

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 [Editorial Policies](#) and the [Editorial Policy Checklist](#).

Statistics

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
<input checked="" type="checkbox"/>	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
<input checked="" type="checkbox"/>	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
<input checked="" type="checkbox"/>	The statistical test(s) used AND whether they are one- or two-sided <i>Only common tests should be described solely by name; describe more complex techniques in the Methods section.</i>
<input checked="" type="checkbox"/>	A description of all covariates tested
<input checked="" type="checkbox"/>	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
<input checked="" type="checkbox"/>	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)
<input checked="" type="checkbox"/>	For null hypothesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted <i>Give P values as exact values whenever suitable.</i>
<input checked="" type="checkbox"/>	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
<input checked="" type="checkbox"/>	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
<input checked="" type="checkbox"/>	Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

Software and code

Policy information about [availability of computer code](#)

Data collection	data is available to download: Egorova, Yulia (2024). Data used in the study. figshare. Dataset. https://doi.org/10.6084/m9.figshare.26360203
Data analysis	The scripts and data used in this study are all available online at https://github.com/yuliaUU/glob-diver-mesopel-zoop . we use R statistical software for our data analysis

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](#)

The data availability for this study is as follows: all additional information about a study can be found in the Supplementary Material. All data can be downloaded from: <https://doi.org/10.6084/m9.figshare.2636023>. The scripts used in this study are all available online at <https://github.com/yuliaUU/glob-diver-mesopel-zoop>.

Research involving human participants, their data, or biological material

Policy information about studies with [human participants or human data](#). See also policy information about [sex, gender \(identity/presentation\), and sexual orientation](#) and [race, ethnicity and racism](#).

Reporting on sex and gender	<input type="checkbox"/> no human participants were involved in the study
Reporting on race, ethnicity, or other socially relevant groupings	<input type="checkbox"/> no human participants were involved in the study
Population characteristics	<input type="checkbox"/> no human participants were involved in the study
Recruitment	<input type="checkbox"/> no human participants were involved in the study
Ethics oversight	<input type="checkbox"/> no human participants were involved in the study

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

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf

Ecological, evolutionary & environmental sciences study design

All studies must disclose on these points even when the disclosure is negative.

Study description	Despite the ecological importance of mesozooplankton, little is known about their global distribution and diversity in the mesopelagic realm. This study examines the spatial distribution and diversity of 861 mesozooplankton species using species distribution models.
Research sample	861 mesozooplankton species
Sampling strategy	output from species distribution model. SDMs were built using published and unpublished literature that provide abundance and biomass of mesopelagic mesozooplankton
Data collection	The database was compiled from a wide range of both published and unpublished sources to offer valuable insights into the distribution, abundance, and quantitative biomass measurements of mesopelagic species. We used records from the database to obtain a list of mesozooplankton taxa occurring in the mesopelagic zone
Timing and spatial scale	The database consists of 256,868 entries collected from 282 different data sources, spanning the years 1880-2016.
Data exclusions	n/a
Reproducibility	all codes and data are provided
Randomization	n/a
Blinding	n/a

Did the study involve field work? Yes No

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials & experimental systems

n/a	Involved in the study
<input checked="" type="checkbox"/>	Antibodies
<input checked="" type="checkbox"/>	Eukaryotic cell lines
<input checked="" type="checkbox"/>	Palaeontology and archaeology
<input checked="" type="checkbox"/>	Animals and other organisms
<input checked="" type="checkbox"/>	Clinical data
<input checked="" type="checkbox"/>	Dual use research of concern
<input checked="" type="checkbox"/>	Plants

Methods

n/a	Involved in the study
<input checked="" type="checkbox"/>	ChIP-seq
<input checked="" type="checkbox"/>	Flow cytometry
<input checked="" type="checkbox"/>	MRI-based neuroimaging

Plants

Seed stocks

not used in this study

Novel plant genotypes

not used in this study

Authentication

not used in this study