

# Package ‘standrecon’

April 8, 2026

**Title** Reconstruct Historical Forest Stand Conditions

**Version** 0.1.0

**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.

**License** MIT + file LICENSE

**Encoding** UTF-8

**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

**NeedsCompilation** no

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**Repository** CRAN

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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 <sup>2</sup> .
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
```

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standrecon\_example\_data

*Example field collected tree data for stand reconstruction*

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**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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