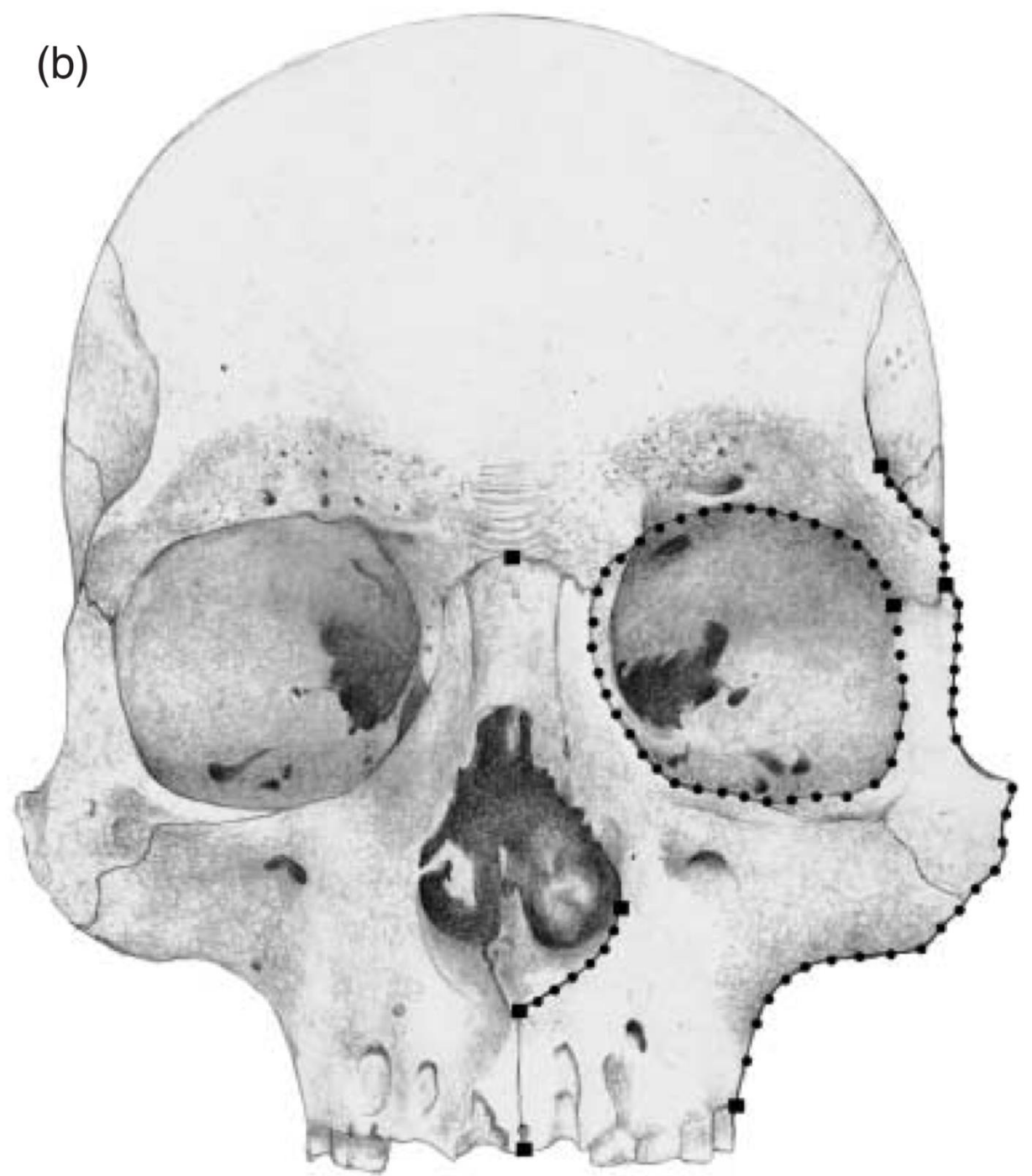


Surface Semi-landmarks



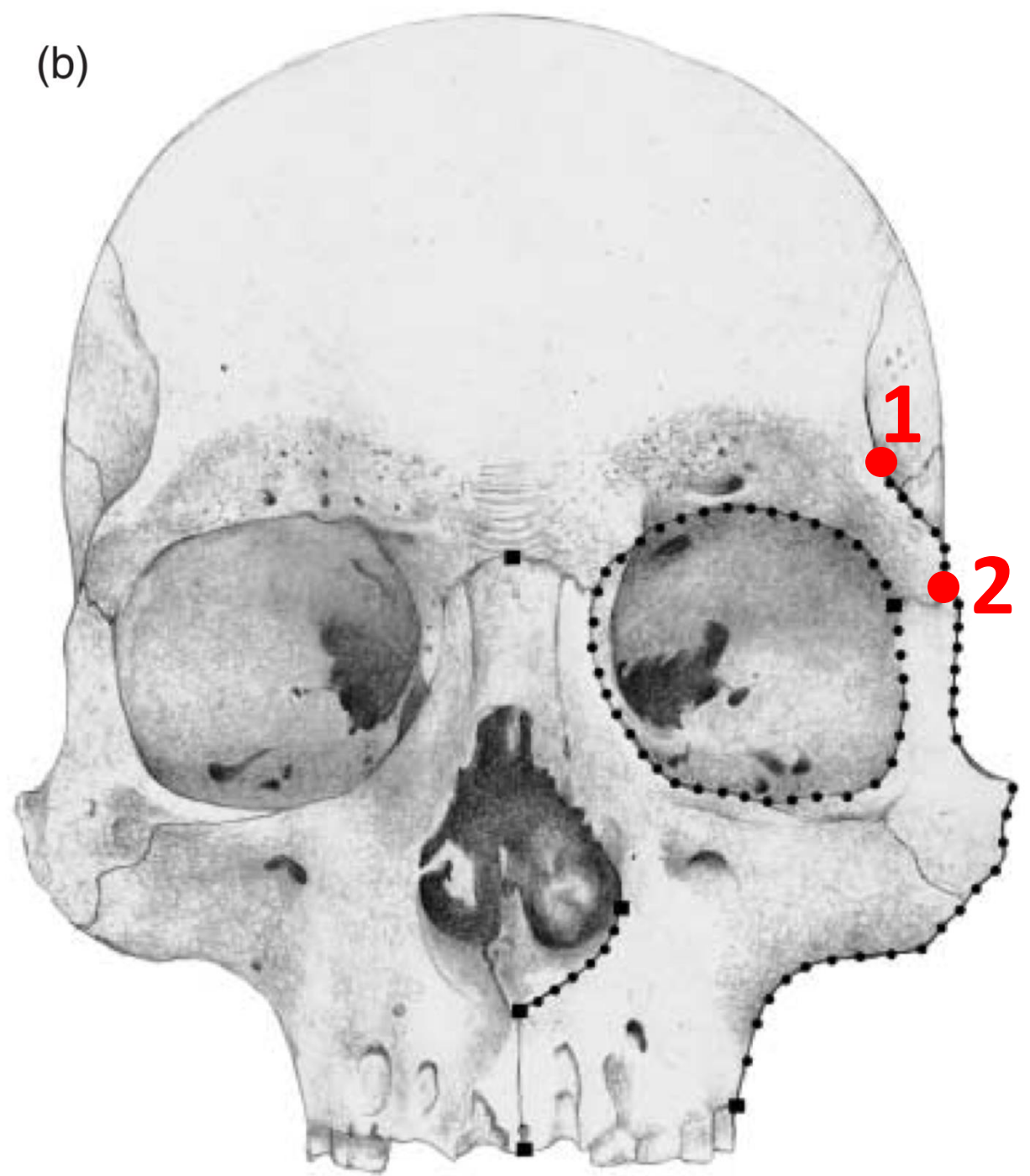
Sliding Semilandmarks

(b)



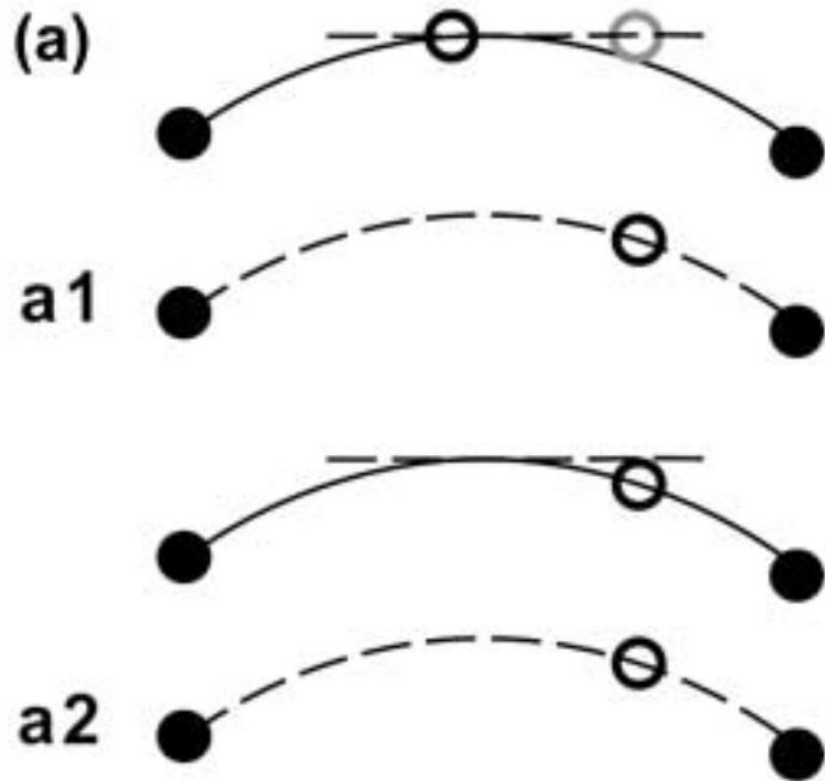
Sliding Semilandmarks

(b)

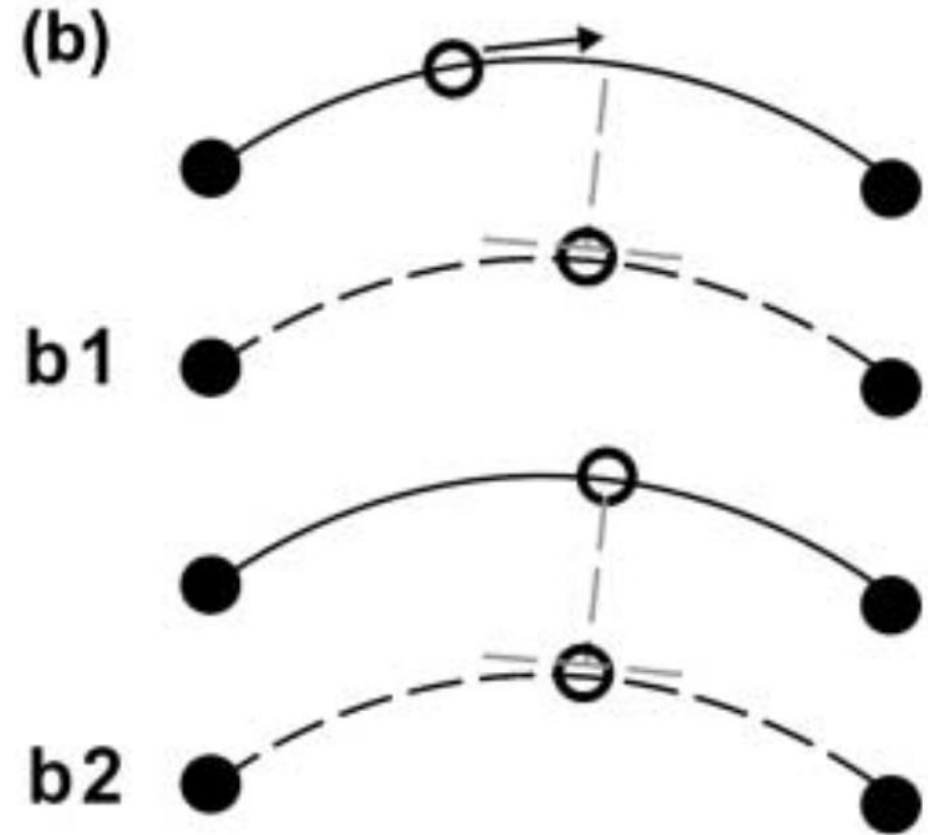


Sliding Semilandmarks

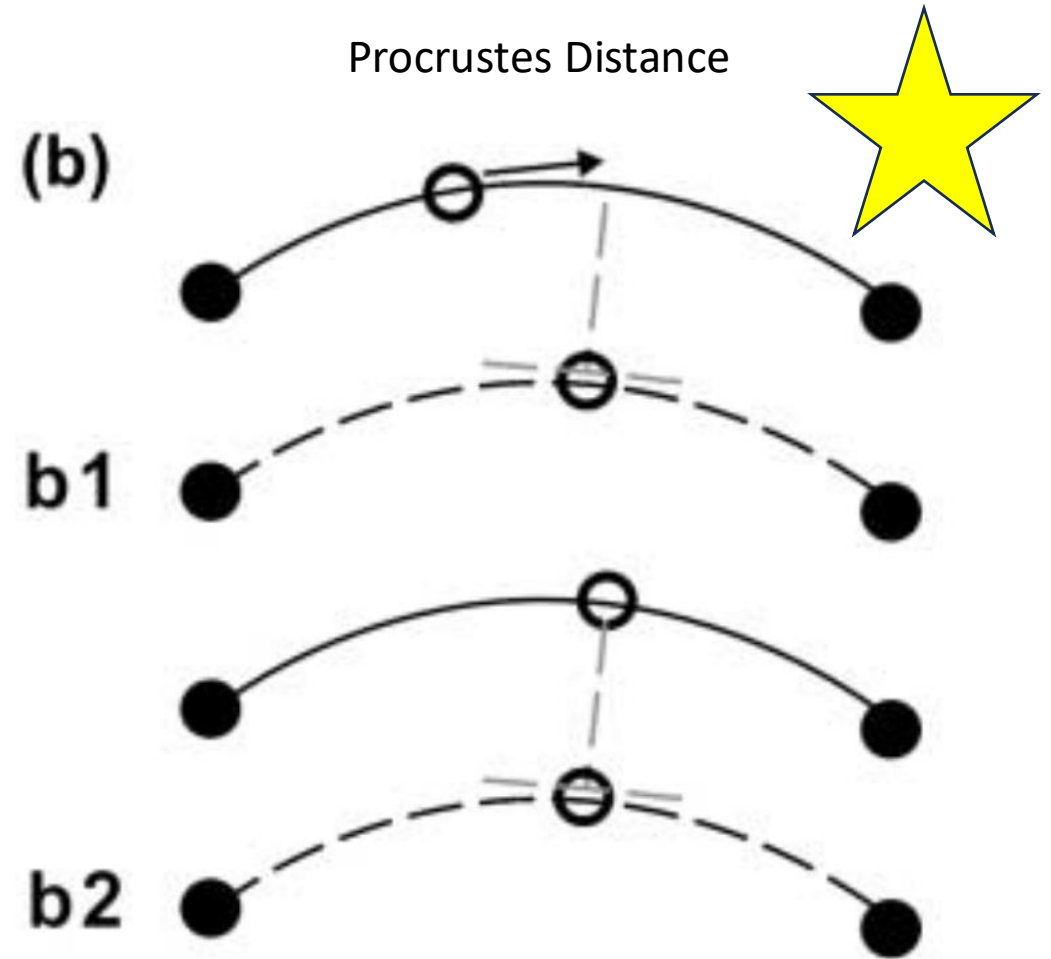
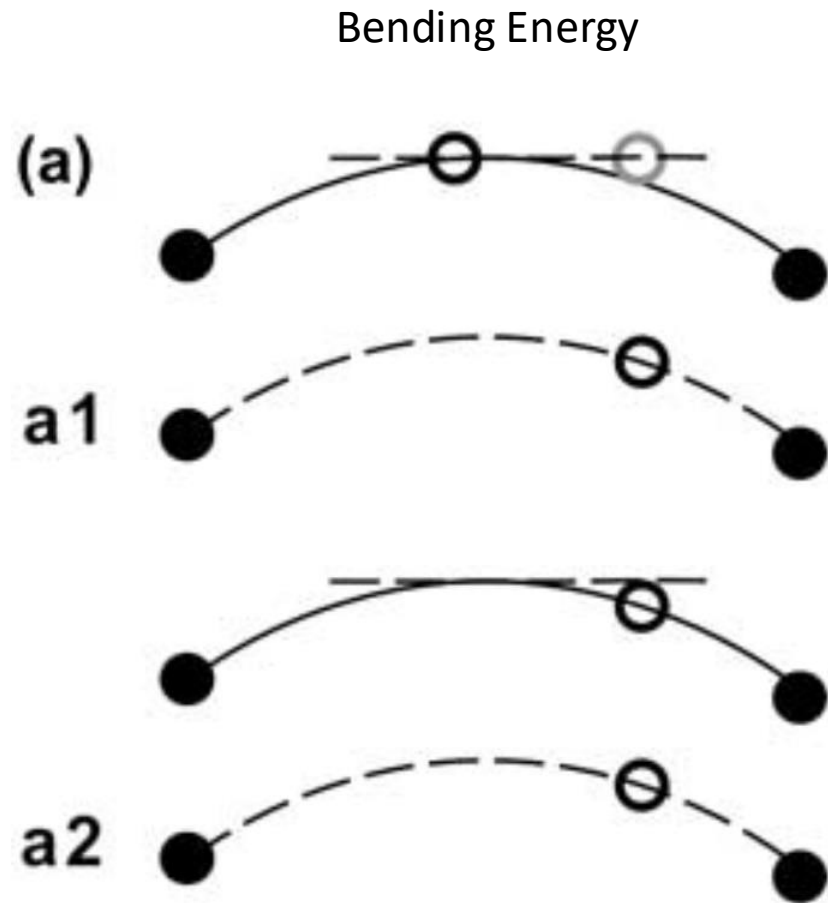
Bending Energy



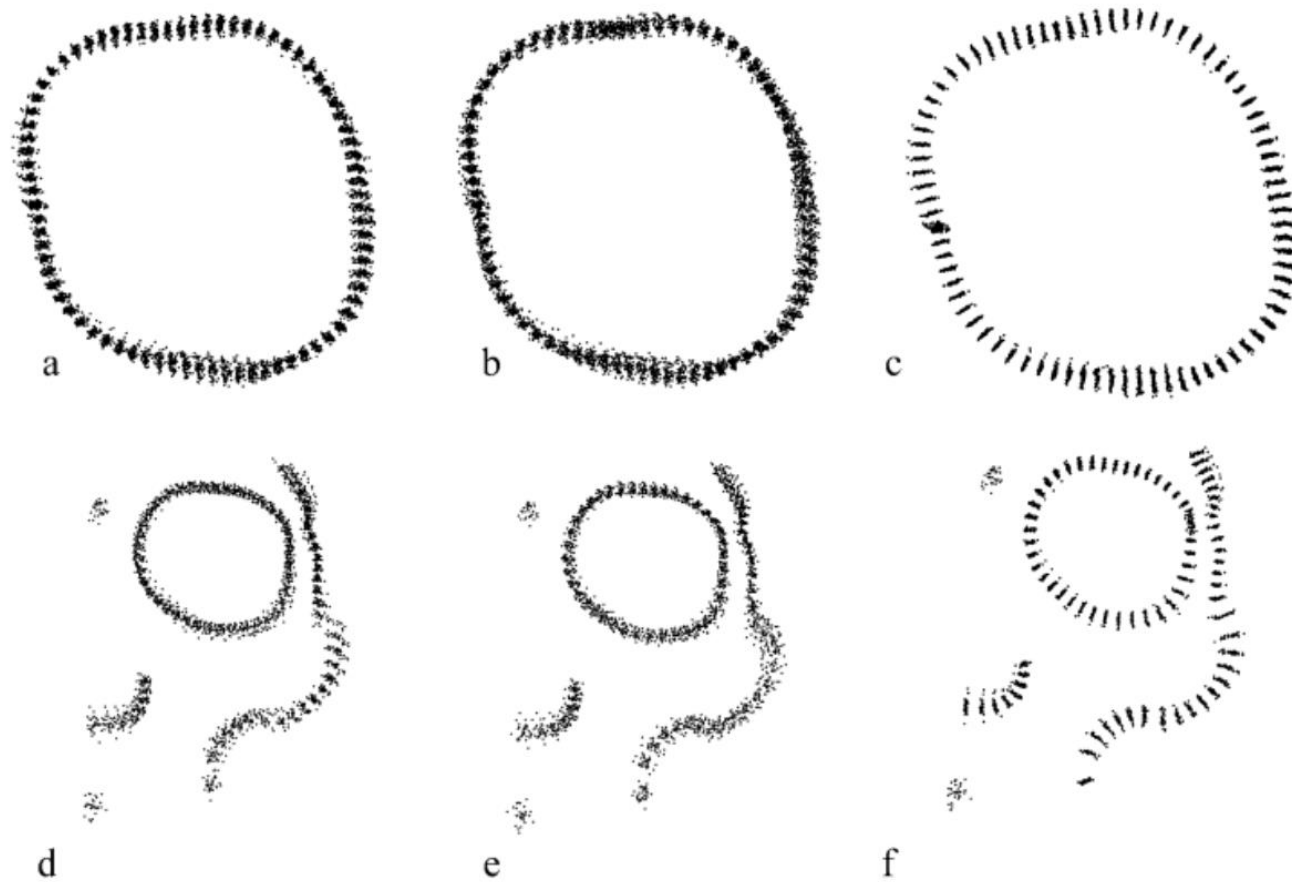
Procrustes Distance



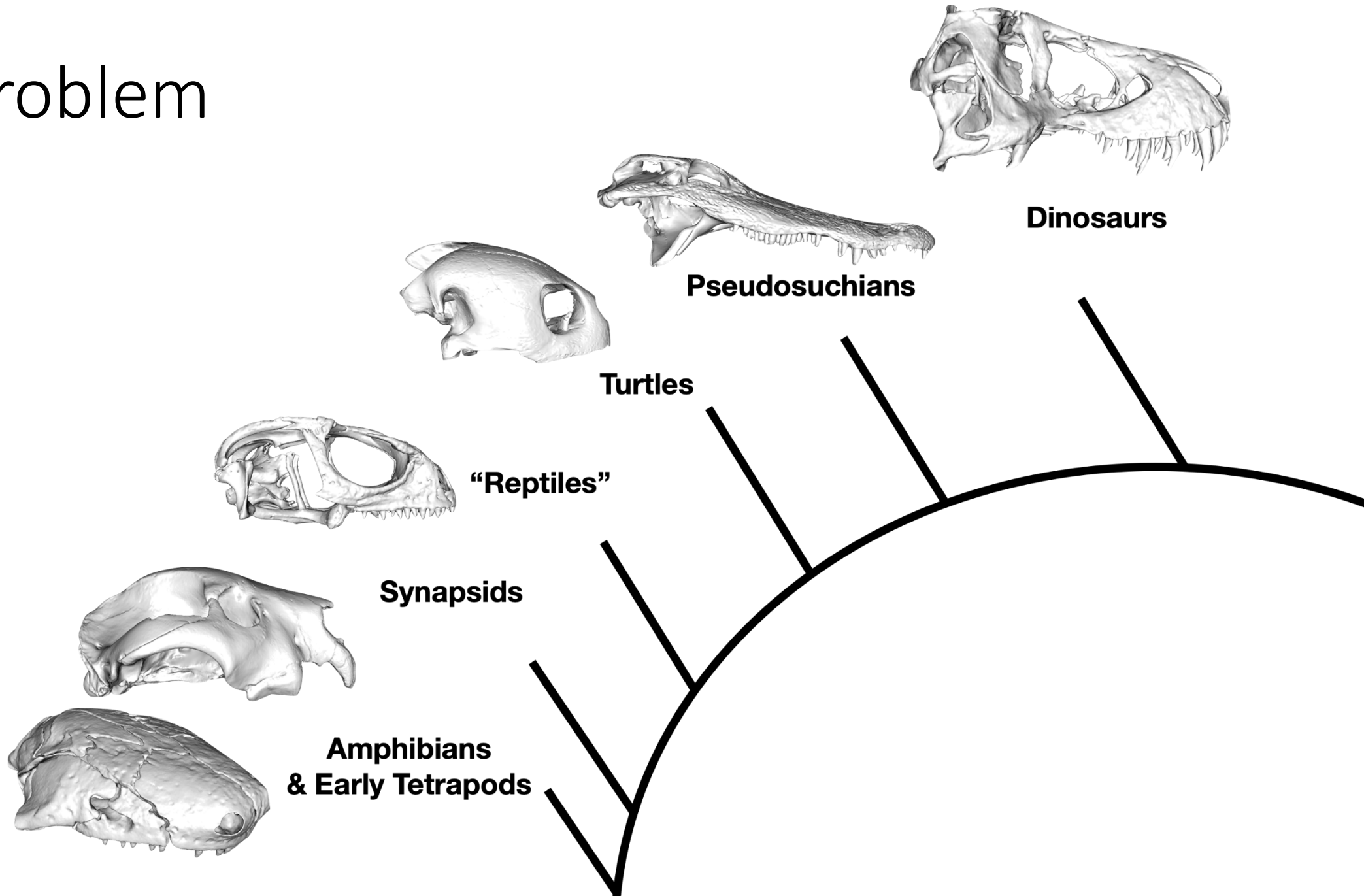
Sliding Semilandmarks

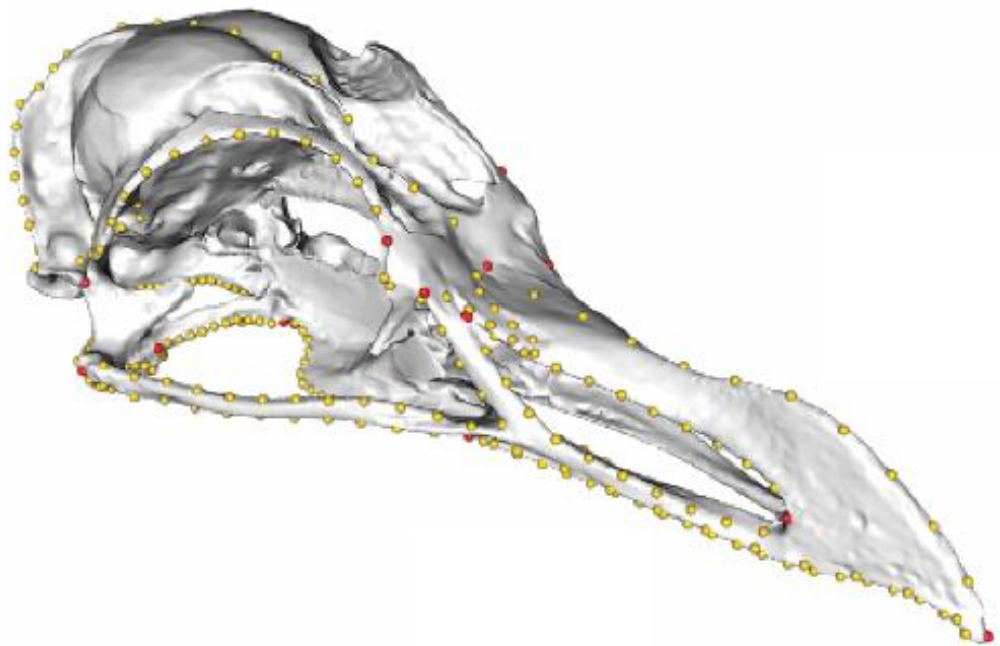


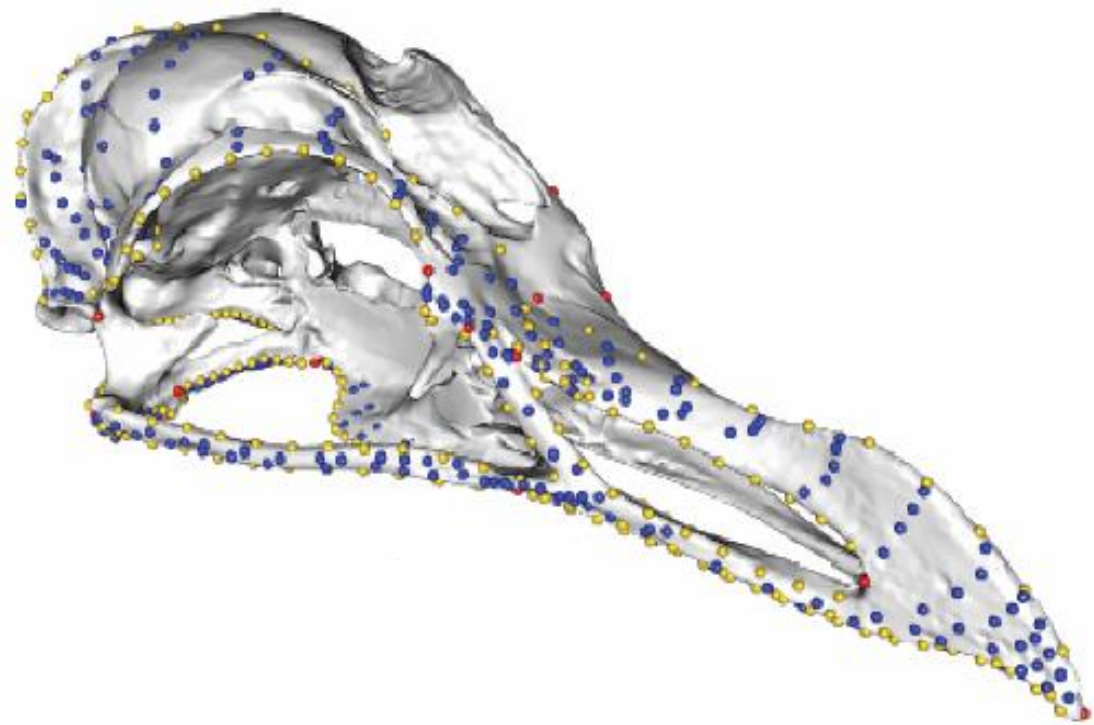
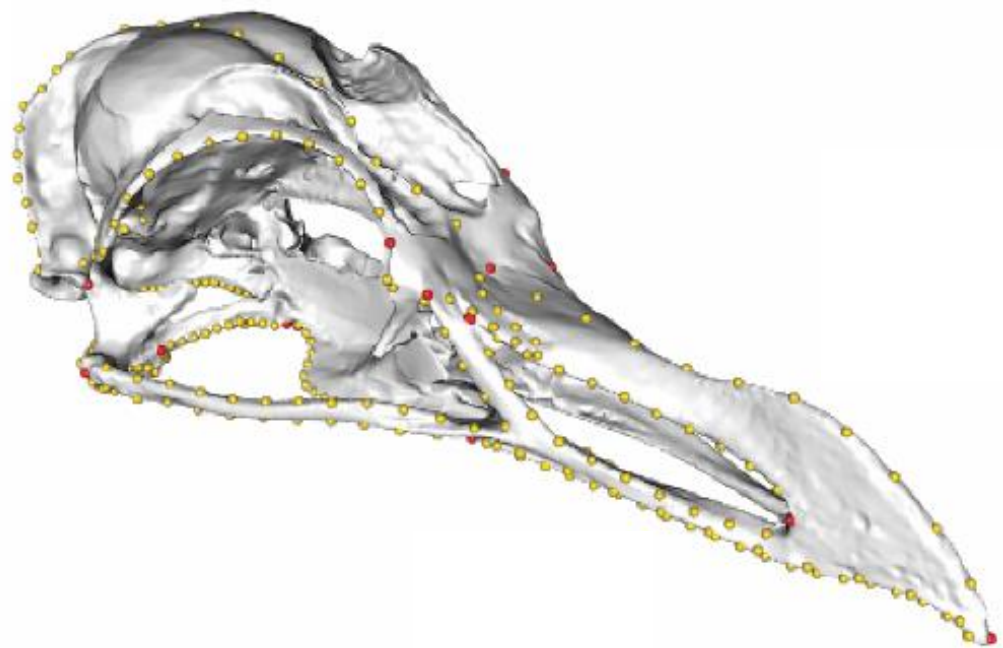
Sliding Semilandmarks



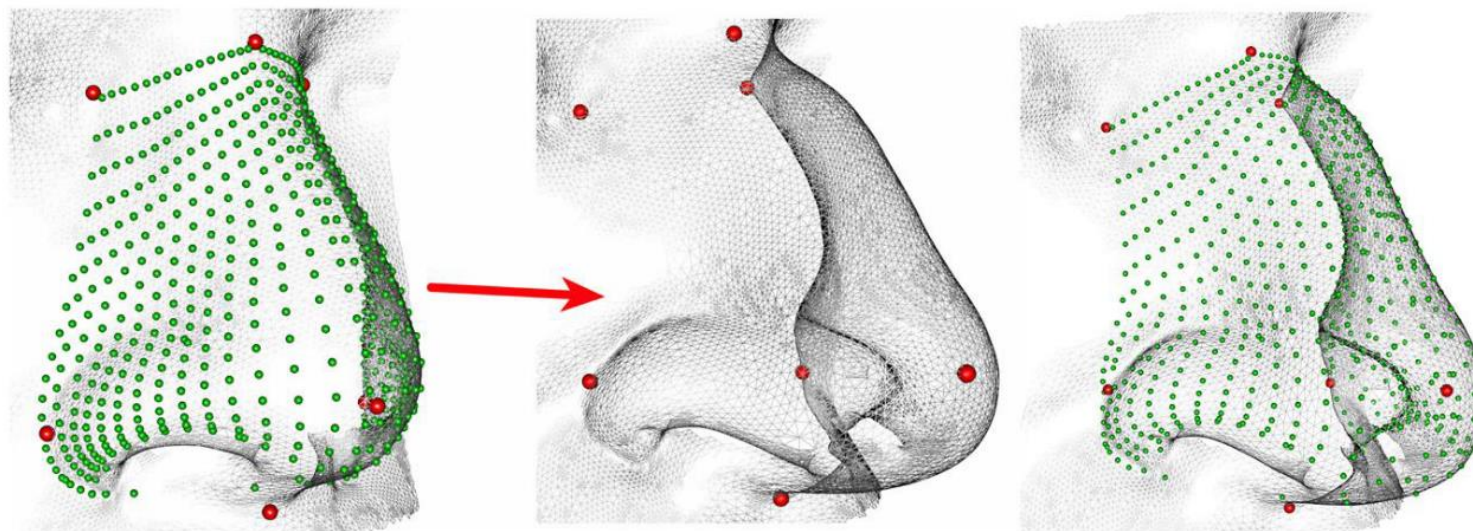
The Problem



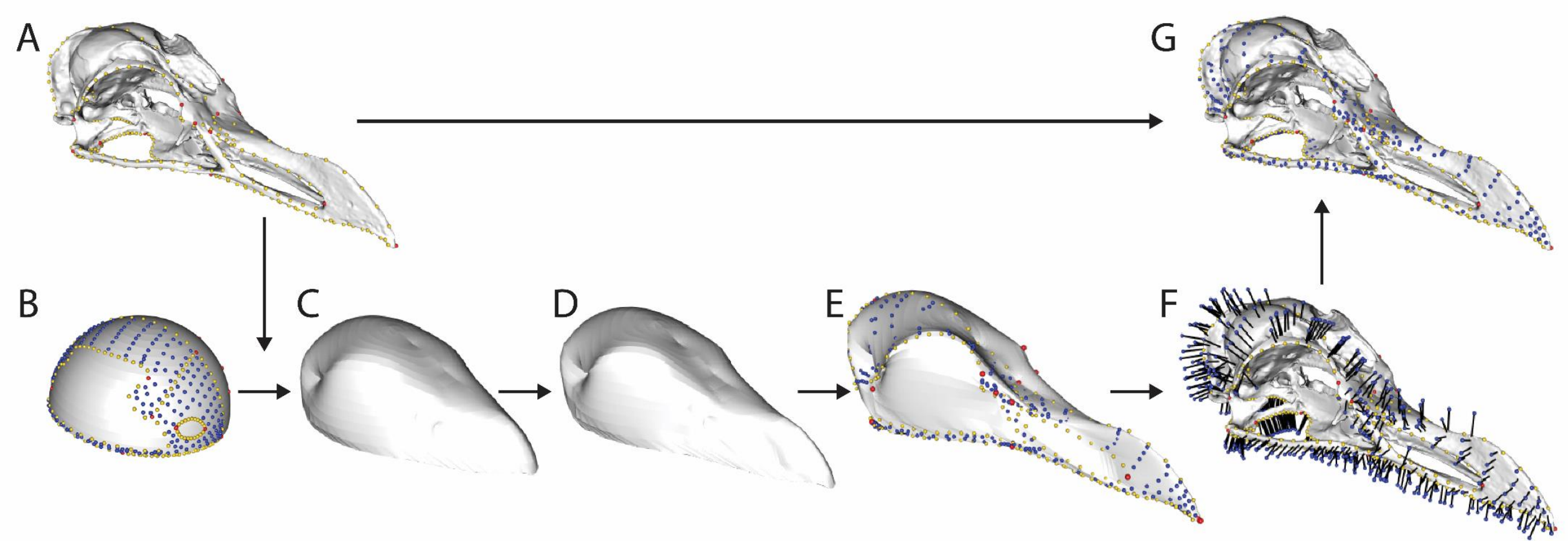


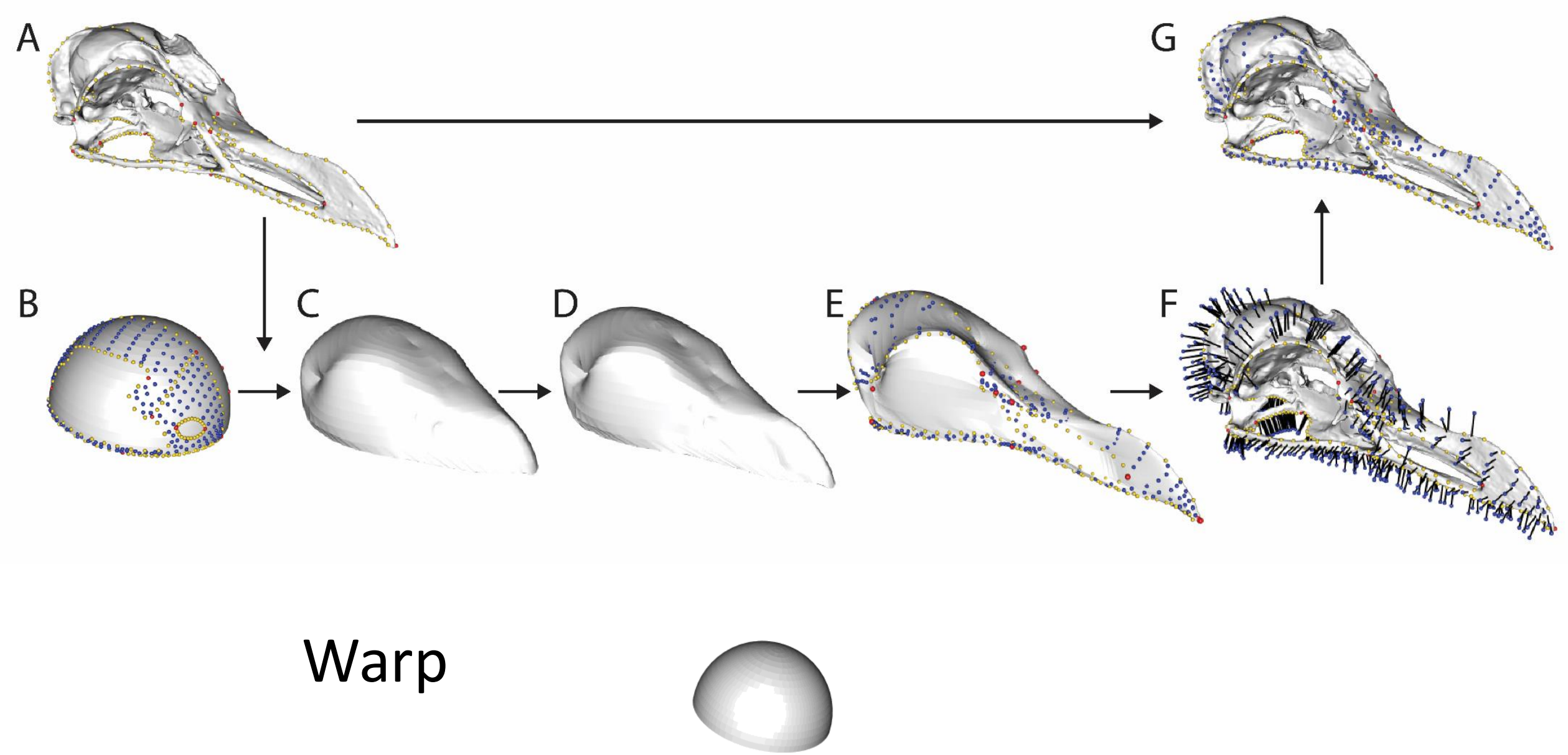


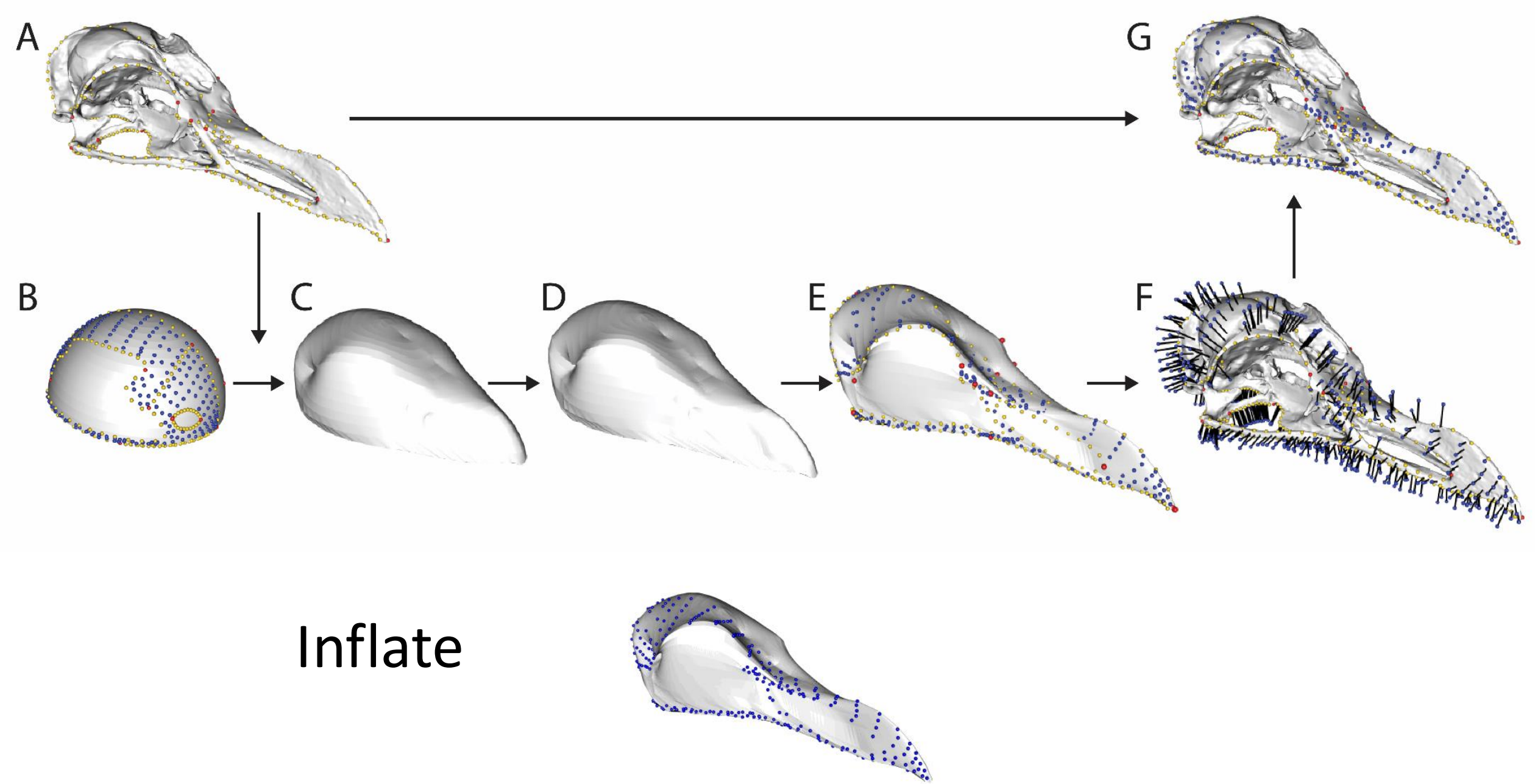
Semi-automated patching

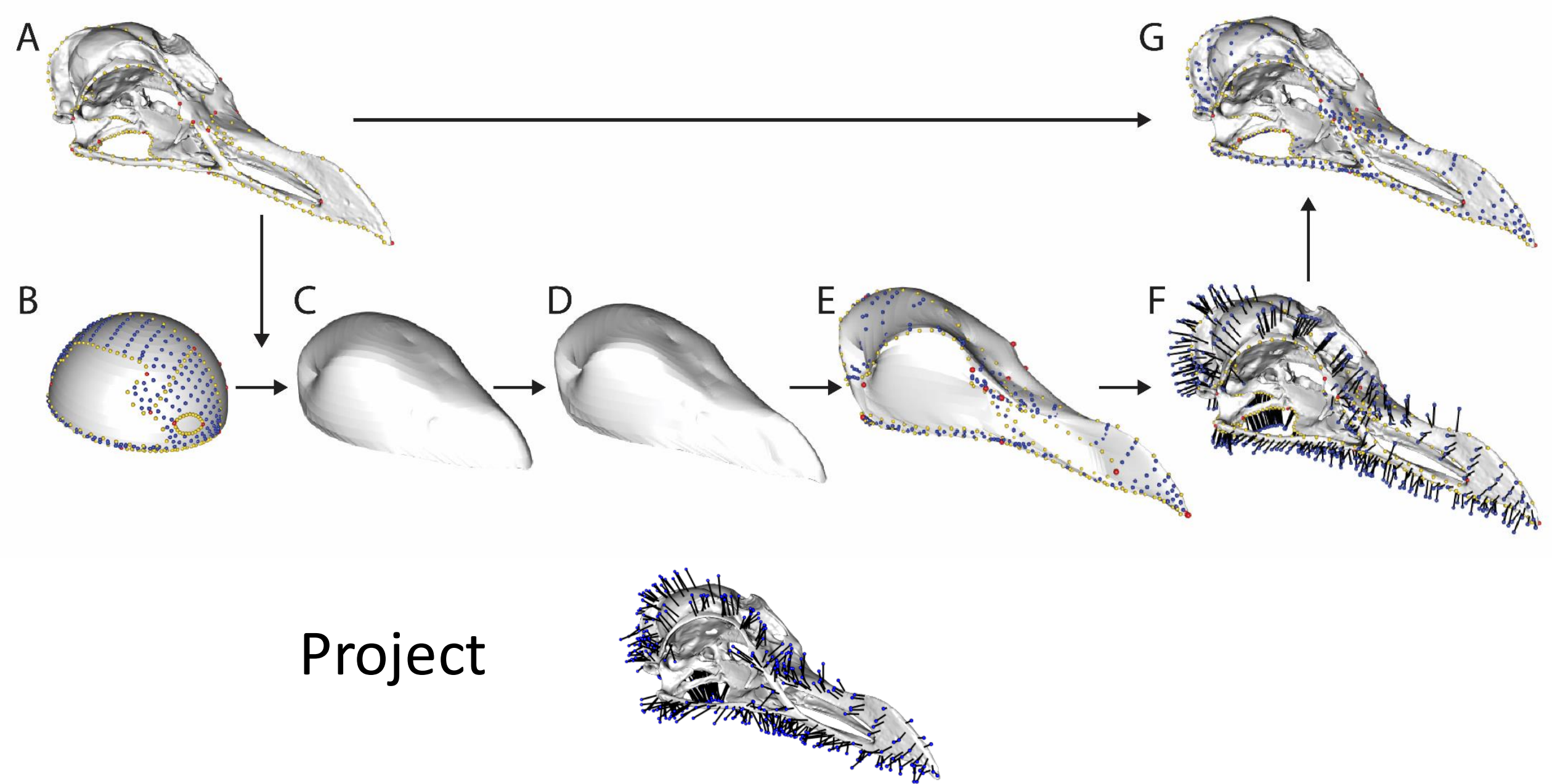


Morpho R package





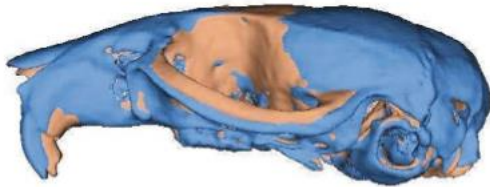




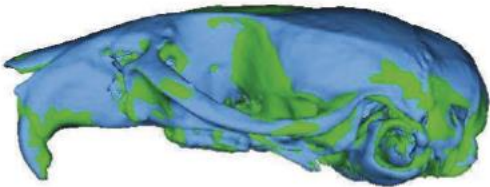
(a)



(b)



(c)



(d)



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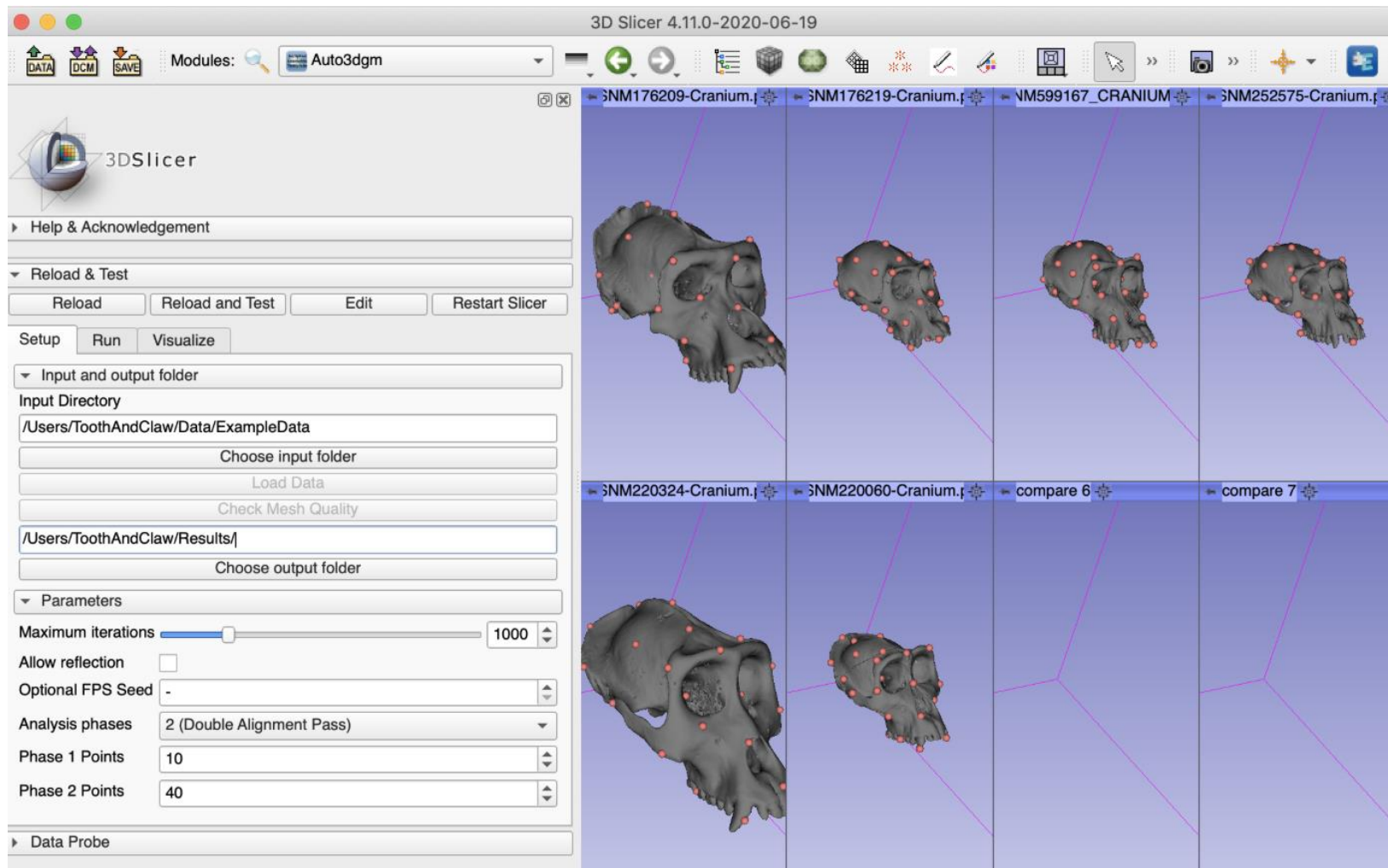
RESEARCH ARTICLE

Methods in Ecology and Evolution



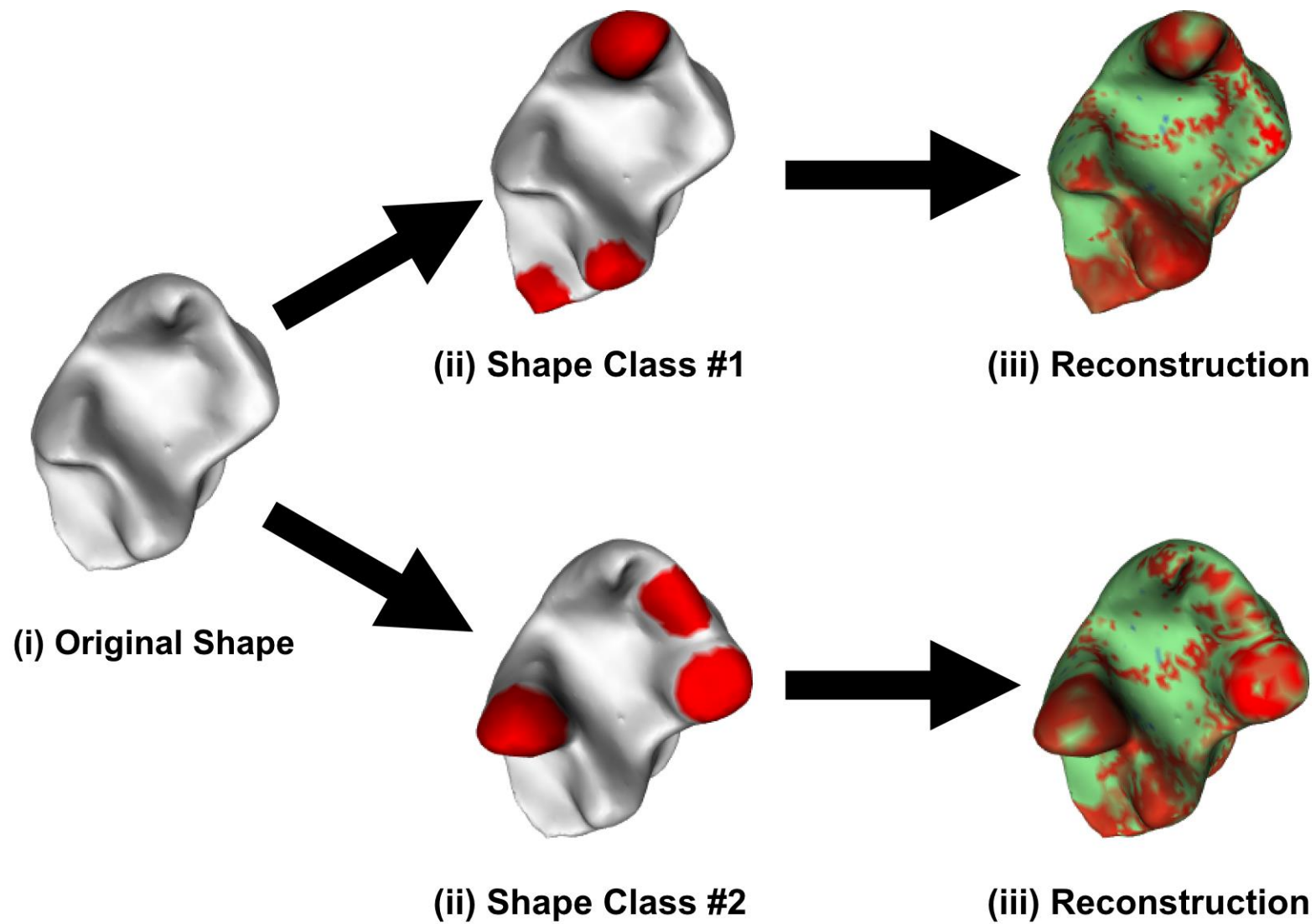
ALPACA: A fast and accurate computer vision approach for automated landmarking of three-dimensional biological structures

Arthur Porto^{1,2}  | Sara Rolfe^{3,4} | A. Murat Maga^{4,5} 



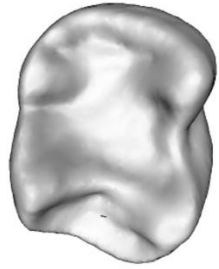
Pseudolandmarks-
Auto3dGm

<https://toothandclaw.github.io>



(a) Input 3D Shapes

Data from Species #1



Outcome: $y_i = 0$

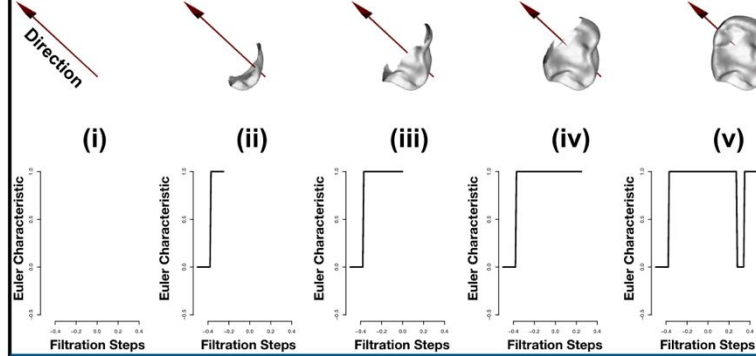
Data from Species #2



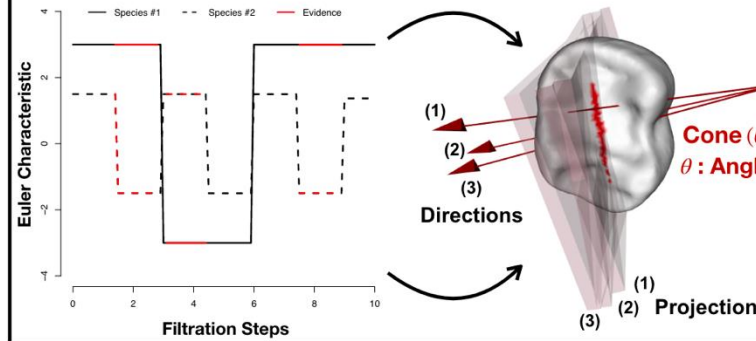
Outcome: $y_i = 1$

(b) Derive Topological Summary Statistics

Steps in Filtration Process:

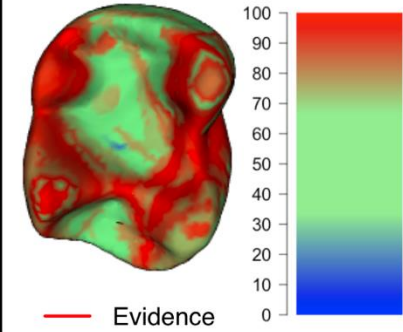


(c) Variable Selection and Reconstruction

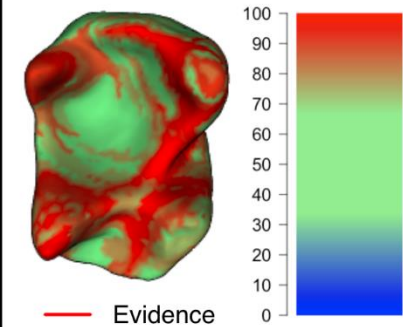


(d) Visualize Enrichment

Enrichment in Species #1



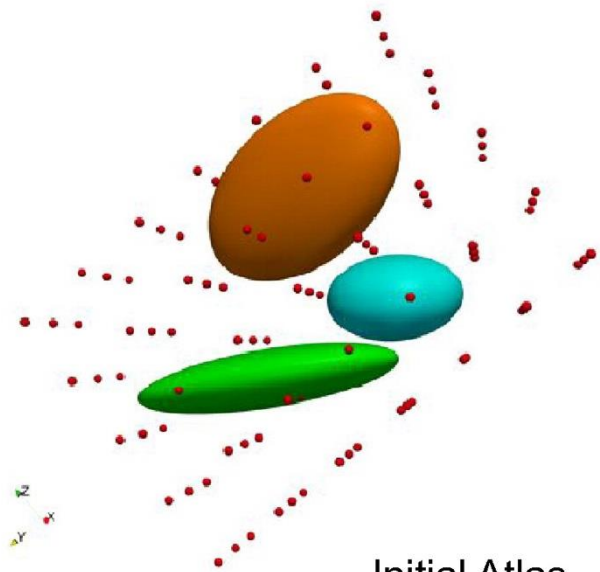
Enrichment in Species #2



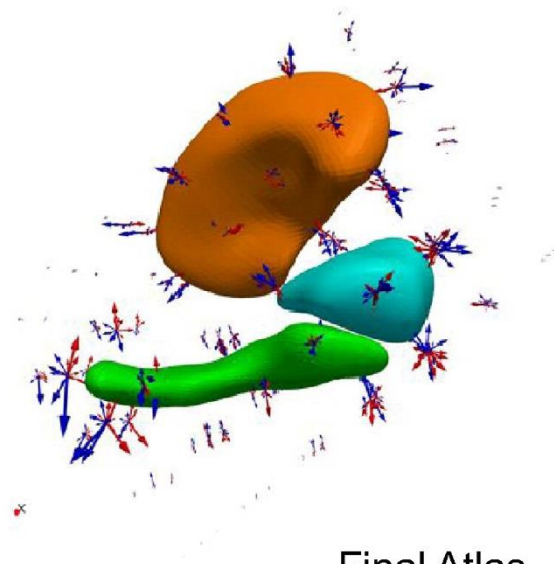
¹ **A Statistical Pipeline for Identifying Physical Features that**
² **Differentiate Classes of 3D Shapes**

³
⁴ Bruce Wang^{1,2,*}, Timothy Sudijono^{3,*}, Henry Kirveslahti^{4,*}, Tingran Gao⁵, Douglas M. Boyer⁶, Sayan
⁵ Mukherjee^{4,7-9,†}, and Lorin Crawford^{10,†}

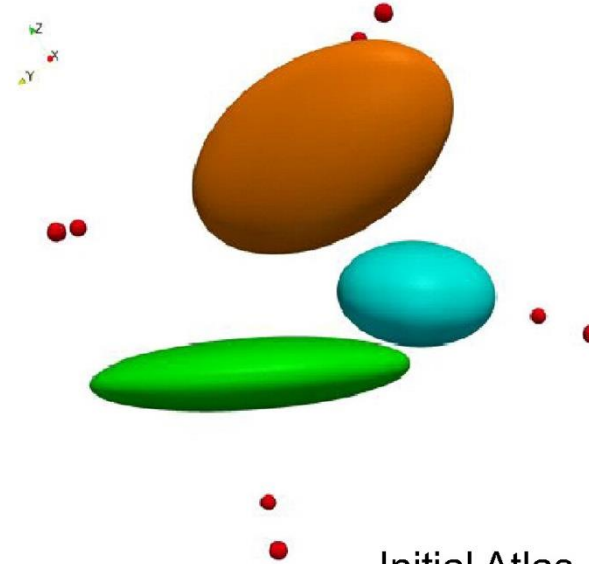
Deterministic Atlas Analysis



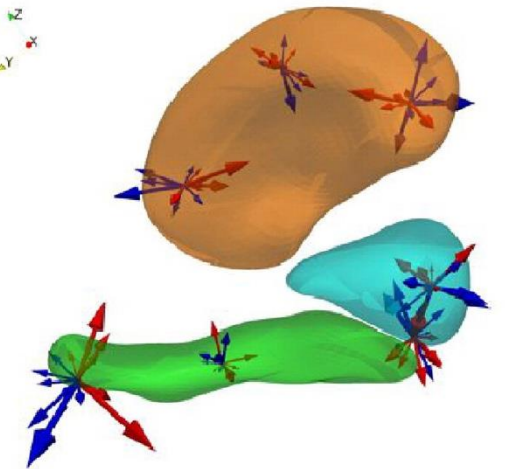
Initial Atlas



Final Atlas

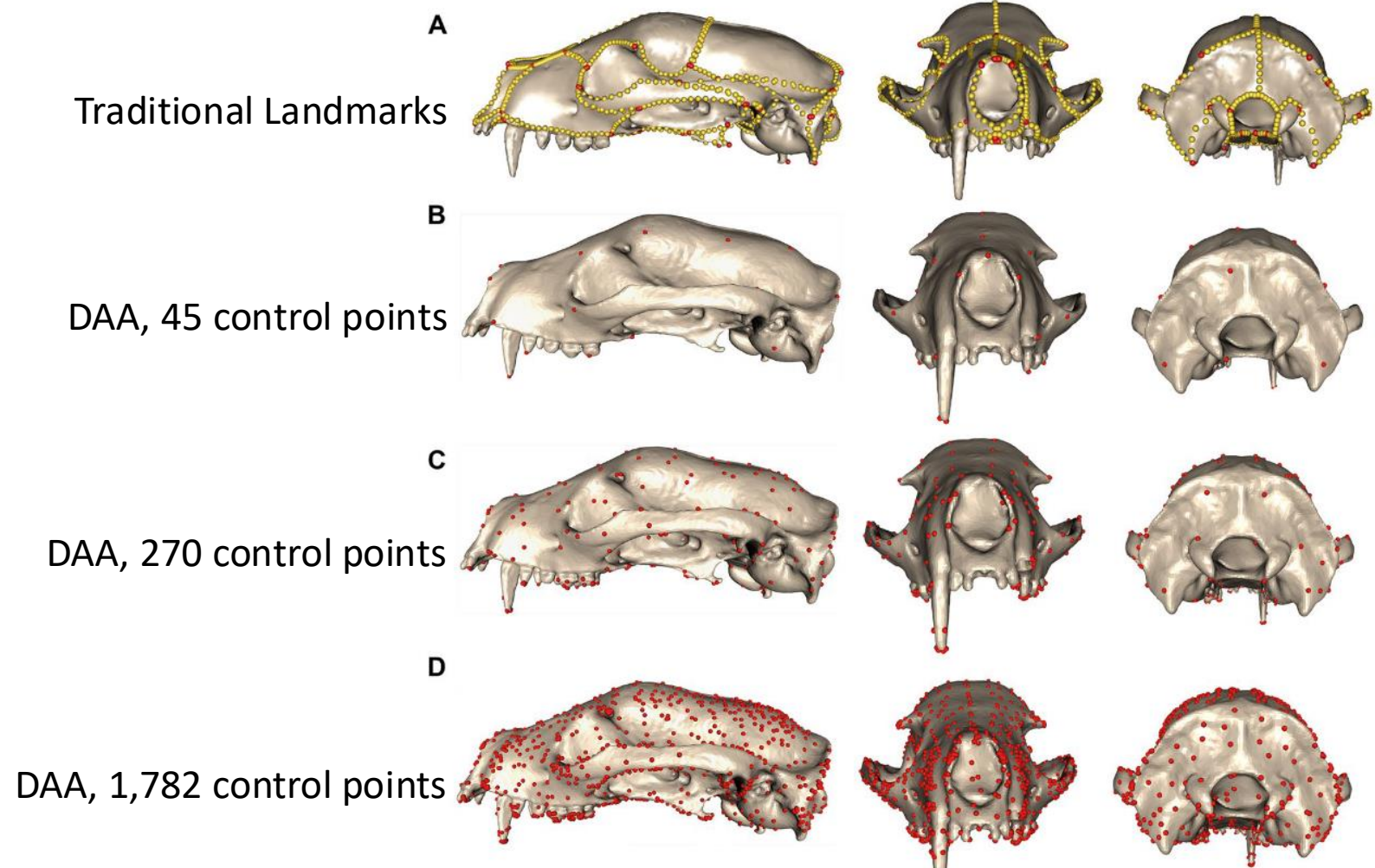


Initial Atlas



Final Atlas

Deterministic Atlas Analysis



RESEARCH

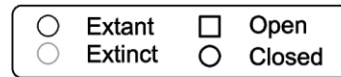
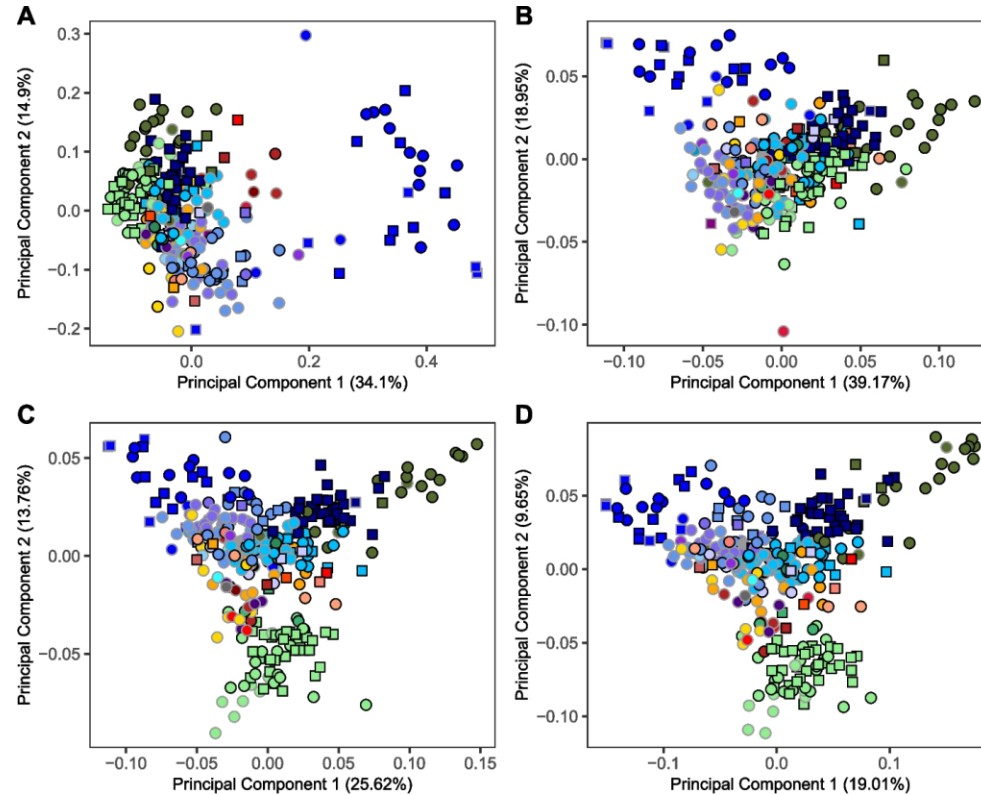
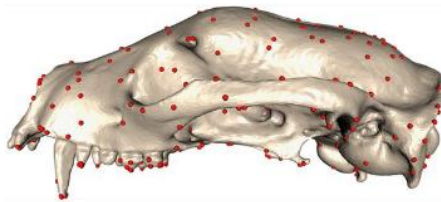
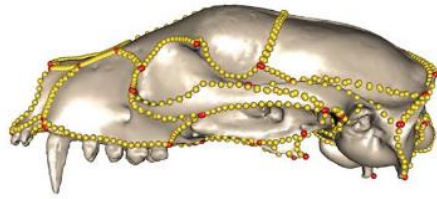
Open Access

Assessing the application of landmark-free morphometrics to macroevolutionary analyses

James M. Mulqueeney^{1,2*}, Thomas H. G. Ezard¹ and Anjali Goswami²



Deterministic Atlas Analysis



Stem

- Zalambdalestidae
- Cimolestia
- Leptictida

Euarchontaglires

- Dermoptera
- Scandentia
- Primates
- Rodentia
- Lagomorpha

Xenarthra

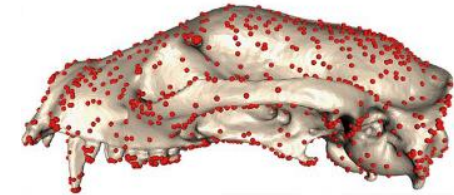
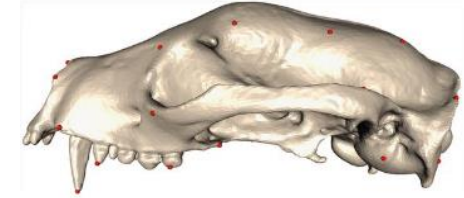
- Cingulata
- Pilosa

Afrotheria

- Afrosoricida
- Macroscelidea
- Tubulidentata
- Hyracoidea
- Proboscidea
- Desmostylia
- Sirenia
- Embrithopoda

Laurasiatheria

- Chiroptera
- Eulipotyphla
- Acreodi
- Amblypoda
- Artiodactyla
- Cetacea
- Litopterna
- Notoungulata
- Astrapotheria
- Perissodactyla
- Creodonta
- Carnivora
- Pholidota



How much firepower do I need?



VS.



Modelling High Dimensional Data

- What do we mean by high dimensional data?
- N = number of observations
- P = number of variables

	Trait 1	Trait 2	Trait 3
Species 1			
Species 2			
Species 3			
Species 4			
Species 5			
Species 6			
Species 7			

	Trait 1	Trait 2	Trait 3	Trait 4	Trait 5	Trait 6
Species 1						
Species 2						
Species 3						
Species 4						

Modelling High Dimensional Data

- Who cares?
- Everything is linear models
- Estimating F-statistic and likelihood of linear models with $p > N$ is impossible (you can't invert a matrix that has $p > N$)
- “Curse of dimensionality”

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \sigma$$

$$F = \frac{\hat{\boldsymbol{\beta}}^\top (\mathbf{X}^\top \mathbf{X}) \hat{\boldsymbol{\beta}} / p}{\hat{\sigma}^2}$$

The Solution:

- RRPP: Residual randomization in permutation procedure
- F-statistic (and significance) is computed by randomizing the residuals of the model and comparing to a null

$$F = \frac{MS_{Model}}{MS_{residuals}}$$

MS = Mean squares

The Solution:

- RRPP: Residual randomization in permutation procedure
- F-statistic (and significance) is computed by randomizing the residuals of the model and comparing to a null

Recommendations:

- 1) Always think about what methods are appropriate for your data
- 2) Use `procD.lm` or `procD.pgls` functions in `geomorph` or `lm.rrpp` in the RRPP package



<https://doi.org/10.1111/2041-210X.13029>

