# Package 'redbookperu'

July 23, 2025

**Title** Access and Analyze Data from the Red Book of Endemic Plants of
Peru

**Version** 0.0.3

**Description** Provides access to and analysis of data from ``The Red Book of En-
demic Plants of Peru'' (León, B., Roque, J., Ulloa, C., Jorgensen, P.M., Pit-
man, N., Cano, A. 2006) <doi:10.15381/rpb.v13i2.1782>. This package offers comprehen-
sive taxonomic, geographic, and conservation information about Peru's en-
demic plant species. It includes functions to verify species inclusion, obtain updated taxo-
nomic details, and explore the dataset.

**License** MIT + file LICENSE

**Suggests** knitr, rmarkdown, testthat (>= 3.0.0)

**Config/testthat/edition** 3

**LazyData** true

**LazyDataCompression** xz

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**URL** <https://github.com/PaulESantos/redbookperu>,

<https://paulesantos.github.io/redbookperu/>

**BugReports** <https://github.com/PaulESantos/redbookperu/issues>

**Depends** R (>= 2.10)

**Maintainer** Paul E. Santos Andrade <paulefrens@gmail.com>

**NeedsCompilation** no

**Author** Paul E. Santos Andrade [aut, cre] (ORCID:
<https://orcid.org/0000-0002-6635-0375>),
Lucely L. Vilca Bustamante [aut] (ORCID:
<https://orcid.org/0000-0002-5559-1296>)

**Repository** CRAN

**Date/Publication** 2024-07-02 07:30:02 UTC

# Contents

---

check_redbooklist *Check Species Names in the Red Book of Endemic Plants of Peru*

---

#### Description

This function checks a list of species names against the Red Book of Endemic Plants of Peru database and provides information about whether a species was recorded as endemic, and checks for misspelling typos (fuzzy match).

#### Usage

```
check_redbooklist(splist, dist = 0.02)
```

#### Arguments

splist        A character vector containing the species names to be checked.

dist          Maximum allowed distance for fuzzy matching of species names.

#### Details

This function checks each species name in the provided list against the Red Book of Endemic Plants of Peru database using fuzzy matching based on the specified maximum distance (`dist`). It provides information about the endemic status of each species and flags if the recorded name needs updating. It also counts the number of exact and fuzzy matches found.

#### Value

A character vector indicating if each input species name is listed as "endemic" in the Red Book of Endemic Plants of Peru database. Returns "endemic" if the species name is listed and "not endemic" if no matching entry is found.

#### References

Red Book of Endemic Plants of Peru The World Checklist of Vascular Plants, a continuously updated resource for exploring global plant diversity. Taxonomic Name Resolution Service - TNRS Plants of the World Online - Facilitated by the Royal Botanic Gardens - Kew.

## Examples

```
# Example usage of the function
splist <- c("Aphelandra cuscoenses",
            "Piper stevensi",
            "Sanchezia ovata",
            "Verbesina andina",
            "Festuca dentiflora",
            "Eucrosia bicolor var. plowmanii",
            "Hydrocotyle bonplandii var. hirtipes",
            "Persea americana")

# Basic usage
check_redbooklist(splist = splist, dist = 0.2)

# Using base R with a data frame
plant_list <- data.frame(splist = splist)
plant_list$label <- check_redbooklist(plant_list$splist, dist = 0.2)
plant_list
```

---

get_redbook_data             *Get Red Book Data for Given Species List*

---

## Description

This function retrieves comprehensive information from the Red Book of Endemic Plants of Peru database for a provided list of species. It associates the provided species names with their corresponding updated taxonomic information and descriptions recorded in the original publication.

## Usage

```
get_redbook_data(splist, dist = 0.1)
```

## Arguments

splist        A character vector containing the species names to be queried.

dist          Maximum allowed distance for fuzzy matching of species names.

## Details

This function checks each species name in the provided list against the Red Book of Endemic Plants of Peru database using fuzzy matching based on the specified maximum distance (dist). For each species, it retrieves and combines taxonomic information (accepted name, accepted family, accepted name author) with additional descriptive data recorded in the original publication, such as IUCN conservation category, bibliographic reference, collector, herbariums, common name, departmental registrations, ecological regions, protected natural areas (SINANPE), Peruvian herbaria, and additional remarks.

## Value

A data frame containing comprehensive information about the provided species, including updated taxonomic details and descriptions.

## References

Red Book of Endemic Plants of Peru The World Checklist of Vascular Plants, a continuously updated resource for exploring global plant diversity. Taxonomic Name Resolution Service - TNRS Plants of the World Online - Facilitated by the Royal Botanic Gardens - Kew.

## See Also

check_redbooklist function for a more focused check of species endemic status.

## Examples

```
# Example illustrating how to use the get_redbook_data function
species_list <- c("Aphelandra cuscoensis", "Sanchezia ovata", "Piper stevensii")
redbook_data <- get_redbook_data(species_list)
head(redbook_data)
```

---

| redbook_position | *List of the number positions of the first 3 letters of the species name in the redbook_tab* |
|---|---|

---

## Description

The 'redbook_position' reports the position (in term of number of rows) of the first three letters (triphthong) for the plant names stored in the variable 'accepted_name' of the table 'redbook_tab'. This indexing system speeds up of the search on the largest list using the package.

## Usage

```
redbook_position
```

## Format

A data frame with 978 observations on the following 3 variables:

**position** A character vector. The position of the first three letters of the species name in the redbook_tab.

**triphthong** A character vector. The first three letters of the species name in the redbook_tab.

**genus** A character vector. The corresponding genus name.

## Details

Positions of Species Names in The Red Book of Endemic Plants of Peru

The `redbook_position` dataset provides the positions of the first three letters of each species name listed in the `redbook_tab`.

## Examples

```
data("redbook_position")
head(redbook_position)
```

---

redbook_sps_class          *List of Species Names in redbook_sps_class Separated by Category*

---

## Description

The `redbook_sps_class` dataset includes all species names separated by genus, epithet, author, subspecies, variety, and their position (ID) in the `redbook_tab`.

## Usage

```
redbook_sps_class
```

## Format

A data.frame with the following columns:

**species**  A character vector. The full species name.

**genus**  A character vector. The genus of the species.

**epithet**  A character vector. The specific epithet of the species.

**input_subspecies_epitheton**  A character vector. The infraspecific epithet of the species, if applicable.

**rank**  A character vector. The taxonomic rank (e.g., "species", "subspecies", "variety").

**subspecies**  A character vector. The subspecies name, if applicable.

**variety**  A character vector. The variety name, if applicable.

**hybrid**  A character vector. Indicates if the species is a hybrid.

**id**  A character vector. The ID of the species in the `redbook_tab`.

## Examples

```
data("redbook_sps_class")
head(redbook_sps_class)
```

---

| redbook_sp_data | *Database of the Red Book of Endemic Plants of Peru* |

---

**Description**

This database contains comprehensive information regarding the endemic plant species listed in the Red Book of Endemic Plants of Peru. Each endemic taxon is accompanied by corresponding variables that detail its taxonomic status, IUCN conservation category, bibliographic references, type collection details, common names, departmental registrations, ecological regions, protected natural areas (SINANPE), and Peruvian herbaria where the specimens are deposited, as recorded in the original book.

**Usage**

```
redbook_sp_data
```

**Format**

A data frame with the following variables:

**redbook_id** Unique identifier for each species in the Red Book of Endemic Plants of Peru.

**redbook_name** Scientific name of the endemic species.

**iucn** Conservation category assigned according to IUCN.

**publication** Bibliographic reference where the taxon was originally described.

**collector** Name(s) of the collector(s) of the type specimen.

**herbariums** Acronyms of the institutions where the type specimens of the taxon are deposited.

**common_name** Common names of the species as mentioned in the literature.

**dep_registry** Abbreviations of the departments where the taxon has been recorded.

**ecological_regions** Abbreviations of the ecological regions proposed by Zamora (1996).

**sinampe** Abbreviation of the Protected Natural Area where the taxon was recorded.

**peruvian_herbariums** Acronyms of the Peruvian institutions where both type and non-type specimens are deposited.

**remarks** Observations and additional information about the endemic taxon.

**Details**

This database provides essential information for research and conservation efforts related to Peru's endemic flora, offering access to the data presented in the corresponding book.

**References**

León, Blanca, et.al. 2006. "The Red Book of Endemic Plants of Peru". Revista Peruana De Biología 13 (2): 9s-22s. https://revistasinvestigacion.unmsm.edu.pe/index.php/rpb/issue/view/153

## Examples

```
# Example illustrating how to load and explore the database
data("redbook_sp_data")
head(redbook_sp_data)
```

---

| redbook_tab | *Species Names Listed in The Red Book of Endemic Plants of Peru* |
|---|---|

---

## Description

The `redbook_tab` contains records for all species listed in The Red Book of Endemic Plants of Peru.

## Usage

```
redbook_tab
```

## Format

A tibble with the following columns:

**redbook_id**  The fixed species ID of the input taxon in The Red Book of Endemic Plants of Peru.

**redbook_name**  A character vector. The species name as listed in The Red Book of Endemic Plants of Peru.

**input_genus**  A character vector. The input genus of the corresponding species name listed.

**input_epitheton**  A character vector. The specific epithet of the corresponding species name listed.

**rank**  A character vector. The taxonomic rank (e.g., "species", "subspecies", "variety") of the corresponding species name listed.

**input_subspecies_epitheton**  A character vector. The infraspecific epithet of the corresponding species name listed, if applicable.

**accepted_name**  A character vector. The accepted plant taxa names according to the World Checklist of Vascular Plants (WCVP).

**accepted_family**  A character vector. The corresponding family name of the accepted name.

**accepted_name_author**  A character vector. The author of the accepted name.

**accepted_name_rank**  A character vector. The rank of the accepted name (e.g., species, subspecies).

**tag_subsp_wcvp**  A character vector. A tag indicating if the subspecies is recognized in the WCVP.

**genus_ephitethon_wcvp**  A character vector. The genus part of the name according to the WCVP.

**species_ephitethon_wcvp**  A character vector. The specific epithet part of the name according to the WCVP.

**subspecies_ephitethon_wcvp**  A character vector. The infraspecific epithet part of the name according to the WCVP, if applicable.

## References

León, Blanca, et.al. 2006. "The Red Book of Endemic Plants of Peru". Revista Peruana De Biología 13 (2): 9s-22s. https://doi.org/10.15381/rpb.v13i2.1782.

## Examples

```
data("redbook_tab")
head(redbook_tab)
```

# Index