

# Package: ggbenjamini (via r-universe)

October 15, 2024

**Title** Generate ficus benjamina leaf shapes with bezier curves

**Version** 0.0.1

**Description** This package can be used to generate shapes in the form of ficus benjamina (weeping fig) leaves with bezier curves. The main output of the package are dataframes containing all the information of these bezier curves.

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**Suggests** flametree, rsvg, minisvg, knitr, rmarkdown, testthat (>= 3.0.0), vdiffr, ambient, covr, usethis

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**Remotes** coolbutuseless/minisvg, djnavarro/flametree

**Depends** R (>= 2.10)

**URL** <https://urswilke.github.io/ggbenjamini/>,

<https://github.com/urswilke/ggbenjamini/>

**Repository** <https://urswilke.r-universe.dev>

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**benjamini\_branch**      *Generate a branch of benjamini leaves*

### Description

Generate a branch of benjamini leaves

### Usage

```
benjamini_branch(
  df_branch = tibble::tibble(x = c(70, 84, 126, 168), y = c(280, 245, 217, 217)),
  leaf_mean_dist_approx = 10,
  leaf_angle = 45,
  first_dir = sample(0:1, 1),
  leave_size_dist = spark_weibull(shape = 1.5, scale_factor = 0.8),
  leaf_angle_dist = spark_norm(mean = 0, sd = 3),
  last_angle_straight = TRUE,
  leaf_size_multipliator = 1
)
```

### Arguments

df_branch	dataframe containing the 4 x & y bezier coordinates of a branch
leaf_mean_dist_approx	approximate distance of two leaves
leaf_angle	angle between the leaf stalks and the branch
first_dir	direction of the first leaf in the branch (0 for right; 1 for left)
leave_size_dist	Manipulate the sizes of the leaves with a spark function This function returns a function which itself returns a numerical vector of length of the number of leaves on the branch. The function will be rescaled (divided by it's maximum value).

```
leaf_angle_dist
    Manipulate the leaf angles with a spark function This function returns a function
    which itself returns a numerical vector of length of the number of leaves on the
    branch.
last_angle_straight
    Logical if the angle of the last leaf on the branch is very sharp (defaults to
    TRUE).
leaf_size_multiplicator
    Multiply leaf size distribution with a factor.
```

**Value**

A dataframe containing the data for leaves on a branch (see example).

**Examples**

```
benjamini_branch() %>%
  tidyr::unite(b, i_part, i_leaf, element, remove = FALSE) %>%
  ggplot2::ggplot() +
  ggforce::geom_bezier(ggplot2::aes(x = x, y = y, group = b)) +
  ggplot2::coord_equal()
```

**benjamini\_leaf**

*Generate bezier curve coordinates of a benjamini leaf*

**Description**

Generate bezier curve coordinates of a benjamini leaf

**Usage**

```
benjamini_leaf(
  leaf_params = gen_leaf_parameters(),
  omega = 0,
  xrot = leaf_params$x0,
  yrot = leaf_params$y0,
  precision = 2
)
```

**Arguments**

<b>leaf_params</b>	parameter that control the leaf shape
<b>omega</b>	rotation angle of the leaf
<b>xrot</b>	x coordinate of pivot point (preset to leaf origin).
<b>yrot</b>	x coordinate of pivot point (preset to leaf origin).
<b>precision</b>	numeric precision of the output

**Value**

A dataframe containing the data for the bezier curves of a leaf (see example).

**Examples**

```
df <- benjamini_leaf()
df
df %>%
  # This generated a unique identifier for the 4 rows of each bezier curve:
  tidyr::unite(b, element, i_part, remove = FALSE) %>%
  ggplot2::ggplot() +
  ggforce::geom_bezier(ggplot2::aes(x = x, y = y, group = b))
```

**bezier\_to\_polygon**

*Transform bezier dataframe to dataframe with path coordinates*

**Description**

Transform bezier dataframe to dataframe with path coordinates

**Usage**

```
bezier_to_polygon(df_benjamini_leaf, ..., n = 100)
```

**Arguments**

df_benjamini_leaf	Dataframe returned by benjamini_leaf()
...	grouping variables in df_benjamini_leaf that will be kept in the transformation.
n	number of points per bezier

**Value**

Dataframe with the coordinates of the bezier curve interpolations.

**Examples**

```
df_coords <- benjamini_leaf() %>%
  tidyr::unite(b, i_part, element, remove = FALSE) %>%
  bezier_to_polygon(b, i_part, element)
df_coords
df_coords %>%
  ggplot2::ggplot(ggplot2::aes(x = x, y = y, group = element, fill = element)) +
  ggplot2::geom_polygon()
```

---

df\_bezier\_skeleton     *Example bezier dataframe to grow leaves on*

---

### Description

Generated with `vignette("import_svg_bezier")`

### Usage

```
df_bezier_skeleton
```

### Format

A data frame with 40 rows and 4 variables:

**i\_branch** branch index

**svg\_point\_type** original point type in the svg file

**x** x coordinate

**y** y coordinate

---

gen\_benjamini\_points     *Generate bezier end points*

---

### Description

Generate bezier end points

### Usage

```
gen_benjamini_points(leaf_params = gen_leaf_parameters())
```

### Arguments

**leaf\_params**     parameters generated by `gen_leaf_parameters()`

### Value

tibble with absolute coordinates

### Examples

```
gen_benjamini_points()
```

`gen_benjamini_slopes`    *Generate bezier slopes coordinates*

### Description

Generate bezier slopes coordinates

### Usage

```
gen_benjamini_slopes(leaf_params = gen_leaf_parameters())
```

### Arguments

`leaf_params`    parameters generated by `gen_leaf_parameters()`

### Value

dataframe with sx1-4 in the x variable and sy1-4 in the y variable

### Examples

```
gen_benjamini_slopes()
```

`gen_leaf_parameters`    *Generate bezier coordinates of a leaf*

### Description

Except the start point coordinates of the leaf origin  $x_0$  &  $y_0$  (arbitrarily set to 10 and 40) all coordinates are relative to this origin. Most of the parameters are random generated, except for some of them (e.g., to close polygons, or to keep the stalk and the mid vein on the same line).

### Usage

```
gen_leaf_parameters(
  x0 = 10,
  y0 = 40,
  dx10 = sample(8:12, 1),
  dy10 = 0,
  dx21 = sample(12:20, 1),
  dy21 = sample(-4:-10, 1),
  dx32 = sample(10:18, 1),
  dy32 = stats::runif(1, 0.92 * (-dy21 - 1), 0.95 * (-dy21 - 1)),
  dx43 = sample(4:6, 1),
  dy43 = y0 + dy10 + dy21 + dy32,
  sx0 = stats::runif(1, 0, 0.1),
```

```

sx1 = sample(1:3, 1),
sx2 = sample(4:6, 1),
sx3 = sample(2:4, 1),
sx4 = stats::runif(1, 0, 0.2),
sy0 = stats::runif(1, 0.5, 1),
sy1 = sample(-4:-6, 1),
sy2 = stats::runif(1, -0.5, 0.5),
sy3 = stats::runif(1, 0.5, 1.5),
sy4 = stats::runif(1, 0.2, 0.7),
smx1 = sample(-5:-15, 1),
smx2 = sample(-5:-15, 1),
smy1 = stats::runif(1, -1, 1),
smy2 = stats::runif(1, -1, 1)
)

```

**Arguments**

`x0, y0` coordinates of the leaf origin  
`dx10, dy10, dx21, dy21, dx32, dy32, dx43, dy43`  
coordinates of the other bezier start & end points  
`sx0, sx1, sx2, sx3, sx4`  
x coordinates of the control points  
`sy0, sy1, sy2, sy3, sy4`  
y coordinates of the control points  
`smx1, smx2, smy1, smy2`  
x & y coordinates of the mid vein control points

**Value**

named list of all parameters

**Examples**

```
gen_leaf_parameters()
```

`gen_middle_line_slopes`

*Generate bezier slopes of the line in the middle of the leaf*

**Description**

Generate bezier slopes of the line in the middle of the leaf

**Usage**

```
gen_middle_line_slopes(leaf_params = gen_leaf_parameters())
```

**Arguments**

`leaf_params` parameters generated by `gen_leaf_parameters()`

**Value**

A dataframe containing the coordinates of the two control points of the bezier curve defining the midvein of the leaf.

**Examples**

`gen_middle_line_slopes()`

`get_one_bezier` *Generate a dataframe of one bezier curve*

**Description**

Generate a dataframe of one bezier curve

**Usage**

`get_one_bezier(i, points_df, slopes_df)`

**Arguments**

`i` number of the bezier  
`points_df` dataframe generated by `gen_benjamini_points()` (see example).  
`slopes_df` dataframe generated by `gen_benjamini_slopes()` (see example).

**Value**

A dataframe containing the information for one bezier curve in the format as needed by `ggforce::geom_bezier`.

**Examples**

```
set.seed(123)
leaf_params <- gen_leaf_parameters()
points_df <- gen_benjamini_points()
slopes_df <- gen_benjamini_slopes()
df_bezier <- get_one_bezier(1, points_df, slopes_df)
ggplot2::ggplot(df_bezier, ggplot2::aes(x = x, y = y)) + ggforce::geom_bezier()
```

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**spark\_norm***Manipulate the angles of the leaves with a normal distribution*

---

## Description

This function returns a function which itself returns a numerical vector of length of the number of leaves on the branch.

## Usage

```
spark_norm(mean = 0, sd = 3)
```

## Arguments

`mean, sd` Parameters passed to `stats::dnorm()`.

## Value

`dnorm()` function with `n_leaves` as one of the arguments

## Examples

```
spark_norm()
```

---

**spark\_weibull***Manipulate the sizes of the leaves with a Weibull distribution*

---

## Description

This function returns a function which itself returns a numerical vector of length `n_leaves`, the number of leaves on the branch. These values serve as relative multiplication factors of the sizes of the leaves of the branch. (The maximum value of this distribution is then normalized to 1.)

## Usage

```
spark_weibull(shape = 1.2, scale_factor = 0.5)
```

## Arguments

`shape, scale_factor`  
Parameters passed to `stats::dweibull()`.

## Value

`dweibull()` function depending on `n_leaves` as one of the arguments

**Examples**

```
#Mark the two consecutive pairs of parentheses:  
spark_weibull()(n_leaves = 10)
```

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