## nature portfolio

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## **Reporting Summary**

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Confirmed
	$\square$ The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
X	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	The statistical test(s) used AND whether they are one- or two-sided  Only common tests should be described solely by name; describe more complex techniques in the Methods section.
	A description of all covariates tested
	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i> ) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>
	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
$\times$	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
	$\boxtimes$ Estimates of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.

## Software and code

Policy information about availability of computer code

Data collection

Landmark Editor (Version 3.0) for landmark capture. http://vertlife.org/sharktree/ for phylogenetic information.

Data analysis

R software (https://www.r-project.org/), in TreeAnnotator version 1.8.2, BayesTraits v 3.0.2 (http://www.evolution.rdg.ac.uk). All relevant packages are described in the Methods section and in the Script in Code Availability section.

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

## Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

All data sources used in the study is indicated in the methods section. The ply files for the shark species can be found at https://doi.org/10.6084/m9.figshare.c.3662366.v1 and 10.18563/journal.m3.133 for C. milii.

Human rese	arch part	icipants		
Policy information about studies involving human research participants and Sex and Gender in Research.				
Reporting on sex	and gender	Not applicable		
Population characteristics		Not applicable		
Recruitment		Not applicable		
Ethics oversight		Not applicable		
Note that full informa	ation on the appi	roval of the study protocol must also be provided in the manuscript.		
Field and	oific ro	porting		
Field-spe		•		
Life sciences		is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.  Behavioural & social sciences  Ecological, evolutionary & environmental sciences		
		Behavioural & social sciences		
Life scier	nces st	udy design		
All studies must dis	sclose on these	points even when the disclosure is negative.		
Sample size	The sample wa	as chosen based on specimen availability in the collections and available CT-Scan Data at the moment.		
Data exclusions	The quality of	the surface reconstruction was considered to exclude specimens.		
Replication		lyses regarding the outliers from the group were conducted. In the case of morphological disparity the groups were also subject cs to check the consistency of the results.		
Randomization	We used a random subsample of 100 trees from the posterior probability to estimate the morphological evolutionary rates and disparity through time.			
Blinding	Blinding does r	not apply here. But different grouping was tested to check for consistency.		
Reportin	a for si	pecific materials, systems and methods		
<u> </u>		about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material,		
•		o your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.		
Materials & ex	perimental s	systems Methods		
n/a Involved in the study		n/a Involved in the study		
Antibodies		ChIP-seq		
Eukaryotic cell lines		Flow cytometry		
	Palaeontology and archaeology MRI-based neuroimaging  MRI-based neuroimaging			
Clinical dat	Animals and other organisms			
Dual use research of concern				
Animals and other research organisms				
, aminais and	. Other rec	zear on organisms		

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals

No laboratory animals were used

Wild animals	All specimens are housed in collections as described in Brazeau, Martin; Kamminga, Pepijn; De Bruin, Paul W.; Geleijns, Jacob (2017): X-ray computed tomography library of shark anatomy and lower jaw surface models. figshare. Collection. https://doi.org/10.6084/m9.figshare.c.3662366.v1 and Dearden, R., Mansuit, R., Cuckovic, A., Herrel, A., Didier, D., Tafforeau, P., & Pradel, A. (2021). 3D models related to the publication: The morphology and evolution of chondrichthyan cranial muscles: a digital dissection of the elephantfish Callorhinchus milii and the catshark Scyliorhinus canicula. MorphoMuseuM.
Reporting on sex	Does not apply. The specimens were selected based on the availability in the museums collections and the literature.
Field-collected samples	No field-collected samples were used.
Ethics oversight	No ethical approval applies since the project involved exclusively museum specimens and non destrutive procedure to analyse their anatomy.

Note that full information on the approval of the study protocol must also be provided in the manuscript.