

# Package ‘AgeBandDecomposition’

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**Title** Age Band Decomposition Method for Tree Ring Standardization

**Version** 2.0.0

**Description** Implements the Age Band Decomposition (ABD) method for standardizing tree ring width data while preserving both low and high frequency variability. Unlike traditional detrending approaches that can distort long term growth trends, ABD decomposes ring width series into multiple age classes, detrends each class separately, and then recombines them to create standardized chronologies. This approach improves the detection of growth signals linked to past climatic and environmental factors, making it particularly valuable for dendroecological and dendroclimatological studies. The package provides functions to perform ABD-based standardization, compare results with other common methods (e.g., BAI, C method, RCS), and facilitate the interpretation of growth patterns under current and future climate variability.

**URL** [https://gitlab.com/Puletti/agebanddecomposition\\_rpackage](https://gitlab.com/Puletti/agebanddecomposition_rpackage)

**Language** en-US

**License** GPL (>= 3)

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ABD	<i>ABD - Age Band Decomposition</i>
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Description

ABD - Age Band Decomposition. This function calculates standardized tree-ring width chronologies by decomposing tree-ring width (stdTRW) data into age bands, detrending each age band separately, and then recombining them to produce the mean standardized chronology. Specifically, ABD standardizes each series within its age band (e.g., each ring width is divided by the corresponding age-specific expected value from the stdTRW). Then, standardized values from all age bands are merged and averaged to produce a composite chronology, preserving both inter-annual and low-frequency climate signals.

Usage

```
ABD(inTRW, min_nTrees_year = 3, pct_stdTRW_th = 0.5, pct_Trees_th = 0.3)
```

Arguments

- inTRW                    tibble. A tibble. The input dataset obtained from the import\_rwl() function.
- min\_nTrees\_year        Numeric. Sets the minimum number of trees per year required within each age class to be included in the analysis. The default is 3. Using less than three trees may result in poor representation of within-class variability and is not recommended unless data availability is limited.
- pct\_stdTRW\_th        Numeric. Sets the threshold for the minimum proportion of standardized tree-ring width values required within a given age class. The default is set to 0.5, meaning that at least 50% of the possible values must be present (e.g., 6 out of 10 values for 10-year age bands, or 11 out of 20 for 20-year bands). For instance, the final age band of a 94-year-old tree (i.e., the 91–100-year band) includes only 4 years of growth. Since this number of years below the threshold, we recommend excluding that tree from the corresponding age band group.

`pct_Trees_th` Numeric. Sets the minimum number of samples (i.e., trees) required to compute the mean standardized tree-ring widths within each age band. The default value is 0.3. However, when working with small sample sizes (approximately 20 trees or fewer), increasing the threshold to 0.5 is recommended to ensure that more trees are included in the analysis.

## Details

The function performs age-band decomposition on `stdTRW` data by filtering out age classes with insufficient observations (`min_nTrees_year`) or excessive within-class variation.

The `stdTRW_th` value is particularly important in small datasets: too strict a threshold may exclude valid data. If working with approximately 20 trees or fewer, a threshold of 0.5 is suggested.

Please, see the Examples section below for a demonstration using a typical input tibble obtained with `import_rwl()`.

## Value

A tibble with the following columns: `year`, `N_ageBands`, `ABD`, and `ABDsd`.

## See Also

[import\\_rwl](#), [stdTRW](#), [plotABD](#)

Other ABD functions: [import\\_rwl\(\)](#), [stdTRW\(\)](#)

## Examples

```
ABD(inTRW)
```

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<code>import_rwl</code>	<i>import_rwl</i>
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## Description

This function imports tree-ring width data from RWL (Tucson format) files and pith offset information, then arranges the dataset for ABD analysis.

## Usage

```
import_rwl(
  rwl_path,
  po_path,
  ageBands,
  first_age_class = NULL,
  zero_as_na = TRUE,
  verbose = TRUE
)
```

## Arguments

<code>rwl_path</code>	Path to the RWL file (Tucson format).
<code>po_path</code>	Path to the pith offset file (tab-delimited text file with columns 'tree_code' and 'pith.offset').
<code>ageBands</code>	character. Setting the age band window. It must be set to '1010' if all the age classes have the same size (10 years). It must be '1020' if the age classes have different sizes: 10 years till 100 and then 20 years size.
<code>first_age_class</code>	numeric. is numeric and specifies the first age band from which the analysis begins. If NULL (default), no filtering is applied. For example, <code>first_age_class = 3</code> excludes the first 20 years of growth (age classes 1 and 2 in '1010' mode).
<code>zero_as_na</code>	logical. If TRUE (default), zero values in TRW are converted to NA. If FALSE, zero values are kept as is.
<code>verbose</code>	logical. If TRUE, prints additional information during import.

## Details

The RWL file must be in Tucson format (readable by `dplR::read.rwl`). The pith offset file must be a tab-delimited text file with header containing columns 'tree\_code' and 'pith.offset', where pith.offset indicates the number of missing rings between the pith and the first measured ring. are available on the package's GitLab page. [https://gitlab.com/Puletti/agebanddecomposition\\_rpackage](https://gitlab.com/Puletti/agebanddecomposition_rpackage) and can be used to test the package's functions.

## Value

A list of two objects. The first object is a tibble representing the imported dataset in long format. In this tibble the last two columns are an identification number (`id_by_years`) and two grouping variables (`age_class` and `ageBands`). The second object in the list is a lookup table (tibble), useful for further steps.

## See Also

[stdTRW](#), [ABD](#)

Other ABD functions: [ABD\(\)](#), [stdTRW\(\)](#)

## Examples

```
# Download example files from the package's GitLab page
package_gitlab_site <- 'https://gitlab.com/Puletti/agebanddecomposition_rpackage'
rwl_url <- "/-/raw/main/studio/dati/TRW_example.rwl"
po_url <- "/-/raw/main/studio/dati/pith.offset.txt"

# Create temporary files
tmpfile_rwl <- tempfile(fileext = ".rwl")
tmpfile_po <- tempfile(fileext = ".txt")

# Download the files
download.file(paste0(package_gitlab_site, rwl_url),
```

```

      tmpfile_rwl,
      mode = "wb")

download.file(paste0(package_gitlab_site, po_url),
             tmpfile_po,
             mode = "wb")

# Import the data
inData <- import_rwl(rwl_path = tmpfile_rwl,
                    po_path = tmpfile_po,
                    ageBands = '1010',
                    first_age_class = 3,
                    zero_as_na = TRUE,
                    verbose = TRUE)

# View the result
inData

```

---

inTRW	<i>AgeBandDecomposition test data</i>
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---

### Description

A tiny test dataset including example, the basic reference for AgeBandDecomposition package.

A list of two test datasets derived from TRW\_readExcel function. The first list-object is a tibble that can be used as the argument of stdTRW function.

### Usage

```
inTRW
```

```
inTRW
```

### Format

A tibble with 85 rows and 16 variables, as derived from the Excel file.

A list of two tibbles

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plotABD	<i>plotABD</i>
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---

### Description

Function for plotting ABD values from inTRW object using the two tibble objects produced by the function import\_rwl.

**Usage**

```
plotABD(
  inTRW,
  min_nTrees_year = 3,
  pct_stdTRW_th = 0.5,
  pct_Trees_th = 0.3,
  linewidth_TRW = 0.7,
  linewidth_Ntrees = 1,
  byYears = 20,
  xlim = NULL,
  ylimABD = NULL,
  ...
)
```

**Arguments**

<code>inTRW</code>	tibble. A tibble. The input dataset obtained from the <code>import_rwl()</code> function.
<code>min_nTrees_year</code>	Numeric. The minimum number of trees per year required within each age class to be included in the analysis. The default is 3. Using less than three trees may result in poor representation of within-class variability and is not recommended unless data availability is limited.
<code>pct_stdTRW_th</code>	Numeric. It defines the threshold for the minimum proportion of standardized tree-ring width values required within a given age class. The default is set to 0.5, meaning that at least 50% of the possible values must be present (e.g., 6 out of 10 values for 10-year age bands, or 11 out of 20 for 20-year bands). For instance, the final age band of a 94-year-old tree (i.e., the 91–100-year band) includes only 4 years of growth. Since this number of years below the threshold, we recommend excluding that tree from the corresponding age band group.
<code>pct_Trees_th</code>	Numeric. It defines the threshold used to calculate the mean standardized tree-ring widths within each age band. The default value is 0.3. However, when working with small sample sizes (approximately 20 trees or fewer), it is advisable to increase the threshold to 0.5. This adjustment helps retain more trees in the analysis while still accounting for natural growth variability.
<code>linewidth_TRW</code>	numeric. line size for TRW.
<code>linewidth_Ntrees</code>	numeric. line size for Ntrees.
<code>byYears</code>	numeric. Spacing (in years) between tick marks on the x-axis of time-related plots.
<code>xlim</code>	Optional numeric vector of length 2 specifying the limits of the x-axis. If NULL the limits are automatically determined.
<code>ylimABD</code>	Optional numeric vector of length 2 specifying the limits of the y-axis for the ABD panel. If NULL the limits are automatically determined.
<code>...</code>	Other arguments passed on to methods. Not currently used.

**Details**

This function produces a multi-panel plot with three graphs: the first displays standardized values by age band; the second shows the final mean chronology (ABD), adjusted for age-related effects; and the third depicts the corresponding number of trees.

**Value**

None. A multipanel plot.

**See Also**

[plotBAI](#), [plotTRW](#), [ABD](#)

Other tree ring plotting: [plotBAI\(\)](#), [plotTRW\(\)](#)

**Examples**

```
plotABD(inTRW,
  min_nTrees_year = 3,
  pct_stdTRW_th = .5,
  pct_Trees_th = .3,
  byYears = 50)
```

---

plotBAI

---

*plotBAI*


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**Description**

Function for plotting basal area increment (BAI) values derived from `inTRW` object using the first tibble produced by the function `import_t_rwl`.

**Usage**

```
plotBAI(
  inTRW,
  linewidth_BAI = 1,
  linewidth_Ntrees = 1,
  byYears = 20,
  xlim = NULL,
  ylim = NULL,
  ...
)
```

**Arguments**

<code>inTRW</code>	tibble. The first object resulting from the <code>import_rwl</code> function.
<code>linewidth_BAI</code>	numeric. line size for BAI.
<code>linewidth_Ntrees</code>	numeric. line size for Ntrees.
<code>byYears</code>	numeric. Spacing (in years) between tick marks on the x-axis of time-related plots.
<code>xlim</code>	Optional numeric vector of length 2 specifying the limits of the x-axis. If NULL the limits are automatically determined.
<code>ylim</code>	Optional numeric vector of length 2 specifying the limits of the y-axis for the ABD panel. If NULL the limits are automatically determined.
<code>...</code>	Other arguments passed on to methods. Not currently used.

**Details**

This function generates a basic plot displaying the mean chronology +/- standard error of raw basal area increments (in squared cm) alongside the corresponding number of trees.

**Value**

None. A plot is produced.

**See Also**

[plotTRW](#), [plotABD](#)

Other tree ring plotting: [plotABD\(\)](#), [plotTRW\(\)](#)

**Examples**

```
plotBAI(inTRW
, linewidth_BAI = .5
, linewidth_Ntrees = 1
)
```

---

plotTRW

*plotTRW*


---

**Description**

Function for plotting TRW values from `inTRW` object using the first tibble produced by the function `import_rwl`.



**Usage**

```
plotTRW(
  inTRW,
  linewidth_TRW = 1,
  linewidth_Ntrees = 1,
  byYears = 20,
  xlim = NULL,
  ylim = NULL,
  ...
)
```

**Arguments**

<code>inTRW</code>	tibble. The first object resulting from the <code>import_rwl</code> function.
<code>linewidth_TRW</code>	numeric. line size for TRW.
<code>linewidth_Ntrees</code>	numeric. line size for Ntrees.
<code>byYears</code>	numeric. Spacing (in years) between tick marks on the x-axis of time-related plots.
<code>xlim</code>	Optional numeric vector of length 2 specifying the limits of the x-axis. If NULL the limits are automatically determined.
<code>ylim</code>	Optional numeric vector of length 2 specifying the limits of the y-axis for the ABD panel. If NULL the limits are automatically determined.
<code>...</code>	Other arguments passed on to methods. Not currently used.

**Details**

This function generates a basic plot displaying the mean chronology +/- standard error of raw tree-ring widths (in mm) alongside the corresponding number of trees.

**Value**

None. A plot is produced.

**See Also**

[plotBAI](#), [plotABD](#)

Other tree ring plotting: [plotABD\(\)](#), [plotBAI\(\)](#)

**Examples**

```
plotTRW(inTRW)
```

stdTRW

*stdTRW***Description**

To remove the influences of local site characteristics on tree growth, this function standardizes the tree-ring width series by dividing each tree-ring width of a particular series by the mean width of that series. In dendroclimatic studies, it is advisable to exclude the initial years or decades of growth (the first “age bands”, i.e., 1-10 and 11-20) as they are less likely to contain climatic signals rather than the influence of strong early-year growth competition within the tree stand (Mazza and Sarris, 2021; Sarris et al., 2007).

**Usage**

```
stdTRW(inTRW)
```

**Arguments**

**inTRW**                      tibble. The first object resulting from the `import_rwl` function.

**Details**

To export a tibble like the one below to an `.xlsx` file, you can use the `writexl` package:

```
mytibble <- stdTRW_df |>
  dplyr::group_by(year) |>
  dplyr::summarise(
    N_trees = dplyr::n(),
    mean_stdTRW = mean(stdTRW, na.rm = TRUE)
  )

writexl::write_xlsx(mytibble, path = "my_stdTRW_df.xlsx")
```

This will create an Excel file named `my_stdTRW_df.xlsx` in your current working directory. Please, check also the **Examples** section below for how to obtain `stdTRW_df` object.

**Value**

A tibble with the following columns: `year`, `tree_code`, `TRW`, `meanTRW`, and `stdTRW`. Together with additional columns of grouping variables based on age class.

**References**

Mazza, G., Sarris, D., 2021. Identifying the full spectrum of climatic signals controlling a tree species' growth and adaptation to climate change. *Ecol. Indic.* 130, 108109. <https://doi.org/10.1016/j.ecolind.2021.108109>.

Sarris, D., Christodoulakis, D., Körner, C., 2007. Recent decline in precipitation and tree growth in the eastern Mediterranean. *Glob. Chang. Biol.* 13 (6), 1187–1200. <https://doi.org/10.1111/j.1365-2486.2007.01348.x>.

**See Also**[import\\_rwl](#), [ABD](#)Other ABD functions: [ABD\(\)](#), [import\\_rwl\(\)](#)**Examples**

```

inTRW_1 <- inTRW[[1]]

stdTRW_df <- stdTRW(inTRW_1)

stdTRW_df |>
  dplyr::group_by(year) |>
  dplyr::summarise(N_trees = dplyr::n(),
    mean_stdTRW = mean(stdTRW, na.rm = TRUE)) |>
  ggplot2::ggplot(ggplot2::aes(year, N_trees)) +
  ggplot2::geom_line()

```

TRW\_readExcel

*TRW\_readExcel (Deprecated)***Description**

**This function is deprecated and will be removed in a future version.** Please use [import\\_rwl](#) instead after converting your data to RWL format.

**Usage**

```
TRW_readExcel(path, sheet, ageBands, limitFirst20y = FALSE, verbose = TRUE)
```

**Arguments**

path	Path to the xls/xlsx file.
sheet	Sheet to read.
ageBands	character. Age band window setting.
limitFirst20y	logical. Remove first 20 years from each tree.
verbose	logical. Print additional information.

**Value**

A list of two objects (see original documentation).

**See Also**[import\\_rwl](#)

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