

Scientific Life

Sustainable Biodiversity
Databasing:
International,
Collaborative, Dynamic,
Centralised

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The World Register of Marine Species (WoRMS) is a sustainable model of international collaboration around a centralised database that provides expert validated biodiversity data freely online. This model could be replicated for the over 1.2 million terrestrial and freshwater species to improve quality control and data management in biology and ecology globally.

Biodiversity Informatics

A natural consequence of the age of information technology has been the emergence of biodiversity informatics and its associated software tools and data systems that transform the speed and capacity to do research [1]. However, such tools are entirely dependent on the quality and quantity of the resource's content. In biology, the key biological standard is the species concept and the primary data are where and when species have occurred.

One of the greatest problems in biological and ecological sciences is that there are at least 1.5 million named species but several times more scientific names than biological species, plus numerous misspellings and conflicting classifications that compromise data management and complicate usage of species data by

non-taxonomists [2]. For example, only one third of over 1 million scientific names of plants are accepted for use (<http://www.theplantlist.org>). These problems are amplified when we try to compare data across studies, geographically, and over time. Society would benefit from increased effort in biodiversity science to describe all species and quality assure taxonomy, but this needs collaboration and coordination [2,3]. The cost of this ignorance due to loss of biodiversity and food production (e.g., over-fishing), and mistakes in natural resource management, is likely to be in US\$ billions per year [3]. Here we use WoRMS as an exemplar to show that collaboration around a centralised dynamic taxonomic database can significantly advance this effort.

Changing Nomenclature and Classification

In addition to correcting errors in the literature, species names and their classification change because new knowledge results in new species, taxonomic revisions that discover synonyms, and reclassification of existing species. Users of taxonomic data find it challenging to keep track of such changes, especially when conflicting information is found in published literature and online, and it can be difficult to access some publications. This can create problems in conservation where a 'species' may receive greater or lesser protection depending on its taxonomic status [4] and in the management of databases that use species names.

Important online nomenclatures capture the names of species, such as the Index of Organism Names (mostly animal names and includes the former Zoological Record), The Plant List, Index Fungorum, MycoBank, and ZooBank. However, with the exception of The Plant List, these are not edited by experts to be comprehensive and to clarify which names are

accepted, and none is continuously updated.

Users of species names include individuals, institutes, and other globally important scientific databases, such as GenBank (<https://www.ncbi.nlm.nih.gov/genbank>), Barcode of Life Database (<http://www.boldsystems.org>), Map of Life (<http://www.mol.org/>), Tree of Life (www.tolweb.org), traits databases (e.g., Try www.try-db.org/), FishBase (www.fishbase.org), the Global Biodiversity Information Facility (www.gbif.org), and Ocean Biogeographic Information System (www.iobis.org). It is essential for such users to know which names are valid, which invalid and which are questionable and how the application of a name has changed over time [5]. Making quality-assured expert knowledge easily accessible is one of the challenges facing science and society in many fields. These kinds of biodiversity databases depend on a dynamic, updated, expert-validated taxonomy of species names. However, they are not designed technically and socially to provide such a service. GBIF and OBIS operate distributed models that publish datasets requiring checking of taxonomic names, but to date have been conveyors of such content rather than validating or curating it themselves [6].

The present system of *ad hoc* publications of taxonomic information and distributed online resources cannot efficiently address this challenge. Publications are often behind a paywall rather than open access and are not easily searched. Moreover, individual papers typically deal with a subset of species of a taxon, making it necessary to extract information from multiple publications that may employ conflicting classifications and different data standards. Online name catalogues produced and maintained by one or a few experts have a high risk of stalling when their champions become unable to contribute further due

to lack of funding, illness, or other factors. Succession planning may start too late and be compromised by the idiosyncratic structure of the database and need for financial support to maintain it. Owners of such resources may lack the professional informatics expertise to manage and future-proof such databases. However, one model is overcoming these challenges in the taxonomic community, particularly for those working on marine species and their relatives.

Benefits of a Dynamic Collaborative Database

WoRMS has now been established for over 10 years [7] (Box 1). WoRMS currently contains over 243 000 valid marine species and 600 000 marine taxonomic names and has over 130 000 unique users per month. It is updated continuously, with amendments typically occurring every few minutes, and archived monthly with a digital object identifier (DOI). Despite its origins, it now also includes freshwater and terrestrial relatives of some taxa, including Mollusca, Porifera, Polychaeta, and Crustacea. Its content is gradually expanding to include geographic information, images, introduced and conservation status, fossils,

and species traits [8]. WoRMS has been meticulous in documentation of data sources. Any information item entered in WoRMS can be linked to one or more sources, providing traceability of decisions and offering another layer for data users to consult.

WoRMS and associated database content are managed by almost 500 invited experts, about 300 of whom are taxonomists [7,9,10]. The ingredients for its sustainability include this social network of experts and a professional data management team and a clear agreement that the intellectual property rights of the content rest with the Editorial Board and are managed by its Steering Committee (Box 1) [11]. Involvement of IT experts allows scientists to focus on their expertise and enables them to reach out for help when needed. Our experience has been that to engage experts from different countries and communities requires clarity on resource ownership, management, succession planning, and peer-to-peer engagement (i.e., community building). Experts are more likely to spend their time on a database if it is open access, easy to use, sustainable, and quality assured, and they get most recognition when it is

considered meritorious and prestigious in their community (Table 1). They, and in some case their employers, are also more willing to provide their time as editors if the product directly contributes to society, other institutions or individuals do not profit by their work (e.g., research funding, authorships), and they have a legal right to influence its management (e.g., electing a steering committee). Thus, a global-scale, community-owned and governed, open-access, continuously edited, centralised database, with professional informatics support, can provide a cost-effective sustainable service to science and society (Table 1). Other community-driven bottom-up efforts, like Wikispecies and its associated Wikidata, also deserve support. The same is true for initiatives driven by the scientific community like the Tree of Life web project, which is an excellent expert-edited introduction to biodiversity and valuable tool for teaching, but comes in a narrative format that precludes automated analysis.

Although the conventional process of peer-reviewed literature serves the taxonomic community well, access to an expert-edited database where mistakes and omissions can be promptly corrected and new knowledge made freely available is widely appreciated by the broader scientific community, as evident from the over 5000 citations of WoRMS in Google Scholar. Other online biodiversity resources, including Species 2000's Catalogue of Life (CoL), Wikispecies, Encyclopedia of Life, Global Biodiversity Information Facility, and Ocean Biogeographic Information System, use information on marine species from WoRMS to support their information systems, demonstrating good cooperation within the biodiversity informatics community. Although the first two of these are expert edited, CoL lacks a centralised database and, with the exception of data provided by WoRMS, most of its datasets are irregularly

Box 1. Evolution of WoRMS

The origin of WoRMS was the European Register of Marine Species (ERMS), a conceptually similar expert-validated checklist of all European marine species. This was freely available online and conventionally published [14]. To avoid complications concerning intellectual property rights with so many editors and authors, the ownership of ERMS was vested in a nonprofit, limited-liability, legally incorporated scientific society: the Society for the Management of Electronic Biodiversity Data Ltd (SMEBD). SMEBD had a deliberately broad remit to support other electronic biodiversity databases (at least in Europe). However, as these did not transpire, its responsibilities were later transferred to the Editorial Board of WoRMS, which includes editors of ERMS and an increasing number of child databases with regional, taxon-specific, or thematic (e.g., introduced species) foci. Instead of supporting core operations, financial contributions from research grants and donations from users are used to fill gaps in and expand WoRMS content.

Shortly after the completion of ERMS, the Flanders Marine Institute (VLIZ) Data Centre offered to host it because having validated species names was critical to its marine biological data management. This relieved the editors of needing to worry about the informatics aspects of a biodiversity database. The hardware, software, and 24-7 online access were now permanently supported by a team of data management specialists at a professional, government-funded data centre. This security encouraged more experts to become editors and provided the computer platform for WoRMS [10]. The involvement of a core group of taxonomists in the Census of Marine Life and its Ocean Biogeographic Information System provided the social network to expand ERMS to WoRMS [15]. WoRMS is now governed by a committee elected by its editors, which has a formal agreement with the database host institute [7].

Table 1. Fifteen Benefits of Collaborative Management of a Biodiversity Database

To science	To users	To the database editors
1. Improved quality control in biodiversity science	1. Ease of access to an electronic, standardised, authoritative species list that is classified hierarchically for use in their own data management	1. Focussed collaboration with colleagues internationally that can aid their personal knowledge and know-how
2. Gaps in knowledge are more visible and encourage researchers and funding agencies to fill them	2. Automated tools to classify and check spelling of species names	2. Information organised and archived, reducing the need for the expert to have his or her own database
3. Reduction in misspellings and incorrect use of species nomenclature	3. More time-efficient to consult a single authoritative source than to research and assess accuracy of numerous disparate sources	3. Citable electronic publication that is digitally archived
4. Rapid conversion of outdated species names into state-of-the-art name, particularly outside a researcher's own field of expertise	4. Contact details of experts easily found	4. Peer recognition; it is prestigious to be invited as an expert to edit the database
5. Easy access to the reference for the original species description, many of which are rarely cited elsewhere	5. Relieves initiatives focussing on other (non-taxonomic) aspects of biodiversity of the need to keep track of taxonomic changes themselves	
6. Standardisation and integration makes it easier to conduct global syntheses of information		

updated. However, the current GBIF implementation plan (<https://www.gbif.org/document/36j6HhbR4kOMY6oqcEMk/gbif-implementation-plan-2017-2021-and-annual-work-programme-2018>) involves collaboration with CoL and others to provide a complete, literature-referenced, automated, and expert-validated world species list (D. Hobern, personal communication.).

We suggest that the WoRMS model of international collaboration should be used to provide a quality-assured taxonomy for all species on Earth and to support other biodiversity-related databases to make expert knowledge openly available to society. An additional benefit of this collaboration is that it becomes easier to conduct global syntheses of taxa because the information is standardised in the database [5]. For example, WoRMS enabled a world synthesis of how many marine species are named and might exist (e.g., [12]) (Table 1). A global-scale, expert-driven, collaborative, and centralised open-access database could thus be available for all species on Earth and was recently called for by conservation biologists and taxonomists [4,13]. This

is essential to provide a current taxonomy for all other biodiversity databases and publications. Following this, perhaps the next gap to be filled will be an identification guide to all life on Earth that links databases with literature and images: a 'key to all life'.

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The Art of Scientific Performance

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