# Viewbox Tutorial Digitization error: Procrustes ANOVA

#### Procrustes ANOVA

Calculation of the digitization error for 3D specimens, as described here, is based on:

- Klingenberg, C. P. MorphoJ: an integrated software package for geometric morphometrics. *Mol. Ecol. Resour.* 11, 353-357 (2011).
- Fruciano, C. Measurement error in geometric morphometrics. *Dev. Genes Evol.* **226,** 139-58 (2016).

MorphoJ is required.

### Step 1 (Viewbox)

Digitize the specimens twice (intra-observer: one investigator, each specimen twice; inter-observer: two investigators, each specimen once) and save them using 'name' as the following:

- -SP001a
- -SP001b
- -SP002a
- -SP002b

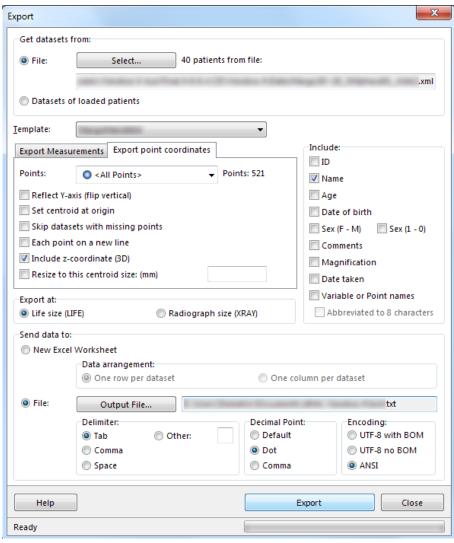
**—**...

where 'a' and 'b' signify the two repeated digitizations.

Note that we pad numbers with '0' (i.e. SP001 and not SP1), so that all names have the same length. This is important later.

### Step 2 (Viewbox)

Export the specimens to a text file using Tools – Export:



Make sure 'Include z-coordinate' is selected, if your data are 3D.

Make sure 'Name' is selected.

Note the settings for the output file:

- Delimiter: Tab
- Decimal Point: Dot (try Comma if Dot does not work when importing into MorphoJ)
- Encoding: ANSI or UTF-8 no BOM

### Step 3 (MorphoJ)

Open MorphoJ and create a Dataset using the file from the previous step:

#### File – Create New Dataset

Make sure you set 2 or 3 dimensions, depending on your data. Then select the txt file and click 'Create Dataset'.

Perform a Procrustes Fit:

**Preliminaries – New Procrustes Fit...** 

#### Step 3 - continued

Extract two classifiers. The first contains the name (e.g. SP001) and the second the repeat (a or b):

#### **Preliminaries – Extract new classifier from ID strings...**

Set the name and the first and last characters (1 and 5, because 'SP001' has 5

characters), click 'Execute':

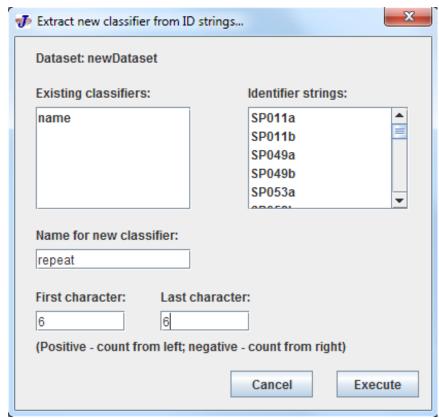
Dataset: newDataset		
Existing classifiers:	Identifier strings:	
	SP011a	•
	SP011b	
	SP049a	
	SP049b	
	00000	
Name for new classifier:	SP053a	~
name  First character: Last character: 5	naracter:	•
name  First character: Last ch	naracter:	•

#### Step 3 - continued

The second classifier is based on the a or b character:

Preliminaries – Extract new classifier from ID strings...

Set the name and the first and last characters (6 and 6, because 'a' and 'b' are always at the 6<sup>th</sup> position), click 'Execute':

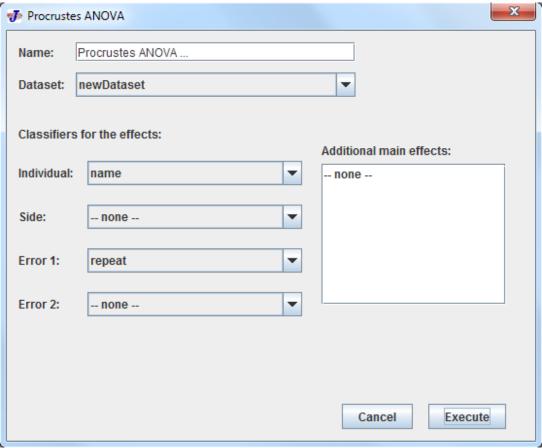


## Step 4 (MorphoJ)

Perform a Procrustes ANOVA:

**Variation – Procrustes ANOVA...** 

Set the 'repeat' classifier as 'Error 1':



### Step 5 (MorphoJ): Results

#### These appear at the end of the 'Results' tab:

```
Procrustes ANOVA: Procrustes ANOVA ...
Dataset: newDataset
Classifiers used for the Procrustes ANOVA:
Individuals: name
Error 1: repeat
Centroid size:
Effect
                                          df
                            MS
                                                              P (param.)
Individual 154155,651890 8113,455363 19
                                                     21240,51 <.0001
Error 1 7,639606 0,381980
                                          20
Shape, Procrustes ANOVA:
Effect
                            MS
                                          df
                                                              P (param.)
             SS
           0,13215122
                            0,0000044700 29564
                                                     242,04
Individual
                                                              < .0001
             0,00057473
                            0,0000000185 31120
Error 1
```

Based on the book "Geometric Morphometrics for Biologists. A Primer, 2nd Edition, Authors: Miriam Zelditch, Donald Swiderski, H. David Sheets", page 258, we calculate reliability of shape as:

```
Ind_var = (MSindividual - MSerror)/2 = (44700 - 185)/2 = 22257.5
Total = 22257.5 + 185 = 22442.5
Ratio = 22257.5 / 22442.5 = 0.992 = repeatability
```