This folder contains the R scripts, nexus file and data used for the phylogenetic comparative methods. When loading the R scripts you’ll notice specific files for analyses. These are also included in this folder.

Basic data:

N=154 LANDMARK Shark MC L OUTPUT (incl. E. zantedeschia).dta is the file containing the 3D landmarkcoordinates generated by Landmark Editor

N=154 MORPHOLOGIKA INPUT Shark MC L (incl. E. zantedeschia).txt is the file where all the coordinates are converted into a MORPHOLOGIKA format that can be read in MORPHOJ

MC L is the MORPHOJ file with all the projects I’ve created to generate PCscore and CVscores

N=154 Classifiers MC dataset (incl. E. zantedeschia).txt is the file containing all the classifiers for each specimen such as species name, family, order, habitat, preychoice and literature references.