

# Package: latbias (via r-universe)

May 19, 2026

**Type** Package

**Title** Calculate the Latitudinal Bias Index

**Version** 1.0.0

**Description** Studies that report shifts in species distributions may be biased by the shape of the study area. The main functionality of this package is to calculate the Latitudinal Bias Index (LBI) for any given shape. The LBI is bounded between +1 (100% probability to exclusively record latitudinal shifts, i.e., range shifts data sampled along a perfectly South-North oriented straight line) and -1 (100% probability to exclusively record longitudinal shifts, i.e., range shifts data sampled along a perfectly East-West oriented straight line).

**Depends** R (>= 3.5.3)

**License** GPL (>= 2)

**Encoding** UTF-8

**LazyData** true

**Imports** dplyr, geosphere, ggplot2, psych, RColorBrewer, reshape2, sf, sp, terra, tidyr, units

**RoxygenNote** 7.3.3

**Suggests** elevatr, knitr, progress, rmarkdown, rnaturalearth, rnaturalearthdata, testthat (>= 3.0.0)

**Config/testthat/edition** 3

**URL** <https://github.com/pierredenelle/latbias>,

**BugReports** <https://github.com/pierredenelle/latbias/issues>

**VignetteBuilder** knitr

**Roxygen** list(markdown = TRUE)

**Config/pak/sysreqs** libabsl-dev cmake libgdal-dev gdal-bin libgeos-dev libicu-dev libssl-dev libproj-dev libsqlite3-dev libudunits2-dev

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

**Date/Publication** 2025-11-03 11:37:16 UTC

**RemoteUrl** <https://github.com/pierredenelle/latbias>

**RemoteRef** HEAD

**RemoteSha** 8ed1af48ecabba1533b2c1928aa642040fd3e639

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LBI	<i>LBI</i>
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## Description

Computes the Latitudinal Bias Index (LBI) for a given shapefile, by calculating the distance between two random locations within the shape multiple times (see details).

## Usage

```
LBI(
  study_area_id,
  study_area_polygon,
  nobs = 250,
  nboot = 1000,
  fact_location = 10,
  elevation = NULL,
  raw_output = FALSE
)
```

## Arguments

<code>study_area_id</code>	Character string. Name ID of the study case area or country name.
<code>study_area_polygon</code>	Polygon shapefile. It should be a <code>sfc</code> object, of class <code>POLYGON</code> or <code>MULTI-POLYGON</code> .
<code>nobs</code>	Numeric. Number of random observations in each sample. 250 by default.
<code>nboot</code>	Numeric. Determines how many times the random shifts are calculated. 1,000 by default.
<code>fact_location</code>	Numeric. <code>fact_location</code> x <code>nobs</code> determine all the possible coordinates that can be sampled within the provided polygon for all bootstraps (faster than to generate a set of random location at each bootstrap)
<code>elevation</code>	Elevation raster. <code>elevation</code> in <code>wgs84</code> ; if not provided, <code>NA</code> will be returned for null-model elevational shifts.
<code>raw_output</code>	Logical. <code>FALSE</code> by default. If <code>TRUE</code> , all bootstraps are returned as a <code>data.frame</code> . => say that the raw outputs are accessible

## Details

The main output contains the following columns:

- `study_area_id`: ID or name of the study case region
- `distance_km`: average expected geographic distance shift between t1 and t2
- `null_mod_SN_shift`: average expected South-North shift between t1 and t2, in absolute values
- `null_mod_EW_shift`: average expected East-West shift between t1 and t2, in absolute values
- `null_mod_elevation_shift`: average expected elevation shift between t1 and t2, in absolute values
- `LBI`: the Latitudinal Bias Index value

LBI formula is

$$LBI = 2 \times \left( \frac{\text{mean}(|\frac{A_n \text{lat}}{A_n \text{lon}}|)}{1 + \text{mean}(|\frac{A_n \text{lat}}{A_n \text{lon}}|)} - 0.5 \right)$$

with  $A_n \text{lat}$  and  $A_n \text{lon}$  denoting the geographic displacement of the centroid positions of both sets of observation in the  $n$ th iteration by means in the latitudinal and the longitudinal dimension.

## Value

A `data.frame` or a list of two `data.frames` if `raw_output` is set to `TRUE`.

## References

Sanczuk et al. submitted.

## Examples

```
study_area <- rnatuarearth::ne_countries(scale = 110, continent = "Europe",
country = "Sweden", type = "map_units", returnclass = "sf")
study_area <- sf::st_union(study_area)
```

```
LBI(study_area_id = "Sweden", study_area_polygon = study_area,
nobs = 10, nboot = 10, fact_location = 5, elevation = NULL)
```

```
# With elevation
elevation_df <- elevatr::get_elev_raster(
locations = sf::st_as_sf(study_area), z = 5)
LBI(study_area_id = "Sweden", study_area_polygon = study_area,
nobs = 10, nboot = 10, fact_location = 5, elevation = elevation_df)
```

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windrose	<i>windrose</i>
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### Description

Plot the directional shifts for a given bootstrap.

### Usage

```
windrose(
  data,
  spd,
  dir,
  spdres = 30,
  dirres = 30,
  spdmin = 0,
  spdmax = 150,
  spdseq = NULL,
  palette = "Spectral",
  countmax = NA,
  debug = 0
)
```

### Arguments

data	ddata frame with two columns including the bearing and distance or speed of shifts
spd	distance or speed column
dir	bearing column
spdres	30 by default. Numeric; resolution for plotting distance of speed categories; adapt here the legend and breaks.
dirres	30 by default. Numeric, resolution for plotting bearings on windrose; adapt here the number of bars plotted on the windrose.
spdmin	0 by default. Numeric, minimal distance/speed for plotting.
spdmax	150 by default. Numeric, maximal distance/speed for plotting.
spdseq	NULL
palette	Character string defining the color palette to be used. Available options are taken from <code>rownames(RColorBrewer::brewer.pal.info)</code> .
countmax	NA by default. Numeric, optional, adjust the y-axis limit (maximum)
debug	0 by default. Numeric, if >0, run debug to find optimal bins of spd and dir.

### Details

This is intended to plot raw simulation output (that is, bearing and distance of random shifts) on a windrose.

**Value**

Windrose plot, ggplot object.

**References**

Sanczuk et al. submitted.

**Examples**

```
study_area <- rnaturalearth::ne_countries(scale = 110, continent = "Europe",
country = "Sweden", type = "map_units", returnclass = "sf")
study_area <- sf::st_union(study_area)
po <- LBI(study_area_id = "Sweden", study_area_polygon = study_area,
nobs = 10, nboot = 10, fact_location = 5, elevation = NULL,
raw_output = TRUE)
test <- as.data.frame(po$all[, c("study_area_id", "distance_km",
"bearing", "rep")])
test$distance_km <- as.numeric(test$distance_km)
pop <- windrose(data = test, spd = "distance_km", dir = "bearing")
```

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