Example Use

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4/13/2021

Introduction

This R Markdown document shows the usage of the package cino1.

```
# Install local package
install.packages("~/Documents/Masterarbeit/Package/nofone/", repos = NULL, type="source")

## Installing package into '/home/thgaertner/R/x86_64-pc-linux-gnu-library/3.6'

## (as 'lib' is unspecified)

# load package
library(cinof1)
```

Data

In this package, a sample data frame is included. It contains data for 300 patients within an n of 1 study. The data has the following structure:

- patient id: Unique patient identifier
- date: Date of data points
- day: Day in study
- Block: identifies treatment block
- Activity: Dummy variable for steps per day
- treatment: Dummy variable for 2 treatments as factors
- Uncertain_Low_Back_Pain: Dummy variable for Uncertain log back pain on scale 1-15

```
load("data/simpatdat.rda")
# Summarize Data
summary(simpatdat)
```

```
##
      patient id
                            date
                                                              treatment
                                           day
##
   Min.
           : 0.00
                    2018-01-01:
                                 20
                                                        Treatment 1:1120
                                             : 1.00
                                      Min.
##
   1st Qu.: 4.75
                    2018-01-02:
                                 20
                                      1st Qu.: 28.75
                                                        Treatment 2:1120
##
   Median: 9.50
                    2018-01-03:
                                 20
                                      Median: 56.50
   Mean
           : 9.50
                    2018-01-04:
                                 20
                                      Mean
                                             : 56.50
##
   3rd Qu.:14.25
                    2018-01-05:
                                 20
                                      3rd Qu.: 84.25
                                 20
##
           :19.00
                    2018-01-06:
                                      Max.
                                              :112.00
                              :2120
##
                    (Other)
   Uncertain_Low_Back_Pain
                                block
                                               Activity
           : 6.000
##
   Min.
                            Min.
                                   :1.00
                                           Min.
                                                  :
                                                       45.19
##
   1st Qu.: 9.000
                            1st Qu.:1.75
                                           1st Qu.: 5563.32
  Median : 9.000
                            Median:2.50
                                           Median: 6910.56
  Mean
         : 9.231
                            Mean
                                   :2.50
                                           Mean
                                                  : 6943.34
                            3rd Qu.:3.25
                                           3rd Qu.: 8290.16
   3rd Qu.:10.000
```

```
## Max. :12.000 Max. :4.00 Max. :14084.38
```

Basic Functions

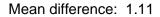
Basic functions for analyse N-of-1 studys are for example wilcox test or comparative plots. These two functions are provided in this package.

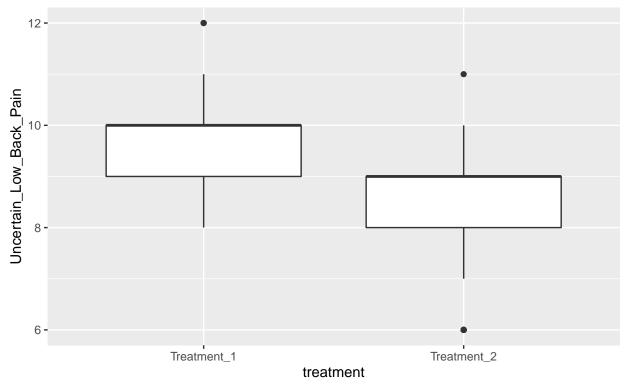
Comparative Plot

To get a first idea about the data and the difference between treatment 1 and treatment 2, a comparative plot could be used. It shows the outcome on the y-Axis against the different treatments on the x-Axis,

```
# Define outcome and exposure column
outcome <- "Uncertain_Low_Back_Pain"
exposure <- "treatment"
# Plot outcome among different exposures
comparative.plot(simpatdat, exposure = exposure, outcome = outcome)</pre>
```

Comparative Plot





Wilcox Test

To validate, that there is a difference in both treatments, the Wilcox test could be used. It calculates the p-value for the null hypothesis, that there location shift is equal to zero.

```
# Define outcome and exposure column
outcome <- "Uncertain_Low_Back_Pain"
exposure <- "treatment"</pre>
```

```
# Perform Wilcox test among different exposures
wilcox.nofone(simpatdat, exposure = exposure, outcome = outcome)

##
## Wilcoxon rank sum test
##
## data: Uncertain_Low_Back_Pain by treatment
## W = 1048691, p-value < 2.2e-16
## alternative hypothesis: true location shift is not equal to 0</pre>
```

Adjust Wash In and Wash Out

```
outcome <- "Uncertain_Low_Back_Pain"</pre>
exposure <- "treatment"</pre>
variables <- c("Activity")</pre>
id <- "patient_id"</pre>
time_col <- "day"
result <- estimate_gamma_tau(data = simpatdat, outcome = outcome, exposure = exposure, variables = vari
fit.adj.lm(data = simpatdat, outcome = outcome, exposure = exposure, variables = variables, effects = r
## Call:
## lm(formula = formula(str_formula), data = data, na.action = na.omit)
## Coefficients:
##
             (Intercept) treatment.Treatment_1
                                                                 Activity
##
               1.219e+01
                                       -1.827e+00
                                                               -8.407e-05
## treatment.Treatment_2
##
              -2.946e+00
```

Bayesian

Bayesian Networks are used to calculated the probability of outcome variables adjusted for confounders. For that, a dag is required, which identifies the relations between the variables. In this implementation, also lags are included and could be specified in the dag by adding .lag= to the variable name.

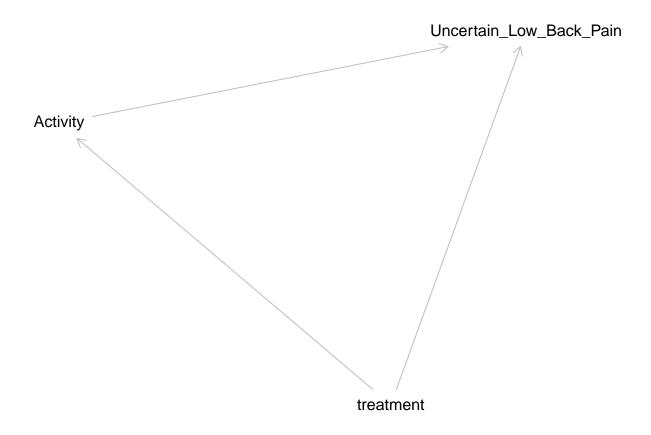
Preprocess Data

```
# specify column names
id <- "patient_id"
time_col <- "day"

# Load data
load("data/simpatdag.rda")
load("data/simpatdat.rda")

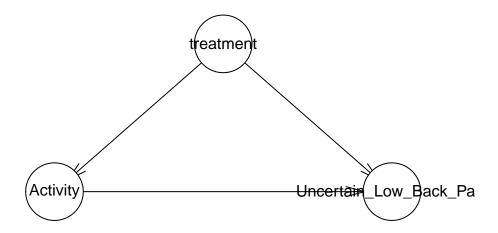
plot(simpatdag)</pre>
```

Plot coordinates for graph not supplied! Generating coordinates, see ?coordinates for how to set you



Dag preprocessing
bn.dag <- bn.prep.dag(simpatdag)</pre>

Transformed Bn DAG

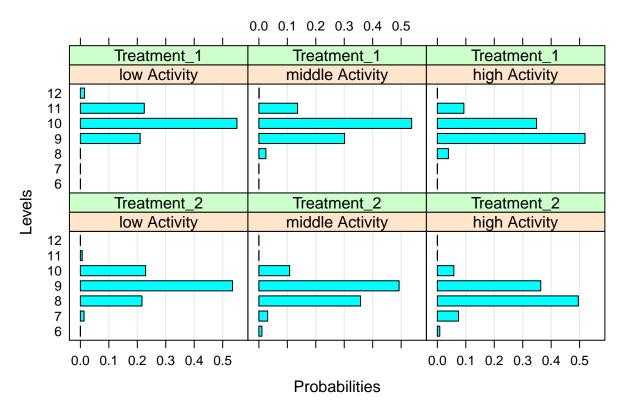


```
# Data Preprocessing (Factorization)
simpatdat$Uncertain_Low_Back_Pain <- as.factor(simpatdat$Uncertain_Low_Back_Pain)
simpatdat$Activity <- cut(simpatdat$Activity, 3, labels=c("low Activity", "middle Activity", "high Activity bn.data <- bn.prep.data(bn.dag, simpatdat, id, time_col)
bn.data <- na.omit(bn.data)</pre>
```

Fit and Plot Results

```
fitted.bn <- bn.fit.dag(bn.data, bn.dag, method="bayes")
library(bnlearn)
bn.fit.barchart(fitted.bn$Uncertain_Low_Back_Pain)</pre>
```

Conditional Probabilities for Node Uncertain_Low_Back_Pain



G-Estimation

G-Estimation is used to adjust the analysis for causal inferences. For that, three different methods are implemented

Load Data

```
load("data/simpatdat.rda")
```

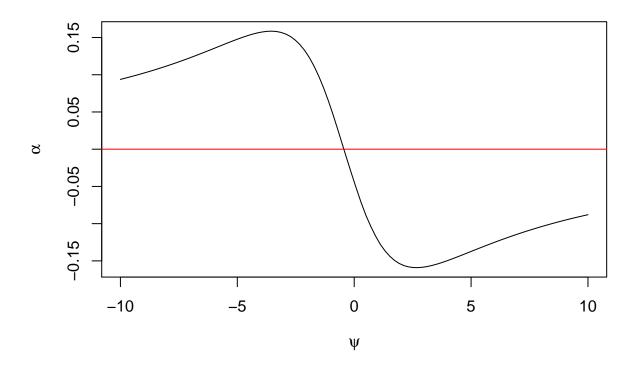
Fit G-Estimation by Iteration

It iterates over several values for ψ and returns a data frame with ψ and corresponding α

```
outcome <- "Uncertain_Low_Back_Pain"
exposure <- "treatment"
confounder <- c("Activity")
id <- "patient_id"
df <- nofgest(simpatdat, outcome, exposure, confounder, id, method="iterate", steps=100, upper_bound_ps</pre>
```

This function is useful to plot a curve for α and ψ .

Plot of $\alpha(\psi)$



Fit G-Estimation by Recursive Mean

This function approximate ψ by an interval search.

```
outcome <- "Uncertain_Low_Back_Pain"</pre>
exposure <- "treatment"</pre>
confounder <- c("Activity")</pre>
id <- "patient_id"</pre>
nofgest(simpatdat, outcome, exposure, confounder, id, method="rec_mean")
## [1] "Converged! Optimal Psi: -0.443288747439986"
## [1] "Number of iterations: 50"
## $upper_psi
## [1] -0.4432887
##
## $upper_beta
## [1] -6.491118e-16
##
## $lower_psi
## [1] -0.4432887
##
## $lower_beta
## [1] 1.881633e-16
##
## $n_it
## [1] 50
```

Fit G-Estimation by Recursive Improved

This function approximate ψ by an optimized interval search.

```
outcome <- "Uncertain_Low_Back_Pain"</pre>
exposure <- "treatment"</pre>
confounder <- c("Activity")</pre>
id <- "patient_id"</pre>
nofgest(simpatdat, outcome, exposure, confounder, id, method="rec")
## [1] "Converged! Optimal Psi: -0.443288747439985"
## [1] "Number of iterations: 8"
## $upper_psi
## [1] -0.4432887
##
## $upper_beta
## [1] -1.980295e-16
##
## $lower_psi
## [1] -0.490945
##
## $lower_beta
## [1] 0.004872248
##
## $n_it
## [1] 8
```