

# Package ‘binaryRL’

May 13, 2025

**Version** 0.8.0

**Title** Reinforcement Learning Tools for Two-Alternative Forced Choice Tasks

**Description** Tools for building reinforcement learning (RL) models specifically tailored for Two-Alternative Forced Choice (TAFC) tasks, commonly employed in psychological research. These models build upon the foundational principles of model-free reinforcement learning detailed in Sutton and Barto (1998) <ISBN:0262039249>. The package allows for the intuitive definition of RL models using simple if-else statements. Our approach to constructing and evaluating these computational models is informed by the guidelines proposed in Wilson & Collins (2019) <doi:10.7554/eLife.49547>. Example datasets included with the package are sourced from the work of Mason et al. (2024) <doi:10.3758/s13423-023-02415-x>.

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**URL** <https://github.com/yuki-961004/binaryRL>

**BugReports** <https://github.com/yuki-961004/binaryRL/issues>

**License** GPL-3

**Encoding** UTF-8

**LazyData** TRUE

**RoxygenNote** 7.3.2

**Depends** R (>= 4.0.0)

**Imports** future, doFuture, foreach, doRNG, progressr

**Suggests** stats, GenSA, GA, DEoptim, mlrMBO, mlr, ParamHelpers, smooof, lhs, pso, cmaes

**NeedsCompilation** no

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add_NA	<i>Create NULL columns and the line 0</i>
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### Description

Create NULL columns and the line 0

### Usage

```
add_NA(data)
```

### Arguments

data	[data.frame] A data frame resulting from the 'step2' process of the 'arrange_data' function.
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**Value**

data frame:

- data: step2 + row[0] for initial value + null cols [Reward, gamma, R\_utility, ...].

---

arrange_data	<i>Arrange Data based on Block and Trial</i>
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---

**Description**

Arrange Data based on Block and Trial

**Usage**

```
arrange_data(data, time_line = c("Block", "Trial"))
```

**Arguments**

data	[data.frame] A data frame resulting from the 'step1' process of the 'unique_choice' function.
time_line	[vector] A vector specifying the name of the column that the sequence of the experiment. This argument defines how the experiment is structured, such as whether it is organized by "Block" with breaks in between, and multiple trials within each block. e.g., 'time_line = c("Block", "Trial")'

**Value**

data frame:

- data: step1 arranged by 'time\_line'.

---

check_dependency	<i>Check Package Dependencies</i>
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**Description**

Checks if one or more specified R packages are installed and available. If any package is missing, it stops execution with an informative error message guiding the user on how to install them.

**Usage**

```
check_dependency(pkg_names, algorithm_name)
```

**Arguments**

`pkg_names` [vector] A character vector containing the names of the packages to check.

`algorithm_name` [character] Optional: The name of the algorithm or feature that requires these packages. Used to make the error message more specific. Defaults to "this functionality".

**Value**

Returns 'invisible(TRUE)' if all specified packages are available, otherwise it stops the execution via 'stop()'.

---

decision_making	<i>Markov Decision Process</i>
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**Description**

Markov Decision Process

**Usage**

```
decision_making(  
  data,  
  options,  
  L_choice = "L_choice",  
  R_choice = "R_choice",  
  L_reward = "L_reward",  
  R_reward = "R_reward",  
  var1 = NA,  
  var2 = NA,  
  seed = 123,  
  initial_value,  
  softmax = TRUE,  
  threshold = 1,  
  gamma,  
  eta,  
  epsilon,  
  tau,  
  lambda,  
  expl_func = func_epsilon,  
  prob_func = func_tau,  
  util_func = func_gamma,  
  rate_func = func_eta  
)
```

**Arguments**

data	[data.frame] A data frame resulting from the 'step4' process of the 'set_initial_value' function.
options	[vector] all alternative options from 'step1' 'unique_choice'
L_choice	[character] column name of left choice. e.g., 'L_choice = "Left_Choice"'
R_choice	[character] column name of right choice. e.g., 'R_choice = "Right_Choice"'
L_reward	[character] column name of the reward of left choice e.g., 'L_reward = "Left_reward"'
R_reward	[character] column name of the reward of right choice e.g., 'R_reward = "Right_reward"'
var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'
var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., 'var2 = "Extra_Var2"'
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
initial_value	[numeric] subject's initial expected value for each stimulus's reward. If this value is not set ('initial_value = NA'), the subject will use the reward received after the first trial as the initial value for that stimulus. In other words, the learning rate for the first trial is 100 e.g., 'initial_value = 0'
softmax	[logical] whether to use the softmax function. When softmax = TRUE, the value of each option influences the probability of selecting that option. Higher values increase the probability of selecting that option. When softmax = FALSE, the subject will always choose the option with the higher value, with no possibility of selecting the lower-value option. default: 'softmax = TRUE'
threshold	[integer] the number of initial trials during which the subject makes random choices rather than choosing based on the values of the options. This occurs because the subject has not yet learned the values of the options. For example, threshold = 20 means the subject will make completely random choices for the first 20 trials. default: 'threshold = 1'
gamma	[vector] Parameters used in the 'util_func' (Utility Function), often referred to as the discount rate. For example, 'utility = gamma * reward', if gamma < 1, it indicates that people tend to discount the objective reward. Provide the value as a vector e.g., 'gamma = c(0.7)'
eta	[vector] Parameters used in the 'rate_func' (Learning Rate Function), representing the rate at which the subject updates the difference (prediction error) between the reward and the expected value in the subject's mind. In the TD model, there is a single learning rate throughout the experiment. In the RSTD model, two different learning rates are used when the reward is higher or lower than the expected value. e.g., 'eta = c(0.3, 0.7)'
epsilon	[vector] Parameters used in the 'expl_func' (Exploration Function), determining whether the subject makes decisions based on the relative values of the left and right options, or chooses completely randomly. For example, when epsilon =

	0.1, it means the subject has a 10 completely random choice and a 90 of the options. e.g., ‘epsilon = c(0.1)’
tau	[vector] Parameters used in the ‘prob_func’ (Soft-Max Function), representing the sensitivity of the subject to the value difference when making decisions. It determines the probability of selecting the left option versus the right option based on their values. A larger value of tau indicates greater sensitivity to the value difference between the options. In other words, even a small difference in value will make the subject more likely to choose the higher-value option. e.g., ‘tau = c(0.5)’
lambda	[vector] Extra parameters that may be used in functions. e.g., ‘lambda = c(0.4, 0.7, 20, 60)’
expl_func	[function] Exploration Function.
prob_func	[function] Soft-Max Function.
util_func	[function] Utility Function.
rate_func	[function] Learning Rate Function.

**Value**

data frame:

- data: step4 + all decisions.

---

digits

*Round Digital*

---

**Description**

Round Digital

**Usage**

```
digits(data, options, digits_1 = 2, digits_2 = 5)
```

**Arguments**

data	[data.frame] A data frame resulting from the ‘step6’ process of the ‘model_fit’ function.
options	[vector] all alternative options from ‘step1’ ‘unique_choice’
digits_1	[integer] The number of decimal places to retain for columns related to the value function The default is 2.
digits_2	[integer] The number of decimal places to retain for columns related to the select function. The default is 5.

**Value**

data frame:

- data: step6 + round(col, digits).

fit\_p

*Fit parameters***Description**

This function optimizes free parameters of reinforcement learning models built with the `'run_m'` function. After constructing a reinforcement learning model (a function with only ONE argument, `'params'`), the `'fit_p'` function searches for the optimal values of these free parameters.

The package provides four optimization algorithms:

1. L-BFGS-B (from `'stats::optim'`);
2. Simulated Annealing (`'GenSA::GenSA'`);
3. Genetic Algorithm (`'GA::ga'`);
4. Differential Evolution (`'DEoptim::DEoptim'`);
5. Bayesian Optimization (`'mlrMBO::mbo'`);
6. Particle Swarm Optimization (`'pso::psoptim'`);
7. Covariance Matrix Adapting Evolutionary Strategy (`'cmaes::cma_es'`);

For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

**Usage**

```
fit_p(
  data,
  id = c(1:40),
  n_trials,
  fit_model = list(TD, RSTD, Utility),
  funcs = NULL,
  model_name = c("TD", "RSTD", "Utility"),
  lower = list(c(0, 0), c(0, 0, 0), c(0, 0, 0)),
  upper = list(c(1, 1), c(1, 1, 1), c(1, 1, 1)),
  initial_params = NA,
  initial_size = 50,
  iteration = 10,
  seed = 123,
  nc = 1,
  algorithm
)
```

**Arguments**

<code>data</code>	[data.frame] raw data. This data should include the following mandatory columns: - "sub", "time_line", "L_choice", "R_choice", "L_reward", "R_reward".
<code>id</code>	[vector] which subject is going to be analyzed. is being analyzed. The value should correspond to an entry in the "sub" column, which must contain the subject IDs. e.g., <code>'id = c(1:40)'</code>
<code>n_trials</code>	[integer] number of total trials
<code>fit_model</code>	[list] A collection of functions applied to fit models to the data.
<code>funcs</code>	[vector] A character vector containing the names of all user-defined functions required for the computation.

model_name	[list] the name of fit modals
lower	[list] The lower bounds for model fit models
upper	[list] The upper bounds for model fit models
initial_params	[vector] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for 'L-BFGS-B', 'GenSA', set 'initial = c(0, 0, ...)'
initial_size	[integer] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for 'Bayesian', 'GA', set 'initial = 50'
iteration	[integer] the number of iteration
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
nc	[integer] Number of CPU cores to use for parallel computation.
algorithm	[character] Choose a algorithm package from 'L-BFGS-B', 'GenSA', 'GA', 'DEoptim', 'Bayesian', 'PSO', 'CMA-ES'

**Value**

binaryRL results for all subjects with all models

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func_epsilon	<i>Epsilon Greedy</i>
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**Description**

Epsilon Greedy

**Usage**

```
func_epsilon(i, var1 = NA, var2 = NA, threshold = 1, epsilon = NA, lambda)
```

**Arguments**

i	row number
var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'
var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., 'var2 = "Extra_Var2"'

threshold	[integer] the number of initial trials during which the subject makes random choices rather than choosing based on the values of the options. This occurs because the subject has not yet learned the values of the options. For example, threshold = 20 means the subject will make completely random choices for the first 20 trials. default: 'threshold = 1'
epsilon	[vector] Parameters used in the 'expl_func' (Exploration Function), determining whether the subject makes decisions based on the relative values of the left and right options, or chooses completely randomly. For example, when epsilon = 0.1, it means the subject has a 10 completely random choice and a 90 of the options. e.g., 'epsilon = c(0.1)'
lambda	[vector] Extra parameters that may be used in functions. e.g., 'lambda = c(0.4, 0.7, 20, 60)'

**Value**

explore or not

**Note**

When customizing these functions, please ensure that you do not modify the arguments. Instead, only modify the 'if-else' statements or the internal logic to adapt the function to your needs.

---

 func\_eta

*Learning Rate*


---

**Description**

Learning Rate

**Usage**

```
func_eta(value, utility, reward, occurrence, var1 = NA, var2 = NA, eta, lambda)
```

**Arguments**

value	The expected value of the stimulus in the subject's mind at this point in time.
utility	The subjective value that the subject assigns to the objective reward.
reward	The objective reward received by the subject after selecting a stimulus.
occurrence	The number of times the same stimulus has appeared.
var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'
var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., 'var2 = "Extra_Var2"'

eta	[vector] Parameters used in the ‘rate_func‘ (Learning Rate Function), representing the rate at which the subject updates the difference (prediction error) between the reward and the expected value in the subject’s mind. In the TD model, there is a single learning rate throughout the experiment. In the RSTD model, two different learning rates are used when the reward is higher or lower than the expected value. e.g., ‘eta = c(0.3, 0.7)’
lambda	[vector] Extra parameters that may be used in functions. e.g., ‘lambda = c(0.4, 0.7, 20, 60)’

**Value**

learning rate eta

**Note**

When customizing these functions, please ensure that you do not modify the arguments. Instead, only modify the ‘if-else‘ statements or the internal logic to adapt the function to your needs.

---

func\_gamma

*Utility Function*

---

**Description**

Utility Function

**Usage**

```
func_gamma(
  value,
  utility,
  reward,
  occurrence,
  var1 = NA,
  var2 = NA,
  gamma = 1,
  lambda
)
```

**Arguments**

value	The expected value of the stimulus in the subject’s mind at this point in time.
utility	The subjective value that the subject assigns to the objective reward.
reward	The objective reward received by the subject after selecting a stimulus.
occurrence	The number of times the same stimulus has appeared.

var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'
var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., 'var2 = "Extra_Var2"'
gamma	[vector] Parameters used in the 'util_func' (Utility Function), often referred to as the discount rate. For example, 'utility = gamma * reward', if gamma < 1, it indicates that people tend to discount the objective reward. Provide the value as a vector e.g., 'gamma = c(0.7)'
lambda	[vector] Extra parameters that may be used in functions. e.g., 'lambda = c(0.4, 0.7, 20, 60)'

**Value**

Discount rate and utility

**Note**

When customizing these functions, please ensure that you do not modify the arguments. Instead, only modify the 'if-else' statements or the internal logic to adapt the function to your needs.

---

 func\_tau

*Soft-Max Function*


---

**Description**

Soft-Max Function

**Usage**

```
func_tau(LR, try, L_value, R_value, var1 = NA, var2 = NA, tau = 1, lambda)
```

**Arguments**

LR	Are you calculating the probability for the left option or the right option?
try	If the choice was random, the value is 1; if the choice was based on value, the value is 0.
L_value	The value of the left option
R_value	The value of the right option
var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'

var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., ‘var2 = "Extra_Var2"‘
tau	[vector] Parameters used in the ‘prob_func‘ (Soft-Max Function), representing the sensitivity of the subject to the value difference when making decisions. It determines the probability of selecting the left option versus the right option based on their values. A larger value of tau indicates greater sensitivity to the value difference between the options. In other words, even a small difference in value will make the subject more likely to choose the higher-value option. e.g., ‘tau = c(0.5)‘
lambda	[vector] Extra parameters that may be used in functions. e.g., ‘lambda = c(0.4, 0.7, 20, 60)‘

### Value

The probability of choosing this option

---

Mason\_2024\_Exp1

*Experiment 1 from Mason et al. (2024)*

---

### Description

This dataset is from Experiment 1 of Mason et al. (2024). (Rare and extreme outcomes in risky choice). Data is publicly available on OSF: <https://osf.io/hy3q4/>. We performed basic cleaning to meet our package needs.

### Format

A data frame with 45000 rows and 11 columns:

**Subject** Subject ID, an integer (16 to 144)

**Block** Block number, an integer (1 to 6)

**Trial** Trial number, an integer (1 to 60)

**L\_choice** Left choice, A = 100% gain 4, B = 90% gain 0 and 10% gain 40, C = 100% lose 4, D = 90% lose 0 and 10% lose 40.

**R\_choice** Right choice, A = 100% gain 4, B = 90% gain 0 and 10% gain 40, C = 100% lose 4, D = 90% lose 0 and 10% lose 40.

**L\_reward** Reward associated with the left choice.

**R\_reward** Reward associated with the right choice.

**Sub\_Choose** The chosen option, either L\_choice or R\_choice.

**Frame** Type of frame, "Gain", "Loss", "Catch".

**NetWorth** The participant’s net worth at the end of each trial.

**RT** The participant’s reaction time (in milliseconds) for each trial.

### Examples

```
# Load the Mason_2024_Exp1 dataset
data(Mason_2024_Exp1)
head(Mason_2024_Exp1)
```

---

Mason_2024_Exp2	<i>Experiment 2 from Mason et al. (2024)</i>
-----------------	----------------------------------------------

---

### Description

This dataset is from Experiment 1 of Mason et al. (2024). (Rare and extreme outcomes in risky choice). Data is publicly available on OSF: <https://osf.io/hy3q4/>. We performed basic cleaning to meet our package needs.

### Format

A data frame with 45000 rows and 11 columns:

**Subject** Subject ID, an integer (16 to 144)

**Block** Block number, an integer (1 to 6)

**Trial** Trial number, an integer (1 to 60)

**L\_choice** Left choice, A = 100% gain 36, B = 90% gain 40 and 10% gain 0, C = 100% lose 36, D = 90% lose 40 and 10% lose 0.

**R\_choice** Right choice, A = 100% gain 36, B = 90% gain 40 and 10% gain 0, C = 100% lose 36, D = 90% lose 40 and 10% lose 0.

**L\_reward** Reward associated with the left choice.

**R\_reward** Reward associated with the right choice.

**Sub\_Choose** The chosen option, either L\_choice or R\_choice.

**Frame** Type of frame, "Gain", "Loss", "Catch".

**NetWorth** The participant's net worth at the end of each trial.

**RT** The participant's reaction time (in milliseconds) for each trial.

### Examples

```
# Load the Mason_2024_Exp2 dataset
data(Mason_2024_Exp2)
head(Mason_2024_Exp2)
```

---

mode	<i>Pretend to be Raw Data</i>
------	-------------------------------

---

### Description

Pretend to be Raw Data

### Usage

```
mode(
  data,
  mode = "fit",
  sub_choose = "Sub_Choose",
  rob_choose = "Rob_Choose",
  raw_cols = c("Subject", "Block", "Trial", "L_choice", "R_choice", "L_reward",
              "R_reward", "Sub_Choose")
)
```

### Arguments

data	[list] a list resulting from the 'step7' process of the 'output' function.
mode	[character] 'fit' or 'simulate' whether to generate raw data. Defaults to FALSE. Set to TRUE to generate fake data. This produces a data frame with the same format as the actual raw data.
sub_choose	[character] column name of choices made by the subject. e.g., 'sub_choose = "Choose"'
rob_choose	[character] column name of choices made by the model. e.g., 'rob_choose = "Rob_Choose"'. you should ignore this argument
raw_cols	[vector] default: c("Subject", "Block", "Trial", "L_choice", "R_choice", "L_reward", "R_reward", "Choose", "Reward") These are the column names of the raw data. Only required when 'back = 'simulate' '.

### Value

binaryRL[list]:

- data: new raw data (decision made by robot)
- params: all parameters value
- numeric: ACC
- numeric: LogL
- numeric: AIC
- numeric: BIC

---

model_fit	<i>Calculate the Model Fit</i>
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---

### Description

Calculate the Model Fit

### Usage

```
model_fit(
  data,
  L_choice = "L_choice",
  R_choice = "R_choice",
  sub_choose = "Sub_Choose"
)
```

### Arguments

data	[data.frame] A data frame resulting from the 'step5' process of the 'decision_making' function.
L_choice	[character] column name of left choice. e.g., 'L_choice = "Left_Choice"'
R_choice	[character] column name of right choice. e.g., 'R_choice = "Right_Choice"'
sub_choose	[character] column name of choices made by the subject. e.g., 'sub_choose = "Choose"'

### Value

data frame:

- data: step5 + ACC + logL.

---

optimize_para	<i>Fit Parameters</i>
---------------	-----------------------

---

### Description

This function is an internal function of 'fit\_p'. We isolate it from direct use by capable users.

The function provides four optimization algorithms:

1. L-BFGS-B (from 'stats::optim');
2. Simulated Annealing ('GenSA::GenSA');
3. Genetic Algorithm ('GA::ga');
4. Differential Evolution ('DEoptim::DEoptim');
5. Bayesian Optimization ('mlrMBO::mbo');
6. Particle Swarm Optimization ('pso::psoptim');
7. Covariance Matrix Adapting Evolutionary Strategy ('cmaes::cma\_es');

For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

**Usage**

```
optimize_para(
  data,
  id,
  obj_func,
  n_params,
  n_trials,
  lower,
  upper,
  initial_params = NA,
  initial_size = 50,
  iteration = 10,
  seed = 123,
  algorithm
)
```

**Arguments**

data	[data.frame] raw data. This data should include the following mandatory columns: - "sub", "time_line", "L_choice", "R_choice", "L_reward", "R_reward".
id	[integer] which subject is going to be analyzed. is being analyzed. The value should correspond to an entry in the "sub" column, which must contain the subject IDs. e.g., 'id = 18'
obj_func	[function] a function with only ONE argument 'params'. Additionally, it is important to note that the data needs to be retrieved from fit_env() and the results passed back to fit_env(). This function returns the log likelihood (logL).
n_params	[integer] The number of free parameters in your model.
n_trials	[integer] The total number of trials in your experiment.
lower	[vector] lower bounds of free parameters
upper	[vector] upper bounds of free parameters
initial_params	[vector] Initial values for the free parameters. automatically generate initial values. for 'L-BFGS-B', 'GenSA', set 'initial = c(0, 0, ...)'
initial_size	[integer] Initial population size for the free parameters. automatically generate initial values. for 'Bayesian', 'GA', set 'initial = 50'
iteration	[integer] the number of iteration
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
algorithm	[character] Choose a algorithm package from 'L-BFGS-B', 'GenSA', 'GA', 'DEoptim', 'Bayesian', 'PSO', 'CMA-ES'

**Value**

the result of binaryRL with optimal parameters

---

 output

*Summary the Results*


---

## Description

Summary the Results

## Usage

```
output(data, n_params, n_trials, gamma, eta, epsilon, tau, lambda)
```

## Arguments

data	[data.frame] A data frame resulting from the 'step7' process of the 'digits' function.
n_params	[integer] The number of free parameters in your model.
n_trials	[integer] The total number of trials in your experiment.
gamma	[vector] Parameters used in the 'util_func' (Utility Function), often referred to as the discount rate. For example, 'utility = gamma * reward', if gamma < 1, it indicates that people tend to discount the objective reward. Provide the value as a vector e.g., 'gamma = c(0.7)'
eta	[vector] Parameters used in the 'rate_func' (Learning Rate Function), representing the rate at which the subject updates the difference (prediction error) between the reward and the expected value in the subject's mind. In the TD model, there is a single learning rate throughout the experiment. In the RSTD model, two different learning rates are used when the reward is higher or lower than the expected value. e.g., 'eta = c(0.3, 0.7)'
epsilon	[vector] Parameters used in the 'expl_func' (Exploration Function), determining whether the subject makes decisions based on the relative values of the left and right options, or chooses completely randomly. For example, when epsilon = 0.1, it means the subject has a 10 completely random choice and a 90 of the options. e.g., 'epsilon = c(0.1)'
tau	[vector] Parameters used in the 'prob_func' (Soft-Max Function), representing the sensitivity of the subject to the value difference when making decisions. It determines the probability of selecting the left option versus the right option based on their values. A larger value of tau indicates greater sensitivity to the value difference between the options. In other words, even a small difference in value will make the subject more likely to choose the higher-value option. e.g., 'tau = c(0.5)'
lambda	[vector] Extra parameters that may be used in functions. e.g., 'lambda = c(0.4, 0.7, 20, 60)'

**Value**

binaryRL[list]:

- data: output data frame with all information
- params: all parameters value
- numeric: ACC
- numeric: LogL
- numeric: AIC
- numeric: BIC

rcv\_d

*Parameter and Model Recovery***Description**

This function fits multiple sets of simulated data using a loop. You need to provide a list of simulation functions, fitting functions, and parameter bounds. If you prefer to handle the process manually, you can use the internal functions ‘simulate\_list’ and ‘recovery\_data’.

For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

**Usage**

```
rcv_d(
  data,
  id = 1,
  n_trials = 288,
  simulate_models = list(TD, RSTD, Utility),
  simulate_lower = list(c(0, 0), c(0, 0, 0), c(0, 0, 0)),
  simulate_upper = list(c(1, 1), c(1, 1, 1), c(1, 1, 1)),
  fit_models = list(TD, RSTD, Utility),
  fit_lower = list(c(0, 0), c(0, 0, 0), c(0, 0, 0)),
  fit_upper = list(c(1, 1), c(1, 1, 1), c(1, 1, 1)),
  model_names = c("TD", "RSTD", "Utility"),
  funcs = NULL,
  initial_params = NA,
  initial_size = 50,
  iteration_s = 10,
  iteration_f = 10,
  seed = 1,
  nc = 1,
  algorithm
)
```

**Arguments**

data	[data.frame] raw data. This data should include the following mandatory columns: - "sub", "time_line", "L_choice", "R_choice", "L_reward", "R_reward".
id	[vector] which subject is going to be analyzed. is being analyzed. The value should correspond to an entry in the "sub" column, which must contain the subject IDs. e.g., 'id = c(1:40)'
n_trials	[integer] number of total trials
simulate_models	[list] A collection of functions used to generate simulated data.
simulate_lower	[list] The lower bounds for simulate models
simulate_upper	[list] The upper bounds for simulate models
fit_models	[list] A collection of functions applied to fit models to the data.
fit_lower	[list] The lower bounds for model fit models
fit_upper	[list] The upper bounds for model fit models
model_names	[list] the names of fit modals
funcs	[vector] A character vector containing the names of all user-defined functions required for the computation.
initial_params	[vector] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for 'L-BFGS-B', 'GenSA', set 'initial = c(0, 0, ...)'
initial_size	[integer] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for 'Bayesian', 'GA', set 'initial = 50'
iteration_s	[integer] the number of iteration in simulation (simulate)
iteration_f	[integer] the number of iteration in algorithm (fit)
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
nc	[integer] Number of CPU cores to use for parallel computation.
algorithm	[character] Choose a algorithm package from 'L-BFGS-B', 'GenSA', 'GA', 'DEoptim', 'Bayesian', 'PSO', 'CMA-ES'

**Value**

a list containing all recovery data

---

recovery_data	<i>recovery_d</i>
---------------	-------------------

---

### Description

This function applies ‘optimize\_para’ to each fake data in the list generated by ‘simulate\_list’. The results can be used for parameter recovery and model recovery, helping evaluate the consistency and validity of the reinforcement learning model.

For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

### Usage

```
recovery_data(
  list,
  id = 1,
  fit_model,
  funcs = NULL,
  model_name,
  n_params,
  n_trials,
  lower,
  upper,
  initial_params = NA,
  initial_size = 50,
  iteration = 10,
  seed = 123,
  nc = 1,
  algorithm
)
```

### Arguments

<code>list</code>	[list] a list generated by function ‘simulate_list’
<code>id</code>	[integer] default = 1
<code>fit_model</code>	[function] fit model
<code>funcs</code>	[vector] A character vector containing the names of all user-defined functions required for the computation.
<code>model_name</code>	[character] the name of your modal
<code>n_params</code>	[integer] The number of free parameters in your model.
<code>n_trials</code>	[integer] The total number of trials in your experiment.
<code>lower</code>	[vector] lower bounds of free parameters
<code>upper</code>	[vector] upper bounds of free parameters
<code>initial_params</code>	[vector] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for ‘L-BFGS-B’, ‘GenSA’, set ‘initial = c(0, 0, ...)’

initial_size	[integer] Initial values for the free parameters. These need to be set only when using L-BFGS-B. Other algorithms automatically generate initial values. for 'Bayesian', 'GA', set 'initial = 50'
iteration	[integer] the number of iteration
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
nc	[integer] Number of CPU cores to use for parallel computation.
algorithm	[character] Choose a algorithm package from 'L-BFGS-B', 'GA', 'GenSA', 'DEoptim'

**Value**

a data frame for parameter recovery and model recovery

---

rev_e	<i>Review Experimental Effect</i>
-------	-----------------------------------

---

**Description**

This function visualizes or extracts the model-estimated effect of reinforcement learning parameters for each subject.

**Usage**

```
rev_e(data, result, model, model_name, param_prefix, n_trials)
```

**Arguments**

data	[data.frame] Raw data. Must include the following columns: - "sub": Subject ID - "time_line": Trial index - "L_choice" / "R_choice": Left and right choice identifiers - "L_reward" / "R_reward": Left and right reward values
result	[data.frame] Output data generated by the 'fit_p()' function. Each row represents model fit results for a subject.
model	[function] A model function to be applied in evaluating the experimental effect.
model_name	[character] A character string specifying the name of the model to extract from the result.
param_prefix	[character] A prefix string used to identify parameter columns in the 'result' data (e.g., "param_").
n_trials	[integer] Number of total trials in the experimental task.

**Value**

A list where each element contains the extracted results for one subject.

---

RSTD	<i>RSTD model for fit</i>
------	---------------------------

---

**Description**

RSTD model for fit

**Usage**

RSTD(params)

**Arguments**

params            [vector] algorithm packages accept only one argument

**Value**

negative log likelihood

---

run_m	<i>Building Reinforcement Learning Model</i>
-------	----------------------------------------------

---

**Description**

This function requires the optimal parameter values obtained through the ‘algorithm’ package. Once the best parameter values are solved for, they are incorporated into the reinforcement learning model, allowing the model to simulate human-like decision-making. The function leverages these optimized parameters to generate choices that mimic the decision-making process of subjects, enabling the study of behavior under varying conditions. By integrating the best-fit parameters from the ‘algorithm’ package, this function offers a powerful tool for simulating human choices in reinforcement learning contexts.

For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

**Usage**

```
run_m(
  data,
  id,
  mode = "fit",
  initial_value = NA,
  softmax = TRUE,
  threshold = 1,
  seed = 123,
  n_params,
  n_trials,
```

```

gamma = 1,
eta,
epsilon = NA,
tau = 1,
lambda = NA,
util_func = func_gamma,
rate_func = func_eta,
expl_func = func_epsilon,
prob_func = func_tau,
sub = "Subject",
time_line = c("Block", "Trial"),
L_choice = "L_choice",
R_choice = "R_choice",
L_reward = "L_reward",
R_reward = "R_reward",
sub_choose = "Sub_Choose",
rob_choose = "Rob_Choose",
raw_cols = NULL,
var1 = NA,
var2 = NA,
digits_1 = 2,
digits_2 = 5
)

```

### Arguments

data	[data.frame] raw data. This data should include the following mandatory columns: - "sub", "time_line", "L_choice", "R_choice", "L_reward", "R_reward".
id	[integer] which subject is going to be analyzed. is being analyzed. The value should correspond to an entry in the "sub" column, which must contain the subject IDs. e.g., 'id = 18'
mode	[character] 'fit' or 'simulate' whether to generate raw data. Defaults to FALSE. Set to TRUE to generate fake data. This produces a data frame with the same format as the actual raw data.
initial_value	[numeric] subject's initial expected value for each stimulus's reward. If this value is not set ('initial_value = NA'), the subject will use the reward received after the first trial as the initial value for that stimulus. In other words, the learning rate for the first trial is 100 e.g., 'initial_value = 0'
softmax	[logical] whether to use the softmax function. When softmax = TRUE, the value of each option influences the probability of selecting that option. Higher values increase the probability of selecting that option. When softmax = FALSE, the subject will always choose the option with the higher value, with no possibility of selecting the lower-value option. default: 'softmax = TRUE'
threshold	[integer] the number of initial trials during which the subject makes random choices rather than choosing based on the values of the options. This occurs because the subject has not yet learned the values of the options. For example, threshold = 20 means the subject will make completely random choices for the first 20 trials. default: 'threshold = 1'

seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'
n_params	[integer] The number of free parameters in your model.
n_trials	[integer] The total number of trials in your experiment.
gamma	[vector] Parameters used in the 'util_func' (Utility Function), often referred to as the discount rate. For example, 'utility = gamma * reward', if gamma < 1, it indicates that people tend to discount the objective reward. Provide the value as a vector e.g., 'gamma = c(0.7)'
eta	[vector] Parameters used in the 'rate_func' (Learning Rate Function), representing the rate at which the subject updates the difference (prediction error) between the reward and the expected value in the subject's mind. In the TD model, there is a single learning rate throughout the experiment. In the RSTD model, two different learning rates are used when the reward is higher or lower than the expected value. e.g., 'eta = c(0.3, 0.7)'
epsilon	[vector] Parameters used in the 'expl_func' (Exploration Function), determining whether the subject makes decisions based on the relative values of the left and right options, or chooses completely randomly. For example, when epsilon = 0.1, it means the subject has a 10 completely random choice and a 90 of the options. e.g., 'epsilon = c(0.1)'
tau	[vector] Parameters used in the 'prob_func' (Soft-Max Function), representing the sensitivity of the subject to the value difference when making decisions. It determines the probability of selecting the left option versus the right option based on their values. A larger value of tau indicates greater sensitivity to the value difference between the options. In other words, even a small difference in value will make the subject more likely to choose the higher-value option. e.g., 'tau = c(0.5)'
lambda	[vector] Extra parameters that may be used in functions. e.g., 'lambda = c(0.4, 0.7, 20, 60)'
util_func	[function] Utility Function.
rate_func	[function] Learning Rate Function.
expl_func	[function] Exploration Function.
prob_func	[function] Soft-Max Function.
sub	[character] column name of subject ID e.g., 'sub = "Subject"'
time_line	[vector] A vector specifying the name of the column that the sequence of the experiment. This argument defines how the experiment is structured, such as whether it is organized by "Block" with breaks in between, and multiple trials within each block. e.g., 'time_line = c("Block", "Trial)'
L_choice	[character] column name of left choice. e.g., 'L_choice = "Left_Choice"'
R_choice	[character] column name of right choice. e.g., 'R_choice = "Right_Choice"'
L_reward	[character] column name of the reward of left choice e.g., 'L_reward = "Left_reward"'
R_reward	[character] column name of the reward of right choice e.g., 'R_reward = "Right_reward"'
sub_choose	[character] column name of choices made by the subject. e.g., 'sub_choose = "Choose"'

rob_choose	[character] column name of choices made by the model. e.g., 'rob_choose = "Rob_Choose"' you should ignore this argument
raw_cols	[vector] default: c("Subject", "Block", "Trial", "L_choice", "R_choice", "L_reward", "R_reward", "Choose", "Reward") These are the column names of the raw data. Only required when 'mode = "simulate"'
var1	[character] column name of extra variable 1. If your model uses more than just reward and expected value, and you need other information, such as whether the choice frame is Gain or Loss, then you can input the 'Frame' column as var1 into the model. e.g., 'var1 = "Extra_Var1"'
var2	[character] column name of extra variable 2. If one additional variable, var1, does not meet your needs, you can add another additional variable, var2, into your model. e.g., 'var2 = "Extra_Var2"'
digits_1	[integer] The number of decimal places to retain for columns related to the value function The default is 2.
digits_2	[integer] The number of decimal places to retain for columns related to the select function. The default is 5.

**Value**

the result of binaryRL with input parameters

**Examples**

```
data <- Mason_2024_Exp1

simulated <- binaryRL::run_m(
  data = data,
  id = 18,
  eta = c(0.321, 0.765),
  n_params = 2,
  n_trials = 360
)

summary(simulated)
```

---

set\_initial\_value      *Set initial values for all options*

---

**Description**

Set initial values for all options

**Usage**

```
set_initial_value(data, options, initial_value = NA)
```

**Arguments**

data	[data.frame] A data frame resulting from the 'step3' process of the 'add_NA' function.
options	[vector] all alternative options from 'step1' 'unique_choice'
initial_value	[numeric] subject's initial expected value for each stimulus's reward. If this value is not set ('initial_value = NA'), the subject will use the reward received after the first trial as the initial value for that stimulus. In other words, the learning rate for the first trial is 100 e.g., 'initial_value = 0'

**Value**

data frame:

- data: step3 + row[0] with initial value.

---

simulate\_list

*simulate\_l*

---

**Description**

This function generates simulated data using a user-defined objective function. You can specify the number of iterations to control how many data are generated. These datasets can be used for parameter recovery and model recovery. For more information, please refer to the GitHub repository: <https://github.com/yuki-961004/binaryRL>

**Usage**

```
simulate_list(
  data,
  id = 1,
  obj_func,
  n_params,
  n_trials,
  lower,
  upper,
  iteration = 10,
  seed = 123
)
```

**Arguments**

data	[data.frame] raw data. This data should include the following mandatory columns: - "sub", "time_line", "L_choice", "R_choice", "L_reward", "R_reward".
id	[vector] which subject is going to be analyzed. is being analyzed. The value should correspond to an entry in the "sub" column, which must contain the subject IDs. e.g., 'id = c(1:40)'

obj_func	[function] a function with only ONE argument 'params'. Additionally, it is important to note that the data needs to be retrieved from parent.frame(). This function returns the binaryRL.res(res).
n_params	[integer] The number of free parameters in your model.
n_trials	[integer] The total number of trials in your experiment.
lower	[vector] lower bounds of free parameters
upper	[vector] upper bounds of free parameters
iteration	[integer] the number of iteration
seed	[integer] random seed. This ensures that the results are reproducible and remain the same each time the function is run. default: 'seed = 123'

**Value**

a list with fake data generated by random free parameters

---

summary.binaryRL	<i>summary</i>
------------------	----------------

---

**Description**

summary

**Usage**

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

**Arguments**

object	binaryRL_res
...	others

**Value**

summary

---

TD	<i>TD model for fit</i>
----	-------------------------

---

**Description**

TD model for fit

**Usage**

TD(params)

**Arguments**

params            [vector] algorithm packages accept only one argument

**Value**

negative log likelihood

---

unique_choice	<i>Figure out how many options exist</i>
---------------	------------------------------------------

---

**Description**

Figure out how many options exist

**Usage**

unique\_choice(data, L\_choice = "L\_choice", R\_choice = "R\_choice")

**Arguments**

data            [data.frame] raw data. This data should include the following mandatory columns:  
- "sub", "time\_line", "L\_choice", "R\_choice", "L\_reward", "R\_reward".

L\_choice        [character] column name of left choice. e.g., 'L\_choice = "Left\_Choice"'

R\_choice        [character] column name of right choice. e.g., 'R\_choice = "Right\_Choice"'

**Value**

list:

- data: raw data frame + null cols [options].
- options: a vector containing all options.

---

Utility

*Utility model for fit*

---

**Description**

Utility model for fit

**Usage**

Utility(params)

**Arguments**

params            [vector] algorithm packages accept only one argument

**Value**

negative log likelihood

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