Shiny app: plot mi-hd results

[DOI: coming soon]

This Shiny app allows you to interact with the results in the study mi-hd.

1. Installing the Shiny app as an R package

1.1 Install from GitHub

• Open an R session and run the following command:

devtools::install_github("https://github.com/EdoardoCostantini/mihd.results")

1.2 Install from a local folder

- 1. Download the package from GitHub or Zendo.
- 2. Unzip the package.
- 3. Open an R session, and run the following command.

```
install.packages(
    "path to the folder containing the package",
    repos = NULL,
    type = "source"
)

For example, on a Windows computer, this could be
install.packages(
    "path to the folder containing the package",
    repos = NULL,
    type = "source"
)
```

2. Using the Shiny app

To start the shiny apps and interact with the plots, open an R session and load the package:

```
library("mihd.results")
```

Then, run the following command in the R console:

```
start_app()
```

The app interface will explain how to interact with it.