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RivFishTIME: A global database of fish time-series as a currency for global change ecology research in riverine systems

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Comte, L, Carvajal-Quintero, J, Tedesco, PA, et al. RivFishTIME: A global database of fish time-series to study global change ecology in riverine systems. Global Ecol Biogeogr. 2020; 30: 38– 50. https://doi.org/10.1111/geb.13210

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This document is the accepted manuscript version of the following article: Comte, L., Carvajal-Quintero, J., Tedesco, P. A., Giam, X., Brose, U., Erős, T., … Olden, J. D. (2021). RivFishTIME: a global database of fish time-series to study global change ecology in riverine systems. Global Ecology and Biogeography, 30(1), 38-50. https://doi.org/10.1111/geb.13210

1 RivFishTIME: A global database of fish time-series to study global change

2 ecology in riverine systems

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- 84 **Running title:** Time-series database of riverine fish communities
- 85

86 ABSTRACT

Motivation: We compiled a global database of long-term riverine fish surveys from 46 regional
and national monitoring programs as well as individual academic research efforts upon which
numerous basic and applied questions in ecology and global change research can be explored.
Such spatially- and temporally-extensive datasets have been lacking for freshwater systems
compared to terrestrial ones.

Main types of variables contained: The database includes 11,386 time-series of riverine fish
 community catch data, including 646,270 species-specific abundance records together with
 metadata related to geographic location and sampling methodology of each time-series.

Spatial location and grain: The database contains 11,072 unique sampling locations (stream
reach), spanning 19 countries, 5 biogeographic realms, and 402 hydrographic basins worldwide.

97 Time period and grain: The database encompasses the period 1951–2019. Each time-series is
98 composed of a minimum of two yearly surveys (mean = 8 years) and represents a minimum time
99 span of 10 years (mean = 19 years).

Major taxa and level of measurement: The database includes 944 species of ray-finned fishes
(Class Actinopterygii).

102 Software format: .csv

Main conclusion: Our collective effort provides the most comprehensive long-term community database of riverine fishes to date. This unique database should interest ecologists who seek to understand the impacts of human activities on riverine fish biodiversity, and model and predict how fish communities will respond to future environmental change. Together, we hope it will promote advances in macroecological research in the freshwater realm.

108 KEYWORDS: species abundance; biodiversity; conservation; freshwater streams and rivers;
 109 Actinopterygii; temporal trends; worldwide

110 1| INTRODUCTION

Increasing awareness of the ongoing biodiversity crisis has motivated global initiatives to 111 112 compile large-scale datasets of population and community abundance records that have been consistently sampled through recent times (Pereira & Cooper, 2006). Included among these are 113 the Global Population Dynamics Database (Inchausti & Halley, 2001), the Living Planet Index 114 115 database (Loh et al., 2005), and more recently, the BioTIME database (Dornelas et al., 2018). 116 These databases have proven extremely useful and allowed major advancements in ecological research (e.g. Kendall, Prendergast, & Bjørnstad, 1998; Sibly, Barker, Denham, Hone, & Pagel, 117 118 2005; Butchart et al., 2010; Dornelas et al., 2014); however, they remain highly biased towards 119 terrestrial and marine assemblages (e.g. only 0.50% of the records concern riverine fishes in 120 BioTIME, the most recent of these initiatives). This is unfortunate as effective strategic plans for conserving water resources that support human well-being and ecosystem integrity rely on access 121 122 to comprehensive, pertinent, quantitative information regarding the status and trends of riverine biodiversity over regional to continental scales (Tickner et al., 2020). 123

Long-term studies of riverine species are limited because they require highly specialized 124 and time-consuming sampling methods. Furthermore, rivers in remote areas are often difficult to 125 126 access (Olden et al., 2010; Radinger et al., 2019). Nevertheless, over the past few decades, large-127 scale policies have been enacted in response to the rapid degradation of freshwater resources, 128 such as the Water Framework Directive in the EU (Hering, Verdonschot, Moog, & Sandin, 2004) and the Clean Water Act in the USA (Paulsen et al., 2008), which require countries to monitor 129 130 and evaluate the biological integrity of surface waters through time to adopt quality standards that restore and maintain ecological integrity (Kuehne, Olden, Strecker, Lawler, & Theobald, 131 132 2017). Beyond these official national and regional monitoring programs, the temporal dynamics of riverine systems and their fish communities have also been assessed through various 133 134 independent, though often local in extent, academic research programs (e.g. Gido, 2017; Matthews & Matthews, 2017). All of these institutional and academic monitoring efforts have 135 produced considerable freshwater fish temporal data that remain largely inaccessible to the 136 broader scientific community due to the inherent difficulty in gathering and harmonizing field 137 data from disparate institutions and sampling protocols (Buss et al., 2015). 138

To fill this important gap, we here present RivFishTIME, a compiled and curated 139 database of long-term (> 10 years) surveys of riverine fish communities at a fine spatial (stream 140 141 reach) and taxonomic (species) resolution, using data mining approaches to harmonize existing but currently fragmented biomonitoring data sets. Riverine fish are extremely diverse in spite of 142 the small surface they inhabit on Earth: they represent about 40% of all known fish species while 143 144 occupying <1% of available aquatic habitat ("the freshwater fish paradox" sensu Lévêque, Oberdorff, Paugy, Stiassny, & Tedesco, 2008 and Tedesco, Paradis, Lévêque, & Hugueny, 145 146 2017). However, they are also among the most threatened taxonomic groups on Earth because of the convergence between the high concentration of biodiversity and the many pressures resulting 147 from human uses of freshwater resources and habitat change (Reid et al., 2019; Tickner et al., 148 2020). The RivFishTIME database provides a unique opportunity to understand the rate, 149 150 magnitude, and geography of biodiversity trends, and to identify opportunities to mitigate human impacts on riverine systems (Pereira & Cooper, 2006; Anderson, 2018). Due to the paucity of 151 152 spatially- and temporally-extensive datasets in freshwater compared to terrestrial systems (Heino, 2011), RivFishTIME should also help ecologists close the gap between these two 153 154 systems and to address a wider range of taxa in unraveling large-scale spatio-temporal 155 biodiversity patterns.

156

157 **2 METHODS**

158 2.1 Data acquisition

159 We gathered time-series of fish community abundance data for riverine (lotic) ecosystems, 160 broadly defined as freshwater bodies that are continually or intermittently flowing. We tried to the extent possible to exclude wetlands and brackish habitats (salinity > 0.5 %). Note, however, 161 162 that due to the complex nature of the datasets, we do not guarantee that sites are located on free-163 flowing river segments (i.e. natural condition without impoundment, diversion, or other modification of the waterway). We used the following criteria for data inclusion: (1) the location 164 of the sampling sites is known and consistent through time, (2) the sampling protocol is known 165 166 and consistent through time, (3) the sampling survey sought to quantify all species in the fish

167 community according to well-established protocols, (4) species-specific abundances are available
168 for each survey, (5) surveys at a given site were conducted over a period of at least 10 years, and
169 (6) at least two yearly surveys with non-null abundance are available. We considered abundance
170 measures derived from direct fish counts, catch-effort indexes such as relative abundances (%)
171 and catch per unit effort (CPUE), abundance classes, as well as statistically estimated
172 abundances (e.g. Leslie method; Ricker, 1975).

173 To identify potential datasets, we used Google Search, Google Scholar and Dataset Search with different combinations of the keywords "time series", "fish", "abundance", 174 "stream", "river", "freshwater", "community", "temporal", and "monitoring" or "monitoring 175 176 program". We screened the scientific as well as the grey literature to identify studies involving 177 temporal datasets of fish communities and conducted similar searches in data repositories such as Dryad (https://datadryad.org/stash) and FigShare (https://figshare.com/). We also conducted 178 179 targeted searches for national and regional monitoring programs by adding country names to the previous keywords. For the European Union, we further used the EuMon database as a reference 180 to identify fish monitoring databases (available at http://eumon.ckff.si/about_daeumon.php). 181

We contacted all the authors and monitoring program coordinators unless the reusability of data was clearly stated on the online repositories where the data were released (e.g., Open Government License, CC0 1.0 Universal). We excluded the datasets for which we did not receive the permission.

186

187 2.2 Quality control

188 *Taxonomy*. We validated species scientific names using the online database Fishbase (Froese &

189 Pauly, 2019). We used the R package *rfishbase* (as of December 2019; Boettiger, Lang, &

190 Wainwright, 2012) and confirmed names with no match manually using the Catalog of Fishes

191 (Fricke, Eschmeyer, & van der Laan, 2018). We then selected only records involving ray-finned

192 fishes (Class Actinopterygii), excluding rays and lampreys, and unidentified species.

193 Coordinates. We harmonized the coordinate system by projecting (if necessary) the 194 coordinates of the individual datasets using the World Geodetic System (WGS84) as reference 195 geographic coordinate system. We visually inspected the spatial distribution of the sites with 196 respect to their respective country, region, or state borders as given in the original data sources 197 and discarded sites with dubious coordinates (e.g. sites located in the ocean). We also removed 198 sites whose coordinates were located outside of any hydrographic basin using the global major 199 river basin GIS layer in HydroSHEDS (Lehner, Verdin, & Jarvis, 2008).

Consistent sampling methods. We excluded surveys lacking information on sampling 200 methods and selected only time-series collected using a consistent sampling protocol through 201 time. The latter evaluation was dataset-specific as dictated by the complexity of the monitoring 202 203 scheme and the available metadata. For instance, surveys were deemed consistent if they did not 204 experience any major deviation in sampling protocol, and disregarded minor variations (e.g. 205 number of anodes or traps, area sampled) due to survey-specific constraints (e.g. water depth, habitat complexity). By contrast, several monitoring programs implemented alternate sampling 206 207 protocols to compare the efficiency of different gears (e.g. seining versus electrofishing) or 208 sampling methods (e.g. continuous versus point electrofishing); these time-series conducted at 209 the same sites but using different sampling protocols were kept separate in the database.

210 *Duplicates*. We removed duplicates within individual datasets based on the coordinates of the 211 sites, date of the survey, and species collected (e.g. due to different name attribution for the same 212 site). We also identified potential duplicates among datasets (e.g. overlap between state-level and 213 national databases) based on the coordinates of the sites rounded to three digits to account for 214 different post-processing of the individual datasets.

215

216 2.3 Database formatting

Each entry (species abundance record) was assigned a unique (1) site, (2) survey, and (3) time-

series identifier. The site ID corresponds to a given pair of coordinates, the survey ID to a

sampling campaign, and the time-series ID to a combination of site \times sampling protocol. We

extracted the names of the sampled water bodies (e.g. creek, stream, river) from the available 220 221 metadata associated with each individual dataset, that we cross-referenced against several 222 continental and national geospatial river networks in GIS (e.g. Australian Hydrological 223 Geospatial Fabric, Ordnance Survey Open Rivers). Additionally, each site ID was assigned to a biogeographic realm following Olson et al. (2001), hydrographic basin following HydroSheds 224 225 (Lehner et al., 2008), and administrative units (country, region and province) based on its coordinates. For each sampling ID, we aggregated abundance records if they were given 226 227 separately for individuals, size classes or sub-species for each validated species name or if different sampling passes, hauls, or sub-sampling areas were considered. We also converted 228 time-series species abundances to densities or CPUE whenever possible. The different surveys 229 were kept independent when conducted on different occasions within the same calendar year. We 230 231 provided the year together with the quarter of the survey (1: January-March; 2: April-June; 3: July-September, 4: October-December). We also provided the associated unit (abundance class, 232 count, CPUE, individuals/100m², Leslie index, relative abundance) for each species abundance 233 record. Finally, we extracted basic information regarding the sampling protocol, including details 234 235 on electrofishing (backpack, shore-based or boat mounted electrofishers), netting (dip nets, gill nets, beach or pelagic seines), trapping (minnow traps, fyke nets or hoop nets), and trawling 236 237 techniques. Many survey protocols involve a combination of sampling approaches, rendering 238 challenging the inclusion of detailed information about the sampling effort in a standardized 239 way. We therefore encourage the data user to refer to each data source for more information on the sampling methods. 240

The database is organized in three tables (.csv format): the time-series table, the survey 241 table, and the information source table. The tables can be linked using the unique dataset source 242 243 ID and time-series ID. The time-series table contains: (1) source ID, (2) site ID, (3) time-series ID, (4) sampling method, (5) latitude (WGS 84), (6) longitude (WGS 84), (7) biogeographic 244 realm, (8) hydrographic basin, (9) country (ISO code), (10) region, (11) province, and (12) water 245 body. The survey table contains: (1) time-series ID, (2) survey ID, (3) sampling year, (4) 246 247 sampling quarter, (5) species scientific name, (6) abundance, and (7) abundance unit. The information source table contains the full citation(s), online link to the raw data when publicly 248 249 available, as well as the name(s) and contact of the data responsible(s) for each individual

dataset. Data curation was performed in the *R* (3.6.0) programming environment (R Core Team,
2019).

A list of the data sources is found in Appendix 1; for further information consult the metadata. A static version of RivFishTIME is available through the iDiv Biodiversity Portal (Comte et al., 2020), but we aim to continue interacting with data contributors to update and add new time-series datasets as they become available (see *Data Availability Statement*).

256

257 **3**| **RESULTS**

Our database includes 11,386 time-series of riverine fish compiled from 46 individual source 258 datasets, representing a total of 106,785 surveys and 646,270 individual species abundance 259 records at 11,072 unique sites. Survey-specific species richness across all time-series ranges 260 from 1 to 50 species, and covers 944 ray-finned fish species. The surveyed sites display a wide 261 262 distribution along longitudinal and latitudinal gradients, spanning 19 countries, 402 hydrographic basins, and 5 biogeographic realms (Fig. 1a). Despite broad geographical coverage, we note a 263 clear spatial bias towards the Palearctic (European Union) and, to a lesser extent, Nearctic (North 264 America) and Australasia realms. The abundance time-series are largely represented by 265 individual counts, followed by densities (individuals/100m²) and CPUE (Fig. 1b). Abundance 266 classes, Leslie index and relative abundance represent < 1% of the time-series. Electrofishing is 267 by far the main sampling technique used to record the time-series, although variations are 268 269 noticeable among biogeographic realms (Fig. 1c). For instance, dipnetting sampling techniques 270 are only represented in the Neotropics, whereas gillnetting is the most common gear in the 271 Afrotropics.

The time-series cover a time period from 1951 to 2019 and are mainly concentrated over the last two decades (average first year = 1996; Fig. 2a). Surveys have been conducted primarily in the 3rd (July-September) and 4th (October-December) quarters of the year, especially in the Palearctic and Nearctic realms (corresponding to periods of low flows), but all quarters are represented in the different biogeographic realms (Fig. 2b). The mean time span of the timeseries is of 19 years and ranges from 10 to 68 years, with the longest time-series located in the Palearctic (Fig. 2c). The sites were sampled from (non-necessarily consecutive) 2 to 52 years, with an average number of yearly surveys of 8 years (Fig 2d). Again, the highest number of yearly surveys was found in the Palearctic. The completeness of the time-series (i.e. ratio of number of yearly surveys to the overall time span) ranges from 4 to 100%, with a mean value of 45% (Figure 2e). Importantly, the degree of completeness is largely uncorrelated to the time span of the time-series (r = 0.05).



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Figure 1. (a) Map showing the distribution of the time-series where each time-series is represented by a dot with colors indicating the biogeographic realm and size representing fish species richness (averaged across surveys). Inset histograms display the distribution of the timeseries according to latitude and longitude. Barplots show the distribution of the time-series with respect to the (b) type of abundance, and (c) primary sampling method. Note the $log_{10}(x+1)$ *y*axes in (b)-(c).



Figure 2. (a) Temporal distribution of the yearly surveys relative to the period covered by the database (1951-2019). Each time-series appears in rows where the background colors correspond to the biogeographical realms and white indicates sampled years. (b) Temporal distribution of the surveys with respect to the quarter of the year. Temporal characteristics of the time-series with respect to the (c) overall time span, (d) number of yearly surveys, and (e) completeness defined as the ratio between the number of yearly surveys and the overall time span (expressed in %). Note the log₁₀(x+1) y-axes in (b)-(e).

299

300 4 CONCLUSIONS

301 Our collective effort provides the most comprehensive long-term community database of riverine 302 fishes to date, spanning large biogeographic, climatic and hydrographic gradients. Almost all biogeographic realms are represented but it is important to note that our database is not exempt 303 from spatial bias. For instance, less than 1% of the time-series belong to the Afrotropic or 304 305 Neotropic realms, whereas 84% belong to the Palearctic realm. These spatial gaps often present 306 in biodiversity-rich regions (tropical areas, southeast Asia) are likely to mirror the current networks of freshwater monitoring programs (Buss et al., 2015; Radinger et al., 2019) as well as 307 308 biodiversity research efforts (Martin, Blossey, & Ellis, 2012), and hence will be prioritized in 309 future updates of RivFishTIME. We also warn data users that species abundance may not be directly comparable across sites without a full understanding of the specifics of sampling 310 approach and effort, with respect to their selectivity and efficiency (Goffaux, Grenouillet, & 311 312 Kestemont, 2005; Portt, Coker, Ming, & Randall, 2006; Oliveira, Gomes, Latini, & Agostinho, 2014; Benejam et al., 2012), and refer to the original data sources for more information about the 313 314 sampling protocols.

Despite these unavoidable limitations associated with secondary datasets collected for 315 316 multiple purposes, we are confident that RivFishTIME will stimulate new research in the fields 317 of global change ecology and macroecology. First and foremost, it will provide the needed 318 baseline information for conservation and restoration efforts to bend the curve of freshwater biodiversity loss (Tickner et al., 2020). For instance, the fish abundance time-series could be 319 320 used to assess population or community trends in different rivers of the world, broadening the taxonomic and spatial representation of existing indicators of the status of global biodiversity 321 322 (e.g. Living Planet Index). Coupled with high-resolution environmental time-series, this unique database could also help to decipher the underlying drivers of biodiversity changes in riverine 323 324 systems, including (but not limited to) habitat fragmentation and destruction, invasive species, pollution, hydrologic alteration and climate change (e.g. Chen & Olden, 2020, Erős et al., 2020). 325 In turn, this knowledge could be integrated into ecological models used to forecast how fish 326 communities will respond to future environmental change, paving the way to mitigate those 327 328 impacts. RivFishTIME could also offer new macroecological insights into the implications of river network complexity on community structure and assembly processes across extensive 329

- environmental gradients (e.g. community composition, population persistence, spatial synchrony
- in community dynamics) questions that have long fascinated ecologists but have been so far
- primarily explored through theoretical approaches.
- 333

334 ACKNOWLEDGMENTS

The data compilation is the result of a joint effort of the international working group "sYNGEO -335 The geography of synchrony in dendritic networks" kindly supported by sDiv, the Synthesis 336 Centre of the German Centre for Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig, 337 funded by the German Research Foundation (FZT 118). We thank the iDiv Biodiversity 338 Informatics unit and Jitendra Gaikwad for the support received to prepare, upload and scrutinize 339 dataset integrity. We also would like to thank Angelo Agostinho, Agencia Catalana del Agua, 340 Agencia Vasca del Agua, Jammes Gammon, Iowa Department of Natural Resources, William 341 Matthews, Edie Marsh-Matthews, Minnesota Pollution Control Agency, Montana Fish, Wildlife 342 & Parks, Murray-Darling Basin Authority, North Carolina Department of Environmental 343 Quality, Okavango Research Institute, Oklahoma Water Resources Board, Office Français de la 344 Biodiversité (OFB), Parks Canada, Vincent Resh, John Rinne, Swedish University of 345 Agricultural Sciences, Toronto and Region Conservation Authority, U.K. Environmental 346 347 Agency, Upper Midwest Environmental Sciences Center, U.S. Environmental Protection Agency, U.S. Geological Survey and Gerlinde Van Thuyne for making their monitoring data 348 available. Data collection for the SourceID #2 was supported by Fapesp - São Paulo Research 349 Foundation through the research grants to J.O.Z. (Processes Numbers: 2012/20280-5 and 350 351 2015/04366-5) and LC (Processes Numbers: 2001/13340-7, 2012/05983-0) and CNPq (Conselho 352 Nacional de Desenvolvimento Científico e Tecnológico) through the research grant provided to LC (Process Number: 301877/2017-3). Data collection for the SourceID #3 was supported by 353 Convenio CT-2017- 001714 between University of Antioquia and Empresas Publicas de 354 Medellin. Part of the data collection for the SourceID #4 was supported by the GINOP-2.3.2.-15-355 356 2016-00004 project. Data for the SourceID #6 were collected by the Queensland Department of 357 Environment and Science in collaboration with Healthy Land and Water Limited. Data collection for the SourceID #10 was supported by the New Mexico Department of Game and Fish. We 358 acknowledge CNPq for research grants provided to J.R.S.V. (Process Numbers: 302367/2018-7 359 360 and 303776/2015-3) [SourceID #13], and the Japan Society for the Promotion of Science KAKENHI Grant 18K06404 to A.T. [SourceID #14]. We also acknowledge the assistance of Dr. 361 Stefano Porcellotti, Dr. Simona Piccini (UTR Grosseto) and Dr. Lorena Di Iulio Chiacchia (UTR 362 Prato), who kindly helped with original data retrieval for the SourceID #16. Data collection for 363 the SourceID #17 (Ivory Coast) was supported by the World Health Organisation (WHO), 364 Onchocerciasis Control Programme (OCP). Data collection for the SourceID #18 was supported 365 366 by the United States Fish and Wildlife Service and Oklahoma Department of Wildlife

Conservation through Endangered Species Act funding (Project E-12). Data collection for the 367 SourceID #19 (Kings Creek watershed) was supported by the NSF Long-Term Ecological 368 Research Program at Konza Prairie Biological Station. Data collection for the SourceID #20 369 (Upper Little Tennessee River Watershed) was supported by Brad Stanback, Janirve Foundation, 370 371 Macon County Community Foundation, Merck Family Fund, National Fish and Wildlife Foundation, National Forest Foundation, NC Division of Water Quality, NC Wildlife Resources 372 Commission, Norcross Foundation, River Network, Southern Appalachian Man and the 373 Biosphere Foundation, Tennessee Valley Authority, World Wildlife Fund, and Z. Smith 374 375 Reynolds Foundation. Data collection for the SourceID #29 was supported by the Companhia Energética de Minas Gerais (P&D ANEEL/CEMIG GT-487 and GT-599), Fundação de Amparo 376 à Pesquisa do Estado de Minas Gerais (FAPEMIG), Coordenação de Aperfeiçoamento de 377 Pessoal de Nível Superior (CAPES) and CNPq. Data collection for the SourceID #35 (Upper 378 Paraná River) was supported by the Long-Term Research Program / CNPq-Sitio 6, carried out by 379 380 Nupélia / UEM (Núcleo de Pesquisas em Limnologia, Ictiologia e Aquicultura/Universidade Estadual de Maringá). R.B.D.C. received a scholarship from CAPES and F.G.B. received a 381 student scholarship from CNPq and financial support from WWF (Programa Natureza e 382 Sociedade) for sampling done in 1999 [SourceID #37]. Data collection for the SourceID #40 was 383 384 supported by the New Mexico Department of Game and Fish. Data collection for the SourceID #43 was supported by various town councils and the Catalan Water Agency through the 385 "Observatori de la Tordera" project (led by Dr Martí Boada). 386

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388 DATA AVAILABILITY

- 389 RivFishTIME is publicly available through the iDiv Biodiversity Portal:
- 390 <u>https://doi.org/10.25829/idiv.1873-10-4000</u>. We kindly ask the users to cite this paper as well as
- the source of each primary dataset in any published material produced using these data. We
- encourage any potential data contributor to contact LC with possible datasets to expand the
- database. Updates of RivFishTIME will be curated through the iDiv Biodiversity Portal and also
- released through the more specialized Freshwater Biodiversity Data Portal
- 395 (https://data.freshwaterbiodiversity.eu/).
- 396

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501 Appendix 1 | Data sources

SourceID	Citations	URLaccess
1	Agència Catalana de l'Aigua (2003); Agència Catalana de l'Aigua (2010); Agència Catalana de l'Aigua (2018)	http://aca.gencat.cat/
2	Zeni, Hoeinghaus, & Casatti (2017); Casatti et al. (2009)	-
3	Universidad de Antioquia-Empresas Publicas de Medellin (2019)	-
4	Erős et al. (2014)	-
5	Gammon (2013)	-
6	Ecosystem Health Monitoring Program Queensland (2019)	https://hlw.org.au/report-card/
7	Finnish electrofishing register Hertta (2019)	-
8	Sigouin (2017)	https://open.canada.ca/data/en/dataset /fe2441a6-8ae4-4884-b181- cd7ec53bd842
9	Whitney, Gido, Martin, & Hase (2016)	-
10	Gido, Propst, Olden, & Bestgen (2013); Gido et al. (2019)	-
11	Kesner & Marsh (2010)	https://www.rosemonteis.us/documen ts/kesner-marsh-2010
12	Griffith (2017); Griffith, Zheng, & Cormier (2018)	https://doi.org/10.23719/1376690
13	Occhi, V. T. & Vitule, J. R. S. (Unpublished data)	-
14	Terui et al. (2018)	-
15	Iowa Department of Natural Resources (2013)	https://data.iowa.gov/Environment/Bi oNet/e7yf-f5fs
16	Milardi et al. (2020)	-
17	Levêque, Hougard, Resh, Statzner, & Yaméogo (2003)	-
18	Pyron, Vaughn, Winston, & Pigg (1998)	-

19	Gido (2017)	https://doi.org/10.6073/pasta/150e218 b069074a8ecede85a7406d43f
20	McLarney, Meador, & Chamblee (2013)	https://www.mainspringconserves.org /what-we-do/aquatic-monitoring/
21	Long Term Resource Monitoring Program (2016)	https://www.umesc.usgs.gov/data_lib rary/fisheries/fish1_query.shtml
22	Matthews & Marsh-Matthews (2017)	https://doi.org/10.5061/dryad.2435k
23	Murray-Darling Basin Authority (2018)	https://data.gov.au/data/dataset/murra y-darling-basin-fish-and- macroinvertebrate-survey
24	Minnesota Pollution Control Agency (2018)	https://www.pca.state.mn.us/water/bi ological-monitoring-water-minnesota
25	Montana, Fish, Wildlife & Parks (2019)	http://gis- mtfwp.opendata.arcgis.com/items/81 92e75218c6460ba97ba3dd0a2fb3a5
26	U.S. Geological Survey (2019)	https://aquatic.biodata.usgs.gov/clear Criteria.action
27	U.K. Environmental Agency (2019)	https://data.gov.uk/dataset/d129b21c- 9e59-4913-91d2-82faef1862dd/nfpd- freshwater-fish-survey-relational- datasets
28	North Carolina Department of Environmental Quality (2018)	https://deq.nc.gov/about/divisions/wa ter-resources/water-resources- data/water-sciences-home- page/ecosystems-branch/fish-stream- assessment-program
29	Fagundes et al. (2015)	-
30	Winston, Taylor, & Pigg (1991); Taylor (2010)	https://onlinelibrary.wiley.com/doi/fu ll/10.1111/fwb.13211
31	Mosie & Makati (2018)	https://www.gbif.org/dataset/77929c0 a-7506-4b2d-a49d-10fc3312d50d
32	Office Français de la Biodiversité (2019)	http://www.naiades.eaufrance.fr/acce s-donnees#/hydrobiologie
33	Oklahoma Water Resources Board (2019)	http://home- owrb.opendata.arcgis.com/search?tag s=fish
34	Agencia Vasca del Agua (2019)	http://www.uragentzia.euskadi.eus/in formazioa/ubegi/u81-0003341/eu/

35	Ortega, Dias, Petry, Oliveira, & Agostinho (2015)	-
36	Davenport, S.R. (Unpublished data)	-
37	Dala-Corte, Becker, & Melo (2017)	-
38	Bêche, Connors, Resh, & Merenlender (2009)	https://nature.berkeley.edu/reshlab/
39	Toronto and Region Conservation Authority (2018)	https://data.trca.ca/dataset/2018- watershed-fish-community
40	U.S. Fish and Wildlife Service (2017)	-
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42	Sers (2013)	https://www.slu.se/en/departments/aq uatic-resources1/databases1/database- for-testfishing-in-streams/
43	Benejam, Angermeier, Munné, & García-Berthou (2010); Merciai, Molons-Sierra, Sabater, & García- Berthou (2017)	_
44	Miyazono & Taylor (2015)	https://bioone.org/journals/The- Southwestern-Naturalist/volume- 60/issue-1/MP-02.1/Long-term- changes-in-seasonal-fish-assemblage- dynamics-in-an/10.1894/MP- 02.1.short
45	Rinne & Miller (2006)	-
46	Van Thuyne et al. (2013); Brosens et al. (2015)	https://ipt.inbo.be/resource?r=vis- inland-occurrences

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