

Using **plantR** to Manage Animal Taxonomy with the CTFB Backbone

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Contents

1	Installing <code>plantR</code>	1
2	A practical example	2
3	Preparing names using <code>fixSpecies()</code>	2
3.1	Internal functions	2
4	Validating taxon names using <code>prepSpecies()</code>	3
4.1	Internal functions	4
5	Validating family names using <code>prepFamily()</code>	5
6	Brief code summary	5
7	Citation	6

1 Installing `plantR`

Install the packages from GitHub if needed and load `plantR`.

```
if (!requireNamespace("remotes"))
  install.packages("remotes")
library(remotes)

if (!requireNamespace("plantR"))
  install_github("LimaRAF/plantR")

if (!requireNamespace("plantRdata"))
  install_github("LimaRAF/plantRdata")

library(plantR)
```

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2 A practical example

We will start with a small list of **names of animal species** that includes common issues (misspellings, synonyms, wrong capitalization, invalid names). We will save it in an objects called `names`:

```
names <- c(  
  "Apis mellifera Linnaeus, 1758",           # name accepted with author  
  "Apis melifera",                          # misspelling  
  "Apis cf. mellifera",                     # open nomenclature  
  "Ancyloscelis armatus",                  # synonym of Ancyloscelis apiformis (Fabricius, 179  
  "Centris aenea",                         # name accepted without author  
  "Centris rufa",                           # synonym of Centris aenea Lepeletier, 1841  
  "Centris Rhodoprocta Moure & Seabra, 1960", # wrong capitalization  
  "Lutjanus purpureus",                    # synonym of Lutjanus campechanus (Poey, 1860)  
  "Parotocinclus amazonensis",            # invalid in CTFB; no synonym for this name  
  "Panthera onca",                        # name accepted without author  
  "Solenopsis bicolor (Emery, 1906)",      # name accepted with author  
  "Eucopricus columbi MacLeay, 1819",       # synonym of Sulcophanaeus columbi (MacLeay, 1819)  
  "Eucopricus sp.1"                       # incomplete identification  
)
```

3 Preparing names using `fixSpecies()`

`fixSpecies()` formats and cleans names (notation, casing, authorship split, notation flags). It accepts either a character vector or a data frame (default `scientificName` column).

```
names_fixed <- fixSpecies(names)  
names_fixed[,-c(2,4)]  
  
#> scientificName  
#> 1 Apis mellifera Linnaeus, 1758  
#> 2 Apis melifera  
#> 3 Apis cf. mellifera  
#> 4 Ancyloscelis armatus  
#> 5 Centris aenea  
#> 6 Centris rufa  
#> 7 Centris Rhodoprocta Moure & Seabra, 1960  
#> 8 Lutjanus purpureus  
#> 9 Parotocinclus amazonensis  
#> 10 Panthera onca  
#> 11 Solenopsis bicolor (Emery, 1906)  
#> 12 Eucopricus columbi MacLeay, 1819  
#> 13 Eucopricus sp.1  
  
#> scientificName.new  
#> 1 Apis mellifera  
#> 2 Apis melifera  
#> 3 Apis mellifera  
#> 4 Ancyloscelis armatus  
#> 5 Centris aenea  
#> 6 Centris rufa  
#> 7 Centris rhodoprocta  
#> 8 Lutjanus purpureus  
#> 9 Parotocinclus amazonensis  
#> 10 Panthera onca  
#> 11 Solenopsis bicolor  
#> 12 Eucopricus columbi  
#> 13 Eucopricus sp.1  
  
#> scientificNameSt  
#> 1 name_w_auth  
#> 2 possible  
#> 3 conf  
#> 4 possible  
#> 5 possible  
#> 6 possible  
#> 7 name_w_wrong_case|name_w_auth  
#> 8 possible  
#> 9 possible  
#> 10 possible  
#> 11 name_w_auth  
#> 12 name_w_auth  
#> 13 i
```

3.1 Internal functions

For this specific list, some functions may not change anything (which is fine). The goal is to illustrate correct usage of the internal functions on the **same** input vector of names.

```
fixIndet(names)      # detects undetermined names (e.g., "sp.", "indet")
```

```

#> [1] "Apis mellifera Linnaeus, 1758"           "Apis mellifera"
#> [3] "Apis cf. mellifera"                      "Ancyloscelis armatus"
#> [5] "Centris aenea"                           "Centris rufa"
#> [7] "Centris Rhodoprocta Moure & Seabra, 1960" "Lutjanus purpureus"
#> [9] "Parotocinclus amazonensis"                "Panthera onca"
#> [11] "Solenopsis bicolor (Emery, 1906)"       "Eucopricus columbi MacLeay, 1819"
#> [13] "Eucopricus sp.1"                         "Eucopricus columbi MacLeay, 1819"

fixCase(names)      # fixes casing (e.g., "Centris Rhodoprocta")

#> Apis mellifera Linnaeus, 1758                 Apis mellifera
#> "Apis mellifera Linnaeus, 1758"               "Apis mellifera"
#> Ancyloscelis armatus                          Centris aenea
#> "Ancyloscelis armatus"                       "Centris aenea"
#> Centris Rhodoprocta Moure & Seabra, 1960     Lutjanus purpureus
#> "Centris rhodoprocta Moure & Seabra, 1960"   "Lutjanus purpureus"
#> Panthera onca                                Solenopsis bicolor (Emery, 1906)
#> "Panthera onca"                             "Solenopsis bicolor (Emery, 1906)"
#> Eucopricus sp.1                               "Eucopricus sp.1"
#> "Eucopricus sp.1"                            "Eucopricus sp.1"

fixAuthors(names)    # splits taxon and author names, if present

#> orig.name          tax.name          tax.au
#> 1 Apis mellifera Linnaeus, 1758 Apis mellifera Linnaeus,
#> 2 Apis mellifera Apis mellifera Mellii
#> 3 Apis cf. mellifera Apis cf. mellifera
#> 4 Ancyloscelis armatus Ancyloscelis armatus
#> 5 Centris aenea Centris aenea
#> 6 Centris rufa Centris rufa
#> 7 Centris Rhodoprocta Moure & Seabra, 1960 Centris Rhodoprocta Moure & Seabra,
#> 8 Lutjanus purpureus Lutjanus purpureus
#> 9 Parotocinclus amazonensis Parotocinclus amazonensis
#> 10 Panthera onca Panthera onca
#> 11 Solenopsis bicolor (Emery, 1906) Solenopsis bicolor (Emery, 1906)
#> 12 Eucopricus columbi MacLeay, 1819 Eucopricus columbi MacLeay,
#> 13 Eucopricus sp.1 Eucopricus sp.1

```

4 Validating taxon names using prepSpecies()

Next, validate names against the CTFB backbone (ctfbNames) from plantRdata by loading it into the Global Environment and passing it to db.

```

# load the CTFB backbone (ctfbNames) into the Global Environment
utils::data("ctfbNames", package = "plantRdata")

# validate against CTFB
names_valid <- prepSpecies(
  names_fixed,
  tax.names = c("scientificName.new", "scientificNameAuthorship.new"),

```

```

db = ctfbNames
)

names_valid[,-c(2,3,4,9,11)]
```

	scientificName	scientificNameStatus	suggestedFamily
#> 1	Apis mellifera Linnaeus, 1758	name_w_authors	Apidae
#> 2	Apis melifera	possibly_ok	Apidae
#> 3	Apis cf. mellifera	conferre	Apidae
#> 4	Ancyloscelis armatus	possibly_ok	Apidae
#> 5	Centris aenea	possibly_ok	Apidae
#> 6	Centris rufa	possibly_ok	Apidae
#> 7	Centris Rhodoprocta Moure & Seabra, 1960	name_w_wrong_case name_w_authors	Apidae
#> 8	Lutjanus purpureus	possibly_ok	Lutjanidae
#> 9	Parotocinclus amazonensis	possibly_ok	Loricariidae
#> 10	Panthera onca	possibly_ok	Felidae
#> 11	Solenopsis bicolor (Emery, 1906)	name_w_authors	Formicidae
#> 12	Eucopricus columbi MacLeay, 1819	name_w_authors	Scarabaeidae
#> 13	Eucopricus sp.1	indet	Scarabaeidae
#>	suggestedAuthorship	tax.notes	scientificNameFull
#> 1	Linnaeus, 1758	name accepted	Apis mellifera Linnaeus, 1758
#> 2	Linnaeus, 1758	name misspelled	Apis mellifera Linnaeus, 1758
#> 3	Linnaeus, 1758	name accepted	Apis mellifera Linnaeus, 1758
#> 4	(Fabricius, 1793)	replaced synonym	Ancyloscelis apiformis (Fabricius, 1793)
#> 5	Lepeletier, 1841	name accepted	Centris aenea Lepeletier, 1841
#> 6	Lepeletier, 1841	replaced synonym	Centris aenea Lepeletier, 1841
#> 7	Moure & Seabra, 1960	name accepted	Centris rhodoprocta Moure & Seabra, 1960
#> 8	(Poey, 1860)	replaced synonym	Lutjanus campechanus (Poey, 1860)
#> 9	Garavello, 1977	synonym not replaced	Parotocinclus amazonensis Garavello, 1977
#> 10	(Linnaeus, 1758)	name accepted	Panthera onca (Linnaeus, 1758)
#> 11	(Emery, 1906)	name accepted	Solenopsis bicolor (Emery, 1906)
#> 12	(MacLeay, 1819)	replaced synonym	Sulcophanaeus columbi (MacLeay, 1819)
#> 13	d'Olsoufieff, 1924	replaced synonym	Sulcophanaeus d'Olsoufieff, 1924

Tip 1: for large name lists, consider altering the argument `split.letters`, `parallel`, and `cores`. The minimal fuzzy similarity is controlled by `sug.dist`.

Tip 2: The maximum distance in fuzzy matching (defaults to 10%) is controlled by the argument `sug.dist`.

4.1 Internal functions

`nameMatching()` is the internal function used for exact and fuzzy matching. Below, we demonstrate it using the same names (reference names are the “accepted/standardized” targets):

```

input_names <- c(
  "Apis mellifera Linnaeus, 1758",
  "Apis mellifica",
  "Ancyloscelis armatus",
  "Centris aenea",
  "Centris rufa",
  "Centris Rhodoprocta Moure & Seabra, 1960",
  "Lutjanus purpureus",
```

```

"Parotocinclus amazonensis",
"Panthera onca",
"Coelonertus baridioides Solari & Solari, 1906",
"Solenopsis bicolor (Emery, 1906)",
"Eucopricus columbi MacLeay, 1819"
)

ref_names <- c(
  "Apis mellifera Linnaeus, 1758",
  "Ancyloscelis apiformis (Fabricius, 1793)",
  "Centris aenea",
  "Centris rhodoprocta Moure & Seabra, 1960",
  "Lutjanus campechanus (Poey, 1860)",
  "Panthera onca",
  "Coelonertus baridioides Solari & Solari, 1906",
  "Solenopsis bicolor (Emery, 1906)",
  "Sulcophanaeus columbi (MacLeay, 1819"
)

nameMatching(input_names, ref_names)

```

```
#> [1] 1 1 2 3 3 4 5 NA 6 7 8 9
```

5 Validating family names using prepFamily()

`plantR` contains an internal dictionary of valid family names which can be used via the function `prepFamily()`. Currently, valid family names are available only for plants. But a similar procedure will be included for animals in the near future. So, for now, the function does not change the input family names.

```

names_valid <- prepFamily(names_valid,
                           fam.name = "suggestedFamily",
                           spp.name = "scientificName.new",
                           kingdom = "animalia")
#> Warning: Synonyms for the input kingdom are currently not available. Returning the input family na

```

6 Brief code summary

A compact two-step workflow (CTFB-only):

```

# 1) Standardize
names_fixed <- fixSpecies(names)

# 2) Validate against the CTFB backbone
utils::data("ctfbNames", package = "plantRdata")
names_valid <- prepSpecies(
  names_fixed,
  tax.names = c("scientificName.new", "scientificNameAuthorship.new"),
  db = ctfbNames

```

```

)
names_valid[, c("scientificName.new", "scientificNameFull", "tax.notes")]

#>      scientificName.new          scientificNameFull      tax.notes
#> 1       Apis mellifera        Apis mellifera Linnaeus, 1758 name accepted
#> 2       Apis melifera        Apis mellifera Linnaeus, 1758 name misspelled
#> 3       Apis mellifera        Apis mellifera Linnaeus, 1758 name accepted
#> 4 Ancyloscelis armatus    Ancyloscelis apiformis (Fabricius, 1793) replaced synonym
#> 5       Centris aenea        Centris aenea Lepeletier, 1841 name accepted
#> 6       Centris rufa         Centris aenea Lepeletier, 1841 replaced synonym
#> 7       Centris rhodoprocta Centris rhodoprocta Moure & Seabra, 1960 name accepted
#> 8       Lutjanus purpureus   Lutjanus campechanus (Poey, 1860) replaced synonym
#> 9 Parotocinclus amazonensis Parotocinclus amazonensis Garavello, 1977 synonym not replaced
#> 10      Panthera onca        Panthera onca (Linnaeus, 1758) name accepted
#> 11      Solenopsis bicolor   Solenopsis bicolor (Emery, 1906) name accepted
#> 12      Eucopricus columbi   Sulcophanaeus columbi (MacLeay, 1819) replaced synonym
#> 13      Eucopricus sp.1       Sulcophanaeus d'Olsoufieff, 1924 replaced synonym

```

Or, even simpler, using the wrapper `formatTax()`:

```

names_df <- data.frame(scientificName = names)
names_df_valid <- formatTax(names_df, db = ctfbNames)
#> The following family names were automatically replaced:
#>
#> |Genus           |Old fam.     |New fam.     |
#> |:-----|:-----|:-----|
#> |Ancyloscelis   |Apidae       |Araceae      |
#> |Apis            |Apidae       |Fabaceae     |
#> |Centris         |Apidae       |Melastomataceae |
#> |Lutjanus        |Lutjanidae   |Fabaceae     |
#> |Panthera        |Felidae      |Fabaceae     |
#> |Parotocinclus  |Loricariidae |Cactaceae    |
#> |Solenopsis      |Formicidae   |Campanulaceae |
#> |Sulcophanaeus   |Scarabaeidae |Elaeocarpaceae |

```

7 Citation

If you use **plantR**, please cite it as:

Lima, R.A.F., Sánchez-Tapia, A., Mortara, S.R., ter Steege, H., Siqueira, M.F. (2021).
plantR: An R package and workflow for managing species records from biological collections.
Methods in Ecology and Evolution 14(2): 332–339. <https://doi.org/10.1101/2021.04.06.437754>

And please also cite the taxonomic backbones that you used:

Boeger, W., & Valim, M. P. (2024). Brazilian Zoology Group 2023 (version 1.1) [Data set]. Zenodo. <https://doi.org/10.5281/zenodo.10498290>