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Towards a new global database of regional red lists (RegRed): metadata

Ivo Kadlec, Adam Uličný, 🕩 Petr Keil, 🕩 Florencia Grattarola

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4 Ivo Kadlec^{1,2}, Adam Uličný¹, Petr Keil¹, Florencia Grattarola¹

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⁶ ¹Deprtment of Spatial Sciences, Faculty of Environmental Sciences, Czech

- 7 University of Life Sciences Prague, Kamýcká 129, Praha Suchdol, 16500, Czech
 8 Republic
- ⁹ ²Department of Ecology, Faculty of Environmental Sciences, Czech University of Life
- 10 Sciences Prague, Kamýcká 129, Praha Suchdol, 16500, Czech Republic
- 11

12 Corresponding author: Ivo Kadlec; <u>kadleci@fzp.czu.cz</u>

13

14 Abstract

Conservation biology depends on an assessment of threats to species. This has 15 been done at the global scale through the IUCN red list, and also locally and 16 17 regionally, e.g. through country-specific regional red lists. The latter quantify the level 18 of threat to a species in a region, irrespectively to its global status (e.g., a species 19 can be non-threatened globally, but threatened or extirpated in a specific region). 20 There are efforts to collate these regional red lists (e.g., the NRL database hosted by ZSL in collaboration with the IUCN National Red List Working Group), but these 21 22 have gaps, and are in the process of redevelopment with increased input from 23 country focal points.

Here, we announce a renewal of the effort to collate regional red lists. To create it we 24 25 searched and compiled sources containing species threat assessments all over the 26 world. As a result, we found 2,093 sources in 172 countries, covering 487 broad taxonomic groups. In this paper, we provide the compiled metadata, enriched with 27 geographical and taxonomic information, and details about the source's title, URL, 28 29 file format, language, and publication date. This is step one in our effort, in which we 30 ultimately plan to digitise all the compiled sources and provide them openly. By 31 announcing this effort here, we aim to actively seek to expand the metadata database and to collate the respective data. Please refer to the section "How to 32 engage?" if you are interested in collaborating with us. 33

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35 Keywords: conservation biology, endangered species, IUCN, data compilation,

36 conservation assessment

37 Introduction

38 Species extinction represents one of the most serious and irreversible consequences of global environmental change, and currently, it is estimated to be between 100 and 39 1000 times higher than the natural rates (Ceballos et al. 2015, 2017). In order to 40 41 monitor and prevent species loss, conservation scientists and practitioners have 42 developed systematic approaches to assess the risk of extinction, with Red Lists emerging as the main tool for this purpose (Mace and Lande 1991). The most 43 44 extensive global Red List, made by the International Union for Conservation of 45 Nature (IUCN), is now the gold standard and reference for determining a species' risk of extinction on a global scale (Rodrigues et al. 2006). 46

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Using standardised criteria based on population size, distribution range, and rate of 48 decline, the IUCN Red List has developed into a sophisticated system since its 49 beginning in 1964 (Mace et al. 2008). To guarantee scientific rigour and uniformity in 50 evaluating extinction risk across various taxonomic groups, these standards have 51 been regularly improved (Mace et al. 2008, IUCN Species Survival Commission 52 2012). However, even though the IUCN Red List has evaluated over 160,000 53 species as of 2025 (IUCN 2025) and successfully captures the risk of *global* 54 55 extinction, it does not represent the complicated reality of species status at the 56 regional and national levels. This is an important limitation because species can face 57 local extinction or significant population decline within specific countries while 58 maintaining populations elsewhere (Gärdenfors 2001, Drago and Vrcibradic 2020, Brodsky et al. 2023). To address this limitation, many countries and regions have 59 developed their own Red Lists following IUCN categories and criteria (IUCN Species 60 61 Survival Commission 2012) or other documents of a similar nature, e.g. government 62 lists and decrees of threatened species (Gärdenfors et al. 2001, Miller et al. 2007).

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There have been several projects concerned with regional red lists. The most 64 notable is the National Red List project (NRL; https://www.nationalredlist.org/). The 65 NRL was created by the IUCN National Red List Working Group, and is hosted by 66 the Zoological Society of London (ZSL). Their updated website provides access to a 67 subset of predominately mammalia species from the original database of 101 68 69 sources from 73 countries (accessed in April 2025) and offers browsing by 70 taxonomic group, species or source. Although the project made a considerable effort 71 to systematise local red lists there was a hiatus in data transfer whilst the database 72 and website underwent significant restructuring to facilitate national focal points in 73 building capacity to contribute and manage their own data (Sophie Ledger from NRL, 74 pers. comm.). Another effort to collate red lists at the sub-global level is the project "PaDRE - Patterns and drivers of regional plant extinctions", jointly led by the 75 Helmholtz Centre for Environmental Reasearch - UFZ, the German Centre for 76 77 Integrative Biodiversity Research Halle-Jena-Leipzig (iDiv), and Martin-Luther 78 University Halle-Wittenberg (Staude et al. n.d.). Other taxon-specific initiatives have also made important contributions, such as the Red Data Book of European 79 80 Butterflies (Swaay et al. 1999), which assessed all 576 butterfly species in each 81 European country against IUCN criteria.

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83 Here, we introduce RegRed, a project that aims to create a comprehensive database 84 of available red lists and sources containing information about threatened or extinct species at the national/regional level. We aim to build on, and significantly expand, 85 the red list data that have already been put together (e.g. in NRL). Our aim is not to 86 supersede other projects, such as the NRL, but rather to complement them, as we 87 88 recognize meaningful differences between our efforts. While the NRL prioritises red 89 lists submitted by national representatives, our objective is to compile a broader range of data from diverse sources to achieve more comprehensive global coverage. 90 91 By consolidating these data in a single open database, we aim to make these 92 essential conservation tools more accessible to both practitioners and other

- 93 stakeholders interested in biodiversity conservation.
- 94
- 95 The RegRed project has two phases:
- 96 1) In the first phase, described in this paper, we have manually searched for, and catalogued, a comprehensive list of known published regional red lists 97 (hereafter metadata). Even though these metadata do not yet provide the per-98 99 species assessments, we see them as large and useful enough to warrant a 100 stand-alone publication. The metadata can serve both researchers, policymakers, and hobbyists to quickly access relevant documents for their 101 102 needs. Some of the documents, mainly Red Books, even contain photos, illustrations, maps and locally relevant species information possibly serving as 103 104 excellent freely available field guides or handbooks.
- 105 2) In the second phase, not yet described here, we plan to digitise and database the complete sources listed in the metadata. Since we are open for coordination and collaborations on this effort, we also see it as important to first release the metadata in this paper (see the "*Future directions and how to engage*" section below).
- 110

111 Methods

Our metadata-gathering workflow primarily relied on search engines such as Google 112 113 Search, Google Scholar and ResearchGate. Both national and university libraries proved valuable, and most official IUCN red lists were sourced from the IUCN library. 114 For the search, we established a set of basic keywords that we consistently used for 115 initial searches, adapting them when necessary to suit specific states/regions (e.g. 116 "red list of...", "endangered fauna of ...", "checklist of...", "list of endangered species 117 of..."). We worked systematically through continents, focusing on one country at a 118 119 time (i.e level 0 in GADM, the Database of Global Administrative Areas) to identify all available resources. When sub-national red lists were found for a given country, we 120 121 recorded them associated with their GADM level 1 or 2. In the cases in which we 122 could not assign sources to standard GADM levels, we categorized them as "custom regions." These custom regions could represent sub-national areas (e.g. Table 123 124 Mountain National Park), large territories spanning administrative boundaries, or 125 collections of several countries (e.g. the Arabian Peninsula or southern Africa).

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Each source containing species threat assessments was entered into our database
with detailed geographical resolution and taxonomic scope. Types of sources
included official red lists, checklists, government documents, scientific publications,
and any other resources where species threat categories were defined. For clarity
and to facilitate subsequent filtering, we assigned each entry to the most specific
geographical level possible (country, state, region) and identified the lowest common
taxonomic unit.

- 134
- 135 In many cases, a single source covered multiple taxonomic groups, resulting in
- 136 multiple database entries (hereafter a "record", a single source per taxonomic
- 137 group). For example, a "Red list of Reptiles and Amphibians" would generate two
- 138 separate records—one for reptiles and one for amphibians. Similarly, documents
- 139 covering "Fungi and Flora" would be separated into two distinct records. This splitting
- approach was applied wherever practical to enhance the database's utility for
- researchers focusing on specific taxonomic groups. Finally, we matched the taxa
- names against GBIF's taxonomic backbone (GBIF Secretariat 2023) to normalise
- names and include the higher taxonomic ranks (i.e., kingdom, phylum, class).

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- 145 For each record, we also collated comprehensive metadata including the source
- 146 URL, original title, file format, language, and publication date (see the full list of data
- 147 columns in Table 1). The database can be accessed at
- 148 <u>https://github.com/petrkeil/RegRed_metadata</u> or through Zenodo (Kadlec et al.,
- 149 2025).

Column Name	Definition
id	Unique identifier for each record.
continent	The name of the continent in which the record occurs.
level_0	The name of the country according to GADM in which the record occurs.
level_1	The name of the first smaller administrative region than country (e.g., states, provinces) according to GADM in which the record occurs.
level_2	The name of the first smaller administrative region than country (e.g., districts, counties) according to GADM in which the record occurs.
region_custom	The name of the non-standard geographic areas not fitting into GADM levels (e.g., Carpathians, Indochina, Dutch Caribbean) in which the record occurs.
region_detail	The list of countries or regions spanning the custom regions in which the record occurs.
iso_2	The two-letter country code according to ISO standards. For non-standard geographic areas (i.e., custom regions) this is a list of the two-letter country codes it overlaps.
taxa	The verbatim taxonomic classification of the record. The value of taxa is not always a scientific name; it may include ecological groups (e.g., "cave animals"), functional groups, or broader categories (e.g., "animals").
kingdom	The full scientific name of the kingdom in which the taxa is classified.
phylum	The full scientific name of the phylum in which the taxa is classified.
class	The full scientific name of the class in which the taxa is classified.
order	The full scientific name of the order in which the taxa is classified.
family	The full scientific name of the family in which the taxa is classified.
source_name	The original name of the source.
source_link	The URL to access the source.
language	The language in which the source is written.
year	The year of publication of the source.

Table 1: Metadata table column names and definitions.

151 Coverage

The metadata identifies a total of 3,200 records derived from 2,064 unique sources, covering 170 countries globally (Fig. 1), 223 sub-national regions (GADM level 1), and 83 regions (i.e., non-standard geographic areas). These sources, produced over the past 50 years (Fig. 2), span 483 distinct taxa (Fig. 3), providing comprehensive coverage across the Tree of Life.

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Geographically, the metadata cover all continents, with the highest density of records
 in Europe, followed by Asia and Africa (Fig. 1). This pattern likely reflects both the
 prevalence of conservation assessment activities in these regions and our search

161 methodology prioritizing more accessible documentation. Conversely, the large

number of sources may only reflect the amount of taxa assessments at sub-national
 levels and not the amount of species covered. For example, Germany has 544 insect

164 aroup assessments spanning a few species, while the United States makes their

165 data accessible as single sources spanning multiple taxa (e.g., fauna).

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Figure 1. Global distribution of regional red list sources by country. Countries
 are colored according to the number of unique source documents identified in our
 metadata collection, ranging from no sources (light gray) to 20 or more sources (dark
 red). Note that each source may contain multiple regional redlists.

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173 Regarding temporal coverage, our database includes sources published between
174 1975 and 2024, with almost half of sources (49%) published within the last decade

175 (Fig. 2). The notable peaks in the number of sources per year around 2003, 2010,

and 2020, possibly follow the establishment of the IUCN's guidelines for regional red

177 listing in 2003, the increased focus on conservation put by the Aichi Biodiversity

178 Targets in 2010, and the Post-2020 Global Biodiversity Framework report in 2020.

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Figure 2. Number of regional red list sources published per year. Sources cover from 1975 to 2024, showing a gradual increase beginning in the early 1990s, with

183 notable peaks around 2003, 2010, and 2020.

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185 Taxonomic distribution within our collection reveals significant variation in coverage (Fig. 3). Globally, our metadata collection contains 2,190 records for Animalia, 898 186 187 records for Plantae, and 105 records for Fungi. The Arthropoda phylum dominates 188 with 978 sources, while Chordata represents 815 sources (with vertebrates accounting for 668 sources total), and Mollusca accounts for 71 sources. Breaking 189 190 down the Chordata sources, we find 202 addressing mammals, 185 covering birds, 191 and the remainder distributed across reptiles (148), amphibians (133), and fish (102). For Arthropoda, insects account for 854 sources and arachnids 47. We also 192 identified 520 broader sources covering multi-taxonomic information. The taxonomic 193 194 coverage varies considerably across geographic regions (Fig. 3).

195

196 Europe has the most comprehensive taxonomic coverage with 2,182 records. predominantly focused on Animalia (1,550 records), followed by Plantae (549 197 records) and Fungi (75 records). Asia has the second highest number of records 198 (452), similarly dominated by animal assessments. In other continents, the pattern of 199 animal-focused assessments prevails, with significantly lower representation of plant 200 and fungal records. The relative scarcity of fungal records is especially notable, with 201 minimal representation outside Europe, reflecting the historical neglect of this 202 kingdom in conservation assessment efforts despite its ecological importance. 203

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Figure 3: Number of records by continent and taxonomic kingdom (Animalia,

Fungi, and Plantae). The three kingdoms are covered in all continents except for

208 South America and Oceania, revealing the predominance of animal records,

209 particularly in Europe, with more balanced kingdom representation in other regions.

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211 Limitations

Our initial assessment revealed that while many countries have published national or regional red lists, their accessibility varies considerably. Some jurisdictions maintain well-organized, freely accessible databases and websites. However, in most cases, these lists are difficult to locate, scattered across various platforms, or hosted on defunct websites. This highlights the value of our compilation effort in making this information more discoverable and accessible to the global conservation community.

Beyond accessibility issues, the collected sources themselves vary in quality, year of publication, language, type, and format. Some are scanned documents and books with little or no optical character recognition (OCR) processing. Not all sources were prepared following the IUCN categories and criteria, which have evolved over time and are currently in version 3.1 (IUCN 2001). Due to language constraints and the project's scale, some records may not include the advertised data, though such cases should be limited.

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We detailed the taxonomic higher hierarchy of the groups covered by the sources.

228 However, some records titled with more general descriptors like "Fauna" may

encompass a mix of taxa. For instance, a "Fauna" red list may include vertebrates
and invertebrates while others include only birds and mammals. The same issue
applies to Flora and other general titles.

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The collection is provided as-is, with corrections and additions to be updated in the GitHub repository. Some links may become inaccessible over time, a common issue with large-scale global data hosted primarily on government websites. Additionally, certain websites may experience intermittent downtime due to server-side issues. We have verified the validity of the URL's at time of publication. If you encounter any dead links, please notify us at the email address provided in the "How to engage?" section of this paper.

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241 Future directions and how to engage

242 We decided to publish these metadata as a means to engage with others,

243 decentralize the effort, and encourage collaboration. Our overarching goal is to

244 compile available per-species national and regional threat data into a database for

further use and sharing. Our team is committed to open science, and the final

246 database will be made available either through a web interface and as a

247 downloadable dataset under a CC-BY license.

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The initial focus of the data compilation will be on vertebrate species. Currently, we are developing efficient methods for large-scale and accurate data digitisation. If these methods prove effective, the database may be expanded to include additional taxa.

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We are actively seeking to expand the metadata database and to collate the respective data. If you have information regarding regional red lists that we may have missed, identify errors in our data or are interested in collaborating, please contact us at RegRed@fzp.czu.cz.

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