A growing interest is devoted to global-scale approaches in ecology and evolution that examine patterns and determinants of species diversity and the threats resulting from global change. These analyses obviously require global datasets of species distribution. Freshwater systems house a disproportionately high fraction of the global fish diversity considering the small proportion of the earth’s surface that they occupy, and are one of the most threatened habitats on Earth. Here we provide complete species lists for 3119 drainage basins covering more than 80% of the Earth surface using 14953 fish species inhabiting permanently or occasionally freshwater systems. The database results from an extensive survey of native and non-native freshwater fish species distribution based on 1436 published papers, books, grey literature and web-based sources. Alone or in combination with further datasets on species biological and ecological characteristics and their evolutionary history, this database represents a highly valuable source of information for further studies on freshwater macroecology, macroevolution, biogeography and conservation.

Design Type(s)

- data integration objective
- database creation objective
- species comparison design

Measurement Type(s)

- biodiversity assessment objective

Technology Type(s)

- digital curation

Factor Type(s)

- geographic location

Sample Characteristic(s)

- Earth
- drainage basin

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Background & Summary
With c. 126,000 already described animal species, freshwater systems host around 10% of all animals described to date\(^1\)–\(^3\) while occupying only 0.8% of the Earth’s surface and 0.02% of available aquatic habitable volume\(^4\). Among aquatic organisms, fishes are a good example of this paradox (the ‘freshwater fish paradox’ sensu Tedesco et al.\(^5\)), with c. 40% of all described species inhabiting freshwaters, while the remaining 60% inhabiting marine habitats that comprise >99% of available aquatic habitat\(^5\). Besides housing a disproportionately high fraction of the global animal diversity considering the small proportion of the earth’s surface that they occupy, freshwater ecosystems are also one of the most threatened habitats on Earth\(^7,8\). Extinction risk for freshwater fishes, for instance, is thought to be higher than that of terrestrial organisms\(^9\) and recent extinction rate estimates are 112 to 855 times higher than natural extinction rates\(^10\)–\(^12\).

Describing global scale freshwater fish diversity patterns, understanding the environmental drivers and evolutionary processes shaping such diversity and revealing the major human-related threats were the major goals that motivated the compilation of the present database. Indeed, global scale datasets allowing for biogeographical, macroecological, macroevolutionary and conservation studies were available for only a few well-documented animal taxa such as birds, mammals and amphibians\(^13\)–\(^15\). The present database increases this list of taxa by providing occurrence data by drainage basin worldwide for the most diverse group of vertebrates (i.e. fishes), with more than 33500 species described to date (FishBase; http://www.fishbase.org), from which c. 40% inhabit permanently freshwater systems.

We conducted an extensive survey of freshwater fish species distribution based on 1436 published papers, books, grey literature, databases and web-based sources, resulting in species lists for 3119 drainage basins covering more than 80% of the Earth surface (Fig. 1). Two important survey efforts were conducted, respectively completed in 2008 (ref. 16) and 2013 (ref. 17). To date, these databases have been used in several studies that have increased our understanding of freshwater fish species distributions. These studies allowed to accurately map global patterns of native\(^18\), endemic\(^19\) and non-native\(^20\) freshwater fish species richness and to reveal their environmental and human-related determinants. The databases were also used to evaluate non-native species influence on native communities structure\(^21\), to forecast climate change effects on species extinction processes\(^11\) and to analyse effects of current and future scenarios of species introductions on fish faunas homogenization processes\(^12\)–\(^25\). Recent studies also focused on analysing the influence of past river connections on the present distribution of native fish species\(^17\), on analysing geographical and trait-based differences in diversification rates and origin of actinopterygian fish families\(^5\), and on evaluating human-related extinction drivers\(^12\).

Although the database has already provided a lot of insightful knowledge, it still represents a valuable source of information for further studies on freshwater macroecology, macroevolution, biogeography and conservation. For instance, the present dataset could serve to identify diversity hotspots and to generate a global map of ichthyogeographic regions by combining data on the distributions and phylogenetic relationships of species, allowing in fine the identification of geographic areas harbouring distinct evolutionary histories. Furthermore, in association with data on the time and place of origin of species or on species functional traits, the global occurrence dataset could provide new insights on the macroevolution of freshwater fishes or approach the functional characteristics of communities.

Figure 1. Global map indicating the drainage basins included in the database with different colours by biogeographic realm\(^27\). The 3119 drainage basins cover more than 80% of the Earth surface (excluding deserts), ranging from 70% for the Indo-Malay region to over 90% for the Afro-tropical region.
forthcoming approaches would surely help designing large scale conservation priorities for freshwater fishes.

The database is organised in three sub-datasets and one shapefile. The first dataset contains the species occurrence records by drainage basin along with their native or non-native status and the corresponding FishBase species code and valid name. The second dataset, which is simply the export of the shapefile attributes table, contains geographic information on the drainage basins (e.g. geographic coordinates, surface area). The third dataset contains the list of references that were used to build the species lists for each of the drainage basins. This reference list is obviously not definitive and updates of the database will be performed regularly to include new occurrence records, the distribution of newly described species, species lists of new drainage basins and nomenclature changes in the always moving taxonomy.

Methods

Information sources

This global database of freshwater fish species distribution results from a joint collaboration between three French research institutes, i.e. the University Paul Sabatier in Toulouse (UPS), the National Museum of Natural History (MNHN) and the Research Institute for Development (IRD). The financial support necessary to build this database mainly came from two projects: the ‘Freshwater Fish Diversity’ (National Agency for Research: ANR-06-BDIV-010) and ‘BioFresh’ (7th Framework European program, Contract N°226874) projects. Starting in 2003, we conducted an extensive survey of literature published from 1960 to 2014 on native and non-native freshwater fish species at the drainage basin grain. This survey was complemented with web-based sources from national and international biodiversity inventory initiatives compiling either or both collection and field sampling data.

Our efforts were mainly devoted to find information sources providing complete fish species lists of a given drainage basin, except for some large basins (e.g. the Amazon basin) where we cumulated sub-drainage basin species lists and point sampling locations to obtain the most complete possible coverage of the entire drainage basin. We also used local or regional check lists such as local inventories of stream reaches or inventories based solely on a given family or genus to complement our species lists and for cross-checking available information at the drainage basin scale. The resulting database was gathered from 1436 sources including published papers, books, grey literature and web-based sources that included museum collections, national or regional initiatives compiling monitoring data (mainly for developed countries), continental scale atlases of species distribution and international biodiversity initiatives. When published information was found in languages not handled by any of the team members (e.g. national inventory reports or books), a translator kindly helped us to ensure the collection of correct information on river basins, species lists and location of the species.

Species, taxonomy and status

Sub-species were not considered due to limited data availability and all occurrences not identified to species level were discarded (i.e. occurrences giving only genus names commonly abbreviated to sp., species affinis commonly abbreviated to: sp. aff., aff., or affin. or species confer abbreviated to cf.). Species migrating between both marine and freshwater environments where systematically included in the database. Concerning marine and estuarine species occasionally occurring in freshwaters, these species may be reported in the database but their distribution information should not be considered as exhaustive in any case, as these systems are not the focus of the database.

All species scientific names are reported in the database as given in each information source. These species names were then carefully checked for typing errors and misspellings. Because taxonomy is a ‘moving target’, species names were standardized based on valid species names and their synonyms reported in FishBase using the ‘fishbase’ package26 from the R environment (http://www.R-project.org). For those species names that did not match with any synonym or valid name from FishBase, a manual search was applied in the Catalogue of Fishes (http://researcharchive.calacademy.org/research/ichthyology/catalog/fishcatmain.asp). This last step allowed finding valid species names and species recently described that are still not included in FishBase. For recently described species not yet validated by FishBase or species only considered valid by the Catalogue of Fishes, a temporary code was created starting by ‘x’ (0.04% of all valid species). The remaining species names were considered as invalid and excluded from the database (only 0.6% of all species names). The final standardized species list has 14953 valid names avoiding biases due to synonyms and uncertain identifications (see ‘Technical Validation’).

According to FishBase, from this list of 14953 valid names our database contains the distribution of 13721 species inhabiting fresh or brackish waters, the remaining being marine species also entering fresh or brackish waters, but not recorded as such by FishBase. As a whole, the database harbours 101779 occurrence records (i.e. single species–drainage basin couples).

A native or exotic status was assigned to each species occurrence record based on the information provided by the sources and further checked (see ‘Technical Validation’). An exotic species is defined as a directly or indirectly (e.g. via artificial channels) introduced species that established in the considered drainage basin. Exotic species included in the database are supposed to complete all their life cycle in the considered basins and to present self-sustaining populations in those basins. When an exotic species occurrence was acknowledged to be unsuccessful (i.e. failure of establishment of that species in the drainage basin) or needing regular release of new individuals to maintain the presence of the introduced
species (i.e. stocking), the species was not included in the basin’s species list. Species that might be
globally extinct or extirpated from a drainage basin were considered as native in the database because the
inventory of freshwater fish diversity loss was not targeted (see for instance Dias et al.\textsuperscript{12} for a compilation
of extirpated freshwater fish species for Western Europe and North America).

Drainage basin location and names
Each drainage basin was assigned a unique name that can be used as an identifier and was characterized
by its location in one of the eight terrestrial biogeographic realms (as described by Olson et al.\textsuperscript{27}; Fig. 1),
the country (or main country for shared drainage basins), its endorheic or exorheic type of water flow, its
geographic coordinates at the river mouth (for exorheic drainage basins), the geographic coordinates of
its centroid and its drainage surface area.

A specific geographic referential (Fig. 1) was built by modifying the 30 sec HydroSheds layer\textsuperscript{28} to
improve the delimitation and accuracy of drainage basins. For instance, some small coastal drainage
basins were included in one single HydroSheds polygon but were considered as separate basins because
having distinct outlets to the sea. Some drainage basins from oceanic islands have no HydroSheds code
simply because not considered in the HydroSheds shapefile. Maps and geographic information available
in the compiled literature and web-based sources were used to locate, name and improve our drainage
basins layer, complemented by country and continental scale geographic data (e.g. Faunafri project for
the African continent; http://www.poissons-afrique.ird.fr/faunafri/) and local topographic maps. This
new geographic referential is provided as a shapefile to facilitate future uses of the database.

Updates and limitations
Species are continuously being discovered and freshwater fishes are no exception, even in well-know
regions\textsuperscript{29}. Rivers are also continuously being explored and re-explored by freshwater scientists. The
database is obviously not complete and definitive, and we aim to support the database with regular
updates, ideally with bi-annual steps, depending on the resources and funding. Three main factors will be
considered in future updates: (1) new or previously non available data sources with species lists or records
for additional drainage basins or drainage basins already present in the database; (2) distribution of newly
described species; and (3) nomenclature changes in the taxonomic classification. The technical validation
procedures described below will also be applied to any new information included in the database.
Researchers having access to new data that want this information to be included in the database can send
the references or data to the corresponding author PAT. This information will be included, after
validation, in the next update release. The resulting new versions of the database will be released through
Figs hare and also through the more specialized Freshwater Biodiversity Data Portal (http://data.
freshwaterbiodiversity.eu/) to ensure the long-term availability of the database.

All biogeographic realms are well represented in terms of surface coverage (Fig. 1). There are however
some regional gaps that will be gradually filled in the next updates of the database. For instance,
Indonesian islands, coastal rivers of Peru and Northeast Brazil are regions where only few drainage basins
are informed in the database. In these regions the scarce existing information is not easily available.
Southeast Asia is the less well represented region in terms of surface coverage (Fig. 1), which is certainly
related to the low number of freshwater taxonomists working in this highly diverse region\textsuperscript{30}. All these
spatial gaps in the database will be prioritized in future updates through literature and web-based sources
monitoring.

Data Records
The database is organised in three datasets and one shapefile: the species occurrence records, the drainage
basins and the information sources table. The three tables are in csv format (columns separated by
comas) and the shapefile in ArcGis shp format (Data Citation 1). The drainage basins table is given in .csv
and shapefile formats. Both formats can be linked to the species occurrence table using the unique
drainage basin names to visualize and analyse species distribution using any adapted software (e.g. R or
QGIS, http://qgis.osgeo.org).

A. The species occurrence records table has six columns: (1) the name of the drainage basin, (2) the
scientific name of the fish species according to the information source, (3) the native or exotic status
of the occurrence records, (4) the taxonomic serial number (TSN) from the Integrated Taxonomic
Information System (ITIS, https://www.itis.gov/) when available, (5) the FishBase code when
available, (6) the FishBase or Catalogue of Fishes valid scientific name at the time of releasing the
database, (7) the occurrence status which can be either ‘valid’ or ‘questionable’ (see Technical
Validation).

B. The geographic information on drainage basins is organised in nine columns given in table and
shapefile formats: (1) the unique drainage basin name, (2) the main country where it belongs, (3) the
corresponding biogeographic region, (4) the endorheic or exorheic status, (5) and (6) the longitude
and latitude coordinates of the drainage basin outlet to the sea (only for exorheic drainages), (7) and
(8) the centroid longitude and latitude coordinates of the drainage basin, (9) the surface area of the
drainage basin.
C. The information sources table has three columns: (1) the drainage basin names, (2) the type of information sources (e.g. published paper, book, report, online database, PhD Thesis), (3) the references used to build the freshwater fish species list for the corresponding drainage basin.

**Technical Validation**

**Taxonomic validation**

Each species name found in a given information source was confronted to the valid and synonym species names lists from FishBase and the Catalogue of Fishes to ensure the validity of the identifications provided in the information source. After taxonomic validation, 103 invalid (unknown) species names were excluded from the database.

**Species distribution and status validation**

Occurrence records were carefully reviewed by the database contributors. When several information sources were used to compile the species list of a given drainage basin, particular attention was given to cross check the occurrence records and ensure a good spatial representation of the drainage basin to avoid (or at least minimize) incomplete species lists. Because occurrence data available in FishBase is often incomplete, FishBase was only used as a secondary source to collect species distribution data and to check that the resulting species distributions from our database corresponded to the broad information given in FishBase.

Because only a few (mostly migratory) freshwater species can occur in more than one biogeographic realm, the distribution of every species occurring in more than one realm was carefully verified. Similarly, for species distributed in a single realm, when one or more occurrences were inconsistent with the actual known distribution of a species (i.e. the presence in a drainage located far away from a group of drainages where the species is known to occur), these occurrences were qualified as ‘questionable’. Particular care was taken with the occurrences of all species considered exotic at least in one drainage basin. The native and exotic distributions of those species were carefully checked to avoid any status error.

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Author Contributions
P.A.T. wrote the first draft of the manuscript, and all authors contributed substantially to finalising this manuscript. P.A.T. and O.B. entered occurrence data, revised the information sources and compiled the database, with contributions of all authors. C.J. and J.-F.C. handled the geographic data related to the location and delimitation of the drainage basins, and all authors contributed to checking the information on distribution and status of the species. T.O. and S.B. initiated and designed the Database.

Additional information
Competing interests: The authors declare no competing financial interests.

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