

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

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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 possible  
#> 12 name_w_auth  
#> 13 name_w_auth
```

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"
#>          Apis cf. mellifera                     Ancyloscelis armatus
#>          "Apis cf. mellifera"                   "Ancyloscelis armatus"
#>          Centris aenea                          Centris rufa
#>          "Centris aenea"                       "Centris rufa"
#> Centris Rhodoprocta Moure & Seabra, 1960        Lutjanus purpureus
#> "Centris rhodoprocta Moure & Seabra, 1960"      "Lutjanus purpureus"
#>          Parotocinclus amazonensis            Panthera onca
#>          "Parotocinclus amazonensis"          "Panthera onca"
#>          Solenopsis bicolor (Emery, 1906)      Eucopricus columbi MacLeay, 1819
#>          "Solenopsis bicolor (Emery, 1906)"    "Eucopricus columbi MacLeay, 1819"
#>          Eucopricus sp.1                        "Eucopricus columbi MacLeay, 1819"
#>          "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
#> 3          Apis cf. mellifera      Apis cf.
#> 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, 1819
#> 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
#>	suggestedName suggestedAuthorship	tax.notes	
#> 1	Apis mellifera Linnaeus, 1758	name accepted	
#> 2	Apis mellifera Linnaeus, 1758	name misspelled	
#> 3	Apis mellifera Linnaeus, 1758	name accepted	
#> 4	Ancyloscelis apiformis (Fabricius, 1793)	replaced synonym	
#> 5	Centris aenea Lepeletier, 1841	name accepted	
#> 6	Centris aenea Lepeletier, 1841	replaced synonym	
#> 7	Centris rhodoprocta Moure & Seabra, 1960	name accepted	
#> 8	Lutjanus campechanus (Poey, 1860)	replaced synonym	
#> 9	Parotocinclus amazonensis Garavello, 1977	synonym not replaced	
#> 10	Panthera onca (Linnaeus, 1758)	name accepted	
#> 11	Solenopsis bicolor (Emery, 1906)	name accepted	
#> 12	Sulcophanaeus columbi (MacLeay, 1819)	replaced synonym	
#> 13	Sulcophanaeus d'Oloufieff, 1924	replaced synonym	
#>	scientificNameFull		
#> 1	Apis mellifera Linnaeus, 1758		
#> 2	Apis mellifera Linnaeus, 1758		
#> 3	Apis mellifera Linnaeus, 1758		
#> 4	Ancyloscelis apiformis (Fabricius, 1793)		
#> 5	Centris aenea Lepeletier, 1841		
#> 6	Centris aenea Lepeletier, 1841		
#> 7	Centris rhodoprocta Moure & Seabra, 1960		
#> 8	Lutjanus campechanus (Poey, 1860)		
#> 9	Parotocinclus amazonensis Garavello, 1977		
#> 10	Panthera onca (Linnaeus, 1758)		
#> 11	Solenopsis bicolor (Emery, 1906)		
#> 12	Sulcophanaeus columbi (MacLeay, 1819)		
#> 13	Sulcophanaeus d'Oloufieff, 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 apiformis",
  "Centris aenea",
  "Lutjanus purpureus",
  "Parotocinclus amazonensis",
  "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 5 NA 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",
                           db = ctfbNames)
#> 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,
                             kingdom = "animalia")
#> Warning: Synonyms for the input kingdom are currently not available. Returning the input family na
```

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>