

# Package ‘StablePopulation’

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**Title** Calculates Alpha for a Stable Population

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**Description** Provides tools to calculate the alpha parameter of the Weibull distribution, given beta and the age-specific fertility of a species, so that the population remains stable and stationary. Methods are inspired by ``Survival profiles from linear models versus Weibull models: Estimating stable and stationary population structures for Pleistocene large mammals" (Martín-González et al. 2019) <[doi:10.1016/j.jasrep.2019.03.031](https://doi.org/10.1016/j.jasrep.2019.03.031)>.

**License** GPL-3

**Encoding** UTF-8

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alpha_objective	<i>Objective Function for uniroot: Finds the Difference Between Births and 1</i>
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**Description**

This function calculates the difference between the number of births, as calculated with the given values of alpha, beta, and fertility\_rates, and the target value of 1.

**Usage**

```
alpha_objective(alpha, beta, fertility_rates)
```

**Arguments**

alpha            A numeric value representing the alpha parameter.

beta             A numeric value representing the beta parameter.

fertility\_rates    A numeric vector containing the fertility rates.

**Details**

Typically used as the objective function in root-finding algorithms such as [uniroot](#), to determine the value of alpha that results in exactly one birth.

This function depends on [calculate\\_population](#), which must be available in your package namespace.

**Value**

A numeric value giving the difference between the number of births (as calculated) and 1.

**See Also**

[uniroot](#)

**Examples**

```
# Basic usage
alpha_objective(0.5, 1.2, c(0.2, 0.3, 0.5, 0.4))

# Example with uniroot:
fertility_rates <- c(0.2, 0.3, 0.5, 0.4)
beta <- 1.2
res <- uniroot(
  alpha_objective,
  interval = c(0.000001, 100),
  beta = beta,
```

```
    fertility_rates = fertility_rates
  )
  res$root
```

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calculate\_population *Calculates the population for each age group*

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### Description

This function calculates the population for each age group and the number of births.

### Usage

```
calculate_population(alpha, beta, fertility_rates)
```

### Arguments

alpha	A numeric value representing the scale parameter ( $\alpha$ ) of the Weibull distribution. Note: In this context, alpha controls the horizontal scaling of the survival curve.
beta	A numeric value representing the shape parameter ( $\beta$ ) of the Weibull distribution. Note: Beta controls the shape of the survival curve (e.g., aging or failure rate).
fertility_rates	A vector of fertility rates for each age group.

### Value

A list with the following elements:

**population** A numeric vector giving the population size for each age group.

**births** A numeric value giving the total number of births.

### Examples

```
calculate_population(0.5, 1.2, c(0.2, 0.3, 0.5, 0.4))
```

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find_alphas	<i>Function to find the value of alpha</i>
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**Description**

This function finds the value of alpha using the uniroot method for a given beta and a vector of fertility rates. If the function values at the interval ends do not have opposite signs, it returns the closest value to 0.

**Usage**

```
find_alphas(beta, fertility_rates, tol = 1e-22)
```

**Arguments**

beta	A numeric value representing the beta parameter of Weibull distribution.
fertility_rates	A numeric vector containing the fertility rates.
tol	A numeric value representing the tolerance for the uniroot method. Default is 1e-22.

**Value**

A numeric value giving the estimated value of alpha, either found by uniroot or selected as the endpoint closest to zero if the root is not bracketed.

**Examples**

```
find_alphas(1.2, c(0.2, 0.3, 0.5, 0.4))
```

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run_analysis	<i>Run Analysis on Excel Data and Export Results</i>
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**Description**

This function reads fertility rate data and Beta value from an Excel file, processes it, and exports the results to a new Excel file for the species, including population matrices and calculated alpha/beta values.

**Usage**

```
run_analysis()
```

## Details

The function relies on functions from the **readxl** and **openxlsx** packages to handle Excel files.

The following external functions are used:

- [excel\\_sheets](#): List all sheet names in an Excel file.
- [read\\_excel](#): Read data from an Excel file.
- [createWorkbook](#): Create a new Excel workbook.
- [addWorksheet](#): Add a worksheet to a workbook.
- [writeData](#): Write data to a worksheet.
- [saveWorkbook](#): Save the workbook to a file.

Please refer to the documentation of those packages for more details.

## Value

No return value. Called for side effects (reading data, writing Excel files, and printing messages).

## See Also

[excel\\_sheets](#), [read\\_excel](#), [createWorkbook](#), [addWorksheet](#), [writeData](#), [saveWorkbook](#)

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weibull\_survival

*Weibull function for the survival rate*

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## Description

This function calculates the survival rate to reach a specific age using the Weibull function.

## Usage

```
weibull_survival(alpha, beta, age)
```

## Arguments

alpha	A numeric value representing the scale parameter of the Weibull distribution.
beta	A numeric value representing the shape parameter of the Weibull distribution.
age	A numeric value representing the age.

## Value

A numeric value giving the survival rate (probability) for reaching the given age.

## Examples

```
weibull_survival(1.5, 0.8, 10)
```

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