

Go-No Go Tutorial

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Go-No Go Task Tutorial

This tutorial will demonstrate how the *gonogo* package is used. The package contains two functions: `play_gonogo()` for playing the Go-No Go Task, and `check_rt()` for checking for irregularities in the output data, specifically the reaction time column.

Go-No Go Task in Short

The Go-No Go Task is a widely used test to measure inhibitory control, a cognitive process that enables humans to cancel motor activity after its initiation. It requires the participant to perform an action given certain stimuli (Go stimuli), and inhibit that action under a different set of stimuli (No Go stimuli).

There are two parameters in the experimental design that are especially important: the length of each trial and the relative proportion of the Go and No-Go trials. Fortunately, both these parameters can be easily manipulated in the `gonogo()` function: length of the trial with the *inter* argument, and the relative proportion of the Go and No-Go trials with the *prb* argument.

In addition to these two arguments, you can specify the participant *id* (name or unique id number), *n_trial* (number of trials), *n_block* (number of blocks), and *stimuli* (the Go and No Go stimuli).

How to Use the `play_gonogo()` Function

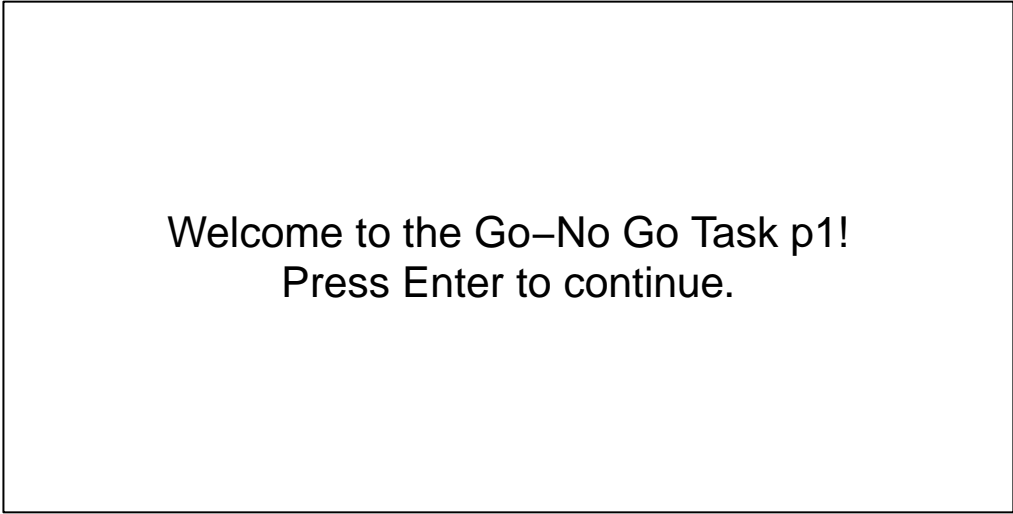
The following code gives an example of how the `play_gonogo()` function can be used.

```
# Load package
# devtools::install_github("Programming-The-Next-Step-2022/GoNoGoTask/gonogo")
library(gonogo)

# Run the Go-No Go Task and save the output in an object
p1_data <- play_gonogo(id = "p1", # id
  n_trial = 5, # number of trials
  n_block = 3, # number of blocks
  stimuli = c("A", "X"), # Go and No Go stimuli
  inter = 0.6, # interval (length of trials)
  prb = c(0.8, 0.2)) # relative proportion
```

Example Screens from the Task

Below you can find examples of how the task looks: the welcome screen, information about the number of trials and blocks, and finally an example of how the stimuli are presented during the task.



Welcome to the Go-No Go Task p1!
Press Enter to continue.

The task consists of 3 blocks, each consisting of 5 trials.
There is a break between each pair of blocks.
Before starting the task, there is one practice block of 10 trials.
Press Enter to start the practice trials.

A

What the Output Data Looks Like

##	id	response	correct	SDT	rt	stimulus	block
## 1	p1	none	1	correctrejection	NA	X	1
## 2	p1	none	1	correctrejection	NA	X	1
## 3	p1	space	1	hit	0.47666382	A	1
## 4	p1	space	1	hit	0.55146193	A	1
## 5	p1	none	0	miss	NA	A	1
## 6	p1	space	0	falsealarm	0.56809711	X	2
## 7	p1	none	0	miss	NA	A	2
## 8	p1	space	1	hit	-0.01646210	A	2
## 9	p1	space	1	hit	0.59251689	A	2
## 10	p1	none	1	correctrejection	NA	X	2
## 11	p1	none	0	miss	NA	A	3
## 12	p1	space	1	hit	-0.01505090	A	3
## 13	p1	none	0	miss	NA	A	3
## 14	p1	space	1	hit	-0.02203418	A	3
## 15	p1	none	0	miss	NA	A	3

How to Read the Output

The `play_gonogo()` function returns a dataframe consisting of `n_trial*n_block` (number of trials times number of blocks) rows and seven columns:

`id` = participant's name or id as specified

response = response key used on the trial (space when participant responded, none when no response was given)

correct = whether the response was correct or not (1=correct, 0=incorrect)

SDT = responses categorized according to Signal Detection Theory; [click here to read more about Signal Detection Theory](#)

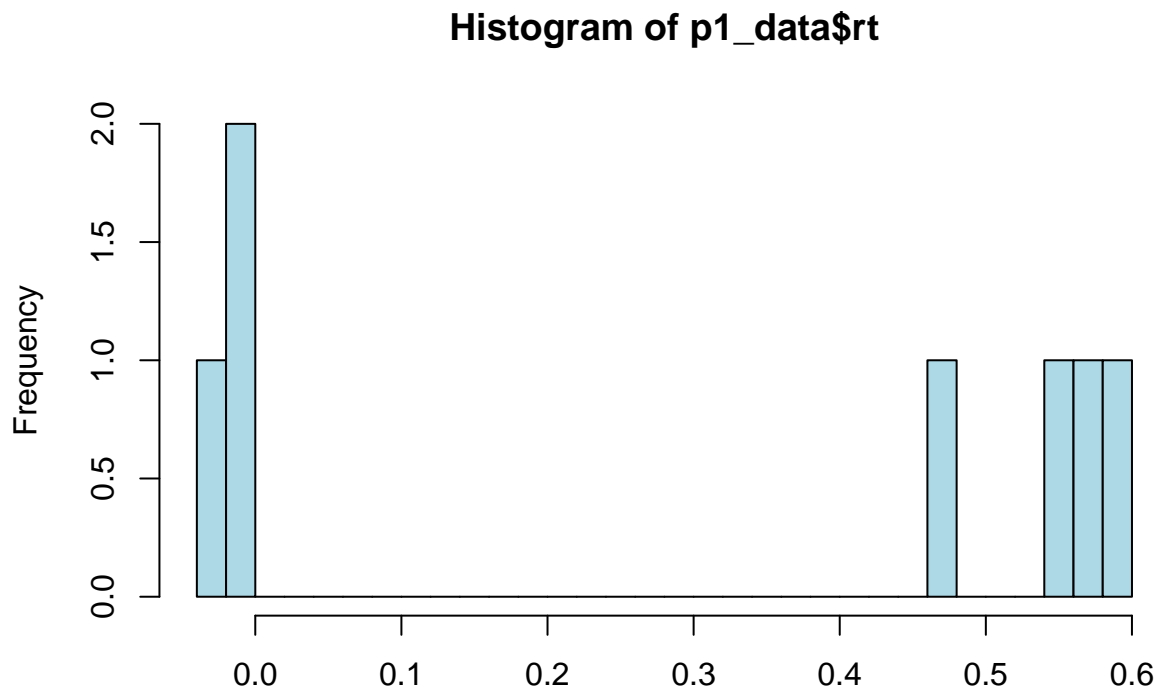
rt = reaction time in seconds (NA when participant did not respond during that trial)

stimulus = the stimulus shown on the trial

block = the block number

The `check_rt()` Function

It appears that there is something weird going on in the reaction time column:

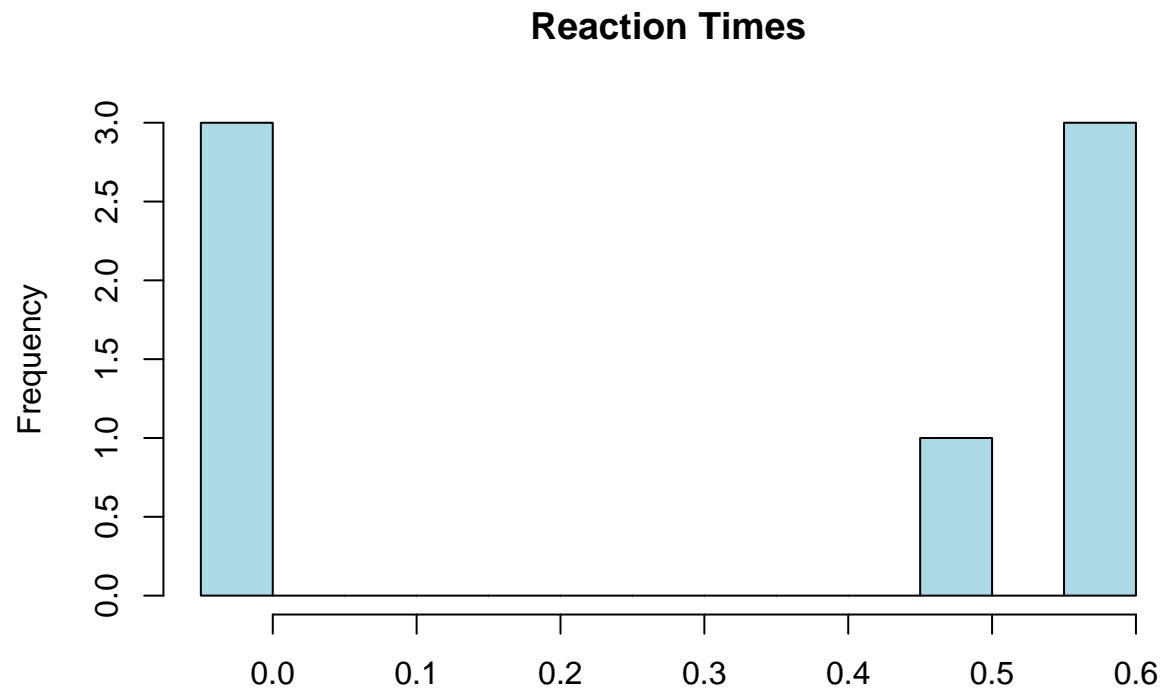


There are some observations that are very close to (or below) 0. This is the case when the participant takes too long to respond to a stimulus, and their response “leaks” into the next trial. As the function corrects for the time that it takes to run the function to gain more accurate reaction times by subtracting a constant from the reaction times, this might sometimes lead to very small or negative values.

Fortunately, the `gonogo` package comes with a function that helps spot irregularities in reaction times: `check_rt()`! The `check_rt()` function points out unusually long distances between sorted observations in your reaction time data. It has two arguments, *data* (specify your dataframe), and *ratio* (the ratio of the longest distance between sorted data points and the range of the data). See below:

```
check_rt(data = p1_data, ratio = 1/3)
```

```
## [1] "There are observations in your reaction times that are unusually far from each other: the ratio
```



End of the Tutorial!