

Contents

We provide the details of the estimation procedure for the LTMLE and the Cheap Bootstrap/Subsampling confidence interval in the following paragraph. The data generating mechanism may be found in the **datageneratingmechanism.R** file, where each regression coefficient corresponds to a regression coefficient in a logistic regression model. Basic implementations of the Cheap Subsampling/Bootstrap confidence interval and the LTMLE are given in the **bootstrapci.R** file. The specific details of how we use this to simulate from the setup in the simulation study are given in the **getlavamodel.R** file. We now demonstrate how to use each of these files in the following R code:

```
suppressMessages(library(data.table))
library(foreach)
library(lava)
library(targets)
source("get_lava_model.R") ## the get_lava_model function and get_sim_data
source("bootstrap_ci.R") ## cheap bootstrap/subsampling confidence interval
source("data_generating_mechanism.R") ## loads data generating mechanism in
tar_source("Ltmle") ## use targets to source all of the files

## make lava model from coefficients
lava_model <- suppressMessages(get_lava_model(3, coefs))

## set up model object
mm <- list()
mm$outcome <- "heart_failure"
mm$censoring <- "Censored"
mm$comp.event <- "Dead"
mm$treatment <- "GS"
mm$time_covariates <- "insulin"
mm$time_horizon <- 3
mm$baseline <- c("sex", "education", "agegroups", "tertile_income", "index_heart")
mm$model <- lava_model

## simulate data
sample_size <- 1000
set.seed(6)
data <- get_sim_data(mm, sample_size)
data
```

pnr	sex	education	agegroups	tertile_income		
	<int>	<fctr>	<fctr>	<fctr>		
1:	1	Female	Medium	60-65		
2:	2	Male	High	60-65		
3:	3	Female	Medium	65-70		
4:	4	Female	Medium	80-85		
5:	5	Male	Medium	70-75		

996:	996	Female	High	below 45		
997:	997	Male	High	70-75		
998:	998	Female	High	45-50		
999:	999	Male	Medium	60-65		
1000:	1000	Female	Medium	50-55		
		index_heart_failure	diabetes_duration			
		<fctr>	<fctr>			
1:		No	above 10			
2:		No	5-10			
3:		No	below 5			
4:		No	5-10			
5:		No	5-10			

996:		No	above 10			
997:		No	below 5			
998:		No	5-10			
999:		No	below 5			
1000:		Yes	above 10			
		secondline_duration	first_2ndline	insulin_0	GS_0	
		<fctr>	<fctr>	<int>	<int>	
1:		above 3	other	1	1	
2:		above 3	other	0	0	
3:		below 1	other	0	0	
4:		1-3	other	0	0	
5:		1-3	other	0	1	

996:		above 3	other	1	1	
997:		above 3	other	0	0	
998:		above 3	glp1	0	0	
999:		1-3	other	0	0	
1000:		below 1	other	0	0	
		Censored_1	heart_failure_1	Dead_1	insulin_1	GS_1

	<int>	<int>	<int>	<int>	<int>
1:	1	0	0	0	0
2:	1	0	0	0	1
3:	1	0	0	0	0
4:	1	0	0	0	0
5:	1	0	0	1	0

996:	1	0	0	0	1
997:	1	0	0	0	1
998:	1	0	0	0	0
999:	1	0	0	0	0
1000:	1	1	NA	NA	NA
Censored_2	heart_failure_2	Dead_2	insulin_2	GS_2	
	<int>	<int>	<int>	<int>	<int>
1:	1	1	NA	NA	NA
2:	1	0	0	0	0
3:	1	0	0	0	0
4:	1	0	0	0	1
5:	1	0	0	1	0

996:	1	0	0	0	0
997:	0	NA	NA	NA	NA
998:	1	0	0	0	0
999:	1	0	0	0	1
1000:	1	1	NA	NA	NA
Censored_3	heart_failure_3				
	<int>	<int>			
1:	1	1			
2:	1	1			
3:	1	0			
4:	1	0			
5:	1	0			

996:	1	0			
997:	0	NA			
998:	1	0			
999:	1	1			
1000:	1	1			

We now demonstrate how we find parameter estimates with the LTMLE

for our simulated data set. The default behavior of the `prepareLtmle` is that it uses the entire history, in an additive manner. Note that the LTMLE code has been taken from an altered version of the `tmle` package (Lendle et al., 2017), which may be found here: [TMLE for breakfast](#).

```
## get data for use in Ltmle, i.e., split up separately into the outcomes, r
data <- get_sim_data(mm, sample_size, TRUE, data)
## get LTMLE estimates
x <- prepare_Ltmle(
  outcome_data = data$outcome,
  regimen_data = data$regimen,
  baseline_data = data$baseline_covariates,
  timevar_data = data$time_covariates,
  time_horizon = 3,
  censored_label = 0,
  name_outcome = "heart_failure",
  name_regimen = "GS",
  name_censoring = "Censored",
  name_competing_risk = "Dead",
  abar = list(treat=c(1,1,1), control = c(0,0,0)),
  SL.library = "glm",
  verbose = FALSE,
  gbounds = c(0,1)
)
f<-summary(do.call("Ltmle", x))
f
```

	Target_parameter	Estimator	estimate	std.err
	<char>	<char>	<num>	<num>
1:	Mean(A=1)	tmle	0.3115874	0.04845545
2:	Mean(A=0)	tmle	0.4487031	0.05120906
3:	ATE	tmle	-0.1371157	0.07003415
4:	Ratio	tmle	0.6944177	0.19167729
	lower	upper	pvalue	
	<num>	<num>	<num>	
1:	0.2166164	0.4065582972	1.272779e-10	
2:	0.3483352	0.5490710128	1.915063e-18	
3:	-0.2743802	0.0001486821	5.024867e-02	
4:	0.4769411	1.0110596389	5.709518e-02	

We also provide a basic example, showing how the Cheap Subsampling/-

Bootstrap confidence intervals may be used in practice for $B = 5$ and $m = \lfloor 0.632n \rfloor$ for the treatment contrast between those that are treated continuously throughout the period and those that are never treated in the period:

```
bs <- 5
## cheap_subsampling_ci
k_m <- 0.632
m_val <- floor(k_m * sample_size)
res_subsampling <- list()
for (b in seq_len(bs)) {
  ## subsample data of size m
  subsample <- sample(1:sample_size, size = m_val, replace = FALSE)
  formatted_data_sub <-
    lapply(data, function(x) {
      x[subsample, ]
    })
  x <- prepare_Ltmle(
    outcome_data = formatted_data_sub$outcome,
    regimen_data = formatted_data_sub$regimen,
    baseline_data = formatted_data_sub$baseline_covariates,
    timevar_data = formatted_data_sub$time_covariates,
    time_horizon = 3,
    censored_label = 0,
    name_outcome = "heart_failure",
    name_regimen = "GS",
    name_censoring = "Censored",
    name_competing_risk = "Dead",
    abar = list(treat=c(1,1,1), control = c(0,0,0)),
    SL.library = "glm",
    verbose = FALSE,
    gbounds = c(0,1)
  )
  f_temp <- do.call("Ltmle", x)
  res_subsampling[[b]] <- summary(f_temp)[,c(1:3)]
}

res_subsampling <- rbindlist(res_subsampling)

## cheap_bootstrap_ci
```

```

print ("95%_Cheap_Subsampling_CI:")
get_cheap_subsampling_ci(f[Target_parameter== "ATE", estimate], res_subsam)

res_non_parametric_bootstrap <- list()
for (b in seq_len(bs)) {
  ## subsample data of size m
  bootstrap_sample <- sample(1:sample_size, size = sample_size, replace = T)
  formatted_data_boot <-
    lapply(data, function(x) {
      temp <- x[bootstrap_sample, ]
      temp[, pnr:= 1:.N]
      temp
    })
  x <- prepare_Ltmle(
    outcome_data = formatted_data_boot$outcome,
    regimen_data = formatted_data_boot$regimen,
    baseline_data = formatted_data_boot$baseline_covariates,
    timevar_data = formatted_data_boot$time_covariates,
    time_horizon = 3,
    censored_label = 0,
    name_outcome = "heart_failure",
    name_regimen = "GS",
    name_censoring = "Censored",
    name_competing_risk = "Dead",
    abar = list(treat=c(1,1,1), control = c(0,0,0)),
    SL.library = "glm",
    verbose = FALSE,
    gbounds = c(0,1)
  )
  f_temp <- do.call("Ltmle", x)
  res_non_parametric_bootstrap[[b]] <- summary(f_temp)[,c(1:3)]
}

## cheap bootstrap ci
res_non_parametric_bootstrap <- rbindlist(res_non_parametric_bootstrap)

print ("95%_Cheap_Bootstrap_CI:")
get_cheap_bootstrap_ci(f[Target_parameter== "ATE", estimate], res_non_param)

[1] "95% Cheap Subsampling CI:"

```

```
[1] -0.33335041  0.05911894  
[1] "95% Cheap Bootstrap CI:"  
[1] -0.22397980 -0.05025168
```

References

Lendle, S. D., J. Schwab, M. L. Petersen, and M. J. van der Laan (2017, October). Ltmle: An R Package Implementing Targeted Minimum Loss-Based Estimation for Longitudinal Data. *Journal of Statistical Software* 81, 1–21.