

Package: solong (via r-universe)

September 3, 2024

Type Package

Title Allometric Equations for Southern Ocean Taxa

Version 0.3.0

Description Provides allometric equations that relate the body size of Southern Ocean taxa to body part measurements.

URL <https://github.com/SCAR/solong>

BugReports <https://github.com/SCAR/solong/issues>

Depends R (>= 3.3.0)

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Encoding UTF-8

LazyData true

Imports assertthat, digest, dplyr, rlang, units (>= 0.5-0), vctrs

Suggests covr, ggplot2, knitr, rfishbase (>= 3.1.7), testthat, worrms

RoxxygenNote 7.2.0

VignetteBuilder knitr

Repository <https://scar.r-universe.dev>

RemoteUrl <https://github.com/SCAR/solong>

RemoteRef HEAD

RemoteSha 63c313dd618066849018e68d63c5bd462fc14bc6

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allometric_equations *Allometric equation data.*

Description

A dataset containing allometric equations (relationships between the size of an organism and the size of its body parts).

Usage

`allometric_equations`

Format

A data frame with variables:

- equation_id** the identifier of this equation
- taxon_name** name of the taxon
- taxon_aphia_id** the Aphia ID of the taxon (identifier within the World Register of Marine Species)
- equation** a function encoding the allometric equation
- inputs** a data.frame specifying (in order) the inputs to the equation, with columns 'property' (the property name of the measurement needed, e.g. 'lower rostral length'), and 'units' (the units of the measurement needed, e.g. 'mm')
- return_property** the name of the body size characteristic that is estimated by this equation (e.g. 'mantle length')
- return_units** the units of measurement of the returned property
- reliability** a data.frame with indicators of the reliability of the equation: type (a description of how the reliability was assessed, e.g. 'R^2' or 'N' the sample size used by the authors of the equation) and value (its value)
- notes** notes
- reference** the source of the equation

solong

solong

Description

Provides allometric equations that relate the body size of Southern Ocean taxa to body part measurements.

sol_allometry	<i>Apply allometric equations</i>
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Description

Apply allometric equations

Usage

```
sol_allometry(data, equation)
```

Arguments

data	data.frame: input data
equation	character or sol_equation object: either the identifier of the equation to apply, or the equation object itself. Can be a single element (this equation will be applied to all rows of the data) or with length matching the number of rows of the data

Value

the input data frame, augmented with columns "allometric_property", "allometric_value", "allometric_value_lower", and "allometric_value_upper"

See Also

[allometric_equations](#)

Examples

```
## Not run:  
x <- data.frame(LRL=c(11.3,13.9),species=c("Architeuthis dux"),  
stringsAsFactors=FALSE)  
## it doesn't matter what the column names are, but we  
## need to set the property types correctly  
x$LRL <- sol_set_property(x$LRL,"lower rostral length")  
  
## apply a single equation to all rows  
sol_allometry(x,c("342218_ML_Roel2000"))  
  
## apply a different equation to each row  
sol_allometry(x,c("342218_ML_Roel2000","342218_ML_Clar1986"))  
  
## End(Not run)
```

sol_equation *Allometric equations*

Description

Allometric equations

Usage

```
sol_equation(id)
sol_equations(id)
```

Arguments

id	string: the identifier of the equation to return
----	--

Value

data.frame

Examples

```
sol_equations()
sol_equation("342218_ML_Roel2000")
```

sol_fb_length_weight *Create allometric equation from Fishbase*

Description

Experimental. Requires the rfishbase package to be installed.

Usage

```
sol_fb_length_weight(
  ...,
  input_properties,
  return_properties,
  worms = requireNamespace("worrms", quietly = TRUE)
)
```

Arguments

... : arguments passed to rfishbase::length_weight
 input_properties character: an optional vector of properties (see link{sol_properties}). Only equations that take an input in input_properties will be returned
 return_properties character: an optional vector of properties (see link{sol_properties}). Only equations that return a value in return_properties will be returned
 worms logical: if TRUE, and if the worrms package is installed, try and find the AphidID for the taxon in the World Register of Marine Species

Value

A tibble of equation(s)

See Also

[sol_equation](#) [sol_equations](#)

Examples

```
## Not run:
library(dplyr)
eq <- sol_fb_length_weight("Electrona antarctica", input_properties = "standard length")
x <- tibble(SL = 10) %>%
  mutate(SL = sol_set_property(SL, "standard length", with_units = "cm"))
sol_allometry(x, eq)

## End(Not run)
```

sol_make_equation *Create an allometric equation object*

Description

Create an allometric equation object

Usage

```
sol_make_equation(
  equation_id,
  taxon_name,
  taxon_aphia_id,
  equation,
  inputs,
  return_property,
```

```

    return_units,
    reliability,
    notes,
    reference,
    check_packaged_ids = TRUE,
    warn_recommended = TRUE
)

```

Arguments

<code>equation_id</code>	string: a unique identifier for the equation (required)
<code>taxon_name</code>	string: the taxon name that the equation applies to (required)
<code>taxon_aphia_id</code>	numeric: the AphiaID of the taxon that the equation applies to (recommended)
<code>equation</code>	function: the equation. Must return a data.frame or tibble, with at least the column "allometric_value", and optionally also "allometric_value_lower" and "allometric_value_upper" (required)
<code>inputs</code>	data.frame: the inputs needed by the equation. Must have columns "property" and "units", with entries that match those in <code>sol_properties</code> . Optionally also "sample_minimum" and "sample_maximum" if known (describing the range of the data used to generate the equation) (required)
<code>return_property</code>	string: the name of the allometric property that the equation returns (required)
<code>return_units</code>	string: the units of measurement of the allometric property that the equation returns. Will be parsed by <code>units::as_units</code> (required)
<code>reliability</code>	data.frame: indicators of reliability of the equation. Must have columns "type" and "value"; see examples (recommended)
<code>notes</code>	string: any notes that users should be aware of (optional)
<code>reference</code>	bibentry: the source of the equation (recommended)
<code>check_packaged_ids</code>	logical: if TRUE, check the <code>equation_id</code> against the package-bundled equations. A warning will be issued if there is a packaged equation with the same ID as <code>equation_id</code>
<code>warn_recommended</code>	logical: issue a warning if "recommended" informations is not supplied?

Value

equation object

See Also

[sol_equation](#) [sol_equations](#) [sol_properties](#)

Examples

```
library(dplyr)
my_ref <- bibentry(bibtype="Article",key="Lake2003",
                    author=c(person("S","Lake"),person("H","Burton"),
                            person("J","van den Hoff")),
                    year=2003,
                    title="Regional, temporal and fine-scale spatial variation in
                           Weddell seal diet at four coastal locations in east Antarctica",
                    journal="Marine Ecology Progress Series",
                    volume=254, pages="293–305", doi="10.3354/meps254293")

eq <- sol_make_equation(equation_id="my_equation_id",
                        taxon_name="Chorisimus antarcticus",
                        taxon_aphia_id=369214,
                        equation=function(L)
                            tibble(allometric_value=0.000943*(L^2.976)),
                            inputs=tibble(property="carapace length",units="mm",
                                         sample_minimum=6,sample_maximum=16),
                            return_property="wet weight",
                            return_units="g",
                            reliability=tribble(~type,~value,
                                         "N",35,
                                         "R^2",0.976),
                            reference=my_ref)
```

`sol_properties`

Properties

Description

Properties

Usage

```
sol_properties(prop)
```

Arguments

prop	string: if provided, return only the property matching this name
------	--

Value

`data.frame`

See Also

[sol_set_property](#)

Examples

```
sol_properties() ## all properties that solong knows about
```

sol_properties_data *Properties relating to allometric equations.*

Description

Properties relating to allometric equations.

Usage

```
sol_properties_data
```

Format

A data frame with variables:

property the property name
units its units of measurement
class_name the corresponding class name used internally by the solong package
notes notes, including a specific definition of the property if appropriate

sol_set_property *Set or get the property name*

Description

Set or get the property name

Usage

```
sol_set_property(x, prop, with_units, ...)
```

```
sol_get_property(x)
```

Arguments

x	vector: data
prop	string: property name
with_units	string: units of measurement to use. If missing, the default units for the property will be used
...	: extra arguments, currently ignored

Value

x with additional class set

See Also

[sol_properties](#)

Examples

```
x <- data.frame(LRL=c(11.3,13.9),species=c("Architeuthis dux"),
  stringsAsFactors=FALSE)
## it doesn't matter what the column names are, but we
## need to set the property types correctly
x$LRL <- sol_set_property(x$LRL,"lower rostral length")

## remove the property
x$LRL <- sol_set_property(x$LRL,NULL)
```

sol_vonbert

Von Bertalanffy growth equation with optional propagation of uncertainty

Description

Intended for use inside of `sol_equation` objects.

Usage

```
sol_vonbert(
  t,
  Linf,
  k,
  t0,
  Linf_se,
  k_se,
  t0_se,
  reps = 1000L,
  ci = 0.95,
  method = "monte carlo"
)
```

Arguments

t	numeric: prediction times
Linf	numeric: Linf parameter estimate
k	numeric: k parameter estimate

<code>t0</code>	numeric: t0 parameter estimate
<code>Linf_se</code>	numeric: Linf parameter standard error estimate
<code>k_se</code>	numeric: k parameter standard error estimate
<code>t0_se</code>	numeric: t0 parameter standard error estimate
<code>reps</code>	integer: number of Monte-Carlo samples to draw
<code>ci</code>	numeric: confidence level
<code>method</code>	string: method to use for uncertainty propagation (only "monte carlo" supported at the moment)

Value

A tibble with columns `allometric_value`, `allometric_value_lower`, `allometric_value_upper`. If any of the standard error inputs are missing, `NULL`, or `NA`, the upper and lower estimates will be `NA_real_`

Examples

```
sol_vonbert(0:7, Linf = 80.7, Linf_se = 0.82, k = 0.25, k_se = 0.01, t0 = -2.31, t0_se = 0.01)
```

strip_units

Remove the units from an object

Description

A convenience function to remove the units assigned to an object.

Usage

```
strip_units(x)
```

Arguments

<code>x</code>	object: with units
----------------	--------------------

Value

`x`, with units removed

Examples

```
x <- data.frame(LRL=c(11.3,13.9),species=c("Architeuthis dux"),  
  stringsAsFactors=FALSE)  
x$LRL <- sol_set_property(x$LRL,"lower rostral length")  
## apply an allometric equation  
xa <- sol_allometry(x,c("342218_ML_Roel2000"))  
  
strip_units(xa$allometric_value)
```

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