

Bootstrap

12/10/2017

```
# lib
library(tidyverse)

## Warning: package 'tidyverse' was built under R version 3.4.2
## -- Attaching packages ----- tidyverse 1.2.1 --
## <U+221A> ggplot2 2.2.1      <U+221A> purrr  0.2.4
## <U+221A> tibble  1.3.4      <U+221A> dplyr  0.7.4
## <U+221A> tidyr   0.7.2      <U+221A> stringr 1.2.0
## <U+221A> readr   1.1.1      <U+221A> forcats 0.2.0
## Warning: package 'tidyr' was built under R version 3.4.2
## Warning: package 'purrr' was built under R version 3.4.2
## Warning: package 'dplyr' was built under R version 3.4.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()

library(boot)

# load the data
data_bootstrap = read.csv("../data/data_no_outliers.csv") %>%
  select(-X, -postalcode, -facilityname, -facilityzip, -patientid, -visitid, -date) %>%
  mutate(is30dayreadmit = as.factor(is30dayreadmit), icu_flag = as.factor(icu_flag)) %>%
  select(log_length_of_stay,
         is30dayreadmit,
         cindex,
         evisit,
         age,
         gender,
         maritalstatus,
         temperature_cat,
         insurancetype,
         heartrate_transformed,
         respirationrate_transformed,
         mean_arterial_pressure)

# final model
# model_sw_0.10 = lm(log_length_of_stay ~ is30dayreadmit + cindex
#                   + evisit + age + gender + maritalstatus + temperature_cat
#                   + insurancetype + heartrate_transformed
#                   + respirationrate_transformed
#                   + mean_arterial_pressure)

# bootstrap to obtain the distribution

# code 1: raw code
# boot_res1 =
```

Table 1: Bootstrap

| term | statistic | bias | std.error |
|-----------------------------|---------------|------------|-------------|
| (Intercept) | 2.5444567 | -0.0002083 | 0.1920116 |
| is30dayreadmit1 | 0.1801712 | -0.0002694 | 0.0421892 |
| cindexmoderate | 0.1323495 | -0.0001276 | 0.0476890 |
| cindexnormal | 0.0251459 | 0.0000472 | 0.0345832 |
| cindexsevere | 0.1775025 | -0.0006796 | 0.0428682 |
| evisit | 0.0642347 | 0.0001129 | 0.0095534 |
| age | 0.0096015 | 0.0000127 | 0.0009287 |
| genderMale | 0.0611153 | 0.0003365 | 0.0295831 |
| maritalstatusNot Married | 0.0885007 | -0.0000767 | 0.0293569 |
| temperature_catlow | -0.0593297 | 0.0005501 | 0.0906106 |
| temperature_catnormal | -0.2456611 | 0.0003272 | 0.0584033 |
| insurancetypeMedicare | -0.1447609 | -0.0009221 | 0.0722351 |
| insurancetypePrivate | -0.1855449 | -0.0002085 | 0.0681488 |
| heartrate_transformed | -2061.1985080 | 0.0079740 | 301.2852763 |
| respirationrate_transformed | -178.7334811 | -0.0165204 | 30.1096161 |
| mean_arterial_pressure | -0.0058527 | -0.0000044 | 0.0008342 |

```

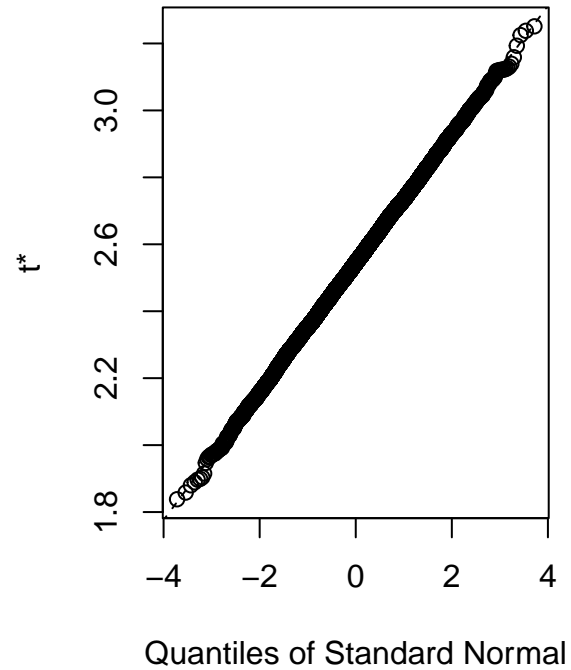
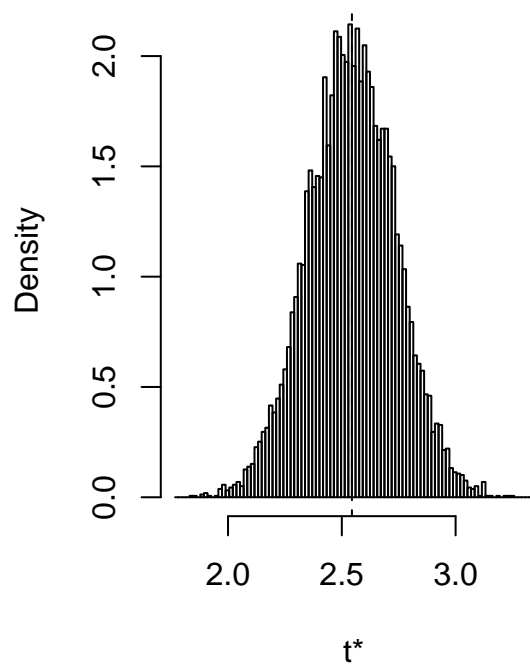
# data_bootstrap %>%
# modelr::bootstrap(n = 10000) %>%
# mutate(models = map(strap, ~lm(log_length_of_stay ~ ., data = .x) ),
#        results = map(models, broom::tidy)) %>%
# select(-strap, -models) %>%
# unnest() %>%
# select(id = `.id`, term, estimate) %>%
# spread(key = term, value = estimate)

# code 2: boot function
boot.fn = function(data, index){
  return(coef(lm(log_length_of_stay ~ ., data = data, subset = index)))
}
boot_res2 = boot(data_bootstrap, boot.fn, 10000)
# table for bootstrapping results
broom::tidy(boot_res2) %>%
  knitr::kable(format = "latex",
               caption = "Bootstrap",
               booktabs = TRUE) %>%
  kableExtra::kable_styling(latex_options = "scale_down")

# plot
plot(boot_res2)

```

Histogram of t



```
# CI
boot.ci(boot.out = boot_res2, type = c("norm", "basic"))

## BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS
## Based on 10000 bootstrap replicates
##
## CALL :
## boot.ci(boot.out = boot_res2, type = c("norm", "basic"))
##
## Intervals :
## Level      Normal          Basic
## 95%   ( 2.168,  2.921 )  ( 2.167,  2.926 )
## Calculations and Intervals on Original Scale
```