

Package: standrecon (via r-universe)

May 14, 2026

Title Reconstruct Historical Forest Stand Conditions

Version 0.1.0.9000

Description Reconstructs forest stand basal area and stem density at user-specified reference years using tree-level inventory data. The method estimates missing tree ages, back-calculates diameters using species-specific growth rates, and incorporates decay-class-based decomposition to infer mortality timing for dead trees. Results are returned in a tidy long format suitable for analysis and visualization.

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Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.3.3

URL <https://github.com/kriggithub/standrecon>

BugReports <https://github.com/kriggithub/standrecon/issues>

Suggests testthat (>= 3.0.0)

Config/testthat/edition 3

Imports stats, utils

Depends R (>= 3.5)

LazyData true

Repository <https://kriggithub.r-universe.dev>

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RemoteUrl <https://github.com/kriggithub/standrecon>

RemoteRef HEAD

RemoteSha 9132a6d606bdeb808e4062f9c5153024906c48af

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standrecon	<i>Reconstruct stand conditions from a given reference year</i>
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Description

Reconstruct stand conditions from a given reference year

Usage

```
standrecon(
  data,
  meas_year,
  ref_year,
  avg_inc_vec,
  plot_size,
  n_plots = 1,
  bark_eq_list = list(),
  species_col = "Species",
  age_col = "Age",
  dbh_col = "DBH",
  status_col = "Status",
  decay_col = "Decay",
  percentiles = c(0.25, 0.5, 0.75),
  min_dbh = 5
)
```

Arguments

data	Data with species, age, DBH, tree status, and tree decay columns.
meas_year	Year that data was measured.
ref_year	Reference years to reconstruct stand conditions.
avg_inc_vec	Vector of average increment growth per species with units mm/year. (PIEN = 0.5, ABBI = 0.3)
plot_size	Size of plots in m ² .
n_plots	Number of plots with plot_size.
bark_eq_list	List of bark correction equations. Default bark corrections for species codes used otherwise. list(PIPO = function(x) x*1.1029 + 0.7162)
species_col	Species column name in data as a string.
age_col	Species column name in data as a string.
dbh_col	DBH column name in data as a string.
status_col	Species status column name in data as a string.
decay_col	Species decay column name in data as a string.
percentiles	Percentiles to be calculated for sensitivity analysis.
min_dbh	Minimum DBH threshold for stand calculations in cm.

Value

A data frame with one row per species, reference year, and percentile. Columns include output type (reconstructed or measured), reference year, percentile, species code, basal area per hectare, and stem density per hectare.

Examples

```
data(standrecon_example_data)

out <- standrecon(
  data = standrecon_example_data,
  meas_year = 2025,
  ref_year = c(1950, 1975),
  avg_inc_vec = c(PIEN = 0.5, ABBI = 0.3, PIPO = 0.4),
  plot_size = 1000)

out
```

standrecon_example_data

Example field collected tree data for stand reconstruction

Description

A small synthetic data set for examples and tests.

Usage

```
standrecon_example_data
```

Format

A data frame with 200 rows and 5 variables:

Species Species code.

Age Tree age in years. Values that are known typically come from cored trees.

Status Tree status code.

Decay Decay class for dead trees.

DBH Diameter at breast height in cm.

Source

Simulated data set.

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