# Package 'EloSteepness'

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Title Bayesian Dominance Hierarchy Steepness via Elo Rating and

```
David's Scores
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     mann and Fischer (2022) <doi:10.1101/2022.01.28.478016>). Steepness estima-
     tion is based on Bayesian implementations of either Elo-rating or David's scores.
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EloSteepness-package The 'EloSteepness' package.

# Description

Dominance Hierarchy Steepness Via Elo Rating

catch\_warnings

catch warnings alongside results without returning warning

# Description

helper function

# Usage

catch\_warnings(expr)

# Arguments

expr

an R expression to evaluate

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#### Value

a list where the first entry is the result of expr and the second provides information about warnings

#### Source

```
demo(error.catching)
```

## **Examples**

```
log(3)
catch_warnings(log(3))

# produces warning
# log(-3)
# catch it
catch_warnings(log(-3))

# produces error
# log("x")
# catch it
catch_warnings(log("x"))
```

davids\_steepness

David's scores and steepness with Bayesian flavor

#### **Description**

David's scores and steepness with Bayesian flavor

# Usage

```
davids_steepness(mat, silent = FALSE, ...)
```

#### **Arguments**

```
mat square interaction matrix
silent logical, suppress warnings (default is FALSE)
... additional arguments for sampling()
```

#### Value

a list with results of the modelling fitting, containing the following list items:

steepness a one-column matrix with the posterior samples for steepness. Each row is one iteration

norm\_ds an matrix with posterior normalized David's scores for each individual. Each column is one individual. Each row is one iteration.

ids a character vector with individual ID codes as supplied in mat diagnostics a list with information regarding sampling problems stanfit the actual stanfit object mat the input matrix

## **Examples**

```
data(dommats, package = "EloRating")
res <- davids_steepness(dommats$elephants, refresh = 0)
plot_steepness(res)</pre>
```

```
{\tt elo\_steepness\_from\_matrix}
```

steepness based on Bayesian Elo-rating

# Description

for interaction data with unknown sequence of observations

## Usage

```
elo_steepness_from_matrix(
  mat,
  algo = c("fixed_sd", "original"),
  n_rand = NULL,
  silent = FALSE,
   ...
)
```

## **Arguments**

mat	square interaction matrix
algo	character, either "fixed_sd" or "original". This determines which algorithm is used. Default is "fixed_sd", which is a slight modification from Goffe et al's original code.
n_rand	numeric, number of randomized sequences. Default is NULL, which uses a rule of thumb to determine the number (see below for more details).
silent	logical, suppress warnings (default is FALSE)
	additional arguments for sampling()

#### **Details**

The number of randomizations is set in the following way, unless a specific number is provided. If there are more than 500 observed interactions,  $n_rand = 5$ . If there are less than 100 interactions,  $n_rand = 50$ . In the remaining cases,  $n_rand = 20$ .

```
If the function call produces warnings about divergent transitions, large Rhat values or low effective sample sizes, increase the number of iterations (via \code{iter=}) and/or adjust the sampling controls (e.g. via \code{control = list(adapt_delta = 0.9)}).
```

#### Value

a list with results of the modelling fitting, containing the following list items:

steepness a matrix with the posterior samples for steepness. Each column corresponds to one randomization (as set via n\_rand). Each row is one iteration.

cumwinprobs an array with posterior cumulative winning probabilities for each individual.

ids a character vector with individual ID codes as supplied in mat

diagnostics a list with information regarding sampling problems

```
stanfit the actual stanfit object
```

mat the input matrix

algo character, describing whether the original fitting algorithm was used ("original") or the one with fixed SD of start ratings ("fixed\_sd")

sequence\_supplied logical, were data supplied as matrix (FALSE) or as sequence via winner/loser vector (TRUE)

```
elo_steepness_from_sequence
```

steepness based on Bayesian Elo-rating

## **Description**

for interaction data with known sequence of observations

## Usage

```
elo_steepness_from_sequence(
  winner,
  loser,
  algo = c("fixed_sd", "original"),
  silent = FALSE,
   ...
)
```

#### **Arguments**

```
winner character (or factor) of winning individuals
loser character (or factor) of losing individuals
algo character, either "fixed_sd" or "original". This determines which algorithm is used. Default is "fixed_sd", which is a slight modification from Goffe et al's original code.
silent logical, suppress warnings (default is FALSE)
... additional arguments for sampling()
```

## Value

a list with results of the model fitting (see elo\_steepness\_from\_matrix) for details

```
generate_interaction_probs
```

generate dyadic interaction probabilities for a group with fixed individual and dyadic biases

## **Description**

generate dyadic interaction probabilities for a group with fixed individual and dyadic biases

## Usage

```
generate_interaction_probs(n_ind, id_bias = 0, rank_bias = 0)
```

## **Arguments**

n_ind	numeric, number of individuals
id_bias	numeric, between 0 and 1. If 0 all individual are equally likely to interact. If 1, some individuals have higher propensities to interact
rank_bias	numeric, between 0 and 1. If 0 there is no relationship between rank distance and interaction propensity. If 1 there is a strong relationship: dyads closer in rank interact more often.

#### Value

a matrix

```
x <- generate_interaction_probs(n_ind = 10, id_bias = 0.2, rank_bias = 1)
rankdiff <- x[, 2] - x[, 1]
interactprob <- x[, "final"]
# closer in rank (smaller rank diff) = interaction more likely
plot(rankdiff, interactprob)

x <- generate_interaction_probs(n_ind = 10, id_bias = 0.2, rank_bias = 0)
rankdiff <- x[, 2] - x[, 1]
interactprob <- x[, "final"]
# approx. equal probs for all dyads regardless of rank diff
plot(rankdiff, interactprob)

x <- generate_interaction_probs(n_ind = 10, id_bias = 0, rank_bias = 0)
interactprob <- x[, "final"]
y <- sample(1:nrow(x), 1000, replace = TRUE, prob = interactprob)
y <- as.numeric(x[y, 1:2])
# approx. equal numbers of interactions per ID</pre>
```

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```
sort(table(y))

# skewed interaction numbers
x <- generate_interaction_probs(n_ind = 10, id_bias = 1, rank_bias = 0)
interactprob <- x[, "final"]
y <- sample(1:nrow(x), 1000, replace = TRUE, prob = interactprob)
y <- as.numeric(x[y, 1:2])
sort(table(y))</pre>
```

plot\_matrix

plot (rather than print) a matrix

# **Description**

a helper function

#### Usage

```
plot_matrix(mat, greyout = NULL, prunkcol = NULL)
```

#### **Arguments**

mat square matrix

greyout numeric, the values to be grayed out

prunkcol color value, which if set to some color will highlight unknown relationships with

rectangles of that color.

## Value

a plot

plot\_scores

plot individual scores (either Elo or David)

#### **Description**

plot individual scores (either Elo or David)

# Usage

```
plot_scores(
    x,
    adjustpar = 4,
    color = TRUE,
    subset_ids = NULL,
    include_others = TRUE
)
```

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# **Arguments**

X	result from elo_steepness_from_matrix, elo_steepness_from_sequence or davids_steepness
adjustpar	numeric, parameter for smoothing posterior of individual scores
color	logical, default is TRUE where individuals get color-coded. If FALSE: a gray scale is used. It is also possible to hand over a vector with colors, which then must correspond in length to the number of individuals.
subset_ids	character, plot only those individual codes. Default is $NULL$ , i.e. all individuals are included in the plot.
include_others	logical, should other IDs (those <i>not</i> in subset_ids) be included as contours. Default is TRUE. This only has an effect if subset_ids is different from NULL,

## Value

a plot

# **Examples**

plot\_steepness

plot steepness density

# Description

plot steepness density

# Usage

```
plot_steepness(x, adjustpar = 1.5, print_numbers = TRUE)
```

# Arguments

X	result from elo_steepness_from_matrix, elo_steepness_from_sequence or
	davids_steepness
adjustpar	numeric, parameter for smoothing posterior of individual scores
<pre>print_numbers</pre>	logical, if TRUE (default) print numeric summaries into into the plot and omit
	them if FALSE

# Value

a plot

# **Examples**

```
plot_steepness_regression plot steepness regression
```

# Description

visually combine individual scores with group-level steepness

## Usage

```
plot_steepness_regression(
    x,
    adjust = 3,
    color = TRUE,
    width_fac = 0.1,
    axis_extend = 0.1
)
```

# Arguments

X	result from elo_steepness_from_matrix, elo_steepness_from_sequence or davids_steepness
adjust	numeric, parameter for smoothing posterior of individual scores
color	logical, default is TRUE where individuals get color- coded. If FALSE: a gray scale is used. It is also possible to hand over a vector with colors, which then must be correspond in length to the number of individuals.
width_fac	numeric, relative width of posterior distributions. This is actually affects the 'height' but since the posteriors are rotated it visually represents width.
axis_extend	numeric, an extension factor to extend the horizontal axis to leave space for the posteriors. When set to $0$ the axis stops at $n$ (the number of individuals, which represents the lowest rank).

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## Value

a plot

## **Examples**

```
data("bonobos", package = "EloRating")
res <- davids_steepness(bonobos, refresh = 0, iter = 1000)
plot_steepness_regression(res, width_fac = 0.5)</pre>
```

prep\_data\_for\_rstan

prepare data for stan call

#### **Description**

prepare data for stan call

#### Usage

```
prep_data_for_rstan(mat, n_rand = 1, silent = FALSE, for_elo_model = TRUE)
```

## **Arguments**

mat square interaction matrix

n\_rand numeric, number of randomizations

silent logical, omit printing messages regarding non-fatal data issues. Default is FALSE,

i.e. do print messages.

for\_elo\_model logical, output ready for Elo steepness (default, TRUE). If FALSE, prep for David's

score steepness.

#### Value

a list that is formatted so that it can be handed over to the respective Stan models

remove\_dyads

remove interactions from matrix to increase sparseness

#### **Description**

remove interactions from matrix to increase sparseness

#### Usage

```
remove_dyads(
   m,
   removal_mode = c("mix", "by_interaction", "by_dyad"),
   stop_at = 0.5,
   max_out = NULL
)
```

#### **Arguments**

m input matrix

removal\_mode character, should interactions be removed interaction by interaction ("by\_interaction"),

or by removing one dyad entirely at a time ("by\_dyad"). Default is "mix", i.e.

a random mix between the two strategies.

stop\_at numeric, fraction of unknown relationships to be reached

max\_out numeric, the number of matrices to be returned maximally. This is useful if the

input matrix is fairly large. If set, this will return the input matrix plus max\_out randomly selected matrices from the remaining produced matrices. So in fact, the output comprises max\_out + 1 matrices (subject to the stop\_at specifica-

tion).

#### Value

a list with two items. \$summary is a data frame with an overview. matrices contains the actual interaction matrices with increasing proportion of unknown relationships.

#### **Examples**

```
data(bonobos)
res <- remove_dyads(bonobos)
res$summary
length(res$matrices)
lapply(res$matrices, prunk)

res <- remove_dyads(bonobos, max_out = 2)
# first plus two randomly selected = 3 matrices
length(res$matrices)
res$summary</pre>
```

repeatability\_steepness

steepness via repeatability (cf aniDom package)

# Description

steepness via repeatability (cf aniDom package)

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## Usage

```
repeatability_steepness(mat, n_rand = 1000)
```

## **Arguments**

mat square interaction matrix

n\_rand numeric, number of randomized sequences (default is 1000)

#### Value

a steepness value

#### References

Sanchez-Tojar et al 2018

## **Examples**

```
data(bonobos, package = "EloRating")
repeatability_steepness(bonobos, n_rand = 20)
```

sampler\_diagnostics

catch Stan sampling issues without throwing a warning

# Description

catch Stan sampling issues without throwing a warning

## Usage

```
sampler_diagnostics(object)
```

# Arguments

object stanfit object

#### Value

a list regarding any sampling issues encountered during fitting

simple\_steep\_gen

scores

numeric summaries of individual Elo-ratings or David's scores

## Description

numeric summaries of individual Elo-ratings or David's scores

## Usage

```
scores(x, quantiles = c(0.045, 0.955))
```

## **Arguments**

```
x result from elo_steepness_from_matrix or davids_steepness
quantiles numeric, the quantiles to be returned
```

#### Value

a data frame with one line per individual, providing summaries of posteriors for individual scores

# **Examples**

```
data("bonobos", package = "EloRating")
res <- davids_steepness(bonobos, refresh = 0, cores = 2)
scores(res)

data("dommats", package = "EloRating")
m <- dommats$elephants
res <- elo_steepness_from_matrix(m, n_rand = 1, refresh = 0, cores = 2)
scores(res)</pre>
```

simple\_steep\_gen

generate dominance interactions with specified steepness

## **Description**

generate dominance interactions with specified steepness

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#### Usage

```
simple_steep_gen(
  n_ind,
  n_int,
  steep,
  id_bias = 0,
  rank_bias = 0,
  sequential = TRUE
)
```

#### **Arguments**

n_ind	integer, the number of individuals
n_int	integer, the number of interactions
steep	numeric (between 0 and 1), the desired steepness value
id_bias	numeric, between 0 and 1. If 0 all individual are equally likely to interact. If 1, some individuals have higher propensities to interact.
rank_bias	numeric, between 0 and 1. If 0 there is no relationship between rank distance and interaction propensity. If 1 there is a strong relationship: dyads closer in rank interact more often.
sequential	logical, default is TRUE. See details.

#### **Details**

Initially (and this is still the default), the function generated interactions and their outcomes sequentially: first a dyad was chosen that interacted and then its winner was determined. This was repeated for as many interactions as set by  $n_i$ t=.

The same results can be achieved much more efficiently by first setting the number of interactions per dyad and then looping through all dyads and then generate the interactions and their outcomes per dyad. This can be achieved by setting sequential = FALSE. In this latter case the 'sequence' of interactions reported in the results is just a randomized version of all interactions, whereas in the former case there is a 'natural sequence' (although it is meaningless because the sequence is irrelevant with respect to outcomes of individual interactions (the system is stable)).

#### Value

a list with the first item being the interactions in sequence form (\$sequence). The second item (\$matrix) is the square interaction matrix and the third item (\$settings) is a list with input settings (including probabilities to interact for each dyad).

```
res <- simple_steep_gen(n_ind = 5, n_int = 30, steep = 0.99)
res$sequence
res$matrix</pre>
```

steepness\_precis

```
library(EloRating)
steeps <- runif(20, 0, 1)
nids <- sample(6:10, length(steeps), TRUE)
mats <- sapply(1:length(steeps), function(x) {
    simple_steep_gen(nids[x], nids[x] ^ 2.5, steeps[x], 0)[[2]]
    })
obs_steeps <- unlist(lapply(mats, function(x)steepness(x)[1]))
plot(steeps, obs_steeps, xlim = c(0, 1), ylim = c(0, 1))
abline(0, 1)</pre>
```

steepness\_precis

numeric summary of steepness

# Description

numeric summary of steepness

#### Usage

```
steepness_precis(x, quantiles = c(0.055, 0.25, 0.75, 0.945))
```

#### **Arguments**

#### Value

a data.frame with one row providing a summary of the steepness posterior

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```
summary.elo_steepness summary
```

# Description

summary

## Usage

```
## S3 method for class 'elo_steepness'
summary(object, ...)
## S3 method for class 'david_steepness'
summary(object, ...)
```

## **Arguments**

 $object \\ result from \verb|elo_steepness_from_matrix|, \verb|elo_steepness_from_sequence| or \\$ 

davids\_steepness

... further arguments passed to or from other methods (ignored)

#### Value

Nothing returned. Called for side effects of textual output to console.

upward\_steepness

proportion of interactions against the rank order

## **Description**

proportion of interactions against the rank order

## Usage

```
upward_steepness(mat)
```

#### **Arguments**

mat

square interaction matrix

## Value

numeric value of upward steepness

```
data(bonobos, package ="EloRating")
upward_steepness(bonobos)
```

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