

# Package ‘bpcs’

December 9, 2020

**Title** Bayesian Paired Comparison Analysis with Stan

**Version** 1.0.0

## Description

Models for the analysis of paired comparison data using Stan. The models include Bayesian versions of the Bradley-Terry model, including random effects (1 level), generalized model for predictors, order effect (home advantage) and the variations for the Davidson (1970) model to handle ties. Additionally, we provide a number of functions to facilitate inference and obtaining results with these models. References: Bradley and Terry (1952) <doi:10.2307/2334029>; Davidson (1970) <doi:10.1080/01621459.1970.10481082>; Carpenter et al. (2017) <doi:10.18637/jss.v076.i01>.

**URL** <https://github.com/davidissamattos/bpcs>,  
<https://davidissamattos.github.io/bpcs/>

**BugReports** <https://github.com/davidissamattos/bpcs/issues>

**License** MIT + file LICENSE

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.1

**Biarch** true

**Depends** R (>= 4.0.0)

**Imports** coda, dplyr, tidyr, stringr, ggplot2, gtools, methods, Rcpp (>= 0.12.0), RcppParallel (>= 5.0.1), rstan (>= 2.20.0), rstantools (>= 2.1.0), tibble, tidyselect, HDInterval, shinystan, loo, magrittr, badger, stats, rlang, knitr

**LinkingTo** BH (>= 1.66.0), Rcpp (>= 0.12.0), RcppEigen (>= 0.3.3.3.0), RcppParallel (>= 5.0.1), rstan (>= 2.20.0), StanHeaders (>= 2.20.0)

**SystemRequirements** GNU make

**Suggests** rmarkdown, testthat, covr, bayesplot, kableExtra

**VignetteBuilder** knitr

**NeedsCompilation** yes

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**Repository** CRAN

**Date/Publication** 2020-12-09 09:20:16 UTC

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bpcs-package *bpcs - A package for Bayesian Paired Comparison analysis with Stan*

**Description**

bpcs - A package for Bayesian Paired Comparison analysis with Stan

**References**

Stan Development Team (2020). RStan: the R interface to Stan. R package version 2.21.2. <https://mc-stan.org>

bpc *Bayesian Paired comparison regression models in Stan*

**Description**

This is the main function of the package. This function utilizes precompiled stan models to sample the posterior distribution of the specified model with the input data. For more information and larger examples of usage see the vignettes.

**Usage**

```
bpc(
  data,
  player0,
  player1,
  player0_score = NULL,
  player1_score = NULL,
  result_column = NULL,
  z_player1 = NULL,
  cluster = NULL,
  predictors = NULL,
  model_type,
```

```

  solve_ties = "random",
  win_score = "higher",
  priors = NULL,
  chains = 4,
  iter = 2000,
  warmup = 1000,
  show_chain_messages = TRUE,
  seed = NA
)

```

## Arguments

<code>data</code>	A data frame containing the observations. The other parameters specify the name of the columns
<code>player0</code>	A string with name of the column containing the players 0. This column should be of string/character type and not be of factor type.
<code>player1</code>	A string with name of the column containing the players 0. This column should be of string/character type and not be of factor type.
<code>player0_score</code>	A string with name of the column containing the scores of players 0
<code>player1_score</code>	A string with name of the column containing the scores of players 1
<code>result_column</code>	A string with name of the column containing the winners. 0 for player 0, 1 for player 1 and 2 for ties
<code>z_player1</code>	A string with the name of the column containing the order effect for player 1. E.g. if player1 has the home advantage this column should have 1 otherwise it should have 0
<code>cluster</code>	A string with the name of the column containing the cluster for the observation. To be used with a random effects model. This column should contain strings
<code>predictors</code>	A data frame that contains the players predictors values when using a generalized model. Only numeric values are accepted. Booleans are accepted but will be cast into integers. The first column should be for the player name, the others will be the predictors. The column names will be used as name for the predictors
<code>model_type</code>	We first add a base model 'bt' or 'davidson' and then additional options with '-' <ul style="list-style-type: none"> <li>• 'bt' for the Bradley Terry model. Ref: Bradley-Terry 1952,</li> <li>• 'davidson' the Davidson model to handle for ties. Ref: Davidson 1970</li> <li>• 'bt-ordereffect' for the Bradley-Terry with order effect, for home advantage. Ref: Davidson 1977</li> <li>• 'davidson-ordereffect' for the Davidson model with order effect, for home advantage, and ties. Ref: Davidson 1977</li> <li>• 'bt-generalized': for the generalized Bradley Terry model for subject specific predictors. Ref: Springall 1973</li> <li>• 'davidson-generalized' for the generalized Davidson model for subject specific predictors</li> <li>• 'bt-U': for the Bradley-Terry with random effects. Ref: Bockenholt 2001</li> <li>• 'davidson-U': For Davidson model with random effects</li> </ul>

	<ul style="list-style-type: none"> <li>• 'bt-ordereffect-U' for Bradley-Terry with order effects and random effects, use similar syntax for other variations by appending the correct options</li> </ul>
solve_ties	<p>A string for the method of handling ties.</p> <ul style="list-style-type: none"> <li>• 'random' for converting ties randomly,</li> <li>• 'remove' for removing the tie occurrences</li> <li>• 'none' to ignore ties. This requires a model capable of handling ties</li> </ul>
win_score	<p>A string that indicates if which score should win</p> <ul style="list-style-type: none"> <li>• 'higher' score is winner</li> <li>• 'lower' score is winner</li> </ul>
priors	<p>A list with the parameters for the priors.</p> <ul style="list-style-type: none"> <li>• 'prior_lambda_mu' Mean value of the lambda parameter in the all models. For the generalized this is also the prior for the B the parameter for lambda <math>\sim</math> normal(mu, std)</li> <li>• 'prior_lambda_std' Standard deviation of the lambda parameter in the all models. <math>\lambda \sim</math> normal(mu, std)</li> <li>• 'prior_nu_mu' Mean value of the nu parameter in the Davidson models. <math>\nu \sim</math> normal(mu, std)</li> <li>• 'prior_nu_std' Standard deviation of nu parameter in the Davidson models. <math>\nu \sim</math> normal(mu, std). Default = 0.3</li> <li>• 'prior_gm_mu' Mean value of the gm in the ordered effect model. <math>gm \sim</math> normal(mu, std). Default = 0</li> <li>• 'prior_gm_std' Standard deviation of the gm parameter in the ordered effect model. <math>gm \sim</math> normal(mu, std). Default =</li> <li>• 'prior_U_std' Standard deviation of the U parameter in the random effects model. <math>U \sim</math> normal(0, std). Default = 3.0</li> </ul>
chains	<p>Number of chains passed to Stan sampling. Positive integer, default=4. For more information consult Stan documentation</p>
iter	<p>Number of iterations passed to Stan sampling. Positive integer, default =2000. For more information consult Stan documentation</p>
warmup	<p>Number of iteration for the warmup passed to Stan sampling. Positive integer, default 1000. For more information consult Stan documentation</p>
show_chain_messages	<p>Hide chain messages from Stan</p>
seed	<p>a random seed for Stan</p>

### Value

An object of the class bpc. This object should be used in conjunction with the several auxiliary functions from the package

### References

1. Bradley RA, Terry ME 1952. Rank Analysis of Incomplete Block Designs I: The Method of Paired Comparisons. *Biometrika*, 39, 324-45.

2. Davidson RR 1970. On Extending the Bradley-Terry Model to Accommodate Ties in Paired Comparison Experiments. Journal of the American Statistical Association, 65, 317-328.
3. Davidson, Roger R., and Robert J. Beaver 1977. "n extending the Bradley-Terry model to incorporate within-pair order effects. Biometrics: 693-702.
4. Stan Development Team 2020. RStan: the R interface to Stan. R package version 2.21.2.
5. Bockenholt, Ulf. Hierarchical modeling of paired comparison data. Psychological Methods 6.1 2001: 49.
6. Springall, A. Response Surface Fitting Using a Generalization of the Bradley-Terry Paired Comparison Model. Journal of the Royal Statistical Society: Series C Applied Statistics 22.1 1973: 59-68.

## Examples

```
#For the simple Bradley-Terry model
bpc(data = tennis_agresti,
     player0 = 'player0',
     player1 = 'player1',
     result_column = 'y',
     model_type = 'bt',
     solve_ties = 'none')
```

---

brasil\_soccer\_league *This is a dataset with the results matches from the first league of the Brazilian soccer championship from 2017-2019. It was reduced and translated from the adadueque/Brasileirao\_Dataset repository*

---

## Description

This is a dataset with the results matches from the first league of the Brazilian soccer championship from 2017-2019. It was reduced and translated from the adadueque/Brasileirao\_Dataset repository

## Usage

```
brasil_soccer_league
```

## Format

Data frame that contains 1140 matches and 9 Columns from the Brazilian soccer championship

- Time: time of the day in 24h format
- DayWeek: day of the week
- Date: date YY-MM-DD
- HomeTeam: name of the team playing home
- VisitorTeam: name of the team playing visitor

- Round: Round number of the championship
- Stadium: Name of the stadium where the game was played
- ScoreHomeTeam: number of goals for the home team
- ScoreVisitorTeam: number of goals for the visitor

### Source

[https://github.com/adaoduque/Brasileirao\\_Dataset](https://github.com/adaoduque/Brasileirao_Dataset)

---

check\_if\_there\_are\_na *Check for NA in the specific columns and returns T or F if there is at least 1 NA in those columns*

---

### Description

Check for NA in the specific columns and returns T or F if there is at least 1 NA in those columns

### Usage

```
check_if_there_are_na(  
  d,  
  player0,  
  player1,  
  player0_score = NULL,  
  player1_score = NULL,  
  result_column = NULL  
)
```

### Arguments

d	a data frame
player0	the name of column for player0
player1	the name of column for player1
player0_score	the name of column for player0 scores
player1_score	the name of column for player1 scores
result_column	the name of column for results

### Value

TRUE (there are NA) or FALSE (no NA)

check\_if\_there\_are\_ties

*Check if a data frame column contains ties*

---

**Description**

Check if a data frame column contains ties

**Usage**

```
check_if_there_are_ties(d_column)
```

**Arguments**

d\_column            a column with the values for the ties

**Value**

T (there are ties) or F (no ties)

---

check\_numeric\_predictor\_matrix

*Check if all values in the predictor matrix are numeric and not NA.  
Note that TRUE will be cast to 1 and FALSE will be cast to 0*

---

**Description**

Check if all values in the predictor matrix are numeric and not NA. Note that TRUE will be cast to 1 and FALSE will be cast to 0

**Usage**

```
check_numeric_predictor_matrix(predictor_matrix)
```

**Arguments**

predictor\_matrix  
                  a predictor matrix generated by the create\_predictor\_matrix\_with\_player\_lookup\_table function

**Value**

TRUE (correct) or FALSE (with problems)



---

check\_predictors\_df\_contains\_all\_players  
*Check if the predictor df contains all players and only those*

---

**Description**

Check if the predictor df contains all players and only those

**Usage**

```
check_predictors_df_contains_all_players(predictor_df, lookup_table)
```

**Arguments**

predictor\_df    the predictor input data frame  
lookup\_table    a lookup table of the players

**Value**

TRUE (correct) or FALSE (with problems)

---

check\_result\_column    *Check if a data frame column contains only the values 1 0 and 2. Used to check the format of the results*

---

**Description**

Check if a data frame column contains only the values 1 0 and 2. Used to check the format of the results

**Usage**

```
check_result_column(d_column)
```

**Arguments**

d\_column            a column from a data frame

**Value**

TRUE (correct) or FALSE (with problems)

---

check_z_column	<i>Check if a data frame column contains only the values 1 or 0. For the z column</i>
----------------	---

---

**Description**

Check if a data frame column contains only the values 1 or 0. For the z column

**Usage**

```
check_z_column(d_column)
```

**Arguments**

d\_column          a column of a data frame to be tested

**Value**

TRUE (correct) or FALSE (with problems)

---

compute_scores	<i>Giving a player0 an player1 scores, this functions adds one column to the data frame containing who won (0= player0 1=player1 2=tie) and another if it was a tie. The ties column superseeds the y column. If it was tie the y column does not matter y column: (0= player0 1=player1 2=tie) ties column (0=not tie, 1=tie)</i>
----------------	--

---

**Description**

Giving a player0 an player1 scores, this functions adds one column to the data frame containing who won (0= player0 1=player1 2=tie) and another if it was a tie. The ties column superseeds the y column. If it was tie the y column does not matter y column: (0= player0 1=player1 2=tie) ties column (0=not tie, 1=tie)

**Usage**

```
compute_scores(
  d,
  player0_score,
  player1_score,
  solve_ties = "random",
  win_score = "higher"
)
```

**Arguments**

d	dataframe
player0_score	name of the column in data
player1_score	name of the column in data
solve_ties	Method to solve the ties, either randomly allocate, or do nothing, or remove the row from the datasetc('random', 'none', 'remove').
win_score	decides if who wins is the one that has the highest score or the lowest score

**Value**

a dataframe with column 'y' that contains the results of the comparison and a ties column indicating if there was ties

---

compute_ties	<i>Giving a result column we create a new column with ties (0 and 1 if it has)</i>
--------------	--

---

**Description**

Giving a result column we create a new column with ties (0 and 1 if it has)

**Usage**

```
compute_ties(d, result_column)
```

**Arguments**

d	data frame
result_column	column where the result is

**Value**

dataframe with a column called ties

---

```
create_array_of_par_names
```

*Create an array with the parameter name and to what player/cluster it refers to in the order stan presents*

---

### Description

Create an array with the parameter name and to what player/cluster it refers to in the order stan presents

### Usage

```
create_array_of_par_names(par, lookup_table, cluster_lookup_table = NULL)
```

### Arguments

par	name of the parameter
lookup_table	lookup table of the players
cluster_lookup_table	a lookup table of the clusters

### Value

a data. frame where we change the names in the variable colum to the corresponding parameter\_name from the lookup table

---

```
create_bpc_object
```

*Defines the class bpc and creates the bpc object. To create we need to receive some defined parameters (the arguments from the bpc function), a lookup table and a the stanfit object generated from the rstan sampling procedure*

---

### Description

Defines the class bpc and creates the bpc object. To create we need to receive some defined parameters (the arguments from the bpc function), a lookup table and a the stanfit object generated from the rstan sampling procedure

**Usage**

```
create_bpc_object(
  stanfit,
  lookup_table,
  model_type,
  standata,
  call_arg,
  cluster_lookup_table = NULL,
  predictors_df = NULL,
  predictors_lookup_table = NULL,
  predictors_matrix = NULL
)
```

**Arguments**

stanfit	Stanfit object returned by rstan::sampling
lookup_table	lookup_table dataframe. Two columns one Index the other Names where each each index will match a string in the names
model_type	the type of the model used to call stan (string)
standata	a list with the data used to call the rstan::sampling procedure
call_arg	a list with the arguments called from the bpc function
cluster_lookup_table	a lookup table with we have random effects
predictors_df	the data frame of the predictors for a generalized model
predictors_lookup_table	a lookup table for generalized models
predictors_matrix	a matrix of predictors for generalized models

**Value**

a bpc object

---

create\_cluster\_index *Create two columns with the indexes for the names of the players Here we create a new lookup table. Should be used when sampling the parameters*

---

**Description**

Create two columns with the indexes for the names of the players Here we create a new lookup table. Should be used when sampling the parameters

**Usage**

```
create_cluster_index(d, cluster)
```

**Arguments**

d	A data frame containing the observations. The other parameters specify the name of the columns
cluster	The name of the column of data data contains player0

**Value**

A dataframe with the additional columns 'cluster\_index'

---

```
create_cluster_index_with_existing_lookup_table
```

*Create two columns with the indexes for the names Here we use an existing lookup table. Should be used in predicting*

---

**Description**

Create two columns with the indexes for the names Here we use an existing lookup table. Should be used in predicting

**Usage**

```
create_cluster_index_with_existing_lookup_table(
  d,
  cluster,
  cluster_lookup_table
)
```

**Arguments**

d	A data frame containing the observations. The other parameters specify the name of the columns
cluster	The name of the column of data data contains player0
cluster_lookup_table	a lookup table for the cluster

**Value**

A dataframe with the additional columns 'player0\_index' and 'player1\_index' that contains the indexes

---

create_index	<i>Create two columns with the indexes for the names of the players Here we create a new lookup table. Should be used when sampling the parameters</i>
--------------	--

---

**Description**

Create two columns with the indexes for the names of the players Here we create a new lookup table. Should be used when sampling the parameters

**Usage**

```
create_index(d, player0, player1)
```

**Arguments**

d	A data frame containing the observations. The other parameters specify the name of the columns
player0	The name of the column of data data contains player0
player1	The name of the column of data data contains player0

**Value**

A dataframe with the additional columns 'player0\_index' and 'player1\_index' that contains the indexes

---

create_index_cluster_lookuptable	<i>Create a lookup table of names and indexes Note that the indexes will be created in the order they appear. For string this does not make much difference but for numbers the index might be different than the actual number that appears in names</i>
----------------------------------	---

---

**Description**

Create a lookup table of names and indexes Note that the indexes will be created in the order they appear. For string this does not make much difference but for numbers the index might be different than the actual number that appears in names

**Usage**

```
create_index_cluster_lookuptable(d, cluster)
```

**Arguments**

d	A data frame containing the observations. The other parameters specify the name of the columns
cluster	A string with the name of the cluster variable

**Value**

A dataframe of a lookup table with columns Names and Index

---

**create\_index\_lookupable**

*Create a lookup table of names and indexes Note that the indexes will be created in the order they appear. For string this doesnt make much difference but for numbers the index might be different than the actual number that appears in names*

---

**Description**

Create a lookup table of names and indexes Note that the indexes will be created in the order they appear. For string this doesnt make much difference but for numbers the index might be different than the actual number that appears in names

**Usage**

```
create_index_lookupable(d, player0, player1)
```

**Arguments**

d	A data frame containing the observations. The other parameters specify the name of the columns
player0	The name of the column of data contains player0
player1	The name of the column of data contains player0

**Value**

A dataframe of a lookup table with columns Names and Index



---

```
create_index_predictors_with_lookup_table
```

*Receives one column with player names and returns a data frame with the relevant index columns based on a given lookup table To be used with the predictors data frame*

---

### Description

Receives one column with player names and returns a data frame with the relevant index columns based on a given lookup table To be used with the predictors data frame

### Usage

```
create_index_predictors_with_lookup_table(d, player, lookup_table)
```

### Arguments

d	a data frame of the predictors
player	The name of the column of data data contains the player
lookup_table	a lookup table data frame

### Value

A dataframe with the additional column 'player\_index'

---

```
create_index_with_existing_lookup_table
```

*Create two columns with the indexes for the names Here we use an existing lookup table. Should be used in predicting*

---

### Description

Create two columns with the indexes for the names Here we use an existing lookup table. Should be used in predicting

### Usage

```
create_index_with_existing_lookup_table(d, player0, player1, lookup_table)
```

### Arguments

d	A data frame containing the observations. The other parameters specify the name of the columns
player0	The name of the column of data data contains player0
player1	The name of the column of data data contains player0
lookup_table	lookup_table a lookup table data frame

**Value**

A dataframe with the additional columns 'player0\_index' and 'player1\_index' that contains the indexes

---

create\_predictors\_lookup\_table

*Receives a vector with predictors strings (the column names) and returns a predictor\_lookup\_table*

---

**Description**

Receives a vector with predictors strings (the column names) and returns a predictor\_lookup\_table

**Usage**

```
create_predictors_lookup_table(predictors_columns)
```

**Arguments**

predictors\_columns

a vector with strings containing the columns for the predictors

**Value**

A matrix to be used in stan

---

create\_predictor\_matrix\_with\_player\_lookup\_table

*Receives a predictor dataframe, a string with the column of the player, a vector of strings with the columns for the predictors and a lookup table and returns an ordered matrix for Stan To be used with the predictors data frame*

---

**Description**

Receives a predictor dataframe, a string with the column of the player, a vector of strings with the columns for the predictors and a lookup table and returns an ordered matrix for Stan To be used with the predictors data frame

**Usage**

```
create_predictor_matrix_with_player_lookup_table(  
  d,  
  player,  
  predictors_columns,  
  lookup_table  
)
```

**Arguments**

d	a data frame of the predictors
player	The name of the column of data data contains the player
predictors_columns	a vector with strings containing the columns for the predictors
lookup_table	a lookup table data frame

**Value**

A matrix to be used in stan

---

expand\_aggregated\_data

*Expand aggregated data Several datasets for the Bradley-Terry Model aggregate the number of wins for each player in a different column. The models we provide are intended to be used in a long format. A single result for each contest. This function expands datasets that have aggregated data into this long format.*

---

**Description**

Expand aggregated data Several datasets for the Bradley-Terry Model aggregate the number of wins for each player in a different column. The models we provide are intended to be used in a long format. A single result for each contest. This function expands datasets that have aggregated data into this long format.

**Usage**

```
expand_aggregated_data(d, player0, player1, wins0, wins1, keep)
```

**Arguments**

d	a data frame
player0	string with column name of player0
player1	string with column name of player1
wins0	string with column name of the number of wins of player 0
wins1	string with column name of the number of wins of player 1
keep	an array of strings with the name of columns we want to keep in the new data frame (and repeat in every expanded row)

**Value**

a data frame with the expanded dataset. It will have the columns player1, player0, y, the keep columns, and a rowid column (to make each row unique)

## Examples

```
#Creating a simple data frame with only one row to illustrate how the function works
df1 <- tibble::tribble(~player0, ~player1, ~wins0, ~wins1, ~cluster, 'A', 'B', 4, 3, 'c1')
df2 <- expand_aggregated_data(df1, 'player0', 'player1', 'wins0', 'wins1', keep=c('cluster'))
print(df2)
```

---

get\_hpdi\_parameters     *Return the mean and the HPDI of the parameters of the model*

---

## Description

Return a data frame with the mean and with high and low 95% hpd interval for all parameters of the model

## Usage

```
get_hpdi_parameters(bpc_object)
```

## Arguments

bpc\_object     a bpc object

## Value

a data frame containing a column with the parameters, a column with mean and two columns with higher and lower hpdi

## Examples

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
hpdi<-get_hpdi_parameters(m)
print(hpdi)
```

---

get_loo	<i>Tiny wrapper for the PSIS-LOO-CV method from the loo package.</i>
---------	--

---

**Description**

This is used to evaluate the fit of the model using entropy criteria

**Usage**

```
get_loo(bpc_object)
```

**Arguments**

bpc\_object      a bpc object

**Value**

a loo object

**References**

Vehtari A, Gelman A, Gabry J (2017). Practical Bayesian model evaluation using leave-one-out cross-validation and WAIC. *Statistics and Computing*, 27, 1413-1432

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
l<-get_loo(m)
print(l)
```

---

get_model_parameters	<i>Return all the name of parameters in a model from a bpc_object. Here we exclude the log_lik and the lp__ since they are not parameters of the model</i>
----------------------	--

---

**Description**

Return all the name of parameters in a model from a bpc\_object. Here we exclude the log\_lik and the lp\_\_ since they are not parameters of the model

**Usage**

```
get_model_parameters(bpc_object)
```

**Arguments**

bpc\_object      a bpc object

**Value**

a vector with the name of the parameters

---

get_probabilities	<i>Get the empirical win/draw probabilities based on the ability/strength parameters. Instead of calculating from the probability formula given from the model we create a predictive posterior distribution for all pair combinations and calculate the posterior wins/loose/draw The function returns the mean value of win/loose/draw for the player i. To calculate for player j the probability is <math>1-p_i</math></i>
-------------------	--

---

**Description**

Get the empirical win/draw probabilities based on the ability/strength parameters. Instead of calculating from the probability formula given from the model we create a predictive posterior distribution for all pair combinations and calculate the posterior wins/loose/draw The function returns the mean value of win/loose/draw for the player i. To calculate for player j the probability is  $1-p_i$

**Usage**

```
get_probabilities(bpc_object, n = 1000)
```

**Arguments**

bpc\_object      a bpc object  
n                number of samples to draw from the posterior

**Value**

a list with data frame table with the respective probabilities and a matrix with the corresponding posterior

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
prob<-get_probabilities(m)
print(prob$Table)
```

---

get_rank_of_players	<i>Generate a ranking of the ability based on sampling the posterior distribution of the ranks.</i>
---------------------	---

---

**Description**

To print this object you should remove the last column PosteriorRank since it contain the whole posterior distribution for each case

**Usage**

```
get_rank_of_players(bpc_object, n = 1000)
```

**Arguments**

bpc_object	a bpc object
n	Number of times we will sample the posterior

**Value**

a data frame. This data frame contains the median of the rank, the mean, the standard deviation and column with a list containing all the posterior values for the rank

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
rank_m<-get_rank_of_players(m,n=100)
rank_table <- dplyr::select(rank_m,-MeanRank, -StdRank, -PosteriorRank)
print(rank_table)
```

---

get\_sample\_posterior *Get the posterior samples for a parameter of the model.*

---

### Description

Return a data frame with the posterior samples for the parameters of the model

### Usage

```
get_sample_posterior(bpc_object, par = "lambda", n = 1000)
```

### Arguments

bpc_object	a bpc object
par	name of the parameters to predict
n	how many times are we sampling? Default 1000

### Value

Return a data frame with the posterior samples for the parameters. One column for each parameter one row for each sample

### Examples

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
s <- get_sample_posterior(m, par='lambda', n=100)
print(head(s))
```

---

get\_stanfit *Retrieve the stanfit object generated by rstan.*

---

### Description

This object can be used with any other function or package that uses stanfit objects from rstan

### Usage

```
get_stanfit(bpc_object)
```



**Arguments**

bpc\_object      a bpc object

**Value**

a stanfit object

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
stanfit<- get_stanfit(m)
print(class(stanfit))
```

---

get\_stanfit\_summary      *Get stanfit summary table of all parameters excluding log\_lik.*

---

**Description**

Important to investigate the neff and the Rhat from the MCMC This excludes the log\_lik paramter

**Usage**

```
get_stanfit_summary(bpc_object)
```

**Arguments**

bpc\_object      a bpc object

**Value**

a data frame with the summary including quantiles, Rhat and neff

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
s <- get_stanfit_summary(m)
```

```
print(s)
```

---

get\_waic

*Tiny wrapper for the WAIC method from the loo package.*

---

### Description

This is used to evaluate the fit of the model using the Watanabe-Akaike Information criteria

### Usage

```
get_waic(bpc_object)
```

### Arguments

bpc\_object      a bpc object

### Value

a loo object

### References

Gelman, Andrew, Jessica Hwang, and Aki Vehtari. Understanding predictive information criteria for Bayesian models. *Statistics and computing* 24.6 (2014): 997-1016.

### Examples

```
m<-bpc(data = tennis_agresti,  
player0 = 'player0',  
player1 = 'player1',  
result_column = 'y',  
model_type = 'bt',  
solve_ties = 'none')  
waic<-get_waic(m)  
print(waic)
```

---

HPDI_from_stanfit	<i>Calculate HPDI for all parameters from a stanfit object Here we use the coda package</i>
-------------------	---

---

**Description**

Calculate HPDI for all parameters from a stanfit object Here we use the coda package

**Usage**

```
HPDI_from_stanfit(stanfit)
```

**Arguments**

stanfit            a stanfit object retrived from a bpc object

**Value**

a data frame with the HPDI calculated from the coda package

**References**

Martyn Plummer, Nicky Best, Kate Cowles and Karen Vines (2006). CODA: Convergence Diagnosis and Output Analysis for MCMC, R News, vol 6, 7-11

---

HPD_higher_from_column	<i>Returns the higher value of the HPD interval for a data frame column</i>
------------------------	---

---

**Description**

Returns the higher value of the HPD interval for a data frame column

**Usage**

```
HPD_higher_from_column(column, credMass = 0.95)
```

**Arguments**

column            the data to calculate the HPDI  
credMass          Credibility mass for the interval (area contained in the interval)

**Value**

the value of the higher HPD interval for that column

**References**

Mike Meredith and John Kruschke (2020). HDInterval: Highest (Posterior) Density Intervals. R package version 0.2.2. <https://CRAN.R-project.org/package=HDInterval>

---

HPD\_lower\_from\_column *Returns the lower value of the HPD interval for a data frame column*

---

**Description**

Returns the lower value of the HPD interval for a data frame column

**Usage**

```
HPD_lower_from_column(column, credMass = 0.95)
```

**Arguments**

column	the data to calculate the HPDI
credMass	Credibility mass for the interval (area contained in the interval)

**Value**

the value of the lower HPD interval for that column

**References**

Mike Meredith and John Kruschke (2020). HDInterval: Highest (Posterior) Density Intervals. R package version 0.2.2. <https://CRAN.R-project.org/package=HDInterval>

---

inv\_logit *Inverse logit function*

---

**Description**

Inverse logit function

**Usage**

```
inv_logit(x)
```

**Arguments**

x	is a real -inf to inf
---	-----------------------

**Value**

a value between 0 and 1

**References**

<https://en.wikipedia.org/wiki/Logit>

**Examples**

```
inv_logit(5)
inv_logit(-5)
inv_logit(0)
```

---

launch_shinystan	<i>Tiny wrapper to launch a shinystan app to investigate the MCMC.</i>
------------------	--

---

**Description**

It launches a shinystan app automatically in the web browser

**Usage**

```
launch_shinystan(bpc_object)
```

**Arguments**

bpc\_object      a bpc object

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
launch_shinystan(m)
```

---

logit	<i>Logit function</i>
-------	-----------------------

---

**Description**

Logit function

**Usage**

logit(x)

**Arguments**

x                    p is a probability 0 to 1

**Value**

a value between -inf and inf

**References**

<https://en.wikipedia.org/wiki/Logit>

**Examples**

```
logit(0.5)
logit(0.2)
```

---

match\_cluster\_names\_to\_cluster\_lookup\_table

*Receives a column with cluster names and returns a data frame with the relevant index column based on a given cluster lookup table*

---

**Description**

Receives a column with cluster names and returns a data frame with the relevant index column based on a given cluster lookup table

**Usage**

```
match_cluster_names_to_cluster_lookup_table(d, cluster, cluster_lookup_table)
```

**Arguments**

d                    a data frame  
cluster              The name of the column of data data contains player0  
cluster\_lookup\_table    a lookup table for the cluster

**Value**

A dataframe with the additional columns 'cluster\_index' that contains the indexes

---

match\_player\_names\_to\_lookup\_table

*Receives two columns with player names and returns a data frame with the relevant index columns based on a given lookup table*

---

**Description**

Receives two columns with player names and returns a data frame with the relevant index columns based on a given lookup table

**Usage**

```
match_player_names_to_lookup_table(d, player0, player1, lookup_table)
```

**Arguments**

d	a data frame
player0	The name of the column of data data contains player0
player1	The name of the column of data data contains player1
lookup_table	a lookup table data frame

**Value**

A dataframe with the additional columns 'player0\_index' and 'player1\_index' that contains the indexes

---

optimization\_algorithms

*Dataset containing an example of the performance of different optimization algorithms against different benchmark functions. This is a reduced version of the dataset presented at the paper: "Statistical Models for the Analysis of Optimization Algorithms with Benchmark Functions.". For details on how the data was collected we refer to the paper.*

---

**Description**

Dataset containing an example of the performance of different optimization algorithms against different benchmark functions. This is a reduced version of the dataset presented at the paper: "Statistical Models for the Analysis of Optimization Algorithms with Benchmark Functions.". For details on how the data was collected we refer to the paper.

**Usage**

```
optimization_algorithms
```

**Format**

This is the expansion of the data where each row contains 1 match only

- Algorithm: name of algorithm
- Benchmark: name of the benchmark problem
- TrueRewardDifference: Difference between the minimum function value obtained by the algorithm and the known global minimum
- Ndimensions: Number of dimensions of the benchmark problem
- MaxFevalPerDimensions: Maximum allowed budget for the algorithm per dimensions of the benchmark problem
- simNumber: id of the simulation. Indicates the repeated measures of each algorithm in each benchmark

**Source**

Mattos, David Issa, Jan Bosch, and Helena Holmstrom Olsson. Statistical Models for the Analysis of Optimization Algorithms with Benchmark Functions. arXiv preprint arXiv:2010.03783 (2020).

---

```
predict.bpc
```

```
Predict results for new data.
```

---

**Description**

This S3 function receives the bpc model and a data frame containing the same columns as the one used to fit the model. It returns another data frame with with the same columns of the new data and n additional columns representing a posterior predictive distribution. See the vignettes for a larger examples with the usage of this function

**Usage**

```
## S3 method for class 'bpc'
predict(object, newdata, predictors = NULL, n = 100, return_matrix = F, ...)
```

**Arguments**

object	a bpc object
newdata	a data frame that contains columns with the same names as used to fit the data in the model.



predictors	A data frame that contains the players predictors values when using a generalized model. Should be set only if using the generalized models. Only numeric values are accepted. Booleans are accepted but will be cast into integers. The first column should be for the player name, the others will be the predictors. The column names will be used as name for the predictors
n	number of time we will iterate and get the posterior. default is 100 so we dont get too many
return_matrix	should we return only a matrix with the predictive values. Default F. Use this to combine with predictive posterior plots in bayesplot This parameter also ignores the n parameter above since it passes all the predictions from stan
...	additional parameters for the generic print function

**Value**

a dataframe or a matrix depending on the return\_matrix parameter

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
predict(m,newdata=tennis_agresti)
```

---

print.bpc	<i>Print method for the bpc object.</i>
-----------	---

---

**Description**

This S3 functions only prints the mean and the HDPI values of all the parameters in the model

**Usage**

```
## S3 method for class 'bpc'
print(x, digits = 3, ...)
```

**Arguments**

x	a bpc object
digits	number of decimal digits in the table
...	additional parameters for the generic print function

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
#' print(m)
```

---

```
replace_parameter_index_with_names
```

*Replace the name of the parameter from index to name using a lookup\_table Receives a data frame and returns a dataframe.*

---

**Description**

Replace the name of the parameter from index to name using a lookup\_table Receives a data frame and returns a dataframe.

**Usage**

```
replace_parameter_index_with_names(
  d,
  column,
  par,
  lookup_table,
  cluster_lookup_table = NULL,
  predictors_lookup_table = NULL
)
```

**Arguments**

d	dataframe
column	name of the colum
par	name of the parameter
lookup_table	lookup table of the players
cluster_lookup_table	a lookup table of the predictors
predictors_lookup_table	a lookup table for the predictors

**Value**

a data. frame where we change the names in the variable column to the corresponding parameter\_name from the lookup table

---

sample_stanfit	<i>Return a data frame by resampling the posterior from a stanfit Here we select a parameter, retrieve the all the posterior from the stanfit and then we resample this posterior n times</i>
----------------	---

---

### Description

Return a data frame by resampling the posterior from a stanfit Here we select a parameter, retrieve the all the posterior from the stanfit and then we resample this posterior n times

### Usage

```
sample_stanfit(stanfit, par, n = 100)
```

### Arguments

stanfit	stanfit object
par	parameter name
n	number of samples

### Value

a dataframe containing the samples of the parameter. Each column is a parameter (in order of the index), each row is a sample

### References

Stan Development Team (2020). RStan: the R interface to Stan. R package version 2.21.2. <http://mc-stan.org/>.

---

summary.bpc	<i>Summary of the model bpc model.</i>
-------------	--

---

### Description

- Table 1: Contains the parameter estimates and respective HPD interval
- Table 2: Contains the posterior probability for the combination of all players
- Table 3: Contains the ranking of the players' abilities based on the posterior distribution of the ranks

### Usage

```
## S3 method for class 'bpc'
summary(object, digits = 2, ...)
```

**Arguments**

object	bpc object
digits	number of decimal digits in the table
...	additional parameters for the generic summary function

**Examples**

```
m<-bpc(data = tennis_agresti,
player0 = 'player0',
player1 = 'player1',
result_column = 'y',
model_type = 'bt',
solve_ties = 'none')
summary(m)
```

---

tennis_agresti	<i>This is the expansion of the tennis data from Agresti (2003) p.449 This data refers to matches for several women tennis players during 1989 and 1990</i>
----------------	---

---

**Description**

This is the expansion of the tennis data from Agresti (2003) p.449 This data refers to matches for several women tennis players during 1989 and 1990

**Usage**

```
tennis_agresti
```

**Format**

This is the expansion of the data where each row contains 1 match only

- player0: name of player0
- player1: name of player1
- y: corresponds to the result of the match: 0 if player0 won, 1 if player1 won.
- id: is a column to make each row unique in the data. It does not have any particular interpretation

**Source**

Agresti, Alan. Categorical data analysis. Vol. 482. John Wiley & Sons, 2003.

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