

**SIBIC 2024**  
X Iberian Congress  
of Ichthyology

# Making sense of the R fishbase package in a macroecological trophic fish perspective

Ignasi Arranz, Universidad Rey Juan Carlos



**Fishes for Future**  
19th Vic



**SIBIC 2024**  
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investigación en  
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Programa Atracción



Lake George Reflection  
Georgia O'Keeffe

**UAB**

Universitat Autònoma  
de Barcelona

Technician era

Universitat  
de Girona

**IGB**  
Leibniz-Institute of  
Freshwater Ecology  
and Inland Fisheries



國立臺灣大學  
National Taiwan University

Predoctoral era



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DE CATALUNYA



AARHUS  
UNIVERSITY



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UNIVERSITÉ  
Concordia  
UNIVERSITY

Postdoctoral era

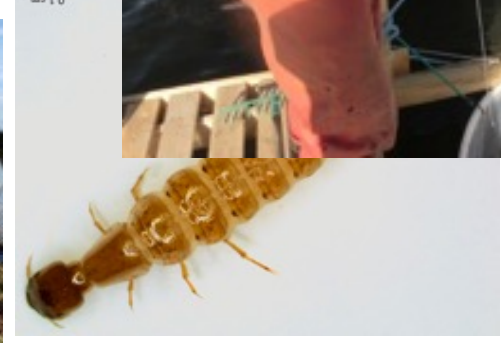
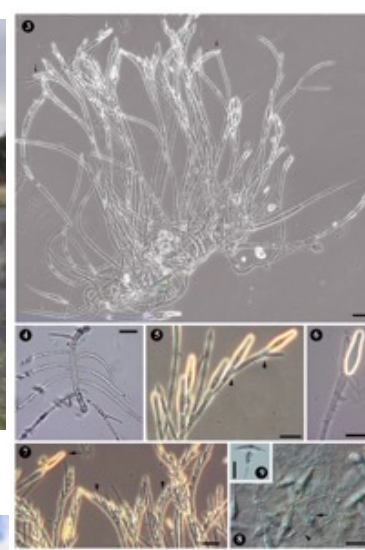


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PAUL SABATIER



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Rey Juan Carlos

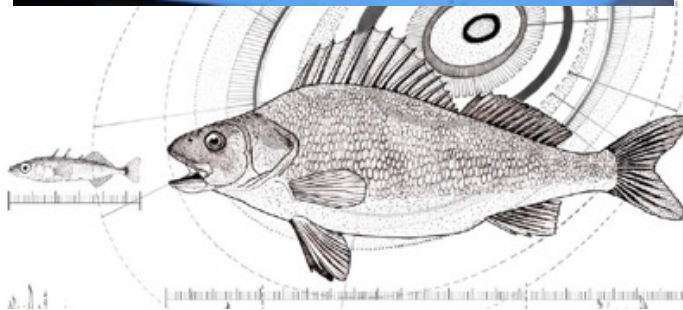




Ignasi Arranz

FotCiencia Ignasi Arranz





# Why this course?

Received 14 April 2021 | Accepted 4 July 2021  
DOI: 10.1111/1365-2450.17047

**RESEARCH HIGHLIGHT**

**Journal of Animal Ecology**

**Reproducibility in ecology and evolution: Minimum standards for data and code**

Gareth B. Jenkins<sup>1</sup> | Andrew P. Beckerman<sup>2</sup> | Céline Bellard<sup>3</sup> | Ana Benítez-López<sup>4</sup> | Aaron M. Ellison<sup>5,6</sup> | Christopher G. Foote<sup>7</sup> | Andrew L. Hufnagel<sup>8</sup> | Marcus A. Lashley<sup>9</sup> | Christopher J. Lortie<sup>9</sup> | Zhaoxue Ma<sup>10</sup> | Allen J. Moore<sup>11</sup> | Shawn R. Narum<sup>12</sup> | Johan Nilsson<sup>13</sup> | Bridget O'Shoye<sup>1</sup> | Diogo B. Provete<sup>14,15</sup> | Orly Razgou<sup>16</sup> | Loren Riseberg<sup>17</sup> | Cynthia Riginos<sup>18</sup> | Luca Santini<sup>19</sup> | Benjamin Sibbett<sup>1</sup> | Pedro R. Peres-Neto<sup>20</sup>

Trends in Ecology & Evolution

CellPress  
REVIEWS

Forum

**Ecological Data Should Not Be So Hard to Find and Reuse**

Timothée Polot<sup>1,2</sup>  
Anne Brunelle<sup>1,2</sup>  
Andrew Gonzalez<sup>3,4</sup>  
Dominique Grais<sup>4</sup> and  
Pedro Peres-Neto<sup>5</sup>

such as DataONE. This model is designed to allow archiving first- and second- order data, and to facilitate the evolution of more robust ways in which new formats can easily incorporate data based on previous ones.

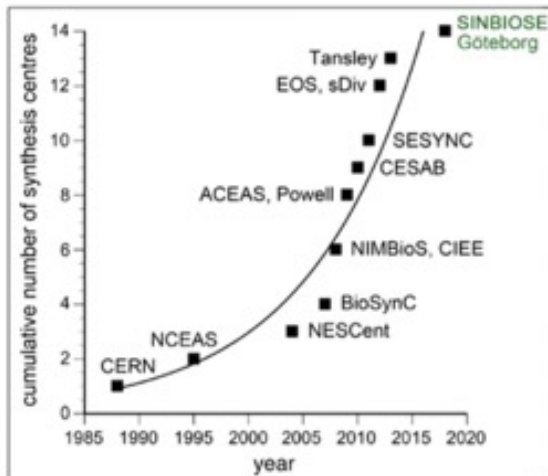
The lack of domain-specific standards leads to a number of challenges faced by synthesis efforts that seek to extract novel knowledge from existing data. Roche and colleagues [8] determined

Review

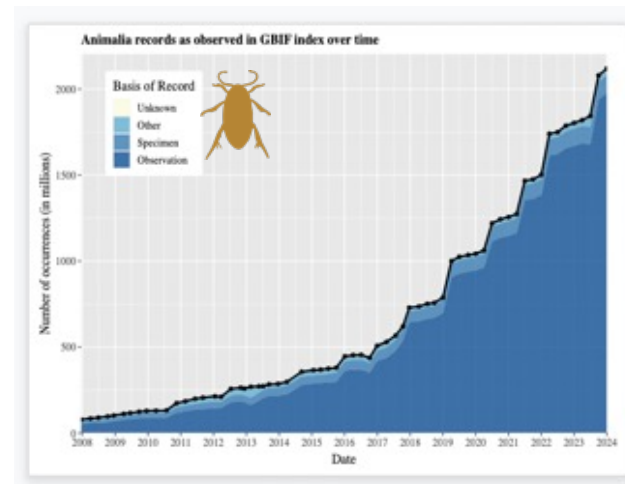
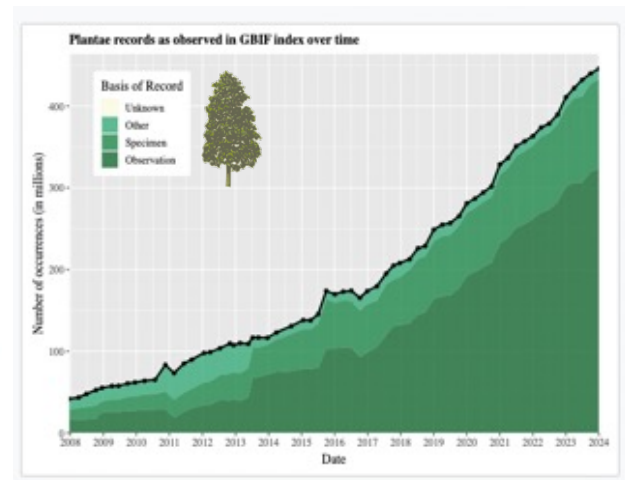
**Special Issue: Long-term ecological research**

**Long-term datasets in biodiversity research and monitoring: assessing change in ecological communities through time**

Anne E. Magurran<sup>1</sup>, Stephen R. Baillie<sup>2</sup>, Stephen T. Buckland<sup>3</sup>, Jan McP. Dick<sup>4</sup>, David A. Elston<sup>5</sup>, E. Marian Scott<sup>6</sup>, Rognvald I. Smith<sup>6</sup>, Paul J. Somerfield<sup>7</sup> and Allan D. Watt<sup>8</sup>



Ref: Nicolas Mouquet



Ref: GBIF

## **Why a macroecological trophic fish perspective?**

Global patterns of fish trophic ecology are scarce

Fish suitable subject group sensitive to human footprint

Answer broad questions in ecology, evolution, and biogeography











*Journal of Fish Biology* (2012) **81**, 2030–2039

doi:10.1111/j.1095-8649.2012.03464.x, available online at [wileyonlinelibrary.com](http://wileyonlinelibrary.com)

## **rfishbase: exploring, manipulating and visualizing FishBase data from R**

C. BOETTIGER<sup>\*†</sup>, D. T. LANG<sup>‡</sup> AND P. C. WAINWRIGHT<sup>\*</sup>

*<sup>\*</sup>Center for Population Biology, University of California, Davis, CA 95616, U.S.A. and*

*<sup>‡</sup>Department of Statistics, University of California, Davis, CA 95616, U.S.A.*

This article introduces a package that provides interactive and programmatic access to the FishBase repository. This package allows interaction with data on over 30 000 fish species in the rich statistical computing environment, R. This direct, scriptable interface to FishBase data enables better discovery and integration essential for large-scale comparative analyses. This article provides several examples to illustrate how the package works, and how it can be integrated into phylogenetics packages such as ape and geiger.

© 2012 The Authors

Journal of Fish Biology © 2012 The Fisheries Society of the British Isles

**Key words:** data access; programmatic; tutorial; XML.

# Package 'rfishbase'

June 3, 2023

**Title** R Interface to 'FishBase'

**Description** A programmatic interface to 'FishBase', re-written based on an accompanying 'RESTful' API. Access tables describing over 30,000 species of fish, their biology, ecology, morphology, and more. This package also supports experimental access to 'SeaLifeBase' data, which contains nearly 200,000 species records for all types of aquatic life not covered by 'FishBase.'

**Version** 4.1.2

**Encoding** UTF-8

**License** CC0

## R topics documented:

rfishbase-package	3
available_releases	3
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# library(tidyverse)

The tidyverse

Components



- Compact format (three digits; only show the first 10 rows...)
- Convenient to see the type of the data

The tidyverse is a collection of R packages that share common philosophies and are designed to work together. This site is a work-in-progress guide to the tidyverse and its packages.

```
# A tibble: 10 × 10
  V1      V2      V3      V4      V5      V6      V7      V8      V9      V10
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1 21.3  0.908  5.59  13.4   2.84  25.8   8.28   2.05  0.537 14.7
2  2.26 37.5    3.74   6.53  11.9  29.5   5.72   6.38  11.4   1.75
3 23.6  11.9    0.400  4.69   1.28   7.64   3.84   4.09  4.34   2.22
4  9.96  2.45  23.2   11.6   3.75  37.5   0.913  8.39  24.1   3.59
5 15.6  0.634  4.98  13.5   1.35   9.98   2.07   4.50  11.8  11.9
6 11.7  3.73   5.51  18.8   9.46   6.75   2.48   7.61  14.9   3.58
7  5.39  9.84   6.38  0.284  3.15   5.16   7.86   3.84  0.282  2.76
8  9.07  7.09   3.52   3.15  0.317  5.10   4.44  10.4   6.17  44.9
9  2.79  7.25  16.7  17.8   3.74  16.0   3.94   8.26  9.20  17.4
10 0.163  7.24  33.4  19.0   2.44   5.15  12.7  10.6   5.68  2.49
```



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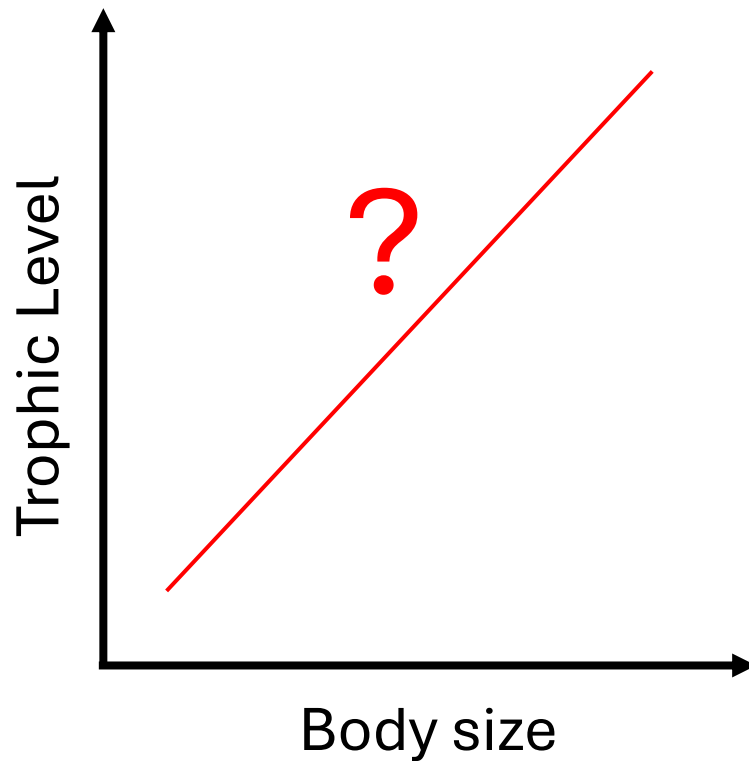
Programa Atracción

# Get started!



## Case study 1

Reveal trophic level vs maximum body size in fish species



## Case study 2

Estimate individual weight from length–weight relationships

$$\text{WEIGHT} = a\text{LENGTH}^b$$



$$\log_{10}(\text{WEIGHT}) = \log_{10}(\text{LENGTH})b + a$$



Data in Brief  
Volume 42, June 2022, 108248



Data Article

### Individual body mass and length dataset for over 12,000 fish from Iberian streams

Ask Copilot: Save time, read 10X faster with AI

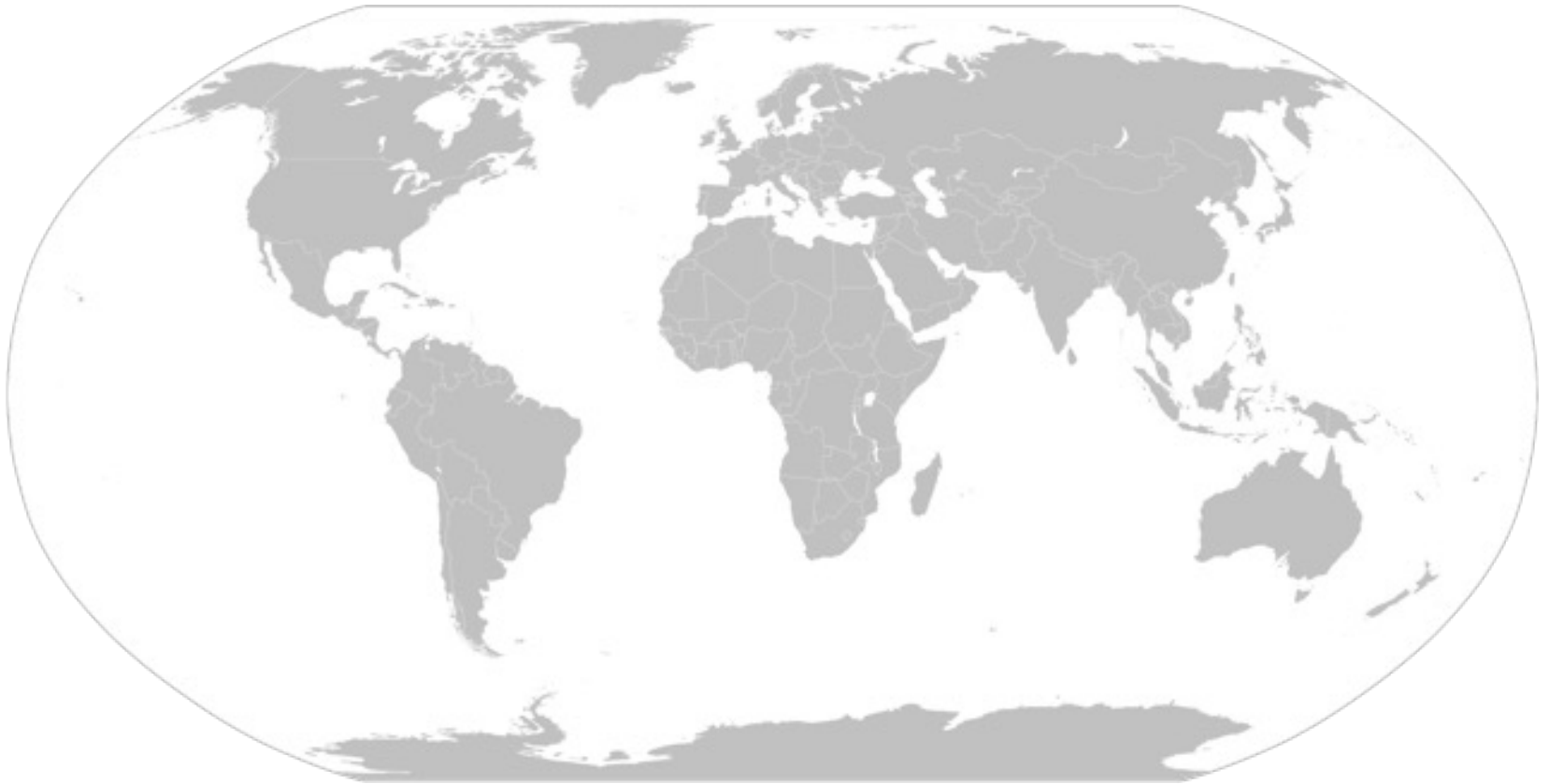
Save Related Papers Purpose Conclusions Evidence/Examples Used  
Points Discussed Biases or Limitations Key Takeaways

Ignasi Arranz<sup>a,b</sup>, Sandra Bruçet<sup>a,c</sup>, Mireia Bartrons<sup>a</sup>, Carmen García-Comas<sup>a,d</sup>,  
Carles Alcaraz<sup>a</sup>, Mònica Bardina<sup>e</sup>, Patricia Navarro-Barovero<sup>a</sup>, Frederic Casals<sup>b,i</sup>,  
Núria Cobiella<sup>j,k</sup>, María Concepción Durán<sup>a</sup>, Emili García-Berthou<sup>l</sup>, Alberto Maceda-Veiga<sup>m</sup>,  
Antoni Munné<sup>f</sup>, María José Rodríguez-Pérez<sup>g</sup>, Carolina Solà<sup>h</sup>, Adolfo de Sostoa<sup>m</sup>, Lluís Benejam<sup>a</sup>



## Case study 3

Create a trophic index at a global scale based on the feeding roles



## Fisheries Magazine

Feature: Fisheries Research | [Full Access](#)

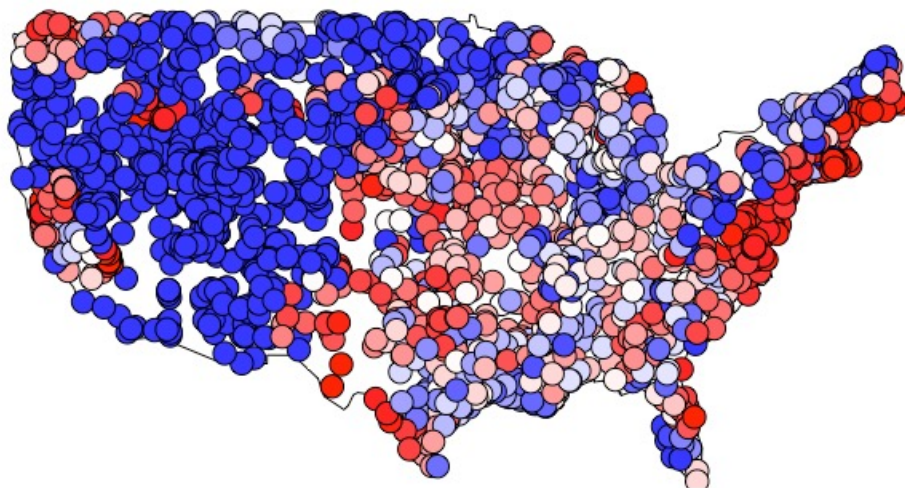
### Fish Traits: A Database of Ecological and Life-history Traits of Freshwater Fishes of the United States

Emmanuel A. Frimpong ✉ Paul L. Angermeier

First published: 26 February 2011 | <https://doi.org/10.1577/1548-8446-34.10.487> | Citations: 195



Planktivorous fish predation  
red = high pressure, blue = low pressure



SCIENTIFIC PAPERS — ANALYSES

### Spatial Data Download

